INSULIN-LIKE GROWTH FACTOR RECEPTORS IN COLORECTAL CANCER

Gemma Victoria Brierley

B.Sc [Hons.]

CSIRO Molecular and Health Technologies and The Preventative Health National Research Flagship

8

Discipline of Biochemistry School of Molecular and Biomedical Science The University of Adelaide Adelaide, South Australia

A thesis submitted to the University of Adelaide, South Australia in fulfilment of the requirements for the degree of **DOCTOR OF PHILOSOPHY**

Abstract

The IGF system is a crucial regulator of normal growth and development, however dysregulation of the system on multiple levels is associated with the incidence of a wide variety of malignancies including the breast, thyroid, lung, and colon, making the IGF system an important anti-cancer therapeutic target. Due to its role in mediating cellular proliferation, protection from apoptosis, and metastasis, traditional focus has been set on examining the role of the type 1 IGF receptor [IGF1R] in cancer. However there is mounting evidence to suggest the insulin receptor [IR] may also be involved in the potentiation and pathogenesis of cancers.

The observation that IGF-II is overexpressed, compared to normal tissues, by cancers suggests signaling via target receptors by this ligand has important implications on cancer pathogenesis. Indeed, both the IGF1R and IR have been demonstrated to be up-regulated in a variety of malignancies. In regards to IR isoform, the IGF-II binding IR-A is preferentially expressed by a number of cancer cell types. Together with the observation that an autocrine proliferative loop exists between IGF-II and the IR-A in malignant thyrocytes and cultured breast cancer cells, suggests signaling via the IR-A may play a role in cancer cell growth and survival. However, very few studies on the IR-A have been conducted in cells co-expressing the IGF1R. This is mainly due to the difficulties associated with discrimination between signaling arising from IGF1R homodimers, IR-A homodimers, and IGF1R/IR-A hybrid receptors.

It is not known how the IR-A interacts, and functions in conjunction with the other receptors of the IGF system to signal biologically relevant outcomes, especially in terms of anti-cancer therapeutics that aim to block and down-regulate the IGF1R. Current anti-cancer therapies targeting the IGF system have concentrated on blocking IGF signaling via the IGF1R, due mostly to the functional properties of the receptor, but also in part due to the metabolic consequences associated with blockade and inhibition of the IR. This individual targeting of the IGF1R potentially leaves a pathway by which IGF-II secreted by the tumour can circumvent current IGF1R based therapies. Consequently, this thesis investigated whether the IR-A could compensate for the targeted loss of the IGF1R and how the IR-A interacts with the IGF1R in cells co-expressing these two receptors. In addition, the individual ability of the IR isoforms to signal biological outcomes in response to IGF stimulation was assessed.

The main experimental techniques used throughout this body of work included; assessment of protein expression and activation by Western blot, siRNA mediated gene silencing, and measures of cell proliferation, survival, and migration.

The key areas of investigation included:

- Investigation of the individual ability of the IR isoforms to signal biological outcomes in response to IGF stimulation
- 2. Identification of an appropriate cell line model in which to investigate the interactions between the IR-A and IGF1R
- Optimisation of siRNA mediated knock-down of the IR-A and IGF1R in SW480 colorectal adenocarcinoma cells
- 4. Determination of the biological role of the IR-A in SW480 cells co-expressing the IGF1R

The key findings from this work included:

- 1. The IR-A could not compensate for IGF1R depletion in SW480 cells
- Dual silencing of the IR-A and IGF1R indicated signaling via the IGF1R was dominant to signaling via the IR-A in SW480 cells
- Signaling via IR-A/IGF1R hybrid receptors may not be as potent as signaling via IGF1R homodimers
- 4. IGF-I at physiological concentrations can stimulate biological responses via both isoforms of the IR.

Declaration

This thesis contains no material that has been accepted for the award of any other degree or diploma in any University. To the best of my knowledge and belief, it contains no material that has previously been published by any other person except where due reference is made.

I give consent for my thesis, when deposited in the library of the University of Adelaide to be available for loan and photocopying

Gemma Victoria Brierley

Acknowledgements

Here we are, at the end of it all. A year ago I thought I would be struggling to write a very different thesis. What now forms this thesis makes for a pleasant and welcome surprise. I'd like to start by acknowledging and thanking Prof. John Wallace [Department of Biochemistry, University of Adelaide], Dr. Leah Cosgrove [CSIRO, Molecular and Health Technologies, Adelaide], and Dr. Lance Macaulay [CSIRO, Molecular and Health Technologies, Melbourne] for their supervision. Particular thanks to Lance for his thoughtful and detailed feedback throughout the preparation of this document. I also acknowledge the input of Dr. Briony Forbes at the beginning of this project.

The majority of the work outlined in this thesis was performed in the laboratory of Dr. Leah Cosgrove, and was funded by the Preventative Health National Research Flagship. I have been financially supported by the University of Adelaide Department of Biochemistry and the Preventative Health National Research Flagship - I am grateful to both institutions for allowing me the opportunity to pursue this research. In addition, I was afforded the wonderful opportunity to conduct six months of research in the laboratory of Dr. Val Macaulay at the Weatherall Institute of Molecular Medicine, University of Oxford. My time spent in Val's laboratory was second to none. My sincere thanks to Val not only for hosting me in her lab, but also for her mentorship. I learnt so much during my visits, I will always reflect on my time in Oxford with great fondness. My thanks also go to Prof. Richard Head, Director of the P-Health Flagship for funding my travel to Oxford.

