

Chondroitin Sulfate E in Signaling of the Growth Factor Midkine

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Author's Profile



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I Introduction

Midkine is a 13-kDa protein that enhances growth, migration and survival of various cells. Midkine has about 50% sequence identity with pleiotrophin / HB-GAM, but is not related to other growth factors or chemokines^{11,12}.

Midkine is essentially composed of two domains tightly held by disulfide bridges, and each domain has three antiparallel β -sheets¹³. Two clusters of basic amino acids are present in the more C-terminally located domain, and serve as binding sites for glycosaminoglycans¹³.

Midkine expression is strong in midgestation embryogenesis, while it is detected only in restricted regions in the adult tissue^{12,14}. However, strong expression of midkine is found in many human carcinomas and also upon inflammation and repair^{12, 15-17}. Midkine antisense oligo DNA inhibits growth of colorectal carcinoma cells in nude mice¹⁸. Therefore, midkine is considered to contribute to the malignant phenotype of tumor cells, probably by enhancing their growth, survival and invasion. Midkine also promotes migration of inflammatory cells. In mice deficient in the midkine gene, neointima formation upon vascular injury¹⁶ and nephritis after ischemic injury¹⁷ are significantly suppressed. Decreased migration of inflammatory cells in the deficient mice is considered to be the reason for these phenomena. On the other hand, survival-promoting activity of midkine is promising in prevention of neuronal degeneration. Retinal degeneration caused by exposure to constant light is prevented by midkine¹⁹. Delayed neuronal death of hippocampal neurons after ischemic injury is also inhibited by midkine²⁰. Furthermore, survival of *in vitro* developed preimplantation embryos is enhanced by midkine²¹. Therefore, midkine is an important factor both as a target of molecular therapy and as a potential drug to treat degenerative diseases.

Midkine receptor is a molecular complex composed of both transmembrane proteins and transmembrane proteoglycans. A transmembrane protein, low density lipoprotein, receptor-related protein (LRP) plays important roles²². In addition, ALK tyrosine kinase has recently been reported to participate in midkine signaling²³. As proteoglycans, both syndecans and a chondroitin sulfate proteoglycan are implicated to play a role in midkine signaling^{6,12}. The intracellular signaling system of midkine includes PI3 kinase and ERK^{12, 24}.

III *Recognition of chondroitin sulfate E by midkine*

The interaction of chondroitin sulfate E with midkine was revealed by two independent approaches. In one side, midkine was found to bind strongly to a chondroitin sulfate proteoglycan, receptor-type protein tyrosine phosphatase ξ (PTP ξ)⁶, (Fig. 2). The following evidence indicates that PTP ξ is a receptor of midkine in midkine-dependent migration of embryonic neurons: 1) digestion of neurons with chondroitinase ABC eliminates the midkine action; 2) antibody to the extracellular portion of PTP ξ inhibits midkine action; 3) PTP ξ binding activity of midkine mutants correlates well with that of migration-promoting activity. PTP ξ is also involved in midkine-dependent migration of osteoblast-like cells²⁵ and midkine-dependent survival of embryonic neurons²⁶.

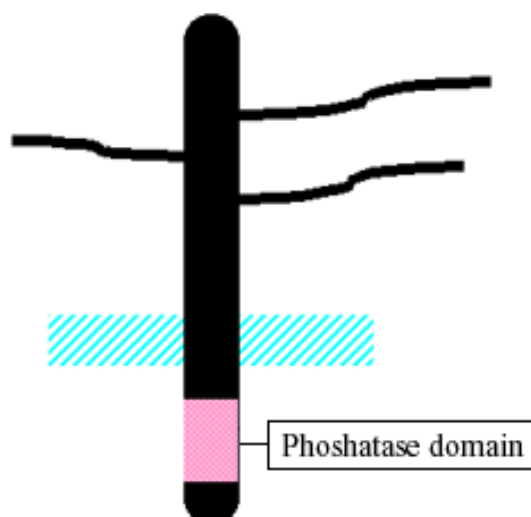


Fig. 2. Receptor-type protein tyrosine phosphatase ζ (PTP ζ)

Midkine binds to the chondroitin sulfate portion of PTP ζ with high affinity and to the protein portion with low affinity⁶. To discover what kind of chondroitin sulfate midkine binds strongly, inhibitory effects of various chondroitin sulfates to midkine-dependent migration were examined. Chondroitin sulfate E was found to inhibit cell migration most strongly among the various chondroitin sulfates investigated.

We also found that midkine binds to a pericellular chondroitin sulfate proteoglycan, PG-M / versican with high affinity²⁷. The binding is inhibited by heparin, chondroitin sulfates D and E. In the chondroitin sulfate from PG-M, small amounts of D and E units have been detected.

