## Electronic supplementary information (ESI)

# Design, synthesis, and biological evaluation of new <br> 6, $N^{2}$-diaryl-1,3,5-triazine-2,4-diamines as anticancer agents selectively targeting triple negative breast cancer cells 

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6-(3-Fluorophenyl)-N ${ }^{2}$-(2-methoxyphenyl)-1,3,5-triazine-2,4-diamine (1)


6-(3-Fluorophenyl)- $\mathrm{N}^{2}$-(2-methoxyphenyl)-1,3,5-triazine-2,4-diamine (1)



6-(4Chloropheryi)- $\mathrm{N}^{2}$-(p-tolyl)-1,3,5-triazine-2,4-diamine (2)




$\mathrm{N}^{2}$-(4-Chlorophenyl)-6-(4-(trifluoromethoxy)phenyl)-1,3,5-triazine-2,4-diamine (4)

$7.51(\mathrm{dd}, \mathrm{J}=0.92,8.96 \mathrm{~Hz}, 2 \mathrm{H})$
7.89 (d, J=8.91 Hz, 2 H$)$
$8.43(\mathrm{~d}, \mathrm{~J}=8.94 \mathrm{~Hz}, 2 \mathrm{H})$

$N^{2}$-(4-Chlorophenyl)-6-(4-(trifluoromethoxy)phenyl)-1,3,5-triazine-2,4-diamine (4)

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## $\mathrm{N}^{2}$-(2-Chlorophenyl)-6-4-(dimethylamino)phenyl)-1,3,5-triazine-2,4-diamine (5)



| $\circ$ |
| :--- |
| - |
| $\dot{\circ}$ |
| 1 |
| 1 |


$6.74(\mathrm{~d}, \mathrm{~J}=9.12 \mathrm{~Hz}, 2$
7.16 (ddd, $J=1.58,7.68,7.68 \mathrm{~Hz}, 1 \mathrm{H})$
7.36 (ddd, $J=1.21,7.75,7.75 \mathrm{~Hz}, 1 \mathrm{H})$
7.50 (dd, $J=1.46,8.00 \mathrm{~Hz}, 1 \mathrm{H})$
7.96 (dd, $J=1.50,8.10 \mathrm{~Hz}, 1 \mathrm{H})$
8.13 (d, $J=9.21 \mathrm{~Hz}, 2 \mathrm{H}$ )

$\mathrm{N}^{2}$-(2-Chlorophenyl)-6-(4-(dimethylamino)phenyl)-1,3,5-triazine-2,4-diamine (5)




$\mathrm{N}^{2}$-(4-Chlorophenyl)-6-(p-tolyl)-1,3,5-triazine-2,4-diamine (7)


## 6-(4-Methoxyphenyl)- $\mathbf{N}^{2}$-phenyl-1,3,5-triazine-2,4-diamine (8)



6-(4-Methoxyphenyl)- $\mathrm{N}^{2}$-phenyl-1,3,5-triazine-2,4-diamine (8)



6-(4-Methoxyphenyl)- $\mathrm{N}^{2}$-(p-tolyl)-1,3,5-triazine-2,4-diamine (9)

$N^{2}$-Phenyl-6-(3,4,5-trimethoxyphenyl)-1,3,5-triazine-2,4-diamine (10)







$\mathrm{N}^{2}$-(3-Chlorophenyl)-6-(3,4,5-trimethoxyphenyl)-1,3,5-triazine-2,4-diamine (14)



## $N^{2}$-(m-Tolyl)-6-(3,4,5-trimethoxyphenyl)-1,3,5-triazine-2,4-diamine (15)


$N^{2}$-(m-Tolyi)-6-(3,4,5-trimethoxyphenyi)-1,3,5-triazine-2,4-diamine (15)

$\mathrm{N}^{2}$-(4-Chlorophenyl)-6-(3,4,5-trimethoxyphenyl)-1,3,5-triazine-2,4-diamine (16)

