

Targeting 3CLpro and SARS-CoV-2 RdRp by *Amphimedon* sp. metabolites: a computational study

Nourhan Hisham Shady ^a, Alaa M. Hayallah ^{b,c}, Mamdouh F. A. Mohamed ^d, Mohammed M. Ghoneim ^{e,f}, Garri Chilingaryan ^g, Mohammad M. Al-Sanea ^h, Mostafa A. Fouad ⁱ, Mohamed Salah Kamel ^{a,i} and Usama Ramadan Abdelmohsen ^{a,i*}

^a Department of Pharmacognosy, Faculty of Pharmacy, Deraya University, Universities Zone, P.O. Box 61111 New Minia City, Minia, Egypt.

^b Department of Pharmaceutical Organic Chemistry, Faculty of Pharmacy, Assiut University, Assiut 71526, Egypt.

^c Pharmaceutical Chemistry Dept. Faculty of pharmacy, Sphinx University. New Assiut. Egypt.

^d Department of Pharmaceutical Chemistry, Faculty of Pharmacy, Sohag University, 82524 Sohag, Egypt.

^e Department of Pharmacy Practice, College of Pharmacy, AlMaarefa University, Ad Diriyah 13713, Saudi Arabia.

^f Department of Pharmacognosy, Faculty of Pharmacy, Al-Azhar University, Cairo, 11371, Egypt

^g Institute of Biomedicine and Pharmacy, Russian –Armenian University, Armenia.

^h Department of Pharmaceutical Chemistry, College of Pharmacy, Jouf University, Aljouf 72341, Saudi Arabia.

ⁱ Department of Pharmacognosy, Faculty of Pharmacy, Minia University, 61519 Minia, Egypt

* Correspondence: usama.ramadan@ mu.edu.eg, Tel.: +2-86-2347759, Fax: +2-86-2369075 (URA);

Table S1. The CDOCKER energy and CDOCKER interaction energy of the virtually screened compounds, Remdesivir and Ligand N3 on RdRp and 3CL pro enzymes.

Compound	RdRp (PDB ID: 6NUR)		3CL protease (PDB ID: 6LU7).	
	CDOCKER energy	CDOCKER interaction energy	CDOCKER energy	CDOCKER interaction energy
1	-45.4891	-51.245	-56.5311	-48.047
7	-24.9113	-55.4676	-22.9961	-46.8449
8	-14.9527	-33.013	-18.0475	-40.973
9	-34.6093	-36.2722	-39.2038	-39.0384
10	-15.8932	-43.8066	-16.4872	-50.1956
11	-40.4527	-55.0797	-40.2121	-50.9693
12	-33.0222	-57.2272	-26.4258	-47.9309
14	-2.54925	-31.7949	-4.5213	-37.3417
Remdesivir	-30.1162	-59.1337	ND	ND
Ligand N3	ND	ND	-70.8463	-79.0435

Table S2. Different interaction energy types and total binding energies of compounds 1 and 12 with SARS-CoV-2 RdRp (PDB ID 6NUR) and 3CL protease (PDB ID 6LU7).

	Van der Waal s (kJ/m ol)	Electrost atic (kJ/mol)	Polar Solvati on (kJ/mo l)	SAS A (kJ/m ol)	Bindi ng Ener gy (kJ/m ol)	Standa rd deviati on (kJ/mo l)
RdRp						
Remdes ivir	- 135.85	-94.41	202.29	-16.83	-44.77	± 2.62
Compo und 1	- 203.89	-32.99	215.63	-23.69	-45.12	± 2.45
Compo und 12	- 148.28	-135.13	255.7	-20.4	-47.86	± 2.33
3CL protease						
Compo und N3	- 284.57	-96.25	251.63	-26.83	- 156.02	± 2.46
Compo und 1	- 149.93	-87.39	183.66	-16.02	-69.81	± 2.03
Compo und 12	-198.6	-33.42	123.96	-19.02	- 127.33	± 1.79

Table S3. Pharmacokinetic properties of compounds.

