

## Oocyte-derived growth factors and ovulation rate in sheep

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The physiological mechanisms controlling ovulation rate in mammals involve a complex exchange of endocrine signals between the pituitary gland and the ovary, and a localized exchange of intraovarian hormones between the oocyte and its adjacent somatic cells. The discoveries in sheep of mutations in bone morphogenetic protein 15 (BMP15) and bone morphogenetic protein receptor type IB (BMPRI-IB) together with recent findings on the physiological effects of growth differentiation factor 9 (GDF9) and BMP15 on follicular development and ovulation rate highlight some important differences in the way in which the oocyte may function in mammals with different ovulation rate phenotypes. In sheep, BMP15 and GDF9 have each been shown to be essential for the early and later stages of follicular development. In addition, ovulation rate is sensitive to changes in the dose of either of these two oocyte-derived growth factors. These findings are in contrast to those reported for mice in which GDF9, but not BMP15, is essential for follicular development. The evidence to date is consistent with the hypothesis that the oocyte plays a central role in regulating key events in the process of follicular development and hence, is important in determining ovulation rate. Moreover, it appears that the mechanisms that the oocyte uses to control these processes differ between species with low and high ovulation rate phenotypes.

## Introduction

Humans, cattle, goats and sheep typically have an ovulation rate of one or two, whereas other mammals, such as rats, mice, dogs and pigs have ovulation rates of four or more. Thus, when asking fundamental questions as to why some individual domestic livestock species, or indeed humans, have a predisposition to produce two or three offspring rather than one, it is useful to study experimental animal models with a low ovulation rate phenotype. In this context, sheep are proving to be remarkably informative experimental models (McNatty *et al.*, 2001; Montgomery *et al.*, 2001).

Control of ovulation rate in mammals not only involves a complex exchange of hormonal signals between the pituitary gland and the ovary, but also is dependent on local communication, via intraovarian hormones and growth factors between oocytes and their adjacent somatic cells (Scaramuzzi *et al.*, 1993; Baird and Campbell, 1998; McNatty *et al.*, 1999; Galloway *et al.*, 2000; Eppig, 2001; Juengel *et al.*, 2002). The temporal interrelationships between the pituitary hormones, FSH and LH, and the ovarian hormones, oestradiol, progesterone and inhibin, are reasonably well understood. Administration of supplementary FSH to predominantly mono-ovulatory species, such as humans, cattle and sheep, can over-ride the endogenous control mechanisms that govern ovulation rate and stimulate multiple ovulations (that is,  $\geq 4$ ). In addition, small temporal changes in plasma FSH concentrations around the time of luteal regression appear to be the physiological basis that explains why some sheep ovulate two or three eggs rather than one (McNatty *et al.*, 1985, 1988; Henderson *et al.*, 1988). Although higher than normal FSH concentrations may explain, in part, why some Booroola ewes have exceptionally high ovulation rates (between 8 and 13; Hudson *et al.*, 1999), it is important to emphasize that changes in plasma FSH concentrations alone do not explain all the observed differences in ovulation rate among breeds of sheep (Driancourt *et al.*, 1988; Shackell *et al.*, 1993; Fry and Driancourt, 1996) or indeed why some species always ovulate several follicles and others mainly one. Results from studies with mice lacking certain growth factor genes or sheep with naturally occurring genetic mutations in certain oocyte-derived growth factors provide evidence that the oocyte plays an essential role not only in regulating the growth and maturation of ovarian follicles, but also in determining ovulation rate (Dong *et al.*, 1996; Galloway *et al.*, 2000). As the oocyte has such a major effect on follicular growth and ovulation rate, it is reasonable to hypothesize that the relative importance, and indeed actions, of certain oocyte-derived growth factors may vary between mammals with different ovulation rate phenotypes.

The purpose of this review is to summarize recent findings on single genetic mutations that affect ovulation rate in sheep. Moreover, these data, together with new evidence from physiological studies, are presented to support the hypothesis that follicular growth and ovulation rate are influenced, in a dose-dependent manner, by oocyte-derived growth factors that influence somatic cells of the ovarian follicle and that the importance of these growth factors differs between mammals with a low or high ovulation rate phenotype.

### Sheep with single genetic mutations affecting ovulation rate

The inheritance patterns of several naturally occurring genetic mutations that cause anovulation or either modest (0.2–0.4) or large ( $\geq 0.8$ ) increases in ovulation rate in sheep have been identified (McNatty *et al.*, 2001; Davis *et al.*, 2002). In several of these sheep families, namely, Inverdale (FecX<sup>I</sup>), Hanna (FecX<sup>H</sup>), Booroola (FecB<sup>B</sup>), Garole (FecB<sup>B</sup>) and Javanese (FecB<sup>B</sup>), the inherited mutation has been mapped to a specific region of the sheep X chromosome (Inverdale, Hanna) or sheep chromosome 6 (Booroola, Garole and Javanese) (Table 1).

