

Supporting Information

Theoretical investigations on interactions of arylsulphonyl indazole derivatives as potential ligands of VEGFR2 kinase

Kornelia Czaja ¹, Jacek Kujawski ^{1*}, Paweł Śliwa ², Rafał Kurczab ³, Radosław Kujawski ⁴, Anna Stodolna ¹, Agnieszka Myślińska ¹ and Marek K. Bernard ^{1*}

¹Chair and Department of Organic Chemistry, Faculty of Pharmacy, Poznan University of Medical Sciences, ul. Grunwaldzka 6, 60-780 Poznań, Poland

²Cracow University of Technology, Faculty of Chemical Engineering and Technology, ul. Warszawska 24, 31-155 Kraków, Poland

³Maj Institute of Pharmacology, Polish Academy of Sciences, ul. Smętna 12, 31-343 Kraków, Poland

⁴Chair and Department of Pharmacology, Faculty of Pharmacy, Poznan University of Medical Sciences, ul. Rokietnicka 5a, 60-806 Poznań, Poland

* Correspondence: mbernard@ump.edu.pl (M.K.B.), jacekkuj@ump.edu.pl (J.K.), phone 48618546670, fax 48618546680

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Fig. S1 Electrostatic potential (ESP) map of docked azole **5** (1st pose) calculated at the B3LYP/6-311++G(2d,3p)//B3LYP-631G(d,p) level of theory (gaseous phase); isovalue = 0.002 a.u.

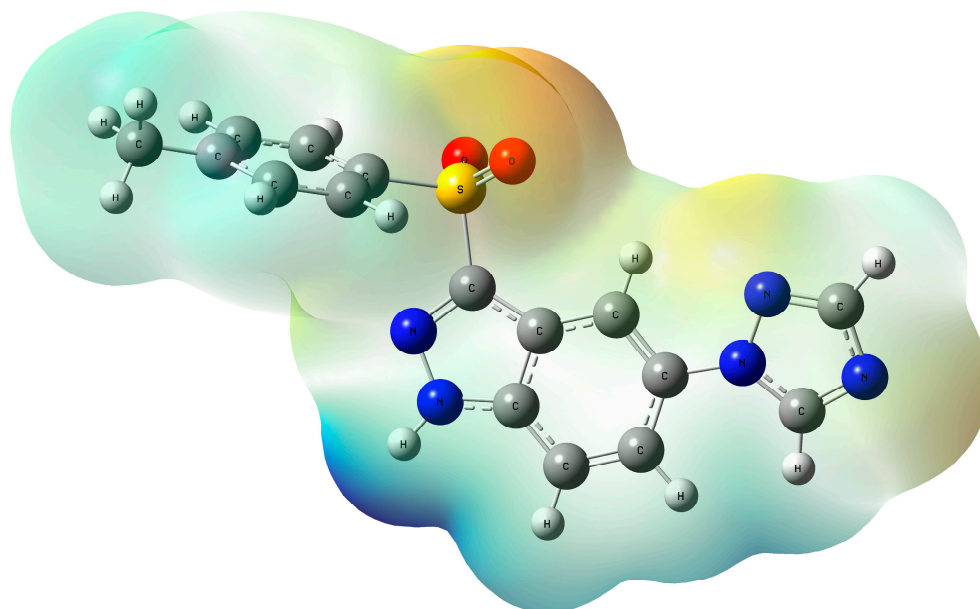
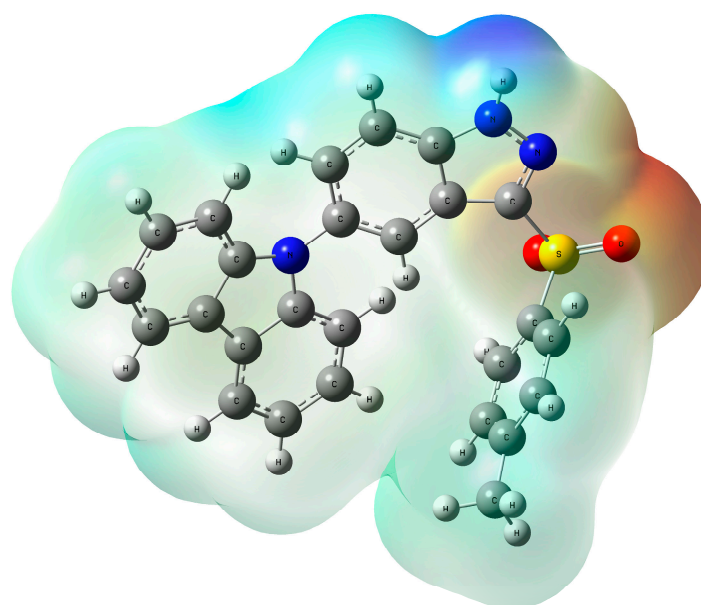


Fig. S2 Electrostatic potential (ESP) map of docked azole **7** (1st pose) calculated at the B3LYP/6-311++G(2d,3p)//B3LYP-631G(d,p) level of theory (gaseous phase); isovalue = 0.002 a.u.



Electrostatic potential (ESP) map of ligands **1–4**, **6**, **8–9** are given in the our previous reports:

Czaja K., Kujawski J., Kamel K., Bernard M.K. Selected arylsulphonyl pyrazole derivatives as potential Chk1 kinase ligands – computational investigations. *J. Mol. Model.* **2020**, *accepted*. doi: 10.1007/s00894-020-04407-3.

Fig. S3: Geometry of the first poses of azoles **1–9** after docking procedure to 3ewh.pdb protein and their Cartesian coordinates (charge=0, multiplicity=1):

Azole 1

Cartesian coordinates:

C	16.051000000	-6.986000000	7.723000000
C	16.267000000	-7.422000000	6.376000000
C	15.464000000	-8.585000000	6.273000000
N	14.861000000	-8.735000000	7.496000000
N	15.217000000	-7.777000000	8.374000000
C	15.387000000	-9.338000000	5.092000000
C	16.137000000	-8.898000000	4.015000000
C	16.940000000	-7.737000000	4.116000000
C	17.028000000	-6.987000000	5.275000000
H	14.227000000	-9.457000000	7.797000000
Cl	17.851000000	-7.251000000	2.718000000
S	16.782000000	-5.579000000	8.557000000
O	15.773000000	-4.511000000	8.458000000
O	18.145000000	-5.394000000	8.045000000
C	16.896000000	-6.080000000	10.275000000
C	16.053000000	-5.490000000	11.216000000
C	16.154000000	-5.878000000	12.551000000
C	17.084000000	-6.845000000	12.956000000
C	17.209000000	-7.237000000	14.407000000
C	17.919000000	-7.420000000	11.985000000
C	17.836000000	-7.044000000	10.648000000
H	14.770000000	-10.222000000	5.026000000
H	16.109000000	-9.448000000	3.086000000
H	17.652000000	-6.108000000	5.332000000
H	15.333000000	-4.744000000	10.915000000
H	15.504000000	-5.426000000	13.286000000
H	18.639000000	-8.168000000	12.281000000
H	18.487000000	-7.490000000	9.911000000
H	16.478000000	-6.684000000	14.997000000
H	18.213000000	-7.004000000	14.761000000
H	17.026000000	-8.306000000	14.511000000

Azole 2

Cartesian coordinates:

C	17.03500	-7.04800	5.32300
C	16.92400	-7.75100	4.15000
C	16.06100	-8.88300	4.05600
C	15.32600	-9.31800	5.11900
C	15.43900	-8.58800	6.31900
N	14.84800	-8.75400	7.57000

H	14.15800	-9.45200	7.85900
N	15.25100	-7.77200	8.46700
C	16.09700	-7.02300	7.78600
C	16.26600	-7.47400	6.42300
N	17.67800	-7.35300	2.98300
C	18.67200	-6.37000	2.93200
C	19.11800	-6.27900	1.66300
C	18.39200	-7.23000	0.87500
C	17.52500	-7.86400	1.69000
S	16.86700	-5.57500	8.51400
O	18.59400	-5.55200	7.84300
O	15.60300	-4.20700	8.45400
C	16.99200	-6.07100	10.24500
C	16.17100	-5.47100	11.18800
C	16.25600	-5.87100	12.51100
C	17.14900	-6.86700	12.90000
C	17.95600	-7.46300	11.93400
C	17.25200	-7.28700	14.36400
C	17.88000	-7.07200	10.60600
H	17.69200	-6.19400	5.40000
H	15.98800	-9.41000	3.11600
H	14.68500	-10.18400	5.04400
H	19.02300	-5.78300	3.76800
H	19.88500	-5.60900	1.30500
H	18.51700	-7.40700	-0.18300
H	16.83000	-8.63600	1.39400
H	15.47300	-4.70100	10.89400
H	15.62200	-5.40500	13.25100
H	18.64900	-8.23900	12.22300
H	18.50500	-7.54200	9.86200
H	18.00000	-8.07300	14.46500
H	17.54400	-6.42800	14.96800
H	16.28600	-7.65900	14.70400

Azole 3

Cartesian coordinates:

C	16.03800	-7.00000	7.70700
C	16.24500	-7.42800	6.35600
C	15.44900	-8.59400	6.25700
N	14.86300	-8.75700	7.48600
N	15.21900	-7.80000	8.36600
C	15.36700	-9.33300	5.06600
C	16.10500	-8.88500	3.98900
C	16.90700	-7.71600	4.08000
C	16.97800	-6.97300	5.24700
H	14.23600	-9.48600	7.78900
S	16.77700	-5.59700	8.53800
O	18.14100	-5.41500	8.02800
O	15.77400	-4.52100	8.43800
C	16.88800	-6.09400	10.25900

C	16.02000	-5.52800	11.19100
C	16.11800	-5.91400	12.52700
C	17.07200	-6.85300	12.94100
C	17.93200	-7.40300	11.97900
C	17.19400	-7.24200	14.39400
C	17.85200	-7.02900	10.64000
N	17.64900	-7.32400	2.93500
C	18.62700	-6.37300	2.84800
C	19.04600	-6.34700	1.53600
C	18.26000	-7.33200	0.89400
N	17.41700	-7.92300	1.73500
H	14.75100	-10.21700	4.99500
H	16.07200	-9.43400	3.05900
H	17.57600	-6.07500	5.30200
H	15.28300	-4.80100	10.88300
H	15.44700	-5.48200	13.25500
H	18.67000	-8.13000	12.28300
H	18.52400	-7.45500	9.90900
H	19.00000	-5.75900	3.65400
H	19.80400	-5.71600	1.09600
H	18.33600	-7.57500	-0.15600
H	16.44200	-6.71100	14.97700
H	18.18800	-6.98000	14.75800
H	17.04100	-8.31600	14.49700

Azole 4

Cartesian coordinates:

C	17.04700	-7.07600	5.28200
C	16.26900	-7.48200	6.38100
C	16.94000	-7.80700	4.10800
C	15.43500	-8.62100	6.28400
C	16.10000	-8.95200	4.02600
C	15.34900	-9.37700	5.10300
C	16.06300	-7.03300	7.72400
N	15.20700	-7.79600	8.38100
N	14.83500	-8.75500	7.51000
H	14.17000	-9.45000	7.80800
S	16.78000	-5.60400	8.52700
O	18.14400	-5.49500	7.97300
O	15.83200	-4.48800	8.43600
C	16.91800	-6.07500	10.25200
C	16.08900	-5.46800	11.19600
C	16.19300	-5.85100	12.53200
C	17.11600	-6.82500	12.93700
C	17.24400	-7.21200	14.39000
C	17.93100	-7.42200	11.96300
C	17.84000	-7.05600	10.62400
N	17.67700	-7.44800	2.95000
N	17.44100	-8.13700	1.79200
C	18.22900	-7.57400	0.88200

