## **Therapeutic Targets Database**



QSAR Model

Target Name	Proto-oncogene tyrosine-protein kinase LCK
Target TTD ID	TTDS00407

Target Species	Human
Chemical Type	Flavonoid derivatives
Mode of Action	Inhibitors
QSAR Model 1	$\log 1/IC_{50} = 0.1586(\pm 0.0784)\log P - 0.0799(\pm 0.0268)He + 0.6499 \times (\pm 0.2148)I_{NH} + 2.3047$ n = 31, r = 0.8257, Se = 0.4067, F = 19.283, Q = 2.030
QSAR Model 2	$log1/IC_{50} = 0.2071(\pm 0.0895)logP - 0.1231(\pm 0.0473)He - 0.3169(\pm 0.2866)I_1 + 2.0029 + 0.5420(\pm 0.2351)I_{NH} n = 31, r = 0.8343, Se = 0.4050, F = 14.887, Q = 2.06$
QSAR Model 3	$\log 1/IC_{50} = 0.1537(\pm 0.0940)\log P - 0.0803(\pm 0.0165)He - 0.4602(\pm 0.1711)I_3 + 2.8095$ n = 39, r = 0.7948, Se = 0.4815, F = 17.125, Q = 1.650
QSAR Model 4	$log1/IC_{50} = 0.1436(\pm 0.0942)logP - 0.0739 \times (\pm 0.0174)He - 0.4184(\pm 0.1748)I_3 + 2.7485 + 0.2110(\pm 0.1930)I_{OH} n = 39, r = 0.8036, Se = 0.48, F = 13.677, Q = 1.674$
QSAR Model 5	$\log 1/IC_{50} = 0.2214(\pm 0.0385)\log P - 0.0901(\pm 0.0102)He - 0.4746(\pm 0.1252)I_3 + 2.8080$ n = 70, r = 0.7896, Se = 0.4603, F = 34.225, Q = 1.710
QSAR Model 6	$log1/IC_{50} = 0.2360(\pm 0.0375)logP - 0.0924(\pm 0.0098)He - 0.3664(\pm 0.1283)I_3 + 2.6814 + 0.3205(\pm 0.1309)I_{NH} n = 70, r = 0.8107, Se = 0.4428, F = 29.238, Q = 1.83$
QSAR Model 7	$\log 1/IC_{50} = 0.0497(\pm 0.0068)$ He $-0.9622(\pm 0.1159)I_{NO_2} - 0.4557(\pm 0.1172)I_{OMe} + 2.9154$ n = 104, r = 0.7783, Se = 0.4611, F = 48.148, Q = 1.688

QSAR Model 8	$log1/IC_{50} = 0.1221(\pm 0.0292)logP - 0.0595(\pm 0.0077)He + 0.5677(\pm 0.1215)I_{OH} + 2.4257 + 0.7633(\pm 0.1145)I_{NH} n = 104, r = 0.8032, Se = 0.4398, F = 42.267, Q = 1.826$
QSAR Model 9	$log1/IC_{50} = 0.1300(\pm 0.0291)logP - 0.0568(\pm 0.0077)He - 0.188(\pm 0.0999)I_3 + 2.5666$ $+ 0.5591(\pm 0.1199)I_{OH} + 0.7137(\pm 0.1159)I_{NH}$ $n = 104, r = 0.8115, Se = 0.4337, F = 35.485, Q = 1.870$
Molecular Descriptor	Access the following web-servers to compute molecular descriptors: <u>MoDel</u> and <u>e-dragon</u> I1 is used for presence of –OH group at any position by 1; I3, indicates presence of substituents at R3position by I3=1 otherwise zero; INH indicates presence of amino group by INH=1, otherwise 0; I1 indicates presence of –OH group at any position by I1=1, otherwise 0; IOH indicates presence of – OH group on phenyl ring by IOH=1 otherwise 0; INO2 indicates presence of nitro group by INO2 =1, otherwise 0; IOMe indicates presence of methoxy group by IOMe=1, otherwise 0; Hydrophobic parameter (logP) and hydration energy (He); n is the number of compounds; Se is the standard error of estimation; r is the correlation coefficient; F is the F-static and Q is the quality factor=R/Se.
Reference	QSAR study of flavonoid derivatives as p56lck tyrosin kinase inhibitors. <i>Bioorganic &amp; Medicinal</i> <i>Chemistry</i> 12 (2004) 1209–1214