

# Pregnancy-specific glycoproteins: complex gene families regulating maternal-fetal interactions

TOM MOORE<sup>1</sup> and GABRIELA S. DVEKSLER<sup>\*,2</sup>

<sup>1</sup>School of Biochemistry and Cell Biology, Western Gateway Building, University College Cork, Cork, Ireland and

<sup>2</sup>Department of Pathology, Uniformed Services University of the Health Sciences, Bethesda, Maryland, USA

**ABSTRACT** The pregnancy-specific glycoproteins (PSGs) are the most abundant trophoblastic proteins in maternal blood during human pregnancy and they appear to be exclusive to species with hemochorial placentation. There are ten protein-coding human *PSG* genes (*PSG1 - PSG9, PSG11*) and also multiple *PSG* genes in non-human primates, rodents and bats. Several studies indicate that PSGs have immunoregulatory, pro-angiogenic, and anti-platelet functions. Some PSGs have been shown to bind different moieties on the surface of cells, including the tetraspanin CD9, heparan sulphate, and specific integrins. Recently, *PSG1* was shown to associate with and activate the anti-inflammatory cytokines transforming growth factor (TGF)- $\beta$ 1 and TGF- $\beta$ 2 making *PSG1* one of the few known biological activators of these important cytokines. TGF- $\beta$ s regulate many biological processes essential for pregnancy success including trophoblast invasion and proliferation, angiogenesis, extracellular matrix formation and tolerance to the fetal semi-allograft. As summarized in this review, progress has been made in recent years towards a better understanding of the functions of these proteins which were originally described in the early 1970s, but more research will likely contribute to demonstrate their importance for a successful pregnancy.

**KEY WORDS:** *trophoblast, placental hormone, immunoregulation, TGF $\beta$ , integrin*

## Introduction

The Pregnancy-specific glycoprotein (*PSG*) and the closely related Carcinoembryonic antigen cell adhesion molecule (*CEACAM*) gene families are members of the immunoglobulin (*Ig*) superfamily (Kammerer & Zimmermann, 2010). The *CEACAM/PSG* ancestral gene is thought to be common to both primates and rodents, but subsequent gene duplications led to considerable diversification of protein structure, expression and function (Kammerer & Zimmermann, 2010, Rudert *et al.*, 1992). The *CEACAMs* are predominately cell membrane-anchored proteins whereas all *PSGs* are secreted (Kammerer & Zimmermann, 2010). However, *CEACAMs* and *PSGs* share several structural features. Both *CEACAMs* and *PSGs* are encoded by multigene families and both families of proteins have an amino terminal *Ig* variable-like domain and a variable number of *Ig* constant-like domains (Kammerer & Zimmermann, 2010, McLellan *et al.*, 2005a, Rudert *et al.*, 1992). There are twelve human and fifteen mouse *CEACAM* genes that are widely expressed in normal and cancerous tissues (Hammarstrom, 1999, Kuespert *et al.*, 2006, Zebhauser *et al.*, 2005). Structural and functional analyses indicate that *CEACAM* extracellular domains are involved in homotypic and

heterotypic adhesion, and the cytoplasmic domains are involved in signal transduction (Huang *et al.*, 2013, Kammerer & Zimmermann, 2010, Obrink, 1997). By virtue of their expression on epithelia and immune cells, *CEACAM* family members act as pathogen receptors, which may be a significant influence on *CEACAM* family evolution (Chang *et al.*, 2013, Kammerer & Zimmermann, 2010). There are ten and seventeen *PSG* protein-coding genes in human and mouse, respectively (Kammerer & Zimmermann, 2010, McLellan *et al.*, 2005a). The phylogenetic distribution and expression of *PSGs* is more restricted than *CEACAMs*, and the evolutionary selective pressures driving *PSG* family expansion and diversification therefore may be different to *CEACAMs* (Ball *et al.*, 2004, McLellan *et al.*, 2005a, Wynne *et al.*, 2006, Zhou *et al.*, 1997).

## PSG locus, gene and protein structure

The *PSG* genes are clustered at chromosome 19q13 in the human and on proximal chromosome 7 in the mouse. The ten protein-

*Abbreviations used in this paper:* *CEACAM*, carcinoembryonic antigen cell adhesion molecule; *PSG*, pregnancy-specific glycoprotein; *TGF*, transforming growth factor.