Throughout the last four years, the encouragement and support I have received from my family and friends has been unrelenting. "Thank you", doesn't quite seem adequate enough. One question I have been asked a lot over the last few weeks is "if you could go back, would you do it all again?" The answer is "yes - but I would approach it differently". My reasons for beginning a PhD in the first place have not changed, and through circumstance of location I have been extremely fortunate to forge strong friendships with several fun-loving, smart, inspirational, like-minded people. To them I am especially thankful and grateful. For the opportunity to have crossed paths with them, I would do it all again.

And finally, to one in particular, who was often my last bastion of sanity over the last two years [especially this year] - Friday nights at the pub are always something special, never underestimate the power of several pints shared in the presence of exceptional company. Thank you.

He who learns must suffer, Even in our sleep, pain which cannot forget, Falls drop by drop upon the heart, Until, in our own despair, Against our will, Comes wisdom

- Aeschylus

Table of Contents

Abstract		ii
Declaration		l
Acknowledgeme	nts	V
•	ts	
	ns	
Abbreviations		X
Ob (4		
	n and Literature Review	1
1.1 The	IGF System	1
1.1.1	Overview of the IGF Axis	
1.1.2	The Insulin-like Growth Factors	
1.1.3	IGF Binding Proteins	
1.1.4	The Type II IGF Receptor	
1.1.5	The Type I IGF Receptor	
1.1.6	The Insulin Receptor	
1.1.7	IGF1R/IR hybrid receptors	
	rectal Cancer	
1.2.1	Environmental risk factors that contribute to colorectal carcinogenesis	
	molecular role of the IGF system and colorectal cancer	
1.3.1	IGF-I and IGF-II expression	
1.3.2	IGF Receptor Expression	
	ent anti-cancer therapeutic strategies aimed at targeting the IGF system	
1.4.1	Strategies aimed at inhibiting IGF1R protein function	
1.4.2	Strategies aimed at targeting the expression of IGF1R	36
1.5 Scor	pe and Aims of this Thesis	
-		40
Chapter 2 Materials a		40
Materials and 2.1 Mate	nd Methods erials	40
Materials at 2.1 Mate 2.1.1	nd Methods erials Mammalian Cell Culture: Materials, Chemicals, and Reagents	40
2.1 Materials at 2.1.1 2.1.2	erials	40 40
2.1 Materials at 2.1.1 2.1.2 2.1.3	erials	40 40 43
2.1 Materials at 2.1.1 2.1.2 2.1.3 2.1.4	erials	40 43 43
Materials at 2.1 Materials 2.1.1 2.1.2 2.1.3 2.1.4 2.1.5	erials	
Materials at 2.1 Materials at 2.1.1 2.1.2 2.1.3 2.1.4 2.1.5 2.1.6	erials	
Materials at 2.1 Materials at 2.1.1	erials	
2.1 Materials at 2.1.1 2.1.2 2.1.3 2.1.4 2.1.5 2.1.6 2.1.7 2.1.8	erials	
2.1 Materials at 2.1.1	erials	
2.1 Materials at 2.1.1 2.1.2 2.1.3 2.1.4 2.1.5 2.1.6 2.1.7 2.1.8 2.2 Meth 2.2.1	erials	
Materials at 2.1 Mate 2.1.1 2.1.2 2.1.3 2.1.4 2.1.5 2.1.6 2.1.7 2.1.8 2.2 Meth 2.2.1 2.2.1	erials	
2.1 Materials at 2.1.1 2.1.2 2.1.3 2.1.4 2.1.5 2.1.6 2.1.7 2.1.8 2.2 Metric 2.2.1 2.	erials	
2.1 Materials at 2.1.1	maching Methods erials	40 40 43 43 44 44 46 46 46 46 46
2.1 Materials at 2.1.1	erials	
Materials at 2.1 Materials at 2.1.1 Materials 2.1.1 2.1.2 2.1.3 2.1.4 2.1.5 2.1.6 2.1.7 2.1.8 2.2 Metricular 2.2.1 2.2.1 2.2.1 2.2.1 2.2.2 2.2 2.2 2	And Methods Prials	
Materials at 2.1 Materials at 2.1.1 Materials 2.1.1 2.1.2 2.1.3 2.1.4 2.1.5 2.1.6 2.1.7 2.1.8 2.2 Metric 2.2.1 2.2.1 2.2.1 2.2.2 2.2 2.2 2.2 2.2	And Methods Prials	
Materials at 2.1 Materials at 2.1.1 Materials 2.1.1 2.1.2 2.1.3 2.1.4 2.1.5 2.1.6 2.1.7 2.1.8 2.2 Metric 2.2.1 2.2	And Methods Prials	
Materials at 2.1 Materials at 2.1.1 Materials 2.1.1 2.1.2 2.1.3 2.1.4 2.1.5 2.1.6 2.1.7 2.1.8 2.2 Metricular 2.2.1	And Methods Prials	
Materials at 2.1 Materials at 2.1.1 Materials 2.1.1 2.1.2 2.1.3 2.1.4 2.1.5 2.1.6 2.1.7 2.1.8 2.2 Metricular 2.2.1	erials	
2.1 Materials at 2.1.1	And Methods Prials	
2.1 Materials at 2.1.1	And Methods Prials	40 40 43 44 44 46 46 46 46 47 47 47 47 47 48 48 48 48
Materials at 2.1 Materials at 2.1.1 Materials 2.1.1 2.1.2 2.1.3 2.1.4 2.1.5 2.1.6 2.1.7 2.1.8 2.2 Metric 2.2.1 2.2.1 2.2.1 2.2.2 2.2.2 2.2.2 2.2.2 2.2	And Methods Prials	
Materials at 2.1 Materials at 2.1.1 Materials 2.1.1 2.1.2 2.1.3 2.1.4 2.1.5 2.1.6 2.1.7 2.1.8 2.2 Metric 2.2.1 2.2.1 2.2.1 2.2.2 2.2 2.2 2.2 2.2	And Methods Prials	
Materials at 2.1 Mate 2.1.1 2.1.2 2.1.3 2.1.4 2.1.5 2.1.6 2.1.7 2.1.8 2.2 Meth 2.2.1 2.2.1 2.2.1 2.2.1 2.2.1 2.2.1 2.2.1 2.2.2 2.2	erials	40 40 43 44 44 46 46 46 46 47 47 47 47 47 48 48 48 49 50
Materials at 2.1 Materials at 2.1.1 Materials 2.1.1 2.1.2 2.1.3 2.1.4 2.1.5 2.1.6 2.1.7 2.1.8 2.2 Meth 2.2.1 2.2.1 2.2.1 2.2.2 2.2 2.2 2.2 2.2	erials	40 43 43 44 44 46 46 46 46 47 47 47 47 48 48 48 49 50 50 50
Materials at 2.1 Materials at 2.1.1 Materials at 2.1.1 2.1.2 2.1.3 2.1.4 2.1.5 2.1.6 2.1.7 2.1.8 2.2 Metricular 2.2.1 2.2.1 2.2.1 2.2.1 2.2.2 2.	erials	
2.1 Materials at 2.1.1	erials	
Materials at 2.1 Materials at 2.1.1 Materials 2.1.1 2.1.2 2.1.3 2.1.4 2.1.5 2.1.6 2.1.7 2.1.8 2.2 Metric 2.2.1 2.2.1 2.2.1 2.2.2 2.2	erials	
Materials at 2.1 Materials at 2.1.1 Materials at 2.1.1 2.1.2 2.1.3 2.1.4 2.1.5 2.1.6 2.1.7 2.1.8 2.2 Metric 2.2.1	And Methods Prials	
2.1 Materials at 2.1.1	And Methods Prials	
2.1 Materials at 2.1.1 (2.1.2) (2.1.3) (2.1.4) (2.1.5) (2.1.6) (2.1.7) (2.1.8) (2.2.1)	mammalian Cell Culture: Materials, Chemicals, and Reagents	40 40 43 44 44 44 46 46 46 46 47 47 47 47 50 50 50 50 50 50 50 50 50 50 50 50 50
2.1 Materials at 2.1.1	And Methods Prials	40 40 43 44 44 44 46 46 46 46 47 47 47 47 50 50 50 50 50 50 50 50 50 50 50 50 50

