

Supporting Information

Multi-Target Chemometric Modelling, Fragment Analysis and Virtual Screening with ERK Inhibitors as Potential Anticancer Agents

Amit Kumar Halder *, Amal Kanta Giri and Maria Natália Dias Soeiro Cordeiro *

Department of Chemistry and Biochemistry, University of Porto, 4169-007 Porto, Portugal

* Correspondence: amit.halder@fc.up.pt (A.K.H.); ncordeir@fc.up.pt (M.N.D.S.C.);

Tel.: +351-22-040-2502 (M.N.D.S.C)

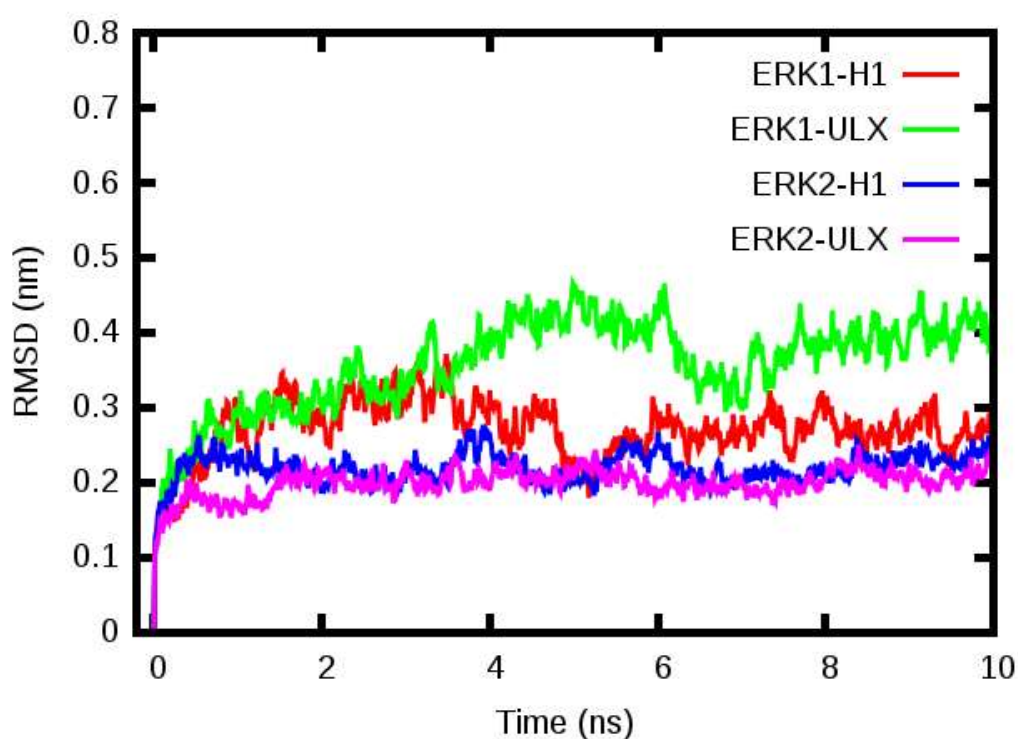


Figure S1. The root mean square deviation (RMSD) plots of backbone atoms of the receptor-ligand complexes.

