

The Effect of Insulin-Like Growth Factor System on Embryo Growth and Development

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Abstract: Insulin like growth factors (IGF-I and IGF-II) are expressed in embryos and reproductive tracts of several species including cow, sheep and swine. They are mitogenic and have endocrine, paracrine and autocrine function infusing cell division, blastocyst formation, implantation and embryo growth. Increase in embryo growth will probably result with a higher implantation rates leading to consequent increases in the number of live offspring. In this review, insulin, IGFs, their receptors and their physiology and function in embryonic growth development were concerned.

Key words: Development, embryo, growth, IGF, IGFBP, insulin

INTRODUCTION

Insulin synthesized in the pancreas within the beta cells (β -cells) of the islets of Langerhans as a 6 kDa endocrine protein composed of two peptide chains referred to as the A and B held together by disulfide bonds. The receptors for insulin have been shown in embryos of rabbit, mouse, rat, cow and sheep (Mattson *et al.*, 1988; Harvey *et al.*, 1995; Kaye and Harvey, 1995; Santos *et al.*, 2004) and the beneficial effect of insulin on development of preimplantation embryos in mice (Gardner and Kaye, 1991), rats (Hertogh *et al.*, 1991), pigs (Lewis *et al.*, 1992) and cattle (Matsui *et al.*, 1995) have been reported.

The amino acid sequences of insulin and insulin-like growth factors (IGF-I and IGF-II), from all vertebrate groups, are currently known and there is a high degree of similarity for each hormone among the different vertebrates (Reinecke and Collet, 1998). Because, insulin, IGF-I and II have been derived from a common ancestral molecule through a series of gene duplications and point mutations.

Insulin like Growth Factors (IGFs) are normally expressed in various tissues and reproductive organs (Table 1) of many species (Carlsson *et al.*, 1993; Spicer, 1995; Daliri *et al.*, 1999; Kowalik *et al.*, 1999; Watson *et al.*, 1999; Wang *et al.*, 2000; Jin and Wang, 2001; Pushpakumara *et al.*, 2002; Hastie *et al.*, 2004; Daftary and Gore, 2005; Hastie and Haresign, 2008; Zakaria *et al.*, 2009). Also, oviductal and endometrial secretions contain IGF-I and IGF-II in many species including pig, cow and sheep (Letche *et al.*, 1989; Simmen *et al.*, 1989; Geisert *et al.*, 1991; Ko *et al.*, 1991; Wiseman *et al.*, 1992; Makarevich and Sirotkin, 1997).

Endocrine IGF-I has been associated with several reproductive traits, such as age at first calving (Yilmaz *et al.*, 2006; Brickell *et al.*, 2007), conception rate to first service (Patton *et al.*, 2007), twin ovulations (Echternkamp *et al.*, 2004), and pre-implantation embryo

development (Velazquez *et al.*, 2005). Insulin-like growth factor-II is essential for normal placental and fetal growth, but not post-natal growth (DeChiara *et al.*, 1990). Excess IGF-II is, however, detrimental to the embryo or fetus as it can lead to over-stimulation of IGF-1R, resulting in somatic overgrowth, cardiac and skeletal abnormalities and perinatal death (Lau *et al.*, 1994).

Increase in the number of live newborn is dependant upon the optimal uterine environment concerted with of optimal physiology mediated by hormones and growth factors. The aim of this review is to outline the impact of insulin and insulin like growth factors on embryo growth and development.

Insulin and its role in embryonic development: Insulin is composed of 51 amino acids. Its amino acid sequence varies among species, but certain segments of the molecule are highly conserved, including the positions of the three disulfide bonds, at the both ends of the A chain and at the C-terminal residues of the B chain.

Insulin signals through the receptor embedded in the plasma membrane. Its presence in embryos of mouse, rat, cow and sheep have been reported (Harvey *et al.*, 1995; Kaye and Harvey, 1995) and the stimulatory effect of insulin on cow (Zhang *et al.*, 1991), rat (Zhang and Armstrong, 1990) and pig (Lewis *et al.*, 1992) embryos have been mentioned. Dose response studies provided evidence that the insulin receptor mediated the stimulatory actions of insulin. Insulin receptor is composed of two extracellular α -subunits and two transmembrane β -subunits linked by disulfide bonds. Binding of insulin to the α -subunit induces a conformational change resulting in the autophosphorylation of a number of tyrosine residues present in the β -subunit (Van Obberghen *et al.*, 2001). Insulin receptor functions just like an enzyme transferring the phosphate groups from ATP to tyrosine residues on intracellular proteins which in turn alters their activity, thereby generating a biological response (Lizcano and Alessi, 2002).