C	18.24800	-8.08700	-0.52500
C	18.97400	-6.50700	1.44500
C	18.60000	-6.44000	2.77200
C	19.02600	-5.46600	3.82600
H	17.70600	-6.22300	5.35000
H	16.04900	-9.50200	3.09800
H	14.72200	-10.25400	5.04100
H	15.37800	-4.71300	10.89600
H	15.55100	-5.38900	13.26800
H	18.64200	-8.17900	12.25800
H	18.47200	-7.52300	9.88400
H	19.68900	-5.87500	0.93900
H	18.95100	-7.50100	-1.11700
H	18.55600	-9.13300	-0.52600
H	17.25100	-8.00200	-0.95600
H	19.79000	-4.80300	3.42100
H	18.16600	-4.87700	4.14500
H	19.43200	-6.00900	4.68000
H	16.53000	-6.64100	14.98300
H	18.25500	-6.99800	14.73500
H	17.04000	-8.27700	14.50200

Azole 5

Cartesian coordinates:

C	16.98300	-6.97900	5.25600
C	16.90800	-7.72700	4.09200
C	16.10300	-8.89100	3.99600
C	15.36000	-9.33500	5.07300
C	15.44500	-8.59400	6.26200
N	14.85500	-8.75000	7.49000
H	14.22300	-9.47500	7.79300
N	15.21300	-7.79400	8.36900
C	16.03900	-6.99800	7.71300
C	16.24600	-7.43000	6.36300
N	17.65800	-7.33800	2.94900
N	18.63000	-6.38300	3.02900
C	19.06400	-6.31700	1.78100
N	18.44400	-7.15300	0.89900
C	17.56800	-7.77600	1.66300
S	16.78700	-5.59700	8.54200
O	18.15000	-5.42500	8.02600
O	15.78700	-4.51900	8.44000
C	16.89900	-6.09300	10.26200
C	17.84700	-7.04600	10.64000
C	17.92900	-7.41700	11.97800
C	17.08400	-6.84900	12.94400
C	16.14500	-5.89300	12.53400
C	17.20700	-7.23800	14.39700
C	16.04600	-5.50800	11.19800
H	17.58500	-6.08400	5.31000

H	16.07000	-9.43900	3.06600
H	14.73900	-10.21600	5.00300
H	19.85600	-5.64700	1.48100
H	16.87600	-8.52900	1.31700
H	18.50500	-7.48700	9.90600
H	18.65700	-8.15600	12.27900
H	15.48700	-5.44700	13.26500
H	15.32100	-4.76900	10.89200
H	16.46800	-6.69200	14.98400
H	18.20700	-6.99400	14.75500
H	17.03500	-8.30900	14.50200

Azole 6

Cartesian coordinates:

C	16.06000	-6.99700	7.72800
C	16.28200	-7.40700	6.37400
C	15.46900	-8.56100	6.24400
N	14.86800	-8.73800	7.46400
N	15.21800	-7.79500	8.36100
C	15.39200	-9.28300	5.04400
C	16.15700	-8.82700	3.98600
C	16.96600	-7.66200	4.09300
C	17.02900	-6.94000	5.27900
H	14.22200	-9.45800	7.74500
S	16.73900	-5.57000	8.57100
O	18.10500	-5.42500	8.03700
O	15.76600	-4.47400	8.48600
C	16.86400	-6.07100	10.28900
C	17.82100	-7.01800	10.65500
C	17.91900	-7.39400	11.99300
C	17.07700	-6.84000	12.96600
C	17.21000	-7.23000	14.41800
C	16.12500	-5.88900	12.56500
C	16.01300	-5.49800	11.23500
N	17.70100	-7.24800	2.95400
C	18.72700	-6.30500	2.97500
C	19.22100	-6.11500	1.71700
C	18.47600	-6.97100	0.83600
C	18.50700	-7.20300	-0.55000
C	17.53100	-7.67100	1.63500
C	17.60500	-8.10200	-1.10300
C	16.66500	-8.77200	-0.29700
C	16.61100	-8.56700	1.07700
H	14.76300	-10.15600	4.95000
H	16.14000	-9.37000	3.05300
H	17.63300	-6.04800	5.35700
H	18.47500	-7.45200	9.91300
H	18.65700	-8.12600	12.28500
H	15.46800	-5.45300	13.30300
H	15.27900	-4.76300	10.93900

H	19.08100	-5.79700	3.86000
H	20.02200	-5.44700	1.43600
H	19.22200	-6.68900	-1.17600
H	17.62500	-8.29000	-2.16600
H	15.97100	-9.46000	-0.75600
H	15.88600	-9.08000	1.69100
H	17.99800	-7.97600	14.52400
H	17.46200	-6.34900	15.00900
H	16.26600	-7.64600	14.77000

Azole 7

Cartesian coordinates:

C	17.39000	-7.13500	1.97600
N	16.52300	-7.95000	1.35600
N	15.78900	-8.59300	2.29400
H	15.08800	-9.24400	2.01200
C	16.17200	-8.20300	3.57800
C	15.72000	-8.59900	4.84600
C	16.32100	-8.02300	5.94500
C	17.35600	-7.06400	5.78100
C	17.80400	-6.66400	4.53100
C	17.21800	-7.23700	3.40300
N	17.93300	-6.48800	6.94200
C	17.22000	-6.29700	8.14700
C	16.09800	-6.97300	8.62200
C	15.62500	-6.62100	9.88100
C	16.25600	-5.63200	10.64900
C	17.38400	-4.96500	10.18100
C	17.87300	-5.28700	8.92200
C	18.98100	-4.80100	8.13100
C	19.94900	-3.83000	8.34800
C	20.88100	-3.58200	7.34400
C	20.85100	-4.29100	6.13500
C	19.89700	-5.27400	5.89100
C	18.96900	-5.52800	6.89900
S	18.53100	-6.15400	1.03400
O	18.05900	-6.13400	-0.33300
O	19.86300	-6.60000	1.36600
C	18.29900	-4.54700	1.71400
C	19.30200	-3.96800	2.49100
C	19.17300	-2.65100	2.92200
C	18.03200	-1.91000	2.58800
C	17.00500	-2.51600	1.86100
C	17.93400	-0.48100	2.99000
C	17.13300	-3.83500	1.43500
H	14.93100	-9.32800	4.95700
H	16.00200	-8.30300	6.93800
H	18.58700	-5.92700	4.43400
H	15.61500	-7.74000	8.03500
H	14.75300	-7.12000	10.27600

H	15.85900	-5.38300	11.62200
H	17.86900	-4.21200	10.78500
H	19.97800	-3.27900	9.27600
H	21.64100	-2.83000	7.49900
H	21.58600	-4.06900	5.37500
H	19.87700	-5.81800	4.95800
H	20.17800	-4.54100	2.75800
H	19.95400	-2.19900	3.51500
H	16.10900	-1.96000	1.62900
H	16.33000	-4.30700	0.88900
H	16.98400	-0.07100	2.64700
H	18.75500	0.07900	2.54200
H	17.99200	-0.40400	4.07600

Azole 8

Cartesian coordinates:

O	14.76600	-6.12700	8.96700
S	16.22000	-6.00400	8.74400
O	16.89000	-4.70000	8.82700
C	16.59500	-6.72200	7.14300
C	17.79200	-6.38800	6.50800
C	18.06800	-6.93900	5.26000
C	17.16500	-7.81600	4.63700
C	17.48800	-8.42600	3.29500
C	15.96700	-8.12100	5.29600
C	15.67400	-7.58200	6.54800
C	16.97000	-7.10900	9.93000
N	17.63600	-8.17700	9.49200
N	18.11700	-8.80300	10.58400
H	18.66100	-9.64000	10.45900
C	17.77600	-8.15000	11.73800
C	17.02100	-7.03700	11.35100
C	17.85700	-8.09700	13.14800
N	16.69400	-6.37500	12.49900
N	17.19500	-7.02500	13.58100
C	15.87000	-5.19200	12.69800
H	18.49300	-5.71300	6.97700
H	18.99300	-6.68800	4.76200
H	15.25800	-8.78600	4.82600
H	14.74900	-7.82700	7.04800
H	18.37300	-8.81100	13.77400
H	16.66800	-9.07300	2.98300
H	18.40300	-9.01200	3.37400
H	17.62600	-7.63400	2.55900
H	15.55600	-4.80100	11.73000
H	14.99100	-5.45600	13.28600
H	16.44600	-4.43300	13.22700

Azole 9

Cartesian coordinates:

C	17.09300	-7.24300	5.16600
C	16.85800	-8.01100	3.98100
C	18.06800	-6.22400	5.09800
N	17.52000	-7.78500	2.81500
C	18.72800	-6.00900	3.90500
C	18.41600	-6.81300	2.79000
C	16.31400	-7.57800	6.32700
C	15.38900	-8.63800	6.25100
C	15.89800	-9.08000	3.97300
C	15.16800	-9.40700	5.07900
C	16.16600	-7.12000	7.67700
N	15.25600	-7.81100	8.34800
N	14.79800	-8.72900	7.47700
H	14.09400	-9.38300	7.78500
S	16.91800	-5.71300	8.50600
O	18.30500	-5.61200	8.01400
O	16.00600	-4.56700	8.36700
C	16.97800	-6.18700	10.23100
C	17.89900	-7.15300	10.63900
C	17.97100	-7.48700	11.98900
C	17.13900	-6.86900	12.93400
C	17.20500	-7.25800	14.39100
C	16.23200	-5.89200	12.49400
C	16.14500	-5.54600	11.14800
H	18.29400	-5.62200	5.96600
H	19.47500	-5.23300	3.82700
H	18.93800	-6.62500	1.86400
H	15.74900	-9.64300	3.06300
H	14.45300	-10.21600	5.06300
H	18.54500	-7.63400	9.91900
H	18.68000	-8.23500	12.31400
H	15.59100	-5.40100	13.21100
H	15.44400	-4.79300	10.81900
H	17.96600	-8.02600	14.52800
H	16.23700	-7.64600	14.70800
H	17.45900	-6.38300	14.98900

Fig. S4 Calculated interaction energies (E_{tot} ; kcal/mol) and the contributions to the total energy (E_{es} , E_{ex} , $E_{\text{ct}} + \text{mix}$, E_{dis} , G_{sol} ; kcal/mol) between docked azole **1** and selected residues of 3ewh.pdb kinase(*GAMESS* program).

