Run parameters

Library screened ChEMBL
Screening method Electroshape

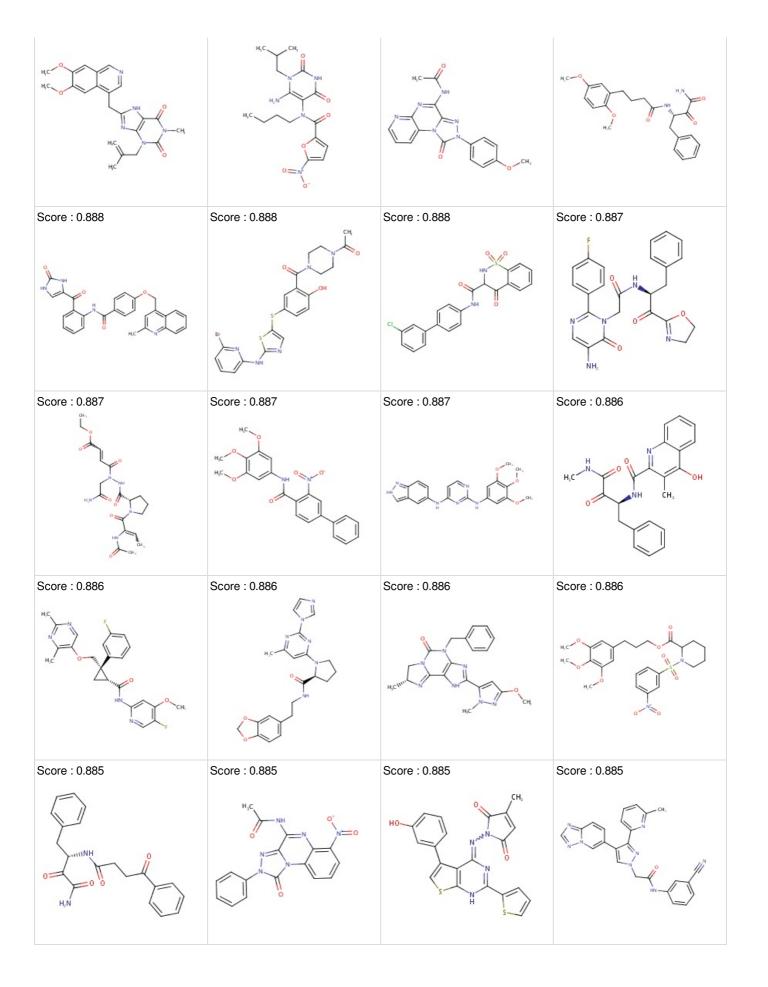
Date Thu Jan 2 11:15:21 2020

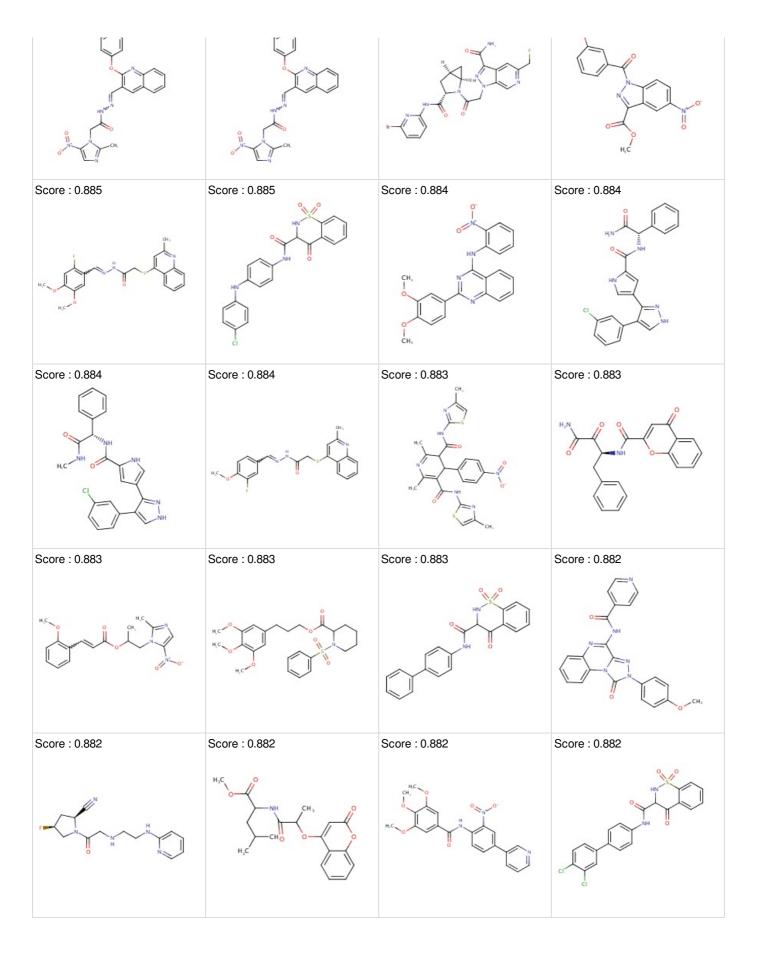
If you publish these results, please cite the SwissSimilarity website.