Property		Compound 1	Compound 7	Compound 8	Compound 9	Compound 10	Compound 11
Absorption	Water Solubility (log mol/L)	-2.94	-6.781	-6.419	-6.309	-7.183	-5.578
	Caco2 permeability (log Papp in 10 ⁻⁶ cm/s)	0.919	1.127	1.575	1.719	1.565	1.264
	Intestinal absorption (%)	88.201	90.9	90.63	92.925	89.723	86.168
	Skin permeability (log Kp)	-2.73	-2.758	-2.661	-2.681	-2.758	-2.734
	P-glycoprotein substrate	Yes	No	No	No	No	Yes
	P-glycoprotein I	No	Yes	No	No	No	No
	P-glycoprotein II	Yes	Yes	No	No	Yes	Yes
Distribution	VDss (log L/kg)	-0.113	0.566	-0.255	0.237	-0.295	-0.644
	Fraction unbound (Fu)	0.349	0	0.085	0.116	0.034	0.098
	BBB permeability (log BB)	-0.477	-0.396	-0.243	0.224	-0.393	-0.731
	CNS permeability (log PS)	-2.709	-2.118	-1.831	-1.984	-1.507	-0.559
Metabolism	CYP2D6 substrate	Yes	No	No	No	No	No
	CYP3A4 substrate	Yes	Yes	Yes	Yes	Yes	Yes
	CYP1A2 inhibitor	No	No	Yes	Yes	Yes	No
	CYP2C19 inhibitor	No	No	No	No	No	No
	CYP2C9 inhibitor	No	No	No	No	No	No
	CYP2D6 inhibitor	No	No	No	No	No	No
	CYP3A4 inhibitor	No	No	No	No	No	No
Excretion	Total Clearance (log ml/min/kg)	1.184	1.605	1.784	1.982	1.996	1.985
	Renal OCT2 substrate	No	No	No	No	No	No
Toxicity	AMES toxicity	No	No	No	No	No	No
	Max. tolerated dose (log mg/kg/day)	-0.512	0.238	-0.305	0.317	-0.676	-1.166
	hERG I inhibitor	No	No	No	No	No	No
	hERG II inhibitor	No	Yes	No	No	Yes	No
	Oral Rat Acute Toxicity (LD50) (mol/kg)	2.719	2.041	1.754	1.67	1.749	3.158
	Oral Rat Chronic Toxicity (LOAEL) (mol/kg_bw/day)	1.823	2.624	3.095	2.727	3.313	3.86
	Hepatotoxicity	Yes	No	No	No	No	No
	Skin Sensitisation	No	No	Yes	Yes	Yes	Yes

	<i>T.Pyriformis</i> toxicity (log µg/L)	0.293	0.336	1.702	1.7	0.932	0.319
	Minnow toxicity (log mM)	-1.235	-4.33	-0.975	-1.335	-1.607	-2.736

Property		Compound 12	Compound 14			
Absorption	Water Solubility (log mol/L)	-3.788	-3.09			
	Caco2 permeability (log Papp in 10 ⁻⁶ cm/s)	0.597	1.389			
	Intestinal absorption (%)	83.364	100			
	Skin permeability (log Kp)	-2.742	-2.733			
	P-glycoprotein substrate	Yes	No			
	P-glycoprotein I	Yes	No			
	P-glycoprotein II	No	Yes			
Distribution	VDss (log L/kg)	-1.3	0.458			
	Fraction unbound (Fu)	0.235	0.336			
	BBB permeability (log BB)	-1.608	0.607			
	CNS permeability (log PS)	-3.491	-2.057			
Metabolism	CYP2D6 substrate	No	No			
	CYP3A4 substrate	Yes	Yes			
	CYP1A2 inhibitor	No	Yes			
	CYP2C19 inhibitor	No	Yes			
	CYP2C9 inhibitor	No	Yes			
	CYP2D6 inhibitor	No	No			
	CYP3A4 inhibitor	No	Yes			
Excretion	Total Clearance (log ml/min/kg)	1.762	0.835			
	Renal OCT2 substrate	No	No			
Toxicity	AMES toxicity	No	No			
	Max. tolerated dose (log mg/kg/day)	-0.058	0.425			
	hERG I inhibitor	No	No			
	hERG II inhibitor	Yes	No			
	Oral Rat Acute Toxicity (LD50) (mol/kg)	4.162	2.61			
	Oral Rat Chronic Toxicity (LOAEL) (mol/kg_bw/day)	4.417	0.649			
	Hepatotoxicity	Yes	Yes			
	Skin Sensitisation	No	No			

	<i>T.Pyriformis</i> toxicity (log µg/L)	0.285	0.285			
	Minnow toxicity (log mM)	2.319	0.612			