



Heparan Sulfate Biosynthesis Enzymes in Embryonic Stem Cell Biology

Christoffer Tamm, Lena Kjellén, and Jin-Ping Li

Department of Medical Biochemistry and Microbiology, Uppsala University, Uppsala, Sweden

Summary

Embryonic stem (ES) cells are derived from the inner cell mass of the blastocyst and can give rise to all cell types in the body. The fate of ES cells depends on the signals they receive from their surrounding environment, which either promote self-renewal or initiate differentiation. Heparan sulfate proteoglycans are macromolecules found on the cell surface and in the extracellular matrix. Acting as low-affinity receptors on the cell surface, heparan sulfate (HS) side chains modulate the functions of numerous growth factors and morphogens, having wide impact on the extracellular information received by cells. ES cells lacking HS fail to differentiate but can be induced to do so by adding heparin. ES cells defective in various components of the HS biosynthesis machinery, thus expressing differently flawed HS, exhibit lineage-specific effects. Here we discuss recent studies on the biological functions of HS in ES cell developmental processes. Since ES cells have significant potential applications in tissue/cell engineering for cell replacement therapies, understanding the functional mechanisms of HS in manipulating ES cell growth in vitro is of utmost importance, if the stem cell regenerative medicine from scientific fiction ever will be made real. (*J Histochem Cytochem* 60:943–949, 2012)

Keywords

proteoglycans, heparan sulfate, embryonic stem cells, development, extracellular signaling

Impact of Heparan Sulfate in Animal Development

Heparan sulfate proteoglycans (HSPGs) at the cell surface and in the extracellular matrix have important functions during embryonic development. They act as essential co-receptors and their interaction with various exogenous factors allows the generation and maintenance of morphogen gradients. The sulfation pattern of the heparan sulfate (HS) side chains, deciding the factors that will bind and the affinity of the interaction, is determined largely during biosynthesis. In addition, after biosynthesis, the HS chains may be modified by the endo-6-*O*-sulfatases (Sulf 1 and 2, see below) or by heparanase, an endo-glycosidase that generates HS fragments. Biosynthesis occurs in the Golgi compartment and involves the action of several different enzymes, some of which occur in more than one isoform. The enzymes that will be mostly discussed here are Ext1, which, together with Ext2, forms the functional HS polymerase, and the glucosaminyl *N*-deacetylase/sulfotransferases (Ndts),

which are responsible for the HS *N*-sulfation that controls the subsequent enzymatic reactions, thereby affecting the overall structure of HS chains. Biosynthesis of HS is covered in two companion mini-reviews (Kreuger & Kjellén 2012; Multhaupt and Couchman 2012).

HS is essential for animal development, as evidenced by mutational studies in model organisms, including *Drosophila melanogaster*, mouse, *Caenorhabditis elegans* and zebrafish. In the mouse, complete interruption of HS biosynthesis by obliteration of the polymerases, Ext1 or Ext2 (Lin et al. 2000; Stickens et al. 2005), led to early termination of mouse embryonic development. Selected elimination of the enzymes involved in modification of HS structure resulted in strikingly varied phenotypes in mice. Inhibition of *N*-sulfation by

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Corresponding Author:

Christoffer Tamm, IMBIM, Box 582, SE-751 23, Uppsala, Sweden.

E-mail: christoffer.tamm@imbim.uu.se

knockout of *Ndst1* caused an overall downregulation of the modification reactions downstream, resulting in HS with greatly reduced charge density (Ringvall et al. 2000). Thus, *Ndst1*^{-/-} mice showed severe developmental defects in multiple organs, including skeleton, brain, and lungs, and died shortly after birth due to respiratory failure (Fan et al. 2000; Ringvall et al. 2000). In comparison, *Ndst2*^{-/-} mice did not show any obvious defect, except an abnormal morphology in mast cells (Forsberg E et al. 1999; Humphries et al. 1999). Elimination of the single gene-coded enzymes in HS biosynthesis, glucuronyl C5-epimerase (*Glce*) and hexuronyl 2-O-sulfotransferase (*Hs2st*), resulted in similar phenotypes, for example, neonatal lethality with kidney agenesis and multiple skeletal malformations (Bullock et al. 1998; Li et al. 2003). Most glucosaminyl 6-O-sulfotransferase 1 (*Hs6st1*) knockout embryos died during late gestation, and those that survived were smaller than wild-type littermates, with various developmental abnormalities (Habuchi et al. 2007). Mice deficient in glucosaminyl 3-O-sulfotransferase 1 (*Hs3st1*) showed intrauterine growth retardation and postnatal lethality, and they displayed a proinflammatory phenotype (Shworak et al. 2002; Shworak et al. 2010).

