



Interferon-stimulated Gene 15 and Interferon-1 Stimulated Gene 17 Messenger RNA-based Detection of Early Pregnancy in Aardi Goats in Saudi Arabia

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ABSTRACT

The accuracy of interferon-stimulated gene 15 (*ISG15*) and interferon-stimulated gene 17 (*ISG17*) mRNA levels in early detection of pregnancy in Aardi goats compared to progesterone and ultrasound (US) were evaluated. Female goats were synchronized using the ovsynch protocol level in combination with natural mating (NM). Blood samples were collected at 1, 7, 15, 23, 35, and 60 days post NM. Levels of ISG15 and ISG17 mRNAs were assayed using real-time PCR, and serum progesterone (P4) concentrations were assayed using an ELISA kit. Pregnancy detection was performed by US on 23, 35, and 60 days post NM. Serum P4 concentration was significantly higher in pregnant than non-pregnant goats at 15, 23, 35, and 60 days post NM. Relative expression of mRNA of ISG15 and ISG17 was significantly higher in pregnant goats at 7, 15, and 35 days post NM. ISG15 and ISG17 were not significantly different compared to P4 and US in the pregnant and non-pregnant goats. The accuracy of ISG15 and ISG17 was not significantly different than P4 on days 7 and 15 or from P4 and US on day 23 post NM. Decreased accuracy of ISG15 and ISG17 on day 35 might be done to lower levels. US provided accurate pregnancy diagnoses on day 35 (96.97%) and 60 (100%) post NM. ISG15 and ISG17 mRNA levels could be considered good indicators of goat pregnancy on day 23 post NM. These methods will provide early and precise detection of pregnancy compared to the routinely used serum P4 and US methods.

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Authors' Contribution

KS and MH designed and performed the experiment. HM and MA analysed the data and helped in reviewing the manuscript. SG and HM finalized the writing.

Key words

Aardi Goats, ISG15, ISG17, P4, Early pregnancy detection

INTRODUCTION

Several methods have been used for detection of early pregnancy in goats over the past few decades. The simplest method involves observation of signs of estrus (Goel and Agrawal, 1992; Ishwar, 1995). Also abdominal auscultation is relatively inexpensive and easy to perform in field conditions with more than 90% accuracy 60 days post gestation (Memon and Ott, 1980). Measuring progesterone (P4) is another method of indirect pregnancy detection in goats. P4 concentrations in serum or milk

samples can be used to determine pregnancy at 19–23 days post insemination (Al-Samawi *et al.*, 2015). Real-time ultrasound scanning (Davey, 1986; Haibel, 1990) had an accuracy of 66% on days 17 to 19 and the accuracy reached 100% on day 34 of pregnancy (Singh *et al.*, 2004). Pregnancy-associated glycoprotein (PAG) is useful for accurate prediction of pregnancy post day 30 (Szenci *et al.*, 1998). The current method for detecting early pregnancy factor (EPF) uses the rosette inhibition test (RIT) which, while accurate, can be difficult to maintain and is not suitable EPF as a diagnostic tool for early pregnancy has been problematic for high-throughput diagnostic applications for example in cattle and dairy bovine (Cordoba *et al.*, 2001; Gandy *et al.*, 2001). Interferon-tau is a major paracrine signal produced by the bovine and ovine conceptus and acts on the endometrium

