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COMPUTATIONAL STUDY ON GPCR – G PROTEIN INTERFACES

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Recent studies show growing interest in targeting the cytoplasmatic interface of GPCRs with drugs [1,2]. New compounds, both petides and small molecules, prove promising pharmacological effects, however the detailed mechanism of interactions with intracellular interface of GPCRs has not been revealed yet.

Since the publication of the β2 Adrenergic receptor in complex with G_s protein is available [3], the aim of this study is to provide an information about other types of GPCRs interacting with G_i (Muscarinic M2 receptor and 5-HT_{1B} receptor) and G_o (5-HT_{2B} receptor) through bioinformatic analysis and molecular dynamics simulations.

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References:

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