$\mathrm{N}^{2}$-(4-Chiorophenyl)-6-(3,4,5-trimethoxyphenyl)-1,3,5-triazine-2,4-diamine (16)


$N^{2}$-4-Bromophenyi)-6-(3,4,5-trimethoxyphemyl)-1,3,5-triazine-2,4-diamine (17)


$N^{2}$-(p-Tolyl)-6-(3,4,5-trimethoxyphenyl)-1,3,5-triazine-2,4-diamine (18)





$\mathbf{N}^{2}$-(4--tsopropylphemyl)-6-(3,4,5-trimethoxyphenyl)-1,3,5-triazine-2,4-diamine (21)



## X-ray structure analysis of

## $N^{2}$-(4-(trifluoromethoxy)phenyl)-6-(3,4,5-trimethoxyphenyl)-1,3,5-triazine-2,4-diamine (20)

The molecular structure of $\mathbf{2 0}$, being representative of the series, has been established by single crystal Xcrystallography and is illustrated in Figure S1. The triazine ring is strictly planar with the r.m.s. deviation of the six atoms comprising the plane being $0.0059 \AA$. The $\mathrm{C}-\mathrm{N}$ bond lengths in the triazine ring lie within the relatively narrow range $1.331(3) \AA$, for $\mathrm{C} 1-\mathrm{N} 3$, to $1.354(3) \AA$, for C3-N3, suggesting substantial delocalisation of $\pi$-electron density over the ring. The appended N4, N5 and C4 atoms are close to co-planar with the ring lying, respectively, $-0.039(3), 0.049(3)$ and $0.012(3) \AA$, out of the plane of the ring. The dihedral angle formed between the central ring and the methoxy-substituted ring of $2.30(8)^{\circ}$ is consistent with a co-planar relationship, and the dihedral angle between the triazine and N5-bound rings of 11.93(7) indicates an inclined relationship. Overall, to a first approximation, the molecule is considered planar as seen in the dihedral angle between the outer rings of $10.84(7)^{\circ}$. The peripheral rings are orientated to the same side of the molecule, so the molecule has the shape of the letter $U$. The methoxy substituents adopt different conformations in order to minimise steric strain. Thus, the central methoxy group lies out of the plane of the ring to which it is connected with the C11-O2-C7-C8 torsion angle being $81.7(2)^{\circ}$ whereas the flanking methoxy groups are co-planar with the ring: $\mathrm{C} 10-\mathrm{O} 1-\mathrm{C} 6-\mathrm{C} 7$ is $179.57(18)^{\circ}$ and $\mathrm{C} 12-\mathrm{O} 3-\mathrm{C} 8-\mathrm{C} 7$ is $177.66(18)^{\circ}$. The trifluoromethoxy group lies out of the plane of the ring to which it is connected as seen in the $\mathrm{C} 19-\mathrm{O} 4-\mathrm{C} 16-\mathrm{C} 15$ torsion angle of $77.9(3)^{\circ}$.


Figure S1. Molecular structure of 20, showing atom labelling scheme and anisotropic displacement parameters at the $70 \%$ probability level.

In the crystal of 20, the primary-amine forms a hydrogen bond with a methoxy-oxygen atom to sustain a linear supramolecular chain along the $b$-axis, as shown in Figure S2(a); geometric parameters characterising the intermolecular interactions are collated in the figure caption. The two remaining acidic hydrogen atoms do not
participate in structure-directing intermolecular interactions. A supramolecular layer in the ac-plane is formed through the cooperativity of $\pi \cdots \pi$ interactions between the triazine-(N1-N3,C1-C3) and appended phenyl-(C4$\mathrm{C} 9)$ rings along with methyl $-\mathrm{C}-\mathrm{H}^{\cdots} \pi(\mathrm{C} 13-\mathrm{C} 18)$ interactions. Layer stack along the $b$-axis with no directional interactions between them.



