

## Structural feature indenture for estrogen analogs as anticancer agents: *De novo* and Hansch Approach

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Quantitative structure activity relationships have been performed on a series of twenty-six compounds of estrogen derivatives, for their anti-proliferative activity, in order to understand the essential structural requirement for inhibition of proliferation in estrogen dependent MCF-7 human breast cancer cells. The quantitative models derived for the study illustrates the significance of the  $\beta$ -hydroxy group at 17<sup>th</sup> position for the drug-enzyme interaction and allowed hydrogen-bonding interaction to estrogenic receptor in optimal manner (free rotation) as compared to oxo group (restricted orientation). The results of the study also reveal the necessity of sulfonamide moiety at 3<sup>rd</sup> position of estrogen. Additionally, presence of smaller substituents at 2<sup>nd</sup> position will be conducive for the activity.

**Keywords:** Estrogen analogs, Hansch approach, Fujita-Ban analysis, QSAR, antiproliferative activity, anticancer agents

Cancer is the second leading cause of mortality in developed countries, and the discovery and development of new treatment is urgently needed due to problem with currently available treatment such as toxicities and drug resistance<sup>1</sup>. Many of current clinically used anticancer agents including paclitaxel, vinblastin<sup>2-4</sup> and compound under clinical evaluation, such as epothiolone<sup>5</sup> are microtubular interfering agents that bind to tubulin. Several new anticancer agents are developed recently such as Gleenec<sup>6</sup> which is kinase inhibitor designed as more selective with less toxicity. Similarly, discovery of the endogenous estrogen metabolite 2-methoxy estradiol (2-MeOE2) would inhibit cancer cell proliferation and angiogenesis<sup>7</sup> and it is 250 times more potent than estradiol. A number of potential mechanisms underlying the reversible mitotic arrest<sup>8</sup> and apoptosis<sup>9-10</sup> induced by 2-methoxy estradiol in various cancer cell lines have been proposed. 2-Substituted estrone sulfamates (EMATEs) displays the similar antiproliferative activity profile to the corresponding estradiol against a range of human cancer cell lines and have 80 fold greater activity than the 2-MeOE2. 2-substituted EMATEs also inhibit the steroid sulfatase, a therapeutic target for the treatment of hormone dependent breast cancer<sup>11</sup>. The 2-substituted EMATEs remain to be elucidated their spectrum of biological activity encouraged us to

explore the closely related structures in the search for the new anticancer compounds.

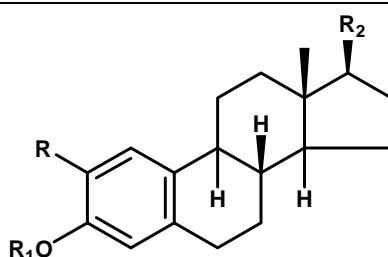
These arouse our interest for designing of novel hormone dependent breast anti-cancer agents. In the present work, we have attempted to quantify the necessary structural and physicochemical requirements for antiproliferative activity against MCF-7 human breast cancer cells.

### Materials and methods

The antiproliferative activity data of estrogen analogs were taken from the reported work of Leese *et. al.*<sup>12</sup> (**Table I**). The antiproliferative activity data against MCF-7 human breast cancer cells ( $GI_{50}$  in  $\mu\text{m}$ ) was converted to negative logarithmic mole dose ( $pGI_{50}$ ) in order to reduce the skewness of the data set, for quantitative structure activity relationship analysis. Initially series was subjected to Fujita-Ban analysis using multiple regression technique in order to estimate the *de novo* contribution of substituents to the activity of the molecules. Further Hansch approach was carried out to establish correlations between antiproliferative activity and various substituents constants at position R,  $R_1$  and indicator variable  $Iv_2$  for the presence of hydroxy group at position  $R_2$  of the molecule (**Figure 1**). The obtained model shows presence of outlier compound on the

