

ELECTRONIC SUPPLEMENTARY DATA

Toward an efficient and eco-friendly route for the synthesis of dimeric 2,4-diacylphloroglucinol and its potential as a SARS-CoV-2 main protease antagonist insight from in silico studies

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S1: EXPERIMENTAL

Materials and instrumentations

All of reagents and solvents were purchased from Merck or Sigma-Aldrich and utilised without additional purification. Branson B1510R-DTH Ultrasonic Cleaner was used to perform any reactions under sonication condition. Silica gel with pore size 60 Å, 230-400 mesh was used to perform column chromatography. Thin Layer Chromatography (TLC), which performed on gel 60 F₂₅₄ (Merck) plate was used to monitor the progress of the reactions. UV light at $\lambda=254$ and 365 nm was used for visualisation of TLC spot. Agilent DD2 Nuclear magnetic resonance (NMR) Spectrometer was used to record NMR spectra, 500 MHz for ¹H and 125 MHz for ¹³C using CDCl₃ as the solvent, operated at 25 °C and using tetramethyl silane as the internal standard. Melting Point Apparatus SMP10 Stuart was used to measure melting points of the isolated product.

Preparation of SAA catalyst

The catalyst was prepared following the previous work¹. A 12.375 g of silica gel was soaked in 50 mL of diethyl ether (Et₂O) and 6.25 mmol of concentrated H₂SO₄. The obtained mixture was stirred for 15 min at ambient temperature and pressure under the atmospheric nitrogen (N₂) and dried at 60 °C for 18 h. The solvent in the resulted residue was then removed by heating at 120 °C for 6 h. Finally, before further use, the resulted **SSA** catalyst was placed in a desiccator for storage.

General procedure for the synthesis of 2,4-diacylphloroglucinols (2a-2c)

The synthesis of monomeric diacylphloroglucinols was performed by following the previous work¹. To the round bottom flask, phloroglucinol **1** (1 mmol) and acetic anhydride or acyl chlorides (2 mmol) were stirred at ambient temperature and pressure for 5 min. Then, into the mixture, 20% (w/w) **SSA** as a catalyst was added and sonicated at 60 °C for 15-20 min. Upon reaction completion, the mixture was cooled, diluted with 15 mL of water, and extracted with ethyl acetate (EtOAc) (3 x 15 mL). Finally, the solvent was evaporated and the crude product was subjected for further purification through short flash column chromatography using n-hexane/EtOAc (95/5 → 70/30 gradient) in silica gel. All of the ¹H NMR and ¹³C NMR spectra of monomers are given in **ESI S2**.

General procedure for the synthesis of 4a via a condensation reaction. In a vial was placed monomer **2a** (0.1 mmol), formaldehyde (0.5 mmol) and **SSA**. The resulted mixture was reacted under ultrasound irradiation. The reaction typically took 10-15 min which monitored by TLC. Upon completion of the reaction, 4 M hydrogen chloride (HCl) was used to neutralise the crude solution. EtOAc (3 x 10 mL) was then used to extract the aqueous layer. The combined organic layers were dried over sodium sulphate anhydrous (Na₂SO₄). Finally, the solvent was removed and the crude product was collected and used without using further purification.

General procedure for the synthesis of 4a via one-pot synthesis using Eschenmoser's salt

To the round bottom flask, a solution of the monomer of diacylphloroglucinol **2a** (0.1 mmol) and the suspension of Eschenmoser's salt (0.5 eq.) was heated at 60 °C in the small amount of chloroform (CHCl₃) under the US method in N₂ atmosphere. The obtained mixture was refluxed until most of diacylphloroglucinol converted to the desired product. To the obtained mixture, 5 mL dichloromethane (CH₂Cl₂) was then added and gently washed with 5 mL of 1 N aqueous HCl and brine solution. CH₂Cl₂ (3 x 5 mL) was used to extract the resulted aqueous layer. The combined organic layers were dried over magnesium sulphate anhydrous (MgSO₄), filtered with filter paper and evaporated using a rotary evaporator. The obtained crude residue was purified by short flash chromatography using n-hexane/EtOAc (95/5 → 80/20 gradient) in silica gel.

Synthesis of arylation of dimethylammonium Salt 3

A solution of the desired acylated phloroglucinol **2a** (1.0 eq.) was reacted with the corresponding dimethylaminomethylated monomer (1.0 eq.) using small amount of toluene under US method at room temperature until most of the acylated phloroglucinol converted to the expected product. Upon completion of the reaction, the solvent was removed. Finally, before further used, CH₂Cl₂ was used to wash the crude product.

1-(3,5-diacetyl-2,4,6-trihydroxyphenyl)-N,N-dimethylmethanaminium (3). Yield: 98%; white coloured solid; ¹H NMR (500 MHz, CDCl₃): δ 9.31 (tt, J =

6.6, 2.9 Hz), 4.45 (d, $J = 6.6$ Hz), 2.67 (t, $J = 1.5$ Hz). ^{13}C NMR (125 MHz, CDCl_3): δ 206.09, 165.70, 104.31, 103.11, 48.75, 40.60, 32.63.

General Synthesis of Dimeric Acylphloroglucinol via Arylation of Dimethylammonium Salt. A solution of diacylphloroglucinol monomer **2a** (1.0 eq.) with the activated dimethylammonium salts from monomer **3** (1.0 eq.) was heated at 80 °C in a small amount of toluene for the designated time. The solvent was then removed and the crude product was collected and used without using further purification.

Methylene-bis-(3,5-diacetyl-2,4,6-trihydroxybenzene) (4a): Yield: 94%; cream coloured solid; mp 279–281 °C; ^1H NMR (CDCl_3 , 500 MHz) δ ppm: 4.03 (s, 2H, $-\text{CH}_2-$), 2.76 (s, 12H, 4 x $-\text{CH}_3$). ^{13}C NMR (CD_3Cl_3 , 125 MHz) δ ppm: 205.7, 166, 164.42, 106, 107.31, 32.63, 16.40.

Methylene-bis-(3,5-di-isobutanoyl-2,4,6-trihydroxybenzene) (4b): Yield: 90%; light orange coloured solid; mp 200–202 °C; ^1H NMR (CDCl_3 , 500 MHz) δ ppm: 3.98 (s, 2H, $-\text{CH}_2-$), 3.51 (m, 2H, $J = 5.4$ Hz, 2 x $-\text{CH}(\text{CH}_3)_2$), 1.2 (d, $J = 5.4$ Hz, 24H, 4 x CH_3). ^{13}C NMR (CDCl_3 , 125 MHz) δ ppm: 212, 171.03, 167.56, 163.37, 108.13, 106.12, 39.74, 18.78, 17.17.

Methylene-bis-(3,5-di-isopentanoyl-2,4,6-trihydroxybenzene) (4c): Yield: 88%; yellow coloured solid; mp 158–160 °C; ^1H NMR (CDCl_3 , 500 MHz) δ ppm: 4.00 (s, 2H, $-\text{CH}_2-$), 3.01 (d, $J = 5.6$ Hz, 8H, 4 x $-\text{CH}_2-\text{CH}(\text{CH}_3)_2$), 2.27 (dt, $J = 11.0, 5.5$ Hz, 4H, 4 x $\text{CH}_2-\text{CH}(\text{CH}_3)_2$), 1.00 (d, $J = 5.4$ Hz, 24H, 8 x CH_3). ^{13}C NMR (CDCl_3 , 125 MHz) δ ppm: 208.00, 171.3, 165.78, 162.82, 108.39, 105.50, 53.50, 25.91, 23.10, 16.89.

Methyl-methylene-bis-(3,5-diacetyl-2,4,6-trihydroxybenzene) (4d): Yield: 82%; yellow coloured solid; mp 206–208 °C. ^1H NMR (CDCl_3 , 500 MHz) δ ppm: 4.78 (d, $J = 5.5$ Hz, 1H, $>\text{CH}-\text{CH}_3$), 2.76 (s, 12H, 4 x $-\text{CH}_3$), 1.56 (d, $J = 5.4$ Hz, 3H, $>\text{CH}-\text{CH}_3$). ^{13}C NMR (CDCl_3 , 125 MHz) δ ppm: 206.10, 171.2, 166.37, 164.51, 112.59, 107.08, 33.06, 26.48, 21.37.

Phenyl-methylene-bis-(3,5-diacetyl-2,4,6-trihydroxybenzene) (4e): Yield: 64%; cream coloured solid; mp 109–111 °C. ^1H NMR (CDCl_3 , 500 MHz) δ ppm: 7.33 – 7.21 (m, 3H, ArH), 7.20 (d, $J = 0.6$ Hz, 2H, ArH), 5.32 (s, 1H, $>\text{CH}$ -phenyl), 2.70 (s, 12H, 4 x CH_3). ^{13}C NMR (CDCl_3 , 125 MHz) δ ppm: δ 206.00, 166.27, 171.6, 164.36, 140.87, 128.9, 128.47, 127.21, 109.78, 107.16, 36.34, 32.67.

Reusability and heterogeneity of SSA catalyst

After the completion of the reaction, the collected **SSA** catalyst was stirred in 10 mL of EtOAc for 5-10 min. Then, it was filtered and washed with 5 mL of EtOAc and acetone, respectively. After that, before further used, the filtered catalyst was dried at 120 °C for 3 h. Additionally, a hot filtration technique was performed following the previous methods¹ to observe the possible leaching of **SSA** active sites.

Environmental assessment

To assess the environmental assessment of the present method, the combination between the Environmental Assessment Tool for Organic Syntheses (EATOS) software established by Eissen and Metzger² and the algorithm by Andraos³ were performed. The necessitated data related to their detail of stoichiometric data was attained from the recent literatures⁴⁻⁷ both for EATOS and Andraos algorithm. For further detail about energy efficiency profile, the energy calculation measurement established by Clark and co-workers⁸ was accomplished. As a note, the details of workup and purification stages were not included in this assessment due to the lack of procedures information.

Preparation of protein targets 3CL^{pro} and ligands

The X-ray crystallographic structures of **3CL^{pro}** in complex with an inhibitor N3 was retrieved from the Protein Data Bank database with PDB ID: 6LU7 (<http://www.rscb.org/>), then, chain A was loaded into the AutoDock Tools (version 1.5.6) software. Through this software, the ligand N3, water, and ions molecules were removed. PrankWeb (<http://prankweb.cz/>) was used to identify the active sites of selected receptors. Finally, before the docking studies were performed, polar hydrogen atoms and Gasteiger charges were added.

Density-functional theory (DFT) studies

All of the studied compounds will be sketched with PerkinElmer Informatics ChemDraw (version 17.1). The quantum chemical calculation for all the investigated molecular structures will be carried out in GAUSSIAN 09 suit of program. The optimisation of all the stationary points of desired molecules was calculated using 6-31G (d, p) basis set with the Lee-Yang-Parr exchange-correlation (B3LYP) density functional in the gas phase. It allows calculating the values of orbital energy HOMO and LUMO, total energy, gap energies (ΔE_{gap}).

Then, based on the calculated orbital energy (HOMO and LUMO), it can be used to calculate several other molecular electronic properties including electronegativity (χ) = $-(E_{\text{HOMO}} - E_{\text{LUMO}})/2$, electron affinity (A) = $-E_{\text{LUMO}}$, ionisation energy (I) = $-E_{\text{HOMO}}$, chemical potential (μ) = $1/2 (E_{\text{LUMO}} + E_{\text{HOMO}})$, chemical softness (S) = $1/2 \eta$, chemical hardness (η) = $1/2 (E_{\text{LUMO}} - E_{\text{HOMO}})$ and electrophilicity index (ω) = $\mu^2/2 \eta^{9,10}$. Additionally, Molecular electrostatic potential (MEP) calculation derived from Mulliken charge was performed to access the detail information of nucleophilic and electrophilic sites.

Molecular docking studies

Molecular docking has extensively used in structure-based drug design¹¹. To assess the binding difference between the synthesised dimeric 2,4-diacetyl phloroglucinol and its derivatives along with N3, Chloroquine, and Remdesivir as a reference with the **3CL^{pro}**, a comparative molecular docking analysis was performed using PyRx software by AutoDock wizard¹², which generated using Grid box centre_x= -14.1744035865, centre_y= 13.2322187073, centre_z= 68.9220623083, size_x= 20.4592380327, size_y= 23.6792741601, size_z= 26.2300864435 and exhaustiveness = 8. The interactions between the receptor and the result of the concerned ligands were compiled, visualised and analysed using PyMol (version 1.7.4) and BIOVIA Discovery Studio Visualizer 2020.

Molecular target prediction

The synthesised **4a** and its derivatives may interact with several numbers of protein bind to proteins or other macro-molecular targets. Accordingly, it is essential to predict the molecular targets for these investigated molecules. It can be easily investigated using an accessible free of charge Swiss Target Prediction (<http://www.swisstargetprediction.ch/index.php>)¹³.