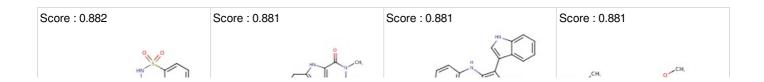
Results

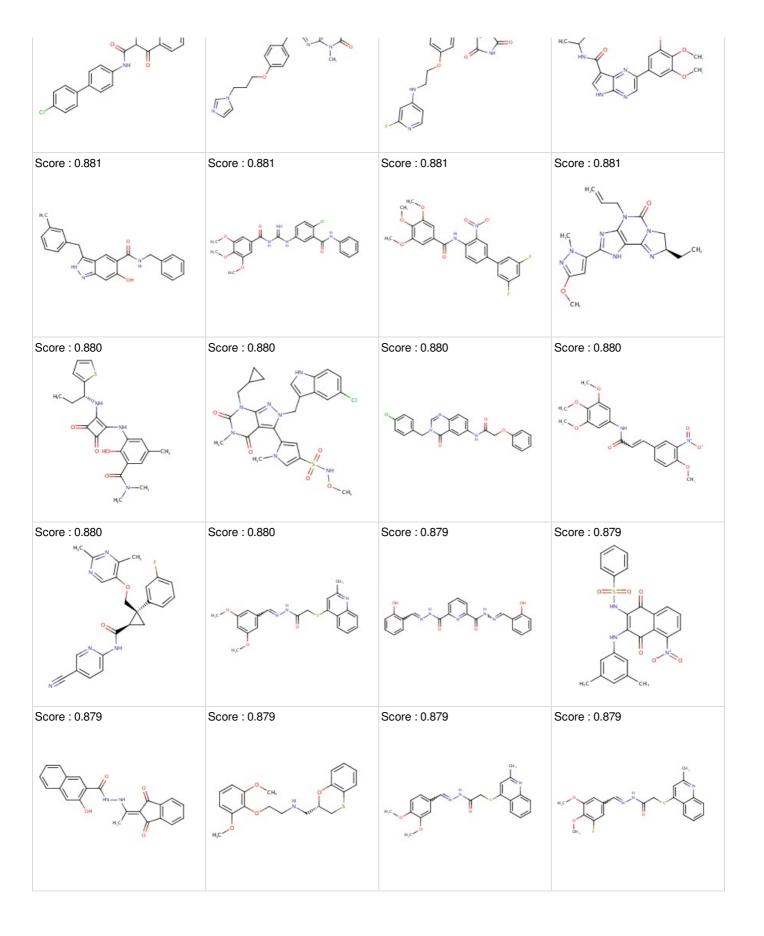
riesuits			
Score : 0.902	Score : 0.900	Score : 0.896	Score : 0.894
H.C. O	NH OCH,	CH, NN	N N N N N N N N N N N N N N N N N N N
Score : 0.894	Score : 0.894	Score : 0.893	Score : 0.893
PRI CH. CH.	H.C. O. S. O	Not the second s	HC CH CH
Score : 0.892	Score : 0.892	Score : 0.891	Score : 0.891
H.C.	H,C H	HI CH.	HC H O CH
Score : 0.891	Score : 0.890	Score : 0.890	Score : 0.889
H ₂ C NH ₂	N O CH ₁	CH ₃ HN NH ₂	MC O CO

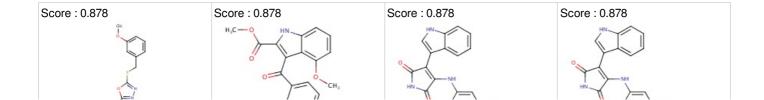
Score : 0.889 Score : 0.888 Score : 0.888

