

Secreted modular calcium-binding proteins in pathophysiological processes and embryonic development

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Abstract

Objective: Secreted modular calcium-binding proteins (SMOCs) are extracellular glycoproteins of the secreted protein, acidic, and rich in cysteine-related modular calcium-binding protein family and include two isoforms, SMOC1 and SMOC2, in humans. Functionally, SMOCs bind to calcium for various cell functions. In this review, we provided a summary of the most recent advancements and findings of *SMOC1* and *SMOC2* in development, homeostasis, and disease states.

Data sources: All publications in the PubMed database were searched and retrieved (up to July 24, 2019) using various combinations of keywords searching, including *SMOC1*, *SMOC2*, and diseases.

Study selection: All original studies and review articles of *SMOCs* in human diseases and embryo development written in English were retrieved and included.

Results: *SMOC1* and *SMOC2* regulate embryonic development, cell homeostasis, and disease pathophysiology. They play an important role in the regulation of cell cycle progression, cell attachment to the extracellular matrix, tissue fibrosis, calcification, angiogenesis, birth defects, and cancer development.

Conclusions: *SMOC1* and *SMOC2* are critical regulators of many cell biological processes and potential therapeutic targets for the control of human cancers and birth defects.

Keywords: Angiogenesis; Cancer; Cell cycle; Embryonic development; *SMOCs*; Secreted protein, acidic and rich in cysteine

Introduction

The secreted protein, acidic and rich in cysteine (SPARC), also known as osteonectin (ON), or basement membrane-40, is an acidic extracellular matrix glycoprotein and plays an important role in bone mineralization, cell-matrix interactions, collagen binding, and bone remodeling.^[1-3] Secreted modular calcium-binding proteins (SMOCs) are extracellular glycoproteins of the SPARC-related modular calcium-binding protein family with two isoforms, *SMOC1* and *SMOC2*, in humans. SMOCs influence cell growth factor signaling, cell migration and proliferation, and angiogenesis.^[4,5] *SMOC1* was discovered in 2002, while *SMOC2* was identified thereafter.^[4,5] To date, their gene structures, patterns of expression, and functions have been precisely identified and studied for their involvement in multiple biological processes including embryonic development,^[6] cell cycle progression,^[7] cell attachment,^[8] tissue fibrosis,^[9,10] calcification,^[11] angiogenesis,^[12] and

tumor development.^[13,14] This review summarizes the recent advancements in *SMOC1* and *SMOC2* and their role in embryonic development and homeostasis as well as human diseases.

SMOC Molecular Structures and Expression in Cells and Tissues

The full-length complementary DNA (cDNA) of *SMOC1* was cloned in 2002 as a secreted modular glycoprotein and expressed in the basement membrane and the extracellular matrix of many tissues.^[13] The *SMOC1* gene is localized at chromosome 14q24.2 and encodes a protein of 434 amino acids with a molecular weight of 48,000.^[14] *SMOC1* protein contains five structural domains, including a follistatin-like domain, two thyroglobulin-like domains, a unique domain, and an helix-loop-helix motif (EF-hand)

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calcium-binding domain (or the extracellular calcium-binding [EC] domain). A subsequent study led by the same group of researchers also identified the *SMOC2* gene by search expressed sequence tag sequence databases.^[5] The *SMOC2* gene is localized at chromosome 6q27 and shares 55% of DNA sequence homology with *SMOC1*.^[5] *SMOC2* has a predicted protein structure that contains similar domains to *SMOC1*, except for its own unique *SMOC* domain.^[5] *SMOC1* protein is highly expressed in the ovary, brain, thymus, heart, skeletal muscle, liver, and lung, while *SMOC2* protein is expressed throughout the human body^[4,5] and on basement membranes and the extracellular matrix.^[7,15] Immunohistochemical analysis showed that *SMOC1* was expressed in myocytes and as a component of the basement membrane in blood capillaries besides the cell surface.^[4] Furthermore, *SMOC1* protein can bind to collagen IV through its EC domain in the basement membrane.^[4] The *SMOC1* interaction with collagen IV indicates that *SMOC1* distributes widely in the basement membrane; however, there are few studies to date demonstrating that *SMOC1* interacts with SPARC in the same manner. In contrast, *SMOC2* protein participates in the assembly of fibrils and is co-localized with fibronectin, the main component of the extracellular matrix.^[5] Thus, *SMOC2* acted as a component of fibrils and the characteristic of *SMOC2* co-localization with fibronectin may not be typical for the extracellular proteins. *SMOC2* has also been identified as a marker of intestinal stem cells,^[16] implying that it may have multiple pleiotropic roles.^[5,16] Taken together, although *SMOC1* and *SMOC2* shared high DNA sequence similarity, they have distinct functions in the human body.