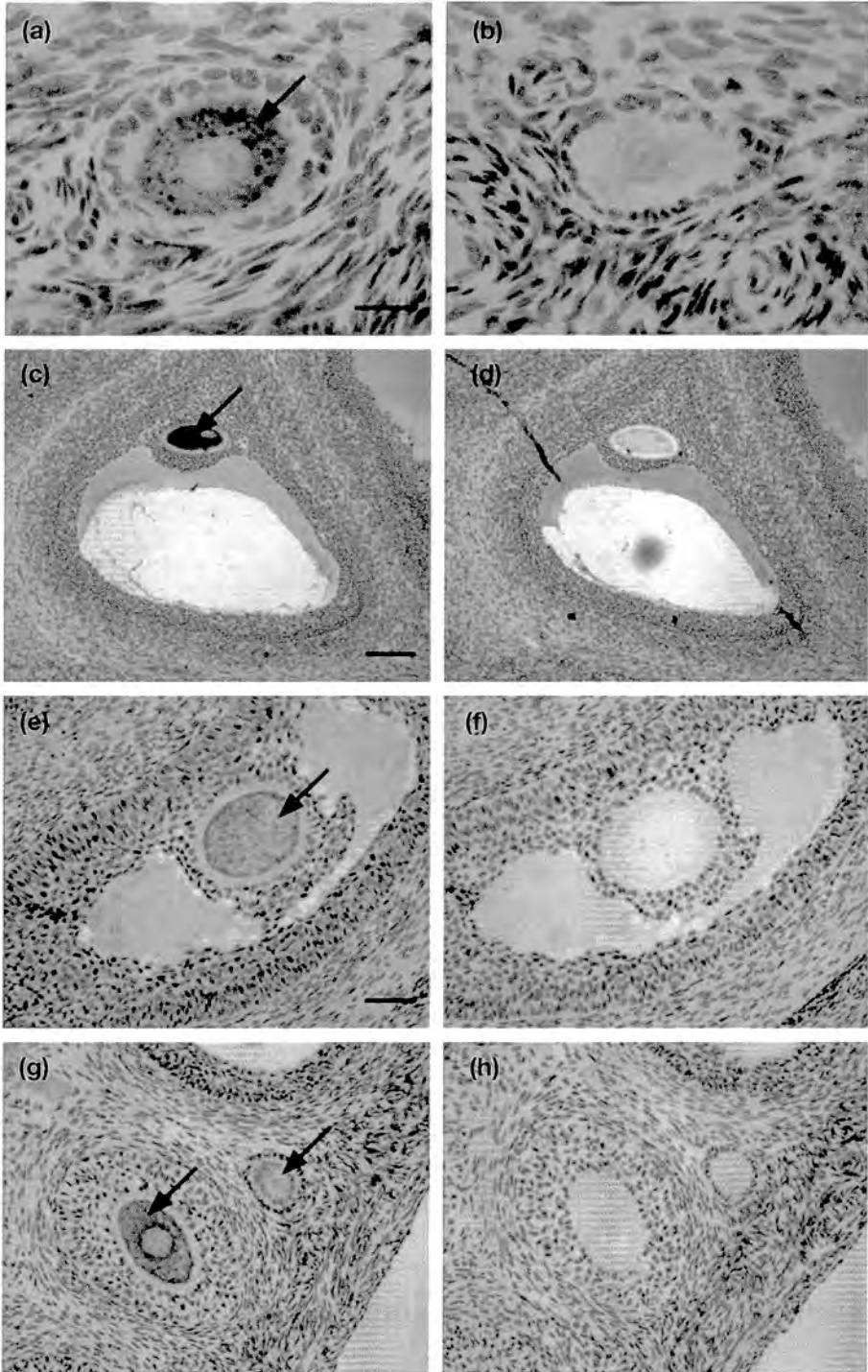
**Table 1.** Mutations in an oocyte-derived growth factor or a growth factor receptor in different sheep families, and the effects of genotype on mean ovulation rate and litter size