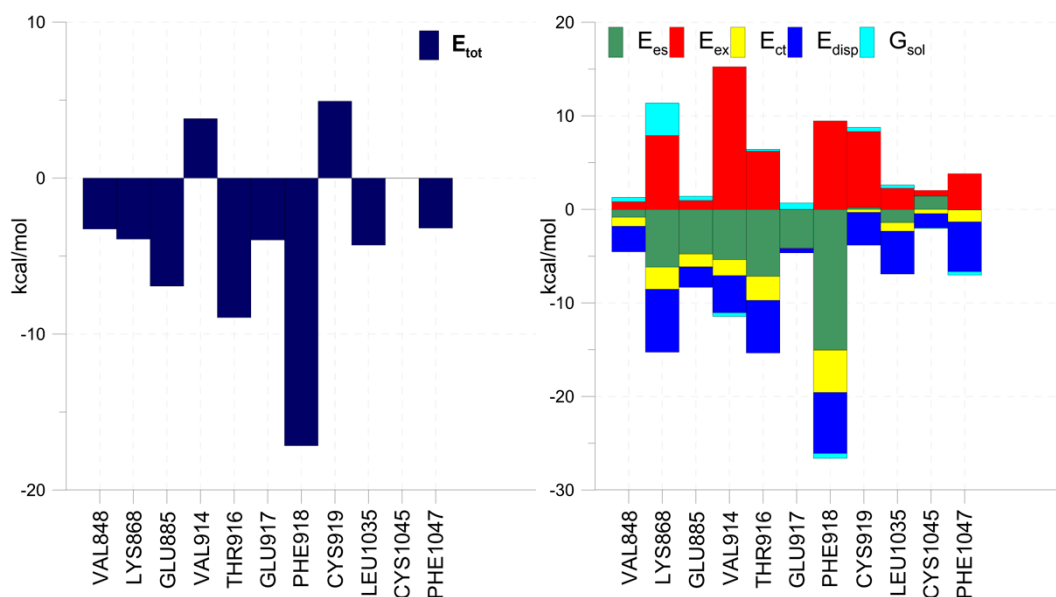


Fig. S5 Calculated interaction energies (E_{tot} ; kcal/mol) and the contributions to the total energy (E_{es} , E_{ex} , $E_{\text{ct}} + \text{mix}$, E_{dis} , G_{sol} ; kcal/mol) between docked azole **2** and selected residues of 3ewh.pdb kinase(*GAMESS* program).

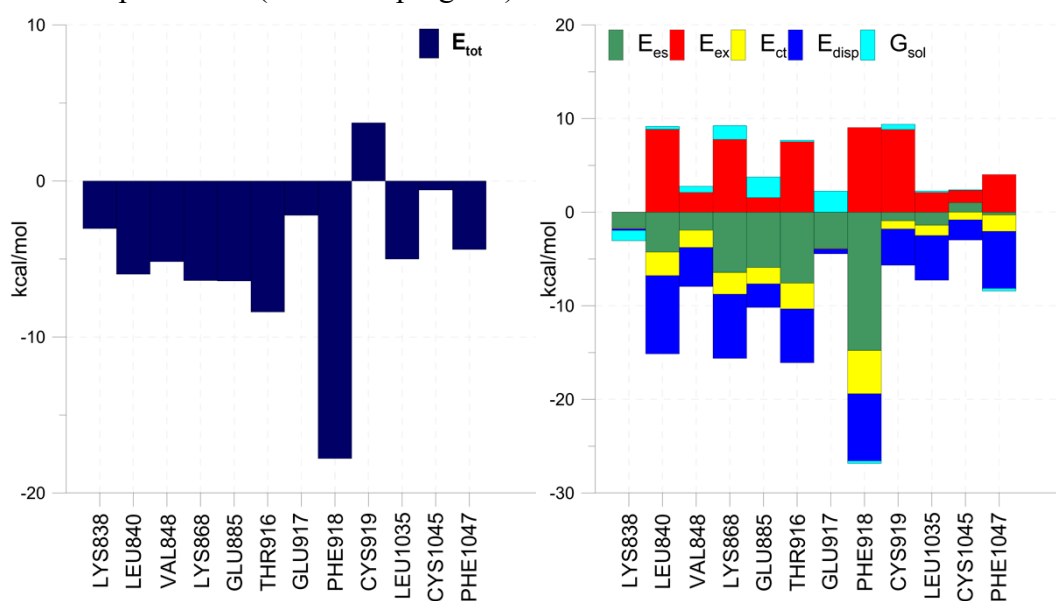


Fig. S6 Calculated interaction energies (E_{tot} ; kcal/mol) and the contributions to the total energy (E_{es} , E_{ex} , $E_{ct} + mix$, E_{dis} , G_{sol} ; kcal/mol) between docked azole **3** and selected residues of 3ewh.pdb kinase(*GAMESS* program).

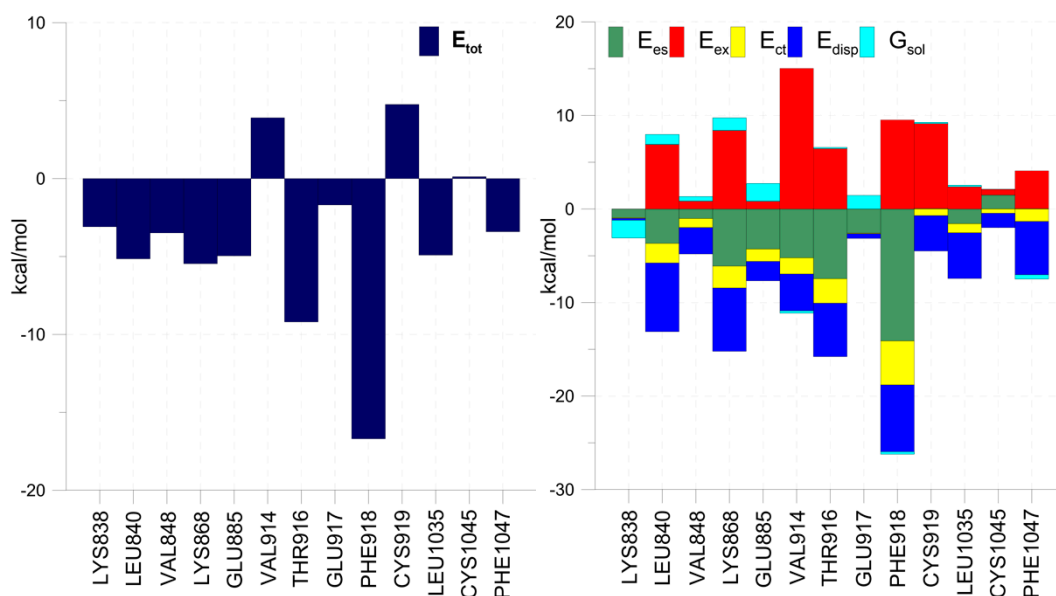


Fig. S7 Calculated interaction energies (E_{tot} ; kcal/mol) and the contributions to the total energy (E_{es} , E_{ex} , $E_{ct} + mix$, E_{dis} , G_{sol} ; kcal/mol) between docked azole **4** and selected residues of 3ewh.pdb kinase(*GAMESS* program).

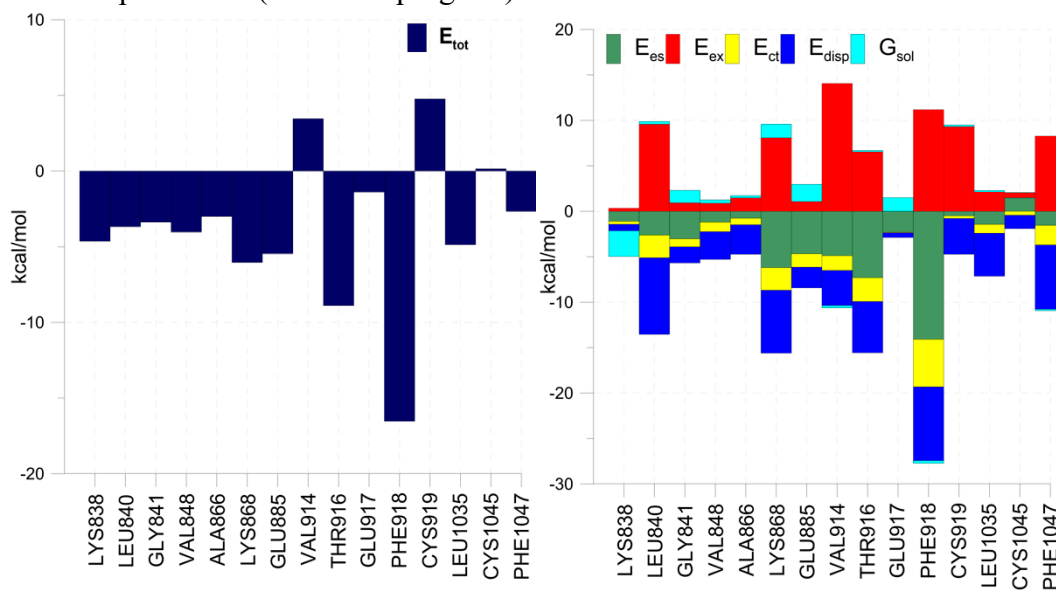


Fig. S8 Calculated interaction energies (E_{tot} ; kcal/mol) and the contributions to the total energy (E_{es} , E_{ex} , $E_{ct} + mix$, E_{dis} , G_{sol} ; kcal/mol) between docked azole **5** and selected residues of 3ewh.pdb kinase(*GAMESS* program).

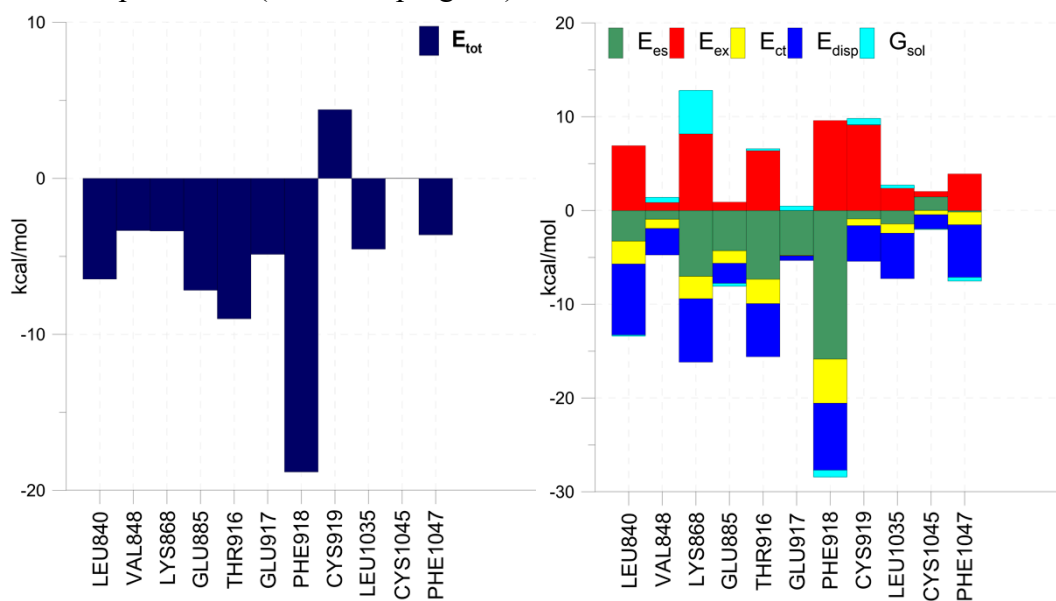


Fig. S9 Calculated interaction energies (E_{tot} ; kcal/mol) and the contributions to the total energy (E_{es} , E_{ex} , $E_{ct} + mix$, E_{dis} , G_{sol} ; kcal/mol) between docked azole **7** and selected residues of 3ewh.pdb kinase(*GAMESS* program).