Although recombinant *SMOC* proteins can be generated in *Escherichia coli*,^[16] such recombinant *SMOC* proteins show little biological activity due to the lack of the glycosylation process *in vitro*. Thus, there is an urgent need to reconstitute functionally glycosylated *SMOC* proteins for future research.

SMOC Functions in Cells and Embryo Development

Embryonic development

A previous study analyzed the expression and regulation of *SMOC1* and *SMOC2* in fetal gonad/mesonephros complexes and revealed their possible role in the gonad and mesonephros development, that is, *SMOC1* expression was elevated at approximately E10.75 in pre-sertoli and pre-granulosa cells but was significantly reduced in the Wilms tumor protein 1, splicing factor 1, and friend of GATA2 mutants. At embryonic day 13.5 (E13.5), *SMOC1* expression was reduced in granulosa cells but persisted in Sertoli cells, indicating a sexually dimorphic requirement in cell lineage differentiation. In contrast, *SMOC2* protein was expressed in Leydig cells and mesonephroi and Wnt4 mutant ovaries, while *SMOC2* expression required functional Hedgehog signaling in the testes, mesonephroi, and kidneys.^[17] *SMOC1* was also reported to be essential for post-gastrulation development in *Xenopus* by inhibition of bone morphogenetic protein (BMP) signaling^[18] and for ocular and limb development in both humans and mice.^[19] Furthermore, during mouse embryogenesis, *SMOC1* is

expressed in the early mouse embryo at E7 in the entire endodermal basement membrane zone of the embryo proper.^[20] *SMOC1* messenger RNA (mRNA) is expressed in mesenchymal as well as epithelial cells deriving from all three germ layers in the early stages of mouse development. This broad and organ-specific *SMOC1* expression underscores *SMOC1*'s multi-functional roles in mouse embryogenesis.^[20] Thomas *et al*^[18] demonstrated in mice that *SMOC2* mRNA was present in mid-gestation embryos, particularly in the forelimb, hindlimb, somites, and branchial arches.^[7,21] *SMOC2* mutations were associated with developmental dental defects.^[22]

Molecularly, a previous study revealed that *SMOCs* function through the Smad (homologies to the *Caenorhabditis elegans* "small" worm phenotype and *Drosophila* mothers against decapentaplegic family of genes) signaling to activate mitogen-activated protein kinase (MAPK) signaling during the joint formation in the linker region.^[23] Similar to the BMP family and its endogenous antagonists that regulate embryonic organ development,^[18] *SMOC1* can act as an antagonist of BMP signaling by mediating the MAPK-mediated Smad phosphorylation in the linker region.^[18] In this regard, *SMOC1* was not only a crucial factor to influence post-gastrulation development, but also regulated the formation of anatomic patterning, cell, or tissue fate specification through modulation of its related signaling.^[18] Indeed, the *C. elegans* *SMOC1* homolog acted as a positive modulator of the BMP signaling during development of *C. elegans*, and human *SMOC1* and *SMOC2* transgenes could each partially rescue phenotypes of *SMOC1* mutants in worms, suggesting that *SMOC1* modulation of BMP signaling was evolutionarily conserved.^[24] The most recent findings of the function and molecular mechanisms of *SMOCs* action in embryo developments and related diseases are summarized in Table 1.

Bone calcification

The extracellular matrix proteins (ECM), like SPARC, play a key role in the mineralization process of bone.^[34] As a member of the SPARC protein family, *SMOC1* is also highly expressed in the bone and the bone marrow mesenchymal stem cells.^[35] Knockdown of *SMOC1* expression inhibited osteoblast differentiation and mineralization, whereas *SMOC1* overexpression enhanced osteoblast differentiation,^[35] indicating that *SMOC1* promotes osteoblast differentiation and mineralization. Furthermore, *SMOC1* expression peaked on day 1 during post-natal induction of osteoblast differentiation, and gradually decreases in the subsequent 7 days. These studies suggest that *SMOC1* positively regulates the early stage of osteoblast differentiation, but it may be not required for the late stage of osteoblast differentiation, similar to other ECM proteins.^[35] Further investigation should focus on bettering our understanding of the underlying molecular mechanisms of *SMOC1* action in calcification, such as *SMOC1* in the regulation of receptor-mediating signaling.

SMOC2 was originally identified in the extracellular matrix of articular cartilage^[5,36] and *SMOC2* inhibited

Table 1: Role of SMOCs in embryo development and diseases.

