QSAR model for BSEP inhibitor (v1.0)



ProtoADME

ProtoADME is a computational (in silico) tool focused on the prediction of endpoints related with the ADME (Absortion, Distribution, Metabolism and Excretion) of chemical substances.

Endpoint

Toxicokinetic: BSEP inhibitor

The bile salt export pump (BSEP) actively transports conjugated monovalent bile acids from the hepatocytes into the bile. This facilitates the formation of micelles and promotes digestion and absorption of dietary fat. Inhibition of BSEP leads to decreased bile flow and accumulation of cytotoxic bile salts in the liver.

Metrics

Training set

experimentai values	QSAR predictions		
	Non-inhibitor	Inhibitor	
Non-inhibitor	256	43	
Inhibitor	12	133	

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Experimental values	QSAR predictions		
	Non-inhibitor	Inhibitor	
Non-inhibitor	84	16	
Inhibitor	8	43	

Parameters	Training	Validation
Accuracy	0.88	0.84
Sensitivity / recall	0.92	0.84
Specificity	0.86	0.84
Precision	0.76	0.73
Negative predictive value	0.96	0.91
F-score	0.83	0.78
Matthews Correlation Coefficient	0.74	0.66
Critical Success Index	0.71	0.64
Area under the ROC	0.89	0.84

ProtoADME is part of



ProtoPRED platform allows the easy, fast and user-friendly prediction of different properties of chemical compounds, using proprietary (Q)SAR models





