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# PM3 Method based QSAR Study of the Derivatives of Thiadiazole and Quinoxaline for Antiepileptic Activity using Topological Descriptors 

Durgesh Kumar Mishra ${ }^{1}$, Ashutosh Singh ${ }^{2, 凶}$, Sunil Mishra ${ }^{1}$, Priti Singh ${ }^{2}$ and Abhishek Singh ${ }^{3}$

## ABSTRACT

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QSAR study of the derivatives of thiadiazole and quinoxaline has been performed for the antiepileptic activity using the topological descriptors viz., molar refractivity, shape index (basic kappa, order 1), shape index (basic kappa, order 2), shape index (basic kappa, order 3), valence connectivity index (order 0 , standard), valence connectivity index (order 1, standard) and valence connectivity index (order 2, standard). In the best QSAR model, the descriptors are molar refractivity, shape index (basic kappa, order 1), shape index (basic kappa, order 3) and valence connectivity index (order 0 , standard). In this QSAR model, the regression coefficient is 0.872435 and cross-validation coefficient is 0.832189 , which indicate that this QSAR model can be used to predict the antiepileptic activity of any compound belonging to this series. QSAR model developed using single descriptor shape index (basic kappa, order 1) or shape index (basic kappa, order 3) or valence connectivity index (order 2, standard) also has good predictive power.

## K EY W ORD S

## Author affiliations:

${ }^{1}$ Department of Chemistry, M.L.K. Post Graduate College, Balrampur-271201, India
${ }^{2}$ Department of Chemistry, K.S. Saket Post Graduate College, Ayodhya-224123, India
${ }^{3}$ Department of Chemistry, Udai Pratap Autonomous College, Varanasi221003, India
${ }^{\boxtimes}$ To whom correspondence to be addressed:
E-mail: asinghkssaket@gmail.com; dmchemistrydada@gmail.com

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QSAR models, Thiadiazole, Quinoxaline, Descriptors, Antiepileptic activity, Computational chemistry.

## INTRODUCTION

QSAR is a process in which the structures of a set of compounds are quantified and then compared to the numerical values of a biological activity or a physical property [1-5]. The challenge here has been to find some numerical code for a molecule or a fragment that is information-rich. This structure information and the measured property or activity is then processed into a mathematical model of relationship. From a good QSAR model, it is possible to predict and design the compounds for synthesis and testing that have a good possibility for activity [4-9].

Computational chemistry [10-16] represents molecular structures as a numerical model and simulates their behaviour with the equations of quantum and classical physics. Available programs enable scientists to easily generate and present molecular data including geometries, energies and associated properties. A QSAR attempts to find consistent relationships between the variations in the values of molecular properties
and the biological activity for a series of compounds.
A QSAR generally takes the form of a linear equation:

$$
\begin{gathered}
\text { Biological activity }=\text { Const }+\left(\mathrm{C}_{1} \mathrm{P}_{1}\right)+ \\
\left(\mathrm{C}_{2} \mathrm{P}_{2}\right)+\left(\mathrm{C}_{3} \mathrm{P}_{3}\right)+\ldots \ldots .+\left(\mathrm{C}_{\mathrm{n}} \mathrm{P}_{\mathrm{n}}\right)
\end{gathered}
$$

where the parameters $P_{1}$ through $P_{n}$ are computed for each molecule in the series and the coefficients $C_{1}$ through $C_{n}$ are calculated by fitting variations in the parameters and the biological activity [17-25].

## EXPERIMENTAL

Molar refractivity [26]: Molar refractivity was calculated by the Lorenz-Lorentz formula:

$$
\mathrm{MR}=\frac{\mathrm{n}^{2}-1}{\mathrm{n}^{2}+1} \times \frac{\mathrm{M}}{\rho}
$$

where $M$ is the molecular weight, $n$ is the refraction index and $\rho$ is the density. For a radiation of infinite wavelength, molar refractivity represents the real volume of the molecules.

Kier's shape indices $\left\{\mathbf{k}_{\mathbf{n}}(\mathbf{n}=\mathbf{1 , 2 , 3})\right\}$ [27-30,35]: These indices compare the molecule graph with "minimal" and "maximal" graphs, where the meaning of "minimal" and "maximal" depends on the order $n$. This is intended to capture different aspects of the molecular shape.

