

Free-energy calculations on potential A₃ adenosine receptors inhibitors

Zuzana Jandova,¹ Willem Jaspers,² Eddy Sotelo,³ Hugo Gutiérrez-de-Terán,² Chris Oostenbrink¹

¹ Institute of Molecular Modeling and Simulation, University of Natural Resources and Life Sciences, Vienna

² Department of Cell and Molecular Biology, Uppsala University, Uppsala SE-75124

³ Centro Singular de Investigación en Química Biolóxica e Materiais Moleculares (CIQUS) and Departamento de Química Orgánica, Facultade de Farmacia, Universidade de Santiago de Compostela, 15782, Santiago de Compostela, Spain

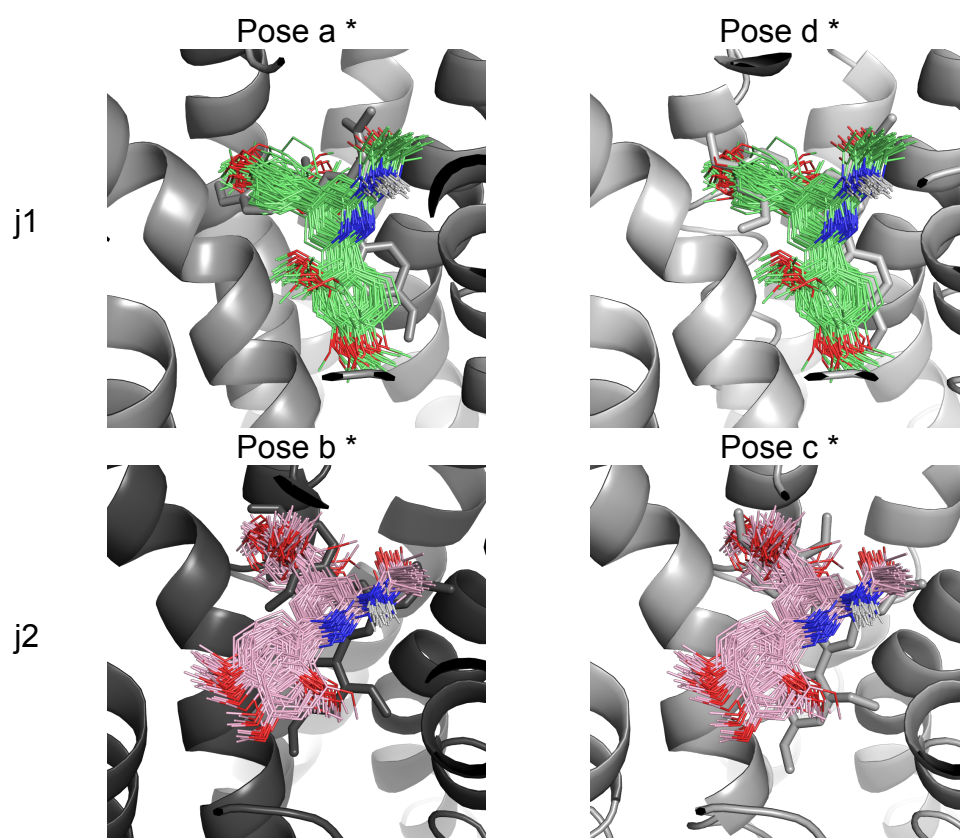


Figure S1: Overlay of trajectory snapshots of 2j1>3j1 and 2j2>3j2 with ligand in lines with poses from Azuaje et al. in grey sticks.

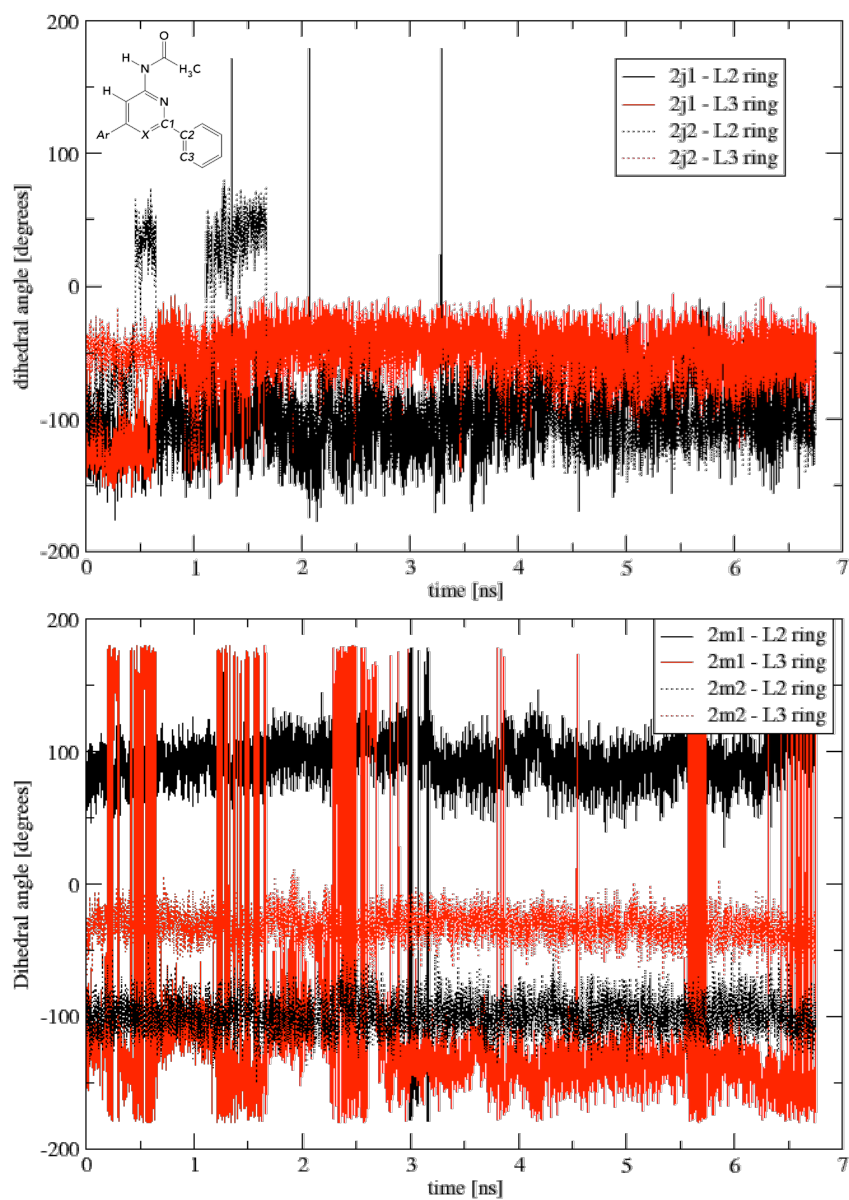


Figure S2: Time series of dihedral angles between X-C1-C2-C3 on both L2 and L3 rings for **m** and **j** compounds in starting poses 1 and 2.