## Run parameters

Library screened ChEMBL
Screening method Electroshape

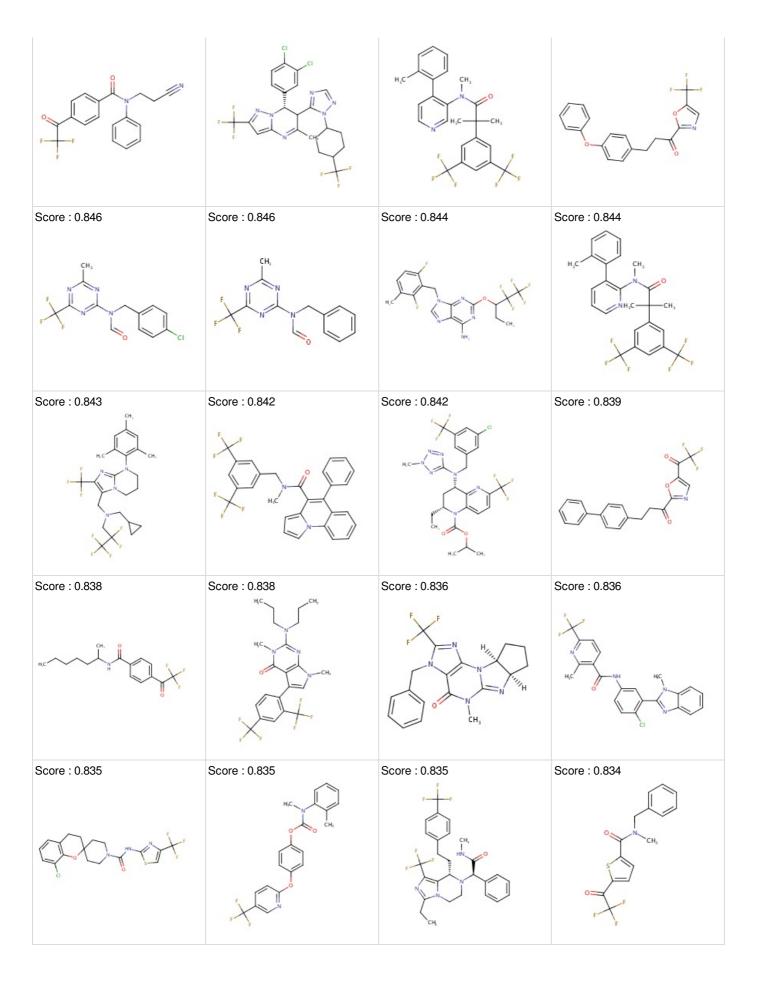
Date Thu Jan 2 11:16:21 2020

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

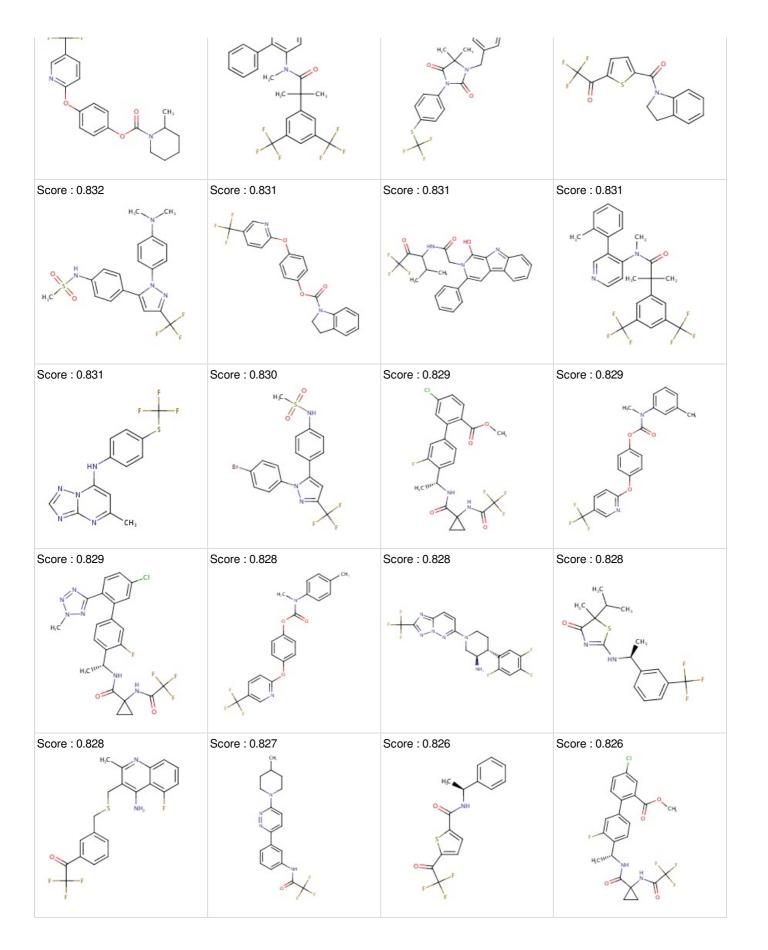
## Results

riesurs			
Score: 0.893	Score : 0.884	Score: 0.874	Score : 0.873
	, X		H,C————————————————————————————————————
Score : 0.871	Score : 0.870	Score : 0.864	Score : 0.862
H,C CH,	F F CH <sub>3</sub>	H <sub>1</sub> C N N CH <sub>3</sub>	N CH,
Score: 0.860	Score : 0.860	Score: 0.860	Score : 0.855
CH, N CH,	F HAC CH CH	CH	H,C N N NH
Score : 0.854	Score : 0.854	Score : 0.852	Score : 0.851
H,C N		H.C. N. H.C. N.	H,C N F F