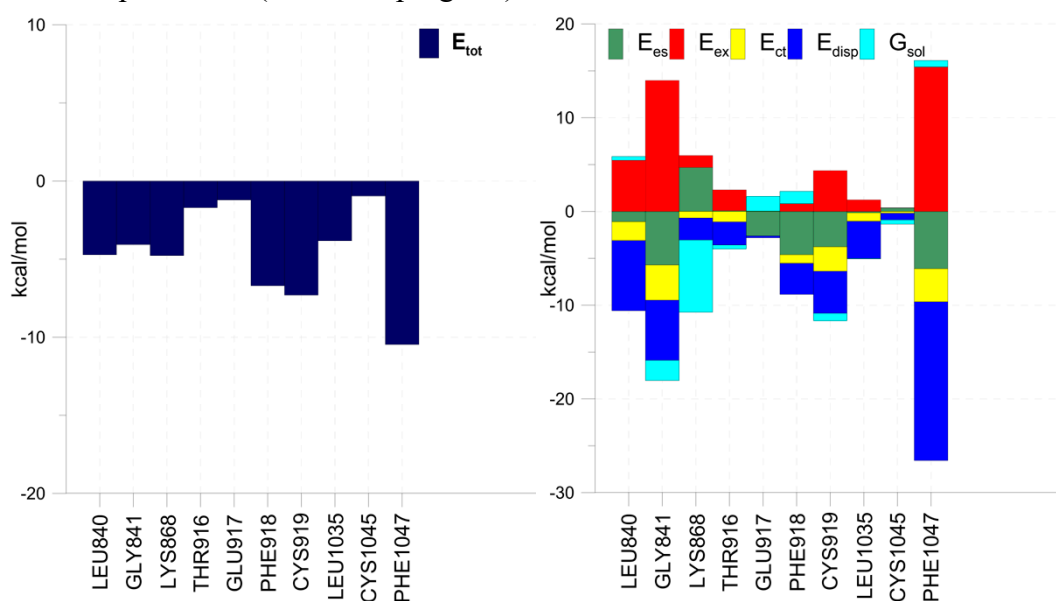


Fig. S10 Calculated interaction energies (E_{tot} ; kcal/mol) and the contributions to the total energy (E_{es} , E_{ex} , $E_{ct} + mix$, E_{dis} , G_{sol} ; kcal/mol) between docked azole **8** and selected residues of 3ewh.pdb kinase(*GAMESS* program).

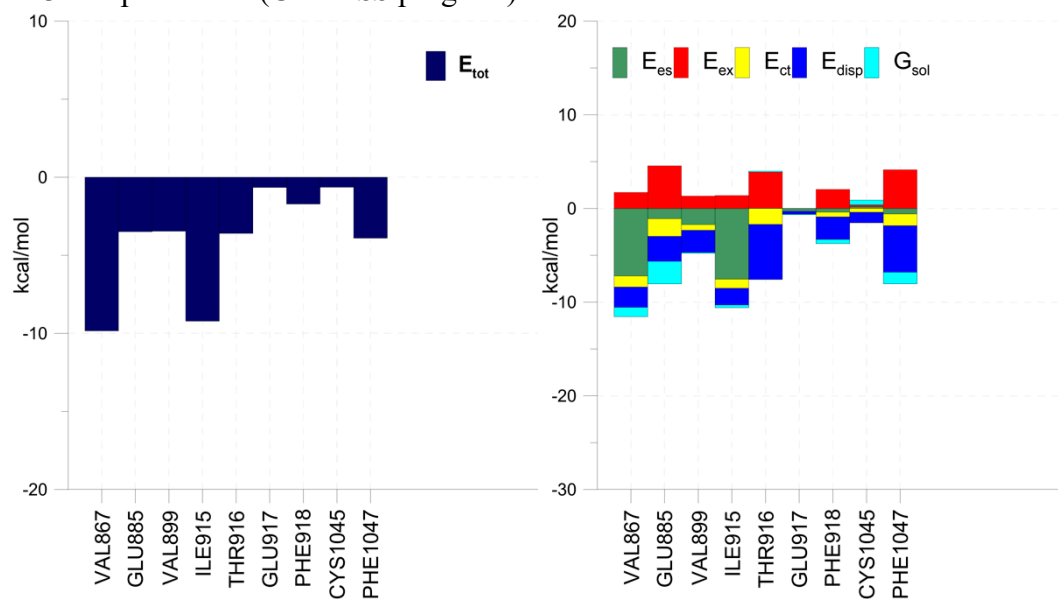


Fig. S11 Calculated interaction energies (E_{tot} ; kcal/mol) and the contributions to the total energy (E_{es} , E_{ex} , $E_{ct} + mix$, E_{dis} , G_{sol} ; kcal/mol) between docked azole **9** and selected residues of 3ewh.pdb kinase(*GAMESS* program).

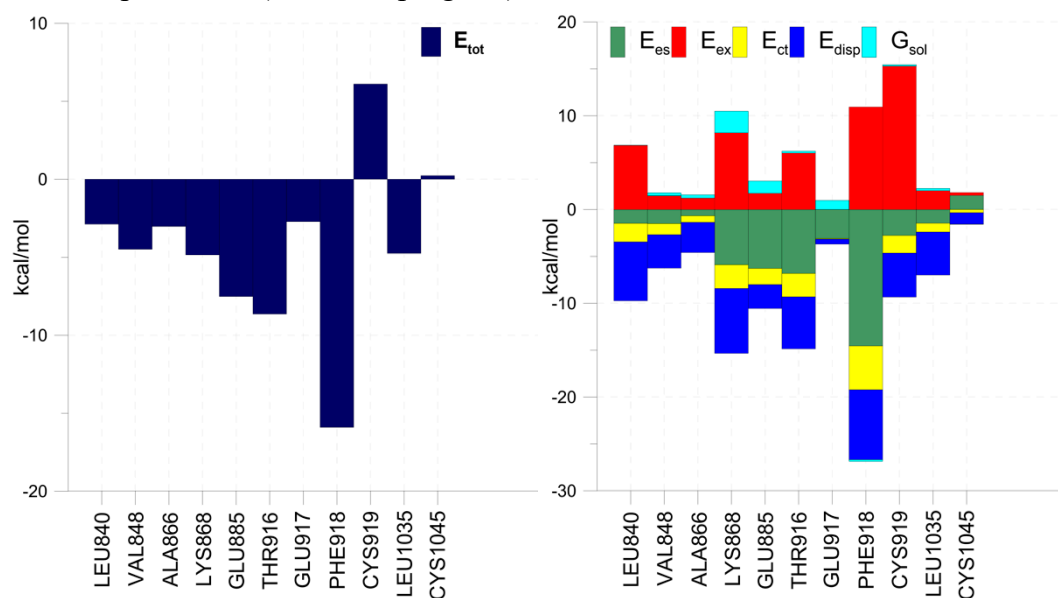


Fig. S12 Example of input file for SAPt calculations regarding the **1–Glu917** complex:
memory 12 Gb

```
molecule {  
0 1  
N    13.09300   -12.93400    9.83300  
H    12.37400   -12.58900   10.46800  
C    12.68400   -13.52800    8.56300  
H    13.03400   -14.58600    8.53500  
C    13.30300   -12.77600    7.38300  
O    13.43000   -11.55100    7.41100  
C    11.16200   -13.51300    8.43700  
H    10.83100   -14.12700    7.56700  
H    10.69100   -14.09500    9.26300  
C    10.57300   -12.10200    8.36200  
H    11.10600   -11.49200    7.59600  
H    10.83000   -11.51800    9.27600  
C    9.07900    -12.08400    8.11700  
O    8.59300    -12.87700    7.28200  
O    8.37100    -11.22400    8.77700  
H    14.07100   -12.86600   10.07800  
H    7.84600    -10.58600    9.26700  
O    13.69900   -13.43600    6.34100  
H    13.99200   -13.92500    5.56900  
--  
C    16.05100   -6.98600    7.72300  
C    16.26700   -7.42200    6.37600  
C    15.46400   -8.58500    6.27300  
N    14.86100   -8.73500    7.49600  
N    15.21700   -7.77700    8.37400  
C    15.38700   -9.33800    5.09200  
C    16.13700   -8.89800    4.01500  
C    16.94000   -7.73700    4.11600  
C    17.02800   -6.98700    5.27500  
H    14.22700   -9.45700    7.79700  
Cl   17.85100   -7.25100    2.71800  
S    16.78200   -5.57900    8.55700  
O    15.77300   -4.51100    8.45800  
O    18.14500   -5.39400    8.04500  
C    16.89600   -6.08000   10.27500  
C    16.05300   -5.49000   11.21600  
C    16.15400   -5.87800   12.55100  
C    17.08400   -6.84500   12.95600  
C    17.20900   -7.23700   14.40700  
C    17.91900   -7.42000   11.98500  
C    17.83600   -7.04400   10.64800  
H    14.77000   -10.22200    5.02600  
H    16.10900   -9.44800    3.08600  
H    17.65200   -6.10800    5.33200
```

H	15.33300	-4.74400	10.91500
H	15.50400	-5.42600	13.28600
H	18.63900	-8.16800	12.28100
H	18.48700	-7.49000	9.91100
H	16.47800	-6.68400	14.99700
H	18.21300	-7.00400	14.76100
H	17.02600	-8.30600	14.51100

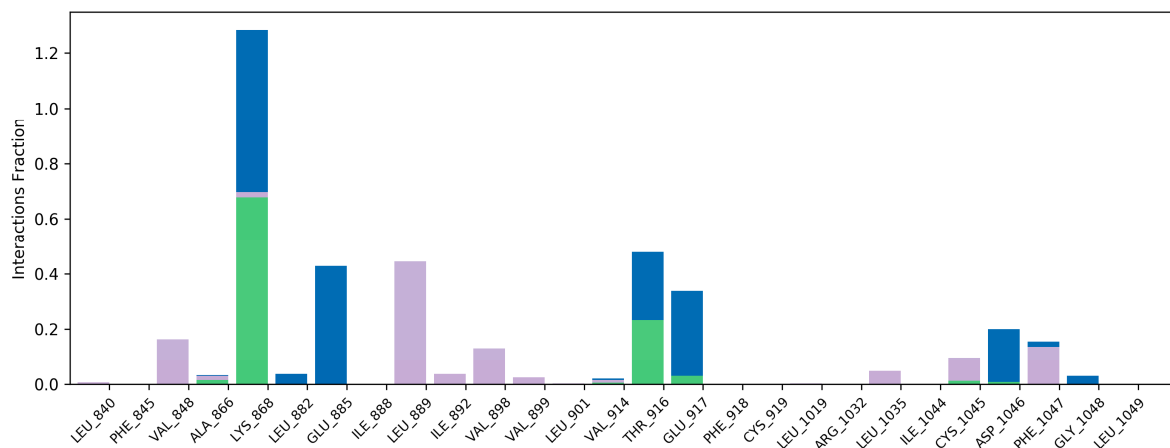
```
units angstrom  
}
```

```
set globals {  
  basis      jun-cc-pvdz  
  df_basis_scf jun-cc-pvdz-jkfit  
  df_basis_mp2 jun-cc-pvdz-ri  
  guess      sad  
  scf_type    df  
}
```

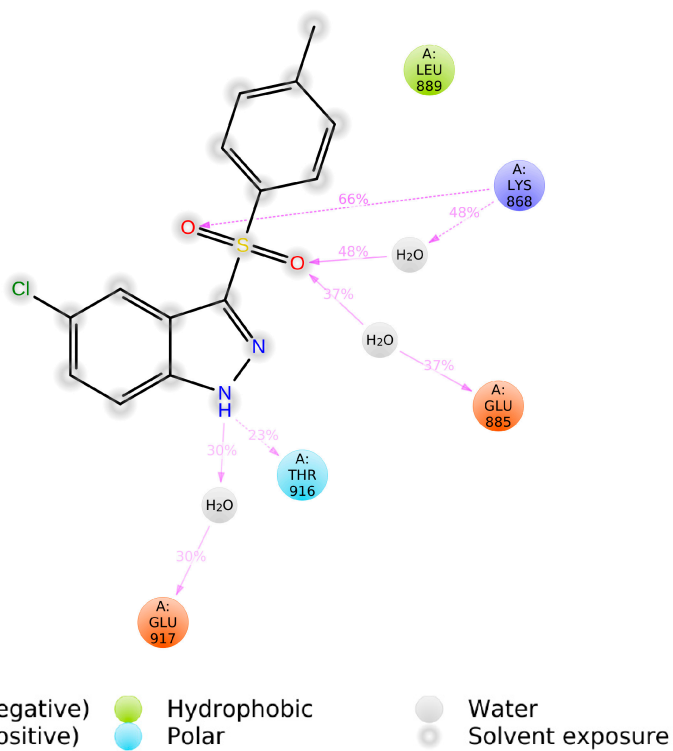
```
set sapt { print 1 }
```

```
energy('sapt0')
```

Fig. S13 The protein–ligand interactions (**a**; hydrogen bonds – green, hydrophobic – white purple, ionic – pink, water bridges – blue) and schematic of detailed ligand atom interactions (**b**) for chlorine derivative **1**.

