

Copyright is owned by the Author of the thesis. Permission is given for a copy to be downloaded by an individual for the purpose of research and private study only. The thesis may not be reproduced elsewhere without the permission of the Author.

# Pre-implantation maternal uterine effects on embryo growth and development: An investigation using models of maternal constraint in sheep

A thesis presented in fulfilment of the requirements for the degree of:

**Doctor of Philosophy** 

in

**Animal Science** 

Massey University, Turitea, Palmerston North

New Zealand.

Lisanne Monique Fermin

2017

### Primary supervisor

Dr. Sarah J. Pain

### Co – supervisors

Professor Paul R. Kenyon

Professor Hugh T. Blair

### Advisors

Professor Frank H. Bloomfield Associate Professor Peter K. Dearden

### Abstract

Prenatal development and growth are critical to survival of the fetus and neonate. Recent evidence suggests that a critical period for determining growth is the pre-implantation period of pregnancy during which differentiation, organogenesis and development of the embryo occur and the embryo is considerably vulnerable to uterine environmental factors. The objectives of the present study were to examine the effects of restrictive uterine environments on embryo development using two sheep models of maternal constraint: litter size and dam size, and to identify embryonic and maternally-driven mechanisms that regulate development of the peri-implantation sheep embryo.

Morphometric analysis (embryo length, width and heart bulge width) of the embryos in peri-implantation single and twin embryos was inconclusive; as was the transcriptomics analysis of whole embryos using RNA-seq to examine differential gene expression that may be responsible for differential regulation of growth.

In a dam size model, large-breed Suffolk embryos gestated in small-breed Cheviot ewes (constrained environment) were smaller than Suffolk embryos gestated in Suffolk ewes (control) at day 19 of pregnancy, confirming previous findings that maternal constraint is evident in early pregnancy when limitations of space are not of consequence. Progesterone administered in the post-ovulatory period, day 0 to 6, alleviates this apparent constraint such that Suffolk embryos gestated in Cheviot ewes that received progesterone are larger than those gestated in Cheviot ewes that did not. Further, differential gene expression analysis of maternal uterine tissues showed that at day 6 and day 19 endometrial genes that encode for histotroph secretion and uterine receptivity are altered by post-ovulatory progesterone administration. Timing of administration of progesterone is critical not only to embryo growth but also to embryo survival. There were lower pregnancy rates in the ewes that received progesterone from day 0 than those that received progesterone from day 2.

The results of this thesis indicate that progesterone exerts its effects by regulation of genes that encode for uterine structural and secretory activity to advance the uterus. This likely forces the asynchronous embryo to accelerate its growth in order to adapt to its environment. These findings contribute to the knowledge of the regulatory mechanisms controlling early embryo growth and present a platform within the livestock industry and human reproductive technology practice to manipulate embryo growth to improve survival of offspring.

## Acknowledgements

With an overwhelming sense of gratitude I would like to thank all those who supported and encouraged me through this journey to the successful completion of this thesis.

I would like to thank my supervisors, Dr Sarah Pain, Professor Paul Kenyon and Professor Hugh Blair, firstly for the opportunity to work with such a great team, but without whose encouragement and support this thesis would not have been completed. Sarah, your patience, advice, support and guidance throughout all the stages of this process, from field work to write up has not gone un-noticed and for all of this I am truly grateful. Paul, your unwavering support, encouragement and advice are appreciated. Hugh, I cannot thank you enough for pushing me just that little bit further, by adding another perspective to my thought process and challenging me to think outside the box. This has made me a better researcher and writer.

Thank you to Professor Frank Bloomfield, Associate Professor Peter Dearden, Professor Patrick Morel, Dr Elizabeth Duncan, Dr Matthew Perrott, Dr Kristene Gedye, Dr Ana Meikle, Dr Mark Oliver, and Hui Hui Phua. All of you provided invaluable support at various levels of this thesis, from technical and laboratory (molecular biology and microscopy) skills, statistical analysis, and write-up. I would not have been able to accomplish this without all of your input.

