

Discovery of Novel Hsp90 C-Terminal Inhibitors Using 3D-Pharmacophores Derived from Molecular Dynamics Simulations

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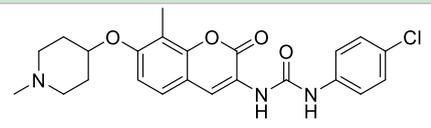
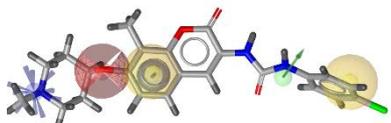
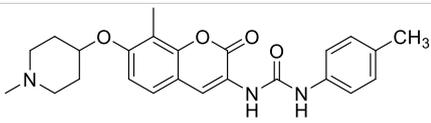
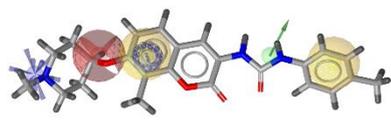
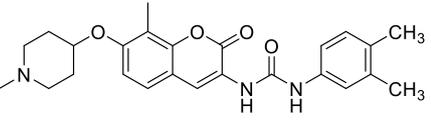
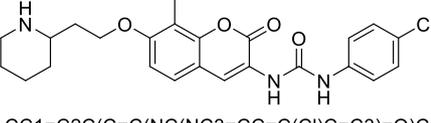
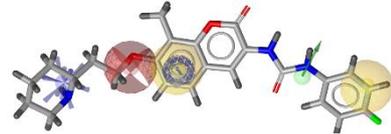
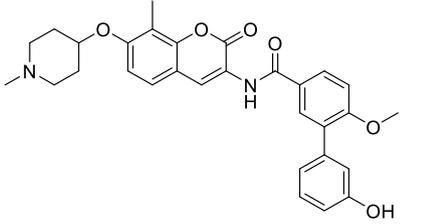
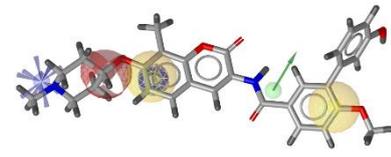
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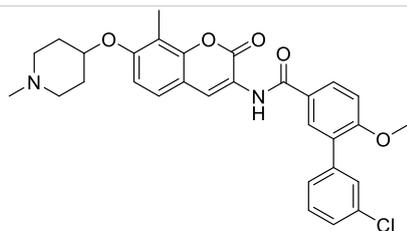
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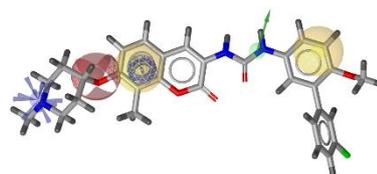
Table S1. Structures and antiproliferative activities in SKBr3 cell line of Hsp90 CTD inhibitors **2** and **S1–S12** used as a training set for ligand-based (LB) pharmacophore model creation. Each ligand is shown aligned to the resulting LB model (exclusion volumes are not displayed).

Cpd	Structure	SKBr3 IC ₅₀ (μM)	Alignment with the 3D Ligand-Based Pharmacophore Model
2	 <chem>CC1=C2C(C=C(NC(NC3=CC=C(Cl)C=C3)=O)C(O2)=O)=CC=C1OC4CCN(C)CC4</chem>	0.17 [1]	
S1	 <chem>CC1=C2C(C=C(NC(NC3=CC=C(C)C=C3)=O)C(O2)=O)=CC=C1OC4CCN(C)CC4</chem>	0.27 [1]	
S2	 <chem>CC1=C2C(C=C(NC(NC3=CC(C)=C(C)C=C3)=O)C(O2)=O)=CC=C1OC4CCN(C)CC4</chem>	0.31 [1]	
S3	 <chem>CC1=C2C(C=C(NC(NC3=CC=C(Cl)C=C3)=O)C(O2)=O)=CC=C1OCCC4NCCCC4</chem>	0.21 [1]	
S4	 <chem>CC1=C2C(C=C(NC(NC3=CC(C4=CC=CC(O)=C4)=C(OC)C=C3)=O)C(O2)=O)=CC=C1OC5CCN(C)CC5</chem>	0.31 [1]	

