

Brief Note on Clinical Significance of Plasma ST6GAL1 as a Biomarker

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DESCRIPTION

ST6GAL1, also known as ST6 beta-galactoside alpha-2,6-sialyltransferase 1, is an enzyme that plays a vital role in the process of glycosylation, specifically sialylation. Sialylation is a post-translational modification of glycoproteins and glycolipids, where sialic acid residues are added to the terminal positions of carbohydrate chains (glycans). This modification has significant implications in various biological processes, including cell-cell interactions, immune responses, and disease states.

Sialylation is an important type of glycosylation, which involves the addition of sialic acid residues to glycoproteins and glycolipids. This modification can significantly influence the function and properties of these molecules. Sialic acids on cell surface glycans play a role in various cellular processes, including cell-cell interactions, adhesion, and signaling. They are involved in modulating immune responses and protecting cells from recognition by the immune system.

Role of ST6GAL1 in plasma

ST6GAL1, is not typically found in significant quantities in the bloodstream or plasma under normal physiological conditions. Instead, it is primarily an intracellular enzyme found in various tissues and cells where it plays a role in sialylation of glycoproteins and glycolipids. Sialyltransferases like ST6GAL1 are primarily involved in the modification of glycoproteins and glycolipids on the cell surface and within intracellular compartments. This modification is crucial for various cellular processes, including cell adhesion, signaling, and immune recognition. Sialic acid residues added by sialyltransferases are generally found on the cell surface, where they influence the properties and functions of cell membrane molecules.

While ST6GAL1 itself is not typically found in plasma, the products of its enzymatic activity, such as sialylated glycoproteins and glycolipids, can be present in the bloodstream. These sialylated molecules have important roles in various physiological and pathological processes, including immune responses, cell signaling, and disease states. However, the presence of ST6GAL1

or its activity in the bloodstream is typically a result of cellular processes and not a direct secretion of the enzyme into the plasma. The measurement of specific sialylated molecules in the bloodstream may provide insights into various health conditions, but the enzyme ST6GAL1 itself is not a common plasma constituent.

Clinical significance

Cancer: ST6GAL1 is often overexpressed in various types of cancer, including breast, colon, and pancreatic cancer. This overexpression leads to increased sialylation of cell surface molecules, which can have several clinical consequences

- Immune evasion sialylation can mask tumor antigens and inhibit immune cell recognition, allowing cancer cells to evade immune surveillance.
- Metastasis altered sialylation can enhance the ability of cancer cells to invade nearby tissues and metastasize to distant organs.
- Prognostic marker elevated ST6GAL1 expression has been associated with a poorer prognosis in cancer patients, making it a potential biomarker for disease progression and survival.

Autoimmune diseases: Abnormal sialylation patterns, potentially driven by ST6GAL1 activity, are observed in autoimmune diseases such as rheumatoid arthritis and systemic lupus erythematosus. These changes in sialylation can affect immune responses and contribute to the pathogenesis of autoimmune disorders.

Neurodegenerative diseases: Sialic acid modifications in the nervous system are essential for proper neuronal development and function. Aberrant sialylation, possibly related to ST6GAL1, has been implicated in neurodegenerative diseases like Alzheimer's and Parkinson's disease. Understanding these glycosylation changes may provide insights into disease mechanisms and potential therapeutic targets.

Cardiovascular Diseases: Altered sialylation patterns in glycoproteins, which may involve the activity of ST6GAL1, have been linked to cardiovascular diseases, such as atherosclerosis and coronary artery disease. These changes can affect the

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properties of proteins involved in blood clotting and inflammation.

Therapeutic development: Researchers are exploring the potential of ST6GAL1 as a target for therapeutic intervention in cancer and other diseases. Inhibiting the enzyme's activity may have clinical benefits, such as reducing immune evasion in cancer or modulating immune responses in autoimmune disorders.

Diagnostic and prognostic biomarker: The expression levels of ST6GAL1 or the sialylation patterns of specific glycoproteins may serve as diagnostic or prognostic biomarkers in various diseases. Detecting abnormal sialylation could provide valuable information about disease progression and response to treatment.

CONCLUSION

Understanding the clinical implications of ST6GAL1 and its role in glycosylation is important for both diagnostic and therapeutic advancements in these conditions. ST6GAL1 is an enzyme involved in sialylation, a type of glycosylation that has far-reaching implications in various physiological and pathological processes. It is particularly important in cell surface interactions, immune responses, and the development and progression of certain diseases. Its ability to modify glycoproteins and glycolipids with sialic acid has far-reaching implications in various aspects of biology, from cellular interactions to disease pathogenesis, making it an important target for research in both basic science and therapeutic development.