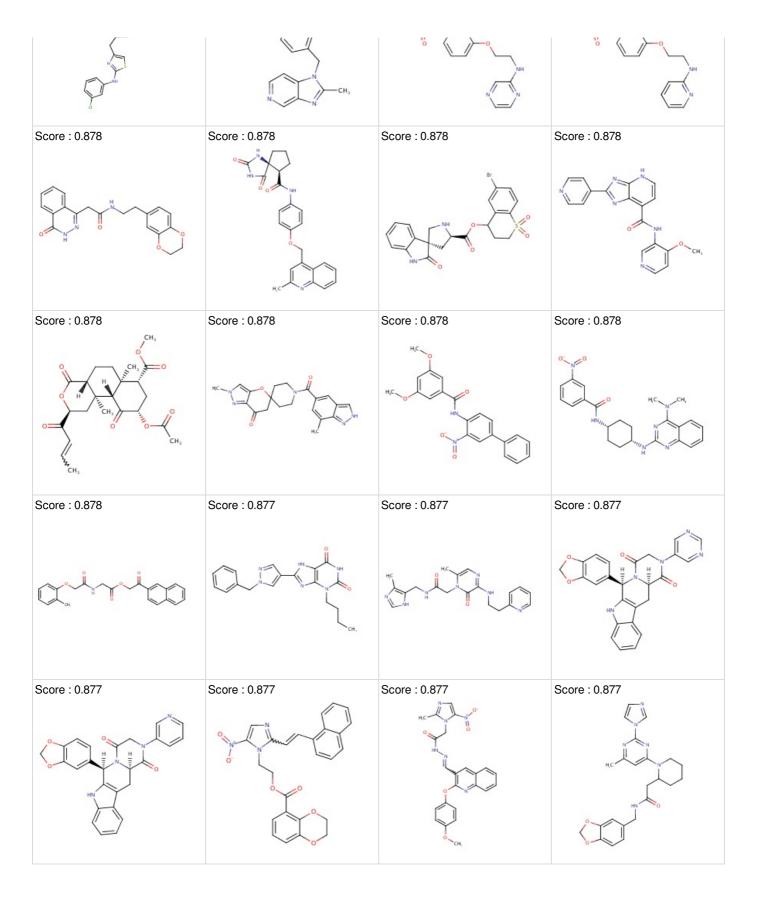


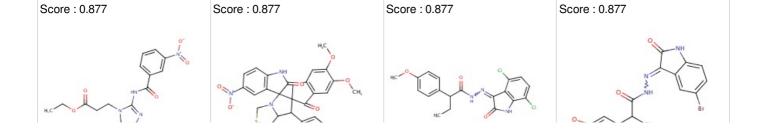


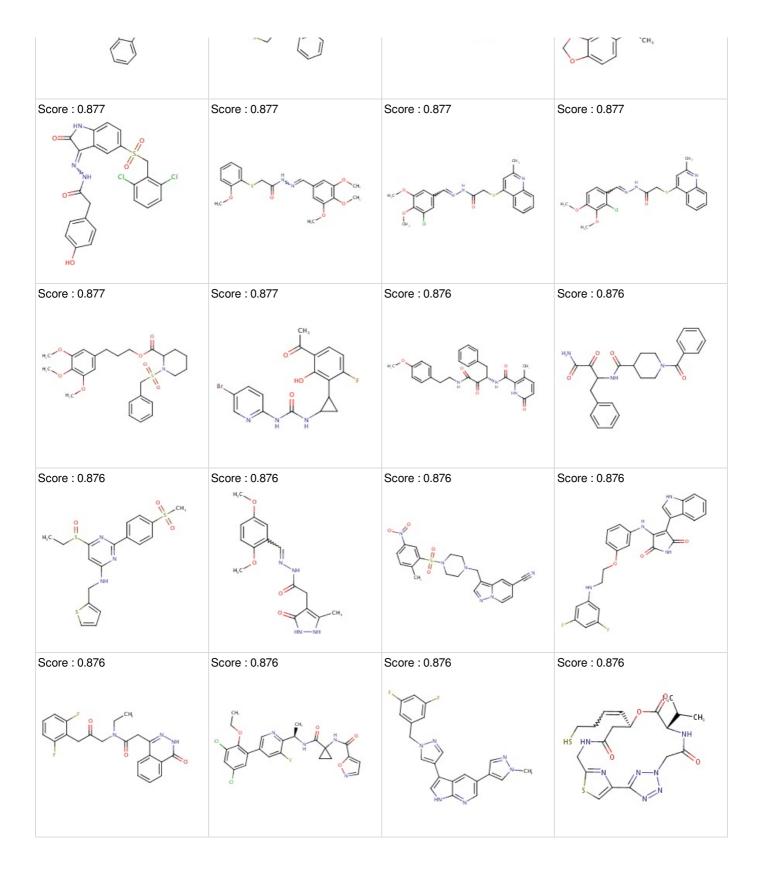


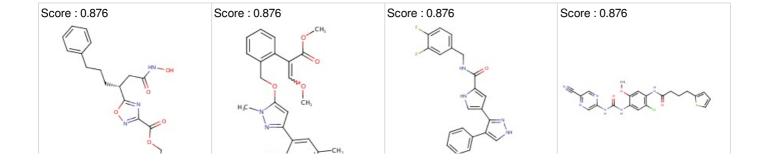


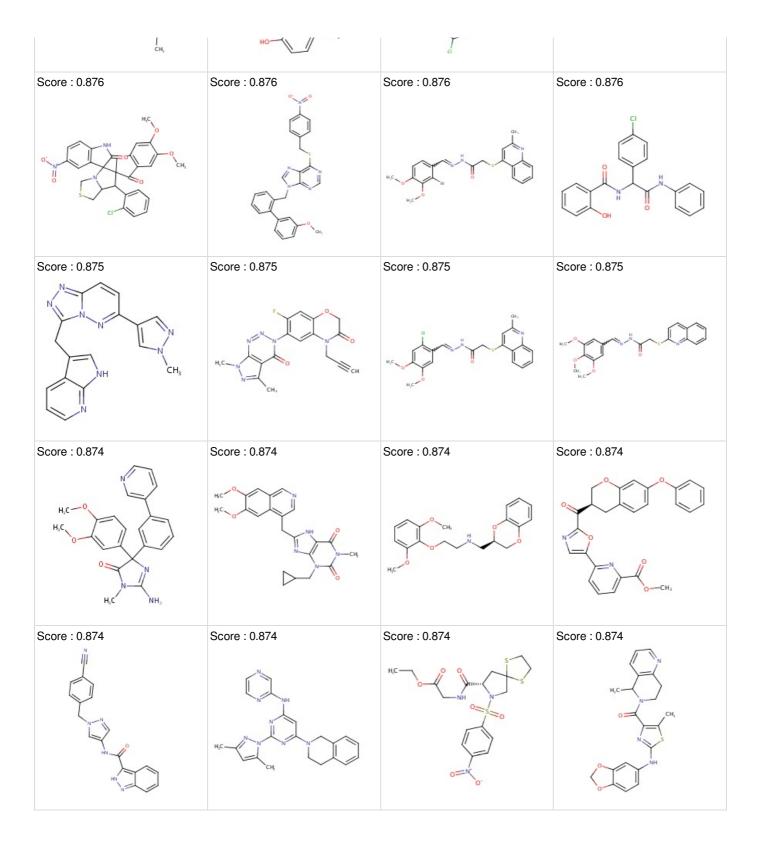


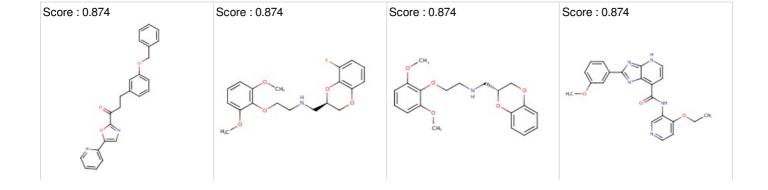