	2.2.2.3	Synthesis of cDNA	59
	2.2.2.4	Polymerase Chain Reaction	
		Agarose gel electrophoresis of RNA/DNA samples	
		ds: Protein Chemistry	
		Lysis of mammalian cells	
	2.2.3.2	Determination of protein concentration	
	2.2.3.3	Immunopreciptiation of proteins expressed in mammalian cells	62
	2.2.3.4	Western blot analysis	63
		Labeling of cell surface proteins with antibodies for detection by flow cytometry	
		IR-A and IR-B phosphorylation and activation of intracellular signaling molecules in respon	
		insulin, IGF-I, IGF-II, and IGF chimeras	
	2.2.3.6.1	Materials	_
	2.2.3.6.2	Methods	
	2.2.3.6.3	Binding analysis of IGF chimeras to insulin receptor isoforms	65
0 1-			07
Ch		centrations of IGF-I can stimulate biological responses via the IR	67
	3.1.1 IGF CI	himeras	68
	3.1.2 Bindin	g specificity of the IGFs to the IR-A and IR-B is regulated by their C and D domains	71
		ion of IR-A and IR-B tyrosine autophosphorylation by insulin, IGF-II, IGF-I, and IGF IGF ch	
		ion of Y960 [IR-A] and Y972 [IR-B] phosphorylation by insulin, IGF-II, IGF-I, and IGF chime	
		, IGF-II, IGF-I, and IGF C domain chimera induced phosphorylation of IRS-1 and IRS-2	
		nd IR-B	
	3.1.6 Ligano	I stimulated activation of Akt/PKB via the IR-A and IR-B	79
	3.1.7 Ligano	stimulated phosphorylation of Erk-1/2 via the IR-A and IR-B	79
	3.2 Results		82
		tion of RIRA and RIRB cells	_
		of ligand to rescue RIRA and RIRB cells from butyrate-induced apoptosis	
		, IGF-II, IGF-I, and IGF C-domain chimera stimulated RTRA chemotaxis	
		le of IRS-2 in IGF-I mediated effects through the IR	90
	3.2.4 The ro	to of the 2 th for a mediated checks through the free transfer	
			90
	3.3 Discussion.	S	
Ch	3.3 Discussion. 3.4 Conclusions	5	95
Ch	3.3 Discussion. 3.4 Conclusions napter 4	S	95
Ch	3.3 Discussion. 3.4 Conclusions napter 4 Identification ar of the IR-A in cell	ond characterisation of a cell line model for the investigation of the biologicals s co-expressing the IGF1R	95 96 al role
Ch	3.3 Discussion. 3.4 Conclusions napter 4 Identification ar of the IR-A in cell. 4.1 Introduction	nd characterisation of a cell line model for the investigation of the biologicals co-expressing the IGF1R	95 96 al role 96
Ch	3.3 Discussion. 3.4 Conclusions napter 4 Identification ar of the IR-A in cell 4.1 Introduction 4.2 Results	nd characterisation of a cell line model for the investigation of the biologicals co-expressing the IGF1R	95 96 96 97
Ch	3.3 Discussion. 3.4 Conclusions hapter 4 Identification ar of the IR-A in cell 4.1 Introduction 4.2 Results 4.2.1 Charac	ond characterisation of a cell line model for the investigation of the biological sco-expressing the IGF1R cterisation of colorectal adenocarcinoma cell IGF1R and IR gene expression by RT-PCR	95 96 96 97 97
Ch	3.3 Discussion. 3.4 Conclusions hapter 4 Identification ar of the IR-A in cell 4.1 Introduction 4.2 Results 4.2.1 Charac	nd characterisation of a cell line model for the investigation of the biologicals co-expressing the IGF1R	95 96 96 97 97
Ch	3.3 Discussion. 3.4 Conclusions napter 4 Identification ar of the IR-A in cell 4.1 Introduction 4.2 Results 4.2.1 Chara 4.2.2 Identification are of the IR-A in cell 4.1 Introduction and the IR-A in cell 4.1 Introduction and the IR-A in cell 4.2 Introduction adeno	od characterisation of a cell line model for the investigation of the biological sco-expressing the IGF1R cterisation of colorectal adenocarcinoma cell IGF1R and IR gene expression by RT-PCR ication of IGF1R, IR, and IGF1R/IR hybrid receptor protein expression in coloractionan cells	95 96 al role 96 97 97 lorectal
Ch	3.3 Discussion. 3.4 Conclusions napter 4 Identification ar of the IR-A in cell 4.1 Introduction 4.2 Results 4.2.1 Chara 4.2.2 Identification are of the IR-A in cell 4.1 Introduction and the IR-A in cell 4.1 Introduction and the IR-A in cell 4.2 Introduction adeno	od characterisation of a cell line model for the investigation of the biological sco-expressing the IGF1R cterisation of colorectal adenocarcinoma cell IGF1R and IR gene expression by RT-PCR ication of IGF1R, IR, and IGF1R/IR hybrid receptor protein expression in coloractionan cells	95 96 al role 96 97 97 lorectal
Ch	3.3 Discussion. 3.4 Conclusions napter 4 Identification ar of the IR-A in cell 4.1 Introduction 4.2 Results 4.2.1 Chara 4.2.2 Identification adeno 4.2.3 Colore	cterisation of IGF1R, IR, and IGF1R/IR hybrid receptor protein expression in coloraction adenocarcinoma cells	95 96 al role 96 97 97 lorectal 99 101
Ch	3.3 Discussion. 3.4 Conclusions napter 4 Identification ar of the IR-A in cell. 4.1 Introduction 4.2 Results 4.2.1 Character of the IR-A in cell. 4.2.1 Character of the IR-A in cell. 4.2.2 Identification are of the IR-A in cell. 4.2.3 Colore of the IR-A in cell.	cterisation of colorectal adenocarcinoma cell IGF1R and IR gene expression by RT-PCR ication of IGF1R, IR, and IGF1R/IR hybrid receptor protein expression in colorectal adenocarcinoma cells	95 96 al role 96 97 97 lorectal 99 101
Ch	3.3 Discussion. 3.4 Conclusions napter 4 Identification ar of the IR-A in cell. 4.1 Introduction 4.2 Results 4.2.1 Characteristics adeno 4.2.2 Identification ar of the IR-A in cell. 4.1 Effect 4.2.3 Colore 4.2.4 Effect 4.2.5 Butyra	cterisation of colorectal adenocarcinoma cell IGF1R and IR gene expression by RT-PCR ication of IGF1R, IR, and IGF1R/IR hybrid receptor protein expression in colorectal adenocarcinoma cells	95 96 al role 97 lorectal 99 101 101
Ch	3.3 Discussion. 3.4 Conclusions napter 4	cterisation of colorectal adenocarcinoma cell IGF1R and IR gene expression by RT-PCR cication of IGF1R, IR, and IGF1R/IR hybrid receptor protein expression in colorectal adenocarcinoma cells	95 96 96 97 97 lorectal 99 101 106 106
Ch	3.3 Discussion. 3.4 Conclusions napter 4	cterisation of colorectal adenocarcinoma cell IGF1R and IR gene expression by RT-PCR cication of IGF1R, IR, and IGF1R/IR hybrid receptor protein expression in coloracrinoma cells	95 96 96 97 97 lorectal 99 101 106 106 106
