

Cure for Primary Dopaminergic Midbrain Neurons by GPR139 Agonists using Structure based Virtual Screening and Molecular Dynamics Simulation

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Abstract

In various organisms, GPCR's are the mostly known cell surface receptors playing vital role among many interacting molecule during signal transduction and metabolic pathways. These are the super largest family which delivers signals from extracellular space to intracellular environment. GPCR'S having structural characteristics in terms of possessing seven transmembrane helices with distinct length of intra and extracellular loops. At both the loops varying terminal are found which usually distinguish the ligand role during the signal transduction pathways. GPCR's are involved in all regulating activities whether immune related or at physiological level in an organism. Through the human genome sequence project it has been reported that around 800 GPCR's sequence exists in human. These GPCR's found most effective physiological role in all peripheral organs and as well as in central nervous system. These also act as a multiple disease target. GPCR's mediate signals from the extracellular environment and then respond toward it to result in effective cellular response. Various ligands (outside signals) such as smaller interacting molecules, peptides (opoids, angiotensins, bradykinins, omatostatin, melanocortin), fatty acids, nitrogeneous substances, photons, proteins, amino acids, carbohydrates, ions and energy variable molecules harbour GPCR's. And then these ligands acknowledge endogeneous signals from the outer space of the cell. Due to the presence of different GPCR's it is not easy to identify the concerned natural ligand of it. So for further understanding of GPCR's, particular identification of endogeneous ligands is done. After knowing the ligand the particular GPCR is being drug targeted for the treatment of disease as they are 40-50% drug effective. GPCR share a common structural feature i.e. the presence of seven membrane spanning helices sharing three each intracellular and extracellular loops with an

intracellular carboxyl terminal and an extracellular amino terminal. Various differences in structure of GPCR can occur during the ligand binding, G protein coupling and interaction with other proteins. In Rhodopsin class of GPCR the basic structure of GPCR is verified by X-ray crystallography. Ligands acts as an extracellular stimulus for GPCR's and then brings physiological changes in organisms. GPR139 found most effective physiological role in all peripheral organs and as well as in central nervous system. In this study GPR139 targeted by virtually screened compounds which inhibit the Dopaminergic Midbrain Neurons. Molecular Dynamics simulation was performed for complex structure of GPR139 to validate binding affinity of compounds, and investigate active site fluctuation of complex structure.

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