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Patent Search

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Abstract:

The method for the development of an AI-enabled approach for identifying patients with breast cancer who would benefit from GPR54 receptor-targeted therapy is presented in this article. Potential as a potential therapeutic target, the GPR54 receptor is implicated in tumour growth and metastasis suppression. Through the integration of multi-modal data, such as imaging features, clinical biomarkers, and gene expression profiles, our machine learning model correctly stratifies patients according to GPR54 receptor status. Through the use of deep neural networks, ensemble learning, and feature selection, the system exhibits strong predictive performance, making it possible to classify patients with increased sensitivity and specificity. The suggested method speeds up clinical decision-making while also improving individualized treatment planning. This AI-powered approach provides a major breakthrough in precision oncology, establishing the foundation for receptor-specific treatments that have the potential to significantly enhance the management of breast cancer. FIG.1

Complete Specification

Description:[0002] The breast cancer is still one of the most common and fatal tumors that affect women globally. The diverse nature of breast cancer makes it difficult to choose the best individualized treatments, even with major advancements in early diagnosis, diagnostic tools, and therapeutic approaches. The intricate molecular landscape of cancers is sometimes missed by conventional diagnostic methods, which results in less-than-ideal therapy choices. Precision medicine has become a game-changer in this regard, customizing treatment plans according to the molecular traits of certain cancers.

[0003] The G-protein-coupled receptor 54 (GPR54), commonly referred to as the kisspeptin receptor, is one promising biomarker being studied for targeted therapy. Through pathways linked to cell migration, invasion, and adhesion, GPR54 plays a crucial role in controlling the spread of cancer cells. Research has shown that in a number of malignancies, including breast cancer, tumor growth is associated with deregulation of the kisspeptin-GPR54 signaling axis. Since different subtypes of breast cancer express the receptor differently, it may be a good target for biomarker-driven therapy. However, there is an increasing need for sophisticated computational algorithms that can precisely identify patients who can benefit from GPR54-targeted medicines because of the complexity of tumor heterogeneity and the shortcomings of traditional biomarker identification techniques. Oncology could undergo a significant transformation thanks to artificial intelligence (AI), especially machine learning (ML) and deep learning (DL) techniques, which make it possible to analyze large, complicated datasets including genomic sequences, histopathological pictures, and patient clinical data. AI models have recently been created to recognize molecular subtypes, forecast the results of treatments, and spot minute patterns that would be missed by human order to help stratify patients with breast cancer for receptor-specific treatments, artificial intelligence (AI) can assist in identifying the expression patterns and signaling pathways linked to GPR54 by combining multi-omics data, such as transcriptomics, proteomics, and genomics.

[0004] Molecular biology and artificial intelligence can be combined to create predictive models that can identify patients who are likely to express GPR54 and receive

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