Order 1: The shape index of order 1 is then defined as:

$$
\kappa_{1}=2 \mathrm{P}_{\min } \frac{\mathrm{P}_{\max }}{\mathrm{P}^{2}}
$$

where P is the number of edges in the graph (edges are paths of length 1 , hence the subscript on the $\kappa_{1}$ ), $\mathrm{P}_{\text {max }}$ is the number of edges in the maximal graph namely $\mathrm{N}(\mathrm{N}-1) / 2$ and $\mathrm{P}_{\text {min }}$ is the number of edges in the minimal graph namely $(\mathrm{N}-1)$.

By inserting the formulas for $\mathrm{P}_{\text {max }}$ and $\mathrm{P}_{\text {min }}$, one obtains the implemented formula:

$$
\begin{equation*}
\kappa_{1}=\frac{\mathrm{N}(\mathrm{~N}-1)^{2}}{\mathrm{P}^{2}} \tag{1}
\end{equation*}
$$

Order 2: The descriptor $\mathrm{k}_{2}$ encodes the branching. $\mathrm{P}, \mathrm{P}_{\text {min }}$ and $\mathrm{P}_{\text {max }}$ now denote the number of paths of length 2 in the corresponding graphs. The maximal graph is taken to be the star graph in which all atoms are adjacent to a common atom. Thus, $\mathrm{P}_{\max }=(\mathrm{N}-1)(\mathrm{N}-2) / 2$. The linear graph is again taken as the minimal graph, so $\mathrm{P}_{\min }=\mathrm{N}-2$. eqn. 1 thus yields:

$$
\begin{equation*}
\kappa_{2}=\frac{(\mathrm{N}-1)(\mathrm{N}-2)^{2}}{\mathrm{P}^{2}} \tag{2}
\end{equation*}
$$

Order 3: For order 3, the counts of paths of length 3 are considered and the maximal graph chosen is a twin-star with $\mathrm{P}_{\max }=(\mathrm{N}-1)(\mathrm{N}-3) / 4$ for N odd and $\mathrm{P}_{\max }=(\mathrm{N}-2)^{2} / 4$ for N even. The minimal graph is again the linear one with $P_{\text {min }}=N$ -3 .

The equation is adjusted by another factor of 2 in the words of the index design "to bring the values into rough equivalence with the other kappa values".

$$
\begin{align*}
\kappa_{3} & =\frac{(\mathrm{N}-1)(\mathrm{N}-3)^{2}}{\mathrm{P}^{2}} \text { for } \mathrm{N} \text { odd } \\
\text { and } \quad \kappa_{3} & =\frac{(\mathrm{N}-2)(\mathrm{N}-3)^{2}}{\mathrm{P}^{2}} \text { for } \mathrm{N} \text { even }
\end{align*}
$$

Valence connectivity indices: Valence connectivity indices are originally defined by Randic [34] and subsequently refined by Kier \& Hall [35], is a series of numbers designated by "order" and "sub-graph type". There are four sub-graph types; path, cluster, path/cluster and chain. These types emphasize different aspects of atom connectivity within a molecule, the amount of branching, ring structures present and flexibility. It is calculated from the hydrogen suppressed molecular graph and defined as follows:

$$
{ }^{\mathrm{m}} \chi^{\mathrm{v}}=\sum_{\mathrm{i}=1}^{\mathrm{NS}} \prod_{\mathrm{k}=1}^{\mathrm{m}+1}\left(\frac{1}{\delta_{\mathrm{k}}^{\mathrm{v}}}\right)^{1 / 2}
$$

where, $\delta_{\mathrm{k}}^{v}=\frac{\left(\mathrm{Z}_{\mathrm{k}}^{v}-\mathrm{H}_{\mathrm{k}}\right)}{\left(\mathrm{Z}_{\mathrm{k}}-\mathrm{Z}_{\mathrm{k}}^{\mathrm{v}}-1\right)}$ - valence connectivity for the kth atom in the molecular graph, $\mathrm{Z}_{\mathrm{k}}=$ the total number of electrons in the kth atom, $\mathrm{Z}_{\mathrm{k}}^{\mathrm{v}}=$ the number of valence electrons in the kth atom, $\mathrm{H}_{\mathrm{k}}=$ the number of hydrogen atoms directly attached to the kth non-hydrogen atom, $\mathrm{m}=0$ - atomic valence connectivity indices (called order-0), $\mathrm{m}=1$ - one bond path valence connectivity indices (called order-1), $m=2$ - two bond fragment valence connectivity indices (called order-2).