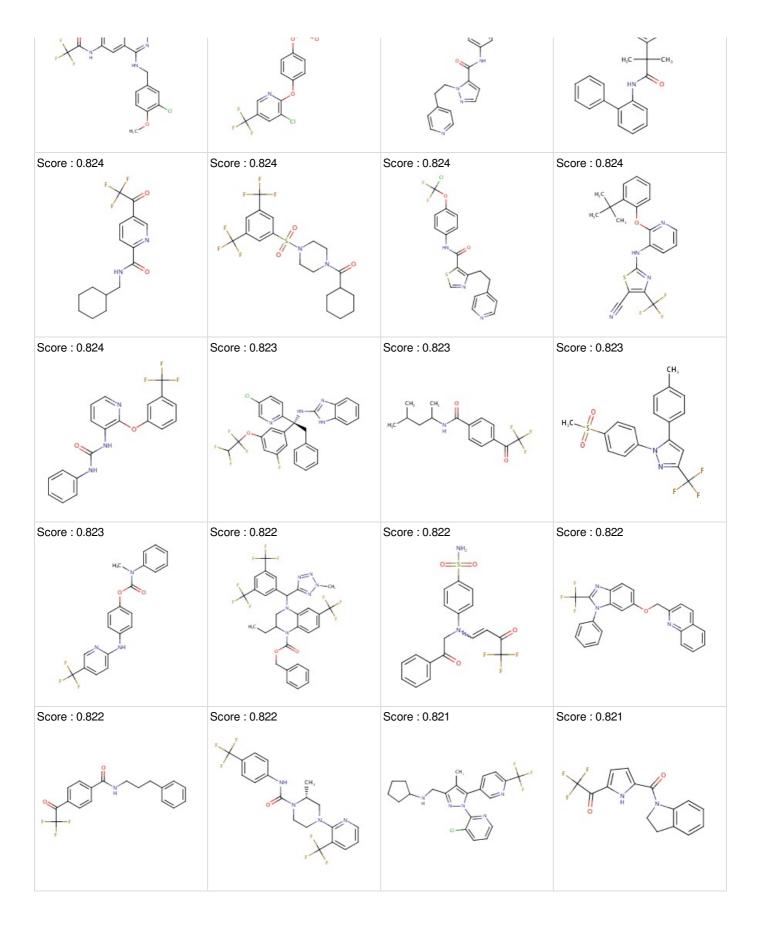
Score : 0.850 Score : 0.848 Score : 0.847 Score : 0.847