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to stimulate secondary responses that are necessary to maintain pregnancy (Roberts *et al.*, 1992; Thatcher *et al.*, 1995; Spencer and Bazer, 2002). Conceptus-derived interferon-tau disrupts the signal transduction pathway that regulates release of prostaglandin F_{2α} (PGF_{2α}), which is the major luteolytic product in sheep, cattle, and goats (Bazer, 1992; Thatcher *et al.*, 1995, 2001). Interferon-tau possesses anti-luteolytic activity and is detected in goat trophoblastic cells from 14 until 17 d of pregnancy (Guillomot *et al.*, 1998). Two proteins have been detected in goats, a non-glycosylated (17 kDa) and a glycosylated (22-24kDa) isoform. Proteins similar to interferon-tau have been identified in other species, including humans, which share 73% identity with ovine interferon-tau (Whaley *et al.*, 1994). Greater expression of ISGs in the endometrium has been recently reported in pregnant cows on day 7 (Sponchiado *et al.*, 2017). Interferon-tau also induces synthesis and secretion of an ubiquitin homolog (ubiquitin cross-reactive protein) that is called ISG15 (as described by HUGO gene nomenclature). Bovine ISG15 is released by the endometrium at times coincident with Interferon-tau release from the conceptus (Austin *et al.*, 1996; Hansen *et al.*, 1997, 1999; Johnson *et al.*, 1998). Likewise, ISG15 is found in significant amount in uterine flushing from day 18 pregnant cows. Because ISG15 functions as an intracellular ubiquitin homolog (Loeb and Haas, 1992; Cunha *et al.*, 1996; Johnson *et al.*, 1998) and an extracellular cytokine (Recht *et al.*, 1991; Cunha *et al.*, 1996), detection of low blood ISG15 mRNA levels during serial collection from days 17 to 25 serves as an accurate indicator of non-pregnant cows, allowing re-synchronization and insemination (Hyungchul *et al.*, 2006). In addition, the sustained presence of conjugated ISG15 through day 50 of pregnancy might reflect stabilization of conjugated proteins in response to implantation and the development of the placenta (Kathy *et al.*, 2004). Johnson *et al.* (2002) have found that ISG17 mRNA increased in the stratum compactum between days 11 and 13, and expression extended into the deep glandular epithelium and stratum spongiosum on day 15 through 17 in pregnant ewes. The present study was conducted to investigate the possibility of using ISG15 and ISG17 for early pregnancy diagnosis in Aardi goats.

MATERIALS AND METHODS

Flock description

The study was conducted for 70 days at the Research Station, Department of Animal Production, King Saud University, Riyadh, Saudi Arabia. Thirty-two healthy mature Aardi female goats (1–1.5-years old) were used in the study and six males were used for natural mating

(NM). Semen was collected and examined before males were introduced to females for NM. Animals were fed a commercial total mixed ration (ME 1950 kcal kg⁻¹1108, crude protein 13%, crude fat 2%, crude fiber 10%, and ash 8% on DM basis; Al-wafi pellets, ARASCO, Riyadh, King Saudi Arabia). Feed was offered twice daily at 07:00 and 15:00 h, and animals had free access to clean fresh tap water throughout the study. Female goats were synchronized using the ovsynch protocol in combination with NM (Al-Samawi *et al.*, 2015). Female goats detected in estrus intramuscular (i/m) injection of 8 µg of GnRH (GnRH Receptal®, MSD Animal Health, Upper Hutt, Wellington, New Zealand) on day 0. On day 7, goats received an i/m injection of 10 mg of prostaglandin F_{2α} (PGF Lulatoryse, Pfizer Animal Health) and 48 h following the PGF_{2α} injection, they received a second i/m injection of 8 µg of GnRH. On the same day of the 2nd GnRH treatment, hot female goats were watched and isolated. Bucks were introduced to hot female goats. Female goats were watched and mating data recorded. Day 1 of pregnancy was defined as 48 h from the estrous beginning.

Blood samples and P4 assay

Blood samples were collected via jugular vein at 1, 7, 15, 23, 35, and 60 days post NM into vacutainer tubes. Serum was separated by centrifugation at 860xg for 30 min at 4°C, transferred into 1.5 mL Eppendorf tubes, and stored at -20°C until assay for P4. Three milliliters of whole blood were collected into Tempus tubes that contained 6 mL of stabilizing reagent for gene expression studies (Applied Biosystems, Foster City, CA, USA), which immediately lysed the blood cells. Tempus tubes with whole blood were vortexed for 10 s and stored at -85°C until isolation of RNA. P4 serum concentrations were assayed after one week from samples collected. P4 serum concentrations were assayed using commercial ELISA kits (Human, Wiesbaden, Germany) and micro-titrimetric plates. The assay procedures were performed according to the manufacturer's instructions, and an automatic photometer plate reader was used for abs gene expression studies orbencies readings.

