

Development and regulation of body weight: A genetic, behavioral and neuro-endocrinological approach

Citation for published version (APA):

Rutters, F. (2009). Development and regulation of body weight: A genetic, behavioral and neuro-endocrinological approach. Maastricht: Maastricht University.

Document status and date:

Published: 01/01/2009

Document Version:

Publisher's PDF, also known as Version of record

Please check the document version of this publication:

- A submitted manuscript is the version of the article upon submission and before peer-review. There can be important differences between the submitted version and the official published version of record. People interested in the research are advised to contact the author for the final version of the publication, or visit the DOI to the publisher's website.
- The final author version and the galley proof are versions of the publication after peer review.
- The final published version features the final layout of the paper including the volume, issue and page numbers.

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Development and regulation of body weight

A genetic, behavioral and neuro-endocrinological approach

The studies presented in this thesis were performed at the Nutrition and Toxicology Research Institute Maastricht (NUTRIM), which participates in the graduate school VLAG (Food Technology, Agro biotechnology, Nutrition and Health Sciences), accredited by the Royal Netherlands Academy of Arts and Sciences.



Printing of this thesis was financially supported by the J.E. Jurriaanse stichting.

Cover design: Femke Rutters and Jurriaan Born

Cover: Jurriaan Born

Layout: Femke Rutters and Datawyse

Printed by: Datawyse, Universitaire Pers Maastricht

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ISBN: 978 90 5278 852 4

Development and regulation of body weight

A genetic, behavioral and neuro-endocrinological approach

Proefschrift

Ter verkrijging van de graad van doctor
aan de universiteit Maastricht,
op gezag van de Rector Magnificus,
Prof. mr. G.P.M.F. Mols,
volgens het besluit van het College van Decanen,
in het openbaar te verdedigen
op vrijdag 4 september 2009 om 14.00 uur

door

Femke Rutters

Geboren te Waalwijk op 1 september 1982



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Chapter 1

Introduction

General

In healthy adults, body weight normally fluctuates around a relatively constant value, and therefore appears to be regulated with remarkable precision [1]. A stable body weight is a reflection of a balance between energy intake and energy expenditure. A chronic deregulation of energy balance, with energy intake exceeding energy expenditure, leads to the storage of the excessive energy as fat, a characteristic feature of overweight and obesity [2]. Overweight and obesity are classified by the body mass index (BMI) and body composition. According to the World Health Organization, adults with a BMI between 20 and 24.9 kg/m² and body fat percentage of 10-15% in men and 20-25% in women are defined as normal weight, with a BMI between 25.0 and 29.9 kg/m² and body fat percentage of > 15% in men and >25% in women as overweight, and with a BMI between 30.0 and 34.9 kg/m² as obese [3]. In children BMI changes substantially with age, therefore Cole et al. (2000) [4] recently developed age and sex specific cut off points for BMI to be able to define overweight and obesity during childhood. Chronic deregulation of body weight, as seen in obesity, reflects a multi-factorial disorder [4]. Development and regulation of body weight are determined by a number of factors namely genetic, parental, behavioral, endocrine, and the interaction between these factors. The first part of this thesis elaborates on the factors involved in body-weight development during childhood and puberty. Thereafter, the second part of this thesis elaborates on the role of the neuro-endocrine system of the hypothalamus/pituitary/adrenal (HPA) axis in body-weight regulation in adults.

Body-weight development during childhood and puberty

First, relevant genetic factors involved in development of body weight during childhood and puberty will be dealt with. The interaction between genetic factors and body-weight development is typically

studied by means of twin studies, adoption studies and/or single nucleotide polymorphism (SNP) association studies [5].

In this thesis, we focus on polymorphisms that previously have been related to variation in body weight, body composition and/or energy intake such as the ciliary neurotrophic factor (CNTF) gene [6, 7], the peroxisome proliferated-activated receptor γ 2 (PPAR γ 2) gene [8], and the fat mass and obesity-associated gene (FTO) [9-11].

CNTF is a neurocytokine that is involved in control of energy balance, as administration of CNTF leads to weight loss in humans [12]. The null mutation G > A of the CNTF gene leads to an inactive CNTF protein, and is related to a higher BMI [7], lower leptin concentrations, and a smaller decrease in leptin concentrations during weight loss in adults [6]. In children, the CNTF G > A null mutation has not been related to obesity yet. PPAR γ is nuclear receptor protein that functions as a transcription factor in the regulation of adipocyte differentiation [13]. The PPAR γ polymorphism Pro12Ala represents a substitution of proline to alanine in exon B, and leads to decreased transcriptional activity [14]. In children, the PPAR γ polymorphism Pro12Ala is related to deficiency in energy storage and utilization, which leads to reduced growth [13, 15], increased obesity risk [16], increased adiposity [17], and insulin resistance [18, 19].

The FTO gene is involved in the control of energy balance, as FTO expression is decreased in the hypothalamus in fasted mice, when compared to fed mice [20]. In children, the A allele of the FTO gene (rs9939609) is related to a number of obesity related characteristics, such as an increased BMI [11, 21], increased hip circumference [22], increased adiposity [9], increased leptin levels [23, 24], increased energy intake [25, 26], and reduced satiety responsiveness [27]. The influence of these SNP's may change over time as gene expression changes over time [28]. The effects of the SNP's during body-weight development therefore need to be assessed over time.

Parental factors

Parental factors are also important in body-weight development during childhood. Parental factors are a combination of genetic and behavioral factors, and for instance, parental obesity is an important parental factor [29], as the BMI of the child is positively related to the BMI of the parent [29, 30]. Also parents' restrained eating behavior affects children's body-weight development, since the child's body weight at age 12y has been shown to be positively related to the restraint score of the mother [30]. The parental association is due to genetic and environmental factors, such as shared lifestyles (i.e. diets, feeding practices and food choices) and patterns of activity [29].

Behavioral factors

Behavioral factors, such as eating behavior and sleep duration are involved in body-weight development during childhood as well. Eating behavior in general can be analyzed using the Three Factor Eating Questionnaire (TFEQ) [31]. The TFEQ assesses three factors involved in eating behavior: dietary restraint, disinhibition and feeling of hunger. Dietary restraint refers to conscious restriction of food intake to achieve or maintain a preferred body weight. Disinhibition reflects individual differences in the extent to which release from the cognitive suppression of eating occurs in response to the presence of palatable food or other disinhibiting stimuli, such as emotional distress. The third factor refers to the general feeling of hunger [31]. In 11-12 y old children dietary restraint only occurs in the overweight children [30], while in adults dietary restraint also occurs in lean people who are called 'successfully dietary restrained eaters' [32]. The development of dietary restraint in lean people may start during puberty.

In children an inverse relationship between the behavioral factor sleep duration and BMI has been observed [36-38], which suggests that short sleep has an effect on body-weight development. Sleep duration could affect body-weight development [33, 34], as short sleep

duration is already present from a very young age onwards [35]. Some studies even observed a curvilinear relationship between sleep duration and BMI [39, 40], which suggests that sleeping too short and too long have a negative effect on body-weight development. Also, negative relationships have been observed between sleep duration and body fat percentage [41, 42], waist circumference [36], leptin levels [42], as well as insulin resistance [42, 43].

Sleep duration during childhood is however subjected to developmental changes, in particular during puberty when a significant decrease in sleep duration has been observed [35, 44]. Previous longitudinal studies on sleep duration and BMI in children have observed a consistent negative linear association between habitual sleep duration and later obesity [44-47]. However they have not investigated whether changes in sleep duration are associated with changes in BMI.

Endocrine factors

Furthermore, the endocrine factor leptin appears to play a remarkable role in body-weight development during childhood, especially during puberty [48, 49], as leptin is thought to affect development of body weight as well as body composition during puberty [50]. In children and adults, concentrations of the hormone leptin are directly related to the amount of adiposity, since leptin is derived from the adipocytes [51, 52]. Leptin is thought to be a permissive factor for the start of puberty, since puberty is suggested to start when a critical amount of body weight or fat mass is achieved [51-53]. Until children reach puberty, body composition in boys and girls are similar, however, during puberty the percentage of fat mass in girls increases when compared to boys [50]. During puberty and adulthood, a 'healthy' body composition beholds a body fat percentage of 10-15% in boys/men and 20-25% in girls/women [50]. The proposed mechanism behind leptin as a permissive factor for the start of puberty, beholds that leptin independently of fat

mass acts on the hypothalamic luteinizing hormone-releasing hormone (LHRH) pulse generator, which stimulates the release of luteinizing hormone (LH) and follicle-stimulating hormone (FSH) from the pituitary [54-56]. LH and FSH in turn stimulate the gonads to release testosterone and estradiol that will form a negative feedback loop, which will inhibit the secretion of LH and FSH [55, 56]. Moreover, testosterone alone will form a negative feedback loop that will inhibit leptin secretion from the adipocytes, and estradiol will form a positive feedback loop, which stimulates leptin secretion from the adipocytes [56-58]. This suggests that leptin, gonadotrophic hormones, and gonadal hormones are temporarily related during puberty and this may alter the relationship between leptin concentrations and body fat during puberty. The role of leptin during puberty is an interesting issue, which deserves more research.

During adulthood genetic, parental, behavioral, physiological i.e. mainly endocrine factors are also involved in body-weight regulation, however it is suggested that the magnitude of involvement of the factors is different. Body-weight regulation during adulthood is for example affected by behavioral and endocrine factors, such as stress and cortisol concentrations [59]. The hypothalamus/pituitary/adrenal (HPA) axis is involved in the stress-response and secretion of cortisol. The second part of this thesis elaborates on the role of the neuro-endocrine system of the HPA axis in the regulation of body weight in adults.

Hypothalamus/pituitary/adrenal (HPA) axis and body-weight regulation

The second part of this thesis elaborates on the role of the neuro-endocrine system of the HPA axis in body-weight regulation in adults. The HPA axis is involved in the stress-response through regulating the physical and behavioral adaptations to stressors [60]. The cascade of the HPA axis beholds that the hypothalamus

produces corticotropin-releasing hormone (CRH), which subsequently stimulates the production of adrenocorticotropin (ACTH) from the anterior pituitary, which in turn will stimulate the synthesis and release of cortisol by the adrenal cortex. Cortisol exerts its actions through binding to and activation of two types of intracellular receptors, the low affinity glucocorticoid receptor (GR) and the high affinity mineralocorticoid receptor. The GR induces negative feedback of the HPA axis through GRs on the hypothalamus and pituitary. The mineralocorticoid receptor is supposed to regulate basal activity of the HPA axis [60]. Support for the involvement of the HPA axis in body-weight regulation is found in two extremes of plasma cortisol levels in humans; Addison's disease (hypocortisolism) that has been related to weight loss, and Cushing's syndrome (hypercortisolism) that has been related to rapid weight gain, particularly of the trunk and face with sparing of the limbs [61].

Genetic factors

Support for the involvement of the HPA axis in body-weight regulation is also derived from genetic studies. In this thesis we focus on the BclI (rs41423247) polymorphism of the GR gene, which has been related to disturbed body-weight regulation and obesity-related characteristics [62]. The BclI G/G genotype has been related to an increased BMI [63, 64], more visceral fat deposition [63], insulin resistance [65], increased blood pressure and cholesterol levels [66, 67], and larger weight loss during a very low calorie diet [6, 67].

With respect to HPA axis functioning, the BclI polymorphism results in decreased GR density, possibly resulting in altered cortisol sensitivity [62]. Consequently, the BclI G/G genotype is related to variation in HPA axis functioning, such as increased salivary cortisol concentrations [68, 69] and increased cortisol concentrations after a dexamethasone suppression test [69-71]. Remarkably, sex differences were shown in the cortisol response to stress, as men with the BclI G/G genotype show a diminished cortisol response to

psychosocial stress [72], while women show an increased cortisol response. This inconsistency may either be explained by different research protocols, or may suggest a sex specific association between the BclI genotype and HPA axis functioning, which has not been investigated before.

Endocrine factors

Additional support for the involvement of the HPA axis in body-weight regulation is found in humans with visceral fat accumulation. Visceral fat accumulation in obese subjects is related to altered HPA axis functioning, such as decreased salivary and serum cortisol levels [73-78], increased urinary secretion of cortisol [76, 79], decreased cortisol variability [73, 74, 80-82], and enhanced cortisol awakening response [83]. In both lean and obese subjects, visceral fat distribution has been associated with increased cortisol secretion after physical and psychological stressors [73-76, 79, 84]. However studies on the relationship between negative feedback of the HPA axis and body fat distribution have yielded inconclusive results. Some studies find an increased capacity of dexamethasone (a GR agonist) to suppress plasma cortisol concentrations [74, 85], while some studies find no differences in the capacity of dexamethasone to suppress plasma cortisol concentrations [73, 75, 76, 80], or some find even a decreased capacity [86]. This inconclusiveness possibly results from differences in techniques to test HPA feedback functioning. Previously, Petrides et al. (1997)[87] developed a technique to test HPA axis feedback functioning under a physical stressor. Using this technique they observed, even in a relatively homogenous population of young men, a large inter-individual variance in HPA axis feedback functioning [87]. It however remains unknown whether intra-individual variation in HPA axis functioning can be related to fat distribution in a normal-weight to obese population, and whether HPA axis feedback functioning tested under a physical stressor is related to fat distribution.

Behavioral factors

Final support for the involvement of the HPA axis in body-weight regulation is found in the involvement of the HPA axis in energy intake. Most of the data directly indicating a crucial role of the HPA axis in the regulation in food intake are derived from animal studies, showing that the anorectic effects of adrenalectomy can be reversed by corticosterone replacement [88, 89]. Human studies support the orexigenic effect of cortisol, as cortisol administration stimulates food intake [90]. The orexigenic effect of cortisol may underlie the reported increase of energy intake after stress, as psychological stress seems to increase the intake frequency, increase the amount of food taken in [91-93], and alter food preference towards intake of sweet and fat foods [94]. The alterations in energy intake during psychological stress have been suggested to relate to non-homeostatic regulatory mechanisms involved in eating behavior, such as reward [95]. Dallman and colleagues [96] even introduced the term 'comfort foods' to emphasize the fact that the drive behind stress-induced eating is not necessarily a homeostatic need for calories, translated into feelings of hunger.

Reward is defined by two distinct processes namely 'liking' and 'wanting' [97], and involves the neural networks of the opioid and dopaminergic system [97, 98]. Disturbance of these neural networks is thought to underlie stress-induced eating, as HPA axis activation leads to a decrease in opioid sensitivity in rats [99], and increased dopamine levels in rats [100]. Altered levels through administration of antagonists of opioid and dopamine lead to a reduction of food intake and reduced pleasantness of food in humans [101, 102]. The involvement of non-homeostatic regulatory mechanisms, such as reward, in stress-induced eating has however never been directly demonstrated in humans. Moreover, previous studies in humans could not exclude a role for hunger [103].

In humans, stress affects eating in a bidirectional way; in a subgroup of about

30% stress decreases food intake, indicating that large inter-individual differences are present. These individual differences in response to stress may relate to eating behavior characteristics [92-94, 104], as these are determined using the Three Factor Eating Questionnaire (TFEQ) [31]. An increase in energy intake during stress has been found in individuals with high scores on dietary restraint and/or disinhibition [92-94, 104]. Moreover, subjects that have a high cortisol response to a laboratory stressor have a larger energy intake after the stressor when compared to subjects with a low cortisol response [92]. These data suggest that there may be a link between cortisol response and eating behavior characteristics. For instance weight loss and/or weight maintenance are experienced as demanding tests, and can therefore be perceived as stressors, even in normal weight women [105-107]. Several studies have shown increased cortisol concentrations (86-88), in normal weight women who attempt to cognitively control their food intake (high dietary restraint score) when compared to those who do not (low dietary restraint score). When dietary restraint is considered as a chronic physiological stressor, it may lead to alterations in HPA axis functioning, such as a decreased sensitivity of the GR to cortisol, which results in decreased negative feedback of the HPA axis [59]. It remains unknown however, whether dietary restraint affects cortisol feedback functioning.

Outline of this thesis

A stable body weight reflects a balance between energy intake and energy expenditure. This thesis encompasses genetic, parental, behavioral, physiological i.e. mainly endocrine factors involved in body-weight development and regulation during childhood, puberty and adulthood.

From a large cohort of Dutch children born between 1990 and 1993 valuable anthropometrical data from birth till age 7y are available. To study body-weight development during puberty, follow-up

studies were performed yearly between 2004 and 2008, when the children had a mean age of 12 to 16y. The roles of relevant genetic factors, such as the FTO gene, in body-weight development are described in **chapter 2**. Subsequently, the assessment of the roles of parental, behavioral, and physiological i.e. mainly endocrine factors in body-weight development during puberty are dealt with in **chapter 3 and 4**.

The amount of adiposity is directly related to leptin concentrations, which is thought to be a permissive factor for the start of puberty, since puberty starts when a critical amount of body weight or fat mass is achieved. The changing relationship between leptin and gonadotropic hormones during puberty is described in **chapter 5**.

During adulthood it is suggested that the magnitude of effect of the factors involved in body-weight regulation is different. Body-weight regulation is for example affected by behavioral and endocrine factors, such as stress and cortisol concentrations. Through the secretion of cortisol, the HPA axis plays an important role in the regulation of the stress response. The present knowledge of the role of the HPA axis in the body-weight regulation in adults is described in **chapter 6**. Subsequently, the role of genetic factors, such as the BclI polymorphism, and physiological factors, such as body composition, in HPA axis functioning are dealt with in **chapter 7 and 8**.

Stress has been related to alterations in eating behavior, such as an increase or decrease in energy intake. Assessment of food intake in the absence of hunger after psychological stress exposure is given in **chapter 9**. The intra-individual differences in the effect of stress on food intake may relate to eating behavior characteristics, such as dietary restraint, which in turn is related to alterations in HPA axis functioning. Weight loss and/or weight maintenance can be perceived as a stressor, even in normal weight women, and therefore dietary restraint may lead to alterations in HPA axis functioning as dealt with in **chapter 10**.

Finally in **chapter 11**, the results and interpretations of the previous described studies are discussed and put into a broader perspective.

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Chapter 2

Associations between the FTO gene (rs9939609) and development of obesity related characteristics over time in a Dutch children cohort

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Submitted

Abstract

Objective: to replicate earlier findings on the relationship between FTO and obesity related characteristics, as well as investigate the effect of FTO on BMI, body composition, leptin concentrations, and physical activity longitudinally.

Subjects and measurements: FTO (rs9939609) was genotyped in 101 children, from whom we have collected anthropometric measurements from birth to age 15y, as well as body composition, leptin concentrations, and physical activity (Baecke questionnaire) yearly from age 12 to 15y.

Results: children with the A/A and A/T genotype had a higher BMI at age 12, 13, 14, 15y (19.5 ± 2.9 vs. 18.3 ± 1.9 , 20.2 ± 3.3 vs. 18.6 ± 2.0 , 21.0 ± 3.2 vs. 20.0 ± 2.1 , 21.9 ± 3.6 vs. 20.8 ± 2.2 kg/m², $p<0.05$), and a larger increase in BMI from age 7-15y (6.3 ± 2.9 vs. 5.4 ± 1.6 kg/m², $p<0.05$), when compared to children with the T/T genotype. Additionally, children with the A/A and A/T genotype had a higher fat mass index (FMI) at age 12 and 13y (4.2 ± 2.1 vs. 3.4 ± 1.5 kg/m² and 4.3 ± 2.3 vs. 3.4 ± 1.5 kg/m², $p<0.03$), a lower Baecke score at age 13y (7.8 ± 0.8 vs. 8.2 ± 0.8 , $p<0.03$), higher leptin concentrations at age 12 and 13y (8.7 ± 7.3 vs. 5.9 ± 4.5 ng/ml, $p<0.02$ and 6.8 ± 6.1 vs. 4.5 ± 3.8 ng/ml, $p<0.05$), as well as a larger leptin peak at age 12y, when compared to children with the T/T genotype. At age 13y, an association was observed between BMI and leptin concentrations (ng/ml), Baecke scores, and the FTO A allele ($R^2=0.523$, $p<0.001$).

Conclusion: the FTO A allele (rs9939609) is associated with higher BMI, FMI, leptin concentrations, and lower activity scores from childhood to puberty; these associations are independent and change in strength over time.

Introduction

The prevalence of obesity is emerging as a major health problem [1] and is associated with several risk factors, such as an increased risk on cardiovascular disease and type 2 diabetes [2, 3]. Individual susceptibility to obesity is determined by interactions between genetic, behavioral, and environmental factors [4]. To study the interaction between genetics and obesity often single nucleotide polymorphism (SNP) association studies are used [5]. Recent SNP association studies identified a relationship between the FTO (fat mass and obesity associated) gene (rs9939609) and the control of energy balance, since FTO expression is decreased in the hypothalamus in fasted mice, when compared to fed mice [6]. SNP association studies have linked the A allele of the FTO gene (rs9939609) to a number of obesity related characteristics, such as an increased BMI [7-10], increased hip circumference [11], increased adiposity [5, 7, 8], increased leptin levels [12, 13], increased C-reactive protein levels [14], elevated dietary fat intake [15, 16], increased energy intake [17, 18], and reduced satiety responsiveness [19]. Additionally, Andreasen et al. observed an interaction between the FTO genotype and physical activity, where physically inactive subjects with the AA genotype had a higher BMI when compared to T/T genotype [7]. Previous studies, however, only assessed the effect of FTO polymorphisms in one single population at one time point [7, 8, 14-17, 19], or assessed only BMI longitudinally [9, 12, 20, 21]. Gene expression however changes developmentally [22], and so the influence of FTO may change over time. It therefore remains unknown whether the effect of FTO is stable during life on other obesity related characteristics, such as body composition, leptin concentrations, as well as physical activity, and whether FTO influences the obesity related characteristics independently.

Consequently, the objective of our study was to replicate earlier findings on the relationship between FTO and obesity related characteristics, as well as investigate the effect of FTO (rs9939609) on BMI, body composition, leptin concentrations, and physical activity longitudinally. Therefore, we determined in 101 children anthropometric measurements yearly from birth to age 15y, as well as body composition, leptin concentrations, and physical activity yearly from age 12 to 15y.

Subjects and methods

Subjects

Subjects were recruited from a Dutch Caucasian cohort of children born between 1990 and 1993 [23, 24]. Anthropometric data were available from these children, no interventions were executed, and follow-up studies were performed with 58 boys and 43 girls. Each child and their parents gave written informed consent to participate in the study, which was approved by the Central Committee Human Research and by the Medical Ethical Committee of Maastricht University.

Study design

Children's body weight (BW), height, body mass index (BMI), waist circumference, body composition [24, 25], leptin concentrations, as well as physical activity were determined in the afternoon after a 3-hour fast.

Measurements

Anthropometry

The children's BW was determined using a digital balance accurate to 0.1 kg (Sauter D7470, Ebingen, Germany) and their height was determined using a wall-mounted stadiometer (Seca, model 220, Hamburg, Germany). Measurements were executed in underwear, after a 3-hour fast, and after voiding the bladder. BMI was calculated by $BW/height^2$ (kg/m^2) and BMI changed substantially with age in childhood. To define normal weight, overweight and obesity in children, we

used the specific cut off points described by Cole et al. 2000 [2]. The waist circumference was measured at the site of the smallest circumference between the rib cage and the ileac crest, with the subjects in standing position. The pubertal stage (development of breast, genitalia, and pubic hair) was determined in all children, according to the classification by Tanner [26]. Anthropometric measurements were determined at age 7, 12, 13, 14 and 15y.

Body composition

Body composition was measured using the deuterium dilution technique (D₂O). D₂O dilution was used to measure total body water (TBW). Subjects were asked to collect a urine sample in the evening just before drinking the deuterium-enriched water solution. After ingestion of this solution, no further consumption was allowed. Ten hours after drinking the water solution, another urine sample was collected. The dilution of the deuterium isotope is a measure of the TBW of the subject. Deuterium was measured in the urine samples with an isotope ratio mass spectrometer (VG-Isogas Aqua Sira, VG Isogas, Middlewich, Cheshire, England). TBW was obtained by dividing the measured deuterium dilution space by 1.04. Fat free mass (FFM) was calculated by dividing TBW by the hydration factor 0.73 [27-29]. Fat mass (FM) was determined as BW-FFM. Fat mass index (FMI) was calculated by fat mass/height² (kg/m²) and fat free mass index (FFMI) was calculated by fat free mass/height² (kg/m²). Body composition was measured at age 12,13,14 and 15y.

Leptin

Plasma leptin concentrations were measured with a double-antibody, sandwich-type enzyme-linked immunosorbent assay that used a monoclonal antibody specific for human leptin. The lower and upper limits of detection were 0.5 µg/L and 50 µg/L respectively. The intra- and inter assay CVs were 9% and 12%, respectively. The leptin concentrations of normal-weight subject's range from 2 to 12 µg/L. Plasma

leptin concentrations were measured at age 7, 12, 13, 14 and 15y.

Physical activity

Physical activity was measured with the Baecke questionnaire. The Baecke questionnaire consists of three components: work activity, sports activity and leisure activity, and has been validated using doubly labeled water [30]. For the children, the work index was replaced by a school index with similar questions (e.g. 'When I am at school, I walk: never/seldom/sometimes/often/all the time?'). The Baecke questionnaire for children was validated by Vogels et al. 2007 [31]. Physical activity was determined at age 12, 13, 14 and 15y.

Determination of FTO genotypes

The genomic DNAs of 101 participants were isolated from peripheral blood leukocytes using a QIAamp kit (QIAGEN, Germany). FTO (rs9939609) genotypes were determined using Taqman allelic discrimination (AB, Applied Biosystems).

Statistical analysis

Student t-tests (for continuous variables) and chi-square tests (for the nominal variables) were executed to determine differences in single variables between groups. Differences over time and between conditions were determined using two-factor ANOVA with repeated measures. Relationships between dependent and independent variables were determined using simple linear regression and multiple regression models.

The genotypic and allelic distributions of the FTO polymorphism are provided in **table 1**.

Table 1: FTO genotype frequencies and the frequency of the A allele in the study sample (n = 101)

Poly-morphism	Genotype	F (n)	F (%)	Allele	F (%)	HWE ^a
Rs9939609	TT	45	44.6	T	67.8	0.93
T > A	TA	47	45.5	A	32.1	
	AA	9	8.9			

F, Frequency, either absolute (n) or relative (%);

^a P-values obtained from the Chi-square test of Hardy Weinberg Equilibrium (HWE)

The genotype frequencies were in Hardy Weinberg Equilibrium. Because of the small study population and since the data of the homozygotes for the minor allele (A/A) did not differ significantly from the heterozygotes (A/T), they were taken together in the analysis and compared to the homozygotes for the major allele (T/T).

All tests were two-sided and differences were considered significant at $P < 0.05$. Values are expressed as mean \pm standard deviation (s.d.).

Results

Data on height, BW, BMI, waist circumference, body composition, leptin concentrations, and Baecke scores were collected at age 7 to 15y from 58 boys and 43 girls. The subject characteristics at age 15y are presented in **table 2**. At age 15y most children were at Tanner stage 3 of the pubertal development, and 11% of the children were overweight. Overweight status measured as BMI and body fat percentage, was defined on the basis of age- and sex-appropriate international standards [2]. With respect to genotypic distribution of the FTO polymorphism more overweight and obese children had the A/A and T/A genotype when compared to the T/T genotype at age 12 (14.3 and 31.4 vs. 4.9%) and 15y (28.6 and 31.3 vs. 5.1%).

Table 2: Characteristics of the boys (n = 58) and girls (n = 43) at the mean age of 15y

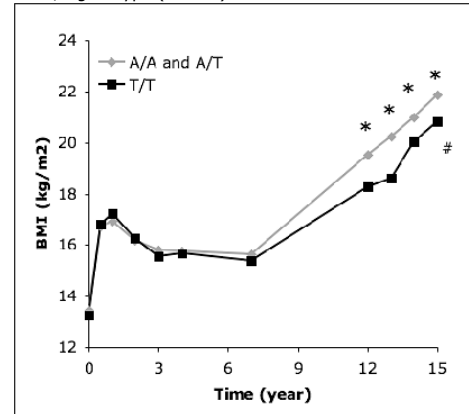
	Boys	Girls	P-value
Body weight (kg)	65.4 \pm 10.8	60.5 \pm 10.8	0.02
Height (cm)	176.1 \pm 7.2	165.8 \pm 6.8	0.01
Body mass index (kg/m ²)	20.9 \pm 2.7	21.9 \pm 3.5	0.18
Waist (cm)	73.4 \pm 6.1	70.4 \pm 7.4	0.01
Body fat (%)	15.9 \pm 6.0	27.5 \pm 5.7	0.01
Leptin (μ g/L)	2.2 \pm 1.9	9.8 \pm 6.5	0.01
Baecke score	8.3 \pm 0.9	8.1 \pm 1.0	0.25

BMI

Figure 1 depicts the development of the BMI (kg/m²) of children with the A/A and A/T genotype versus the T/T genotype over time. BMI was not different between birth and age 7, however BMI was significantly higher at age 12, 13, 14, and 15y in children with the A/A and A/T genotype, when compared to children with the T/T genotype. Over time, a significant

increase in BMI was seen in all groups from age 7 to 15y, and in children with the A/A and A/T genotype a stronger increase was observed, when compared to children with the T/T genotype (6.3 \pm 2.9 vs. 5.4 \pm 1.6 kg/m², $p < 0.05$).

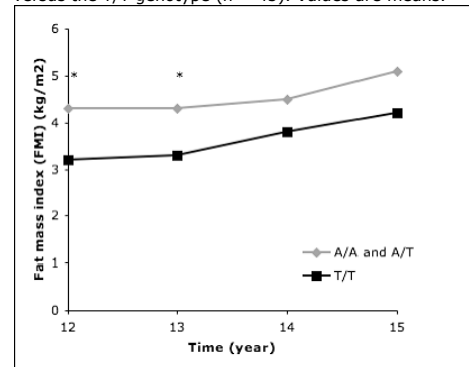
Figure 1: Development of body mass index (kg/m²) in children with the A/A and A/T genotype (n = 56) versus the T/T genotype (n = 45). Values are means.



$P < 0.05$ overall group \times time interaction between A/A and A/T genotype vs. T/T genotype (2- factor ANOVA repeated measures)

* $P < 0.05$ for differences in children with the A/A and A/T genotype vs. T/T genotype (2- factor ANOVA repeated measures)

Figure 2: Development of fat mass index (FMI)(kg/m²) in children with the A/A and A/T genotype (n = 56) versus the T/T genotype (n = 45). Values are means.



* $P < 0.05$ for differences in children with the A/A and A/T genotype vs. T/T genotype (2- factor ANOVA repeated measures)

Fat Mass Index (FMI)

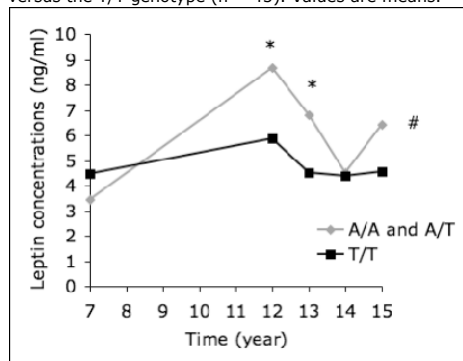
Fat mass index (FMI) was calculated by fat mass/height² (kg/m²). **Figure 2** depicts the development of the FMI (kg/m²) of children with the A/A and A/T genotype versus the T/T genotype over time. FMI was significantly higher at age

12 and 13y in children with the A/A and A/T genotype, when compared to children with the T/T genotype. Over time no significant changes in FMI were present. Surprisingly, no differences over time were found in FFMI (kg/m^2) between the A/A and A/T genotype versus the T/T genotype.

Leptin concentrations

Figure 3 depicts the development of the leptin concentrations (ng/ml) of children with the A/A and A/T genotype versus the T/T genotype over time. Leptin concentrations were significantly higher at age 12 and 13y in children with the A/A and A/T genotype, when compared to children with the T/T genotype. A significant group x time effect was found in leptin concentration measurements ($p < 0.03$, two-factor ANOVA repeated measures). A significant higher peak value was observed in children with the A/A and A/T genotype, when compared to children with the T/T genotype, as a stronger increase between 7 and 12y (5.7 ± 8.3 vs. 1.0 ± 5.3 ng/ml , $p < 0.01$) and a stronger decrease between 12 and 14y (-2.2 ± 3.2 vs. -1.2 ± 5.3 ng/ml , $p < 0.10$) was observed in children with the A/A and A/T genotype.

Figure 3: Development of leptin concentrations (ng/ml) in children with the A/A and A/T genotype ($n = 56$) versus the T/T genotype ($n = 45$). Values are means.



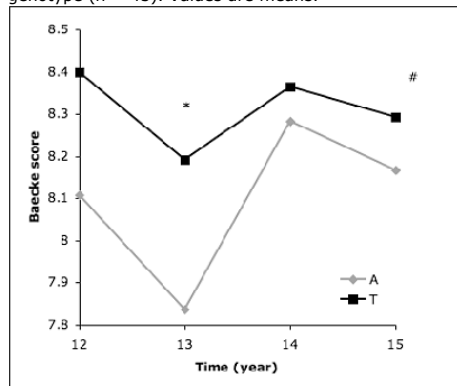
$P < 0.05$ overall group x time interaction between A/A and A/T genotype vs. T/T genotype (2- factor ANOVA repeated measures)

* $P < 0.05$ for differences in children with the A/A and A/T genotype vs. T/T genotype (2- factor ANOVA repeated measures)

Activity scores

Figure 4 depicts the development of the Baecke scores of children with the A/A and A/T genotype versus the T/T genotype over time. Baecke scores were significantly higher at age 13y in children with the T/T genotype, when compared to children with the A/A and A/T genotype. When adjusted for BMI, the association between FTO and Baecke scores remained. Over time Baecke scores in children with the T/T genotype were significantly higher, when compared to children with the A/A and A/T genotype ($p < 0.05$). No alterations over time in Baecke scores were observed in both groups.

Figure 4: Development of Baecke scores in children with the A/A and A/T genotype ($n = 56$) versus the T/T genotype ($n = 45$). Values are means.



$P < 0.05$ overall group x time interaction between A/A and A/T genotype vs. T/T genotype (2- factor ANOVA repeated measures)

* $P < 0.05$ for differences in children with the A/A and A/T genotype vs. T/T genotype (2- factor ANOVA repeated measures)

Table 3A depicts linear regression analyses with FTO allele genotypes ($A=1/T=0$) at a certain age as the dependent variable and BMI (kg/m^2) at that age as the independent variable. The association between FTO allele genotype and BMI were not significant until age 12y and became weaker again from age 13y. Per year, multiple regression analyses were performed with BMI (kg/m^2) as the dependent variable and leptin concentrations (ng/ml), Baecke scores, and FTO allele as the independent variables. Multiple regression analyses with BMI (kg/m^2) at a certain age as the

dependent variable showed that only the leptin concentrations contributed significantly to the model at age 12y ($R^2 = 0.45$, $p < 0.01$), at age 14y ($R^2 = 0.17$, $p < 0.05$), and at age 15y ($R^2 = 0.44$, $p < 0.01$).

Only at age 13y a significant association was observed between all 3 independent variables, and therefore **table 3B** depicts the multiple regression analysis with BMI (kg/m^2) at age 13y as the dependent variable and leptin concentrations (ng/ml), Baecke scores, and FTO allele genotypes at age 13y as the independent variables ($R^2 = 0.523$, $p < 0.001$).

