

**The effects of diet-induced obesity on metabolic
and vascular functions: role of insulin signalling
and insulin resistance**

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University of Liverpool for the degree of Doctor in Philosophy**

By

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Declaration

I declare that the work presented in this thesis is all my own and has not been previously submitted for any other degree.

Sameer Fatani

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I would like to dedicate the whole of this work and thesis to my great mother and honest wife "Amal Bafageeh", the two persons whom encouraged me by all means to finish this work successfully.

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Abstract

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High energy diets and diet-induced obesity are known to induce detrimental effects on various physiological functions. However, insulin resistance is postulated to play a central role in mediating and linking these detrimental effects with related clinical symptoms which include cardiovascular disorders. Therefore, the present study aimed to investigate the relationship between insulin resistance and "different negative effects and pathological signs" in diet-induced obese rats.

Long term feeding of a highly-palatable diet induced obesity, with concomitant alterations in metabolic, vascular, and molecular functions, namely insulin resistance and attenuated endothelial-dependent vasorelaxation. At the molecular level, several abnormalities were detected in protein expression levels of both PI 3-kinase and MAP-kinase insulin signalling pathways in vascular, hepatic and skeletal muscular tissues. In all tissues the greater changes were in the level of PI 3-kinase. These defects in PI 3-kinase levels may play a crucial role in inducing local and systemic insulin resistance, and subsequent pathological symptoms.

Consumption of all types of high energy diets in the present study (such as biscuit, chocolate or high-fat diets) induced vascular dysfunctions mainly the attenuation of endothelium-dependent vasorelaxation. Moreover, such diets did not induce obesity or metabolic changes, but were associated with a changed lipid profile, in particular hypertriglyceridaemia, suggesting a role for triglyceride in dietary-induced vascular abnormality.

Chronic withdrawal of the highly-palatable diet reversed body weight, and was associated with normalisation of metabolic, vascular, and molecular abnormalities. Although, fenofibrate treatment caused a remarkable reduction in body weight and metabolic improvement, it had very limited beneficial effects on vascular function.

In conclusion, highly-palatable diet-induced obesity is associated with metabolic and vascular related clinical symptoms. Molecular defects in insulin signalling pathways could participate in the induction of such pathological signs. Principally, PI 3-kinase seems to be the key controlling step of insulin transduction, having a major role in inducing local insulin resistance and the subsequent relevant clinical symptoms. Moreover, the augmented levels of circulating and tissue accumulation of triglyceride in diet-induced obese animals could have a central role in mediating the local and systemic insulin resistance. Long term consumption of a high calorie diet irrespective of developing obesity, appears to be the major causative factor in inducing vascular defects leading to cardiovascular disease. Physiological but not pharmacological management of obesity appears to be more effective in correcting vascular abnormalities associated with dietary obesity.

Abbreviations

| | |
|------------------|--|
| ADP | Adenosine-Di-Phosphate |
| AMP | Adenosine-5'-Monophosphate |
| ATP | Adenosine-Tri Phosphate |
| BCA | Bicinconinic Acid |
| BMI | Body Mass Index |
| C | Chow-fed group |
| CCh | Carbamylcholine |
| CCRC | Cumulative Concentration Response Curve |
| CNS | Central Nervous System |
| CRP | C-Reactive Protein |
| DC | Diet-to-Chow group |
| EC ₅₀ | Effective Concentration ₅₀ |
| ELISA | Enzyme-Linked ImmunoSorbent Assay |
| ET-1 | Endothelin-1 |
| eNOS | Endothelial Nitric Oxide Synthase |
| ERK1/2 | Extracellular Regulated Kinase 1/2 |
| F | Fenofibrate |
| FBF | Forearm Blood Flow |
| FFA | Free Fatty Acid |
| G6P DH | Glucose-6-Phosphate Dehydrogenase |
| GRB-2 | Growth Factor Receptor-protein Bound-2 |
| GTP | Gaunosine-Tri-Phosphate |
| HK | Hexokinase |
| HOMA | Homeostasis Model Assessment |
| HPD | Highly-Palatable Diet |
| HRP | Horse-reddish Peroxidase |
| IGT | Impaired Glucose Tolerance |
| IL-6 | Interlukin-6 |
| IR- α | Insulin Receptor- α |
| IR- β | Insulin Receptor- β |
| IRS-1 | Insulin Receptor-Substrate-1 |
| IRS- 2 | Insulin Receptor-Substrate-2 |
| KPSS | Potassium Physiological Salt Solution |
| LCD | Low Calorie Diet |
| LDL | Low-Density Lipoprotein |
| MAP | Mitogen Activated-Protein |
| NA | Nor-Adrenaline |
| NAFLD | Non Alcoholic Fatty Liver Disease |
| NBF | Neutral Buffered Formalin |
| NEFA | Non-Esterified Fatty Acid |
| NIDDM | Non-Insulin Dependent Diabetes Mellitus |
| NO | Nitric Oxide |
| PAI-1 | plasminogen activator inhibitor-1 |
| PBS | Phosphate Buffered saline |
| PI 3-kinase | Phosphatidylinositol 3-kinase |
| PKC- β | Protein Kinase C- β |
| POD | Peroxidase |
| PPAR- α | Peroxisome Proliferator Activated Receptor- α |
| PSS | Physiological Salt Solution |
| RIA | Radio-Immuno Assay |
| RER | Rough Endoplasmic Reticulum |
| ROS | Reactive Oxygen Species |

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|---------------|--|
| SEM | Standard Error of Mean |
| SDS-PAGE | Sodium-Dodecyl-Sulfate Poly-Acrylamide Gel Electrophoresis |
| SNP | Sodium Nitroprusside |
| TBW | Total Body Water |
| TG | Triglyceride |
| TBHB | 2,4,6-tr-bromo-3-hydroxy-benzoic acid |
| TNF- α | Tumour-Necrosis Factor- α |
| TPS | Tris Phosphate Saline |
| UV | Ultra-Violet |
| VLCD | Very low Calorie Diet |
| V | Vehicle group |
| VSMC | Vascular Smooth Muscle Cells |
| V / V | Volume / Volume |
| W / V | weight / Volume |
| WAT | White Adipose Tissue |
| WHO | World Health Organization |
| ZDF | Zucker Diabetic Fatty rats |

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Chapter 1

Introduction

1.1 INTRODUCTION

Obesity is a disease of the life style that was largely neglected. In 1997, the world health organization (WHO), issued a statement indicating that "obesity's impact is so diverse and extreme that it should now be regarded as one of greatest neglected public health problems of our time (WHO, 1997). In fact, overweight and obesity is forecast to increase by 50% by 2015 according to a new report from the WHO. Obesity is a worldwide epidemic (James *et al.*, 2001) that is characterized by excess adipose tissue contributing to numerous chronic diseases (Must *et al.*, 1999) and early mortality (Simopoulous and Van Itallie, 1984; Kushner, 1993). This epidemic has received both national and international attention because of its increasing prevalence, its detrimental impact on health, and subsequent enormous economic burden it imposes (Wolf and Colditz, 1998).

Obesity is a chronic (NHLBI, 1998; WHO, 1998) condition that develops as a result of an interaction between a person's social, behavioral, cultural, psychological, metabolic, and genetic factors (Stunkard *et al.*, 1986; WHO, 1998). It causes or exacerbates many health problems both independently and in association with other diseases. In particular, it is associated with the development of coronary heart disease (CHD), non-insulin-dependent diabetes (NIDD), some forms of cancers (Mark *et al.*, 1999), non-alcoholic fatty liver disease (NAFLD), in addition to many other chronic diseases. In 1998, the American Heart Association issued a "call to action" and reclassified obesity as a major modifiable risk factor for CHD (Eckel and Krauss, 1998). Increased intra-abdominal fat, or waist circumference, as a consequence of obesity is related to a constellation of risk factors, the so-called insulin resistance syndrome. Therefore, breakthroughs in our understanding of the molecular mechanisms of insulin resistance and endothelial dysfunction in vasculature may provide potential opportunities for therapeutic intervention and bring renewed hope and vitality to the management of obesity and its complications.

1.2 OBESITY

1.2.1 Definition and measurement

Obesity, defined as excess adiposity for a given body size, results from an imbalance between energy intake and energy expenditure (Kuczmarski *et al.*, 1997). It is assumed that both a predisposition to excess energy intake and a reduced resting metabolic rate contribute to obesity (Ravussin *et al.*, 1988; Roberts and Leibel, 1998). The concept of energy balance is easy to understand. A typical obese patient will have increased body weight by some 20 kg over 10 years. This means a daily energy excess of 30-40 kcal per day at the start of the development of obesity, which gradually has to be increased to maintain the enlarged body mass. The positive energy balance needed for obesity to develop over the years is so small that the patient does not notice it (Bjorntorp, 1997).

Weight gain occurs differently in men and women. In women, the greatest weight gain is in the younger age groups, while, men generally develops obesity at later in life. Furthermore, as women's educational level raises, their obesity decreases, whereas in men, educational level appears not to be related to obesity (Krauss *et al.*, 1998).

The most commonly used method today for classifying an individual as overweight or obese is based on body mass index (BMI), a value that is measured as body weight in kilograms over the square of height in meters (kg/m^2). In adults, BMI between 20 and 25 is considered normal, individuals with a BMI between 25 and 30 are considered overweight and those with a BMI over 30 are obese (Kuczmarski *et al.*, 1997; NHLBI, 1998), regardless of sex. (Table 1.1).

Table 1.1 World health organization's Body Mass Index categories based on increasing health risks.

| BMI (kg/m ²) | WHO classification | popular description |
|--------------------------|--------------------|-----------------------------|
| < 18.5 | Underweight | Thin |
| 18.5-24.9 | Normal | Healthy normal "acceptable" |
| 25.0-29.9 | Grade 1 overweight | Overweight |
| 30.0-39.9 | Grade 2 overweight | Obesity |
| ≥ 40.0 | Grade 3 overweight | Morbid obesity |

The use of BMI to assess weight-related health risk has gained international acceptance because of the associations between BMI and adiposity (Gallagher *et al.*, 2000), BMI and disease risk (James *et al.*, 2001), and BMI and mortality (Calle *et al.*, 1999). BMI values provide valuable information about increasing body fatness. It allows meaningful comparisons of weight status within and between populations and the identification of individuals and groups at risk of morbidity. It also permits identification of priorities of intervention at an individual or community level and for evaluating the effectiveness of such interventions (WHO, 1995; WHO, 1997).

Waist circumference is another clinically feasible measurement that may be used independently or in addition to BMI (Janssen *et al.*, 2002) to assess weight-related health risk. Although, BMI and waist circumference are the recommended (USDHHS, 1998) and most clinically feasible means of identifying patients who are overweight or obese in clinical practice, there are several other methods used in clinical practice to assess body adiposity such as: (1) Skinfold test; a simpler test for measuring body fat, in which a pinch of skin is precisely measured by calipers at standardized points on the body to determine the thickness of the subcutaneous fat layer, and converted to body fat percentage by an equation; (2) Bioelectrical impedance analysis (BIA). is

a commonly used process for estimating body composition. It has come into favor in recent years due to its role in determining an estimate of total body fat. BIA actually determines the electrical impedance of body tissues, providing an estimate of total body water (TBW). TBW in turn, can be used to estimate fat-free body mass and subsequent body fat (WHO, 1995; WHO, 1997).

1.2.2 Prevalence and epidemiology

Obesity should no longer be regarded simply as a cosmic problem affecting certain individuals, but as epidemic that threatens global well being. It is affecting an estimated 300 million people worldwide (WHO, 2002). Its prevalence is increasing in both developed and developing countries throughout the world. In the United States, the prevalence of obesity is greater than it has ever been, with striking increases observed during the past 2 decades (Flegal *et al.*, 1998). Approximately, 64.5 % of adult population in the United States is estimated to be overweight in the period 1999-2000 as compared with 46 % in 1976 to 1980. Most of this increase is attributable to a dramatic rise in the prevalence of obesity from 15% to the recent estimate of nearly 31%.(Flegal, 2002). Obesity is pervasive, affecting people of all ages and at all socioeconomic level. Moreover (Mokdad *et al.*, 1999), in addition to age and sex, race and ethnicity are important factors in the prevalence of obesity (Williamson, 1993; Lovejoy *et al.*, 1996; USDHHS, 2000). The prevalence of obesity increased from 12% in 1991 to 18% in 1998. Increases were seen in both sexes and all socioeconomic classes, with the greatest increase seen in 18-29 year olds (Mokdad *et al.*, 1999). Conservative estimates of the economic costs of obesity in developed countries are between 2 and 7 % of the total health costs, which represents a significant expenditure of national health-care budgets (Seidell, 1996).

Increasing age is associated with an increase in obesity. Body weights of men and women in the United states increase approximately 9.1 kg between the ages of 25 and 55 years (Belanger *et al.*, 1988; Wang *et al.*, 1994). These increases in body weight generally are not explained by increases in fat-free mass, because bone mass peaks around 30 years of age and muscle mass

plateaus and later declines unless the individual engages in the strengthening activities. These changes in body weight and body composition are attributable, in part, to the natural declines in growth hormone, dehydroepandrosterone, and testosterone with aging. In addition, reductions in resting metabolism alter energy balance and contribute to weight gain.

In Europe, on average, 15% of men and 22% of women are considered to be obese, with overweight also being more common among women than men. More than half the adult population between 35 and 65 years of age were either overweight or obese. In England and Wales, the most recent health survey has confirmed an increase in the prevalence of obesity in adults from 6% in men and 8% in women in 1980 to 23% of men and 25% of women in 2002 (Prescott-Clarke and Primatesta, 1998; Rennie and Jebb, 2005). Furthermore, the increasing prevalence of obesity is not confined merely to Europe and America. In Southeast Asia a marked rise is being seen in all populations, and in Japan and China a pronounced increase in the prevalence of overweight and obesity has been observed during the past two decades. The most striking figures come from the Pacific region. In urban Samoa the prevalence of obesity is estimated as greater than 75% of adult women and 60% of adult men (Hodge *et al.*, 1995). High prevalence rates also occur in the Middle East, in the United Arab Emirates is recognized as a major public-health problem that maybe an important factor in the increasing occurrence of other chronic diseases (Musaiger, 1996).

1.2.3 Etiology of obesity

Overweight and obesity result from the interaction of many factors, including genetic, behavioural, environmental, physiological, social and cultural influences that result in energy imbalance and promote excessive fat deposition. The relative contribution of each of these factors has been studied extensively, and although genes play an important role in the regulation of body weight, the WHO consultation on obesity (WHO, 1998) conducted that behavioral and environmental factors (i.e., sedentary lifestyles combined with excess energy intake) are primarily responsible for the dramatic increase in obesity during the past two decades. Moreover, these factors may interact

with genetic susceptibility to produce obesity (Bouchard and Tremblay, 1990). Increasing energy consumption, decreasing energy expenditure, or a combination of both leads to a positive energy balance and consequently increasing body mass of the individuals promoting induction of obesity.

1.2.4 Factors influencing obesity

1.2.4.1 Genetic susceptibility

Although, obesity runs in families, the influence of genotype on the etiology of obesity may be attenuated or exacerbated by nongenetic factors. Apart from rare obesity-associated syndromes, the genetic influences seem to operate through susceptible genes. Such genes increase the risk of developing obese characteristics but are not essential for their expressions or, by themselves, sufficient to explain the development of the disease. Moreover, it is estimated that approximately 40% of the variance in daily energy expenditure (excluding vigorous physical activity) is attributable to genotype (Bouchard and Tremblay, 1990). Thus, there is substantial evidence implicating the role of genetics in body weight regulation. These suggest that differences in genetic susceptibility within population determine those who are most likely to become obese in any given set of environmental circumstances (Bouchard *et al.*, 1990). In other words; implicit to the susceptible-gene hypothesis is the role of environmental factors that unmask latent tendencies to develop obesity.

1.2.4.2 Environmental factors

Energy expenditure. The most variable component of energy expenditure is physical activity, representing 20-50% of total energy expenditure (Prentice *et al.*, 1996). In developed countries there is a relationship between low level of physical activity and obesity (Rissanen *et al.*, 1991). Longitudinal studies of Pima Indians indicate that the risk of 10-kg weight gain during a 4-year follow-up is a seven-fold higher in those in the lowest tertile of relative resting metabolic rate, compared with those in the highest tertile (Ravussin *et al.*, 1988).

Energy intake. It seems that environmental influences act through increasing energy intake and / or decreasing energy expenditure. There is some

evidence that high-fat diets are associated with an increased risk of obesity within populations, but cross-cultural studies have failed to show any consistent relationship between nutritional factors and relative weights (Blundell and Macdiarmid, 1997).

Culture. Evidence for the critical role of environmental factors in the development of obesity comes from the migrant studies and the 'westernisation' of diet and lifestyles in developing countries. Study of Nigerians living in the United States, found that mean BMI for men and women in Nigeria is 21.7 and 22.6, respectively, while those living in United States have an average BMI of 27.1 for men and 30.8 for women. Moreover, increasing prevalence of obesity is associated with adverse health consequences such as hypertension (adult Nigerian living in Africa is 15%, whereas it is as high as 30% among those living in United States) (Wilks, 1996).

In industrialised countries, a higher prevalence of overweight and obesity is observed in those with lower educational attainments and low income, although the reverse may be seen in developing countries. There is tendency for overweight to increase after marriage and with increasing parity. Hence, a balanced dietary intake and physical activity are crucial in increasingly affluent societies (Kuczmarski *et al.*, 1994).

1.2.5 Complications of obesity

Even small increases in weight across a population can have devastating impact on public health. Close to 300,000 deaths each year in the United States may be attributable to obesity (Allison *et al.*, 1999). Excess weight increases the risk of multiple pathological conditions, including cardiovascular disease, type 2 diabetes, non-alcoholic fatty liver disease (NAFLD), hypertension, certain forms of cancer, cerebrovascular disease, gallstones, osteoarthritis, dyslipidemia, sleep apnea, degenerative arthritis (Mark *et al.*, 1999), asthma (USDHHS, 2001), cataracts (Weintraub *et al.*, 2002), benign prostatic hypertrophy (Giovannucci *et al.*, 1994), menstrual irregularities, and pregnancy complications resulting in reduced life expectancy. Moreover,

obesity may lead to depression and social discrimination (USDHHS, 2001), having negative effects on physical functioning, vitality (Coakley *et al.*, 1998), and general quality of life (Fine *et al.*, 1999). The adverse health consequences occur not only in individuals who are in the overweight and obese categories, but disease risk also starts to increase even for those at the upper end of the normal range of BMI (22.0-24.9) (Flegal *et al.*, 1998).

Increasing body fat mass is accompanied by profound changes in physiological function. These changes are, to a certain extent, dependent on the regional distribution of adipose tissue. Generalized obesity results in alterations in total blood volume and cardiac function, whereas the distribution of fat around the thoracic cage and abdomen restricts respiratory excursion and alters respiratory function. The intra-abdominal visceral deposition of adipose tissue, which characterises upper body obesity, is a major contributor to the development of hypertension, elevated plasma insulin concentrations and insulin resistance, diabetes mellitus and hyperlipidaemia.

Obesity is an important determinant of cardiovascular disease. Moreover, there is strong link between obesity and a generalised metabolic disorder of which insulin resistance is an indicator. It is difficult to define the precise contribution of obesity to insulin resistance, but most analyses suggest that it can account for $\geq 50\%$ of the variance in insulin sensitivity in the general population. Insulin resistance is associated with a constellation of metabolic abnormalities, including obesity, diabetes, dyslipoproteinaemia, hypertension, and atherosclerosis. The complex nature of insulin resistance, and its relation to atherogenesis is not fully understood yet. Therefore further studies are needed to explore whether insulin resistance can promote atherosclerosis independently of other risk factors.

1.2.6 Treatment of obesity

The goals of obesity treatment are to achieve and then to maintain clinically meaningful weight loss, with ultimate goal of reducing the risk for or severity of obesity-related diseases, impairments, and functional limitations. Weight losses of 5% to 10% of initial body weight produce health benefits and are deemed by many health care practitioners to represent a clinical success. Long-term success, however, is dependent on maintenance of a 10% weight loss for at least 1 year (Wing and Hill, 2001). Wing and Hill estimated that approximately 21% of adults who are overweight or obese are successful at 1 year, but longer-term success generally is lower (Wing and Hill, 2001). Effective therapeutic regimens for treating obesity should incorporate multiple approaches to encourage behavioural change or modification and creative strategies to facilitate consistent and long term follow-through. Numerous options are available today, (Rosenbaum *et al.*, 1997; NAASO, 2000) including low-energy diets, physical activity/exercise, behavioural modification, (Wadden and Foster, 2000), pharmacotherapy, (Klein, 1999; Bray and Tartaglia, 2000) and surgery (NIH, 1991). The treatment choice depends on the degree of obesity, the presence of co-morbidities, previous weight loss therapies utilised and the relative success of each, and specific characteristics of an individual's personal life.

1.2.6.1 Behavior and lifestyle modifications:

Behavior modification is an important component of all weight loss programmes (Wadden and Foster, 2000). Behavior strategies frequently are targeted toward identifying stimuli that signal unhealthy behaviours (e.g., binge eating), learning about the role of readiness, in initiating or continuing positive behaviours (Prochaska and Velicer, 1997) and recognising barriers that may compromise healthy pursuits. Most of the various lifestyle approaches have several factors in common, which includes: the use of self-monitoring and goal setting; stimulus control; modification of eating style habits; the use of reinforcement for healthy behaviours; nutrition education; moderate physical activity; and cognitive restructuring, including stress management, relaxing skills, meditation, and relapse-prevention training. In

general, these approaches produce moderate weight loss and have minimal side effects. They are most useful for individuals with mild obesity (BMI of 27 to 30). Individuals who follow such approaches to weight loss maintain on average about two thirds of their initial weight loss 12 months after treatment termination (Krauss *et al.*, 1998). Although no single theoretical framework for behavioural intervention has been shown to be superior, success in modifying patterns of eating and physical activity generally is dependent on consistency, support, and long-term modification of lifestyle, rather than on one specific diet or exercise program (Racette *et al.*, 2003).

1.2.6.2 Diet:

Dietary approaches form the basis of most weight loss interventions and rely on a reduction in total energy intake. Although many diets focus on dietary fat reductions, the main determinant of weight loss is the total energy content of the diet (Kennedy *et al.*, 2001) (relative to total energy expenditure), rather than the composition of macronutrients. Therefore, whether diet therapy is based on a high complex-carbohydrate, low-fat meal plan, as advocated by the American Heart Association (Krauss *et al.*, 2000) and many nutrition professionals, (NAASO, 2000; St Jeor *et al.*, 2001) or relies primarily on carbohydrate restriction, its success is dependent on a relative energy deficit. Weight-reducing diets may be very-low-calorie diets (VLCDs, 800 kcal/d) or low-calorie-diets (LCDs, 800/1500 kcal/d) and may consist of liquid formulas, pre-packaged meals, nutritional bars, regular foods, or a combination. Very-low-calorie diets are very effective for weight reduction, (Saris, 2001) and the nutritionally adequate formulas used today have fewer associated health problems (e.g., gout, cholelithiasis, hair loss) as compared with the VLCD formulas used in the 1970s.

Despite rapid and clinically meaningful weight loss with VLCDs, however, maintenance of the reduced weight is variable and generally poor. Instead, LCDs are used more commonly because they are safer, have fewer side effects, enable better adherence, and therefore may result in comparable weight loss as VLCD regimens (Rossner and Flaten, 1997). Diets containing more than 1,200 kcal/day produce slower weight losses, but they are

advantageous because they can be incorporated more easily into individual lifestyles and generally can be followed for longer periods of time without adverse health effects (Wing, 2001).

1.2.6.3 Exercise and physical activity:

Exercise generally does not produce considerable weight loss when used independently, but is a very important adjunct to a weight-reducing diet (Jakicic *et al.*, 1999) because it increases energy expenditure, enhances loss of adipose tissue, (Racette *et al.*, 1995) and improves dietary adherence. (Racette *et al.*, 1995). Although aerobic exercise has been used most frequently for weight loss and control because of the caloric expenditure required, strength training has numerous benefits and may help to preserve fat-free mass during diet-induced weight loss (Geliebter *et al.*, 1997). In addition to formal exercise, daily physical activity plays critical role in energy balance, (Saltzman and Roberts, 1995) weight control, disease prevention, and achievement and maintenance of overall health. Prospective studies indicate that physical activity protects against obesity, (DiPietro *et al.*, 1998) specifically influencing abdominal obesity, (Tremblay *et al.*, 1990) and reduces the incidence of several diseases associated with obesity (such as cardiovascular disease, (Leon *et al.*, 1997; Blair *et al.*, 2001) type 2 diabetes (Eriksson and Lindgarde, 1991; Helmrich *et al.*, 1991), depression, (Hughes, 1984; Taylor *et al.*, 1985) and premature death (Leon *et al.*, 1997; Lee and Paffenbarger, 2000). Furthermore, exercise adherence and habitual physical activity *are the* greatest determinants of weight maintenance following weight loss (Andersen *et al.*, 1999; Jakicic *et al.*, 1999). The effects of physical activity on fitness and health are dramatic (Blair *et al.*, 2001) and there is evidence that physically active individuals who are obese have a lower risk for morbidity and mortality than sedentary individuals of normal weight. (Bray, 2000).

1.2.6.4 Pharmacotherapy:

Pharmacological agents may be used in conjunction with diet, exercise, and behavioural strategies when non-pharmacological approaches alone fail to produce or sustain meaningful weight loss. Several appetite suppressant

drugs are approved for weight loss, (Bray, 2000; Yanovski and Yanovski, 2002) but the treatment duration for most is limited to twelve weeks or less. The chronic nature of obesity necessitates longer-term therapy, because drug cessation usually leads to weight regain. Two drugs approved for the long-term treatment of obesity include:

Orlistat (Xenical): is indicated with meals (120 mg) and acts to reduce the absorption of fatty acids by inhibition of triglyceride hydrolysis through its action as a gastric and pancreatic inhibitor (Hill *et al.*, 1999; Finer *et al.*, 2000). Orlistat proved more effective than diet alone for weight loss, with improvements in total cholesterol, low density lipoprotein and the low density lipoprotein to high density lipoprotein ratio and glycemic control. Orlistat represents an overall safe treatment for obesity, given that the drug is minimally absorbed. Side effects of Orlistat include malabsorption of fat-soluble vitamins and steatorrhea.

Sibutramine (meridia): acts on CNS and exerts its effects by acting as a norepinephrine, serotonin, and dopamine reuptake inhibitor (Luque and Rey, 1999). Sibutramine treatment is indicated for weight loss, and applied in combination with reduced-calorie diet. Daily doses of 10 mg and 15 mg sibutramine lead to sustained weight loss over a period of about six months. Sibutramine is contra-indicated in patients with poorly controlled hypertension and patients with a history of cardiovascular heart disease (weber, 1999).

It is worth noting that there are many other compounds and hormones in clinical trials such as: Axokine, β 3-adrenergic agonists, cannabinoid receptor antagonists, and leptin.

1.2.6.5 Surgery:

Surgery is reserved for cases of extreme obesity (BMI of $\geq 40 \text{ kg/m}^2$) or for more moderate obesity (BMI of $\geq 35 \text{ kg/m}^2$) when obesity-related comorbidities are present. The most commonly surgical procedure is the gastric bypass, in which the upper portion of the stomach is stapled to create a small (10-30 ml) reservoir (Deitel and Shikora, 2002). The net result is a substantial weight loss within 6 months. Weight losses of 45 kg (100 lb) (Pories *et al.*,

1995), or 60 to 70% of excess body weight (Sugerman *et al.*, 1987), have been observed 1 year after gastric bypass, and large losses have been maintained for up to 15 years (Pories *et al.*, 1995). Vertical banded gastroplasty is another, less commonly used surgical procedure in which a band constricts the upper portion of the stomach, effectively reducing its capacity. In contrast to the surgical procedures used in the past decades, the current techniques have low mortality rates of approximately 1.3% (Melinek *et al.*, 2002) to 1.5% (Pories *et al.*, 1995). However, morbidity associated with wound infections, incisional hernia, and anastomotic leak with peritonitis may be higher, particularly when additional surgery is necessary. Additional risks of surgical treatment for obesity include steatorrhea, vitamin and mineral deficiencies, and osteoporosis (Sugerman, 2000).

1.3 INSULIN AND INSULIN RESISTANCE

With obesity and diabetes epidemic proportions in the developed world (WHO, 2000), the role of insulin resistance and its sequelae is gaining prominence. Understanding the roles of insulin across a wide range of physiological process and the influences on its synthesis and secretion, alongside its actions from the molecular to the whole body level, has significant implications for much chronic disease seen in westernised populations today. Therefore, the study of insulin and insulin resistance remain in the forefront of medical research, relevant at all levels from bench to bedside to public health policy.

1.3.1 The discovery of Insulin

In 1889 two Germans, Minkowski and Mering noted, from their experimental work with animals, that total pancreatectomy led to the development of severe diabetes (Bliss, 1993). They hypothesized that a substance secreted by the pancreas was responsible for the metabolic control. In 1921, Banting and McLeod, administered chilled saline extracts of pancreas intravenously to dogs rendered diabetic by pancreatectomy and observed lowering of blood glucose. Insulin was finally isolated, purified and made available in a form capable of therapeutic administration, which lead to the award of Nobel Prize, in 1923 to Banting and McLeod.

1.3.2 Definition

Insulin is a peptide hormone secreted by the β -cells of the pancreatic islets of Langerhans and maintains normal blood glucose levels by facilitating cellular glucose uptake, regulating carbohydrate, lipid and protein metabolism, promoting cell division and growth via mitogenic effects.

1.3.3 Structure of Insulin

In 1928 polypeptide nature of insulin was defined and its amino acid sequence was identified in 1952. It is in fact a dipeptide, containing A and B chains respectively, linked by disulphide bridges, and containing 51 amino acids, with a molecular weight of 5802 (Home, 1997). The A chain comprises 21 amino acids and the B chain 30 amino acids (Dodson and Steiner, 1998).

1.3.4 Synthesis and release of Insulin

Insulin is synthesized in the β -cells of the pancreatic islets of Langerhans as its precursor, proinsulin. Proinsulin is synthesized in the ribosomes of the rough endoplasmic reticulum (RER) from the mRNA as pre-proinsulin. Pre-proinsulin is formed by sequential synthesis of signal peptide, the B chain, the connecting (C-) peptide and then the A chain comprising a single chain of 100 amino acids. Removal of the signal peptide forms proinsulin, which acquires its characteristic three dimensional structure in the endoplasmic reticulum. Secretory vesicles transfer proinsulin from the RER to the Golgi apparatus, whose aqueous zinc-containing and calcium rich environment favours formation of soluble zinc-containing proinsulin hexamers (Dodson and Steiner, 1998). As immature storage vesicles form from Golgi, enzymes acting outside the Golgi convert proinsulin to insulin and C-peptide (Malaisse, 1997). Insulin forms zinc-containing hexamers which are insoluble, precipitating as chemically crystals at pH 5.5. When mature granules secreted into circulation by exocytosis, insulin and an equimolar ratio of C-peptide are released. Proinsulin and zinc typically comprise no more than 6% of the islet cell secretion (Dodson and Steiner, 1998).

1.3.5 Mechanism of action of insulin secretion

Increased levels of glucose induce the “first phase” of the glucose-mediated insulin secretion by release of insulin from secretory granules in the β -cell. Glucose entry into the β -cell is sensed by glucokinase, which phosphorylates glucose to glucose-6-phosphate (G6P), generating ATP (De Lonlay and Saudubray, 2000). Closure of K^{+} -ATP-dependent channels results in membrane depolarization and activation of voltage dependent calcium channels leading to an increase in intracellular calcium concentrations; this triggers pulsatile insulin secretion (Soria *et al.*, 2004). Other mediators of insulin release include activation of phospholipases and protein kinase A, (e.g., by acetylcholine) and by stimulation of adenylyl cyclase activity and activation of β -cell protein kinase A (Bratanova-Tochkova *et al.*, 2002).

1.3.6 Sites of action of insulin

1.3.6.1 Vasculature

In addition to conventional responses on skeletal muscle and adipose tissue, insulin also exerts important biological effects on the vasculature (Baron and Brechtel, 1993; Baron, 1994).

Insulin-stimulated vasodilation. Insulin-stimulated vasodilation has been attributed to several mechanisms, including interaction with the sympathetic nervous system at the vascular level, the activation of ion channels (Hasdai *et al.*, 1998), the release of adenosine (McKay and Hester, 1996) and increase in the generation of NO by the vascular endothelium (Scherrer *et al.*, 1994; Steinberg *et al.*, 1994). Studies in isolated vessels and cultured endothelial cells have improved our understanding of the molecular mechanisms by which insulin stimulates relaxation of isolated vascular strip preparation (Laight *et al.*, 1998), and blunts the response to a number of vasoconstricting reagents (Wambach and Liu, 1992). Therefore variant insulin signalling proteins may contribute to the genetic predisposition to develop endothelial dysfunction and cardiovascular disease. Insulin-stimulated PI 3-kinase activity results in activation of PKB (protein kinase B)/AKt, which has been demonstrated to mediate many metabolic and vascular effects of insulin

(Whiteman *et al.*, 2002). In cultured endothelial cells, stimulation of AKt (PKB) by insulin has been proposed to phosphorylate eNOS at specific residue, Ser¹¹⁷⁷ (human sequence), increasing its activity several folds (Michell *et al.*, 1999; Kim *et al.*, 2001). Majority of studies have demonstrate that insulin stimulate NO production in endothelial cells. In a number of insulin-resistant states, the ability of supraphysiological concentrations of insulin to stimulate blood flow is attenuated (Yki-Jarvinen and Utriainen, 1998), indicating that direct effects of insulin on NO production is of importance in the vascular endothelium (Yki-Jarvinen, 2003).

Vasoconstrictive actions of insulin. It has also been suggested that insulin has vasoconstrictor action, thought to be principally mediated by Endothelin-1 (ET-1) (Webb, 1997). Insulin stimulate ET-1 production in endothelial cells both *in vivo and in vitro*, and this is thought to influence both normal blood flow and changes in blood pressure in patients with hyperinsulinaemia and insulin resistance (Piatti *et al.*, 2000). During eNOS inhibition and hyperglycemia, insulin stimulates vasoconstriction (Schroeder *et al.*, 1999), mediated by ET-1 (Eringa *et al.*, 2002). NO-depending vasodilation is attenuated in insulin-resistant subjects, whereas acute stimulation of ET-1 release is unaffected (Ferri *et al.*, 1995). Insulin also activates MAPK (mitogen-activated protein kinase) which is thought to be a key step in vasoconstriction and ET-1 activation in skeletal muscle arterioles, an effect to antagonize PI 3-kinase-depending vasodilation (Eringa *et al.*, 2004).

Other vascular effects of insulin. Studies in cultured endothelial cells, have shown that, insulin up-regulates eNOS expression (Aljada *et al.*, 2000; Ding *et al.*, 2000) and inhibits the expression of the pro-atherogenic transcription factor NF κ B (nuclear factor κ B).

1.3.6.2 Muscle

Glucose uptake into muscle is essentially insulin dependent via GLUT 4, and muscle accounts for about 60-70% of whole-body insulin mediated uptake (Smith, 2002). In fed state, insulin promotes glycogen synthesis via activation of glycogen synthase, enabling energy release *via* glycolysis (e.g. during

intense muscular activity). Muscle cells do not rely on glucose (or glycogen) for energy during basal state, when insulin levels are low. Insulin suppresses protein catabolism while insulin deficiency promotes release of amino acids for gluconeogenesis (Giorgino *et al.*, 2005). In insulin resistance, muscle glycogen synthesis is impaired; this appears mediated by reduced intracellular glucose translocation (Hunter and Garvey, 1998).

1.3.6.3 Liver

While glucose uptake into the liver is not insulin dependent, it accounts for about 30 % of whole body insulin-mediated glucose disposal (Smith, 2002). In liver glycogen synthesis is stimulated and protein synthesis and lipoprotein metabolism are modulated (Denton and Tavare, 1997). Gluconeogenesis and ketone body production are inhibited (Hunter and Garvey, 1998). Alterations of lipoprotein metabolism represent a major manifestation of insulin resistance. Increased free fatty acid delivery, and reduced VLDL catabolism by insulin resistant adipocytes, results in increased hepatic triglyceride content and VLDL secretion (Krauss, 2004). Moreover, synthesis of CRP in hepatocytes is induced in response to adipocyte-derived pro-inflammatory cytokines such as IL-6 and TNF- α (Devaraj *et al.*, 2004).

1.3.7 Physiological role of Insulin

Insulin is the pivotal hormone regulating cellular energy supply and macronutrient balance, thus directing anabolic processes of the fed state (Burks and White, 2001). Insulin is essential for the intracellular transport of glucose into insulin-dependent organs such as muscle and adipose tissue. Signalling abundance of exogenous energy, adipose tissue breakdown is suppressed and its synthesis promoted. In muscle cells, glucose entry promotes glycogen synthesis and storage, and for carbohydrates, rather than fatty acids, (or amino acids) to be utilized as the immediately available energy source for muscle contraction. Insulin therefore, promotes glycogen and lipid synthesis in muscle cells, while suppressing lipolysis and gluconeogenesis from muscle amino acids. In the presence of an adequate supply of amino acids, insulin is anabolic in muscle (Karam, 1997). In vascular tissue, insulin has several specific physiological roles which includes induction of

vasodilation, *via* insulin signalling pathway "PI 3-kinase" in which activated Akt enzyme phosphorylate and activate eNOS enzyme leading to increased production of NO which act as a vasodilator (Zeng and Quon, 1996; Fleming and Busse, 2003).

1.3.8 Insulin signalling

Insulin mediates its actions through binding to insulin receptor (Fig. 1.1). The insulin receptor was first described in 1971. It consists of a heterotetramer consisting of 2 α and 2 β glycoprotein subunits linked by disulphide bonds and is located on the cell membrane (Kido *et al.*, 2001). Insulin binds to the extracellular α subunit, resulting in conformational change enabling binding of ATP to intracellular component of the β subunit (Wolever, 1990). ATP binding in turn, triggers phosphorylation of β subunit conferring tyrosine kinase activity. This enables tyrosine phosphorylation of intracellular substrate proteins known as insulin receptor substrates (Rosenbaum *et al.*, 1997; Kido *et al.*, 2001). There are four known specifically-named IRS proteins: IRS 1 and 2 have widely overlapping tissue distribution (Withers and White, 2000). Based on results from specific knockout models, the most important proteins in the regulation of carbohydrate metabolism appear to be IRS-1 and -2 (White, 2002). In humans, rare mutations of the IRS-1 protein are associated with insulin resistance (Whitehead *et al.*, 1998) and disruption of the IRS-1 gene in mice results in insulin resistance mainly of muscle and fat (Yamauchi *et al.*, 1996). Interestingly, knockout mice not only show insulin resistance of muscle, fat and liver, but also develop and manifest diabetes as a result of β -cell failure (Previs *et al.*, 2000). Therefore, it is tempting to speculate that dysfunction of IRS-2 and its downstream targets might represent a common feature of both peripheral insulin resistance and β -cell failure. Furthermore, circulating free fatty acids and the adipokine TNF- α increase serine phosphorylation of IRS proteins, thereby causing impaired insulin signal transduction (White, 2002). In addition, prolonged stimulation with insulin as commonly found in diabetic patients with hyperinsulinaemia may result in regulated degradation of IRS proteins (Rui *et al.*, 2001).

Down-stream of IRS proteins, the phosphatidylinositol-3-kinase (PI 3-kinase) is a central mediator of the effect of insulin. PI 3-kinase isoforms have been subdivided into three classes Ia, Ib, and II. Class Ia PI 3-kinases are thought to be the major effector of insulin signaling and activation of protein kinase B (PKB) also called (Akt) (Alessi and Cohen, 1998). Tyrosine phosphorylated IRS-1 or IRS-2 then binds to src-homology 2 (SH2) domains of intracellular proteins, including the p85 regulatory subunit of phosphatidylinositol (PI) 3-kinase. The interactions of IRS and p85 subunit of PI 3-kinase results in the activation of p110 catalytic subunit of PI 3-kinase. Activation of PI 3-kinase increases serine phosphorylation of Akt which in turn, will mediate the effects of insulin on glucose transport, glycogen synthesis, protein synthesis, lipogenesis, and suppression of hepatic gluconeogenesis. Akt regulates both, glucose uptake via facilitated glucose transporters (GLUTs) and intracellular glucose metabolism in insulin sensitive tissue such as skeletal muscle (Fig. 1.1) (Alessi and Cohen, 1998).

As is the case for other growth factors, insulin stimulates the mitogen-activated protein (MAP) kinase extracellular signal regulated kinase (ERK) (Fig. 1.1). This pathway involves the tyrosine phosphorylation of IRS proteins and/or Shc, which in turn interact with the adaptor protein Grb2, recruiting the Son-of-sevenless (SOS) exchange protein to the plasma membrane for activation of Ras. The activation of Ras also requires stimulation of the tyrosine phosphatase SHP2, through its interaction with receptor substrates such as Gab-1 or IRS1/2 (Fig. 1.1). Once activated, Ras operates as a molecular switch, stimulating a serine kinase cascade through the stepwise activation of Ras, MEK, and ERK. Activated ERK can translocate into the nucleus, where it catalyses the phosphorylation of transcription factors such as P62, initiating a transcriptional programme that leads to cellular proliferation or differentiation (Boulton *et al.*, 1991). Blockade of the pathway with dominant negative mutants or pharmacological inhibitors prevents the stimulation of cell growth by insulin, but no effect on the metabolic actions of the hormone (Lazar *et al.*, 1995).

In vascular tissue, in addition to the above insulin signalling pathways and metabolic effects, the activated AKT " by phosphorylated PI 3-kinase" will directly phosphorylates eNOS on serine 1177 and activates the enzyme, leading to increased NO production and thus providing vascular protection (normal vasodilation) (Fig. 1.1) (Zeng and Quon, 1996; Fleming and Busse, 2003).

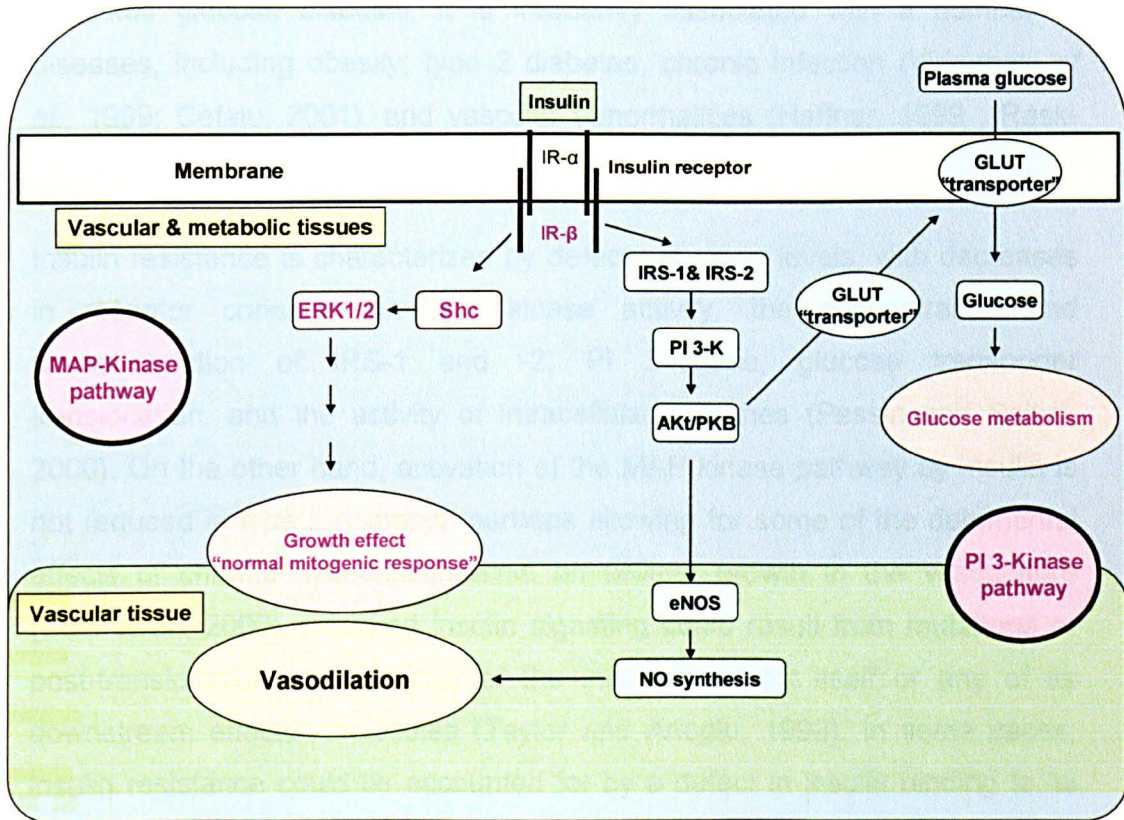


Figure 1.1 Insulin signalling pathways in vascular and metabolic tissues

1.3.9 Insulin resistance

Insulin resistance is a common pathologic state in which target cells fail to respond to ordinary levels of circulating insulin, thus producing an attenuated biological response. Classically this refers to impaired sensitivity to insulin mediated glucose disposal. It is frequently associated with a number of diseases, including obesity; type 2 diabetes, chronic infection (Virkamaki *et al.*, 1999; Cefalu, 2001), and vascular abnormalities (Haffner, 1999 ; Rask-Madsen and King, 2005).

Insulin resistance is characterized by defects at many levels, with decreases in receptor concentration and kinase activity, the concentration and phosphorylation of IRS-1 and -2, PI 3-kinase, glucose transporter translocation, and the activity of intracellular enzymes (Pessin and Saltiel, 2000). On the other hand, activation of the MAP kinase pathway by insulin is not reduced in type 2 diabetes, perhaps allowing for some of the detrimental effects of chronic hyperinsulinaemia on cellular growth in the vasculature (Cusi *et al.*, 2000). Impaired insulin signaling could result from mutations or post-translocation modifications of the insulin receptor itself or any of its downstream effector molecules (Taylor and Arioglu, 1998). In some cases, insulin resistance could be accounted for by a defect in insulin binding to its receptor (Roach *et al.*, 1994); however, insulin resistance is most often attributed to post binding defect in insulin action. A marked reduction in the receptor kinase activity was observed in several patients (type A) with extreme resistance to insulin, but with normal binding (Grunberger *et al.*, 1984; Grigorescu *et al.*, 1986). Similarly, severe defects in receptor kinase activity are associated with naturally occurring mutations of the insulin receptor gene (Taylor *et al.*, 1992), indicating that genetic and acquired factors can profoundly influence insulin sensitivity. Nonetheless, genetic defects in the insulin receptor are relatively rare, but represent the most severe forms of insulin resistance and are exemplified by leprechaunism, the Rabson Mendenhall, and the type A syndrome of insulin resistance (Taylor and Arioglu, 1998).

1.3.9.1 Insulin resistance and metabolic function:

Insulin resistance affects many tissue functions and a variety of metabolic processes in the body. Hyperinsulinaemia causes exaggerated responses in tissues that remain sensitive to insulin. For example, hyperinsulinaemia stimulates the sympathetic nervous system, which may contribute to hypertension (Reaven *et al.*, 1996; Ginsberg, 2000), or increasing ovarian androgen production, which is integral to hormonal imbalances observed in the polycystic ovary syndrome (Franks, 1995; Dunaif and Thomas, 2001). Similar to the muscle and liver cells, there are various cellular functions that also exhibit resistance to insulin. In particular, insulin resistance in fat cells leads to increased lipolysis with release of fatty acids and a variety of sequelae, including dyslipidaemia and vascular abnormalities caused by excessive amounts of circulating free fatty acids (Ginsberg, 2000). High free fatty acid concentrations also contribute to resistance to insulin's action by enhancing glucose output from the liver and reducing glucose disposal in skeletal muscle (Boden, 1999).

1.4 INSULIN RESISTANCE AND OBESITY

Increased adipose tissue, especially that in an upper body or "android" deposition, was first associated with diabetes and vascular disease by French endocrinologist Jean Vague in 1956 (Vague, 1996). Insulin resistance increases with increasing body mass index, waist circumference, and in particular waist-hip ratio (Aronne and Segal, 2002), reflecting increased adiposity especially increased levels of visceral adipose tissue. Visceral adipose tissue refers to intra-abdominal fat around the intestines and correlates closely with liver fat accumulation.

1.4.1 Fatty acid-induced insulin resistance

Metabolic characteristics of visceral adipose tissue differs from that of the subcutaneous fat. Visceral adipose tissue is metabolically more active; increased free fatty acid turnover that promotes insulin resistance at cellular level (Giorgino *et al.*, 2005). The insulin resistance seen in obesity is believed to involve primarily muscle and liver, with increased adipocyte-derived free

fatty acids promoting triglyceride accumulation in these tissues. (Perseghin *et al.*, 2003). Circulating FFAs derived from adipocytes are elevated in many insulin-resistant states and have been suggested to contribute to the insulin resistance of diabetes and obesity by inhibiting glucose uptake, glycogen synthesis, glucose oxidation, and by increasing hepatic glucose output (Bergman and Ader, 2000). Elevated FFAs are also associated with a reduction in insulin-stimulated IRS-1 phosphorylation and IRS-1 associated PI 3-kinase activity (Shulman, 2000). There are two hypotheses that links increased circulating FFAs with insulin resistance:

1) Randle *et al.* demonstrated that fatty acids compete with glucose for substrate oxidation in isolated rat heart muscle and rat diaphragm muscle (Fig. 1.2). They speculated that increased fat oxidation causes the insulin resistance associated with obesity (Randle *et al.*, 1963; Randle *et al.*, 1965). The mechanism they proposed to explain the insulin resistance was that an increase in fatty acids caused an increase in the intramitochondrial acetyl CoA/CoA and NADH⁺/NAD⁺ ratios, with subsequent inactivation of pyruvate dehydrogenase (Fig. 1.2). This in turn would cause intracellular citrate concentrations to increase, leading to inhibition of phosphofructokinase, a key rate-controlling enzyme in glycolysis. Subsequent accumulation of glucose-6-phosphate would inhibit hexokinase II activity, resulting in an increase in intracellular glucose concentrations and decreased glucose uptake.

2) Alternative mechanism for fatty acid-induced insulin resistance has been proposed based on studies on human skeletal muscle (Fig. 1.3). An increase in delivery of fatty acids to muscle or a decrease in intracellular metabolism of fatty acids, leads to an increase in intracellular fatty acid metabolites such as diacylglycerol, fatty acyl CoA, and ceramides. These metabolites activate a serine/threonine kinase cascade (possibly initiated by protein kinase C) leading to phosphorylation of serine/threonine sites on insulin receptor substrates (IRS-1 and IRS-2), which in turn reduces the ability of the insulin receptor substrates to activate PI 3-kinase. As a consequence, glucose transporter activity and other events downstream of insulin receptor signaling are diminished (Gavrilova, 2000; Kim *et al.*, 2000).

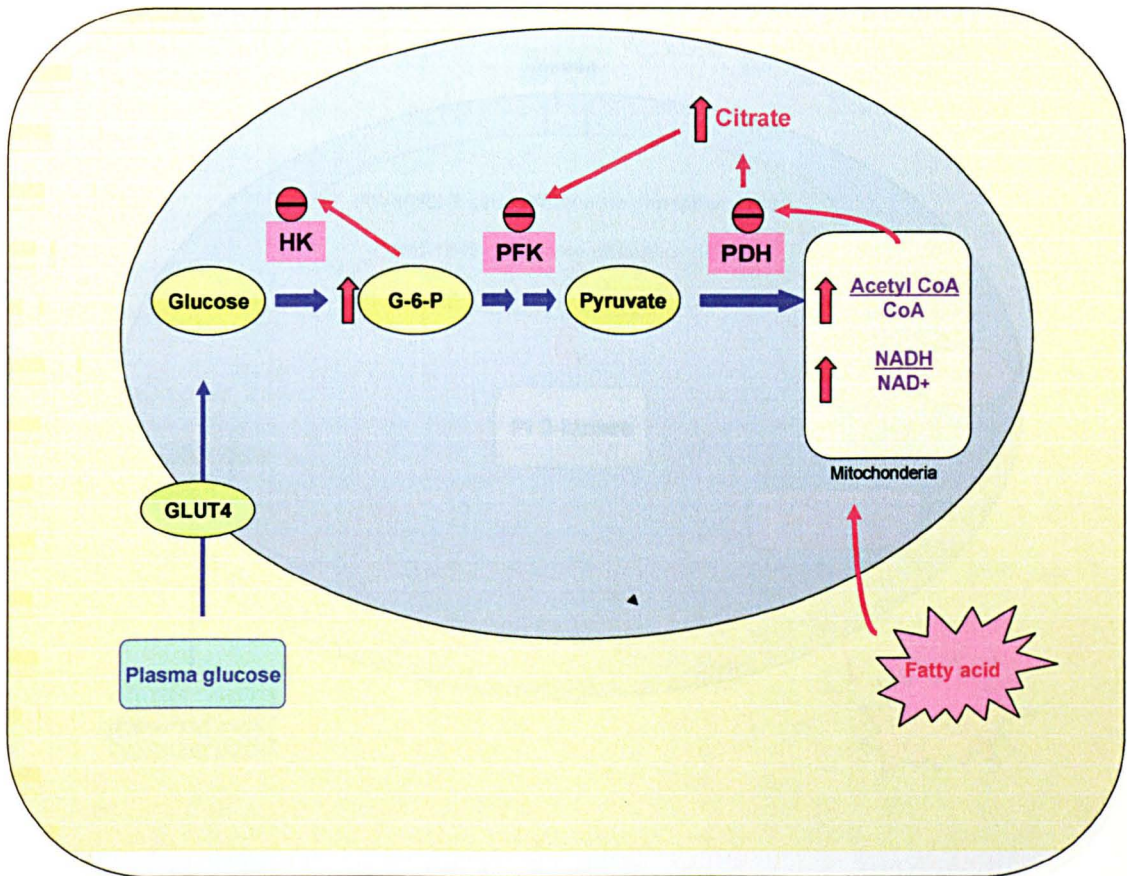


Figure 1.2 Mechanism of fatty acid-induced insulin resistance in skeletal muscle as proposed by Randle et al.

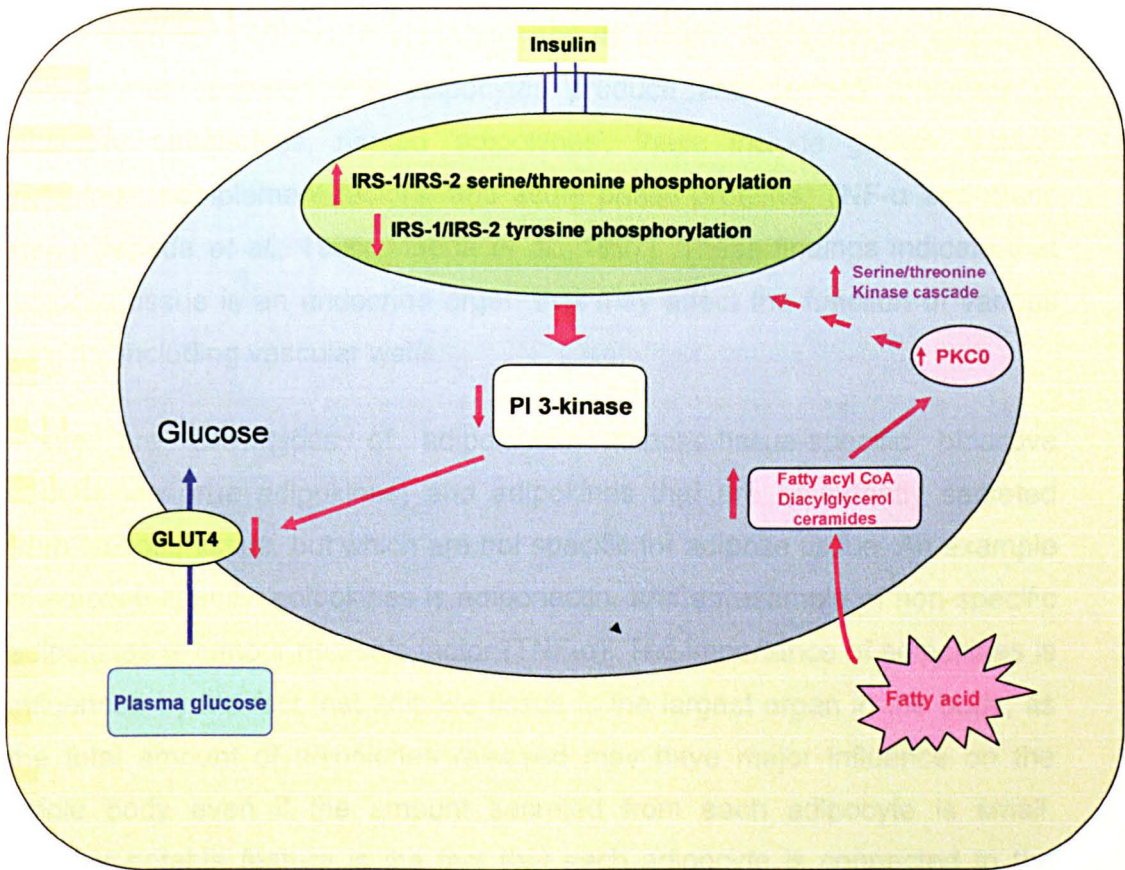


Figure 1.3 Proposed alternative Mechanism for fatty acid-induced insulin resistance in skeletal muscle.

1.4.2 Adipokines and insulin resistance

There is a close relationship between insulin resistance and visceral adiposity. It has become apparent that insulin sensitivity is modulated by adiposity, and this has led to significant interests in adipokines, which may play a pivotal role in the regulation of insulin signalling and its action. Research on adipocyte biology has revealed that adipocytes produce and secrete a variety of bioactive substances, named 'adipokines'; these include growth factors, cytokines, complement factors, and acute phase proteins, TNF- α and many more (Maeda *et al.*, 1996; Maeda *et al.*, 1997). These findings indicate that adipose tissue is an endocrine organ that may affect the function of various organs, including vascular walls.

There are two types of adipokines: adipose-tissue-specific bioactive substances (true adipokines) and adipokines that are abundantly secreted from adipose tissue, but which are not specific for adipose tissue. An example of adipose-specific adipokines is adiponectin, and an example of non-specific adipokines is tumour necrosis factor (TNF- α). The importance of adipokines is influenced by the fact that adipose tissue is the largest organ in the body, as the total amount of adipokines released may have major influence on the whole body even if the amount secreted from each adipocyte is small. Another notable feature is the fact that each adipocyte is connected to the vascular network, therefore, released adipokines can easily reach the systemic circulation.

Dysregulation of adipokine production and secretion is involved in the development of metabolic (including insulin resistance) and cardiovascular diseases. Metabolic syndrome has been receiving an increased interest recently, as a highly atherogenic state that is characterized by the clustering of multiple risk factors such as impaired glucose metabolism, hyperlipidaemia and hypertension in a given individual. In metabolic syndrome, intra-abdominal visceral fat accumulation is shown to play a key role in the

development of a variety of metabolic and circulatory disorders through the dysregulation of adipokines.

1.4.2.1 TNF- α :

Is expressed as a 26 kDa cell surface transmembrane protein, subsequently cleaved to a 17 kDa active biological form, produced by a broad range of tissues, but mainly by adipocytes (Kern *et al.*, 1995; Wajant *et al.*, 2003). It has been shown that adipose TNF- α mRNA and plasma TNF- α protein are increased in most animal models and human subjects with obesity and insulin resistance. Neutralising the blood TNF- α in obese rats with a soluble TNF- α receptor, immunoglobulin G fusion protein, markedly improves insulin resistance. This finding indicates that higher production of TNF- α accumulated in adipose tissue may be causative for obesity-associated insulin resistance. Furthermore, vascular endothelium is a principal target of actions of TNF- α (Locksley *et al.*, 2001; Wajant *et al.*, 2003). Bioavailability of NO is reduced in cultured endothelial cells incubated with TNF- α (Myers *et al.*, 1994). In addition, endothelium-dependent vasodilatation is impaired in animals and human subjects after infusion with TNF- α (Wang *et al.*, 1994; Bhagat and Valance, 1997). Possibly by reducing eNOS mRNA stability or increasing super oxide production (Yoshizumi *et al.*, 1993; De Keulenaer *et al.*, 1998).

1.4.2.2 IL-6:

Is a 26 kDa protein produced by several tissues, including adipose and endothelium (Mohamed-Ali *et al.*, 1997). Adipose tissue has been postulated to contribute a significant production (approx. 30 %) of the total circulating IL-6 (Mohamed-Ali *et al.*, 1997; Yudkin *et al.*, 1999). Elevated IL-6 concentrations is associated with insulin resistance and obesity (Yudkin *et al.*, 1999). Furthermore, IL-6 results in impaired NO-mediated relaxation of systemic vessels in pregnant rats (Orshal and Khalil, 2004), and stimulates endothelial cell proliferation and angiogenesis (Holzinger *et al.*, 1993; Hernandez-Rodriguez *et al.*, 2003). Moreover, IL-6 has been reported to stimulate the synthesis and secretion of CRP, that contributes to atherosclerosis (Bhakdi *et al.*, 1999).

1.4.2.3 Leptin:

Leptin, the product of the *ob* gene, is a plasma protein secreted by adipocytes and is involved in the control of body weight, mainly through its hypothalamic effects. Obese subjects have been shown to have up to a 10-fold greater plasma leptin concentration than lean subjects (Considine *et al.*, 1996), but the anorexigenic effects of leptin become less pronounced indicating development of leptin resistance. However, the exact mechanism of this proposed 'leptin resistance' is not fully clear yet (Bjorbaek *et al.*, 1999; Banks and Farrell, 2003).

The presence of leptin receptors on endothelial cells, (Bouloumie *et al.*, 1998) has led to the hypothesis that leptin interacts with the vascular wall and may contribute to atherosclerotic change and angiogenesis. Atherosclerosis is initiated by endothelial cell damage. As the atherosclerotic plaque progresses it develops its own microvascular network (neovascularisation), the disruption of which is thought to play a role in plaque instability. Leptin enhances formation of capillary-like tubes and neovascularization *in vitro* and *in vivo* (Bouloumie *et al.*, 1998; Sierra-Honigmann *et al.*, 1998). Moreover, studies in normal adolescents with variable BMIs found increased leptin concentration to be associated with impaired vascular function independent of the metabolic and inflammatory disturbances associated with obesity (Singhal *et al.*, 2002). Furthermore, *in vitro* stimulation of cultured endothelial cells with leptin leads to an increase in the production of (Reactive oxygen Species) ROS, *via* hydrogen peroxide (Bouloumie *et al.*, 1999) and superoxide production from mitochondria (Bouloumie *et al.*, 1999). Migration of endothelial cells is a key event in angiogenesis contributing to pathological states such as diabetic vasculopathy. Leptin has been shown to induce cell migration in human umbilical vein endothelial cells (HUVECs), mediated by the PI 3-K/Akt (pkB)/eNOS and MAPK pathways (Goetze *et al.*, 2002). In contrast, number of studies have indicated that leptin stimulates vasodilation (Lembo *et al.*, 2000; Matsuda *et al.*, 2003). Despite this insulin-mimetic action of leptin that may mediate beneficial cardiovascular effects, the majority of studies

performed to date indicate that hyperleptinaemia is atherogenic. Therefore, to further elucidate the exact role of leptin in obesity-related endothelial dysfunction and its mechanisms, more *in vivo* and *ex vivo* studies are needed.

1.4.2.4 C-reactive protein (CRP):

An acute phase reactant marker produced by the liver and adipose tissue in response to systemic inflammation. The circulating level of CRP rises with BMI (Visser *et al.*, 1999; Pannacciulli *et al.*, 2001; Bullo *et al.*, 2003; Anty *et al.*, 2006), and elevated levels of this inflammatory marker have been associated with both obesity and diabetes, while its levels fall with a weight loss (Tchernof *et al.*, 2002). A very recent study, (unpublished results, IS Wood and P Trayhurn) using conventional RT-PCR have found that there are very small amount of CRP levels in human white adipose tissue (WAT), or adipocytes. The very low levels of CRP expression would suggest that adipocytes are unlikely to be a significant contributor of circulating CRP levels. Secretion of IL-6 from adipose tissue is increased in obesity. IL-6 is the major cytokine regulating the hepatic production of CRP (Heinrich *et al.*, 1990; Yudkin *et al.*, 1999). Thus, WAT may be a major player in the raised circulating levels of CRP in obesity, but through the indirect route of adipocyte-derived IL-6. CRP levels are frequently elevated in patients with acute ischaemia and myocardial infarction (Biasucci *et al.*, 1999; Buffon *et al.*, 1999) indicating its active role in the atherosclerosis process (Yeh *et al.*, 2001; Verma *et al.*, 2002). Thus, CRP should be regarded as an indirect, albeit important measure of endothelial function.

1.5 INSULIN RESISTANCE AND VASCULAR SYSTEM

1.5.1 Pathways of insulin action in vasculature

When insulin activates its receptor, 2 major pathways are switched on (Fig. 1.1). The first pathway involves activation of phosphatidylinositol 3-kinase, which is an important mechanism for glucose transport in skeletal muscle. This pathway has also recently been shown to be an important factor for endothelial nitric oxide production that leads to insulin-induced vasodilation. The second insulin-activated pathway is the MAP-K pathway, which promotes insulin-mediated growth of the vascular smooth muscle cells (VSMC).

Therefore, activated ERK can translocate into the nucleus, thus initiating a transcriptional programme that lead to cellular proliferation or differentiation (Boulton *et al.*, 1991; Lazar *et al.*, 1995). Moreover, MAP-K also can stimulates migration induced by insulin (Wang *et al.*, 2003).

1.5.2 Endothelium and endothelial function.

The endothelium is the layer of thin, flat cells that lines interior surface of blood vessels, forming an interface between circulating blood in the lumen and the rest of the vessel wall. Endothelial cells line the entire circulatory system, from the heart to the smallest capillary. The importance of endothelium in vascular function has been known for decades. Endothelial cells not only provide the physical lining of the blood vessels but also play a vital role in vascular homeostasis, vascular tone regulation, vascular smooth muscle cells proliferation, trans-endothelial leukocyte migration, thrombosis, and thrombolysis. Endothelial cells are involved in many aspects of vascular biology including:

- vasoconstriction & vasodilation, and hence the control of blood pressure
- blood clotting
- atherosclerosis
- formation of new blood vessels (angiogenesis)
- inflammation and swelling (oedema)

Endothelial cells also control the passage of materials and the transit of white blood cells into and out of the bloodstream. In some organs, there are highly differentiated endothelial cells to perform specialised 'filtering' functions. Examples of such unique endothelial structures include renal glomerulus and blood-brain barrier. Endothelial tissue is a specialised type of epithelium tissue (one of the four types of biological tissue in animals (Alberts *et al.*, 2002). In response to various mechanical and chemical stimuli, endothelium synthesise and release a large number of vasoactive substances; that include NO and prostaglandins (Quyyumi, 1998). Clinical problems may develop when these functions or processes are in imbalance.

1.5.3 Endothelial dysfunction

It can be defined as the partial or complete loss of balance between vasoconstrictor and vasodilators, growth promoting and inhibiting factors, pro-atherogenic and anti-atherogenic factors, and pro-coagulant and anti-coagulant factors (Quyyumi, 1998). In other words, endothelial dysfunction is a physiological dysfunction of normal biochemical processes carried out by the endothelium, the cells that line the inner surface of all blood vessels including arteries and veins. Compromise of normal function of endothelial cells is characteristic of endothelial dysfunction. Endothelial dysfunction can result from disease processes, as occurs in septic shock, hypertension, hypercholesterolaemia, diabetes as well as from environmental factors, such as smoking tobacco products. Endothelial dysfunction is now regarded as an early pivotal event in atherogenesis (Ross, 1999) and has been shown to precede the development of clinically detectable atherosclerotic plaques in the coronary arteries (Mano *et al.*, 1996). It has also been considered an important event in the development of microvascular complications in diabetes (Tooke, 1995).

A key feature of endothelial dysfunction is the inability of arteries and arterioles to dilate fully in response to appropriate stimuli. In animal models, this can be tested using myograph study. This technique is thought to stimulate the endothelium to release nitric oxide (NO), which diffuse into the surrounding vascular smooth muscle causing vasodilatation.

Dysfunctional endothelial cells are unable to produce NO to the same extent (or there maybe an increased and rapid destruction of NO) as healthy endothelial cells and therefore vasodilatation is reduced. Endothelial function can be improved significantly by exercise and improved diet. Other factors have been identified as improving endothelial function includes cessation of smoking, loss of weight and treatment of hypertension and hypercholesterolaemia amongst various other measures.

1.5.4 Nitric Oxide synthesis

Nitric oxide is the major factor in arteries mediating endothelial dependent relaxation. NO is produced in endothelial cells through the conversion of L-arginine to L-citrulline (Fig. 1.4), in the presence of molecular oxygen and NADPH, via the activity of endothelial enzyme Nitric Oxide Synthase (eNOS), and its cofactors tetrahydrobiopterin, flavin adenine dinucleotide and flavin mononucleotide (Fig. 1.4). Insulin enhances tetrahydrobiopterin production by stimulating its biosynthetic enzyme GTP cyclohydrolase, and thus stimulates eNOS activity by calcium-independent phosphorylation of eNOS at serine and threonine residues via PI 3-kinase/Akt pathway, resulting in enhanced nitric oxide production. Nitric oxide then diffuses to the underlying vascular smooth muscle cells to stimulate the production of cyclic guanosine monophosphate (cGMP) through the activation of enzyme guanylate cyclase. The increase in cGMP will lead to a reduction in intracellular calcium ions concentration, inducing vasodilation (Fig. 1.4).

There are various pharmacological and non-pharmacological stimuli that can be used to assess the production of nitric oxide in endothelial cells *in vivo* as well as its effect on the underlying vascular smooth muscle cells (endothelium-dependent and independent vasodilation, respectively). The intra-arterial infusion of acetylcholine or methacholine has widely been used to stimulate endothelial cell production of nitric oxide, whereas sodium nitroprusside has been used to directly stimulate vascular smooth muscle cells. Another approach to the assessment of endothelial function is to measure plasma levels of markers of low-grade inflammation such as C-reactive protein (CRP), IL-6, and TNF- α . Apart from vasodilatory effects, NO reduces vascular smooth muscle cell migration and growth (Sarkar *et al.*, 1996), platelet aggregation and thrombosis (Antl *et al.*, 2006), monocyte and macrophage adhesion, and inflammation (Peng *et al.*, 1998).

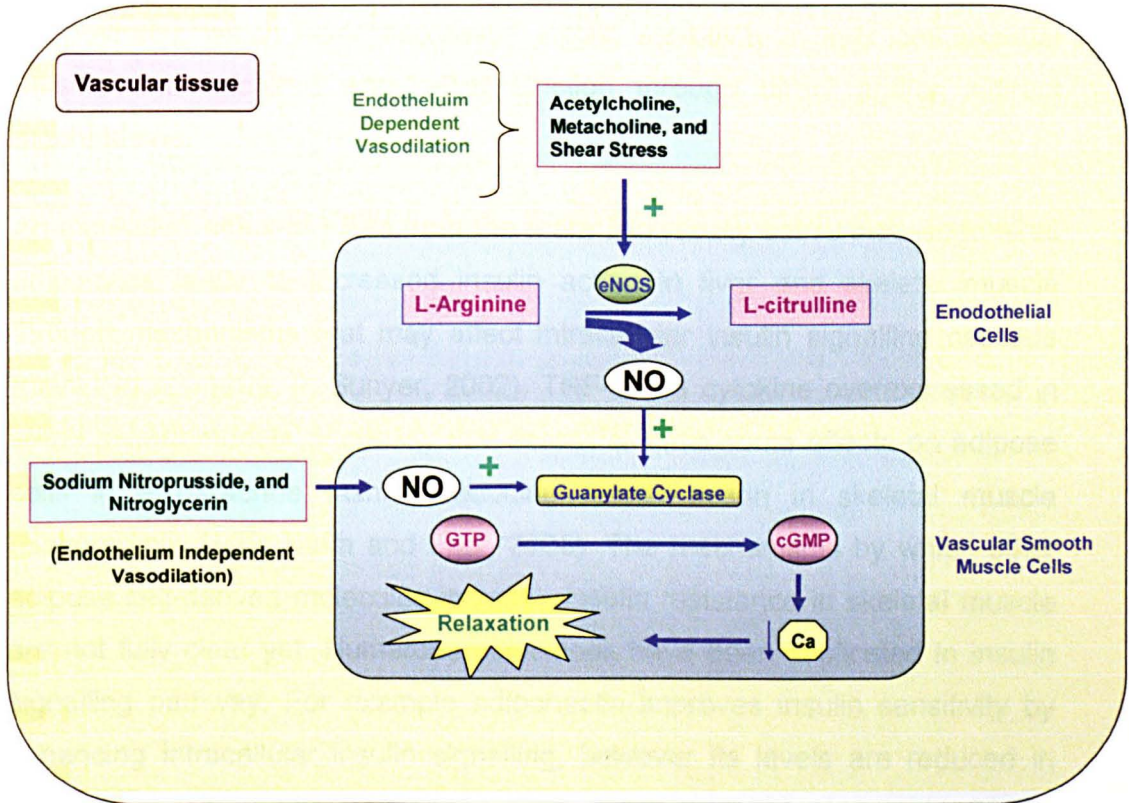


Figure 1.4 Synthesis and function of Nitric oxide as vasodilator in vascular system

1.6 OBESITY, INSULIN RESISTANCE, ENDOTHELIAL DYSFUNCTION, AND ATHEROSCLEROSIS: THE LINK

Obesity, particularly "central or abdominal" adiposity, leads to an imbalanced production of several metabolic products, hormones, and cytokines (adipokines), which favor decreased insulin sensitivity in liver and skeletal muscle and impaired endothelial function through direct and/or indirect mechanisms.

An increased efflux of FFAs from the more lipolytically active intra-abdominal adipocytes leads to increased insulin action in liver and skeletal muscle through mechanisms that may affect intracellular insulin signalling cascade (Griffin *et al.*, 1999; Pi-Sunyer, 2002). TNF- α is a cytokine overexpressed in some animal models of obesity that seems to have some effects on adipose cells in a paracrine fashion reducing insulin action in skeletal muscle (Hotamisligil, 1999; Kahn and Flier, 2000). The mechanisms by which other adipose cell-derived molecules increase insulin resistance in skeletal muscle are not fully clear yet. Numerous adipokines have been implicated in insulin signalling pathway. For example adiponectin improves insulin sensitivity by enhancing intracellular insulin signalling, however its levels are reduced in states of obesity/insulin resistance (Yamauchi *et al.*, 2001; Yamauchi, 2003). Resistin, another fat-derived hormone, has also been found to be linked to obesity and insulin resistance in animal models; yet, its role in human obesity seems to be insignificant (Nagaev and Smith, 2001; Stepan *et al.*, 2001).