RNA extraction

Tempus tube contents were transferred to 50-mL falcon tubes. One molar phosphate buffer saline (PBS) (Ca²⁺/Mg²⁺-free) was added into the tube to bring the total volume to 12 mL. The diluted sample was vortexed for at least 30 s then centrifuged at 4°C at 860xg for 60 min. The supernatant was carefully poured off to avoid RNA pellet loss. The tube was left inverted on absorbent paper for 1 to 2 min. An RNA isolation kit (Promega®, Wisconsin, USA) was used for RNA extraction, and 175 µL RNA lysis buffer

was immediately added to samples and mixed thoroughly by inversion. Next, 350 μ L RNA dilution buffer was added and mixed by inverting 3–4 times and then heating at 70°C for 3 min. The solution was then centrifuged at room temperature at 12,000 \times g for 10 min; the clear lysate was transferred into a fresh tube and 200 μ L 95% ethanol was added and mixed well. The mixture was transferred to a spin basket assembly and centrifuged at room temperature at 12,000 \times g for 1 min; the eluate was discarded and 600 μ L of RNA wash solution was added. Again, the tubes were centrifuged at room temperature at 12,000 \times g for 1 min, the eluate discarded, and 50 μ L of DNase mix was added to the membrane and incubated at room temperature for 15 min. Then, 200 μ L DNase stop solution was added and centrifuged at room temperature at 12,000 \times g for 1 min. RNA wash solution (600 μ L) was added and centrifuged at room temperature at 12,000 \times g for 1 min and discarded by pouring off. Another 250 μ L of RNA wash solution was added, centrifuged at room temperature at 12,000 \times g for 2 min and transferred from the spin basket to the elution tube. Nuclease-free water (100 μ L) was added to the membrane, centrifuged at room temperature at 12,000 \times g for 1 min to elute the RNA. RNA quality was assessed using agarose gel electrophoresis; two bands appeared as shown in [Supplementary Figure 1](#). RNA samples were stored at -85°C. Total RNA concentration was estimated using Nanodrop 2000 (Thermo Fisher Scientific, Inc. Waltham, MA, USA). Single-stranded cDNA was synthesized from total cellular RNA (equal volumes) using the Reverse Transcription System (Promega®, Wisconsin, USA). Total RNA samples were placed in micro-centrifuge tubes and incubated at 70°C for 10 min, spin at 860 \times g, and placed on ice. Master mix was prepared and RNA samples were added to the master mix for a final volume of 20 μ L, and then incubated at 42°C for 30 min. The samples were heated at 95°C for 5 min, and incubated at 0–5°C for 5 min. The first-strand cDNA synthesis reaction was diluted to 100 μ L with Nuclease-free water. The polymerase chain reaction (PCR) cycling program was set for 35 cycles as follows: initial denaturing at 94°C for 5 min, followed by a denaturation at 94°C for 30 s, then annealing at 72°C for 1 min. A final extension step was performed at 72°C for 20 min. The PCR product was resolved as a single band using agarose gel electrophoresis using Mini-Sub Cell Gt electrophoresis Cell (Bio-rad, Hercules, California, USA) ([Supplementary Fig. 2](#)).

Quantitative real-time PCR

Diluted cDNA (10 μ L) was used as a template for quantitative real-time PCR amplification using SYBR Green (Applied Biosystems, Foster City, CA, USA). The GAPDH gene was used as an internal reference for

normalization (NCBI Accession no.: BC102589) for ISG15 mRNA (NCBI Accession no.: NM_174366) and ISG17 (NCBI Accession no.: NM_001009735) expression. Bovine ISG15 (forward; 5'GGTATCCGAGCTGAAGCAGTT3', reverse; 5'ACCTCCCTGCTGTCA AGGT3'), ISG17 (forward; 5'GGTATCTGAGCTGAAGCAGTT3', reverse; 5'ACTTCCCTGCTGTCAAGGT3'), and GAPDH (forward; 5'GATTGTCAG CAATGCCTCCT3') reverse; (5'GGTCATAAGTCCCTCCACGA3') primers were designed to generate an amplicon size of 87 and 94 bp, respectively. ISG15 and ISG17 mRNA expression level was reported relative to GAPDH. Prediction of pregnancy based on blood ISG15 and ISG17 mRNA levels determined using -0.62 and -0.58, respectively as the arbitrary threshold levels relative to GAPDH expression. Goats with ISG15 mRNA levels over -0.62 (relative to GAPDH) were considered pregnant, whereas those with mRNA levels under -0.62 were considered non-pregnant. Goats with ISG17 mRNA levels over -0.58 (relative to GAPDH) were considered pregnant, whereas those with mRNA levels under -0.58 were considered non-pregnant. This level of ISG15 and ISG17 mRNA was sustained 1, 7, 15, 23, 35, and 60 days post natural mating (NM). RT-PCR for ISG15 and GAPDH cDNA amplification was performed using 40 cycles of 95°C for 30 s, 62°C for 30 s, and 72°C for 15 s (Applied Biosystems, Foster City, CA, USA). Following RT-PCR, cDNAs were melted (melting curve) to ensure the quality of amplification. For melting curve analysis, RT-PCR products were incubated for 10 s at each step with an increase in temperature of 0.5°C from 55 to 95°C in each cycle.

