

Marine-Derived Compounds as Potential Inhibitors of Hsp90 for Anticancer and Antimicrobial Drug Development: A Comprehensive In Silico Study

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Table S1. Properties and Definitions of Selected Marine-Derived Compounds as Potent Inhibitors of Hsp90 Protein.

CMNPD	22591	9335	10015	360799	15115	20988
Structure						
Canonical SMILES	c1ccc(ccc1O)C2=C([C@H](OC)c3ccc(O)cc3)C(=O)NC(=O)C(c2c[nH]c(c4ccc3O)c2)C(=O)c4ccc(c4)N	C1(=O)NC(=O)C(c2c[nH]c(c4ccc3O)c2)C(=O)c3ccc(O)cc3	C1=CC=C2C(=C1)C(=CN2)C(=O)C3=N	c12c(ccc1)[nH]cc2[C@H]3CNC(=O)C(c4c[nH]c(c4ccc5)c5)=N3	CN(CCC1=CNC2=C1C(=O)C3=C(C(=O)NCC)C=C2C(=O)C(C=C13)C=C(C=C3)O	COCl=C2C(=C(C=C1C3=CC=C(C=C3)O)O)C4=CC(=C(C=C4O2)O)O
PubChem CID	162904707	398493	136691197	16104844	11717517	53262759
Molecular formula	C ₂₀ H ₁₅ NO ₆	C ₁₅ H ₁₀ N ₄ O ₂	C ₂₁ H ₁₄ N ₄ O ₂	C ₂₀ H ₁₆ N ₄ O	C ₂₀ H ₁₇ N ₃ O ₄	C ₁₉ H ₁₄ O ₆
IUPAC name	methyl (2R)-2-ethoxy-3,4-bis(4-hydroxyphenyl)-5-oxo-1H-pyrrole-2-carboxylate	3-(1H-imidazol-5-yl)-4-(1H-indol-3-yl)pyrrole-2,5-dione	[4-hydroxy-5-(indol-3-ylidenemethyl)-1H-imidazol-2-yl]-[1H-indol-3-yl]methanone	((3R)-3,5-bis(1H-indol-3-yl)-2,3-dihydro-1H-pyrazin-6-one	N-[2-[3-(4-hydroxyphenyl)-4,8-dioxo-1,7-dihydropyrrrolo[3,2-f]indol-5-yl]ethyl]-N-methylformamide	3-(4-hydroxyphenyl)-4-methoxydibenzofuran-1,7,8-triol
Synonym	Ianthellidone B	Didemnimide A	Rhopaladins D	(R)-6'-debromohamacanthin B	Zyzzyanone D	3''-deoxy-6'-O-desmethylcandidusin B
Resource	Demospongiae	Didemnum granulatum	tunicate Rhopalaea sp	Spongisorites	Zyzya fuliginosa	Penicillium chermesinum

Classy fire subclass	Stilbenes	Indoles	Indoles	Indoles	Pyrroloindoles	Phenylbenzofurans
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Table S2. Result Of Enrichment Study.

Pharmacophore model	AUC	RIE	ROC	BedROC160.9	Total active	Number of ranked actives
DDRRR	0.84	7.18	0.78	0.58	35	29

Table S3. Organ and End Points Toxicity for Top Leads Compound.

		22591		9335		10015		360799		15115		20988		MEY		
		Probability	Prediction													
Organ toxicity	Hepatotoxicity	dili	0.50	inactive	0.51	inactive	0.50	inactive	0.72	inactive	0.62	inactive	0.67	inactive	0.55	active
	Carcinogenicity	carcino	0.58	inactive	0.51	active	0.55	active	0.56	inactive	0.59	active	0.62	active	0.71	inactive
	Immunotoxicity	immuno	0.95	inactive	0.73	inactive	0.7	inactive	0.98	inactive	0.79	inactive	0.8	inactive	0.54	inactive
	Mutagenicity	mutagen	0.67	inactive	0.61	inactive	0.6	active	0.67	inactive	0.54	inactive	0.55	inactive	0.78	inactive
	Cytotoxicity	cyto	0.66	inactive	0.7	inactive	0.74	inactive	0.77	inactive	0.55	inactive	0.95	inactive	0.64	inactive

