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Title: Novel Cleavage-Dependent Interaction of Proteoglycan Receptors with

Neuropilin-1

Abstract

Receptor tyrosine phosphatases (RPTPs) play a critical role in regulating signaling events in coordination with tyrosine kinases. RPTPs not only regulate tyrosine phosphorylation of intracellular proteins at the intersection of many signaling pathways but also play key roles in transmitting extracellular signals into the cytosol. The type IIa subfamily of RPTPs (RPTP σ , RPTP δ , and RPTP-LAR) serves as functional receptors for proteoglycan glycosaminoglycans (GAGs).

In this study, we compared the characteristics of RPTP σ and RPTP δ . We found that the extracellular domains (ECDs) of both RPTP σ and RPTP δ distinguished the extent of sulfation on GAG chains based on solid-phase binding assays and Glycan Microarray. Notably, we observed striking differences: (1) RPTP δ possessed a single GAG binding site, whereas RPTP σ exhibited an additional heparin binding site in the juxtamembrane domain; (2) complete cleavage was observed within the ECD of RPTP δ , whereas only partial cleavage occurred in the RPTP σ ECD; and (3) the novel heparin binding site and the potential cleavage site on RPTP σ overlapped.

Identification of the exact cleavage sites within the ECD of RPTP σ and RPTP δ in this study led us to conclude that furin was the responsible enzyme. This conclusion is based on the in vitro cleavage of recombinant RPTP σ and RPTP δ by the recombinant furin (rFurin), and the inhibition of the cleavage by the furin inhibitor in HEK293 cells and no cleavage in the furin-deficient cells. Furthermore, exogenous heparin partially blocked the cleavage of RPTP σ by rFurin, suggesting that heparan sulfate acts as a potential regulator of RPTP σ ECD cleavage.

We discovered that the cleavage products of RPTP σ and RPTP δ bound to neuropilin-1 (NRP1) through their C-terminal end, known as the "C-end Rule [CendR] motif". Solid-phase binding assays revealed that both cleavage products exhibited very high-affinity binding (K_d=~10⁻⁹M) to NRP1, with 2 binding sites on RPTP δ and a single binding site on RPTP σ .

Since NRP1 plays a role as a co-receptor in multiple signaling pathways, including VEGF-A and TGF β , we sought to identify the pathways that are potentially affected by the cleavage products of RPTP σ and RPTP δ . Recombinant ECDs of RPTP σ and RPTP δ reduced the proliferation of human umbilical vein endothelial cells (HUVEC) induced by VEGF-A in a CendR motif-dependent manner, showing that the recombinant fragments were biologically active. Further, the exogenous ECD of RPTP δ reduced TGF β -induced transcription in a CendR motif-dependent manner. In contrast, RPTP σ reduced TGF β -induced transcription independently of CendR.

While the ECD cleavage of type IIa RPTP has long been known but often overlooked, our findings shed light on the physiological relevance of this process.