Therapeutic Targets Database



QSAR Model

Target Name	CYP3A4
Target TTD ID	TTDS00418

Target Species	Human		
Chemical Type	Azole compounds		
Mode of Action	Inhibitor		
QSAR Model 1	Human CYP3A4: pIC_{50} (Inhibitory activity) = $1.37(\pm 0.34)\log P - 2.31(\pm 0.79)\log (\beta P + 1)$ $-1.59(\pm 0.77)I_{OH} + 2.45(\pm 0.83)$ $n = 17, r = 0.927, s = 0.609, F = 26.7, \log \beta = -3.87, \log P_{opt} = 4.03,$ Outlier = flusilazole (6)		
QSAR Model 2	Human CYP3A4: $pIC_{50}(Inhibitory) = 1.35(\pm 0.41)pIC_{50}(Binding) -1.07(\pm 0.82)I_{OH}$ $-1.79(\pm 2.21)$ n = 18, r = 0.875, s = 0.744, F = 24.4		
Molecular Descriptor	Access the following web-servers to compute molecular descriptors: MoDel and e-dragon P is the 1-octanol-water partition coefficient; β is the value related to the location of the optimum log P value. X is an indicator variable taking the value of unity for the presence and zero for the absence of certain structural features. E_{HOMO} is the energy level of HOMO. I_{azole} is the indicator variable for unity for imidazoles and zero for triazoles. I_{OH} is the indicator variable taking unity and zero for compounds with and without a hydroxyl group in the molecule.		
Reference	Quantitative Structure – Activity Relationship for Inhibition of CYP2B6 and CYP3A4 by Azole Compounds – Comparison with Their Binding Affinity. <i>QSAR Comb. Sci.</i> 28, 2009, No. 6-7, 629 –		

636	

Target Species	Rat		
Chemical Type	Azole compounds		
Mode of Action	Inhibitor		
QSAR Model 1	Rat CYP3A: $pIC_{50}(Binding) = 1.07(\pm 0.11)\log P - 1.34(\pm 0.19)\log (\beta P + 1)$ $+0.23(\pm 0.20)I_{azole} + 2.90(\pm 0.29)$ $n = 18, r = 0.985, s = 0.185, F = 157, \log \beta = -3.12, \log P_{opt} = 3.73$		
QSAR Model 2	Rat CYP3A: $pIC_{50}(Binding) = 1.01(\pm 0.11)\log P - 1.25(\pm 0.19)\log (\beta P + 1)$ $+7.03(\pm 5.29)E_{HOMO} + 5.48(\pm 1.85)$ $n = 18, r = 0.987, s = 0.179, F = 169, \log \beta = -3.15, \log P_{opt} = 3.76$		
Molecular Descriptor	Access the following web-servers to compute molecular descriptors: <u>MoDel</u> and <u>e-dragon</u> <i>P</i> is the 1-octanol-water partition coefficient; β is the value related to the location of the optimum log <i>P</i> value. <i>X</i> is an indicator variable taking the value of unity for the presence and zero for the absence of certain structural features. <i>E</i> _{HOMO} is the energy level of HOMO. <i>I</i> _{azole} is the indicator variable for unity for imidazoles and zero for triazoles. <i>I</i> _{OH} is the indicator variable taking unity and zero for compounds with and without a hydroxyl group in the molecule.		
Reference	Quantitative Structure – Activity Relationship for Inhibition of CYP2B6 and CYP3A4 by Azole Compounds – Comparison with Their Binding Affinity. <i>QSAR Comb. Sci.</i> 28, 2009, No. 6-7, 629 – 636		