Table 3A: linear regression analysis with FTO allele genotypes (A=1/T=0) at a certain age as the dependent variable and BMI (kg/m^2) at that age as the independent variable

Age (year)	Correlation coefficient (r)	R ²	P-value
0	0.09	0.008	0.38
1	0.10	0.01	0.34
2	0.02	0.001	0.81
3	0.10	0.01	0.37
4	0.04	0.002	0.71
7	0.08	0.007	0.42
12	0.24	0.058	0.02
13	0.28	0.078	0.001
14	0.17	0.029	0.09
15	0.17	0.030	0.09

Table 3B: multiple regression with BMI (kg/m^2) at age 13y as the dependent variable and leptin concentrations (ng/ml), Baecke scores, and FTO allele genotypes at age 13y as the independent variables

	Partial β	Std. Error	P value
Intercept	10.32	2.5	0.001
Leptin concentrations (ng/ml)	0.36	0.047	0.001
Baecke scores	0.82	0.30	0.007
FTO allele genotype (A=1/T=0)	1.30	0.52	0.01

$R^2 = 0.523$, $p < 0.001$

Discussion

The objective of our study was to replicate earlier findings on the relationship between FTO and obesity related characteristics, as well as investigate the effect of FTO (rs9939609) on BMI, body composition, leptin concentrations, and physical activity longitudinally. We observed that children with the A/A and A/T genotype had a higher BMI at age 12 to 15y, and a larger increase in BMI from age 7-15y, when compared to children with the T/T genotype. Children with the A/A and A/T genotype also had a higher

FMI at age 12 and 13y, lower Baecke score at age 13y, higher leptin concentrations at age 12y, as well as a larger leptin peak at age 12y when compared to children with the T/T genotype. Additionally, the association between the FTO gene and BMI increased with age, and only at age 13y BMI was associated with leptin concentrations, Baecke scores and the FTO A allele.

The strengths of our study is the fact that we have longitudinal data from a cohort of solely Caucasian children, thereby eliminating population stratification, as well as determination of body fat percentage using a high-quality method, such as the deuterium dilution technique [29]. Although the sample seems rather small, the genotype frequency distribution of our cohort was consistent with the frequency distributions in other Caucasian populations [7, 17], and because the minor allele frequency is over 30%, sufficient power will be present to study differences in physiological parameters. This reassured us that the use of the FTO genotype in this smaller cohort was a valid approach. In concordance with previous studies [9, 10, 20, 21], we observed a significant association between FTO and BMI during childhood, and no association at birth. Furthermore, we observed a higher BMI from 7y onwards in subjects with the A/A and A/T genotype, which is supported by two other longitudinal studies [20, 21], but is in contrast to the longitudinal study by Jess et al. 2008 [9]. Differences may be explained by time of measurement, while we measured the children yearly, Jess et al. 2008 [9] only measured the children at birth, 7, 10, and 13y, thereby they possibly did not observe changes in BMI during puberty. We however observed that the association between the FTO gene and BMI increased with age, and became significant at age 12y, which has been reported by Haworth et al [20]. Overall, these results imply that the FTO A allele is associated with a higher BMI during early childhood, and that the association becomes stronger during puberty [20].

To determine if the FTO polymorphism was associated with differences in development of body composition and physical activity, we measured body composition and Baecke scores from age 12 to 15y. Children with the A/A and A/T genotype had a significantly higher FMI at age 12 and 13y, as well as lower Baecke scores over time, when compared to children with the T/T genotype. These results confirm several cross-sectional studies on FTO and body composition [5, 7, 8]. Previous studies observed inconsistent results on the relationship between FTO genotypes and physical activity; Andreasen et al [7] in concordance to our results observed an association, while others have observed no associations [21, 32]. Differences between the studies may be explained different study populations and different measurement techniques. Additionally, we measured leptin concentrations from age 7 to 15y and observed that children with the A/A and A/T genotype showed higher leptin concentrations at age 12y, which confirms several cross-sectional studies [12, 13]. Furthermore, we observed that children with the A/A and A/T genotype had a larger leptin peak at age 12y. The leptin peak has been observed to be the onset of sexual maturation [33, 34], and a larger leptin peak may result in increased gonadal hormones concentrations that in turn alter the relationship between leptin concentrations and body fat [35]. Together, these results suggest that the associations between the FTO A allele and decreased physical activity, increased FMI along with increased leptin concentrations are stable during puberty.

Multiple linear regression analysis showed that at age 13y a larger BMI was associated to the FTO A allele, larger leptin concentrations, and decreased physical activity. These results implicate that at age 13y genetic, physiological and

psychological factors are independent predictors for BMI. At age 12, 14, and 15y BMI was only associated to leptin concentrations, which implicates that during puberty BMI is only temporarily predicted by FTO. This has previously been suggested by Haworth et al. [20], which implicates that during puberty BMI is predominately predicted by physiological factors. The strong point of the present study is the fact that we found associations that change in strength over time, between FTO and a number of obesity related characteristics in this relatively small cohort. Our study can however not elucidate causality concerning the development of the obesity related characteristics. We conclude that the FTO A allele (rs9939609) is associated with higher BMI, FMI, leptin concentrations, and lower activity scores from childhood to puberty; these associations are independent and change in strength over time.

Acknowledgements

We want to thank our subjects for their participation in our cohort study. We gratefully thank Loek Wouters and Wendy Sluijsmans for their assistance. FR carried out the study, collected and analyzed the data, and wrote the largest part of the manuscript. FB (supervised by EM) contributed to the practical work. AN and MW supervised FR. Planning, processing the results and writing the manuscript were done under general supervision by AN and MW. The authors had no conflict of interest.

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Chapter 3

Leptin-adiposity relationship changes, plus behavioral and parental factors, are involved in the development of body weight in a Dutch children cohort

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Physiology and behavior 2008 March 18; 93 (4-5); 967-74

Abstract

Background: the development of body weight is determined by different factors, namely genetic, behavioral, parental and physiological.

Objective: to investigate whether genetic, behavioral, parental and physiological factors are involved and the extent of involvement in the development of body weight at age 12 and 13y in a Dutch children cohort.

Methods: In a Dutch cohort of 94 children at age 12 and 13y, we determined anthropometric measurements, body composition, leptin concentrations, TFEQ scores, physical activity, as well as 3 polymorphisms, and determined anthropometric measurements and TFEQ scores in the parents.

Results: 11% of the children in the cohort were classified as overweight. The genotype frequency distributions of the PPAR γ 2, BCL1 and CNTF genes) at age 12 and 13y were not significantly different for the overweight children compared to the lean children. Overweight children showed higher dietary restraint and disinhibition scores. Overweight children's parents had a higher BMI, dietary restraint and disinhibition scores, compared to lean children's parents. A peak in leptin concentrations between 7 to 13y was shown at 12y. In lean boys, the decrease in leptin concentrations between 12 and 13y was related to an increase in fat free mass. At the age of 12y predominantly the physiological factors were predictors for body weight, and at the age of 13y both the physiological and behavioral factors were predictors for body weight.

Conclusion: we conclude from this longitudinal study, that leptin appeared to play an important role in the development of body weight during puberty, in addition to behavioral and parental factors.

Introduction

The prevalence of childhood overweight is emerging as a major health problem [1] and is associated with several risk factors, such as the increased risk on overweight during adulthood [2, 3]. The development of body weight is determined by different factors, namely genetic, behavioral, parental and physiological factors [3-6]. Several genetic factors have been linked to the development of body weight, three of them being the polymorphisms of the peroxisome proliferators-activated receptor (PPAR), the glucocorticoid receptor (GR) and the neurocytokine ciliary neurotrophic factor (CNTF). PPAR γ is a nuclear fatty acid receptor that is involved in adipocyte differentiation and represents a direct link between adiposity, food response and control of appetite in adults [7-9]. In children a link was found between the PPAR γ 2 polymorphism Pro12Ala and obesity, as Cecil et al. found that the Proala12 mutation is related to deficiency in energy storage and utilization, leading to reduced growth [10]. The GR has an important role in the metabolism of adipose tissue and in the regulation of abdominal fat distribution. Mutations in the GR gene may result in perturbation of the glucocorticoid receptor, leading to diminished sensitivity to cortisol, which could lead to stress-induced obesity. Adults with the GR polymorphism IVS2-BclI have been associated with increased BMI, waist-to-hip circumference and leptin concentrations [11, 12]. In children the BclI polymorphism has been related to a larger increase in adiposity during puberty [13]. CNTF is a neurocytokine that works via the signaling pathway normally activated by leptin. The null mutation G > A leads to an inactive CNTF protein, resulting in diminished initiation of anorectic pathways, which leads to an increase in body mass. In adults with the G > A null mutation, a higher BMI and a smaller decrease in leptin concentrations during weight loss, have been found [8, 14]. In children, no associations between

the CNTF G > A null mutation and obesity have been found yet.

Several behavioral factors such as eating behavior and physical activity have been related to the development of body weight. Eating behavior can be analyzed using the Three Factor Eating Questionnaire (TFEQ) and assesses 3 factors involved in eating behavior: dietary restraint, disinhibition and feeling of hunger. Dietary restraint refers to conscious restriction of food intake to achieve or maintain a preferred bodyweight. Disinhibition reflects individual differences in the extent to which release from the cognitive suppression of eating occurs in response to the presence of palatable food or other disinhibiting stimuli, such as emotional distress [15, 16]. Before, we have shown that dietary restraint only occurred in the 11-12 y old overweight children, thus it appeared to be an effect rather than a cause of overweight [6]. Additionally, physical activity can be analyzed using the BAECKE questionnaire, as several studies have shown that increased physical activity levels are protection against overweight development [17, 18].

Furthermore, several parental factors have been related to the development of body weight, like the fact that children with obese parents are more likely to become obese themselves [19]. It is likely that the family association is due to genetic factors and partly to shared lifestyles (i.e. diets and patterns of activity). Early experiences with food, feeding practices and family food choices affect children's nutritional habits. Additionally, parents' restrained eating behavior may affect children's body-weight development as dietary restraint of mothers, and their perception about their daughters' risk of overweight, predicted maternal child feeding practices. This in turn predicted the daughters' eating and relative weight [20].

In addition several physiological, especially endocrine factors have been

related to the development of body weight. The start of puberty around the age of 12y is a period of pronounced changes in the endocrine milieu. For the start of puberty, the achievement of a critical body weight or fat mass is thought to play a prominent role. In children and adults, leptin concentrations are directly related to the amount of adiposity, consequently the hormone leptin has been proposed as a physiological link between adiposity status and the start of sexual maturation [21, 22]. However, most studies investigating this physiological link were done in transversal cohorts, and not yet in a longitudinal cohort.

Overall, the objective of our study was to investigate whether genetic, behavioral, parental and physiological factors are involved and the extent of involvement in the development of body weight at age 12 and 13y in a Dutch children cohort. Therefore, we determined in a Dutch cohort of 94 children at age 12 and 13y anthropometric measurements, body composition, leptin concentrations, TFEQ scores, physical activity, as well as 3 polymorphisms, and determined anthropometric measurements and TFEQ scores in the parents. We hypothesized that at the age of 12 and 13y body weight may be influenced by predominantly physiological factors, in addition to the genetic, behavioral, and parental factors that we previously described before in Vogels et al. 2006 [6].

Subjects and methods

Subjects

Subjects were recruited from a Dutch Caucasian cohort of children born between 1990 and 1993 [6, 23]. As infants, these children and their mothers participated in i) a study of essential fatty acids during pregnancy and pregnancy outcome [23], and ii) a study, performed between 1997 and 2000, about the long-term effects of fetal essential fatty acid availability [24]. Anthropometric data were available from these children and no interventions were provided. To evaluate the development of obesity and related parameters, follow-up

studies were performed in 2004 and 2005 with 94 children [6]. Each child and one of his or her parents gave written informed consent to participate in the study, which was approved by the CCMO (Central Committee Human Research) and by the MEC (Medical Ethical Committee) of the University Maastricht.

Study design

Children's body weight (BW), height, body mass index (BMI), waist circumference, body composition, leptin concentrations, Three Factor Eating Questionnaire (TFEQ) scores and physical activity were determined in 2004 [6] and 2005, when the mean age was 12.4 and 13.4 years (range 11-15 years), respectively. Moreover, the polymorphisms of three relevant genes (PPAR γ 2, BclI and CTNF) were determined [23, 24]. BW, height, BMI and TFEQ scores of both parents were determined in 2005.

Measurements

Anthropometry

At the mean age of 12 and 13 year, the children's BW was measured using a digital balance accurate to 0.1 kg (Sauter D7470, Ebingen, Germany) and height was measured using a wall-mounted stadiometer (Seca, model 220, Hamburg, Germany). Measurements were executed in underwear, after an overnight fast and after voiding the bladder. BMI was calculated by BW/height² (kg/ m²) and BMI in childhood changes substantially with age. To define normal, overweight and obesity in children, we used the specific cut of points described by Cole et al. 2000 [2]. The waist circumference was measured at the site of the smallest circumference between the rib cage and the ileac crest, with the subjects in standing position. The pubertal stage (development of breast, genitalia, and pubic hair) was documented in all children, according to the classification by Tanner.

Body composition

Body composition was measured using the deuterium dilution technique (D₂O). D₂O dilution was used to measure total body

water (TBW). Subjects were asked to collect a urine sample in the evening just before drinking the deuterium-enriched water solution. After ingestion of this solution, no further consumption was allowed. Ten hours after drinking the water solution, another urine sample was collected. The dilution of the deuterium isotope is a measure of the TBW of the subject. Deuterium was measured in the urine samples with an isotope ratio mass spectrometer (VG-Isogas Aqua Sira, VG Isogas, Middlewich, Cheshire, England). TBW was obtained by dividing the measured deuterium dilution space by 1.04. Fat free mass (FFM) was calculated by dividing TBW by the hydration factor 0.73 [25-27]. Fat mass (FM) was determined as BW-FFM.

Leptin

Serum leptin concentrations were measured with a double-antibody, sandwich-type enzyme-linked immunosorbent assay that used a monoclonal antibody specific for human leptin. The lower limit of detection is 0.5 µg/L and the upper limit is 50 µg/L. The intra- and inter assay CVs were 9% and 12%, respectively. The leptin concentrations of normal-weight subjects range from 2 to 12 µg/L. Serum leptin concentrations were also measured at age 7, 12 and 13y.

Attitude towards eating

Eating behavior was analyzed using a validated Dutch translation of the TFEQ [15], which was translated at some questions into an easier, understandable Dutch language for children. The TFEQ consists of 3 factors measuring a person's attitude towards eating. Dietary restraint reflects the extent to which individuals attempt to cognitively control their food intake [15]. Inhibition of restraint reflects individual differences in the extent to which release from the cognitive suppression of eating occurs in response to the presence of palatable food or other disinhibiting stimuli, such as emotional distress [15]. The third factor refers to the subjective feeling of hunger [15].

Physical activity

Physical activity was estimated with the Baecke Questionnaire. This questionnaire consists of three components: work activity, sports activity and leisure activity [28] and has been validated using doubly labelled water [18, 29]. For the children the work index was replaced by a school index with exactly the same questions.

Determination of the genotypes

The genomic DNAs of the 94 subjects were isolated from peripheral blood leukocytes using a QIAamp kit (Qiagen, Hilden, Germany). Followed by genotyping of the PPAR γ 2, BCL1 and CNTF polymorphisms, which has been previously described by Vogels 2006 et al [6].

Parental characteristics

Both the parents reported actual BW measured at home according to our standard instructions (as previously described). Height was copied from their passports, originally measured using a wall-mounted stadiometer. BMI was calculated by BW/height² (kg/m²). In addition, attitude towards eating was measured with the TFEQ.

Statistical analysis

Student t-tests and ANOVA-tests (for the continuous variables), and chi-square tests (for the nominal variables) were executed to determine differences in single variables between groups (i.e. lean vs. overweight) and between polymorphisms. The ANOVA tests were followed by Scheffe's test or a Fishers PLSD test. All tests were two-sided and differences were considered significant at P<0.05. Relationships between dependent and independent variables were tested using simple linear regression and multiple linear regression models. Values are expressed as mean \pm standard deviation (s.d.).

Results

In our cohort 11% of the children were overweight, which was determined using two measures, BMI and body fat

percentage. BMI strongly related to body fat percentage ($R^2 = 0.60$, $p < 0.001$), and also leptin concentrations were related to body fat percentage ($R^2 = 0.825$, $p < 0.001$), which has been shown previously [6, 30].

The characteristics of the boys ($n = 55$) and girls ($n = 39$), divided by overweight status at age 12 and 13y are presented in **table 1**. At age 12 and 13y overweight boys and girls showed significantly higher BW, BMI, waist circumference, body fat percentage, fat mass and leptin levels when compared to lean children. At age 12 and 13y no significant differences were shown in height. At age 12y no significant difference was shown in fat free mass, however, at age 13y overweight boys showed a significant higher fat free mass compared to lean girls.

With respect to the genetic parameters i.e. analysis of the PPARy2, BclI and CNTF genotypes of the subjects, the overall genotype frequencies were in Hardy-Weinberg equilibrium. The genotype frequency distributions of the PPARy2, BclI and CNTF genes (**table 2**) at age 12 and 13y were not significantly different for the overweight children compared to the lean children.

The behavioral characteristics (TFEQ and Baecke scores) of the boys and girls, divided by overweight status at age 12

and 13y are presented in **table 3**. At age 12y no differences were found in restraint score between the four groups, however, at age 13 lean boys showed significant lower restraint scores in comparison to other three groups. At age 12 and 13y the disinhibition scores were significantly higher in overweight boys compared to the other three groups. Overall, no differences between groups were found for feeling of hunger scores and Baecke scores.

The characteristics of the parents of the lean and overweight children are presented in **table 4**. The fathers of overweight children have a higher BW, BMI and restraint score compared to the fathers of lean children. The mothers of the overweight children have a higher disinhibition score compared to the mothers of lean children. A simple linear regression was used to test the association between BMI and restraint or disinhibition scores of the children, fathers or mothers. A significant correlation was found between the restraint score and the BMI of the children ($R^2 = 0.131$, $p < 0.0001$). Also, significant associations were found in the fathers and mothers between the disinhibition score and their BMI ($R^2 = 0.252$, $p < 0.001$) ($R^2 = 0.168$, $p < 0.001$). No correlations were found between the TFEQ scores of the parents and the BMI of the children.

Table 1: characteristics of the boys ($n = 55$) and girls ($n = 39$), divided by overweight status¹ at age 12 and 13y

12 year	Total (n=94)	Lean girls (n = 33)	Overweight girls (n = 6)	Lean boys (n = 51)	Overweight boys (n = 4)	P-value ²
BW (kg) ³	48.4±10.3	47.3±8.2 ^{ab}	61.4±11.2 ^{ac}	46.6±10.1 ^{cd}	60.7±6.4 ^{bd}	0.01
Height (cm)	159.2±10.2	158.7±9.7	160.7±10.3	159.2±10.6	161.6±12.5	0.94
BMI (kg/m ²) ³	18.9±2.6	18.6±1.8 ^a	23.7±2.7 ^{a,b}	18.3±2.2 ^b	23.3±2.1 ^b	0.01
Waist (cm)	66.4±6.2	64.4±5.1 ^{ab}	73.5±5.0 ^{ac}	65.5±5.7 ^{cd}	74.8±3.6 ^{bd}	0.01
Body fat (%)	19.6±7.6	21.7±5.2 ^{ab}	29.7±6.1 ^{ac}	16.6±6.6 ^{cd}	29.1±10.5 ^{bd}	0.01
Fat mass (kg)	9.69±4.9	10.5±3.4 ^{ab}	18.1±5.3 ^{ab}	7.8±3.9 ^a	17.4±5.8 ^b	0.01
Fat free mass (kg)	38.4±7.7	37.4±6.3	43.1±8.1	38.5±8.0	43.3±9.4	0.26
13 Year						0.26
BW (kg) ³	53.1±10.6	50.9±8.3 ^{ab}	66.1±9.8 ^{ac}	51.8±10 ^{cd}	69.0±8.7 ^{ab}	0.01
Height (cm)	164.9±10.1	163.1±8.6	163.7±8.4	165.8±10.9	168.9±14.4	0.54
BMI (kg/m ²) ³	19.5±2.9	19.0±1.9 ^a	26.2±2.8 ^{a,b}	18.7±2.1 ^b	24.1±0.9 ^b	0.01
Waist (cm)	68.0±6.2	65.6±5 ^{ab}	76.2±5.8 ^{ac}	67.6±5.2 ^{cd}	80.7±2.9 ^{bd}	0.01
Body fat (%)	19.8±7.5	21.8±4.4 ^{ab}	31.3±6.9 ^{ac}	15.9±5.9 ^{cd}	27.9±10 ^{bd}	0.01
Fat mass (kg)	10.6±5.2	11.4±3.7 ^{ab}	20.6±6.6 ^{ac}	8.3±3.5 ^{cd}	18.6±5.3 ^{bd}	0.01
Fat free mass (kg)	42.1±8.6	40±4.9 ^a	44.4±6.9	43.8±8.6	50.4±13.5 ^a	0.05

¹ Overweight as defined by age and sex specific cut off points described by Cole et al.

² Differences between lean vs. overweight boys and girls using ANOVA

³ BW, body weight; BMI, body mass index

^a $p < 0.05$ (2-factor ANOVA with group x group interaction); ^b $p < 0.05$ (2-factor ANOVA with group x group interaction)

^c $p < 0.05$ (2-factor ANOVA with group x group interaction); ^d $p < 0.05$ (2-factor ANOVA with group x group interaction)

Table 2: genotype frequency distributions of the PPAR γ 2, GRL and CNTF genes in lean and overweight¹ children at age 12 and 13y

12 and 13 year	Total (n =92)	Lean (n=84)	Overweight (n=10)	P-value ²
PPAR γ 2 ³ (PP/PA/AA)	75/24/1 %	77/22/1 %	60/40/0 %	0.43
GRL ³ (CC/CG/GG)	49/36/15 %	47/37/16 %	10/30/60 %	0.74
CNTF ³ (GG/GA/AA)	78/19/2 %	79/18/ 2%	70/30/0 %	0.96

¹ Overweight as defined by age and sex specific cut off points described by Cole et al.² Differences between lean vs. overweight children using chi-square test³ PPAR γ 2, peroxisome proliferator-activated receptor γ 2; GRL, glucocorticoid receptor; CNTF, ciliary neurotrophic factor**Table 3:** behavioral characteristics of the boys and girls, divided overweight status¹ at age 12 and 13y

12 year	Total (n =94)	Lean girls (n = 33)	Overweight girls (n = 6)	Lean boys (n = 51)	Overweight boys (n=4)	P-value ²
Factor 1 (TFEQ) ³	5.1 \pm 3.5	4.8 \pm 3.5	7.2 \pm 3.1	4.9 \pm 3.2	7.5 \pm 5.9	0.22
Factor 2 (TFEQ) ³	2.7 \pm 1.3	2.6 \pm 1.4 ^a	3.3 \pm 1.5	2.6 \pm 1.1 ^b	4.3 \pm 0.9 ^{ab}	0.04
Factor 3 (TFEQ) ³	3.0 \pm 2.4	2.7 \pm 2.5	3.0 \pm 3.3	5.3 \pm 3.8	5.3 \pm 3.8	0.23
Physical activity (Baecke)	8.2 \pm 1.0	8.3 \pm 0.8	7.6 \pm 1.1	8.3 \pm 0.9	8.2 \pm 2.2	0.38
13 Year						
Factor 1 (TFEQ) ³	5.7 \pm 3.2	6.7 \pm 3.1 ^a	8.3 \pm 4.1 ^b	4.7 \pm 2.6 ^{bcd}	9.5 \pm 4.2 ^c	0.01
Factor 2 (TFEQ) ³	2.4 \pm 1.4	2.5 \pm 1.5 ^a	2.7 \pm 1.3 ^b	2.2 \pm 1.0 ^c	5.0 \pm 2.7 ^{abc}	0.01
Factor 3 (TFEQ) ³	3.1 \pm 2.3	2.8 \pm 2.1	2.5 \pm 0.8	3.5 \pm 2.2	4.5 \pm 5.8	0.35
Physical activity (Baecke)	7.9 \pm 0.8	8.1 \pm 0.9	7.9 \pm 0.5	7.9 \pm 0.8	8.1 \pm 0.9	0.96

¹ Overweight as defined by age and sex specific cut off points described by Cole et al.² Differences between lean vs. overweight children using ANOVA³ TFEQ, three-factor eating questionnaire; Factor 1, dietary restraint eating behavior; Factor 2, inhibition of restraint; Factor 3, general feelings of hunger^a p < 0.05 (2-factor ANOVA with group x group interaction); ^b p < 0.05 (2-factor ANOVA with group x group interaction)^c p < 0.05 (2-factor ANOVA with group x group interaction); ^d p < 0.05 (2-factor ANOVA with group x group interaction)**Table 4:** Parental characteristics of lean and overweight children

	All children		Lean children		Overweight children	
	Father	Mother	Father	Mother	Father	Mother
BW (kg) ²	84.9 \pm 15.8	69.3 \pm 13.8	83.7 \pm 13.5	69.2 \pm 14.2	94.5 \pm 27.7 ¹	69.9 \pm 10.7
Height (cm)	179.9 \pm 6.8	166.9 \pm 5.9	180.0 \pm 6.9	166.9 \pm 5.9	179.9 \pm 6.3	167.3 \pm 5.9
BMI (kg/m ²) ²	26.1 \pm 4.4	24.4 \pm 3.9	25.7 \pm 3.6	24.4 \pm 4.0	29.1 \pm 8.3 ¹	25.0 \pm 3.7
Factor 1 (TFEQ) ²	4.8 \pm 3.2	7.3 \pm 3.7	4.6 \pm 3.1	7.4 \pm 3.8	5.8 \pm 3.7 ¹	6.4 \pm 2.9
Factor 2 (TFEQ) ²	2.7 \pm 1.5	3.4 \pm 2.2	2.8 \pm 1.6	3.4 \pm 2.1	2.3 \pm 0.8	3.5 \pm 2.8 ¹
Factor 3 (TFEQ) ²	2.5 \pm 2.4	2.6 \pm 2.2	2.6 \pm 2.5	2.6 \pm 2.0	2.3 \pm 1.8	2.4 \pm 3.5

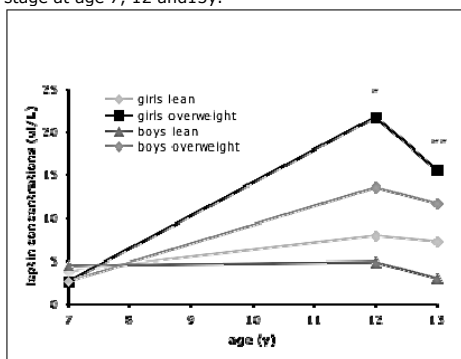
¹ Significant differences in characteristic between the father/mother of overweight children, compared to the father/mother of lean children (p < 0.05)² BW, body weight; BMI, body mass index; TFEQ, three-factor eating questionnaire; Factor 1, dietary restraint eating behavior; Factor 2, inhibition of restraint; Factor 3, general feelings of hunger**Table 5:** Multiple linear regression analyses with body weight (kg) as the dependent variable at age 12 and 13y (n =94)

12 year	Partial β	Std. Error	P-value
(Constant)	39.534	6.376	0.0001
Leptin concentrations (μ L)	0.450	0.170	0.0098
Factor 1 child (TFEQ) ¹	0.282	0.325	0.3883
Factor 2 child (TFEQ) ¹	1.142	0.800	0.1574
BMI father (kg/m ²) ¹	-0.036	0.239	0.8790
Factor 2 mother (TFEQ) ¹	0.692	0.460	0.1363
13 Year			
(Constant)	43.559	6.687	0.0001
Leptin concentrations (μ L)	0.514	0.226	0.0253
Factor 1 child (TFEQ) ¹	-0.132	0.381	0.7305
Factor 2 child (TFEQ) ¹	1.811	0.767	0.0206
BMI father (kg/m ²) ¹	0.027	0.249	0.9147
Factor 2 mother (TFEQ) ¹	0.748	0.472	0.1165

¹ TFEQ, three-factor eating questionnaire; Factor 1, dietary restraint eating behavior; Factor 2, inhibition of restraint; BMI, body mass index. Model 12 year, R² = 0.187; Model 13 year, R² = 0.181

The mean leptin concentrations (μL) of the boys and girls, divided by overweight status at age 7, 12 and 13y are presented in **figure 1**.

Figure 1: Mean of plasma leptin concentrations of the boys ($n = 55$) and girls ($n = 39$), divided by overweight stage at age 7, 12 and 13y.



* overall group \times group interaction at age 12, was significant ($p < 0.05$)

** overall group \times group interaction at age 13, was significant ($p < 0.05$)

At age 7 no significant differences in leptin levels were shown between the four groups. At age 12 and 13y leptin levels were significantly different between all four groups ($p < 0.001$), except between the overweight boys and girls at age 13. The increase in leptin concentrations between 7 to 12y was significant in lean girls ($4.3 \pm 3.9 \mu\text{L}$), overweight girls ($19.2 \pm 8.6 \mu\text{L}$), overweight boys ($10.9 \pm 6.2 \mu\text{L}$) ($p < 0.001$), except in the lean boys ($0.5 \pm 6.4 \mu\text{L}$). The decrease in leptin concentrations from 12 to 13y was significant in the lean boys ($-2.1 \pm 3.3 \mu\text{L}$) ($p < 0.01$), but not in lean girls ($-0.6 \pm 2.9 \mu\text{L}$), overweight girls ($-6.3 \pm 6.1 \mu\text{L}$) and overweight boys ($-1.8 \pm 3.6 \mu\text{L}$). The differences between the increase from 7 to 12y and the decrease from 12 to 13y were significant ($p < 0.01$) in all four groups, suggesting a peak value of leptin at 12y in boys and girls.

A simple linear regression was used to test the association between the leptin concentrations (μL) and fat mass (kg) at age 12 and 13y for lean overweight boys and girls. Leptin was associated to fat mass at age 12 and 13y in lean boys ($R^2 = 0.36$, $p < 0.01$ and $R^2 = 0.35$, $p < 0.01$) and in lean girls ($R^2 = 0.31$, $p <$

0.01 and $R^2 = 0.35$, $p < 0.01$). No associations were found in overweight boys and girls.

The multiple linear regression analyses at age 12 and 13y with body weight (kg) as the dependent variable and the behavioral, parental and physiological factors that were related to body weight (leptin concentrations, dietary restraint score of the child, disinhibition score of the child, BMI of the father and disinhibition score of the mother) as the independent variables are presented in **table 5**. From the model at age 12y, we concluded that the five variables together predicted 18,7% of the variance in body weight at age 12y, however only the leptin concentrations ($p < 0.009$) at age 12y contributed significantly to the model. From the model at age 13y, we concluded that the five variables together predicted 18,1% of the variance in body weight at age 13y, however only the leptin concentrations ($p < 0.03$) and disinhibition score ($p < 0.02$) at age 13y contributed significantly to the model.

Discussion

The objective of our study was to investigate whether genetic, behavioral, parental and physiological factors are involved and the extent of involvement in the development of body weight at age 12 and 13y in a Dutch children cohort. We showed that in our cohort, 11% of the children were overweight. In 2006, the overall prevalence of childhood obesity in the Netherlands was 14.3%, which is the still the lowest in Europe. Like in adults [8], overweight in the children coincided with higher BW, waist circumference, BF%, fat mass and leptin levels.

With respect to the genetic influences on the development of body weight, previous studies showed in adults an association between body weight and the genetic polymorphisms PPAR γ 2, BCL1 and CNTF [7, 9, 10], however the genotype frequency distributions of the genes at age 12 and 13y were not significantly different for the overweight children compared to the lean children. Possible

reasons why we did not observe such associations could be the rather small sample size of the cohort, the small percentage of overweight in this cohort, or age-dependent expression of genotype.

With respect to the behavioral factors involved in the development of body weight, previous studies showed that adults with high levels of both restraint and disinhibition are often overweight, since they are 'unsuccessfully dietary restraint', which beholds that their high restraint scores do not help them to control their weight [31]. In our cohort overweight boys and girls showed higher dietary restraint and disinhibition scores, which suggest they are also 'unsuccessfully dietary restraint'. Consequently, since overweight leads to dietary restraint, subsequent disinhibition of restraint leads to overeating, which leads to storing of excessive energy and thereby results in a vicious circle in which the children are caught that will lead to further increases in body weight. Lean girls showed an increase in dietary restraint scores between 12 and 13y, but since they have a normal weight, this might indicate that they are 'successfully dietary restraint, which beholds that their high restraint scores help them to control their weight [6, 31]. No differences were found however, between lean and overweight boys and girls in physical activity measured by the Baecke questionnaire. We have shown previously in a study by Vogels 2007 et al. that this was caused by overestimation of activity by overweight children [18], when daily physical activity measured by tri-axial accelerometers was compared to estimation of physical activity measured by the Baecke questionnaire.

With respect to the parental influences on the development of body weight, previous studies showed that children of overweight parents are more likely to be overweight themselves [6, 19]. In the current study the fathers of overweight children showed higher body weight, BMI and dietary restraint score; having an overweight and restraint father in the

current cohort of children is therefore an important predictor for children to become overweight. In accordance to previous findings [6, 32] the mothers of overweight children showed a higher disinhibition score. The overall high restraint scores of the mothers and the association between the mother's dietary restraint score and the BMI of the child may be a possible explanation for the fact that we found no association between BMI of the mother and the weight status of the child [6]. In general, one could state that the eating behavior of the parents may serve as a source of information for children what cues should trigger eating and how much to eat [32].

With respect to the physiological influences on the development of body weight, literature showed these are more pronounced during puberty. The hormone leptin is thought to be involved in the start of the changes in the endocrine milieu during puberty. Several transversal studies showed [22, 33-37] a rise in leptin levels progressively from 5 to 15y in girls, whereas in boys, levels increased between 5 and 10y and declined thereafter. In the current longitudinal study we described the leptin levels at age 7, 12 and 13y, and showed an increase in leptin levels between 7 and 12y and a decrease, although only significant for the lean boys between 12 and 13y. Further analysis suggested a peak value of leptin at 12y in all children, which has been suggested to be a physiological trigger to the start of the increase in LH and FSH production; the first recognizable hormonal changes of puberty [38].

In the current longitudinal study we also showed that the positive association between leptin concentrations and fat mass did not change in the lean boys between age 12 and 13y, but became stronger in the lean girls. No significant relationship between leptin concentrations and fat mass at age 12 and 13y was found in the overweight boys and girls, because the small percentage of overweight children in this cohort. Several explanations have been suggested for the

divergent correlations between leptin concentrations and fat mass in lean boys and girls during pubertal development. One explanation is the difference in body fat distribution, since in adipocytes taken from subcutaneous abdominal fat produce greater leptin concentrations compared to adipocytes taken from visceral fat. Therefore, increased serum concentrations of leptin in pubertal girls may be due to greater amounts of subcutaneous fat [39]. A second explanation is the rise in androgens levels in boys at the start of puberty. In vitro, when androgens are added to a culture medium of adipocytes, the leptin secretions are reduced by 62%, which suggests that in androgens suppresses leptin concentrations in boys during puberty [34]. A third explanation is that in boys the ratio of fat free mass/fat mass changes and becomes higher during puberty, when compared to girls [40], which is also shown in our cohort.

With respect to the genetic, parental, behavioral and physiological factors only the last three factors seemed to have an effect on the development of body weight at age 12 and 13y. To determine whether these factors were predictors for body weight, we performed multiple linear regression analyses at age 12 and 13y with body weight as the dependent variable and the behavioral, parental and physiological factors that were related to body weight (leptin concentrations, dietary restraint score of the child, disinhibition score of the child, BMI of the father and disinhibition score of the mother) as the independent variables.

