

Computational Epidemiology – PS3

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Summary

- The descriptors for the compounds provided was obtained using the ChemDes web server.
- 1875 descriptors (PaDEL) were obtained after giving the SMILES data as input.
- A regression QSAR model was built using Python based on the descriptors obtained and the pIC₅₀ data.
- After training and validation of the model, the pIC₅₀ for the 'BLINDED' compounds was found and analysed.

BLINDED pIC₅₀ Scores

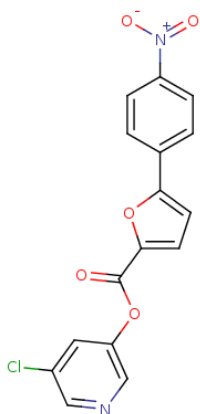
Compound No.	SMILES	Predicted pIC ₅₀
6	<chem>ClC1=CC(OC(=O)C2=C3C=CC=CC3=CC=C2)=CN=C1</chem>	-0.60950964
12	<chem>CC1=CC(O)=NC(SCC(=O)NC2=CC=C(OC3=CC=C(Cl)C=C3)C=C2)=N1</chem>	-2.183443378
24	<chem>CC1=C(C=C(O1)C(C)(C)C1=NNC(NS(=O)(=O)C2=CC=CS2)=C1</chem>	-1.568560844
25	<chem>ClC(Cl)=C(Cl)C(=O)OC1=CC=C(C=C1)S(=O)(=O)C1=CC=C(OC(=O)C(Cl)=C(Cl)Cl)C=C1</chem>	-1.283476567
31	<chem>O=C1N(CC2=CC=C3C=CC=CC3=C2)C2=CC=C(C=C2C1=O)S(=O)(=O)N1CCCCC1</chem>	-0.839339837
37	<chem>IC1=CC=C2N(C\C=C\C3=CC4=CC=CC=C4S3)C(=O)C(=O)C2=C1</chem>	-0.436155768
43	<chem>CC1CCCCN1S(=O)(=O)C1=CC2=C(NC(=O)C2=O)C=C1</chem>	-0.359989254
59	<chem>[O-][N+](=O)C1=CC=C(C=C1)C1=CC=C(O1)C(=O)OC1=CN=CC(Cl)=C1</chem>	0.478992592
70	<chem>CC(SC1=NC(C2=CC=CC=C2)=C(C#N)C(=O)N1)C(=O)NC1=CC=C(Cl)C=C1</chem>	-2.05000698
80	<chem>CC(C)C1=CC=C(NC(=O)CSC2=NC(=CC=N2)C2=CC=CS2)C=C1</chem>	-1.601485981

IC₅₀ is the half maximal inhibitory concentration. A lower IC₅₀ is desirable for an inhibitor as it achieves half of the maximum possible inhibition at a lower concentration.

pIC₅₀ is the negative logarithm of IC₅₀. Thus, a higher value of pIC₅₀ implies a greater potency of the inhibitor.

From the predicted pIC₅₀ values, **Compound No. 59** seems to be the most potent inhibitor among the 'BLINDED' compounds.

Using the Smi2DDepict web server, the structure of Compound No. 59 is obtained.



Model Validation

After performing a train-test split on the ‘unblinded’ compounds,

- the R^2 score for the model was found to be 0.774
- the LOO- q^2 score for the model was found to be 0.485

According to the Golbraikh–Tropsha model acceptability criteria¹,

- The model should have a high LOO- q^2 score (> 0.5).

Our model has a score of 0.485, which doesn’t meet this criterion, but comes very close to it.

- Correlation coefficient R between the predicted and actual activities for an external test set should be close to 1.

Our model’s coefficient is 0.88, which is quite close to 1.

- The slope of regression line should be close to 1.

For our model, the slope is 0.8, which is again close to 1.

Thus, our model doesn’t completely satisfy the Golbraikh–Tropsha criteria, but comes very close to it.

¹ Golbraikh, A., & Tropsha, A. (2002). Beware of q^2 ! Journal of Molecular Graphics and Modelling, 20(4), 269–276. doi:10.1016/s1093-3263(01)00123-1