

Supplementary data

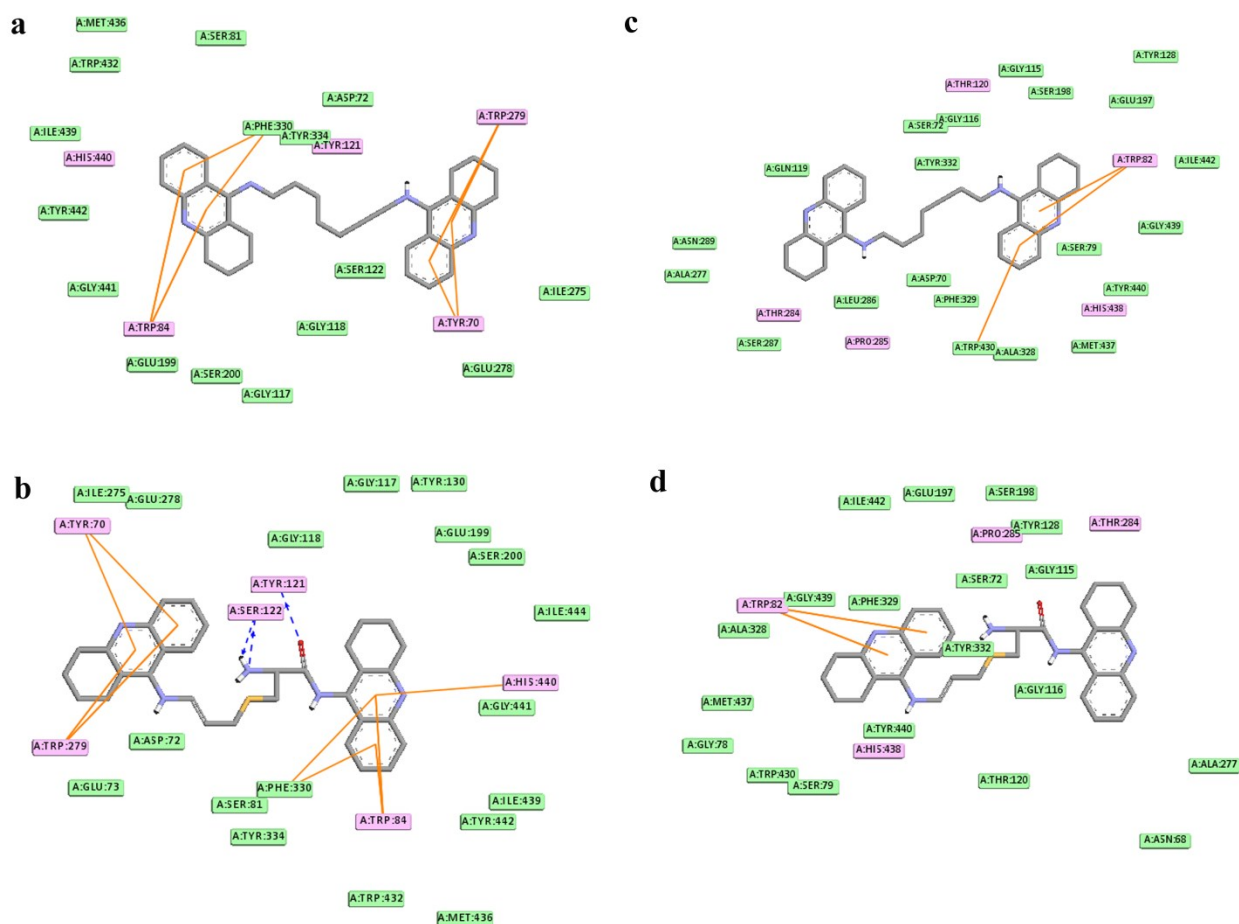


Figure S1: Two-dimensional representations of the observed interactions between both enzymes and both ligands after docking procedure; residues in a green frame: van der Waals interaction, residues in a magenta frame: electrostatic interaction, Orange lines: π - π interaction, blue dashed line with arrow at H-bond acceptor: hydrogen bond with amino acid side-chain; (a) AChE-bis(7)tacrine, (b) AChE-bis(7)tacrine(7)-SAC, (c) AChE-bis(7)tacrine, (d) BChE-bis(7)tacrine-SAC.