From the model at age 12y, we concluded that only the leptin concentrations at age 12y contributed significantly to the model and from the model at age 13y, that only the leptin concentrations and disinhibition scores of the child at age 13y contributed significantly to the model. These results imply that at the age of 12y predominantly the physiological factors were predictors for body weight, and at the age of 13y both the physiological and behavioral factors were predictors for body weight.

The strengths of the present study is that we found associations between several factors and childhood overweight in this relatively small cohort and the accurate measurement method to determine body composition. We conclude from this longitudinal study, that leptin appeared to play an important role in the development of body weight during puberty, in addition to behavioral and parental factors.

Acknowledgements

We want to thank our subjects for their participation in our cohort study. We gratefully thank Loek Wouters and Wendy Sluijsmans for their assistance. FR and NV carried out the study, collected and analyzed the data, and wrote the largest part of the manuscript. FB (supervised by EM) contributed to the practical work. MW and AN supervised FR. Planning, processing the results and writing the manuscript were done under general supervision by MW. The authors had no conflict of interest.

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Chapter 4

Sleep duration and body-weight development during puberty in a Dutch children cohort

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Submitted

Abstract

Background: Short sleep duration is associated with obesity during childhood and adulthood.

Objective: the objective of our study was to investigate the relationship between sleep duration and body mass index (BMI) from Tanner stage 1 to 5 in a Dutch children cohort.

Design: we determined in 96 children, sleep duration, anthropometric measurements, body composition, physical activity (Baecke questionnaire), and leptin concentrations, from age 7 to 16y.

Results: at Tanner stage 1, 2, 3, 4 and 5 sex differences were observed in height, body weight, waist circumference, fat mass per squared meter (FMI), and leptin concentrations per kg fat mass. From Tanner stage 1 to 5 an increase was observed in both boys and girls in height (0.25 ± 0.15 and 0.25 ± 0.15 m), body weight (24 ± 12.5 and 26.4 ± 12.5 kg), BMI (3.5 ± 2.7 and 4.9 ± 3.1 kg/m²), waist circumference (7.6 ± 11.5 and 11.6 ± 7.7 cm), as well as an increase over time in FMI (1.8 ± 2.3 kg/m²) in girls. A decrease was observed in boys and girls in hours sleep per night (-1.2 ± 0.4 and -1.4 ± 0.5 h) and leptin concentrations per kg fat mass (-0.33 ± 0.15 and -0.17 ± 0.2 ng/ml/kg). Inverse relationships were observed between the change in BMI (kg/m²) and the change in hours of sleep per night (h) from Tanner stage 1 to 4 ($r = 0.68$, $p < 0.001$), from Tanner stage 2 to 5 ($r = 0.35$, $p < 0.05$), and from Tanner stage 1 to 5 ($r = 0.33$, $p < 0.05$).

Conclusion: Development of BMI during puberty is inversely related to change in sleep duration.

Introduction

The prevalence of childhood obesity is emerging as a major health problem (1). The increase in childhood obesity has developed over the same time period as the progressive decrease in self-reported sleep duration (2, 3). Several cross-sectional studies reported on a negative relationship between sleep duration and body mass index (BMI) during childhood (4-9), while some even reported a curvilinear relationship between sleep duration and BMI (10, 11). Additionally, short sleep duration in children was linked to insulin resistance, a larger body fat percentage, a larger waist circumference, decreased physical activity and increased television watching (12-16). Sleep duration during childhood is however subjected to developmental changes, in particular during puberty when a significant decrease in sleep duration has been observed (17, 18). Until now only 4 studies have been described that study sleep duration and BMI development longitudinally (1, 19-21), which measured either only children from age 3 to 9y (1, 20, 21), or measured a group of 3 to 12 year old children twice with a 5-year interval not taking pubertal stage into account (19). Pubertal stage may however alter the relationship between sleep duration and BMI as well as other determinants of overweight in children, such as body composition, and waist circumference (22). This however has not been investigated until now. Consequently, the objective of our longitudinal study was to investigate the relationship between sleep duration and BMI from Tanner stage 1 to 5. Therefore, we determined in a Dutch cohort of 96 children, sleep duration, anthropometric measurements, body composition, Baecke questionnaire, and leptin concentrations, from age 7 to 16y.

Subjects and methods

Subjects

Subjects were recruited from a Dutch Caucasian cohort of children born between 1990 and 1993 (23, 24). As infants, these

children and their mothers participated in i) a study of essential fatty acids during pregnancy and pregnancy outcome (23), and ii) a study, performed between 1997 and 2000, about the long-term effects of fetal essential fatty acid availability (25). Anthropometric data were available from these children, no interventions were executed, and follow-up studies were performed with 96 children (24). Exclusion criteria for follow-up measurements were chronic illness and depression. Each child and their parents gave written informed consent to participate in the study, which was approved by the CCMO (Central Committee Human Research) and by the MEC (Medical Ethical Committee) of the Maastricht University.

Study design

Every year the follow-up measurements were performed in the months November and December to exclude seasonal effects. At 4 p.m. after a 3-hour fast children's body weight (BW), height, body mass index (BMI), waist circumference, body composition (24, 26), leptin concentrations, physical activity, as well as sleep duration were determined at ages 7 to 16y.

Measurements

Anthropometry

The children's BW was determined using a digital balance accurate to 0.1 kg (Sauter D7470, Ebingen, Germany) and height was determined using a wall-mounted stadiometer (Seca, model 220, Hamburg, Germany). Measurements were executed in underwear, after an overnight fast and after voiding the bladder. BMI was calculated by $BW/height^2$ (kg/m²) and BMI in childhood changes substantially with age. To define normal, overweight and obesity in children, we used the specific cut off points described by Cole et al. 2000 (27). The waist circumference was measured at the site of the smallest circumference between the rib cage and the ileac crest, with the subjects in standing position. The pubertal stage was documented in all children according to

the classification by Tanner (28). The Tanner stage is defined based on physical measurements of external primary and secondary sex characteristics, namely development of breast and pubic hair in girls or development of genitalia and pubic hair in boys (28). Boys with Tanner stage 1 had LH and FSH levels of 1.7 ± 1.4 and 2.9 ± 2.3 IU/l and boys with Tanner stage 5 had LH and FSH levels of 5.7 ± 2.8 and 3.3 ± 1.7 IU/l. Girls with Tanner stage 1 had LH and FSH levels of 3.0 ± 2.9 and 3.8 ± 2.1 IU/l and girls with Tanner stage 5 had LH and FSH levels of 4.4 ± 4.6 and 5.0 ± 4.4 IU/l. Tanner stage definition was thus supported by plasma LH and FSH measurements. Anthropometric measurements were determined at age 7, 12, 13, 14, 15 and 16y.

Body composition

Body composition was measured using the deuterium dilution technique (D_2O). D_2O dilution was used to measure total body water (TBW). Subjects were asked to collect a urine sample in the evening just before drinking the deuterium-enriched water solution. After ingestion of this solution, no further consumption was allowed. In the morning the first urine sample was discarded, the whole second urine sample (about ten hours after drinking the water solution) was collected. The dilution of the deuterium isotope is a measure of the TBW of the subject. Deuterium was measured in the urine samples with an isotope ratio mass spectrometer (VG-Isogas Aqua Sira, VG Isogas, Middlewich, Cheshire, England). TBW was obtained by dividing the measured deuterium dilution space by 1.04. Fat free mass (FFM) was calculated by dividing TBW by the hydration factor 0.73 as defined by Fomon et al. 1982 (29). Fat mass (FM) was determined as $BW - FFM$. Fat mass index (FMI) was calculated by $\text{fat mass}/\text{height}^2$ (kg/m^2) and fat free mass index (FFMI) was calculated by $\text{fat free mass}/\text{height}^2$ (kg/m^2) (30-32). Body composition was measured at age 12, 13, 14, 15 and 16y.

Leptin

Plasma leptin concentrations were measured with a double-antibody, sandwich-type enzyme-linked immunosorbent assay that used a monoclonal antibody specific for human leptin. The lower limit of detection is $0.5 \mu\text{g}/\text{L}$ and the upper limit is $50 \mu\text{g}/\text{L}$. The intra- and inter assay CVs were 9% and 12%, respectively. The leptin concentrations of normal-weight subject's range from 2 to $12 \mu\text{g}/\text{L}$. Plasma leptin concentrations were measured at age 7, 12, 13, 14, 15 and 16y.

Physical activity

Physical activity was measured with the Baecke questionnaire. Responses to 19 questions were scored on a five-point scale and resulted in three scores: work activity score, sports activity score and leisure activity score, and has been validated using doubly labeled water (33). For the children, the work index was replaced by a school index with similar questions (e.g. 'When I am at school, I walk: never/seldom/sometimes/often/all the time?'). The Baecke questionnaire for children was validated by Vogels et al. 2007 (34). Physical activity was determined at age 12, 13, 14, 15 and 16y.

Sleep duration

Sleep duration was measured using the question: 'how many hours do you usually sleep per night?' which would represent their habitual sleep duration (7, 9, 21, 35). Sleep duration was determined at age 12, 13, 14, 15, and 15y.

Statistical analysis

Differences between boys and girls per Tanner stage were determined using unpaired t-tests. As hours of sleep per night were measured at age 12 to 16y, and not all children were at Tanner stage 1 at age 12y or Tanner stage 5 at age 16y, three groups appeared, namely from one with data from Tanner stage 1 to 4 (17m/7f), one with data from Tanner stage 2 to 5 (23m/19f), and one with data from Tanner stage 1 to 5 (14m/18f).

Relationships between the changes in BMI

and changes in sleep duration during the progressive Tanner stages were assessed using simple linear regression models. Relationships between the changes in BMI and changes in waist circumference and FMI during the progressive Tanner stages were assessed using simple linear regression models. Statistical analyses were performed with Statview SE Graphics™ for Macintosh. All tests were two-sided and differences were considered significant at $P < 0.05$. Values are expressed as mean \pm standard deviation (s.d.).

Results

Data from 57 boys and 39 girls on height, body weight, BMI, waist circumference, fat mass per squared meter (FMI), leptin concentrations per kg fat mass, Baecke sport score, and hours sleep per night were collected from age 7 to 16y. The characteristics of the boys and girls from Tanner stage 1 to 5 are represented in **table 1**. At Tanner stage 1, 2, 3, 4 and 5 sex differences were observed in height, body weight, waist circumference, fat mass per squared meter (FMI), and leptin concentrations per kg fat mass. No differences were found in BMI, Baecke sport score, and hours sleep per night between boys and girls. From Tanner stage 1 to 5 an increase was observed in both boys and girls in respectively height (0.25 ± 0.15 and 0.25 ± 0.15 m), body weight (24 ± 12.5 and 26.4 ± 12.5 kg), BMI (3.5 ± 2.7 and 4.9 ± 3.1 kg/m²), waist circumference (7.6 ± 11.5 and 11.6 ± 7.7 cm), as well as an increase over time in

FMI (1.8 ± 2.3 kg/m²) in girls. A decrease was observed in boys and girls in hours sleep per night (-1.2 ± 0.4 and -1.4 ± 0.5 h) and leptin concentrations per kg fat mass (-0.33 ± 0.15 and -0.17 ± 0.2 ng/ml/kg). No alterations over time were observed in FMI and Baecke sport scores in boys, while in girls no alterations were observed in Baecke sport score.

As hours of sleep per night were measured at age 12 to 16y, and not all children were at Tanner stage 1 at age 12y or Tanner stage 5 at age 16y, three groups appeared, namely from one with data from Tanner stage 1 to 4 (17m/7f), one with data from Tanner stage 2 to 5 (23m/19f), and one with data from Tanner stage 1 to 5 (14m/18f). In all three groups we tested whether changes in BMI over time were related to changes in hours of sleep per night over time. Inverse relationships were observed between the change in BMI (kg/m²) and the change in hours of sleep per night (h) from Tanner stage 1 to 4 ($r = 0.68$, $p < 0.001$), from Tanner stage 2 to 5 ($r = 0.35$, $p < 0.05$), and from Tanner stage 1 to 5 ($r = 0.33$, $p < 0.05$).

In all three groups positive relationships were found between the change in BMI over time and the change in waist circumference over time ($r = 0.79$, $p < 0.0001$; $r = 0.85$, $p < 0.0001$; $r = 0.81$, $p < 0.0001$), and the change in FMI over time ($r = 0.71$, $p < 0.001$; $r = 0.62$, $p < 0.001$; $r = 0.85$, $p < 0.0001$).

Table 1: Characteristics of the boys and girls at Tanner stage 1 to 5

	Tanner stage 1		Tanner stage 2		Tanner stage 3		Tanner stage 4		Tanner stage 5	
	Boys	Girls	Boys	Girls	Boys	Girls	Boys	Girls	Boys	Girls
Number	57	39	53	40	52	40	51	41	35	36
Height (m)	1.51 \pm 0.2	1.43 \pm 0.2 [#]	1.66 \pm 0.1	1.61 \pm 0.1 [#]	1.72 \pm 0.1	1.64 \pm 0.1 [#]	1.74 \pm 0.1	1.67 \pm 0.1 [#]	1.76 \pm 0.1	1.68 \pm 0.1 [#]
Body weight (kg)	42.9 \pm 13	37.2 \pm 14 [#]	54.6 \pm 11	51.5 \pm 10 [#]	60.7 \pm 9	55.6 \pm 10 [#]	65.3 \pm 11	60.2 \pm 10 [#]	66.9 \pm 12	63.6 \pm 11 [#]
BMI (kg/m ²)	18.0 \pm 2.5	17.4 \pm 2.9	19.5 \pm 2.5	19.7 \pm 2.9	20.4 \pm 2.5	20.5 \pm 3.0	21.3 \pm 2.4	21.3 \pm 2.8	21.5 \pm 2.8	22.3 \pm 3.3
Waist circumference (cm)	66.7 \pm 17	60.6 \pm 7.5 [#]	69.7 \pm 5.8	66.4 \pm 5.9 [#]	71.7 \pm 5.5	68.1 \pm 6.5 [#]	74.0 \pm 6.9	70.4 \pm 5.9 [#]	74.3 \pm 6.1	72.2 \pm 7.8 [#]
Fat mass Index (kg/m ²)	3.3 \pm 1.9	4.7 \pm 1.8 [#]	3.6 \pm 1.9	4.6 \pm 1.9 [#]	3.2 \pm 1.6	5.2 \pm 2.1 [#]	3.7 \pm 1.9	5.6 \pm 1.9 [#]	3.7 \pm 1.6	6.5 \pm 2.8 [#]
Leptin/fat mass (ng/ml/kg)	0.50 \pm 0.2	0.73 \pm 0.2 [#]	0.51 \pm 0.4	0.73 \pm 0.3 [#]	0.38 \pm 0.4	0.55 \pm 0.3 [#]	0.26 \pm 0.3	0.39 \pm 0.2 [#]	0.17 \pm 0.1	0.56 \pm 0.2 [#]
Baecke sport score	2.7 \pm 1.2	2.6 \pm 1.6	2.7 \pm 0.9	2.9 \pm 1.1	3.1 \pm 0.6	2.9 \pm 1.1	3.1 \pm 0.6	2.8 \pm 0.6	3.1 \pm 0.7	2.8 \pm 0.7
Hours sleep per night (h)	8.9 \pm 0.2	8.9 \pm 0.3	8.6 \pm 0.6	8.7 \pm 0.6	8.3 \pm 0.7	8.3 \pm 0.8	7.8 \pm 0.9	7.8 \pm 0.8	7.7 \pm 0.7	7.5 \pm 0.8

#Differences between boys and girls per Tanner stage (unpaired t-test) ($p < 0.05$)

Discussion

The objective of our longitudinal study was to investigate the relationship between sleep duration and BMI from Tanner stage 1 to 5. We observed sex differences from Tanner stage 1 onwards in height, body weight, waist circumference, body composition, and leptin concentrations per kg fat mass. This was in concordance with previous literature on developmental changes in anthropometry and leptin concentrations normalized to fat mass during puberty (36, 37).

In concordance with previous longitudinal studies on sleep duration (17-19), a significant decrease in sleep duration during puberty was observed in our cohort. Previous studies however have not corrected for pubertal stage, and we observed that the decrease in sleep duration is related to increasing Tanner stage. These results illustrate the importance for longitudinal and transversal studies on sleep duration to correct for pubertal stage and not for age.

Previous longitudinal studies on sleep duration and BMI in children have observed a consistent negative linear association between baseline habitual sleep duration and later obesity (1, 19-21). None of the longitudinal studies in children have however investigated whether changes in sleep duration are associated with changes in BMI. Our study is therefore the first to show that with progressive Tanner stages BMI increases and sleep duration decreases in an interrelated way. The increase in BMI from Tanner stage 1 to 5 is furthermore related to the increase in waist circumference and FMI, which suggest that the decrease in sleep duration leads to an increase in obesity related characteristics.

From the observed correlations between development of BMI and changes in sleep duration cause and effect cannot be completely disentangled. Nevertheless, a larger increase in BMI following a larger reduction in sleep duration seems to be plausible based upon the longitudinal study from Snell et al. 2002 (19) that showed a relationship between sleep duration measured at the first time point and BMI measured after a 5-year interval, and not vice versa. On the other hand if sleep duration were to be reduced following an increase in BMI, a plausible theory would be sleep apnea. Sleep apnea however is only present in 1-3% of the children in a normal population and strongly related to obesity (38), and is therefore an unlikely explanation in our cohort as only 11% of the children were overweight and with no known cases of sleep apnea.

We therefore suggest that based upon our data, that reduction of sleep duration may contribute to development of overweight during puberty. Taken together, we conclude that development of BMI during puberty is inversely related to change in sleep duration.

Acknowledgements

We want to thank our subjects for their participation in our cohort study. We gratefully thank Loek Wouters and Wendy Sluijsmans for their assistance. FR and SV carried out the study, collected and analyzed the data, and wrote the largest part of the manuscript. MW and AN supervised FR and SV. Planning, processing the results and writing the manuscript were done under general supervision by AN and MW. The authors had no conflict of interest.

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Chapter 5

The relationship between leptin, gonadotropic hormones, and body composition during puberty in a Dutch children cohort

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European Journal of endocrinology 2009 Jun;160(6):973-8

Abstract

Objective: to investigate the relationship between leptin concentrations, gonadotropic hormone concentrations, and body composition during puberty in a Dutch children cohort.

Design: In a cohort of 98 children, we determined anthropometric measurements, body composition, and concentrations of leptin, FSH, and LH.

Results: sex differences were observed from Tanner stage 1 onwards in weight, body fat percentage, and leptin/fat mass ratio. In boys and girls the relationship between leptin concentrations and fat mass was weaker at Tanner stage 2 ($R^2=0.33$ and $R^2=0.39$, $p<0.001$), 3 ($R^2=0.27$ and $R^2=0.36$, $p<0.002$), and 4 ($R^2=0.21$ and $R^2=0.28$, $p<0.03$) than at Tanner stage 1 ($R^2=0.51$ and $R^2=0.67$, $p<0.001$) and 5 ($R^2=0.46$ and $R^2=0.78$, $p<0.01$). In girls, a peak in leptin concentrations (8.5 ± 6.0 ng/ml) preceded a peak in LH and FSH concentrations (15.1 ± 3.5 and 5.0 ± 4.5 IU/l). A lead/lag relationship was observed of leptin at Tanner stage 1 to LH and FSH at Tanner stage 2 ($R^2=0.12$, $p<0.05$ and $R^2=0.18$, $p<0.05$). In boys, there was no peak in leptin, LH, and FSH and additionally, leptin at Tanner stage 3 was related FSH at Tanner stage 4 ($R^2=0.17$, $p<0.04$).

Conclusion: in boys and girls during puberty factors independent of fat mass become (transiently) more important in the regulation of plasma leptin concentrations. Moreover, in girls leptin is suggested to act as a permissive factor for the onset of puberty, while in boys leptin has a different timing and possibly different function.

Introduction

The achievement of a critical body weight or fat mass is thought to play a prominent role in the start of sexual maturation [1, 2], since underfeeding and malnutrition in humans, as seen in anorexia, is related to a delay in the onset puberty [1-3]. Leptin has been proposed as a physiological link between adiposity status and the start of sexual maturation [1-5]. Several cross-sectional studies and three small longitudinal studies support this theory since they have found that leptin concentrations rose prepubertally, but then decreased at the initiation of puberty, as gonadal hormone concentrations began to increase [1, 4, 6-11].

The proposed mechanism behind leptin as a trigger to puberty, beholds that leptin independently of fat mass acts on the hypothalamic luteinizing hormone-releasing hormone (LHRH) pulse generator [12-14]. Consequently, LHRH stimulates the release of luteinizing hormone (LH) and follicle-stimulating hormone (FSH) from the pituitary, which in turn stimulate the gonads to release testosterone and estradiol [12-14]. Testosterone and estradiol inhibit the secretion of LHRH and respectively LH and FSH, which thereby form a negative feedback loop [13, 14]. Moreover, testosterone inhibits leptin secretion from the adipocytes, which forms a second negative feedback loop [14-16]. A positive feedback loop is formed by estradiol, which stimulates leptin secretion from the adipocytes [14, 16]. This proposed mechanism suggests that increased gonadotrophic and gonadal hormones concentrations may alter the relationship between leptin concentrations and body fat during puberty.

Most studies, however, that investigated the role of leptin as a trigger to puberty, did not determine body composition [6], or were done in transversal cohorts [1, 6, 8, 9] and relatively small longitudinal cohorts [4, 10, 11]. Consequently, previous studies have been unable to test the hypothesis that leptin and

gonadotropic hormones are temporarily related during puberty, and whether consequently the relationship between leptin and body composition is temporarily altered. Therefore, the objective of our longitudinal study was to investigate the relationship between leptin concentrations, gonadotropic hormone concentrations, and body composition during puberty in a Dutch children cohort. Therefore, we determined in 98 children anthropometric measurements, body composition, and concentrations of leptin, FSH, LH, testosterone and estradiol.

Subjects and methods

Subjects

Subjects were recruited from a Dutch Caucasian cohort of children born between 1990 and 1993 [17, 18]. As infants, these children and their mothers participated in i) a study of essential fatty acids during pregnancy and pregnancy outcome [17], and ii) a study, performed between 1997 and 2000, about the long-term effects of fetal essential fatty acid availability [19]. Anthropometric data were available from these children and no interventions were implemented. To evaluate the development of obesity and related parameters, follow-up studies were performed with 98 children, and the dropout rate was 0% [18]. Each child and one of his or her parents gave written informed consent to participate in the study, which was approved by the Central Committee Human Research and by the Medical Ethical Committee of the Maastricht University.

Study design

At 4 p.m. after a 3-hour fast children's age, body weight (BW), height, body mass index (BMI), body composition [7, 18], leptin concentrations, follicle-stimulating hormone (FSH) concentrations, luteinizing hormone (LH) concentrations, testosterone concentrations and estradiol concentrations were determined.

Measurements

Anthropometry

The children's BW was determined using a digital balance accurate to 0.1 kg (Sauter D7470, Ebingen, Germany) and height was determined using a wall-mounted stadiometer (Seca, model 220, Hamburg, Germany). Measurements were executed in underwear and after voiding the bladder. BMI in childhood changes substantially with age and was calculated by $BW/height^2$ (kg/m^2). Therefore to define normal, overweight and obesity in children, we used the specific cut off points described by Cole et al. 2000 [20]. For example, at age 12y a BMI of 21.22 and 21.68 kg/m^2 in boys and girls is defined as overweight, and a BMI of 26.02 and 26.67 kg/m^2 as obese. The pubertal stage was documented in all children according to the classification by Tanner [21]. The Tanner stage is defined based on physical measurements of external primary and secondary sex characteristics, namely development of breast and pubic hair in girls or development of genitalia and pubic hair in boys [21]. At age 15y, boys in Tanner stage 3 had testosterone levels of 8.3 ± 5.1 ng/l and boys with Tanner stage 5 had testosterone levels of 16.5 ± 5.4 ng/l. At age 15, girls in Tanner stage 3 had estradiol levels of 14.8 ± 15.8 ng/l and girls with Tanner stage 5 had estradiol levels of 32.7 ± 32.4 ng/l. Tanner stage definition was thus supported by plasma testosterone and estradiol measurements (ng/l) at age 15. Anthropometric measurements were determined at ages 7, 12, 13, 14 and 15y.

Body composition

Body composition was measured using the deuterium dilution technique (D_2O). D_2O dilution was used to measure total body water (TBW). Subjects were asked to collect a urine sample in the evening just before drinking the deuterium-enriched water solution. After ingestion of this solution, no further consumption was allowed. Ten hours after drinking the water solution, another urine sample was collected. The dilution of the deuterium isotope is a measure of the TBW of the

subject. Deuterium was measured in the urine samples with an isotope ratio mass spectrometer (VG-Isogas Aqua Sira, VG Isogas, Middlewich, Cheshire, England). TBW was obtained by dividing the measured deuterium dilution space by 1.04. Fat free mass (FFM) was calculated by dividing TBW by the hydration factor 0.73 [22-24]. Fat mass (FM) was determined as $BW-FFM$. Body composition was measured at ages 12,13,14 and 15y.

Leptin

At 4 p.m. after a 3-hour fast blood was drawn for determination of leptin concentrations. Plasma leptin concentrations were measured with a double-antibody, sandwich-type enzyme-linked immunosorbent assay that used a monoclonal antibody specific for human leptin. The lower limit of detection is 0.5 $\mu g/L$ and the upper limit is 50 $\mu g/L$. The intra- and inter assay CVs were 9% and 12%, respectively. The leptin concentrations of normal-weight subjects' range from 2 to 12 $\mu g/L$. Plasma leptin concentrations (ng/ml) were measured at ages 7, 12, 13, 14 and 15y.

Follicle-stimulating hormone (FSH)

At 4 p.m. after a 3-hour fast blood was drawn for determination of FSH concentrations. Plasma FSH concentrations were measured with a double-antibody, sandwich-type enzyme-linked immunosorbent assay that used a monoclonal antibody specific for human FSH. The lower limit of detection is 0.100 IU /L and the upper limit is 200.0 IU/L. The intra- and inter assay CVs were 1.8% and 5.1%, respectively. The FSH concentrations range from 1.5 to 12.4 IU /L in boys and from 1.7 to 21.5 IU /L in girls. Plasma FSH concentrations (IU /L) were measured at ages 12, 13, 14 and 15y.

Luteinizing hormone (LH)

At 4 p.m. after a 3-hour fast blood was drawn for determination of LH concentrations. Plasma LH concentrations were measured with a double-antibody, sandwich-type enzyme-linked immunosorbent assay that used a

monoclonal antibody specific for human LH. The lower limit of detection is 0.100 IU /L and the upper limit is 200 IU /L. The intra- and inter assay CVs were 0.8% and 2.0%, respectively. The LH concentrations range from 1.7 to 8.6 IU /L in boys and from 1.0 to 95.6 IU /L in girls. Plasma LH concentrations (IU /L) were measured at ages 12, 13, 14 and 15y.

Statistical analysis

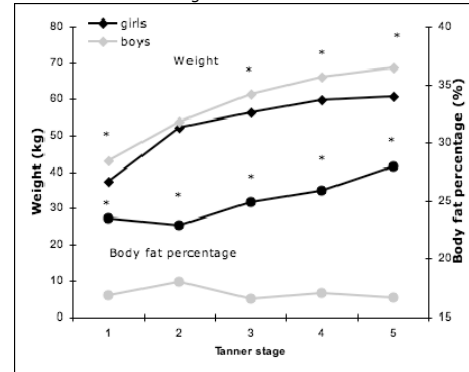
Differences between two groups were determined using unpaired t-tests. Differences over time and between conditions were determined using two-factor ANOVA with repeated measures. Relationships between dependent and independent variables were determined using simple linear regression models. To investigate a time-dependent relationship in the sequence of the hormonal changes during puberty, we used the lead/lag technique, which correlates hormones secreted in one stage, with those secreted in the next stage [11]. All tests were two-sided and differences were considered significant at $P < 0.05$. Values are expressed as mean \pm standard deviation (s.d.).

Results

At Tanner stage 1 we collected data from 95 (57 m/38f) children, at Tanner stage 2 from 95 children (54 m/41f), at Tanner stage 3 from 95 children (55 m/40f), at Tanner stage 4 from 63 children (35m/28f), as well as from 33 children (14m/19f) at Tanner stage 5. 11% of the children were classified as overweight using specific cut off points described by Cole et al. 2000 [20]. In our cohort annual height velocity in boys and girls was 5.7 ± 1.3 and 5.7 ± 1.2 cm/y at Tanner stage 1; 6.5 ± 1.8 and 5.7 ± 1.2 cm/y at Tanner stage 2; 6.9 ± 2.6 and 5.7 ± 2.4 cm/y at Tanner stage 3; 6.1 ± 3.0 and 3.1 ± 2.3 cm/y at Tanner stage 4; and 3.7 ± 3.1 and 1.5 ± 1.4 cm/y Tanner stage 5. **Figure 1** shows sex differences (boys vs. girls) in weight at Tanner stage 1, 3, 4, and 5 (43.1 ± 12.5 vs. 36.9 ± 14.2 kg, $p < 0.02$; 61.5 ± 10.5 vs. 56.2 ± 9.7 kg, $p < 0.01$; 66.1 ± 9.5

vs. 59.9 ± 9.7 kg, $p < 0.01$; 68.5 ± 9.2 vs. 60.7 ± 8.8 kg, $p < 0.02$). Differences were absent at Tanner stage 2, caused by a larger increase in weight in girls when compared to boys between Tanner stage 1 to 2 (15.6 ± 12.4 vs. 12.4 ± 10.4 kg, $p < 0.18$), and a significant larger increase in weight in boys when compared to girls between Tanner stage 2 to 3 (6.6 ± 3.2 vs. 3.9 ± 2.2 kg, $p < 0.001$). Additionally, sex differences in body fat percentage were shown at all Tanner stages (16.9 ± 7.2 vs. $23.8 \pm 8.1\%$, $p < 0.05$; 18.1 ± 7.9 vs. $22.9 \pm 5.9\%$, $p < 0.002$; 16.6 ± 6.7 vs. $24.9 \pm 6.3\%$, $p < 0.001$; 17.1 ± 6.9 vs. $25.9 \pm 5.7\%$, $p < 0.001$; 16.7 ± 5.9 vs. $27.9 \pm 6.3\%$, $p < 0.001$). Because of sex differences, the results on body composition, leptin concentrations, and gonadotropic hormone concentrations were analyzed separately for boys and girls.

Figure 1: Mean of weight (kg) and body fat percentage (%) in the boys (n = 57) and girls (n = 41) as a function of Tanner stage.

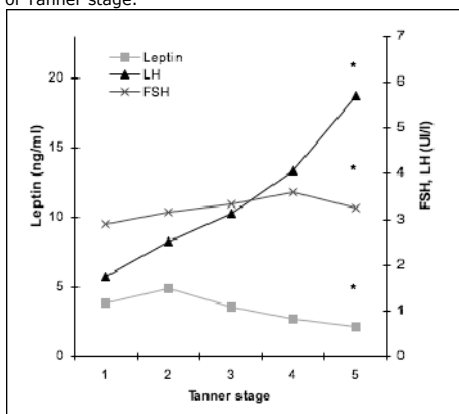


* $p < 0.05$ for differences between boys and girls (ANOVA repeated measures)

Figure 2a shows the mean of plasma leptin concentrations (ng/ml), LH concentrations (IU/l) and FSH concentrations (IU/l) in the boys (n = 57) as a function of Tanner stage. In lean and overweight boys no differences were found in the leptin and gonadotropic hormone patterns. The leptin concentrations increased from Tanner stage 1 to 2 (3.9 ± 2.3 to 4.9 ± 5.1 ng/ml, $p < 0.5$), and decreased significantly from Tanner stage 2 to 5 (4.9 ± 5.1 to 2.1 ± 1.9 ng/ml, $p < 0.01$). FSH

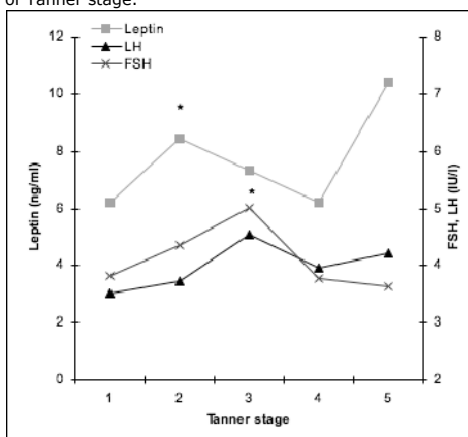
concentrations increased from Tanner stage 1 to 4 (2.9 ± 2.3 to 3.6 ± 1.9 IU/l, $p < 0.09$), and LH concentrations significantly increased from Tanner stage 1 to 5 (1.8 ± 1.4 to 5.7 ± 2.8 IU/l, $p < 0.01$).

Figure 2a: Mean of plasma leptin concentrations (ng/ml), LH concentrations (IU/l), and FSH concentrations (IU/l) in the boys ($n = 57$) as a function of Tanner stage.



* $p < 0.05$ for differences in time in the leptin and LH concentrations (ANOVA repeated measures)

Figure 2b: Mean of plasma leptin concentrations (ng/ml), LH concentrations (IU/l), and FSH concentrations (IU/l) in the girls ($n = 41$) as a function of Tanner stage.



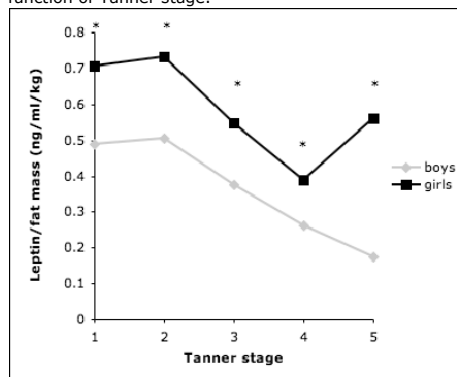
* $p < 0.05$ for differences in time in the leptin and LH concentrations (ANOVA repeated measures)

Figure 2b shows the mean of plasma leptin concentrations (ng/ml), LH concentrations (IU/l) and FSH concentrations (IU/l) in girls ($n = 41$) as a function of Tanner stage. In lean and overweight girls no differences were found in the leptin and gonadotropic hormone

patterns. The leptin concentrations increased significantly from Tanner stage 1 to 2 (6.2 ± 4.9 to 8.5 ± 6.0 ng/ml, $p < 0.02$), decreased significantly from Tanner stage 2 to 4 (8.5 ± 6.0 to 6.2 ± 3.3 ng/ml, $p < 0.04$), and significantly increased again from Tanner stage 4 to 5 (6.2 ± 3.3 vs. 10.4 ± 6.6 ng/ml $p < 0.01$). The peak in leptin concentrations was followed by an increase LH and FSH concentrations from Tanner stage 2 to 3 (3.4 ± 2.3 to 5.1 ± 3.5 IU/l, $p < 0.05$ and 5.1 ± 3.5 to 3.9 ± 4.3 IU/l, $p < 0.14$), and a decrease in LH and FSH concentrations from Tanner stage 3 to 4 (4.4 ± 5.0 to 5.0 ± 4.5 IU/l, $p < 0.24$ and 5.0 ± 4.5 to 3.8 ± 2.1 IU/l, $p < 0.33$).

