Research Article

The Effect of Leptin on Luteal Angiogenic Factors in the Developing Porcine Corpus Luteum

Robin A. Katchko¹, Jessica R. Wiles¹, Martha A. Ramirez¹, Luis Ayala¹, Fang Xie¹, Chad W. O'Gorman¹, Duane H. Keisler², Randy L. Stanko^{1,3}, Michelle R. Garcia^{1*}, and Elizabeth A. Benavides⁴

¹Department of Animal, Rangeland & Wildlife Science, Texas A&M University-Kingsville, USA

²Division of Animal Sciences, University of Missouri, USA

³Animal Reproduction Laboratory, Texas A&M University, USA

⁴Department of Agriculture, Texas State University, USA

Abstract

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*Corresponding author

Michelle R Garcia, Department of Animal, Texas A&M University-Kingsville, Rangeland & Wildlife Sciences, Kleberg Ag. Bldg. Rm 133, Kingsville, TX 78363, USA, Tel: 1-361-593-3197; Fax: 1-361-593-3788; Email: michelle. aarcia@tamuk.edu

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Keywords

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Defects or abnormalities in a CL are believed to account for approximately 65% of recurrent miscarriages. Abnormalities can be associated with incomplete angiogenesis, which leads to a decrease in progesterone production and potential loss of a conceptus. Luteal angiogenesis is regulated by factors, such as fibroblast growth factor-2 (FGF2), vascular endothelial growth factor (VEGF), and angiopoietin-1 (Ang1). Leptin can influence the expression of VEGF, Ang1, and FGF2 in a caprine CL. Therefore, it is hypothesized that leptin is involved in luteal angiogenesis during CL development in the porcine species. Twenty-four mature crossbred gilts were allocated to one of four groups relative to day of the estrous cycle in preparation for CL collection: day (D) 4, 5, 6, or 7 of the estrous cycle (n = 6 gilts/D). Luteal tissue was divided and either frozen in liquid nitrogen for RNA analysis or enzymatically dispersed for cell culture. Gene expression for VEGF, Ang1, FGF2, leptin (tissue only), and leptin receptor (ObRb, tissue only) in CL tissue and cell cultures was determined by qPCR. Dispersed cells were cultured with or without leptin (0, 10^{-12} , 10^{-11} , 10^{-9} , 10^{-8} M; n = 3 wells/dose/gilt) for 24 hours. The expression of FGF2 in luteal tissue significantly (P<0.003) varied relative to D and Ang1 tended (P=0.1) to be highest in D 6 CL. Leptin increased (P<0.05) Ang1 in D4 cultured dispersed lutea. Collectively, leptin appears to be involved in the angiogenic process in the developing CL.

ABBREVIATIONS

VEGF: Vascular Endothelial Growth Factor; FGF2: Fibroblast Growth Factor; Ang1: Angiopoietin 1; D: Day; CL: Corpus Luteum; IGF-1: Insulin-Like Growth Factor 1

INTRODUCTION

Abnormal vasculature of the CL is a defect that leads to abnormal development, decreased progesterone production, and subsequent loss of a conceptus [1,2]. Identifying and understanding the underlying mechanisms of luteal angiogenesis may lead to a reduction in luteal-related infertility and alternate methodology for regulation of reproductive cycles. The luteal neo-vascularization process is attributed to the biological activity of fibroblast growth factor 2 (FGF2), angiopoietin 1 (Ang1), and vascular endothelial growth factor (VEGF). Both VEGF and FGF2 promote capillary membrane destabilization, endothelial cell differentiation, proliferation, migration, and vascular tube formation in human, bovine, and ovine luteal tissue [3,4]. Maturation and stabilization of nascent vessels is then promoted by Ang1 through the recruitment of stromal support cells, including pericytes and smooth muscle cells [5]. Each of these angiogenic regulatory factors is regulated by leptin in both nonovarian and ovarian tissues [6-8].