In addition to inducing insulin resistance, certain fat-derived products might have a direct effect on vascular function. FFAs (Steinberg and Baron, 2002), TG (Naderali *et al.*, 2001) and adipokines IL-1, IL-6, and TNF- α been closely linked to endothelial dysfunction and subclinical inflammation (Winkler *et al.*, 1999; Yudkin *et al.*, 2002). IL-6 is a potent stimulus for the production of CRP in the liver which, in turn, may have some direct deleterious effects in the vascular wall (Yudkin *et al.*, 1999). CRP is considered an excellent marker of low-grade inflammation in the vascular wall, a well recognized mechanism in the development of atherosclerosis (Libby *et al.*, 2002).

The mechanisms by which insulin resistance leads to endothelial dysfunction are certainly multiple and complex. All major abnormalities that are part of the insulin resistance syndrome, such as hyperglycaemia, hypertension, dyslipidaemia, and altered coagulation/fibrinolysis, are directly and independently linked to endothelial dysfunction. Despite that, it is still not clear whether insulin resistance *per se* causes endothelial dysfunction and ultimately atherosclerosis. Nonetheless, hyperinsulinaemia, a surrogate marker for the presence of insulin resistance, was found to be an independent risk factor for cardiovascular disease in nondiabetic individuals (Despres *et al.*, 1996).

New insights into a more direct link between insulin resistance and the vasculature come from the observation that insulin has direct vascular effects. Insulin is known to have a direct vasodilatory effect mediated through stimulation of nitric oxide production in endothelial cells (Kuboki *et al.*, 2000). In insulin-resistant state, the ability of insulin to stimulate nitric oxide production in the endothelium is diminished. Interestingly, the stimulation of nitric oxide production in endothelial cells and the stimulation of glucose uptake in muscle and fat tissue by insulin occur through the phosphatidylinositol 3-kinase (PI3-K) and Akt pathway (Cusi *et al.*, 2000; Montagnani and Quon, 2000). In contrast, other effects of insulin action on the vasculature, including the stimulation of migration and growth of smooth muscle cells and the production of plasminogen activator inhibitor (PAI-1), are mediated through the mitogen-activated protein kinase (MAP-K) pathway (Hsueh and Law, 1999).

Many other abnormalities may certainly contribute to the link between insulin resistance and endothelial dysfunction, such as the production of reactive oxygen species, i.e., superoxide (Cai and Harrison, 2000), inflammation (Yudkin *et al.*, 1999), production of cytokines such as TNF- α (Winkler *et al.*, 1999), activation of the renin-angiotensin system (Folli *et al.*, 1999), and elevation of ET-1 (Piatti *et al.*, 2000). Therefore, a large number of mediators with complex mechanisms closely link obesity, insulin resistance, and endothelial dysfunction in a multidirectional and dynamic framework.

Jiang et al. found that Zucker rats, a well-established model of insulin resistance, decreased insulin-mediated glucose uptake in muscle and decreased insulin-mediated nitric oxide production in endothelial cells (Jiang *et al.*, 1999) are mediated through the PI3-K pathway, while insulin-mediated stimulation of VSMC migration and growth and PAI-1 production, *via* MAP-K pathway, is preserved. Extrapolating these findings to a theoretical framework in humans, could suggest that individuals who exhibit the insulin resistance syndrome may have an abnormality in nitric oxide production by endothelial cells (Arcaro *et al.*, 2002). However, it is not known if the mechanism of abnormal selective signalling pathways described in animal model of genetic obesity would also operate in dietary obese animals. Furthermore, whether or not there is an increased secretion of adipocytokines and inflammatory markers in dietary-obese animals merits further investigation.

Support for a closer association of obesity, insulin resistance, and endothelial dysfunction comes from studies that have assessed the effects of weight loss on an improvement in insulin sensitivity and endothelial function. As described above increased adipose tissue, particularly in the abdominal region, is associated with the production of several hormones, mediators, and cytokines that impact insulin sensitivity in liver and skeletal muscle and vascular function. Thus, weight loss and/or an improvement in insulin sensitivity whether physiologically or pharmacologically may lead to a more favorable profile of these fat-derived substances, which would translate into an improvement of metabolic and endothelial function and a reduction of the low-grade inflammatory state. In fact, several studies now support this concept (Dandona *et al.*, 1998; Ziccardi *et al.*, 2002).

1.7 THE PEROXISOME PROLIFERATORS ACTIVATED RECEPTOR- α AGONIST (FENOFIBRATE)

Peroxisome proliferators are a group of chemicals that have several applications such as: therapeutically used drugs (e.g., hypolipidaemic agents), plasticizers, and organic solvents used in the chemical industry, herbicides, and naturally occurring hormones. Peroxisome proliferators, as the name implies cause an increase in the number and size of peroxisomes in the liver

(associated with increased liver size), kidney and heart tissue of susceptible species, such as rats and mice. Long-term administration of peroxisome proliferators can cause liver cancer in these animals (Reddy and Rao, 1992). Whether frequent exposure to peroxisome proliferators represents a hazard to human subjects is unknown; however, increased cancer risk has been shown to be associated with long-term therapeutic administration of the hypolipidemic drugs gemfibrozil, fenofibrate, and clofibrate. The peroxisome proliferator activated receptor- α (PPAR- α), a member of the nuclear receptor superfamily, has been found to be activated by different ligands including naturally occurring eicosanoids and certain long-chain fatty acids, as well as synthetic compounds such as lipid-lowering fibric acids (i.e. fenofibrate, gemfibrozil and the research compound WY14643). Moreover, recent studies have revealed that humans have considerably lower levels of PPAR- α in liver than rodents, and this difference may, in part, explain the species differences in the carcinogenic response to peroxisome proliferators (Reddy and Rao, 1992; Gonzalez *et al.*, 1998).

1.7.1 PPAR- α and peroxisome proliferation in the liver

In hepatocytes, peroxisomes are membrane-bound organelles that contain enzymes responsible for β -oxidation of fatty acids, the biosynthesis of cholesterol, and other biochemical pathways (Mannaerts and Van Veldhoven, 1993). Peroxisome proliferation coincides with increase in many peroxisomal and mitochondrial enzymes involved in fatty acid β -oxidation (Reddy and Chu, 1996). PPAR- α mediates the activation of genes encoding these enzymes through dimerization with the retinoid X receptor α and binding to cis-acting regulatory elements upstream of the promoter regions in target genes (Green and Wahli, 1994; Schoonjans *et al.*, 1997). In rats and mice, PPAR- α is expressed at high levels in the liver and kidney, the primary sites of peroxisome proliferation. (Lee *et al.*, 1995).

In addition to PPAR- α , two other nuclear receptors share significant sequence similarity to PPAR- α . PPAR- γ is highly expressed in white and brown adipose tissues and macrophages; it is not found at significant levels in the liver (Lemberger *et al.*, 1996). PPAR- γ is required for the differentiation of

adipocytes and the tissue-specific regulation of genes in white and brown fat cells (Spiegelman and Flier, 1996). It is the target for the thiozolidinedione drugs used to control adult-onset diabetes (Lehmann *et al.*, 1995). In contrast to PPAR- α and PPAR- γ , PPAR- β is more ubiquitously expressed and is most abundant in the central nervous system (Lemberger *et al.*, 1996). The target genes and physiological functions of PPAR- β are currently unknown.

1.7.2 Physiological role of PPAR- α

On the basis of target gene expression patterns, PPAR- α appears to have critical roles in regulation of fatty acid metabolism, including fatty acid β -oxidation, apolipoproteins, and fatty acid transport proteins (Lee *et al.*, 1995; Auwerx *et al.*, 1996; Aoyama *et al.*, 1998). Moreover, PPAR- α regulates serum cholesterol, in particular, high-density lipoprotein cholesterol, and triglyceride levels, as indicated by the results obtained in mice lacking PPAR- α . Constitutive levels of serum cholesterol are markedly higher in PPAR- α -null mice than in wild-type controls, and fibrate-induced reductions of serum triglycerides do not occur in these mice as well. PPAR- α affects constitutive expression of mitochondrial enzymes involved in fatty acid catabolism as well as induction of mitochondrial and peroxisomal fatty acid β -oxidation pathways by peroxisome proliferators and perhaps other fatty acid metabolites (Aoyama *et al.*, 1998). Thus, genes encoding peroxisomal and mitochondrial fatty acid-metabolising enzymes, apolipoproteins, and fatty acid transport proteins are all regulated by PPAR- α , indicating that this receptor has important physiological roles in control of fatty acid metabolism.

1.7.3 Functions of peroxisome proliferator-activated receptor- α in the endothelium

PPAR- α is expressed in monocytes/macrophages, vascular smooth muscle cells (VSMCs), and endothelial cells, with evidence for regulation by PPARs of relevant vascular target genes (Marx *et al.*, 1999; Marx *et al.*, 2000). There are several studies argue for anti-inflammatory and anti-atherosclerotic effects through PPAR- α . It has been demonstrated that fenofibrate (PPAR- α activator) caused activation for eNOS expression in endothelial cells leading to increased NO production and finally vasodilation (Goya *et al.*, 2004;

Murakami *et al.*, 2006). Moreover, in VSMCs, PPAR- α activators (WY14643, fenofibrate) inhibites interleukin1-induced production of interleukin-6 and prostaglandin as well as expression of cyclo-oxygenase-2 (Staels *et al.*, 1998). Induction of adhesion molecule expression in the endothelium is considered one of the earliest steps in atherogenesis. The endothelium has also been suggested to defend against oxidative stress and to maintain appropriate redox status (Kunsch and Medford, 1999; Inoue *et al.*, 2001). In this sense, data implicating PPARs in redox responses are also relevant. Inoue *et al.* (Inoue *et al.*, 2001) found that both PPAR- α and PPAR- γ activators induced expression of the superoxide scavenger enzyme Cu²⁺, Zn²⁺ - superoxide dismutase (Inoue *et al.*, 2001).

Interestingly, despite the extensive PPAR- α mediated anti-inflammatory and anti-atherosclerotic effects, recent studies crossing PPAR- α -deficient mice with apolipoprotein-E-deficient mice found less, rather than more, atherosclerosis (Tordjman *et al.*, 2001). The reasons for these differences between *in-vitro* and *in-vivo* findings are unclear. Certainly, the available evidence with PPAR agonists in clinical trials appears to support beneficial effects, with no obvious evidence for an increase in atherosclerosis among such patients.

1.8 ANIMAL MODELS OF OBESITY

1.8.1 Obesity induced by spontaneous mutations:

Several strains of rodents have been identified and propagated because of spontaneous mutations that resulted in an obese phenotype. The animals are not only useful when studying biology of the underlying a specific mutation, but also studying the mechanism of obesity itself.

1.8.1.1 *ob/ob* (*Lepr^{ob}* / *Lepr^{ob}* mice):

The leptin gene is one of the most extensively studied in obesity research because a spontaneous mutation of this gene that prohibits secretion of mature leptin causing hyperphagia, hypothermia, hypercorticosteronaemia, hyperglycaemia, hypothyroidism, growth hormone deficiency,

hyperinsulinaemia, reduced energy expenditure, decreased linear growth, infertility, and early-onset morbid obesity (Friedman, 1998).

This mutation is a single base substitution that results in premature termination of leptin synthesis. This mouse serves as an ideal model to study both leptin function and morbid early onset obesity. The mutation is rare in humans (Tschop and Heiman, 2001).

1.8.1.2 *db/db (Lepr^{db}/Lepr^{db} mice):*

The Lepr^{db}/Lepr^{db} mice have a phenotype similar to Lepr^{ob}/Lepr^{ob} mice, but they are resistant to leptin because of a mutated leptin receptor. Lepr^{db}/Lepr^{db} mice suffer from morbid obesity with hyperleptinaemia, insulin resistance, and diabetes (Coleman, 1978; Halaas *et al.*, 1995). This mouse provides an excellent model to study leptin resistance, early onset morbid obesity, and is also a useful model to study type II diabetes (Friedman *et al.*, 1997; Tschop and Heiman, 2001).

1.8.1.3 *The fa/fa mutation in Zucker, Koletsky- and Zucker Diabetic Fatty (ZDF) rats:*

Obese Zucker and Koletsky rats have mutations in the extracellular domain of the leptin receptor and exhibit morbid obesity, the *fa/fa* mutation observed in Zucker rats reduces the expression of the leptin receptor on the cell surface with marked intracellular retention, decreased leptin binding and diminished signal transduction. ZDF rats were isolated from a colony of obese Zucker rats and become diabetic after about 8 weeks when presented with a high fat diet (Chua *et al.*, 1996; Friedman *et al.*, 1997). These rats provide an excellent model to study leptin resistance and early onset morbid obesity (Tschop and Heiman, 2001).

1.8.2 Environmentally induced obesity:

In general, genetic models of rodents' obesity show severe insulin resistance and β -cell dysfunction, features which are not usually typical of human type 2 diabetes. This has led to the search for a model more representative of the human form of the disease; i.e. one which has impaired glucose tolerance

(IGT) and mild insulin resistance as a consequence increased intake of a high-energy, highly palatable diet-induced obesity.

Many investigators use wild-type rodents on a high fat diet as a model to study obesity. The rats can be weaned onto obesity-inducing diet (a highly-palatable diet; HPD). Hundred days is the recommended age to start an obesity study, because at this age rats cease to gain lean mass but readily accrue fat mass. Throughout the present study this type of obesity model has been used, which resembles common human obesity better than monogenic rodent models. The increased energy intake is not associated with abnormal leptin or leptin signalling (i.e. absence of a genetic defect in the production of leptin or its receptor). There is extensive literature to show that rats fed high-fat or highly palatable diet (HPD) become progressively more obese than chow-fed littermates, developing hyperleptinaemia and insulin resistance similar to obese human subjects (Chang *et al.*, 1990; Kraegen *et al.*, 1991; Pickavance *et al.*, 1999 ; Harrold *et al.*, 2000). Moreover, the development of obesity on a polygenic background makes this an attractive model.

1.9 AIMS OF THE PROJECT

Our understanding of insulin signal transduction pathways and insulin resistance has evolved extremely rapidly over the past few years. Still, further studies are required to unravel the mechanisms controlling these intricate regulatory processes that presumably mediate, at least in part, the insulin resistance associated with obesity and hyperinsulinaemia.

Metabolic factors associated with obesity, such as free fatty acids can modulate insulin signalling pathways, so this study speculate that there are molecular processes in insulin signalling cascade that are susceptible to inhibition or activation by metabolic factors associated with obesity and thus the spectrum of responses results from the interactions between these factors yielding insulin resistance. Moreover, insulin resistance is associated with accelerated atherosclerosis and a major question is whether or not insulin resistance occurs in cardiovascular tissues and whether this is the driving force for accelerated cardiovascular disease. Furthermore, hepatic and

muscular resistance to the actions of insulin considered the major causative factors in inducing systemic insulin resistance. However, the molecular events of insulin signalling cascades and its correlation to metabolic and vascular abnormalities merit further investigation.

Visceral fat-derived metabolic products, hormones, and cytokines play a major role in effecting insulin action in skeletal muscle and in creating a state of low-grade inflammation and endothelial dysfunction. Some specific and shared insulin signalling abnormalities in muscle, fat, and endothelial cells may also contribute to this rather complex association. Further studies are needed to define the missing steps in the insulin-signalling network, elucidate the mechanisms of cross-communication in the system, and determine the molecular defects that give rise to insulin resistance. Such studies will provide new insights into diabetes and insulin resistance, perhaps allowing a more focused and individualised approach to therapy or prevention of these disorders. Furthermore, Nonpharmacological and pharmacological interventions targeting obesity and/or insulin resistance demonstrate an amelioration of endothelial dysfunction and low-grade inflammation. These strategies, particularly in individuals at risk of type 2 diabetes, may improve clinical cardiovascular outcomes. Investigating the molecular mechanisms related to protein expression of insulin signalling pathways, in addition to determining the vascular and metabolic changes associated with both therapeutical approaches may provide valuable information on the mechanisms and strategies of both types of treatment.

That obesity is associated with vascular dysfunction is well known, but, importantly, such dysfunction can develop after only short-term consumption of an energy-rich diet, long before the onset of frank obesity (Naderali and Williams, 2001b). However, the roles played by different dietary components remains unclear. Thus, the first chapters of the present study investigated the effects of consumption of different high-energy diets of differing macronutrient compositions on the genesis of obesity, induction of vascular abnormalities and metabolic changes.

Also, this study will focus on investigating the protein expressions and deficiencies in the cascade of insulin signalling pathways of insulin resistance vascular and metabolic tissues (aorta, liver, and skeletal muscle) under circumstances related to diet-induced obesity and consumption of high energy diets. Furthermore, the present study will investigate possible association of the above detrimental factors with the development of related clinical disorders such as atherosclerosis and cardiovascular abnormalities, and non-alcoholic fatty liver disease. Moreover, the beneficial effects of the therapeutical approaches of obesity such as non-pharmacological (lifestyle modification, diet withdrawal) and pharmacological (fenofibrate administration) will also be investigated.

It is possible that changes in the levels of protein expression of insulin signaling pathway in vascular and metabolic tissues may induce insulin resistance and alter vasorelaxant property of insulin. Moreover, the level of protein expression of these kinases could fluctuate and respond differently to any pathological, physiological, or pharmacological conditions, such as dietary obesity and its treatment. Therefore, investigating the protein expression of these kinases under certain obesity-related experimental conditions could provide vital information about obesity-induced insulin resistance in vascular and metabolic tissues. Consequently, this study aims to investigate the protein expression of insulin signalling components in aorta, liver and skeletal muscle different experimental groups.

Chapter 2

Methods

2. 1 ANIMAL MODEL

Diet-induced obesity resembles common human obesity better than monogenic rodent models; the increased energy intake is not associated with abnormal leptin or leptin signalling. There is extensive literature to show that rats fed high-fat or highly-palatable diets (HPD) become progressively more obese than chow-fed littermates, and develop insulin resistance and hyperleptinaemia (Wilding *et al.*, 1992a; Pickavance *et al.*, 1999; Harrold *et al.*, 2000) similar to obese humans.

2.1.1 Induction of obesity and energy distribution of the various diets

Out bred male Wistar rats were obtained from Charles River UK Ltd. (Margate, UK), and were selected with a starting weight ranging between 170 and 200 g.

For all experimental protocols unless otherwise stated, animals were fed a highly-palatable diet (HPD) for 7 to 15 weeks to induce obesity (under several experimental conditions), which was defined as a significant increase in fat-pad masses than those of the chow fed controls. The highly-palatable diet provided 1010 KJ/100 g as total energy content, consisted of 33% powdered chow, 33% condensed milk (Nestle UK Ltd., York, UK) and 7% sucrose (Tate & Lyle, London, UK) by weight with the remainder being the tap water. This provided 65% energy as carbohydrate, 19% as protein and 16% as fat (Wilding *et al.*, 1992a). Age-matched controls were fed standard laboratory pelleted chow (chow-fed group)(CRM (P), Biosure, Cambridge, UK) which provide 1310 KJ/100 g as total energy content, consisted of 60% energy as carbohydrate, 30% as protein, and 10% as fat. The diet was made by hand and stored at 4°C. On the other hand, several other types of diets were also included in this study. The High-energy Diet ("Galaxy", milk smooth and creamy chocolate (Mars)) which provided 2256 KJ/100 g as total energy content (42.2% as carbohydrate, 52.7% as fat, and 5.1% as protein) (chocolate-supplemented group). Chocolate-Biscuit supplemented diet ("McVitie's Hobnobs", biscuit covered one side with chocolate) (Biscuit-fed group) which provided 2035 KJ/100g as total energy content (55% as

carbohydrate, 15% as protein and 30% as fat). And fat-enriched diet (Research Diets Inc, New Brunswick, NJ, USA) (high-fat fed group) providing 1980 KJ/100 g as total energy content (35% of energy as carbohydrate, 20% as protein, and 45% as fat) (Table 2.1).

Table 2.1 Summary for energy and micronutrient distribution for the different diets Used in this study

| Type of diet | Total energy (KJ) | | Energy distribution (% & KJ) per 100g | | |
|--|-------------------|------------|---------------------------------------|-----------|----------|
| | Per 1 gm | Per 100 gm | Carbo. | Fat | Protein |
| Standard lab. chow | 13.1 | 1310 | 60 % | 10 % | 30 % |
| | | | 786 KJ | 131 KJ | 393 KJ |
| Highly-palatable diet (HPD), obesity-inducing diet | 10.1 | 1010 | 65 % | 16 % | 19 % |
| | | | 656.5 KJ | 161.6 KJ | 191.9 |
| Highly-energy diet (HED), "Galaxy" | 22.56 | 2256 | 42.2 % | 52.7 % | 5.1 % |
| | | | 952.032 KJ | 1188.9 KJ | 115.1 KJ |
| Chocolate-Biscuit Diet (CBD), "McVitie's" | 20.35 | 2035 | 56 % | 30 % | 15 % |
| | | | 1119.25 KJ | 610.5 KJ | 305.25KJ |
| Fat-Enriched Diet (FED) | 19.8 | 1980 | 35 % | 45 % | 20 % |
| | | | 693 KJ | 891 KJ | 396 KJ |

2.2 EXPERIMENTAL PROTOCOLS

2.2.1 Experimental protocol No. 1

Adult male Wistar rats were randomly assigned to three groups ($n=6/\text{group}$) which, throughout a 10-week period, were each fed diets of differing compositions. This duration was chosen to model human chronic consumption of high-energy food inducing excessive weight gain. Control animals received standard pelleted laboratory chow (chow-fed group). One of the experimental groups received chow supplemented with chocolate biscuit (McVitie's 'HobNobs') (biscuit-fed group), and the second received a fat-enriched diet (high-fat-fed group). Animals were housed singly under controlled environmental conditions (19-22°C; 30-40% humidity) on a 12-h light/dark cycle (lights on at 07:00 h) and had free access to food and water throughout the study. Food intake was measured daily and the body weights weekly. On the day of experiment, the rats were killed by CO₂ inhalation after 2 hours of fasting (Fig. 2.1)

2.2.2 Experimental protocol No. 2

Adult male Wistar rats ($n=21$) were randomized and assigned to two groups which were fed on either a standard pelleted laboratory chow throughout ($n=7$; control group), or given chow plus "Galaxy" chocolate ($n=14$; chocolate-supplemented group). After 14 weeks, half of the chocolate-supplemented group was given fenofibrate (50 mg/kg/day) and the remainder was given vehicle (1% carboxymethyl cellulose at 1 ml/kg body weight; Sigma, UK) by oral gavage daily for 7 days, before being sacrificed. Daily energy intake was determined in all three groups for the last 7 days of the study. On the day of experiment, the rats were killed by CO₂ inhalation after 2 hours of fasting (Fig.2.2)

2.2.3 Experimental protocol No. 3

Adult male Wistar rats ($n=30$) were randomised and assigned to a control group ($n=10$) and a test group ($n=20$). Controls were fed a standard laboratory pelleted diet (chow-fed group), while test group had free access to a highly-

palatable diet. Chow-fed controls remained on their prospective diet for 15 weeks, while after 8 weeks, highly-palatable-diet fed animals were subdivided into two groups. The highly-palatable diet was removed from one group and the standard chow diet was reintroduced (diet-to-chow) for further 7 weeks, while the remaining subgroup was maintained on the highly-palatable diet (diet-fed) for another 7 weeks. After 15 weeks from the start of the study, rats were killed by CO₂ inhalation after 2 hours of fasting (Fig. 2.3).

2.2.4 Experimental protocol No. 4

Adult male Wistar rats (n=30) were randomised and assigned to a control group (n=10) and a test group (n=20) that was fed a highly palatable diet for 15 weeks. Controls were fed a standard laboratory pelleted diet (chow-fed). The test group had free access to a highly-palatable diet. After 8 weeks, half of the test group were given fenofibrate (50 mg/kg/day) and the remainder were given vehicle (1% carboxymethyl cellulose at 1 ml/kg body weight; Sigma, UK) by oral gavage, daily for 7 weeks. On the day of experiment, the rats were killed by CO₂ inhalation after 2 hours of fasting (Fig. 2.4).

2.2.5 Experimental protocol No. 5

Adult male Wistar rats (n=40) were randomized and assigned in the first period of the experiment (8 weeks) to a control group (n=10) and a test group (n=30). Controls were fed standard laboratory pelleted chow (chow-fed group), while the test group had free access to a highly-palatable diet. In the second period (7 weeks), chow-fed controls remained on their prospective diet, while the highly-palatable diet fed animals were subdivided into three subgroups (each subgroup 10 animals). In the first subgroup, highly-palatable diet was removed and the standard chow diet was re-introduced (diet-to-chow), while the second subgroup was given fenofibrate (fenofibrate-treated group, 50 mg/kg/day) and the third group (diet-fed group) was given vehicle (1% carboxymethyl cellulose at 1 ml/kg body weight; Sigma, UK), by oral gavage daily for 7 weeks. On the day of experiment, the rats were killed by CO₂ inhalation after 2 hours of fasting (Fig. 2.5).

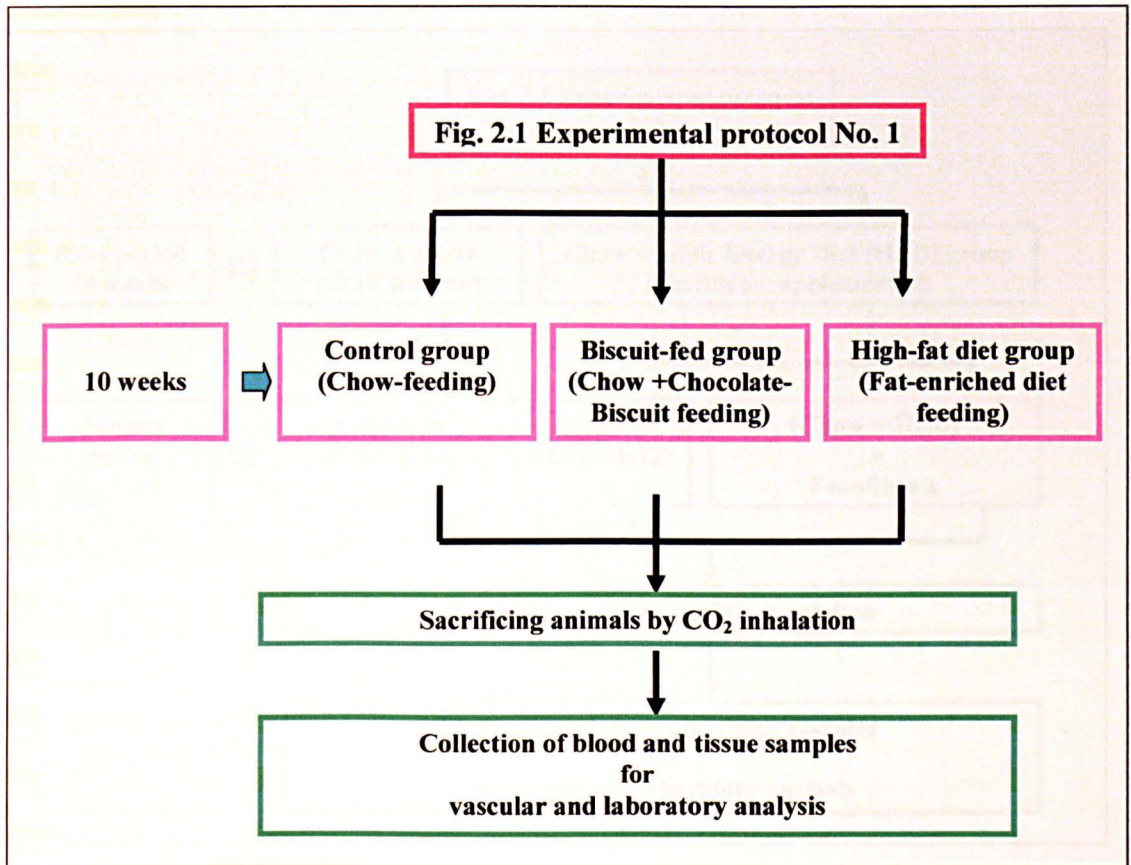


Fig. 2.1. Experimental protocol No. 1

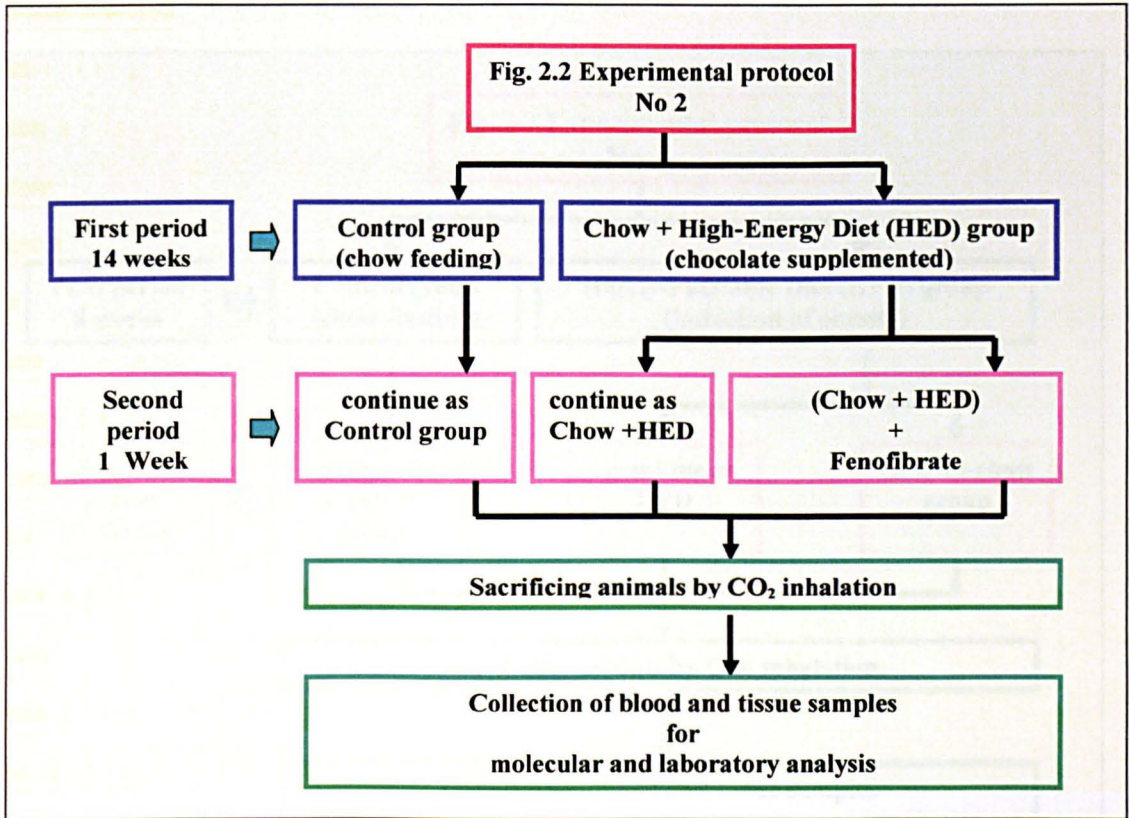


Fig. 2.2. Experimental protocol No. 2

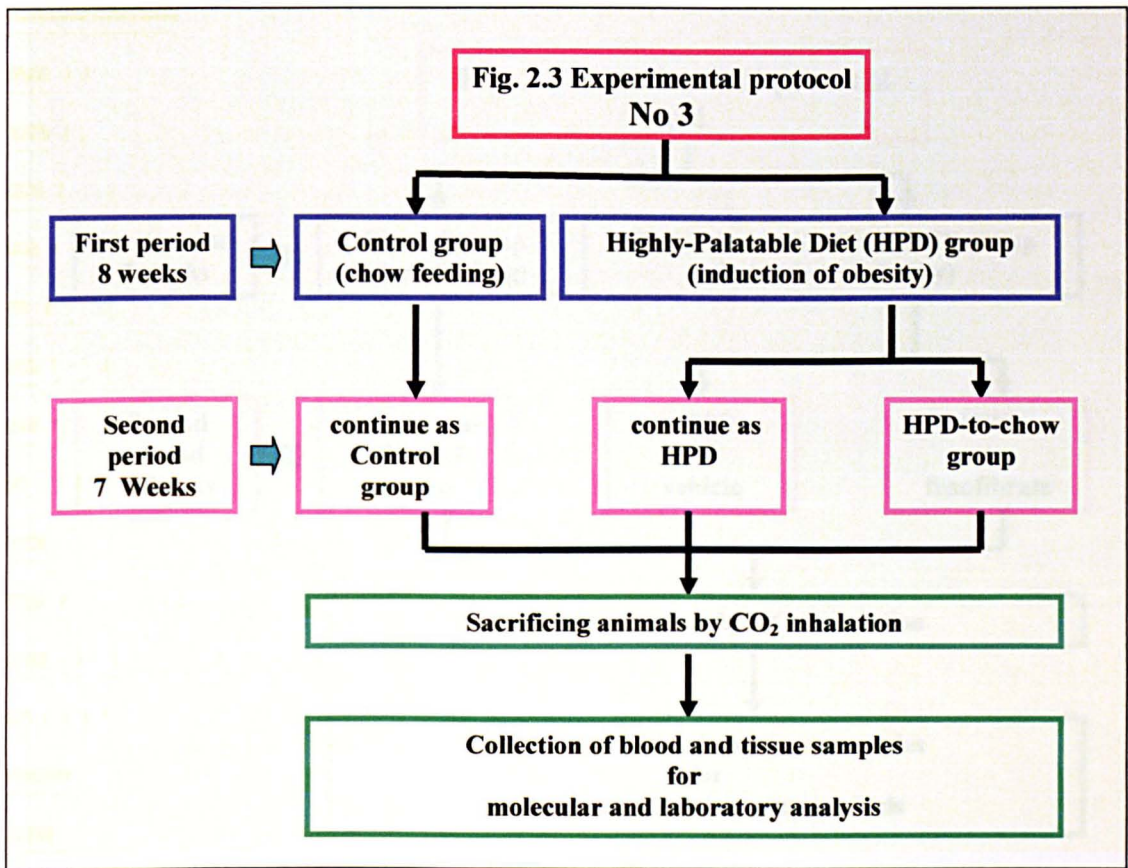


Fig. 2.3. Experimental protocol No. 3

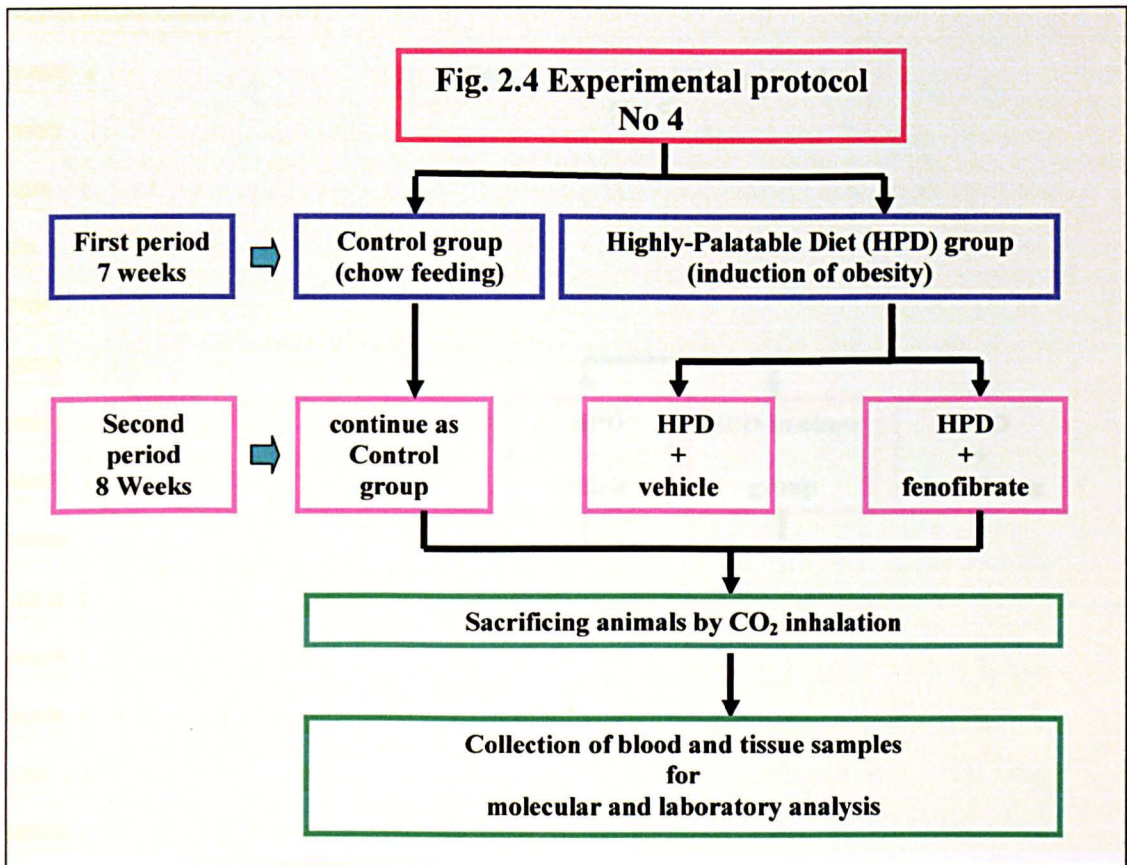


Fig. 2.4. Experimental protocol No. 4

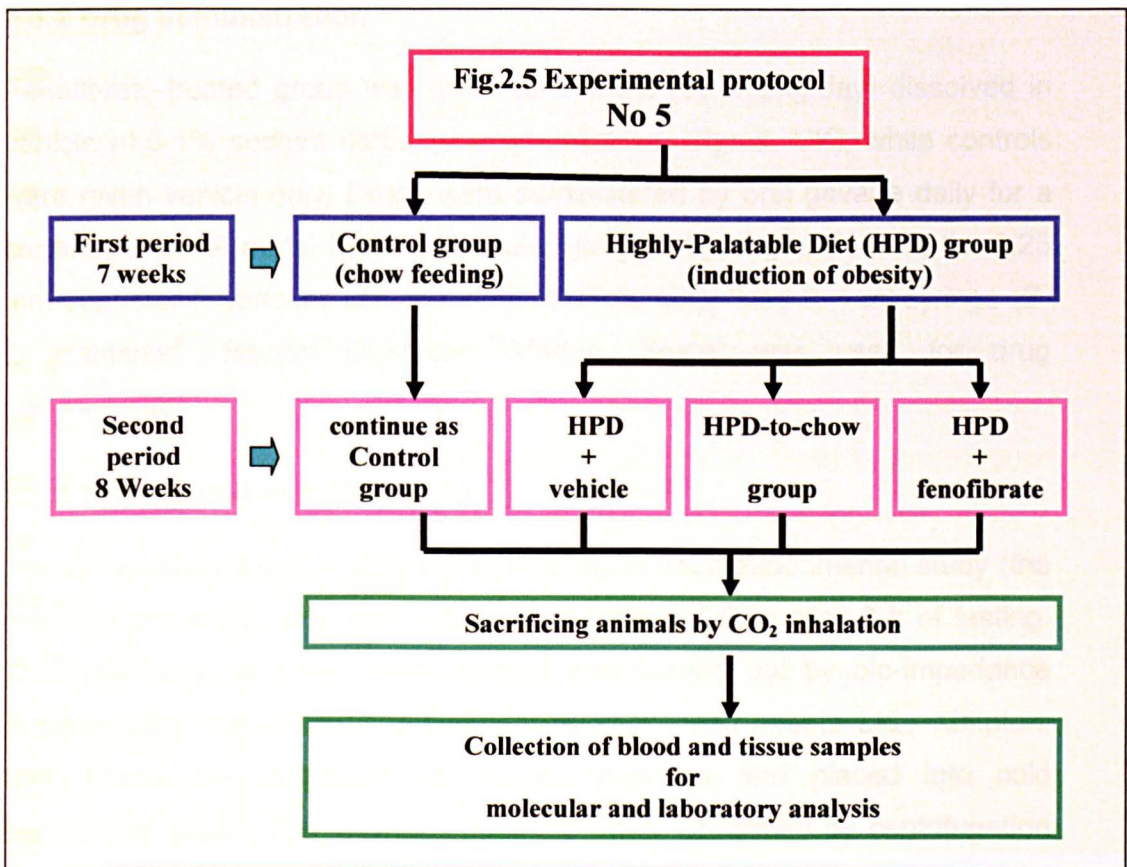


Fig. 2.5. Experimental protocol No. 5

2.3 HOUSING, MAINTENANCE, AND ANIMAL TERMINATION

2.3.1 Housing and maintenance

All animals had free access to tap water throughout the experiment and were housed individually or in groups of 2 in solid-bottomed cages during the period of the experiment, under controlled environmental conditions (19-22 °C; 30-40% humidity) and a 12 h light-dark cycle (lights on at 07:00h).

2.3.2 Drug administration

Fenofibrate-treated group was given fenofibrate (50 mg/kg/day) dissolved in vehicle (0.5-1% sodium carboxymethyl cellulose, Sigma, UK), while controls were given vehicle only. Drugs were administered by oral gavage daily for a certain period. A metal feeding cannula (gauge: 18; length: 50 mm; tip: 2.25 mm diameter (Interfocus Ltd., Haverhill, Suffolk, UK)] fixed to 1-ml syringe (B-D plastipak[®], Becton Dickinson, Madrid, Spain) was used for drug administration.

2.3.3 Termination and dissection

For all experimental protocols, In the last day of each experimental study (the day of termination), rats were killed by inhalation of CO₂ after 2 h of fasting. The total body fat mass measurement was carried out by bio-impedance method using the TOBEC[®] equipment (biotech. Instruments Ltd., Kimpton, UK). Blood was withdrawn by cardiac puncture and placed into cold heparinized tubes. The plasma was immediately separated by centrifugation before being frozen for later measurements of glucose, insulin, leptin, nonesterified free fatty acids (NEFA), triglycerides (TG), C-reactive protein (CRP), adiponectin, and tumour necrosis factor-alpha (TNF-alpha). The aorta, as well as gonadal, epididymal and perirenal fat pads, gastrocnemius and soleus muscles and liver were dissected, weighed and snap frozen in liquid nitrogen for later molecular and experimental studies. The heart and lungs were also dissected and their weights were determined. Six third-order mesenteric arteries were also carefully dissected from each animal for assessment of vascular contractility.

2.4 MOLECULAR BIOLOGY TECHNIQUES

2.4.1 Protein extraction

Fifty milligram of tissue (aorta; liver or gastrocnemius muscle) was homogenized at 4 °C in 500 µl buffer containing 120 mM NaCl, 10% glycerol, 2 mM Na₃VO₄, 1% Nonidet-p40, 1 mM PMSF, 10 mM Na₄p₂O₇, 100 mM NaF, 20 mM Tris (pH 7.6) and a complete mini® protease inhibitor cocktail with polytron homogenizer. The homogenates were then incubated on a rocking platform at 4°C for 30 min. After 3 x 10 second bursts of sonication, tubes were subsequently centrifuged for 45 min at 13000 x g at 4°C. Supernatants were collected, and protein concentrations were determined by Bicinchoninic acid (BCA) method (as per kit instructions-Sigma, UK).

2.4.2 Western immunoblotting

2.4.2.1 SDS-PAGE

A discontinuous acrylamide gel system was used. A stacking gel (5%) was set above a 10% (depending on protein of interest) resolving gel. Samples were standardized to 4-6 mg protein/ml with a lysis buffer. Twenty microlitre of sample protein was boiled in 20 µl 2x electrophoresis sample buffer, for 10 min and then subjected to SDS-PAGE (Tris-glycine buffer, 100 V). Resolved proteins were electro blotted onto nitrocellulose membranes in a buffer containing 25 mM Tris, 190 mM glycine, 1% SDS (W/V) and 20% (V/V) methanol (100 V for 1 hour). Staining immobilized proteins on nitrocellulose with Ponceau S assessed a successful transfer. Nitrocellulose blots were submerged in 0.2% Ponceau S for 15 min with agitation. Blots were then washed with 1 x TBS containing 1% Tween-20 until proteins could be visualized, after which blots were further washed until protein bands had disappeared.

2.4.2.2 Immunoblotting

Non-specific binding proteins were discriminated by incubating the blot with a blocking buffer (5% milk powder, 1x PBS) at room temperature for one hour, followed by immunoblotting with appropriate primary antibody (1:500 dilution) made up in blocking buffer, left overnight at 4 °C. The following morning, blots were washed in 1 x PBS with 1% Tween and incubated with secondary antibody (1:1000), a Horse-reddish peroxidase (HRP) linked anti-rabbit (or

anti-goat, depending on the source of the species used for primary antibody) for 1 hour at room temperature. Proteins were detected using enhanced chemiluminescence method. Positive controls were included for standardization of samples between blots, while molecular weights markers were used for sizing the bands. Densitometry was used to quantify the bands obtained.

2.5 CHEMICALS AND ANTIBODIES

SDS-PAGE and immunoblotting equipments were obtained from Bio-Rad (Richmond, California, USA). Tris, phenylmethyl-sulfonylfluoride (PMSF), aprotinin, dithiothreitol, Tween-20, glycerol, SDS "Lauryl sulfate", B-mercaptoethanol, Bromophenol blue, Temed, ammonium persulfate, and Ponceau S stain, Noradrenaline (NA), carbamylcholine (CCh), sodium nitroprusside (SNP), indomethacin, carboxymethyl cellulose, fenofibrate, and BCA (Bicinchoninic acid) method kit for protein determination were all obtained from Sigma Chemicals (Sigma Ltd., UK). The commercial ELISA kit for measuring plasma TNF- α was obtained from PeproTechEC Ltd. Plasma insulin was measured using ELISA kit from Mercodia. Complete mini® protease inhibitor cocktail, the commercial kits for measuring plasma glucose and triglycerides were all obtained from Roche Diagnostics Ltd. (East Sussex, UK). NaCl, KCl, CaCl₂, MgSO₄, NaCO₃, KH₂PO₄, and EDTA, were all obtained from BDH laboratory supplies (Poole, UK). The commercial kit for measuring plasma C-reactive protein was obtained from Life Diagnostics, Inc. (West Chester (PA), USA). Acrylamide and molecular weight marker were obtained from Bio-Rad laboratories, Inc. (Hercules (CA), USA). Running buffer (Glycine/SDS) and transferring buffer (Tris/glycine) were obtained from National Diagnostics (Atlanta, Georgia, USA). Secondary antibody, an HRP-linked anti-rabbit was purchased from Serotec (Oxford, UK). Polyclonal antibodies against beta subunit of insulin receptor (C-19, sc-711), IRS-1 (C-20, sc-559), IRS-2 (A-19, sc-1556), eNOS (NOS3, C-20, sc-654), Akt1 (C-20, sc-1618), Shc (C-20, sc-288: specific for Shc p46, p52 and p66) and ERK1 (C-16, sc-93: reactive with ERK1 p44 and, to lesser extent, ERK2 p42) and PI 3-kinase were purchased from Santa Cruz Biotechnology (Santa Cruz,

California, USA). Polyclonal antibody against PKC- β was purchased from UPSTATE (UPSTATE house, Dundee, UK).

NA, CCh and SNP were dissolved in double distilled water, while indomethacin was dissolved in ethanol. NA and SNP were kept away from the light throughout the experiment. All water-soluble solutions were freshly made on the day of experiment.

2.6 HOMEOSTASIS MODEL ASSESSMENT (HOMA)

HOMA, an index of insulin resistance, employs the measures of fasting plasma concentrations of glucose and insulin. Thus, terminal concentrations of glucose and insulin were used to calculate an index of insulin resistance where insulin concentration is expressed in units of $\mu\text{U/ml}$ and glucose in mM.

$$[\text{insulin} \times \text{glucose}] / 22.5$$

2.7 ASSESSMENT OF VASCULAR FUNCTION

2.7.1 Myograph technique set up

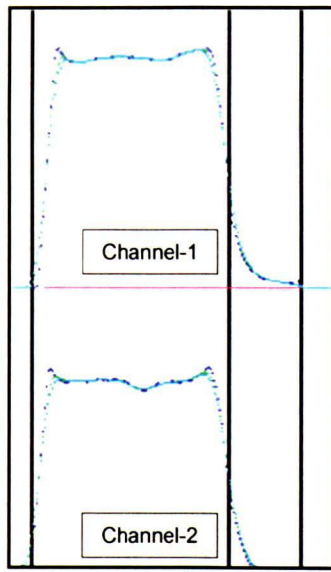
Six third-order mesenteric arteries (150-220 μm diameter, 2 mm length) were carefully dissected from each animal. Each artery was freed of fat and connective tissue and mounted on two 40- μm diameter stainless-steel wires in an automated myograph (Cambustion, Cambridge, UK), based on the principle of Mulvany's myograph which measures isometric tension generated in response to various stimuli. Arterial segments (two at a time) were incubated in a 5-ml organ bath containing physiological salt solution (PSS; composition [in mM]: NaCl 119; KCl 4.7; CaCl_2 2.5; MgSO_4 1.17; NaHCO_3 25; KH_2PO_4 1.18; EDTA 0.026; and glucose 5.5) gassed with 95% O_2 and 5% CO_2 at 37 °C. After 30 min equilibration, the length-tension characteristics for each vessel were determined using the law of Laplace ($P = T/r$, where P is the transmural pressure, T is the tension, and r is the vessel radius). Each vessel was then set to its normalized diameter, i.e. the diameter it would achieve at rest *in vivo* under a transmural pressure of 100 mmHg. The computer also calculated the target tension that each vessel should develop in response to a

maximal stimulus. Arteries were then allowed a further 30 min to equilibrate before being depolarised twice with high-potassium physiological salt solution (KPSS) in which the NaCl in normal PSS was replaced by an equimolar concentration of KCl (125 mM). Cumulative concentration–response curves to either KCl (10–125 mM) or noradrenaline (NA, 0.5–6 μ M) were then carried out. Any vessel failing to reach its predetermined target tension in response to vasoconstriction with KCl (125 mM) was discarded.

However, examples of the original myograph tracing involving the administration of vasoconstrictors and vasodilators to illustrate the experiment and data acquisition have been shown in the figures (2.6 to 2.10).

2.7.2 Assessment of defect in vascular relaxation

Endothelium-dependent and -independent defects were investigated by exposing NA-precontracted arteries to carbamylcholine (CCh) or sodium nitroprusside (SNP), respectively. To eliminate possible interference by vasoactive prostanoids, 10 mM indomethacin was added to the organ bath, prior to contraction with NA (8 μ M). When contraction reached a plateau (after 2 min), concentration–response curves were performed to either CCh, histamine or SNP (for all, 10 nM–100 μ M).



[KCl] 125 mM

Fig. 2.6. An example for the original myograph tracing for KCl (125mM).

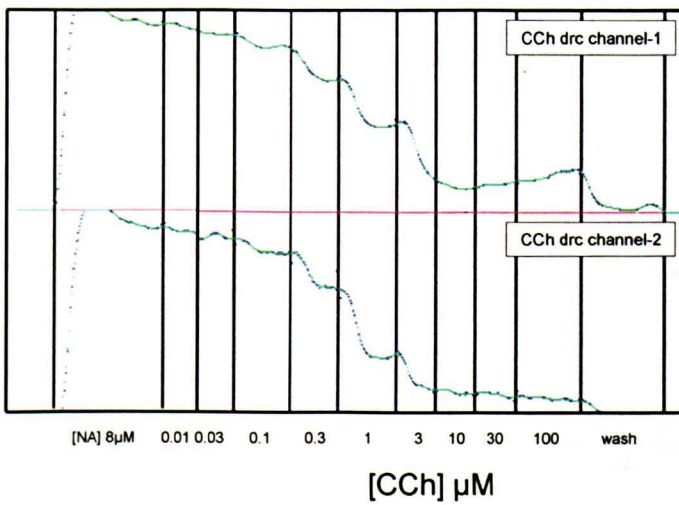


Fig. 2.7. An example of the original myograph tracing for CCh (10nM -100µM) dose-response curve.

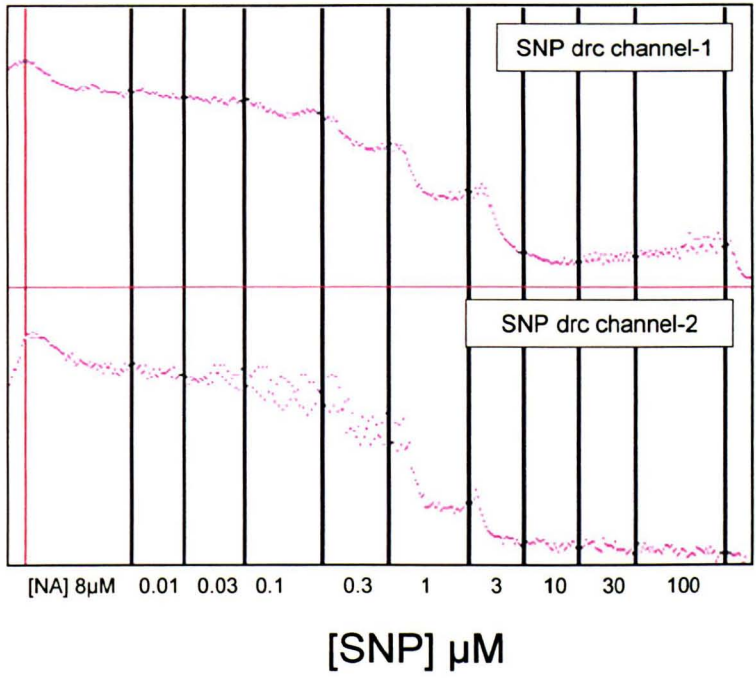


Fig. 2.8. An example of the original myograph tracing for SNP (10nM-100 μM) dose-response curve.

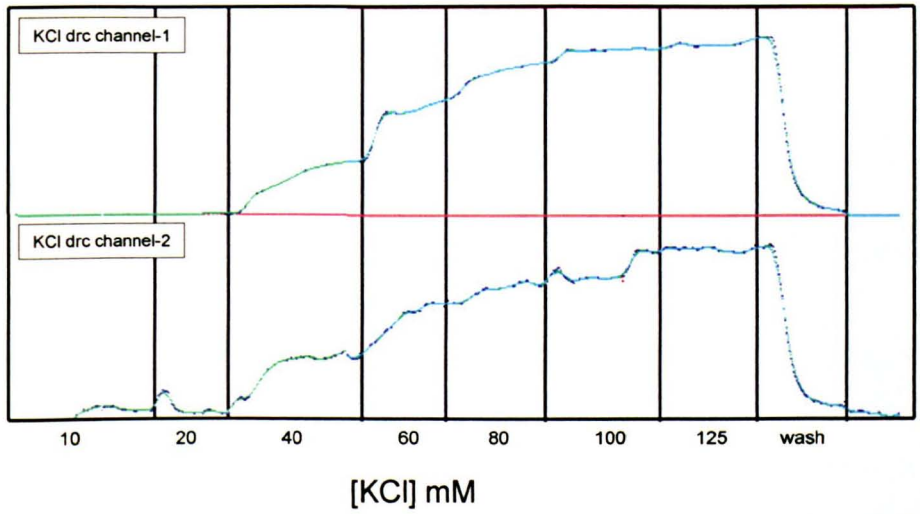


Fig. 2.9. An example of the original myograph tracing for KCl (10-125mM) dose-response curve.

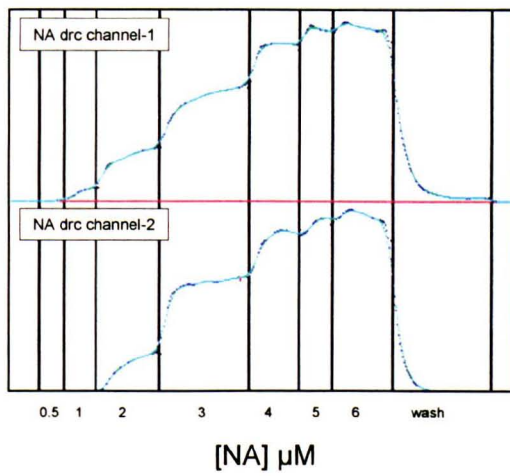


Fig.2.10. An example of the original myograph tracing for NA (0.5-6 μM) dose-response curve.

2.8 DATA INTERPRETATION AND STATISTICAL ANALYSIS

Changes in body weights of each group were collected weekly and are expressed as absolute total body weights. For Western blotting, the data from chow-fed (control) animals were expressed as 100% response, and the results from other test groups (in each experiment) were normalized and subsequently expressed as the percentage of their respective controls.

In myograph study, vasoconstriction in response to NA and KCl were expressed as absolute force generated. Relaxation in responses to CCh, SNP, or histamine were calculated as the percentage reduction from the maximal tension generated at the supramaximal concentration of NA (8 μ M). The average response for all the vessels from a given animal was determined before group analysis.

In the vascular experiment (myograph), in order to determine the statistically significant difference between the experimental groups, ANOVA test was used, specifically, MANOVA tests were applied to perform multiple dataset analysis. Multivariate analysis of variance (MANOVA) is simply ANOVA with several dependent variables. However, the main objective of using MANOVA is to determine if the response variables are altered by the observer's manipulation of the independent variables. If the overall multivariate test is significant, we could conclude that the respective effect is significant.

In all experiments, data are expressed as mean \pm standard error of mean (SEM). Data have normal distribution (Shapiro Wilk W test). Statistical significance was tested using Student t-test' or repeated-measures (Bratanova-Tochkova *et al.*, 2002), Bonferroni "t-test" or the Mann-Whitney test, as appropriate. Results were considered statistically significant at the $p < 0.05$ levels. All statistical analysis were performed using Microsoft Office Excel 2003.

2.9 LIVER HISTOLOGY

The tissue samples of liver were prepared for histological examinations using ordinary tissue processing and stained by Haematoxylin and Eosin (H and E) stains protocol. Ten liver specimens (10 animals) of each experimental group were interpreted by a single pathologist who was blinded to all data of liver samples and the experimental groups under investigation. The liver samples were analysed microscopically for the presence or absence of the following histological features: steatosis, steatohepatitis, fibrosis and hepatocellular carcinoma. Histological features were interpreted according to the schema outlined by Brunt et al. (Brunt *et al.*, 1999). Briefly, steatosis was graded on a 3-point scale: grade 1=steatosis involving < 33 % of hepatocytes, grade 2= 33-66%, grade 3 > 66%. Inflammation (steatohepatitis) was graded on a 4 point scale: grade 0=no or negligible inflammation, grade 1=mild, grade 2=moderate, grade 3=severe. Fibrosis was staged on a 5 point scale: stage 0=no fibrosis, stage 1=zone 3 perisinusoidal/ perivenular fibrosis, stage 2=zone 3 and periportal fibrosis, stage 3=septal/bridging fibrosis, stage 4=cirrhosis.

2.9.1 Procedure of liver histology

2.9.1.1 Tissue processing:

- 1- Liver tissues were fixed overnight in 10% neutral buffered formalin (NBF).
- 2- Tissue samples were dehydrated with ascending grade of alcohol (70%, 80%, 90%, and 100%). Each for an hour.
- 3- Tissue were then cleared with xylene; two changes each in an hour.
- 4- Tissues were impregnated with molten paraffin wax two changes each in an hour.
- 5- Tissues were then picked from molten paraffin and embedded in embedding paraffin to obtain tissue embedded paraffin blocks.
- 6- Tissues were then cut using microtome into 4 μ m thickness and now the samples ready for H and E staining.

2.9.1.2 Haematoxylin and Eosin staining protocol:

- 1- Slides were deparaffinised by immersing in xylene, two changes 3 minutes each.
- 2- Slides were rehydrated in descending grades of alcohol (100% two changes 5 minutes each, 90%, 70%, 5 minutes each).
- 3- Sections were left in running water for 5 minutes.
- 4- Sections were stained in haematoxylin by dipping for few minutes.
- 5- Sections were placed in running water until water looks clean and sections became blue.
- 6- Sections were placed in 1% acid alcohol (99 ml 70% alcohol + 1 ml concentrated HCl) for two seconds.
- 7- Sections were washed in running tap water.
- 8- Sections were stained with eosin for 2 minutes.
- 9- Sections were washed in running tap water.
- 10- Sections were dehydrated in ascending grades of alcohol, cleared in xylene two changes.
- 11- Sections were mounted with DPX.

2.10 PLASMA AND TISSUE MEASUREMENTS AND ASSAYS

2.10.1 Plasma Insulin

2.10.1.1 Principle

Mercodia rat insulin ELISA is a solid phase two-site enzyme immunoassay. It is based on the direct sandwich technique in which two monoclonal antibodies are directed against separate antigenic determinants on the insulin molecule. During incubation, insulin in the sample react with peroxidase-conjugate anti-insulin antibodies and anti-insulin antibodies bound to microtitration well. A simple washing step removes unbound enzyme labelled antibody. The bound conjugate is detected by reaction with 3,3',5,5'-tetramethylbenzidine. The reaction is stopped by adding acid to give colorimetric endpoint that is read spectrophotometrically.

2.10.1.2 Assay procedure

Plasma insulin concentrations were measured using a commercial ELISA kit. Standards (0.0; 0.15; 0.4; 1; 3; and 5.5 µg/l) and samples were added in duplicate to 96 well microtitre plate coated with anti-insulin antibodies and mixed with 50 µl conjugate solution, and incubated in a shaker for 2 hours at room temperature. The reaction volume was aspirated, and 350 µl washing solution was added to each well. The washing solution was then aspirated completely. The aspiration and washing procedure was repeated 5 times, after which to all wells, 200 µl of peroxidase substrate was added, and the plate was then incubated for further 15 minutes. Finally, 50 µl of stop solution was added to the wells. The plate was shaken for approximately 5 seconds to ensure mixing of substrate and stop solution. The absorbance was measured at 450 nm using 96 well plate reader. The concentrations of the samples were measured as µg/l from the obtained standard curve.

2.10.2 Plasma Leptin

2.10.2.1 Principle

Plasma leptin concentrations were measured with a commercial radioimmunoassay (RIA) kit (mouse leptin RIA; Linco, Dorset, UK). Mouse anti-leptin serum exhibits 100% cross-reactivity with rat leptin.

2.10.2.2 Assay procedure

Standards and samples (100 µl) were added to GL4 tubes (Beaumont, Leicester, UK) in duplicate and mixed with 100 µl of mouse leptin antibody. Tubes were vortexed, covered and left to incubate overnight at 4 °C. 100 µl of ¹²⁵I-mouse leptin was added; tubes were vortexed and incubated overnight at 4°C. Precipitating reagent (1 ml) was then added, and tubes were vortexed and incubated for 20 min at room temperature. After which tubes were centrifuged at 3000 x g at 4°C for 40 min. the tubes were decanted in one movement, careful not to lose the pellet. Bound radioactivity was counted on an automated gamma counter (Cobra auto-gamma; Berks, UK). The intra-assay coefficient of variation was below 10% and the detection limit was 0.2 ng/ml.

2.10.3 Plasma TNF- α

2.10.3.1 Principle

Plasma TNF- α concentrations were measured using a commercial ELISA kit (PeproTechEC Ltd, London, UK). Rat TNF- α development kit contains the key components required for the quantitative measurement of rat TNF- α in a sandwich ELISA format within the range of 32 - 3000 pg/ml.

2.10.3.2 Assay procedure

The standard was diluted from 3 ng/ml to zero in diluent. 100 μ l of standard or sample was added to each well in duplicate, and incubated at room temperature for at least 2 hours. The solution in the plate was then aspirated and the plate washed 4 times. 100 μ l detection antibody, diluted in diluent to a concentration of 0.5 μ g/ml, was added to each well. The plate was then incubated for 2 hours at room temperature. The solution was aspirated again and the wells washed further 4 times. 100 μ l of avidin peroxidase (1: 2000 diluted) was added to each well. The plate was then incubated at room temperature for further 30 minutes.

The plate then aspirated and washed further 4 times. Substrate solution (100 μ l) was added to each well and the plate incubated at room temperature for colour development. The colour development was monitored with an ELISA plate reader at 405 nm. The concentrations of the samples were measured as pg/ml from the obtained standard curve.

2.10.4 Plasma C-Reactive Protein (CRP)

2.10.4.1 Principle

The rat CRP ELISA is based on a solid phase enzyme-linked immunosorbent assay (Filippatos *et al.*, 2005). The assay uses affinity purified anti-rat CRP antibodies for solid phase (microtiter wells) immobilization and HRP conjugated anti-rat CRP antibodies for detection. The test sample is diluted and incubated in the microtiter wells for 45 minutes. The microtiter wells are subsequently washed and HRP conjugate is added and incubated for 30 minutes. These results in CRP molecules being sandwiched between the

immobilization and detection antibodies. The wells are then washed to remove unbound HRP-labeled antibodies and TMB reagent is added and incubated for 20 minutes at room temperature. This results in the development of blue colour. Colour development is stopped by the addition of stop solution, changing the colour to yellow, and optical density is measured spectrophotometrically at 405 nm. The concentration of CRP is proportional to the optical density of test sample.

2.10.4.2 Assay procedure

Standards (31.25, 15.6, 7.8, 3.9, 1.95, 0.98, and 0 ng/ml) and samples were added in duplicate to the wells of microtiter plate which coated with anti-rat CRP antibody, and incubated on an orbital micro-plate shaker at 100-150 rpm at room temperature (18-25 °C) for 45 minutes. The incubation mixture then removed by flicking plate into an appropriate Bio-waste container. The microtiter wells were rinsed and flicked 6 times with distilled water. The wells were then struck sharply onto absorbent paper towels to remove all residual water droplets. Enzyme conjugate reagent (100 µl) was added to each well and incubated on an orbital micro-plate shaker at 100-150 rpm at room temperature (18-25 °C) for further 30 minutes. The wells were washed as detailed above (by flicking, rinsing, and striking the wells sharply onto absorbent paper or towels to remove all residual water droplets). TMB reagent (100 µl) was then added to each well and gently mixed on an orbital micro-plate shaker at 100-150 rpm at room temperature for another 20 minutes. The reaction was stopped by adding 100 µl of stop solution to each well and gently mixing (it's important to make sure that all the blue colour changes to yellow). The optical density was read at 405 nm with a microtiter plate reader within 15 minutes. The concentrations of the samples were measured as ng/ml from the obtained standard curve.

2.10.5 Triglyceride content of muscle and liver tissues

Muscle tissue (160-200 mg) or liver tissue (80-100 mg) were homogenised in 500 µl distilled water with polytron homogeniser. The homogenate were centrifuged for 5 minutes at 13,000 x g, and the supernatants were transferred

to eppendurff tubes. The triglycerides assay was performed on the supernatant to detect the concentration of TG in muscle or liver tissue.

2.10.6 Plasma triglycerides:

2.10.6.1 Principle

The series of reactions involved in the assay system are as follows:

Triglycerides + H₂O ----- Lipase ----- Glycerol + Free Fatty Acids

The glycerol is phosphorylated by adenosine triphosphate (ATP) with glycerol kinase (GK) to produce glycerol-3-phosphate and adenosine diphosphate (ADP).

Glycerol + ATP ----- GK ----- Glycerol-3-phosphate ADP

Glycerol-3-phosphate is oxidized to dihydroxyacetone phosphate (DAP) by glycerol phosphate oxidase (GPO) producing hydrogen peroxide (H₂O₂).

Glycerol-3-phosphate + O₂ ----- GPO ----- DAP + H₂O₂

in a trinder type colour reaction catalyzed by peroxidase (POD), the H₂O₂ reacts with 4-aminoantipyrine (4 AAP) and 3,5 dichloro-2 hydroxybenzene sulfonate (3,5 DHBS) to produce a red colored dye. The absorbance of this dye is proportional to the concentration of triglycerides present in the sample.

H₂O₂ + 4AAP + 3,5DHBS ----- POD ----- Quinoneimine dye + 2H₂O

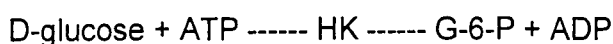
2.10.6.2 Assay procedure

Serial dilutions of standards were prepared as following: 125; 62.5; 31.25; 15.512; 7.756; and 0.0 mg/dl. In 96 well micro titre plate, 3 µl of each sample or standard in duplicate plus 300 µl of reagent solution was added to the wells. The plate was then incubated for 5 minutes at room temperature. Finally, the absorbance was read at 520 nm using micro titre plate reader and the concentrations of unknown samples were obtained from the standard curve.

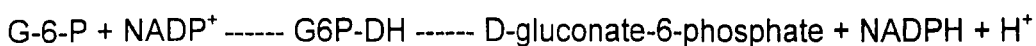
2.10.7 Plasma Glucose

2.10.7.1 Principle

The test kit for glucose, non-colorimetric (Roche Diagnostics; Milton Keynes, UK) is based on the following principle: D-glucose is phosphorylated to D-glucose-6-phosphate (G-6-P) in the presence of the enzyme hexokinase (Bratanova-Tochkova *et al.*, 2002) and adenosine-5'phosphate (ATP), with the simultaneous formation of adenosine-5'-diphosphate (ADP):



In the presence of the enzyme glucose-6-phosphate dehydrogenase (G6P-DH), G-6-P is oxidised by nicotinamide-adenine dinucleotide phosphate (NADP) to D-gluconate-6-phosphate with the formation of reduced nicotinamide-adenine dinucleotide phosphate (NADPH):



The amount of NADPH formed in this reaction is stoichiometric to the amount of D-glucose. The increase in NDAPH is measured by means of its light absorbance at 340 nm in the UV range.

2.10.7.2 Assay procedure

First, plasma samples had to be de-proteinised. This involved mixing 1 ml of 3% perchloric acid with 100 μ l of samples in a 1.5-ml Eppendorf tube and centrifuging for 5 minutes at 5000 rpm. Then, 1 ml of a solution containing NADP and ATP was added to an empty 3-ml plastic cuvette (the "blank") as well as to all the cuvettes containing 100- μ l volumes of the de-proteinised plasma solution. The volume was made up to 3 ml by adding 2 ml of distilled water to all cuvettes. The final concentrations of NADP and ATP were 0.8 mg/ml and 1.9 mg/ml, respectively. The solutions were mixed and the first set of absorbencies (A1) read at 340 nm at room temperature after about 3 min, referencing against the blank. Next, 20 μ l of a suspension containing Hexokinase (Bratanova-Tochkova *et al.*, 2002) and glucose-6-phosphate-dehydrogenase (G6P-DH) (final concentrations 1.94 U/ml and 0.97 U/ml, respectively) we added to each cuvette and the solutions mixed. After allowing

the reaction to proceed for 15 min (the endpoint of reaction). The second set of absorbencies were read (A2) at 340 nm. These absorbencies were then applied to the general equation for calculating concentrations (C):

$$C = V \times \Delta A / v \times d \times \epsilon$$

Where

V = final volume (ml)

v = sample volume (ml)

d = light path (cm)

ϵ = absorption coefficient of the dyestuff at the wavelength,; in this case,
 ϵ =extinction coefficient of NADPH at 340 nm = 6.3

ΔA = the absorbance difference (A2 – A1)

This reduces to the formula

$$52.49 \times \Delta A$$

To yield the concentration of glucose in plasma in mM units.

2.10.8 Free Fatty Acid (FFA)

2.10.8.1 Principle

FFAs are, in the presence of the enzyme acyl-CoA synthetase (acyl CS), converted by ATP and coenzyme A (CoA) into acyl-coenzyme A (acyl-CoA), resulting in adenosine-5'-monophosphate (AMP) and pyrophosphate. Acyl CoA reacts with oxygen (O₂) in the presence of acyl-CoA oxidase to form 2,3-enoyl-coenzyme A (enoyl-CoA).

The resulting hydrogen peroxide (H₂O₂) converts 2,4,6-tribromo-3-hydroxybenzoic acid (TBHB) and 4-aminoantipyrine to a red dye in the presence of peroxidase (POD). The dye is measured in the visible range at 546 nm.

FFAs + CoA + ATP ----- Acyl CS ----- acyl-CoA + AMP + pyrophosphate



2.10.8.2 Assay procedure

One milliliter of a solution containing ATP, CoA, acyl CS, POD and 4-AAp and TBHB was added to an empty 2-ml plastic cuvette (the blank) as well as to cuvettes containing 50 µl volumes of sample. The volume of the blank was made up with 50 µl of distilled water. The solutions were mixed and allowed to sit for 10 minutes at room temperature. Fifty µl of a solution containing N-ethyl-maleinimide was then added to all cuvettes for the removal of an existing surplus of CoA before the oxidation of the activated FFAs by ACOD. The solutions were then mixed and the first set of absorbencies (A1) read at 546 nm measured against water. The reaction was started by adding 50 µl of a solution containing ACOD to all cuvettes. The solutions were mixed again and the end of the reaction reached at room temperature after 15 min. the second set of absorbencies (A2) were then read, again at 546 nm. The absorbance differences (A2 – A1) were calculated for both blank and unknown (plasma) and applied to the formula:

$$\Delta A = \Delta A_s - \Delta A_b,$$

Where ΔA_s and ΔA_b are the absorbance differences of each unknown sample and the blank respectively. These volumes were then applied to the formula

$$1.192 \times \Delta A$$

Derived from the general formula for calculating the concentration outlined above (in glucose assay) and where:

$$\epsilon = \text{absorption coefficient of the red dye at 546 nm} = 19.3$$

to yield the concentration of FFAs in plasma in mM units.

Chapter 3

Results: Differential effects of high-energy diets on vascular reactivity precede significant obesity: Combined high-carbohydrate and high-fat induces greater vascular dysfunction than high-fat alone

3.1 Introduction

Obesity is an important risk factor for atheroma, and is thought to be a major factor in the development of hypertension (Lakka *et al.*, 2002). Arterial dysfunction and structural abnormalities, ranging from alterations in release of vasoactive substances from endothelial cells to changes in arterial wall elasticity and agonist-induced vasodilatory responses, have been seen in obese human subjects and in animal models that reflect some aspects of human obesity (Cowan *et al.*, 1991; Steinberg *et al.*, 1996; Arcaro *et al.*, 1999; Dobrian *et al.*, 2000; Perticone *et al.*, 2001; Naderali, 2001b). Thus, the link between obesity and vascular dysfunction is well-established. Critically, however, many of the components of the metabolic syndrome, including cardiovascular disorders, are manifest in preobesity (Ota *et al.*, 2002). Indeed, acute vascular dysfunction is evident in humans after only a couple of high-energy meals (Ong *et al.*, 1999), and animal models reveal that irreversible vascular impairment can arise as early as two days after consumption of a high-energy diet (Naderali, 2001b; Naderali, 2003).