**Table I**—Analogs of estrogen and their antiproliferative activity data

Compd	R	R <sub>1</sub>	R <sub>2</sub>	R <sub>3</sub>	GI <sub>50</sub> <sup>a</sup>	pGI <sub>50</sub> <sup>b</sup>
<b>1</b>	CH <sub>3</sub> O	H	O	O	21.3	4.672
<b>2</b>	CH <sub>3</sub> O	H	OH	H	2.35	5.629
<b>3</b>	CH <sub>3</sub> O	SO <sub>2</sub> NH <sub>2</sub>	O	O	0.30	6.523
<b>4</b>	CH <sub>3</sub> O	SO <sub>2</sub> NH <sub>2</sub>	OH	H	0.36	6.444
<b>5</b>	CH <sub>3</sub> O	SO <sub>2</sub> N(CH <sub>3</sub> ) <sub>2</sub>	O	O	>10	<5.000
<b>6</b>	CH <sub>3</sub> O	SO <sub>2</sub> NHCOCH <sub>3</sub>	O	O	168	3.775
<b>7</b>	C <sub>2</sub> H <sub>5</sub> O	H	O	O	9.04	5.044
<b>8</b>	C <sub>2</sub> H <sub>5</sub> O	SO <sub>2</sub> NH <sub>2</sub>	O	O	4.84	5.315
<b>9</b>	C <sub>2</sub> H <sub>5</sub> O	SO <sub>2</sub> NH <sub>2</sub>	OH	H	0.61	6.215
<b>10</b>	CH <sub>3</sub> S	H	O	O	33.4	4.476
<b>11</b>	CH <sub>3</sub> S	H	OH	H	3.96	5.402
<b>12</b>	CH <sub>3</sub> S	SO <sub>2</sub> NH <sub>2</sub>	O	O	0.40	6.398
<b>13</b>	CH <sub>3</sub> S	SO <sub>2</sub> NH <sub>2</sub>	OH	H	0.43	6.367
<b>14</b>	CH <sub>3</sub> SO	H	O	O	>10	<5.000
<b>15</b>	CH <sub>3</sub> SO <sub>2</sub>	H	O	O	66	4.180
<b>16</b>	C <sub>2</sub> H <sub>5</sub> S	H	O	O	31.9	4.496
<b>17</b>	C <sub>2</sub> H <sub>5</sub> S	H	OH	H	23.2	4.635
<b>18</b>	C <sub>2</sub> H <sub>5</sub> S	SO <sub>2</sub> NH <sub>2</sub>	O	O	35.2	4.453
<b>19</b>	C <sub>2</sub> H <sub>5</sub> S	SO <sub>2</sub> NH <sub>2</sub>	OH	H	23.4	4.631
<b>20</b>	C <sub>2</sub> H <sub>5</sub>	H	O	O	57.6	4.240
<b>21</b>	C <sub>2</sub> H <sub>5</sub>	H	OH	H	10.5	4.979
<b>22</b>	C <sub>2</sub> H <sub>5</sub>	SO <sub>2</sub> NH <sub>2</sub>	O	O	0.34	6.469
<b>23</b>	C <sub>2</sub> H <sub>5</sub>	SO <sub>2</sub> NH <sub>2</sub>	OH	H	0.07	7.155
<b>24</b>	CH <sub>3</sub>	SO <sub>2</sub> NH <sub>2</sub>	O	O	0.26	6.585
<b>25</b>	Br	H	O	O	33	4.481
<b>26</b>	Br	SO <sub>2</sub> NH <sub>2</sub>	O	O	6.95	5.158
<b>27</b>	I	H	O	O	38	4.420
<b>28</b>	I	SO <sub>2</sub> NH <sub>2</sub>	O	O	1.59	5.799

<sup>a</sup> Concentration of 50 percent antiproliferative activity data against MCF-7 human breast cancer cells in mole.<sup>b</sup> Negative logarithm of GI<sub>50</sub>**Figure 1**—Lead structure of estrogen analogs used in Fujita-Ban analysis

basis of Z-score value. For better understanding of drug receptor interaction removing outlier compound and incorporated indicator variable  $I_{V1}$  for the presence of sulfonamide moiety at position  $R_1$ . Values of the substituents constants like hydrophobic ( $\pi$ ), steric (molar refractivity or  $MR$ ), hydrogen acceptor ( $HA$ ), hydrogen donor ( $HD$ ) and electronic (field effect or  $\mathcal{F}$ , resonance effect or  $\mathcal{R}$  and Hammett's constant or  $\sigma$ ), taken from the reported data<sup>13,14</sup> and Verloop parameters (value of shape of each substituent) like  $L$ ,  $B_1$ ,  $B_2$ ,  $B_3$ ,  $B_4$  were taken from reported work of Skagerberg *et al*<sup>15</sup>. The data was transferred to statistical program VALSTAT<sup>16</sup> in order to establish the correlation between substituent constants as independent variable and antiproliferative activity as dependent variable using sequential multiple linear regression analysis method (in sequential multiple regression the program searches all permutation and combination sequentially for the