Pregnancy detection using ultrasound

Pregnancy was diagnosed by ultrasonography (US) (Prosound 2, ALOKA, Tokyo, Japan) on day 23 post NM using a multi-frequency linear trans-rectal probe (UST 660-7.5, ALOKA, Tokyo, Japan) and confirmed on day 35 and 60 using a multi frequency convex abdominal probe (UST-9137C, ALOKA, Tokyo, Japan). Number and sex of embryos were recorded. All pregnancies were tracked until kidding and the number and sex of kids were confirmed.

Statistical analysis

Predictions of pregnancy status using RT-PCR of ISG15 and ISG17 were compared with P4 concentration in serum, and the transrectal and transabdominal US data. The results were presented as (i) true pregnant (TP): the goat was diagnosed as pregnant and was already pregnant. (ii) False pregnant (FP): the goat was diagnosed as pregnant while it was not pregnant. (iii) True non-pregnant (TN): the goat was diagnosed as non-pregnant and was already non-pregnant. (iv) False non-pregnant (FN): the

goat was diagnosed as non-pregnant while it was pregnant. Probability that the diagnosis of goats as non-pregnant, which is truly non-pregnant.

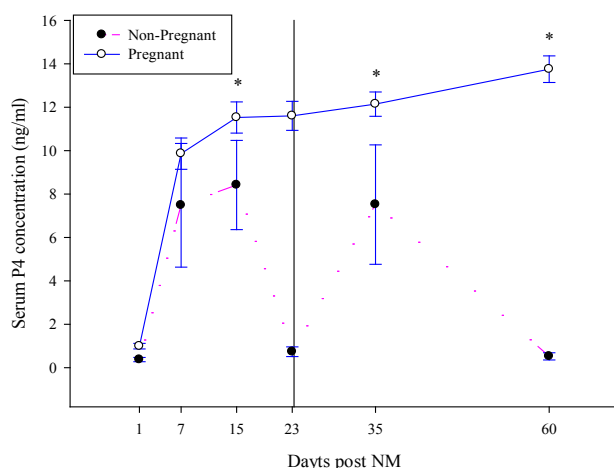


Fig. 1. Concentrations of progesterone hormone (P4) in pregnant and non-pregnant Aardi goats at 1, 7, 15, 23, 35, and 60 days post natural mating (NM), blood samples were collected from Aardi goats, (means \pm SEM), *denotes significant difference ($P < 0.01$) between pregnant and non-pregnant goats, ($n = 32$).

Sensitivity was defined as the probability that the goat diagnosed pregnant was truly pregnant, where Sensitivity = $TP / (TP + FN) \times 100$. Specificity was defined as the probability that the goat diagnosed non-pregnant was truly non-pregnant, where Specificity = $TN / (TN + FP) \times 100$. The positive predictive value (PPV): (probability that the goats are truly pregnant if the diagnosis is pregnant) was calculated as $PPV = (TP / [TP + FP]) \times 100$. The negative predictive value (NPV): (probability that the goats are truly non-pregnant if the diagnosis is non-pregnant) was calculated as $NPV = (TN / [TN + FN]) \times 100$ (Hyungchul *et al.*, 2006). Accuracy of the diagnosis test was calculated as Accuracy = $([TP + TN] / [TP + TN + FN + FP]) \times 100$ (Karen *et al.*, 2006). A general linear model (GLM) in SAS® (SAS, 2003, Cary, NC, USA) was used to conduct an analysis of variance (ANOVA). A completely randomized design (CRD) was used to examine effects of pregnant and non-pregnant on ISG15, ISG17, and P4 concentrations and to examine effects of day of pregnancy on ISG15, ISG17, P4, and US. Student's *t* test was used to examine the sensitivity, specificity, accuracy, PPV, and NPV among the early pregnancy diagnosis methods. Data were expressed as mean \pm standard error of the mean (SEM) for all parameters at a level of $P < 0.05$.