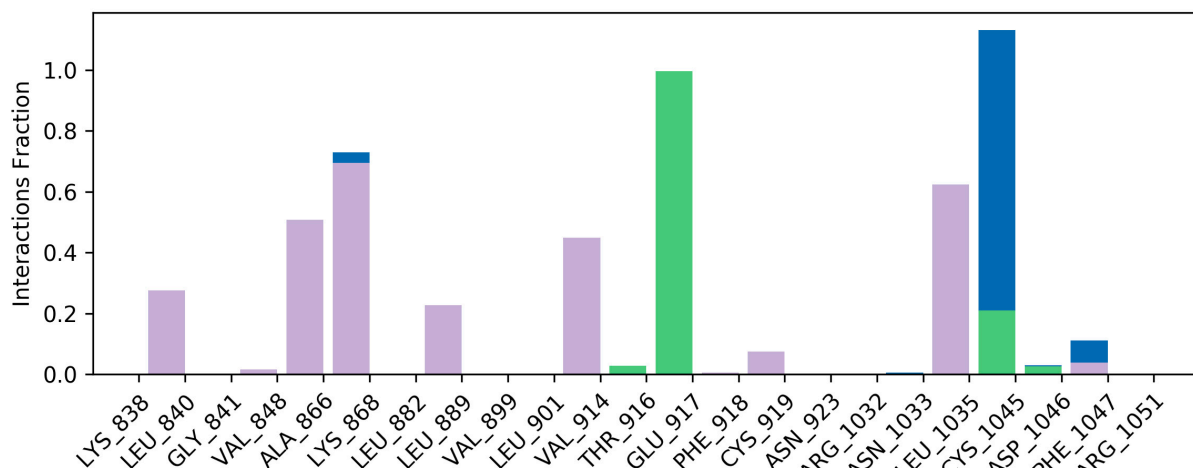


a

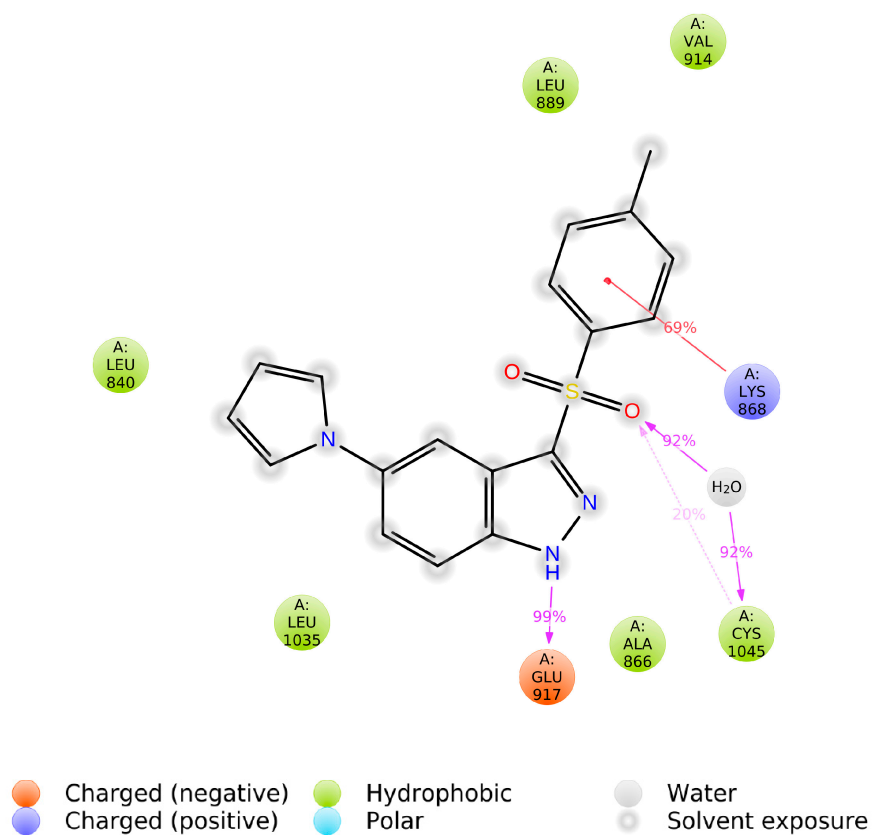


b

Fig. S14 The protein–ligand interactions (**a**; hydrogen bonds – green, hydrophobic – white purple, ionic – pink, water bridges – blue) and schematic of detailed ligand atom interactions (**b**) for pyrrole derivative **2**.

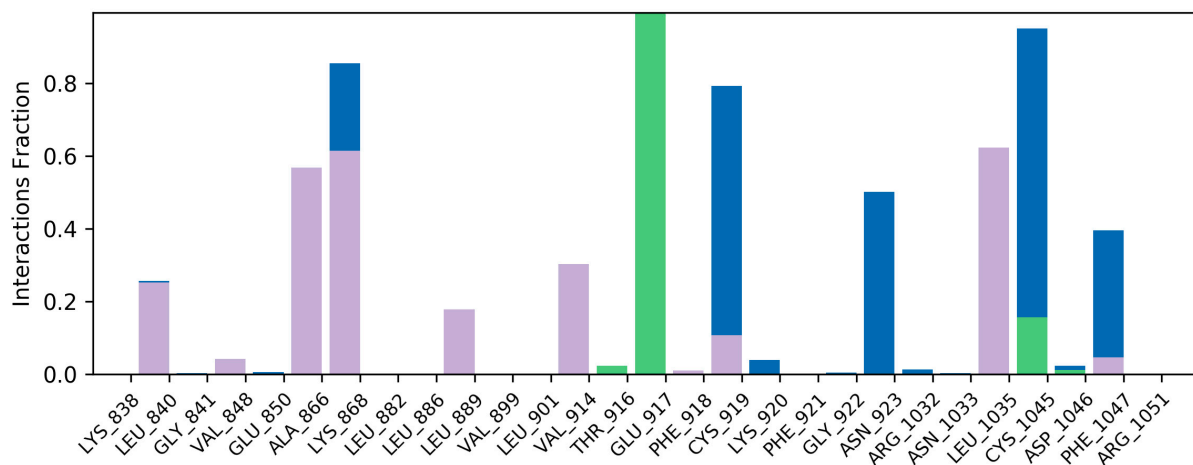


a

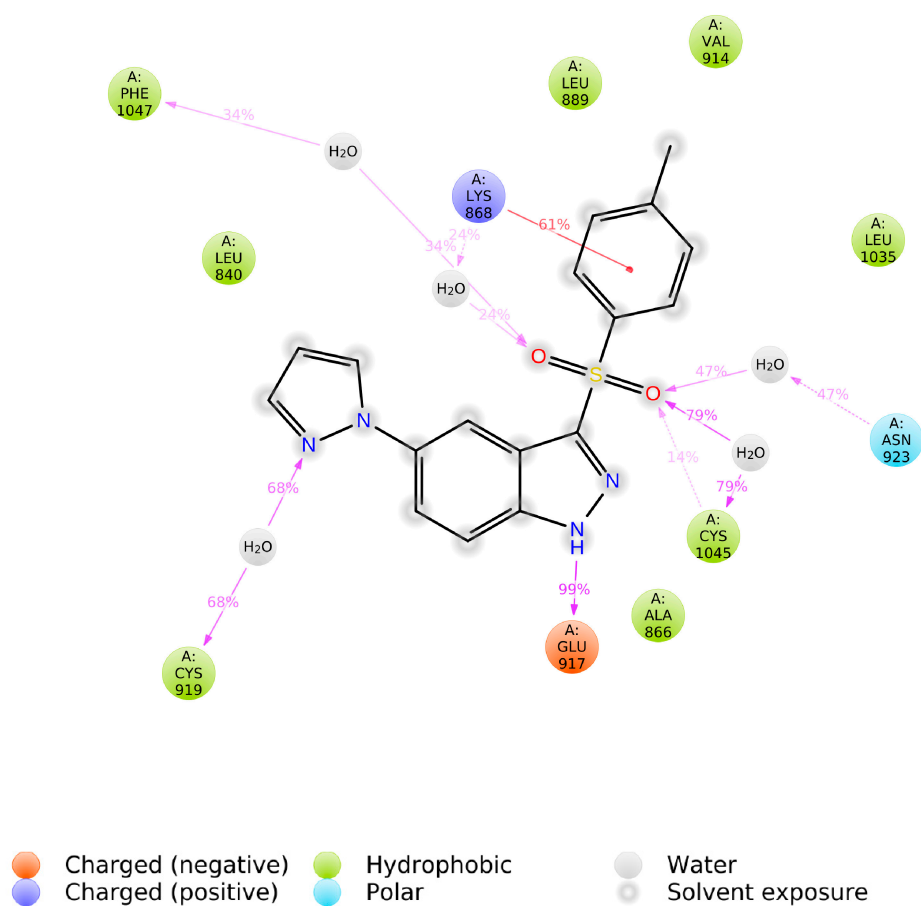


b

Fig. S15 The protein–ligand interactions (**a**; hydrogen bonds – green, hydrophobic – white purple, ionic – pink, water bridges – blue) and schematic of detailed ligand atom interactions (**b**) for pyrazole derivative **3**.