\*Address correspondence to: Gabriela S. Dveksler, Department of Pathology, Uniformed Services University of the Health Sciences, Bethesda, Maryland, USA.  
e-mail: gabriela.dveksler@usuhs.edu

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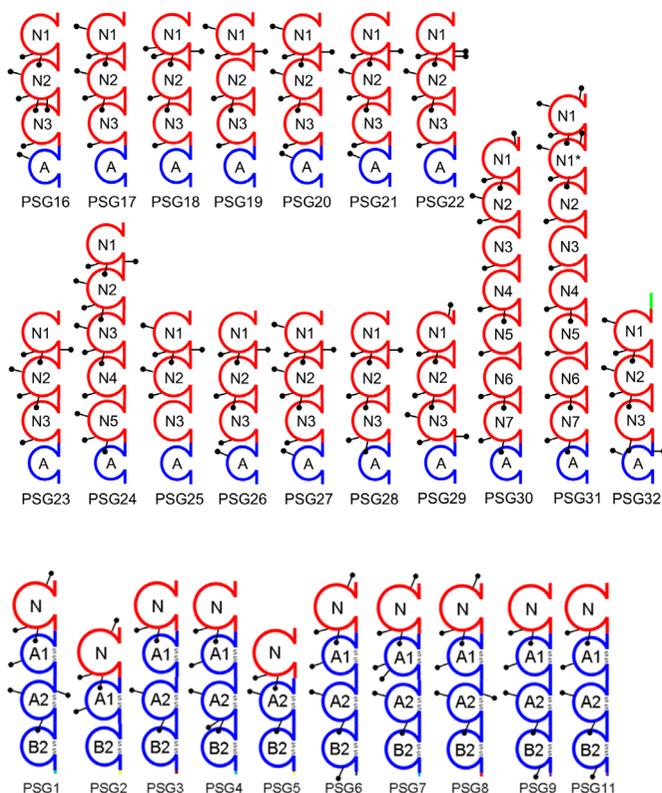
coding human *PSG* genes, *PSG1* - *PSG9*, and *PSG11* (*PSG10* is a non-coding pseudogene), are arrayed within a genomic region of 0.55 Mb (www.ensembl.org, Teglund *et al.*, 1994). The mouse locus is approximately 1.74 Mb and contains six *Psg* genes in the A2 chromosome band and eleven genes in the A3 band, the latter flanked by the unrelated *Mill1* and *Mill2* genes; we note that the gene order in the *Psg* locus described in McLellan *et al.*, (2005a) is inaccurate and the correct map is available at www.ensembl.org. Human and mouse loci contain a variety of *CEACAM* and *PSG* pseudogenes and non-coding RNAs (Kammerer & Zimmermann, 2010, www.ensembl.org). Orthologous relationships cannot be determined between human and mouse *PSG* genes indicating that these gene clusters have evolved independently since the common ancestor (McLellan *et al.*, 2005a), although orthologous relations can be determined for a subset of mouse and rat *PSGs* (T. Moore & colleagues, unpublished). The human *PSG* genes appear to exhibit a high level of copy number variations, rapid coding sequence divergence, and a relatively high frequency of gene conversion between different *PSG* family members compared to the genome average (Chang *et al.*, 2013, Dumont & Eichler, 2013).

The domain organization of the human and mouse *PSG* proteins exhibit pronounced divergence (see Fig. 1). Human *PSGs* contain one Ig variable-like domain (N) and, usually, three Ig constant-like domains (Ig C2-like domains of type A or B), and a relatively hy-