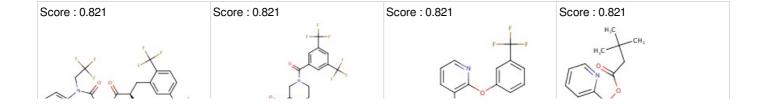


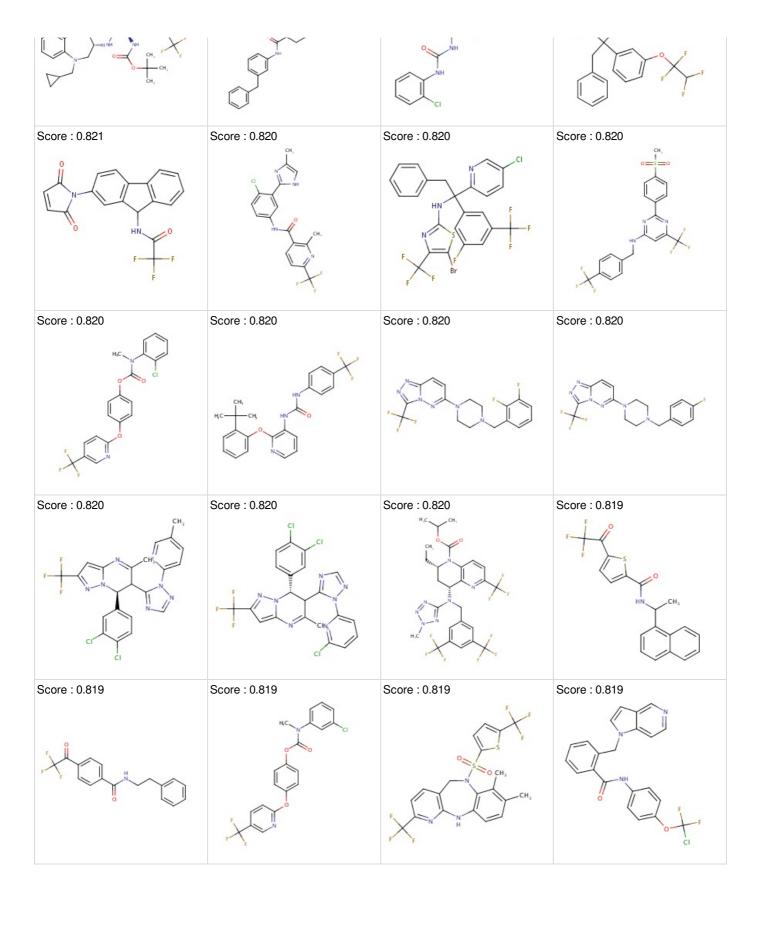


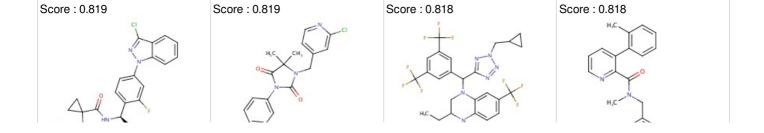


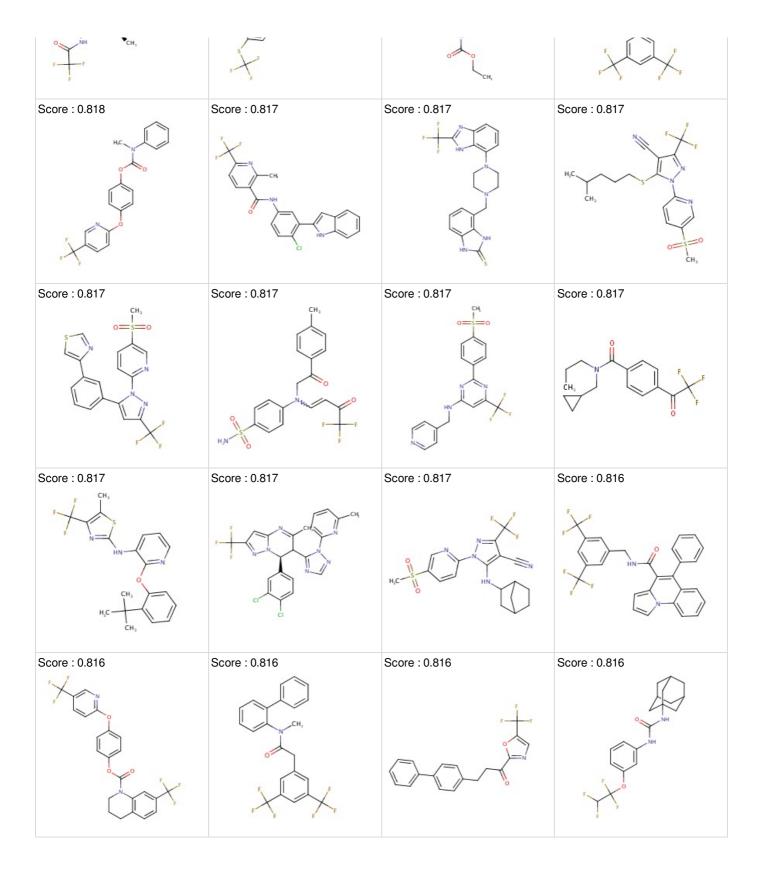


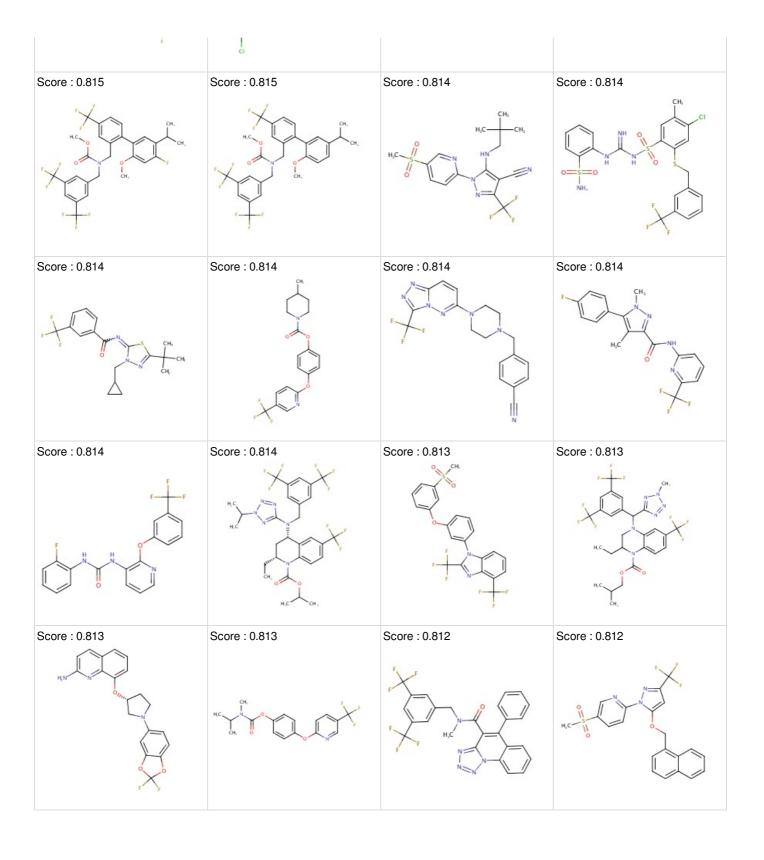