Ch	3.3 Discussion. 3.4 Conclusions napter 4	cterisation of colorectal adenocarcinoma cell IGF1R and IR gene expression by RT-PCR cication of IGF1R, IR, and IGF1R/IR hybrid receptor protein expression in colorectal adenocarcinoma cells	95 96 96 97 97 lorectal 99 101 106 106 106
Ch	3.3 Discussion. 3.4 Conclusions napter 4	cterisation of colorectal adenocarcinoma cell IGF1R and IR gene expression by RT-PCR cication of IGF1R, IR, and IGF1R/IR hybrid receptor protein expression in coloracrinoma cells	95 96 al role 96 97 97 lorectal 99 101 106 106 109 110
Ch	3.3 Discussion. 3.4 Conclusions napter 4	cterisation of colorectal adenocarcinoma cell IGF1R and IR gene expression by RT-PCR cication of IGF1R, IR, and IGF1R/IR hybrid receptor protein expression in coloracrinoma cells	95 96 al role 96 97 97 lorectal 101 106 106 109 110
Ch	3.3 Discussion. 3.4 Conclusions napter 4	cterisation of colorectal adenocarcinoma cell IGF1R and IR gene expression by RT-PCR cication of IGF1R, IR, and IGF1R/IR hybrid receptor protein expression in coloracrinoma cells	95 96 96 97 97 lorectal 101 106 106 109 110
Ch	3.3 Discussion. 3.4 Conclusions napter 4	cterisation of colorectal adenocarcinoma cell IGF1R and IR gene expression by RT-PCR cication of IGF1R, IR, and IGF1R/IR hybrid receptor protein expression in coloracrinoma cells	95 96 al role 96 97 97 lorectal 101 106 106 109 110
Ch	3.3 Discussion. 3.4 Conclusions napter 4	cterisation of colorectal adenocarcinoma cell IGF1R and IR gene expression by RT-PCR cication of IGF1R, IR, and IGF1R/IR hybrid receptor protein expression in coloracrinoma cells	95 96 al role 96 97 97 lorectal 101 106 106 109 110
	3.3 Discussion. 3.4 Conclusions hapter 4	cterisation of colorectal adenocarcinoma cell IGF1R and IR gene expression by RT-PCR cterisation of colorectal adenocarcinoma cell IGF1R and IR gene expression by RT-PCR cterisation of IGF1R, IR, and IGF1R/IR hybrid receptor protein expression in colorarcinoma cells ctal adenocarcinoma cell surface expression of IGF1R and IR	9596 al role969797 lorectal101106109110
	3.3 Discussion. 3.4 Conclusions hapter 4	cterisation of colorectal adenocarcinoma cell IGF1R and IR gene expression by RT-PCR cterisation of colorectal adenocarcinoma cell IGF1R and IR gene expression by RT-PCR cterisation of IGF1R, IR, and IGF1R/IR hybrid receptor protein expression in colorarcinoma cells ctal adenocarcinoma cell surface expression of IGF1R and IR	9596 al role969797 lorectal99101106109110
	3.3 Discussion. 3.4 Conclusions hapter 4	cterisation of colorectal adenocarcinoma cell IGF1R and IR gene expression by RT-PCR cication of IGF1R, IR, and IGF1R/IR hybrid receptor protein expression in coloracrinoma cells cital adenocarcinoma cell surface expression of IGF1R and IR of IGF-II, IGF-II, and insulin on butyrate-treated colorectal adenocarcinoma cell viability te induces apoptosis in SW480 cells with concomitant decrease in cell viability IGF-II, and insulin stimulates SW480 cell chemotaxis tion of IR and IGF1R by ligand in SW480 cells	9596 al role969797 lorectal99101106109110
	3.3 Discussion. 3.4 Conclusions hapter 4	cterisation of colorectal adenocarcinoma cell IGF1R and IR gene expression by RT-PCR cterisation of colorectal adenocarcinoma cell IGF1R and IR gene expression by RT-PCR cterisation of IGF1R, IR, and IGF1R/IR hybrid receptor protein expression in colorarcinoma cells ctal adenocarcinoma cell surface expression of IGF1R and IR	9596 al role969797 lorectal99101106109110
	3.3 Discussion. 3.4 Conclusions hapter 4	cterisation of colorectal adenocarcinoma cell IGF1R and IR gene expression by RT-PCR ication of IGF1R, IR, and IGF1R/IR hybrid receptor protein expression in coloractal adenocarcinoma cells ication of IGF-I, IGF-II, and insulin on butyrate-treated colorectal adenocarcinoma cell viability ite induces apoptosis in SW480 cells with concomitant decrease in cell viability ite induces apoptosis in SW480 cells with concomitant decrease in cell viability ite induces apoptosis in SW480 cells cells with concomitant decrease in cell viability item of IGF1R and IGF1R by ligand in SW480 cells	95 96 96 97 97 lorectal 101 106 109 112
	3.3 Discussion. 3.4 Conclusions hapter 4	cterisation of colorectal adenocarcinoma cell IGF1R and IR gene expression by RT-PCR cication of IGF1R, IR, and IGF1R/IR hybrid receptor protein expression in coloracrinoma cells cital adenocarcinoma cell surface expression of IGF1R and IR of IGF-II, IGF-II, and insulin on butyrate-treated colorectal adenocarcinoma cell viability te induces apoptosis in SW480 cells with concomitant decrease in cell viability IGF-II, and insulin stimulates SW480 cell chemotaxis tion of IR and IGF1R by ligand in SW480 cells	95 96 96 97 97 lorectal 101 106 109 112
	3.3 Discussion. 3.4 Conclusions hapter 4	cterisation of colorectal adenocarcinoma cell IGF1R and IR gene expression by RT-PCR ication of IGF1R, IR, and IGF1R/IR hybrid receptor protein expression in coloractal adenocarcinoma cells ication of IGF-I, IGF-II, and insulin on butyrate-treated colorectal adenocarcinoma cell viability ite induces apoptosis in SW480 cells with concomitant decrease in cell viability ite induces apoptosis in SW480 cells with concomitant decrease in cell viability ite induces apoptosis in SW480 cells cells with concomitant decrease in cell viability item of IGF1R and IGF1R by ligand in SW480 cells	95 96 96 97 97 lorectal 101 106 109 112 113
	3.3 Discussion. 3.4 Conclusions hapter 4	and characterisation of a cell line model for the investigation of the biologicals co-expressing the IGF1R cterisation of colorectal adenocarcinoma cell IGF1R and IR gene expression by RT-PCR ication of IGF1R, IR, and IGF1R/IR hybrid receptor protein expression in coloractinoma cells ctal adenocarcinoma cell surface expression of IGF1R and IR of IGF-I, IGF-II, and insulin on butyrate-treated colorectal adenocarcinoma cell viability te induces apoptosis in SW480 cells with concomitant decrease in cell viability IGF-II, and insulin stimulates SW480 cell chemotaxis tion of IR and IGF1R by ligand in SW480 cells	95 96 al role 96 97 97 lorectal 101 106 109 112 113 113