Gene identification (Chromosome number)	Sheep family (Allele)	Phenotype		Reference
		(Ovulation rate)	(Litter size)	
BMP15 (X)	Inverdale (FecX <sup>I</sup> )	++ (1.8)	++ (1.6)	Galloway <i>et al.</i> (2000)
		I+ (2.9)	I+ (2.3)	Davis <i>et al.</i> (2001a)
		II (0)	II (0)	
	Hanna (FecX <sup>H</sup> )	++ (1.8)	++ (1.6)	Galloway <i>et al.</i> (2000)
		H+ (2.9)	H+ (2.2)	Davis <i>et al.</i> (2001a)
		HH (0)	HH (0)	
BMPR-IB (6)	Booroola (FecB <sup>B</sup> )	++ (1.5)	++ (1.3)	Wilson <i>et al.</i> (2001)
		B+ (2.8)	B+ (2.2)	Mulsant <i>et al.</i> (2001)
		BB (4.6)	BB (2.7)	Souza <i>et al.</i> (2001)
				Davis <i>et al.</i> (1982)
	Garole (FecB <sup>B</sup> )	Unknown	BB (2.2+)	Davis <i>et al.</i> (2002)
	Javanese (FecB <sup>B</sup> )	++ (1.4)	++ (1.2)	Davis <i>et al.</i> (2002)
	B+ (2.7)	B+ (1.9)		
	BB (unknown)	BB (2.6)		

BMP15: bone morphogenetic protein 15; BMPR-IB: bone morphogenetic protein IB receptor.

In each of these animals, a point mutation was identified in genes from the transforming growth factor  $\beta$  (TGF $\beta$ ) superfamily or their receptors (Table 1). In Inverdale and Hanna ewes, separate point mutations were identified in the bone morphogenetic protein 15 (BMP15) gene corresponding to sites in the mature peptide coding region of the BMP15 growth factor (also known as growth differentiation factor 9B; GDF9B) (Galloway *et al.*, 2000). The ovine BMP15 mature protein consists of 125 amino acids. In FecX<sup>H</sup> sheep, a single C to T transition at nucleotide position 67 of the mature peptide coding region introduces a premature stop codon in the place of glutamic acid at amino acid residue 23 leading to a severe truncation and likely total loss of BMP15 bioactivity. In FecX<sup>I</sup> ewes, a single T to A transition occurs at nucleotide position 92 of the mature peptide, thereby substituting a valine with aspartic acid at amino acid residue 31 of the mature protein. This mutation represents a non-conservative change in a highly conserved region of the protein and probably results in a marked reduction in BMP15 bioactivity in FecX<sup>I</sup> ewes. Evidence to support this claim comes from phenotypic evaluation of the Hanna (H)–Inverdale (I) crossed ewe (HI). As with the homozygous Inverdale and Hanna ewes (Fig. 1), the HI animals are sterile (Galloway *et al.*, 2000). A remarkable characteristic of Hanna or Inverdale ewes is that ewes that are heterozygous for the FecXH or FecXI mutation, have higher than normal ovulation rates and litter sizes (Davis *et al.*, 2001a). Thus, for some reason, half the normal genetic dose of BMP15 leads to a higher than normal ovulation rate. No major effects of the FecX<sup>H</sup> or FecX<sup>I</sup> mutation are evident in males.

In Booroola, Garole and Javanese sheep, a point mutation was identified in the highly conserved intracellular serine threonine kinase signalling domain of the BMP-IB receptor (BMPR-IB) (Mulsant *et al.*, 2001; Souza *et al.*, 2001; Wilson *et al.*, 2001; Davis *et al.*, 2002). In FecB<sup>B</sup> sheep, a single A to G transition occurs at nucleotide position 830, thereby substituting a glutamine (neutral/polar amino acid) for arginine (basic amino acid) at position 249 of the protein. This transition represents a non-conservative change in a highly conserved region of



**Fig. 1.** Immunohistochemical evidence for the localization of growth differentiation factor 9 (GDF9) protein and bone morphogenetic protein 15 (BMP15) protein in oocytes (arrows) within developing ovarian follicles of 4-week-old ewe lambs. (a) GDF9

a type 1 BMP receptor. The consequences are that females with one copy of this mutation have an ovulation rate of about one or two eggs more than the non-carriers and those with two copies have ovulation rates of about three to ten eggs more than non-carriers (McNatty *et al.*, 2001; Davis *et al.*, 2002). No major effects of the *FecB<sup>B</sup>* mutation are evident in male sheep (Hochereau-de Reviers and Seck, 1991; Smith *et al.*, 1996).

### BMP growth factors, BMP receptors and ovarian follicular development in sheep

For the purposes of this review, the terminology described by Lundy *et al.* (1999) will be used to define the specific stages of ovarian follicular development in sheep, namely types 1, 1a, 2, 3, 4 and 5 for primordial, transitory, primary, small preantral, large preantral and antral follicles, respectively.