Figure 3 shows the mean leptin concentrations per kilogram of fat mass (FM) of the boys and girls as a function of Tanner stage. Sex differences in leptin/fat mass ratio were observed at Tanner stage 1, 2, 3, 4, and 5 (0.49 ± 0.2 vs. 0.70 ± 0.2 ng/ml/kg, $p < 0.009$; 0.51 ± 0.4 vs. 0.73 ± 0.3 ng/ml/kg, $p < 0.005$; 0.38 ± 0.4 vs. 0.55 ± 0.3 ng/ml/kg, $p < 0.04$; 0.26 ± 0.3 vs. 0.39 ± 0.2 ng/ml/kg, $p < 0.04$; 0.17 ± 0.1 vs. 0.56 ± 0.2 ng/ml/kg, $p < 0.001$).

Figure 3: Mean leptin concentrations (ng/ml) per kilogram of fat mass (FM) of the boys and girls as a function of Tanner stage.



* $p < 0.05$ for differences between boys and girls (ANOVA repeated measures)

Table 1 shows regression analyses between plasma leptin concentrations (ng/ml) as the dependent variable and fat mass (kg) as the independent variable in boys ($n = 57$) and girls ($n = 41$) at

Tanner stage 1-5. In both boys and girls a positive relationship was observed between leptin concentrations and fat mass from Tanner stage 1 onwards. In the girls only at Tanner stage 2, 3, and 4 a smaller proportion of the variance in the leptin concentrations was accounted for by using fat mass as the independent variable.

Table 1: Regression between plasma leptin concentrations (ng/ml) as the dependent variable and fat mass (kg) as the independent variable in boys (n = 57) and girls (n = 41) at Tanner stage 1-5.

Boys		
Tanner stage	R ²	P value
1	0.51	0.001
2	0.33	0.001
3	0.27	0.001
4	0.21	0.01
5	0.46	0.01
Girls		
Tanner stage	R ²	P value
1	0.67	0.001
2	0.39	0.001
3	0.36	0.002
4	0.28	0.03
5	0.78	0.001

To test whether leptin can be considered as the 'lead' hormone in relation to LH, and FSH or vice versa, in boys and girls leptin concentrations in a certain Tanner stage were paired with the other hormones in the following Tanner stage and vice versa, which results in 32 sets of paired data per child. In boys at Tanner stage 1 LH and FSH concentrations were related to Tanner stage 2 leptin concentrations ($R^2 = 0.16$, $p < 0.02$ and $R^2 = 0.13$, $p < 0.03$), in turn Tanner stage 3 leptin concentrations were related to Tanner stage 4 FSH concentrations ($R^2 = 0.17$, $p < 0.04$). In girls at Tanner stage 1 leptin concentrations were related to Tanner stage 2 LH and FSH concentrations ($R^2 = 0.12$, $p < 0.05$ and $R^2 = 0.18$, $p < 0.05$).

Discussion

The objective of our longitudinal study was to investigate the relationship between leptin concentrations, gonadotropic hormone concentrations, and body composition during puberty in a Dutch children cohort. We observed sex differences from Tanner stage 1 onwards in weight, body composition, and leptin

concentrations per kg fat mass. This was in concordance with previous literature on developmental changes in anthropometry and leptin concentrations normalized to fat mass during puberty [8, 25].

Sex differences were also observed in the development of leptin, LH, and FSH concentrations over time. In girls a peak in leptin concentrations was observed at Tanner stage 2, followed by a peak in LH and FSH concentrations at Tanner stage 3, thereby confirming results from previous cross-sectional and small longitudinal studies [1, 4, 6-11]. In boys no peak in leptin was observed at Tanner stage 2, as leptin decreased from Tanner stage 2 onwards and LH and FSH concentrations increased from Tanner stage 1 to 4, thereby confirming results from previous small longitudinal studies [4, 26]. Transversal studies, however, did show a leptin peak in boys during puberty [1, 6], which may be explained by methodological differences as well as presentation of the data by age group instead of by pubertal stage.

The observations of our longitudinal study shed more light on the temporal alterations in the relationship between leptin and body composition [8]. Both in boys and girls, the leptin/fat mass ratio decreased from Tanner stage 2 onwards. In boys, this decrease continued throughout puberty, while in girls this ratio increased again at Tanner stage 5, which confirms results by Horlick et al [8]. These results imply that during puberty factors independent of fat mass become (transiently) more important in the regulation of plasma leptin concentrations.

To investigate a time-dependent relationship in the sequence of the hormonal changes during puberty, we used the lead/lag technique, which correlates hormones secreted in one stage, with those secreted in the next stage [11]. In girls we observed that prepubertal leptin concentrations (Tanner stage 1), when acting as a lead, related to early pubertal LH and FSH concentrations (Tanner stage 2), as observed in both

boys and girls by Masqood et al [11]. These results imply a temporal relationship between leptin and gonadotropic hormones during early puberty.

In boys, opposing results were found: prepubertal LH and FSH concentrations (Tanner stage 1) were related to early pubertal leptin concentrations (Tanner stage 2), as observed in boys and girls by Masqood et al [11]. They however observed prepubertal leptin concentrations to be related to early pubertal gonadotropic hormones and vice versa in both boys and girls [11], while we observed the first relationship in girls and the second one in boys. The discrepancy between the two studies is presumably caused by the difference in study population, because we analyzed boys and girls separately, each group consisting of about 40 children, and Masqood et al. [11] analyzed only 13 boys and 7 girls, 20 children together as one group, thereby being unable to separate the two different relationships. It should be noted that in both studies LH and FSH concentrations were determined in morning urine or plasma samples. Both methods do not provide information on the night-day rhythm and on pulsatile secretory patterns, typical for the gonadotropic hormones [27]. Moreover, we observed in boys a lead/lag relationship of leptin concentrations Tanner stage 3 and FSH concentrations at Tanner stage 4. Our results show that during puberty the relationship between leptin and gonadotropic hormones is sex specific, since during puberty in boys leptin seems to have a different function and different timing. Still, previous studies have shown that leptin is essential in reproductive functioning in both boys and girls, as male and female sterile

ob/ob mice that became fertile again by leptin treatment [28]. The observations from our and other studies show that in a deficiency in leptin and fat mass as seen in anorexia nervosa or hyperleptinemia as seen in morbid obesity, will disturb the role of leptin during puberty, and thereby the start and progression of puberty [3].

Thus, we observed in our longitudinal cohort that during puberty the leptin/fat mass ratio decreased from Tanner stage 2 onwards. In girls, a peak in plasma leptin concentrations precedes a peak in LH and FSH concentrations, which supports a permissive role for leptin in the onset of puberty in girls. In girls, temporal relationships were observed between leptin and gonadotropic hormones during early puberty. In boys, however there were no peaks in leptin, LH, and FSH, and leptin was only related to LH and FSH during late puberty. We therefore conclude that in boys and girls during puberty factors independent of fat mass become (transiently) more important in the regulation of plasma leptin concentrations. Moreover, in girls leptin is suggested to act as a permissive factor for the onset of puberty, while in boys leptin has a different timing and possibly different function.

Acknowledgements

FR, SV, SL and NV carried out the study, collected and analyzed the data, and wrote the largest part of the manuscript. MW and AN supervised FR, SV, SL and NV. Planning, processing the results and writing the manuscript were done under general supervision by MW and AN. We want to thank our subjects for their participation in our cohort study. We gratefully thank Loek Wouters and Wendy Sluijsmans for their assistance.

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Chapter 6

The hypothalamic-pituitary-adrenal axis in the regulation of energy balance

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Physiology and Behavior 2008 May 23; 94 (2): 169-77

Abstract

Human (visceral) obesity is associated with alterations hypothalamus-pituitary-adrenal (HPA) axis functioning. It is however not completely clear whether the HPA axis is causally or co-incidentally related to (visceral) obesity. This review summarizes supporting data of an involvement of the HPA axis in the development of (visceral) obesity. First, several DNA polymorphisms related to HPA axis functioning are correlated to the development of obesity. Second, chronic elevation of circulatory glucocorticoid concentrations, as in Cushing's disease, results in increased abdominal adiposity. Third, (visceral) obesity is associated with a diminished capacity of cortisol to suppress its own secretion. HPA axis functioning might affect energy balance through affecting energy intake. Both CRH and cortisol influence physiological, central mechanisms involved in the regulation of food intake. Still, general activation of the HPA axis has shown to have inconsistent effects on food intake in humans. This inconsistency may partially be explained by gender differences, individual differences in the functioning of the HPA axis, as well as differences in attitude towards eating. In particular, women with high scores on dietary restraint are prone to stress-induced hyperphagia. Dietary restraint scores, in turn, are positively correlated to basal and dexamethasone-suppressed cortisol levels, indicating a complex dual relationship between stress, HPA axis functioning, attitude towards eating and the risk for stress-induced hyperphagia. In the Western society, with chronically high ambient levels of stress and the availability of high caloric foods, this relationship may imply a risk for the development of (visceral) obesity and the metabolic syndrome.

Introduction

Evidence suggests that the HPA axis is involved in the pathogenesis of human obesity, in particular characterized by visceral fat distribution. The HPA axis functioning plays an important role in the regulation of energy balance and body weight; it is, however, not completely clear whether the HPA axis is causally or co-incidentally related to the development of (visceral) obesity.

The hypothalamus-pituitary-adrenal (HPA) axis is a neuro-endocrine system involved in the stress-response, by regulating the secretion of cortisol [1]. The cascade of the HPA axis beholds that first the hypothalamus produces and releases corticotropin-releasing hormone (CRH), which subsequently stimulates the synthesis and release of adrenocorticotropin (ACTH) from the anterior pituitary. ACTH is produced from a larger precursor namely the proopiomelanocortin (POMC) protein, and stimulates the synthesis and release of cortisol by the adrenal cortex. In the circulation, cortisol is bound to corticosteroid-binding globulin (CBG) with high affinity, facilitating transport of cortisol in blood, followed by conversion of cortisol at the peripheral level by 11 β -hydroxysteroid dehydrogenase (11 β -HSD) 1 and 11 β -HSD 11. The 11 β -HSD 1 enzyme, converts cortisone into the active form, and 11 β -HSD 11 inactivates cortisol by conversion into cortisone. Cortisol exerts his actions through binding to and activation of two types of intracellular receptors, the glucocorticoid receptor (GR) and mineralocorticoid receptor (MR). The GR initiates or represses gene transcription, and induces negative feedback of the HPA axis through GR on the hypothalamus and pituitary level. The MR regulates basal activity of the HPA axis. Physiological cortisol levels follow a circadian rhythm; an early morning peak after awakening, a rapid decrease over the next few hours, and then a more gradual decline over the course of the day, to very low levels by bedtime [2].

Addison's disease and Cushing's syndrome represent two extremes of plasma cortisolism in human pathology. Addison's disease shows hypocortisolism, which causes symptoms such as fatigue, muscle weakness, changes in mood, and weight loss [3]. Also hypocortisolism caused by adrenalectomy will induce body weight loss in humans [4]. In animals, adrenalectomy will even prevent genetic as well as diet-induced form of obesity [5]. Cushing's syndrome is characterized by hypercortisolism, which causes symptoms such as hypertension, insulin resistance, hyperglycemia and rapid weight gain, particularly of the trunk and face with sparing of the limbs [6]. In the adipose tissue the glucocorticoids promote differentiation of pre-adipocytes into mature adipocytes and increase lipoprotein lipase activity, thus promoting adipose tissue increase and weight gain [7]. The clinical observations in Addison's disease and Cushing's syndrome clearly demonstrate a role for the HPA axis in the regulation of energy balance [8]. The pathology of visceral obesity shows a marked resemblance with Cushing's syndrome, including a central fat distribution of excess body-fat mass, elevated blood pressure, insulin resistance, and dyslipidemia. This form of obesity is associated with hypercortisolism, which becomes more pronounced when the HPA axis is stimulated (physiological or psychological) [9-11] and is associated with a diminished capacity of a low-dose (0.5 mg) dexamethasone (a GR agonist) to suppress plasma cortisol concentrations [12]. The latter finding suggest that the negative feedback loop, with cortisol activity on GR in both the hypothalamus and pituitary to inhibit HPA axis activity, is impaired in visceral obese subjects. This might explain (partially) the HPA hyperactivity and hypercortisolism of visceral obesity. In contrast, obese with a more peripheral body fat distribution do not show health risk markers, hypercortisolism, and diminished feedback capacity.

This review summarizes supporting data of an involvement of HPA axis hyperactivity in the development of (visceral) obesity. We start by describing physiological consequences of polymorphisms related to HPA axis functioning and the development of obesity. This will be followed by description of pathways through which the HPA axis might affect energy balance, and ultimately body weight, which will lead to a proposal of a model on the relationship between alterations in HPA axis functioning and (visceral) obesity.

Physiological consequences of polymorphisms related to HPA axis functioning

Visceral obesity shows considerable heritability, which results from contributions from many genes, so it is plausible that genetic variation in the cascade of the HPA axis may be involved in visceral obesity development [13]. Several polymorphisms in the cascade of the HPA axis have been described in the literature, which cause differences in HPA functioning and/or are involved in obesity development. In **table 1** the polymorphisms, the functional variation in HPA axis functioning caused by the mutations, and the possible association to obesity have been described. Regarding the initial step of the HPA axis cascade, a variant in the 5'-flanking region of the corticotropin-releasing hormone (CRH) gene (T255G) the of (CRH) has been found. This polymorphism has been associated with increased cortisol levels during physiological stress and total diurnal cortisol secretion, when combined with a variant in the gene of the GR. Still, no association between the T255G mutation and obesity has been shown yet [14].

With respect to functional variations of the POMC gene related to ACTH production, a rare mutation in exon 2 (C3804A) of the POMC gene, that causes ACTH insufficiency, has been associated with early-onset obesity [15].

Several polymorphisms have been described that can be linked to

glucocorticoid action and may be related to specific aspects of obesity. Thus, a (GTTT)_n repeat in intron 1 of the corticosteroid-binding globulin (CBG) has been associated with increased proliferation/differentiation of pre-adipocytes, higher salivary cortisol after dexamethasone suppression test, higher waist-to-hip ratio and a higher risk on obesity development [16, 17].

At the level of cortisol conversion, the ns4436A polymorphism of the 11b-hydroxysteroid dehydrogenase (11b-HSD) 1 gene has been associated with increased waist-to-hip ratio in women and pediatric obesity. No functional variation in HPA axis functioning caused by the ns4436A polymorphism has been described yet [18]. Additionally, a polymorphism at the (CA)_n repeat in the first intron of the 11b-HSD 11 gene has been associated with increased activity of the 11b-HSD 11 enzyme in vitro, which was observed in the subcutaneous adipose tissue of obese men. This results in higher local cortisol levels in adipose tissue and an increase in the ratio of urinary free cortisol/urinary free cortisone [19].

At the level of cortisol binding by the glucocorticoid receptor (GR) and mineralocorticoid receptor (MR) several polymorphisms have been described [20]. Three of the most prominent polymorphisms in the GR gene are the N363S, ER22/23EK and BclI polymorphisms, which all have been associated to obesity. The N363S polymorphism has been associated with increased glucocorticoid sensitivity, increased BMI, increased waist-hip ratio, and increased cholesterol levels [21, 22]. Secondly, the ER22/23EK polymorphism, which has been associated with decreased glucocorticoid sensitivity, decreased cholesterol levels, decreased insulin levels, and increased fat free mass [23]. Thirdly, the BclI polymorphism which has been associated with increased glucocorticoid sensitivity, increased insulin levels, increased blood pressure, increased abdominal visceral fat, and increased cholesterol levels [14, 24-26]. The most common described polymorphism in the MR gene is the

Table 1: Description of polymorphisms in the cascade of the HPA axis

Gene	Name	Obesity	Functional genetic variant
Corticotropin-releasing hormone (CRH)	T255G	No	increase cortisol levels
Proopiomelanocortin (POMC)	Exon 3 (AAC)(AGC) ₂ (GGC)	No	No influence
Proopiomelanocortin (POMC)	C3804A	yes	ACTH insufficiency
Corticosteroid binding globulin (CBG)	Intron 1 (GTTT) _n repeat	yes	Decrease feedback functioning
11b-HSD 1	ns4436A	yes	No influence
11b-HSD 11	Intron 1 (CA) _n repeat	yes	increase local cortisol levels
Glucocorticoid receptor (GR)	N363S	yes	Increase cortisol sensitivity
Glucocorticoid receptor (GR)	ER22/23EK	yes	Decrease cortisol sensitivity
Glucocorticoid receptor (GR)	BcI	yes	increase cortisol sensitivity
Mineralocorticoid receptor (MR)	M180V	no	increase cortisol secretion

MR180V polymorphism, which has been shown to enhance responses in cortisol secretion to a psychological stressor. In vitro studies showed a mild loss of function of the MR180V polymorphism, when cortisol was used as a ligand [27]. No associations have been found between the MR180V allele and the prevalence of obesity.

One can conclude that genetic variation in the cascade of the HPA axis does not only influence HPA axis functioning, but is also frequently related to differences in body composition and metabolic parameters. These findings stress that the HPA axis plays an essential role in the regulation of body weight, body composition and energy balance. The next paragraphs will extend on the physiological role of the HPA axis in the regulation of energy balance.

HPA axis functioning and energy balance

In the literature several possible mechanisms have been described through which the HPA axis might affect energy balance, which is composed of energy intake and energy expenditure. Obesity results from a chronic deregulation of energy balance, with energy intake exceeding energy expenditure, which leads to the storage of the excessive energy as fat.

HPA axis functioning and energy intake

The literature consists of ample human data on the effects of stress on food intake [28]; however, these data are difficult to interpret with respect to the role of the HPA axis for various reasons.

First of all, a wide variation in the type of stressor is used, resulting in different outcomes. Thus, stress in the workplace has been associated with higher energy intake [29, 30] while examination stress and surgical stress have produced mixed results [31, 32]. Secondly, there is a large inter-individual variance in HPA axis responses, i.e., plasma ACTH and cortisol production, to stress [8, 33]. A few studies on the effects of stress on food intake included measurements of plasma cortisol levels [34, 35]. Both studies showed a higher increase in stress-induced food intake in women who showed high cortisol responses to the stressor, suggesting a relationship between cortisol responses and food intake.

Most of the data directly indicating a crucial role of the HPA axis in the regulation of food intake are derived from animal studies, showing that the anorectic effects of adrenalectomy can be reversed by glucocorticoid (corticosterone) replacement [36, 37]. Human studies on the effects of glucocorticoids on food intake are limited, but generally support the findings in laboratory animals of an orexigenic effect of glucocorticoids [7, 33]. The stimulation of food intake by glucocorticoids seems to be macronutrient specific: when rats had a free choice for different chow compositions, corticosterone withdrawal (adrenalectomy) and subsequent replacement principally affected carbohydrate intake [38, 39]. In another paradigm, where adrenalectomized rats were offered a choice between chow and lard, corticosterone replacement dose-dependently increased intake of the more

palatable lard, but only in the presence of insulin [37]. These actions of glucocorticoids may underlie the preference for certain macronutrients and kinds of foods in humans after stress, in particular high saturated fat and sweet food items [30, 34, 40, 41]. Additionally, studies have shown that snack consumption may be more susceptible to stress-induced eating than meals [42]. Such foods may be preferred during stressful conditions, through learning that small energy-dense snacks are more easily ingested and digested when gut activity is suppressed by sympathetic arousal [43]. Several homeostatic and non-homeostatic mechanisms have been proposed through which HPA axis functioning may regulate food intake, and they will be discussed below.

HPA axis functioning and homeostatic regulation of food intake

The HPA axis seems to have many cross-links with the neuro-endocrine pathways that control the homeostatic regulation of food intake [44]. First of all, the CRH containing neurons, which comprise the initial component of the HPA axis, are located in the paraventricular nucleus (PVN) of the hypothalamus, which is also considered to be a major centre in the control of feeding behavior [45]. Central (intraventricular) administration of CRH inhibits feeding in rats [45, 46], and CRH is suggested to be an important intermediate in the anorectic effects of leptin [46-48]. The hypophagic effect of central CRH may (in part) result from inhibitory control of the orexigenic neural pathways involving neuropeptide Y (NPY), as paraventricular administration of the CRH-antagonist, alpha-helical CRF9-41, potentiated NPY-induced food intake [49]. CRH may be involved in the pathogenesis of obesity, at least in specific animal models: in the obese *fa/fa* Zucker rat a reduced level of CRH mRNA expression has been found [50]. Whether CRH is also involved in the pathogenesis of human obesity is unknown, but when subcutaneous fat cells from humans were incubated *in vitro* with CRH, it decreased 11 β -HSD 11 enzyme activity, thereby

reducing local cortisol production and reducing adiposity [48].

Thus, CRH exerts an opposite effect on appetite when compared to glucocorticoids. Physiologically, the (later) orexigenic effect of glucocorticoids, following the anorectic effect of CRH at the beginning of the stress response, may play a role in the recovery stage for the replacement of the replenishment of energy required during the 'fight or flight response'. The inhibitory effects of glucocorticoids on hypothalamic CRH release [46] may be involved in the stimulatory effect of glucocorticoids on food intake.

The opposite effects of CRH and glucocorticoids on food intake also emerge at a mechanistic level: where CRH exerts inhibitory control on NPY-induced food intake [49], (central) glucocorticoids potentiate the orexigenic actions of NPY. Additionally, dexamethasone stimulated NPY release in the mediobasal hypothalamus of female rats, and NPY production of cultured hypothalamic neurons [47]. In addition, recent research shows that cortisol up regulates the NPY Y2 receptor in the abdominal fat. Release of NPY and activation of the NPY Y2 receptor stimulates fat angiogenesis, proliferation and differentiation of new adipocytes, thereby linking HPA axis activation, NPY and increased abdominal fat storage [51].

The HPA axis interferes with leptin release, in that glucocorticoids stimulate secretion of leptin by (rat) adipocytes [52]. This stimulatory effect of cortisol on leptin secretion has also been shown *in vivo* [53, 54]. On the other hand, glucocorticoids may reduce the efficacy of leptin to suppress food intake, hence, induce leptin resistance. In rats, adrenalectomy has shown to increase the anorectic potency of leptin, while this effect was reversed by glucocorticoid replacement [55]. Leptin exerts its anorectic actions in part through activation of the central melanocortin system [56]. Similar to the results on leptin, the anorectic response to the melanocortin agonist, melanotan II, in

rats is enhanced by adrenalectomy, which can be reversed by glucocorticoid replacement [39]. These data suggest that glucocorticoids potentially decrease leptin sensitivity, partially through inducing insensitivity to central melanocortins. Induction of leptin resistance may also explain why glucocorticoid administration in humans results in elevated leptin concentrations on the one hand, and an increased food intake on the other [7]. Indeed, HPA axis hyperactivity, and a subsequent elevation of glucocorticoid concentrations, has been hypothesized to be involved in the leptin resistance of human obesity [57, 58].

Glucocorticoids also interact with the action and secretion of insulin, another endocrine adiposity signal. In vitro, glucocorticoids interfere with insulin-induced glucose uptake and metabolism in both cultured myocytes and adipocytes [59-62]. In vivo, dexamethasone administration to humans decreases whole body insulin sensitivity [63, 64]. This is associated with a compensatory increase in plasma insulin concentrations [65]. Since intraventricular administration of insulin has shown to suppress food intake [66], one might speculate that insulin (or insulin resistance) is involved in the effects of glucocorticoids on energy intake. Still, studies in glucocorticoid-treated adrenalectomized, streptozotocin-diabetic rats showed that total energy intake was not affected by insulin supplementation [36, 37], although insulin did dose-dependently increase the relative contribution of palatable high-fat food to total energy consumption [37]. Thus, insulin is likely to be involved in the effects of glucocorticoids on feeding behavior through modulation of food choice, rather than energy intake. Furthermore, through its anabolic actions on adipocytes, the hyperinsulinaemia caused by glucocorticoids may facilitate fat deposition. Remarkably, dexamethasone and insulin synergistically stimulate lipoprotein lipase activity in human adipose tissues, thereby facilitating free fatty acid uptake and lipogenesis [67].

HPA axis functioning and non-homeostatic regulation of food intake

In addition to the hypothalamic homeostatic pathways involved in food intake regulation, cortico-limbic brain areas are important structures for determining non-homeostatic regulation of food intake. These brain areas are involved in processes such as cognition, reward, memory and cognition, which are able to overrule the above-mentioned homeostatic regulatory mechanisms [44].

To a certain extent, humans show tendencies to cognitively moderate their food intake, a phenomenon that has been called "dietary restraint". Under some conditions, however, some subjects lose their cognitive control of caloric intake, which is called "disinhibition". The level of dietary restraint and disinhibition can be assessed by using specific questionnaires, such as the Three Factor Eating Questionnaire (TFEQ) [68]. We previously showed [Rutters, submitted] that the level of dietary restraint was positively correlated to 5-hour plasma cortisol patterns, and negatively correlated to the ability of the GR-agonist, dexamethasone, to suppress cortisol release under a strenuous exercise protocol. Accordingly, three studies reported that salivary cortisol at one time point (time point not specified) and 24-hour cortisol excretion were significantly higher in restrained women compared to unrestrained women [69-71]. Notably, some other studies did not find this relationship [72-74], possibly because cortisol concentrations were only measured at one time point or during the night.

One can speculate about the causality of this relationship. Although it cannot be excluded that high ambient cortisol levels increase cognitive awareness of caloric intake, the opposite (i.e., high level of dietary stress causes HPA axis activity) seems a somewhat more tempting hypothesis. Dietary restraint is positively correlated with body fat percentage [73, 75, 76], and the load of continuously restrained eating behavior is reportedly perceived as stressful [77]. Still, dietary restraint may be a risk factor for stress-

induced hyperphagia, as studies showed that restrained eaters experiencing psychological stress increased their food intake, while unrestrained eaters decreased food intake [30, 78-80]. Also, women with high disinhibition scores showed an increase in food intake following psychological stress [81, 82].

In the concept of non-homeostatic regulation of food intake, as lined out by Berthoud [44], food reward plays an important role. Like addictive drugs, palatable foods may act as a reinforcer, thereby influencing their own intake. Berridge and Robinson defined two distinct psychological processes determining the reinforcing value: "liking" and "wanting" [83]. These processes are regulated by different neural networks and mediated by different neurotransmitters [84].

In the process of "liking", mu-opioid systems in the nucleus accumbens seem to play an important role [85, 86]. Thus, local administration of the opioid agonist, morphine, or the mu-opioid agonist, (d-Ala(2), N-Me-Phe(4), Glycinol(5))-enkephalin (DAMGO), into the nucleus accumbens of rats increased food intake [85, 87]. In humans, the opioid antagonist, naltrexone, results in a reduction of both the intake and the reported pleasantness of food [88-90]. The expression of the mu-opioid receptor, and, hence, opioid sensitivity, is modulated by glucocorticoids: its expression is diminished in CRH-deficient mice and increased by corticosterone administration [91]. Indeed, activation of the endogenous opioid system has been suggested to be involved in stress-induced eating [92].

The endogenous opioid system may exert its effects in interaction with the endocannabinoid system, which has also been shown to be involved in the regulation of feeding behavior [93] and reward [94]. Endocannabinoid receptors of the CB1 type are abundantly expressed in the nucleus accumbens [95], and combined administration of the opioid antagonist, naloxone, and the CB1 receptor antagonist SR141716 decrease

food intake in rats in a synergistic fashion [96]. Direct evidence for a role of endocannabinoids in the orexigenic effects of glucocorticoids is lacking at present, but it is noteworthy that the endocannabinoid antagonist, AM281, antagonizes the effects of corticosterone on neuronal activity and on sexual behavior in amphibians [97]. With respect to the "wanting" or motivational aspect of reward, it is thought that dopaminergic neurons, originating in the ventral tegmental area (VTA) of the mid brain, and projecting to forebrain areas such as the orbitofrontal cortex and the nucleus accumbens or ventral striatum play a crucial role [83]. In rats, peripheral administration of corticosterone increases dopamine outflow in the nucleus accumbens [98]. Hence, both in rats [99] and in humans [100] glucocorticoids are suggested to contribute to the stress-induced increase of dopamine release in this brain area. The relationship between (striatal) dopamine and food intake seems, however, rather complex. Dopamine acts through several dopamine receptors (D1-D5), which seem to mediate distinct effects on food intake and food preference: selective D1 receptor activation resulted in increased caloric intake and preference for highly palatable foods, whereas combined D2/D3 activation showed an opposite effect [101]. Human studies showed that a single bolus administration of the dopamine transporter protein (DAT)-inhibitor, methylphenidate, results in a reduction of food intake, suggesting that high synaptic dopamine levels reduce the reinforcing value of food [102]. In contrast to acute effects, chronic elevation of synaptic dopamine may result in an increased wanting for rewarding foods, as indicated by the increased efforts and motivation of DAT-knockdown mice to obtain a sweet reward [103]. Possibly, shifts in the sensitivity and/or distribution of the different dopamine receptors may underlie this apparent discrepancy. In this case, it is noteworthy that obese subjects exhibit lower striatal D2 receptor availability [104], hence, it cannot be excluded that

in obesity the orexigenic effect of D1 activation may become considerably more predominant in the effects of dopamine on food intake. Whether this mechanism is also involved in the effects of glucocorticoids or stress on food intake in obesity, however, needs to be established. It should be noted, however, that above homeostatic and non-homeostatic pathways regulating feeding behavior are not 2 completely separate neural systems: significant interaction at different levels exists. In this respect, the adipocyte-derived hormone, leptin, has gained a particular interest. Leptin receptors have been found in the VTA, leptin has been shown to reduce firing rate of VTA dopaminergic neurons in brain slices, and direct administration of leptin into the VTA of rats decreased food intake [105]. Notably, a decreased performance in behavioral paradigms that assess the rewarding properties of food has also been observed when leptin was administered intracerebroventricular [106] and even intrahypothalamic [107]. Thus, leptin may reduce food intake by lowering the rewarding value of food. It is tempting to speculate that the leptin resistance following glucocorticoid supplementation [55] is also represented by a diminished influence of leptin on the neuronal pathways involved in food reward [43, 56]. In conclusion we described several mechanisms such as hormonal pathways, food reward and dietary restraint, through which HPA axis functioning influences energy intake, but also energy expenditure is influenced.

HPA axis functioning and energy expenditure

With respect to the effect of HPA axis functioning on energy expenditure, the literature is not as extensive as on the effect on energy intake. Like with the regulation of food intake, CRH and cortisol may exert distinct effects on energy expenditure. CRH infusion in humans resulted in a 14% increase in resting energy expenditure [108]. CRH may act through activation of the central sympathetic system, both in the hypothalamus as well as in the locus

ceruleus of the brain stem [1]. However, as the increase in energy expenditure following CRH infusion was not associated with increases in plasma catecholamines, other mechanisms may also be involved.

With respect to the effects of glucocorticoids on energy expenditure, the literature shows a dichotomous response. When hydrocortisone was infused for 60 h, it led to an increase in resting energy expenditure [109]. Also, when hydrocortisone was infused for 16h at two different concentrations, it led to an increase in resting energy expenditure of 9-15% [110]. On the contrary, two other studies showed that infusion of glucocorticoids for 168 h did not alter energy expenditure [111]. In addition, chronic dexamethasone treatment in infants did not affect energy expenditure [112]. Also, no difference in resting energy expenditure (corrected for the amount of lean body mass) was found between normal and patients with Cushing's syndrome [113, 114]. It may be hypothesized that glucocorticoids exert dual effects on energy expenditure, and that the stimulatory actions of glucocorticoids on energy expenditure [109, 110] are overruled by other, inhibitory actions during prolonged or chronic exposure [115].

Several mechanisms may underlie the inhibition of energy expenditure by glucocorticoids. Firstly, suppression of CRH release by glucocorticoids as part of the negative feedback regulation of HPA axis functioning [1]. Secondly, it has been shown that chronic increased levels of cortisol in Cushing's disease leads to inhibition and promotes breakdown of muscle, resulting in muscle wastage. Since the amount of muscle mass is positively correlated to (resting) energy expenditure, chronic hypercortisolism, as in patients with Cushing's disease, may ultimately result in lower energy expenditure [61]. Thirdly, glucocorticoids have shown to affect the hypothalamus-pituitary-thyroid axis. Glucocorticoids inhibit the production of thyrotropin-releasing hormone as well as thyroid-

stimulating hormone, and the conversion of thyroxine (T4) to triiodothyronine (T3), ultimately resulting in decreased plasma T3 levels [1, 116], which may subsequently result in a reduced basal metabolic rate [115].

In summary we described that glucocorticoids exert dual effects on energy expenditure, and that the stimulatory actions of glucocorticoids on energy expenditure are overruled by other, inhibitory actions during prolonged or chronic exposure.

Summary and conclusions

The hypothalamus-pituitary-adrenal axis affects energy balance on different levels, at different ways, and with different underlying mechanisms. All in all, prolonged exposure to elevated glucocorticoid levels may result in a positive energy balance, by increasing energy intake, without affecting resting energy expenditure. The stimulatory effects of glucocorticoids on energy intake may involve both homeostatic and non-homeostatic pathways involved in the regulation of eating behavior. The homeostatic pathways may include a suppression of CRH release, induction of leptin resistance and the stimulation of NPY release. Non-homeostatic pathways may include the individuals' attitude towards eating, as well as neural systems involved in both the "wanting" (dopaminergic) and "liking" (opioidergic) aspects of food reward. The latter systems may, just such as insulin [37], also be involved in a shift in food choice towards high energetic sweet and high fat foods. The effects of glucocorticoids on energy expenditure may depend on the duration of exposure, with mechanisms such as a decrease in CRH release, increased loss of lean body mass and decreased circulating T3 levels ultimately counteracting an initial stimulation of resting metabolic rate.