The causes of vascular dysfunction in obesity and related disorders remain uncertain, but plasma non-esterified fatty acids (NEFAs), triglycerides (TGs), leptin and insulin, all of which are increased in obesity, are prime candidates (Despres *et al.*, 1996; Steinberg, 1997; Tka *et al.*, 1997; Glueck *et al.*, 1999; Soderberg *et al.*, 1999; Wallace *et al.*, 2001; Naderali, 2001b). Of these, only NEFAs and TGs are raised by short-term high-energy feeding, suggesting that they may play the more critical role (Naderali, 2001b; Naderali, 2003). Consumption of diets high in either carbohydrate or saturated fat results in atherogenic dyslipidaemia (German and Dillard, 2004), further suggesting a link between dietary components and vascular abnormalities. It is less clear whether or not these components have different effects on vascular function and, if so, what these are.

It is reasonable to suggest that the current obesity epidemic, at least in the Western world, is due to consumption of energy-dense foods which are relatively high in both carbohydrate and fat, rather than one or the other.

Thus, it was of value to test the vascular effects of such a diet. As a secondary aim, it was also of value, whilst exploring the obesity-inducing effects of various diets in our laboratory, to identify those which, after a prolonged period of consumption, failed to induce significant obesity.

The hypothesis, therefore, is that different dietary components would differentially effect vascular function, that a combination of such components would exacerbate dysfunction, and that these effects would be measurable in a prevailing phase of overweight. This would be shown by measuring vascular function in rats fed diets high in various atherogenic components for a prolonged period.

3.2 METHODS

3.2.1 Experimental protocol

Adult male Wistar rats (192 ± 4 g) were randomly assigned to three groups ($n=6$ /group) which, throughout a 10-week period, were each fed diets of differing compositions. This duration was chosen to model human chronic consumption of high-energy food while inducing weight gain, representing a state of obesity. Control animals received standard pelleted laboratory chow, providing 63% of energy as carbohydrate, 28% as protein and 10% as fat (chow group). One of the experimental groups received chow supplemented with chocolate biscuit, providing 56% of energy as carbohydrate, 15% as protein and 30% as fat (biscuit-fed group), and the second received a fat-enriched diet, providing 35% of energy as carbohydrate, 20% as protein and 45% as fat (high-fat-fed group). A more detailed breakdown of the macronutrient composition of the diets is given in Table 1. Animals were housed singly under controlled environmental conditions. Food intake was measured daily and body weight weekly.

At the end of the experiment, after two hours fasting, the rats were killed by CO₂ inhalation. A single epididymal fat pad and gastrocnemius muscle was dissected free from each animal and weighed. As an index of body composition, these values were expressed as a ratio. Blood was collected into cold heparinized tubes and plasma separated by centrifugation and stored

frozen for later measurement of analytes such as: glucose, insulin, leptin, non-esterified fatty acids (NEFA) and triglyceride (TG).

3.2.2 Assessment of vascular function

Six third-order mesenteric arteries were carefully dissected from each animal and mounted on two 40- μm diameter stainless-steel wires in an automated myograph (Cambustion, Cambridge, UK). Pairs of arteries were incubated in a 5-ml organ bath containing physiological salt solution (PSS) gassed with 95% O_2 and 5% CO_2 at 37 °C.

After a 30-min equilibration, the length-tension characteristics for each vessel were determined, as described previously in chapter two, section 2.7.1. Arteries were allowed a further 30 min to equilibrate before being depolarized twice with high-potassium PSS (KPSS), in which NaCl in normal PSS was replaced by an equimolar concentration of KCl (final $[\text{K}^+] = 125 \text{ mM}$). Any vessel failing to reach its predetermined target tension in response to KPSS was discarded. Arterial contractility was evaluated from concentration-response curves to KCl (10-125 mM) and noradrenaline (NA; 0.5-6 μM).

Arterial relaxation was measured in NA-pre-constricted arteries following exposure to two endothelium-dependent vasodilators, carbamylcholine (CCh) and histamine, and an endothelium-independent vasodilator, sodium nitroprusside (SNP). Arterial segments were contracted with a supramaximal concentration of NA (8 μM). When contraction reached a plateau (after 2 min), responses to CCh, histamine and SNP over the range of 10 nM-100 μM were carried out in random order (for more details about the assessment of vascular function, see chapter two, sections 2.7.1 & 2.7.2).

3.2.3 Statistical analyses

Vasoconstriction in response to NA and KCl was expressed as absolute force generated, whilst relaxation in response to CCh, histamine and SNP concentrations was calculated as percentage reduction from the maximal tension induced by 8 μM NA.

Data are expressed as mean \pm S.E.M. An average response for all the vessels from a given animal was determined before group analysis. EC₅₀ values were determined to evaluate changes in agonist-induced responses. Significance of statistical differences between vascular responses were tested using repeated measures analysis of variance (Bratanova- *et al.*, 2002), followed by *post-hoc* Bonferroni *t*-tests. Differences between diet-fed groups and chow-fed controls for all other parameters were tested by student's (unpaired) "*t*-test" or Mann-Whitney test, as appropriate. Within-group differences were analysed by paired "*t*-test". Vasorelaxation data were first quantified by calculation of area under the curve (AUC) using the trapezoidal rule. Differences were considered statistically significant at $p < 0.05$.

3.3 RESULTS

3.3.1 Body weight, food intake and metabolic data

Animals fed the biscuit- or high-fat diet gained more weight than their chow-fed counterparts, although this did not reach significance (+13% and 16%, respectively; both $p > 0.05$; Fig. 3.1). This was in accordance with higher daily energy intakes for both groups (+27% and 8%), but only that of biscuit-fed animals rose significantly ($p < 0.01$). Consistent with these findings, both biscuit- and high-fat-fed groups had increased epididymal fat pad masses (+72% and 48% vs controls, Fig. 3.4), although only significantly so in the former ($p < 0.05$). In contrast, gastrocnemius muscle mass was comparable in all three groups (Fig. 3.5), resulting in a significant change in fat/lean ratio (body composition) only on the biscuit-supplemented diet ($p < 0.05$ vs controls; Fig. 3.6).

Although neither diet affected circulating concentrations of glucose, insulin or leptin (all $p > 0.05$ vs chow-fed controls, Figs. 3.7 – 3.9), triglycerides rose by 41% in the biscuit-fed group ($p < 0.05$). The high-fat diet increased circulating NEFAs by twice the proportion of the biscuit-supplemented diet (+38% and 17% vs controls; $p < 0.01$ and $p < 0.05$, respectively; Fig. 3.11).

3.3.2 Arterial contractile responses

Neither of the diets altered vessel diameter (chow-fed: $190 \pm 10 \mu\text{m}$; biscuit-fed: $200 \pm 10 \mu\text{m}$; high-fat-fed: $200 \pm 15 \mu\text{m}$; both $p > 0.05$ vs chow-fed). KCl- and NA-induced vasoconstriction in arteries from all three groups produced a characteristic sigmoid relationship. Both agonists increased vasoconstriction in the biscuit-fed group ($p < 0.01$ and $p < 0.001$, respectively), but had no effect in the high-fat-fed group (both $p > 0.05$). Maximal KCl- and NA-induced contractions induced were the following, respectively: chow-fed, 5.29 ± 0.52 and 6.58 ± 0.89 mN/mm; biscuit-fed, 6.79 ± 0.58 and 11.14 ± 0.11 mN/mm; high-fat-fed, 5.25 ± 0.21 and 6.55 ± 1.1 mN/mm (Figs. 3.12 & 3.13).

3.3.3 Endothelium-dependent vasorelaxation

NA-precontracted arteries from chow-fed rats demonstrated progressive relaxation to cumulative addition of CCh (10 nM-100 μM), achieving a maximum of $89 \pm 5\%$ at a concentration of 100 μM , with EC_{50} values of $0.11 \pm 0.01 \mu\text{M}$. Arteries from biscuit-fed and high-fat-fed rats that were similarly exposed to CCh displayed a significant rightward shift of concentration-response curves compared with those of chow-fed controls (EC_{50} biscuit-fed: 0.28 ± 0.01 ; high-fat-fed: $0.52 \pm 0.04 \mu\text{M}$; both $p < 0.001$; Fig. 3.14). Maximal vasorelaxation to 100 μM CCh was reduced by 13% in both biscuit- and high-fat-fed rats (both $p < 0.01$ vs chow-fed), but was more attenuated in arteries from high-fat- than biscuit-fed rats at lower CCh concentrations (3.16-316 nM; $p < 0.01$; Fig. 3.14). Histamine-induced vasorelaxation was similar in all three groups, the concentration-response curves being virtually superimposable. Maximum vasorelaxations induced (at 100 μM) were $81 \pm 3\%$, $86 \pm 5\%$ and $75 \pm 7\%$ for chow-, biscuit- and high-fat-fed rats, respectively (Fig. 3.15).

3.3.4 Endothelium-independent vasorelaxation

The endothelium-independent vasodilator, SNP, also induced concentration-dependent vasorelaxation in arteries from all three groups, the curves being significantly shifted to the right in the biscuit- and high-fat-fed groups (EC_{50} biscuit-fed: 1.58 ± 0.02 ; high-fat-fed: 0.24 ± 0.01 ; both $p < 0.01$ vs chow-fed: 0.13 ± 0.02). However, maximum SNP-induced responses were similar in all

three groups. Compared with the chow-fed control and high-fat-fed groups, SNP-induced relaxation in biscuit-fed rats was attenuated by more than 2-fold (AUC for chow-fed: 10.89 ± 0.31 ; biscuit-fed: 3.59 ± 0.23 ; high-fat-fed: 8.91 ± 0.07 ; $p < 0.001$, biscuit-fed vs both control and high-fat-fed). Although the attenuation of SNP responses was less pronounced in high-fat-fed animals, it nevertheless reached statistical significance (AUC for high-fat vs control: $p < 0.05$; Fig. 3.16).

Table 3.1 Breakdown of digestible macronutrients (% of energy) in different diets. Values were calculated from manufacturers' nutritional analyses. Summed macronutrient values may not exactly match total values due to small amounts of calorific ingredients which do not fit under these categories and/or to slightly discrepant values given in manufacturers' analysis sheets. Abbreviations: CHO = carbohydrate.

| Groups | Total CHO | Complex CHO | Simple CHO | Total fat | Saturated fat | MUFA | PUFA | Total protein |
|-----------------|-----------|-------------|------------|-----------|---------------|------|------|---------------|
| Chow | 60.0 | 55.7 | 4.3 | 10.0 | 1.7 | 3.0 | 5.3 | 30.0 |
| Biscuit | 55.0 | 37.0 | 18.0 | 30.0 | 12.9 | 10.6 | 5.4 | 15.0 |
| High-fat | 35.0 | 16.9 | 16.9 | 45.0 | 39.1 | 0.0 | 5.5 | 20.0 |

Table 3.2 Daily macronutrient intake (kJ) in male Wistar rats fed different diets for 10 weeks. Statistical significance vs chow controls: * $p < 0.05$; ** $p < 0.01$; *** $p < 0.0001$ (unpaired t -test; $n = 6$ /group). Abbreviations: CHO = carbohydrate; MUFA = monounsaturated fatty acid; PUFA = polyunsaturated fatty acid.

| Diet | Overall total | Total CHO | Complex CHO | Simple CHO | Total fat | Saturated fat | MUFA | PUFA | Total protein |
|-----------------|---------------|--------------|-------------|-------------|--------------|---------------|-------------|-----------|---------------|
| Chow | 361.1+15.3 | 227.6+9.6 | 208.5+8.8 | 19.1+0.8 | 37.2+1.6 | 6.1+0.2 | 11.0+0.5 | 20.0+0.8 | 101.6+4.3 |
| Biscuit | 459.0+13.6** | 255.6+6.8 | 167.8+3.2** | 88.1+4.2*** | 140.6+6.5*** | 60.8+3.1*** | 50.0+2.4*** | 24.8+0.7* | 67.0+1.2*** |
| High-fat | 390.2+13.3 | 136.6+4.7*** | 66.3+2.3*** | 66.3+2.3*** | 175.6+6.0*** | 152.5+5.2*** | 0.0 | 21.5+0.7 | 78.0+2.7** |

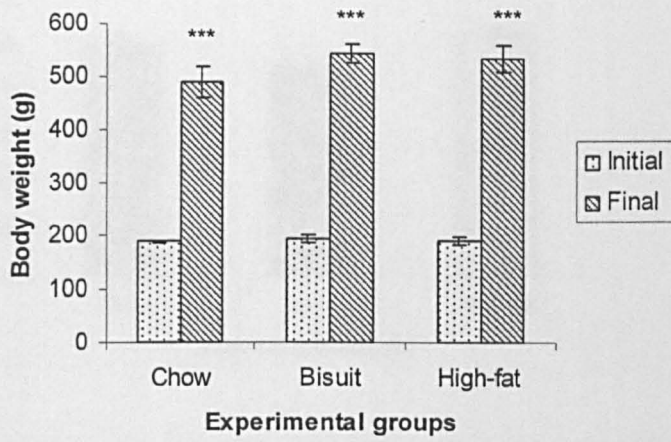


Fig. 3.1. Comparison between initial and final body weight of three experimental groups: chow, biscuit, and high-fat feeding. Statistical significance vs initial body weight: *** $p < 0.0001$ (paired t-test; $n=6$ /group).

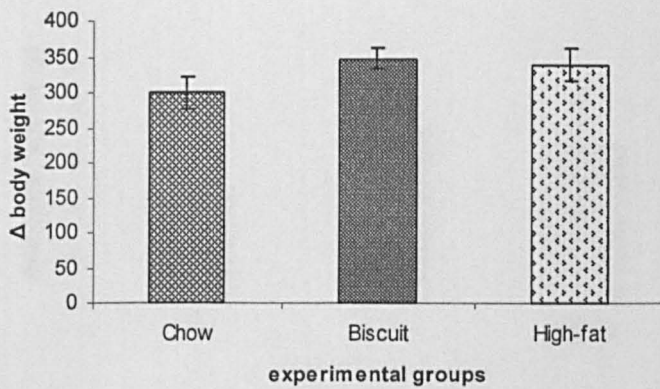


Fig. 3.2. Δ body weight of the three experimental groups: chow, biscuit, and high-fat.

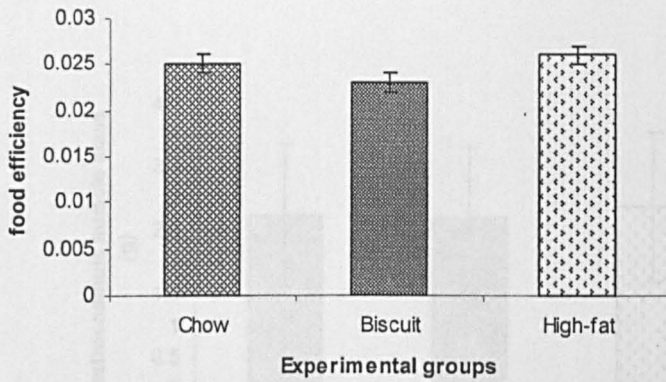


Fig. 3.3. Food efficiency for the three experimental groups: chow, biscuit, and high-fat feeding. There were no significant differences in comparison to chow feed control.

Fig. 3.5. Gastrocnemius muscle mass was measured in all three experimental groups: chow, biscuit, and high-fat. No significant differences were seen between the three groups.

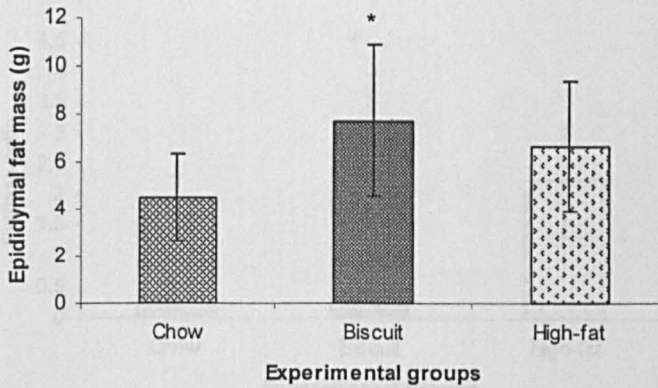


Fig. 3.4. Epididymal fat mass was measured in all three experimental groups: chow, biscuit, and high-fat feeding. The fat mass in biscuit group was significantly more heavier than that of chow-fed control.

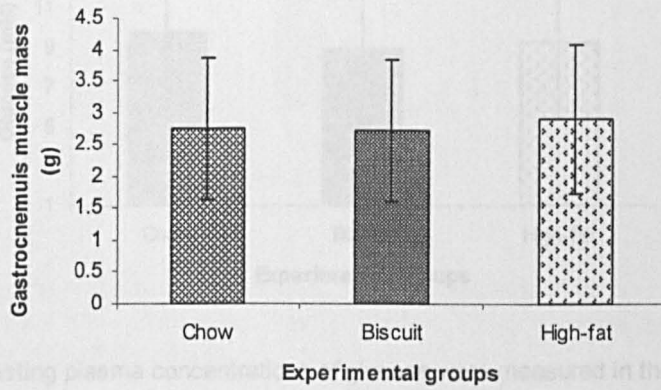


Fig. 3.7. Fasting plasma concentration of insulin was measured in the experimental groups: chow, biscuit, and high-fat feeding. No significant differences were seen between the three groups.

Fig. 3.5. Gastrocnemius muscle mass was measured in all three experimental groups: chow, biscuit, and high-fat. No significant differences were seen between the three groups.

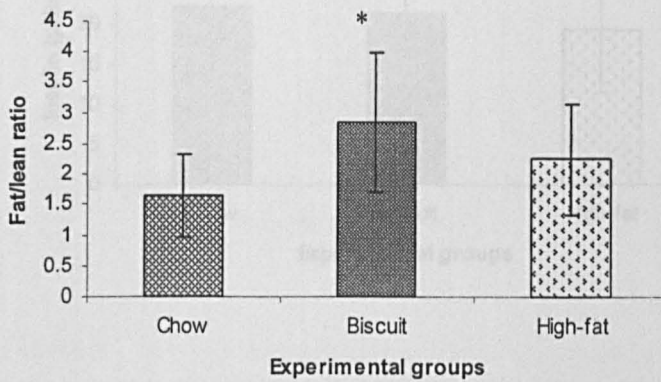


Fig. 3.6. Fasting plasma concentration of insulin was measured in the experimental groups: chow, biscuit, and high-fat feeding. No significant differences were seen between the three groups.

Fig. 3.6. Fat/lean ratio for the three experimental groups: chow, biscuit, and high-fat feeding. There was significant increase in biscuit group in comparison with chow fed control.

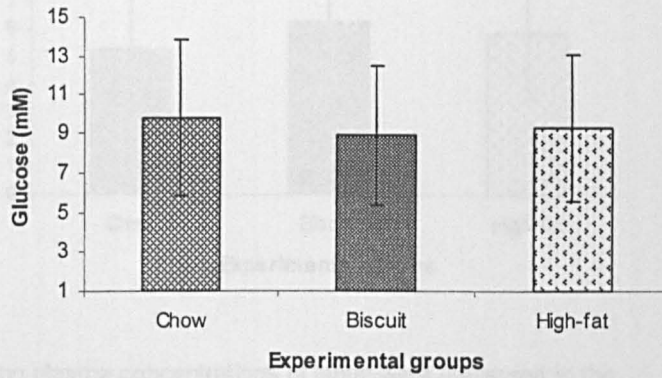


Fig. 3.7. Fasting plasma concentrations of glucose were measured in the experimental groups: chow, biscuit, and high-fat feeding. No significant differences were seen between the three groups.

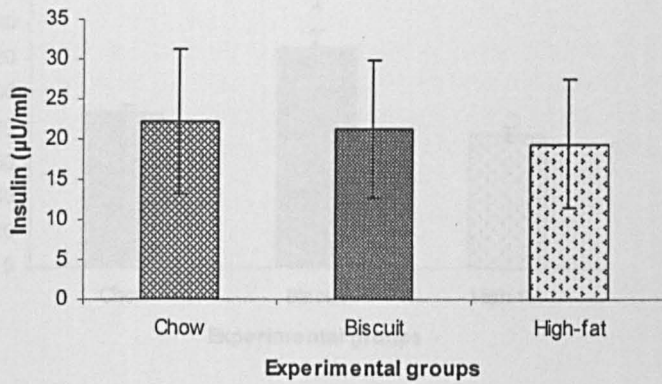


Fig. 3.8. Fasting plasma concentrations of insulin were measured in the experimental groups: chow, biscuit, and high-fat feeding. No significant differences were seen between the three groups.

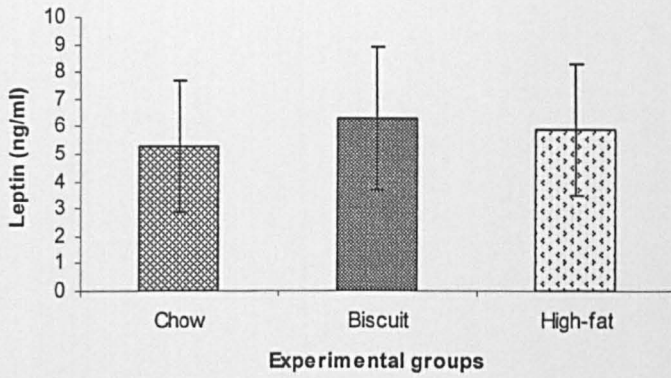


Fig. 3.9. Fasting plasma concentrations of leptin were measured in the experimental groups: chow, biscuit, and high-fat feeding. No significant differences were seen between the three groups.

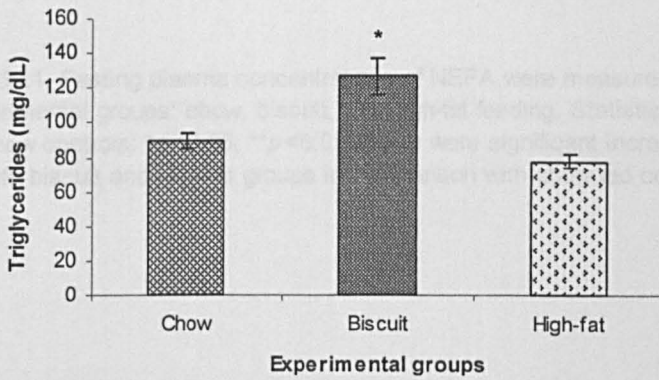


Fig. 3.10. Fasting plasma concentrations of triglycerides were measured in the experimental groups: chow, biscuit, and high-fat feeding. Statistical significance vs chow controls: * $p < 0.05$. Biscuit- but not high-fat-fed group had significantly higher TG than their chow-fed counterpart.

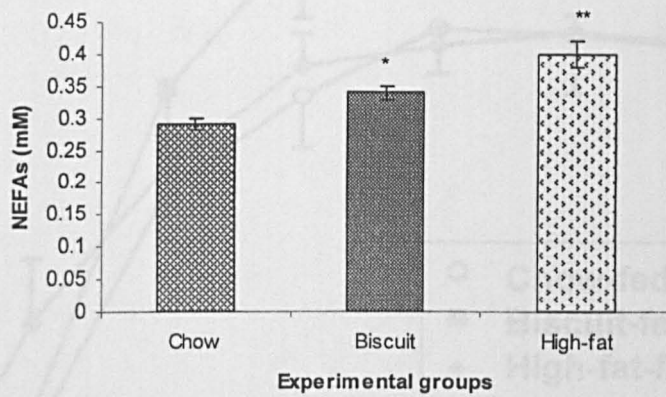


Fig. 3.11. Fasting plasma concentrations of NEFA were measured in the three experimental groups: chow, biscuit, and high-fat feeding. Statistical significance vs chow controls: * $p < 0.05$; ** $p < 0.01$. There were significant increase in the levels of both biscuit and high-fat groups in comparison with chow fed control.

Fig. 3.12 Concentration-response curves for KCl (10-125 mM) in arteries from male Wistar rats fed chow, biscuit-supplemented or high-fat diet for 10 weeks. Data are expressed as mean \pm S.E.M. Compared with chow-fed controls, KCl-induced vasoconstriction was significantly augmented in arteries from biscuit-fed animals (* $p < 0.01$, ANOVA (Bonferroni test); $n = 6$ /group).

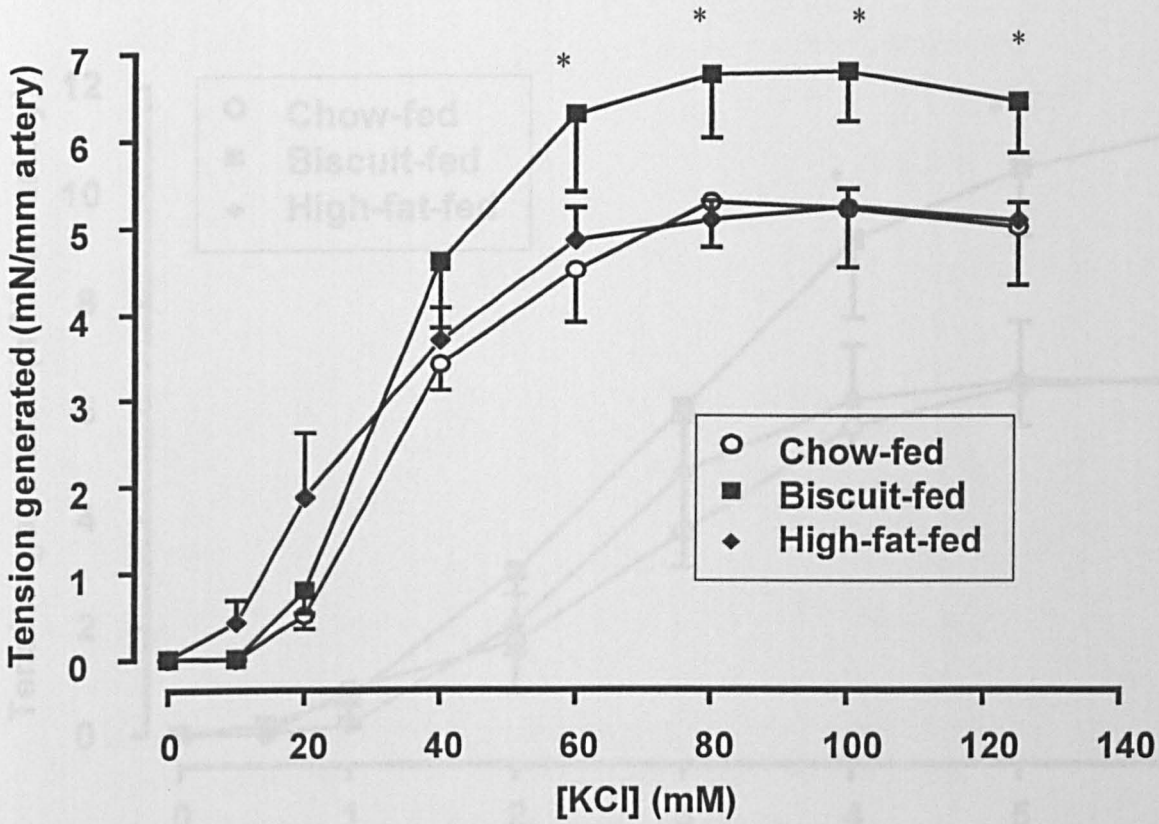


Fig. 3.12 Concentration-response curves for KCl (10-125 mM) in arteries from male Wistar rats fed chow, biscuit-supplemented or high-fat diet for 10 weeks. Data are expressed as mean \pm S.E.M. Compared with chow-fed controls, KCl-induced vasoconstriction was significantly augmented in arteries from biscuit-fed animals (* $p < 0.01$; ANOVA (Bonferroni test); $n = 6$ /group).

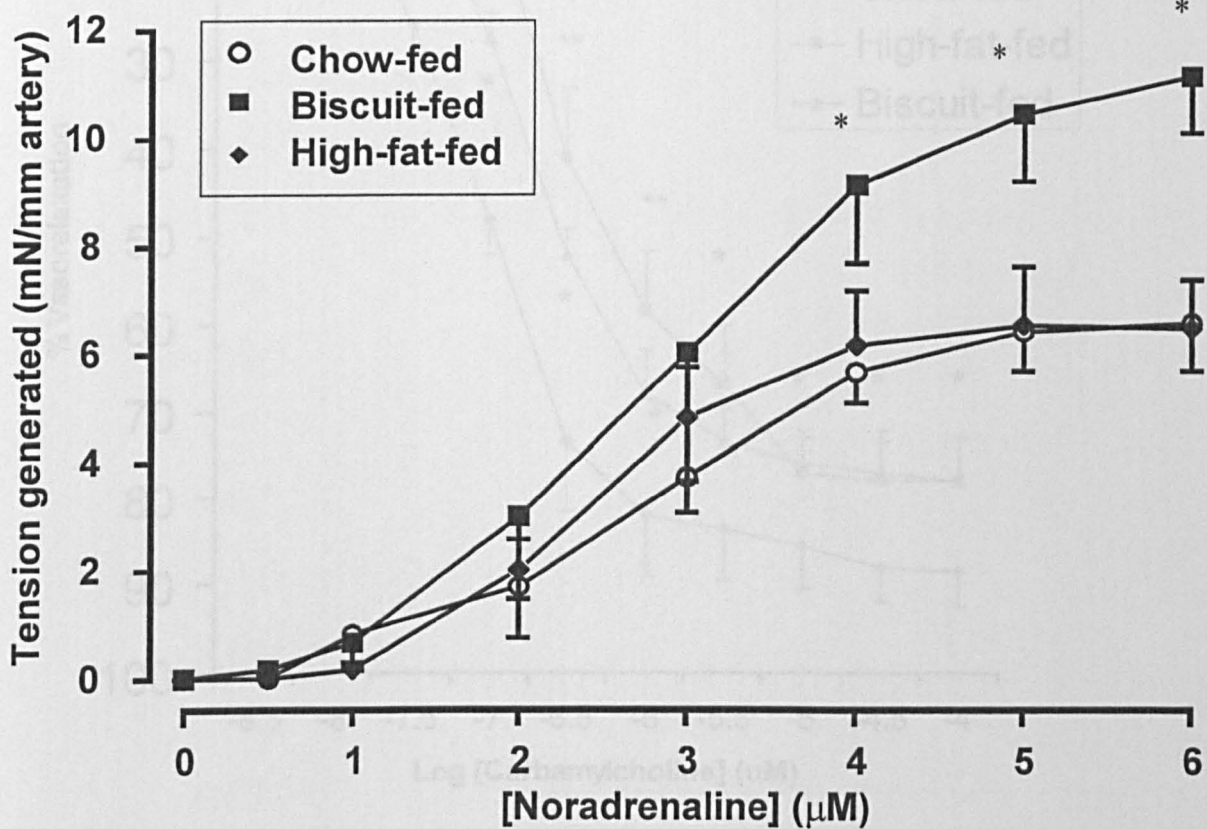


Fig. 3.13 Concentration-response curves for noradrenaline (NA; 0.5-6 μM) in arteries from male Wistar rats fed chow, biscuit-supplemented or high-fat diet for 10 weeks. Data are expressed as mean \pm S.E.M. Compared with chow-fed controls, NA-induced vasoconstriction was significantly augmented in arteries from biscuit-fed animals ($*p < 0.01$; ANOVA (Bonferroni test); $n = 6/\text{group}$).

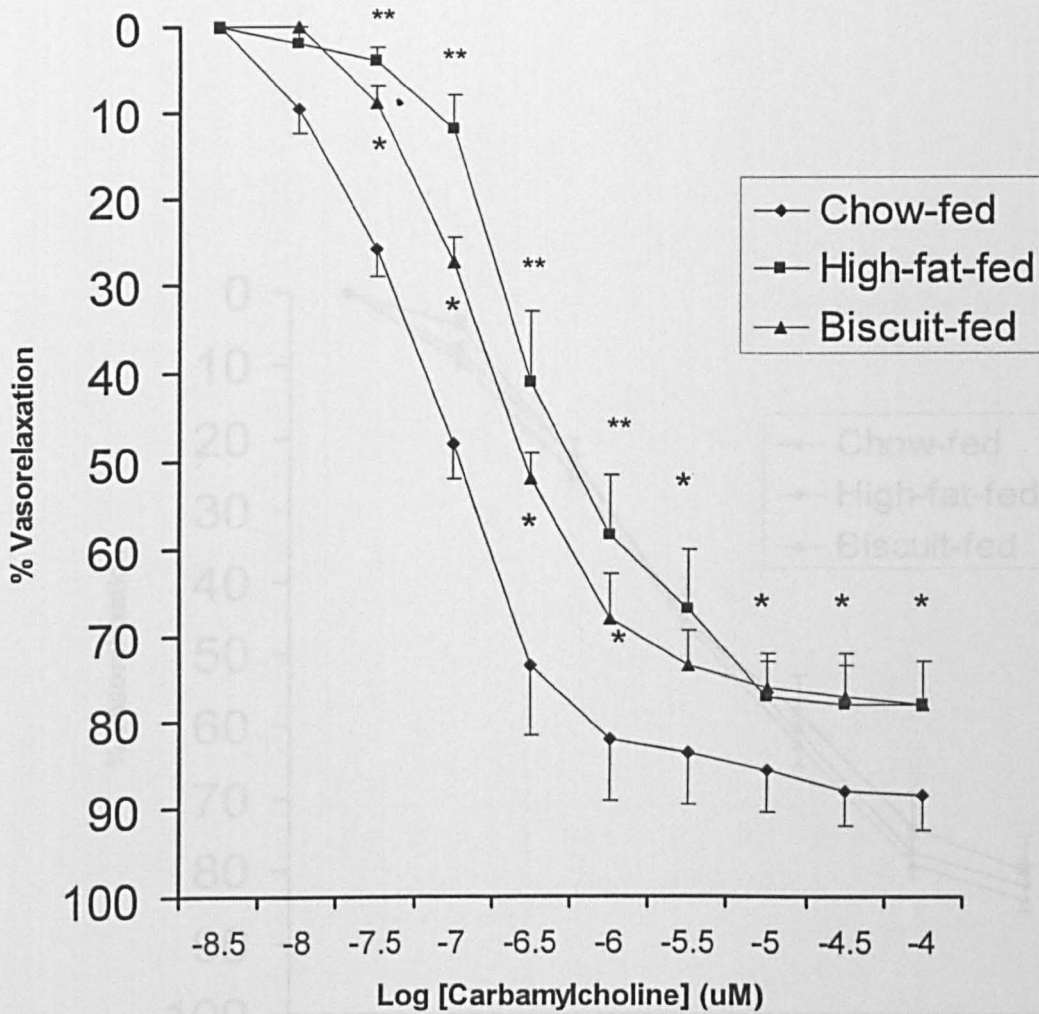


Fig. 3.14 Concentration-response curves for carbamylcholine in noradrenaline (NA)-precontracted arteries from male Wistar rats fed chow, biscuit-supplemented or high-fat diet for 10 weeks, as described in Methods. Data represent mean \pm S.E.M. Compared with chow-fed controls, vasorelaxation in response to CCh was significantly attenuated in biscuit-fed and high-fat-fed animals (* $p < 0.01$ and ** $p < 0.001$, respectively; ANOVA (Bonferroni test); $n = 6$ /group).

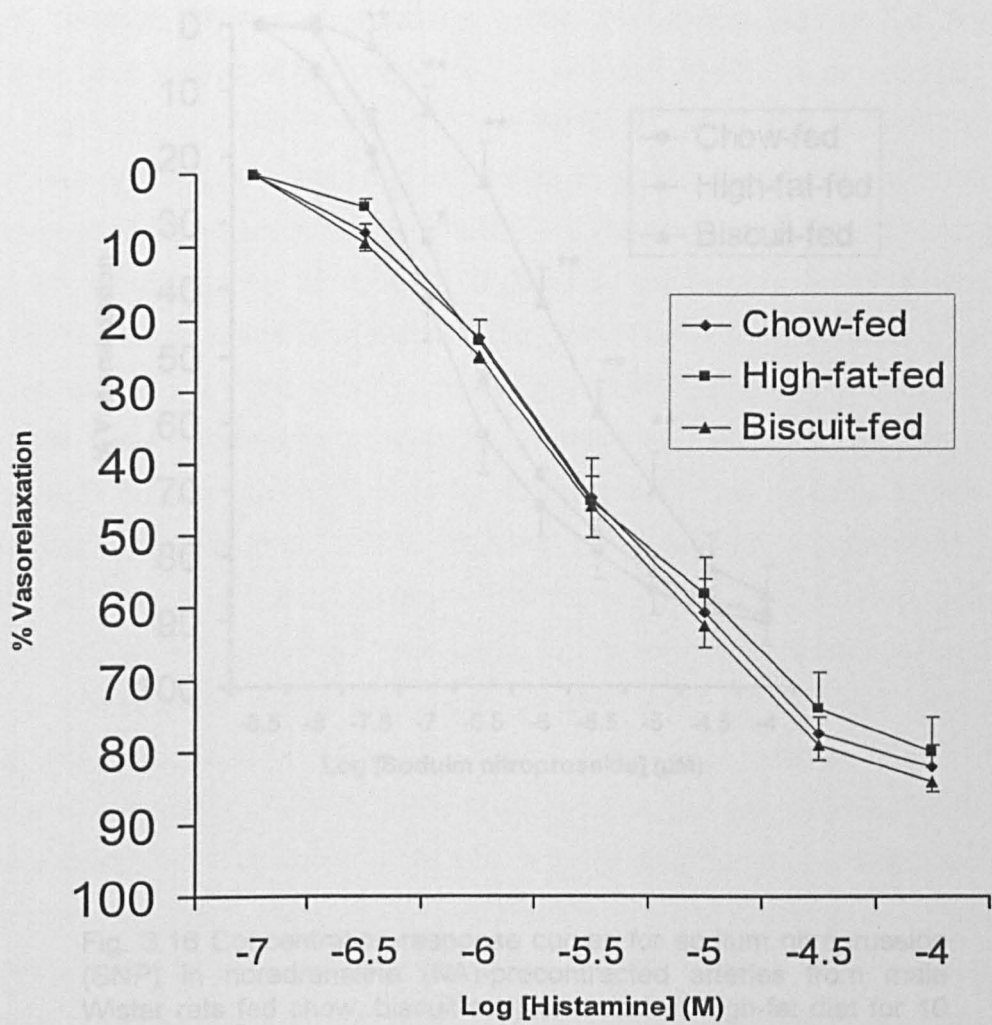


Fig. 3.15 Concentration-response curves for histamine in noradrenaline (NA)-precontracted arteries from male Wistar rats fed chow, biscuit-supplemented or high-fat diet for 10 weeks, as described in Methods. Data represent mean \pm S.E.M. No significant changes were seen between all three groups (n=6/group).

3.4 DISCUSSION

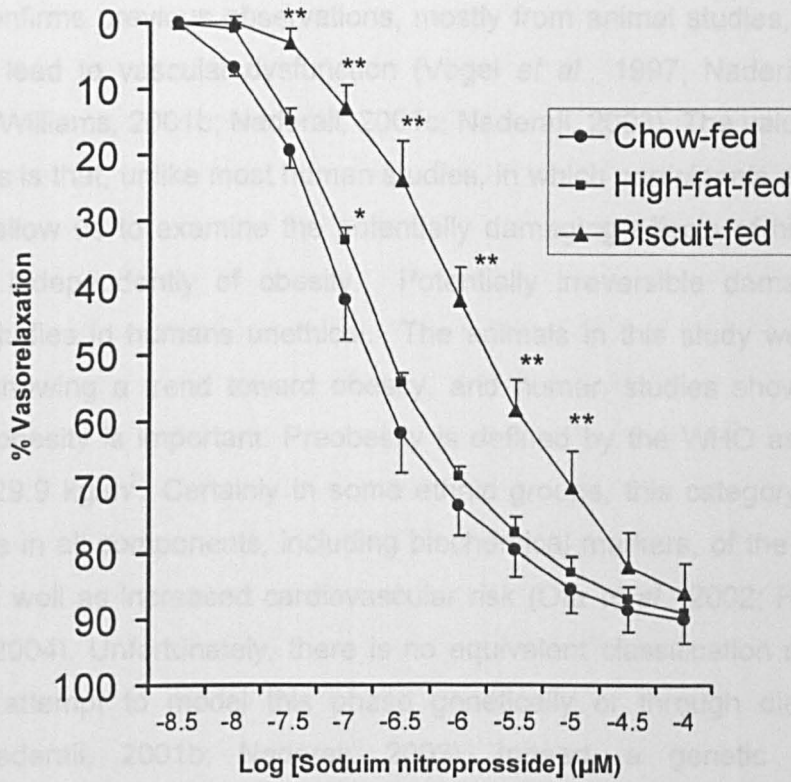


Fig. 3.16 Concentration-response curves for sodium nitroprusside (SNP) in noradrenaline (NA)-precontracted arteries from male Wistar rats fed chow, biscuit-supplemented or high-fat diet for 10 weeks, as described in Methods. Data represent mean \pm S.E.M. SNP-induced relaxation was greatly attenuated in biscuit-fed and, to a lesser extent, in high-fat-fed rats (* P <0.05 and ** p <0.01, respectively; ANOVA (Bonferroni test); n =6/group).

3.4 DISCUSSION

This study confirms previous observations, mostly from animal studies, that high-energy diets lead to vascular dysfunction (Vogel *et al.*, 1997; Naderali, 2001b; Naderali and Williams, 2001b; Naderali, 2001c; Naderali, 2003). The value of these animal studies is that, unlike most human studies, in which participants are already obese, they allow us to examine the potentially damaging effects of high-energy consumption independently of obesity. Potentially irreversible damage would make such studies in humans unethical. The animals in this study were clearly overweight, showing a trend toward obesity, and human studies show that this period of preobesity is important. Preobesity is defined by the WHO as having a BMI of 25.0-29.9 kg/m². Certainly in some ethnic groups, this category shows a significant rise in all components, including biochemical markers, of the metabolic syndrome, as well as increased cardiovascular risk (Ota *et al.*, 2002; Hop le and Xuan Ngoc, 2004). Unfortunately, there is no equivalent classification system for rodents, but attempt to model this phase genetically or through diet can be achieved (Naderali, 2001b; Naderali, 2003). Indeed, a genetic model of atherosclerosis, the JCR:LA-cp rat, prior to development of frank obesity, shows profound alterations in fat and muscle LDL activity (Mantha *et al.*, 2002). It is arguably more important, however, to demonstrate relevant changes in a model more closely resembling the multifactorial nature of most forms of human obesity, which arise on a polygenic background in response to consumption of energy-dense food. In previous studies, vascular damage has been induced prior to obesity onset with short-term feeding of high-energy diets (Naderali, 2001b; Naderali, 2003). It is logical that this would therefore also be true after chronic consumption of a high-energy diet, but this has not been demonstrated until now, as most feeding paradigms of this nature would be expected to induce frank obesity. This study shows, how intriguing it is that, depending on the particular dietary stimulus (in this case, composition), vascular damage can prevail in the absence of significant phenotypic changes reflecting obesity. Thus, one could suggest that some humans are consuming atherogenic diets for prolonged periods,

but are failing to be alerted to their increased cardiovascular risk because they remain relatively slim.

The high-fat and biscuit-supplemented diets used in this study induced similar degrees of weight gain and associated metabolic abnormalities despite having different macronutrient compositions. This suggests that these differences in composition account for the differences observed in vascular reactivity. The biscuit-supplemented diet exerted more pronounced effects on vascular reactivity overall. It heightened contractility, as shown by augmentation of both KCl- and NA-induced vasoconstriction, whereas the high-fat diet had no effect on either of these responses. It also attenuated vasorelaxation in response to the cGMP-mediated actions of both CCh and SNP, as did the high-fat diet. This means that this effect of both diets is mediated through both endothelium and vascular smooth muscle. Consumption of other high-energy diets has shown this dual mechanism (Naderali, 2001b; Naderali and Williams, 2001b; Naderali, 2003), although some such diets exert their effects only through one mechanism (see chapter four), further highlighting the specificity of effects depending on dietary components. The failure of either diet to alter cAMP-mediated vasorelaxation (i.e., histamine-induced responses) also appears to be a common feature of high-energy diets (Naderali, 2001b; Naderali, 2003).

Intriguingly, the degree of abnormality seen in both endothelium- and smooth muscle-mediated vasodilation also differed according to diet. CCh-induced vasorelaxation was more severely attenuated by the high-fat diet, whilst the SNP response more so by the biscuit-supplemented diet. These findings suggest that different dietary components affect vasorelaxation by different mechanisms, that a diet high in fat has greater deleterious effects on the arterial endothelial lining, whereas a diet relatively rich in carbohydrate and fat largely compromises the vascular smooth muscle. There is substantial supporting evidence that dietary components play a role in vascular function. Humans and animals fed diets high in fat, particularly saturated fat (Vogel *et al.*, 1997; German and Dillard, 2004; Seo *et al.*, 2005; Verhamme *et al.*, 2002), or low in protein (Koumentaki *et al.*, 2002) show endothelial abnormalities. On the other hand, the protective effects of high-

carbohydrate (low-saturated fat) diets and diets high in unsaturated fats, are contentious (Ong *et al.*, 1999; German and Dillard, 2004; Pelkman *et al.*, 2004; Keogh *et al.*, 2005; Volek and Feinman, 2005). The biscuit-supplemented diet, although lower in both total and saturated fat than the high-fat diet, contained a significantly greater amount of monounsaturated fats (MUFAs). Although these are associated with improved cardiovascular risk (Pelkman *et al.*, 2004; Appel *et al.*, 2005), at high levels they can acutely impair endothelial function (Ong *et al.*, 1999). Recent reviews of the literature note that stimulation of insulin release by increased carbohydrates promotes adipogenesis, weight gain and atherogenesis, all associated with the metabolic syndrome (German and Dillard, 2004; Suter, 2005; Volek and Feinman, 2005). The biscuit-supplemented diet was higher in overall carbohydrate content than the high-fat diet, and it may be this, in combination with relatively high fat content, that accounts for its more detrimental effects. The relatively lower saturated fat and higher carbohydrate content may also be more palatable and account for the increased daily intake in the biscuit-fed animals, and, hence, epididymal adiposity and overall greater hyperlipidaemia. This would have to be more carefully analysed in future, however, by matching intake volume and caloric content between groups. An adipocentric view would suggest that the increased adiposity *per se* of biscuit-fed animals is critically linked with the worsened vascular defects in this group, with excessive fat mass in this depot resulting in fat cell dysfunction, which in turn contributes to the metabolic disorders that increase the risk of atherosclerosis (Bays and Dujovne, 2006).

High-carbohydrate meals may also stimulate sympathetic nervous system activity *in vivo* (Marques-Lopes *et al.*, 2003), and counteract insulin-induced vasodilation. Although fasting insulin levels were similar in the three groups studied, it is possible that differences in post-prandial insulin (which would be expected to be higher in rats fed a carbohydrate-enriched diet) and triglycerides may account for the differences observed. Exactly which dietary factors are directly responsible for atherosclerosis and how, is the subject of ongoing debate and discovery, but triglycerides appear to play an important role in both humans and animals (Naderali and Williams, 2001b; German and Dillard, 2004; Barbagallo *et al.*, 2006) in that hypertriglyceridaemia is a recognized atherogenic risk factor (for review, see

German 2005). Indeed, this is even the case in the absence of insulin resistance (Naderali and Williams, 2001b) and in subjects asymptomatic for atherosclerosis (Corrado *et al.*, 2005), emphasising once again the insidious nature of the disorder.

Finally, although % energy derived from protein was lower in both biscuit-supplemented and high-fat diets compared to chow, being lowest in the former, the total average daily protein intake was similar (67 g and 78 g/day, respectively; biscuit-supplemented vs high-fat: $p>0.05$). Hence, although very low protein diets are known to be associated with vascular dysfunction (Koumentaki *et al.*, 2002), we consider it unlikely that protein deficiency explains the differences in vascular function seen in this study. Differences in antioxidant levels in the diets may also play a role (Kuno *et al.*, 1998; Sato *et al.*, 2002), but it is not possible to comment further on these as they were not measured in this study.

In conclusion, this study shows that high-energy diets of varying compositions can induce vascular dysfunction in the rat over a protracted period prior to the onset of obesity. They do this by distinct mechanisms involving different layers of the blood vessel wall, and it appears that combining atherogenic dietary components is particularly damaging, possibly through increased hyperlipidaemia. This clarifies to some extent our understanding of the nature of the vascular pathology of high-energy diets blamed for the current obesity epidemic. Clinicians now must be watchful not only of overtly obese patients, but also of those who display body weight lying persistently in the overweight category. A detailed review of dietary intake is thus important regardless of body type or individual phenotypic response to high-energy diets.

Chapter 4

Results: The effect of high-energy diet supplementation on changes in metabolic parameters and vascular reactivity in male Wistar rats.

4.1 INTRODUCTION

Acceleration in the rate of diet-induced obesity throughout the world is becoming a major health problem. Excess energy intake in the form of refined carbohydrates and fatty acids is one of the main causes of the obesity in human. Obesity is a major risk factor for cardiovascular disorders, which include hypertension, atherosclerosis and ultimately myocardial infarction (Andres, 1980; Lakka *et al.*, 2002). Arterial function and structural abnormalities which include alterations of the release of vasoactive substances from endothelial cells as well as changes in arterial wall elasticity have been seen in obese human subjects and certain animal models that reflect some aspects of human obesity (Cowan *et al.*, 1991; Dobrian *et al.*, 2000). Moreover, human and animal studies have shown various degrees of endothelial dysfunction, namely a reduction to vasorelaxants, in obesity (Steinberg *et al.*, 1996; Abram, 1997; Perticone *et al.*, 2001; Naderali and Williams, 2001b). Furthermore, diet rich in fat and refined carbohydrate induces insulin resistance (Abram, 1997; Barnard *et al.*, 1998), endothelial dysfunction, and oxidant/antioxidant imbalance (Roberts *et al.*, 2005), which may increase morbidity and premature mortality.

Increasing reports suggest that diet and its variant contents and/or components plays important roles in arterial function. Studies have shown that consumption of a specific diet of high fat or low protein causes endothelial abnormalities in humans and animals, while a low fat diet may produce a protective effect on vascular reactivity as shown in chapter three and other reports (Vogel *et al.*, 1997; Hannum and Erdman, 2000; Verhamme *et al.*, 2002; Serafini *et al.*, 2003), high fat diet have detrimental effects on vascular function (Vogel *et al.*, 1997; Naderali and Williams, 2001b; Naderali, 2003). Moreover, pharmacological lowering of plasma lipids by fenofibrate a PPAR- α agonist, retards atheroma developments (Lee *et al.*, 1995; Auwerx *et al.*, 1996), perhaps by stimulating the expression of mitochondrial enzymes involved in fatty acid catabolism as well as induction of mitochondrial and peroxisomal fatty acid β -oxidation pathways (Aoyama *et al.*, 1998). Therefore, the aim of this study was to investigate the effects of supplementation of high-fat, refined carbohydrate diet (Galaxy creamy milk chocolate supplemented) in the

absence or presence of short-term (7 days) administration of fenofibrate on the genesis of obesity, metabolic parameters, vascular functions in the rat.

4.2 METHODS

4.2.1 Experimental protocol

Adult male Wistar rats (n=21) were randomized and assigned to three groups which were fed on either a standard pelleted laboratory chow (which provided 60% of energy as carbohydrate, 30% as protein and 10% as fat) throughout (n=7; control, providing total energy content of 1310 KJ/100 g), or given chow plus chocolate (Galaxy; which provided 42.2% of energy as carbohydrate, 5.1% as protein and 52.7% as fat) (n=14; chocolate supplemented, providing total energy content of 2256 KJ/100 g). After 14 weeks, half of the chocolate supplemented group was given fenofibrate (50 mg/kg/day) and the remainder were given vehicle (1% carboxymethyl cellulose at 1 ml/kg body weight) by oral gavage daily for 7 days, before being sacrificed. All animals had free access to their respective diets and water, and were singly housed under controlled environmental conditions, and a 12-h light/dark cycle. Daily energy-intake was determined in all three groups for the last 7 days of the study. The rats were killed by CO₂ inhalation, and the epididymal and perirenal fat pads and the gastrocnemius and cardiac muscles were dissected and weighed. Blood was collected for haematocrite studies and for the measurement of glucose, insulin, leptin, non-esterified fatty acids (NEFA) and triglyceride (TG).

4.2.2 Assessment of vascular function

Six third-order mesenteric arteries were carefully dissected from each animal and mounted on two 40- μ m diameter stainless-steel wires in an automated myograph (Cambustion, Cambridge, UK). A Pair of arteries were incubated in a 5-ml organ bath containing physiological salt solution (PSS; composition [in mM]: NaCl 119, KCl 4.7, CaCl₂ 2.5, MgSO₄ 1.17, NaHCO₃ 25, KH₂PO₄ 1.18, EDTA 0.026 and glucose 5.5) gassed with 95% O₂ and 5% CO₂ at 37 °C.

After 30 min equilibration, the length-tension characteristics for each vessel were determined. Arteries were allowed a further 30 min to equilibrate before being

depolarised twice with high-potassium physiological salt solution (KPSS) in which NaCl in normal PSS was replaced by an equimolar concentration of KCl (125 mM). Cumulative concentration-response curves to either KCl (10-125 mM) or noradrenaline (NA, 0.5-6 μ M) were then carried out. Any vessel failing to reach its predetermined target tension in response to KCl (125 mM) was discarded. The effects of selected endothelium- dependent vasodilator, carbamylcholine (CCh), and endothelium-independent vasodilator, sodium nitroprusside (SNP) on arteries precontracted with 8 μ M NA, were also measured.

4.2.3 Data interpretation and statistical analyses

KCl- and NA-induced vasoconstriction responses are expressed as absolute force generated. Relaxation in response to CCh, and SNP are calculated as the percentage reduction from the maximal tension generated in response to a supramaximal concentration of NA (8 μ M).

Data are means \pm S.E.M. Statistical significance was tested using repeated-measures ANOVA or the Mann-Whitney test, as appropriate. Differences were considered statistically significant at the $p < 0.05$ levels.

4.3. RESULTS

4.3.1 Metabolic data

Supplementation of chocolate did not alter total body weight gain. In fact throughout the experiment there were no significant differences in total body weight at any given time between chow-fed control and chocolate supplemented test groups (Fig. 4.1). Furthermore, the epididymal and perirenal fat pad masses as well as gastrocnemius muscle mass were comparable in all three groups, indicating the absence of any obesity (Figs. 4.2 A, B, & C). Analysis of food intake during the last 7 days showed that compared with control chow-fed rats, total weekly energy intake were slightly increased in chocolate supplemented group, but this increased were not statistically significant (Fig. 4.3), whereas, fenofibrate-treated rats, had significantly ($p < 0.05$) higher energy intake than control chow-fed group (Fig. 4.3).

While supplementation of chocolate had no significant effect on cardiac muscle mass or haematocrit values, fenofibrate treated animals had significantly ($p<0.01$) higher cardiac muscle mass and reduced haematocrit values (Fig. 4.4 A, & B). At the end of the experiment, fasting plasma glucose, insulin levels were comparable between control chow-fed and chocolate supplemented groups, whereas, fenofibrate treated animals had significantly ($p<0.01$) lower insulin levels than their control counterparts (Figs. 4.5 A, & B). Furthermore, chocolate supplemented animals had significantly higher fasting concentrations of plasma leptin (Fig. 4.6), TG, and NEFA (Figs. 4.7 A, & B), than chow-fed controls. These raises were almost completely reversed by fenofibrate-treatment.

4.3.2 Agonist-induced contractile responses

KCl (10-125 mM)- and NA (0.5-6 μ M)-induced vasoconstriction in arteries from all three groups produced a characteristic sigmoid relationship. The contractile responses to KCl and NA in arteries from chocolate supplemented animals were significantly ($p<0.001$) increased in comparison to control groups (Figs. 4.8 & 4.9). The maximal KCl-induced contractions were 5.78 ± 0.54 control, 7.14 ± 0.57 chocolate supplemented, and 7.07 ± 0.52 mN/mm artery fenofibrate-treated, respectively (Fig. 4.8). A similar picture was also seen with NA-induced vasoconstriction giving maximal contraction of 7.68 ± 1.02 control, 10.91 ± 0.96 supplemented, and 10.34 ± 0.66 mN/mm artery fenofibrate-treated, respectively (Fig. 4.9).

4.3.3 Endothelium-dependent and -independent vasorelaxation

NA-precontracted arteries from chow-fed rats demonstrated progressive relaxation to cumulative addition of CCh (10 nM-100 μ M), achieving a maximum of $87 \pm 2\%$ at a CCh concentration of 100 μ M. Arteries from test and fenofibrate-treated groups that were similarly exposed to CCh displayed similar sigmoid responses with maximal relaxation of $72 \pm 6\%$ and $92 \pm 2\%$ (Fig. 4.10). However, arteries from chocolate supplemented rats had significantly ($p<0.01$) higher EC_{50} than control chow-fed animals (EC_{50} : 1.86 ± 0.02 chocolate supplemented, vs EC_{50} : 0.28 ± 0.01 μ M Chow-fed). Fenofibrate-treatment significantly ($p<0.05$) improved

EC₅₀ compared with chocolate supplemented group (EC₅₀: 0.68 ± 0.01 μM fenofibrate-treatment vs EC₅₀: 1.86 ± 0.02 μM chocolate supplemented).

Endothelium-independent vasodilator (SNP; 10 nM-100 μM) also induced concentration dependent vasorelaxation of arteries from all three groups with no significant differences in arterial vasorelaxation between the three groups achieving a maximal relaxation of 85 ± 3 control, 82 ± 2 chocolate supplemented, and 84 ± 3% fenofibrate-treated groups, respectively (Fig. 4.11).

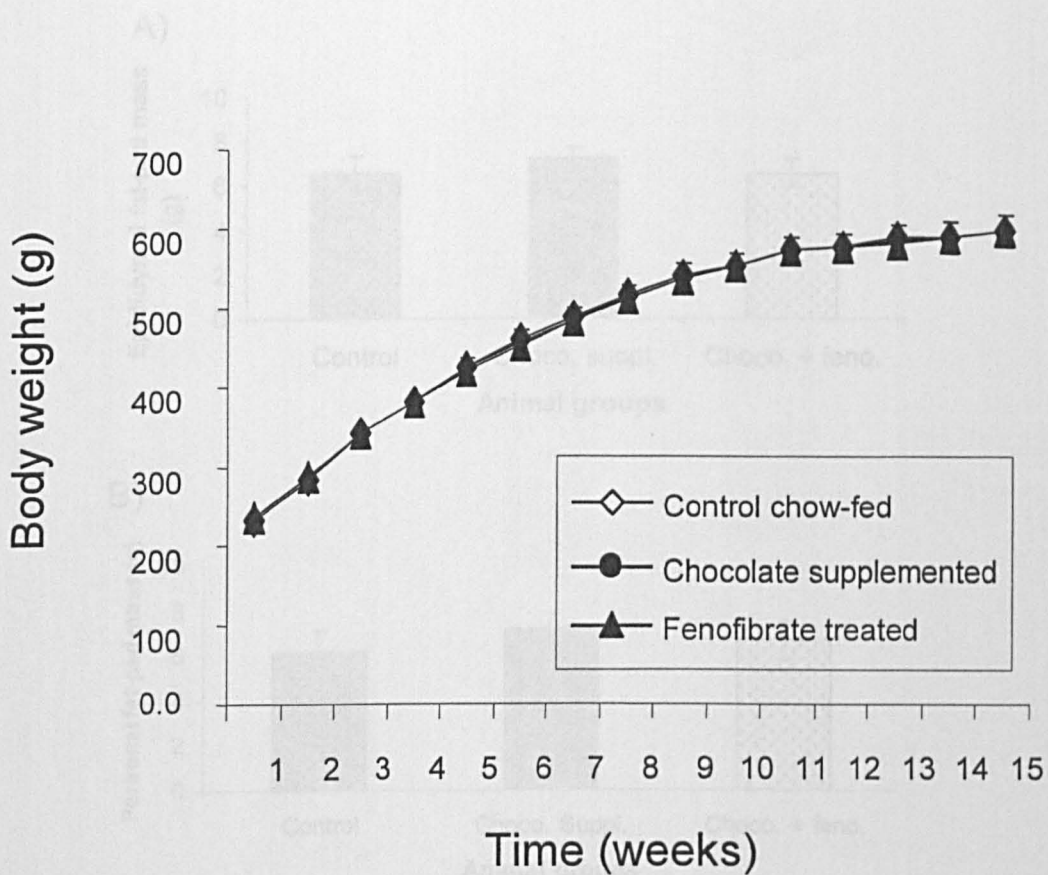


Figure 4.1. The effect of chocolate supplementation and fenofibrate treatment on total body weight in the male wistar rats. Control animals were fed standard chow, while chocolate supplemented groups was given chow plus chocolate (galaxy) for 15 weeks. Fenofibrate-treated rats were given fenofibrate (50 mg/kg/day) daily for 7 days. At the end of 15 week, there were no significant changes in total body weight between the three groups. In fact weight increase in all three groups were superimposable. Data are expressed as mean \pm S.E.M.

Fig. 4.2. Effect of high energy diet (chocolate supplemented) and fenofibrate treatment on A) epididymal fat-pad, B) perirenal fat-pad mass, and C) gastrocnemius muscle mass. Data are presented as mean \pm S.E.M. No significant differences were seen between all three groups. Choco. suppl. = chocolate supplemented, and Choco. + fenof. = Chocolate supplemented + Fenofibrate.

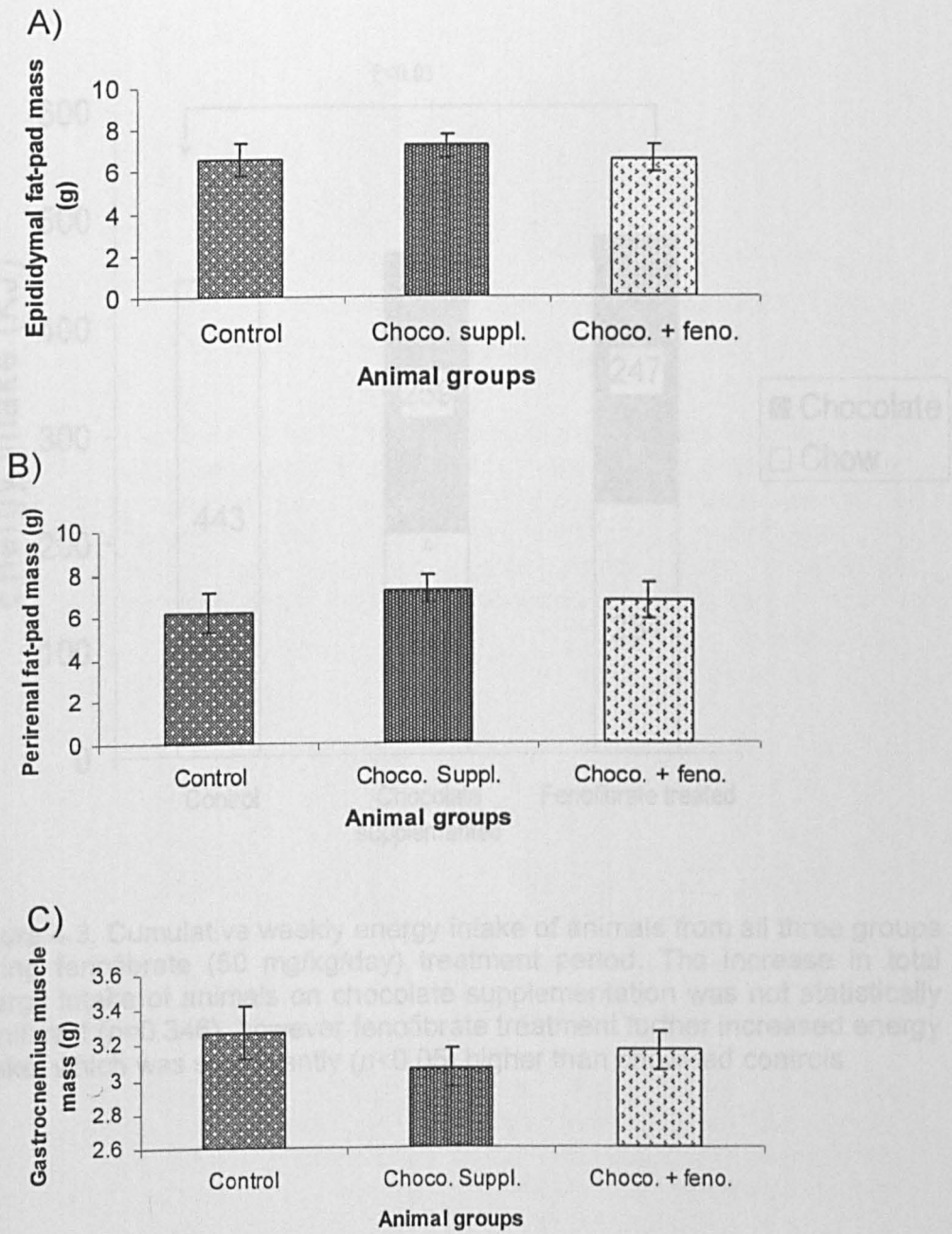


Fig. 4.2. Effect of high energy diet (chocolate supplemented) and fenofibrate treatment on A) epididymal fat-pad, B) perirenal fat-pad mass, and C) gastrocnemius muscle mass. Data are presented as mean \pm S.E.M. No significant differences were seen between all three groups. Choco. suppl. = chocolate supplemented, and Choco. + feno. = Chocolate supplemented + Fenofibrate.

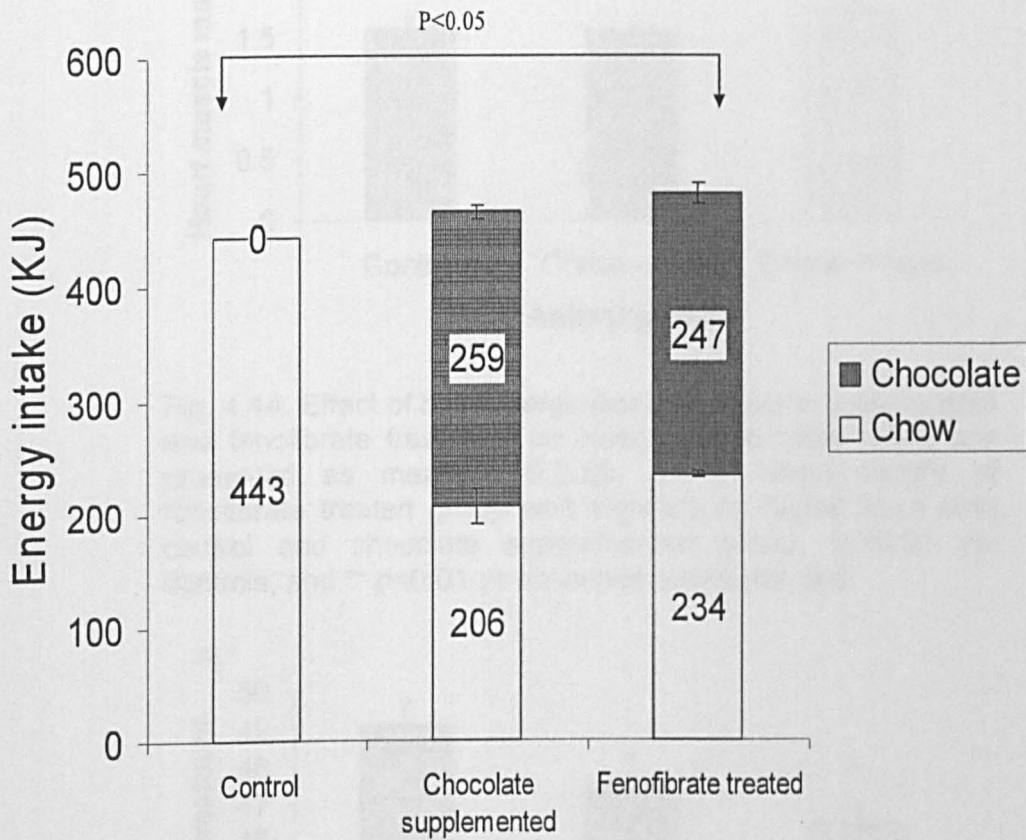


Figure 4.3. Cumulative weekly energy intake of animals from all three groups during fenofibrate (50 mg/kg/day) treatment period. The increase in total energy intake of animals on chocolate supplementation was not statistically significant ($p=0.346$), however fenofibrate treatment further increased energy intake, which was significantly ($p<0.05$) higher than chow-fed controls.

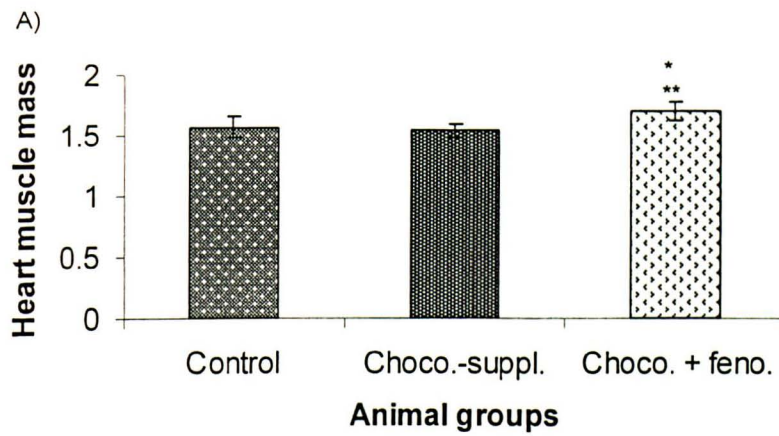


Fig. 4.4A. Effect of high energy diet (chocolate supplemented) and fenofibrate treatment on heart muscle mass. Data are presented as mean \pm S.E.M., (n=7). Heart weight of fenofibrate treated group was significantly higher than both control and chocolate supplemented group. * $p < 0.01$ vs. controls, and ** $p < 0.01$ vs chocolate supplemented.

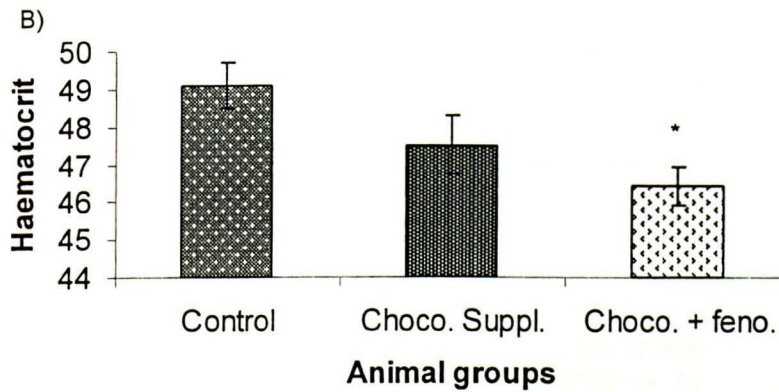


Fig. 4.4B. Effect of high energy diet (chocolate supplemented) and fenofibrate treatment on haematocrit. Data are presented as mean \pm S.E.M. (n=7). Fenofibrate treated group had significantly lower haematocrit than chow-fed control. * $p < 0.01$ vs control.

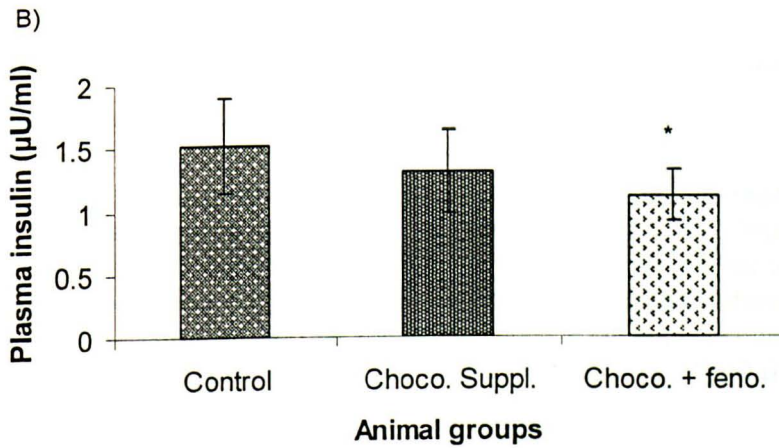
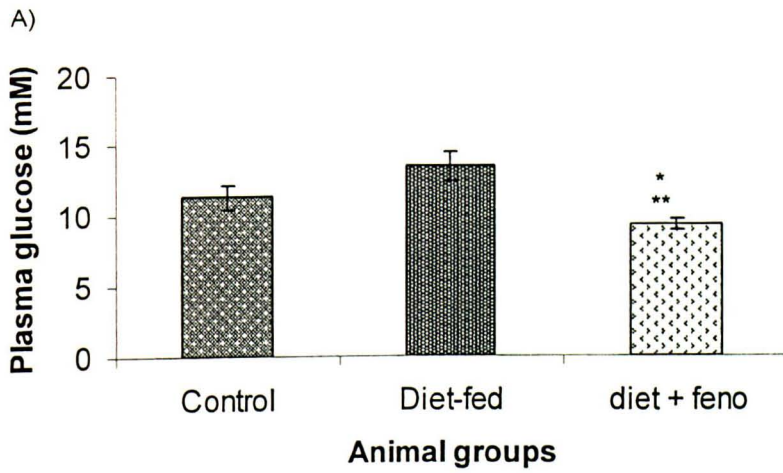


Fig. 4.5. The effects of high energy diet (chocolate supplementation) and fenofibrate-treatment on plasma A) glucose, B) insulin levels. Fenofibrate-treated animals had significantly lower plasma glucose and insulin than their chow-fed counterparts. Data are presented as means \pm S.E.M. * $p < 0.01$ vs control, ** $p < 0.01$ vs chocolate supplementation.

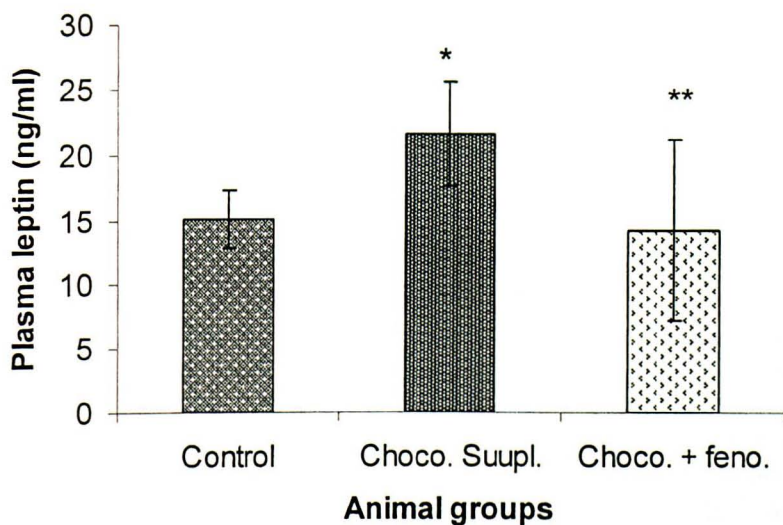


Fig. 4.6. Effect of high energy diet (chocolate supplemented) and fenofibrate treatment on plasma leptin. The level of chocolate supplemented group had significantly higher leptin than control, while there was no significant difference between control and fenofibrate groups. Data are presented as mean \pm S.E.M. * $p < 0.01$ vs controls, and ** $p < 0.01$ vs chocolate supplemented.