Table 1: Insulin-like growth factor ligand and receptor expression in reproductive organs of bovine, ovine and swine

Ligands and receptors	Bovine			Ovine			Swine		
	Ovary	Oviduct	Uterus	Ovary	Oviduct	Uterus	Ovary	Oviduct	Uterus
IGF-I	+	+	+	+	+	+	+	+	+
IGF-II	+	+	+	+	+	+	+	+	+
IGF-IR	+	+	+	+	+	+	+	+	+
IGF-IIR	-	-	-	+	-	-	+	-	-

+: Indicates the presence of expression, references in text above; -: Indicates that the data is not present, references in text above

Insulin promotes growth of pre-implantation embryos in several species (Susa *et al.*, 1984; Kane *et al.*, 1997). Insulin response arose during compaction (Harvey and Kaye, 1988) and persisted in both trophoctoderm and inner cell mass lineages of the blastocyst (Harvey and Kaye, 1991). The number of cells within bovine (Zhang *et al.*, 1991) and murine (Gardner and Kaye, 1991) blastocysts is increased by culturing in medium supplemented with insulin. However, insulin did not stimulate cell proliferation in trophoctoderm, as it did in the inner cell mass (Harvey and Kaye, 1991). The resistance of trophoctoderm proliferation to insulin suggests that the receptors in these cells are coupled to a different set of signaling pathways from those in the inner cell mass. Apical and basolateral membranes of trophoctoderm cells and the inner cell mass membranes seemed to express both insulin and IGF-I receptors. Therefore, it is controversial whether insulin is capable of inducing mitogenic effects through its own receptor, or whether the growth-promoting effects of insulin result from its weak interaction with the IGF-I receptor or occur within insulin/IGF-I receptor hybrids (Sweet *et al.*, 1987; Soos *et al.*, 1993).

Insulin-like growth factors and their role in embryo development: Insulin and IGF-I, II act endocrine as well as paracrine/autocrine manner mediated by, two type of receptors (IGF-IR, IGF-IIR) and six IGF binding proteins (IGFBPs) (Jones and Clemmons, 1995; Pavelic *et al.*, 2007). Mature IGF-I and IGF-II consist of A, B, C, and D-domains. The A- and B-domains of IGFs are homologous to those of insulin. Unlike in the case of insulin, the C-domain is not cleaved off in mature IGFs. Insulin-like growth factors contain an additional D-domain, which is not present in insulin (Le-Roith *et al.*, 2001).

The IGF peptides (IGF-I and -II) mediate their effects through IGF-IR receptor, which is structurally related to the insulin receptor and binds both IGF-I and IGF-II with high affinity and insulin with lower affinity. Insulin-like growth factors-IIR receptor, also known as the mannose-6-phosphate receptor, binds IGF-II with high affinity but will not bind IGF-I or insulin (Fig. 1).

The expression of IGF-I, II and their receptor proteins (IGF-1R and IGF-IIR) have been detected in ovary, oviducts and uterus of bovine (Table 1) (Daliri *et al.*, 1999; Armstrong *et al.*, 2000; Llewellyn *et al.*, 2007; Sudo *et al.*, 2007; Winger *et al.*, 1997; Fenwick

et al., 2008a, b), ovine (Teissier *et al.*, 1994; Stevenson and Wathes, 1996; Taylor *et al.*, 2001; Hastie *et al.*, 2004; van Lier *et al.*, 2006) and swine (Simmen *et al.*, 1992; Liu *et al.*, 2000).

The presence of IGF ligands and receptors in pre-implantation embryos from the different species, including sheep, have also been mentioned (Watson *et al.*, 1994, 1999; Lonergan *et al.*, 2000; Lighten *et al.*, 1997; Wang *et al.*, 2009).

It has been reported that, Addition of IGF-I, IGF-II to the culture of bovine embryos significantly accelerated embryonic development, especially the change from the expanded blastocyst to hatched blastocyst stages (Neira *et al.*, 2010). In bovine, addition of IGF-I and II to the *in vitro* maturation or culture media increased embryo development (Bonilla *et al.*, 2011). The similar result obtained from rabbit, When rabbit embryos incubated with IGF-I progressed to the blastocyst stage (Herrler and Beier, 1994). In mouse embryo culture, when IGF-II expression was reduced by the presence of antisense IGF-II oligonucleotides, the rate of embryo development was inhibited. This effect was abolished by the addition of IGF-II to the culture medium (Rappolee *et al.*, 1992).