The mechanisms by which mutations in BMP15 or BMPR-IB influence ovulation rate in sheep are not well understood. Results from gene or protein expression studies in sheep ovaries show that BMP15, GDF9 and BMPR-IB are all expressed in oocytes (Fig. 1). However, whereas BMP15 and GDF9 are expressed exclusively in oocytes, BMPR-IB is expressed in oocytes, granulosa cells and luteal cells (Bodensteiner *et al.*, 1999, 2000; Galloway *et al.*, 2000; Wilson *et al.*, 2001; Juengel *et al.*, 2002; Souza *et al.*, 2002). GDF9 mRNA and protein are expressed in most oocytes from the type 1 to type 1a stage of follicular development (Bodensteiner *et al.*, 1999, 2000; Fig. 1) and BMP15 from the type 2 stage (Galloway *et al.*, 2000; Fig. 1). Although both BMPR-IB and BMPR-II are expressed in oocytes and BMPR-II is expressed in granulosa cells from the type 1 stage, expression of BMPR-IB in granulosa cells is not evident until the type 2 stage of development (Wilson *et al.*, 2001; Souza *et al.*, 2002). Other TGF $\beta$  growth factors in sheep, such as  $\beta_B$  inhibin-activin subunit or anti-Müllerian hormone and  $\alpha$  inhibin-activin subunit, are expressed in granulosa cells from the type 2 and type 3 stages of growth, respectively (Montgomery *et al.*, 2001).

In homozygous carriers of the Inverdale (II) and Hanna (HH) mutations, normal ovarian follicular development is blocked from the type 2 stage of growth (Braw-Tal *et al.*, 1993; Galloway *et al.*, 2000). However, in II and HH ewes, oocytes can continue to grow from approximately 50  $\mu\text{m}$  in diameter in type 2 follicles to approximately 80–100  $\mu\text{m}$ , which is typical of the diameter observed in type 4 or type 5 follicles without a concomitant increase in the number of granulosa cells (Braw-Tal *et al.*, 1993; Galloway *et al.*, 2000). These results imply that the oocyte is capable of growing independently of granulosa cells but that

#### Fig. 1. continued

localization in an oocyte of a type 2 (primary) follicle identified using a monoclonal antibody to GDF9 (Clone 37, 5  $\mu\text{g ml}^{-1}$ ); (b) non-specific binding on an adjacent tissue section to (a) of an irrelevant polyclonal mouse antibody (that is, bovine  $\alpha$  lactalbumin (M1-Lm, 1.4  $\mu\text{g ml}^{-1}$ )); (c) GDF9 localization in an oocyte of a type 5 (antral) follicle identified using Clone 37 at 5  $\mu\text{g ml}^{-1}$ ; (d) non-specific binding on an adjacent tissue section using an irrelevant monoclonal mouse antibody (YD3; 5  $\mu\text{g ml}^{-1}$ ); (e) BMP15 localization in a type 5 follicle identified using affinity-purified polyclonal anti-BMP15 (M10-9B, 2  $\mu\text{g ml}^{-1}$ ); (f) non-specific binding in an adjacent tissue section after preincubating M10-9B (2  $\mu\text{g ml}^{-1}$ ) with an *Escherichia coli* generated oBMP15 mature protein (10  $\mu\text{g ml}^{-1}$ ); (g) BMP15 localization in oocytes (arrows) of a type 2 and type 4 (large preantral) follicle identified using M10-9B (2  $\mu\text{g ml}^{-1}$ ) antibodies; and (h) non-specific binding on an adjacent tissue section after preincubating M10-9B and oBMP15 mature protein. For experimental details see Juengel *et al.* (2002). Scale bars represent (a,b) 20  $\mu\text{m}$ , (c,d) 100  $\mu\text{m}$  and (e–h) 50  $\mu\text{m}$ .

oocyte-derived BMP15 is essential for the proliferation of granulosa cells for the advancement of follicles beyond the type 2 stage of growth.

Currently, there is uncertainty about the nature of molecular forms of BMP15 in biological fluids. It is thought that BMP15 and GDF9 function as homodimers or heterodimers, but until appropriate biologically active standards and highly specific antibodies become available, this question cannot be answered. However, it is known that GDF9 and BMP15 protein can be detected in oocytes and that the protein cannot be localized within any other type of ovarian cell (Fig. 1). Current evidence indicates that BMP15, in the absence of other TGF $\beta$  superfamily members, acts to stimulate DNA synthesis in granulosa cells of mammals with low and high ovulation rates. The addition of recombinant human BMP15 to oestrogen-stimulated rat granulosa cells for 24 h (Otsuka *et al.*, 2000) or a recombinant ovine BMP15 to ovine granulosa cells for 48 h (K. Reader, S. Lun, A. Western, S. Lawrence and K. P. McNatty, unpublished) stimulates a 2.0–2.5-fold increase in [ $^3$ H]thymidine incorporation *in vitro*. In addition, Otsuka *et al.* (2001) reported that recombinant BMP15 inhibits the biological actions of FSH by suppression of FSH receptor gene expression. They hypothesized from this finding that the higher ovulation rate in heterozygous Inverdale and Hanna ewes is due to a greater responsiveness of granulosa cells to a given amount of FSH because of a 'lowered dose' of oocyte-derived BMP15. The evidence from studies of granulosa cells in Inverdale ewes is that the onset of FSH receptor gene expression, which begins at the type 3 stage of follicle development, remains unchanged between the heterozygotes and non-carriers of the mutated BMP15 gene (Juengel *et al.*, 2000). However, it is known that the FSH-induced cAMP responsiveness of granulosa cells from heterozygous Inverdale ewes *in vitro* is higher than in non-carriers and that granulosa cells from heterozygous animals acquire cAMP responsive LH receptors earlier in follicular development (Shackell *et al.*, 1993). Whether this increased responsiveness to FSH in granulosa cells from Inverdale ewes is the result of an increased number of receptors or changes in the ability of the receptor to stimulate cAMP has not been determined.