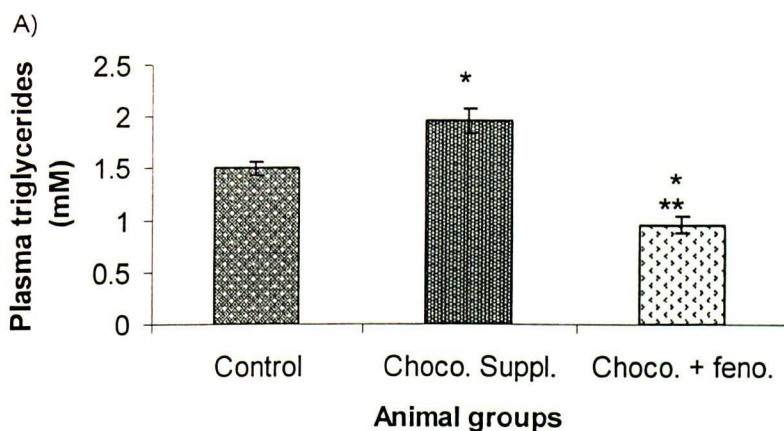


Fig. 4.7A. Effect of high energy diet (chocolate supplemented) and fenofibrate treatment on plasma triglycerides. The level of chocolate supplemented group was significantly higher than control, while fenofibrate treated group was significantly lower than both control and chocolate supplemented groups. Data are presented as mean \pm S.E.M. * p <0.01 vs. controls, and ** p <0.01 vs chocolate supplemented.

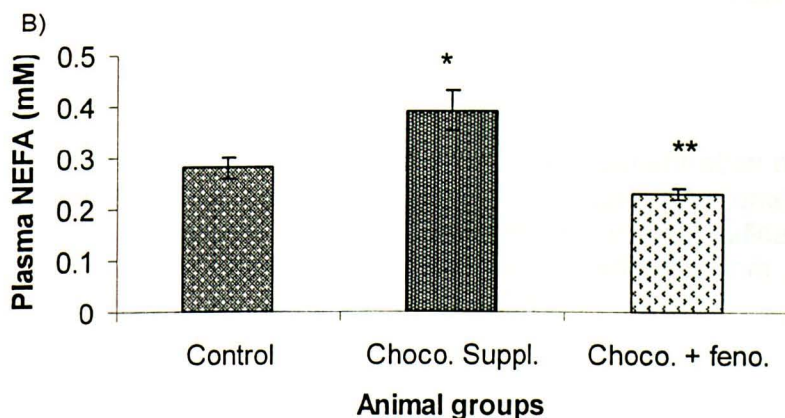


Fig. 4.7B. Effect of high energy diet (chocolate supplemented) and fenofibrate treatment on plasma NEFA. The level of chocolate supplemented group was significantly higher than control, while there was no significant difference between control and fenofibrate groups. Data are presented as mean \pm S.E.M. * p <0.01 vs controls, and ** p <0.01 vs chocolate supplemented.

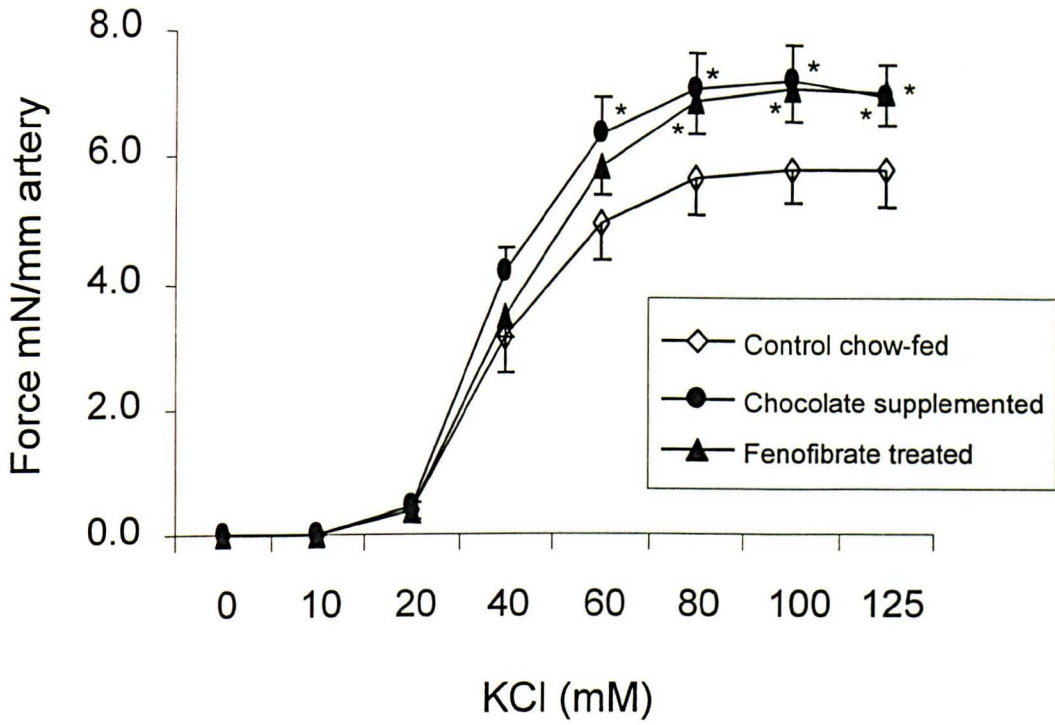


Figure 4.8. Vasoconstriction responses to cumulative concentration of KCl (10-125 mM) in arteries from chow-fed control, chocolate supplemented and fenofibrate-treated animals. KCl-induced contractions were significantly (n=7; ANOVA (Bonferroni test), * $p < 0.001$) augmented in arteries from chocolate supplemented animals. Data represent mean \pm S.E.M.

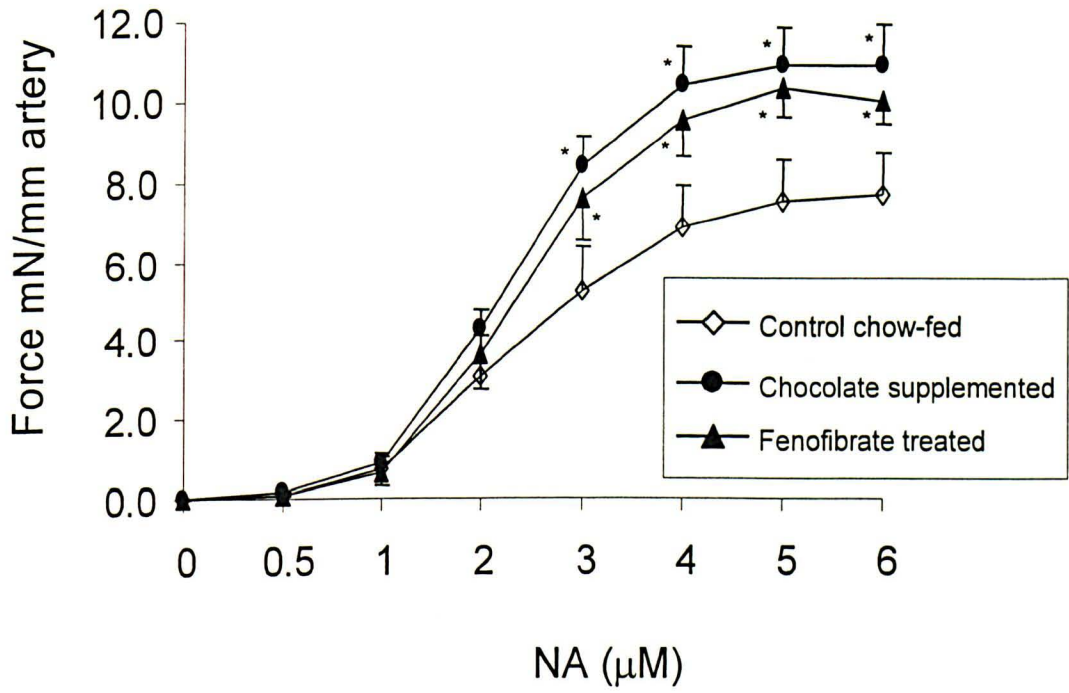


Figure 4.9. Vasoconstriction responses to cumulative concentration of noradrenaline (NA; 0.5-6 µM) in arteries from chow-fed control, chocolate supplemented and fenofibrate-treated animals. NA-induced contractions were significantly (n=7; ANOVA (Bonferroni test), * $p < 0.01$) augmented in arteries from chocolate supplemented animals. Data represent mean \pm S.E.M.

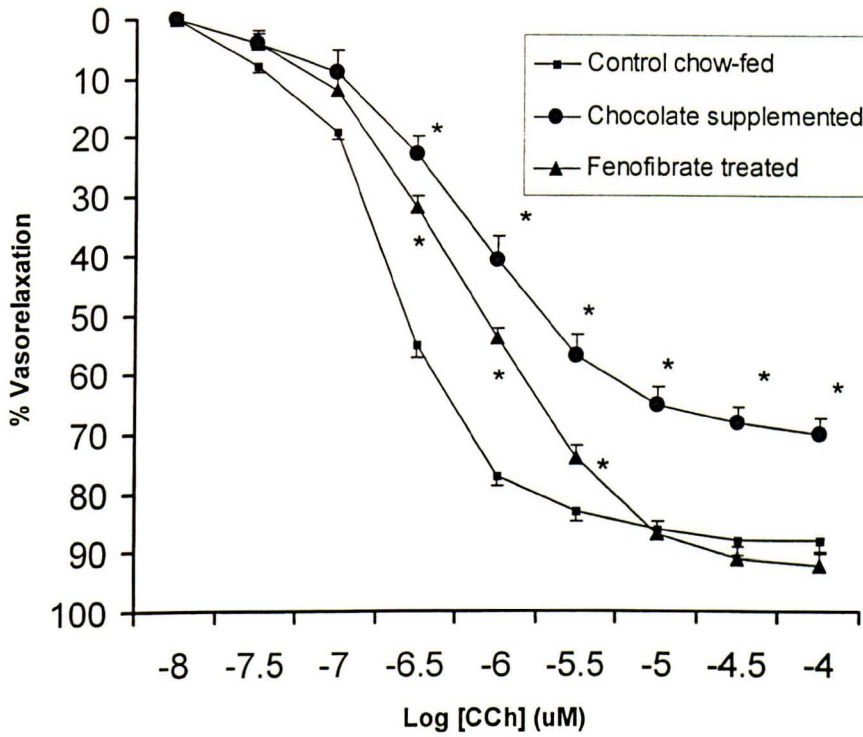


Figure 4.10. The effects of carbamylcholine on NA-precontracted ($8\mu\text{M}$) arteries from three groups of rats. When contraction reached a plateau after 2 minutes, concentration-response curves to carbamylcholine (CCh) were carried out. Data represent mean \pm S.E.M. The concentration-response curves to CCh, were significantly reduced in chocolate supplemented group ($n=7$; ANOVA (Bonferroni test), $p<0.01$), while fenofibrate-treatment significantly ($p<0.01$) improved vasorelaxation in chocolate supplemented animals.

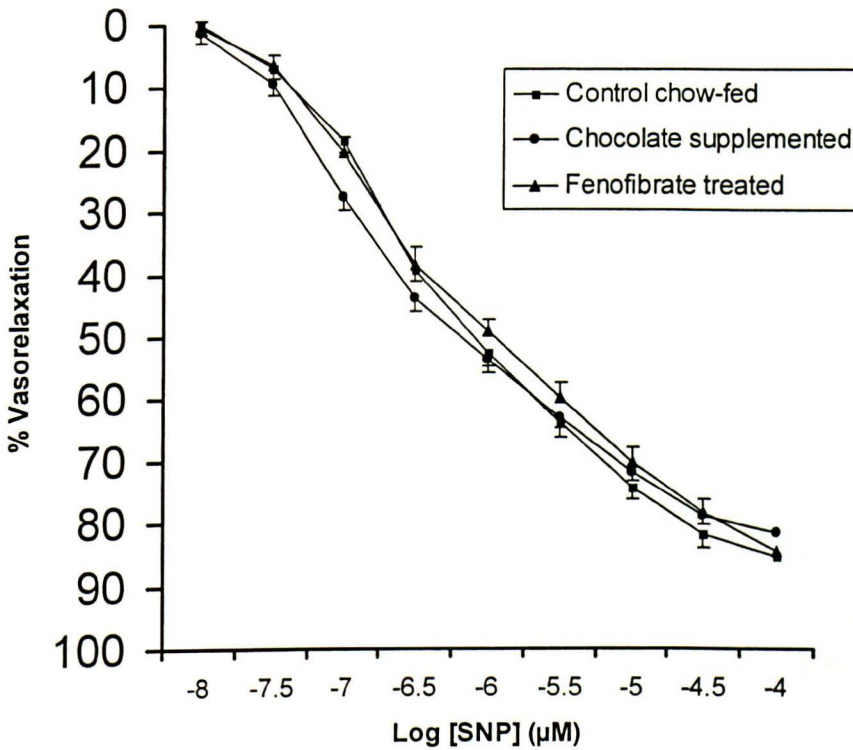


Figure 4.11. The effects of sodium nitroprusside (SNP) on NA-precontracted (8µM) arteries from three groups of rats. When contraction reached a plateau after 2 minutes, concentration-response curves to SNP were carried out. Data represent mean \pm S.E.M. The concentration-response curves to SNP, were not significantly different in both chocolate supplemented and fenofibrate-treatment groups, when compared with control chow-fed.

4.4 DISCUSSION

Excessive availability of high-energy diet in many industrialised societies, is producing a health hazard, namely obesity and obesity related secondary disorders, which ultimately leads to the deterioration of quality of life, and in many cases to premature and sudden death. Numerous experimental studies have reported severe detrimental outcome of excessive high-energy diets on well being. For example, high-fat diet impairs endothelium-dependent vasorelaxation (Steinberg *et al.*, 1996; Vogel *et al.*, 1997; Naderali and Williams, 2001b), while removal of high-energy diet provides some protection against obesity-induced metabolic and cardiovascular abnormalities (Sasaki *et al.*, 2002; Verhamme *et al.*, 2002). In hypertensive obese patients, ACh-induced forearm blood flow (FBF) is significantly lower than healthy individuals. Caloric restriction decreases body weight and the mean blood pressure, fasting plasma insulin, total cholesterol, triglycerides, low-density lipoproteins (LDL) as well as enhancing ACh-induced FBF (Sasaki *et al.*, 2002). Moreover, animal studies have shown that withdrawal of high-fat, high-palatable diet significantly reduces plasma lipid profile and markedly improves vascular reactivity to acetylcholine (Verhamme *et al.*, 2002). Furthermore, Pharmacological lowering of circulating plasma lipids has also been shown to reduce the risk for coronary events, cardiovascular morbidity and mortality in a broad range of patients (Brown *et al.*, 1993 ; Bucher *et al.*, 1999). In this study, supplemental feeding of animals with chocolate did not resulted in obesity (characterised with an excessive increase in total body weight and fat pad mass), however, there were significant metabolic abnormalities, namely higher plasma lipids of TG and NEFA. Higher plasma lipids seen in this study were similar to those achieved by chronic consumption of high-fat, high-energy diets and obesity, as shown in chapter three, and other reports (Steinberg *et al.*, 1996; Perticone *et al.*, 2001; Naderali *et al.*, 2001c; Verhamme *et al.*, 2002), indicating that adipose accumulation is not a pre-requisite for abnormal lipid profile (Vogel *et al.*, 1997; Naderali and Williams, 2003). Fenofibrate treatment in the presence of chocolate diet completely reversed lipid profile, indicating beneficial effects of fenofibrate in correcting abnormal lipid profile. Moreover, treatment of animals with fenofibrate corrected all metabolic changes associated with diet and obesity.

Examination of contractile and relaxation properties of arteries showed striking differences in responses to various stimuli between the three groups. In contrast to the results reported in chapter three and previous studies (Walker *et al.*, 1997; Fontes *et al.*, 1998; Naderali and Williams, 2001b), contractile responses to KCl and NA were significantly augmented in chocolate supplemented diet groups (with or without fenofibrate administration). Although from this study it is not possible to contemplate the mechanism of hyperreactivity to KCl and NA, nonetheless, possibilities are that a) long-term consumption of chocolate may have a direct effect on the activity of the contractile apparatus, or b) it may have increased/eased the availability of the calcium for contraction required by KCl and NA. The latter hypothesis is more plausible as chocolate has been shown to markedly increase calciuria (Nguyen *et al.*, 1994). However, these hypothesis merits further investigation.

Results in chapter three and other numerous studies have shown that excessive high-energy diet impairs endothelium-dependent arterial function (Brown *et al.*, 1993 ; Walker *et al.*, 1997; Fontes *et al.*, 1998; Bucher *et al.*, 1999; Naderali *et al.*, 2001c; Sasaki *et al.*, 2002). Moreover, defect in vasorelaxation can also be seen with short-term consumption of high-energy diet in the absence of any obesity (Vogel *et al.*, 1997; Naderali and Williams, 2001b; Naderali and Williams, 2003). In agreement with previous reports, in this study there was a marked attenuation of endothelial mediated CCh-induced vasorelaxation in the absence of any obesity, suggesting a significant defect in either CCh-induced NO production via eNOS activity (Boulanger, 1999) or the release of endothelium-derived hyperpolarizing factor (EDHF) (Gerber *et al.*, 1998). On the other hand, vasorelaxant effect of SNP is due to its direct stimulatory action on cGMP and thereby induction of NO generation (Azula *et al.*, 1996) in vascular smooth muscle cells. In present study, chocolate supplementation did not alter SNP-induced vasorelaxation, suggesting that there were no changes in arterial smooth muscle activity.

A number of various hypotheses have been put forward as the mechanism(s) of action of high-energy diet and obesity related vascular dysfunction, which include a rise in plasma NEFA (Steinberg *et al.*, 1997) hypertriglyceridaemia (Tka *et al.*,

1997; Lewis *et al.*, 1999; Naderali *et al.*, 2001c). This rise in circulating plasma lipids has been, at least partly, attributed to inhibition of eNOS activity and thereby reduction in NO production (Liao *et al.*, 1995), ultimately resulting in a reduced CCh-induced vasorelaxation. Raised NEFA and TG levels in chocolate supplemented animals seen in this study, argue for the role of plasma NEFA and TG as the possible agents affecting vasorelaxation. However, previous reports have failed to show any relationship with raised plasma lipids and increased vasoconstriction (Fontes *et al.*, 1998; Naderali and Williams, 2001b; Naderali *et al.*, 2001c; Naderali and Williams, 2003), therefore, it is unlikely that augmented responses seen to KCl and NA in this study, might have been as a result of changes in circulating plasma lipids. In this study, despite no obesity or weight gain plasma leptin levels were significantly augmented in chocolate supplemented group. Moreover, Leptin has been shown to induce oxidative stress in human endothelial cells (Bouloumie *et al.*, 1998), resulting in activation of atherogenic process, which may contribute to the development of vascular pathology. In addition, leptin has been shown to induce vasoconstriction (Mark *et al.*, 1999 ; Correia *et al.*, 2001; Hall *et al.*, 2001) and exacerbate arterial contractility (Shek *et al.*, 1998). Therefore, raised leptin levels might participate an increase in contractility of arteries in response to NA and KCl.

In this study, although fenofibrate had very little or no effects on KCl- and NA-induced vasoconstriction, but improved CCh-induced responses, indicating improvement of endothelial function in fenofibrate-treated animals. The mechanism of action of fenofibrate on vascular function is not fully known. However, it is possible that reduction in circulating TG may play a role in fenofibrate-induced improvement of endothelial function.

Although, the component of excess energy content of high-fat, high-carbohydrate diet well known to increase weight gain and cause obesity (Matsui *et al.*, 2005), in the present study this component did not induce weight gain or obesity, and this could be related to the action of elevated plasma leptin levels, which may increase thermogenesis and suppress appetite (Pelleycounter *et al.*, 1995), leading to reduced body fat mass (Schwartz *et al.*, 2000).

Overall, this study indicates that consumption of high-energy diet (high-fat, refined carbohydrate diet, milk chocolate supplemented) attenuates arterial relaxation in the absence of any obesity in the rat. The arterial dysfunction may be due to an induction of hyperlipidaemia in general or hypertriglyceridaemia in particular. Lowering circulating plasma NEFA and/or TG by fenofibrate improved endothelial function but not increased vasoconstriction. Furthermore, increases in contractile responses of arteries from chocolate supplemented groups may not be due to the changes in circulating lipid profile

Chapter 5

Results: Palatable diet-induced obesity: changes in metabolic and vascular functions.

5.1 INTRODUCTION

Throughout the world, consumption of high-energy diets together with a general reduction in levels of physical activity causing an alarming health problem, namely obesity. Obesity is a major cardiovascular risk factor and the prevalence of atherosclerotic disease in obese human has been recognised for more than two decades (Andres, 1980; Barret-Conner, 1985). An association has been described between obesity, arterial hypertension, insulin resistance and dyslipidaemia, which comprise core features of the 'metabolic syndrome' or Syndrome X (Reaven, 1988). Many obese patients accomplish weight loss with diet, exercise and lifestyle modification, achieving substantial reductions in central obesity and associated disturbances corresponding to a significant decrease in cardiac events and mortality (Singh *et al.*, 1996; Jen *et al.*, 2002; Gutin *et al.*, 2002).

Obesity induces abnormalities of arterial function and structure in human and in certain animal models that reflect some aspects of human obesity (Cowan *et al.*, 1991; Singh *et al.*, 1996; Naderali *et al.*, 2001; Sasaki *et al.*, 2002). These include alterations of the release of vasoactive substances from endothelial cells as well as changes in arterial wall elasticity. A low-caloric diet improves (Sasaki *et al.*, 2002), while elevating plasma lipid profile further deteriorate endothelium-dependent vasodilatation in obese subjects (Steinberg *et al.*, 1997), and in patients with coronary artery disease (Zhao *et al.*, 2001). It has been shown that, acute consumption of a palatable diet causes endothelial dysfunction long before development of any significant obesity (Naderali and Williams, 2001b). Moreover, endothelial function was only partially corrected with short-term withdrawal of the palatable diet (Naderali and Williams, 2003). Thus, the aims of this study were to a) investigate the effects of long term highly-palatable diet on the genesis of obesity, metabolic abnormalities, and vascular dysfunction, b) to determine whether lifestyle modification (chronic withdrawal of highly-palatable diet) can improve the palatable diet-induced vascular and metabolic abnormalities.

5.2 METHODS

5.2.1 Experimental protocol

Adult male Wistar rats (n=24) were randomised and assigned to a control group (n=8) and a test group (n=16). All animals had free access to water and were housed individually under controlled environmental conditions (19-22 °C; 30-40% humidity) and a 12-hour light/dark cycle (lights on at 07:00h). Controls were fed a standard laboratory pelleted diet (chow-fed), while test group had free access to a highly-palatable diet as described in chapter 2, section 2.1.1. Chow-fed controls remained on their prospective diet for 15 weeks, while after 7 weeks, palatable-diet fed animals were subdivided into two subgroups. Palatable diet was removed from one group and the standard chow diet was reintroduced (diet-to-chow) for further 8 weeks, while the remaining subgroup was maintained on the palatable diet (diet-fed) for another 8 weeks. After 15 weeks from the start of the study, rats were killed by CO₂ inhalation after 2 hours of fasting, and the total body fat mass was measured by Bioimpedence Method using the TOBEC® equipment (Biotech Instrument Ltd.).

5.2.2 Biochemical analysis

Blood was withdrawn by cardiac puncture and removed into cold heparinised tubes and the gonadal and perirenal fat pads, as well as the gastrocnemius muscles, heart, liver, and lung were dissected and weighed. The plasma was immediately separated by centrifugation before being frozen for later measurements of blood analytes (glucose, insulin, leptin, TNF- α , CRP, nonesterified free fatty acids (NEFA) and triglycerides (TG)). Plasma glucose concentration was determined using a glucose oxidase method, and NEFA and TG concentrations were measured using commercial diagnostic kits (Boehringer Mannheim, and Sigma Diagnostics.). insulin and leptin concentrations were measured by radioimmunoassay (RIA) kits, While, TNF- α and CRP levels were measured using Enzyme-immunoassay kits.

5.2.3 Assessment of vascular contractility

Six third-order mesenteric arteries (>200 µm diameter, 2 mm lengths) were carefully dissected from each animal. Each artery was freed of fat and connective tissue and mounted on two 40-µm diameter stainless-steel wires in an automated myograph (Cambustion Ltd). The vessels (in duplicate) were incubated in a 5-ml organ bath containing physiological salt solution. Cumulative concentration-response curves to either KCl (10-125 mM) or noradrenaline (NA, 0.5-6 µM) were then carried out. Any vessel failing to reach its predetermined target tension in response to vasoconstriction with KCl (125 mM) was discarded.

5.2.4 Assessment of vascular relaxation

To investigate defects in endothelial-dependent and –independent vasorelaxation, relaxation to endothelium-dependent (carbamylcholine; CCh) or endothelium-independent (sodium nitroprusside; SNP) vasodilators were measured in NA-precontracted arteries. To eliminate interference of vasoactive prostanoids, 10 mM indomethacine was added to the organ bath prior to contraction with NA (8 µM). When contraction reached a plateau after 2 minutes, concentration-response curves were carried out to either CCh or SNP (for both, 10 nM-100 µM).

5.2.5 Homeostasis model assessment (HOMA)

HOMA, an index of insulin resistance, which employs measures of fasting plasma concentrations of glucose and insulin, was calculated according to the method described previously (Matthews *et al.*, 1985).

5.2.6 Data interpretation and statistical analysis

Vasoconstriction in response to NA and KCl were expressed as absolute force generated. Relaxation in response to CCh and SNP was calculated as the percentage reduction from the maximal tension generated in response to the supramaximal concentration of NA (8 µM). Data are expressed as mean ± SD or where stated as SEM for more clarity. An average response for all the vessels from a given animal was determined before group analysis. Data have normal distribution (Shapiro Wilk W test). Statistical significance was tested using Student

't-test' or repeated-measures (ANOVA; Bonferroni t-test) or the Mann-Whitney test, as appropriate. Results were considered statistically significant at the $p < 0.05$ levels. Univariate correlation between maximal CCh-induced vasorelaxation and fasting terminal insulin, leptin, TNF- α , CRP, and TG levels were determined by linear regression followed by two-tailed analyses, using ARCUS PRO-STAT (version 3.23).

5.3 RESULTS

5.3.1 Metabolic data

Animals given palatable diet progressively gained more weight than their chow-fed counterparts. A significant difference in total body weight was observed after 5th week of feeding and further increased after 8th week where diet-fed animals had significantly ($>10\%$, $p=0.0059$) higher total body weight than chow-fed controls (Fig. 5.1). Following the next 7 weeks, average weekly weight gain of diet-fed animals was 1.8-fold greater than chow-fed controls (18.43 ± 0.50 vs 10.34 ± 0.23 g control), while average weekly weight gain of diet-to-chow animals was significantly lower ($p < 0.001$, 10.04 ± 0.06 vs 18.43 ± 0.50 g diet-fed) than their diet-fed counterparts (Fig. 5.1). However, at the end of the experiment, total body weight of diet-to-chow animals was still significantly ($p=0.043$) higher than control chow-fed but was markedly ($p < 0.006$) lower than the diet-fed animals (Fig. 5.1).

At the end of the experiment, Bioimpedance measurements showed that diet-fed animals had 70 and 27% more total body fat than chow-fed and diet-to-chow groups, respectively (Figs. 5.2 C & D). Diet-fed animals had significantly higher epididymal and perirenal fat pad mass than both chow-fed (by $>190\%$, $p < 0.0001$) and diet-to-chow (by $>165\%$, $p < 0.0002$) groups (Fig. 5.2 C). Diet-fed rats had significantly (for both, $p < 0.023$) higher heart weights than both chow-fed and diet-to-chow rats (Fig. 5.2 A). There were no significant differences between fat pad depots and heart weights of diet-to-chow and chow-fed groups, respectively; nor there were any significant differences in gastrocnemius muscle weights between the three groups (Fig. 5.2 B). However, diet-fed animals had significantly heavier (122% , $p < 0.01$) liver weight than their chow-fed counterparts. Furthermore, liver weights of diet-to-chow animals showed marked reduction (18% , $p < 0.01$)

compared to diet-fed animals with no difference with that of chow-fed animals (Fig. 5.3 A). Lung weights were also comparable in all three groups with no significant difference between them (Fig. 5.3 B).

While fasting plasma glucose levels were comparable in all three groups (12.42 ± 1.12 chow-fed; 14.40 ± 0.362 diet-fed; 14.43 ± 0.42 mM diet-to-chow) (Fig. 5.4 A), diet-fed rats had significantly higher fasting terminal levels of plasma insulin than both chow-fed and diet-to chow group (2.83 ± 0.18 diet-fed vs 1.22 ± 0.18 chow-fed and 1.22 ± 0.22 mM diet-to-chow; for all $p < 0.009$) (Fig. 5.4 B). As a result, diet-fed rats had significantly (for both, $p < 0.001$) higher HOMA index, indicating development of insulin resistance in these animals. Moreover, diet-fed rats had significantly higher plasma leptin levels than both chow-fed control and diet-to-chow groups (for both $p < 0.001$; 9.35 ± 0.25 diet-fed vs 7.53 ± 0.52 chow-fed and 6.90 ± 0.24 mM diet-to chow) (Fig. 5.4 C).

Fasting plasma levels of NEFA were comparable in all these groups (0.45 ± 0.24 chow-fed; 0.48 ± 0.41 diet fed and 0.40 ± 0.24 mM diet-to-chow), while diet-fed rats had significantly (>2 -fold, $p < 0.001$ for both) higher levels of fasting TG than both chow-fed and diet-to chow groups (Fig. 5.5 A & B). Compared with chow-fed group, plasma levels of TNF- α in both diet-fed and diet-to-chow groups were significantly elevated (72% and 66% respectively, $p < 0.05$ for both). There was no significant difference between diet-fed and diet-to-chow groups. Similarly, plasma levels of CRP in diet-fed animals were significantly higher than chow-fed group (> 3 fold, $p < 0.05$), an effect which was not seen in diet-to-chow group (Fig. 5.6 A & B).

5.3.2 Vascular data

There were no significant differences in vessel diameter between the three groups (189 ± 20 chow-fed; 200 ± 15 diet-fed and 200 ± 10 μ m diet-to-chow; $p > 0.5$ for all).

5.3.3 Agonist-induced vasoconstriction responses

Cumulative concentration response curves (CCRC) to KCl (10-125 mM) and NA (0.5-6 μ M) in arteries from all three groups produced characteristic sigmoid

relationship. KCl-induced contractions were similar in all three groups such that CCRC were virtually superimposable, giving maximum contractions of 5.90 ± 0.41 chow-fed, 6.16 ± 0.31 diet-fed and 6.32 ± 0.26 mN/mm artery for diet-to-chow groups (Fig. 5.7), with no significant differences between the groups, either overall (by ANOVA) or at any given KCl concentration. However, arteries from diet-fed group had significantly ($>30\%$; $p<0.01$ for all) greater contractile responses to 3 and 4 μM NA than chow-fed and diet-to-chow groups. This difference was not seen at lower NA concentrations and was abolished at higher NA concentrations (Fig. 5.8).

5.3.4 Endothelium-dependent and -independent vasorelaxation

NA-precontracted arteries from chow-fed rats demonstrated progressive relaxation to cumulative addition of CCh (10 nM- 100 μM), achieving a maximum of $90 \pm 2\%$ at a CCh concentration of 100 μM , with EC_{50} values of 0.14 ± 0.02 μM . Arteries from diet fed rats that were similarly exposed to CCh displayed a significant rightward shift of CCRC compared with chow-fed control arteries (EC_{50} diet-fed: 0.75 ± 0.03 , $p<0.002$ vs chow-fed) (Fig. 5.9). Maximal vasorelaxation to 100 μM CCh was reduced by 23% in diet-fed rats ($p<0.004$ vs chow-fed). However, CCh-induced vasorelaxation curves in arteries from diet-to-chow group were virtually identical to those in chow-fed group, achieving maximal relaxation of $95 \pm 1\%$ at 100 μM CCh, with EC_{50} values of 0.19 ± 0.04 μM .

SNP (10 nM-100 μM) also induced concentration dependent vasorelaxation of arteries from all three groups. Concentration-relaxation response curves were similar and superimposable in all three groups (Fig. 5.10), with no statistically significant differences between the groups, either overall (by ANOVA) or at any given SNP concentration.

5.3.5 Metabolic determinants of vascular dysfunction

Of the various blood analytes measured, insulin, leptin, TNF- α , CRP, and TG showed significant increases in diet-fed animals. Therefore, the relationship between endothelial dysfunction (measured as the percentage reduction of 100 μ M CCh-induced vasorelaxation) and terminal plasma levels of insulin, leptin, TNF- α , CRP, and TG were examined. In each case, univariate correlation analysis showed an inverse correlation between fasting terminal levels of TG ($r^2 = 0.4787$; $p = 0.0013$), insulin ($r^2 = 0.4237$; $p = 0.0023$) and leptin ($r^2 = 0.609$; $p = 0.0001$) and maximal CCh-induced vasorelaxation.

The relationship of adipokines levels TNF- α and CRP with maximum CCh-induced vasorelaxation was also measured, and no significant correlations were found for both adipokines ($r^2 = 0.0108$; $p \geq 0.05$, and $r^2 = 0.1402$; $p \geq 0.05$, respectively). We also analysed the relationship between total body weight and total fat pad mass with vascular reactivity. Univariate correlation analysis once again indicated an inverse correlation relationship between total body weight ($r^2 = 0.3751$; $p = 0.0043$) and fat pad mass ($r^2 = 0.3565$; $p = 0.0048$) and maximum carbamylcholine-induced vasorelaxation. Furthermore, analysis of blood analytes showed significant positive correlations between fasting plasma insulin ($r^2 = 0.3043$; $p = 0.0025$) and leptin ($r^2 = 0.4189$; $p = 0.0020$) with that of circulating triglycerides.

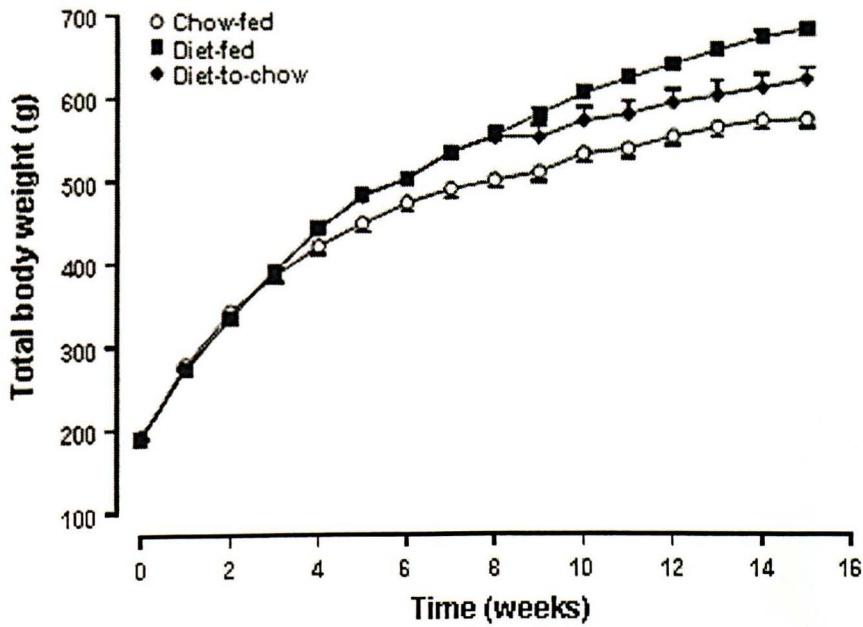


Fig. 5.1. The effect of a highly palatable-diet and its removal on total body weight in Wistar rats. Animals were fed either standard chow (chow-fed) or a palatable diet (diet-fed) for 15 weeks, or fed the palatable diet for 8 weeks and then returned to standard chow for further 7 weeks (diet-to-chow). Diet-fed animals were significantly ($p < 0.00001$) heavier than both chow-fed and diet-to-chow groups. Diet-to-chow group was significantly heavier ($p < 0.04$) than chow-fed but lighter ($p < 0.006$) than diet-fed groups. Data are expressed as mean \pm S.E.M.

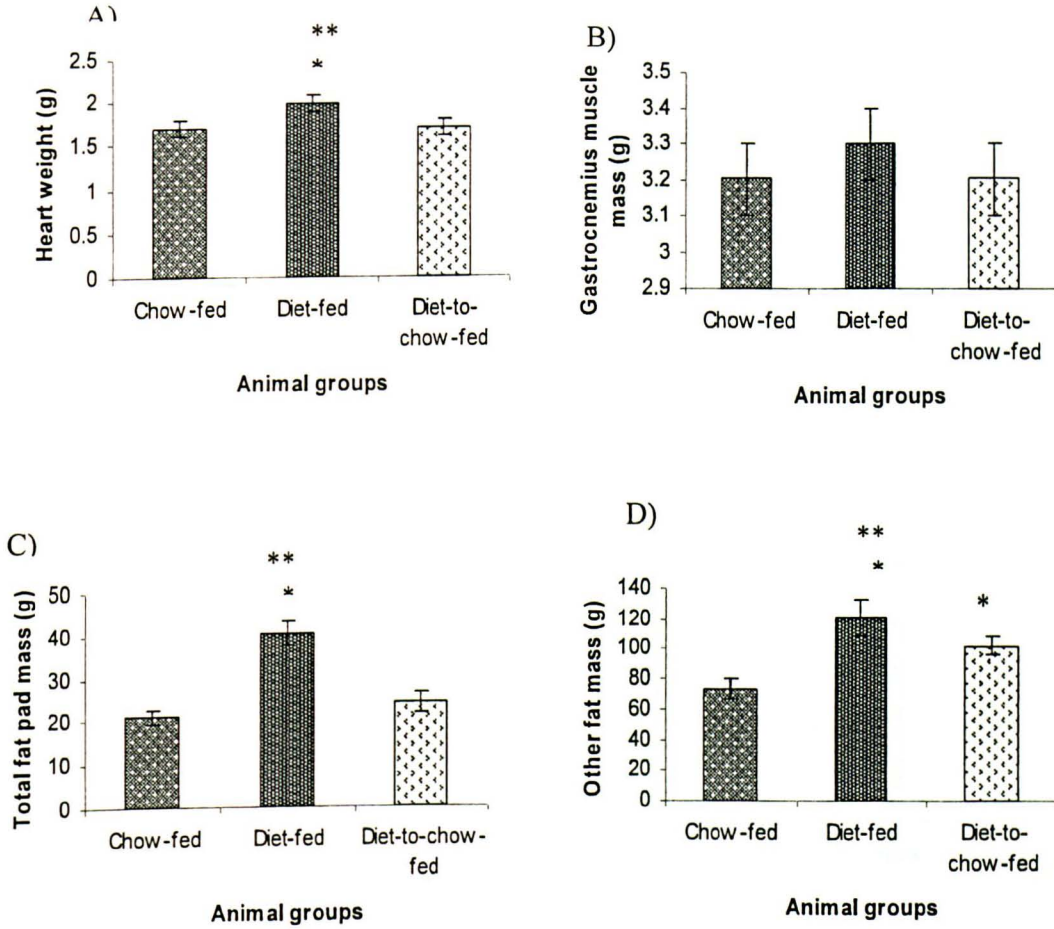


Fig. 5.2. The physiological characteristics: A) heart weight, B) gastrocnemius muscle mass, C) total fat pad mass, and D) other fat mass, of the three experimental groups at the end of the experiment. Animals were either fed standard chow (chow-fed) or a palatable diet (diet-fed) for 15 weeks, or fed the palatable diet for 7 weeks and then returned to standard chow for further 8 weeks (diet-to-chow). Data are expressed as mean \pm S.D. $p < 0.001$ for all. *Diet-fed or diet-to-chow vs chow-fed. **diet-fed vs diet-to-chow-fed.

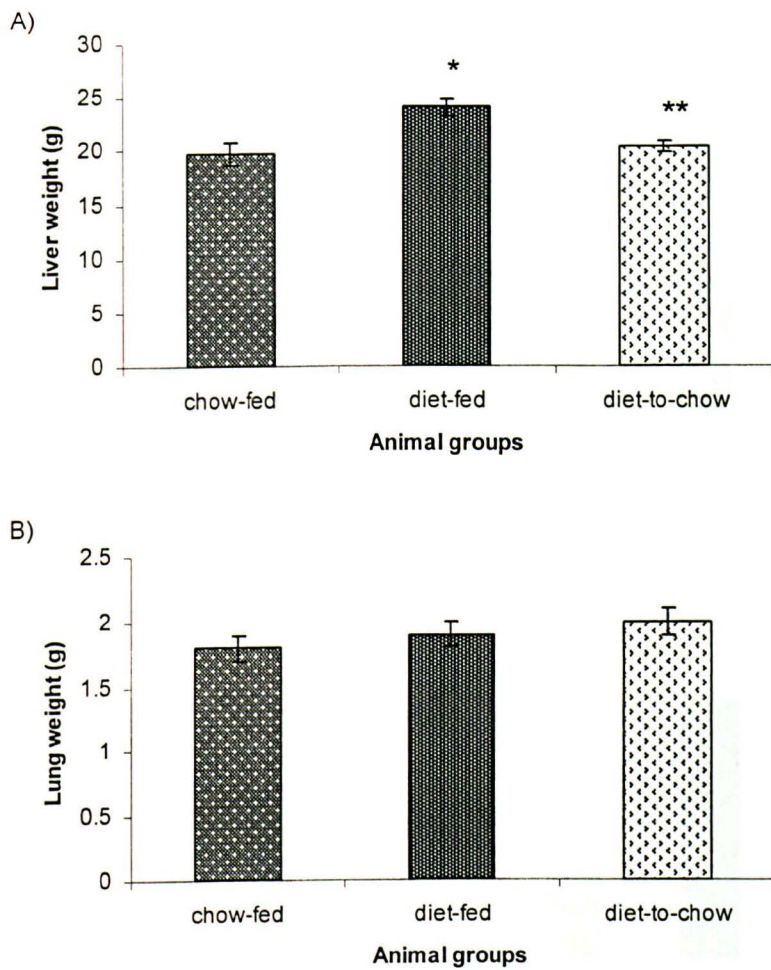


Fig. 5.3. The effect of diet-induced obesity and chronic withdrawal of the palatable diet on the weights of A) liver, and B) lung, in male Wistar rats. Lean control animals were fed a standard laboratory chow, while the second group was given high-palatable diet for 15 weeks (diet-fed). In the third group, the palatable-diet was removed after 7 weeks and replaced by standard laboratory chow for the last 8 weeks (diet-to-chow). Data are expressed as mean \pm S.E.M. * $p < 0.008$ vs chow-fed, and ** $p < 0.05$ vs diet-fed.

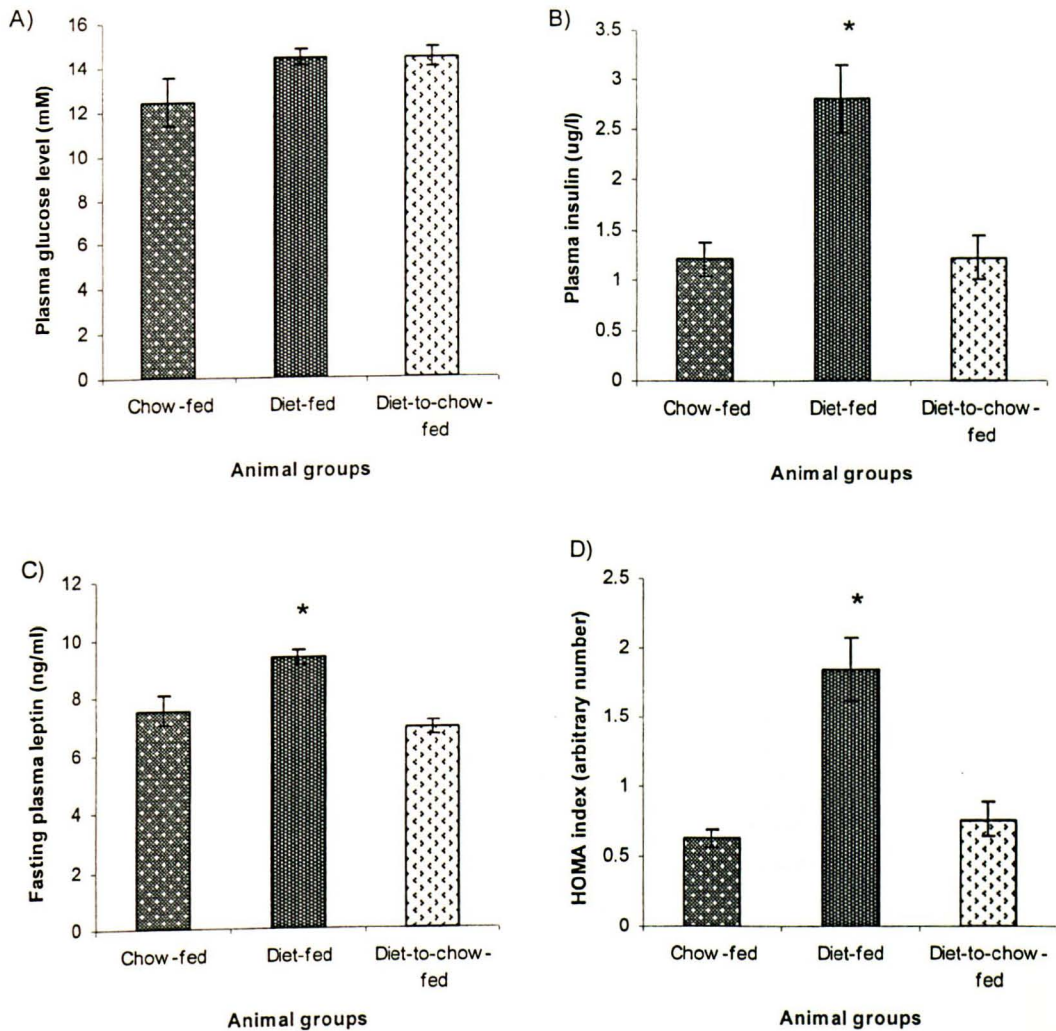


Fig. 5.4. Effects of highly-palatable diet feeding and its withdrawal on fasting plasma levels of A) glucose, B) insulin, C) leptin, and D) HOMA index in three experimental animals. There were no significant changes in glucose levels in three groups. Diet-fed rats had significantly (for all $p < 0.01$) higher fasting insulin, and leptin levels than both chow-fed and diet-to-chow groups. Diet-fed animals were also insulin resistance, indicated by a higher HOMA index. Fasting insulin and leptin levels returned to normal levels by long-term removal of the highly-palatable diet and insulin resistance was corrected subsequently. Data are presented as mean \pm S.E.M. * $p < 0.01$ vs chow-fed.

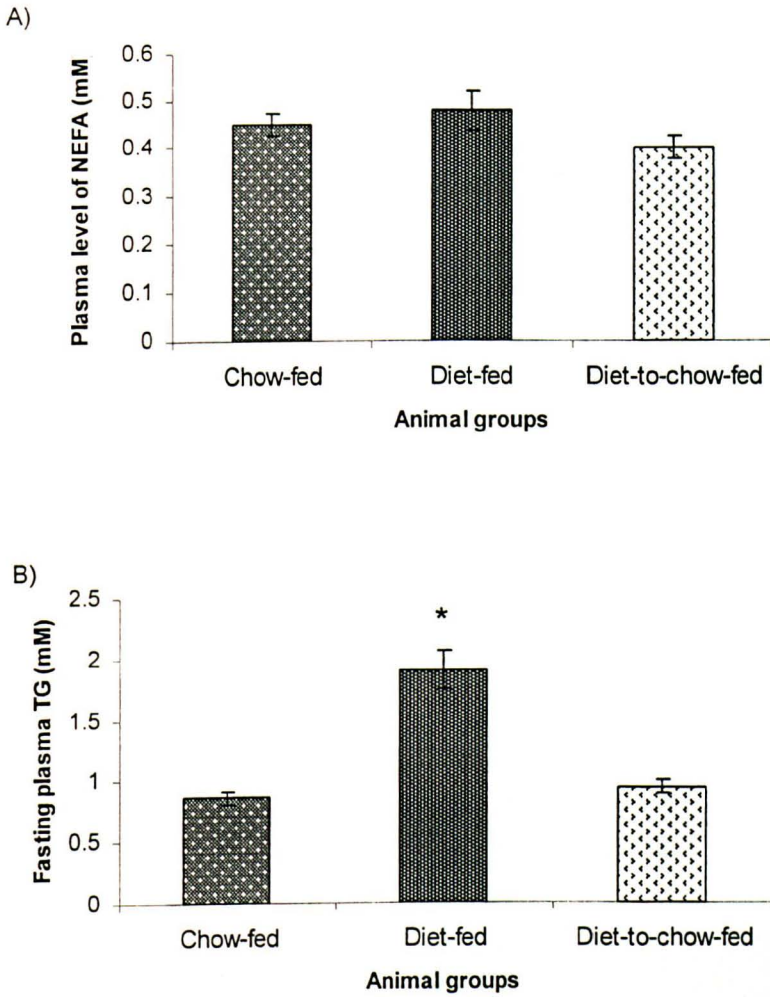


Fig. 5.5. Effects of highly-palatable diet feeding and its withdrawal on fasting plasma levels of A) Non-esterified fatty acid (NEFA), and B) triglyceride (TG) in rats. Levels of NEFA were comparable in all three groups. Diet-fed rats had significantly (for both, $*p < 0.001$) higher fasting TG levels than both chow-fed and diet-to-chow groups. Data are expressed as mean \pm S.E.M.

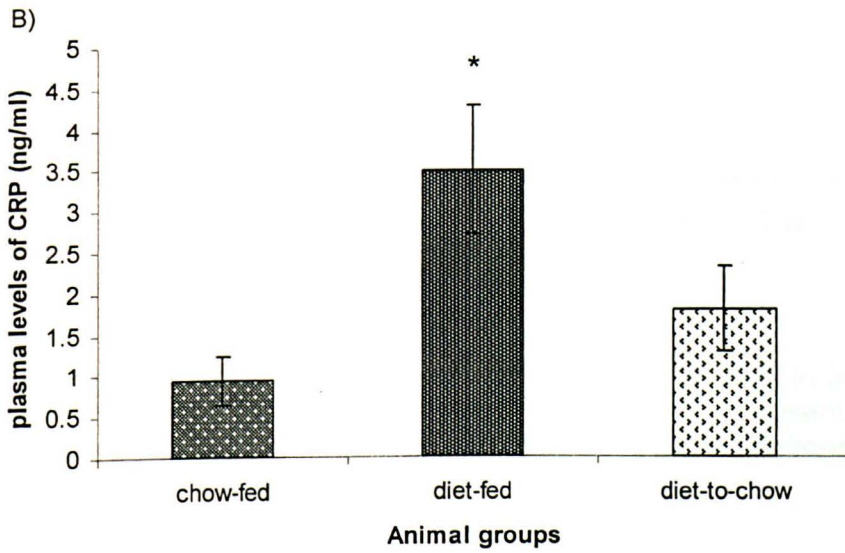
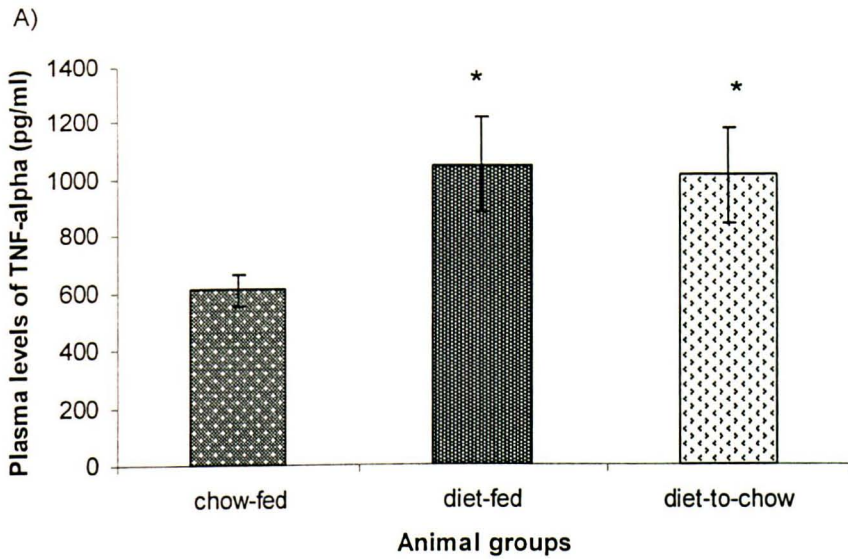


Fig. 5.6. the effects of diet-induced obesity and chronic withdrawal of the palatable diet on plasma levels of A) TNF- α , and B) C-reactive protein (CRP) in male Wistar rats. Lean control animals were fed a standard laboratory chow, while the second group was given high-palatable diet for 15 weeks (diet-fed). In the third group, the palatable-diet was removed after 7 weeks and replaced by standard laboratory chow for the last 8 weeks (diet-to-chow). Data are expressed as mean \pm S.E.M. * $p < 0.05$ vs chow-fed.

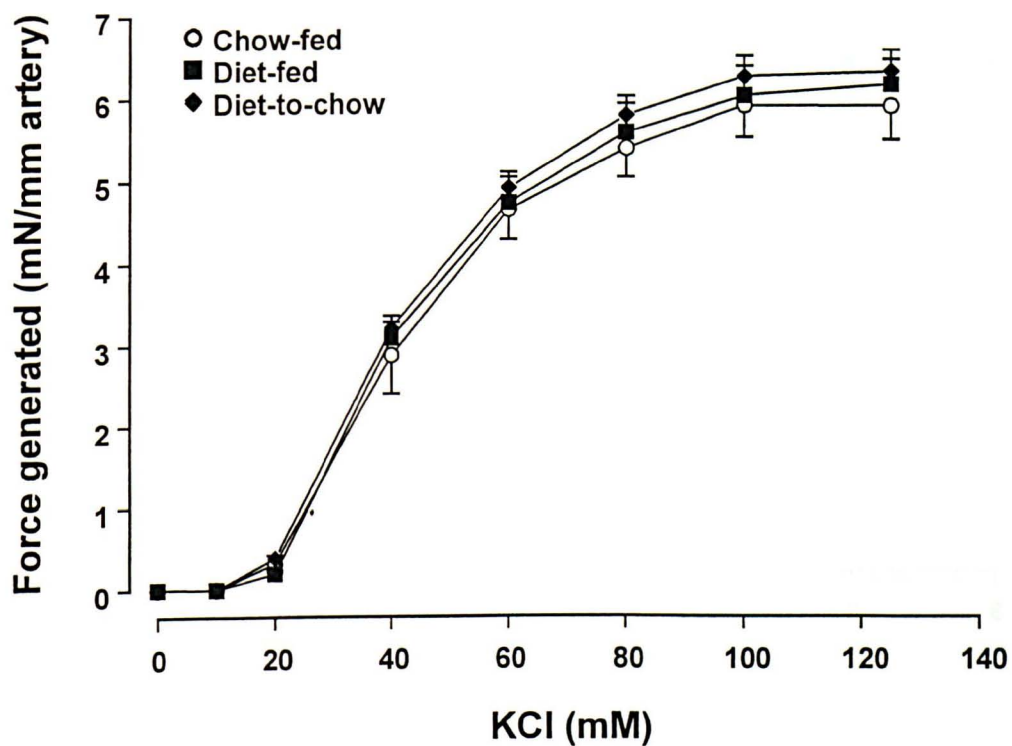


Fig. 5.7. Concentration-response curves for KCl (10 – 125 mM) in arteries from chow-fed, diet-fed and diet-to-chow-fed animals. Data represent mean \pm S.E.M for 6 animals in each group. KCl-induced vasoconstrictions were similar between all three groups.

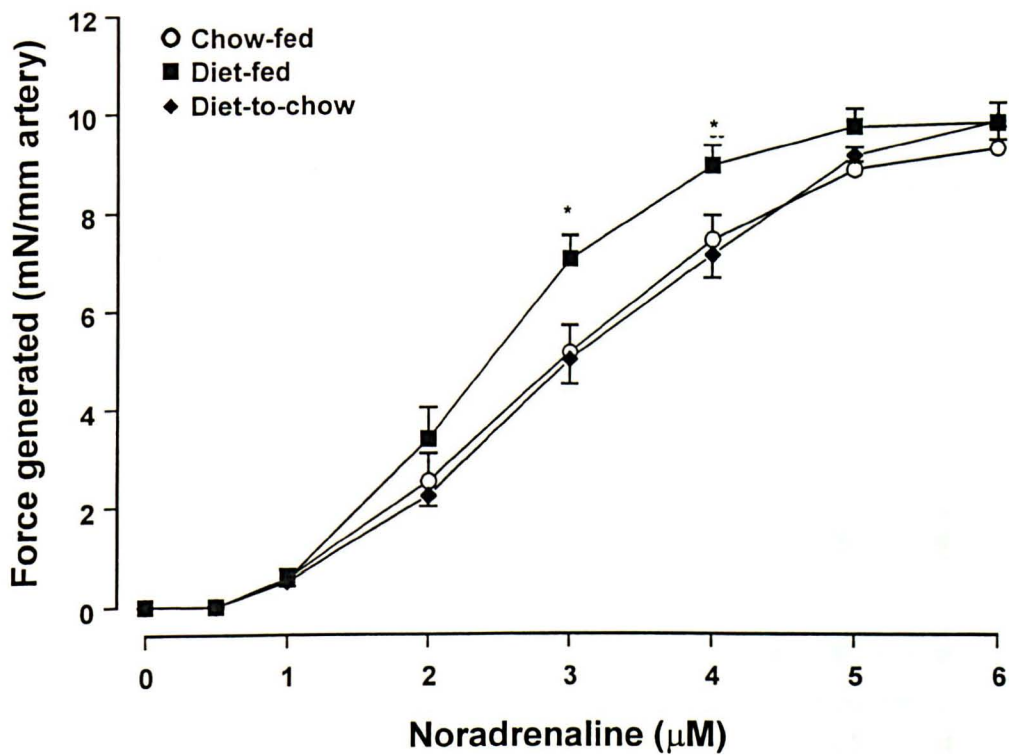


Fig. 5.8. Concentration-response curves for noradrenaline (NA; 0.5 – 7 µM) in arteries from chow-fed, diet-fed and diet-to-chow-fed animals. Data represent mean ± S.E.M. for 6 animals in each group. However, 3 and 4 µM NA-induced contractions significantly ($*p < 0.01$; ANOVA, Bonferroni test) were greater than both chow-fed and diet-to-chow-fed groups.

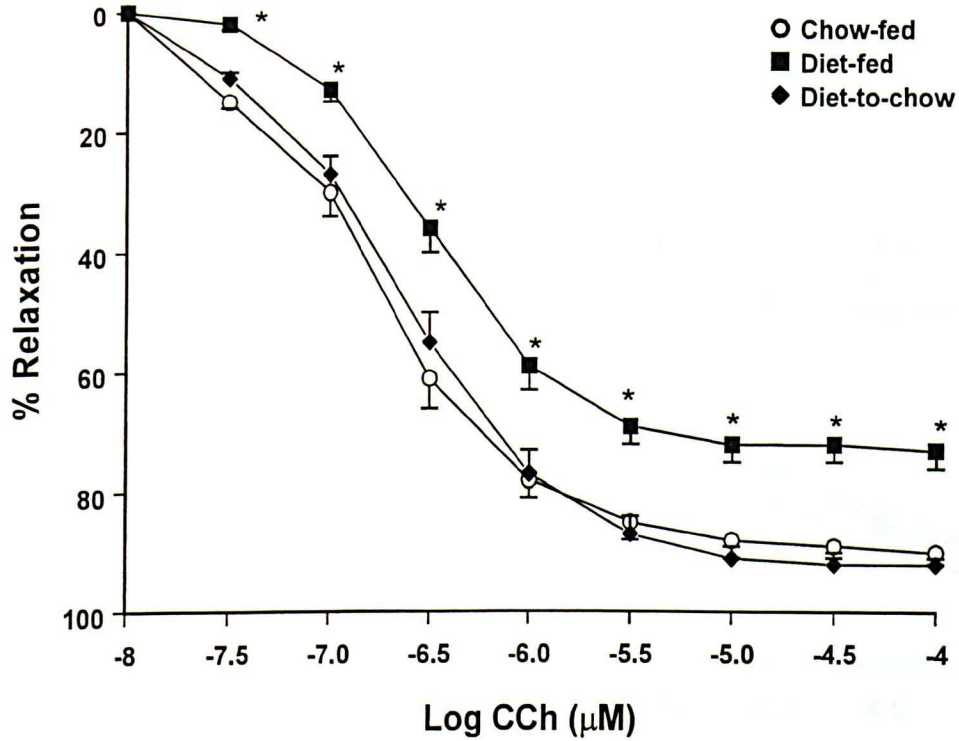


Fig. 5.9. Relaxation response curves (mean \pm S.E.M) of carbamylcholine (CCh) in the 3 groups. Arteries were contracted with 8 μM Noradrenaline. When contractions reached a plateau cumulative concentration responses to CCh were carried out. CCh-induced vasorelaxation was significantly reduced in diet-fed animals ($*p, 0.001$; ANOVA, Bonferroni test) in comparison to chow-fed group. This abnormality was completely abolished by the chronic removal of the palatable diet.

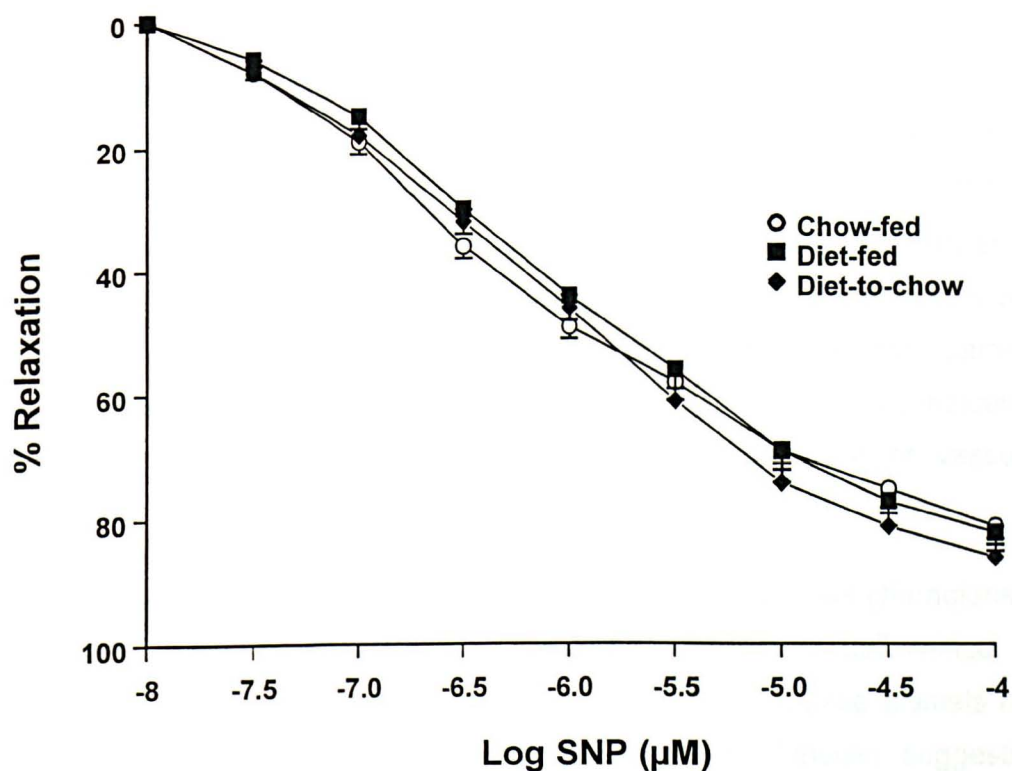


Fig. 5.10. Relaxation response curves (mean \pm S.E.M) of sodium nitroprusside (SNP) in the 3 groups. Arteries were contracted with 8 μ M Noradrenaline. When contractions reached a plateau cumulative concentration responses to SNP were carried out. SNP-induced vasorelaxations were similar in all three groups.

5.4 DISCUSSION

The present study, demonstrates for the first time that chronic removal of an obesity-inducing palatable diet corrects all the metabolic abnormalities, cardiac hypertrophy and vascular dysfunctions associated with obesity. It has been demonstrated previously that chronic consumption of a palatable diet induces significant level of obesity and markedly effects metabolic parameters and vascular reactivity in the absence of any visible insulin resistance (Naderali *et al.*, 2001a). Moreover, short-term feeding (2-days) of the same palatable diet significantly alters lipid profile and markedly attenuates endothelial function in the absence of any significant obesity (Naderali *et al.*, 2001c). These defects were only partially corrected by acute removal of the diet (Naderali and Williams, 2003), indicating that diet *per se* may have a major role in the development of vascular abnormalities in animals.

In this study, diet-fed animals developed marked obesity, which was characterised by more than 2-fold increase in total body fat content as determined by Bioimpedence methods and measurements of fat pad mass. Obese animals had significantly higher heart weight and fasting plasma levels of insulin, suggesting cardiac hypertrophy and development of insulin resistance in dietary-obese animals. These findings are similar to those seen in human obesity (Andres, 1980; Barret-Conner, 1985; Reaven, 1988; Singh *et al.*, 1996). In human, obesity leads to dyslipidaemia and type 2 diabetes and hypertension (Lind *et al.*, 1995; Carey *et al.*, 1997; Parker *et al.*, 2003), while changes in life style improves obesity-induced abnormalities to a certain degree (Singh *et al.*, 1996; Sasaki *et al.*, 2002; Gutin *et al.*, 2002).

A number of experimental studies have reported severe detrimental outcome of excessive high-energy diet on vascular reactivity. For example, high-fat diet impairs endothelium-dependent vasorelaxation (Vogel *et al.*, 1997; Naderali *et al.*, 2001a; Naderali and Williams, 2001b; Naderali *et al.*, 2001c), while antioxidant therapy improves endothelial function (Sato *et al.*, 2002; Plotnick *et al.*, 1997), suggesting presence of diet-induced oxidative stress in the vasculature. In obesity,

plasma antioxidant levels are decreased (Kuno *et al.*, 1998; Sato *et al.*, 2002; Plotnick *et al.*, 1997), whereas, myocardial oxidative stress is increased (Moor-de Burgos *et al.*, 1992; Vincent *et al.*, 1999). These changes may, at least in part, explain the increased risk of vascular disease and the abnormal endothelium-dependent vasorelaxation found in obese subjects and animals. In obese patients with hypertension, ACh-induced forearm blood flow (FBF) is significantly reduced compared to healthy individuals. Caloric restriction decreases body weight and the mean blood pressure, fasting plasma insulin, total cholesterol, triglycerides, low-density lipoproteins (LDL) as well as enhancing ACh-induced FBF, while nitric oxide synthase inhibitors mask increase in ACh-induced FBF, indicating that reduction in energy intake improves endothelial function possibly by increasing endogenous NO production in obese hypertensive patients (Sasaki *et al.*, 2002). Lipid lowering also reduces the risk for coronary events, cardiovascular morbidity and mortality in a broad range of patients (Bucher *et al.*, 1999). Similar observations have also been reported in animal studies. In miniature pigs, 6 to 9 month of cholesterol-rich diet induces LDL-hypercholesterolaemia and advanced coronary atherogenesis. Withdrawal of dietary cholesterol reverses hypercholesterolaemia and stabilises coronary plaque in miniature pigs (Verhamme *et al.*, 2002) and partially restores endothelial function in rats (Naderali and Williams, 2003).

In the present study, diet-fed obese rats had significant vascular abnormalities, in particular a marked endothelial dysfunction. The potential factors for vascular abnormalities are obesity *per se* (Andres, 1980), insulin resistance (Reaven, 1988), increased leptin levels (Wallace *et al.*, 2001), and higher level of plasma triglyceride (Naderali *et al.*, 2001a; Naderali and Williams, 2001b; Naderali and Williams, 2003). In fact, there were negative correlations between endothelial function and plasma levels of insulin, leptin and triglycerides. The findings that changes in vascular function occurs in the absence of obesity or rise in plasma leptin levels and insulin resistance (Naderali *et al.*, 2001c; Naderali and Williams, 2003) argue against major roles of obesity *per se*, insulin and leptin resistance in vascular dysfunction. Moreover, in this study diet-to-chow rats were significantly obese compared with chow-fed controls, yet their vascular function was similar to

that of chow-fed group, further arguing, at least in the rat, against the role of obesity *per se* in the genesis of vascular abnormalities. On the other hand, recent human and animal studies point to the role of circulating triglycerides in obesity-induced vascular dysfunction (Tka *et al.*, 1997; Krauss, 1998; Zhao *et al.*, 2001; Naderali *et al.*, 2001a; Naderali and Williams, 2001b; Naderali and Williams, 2003). In this study, circulating triglyceride levels were raised more than 115% and a significant negative correlation was seen between fasting plasma triglyceride levels and endothelium-dependent vasorelaxation. The plasma TG levels of chow-fed and diet-to-chow groups were similar as were CCh-induced vasorelaxation between the two groups, suggesting that changing diet (i.e. removing palatable diet and replacing it with the standard chow) resulted in lowering plasma triglyceride levels and thereby improving vascular function.

The adipocytokines, TNF- α (Myers *et al.*, 1994; Bhagat and Valance, 1997; Locksley *et al.*, 2001; Wajant *et al.*, 2003) and CRP (Biasucci *et al.*, 1999; Buffon *et al.*, 1999; Verma *et al.*, 2002) have been documented to induce vascular and metabolic defects. TNF- α and CRP were significantly increased in diet-fed animals, suggesting possible role for these adipocytokines in obesity-induced vascular dysfunction.

Overall, this study indicates that prolonged consumption of a high-palatable, high-energy diet induces metabolic and cardiovascular disorders. Nonetheless, these abnormalities are completely reversible by chronic withdrawal of the palatable diet.

Chapter 6

Results: The effects of fenofibrate on adiposity, metabolic abnormalities, and endothelial dysfunction in dietary obese rats

6.1 INTRODUCTION

Obesity and type 2 diabetes are strong risk factors for the development of atherosclerosis. The underlying pathophysiological mechanisms are not fully understood, but may include hyperlipidaemia and increased oxidation of low-density lipoproteins (Renier *et al.*, 2000), hypertriglyceridaemia (Naderali *et al.*, 2001a), insulin resistance (Reaven, 1988) and hyperleptinaemia (Wallace *et al.*, 2001). Obesity, type 2 diabetes and the above risk factors are also associated with marked defects in vascular function. Endothelial dysfunction in particular is prominent and may contribute to atherogenesis (Renier *et al.*, 2000; Suwaidi *et al.*, 2000; Mancini *et al.*, 2001; Naderali *et al.*, 2001; Naderali *et al.*, 2001a; Naderali *et al.*, 2001c).

The lipid-lowering fibrates, such as fenofibrate and ciprofibrate have been shown to impede atheroma development. These drugs bind to peroxisome proliferator-activated receptor- α (PPAR- α), leading to enhanced lipid catabolism in the liver (Mancini *et al.*, 2001) and reductions in total body fat as well as the circulating plasma lipids. Fenofibrate has been reported to retard angiographic progression of coronary atherosclerosis in diabetic patients (Steiner *et al.*, 2001), and improve the microcirculation in patients with hyperlipidaemia (Liang *et al.*, 2000). The mechanism(s) of the beneficial effects of fenofibrate on vascular function has not been fully elucidated yet, but direct activation of PPAR- α in arterial wall (Fruchart *et al.*, 1999), correction of lipid abnormalities (Despr es *et al.*, 2002; Capell *et al.*, 2003), and increasing the formation, availability and action of NO (Haak *et al.*, 1998; Playford *et al.*, 2002) have all been postulated.

Consequently, this study aimed to further evaluate the effect of fenofibrate on adiposity, metabolic, and vascular abnormalities seen with obesity-inducing diet in obese rats.

6.2 METHODS

6.2.1 Experimental protocol

Adult male Wistar rats (n=27) were randomised and assigned to a control group (n=9, 191 ± 3 g) and a test group (n=18, 190 ± 1 g) that was fed a highly palatable diet for 15 weeks.

All animals had free access to water and were housed individually under controlled environmental conditions (19-22 °C; 30-40% humidity) and a 12-hour light/dark cycle (lights on at 07:00h). Controls were fed a standard laboratory pelleted diet (chow-fed). The test group had free access to a highly-palatable diet. After 8 weeks, half of the test group was given fenofibrate (50 mg/kg/day) and the remainder were given vehicle (1% carboxymethyl cellulose at 1 ml/kg body weight; Sigma, Pool, UK) by oral gavage, daily for 7 weeks (for more details see chapter 2, section 2.3.2). On the day of experiment, the rats were killed by CO₂ inhalation after 2 hours of fasting. The total body fat mass was measured immediately by Bioimpedance Method using the TOBEC® equipment (Biotech Instruments Ltd, Kimpton, UK). The gonadal and perirenal fat pads, the gastrocnemius and soleus muscles, liver, heart, and lungs were dissected and weighed.

6.2.2 Biochemical analysis

Blood was removed by cardiac puncture into cold heparinised tubes. The plasma was immediately separated by centrifugation before being frozen for later measurements of glucose, insulin, leptin, TNF- α , CRP, NEFA and TG. Plasma glucose concentration was determined using a glucose oxidase method, and NEFA and TG concentrations using commercial diagnostic kits. Insulin and leptin concentrations were measured by RIA kits, while plasma TNF- α and CRP concentrations were measured using a commercially available ELISA kits.

6.2.3 Homeostasis Model Assessment (HOMA)

HOMA, an index of insulin resistance, which employs measures of fasting plasma concentrations of glucose and insulin, was calculated according to the method described previously (Matthews *et al.*, 1985).

