



In Silico Study of Neurokinin-1 Receptor Antagonists by QSAR Modeling

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ABSTRACT

QSAR studies were performed on a series neurokinin-1 receptor. neurokinin-1 receptor have been analyzed in relation to their physicochemical and molecular properties. On the basis of QSAR studies presented here, the coefficients of Id, IOR, MR, MW, ST, MV, Pz are positive and therefore it may be suggested that denser, bulkier, more polar substituents with more vertices should have positive influence on the activity of NK₁ receptor. and the positive coefficients of $^0\chi$, $^1\chi$, $^2\chi$, $^3\chi$, $^4\chi$, $^5\chi$, W, W_A, and BAC parameters indicate that compounds, which are sterically hindered, are preferable for the activity of NK₁ receptor. The coefficient of hydrophobic parameter (LogP) was found to be negative and this clearly indicates that this parameter has negative influence in determining the activity, also the presence of  group at R₁-position and (-CH₃) group at Z-position should be preferable and the presence of  group at Y-position causes a substantial decrease in activity. The results are critically discussed on the basis of regression data and cross validation techniques. Poglani factor Q and the results of LOO (leave one out) method confirms the reliability and predictability of the proposed models.

Keywords: QSAR, BAC, PRESS/SSY

I. INTRODUCTION

Neurokinin (NK) receptors belong to G-protein coupled receptor and can be divided into three subtypes: NK₁, NK₂ and NK₃. They are located on submucosal enterocytes through the release of the both cholinergic and non cholinergic neurotransmitters. These NK₁ receptors have been implicated in many pathophysiological effects related to pulmonary dysfunction and many mental disorders such as anxiety and depression¹. The current study represents the QSAR analysis of a series of NK₁ receptor antagonists² (K_i which represents NK₁ receptor affinities are reported in the literature) using a number of structural parameters alongwith different indicator parameters at different substitution sites.

II. Experimental

Hansch analysis³ was used for multiparametric regression analysis. The multiparametric regression analysis used to derive the correlation was executed



with the SPSS (7.5) programme. For QSAR studies, different analogues of parent structure and values of biological activity (pK_i) are reported in Table1. The structural parameters used are equalized electronegativity (X_{eq})⁴, Polarizability (Pz) as electronic parameters, Partition coefficient (LogP)⁵ as hydrophobic parameter, Density (D), Index of refraction (IOR), Molar refractivity (MR), Molecular weight (MW), Molar Volume (MV), Parachor (Pc), Surface tension (ST), Information Theoretic Index (Id), Weiner Index (W), Mean Weiner Index (W_A), Balaban Index (J), Balaban centric index (BAC) and Conectivity indices($^0\chi$, $^1\chi$, $^2\chi$, $^3\chi$, $^4\chi$, $^5\chi$)⁶ as steric parameters. The structural parameters are calculated with the help of ACD Lab⁶ and DRAGON⁷ software. A total of 21 descriptors were chosen for the QSAR analysis are listed in Table-2.

III. Monoparametric Models

In silico study of neurokinin-1 receptor antagonists by QSAR Modeling

Where n is the number of data points, R^8 is the correlation coefficient, R^2 is coefficient of determination, R^2_A represents adjusted R^2 or explained variance in activity that can be accounted by the equation, SE is the standard error of estimate, F is the variance ratio between observed and calculated activities and data with in parenthesis are confidence interval at 95% level. On the basis of statistical parameters, it can be said that all the above equations in their present form are insignificant. All the physicochemical parameters and their square terms were taken together to check the parabolic relationship which was found to be absent.