Figure S2. Molecular packing in the crystal of 20:
(a) supramolecular linear chain sustained by amine- $\mathrm{N}-\mathrm{H}^{\cdots} \mathrm{O}$ (methoxy) hydrogen bonds (shown as orange dashed lines) $\left[\mathrm{N} 4-\mathrm{H} 2 \mathrm{n} \cdots \mathrm{O} 1^{\mathrm{i}}: \mathrm{H} 2 \mathrm{n} \cdots \mathrm{O} 1^{\mathrm{i}}=2.08(3) \AA, \mathrm{N} 4 \cdots \mathrm{O} 1^{\mathrm{i}}=2.941(3) \AA\right.$ and angle at $\mathrm{H} 2 \mathrm{n}=164(3)^{\circ}$ for symmetry operation (i) $x, 1+y, z]$;
(b) a view of the unit contents in projection down the a-axis. The $\pi \cdots \pi$ $\left[\mathrm{Cg}\right.$ (triazine) ${ }^{\cdots} \mathrm{Cg}(\text { (trimethoxyphenyl })^{\mathrm{ii}}=3.5987(13) \AA$ and angle of inclination $=2.30(11)^{\circ}$ for symmetry operation (ii) $2-\mathrm{x}, \quad 1-\mathrm{y}, \quad 1-\mathrm{z}]$ and methyl-C-H${ }^{\cdots} \pi \quad\left[\mathrm{C} 12-\mathrm{H} 12 \mathrm{a} \cdots \mathrm{Cg}(\text { trifluoromethoxyphenyl })^{\mathrm{iiii}}=2.88 \AA\right.$, $\mathrm{C} 12 \cdots \mathrm{Cg}\left(\right.$ trifluoromethoxyphenyl ${ }^{\text {iii }}=3.461(3) \AA \quad$ with $\quad$ angle at $\mathrm{H} 12 \mathrm{a}=119^{\circ} \quad$ and $\mathrm{C} 12-\mathrm{H} 12 \mathrm{c} \cdots \mathrm{Cg}(\text { trifluoromethoxyphenyl })^{\mathrm{iv}}=2.90 \AA, \mathrm{C} 12{ }^{\cdots} \mathrm{Cg}(\text { trifluoromethoxyphenyl })^{\mathrm{iv}}=3.821(3) \AA$ with angle at $\mathrm{H} 12 \mathrm{c}=157^{\circ}$ for symmetry operations (iii) $2-\mathrm{x}, 1-\mathrm{y}, 1-\mathrm{z}$ and (iv) $\left.1-\mathrm{x}, 1-\mathrm{y}, 1-\mathrm{z}\right]$ interactions are shown as blue and purple dashed lines, respectively.

Concentration-response curves for the compounds tested against
MDA-MB231 breast cancer cells

## Compound 1



Compound 3


Compound 5


Compound 2


Compound 4


Compound 6


Compound 7


Compound 9



Compound 11


Compound 12


Compound 13



## Compound 15



Compound 17


Compound 16




Compound 21


## Nilotinib



Concentration-response curves for the most active compounds tested against SKBR-3 breast cancer cells


Compound 8


Compound 13


Compound 5


Compound 10


Compound 14


## Compound 15



Compound 17



Compound 18





Concentration-response curves for the most active compounds tested against
MCF-7 breast cancer cells


Time-dependent concentration-response curves for compounds 16 and 18 tested against MDA-MB231 breast cancer cells