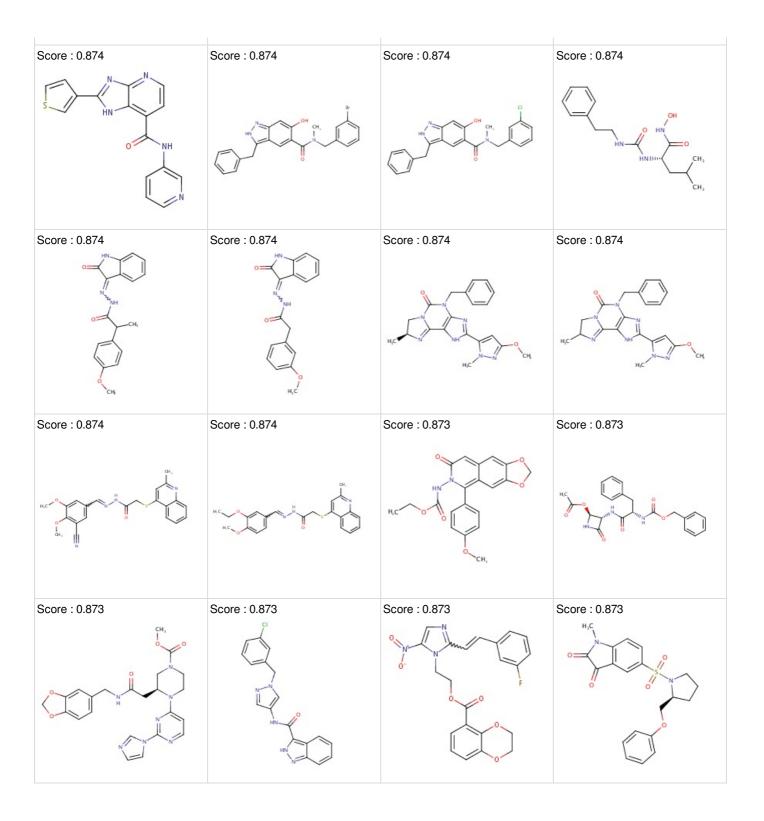


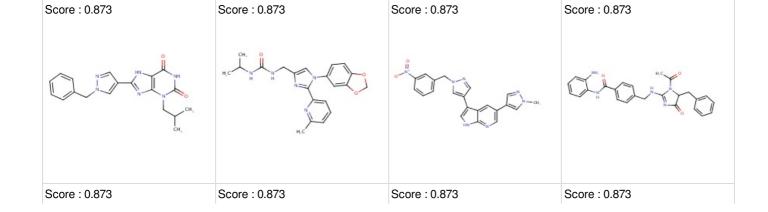


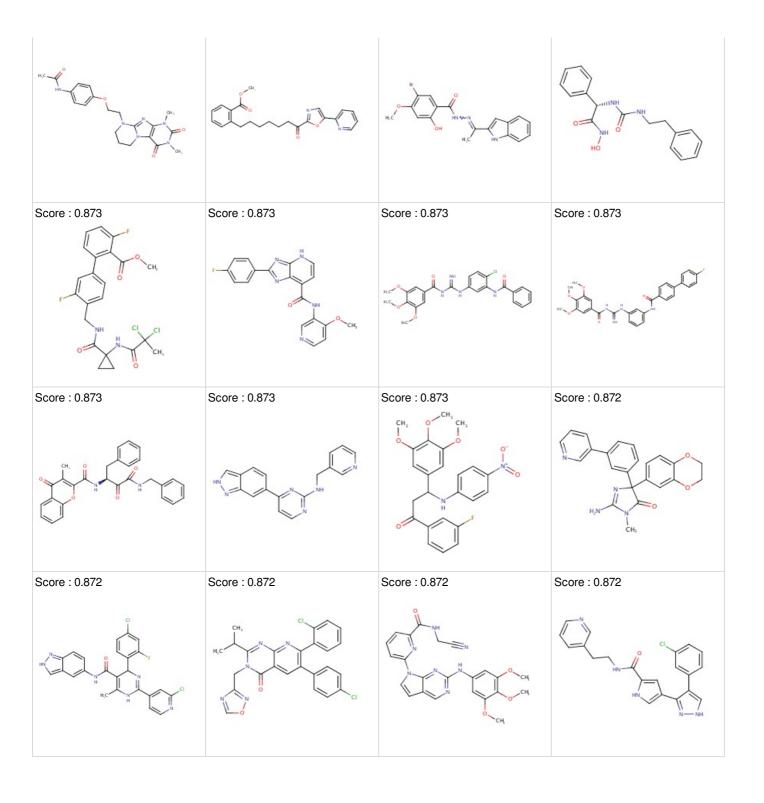


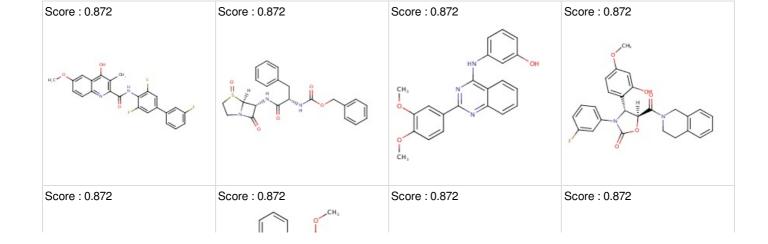


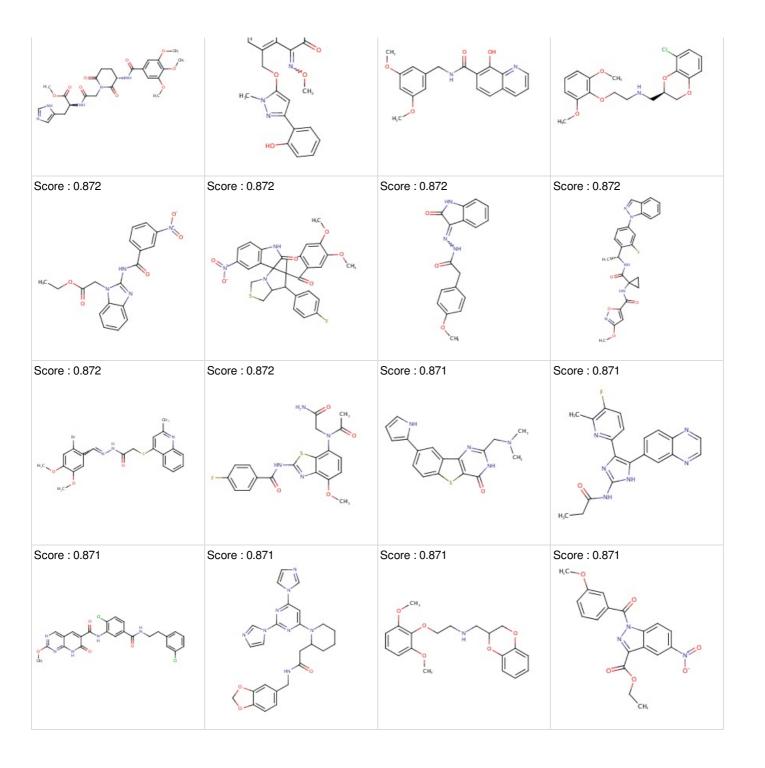