References	Species	Embryo development/ diseases	SMOCs	Brief descriptions
Thomas <i>et al</i> (2009) ^[18]	Xenopus	Post-gastrulation development	SMOC1	<i>SMOC1</i> worked as an antagonist of BMP
Thomas <i>et al</i> (2017) ^[23]	Mammalian	Mammalian joint formation	SMOC	MAPK signaling was activated by <i>SMOC</i> via Smad phosphorylation in the linker region
Marchant <i>et al</i> (2017) ^[25]	Brachycephalic dogs	Facial form	SMOC2	Effects of <i>SMOC2</i> on the facial skeleton are in a dose- dependent manner.
Mommaerts <i>et al</i> (2014) ^[6]	Zebrafish	Myeloid cells of the anterior lateral plate mesoderm development	SMOC2	<i>SMOC2</i> is involved in the development of myeloid cells of the anterior lateral plate mesoderm of zebrafish via <i>BMP</i> related gene transcription,
Birlea <i>et al</i> (2010) ^[26]	Isolated Romanian community	Generalized vitiligo, autoimmune thyroid disease, rheumatoid arthritis	SMOC2	<i>SMOC2</i> is a risk locus for generalized vitiligo, in addition to other autoimmune diseases. The odds ratio is 7.445.
Alkhateeb <i>et al</i> (2010) ^[27]	Jordanian Arab patients	Vitiligo	SMOC2	The variant rs13208776 in the <i>SMOC2</i> gene is not correlated with the risk of vitiligo in Jordanian Arab patients
Jamshidi <i>et al</i> (2017) ^[28]	Iranian patients	WAS	SMOC1	Causative mutation of <i>SMOC1</i> is the homozygous missense mutation (NM_001034852.2 (<i>SMOC1</i>):c.367T>C) in exon 3.
Alfawaz <i>et al</i> (2013) ^[29]	Consanguineous Pakistan family	Oligodontia and microdontia	SMOC2	<i>SMOC2</i> mutation in exon 8, c.681T>A (p.C227X) is critical for tooth development
Rainger <i>et al</i> (2011) ^[30]	Human and mice	WAS	SMOC1	<i>SMOC1</i> is crucial for the developments of eyes and limbs via regulation of BMP signaling.
Okada <i>et al</i> (2011) ^[19]	Human and mice	MLA	SMOC1	The MLA locus is mapped to 14q24 and three homozygous (one nonsense and two splice site) mutations are identified in <i>SMOC1</i> . It is also detected in the optic stalk, ventral optic cup, and limbs of mouse embryos during development.
Alkhateeb <i>et al</i> (2013) ^[31]	Jordanian Arab population	Autoimmune thyroid diseases	SMOC2	<i>SMOC2</i> single nucleotide polymorphisms may act as the dominant polymorphism in autoimmune thyroid diseases susceptibility.
Lawrance-Owen <i>et al</i> (2013) ^[32]	Human	Digit ratio	SMOC1	Exposure to prenatal sex hormones influences the digit ratio possibly by <i>SMOC1</i> mediation.
Al-Dabbagh <i>et al</i> (2017) ^[33]	Saudi population	PACG	SMOC2	Polymorphism of <i>SMOC2</i> (G>A) may be associated with PACG, and probably is a risk factor.

SMOCs: Secreted modular calcium-binding proteins; WAS: Waardenburg anophthalmia syndrome; MLA: Microphthalmia with limb anomalies; PACG: Primary angle-closure glaucoma; BMPs: Bone morphogenetic proteins; MAPK: Mitogen-activated protein kinase[.

osteogenic differentiation and extracellular matrix mineralization through the *SMOC2* EC domain.^[11] Ectopic or excessive calcification contributes to diseases like chondrocalcinosis by calcium deposition in the skin or blood vessels. *SMOC2* was able to downregulate the mineralization process in human umbilical vein endothelial cells, whereas *SMOC-ΔEC*, which lacks the EC domain, was unable to inhibit BMP2 signaling and *SMOC-EC* (the EC domain only)-induced activity of the BMP2 signaling.^[23] However, knockdown of *SMOC2* expression did not affect MC3T3-E1 cell calcification.^[23] This implies that an increase in *SMOC2* levels might result from cell mineralization, rather than as a cause. These findings are contradictory to a previous observation showing that *SMOC2* was required for bone development.^[37] One of the plausible reasons may be because MC3T3-E1 cells were derived from neonatal mouse calvaria and are already committed to calcification,^[38] in which *SMOC2* may have a less regulatory role.

Altogether, *SMOCs* are required for the tissue calcification process, although the regulatory functions of *SMOC1* are different from *SMOC2*. To date, there are few publications that describe the role of other *SMOC* domains, especially how these unique domains participate in calcification and BMP-related signaling, which warrants further investigation.