S5

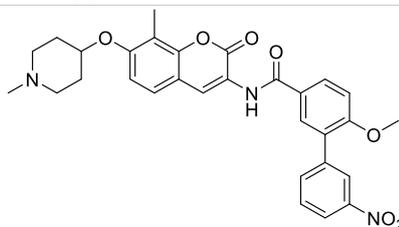


0.13 [1]

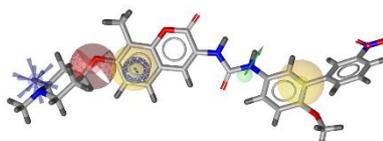


CC1=C2C(C=C(NC(C3=CC(C4=CC=CC(Cl)=C4)=C(OC)C=C3)=O)C(O2)=O)=CC=C1OC5CCN(C)CC5

S6

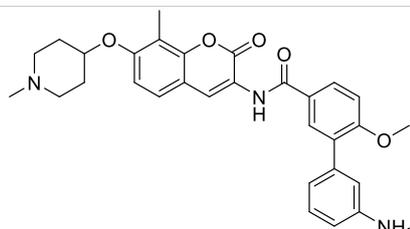


0.16 [1]

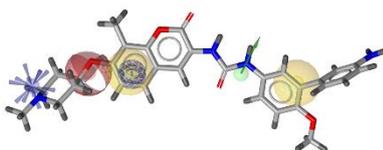


CC1=C2C(C=C(NC(C3=CC(C4=CC=CC([N+](O-))=O)=C4)=C(OC)C=C3)=O)C(O2)=O)=CC=C1OC5CCN(C)CC5

S7

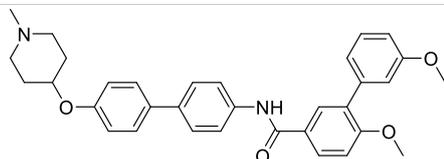


0.36 [1]



CC1=C2C(C=C(NC(C3=CC(C4=CC=CC(N)=C4)=C(OC)C=C3)=O)C(O2)=O)=CC=C1OC5CCN(C)CC5

S8

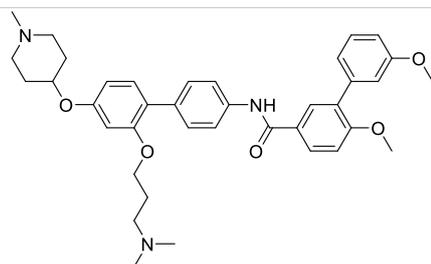


0.47 [2]

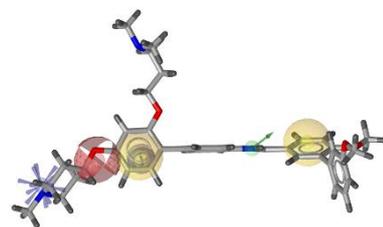


CN(CC1)CCC1OC(C=C2)=CC=C2C3=CC=C(NC(C4=CC(C5=CC=CC(OC)=C5)=C(OC)C=C4)=O)C=C3

S9

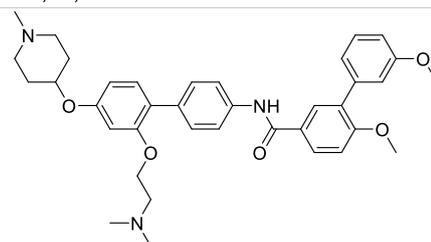


0.39 [2]