Also, insulin-like growth factors are important for the development and functional maturation of the Central Nervous System (CNS), skeletal tissues, and reproductive organs. In mice, knock out of the IGF-I gene causes infertility (Baker *et al.*, 1996), underdevelopment of muscle tissue (Powellbraxton *et al.*, 1993), significant decrease in auditory neuron number and increase in apoptosis of cochlear neurons (Camarero *et al.*, 2001).

Insulin-like growth factor-I (GF-1) and its impact on embryo development: Insulin-like growth factor-I (GF-I) is a single-chain polypeptide with a molecular mass of 7649 Da and shares 43% amino acid sequence homology with insulin. This peptide is produced in organs of reproductive significance such as hypothalamus, ovaries, oviducts, and uterus (Spicer, 1995; Watson *et al.*, 1999; Daftary and Gore, 2005). The IGF-I exerts its effect by binding to high-affinity membrane receptors. The IGF-I receptor (IGF-IR) binds to IGF-I with the highest affinity and there is a 60% homology between IGF-1R and the insulin receptor (DeMeyts and Whittaker, 2002). The IGF-IR has two α -subunits and two β -subunits linked by disulfide bonds (Fig. 1). The α -chains are located extracellularly while the β -subunit spans the membrane

and is responsible for intracellular signal transduction upon ligand stimulation. Both of the subunits (α and β) are synthesized from a single mRNA precursor. The α -subunit contains a cysteine-rich ligand-binding site. The precursor is glycosylated, proteolytically cleaved, and cross-linked by cysteine bonds to form a functional transmembrane $\alpha\beta$ -chain. The β -subunit has tyrosine kinase activity. Insulin and IGF-1 can cross-activate the receptors, due to the exhibition of high sequence and structural similarity, when they added together at high concentrations in cell culture studies. Recent genetic studies in *Xenopus* and zebrafish suggest that the structure and function of the IGF-IR is evolutionarily conserved (Pera *et al.*, 2001; Richard-Parpaillon *et al.*, 2002; Eivers *et al.*, 2004; Schlueter *et al.*, 2006, 2007). Using antisense Morpholino Oligonucleotide (MO)-based target gene knockdown approach and by specific inhibiting IGF-1R-mediated signaling using a dominant-negative IGF-1R fusion protein, Schlueter *et al.* (2007) have shown that IGF-1Rs in zebra fish are required for embryo viability and proper growth.

Most of the IGF-I measured in blood is produced by the liver (Pfaffl *et al.*, 1998; Yakar *et al.*, 1999; Fenwick *et al.*, 2008a, b). Endocrine IGF-I has been associated with several reproductive traits, such as age at first calving (Liu *et al.*, 1993; Baker *et al.*, 1996; Zhou *et al.*, 1997; Kadakia *et al.*, 2001; Yilmaz *et al.*, 2006; Brickell *et al.*, 2007), conception rate to first service (Patton *et al.*, 2007), twin ovulations (Echternkamp *et al.*, 2004) and preimplantation embryo development (Velazquez *et al.*, 2005).

Locally produced IGF-I in ovary has various actions including enhancement of cell proliferation, aromatase activity, and progesterone biosynthesis (Adashi *et al.*, 1985; Kamada *et al.*, 1992; Savion *et al.*, 1981). Supplementation of bovine embryo culture with IGF1 has been reported to increase rate of blastocyst development, embryo survival as well as reducing the effects of heat shock on embryos (Bonilla *et al.*, 2011).

Insulin-like growth factor-II (IGF-II) and embryo development: Insulin-like growth factor-II (IGF-II) is a single chain polypeptide. Mature IGF-II is a 7.5 kDa protein (Liu *et al.*, 1993). It binds to the IGF-IR and with the highest affinity to type IGF-II receptors (IGF-IIR). IGF-IIR is a polypeptide hormone and sharing approximately 70% sequence identity with IGF-IR. But, IGF-IIR is also acts as a mannose-6-phosphate (M6P) receptor. Therefore, it is structurally and functionally distinct from the IGF-IR. The IGF-IIR/ M6P are a monomeric transmembrane protein with an extracellular domain composed of 15 cysteine-rich repeats. Mammalian IGF-IIR/ M6P has about 100 times higher affinity for IGF-II than IGF-I. Knockout of the IGF-IIR gene or loss of the imprinted IGFIIR results in fetal

overgrowth and perinatal lethality (Lau *et al.*, 1994; Wylie *et al.*, 2003).