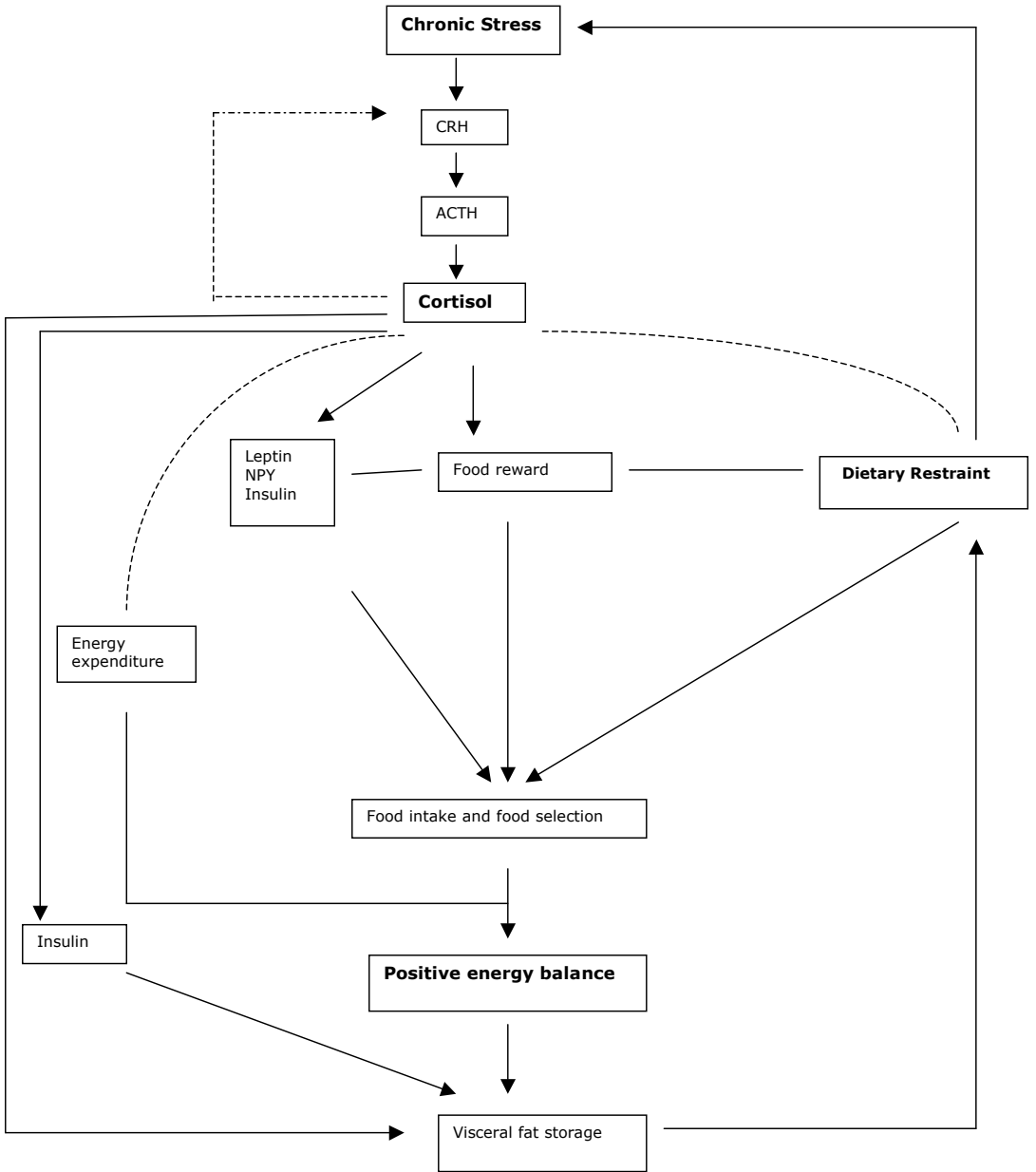
The resulting positive energy balance is likely to result in increased lipogenesis and fat storage, under influence of glucocorticoids and elevated concentrations of insulin. In addition,

glucocorticoids may even increase adipogenesis due to an up-regulation of the NPY Y2 receptor in abdominal adipose tissue [51]. Notably, visceral adipocytes have a fourfold higher number of glucocorticoid receptors than adipocytes in other fat depots [8]. Accordingly, the stimulatory effects of dexamethasone on lipoprotein lipase activity were more pronounced in visceral adipocytes than in subcutaneous adipocytes [67]. Thus, the increased fat storage due to prolonged hypercortisolism may show some site-specificity, with a preference for the visceral regions.

The increased adiposity may lead to an increase in weight dissatisfaction [75], which will lead to dieting in an attempt to achieve and maintain desired body weight. Dieting is done through cognitively controlling eating behavior and, hence, is accompanied with an increase in dietary restraint [117]. However, dietary restraint can act as a chronic physiological stressor, and like all other chronic stress will lead to alterations in HPA axis functioning. We found that the level of dietary restraint was positively correlated to 5-hour plasma cortisol patterns, and negatively correlated to the ability of the GR-agonist, dexamethasone, to suppress cortisol release under a strenuous exercise protocol [Rutters, in press].

Therefore, we propose that chronic hyperactivity of the HPA axis initiates a vicious circle (**figure 1**), which puts chronic stress as a major risk factor for excessive weight gain and (visceral) obesity. Whether this mechanism is the primary cause for the obese state in the visceral obese remains to be elucidated, but it should be noted that HPA axis hyperactivity is typical in visceral obese subjects [9-11] and that HPA axis functioning in obese women was not significantly affected by weight loss [118]. Therefore, the HPA axis may therefore be an important causal factor in to the obesity epidemics of the Western society, where high levels of ambient stress and availability of high fat, sweet foods are abundantly present.

Figure 1: a model for the *possible* relationship between HPA axis functioning, eating behavior, energy balance, and body composition (solid lines represent stimulation and dashed lines represent inhibition).



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Chapter 7

Sex specific associations between the BclI polymorphism in the glucocorticoid receptor gene and HPA axis exposure as well as feedback sensitivity under stress

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Submitted

Abstract

Objective: to investigate, in men and women, the relationship between the BclI genotypes and circulating cortisol concentrations, HPA axis feedback sensitivity, as well as HPA axis feedback sensitivity under stress.

Methods: we assessed in 77 men and 89 women, aged between 18 to 55 y and BMI between 19.8 and 41.2 kg/m², HPA axis functioning through measuring 5-hour cortisol exposure as well as cortisol feedback sensitivity under stress through a standardized high intensity test with 4 mg dexamethasone.

Results: the BclI genotype frequency distribution (CC/CG/GG) was 41/42/17 in men and 45/42/13 in women. The G/G BclI genotype had higher cortisol concentrations after dexamethasone exposure when compared to the C/C genotype in men (42.2 ± 8.7 vs. 27.3 ± 2.2 nmol/L, $p < 0.05$), and to the C/G genotypes in women (70.5 ± 14.6 vs 47.1 ± 9.1 nmol/L, $p < 0.05$). Only in women, the G/G BclI genotype had a higher area under the curve (AUC) of the cortisol concentrations over a 5-hour time period when compared to the C/G genotype (178458 ± 85818 vs. 135063.7 ± 65511 nmol/L.min, $p < 0.05$). Additionally, women with the G/G BclI genotype had a higher AUC of the cortisol concentrations over a 70 minutes time period after a high intensity test with dexamethasone suppression when compared to the C/G and C/C genotypes (3788.7 ± 2248 vs. 3323.2 ± 1716 and 5392.9 ± 3545 nmol/L.min, $p < 0.05$).

Conclusion: the BclI genotype is sex specifically associated with increased cortisol exposure as well as decreased cortisol feedback sensitivity under stress.

Introduction

The hypothalamus/pituitary/adrenal (HPA) axis is a neuro-endocrine system involved in the stress-response and homeostasis, by regulating the secretion of cortisol [1]. Cortisol affects several systems, such as the cardiovascular system, the immune system, metabolism, cell growth, behavior, and the HPA axis itself through negative feedback [2]. Literature has shown large intra-individual and inter-individual variation in HPA axis functioning, such as the efficacy of cortisol to suppress its own secretion (negative feedback). This results in considerable variation in circulation cortisol concentrations [3]. The variation has been related to several polymorphisms of the glucocorticoid receptor gene, such as N363S [4], the ER22/23EK [5], Tth111I [6], and BclI polymorphisms [3]. The latter polymorphism, BclI (rs41423247), has been related to insulin resistance in obese women [7], increased BMI, waist-hip ratio, leptin levels [6, 8-10], as well as better weight loss during a very low calorie diet [11, 12]. Additionally, in men, the BclI G/G genotype has been associated with increased salivary concentrations of cortisol [6], increased levels of salivary cortisol after a standardized lunch [6], increased levels of cortisol after a 0.5 mg dexamethasone suppression test [6], as well as a diminished cortisol response to psychosocial stress [2]. In women, however the BclI G/G genotype has been associated with an increased cortisol response to psychosocial stress [13], which suggests possible sex specific associations between the BclI genotypes and HPA axis functioning. Besides the study from Kumsta et al. 2007 [13], no previous studies have examined the possible sex specific associations, since other studies only included males [2, 6]. Furthermore, HPA axis functioning was studied via basal cortisol levels, a dexamethasone suppression test, or subjection to a psychosocial test, never via a combination of physical stress and dexamethasone suppression, which

indicates sensitivity of the HPA axis to negative feedback under stress [14]. Consequently, the objective of our study was to investigate, in men and women, the relationship between the BclI genotypes and circulating cortisol concentrations, HPA axis feedback sensitivity, as well as HPA axis feedback sensitivity under stress.

Materials and methods

Protocol

The Medical Ethical Committee of the University of Maastricht approved of the study, and written consent was obtained from all participants. Healthy, medication-free, non-smoking men (n = 80) and women (n = 90) aged between 18 to 55 y and BMI between 19.8 and 41.2 kg/m² were recruited to complete all phases of the study. A medical history was obtained from each participant before entry into the study and exclusion criteria were menopause, chronic illness, depression, and a history of eating disorders or current dieting. From 170 participants recruited, 77 men and 89 women completed the study, 4 participants dropped out before the actual start of the study.

From each participant we obtained: 5-hour cortisol exposure, a progressive maximal cycling test to volitional exhaustion and a high intensity test under 4 mg dexamethasone, anthropometry, and BclI genotypes.

Measurements

5-hour cortisol exposure

At 8 a.m. after an overnight fast an intravenous catheter for blood sampling was inserted into the forearm vein. Direct after insertion of the catheter the first blood sample (9 ml) was drawn (0 minutes), and after 15 minutes the participants received their breakfast (20 % of the participants' daily energy requirement). At 45, 90 and 210 minutes the second, third and fourth blood samples were drawn. At 225 minutes the participants received their lunch (30 % of the participants' daily energy

requirement) and at 255 and 300 minutes the fifth and sixth blood samples were drawn. During the test participants were able to drink water ad libitum. Breakfast and lunch consisted of typical Dutch products namely brown bread (1.0 MJ per 100 g; en% C/P/F: 74/15/11) with young cheese (1.6 MJ per 100 g; en% C/P/F: 0/26/74) as well as ab libitum tea or water consumption. The daily energy requirements were calculated by multiplying the basal metabolic rate (BMR) by an activity index of 1.75. The BMR (MJ/day) was calculated according to the equations of Harris-Benedict [15]. Participants remained seated during sampling, and daytime naps were not allowed. The participants' 5-hour cortisol exposure was measured by calculating the area under the curve (AUC) (0-300 minutes).

Progressive maximal cycling test to volitional exhaustion

The 5-hour cortisol exposure measurement and the progressive maximal cycling test (VO^2_{max}) were separated by at least one day. Each participant underwent a progressive maximal cycling test to volitional exhaustion for quantification of VO^2_{max} [16]. Participants cycled on an electromagnetically braked bicycle ergometer (Lode, Groningen, The Netherlands) at 80 rpm, at a workload of 75 W for women and 100 W for men [16]. After 5 minutes, the workload was increased every 2.5 minutes with 50 W, until participants were no longer able to maintain a pedaling frequency above 80 rpm. Open-circuit spirometry was performed with an Oxycon (Mijnhardt Jaeger, Mannheim, Germany), and oxygen consumption (VO^2) and carbon dioxide production (CO^2) were measured in the breath-by-breath mode and averaged over 15 seconds [16]. Heart rate was monitored continuously throughout all exercise protocols (Polar s610). Verification that each participant actually achieved VO^2_{max} consisted of meeting the following criteria: 1) achieving predicted maximal heart rate and 2) a respiratory exchange ratio of 1.10 or more

[16]. The results of the progressive maximal cycling test were used to measure the resistance to elicit an exercise intensity of 90 % of each participants maximal workload (W_{max} , V) for the high intensity test under 4 mg dexamethasone.

High intensity test under 4 mg dexamethasone

The progressive maximal cycling test and high intensity test under 4 mg dexamethasone [14] were separated by at least one week. Each participant received 4 mg dexamethasone 8 hours before the high intensity test. Each participant participated in the treatment, and no adverse reactions were reported. Participants abstained from alcohol consumption and strenuous exercise 15 h before testing. At 8 a.m. after an overnight fast, an intravenous catheter for blood sampling was inserted into the forearm vein and the first blood sample was drawn for a baseline measurement (0 minutes). After 5 minutes the participants received their breakfast (20 % of the participants' daily energy requirement). At 15 minutes the second blood sample was taken and the high intensity test was started. At 30 minutes the high intensity test ended and the third blood sample was taken. At 40, 50, 60 and 70 minutes, the fourth, fifth, sixth and seventh blood samples were taken [14]. The high intensity test consisted of 15 minutes of cycling. The initial 5 minutes served as a warming-up, during which each participant cycled at a workload of 40 % of his/her W_{max} (V), at 80 rpm. After the warming-up, the workload was increased with 50 % of his/her W_{max} (V) for 30 sec at 80 rpm, followed by a decrease of the workload with 50 % of his/her W_{max} (V) for 30 sec at 80 rpm. The increase and decrease of workload were repeated 10 times. After the third blood sample was drawn a 5 minutes cooling-down was performed at 30 % of his/her W_{max} (V) at 60 rpm [14].

Blood samples

Blood samples were collected and mixed in serum tubes (BD Vacutainer, 9.5 ml).

Serum was obtained by centrifugation (4°C, 3000 rpm, 10 minutes), frozen in liquid nitrogen and stored at -80°C until analysis of cortisol concentrations. Serum cortisol concentrations were measured using RIA (Buhlmann Laboratories) and detection limits for cortisol concentrations were 2.9 ng/ml. Intra-assay coefficient variation for cortisol concentrations was less than 6 % for all samples.

Anthropometry

Measurements were carried out in the morning after voiding the bladder and before breakfast. Body weight and height were measured to the nearest 0.01 kg and 0.1 cm respectively. BMI (kg/m²) was calculated as body weight divided by height (m) squared.

Determination of BclI genotypes

The genomic DNAs of 166 participants were isolated from peripheral blood leukocytes using a QIAamp kit (QIAGEN, Germany). A 87 bp fragment of the GRL gene was generated from genomic DNA by PCR using forward primer 5'-GCTCACAGGGTTCTTGCCATA-3' and reverse primer 5'-TTGACCATGTTGACACCAAT-3', which includes a C/G polymorphism in intron 2, 646 nucleotides downstream from exon 2 [11]. The PCR products were digested with BclI at 50°C for 60 minutes, electrophoresed on a 3% agarose gel and stained with ethidium bromide. The expected products after digestion with BclI are 87 bp for G/G homozygotes, 47 and 40 bp for C/C homozygotes, and 87, 47, and 40 bp for G/C heterozygotes.

Statistical analysis

Student t-test and ANOVA (for continuous variables) were executed to determine differences in single variables between groups. Hardy-Weinberg equilibrium for BclI was calculated using the chi-square test. The BclI polymorphism was in Hardy-Weinberg equilibrium. The differences over time and between conditions were determined using three-factor ANOVA with repeated measures. All statistical tests

were two-tailed, differences were considered significant at $p < 0.05$ and values are expressed as mean \pm standard deviation.

Results

Table 1 shows the characteristics of the male (n = 77) and female (n = 89) participants. No significant differences were shown in age, BMI, and BclI genotype frequency distribution between men and women. Men showed a higher body weight, height, and maximal workload during the progressive maximal cycling test when compared to women. Women showed a higher area under the curve (AUC) of the cortisol concentrations over a 5-hour time period, as well as a higher AUC of the cortisol concentrations over a 70 minutes time period after a high intensity test with dexamethasone suppression. Therefore, the results of men and women were analyzed separately.

Table 1. Characteristics of the male (n = 77) and female (n = 89) participants

	Male (n = 77)	Female (n = 89)	P value ¹
Age (year)	29 \pm 10	26.9 \pm 9	0.16
Body weight (kg)	81.4 \pm 13	70.1 \pm 12.4	0.001
Height (cm)	181.6 \pm 6.8	169.3 \pm 6.6	0.001
BMI (kg/m ²)	24.4 \pm 23.6	24.4 \pm 4.1	0.99
Maximal workload (V)	299.2 \pm 58	208.9 \pm 43	0.001
AUC ² of 5-hour cortisol exposure (nmol/L.min)	9.8 (10 ⁵) \pm 2.9	14.9 (10 ⁵) \pm 7.3	0.001
AUC ² cortisol after high intensity test (nmol/L.min)	2556 \pm 1970	4217 \pm 4445	0.004
BclI genotype frequency distribution (CC/CG/GG) (%)	41/42/17	45/42/13	0.80

¹ Differences between men vs. women (t-test for continuous variables or Chi-square test for nominal variables)

²AUC, area under curve

Figure 1A shows the changes in cortisol concentrations (nmol/L.min) over a 5-hour time period in men. No differences in cortisol concentrations were observed between the three BclI genotypes. Over time there was a significant decrease of cortisol concentrations in all men, however, over time no differences were seen between the three BclI genotypes.

Figure 1A. Changes in cortisol concentrations (nmol/l) over a 5-hour time period in men with the BclI G/G (n = 13), CC (n = 32) or C/G (n = 33) genotype. Values are means with standard errors.

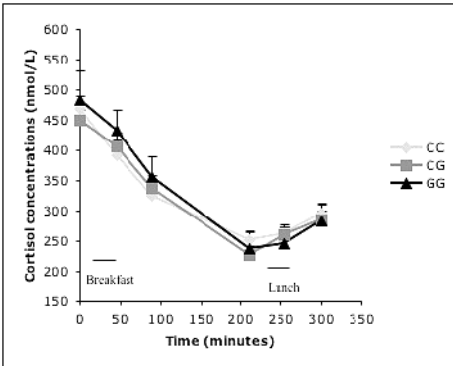
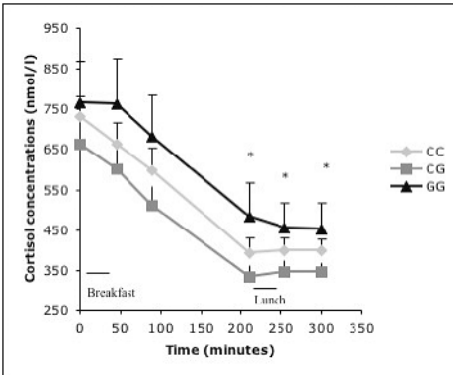


Figure 1B. Changes in cortisol concentrations (nmol/l) over a 5-hour time period in women with the BclI G/G (n = 12), CC (n = 39) or C/G (n = 38) genotype. Values are means with standard errors.

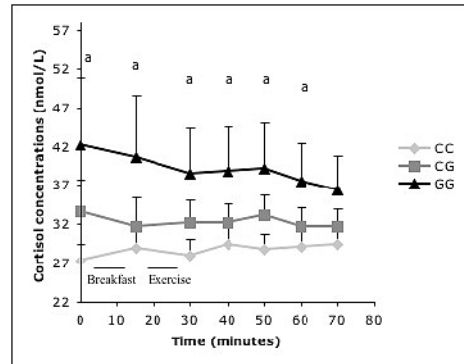


* P < 0.05 for differences between women with the BclI G/G and the C/G genotype (ANOVA)

Figure 1B shows the changes in cortisol concentrations (nmol/L.min) over a 5-hour time period in the women. The cortisol concentrations were significantly higher at 210, 255, and 300 minutes in women with the G/G genotype when compared to the C/G genotype. Over time there was a significant decrease of cortisol concentrations in all women, however, over time no differences were seen between the three BclI genotypes. The area under the curve (AUC) of the cortisol concentrations over a 5-hour time period was significantly higher in women with the G/G genotype when compared to women with the C/G genotype (178458 ± 85818 vs. 135063.7 ± 65511 nmol/L.min, $p < 0.05$).

Figure 2A shows the changes in cortisol concentrations (nmol/L.min) over time after a high intensity test with dexamethasone suppression in men. After dexamethasone exposure ($t=0$), the cortisol concentrations were significantly higher in men with the G/G genotype when compared to men with the C/C genotype (42.2 ± 8.7 vs. 27.3 ± 2.2 nmol/L, $p < 0.05$). After the high intensity test in combination with dexamethasone suppression, the cortisol concentrations were significantly higher at 15, 30, 40, 50, and 60 minutes in men with the G/G genotype when compared to men with the C/C genotype. Over time as well as between the three genotypes no differences were observed in the men.

Figure 2A: Cortisol concentrations (nmol/l) over time after a high intensity test with dexamethasone suppression in men with the BclI G/G (n = 13), CC (n = 32) or C/G (n = 33) genotype. Values are means with standard errors.

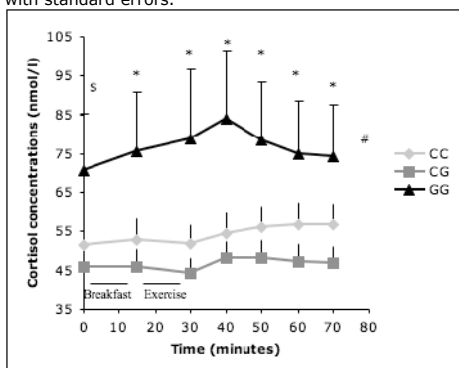


^a P < 0.05 for differences between men with the BclI G/G and the C/C genotype (ANOVA)

Figure 2B shows the changes in cortisol concentrations (nmol/L.min) over time after a high intensity test with dexamethasone suppression in women. After dexamethasone exposure ($t=0$), the cortisol concentrations were significantly higher in women with the G/G genotype when compared to women with the C/G genotype (70.5 ± 14.6 vs. 47.1 ± 9.1 nmol/L, $p < 0.05$). After the high intensity test in combination with dexamethasone suppression, the cortisol concentrations were significantly higher at all time points in women with the G/G genotype when compared to women with the C/G and C/C genotype. From 0 to 40 minutes a

significant increase in cortisol was seen in women with the G/G genotype, while this was absent in women with the C/G and C/C genotypes (13.1 ± 15.1 vs. 8.5 ± 38 and 3.1 ± 9.4 nmol/L, $p < 0.05$). Additionally, from 40 to 70 minutes a significant decrease in cortisol was seen in women with the G/G genotype, while an increase in cortisol was seen in women with the C/G and C/C genotype (-8.2 ± 21.6 vs. 0.6 ± 12.6 and 2.2 ± 9.8 nmol/L, $p < 0.01$).

Figure 2B: Cortisol concentrations (nmol/l) over time after a high intensity test with dexamethasone suppression in women with the BclI G/G (n = 12), CC (n = 39) or C/G (n = 38) genotype. Values are means with standard errors.



* $P < 0.05$ for differences between women with the BclI G/G and the C/G as well as C/C genotype (ANOVA)

§ $P < 0.05$ for differences between men with the BclI G/G and the C/C genotype (ANOVA)

$P < 0.05$ overall group x time interaction

The AUC of the cortisol concentrations corrected for baseline, over a 70 minutes time period after a high intensity test with dexamethasone suppression was significantly higher in women with the G/G genotype when compared to women with the C/G genotype (460 ± 408 vs. 38.3 ± 666 nmol/L.min, $p < 0.03$).

Discussion

The objective of our study was to investigate, in men and women, the relationship between the BclI genotypes and circulating cortisol concentrations, HPA axis feedback sensitivity, as well as HPA axis feedback sensitivity under stress. In both men and women the G/G BclI genotype was associated with increased cortisol concentrations after a dexamethasone exposure test. Only in

women the G/G BclI genotype was associated with increased cortisol concentrations over a 5-hour time period, and increases in cortisol concentrations after a high intensity test with dexamethasone suppression.

With these results, our study adds to the growing literature on the relationship between the BclI genotypes and HPA axis functioning, and this study is the first to analyze the effect of the BclI genotypes on HPA axis feedback sensitivity under stress, as well as sex specific associations. In our study, we measured cortisol concentrations (nmol/L.min) over a 5-hour time period to determine cortisol exposure under resting conditions. We showed that women with the G/G BclI genotype had significant higher cortisol concentrations just before and after lunch, in addition to a higher AUC of the cortisol concentrations over a 5-hour time period when compared to the C/G genotype. This suggests that women with the G/G BclI genotype have increased daily cortisol exposure.

Additionally, we measured feedback sensitivity through a dexamethasone exposure test [17], which determines the amount of cortisol that escapes from dexamethasone suppression (a type II glucocorticoid receptor agonist) [6]. When cortisol feedback is strong low amounts of cortisol will be present, and when high amounts of cortisol escape, cortisol feedback is decreased [17]. We showed that both in men and women the G/G BclI genotype was associated with higher cortisol concentrations after a dexamethasone exposure test, when compared to other genotypes. The results in men are in accordance with literature [6], and together these results imply that men and women with the G/G BclI genotype have decreased sensitivity to negative feedback of the HPA axis.

Furthermore, to assess HPA axis feedback sensitivity under stress, we measured cortisol concentrations (nmol/L.min) corrected for baseline, over a 70 minutes time period after a high intensity test in

combination with dexamethasone suppression [14]. The high intensity exercise promotes the escape of cortisol from dexamethasone suppression, which indicates cortisol feedback sensitivity under stress [14]. We showed that after the high intensity test with dexamethasone suppression women with G/G BclI genotype had higher amounts of cortisol that escaped, as well as higher AUC of the cortisol concentrations, when compared to women with the C/G genotype. These results imply that women with the G/G BclI genotype have decreased sensitivity of the HPA axis to negative feedback under stress.

Besides our study and the study from Kumsta et al. 2007 [13], no previous studies examined possible sex specific associations between BclI polymorphisms and HPA axis functioning, since other studies only included males [2, 6]. Kumstra et al. 2007 [13] have shown a sex specific association between BclI and HPA axis responses to psychosocial stress; together with the present study, these results point towards sex by genotype interactions, which suggests that the same genetic variant of the GR gene can have differential, or perhaps even

opposite effects on HPA axis functioning depending on sex. The BclI polymorphism of the glucocorticoid receptor is thought to impact the efficacy of cortisol signaling and thereby influences the downstream biology of peripheral and central cortisol responsive systems [6]. The BclI polymorphism might therefore be one of the factors that underlie individual susceptibility to stress related disorders, such as depression and the metabolic syndrome [18]. From our study we conclude that the BclI genotype is sex specifically associated with increased cortisol exposure as well as decreased cortisol feedback sensitivity under stress.

Acknowledgements

We want to thank our subjects for their participation in this study. We gratefully thank José Sulon for his assistance. FR carried out the study, collected and analyzed the data, and wrote the largest part of the manuscript. FB (supervised by EM) contributed to the practical work. SL and JB reviewed the manuscript. AN and MW supervised FR. Planning, processing the results and writing the manuscript were done under general supervision by AN and MW. The authors had no conflict of interest.

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Chapter 8

Hypothalamus/Pituitary/Adrenal (HPA) axis functioning in relation to body fat distribution

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Submitted

Abstract

Objective: to relate HPA axis functioning and HPA feedback functioning to body fat distribution in normal weight to obese subjects.

Patients: 91 men and 103 women (age 18-45y, BMI 19-35 kg/m², waist-to-hip ratio (WHR) 0.6-1.1)

Measurements: we assessed anthropometry, body composition through hydrodensitometry and deuterium dilution method, cortisol variability through measuring 5-hour cortisol concentrations, HPA axis feedback functioning through a dexamethasone suppression test, and HPA axis functioning under in a challenged condition through a standardized high intensity test under 4 mg dexamethasone.

Results: in men an inverse relationship was observed between 5h cortisol exposure (nmol/ml) and fat mass index (FMI) (kg/m²) ($r=0.55$, $p<0.001$). In women inverse relationships were observed between 5h cortisol exposure (nmol/ml) and WHR ($r=0.49$, $p<0.001$), maximal workload ($r=0.32$, $p<0.001$) as well as oral contraceptive use ($r=0.38$, $p<0.001$). Similarly, in men an inverse relationship was observed between negative feedback expressed as baseline cortisol concentrations minus post dexamethasone cortisol concentrations (nmol/ml) and FMI ($r=0.53$, $p<0.001$). In women an inverse relationship was observed between negative feedback expressed as baseline cortisol concentrations minus post dexamethasone cortisol concentrations (nmol/ml) and WHR ($r=0.43$, $p<0.001$), maximal workload ($r=0.30$, $p<0.001$) as well as oral contraceptive use ($r=0.43$, $p<0.001$) in women. Moreover, an inverse relationship was observed between HPA axis functioning in a challenged condition expressed as percentage increase of cortisol concentrations after standardized high intensity test under 4 mg dexamethasone (%) and waist ($r=0.21$, $p<0.10$) in men and WHR ($r=0.21$, $p<0.05$) in women. In male subjects, strong positive relationships were observed between FMI and waist circumference ($r = 0.85$, $p<0.001$), as well as waist-to-hip ratio ($r = 0.70$, $p<0.001$).

Conclusion: disturbance of HPA axis functioning under basal and challenged conditions is related to visceral fat accumulation.

Introduction

The hypothalamus-pituitary-adrenal (HPA) axis is a neuro-endocrine system that plays an important role in the regulation of the stress-response [1]. The HPA axis is suggested to be involved in the pathogenesis of human obesity, in particular characterized by visceral fat distribution (visceral obesity). Support for the involvement of the HPA axis in the regulation of body weight and body fat distribution is found in two extremes of plasma cortisol levels in humans; Addison's disease (hypocortisolism) that is related to weight loss, and Cushing's syndrome (hypercortisolism) that is related to rapid weight gain, particularly of the trunk and face with sparing of the limbs [2].

Additional support is found in obese subjects with visceral fat accumulation that show decreased salivary and serum cortisol levels [3-8], increased urinary secretion of cortisol [6, 9], decreased cortisol exposure [3, 4, 10-12], and enhanced awakening cortisol response [13]. In both lean and obese subjects visceral fat distribution has been associated with increased cortisol secretion after physical and psychological stressors [3-6, 9, 14]. These conflicting findings, such as decreased exposure and increased secretion after stressors, suggest a reset of the HPA axis toward chronic hyper-activation [15]. The ability of cortisol to suppress its own secretion through activation of hypothalamic and pituitary glucocorticoid receptors (negative feedback) is an important part of adequate HPA axis functioning. Studies on the relationship between negative feedback and body fat distribution have however yielded inconclusive results. Some studies found an increased capacity of dexamethasone (a glucocorticoid receptor agonist) to suppress plasma cortisol concentrations [4, 16], some studies found no differences in the capacity of dexamethasone to suppress plasma cortisol concentrations [3, 5, 6, 10] and some even a decreased capacity [17, 18]. This inconclusiveness

possibly results from differences in techniques to test HPA feedback functioning.

Ten years ago, Petrides et al. (1997) [19] developed a more elaborate technique to test HPA axis functioning, using a physical challenge to promote escape of cortisol from dexamethasone suppression. Using this technique they observed even in a relative homogenous population of young men a large inter-individual variance in HPA functioning [19]. It remains unknown whether inter-individual variation in HPA axis functioning can be related to fat distribution in a normal-weight to obese population, and whether HPA axis functioning tested under 4 mg dexamethasone in combination with a physical stressor is related to fat distribution. Therefore, the objective of our study was to relate HPA axis functioning and HPA feedback functioning to body fat distribution in normal weight to obese subjects.

Subjects and methods

Subjects and protocol

The study was approved by the Medical Ethical Committee of Maastricht University, and a written consent was obtained from all subjects. Healthy, medication-free, non-smoking men ($n = 91$) and women ($n = 105$) aged between 18 to 45 y and with a BMI between 19 and 35 kg/m² were recruited to complete all phases of this study. From 196 subjects recruited, 91 men and 103 women completed the study, 2 subjects dropped out before the actual start of the study. A medical history was obtained from each subject before entry into the study containing questions on illness, medicine use, smoking habits, daily coffee intake, weekly alcohol intake and food allergies. Exclusion criteria for our study were: smoking, excessive alcohol consumption, chronic illness, depression, and a history of eating disorders or current dieting. The women were examined regardless of the phase of the menstrual cycle. From each subject we obtained:

anthropometry as well as body composition, 5-hour cortisol exposure measurement, a progressive maximal cycling test to volitional exhaustion and a high intensity test under 4 mg dexamethasone.

Measurements

Anthropometry and body composition

Measurements were carried out in the morning after voiding the bladder and before breakfast. Body weight (Sauter D7470, Ebingen, Germany) and height (Seca, model 220, Hamburg, Germany) were measured to the nearest 0.01 kg and 0.1 cm respectively. BMI (kg/m^2) was calculated as body weight (kg) divided by height (m) squared. The waist circumference was measured to the nearest 0.1 cm at the site of the smallest circumference between the rib cage and the iliac crest, whereas the hip circumference was measured to the nearest 0.1 cm at the level of maximum extension of the buttocks. The waist-to-hip ratio (WHR) was calculated accordingly. Both measurements were done with the subjects in standing position. Additionally, both waist circumference and the WHR were used to define different patterns of body fat distribution. Waist circumference and waist-to-hip measurements were used as surrogates for body fat distribution measurements, as anthropometric measurements have been shown to predict total and abdominal fat content [20-22].

Body density was determined by hydrodensitometry with simultaneous measurements of residual lung volume with the helium dilution technique. Total body water (TBW) was determined with the deuterium dilution method following the Maastricht protocol [23]. Body composition was calculated from body density and TBW using the three-compartment model of Siri. Fat free mass (FFM) was calculated by dividing TBW by the hydration factor 0.73. Fat mass (FM) was determined as $\text{BW} - \text{FFM}$. Fat mass index (FMI) was calculated by fat mass/height² (kg/m^2) and fat free mass

index (FFMI) was calculated by fat free mass/height² (kg/m^2).

5-hour cortisol exposure measurement

At 8 a.m. after an overnight fast an intravenous catheter for blood sampling was inserted into the forearm vein. Direct after insertion of the catheter the first blood sample (9 ml) was drawn (0 minutes), and after 15 minutes the subjects received their breakfast (20 % of the subjects' daily energy requirement). The daily energy requirements were calculated by multiplying the basal metabolic rate (BMR) by an activity index of 1.75. The BMR (MJ/day) was calculated according to the equations of Harris-Benedict [24]. At 45, 90 and 210 minutes the second, third and fourth blood samples were drawn. At 225 minutes the subjects received their lunch (30 % of the subjects' daily energy requirement) and at 255 and 300 minutes the fifth and sixth blood samples were drawn. During the test subjects were able to drink water ad libitum. The participants' 5-hour cortisol exposure was measured by calculating the area under the curve (AUC) (0-300 minutes).

Progressive maximal cycling test to volitional exhaustion

Each subject underwent a progressive maximal cycling test to volitional exhaustion for quantification of maximal workload (W_{max}) [25]. Subjects cycled on an electromagnetically braked bicycle ergometer (Lode, Groningen, The Netherlands) at 80 rpm, at a workload of 75 W for women and 100 W for men. After 5 minutes, the workload was increased every 2.5 minutes with 50 W, until subjects were no longer able to maintain a pedaling frequency above 80 rpm. Heart rate was monitored continuously throughout all exercise protocols (Polar s610). Verification that each subject actually achieved W_{max} consisted of meeting the criteria of achieving predicted maximal heart rate. The results of the progressive maximal cycling test were used to measure the resistance to elicit an exercise intensity of

90 % of each W_{max} (V) for the high intensity test under 4 mg dexamethasone.

High intensity test under 4 mg dexamethasone

The maximal cycling test and high intensity test under 4 mg dexamethasone [19] were separated by at least one week. Each subject received 4 mg dexamethasone 8 hours before the high intensity test. Each subject participated in the treatment, and no adverse reactions were reported. Subjects abstained from alcohol consumption and strenuous exercise 15 h before testing. At 8 a.m. after an overnight fast, an intravenous catheter for blood sampling was inserted into the forearm vein and the first blood sample was drawn for a baseline measurement (0 minutes), and after 5 minutes the subjects received their breakfast (20 % of the subjects' daily energy requirement). At 15 minutes the second blood sample was taken and the high intensity test was started. At 30 minutes the high intensity test ended and the third blood sample was taken. At 40, 50, 60 and 70 minutes, the fourth, fifth, sixth and seventh blood samples were taken.

The high intensity test consisted of 15 minutes of cycling. The initial 5 minutes served as a warming-up, during which each subject cycled at a workload of 40 % of his/her W_{max} (V), at 80 rpm. After the warming-up, the workload was increased with 50 % of his/her W_{max} (V) for 30 sec at 80 rpm, followed by a decrease of the workload with 50 % for 30 sec at 80 rpm. The increase and decrease of workload was repeated 10 times. After the third blood sample was drawn a 5 minutes cooling-down was performed at 30 % of W_{max} (V) at 60 rpm.