Both leptin and its receptor, ObRb, have been identified in normal and polycystic ovaries of various species [9-12], particularly in steroidogenically active follicles and luteal tissue where it has been reported to influence the synthesis and secretion of progesterone [8,13]. However, in order to substantially modify steroid production cells are usually incubated with leptin in the presence of an additional growth promotant, such as IGF-I [13-17], which suggests that a potential ancillary role for leptin in ovarian tissue, such as angiogenesis, may exist. It has recently been reported that leptin is present in luteal vascular endothelial cells and in small and large luteal cells of an early stage CL [8]. Furthermore, leptin upregulates the expression of angiogenic factors in the early caprine CL [8]. Collectively, the evidence supports the supposition that leptin may be involved in the development of a CL, which may include proper vascularization of the tissue. Understanding the mechanisms of the development of luteal tissue may lead to a future reduction in the occurrence of luteal-associated infertility. Therefore, it is hypothesized that leptin is involved in luteal angiogenesis during CL development in the porcine species. The study herein characterizes the primary angiogenic growth promotants VEGF, Ang1, FGF2, leptin, ObRb in developing lutea and investigates the effect of leptin on the production of the angiogenic factors throughout early luteal development.

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MATERIALS AND METHODS

Animals and management

Twenty-four, mature crossbred (Yorkshire x Landrace) gilts, Sus scrofa, of similar age and weight, from the Texas A&M University-Kingsville (TAMUK) Farm were utilized. Gilts were randomly allocated to one of four groups relative to day of the estrous cycle in preparation for CL collection: day (D) 4, 5, 6, or 7 of the estrous cycle (n = 6 gilts/D). Ovulation in the porcine species occurs between 10 and 85 hours after the onset of estrus [18], therefore, CL collection began on D4 of the estrous cycle to allow for completion of the ovulation process. Animals were housed in concrete pens (7 m x 13 m) in an outdoor, sheltered facility and had ad libitum access to water and a corn-soy based diet [19]. Females were checked twice daily (30 minutes) for classical, behavioral estrus using an intact boar. Tissue collection occurred on the third cycle after animals exhibited two consistent consecutive cycles. On the day of CL tissue collection, blood samples were collected, via jugular venipuncture, for analysis of serum progesterone for confirmation of normal luteal function.

Corpus luteum collection

Feed was removed from the gilts for a period of 24 hours prior to tissue collection. On the day of CL tissue collection, body weight was recorded and the gilts were transported to a surgical facility located on the TAMUK farm. Gilts were pre-anesthetized with sodium thiopental (2.5%) administered intravenously (ear vein) using a butterfly infusion set (25G x 1.65 cm needle, 30.48 cm tubing; Abbott Laboratories, Chicago, IL). For a surgical plane of anesthetization, the inhalation agent, isoflurane, was utilized. Gilts received isoflurane gas from an isoflurane vaporizer that was set to 3-4%. Corpora luteal were harvested from the ovaries and placed in ice-cold Hank's Balanced Salt Solution (1X) (HyQ HBSS; HyClone, Logan, UT) for transportation. Harvested tissue was divided and processed for RNA extraction, or enzymatically digested and dispersed for cell culture.

Serum analyses

Serum and culture media were analyzed for progesterone concentration by radioimmunoassay (Coat-A-Count®; Diagnostics Product Corporation, Los Angeles, CA) [20,21] to confirm a functional CL at the time of tissue harvest, cell viability during culture process, and response to culture treatment. Serum concentrations of leptin were determined using ovine specific leptin RIA validated for porcine serum [19,22]. Inter- and intraassay coefficients of variation for progesterone and leptin assays were < 10% as previously reported [19,23].