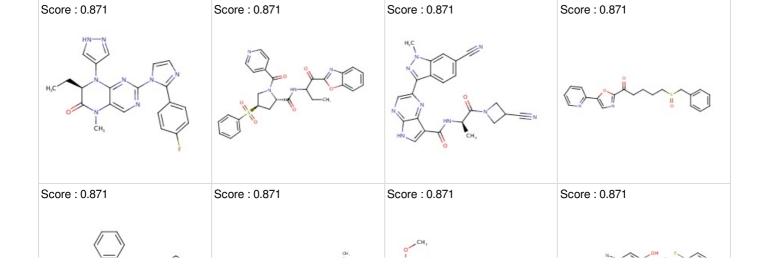


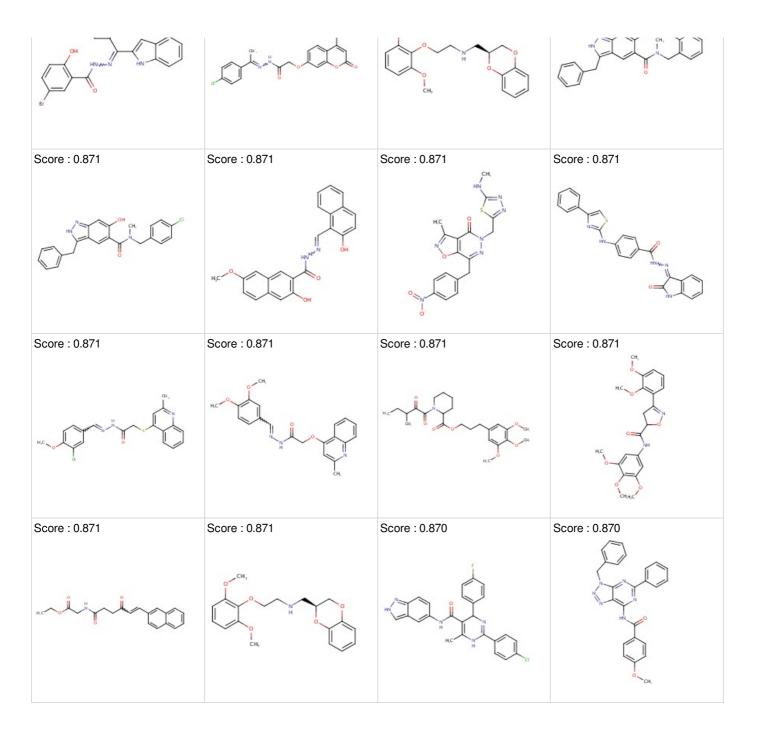


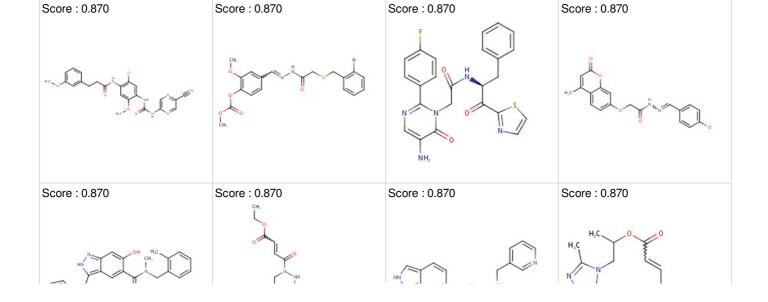


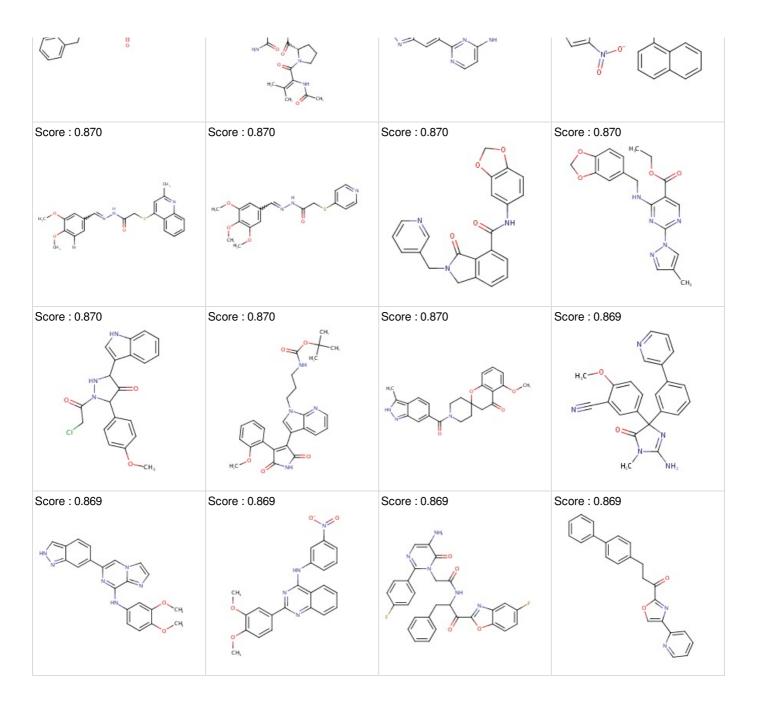


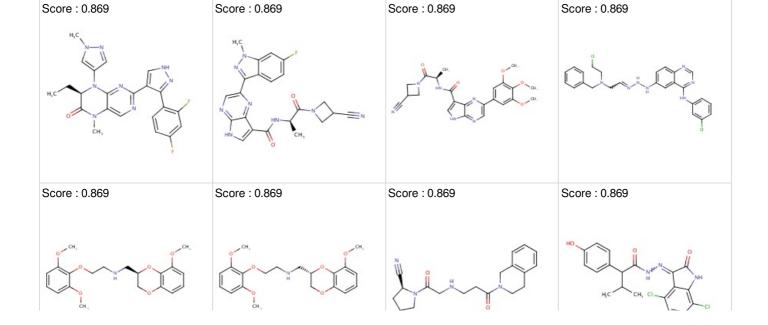


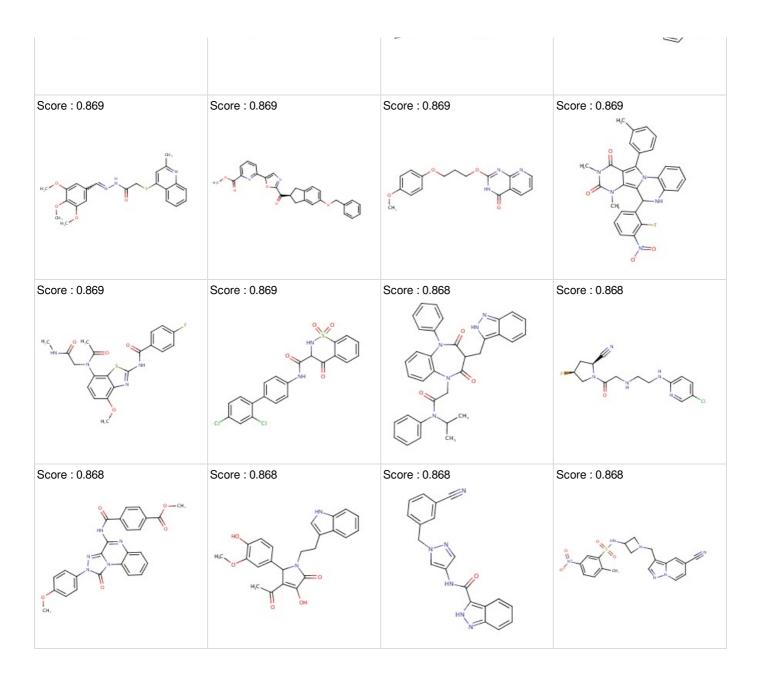