Insulin-like growth factor-II is expressed in ovary, oviduct and uterus of human, bovine and sheep (Watson *et al.*, 1999; Wang *et al.*, 2000; Jin and Wang, 2001; Hastie *et al.*, 2004; Hastie and Haresign, 2008; Zakaria *et al.*, 2009). In mice, use of homologous recombination technology proved conclusively that IGF-II was required for normal embryonic development, as IGF-II null mice were 60% smaller than their wild-type littermates (DeChiara *et al.*, 1990, 1991). IGF-II has been reported to be essential for normal placental and fetal growth, but not post-natal growth (DeChiara *et al.*, 1990). Excess IGF-II is, however, detrimental to the embryo or fetus as it can lead to over-stimulation of IGF-1R, resulting in somatic overgrowth, cardiac and skeletal abnormalities and perinatal death (Lau *et al.*, 1994).

The effect of IGF-binding proteins and their functional role: There are a family of secreted proteins, named as Insulin-like Growth Factors Binding Proteins (IGFBPs) that specifically bind IGF-I and IGF-II with affinities that are equal to or greater than those of the IGF-IR (Fig. 1). There are six well characterized mammalian IGFBPs, designated IGFBP-1 to -6 (Allan *et al.*, 2001). Most IGFBPs, including IGFBP-2 to -6, are expressed in peripheral tissues and most mammalian cells express more than one form of IGFBPs (Duan and Xu, 2005; Firth and Baxter, 2002; Jogie-Brahim *et al.*, 2009; Jones and Clemmons, 1995; Yamada and Lee, 2009). Insulin-like growth factors binding proteins function as carrier proteins in the circulation and regulate IGF turnover, transport, and half-life of circulating IGFs (Jones and Clemmons, 1995). The affinity of IGFBPs for IGFs is controlled by phosphorylation, glycosylation and specific proteolysis (Clemmons, 1998). The IGF/IGFBP complexes help to prevent potential hypoglycemic effect of circulating IGFs by preventing possible cross-binding of IGFs with the insulin receptor (Rajaram *et al.*, 1997). In addition to functioning as carrier proteins, IGFBPs also have their own receptors mediating IGF-independent actions. Cell surface receptors for IGFBP-1 (Jones *et al.*, 1993), IGFBP-2 (Rauschnabel *et al.*, 1999), IGFBP-3 (Oh *et al.*, 1993), IGFBP-5 (Andress, 1995, 1998), IGFBP-6 (Bach *et al.*, 1992) and a low-affinity IGFBP, namely IGFBP-7 and 8 bearing structural homology to the classic IGFBPs was recently described (Oh *et al.*, 1996; Kim *et al.*, 1997) although, to date, none of these proteins have been cloned.

Insulin-like growth factor binding proteins shares a common domain structure arrangement. They all have a highly conserved N-terminal domain (N-domain) and C-terminal domain (C-domain), and a variable central domain (L-domain). The N- and C-domain contain multiple conserved cysteine residues, which form intra-

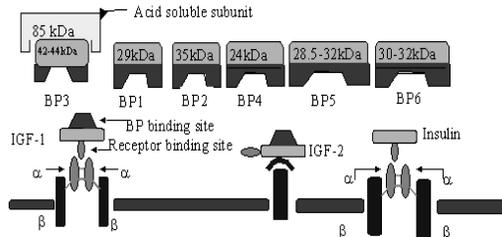


Fig. 1: The IGF system consist of three different parts, two ligands (IGF-I, IGF-II), which are structurally related to insulin, two receptors and six binding proteins. Insulin like Growth Factor-I and II are complexed with the family of binding proteins (IGFBP's) in serum, amniotic and other body fluids. These binding proteins modulate the autocrine and paracrine action of the ligands. They inhibit the mitogenic effect of IGFs (IGF1 and IGF-2) by limiting peptide access to specific cell surface receptors (Modified from Fig. 1 of Monget and Monniaux, 1995).

domain disulfide bonds within the N-domain and C-domain, thereby defining their overall globular structure.

The highly variable L-domain is considered as a flexible linker region connecting the N- and C- domain (Chelius *et al.*, 2001; Forbes *et al.*, 1998; Neumann and Bach, 1999). The N-domain contains the high affinity IGF-binding site, but the C-domain also contributes to IGF binding to some degree (Brinkman *et al.*, 1991; Clemmons, 2001; Hobba *et al.*, 1998; Zeslawski *et al.*, 2001). The C-domain of an IGFBP often mediates its interactions with other proteins. For instance, both IGFBP-3 and IGFBP-5 bind to the acid-labile subunit (ALS) through their C-domains (Firth and Baxter, 2002; Guler *et al.*, 1987). This ternary complex (IGF-IGFBP-ALS) greatly increases the half-life of IGFs in circulation.