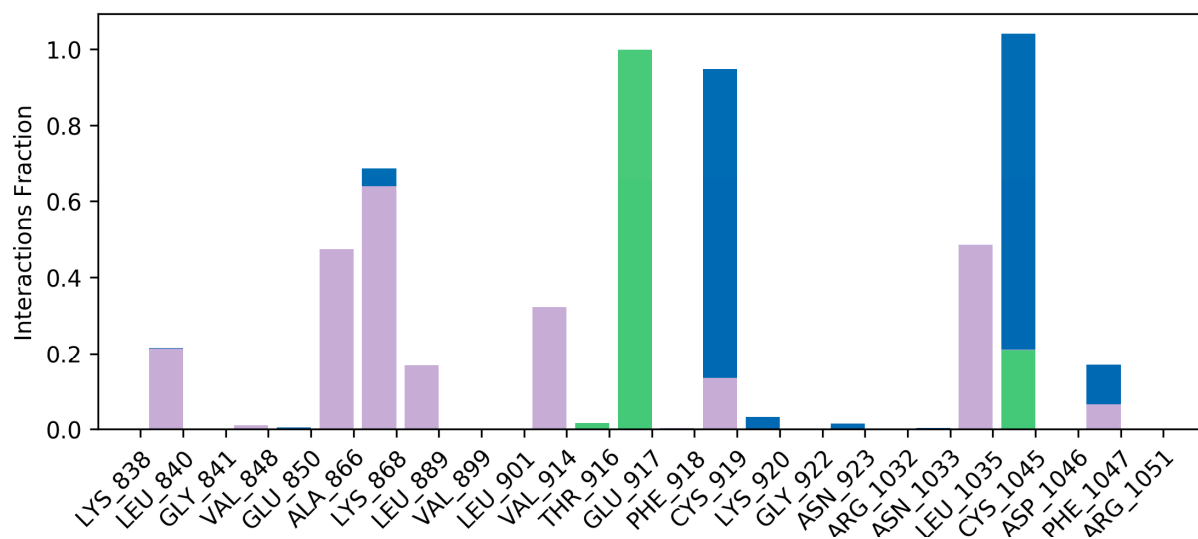


a

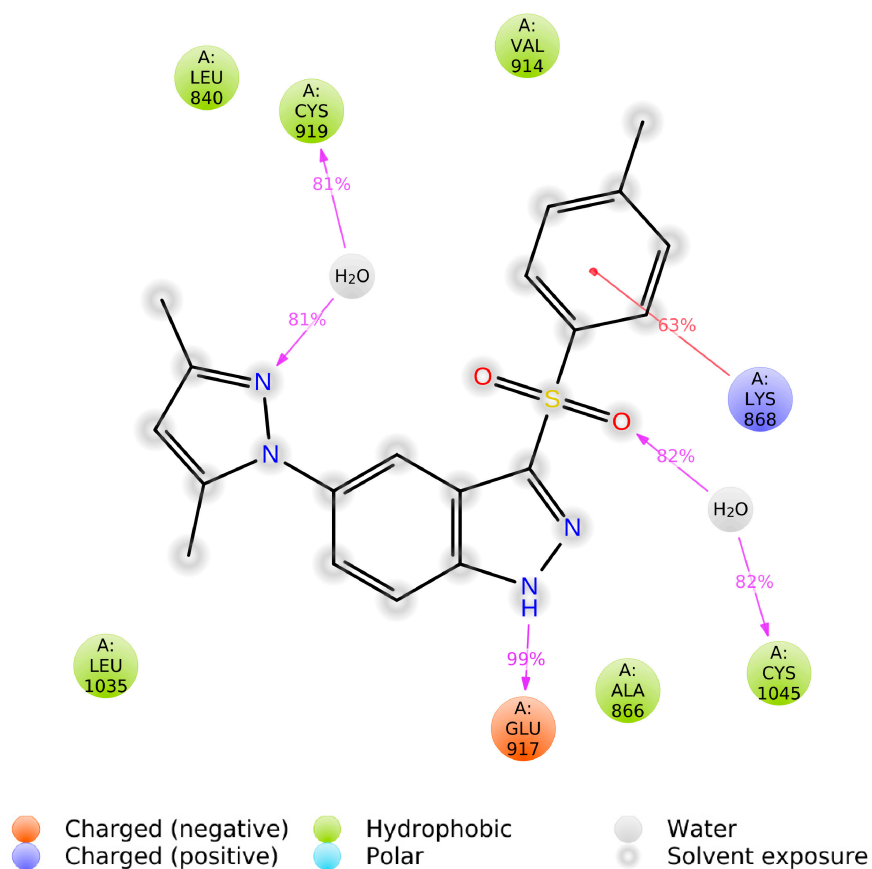


b

Fig. S16 The protein–ligand interactions (**a**; hydrogen bonds – green, hydrophobic – white purple, ionic – pink, water bridges – blue) and schematic of detailed ligand atom interactions (**b**) for dimethyl pyrazole derivative **4**.

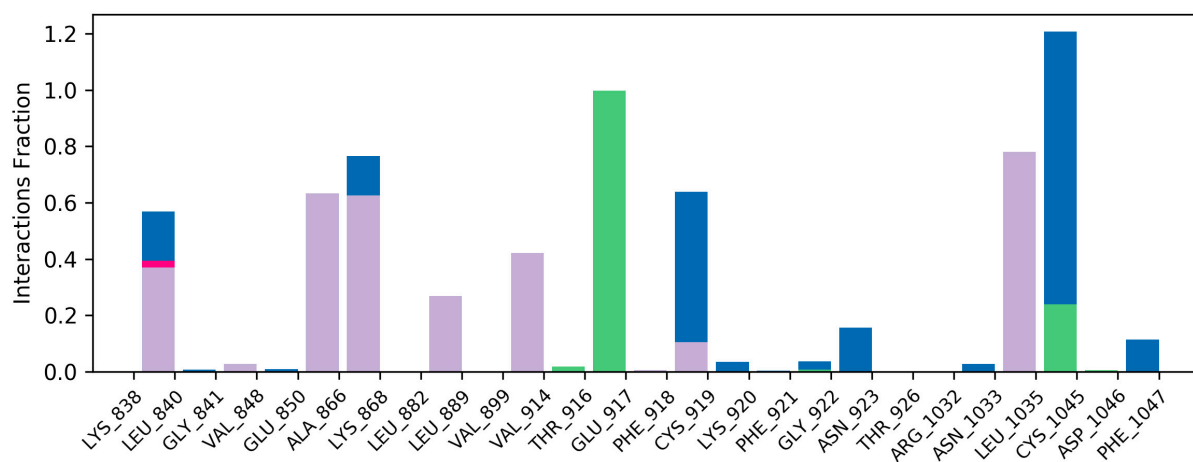


a

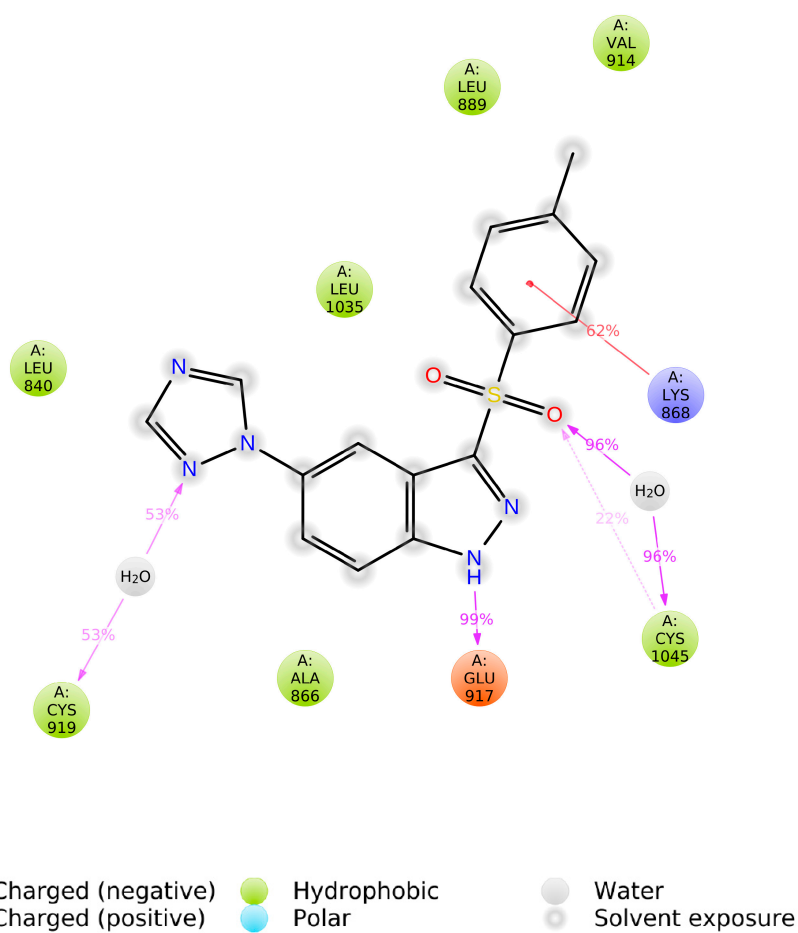


b

Fig. S17 The protein–ligand interactions (**a**; hydrogen bonds – green, hydrophobic – white purple, ionic – pink, water bridges – blue) and schematic of detailed ligand atom interactions (**b**) for triazole derivative **5**.

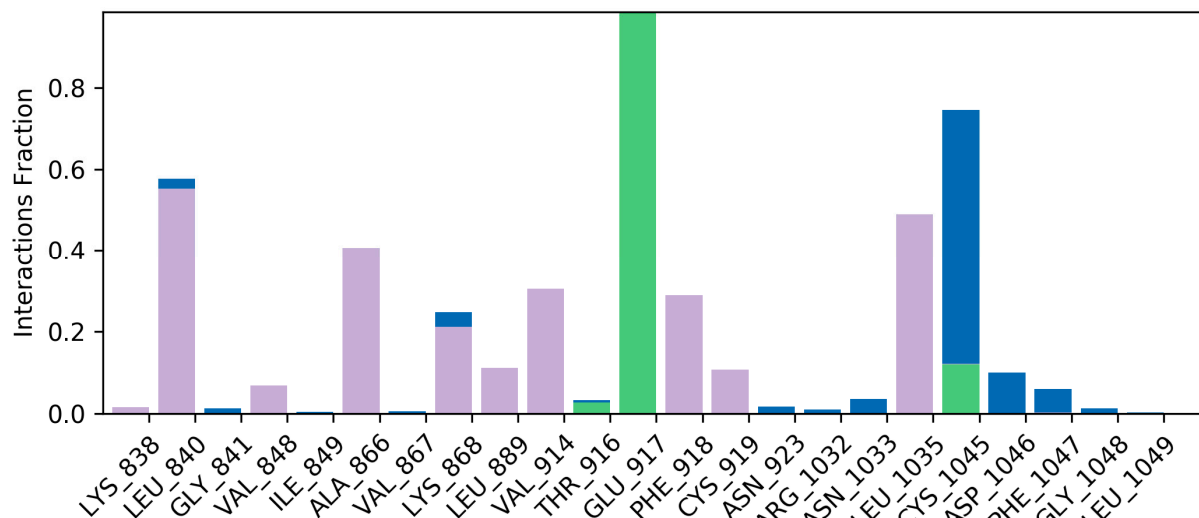


a

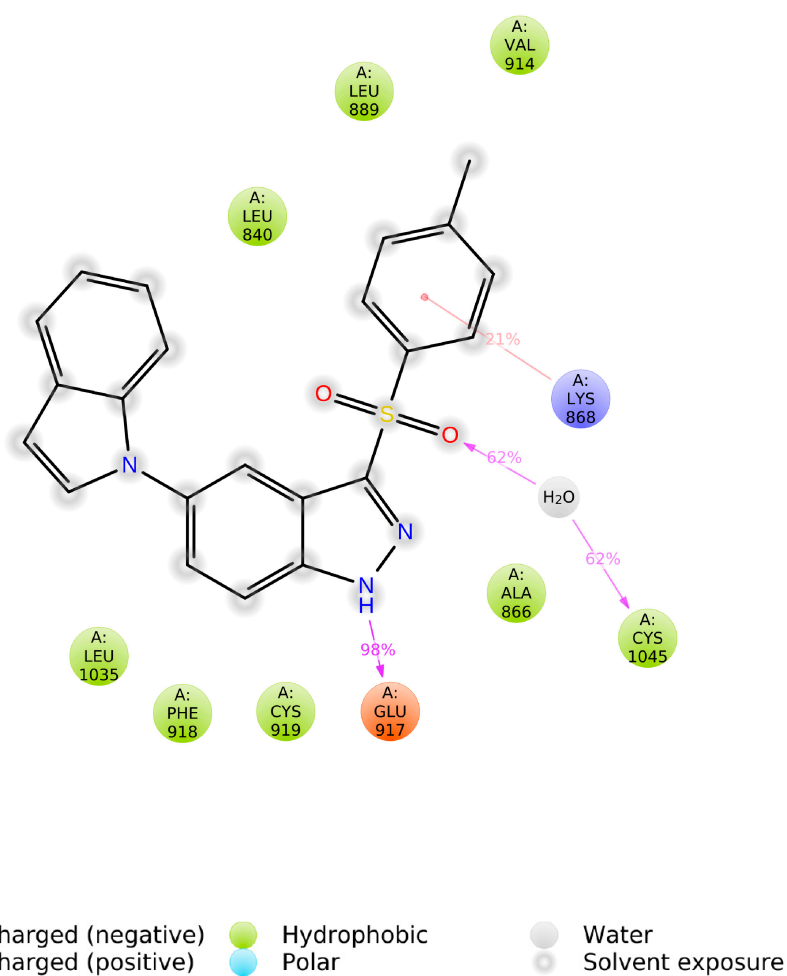


b

Fig. S18 The protein–ligand interactions (**a**; hydrogen bonds – green, hydrophobic – white purple, ionic – pink, water bridges – blue) and schematic of detailed ligand atom interactions (**b**) for indole derivative **6**.