drophilic tail (www.carcinoembryonic-antigen.de, McLellan *et al.*, 2005a). In addition, there are a number of variations, including splice variants, particularly at the carboxy terminus. However, many of these are poorly characterised and may reflect cloning and annotation of rare, functionally irrelevant variants. In contrast, rodent *PSGs* typically have three or more Ig variable-like (N) domains followed by a single Ig constant-like (A) domain (McLellan *et al.*, 2005). All eight rat *PSGs*, with the exception of *PSG36* (N1-N2-N3-N4-N5-A), are of the N1-N2-N3-A domain arrangement (McLellan *et al.*, 2005b). The mouse *Psg* protein family has seventeen members, fourteen with three N-domains and a single A-domain (N1-N2-N3-A) arrangement, and three - *Psg24*, *Psg30* and *Psg31*, which have an expanded structure created by domain duplications, as follows: *Psg24* (N1-N2-N3-N4-N5-A), *Psg30* (N1-N2-N3-N4-N5-N6-N7-A) and *Psg31* which has a recently duplicated N1 domain and therefore a (N1-N1-N2-N3-N4-N5-N6-N7-A) domain arrangement (McLellan *et al.*, 2005a). However, notwithstanding these structural differences, there is evidence that human *PSG* N-domains and mouse N1-domains have conserved structures as evidenced by branching of the mouse N1-domains with human N-domain, in preference to mouse N2 and N3-domains, in a phylogenetic tree (McLellan *et al.*, 2005b). This is consistent with the apparently conserved functions of human and mouse *PSG* proteins in immune and vascular regulation (see below). The common ancestor of human and mouse *CEACAMs* and *PSGs* was most likely similar to *CEACAM1*, the only family member with a homologous gene structure in the human, rat and mouse that encodes all types of extracellular domains present in *CEACAM* and *PSG* proteins (McLellan *et al.*, 2005a, Rudert *et al.*, 1992, Teglund *et al.*, 1995).

## PSG expression pattern

*PSGs* may be the most abundant trophoblastic proteins in maternal blood during human pregnancy. The maternal serum level increases as gestation progresses with concentrations of up to 200-400 µg/ml reported at term, far exceeding levels of other well-known placental hormones such as human chorionic gonadotropin (Lee *et al.*, 1979, Lin *et al.*, 1974, Towler *et al.*, 1976). Mouse *Psg* proteins do not appear to exhibit similarly high levels of expression (J. Baensinger, Pers. Commun.). In the first half of mouse gestation, *Psg* mRNA expression is comprised almost entirely of *Psg22* expressed in trophoblast giant cells (Blois *et al.*, 2012, Wynne *et al.*, 2006); however, a subset of *Psg* mRNAs are highly expressed in spongiotrophoblast during the second half of mouse gestation, with *Psg16*, *Psg21* and *Psg23* mRNAs being particularly abundant (Ball *et al.*, 2004, Kromer *et al.*, 1996, Wynne *et al.*, 2006). Given the large number of *PSG* genes in the human, and the evidence of rapid evolution at this locus (Chang *et al.*, 2013, Dumont *et al.*, 2013), the *PSG* expression pattern appears remarkably constant in individual samples: in a series of placental samples from two male and two female pregnancies, *PSG1* and *PSG3* comprised the bulk of expression at the end of the first trimester, whereas *PSG1*, *PSG3*, *PSG4*, *PSG5*, and *PSG6* were approximately equally expressed at term, with relatively low expression of the other *PSGs* (Shanley *et al.*, 2013). Human *PSG* transcripts and proteins increase in trophoblast cells undergoing differentiation (Aronow *et al.*, 2001, Camolotto *et al.*, 2010). However, reports that *PSGs* are secreted by cultured human preimplantation embryos must be questioned given the apparent absence of *PSG*



**Fig. 1. Structures of mouse (PSG16 - PSG32) and human (PSG1 - PSG9, PSG11) PSG proteins.** (Adapted from McLellan *et al.*, 2005, and <http://www.carcinoembryonic-antigen.de/index.html>). Red color-coded 'N' domains are Ig variable-like domains, and blue 'A' and 'B' domains are Ig constant-like domains. Black 'lollipops' indicate potential N-linked glycosylation sites. Note that atypical amino and carboxy terminal sequences are also color-coded.

transcripts in recent RNA-seq datasets from preimplantation-stage human embryos (Dimitriadou *et al.*, 1992, T. Moore & K. Niakan, unpublished observations). We suggest that the antibodies used in some early studies of PSG expression may not have been specific to PSGs and that expression commences with the differentiation of syncytiotrophoblast as suggested by studies involving *in vitro* differentiation of cytotrophoblasts and the JEG-3 cell line (Aronow *et al.*, 2001, Camolotto *et al.*, 2010).

There is some evidence for PSG expression in non-trophoblastic tissues; however, much of the older literature may over-estimate the extent of PSG expression in adult tissues due to technical issues such as use of poorly validated antibodies, or reporting of rare or aberrant transcripts from cDNA libraries.