Study material: Derivatives of thiadiazole and quinoxaline have been used as the study material. These derivatives are presented in Table-1 along with their antiepileptic activities $\left(\mathrm{pED}_{50}\right)$. For QSAR analysis, the 3D modeling and geometry optimization $[22,23]$ of all the compounds have been done with the help of CAChe software. The values of descriptors used for QSAR models have been evaluated using the CAChe software by PM3 [24,25] methods. The descriptors that have been used for the development of QSAR models are given below:

1. Molar refractivity
2. Shape index (basic kappa, order 1)
3. Shape index (basic kappa, order 2)
4. Shape index (basic kappa, order 3)
5. Valence Connectivity Index (order 0, standard)
6. Valence connectivity index (order 1 , standard)
7. Valence connectivity index (order 2, standard)

## RESULTS AND DISCUSSION

QSAR studies of the compounds listed in Table-1 were made with the help of topological descriptors in the different combinations. The outlier compounds were TD14, TD23 and TD37. The values of the descriptors have been calculated using PM3 method with the help of CAChe software and are given in Table-2. The values of the descriptors have been used to develop ninety QSAR models using different combinations of descriptors. With the help of good QSAR models, the activity of any unknown compound of the series can be calculated and the then synthesis may be done if the activity is found good. A QSAR model is said to have good predictive power if the regression coefficient $\left(r^{2}\right)$ is greater than 0.5 and the value of cross-validation coefficient $\left(\mathrm{rCV}^{2}\right)$ is greater than or equal to 0.2 . As the value of regression coefficient increases, the predictive power of the QSAR model increases. A QSAR model is said to have $100 \%$ predictive power if the value of

TABLE-1
ANTIEPILEPTIC ACTIVITIES $\left(\mathrm{pED}_{50}\right)$ OF THIADIAZOLES AND QUINOXALINE DERIVATIVES

## Compound

Compound structure
Antiepileptic activity $\left(\mathrm{pED}_{50}\right)$

TD 1


TD 2


TD 3


TD 5


TD 4


TD 6


TD 7


TD 8



TD 10


TD 11


TD 12


TD 13


TD 14


TD 15


TD 16


TD 17


TD 19


TD 18


TD 20




TD 21


TD 22


| TD 23 |  | 0.34 |
| :---: | :---: | :---: |
| TD 24 |  | 0.47 |
| TD 25 |  | 0.81 |
| TD 26 |  | 0.74 |
| TD 27 |  | 0.91 |
| TD 28 |  | 1.60 |
| TD 29 |  | 1.90 |