Drug likeness prediction and ADMET analysis

Drug-likeness properties of ligands were assessed using SwissADME (<http://www.swissadme.ch/>). The ADMET ((i.e., absorption, distribution, metabolism, excretion, and toxicity) analysis was accessed using admetSAR (<http://lmm.d.ecust.edu.cn/admetSar1>) and pKCSM approach (<http://biosig.unimelb.edu.au/pkcsM/>)¹⁴.

Prediction of activity spectra for substances (PASS)

The prediction of activity, particularly antiviral activities of the **4a** and its derivatives were assessed using PASS (<http://www.way2drug.com/>). This is an accessible free of charge a computer program, which can estimate the activity of molecules¹⁵. Through the basic structural formula, it could predict about 4,000 types of biological activity with an average accuracy of about 90%¹⁶, therefore, the use of PASS to discover and optimize new lead compounds is very reasonable.

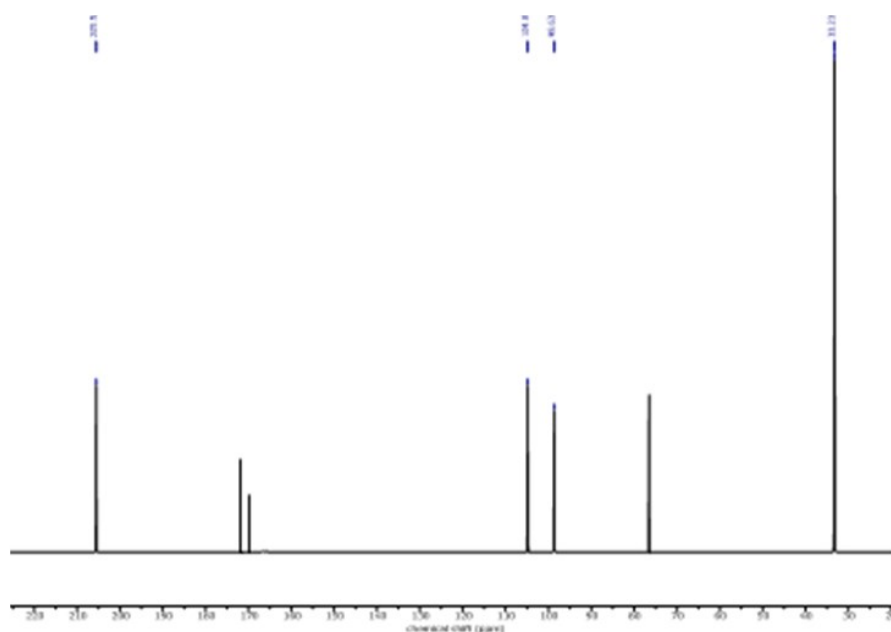
S2: ^1H NMR AND ^{13}C NMR SPECTRA

After purification and characterization, all of the products were characterized using NMR and the spectral data of ^1H NMR and ^{13}C NMR were compared with the relevant literature^{4,17}.

1,1'-(2,4,6-trihydroxy-1,3-phenylene)bis(ethan-1-one) (2a). ^1H NMR (CD_3Cl_3 , 500 MHz) δ ppm: 6.19 (s, 1H, ArH), 2.64 (s, 6H, 2 x CH_3). ^{13}C NMR (CD_3Cl_3 , 125 MHz) δ ppm: 205.56, 173.56, 170.65, 104.83, 98.63, 33.23.



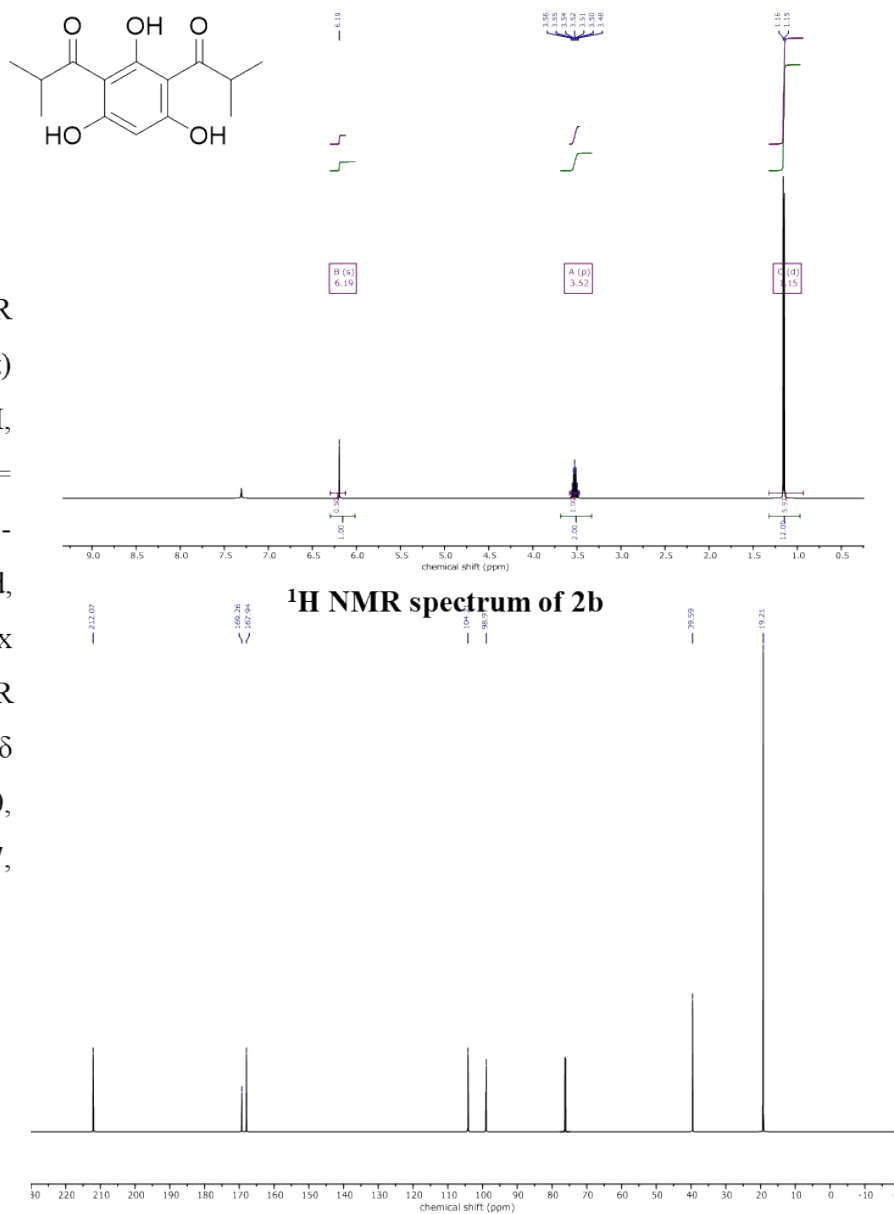
^1H NMR spectrum of 2a



^{13}C NMR spectrum of 2a

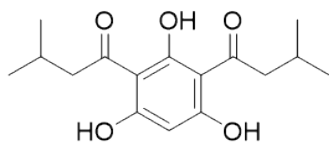
Yield: 95%

1,1'-(2,4,6-trihydroxy-1,3-phenylene)bis(2-methylpropan-1-one) (2b). ^1H NMR (CD_3Cl_3 , 500 MHz) δ ppm: 6.19 (s, 1H, ArH), 3.52 (p, $J = 5.4$ Hz, 2H, 2 x - $\text{CH}(\text{CH}_3)_2$), 1.15 (d, $J = 5.4$ Hz, 12H, 4 x CH_3). ^{13}C NMR (CD_3Cl_3 , 125) δ ppm: 212.07, 170, 168, 104.21, 98.97, 39.59, 19.21.

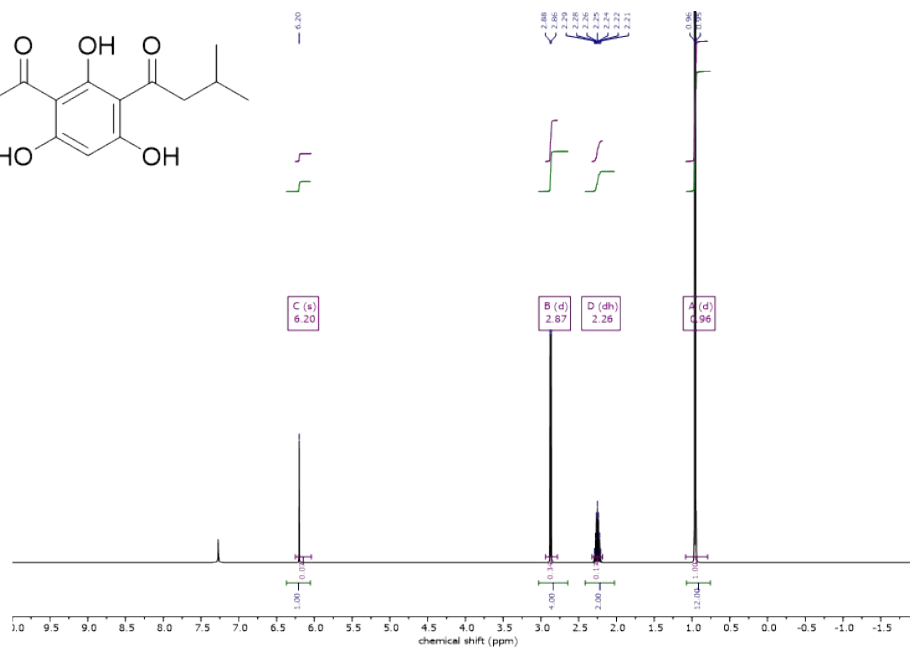


Yield: 89%

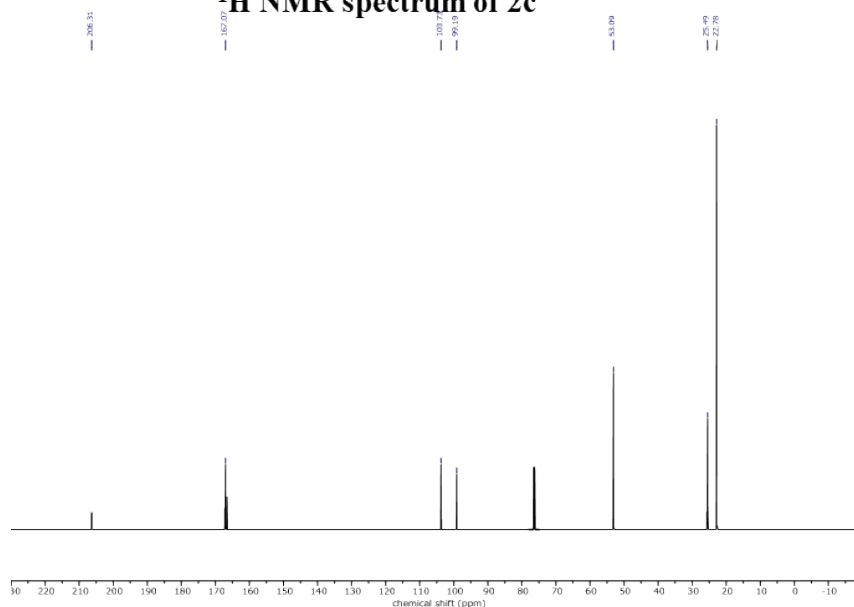
1,1'-(2,4,6-trihydroxy-1,3-phenylene)bis(3-methylbutan-1-



one) (2c). ^1H NMR (CD_3Cl_3 , 500 MHz) δ ppm: : 6.20 (s, 1H, ArH), 2.9 (d, $J = 5.6$ Hz, 4H, 2 x $-\text{CH}_2-\text{CH}(\text{CH}_3)_2$), 2.26 (dh, $J = 11.0, 5.5$ Hz, 2H, 2 x $\text{CH}_2-\text{CH}(\text{CH}_3)_2$), 0.97 (d, $J = 5.4$ Hz, 12H, 4x CH_3). ^{13}C NMR (CD_3Cl_3 , 125 MHz) δ ppm: 205.98, 167.07, 103.73, 99.19, 53.09, 24.94, 22.81.