Collectively, studies on Inverdale ewes heterozygous for an inactivating BMP15 mutation show that there are a greater number of smaller diameter antral follicles and that some of these follicles can undergo preovulatory maturation prematurely (that is, relative to wild-type ewes) and ovulate at a smaller follicular diameter (Shackell *et al.*, 1993). The consequence of these changes is, on average, one more ovulation than in control sheep and the formation of smaller corpora lutea. The heterozygous Inverdale, and presumably Hanna, ewes are examples of sheep that have a higher ovulation rate than their littermate controls without any significant change in plasma FSH concentration. It is reasonable to hypothesize that the difference in ovulation rate in these genotypes is due entirely to a change in the intrafollicular concentrations of the oocyte-derived BMP15. Although in mice with heterozygous or homozygous inactivations of the BMP15 gene, there are no major effects on follicular development or ovulation rate, there is a modest reduction in litter size in knockout mice that is probably due to fertilization failure (Yan *et al.*, 2001). These findings represent a fundamental difference between sheep and rodents.

The BMPR-IB mutation identified in Booroola and other breeds of sheep (Table 1) is a clear example of how a change in BMP signalling can lead to a significantly earlier maturation of differentiative functions in ovarian follicles. As a result, follicles become capable of ovulating after 16–17 doublings of the granulosa cell population instead of the normal 18–19 doublings observed in the wildtype (for a review, see Montgomery *et al.*, 2001; Mulsant *et al.*, 2001). Earlier maturation in follicles of Booroola ewes is evident by the type 3 stage of follicular development (Wilson *et al.*, 2001). Moreover, from studies in hypophysectomized Booroola ewes, it is evident that follicles become gonadotrophin-dependent once they

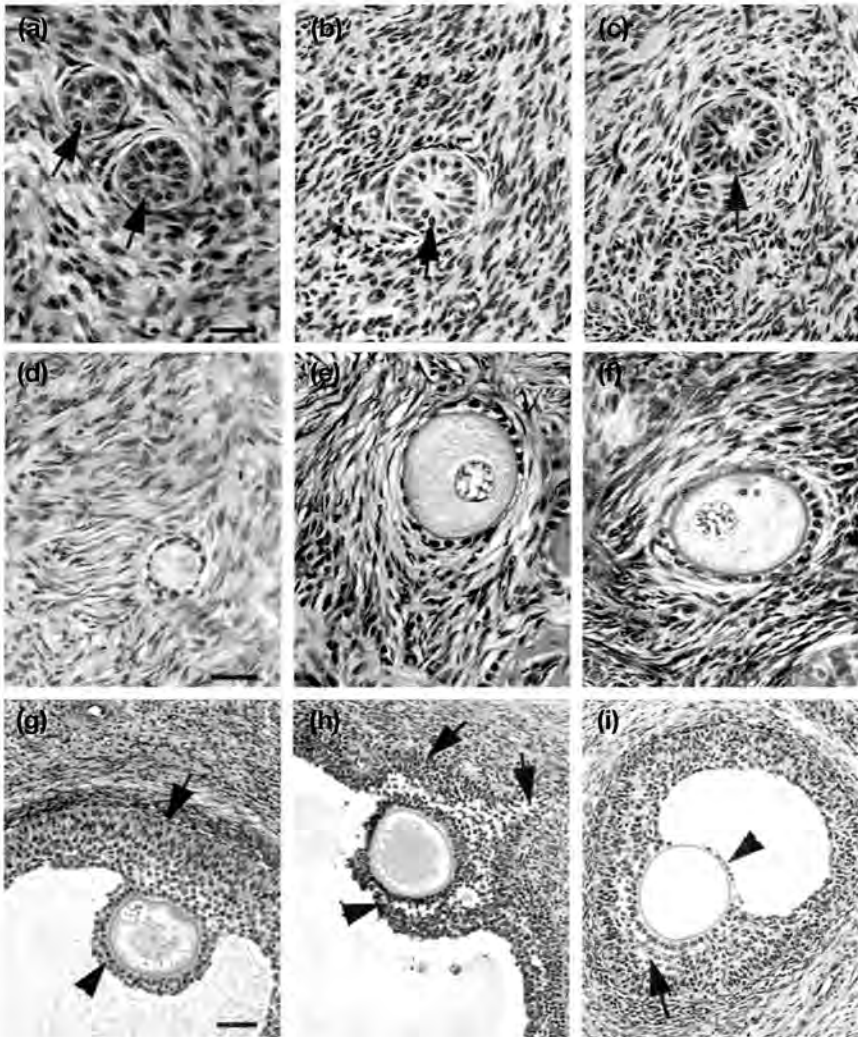
reach 2 mm in diameter, whereas in wild-type hypophysectomized Booroola ewes, follicles become gonadotrophin-dependent at about 3 mm in diameter, which is the equivalent of one granulosa cell doubling cycle later (McNatty *et al.*, 1990). It remains unclear at present whether the earlier maturation process in follicles or the increased ovulation rate from the BMPR-IB mutation is driven by changes in oocyte regulation of granulosa cell function or through an interactive signalling exchange between these two types of cell. BMPR-IB mRNA has been identified in many tissues other than the ovary, such as brain, pituitary gland, skeletal muscle, testis, prostate gland and reproductive tract (Wilson *et al.*, 2001). Thus, it is not surprising that the Booroola mutation has marked effects on organ development and on pituitary gland function (McNatty *et al.*, 1995; Montgomery *et al.*, 2001). Unlike the Inverdale and Hanna genotypes, in which all the major effects of the mutations are localized to the ovary, some effects in Booroola may be mediated via the pituitary gland as a result of higher than normal secretions of FSH (McNatty *et al.*, 1991, 1993). At present, there is no equivalent mutation observed in mice or other high ovulation rate phenotypes with which to compare Booroola.