The other line of work was started when chondroitin sulfate E was found to inhibit neurite outgrowth on dishes coated with midkine⁷. Other chondroitin sulfates showed no inhibitory effects in this system²⁸. We compared the binding affinity of chondroitin sulfate E to midkine with that of heparin to midkine, since heparin is known to bind strongly to midkine. The methods of analysis were surface plasmon resonance⁷ and midkine Sepharose-affinity chromatography⁸, in addition to inhibition of neurite outgrowth^{7,28}. Consequently, we have reached to the conclusion that chondroitin sulfate E binds to midkine with an intensity similar to that of heparin.

We also questioned whether midkine responding cells have chondroitin sulfate rich in E units. Analysis of chondroitin sulfate synthesized by the brain from day-13 mouse embryos has revealed that a fraction of chondroitin sulfate indeed binds strongly to midkine and contains large amounts of E units⁸.

The other question is whether modification of chondroitin sulfate E, such as 3-*O*-sulfate of gluconic acid²⁹, is involved in strong binding to midkine. We utilized artificial chondroitin sulfate E, which was synthesized from chondroitin 4-sulfate using purified or recombinant chondroitin 4-sulfate 6-sulfotransferase³⁰. The artificial chondroitin sulfate E was found to bind to midkine strongly, indicating that modifications such as 3-*O*-sulfation are not essential for strong binding⁸. We

also noted that a cluster of E units is required for strong binding.

In case of the binding of midkine to heparin or heparan sulfate, all three sulfate groups in the heparin trisulfated unit are required for strong binding³¹. It is interesting that the number of sulfate residues per disaccharide unit required for strong binding to midkine is less in chondroitin sulfate E than in heparin (Fig. 3).

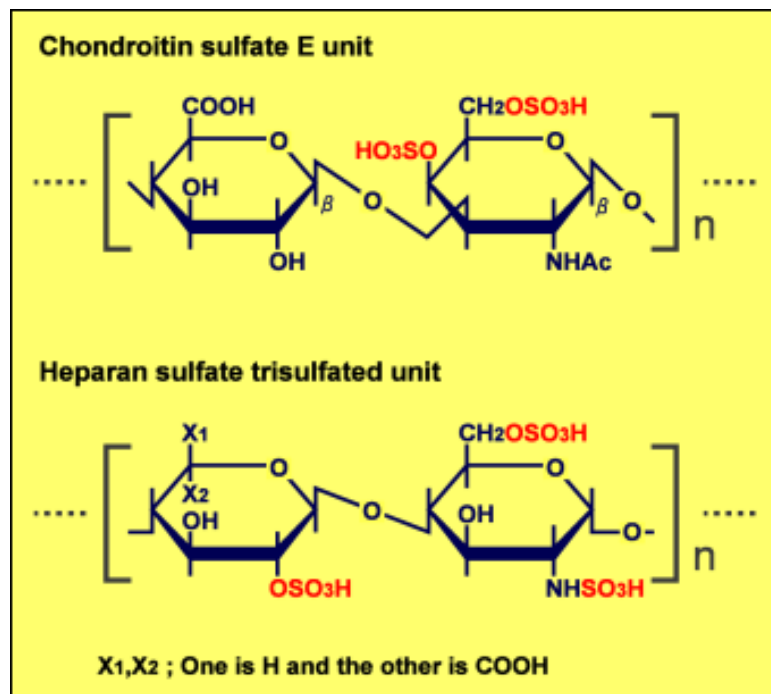


Fig. 3. Comparison of two structures with strong binding capability to midkine

Clusters of the E units are not widely distributed in the chondroitin sulfate chain. Therefore this structure can serve as a specific signal for binding to midkine as well as to other molecules.

IV Comments

Our studies established that clusters of chondroitin sulfate E units bind strongly to midkine and are involved in the signaling of midkine. However, the exact role of the binding in midkine signaling remains to be clarified by further studies. Most probably, midkine binding to a chondroitin sulfate proteoglycan, PTP ζ , induces dimerization of the molecule, leading to inactivation of the phosphatase activity. Since the midkine receptor is a molecular complex, the possibility that the binding to the chondroitin sulfate portion alters the relationship of the chondroitin sulfate proteoglycan to other components of the complex cannot be excluded.

The finding that chemokines⁹ and selectins⁹ also bind to chondroitin sulfate E indicates that a significant number of recognition molecules bind to oversulfated chondroitin sulfates. The classification of heparin-binding recognition molecules to those specific to heparin and heparan sulfate and those with dual recognition capability to heparin/heparan sulfate and oversulfated chondroitin sulfates is an urgent task. It has recently been reported that FGF-16, FGF-18 and HB-EGF also

strongly bind to chondroitin sulfate E³².

Equally important is the establishment of the *in vivo* significance of recognition of oversulfated chondroitin sulfates. Chondroitin sulfates D and E are synthesized as outlined in Fig. 4. Indeed, knockout of chondroitin 6-sulfotransferase resulted in the absence of chondroitin sulfate D in the brain³³. So far, no abnormalities have been found in the brain of the deficient mice. Most probably, the chondroitin sulfate E structure, which remained in the brain of the knockout mice compensated for the loss of the D structure.