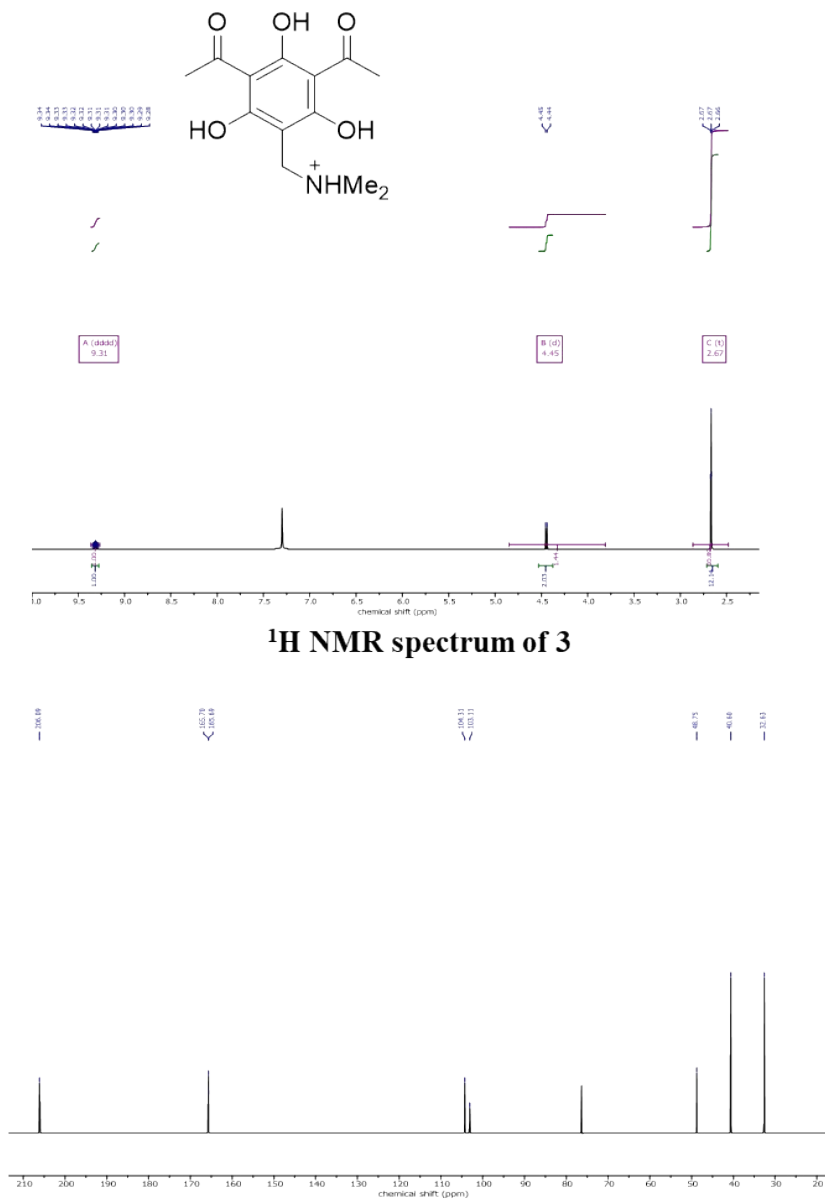
^1H NMR spectrum of 2c



^{13}C NMR spectrum of 2c

Yield: 88%

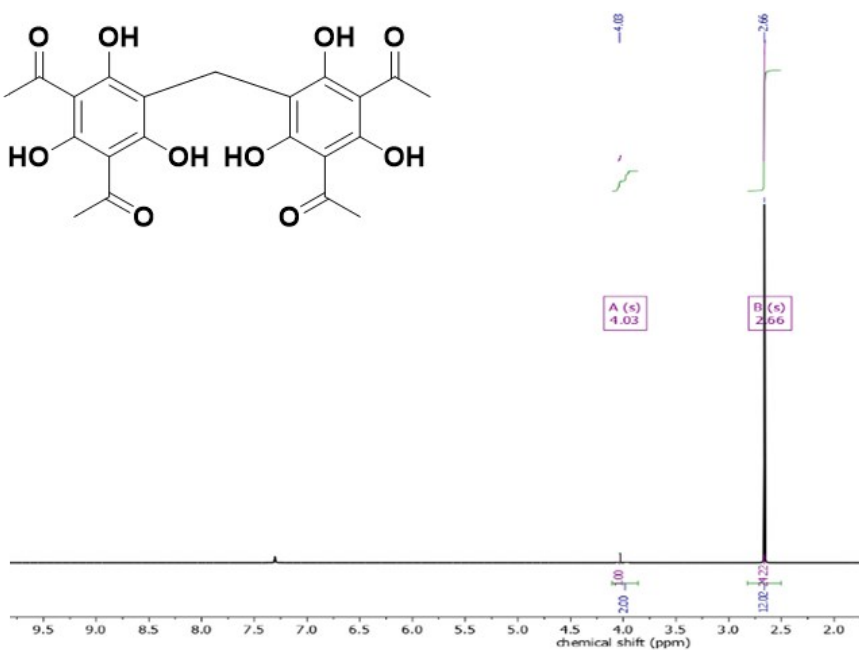
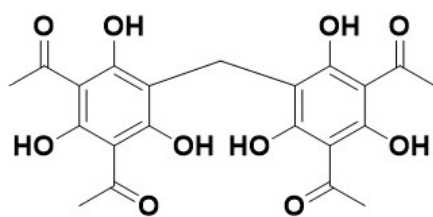
1-(3,5-diacetyl-2,4,6-trihydroxyphenyl)-N,N-dimethylmethanaminium **3.** ^1H NMR (CD_3Cl_3 , 500 MHz) δ ppm: 9.31 (dddd, $J = 9.5, 6.6, 5.9, 3.0$ Hz), 4.45 (d, $J = 6.6$ Hz), 2.67 (t, $J = 1.5$ Hz). ^{13}C NMR (CD_3Cl_3 , 125 MHz) δ ppm: 206.09, 165.70, 165.69, 104.31, 103.11, 48.75, 40.60, 32.63.



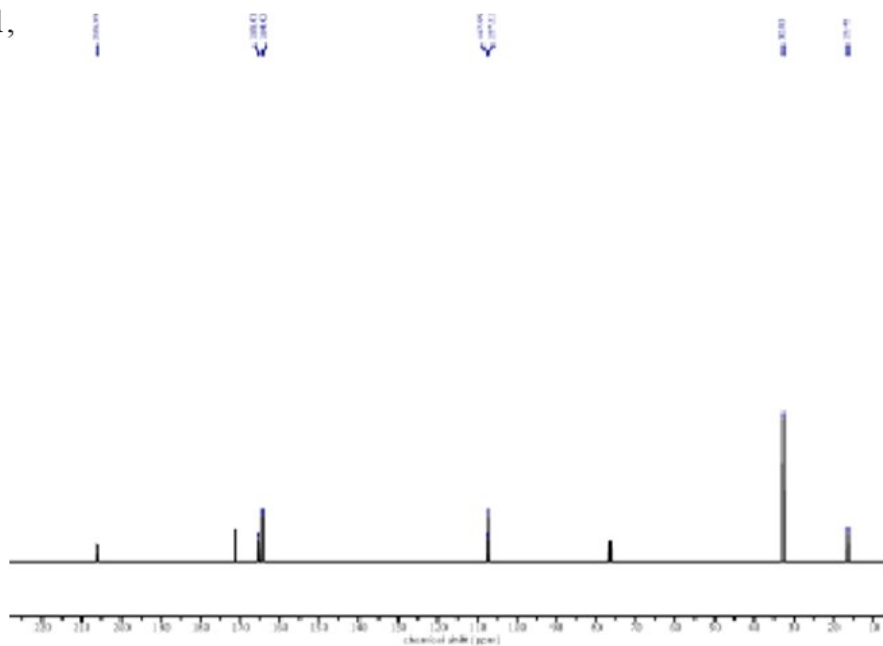
Yield: 98%

**Methylene-bis-
(3,5-diacetyl-2,4,6-
trihydroxybenzene**

) (4a): ^1H NMR
(CD_3Cl_3 , 500 MHz)
 δ ppm: 4.03 (s, 2H, -
 CH_2 -), 2.76 (s, 12H,
4 x $-\text{CH}_3$). ^{13}C NMR
(CD_3Cl_3 , 125 MHz)
 δ ppm: 205.7, 166,
164.42, 106, 107.31,
32.63, 16.40.



^1H NMR spectrum of 4a

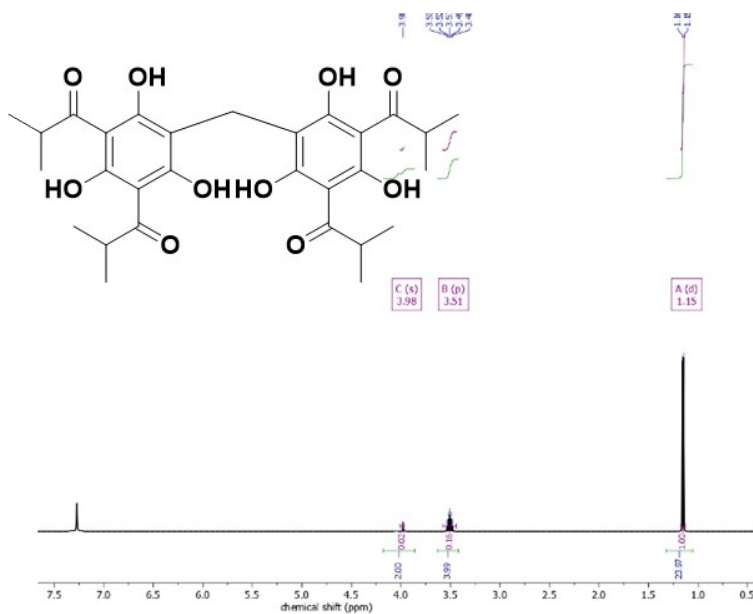


^{13}C NMR spectrum of 4a

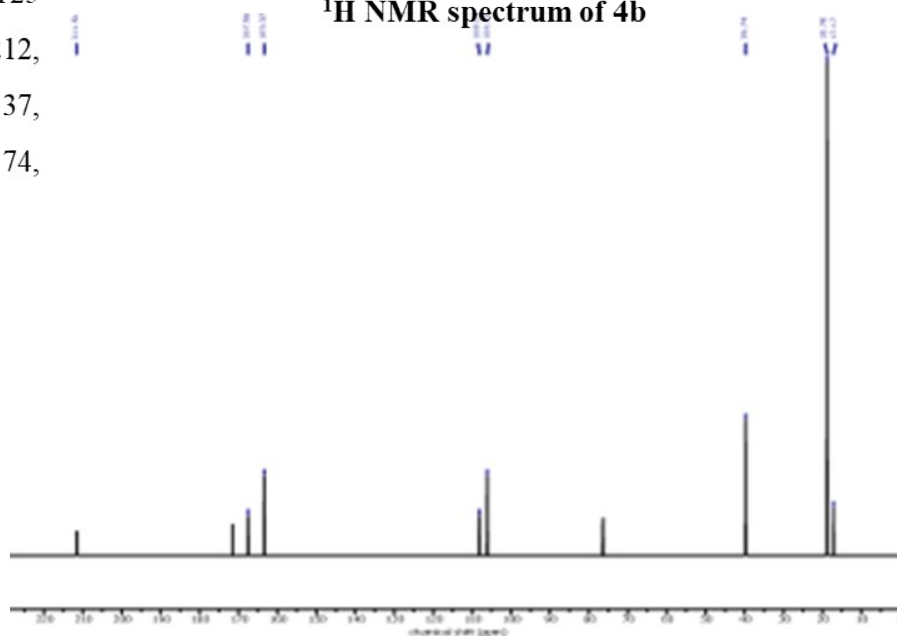
Yield: 94%

Methylene-bis-(3,5-di-isobutanoyl-2,4,6-trihydroxybenzene)

(4b): ^1H NMR (CD_3Cl_3 , 500 MHz) δ ppm: 3.98 (s, 2H, $-\text{CH}_2-$), 3.51 (m, 2H, $J = 5.4$ Hz, 2 x $-\text{CH}(\text{CH}_3)_2$), 1.2 (d, $J = 5.4$ Hz, 24H, 4 x CH_3).
 ^{13}C NMR (CD_3Cl_3 , 125 MHz) δ ppm: 212, 171.03, 167.56, 163.37, 108.13, 106.12, 39.74, 18.78, 17.17.