### Immunization of sheep against the oocyte-derived growth factors GDF9 and BMP15

It has been suggested that some of the differences in the roles of BMP15 between rodents and sheep are due to species differences in the actions of GDF9 and BMP15: GDF9 is essential for follicular development in rodents and BMP15 is essential in sheep (Yan *et al.*, 2001). However, this notion presupposes that GDF9 is ineffectual or less important in sheep.

The importance of GDF9 in sheep was tested by Juengel *et al.* (2002) who immunized ewes (9–10 per treatment group) with keyhole limpet haemocyanin (KLH) as a control, an ovine GDF9-specific peptide conjugated to KLH (GDF9 peptide) or an ovine BMP15 specific peptide conjugated to KLH (BMP15 peptide) at monthly intervals before and during the breeding season for 7 months. Ewes immunized with KLH maintained regular oestrous cycles (that is,  $17 \pm 2$  days) during the breeding season. In contrast, none of the ewes immunized with GDF9 peptide experienced normal oestrous cycles of 17 days and most ewes (seven of ten) had no evidence of corpora lutea at the time the ovaries were recovered. Plasma concentrations of progesterone in the three ewes with corpora lutea were highly irregular and abnormal and not characteristic of a normal oestrous cycle. Many of the ewes without corpora lutea had no visible antral follicles at ovarian collection. Only one of ten of the ewes immunized with BMP15 peptide displayed regular oestrous cycles and this ewe had an average ovulation rate of six. In none of the other ewes immunized with BMP15 peptide was a corpus luteum visible on the surface of the ovary, and as with the GDF9 peptide-immunized group, many had no visible antral follicles at the time the ovaries were collected.

When the ovaries of these animals were examined histologically and subjected to morphometric analysis, it was evident that in both the GDF9 peptide and BMP15 peptide immunized groups, most ewes had few, if any, normal follicles beyond the type 2 stage of development. Of particular interest was the finding of numerous oocyte-free nodules of granulosa cells in both the GDF9- and BMP15-peptide immunized groups. These nodules were identical to those typically observed in homozygous Inverdale or Hanna ewes (Fig. 2) (Braw-Tal *et al.*, 1993; Smith *et al.*, 1997; Galloway *et al.*, 2000). Although follicles beyond the type 2 stage of development were observed, these follicles often had unusually large oocytes relative to those in the control (that is, KLH treated) animals and the organization of the granulosa cells or the cumulus cells around the oocyte was abnormal (Fig. 2). These abnormal changes in morphology of the oocyte and granulosa cells in the GDF9- and