Dispersed CL cell cultures

Luteal tissue (n=6 animals/per day assignment) was processed as previously reported [8]. Briefly, tissue was mechanically minced and enzymatically dissociated in digestion media (Type IA collagenase, 1.27 mg/ml, 0.5% BSA in Dulbecco's Modified Eagle's Medium; DMEM) for 90 minutes at 37°C with gentle agitation in a shaking water bath. Cells were filtered through a metal mesh (80 μ m) and centrifuged twice at 150 X *g* for 4 minutes and the enzymatic supernatant was removed. Cells were re-suspended and the numbers of viable cells were determined using the Trypan Blue Exclusion Test. Cell viability was greater than 85% for each cell culture. The dispersed cells were seeded in a treated polystyrene 24-well cluster dishes (Sigma) at 3 X 10⁶ cells/well in attachment serum media [1 ml/ well; DMEM supplemented with 10% (v/v) charcoal stripped fetal calf serum and 5,000 µg/mL penicillin/streptomycin] and incubated for 24 hours in a humidified atmosphere of 5% CO₂ in air at 37°C to equilibrate cells to culture condition, allow aggregation, and promote adherence. Following the 24 hour incubation, attachment serum media was aspirated and replaced with culture media (1 ml/well; DMEM, 0.1% (w/v) BSA, 2.0% (v/v) charcoal stripped fetal calf serum, 0.5 mM ascorbic acid, and 5,000 $\mu g/mL$ penicillin/streptomycin) containing recombinant ovine leptin (0, 10⁻¹², 10⁻¹¹, 10⁻¹⁰, 10⁻⁹, 10⁻⁸ M; Cell Sciences) in triplicate (3 wells/dose/gilt) for 24 hours in a humidified atmospere of 5% $\mathrm{CO}_{\scriptscriptstyle 2}$ in air at 37°C. Doses of leptin were based on previous studies examining the in vitro effect of leptin in metabolically active cells [8,24] and conditions where leptin stimulates the MAPK signal transduction cascade [25]. Biologically active ovine leptin protein, which is >92% identical porcine leptin (UniProt access. #Q29406 and # G8BLB2), was substituted due to availability. After 24 hours, the cell culture media was aspirated and stored at -80°C for measurement of secreted progesterone as a marker for cell responsiveness to leptin in vitro [8,26,27]. Denaturing solution [1 ml/well; 25 mM sodium citrate (pH 7.0), 0.5% (w/v) N-laurylsarcosine, 4 M guanidine thiocyanate with 0.7% (v/v) 2-mercaptoethanol] was added to cells, aspirated, pooled by dose, and stored at -80°C for total RNA extraction and mRNA analysis of VEGF, Ang1, and FGF2.

RNA extraction and qPCR

Total RNA was extracted from CL tissue (~20 mg) and denatured cells using a phenol: chloroform based procedure [8,19,28]. Total RNA was quantified using a spectrophotometer (640 UV/Visible Beckmann) at UV absorbencies (A) of 260 to 280. Purity of the extracted RNA was evaluated by determining the A260/A280 ratio. The extracted total RNA was then analyzed for VEGF, Ang1, FGF2, leptin (tissue only), and ObRb (tissue only) mRNA using quantitative polymerase chain reaction (qPCR) with the DNA Engine Opticon II (MJ Research, Inc., Miami, FL). Two µg of extracted total RNA was DNase (RQ1 DNase) (Promega, Madison, WI) treated and reverse transcribed using moloney murine leukemia virus reverse transcriptase (M-MLV RT; Promega). Briefly, total RNA was incubated at 37°C for 30 minutes for DNase 1 treatment, followed by 75°C incubation for 8 minutes to terminate DNase 1 activity. Samples were placed on ice and oligo dt15 (Promega) and M-MLV RT (Promega) were added. Samples were incubated at 37°C for 1 hour to RT mRNA into cDNA. Samples were then incubated at 85°C for 5 minutes to terminate the RT.