^1H NMR spectrum of 4b

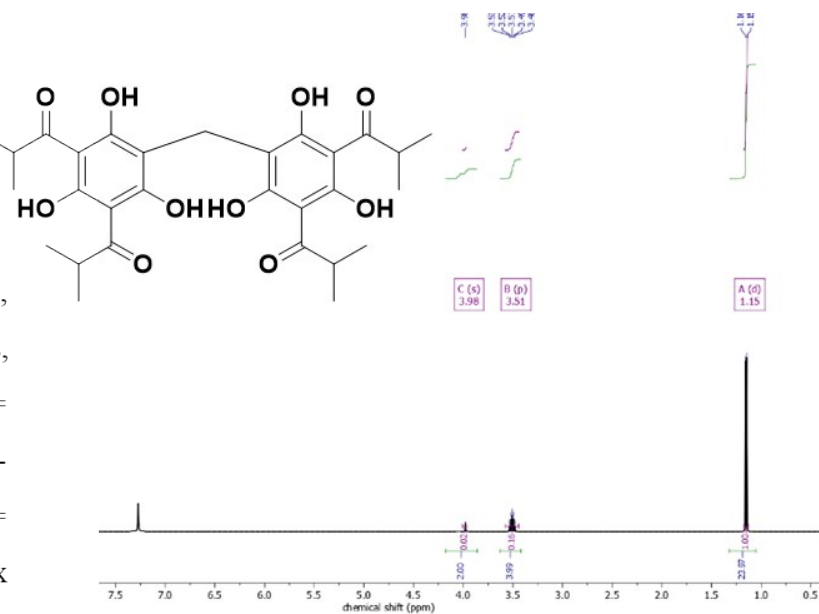


^{13}C NMR spectrum of 4b

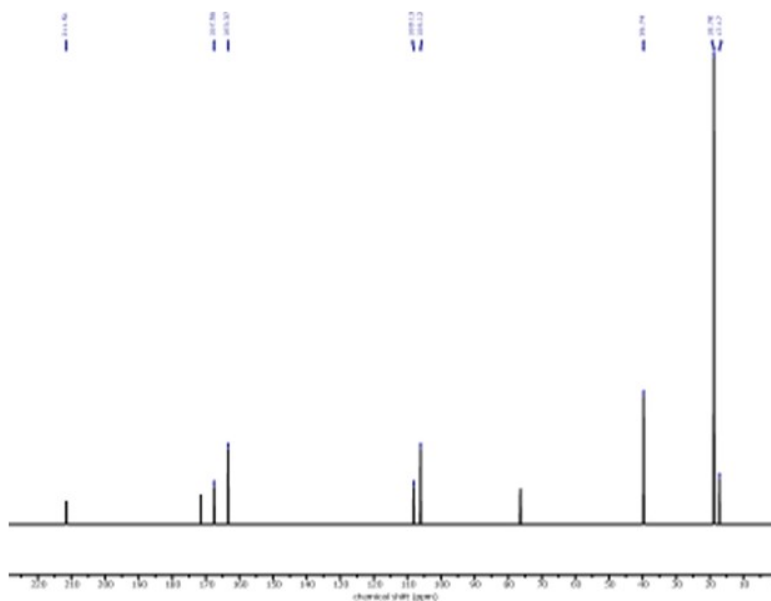
Yield: 90%

Methylene-bis-(3,5-di-isopentanoyl-2,4,6-trihydroxybenzene)

(4c): ^1H NMR (CD_3Cl_3 , 500 MHz) δ ppm: 4.00 (s, 2H, $-\text{CH}_2-$), 3.01 (d, $J = 5.6$ Hz, 8H, 4 x $-\text{CH}_2-\text{CH}(\text{CH}_3)_2$), 2.27 (dt, $J = 11.0, 5.5$ Hz, 4H, 4 x $\text{CH}_2-\text{CH}(\text{CH}_3)_2$), 1.00 (d, $J = 5.4$ Hz, 24H, 8 x CH_3). ^{13}C NMR (CD_3Cl_3 , 125 MHz) δ ppm: 208.00, 171.3, 165.78, 162.82, 108.39, 105.50, 53.50, 25.91, 23.10, 16.89.



^1H NMR spectrum of 4c



^{13}C NMR spectrum of 4c

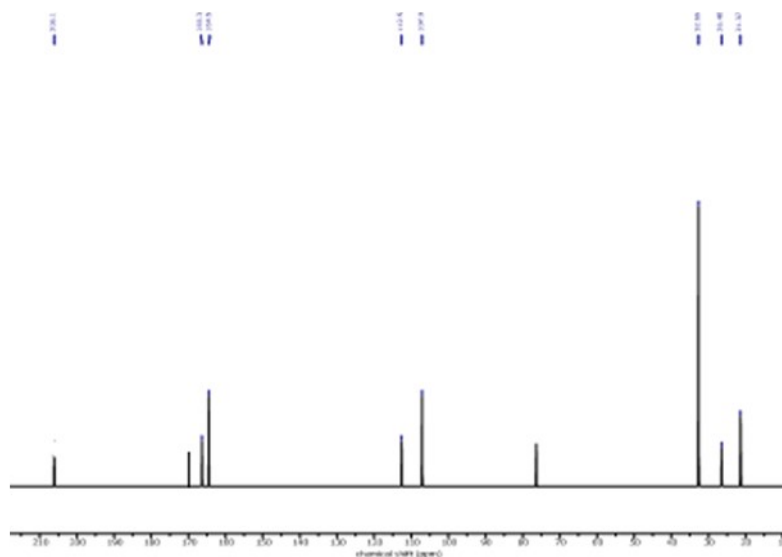
Yield: 88%

**Methyl-methylene-
bis-(3,5-diacetyl-2,4,6-
trihydroxybenzene)**

(4d): ^1H NMR (CD_3Cl_3 , 500 MHz) δ ppm: 4.78 (d, $J = 5.5$ Hz, 1H, $>\text{CH}-\text{CH}_3$), 2.76 (s, 12H, 4 x $-\text{CH}_3$), 1.56 (d, $J = 5.4$ Hz, 3H, $>\text{CH}-\text{CH}_3$). ^{13}C NMR (CD_3Cl_3 , 125 MHz) δ ppm: 206.10, 171.2, 166.37, 164.51, 112.59, 107.08, 33.06, 26.48, 21.37.



^1H NMR spectrum of 4d

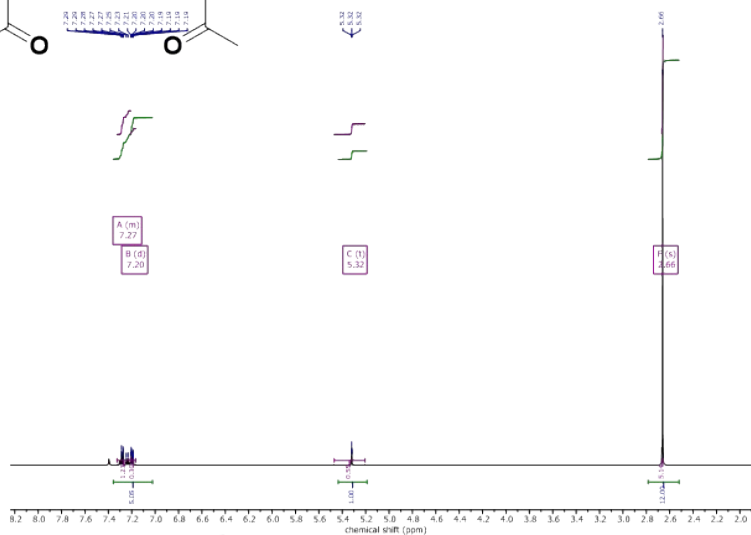
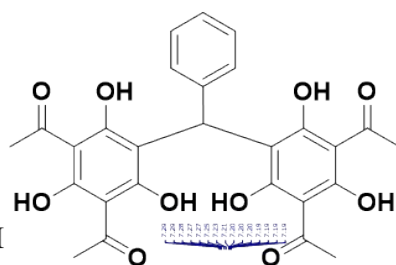


^{13}C NMR spectrum of 4d

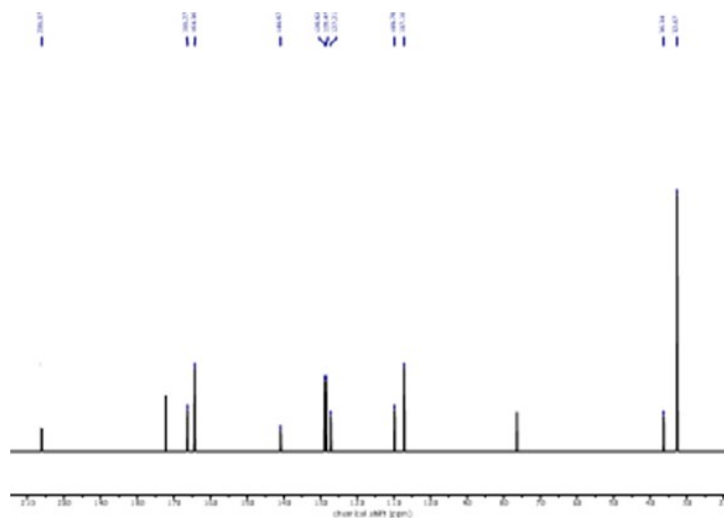
Yield: 82%

Phenyl-methylene-bis-(3,5-diacetyl-2,4,6-trihydroxybenzene)

(4e): 109–111 °C. ^1H NMR (CD_3Cl_3 , 500 MHz) δ ppm: 7.33 – 7.21 (m, 3H, ArH), 7.20 (d, $J = 0.6$ Hz, 2H, ArH), 5.32 (s, 1H, $>\text{CH}$ -phenyl), 2.70 (s, 12H, 4 x CH_3). ^{13}C NMR (CD_3Cl_3 , 125 MHz) δ ppm: δ 206.00, 166.27, 171.6, 164.36, 140.87, 128.9, 128.47, 127.21, 109.78, 107.16, 36.34, 32.67