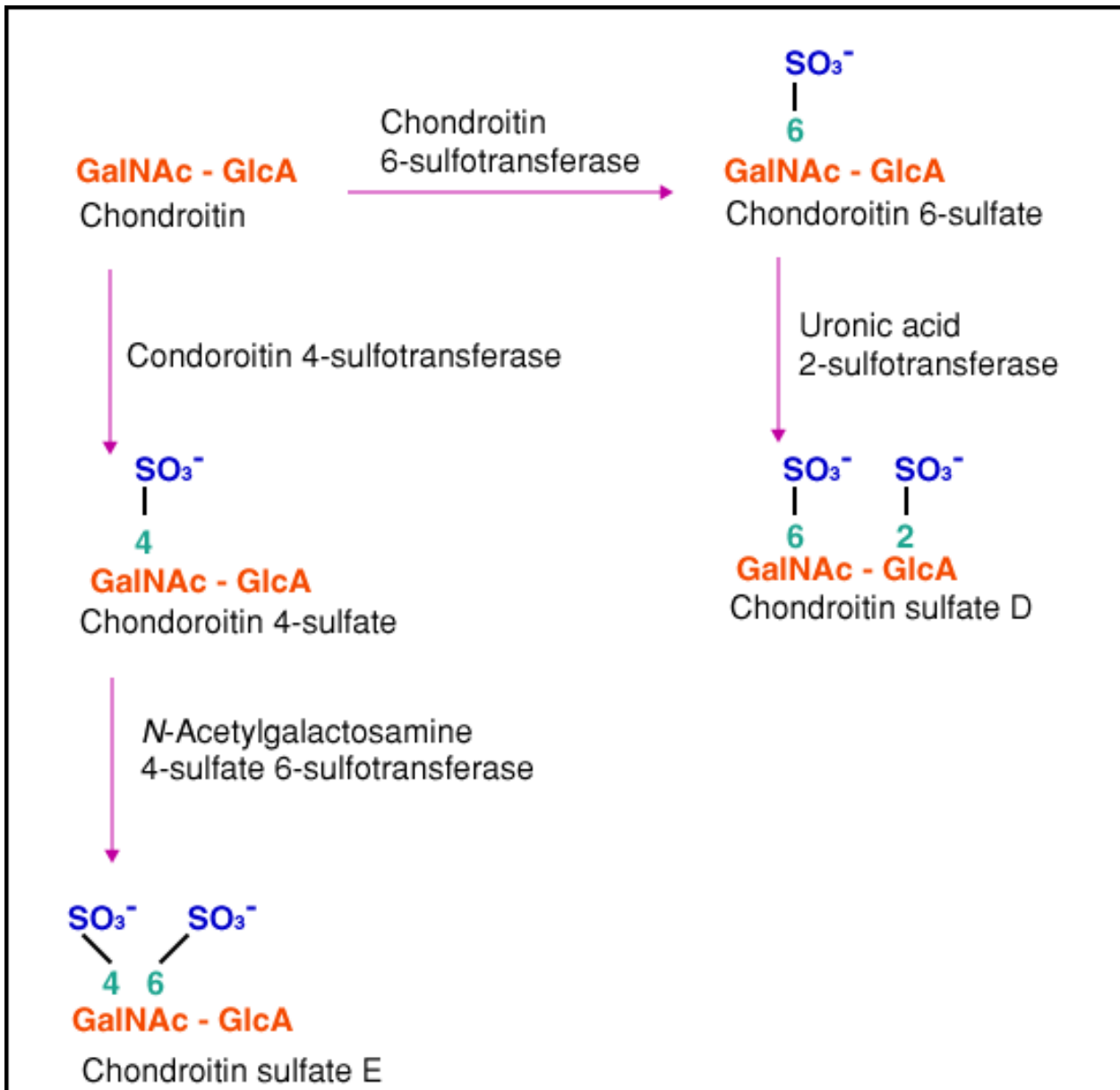


Fig. 4 Biosynthetic pathway for the formation of oversulfated chondroitin sulfates

Keyword

Chondroitin sulfates

The fundamental structure of chondroitin sulfate is chondroitin, whose repeating unit is a disaccharide composed of *N*-acetylgalactosamine and glucuronic acid. Sulfation at different hydroxyl groups yields different chondroitin sulfates. Epimerization of the C-5 hydroxyl group in glucuronic acid of chondroitin 4-sulfate yields dermatan sulfate. Microheterogeneity in chondroitin sulfate structures is an important characteristic. As an example, oversulfation occurs in varying degrees in chondroitin sulfates from different sources. Structural analysis of chondroitin sulfates depends on chondroitinases, a method developed by Prof. S. Suzuki. After chondroitinase ABC digestion, chondroitin sulfates are depolymerized to unsaturated disaccharides, which are separated by HPLC and quantitated.

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