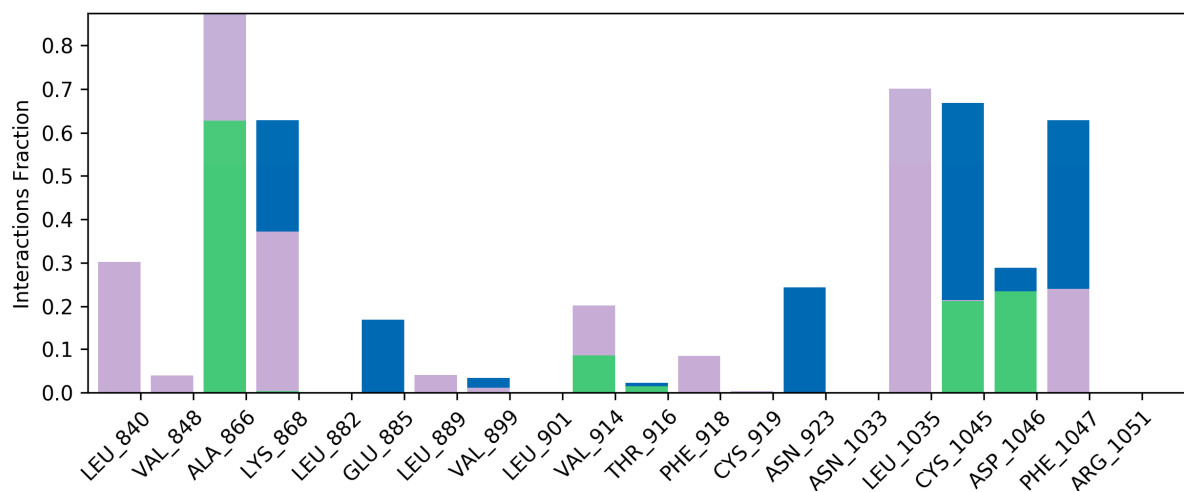


a



b

Fig. S19 The protein–ligand interactions (**a**; hydrogen bonds – green, hydrophobic – white purple, ionic – pink, water bridges – blue) and schematic of detailed ligand atom interactions (**b**) for carbazole derivative **8**.



a

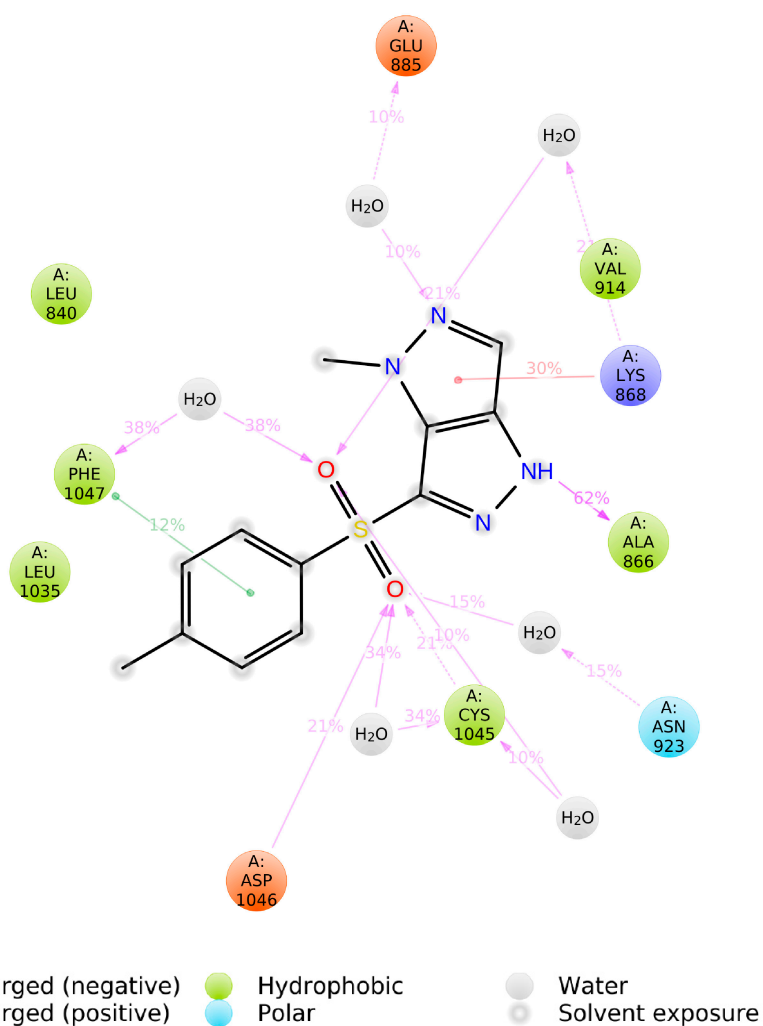


Fig. S20 Atoms numbering for RMSF calculations of the ligand **1** and the RMSF plot for **1** within ligand–protein complex during the MD productive phase calculated complex of kinase with **1**.

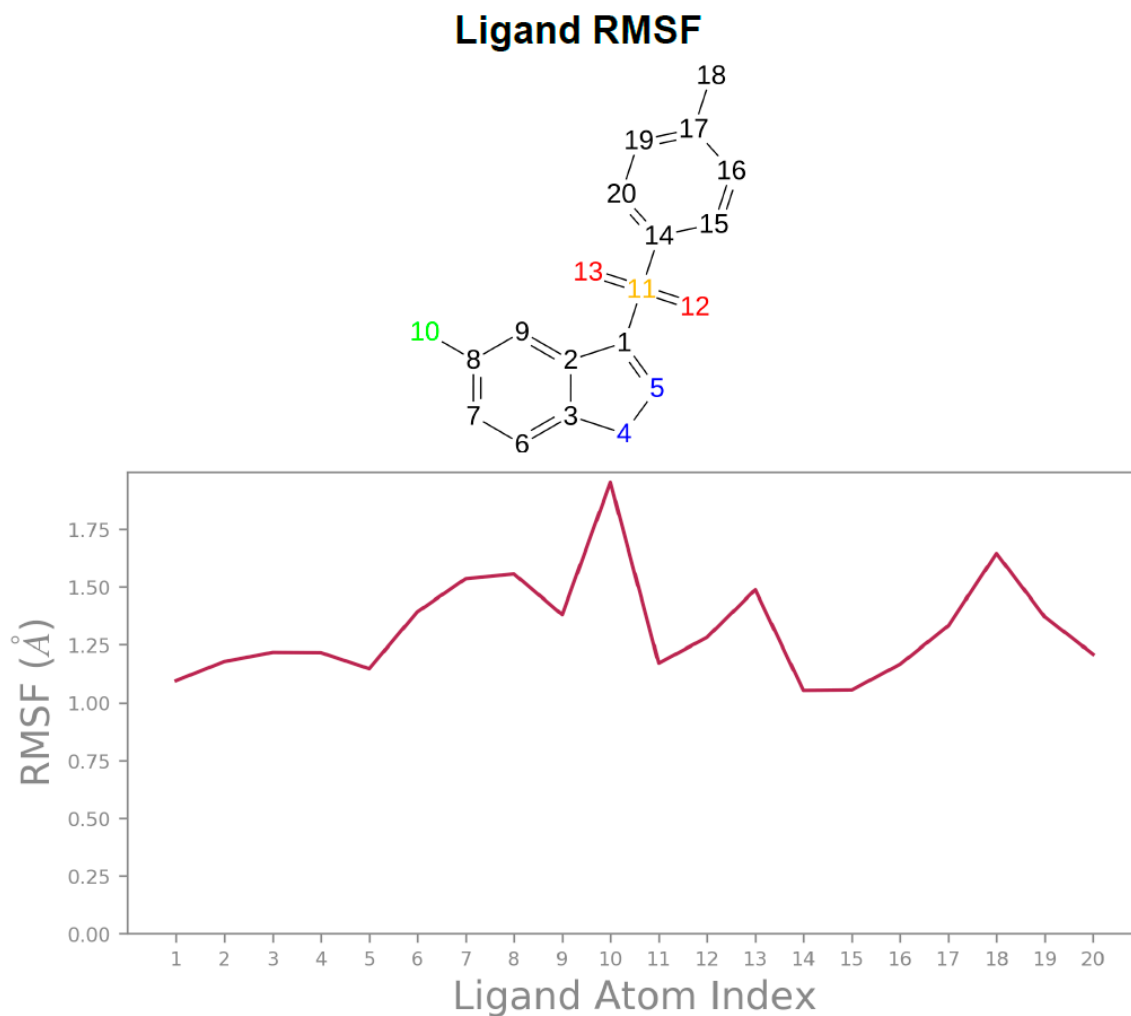


Fig. S21 The RMSD plot for the backbone within ligand-protein complex during the MD productive phase calculated complex of kinase with: **1–9** (Y-axis in Å); colors related with **Fig. 4** given in the manuscript.