Further functional regulation of HS in embryo development appears linked to the postsynthesis modification by *Sulf1* and *2* and the endo-glucuronidase heparanase (*Hpse*). Although *sulf1* and *sulf2* single as well as double mutants appear normal at birth, the double mutants die shortly after birth (Ai et al. 2007; Langsdorf et al. 2007; Lamanna et al. 2008). Surprisingly, neither complete elimination of the heparanase gene nor overexpression of the enzyme affected embryo development (Zcharia et al. 2004; Zcharia et al. 2009).

Embryonic Stem Cells and Induced Pluripotent Stem Cells

Pluripotent embryonic stem (ES) cells are permanent cell lines derived from the inner cell mass of the blastocyst (Evans and Kaufman 1981; Martin 1981) and can be maintained and expanded in culture by addition of factors that promote proliferation in the absence of differentiation, also known as self-renewal. They retain the pluripotency of the cells in the early embryo when reintroduced into the blastocyst (Beddington and Robertson 1989) and can give rise to all cell types in the body (Keller 2005). Although some argue that ES cells do not occur in vivo as such, the in vitro differentiation of ES cells can be achieved by reproducing the developmental signaling pathways identified in vivo (Keller 2005). Thus, human ES cells have significant potential applications in tissue and cell engineering and as tools in drug discovery. Applications include the generation of blood cells for blood transfusions, replacement of damaged neurons in Parkinson disease, replenishment of insulin-secreting beta cells in diabetes mellitus, and bone formation

in osteoporosis. In addition, the newly discovered means of reprogramming postmitotic cells into induced pluripotent stem (iPS) cells (Takahashi and Yamanaka 2006; Takahashi et al. 2007; Yu et al. 2007) not only offers the possibility to generate disease-specific stem cells for derivation of novel treatment targets but also the production of patient-specific cells that will not cause immune rejection when transplanted back into the patient.

Classically, ES cells are derived and maintained in vitro using combinations of feeder cells, conditioned media, cytokines, growth factors, serum (primarily fetal bovine serum), and serum extracts as multifactorial stimulation of dedicated transcriptional circuitries that lead to the constant transcriptional activation of pluripotency-linked transcription factors such as Oct4, Sox2, and Nanog. Although the cytokine LIF (leukemia inhibitory factor) alone is insufficient to support self-renewal in serum- and feeder cell-free conditions, the activation via the LIF and gp130 receptors of at least four different downstream signal transduction pathways—JAK/STAT (Janus tyrosine kinase/signal transducer and activator of transcription), Ras/ERK1/2 (extracellular signal-related kinases), PI3K (phosphoinositide-3 kinase), and SFK (Src family kinase) pathway—are generally considered most critical for mouse ES (mES) cell maintenance (Anneren 2008; Burdon et al. 2002). Although human ES (hES) cells respond to LIF, the cytokine does not maintain their self-renewal capacity. Instead, hES cells require supplementation with fibroblast growth factor 2 (FGF2) and signaling via members of the FGF receptor (FGFR) tyrosine kinase family (Levenstein et al. 2006). Interestingly, recent breakthroughs have shown that extrinsic stimuli in many respects are dispensable for propagation and self-renewal of mES cells. For example, mES cell self-renewal can be maintained by the use of small-molecule inhibitors to suppress differentiation-inducing signaling from mitogen-activated protein kinase (MAPK) and glycogen synthase kinase 3 (GSK-3) (Ying et al. 2008).