I am very grateful to IVABS/International Sheep Research Centre team for all their help, particularly with the field work: Dean Burnham, Catriona Jenkins and Geoff Purchas for their assistance in managing those crazy Cheviot and Suffolk sheep and your dedication to the smooth running of my field trials. Thanks to Stephan Smith for your assistance with semen collection for artificial insemination procedures, and my fellow post-graduates and colleagues who assisted in field work. Special thanks to Mr Ross Edwards and Mr Trevor Cook for their assistance with the AI and embryo transfer procedures. This project would not have been possible without your valuable expertise. I would also like to acknowledge Eric Thorstensen who did the analysis on the plasma samples. Debbie Hill I really appreciate your assistance with administrative work.

To numerous other colleagues who I had the opportunity to work with and learn new approaches or who just offered advice: Professor Tim Parkinson, Dr Rebecca Hickson, Dr Rene Corner-Thomas, Dr Anne Ridler and Dr Penny Back, thank you.

To the friends I made here in New Zealand: Claire Maxwell, Gaby Gronqvist, Antoinette Danso, Maria Loureiro, Lydia Cave, Heidi Jack, Amy Paten and Javier Roca. Thank you for sharing this journey with me. Many of you have also been in the pursuit of a PhD, and the support and understanding, the laughs, the tears, the hugs and the words of encouragement will not be forgotten.

The financial assistance provided through the funding of this project by grants from Gravida (National Centre for Growth and Development) and Massey University, Palmerston North, and PhD stipend from Gravida is acknowledged.

To my New Zealand "Trini" family: Marie Anne and Derek, thanks for the support and friendship. To my friends in Trinidad and the world over: Kathy, Shelly, Ria, Lisa- Marie, Jen, Heather, you ladies are amazing and I am grateful for your support and encouragement.

Finally, so much thanks and appreciation goes to my family, my parents: Margaret and Leroy, my brothers: Marc, Maurice and Martin, and my aunt Marilyn. Your constant and never-ending love and support in spite of the distance means more to me than you can ever

know. Thank you for believing in me!

# **Table of Contents**

Abstract	. iii
Acknowledgements	v
Table of Contents	/iii
List of tables	kiv
List of figures	/iii
List of Abbreviationsx	xii
1 Introduction	1
2 Embryonic development and maternal-embryonic interactions during ea pregnancy: A review of literature	-
2.1 Preamble	9
2.2 Embryonic development in sheep: day 0 to 34	11
2.2.1 Period of the ovum: day 0 to 10	12
2.2.2 Embryonic period: day 11 to 34	14
2.2.3 Extra-embryonic (fetal) membranes	21
2.3 Embryo- maternal interactions: Factors involved in maternal constraint	27
2.3.1 Dam nutrition	28
2.3.2 Dam Age and Parity	29
2.3.3 Dam size	30
2.3.4 Litter size	35
2.4 Uterine adaptions to pregnancy: Structural, secretory and biochemical (cell signallin function during embryo development	• •
2.4.1 Structural and secretory adaptations	37
2.4.2 Cell signalling during embryo development	40
2.5 Progesterone regulated embryo maternal interactions	46
2.6 Summary	51
Foreword to Chapter 3	55
3 Comparison of pre-implantation single and twin embryo size and embryonic ge expression at day 21 of gestation	
3.1 Abstract	59
3.2 Introduction	61