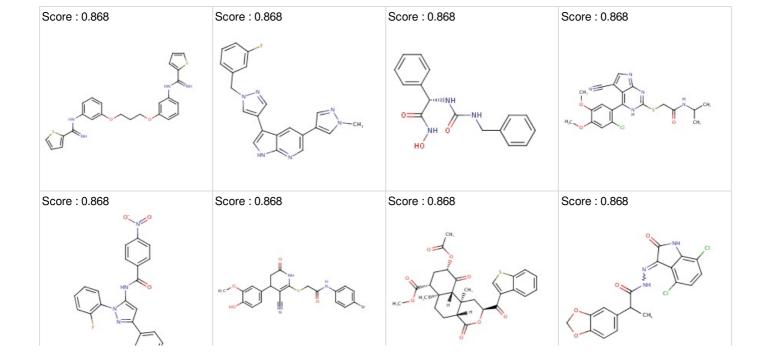


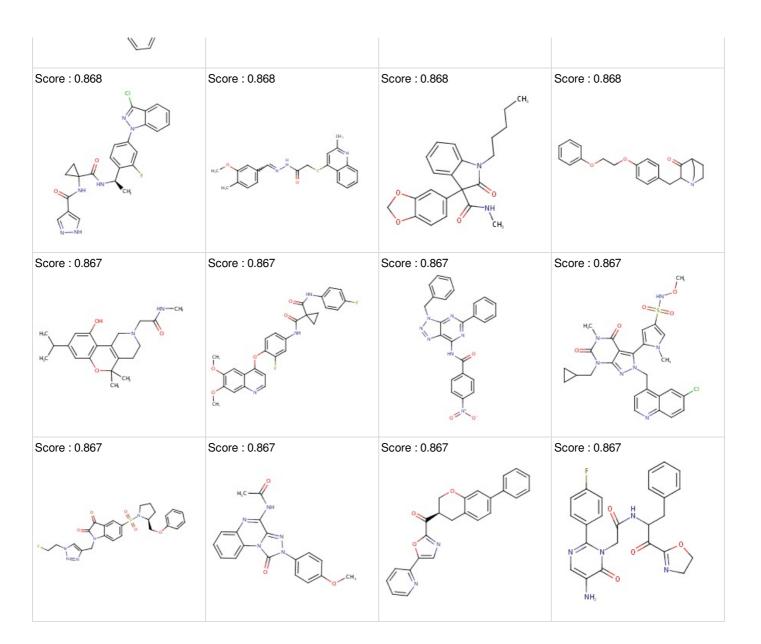


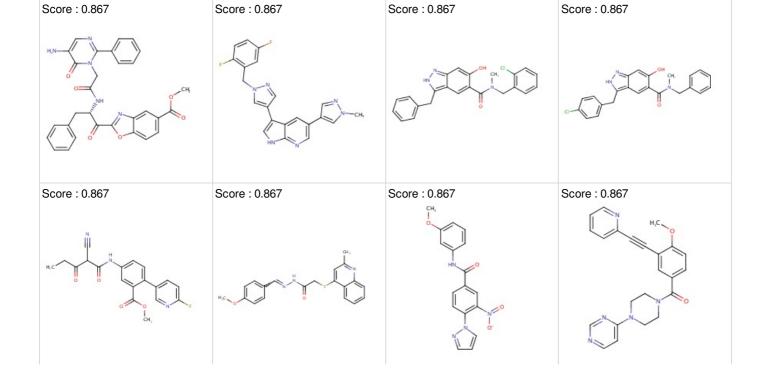


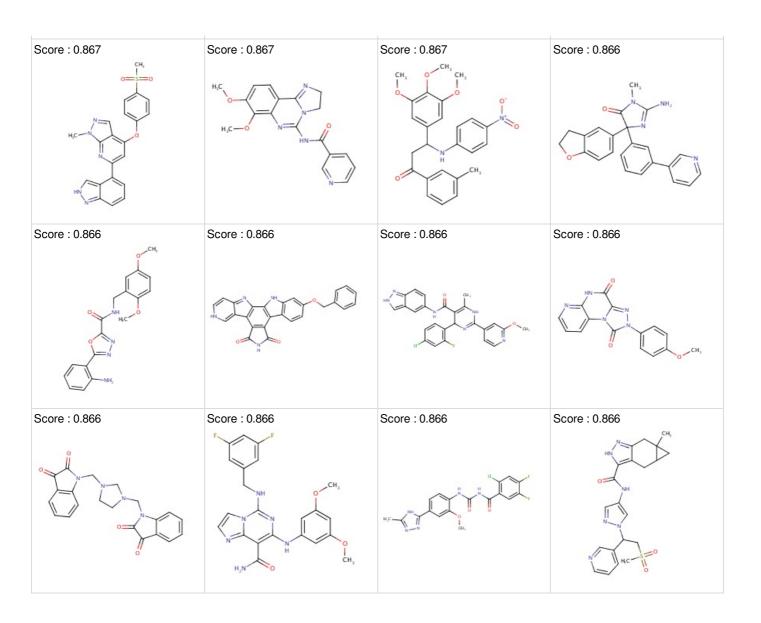


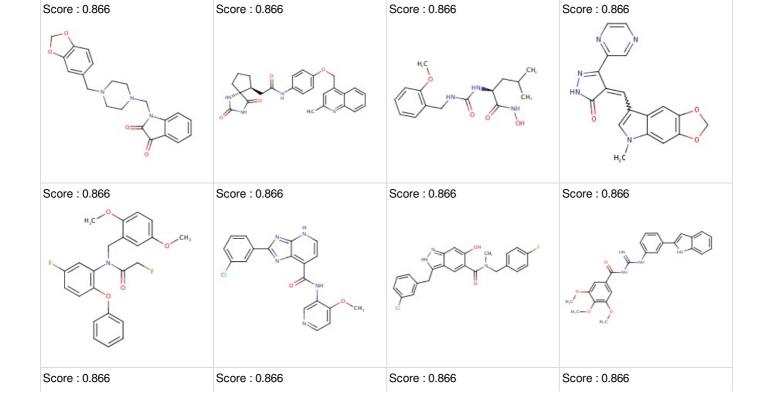