Obviously, significant hurdles remain to be conquered before ES and iPS cells can be used in large-scale applications. Key issues, such as defining and re-creating a suitable culture environment to allow efficient differentiation of the cells, remain to be resolved before the stem cell regenerative medicine from scientific fiction becomes a medical reality.

The Effect of Altered HS in ES Cells

Over the years, the scientific community has focused more on describing the microenvironments, or so-called niches, surrounding stem cells. Numerous studies have shown that there is a constant signaling (e.g., by auto-, para-, and exocrine secreted factors) through cell-to-cell interactions between stem cells or with the neighboring tissue, as well as through interactions with the extracellular matrix that

promote either self-renewal or differentiation initiation. HSPGs are considered the most common and widely acting low-affinity receptors on the cell surface and play a central role in the reception and modulation of a wide range of growth factors, morphogens, and chemokines. Compared with a variety of murine tissues, mouse ES cells express fairly low-sulfated HS (Ledin et al. 2004; Johnson et al. 2007). However, a clear increase in the expression of HS biosynthetic enzymes, which results in increased HS chain sulfation, has been observed during ES cell differentiation (Johnson et al. 2007). Although *in vivo* studies have shown that disruption of HS biosynthesis, by mutating a good number of the genes involved in this process, clearly affects various stages during animal development, so far studies on ES cells mainly have been done using cell lines deficient in the *Ext* and *Ndst* enzymes. In addition, the role of HS in ES cell biology so far has been studied mainly using mouse ES cells, and the studies discussed in this review, if not stated otherwise, have been done in these cells.

Progression from self-renewal to differentiation and the subsequent lineage commitment is the main focus of most recent studies regarding the role and function of HS in ES cells. As stated above, HS sulfation is upregulated upon ES cell differentiation, which in turn alters the ability of the cells to interact with extrinsic factors. Numerous studies have identified various signaling networks, including those initiated by FGF, platelet-derived growth factor (PDGF), vascular endothelial growth factor (VEGF), Wnt, and bone morphogenetic protein (BMP) (all of which are modulated by HS interactions), involved in the regulation of cell fates during embryogenesis. Notably, these growth factors are also well established to affect ES cells, either to maintain self-renewal or to induce differentiation. Consequently, an impaired HS function inhibits the transition to differentiation probably by “numbing” the ES cells to their environment and thus keeping the cells in a naive ES cell state.

Studies Using *Ext1*^{-/-} ES Cells

ES cells lacking *Ext1* do not synthesize any HS and are therefore excellent tools for studying the role of HS in differentiation. Several studies have shown that these cells can form phenotypically correct colonies and express high levels of ES cell markers (Johnson et al. 2007; Kraushaar et al. 2010). However, Sasaki and coworkers (2008) show that RNAi silencing of *Ext1* leads to a reduced proliferation with subsequent spontaneous differentiation toward extra-embryonic endoderm. Maybe the low level of *Ext1* expressed by these cells is enough to support HS-mediated signaling.

In contrast, a total lack of *Ext1* prevents both spontaneous differentiation as well as directed neural differentiation (Johnson et al. 2007; Kraushaar et al. 2010; Holley et al. 2011). A recent study by Holley and coworkers (2011)

showed that *Ext1*^{-/-} cells do form embryonic bodies (EBs), an *in vitro* three-dimensional (3D) model for ES cell differentiation, with a subsequent increase in early mesodermal markers. However, these cells failed to terminally differentiate and entirely adopt full lineage restrictions. Notably, the addition of soluble heparin efficiently stimulated these cells to differentiate in a similar manner as their wild-type counterparts by overcoming insufficient signaling by FGF, BMP, and Wnt. In contrast, the addition of heparin to wild-type cells impaired differentiation, most likely by outcompeting cell surface HS for growth factor binding (Holley et al. 2011). Kraushaar and coworkers (2012) showed that *Ext1*^{-/-} ES cells were able to differentiate in serum-containing medium supplemented with high levels of FGF2 and exhibited morphological changes associated with differentiation, followed by downregulation of ES pluripotency genes as well as upregulation of endoderm-associated markers. Early markers for neuroectoderm were also detected, albeit at lower levels and at later stages compared with differentiating wild-type ES cells. This is in contrast to the study by Holley and colleagues (2011), who could not detect any mesoderm differentiation. Again, the addition of heparin fully restored differential efficacy, clearly showing the importance of HS in FGF-induced mesoderm differentiation. Similar results were obtained using BMP4, but not Wnt, to overcome the impaired differentiation potential of *Ext1*^{-/-} ES cells and induce mesoderm. Their results suggest that HS stabilizes BMP4 levels in the medium, thus maintaining and facilitating signaling during the differentiation process.