TD 31


TD 32


TD 33


TD 34


TD 35


TD 36



TABLE-2
VALUES OF TOPOLOGICAL DESCRIPTORS OF THIADIAZOLE AND QUINOXALINE DERIVATIVES ALONG WITH ANTIEPILEPTIC ACTIVITY

| Compound | Molar refractivity | Shape index (basic kappa) |  |  | Valence connectivity index (standard) |  |  | Activity |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
|  |  | Order 1 | Order 2 | Order 3 | Order 0 | Order 1 | Order 2 |  |
| TD 1 | 134.579 | 26.601 | 12.240 | 6.533 | 20.114 | 13.077 | 10.610 | 0.810 |
| TD 2 | 119.525 | 24.135 | 10.508 | 5.742 | 18.650 | 11.833 | 9.957 | 0.870 |
| TD 3 | 92.246 | 18.781 | 8.131 | 5.019 | 14.534 | 9.839 | 8.154 | 0.910 |
| TD 4 | 117.390 | 23.168 | 9.868 | 5.333 | 18.988 | 12.281 | 11.162 | 0.300 |
| TD 5 | 171.721 | 33.161 | 13.278 | 5.781 | 26.567 | 18.179 | 15.967 | 0.830 |
| TD 6 | 105.816 | 22.203 | 9.647 | 5.087 | 16.851 | 10.775 | 8.952 | 0.540 |
| TD 7 | 121.390 | 29.970 | 11.852 | 7.398 | 19.918 | 12.461 | 10.461 | 0.240 |
| TD 8 | 114.484 | 23.168 | 10.292 | 5.689 | 17.727 | 11.416 | 9.533 | 0.900 |
| TD 9 | 103.469 | 21.703 | 8.789 | 5.136 | 16.918 | 10.595 | 10.585 | 0.900 |
| TD 10 | 115.416 | 26.074 | 10.948 | 6.416 | 18.361 | 11.733 | 9.745 | 0.400 |
| TD 11 | 121.851 | 23.168 | 10.292 | 5.689 | 19.263 | 12.184 | 10.419 | 0.670 |
| TD 12 | 117.065 | 23.168 | 10.292 | 5.689 | 18.691 | 11.898 | 10.089 | 0.910 |
| TD 13 | 122.741 | 28.994 | 12.027 | 6.966 | 19.548 | 12.239 | 10.125 | 0.320 |
| TD 15 | 115.416 | 26.074 | 10.948 | 6.416 | 18.361 | 11.733 | 9.742 | 0.470 |
| TD 16 | 141.967 | 31.426 | 12.545 | 6.745 | 22.125 | 13.790 | 11.524 | 0.330 |
| TD 17 | 140.616 | 32.395 | 12.416 | 7.131 | 22.495 | 14.012 | 11.860 | 0.250 |
| TD 18 | 194.884 | 36.950 | 15.264 | 7.275 | 29.067 | 19.385 | 16.499 | 0.810 |
| TD 19 | 136.616 | 25.641 | 10.401 | 5.218 | 21.565 | 13.832 | 12.561 | 0.320 |
| TD 20 | 141.076 | 25.641 | 10.776 | 5.496 | 21.840 | 13.736 | 11.818 | 0.800 |
| TD 21 | 136.291 | 25.641 | 10.776 | 5.496 | 21.269 | 13.450 | 11.488 | 0.790 |
| TD 22 | 133.709 | 25.641 | 10.776 | 5.496 | 20.305 | 12.968 | 10.931 | 0.840 |
| TD 24 | 134.642 | 28.526 | 11.472 | 6.189 | 20.939 | 13.285 | 11.140 | 0.470 |
| TD 25 | 134.642 | 28.526 | 11.472 | 6.189 | 20.939 | 13.285 | 11.144 | 0.810 |
| TD 26 | 134.406 | 26.601 | 11.396 | 5.723 | 20.252 | 12.941 | 10.754 | 0.740 |
| TD 27 | 108.891 | 21.240 | 8.626 | 4.694 | 16.995 | 11.259 | 9.421 | 0.910 |
| TD 28 | 94.726 | 17.416 | 8.131 | 4.066 | 13.402 | 8.359 | 6.097 | 1.600 |
| TD 29 | 74.536 | 13.959 | 6.185 | 2.880 | 10.593 | 6.487 | 4.750 | 1.900 |
| TD 30 | 91.476 | 17.416 | 7.319 | 3.486 | 13.525 | 8.230 | 6.294 | 1.600 |
| TD 31 | 107.504 | 21.240 | 10.347 | 5.587 | 16.037 | 10.257 | 7.665 | 1.500 |
| TD 32 | 98.511 | 18.781 | 9.087 | 4.803 | 14.448 | 8.621 | 6.124 | 1.500 |
| TD 33 | 112.055 | 21.240 | 10.347 | 5.587 | 16.127 | 9.721 | 7.085 | 1.500 |
| TD 34 | 82.918 | 16.844 | 7.713 | 4.110 | 12.424 | 7.402 | 5.412 | 1.500 |
| TD 35 | 104.664 | 19.322 | 9.467 | 4.899 | 14.427 | 8.879 | 6.387 | 1.600 |
| TD 36 | 106.358 | 20.280 | 9.667 | 5.136 | 14.797 | 9.013 | 6.571 | 1.600 |

regression coefficient becomes unity. QSAR model has no predictive power if either the value of cross-validation coefficient $\left(\mathrm{rCV}^{2}\right)$ is less than 0.2 or the value regression coefficient $\left(\mathrm{r}^{2}\right)$ is less than 0.5 . Predicted antiepileptic activities PB1-PB8 of the compounds have been obtained by substituting the values of descriptors in the MLR equations and are given in Table-3.

Out of 90 QSAR models developed with the help of topological descriptors, the good QSAR models are 80. Five good QSAR models in decreasing order of predicted epileptic activity are listed in Table-4 along with the descriptors used. These good QSAR models are PB1, PB2, PB3, PB4 and PB5 in the decreasing order of their predictive power.