^1H NMR spectrum of 4e



^{13}C NMR spectrum of 4e

Yield: 64%

S3: ENVIRONMENTAL ASSESSMENT

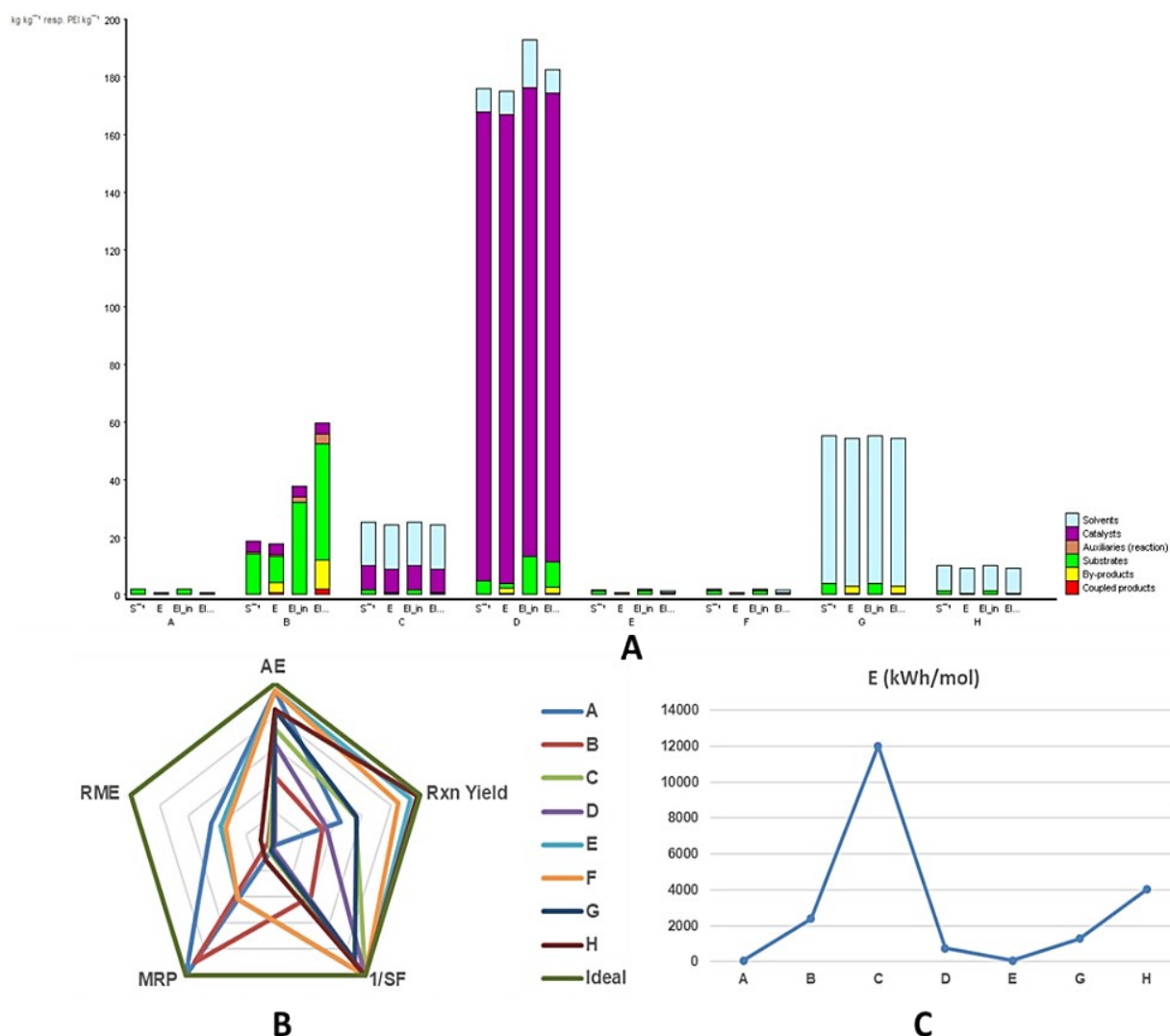


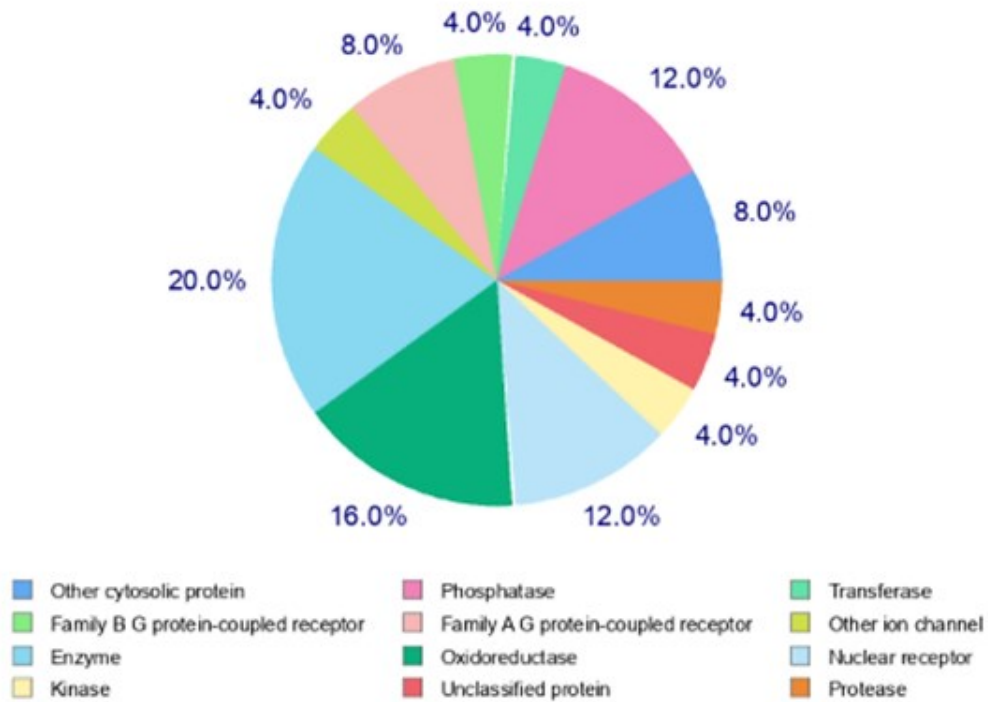
Figure S1. Environmental assessment results investigated by EATOS software (**A**), Andraos's algorithm (**B**), and energy intake requirement (**C**). Method A: MW⁶, Method B: Oil-bath⁷, Method C: Oil-bath⁷, Method D: Oil-bath⁵, Method E: present US, Method F: present LAG, Method G: Eschenmoser's salt, Method H: Dimethylammonium salt.

S4: ENERGY REQUIREMENTS CALCULATION RESULTS

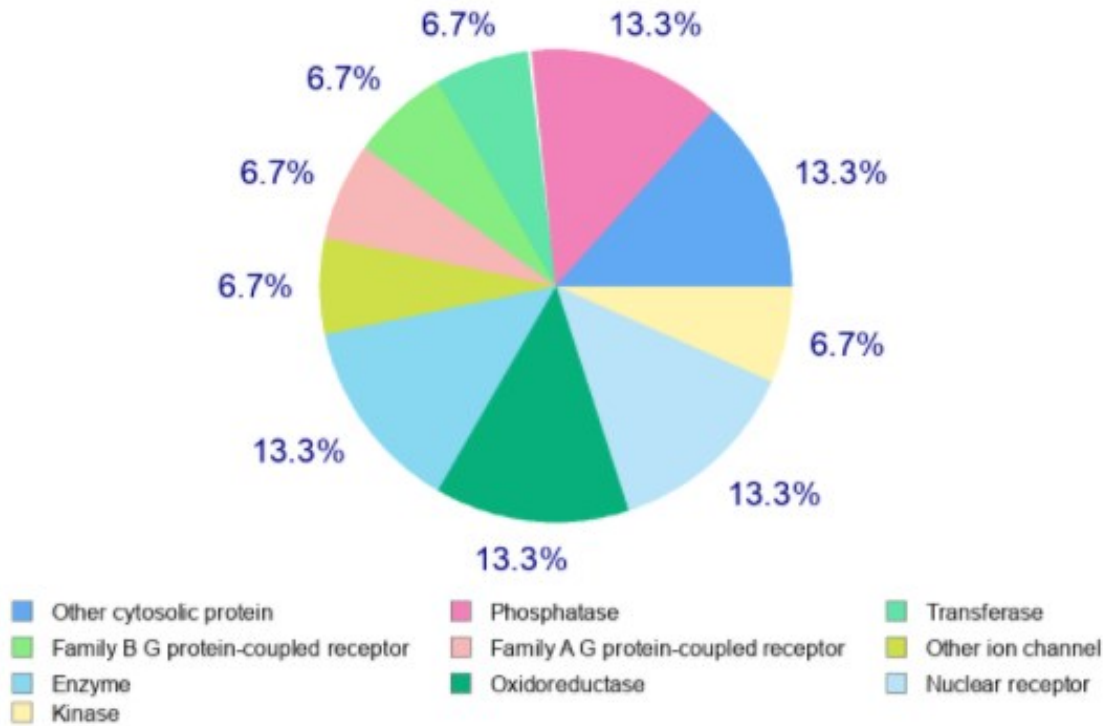
Table S1. Energy requirements calculation results for the synthesis of **4a**

Method	Heating method	mmol	Yield (%)	P (kW)	t (h)	E (kWh)	Reference
A	MW	1.00	88.00	0.70	0.08	0.06	6
B	Oil bath	1.00	30.00	1.44	0.50	0.72	7
C	Oil bath	2.50	60.00	1.44	2.00	2.88	7
D	Oil bath	1.00	95.00	1.44	0.50	0.72	5
E	US	1.00	95.00	0.16	0.33	0.05	Present: US
G	Oil bath	1.00	56.00	1.44	0.50	0.72	Present: Eschenmoscher's salt
H	Oil bath	1.00	98.00	3.16	1.25	3.95	Present: Dimethylammonium salt

