

Computational Analysis of nAChR α 4 and β 2 Subunit Stability and NMR Study of Protein Anesthetic Interaction

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Abstract: Because the α 4 and β 2 subunits of the transmembrane domain of nAChRs are naturally unstable in solution suitable for NMR experimentation and structural determination, mutation of the subunit sequences has been performed to lower subunit pI. However, as α 4 stability is much greater than β 2 stability, further mutation of the β 2 sequence at key residues has been attempted to increase β 2 stability. Computer modeling and simulation of the α 4 and β 2 subunits provide a basis for assessing the mutant subunit stability. NMR experiments run both with and without anesthetic were performed to provide insight as to which specific residues within the α 4 subunit interact with anesthetic based on observed differences in chemical shifts.