## Description of first five good QSAR models

Best QSAR model: The best QSAR model is PB1 in which the descriptors are molar refractivity, shape index (basic kappa, order 1), shape index (basic kappa, order 3) and valence connectivity index (order 0, standard). Multi linear regression (MLR) equation is given below:

PB1 $=0.0585933 *$ Molar refractivity- 0.0187282 * Shape index (basic kappa, order 1)-0.111044 * Shape index (basic kappa, order 3)-0.392556 * Valence connectivity index (order 0 , standard) +2.18438
$\mathrm{rCV}^{2}=0.832189$
$\mathrm{r}^{2}=0.872435$

TABLE-3
VALUES OF THE PREDICTED ANTIEPILEPTIC ACTIVITIES FROM PB1-PB8

| Compound | PB1 | PB2 | PB3 | PB4 | PB5 | PB6 | PB7 | PB8 |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| TD 1 | 0.950 | 0.938 | 0.940 | 0.929 | 0.931 | 0.740 | 0.787 | 0.561 |
| TD 2 | 0.777 | 0.767 | 0.765 | 0.765 | 0.765 | 0.907 | 0.870 | 0.836 |
| TD 3 | 0.975 | 0.929 | 0.980 | 0.955 | 0.957 | 1.271 | 1.099 | 1.087 |
| TD 4 | 0.583 | 0.568 | 0.566 | 0.562 | 0.563 | 0.973 | 0.717 | 0.978 |
| TD 5 | 0.554 | 0.615 | 0.595 | 0.578 | 0.579 | 0.294 | 0.106 | 0.822 |
| TD 6 | 0.789 | 0.824 | 0.798 | 0.796 | 0.795 | 1.038 | 0.998 | 1.063 |
| TD 7 | 0.095 | 0.108 | 0.112 | 0.114 | 0.113 | 0.511 | 0.806 | 0.260 |
| TD 8 | 0.868 | 0.857 | 0.861 | 0.854 | 0.855 | 0.973 | 0.924 | 0.854 |
| TD 9 | 0.629 | 0.611 | 0.620 | 0.620 | 0.623 | 1.072 | 0.790 | 1.046 |
| TD 10 | 0.538 | 0.548 | 0.549 | 0.544 | 0.544 | 0.776 | 0.897 | 0.602 |
| TD 11 | 0.697 | 0.650 | 0.656 | 0.659 | 0.659 | 0.973 | 0.811 | 0.854 |
| TD 12 | 0.641 | 0.618 | 0.613 | 0.613 | 0.613 | 0.973 | 0.853 | 0.854 |
| TD 13 | 0.386 | 0.425 | 0.405 | 0.407 | 0.407 | 0.577 | 0.849 | 0.410 |
| TD 15 | 0.538 | 0.548 | 0.549 | 0.544 | 0.544 | 0.776 | 0.897 | 0.602 |
| TD 16 | 0.480 | 0.520 | 0.500 | 0.512 | 0.511 | 0.412 | 0.671 | 0.487 |
| TD 17 | 0.194 | 0.215 | 0.214 | 0.226 | 0.224 | 0.346 | 0.628 | 0.353 |
| TD 18 | 0.693 | 0.681 | 0.699 | 0.691 | 0.692 | 0.037 | 0.038 | 0.303 |
| TD 19 | 0.664 | 0.642 | 0.646 | 0.652 | 0.652 | 0.805 | 0.539 | 1.018 |
| TD 20 | 0.787 | 0.737 | 0.747 | 0.761 | 0.759 | 0.805 | 0.633 | 0.921 |
| TD 21 | 0.730 | 0.705 | 0.704 | 0.714 | 0.713 | 0.805 | 0.675 | 0.921 |
| TD 22 | 0.957 | 0.944 | 0.952 | 0.955 | 0.955 | 0.805 | 0.746 | 0.921 |
| TD 24 | 0.632 | 0.644 | 0.645 | 0.649 | 0.649 | 0.609 | 0.720 | 0.680 |
| TD 25 | 0.632 | 0.644 | 0.645 | 0.649 | 0.649 | 0.609 | 0.719 | 0.680 |
| TD 26 | 0.976 | 0.992 | 0.982 | 0.984 | 0.984 | 0.740 | 0.769 | 0.842 |
| TD 27 | 0.974 | 0.952 | 0.985 | 0.973 | 0.974 | 1.104 | 0.938 | 1.200 |
| TD 28 | 1.696 | 1.702 | 1.706 | 1.704 | 1.704 | 1.364 | 1.361 | 1.418 |
| TD 29 | 1.812 | 1.853 | 1.846 | 1.846 | 1.846 | 1.599 | 1.532 | 1.830 |
| TD 30 | 1.522 | 1.552 | 1.543 | 1.551 | 1.550 | 1.364 | 1.336 | 1.619 |
| TD 31 | 1.170 | 1.170 | 1.160 | 1.152 | 1.152 | 1.104 | 1.162 | 0.889 |
| TD 32 | 1.400 | 1.399 | 1.382 | 1.392 | 1.390 | 1.271 | 1.358 | 1.162 |
| TD 33 | 1.401 | 1.378 | 1.374 | 1.382 | 1.381 | 1.104 | 1.235 | 0.889 |
| TD 34 | 1.394 | 1.420 | 1.402 | 1.408 | 1.407 | 1.403 | 1.448 | 1.403 |
| TD 35 | 1.748 | 1.738 | 1.744 | 1.743 | 1.744 | 1.234 | 1.324 | 1.128 |
| TD 36 | 1.657 | 1.646 | 1.655 | 1.656 | 1.657 | 1.169 | 1.301 | 1.046 |