data set). The  $\pm$  data within the parentheses are the standard deviation, associated with coefficient of descriptors in regression equations. The best model was selected from the various statistically significant equations on the basis of observed squared correlation coefficient ( $r^2$ ), standard error of estimation (SE), sequential Fischer test (F), bootstrapping squared correlation coefficient ( $r_{bs}^2$ ), bootstrapping standard deviation ( $S_{bs}$ ), cross validated squared correlation coefficient using leave one out procedure ( $q^2$ ), chance statistics (evaluated as the ratio of the equivalent regression equations to the total number of randomized sets; a chance value of 0.001 corresponds to 0.1% chance of fortuitous correlation) and outliers (on the basis of Z-score value).

## Results and Discussion

Fujita-Ban analysis was carried out in order to find out *de novo* contribution of the substituents to the activity of the molecules (**Table II**). The multivariant

**Table II**—Fujita-Ban matrix, calculated  $pGI_{50}$ , residual and Z-score data of estrogens analogs

Compd	R <sub>1</sub>						R <sub>2</sub>			R <sub>3</sub>			Fujita-Ban Approach (eqn 2)		
	2C <sub>2</sub> H <sub>5</sub> O	2CH <sub>3</sub> S	2CH <sub>3</sub> SO <sub>2</sub>	2C <sub>2</sub> H <sub>5</sub> S	2C <sub>2</sub> H <sub>5</sub>	2CH <sub>3</sub>	2Br	2I	3SO <sub>2</sub> NH <sub>2</sub>	3SO <sub>2</sub> NHAc	17O	Cal. <sup>a</sup>	Cal <sub>res</sub> <sup>b</sup>	Z <sup>c</sup>	
<b>1</b>	0	0	0	0	0	0	0	0	0	0	1	4.610	0.062	0.124	
<b>2</b>	0	0	0	0	0	0	0	0	0	0	0	5.237	0.392	0.779	
<b>3</b>	0	0	0	0	0	0	0	0	1	0	1	5.875	0.648	1.288	
<b>4</b>	0	0	0	0	0	0	0	0	1	0	0	6.503	-0.059	-0.117	
<b>6</b>	0	0	0	0	0	0	0	0	0	1	1	4.610	-0.835	-1.659	
<b>7</b>	1	0	0	0	0	0	0	0	0	0	0	1	4.610	0.434	0.864
<b>8</b>	1	0	0	0	0	0	0	0	1	0	1	5.875	-0.560	-1.114	
<b>9</b>	1	0	0	0	0	0	0	0	1	0	0	6.503	-0.288	-0.572	
<b>10</b>	0	1	0	0	0	0	0	0	0	0	0	1	4.610	-0.134	-0.266
<b>11</b>	0	1	0	0	0	0	0	0	0	0	0	0	5.237	0.165	0.328
<b>12</b>	0	1	0	0	0	0	0	0	1	0	1	5.875	0.523	1.039	
<b>13</b>	0	1	0	0	0	0	0	0	1	0	0	6.503	-0.136	-0.270	
<b>15</b>	0	0	1	0	0	0	0	0	0	0	1	4.610	-0.430	-0.854	
<b>16</b>	0	0	0	1	0	0	0	0	0	0	1	3.607	0.889	1.767	
<b>17</b>	0	0	0	1	0	0	0	0	0	0	0	4.235	0.400	0.796	
<b>18</b>	0	0	0	1	0	0	0	0	1	0	1	4.873	-0.420	-0.835	
<b>19</b>	0	0	0	1	0	0	0	0	1	0	0	5.500	-0.869	-1.729	
<b>20</b>	0	0	0	0	1	0	0	0	0	0	1	4.610	-0.370	-0.735	
<b>21</b>	0	0	0	0	1	0	0	0	0	0	0	5.237	-0.258	-0.513	
<b>22</b>	0	0	0	0	1	0	0	0	1	0	1	5.875	0.594	1.180	
<b>23</b>	0	0	0	0	1	0	0	0	1	0	0	6.503	0.652	1.297	
<b>24</b>	0	0	0	0	0	1	0	0	1	0	1	5.875	0.710	1.411	
<b>25</b>	0	0	0	0	0	0	1	0	0	0	1	4.610	-0.129	-0.256	
<b>26</b>	0	0	0	0	0	0	1	0	1	0	1	5.875	-0.717	-1.426	
<b>27</b>	0	0	0	0	0	0	0	1	0	0	1	4.610	-0.190	-0.377	
<b>28</b>	0	0	0	0	0	0	0	1	1	0	1	5.875	-0.076	-0.152	

