

عنوان مقاله:

MiR-14b and let-Ya as Non-invasive Diagnostic Biomarkers of Alzheimer's Disease Using an Artificial Neural Network

محل انتشار: مجله بیوشیمی پزشکی, دوره 11, شماره 2 (سال: 1402)

تعداد صفحات اصل مقاله: 8

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خلاصه مقاله:

Background : Gaining insight into the underlying molecular mechanisms of Alzheimer's disease (AD) is crucial. Objectives : This study aimed to employ a systems biology approach to identify new non-invasive diagnostic biomarkers for AD. Methods : Gene expression data series GSE $177 \cdot 57$ and microRNA (miRNA) expression data series GSE $177 \cdot 57$ and microRNA (miRNA) expression data series GSE $177 \cdot 57$ and differentially expressed from the Gene Expression Omnibus database. The Limma package under R software was used to assess differentially expressed miRNAs and differentially expressed genes (DEGs). Afterward the protein-protein interaction (PPI) network was constructed by the STRING software and evaluated with Cytoscape software. The multilayer perceptron neural network (MLP-NN), a widely used artificial neural network (ANN), was employed to classify two groups. Results : A total of 174A DEGs were identified in AD patients compared to the control group, and 11 differentially expressed miRNAs were found in patients with mild cognitive impairment (MCI) in comparison to the control group. The results revealed that EGFR, identified as a hub gene, was targeted by miR $-1\Delta b-Tp$ and let $-Ya-\Delta p$, while TLRF, another hub gene, was targeted by miR $-1\Delta b-Tp$. The MLP-NN constructed using both hsa-let $-Ya-\Delta p$ and hsa-miR $-1\Delta b-Tp$, by targeting EGFR and TLRF, and let $-Ya-\Delta p$, by targeting EGFR, may play a significant role in AD. Additionally, the constructed ANN utilizing the expression levels of plasma miR $-1\Delta b-Tp$ and let $-Ya-\Delta p$ could .serve as a potential non-invasive diagnostic tool with high sensitivity for AD detection

كلمات كليدى:

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