TABLE-4

| TABLE-4 <br> GOOD QSAR MODELS IN THE DECREASING ORDER OF PREDICTIVE EPILEPTIC ACTIVITIES ALONG WITH THE DESCRIPTORS USED |  |  |  |
| :---: | :---: | :---: | :---: |
| Predicted activity | Descriptors used in the predicted activity | Cross-validation coefficient ( $\mathrm{rCV}^{\wedge} 2$ ) | Regression coefficient ( $\mathrm{r}^{\wedge} 2$ ) |
| PB1 | Molar refractivity, shape index (basic kappa, order 1), shape index (basic kappa, order 3 ), valence connectivity index (order 0 , standard) | 0.832189 | 0.872435 |
| PB2 | Molar refractivity, shape index (basic kappa, order 2), shape index (basic kappa, order 3 ), valence connectivity index (order 0 , standard) | 0.787784 | 0.871703 |
| PB3 | Molar refractivity, shape index (basic kappa, order 3), valence connectivity index (order 0 , standard), valence connectivity index (order 1 , standard) | 0.813375 | 0.871128 |
| PB4 | Molar refractivity, shape index (basic kappa, order 3), valence connectivity index (order 0 , standard), valence connectivity index (order 2, standard) | 0.800679 | 0.870877 |
| PB5 | Molar refractivity, shape index (basic kappa, order 3), valence connectivity index (order 0 , standard) | 0.829812 | 0.870871 |

The value of regression coefficient is 0.872435 and the value of cross-validation coefficient is 0.872435 . These values of regression and cross-validation coefficients indicate that the QSAR model possesses very good predictive power and can be used to predict the antiepileptic activity of any compound of this series. Graph between observed and predicted antiepileptic activities is shown in Fig. 1. The difference between observed antiepileptic activity and predicted antiepileptic activity PB1 is shown in Fig. 2.

Second best QSAR model: The second best QSAR model is PB2 whose MLR equation is given by:

PB2 $=0.0532403 *$ Molar refractivity $+0.060905 *$ Shape index (basic kappa, order 2)-0.208831 * Shape index (basic kappa, order 3)-0.390519 * Valence connectivity index (order 0 , standard) +2.24621
$\mathrm{rCV}^{2}=0.787784$ $r^{2}=0.871703$
These values of regression and cross-validation coefficients indicate that this QSAR model possesses very good predictive power and can be used to predict the antiepileptic activity of any compound of this series. Graph between observed and predicted antiepileptic activities is shown in Fig. 3. The


Fig. 1. Graph between observed activity and predicted activity PB1


Fig. 2. Graph showing the difference between observed activity and predicted activity PB1


Fig. 3. Graph between observed activity and predicted activity PB2
difference between observed antiepileptic activity and predicted antiepileptic activity PB2 is shown in Fig. 4.