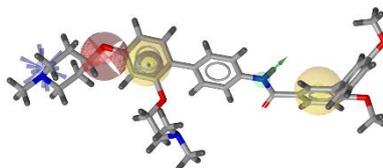


CN(CC1)CCC1OC(C=C2)=CC(OCCN(C)C)=C2C3=CC=C(NC(C4=CC(C5=CC=CC(OC)=C5)=C(OC)C=C4)=O)C=C3

S10



0.34 [2]



CN(CC1)CCC1OC(C=C2)=CC(OCCN(C)C)=C2C3=CC=C(NC(C4=CC(C5=CC=CC(OC)=C5)=C(OC)C=C4)=O)C=C3

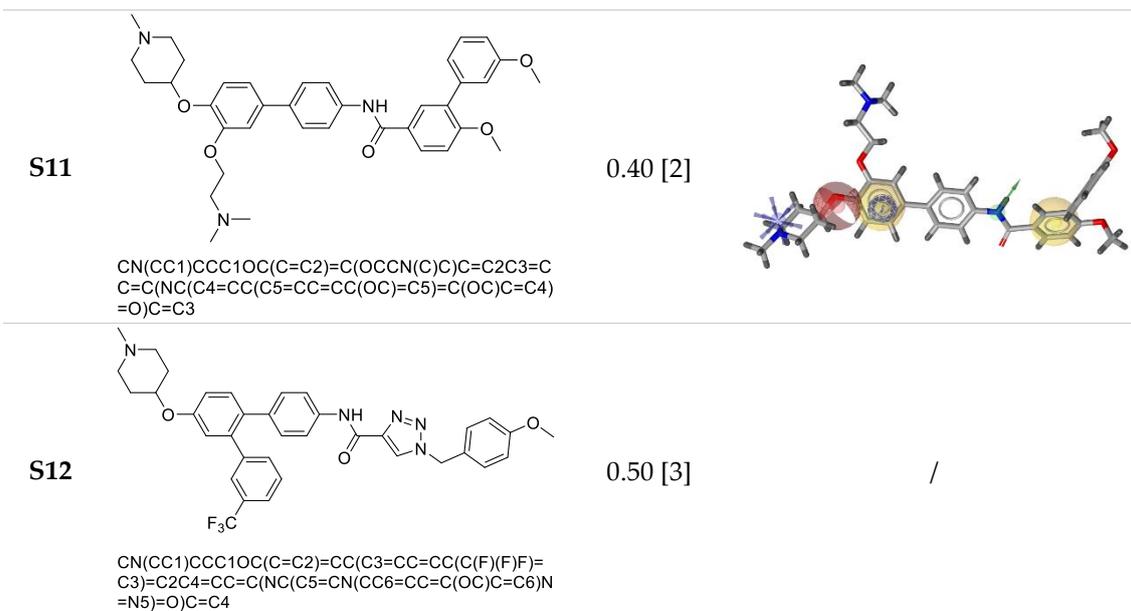


Table S2. Cell viability of Hep G2 and MCF-7 cells in MTS assay after treatment with compounds 5–12. Data are means \pm SD of three independent experiments performed in triplicates.

Compound	Hep G2 % Viability	MCF-7 % Viability
5	86.8 \pm 7.6	93.5 \pm 4.3
6	67.0 \pm 3.7	71.0 \pm 9.5
7	100.2 \pm 6.0	99.5 \pm 5.8
8	67.2 \pm 3.0	65.2 \pm 20.8
9	86.2 \pm 12.8	-0.53 \pm 1.5
10	97.7 \pm 7.5	116.5 \pm 41.4
11	1.38 \pm 0.1	31.4 \pm 11.9
12	92.6 \pm 4.3	93.8 \pm 1.4

