Therapeutic Targets Database



QSAR Model

Target Name	Transcription factor $NF - \kappa B$
Target TTD ID	TTDR00250

Target Species	Human
Chemical Type	Sesquiterpene lactones
Mode of Action	Inhibitor
QSAR Model 1	$ pIC_{100} = 0.239 \ (\pm 0.054) \ \text{UNC} + 0.431 \ (\pm 0.164) \ \text{ML} + 0.648 \ (\pm 0.143) \ \text{EA} + 0.050 \ (\pm 0.023) \ \text{SH3} - 2.861 \ (\pm 0.169) \ n = 103, R = 0.772, R^2 = 0.596, s = 0.381, F = 36.2, P < 0.0001 $
QSAR Model 2	$ pIC_{100} = 0.271 \ (\pm 0.068) \ \text{UNC} + 0.452 \ (\pm 0.226) \ \text{ML} + 0.518 \ (\pm 0.163) \ \text{EA} - 2.597 \ (\pm 0.224) \\ n = 60, R = 0.779, R^2 = 0.607, s = 0.329. \ F = 28.8, P < 0.0001 $
QSAR Model 3	$ pIC_{100} = 1.001 \ (\pm 0.256) \ ML + 0.327 \ (\pm 0.082) \ UNA + 0.977 \ (\pm 0.162) \ EA - 2.913 \ (\pm 0.261) \\ n = 44, R = 0.848, R^2 = 0.718, s = 0.251, F = 34.0, P < 0.0001 $
QSAR Model 4	$ pIC_{100} = 0.239 \ (\pm 0.061) \ \text{UNC} + 0.913 \ (\pm 0.262) \ \text{ML} + 0.599 \ (\pm 0.218) \ \text{EA} - 2.983 \ (\pm 0.262) \ n = 44, R = 0.846, R^2 = 0.715, s = 0.252, \ F = 33.5, P < 0.0001 $
QSAR Model 5	$ pIC_{100} = 1.012 (\pm 0.206) \text{ ML} + 0.347 (\pm 0.071) \text{ UNA} + 0.933 (\pm 0.134) \text{ EA} - 2.893 (\pm 0.210) $ $ n = 37, R = 0.904, R^2 = 0.812, s = 0.201, F = 49.06, P < 0.0001 $
QSAR Model 6	
QSAR Model 7	$ pIC_{100} = 0.303 (\pm 0.088) \text{ UNC} + 0.639 (\pm 0.174) \text{ CN2} - 0.109 (\pm 0.027) \text{ MR} - 0.346 (\pm 0.114) \text{ OH} + 0.053 (\pm 0.577) \text{ m} = 22, R = 0.818, R^2 = 0.669, s = 0.282, F = 8.58, P = 0.0006 $
QSAR Model 8	$ pIC_{100} = 0.984 \ (\pm 0.097) \ \text{UNC} - 1.233 \ (\pm 0.286) \text{HOM} - 26.574 \ (\pm 3.052) \\ n = 9, R = 0.975, R^2 = 0.950, \text{ s} = 0.128, \text{F} = 57.30, P = 0.0001 $
QSAR Model 9	$ pIC_{100} = -1.067 \ (\pm 0.325) \ SH1 + 1.862 \ (\pm 0.531) \ SH2 + 0.804 \ (\pm 0.286) \ ATOM + 1.195 \ (\pm 1.153) \\ n = 10, R = 0.860, \ R^2 = 0.740, \ s = 0.145, \ F = 5.70, \ P = 0.0343 $

	Access the following web-servers to compute molecular descriptors: MoDel and e-dragon
	The Calculated Descriptors and the Abbreviation Used for the Calculations:
	Structural descriptor (Abbreviation): Total number of α , β -unsaturated carbonyl structures in the
Molecular	molecule (sum of ML, ENON and ACYL) (UNC); α -methylene- γ -lactone (ML); Conjugated ester
Descriptor	groups (UNA); Conjugated keto or aldehyde functions (ENONE); Number of oxygen atoms
	(ATOM); Number of hydroxyl groups (OH); Octanol water partition coefficient (LOGP); Electron
	affinity (EA); Dipole moment (DIPOL); Molar refractivity (MR); Connectivity Indices (CN0-2);
	Shape Indices (SH1-3); Highest occupied molecular orbital (HOMO); Lowest unoccupied molecular
	orbitals (LUMO1-3).
Reference	Quantitative Structure-Activity Relationship of Sesquiterpene Lactones as Inhibitors of the
	Transcription Factor NF – κB. J. Med. Chem. 2004, 47, 6042-6054