**Fig. 2.** Photomicrographs of sections through normal and abnormal follicles from carriers and non-carriers of the Inverdale gene or after long-term immunization with growth differentiation factor 9 (GDF9) or bone morphogenetic protein 15 (BMP15) peptide (Juengel *et al.*, 2002). (a) Two oocyte-free follicles (arrows) in a ewe homozygous for the *FecX*<sup>1</sup> gene; (b) oocyte-free follicle (arrow) in a GDF9 peptide immunized ewe; (c) oocyte-free follicle (arrow) in a BMP15 peptide immunized ewe; (d) a type 1a (transitory) follicle in an Inverdale wild-type ewe; (e) an abnormal type 1 (primordial) follicle in an Inverdale ewe homozygous for the *FecX*<sup>1</sup> gene; (f) an abnormal type 1a follicle in a BMP15 peptide immunized ewe; (g) a normal antral follicle in an Inverdale wild-type ewe with a uniform distribution of cumulus cells around the oocyte (arrowhead) and an organized and uniform arrangement of granulosa cells around the basement membrane (arrow); (h) an antral follicle with abnormal appearance in a GDF9 peptide immunized ewe with a disorganized arrangement of cumulus cells (arrowhead) and also a disorganized array of granulosa cells around the basement membrane which appears highly irregular (arrows). (i) An abnormal-looking antral follicle in a BMP15 peptide immunized ewe with an abnormally enlarged oocyte, incomplete arrangement of cumulus cells (arrowhead) and extensive intercellular spaces (arrow) between cumulus cells. Scale bars represent (a–c) 50  $\mu$ m and (d–i) 30  $\mu$ m.



BMP15 peptide-immunized sheep were first evident at the type 1a and 2 stages of development (Fig. 2). Morphometric analysis revealed that the mean total numbers of types 4 and 5 follicles were significantly lower in the GDF9 peptide- and BMP15 peptide-immunized ewes compared with the KLH group. In the KLH-immunized ewes, the mean number of follicles  $\geq 1$  mm in diameter was ten, whereas in the GDF9 peptide and BMP15 peptide immunized groups, the mean numbers were two and  $< 1$ , respectively. In these studies, evidence was found that the ewes immunized with the GDF9 peptide contained antibodies against *Escherichia coli* generated ovine (o)GDF9 protein but no crossreactivity (that is,  $< 1\%$ ) to *E. coli* generated oBMP15 protein. Likewise, ewes immunized with the BMP15 peptide contained antibodies against oBMP15 protein but not against oGDF9 protein (that is,  $< 1\%$ ).

Collectively, these data indicate that GDF9, as well as BMP15, is essential for ovarian follicular development in sheep. Moreover, both growth factors are required very early in follicular development, namely from the type 1a and type 2 stage of development. These findings are in contrast to those in mice in which GDF9, but not BMP15, is required for the early stages of follicular development. After these observations, the question arose as to whether GDF9 and BMP15 are of any importance after follicles reach the antral stages of growth and become totally dependent on gonadotrophins for subsequent maturation and ovulation. Studies in rodents indicate that GDF9 is important for establishing the cumulus cell phenotype and that both GDF9 and BMP15 modulate the actions of FSH on differentiation of mural granulosa cells (Elvin *et al.*, 1999; Erickson and Shimasaki, 2000; Vitt *et al.*, 2000; Eppig, 2001; Otsuka *et al.*, 2001). The importance of GDF9 and BMP15 in sheep was tested by giving each ewe (five per group) 100 ml of a pool of antiplasma to GDF9 peptide, BMP15 peptide or KLH. The antiplasma for GDF9 or BMP15 peptide was collected from the aforementioned GDF9 or BMP15 immunized ewes that were anovulatory (Juengel *et al.*, 2002). The antiplasmas were administered i.v. at day 4 before the induction of luteal regression or approximately 6–7 days before ovulation and all ewes were subjected to laparoscopy 8 days after the expected day of ovulation. The ewes that received the antiplasma against the GDF9 peptide formed one to two corpora lutea but three of five ewes did not display normal luteal phase patterns of progesterone concentrations. Four of the five ewes that received the antiplasma against BMP15 peptide failed to ovulate and the ovaries of three of five ewes were devoid of visible follicles. By contrast, ewes that were given KLH antiplasma ovulated with normal luteal function as assessed from the plasma progesterone concentrations. These data indicate that both GDF9 and BMP15 are important in regulating the later stages of follicular maturation as well as being essential for normal follicular growth.