	3.3 Discussion. 3.4 Conclusions hapter 4	cterisation of colorectal adenocarcinoma cell IGF1R and IR gene expression by RT-PCR cation of IGF1R, IR, and IGF1R/IR hybrid receptor protein expression in colorarcinoma cells cation acells ctal adenocarcinoma cell surface expression of IGF1R and IR. of IGF-I, IGF-II, and insulin on butyrate-treated colorectal adenocarcinoma cell viability IGF-II, and insulin stimulates SW480 cell with concomitant decrease in cell viability IGF-II, and IGF1R by ligand in SW480 cells RNA mediated knock-down of the IGF1R and IR-A in SW480 cells B Discussion validation of siRNAs targeting the IGF1R and IR.	9596 al role969797 lorectal101106109112113
	3.3 Discussion. 3.4 Conclusions hapter 4	cterisation of colorectal adenocarcinoma cell IGF1R and IR gene expression by RT-PCR cation of IGF1R, IR, and IGF1R/IR hybrid receptor protein expression in colorarcinoma cells ctal adenocarcinoma cell surface expression of IGF1R and IR of IGF-I, IGF-II, and insulin on butyrate-treated colorectal adenocarcinoma cell viability te induces apoptosis in SW480 cells with concomitant decrease in cell viability IGF-II, and insulin stimulates SW480 cell chemotaxis tion of IR and IGF1R by ligand in SW480 cells BY A mediated knock-down of the IGF1R and IR-A in SW480 cells I Discussion validation of siRNAs targeting the IGF1R and IR sation of non-silencing control siRNAs and transfection reagent used for delivery of siRN.	95 96 al role 96 97 97 lorectal 101 106 109 112 113 115 As into
	3.3 Discussion. 3.4 Conclusions hapter 4	cterisation of colorectal adenocarcinoma cell IGF1R and IR gene expression by RT-PCR ication of IGF1R, IR, and IGF1R/IR hybrid receptor protein expression in colorectal adenocarcinoma cells ication of IGF1, IGF-II, and insulin on butyrate-treated colorectal adenocarcinoma cell viability ite induces apoptosis in SW480 cells with concomitant decrease in cell viability IGF-II, and insulin stimulates SW480 cells chemotaxis ition of IR and IGF1R by ligand in SW480 cells RNA mediated knock-down of the IGF1R and IR-A in SW480 cells I Discussion I Discussion I Discussion I Discussion control siRNAs targeting the IGF1R and IR I Discussion consideration of siRNAs targeting the IGF1R and IR I Discussion consideration of siRNAs targeting the IGF1R and IR I Discussion consideration of siRNAs and transfection reagent used for delivery of siRN.	9596 al role969797 lorectal101106106112113115 As into121
	3.3 Discussion. 3.4 Conclusions napter 4	and characterisation of a cell line model for the investigation of the biological sco-expressing the IGF1R cterisation of colorectal adenocarcinoma cell IGF1R and IR gene expression by RT-PCR cterisation of IGF1R, IR, and IGF1R/IR hybrid receptor protein expression in coloraction cells ctal adenocarcinoma cell surface expression of IGF1R and IR ctal adenocarcinoma cell surface expression of IGF1R and IR ctal adenocarcinoma cell viability te induces apoptosis in SW480 cells with concomitant decrease in cell viability IGF-II, and insulin stimulates SW480 cell chemotaxis tion of IR and IGF1R by ligand in SW480 cells	95 96 al role 96 97 lorectal 101 106 109 113 112 115 115 115 121 124
	3.3 Discussion. 3.4 Conclusions napter 4	and characterisation of a cell line model for the investigation of the biologics is co-expressing the IGF1R cterisation of colorectal adenocarcinoma cell IGF1R and IR gene expression by RT-PCR ctation of IGF1R, IR, and IGF1R/IR hybrid receptor protein expression in coloraction carcinoma cells ctal adenocarcinoma cell surface expression of IGF1R and IR of IGF-II, and insulin on butyrate-treated colorectal adenocarcinoma cell viability te induces apoptosis in SW480 cells with concomitant decrease in cell viability IGF-II, and insulin stimulates SW480 cell chemotaxis tion of IR and IGF1R by ligand in SW480 cells RNA mediated knock-down of the IGF1R and IR. Salidation of siRNAs targeting the IGF1R and IR sation of non-silencing control siRNAs and transfection reagent used for delivery of siRN. ocells sation of HiPerFect transfection conditions sation of Post-transfection survival assays	95 96 al role 96 97 97 lorectal 101 106 109 113 115 115 As into 121 124 133
	3.3 Discussion. 3.4 Conclusions napter 4	and characterisation of a cell line model for the investigation of the biologics is co-expressing the IGF1R cterisation of colorectal adenocarcinoma cell IGF1R and IR gene expression by RT-PCR ctation of IGF1R, IR, and IGF1R/IR hybrid receptor protein expression in coloraction carcinoma cells ctal adenocarcinoma cell surface expression of IGF1R and IR of IGF-II, and insulin on butyrate-treated colorectal adenocarcinoma cell viability te induces apoptosis in SW480 cells with concomitant decrease in cell viability IGF-II, and insulin stimulates SW480 cell chemotaxis tion of IR and IGF1R by ligand in SW480 cells RNA mediated knock-down of the IGF1R and IR. Salidation of siRNAs targeting the IGF1R and IR sation of non-silencing control siRNAs and transfection reagent used for delivery of siRN. ocells sation of HiPerFect transfection conditions sation of Post-transfection survival assays	95 96 al role 96 97 97 lorectal 101 106 109 113 115 115 As into 121 124 133
	3.3 Discussion. 3.4 Conclusions napter 4	and characterisation of a cell line model for the investigation of the biological sco-expressing the IGF1R cterisation of colorectal adenocarcinoma cell IGF1R and IR gene expression by RT-PCR cterisation of IGF1R, IR, and IGF1R/IR hybrid receptor protein expression in coloraction cells ctal adenocarcinoma cell surface expression of IGF1R and IR ctal adenocarcinoma cell surface expression of IGF1R and IR ctal adenocarcinoma cell viability te induces apoptosis in SW480 cells with concomitant decrease in cell viability IGF-II, and insulin stimulates SW480 cell chemotaxis tion of IR and IGF1R by ligand in SW480 cells	95 96 al role 96 97 lorectal 101 106 109 113 115 115 As into 121 123 133 133