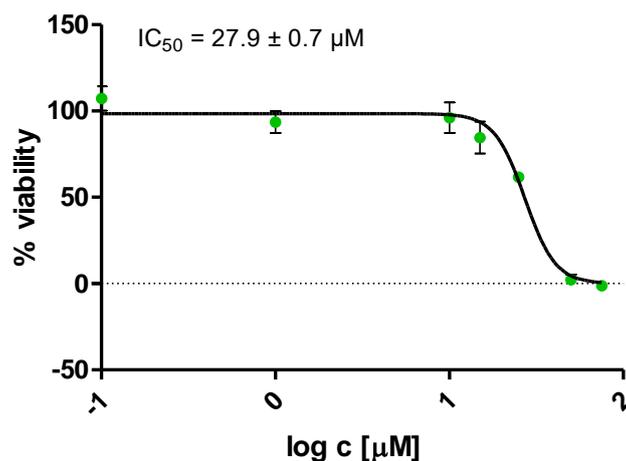


Figure S1. Dose-response curve for compound 11 in MTS assay in Hep G2 cell line, shown for an independent measurement in triplicate. The IC₅₀ value (mean \pm SD) is a result of three independent measurements.

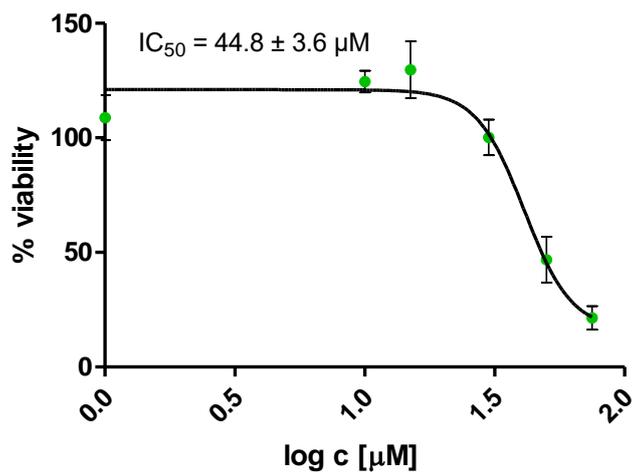


Figure S2. Dose-response curve for compound **11** in MTS assay in MCF-7 cell line, shown for an independent measurement in triplicate. The IC₅₀ value (mean ± SD) is a result of three independent measurements.

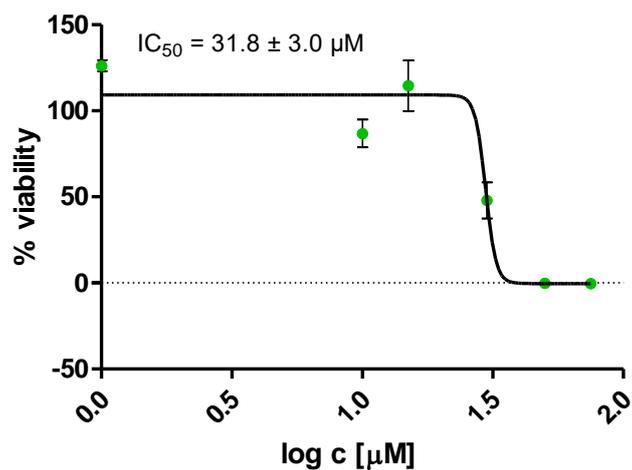


Figure S3. Dose-response curve for compound **9** in MTS assay in MCF-7 cell line, shown for an independent measurement in triplicate. The IC₅₀ value (mean ± SD) is a result of three independent measurements.