RESULTS

The results showed no significant difference in serum P4 concentrations in pregnant (0.38 ± 0.10 , 0.99 ± 0.13 ng/ml) and non-pregnant (7.48 ± 2.85 , 9.86 ± 0.72 ng/ml) goats at 1 and 7 days post NM, respectively. Serum P4 concentrations were significantly higher ($P < 0.01$) in pregnant (11.53 ± 0.72 , 11.60 ± 0.67 , 12.14 ± 0.56 , and 13.75 ± 0.61 ng/ml) than in non-pregnant (8.42 ± 2.06 , 0.74 ± 0.22 , 7.51 ± 2.75 , and 0.52 ± 0.16 ng/ml) goats at 15, 23, 35, and 60 days post NM, respectively (Fig. 1). ISG15 mRNA expression in the blood increased post day 1, peaked on day 23, and then declined post 35 days of NM (Fig. 2A). ISG17 mRNA expression peaked on day 35 and then declined post 35 days of NM (Fig. 2B). Relative expression mRNA for ISG15 and ISG17 were significantly higher ($P < 0.01$) in pregnant goats at 7, 15, and 35 days post NM compared to non-pregnant goats. No significant differences between pregnant and non-pregnant goats at 1 and 60 days post NM were detected (Fig. 2A, B). There were no significant differences in relative expression of mRNA for ISG15 or ISG17 between pregnant (Fig. 3A) and non-pregnant Aardi goats (Fig. 3B). ISG15, ISG17, P4, and US can be used to calculate PPV and NPV (Table I and II). Our results indicated non-significant differences in ISG15, ISG17, and P4 in PPV and NPV at 1, 7, and 15 days post NM. In addition, PPV and NPV were not significantly different in ISG15, ISG17, P4, and US at 23, 35, and 60 days post NM. Sensitivity of ISG15, ISG17, and P4 were not significantly different at 1, 7, and 15 days post NM. ISG15, ISG17, P4, and US sensitivity were not significantly different at 23 days (58.33, 58.33, 76.19, and 80.95%, respectively) post NM. US sensitivity (100%) was significantly higher ($P < 0.05$) than sensitivity of ISG15, ISG17, and P4 (50.0, 50.0, and 42.86%, respectively) at 35 days post NM. P4 sensitivity was significantly higher ($P < 0.05$) (90.48%) at 60 days post NM. In contrast, ISG15 and ISG17 sensitivity were significantly lower ($P < 0.05$) at 60 days post NM (Table III). Specificity of ISG15, ISG17, and P4 were not significantly different at 1, 7, and 15 days post NM. In addition, specificity of ISG15, ISG17, P4, and US were not significantly different at 23, 35, and 60 days post NM (Table IV). The results show that there was no significant difference in accuracy of ISG15, ISG17, and P4 at 1, 7, and 15 days post NM, and there were no significant differences in accuracy of ISG15, ISG17, P4, and US (65.56, 65.56, 75.61, and 80.91%, respectively) at 23 days post NM. However, US accuracy was significantly higher (96.97%) on day 23 ($P < 0.01$) than accuracy of ISG15, ISG17, and P4 (53.33, 46.67, and 50.61%, respectively) at 35 and 60 days post NM (100%) than ISG15 and ISG17 (37.78 and 55.56%, respectively).

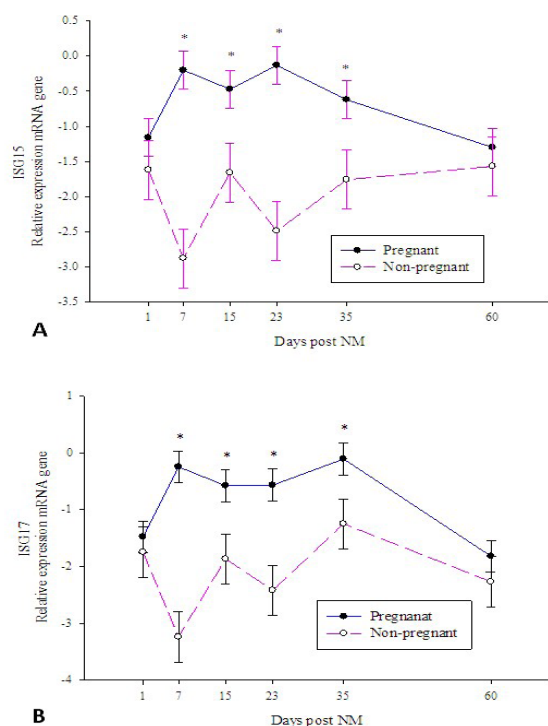


Fig. 2. Relative expression of mRNA of interferon-stimulated gene 17 (ISG17) in pregnant and non-pregnant Aardi goats at 1, 7, 15, 23, 35, and 60 days post natural mating (NM), blood samples were collected from Aardi goats, (means \pm SEM), *denotes significant difference ($P < 0.01$) between pregnant and non-pregnant goats, (n=32).