Conversely, a more recent study reported up-regulation of PSG9 expression in colorectal carcinogenesis (Salahshor *et al.*, 2005), and using a panel of novel well characterized monoclonal antibodies to PSG1, PSG expression was found in intestinal epithelium (T. Moore & colleagues, unpublished observations).

Mouse Psg17, Psg18 and Psg19 mRNAs were found using RT-PCR in the placenta and in pooled tissues of embryonic, but not in adult, tissues (Kromer *et al.*, 1996). However, Psg18 mRNA and protein were subsequently found in the follicle-associated epithelium overlaying intestinal Peyer's patches in the mouse (Kawano *et al.*, 2007), and a brain-specific transcript of *Psg16* has been reported (Phillips *et al.*, 2012), suggesting that some PSGs may have evolved specific functions in adult tissues.

### PSG gene transcriptional regulation

Relatively little is known about regulation of PSG expression. The region upstream of the *PSG5* gene transcriptional start site has been characterized and, given the similarity to the paralogous region of other *PSG* genes, these studies may be applicable to understanding regulation of the entire family (Blanchon *et al.*, 2006, Chamberlin *et al.*, 1994, Nores *et al.*, 2004). The human *PSG* gene promoters lack an obvious TATA box, typical initiator elements, or GC-rich regions (Frangmyr *et al.*, 2000, Panzetta-Dutari *et al.*, 1992). *PSG5* expression depends on a ubiquitous specificity protein 1 (Sp1) binding site located in the minimal core promoter region of all human *PSGs*. This SP1 site activates *PSG5* promoter constructs, and SP1 is co-expressed with *PSGs* in the syncytiotrophoblast (Nores *et al.*, 2004). Using *PSG5* promoter-reporter transfections and ChIP assays, Kruppel-like factor 4 (KLF4) was shown to be an activator of the *PSG5* promoter by binding to a consensus site in the core promoter (Blanchon *et al.*, 2006). Kruppel-like factor 6 (KLF6) was shown to activate *PSG3* and *PSG5* gene promoters in JEG-3 cells, further supporting an important role for the KLF family of transcription factors in *PSG* gene regulation (Racca *et al.*, 2011). Additionally, a retinoic acid response element (RARE) /CACCC box composite element is highly conserved in the core promoters of all human *PSG* genes. Consistent with a role for this element in *PSG* expression, it was shown that RXR $\alpha$  binds the *PSG5* core promoter and 9-cis retinoic acid (RA) induces *PSG5* expression in JEG-3 cells (Lopez-Diaz *et al.*, 2007). Similarly, a RARE site in the *PSG3* promoter was shown to be required for basal promoter activity using reporter assays (Camolotto *et al.*, 2010). These observations are consistent with the known roles for retinoic acid (RA) derivatives and their receptors (RAR and RXR) in placental development (Sapin *et al.*, 1997, Yan *et al.*, 2001).

The *PSG3* promoter also contains a putative binding site for the Ets family transcription factor GA-binding protein (GABP). This binding site was shown to be involved in *PSG3* gene activation during trophoblast differentiation (Camolotto *et al.*, 2010). Another Ets family member (PEA3) is similarly implicated in expression of multiple human *PSGs* (Chamberlin *et al.*, 1994).

It has been suggested that *PSG5*, and presumably all *PSGs* by extrapolation, may be regulated by release from transcriptional repression. The *PSG5* core promoter drives transcription in a reporter assay in a variety of cell types and tissue-specific expression may be mediated by repressive factors binding further upstream from the transcriptional start site (Panzetta-Dutari *et al.*, 2000). Interestingly, induction of replicative senescence in HeLa cells by treatment with 5-bromodeoxyuridine resulted in upregulation of all *PSGs*; likewise, in normal human fibroblasts undergoing replicative senescence (Minagawa *et al.*, 2005, Endoh *et al.*, 2009). We suggest that replicative senescence may mimic aspects of terminally differentiated trophoblast cell lineages such as the syncytiotrophoblast thereby resulting in *PSG* expression.