3.3 Materials and Methods	62
3.3.1 Experimental animals and design	63
3.3.2 Embryo morphometric measurements	66
3.3.3 Transcriptomic analysis of embryos	68
3.3.4 Quantitative real time PCR (qPCR) validation of transcriptomics res	ults73
3.4 Results	82
3.4.1 Pregnancy rates	82
3.4.2 Embryo morphometric data	85
3.4.3 RNA integrity	85
3.4.4 RNA-seq analysis	88
3.4.5 qPCR validation of RNA-seq data	88
3.4.6 Differentially expressed genes: qpCR analysis	89
3.5 Discussion	92
3.6 Summary and conclusions	96
Foreword to Chapters 4 and 5	99
4 Morphometric examination of pre-implantation embryos and mat	ernal hormone
	101
profiles in Cheviot and Suffolk breeds of sheep	
4.1 Abstract	
4.1 Abstract	
4.1 Abstract	103 104 106
<ul><li>4.1 Abstract</li><li>4.2 Introduction</li><li>4.3 Materials and Methods</li></ul>	
<ul> <li>4.1 Abstract</li> <li>4.2 Introduction</li> <li>4.3 Materials and Methods</li> <li>4.3.1 Experimental Animals and Design</li> </ul>	
<ul> <li>4.1 Abstract</li> <li>4.2 Introduction</li> <li>4.3 Materials and Methods</li></ul>	
<ul> <li>4.1 Abstract</li> <li>4.2 Introduction</li> <li>4.3 Materials and Methods</li> <li>4.3.1 Experimental Animals and Design</li> <li>4.3.2 Oestrus synchronisation and artificial insemination of recipients</li> <li>4.3.3 Embryo harvest (day 19 and 21)</li> </ul>	
<ul> <li>4.1 Abstract</li> <li>4.2 Introduction</li> <li>4.3 Materials and Methods</li> <li>4.3.1 Experimental Animals and Design</li> <li>4.3.2 Oestrus synchronisation and artificial insemination of recipients</li> <li>4.3.3 Embryo harvest (day 19 and 21)</li> <li>4.3.4 Embryo measurements</li> </ul>	
<ul> <li>4.1 Abstract</li> <li>4.2 Introduction</li> <li>4.3 Materials and Methods</li> <li>4.3.1 Experimental Animals and Design</li> <li>4.3.2 Oestrus synchronisation and artificial insemination of recipients</li> <li>4.3.3 Embryo harvest (day 19 and 21)</li> <li>4.3.4 Embryo measurements</li> <li>4.3.5 Blood sampling and hormonal assays</li> </ul>	
<ul> <li>4.1 Abstract</li> <li>4.2 Introduction</li> <li>4.3 Materials and Methods</li> <li>4.3.1 Experimental Animals and Design</li> <li>4.3.2 Oestrus synchronisation and artificial insemination of recipients</li> <li>4.3.3 Embryo harvest (day 19 and 21)</li> <li>4.3.4 Embryo measurements</li> <li>4.3.5 Blood sampling and hormonal assays</li> <li>4.3.6 Statistical analysis</li> </ul>	
<ul> <li>4.1 Abstract</li> <li>4.2 Introduction</li> <li>4.3 Materials and Methods</li> <li>4.3.1 Experimental Animals and Design</li> <li>4.3.2 Oestrus synchronisation and artificial insemination of recipients</li> <li>4.3.3 Embryo harvest (day 19 and 21)</li> <li>4.3.4 Embryo measurements</li> <li>4.3.5 Blood sampling and hormonal assays</li> <li>4.3.6 Statistical analysis</li> <li>4.4 Results</li> </ul>	103 104 106 106 107 107 107 108 109 110 111 111
<ul> <li>4.1 Abstract</li> <li>4.2 Introduction</li> <li>4.3 Materials and Methods</li> <li>4.3.1 Experimental Animals and Design</li> <li>4.3.2 Oestrus synchronisation and artificial insemination of recipients</li> <li>4.3.3 Embryo harvest (day 19 and 21)</li> <li>4.3.4 Embryo measurements</li> <li>4.3.5 Blood sampling and hormonal assays</li> <li>4.3.6 Statistical analysis</li> <li>4.4 Results</li> <li>4.4.1 Embryo morphometry</li> </ul>	103 104 106 106 107 107 107 108 109 110 111 111 111
<ul> <li>4.1 Abstract</li> <li>4.2 Introduction</li> <li>4.3 Materials and Methods</li> <li>4.3.1 Experimental Animals and Design</li> <li>4.3.2 Oestrus synchronisation and artificial insemination of recipients</li> <li>4.3.3 Embryo harvest (day 19 and 21)</li> <li>4.3.4 Embryo measurements</li> <li>4.3.5 Blood sampling and hormonal assays</li> <li>4.3.6 Statistical analysis</li> <li>4.4 Results</li> <li>4.4.1 Embryo morphometry</li> <li>4.4.2 Uterine and ovarian weights and morphometric data</li> </ul>	103 104 106 106 107 107 107 108 109 110 111 111 111 111 111