Subjects' negative feedback functioning (nmol/ml) was calculated as the cortisol concentrations of the baseline sample of the cortisol exposure measurement (0 min) minus the cortisol concentrations of the baseline sample after dexamethasone ingestion (0 min). When low amounts of cortisol were present after dexamethasone suppression, cortisol feedback was strong and when high

amounts of cortisol were present, cortisol feedback was decreased.

Subjects' HPA axis functioning in a challenged condition under 4 mg dexamethasone (%) was calculated as the difference between the highest cortisol concentrations after the high intensity test (30 or 40 minutes) and the cortisol concentration of the baseline sample after dexamethasone ingestion (0 min), subsequently divided by the cortisol concentrations of the baseline sample after dexamethasone ingestion (0 min). When low amounts of cortisol escaped during the high intensity test, cortisol feedback was strong and/or cortisol production was low, when high amounts of cortisol escaped, cortisol feedback was decreased and/or cortisol production was high [19].

Blood samples

Serum cortisol concentrations were measured using RIA (Buhlmann Laboratories) with an intra-assay coefficient variation of < 6%.

Statistical analysis

Differences between two groups were determined using t-tests. Relationships between dependent and independent variables were determined using simple linear and multiple regression models. All statistical tests were two-tailed, differences were considered significant at $p < 0.05$ and values are expressed as mean \pm standard deviation.

Results

Table 1 shows the characteristics of the male ($n = 91$) and female ($n = 103$) subjects. No significant differences were observed in age, BMI and hip circumference. The men showed significantly larger height, weight, waist circumference, waist-to-hip ratio (WHR) and maximal workload when compared to the women. The women showed significantly larger body fat percentage, and fat mass index when compared to the men. FMI (kg/m^2) was used to define body composition, and waist circumference (cm) as well as WHR was

Table 1: Characteristics of the male (n = 91) and female subjects (n = 103)

	Men	Women	P value ¹
Age (year)	28.1 ± 8.9	27.1 ± 8.6	0.84
Height (cm)	181.4 ± 6.9	169.0 ± 6.4	0.01
Body weight (kg)	82.4 ± 11.9	72.3 ± 12.5	0.01
Body mass index (kg/m ²)	25.1 ± 3.6	25.3 ± 4.1	0.68
Body fat percentage (%)	20.3 ± 9.6	32.0 ± 8.9	0.01
Fat Mass Index (FMI) (kg/m ²)	5.4 ± 3.2	8.5 ± 3.5	0.01
Waist circumference (cm)	88.1 ± 10.1	79.9 ± 10.3	0.01
Hip circumference (cm)	103.4 ± 6.6	105.3 ± 9.5	0.14
Waist hip ratio	0.85 ± 0.07	0.75 ± 0.06	0.01
Maximal workload (Wmax) (V)	292.8 ± 56	204.5 ± 47	0.01
5-hour cortisol exposure (nmol/L.min)	0.9 (10 ⁵) ± 0.2	1.3 (10 ⁵) ± 0.7	0.01
Baseline cortisol minus post dexamethasone cortisol concentrations (nmol/ml)	432.3 ± 103.9	624.7 ± 279.6	0.05
Percentage increase of cortisol concentrations after standardized high intensity test under 4mg dexamethasone (%)	17.7 ± 12.6	20.8 ± 19.9	0.05

Values are means ± sd.

¹ Differences between men vs. women (t-test)

used to define different patterns of body fat distribution. Additionally, the women showed significantly higher 5-hour cortisol exposure (nmol/L.min), baseline cortisol minus post dexamethasone cortisol concentrations (nmol/ml), and percentage increase of cortisol concentrations after standardized high intensity test under 4 mg dexamethasone (%) when compared to the men. Because of differences in HPA axis functioning the results for male and female subjects were analyzed separately. Subsequently, to study the relationship between HPA axis functioning and body fat distribution, the three parameters of HPA axis functioning were related to FMI, waist circumference, waist-to-hip ratio and possible confounders which have been related previously to HPA axis functioning, such as maximal workload, oral contraceptive intake and daily coffee intake.

Subjects' 5-hour cortisol exposure (nmol/L.min)

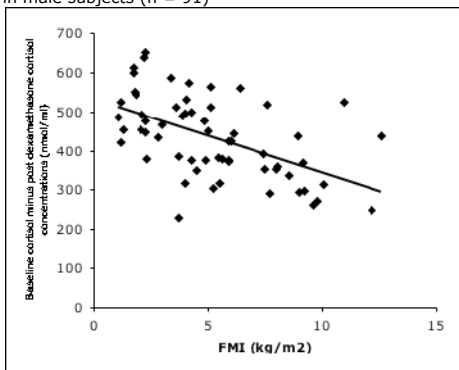
In male subjects, inverse relationships were observed between 5h cortisol exposure and BMI ($r = 0.43$, $p < 0.001$), FMI ($r = 0.55$, $p < 0.001$), waist circumference ($r = 0.23$, $p < 0.001$) and waist-to-hip ratio ($r = 0.27$, $p < 0.02$). Multiple regression analysis with 5h cortisol exposure as the dependent variable showed that only FMI significantly contributed to the explained variance.

In female subjects, inverse relationships were observed between 5h cortisol exposure and BMI ($r = 0.39$, $p < 0.001$), FMI ($r = 0.38$, $p < 0.001$), waist circumference ($r = 0.47$, $p < 0.001$), and waist-to-hip ratio ($r = 0.49$, $p < 0.001$). Positive relationships were observed between 5h cortisol exposure and maximal workload ($r = 0.32$, $p < 0.001$) and oral contraceptive use ($r = 0.39$, $p < 0.001$). Multiple regression analysis with 5h cortisol exposure as the dependent variable showed that only waist-to-hip ratio, maximal workload and oral contraceptive use significantly contributed to the explained variance ($r = 0.60$, $p < 0.001$).

Subjects' negative feedback functioning (nmol/ml)

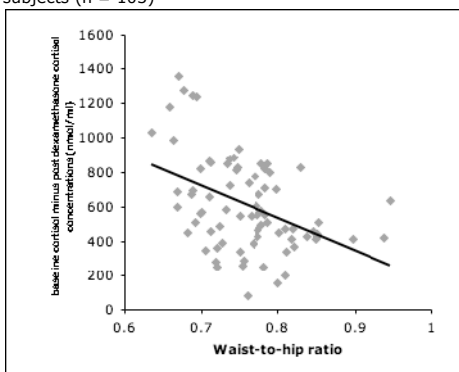
In male subjects, inverse relationships were observed between baseline cortisol minus post dexamethasone cortisol concentrations and BMI ($r = 0.43$, $p < 0.001$), FMI ($r = 0.53$, $p < 0.001$) (**figure 1a**), waist circumference ($r = 0.48$, $p < 0.001$) and waist-to-hip ratio ($r = 0.36$, $p < 0.002$). Multiple regression analysis with baseline cortisol minus post dexamethasone cortisol concentrations as the dependent variable showed that only FMI significantly contributed to the explained variance.

Figure 1a: relationship between baseline cortisol minus post dexamethasone cortisol concentrations (nmol/ml) and fat mass index (FMI) (kg/m²) ($r = 0.43$, $p < 0.001$) in male subjects ($n = 91$)



In female subjects, inverse relationships were observed between baseline cortisol minus post dexamethasone cortisol concentrations and BMI ($r = 0.34$, $p < 0.001$), FMI ($r = 0.32$, $p < 0.001$), waist circumference ($r = 0.41$, $p < 0.001$), and waist-to-hip ratio ($r = 0.43$, $p < 0.001$)(**figure 1b**). Positive relationships were observed between baseline cortisol minus post dexamethasone cortisol concentrations and maximum workload ($r = 0.30$, $p < 0.001$) and oral contraceptive use ($r = 0.43$, $p < 0.001$).

Figure 1b: relationship between baseline cortisol minus post dexamethasone cortisol concentrations (nmol/ml) and waist-to-hip ratio ($r = 0.43$, $p < 0.001$) in female subjects ($n = 103$)



Multiple regression analysis with baseline cortisol minus post dexamethasone cortisol concentrations as the dependent variable showed that only waist-to-hip ratio, maximum workload and oral

contraceptive use significantly contributed to the explained variance ($r = 0.59$, $p < 0.001$).

Subjects' HPA axis functioning in a challenged condition under 4 mg dexamethasone (%)

In male subjects, an inverse relationship was observed between percentage increase of cortisol concentrations after standardized high intensity test under 4 mg dexamethasone and waist circumference ($r = 0.21$, $p < 0.10$).

In female subjects, inverse relationships were observed between percentage increase of cortisol concentrations after standardized high intensity test under 4 mg dexamethasone and BMI ($r = 0.19$, $p < 0.05$), waist circumference ($r = 0.23$, $p < 0.04$), and waist-to-hip ratio ($r = 0.21$, $p < 0.05$). Multiple regression analysis with percentage increase of cortisol concentrations after standardized high intensity test under 4 mg dexamethasone as the dependent variable showed that only waist-to-hip ratio significantly contributed to the explained variance.

In both male and female subjects, a positive relationship was observed between 5-hour cortisol exposure (nmol/ml) and baseline cortisol minus post dexamethasone cortisol concentrations (nmol/ml) ($r = 0.68$, $p < 0.001$ and $r = 0.86$, $p < 0.001$).

In male subjects, strong positive relationships were observed between FMI and waist circumference ($r = 0.85$, $p < 0.001$), as well as waist-to-hip ratio ($r = 0.70$, $p < 0.001$).

Discussion

The objective of our study was to relate HPA axis functioning and HPA feedback functioning to body fat distribution in normal weight to obese subjects. We therefore tested the relationship between several parameters of HPA axis functioning, body composition and body fat distribution measured as waist circumference and waist-to-hip ratio. Waist circumference and waist-to-hip

measurements were used as surrogates for body fat distribution measurements, as anthropometric measurements have been shown to predict total and abdominal fat content [20-22].

Firstly, we investigated the relationships between cortisol exposure measured over 5 hours, body composition and body fat distribution. In accordance to the literature [3, 4, 10-12] we observed that cortisol exposure was related to a higher FMI in men and a higher waist-to-hip ratio in women. This sex difference is likely to be caused by the fact that men tend to progressively increase visceral fat depots with increasing adiposity, while women tend to develop different types of body fat distribution (visceral or peripheral) [26], which is supported by the strong relationship between FMI and waist circumference as well as waist-to-hip ratio in men in our study. This suggests that in both men and women low cortisol exposure is related to visceral fat distribution. Low cortisol exposure is a sign of a decreased HPA axis activity, which is for instance also observed in subjects with chronic stress and depression [4, 27]. As expected and in concordance with previous studies we observed that in the women HPA axis functioning was related to oral contraceptive use [28, 29], but also to baseline exercise levels. These relationships were however independent of the relationship between HPA axis functioning and visceral fat disposition. A limitation of the present study is that the women were examined regardless of the phase of their menstrual cycle, as previous studies have shown differences in basal cortisol concentrations between the luteal and follicular phase, which is about 3% [28]. The women being in different phases of the menstrual cycle can thus not explain the differences in basal cortisol concentrations between women with a high or low waist-to-hip ratio. However, for future studies information on the menstrual cycle should be taken into account.

Secondly, we investigated the relationships between HPA axis feedback functioning, body composition and body fat distribution, measured through a 4 mg dexamethasone suppression test. In accordance to part of the literature [17, 18] we observed that reduced negative feedback of the HPA axis in a non-challenged condition (i.e. high amounts of cortisol present after dexamethasone suppression) was related to a higher FMI in men and a higher waist-to-hip ratio in women. Other studies however have shown an inverse [4, 16], no relationship [3, 5, 6, 10] or sex specific associations [16]; this inconclusiveness may result from different dosages of dexamethasone, as they used smaller dosages of dexamethasone (0.05 to 0.25 mg). Low dosages of dexamethasone may not be potent enough to detect possible differences in HPA axis feedback functioning, as Ljung et al 1996 [17] observe no differences in inhibition when using low dosages (0.05 and 0.125 mg) but did observe differences when using a higher dosage (0.5 mg). We therefore suggest that a protocol that includes the range of dexamethasone dosage of 0.5 to 4 mg is necessary to detect subtle differences in HPA axis feedback functioning. The relationship between 5-h cortisol exposure and the suppression of non-challenged cortisol levels by dexamethasone that was observed in both men and women, strongly suggest that a diminished negative HPA axis feedback is one of the mechanisms that is responsible for the relatively low 5-h cortisol exposure in visceral obesity.

Thirdly, we investigated the relationship between HPA axis functioning in a challenged condition under 4 mg dexamethasone, body composition and body fat distribution, measured through a high intensity test under 4 mg dexamethasone [19]. In accordance to previous studies that only measured lean subjects [19, 30] a significant sex difference was observed in HPA axis functioning in a challenged condition under 4 mg dexamethasone. Moreover, we observed differences in HPA axis

functioning i.e. relatively lower cortisol levels after a challenged condition under dexamethasone that related to a higher waist circumference in men and a higher waist-to-hip ratio in women. This is in contrast with the non-challenged condition under dexamethasone, where relatively higher cortisol levels after dexamethasone suppression were related to a higher fat mass index in men and a higher waist-to-hip ratio in women. Several mechanisms are hypothesized that could underlie the discrepancy between the challenged and the non-challenged condition. Since dexamethasone is unable to cross the blood-brain barrier [31], cortisol levels after dexamethasone suppression reflect negative feedback functioning only at the level of the pituitary. Therefore, it cannot be excluded that the negative feedback mechanisms at the level of the pituitary and the hypothalamus are distinctly influenced by visceral obesity, and that visceral obesity is associated with an increased negative feedback at the level of the hypothalamus, which in the current study became predominant in the exercise-challenged condition. Alternatively, it can be hypothesized that the decreased negative feedback signaling in visceral obesity, as shown in our non-challenged condition and confirmed by others [17, 18], is associated with a decreased responsiveness of the HPA axis to (physical) stress. Under normal, dexamethasone-free conditions, where differences in feedback sensitivity may play a role, the increased cortisol

response to stress [3-6, 9, 14] could then be attributed to the decreased efficacy of cortisol to suppress its own secretion.

Taken together all our observations pointed into the same direction, as we observed that in lean to overweight subjects, low cortisol exposure, enhanced negative feedback, and altered HPA axis functioning in a challenged condition under 4 mg dexamethasone is related to waist circumference in men and to waist-to-hip ratio in women. Our results are thus consistent with previous work that suggests a deregulation of the HPA axis in humans with relative more visceral fat. It is however still not clear whether the HPA axis is causally or co-incidentally related to (visceral) obesity, which only can be determined using a longitudinal study. From our results, we conclude that disturbance of HPA axis functioning under basal and challenged conditions is related to visceral fat accumulation.

Acknowledgements

We want to thank our subjects for their participation in this study. We gratefully thank Loek Wouters and Jose Sulon for their assistance. FR (supervised by AN and MW) carried out the study, collected and analyzed the data, and wrote the largest part of the manuscript. SL and JB reviewed the manuscript. Planning, processing the results and writing the manuscript were done under general supervision by AN and MW. The authors had no conflict of interest.

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Chapter 9

Acute stress-related changes in eating in the absence of hunger

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Obesity 2009 January; 17 (1): 72-7

Abstract

Background: obesity results from chronic deregulation of energy balance, which may in part be caused by stress.

Objective: to investigate the effect of acute and psychological stress on food intake, using the eating in the absence of hunger paradigm, in normal- and overweight men and women, while taking dietary restraint and disinhibition into account. In 129 subjects (BMI = 24.5 ± 3.4 kg/m² and age = 27.6 ± 8.8 y), scores were determined on the Three Factor Eating Questionnaire (dietary restraint = 7.2 ± 4.4 ; disinhibition = 4.5 ± 2.6 ; feeling of hunger = 3.9 ± 2.6) and State-Trait Anxiety Inventory (trait score = 31.7 ± 24.2). In a randomized crossover design, the 'eating in absence of hunger' protocol was measured as a function of acute stress versus a control task and of state anxiety scores.

Results: energy intake from sweet foods (708.1 vs. 599.4 kJ, $p < 0.03$) and total energy intake (965.2 vs. 793.8 kJ, $p < 0.01$) were significantly higher in the stress compared to the control condition. Differences in energy intake between the stress and control condition were a function of increase in state anxiety scores during the stress task ($R^2 = 0.05$, $p < 0.01$). This positive relationship was stronger in subjects with high disinhibition scores ($R^2 = 0.12$, $p < 0.05$). Differences in state anxiety scores were a function of trait anxiety scores ($R^2 = 0.07$, $p < 0.05$).

Conclusion: acute psychological stress is associated with eating in the absence of hunger, especially in vulnerable individuals characterized by disinhibited eating behavior and sensitivity to chronic stress.

Introduction

The prevalence of overweight and obesity has increased worldwide to epidemic proportions [1]. Obesity results from a chronic deregulation of energy balance, with energy intake exceeding energy expenditure, leading to the storage of the excessive energy as fat [2]. This chronic deregulation of energy balance may be caused, in part, by stress, since in Western society high levels of ambient stress are abundantly present [3]. Several human studies in laboratory settings indicate that, in adults, acute stress may influence energy intake [3-7]. Acute stress alters food preference, eating frequency, and amount of energy intake [3, 5]. During stress, food preference is altered towards intake of sweet foods and foods with saturated fats, next to an increase in the intake frequency and amount of food intake [4, 6, 7]. However, not all studies that investigated the relationship between energy intake and stress have yielded conclusive results [8, 9]. Stress may also result in a decrease in energy intake [8, 9]. These individual differences in response to stress may relate to eating behavior characteristics [4-7], as are determined using the Three Factor Eating Questionnaire (TFEQ). The TFEQ assesses three factors involved in eating behavior: dietary restraint; disinhibition; and hunger [10]. An increase in energy intake during stress has been found in individuals with high scores on dietary restraint and/or disinhibition [4-7].

A possible technique to study the effect of stress on food intake is the 'Eating in the absence of hunger' paradigm [11]. Eating in the absence of hunger is the behavioral phenotype in which individuals eat when exposed to large portions of palatable foods in the absence of hunger. The paradigm has primarily been used in children [11] and was thought to resemble disinhibited eating patterns observed in adults [11]. Therefore, the objective of our study was to investigate the effect of acute psychological stress on food intake, using the eating in the

absence of hunger paradigm, in normal- and overweight men and women, while taking dietary restraint and disinhibition into account.

Subjects and methods

Subjects

The study was approved by the Medical Ethical Committee of Maastricht University, and informed, written consent was obtained from all subjects. Healthy, medication-free, non-smoking men ($n = 65$) and women ($n = 65$), 18 to 45 y and with a BMI of 20 to 35 kg/m² were recruited to complete all phases of this study. A medical history was obtained from each subject before entry into the study, and exclusion criteria were chronic illness, depression, and a history of eating disorders or current dieting. From 130 subjects recruited, 65 men and 64 women completed the study, and 1 subject dropped out before the actual start of the study. Subjects were not told the real purpose of the study but instead were given a cover story on the effect of activity on leptin concentrations.

Experimental design

The study was performed in a within-subjects, randomized, crossover design. Each subject reported to the laboratory on 3 occasions for: 1) measurement of anthropometry and subject characterization; 2) measurement of 'eating in absence of hunger' protocol in combination with a stress task; and 3) measurement of 'eating in absence of hunger' protocol in combination with a control task.

The anthropometry measurements consisted of body weight, height, waist and hip circumference determination. Subjects also completed the State-Trait Anxiety Inventory (STAI trait) and the Three Factor Eating Questionnaire (TFEQ), which assesses restraint, disinhibition, and feeling of hunger.

'Eating in absence of hunger' was measured as a function of acute stress versus control. Analog scales (VAS) that measure hunger and satiety were used to

determine if subjects ate beyond fullness. A mental arithmetic task with sums that subjects could (control task) or could not solve (stress task) was used as a stress manipulation. To investigate whether the stress condition inflicted psychological or physiological changes, we used the profile of mood states (POMS) and state-anxiety (STAI state) questionnaires, as well as heart rate and blood pressure measurements. Before the eating in absence of hunger measurements, subjects were made familiar with the VAS, POMS, and STAI state questionnaires and were trained to complete them routinely, in order not to disturb the actual activity they were involved in during the measurements.

Measurements

Anthropometry

Measurements were obtained in the morning after voiding the bladder and before breakfast. Body weight (Sauter D7470, Ebingen, Germany) and height (Seca, model 220, Hamburg, Germany) were measured to the nearest 0.01 kg and 0.1 cm, respectively. BMI (kg/m^2) was calculated as body weight (kg) divided by height (m) squared. The waist circumference was measured to the nearest 0.1 cm at the site of the smallest circumference between the rib cage and the iliac crest, whereas the hip circumference was measured to the nearest 0.1 cm at the level of maximum extension of the buttocks. The waist and hip circumference measurements were obtained with subjects in the standing position [12].

Eating behavior

Eating behavior was analyzed using a validated Dutch translation of the TFEQ [10, 13]. The TFEQ measures three components of eating behavior. Dietary restraint reflects the extent to which individuals attempt to cognitively control their food intake. Disinhibition of restraint reflects loss of control over eating in response to the presence of palatable food or other disinhibiting stimuli, such as emotional distress. The third factor measures the subjective feeling of hunger.

Based upon the median for the TFEQ scores in the South of the Netherlands, subjects were characterized as unrestrained when dietary restraint scores were < 9 , and as restrained when scores were ≥ 9 . Subjects were characterized as having low disinhibition when disinhibition scores were < 5 , and as having high disinhibition when scores were ≥ 5 [13].

Visual Analog Scales (VAS)

Aspects of appetite were assessed using 100 mm visual analog scales (VAS) with questions about feelings of hunger, satiety, thirst, and desire to eat. Opposing extremes of each feeling were described at either end of the 100-mm horizontal line, and subjects marked the line to indicate how they felt at that moment [14]. Completion of the VAS questionnaire took our experienced subjects about 0.5 minute. During the 'eating in absence of hunger' protocol, appetite profiles were assessed three times: before lunch, 5 minutes after lunch, and 30 minutes after the stress or control task.

Profile of Mood States (POMS)

Aspects of mood were assessed using the Dutch translation of the Profile of Mood States (POMS) [15]. This questionnaire contains 70 adjectives that are rated on a five-point scale, anchored by "much like this" to "much unlike this" and is divided into five subscales (depression, tension, confusion, fatigue and anger), each scoring a maximum of 35 points. An increase in POMS scores is associated with a worsening in mood, except in the case of "vigor" [15]. Completion of the POMS questionnaire took our experienced subjects only 3 minutes. During the 'eating in absence of hunger' protocol, mood profiles were assessed five times: 5 minutes after lunch, and at 0, 10, 20 and 30 minutes after the stress or control task.

State-Trait Anxiety Inventory (STAI)

Anxiety was assessed using the Dutch translation of the State-Trait Anxiety Inventory (STAI) [16]. This questionnaire consists of two subscales: the trait scale (anxiety-trait) and the state scale

(anxiety-state). The trait scale refers to individual differences in the likelihood that a person would experience state anxiety in a stressful situation, and thereby refers to chronic feelings of anxiety. The state scale refers to the transitory emotional response involving unpleasant feelings of tension and apprehensive thoughts. Both scales are composed of 20 questions rated on a four-point scale, ranging from "much like this," to "much unlike this" and require that subjects describe how they feel generally, on the anxiety-trait scale, and how they feel at a specific moment, on the anxiety-state scale [16]. Each subscale of the questionnaire can score a maximum of 80 points and an increase in STAI trait or state scores is associated with an increase in anxiety. Completion of the STAI state questionnaire took our experienced subjects only 2 minutes. At the start of the experiment subjects filled in the trait scale and during the 'eating in absence of hunger' protocol, state anxiety was assessed five times: 5 minutes after lunch, and at 0, 10, 20 and 30 minutes after the stress or control task.

Stress/control task

During the 'eating in the absence of hunger' protocol subjects received a mental arithmetic task with sums that they could or could not solve (control versus stress task). Subjects were told to solve the sums within the time limits specified by the assignment. The control and stress tasks were presented in random order to the subjects. The arithmetic task was a modified and improved version of the computer program previously described and validated by Peters et al. 1998 [17].

'Eating in the absence of hunger' protocol

The 'eating in the absence of hunger' protocol consisted of consumption of a standard lunch, followed by completion of a stress or control task, and was concluded with half an hour in which subjects had free-access to snacks. The 'eating in the absence of hunger' protocol was started by completing a VAS, and followed by a standard lunch to minimize the influence of hunger on the subjects'

snack intake. The lunch contained 30% of the subjects' daily energy requirement and consisted of typical Dutch lunch products, namely brown bread (1.0 MJ per 100 g; en% C/P/F: 74/15/11) with young cheese (1.6 MJ per 100 g; en% C/P/F: 0/26/74), as well as ab libitum tea or water consumption. The daily energy requirements were calculated by multiplying the basal metabolic rate (BMR) by an activity index of 1.75. The BMR (MJ/day) was calculated according to the equations of Harris-Benedict [18]. Subjects were given 15 minutes to consume the lunch and were instructed to eat until they were satiated. Immediately after lunch, subjects completed POMS and state anxiety questionnaires and completed a stress or control task. After completing VAS, POMS and state anxiety questionnaires, which took about 5 minutes, subjects were retained in the research room for half an hour in which they were allowed to relax and read magazines. During this half hour the subjects were presented a tray with containers holding generous pre-weighed portions of sweet snack foods, including 100 g chocolate (2.2 MJ per 100 g) and 200 g fruit-chew candy (1.5 MJ per 100 g), as well as salty snack foods that included 100 g potato chips (2.4 MJ per 100 g), 50 g pretzels (2.1 MJ per 100 g), and 200 g nuts (2.7 MJ per 100 g). The subjects were instructed to eat as much as they wished of any of the foods. Every 10 minutes during the half hour, subjects completed POMS and state anxiety questionnaires. At the end of the half hour the subjects completed VAS, POMS and state questionnaires one more time. After the subjects left, each of the containers holding the snack foods were weighed to determine food choice and the amount eaten. Before and after completing the stress or control tasks, heart rate (bpm), and systolic and diastolic blood pressure (mmHg) were measured using an automated device (WelclAllyn OSZ 5 easy, Spreidel and Keller GmbH and Co, KG, Jungingen, Germany).

Statistical analysis

Differences between men and women, as well as treatments (control and stress), were determined using unpaired t-tests. Differences over time and between treatments (control and stress) were determined using two-factor ANOVA with repeated measures. To analyze whether the amount of stress experienced during the stress task predicted the difference in energy intake between the stress and control condition, a simple linear regression model was used, depicting the differences in energy intake between the stress and control condition (Δ energy intake) as the dependent variable and the differences in state anxiety scores immediately after compared to before the stress task (Δ state anxiety scores) as the independent variable. The relation between dependent (Δ state anxiety scores) and independent (trait anxiety scores) variable was determined using a linear regression model. All statistical tests were two-sided. Differences were considered significant at $p < 0.05$, and values were expressed as mean \pm standard deviation.

Results

Subject characteristics

The characteristics of the subjects are summarized in **table 1**. Participants were 18 to 48 years old, and their BMI ranged from 19.8 to 34.8 kg/m². Men had a predictably higher height, body weight, and waist circumference than women. Women had higher dietary restraint, disinhibition, and feeling of hunger scores than men. The age, BMI, hip circumference and trait anxiety scores did not differ between the sexes. We did not analyze the results of men and women separately, because no significant differences were found between men and women in VAS, POMS, and state anxiety scores, or in energy intake during the eating in absence of hunger protocol (data not shown).

Table 1: Characteristics of subjects (n = 129)

	Men (n = 65)	Women (n = 64)	P value ¹
Age (year)	28.1 \pm 8.9	27.1 \pm 8.6	0.84
Height (cm)	181.2 \pm 7.5	168.7 \pm 6.2	0.01
Body weight (kg)	79.8 \pm 10.2	69.9 \pm 10.4	0.01
Body mass index (kg/m ²)	24.4 \pm 3.3	24.6 \pm 3.6	0.74
Waist circumference (cm)	85.7 \pm 9.5	78.8 \pm 7.9	0.01
Hip circumference (cm)	102.2 \pm 6.2	104.7 \pm 8.5	0.12
Trait anxiety score	31.6 \pm 6.7	31.9 \pm 6.7	0.77
Dietary restraint score	5.8 \pm 3.9	8.8 \pm 4.4	0.01
Disinhibition score	3.4 \pm 1.7	5.6 \pm 2.8	0.01
Feeling of hunger score	3.5 \pm 2.5	4.4 \pm 2.6	0.03

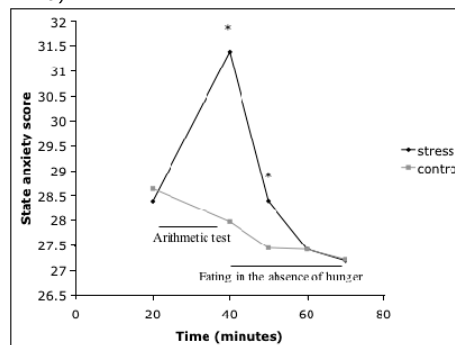
Values are means \pm sd.

¹ Differences between men vs. women (t-test)

State anxiety scores

The changes in state anxiety scores over time in the stress and control condition are depicted in **figure 1**. The state anxiety scores were significantly higher in the stress condition compared to the control condition, immediately (31.3 \pm 8.8 vs. 27.9 \pm 6.4, $p < 0.001$) and 10 minutes (28.4 \pm 7.7 vs. 27.3 \pm 6.9, $p < 0.05$) after receiving the stress or control task (figure 1). After receiving the stress task, state anxiety scores increased significantly, while in the control condition no significant changes were observed (4.2 \pm 5.7 vs. -0.8 \pm 0.3, $p < 0.001$) (figure 1).

Figure 1: Changes in state anxiety scores (means) over time in the stress and control condition in all subjects (n = 129)



* $p < 0.05$ for differences in time in the stress compared to the control condition (ANOVA repeated measures)

POMS scores

The POMS scores were significantly higher in the stress condition compared to the control condition, namely immediately after receiving the stress task in four out of five POMS categories, which are depression (15.7 \pm 4.1 vs. 14.2 \pm 3.5, p

< 0.05), tension (14.7 ± 5.1 vs. 12.7 ± 3.7 , $p < 0.05$), confusion (15.9 ± 4.3 vs. 14.7 ± 3.1 , $p < 0.05$) and anger (14.6 ± 4.8 vs. 12.6 ± 3.3 , $p < 0.05$). After receiving the stress task POMS scores increased significantly, while in the control condition no significant alterations were shown in the categories depression (0.5 ± 3.2 vs. -1.0 ± 2.5 , $p < 0.05$), tension (1.8 ± 3.9 vs. -0.7 ± 2.5 , $p < 0.05$), confusion (0.9 ± 3.2 vs. -0.7 ± 2.4 , $p < 0.05$) and anger (1.7 ± 3.9 vs. -0.4 ± 1.8 , $p < 0.05$).

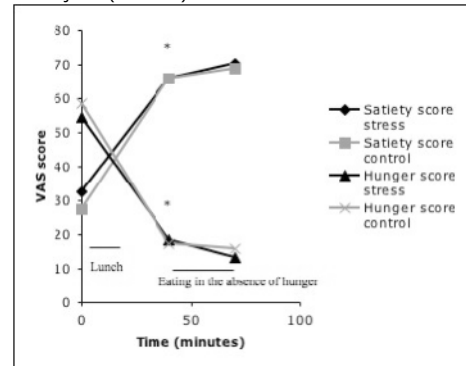
Heart rate and blood pressure measurements, immediately before and directly after completion of the arithmetic task, showed no changes over time.

Additionally, no changes were seen between the stress and control condition in heart rate (70.3 ± 13 vs. 72.3 ± 12.3 bpm, $p < 0.86$) and systolic (123.7 ± 14.1 vs. 125.7 ± 12.9 mmHg, $p < 0.45$) and diastolic (73.5 ± 10.2 vs. 72.3 ± 12.3 mmHg, $p < 0.89$) blood pressure.

VAS hunger and satiety scores

During the lunch the average energy intake for men was 4.0 ± 0.5 MJ and for women 3.5 ± 0.4 MJ. The changes in satiety and hunger scores over time in the stress and control condition are depicted in **figure 2**. No significant differences in satiety scores were seen between the stress and control condition before lunch (32.5 ± 21.1 vs. 27.6 ± 15.8), directly after lunch (65.6 ± 17.7 vs. 65.9 ± 18.3) and after the eating in absence of hunger protocol (70.7 ± 16.9 vs. 69.1 ± 18.5). No significant differences in hunger scores were seen between the stress and control conditions before lunch (54.2 ± 22.9 vs. 58.5 ± 19.8), directly after lunch (18.6 ± 13.9 vs. 17.5 ± 13.8) and after the eating in absence of hunger protocol (13.5 ± 19.5 vs. 15.8 ± 13.7). A significant increase in satiety scores and a significant decrease in hunger scores were seen over time in both conditions.

Figure 2: Changes in satiety and hunger scores (means) over time in the stress and control condition in all subjects (n = 129)

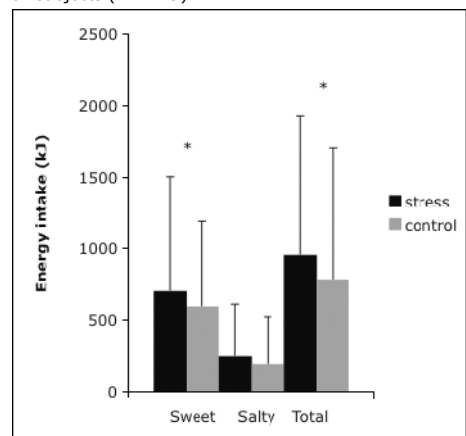


* $p < 0.05$ for differences in time in both the stress and control condition (ANOVA repeated measures)

Differences in energy intake in stress/control condition

The differences in energy intake (kJ) during stress and control condition are depicted in **figure 3**. Significant differences were found between the stress and control conditions in energy intake from sweet snack foods (708.1 ± 798.8 vs. 599.4 ± 734.4 kJ, $p < 0.03$) and total energy intake (965.2 ± 970.6 vs. 793.8 ± 912.5 kJ, $p < 0.01$).

Figure 3: Differences in means (\pm sd) of energy intake of snack foods (kJ) during stress and control condition in all subjects (n = 129)

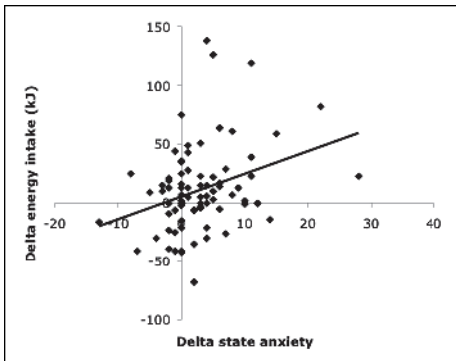


* $P < 0.05$ for stress versus control condition, t-test

Figure 4 depicts the differences in energy intake of snack foods between the stress and control conditions (Δ energy intake) as a function of the differences in STAI state scores between immediately after

compared to before the stress task (Δ state anxiety scores). A positive relationship was shown between Δ energy intake (kJ) and Δ state anxiety scores ($R^2 = 0.05$, $p < 0.01$). Apart from that, no relationship between Δ energy intake (kJ) and trait anxiety scores was present.

