Therapeutic Targets Database



QSAR Model

Target Name	VCAM-1
Target TTD ID	TTDR00453

Target Species	Human
Chemical Type	Piperazinyl phenylalanine derivatives
Mode of Action	Inhibitor
QSAR Model 1	-logIC ₅₀ = [2.93] + SaaNEindex [-0.037] + SsClcount [0.77] + 4path count [0.064] + SlogP [-0.69]
	n=21, r=0.93, r ² =0.8567, r ² se=0.25, F=33.89, F tabulated=7.46, q ² =0.76, q^{2} _se=0.33, pred_r ² =0.42, pred_r ² se=0.24, degree of freedom=17
Molecular Descriptor	Access the following web-servers to compute molecular descriptors: <u>MoDel</u> and <u>e-dragon</u> SaaNEindex, electro-topological state indices for number of nitrogen atoms connected with two aromatic double bonds; SsClcount, electro-topological parameter defining total number of chlorine atoms connected with one single bond; 4pathcount, topological parameter signifying total number of fragments of fourth order (four bond path) in compound; SlogP, signify log of the octanol/water partition coefficient; n, number of samples, r, correlation coefficient; r^2 se, standard error of squared correlation coefficient; F, F-ratio; F tabulated, tabulated F-ratio; q^2 , square of the correlation coefficient; q^2 se, standard error of square of the correlation coefficient; pred r^2 , predicted squared correlation coefficient; pred r^2 se, standard error of predicted squared correlation coefficient
Reference	Quantitative Structure Activity Relationship Studies of Piperazinyl Phenylalanine Derivatives as VLA-4/VCAM-1 Inhibitors. <i>Medicinal Chemistry</i> , 2009, <i>5</i> , 446-454