Quantitative Structure Activity Relationship of Riluzole Series as Anticonvulsants

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24 Riluzole series, a blocker of excitatory amino acid (glutamic acid) mediated transmission, 2-benzothiazol-amines and 3-substituted 2-imino benzothiazolines have been subjected to quantitative structure activity relationship studies by step wise multiple regression analysis using steric, electronic descriptors. Result of this study has given 12 QSAR models and among all models one has statistically significant result (r > 0.97, F = 133.4899, $q^2 = 0.97753$). Descriptors used in this study are: polarizability, density, molar refractivity, molar volume, average mass, parachor. Predictive ability of proposed model was assessed on the basis of regression data and cross validation parameters. This validated model brings important structural insight to model better anticonvulsant containing benzothiazole nucleus and more potent than riluzole.

Key Words: QSAR, Riluzole series, Benzothiazoles analogs, Anticonvulsants, Antiglutamate.

INTRODUCTION

There is currently need for improved agents for the treatment of seizure disorders, since available drugs are effective in only 60-80% of epileptic patients^{1,2}. Absence seizures are well controlled in most cases but significant therapeutic advancement is required for the treatment of partial-complex (focal) seizures and generalized tonic-clonic (grand mal) epilepsy. In pharmacological experiments show that riluzole possessed anticonvulsant properties in different models including maximal electroshock in mice, convulsion induced by glutamic acid decarboxylase inhibitors in mice and rats, sound-induced convulsions in DBA/2 mice, photo sensitive epilepsy in baboons. However unlike benzodiazepines, barbiturates, valproic acid, riluzole was ineffective against seizures induced by pentylenetetrazole and picrotoxin. Riluzole has been shown to have interesting neuroprotective effects *in vitro*, protecting primary culture of rat cortical neurons against anoxic stress. Furthermore rat motoneurons were protected by riluzole from the excitotoxocity effects of glutamic acid

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uptake inhibitors. These studies suggested that riluzole may have neuroprotective effects in whole animal models of cerebral ischemia. Clinical studies have shown that riluzole can slow down the disease progression in amylotropic lateral sclerosis.

Thus, there is hope to find more potent anticonvulsant drug from riluzole series. The present study has concerned with bringing up new QSAR model to develop better anticonvulsant containing benzothiazole nucleus. QSAR study already reported in substituted 2-benzothiazolamine as sodium flux inhibitors series with anticonvulsant property. In the present study concerning with different descriptors and has developed statistically significant results (Table-1) when comparing to already reported QSAR model¹, however for this study compounds are taken from different source and 3-substituted imino benzothiazolines were used with their ED_{50} values (protection against glutamic acid evoked convulsion).

TABLE-1

EXPERIMENTAL

Anticonvulsant ED_{50} value of glutamic acid evoked convulsion values and their corresponding structures are given in Table-2. These values are work of Patrick Jiminet et al.². 3-Substituted imino benzothiazolines were taken in present studies. These ED_{50} values are converted into $log (ED_{50})$ and all the compounds in both training and test set were drawn in chem. Sketch physiochemical properties were calculated by ACD lab freeware release 8.00 version 8.17, only 2D descriptors were calculated from drawing the molecules in chemsketch. The following descriptors were calculated *viz.*, polarizability(pol), average mass (AM), parachor (Pc), molar refractivity (MR), molar volume (MV), surface tension (ST), index of refraction (η), density (D) and indicator parameter (I) (presence of 6 trifluoromethoxy group in benzothiazole $)^3$.

TABLE-2 6-TRIFLUOROMETHOXY 3-IMINO BENZOTHIAZOLINES

#compounds taken from reference no.2, Predicted by Equation no.12, the compound no superscripted by T are known as external validation set (not included in developing model), compound no superscripted by *are not included in study.

All the calculated parameters in the ACD labs freeware were considered as independent variables and biological activity [log ED₅₀] was taken as dependent variable. Stepwise regression analysis method was used to develop equations. All the calculated parameters are listed in Table-4. 2846 Gunakkunru *et al. Asian J. Chem.*

The statistical parameters were used to find out best regression model among all equation obtained in regression analysis coefficient of correlation (R), F-statistics (F), quality factor (Q) and standard error (Se). In addition to that the best QSAR equation further subjected to internal validation by leave-one-out (LOO) method, external validation by taking 5 compounds in test set, these compounds are,in compound number, superscripted with 'T' and cross validation process^{4,5}.

RESULTS AND DISCUSSION

When training set was subjected to stepwise regression multiple linear analysis, in order to develop α SAR model⁶⁻⁸.

