

Target Name	Cyclooxygenase (COX)
Target TTD ID	TTDS00039

Target Species	Human
Chemical Type	4',5 di-substituted biphenyl acetic acid molecules
Mode of Action	Inhibitor
QSAR Model 1	$\log \text{CPE}\% = -0.266 - 0.0085f_3^N + 0.0488f_{10}^E + 0.0429 \frac{\mu_k^+}{\mu_k^-} (k = 3)$ $r^2 = 0.854 \quad q^2 = 0.776 \quad F = 27.257 \quad \sigma = 0.127$
QSAR Model 2	$\log \text{CPE}\% = 6.806 - 0.01474f_3^N + 0.050f_{10}^E + 0.634 \text{NICS}(1)$ $+ 0.024 \frac{\mu_k^+}{\mu_k^-} (k = 3)$ $r^2 = 0.916 \quad q^2 = 0.867 \quad F = 35.041 \quad \sigma = 0.100$
QSAR Model 3	$\log \text{CPE}\% = 9.762 - 0.019f_3^N + 0.036f_{10}^E + 0.818 \text{NICS}(1)$ $r^2 = 0.895 \quad q^2 = 0.774 \quad F = 38.818 \quad \sigma = 0.108$
Molecular Descriptor	<p>Access the following web-servers to compute molecular descriptors: <a href="#">MoDel</a> and <a href="#">e-dragon</a></p> <p>Chemical potential (<math>\mu</math>): a useful global index of reactivity in atoms, molecules and clusters. <math>f_k^a</math>: a single value around each atomic site that characterize the atomic contribution in a molecule.</p> <p><math>\log \text{CPE}\%</math> refers to the anti-inflammatory activity. <math>f_3^N</math>, <math>f_{10}^E</math> and <math>\frac{\mu_k^+}{\mu_k^-} (k=3)</math>: weighted nucleophilic frontier electron densities on atom 3, weighted electrophilic frontier electron densities on atom 10 and relative nucleophilic chemical potential on atom 3, respectively. <math>r^2</math>: the correlation coefficient. <math>q^2</math>: the cross-</p>

	validated correlation coefficient using leave-one-out (LOO) method. $\sigma$ : the standard error. F: F-test value.
<b>Reference</b>	Towards the design of Cyclooxygenase (COX) inhibitors based on 4',5 di-substituted biphenyl acetic acid molecules: a QSAR study with a new DFT based descriptor – nucleus independent chemical shift. <i>J Mol Model</i> (2009) 15:1221–1228