<sup>a</sup> calculated negative logarithm of concentration of 50 percent antiproliferative activity data against MCF-7 human breast cancer cells in mole. <sup>b</sup> calculated residual value. <sup>c</sup> Z-score value

regression expression (Eqn 1) indicated that the many substituent having poor contribution to the antiproliferative activity, which is further supported by high standard error of the substituent coefficient.

$$\begin{aligned}
 pGI_{50} = & 5.482(\pm 0.309) - 0.404(\pm 0.400)2C_2H_5O \\
 & 0.156(\pm 0.367)2CH_3S - 0.857(\pm 0.601) \\
 & 2CH_3SO_2 - 1.263(\pm 0.367)2C_2H_5S - \\
 & 0.106(\pm 0.367)2C_2H_5 + 0.433(\pm 0.604)2CH_3 - \\
 & 0.775(\pm 0.465)2Br - 0.485(\pm 0.465)2I \\
 & + 1.114(\pm 0.219)3OSO_2NH_2 - 1.262(\pm 0.601) \\
 & 3OSO_2NHCOCH_3 - 0.445(\pm 0.241)17O
 \end{aligned}$$

n=26, r=0.912, r<sup>2</sup>=0.832, SE=0.519, F=6.306 ... (1)

The Eqn 1 was further optimized by removing the substituents which are insignificant at 95% confidence level; a statistical significant trivalent expression (Eqn 2) was obtained.

$$\begin{aligned}
 pGI_{50} = & 5.237(\pm 0.224) + 1.266(\pm 0.211)3SO_2NH_2 - \\
 & 1.002(\pm 0.294)2C_2H_5S - 0.627(\pm 0.224)17O
 \end{aligned}$$

n=26, r=0.848, r<sup>2</sup>=0.719, SE=0.536, F=18.731 ... (2)

Fujita-Ban analysis of antiproliferative activity of estrogen analogs inferred that the presence of *o*-sulfamate moiety at 3<sup>rd</sup> position of the A ring of estrogen is essential for the activity as compared to non substituted estrogen or *N*-acetylated *o*-sulfamate analogs. *De novo* techniques also suggested that the presence of  $\beta$ -hydroxy group at 17<sup>th</sup> position is favorable as compared to oxo group.

Fujita-Ban expression gave insight to some important structural features i.e. the  $\beta$ -hydroxy group of 17<sup>th</sup> position is optimal for the activity and allowed hydrogen-bonding interaction to estrogenic receptor in optimal manner (free rotation) as compared to oxo group (restricted orientation).

Series was further subjected to Hansch approach, using stepwise multiple linear regression method, in order to develop 2D-QSAR between inhibition of proliferation of MCF-7 cells an estrogen dependent human breast cancer cell line as dependent variable and substituents constants as independent variables. A tri-varient statistically significant equation was obtained (Eqn 3).

$$\begin{aligned}
 pGI_{50} = & -0.766(\pm 0.404) MR - 0.481(\pm 0.157)\pi_1 + \\
 & 0.693(\pm 0.306) Iv_2 + 5.502
 \end{aligned}$$

n=26, r=0.684, r<sup>2</sup>=0.468, SE=0.737, F=6.442, outlier=1 ... (3)