It was recently shown that *Ext1*^{-/-} EBs respond to VEGF stimulation and form angiogenic sprouts (Le Jan et al. 2012). Although the subsequent pericyte attachment was slightly defective, this finding suggests that early angiogenic sprouting and tube formation can occur in the absence of HS. Interestingly, the complete loss of HS in the *Ext1*^{-/-} cells led to increased chondroitin sulfate (CS) biosynthesis, which most likely can compensate for the lack of HS. Although CS was shown to partly substitute for HS deficiency-induced aberrant PDGFB and TGF signaling, both known regulators of pericyte differentiation and attachment (Hirschi et al. 1998; Hellstrom et al. 1999), the signaling input was not sufficient to warrant a complete rescue.

Studies Using *Ndst1/2*^{-/-} ES Cells

ES cells devoid of both *Ndst1* and *Ndst2* also have been used to study the role of HS in differentiation. These cells produce HS that completely lacks *N*-sulfation. Apart from a low degree of 6-*O*-sulfation, the HS was unmodified (Holmborn et al. 2004). Similar to *Ext1*^{-/-} ES cells, *Ndst1/2*^{-/-} ES cells also maintained their self-renewal and pluripotency capacity under normal ES cell culture conditions (Holmborn et al. 2004; Lanner et al. 2010). In contrast, angiogenesis was