Compound 16


Compound 18


## Predicted ADME properties of compounds 1-21

Table S1(a). Properties calculated / predicted by QikProp for compounds 1-7

| Property / Descriptor | Compounds |  |  |  |  |  |  |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
|  | 1 | 2 | 3 | 4 | 5 | 6 | 7 |
| \#stars | 0 | 0 | 1 | 3 | 1 | 0 | 0 |
| \#rotor | 5 | 4 | 4 | 5 | 5 | 4 | 4 |
| CNS | -1 | -1 | -1 | -1 | -1 | -1 | -1 |
| mol_MW | 311.318 | 311.773 | 365.745 | 381.744 | 340.814 | 311.773 | 311.773 |
| dipole | 3.52 | 4.139 | 5.015 | 6.86 | 3.999 | 2.649 | 5.011 |
| SASA | 663.942 | 659.67 | 642.845 | 696.372 | 709.92 | 664.899 | 668.187 |
| FOSA | 88.845 | 87.813 | 22.596 | 2.005 | 153.168 | 87.856 | 87.857 |
| FISA | 112.503 | 116.153 | 108.718 | 115.201 | 109.667 | 108.614 | 116.165 |
| PISA | 420.217 | 384.054 | 357.419 | 384.981 | 381.353 | 402.696 | 392.521 |
| WPSA | 42.377 | 71.65 | 154.112 | 194.184 | 65.732 | 65.732 | 71.644 |
| volume | 1071.642 | 1076.618 | 1054.303 | 1114.602 | 1163.85 | 1074.633 | 1078.449 |
| donorHB | 3 | 3 | 3 | 3 | 3 | 3 | 3 |
| accptHB | 4.75 | 4 | 4 | 4 | 5 | 4 | 4 |
| $\operatorname{dip}^{\wedge} 2 / \mathrm{V}$ | 0.011561 | 0.015911 | 0.023854 | 0.042222 | 0.013738 | 0.006528 | 0.023283 |
| ACxDN^. ${ }^{\wedge}$ /SA | 0.012392 | 0.010503 | 0.010777 | 0.009949 | 0.012199 | 0.01042 | 0.010369 |
| glob | 0.762784 | 0.770098 | 0.779296 | 0.74657 | 0.753738 | 0.763102 | 0.761144 |
| QPpolrz | 37.536 | 38.032 | 36.884 | 38.918 | 40.855 | 38.13 | 38.186 |