			135
Asse	ssme	nt of the biological role of the IR-A in cells co-expressing the IGF1R	
0.4			405
6.1		oduction	
6.2		sults	
6.2 6.2		The effect of transient siRNA transfection on SW480 cell surface expression of IGF1R and IR-A The effect of IGF1R and IR-A knock-down on clonogenic survival and sensitivity to butyrate	
6.2	2.3	The effect of knocking-down IGF1R expression on ligand stimulated cell survival from butyrate ind apoptosis	.142
6.2 6.2		The effect of silencing the IR-A on ligand stimulated cell survival from butyrate-induced apoptosis The effect of dual knock-down of IGF1R and IR-A on ligand stimulated cell survival from buty induced apoptosis	rate-
6.2	2.6	The effect of IGF1R and IR-A knock-down on hybrid receptor formation	
6.2	2.7	The effect of siRNA mediated receptor knock-down on ligand stimulated receptor tyro	
6.3	Disc	cussion	153
6.4		iclusions	
		ussion	164
7.1	Sco	pe and Aims of Thesis	164
7.2	Sun	nmary of Findings	164
7.3	Disc	cussion	166
7.4		ure Directions	
7.5	Cor	clusions	171
Referenc	es		172
Appendix	(188
Publi	icatio	ns arising from this thesis	