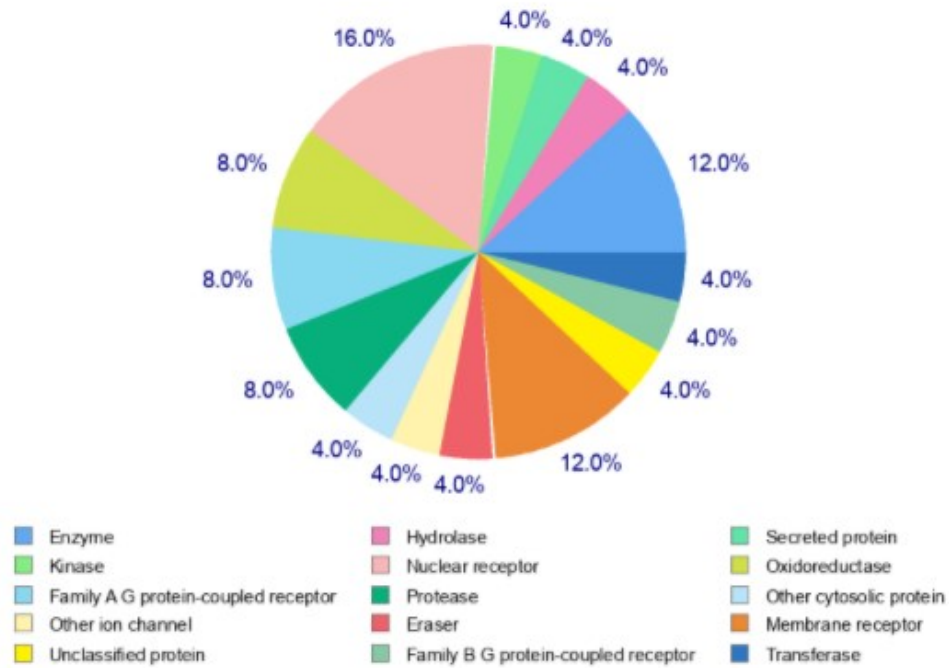
S5: MOLECULAR TARGET PREDICTION



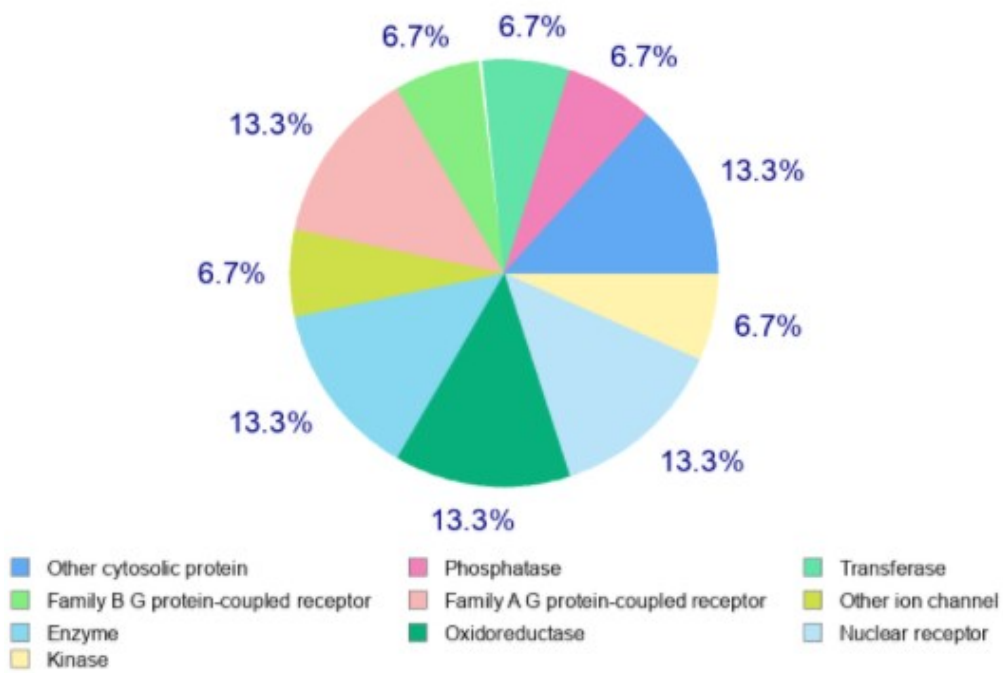
Molecular target prediction of 4a



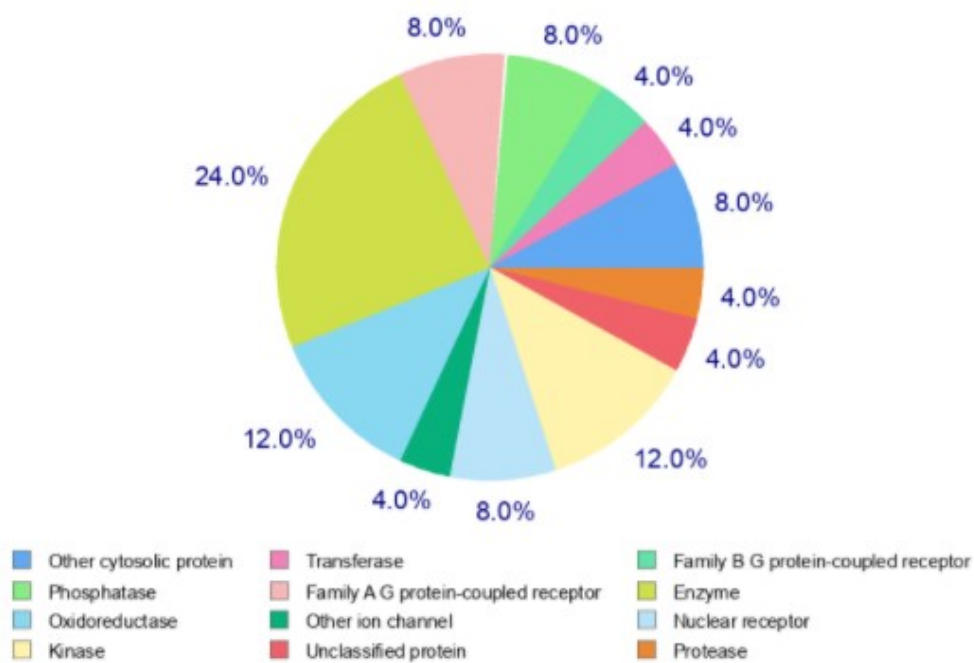
Molecular target prediction of 4b



Molecular target prediction of 4c



Molecular target prediction of 4d



Molecular target prediction of 4e

Figure S2. Molecular target prediction of 4a-e

S6: SURFACE REPRESENTATION

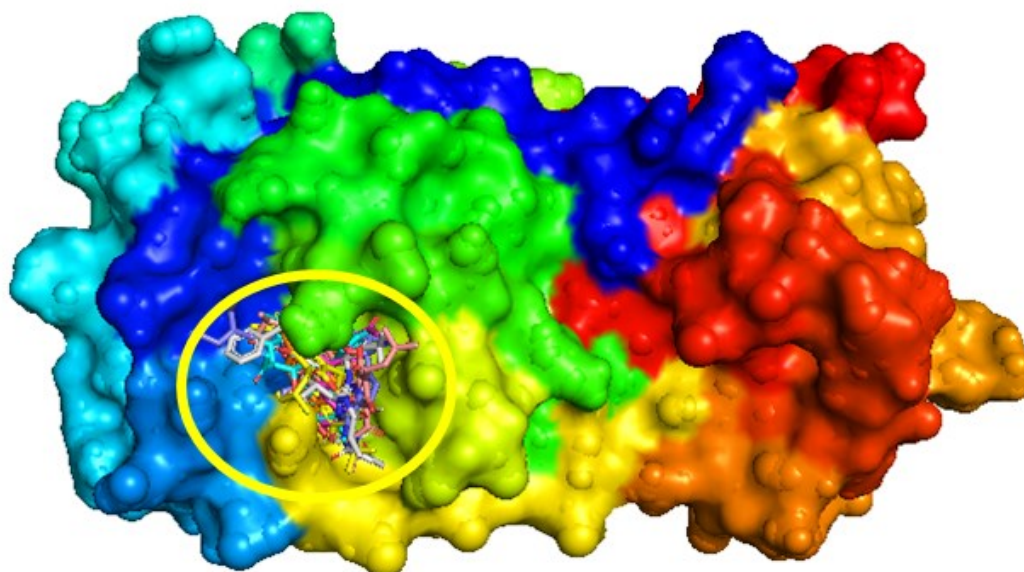
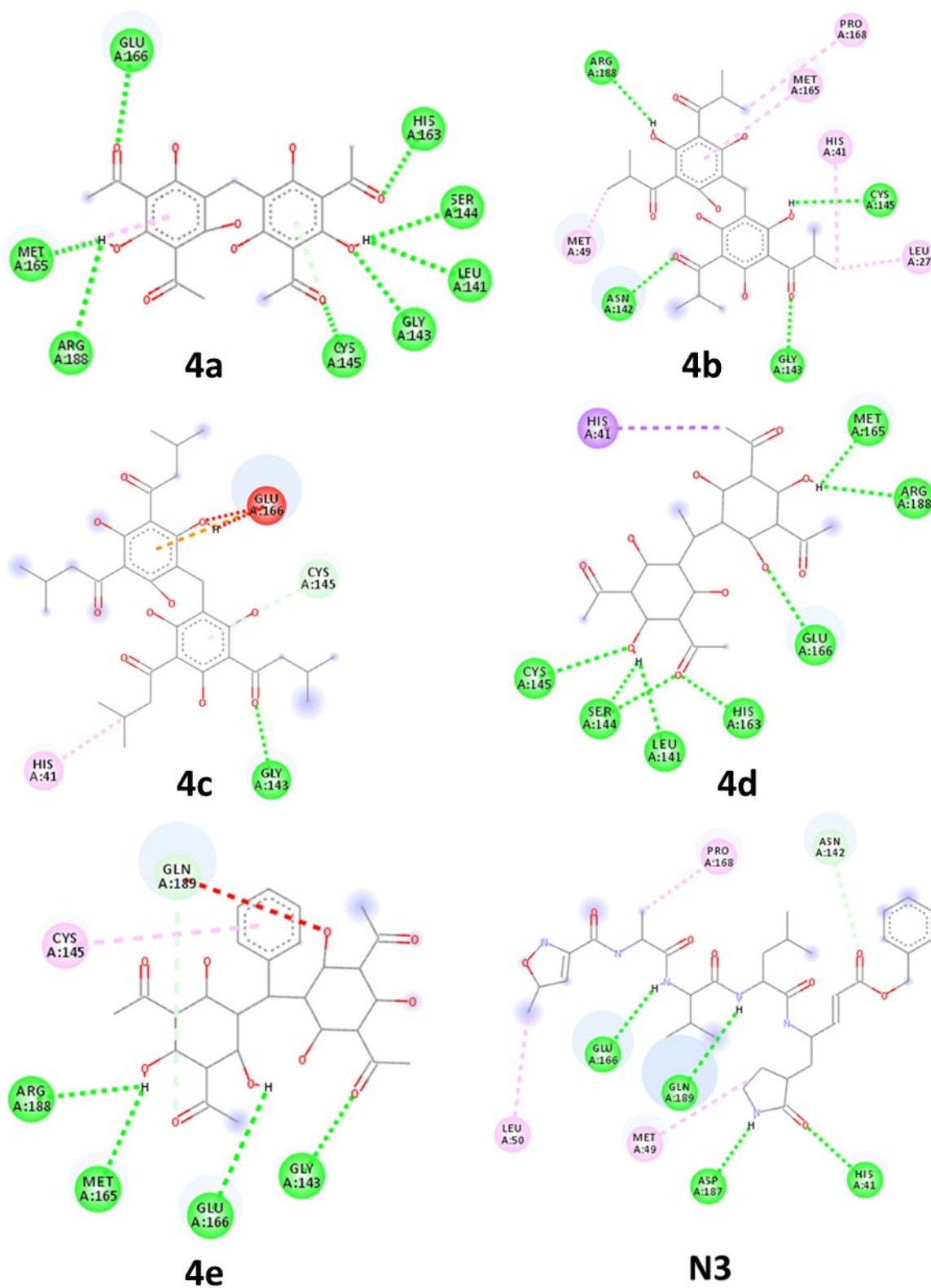


Figure S3. Surface representation showing the interaction of compound **4a-e**, N3, Chloroquine and Remdesivir at the substrate-binding region of 3CL^{PRO} of SARS-CoV-2 (**A**),

S7: 2D STRUCTURE DEMONSTRATING THE BINDING INTERACTIONS



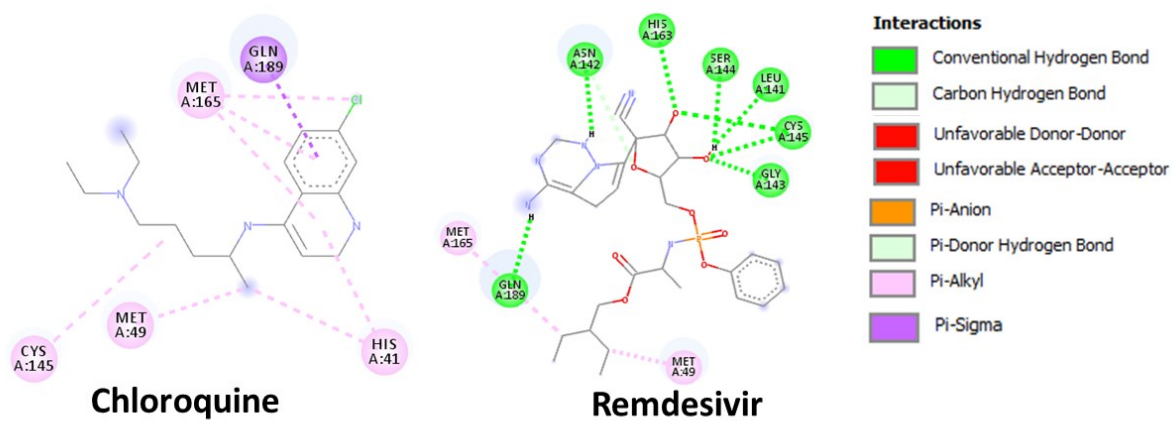
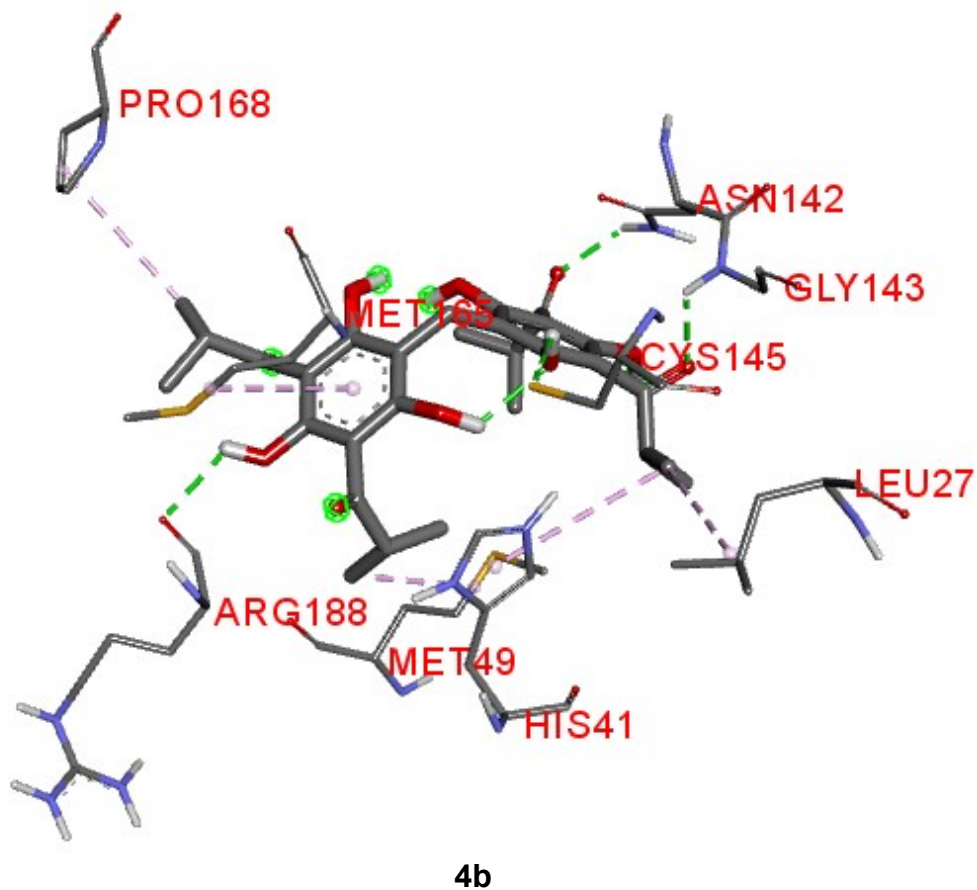
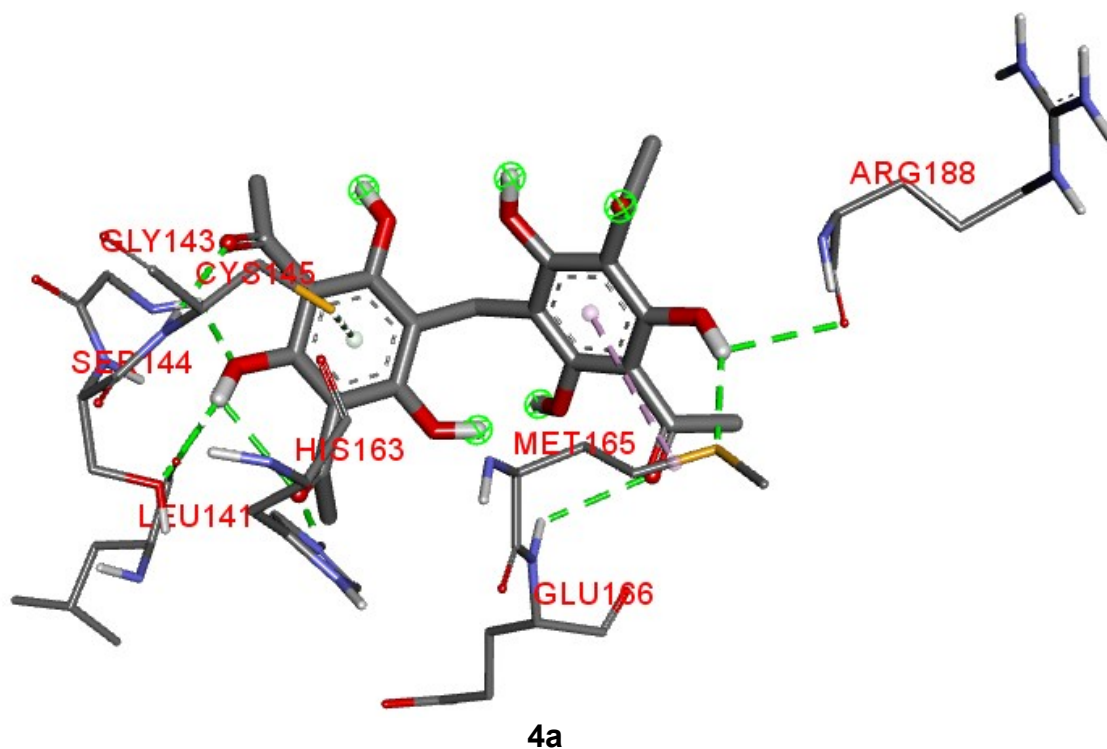
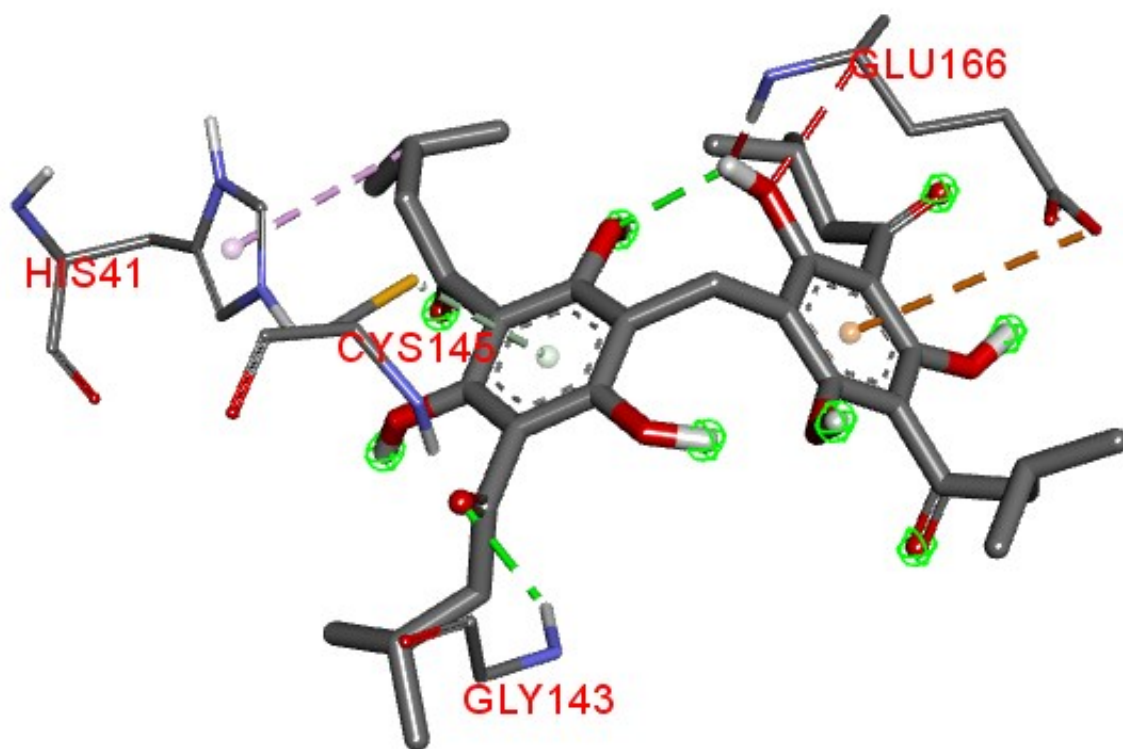


