Supplementary Data

Identification of Putative Molecular Imaging Probes for BACE-1 by Accounting for Protein Flexibility in Virtual Screening

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Excel Sheet 1, 143*283 binding affinity matrix for virtual screening of the focused ligand library against the BACE-1 ensemble structures; Sheet 2, the ranked list of ligands in each category according to the best accommodating affinity with BACE-1 receptors.

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