List of Publications

Peer reviewed publications [see appendix]

* Denotes joint first authorship

Denley A*, Carroll JM*, **Brierley GV**, Cosgrove LJ, Wallace JC, Forbes BE, Roberts CT Jr. Differential activation of insulin receptor substrates 1 and 2 by insulin-like growth factor-activated insulin receptors. *Molecular Cell Biology* 2007 May; 27[10]: 3569 - 77

Denley A, **Brierley GV**, Carroll JM, Lindenberg A, Booker GW, Cosgrove LJ, Wallace JC, Forbes BE, Roberts CT Jr. Differential activation of insulin receptor isoforms by insulin-like growth factors is determined by the C domain. *Endocrinology* 2006 Feb; 147[2]: 1029 – 1036

Conference Proceedings

Brierley GV, Forbes BE, Macaulay L, Siddle K, Wallace JC, Cosgrove LJ. The biological role of insulin-like growth factors signalling via IR-A and IR-A/IGF1R hybrid receptors in cell survival, migration, and invasion. *The role of the IGF system in Cancer – Taormina, Sicily, Italy. November* 2005

Brierley GV, Forbes BE, Wallace JC, Cosgrove LJ. The biological role of insulin-like growth factors signalling via IR-A and IR-A/IGF1R hybrid receptors in cell survival, migration, and invasion. Gordon Research Conference: Insulin-like growth factors in physiology and disease – Ventura, California, U.S.A. April 2005

Abbreviations

٥С degrees celcius microgram μg microlitre μΙ micrometre μm . 3' three prime 5' five prime

95% CI ninety five percent confidence interval

bp base pair antibody Ab **ACF** aberrant crypt foci

APC Adenomatosis Polyposis coli ASO anti-sense oligonucleotide ATP adenosine triphosphate BMI body mass index

butyrate BuA

BRET bioluminescence resonance energy transfer

BSA bovine serum albumin CO_2 carbon dioxide cDÑA complementary DNA compared with c.f COX-2 cyclooxygenase-2 cystine-rich CR CRC colorectal cancer carboxy-terminal tail CT C-terminus Carboxyl-terminus