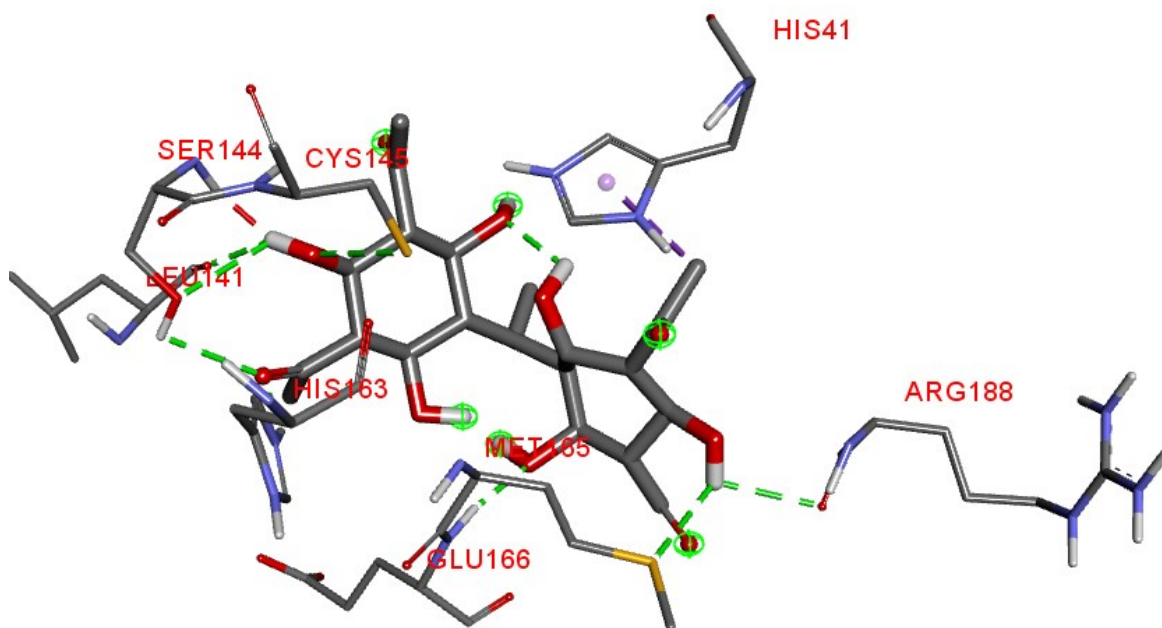
Figure S4. A 2D structure demonstrating the binding interactions between synthesised compounds along with N3, Chloroquine, and Remdesivir against 3CL^{pro}.