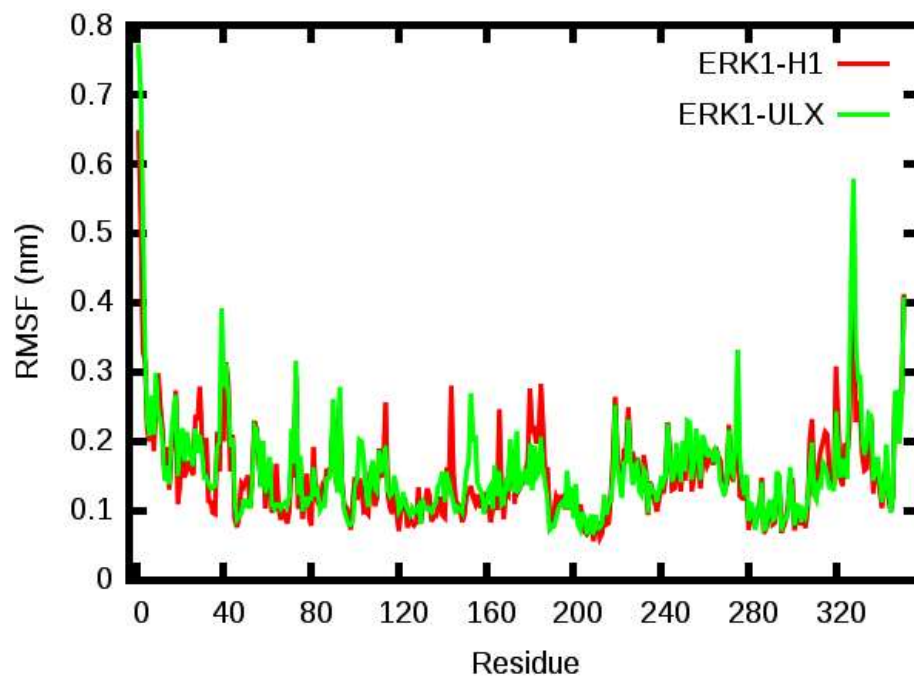


Figure S2. The root mean square fluctuation (RMSF) plots of ERK-1 complexes.

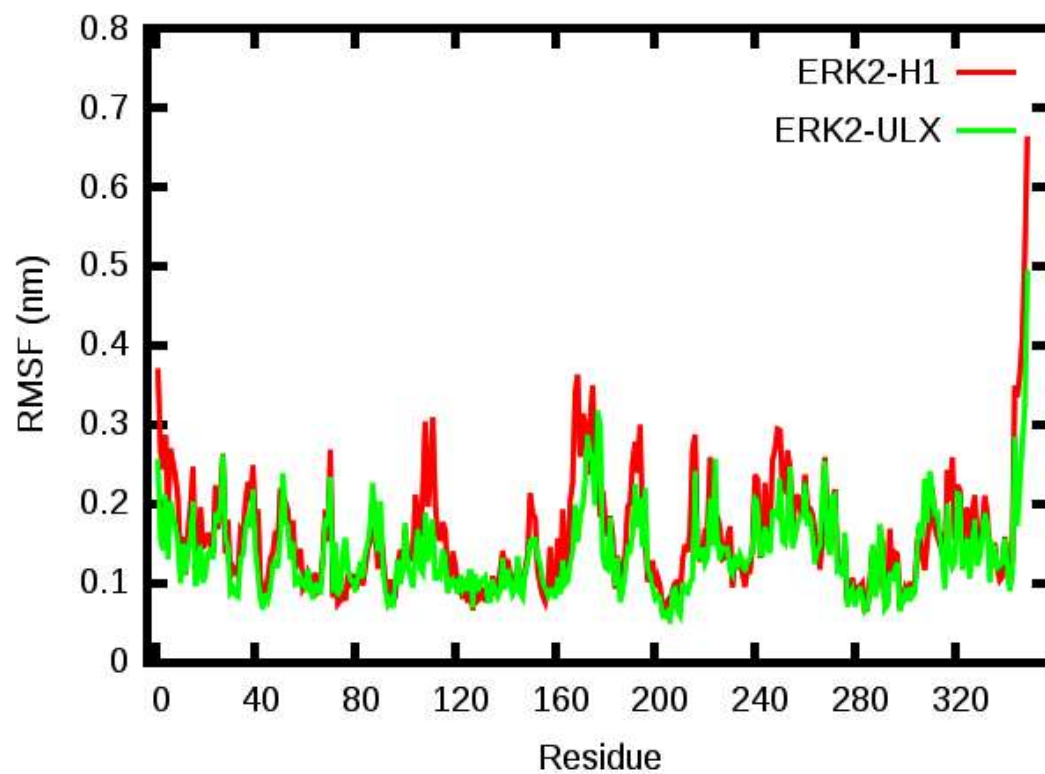


Figure S3. The root mean square fluctuation (RMSF) plots of ERK-2 complexes.