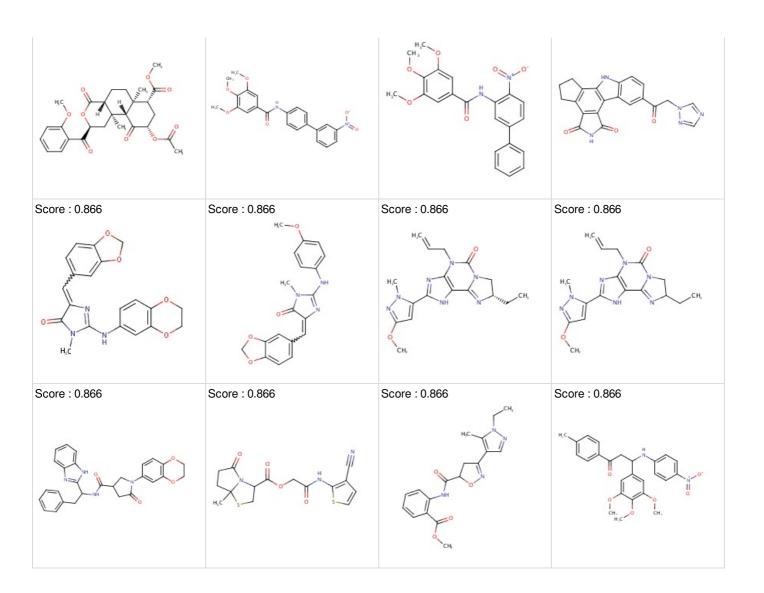


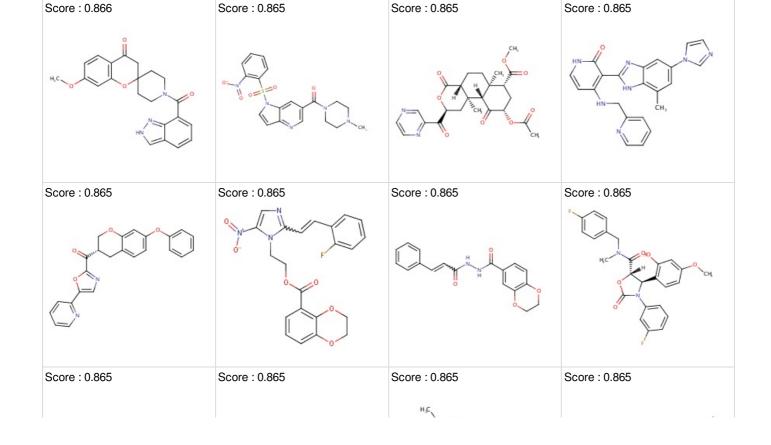


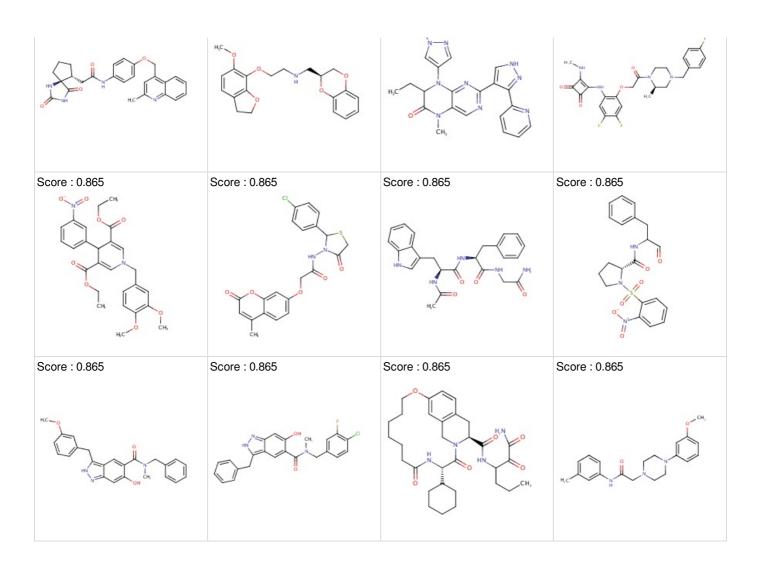


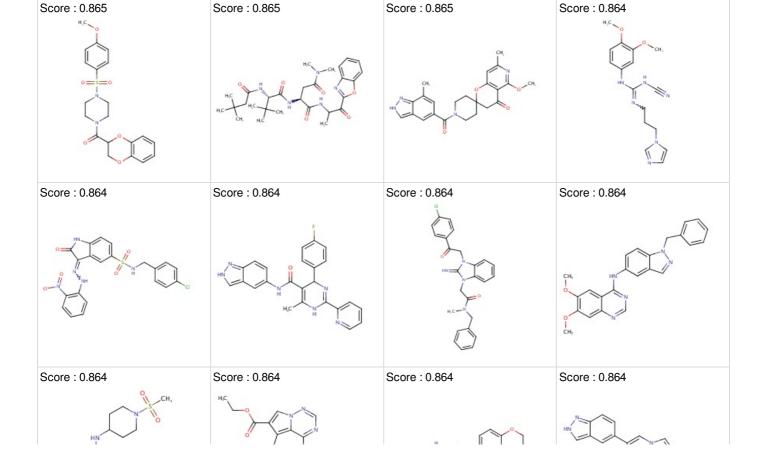


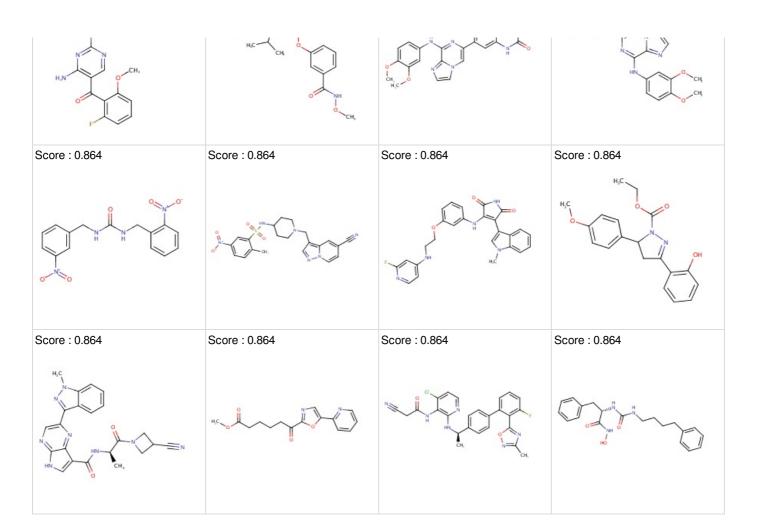


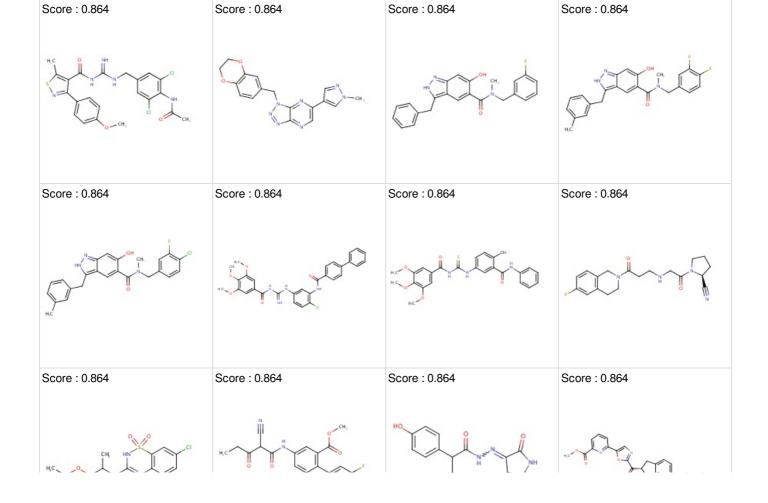