The autocorrelation between all the structural parameters is shown in Table-4. A close study of this table makes it clear that there are a number of parameters which are orthogonal and, therefore, can be used together in multiparametric regression studies. Multiparametric Modeling⁹ Using Indicator parameters-

In order to study the role of specific substituents at a particular position, three indicator parameters IR for  at R₁- position, IY for  at Y-position and IZ for (-CH₃) at Z-position were used.. The values for these dummy parameters are also listed in Table-1. It is taken one for that particular group or substituent and zero for all such cases were that group or substituent is absent. Different combination of physicochemical parameters and indicator parameters which have no autocorrelation were used and the statistically significant models obtained are summarized in Table-5 in the form of equations.

The statistical data listed in table show that all the given equations in Table-5 are significant and the set of physicochemical parameters and indicator parameters used may be useful for the future drug designing.

On the basis of QSAR studies presented above (Table-5), the coefficients of Id, IOR, MR, MW, ST, MV, Pz are positive and therefore it may be suggested that denser, bulkier, more polar substituents with more vertices should have positive influence on the activity of NK₁ receptor. Further, positive coefficients of $\chi^0, \chi^1, \chi^2, \chi^3, \chi^4, \chi^5$, W, W_A, and BAC parameters indicate that compounds, which are sterically hindered, are preferable for the activity of NK₁ receptor. The coefficient of hydrophobic parameter (Log P) was found to be negative which clearly indicates that this

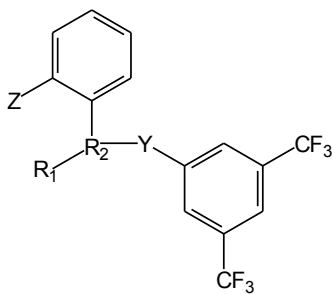
parameter has negative influence in determining the activity.

A very interesting and striking feature of the QSAR modeling¹⁰⁻¹⁶ presented in this section reveals that everywhere the coefficient of IR and IZ are positive and for IY the coefficient is negative. This can be explained by suggesting that the presence of  group at R₁- position and (-CH₃) group at Z-position should be preferable and at the same time while some other substituent may be tolerated. The presence of  group at Y-position definitely causes a substantial decrease in activity.

In order to examine the relative potential of models, predictive correlation coefficient (R^2_{Pred})¹¹ were estimated by plotting graphs between observed and calculated pKi values obtained with the help of eqs 1 and eqs 2. The comparison between observed and predicted activities is listed in Table- 6. Predictive ability was also evaluated by the LOO¹² (Leave one out) cross-validation procedure. This method systematically removes one data point at a time a model is constructed on the basis of this reduce data set and is subsequently used to predict the removed sample. This procedure was repeated for all the points until a complete set of predicted values was obtained. Various cross-validation parameters calculated for the proposed models are presented in Table- 8. PRESS (Predicted residual sum of squares) appears to be the most important cross-validation parameters accounting for good estimate of the real predictive error of the models. In case its value is less than SSY (Sum of the square of all response value), it will mean that the predictive power of the model is good and is not based upon chance and therefore, can be considered statistically significant. For a reasonable QSAR model¹³⁻¹⁷, PRESS/SSY should be smaller than 0.400. In the present case, the ratio PRESS/SSY ranges between 0.108-0.131 indicating that all the proposed models are reliable. The PSE as well as S_{PRESS} are good parameters to be used for discussing the uncertainty in prediction. The lower the value of these parameters, the better will be the predictive ability of the model.

The indication of the performance of the model was also obtained from R^2_{cv} (the overall predictive ability). The highest R^2_{cv} (0.892) was found for QSAR model 2, indicating that it has an outstanding predictive power.