The central L-domain is the least conserved and often contains sites for post-translational regulations (Clemmons, 2001; Firth and Baxter, 2002). The post-translational regulations (such as phosphorylation, glycosylation and proteolysis) influence the affinity of IGFBPs to IGFs and regulate IGF availability.

Insulin-like binding protein-1 (IGFBP-1) is a 25-34 kDa protein and isolated in two forms. Placental form was originally isolated from human placenta as protein 12 and it is growth hormone independent. Amniotic form is 28 kDa protein isolated from human amniotic fluid. Both forms have the same amino acid sequence at the N-terminal (Koistinen *et al.*, 1986). Amniotic form of IGFBP-1 binds to both IGF-I and IGF-II with high affinity. Using the zebra fish model, it was hypothesised that elevated IGFBP-1 mediates hypoxia-induced embryonic growth retardation and developmental delay by binding to and inhibiting the activities of IGFs using loss-and gain of function approaches. Knockdown of IGFBP-1 using antisense oligonucleotides (MOs) significantly alleviated the hypoxia-induced growth retardation and

developmental delay. Over-expression of IGFBP-1 caused growth and developmental retardation under normoxia. Furthermore, re-introduction of IGFBP-1 to the IGFBP-1 knocked down embryos restored the hypoxic effects on embryonic growth and development (Kajimura *et al.*, 2005).

Insulin-like binding protein-2 (IGFBP-2) is a 32-34 kDa protein present in fetal tissue, serum lymph, amniotic fluid, follicular fluid and cerebrospinal fluid (Mondschein *et al.*, 1990; Schoen *et al.*, 1992). IGFBP-2 preferently binds to IGF-II.

Insulin-like binding protein-3 (IGFBP-3) is a 53 kDa protein binds to IGF-I and II with high affinity. It can function either as inhibitor or activator of IGF-I stimulated DNA synthesis. Molecular weight of IGFBP-3 depends on glycosylation degree (Russell and Van Wyk, 1995). Glycosylated IGFBP-3 is able to bind to cell surface by a weak non-covalent sugar-sugar interaction. Free IGFBP-3 has 3 to 10-fold higher affinity to ligand than cell surface-associated IGFBP-3 (McCusker and Clemmons, 1992). The concentration of IGFBP-3, in blood, is 40-fold higher than IGFBP-1 and also has higher affinity to IGF-I. Therefore, the majority of IGF-I, in circulation, is bound to IGFBP-3.

Insulin-like binding protein-4 (IGFBP-4) was isolated in two forms with different molecular weight (29 kDa and 24 kDa) from ovine blood plasma. It blocks the effect of exogenously added IGF-I to the cells (Russell and Van Wyk, 1995). Although a large number of *in vitro* studies have shown IGFBP-4 to be an inhibitory IGFBP, knockout of the IGFBP-4 gene in mice reduces, rather than increases, prenatal growth (Ning *et al.*, 2008).

Insulin-like binding protein-5 is a 23 kDa protein and originally purified from human osteoblast derived culture. It is present in endocrine tissues.

Insulin-like binding protein-6 is a 23 kDa protein was originally isolated from human cerebrospinal fluid). Relatively little is currently known about its role and regulation. IGFBP-6 is O-glycosylated and its glycosylated form exhibits much greater resistance to proteolysis than its nonglycosylated counterpart (Neumann *et al.*, 1998). IGFBP-6 is distinct in its preferential affinity for IGF-II relative to IGF-I, which, unlike any other IGFBP, is 20- to 100-fold higher (Neumann and Bach, 1999). IGFBP-6 functions to inhibit the actions of IGFII, with inhibition thought to result from the formation of high-affinity IGF-IGFBP complexes that prevent IGF-II from binding to the IGF receptors (Neumann and Bach, 1999). Several hormones, including a number of growth factors, have been shown to regulate IGFBP-6, but the manner in which they act appears to be very cell-specific, and the mechanisms at this time are not fully understood (Neumann and Bach, 1999). Insulin-like binding protein-7 together with IGFBP-8 show a low

affinity to IGFs. IGFBP-8 is probably connective tissue growth factor (Kim *et al.*, 1997).

CONCLUSION

Insulin-like growth factors and their receptor are expressed in embryos and in reproductive organs of farm animal. Although there are many published reports concerning importance of IGF system and their receptors in blastocyst formation, implantation and embryo growth, there is not enough information about the exact physiologic regulation and function of some components of the IGFs system (such as GFbps). The regulation of IGF system is complicated.

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