Angiogenesis

SMOC1 possesses a pro-angiogenic activity and is a target of anti-angiogenic microRNA-223 (miR-223).^[12] Knockdown of *SMOC1* expression significantly attenuated the sprouting of endothelial cells in the aortic rings and delayed retinal angiogenesis in *SMOC1^{+/-}* mice.^[12] The inhibitory effect of *SMOC1* silencing on angiogenesis was associated with upregulation of endoglin expression, which is an endothelium-specific type III auxiliary receptor in the transforming growth factor- β (TGF- β) receptor family and can induce Smad2 phosphorylation. Mouse endothelial cells were able to secrete *SMOC1* protein to inhibit expression of the activin-like kinase (ALK) 5 and promote TGF- β signaling and ALK1 activation, resulting in endothelial cell proliferation and angiogenesis.^[12] Furthermore, hypoxia can stimulate transcription of *SMOC1* mRNA in human endothelial cells by downregulation of miR-223 expression.^[12] *SMOC2* overexpression acts in parallel synergize with vascular endothelial growth factor (VEGF) or basic fibroblast growth factor to stimulate DNA synthesis and the formation of endothelial cell network-like structures, whereas *SMOC1* small interfering RNA (siRNA) treatment inhibited endothelial cell network formation and proliferation. These *in vitro* data were confirmed in mice with sub-dermal implantation of Matrigel plugs containing *SMOC2* cDNA adenovirus.^[39] Similarly, *SMOC2* can promote cell migration, angiogenesis, and migration beyond the enhancement of the neointima formation.^[8] Accordingly, *SMOC2* may promote the pathological progression of endometriosis.^[40]

Although there is high sequence similarity between *SMOC* and SPARC, these proteins have opposing effects on angiogenesis.^[7] For example, SPARC was able to inhibit

gastric cancer angiogenesis by attenuating expression of VEGF and matrix metalloproteinase 7.^[41] The functional balance between the *SMOCs* and SPARC ultimately determines their effects on endothelial cell proliferation, angiogenesis, and other pathogenic processes. Further studies should focus on the molecular mechanisms by which *SMOCs* regulate angiogenesis.

In addition, a previous study reports that the smooth muscle associated protein 2 (*SMAP2*), originally isolated from human aortic 3V cDNA library, showed a similarity to *SMOC2*.^[42] The *SMAP2* protein contains two thyroglobulin type-1 (Tg1) domains, two EF-hand calcium-binding domains, and a putative signal sequence domain. The Tg1 domain can regulate proteolysis.^[42] *SMAP2* mRNA transcripts are detected predominantly in the aorta, skeletal muscle and heart, ovary, testis, stomach, small intestine and colon, thyroid gland, mammary gland, and prostate. *SMAP2* expression was upregulated by 5.4 fold at 7 days post-surgery in a rat model of intraluminal balloon injury in the right carotid artery. Its expression was activated at day 3 and remained up to 14 days after surgery, suggesting that *SMAP2* may be closely related to the proliferation and migration of vascular smooth muscle cell after vascular injury.

Cell cycle regulation

SPARC can directly bind to various growth factors, including the platelet-derived growth factor (PDGF),^[43] VEGF,^[44] TGF- β ,^[45] or fibroblast growth factor 2^[46] to inhibit receptor-mediated signaling. In contrast, SPARC deficiency promoted the proliferation of mesangial cells, fibroblasts, and smooth muscle cells.^[47] Similarly, high levels of *SMOC2* expression were detected in the G1/S phase of the cell cycle in fibroblasts but these were reduced during the process of serum-stimulated cell cycle progression, suggesting that *SMOC2* inhibits cell cycle progression and cell proliferation in fibroblasts.^[7] In contrast, serum stimulation did not change *SMOC1* expression in fibroblasts Swiss 3T3. These data suggest that both *SMOC1* and *SMOC2* have differing roles in the regulation of the cell cycle although the underlying molecular mechanisms are unclear. Furthermore, *SMOC2* is required for growth factor-induced mitogenesis, especially for TGF- β .^[7] *SMOC2* ablation abrogated cyclin D1 levels in the G1 phase of the cell cycle in 3T3 cells.^[7] Interestingly, *SMOC2* ablation also reduced PDGF-induced DNA synthesis by 46% in cytomegalovirus-green fluorescent protein (GFP)-transfected cells, while *SMOC2* ablation only decreases the PDGF-induced DNA synthesis by 4% in cyclin D1-overexpressing Swiss 3T3 cells.^[7] These studies suggest that cyclin-D1-overexpressing cells had better responses to growth factor(s), independent of *SMOC2*. Actually, *SMOC2* deficiency did not significantly affect the PDGF-mediated PDGF β R phosphorylation, regardless of PDGF addition,^[7] or the PDGF-induced MAPK and protein kinase B (Akt) signaling and DNA synthesis.^[7] Hence, *SMOC2* may not be required for the PDGF β R signaling. *SMOC2* is required for the effective activation of integrin-linked kinase (ILK). However, the precise mechanisms by which *SMOC2* regulates ILK activation remain to be clarified. Defective ILK signaling due to the *SMOC2* deficiency could retard the