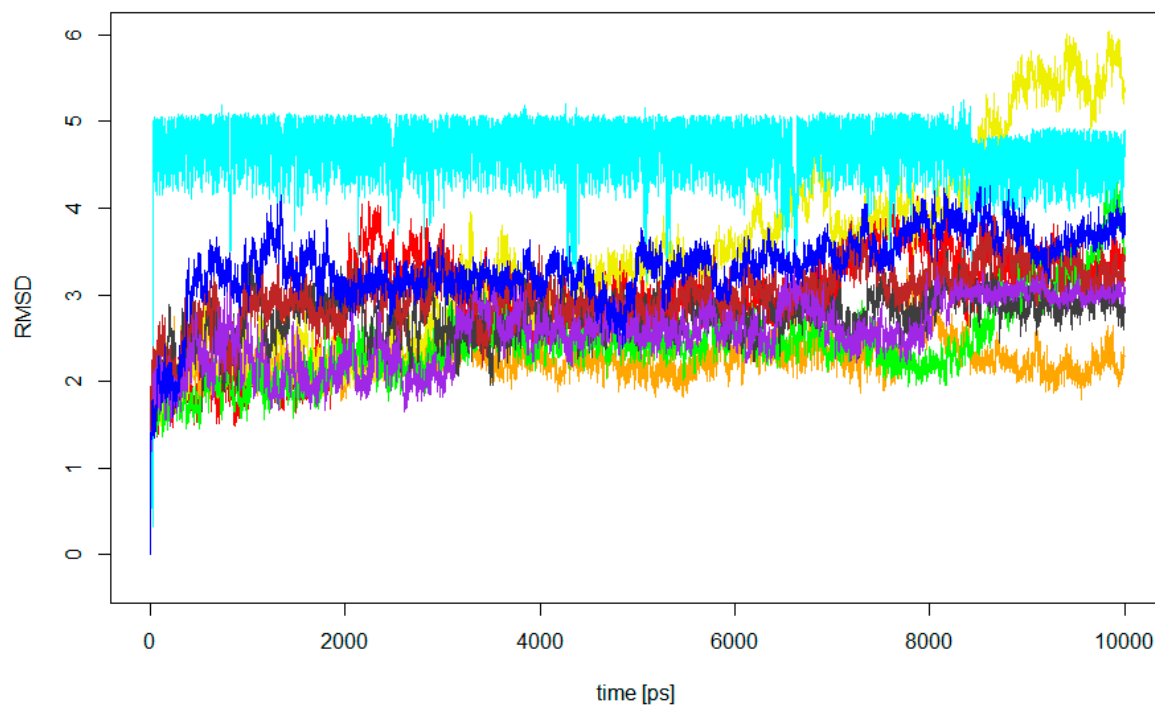


Fig. S22 Representative MD cluster for binding mode of **1** within the 1–3ewh complex (A) and its comparison with results using docking protocol (B).

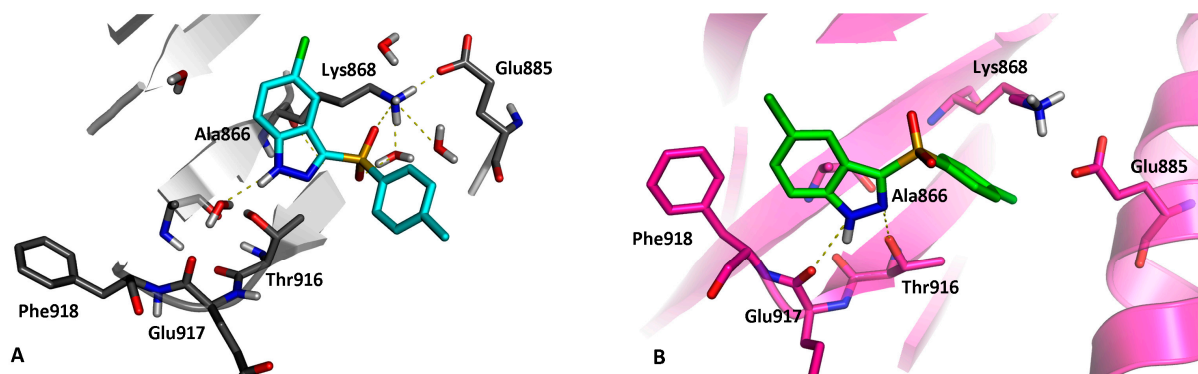


Fig. S23 Hydrogen contacts with Thr916 during the MD productive phase calculated complex of kinase with derivative **3**; red – angles [°], blue – distances [Å].

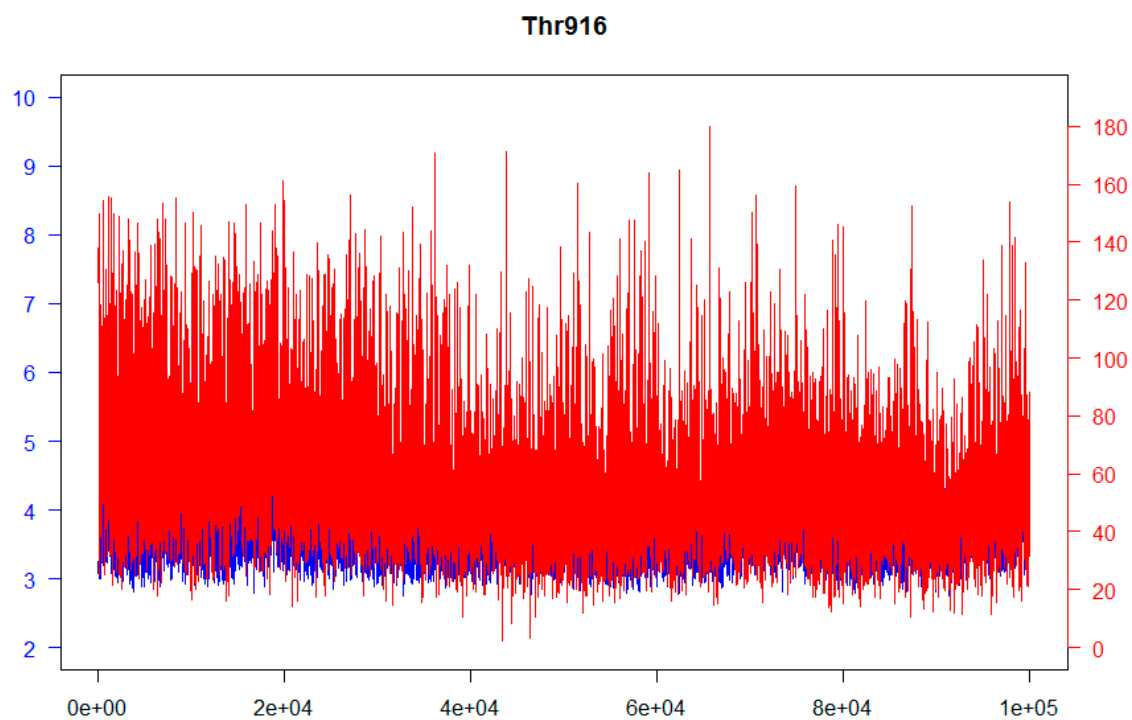


Fig. S24 Hydrogen contacts with Glu917 during the MD productive phase calculated complex of kinase with derivative **3**; red – angles [°], blue – distances [Å].

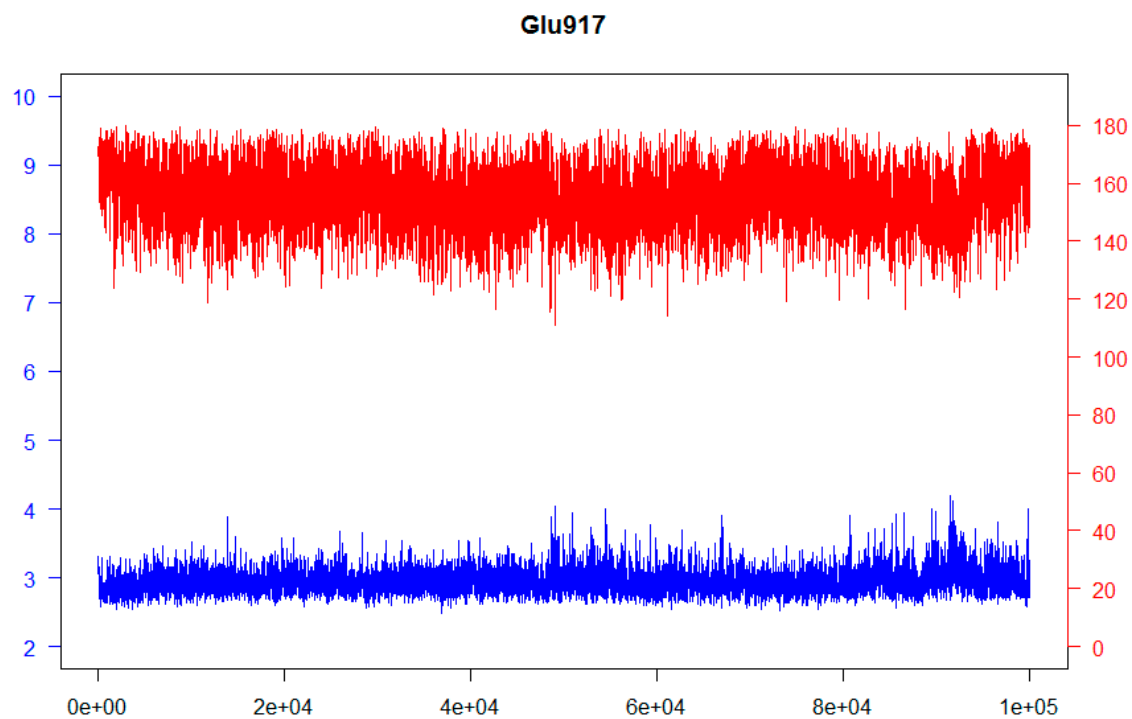


Fig. S25 Hydrogen contacts with Thr916 during the MD productive phase calculated complex of kinase with derivative **6**; red – angles [°], blue – distances [Å].

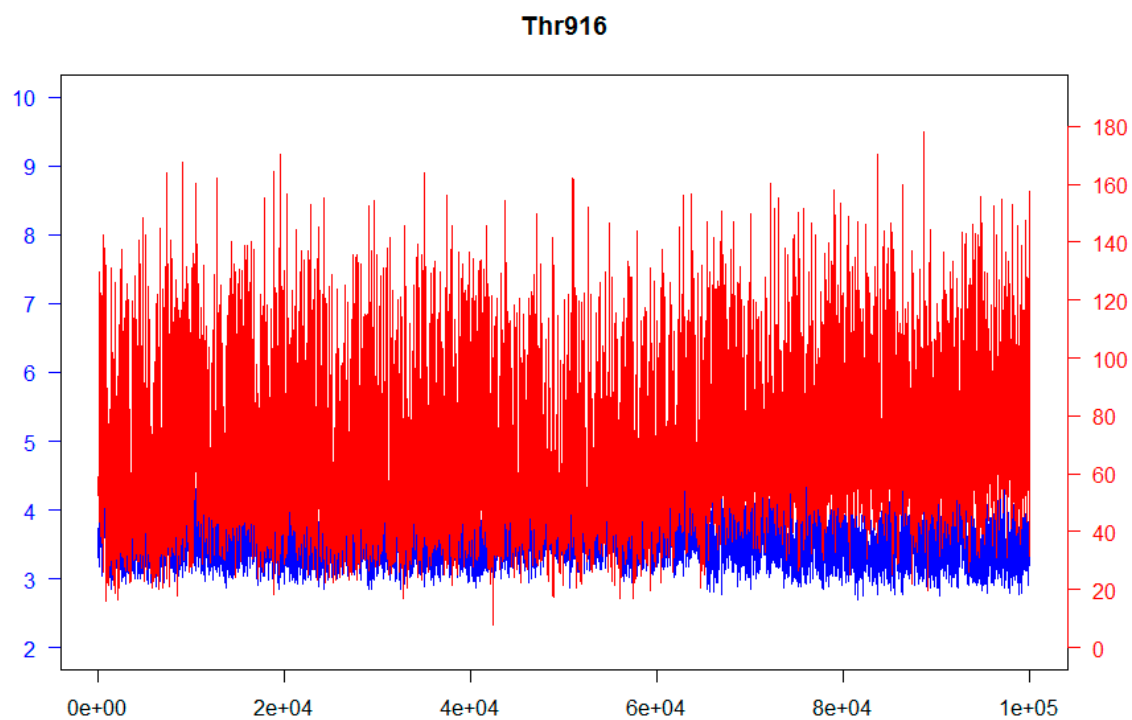


Fig. S26 Hydrogen contacts with Glu917 during the MD productive phase calculated complex of kinase with derivative **6**; red – angles [°], blue – distances [Å].