| QPlogPC16 | 12.329 | 12.918 | 11.82 | 12.583 | 13.76 | 12.947 | 13.014 |
| QPlogPoct | 19.513 | 19.512 | 19.584 | 20.655 | 20.85 | 19.351 | 19.643 |
| QPlogPw | 12.925 | 12.186 | 11.942 | 12.096 | 13.007 | 12.272 | 12.264 |
| QPlogPo/w | 3.498 | 3.856 | 4.058 | 4.555 | 4.033 | 3.891 | 3.872 |
| QPlogS | -5.716 | -6.13 | -6.198 | -7.141 | -6.539 | -6.195 | -6.278 |
| CIQPlogS | -4.59 | -4.844 | -5.91 | -6.23 | -5.076 | -4.844 | -4.844 |
| QPlogHERG | -7.374 | -7.084 | -6.808 | -7.469 | -7.335 | -7.286 | -7.269 |
| QPPCaco | 849.238 | 784.193 | 922.413 | 800.653 | 903.506 | 924.508 | 783.98 |
| IP(eV) | 8.288 | 8.467 | 8.728 | 8.652 | 8.115 | 8.546 | 8.546 |
| EA(eV) | 0.659 | 0.674 | 3.22 | 0.846 | 0.349 | 0.549 | 0.567 |
| \#metab | 2 | 1 | 1 | 1 | 2 | 2 | 1 |
| QPlogKhsa | 0.229 | 0.405 | 0.352 | 0.448 | 0.421 | 0.395 | 0.404 |
| HumanOralAbsorption | 3 | 3 | 3 | 1 | 1 | 3 | 1 |
| PercentHumanOralAbsorption | 100 | 100 | 100 | 100 | 100 | 100 | 100 |
| SAfluorine | 42.377 | 0 | 88.37 | 122.533 | 0 | 0 | 0 |
| SAamideO | 0 | 0 | 0 | 0 | 0 | 0 | 0 |
| PSA | 73.984 | 66.209 | 64.771 | 74.891 | 67.919 | 64.702 | 66.211 |
| \#NandO | 6 | 5 | 5 | 6 | 6 | 5 | 5 |
| RuleOfFive | 0 | 0 | 0 | 0 | 0 | 0 | 0 |
| \#ringatoms | 18 | 18 | 18 | 18 | 18 | 18 | 18 |
| \#in56 | 18 | 18 | 18 | 18 | 18 | 18 | 18 |
| \#noncon | 0 | 0 | 0 | 0 | 0 | 0 | 0 |
| \#nonHatm | 23 | 22 | 25 | 26 | 24 | 22 | 22 |
| RuleOfThree | 1 | 1 | 1 | 1 | 1 | 1 | 1 |