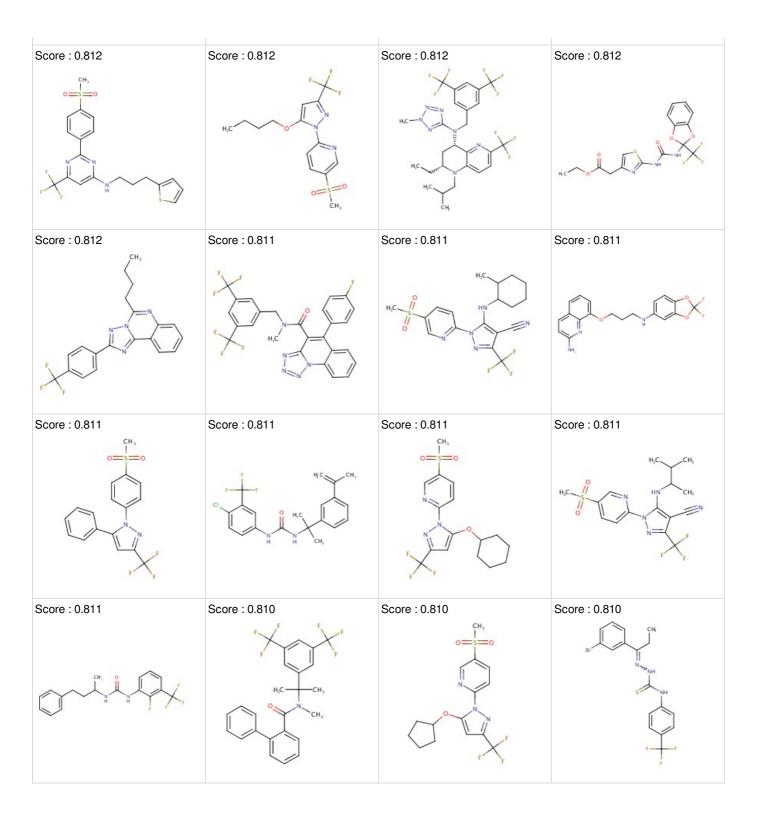


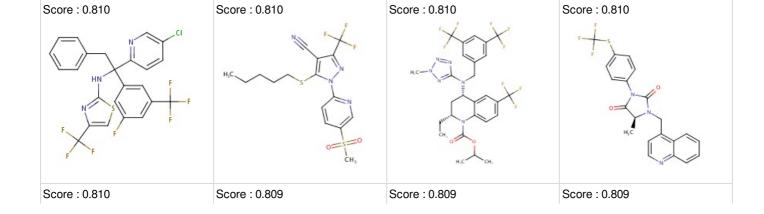


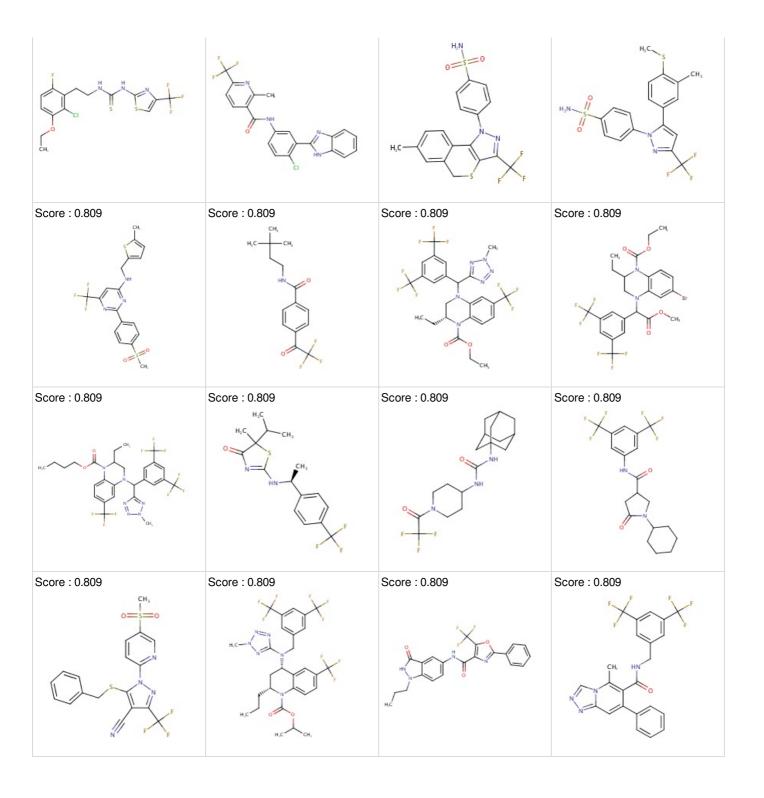


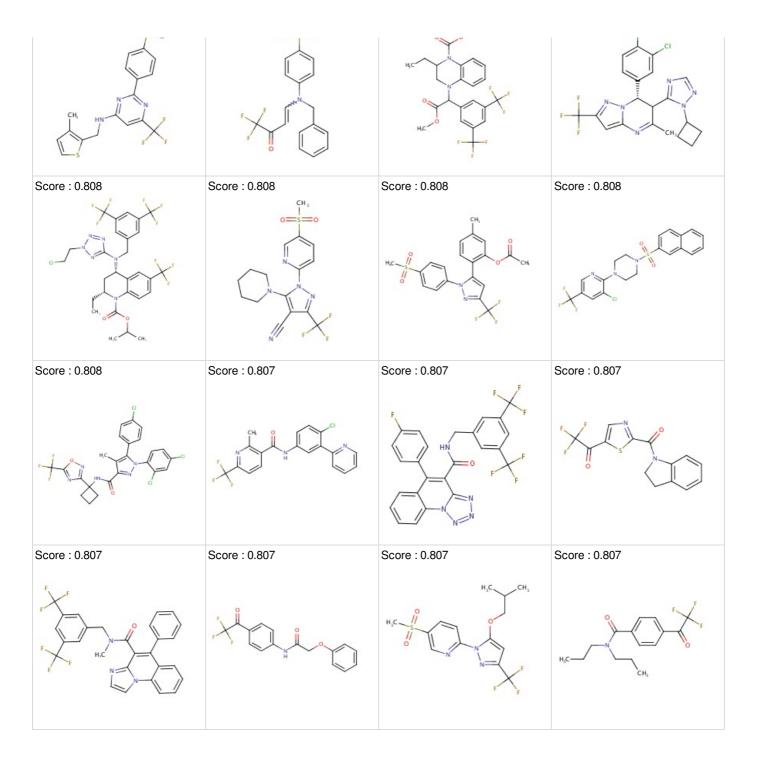


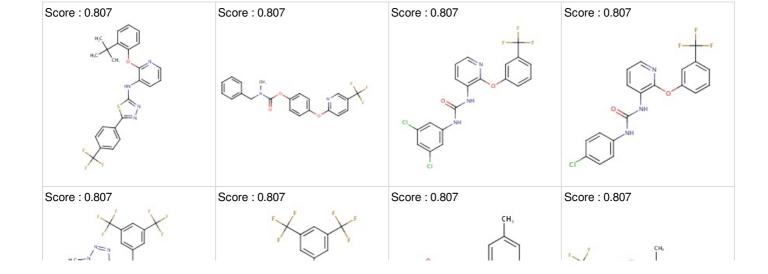


