Therapeutic Targets Database



QSAR Model

Target Name	5-HT3 receptor
Target TTD ID	TTDS00106

Target Species	Human
Chemical Type	Arylguanidines
Mode of Action	Binder
QSAR Model 1	$pK_i = 2.30(\pm 0.81)\sigma_m + 5.53$ n = 8, r = 0.760, F = 8.17
Molecular Descriptor	Access the following web-servers to compute molecular descriptors: <u>MoDel</u> and <u>e-dragon</u> No correlation was found between 5-HT3 affinity (pKi) and p (r=0.132), but a better relationship exists with sm (r=0.760) (eq 1). Interestingly, if compound 4 is excluded, r=0.870
Reference	Arylguanidine and Arylbiguanide Binding at 5-HT3 Serotonin Receptors: A QSAR Study. Bioorganic & Medicinal Chemistry 11 (2003) 4449–4454

Target Species	Human
Chemical Type	3-monosubstituted compounds
Mode of Action	Binder
QSAR Model 1	$pK_i = 2.30(\pm 0.81)\sigma_m + 5.53$ n = 8, r = 0.760, F = 8.17
Molecular Descriptor	Access the following web-servers to compute molecular descriptors: <u>MoDel</u> and <u>e-dragon</u> No correlation was found between 5-HT3 affinity (pKi) and p (r=0.132), but a better relationship

	exists with sm (r=0.760) (eq 1). Interestingly, if compound 4 is excluded, r=0.870
Reference	Arylguanidine and Arylbiguanide Binding at 5-HT3 Serotonin Receptors: A QSAR Study.
	Bioorganic & Medicinal Chemistry 11 (2003) 4449–4454

Target Species	Human
Chemical Type	Arylguanidines
Mode of Action	Binder
QSAR Model 1	$pK_i = 1.80(\pm 0.23)\pi_4 + 5.69$ n = 7, r = 0.906, F = 23.0
Molecular Descriptor	Access the following web-servers to compute molecular descriptors: <u>MoDel</u> and <u>e-dragon</u> No correlation was found between 5-HT3 affinity (pKi) and p (r=0.132), but a better relationship exists with sm (r=0.760) (eq 1). Interestingly, if compound 4 is excluded, r=0.870
Reference	Arylguanidine and Arylbiguanide Binding at 5-HT3 Serotonin Receptors: A QSAR Study. Bioorganic & Medicinal Chemistry 11 (2003) 4449–4454

Target Species	Human
Chemical Type	4-monosubstituted compounds
Mode of Action	Binder
QSAR	$pK_i = 1.80(\pm 0.23)\pi_4 + 5.69$
Model 1	n = 7, r = 0.906, F = 23.0
Molecular	Access the following web-servers to compute molecular descriptors: MoDel and e-dragon
Descriptor	No correlation was found between 5-HT3 affinity (pKi) and p (r=0.132), but a better relationship exists with sm (r=0.760) (eq 1). Interestingly, if compound 4 is excluded, r=0.870

Reference	Arylguanidine and Arylbiguanide Binding at 5-HT3 Serotonin Receptors: A QSAR Study.
	Bioorganic & Medicinal Chemistry 11 (2003) 4449–4454

Target Species	Human
Chemical Type	Di-substituted compounds
Mode of Action	Binder
QSAR Model 1	$pK_i = 2.59(\pm 0.59)\sigma_m + 1.38(\pm .24)\pi_4 + 5.52$ n = 18, r = 0.852, F = 19.9
Molecular Descriptor	Access the following web-servers to compute molecular descriptors: <u>MoDel</u> and <u>e-dragon</u> No correlation was found between 5-HT3 affinity (pKi) and p (r=0.132), but a better relationship exists with sm (r=0.760) (eq 1). Interestingly, if compound 4 is excluded, r=0.870.
Reference	Arylguanidine and Arylbiguanide Binding at 5-HT3 Serotonin Receptors: A QSAR Study. Bioorganic & Medicinal Chemistry 11 (2003) 4449–4454

Target Species	Human
Chemical Type	Lipophilic 3-position substituents
Mode of Action	Binder
QSAR Model 1	$pK_i = 2.48(\pm 0.90)\sigma_m + 1.27(\pm 0.25)\pi_4 - 2.40(\pm .55)I_4 + 5.60$ n = 21, r = 0.824, F = 11.9
QSAR Model 2	$pK_{i} = -0.68(\pm 0.15)P + 2.04(\pm 0.84)^{8}\chi_{P} + 0.39(\pm 0.05)SsCl + 0.012(\pm 0.008)Vol + 8.53$ n = 24, r = 0.908, F = 22.3
QSAR Model 3	$pK_{i} = -125.9(\pm 27.9)\text{SpcPol} + 1.69(\pm 0.44)^{8}\chi_{P} + 0.39(\pm 0.05)\text{SsCl} + 8.53$

	n = 24, r = 0.902 $F = 29.2$
	Access the following web-servers to compute molecular descriptors: MoDel and e-dragon
	I4 was set at 0 for 4-position substituents with a Verloop B2 value 16 of <2.5, and I4=1 for substituents where $P_2 > 2.5$, 3 position substituent 4 position substituent P is the polarizability of
Molecular	the molecule, $8\chi_P$ is the simple 8th order chi index, SsCl is the sum of all (–Cl) E-State values in
Descriptor	the molecule, and Vol=the molecular volume.
	The $8\chi_P$ descriptor is a count of paths of eight bonds; substitution at one of the aryl meta positions
	by a monoatomic substituent would be expected to add an additional path. specific
	polarizability (SpcPol)=P/Vol
Poforonco	Arylguanidine and Arylbiguanide Binding at 5-HT3 Serotonin Receptors: A QSAR Study.
Reference	Bioorganic & Medicinal Chemistry 11 (2003) 4449-4454