Table S1(b). Properties calculated / predicted by QikProp for compounds 8-14

| Property / Descriptor | Compounds |  |  |  |  |  |  |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
|  | 8 | 9 | 10 | 11 | 12 | 13 | 14 |
| \#stars | 0 | 0 | 1 | 1 | 2 | 1 | 2 |
| \#rotor | 5 | 5 | 7 | 7 | 7 | 8 | 7 |
| CNS | -2 | -2 | -2 | -2 | -2 | -2 | -2 |
| mol_MW | 293.327 | 307.354 | 353.38 | 371.37 | 387.825 | 383.406 | 387.825 |
| dipole | 4.364 | 3.967 | 5.125 | 4.618 | 4.676 | 3.498 | 6.725 |
| SASA | 644.114 | 666.657 | 735.081 | 742.371 | 755.895 | 766.443 | 759.212 |
| FOSA | 88.878 | 176.695 | 258.471 | 258.445 | 258.533 | 347.289 | 258.535 |
| FISA | 115.866 | 115.959 | 113.811 | 110.489 | 106.442 | 110.196 | 113.995 |
| PISA | 439.37 | 374.003 | 362.799 | 331.748 | 325.172 | 308.958 | 315.079 |
| WPSA | 0 | 0 | 0 | 41.689 | 65.749 | 0 | 71.603 |
| volume | 1043.191 | 1099.821 | 1194.192 | 1208.377 | 1234.265 | 1263.357 | 1238.065 |
| donorHB | 3 | 3 | 3 | 3 | 3 | 3 | 3 |
| accptHB | 4.75 | 4.75 | 6.25 | 6.25 | 6.25 | 7 | 6.25 |
| $\operatorname{dip}^{\wedge} 2 / \mathrm{V}$ | 0.018256 | 0.014308 | 0.021993 | 0.01765 | 0.017719 | 0.009684 | 0.036532 |
| ACxDN^. ${ }^{\wedge} / \mathrm{SA}$ | 0.012773 | 0.012341 | 0.014727 | 0.014582 | 0.014321 | 0.015819 | 0.014259 |
| glob | 0.772286 | 0.772937 | 0.740536 | 0.739058 | 0.736166 | 0.737399 | 0.734454 |
| QPpolrz | 36.58 | 38.222 | 40.608 | 40.879 | 41.852 | 42.219 | 41.908 |
| QPlogPC16 | 12.45 | 12.575 | 13.569 | 13.212 | 14.156 | 13.898 | 14.229 |
| QPlogPoct | 19.145 | 19.571 | 21.089 | 21.259 | 21.678 | 21.744 | 21.997 |
| QPlogPw | 13.016 | 12.626 | 13.736 | 13.534 | 13.503 | 13.939 | 13.496 |
| QPlogPo/w | 3.167 | 3.44 | 3.433 | 3.665 | 3.934 | 3.337 | 3.914 |
| QPlogS | -5.194 | -5.578 | -6.002 | -6.308 | -6.643 | -6.15 | -6.725 |
| CIQPlogS | -4.248 | -4.514 | -4.891 | -5.237 | -5.556 | -5.205 | -5.556 |
| QPlogHERG | -7.312 | -6.99 | -7.461 | -7.336 | -7.352 | -7.255 | -7.331 |
| QPPCaco | 789.122 | 787.511 | 825.332 | 887.434 | 969.421 | 893.131 | 822.034 |
| IP(eV) | 8.492 | 8.399 | 8.468 | 8.579 | 8.528 | 8.198 | 8.625 |
| EA(eV) | 0.496 | 0.486 | 0.572 | 0.634 | 0.625 | 0.5 | 0.646 |
| \#metab | 2 | 2 | 4 | 4 | 4 | 5 | 4 |
| QPlogKhsa | 0.163 | 0.309 | 0.2 | 0.235 | 0.299 | 0.205 | 0.307 |
| HumanOralAbsorption | 3 | 3 | 3 | 1 | 1 | 3 | 1 |
| PercentHumanOralAbsorption | 100 | 100 | 100 | 100 | 100 | 100 | 100 |
| SAfluorine | 0 | 0 | 0 | 41.689 | 0 | 0 | 0 |
| SAamideO | 0 | 0 | 0 | 0 | 0 | 0 | 0 |
| PSA | 75.304 | 75.314 | 85.119 | 84.546 | 83.623 | 92.902 | 85.146 |
| \#NandO | 6 | 6 | 8 | 8 | 8 | 9 | 8 |
| RuleOfFive | 0 | 0 | 0 | 0 | 0 | 0 | 0 |
| \#ringatoms | 18 | 18 | 18 | 18 | 18 | 18 | 18 |
| \#in56 | 18 | 18 | 18 | 18 | 18 | 18 | 18 |
| \#noncon | 0 | 0 | 0 | 0 | 0 | 0 | 0 |
| \#nonHatm | 22 | 23 | 26 | 27 | 27 | 28 | 27 |
| RuleOfThree | 0 | 0 | 1 | 1 | 1 | 1 | 1 |