Figure 4: differences in energy intake of snack foods (kJ) between the stress and control condition (Δ energy intake) as a function of the differences in state anxiety scores between immediately after compared to before the stress task (Δ state anxiety scores) in all subjects ($n = 129$)



$R^2 = 0.05$, $p < 0.01$

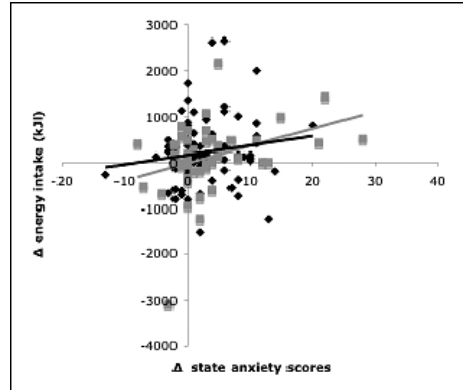
Differences in energy intake in stress/control conditions combined with eating behavior

First, linear regression analyses with Δ energy intake (kJ) as the dependent variable and Δ state anxiety scores as the independent variable were performed for subjects with low dietary restraint scores ($R^2 = 0.05$, ns) and high dietary restraint scores ($R^2 = 0.01$, ns).

Second, a multiple linear regression analysis was performed with Δ energy intake (kJ) as the dependent variable and Δ state anxiety scores, disinhibition scores and interaction between the Δ state anxiety scores and the disinhibition scores, as the three independent variables. A significant interaction between Δ state anxiety scores and disinhibition scores was found, indicating that the explained variance of Δ of energy intake by Δ state anxiety scores was different in subjects, namely with low or high disinhibition scores. Therefore, **figure 5** depicts the Δ energy intake (kJ) as a function of Δ state anxiety scores plotted for two groups of subjects namely

those with a low disinhibition score ($R^2 = 0.02$, $p = 0.23$) and a high disinhibition score ($R^2 = 0.12$, $p < 0.05$).

Figure 5: differences in energy intake (kJ) between the stress and control condition (Δ energy intake) as a function of the differences in state anxiety scores between immediately after compared to before the stress task (Δ state anxiety scores), plotted for a low disinhibition scores group (in black, $R^2 = 0.02$, $p = 0.23$) and a high disinhibition scores group (in grey, $R^2 = 0.12$, $p < 0.05$), in all subjects ($n = 129$)



Linear regression model with Δ state anxiety scores

A linear regression model showed a positive relationship between the Δ state anxiety scores as the dependent variable and the trait anxiety scores as the independent variable ($R^2 = 0.07$, $p < 0.01$).

Discussion

The objective of our study was to investigate the effect of acute psychological stress on food intake, using the eating in the absence of hunger paradigm, in normal- and overweight men and women, taking dietary restraint and disinhibition into account. Acute psychological stress, indicated as an increase in state anxiety and POMS scores, was related to an increase in energy intake in the absence of hunger in adults. Moreover, subject specific features including higher trait anxiety scores, were related to a larger increase in state anxiety scores during the stress condition.

With these results, our study adds to the growing literature on the effect of stress on energy intake. To investigate whether

the stress condition induced stress, we measured anxiety (STAI questionnaire) [16] mood profiles (POMS questionnaire) [15] over time and heart rate and blood pressure. We showed an increase in anxiety and a deterioration of mood during the stress condition, which confirms previous findings [19, 20] and thereby confirms stress induction. The changes in anxiety and mood after receiving the stress task compared to the control were observed immediately and after 10 minutes, which indicates acute psychological stress induction that has been shown in previous studies using this particular stress task [17]. Completion of the various questionnaires could be considered a potential stressor. Nevertheless, the number of questionnaires was identical in both conditions, and the control condition showed a decrease in state anxiety and POMS scores, indicating no stressful experience.

However, contrary to previous studies [17, 21], no differences were found in heart rate and blood pressure over time and between the stress and control condition. A possible explanation may be the frequency of the measurements; we measured at two time points (before and after stress/control task), thereby possibly running the risk of missing some changes in heart rate and blood pressure. For future experiments we recommend to focus on heart rate variability [22].

To determine whether subjects were eating beyond fullness, satiety and hunger scores (VAS questionnaire) were measured over time, and in both conditions (stress and control) an increase in satiety scores and a decrease in hunger scores over time were shown. These satiety and hunger scores indicate a state of satiety after the consumption of the lunch, as confirmed by previous research on *ab libitum* food intake [23], and suggested that possible snack intake was not hunger based. No differences in satiety and hunger scores were seen between the stress and control conditions, suggesting that acute psychological stress

did not affect appetite, which is contrary to other studies that did show a decrease in hunger ratings after stress [5]. The contradiction may be due to the fact that in our experiment, the subjects were fed before the exposure to stress, while in the other studies subjects were hungry during stress exposure. The observed food intake, ingested in the absence of hunger after stress exposure, hardly led to further decreases in hunger, underscoring that this may not be a homeostatic mechanism, yet rather a non-homeostatic regulatory mechanisms involved in feeding behavior, such as reward [9, 24].

In accordance with the literature [4-7], we found that (acute psychological) stress altered food preference and amount of energy intake; namely, it increased intake of sweet foods and increased total amount of energy intake. Our study is the first to show that, in adults, eating beyond fullness ('eating in absence of hunger') is affected by acute stress. Before, the paradigm of eating in the absence of hunger only has been used in children [11]. Here it appears to be a sensitive paradigm in adults as well. Since there is no literature on the validity of the paradigm in adults, the paradigm should be used in different studies in order to show reproducibility and sensitivity. A linear regression showed that differences in energy intake between the stress and control conditions was a function of the differences in state anxiety scores between immediately after compared to before the stress task. These results imply that subjects who experienced a greater amount of acute stress had the largest increase in energy intake. Here trait anxiety scores did not play a role, since these did not contribute to the explanation of Δ energy intake.

Taking subject characteristics into account, it has been suggested that dietary restraint may account for the differences in reaction to stress with respect to energy intake. In contrast to other studies [5, 6], our study did not show a relationship between energy intake, stress and dietary restraint.

Instead, an effect of disinhibition was shown. The relationship between differences in energy intake between the stress and control conditions and differences in state anxiety scores between immediately after compared to before the stress task, was only significant in subjects with a high disinhibition scores, when compared to the subjects with a low disinhibition scores. Thus, disinhibition scores appeared to be a subject specific feature that contributed to increased energy intake during acute stress.

In addition, a linear regression analysis showed that subjects who experienced acute stress had higher trait anxiety scores. These results implicate that a higher likelihood to experience state anxiety in a stressful situation [16],

contributes to a higher sensitivity to acute stress from the stress task. Therefore, we conclude that acute psychological stress is associated to eating in absence of hunger, especially in vulnerable individuals characterized by disinhibited eating behavior and sensitivity to chronic stress.

Acknowledgements

We want to thank our subjects for their participation in this study. FR (supervised by AN and MW) carried out the study, collected and analyzed the data, and wrote the largest part of the manuscript. SL and JB reviewed the manuscript. Planning, processing the results and writing the manuscript were done under general supervision by AN and MW. The authors had no conflict of interest.

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Chapter 10

Hyperactivity of the HPA axis is related to dietary restraint in normal weight women

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Physiology and Behavior 2009 February 16; 96 (2): 315-9

Abstract

Objective: to investigate the relationship between hypothalamus/pituitary/adrenal (HPA) axis functioning and dietary restraint in normal weight (BMI between 20 and 25 kg/m²) men and women.

Methods: we assessed in 38 men and 38 women HPA axis functioning, through measuring 5-hour cortisol exposure and cortisol feedback functioning through a dexamethasone (4 mg) suppression test. Eating behavior was assessed through the Three Factor Eating Questionnaire and body composition through hydro densitometry and deuterium dilution method.

Results: no relationship between HPA axis functioning and dietary restraint was found in men. Normal weight women with a restraint score ≥ 9 showed increased cortisol concentrations over a 5-hour time period, increased cortisol concentrations after a dexamethasone (4 mg) suppression test, higher BMI, and higher body fat percentage, when compared to women with a restraint score < 9 . Moreover, a positive relationship was found between cortisol concentrations over a 5-hour time period and dietary restraint in combination with the disinhibition score ($R^2=0.23$, $p<0.001$).

Conclusion: in normal weight women hyperactivity of the HPA-axis is related to dietary restraint especially in combination with disinhibition.

Introduction

In order to either prevent weight gain or induce weight loss, people attempt to restrict food intake through controlled eating behavior [1]. Eating behavior characteristics can be determined using the Three Factor Eating Questionnaire (TFEQ) [2]. The TFEQ assesses 3 factors involved in eating behavior: dietary restraint, disinhibition and feeling of hunger. Dietary restraint reflects the extent to which individuals attempt to cognitively control their food intake. Disinhibition or inhibition of restraint reflects individual differences in the extent to which release from the cognitive suppression of eating occurs in response to the presence of palatable food or other disinhibiting stimuli, such as emotional distress. The third factor refers to the general feeling of hunger [2]. Studies have shown behavioral, metabolic and endocrinological differences between normal weight women with a low score and a high score on dietary restraint. Normal weight dietary restrained women showed behavioral and metabolic differences such as lower self-reported energy consumption, lower energy expenditure, higher body mass index (BMI) and higher body fat percentage when compared to the dietary unrestrained women [3, 4]. Furthermore, a number of studies showed that normal weight dietary restrained women showed endocrinological differences such as lower post-meal insulin levels [5-7] and increased cortisol levels compared to dietary unrestrained women [8-11]. The increased cortisol levels in dietary restrained women may be secondary to the occurrence of psychological stress, since dietary restraint is experienced as a demanding task [8-11]. However, not all studies investigating the association between cortisol levels and dietary restraint have yielded conclusive results, since some studies found a positive association in the dietary restrained women [8-11], whereas others did not [4-6]. This inconclusiveness possibly resulted from study limitations such as cortisol

measurements at only one (sometimes unspecified) time point during the day or several time points during the night. Moreover, all cortisol measurements were done only under basal conditions. However basal cortisol levels also depend on cortisol conversion, and therefore cortisol measurements under basal condition do not reflect hypothalamic/pituitary/adrenal (HPA) axis functioning. The ability of cortisol to suppress its own secretion, mediated by the glucocorticoid receptor in both the hypothalamus and pituitary (negative feedback) is an important determinant of the HPA axis functioning [12]. However, to our knowledge no studies reported adequately on the possible association between HPA axis functioning, measuring both cortisol exposure and cortisol feedback functioning, and dietary restraint. Consequently, the objective of our study was to investigate the relationship between hypothalamus/pituitary/adrenal (HPA) axis functioning and dietary restraint in normal weight (BMI between 20 and 25 kg/m²) men and women.

Participants and methods

Participants and protocol

The study was approved by the Medical Ethical Committee of the Maastricht University, and a written consent was obtained from all participants. Healthy, medication-free, non-smoking men (n = 40) and women (n = 40) aged between 18 to 45 y and with a BMI between 20 and 25 kg/m² were recruited to complete all phases of this study. A medical history was obtained from each participant before entry into the study and exclusion criteria were chronic illness, depression, and a history of eating disorders or current dieting. From 80 participants recruited, 38 men and 38 women completed the study, 4 participants dropped out before the actual start of the study. From each participant we obtained: 5-hour cortisol exposure, a progressive maximal cycling test to volitional exhaustion, a dexamethasone (4 mg)

suppression test, Three Factor Eating Questionnaire (TFEQ) scores, anthropometry and body composition.

Measurements

5-hour cortisol exposure

At 8 a.m. after an overnight fast an intravenous catheter for blood sampling was inserted into the forearm vein. Directly after insertion of the catheter the first blood sample (9 ml) was drawn (0 minutes), and after 15 minutes the participants received their breakfast (20 % of the participants' daily energy requirement). At 45, 90 and 210 minutes the second, third and fourth blood samples were drawn. At 225 minutes the participants received their lunch (30 % of the participants' daily energy requirement) and at 255 and 300 minutes the fifth and sixth blood samples were drawn. During the test, participants were able to drink water ad libitum. Breakfast and lunch consisted of typical Dutch products, namely brown bread (1.0 MJ per 100 g; en% C/P/F: 74/15/11) with young cheese (1.6 MJ per 100 g; en% C/P/F: 0/26/74) as well as ab libitum tea or water consumption. The daily energy requirements were calculated by multiplying the basal metabolic rate (BMR) by an activity index of 1.75. The BMR (MJ/day) was calculated according to the equations of Harris-Benedict [13]. Participants remained seated during sampling and daytime naps were not allowed. The participants' 5-hour cortisol exposure was measured by calculating the area under the curve (AUC) (0-300 minutes).

Progressive maximal cycling test to volitional exhaustion

Each subject underwent a progressive maximal cycling test to volitional exhaustion for quantification of VO_2^{\max} [14]. Subjects cycled on an electromagnetically braked bicycle ergometer (Lode, Groningen, The Netherlands) at 80 rpm, at a workload of 75 W for women and 100 W for men. After 5 minutes, the workload was increased every 2.5 minutes with 50 W, until subjects were no longer able to

maintain a pedaling frequency above 80 rpm. Open-circuit spirometry was performed with an Oxycon (Mijnhardt Jaeger, Mannheim, Germany), and oxygen consumption (VO_2) and carbon dioxide production (CO_2) were measured in the breath-by-breath mode and averaged over 15 seconds. Heart rate was monitored continuously throughout all exercise protocols (Polar s610). Verification that each subject actually achieved VO_2^{\max} consisted of meeting the following criteria: 1) achieving predicted maximal heart rate and 2) a respiratory exchange ratio of 1.10 or more.

Dexamethasone (4 mg) suppression test

Each participant received 4 mg dexamethasone 8 hours before the suppression test, all participants participated in the treatment, and no adverse reactions were reported. Participants abstained from alcohol consumption and strenuous exercise 15 h before testing. At 8 a.m. after an overnight fast, an intravenous catheter for blood sampling was inserted into the forearm vein and a blood sample was drawn [12].

Blood samples

Blood samples were collected and mixed in serum tubes (BD Vacutainer, 9.5 ml). Serum was obtained by centrifugation (4°C, 3000 rpm, 10 minutes), frozen in liquid nitrogen and stored at -80°C until analysis of cortisol concentrations. Serum cortisol concentrations were measured using RIA (Buhlmann Laboratories) and detection limits for cortisol concentrations were 2.9 ng/ml. Intra-assay coefficient variation for cortisol concentrations was less than 6 % for all samples.

Attitude towards eating

Eating behavior was analyzed using a validated Dutch translation of the TFEQ [2, 15]. The TFEQ consists of 3 factors measuring a person's attitude towards eating. Dietary restraint (Factor 1) reflects the extent to which individuals attempt to cognitively control their food intake. Inhibition of restraint (Factor 2 or

disinhibition) reflects individual differences in the extent to which release from the cognitive suppression of eating occurs in response to the presence of palatable food or other disinhibiting stimuli, such as emotional distress. Factor 3 refers to the subjective feeling of hunger. Based upon the median for the TFEQ scores in the South of the Netherlands, participants were characterized as unrestrained when dietary restraint scores were < 9 , and as restrained when scores ≥ 9 . Participants were characterized as having low disinhibition when disinhibition scores were < 5 , and as having high disinhibition when scores were ≥ 5 [1].

Anthropometry and body composition

Measurements were carried out in the morning after voiding the bladder and before breakfast. Body weight (BW) (Sauter D7470, Ebingen, Germany) and height (Seca, model 220, Hamburg, Germany) were measured to the nearest 0.01 kg and 0.1 cm respectively. BMI (kg/m^2) was calculated as body weight divided by height (m) squared. The waist circumference was measured to the nearest 0.1 cm at the site of the smallest circumference between the rib cage and the ileac crest, whereas the hip circumference was measured to the nearest 0.1 cm at the level of maximum extension of the buttocks. Both measurements were done with the participants in standing position. Body density was determined by hydro densitometry with simultaneous measurements of residual lung volume with the helium dilution technique. Total body water (TBW) was determined with the deuterium dilution method following the Maastricht protocol [16]. Body composition was calculated from body density and TBW using the three-compartment model of Siri. Fat free mass (FFM) was calculated by dividing TBW by the hydration factor 0.73. Fat mass (FM) was determined as $\text{BW} - \text{FFM}$.

Statistical analysis

Differences between two groups were determined using unpaired t-tests.

Differences over time and between conditions were determined using two-factor ANOVA with repeated measures. Relationships between dependent and independent variables were determined using simple linear, or multiple regression models. All statistical tests were two-tailed, differences were considered significant at $p < 0.05$ and values are expressed as mean \pm standard deviation.

Results

Table 1 shows the characteristics of all participants ($n = 76$). No significant differences were shown between men and women in age, BMI, hip circumference and feeling of hunger score. The male participants showed significantly higher height, body weight, waist circumference, waist-to-hip ratio, and maximal workload during the progressive maximal cycling test when compared to the female participants. The female participants showed significantly higher body fat percentage, area under the curve (AUC) of 5-hour cortisol exposure, cortisol concentrations after a dexamethasone (4 mg) suppression test, dietary restraint score, and disinhibition score when compared to male participants. Therefore, the results of the male and female participants were analyzed separately.

Based upon the median for the TFEQ scores in the South of the Netherlands the female participants were characterized as unrestrained when dietary restraint scores were < 9 , and as restrained when score ≥ 9 [1]. **Table 2** shows the characteristics of the dietary unrestrained ($n = 26$) and the dietary restrained ($n = 12$) women. The dietary restrained women showed significantly higher BMI and body fat percentage when compared to the dietary unrestrained women. No significant differences were shown between dietary unrestrained and dietary restrained women in age, height, body weight, waist circumference, hip circumference, waist-to-hip ratio, maximal workload during the progressive maximal cycling test, disinhibition score and feeling of hunger score.

Table 1. Characteristics of participants (n = 76)

	Men (n = 38)	Women (n = 38)	P value ¹
Age (year)	31.2 ± 5.2	25.9 ± 7.5	0.17
Height (cm)	183.5 ± 7.3	170.4 ± 6.6	0.01
Body weight (kg)	75.2 ± 7.9	65.6 ± 7.5	0.01
Body mass index (kg/m ²)	22.3 ± 1.5	22.6 ± 1.9	0.99
Body fat (%)	11.6 ± 5.2	25.2 ± 5.5	0.01
Waist circumference (cm)	80.6 ± 5.5	74.4 ± 5.8	0.01
Hip circumference (cm)	100.8 ± 5.8	99.6 ± 5.7	0.12
Waist-to-hip ratio	0.80 ± 0.04	0.74 ± 0.06	0.01
Maximal workload (V)	323 ± 55	216 ± 44	0.01
AUC ² of 5-hour cortisol exposure (nmol/L.min)	1.1(10 ⁵) ± 0.2	1.6(10 ⁵) ± 0.7	0.01
Cortisol after dexamethasone (4 mg) (nmol/L)	33.9 ± 12.6	53.7 ± 28.7	0.01
Dietary restraint score	4.7 ± 3.5	6.5 ± 3.3	0.08
Disinhibition score	2.9 ± 1.2	4.2 ± 1.9	0.02
Feeling of hunger score	3.8 ± 2.5	3.9 ± 2.4	0.84

¹ Differences between men vs. women (t-test)

²AUC, area under curve

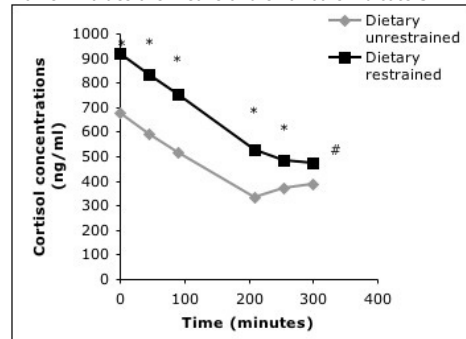
Table 2. Characteristics of dietary unrestrained (score < 9) and dietary restrained (score ≥ 9) women (n = 38)

	Dietary unrestrained (n = 26)	Dietary Restrained (n = 12)	P value ¹
Age (year)	26.1 ± 6.2	25.9 ± 7.5	0.87
Height (cm)	171.2 ± 6.2	168.6 ± 7.2	0.26
Body weight (kg)	64.8 ± 7.8	67.2 ± 6.6	0.35
Body mass index (kg/m ²)	22.1 ± 1.8	23.7 ± 1.6	0.01
Body fat (%)	24.2 ± 4.5	28.6 ± 5.6	0.02
Waist circumference (cm)	74.9 ± 6.8	73.4 ± 3.7	0.57
Hip circumference (cm)	98.6 ± 5.9	101.7 ± 5.2	0.22
Waist-to-hip ratio	0.75 ± 0.06	0.72 ± 0.04	0.22
Maximal workload (V)	214 ± 46	218 ± 41	0.78
Dietary restraint score	4.7 ± 2.1	10.5 ± 1.5	0.01
Disinhibition score	4.3 ± 2.1	3.9 ± 1.8	0.58
Feeling of hunger score	4.0 ± 2.5	3.8 ± 2.2	0.84

¹ Differences between dietary unrestrained vs. dietary restrained women (t-test)

Figure 1 shows the changes in cortisol concentrations (nmol/L.min) over a 5-hour time period in the dietary unrestrained and dietary restrained women. The cortisol concentrations were significantly higher at all time points except at 300 minutes in the dietary restrained women when compared to the dietary unrestrained women. Over time there was a significant decrease of cortisol in both groups of women, however, the decrease was significantly stronger in the dietary restrained women. The AUC over a 5-hour time period of the cortisol concentrations was significantly higher in the dietary restrained women when compared to the dietary unrestrained women (136429.8 ± 57665.9 vs. 193622.5 ± 89322.2 nmol/L.min, p <

0.02). Additionally, the cortisol concentrations after a dexamethasone (4 mg) suppression test were significantly higher in the dietary restrained women when compared to the dietary unrestrained women (70.7 ± 36.6 vs. 45.8 ± 20.6 nmol/L, p < 0.01).

Figure 1. Changes in cortisol concentrations (nmol/L) over a 5-hour time period in the dietary unrestrained (grey, n = 26) and dietary restrained (black, n = 12) women. Values are means and error bars indicate SD.

* P < 0.05 for differences in dietary unrestrained women compared to dietary restrained women (2- factor ANOVA repeated measures)

P < 0.05 overall group x time interaction was significant

Table 3 shows a positive relationship between the cortisol concentrations after a dexamethasone (4 mg) suppression test and the AUC over a 5-hour time period of the cortisol concentrations, the dietary restraint score, and the disinhibition scores as the independent variables ($R^2 = 0.62$, p < 0.01) in normal weight women.

Table 4 shows a positive relationship between the AUC over a 5-hour time period of the cortisol concentrations as the dependent variable and the dietary restraint scores and the disinhibition scores as the independent variables ($R^2 = 0.23$, p < 0.01) in normal weight women.

Furthermore, in normal weight men no relationship was seen between HPA axis functioning and eating behavior. Here different relationships appeared that were not primarily related to HPA axis functioning. Dietary restraint scores were < 9 in the men, except 3 outliers, who were athletes. In these normal weight men a linear regression model showed a

positive relationship between the dietary restraint score as the dependent variable and the maximal workload during the progressive maximal cycling test as the independent variable ($R^2 = 0.12$, $p < 0.03$). Additionally, a linear regression model showed a negative relationship between the AUC over a 5-hour time period of the cortisol concentrations as the dependent variable and the dietary restraint score as the independent variable ($R^2 = 0.13$, $p < 0.03$).

Table 3: Multiple regression model with the cortisol concentrations after a dexamethasone (4 mg) suppression test (nmol/L) as the dependent variable and the area under the curve of cortisol concentrations over a 5-hour time period (nmol/L), dietary restraint score and disinhibition score as the independent variables in normal weight women ($n = 38$)
 $R^2 = 0.53$, $p < 0.01$

	Partial β	Std. Error	P value
Intercept	9.8	11.05	0.38
AUC ¹ of cortisol concentrations over a 5-hour time period	0.002	0.69	0.001
Dietary restraint score	1.1	0.1	0.32
Disinhibition score	-1.3	-0.09	0.49

¹ AUC; area under the curve

Table 4: Multiple regression model with the area under the curve of cortisol concentrations over a 5-hour time period (nmol/L) as the dependent variable and dietary restraint score and disinhibition score as the independent variables in normal weight women ($n = 38$)
 $R^2 = 0.23$, $p < 0.01$

	Partial β	Std. Error	P value
Intercept	49329.5	34584.1	0.16
Dietary restraint score	8031.7	3273.2	0.01
Disinhibition score	12605.3	5500.5	0.01

¹ AUC; area under the curve

Discussion

The objective of our study was to investigate the relationship between HPA axis functioning and dietary restraint in normal weight (BMI between 20 and 25 kg/m²) men and women. No such relationship was found in men. Dietary restrained women showed increased cortisol concentrations over a 5-hour time period, increased cortisol concentrations after a dexamethasone (4 mg) suppression test, higher BMI, and higher body fat percentage. Moreover, a positive relationship was found between cortisol concentrations over a 5-hour time period and dietary restraint alone or in combination with the disinhibition score.

Our study contributes to insight in the relationship between HPA axis functioning and dietary restraint in normal weight women [8-11]. Our study is the first to analyze HPA axis functioning elaborately, since most studies only measured cortisol exposure at only one (sometimes unspecified) time point during the day or the night, and measured no HPA axis feedback functioning. In accordance to the literature [3, 4] we found that dietary restrained women have a higher BMI and body fat percentage compared to dietary unrestrained women. To determine cortisol exposure, we measured cortisol concentrations (nmol/L.min) over a 5-hour time period. We showed that the dietary restrained normal weight women had a significant higher AUC over a 5-hour time period of the cortisol concentrations and a stronger decrease in cortisol concentrations compared to the dietary unrestrained women. This suggests that normal weight women, who attempt more to restrict food intake through dietary restraint, have increased cortisol exposure and stronger decline in cortisol levels over time.

Additionally, we measured feedback functioning through a dexamethasone (4 mg) suppression test [12], which measures the amount of cortisol that escaped from dexamethasone suppression. When cortisol feedback is strong low amount of cortisol will be present, and when high amounts of cortisol escaped, cortisol feedback is decreased [12]. We showed that the dietary restrained normal weight women had higher cortisol concentrations after dexamethasone suppression test, when compared to the dietary unrestrained women. These results imply that normal weight women, who attempt more to restrict food intake through dietary restraint, have decreased cortisol feedback functioning.

Multiple regression analysis showed a positive relationship between cortisol feedback functioning and cortisol exposure as well as dietary restraint score. Additionally, we showed a positive

relationship between cortisol exposure and dietary restraint in combination with disinhibition score. This suggests that especially normal weight women, who attempt to restrict food intake through dietary restraint but who have high inhibition of restraint show increased cortisol exposure, and decreased cortisol feedback. However, the direction of causality cannot be inferred from this research.

The lack of a direct relationship between HPA axis functioning and dietary restraint in normal weight men is consistent with the literature [8-11]. Here different relationships appeared that were not primarily related to HPA axis functioning. A possible explanation may be the gender difference in stress sensation [12]. In men a positive relationship was shown between the dietary restraint score and the maximal workload during the progressive maximal cycling test, which suggests that only men who are sincerely engaged in sports are dietary restraint. Additionally, a negative relationship was shown between the maximal workload during the progressive maximal cycling test and AUC over a 5-hour time period of the cortisol concentrations, which suggests that men who are sincerely engaged in sports have decreased cortisol exposure, as previously shown [12]. In general, only men who are overweight or who are sincerely engaged in sports are dietary restraint. Overweight men were not included in our study and when the normal weight men who are sincerely engaged in sports were omitted from the analysis, no relationship between HPA axis functioning and dietary restraint

was found. These results suggest that dietary restraint is not perceived as a stressor in normal weight men.

Increased cortisol exposure, heightened cortisol responsiveness and decreased cortisol feedback functioning are characteristics of hypercortisolism [12]. Hypercortisolism is a state in which the HPA axis responsiveness is preserved, which is also seen in psychological or physiological chronic stressed humans [17-19]. It may therefore be hypothesized that the association between increased cortisol levels and dietary restraint may be mediated through psychological stress [8-11]. Sustained hypercortisolism however may predispose individuals to many illnesses, ranging from inflammatory diseases to major depression, thereby indicating that dietary restraint in combination with disinhibition is a possible health risk [20]. In conclusion, hyperactivity of the HPA-axis is related to dietary restraint especially in combination with disinhibition, in normal weight women.

Acknowledgements

We want to thank our participants for their participation in this study. We gratefully thank Loek Wouters and Jose Sulon for their assistance. FR (supervised by AN and MW) carried out the study, collected and analyzed the data, and wrote the largest part of the manuscript. SL and JB reviewed the manuscript. Planning, processing the results and writing the manuscript were done under general supervision by AN and MW. The authors had no conflict of interest.

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Chapter 11

Discussion

Body weight appears to be regulated with remarkable precision in healthy adults, and a stable body weight reflects a balance between energy intake and energy expenditure [1]. Chronic deregulation of energy balance with energy intake exceeding energy expenditure, leads to storage of excessive energy as fat, a characteristic of overweight and obesity [2]. The first aim of the studies presented in this thesis was to identify genetic, parental, behavioral, and endocrine factors involved in body-weight development in children, and the interaction between these factors. The second aim was to assess genetic, behavioral, and endocrine factors and their interaction in body-weight regulation in adults.

Body-weight development during childhood and puberty

Genetic factors

With respect to genetic factors involved in body-weight development, we observed that in our Dutch children cohort the A allele of the fat mass and obesity-associated gene (FTO) (rs9939609) was associated with higher BMI, fat mass index (FMI), leptin concentrations, and lower activity scores from childhood to puberty [3]. In the last decades a large number of polymorphisms have been related to variation in body weight [4]. For our studies, we selected polymorphisms with a high allelic frequency distribution and a proposed biological function that could influence body-weight development. The FTO polymorphism is related to altered DNA methylation [5, 6], which results in altered biological function. Studies in humans and rodents have suggested a central role for FTO through an effect on regulation of food intake, such as decreased feelings of satiety [7, 8], and a peripheral role through an effect on lipolytic activity in adipose tissue [6]. Additionally, we observed in our cohort that at age 13y a larger BMI is associated to the FTO A allele, larger leptin concentrations, and decreased physical activity. These results implicate that at

age 13y genetic, endocrine and psychological factors are independent predictors for BMI [3]. At age 12, 14, and 15y BMI was associated to larger leptin concentrations, which implicates that during puberty BMI is temporarily predicted by FTO and predominately predicted by endocrine factors [3]. Furthermore, in our cohort polymorphisms of the ciliary neurotrophic factor (CNTF) gene and the peroxisome proliferated-activated receptor γ 2 (PPAR γ 2) gene did not show similar associations during puberty [9], although this was expected based upon studies in adults [10]. These results suggest that especially polymorphisms with a strong influence on body weight, such as FTO [6], affect body weight during puberty, and that the influence of polymorphisms, such as CNTF and PPAR γ 2, are stronger during early childhood and late adolescence, as observed by previous studies [11, 12]. Our results implicate that the changes in the associations between body-weight development and polymorphisms over time may be explained by the fact that body weight is predominately regulated by physiological and endocrine factors during puberty.

Parental factors

With respect to parental factors involved in body-weight development, we observed that in our cohort a high BMI of the father and a high disinhibition score of the mother were related to an increased BMI in the child [9]. These observations are in concordance with previous findings, which observed that the child's BMI is positively related to the parents' BMI [13, 14]. In contrast to previous findings we did not show a relationship between mothers cognitively restrained eating behavior and the body-weight development of the children [14]. A possible explanation may be that the mothers were successfully restrained, so their high restraint scores helped them to control their weight successfully [15], and only when a high disinhibition score was present this resulted in a relationship with the weight of the child [9]. The parent association is due to genetic and environmental factors,

such as shared lifestyles (i.e. diets, feeding practices and food choices) and patterns of activity. For future research, it will be of interest to observe whether the role of these genetic and parental factors in body-weight development will become stronger in our cohort as well as in other cohorts during late adolescence.

Behavioral factors

With respect to the behavioral factors involved in body-weight development, we were the first to show that development of BMI during puberty is inversely related to change in sleep duration [16]. We focused on short sleep duration, as a number of cross-sectional and longitudinal studies have shown a negative relationship between habitual sleep duration and body weight in children [17, 18]. None of the four longitudinal studies in children however investigated whether changes in sleep duration were associated with changes in body weight [19-22]. Although cause and effect cannot be completely disentangled from the observed inverse relationship between changes in body weight and sleep duration, we carefully suggest that reduction of sleep duration contributes to development of overweight during puberty.

Short sleep has been suggested to result from psychological distress, and the physiological disturbances related to psychological distress, such as hypercortisolemia [23, 24]. Studies investigating the physiological mechanisms on how sleep duration affects body-weight development in humans are however limited. Laboratory based studies have observed that short sleep (4 hours) lead to lower levels of the anorexigenic hormone leptin and higher levels of the orexigenic hormone ghrelin [25, 26]. Short sleep has also been related to reduced insulin sensitivity and increased glucose levels [27]. Other physiological alterations that result from disturbed energy balance are suggested to be involved in body-weight development, but have not been investigated yet. For future research, it will therefore be of interest to identify the physiological mechanisms

behind the relationship between short sleep and altered body weight.

Endocrine factors

With respect to endocrine factors involved in body-weight development, leptin is suggested to be a permissive factor for the start of puberty [28-30], which would be mediated through the stimulatory effect of leptin on the release of gonadotrophic hormones, and consequently gonadal hormones. We observed that in boys the leptin/fat mass ratio decreased from Tanner stage 2 onwards, while in girls this ratio decreased from Tanner stage 2 and increased again at Tanner stage 5 [31]. These results show that factors independent of fat mass become (transiently) more important in the regulation of plasma leptin concentrations in boys and girls [31]. The proposed mechanism behind the temporal relationships between leptin, gonadotrophic hormones, and gonadal hormones during puberty, beholds that leptin independently of fat mass acts on the hypothalamic luteinizing hormone-releasing hormone pulse generator [32-34]. Consequently, the release of LH and FSH from the pituitary is stimulated. LH and FSH in turn stimulate the gonads to release testosterone and estradiol, which will form a negative feedback loop, which will inhibit the secretion of LH and FSH. Moreover, testosterone alone will form a negative feedback loop, which will inhibit leptin secretion from the adipocytes, and estradiol will form a positive feedback loop, which stimulates leptin secretion from the adipocytes [34-36]. As a result, the relationship between leptin concentrations and body fat is altered during puberty.