6.2.4 Assessment of vascular function

Six third-order mesenteric arteries (<250 μm diameter, 2-mm lengths) were carefully dissected from each animal. Each artery was freed of fat and connective tissue and mounted on two 40- μm diameter stainless-steel wires in an automated myograph (Cambustion, Cambridge, UK), based on the principle of the Mulvany myograph which measures isometric tension generated in response to various stimuli. Arterial segments (two at a time) were incubated in a 5-ml organ bath containing physiological salt solution gassed with 95% O_2 and 5% CO_2 at 37 °C.

Cumulative concentration-response curves to either KCl (10-125 mM) or noradrenaline (NA, 0.5-6 μM) were then carried out. Any vessel failing to reach its predetermined target tension in response to vasoconstriction with KCl (125 mM) was discarded (for more details see chapter 2).

6.2.5 Assessment of defect in vascular relaxation

Endothelial-dependent and –independent defects were investigated by exposing NA-precontracted arteries to CCh or SNP, respectively. To eliminate possible interference by vasoactive prostanoids, 10 mM indomethacin was added to the organ bath prior to contraction with NA (8 μM). When contraction reached a plateau (after 2 minutes) concentration-response curves were performed to either CCh or SNP (for both, 10 nM-100 μM).

6.2.6 Data interpretation and statistical analyses

Vasoconstriction in response to NA and KCl were expressed as absolute force generated (mN/mm artery). Relaxation in response to CCh and SNP was calculated as the percentage reduction from the maximal tension generated at the supramaximal concentration of NA (8 μM). The average response for all the

vessels from a given animal was used in the group analysis. EC₅₀ values were determined to evaluate changes in agonist-induced responses.

Data showed a normal distribution (Shapiro-Wilkes W test) and are therefore expressed as mean \pm S.E.M. Statistical significance was tested using Student 't-test' or repeated-measures (ANOVA; Bonferroni t-test) or the Mann-Whitney test, as appropriate. Differences were considered statistically significant at the $p < 0.05$ levels.

6.3 RESULTS

6.3.1 Effects of fenofibrate on body weight and adiposity

Animals fed palatable diet progressively gained more weight than chow-fed controls. This difference became significant after the third week of feeding and was maintained thereafter. The final body weight of the diet-fed animals was significantly higher than that of chow-fed controls ($>21\%$; 695 ± 17 vs 572 ± 8 g, $p < 0.0001$). Overall, the weekly weight gain of diet-fed animals was significantly greater than that of chow-fed controls (diet-fed: 20.1 ± 4.77 vs control: 11.98 ± 4.65 g, $p = 0.013$) (Fig. 6.1). Administration of fenofibrate significantly reduced weekly weight gain of diet-fed animals and final total body weight (Fig. 6.1). After 7 weeks of treatment, total body weight of the diet-fed fenofibrate-treated animals was not significantly different from that of chow-fed controls (fenofibrate treated: 593 ± 16 vs chow-fed: 572 ± 8 g, $p < 0.2$), but it was significantly lower ($p < 0.0007$) than that of untreated diet-fed animals (Fig. 6.1).

After 15 weeks of study, gastrocnemius and soleus muscle masses were comparable between all three groups. However, diet-fed animals had significantly higher total white adipose tissue (sum of epididymal and perirenal fat pads) mass than both chow-fed (by >2 -fold, $p < 0.0002$) and fenofibrate-treated (by >1.5 -fold, $p < 0.01$) groups (Figs. 6.2 a, b, c & d). Fenofibrate-treated animals also had significantly (by >1.4 -fold $p < 0.02$) greater white adipose tissue than chow-fed controls. Bioimpedance measurements indicated higher percentage of body-fat in diet-fed animals compared with chow-fed and fenofibrate-treated groups. Conversely, diet-fed rats had significantly lower percentage of lean mass than

chow-fed and fenofibrate-treated animals (Fig. 6.2 e, & f). Liver weight increased significantly in fenofibrate treated group compared with chow-fed (by > 213%, $p < 0.00001$) and diet-fed groups (by > 174%, $p < 0.008$), while there was no significant difference between chow-fed and diet-fed groups. Moreover, heart weight also increased significantly in fenofibrate-treated group, compared with chow-fed group (by > 111%, $p < 0.01$), whereas lungs weight was comparable in all three groups with no significant difference (Fig. 6.3 a, b, and c).

6.3.2 Effects of diet and fenofibrate on blood metabolites

Palatable diet-feeding did not effect fasting plasma glucose levels (11.7 ± 1.2 chow-fed; 14.3 ± 0.3 mM diet-fed; $p < 0.06$) (Fig. 6.4 a). However, diet-fed rats had significantly higher fasting terminal insulin (3.8 ± 0.8 diet-fed vs 1.6 ± 0.3 mM chow-fed; $p < 0.002$) than the chow-fed group (Fig.6.4 b), with a significantly higher HOMA index value (2.40 ± 0.52 diet-fed vs 0.73 ± 0.10 chow-fed; $p < 0.008$) indicating development of insulin resistance in diet-fed animals (Fig.6.5 a). Fenofibrate treatment significantly lowered levels of both glucose (10.8 ± 0.3 mM; $p < 0.0001$ vs untreated diet-fed group) and insulin (1.4 ± 0.2 mM; $p < 0.001$ vs untreated diet-fed group) levels. Consequently, HOMA index was also reduced (0.67 ± 0.10 ; $p < 0.007$ vs untreated diet-fed group), indicating correction of insulin resistance (Fig. 6.5 a).

Diet-fed rats had significantly higher plasma leptin levels than chow-fed control (9.4 ± 0.7 diet-fed vs 6.9 ± 0.5 ng/ml chow-fed; $p < 0.0131$) (Fig. 6.5 b). The rise in leptin levels was not affected by fenofibrate treatment (8.5 ± 0.4 ng/ml; $p < 0.03$ vs chow-fed). Fasting plasma levels of NEFA were comparable in diet-fed and chow-fed groups (0.44 ± 0.02 chow-fed; 0.45 ± 0.04 mM diet fed) (Fig. 6.6 a). However, diet-fed rats had significantly higher (>110%) levels of fasting TG than chow-fed (chow-fed: 0.79 ± 0.04 ; diet-fed: 1.68 ± 0.17 mM; $p < 0.0003$) (Fig. 6.6 b). Fenofibrate treatment significantly lowered fasting plasma levels of both NEFA (0.31 ± 0.02 mM; $p < 0.001$ vs chow-fed and untreated diet-fed groups) and TG (0.80 ± 0.06 mM $p < 0.001$ vs untreated diet-fed group) below those of chow-fed group.

6.3.3 Effects on TNF- α and CRP

Diet-fed animals had significantly ($p < 0.05$) higher TNF- α than their chow-fed counterparts (608.6 ± 56 chow-fed vs 1046.9 ± 166.9 pg/ml diet-fed). Fenofibrate-treated animals had significantly ($p < 0.04$) lower TNF- α (532 ± 91 pg/ml) than their obese untreated counterparts. There were no significant differences between TNF- α level of chow-fed and fenofibrate treated animals (Fig. 6.7 a). Similarly, compared to chow-fed group, CRP levels increased significantly ($p < 0.05$) in diet-fed animals (3.5 ± 0.8 diet-fed vs. 0.9 ± 0.3 ng/ml chow-fed). Interestingly, CRP levels further increased significantly ($p < 0.001$) by fenofibrate-treatment (6.3 ± 0.7 ng/ml) (Fig. 6.7 b).

6.3.4 Vascular data

Diet-induced obesity did not alter vessel diameter, and there were no significant differences in arterial diameter between the three groups (200 ± 15 in chow-fed; 205 ± 15 in diet-fed; 200 ± 10 μm in fenofibrate-treated).

6.3.5 Agonist-induced vasoconstriction responses

Cumulative concentration response curves to KCl (10-125 mM) and NA (0.5-6 μM) in arteries from all three groups showed the characteristic sigmoid shape. KCl-induced contraction was similar in all three groups (Fig. 6.8 a), and the concentration response curves were virtually superimposable, reaching maximum force of 5.68 ± 0.44 (chow-fed), 6.07 ± 0.33 (diet-fed) 5.35 ± 0.37 (fenofibrate-treated) mN/mm artery. There was no significant differences between the groups, either overall (by ANOVA) or at any given KCl concentration. A similar pattern was also seen with NA-induced contractions (Fig. 6.8 b), giving maximum force of 9.14 ± 0.63 (chow-fed), 9.94 ± 0.43 (diet-fed) and 9.70 ± 0.84 (fenofibrate-treated) mN/mm artery.

6.3.6 Endothelium-dependent and -independent vasorelaxation

NA-precontracted arteries from chow-fed rats demonstrated progressive relaxation to cumulative addition of CCh (10 nM- 100 μM), achieving a maximum of $88 \pm 2\%$ at a CCh concentration of 100 μM (Fig. 6.9 a), with EC_{50} values of 0.28 ± 0.03 μM .

Arteries from diet fed rats that were similarly exposed to CCh displayed a significant rightward shift of the concentration-response curve, as compared with chow-fed control arteries (EC_{50} diet-fed: $0.94 \pm 0.04 \mu\text{M}$ $p < 0.01$ vs. chow-fed). Maximal vasorelaxation to $100 \mu\text{M}$ CCh was reduced by 17% in diet-fed rats ($p < 0.005$ vs. chow-fed). Fenofibrate-treatment did not improve EC_{50} values ($0.86 \pm 0.04 \mu\text{M}$, $p < 0.002$ vs. chow-fed), but it enhanced maximal CCh-induced vasorelaxation at higher concentrations ($>10 \mu\text{M}$), achieving $86 \pm 3\%$ relaxation at $100 \mu\text{M}$ CCh (Fig. 6.9 a).

Feeding the high-palatable diet did not effect concentration-dependent vasorelaxation induced by SNP (10 nM - $100 \mu\text{M}$) (Fig. 6.9 b). The concentration-relaxation response curves were identical and virtually superimposable in all three groups, with EC_{50} values of 2.08 ± 0.07 (chow-fed); 1.73 ± 0.12 (untreated obese), and $1.46 \pm 0.10 \mu\text{M}$ (fenofibrate-treated). There were no statistically significant differences between all three groups, either overall (by ANOVA) or at any given SNP concentration.

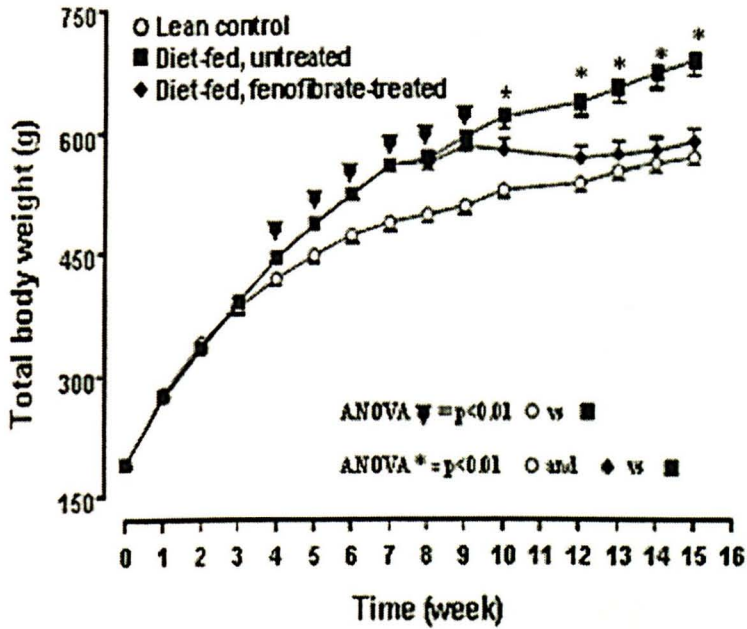


Fig. 6.1. The effects of highly-palatable-diet and fenofibrate on total body weight in male Wistar rats. Lean control animals were fed a standard laboratory chow, while diet-fed group was given a palatable diet for 15 weeks. Fenofibrate-treated animals were given fenofibrate (50 mg/kg/day) daily for the last 8 weeks of the experiment. At the end of 15 weeks, untreated diet-fed animals were significantly ($p < 0.0001$) heavier than lean control and fenofibrate-treated groups. Data are expressed as mean \pm S.E.M.

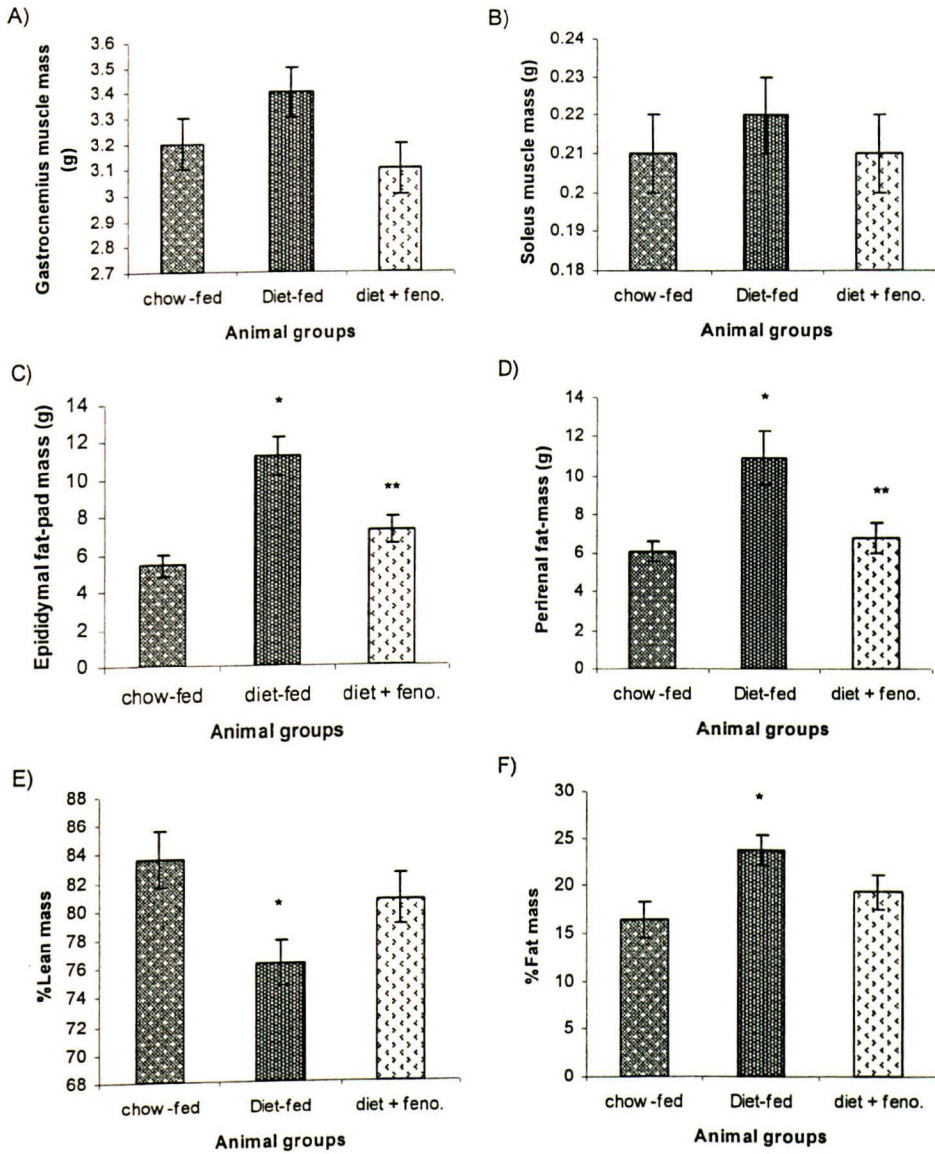


Fig. 6.2. The effects of fenofibrate on the weight of: A) Gastrocnemius muscle mass, B) Soleus muscle mass, C) Epididymal fat-pad mass (* $p < 0.001$ vs chow-fed, ** $p < 0.001$ vs diet-fed), D) Perirenal fat-mass (* $p < 0.001$ vs chow-fed, ** $p < 0.001$ vs. diet-fed), E) % Lean mass (* $p < 0.01$ vs chow-fed and diet + fenofibrate treated groups), and F) % fat mass (** $p < 0.01$ vs chow-fed and diet + fenofibrate treated groups), in dietary-induced obese rats. Lean control animals were fed a standard laboratory chow while the two diet-fed groups were given a palatable diet for 15 weeks. Fenofibrate was given to one of the diet groups (50 mg/kg/day) daily for the last 8 weeks. Data are mean \pm S.E.M.

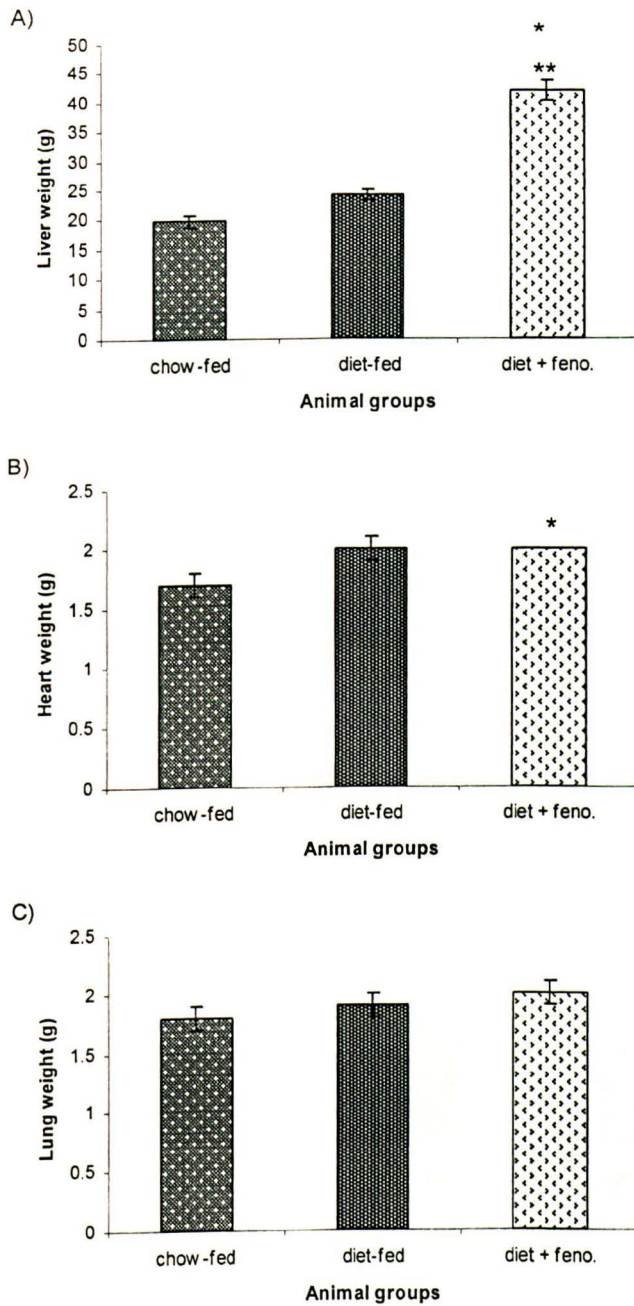


Fig. 6.3. The effect of fenofibrate treatment on the weights of: A) liver, ($*p < 0.0001$ vs chow-fed, and $**p < 0.0001$ vs diet-fed), B) heart, ($*p < 0.02$ vs chow-fed) and C) lung in dietary-induced obese rats. Lean control animals were fed a standard laboratory chow, while the two diet-fed groups were given a highly-palatable diet for 15 weeks. Fenofibrate was given to one of the diet groups (50 mg/kg/day) daily for the last 8 weeks. Data are expressed as mean \pm S.E.M.

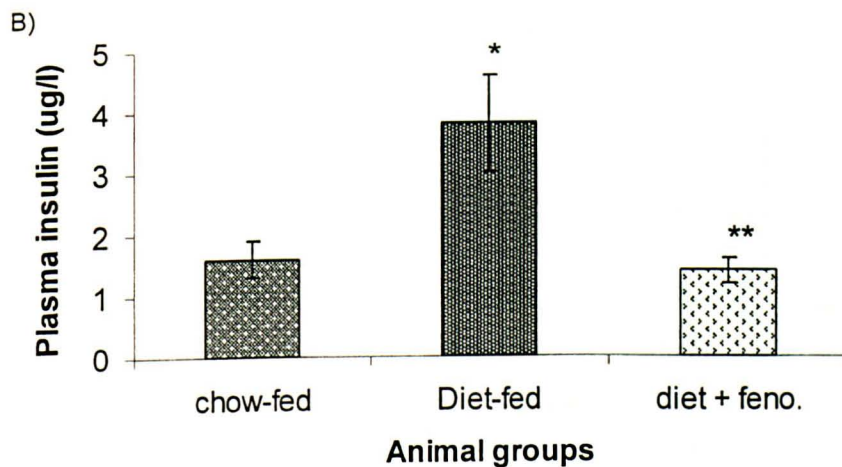
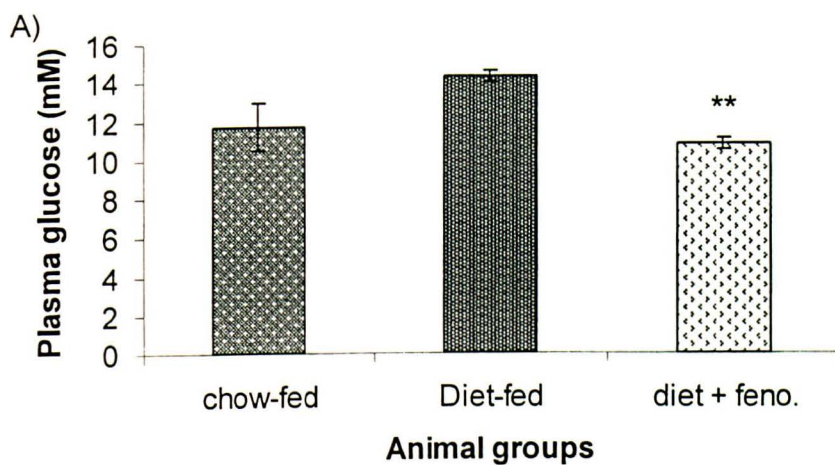


Fig. 6.4. The effects of fenofibrate on the concentration of: A) Plasma glucose, and B) Plasma insulin, in dietary-induced obese rats. Lean control animals were fed a standard laboratory chow while the two diet-fed groups were given a palatable diet for 15 weeks. Fenofibrate was given to one of the diet groups (50 mg/kg/day) daily for the last 8 weeks. Data are mean \pm S.E.M. * $p < 0.001$ vs chow-fed, and ** $p < 0.001$ vs diet-fed.

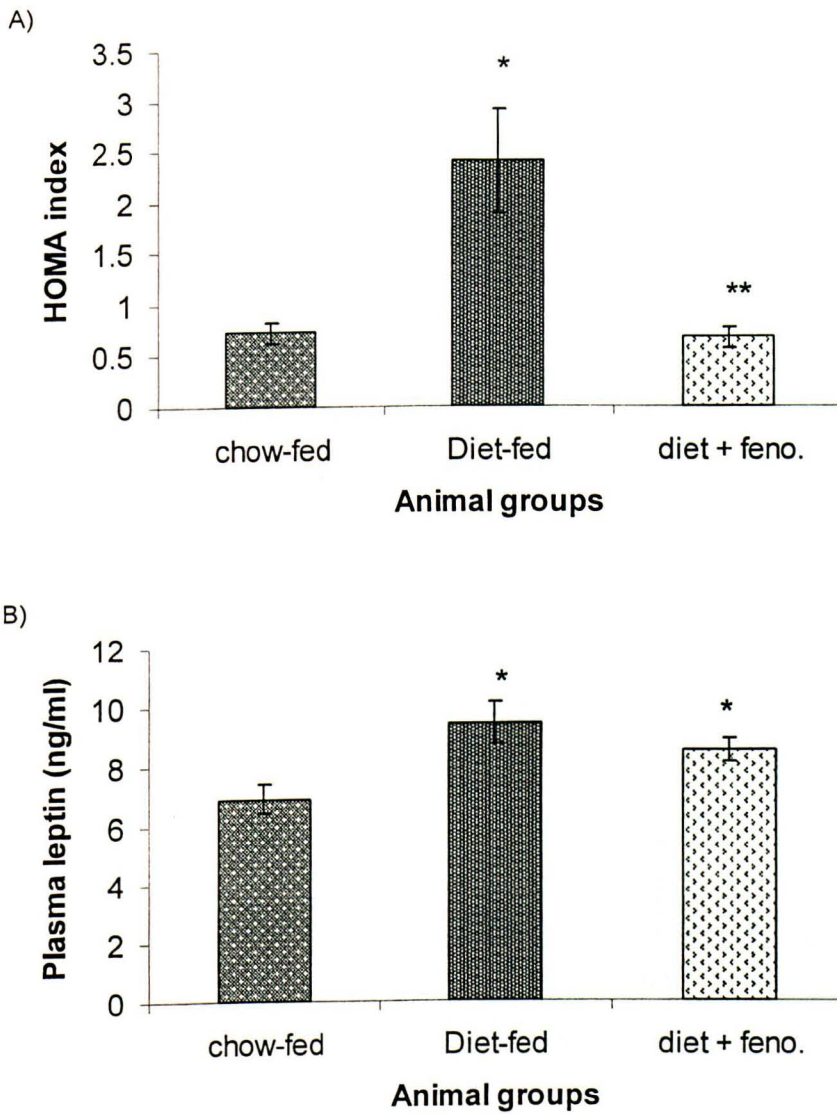


Fig. 6.5. The effects of fenofibrate on: A) HOMA index, and B) the concentration of Plasma leptin, in dietary-induced obese rats. Lean control animals were fed a standard laboratory chow, while the two diet-fed groups were given a palatable diet for 15 weeks. Fenofibrate was given to one of the diet groups (50 mg/kg/day) daily for the last 8 weeks. Data are mean \pm S.E.M. * p <0.001 vs chow-fed, and ** p <0.001 vs diet-fed.

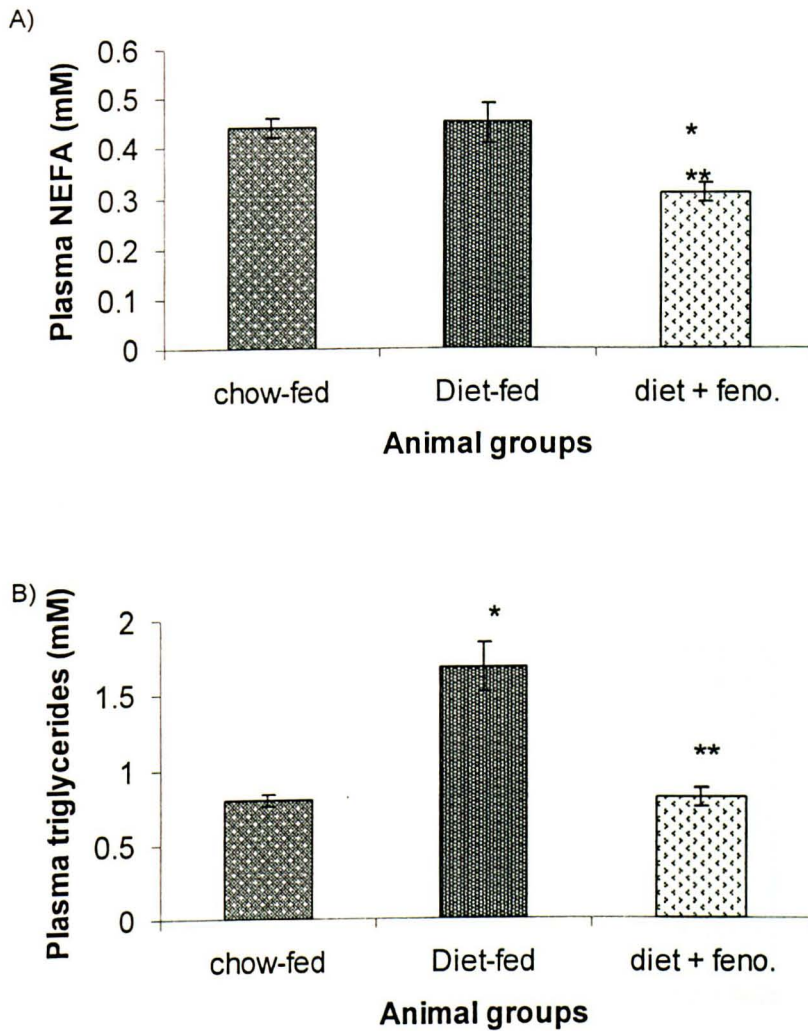


Fig. 6.6. The effects of fenofibrate on the concentration of: A) Plasma NEFA, and B) Plasma triglycerides, in dietary-induced obese rats. Lean control animals were fed a standard laboratory chow while the two diet-fed groups were given a palatable diet for 15 weeks. Fenofibrate was given to one of the diet groups (50 mg/kg/day) daily for the last 8 weeks. Data are mean \pm S.E.M. * $p < 0.001$ vs chow-fed, and ** $p < 0.001$ vs diet-fed.

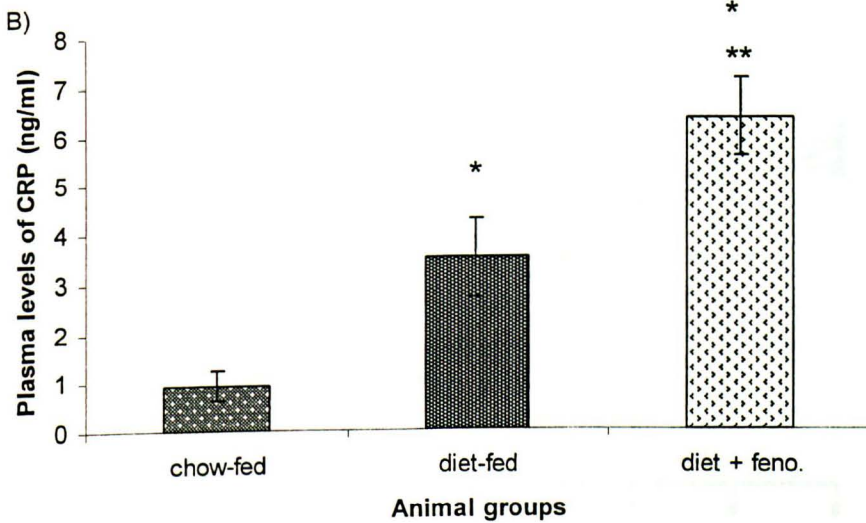
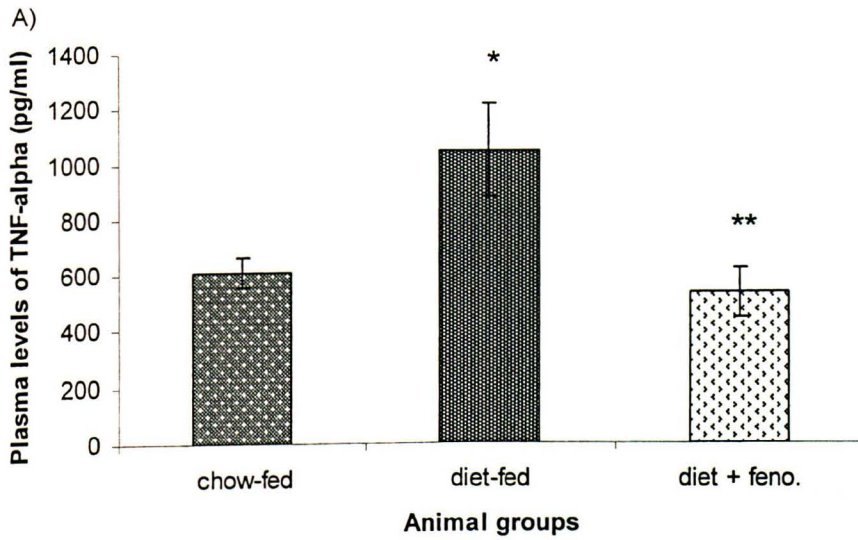
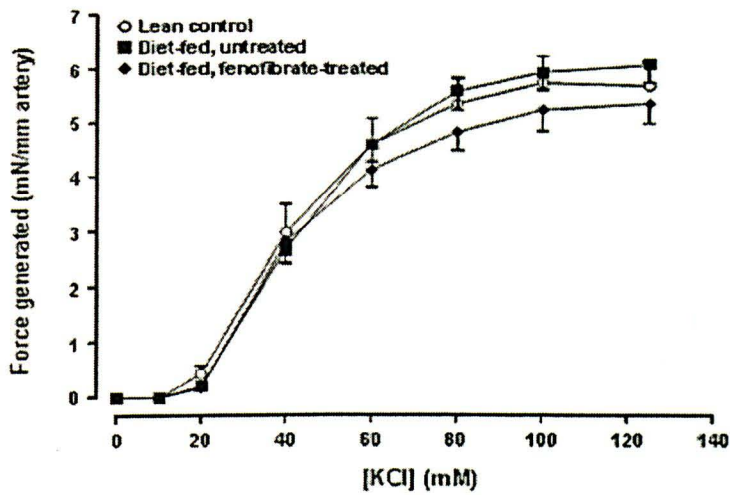


Fig. 6.7. The effect of fenofibrate treatment on plasma levels of A) TNF- α , ($*p < 0.05$ vs chow-fed and $**p < 0.05$ vs diet-fed) and B) C-reactive protein (CRP) ($*p < 0.0001$ vs chow-fed and $**p < 0.02$ vs diet-fed) in dietary-induced obese rats. Lean control animals were fed a standard laboratory chow while the two diet-fed groups were given a palatable diet for 15 weeks. Fenofibrate was given to one of the diet groups (50 mg/kg/day) daily for the last 8 weeks. Data are expressed as mean \pm S.E.M.

a)



b)

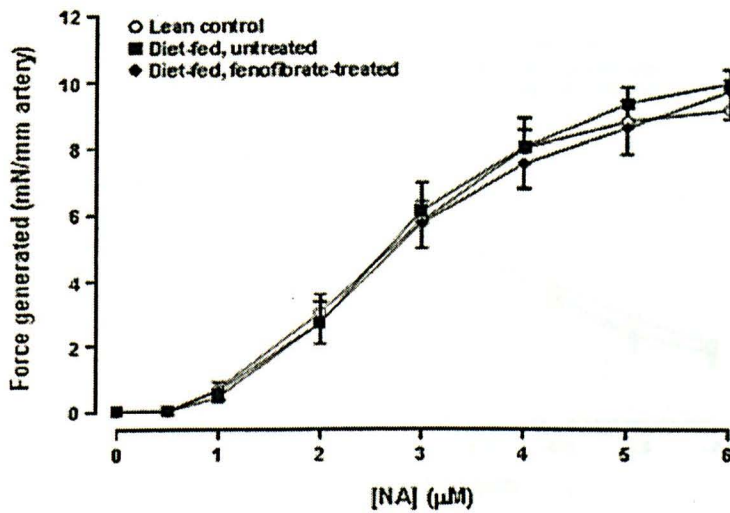


Fig. 6.8. Vasoconstriction responses to cumulative concentration of: a) KCl (10 – 125 mM) and b) noradrenaline (0.5 – 7 μM) in arteries from lean control, untreated diet-fed and fenofibrate-treated diet-fed animals. Data represent mean \pm S.E.M for 9 animals in each group. There were no significant differences in agonist-induced vasoconstriction between the three groups.

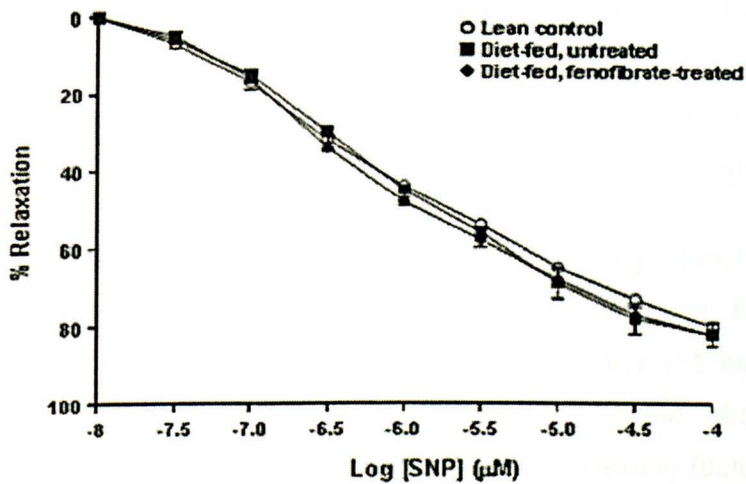
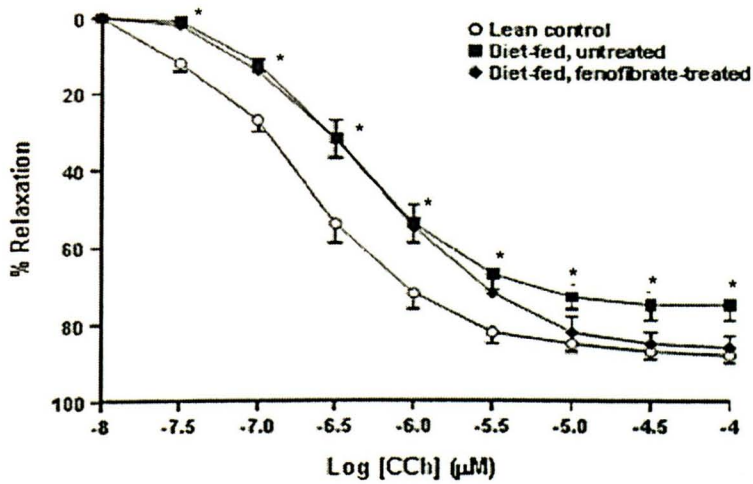


Fig. 6.9. The effects of (a) carbamylcholine (CCh), (b) sodium nitroprusside (SNP) on arteries from lean control, untreated diet-fed and fenofibrate-treated diet-fed rats. Noradrenaline ($8 \mu\text{M}$) precontracted arteries challenged with cumulative concentrations of CCh or SNP. CCh-induced vasorelaxation was significantly reduced in diet-fed animals ($*p < 0.001$; ANOVA, Bonferroni test) in comparison to lean control group. Moreover, vasorelaxation curves were shifted to the right in both diet-fed groups. Fenofibrate-treatment only partially restored CCh-induced vasorelaxation. Data represent mean \pm S.E.M for 9 animals in each group.

6.4 DISCUSSION

Various studies in animal models and in subjects with diabetes have shown that PPAR- α activation reduces diet-induced weight gain and visceral fat mass and improves insulin resistance (Guerre-Millo *et al.*, 2000; Lee *et al.*, 2002). The precise mechanism(s) by which PPAR- α improves insulin resistance is not fully known yet, but increasing fatty acid catabolism in liver, decreasing skeletal muscle TG content or decreasing production of cytokines associated with insulin resistance pathways (Wang *et al.*, 2001), counteracting hypertriglyceridaemia by increasing hepatic oxidation of fatty acids and reduced synthesis and secretion of TG (Staels *et al.*, 1998) have all been postulated as possible mechanisms. Similarly, the mechanism of PPAR- α -induced reduction in weight gain also remains controversial. There is a suggestion that a reduction in fat mass and consequential decrease in leptin production may ultimately induce hypophagia and the resultant decrease in weight gain (Kubota *et al.*, 1999), while another study reported lack of any association between plasma levels of leptin, food intake and inhibition of weight gain by PPAR- α agonist, fenofibrate treatment (Guerre-Millo *et al.*, 2000).

The present study and other reports have shown that dietary obesity not only causes metabolic abnormalities, it also markedly affects vascular function, in particular endothelium-dependent agonist-induced vasorelaxation (Steinberg *et al.*, 1996; Naderali *et al.*, 2001a; Naderali *et al.*, 2001c). The vascular abnormalities induced by dietary obesity are complex and multifactorial. Possible factors include fat mass *per se*, insulin resistance (Reaven, 1988), increased in NEFA levels (Steinberg *et al.*, 1996) hypertriglyceridaemia (Naderali *et al.*, 2001a) decrease in plasma HLD-c levels (Despr es *et al.*, 2002), and hyperleptinaemia (Wallace *et al.*, 2001). The key players could, in theory be identified through interventions that aim to correct these abnormalities and findings which leads to the amelioration of vascular dysfunction.

In this study, in agreement with previous reports (Furuhashi *et al.*, 2002) 8 weeks treatment of fenofibrate reduced total body weight, weekly weight gain and adiposity. Moreover, fenofibrate-treatment completely corrected dietary-induced insulin-resistance, hyperinsulinaemia, and hypertriglyceridaemia but not

hyperleptinaemia. By contrast, the beneficial effects of fenofibrate on endothelial function were relatively minor and only appeared at higher concentration of carbamylcholine (>31.6 μ M CCh). The failure of fenofibrate to completely correct dietary-induced endothelial dysfunction seen in this study is in agreement with previous reports. Studies by others have not shown any significant changes in endothelial function after acute treatment with 250 mg fenofibrate (Bae *et al.*, 2001), nor were there any beneficial effects on blood pressure or progression of atherosclerosis process (Wilmink *et al.*, 2001; Furuhashi *et al.*, 2002), despite improving insulin sensitivity and hypertriglyceridaemia. However, others have shown significant improvement of endothelial function after fenofibrate-treatment in human subjects (Idzior-Walus *et al.*, 2000; Malik *et al.*, 2001; Capell *et al.*, 2003). These contrasting findings could be explained by differences in experimental conditions. For example, subjects in Idzior-Walus and colleagues' (Idzior-Walus *et al.*, 2000) study had been under lipid management for at least 12 months, establishing a steady state of weight and lipid parameters and those in Malik and colleagues (Malik *et al.*, 2001) study did not include obese or insulin resistance subjects. Another recent study (Capell *et al.*, 2003) reported that fenofibrate treatment (2 weeks) significantly reduced fasting TG levels and improved both endothelium-dependent and -independent vasorelaxation. Once again, these patients underwent specific dieting for at least 2 weeks before any vascular analyses were carried out. Taken together, these observations, it is possible to hypothesise that withdrawal of a fat-enriched diet attenuated vascular dysfunction in these subjects and that this may have further improved by administration of fenofibrate, as seen in the present study. Consistent with an effect of diet *per se*, it has been reported recently that acute withdrawal of an obesity-inducing diet partially restores normal endothelial function (Naderali and Williams, 2003). While, data in previous chapter (chapter 5) have demonstrated that chronic withdrawal completely corrects dietary-obesity-induced vascular dysfunction, indicating that diet *per se* rather than dietary-induced obesity plays a key role in vascular abnormalities. However, it is also possible to argue that hyperleptinaemia (Wallace *et al.*, 2001) seen in diet-fed rats (both vehicle- and fenofibrate-treated rats) may at

least in part, explain the limited beneficial effects of fenofibrate on vascular function.

The present study, identifies for the first time the possible causative factors and mechanisms which might be involved in "the limited" beneficial effects of fenofibrate treatment on the vascular abnormalities caused by diet-induced obesity (Fig. 6.10). In this study, fenofibrate was administered in the presence of high palatable diet feeding, and was associated with elevated levels of plasma CRP and leptin. A recent study has demonstrated that hyperleptinaemia is associated with increased CRP, and that a significant correlation between plasma leptin and CRP is evident, suggesting involvement of leptin and CRP in inflammatory and cardiovascular disease processes (Shamsuzzaman *et al.*, 2004). Various studies reported that highly-palatable diet enhances leptin production well before a significant weight increase (Harrold *et al.*, 2000; Archer *et al.*, 2005). Similarly, in the present study increase in plasma leptin could be attributed to diet-induced obesity and not to fenofibrate-treatment, since fenofibrate administration (in the absence of high palatable diet feeding) induces significant reduction in plasma levels of leptin (Damci *et al.*, 2003).

This study also hypothesises that the increased concentration of plasma CRP seen in fenofibrate treated group might be induced by hyperleptinaemia. In fact physiological concentrations of leptin stimulates expression of CRP in human primary hepatocytes (Chen *et al.*, 2006). Various studies have demonstrated that CRP not only has pro-inflammatory effects (Leinonen *et al.*, 2003; Lin *et al.*, 2004; Nesto, 2004; Garanty-Bogacka *et al.*, 2005), it also causes insulin resistance, endothelial dysfunction and involved in the pathogenesis of atherosclerosis (Lin *et al.*, 2004). Incubation of cultured human coronary artery smooth muscle cells with CRP resulted in reactive oxygen species (ROS) generation, suggesting that CRP might play a crucial role in plaque instability and in the pathogenesis of acute coronary syndrome (Kobayashi *et al.*, 2003). Therefore, the vascular abnormalities seen in this study could be caused by the elevated levels of plasma CRP *via* its pro-inflammatory, atherogenic, and pro-oxidative effects.

Leptin affects vascular system, promoting cardiovascular disease (Wallace *et al.*, 2001). Leptin increases CRP production which, in turn, leads to induction of leptin-resistance (Chen *et al.*, 2006), and subsequent cardiac hypertrophy (Barouch *et al.*, 2003). Similarly, in this study diet-induced obese animals had higher leptin levels, reduced vascular function, and increased cardiac mass, suggesting involvement of leptin-CRP axis in cardiovascular pathology. Both leptin and CRP have direct and indirect detrimental effects on vascular function. In the presence of highly-palatable diet, production of leptin and CRP continued such that beneficial effects of fenofibrate failed to counter their negative effects on vasculature.

Fenofibrate protects vascular function and improves insulin resistance (Badiou *et al.*, 2004; Koh *et al.*, 2005), by lowering lipid profile (Damci *et al.*, 2003; Empen *et al.*, 2003; Badiou *et al.*, 2004), total body weight and adiposity (Filippatos *et al.*, 2005; Ji *et al.*, 2005). This in turn would result in reduction of adipocytokines such as TNF- α (Skoog *et al.*, 2002 ; Yamaguchi *et al.*, 2003; Rask-Madsen *et al.*, 2003 ; Matsuda and Shimomura, 2004; Recasens *et al.*, 2004). In the present study, fenofibrate reduced plasma concentration of TNF α to levels similar to chow-fed groups, indicating anti-inflammatory effects of fenofibrate (Lin *et al.*, 2004). However, despite complete improvement of systemic insulin resistance, fenofibrate only partly restored obesity-induced vascular abnormalities (Fig. 6.10).

In conclusion, this study indicates that fenofibrate treatment in the presence of obesity-inducing diet, decreases adiposity, lowers plasma TG and NEFA levels and improves insulin resistance in dietary-obese animals, but has little effects on vascular function. The lack of marked beneficial effects of fenofibrate on vascular function may, at least in part, be due to the continuous presence of detrimental factors in obesity-inducing diet, of which high sucrose and/or fats may play important roles. In particular, the diet-induced elevated levels of CRP might play a crucial role in the development of vascular abnormalities. Furthermore, findings from this study, indicate that pharmacological attenuation of adiposity alone does not alleviate obesity-induced endothelial dysfunction, and that removal of obesity-inducing diet (as shown in chapter five) is an important step in correcting vascular dysfunction.

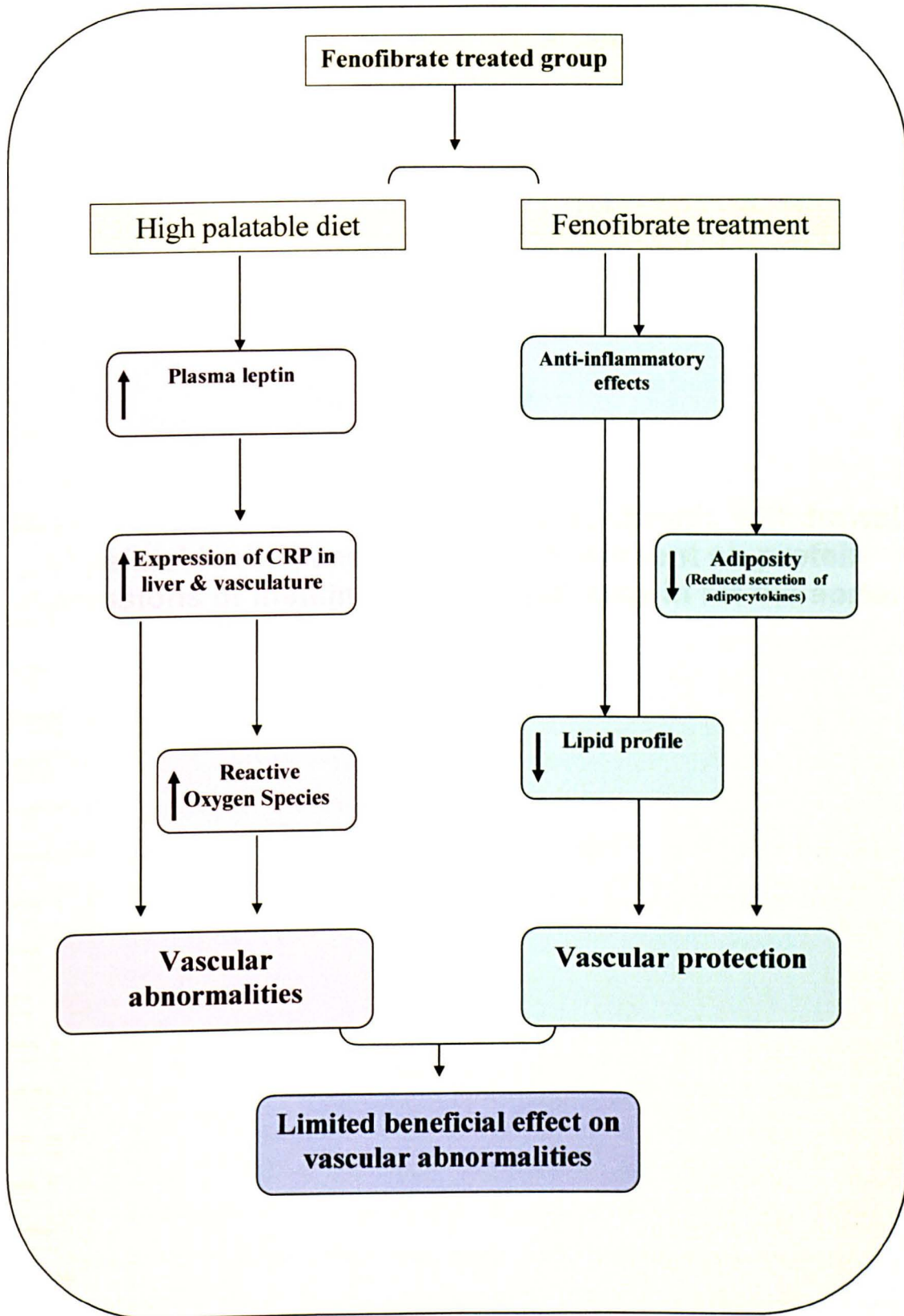


Fig. 6.10. Analysis for the vascular effects of fenofibrate treatment in diet-induced obesity “in the present study”.

Chapter 7

Results: the effects of dietary obesity, chronic withdrawal of palatable diet, and fenofibrate treatment on protein expressions of insulin signalling pathway in rat the aorta.

7.1 INTRODUCTION

Obesity, characterised by excess adipose tissue is now becoming a worldwide epidemic (Staels *et al.*, 1998; James *et al.*, 2001). Various studies have suggested that obesity *per se* is an independent cardiovascular risk factor (Duflou *et al.*, 1995), as well as predisposing to type 2 diabetes, hypertension and dyslipidaemia (Reaven, 1988). Furthermore, obesity induces insulin resistance which is associated with development of cardiovascular diseases that include hypertension (Reaven, 1993), and reduced endothelial function (Steinberg, 1996). Insulin has a protective role in vascular function. It stimulates nitric oxide (Mancini *et al.*, 2001) production, leading to vasorelaxation. Insulin-induced NO-dependent vasorelaxation is markedly decreased in obesity (Walker *et al.*, 1997), however, the mechanism(s) of decrease in insulin-induced vasorelaxation is not fully understood. Insulin has other major physiological roles that include facilitation and increase of amino acid transport, glycogen synthesis, DNA synthesis and gene expression (Obata, 1996). Moreover, it specifically enhances release of nitric oxide (Zeng and Quon M, 1996; Mancini *et al.*, 2001), regulates mRNA matrix proteins (Tamaroglio and Lo, 1994) and constitutive endothelial NO synthase (Kuboki, 1998) activity in vasculature.

In vascular cells, the effects of insulin are initiated through binding to the insulin receptor alpha subunit (IR- α), which activates the intrinsic receptor tyrosine kinase (Kasuga *et al.*, 1982), resulting in autophosphorylation of insulin receptor beta subunit and tyrosine phosphorylation of intracellular adaptor proteins - insulin receptor substrates (IRS-1 and IRS-2) (Sun, 1991) and Shc (Skolink, 1993). Tyrosine phosphorylated IRS-1 or IRS-2 then binds to src-homology 2 (SH2) domains of intracellular proteins, including the p85 regulatory subunit of phosphatidylinositol (PI) 3-kinase (Myers and Jr., 1992). The interactions of IRS and p85 subunit of PI 3-kinase results in the activation of p110 catalytic subunit of PI 3-kinase. Activation of PI 3-kinase increases serine phosphorylation of Akt, which in turn, directly phosphorylates eNOS on serine 1177 and activates the enzyme, leading to increased NO production and thus providing vascular protection (Dimmeler *et al.*, 1999). Tyrosine phosphorylated Shc and IRS proteins

can also bind to SH2 domain of Growth factor Receptor-protein Bound 2 (GRB2), leading to the activation of the Ras-Raf-MAP kinase signal pathway that is associated with gene expression and cell growth (Skolink, 1993). PI 3-kinase expression and activation has been linked to NO production, whereas, Raf-MAP kinase pathway is associated with the growth of vascular cells and the expression of extracellular matrix proteins (Zeng and Quon M, 1996). Furthermore, various studies (Tamaroglio and Lo, 1994; Obata, 1996; Zeng and Quon M, 1996; Walker *et al.*, 1997; Kuboki, 1998; Mancini *et al.*, 2001) have shown attenuation of insulin-induced vasorelaxation in obesity, however, there is little information on the mechanism(s) of diet-induced changes in insulin signalling pathway in vasculature. Therefore, it is possible that changes in any component of insulin signalling pathway may alter vasorelaxant property of insulin. Furthermore, the level of protein expression of these kinases could fluctuate and respond differently to any pathological, physiological, or pharmacological conditions, such as dietary obesity and its treatment. Therefore, investigating the protein expression of these kinases under certain obesity-related experimental conditions could provide vital information about obesity-induced insulin resistance in vascular tissues. Consequently, this study aimed to investigate the protein expression of insulin signalling components in aorta of four different experimental groups, namely, 1) control chow-fed lean animals, 2) untreated diet-induced obese animals, 3) obese animals following chronic withdrawn of palatable diet, and 4) obese animals treated with fenofibrate. Fenofibrate has been shown to improve lipid profiles, reduce adiposity, and may have a role in correction of vascular function in obesity.

7.2 METHODS

7.2.1 Experimental protocol

Adult male Wistar rats ($n = 28$, 190 ± 3 g) were randomised and assigned to a control group ($n = 10$) and a test group ($n = 20$). Controls were fed standard laboratory pelleted diet (Chow-fed), while test group had free access to a highly-palatable diet. Chow-fed controls remained on their prospective diet for 15 weeks, while after 8 weeks palatable-diet-fed animals were subdivided into three subgroups (each group 7 animals). In the first subgroup, palatable diet was removed and the standard chow diet was re-introduced (diet-to-chow), while the

second subgroup remain on palatable diet and were given fenofibrate (fenofibrate-treated, 50 mg/kg/day) by oral gavage for further 7 weeks and the third subgroup (diet-fed) was given vehicle (1% carboxymethyl cellulose at 1 ml/kg body weight; Sigma, UK), by oral gavage daily for 7 weeks. On the day of experiment (after 15 weeks), the rats were killed by CO₂ inhalation after 2 hours of fasting, the animal body was weighed, and the aorta was dissected and snap frozen in liquid nitrogen.

7.2.2 Molecular analysis of Aorta

Western-immunoblotting technique was then carried out (as described in chapter 2) on the four experimental groups (chow-fed, vehicle, diet-to-chow, and fenofibrate treated groups) for measuring protein expressions of insulin signaling pathway in the aorta (IR- β , IRS-1, IRS-2, PI 3-kinase, AKt, eNOS, SHC, and ERK1/2). The method which described in the previous paper by Iartey et.al. (2006) retrospectively should have been used for comparing the blotting data statistically.

7.2.3 Data interpretation and statistical analysis

Changes in body weights of each group were collected weekly and are expressed as absolute total body weight, respectively. For Western blotting, the data from chow-fed (control) animals were expressed as 100% response, and the results from other three groups were normalized and subsequently expressed as the percentage of their respective controls. Data are expressed as mean \pm S.E.M. Data have normal distribution (Shapiro Wilk W test). Statistical significance was tested using Student 't-test' or repeated-measures (ANOVA; Bonferroni t-test) or the Mann-Whitney test, as appropriate. Results were considered statistically significant at the $p < 0.05$ levels.

7.3 RESULTS

7.3.1 Changes in total body weight

Animals given palatable diet progressively gained more weight than their chow-fed counterparts. Untreated diet-fed animals were significantly heavier than chow-fed (>21%; 695 \pm 17 versus 572 \pm 8 g, $p < 0.0001$); diet-to-chow (>11%; 695 \pm 17 versus 625 \pm 16 g, $p < 0.01$) and fenofibrate treated (>17%; 695 \pm 17 versus 593 \pm 17 g, $p < 0.001$) animals. However, at the end of the experiment, body weight of

diet-to-chow and fenofibrate treated animals were significantly ($p < 0.01$) lower than untreated diet-fed animals (Fig. 7.1).

7.3.2 Protein expression studies

IR- β . There were no significant differences in IR- β protein levels in aorta from chow-fed, untreated diet-fed and diet-to-chow animals, while aorta from fenofibrate-treated animals had significantly (up to 30%, $p < 0.001$) higher IR- β protein levels than chow-fed group (Fig. 7.2).

IRS-1. and IRS-2. As with IR- β , the IRS-1 and IRS-2 levels in chow-fed, untreated diet-fed and diet-to-chow fed groups were comparable, while aorta from fenofibrate-treated animals had significantly higher levels of both IRS-1 (26%, $p < 0.05$) and IRS-2 (25 %, $p < 0.05$) compared to chow-fed group (Fig. 7.3 & 7.4).

PI 3-kinase. Compared with chow-fed animals, there was a marked reduction in PI 3-kinase levels in untreated diet-fed animals (59%, $p < 0.001$). PI 3-kinase levels were further reduced by fenofibrate-treatment (92%, $p < 0.0001$), while removal of palatable diet completely reversed the reduction in PI 3-kinase levels seen in diet-fed group (Fig. 7.5).

Akt. The protein levels of Akt were similar in chow-fed, untreated diet-fed and fenofibrate-treated animals, while aorta from diet-to-chow group had significantly lower (13%, $p < 0.01$) levels of Akt protein (Fig. 7.6).

eNOS. There was a significant (25%, $p < 0.001$) elevation of eNOS levels in aorta from untreated diet-fed animals compared with that of chow-fed control, while removal of diet or fenofibrate-treatment markedly reduced the elevation of eNOS protein concentration seen in untreated diet-fed animals (Fig. 7.7).

Shc. Although, Shc protein levels were not significantly altered by diet feeding but removal of the diet or fenofibrate-treatment significantly (by up to 20%, $p < 0.05$) attenuated Shc levels in comparison to that of chow-fed controls (Fig. 7.8).

ERK1/2. Immunoblotting analysis of ERK1/2 (MAP kinase pathway) protein in all four groups was remarkably similar to each other. Palatable-diet feeding in the presence or absence of fenofibrate had no effect on ERK1/2 levels nor did the removal of the palatable diet altered ERK1/2 levels in aorta (Fig. 7.9).

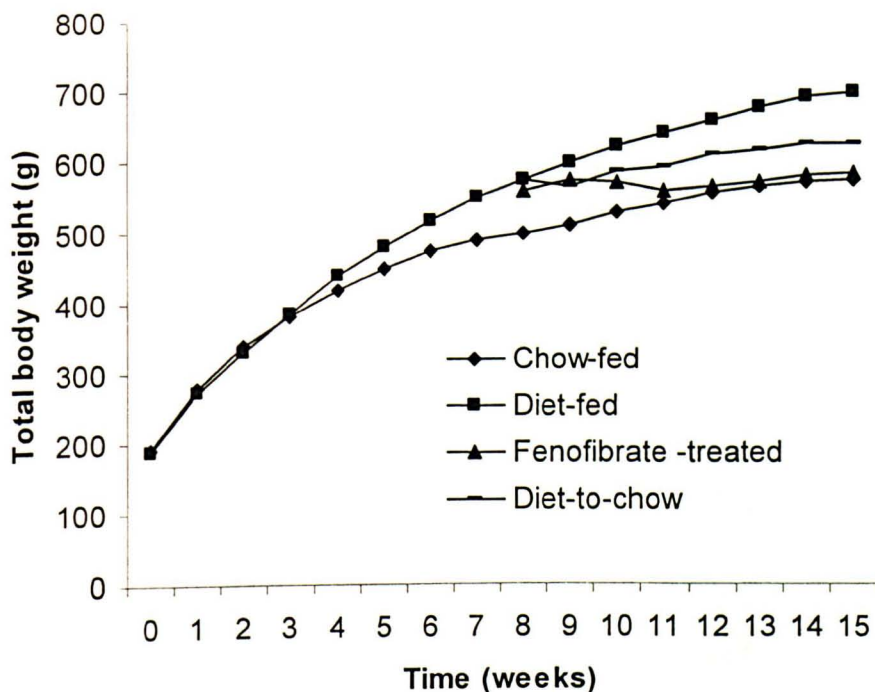


Fig. 7.1. The effects of palatable-diet, its removal, and fenofibrate treatment on total body weight in the rat. Animals were fed either standard chow (chow-fed) or a palatable diet (diet-fed) for 15 weeks. Fenofibrate treated group was fed palatable diet for 15 weeks and received fenofibrate (50 mg/kg/day) for the last 7 weeks, while diet-to-chow group was fed palatable diet for the first 8 weeks and chow for the last 7 weeks of the experiment. At the end of the experiment (15 weeks) diet-fed animals weighed significantly greater than the chow-fed, fenofibrate-treated, and diet-to-chow groups ($p < 0.0002$, 0.002 , and 0.02 respectively). The body weight of diet-to-chow group was significantly greater ($p < 0.02$) than the chow-fed group, while that of the fenofibrate-treated group was not significantly ($p < 0.66$) different from the chow-fed group. Data are expressed as mean \pm S.E.M.

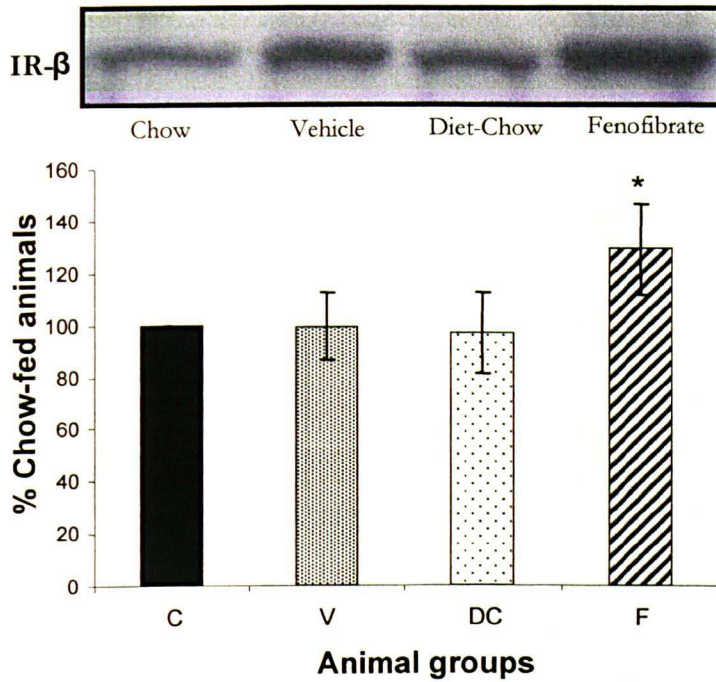


Fig. 7.2. Protein expression of IR- β in the rat aorta. Equal amounts (40 μ g / well) of protein were separated by SDS-PAGE and immunoblotted with IR- β antibody. The animal groups were: C) chow-fed lean control, V) Vehicle group (untreated dietary obese), DC) Diet-to-Chow group, and F) fenofibrate treated group. Data (mean \pm S.E.M; n=7) are normalised to that of control group and expressed as a percentage of the chow fed (control) rats. * $p < 0.05$ vs chow fed.

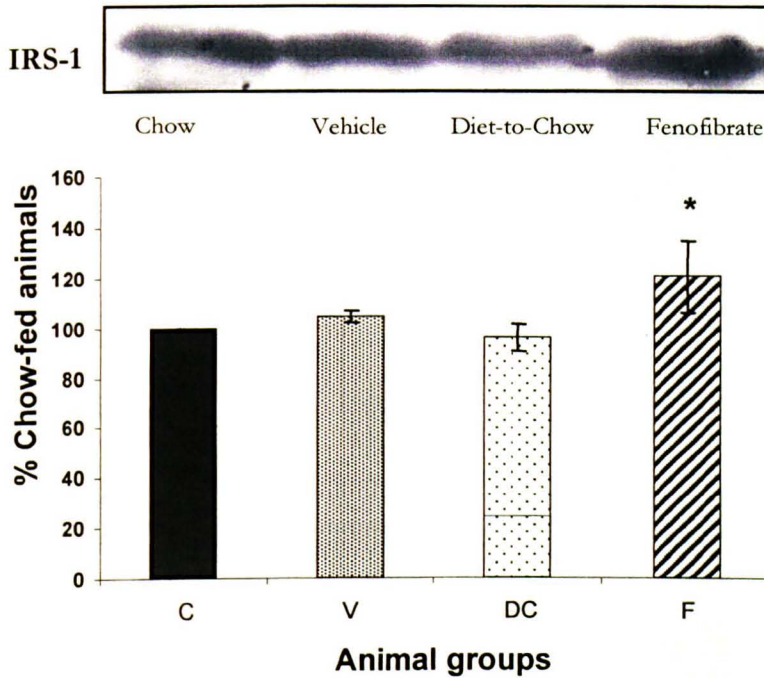


Fig. 7.3. Protein expression of IRS-1 in the rat aorta. Equal amounts (40 μg / well) of protein were separated by SDS-PAGE and immunoblotted with IRS-1 antibody. The animal groups were: C) chow-fed lean control, V) Vehicle group (untreated dietary obese), DC) Diet-to-Chow group, and F) fenofibrate treated group. Data (mean \pm S.E.M; n=7) are normalised to that of control group and expressed as a percentage of the chow fed (control) rats. * $p < 0.05$ vs chow fed.

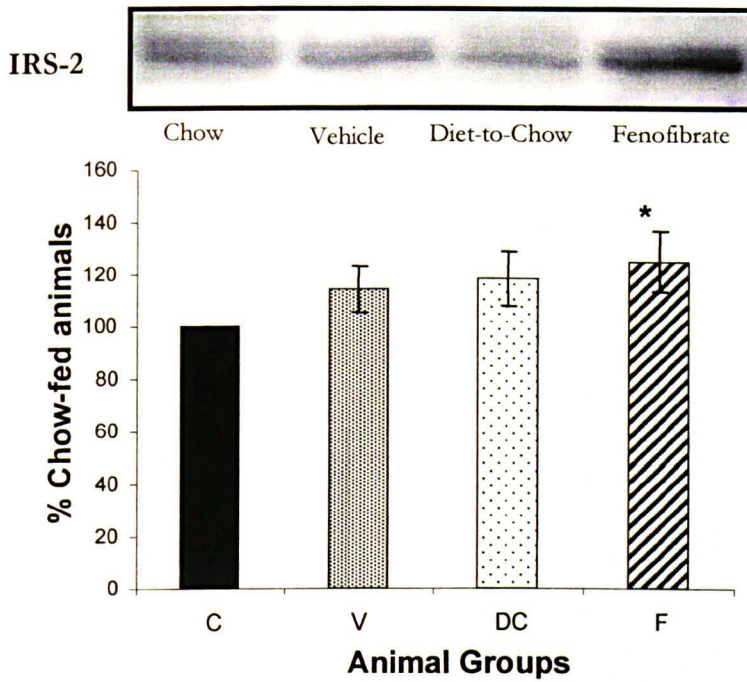


Fig. 7.4. Protein expression of IRS-2 in the rat aorta. Equal amounts (40 μ g / well) of protein were separated by SDS-PAGE and immunoblotted with IRS-2 antibody. The animal groups were: C) chow-fed lean control, V) Vehicle group (untreated dietary obese), DC) Diet-to-Chow group, and F) fenofibrate treated group. Data (mean \pm S.E.M; n=7) are normalised to that of control group and expressed as a percentage of the chow fed (control) rats. * $p < 0.05$ vs chow fed.

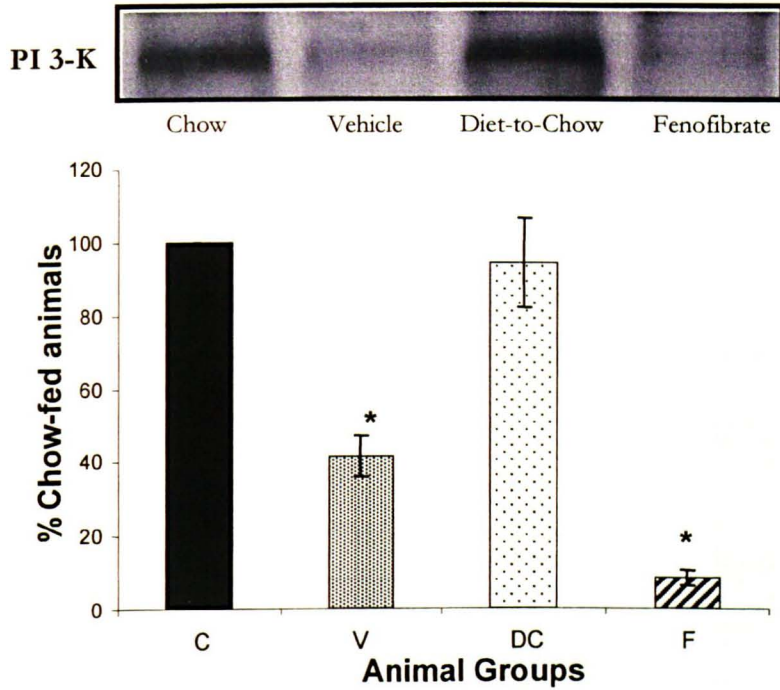


Fig. 7.5. Protein expression of PI 3-K in the rat aorta. Equal amounts (40 μg / well) of protein were separated by SDS-PAGE and immunoblotted with PI 3-kinase antibody. The animal groups were: C) chow-fed lean control, V) Vehicle group (untreated dietary obese), DC) Diet-to-Chow group, and F) fenofibrate treated group. Data (mean \pm S.E.M; n=7) are normalised to that of control group and expressed as a percentage of the chow fed (control) rats. * $p < 0.0001$ vs chow fed.

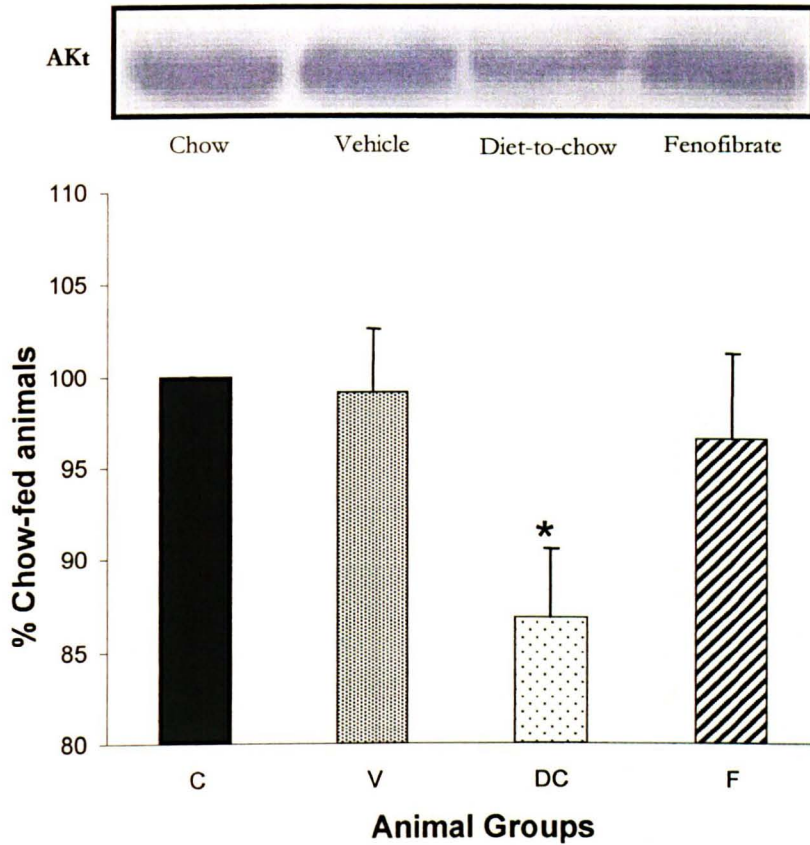


Fig. 7.6. Protein expression of Akt in the rat aorta. Equal amounts (40 μ g / well) of protein were separated by SDS-PAGE and immunoblotted with Akt antibody. The animal groups were: C) chow-fed lean control, V) Vehicle group (untreated dietary obese), DC) Diet-to-Chow group, and F) fenofibrate treated group. Data (mean \pm S.E.M; n=7) are normalised to that of control group and expressed as a percentage of the chow fed (control) rats. * $p < 0.05$ vs chow fed.