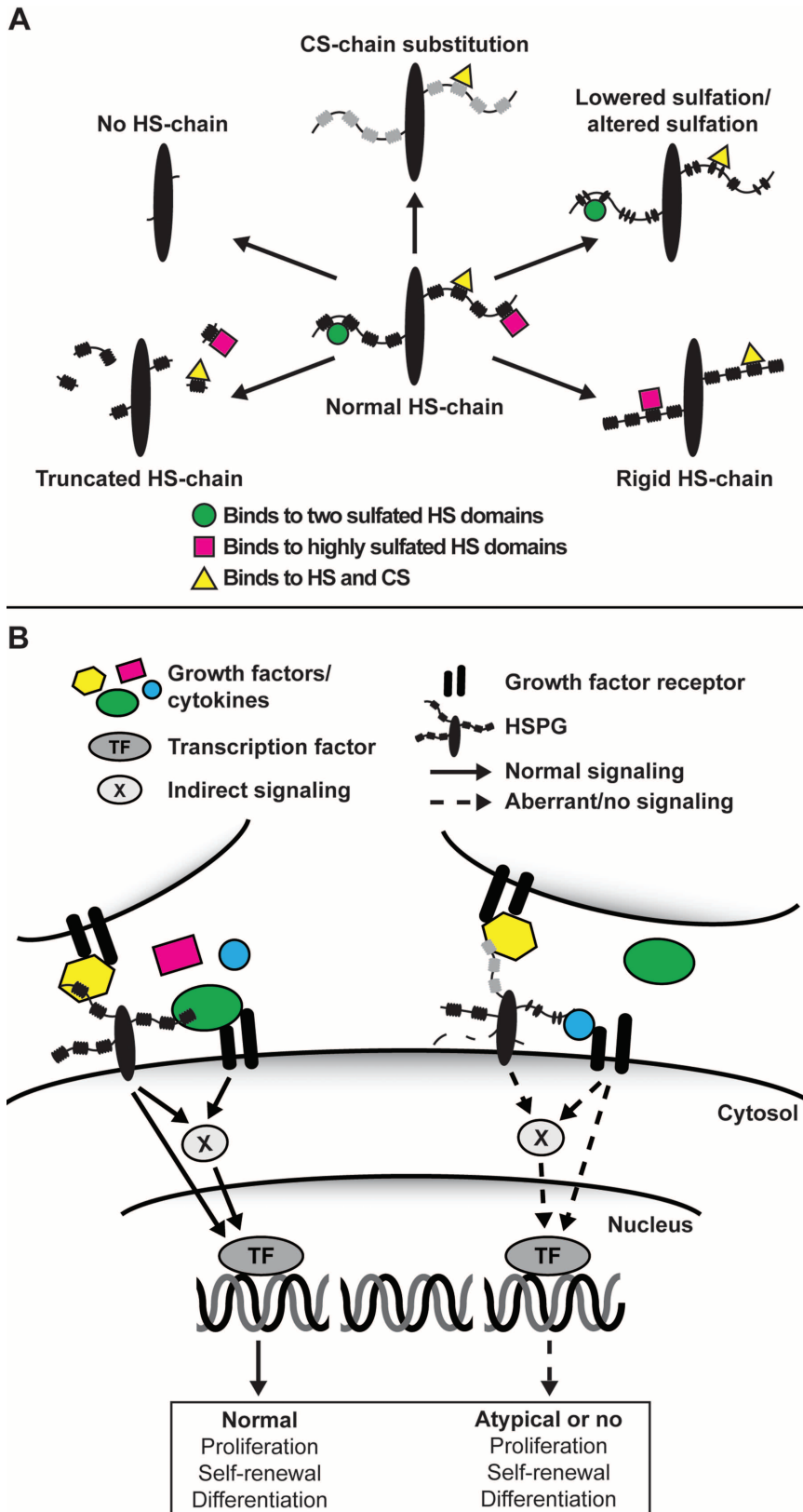


Figure 1. (A) Hypothetical heparan sulfate (HS) chain structural consequences and effects on ligand interaction upon various mutations in the HS biosynthesis machinery. Depending on the enzyme being modified, various degrees of structural defects in the HS chain will occur ranging from a complete lack of HS chains (*Ext1^{-/-}*) and exhibiting different degrees of sulfation (various HS sulfotransferase knockouts) to chain length (*Hpse^{-/-}*) and HS chain rigidity (*Glce^{-/-}*). The loss of HS in the *Ext1^{-/-}* cells can also lead to a compensatory increase in chondroitin sulfate (CS) biosynthesis. (B) In wild-type embryonic stem (ES) cells (left), heparan sulfate proteoglycans (HSPGs) located on the same or an adjacent cell function as co-receptors for growth factors and their receptors or, upon ligand binding, induce downstream signal transduction on their own. Although not shown in the figure, HSPGs also facilitate cell adhesion to the extracellular matrix and, when secreted, are involved in the formation of the extracellular matrices that, for example, sequester growth factors and morphogens. However, in the various mutant ES cells (right), in which the HS synthesis and modification machinery have been targeted, an altered HS chain will change the extracellular information input received by the cell, subsequently affecting processes such as self-renewal capacity and pluripotency.

severely delayed in *Ndst1/2*^{-/-} EB cells, and pericyte formation completely arrested (Le Jan et al. 2012). In comparison to *Ext1*^{-/-} ES cells, virtually no CS could be detected in *Ndst1/2*^{-/-} ES cells, which further supports a functional overlap between HS and CS during angiogenesis.