However, in the literature there is also evidence that does not support the role of leptin as a permissive factor for the start of puberty [37]. A sex specific association may explain contradicting findings, as we found that in girls a peak in leptin was present at the start of puberty followed by a peak in the gonadotropic hormone, and temporal relationships were observed between leptin and gonadotropic

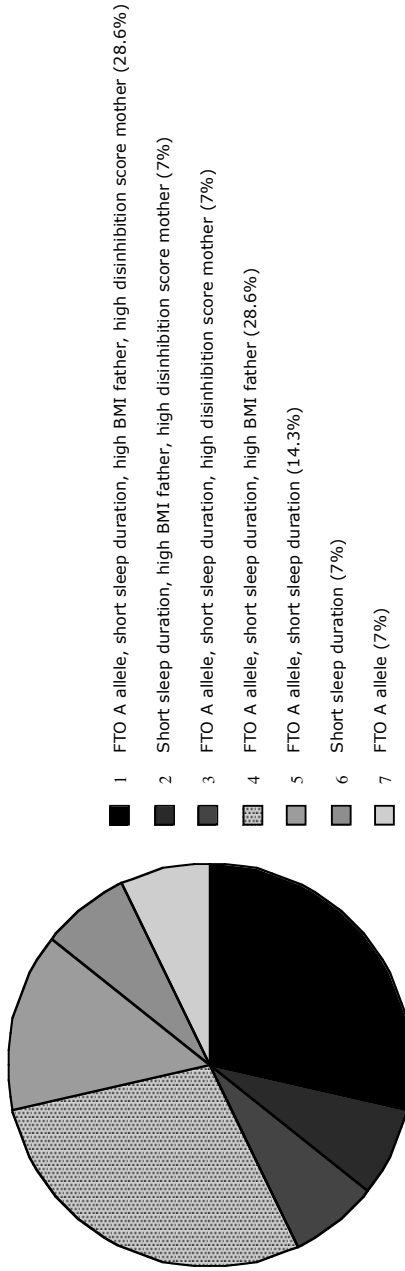
hormones during early puberty. In boys, however there was no peak in leptin, LH, and FSH, and leptin was only related to LH and FSH during late puberty [31]. We therefore suggested that in girls leptin acts as a permissive factor for the onset of puberty, while in boys leptin has a different function and different timing [31]. Why it is more important to have a permissive factor for the start of puberty in girls than in boys may be explained from an evolutionary point of view; girls need enough energy reserves in the form of body fat for the start of puberty for reproduction [38]. The sex differences in puberty timing may result from differences in expression of genes involved in puberty timing [39]. For future research, it will therefore be of interest to identify the mechanisms behind sex differences in puberty timing.

overweight during puberty. In contrast, most of the normal-weight children still appeared to have one or more of these factors that are associated with overweight.

To summarize the previous described results, we included a pie chart (**figure 1**) that describes the factors involved in body-weight development in overweight children, and the interactions between the factors in our Dutch children cohort during puberty at age 15y. The results from our studies implicate that body-weight development during puberty is related to the FTO polymorphism, as 80 percent of overweight children had the A allele [3]. Additionally, the behavioral factor sleep duration influenced body-weight development, as 93 percent of the overweight children had short sleep duration [16]. Finally, the parental factors high BMI of the father and high disinhibition score of the mother influenced body-weight development in overweight children. The effects of the genetic, parental and behavioral factors are largely combined, as 28% of the overweight children have all 4 factors related to body-weight development, and about 43% of the overweight children have 3 out of 4 factors. Only 28% of the overweight children have 1 or 2 factors related to body-weight development. Overall, the model (figure 1) suggests that the FTO A allele, short sleep duration, and the parental factors interact, and together are involved in development of

Figure 1: a pie chart describing the factors involved in body-weight development and the interactions between the factors in the overweight children (n=14) of our Dutch children cohort at age 15y.

Overweight children



Hypothalamus/pituitary/adrenal (HPA) axis and body-weight regulation

The HPA axis is involved in body-weight regulation, and alterations in HPA axis functioning are suggested to be involved in the pathogenesis of obesity, in particular obesity characterized by visceral fat distribution [40]. Support for the involvement of the HPA axis in body-weight regulation is found in two extremes of plasma cortisol levels in humans; Addison's disease (hypocortisolism) that is related to weight loss, and Cushing's syndrome (hypercortisolism) that is related to rapid weight gain, particularly of the trunk and face with sparing of the limbs [41]. It however is not completely clear to what extent the HPA axis is physiologically involved in the regulation of body weight [42]. Evidence for a role of the HPA axis in body-weight regulation is provided via the genetic, behavioral, and endocrine factors described in this thesis.

Genetic factors

With respect to the genetic factors, several polymorphisms of the glucocorticoid receptor, such as the BclI (rs41423247) polymorphism, have been related to body-weight regulation, body fat distribution and altered HPA axis functioning [43]. We observed that the BclI G/G genotype is associated with increased cortisol exposure as well as decreased cortisol feedback sensitivity under stress in women but not in men [44]. These findings are in concordance with the sex specific associations of the BclI polymorphism observed in other studies, which showed a diminished cortisol response to psychosocial stress in men with the BclI G/G genotype and an increased cortisol response in women [45]. The sex specific associations are presumably caused by the differences in cellular hormonal milieu, which lead to differential expressivity of the underlying genetic networks [46].

The BclI polymorphism is also involved in the efficacy of cortisol signaling and thereby influencing the downstream biology of peripheral and central cortisol

responsive systems. Therefore the BclI polymorphism is suggested to be one of the factors that underlie individual susceptibility to stress related disorders, such as depression and the metabolic syndrome. The associations between the BclI G/G genotype and altered body weight and body fat distribution were only found in a few studies in men, while these relationships were always present in women [47-49]. Together with the earlier discussed sex specific associations on altered HPA axis functioning in the BclI G/G genotype [44, 45], this suggests that women with the BclI G/G genotype are more susceptible to disturbed regulation of body weight and HPA axis functioning compared to men with the same genotype. For future research on the role of HPA axis related polymorphism in body-weight regulation and HPA axis functioning, it is therefore strongly suggested to take possible sex differences into account.

Endocrine factors

With respect to the endocrine factors, we observed that low cortisol variability, enhanced negative feedback, and impaired HPA axis functioning in a challenged condition under 4 mg dexamethasone were related to visceral fat accumulation in both men and women [50]. These findings add to evidence from previous studies that observed associations between visceral fat distribution and alterations in HPA axis functioning, such as decreased cortisol variability [51-53], and increased cortisol secretion after physical and psychological stressors [51, 52, 54]. Studies on whether body fat distribution is related to feedback functioning, thereby using dexamethasone (a glucocorticoid receptor agonist), have however yielded inconclusive results [52, 53, 55]. Our results are consistent with previous work that suggests a deregulation of the HPA axis in humans with relative more visceral fat [50].

The disturbed state of low cortisol variability related to visceral fat distribution possibly results from a combination of alterations of the HPA axis.

Firstly, as often stated by the literature, the alterations result from decreased negative feedback mediated through a decrease in glucocorticoid receptors of the pituitary [52, 56] and through a decrease in the mineralocorticoid receptor response [57, 58], which leads to higher cortisol levels after a stressor [51, 52, 54] and after dexamethasone suppression in unchallenged conditions [50, 55]. Secondly, we suggest that lower cortisol levels in a challenged condition under 4 mg dexamethasone [50] result from decreased sensitivity of the adrenal gland to ACTH as suggested by Mattsson et al 2009 [58]. These suggested alterations have until now not been directly tested, as most studies examine HPA axis functioning under basal conditions or via dexamethasone, an agonist of the GR receptor. For future research it is of interest to study the role of the mineralocorticoid receptor and the decreased sensitivity of the adrenal gland in the disturbance of the HPA axis related to visceral fat distribution.

Behavioral factors

With respect to the behavioral factors involved in body-weight regulation, we observed that acute psychological stress results in eating in the absence of hunger [60]. In our study, the observed food intake in the absence of hunger after stress exposure, hardly led to further decreases in hunger, underscoring the involvement of a non-homeostatic regulatory mechanism, such as reward. Previous studies on stress-induced eating [61-64] proposed the involvement of reward, and Dallman et al [65] even introduced the term 'comfort foods' to emphasize the idea that the drive behind stress-induced eating is not homeostatic. These studies could however not directly demonstrate the role of reward, as a role for hunger could not be excluded [65, 66]. This is possible when using the 'eating in the absence of hunger' paradigm [60]. Stress-induced eating is suggested to be caused by disturbance of several neural networks related to reward, such as the opioid and NPY system [67, 68]. Recently, researchers from our lab [69] observed

that stress prevented a meal-induced decrease of wanting and energy intake in visceral obese objects and not in normal weight objects. Furthermore, Born et al. [70] from our lab observed in an fMRI study using the 'eating in absence of hunger' paradigm that activity in reward-related brain areas were significantly reduced under stress. These findings further support the role of disturbed reward in stress-induced eating.

Large inter-individual differences are present in the effect of stress on food intake; stress can increase, decrease or does not alter food intake. These individual differences in response to stress were also present in our eating in the absence of hunger paradigm [60], and may relate to eating behavior characteristics [71], as an increase in energy intake during stress has been found in individuals with high scores on dietary restraint and/or emotional eating [61-64]. We however observed that stress induced an increase in eating in the absence of hunger especially in vulnerable individuals, which are characterized by disinhibited eating behavior and sensitivity to chronic stress [60]. High dietary cognitive restraint scores did not result in stress induced eating in the absence of hunger, which is contradictory to previous findings [61-64]. The contradicting results may be explained by the fact that previous studies used the Herman-Polivy restraint scale [72], in which high restraint score is related to weight concern, such as disinhibited eating and weight fluctuations, in contrast to the TFEQ [71] that we used in our study, in which high restraint score is related to (successful) control of food intake [15, 73]. It therefore is suggested that stress-induced eating is caused by disinhibition, which reflects individual differences in the extent to which release from the cognitive suppression of eating occurs in response to the presence of palatable food or other disinhibiting stimuli, such as emotional distress [71].

Several studies have suggested that dietary restraint can be experienced as a

chronic physiological stressor. We observed that in normal weight women hyperactivity of the HPA-axis is related to dietary restraint especially in combination with disinhibition [74]. This is in contrast to previous studies that have shown that only dietary restraint scores were related to alterations in HPA axis functioning, such as increased cortisol levels [75-77]. Our results suggest that dietary restraint in combination with disinhibited eating behavior can be considered as a chronic stressor. Together with the results from our study on the relationship between stress and eating in the absence of hunger [60], our results imply that disinhibition is an important mediator in the relationship between HPA axis functioning and body-weight regulation. For future research it will be of interest to investigate whether disinhibited eaters have a higher cortisol response to stress, which in turn may lead to larger food intake, and whether these differences are related to altered feelings of reward.

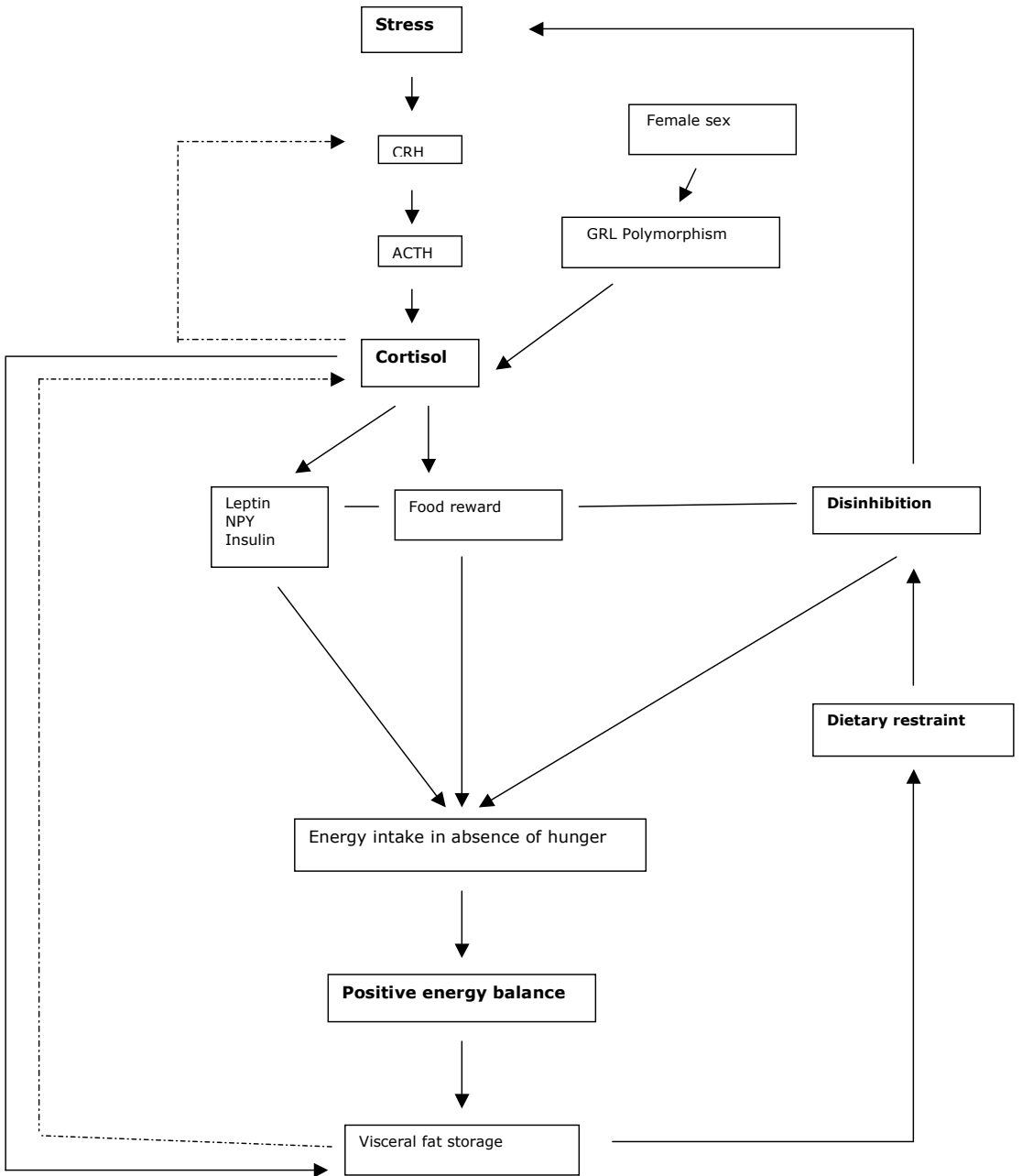
To summarize, we propose a model (figure 2) on how disturbance of the HPA axis is involved in body-weight regulation. Stress (psychological or physical) results via secretion of CRH and ACTH in production of cortisol [40]. HPA axis activation is larger in women with the BclI G/G genotype, which results in higher cortisol levels in these women compared to men with the same genotype [44, 45]. In turn, cortisol interacts with the neuro-endocrine pathways that control the homeostatic regulation of food intake, as

it potentiates the orexigenic actions of NPY [68], induces leptin resistance resulting in reduced suppression of food intake [78], and interacts with insulin that leads to high-fat food choice [79]. Cortisol also interacts with the neuro-endocrine pathways that control the non-homeostatic regulation of food intake, which results in a decrease in opioid levels [67].

The interaction of cortisol with the pathways that control the homeostatic and non-homeostatic regulation of food intake results in stress-induced eating in the absence of hunger [60], especially in disinhibited eaters [60]. Stress-induced eating results in a positive energy balance that leads to visceral fat storage, as visceral adipocytes have a four-fold of GR receptors that facilitate fat storage in the presence of high cortisol levels [80, 81]. Visceral fat distribution in turn has been related to the disturbance of HPA axis functioning in basal and challenged conditions [52, 56, 57].

Moreover, increased fat storage may lead to increased weight dissatisfaction, which results in altered eating behavior [71], such as increased dietary restraint to decrease or prevent further weight gain. Dietary restraint in combination with disinhibited eating behavior is considered as a chronic stressor [74], which leads to further activation of the HPA axis. Therefore, we propose that chronic disturbance of the HPA axis initiates a vicious circle (**figure 2**), which puts stress as a major risk factor for excessive visceral weight gain.

Figure 2: a *proposed* model on how disturbance of the HPA axis is involved in body-weight regulation (solid lines represent stimulation and dashed lines represent inhibition).



Conclusions

In children, body-weight development is regulated by various factors and interactions between these factors, such as the A allele of the FTO gene, which was associated with higher BMI, FMI, leptin concentrations, and lower activity scores from childhood to puberty. The influence of the FTO polymorphism however changed over time, as at age 13y the FTO A allele, larger leptin concentrations and decreased physical activity were independent predictors for a larger BMI, while at age 12, 14, and 15y BMI was only predicted by larger leptin concentrations. Polymorphisms of the CNTF and PPAR γ 2 genes did not show an association with body-weight development during puberty, which implicates that during puberty BMI is only temporarily predicted by FTO and predominately predicted by endocrine factors. Additionally, the behavioral factor sleep duration was involved in body-weight development, as changes in BMI during puberty were inversely related to changes in sleep duration. Subsequently, a higher BMI is associated with an earlier leptin peak in girls, and thereby an earlier start of puberty, as in girls leptin acts as a permissive factor for the onset of puberty. In boys leptin however has a different function and different timing. After the start of puberty factors independent of fat mass become (transiently) more important in the regulation of plasma leptin concentrations in boys and girls.

In adults, stress results in eating in the absence of hunger, which demonstrates for the first time that non-homeostatic mechanisms, such as reward, are involved in stress-induced eating. Stress-induced eating was present especially in subjects with disinhibited eating behavior and sensitivity to chronic stress. Disinhibited eating behavior in combination with dietary restraint is related to hyperactivity of the HPA axis. Disturbance of HPA axis

functioning under basal and challenged conditions is related to visceral fat accumulation. Therefore, we propose that chronic disturbance of the HPA axis initiates a vicious circle, which puts stress as a major risk factor for excessive visceral weight gain.

Future research

A further follow-up of our Dutch children cohort is strongly recommended, since this cohort has provided valuable data on body-weight development during childhood and puberty. It is of interest to observe whether the role of endocrine factors will diminish once the children are in late adolescence, and whether the role of genetic factors in body-weight development will become stronger during this period.

Furthermore, for future research, it will be of interest to identify the physiological mechanisms behind the relationship between short sleep and altered body weight, and to investigate whether weight loss or weight stabilization can be achieved via sleep duration modification. Additionally, future experiments on the role of HPA axis functioning in body-weight regulation should be conducted to further disentangle this relationship. It remains to be elucidated, whether the HPA axis is the primary cause for the obese state in the visceral obese, which can only be investigated via longitudinal studies. Furthermore, the disturbance of several neural homeostatic and non-homeostatic networks involved in energy intake by HPA axis activation, such as the opioid and NPY system, are thought to be involved in stress-induced eating, however no direct evidence is found in humans yet. For future research it is of interest to identify brain regions involved in stress-induced eating and to evaluate if these regions are involved in homeostatic and non-homeostatic regulated food intake.

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Summary

A stable body weight reflects a balance between energy intake and energy expenditure. Chronic deregulation of energy balance leads to storage of excessive energy as fat, a characteristic of overweight and obesity. The studies described in this thesis encompass genetic, parental, behavioral, and endocrine factors involved in body-weight development and regulation during childhood, puberty and adulthood. The first part elaborates on the factors involved in body-weight development during childhood and puberty, studied in a large cohort of Dutch children, from which valuable anthropometrical data from birth till age 7y were available. To study body-weight development during puberty, follow-up studies were performed yearly between 2004 and 2008, when the children had a mean age of 12 to 16y. In this cohort the A allele of the FTO gene (rs9939609) was associated with higher BMI, fat mass index, leptin concentrations, and lower activity scores from childhood to puberty. The influence of the FTO polymorphism however changed over time, as at age 13y the FTO A allele, larger leptin concentrations and decreased physical activity were independent predictors for a larger BMI, while at age 12, 14, and 15y BMI was only predicted by larger leptin concentrations (**chapter 2**).

In our cohort, polymorphisms of the CNTF and PPAR γ 2 genes did not show an association with body-weight development during puberty. Our results implicate that BMI is only temporarily predicted by the FTO A allele and predominately predicted by endocrine factors. Next to the genetic factors, parental factors were involved in body-weight development during puberty, as a high BMI of the father and a high disinhibition score of the mother were related to an increased BMI in the child at age 12y (**chapter 3**).

To investigate whether changes in sleep duration were associated with changes in body weight, we measured average sleep duration and body weight from Tanner

stage 1 to 5 in our cohort. We were the first to show that development of BMI during puberty is inversely related to change in sleep duration, and although cause and effect cannot be completely disentangled from the observed inverse relationship, we carefully suggest that reduction of sleep duration contributes to development of overweight during puberty (**chapter 4**).

With respect to endocrine factors, we studied the hormone leptin, as leptin is suggested to be a permissive factor for the start of puberty, which would be mediated through the stimulatory effect of leptin on the release of gonadotrophic hormones, and consequently gonadal hormones. We observed in our cohort that in boys the leptin/fat mass ratio decreased from Tanner stage 2 onwards, while in girls this ratio decreased from Tanner stage 2 and increased again at Tanner stage 5, which shows that factors independent of fat mass become (transiently) more important in the regulation of plasma leptin concentrations in boys and girls. Additionally, we found that in girls a peak in leptin was present at the start of puberty followed by a peak in LH and FSH, and during early puberty temporal relationships were observed between leptin and gonadotropic hormones. In boys, however there was no peak in leptin, LH, and FSH, and leptin was only related to gonadotropic hormones during late puberty. Leptin is therefore suggested to act as a permissive factor for the onset of puberty in girls, while in boys leptin has a different function and different timing (**chapter 5**). To conclude, the data from our cohort suggests that the FTO A allele, short sleep duration, and the parental factors interact, and together are involved in development of overweight during puberty.

The second part of this thesis elaborates on the role of the neuro-endocrine system of the HPA axis in body-weight regulation in adults. The HPA axis has been suggested to be involved in body-weight

regulation, however, to what extent the HPA axis is physiologically involved in the regulation of body weight is not completely clear (**chapter 6**). The physiological role of the HPA axis in body-weight regulation was studied in a cohort of healthy adults, 190 men and women, who had a BMI between 20 and 35 kg/m². Evidence for involvement of the HPA axis in body-weight regulation is provided by genetic factors, such as the BclI polymorphism. In our cohort the BclI G/G genotype was associated with increased cortisol exposure as well as decreased cortisol feedback sensitivity under stress in women but not in men. Together with results on associations between the BclI G/G genotype and altered body weight and body fat distribution in women, this suggests that women with the BclI G/G genotype are more susceptible to disturbed regulation of body weight and HPA axis functioning compared to men with the same genotype (**chapter 7**). Additional evidence for involvement of the HPA axis in body-weight regulation is found in endocrine factors, as we observed that low cortisol exposure, enhanced negative feedback, and impaired HPA axis functioning in a challenged condition under 4 mg dexamethasone were related to visceral fat accumulation in both men and women (**chapter 8**). With respect to the behavioral factors involved in body-weight regulation, we observed that acute psychological stress results in eating in the absence of hunger. The observed food intake in the absence of hunger after stress exposure, hardly led to further decreases in hunger, which demonstrates for the first time that non-homeostatic mechanisms, such as reward, are involved in stress-induced eating.

Previous studies on stress-induced eating propose the involvement of reward, but they could not directly demonstrate the role of reward, as a role for hunger could not be excluded. Large inter-individual differences were present in our eating in the absence of hunger paradigm; stress could increase, decrease or did not alter food intake. These individual differences in response to stress relate to eating behavior characteristics, as we observed a stress-induced increase in eating in the absence of hunger especially in vulnerable individuals, who are characterized by disinhibited eating behavior and sensitivity to chronic stress (**chapter 9**). In our cohort disinhibited eating behavior in combination with dietary restraint was related to hyperactivity of the HPA-axis in normal weight women. Our results suggest that disinhibition in combination with dietary restraint can be considered as a chronic stressor. Together with the results from our study on the relationship between stress and eating in the absence of hunger, our results imply that disinhibition is an important mediator in the relationship between HPA axis functioning and body-weight regulation (**chapter 10**). To conclude, we propose that in adults chronic disturbance of the HPA axis initiates a vicious circle, which puts stress as a major risk factor for excessive visceral weight gain. Stress results in eating in the absence of hunger, especially in subjects with disinhibited eating behavior and sensitivity to chronic stress. Disinhibited eating behavior in combination with dietary restraint is related to hyperactivity of the HPA axis. In turn, disturbance of HPA axis functioning under basal and challenged conditions is related to visceral fat accumulation.

Samenvatting

Een stabiel lichaamsgewicht is een weerspiegeling van de balans tussen energie inname en energiegebruik. Als het evenwicht tussen energie inname en energie gebruik doorslaat naar een positieve balans wordt het teveel aan energie opgeslagen als lichaamsvet. Wanneer deze disbalans lange tijd aanhoudt, ontstaan overgewicht. Dit proefschrift richt zich op erfelijke, ouderlijke, gedragsmatige en endocriene factoren die invloed hebben op de ontwikkeling en regulatie van lichaamsgewicht bij kinderen en volwassenen.

Het eerste deel van dit proefschrift richt zich op de ontwikkeling van lichaamsgewicht zoals dat tijdens de puberteit gemeten is bij een Nederlands kindercohort. Van deze kinderen is vanaf de geboorte tot en met de leeftijd van 7 jaar jaarlijks lengte en gewicht gemeten. Om de ontwikkeling van lichaamsgewicht tijdens de puberteit te bepalen werden tussen 2004 en 2008 vervolgmetingen uitgevoerd, toen deze kinderen gemiddeld een leeftijd van 12 tot 18 jaar hadden. Voor wat betreft genetische invloeden, bleek het voorkomen van het A allel van het fat mass and obesity-associated (FTO) gen (rs9939609) gerelateerd te zijn aan een hogere Body Mass Index (BMI), een hogere vetmassa, hogere leptine concentraties en verlaagde lichamelijke activiteit. De invloed van het FTO A allel bleek echter niet ieder jaar even sterk, want alleen op 13 jarige leeftijd was het FTO A allel een voorspeller voor een hogere BMI. Op 12, 14 en 15 jarige leeftijd waren voornamelijk leptine concentraties voorspellend voor een hogere BMI (**hoofdstuk 2**).

De verschillende genotypen van de ciliary neurotrophic factor (CNTF) en peroxisome proliferated-activated receptor (PPAR γ 2) genen bleken niet geassocieerd te zijn met de ontwikkeling van lichaamsgewicht in het kindercohort. Voor wat betreft de invloed van de ouders, bleek dat op 12 jarige leeftijd een hogere BMI van de

vader en ontremd eetgedrag van de moeder gerelateerd te zijn aan de ontwikkeling van overgewicht bij het kind (**hoofdstuk 3**).

De ontwikkeling van het lichaamsgewicht tijdens de puberteit werd tevens beïnvloed door gedragsmatige factoren zoals het aantal uren slaap per nacht. Wij hebben met de data van ons cohort als eerste aangetoond dat tijdens de puberteit de daling van het aantal uren slaap per nacht gerelateerd is aan de stijging van het lichaamsgewicht. En hoewel het moeilijk is om in deze relatie oorzaak en gevolg te scheiden, suggereren wij dat een verlaging van het aantal uren slaap tijdens de puberteit zou kunnen bijdragen aan de ontwikkeling van overgewicht (**hoofdstuk 4**).

Vervolgens werd ingegaan op de rol van de endocriene factor leptine in de ontwikkeling van lichaamsgewicht. Leptine wordt verondersteld een belangrijke rol te spelen in de aanzet tot de puberteit, omdat leptine mogelijk de productie van de gonadotrope hormonen en geslachtshormonen stimuleert. In ons cohort bleek dat de relatie tussen leptine en vetmassa minder sterk werd vanaf Tanner stadium 2 bij jongens; bij meisjes gebeurde dit ook maar werd de relatie weer sterker in Tanner stadium 5. Dat de relatie verandert, laat zien dat tijdens de puberteit de regulatie van de afgifte van leptine (tijdelijk) bepaald wordt door factoren die niet gerelateerd zijn aan de vetmassa. Verder vond er bij meisjes tijdens de start van de puberteit een piek in leptine concentraties plaats, welke werd gevolgd door een piek in de concentraties van de gonadotrope hormonen. Deze pieken waren afwezig bij de jongens en de relatie tussen leptine en de gonadotrope hormonen was alleen aanwezig tijdens de late stadia van de puberteit bij de jongens. Leptine lijkt dus een belangrijke rol te spelen in de aanzet tot de puberteit bij meisjes, maar lijkt een andere rol en andere timing te hebben bij jongens (**hoofdstuk 5**).

Uit de resultaten van ons kindercohort beschreven in het eerste deel van het proefschrift, kan worden geconcludeerd dat het FTO A allel, weinig uren slaap per nacht, en ouderlijke factoren samen betrokken zijn bij de ontwikkeling van overgewicht tijdens de puberteit.

Het tweede deel van dit proefschrift richt zich op de rol van de hypothalamus-hypofyse-bijnier as (HPA-as) in de regulatie van lichaamsgewicht bij volwassenen (**hoofdstuk 6**).

De fysiologische rol van de HPA-as in de regulatie van lichaamsgewicht werd onderzocht in een cohort van gezonde volwassenen: 190 mannen en vrouwen met een BMI tussen de 20 en 35 kg/m². Genetische factoren, zoals het BclI polymorfisme, bevestigen de rol van de HPA-as in gewichtsregulatie in ons cohort. Het BclI G/G genotype bleek namelijk gerelateerd te zijn aan verhoogde cortisol concentraties en verminderde remming van de HPA-as tijdens stress bij vrouwen, maar niet bij mannen. Uit de literatuur bleek dat het BclI G/G genotype bij vrouwen is gerelateerd aan een hoger lichaamsgewicht en meer visceraal vet. Het totaalbeeld dat ontstaat is dat vrouwen met het BclI G/G genotype gevoeliger zijn voor verstoorde regulatie van lichaamsgewicht en van de HPA-as, dan mannen met het BclI G/G genotype (**hoofdstuk 7**).

Ook endocriene factoren ondersteunen de rol van de HPA-as bij gewichtsregulatie, zoals bleek in ons cohort. Lage exposure van cortisol concentraties, sterkere remming van de HPA-as en verminderd functioneren van de HPA-as bij toediening van 4 mg dexamethasone bleken gerelateerd te zijn aan viscerale vetaccumulatie bij mannen en vrouwen (**hoofdstuk 8**).

Voorts bevestigen gedragsmatige factoren de rol van de HPA-as bij gewichtsregulatie, zoals bleek uit het verschijnsel dat in ons cohort psychologische stress leidde tot voedselinname zonder dat iemand honger heeft. Bovendien werden de hongergevoelens niet verder verlaagd door de voedselinname, hetgeen voor het eerst aantoont dat niet-homeostatische

mechanismen zoals de belonende waarde van voedsel betrokken zijn bij de regulatie van voedselinname tijdens stress. Vorige studies suggereerden een rol voor de belonende waarde van voedsel bij eten als gevolg van stress, maar dit kon niet met zekerheid vastgesteld worden omdat het effect van stress op hongergevoel niet uitgesloten kon worden. Niet iedereen verhoogt zijn voedselinname tijdens stress en grote verschillen bleken dan ook aanwezig te zijn in ons cohort.

Voornamelijk mensen met ontremd eetgedrag en een grote gevoeligheid voor chronische stress eten meer tijdens stress ongeacht hun hongergevoel (**hoofdstuk 9**).

Ontremd eetgedrag afgewisseld met geremd eetgedrag bleek echter weer gerelateerd te zijn aan hyperactiviteit van de HPA-as bij slanke vrouwen. Dit suggereert dat ontremd eetgedrag afgewisseld met geremd eetgedrag chronische stress veroorzaakt. Samen suggereren deze resultaten dat ontremd eetgedrag een belangrijke factor is in de relatie tussen de HPA-as en gewichtsregulatie (**hoofdstuk 10**).

Uit de resultaten beschreven in het tweede deel van het proefschrift kan worden geconcludeerd dat chronische activiteit van de HPA-as als gevolg van stress een groot risico vormt voor de regulatie van lichaamsgewicht. Stress veroorzaakt namelijk voedselinname zonder dat iemand honger heeft, en wel voornamelijk in mensen met ontremd eetgedrag en een grote gevoeligheid voor chronische stress. Ontremd eetgedrag afgewisseld met geremd eetgedrag is gerelateerd aan hyperactiviteit van de HPA-as. Verstoring van HPA-as leidt weer tot opslag van visceraal vet, daarbij een vicieuze cirkel vormend.

Uit de resultaten beschreven in dit proefschrift wordt geconcludeerd dat erfelijke, ouderlijke, gedragsmatige en endocriene factoren invloed hebben op de ontwikkeling en regulatie van lichaamsgewicht bij kinderen en volwassenen.

Dankwoord

Een proefschrift schrijf je niet alleen en dit is de plek om de mensen te bedanken die hieraan hebben bijgedragen.

Allereerst wil ik mijn promotor Margriet en copromotor Arie bedanken. Margriet bedankt voor de snel nagekeken manuscripten, interesse in mijn bezigheden, de vaart erin te houden, het vertrouwen in mijn kunnen en zoveel meer. Arie bedankt voor je heldere momenten als ik weer vast zat, gezellige gesprekken en ongeorganiseerdheid (dan valt dat van mij nog wel mee).

De leden van de beoordelingscommissie bestaande uit prof. Dr. K. Westerterp, prof. Dr. N. Schaper, Dr. W.J. Gerver, Dr. S.E. La Fleur en prof. Dr. J. Wardle wil ik graag bedanken voor het lezen van mijn proefschrift.

Mijn paranimfen Sofie en (super stagiaire) Sanne wil ik bedanken voor alle hulp tijdens experimenten, doorlezen van mijn artikelen, super gezelschap tijdens congressen en natuurlijk het feit dat jullie achter me staan tijdens mijn promotie.

Mijn clustergenoten (Jurriaan, Mieke, Ananda, Freeha, Anneke, Margriet, Rick, Marcel, Alberto, Stijn, Astrid, Kristel, Neeltje, Manuela, Annemiek, Mirjam, Carolien, Natalie, Helene) wil ik bedanken voor de fijne samenwerking en gezellige theepauzes. Verder wil ik ook alle stagiaires bedanken voor jullie hulp bij mijn studies.

Paul en Loek wil ik bedanken voor de technische hulp en Jos, Wendy, Loek en Freek voor het uitvoeren van alle analyses. Natuurlijk wil ik ook de HB secretaresses bedanken voor hun administratieve hulp. And although we only met once in person, large part of my thesis would be absent if not for the analyses of cortisol by Jose Sulon, University of Liege.

Mijn 'nieuwe' vrienden en vriendinnen van de keet, Dames 2 van hockeyclub Horst en mijn 'oude' vriendinnen (Laura, Nathalie, Anouk, Yolanda, Barbara en Atsuko) wil ik bedanken voor de nodige ontspanning in de vorm van gezellige stapavonden, hockeywedstrijden en leuke uitjes.

Wim, Petra en Sandra bedankt voor mijn 'derde' huis in Hegelsom.

Pap, mam, Ilke, Yannicke en schoonbroer Ruud bedankt voor mijn 'tweede' huis in Kaatsheuvel en stiekem doe ik dit allemaal voor jullie zodat jullie trots op mij kunnen zijn.

Lieve Danny, jij bent mijn eerste thuis...

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Martens MJI, Nieuwenhuizen AG, **Rutters F**, Westerterp-Plantenga MS

Curriculum vitae

Femke Rutters was born on September 1st 1982 in Waalwijk, the Netherlands. She completed secondary school at the 'Dr. Moller College' in Waalwijk in 2000. In the same year, she started her study Medical Biology at Radboud University Nijmegen, where she graduated with honor in 2005.

She started working as a PhD-student at the department of Human biology of Maastricht University in November 2005 on a project called 'The role of the HPA axis in the regulation of energy balance', guided by Prof. dr. MS Westerterp-Plantenga and dr. AG Nieuwenhuizen. In July 2007, she was awarded with a New Investigator Award at the Annual Meeting for the Society for the Study of Ingestive Behavior (SSIB) in Steamboat Springs, America. In February 2009, she was awarded a Travel Award of the Van Walree Fonds to visit the Annual Meeting SSIB meeting in Portland, America.

In May 2009 she finished her thesis entitled "Development and regulation of body weight: a genetic, behavioral and neuro-endocrinological approach". In May 2009, she was granted a 'Kootstra Talent Fellowship' scholarship from the Faculty of Health Sciences of Maastricht University to continue her work in the field of body-weight regulation.