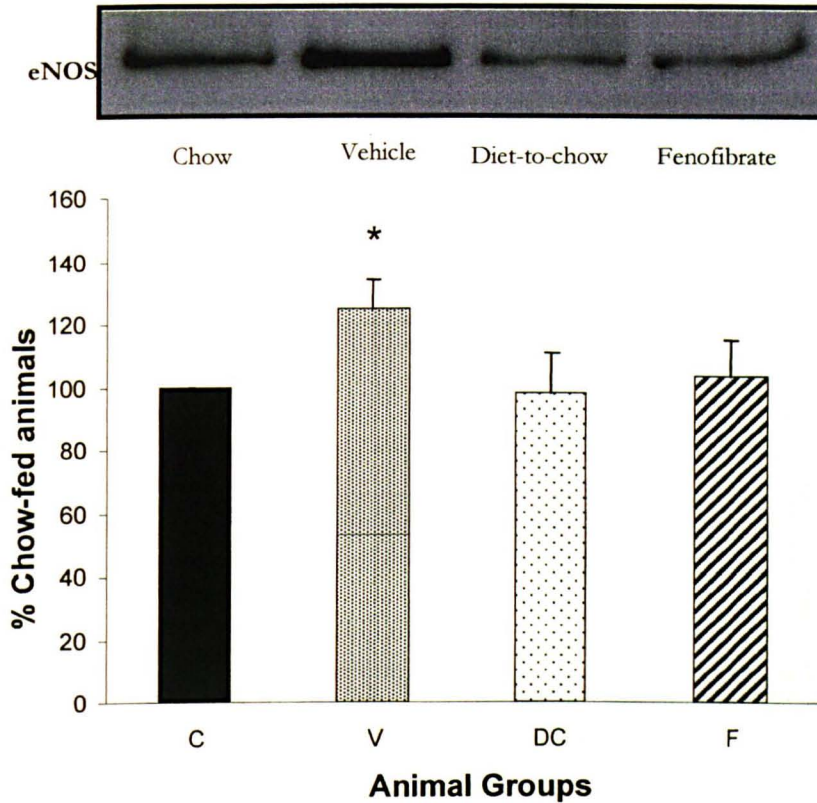


Fig. 7.7. Protein expression of eNOS in the rat aorta. Equal amounts (40 μg / well) of protein were separated by SDS-PAGE and immunoblotted with eNOS antibody. The animal groups were: C) chow-fed lean control, V) Vehicle group (untreated dietary obese), DC) Diet-to-Chow group, and F) fenofibrate treated group. Data (mean \pm S.E.M; n=7) are normalised to that of control group and expressed as a percentage of the chow fed (control) rats. * $p < 0.05$ vs chow fed.

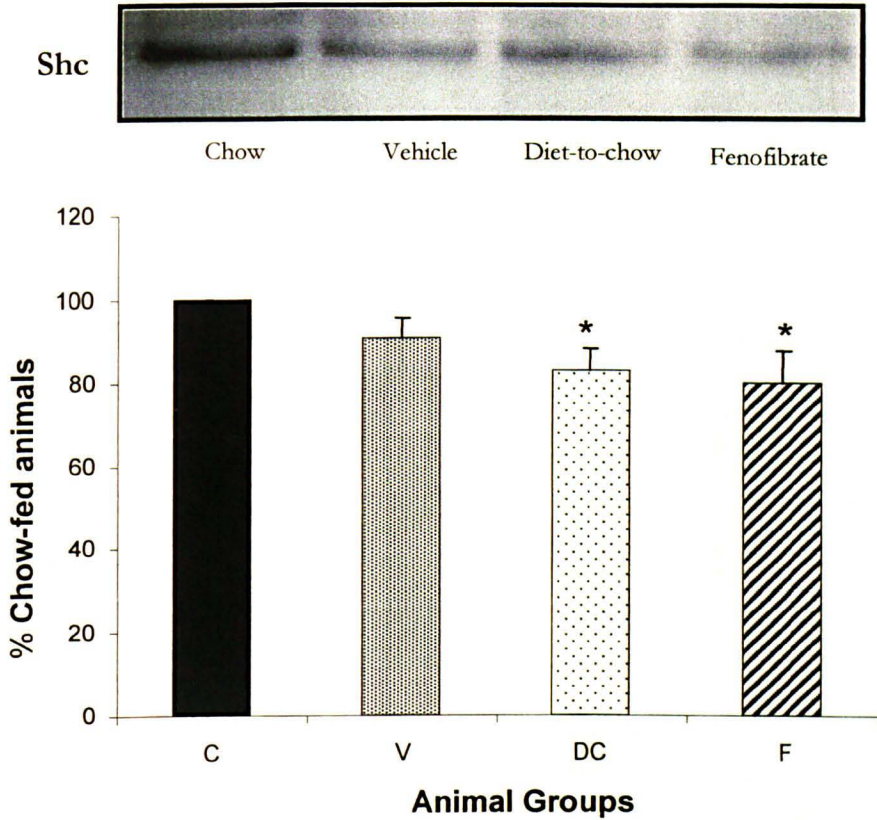


Fig. 7.8. Protein expression of Shc in the rat aorta. Equal amounts (40 μg / well) of protein were separated by SDS-PAGE and immunoblotted with Shc antibody. The animal groups were: C) chow-fed lean control, V) Vehicle group (untreated dietary obese), DC) Diet-to-Chow group, and F) fenofibrate treated group. Data (mean \pm S.E.M; $n=7$) are normalised to that of control group and expressed as a percentage of the chow fed (control) rats. * $p < 0.05$ vs chow fed.

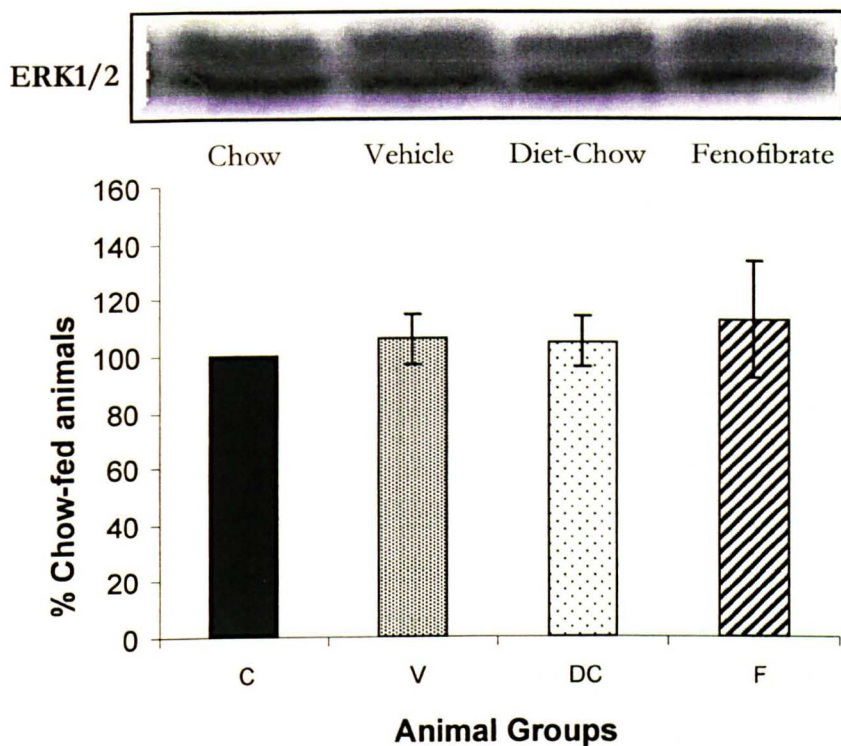


Fig. 7.9. Protein expression of ERK1/2 in the rat aorta. Equal amounts (40 μg / well) of protein were separated by SDS-PAGE and immunoblotted with PERK1/2 antibody. The animal groups were: C) chow-fed lean control, V) Vehicle group (untreated dietary obese), DC) Diet-to-Chow group, and F) fenofibrate treated group. Data (mean \pm S.E.M; n=7) are normalised to that of control group and expressed as a percentage of the chow fed (control) rats.

7.4 DISCUSSION

Interactions between insulin, IR- β , IRS-1, and IRS-2 activates PI 3-kinase pathway resulting in vasorelaxation. Therefore, the integrity of protein levels and activation of the cascade in insulin signaling pathway in vasculature is a crucial factor in mediating normal vascular functions. In the present study, dietary-obesity did not adversely alter the concentrations of IR- β , IRS-1, and IRS-2, suggesting that reduced vasorelaxation to insulin seen in dietary-obesity (Walker *et al.*, 1997; Naderali *et al.*, 2001a), may not be due to the changes in protein levels of IR- β , IRS-1, and IRS-2. This raises possibility that attenuated insulin responses seen in obesity (Walker *et al.*, 1997; Naderali *et al.*, 2001a) may be due to changes in protein levels beyond membrane receptors. In fact, the present study indicates a marked decrease in PI 3-kinase levels in aorta from dietary-obese rats, suggesting that dietary-obesity adversely affects PI3-kinase protein level and thereby reducing insulin-induced vasorelaxation. Studies on genetically obese animals have reported similar observation. Jiang and colleagues have shown reductions in IRS-1, and IRS-2 but not IRS- β protein levels in obese Zucker rats in comparison with lean animals (Jiang *et al.*, 1999). Moreover, the same study reported a marked inhibition of PI 3-kinase activation in aorta of Fatty-Zucker rats. Although, the present study did not measure insulin-induced phosphorylation of signalling components, however, low level of PI 3-kinase seen in this study together with reduced functionality of PI 3-kinase reported on Zucker rats (Jiang *et al.*, 1999) may play an important role in integrity of endothelial function. A similar finding has also been reported in human umbilical vein endothelial cells where inhibition of PI 3-kinase markedly attenuated insulin-stimulated NO production (Zeng *et al.*, 2000), further arguing for a significant role of PI 3-kinase-changes in endothelial dependent insulin-induced vasorelaxation.

Earlier experiments (chapter five) indicated that in obese animals, removal of obesity-inducing diet completely restores endothelial function, suggesting that reversal of adiposity might be of benefit in correcting obesity-induced attenuation of PI 3-kinase. In fact, in this experiment, PI 3-kinase levels were restored in diet-to-chow group, further strengthening the hypothesis that PI 3-kinase levels may

determine the magnitude of endothelial function in obesity. Therefore, it is possible that any deficiency in the concentrations of PI 3-kinase may participate in inducing insulin resistance in vascular system accompanied with reduced NO production, leading to endothelial dysfunction.

Peroxisome proliferator-activated receptor- α (PPAR- α) plays a crucial role in the control of mitochondrial β -oxidation of fatty acids (Pyorala *et al.*, 2000; Eringa *et al.*, 2004). PPAR- α expressed in the vascular tissue, mainly the endothelial cells (Marx *et al.*, 1999). Hence, it is plausible to assume that the increased intracellular fatty acids and the presence of abundant amount of (PPAR)- α agonist (fenofibrate) may enhance fatty acid oxidation leading to increase in the intracellular fatty acid metabolites. Fatty acid metabolites such as fatty acyl-CoA, diacylglycerols, and ceramides shown to alter insulin signaling and induce insulin resistance (Schmitzpeiffer, 2002) thus causing reduction in PI3 kinase levels in these animals. Therefore, it is reasonable to assume that further reduction of PI 3-kinase levels seen in fenofibrate-treated animals may be the consequence of increased fatty-acid oxidation in these animals. However, this hypothesis merits further investigation. Fenofibrate-treated group had deficient protein expression of PI 3-kinase in vasculature, "the key controlling step of insulin signalling pathway", possibly contributing to insulin resistance, and subsequent endothelial dysfunction (chapter six).

Functionally active PI 3-kinase stimulates Akt, which in turn activates eNOS, leading to production of NO and subsequent vasorelaxation (Zeng *et al.*, 2000). In this study, protein levels of Akt were similar between dietary-obese and lean aorta, while removal of palatable diet caused a marked decrease in Akt levels. Surprisingly, on the other hand, there was a significant increase of eNOS levels in the obese group, whereas removal of palatable diet or fenofibrate treatment restored eNOS levels to that of lean control group. Although, increased eNOS level in this study is suggestive of augmented endothelial-dependent vasorelaxation, but results in chapter five in addition to numerous human and animal studies have shown attenuated endothelial function in obesity (Steinberg, 1996; Naderali *et al.*, 2001; Naderali *et al.*, 2001a). There are several possibilities on mechanisms of

increased level of eNOS in obese animals. One possible explanation, is activation of a compensatory mechanism to overcome the decrease in NO production or increase inactivation of NO seen in arteries of dietary obese rats as a result of increased oxidative stress (Katakam *et al.*, 2005). Moreover, in obese animals there is an elevation of insulin concentration and endothelial dysfunction (Naderali *et al.*, 2004a). Furthermore, insulin is a potent enhancer of reactive oxygen species (ROS) synthesis in endothelial and vascular smooth muscle cells, and superoxide is known to reduce NO and subsequently increased vasoconstriction (Cai and Harrison, 2000; Griendling and FitzGerald, 2003), thus, the elevation of eNOS could compensate this reduction. Insulin is also shown to induce eNOS expression in endothelial cells (Aljada and Dandona, 2000; Mather *et al.*, 2001) and therefore, insulinaemia which observed in obese rats could directly be responsible for the enhancement of eNOS expression. Similar results and hypothesis have also been reported by others indicating an increase in eNOS levels in Zucker obese coronary (Katakam *et al.*, 2005) and cerebral (Erdos *et al.*, 2004) arteries. However, the function of eNOS maybe regulated by PI 3-kinase or MAP-kinase pathway (Park *et al.*, 2002), and thus, decrease in PI 3-kinase seen in untreated obese rats may cancel any beneficial effects of increased eNOS. Furthermore, activated vascular endothelial growth factor (VEGF) stimulates eNOS expression at both mRNA and protein levels in a dose-dependent manner (Park *et al.*, 2002), suggesting that elevated levels of eNOS in dietary-obese group seen in the present study maybe due to an increased level of VEGF through MAP-kinase pathway, correlating positively with the dietary-obese subjects (Miyazawa-Hoshimoto *et al.*, 2003). Therefore, restoration (i.e. reduction) of total body fat seen in diet-to-chow and fenofibrate-treated animals, may have resulted in reduced effect of VEGF on eNOS expression (Miyazawa- Hoshimoto *et al.*, 2003) and thus improved vascular function (Naderali *et al.*, 2004a; Naderali *et al.*, 2004b). Inhibition of PI 3-kinase pathway enhances the mitogenic actions of insulin through MAP kinase pathway (Montagnani *et al.*, 2002). Hence, it is possible that inhibition of PI 3-kinase pathway seen in both untreated dietary-obese and fenofibrate treated groups may cause over activation of other signalling pathway such as MAP kinase. This result is compatible with the result in chapter five where endothelial

dysfunction in untreated-diet fed group was evident, indicating that deficient PI 3-kinase pathway in this group and over-activated MAP kinase pathway may result in endothelial dysfunction.

In conclusion, the present study demonstrates for the first time that long term feeding (15 weeks) of animals with an obesity-inducing palatable diet causes selective changes in protein levels of PI 3-kinase-dependent signalling pathway in aorta. Chronic withdrawal of obesity-inducing diet causes a complete normalization of PI 3-kinase, while fenofibrate treatment failed to improve PI 3-kinase concentrations in dietary-obese animals. A reduction in PI 3-kinase levels may have a role in inducing insulin resistance in vasculature, contributing to increased incidence of cardiovascular events seen in obese subjects. Furthermore, it can be postulate that, the inhibition of PI 3-kinase pathway may result in increased activation of MAP kinase pathway leading to an increased proliferation and migration of endothelial cells, thereby increasing the risk of cardiovascular events. However, this hypothesis merits further investigation. Furthermore, the adverse effects of dietary obesity on insulin transduction in vasculature are post receptor, on the level of PI3-kinase and downstream, and reversible, mainly by removal of obesity-inducing diet, which may have a role in combating diet-induced obesity-related cardiovascular dysfunction

Chapter 8

**Results: effects of diet-induced obesity on liver:
alterations in insulin signaling pathway and formation of
non-alcoholic fatty liver**

8.1 INTRODUCTION

Insulin is the primary anabolic hormone promoting the storage of energy in the fed state. In response to nutrient secretagogues, insulin is secreted from pancreatic β cells directly into the portal circulation. The resulting high portal insulin levels prime the liver for rapid alterations in hepatic carbohydrate and lipid homeostasis, such as stimulation of glycogen synthesis, lipogenesis, lipoprotein synthesis and suppression of gluconeogenesis / glycogenolysis and VLDL secretion (Cherrington *et al.*, 1978; Sparks and Sparks, 1994; Zammit, 2000). In fact, the liver is also a major depot for glucose uptake and storage, and may account for disposal of up to one third of an oral glucose load (pagliassotti and Horton, 1994).

Obesity has been associated with glucose intolerance, insulin resistance, dyslipidemia, and cardiovascular disease (Evans *et al.*, 1984; Golay *et al.*, 1990; DeFronzo and Ferrannini, 1991). The liver has specifically been implicated as a primary site of insulin resistance observed with visceral obesity. The mechanism(s) relating visceral fat accumulation and hepatic insulin resistance are not well understood yet, but several possible factors have been implicated. For example, increased release of adipokines such as TNF- α , resistin, and adiponectin have been implicated in the development of hepatic insulin resistance (Hotamisligil, 1999; Stepan *et al.*, 2001). Alternatively "portal hypothesis" posits a high rate of lipolysis of visceral adipose tissue leading to increased delivery of free fatty acids to the liver *via* portal vein, thus contributing to increased fat accumulation and liver insulin resistance (Bjorntorp, 1990; Boden, 1997; Bergman and Ader, 2000).

Chronic exposure of the liver to elevated FFA can promote liver gluconeogenesis (Williamson, 1966), deplete enzymes involved in FFA oxidation, and increase hepatic lipogenesis (Xu *et al.*, 1999; Clarke, 2000). In addition, elevated FFA are known to increase liver triglyceride content (Oakes *et al.*, 1997), and decrease insulin clearance (Mittelman *et al.*, 2000), both of which are associated with insulin resistance. Thus, insulin signalling in liver is critical in regulating glucose homeostasis and maintaining normal hepatic function and insulin resistance in hepatocytes leads to sever systemic insulin resistance and progressive hepatic

dysfunction (Michael *et al.*, 2000). It is therefore, possible that defective levels in protein expression of insulin signalling components in liver and/or inhibition of their activation will participate in insulin resistance.

A complex interaction of insulin resistance, deranged lipid profile, and vascular dysfunction is known as metabolic syndrome X (Zeman *et al.*, 2005). Patients with metabolic syndrome develop non-alcoholic fatty liver disease (NAFLD). As obesity is closely associated with NAFLD (Marchesini and Babini, 2006), it is expected that its incidence and prevalence would become more common. NAFLD is characterized by fatty liver and hepatitis, thus it is a spectrum of disorders that range from simple hepatic steatosis to steatohepatitis, cirrhosis, and hepatocellular carcinoma (Ludwig *et al.*, 1980; Luyckx, 2000). Various studies have pointed to hyperinsulinaemia and insulin resistance as pathogenic factors in NAFLD. Using the homeostasis model assessment (HOMA) method (Matthews *et al.*, 1985), Marchesini *et al.* (Marchesini *et al.*, 1999) shows that insulin resistance was the laboratory finding most closely associated with the presence of NAFLD in large series of patients, irrespective of BMI, fat distribution, or glucose tolerance. The strong association of NAFLD with other features of the metabolic syndrome (obesity, central fat, distribution, diabetes, dyslipidaemia, hypertension, and atherosclerotic cardiovascular disease) supports the hypothesis (Cortez-Pinto *et al.*, 1999) that NAFLD might represent another feature of the metabolic syndrome, with decreased insulin sensitivity being the common factor (DeFronzo and Ferrannini, 1991).

The pathogenesis of NAFLD is not fully understood, and the 'the two-hit' hypothesis proposed by Day *et al.* remains the prevailing theory (Day and James, 1998).

First hits: insulin resistance and fatty acid accumulation. It is believed that, resistance leads to the accumulation of triglycerides in hepatocytes. As more fatty acids being synthesised, more free fatty acids being delivered to the liver, less fatty acids being degraded, and less triglycerides being released from the liver. Lipid accumulation in hepatocytes (steatosis) is a characteristic histological feature of

NAFLD. This link is supported by findings that many patients with NAFLD have hyperinsulinaemia, insulin resistance, and the metabolic syndrome. (Marchesini, 2001; Chitturi *et al.*, 2002). There are two major mechanisms of fatty acid toxicity in the pathogenesis of NAFLD, namely the “direct” and “indirect” mechanisms. The “direct” mechanism involves direct toxicity such as membrane disruption at very high concentrations (detergent effects), apoptosis, and necrosis, while the “indirect” mechanism involves cytotoxic effects of lipid peroxidation of fatty acids.

Second hits: Oxidative stress, cytokines. A variety of second hits could account for the progression from simple steatosis to steatohepatitis. Oxidative stress occurs when more oxidant substances are produced than antioxidant processes of the liver can handle. Oxidative stress can cause lipid preoxidation, leading to activation of hepatic stellate cells and hepatocyte death contributing to hepatocellular injury and fibrosis. Sources of oxidative stress in steatohepatitis include reactive oxygen species that leak from the mitochondria during oxidation of fatty acids, cytochrome P450 enzymes, and hepatic iron (George *et al.*, 1998; Younossi *et al.*, 1999; Pessayre *et al.*, 2001).

Cytokine production is increased in nonalcoholic steatohepatitis and is believed to play a role in its pathogenesis. In the liver, TNF- α can contribute to oxidative stress (Tilg and Diehl, 2000), and may contribute to insulin resistance through activation of the inhibitor of Kappa kinase beta (IKK beta) (Yuan *et al.*, 2001). Increased free fatty acid levels, in addition to mediating insulin resistance and causing oxidative stress, can be directly hepatotoxic, leading to cellular injury (Neuschwander-Tetri and Caldwell, 2003).

Leptin: a possible third hit? It may promote insulin resistance, contributes both to oxidative stress and to enhanced secretion of inflammatory cytokines (Loffreda *et al.*, 1998; Chitturi, 2002) and may play a role in causing fibrosis.

Biopsy is the definitive test. Because the clinical and laboratory features of NAFLD are not specific and imaging studies cannot reliably distinguish nonalcoholic steatohepatitis from other forms of NAFLD, liver biopsy from human or samples of histological sectioning from animal models are the only ways to accurately

diagnose NAFLD, particularly steatohepatitis. Histological features include steatosis, lobular inflammation, ballooning degeneration, perisinusoidal fibrosis, (Younossi *et al.*, 1998; Brunt *et al.*, 1999; Neuschwander-Tetri and Caldwell, 2003), and hepatocellular carcinoma (Bugianesi *et al.*, 2002).

Treatment. There currently is no proven effective therapy for nonalcoholic steatohepatitis. Early efforts focused on modifying the associated conditions such as obesity, hypertriglyceridaemia, and diabetes mellitus. More recent efforts have targeted theoretical aspects of the pathogenesis, such as insulin resistance and oxidative stress.

a) Weight loss, particularly of 10% or more, can lead to improvement in hepatic steatosis (Nomura *et al.*, 1987). However, there has been very little histological follow-up in obese subjects who have lost weight. One study reported that two patients with nonalcoholic steatohepatitis who lost weight had significant histological improvement on follow-up liver biopsies (Eriksson *et al.*, 1986; Andersen *et al.*, 1991). A very rapid weight loss, however, may lead to increased portal inflammation and fibrosis (Andersen *et al.*, 1991; Luyckx *et al.*, 1998), while the optimum rate of weight loss is not clear, gradual loss of 10% of baseline weight seems to be a reasonable recommendation (Sanyal, 2002).

b) Lipid-lowering agents, have been evaluated in patients with nonalcoholic steatohepatitis. Patients did not appear to benefit from 12 months clofibrate (Laurin *et al.*, 1996). On the other hand, atorvastatin for 1 year produced significant improvement in ballooning degeneration, and inflammatory scores in NAFLD patients (Horlander and Kwo, 1997; Brunt *et al.*, 1999). While, there is very small number of studies in literature regarding the effect of fenofibrate treatment on the progression of NAFLD, many of these reports indicated an improvement in NAFLD, specifically steatohepatitis as result of fenofibrate administration (Shiri-Sverdlov *et al.*, 2006).

Therefore, in this experiment, the effects of dietary-induced obesity and physiological and pharmacological correction of obesity on liver insulin signalling pathway as well as histological features of the liver were investigated. This study

provides valuable information on the mechanism of obesity-induced hepatic insulin resistance.

8.2 METHODS

8.2.1 Experimental protocol

Animals used in this study were those described in chapter 7. At the end of experimental protocol (as described in chapter 7), rats were killed by CO₂ inhalation after 2 hours of fasting, and the liver was dissected and snap frozen in liquid nitrogen, for later molecular and histological techniques.

8.2.2 Experimental procedures

8.2.2.1 Western-immunoblotting technique

Western-immunoblotting technique was carried out (as described in chapter two) on the four experimental groups (chow-fed, vehicle, diet-to-chow, and fenofibrate treated groups) for measuring the levels of protein expressions for kinases such as: IR- β , IRS-1, IRS-2, PI 3-kinase, AKt, eNOS, Shc, and ERK1/2 in insulin signaling pathways in the liver.

8.2.2.2 Liver triglyceride content

Triglyceride content was assessed in liver samples of all experimental groups: chow-fed, diet-fed, diet-to-chow, and fenofibrate treated groups, using a procedure described in chapter two.

8.2.2.3 Liver histology

The tissue samples of liver were prepared for histological examinations using ordinary tissue processing and stained by Haematoxylin and Eosin staining protocol which all described in chapter two. Ten liver specimens (10 animals) of each experimental group were interpreted by a single pathologist who was blinded to all data of liver samples and the experimental groups under investigation. The Liver samples were analyzed microscopically for the presence or absence of the following histological features: steatosis, steatohepatitis, fibrosis and hepatocellular carcinoma. Histological features were interpreted according to the schema outlined by Brunt et al. (Brunt *et al.*, 1999). Briefly, steatosis was graded on a 3-point scale:

grade 1 = steatosis involving <33% of hepatocytes, grade 2=33-66%, grade 3>66%. Inflammation (steatohepatitis) was graded on a 4 point scale: grade 0=no or negligible inflammation, grade 1 = mild, grade 2 = moderate, grade 3 = severe. Fibrosis was staged on a 5 point scale: stage 0 = no fibrosis, stage 1 = zone 3 perisinusoidal/ perivenular fibrosis, stage 2 = zone 3 and periportal fibrosis, stage 3 = septal/bridging fibrosis, stage 4 = cirrhosis.

8.2.3 Data interpretation and statistical analysis

For Western blotting, the data from chow-fed (control) animals were expressed as 100% response, and the results from other three groups were normalized and subsequently expressed as the percentage of their respective controls. Data are expressed as mean \pm S.E.M. Data have normal distribution (Shapiro Wilk W test). Statistical significance was tested using Student 't-test' or repeated-measures (ANOVA; Bonferroni t-test) or the Mann-Whitney test, as appropriate. Results were considered statistically significant at the $p<0.05$. The histological data are presented as number (percentage), each sample represent 10%. There are ten histological samples (10 animals) in each experimental group.

8.3 RESULTS

8.3.1 Body weight

By referring to the results of changes in total body weight in chapter 7, animals given palatable diet progressively gained more weight than their chow-fed counterparts. Untreated diet-fed animals were significantly heavier than chow-fed, diet-to-chow, and fenofibrate treated animals. However, at the end of the experiment, body weight of diet-to-chow and fenofibrate treated animals were significantly ($p<0.01$) lower than untreated diet-fed animals.

8.3.2 Liver weight

Liver weight of untreated obese animals (vehicle) was elevated significantly ($p<0.01$, 1.22 folds) compared with chow-fed group (control). Replacing the highly-palatable diet with standard laboratory chow (diet-to-chow group) completely restored the liver weight to normal range, with no significant difference in comparison to chow-fed group. However, compared with chow-fed and vehicle

groups fenofibrate treatment, produced a marked increase in liver weight up to 2.13-folds (for both, $p < 0.01$) (Fig. 8.1).

8.3.3. Triglyceride content of liver

Compared with chow-fed group, untreated obese animals (vehicle) had significantly ($p < 0.05$) higher TG levels. Diet-to-chow animals had liver TG similar to that of chow-fed, while TG levels in fenofibrate treated group were significantly reduced ($p < 0.01$) compared with chow-fed group (Fig. 8.2).

8.3.4 Expression of protein kinases of insulin signalling pathways in hepatocytes

IR- β . Diet-induced obesity and chronic withdrawal of highly-palatable diet did not affect IR- β levels in hepatocytes with no significant difference compared with chow-fed group. However, fenofibrate treatment induced a significant reduction ($p < 0.001$) in the concentration of IR- β (Fig. 8.3).

IRS-1. Similar to IR- β , the levels of IRS-1 in both vehicle and diet-to-chow groups were not affected and there was no significant difference compared with chow-fed group, whereas, fenofibrate treatment produced marked decrease ($p < 0.001$) in the concentration of IRS-1 in hepatocytes (Fig. 8.4a).

IRS-2. Concentration of IRS-2 was significantly reduced ($p < 0.05$) in vehicle group, while its levels were completely restored to the level of chow-fed group. However, compared with chow-fed group, fenofibrate-treated group had significantly reduced ($p < 0.0005$) concentration of IRS-2 (Fig. 8.4b).

PI 3-kinase. Similar to IRS-2, there was a significant decrease ($p < 0.05$) in the levels of this protein kinase in vehicle group, while its levels were similar to chow-fed group in diet-to-chow group. fenofibrate treatment induced significant decrease ($p < 0.0005$) in PI 3-kinase levels (Fig. 8.5a).

AKt. Although, diet-induced obesity did not affect the levels of AKt, chronic withdrawal of the highly-palatable diet caused marked decrease in AKt protein

($p < 0.05$). Furthermore, fenofibrate treatment induced a significant reduction ($p < 0.000005$) for the levels of Akt (Fig. 8.5b).

eNOS. The expression of eNOS protein was not affected by the three experimental groups: vehicle, diet-to-chow, and fenofibrate treatment groups. eNOS levels were similar in all groups with no significant difference between the groups (Fig. 8.6).

Shc. The expression of Shc was not affected in both vehicle and diet-to-chow groups, with no significant difference compared with chow-fed group. However, fenofibrate treatment induced significant decrease ($p < 0.005$) in Shc levels compared with chow-fed group (Fig. 8.7a).

ERK1/2. Similar to Shc, the levels of ERK1/2 were not affected by both diet-induced obesity and chronic withdrawal of the highly-palatable diet. However, fenofibrate treatment caused significant reduction ($p < 0.01$) in Akt protein levels (Fig. 8.7b).

8.3.5. The histological features of the liver (non-alcoholic fatty liver disease)

The histological slides of liver tissue of each animal were examined for the presence or absence for the histological features of the different stages of non-alcoholic fatty liver disease (Figs. 8.8a, b, c, and d). Ten liver samples (10 animals) in each experimental group were analysed microscopically, and each sample represent 10% of the total percentage of each group.

Steatosis (accumulation of fat in the liver) was found in 50% (grade 1 of steatosis) of chow-fed control group, while untreated obese animals (vehicle group) having steatosis in 80% of the total samples (40% in grade 1, 30% in grade 2, and 10% in grad 3). Steatosis was present in 50% of the samples of diet-to-chow group (40% in grade 1, and 10% in grade 2). Fenofibrate-treated group having steatosis in only 10% of the total samples of this group (which was classified as grade 2) (Table 8.1).

Steatohepatitis was present in 40% of the samples in chow-fed control, whereas in vehicle group, 70% of the samples showed steatohepatitis. Diet-to-chow group

had 30% of the samples steatohepatitis, while in fenofibrate-treated group only 10% of the samples were positive for steatohepatitis. The positive samples in all groups were of grade one steatohepatitis (Table 8.2).

Fibrosis and hepatocellular carcinoma. The histological examination of samples in all experimental groups did not show signs for the presence of advanced features of NAFLD: fibrosis and hepatocellular carcinoma (Table 8.3). In comparison with chow-fed group, obese animals had 30% greater steatosis, while the percentage of samples with steatosis in diet-to-chow group was similar to that of diet-fed “control”. Furthermore, steatosis in fenofibrate treated group was 40% less than that of chow-fed group.

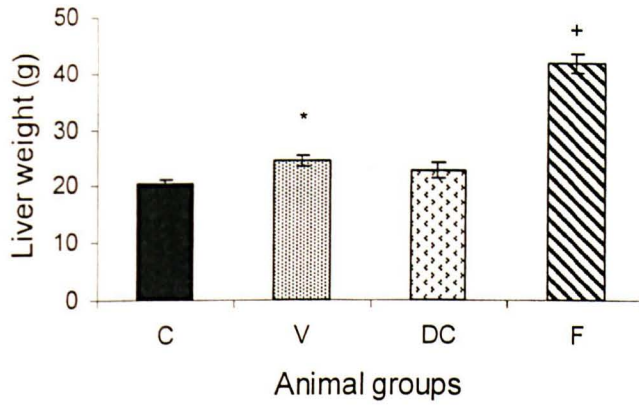


Fig. 8.1. Liver weight of animals at the end of the experiment in the four experimental groups: C) Chow-fed, V) Vehicle, DC) Diet-to-Chow, and F) Fenofibrate-treated groups. Data were represented as mean \pm S.E.M. There were significant increase in both vehicle and fenofibrate -treated groups compared with chow-fed group. * = p <0.01, += p <0.0001 vs chow-fed group.

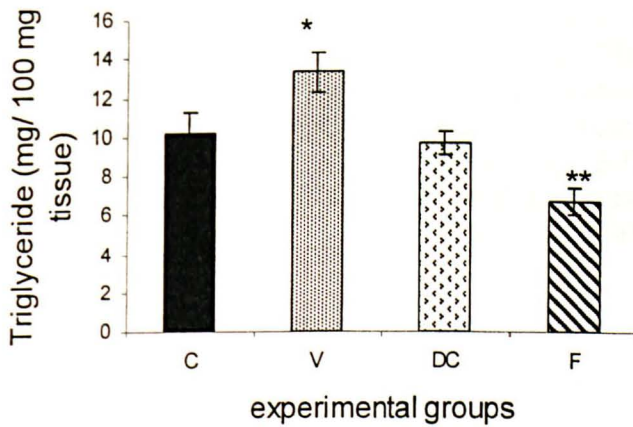


Fig. 8.2. Content of hepatic triglycerides in four experimental groups: C) Chow-fed, V) Vehicle, DC) Diet-to-Chow, and F) Fenofibrate-treated groups. Data are represented as mean \pm S.E.M. *= p <0.05, **= p <0.01 both vs. chow-fed group.

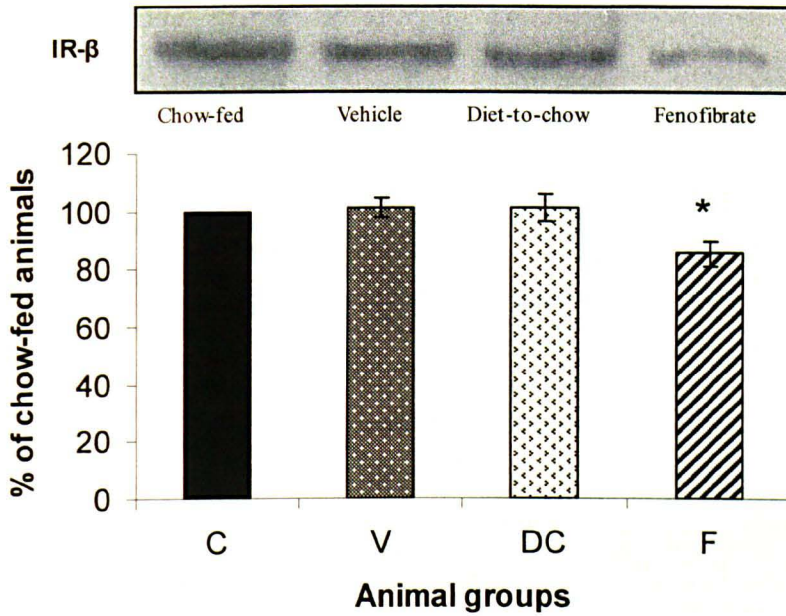


Fig. 8.3 Protein expression of IR- β in the rat liver. Equal amounts (50 μ g/well) of protein were separated by SDS-PAGE and immunoblotted with IR- β antibody. The animal groups are: C) chow-fed lean control, V) Vehicle group (untreated dietary obese), DC) Diet-to-Chow group, and F) fenofibrate treated group. Data (mean \pm S.E.M) are normalised to that of control group and expressed as a percentage of the chow fed (control) rats. * = p <0.00 vs chow fed.

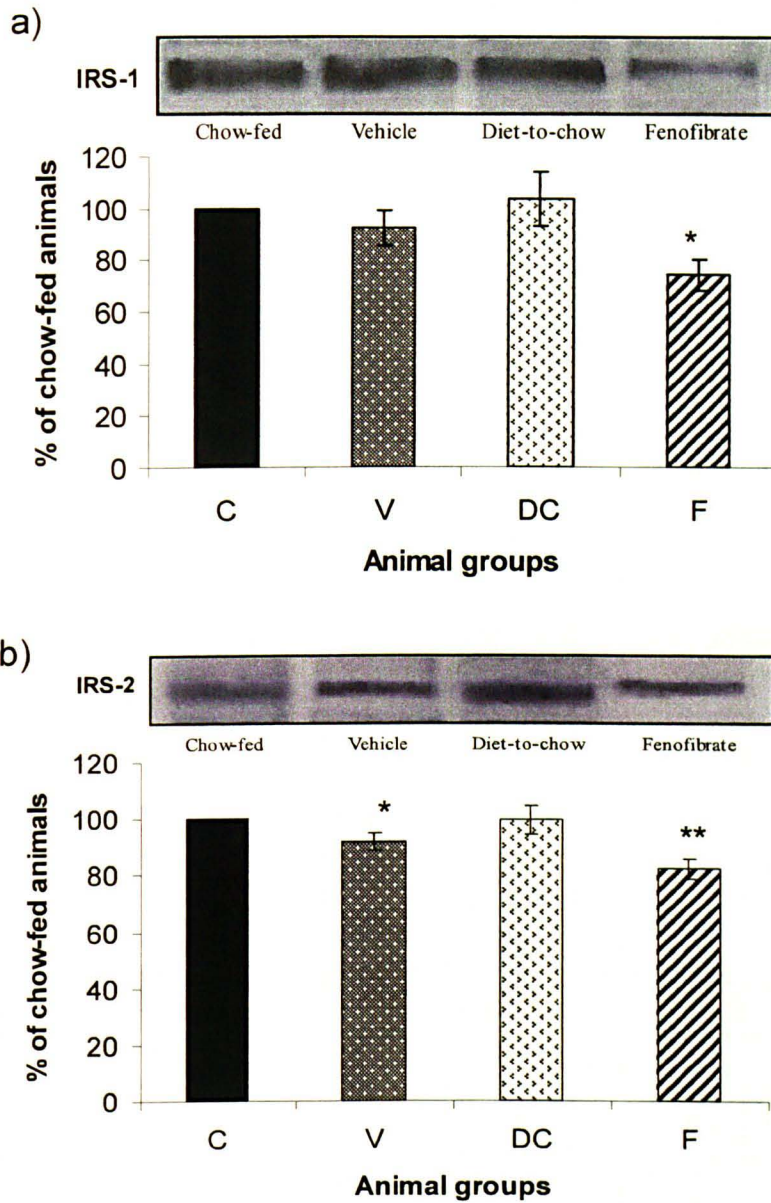


Fig. 8.4. Protein expression of a) IRS-1, and b) IRS-2 in the rat liver. Equal amounts ($50\mu\text{g}/\text{well}$) of protein were separated by SDS-PAGE and immunoblotted with IRS-1 and IRS-2 antibodies. The animal groups are: C) chow-fed lean control, V) Vehicle group (untreated dietary obese), DC) Diet-to-Chow group, and F) fenofibrate treated group. Data (mean \pm S.E.M) are normalised to that of control group and expressed as a percentage of the chow fed (control) rats. * = $p < 0.05$, ** = $p < 0.0005$ vs. chow-fed.

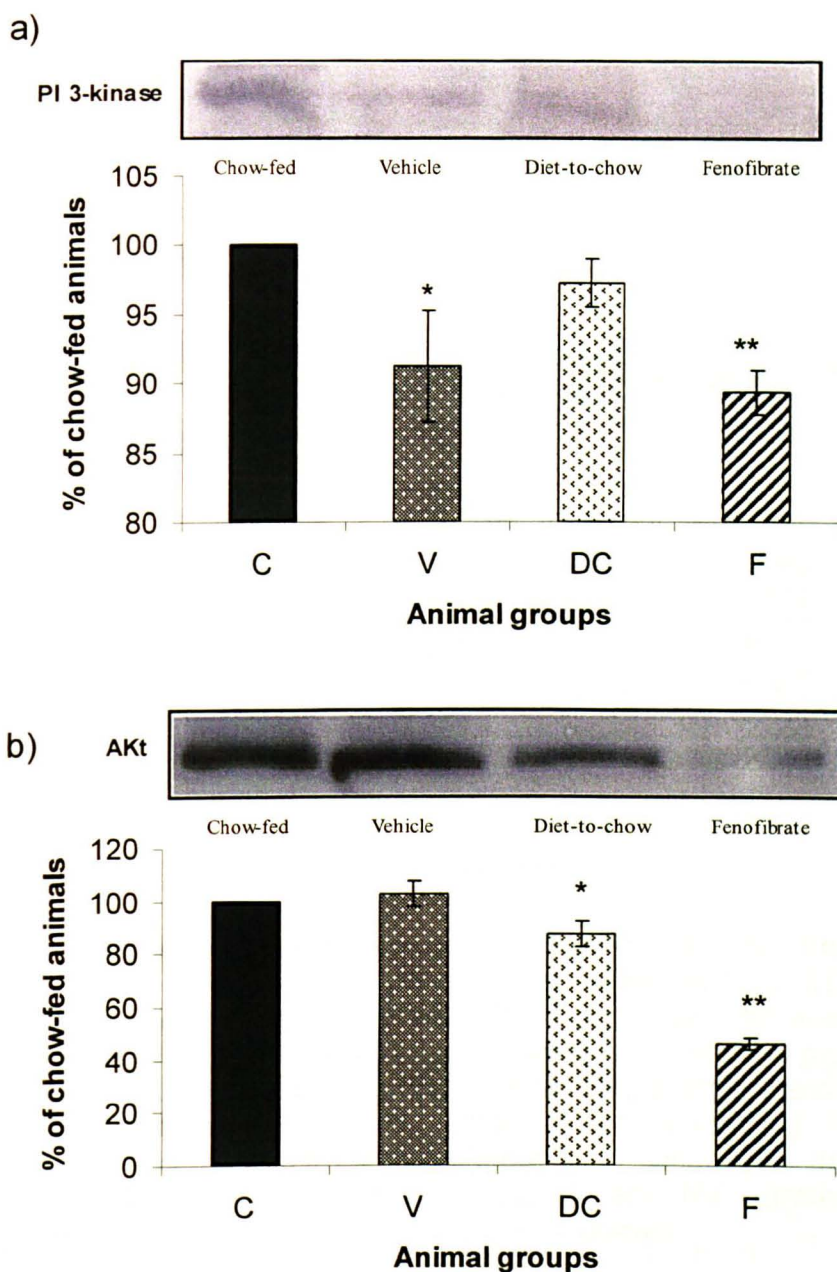


Fig. 8.5. Protein expression of a) PI 3-kinase, and b) AKT in the rat liver. Equal amounts (50 μ g/well) of protein were separated by SDS-PAGE and immunoblotted with AKT and PI 3-kinase antibodies. The animal groups are: C) chow-fed lean control, V) Vehicle group (untreated dietary obese), DC) Diet-to-Chow group, and F) fenofibrate treated group. Data (mean \pm S.E.M; n=10) are normalised to that of control group and expressed as a percentage of the chow fed (control) rats. * = $p < 0.05$, ** = $p < 0.000005$ vs. chow-fed.

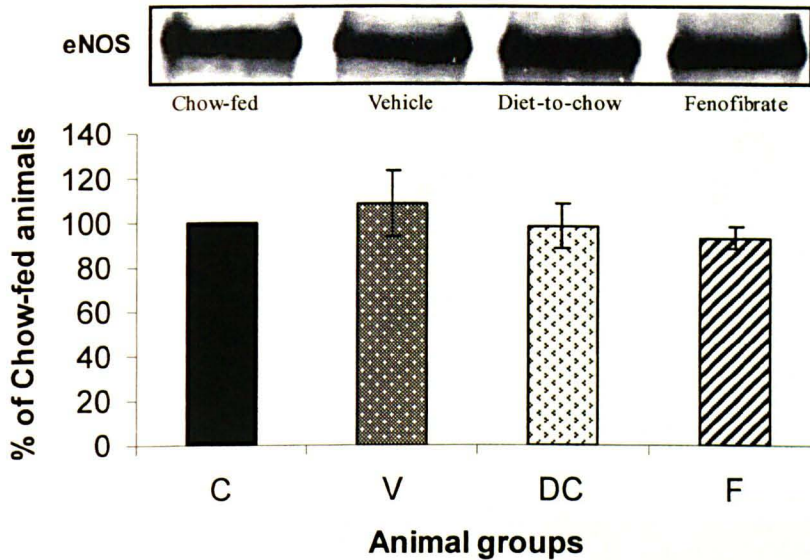


Fig. 8.6. Protein expression of eNOS in the rat liver. Equal amounts (50µg/well) of protein were separated by SDS-PAGE and immunoblotted with eNOS antibody. The animal groups are: C) chow-fed lean control, V) Vehicle group (untreated dietary obese), DC) Diet-to-Chow group, and F) fenofibrate treated group. Data (mean ± S.E.M) are normalised to that of control group and expressed as a percentage of the chow fed (control) rats. No significant differences were seen between the four groups.

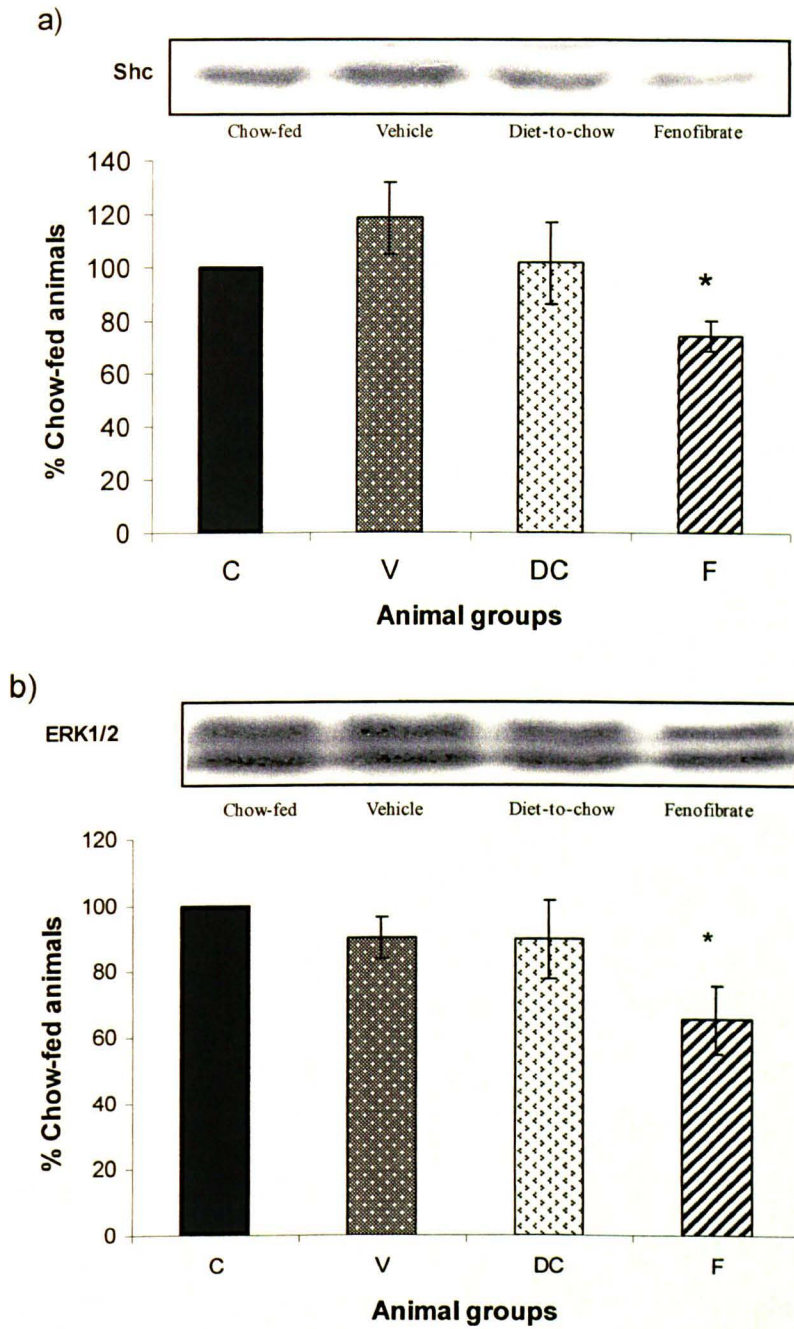


Fig. 8.7. Protein expression of a) Shc, and b) ERK1/2 in the rat liver. Equal amounts (50 μ g/well) of protein were separated by SDS-PAGE and immunoblotted with Shc and ERK1/2 antibodies. The animal groups are: C) chow-fed lean control, V) Vehicle group (untreated dietary obese), DC) Diet-to-Chow group, and F) fenofibrate treated group. Data (mean \pm S.E.M) are normalised to that of control group and expressed as a percentage of the chow fed (control) rats. * = $p < 0.01$ vs chow-fed.

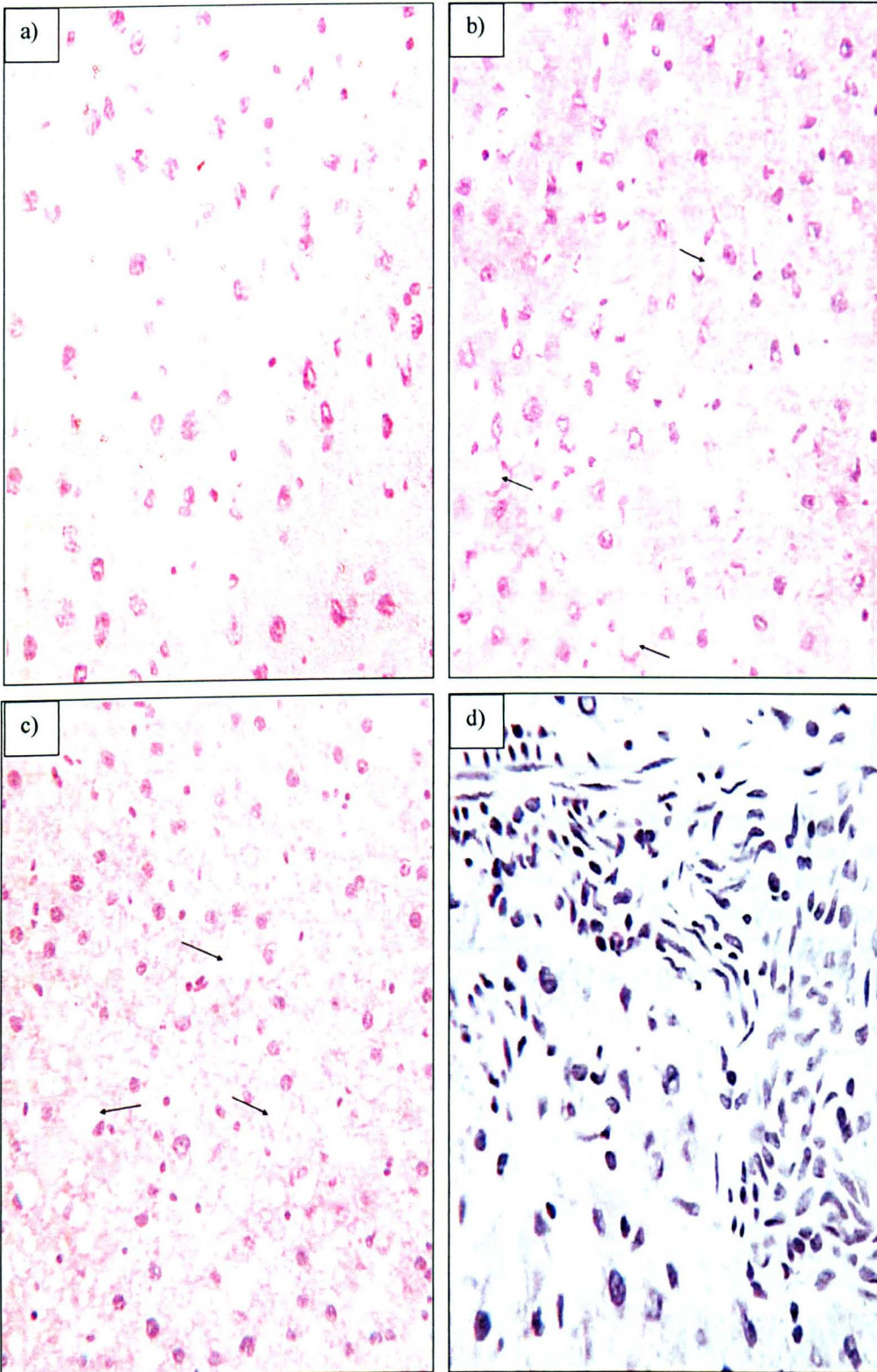


Fig. 8.8. Histological sectioning for the liver: a) normal liver, b) steatosis fatty liver grade one, c) steatosis fatty liver grade two, and d) steatohepatitis grade one.

| Animal groups | STEATOSIS (FATTY CHANGE IN THE LIVER) | | | | Total % of fatty change in each group |
|--|--|-----|-----|-----|---------------------------------------|
| | Grade | | | | |
| | 0 Negative samples | 1 | 2 | 3 | |
| Chow-fed (control) (10 animals=100%) | 50% | 50% | | | 50% |
| Vehicle (obese animals) (10 animals=100%) | 20% | 40% | 30% | 10% | 80% |
| Diet-to-chow (10 animals=100%) | 50% | 40% | 10% | | 50% |
| Fenofibrate treatment (10 animals=100%) | 90% | | 10% | | 10% |

Table 8.1 Total percentage of hepatic fatty change (steatosis). Stage I of non-alcoholic fatty liver disease: simple steatosis (fat accumulation in hepatocytes) in the four experimental groups: Chow-fed, Vehicle, Diet-to-chow, and Fenofibrate treated group. Tissue liver of 10 male Wister rats from each group were analyzed microscopically for the presence of different grades of Steatosis. Each analyzed animal represent 10% of the total number of animals in each group.

| Animal groups | INFLAMMATION | | | | Total % of inflammation in each group |
|--|-----------------------|------|---|---|---------------------------------------|
| | Grade | | | | |
| | 0 Negative samples | 1 | 2 | 3 | |
| Chow-fed Control (10 animals=100%) | 60% | 40 % | | | 40 % |
| Vehicle (obese animals) (10 animals=100%) | 30% | 70 % | | | 70 % |
| Diet-to-chow (10 animals=100%) | 70% | 30 % | | | 30 % |
| Fenofibrate treatment (10 animals=100%) | 90% | 10 % | | | 10 % |

Table 8.2 Total percentage of hepatic inflammation. Stage II of non-alcoholic fatty liver disease: inflammation (steatohepatitis) in the four experimental groups: Chow-fed, Vehicle, Diet-to-chow, and Fenofibrate treated group. Tissue liver of 10 male Wister rats from each group were analyzed microscopically for the presence of different grades of inflammation. Each analyzed animal represent 10% of the total number of animals in each group.

| Animal groups | Fibrosis | | Hepatocellular carcinoma | |
|---------------------|-----------------------|-----------------------|--------------------------|-----------------------|
| | % of negative samples | % of positive samples | % of negative samples | % of positive samples |
| Chow-fed | 100% | - | 100% | - |
| Vehicle | 100% | - | 100% | - |
| Diet-to-chow | 100% | - | 100% | - |
| Fenofibrate-treated | 100% | - | 100% | - |

Table. 8.3 Advanced histological features of non-alcoholic fatty liver disease. Ten liver samples (100%) of each group were analysed microscopically for the presence of: fibrosis and hepatocellular carcinoma. All of the samples were negative for both histological features of non-alcoholic fatty liver disease.

8.4. DISCUSSION

In liver, Insulin has a crucial role in mediating carbohydrate and lipid homeostasis. Therefore, the integrity of hepatic insulin signalling is important for the actions of insulin in hepatocytes which includes stimulation of glycogen synthesis, lipogenesis, and lipoprotein synthesis and suppression of gluconeogenesis / glycogenolysis and VLDL secretion in fed state (Sparks and Sparks, 1994; Zammit, 2000). Insulin also plays a role in hepatic growth and regeneration (Starzl *et al.*, 1976). Deficiency in protein expression of any components of insulin transduction may induce hepatic insulin resistance and subsequently participate induction of systemic insulin resistance and type II diabetes, while local hepatic insulin resistance may cause nonalcoholic fatty liver disease.

Insulin has a major role in glucose homeostasis by uploading circulating glucose into skeletal muscle cells. Therefore, it is expected that malfunctioning of insulin-receptors in skeletal muscle cells may cause imbalance of whole-body glucose homeostasis. However, mice with severe insulin resistance in skeletal muscle due to a muscle-specific insulin receptor knockout have surprisingly normal whole-body glucose homeostasis (Bruning *et al.*, 1998).

In general, protein kinase phosphorylation and activation levels depend on the concentration of the expressed protein. Several studies demonstrated direct relationship between concentration of expressed protein kinase and levels of its activation (Anai *et al.*, 1998). Therefore, in hepatocytes, defective protein kinase expression may cause insulin resistance, with subsequent induction of hepatic insulin resistance-related clinical disorders, namely nonalcoholic fatty liver disease.

In the present study diet-induced obesity did not affect expression of insulin receptor- β (IR- β) and insulin receptor substrate-1 (IRS-1), indicating that the detrimental effects of diet-induced obesity on hepatic insulin signalling is at postreceptor levels. Furthermore, previous studies have demonstrated that levels of both proteins IR- β and IRS-1 in the different types of genetic obesity could be modified that may contribute to inducing hepatic insulin resistance. Friedman and colleagues (Friedman *et al.*, 1997) detected that the genetically obese

hypertensive strain (SHROB) is characterised by reduced IR- β auto-phosphorylation, and this reduction of protein activation was due to inhibited expression of IR- β itself in hepatocytes. Another study revealed that in the genetic obesity *ob/ob* mice, levels and activation of IRS-1 is reduced significantly (Saad *et al.*, 1992). Taken together, these reports have indicated that the molecular mechanism of hepatic insulin resistance in genetic obesity may involve modification in the early steps (IR- β and IRS-1 proteins) of insulin signalling, and that the amount of protein expression may correlate directly with the levels of phosphorylation and activation of the protein (Kim *et al.*, 2000). However in the present study of diet-induced obesity, unlike genetic obesity study, IR- β and IRS-1 did not involve in genesis of hepatic insulin resistance.

Insulin receptor substrate-2 (IRS-2) appears to have a major role in liver by regulating hepatic insulin action, whereas IRS-1 appears to have important functions in skeletal muscle (Sesti *et al.*, 2001). Cell-based experimentation with hepatocytes suggests that IRS-2 is the major effector of both the metabolic and the growth-promoting actions of insulin (Rother *et al.*, 1998). Moreover IRS-2 signalling is essential for the suppression of gluconeogenesis and apoptosis in hepatocytes. IRS-2 is an important regulatory point for the PI 3-kinase-Akt cascade in hepatocytes. PI 3-kinase activity associated with IRS-2 depends mainly on the level of IRS-2 expression and degree of phosphorylation (Rother *et al.*, 1998; Ide *et al.*, 2004). In fact, total insulin-induced PI 3-kinase activity decreased by 50% in IRS-2 (-/-) hepatocytes (Valverde *et al.*, 2003), while other reports have concluded that reduced PI 3-kinase activity may play a role in the pathophysiology of insulin resistant diabetic states, such as that seen in the *ob/ob* mouse (Folli *et al.*, 1993; Anai *et al.*, 1998). In the present study, levels of expression of both proteins (IRS-2 and PI 3-kinase) were significantly decreased in diet-induced obese animals, suggesting that diet-induced obesity have similar detrimental effects as genetic obesity on IRS-2 and PI 3-kinase proteins in hepatocytes.

The normal concentration of Akt protein in hepatocytes of diet-induced obese animals in the current study is consistent with other reports (Kim *et al.*, 2000) where they found unchanged levels of protein expression and slightly elevated

phosphorylation of Akt in genetic obesity. Therefore, both types of obesity might have similar effects on Akt protein expression.

Much less is known about the expression of eNOS in normal, diseased, and treated livers. eNOS in hepatocytes may regulate hepatic vasculature, regulating blood flow and blood cell interactions (Li and Billiar, 1999). In the present study, expression of eNOS in liver was not affected by the different dietary manipulations or by disease states (normal, steatosis, or steatohepatitis). This finding is in agreement with others demonstrating that the expression of eNOS protein and its activity in hepatocytes was not changed in chronic liver disorders (such as alcoholic cirrhosis, viral hepatitis, cholestasis, and fulminant liver failure) (McNaughton *et al.*, 2002). On the other hand, normal expression of MAP-kinase pathway is a crucial factor in mediating the mitogenic response of insulin. A defect in PI 3-kinase may overactivate MAP-kinase pathway. In the present study, diet-induced obesity did not alter the expression of two important protein kinases (Shc and ERK1/2) in MAP-kinase pathway, despite significant protein deficiencies in PI 3-kinase pathway in hepatocytes.

Interestingly, in the present study the effect of palatable-diet removal on expression of AKt protein in hepatocytes was similar to its affect on Akt in aorta (chapter 7). Moreover, this reduction in Akt expression in hepatocytes might be related to the sharp decrease in calorie intake. It is possible to correlate the reduced AKt levels to improvement in liver histology seen in this study. Therefore, it is possible that, the presence of other isoform(s) of Akt might have greater effect on insulin transduction and glucose metabolism than the studied isoform of AKt in the present study (AKt1). However, other studies have detected the importance of AKt2 protein in mediating the effect of calorie restriction on insulin signalling and glucose homeostasis (McCurdy *et al.*, 2003; McCurdy, 2005). McCurdy and colleagues demonstrated that moderate calorie restriction for 20 days induced a decrease in AKt1 levels. However, in skeletal muscle of rat, there was no change in AKt2 protein abundance but its activation was increased significantly, indicating the importance of (AKt2) in insulin sensitivity (McCurdy *et al.*, 2003). Taking together, the results of the present study and the above reports would indicate that

the levels of AKt1 protein can be affected by the amount of ingested calorie (calorie intake), and might have no major effect on insulin sensitivity in liver and vasculature.

Upregulation of TNF- α is associated with reduced activation of AKt (Gupta and Khandelwal, 2004; Grzelkowska-Kowalczyk and Wieteska-Skrzeczynska, 2006; Tazi *et al.*, 2006). Hence, this study and the other reports together may indicate that overexpression of TNF- α might have inhibitory effects on both protein expression and activity of AKt.

Fatty liver (steatosis), the first stage of NAFLD was apparent in this study. Histological sectioning of liver samples of diet-induced obese animals showed a considerable amount of fat droplets, which was further confirmed by a significant elevation of triglyceride content of the liver samples. Steatohepatitis (inflammation of the liver), the second stage of NAFLD, in untreated obese animals was also confirmed by the histological sectioning of the liver, indicating that steatosis was induced as result of increasing influx of lipid into the liver (Gauthier *et al.*, 2006) in obese animals. Simple steatosis might have also progressed into steatohepatitis due to the increased secretion of inflammatory markers from adipose tissue, which include TNF- α , (Crespo *et al.*, 2001; Li *et al.*, 2003) and CRP (Koruk *et al.*, 2003; Uchihara and Izumi, 2006). In the presence of hepatic insulin resistance, where PI 3-kinase pathway is inhibited and MAP-kinase (ERK1/2) is overactivated liver disease may progress further. Previous studies, have shown that regenerative response to liver injury is impaired in *ob/ob* obese mice with fatty livers (Yang *et al.*, 2001), providing further support for the concept that inhibited regeneration contributes to disease progression in NAFLD. Their results indicate that fatty hepatocytes in *ob/ob* mice become trapped in G₁, the prereplicative phase of the cell cycle. They identified a mechanism that is likely to contribute to this cell cycle arrest. The coordinated activation of the oxidant-sensitive, stress and mitogen-activated protein kinase cascades that normally occurs early in G₁ (Lee *et al.*, 1999) does not occur in fatty hepatocytes, which have adapted to a state of chronic oxidant stress (Yang *et al.*, 2000). Rather, in fatty liver, activation of the mitogen-activated kinase, ERK-1 and ERK-2, is enhanced, whereas induction of the stress-

activated kinase, Jnk, is abolished. ERK and Jnk have multiple cellular targets and interact to orchestrate the complex hepatic response to injury, inducing genes that regulate cell survival, proliferation, and certain-tissues-specific functions (Yan *et al.*, 1994; Fan and Derynck, 1999; Talarmin *et al.*, 1999). Therefore, fatty *ob/ob* hepatocytes fail to proliferate despite robust ERK activation when Jnk induction is abolished (Bost *et al.*, 1997; Lee *et al.*, 1999). Thus, these experiments identify a mechanism (i.e. activation of ERK and Jnk inhibition) that is likely to inhibit proliferation in fatty hepatocytes. Furthermore, TNF- α has been shown to enhance apoptosis, an action which can be inhibited by Jnk (Ventura *et al.*, 2004). Hence, inhibition of Jnk in obesity (Yang *et al.*, 2001) and increased TNF- α in this study may suggest an enhanced apoptosis in obese livers which may contribute to the progression of NAFLD.

A key feature in the molecular pathogenesis of liver fibrosis requires maintenance of the activated hepatic stellate cells phenotype by both proliferation and inhibition of apoptosis. Another study has demonstrated that leptin is a potent hepatic stellate cell mitogen and dramatically inhibits stellate cell apoptosis *via* ERK1/2 and AKt kinases (Saxena *et al.*, 2004). In this study, plasma leptin concentration is elevated in diet-induced obese animals, a finding which may promote the implication of leptin and ERK1/2 in NAFLD progression. Similarly, obese animals had elevated plasma CRP, a protein which has been shown to be an independent risk factor for NAFLD (Park *et al.*, 2004).

The above results of the current study, indicated that the different detrimental factors associated with diet-induced obesity such as increased plasma and liver lipid profile, increased secretion of the different inflammatory markers and adipocytokines (Table 8.4), in addition to hepatic insulin resistance (inactivation of PI 3-kinase and overactivation of MAP-kinase (ERK1/2) pathway) may involve “directly or indirectly” in inducing the different stages of NAFLD.

Lifestyle modification of diet-induced obesity, which represented in the present study by chronic withdrawal of highly-palatable diet (diet-to-chow group), retarded the progression of NAFLD and improved the histological features of this disease to

similar levels of chow-fed animals (Table 8.4). On the other hand, the pharmacological treatment (fenofibrate administration) further improved histological features to levels better than that of chow-fed group. Interestingly, improvements in histological features were paralleled with similar improvement in three detrimental factors associated with diet-induced obesity; namely, TG content of the liver, plasma concentration of TNF- α , and total body weight, suggesting their role in initiating and inducing the different histological features of NAFLD. Moreover, triglyceride content of plasma and liver appears to have a major role in normalising TG content in diet-to-chow and fenofibrate-treated groups, and this may participate in restoring the histological features. Plasma CRP and leptin may also have affected progression or retardation of the histological features of NAFLD, but to a lesser extent. In fenofibrate-treated group, levels of both CRP and leptin were highly elevated, while the histological features were reduced to the minimum level, indicating absence of any considerable correlation between leptin and CRP with development of NAFLD.

Overall, this study suggests that diet-induced obesity results in hepatic insulin resistance possibly by affecting expression and activation of insulin signaling proteins. Moreover, dietary obesity affects liver histology leading to development of NAFLD. Development of NAFLD may be associated by accumulation of triglycerides in liver and/or changes in insulin signaling pathway in the liver.

Table 8.4. Summary for biochemical and hepatic features of the three experimental groups

| Biochemical & pathological parameters | | Experimental groups | | | |
|---------------------------------------|-----------------------------|---------------------|----------|--------------|--------|
| | | Chow-fed | Diet-fed | Diet-to-chow | Feno. |
| Metabolic Parameters | Insulin | Normal | ↑↑ | Normal | Normal |
| | Glucose | Normal | ↑↑ | Normal | Normal |
| | Homa | Normal | ↑↑ | Normal | Normal |
| | TG | Normal | ↑↑ | Normal | Normal |
| | NEFA | Normal | Normal | Normal | Normal |
| | Body Wt. | Normal | ↑↑↑ | ↑ | Normal |
| Hepatic features | Steatosis +ve samples | 50% | 80% | 50% | 10% |
| | Steatohepatitis +ve samples | 40% | 70% | 30% | 10% |
| | Cirrhosis | 0% | 0% | 0% | 0% |
| | Carcinoma | 0% | 0% | 0% | 0% |
| | TG content | Normal | ↑↑↑ | Normal | ↓↓ |
| | Liver weight | Normal | ↑ | Normal | ↑↑↑ |
| Inflammatory markers | Plasma TNF-α | Normal | ↑↑ | ↑↑ | Normal |
| | Plasma CRP | Normal | ↑↑ | Normal | ↑↑ |
| | Plasma leptin | Normal | ↑↑ | Normal | ↑↑ |

Chapter 9

RESULTS: Effects of diet-induced obesity on protein expressions of insulin signalling pathways in the skeletal muscle of male Wistar rats

9.1 INTRODUCTION

Insulin resistance is a co-morbidity of overweight and obesity and is a risk factor for development of type 2 diabetes and cardiovascular disease. In obese individuals and patients with type 2 diabetes, skeletal muscle insulin resistance is characterised by markedly blunted insulin-stimulated glucose uptake and metabolism (Dohm *et al.*, 1988). Since skeletal muscle is the primary site of glucose disposal in the human body, the inability of this tissue to uptake glucose in response to insulin most probably explains the reduced *in vivo* disposal of glucose observed in insulin-resistant individuals. At the cellular level, metabolic insulin resistance in diet-induced obesity displays a disturbed protein expression and activation (mainly in skeletal muscle and liver) of signalling *via* the IRS-phosphatidylinositol (PI) 3-kinase pathway, accompanied by a reduction in glucose uptake and utilization (Shulman, 1999).

Systemic lipid accumulation (McGarry, 2002) and increase in the intramuscular concentration of lipid intermediates such as fatty acyl-CoA, ceramides, and diacylglycerols (Despres *et al.*, 1996) not only correlate with insulin resistance, but also directly and indirectly alter insulin signalling (Schmitz-Peiffer, 2002). Considerable evidence linking increased skeletal muscle lipid content to insulin resistance has been derived from animal studies employing acute and chronic high-fat diets (Cooney *et al.*, 2002). Moreover, it has been documented that Protein Kinase-C (PKC) causes insulin resistance in skeletal muscle (Bossenmaier *et al.*, 1997). Furthermore, among the eight PKC isoforms, PKC- β is the only isoform increased in insulin-resistant muscle. Basal PKC- β is higher in the *in vitro* incubated muscles from obese individuals, and insulin increases PKC- β in muscle from obese patients but not from lean individuals (Itani *et al.*, 2000).

Various intramuscular lipid intermediates such as fatty acyl-CoA, ceramides, and DAG inhibit the insulin signalling cascade steps (Schmitz-Peiffer, 2002). Ceramide activates a protein phosphatase that dephosphorylates Akt (protein kinase B), resulting in inhibition of GLUT4 translocation and glycogen synthesis (Long and Pekala, 1996; Chavez *et al.*, 2003).

There is a close relationship between insulin resistance and visceral adiposity. It has become apparent that insulin sensitivity is modulated by adiposity, and this has led to a significant interest in adipocytokines such as TNF- α , leptin, and CRP, which may play pivotal role in skeletal muscle insulin signalling and resistance (Bastard *et al.*, 2006). Moreover, insulin sensitivity is restored by treatments that reduce intramuscular lipid accumulation such as lifestyle modification (i.e. low-fat feeding, fasting and exercise) (Oakes *et al.*, 1997; Monzillo *et al.*, 2003), and pharmacological therapy (e.g. fenofibrate, a PPAR- α activator), which improve lipid profile and insulin resistance (Ye *et al.*, 2001; Lee *et al.*, 2002). However, the exact molecular mechanism(s) involved in insulin transduction in skeletal muscle still not well understood. Therefore, the present study aims to investigate the protein expressions of PI 3-kinase and MAP-kinase insulin signalling pathways in skeletal muscle of diet-induced obese rats, and analysing the relationship between lipid profile, protein expression, and induction of skeletal muscle insulin resistance.

9.2 METHODS

9.2.1. Experimental protocol

Animals used in this study were those described in chapter 7. At the end of experimental protocol (as described in chapter 7), the rats were killed by CO₂ inhalation after 2 hours of fasting, skeletal muscle gastrocnemius was dissected and snap frozen in liquid nitrogen, for later molecular techniques.

9.2.2 Experimental procedures

9.2.2.1 Western-immunoblotting technique

Western-immunoblotting technique was carried out (as described in chapter two, section 2.4) on the four experimental groups (chow-fed, vehicle, diet-to-chow, and fenofibrate treated groups) for measuring the expression of proteins involved in insulin signalling pathways in skeletal muscle. The measured protein kinases were: PKC- β , IR- β , IRS-1, PI 3-kinase, AKt, Shc, ERK1/2 and eNOS.

9.2.2.2 Triglyceride content of gastrocnemius skeletal muscle

A small sample of gastrocnemius muscle (160-200 mg) was homogenised in 500 μ l distilled water with polytron homogeniser. The homogenate were centrifuged for

5 minutes at 13000 x g, and the supernatants were transferred to eppendurff tubes. The triglycerides assay was performed on the supernatant to detect the concentration of TG in muscle.

9.2.3 Data interpretation and statistical analysis

For Western blotting, the data from chow-fed (control) animals were expressed as 100% response, and the results from other three groups were normalised and subsequently expressed as the percentage of their respective controls. Data are expressed as mean \pm S.E.M. Data have normal distribution (Shapiro Wilk W test). Statistical significance was tested using Student 't-test' or repeated-measures (ANOVA; Bonferroni t-test) or the Mann-Whitney test, as appropriate. Results were considered statistically significant at the $p < 0.05$.

9.3. RESULTS

9.3.1 Body weight

By referring to the results of changes in total body weight in chapter 7, animals given palatable diet progressively gained more weight than their chow-fed counterparts. Untreated diet-fed animals were significantly heavier than chow-fed, diet-to-chow, and fenofibrate treated animals. However, at the end of the experiment, body weight of diet-to-chow and fenofibrate treated animals were significantly ($p < 0.01$) lower than untreated diet-fed animals (for more statistical details see results in chapter 7).

9.3.2 Triglyceride content of gastrocnemius skeletal muscle

Triglyceride concentration was measured in tissue of skeletal muscle (Fig. 9.1) in all three experimental groups: vehicle, diet-to-chow, and fenofibrate-treated, groups. TG levels in muscle of vehicle group were significantly increased (70%, $p < 0.05$) in comparison with chow-fed control group, while TG concentrations of the other two groups (diet-to-chow and fenofibrate-treated) were comparable with chow-fed control group, with no significant changes (Fig. 9.1).

