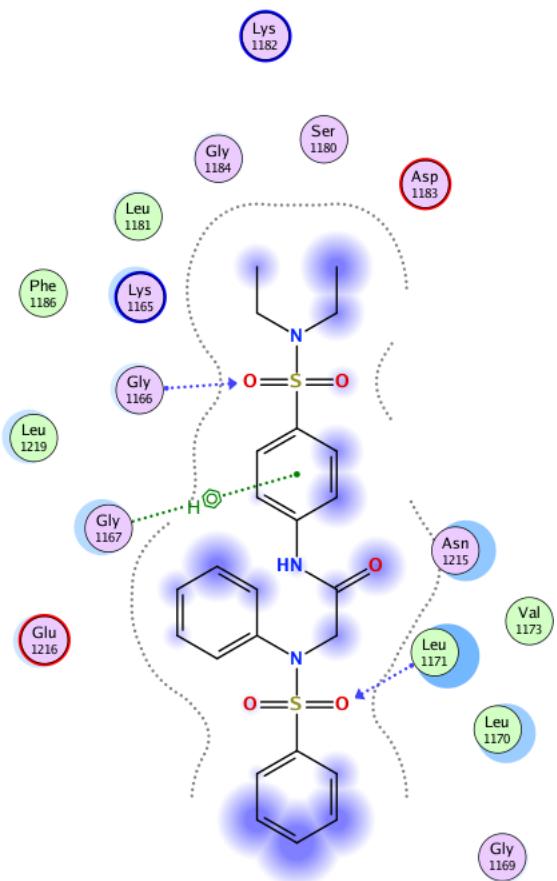


**Table S1.** Ten candidate compounds identified by *in silico* SBDS.

| compound | ChemBridge IDs | IUPAC name   | M.W. | cLogP | tPSA | GOLD score |
|----------|----------------|--|------|-------|------|------------|
| 1        | 7389795        | 4-methoxy-N-(4-[(2-phenylethyl)amino]sulfonyl]phenyl)benzenesulfonamide  | 447  | 3.86  | 102  | 75.9       |
| 2        | 7788851        | <i>N</i> <sup>1</sup> -{4-[(diethylamino)sulfonyl]phenyl}- <i>N</i> <sup>2</sup> -phenyl- <i>N</i> <sup>2</sup> -(phenylsulfonyl)glycinamide               | 502  | 4.46  | 104  | 75.8       |
| 3        | 7780058        | 2-({3-[(2-ethoxybenzoyl)amino]benzoyl}amino)benzoic acid   | 404  | 4.70  | 105  | 72.4       |
| 4        | 7972614        | <i>N,N'</i> -bis(4-methoxybenzyl)-1-[(4-methylphenyl)sulfonyl]-1 <i>H</i> -1,2,4-triazole-3,5-diamine  | 494  | 3.18  | 107  | 71.1       |
| 5        | 6824862        | <i>N</i> -(1,5-dimethyl-3-oxo-2-phenyl-2,3-dihydro-1 <i>H</i> -pyrazol-4-yl)-2-[(5-(phenoxy)methyl)-4-phenyl-4 <i>H</i> -1,2,4-triazol-3-yl]thio)acetamide | 527  | 1.42  | 96   | 70.8       |
| 6        | 7968391        | <i>N</i> -1,3-benzothiazol-2-yl-2-((5-[3-(4-methoxy-2-methylphenyl)propyl]-4-methyl-4 <i>H</i> -1,2,4-triazol-3-yl)thio)acetamide                          | 468  | 4.31  | 82   | 70.4       |
| 7        | 6945187        | 3-[4-(4-bromophenyl)-1,3-thiazol-2-yl]-7-hydroxy-8-[[4-(2-hydroxyethyl)-1-piperazinyl]methyl]-2 <i>H</i> -chromen-2-one                                    | 542  | 4.92  | 90   | 68.8       |
| 8        | 7355746        | <i>N</i> -(2,4-dichlorobenzyl)- <i>N</i> -(4-methylphenyl)sulfonyl)glycine   | 388  | 4.86  | 75   | 68.3       |
| 9        | 7917692        | <i>N</i> -[(1-ethyl-2-oxo-1,2-dihydrobenzo[cd]indol-6-yl)sulfonyl]tryptophan   | 464  | 3.26  | 120  | 68.1       |
| 10       | 6082385        | 4-[4-[(phenoxyacetyl)amino]phenoxy]phthalic acid   | 407  | 3.58  | 122  | 68.0       |



**Figure S1.** Ligand Interaction (LI) between IR $\beta$  and compound 2. The data shows predicted interaction at 50 ns in the MD simulation. The dotted blue allows and green line represent hydrogen bonds and cation- $\pi$  interaction, respectively. The blue crowds on compound 2 indicate hydrophobic interactions with the binding pocket.