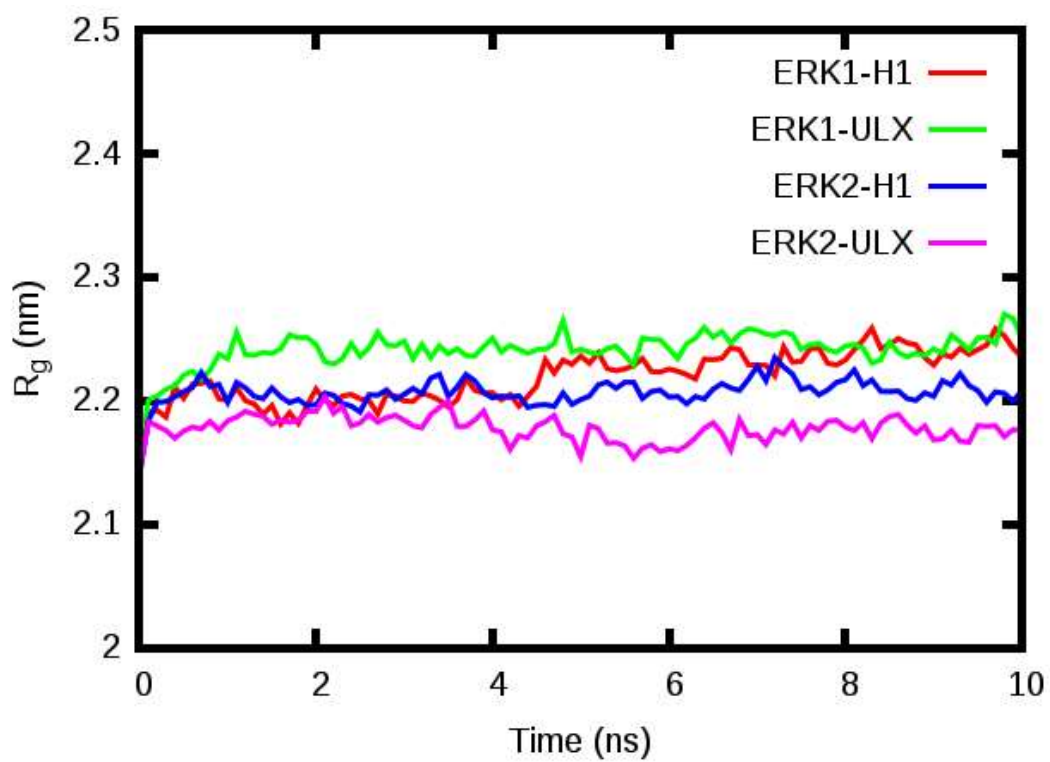


Figure S4. The radius of gyration (R_g) plots of ERK-1 and ERK-2 complexes

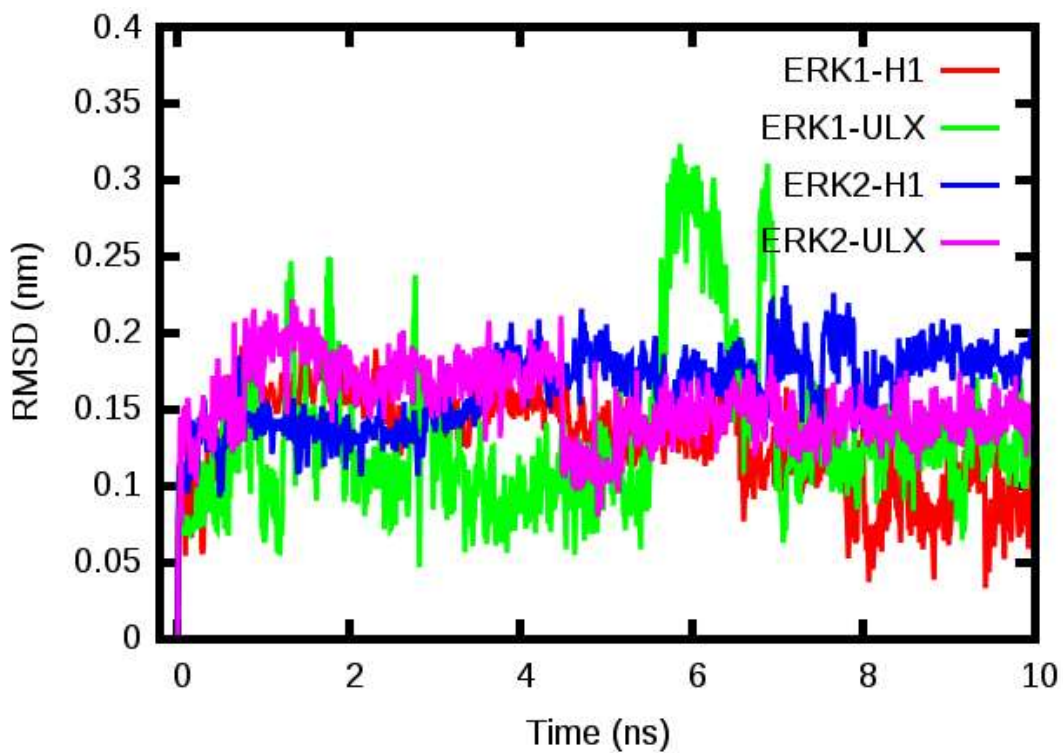