9.3.3 Expression of protein kinases of insulin signalling pathways in skeletal muscle

PKC- β . Compared with chow-fed group, expression of PKC- β levels were significantly increased in both; vehicle (113%, $p < 0.01$) and fenofibrate-treated (137%, $p < 0.05$) groups, but not in diet-to-chow group (Fig. 9.2).

IR- β . The expression of IR- β protein was not affected by the three experimental groups: vehicle, diet-to-chow, and fenofibrate-treated groups. IR- β levels were similar in all groups with no significant difference between the groups (Fig. 9.3a).

IRS-1. Similar to IR- β , the expression of IRS-1 was comparable in all three experimental groups; vehicle, diet-to-chow, and fenofibrate-treated, with no significant differences in comparison with chow-fed control group (Fig. 9.3b).

PI 3-kinase. Surprisingly, the concentrations of the regulatory subunit p85 of PI 3-kinase were significantly increased in both vehicle (77%, $p < 0.005$), and fenofibrate-treated (68%, $p < 0.05$) groups, but not diet-to-chow group compared with chow-fed control (Fig. 9.4a).

AKt. The expression of AKt protein in vehicle group (diet-induced obese animals) did not change significantly in comparison with control group (Fig. 9.4b). However, there were marked reduction in AKt levels in both groups: diet-to-chow (38%, $p < 0.001$) and fenofibrate-treated (24%, $p < 0.05$) groups when compared with chow-fed group. Moreover, AKt levels in diet-to-chow group were significantly lower ($p < 0.05$) than that of vehicle group (Fig. 9.4b).

Shc. In MAP-kinase pathway, expression of two important protein kinases "Shc and ERK1/2" were measured. Compared with chow-fed control group protein levels of Shc in vehicle group were significantly increased (65%, $p < 0.01$) (Fig. 9.5a), while the expression of Shc in the other two experimental groups: diet-to-chow and fenofibrate-treated groups were not significantly altered (Fig. 9.5a).

ERK1/2. The expression of ERK1/2 protein in vehicle group was significantly increased (50%, $p < 0.005$), while diet-to-chow group was decreased markedly

(28%, $p < 0.05$) when compared with control group (Fig. 9.5b). Moreover, ERK1/2 levels in skeletal muscle were not significantly altered by fenofibrate-treatment (Fig. 9.5b).

eNOS. Similar to PI 3-kinase, eNOS protein levels were remarkably augmented in both groups; vehicle (98%, $p < 0.001$) and fenofibrate-treated (115%, $p < 0.001$) groups in comparison with chow-fed control. However, there were no significant changes in eNOS expression in tissues from diet-to-chow group compared with chow-fed control (Fig. 9.6).

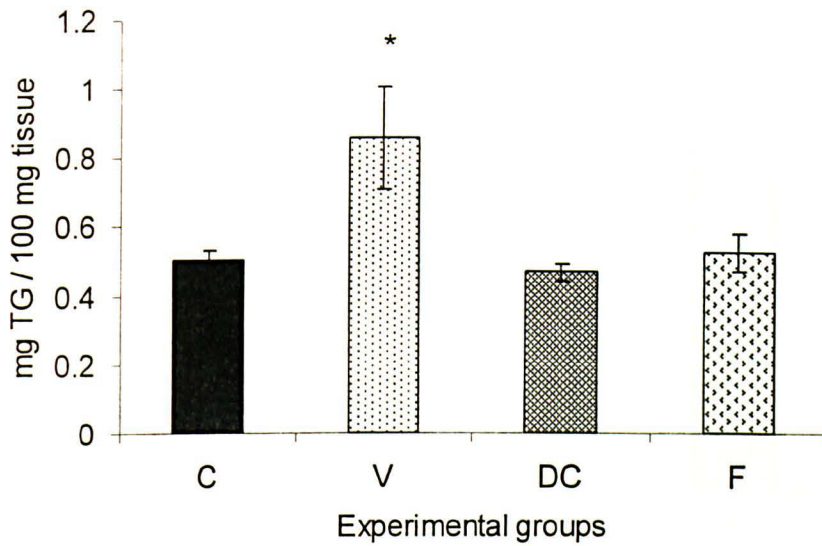


Fig. 9.1. Triglyceride (TG) content of gastrocnemius skeletal muscle in three experimental groups: V) vehicle, DC) diet-to chow, and F) fenofibrate-treated, groups. Data are presented as mean \pm S.E.M. * $p < 0.05$ vs chow-fed control. TG content of skeletal muscle in vehicle group was significantly higher than chow-fed group.

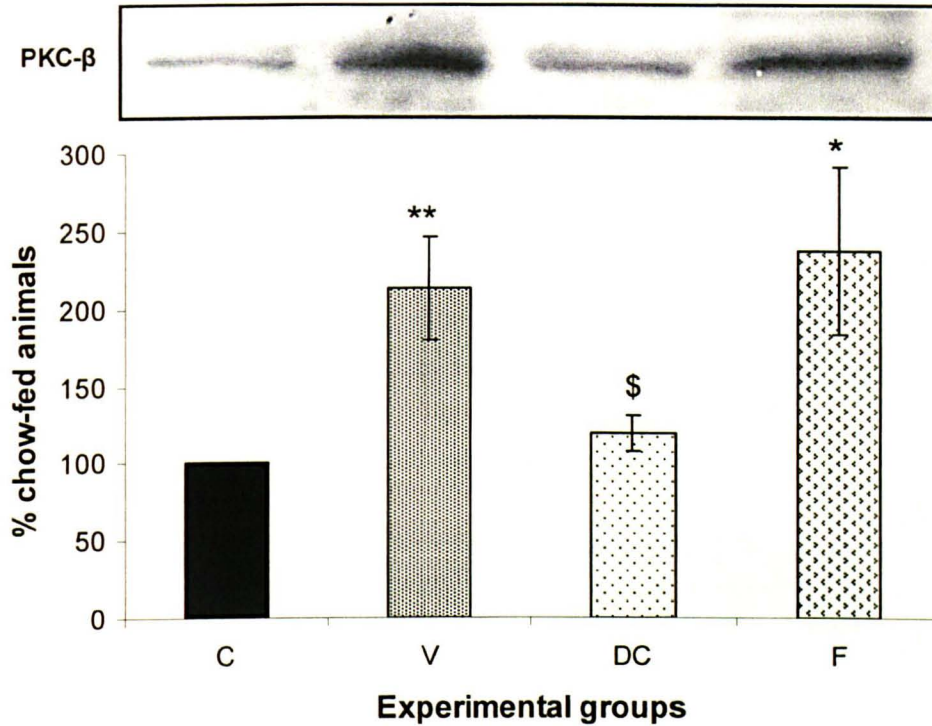


Fig. 9.2. Protein expression of PKC- β in the rat skeletal muscle. Equal amounts (40 μ g / well) of protein were separated by SDS-PAGE and immunoblotted with PKC- β antibody. The animal groups were: C) chow-fed lean control, V) Vehicle group (untreated dietary obese), DC) Diet-to-Chow group, and F) fenofibrate treated group. Data (mean \pm S.E.M) are normalised to that of control group and expressed as a percentage of the chow fed (control) rats. *= p <0.05, **= p <0.01 vs chow-fed, \$= p <0.05 vs vehicle.

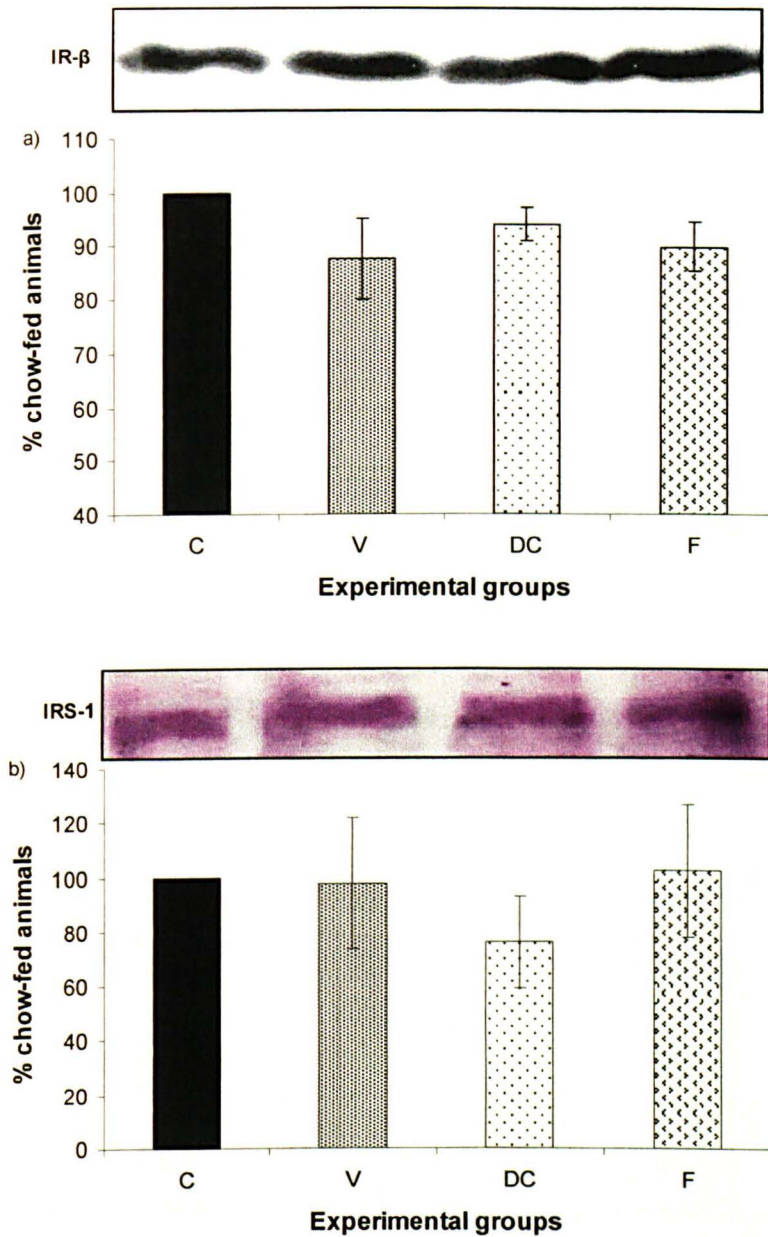


Fig 9.3. Protein expression of a) IR- β , and b) IRS-1 in the rat skeletal muscle. Equal amounts (40 μ g / well) of protein were separated by SDS-PAGE and immunoblotted with IR- β or IRS-1 antibodies. The animal groups are: C) chow-fed lean control, V) Vehicle group (untreated dietary obese), DC) Diet-to-Chow group, and F) fenofibrate treated group. Data (mean \pm S.E.M) are normalised to that of control group and expressed as a percentage of the chow fed (control) rats. There was no significant difference in protein expression between any group.

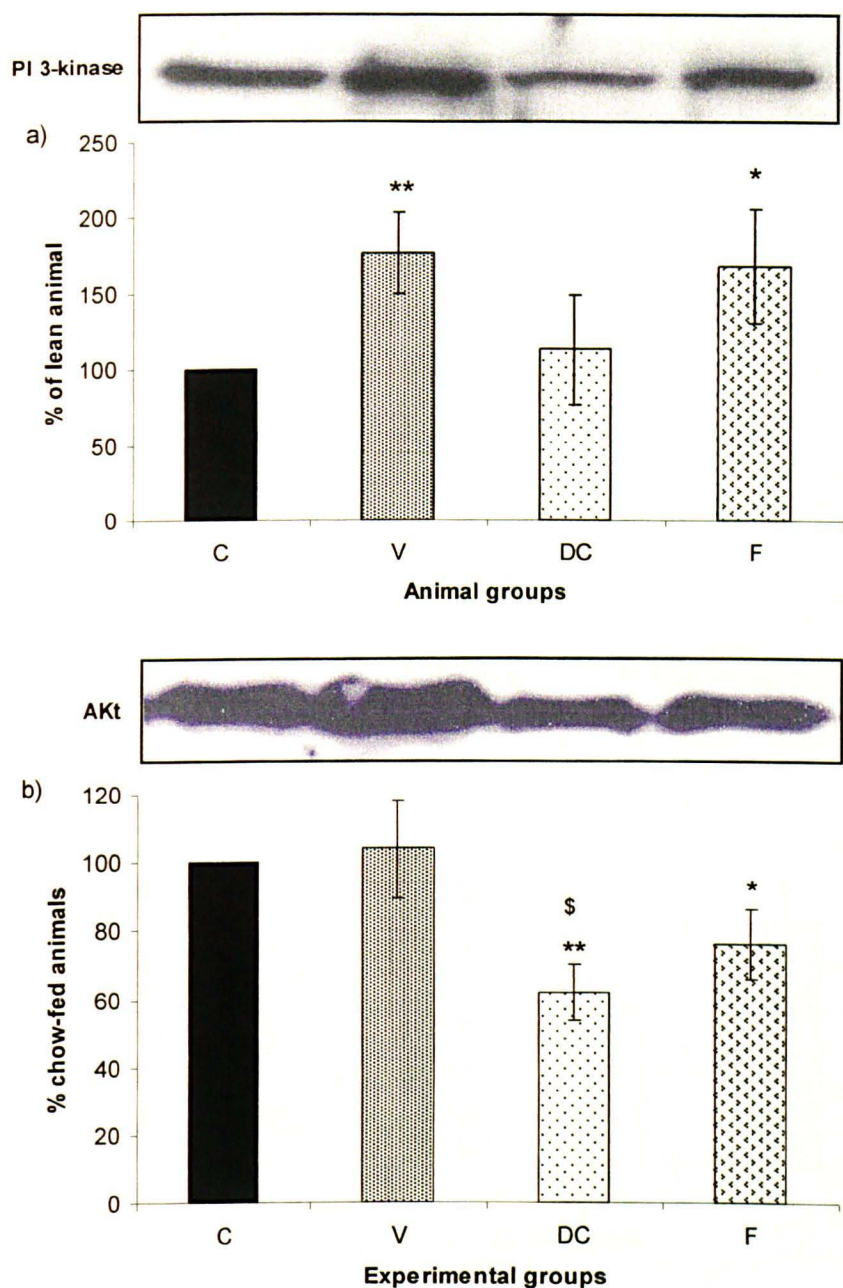


Fig. 9.4. Protein expression of a) PI 3-kinase, and b) AKt in the rat skeletal muscle. Equal amounts (40 μ g / well) of protein were separated by SDS-PAGE and immunoblotted with PI 3-kinase or AKt antibodies. The animal groups are: C) chow-fed lean control, V) Vehicle group (untreated dietary obese), DC) Diet-to-Chow group, and F) fenofibrate treated group. Data (mean \pm S.E.M) are normalised to that of control group and expressed as a percentage of the chow fed (control) rats. * $p < 0.05$, ** $p < 0.005$ vs chow-fed. \$ $p < 0.05$ vs vehicle.

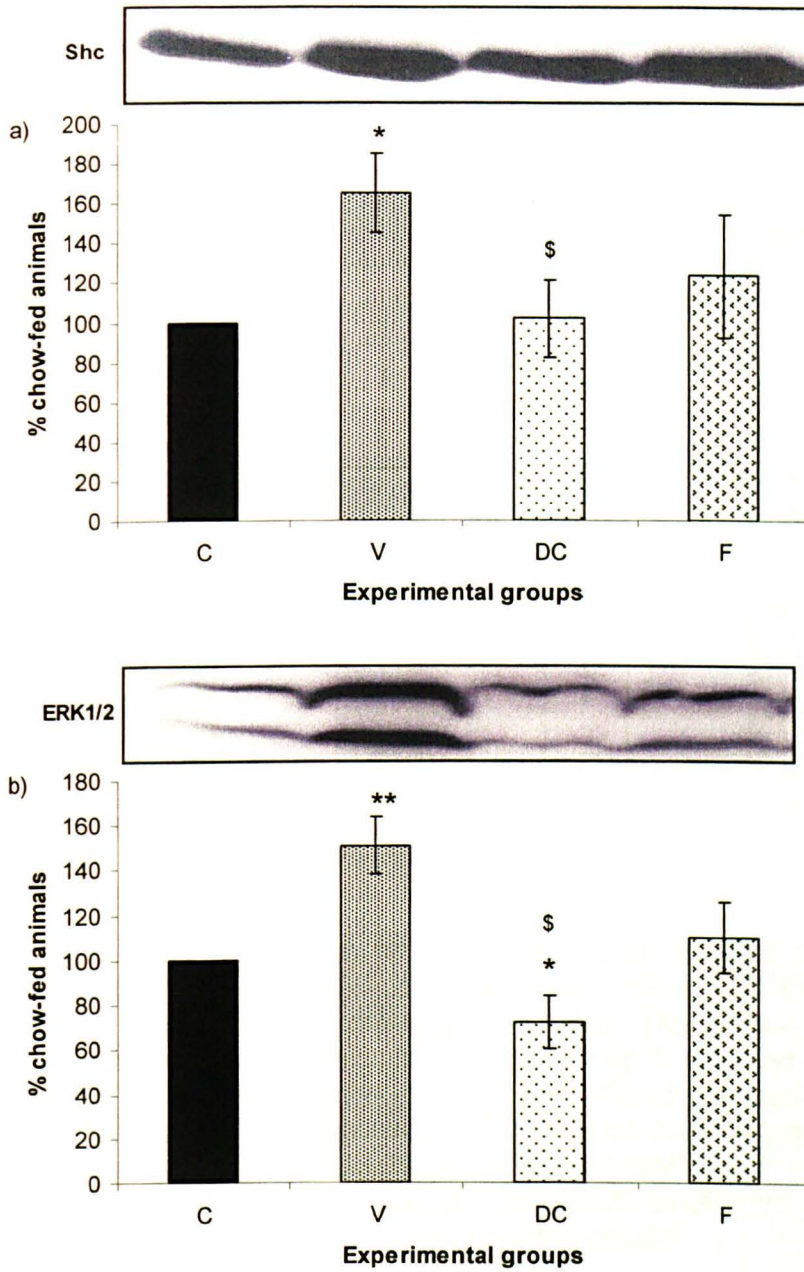


Fig. 9.5. Protein expression of a) Shc, and b) ERK1/2, in the rat skeletal muscle. Equal amounts (40 μg / well) of protein were separated by SDS-PAGE and immunoblotted with Shc or PERK1/2 antibodies. The animal groups are: C) chow-fed lean control, V) Vehicle group (untreated dietary obese), DC) Diet-to-Chow group, and F) fenofibrate treated group. Data (mean \pm S.E.M) are normalised to that of control group and expressed as a percentage of the chow fed (control) rats. *= $p < 0.05$, ** = $p < 0.005$ vs chow-fed. \$ = $p < 0.05$ vs vehicle.

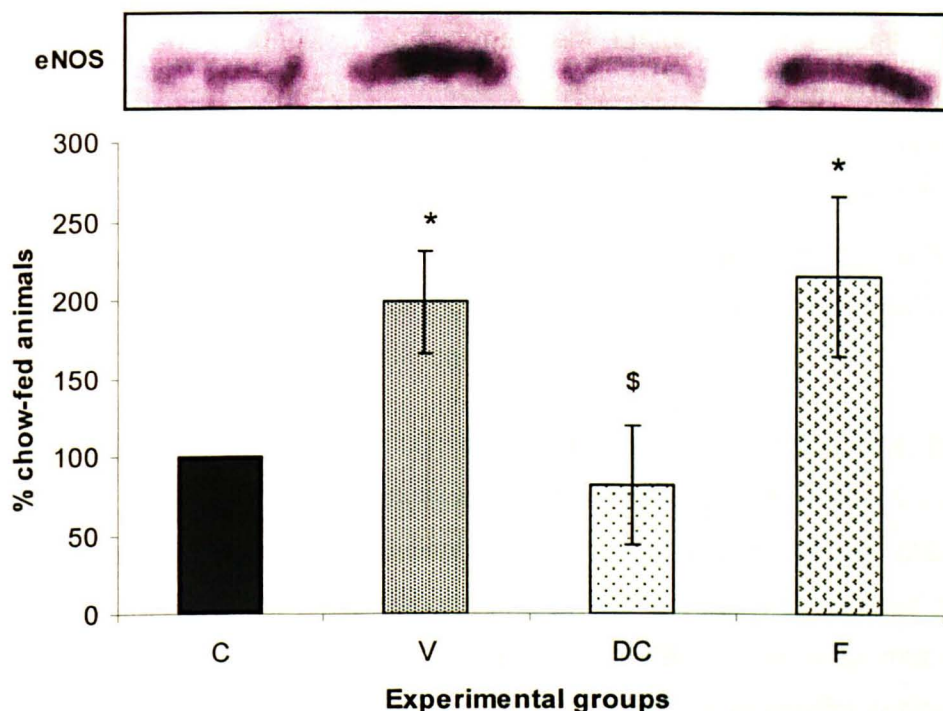


Fig. 9.6. Protein expression of eNOS in the rat skeletal muscle. Equal amounts (40 μg / well) of protein were separated by SDS-PAGE and immunoblotted with eNOS antibody. The animal groups are: C) chow-fed lean control, V) Vehicle group (untreated dietary obese), DC) Diet-to-Chow group, and F) fenofibrate treated group. Data (mean \pm S.E.M) are normalised to that of control group and expressed as a percentage of the chow fed (control) rats. * = $p < 0.05$ vs chow-fed. \$ = $p < 0.05$ vs vehicle. Vehicle & fenofibrate treated groups are significantly higher than chow-fed control.

9.4 DISCUSSION

Normal muscular insulin signalling is an important factor in the regulation of glucose metabolism. In fact, muscle insulin resistance is one of the major contributing factors to the development of diabetes mellitus type 2. In muscle, insulin signalling pathway is composed of cascade of protein kinases. Their expression and activation play crucial roles in maintaining normal glucose homeostasis. Therefore, any reduced strength in this insulin transduction may lead to skeletal muscle and whole body insulin resistance.

In diet-induced obesity, intramuscular TG (and its intermediates: ceramides, fatty acyl-CoA, and diacyl glycerol), hyperinsulinaemia, and the elevated levels of some adipocytokines, in cooperative or separate manner induce muscular insulin resistance by disturbing protein expressions and phosphorylation of kinases that belong to PI 3-kinase and MAP-kinase signalling pathways in skeletal muscle. Disruption of protein phosphorylation of the above cascades in insulin resistant skeletal muscle and obesity has received an adequate attention in literature and scientific research, while the associated disruption in the levels of their protein expressions still not well understood and merits further investigation.

The present study demonstrated that increased intramuscular triglyceride content was detected in diet-induced obese rats, and the greater lipid uptake by the muscle may be a potential mechanism responsible for this accumulation rather than due to a decrease in fatty acid oxidation (Hulver *et al.*, 2003). This hypothesis can be strengthened by a result in chapter five, where it has been detected that plasma levels of triglyceride in untreated diet-fed group (obese rats) were markedly elevated. Furthermore, reduction in fatty acid oxidation is not a necessity for increased intramuscular triglyceride intermediate (fatty acyl-CoA) accumulation (Oakes *et al.*, 1997; Hegarty *et al.*, 2003; Hulver *et al.*, 2003). Therefore, this study suggests that increased influx of lipid into skeletal muscle might play a major role in inducing augmentation of intramuscular triglyceride in diet-induced obesity. Moreover, both types of obesity treatment, chronic removal of highly-palatable diet and fenofibrate administration, caused remarkable reductions in the elevated levels

of intramuscular and plasma triglyceride. These improvements were occurred through two different mechanisms: in removal of highly-palatable diet, reduction in energy (calorie) intake was involved, and thus improving plasma and muscle lipid profile, while in fenofibrate treatment, increasing muscular fatty acid β -oxidation lead to raised levels of fatty acid utilization and clearance.

In the present study, PKC- β expression in gastrocnemius muscle of diet-induced obese rats were remarkably increased; and was associated with insulin resistance (as shown in chapter five), suggesting a role for PKC- β protein expression in glucose uptake in skeletal muscle (Bossenmaier *et al.*, 1997; Itani *et al.*, 2000). Triglyceride intermediates “fatty acyl-CoA, and diacylglycerol” might directly activate PKC- β (Shulman, 2000; Cooney *et al.*, 2002; Itani *et al.*, 2002; Schmitz-Peiffer, 2002). PKC- β activation in turn, may lead to negative effects on insulin transduction cascade in muscle. In this study, treating obesity by lifestyle modification “chronic withdrawal of highly-palatable diet” induced complete normalisation of PKC- β protein expression as well as an improvement in the intramuscular lipid accumulation and subsequent decrease in muscular and whole body insulin resistance. However, pharmacological treatment of obesity in this study (fenofibrate administration) failed to reduce or normalise the muscular expression of PKC- β , despite a remarkable improvement in intramuscular and plasma lipid concentrations and insulin resistance. Elevation of PKC- β levels in the muscle of fenofibrate-treated group could be due to excessive production of reactive oxygen species (ROS) *via* increased fatty acid- β oxidation. The Principal mechanism of lipid reduction by fenofibrate is via stimulating fatty acid- β oxidation in skeletal muscle and other tissues (Furuhashi *et al.*, 2002). ROS is one of the end products of fatty acid- β oxidation (Kudin *et al.*, 2005) which may have a major role in increasing expression and the phosphorylation of PKC- β (DeCarlo and Loeser, 2006). This hypothesis could be strengthened further by reports that fenofibrate-treatment is associated with increased ROS production and availability (Arnaiz *et al.*, 1995; Lores Arnaiz *et al.*, 1997; Jiao and Zhao, 2002; Jiao *et al.*, 2003). Hence, these results may indicate that diet-induced obesity can cause an increase in expression of protein kinase PKC- β in skeletal muscle, and this augmented level may participate in inducing muscular and subsequent whole body

insulin resistance. Furthermore, in the present study, the increased level of PKC- β was reversed completely by “chronic withdrawal of highly-palatable diet” but not pharmacologically (fenofibrate treatment), indicating a key role for the diet-content in the elevation of PKC- β .

The protein expression of the early steps of PI 3-kinase insulin signalling pathway such as IR- β and IRS-1 were not affected by diet-induced obesity. However, others have detected that tyrosine phosphorylation of IRS-1 and IR- β in insulin resistant skeletal muscle in obesity is reduced significantly (Goodyear *et al.*, 1995). These deficiencies in phosphorylation could be mediated by increased protein expression and activation of PKC- β , which in turn phosphorylate IR- β and IRS-1 in serine residue, consequently reducing their ability for tyrosine phosphorylation and development of subsequent insulin resistance (Caro *et al.*, 1987; Bossenmaier *et al.*, 1997; Zhou *et al.*, 1999; Kruszynska *et al.*, 2002).

Although, in the current study we didn't measure the protein expression of IRS-2 in skeletal muscle, several studies have indicated that IRS-1 and IRS-2 are not functionally interchangeable in tissues that are responsible for glucose production (Lehmann *et al.*, 1995), and glucose up take (skeletal muscle). Thus, IRS-1 appears to have its major role in skeletal muscle, whereas IRS-2 appears to regulate hepatic insulin action (Yamauchi *et al.*, 1996; Kido *et al.*, 2000; Previs *et al.*, 2000; Sesti *et al.*, 2001).

Interestingly, in this study, diet-induced obesity and/or overfeeding of highly-palatable diet induced a significant increase in protein expression of the regulatory subunit p85 of PI 3-kinase in skeletal muscle. Moreover, this increase in protein expression was synchronised with muscular and whole body insulin resistance. Therefore, it could be postulated that increased level of the p85 regulatory subunit might inhibit PI 3-kinase activity by competing with phosphotyrosine targets. Hence, disruption of the balance between the amounts of PI 3-kinase subunits and excess levels of p85 regulatory subunit could contribute to insulin resistance. The possible mechanism(s) of insulin resistance could be explained as follow: PI 3-kinase consists of a regulatory subunit, p85, and a catalytic subunit, p110

(Shepherd *et al.*, 1998). Normally, the regulatory subunit exists in excess to the catalytic one, resulting in a pool of free p85 monomers not associated with the p110 catalytic subunit. Thus, there is a balance between the free p85 monomer and the complex p85-p110 heterodimer, the latter being responsible for the PI 3-kinase activity. Increases or decreases in expression of p85 shift this balance in favour of either free p85 or p85-p110 complexes (Terauchi *et al.*, 1999; Ueki *et al.*, 2003). Because the monomer and the heterodimer compete for the same binding sites on the IRS proteins, an imbalance could cause either increased or decreased PI 3-kinase activity. This possibility has recently been supported by studies of the insulin-resistant states induced by human placental growth hormone (Barbour *et al.*, 2004), obesity and type 2 diabetes (Bandyopadhyay *et al.*, 2005). If the production of p85 can be enhanced nutritionally, the resulting changes in the ratio of p85 to p110 may be the earliest manifestation of the ensuing insulin resistance. This could also explain how overfeeding and weight gain trigger insulin resistance.

The levels of the protein expression of AKt in skeletal muscle did not change by diet-induced obesity and/or overfeeding of highly-palatable diet for fifteen week. A similar result has been also reported where the phosphorylation of AKt in skeletal muscle of insulin-resistant obese-diabetic subjects was not changed significantly (Kim *et al.*, 1999). Taken together, it can be argued that AKt in skeletal muscle has no role in inducing muscular insulin resistance. However, chronic withdrawal of the highly-palatable diet caused significant reduction in the protein levels of AKt compared to that of control group, despite the complete normalisation in the muscular and whole body insulin resistance, further indicating lack of a significant role for AKt in skeletal muscle insulin resistance. This in agreement with other studies, where Akt had no major role in resistance to insulin's action in muscle of obese type 2 diabetic subjects (Kim *et al.*, 1999). On the other hand, fenofibrate – treated animals, had lower AKt levels in skeletal muscle, despite an improved insulin signalling and muscular lipid concentration. Once again, this is in agreement with others showing that fenofibrate remarkably reduces AKt activity (Kubota *et al.*, 2005; Grabacka *et al.*, 2006). Therefore, it could be postulated that, the triglyceride-induced muscular insulin resistance, appears to be the consequence of a defect in insulin signalling pathway leading to impaired PI 3-

kinase activation. This in turn, may lead to impaired glucose transport through an AKt-independent pathway.

Lifestyle (withdrawal of highly-palatable diet), but not pharmacological modification of obesity improves protein expression of PI 3-kinase cascade of insulin signalling transduction in skeletal muscle. There is little or no information about mechanism of insulin-induced expression and activation of ERK1/2 MAP-kinase in insulin-resistant obese or diabetic animals. PI 3-kinase mediates most of the metabolic effects of insulin and its downstream signalling events (Kanai *et al.*, 1993; Shepherd *et al.*, 1995), while Shc and Grb2/Sos link insulin receptor signalling to activation of the mitogen-activated protein (MAP) kinase (MAPK) isoform p42MAPK/p44MAPK (ERK1/ERK2) cascade. The later is not required for insulin's metabolic effects, but mediates mitogenic signalling (Lazar *et al.*, 1995). The increased expression of both Shc and ERK1/2 in diet-induced obese animals in the present study, suggest a role for this pathway in insulin resistance. However, several studies have demonstrated that activation of MAP-kinase pathway by insulin is not reduced in insulin resistance (Paez-Espinosa *et al.*, 1999 ; Krook *et al.*, 2000; Cusi *et al.*, 2000 ; Paez-Espinosa *et al.*, 2001 ; Zecchin *et al.*, 2003). Furthermore, PKC- β may activate MAP-kinase pathway (Formisano *et al.*, 2000). Counteracting obesity by withdrawing highly-palatable diet or by fenofibrate administration, caused reduction in the expression of Shc and ERK1/2 in skeletal muscle, indicating that the augmented levels of these protein kinases in diet-induced obesity could be reversed by losing weight and subsequent improvement in lipid profile and insulin signalling. In fact, previous reports have indicated that lowering intramuscular lipid accumulation by removing fat-enriched diet or fenofibrate treatment improve muscular insulin resistance (Oakes *et al.*, 1997; Hegarty *et al.*, 2003; Ide *et al.*, 2004; Jazet *et al.*, 2005), perhaps by effecting Shc and ERK1/2 expression.

The protein kinase eNOS is present in endothelial cells of capillaries and larger blood vessels (Segal *et al.*, 1999) of skeletal muscle and in the fibre of skeletal muscle itself. Furthermore, insulin signalling shown to induce eNOS expression and activation (Fisslthaler *et al.*, 2003; Toba *et al.*, 2006), an effect which may not

be related to its metabolic functions (Lau *et al.*, 2000; Baum *et al.*, 2004; Williams *et al.*, 2006). The angiogenic factors such as VEGF and the process of angiogenesis (Silha *et al.*, 2005) are elevated in overweight and obese individuals. In the present study, hyperinsulinaemia (results in chapter 5) might be involved in inducing augmented levels of eNOS expression in skeletal muscle of diet-induced obese rats. Therefore, taken together these results, it is possible that elevated levels of eNOS in skeletal muscle from obese rats could have a major role in inducing the angiogenesis in this tissue. Treating obesity by removing the highly-palatable diet had reversed eNOS levels, while fenofibrate treatment was unable to reduce or improve the raised levels of eNOS.

In conclusion, this study indicate that diet-induced obesity results in insulin resistance in skeletal muscle mainly by increasing the protein expression of the regulatory subunit p85 of PI 3-kinase. Moreover, this augmented level was reversible by physiological but not pharmacological manipulation of obesity. This study also indicates that increased intramuscular triglyceride accumulation may play a principal role in triggering protein expression modification and possible subsequent muscular insulin resistance. Furthermore, protein expressions of early steps (IR- β and IRS-1) of PI 3-kinase pathway were not affected by muscular insulin resistance.

Chapter 10

General discussion

The current obesity epidemic is partly due to the consumption of a high energy diet, which is relatively abundant in atherogenic components such as carbohydrates and fat, leading to dyslipidaemia and vascular abnormalities. The atherogenic dietary components have different detrimental effects on various vascular functions. In fact, the present study indicates that all high energy diets, regardless of their varying compositions, were able to induce varieties of metabolic abnormalities and vascular dysfunctions. Moreover, combination of high amounts of atherogenic components (fat and carbohydrates) in one diet would exacerbate vascular dysfunction. For example: diets of combined atherogenic components such as chocolate, and biscuit had the most diverse detrimental effects on the vascular system (Table 10.1), causing increased KCl- and NA-induced vasoconstriction, and attenuation of vasodilatation in response to carbamylcholine (endothelium-dependent vasorelaxation), indicating endothelial dysfunction. On the other hand, other high energy diets such as highly-palatable and high-fat diets, only attenuated endothelium-dependent vasorelaxation (Table 10.1) with no effect on vascular contractility. These general results further argue that, all types of high energy diets (high-fat, high-carbohydrate, or combination of both macronutrients), at least in the present work are directly or summatively involved in inducing vascular abnormalities. Interestingly, by analysing the results of studies in this project, potential role of obesity and atherogenic diets on the induction of vascular insulin resistance and subsequent endothelial dysfunction, leading to cardiovascular diseases and atherosclerosis becomes more evident. However, the degree of vascular abnormalities in response to long term consumption of a certain high energy diet may depend on the combination and/or amounts of the two atherogenic components (fat and carbohydrate) in any given diet. For example, in the present work, the greatest diversity in vascular defects was present in biscuit, and chocolate supplemented diets, where both diets contained high amounts of atherogenic macronutrients (Table 10.1).

Induction of vascular dysfunction by atherogenic macronutrient components of high energy diet is independent of BMI, and occurs well before appearance of any visible sign of obesity. For example, in chapter four, vascular abnormalities induced by high energy diet (chocolate supplementation) were occurred in the

absence of both hyperphagia and/or obesity. Similarly, in chapter three, long term consumption of biscuit diet induced the greatest variation in vascular functions without any remarkable obesity. These indicate that the induced vascular defects could be more related to the types of the macronutrient components in each diet rather than obesity *per se* or the amount of calorie intake (i.e hyperphagia). Therefore, relying on the amount of calorie intake as indicator of vascular defect would not be a plausible way to predict vascular abnormality. Moreover, these studies reveal that obesity *per se* might not have a major role in inducing vascular dysfunction. The presence of attenuated endothelium-dependent vasorelaxation as one of the vascular detrimental effects of the all four types of high energy diets in the present study may indicate that these types of the diets might have induced their vascular dysfunctions, at least partly, *via* involving the disruption and the inhibition of insulin action in vascular system by disturbing the levels of protein expression and/or activation of insulin signalling pathways in vasculature.

The highly-palatable diet was the only diet among the others, associated with a significant obesity, elevated plasma insulin level and subsequent systemic insulin resistance, in addition to attenuated endothelium-dependent vasorelaxation (Table 10.1 & 10.2), indicating a major role for palatable diet in inducing systemic insulin resistance and metabolic changes. Vascular insulin resistance might also be induced by atherogenic components of the diet itself. However, elevated levels of plasma triglyceride in these animals are the principal link to vascular dysfunction. Once again, these studies reveal that combined macronutrients of high fat and carbohydrate are mostly associated with a significant obesity. Nonetheless, long term consumption of any type of high energy diets resulted in synchronisation of both; elevated levels of plasma triglyceride and attenuated endothelial-dependent vasorelaxation. These parallel events may indicate that triglyceride have a major role in disturbing insulin action in vasculature (vascular insulin resistance) with subsequent vascular defect (attenuated endothelium-dependent vasorelaxation) (Table 10.1 & 10.2).

At the molecular level, protein expressions of PI 3-kinase and MAP kinase pathways were demonstrated in three different insulin-sensitive organs namely,

vascular system, liver, and skeletal muscle. The presence of local insulin resistance in each of these organs might lead to induction of a particular systemic disease. For example vascular insulin resistance can cause endothelial dysfunction and subsequent cardiovascular disease and atherosclerosis, liver insulin resistance can induce fatty liver problems such as non-alcoholic fatty liver disease, while insulin resistance in skeletal muscle may cause systemic insulin resistance and subsequently type II diabetes. Unfortunately, exact molecular mechanisms linking local insulin resistance with the clinical disease still not well understood.

In the aorta, obesity-inducing palatable diet caused selective changes in protein levels of PI 3-kinase-dependent signalling pathway. The reduced levels of PI 3-kinase may have a role in inducing insulin resistance in vascular system, contributing to increased incidence of cardiovascular events seen in obese subjects. Chronic withdrawal of obesity-inducing diet caused a complete normalisation of PI 3-kinase, while fenofibrate treatment failed to improve PI 3-kinase concentrations in dietary-obese animals, indicating that lifestyle modification may have a major role in combating diet-induced obesity-related cardiovascular dysfunction. In the skeletal muscle, protein expression of the regulatory subunit p85 of PI 3-kinase was augmented and were reversed by physiological but not pharmacological manipulation of obesity. The protein expressions of Shc and ERK1/2 in MAP-kinase pathway did not change in vascular and hepatic tissues, but were increased in skeletal muscle. Similarly, in the liver, reduction in protein levels of IRS-2 and PI 3-kinase were the only detrimental effects induced by dietary obesity, and these defects are suggested to be the principal causatives in induction of hepatic insulin resistance. Furthermore, disturbed levels of circulating and hepatic content of triglyceride may have crucial role in insulin resistance. Nonetheless, these molecular abnormalities are reversible by physiological but not pharmacological modification. Moreover, the elevated levels of plasma and liver triglyceride in addition to hepatic insulin resistance in diet-induced obesity might have a major role in inducing the detected histological features of non-alcoholic fatty liver disease, which were normalised by both physiological and pharmacological reversals of the obesity.

Chronic removal of highly-palatable diet for seven weeks was relatively the best therapeutical approach for obesity, as it completely reversed all diet-induced metabolic changes, increased lipid profile, insulin resistance, and vascular defects. Pharmacological treatment (fenofibrate administration), despite improving metabolic and insulin resistance, had very limited beneficial effects on vasculature. In general, it is well known that the application of non-pharmacological approaches such as "lifestyle modification" for treating obesity is both easy and financially more acceptable. The present work provides evidence that pharmacological intervention is much less preferable than non-pharmacological approaches in treating obesity, because of the undesirable effects that produced by the pharmacological agents. Non-pharmacological treatment "withdrawal of highly-palatable diet " reduced obesity by removing the source of obesity, and thus reducing the ingestion of high calorie intake, leading to near perfect reversal and normalisation of all altered physiological functions and mechanisms. Pharmacological intervention involves inhibition or acceleration of certain biochemical steps or functions that lead to increased accumulation of harmful end-products. It has been reported that fenofibrate completely improves the vascular dysfunction (Liang *et al.*, 2000; Steiner *et al.*, 2001). However, in the present work, fenofibrate administration was associated with molecular defects, and limited vascular improvement. It is possible that fenofibrate-induced side-effects are result of ROS production *via* accelerating the oxidation of lipid profile. Hence, it is plausible to recommend application of lifestyle modification such as reducing calorie intake for more convenience and satisfactory result, for treating obesity and its associated complications.

Conclusion. Long term consumption of a high calorie diet could be the major causative agent in inducing vascular defects leading to cardiovascular disease and atherosclerosis, regardless of obesity induction, or the amount of calorie intake. However, the consumption of any type of high calorie diets effects vascular functions. Moreover, the diversity of vascular dysfunction-induced by any high energy diet might not depend on the amount of ingested calorie or diet-induced obesity *per se*, rather it may depend on the combination of atherogenic macronutrients such as carbohydrate and fat in a given diet. The major effects of

the diet were on endothelium-dependent vasorelaxation, probably involving triglyceride-induced vascular insulin resistance.

Chocolate bars (Galaxy) supplementation for long period in addition to the chow-diet may induce wide range of vascular defects (increased vasoconstrictions in response to both KCl and NA, and attenuated endothelium-dependent vasorelaxation), while positive effects of fenofibrate treatment on vascular function may be due to its ability to reduce TG levels.

At the molecular level, it is plausible to conclude that: diet-induced obesity in combination with a highly-palatable diet results in significant alterations in the levels of protein expressions of both PI 3-kinase and MAP kinase pathways in the three different insulin-sensitive organs (vascular system, liver, and skeletal muscle). Furthermore, some of these alterations in protein levels are principally participated in triggering the induction of local and systemic insulin resistance. PI 3-kinase, in particular appears to be the key controlling step of insulin transduction in these organs. Moreover, abnormalities in expression of PI 3-kinase could be the main trigger for the induction of insulin resistance in these organs.

In diet-induced obese animals, MAP-kinase pathway does not appear to be deficient in insulin-sensitive tissues, thus allowing more mitogenic response, ultimately promoting the pathological progression. Moreover, the protein expressions of the early steps of insulin signalling were intact, indicating that the molecular defects in insulin transduction are post receptor. Moreover, differences in the molecular effects of diet-induced obesity on insulin-sensitive tissues may depend on the nature of the cellular function(s) in each organ. For example, in vascular tissue the protein expression of eNOS plays a major role in producing NO and subsequent facilitation of the vascular functions (endothelium-dependent vasorelaxation). Therefore, deficient PI 3-kinase pathway leads to abnormal eNOS expression and activation causing reduced NO production and subsequent attenuation of vasorelaxation.

Long term consumption of any type of high energy diet may not induce obesity but it generally associated with a wide range of vascular abnormalities, which is

depended on the type of the macronutrient present in the diet. In the present work, two approaches of obesity treatment were included for investigation. However, the sum of these studies have shown clearly that lifestyle modification, which represented in this work by removal of highly-palatable obesity-inducing diet for treating obesity, has no pronounced side effects. Moreover, it is a safe method, does not induce detrimental effects on the various physiological functions of the body such as: metabolic parameters, vascular functions, and molecular mechanisms (protein expressions), and has convenient and satisfactory results.

Final comment. Obesity is a neglected epidemic disease that is frequently thought of it as a self-inflicted condition. Unfortunately, it can represent as a network or a group of pathological illnesses and symptoms that grow up, spread, and detrimentally effects every organ of the body. Therefore, I would like to have this opportunity to express my recommendations and advices for over weight, obese, and non-obese high calorie diet loving individuals, informing them that obesity is a complicated and harmful than it appears, especially at vascular, and metabolic levels, advising them to rethink carefully and have a more healthy lifestyle. It is plausible to change lifestyle by modifying eating behaviour such as by decreasing calorie intake, avoiding overfeeding, and applying more healthy, convenient, and natural therapeutical mechanisms to reduce body weight. I would also like to advise those apparently healthy non-obese consumers of high calorie diets such as biscuit, chocolate, and cakes to reconsider the pathological “vascular” hazards of such diets, regardless of whether they are obese or not. These diets are capable of exposing individuals to incidence of cardiovascular diseases and atherosclerosis even in the absence of high BMI or a significant obesity. Last advice will be for both groups of individuals, to perform a regular medical checkup, especially of the cardiovascular system, liver function tests, and metabolic functions. By doing this medical checkup as early as possible, they could save their vital organs from a permanent pathological damage. Finally, it is definitely the time for all categories of people all over the world, to unite, combat and defeat such silent devastating monster, namely the obesity.

Table 10.1 Relationship between the type of high-energy diet and the type of vascular dysfunction

| Type of high energy diet | Type of vascular dysfunction | | | |
|--------------------------|---|----------|---|----------|
| | increase vasoconstriction in response to: | | Attenuation of vasorelaxation In response to: | |
| | KCl | NA | CCh | SNP |
| Biscuits | Sig. | Sig. | Sig. | Sig. |
| High-fat | Not sig. | Not sig. | Sig. | Not sig. |
| Chocolate | Sig. | Sig. | Sig. | Not sig. |
| High-palatable | Not sig. | Not sig. | Sig. | Not sig. |

KCl = potassium chloride, NA = noradrenaline, CCh = carbamylcholine, SNP = sodium nitroprosside, Sig. = significant increase or attenuation, Not sig. = not significant

Table 10.2 Relationship between metabolic changes and the different types of high energy diets

| Type of high energy diet | Metabolic parameters "Plasma levels" | | | | | |
|--------------------------|--------------------------------------|---------|--------|----|--------|------------|
| | Glucose | Insulin | leptin | TG | NEFA | HOMA INDEX |
| Biscuits | normal | normal | normal | ↑ | normal | normal |
| High-fat | normal | normal | normal | ↑ | ↑ | normal |
| Chocolate | normal | normal | ↑ | ↑ | ↑ | normal |
| High-palatable | normal | ↑ | ↑ | ↑ | normal | ↑ |

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Appendix

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Differential vascular dysfunction in response to diets of differing macronutrient composition: a phenomenological study

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Brief Communication

Differential vascular dysfunction in response to diets of differing macronutrient composition: a phenomenological study

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Abstract

Background: Vascular dysfunction can develop from consumption of an energy-rich diet, even prior to the onset of obesity. However, the roles played by different dietary components remain uncertain. While attempting to develop models of obesity in a separate study, we observed that two high-energy diets of differing macronutrient compositions affected vascular function differently in overweight rats.

Methods: Male Wistar rats ($n=6/\text{group}$) were fed diets providing varying percentages of energy from fat and carbohydrate (CHO). For 10 weeks, they were fed either chow, as control diet (10% of energy from fat; 63% from CHO), chow supplemented with chocolate biscuit (30% fat; 56% CHO) or a high-fat diet (45% fat; 35% CHO). Blood concentrations of biochemical markers of obesity were measured, and epididymal fat pads weighed as a measure of adiposity. Mesenteric arteries were dissected and their contractile and relaxant properties analysed myographically. Data were tested by analysis of variance (ANOVA).

Results: Weight gain and plasma concentrations of glucose, insulin and leptin were similar in all groups. However, biscuit-fed animals showed increased food intake (+27%; $p<0.01$) and elevated concentrations of TGs and non-NEFAs (+41% and +17%; both $p<0.05$). High-fat-fed animals showed an increase only in NEFAs (+38%; $p<0.01$). Arterial vasoconstriction in response to NA and KCl increased only in biscuit-fed rats (both $p<0.01$), while vasorelaxation in response to CCh and SNP, but not histamine, was attenuated in both groups (both $p<0.01$). Furthermore, whereas the effect of the high-fat diet was most pronounced in endothelium-dependent vasorelaxation, the biscuit diet had the greater effect on endothelium-independent vasorelaxation.

Conclusions: Vascular dysfunction resulting from consumption of a high-fat or combined relatively high-fat/high-CHO diet occurs through different physiological processes, which may be attributable to their differing macronutrient compositions. Combining potentially atherogenic macronutrients induces more extensive vascular impairment than that of high-fat alone, and may be attributable to the more marked dyslipidaemia observed with such a diet. Thus, these findings help clarify the role of dietary components in vascular impairment, which has implications for clinical approaches to preventing cardiovascular disease.

Background

The link between obesity and vascular dysfunction is well-established [1-8], but its causes remain uncertain. Of the circulating factors increased in obesity – including leptin, insulin, NEFAs and TGs [5, 6, 9-14] – only the latter two are raised by short-term high-energy feeding (prior to obesity onset), suggesting they may play the more critical role [15-18]. Dissecting out the relative importance of dietary macronutrients in atherogenic dyslipidaemia is complex [19], but it is believed that carbohydrate plays the primary role by stimulating insulin secretion, downstream of which fatty acid metabolism is determined [20, 21]. The mechanisms by which these dietary constituents' effects alter vascular function, particularly in animal models, is less clear, although some attempts have been made by different methods in humans [16, 22]. In view of this, we took the opportunity to investigate these mechanisms during an exploratory study in which we compared a number of high-energy diets for their abilities to induce weight gain in rats.

Methods

Male Wistar rats (192 ± 4 g) were fed diets of differing compositions for 10 weeks ($n=6/\text{group}$). Controls received standard chow (CRM Biosure, Cambridge, UK; 'chow' group). One experimental group received chow supplemented with chocolate biscuit (McVitie's 'HobNobs', Ashby de la Zouch, Leics, UK; 'biscuit' group), and the other received a fat-enriched diet (Research Diets, Inc, New Brunswick, NJ, USA; 'high-fat' group; Table 1). Food intake was measured daily and body weight weekly.

At termination, fasting blood samples were collected and plasma glucose, NEFA and TG concentrations determined by diagnostic kit, and insulin and leptin by RIA. A

single epididymal fat pad was dissected from each animal and weighed as a measure of adiposity. Length-tension characteristics of mesenteric arteries (6/rat) were determined, followed by evaluation of arterial contractility and relaxation, as described previously [18].

Before statistical analysis, vascular reactivity data were quantified as AUC. Experimental groups were compared on all parameters to controls by one-way ANOVA followed by *post hoc* analysis.

Results

Biscuit-fed rats showed a higher daily energy intake and increased TG concentrations. The high-fat diet increased NEFAs by twice the proportion of the biscuit diet. There were non-significant rises in body weight and epididymal fat mass on both experimental diets, but neither affected glucose, insulin or leptin concentrations (Table 2).

The experimental diets did not alter vessel diameter (data not shown; both $p > 0.05$ vs chow). Both KCl and NA, however, increased vasoconstriction in the biscuit group ($p < 0.01$ and $p < 0.001$), but had no effect in the high-fat group (both $p > 0.05$; Fig. 1).

In response to cumulative addition of CCh (10 nM-100 μ M), NA-pre-constricted arteries from biscuit- and high-fat-fed rats displayed a significant rightward shift of concentration-response curves, compared with controls (EC_{50} biscuit: 0.28 ± 0.01 ; high-fat: 0.52 ± 0.04 μ M; both $p < 0.001$ vs chow: 0.11 ± 0.01 μ M; Fig. 2a). Maximal vasorelaxation to 100 μ M CCh was reduced by 13% in both biscuit- and high-fat-fed

rats (both $p < 0.01$ vs chow-fed), but was more attenuated in the latter at lower CCh concentrations (3.16-316 nM; $p < 0.01$ vs biscuit; Fig. 2a). Histamine-induced vasorelaxation was similar in all groups (Fig. 2b).

The endothelium-independent vasodilator, SNP, also induced concentration-dependent vasorelaxation in arteries from all groups, the curves shifted to the right in the biscuit- and high-fat-fed groups (EC_{50} biscuit: 1.58 ± 0.02 ; high-fat: 0.24 ± 0.01 ; both $p \leq 0.01$ vs chow: 0.13 ± 0.02). However, maximum SNP-induced responses were similar in all groups. Compared with the chow- and high-fat-fed groups, SNP-induced relaxation in biscuit-fed rats was attenuated by more than 2-fold (both $p < 0.001$). Attenuation of SNP responses was also significant, though less pronounced, in high-fat-fed animals (vs control: $p < 0.05$; Fig. 2c).

Discussion

This study confirms previous observations that high-energy diets lead to vascular dysfunction [5, 6, 17, 18, 23, 24]. The high-fat and biscuit diets induced similar degrees of modest weight gain and adiposity despite different macronutrient compositions, suggesting these account for the differences observed in vascular reactivity. The biscuit diet had a detrimental effect on both vasoconstriction, and – relaxation, whereas the high-fat diet affected only the latter. The attenuation of vasorelaxation in response to the cGMP-mediated actions of both CCh and SNP shows that this effect of both diets is mediated through both endothelium and vascular smooth muscle. Consumption of other high-energy diets has shown this dual process [5, 17, 18], although some such diets exert their effects only through one mechanism [23], further highlighting the specificity of effects depending on dietary components.

The failure of either diet to alter cAMP-mediated vasorelaxation (i.e., histamine-induced responses) also appears to be a common feature of high-energy diets [17, 18].

Intriguingly, the degree of abnormality seen in both endothelium- and smooth muscle-mediated vasodilation also differed according to diet. CCh-induced vasorelaxation was more severely attenuated by the high-fat diet, and the SNP response more so by the biscuit diet. These findings suggest that different dietary components affect vasorelaxation by different processes, that a diet high in fat has greater deleterious effects on the arterial endothelial lining, whereas a diet relatively rich in carbohydrate *and* fat largely compromises the vascular smooth muscle. There is substantial supporting evidence that dietary components play a role in vascular function. Humans and animals fed diets high in fat, particularly saturated fat [19, 24-26], show endothelial abnormalities. On the other hand, reports of the protective effects of high-carbohydrate (low-saturated fat) diets and diets high in unsaturated fats are contentious [16, 19, 22, 27, 28]. The biscuit diet, although lower in both total and saturated fat than the high-fat diet, contained a significantly greater amount of MUFAs. Although these are associated with improved cardiovascular risk [27, 29], at high levels they can acutely impair endothelial function [16]. Recent reviews of the literature note that stimulation of insulin release by increased carbohydrates promotes adipogenesis, weight gain and atherogenesis, all associated with the metabolic syndrome [19, 28, 30]. The biscuit diet was higher in overall carbohydrate content than the high-fat diet, and it may be this, in combination with relatively high fat content, that accounts for its more detrimental effects. The relatively lower saturated fat and higher carbohydrate content may also be more palatable and account for the increased daily intake in the biscuit-fed animals, and, hence, epididymal adiposity

(though short of significance) and overall greater hyperlipidaemia. This would have to be more carefully analysed in future, however, by matching intake volume and overall energy content between groups. An adipocentric view would suggest that the increased adiposity *per se* of biscuit-fed animals is critically linked with the worsened vascular defects in this group, with excessive fat mass in this depot resulting in fat cell dysfunction, which in turn contributes to the metabolic disorders that increase the risk of atherosclerosis [31].

High-carbohydrate meals may also stimulate sympathetic nervous system activity *in vivo* [32], and counteract insulin-induced vasodilation. Although fasting insulin levels were similar in all groups, it is possible that differences in post-prandial insulin (which would be expected to be higher in rats fed a carbohydrate-enriched diet and resulting adipogenesis) and then raised plasma TGs, resulting in part from *de novo* lipogenesis, may account for the vascular differences. Indeed, hypertriglyceridaemia is a recognized atherogenic risk factor (e.g., [5, 19, 33]). This can be the case even in the absence of insulin resistance [5] or symptoms of atherosclerosis [34]. Although our animal model did not adequately mimic human obesity in terms of weight gain and some of its metabolic disturbances – probably due in part to a lack of statistical power - it remains predictive of vascular dysfunction in the absence of these and therefore further highlights the insidiousness of the disorder.

Finally, although % energy derived from protein was lower in both biscuit and high-fat diets compared to chow, the total average daily protein intake was similar (67 g and 78 g/day, respectively; biscuit *vs* high-fat: $p>0.05$). Hence, it is unlikely that protein deficiency explains the differences in vascular function. Dietary antioxidant

levels may also play a role (e.g., [35, 36]), but it is not possible to comment further on these as they were not measured.

Conclusion

The limitations of this study, therefore, revolve around the fact that the nutrient contents of the diets were not sufficiently controlled to draw very precise conclusions regarding their comparative effects. We cannot know whether their differential effects on vascular function were direct effects of nutrient content or indirect effects of resulting adiposity and/or circulating lipids. However, this study does show that high-energy diets of varying compositions can induce vascular dysfunction to varying degrees in the rat via mechanisms involving different layers of the vascular wall. The combination of high-fat and high-carbohydrate diets may be particularly damaging, possibly through increased hyperlipidaemia.

Abbreviations

| | |
|--------|----------------------------|
| ANOVA: | analysis of variance |
| AUC: | area under the curve |
| CCh: | carbamylcholine |
| CHO: | carbohydrate |
| KCl: | potassium chloride |
| MUFA: | monounsaturated fatty acid |
| NA: | noradrenaline |
| NEFA: | non-esterified fatty acid |
| PUFA: | polyunsaturated fatty acid |
| RIA: | radioimmunoassay |

TG: triglyceride
SNP: sodium nitroprusside

Competing interests

The authors declare that they have no competing interests.

Authors' contributions

SF carried out the myography and participated in tissue collection, statistical analysis and drafting of the manuscript.

LCP calculated the diet composition and participated in data collection, tissue sampling, statistical analysis and drafting of the manuscript.

CJS participated in data collection, tissue sampling and statistical analysis, and carried out metabolic assays.

JAH participated in data collection, tissue sampling and statistical analysis, and carried out metabolic assays.

RC participated in data collection and tissue sampling.

JPHW participated in drafting of the manuscript.

EKNA participated in the conception and design of the vascular study, the myography and in drafting of the manuscript.

All authors read and approved the final manuscript.

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Figure legends

Figure 1 - Arterial contractility

Figure 2 – Arterial relaxation

Table 1 Dietary breakdown of macronutrients (% of energy)

| Diet | Total CHO | Complex CHO | Simple CHO | Total Fat | Saturated fat | MUFA | PUFA | Total protein |
|-----------|-----------|-------------|------------|-----------|---------------|------|------|---------------|
| Chow | 63.0 | 57.7 | 5.3 | 10.3 | 1.7 | 3.0 | 5.6 | 28.1 |
| 'Biscuit' | 55.9 | 37.2 | 18.7 | 30.0 | 12.9 | 10.6 | 5.4 | 15.0 |
| High-fat | 35.0 | 16.9 | 16.9 | 45.0 | 39.1 | 0.0 | 5.5 | 20.0 |

Table 2 Body weight, energy intake and metabolic parameters

| | Chow | Biscuit | High-fat |
|----------------------------|-----------|-------------|-------------|
| Daily energy intake (kJ/d) | 361±15 | 459±14** | 390±13 |
| Body weight gain (g) | 299±22 | 347±15 | 339±22 |
| Epididymal fat mass (g) | 4.5±0.6 | 7.7±0.8 | 6.6±2.2 |
| Glucose (mM) | 9.8±1.0 | 8.9±1.7 | 9.3±1.0 |
| Insulin (µU/ml) | 22.2±2.6 | 21.1±1.2 | 19.4±1.6 |
| Leptin (ng/ml) | 5.3±0.9 | 6.3±0.3 | 5.9±0.4 |
| Triglycerides (mg/dL) | 89.4±4.8 | 126.5±10.1* | 77.7±3.6 |
| NEFAs (mM) | 0.29±0.01 | 0.34±0.01* | 0.40±0.02** |

* $p < 0.05$; ** $p < 0.01$ vs chow

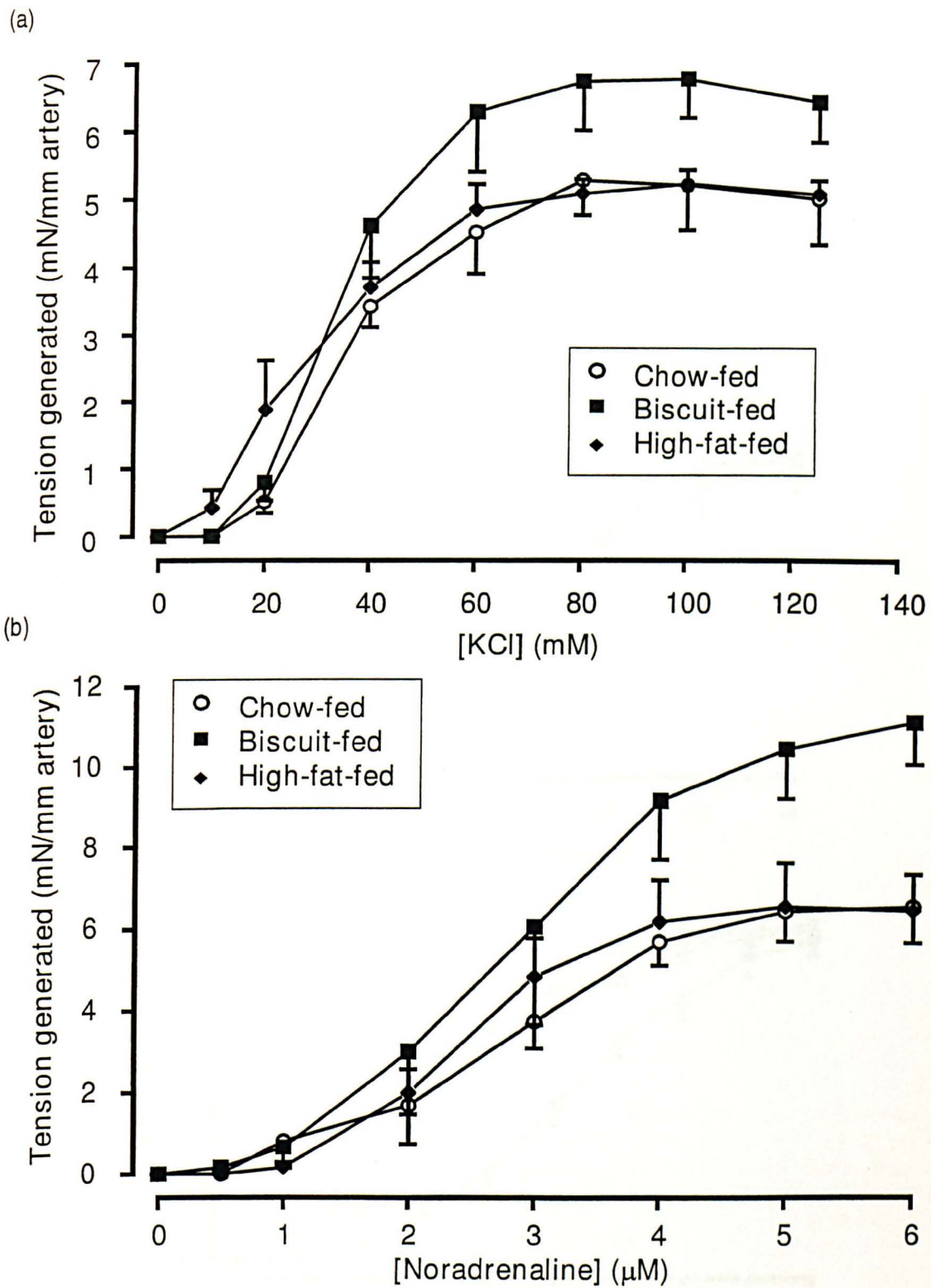


Figure 1

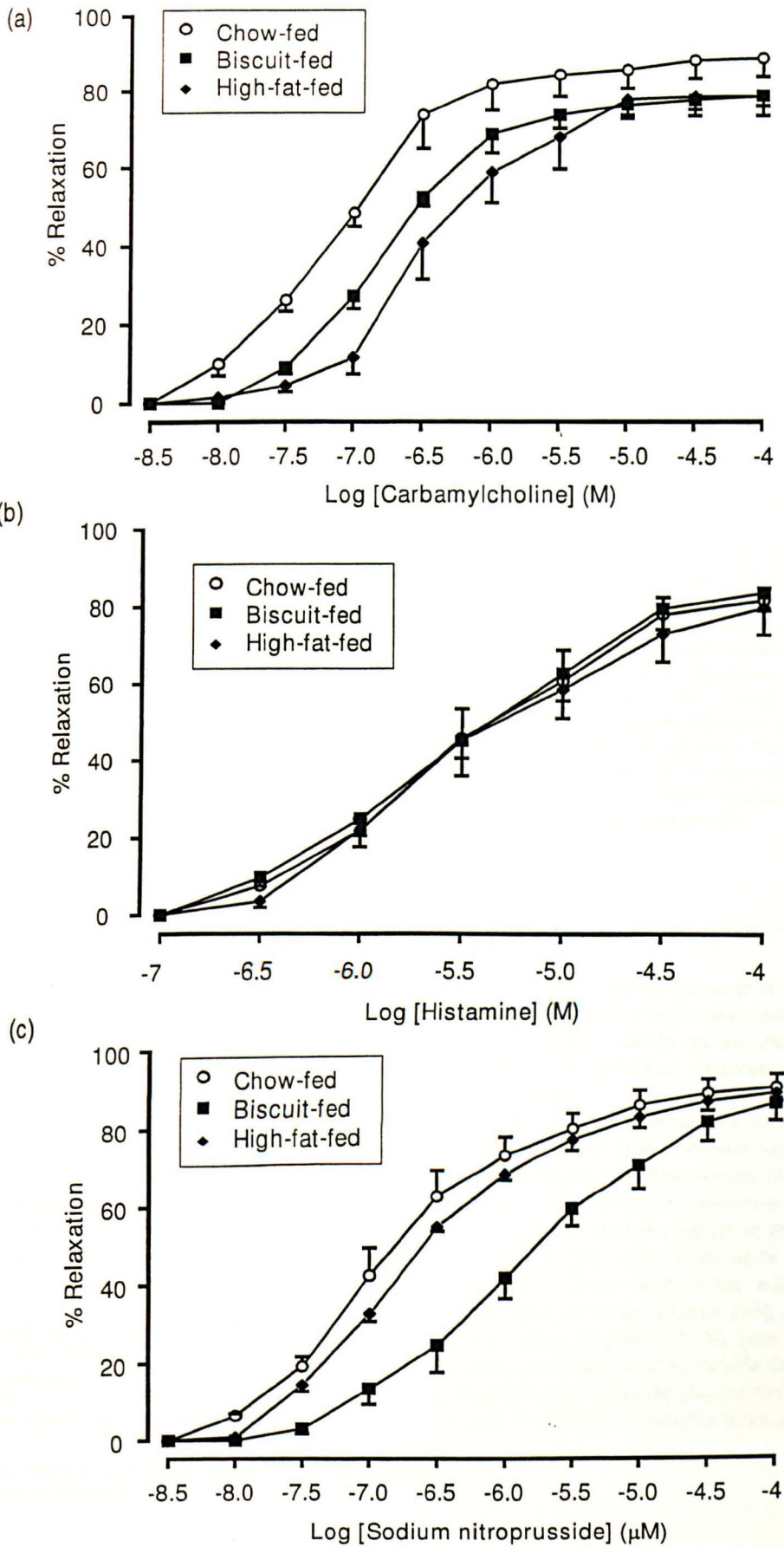


Figure 2

Chronic withdrawal of a high-palatable obesity-inducing diet completely reverses metabolic and vascular abnormalities associated with dietary-obesity in the rat[☆]

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Abstract

Chronic consumption of a high-palatable diet induces obesity and markedly impairs arterial relaxation. We have recently reported that endothelial function is only partially restored after acute withdrawal of palatable diet. Therefore, this study was designed to investigate the effects of chronic withdrawal of high-palatable obesity-inducing diet on metabolic and vascular function in rats. Wistar rats were fed either standard laboratory chow throughout (controls) or given a highly-palatable diet (diet-fed) for 15 weeks; or fed the diet for 8 weeks and then returned to chow (diet-to-chow) for further 7 weeks before sacrifice. Diet-fed rats had higher body weight, fat mass, liver and heart weight than both chow-fed and diet-to-chow groups ($P < 0.01$ for all). Compared with chow-fed and diet-to-chow groups, diet-fed rats had significantly raised fasting plasma levels of insulin, leptin and triglycerides levels (each +180%; $P < 0.0001$), but not glucose or non-esterified fatty acids. There were no significant differences between any metabolic parameters between chow-fed and diet-to-chow groups. Mesenteric arteries showed no significant differences between any groups in KCl-induced tension generation, while diet-fed groups had significantly higher noradrenaline-induced vasoconstriction than both chow-fed and diet-to-chow groups. Maximum endothelial-dependent vasorelaxation responses to carbamylcholine (CCh) were significantly (by 23%; $P < 0.001$) attenuated in the diet-fed group. This defect was abolished in the diet-to-chow group. There were no significant differences in endothelium-independent vasorelaxation responses to sodium nitroprusside between the three groups. In conclusion, palatable diet induces obesity and metabolic abnormalities as well as a marked endothelial dysfunction. These abnormalities are completely reversed by chronic withdrawal of the obesity-inducing high-palatable diet.

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Keywords: Diet; Obesity; Insulin resistance; Triglycerides; Vascular function

1. Introduction

Throughout the world, consumption of high-energy diets together with a general reduction in the levels of physical activity are causing an alarming health problem, namely obesity. Obesity is a major cardiovascular risk factor and the prevalence of atherosclerotic disease in obese humans has been recognised for more than two decades [1,2]. An association has been described between obesity, arterial hypertension, insulin resistance and dyslipidaemia, which comprise core features of the 'metabolic syndrome' or Syndrome X

[3]. Many obese patients accomplish weight loss with diet, exercise and lifestyle modification, achieving substantial reductions in central obesity and associated disturbances corresponding to a significant decrease in cardiac events and mortality [4–6].

Obesity induces abnormalities of arterial function and structure in human and in certain animal models that reflect some aspects of human obesity [4,7–9]. These include alterations of the release of vasoactive substances from endothelial cells as well as changes in arterial wall elasticity. A low-caloric diet improves [9], while elevating the plasma lipid profile further deteriorates endothelium-dependent vasodilatation in obese subjects [10], and in patients with coronary artery disease [11]. We have also recently shown that acute consumption of a palatable diet causes endothelial dysfunction long before development of any significant obesity [12]. Moreover, endothelial function was only partially

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corrected with short-term withdrawal of the palatable diet [13]. Thus, in this study we aimed to investigate whether diet- and obesity-induced endothelial abnormalities are fully reversible after chronic withdrawal of the high-energy, palatable diet.

2. Methods

2.1. Animals and experimental protocol

All procedures were approved by the Institutional Committee and accord with current UK legislations. Adult male Wistar rats ($n = 18$) were randomised and assigned to a control group ($n = 6$) and a test group ($n = 12$). All animals had free access to water and were housed individually under controlled environmental conditions (19–22 °C; 30–40% humidity) and a 12 h light–dark cycle (lights on at 07:00 h). Controls were fed a standard laboratory pelleted diet (chow-fed; CRM Biosure, Cambridge, UK), while test group had free access to a highly-palatable diet consisting of (by weight) 33% ground pellet diet, 33% Nestlé condensed milk, 7% sucrose and 27% water. Chow-fed controls remained on their prospective diet for 15 weeks, while after 8 weeks, palatable-diet-fed animals were subdivided into two groups. Palatable diet was removed from one group and the standard chow diet was reintroduced (diet-to-chow) for further 7 weeks, while the remaining subgroup was maintained on the palatable diet (diet-fed) for another 7 weeks. After 15 weeks from the start of the study, rats were killed by CO₂ inhalation after 2 h of fasting, and the total body fat mass was measured using the Bioimpedance method using the TOBEC[®] equipment (Biotech Instrument Ltd.).

2.2. Biochemical analyses

Trunk blood was removed into cold heparinized tubes and the gonadal and perirenal fat pads, as well as the gastrocnemius muscles and heart were dissected and weighed. The plasma was immediately separated by centrifugation before being frozen for later measurements of blood analytes (glucose, insulin, leptin, non-esterified free fatty acids (NEFA) and triglycerides (TG)). Plasma glucose concentration was determined using a glucose oxidase method, and NEFA and TG concentrations were measured using commercial diagnostic kits (Boehringer Mannheim, and Sigma Diagnostics.). Insulin and leptin concentrations were measured by radioimmunoassay (RIA) kits (Pharmacia/Upjohn Diagnostics, and Linco Research, respectively).

2.3. Assessment of vascular contractility

Six third-order mesenteric arteries (>200 µm diameter and 2 mm lengths) were carefully dissected from each animal. Each artery was freed of fat and connective tissue and mounted on two 40 µm diameter stainless-steel wires

in an automated myograph (Cambustion Ltd.). The vessels (in duplicate) were incubated in a 5 ml organ bath containing physiological salt solution (PSS; composition [in mM]: NaCl 119, KCl 4.7, CaCl₂ 2.5, MgSO₄ 1.17, NaHCO₃ 25, KH₂PO₄ 1.18, EDTA 0.026 and glucose 5.5) gassed with 95% O₂ and 5% CO₂ at 37 °C.

After 30 min equilibration, the length–tension characteristics for each vessel were determined as described previously [14]. The computer also calculated the target tension that each vessel should develop in response to a maximal stimulus. Arteries were then allowed a further 30 min to equilibrate before being depolarised twice with a high-potassium physiological salt solution (KPSS, 125 mM) in which NaCl in normal PSS was replaced by an equimolar concentration of KCl. Cumulative concentration–response curves to either KCl (10–125 mM) or noradrenaline (NA, 0.5–6 µM) were then carried out. Any vessel failing to reach its predetermined target tension in response to vasoconstriction with KCl (125 mM) was discarded.

2.4. Assessment of vascular relaxation

To investigate defects in endothelial-dependent and -independent vasorelaxation, we measured relaxation induced in NA-precontracted arteries following exposure to endothelium-dependent (carbamylcholine (CCh)) or endothelium-independent (sodium nitroprusside (SNP)) vasodilators. To eliminate interference of vasoactive prostanoids, 10 mM indomethacin was added to the organ bath prior to contraction with NA (8 µM). When contraction reached a plateau after 2 min, concentration–response curves were carried out to either CCh or SNP (for both, 10 nM to 100 µM).

2.5. Homeostasis model assessment (HOMA)

HOMA, an index of insulin resistance, which employs measures of fasting plasma concentrations of glucose and insulin, was calculated according to the method described previously [15].