Table I. Positive predictive value (PPV) of interferon-stimulated gene 15 (ISG15), interferon-stimulated gene 17 (ISG17), progesterone hormone (P4) and ultrasound (US) for pregnancy diagnosis in goats.

Days	PPV (%) ^a			
	ISG15 (n=17)	ISG17 (n=17)	P4 (n=32)	US (n=32)
1	66.67	-	-	-
7	66.67	66.67	80.16	-
15	66.67	66.67	64.72	-
SEM	29.03	29.03	9.92	
23	66.67	66.67	83.61	90.28
35	66.67	66.67	64.44	95.83
60	50.00	100.00	87.03	100
SEM	22.29	22.29	19.93	19.93

Means \pm SEM Means within the same row with different superscript are different ($p < 0.01$), (n=32). ^aPPV (positive predictive value): probability that the goats are truly pregnant if the diagnosis is pregnant; correct pregnant/(correct pregnant + incorrect pregnant)*100. (Karen *et al.*, 2006), blood samples were collected from Aardi goats, (Goats are pregnant at P4 \geq 7.5 ng/ml).

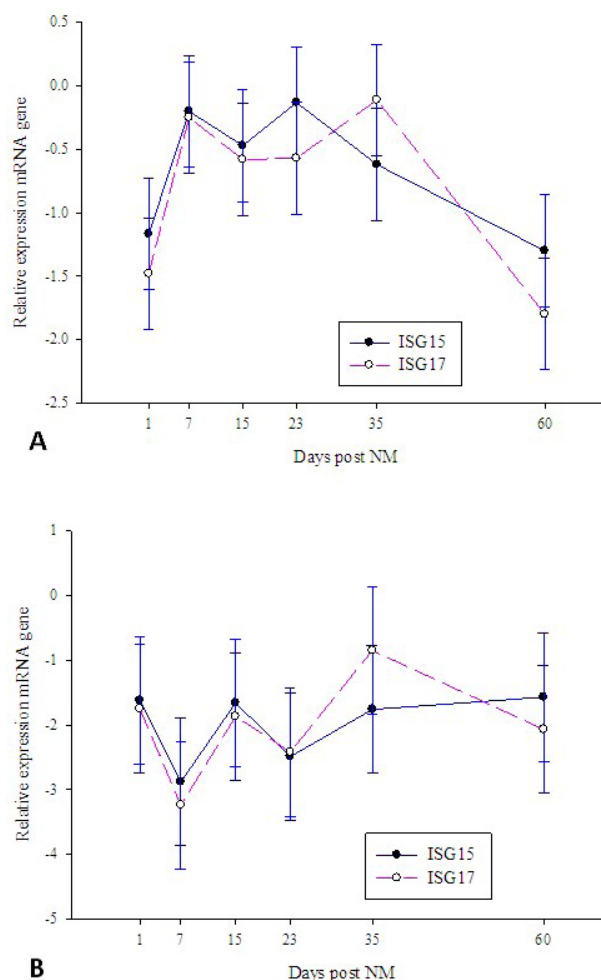


Fig. 3. Relative expression of mRNA of interferon-stimulated gene 15 (ISG15) and interferon-stimulated gene 17 (ISG17) in pregnant (A) and non pregnant Aardi goats at 1, 7, 15, 23, 35, and 60 days post natural mating (NM), blood samples were collected from Aardi goats, (means \pm SEM), (n=32).

DISCUSSION

Interferons are cytokines that communicate signals for a broad spectrum of cellular activities that encompass antiviral and immunomodulatory responses, as well as growth regulation. These pleiotropic cellular activities are mediated through a large number of proteins whose expression is triggered by activated interferon receptors present on almost all cells (Boehm *et al.*, 1997; Stark *et al.*, 1998). All effects of interferon τ on ovine endometrial gene expression, both suppression of hormone receptors and up regulation of interferon-induced genes, appear to be mediated by intracellular signal transduction systems involving type I interferon receptors (Han *et al.*, 1997;

Table II. Negative predictive value (NPV) of interferon-stimulated gene 15 (ISG15), interferon-stimulated gene 17 (ISG17), progesterone hormone (P4) and ultrasound (US) for pregnancy diagnosis in goats.