Fig. 4. Graph showing the difference between observed activity and predicted activity PB2

Third best QSAR model: The third best QSAR model is PB3 whose MLR equation is given by:

PB3 $=0.0585114 *$ Molar refractivity- $0.142787 *$ Shape index (basic kappa, order 3)-0.42761 * Valence Connectivity index (order 0, standard) +0.0262424 * Valence connectivity index (order 1, standard) + 2.25573

$$
\begin{aligned}
& \mathrm{rCV}^{2}=0.813375 \\
& \mathrm{r}^{2}=0.871128
\end{aligned}
$$

These values of regression and cross-validation coefficients indicate that this QSAR model possesses very good predictive power and can be used to predict the antiepileptic activity of any compound of this series. Graph between observed and predicted antiepileptic activities is shown in Fig. 5. The difference between observed antiepileptic activity and predicted antiepileptic activity PB3 is shown in Fig. 6.


Fig. 5. Graph between observed activity and predicted activity PB3


Fig. 6. Graph showing the difference between observed activity and predicted activity PB3

Fourth best QSAR model: The fourth best QSAR model is PB4 whose MLR equation is given by:

PB4 $=0.0581155 *$ Molar refractivity-0.14874 * Shape index (basic kappa, order 3)-0.403869 * Valence connectivity index (order 0 , standard)0.00291618 * Valence connectivity index (order 2, standard) +2.2343
$\mathrm{rCV}^{2}=0.800679$
$\mathrm{r}^{2}=0.870877$
These values of regression and cross-validation coefficients indicate that this QSAR model possesses very good predictive power and can be used to predict the antiepileptic activity of any compound of this series. Graph between observed and predicted antiepileptic activities is shown Fig. 7. The difference between observed antiepileptic activity and predicted antiepileptic activity PB4 is shown in Fig. 8.

Fifth best QSAR model: The Fifth best QSAR model is PB5 whose MLR equation is given by:

PB5 $=0.0583898 *$ Molar refractivity-0.147308 *
Shape index (basic kappa, order 3)-0.407782 *
Valence connectivity index (order 0 , standard) + 2.2373
$\mathrm{rCV}^{2}=0.829812$
$\mathrm{r}^{2}=0.870871$
These values of regression and cross-validation coefficients indicate that this QSAR model possesses very good


Fig. 7. Graph between observed activity and predicted activity PB4


Fig. 8. Graph showing the difference between observed activity and predicted activity PB4
predictive power and can be used to predict the antiepileptic activity of any compound of this series.

Good QSAR model using single descriptors: Descriptors shape index (basic kappa, order 1), valence connectivity index (order 2, standard) and shape index (basic kappa, order 3) are individually capable to produce good QSAR models. These QSAR models are discussed below:

QSAR model using single descriptor shape index (basic kappa, order 1): The predicted activity PB6 of the QSAR model developed using the descriptor Shape Index (basic kappa, order 1) is as followsPB6 $=-0.0679319 *$ Shape Index (basic kappa, order 1) +2.54677 $\mathrm{rCV}^{2}=0.515348$ $\mathrm{r}^{2}=0.536487$
The values of regression and cross-validation coefficients suggest that the QSAR model possesses good predictive power.

QSAR model using single descriptor valence connectivity index (order 2, standard): The predicted activity PB7 of the QSAR model developed using the descriptor Valence Connectivity Index (order 2, standard) is as follows-

PB7=-0.127179 * Valence connectivity index (order 2 , standard) +2.13646 $\mathrm{rCV}^{2}=0.476899$ $r^{2}=0.509596$
The values of regression and cross-validation coefficients suggest that the QSAR model possesses good predictive power.

QSAR model using single descriptor shape index (basic kappa, order 2): The predicted activity PB8 of the QSAR model developed using the descriptor shape index (basic kappa, order 3 is as follows:

PB8 $=-0.347358 *$ Shape index (basic kappa, order 3) +2.83016
$\mathrm{rCV}^{2}=0.522455$
$\mathrm{r}^{2}=0.549397$

The values of regression and cross-validation coefficients suggest that the QSAR model possesses good predictive power.

## Conclusion

Best QSAR model is PB1 in which the descriptors are molar refractivity, shape index (basic kappa, order 1), shape index (basic kappa, order 3) and valence connectivity index (order 0, standard). In this QSAR model, the regression coefficient is 0.872435 and cross-validation coefficient is 0.832189 . These values of regression and cross-validation coefficients indicate that the predictive power of the QSAR model PB1 is excellent. Shape index (basic kappa, order 1) alone produces QSAR model having good predictive power in which regression and cross-validation coefficients are 0.515348 and 0.536487 respectively. The single descriptor shape index (basic kappa, order 3) and the single descriptor valence connectivity index (order 2, standard) also produce the QSAR models having good predictive power.

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