5 Assessment of Cheviot and Suffolk embryo size and somite count as in developmental stage and embryo growth	
5.1 Abstract	123
5.2 Introduction	124
5.3 Materials and Methods	125
5.3.1 Experimental animals and design	125
5.3.2 Day 19 and Day 20 embryo harvest	126
5.3.3 Analysis of embryo developmental stage	126
5.3.4 Statistical analysis of embryo developmental stages and somite count	127
5.4 Results	130
5.5 Discussion	132
5.6 Summary and conclusions	134
Foreword to Chapters 6 to 9	135
6 Effect of administration of exogenous progesterone on embryo size and plasma hormone concentrations in a dam size model of maternal constraint	
6.1 Abstract	139
6.2 Introduction	140
6.3 Materials and Methods	142
6.3.1 Experimental animals and design	142
6.3.2 Donor protocol: Oestrus synchronisation, superovulation, artificial ins day 0 and embryo recovery-day 6	
6.3.3 Recipient Protocol: Oestrus synchronisation, P4 treatment application transfer- day 6	
6.3.4 Embryo Harvest- day 19	146
6.3.5 Blood sampling and hormone assays	147
6.3.6 Statistical analysis	148
6.4 Results	150
6.4.1 Embryo dimensions	150
6.4.2 Uterine and corpus luteum weights and morphometric data	151
6.4.3 Ewe hormonal measurements	151
6.5 Discussion	157
6.6 Summary and conclusions	160

7 Comparison of uterine gene expression in a Cheviot-Suffolk n	
constraint in response to exogenous administration of progesterone	
7.1 Abstract	
7.2 Introduction	
7.3 Materials and Methods	
7.3.1 Experimental animals and design	169
7.3.2 Uterine tissue collection- day 19	170
7.3.3 RNA preparation	170
7.3.4 Designing of primers and probes	174
7.3.5 Quantitative PCR reactions	
7.3.6 Statistical analysis of RT-qPCR data	
7.4 Results	
7.5 Discussion	
7.6 Summary and conclusions	
8 Effect of timing of exogenous progesterone administration on maternal progesterone concentration and pregnancy rate in a dam size constraint.	model of maternal
8.1 Abstract	203
8.2 Introduction	205
8.3 Materials and Methods	206
8.3.1 Experimental animals and design	207
8.3.2 Donor protocol: Oestrus synchronisation, superovulation, arti day 0 and embryo recovery-day 6	
8.3.3 Recipient Protocol: Oestrus synchronisation, P4 treatment a transfer- day 6	••••••
8.3.4 Embryo Harvest- day 19	210
8.3.5 Blood sampling and hormone assays	212
8.3.6 Statistical analysis	212
8.4 Results	213
8.4.1 Embryo morphometric measurements	213
8.4.2 Uterine and corpus luteum weights and morphometric data	214
8.4.3 Ewe progesterone concentrations	215
8.5 Discussion	