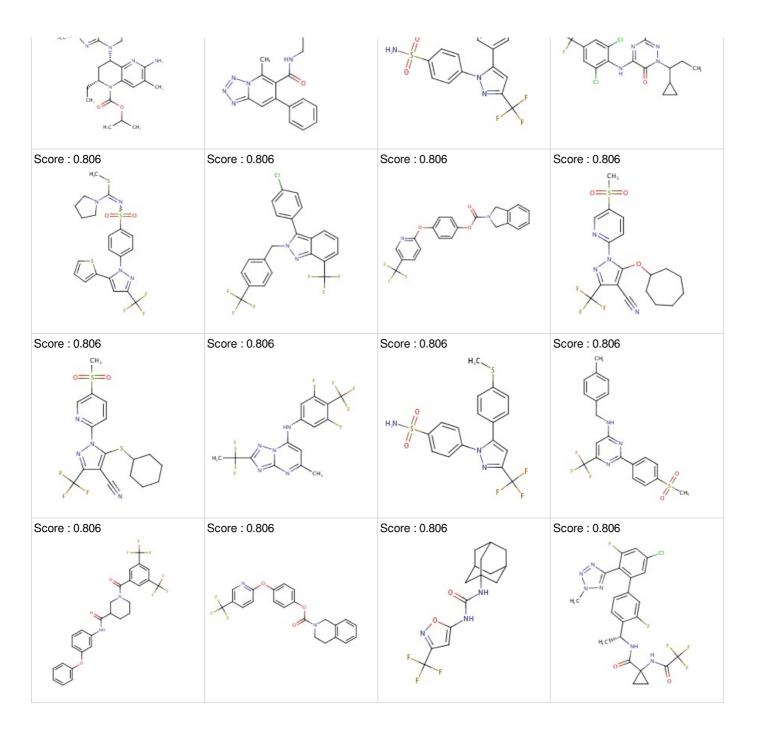


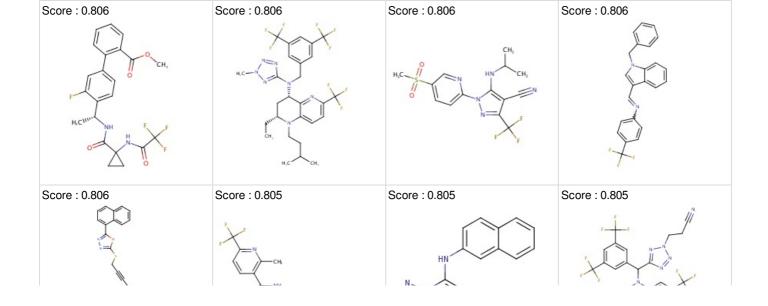


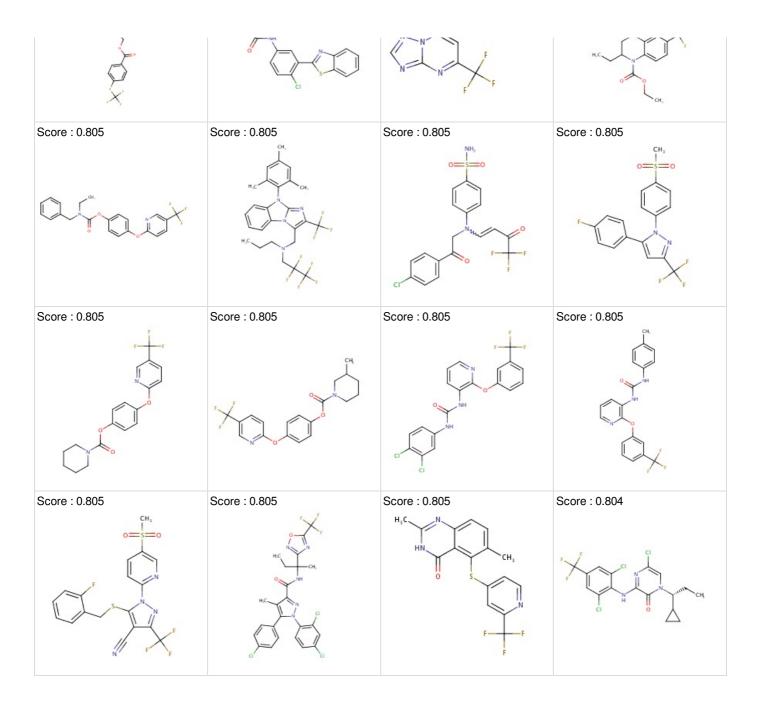


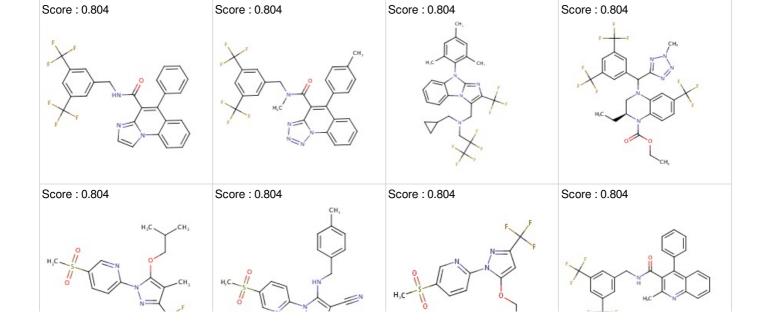


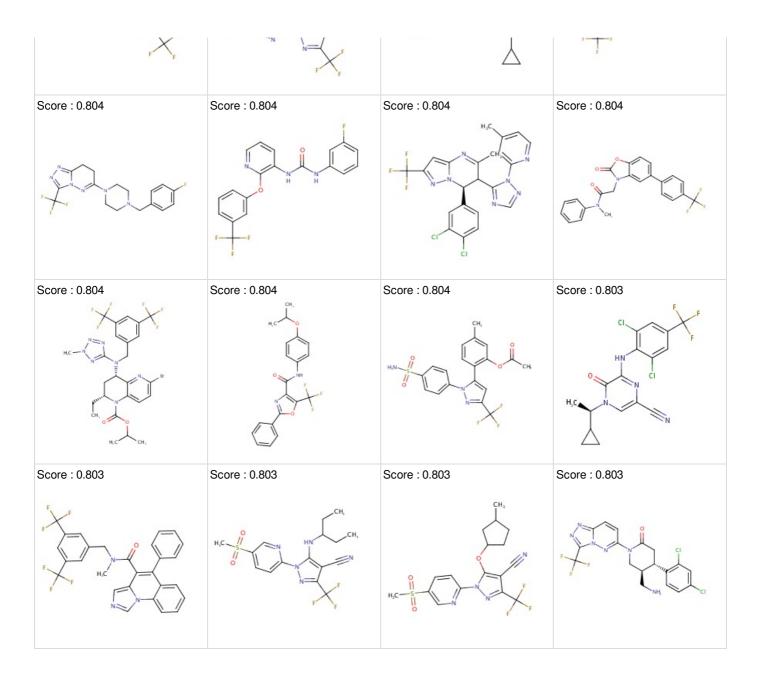


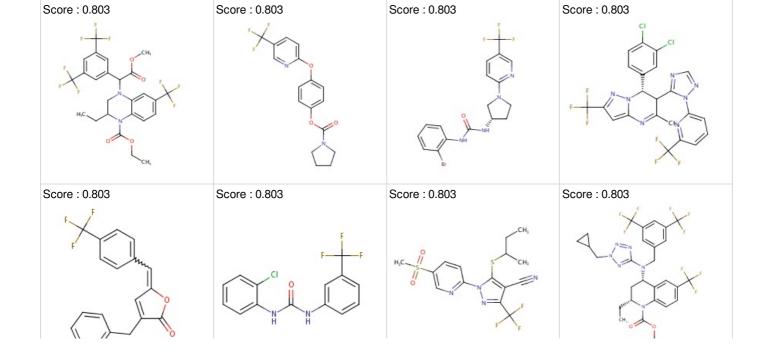