Days	NPV (%) ^a			
	ISG15 (n=17)	ISG17 (n=17)	P4 (n=32)	US (n=32)
1	33.33	100	34.24	-
7	100	100	37.50	-
15	33.33	33.33	51.39	-
SEM	25.87	25.87	10.95	
23	33.33	33.33	70.00	72.22
35	33.33	33.33	38.89	100
60	33.33	33.33	86.67	100
SEM	18.24	18.24	18.24	18.24

Means \pm SEM. Means within the same row with different superscript are different ($p < 0.01$), (n=32). ^aNPV (negative predictive value): probability that the goats are truly non-pregnant if the diagnosis is non-pregnant; correct non-pregnant / (correct non-pregnant + incorrect non-pregnant) * 100 (Karen *et al.*, 2006), blood samples were collected from Aardi goats, (Goats are pregnant at P4 \geq 7.5 ng/ml).

Table III. Sensitivity of interferon-stimulated gene 15 (ISG15), interferon-stimulated gene 17 (ISG17), progesterone hormone (P4) and ultrasound (US) for pregnancy diagnosis in goats.

Days	Sensitivity (%) ^a			
	ISG15 (n=17)	ISG17 (n=17)	P4 (n=32)	US (n=32)
1	33.33	0	0	-
7	41.67	41.67	55.36	-
15	66.67	66.67	52.38	-
SEM	17.12	17.12	13.98	
23	58.33	58.33	76.19	80.95
35	50.00 ^b	50.00 ^b	42.86 ^b	100.00 ^a
60	16.67 ^b	33.33 ^b	90.48 ^a	100.00 ^a
SEM	14.22	14.22	11.61	11.61

Means \pm SEM. Means within the same row with different superscript are different ($p < 0.05$), (n=32). ^aSensitivity: probability that diagnosis is pregnant among goats which are truly pregnant; correct pregnant / (correct pregnant + incorrect non-pregnant) * 100 (Karen *et al.*, 2006), blood samples were collected from Aardi goats, (Goats are pregnant at P4 \geq 7.5 ng/ml).

Stewart *et al.*, 2001). Interferon stimulated genes 17 (ISG17) and 15 (ISG15) are up-regulated in the uterus of ruminants, and mammals in general, in response to Interferon τ , the pregnancy recognition signal in ruminants (Charleston and Stewart, 1993; Ott *et al.*, 1998; Bebington *et al.*, 1999; Johnson *et al.*, 1999). Yankey *et*

al. (2001) reported that peripheral blood mononuclear cells were activated by conceptus-derived interferon-tau and expressed higher levels of MX mRNA in pregnant compared with non-pregnant sheep. It has been known for several years that ISG15 is expressed in greater levels in the endometrium of pregnant compared to non-pregnant cows (Austin *et al.*, 1996, 2004; Johnson *et al.*, 1998). In a study conducted to determine the impact interferon-induced gene expression in white blood cells from early pregnant cows, results showed that SG15 mRNA was up-regulated in blood from pregnant to non-pregnant cows (Hyungchul *et al.*, 2006). Study of interferon-induced gene expression in white blood cells from early pregnant goats has not been conducted until the present experiments. We tested the hypothesis that ISG15 and ISG17 mRNA were up-regulated in blood from pregnant compared to non-pregnant goats. Our results revealed that there were significantly higher ISG15 and ISG17 mRNA levels in pregnant than non-pregnant goats at 7, 15, and 35 days post NM, although no significant difference was noticed at 1 and 60 days post NM. Unfortunately, there is a lack of preexisting data regarding goats to compare with our results. Our results were in agreement with Margaret *et al.* (2005) who reported ISG15 mRNA was increased markedly on day 15, concurrent with maximal secretion of interferon-tau from the conceptus in sheep. In addition, our results agree with Johnson *et al.* (2002) who reported both ISG17 and Mx mRNA increased between days 11 and 13, and expression extended to days 15 through 17 in pregnant ewes. Furthermore, our results are in agreement with those of Hyungchul *et al.* (2006), who reported that ISG15 mRNA levels increased post day 16, peaked on day 20, and declined to day 16 levels at 32 days post artificial insemination in cows. We could not find any reference to ISG15 mRNA and ISG17 mRNA levels at 7 days post NM. Blood concentration of P4 in estrus is low (less than 1.0 ng/ml) through day 2 of diestrus and then quickly increases to maximum concentrations at 7 days, and remains elevated for up to days 13–15 following estrus (Rahman, 2006). Therefore, no difference between concentrations of serum P4 of both pregnant and non-pregnant occurs on days 1–7 from the estrous cycle, but regression of the corpus luteum (luteolysis), induced by prostaglandin F₂, (PGF_{2 α}), occurs if an embryo is not present in the uterus, which leads to rapid decline in plasma P4 (Rahman, 2006). Conversely, when an embryo is present in the uterus PGF_{2 α} secretion is inhibited. Our results showed no significant differences in P4 concentrations between non-pregnant and pregnant goats at 1 and 7 days post NM, whereas P4 concentrations were significantly higher ($P < 0.01$) in pregnant than non-pregnant at 15, 23, 35, and 60 days post NM, in agreement with the Zamfirescu *et al.* (2011). No significant differences