8.6 Summary and conclusions	222
9 Effect of timing of exogenous progesterone administration during the period of pregnancy on day 6 and day 19 uterine gene expression in a dar	n size model of
maternal constraint	225
9.1 Abstract	227
9.2 Introduction	229
9.3 Materials and Methods	231
9.3.1 Experimental animals and design	231
9.3.2 Uterine tissue collection- day 6	233
9.3.3 Uterine tissue collection- day 19	233
9.3.4 RNA preparation	233
9.3.5 Designing of primers and probes	237
9.3.6 Quantitative PCR reactions	247
9.3.7 Statistical analysis of RT-qPCR data	248
9.4 Results	249
9.4.1 Differential gene expression of Day 6 uterine endometrium	249
9.4.2 Differential expression of Day 19 uterine horns	252
9.5 Discussion	258
9.6 Summary and conclusion	268
10 General Discussion	271
10.1 Overview of thesis	273
10.2 Summary of main findings and conclusions drawn	274
10.2.1 Embryo size and development in the peri-implantation period	274
10.2.2 Embryonic gene expression	275
10.2.3 Effects of progesterone on embryo growth	276
10.3 Methodological considerations	277
10.4 Recommendations for future research	280
10.5 Practical implications	
10.6 Overall summary and conclusions	284
References	
Appendices	
Appendix I	

Appendix II	
Appendix III	
Appendix IV	
Appendix V	
Appendix VI	324
Appendix VII	
Appendix VIII	
Appendix IX	336
Appendix X	
Appendix XI	

# List of tables

**Table 2.2** The effect of ewe and embryo genotype combination on offspring weight and size following reciprocal embryo transfer between large genotype Suffolk sheep and small genotype Cheviot sheep at day 19 and 90 of gestation and at birth. CinC (Cheviot in Cheviot =small control), CinS (Cheviot in Suffolk = luxurious environment), SinS (Suffolk in Suffolk = large control), SinC (Suffolk in Cheviot = restricted environment). Table shows least square means ( $\pm$  SEM). Within a row, means with differing superscripts are different from each other (p<0.05).

**Table 3.2** Candidate target and reference genes tested by qPCR. Gene ID, NCBI accessionnumber, forward, reverse primer and probe sequences, amplicon sizes (base pairs, bp) andprimer efficiency.78

**Table 3.3** Pregnancy rates (%) of embryo transfer groups and percentage of pregnant ewesthat satisfied the requirements of the experimental group that they were allocated to.......84

**Table 3.7 (A):** Embryonic mRNA gene expression levels in single embryos transferred to ewes with 1CL (previously singleton bearing) (1E1CL) experimental ET group compared with control group singleton bearing ewes (Con1E1CL), and single embryos transferred to ewes

**Table 3.7 (B):** Embryonic mRNA gene expression levels in twin embryos transferred to ewes with 2CLs (previously twin bearing) (2E2CL) experimental ET group compared with control group twin bearing ewe (Con2E2CL), and twin embryos transferred to ewe with 1CL (previously singleton bearing) (2E1CL), single embryos transferred to a ewe with 2CLs (1E2CL), and twin embryo transferred to ewe with 1CLs (2E1CL) experimental ET groups...91

**Table 4.2** Suffolk and Cheviot uterine weight, uterine body length and body width at day 19and 21 of gestation.112

 Table 5.1 Characteristics of embryos used to assign a developmental score.
 128

**Table 6.1** The effect of recipient ewe breed, and progesterone (P4) treatment combinationon embryo morphometry in sheep151

**Table 6.2** Plasma insulin and IGF1 concentrations (ng/ml) in Cheviot (C, n=20) and Suffolk (S,n=20) recipient ewes, in untreated (nP4, n=20) or progesterone supplemented (P4, n=20)recipient ewes, and on days 0, 3 and 6 (n=40).156

**Table 7.1** Candidate and reference gene ID, accession number, forward and reverse primersequences, amplicon sizes (base pairs, bp) and primer efficiency tested by reverse-transcription PCR (RT-qPCR).176