The transcriptional regulation of mouse *Psg* genes is virtually unexplored. However, both *Ceacam* and *Psg* gene expression is greatly attenuated in the placentas of Inositol requiring enzyme-1a (IRE1a) and X-box binding protein 1 (XBP1) knockout mice, which are implicated in endoplasmic reticulum (ER) stress responses and in placental development (Oikawa *et al.*, 2010). Treatment of SM10 trophoblast cells with thapsigargin, an ER stressor that activates the IRE1a-XBP1 pathway, or overexpression of wild type IRE1a or XBP1, upregulated both *Psg18* and *Psg28*. Use of *Psg28* promoter deletion constructs identified two regions whose deletion reduced this response, but specific binding sites for XBP1 binding sites were not identified (Oikawa *et al.*, 2010).

### PSG receptors

While the cellular receptors for all members of the *PSG* family have yet to be identified, some progress has been made in the identification of the moieties that bind *PSGs* on the cell surface for some murine *PSGs* and for human *PSG1*. Murine *PSG17* and *PSG19* bind to CD9 (Ha *et al.*, 2008, Waterhouse *et al.*, 2002). CD9 is a member of the tetraspanin family, which consists of four transmembrane domains delimiting two extracellular loops or domains (Charrin *et al.*, 2009). Over thirty different tetraspanins have been identified in humans and mouse. So far, the only interaction between the *PSG* and tetraspanin families that has been described is that of murine *PSG17* and *PSG19* with the extracellular domain 2 of CD9 (Ellerman *et al.*, 2003). At this time, we have found no evidence for an interaction between mouse *PSG22* and *PSG23* or human *PSG1* with CD9 but whether other members of the *PSG* family interact with CD9 or other tetraspanins remains to be explored. Expression of CD9 in macrophages was required for the ability of a protein consisting of the N-terminal domain of *PSG17* to induce the secretion of cytokines by these cells, indicating that the *PSG17*-CD9 interaction has physiological consequences (Ha *et al.*, 2005).

Recently, human *PSG1* and murine *PSG17*, *PSG22* and *PSG23* were shown to interact with heparan sulfate (Blois *et al.*, 2012, Lisboa *et al.*, 2011, Sulkowski *et al.*, 2011). These interactions were demonstrated in solid phase assays, by affinity chromatography and in FACS analysis using Chinese hamster ovary (CHO) cells

lacking surface expression of the glycosaminoglycans (GAGs) heparin or chondroitin sulfate. In addition, these PSGs bound to the pre-B cell line Namalwa only when the cell line was transfected with syndecans or glypican-1, cell surface proteins which contain GAG chains. (Lisboa *et al.*, 2011). Recently, we also determined that human PSG9 binds to syndecan-transfected Namalwa cells but not to the parental cell line, indicating that like PSG1 and murine PSGs-17, 22 and 23, PSG9 binds to GAGs (Dveksler, unpublished data).

Besides binding to heparan and chondroitin sulfate, recent studies indicate that PSG1 interacts with members of the integrin family. The tripeptide sequence Arg/Gly/Asp (RGD) found in the N domain of most human PSGs led to the suggestion that PSGs, similar to snake venom disintegrins, might bind integrins and disrupt cell - extracellular matrix interactions or modify other integrin-mediated functions, thereby facilitating trophoblastic invasion of maternal tissues (Rutherford *et al.*, 1995). The disintegrin hypothesis was supported by an evolutionary analysis suggesting conservation of RGD-like tri-peptides at the homologous position of mouse Psg N1 domains (McLellan *et al.*, 2005b).

Specifically, mouse Psg N1 domains do not possess an RGD tri-peptide motif, but do contain RGD-like motif sequences, which are not found in the N2 and N3 domains (Rudert *et al.*, 1992, McLellan *et al.*, 2005b). Additionally, PSG1 contains a KGD tripeptide motif overlapping, but one residue out of phase, with the 'RGD' position, suggesting an analogy to the snake venom disintegrin barbourin, which is a specific inhibitor of fibrinogen binding to the  $\alpha$ IIb $\beta$ 3 integrin on platelets mediated by a KGD motif (Scarborough *et al.*, 1991). All PSGs tested (PSG1, PSG9, and mouse Psg23) inhibited platelet - fibrinogen interactions suggesting an anti-thrombotic function for PSGs (Shanley *et al.*, 2013). PSG1 binding to integrin  $\alpha$ IIb $\beta$ 3 does not result in signaling indicating that the sole function of PSG binding appears to be inhibition of fibrinogen binding (Shanley *et al.*, 2013). However, the mode of binding of PSG1 to the integrin is complex and mutation of the KGD sequence or deletion of the entire N domain did not abolish this function, suggesting that more than one of the PSG1 protein domains can independently mediate binding (Shanley *et al.*, 2013).