were found between ISG15 and ISG17 when compared to P4 and US in the prediction of pregnant and non-pregnant goats. Additionally, no significant differences were found between ISG15 and ISG17 when compared to P4 and US in accuracy at 7, 15, and 23 days post NM. Decreased accuracy of ISG15 and ISG17 at 35 days post NM might be caused by low levels of ISG15 and ISG17.

Table IV. Specificity of interferon-stimulated gene 15 (ISG15), interferon-stimulated gene 17 (ISG17), progesterone hormone (P4) and ultrasound (US) for pregnancy diagnosis in goats.

Days	Specificity (%) ^a			
	ISG15 (n=17)	ISG17(n=17)	P4 (n=32)	US (n=32)
1	80.00	80.00	100	-
7	80.00	80.00	63.89	-
15	80.00	80.00	55.56	-
SEM	26.64	26.64	15.38	
23	80.00	80.00	72.22	80.56
35	60.00	40.00	63.89	91.67
60	80.00	100.00	75.00	100.00
SEM	14.83	14.83	8.56	8.56

Means \pm SEM within the same row with different superscript are different ($p < 0.01$), (n=32). ^aSpecificity: probability that diagnosis is non-pregnant among goats which are truly non-pregnant; correct non-pregnant/(correct non-pregnant + incorrect pregnant)*100 (Karen *et al.*, 2006), blood samples were collected from Aardi goats, (Goats are pregnant at P4 \geq 7.5 ng/ml).

Table V. Accuracy of interferon-stimulated gene 15 (ISG15), interferon-stimulated gene 17 (ISG17), progesterone hormone (P4) and ultrasound (US) for pregnancy diagnosis in goats.

Days	Accuracy (%) ^a			
	ISG15 (n=17)	ISG17(n=17)	P4 (n=32)	US (n=32)
1	48.89	26.67	34.24	-
7	54.44	54.44	56.67	-
15	71.11	71.11	54.24	-
SEM	15.75	15.75	15.75	
23	65.56	65.56	75.45	80.91
35	53.33 ^b	46.67 ^b	50.61 ^b	96.97 ^a
60	37.78 ^b	55.56 ^b	84.85 ^{ab}	100 ^a
SEM	12.71	12.71	12.71	12.71

Means \pm SEM within the same row with different superscript are different ($p < 0.01$), (n=32). ^aAccuracy: The accuracy of the diagnosis test; (correct pregnant + correct non-pregnant)/(correct pregnant + correct non-pregnant + incorrect non-pregnant + incorrect pregnant)*100 (Karen *et al.*, 2006), blood samples were collected from Aardi goats, (Goats are pregnant at P4 \geq 7.5 ng/ml).

CONCLUSION

ISG15 and ISG17 were used in the diagnosis of early pregnancy in sheep and cattle. To our knowledge, this is the first time that ISG15 and ISG17 were used for the diagnosis of early pregnancy in goats. ISG15 and ISG17 provide accurate pregnancy diagnoses at 15 and 23 days post natural mating.

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Supplementary material

There is supplementary material associated with this article. Access the material online at: <https://dx.doi.org/10.17582/journal.pjz/20180329180335>

Statement of conflict of interest

The authors have declared no conflict of interest.

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