Figure S5. The root mean square deviation (RMSD) plots of bound ligands of the receptor-ligand complexes.

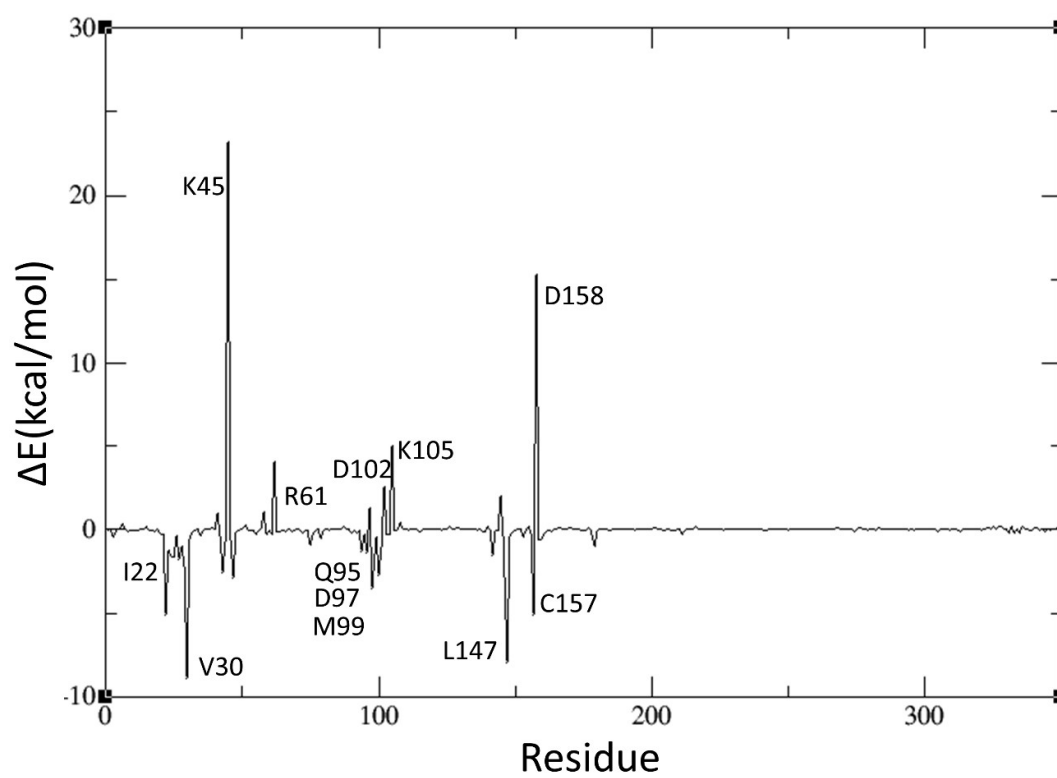


Figure S6. Per-residue decomposition profile of ERK2-ULX complex