Table S1(c). Properties calculated / predicted by QikProp for compounds 15-21

| Property / Descriptor | Compounds |  |  |  |  |  |  |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
|  | 15 | 16 | 17 | 18 | 19 | 20 | 21 |
| \#stars | 2 | 2 | 2 | 1 | 0 | 2 | 2 |
| \#rotor | 7 | 7 | 7 | 7 | 8 | 8 | 8 |
| CNS | -2 | -2 | -2 | -2 | -2 | -2 | -2 |
| mol_MW | 367.407 | 387.825 | 432.276 | 367.407 | 383.406 | 437.378 | 395.46 |
| dipole | 4.992 | 7.55 | 7.209 | 4.732 | 5.887 | 8.48 | 4.615 |
| SASA | 766.568 | 759.198 | 764.151 | 758.052 | 750.265 | 803.458 | 800.441 |
| FOSA | 346.307 | 258.528 | 258.531 | 346.327 | 347.388 | 260.791 | 422.065 |
| FISA | 113.872 | 113.995 | 114.023 | 113.955 | 114.03 | 115.09 | 114.194 |
| PISA | 306.388 | 315.035 | 314.274 | 297.77 | 288.846 | 306.026 | 264.182 |
| WPSA | 0 | 71.64 | 77.322 | 0 | 0 | 121.551 | 0 |
| volume | 1253.024 | 1238.006 | 1246.71 | 1250.965 | 1255.543 | 1349.821 | 1348.757 |
| donorHB | 3 | 3 | 3 | 3 | 3 | 3 | 3 |
| accptHB | 6.25 | 6.25 | 6.25 | 6.25 | 7 | 6.25 | 6.25 |
| $\operatorname{dip}^{\wedge} 2 / \mathrm{V}$ | 0.019885 | 0.046045 | 0.041685 | 0.017901 | 0.027605 | 0.053274 | 0.015789 |
| ACxDN^. $5 / \mathrm{SA}$ | 0.014122 | 0.014259 | 0.014167 | 0.01428 | 0.01616 | 0.013473 | 0.013524 |
| glob | 0.733253 | 0.734444 | 0.733099 | 0.740678 | 0.750191 | 0.735167 | 0.73755 |
| QPpolrz | 42.423 | 41.905 | 42.246 | 42.259 | 41.715 | 45.652 | 45.209 |
| QPlogPC16 | 13.825 | 14.229 | 14.363 | 13.698 | 13.629 | 14.019 | 14.477 |
| QPlogPoct | 21.575 | 22.127 | 22.208 | 21.515 | 21.863 | 23.765 | 22.433 |
| QPlogPw | 13.429 | 13.496 | 13.505 | 13.35 | 13.789 | 13.345 | 13.005 |
| QPlogPo/w | 3.723 | 3.914 | 3.987 | 3.706 | 3.443 | 4.781 | 4.27 |
| QPlogS | -6.538 | -6.725 | -6.834 | -6.393 | -5.878 | -7.54 | -6.958 |
| CIQPlogS | -5.16 | -5.556 | -6.429 | -5.16 | -5.205 | -6.53 | -5.705 |
| QPlogHERG | -7.309 | -7.331 | -7.349 | -7.148 | -6.952 | -7.277 | -7.039 |
| QPPCaco | 824.23 | 822.033 | 821.517 | 822.742 | 821.392 | 802.602 | 818.458 |
| IP(eV) | 8.442 | 8.52 | 8.624 | 8.372 | 8.302 | 8.661 | 8.357 |
| EA(eV) | 0.561 | 0.642 | 0.655 | 0.559 | 0.594 | 0.684 | 0.554 |
| \#metab | 5 | 3 | 3 | 4 | 4 | 4 | 4 |
| QPlogKhsa | 0.344 | 0.307 | 0.328 | 0.345 | 0.197 | 0.555 | 0.554 |
| HumanOralAbsorption | 1 | 1 | 1 | 1 | 3 | 1 | 1 |
| PercentHumanOralAbsorption | 100 | 100 | 100 | 100 | 100 | 100 | 100 |
| SAfluorine | 0 | 0 | 0 | 0 | 0 | 121.551 | 0 |
| SAamideO | 0 | 0 | 0 | 0 | 0 | 0 | 0 |
| PSA | 85.128 | 85.141 | 85.146 | 85.136 | 94.304 | 94.193 | 85.163 |
| \#NandO | 8 | 8 | 8 | 8 | 9 | 9 | 8 |
| RuleOfFive | 0 | 0 | 0 | 0 | 0 | 0 | 0 |
| \#ringatoms | 18 | 18 | 18 | 18 | 18 | 18 | 18 |
| \#in56 | 18 | 18 | 18 | 18 | 18 | 18 | 18 |
| \#noncon | 0 | 0 | 0 | 0 | 0 | 0 | 0 |
| \#nonHatm | 27 | 27 | 27 | 27 | 28 | 31 | 29 |
| RuleOfThree | 1 | 1 | 1 | 1 | 1 | 1 | 1 |