G1/S transition of the cell cycle and may be involved in SMOC2-regulated cyclin D1 expression.^[7]

Cell attachment

The EC domain of SMOC proteins is an autonomously folding domain and functions to promote the adhesion of skin HaCaT cells in a heparin sulfate-dependent manner.^[48] Downregulation of SMOC1 expression reduces endothelial cell adhesion to collagen I, laminin-III, and/or Matrigel.^[12] The EC domain of SMOC1 contains unique basic amino acids, the glycosaminoglycan-binding site, which are not present in other SPARC family proteins.^[48] However, heparin sulfate proteoglycans on the cell surface do not guarantee the successful binding of SMOCs to cell adhesion molecules because HaCaT cells do not adhere to the EC domain of SMOC1 in the presence of a chelating reagent, even if the recombinant EC domain can maintain such a heparin-binding activity.^[48] SMOC2, unlike SMOC1, does not promote adhesion of HaCaT cells, although there are a high sequence similarity of heparin-binding sites between SMOC1 and SMOC2.^[48] Maier *et al*^[8] previously reported that SMOC2 was able to selectively promote the adhesion of epidermal cells, dependent on the EC domain of SMOC2 and the integrins $\alpha v \beta 6$ and $\alpha v \beta 1$, but did not affect fibroblast adhesion. Moreover, the EC domain of SMOC2 can induce cell migration and wound healing.^[8] In basement membranes, there are many heparin sulfate proteoglycans, such as collagen XVIII and perlecan. There is a need for further investigation as to how SMOCs interact with them and regulate cell adhesion and migration.

Alteration of SMOCs in human diseases

Thus far, we reviewed and discussed SMOCs and their function in cells and tissues. Some studies have also revealed the alteration of SMOC1 and SMOC2 functions in the pathogenesis and progression of human diseases. Generally, alterations of SMOC1 and SMOC2 can occur at the genetic, transcriptional, and post-translational levels and the aberrant SMOC1 and SMOC2 expression can lead to the development of human diseases. We discuss them in more detail below.

Birth defects

SMOC1 homozygosity mapping and subsequent targeted mutation analysis revealed that SMOC1 mutations were associated with ophthalmo-acromelic syndrome (OAS), also known as Waardenburg Anophthalmia syndrome. OAS is defined by a combination of eye malformations, most commonly bilateral anophthalmia, with post-axial oligosyndactyly.^[66] SMOC1 mutations include nonsense, frame-shift, and missense mutations. The latter mutations occurred in the second Tg1 domain of the protein, while the targeted pre-conditional gene-trap mutation of Smoc1 [Smoc1(tm1a)] results in a reduction of SMOC1 mRNA levels to 10% of the wild-type (WT) SMOC1 levels.^[66] A recent study also reported a novel homozygous missense SMOC1 mutation (c.812G>A; p.Cys271Tyr) in a family of patients with Waardenburg anophthalmia syndrome.^[49]

Tissue inflammation and fibrosis

Persistent chronic inflammation can lead to tissue fibrosis. A previous study showed that treatment of rat mesangial cells with interleukin 1 beta induced nitric oxide (NO) production, but reduced SMOC1 expression in rat mesangial cells.^[50] The blockade of the inducible nitric oxide synthases activity upregulates SMOC1 expression to enhance rat glomerular inflammation, which is accompanied by an increase in the deposition of fibrin in the glomeruli in a rat model of anti-Thy1.1-induced chronic glomerulonephritis.^[51] Furthermore, knockdown of SMOC1 expression using SMOC1 siRNA decreased TGF- β expression and activity of the TGF- β signaling, which in turn reduced Smad activation and its target gene expression.^[51] Interestingly, NO was able to inhibit SMOC1 expression and attenuate TGF- β -mediated signaling.^[51] However, the specific mechanisms underlying the regulation of NO on SMOC1 expression remains unclear. SMOC2 has similar biological effects on the pathological process of fibrosis. For example, Gerarduzzi *et al*^[9] demonstrated that SMOC2 is a key regulator of the pathological secretome in a damaged kidney and that SMOC2 promoted kidney fibrosis. In contrast, neutralization of SMOC2 attenuated the TGF- β -induced fibrosis in NIH 3T3 cells.^[9] Intriguingly, SMOC2 overexpression was able to induce mouse kidney fibrosis and deteriorated inflammation-induced kidney fibrosis.^[9]