**Table 7.2** Pregnancy day 19 uterine horn mRNA expression levels in pregnant Cheviot ewes that were and were not administered exogenous progesterone from day 0-6 (CP4 and CnP4) and pregnant Suffolk ewes that were administered exogenous progesterone from day 0-6 (SP4) for combined ipsilateral and contralateral to CL horns (left) and for horn ipsilateral to the CL only (right). Fold change is expressed relative to levels in control Suffolk ewes that were not administered exogenous progesterone (SnP4, n=18; n=9 for combined and horn ipsilateral to the CL respectively). Data is normalised with *RPL19, SF1* and *TBP*. Data is shown as fold change with 95% confidence intervals (given in parenthesis). If confidence intervals do not include 1, then mRNA expression levels are significantly different from SnP4

 Table 8.1 The effect of time of progesterone (P4) treatment on day 19 embryo

 morphometry in sheep
 214

**Table 8.2** Day 0-6 plasma progesterone (P4) concentrations (ng/mL) in Cheviot ewes that did and did not receive exogenous P4 and Suffolk ewes that did not receive exogenous P4 via intravaginal CIDR for various time periods from day 0 to day 6......217

**Table 9.1** Candidate and reference gene ID, accession number, forward and reverse primersequences, amplicon sizes (base pairs, bp) and primer efficiency tested by reverse-transcription PCR (RT-qPCR).239

**Table 9.2** Pregnancy day 6 uterine horn mRNA expression levels in pregnant Cheviot ewes that were and were not administered exogenous progesterone for various time periods from day 0-6 (CP40-3, CP40-6 CP42-4, CP43-6 and CP4n) for horns contralateral to ovary containing the CL. Fold change is expressed relative to levels in control Suffolk ewes that were not administered exogenous progesterone (SP4n, n=7). Data is normalised with RPL19, SF1 and TBP. Data is shown as fold change with 95% confidence intervals (given in parenthesis). If confidence intervals do not include 1, then mRNA expression levels are significantly different from SnP4 control (bold). Different superscripts indicate that mRNA expression levels differ between CP40-3, CP40-6 CP42-4, CP43-6 and CP4n treatment groups (p<0.05).

**Table 9.3** Pregnancy day 19 uterine horn mRNA expression levels in in pregnant Cheviot (C) ewes that were and were not administered exogenous progesterone (P40 for various time periods from day 0-6 (CP4<sup>0-3,</sup> CP4<sup>0-6</sup> CP4<sup>2-4,</sup> CP4<sup>3-6</sup> and CnP4) for combined uterine horns. Fold change is expressed relative to levels in control Suffolk (S) ewes that were not administered exogenous progesterone (SnP4, n=14). Data is normalised with *RPL19, SF1* and *TBP*. Data is shown as fold change with 95% confidence intervals (given in parenthesis). If confidence intervals do not include 1, then mRNA expression levels are significantly

different from SnP4 control (bold). Different superscripts indicate that mRNA expression levels differ between  $CP4^{0-3}$ ,  $CP4^{0-6}$ ,  $CP4^{2-4}$ ,  $CP4^{3-6}$  and CnP4 treatment groups (p<0.05)......254

# **List of figures**

**Figure 2.2** Illustration of mammalian gastrulation. Cells separate from the central part of the ectoderm and move into the interior of the embryo, becoming endoderm and mesoderm. (Adapted from College of Arts and Science, The University of Tokyo, 2011)......16