 $log ED_{50} = 12.68 (\eta) + 17.08 (Pol) - 0.044 (ST) - 6.777 (MR) - 17.18$ (1) $R = 0.439417$, $Q = 1.156385$, $F = 0.897344$, $n = 20$

 $log ED_{50} = 8.871 (Pol) + 0.036 (ST) - 2.88 (D) - 3.534 (MR) + 4.346$ (2) $R = 0.762222$, $Q = 2.783585$, $F = 5.199512$, $n = 20$

 $log ED_{50} = 27.603 (Pol) - 10.855 (MR) - 1.930 (D) - 0.013 (Pc) + 4.733 (3)$ $R = 0.730839$, $Q = 2.531197$, $F = 4.299375$, $n = 20$

 $log ED_{50} = 8.158 (Pol) + 0.015 (ST) - 0.0131 (AM) - 3.198 (MR) + 1.329 (4)$ R= 0.859434, Q= 3.973936, F= 10.5973, n= 20

 $log ED_{50} = 5.9639 (\eta) + 0.0089 (ST) + 0.012 (MV) - 0.013 (AM) - 7.975 (5)$ $R = 0.855956$, $Q = 3.913436$, $F = 10.27709$, $n = 20$

log ED₅₀ = 5.682 (η) + 0.0045 (Pc) - 0.012 (AM) - 0.0038 (ST) - 6.914 (6) $R = 0.855597$, $Q = 3.907308$, $F = 10.24492$, $n = 20$

 $log ED_{50} = 0.0014$ (Pc) - 0.0061 (ST) - 0.003 (MV) - 0.789 (I) + 1.187 (7) $R = 0.951198$, $Q = 7.286867$, $F = 35.63159$, $n = 20$

 $log ED_{50} = 10.671 (D) + 0.0302 (Pc) - 0.058 (AM) - 0.074 (ST) - 10.372 (8)$ $R = 0.927453$, $Q = 5.863123$, $F = 23.06808$, $n = 20$

 $log ED_{50} = 3.78 (Pol) + 0.0999 (D) - 1.4956 (MR) - 0.795 (I) + 0.707 (9)$ $R = 0.95213$, $Q = 7.36288$, $F = 36.37885$, $n = 20$

$$
log ED50 = 1.563 (\eta) + 0.001 (AM) - 0.01 (ST) - 0.822 (I) - 1.247
$$
 (10)
R = 0.953991, Q = 7.521439, F = 37.96255, n = 20

$$
log ED_{50} = 1.062506 \text{ (}\eta\text{)} + 0.001 \text{ (AM)} - 0.01 \text{ (ST)} - 0.8148 \text{ (I)} - 0.44404 \qquad (11)
$$

R = 0.97557, Q = 10.43626, F = 64.0771, n = 18

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 $log ED_{50} = 0.5473 (\eta) + 0.001 (AM) - 0.002 (ST) - 0.7739 (I) + 0.094609 (12)$ $R = 0.9889$, $Q = 16.1808$, $F = 133.4899$, $n = 17$

Fig.1 Plot of observed log ED_{50} *vs.* predicted log ED_{50}

All the equations were screened for inter correlation with in descriptors by correlation matrix (Table-3). Internal validation was done by leaveout-one validation method. Among above 12 equations, equation 12th has good statistical significance; equation 1-5 did not comply with cross validation parameter. Hence equation 12 considered to be the best model for predicting the activity. The presence of outliers was tested and 3 outliers were removed equation 11 and 12 are best equation without outliers. Outliers are nothing but values which are higher than two times of standard deviation of predicted value. But simply getting good statistics is not only enough for dealing with biological studies when no proper predictivity is possible. In view of this quality factor Q was used in this study. Use of quality factor Q indicates that all the models under the referred condition have the best predictive power too. In this view we have under gone cross validation methodology for deciding predictive power of the proposed models.

Predicted residual sum of squares (PRESS) appears to be the most important cross-validation parameter. Its value less than SSY (sum of squares of response value) indicate that the model predicts better than chance and can be considered statistically significant. In this model PRESS << SSY indicates its predictive power better than chance. To be a reasonable QSAR model, PRESS/SSY $^{7-9}$ should be smaller than 0.4.

The satisfactory values of internal validation cross validation parameters (PRESS/SSY ratio < 0.3), standard deviation of prediction (SPRESS) 0.126, standard deviation of error of predictions (SDEP) 0.1098 are supporting the predictive ability of this model.

Present study shows that index of refraction (η) and average mass (AM) positively correlating with activity and indicator parameter (I) and surface tension (ST) are negatively correlating with activity. From the above model equation can be used to predict the activity for new models.

Conclusion

This study led to the identification of important physiochemical properties in explaining the variation in activity in both training and test set molecules. 3-Substituted imino benzothiazolines with anticonvulsant (antiglutamate) can be modeled excellently by the above equations using index of refraction, surface tension and average mass and indicator parameter. Indicator parameter is important parameter which gives best results, in this case presence of 6-trifluoromethoxy is influencing biological activity. The substitutions at 3-increasing the index of refraction may increase biological activity and presence of 6-trifluoromethoxy increases the activity.

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