From the long-term immunization studies, it was unclear whether partial neutralization of GDF9 was able to enhance ovulation rates (Juengel *et al.*, 2002). This question was addressed in a preliminary study using a short-term immunization regimen with a weaker adjuvant than the Freund's complete or incomplete system used in the study by Juengel *et al.* (2002). It was hypothesized that a limited number of immunizations of ewes with GDF9 or BMP15 in a diethylaminoethyl dextran 5% (w/v) PBS adjuvant would lead to a more modest antibody response and, thus, only a partial neutralization of GDF9 or BMP15. It was anticipated from such a treatment that some of the ewes immunized with BMP15 might have a higher than normal ovulation rate for BMP15-treated animals and the question as to whether GDF9 caused increases in ovulation rate might be resolved. Accordingly, ewes (eight per treatment group) were immunized on three occasions, at 1 month intervals, with KLH or KLH conjugated to GDF9 peptide (GDF9 peptide), BMP15 peptide (BMP15 peptide), oGDF9 protein (encoding the entire mature region) or oBMP15 protein (encoding the entire mature region). The ewes received 0.4 mg antigen at the primary injection and 0.2 mg boosters at the second and third injections. The ovaries of all ewes were examined by laparoscopy 14 days after the

**Table 2.** Number of ewes with average ovulation rates of 0, 1–3 or  $\geq 4$  after immunization with KLH or KLH conjugated to GDF-9 peptide, oGDF9 protein, BMP15 peptide or oBMP15 protein

Treatment	Mean ovulation rate		
	0	1–3	$\geq 4$
KLH	0	8	0
GDF9 peptide	0	7	1
oGDF9 protein	3	1	4
BMP15 peptide	0	2	6
oBMP15 protein	1	4	3

BMP: bone morphogenetic protein 15; GDF9: growth differentiation factor 9; KLH: keyhole limpet haemocyanin. H. Sawyer, C. Moeller, N. Hudson, J. Juengel, K. McNatty, unpublished.

administration of the third injection when the ewes were also given a second prostaglandin  $F_{2\alpha}$  injection (10 days after the first). In addition, the ovaries of all ewes were examined at slaughter at day 19 after laparoscopy. The range of ovulation rates recorded after first taking the average of the number of corpora lutea from the two successive ovulations for each ewe are shown (Table 2). Immunization of ewes with GDF9 peptide resulted in one ewe having an ovulation rate of 10, which was the average of two successive ovulations of 12 and 8. Immunization of ewes with the oGDF9 protein either resulted in most ewes having a higher than normal ovulation rate ( $\geq 4$ ) or anovulation. Treatment with either BMP15 peptide or oBMP protein led to 75% and 37.5% of ewes, respectively, having ovulation rates of  $\geq 4$ .

Examination of sera from ewes immunized against GDF9 peptide or protein revealed the presence of antibodies against oGDF9 protein, but no crossreactivity (<2%) to *E. coli* generated oBMP15 protein. Likewise, ewes immunized with the BMP15 peptide or protein contained antibodies against oBMP15 protein, but not against oGDF9 protein (<2%).

In all the aforementioned immunization studies particular attention was paid to whether ewes immunized with GDF9 had crossreactivities with the closely related BMP15 and vice versa. In no instance were measurable crossreactivities noted. As these proteins are much more closely related to each other than to other members of the TGF $\beta$  superfamily it is likely that the effects observed were specific for neutralization of the target protein.

These data indicate that GDF9 as well as BMP15 may enhance ovulation rate in sheep. Moreover, it appears that the dose of oocyte-derived GDF9 or BMP15 acting on the adjacent granulosa cells has profound effects on ovulation rate. These findings, taken together with the long-term immunization studies, support the hypothesis that in mammals with a low ovulation rate phenotype (for example, sheep), the oocyte has the ability to regulate both follicular development and ovulation rate through at least two different growth factors, GDF9 and BMP15.

## Conclusions

From studies on sheep with unusual ovulation rates or fecundity, namely Booroola, Inverdale, Hanna, Garole and Javanese (Montgomery *et al.*, 2001; Davis *et al.*, 2001a,b; 2002), it is evident that there are now several major genetic mutations in sheep that lead to increased fecundity. In addition, it seems likely that several more fecundity-enhancing mutations will be identified in the future (for example, Woodlands: Davis *et al.*, 2001b; for a review, see McNatty *et al.*, 2001). When these genes and the mutations are identified, appropriate genetic marker tests can be developed (Davis *et al.*, 2002). Collectively, these diagnostic tests have the potential to change farm management practices profoundly through rapid improvements in reproductive efficiency.

The recent discoveries in sheep of the mutations in BMP15 and BMP15 together with recent results on the physiological effects of GDF9 and BMP15 on ovulation rate highlight some important differences in the way the oocyte may function in mammals with different ovulation rates. The evidence to date is consistent with the hypothesis that both follicular growth and ovulation rate are influenced by the dose of BMP15 or GDF9 delivered to the somatic cells of the follicle in mammals with a low ovulation rate phenotype, whereas in mammals with a high ovulation rate phenotype, the follicular somatic cells are relatively insensitive to changes in the dose of BMP15 but have an absolute requirement for GDF9.

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