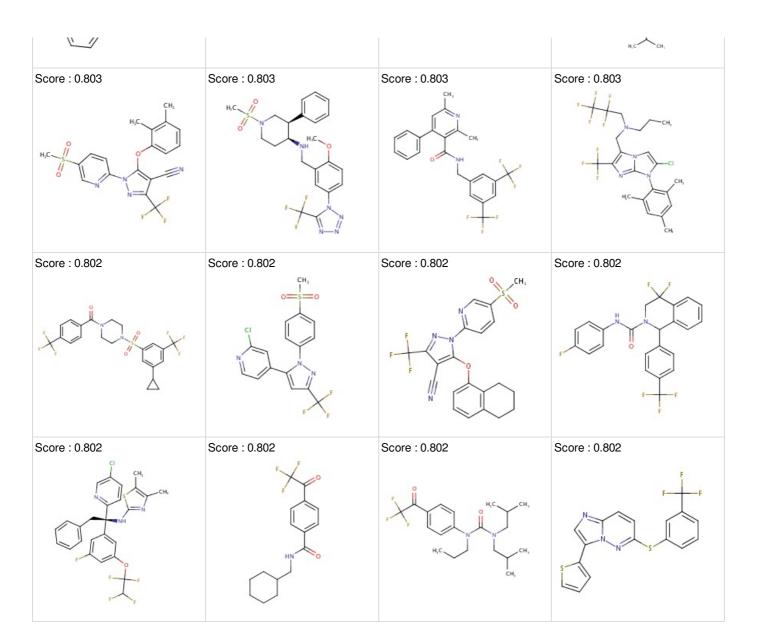


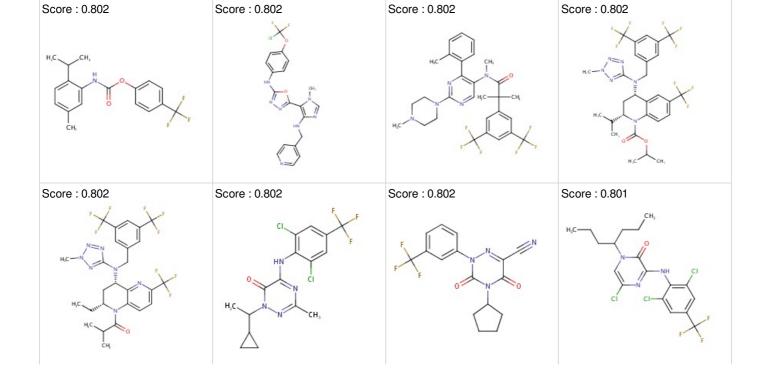


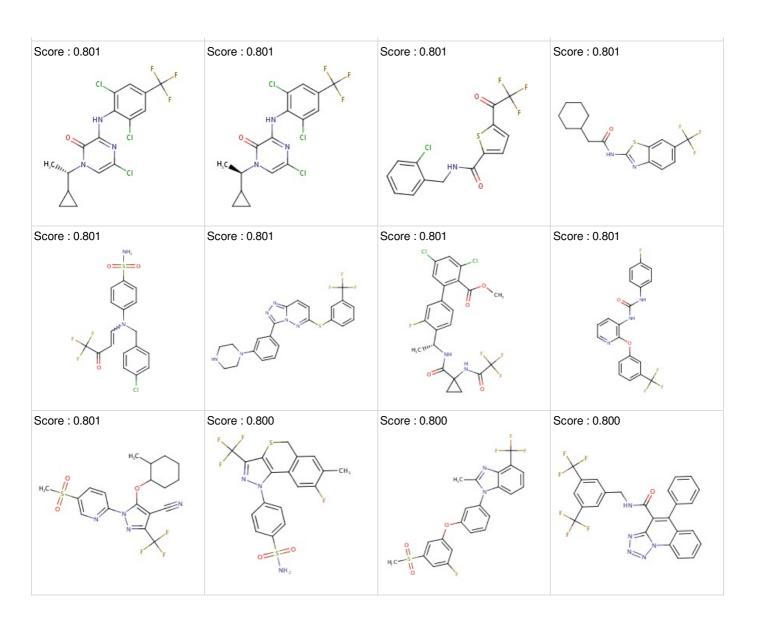


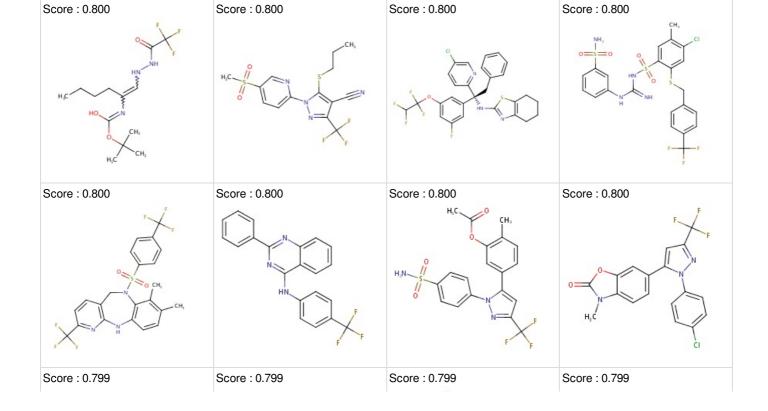


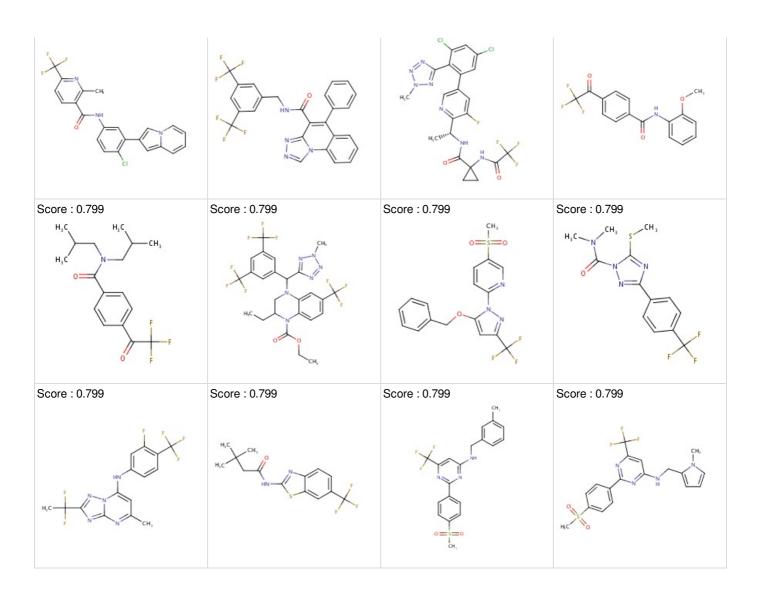