Bleomycin, a cytotoxic anti-cancer agent, can inhibit DNA synthesis but induce reactive oxygen species production, which may also cause pulmonary fibrosis.^[10] A recent study showed that reduction of SMOC2 expression was able to attenuate bleomycin-induced pulmonary fibrosis through inhibition of TGF- β signaling.^[8] Moreover, SMOC2 suppression also attenuated the nuclear factor kappa-B (NF- κ B) signaling and inflammatory responses.^[10] Besides, SMOC2 deficiency can ablate the bleomycin-induced macrophage activation and neutrophil infiltration *in vivo* and *in vitro*.^[10] The pathological score, collagen accumulation, and pulmonary fibrosis degrees were all lower in SMOC2^{-/-} mice than in WT mice after bleomycin treatment.^[10] Similarly, the levels of NF- κ B phosphorylation and TGF- β 1 expression in the lung were also significantly lower in the SMOC2^{-/-} mice than in SMOC2 WT mice.^[10] These data suggest that SMOC2 participates in the pathogenic process of pulmonary fibrosis and that further study could lead to the use of SMOC2 as a potential therapeutic target to treat pulmonary fibrosis. Also, SMOC2 overexpression exacerbated the fibrotic process, whereas knockdown of SMOC2 expression attenuated TGF- β 1-mediated fibrosis.^[10] These data indicate that SMOC2 could be a potential regulator of TGF- β 1 signaling and fibrosis.

However, a previous study of indomethacin and retinoic acid (RA) in the modification of mouse intestinal inflammation and fibrosis revealed that indomethacin/2,4,6-trinitrobenzene sulfonic acid (TNBS) enhanced and RA reduced inflammation, tissue destruction, and fibrosis in a mouse model of TNBS-induced intestinal fibrosis, in which SPARC expression was inversely associated with fibrosis, but not with inflammation.^[52] This study may indicate the differential role of SMOCs *vs.* SPARC in the regulation of tissue fibrosis.

Cancer development and progression

It is well known that interactions between malignant cells and the extracellular milieu are critical for cancer development and progression.^[53] SPARC, as a set of ECM proteins, can affect stromal and desmoplastic responses to tumor cells.^[54] The role of SPARC in the regulation of ECM production could make it a candidate conditioner of tumors.^[54] Consistent with this speculation, SPARC deficiency promoted tumor growth in mice,^[55] and SMOCs also regulate malignant cell proliferation.^[56] At the chromosome level, the *SMOC2* gene region contains several tumor suppressor genes.^[57,58] Previous studies revealed that *SMOC2* expression was downregulated in gall bladder carcinoma^[59] and advanced breast cancer.^[60] In contrast, levels of *SMOC2* mRNA were upregulated by 36.55 fold in metastatic head and neck squamous cell carcinoma tissues.^[61] *SMOC2* modulated keratinocyte adhesion through binding to integrins^[8] and the latter is important in the regulation of cell anoikis resistance and maintenance of cancer stem cell phenotypes.^[62,63] By targeting a cancer stem cell signature gene, *SMOC2* was able to overcome chemoresistance and inhibit proliferation of endometrial carcinoma cells.^[64] Furthermore, *SMOC1* was thought to be a cancer-related protein related to tenascin-C expression, which is an ECM protein and is highly expressed in a large variety of human cancers.^[65] *SMOC1* expression was upregulated in cancer tissues, such as oligodendrogliomas and astrocytic tumors.^[66] *SMOC1* was able to inhibit tenascin-C-induced migration of glioma U87 cells.^[65] In this section, we further discuss the role of SMOCs in the regulation of development and progression of hepatocellular carcinoma (HCC), colorectal cancer, and lung adenocarcinoma.

Hepatocellular carcinoma

To date, the role of *SMOC2* on HCC development and progression is controversial. For instance, *SMOC2* expression was significantly upregulated in HCC tissues and upregulated *SMOC2* expression induced expression of the MAPK/extracellular regulated protein kinases and AKT signaling.^[14] Moreover, upregulated *SMOC2* expression was able to promote cell cycle progression and tumor cell proliferation through an increase in cyclin D1 expression.^[7] Hence, *SMOC2* may be further studied as a potential therapeutic target to treat HCC clinically. In contrast, Huang *et al*^[13] reported that *SMOC2* expression was downregulated in HCC tissues and *SMOC2* expression was associated with a favorable overall survival and disease-free survival of patients with HCC. The further experimental study indicated that *SMOC2* overexpression suppressed HCC cell migration, proliferation, and cell cycle progression.^[13] Thus, future study is needed to clarify the role of *SMOC2* in HCC; however, to date, there is no single study of *SMOC1* in HCC reported in the literature.