2'-Deoxyadenosine 5'-triphosphate dATP

DEPC diethyl pyrocarbonate

DEPC-MQ H₂O dethyl pyrocarbonate milliQ water **DMEM** Dulbecco's modified eagles medium

DMSO dimethyl sulfoxide deoxyribonucleic acid DNA

decilitre dL

dATP 2'-deoxyadenosine 5'-triphosphate 2'-deoxycytidine 5'-triphosphate dCTP dGTP 2'-deoxyguanosine 5'-triphosphate dTTP 2'-deoxythymidine 5'-triphosphate deoxynucleotide triphosphates dNTPs DTT dithiothreitol

 EC_{50} 50% effective concentration **EDTA** ethylene diamine tetra-acetic acid Epidermal Growth Factor **EGF EGFR** Epidermal Growth Factor Receptor

EPIC European Prospective Investigation into Cancer and Nutrition

extracellular signalling-related kinase - 1/2 Erk-1/2

Europium Eu

fluorescent-activated cell sorting **FACS**

foetal calf serum **FCS** fluorescein isothiocyanate FITC fibronectin domain F_{n} GH growth hormone

Grb2 Growth factor receptor-bound protein 2

 H_2O water

HEPES 4-(2-hydroxyethyl)-1-piperazineethanesulfonic acid

HER-2 Human Epidermal Growth Factor-2 Hybrid-A IGF1R/IR-A hybrid receptor Hybrid-B IGF1R/IR-B hybrid receptor

ΙŔ immunoblot

 IC_{50} IGF1R concentration of inhibitor which reduces binding by 50%

Type 1 IGF receptor
Type 1 IGF receptor alpha-subunit
Type 1 IGF receptor beta-subunit $\mathsf{IGF1R}_{\alpha}$ IGF1R₈

IGF2R

Type 2 IGF receptor
Insulin-like growth factor binding proteins **IGFBPs**

Insulin-like growth factor I IGF-I

IGF-I containing the IGF-II C domain IGF-I CII IGF-I containing the IGF-II D domain IGF-I containing the IGF-II C and D domains IGF-I DII IGF-I CIIDII

IGF-II Insulin-like growth factor II

IGF-II CI IGF-II containing the IGF-I C domain IGF-II containing the IGF-I D domain IGF-II DI IGF-II containing the IGF-I C and D domains IGF-II CIDI

insert domain Ins ΙP immunoprecipitate IR Insulin receptor

 $\begin{array}{ll} IR_{\alpha} & \quad Insulin \ receptor \ alpha-subunit \\ IR_{\beta} & \quad Insulin \ receptor \ beta-subunit \end{array}$

IR-A A isoform of the human insulin receptor [exon 11-]
IR-B B isoform of the human insulin receptor [exon 11+]

IRS-1 Insulin Receptor Substrate -1
IRS-2 Insulin Receptor Substrate -2
JM juxtamembrane domain

kDa kilodalton
L1 large domain 1
L2 large domain 2
L-chain light chain
LHS left-hand side
LOI loss of imprinting

M molar

mAb monoclonal antibody

MALDI-TOF matrix-assisted laser desorption/ionisation – time of flight

MAPK Mitogen-activated protein kinase

Max. maximum mg milligram

MgCl₂ magnesium chloride

ml millilitre mM millimolar

MMP matrix metalloprotienase

MOPS 3-(N-morpholino)propanesulfonic acid

mRNA messenger ribonucleic acid

N/A not applicable
NaCl sodium chloride
NaF sodium fluoride
Na₄O₇P₂ sodium orthovanidate
Na₃PO₄ sodium pyrophosphate

ng nanogram
NH₄ ammonia
nm nanometer
nM nanomolar
n/s not significant

NSAIDs non-steroidal anti-inflammatory drugs

nt nucleotide N-terminus amino-terminus

OD₂₆₀ optical density at 260 nm optical density at 280 nm

p53 protein 53 pAb polyclonal antibody

PAGE polyacrylamide gel electrophoresis
PBS phosphate buffered saline
PCR polymerase chain reaction
PFA paraformaldehyde
PI3K phosphoinositide kinase-3
pIRS-1 phosphorylated IRS-1
pIRS-2 phosphorylated IRS-2
PKB protein kinase B

PMSF phenylmethanesulphonyl fluoride PTEN phosphatase and tensin homologue

R IGF1R negative NIH 3T3-like mouse fibroblasts

RIGF1R IGF1R negative NIH 3T3-like mouse fibroblasts recombinantly expressing human IGF1R RIRA IGF1R negative NIH 3T3-like mouse fibroblasts recombinantly expressing human IR-A RIRB IGF1R negative NIH 3T3-like mouse fibroblasts recombinantly expressing human IR-B

RHS right-hand side

RISC RNA-inducing silencing complex

RNA ribonucleic acid RNAi RNA interference rpm revolutions per minute

RR relative risk

RT-PCR reverse-transcriptase PCR SCFA short-chain fatty acid SDS sodium dodecyl sulphate SH2 Src-homology domain siRNA short interfering RNA SEM standard error of the mean TAE Tris-acetate EDTA TK tyrosine-kinase domain

Tris-Cl Tris (hydroxymethyl) aminomethane

Tyr tyrosine

VEGF vascular epidermal growth factor

vs. versus
WB Western blot
WCL Whole cell lysate