Table – 1. Biological activities and indicator Parameters of Neurokinin-1 Receptor Antagonists



S. No	Substituents		R ₂	Y	Observed pK _i	Indicator Parameters		
	Z	R ₁				IZ	IR	IY
1.	H	H			7.2924	0	0	0
2.	CH ₃	H			7.770	1	0	0
3.	Cl	H			7.658	0	0	0
4.	Br	H			7.347	0	0	0
5.	H	H			6.530	0	0	0
6.	H	H			6.250	0	0	0
7.	H	H			7.168	0	0	0
8.	H	H			5.860	0	0	1
9.	H	H			6.140	0	0	0
10.	H	H			6.430	0	0	0
11.	H	H			6.650	0	0	0
12.	CH ₃	H			8.854	1	0	0
13.	CH ₃	H			8.0710	1	0	0
14.	CH ₃	H			8.469	1	0	0


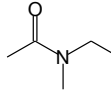

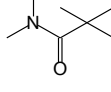

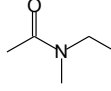

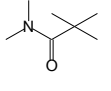
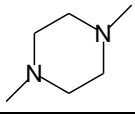

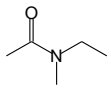
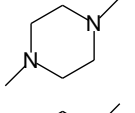

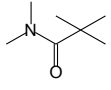
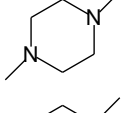

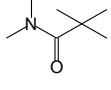
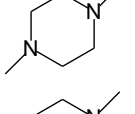

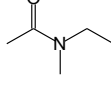
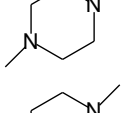

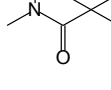
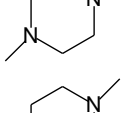

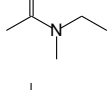
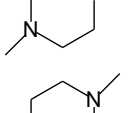

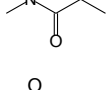
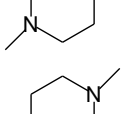

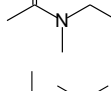
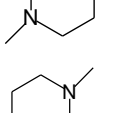

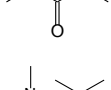
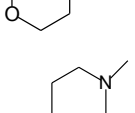

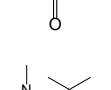
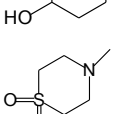

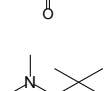
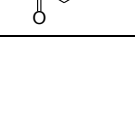

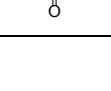
S. No	Substituents				Observed pKi	Indicator Parameters		
	Z	R ₁	R ₂	Y		IZ	IR	IY
15.	CH ₃	H			8.337	1	0	0
16.	CH ₃	H			4.402	1	0	0
17.	CH ₃	H			8.420	1	0	0
18.	CH ₃	H			7.776	1	0	0
19.	CH ₃				9.237	1	0	0
20.	CH ₃				9.022	1	0	0
21.	CH ₃				8.770	1	0	0
22.	CH ₃				8.208	1	0	0
23.	CH ₃				8.699	1	0	0
24.	CH ₃				8.456	1	0	0
25.	CH ₃				8.959	1	0	0
26.	CH ₃				8.770	1	0	0
27.	CH ₃				9.357	1	0	0
28.	CH ₃				9.699	1	1	0
29.	CH ₃				9.301	1	0	0
30.	CH ₃				9.097	1	0	0

Table 5. Multiparametric Models Using Indicator Parameters

S. No.	Equation	n	R	R ²	R _A	SE	F _{5,24}
1.	pKi= 3.754 (±3.601) D + 2.377 (±1.156) Id + 1.313 (± 0.419) IZ - 1.078 (±0.809) IY + 0.789 (±0.795) IR - 9.557	30	0.950	0.903	0.883	0.370	44.595
2.	pKi= -8.39 × 10 ⁻² (±4.342) D + 39.466 (±21.259) IOR + 1.415 (± 0.413) IZ - 1.225 (±0.854) IY + 1.001 (±0.817) IR - 53.269	30	0.946	0.894	0.872	0.386	40.667
3.	pKi= 4.005 (±4.481) D - 1.433 (±1.675) J + 1.796 (± 0.405) IZ - 0.827 (±1.009) IY + 0.996 (±0.979) IR + 3.986	30	0.922	0.849	0.818	0.460	27.078
4.	pKi= 3.409 (±4.383) D + 8.853 × 10 ⁻² (±8.353 × 10 ⁻²) 1χ + 1.655 ± 0.437) IZ - 0.960 (±0.974) IY + 0.915 (±0.958) IR + 1.109	30	0.926	0.858	0.828	0.447	29.012
5.	pKi= 1.102 (±1.763) Id + 26.638 (±26.896) IOR + 1.282(± 0.418) IZ - 1.210 (±0.819) IY + 0.894 (±0.808) IR - 39.138	30	0.949	0.901	0.881	0.373	43.817