## QikProp descriptors and properties

\#stars = Number of property or descriptor values that fall outside the $95 \%$ range of similar values for known drugs.
\#rotor = Number of non-trivial (not CX3), non-hindered (not alkene, amide, small ring) rotatable bonds.
CNS $=$ Predicted central nervous system activity on a -2 (inactive) to +2 (active) scale.
Dipole $=$ Computed dipole moment of the molecule.
SASA = Total solvent accessible surface area (SASA) in square angstroms using a probe with a $1.4 \AA$ radius.
FOSA = Hydrophobic component of the SASA (saturated carbon and attached hydrogen).
FISA = Hydrophilic component of the SASA (SASA on N, O, and H on heteroatoms).
PISA $=\pi$ (carbon and attached hydrogen) component of the SASA.
WPSA = Weakly polar component of the SASA (halogens, P, and S).
Volume $=$ Total solvent-accessible volume in cubic angstroms using a probe with a $1.4 \AA$ radius. $500.0-$ 2000.0
donorHB = Estimated number of hydrogen bonds that would be donated by the solute to water molecules in an aqueous solution.
accptHB $=$ Estimated number of hydrogen bonds that would be accepted by the solute from water molecules. $\boldsymbol{d i p}^{\wedge} \mathbf{2} / \mathbf{V}=$ Square of the dipole moment divided by the molecular volume. This is the key term in the Kirkwood-Onsager equation for the free energy of solvation of a dipole with volume V .
$\mathbf{A C x D N} \wedge .5 / \mathbf{S A}=$ Index of cohesive interaction in solids, $\left.(\operatorname{accptHB})(\text { donorHB })^{1 / 2}\right) /(\mathrm{SA})$
glob $=$ Globularity descriptor, $\left(4 \pi r^{2}\right) /(\mathrm{SASA}), \mathrm{r}$ is the radius of a sphere with a volume equal to the molecular volume. Globularity is 1.0 for a spherical molecule.
QPpolrz = Predicted polarizability in cubic angstroms.
QPlogPC16 = Predicted hexadecane/gas partition coefficient.
QPlogPoct $=$ Predicted octanol/gas partition coefficient.
QPlogPw = Predicted water/gas partition coefficient.
QPlogPo/w = Predicted octanol/water partition coefficient.
QPlogS = Predicted aqueous solubility, $\log \mathrm{S} . \mathrm{S}$ in mol dm-3 is the concentration of the solute in a saturated solution that is in equilibrium with the crystalline solid.
CIQPlogS $=$ Conformation-independent predicted aqueous solubility, $\log \mathrm{S} . \mathrm{S}$ in mol dm-3 is the concentration of the solute in a saturated solution that is in equilibrium with the crystalline solid.
QPlogHERG = Predicted IC50 value for blockage of HERG K+ channels.
QPPCaco $=$ Predicted apparent Caco-2 cell permeability in nm/sec. Caco-2 cells are a model for the gut blood barrier.
$\mathbf{I P}(\mathbf{e v})=$ PM3 calculated ionization potential.
$\mathbf{E A}(\mathbf{e V})=$ PM3 calculated electron affinity.
\#metab = Number of likely metabolic reactions.

QPlogKhsa = Prediction of binding to human serum albumin.
HumanOralAbsorption $=$ Predicted qualitative human oral absorption:
PercentHuman-OralAbsorption $=$ Predicted human oral absorption on 0 to $100 \%$ scale. The prediction is based on a quantitative multiple linear regression model. This property usually correlates well with HumanOral-Absorption, as both measure the same property.
SAFluorine $=$ Solvent-accessible surface area of fluorine atoms.
SAamideO = Solvent-accessible surface area of amide oxygen atoms.
PSA = Van der Waals surface area of polar nitrogen and oxygen atoms.
\#NandO = Number of nitrogen and oxygen atoms.
RuleOfFive $=$ Number of violations of Lipinski's rule of five. The rules are: mol_MW $<500$, QPlogPo/w $<5$, donorHB $\leq 5$, accptHB $\leq 10$. Compunds that satisfy these rules are considered drug-like.
RuleOfThree $=$ Number of violations of Jorgensen's rule of three. The three rules are: QPlogS >-5.7, QP PCaco > $22 \mathrm{~nm} / \mathrm{s}$, \# Primary Metabolites < 7. Compounds with fewer (and preferably no) violations of these rules are more likely to be orally available.
\#ringatoms = Number of atoms in rings.
\#in56 = Number of atoms in 5- or 6-membered rings.
\#noncon = Number of ring atoms not able to form conjugated aromatic systems (e.g. sp3 C).
\#nonHatm = Number of heavy atoms (nonhydrogen atoms).


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