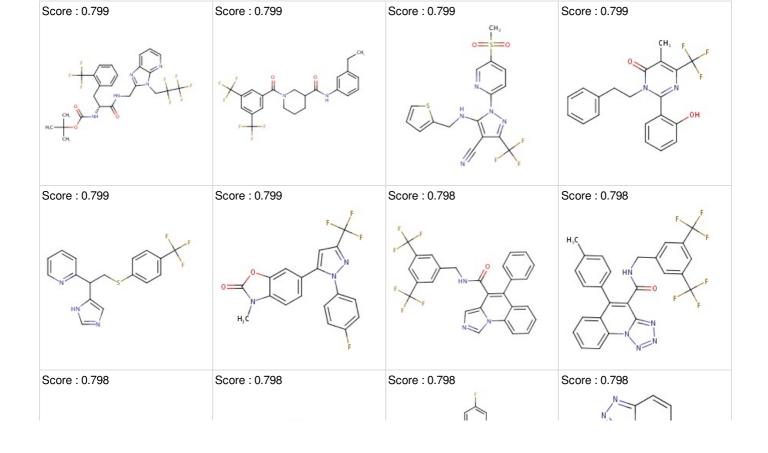


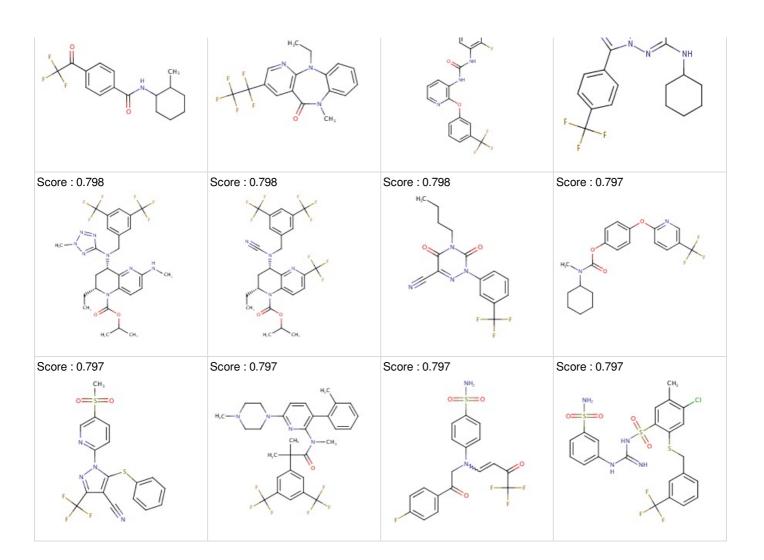


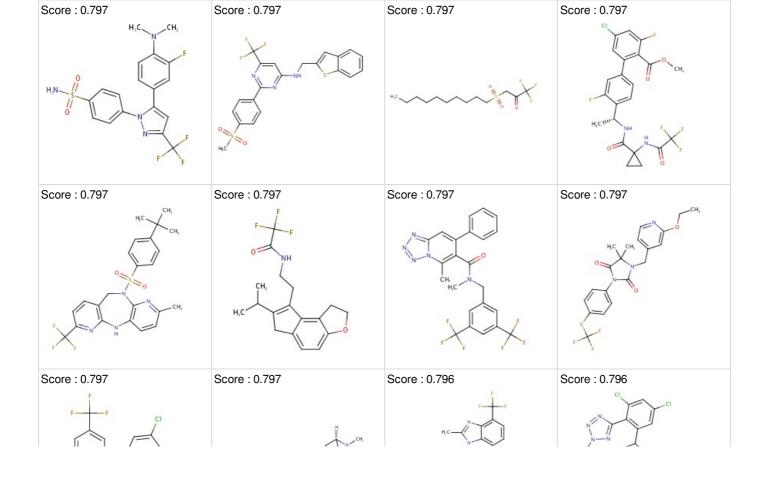


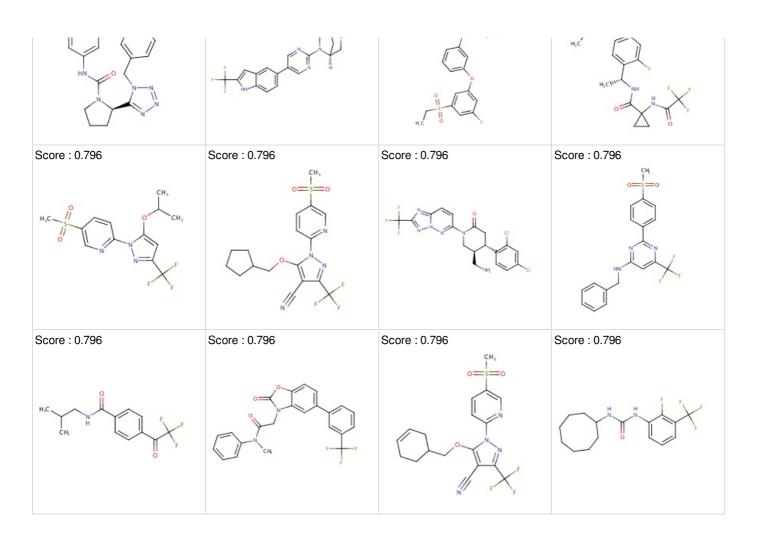


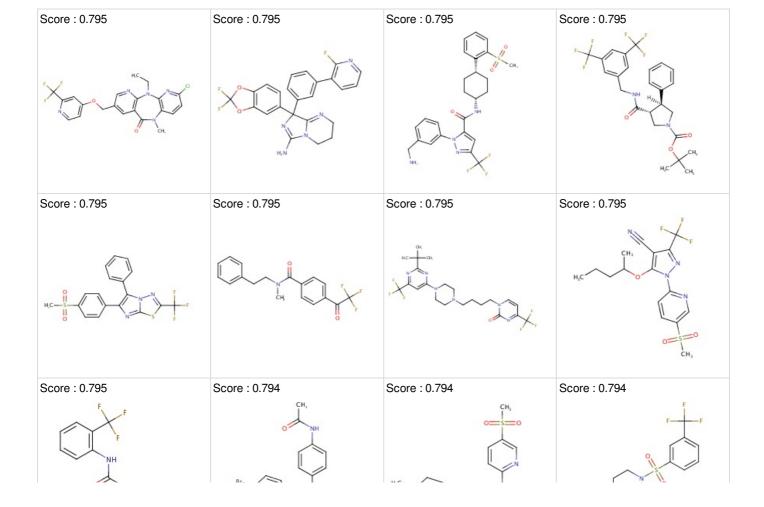