Table 6. Comparison between observed and predicted values for proposed models

PKi					
S.No.	Observed	Equation – 1		Equation - 2	
		Predicted	Residual	Predicted	Residual
1.	7.292	6.790	0.502	6.847	0.446
2.	7.770	8.119	-0.350	8.226	-0.457
3.	7.658	7.084	0.574	7.122	0.536
4.	7.347	7.493	-0.146	7.436	-0.889
5.	6.530	6.791	-0.261	6.493	0.037
6.	6.250	6.843	-0.592	7.239	-0.989
7.	7.168	6.840	0.327	6.847	0.321
8.	5.860	5.860	.000	5.860	0.000
9.	6.140	6.520	-0.380	6.611	-0.471
10.	6.430	6.577	-0.147	6.572	-0.143
11.	6.650	6.526	0.124	6.297	0.353
12.	8.854	8.750	0.104	8.736	0.117
13.	8.071	8.221	-0.150	8.265	-0.195
14.	8.469	8.267	0.202	8.265	0.203
15.	8.337	8.221	0.117	8.265	0.072
16.	8.402	8.267	0.135	8.265	0.136
17.	8.420	8.221	0.200	8.265	0.155
18.	7.796	8.267	-0.471	8.265	-0.469
19.	9.237	8.833	0.404	8.776	0.461
20.	9.022	8.876	0.146	8.776	0.247
21.	8.770	8.746	0.023	8.776	-0.006
22.	8.208	8.833	-0.625	8.776	-0.568
23.	8.699	8.876	-0.177	8.776	-0.077
24.	8.456	8.919	-0.483	8.815	-0.359
25.	8.959	9.312	0.647	8.815	0.144
26.	8.770	8.919	-0.149	8.815	-0.045
27.	9.357	8.955	0.401	8.815	0.542
28.	9.699	9.699	0.000	9.699	0.000
29.	9.301	9.124	0.177	9.208	0.093
30.	9.097	9.267	-0.170	9.090	0.007

Table 7. Some Statistical techniques prove predictive ability for proposed models

S.No.	Model No.	Q	K	E	PE	6PE
1.	1	2.503	0.326	67.4	0.013	0.078
2.	2	2.568	0.311	68.9	0.012	0.072
3.	3	2.451	0.326	67.4	0.013	0.078
4.	4	2.544	0.315	68.5	0.012	0.072
5.	5	2.327	0.341	65.9	0.014	0.084

Table 8. Cross validation parameters for proposed models

S.No.	Model No.	PRESS	SSY	PRESS/SSY	R ² _{cv}	PSE	S _{PRESS}
1.	1	3.568	30.224	0.118	0.882	0.345	0.378
2.	2	3.284	30.501	0.108	0.892	0.331	0.370
3.	3	3.567	30.224	0.118	0.882	0.345	0.386
4.	4	3.336	30.456	0.110	0.890	0.333	0.373
5.	5	3.908	29.884	0.131	0.869	0.361	0.404

IV. CONCLUSION

On the basis of above discussions, it becomes evident that multiparametric models represented by QSAR equations are statistically significant models and they are reliable and have good predictive ability and therefore it can be concluded that the coefficients of Id, IOR, D are positive and therefore denser, bulkier should have positive influence on the activity of NK₁ receptor whereas the positive coefficient of ¹χ parameter indicate that compounds, which are sterically hindered, are preferable for the activity of NK₁ receptors.

V. REFERENCES

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