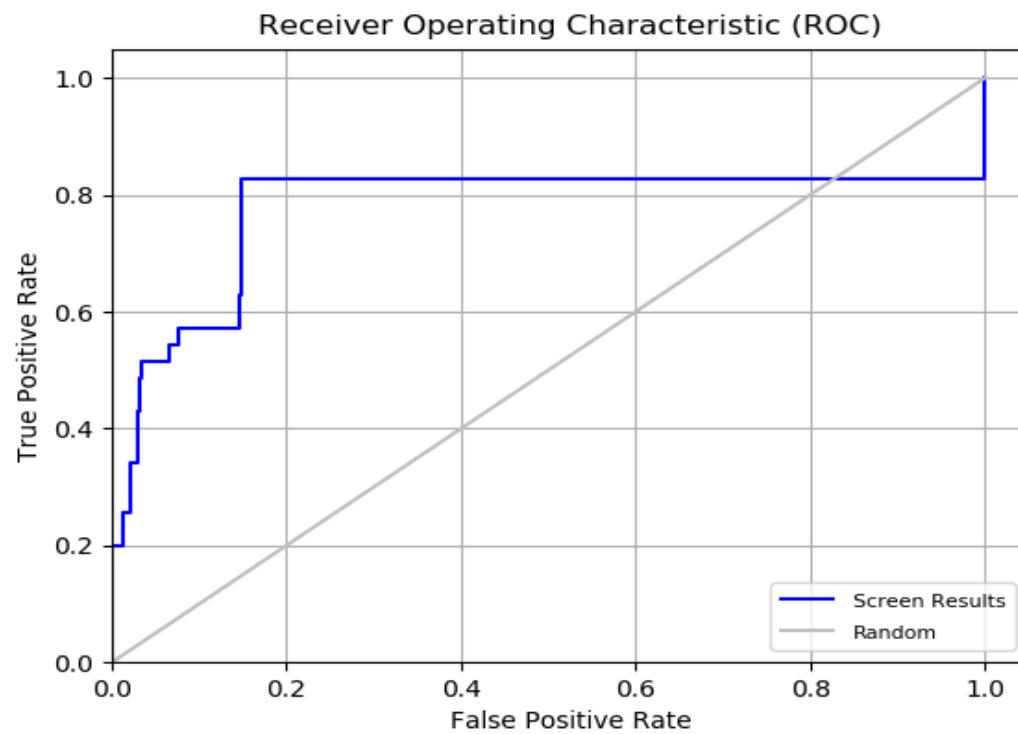
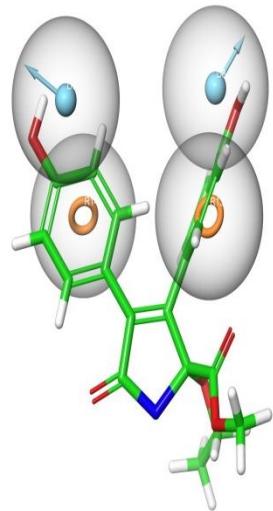
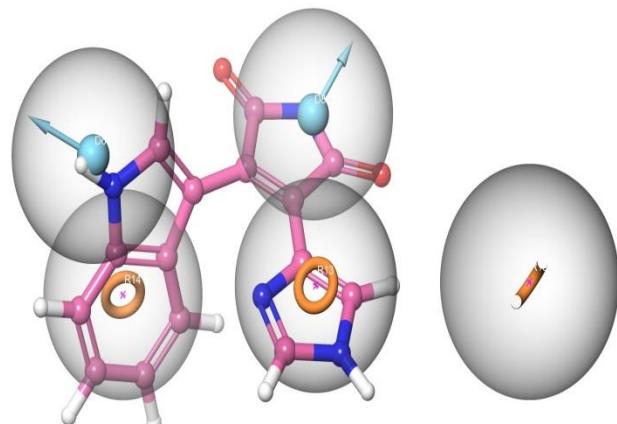