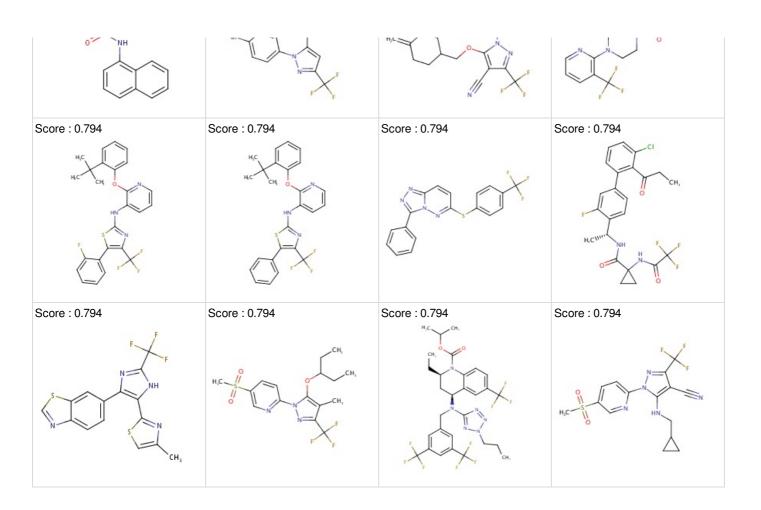


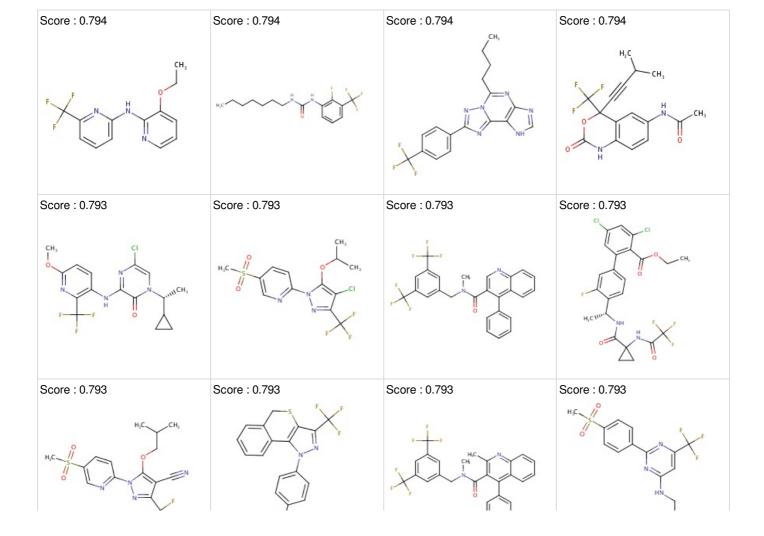


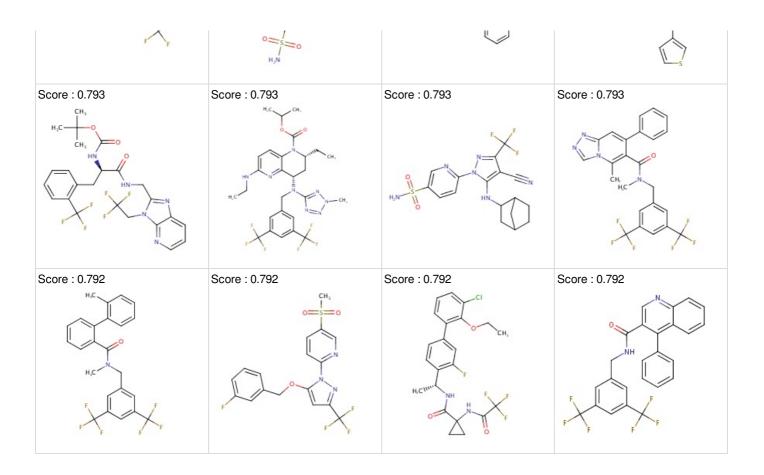


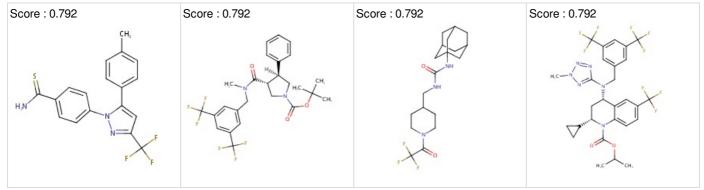












Swiss Institute of Bioinformatics -  $\odot$  2015