 Figure 2.3
 Arrangement of mid-gestation extraembryonic membranes of the sheep

 (Latshaw, 1987).
 23

**Figure 2.6** Illustration of hormonal regulation and integrated signalling between embryouterine inter-face during early pregnancy. (Spencer *et al.*, 2006). In cyclic ewes circulating oestrogen increases expression of oestrogen receptor (*ESR1*) and oxytocin receptor (*OXTR*) present on the luminal epithelium (LE) and superficial glandular epithelium (sGE) during oestrus and metoestrus. At the same time circulating levels of progesterone are inadequate to activate progesterone receptors (*PGR*) to cause the suppression of *ESR1* and *OXTR*. Maturation of the corpus luteum (CL) during early dioestrus increases circulating progesterone, activating PGR with resulting suppression of ESR1 and OXTR for 8 to 10 days, in combination with low oestrogen. Continuous progesterone exposure results in downregulation of *PGR* in LE and sGE (days 11 and 12) ending the progesterone block of *ESR1* and *OXTR*. This is followed by increased ESR1 and subsequent induction of *OXTR* by oestrogen (day 13 and 14), allowing oxytocin secreted by the pituitary and CL to bind to *OXTR* resulting in luteolytic pulses of PGF<sub>2α</sub> via a prostaglandin synthase 2 (*PTGS2*) pathways. In pregnant sheep, interferon tau (INF- $\tau$ ), secreted by the elongating conceptus from day 11 to 25 of Figure 2.7 Mean progesterone concentration in the peripheral plasma of sheep during the oestrous cycle (Thorburn *et al.*, 1969)......47

Figure 2.8 Mean progesterone concentration in the peripheral plasma of sheep throughout pregnancy (Bassett *et al.*, 1969)......47

**Figure 4.2** Plasma progesterone concentrations in Suffolk and Cheviot ewes from day 0 to day 21 of pregnancy. Values are least squares means with standard error of the means...114

**Figure 4.4** Plasma insulin concentrations in Suffolk and Cheviot ewes from day 0 to day 21 of pregnancy. Values are least squares means with standard error of the means......115

**Figure 7.1** LB agar plate of *E.coli*-plasmid suspension. White colonies are the transformed competent cells with the inserted plasmid. Blue colonies are non-transformed cells........180

**Figure 8.1** Day 0-6 plasma progesterone (P4) concentrations (ng/mL) in Cheviot ewes that did and did not receive exogenous P4 for various time periods from day 0 to day 6 (CP40-3

# **List of Abbreviations**

ACTB = Beta actin

- AI = Artificial insemination
- BNC = Binucleate cells

Bp = Base pair

C = Cheviot

cDNA = Complementary Deoxyribonucleic acid

CIDR = Controlled intravaginal progesterone drug releasing device

CL/s = corpus luteum/corpora lutea

Con1E1CL = Control singleton bearing ewe (single CL), no embryo transfer

Con 2E2CL = Control twin bearing ewe (two CLs), no embryo transfer

COX2 = Cyclooxygenase 2

CP4 = Cheviot ewe that receive progesterone from day 0 to day 6 of pregnancy

CnP4 = Cheviot ewe that did not receive progesterone from day 0 to day 6 of pregnancy

Ct = Quantification cycle

CTSL = Cathepsin L

CV% = coefficient of variance

DEG = differentially expressed gene

DGAT2 = diacylglycerol-O-acyltransferase

DKK4 = Dickkopf WNT signalling pathway inhibitor 4

EL = embryo length: distance from the medial aspect of the head to the tip of the embryonic

tail, following the outer curvature of the embryo

ET = Embryo transfer

EGF = Epidermal growth factor

ER/ESR1 = Estrogen receptors

EW = Embryo width: distance between the two widest points of the embryo with the line passing just below the heart bulge

FDR = False discovery rate

FGF1 = Fibroblast growth factor 1

FGF2 = Fibroblast growth factor 2

FGF7 = Fibroblast growth factor 7

FGF10 = Fibroblast growth factor 10

GAPDH = Glyceralydehyde-3-phosphate dehydrogenase

GE = Glandular epithelium

GH = placental growth hormone

HB = Heart bulge width: distance between the two widest points of the heart bulge with the line passing through the midsection of the heart bulge

HGF = Hepatocyte growth factor

- HPRT = Hypoxanthine phosphoribosyltransferase 1
- IGFs = Insulin like growth factors
- IGF1 = Insulin like growth factor 1
- IGF2 = Insulin like growth factor 2
- IGF1R = Insulin like growth factor 1 receptor