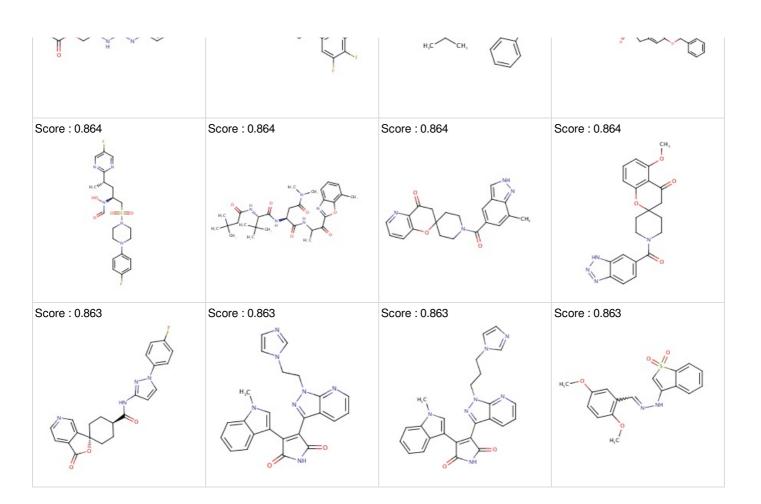


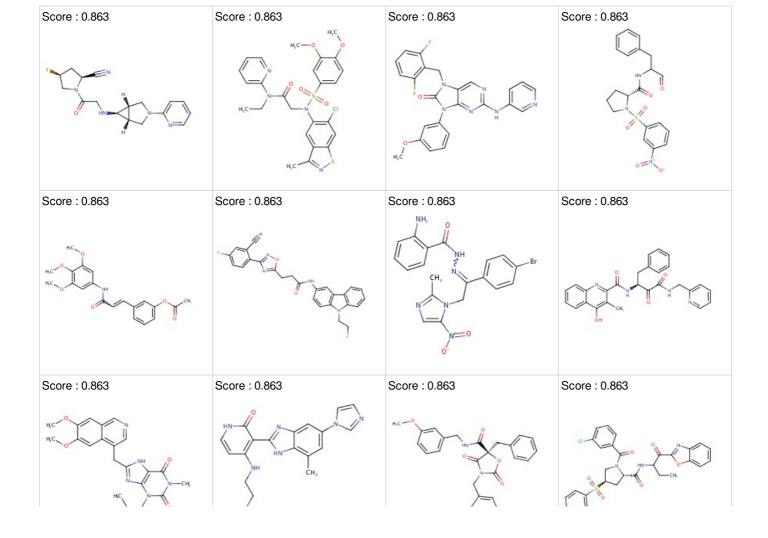


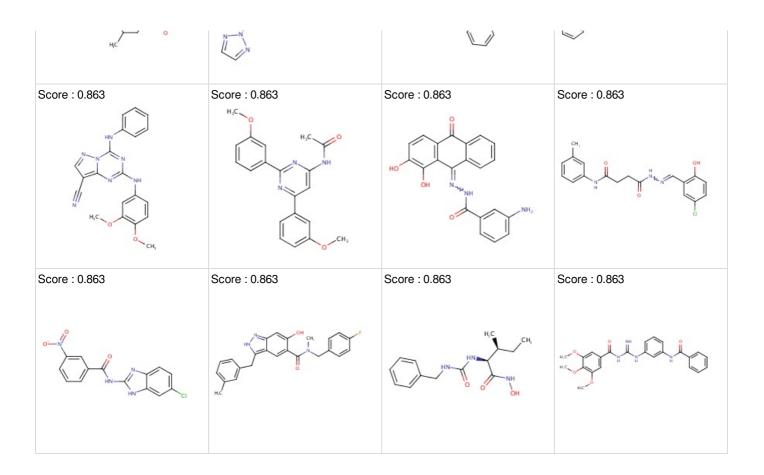


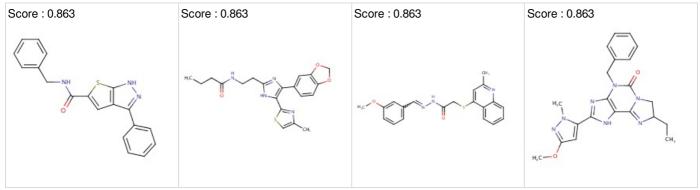












Swiss Institute of Bioinformatics - @2015