Preliminary studies from the Dveksler laboratory indicate that  $\beta$ 1 integrin mediates the adhesion of endothelial cells and a trophoblast cell line (HTR-8) to PSG1-coated wells. Several members of the integrin family including  $\alpha$ 5 $\beta$ 1,  $\alpha$ v $\beta$ 3 and  $\alpha$ IIb $\beta$ 3 are involved in modulation of trophoblast migration, invasion and adhesion and also regulate the interaction of endovascular trophoblasts with endothelial cells (Harris *et al.*, 2009, Rout *et al.*, 2004). Further studies are required to identify the integrin  $\alpha$  chain, which together with the  $\beta$ 1 integrin subunit is involved in the adhesion to PSG1 and the physiological relevance of this interaction. Other human PSGs could potentially interact with integrins, which has been hypothesized based on the presence of an RGD tripeptide sequence in the N-domain of some human PSGs with the exception of PSG1, 4 and 8. At this time, however, there is no evidence for an interaction of the N-terminal domain of the RGD-containing PSGs with integrins. Furthermore, we have no evidence that the N-terminal domain is involved in the PSG1- $\beta$ 1 integrin interaction, which was observed with recombinant PSG1 composed of all 4 domains as well as the recombinant protein containing just 3 domains (N, A2, and B2).

## PSGs and angiogenesis

The success of pregnancy in mammals relies on the spatio-temporal coordination of several vascular processes at the fetal-maternal interface. During normal pregnancy, increased placental blood flow is critical for fetal growth and survival. This increased blood flow results from angiogenesis, vasodilatation, and vascular remodeling. Placental trophoblasts are key regulators of placental endothelial functions. We found that PSG1 and murine PSG22 and PSG23 induced endothelial tube formation on Matrigel and type I collagen. Enzymatic removal of GAGs from the surface of endothelial cells negated the ability of PSG1 to induce tube formation demonstrating the importance of the PSG1-heparan/chondroitin sulfate interaction. On the other hand, endothelial cells lacking the Vascular endothelial growth factor (VEGF) kinase insert domain receptor formed tubes in response to PSG1 treatment indicating that VEGF signaling is not required for the response to PSG1 (Lisboa *et al.*, 2011). While PSG22 and PSG23 also induce endothelial tubulogenesis and bind to GAGs, the requirement for the interaction of these PSGs with GAGs for tube formation has not been studied (Blois *et al.*, 2012 and G. Dveksler, unpublished data). In addition, whether PSG17 or other PSGs can induce the formation of tubes by endothelial cells remains unknown. VEGF-A is a key mediator in vasculogenesis and angiogenesis (Ferrara and Davis-Smyth, 1997). TGF- $\beta$ 1 has been shown to regulate the production of VEGF-A (Jeon *et al.*, 2007, Pertovaara *et al.*, 1994). We found that PSG1 treatment of human extravillous trophoblast cell lines and primary monocytes resulted in an increase of VEGF-A secretion (Ha *et al.*, 2010). The PSG1-mediated increase in VEGF secretion was mediated by TGF- $\beta$  since a neutralizing Ab to TGF- $\beta$  inhibited VEGF secretion from PSG1-treated extravillous trophoblast cells. Therefore at least some PSGs have pro-angiogenic properties which are mediated by two different mechanisms; the TGF- $\beta$ -mediated induction of VEGF-A, and the direct interaction of PSGs with GAGs on the surface of endothelial cells.

The contribution of the PSG1-integrin  $\beta$ 1 interaction in the pro-angiogenic activity of this protein remains to be investigated as endothelial cells express high levels of this integrin.