INFτ = Interferon tau

- INFAR = Type 1 interferon receptors
- IRF2 = Interferon regulatory factor 2
- ISG17 = Interferon stimulated gene 17
- IUGR = Intrauterine growth restriction
- IV = Intravenous
- LAPTM5 = Lysosomal-associated protein transmembrane 5
- LE = Endometrial luminal epithelium
- LGALS3 = Lectin galactoside-binding, soluble 3
- LGALS15 = Endometrial galectin 15/ Lectin galactoside-binding soluble 15
- LOC101103603 = Pregnancy associated glycoprotein-4 like
- LOC101117738 = Pregnancy associated glycoprotein-1 like
- LRRC32 = Leucine rich repeat containing 32

#### MET = C-met proto-oncogene

- mRNA = Messenger RNA
- MSTN = Myostatin
- MUC1 = Mucin glycoprotein 1
- NFW = Nuclease free water
- OXTR = Oxytocin receptor
- PBS = Phosphate buffered saline
- PCR = Polymerase chain reaction
- $PGF_{2\alpha} = Prostaglandin F_{2\alpha}$
- PL = Placental lactogen
- PGR = Progesterone receptors
- PTGS2 = Prostaglandin endoperoxide synthase 2
- P4 = Progesterone
- qPCR = Quantitative real time PCR
- RIN = RNA Integrity number
- RNA = Ribonucleic acid
- RPL19 = Ribosomal protein L 19
- RSAD2 = Radical S-adenosyl methionine domain containing 2

RT = Reverse transcriptase

RT-qPCR = Reverse transcriptase quantitative PCR

S = Suffolk

SinCP4 = Suffolk embryo that was gestated in a Cheviot ewe that receive progesterone from day 0 to day 6 of pregnancy

 $SinCP4^{0-3}$  = Suffolk embryo that was gestated in a Cheviot ewe that receive progesterone from day 0 to day 3 of pregnancy

SinCP4<sup>0-6</sup> = Suffolk embryo that was gestated in a Cheviot ewe that receive progesterone from day 0 to day 6 of pregnancy

SinCP4<sup>2-4</sup> = Suffolk embryo that was gestated in a Cheviot ewe that receive progesterone from day 2 to day 4 of pregnancy

SinCP4<sup>3-6</sup> = Suffolk embryo that was gestated in a Cheviot ewe that receive progesterone from day 3 to day 6 of pregnancy

SinCnP4 = Suffolk embryo that was gestated in a Cheviot ewe that did not receive progesterone from day 0 to day 6 of pregnancy

SinSP4 = Suffolk embryo that was gestated in a Suffolk ewe that received progesterone from day 0 to day 6 of pregnancy

SinSnP4 = Suffolk embryo that was gestated in a Suffolk ewe that did not receive progesterone from day 0 to day 6 of pregnancy

SERPIN = Uterine serine proteinase inhibitor/ Uterine milk proteins

sGE = Superficial glandular epithelium

SPP1 = Secreted phosphoprotein 1/osteopontin

SP4 = Suffolk ewe that received progesterone from day 0 to day 6 of pregnancy

xxvi

SnP4 = Suffolk ewe that did not receive progesterone from day 0 to day 6 of pregnancy

TGF = Transforming growth factor

TKDP = Trophoblast Kunitz domain protein-1

TP1 = Trophoblast protein 1

UGKO = Uterine gland knock out

UTMP = Uterine Milk Proteins

YWHAZ = Tyrosine 3-monooxygenase/tryptophan 5-monooxygenase activation protein zeta.

1E1CL = Singleton embryos (harvested from a ewe with a single CL) transferred into a ewe that was also identified as having a single CL, and a single embryo that was removed

1E2CL = Singleton embryos (harvested from a ewe with a single CL) transferred into a ewe that was identified as having two CLs, and twin embryos that were removed

2E1CL = Twin embryos (harvested from a ewe with two CLs) transferred into a ewe that was identified as having a single CL, and single embryo that was removed

2E2CL = Twin embryos (harvested from a ewe with two CLs) transferred into a ewe that was

also identified as having two CLs, and twin embryos that were removed