Four μ l of cDNA were amplified by qPCR using 25 pmol of VEGF, Ang1, FGF2, leptin, ObRb, and cyclophilin primers previously published [8], all synthesized by Integrated DNA Technologies Inc. (Coralville, IA), using a SYBR® green kit (Takara Mirus Bio®, Madison, WI) and diethyl pyrocarbonate (Sigma) treated water in a 50 μ l reaction. The PCR cycles were as follows: a 10 minute hotstart at 94°C, then 94°C for 1 minute, anneal temperature of 50°C for 30 seconds, 72°C for 1 minute for 35 cycles followed by

a 10 minute extension at 72°C that was conducted at the end of the PCR. To control for potential primer-dimer formation, control tubes containing everything except total RNA were subjected to the same PCR conditions. Relative quantities of VEGF, Ang1, FGF2, leptin, and ObRb, and cyclophilin (housekeeping gene) were quantitated from a relative standard curve (1 µg to 10 pg of total RNA) obtained from qPCR of cyclophilin mRNA in porcine liver tissue [8,19,23,29]. Each qPCR was performed in duplicate to reduce experimental errors. Cycle threshold (C,), the cycle at which the amplification curve crosses threshold, was set to standardize each PCR run. Relative values obtained from the relative standard curve were transformed to $\log_{\rm 10}$ and normalized with cyclophilin. Cyclophilin mRNA was used to account for variability in sample preparation and procedural variability to normalize targeted amplicons. Dispersed luteal cell values are represented as a % of the control (0M) as previously reported [8]. PCR products were then separated on a 2% agarose gel by electrophoresis and visualized with ethidium bromide staining for estimated size verification.

Statistical analyses

qPCR values of VEGF, Ang1, FGF2, leptin, and ObRb gene expression in luteal tissue were transformed to \log_{10} and analyzed using the PROC MIXED procedure of SAS with day of development as the source of variation. qPCR values of VEGF, Ang1, and FGF2 obtained in cultured cells were transformed to log₁₀ and expressed as a percentage of 0M treatment dose. The effect of leptin on each of the angiogenic factors in vitro was analyzed using the PROC MIXED procedure of SAS. Leptin dose was the source of variation. Animal was used as the subject to account for correlated variation within all MIXED analysis procedure. The PDIFF option of least square mean (LS Mean) values was conducted to compare means where significance of the model was detected to determine the specific day (tissue and serum analyses) or doses (cell culture) that were significantly different than 0M. The values are represented as LS Means ± standard error of the mean.

RESULTS

Neither serum concentrations of progesterone or leptin (Figure 1) significantly differed relative to day as the CL developed over the 4 day sampling period. However, serum progesterone values are indicative of a normal, functional CL at the time of tissue collection.

Vascular endothelial growth factor, ObRb, and leptin gene expression did not significantly differ by day of developing luteal tissue (Figure 2). However, FGF2 varied significantly (P<0.0003) by day, decreasing between D4 and D5, increasing between D5 and D6, and decreasing on D7 to concentrations detected in D4 tissue. Angiopoietin 1 tended (P=0.1) to differ relative to day increasing on D6 and, similar to FGF2, decreasing on D7 to concentrations detected in D4 tissue.

Leptin treatment (Figure 3) stimulated (P=0.05) Ang1 gene expression in D4 dispersed cell cultures at the -11M dose compared to all other doses, but did not upregulate VEGF or FGF2 at any dose. Leptin did not significantly influence gene expression of VEGF, Ang1, or FGF2 in the D5 or D6 cell cultures; however, leptin did significantly reduce (P<0.03) FGF2 in D7 cell cultures

across all doses compared to 0M. Leptin influenced ($P \le 0.05$) progesterone secretion in D4, 5, 6, and 7 cell cultures (Figure 4) verifying cellular responsiveness to leptin as previously reported [8,27].

DISCUSSION

Gene expression of VEGF, leptin, and ObRb did not differ by day in luteal tissue, which is not attributed to abnormal CL function since circulating concentrations of progesterone were normal for each day of the porcine estrous cycle [23] and serum leptin remained constant. Insignificant changes during the 4 days of early development does not necessarily imply that each factor would not have differed prior to or after the collection period as previously reported in the caprine [8], porcine [10] and human [30] CL. Vascular endothelial growth factor is reported to increase and remain elevated throughout the luteal lifespan to support angiogenic processes [3,4] and maintain vascular viability [31]. The expression of leptin and ObRb has also been reported to vary throughout the luteal phase in the caprine [8] and porcine [10] CL. However, leptin only appears to influence angiogenic factor expression in the early stage caprine CL [8], which implies leptin may potentially have an alternate role in the mature CL. Both Ang1 and FGF2 exhibited a similar pattern of expression in CL tissue where D7 expression appeared to be lower than in D6 CL.