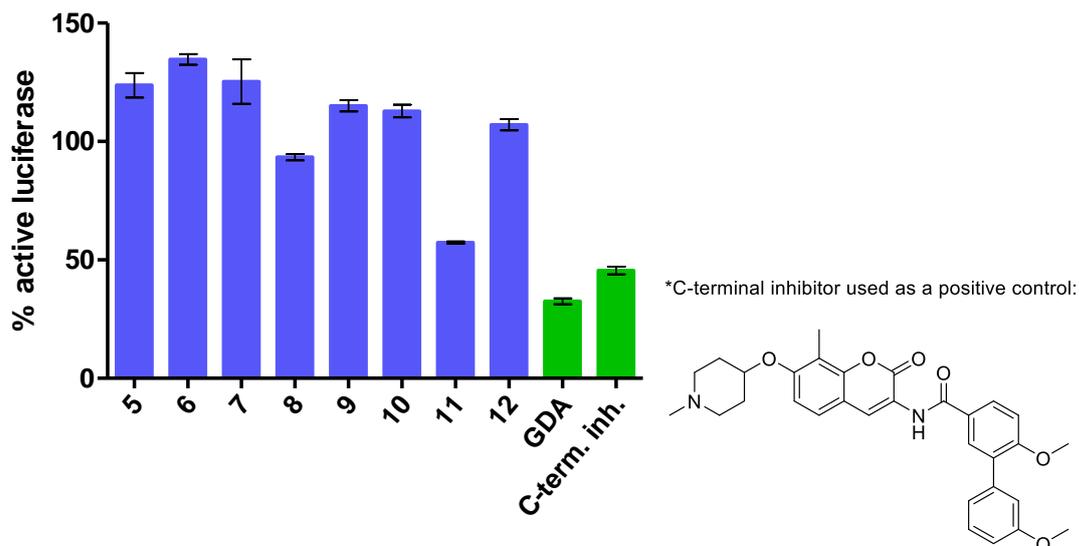


Figure S4. Luciferase refolding activity of Hsp90 in PC3 MM2 cells after treatment with compounds 5–12, C-terminal inhibitor* and geldanamycin (GDA) at 50 μM concentration. Data are means \pm SD of three independent experiments performed in triplicates.

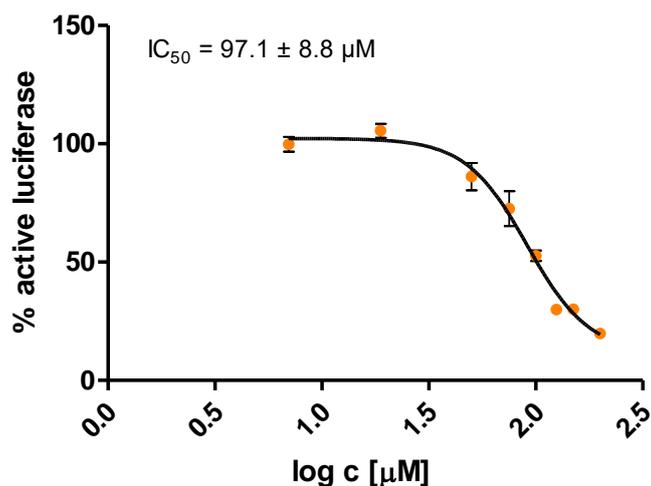


Figure S5. Dose-response curve for compound 11 in luciferase refolding assay on PC3 MM2 cell line, shown for an independent measurement in triplicate. The IC_{50} value (mean \pm SD) is a result of three independent measurements.

References

1. Moroni, E.; Zhao, H.; Blagg, B.S.J.; Colombo, G. Exploiting Conformational Dynamics in Drug Discovery: Design of C-Terminal Inhibitors of Hsp90 with Improved Activities. *J. Chem. Inf. Mode.* **2014**, *54*, 195–208, doi:10.1021/ci4005767.
2. Garg, G.; Zhao, H.; Blagg, B.S.J. Design, Synthesis and Biological Evaluation of Alkylamino Biphenylamides as Hsp90 C-Terminal Inhibitors. *Bioorg. Med. Chem.* **2017**, *25*, 451–457, doi:10.1016/j.bmc.2016.11.030.
3. Davis, R.E.; Zhang, Z.; Blagg, B.S.J. A Scaffold Merging Approach to Hsp90 C-Terminal Inhibition: Synthesis and Evaluation of a Chimeric Library. *Med. Chem. Commun.* **2017**, *8*, 593–598, doi:10.1039/C6MD00377J.