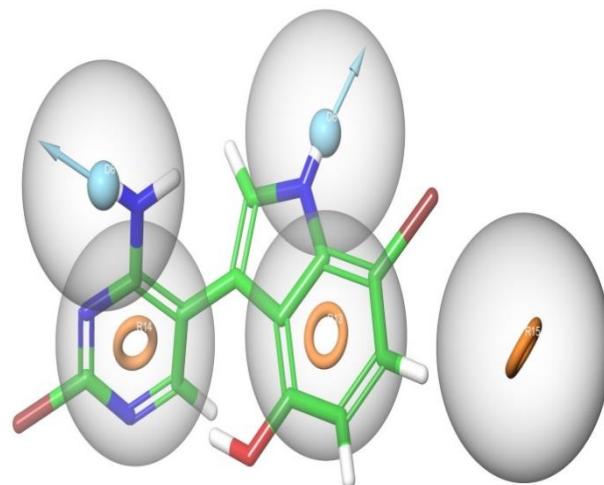
Figure S1. ROC plot of the pharmacophore model.



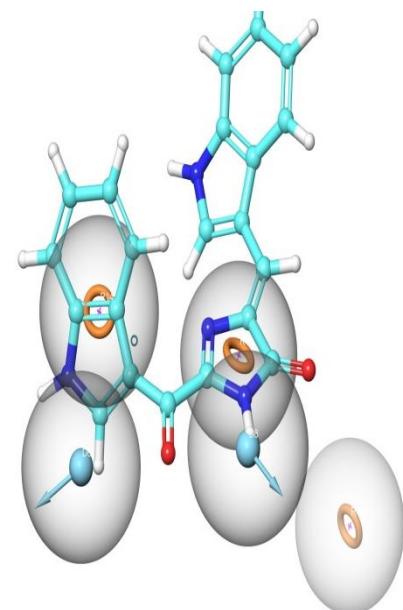
22591



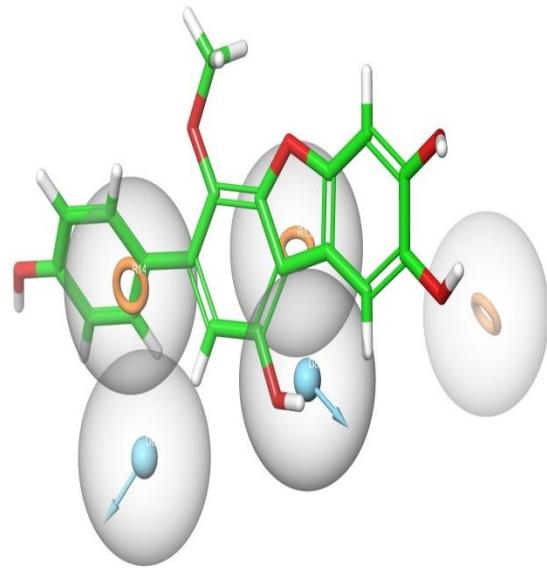
9335



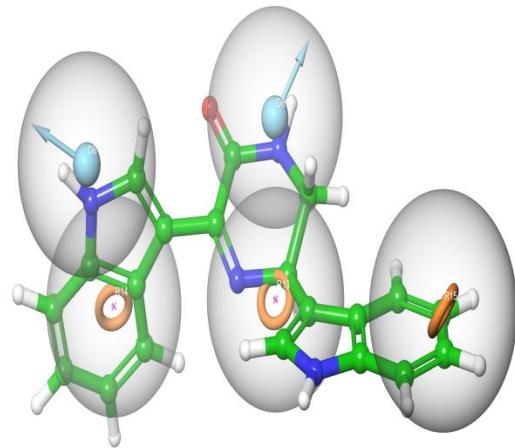
15115



10015



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360799

Figure S2. The model map onto top leads.