Colorectal cancer

SMOCs were shown to be colorectal cancer biomarkers and potential therapeutic targets for colorectal cancer patients.^[67] For example, recent studies revealed that high levels of *SMOC2* and leucine-rich repeat containing G

protein-coupled receptor 5 (*Lgr5*) expression were observed in human colorectal cancer tissues^[68] and that *SMOC2* overexpression was associated with the potent metastatic capacity of *Lgr5*-upregulated colorectal cancer cells.^[67] Elevated *SMOC2* expression was detected in the invasive front of tumor tissues and *SMOC2* expression is indispensable for L1-mediated invasion of colorectal cancer.^[68] In parallel, *SMOC2* overexpression was able to induce colorectal cancer motility and liver metastasis and the oncogenic effect of *SMOC2* was associated with modulation of ILK activity, whereas treatment of colorectal cancer cells with QLT0267, a selective p38 MAPK inhibitor or knockdown of ILK expression reduced the *SMOC2*-mediated oncogenic effects.^[68]

The “serrated neoplasia pathway” was considered a route of colorectal cancer development,^[69] and sessile serrated adenoma/hyperplastic polyps and traditional serrated adenomas (TSAs) are different premalignant lesions in colorectal cancer development. Methylation of the *SMOC1* gene gradually increased during TSA development, but rarely occurred in sessile serrated adenoma/hyperplastic polyps.^[69] Consistent with the elevated *SMOC1* gene methylation, *SMOC1* expression was downregulated in TSAs, but not in sessile serrated adenoma/hyperplastic polyps.^[70] Therefore, *SMOC1* might serve as a diagnostic marker for serrated lesions.

Lung adenocarcinoma

Lung adenocarcinoma is one of the most commonly diagnosed malignancy in the world and lung cancer metastasis usually leads to a poor prognosis.^[71] Currently, the molecular mechanisms underlying lung cancer metastasis remain to be defined. *SMOC2* has been identified as a pro-metastatic matricellular protein and knockdown of *SMOC2* expression inhibited lung cancer metastasis.^[72] Brady *et al*^[72] showed that expression of Aryl hydrocarbon receptor nuclear translocator-like 2 (*Arntl2*), a paralog of the circadian transcription factor *Arntl*,^[73] was upregulated in metastatic lung adenocarcinoma and associated with poor prognosis. Another study revealed that the pro-metastatic effects of *Arntl2* are attributed to a ten-fold increase in *SMOC2* expression and that knockdown of *Arntl2* expression significantly reduced *SMOC2* expression in metastatic lung cancer cells.^[72] Their additional data showed that both *Arntl2* and *Clock* were able to bind to the proximal *SMOC2* gene promoter, implying that both *Arntl2* and *Clock* can directly regulate *SMOC2* expression in metastatic lung adenocarcinoma.^[74,75] Given that there are multiple extracellular proteins involved in metastasis of different human cancers,^[74,75] further studies are needed to better understand how these genes interact with SMOCs to impact lung cancer metastasis.

Conclusions

SMOC1 and *SMOC2* are ECM proteins of the SPARC family with differential and unique characteristics in tissue distribution and cell functions. *SMOC1* was necessary for the early stage of osteoblast differentiation, while *SMOC2* inhibited osteoblast differentiation.^[11,35] SMOC proteins