Equation 3 showed moderate correlation coefficient value 0.684 with one outlier (Compd No 6). The unexplained variance in calculated activity of

compd no. 6 might be due to the steric hindrance of *N*-substitution on sulfamate analogs. The fitness of the model can be improved by removing outlier. Hence remaining 25 compounds were considered for the QSAR analysis. The 1:1 correlation study revealed that indicator variable (*Iv*<sub>1</sub>) contributed positively and linearly to the activity (Eqn 4) with 48.0% variance and statistical significance level better than 99.9% as it exceeded the student t-value 4.606 against tabulated t<sub>0.001(2), 23</sub> = 3.768, while molar refractivity contributed negatively to the activity (Eqn 5) with correlation coefficient value 0.437 and internal statistical significance level better than 95.0% (t<sub>0.05(2), 23</sub>=2.328 as it exceeded the tabulated value 2.069) and suggested that bulkier substitution at the parent structure is unfavorable for the inhibitory action.

$$pGI_{50} = 1.241(\pm 0.269) Iv_1 + 4.721$$

$$n=25, r=0.693, r^2=0.480, SE=0.673, F=21.212$$

$$pGI_{50} = -1.080(\pm 0.464) MR + 6.679 \quad \dots (4)$$

$$n=25, r=0.437, r^2=0.191, SE=0.840, F=5.420 \quad \dots (5)$$

Stepwise multiple linear regression analysis method was employed for improving the fitness of expression, several multivariant significant equations were obtained, and equation 6 was considered as model on the basis of statistical parameters (**Table III**).

$$\begin{aligned}
 pGI_{50} = & -1.014(\pm 0.292) MR + 1.152(\pm 0.211) Iv_1 + \\
 & 0.553(\pm 0.220) Iv_2 + 5.801
 \end{aligned}$$

$$\begin{aligned}
 n = 25, r = 0.843, r^2 = 0.711, SE = 0.525, F = 17.245, \\
 r_{bs}^2 = 0.729, S_{bs} = 0.089, \text{Chance} < 0.001, q^2 = 0.577, \\
 S_{PRESS} = 0.635, S_{DEP} = 0.582 \quad \dots (6)
 \end{aligned}$$

Equation 6 has better correlation coefficient (r=0.843), which accounts for more than 71.0% of the variance in the activity; also the inter-correlation among the parameters is less <0.100 (**Table IV**). The P value is less than 0.001 for each physiochemical parameters involved in the multivariant model generation, suggested that each independent variable contributed linearly. The data showed overall internal statistical significance level better than 99.9% as it exceeded the tabulated F<sub>(3,21 α 0.001)</sub> = 8.99. The model was further subjected for outlier by Z-score method and no compound was found to be an outlier, which suggested that the model is able to explain the structurally diverse analogs, and is helpful in designing of more potent compounds using physiochemical

**Table III**—Substituent constants, calculated, calculated (loo) value with residual and Z-score data of estrogen analogs used in QSAR study

Compd	Parameters			QSAR Model				
	MR	$Iv_1$	$Iv_2$	Cal. <sup>a</sup>	Cal <sub>res</sub> <sup>b</sup>	Z-Score	Cal <sub>(loo)</sub> <sup>c</sup>	Cal <sub>(loo)res</sub> <sup>d</sup>
<b>1</b>	0.787	0	0	5.003	-0.331	-0.675	5.067	-0.395
<b>2</b>	0.787	0	1	5.556	0.073	0.149	5.533	0.096
<b>3</b>	0.787	1	0	6.155	0.368	0.750	6.092	0.431
<b>4</b>	0.787	1	1	6.707	-0.264	-0.537	6.775	-0.331
<b>7</b>	1.247	0	0	4.537	0.507	1.033	4.478	0.565
<b>8</b>	1.247	1	0	5.688	-0.373	-0.760	5.732	-0.417
<b>9</b>	1.247	1	1	6.241	-0.026	-0.054	6.245	-0.031
<b>10</b>	1.382	0	0	4.400	0.077	0.156	4.390	0.086
<b>11</b>	1.382	0	1	4.952	0.450	0.916	4.865	0.537
<b>12</b>	1.382	1	0	5.551	0.847	1.724	5.438	0.960
<b>13</b>	1.382	1	1	6.104	0.262	0.534	6.058	0.309
<b>15</b>	1.349	0	0	4.433	-0.253	-0.515	4.464	-0.283
<b>16</b>	1.842	0	0	3.933	0.563	1.146	3.773	0.723
<b>17</b>	1.842	0	1	4.486	0.148	0.302	4.436	0.199
<b>18</b>	1.842	1	0	5.085	-0.632	-1.286	5.293	-0.839
<b>19</b>	1.842	1	1	5.638	-1.007	-2.051	5.990	-1.359
<b>20</b>	1.030	0	0	4.757	-0.517	-1.053	4.823	-0.584
<b>21</b>	1.030	0	1	5.309	-0.331	-0.673	5.383	-0.404
<b>22</b>	1.030	1	0	5.908	0.560	1.141	5.841	0.628
<b>23</b>	1.030	1	1	6.461	0.694	1.413	6.333	0.821
<b>24</b>	0.565	1	0	6.380	0.205	0.418	6.324	0.261
<b>25</b>	0.888	0	0	4.901	-0.419	-0.854	4.967	-0.486
<b>26</b>	0.888	1	0	6.052	-0.894	-1.821	6.181	-1.022
<b>27</b>	1.394	0	0	4.388	0.033	0.067	4.383	0.037
<b>28</b>	1.394	1	0	5.539	0.259	0.528	5.504	0.295