2.6. Data interpretation and statistical analyses

Vasoconstriction in response to NA and KCl were expressed as absolute force generated. Relaxation in response to CCh and SNP was calculated as the percentage reduction from the maximal tension generated in response to the supra-maximal concentration of NA (8 µM). Data are expressed as mean ± S.D. or where stated as S.E.M. for more clarity. An average response for all the vessels from a given animal was determined before group analysis. Data have normal distribution (Shapiro Wilk *W*-test). Statistical significance was tested using Student '*t*-test' or repeated-measures (ANOVA; Bonferroni *t*-test) or the Mann-Whitney *U*-test, as appropriate. Results were considered statistically significant at $P < 0.05$. Univariate correlation between maximal CCh-induced vasorelaxation and fasting terminal insulin,

leptin and TG levels were determined by linear regression followed by two-tailed analyses, using ARCUS PRO-STAT (version 3.23).

3. Results

3.1. Metabolic data

Animals giving palatable diet progressively gained more weight than their chow-fed counterparts. A significant difference in total body weight was observed after 5th week of feeding and further increased after 8th week where diet-fed animals had significantly ($>10\%$, $P = 0.0059$) higher total body weight than chow-fed controls (Fig. 1). Following the next 7 weeks, average weekly weight gain of diet-fed animals was 1.8-fold greater than chow-fed controls (18.43 ± 0.50 g versus 10.34 ± 0.23 g control), while average weekly weight gain of diet-to-chow animals was significantly lower ($P < 0.001$, 10.04 ± 0.06 g versus 18.43 ± 0.50 g diet-fed) than their diet-fed counterparts (Fig. 1). However, at the end

of the experiment, total body weight of diet-to-chow animals was still significantly ($P = 0.0431$) higher than control chow-fed but was markedly ($P < 0.0062$) lower than the diet-fed animals (Fig. 1).

At the end of the experiment, Bioimpedance measurements showed that diet-fed animals had 70 and 27% more total body fat than chow-fed and diet-to-chow groups, respectively. Diet-fed animals had significantly higher epididymal and perirenal fat pad mass than both chow-fed (by $>180\%$, $P < 0.0001$) and diet-to-chow (by $>175\%$, $P < 0.0002$) groups (Table 1). Diet-fed rats had significantly (for both, $P < 0.0230$) higher heart weights than both chow-fed and diet-to-chow rats (Table 1). There were neither significant differences between fat pad depots and heart weights of diet-to-chow and chow-fed groups, respectively; nor there were any significant differences in gastrocnemius muscle weights between the three groups (Table 1).

While fasting plasma glucose levels were comparable in all these groups (12.42 ± 1.12 mM chow-fed; 14.40 ± 0.362 mM diet-fed; 14.43 ± 0.42 mM diet-to-chow), diet-fed rats had significantly higher ($>130\%$, $P < 0.0093$) fasting

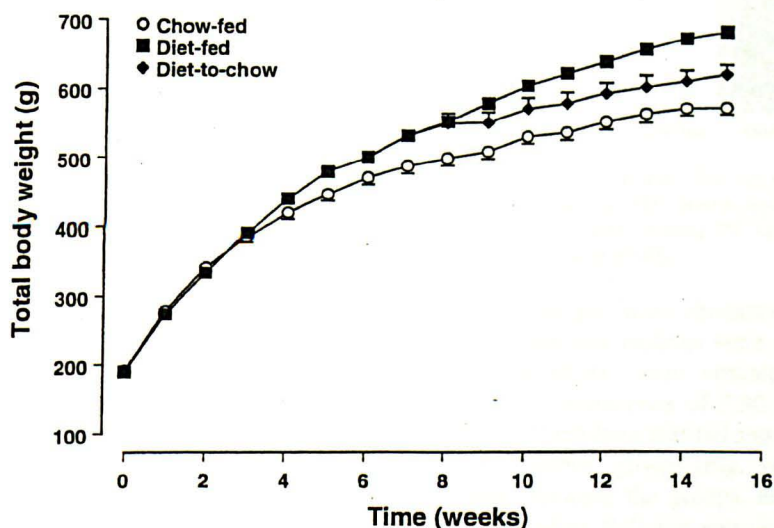


Fig. 1. The effect of a palatable-diet and its removal on total body weight in the rat. Animals were fed either standard chow (chow-fed) or a palatable diet (diet-fed) for 15 weeks, or fed the palatable diet for 8 weeks and then returned to standard chow for further 7 weeks (diet-to-chow). Diet-fed animals were significantly ($P < 0.00001$) heavier than both chow-fed and diet-to-chow groups. Diet-to-chow group was significantly heavier ($P < 0.04$) than chow-fed but lighter ($P < 0.006$) than diet-fed groups. Data are expressed as mean \pm S.E.M.

Table 1
Physiological and metabolic characteristics of the three experimental groups at the end of the experiment

| | Chow-fed ($n = 6$) | Diet-fed ($n = 6$) | Diet-to-chow-fed ($n = 6$) |
|-------------------------------|----------------------|------------------------|------------------------------|
| Total body weight (g) | 570.2 ± 9.7 | $680.2 \pm 8.7^{a,b}$ | 619.2 ± 7.7^a |
| Total fat pad mass (g) | 21.0 ± 1.8 | 40.5 ± 2.9^a | 24.1 ± 2.4 |
| Other fat mass (g) | 73.8 ± 6.9 | $121.3 \pm 11.9^{a,b}$ | 102.5 ± 6.6^a |
| Gastrocnemius muscle mass (g) | 3.2 ± 0.1 | 3.3 ± 0.1 | 3.2 ± 0.1 |
| Heart weight (g) | 1.7 ± 0.1 | $2.0 \pm 0.1^{a,b}$ | 1.7 ± 0.1 |

Data are mean \pm S.D. $P < 0.001$ for all.

^a Diet-fed and diet-to-chow-fed vs. chow-fed.

^b Diet-fed vs. diet-to-chow-fed.

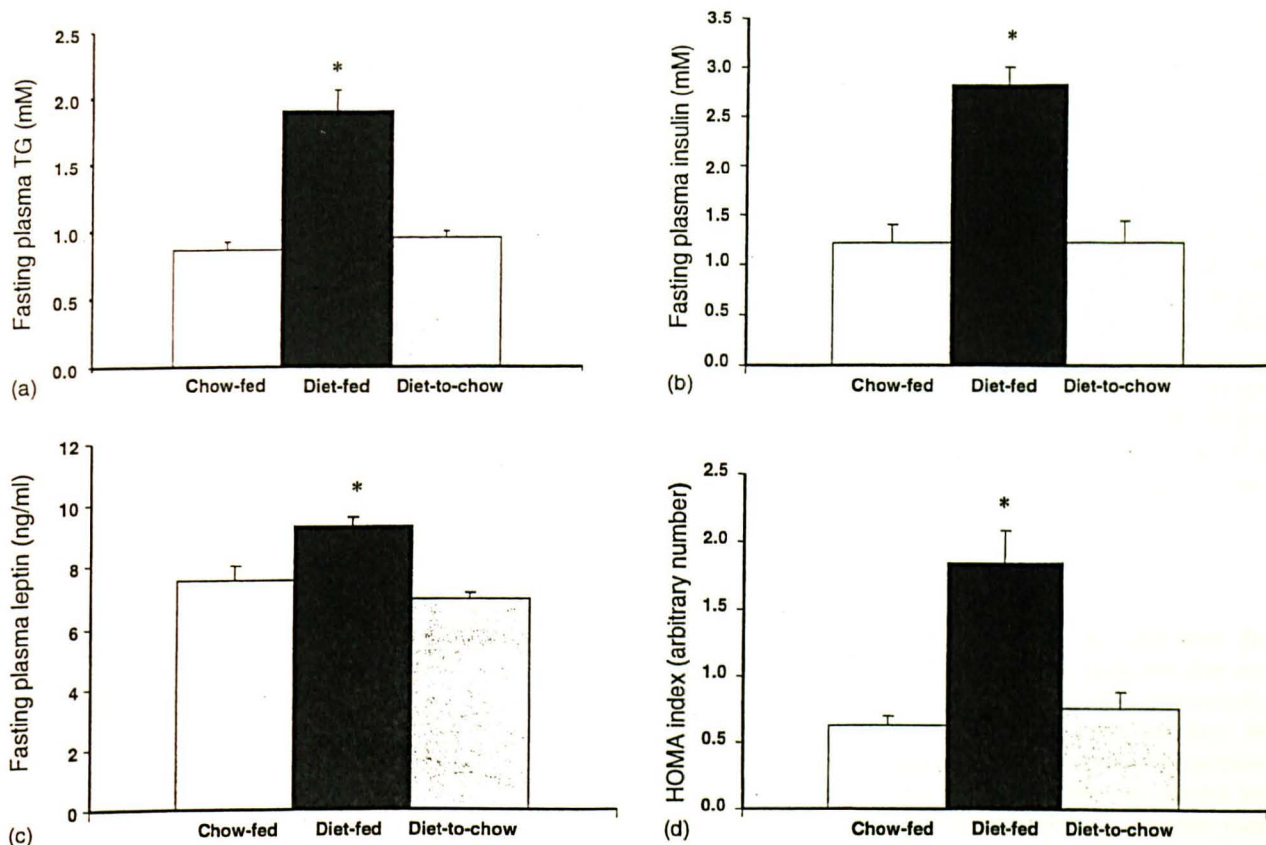


Fig. 2. Effects of highly-palatable diet feeding and its withdrawal on fasting plasma levels of: (a) triglyceride (TG), (b) insulin, (c) leptin, and (d) HOMA index in three experimental animals. Diet-fed rats had significantly (for all, $P < 0.01$) higher fasting (TG), insulin, leptin levels than both chow-fed and diet-to-chow groups. Diet-fed animals were also insulin resistance, indicated by a higher HOMA index. Fasting TG, insulin and leptin levels returned to normal levels by long-term removal of the palatable diet and insulin resistance was corrected subsequently.

terminal levels of plasma insulin than both chow-fed and diet-to-chow group. As a result, diet-fed rats had significantly (for both, $P < 0.001$) higher HOMA index, indicating development of insulin resistance in these animals (Fig. 2). Moreover, diet-fed rats had significantly higher (up to 35%, $P < 0.001$) plasma leptin levels than both chow-fed control and diet-to-chow groups (Fig. 2).

Fasting plasma levels of NEFA were comparable in all three groups (0.45 ± 0.24 mM chow-fed; 0.48 ± 0.41 mM diet-fed and 0.40 ± 0.24 mM diet-to-chow), while diet-fed rats had significantly (greater than two-fold, $P < 0.001$, for both) higher levels of fasting TG than both chow-fed and diet-to-chow groups (Fig. 2).

3.2. Vascular data

There were no significant differences in vessel diameter between the three groups (189 ± 20 μ m chow-fed, 200 ± 15 μ m diet-fed and 200 ± 10 μ m diet-to-chow; $P > 0.5$ for all).

3.3. Agonist-induced vasoconstriction responses

Cumulative concentration response curves (CCRC) to KCl (10–125 mM) and NA (0.5–6 μ M) in arteries from all

three groups produced characteristic sigmoid relationship. KCl-induced contractions were similar in all three groups such that CCRC were virtually superimposable, giving maximum contractions of 5.90 ± 0.41 mN/mm chow-fed, 6.16 ± 0.31 mN/mm diet-fed and 6.32 ± 0.26 mN/mm artery for diet-to-chow groups (Fig. 3a), with no significant differences between the groups, either overall (by ANOVA) or at any given KCl concentration. However, arteries from the diet-fed group showed greater contractile reactivity to NA administration, reaching significance (>30%, $P < 0.01$ for all) at 3 and 4 μ M NA than chow-fed and diet-to-chow groups (Fig. 3b).

3.4. Endothelium-dependent and -independent vasorelaxation

NA-precontracted arteries from chow-fed rats demonstrated progressive relaxation to cumulative addition of CCh (10 nM to 100 μ M), achieving a maximum of $90 \pm 2\%$ at a CCh concentration of 100 μ M, with EC_{50} values of 0.14 ± 0.02 μ M. Arteries from diet-fed rats that were similarly exposed to CCh displayed a significant rightward shift of CCRC compared with chow-fed control arteries (EC_{50} diet-fed: 0.75 ± 0.03 , $P = 0.0023$ versus chow-fed; Fig. 4a).

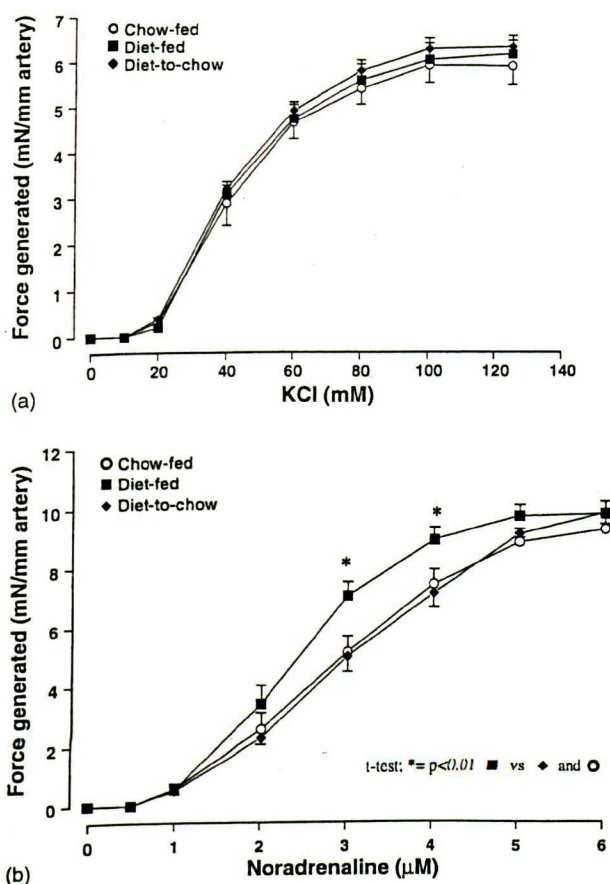


Fig. 3. Concentration–response curves for: (a) KCl (10–125 mM) and (b) noradrenaline (0.5–6 μM) in arteries from chow-fed, diet-fed and diet-to-chow-fed animals. Data represent mean \pm S.E.M. for six animals in each group. KCl-induced vasoconstrictions were similar between all three groups. However, 3 and 4 μM NA-induced contractions significantly ($P < 0.001$) were greater than both chow-fed and diet-to-chow-fed groups. Data represents mean and S.E.M. for $n = 6$ in each groups.

Maximal vasorelaxation to 100 μM CCh was reduced by 23% in diet-fed rats ($P = 0.0045$ versus chow-fed). However, CCh-induced vasorelaxation curves in arteries from diet-to-chow group were virtually identical to those in chow-fed group, achieving maximal relaxation of $95 \pm 1\%$ at 100 μM CCh, with EC_{50} values of 0.19 ± 0.04 μM.

SNP (10 nM to 100 μM) also induced concentration-dependent vasorelaxation of arteries from all three groups. Concentration-relaxation–response curves were similar and superimposable in all three groups (Fig. 4b), with no statistically significant differences between the groups, either overall (by ANOVA) or at any given SNP concentration.

3.5. Metabolic determinants of vascular dysfunction

Of the various blood analytes measured, insulin, leptin and TG showed significant increases in diet-fed animals. Therefore, the relationship between endothelial dysfunction (measured as the percentage reduction of 100 μM

CCh-induced vasorelaxation) and terminal plasma levels of insulin, leptin and TG were examined. In each case, univariate correlation analysis showed an inverse correlation between fasting terminal levels of TG ($r^2 = 0.4787$; $P = 0.0013$), insulin ($r^2 = 0.4237$; $P = 0.0023$) and leptin ($r^2 = 0.609$; $P = 0.0001$) and maximal CCh-induced vasorelaxation. We also analysed the relationship between total body weight and total fat pad mass with vascular reactivity. Univariate correlation analysis once again indicated an inverse correlation relationship between total body weight ($r^2 = 0.3751$; $P = 0.0043$) and fat pad mass ($r^2 = 0.3565$; $P = 0.0048$) and maximum carbamylcholine-induced vasorelaxation. Furthermore, analysis of blood analytes showed significant positive correlations between fasting plasma insulin ($r^2 = 0.3043$; $P = 0.0025$) and leptin ($r^2 = 0.4189$; $P = 0.0020$) with that of circulating triglycerides.

4. Discussion

The present study, demonstrates for the first time that chronic removal of an obesity-inducing palatable diet corrects all the metabolic abnormalities, cardiac hypertrophy and vascular dysfunctions associated with obesity. We have previously demonstrated that chronic consumption of a palatable diet induces significant level of obesity and markedly affects metabolic parameters and vascular reactivity in the absence of any visible insulin resistance [8]. Moreover, short-term feeding (2 days) of the same palatable diet significantly alters lipid profile and markedly attenuates endothelial function in the absence of any significant obesity [12]. These defects were only partially corrected by acute removal of the diet [13], indicating that diet per se may have a major role in the development of vascular abnormalities in animal models of obesity.

In this study, diet-fed animals developed marked obesity, which was characterised by more than two-fold increase in total body fat content as determined by Bioimpedance methods and measurements of fat pad mass. Obese animals had significantly higher heart weight and fasting plasma levels of insulin, suggesting cardiac hypertrophy and development of insulin resistance in dietary-obese animals, respectively. These findings are similar to those seen in human obesity [1–4]. In humans, obesity leads to dyslipidaemia and type 2 diabetes and hypertension [16–18], while changes in life style improve obesity-induced abnormalities to a certain degree [4,6,9].

A number of experimental studies have reported severe detrimental outcome of excessive high-energy diet on vascular reactivity. For example, high-fat diet impairs endothelium-dependent vasorelaxation [8,12–14,19,20] while antioxidant therapy improves endothelial function [21,22], suggesting a diet-induced oxidative stress in vasculature. In obesity, plasma antioxidant levels are decreased [21–23], whereas myocardial oxidative stress is increased [24,25]. These changes may, at least in part, explain

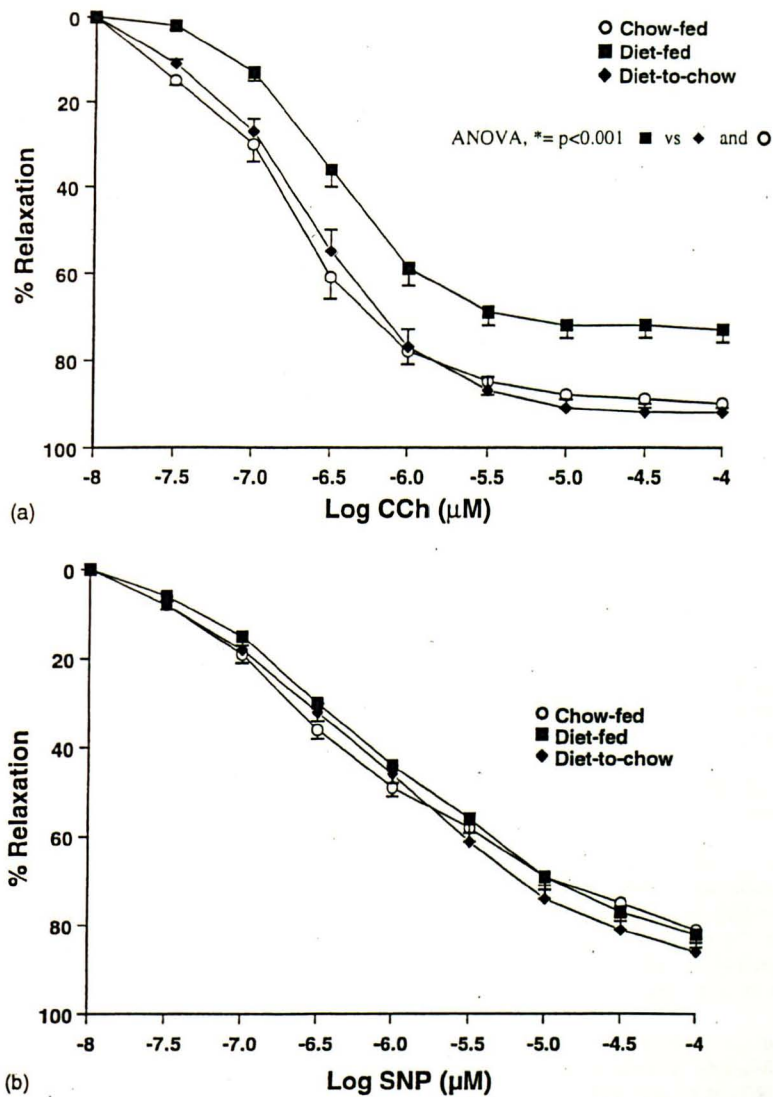


Fig. 4. Concentration-relaxation-response curves (mean \pm S.E.M.) of: (a) carbamylcholine (CCh), (b) sodium nitroprusside (SNP) in the three groups. Arteries were contracted with $8 \mu\text{M}$ noradrenaline. When contractions reached a plateau cumulative concentration responses to CCh or SNP were carried out. CCh-induced vasorelaxation was significantly reduced in diet-fed animals (ANOVA $P < 0.001$) in comparison to chow-fed group. This abnormality was completely abolished by the chronic removal of the palatable diet. SNP-induced vasorelaxations were similar in all three groups.

the increased risk of vascular disease and the abnormal endothelium-dependent vasorelaxation found in obese subjects and animals. In obese patients with hypertension, ACh-induced forearm blood flow (FBF) is significantly reduced compared to healthy individuals. Caloric restriction decreases body weight and the mean blood pressure, fasting plasma insulin, total cholesterol, triglycerides, low-density lipoproteins (LDL) as well as enhancing ACh-induced FBF, while nitric oxide synthase inhibitors mask increase in ACh-induced FBF, indicating that reduction in energy intake improves endothelial function possibly by increasing endogenous NO production in obese hypertensive patients [9]. Lipid lowering also reduces the risk for coronary events, cardiovascular morbidity and mortality in a broad range of patients [26]. Similar observations have also been

reported in animal studies. In miniature pigs, 6–9 month of cholesterol-rich diet induces LDL-hypercholesterolaemia and advanced coronary atherogenesis. Withdrawal of dietary cholesterol reverses hypercholesterolaemia and stabilizes coronary plaque in miniature pigs [27] and partially restores endothelial function in rats [13].

In the present study, diet-fed obese rats had significant vascular abnormalities, in particular a marked endothelial dysfunction. The potential factors for vascular abnormalities are obesity per se [1], insulin resistance [3], increased leptin levels [28], and higher level of plasma triglyceride [8,12,13], as well as total body weight and an increase in fat-pad mass. In fact, there were negative correlations between endothelial function and plasma levels of insulin, leptin and triglycerides. Previous studies reporting occurrence

of changes in vascular function in the absence of obesity or rise in plasma leptin levels and insulin resistance [12–14] argue against major roles of obesity per se, insulin and leptin resistance in vascular dysfunction. Moreover, in this study diet-to-chow rats were significantly obese compared with chow-fed controls, yet their vascular function was similar to that of chow-fed group, further arguing, at least in the rat, against the role of obesity per se in the genesis of vascular abnormalities. On the other hand, recent human and animal studies point to the role of circulating triglyceride in the genesis of vascular dysfunction [8,11–13,29,30]. In this study, circulating triglyceride levels were raised more than 115% and a significant negative correlation was seen between fasting plasma triglyceride levels and endothelium-dependent vasorelaxation. The plasma TG levels of chow-fed and diet-to-chow groups were similar as were CCh-induced vasorelaxation between the two groups, suggesting that changing diet (i.e. removing palatable diet and replacing it with the standard chow) resulted in lowering plasma triglyceride levels and thereby improving vascular function.

Overall, this study indicates that prolonged consumption of a high-palatable, high-energy diet induces metabolic and cardiovascular disorders. Nonetheless, these abnormalities are completely reversible by chronic withdrawal of the palatable diet.

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The effects of fenofibrate on metabolic and vascular changes induced by chocolate-supplemented diet in the rat

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Abstract

Current evidences suggest that diet per se plays an important role in genesis of metabolic and vascular function abnormalities. We have investigated the effects of addition of a high-fat diet (chocolate) in the presence or absence of short-term (7 days) administration of fenofibrate on metabolic and vascular changes in adult male Wistar rats. Despite similarities in total body weight in all groups, compared with control fed groups, chocolate-supplemented animals had significantly higher plasma triacylglyceride and non-esterified fatty acids and leptin (for all, $P < 0.01$), but not glucose or insulin levels. Fenofibrate treatment corrected metabolic changes. In the mesenteric arteries, responses to KCl and noradrenaline were significantly (for both, $P < 0.01$) higher in chocolate-supplemented group. Furthermore, vasorelaxant responses to carbamylcholine, but not to sodium nitroprusside, were significantly ($P < 0.01$) reduced in the chocolate-supplemented group. Although fenofibrate failed to improve noradrenaline and KCl responses, it was effective in improving carbamylcholine-induced vasorelaxation. These data suggest that high-fat diet has a profound effect on plasma lipid profile and vascular function. Furthermore, fenofibrate treatment may ameliorate high-fat diet effects on vascular function and metabolic changes.

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Keywords: High-fat diet; Vascular contractility; Endothelial dysfunction; Fenofibrate; Hypertriglyceridaemia

1. Introduction

Acceleration in the rate of diet-induced obesity throughout the world is becoming a major health problem. Excess energy intake in the form of refined carbohydrates and fatty acids is one of the main causes of human obesity. Obesity is a major risk factor for cardiovascular disorders, which include hypertension, atherosclerosis and ultimately myocardial infarction (Andres, 1980; Lakka et al., 2002). Arterial function and structural abnormalities which include alterations of the release of vasoactive substances from endothelial cells as well as changes in arterial wall elasticity have been seen in obese human subjects and certain animal models that reflect some aspects of human obesity (Cowan et al., 1991; Dobrian et al., 2000).

Moreover, human and animal studies have shown various degrees of endothelial dysfunction in obesity, namely a reduction to vasorelaxant's response (Abram, 1997; Naderali et al., 2001; Perticone et al., 2001; Steinberg et al., 1996).

Increasing reports suggest that diet and its variant component play important roles in arterial function. Studies have shown that consumption of a specific diet of high fat or low protein causes endothelial abnormalities in humans and animals, while a low fat diet has a protective effect on vascular reactivity (Vogel et al., 1997; Verhamme et al., 2002; Naderali et al., 2004). Moreover, pharmacological lowering of plasma lipids by fenofibrate, retards atheroma developments (Renier et al., 2000), indicating a need to control circulating plasma lipids. Therefore, the aim of this study was to investigate the effects of supplementation of a high-fat diet in the presence or absence of short-term (7 days) administration of fenofibrate on metabolic changes and vascular function.

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2. Materials and methods

2.1. Animals and experimental protocol

Adult male Wistar rats ($n=21$) were randomized and assigned to two groups which were fed on either a standard pelleted laboratory chow (CRM Biosure, Cambridge, UK; which provided 60% of energy as carbohydrate, 30% as protein and 10% as fat) throughout ($n=7$; control), or given chow plus Galaxy smooth and creamy milk chocolate (Mars) (which provided 42.2% of energy as carbohydrate, 5.1% as protein and 52.7% as fat) ($n=14$; chocolate-supplemented). After 14 weeks, half of the chocolate-supplemented group was given fenofibrate (50 mg/kg/day) and the remainder were given vehicle (1% carboxymethyl cellulose at 1 ml/kg body weight; Sigma, UK) by oral gavage daily for 7 days, before being sacrificed. All animals had free access to their respective diets and water, and were housed in groups of 2 under controlled environmental conditions (19–22 °C; 30–40% humidity) and a 12-h light/dark cycle (lights on at 07:00 h). Daily energy intake was determined in all three groups for the last 7 days of the study.

The rats were killed by CO₂ inhalation, and the epididymal and perirenal fat pads and the gastrocnemius and cardiac muscles were dissected and weighed. Blood was collected for haematocrite studies and for the measurement of glucose, insulin, leptin, non-esterified fatty acids and triacylglycerol. Plasma glucose concentration was determined by the glucose oxidase method, while non-esterified fatty acids and triacylglycerol concentrations were determined by commercial diagnostic kits (Boehringer Mannheim, U.K and Sigma, U.K, respectively). Insulin and leptin concentrations were measured by radioimmunoassay (RIA) kits (Pharmacia/Upjohn Diagnostics, U.K. and Linco Research, U.K, respectively).

2.2. Assessment of vascular function

Six third-order mesenteric arteries from each animal (150–180 μ m diameter, 2 mm lengths) were carefully dissected from each animal and mounted on two 40- μ m diameter stainless-steel wires in an automated myograph (Cambustion, Cambridge, UK). Pairs of arteries were incubated in a 5-ml organ bath containing physiological salt solution (PSS; composition [in mM]: NaCl 119, KCl 4.7, CaCl₂ 2.5, MgSO₄ 1.17, NaHCO₃ 25, KH₂PO₄ 1.18, EDTA 0.026 and glucose 5.5) gassed with 95% O₂ and 5% CO₂ at 37 °C.

After 30 min equilibration, the length–tension characteristics for each vessel were determined (Naderali et al., 2000). Arteries were allowed a further 30 min to equilibrate before being depolarised twice with high-potassium physiological salt solution (KPSS) in which NaCl in normal PSS was replaced by an equimolar concentration of KCl (125 mM). Cumulative concentration–response curves to either KCl (10–125 mM) or noradrenaline (0.5–6 μ M) were then

carried out. Any vessel failing to reach its predetermined target tension in response to KCl (125 mM) was discarded. We have also measured the effects of selected endothelium-dependent vasodilator, carbamylcholine, and endothelium-independent vasodilator, sodium nitroprusside on arteries precontracted with 8 μ M noradrenaline.

2.3. Reagents

Noradrenaline, carbamylcholine, sodium nitroprusside, and fenofibrate were all obtained from Sigma Chemicals. All solutions were prepared freshly in distilled water prior to use.

2.4. Data interpretation and statistical analyses

KCl- and noradrenaline-induced vasoconstriction responses were expressed as absolute force generated. Relaxation in response to carbamylcholine and sodium nitroprusside were calculated as the percentage reduction from the maximal tension generated in response to a supramaximal concentration of noradrenaline (8 μ M). An average response for all the vessels from a given animal was determined before group analysis. Data are presented as mean \pm S.E.M. Statistical significance was tested using repeated-measures analysis of variance (ANOVA) or the Mann–Whitney test, as appropriate. Differences were considered statistically significant at values of $P < 0.05$.

3. Results

3.1. Metabolic data

Supplementation of chocolate did not alter total body weight gain. In fact throughout the experiment there were no significant differences in total body weight at any given time

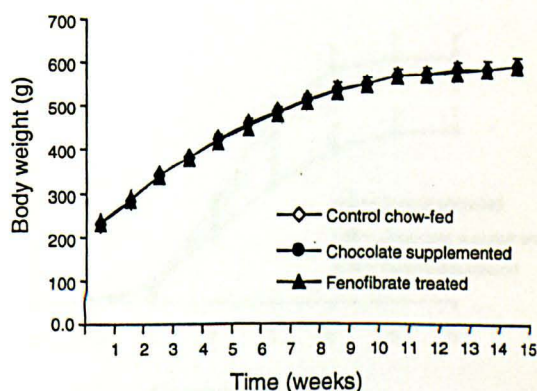


Fig. 1. The effect of chocolate supplementation and fenofibrate treatment on total body weight in the rat. Control animals were fed standard chow while chocolate-supplemented groups was given chow plus chocolate (Galaxy) for 15 weeks. Fenofibrate-treated rats were given fenofibrate (50 mg/kg/day) daily for 7 days. At the end of 15 weeks, there were no significant changes in total body weight among the three groups. In fact weight increase in all three groups were superimposable. Data are expressed as mean \pm S.E.M.

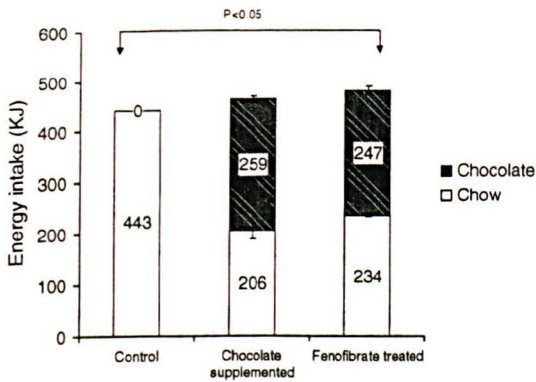


Fig. 2. Cumulative weekly energy intake of animals from all three groups during fenofibrate (50 mg/kg/day) treatment period. The increase in total energy intake of animals on chocolate supplementation was not statistically significant ($P=0.346$), however fenofibrate treatment further increased energy intake, which was significantly ($P<0.05$) higher than chow-fed controls.

between chow-fed control and chocolate-supplemented test groups (Fig. 1). Furthermore, the epididymal and perirenal fat pad masses as well as gastrocnemius muscle mass were comparable in all three groups, indicating the absence of any obesity. Analysis of food intake during the last 7 days showed that compared with control chow-fed rats, increase in total weekly energy intake of chocolate-supplemented group was insignificant (Fig. 2). However, fenofibrate-treated rats had significantly ($P<0.05$) higher energy intake than control chow-fed groups (Fig. 2). Nonetheless, this

Table 1
Physiological and metabolic characteristics of the 3 experimental groups

| | Chow-fed controls ($n=7$) | Chocolate-supplemented ($n=7$) | Fenofibrate-treated ($n=7$) |
|-------------------------------|-----------------------------|----------------------------------|-------------------------------|
| Body weight (g) | 598±23 | 606±19 | 611±21 |
| Epididymal fat pad mass (g) | 6.56±0.80 | 7.20±0.59 | 6.56±0.64 |
| Perirenal fat pad mass (g) | 6.20±0.94 | 7.24±0.66 | 6.69±0.84 |
| Gastrocnemius muscle mass (g) | 3.27±0.15 | 3.06±0.11 | 3.16±0.11 |
| Heart muscle mass (g) | 1.57±0.09 | 1.54±0.05 | 1.70±0.08 ^{a,b} |
| Haematocrit | 49.1±0.6 | 47.5±0.8 | 46.4±0.5 ^a |
| Plasma glucose (mM) | 11.3±0.9 | 13.5±1.0 | 9.3±0.4 ^{a,b} |
| Plasma insulin (μU/ml) | 1.52±.38 | 1.31±0.33 | 1.12±0.21 ^a |
| Plasma leptin (ng/ml) | 15.1±2.2 | 21.6±4.0 ^a | 14.2±6.9 ^b |
| Plasma triglycerides (mM) | 1.50±0.06 | 1.96±0.12 ^a | 0.96±0.08 ^{a,b} |
| Plasma NEFA (mM) | 0.28±0.02 | 0.39±0.02 ^a | 0.23±0.01 ^b |

Data are mean±SEM.

^a $P<0.01$ vs controls.

^b $P<0.01$ vs chocolate-supplemented.

increase in energy intake did not translate to a significant increase in total body weight (Table 1).

While supplementation of chocolate had no significant effect on cardiac muscle mass or haematocrit values, fenofibrate-treated animals have significantly ($P<0.01$) higher cardiac muscle mass and reduced haematocrit values (Table 1). At the end of the experiment, fasting plasma glucose, insulin levels were comparable between control chow-fed and chocolate-supplemented groups, however, fenofibrate-treated animals had significantly ($P<0.01$) lower glucose and insulin levels than control counterparts. Furthermore, chocolate-supplemented animals had significantly higher fasting concentrations of plasma leptin, triacylglycerol and non-esterified fatty acids than chow-fed controls. These changes were almost completely reversed by fenofibrate treatment (Table 1).

3.2. Agonist-induced contractile responses

KCl (10–125 mM)- and noradrenaline (0.5–6 μM)-induced vasoconstriction in arteries from all three groups

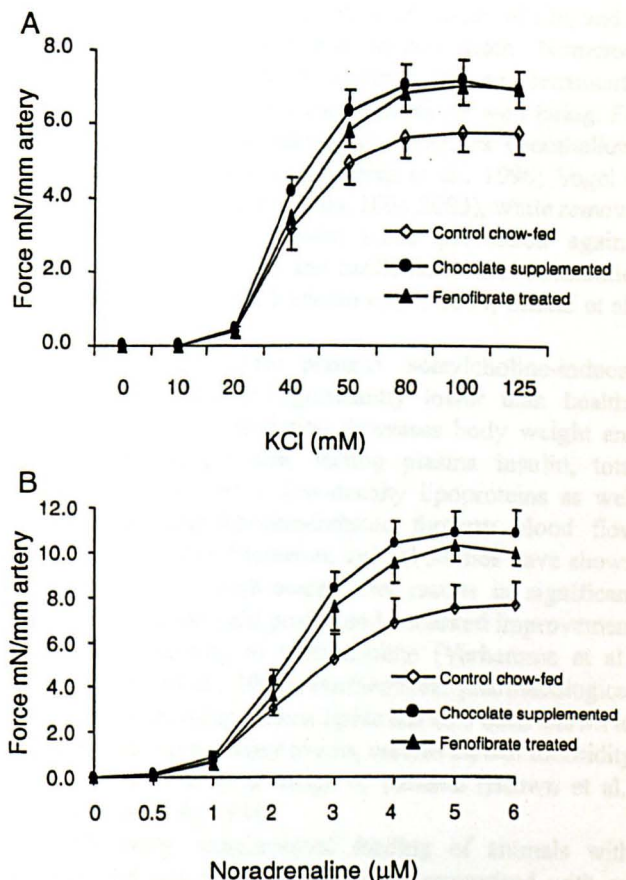


Fig. 3. Vasoconstriction responses to cumulative concentration of (A) KCl (10–125 mM) and (B) noradrenaline (0.5–6 μM) in arteries from chow-fed control, chocolate-supplemented and fenofibrate-treated animals. Both KCl and noradrenaline-induced contractions were significantly (ANOVA, $P<0.01$) augmented in arteries from chocolate-supplemented animals. Data represent mean±S.E.M.

produced a characteristic sigmoid relationship. The contractile responses to KCl and noradrenaline in arteries from chocolate-supplemented animals (with or without fenofibrate treatment) were significantly ($P < 0.001$) increased in comparison to control groups (Fig. 2). The maximal KCl-induced contractions were 5.78 ± 0.54 control, 7.14 ± 0.57 chocolate-supplemented, and 7.07 ± 0.52 mN/mm artery fenofibrate-treated, respectively (Fig. 3A). A similar picture was also seen with noradrenaline-induced vasoconstriction giving maximal contraction of 7.68 ± 1.02 control, 10.91 ± 0.96 supplemented, and 10.34 ± 0.66 mN/mm artery fenofibrate-treated, respectively (Fig. 3B).

3.3. Endothelium-dependent and -independent vasorelaxation

Noradrenaline precontracted arteries from chow-fed rats demonstrated progressive relaxation to cumulative addition of carbamylcholine (10 nM–100 μ M), achieving a maximum of $87 \pm 2\%$ at a carbamylcholine concentration of 100 μ M. Arteries from test and fenofibrate-treated groups that were similarly exposed to carbamylcholine displayed similar responses with maximal relaxation of $85 \pm 6\%$ and

$92 \pm 2\%$ (Fig. 4A). However, arteries from chocolate-supplemented rats had significantly ($P < 0.01$) higher EC_{50} than control chow-fed animals (EC_{50} : 1.86 ± 0.02 chocolate-supplemented, vs EC_{50} : 0.28 ± 0.01 μ M chow-fed). Fenofibrate treatment significantly ($P < 0.05$) improved EC_{50} compared with chocolate-supplemented group (EC_{50} : 0.68 ± 0.01 μ M fenofibrate treatment vs EC_{50} : 1.86 ± 0.02 μ M chocolate-supplemented).

Endothelium-independent vasodilator (sodium nitroprusside; 10 nM–100 μ M) also induced concentration dependent vasorelaxation of arteries from all three groups with no significant differences in arterial vasorelaxation between the three groups achieving a maximal relaxation of $85 \pm 3\%$ control, $82 \pm 2\%$ chocolate-supplemented, and $84 \pm 3\%$ fenofibrate-treated groups, respectively (Fig. 4B).

4. Discussion

Excessive availability of high-energy diet in many industrialised societies, is producing a health hazard, namely obesity and obesity related secondary disorders, which ultimately leads to the deterioration of quality of life, and in many cases to premature and sudden death. Numerous experimental studies have reported severe detrimental outcome of excessive high-energy diets on well being. For example, high-fat, high-energy diet impairs endothelium-dependent vasorelaxation (Steinberg et al., 1996; Vogel et al., 1997; Naderali and Williams, 2001, 2003), while removal of high-energy diet provides some protection against obesity-induced metabolic and cardiovascular abnormalities (Verhamme et al., 2002; Naderali et al., 2004; Sasaki et al., 2002).

In hypertensive obese patients, acetylcholine-induced forearm blood flow is significantly lower than healthy individuals. Caloric restriction decreases body weight and the mean blood pressure, fasting plasma insulin, total cholesterol, triglycerides, low-density lipoproteins as well as enhancing acetylcholine-induced forearm blood flow (Sasaki et al., 2002). Moreover, animal studies have shown that withdrawal of high-energy diet results in significant reduction of plasma lipid profile and a marked improvement of vascular reactivity to acetylcholine (Verhamme et al., 2002; Naderali et al., 2004). Furthermore, pharmacological lowering of circulating plasma lipids has also been shown to reduce the risk for coronary events, cardiovascular morbidity and mortality in a broad range of patients (Brown et al., 1993; Bucher et al., 1999).

In this study, supplemental feeding of animals with chocolate did not result in obesity (characterised with an excessive increase in total body weight and fat pad mass). This may be due to comparable caloric intake between chow-fed control and chocolate-supplemented groups. However, there were significant metabolic abnormalities, namely higher plasma lipids of triacylglycerol and non-esterified fatty acids, in chocolate-supplemented animals. In agreement

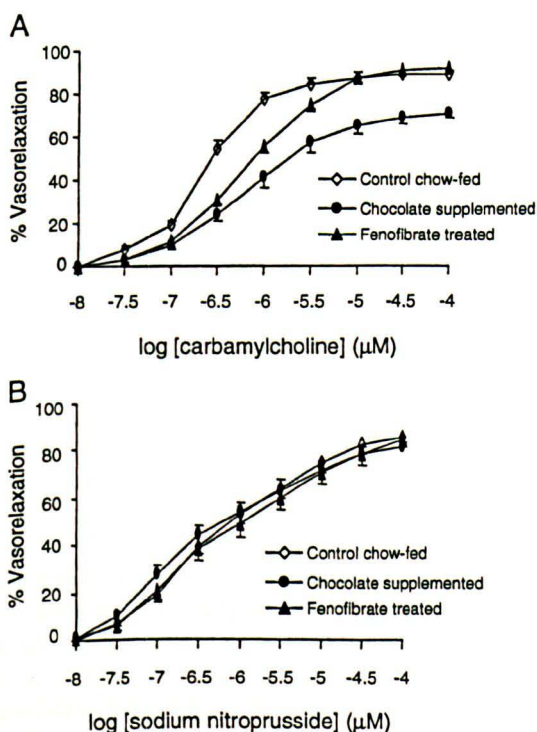


Fig. 4. The effects of (A) carbamylcholine, and (B) sodium nitroprusside on NA-precontracted (8 μ M) arteries from three groups of rats. When contraction reached a plateau after 2 min, concentration–response curves to carbamylcholine or sodium nitroprusside were carried out. Data represent mean \pm S.E.M. The concentration–response curves to carbamylcholine but not to sodium nitroprusside, were significantly reduced in chocolate-supplemented group (ANOVA, $P < 0.01$), while fenofibrate treatment significantly ($P < 0.01$) improved carbamylcholine-induced vasorelaxation in chocolate-supplemented animals.

with previous reports (Despres et al., 2002; Capell et al., 2003), fenofibrate treatment in the presence of chocolate diet completely reversed lipid profile. Higher plasma lipids seen in this study were similar to those achieved by chronic consumption of high-fat, high-energy diets and obesity (Naderali et al., 2001, 2004; Perticone et al., 2001; Steinberg et al., 1996; Verhamme et al., 2002) indicating that adipose tissue accumulation is not a pre-requisite for abnormal lipid profile (Vogel et al., 1997; Naderali and Williams, 2003).

Examination of contractile and relaxation properties of arteries showed striking differences in responses to various stimuli between the three groups. In contrast to previous reports (Walker et al., 1997; Fontes et al., 1998; Naderali et al., 2001, 2004), contractile responses to KCl and noradrenaline were significantly augmented in chocolate-supplemented diet. Although from our study it is not possible to contemplate the mechanism of hyperreactivity to KCl and noradrenaline, nonetheless, one possibility is that long-term consumption of chocolate may have a direct effect on the activity of the contractile apparatus, or it may have increased/eased the availability of the calcium for contraction required by KCl and noradrenaline. The latter theory is more plausible as chocolate has been shown to markedly increase calciuria (Nguyen et al., 1994). However, these hypotheses merit further investigation.

Numerous studies have shown that excessive high-energy diet impairs endothelium-dependent arterial function (Sasaki et al., 2002; Brown et al., 1993; Bucher et al., 1999; Naderali et al., 2001, 2004; Walker et al., 1997; Fontes et al., 1998). Moreover, defects in vasorelaxation can also be seen with short-term consumption of high-energy diet in the absence of any obesity (Vogel et al., 1997; Naderali and Williams, 2001, 2003). In agreement with previous reports, in this study there was a marked attenuation of endothelial mediated carbamylcholine-induced vasorelaxation in the absence of any obesity, suggesting a significant defect in either carbamylcholine-induced NO production via eNOS activity (Boulanger, 1999) or the release of endothelium-derived hyperpolarizing factor (Gerber et al., 1998). On the other hand, vasorelaxant effect of sodium nitroprusside is due to its direct stimulatory action on cGMP and thereby induction of NO generation (Azula et al., 1996) in vascular smooth muscle cells. Failure of the chocolate-supplemented diet in altering sodium nitroprusside-induced vasorelaxation, argues against any detrimental changes in arterial smooth muscle activity.

A number of various hypotheses have been put forward as the mechanism(s) of action of high-fat diet and obesity related vascular dysfunction, which include a rise in plasma non-esterified fatty acids (Steinberg et al., 1997) hypertriglyceridaemia (Tka et al., 1997; Lewis et al., 1999; Naderali et al., 2001). The detrimental effects of increased circulating plasma lipids has been, at least partly, attributed to inhibition of eNOS activity and thereby reduction in NO synthesis (Liao et al., 1995), ultimately resulting in a reduced endothelial-dependent vasorelaxation. Raised non-esterified

fatty acids and triacylglycerols levels in chocolate-supplemented animals seen in this study, argue for the role of plasma non-esterified fatty acids and triacylglycerols as the possible agents effecting vasorelaxation. These hypotheses are further strengthened by the finding that fenofibrate treatment in our study improved carbamylcholine-induced responses. Although the mechanism of action of fenofibrate on vascular function is not yet fully known, the protective effects of fenofibrate on vascular function may be due to correction of lipid profile (Despres et al., 2002; Capell et al., 2003) or increasing formation, availability and duration of NO action (Haak et al., 1998; Playford et al., 2002), indicating improvement of endothelial function in fenofibrate-treated animals. Overall, this study confirms that consumption of a high-fat diet attenuates arterial relaxation in the absence of any obesity in the rat. The arterial dysfunction may be due to an induction of hyperlipidaemia in general or hypertriglyceridaemia in particular. Moreover, lowering circulating plasma non-esterified fatty acids and/or triacylglycerols by fenofibrate treatment may have beneficial effects on endothelial function.

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Fenofibrate lowers adiposity and corrects metabolic abnormalities, but only partially restores endothelial function in dietary obese rats

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Abstract

In humans, dietary-induced obesity markedly increases plasma lipid profile and impairs vascular function leading to increased incidence of cardiovascular events. We have recently reported that chronic withdrawal of obesity-inducing diet attenuates obesity and completely corrects endothelial function. The aim of this study was to investigate whether fenofibrate-induced decrease in adiposity would also correct vascular function in the presence of obesity-inducing diet. Wistar rats were fed with either standard laboratory chow (lean, $n = 9$) or given a highly palatable diet (diet-fed, $n = 18$) for 15 weeks. After 7 weeks, half of the diet-fed group was treated with fenofibrate (fenofibrate-treated, $n = 9$) for 8 weeks before being sacrificed. Untreated diet-fed ($n = 9$) rats had significantly higher body weight, total fat mass (by up to two-fold, $p < 0.001$ for both), and raised fasting plasma levels of insulin, leptin and triglycerides (up to 110%; $p < 0.001$), but not glucose or nonesterified fatty acids (NEFA) than both lean control and fenofibrate-treated groups. Resistance mesenteric arteries responses to KCl- and noradrenaline-induced vasoconstriction were similar in all three groups. However, compared with lean controls, endothelium-dependent vasorelaxation responses were shifted to the right in both untreated and fenofibrate-treated diet-fed groups. Fenofibrate treatment improved endothelium-dependent vasorelaxation at only high carbamylcholine concentrations (10 μM). There were no differences in endothelium-independent vasorelaxation between the three groups. These results indicate that, in the presence of obesity-inducing diet, fenofibrate markedly reverses obesity and corrects insulin resistance and lipid profile, but it only has a limited beneficial effect on vascular function. Therefore, it seems that diet component rather than obesity per se plays a key role in the genesis of vascular abnormalities.

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Keywords: Diet; Obesity; Insulin resistance; Fenofibrate; Endothelial function

1. Introduction

Obesity and type 2 diabetes are strong risk factors for the development of atherosclerosis. The underlying pathophysiological mechanisms are not fully understood, but may include hyperlipidaemia and increased oxidation of low-density lipoproteins [1], hypertriglyceridaemia [2], insulin resistance [3] and hyperleptinaemia [4]. Obesity, type 2 diabetes and the above risk factors are also associated with marked defects in vascular function. Endothelial dysfunction in particular is prominent and may contribute to atherogenesis [1,2,5,6]. The lipid-lowering fibrates, such as fenofibrate and ciprofibrate, have been shown to impede the development

of atheroma. These drugs bind to peroxisome proliferator-activated receptor α (PPAR α), leading to enhanced lipid catabolism in the liver [7] and reductions in total body fat as well as the circulating plasma lipids.

Fenofibrate has been reported to retard angiographic progression of coronary atherosclerosis in diabetic patients [8], and improve the microcirculation in patients with hyperlipidaemia [9]. The mechanism(s) of the beneficial effects of fenofibrate on vascular function has not been fully elucidated yet, but direct activation of PPAR α in arterial wall [10], correction of lipid abnormalities [11,12], and increasing the formation, availability and action of NO [13,14] have all been postulated.

In this study we aimed to evaluate further the effect of fenofibrate on vascular abnormalities in the presence of obesity-inducing diet in obese rats.

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2. Material and methods

2.1. Animals

Adult male Wistar rats ($n = 27$) were randomised and assigned to a control group ($n = 9$, 191 ± 3 g) and a test group ($n = 18$, 190 ± 1 g) that was fed a highly palatable diet for 15 weeks.

All animals had free access to water and were housed individually under controlled environmental conditions ($19\text{--}22^\circ\text{C}$; $30\text{--}40\%$ humidity) and a 12-h light/dark cycle (lights on at 07:00 h). Controls were fed a standard laboratory pelleted diet (CRM Biosure, Cambridge, UK), which provides 60% of energy as carbohydrate, 30% as protein and 10% as fat. The test group had free access to a highly palatable diet consisting of 33% (by weight) ground pellet diet, 33% Nestlé condensed milk, 7% sucrose and 27% water. This provided 65% of energy as carbohydrate, 19% as protein and 16% as fat. After 7 weeks, half of the test group were given fenofibrate (50 mg/kg/day) and the remainder were given vehicle (1% carboxymethyl cellulose at 1 ml/kg body weight; Sigma, Pool, UK) by oral gavage, daily for 8 weeks.

On the day of experiment, the rats were killed by CO_2 inhalation after 2 h of fasting. The total body fat mass was measured immediately by bioimpedance method using the TOBEC[®] equipment (Biotech. Instruments Ltd., Kimpton, UK). Blood was removed by cardiac puncture into cold heparinized tubes, and the gonadal and perirenal fat pads as well as the gastrocnemius and soleus muscles were dissected and weighed. The plasma was immediately separated by centrifugation before being frozen for later measurements of glucose, insulin, leptin, nonesterified free fatty acids (NEFA) and triglycerides (TG). Plasma glucose concentration was determined using a glucose oxidase method, and NEFA and TG concentrations using commercial diagnostic kits (Boehringer Mannheim, Milton Keynes, Bucks and Sigma Diagnostics, Poole, UK). Insulin and leptin concentrations were measured by radioimmunoassay (RIA) kits (Pharmacia/Upjohn Diagnostics, Lewes, Sussex and Linco Research, Biogenesis, Poole, Dorset, UK, respectively).

2.2. Homeostasis model assessment (HOMA)

HOMA, an index of insulin resistance, which employs measures of fasting plasma concentrations of glucose and insulin, was calculated according to the method described previously [15].

2.3. Assessment of vascular function

Six third-order mesenteric arteries (>200 μm diameter, 2-mm length) were carefully dissected from each animal. Each artery was freed of fat and connective tissue and mounted on two 40- μm diameter stainless-steel wires in an automated myograph (Cambustion, Cambridge, UK), based on the principle of the Mulvany myograph which measures isometric

tension generated in response to various stimuli. Arterial segments (two at a time) were incubated in a 5-ml organ bath containing physiological salt solution (PSS; composition [in mM]: NaCl, 119; KCl, 4.7; CaCl_2 , 2.5; MgSO_4 , 1.17; NaHCO_3 , 25; KH_2PO_4 , 1.18; EDTA, 0.026; and glucose 5.5) gassed with 95% O_2 and 5% CO_2 at 37°C .

After 30 min equilibration, the length–tension characteristics for each artery were determined as described previously [16]. The computer also calculated the target tension that each vessel should develop in response to a maximal stimulus. Arteries were then allowed a further 30 min to equilibrate before being depolarised twice with high-potassium physiological salt solution (KPSS, 125 mM) in which the NaCl in normal PSS was replaced by an equimolar concentration of KCl. Cumulative concentration–response curves to either KCl (10–125 mM) or noradrenaline (NA, 0.5–6 μM) were then carried out. Any vessel failing to reach its predetermined target tension in response to vasoconstriction with KCl (125 mM) was discarded.

2.4. Assessment of defect in vascular relaxation

Endothelium-dependent and -independent defects were investigated by exposing NA-precontracted arteries to carbamylcholine (CCh) or sodium nitroprusside (SNP), respectively. To eliminate possible interference by vasoactive prostanoids, 10 mM indomethacin was added to the organ bath, prior to contraction with NA (8 μM). When contraction reached a plateau (after 2 min), concentration–response curves were performed to either CCh or SNP (for both, 10 nM–100 μM).

2.5. Reagents

Noradrenaline (NA), carbamylcholine (CCh), sodium nitroprusside (SNP), indomethacin, carboxymethyl cellulose and fenofibrate were all obtained from Sigma (Poole, Dorset, UK). NA, CCh and SNP were dissolved in double distilled water while indomethacin was dissolved in ethanol. NA and SNP were kept away from light throughout the experiment. All water-soluble solutions were freshly made on the day of experiment.

2.6. Data interpretation and statistical analyses

Vasoconstriction in response to NA and KCl were expressed as absolute force generated (mN/mm artery). Relaxation in response to CCh and SNP was calculated as the percentage reduction from the maximal tension generated at the supramaximal concentration of NA (8 μM). The average response for all the vessels from a given animal was used in the group analysis. EC_{50} values were determined to evaluate changes in agonist-induced responses.

Data showed a normal distribution (Shapiro–Wilkes *W* test) and are therefore expressed as mean \pm S.E.M. Statistical significance was tested using Student 't-test' or repeated-

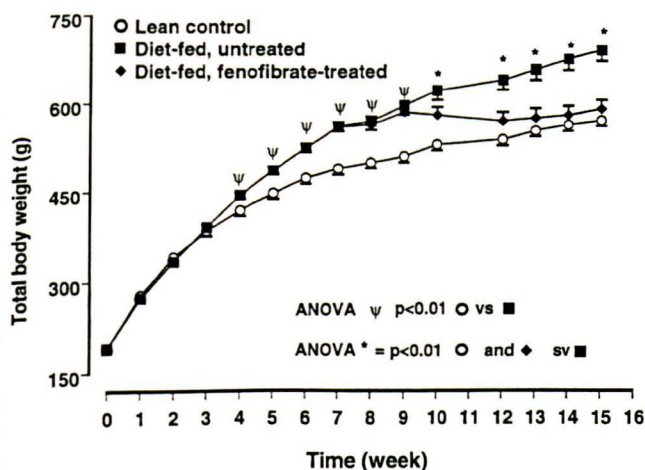


Fig. 1. The effects of palatable diet and fenofibrate on total body weight in male Wistar rats. Lean control animals were fed a standard laboratory chow while diet-fed groups was given a palatable diet for 15 weeks. Fenofibrate-treated rats were given fenofibrate (50 mg/kg/day) daily for 8 weeks. At the end of 15 weeks, untreated diet-fed animals were significantly ($p < 0.0001$) heavier than lean control and fenofibrate-treated groups. Data are expressed as mean \pm S.E.M.

measures (ANOVA; Bonferroni *t*-test) or the Mann–Whitney test, as appropriate. Differences were considered statistically significant at the $p < 0.05$ levels.

3. Results

3.1. Effects of fenofibrate on body weight and adiposity

As shown in Fig. 1, rats fed palatable diet progressively gained more weight than chow-fed controls. This difference became significant after the third week of feeding and was maintained thereafter. The final body weight of the diet-fed animals was significantly higher than that of chow-fed controls ($>21\%$; 695 ± 17 versus 572 ± 8 g, $p < 0.0001$). Overall, the weekly weight gain of diet-fed animals was significantly

greater than that of chow-fed controls (diet-fed: 20.1 ± 4.77 versus control: 11.98 ± 4.65 g, $p = 0.0135$).

Administration of fenofibrate significantly reduced weekly weight gain of diet-fed animals and final total body weight (Fig. 1). After 8 weeks of treatment, total body weight of the diet-fed fenofibrate-treated animals was not significantly different from that of chow-fed controls (fenofibrate treated: 593 ± 16 versus chow-fed: 572 ± 8 g, $p = 0.2738$), but it was significantly lower ($p = 0.00071$) than that of untreated diet-fed animals (Fig. 1).

After 15 weeks of study, gastrocnemius and soleus muscle masses were comparable between all three groups. However, diet-fed animals had significantly higher total white adipose tissue (sum of epididymal and perirenal fat pads) mass than both chow-fed (by >2 -fold, $p < 0.0002$) and fenofibrate-treated (by >1.5 -fold, $p = 0.0114$) groups (Table 1). However, fenofibrate-treated animals had significantly (by 1.4-fold; $p = 0.0264$) heavier white adipose tissue than chow-fed controls. Bioimpedance measurements indicated higher percentage of body fat in diet-fed animals compared with chow-fed and fenofibrate-treated groups. Conversely, diet-fed rats had significantly lower percentage of lean mass than chow-fed and fenofibrate-treated animals (Table 1).

3.2. Effects of diet and fenofibrate on blood metabolites

Palatable diet feeding did not affect fasting plasma glucose levels (11.7 ± 1.2 chow-fed; 14.3 ± 0.3 mM diet-fed; $p = 0.0625$). However, diet-fed rats had significantly higher fasting terminal insulin (3.8 ± 0.8 diet-fed versus 1.6 ± 0.3 chow-fed; $p = 0.0025$) than the chow-fed group, with a significantly higher HOMA index value (2.40 ± 0.52 diet-fed versus 0.73 ± 0.10 chow-fed; $p = 0.0086$) indicating development of insulin resistance in diet-fed animals. Fenofibrate treatment significantly lowered levels of both glucose (10.8 ± 0.3 mM; $p < 0.0001$ versus untreated diet-fed group) and insulin (1.4 ± 0.2 mM; $p = 0.0014$ versus untreated diet-fed group) levels. Consequently, HOMA index was also reduced

Table 1
The effects of fenofibrate on physiological and metabolic profiles of dietary-induced obese rats

| | Lean controls (n = 9) | Diet-fed, untreated (n = 9) | Diet-fed, fenofibrate-treated (n = 9) |
|-------------------------------|-----------------------|-----------------------------|---------------------------------------|
| Epididymal fat-pad mass (g) | 5.4 ± 0.6 | 11.2 ± 1.0^a | 7.2 ± 0.7^b |
| Perirenal fat pad mass (g) | 6.1 ± 0.5 | 10.9 ± 1.4^a | 6.8 ± 0.8^b |
| % Lean mass | 83.6 ± 2.0 | 76.3 ± 1.6^c | 80.7 ± 1.8 |
| % Fat mass | 16.4 ± 2.0 | 23.7 ± 1.6^c | 19.3 ± 1.8 |
| Gastrocnemius muscle mass (g) | 3.2 ± 0.1 | 3.4 ± 0.1 | 3.1 ± 0.1 |
| Soleus muscle mass (g) | 0.21 ± 0.01 | 0.22 ± 0.01 | 0.21 ± 0.01 |
| HOMA index | 0.73 ± 0.10 | 2.40 ± 0.52^a | 0.67 ± 0.10^b |
| Plasma glucose (mM) | 11.7 ± 1.2 | 14.3 ± 0.3 | 10.8 ± 0.3^b |
| Plasma insulin (μ g/l) | 1.6 ± 0.3 | 3.8 ± 0.8^a | 1.4 ± 0.2^b |
| Plasma leptin (ng/ml) | 6.9 ± 0.5 | 9.4 ± 0.7^a | 8.5 ± 0.4^a |
| Plasma triglycerides (mM) | 0.79 ± 0.04 | 1.68 ± 0.17^a | 0.80 ± 0.06^b |
| Plasma NEFA (mM) | 0.44 ± 0.02 | 0.45 ± 0.04 | $0.31 \pm 0.02^{a,b}$ |

Data are mean \pm S.E.M.

^a $p < 0.001$ vs. lean control.

^b $p < 0.001$ vs. untreated diet-fed.

^c $p < 0.01$ vs. lean control and fenofibrate treated diet-fed.

(0.67 ± 0.10 ; $p = 0.0073$ versus untreated diet-fed group), indicating correction of insulin resistance (Table 1).

Diet-fed rats had significantly higher plasma leptin levels than chow-fed control (9.4 ± 0.7 diet-fed versus 6.9 ± 0.5 ng/ml chow-fed; $p = 0.0131$). The rise in leptin levels was not affected by fenofibrate treatment (8.5 ± 0.4 ng/ml; $p = 0.03612$ versus chow-fed). Fasting plasma levels of NEFA were comparable in diet-fed and chow-fed groups (0.44 ± 0.02 chow-fed; 0.45 ± 0.04 mM diet fed). However, diet-fed rats had significantly higher (>110%) levels of fasting TG than chow-fed (chow-fed: 0.79 ± 0.04 ; diet-fed: 1.68 ± 0.17 mM; $p = 0.0004$). Fenofibrate treatment significantly lowered fasting plasma levels of both NEFA (0.31 ± 0.02 mM; $p = 0.0012$ versus chow-fed and untreated diet-fed groups) and TG (0.80 ± 0.06 mM; $p = 0.0001$ versus untreated diet-fed group) below those of chow-fed group (Table 1).

3.3. Vascular data

Diet-induced obesity did not alter vessel diameter, and there were no significant differences in arterial diameter between the three groups (200 ± 15 in chow-fed; 205 ± 15 in diet-fed; 200 ± 10 μ m in fenofibrate-treated).

3.4. Agonist-induced vasoconstriction responses

Cumulative concentration–response curves of KCl (10–125 mM) and NA (0.5–6 μ M) in arteries from all three groups showed the characteristic sigmoid form. KCl-induced contraction was similar in all three groups, and the concentration–response curves were virtually superimposable, reaching maximum force of 5.68 ± 0.44 (chow-fed), 6.07 ± 0.33 (diet-fed) and 5.35 ± 0.37 (fenofibrate-treated) mN/mm artery (Fig. 2a). There was no significant differences between the groups, either overall (by ANOVA) or at any given KCl concentration. A similar pattern was also seen with NA-induced contractions, giving maximum force of 9.14 ± 0.63 (chow-fed), 9.94 ± 0.43 (diet-fed) and 9.70 ± 0.84 (fenofibrate-treated) mN/mm artery (Fig. 2b).

3.5. Endothelium-dependent and -independent vasorelaxation

NA-precontracted arteries from chow-fed rats demonstrated progressive relaxation to cumulative addition of CCh (10 nM–100 μ M), achieving a maximum of $88 \pm 2\%$ at a CCh concentration of 100 μ M, with EC_{50} values of 0.28 ± 0.03 μ M. Arteries from diet-fed rats that were similarly exposed to CCh displayed a significant rightward shift of the concentration–response curve, as compared with chow-fed control arteries (EC_{50} diet-fed: 0.94 ± 0.04 μ M, $p = 0.0023$ versus chow-fed) (Fig. 3a). Maximal vasorelaxation to 100 μ M CCh was reduced by 17% in diet-fed rats ($p = 0.0069$ versus chow-fed). Fenofibrate treatment did not improve EC_{50} values (0.86 ± 0.04 μ M, $p = 0.0025$ versus chow-

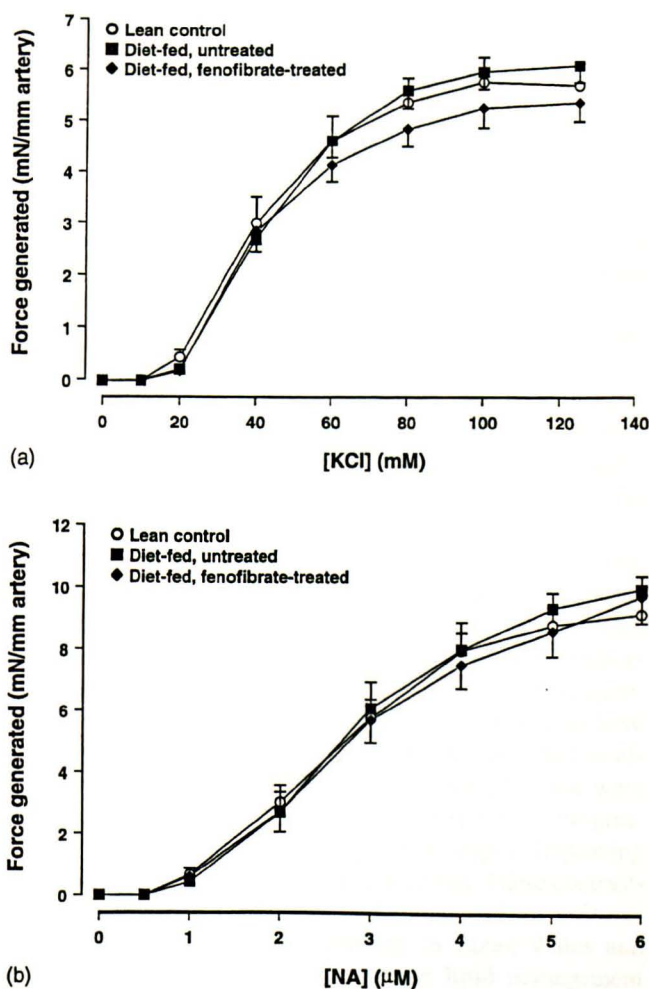


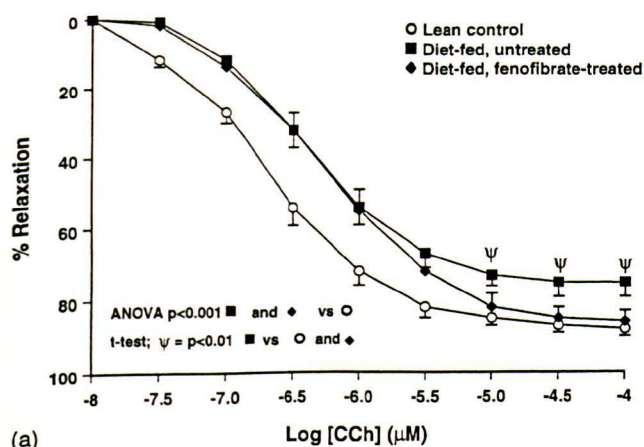
Fig. 2. Vasoconstriction responses to cumulative concentration of (a) KCl (10–125 mM) and (b) noradrenaline (0.5–7 μ M) in arteries from lean control, untreated diet-fed and fenofibrate-treated diet-fed animals. Data represent mean \pm S.E.M. for nine animals in each group. There were no significant differences in agonist-induced vasoconstriction between the three groups.

fed), but it enhanced maximal CCh-induced vasorelaxation at higher concentrations (>10 μ M), achieving $86 \pm 3\%$ relaxation at 100 μ M CCh (Fig. 3a).

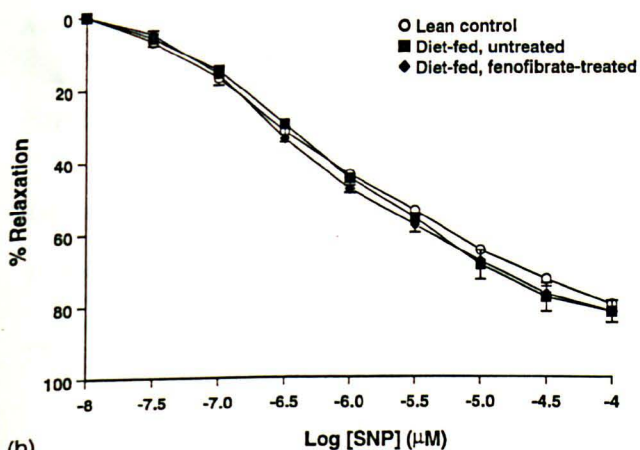
Feeding the high-palatable diet did not affect concentration-dependent vasorelaxation induced by SNP (10 nM–100 μ M). The concentration–relaxation response curves were identical and virtually superimposable in all three groups, with EC_{50} values of 2.08 ± 0.07 (chow-fed), 1.73 ± 0.12 (untreated obese) and 1.46 ± 0.10 μ M (fenofibrate-treated) (Fig. 3b). There were no statistically significant differences between all three groups, either overall (by ANOVA) or at any given SNP concentration.

4. Discussion

Recent studies in animal models and in subjects with diabetes have shown that PPAR α activation reduces diet-induced weight gain and visceral fat mass and improving insulin



(a)



(b)

Fig. 3. The effects of (a) carbamylcholine (CCh), (b) sodium nitroprusside (SNP) on arteries from lean control, untreated diet-fed and fenofibrate-treated diet-fed rats. Noradrenaline ($8 \mu\text{M}$) precontracted arteries challenged with cumulative concentrations of CCh or SNP. CCh-induced vasorelaxation was significantly reduced in diet-fed animals (ANOVA, $p < 0.001$) in comparison to lean control group. Moreover, vasorelaxation curves were shifted to the right in both diet-fed groups. Fenofibrate treatment only partially restored CCh-induced vasorelaxation. Data represent mean \pm S.E.M. for nine animals in each group.

resistance [17,18]. The precise mechanism(s) by which PPAR α improves insulin resistance is not fully known yet, but increasing fatty acid catabolism in liver, decreasing skeletal muscle TG content or decreasing production of cytokines associated with insulin resistance pathways [19], counteracting hypertriglyceridaemia by increasing hepatic oxidation of fatty acids and reduced synthesis and secretion of TG [20] have all been postulated as possible mechanisms by which PPAR α activation improves insulin resistance. Moreover, The mechanism of PPAR α -induced reduction in weight gain also remains controversial. There is a suggestion that a reduction in fat mass and consequential decrease in leptin production may ultimately induce hypophagia and the resultant decrease in weight gain [21], while another study reported lack of any association between plasma levels of leptin, food intake and inhibition of weight gain by PPAR α agonist, fenofibrate treatment [18].

We and others [2,6,22] have shown that dietary obesity not only causes metabolic abnormalities but also markedly affects vascular function, in particular endothelium-dependent agonist-induced vasorelaxation. The vascular abnormalities induced by dietary obesity are complex and multifactorial. Possible factors include fat mass per se, insulin resistance [3], increased NEFA levels [22] hypertriglyceridaemia [2] decrease in plasma HLD-c levels [11] and hyperleptinaemia [4]. The key players could, in theory, be identified through interventions that aim to correct these abnormalities and findings, which leads to the amelioration of vascular dysfunction.

In this study, in agreement with previous reports [16,23] 8-week treatment of fenofibrate reduced total body weight, weekly weight gain and adiposity. Moreover, fenofibrate treatment completely corrected dietary-induced insulin-resistance, hyperinsulinaemia and hypertriglyceridaemia but not hyperleptinaemia.

By contrast, the beneficial effects of fenofibrate on endothelial function were relatively minor and only appeared at higher concentration of carbamylcholine ($>31.6 \mu\text{M}$ CCh). The failure of fenofibrate to completely correct dietary-induced endothelial dysfunction seen in our study is in agreement with various previous reports. Studies by others have not shown any significant changes in endothelial function after acute treatment with 250 mg fenofibrate [24], nor were there any beneficial effects on blood pressure or progression of atherosclerosis process [23,25] despite improving insulin sensitivity and hypertriglyceridaemia. These contrasting findings could be explained by differences in experimental conditions. For example, subjects in Idzior-Walus and colleagues' [26] study had been under lipid management for at least 12 months, establishing a steady state of weight and lipid parameters and those in Malik and colleagues' [27] study did not include obese or insulin-resistant subjects. Another recent study [12] reported that fenofibrate treatment (2 weeks) significantly reduced fasting TG levels and improved both endothelium-dependent and -independent vasorelaxation. Once again, these patients underwent specific dieting for at least 2 weeks before any vascular analyses were carried out. Taking these observations together, it is possible to hypothesise that withdrawal of a fat-enriched diet attenuated vascular dysfunction in these subjects and that this may have further improved by administration of fenofibrate, as seen in our study. Consistent with an effect of diet per se, we have recently reported that acute withdrawal of an obesity-inducing diet partially restores normal endothelial function [28] while chronic withdrawal completely corrects dietary-obesity-induced vascular dysfunction [29], indicating that diet per se rather than dietary-induced obesity plays a key role in vascular abnormalities. However, it is also possible to argue that hyperleptinaemia [4] seen in diet-fed rats (both vehicle- and fenofibrate-treated rats) may, at least in part, explain the limited beneficial effects of fenofibrate on vascular function. Nonetheless, we have previously shown that diet-induced vascular dysfunction occurs irrespective of circulating plasma leptin levels [30].

In conclusion, our study indicates that fenofibrate treatment in the presence of obesity-inducing diet decreases adiposity, lowers plasma TG and NEFA levels and improves insulin resistance in dietary-obese animals, but has little effect on vascular function. The lack of marked beneficial effects of fenofibrate on vascular function may be, at least in part, due to the continuous presence of detrimental factors in obesity-inducing diet, of which high sucrose and/or fats may play important roles. Furthermore, findings from this study indicate that pharmacological attenuation of adiposity alone does not alleviate obesity-induced endothelial dysfunction, and that removal of obesity-inducing diet is an important step in correcting vascular dysfunction.

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