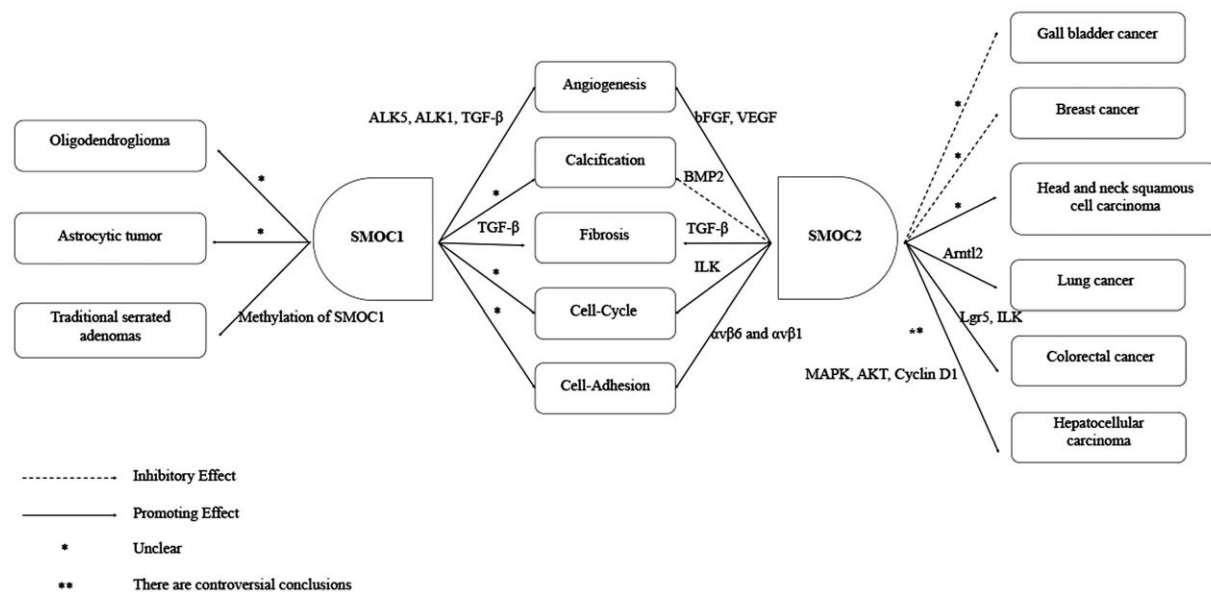


Figure 1: Illustration of basic SMOC functions in cells. Both SMOC1 and SMOC2 proteins are differentially expressed in various cells and tissues, which also function differently for embryonic development, homeostasis, and disease processes, that is, both SMOC1 and SMOC2 can promote angiogenesis, fibrosis, cell cycle progression, and cell adhesion. Molecularly, SMOC1 was able to regulate ALK5, ALK1, TGF- β , whereas SMOC2 regulated bFGF, VEGF in the angiogenesis process, although both SMOC1 and SMOC2 could regulate TGF- β in tissue fibrosis. SMOC2 can also regulate ILK in cell-cycle progression and α v β 6 and α v β 1 in the cell adhesion process, whereas the molecular mechanisms of SMOC1 in these two processes are not currently clear. SMOC1 enhanced calcification, but the underlying molecular mechanism remains to be defined. SMOC2 was able to inhibit calcification via the BMP2 signaling. Furthermore, SMOC1 deteriorated progression of oligodendroglomas, astrocytic tumors, and traditional serrated adenoma, while SMOC2 is a risk factor for the development of head and neck squamous cell carcinoma, lung cancer, and colorectal cancer. SMOC2 can also act as a tumor suppressor in gallbladder and breast cancers. The potential mechanism involved in traditional serrated adenoma development was due to SMOC1 methylation, but the molecular mechanisms of SMOC1 involved in oligodendrogloma and astrocytic tumor are unclear. SMOC2 might regulate Arntl2 in lung cancer, Lgr5, and ILK in colorectal cancer development and interact with MAPK, AKT, and Cyclin D1 in HCC, but the role of SMOC2 in HCC development remains controversial. The potential mechanisms of SMOC2 involved in gallbladder and breast cancers as well as head and neck squamous cell carcinoma remains unknown. ALK: Activin-like kinase; Akt: protein kinase B; Arntl2: Aryl hydrocarbon receptor nuclear translocator-like 2; bFGF: Basic fibroblast growth factor; BMP: Bone morphogenetic protein; HCC: Hepatocellular carcinoma; ILK: Integrin-linked kinase; Lgr5: Leucine-rich repeat containing G protein-coupled receptor 5; MAPK: Mitogen-activated protein kinase; SMOC: Secreted modular calcium-binding protein; TGF- β : Transforming growth factor- β ; VEGF: Vascular endothelial growth factor.

were able to promote angiogenesis, fibrosis, and cell motility and adhesion. The basic SMOCs functions and signaling are summarized in Figure 1. It should be noted that the pro-angiogenic effects of SMOCs oppose those of SPARC and the imbalance between the SMOCs and SPARC might determine the outcome and potential pathological changes during disease.^[7] Thus, although the EC domain of SMOCs can regulate the mineralization process and cell attachment, functions of other SMOC protein domains, especially their unique domains, require further investigation. Furthermore, the regulatory effect of SMOC2 on cell adhesion depends on the context of individual cell types.^[48] The mechanisms by which SMOCs as the soluble matrix proteins enter into their targeted cells and bind to their related receptor(s) to regulate cell signaling remain unstudied. Besides, the molecular mechanisms underlying how SMOCs regulate the cell cycle, interact with growth factors, and contribute to cancer development and progression as well as regulate embryonic development should be clarified in future research.

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Conflicts of interest

None.

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