<sup>a</sup> calculated negative logarithm of concentration of 50 percent antiproliferative activity data against MCF-7 human breast cancer cells in mole.

<sup>b</sup> calculated residual value.

<sup>c</sup> calculated (leave-one-out) negative logarithm of concentration of 50 percent antiproliferative activity data against MCF-7 human breast cancer cells in mole.

<sup>d</sup> calculated (leave-one-out) residual value.

**Table IV**—Inter-correlation matrix of substituent constants used in QSAR analysis

	$\pi$	MR	$L$	$B_1$	$B_2$	$B_3$	$B_4$	HA	$\mathcal{R}$	$\mathcal{F}$	$Iv_1$	$Iv_2$
$\pi$	1.000											
MR	0.307	1.000										
$L$	0.176	0.866	1.000									
$B_1$	0.124	0.377	0.007	1.000								
$B_2$	0.017	0.718	0.839	0.309	1.000							
$B_3$	0.281	0.046	0.300	0.693	0.260	1.000						
$B_4$	0.590	0.110	0.045	0.634	0.188	0.719	1.000					
HA	0.759	0.352	0.039	0.521	0.067	0.266	0.177	1.000				
$\mathcal{R}$	0.082	0.313	0.042	0.674	0.027	0.647	0.538	0.628	1.000			
$\mathcal{F}$	0.395	0.192	0.151	0.575	0.183	0.353	0.569	0.276	0.042	1.000		
$Iv_1$	0.133	0.083	0.059	0.138	0.052	0.205	0.205	0.027	0.128	0.145	1.000	
$Iv_2$	0.060	0.090	0.157	0.280	0.309	0.147	0.255	0.021	0.135	0.233	0.053	1.000

parameters. The leave-one-out cross validation method was employed for the prediction of activity, and a  $q^2$  value (in the biological activity data of leave one compound) of 0.3 corresponds to a confidence limit greater than 95%, which minimizes the risk of finding significant explanatory equation for the biological activity just by mere chance. The Cross-validated squared correlation co-efficient ( $q^2=0.577$ ) suggested a good internal consistency as well as predictive ability of the biological activity. The  $r^2_{bs}$  is at par with the conventional squared correlation coefficient ( $r^2$ ). Randomized biological activity test (Chance < 0.001) revealed that the results were not based on chance correlation. In general, the model fulfills the statistical validation criteria to a significant extent to be a useful theoretical base for proposing more active compounds. The Hansch analysis indicated that both indicator variables  $Iv_1$  &  $Iv_2$  contributed positively while molar refractivity contributed negatively to the expression. Analysis revealed that the presence of sulfonamide moiety at 3<sup>rd</sup> position ( $Iv_1$ ) and  $\beta$ -hydroxy group ( $Iv_2$ ) at 17<sup>th</sup> position is essential for the optimum activity. Molar refractivity (MR) which is representative of bulkiness/molar volume of the substituents play key role at 2<sup>nd</sup> position of the ring and suggested that small substituents are more favorable as compared to the larger one.

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