extracted and relatively quantitated using qPCR. Relative quantities of PCR amplicons were transformed to \log_{10} and normalized with cyclophilin. No differences in VEGF, leptin, or ObRb gene expression was detected between D of the developing CL. FGF2 varied significantly (P<0.003) as the CL developed and Ang1 tended (P=0.1) to vary by D decreasing between D6 and 7. ^{abc}Different superscript represent LS means that significantly differ ($P\leq0.05$) within target gene by D.

Angiopoietin 1 is important in the stabilization of nascent vessels [5], therefore, a tendency towards a decrease in Ang1 availability in D7 tissue may imply a vascular stabilization event in D6 tissue. The variation in FGF2 during the 4-day window of CL development suggests that as development occurs the requirement for FGF2 may alter in availability. Luteal FGF2 is believed to be a primary regulator of luteal angiogenesis during CL development [4,32,33]. However, FGF2 can mitigate cellular apoptosis in the ovary [34-36], which may support the supposition that FGF2 is important for luteal angiogenic processes as well as the maintenance of forming luteal tissue. Therefore, the required availability of FGF2 may vary daily depending on the functional requirements of the hormone. Interestingly, leptin reduced FGF2 in D7 dispersed cell cultures rather than increase it, as observed for Ang1 in D4 cultures. A previous report in the caprine CL indicated that leptin stimulated VEGF, Ang1, and FGF2 in the early stage CL [8]. However, the development of the caprine CL is a slower process

study may have been too late for detecting the stimulatory affect of leptin on all angiogenic factors in the porcine CL. The reduction of FGF2 in D7 cultures may explicate, in part, the decrease in FGF2 in D7 CL tissue, but this does not explicate the inhibitory rather than a stimulatory affect in vitro as previously reported [8]. This inhibitory and stimulatory variation was also observed in secreted progesterone in cell culture media where leptin increased progesterone secretion in dispersed cell cultures in the study herein, which is in contrast to previous reports. Leptin decreases progesterone secretion in luteinized bovine granulosa cells [14] and in dispersed caprine lutea [8], however, leptin can both increase or decrease progesterone secretion in cultured porcine luteinized granulosa cells [27]. The direction and magnitude through which leptin influences progesterone secretion can be altered by dose of leptin and/or the addition of growth promoting hormones such as IGF-I [18,27]. Collectively,

than in the porcine species [37]. Therefore, D4 in the current



Figure 3 The gene expression of VEGF, Ang1, and FGF2 gene in cultured dispersed lutea from mature gilts on days (D) 4, 5, 6, and 7 of the estrous cycle (n=6 gilts/D). Cells were incubated with or without leptin (0, 10^{-12} , 10^{-11} , 10^{-0} , 10^{-9} M) for 24 hours. Leptin increased (P=0.05) Ang1 in the -11M dose compared to all other doses in the D4 cells and significantly decreased (P<0.03) FGF2 in D7 cultures across all doses compared to 0M. ^{abc}Different superscript represent LS means that significantly differ (P<0.05) from control dose (0M).

leptin influences progesterone secretion in luteinized cells, which may be influenced by various factors as the CL experiences the daily biochemical and biophysical modifications that occur as a CL develops [38]. Therefore, variations in endogenous parameters that influence the affect of leptin on progesterone secretion may also influence the affect of leptin on luteal angiogenic factor expression.

CONCLUSION

Leptin influences the expression of angiogenic factors during early luteal development supporting a potential role for





leptin in the luteal angiogenic process of the porcine CL. Further investigation into the functional role of leptin in early luteal development is needed to definitively determine how leptin influences angiogenic processes in the developing CL.

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