## PSGs and TGF- $\beta$

A relationship between PSGs and TGF- $\beta$  was first described *in vitro* by the Dveksler laboratory (Snyder *et al.*, 2001). Treatment of human monocytes with recombinant PSG1, PSG6, and PSG11 generated in insect cells induced the secretion of TGF- $\beta$ 1 in a dose-dependent manner. In addition, a protein composed of the N-terminal domain of PSG17, PSG17N, generated in insect or CHO-K1 cells and purified following SDS-PAGE electrophoresis under denaturing conditions, was also shown to induce the secretion of TGF- $\beta$ 1 by murine bone marrow derived macrophages and the murine macrophage cell line RAW 264.7 (Ha *et al.*, 2005). Subsequent studies using other preparations of recombinant PSGs, including PSG1 composed of the N-A2-B2 domains fused to the FLAG, V5 and His tags (PSG1-FLAG-V5- His) or to the Fc portion of the IgG1 heavy chain (PSG1-Fc), and two murine PSGs composed of the N1 and A-domains fused to the His and FLAG tags (PSG22N1A-His-FLAG and PSG23N1A-His-FLAG) were shown to induce the secretion of TGF- $\beta$ 1 by several cells including dendritic cells, extravillous trophoblast cell lines, endothelial cells

and NK cells (Blois *et al.*, 2012, Ha *et al.*, 2010, Wu *et al.*, 2008).

Recently, we found that recombinant PSG1-Fc, PSG1-FLAG-V5-His, murine PSG22N1A-His-FLAG and PSG23N1A-His-FLAG purified from the supernatant of transfected CHO-K1 cells are associated with TGF- $\beta$ 1, which is normally produced by this cell line (Blois *et al.*, 2013; G.Dveksler, unpublished results). Therefore, while it is still possible that these PSGs induce TGF- $\beta$ 1 in the cells listed above, new experiments should be performed with preparations of these proteins lacking associated TGF- $\beta$ 1 before a definitive conclusion can be reached regarding their ability to induce the secretion of this cytokine.

Importantly, proteins generated in insect cells or after separation by SDS-PAGE do not have associated TGF- $\beta$ 1, indicating that the initial reports on the ability of some human PSGs and PSG17N to induce TGF- $\beta$ 1 are not compromised by our recent discovery.

Since native PSG1 purified from the serum of pregnant women and recombinant PSG1 were both shown to associate with TGF- $\beta$ 1, we asked whether PSG1 is a novel TGF- $\beta$ 1 binding protein and whether it has the potential to activate latent TGF- $\beta$ . Most cells secrete at least one TGF- $\beta$  isoform (TGF- $\beta$ 1, -2 and -3) in the non-biologically active/latent form (Annes *et al.*, 2003). The non-covalent interaction of mature TGF- $\beta$  with the isoform-specific latency-associated peptide (LAP) blocks binding of mature TGF- $\beta$ s to the specific receptors. Therefore activation of latent TGF- $\beta$  requires agents that disrupt the association of LAP with the mature domain either by cleaving LAP or by altering its conformation (Shi *et al.*, 2011). An experimental approach employing a cell-free system indicated that PSG1 is able to activate the small latent complex (SLC) of TGF- $\beta$ 1 in a dose-dependent manner (Blois *et al.*, 2013). Because the small latent form of TGF- $\beta$ 2 is not commercially available, reporter assays were used and strongly suggested that PSG1 can also activate this isoform (Blois *et al.*, 2013). Whether PSG1 can activate the latent TGF- $\beta$ 3 isoform remains to be investigated. The Dveksler laboratory is currently testing whether other members of the human PSG family share with PSG1 the ability to activate TGF- $\beta$ 1 and TGF- $\beta$ 2.

The increased availability of biologically active TGF- $\beta$  mediated by PSG1 administration has biological consequences *in vivo*. Administration of PSG1 protected mice from dextran sulfate induced colitis and the observed protective effect of PSG1 was inhibited by co-administration of a neutralizing anti-TGF- $\beta$  antibody (Blois *et al.*, 2013). PSG1-treated mice expressed lower levels of the pro-inflammatory cytokines TNF- $\alpha$  and IFN- $\gamma$  compared to protein control-treated mice. In addition, mice receiving PSG1 had increased numbers of FoxP3+ LAP+ regulatory T cells (Tregs) in the colonic lamina propria. An increase in the number of Tregs upon PSG1 administration was also observed in a different *in vivo* experimental system. Mice injected with a vaccinia virus expressing PSG1 and challenged with *Listeria monocytogenes* had expanded numbers of CD25+ Foxp3+ cells when compared to vaccinia virus control-treated mice (Martinez *et al.*, 2012). We and others found that conditioned media obtained from the extravillous trophoblast cell line HTR-8 contains the active form of TGF- $\beta$  (Graham and Lala, 1991). The mechanism by which TGF- $\beta$  is activated in the media of these cells remains unknown but HTR-8 cells express PSG mRNA and PSG1 can be detected in the supernatant of these cells by ELISA (G. Dveksler, unpublished data). Therefore, it is possible that the reported HTR-8-mediated induction of inducible Tregs is associated with the production of PSG1 by this cell line

(Ramhorst *et al.*, 2012).