Although it has been suggested that *Ndst1/2*^{-/-} ES cells generally fail to properly differentiate upon EB formation (Lanner et al. 2010), Forsberg and coworkers show that a stepwise differentiation approach allowing for the formation of intermediate progenitors renders the formation of mesoderm, as well as primitive endo- and ectoderm (Forsberg M et al. 2012). In their study, the cells formed osteoblasts, although with a lower efficacy compared with wild-type ES cells. However, no formation of adipocytes, which is usually found in concert with osteoblasts under in vitro differentiation, was detected. Under neural-inducing conditions, the *Ndst1/2*^{-/-} ES cells adapted a primitive ectoderm-like state expressing FGF5, but the cells were blocked and could not proceed to neural differentiation. Similar results were obtained when looking at endoderm differentiation where the cells seemed to halt in a Gata4-expressing primitive state. The addition of low levels of heparin induced neural differentiation, presumably by enhancing FGF4 signaling.

Future Studies and Available ES Cell Models

Several key ES cell transcription factors have recently been reported to show diverse but dynamic expression during normal ES cell maintenance in serum (Wray et al. 2010). Functionally, this heterogeneous mixture of cells has distinct differential potential and capacity to form chimeras upon blastocyst injection. When shielding the cultures from exogenous differentiation-inducing serum factors with small-molecule kinase inhibitors, the transcription factor levels are stably high and fluctuate less. We and others have observed that ES cells grown with MAPK and GSK-3 inhibitors form colonies that phenotypically appear more compact than the cells cultured under normal conditions and show very little spontaneous differentiation (C. Tamm, S. P. Galitó, and C. Annerén unpublished observations; Marks et al. 2012). Interestingly, similar phenotypes were found for *Ext1*^{-/-} ES cells, which lack cell surface HS and thus repress HS-regulated extracellular signaling (Holley et al. 2011). Hence, although not fully investigated, it has been suggested that the pluripotency transcription factor network is inherently stable but is easily altered by the vast amount of extrinsic signaling factors present in serum (Wray et al. 2010). As described above, many of these factors have been shown to rely on HS to function properly, and it is therefore clear that any alterations in the HS chains will have a wide impact on the extracellular information to be received by a cell.

Altogether, the results underscore the need for functional HS for inclusive and committed differentiation. As stated

above, the functions of most genes involved in HS biosynthesis have been studied for animal development through targeted mutation of the genes in mice. So far, only a few ES cell lines (i.e., *Ext1/2* and *Ndst1/2*) derived from the mutant mice have been investigated for the functional roles of HS in ES cell biology, and the evidence clearly shows a fundamental role in the mere existence and overall design of the sulfation pattern of HS chains. Further studies by application of ES cells derived from mutant mice will reveal how more subtle and defined changes, brought on by mutations of downstream regulators involved in fine-tuning the definitive HS chains' architecture, will affect the ES cell self-renewal and differentiation capacity.

To facilitate the investigation for the roles of HS in ES cell differentiation, apart from the *Ext1*^{-/-} and *Ndst1/2*^{-/-} cells discussed above, we have derived *Ndst1*^{-/-}, *Ndst2*^{-/-}, *Glyce*^{-/-}, and *Hpse*^{-/-} ES cells from the corresponding mouse strains generated in our lab. In fact, ES cell lines are being established from most mutant mice strains known to the HS field. Interestingly, our mutant cells, which all express HS with various flaws, demonstrate dissimilar differentiation capacity and lineage commitment. This unique panel of mutant ES cells will allow us to study and compare the functional consequences of completely lacking HS (*Ext1*^{-/-}) or exhibiting various degrees of structural defects in HS (Figure 1). Elimination/neutralization of extrinsic signaling has revolutionized in vitro maintenance of ES cells; however, the role and function of extracellular signaling in ES cells and early embryogenesis, as well as how it is controlled by HSPGs, still remain of great interest for developmental studies and, maybe more important, for the regulation and harnessing of ES pluripotency and differentiation.

Finally, although the role of HS in ES cell biology mainly has been studied in mES cells, a recent study by Levenstein et al. (2008) showed that mouse embryonic fibroblast-secreted HSPGs stabilize basic fibroblast growth factor (FGF2) and directly mediate binding of FGF2 to hES cells, and HS removal impaired proliferation. No doubt, HS plays an essential role also in hES cell biology, fine-tuning the extracellular input received by the cells and thus directing the interaction with adjacent cells and matrices in its surrounding niche.

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