Target Species	Human
Chemical Type	Lipophilic 4-position substituents
Mode of Action	Binder
QSAR Model 1	$pK_{i} = 2.48(\pm 0.90)\sigma_{m} + 1.27(\pm 0.25)\pi_{4} - 2.40(\pm .55)I_{4} + 5.60$ n = 21, r = 0.824, F = 11.9
QSAR Model 2	$pK_{i} = -0.68(\pm 0.15)P + 2.04(\pm 0.84)^{8}\chi_{P} + 0.39(\pm 0.05)SsCl + 0.012(\pm 0.008)Vol + 8.53$ n = 24, r = 0.908, F = 22.3
QSAR Model 3	$pK_{i} = -125.9(\pm 27.9)\text{SpcPol} + 1.69(\pm 0.44)^{8}\chi_{P} + 0.39(\pm 0.05)\text{SsCl} + 8.53$ $n = 24, r = 0.902 F = 29.2$
Molecular Descriptor	Access the following web-servers to compute molecular descriptors: <u>MoDel</u> and <u>e-dragon</u> I4 was set at 0 for 4-position substituents with a Verloop B2 value16 of <2.5, and I4=1 for substituents where B2 >2.5. 3-position substituent 4-position substituent P is the polarizability of the molecule, $8\chi_P$ is the simple 8th order chi index, SsCl is the sum of all (–Cl) E-State values in

	the molecule, and Vol=the molecular volume.
	The $8\chi_P$ descriptor is a count of paths of eight bonds; substitution at one of the aryl meta positions by a monoatomic substituent would be expected to add an additional path. specific
	polarizability (SpcPol)=P/Vol
Reference	Arylguanidine and Arylbiguanide Binding at 5-HT3 Serotonin Receptors: A QSAR Study. <i>Bioorganic & Medicinal Chemistry</i> 11 (2003) 4449–4454

Target Species	Human
Chemical Type	Arylbiguanide derivatives
Mode of Action	Binder
QSAR Model 1	$pK_{i} = -0.64(\pm 0.14)P + 2.74(\pm 0.53)^{8}\chi_{P} + 0.37(\pm 0.05)SsCl + 8.03$
	n = 35, r = 0.907, F = 47.7
QSAR Model 2	$pK_{i} = -0.63(\pm 0.13)P + 2.75(\pm 0.43)^{8}\chi_{P} + 0.37(\pm 0.04)SsCl + 8.03$
	n = 35, r = 0.907, F = 47.7
QSAR Model 3	$pK_i = 1.81(\pm 0.28)\Sigma\sigma + 1.02(\pm 0.22)\pi_3 + 6.19$
	n = 20, r = 0.924, F = 49.3
Molecular Descriptor	Access the following web-servers to compute molecular descriptors: MoDel and e-dragon
	The $8\chi_P$ descriptor is a count of paths of eight bonds; substitution at one of the aryl meta positions
	by a monoatomic substituent would be expected to add an additional path. specific
	polarizability (SpcPol)=P/Vol
Reference	Arylguanidine and Arylbiguanide Binding at 5-HT3 Serotonin Receptors: A QSAR Study.
	Bioorganic & Medicinal Chemistry 11 (2003) 4449–4454

Target Species	Human
Chemical	Arylguanidine derivatives

Туре	
Mode of Action	Binder
QSAR Model 1	$pK_i = -0.64(\pm 0.14)P + 2.74(\pm 0.53)^8 \chi_P + 0.37(\pm 0.05)SsCl + 8.03$ n = 35, r = 0.907, F = 47.7
QSAR Model 2	$pK_{i} = -0.63(\pm 0.13)P + 2.75(\pm 0.43)^{8}\chi_{P} + 0.37(\pm 0.04)SsCl + 8.03$ n = 35, r = 0.907, F = 47.7
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Molecular Descriptor	Access the following web-servers to compute molecular descriptors: <u>MoDel</u> and <u>e-dragon</u> The $8\chi_P$ descriptor is a count of paths of eight bonds; substitution at one of the aryl meta positions by a monoatomic substituent would be expected to add an additional path. specific polarizability (SpcPol)=P/Vol
Reference	Arylguanidine and Arylbiguanide Binding at 5-HT3 Serotonin Receptors: A QSAR Study. Bioorganic & Medicinal Chemistry 11 (2003) 4449–4454