The actions of PSG1 on cells of the innate and adaptive immune system have been recently reviewed in (Martinez *et al.*, 2013). IL-10 and TGF- $\beta$  are suppressor cytokines, which are interrelated (Kitani *et al.*, 2003, Li *et al.*, 2006, Moore *et al.*, 2001). PSG1 has been shown to induce the secretion of IL-10 and the alternative activation of macrophages (Motran *et al.*, 2002). Whether these PSG1 functions and the reported modulation of dendritic cell maturation by PSG1 are a result of its newly discovered ability to activate TGF- $\beta$  remains to be elucidated (Martinez *et al.*, 2012). In addition, while some of the reported activities of human PSG1, -6 and -11 such as the ability to induce the secretion of IL-10 and IL-6 by monocytes, have also been reported for murine PSG17; the mechanism and signaling pathways behind the responses to human PSGs have not been elucidated and may differ from the signaling events involved in the response to murine PSGs, as different receptors and/or signaling mechanisms may have evolved separately resulting in similar functions (Ha *et al.*, 2005, Snyder *et al.*, 2001, Wessells *et al.*, 2000).

### PSGs and pregnancy pathologies

Several reports indicate that lower than normal serum concentrations of PSGs are associated with fetal growth restrictions (Grudzinskas *et al.*, 1983, Pihl *et al.*, 2009, Salem *et al.*, 1981, Towler *et al.*, 1977, Wurz *et al.*, 1981). In addition, a correlation between abnormally low PSG levels and preeclampsia was found in some studies but not others (Bersinger and Odegard, 2004, Pihl *et al.*, 2009, Silver *et al.*, 1993, Towler *et al.*, 1977). A critical evaluation of the published studies on the association of PSG levels and pregnancy complications is required based on the complexity of the CEACAM/PSG family. First, all measurements of PSGs reported in the cited publications were performed with polyclonal anti-PSG antibodies incapable of differentiating between the different gene products and the measurements of PSG levels sometimes differ in the trimester of pregnancy at which they were conducted. In addition, *in vitro* studies using sequence specific primers for different PSG family members indicate that PSGs are expressed at different levels (Camolotto *et al.*, 2010), which is supported by differences in the promoter sequences (Chamberlin *et al.*, 1994). Therefore measuring the concentration of all PSGs may not accurately reflect a possible association of abnormal concentrations of some members of the family with pregnancy pathology. Adding to these complications, is new information indicating that there is a big range of what may be considered "normal PSG levels". PSG gene copy numbers are polymorphic in the human population and they range from 12 to 30 in normal individuals (Chang *et al.*, 2013). Recently, Dewan and co-workers reported that a copy-number deletion of PSG11 and its alternatively spliced variants may confer risk for preeclampsia but further studies are required to confirm their findings (Zhao *et al.*, 2012). Expression of some PSGs has been reported in molar trophoblastic tissue (Hagiwara *et al.*, 1986, Leslie *et al.*, 1990) and differential expression of PSGs in intrauterine pregnancy and ectopic pregnancy have been recently suggested to have useful diagnostic value as part of a multiple biomarker test (Rausch *et al.*, 2011). In summary, while low PSG concentrations appear associated with some pregnancy complications, studies with more specific reagents are required as PSG levels could potentially be used as useful markers of placental function.

## Conclusions

Further studies are required to understand the functions and characterize the receptors and modes of action of the different PSGs expressed in humans, non-human primates, and rodents. In addition, the recently discovered association between PSG1 and TGF- $\beta$  suggests that PSGs may play a role in the well-established beneficial effects of pregnancy in patients suffering from autoimmune diseases such as multiple sclerosis and rheumatoid arthritis. This indicates that administration of PSG1 should be explored as a novel therapeutic agent in diseases related to undesired immune activation in non-pregnant individuals.

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