S8: 3D REPRESENTATION OF COMPOUND 4A-D DOCKED INTO THE ACTIVE POCKET OF 3CL^{pro}

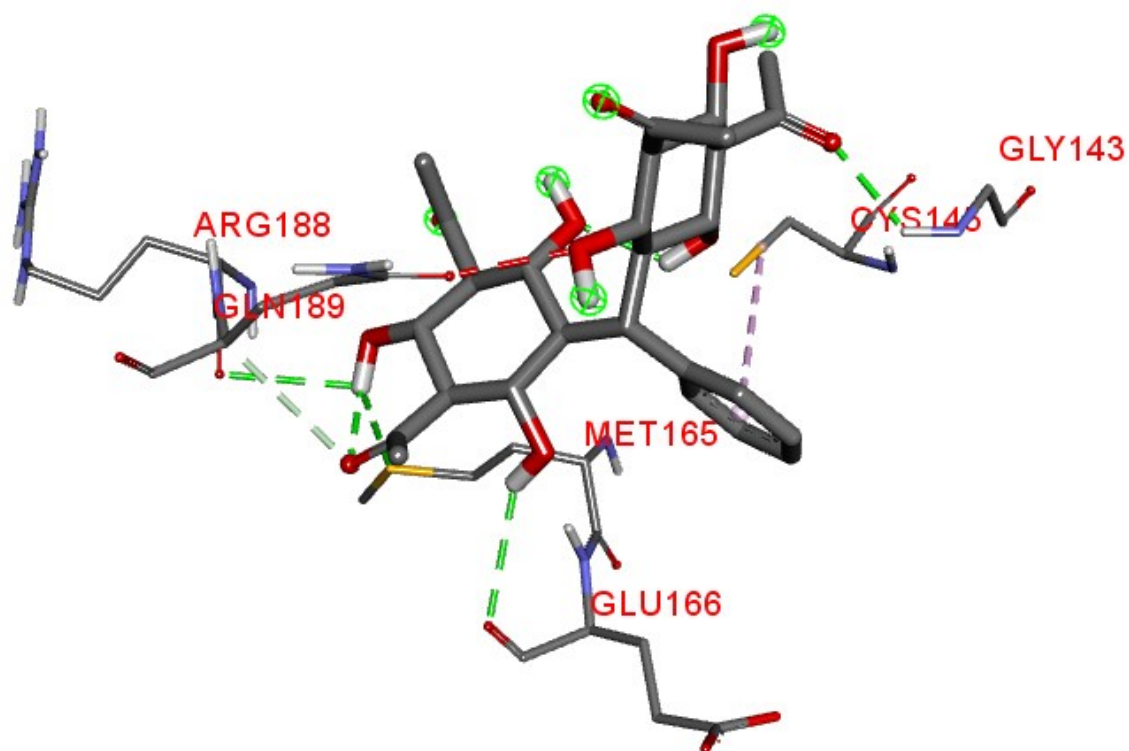




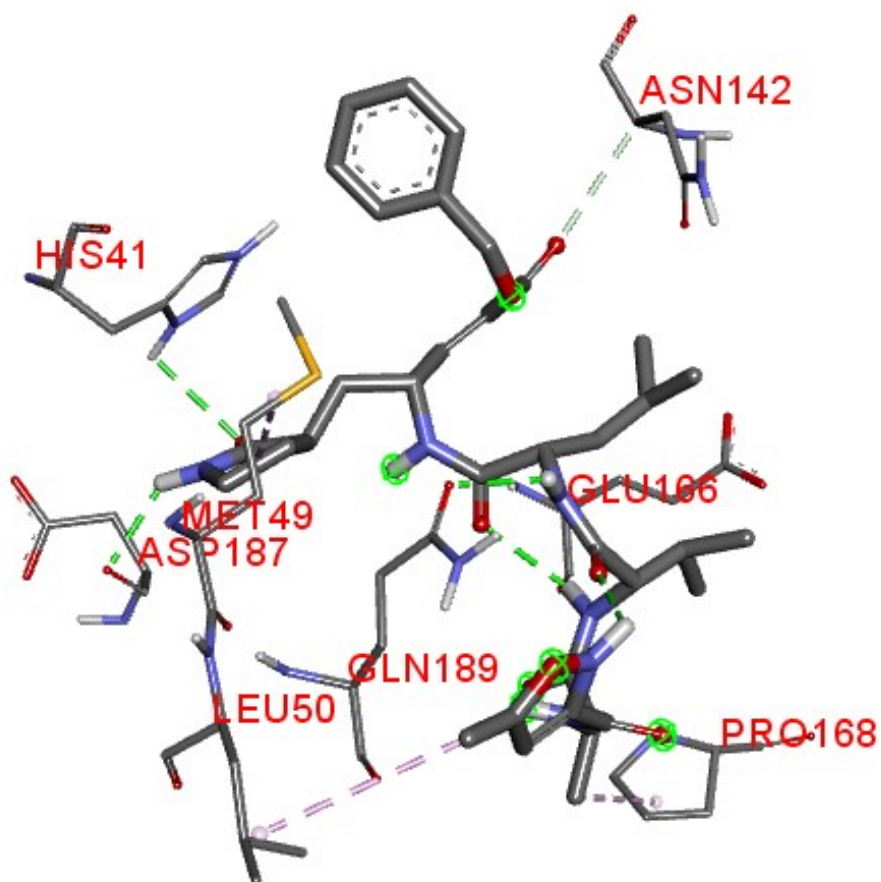
4c



4d



4e



N3

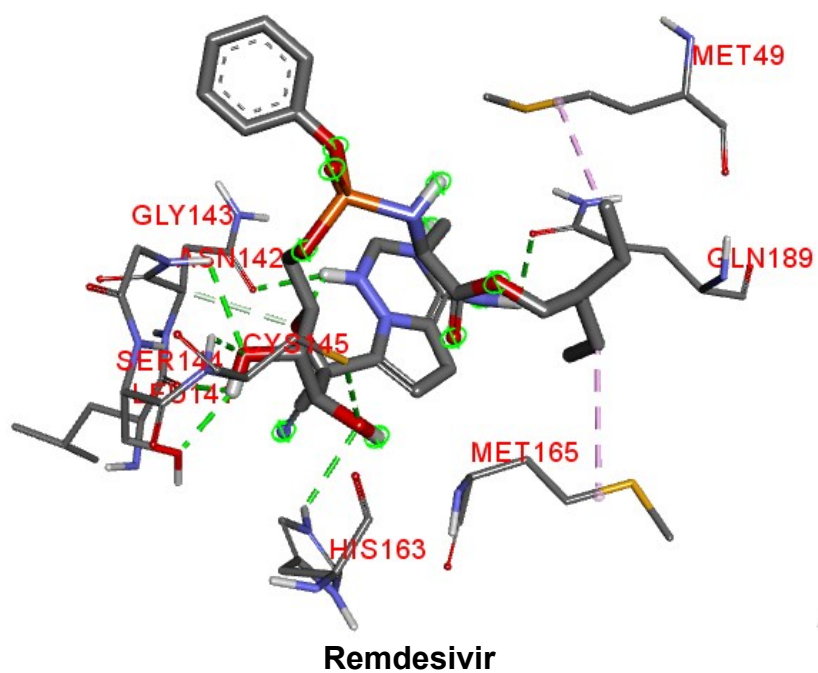
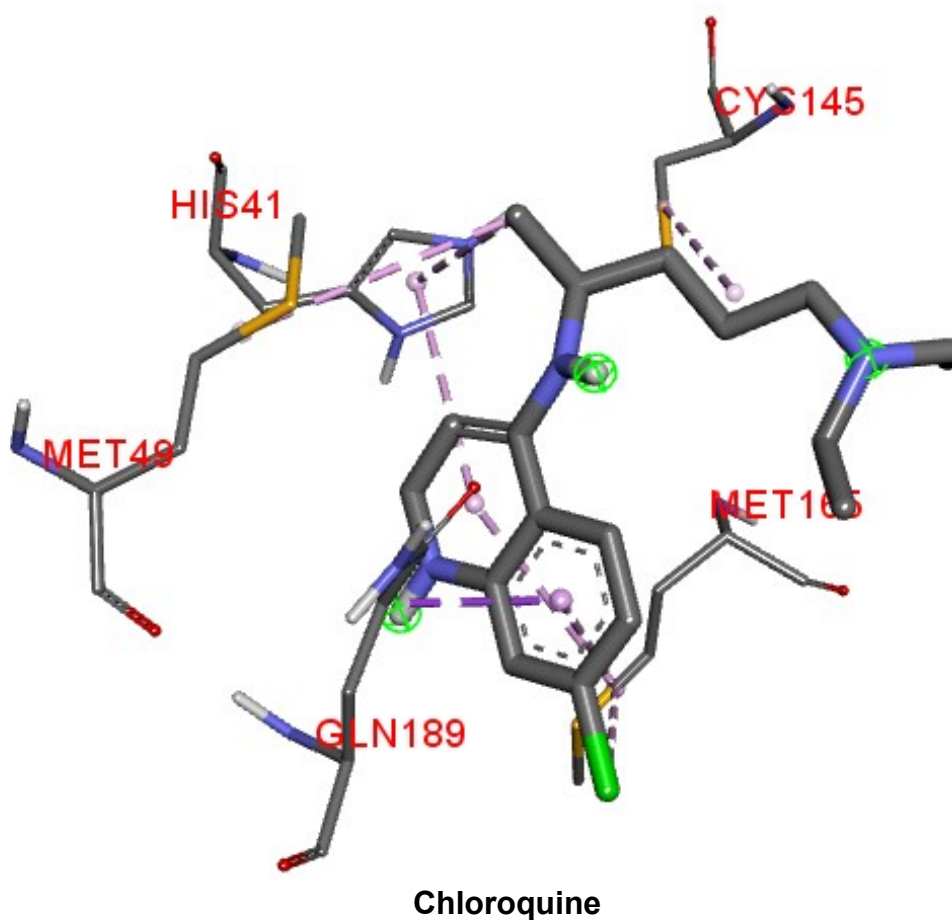


Figure S5. 3D representation of compound **4a-d** docked into the active pocket of 3CL^{pro}

S9: PHARMACOKINETIC PARAMETERS

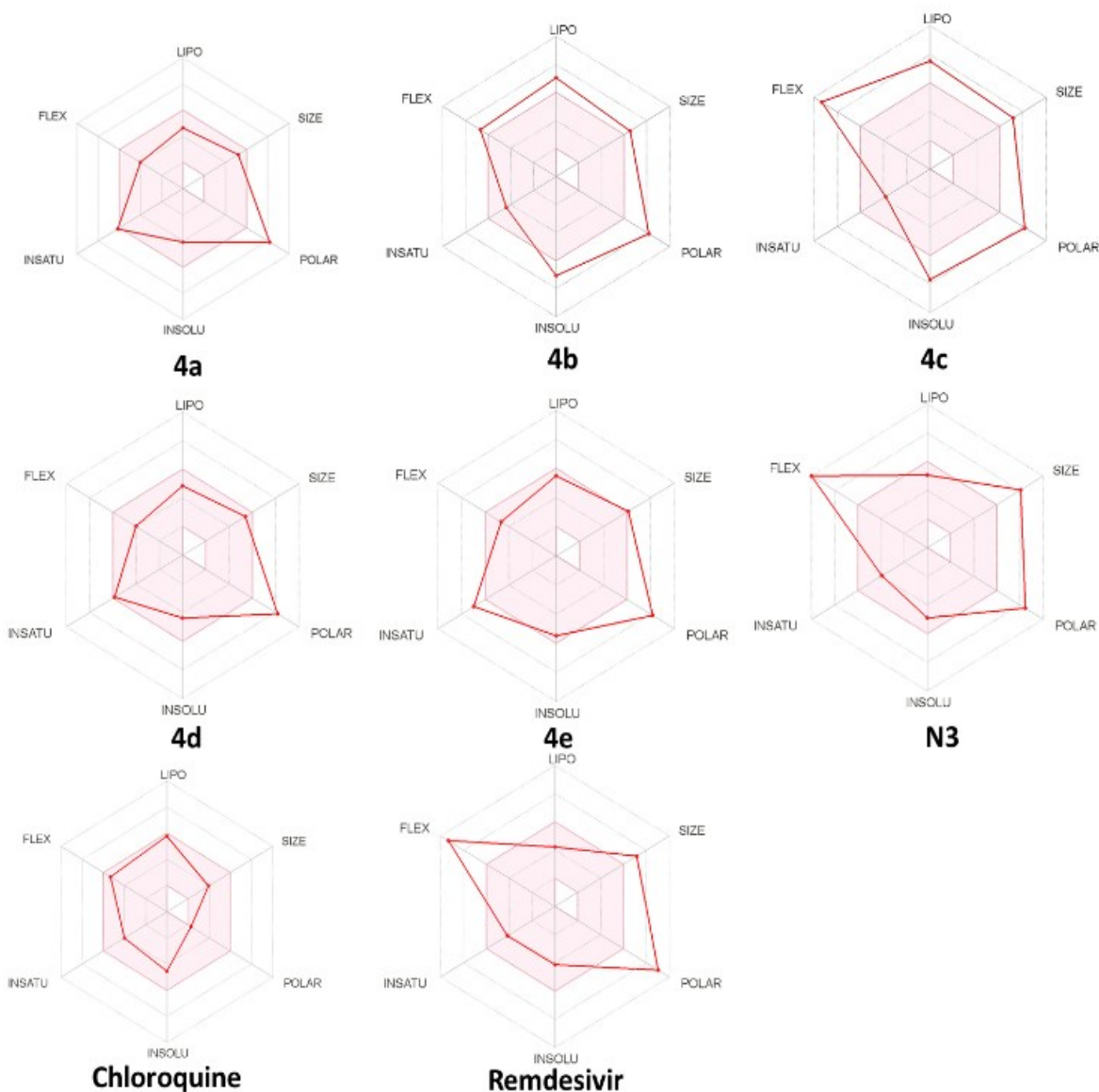


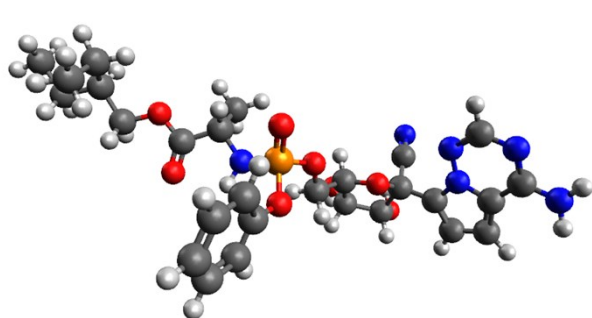
Figure S6. The analysis of pharmacokinetic parameters of the 5 synthesised compounds along with N3, Chloroquine, and Remdesivir using SwissADME server (<http://www.swissadme.ch/>). The colour space is the suitable physiochemical space for oral bioavailability. LIPO Lipophilicity: $-0.7 < XLOGP3 < 5.0$. SIZE: $150 \text{g/mol} < MW < 500 \text{g/mol}$. POLAR (Polarity): $20 \text{\AA}^2 < TPSA < 130 \text{\AA}^2$. INSOLU (insolubility): $0 < \text{Log S (ESOL)} < 6$. INSATU (insaturation): $0.25 < \text{Fraction Csp3} < 1$. FLEX (Flexibility): $0 < \text{Num. rotatable bonds} < 9$.

S10: PART OF THE PREDICTED ANTIVIRAL ACTIVITY SPECTRA

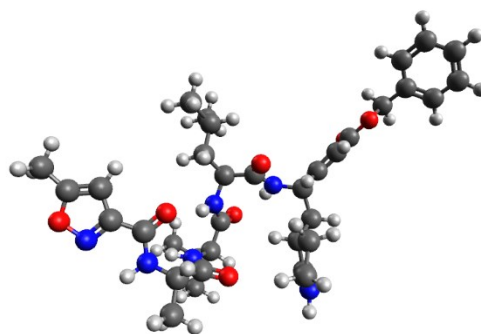
Table S2. Part of the predicted antiviral activity spectra for the synthesised compounds **4a-e** along with N3, Chloroquine, and Remdesivir

Ligands	4a		4b		4c		4d		4e		N3		Chloroquine		Remdesivir	
	Pa	Pi	Pa	Pi	Pa	Pi	Pa	Pi	Pa	Pi	Pa	Pi	Pa	Pi	Pa	Pi
Antiviral (Adenovirus)	0.341	0.059	0.342	0.059	0.352	0.053	0.314	0.078	0.289	0.099	-	-	-	-	-	-
Antiviral (CMV)	0.281	0.042	0.277	0.046	0.265	0.058	0.236	0.098	0.228	0.114	-	-	-	-	-	-
Antiviral (HIV)	0.229	0.020	0.249	0.016	0.262	0.013	0.205	0.028	0.554	0.004	0.121	0.113	-	-	0.272	0.012
Antiviral (Herpes)	0.303	0.090	0.311	0.084	0.335	0.068	0.251	0.129	0.221	0.155	-	-	-	-	0.466	0.015
Antiviral (Herpesvirus 3)	0.011	0.008	-	-	-	-	-	-	-	-	-	-	-	-	-	-
Antiviral (Influenza A)	0.243	0.111	0.217	0.172	0.232	0.133	0.235	0.126	0.225	0.150	0.234	0.129	-	-	0.216	0.174
Antiviral (Influenza)	0.317	0.079	0.242	0.141	0.230	0.155	0.412	0.043	0.259	0.122	-	-	-	-	-	-
Antiviral (Parainfluenza)	0.042	0.032	0.047	0.025	0.040	0.034	0.039	0.037	-	-	-	-	-	-	-	-
Antiviral (Picornavirus)	0.429	0.088	0.425	0.091	0.388	0.120	0.482	0.059	0.347	0.162	-	-	-	-	-	-
Antiviral (Poxvirus)	0.240	0.105	0.192	0.171	-	-	0.216	0.135	0.189	0.176	-	-	-	-	-	-
Antiviral (Rhinovirus)	0.453	0.047	0.385	0.108	0.426	0.066	0.427	0.065	0.417	0.074	0.337	0.181	-	-	-	-
Antiviral (Hepatitis C)	-	-	0.104	0.054	0.116	0.039	-	-	-	-	0.119	0.036	-	-	0.509	0.003
Antiviral (Hepatitis)	-	-	0.110	0.086	0.125	0.062	-	-	-	-	0.136	0.047	-	-	0.544	0.003
Antiviral (Hepatitis B)	-	-	-	-	-	-	-	-	-	-	-	-	-	-	0.344	0.024

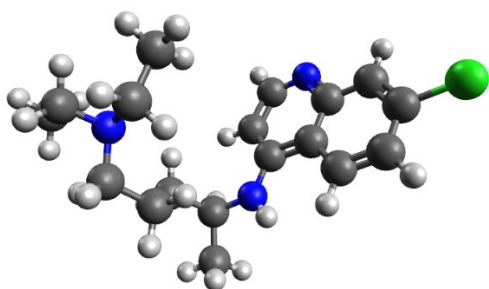
S11: DFT GEOMETRY OPTIMISATION



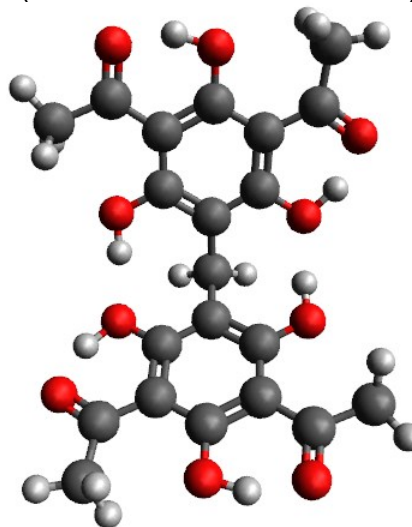
Chloroquine
CQ_TD_chelpg B3LYP/6-31+G(d,p)
E=-2321.7550354 H TD DFT
(-1456923.59220691 kcal/mol)
E=-2321.7551045 H DFT
(-1456923.6355678 kcal/mol)



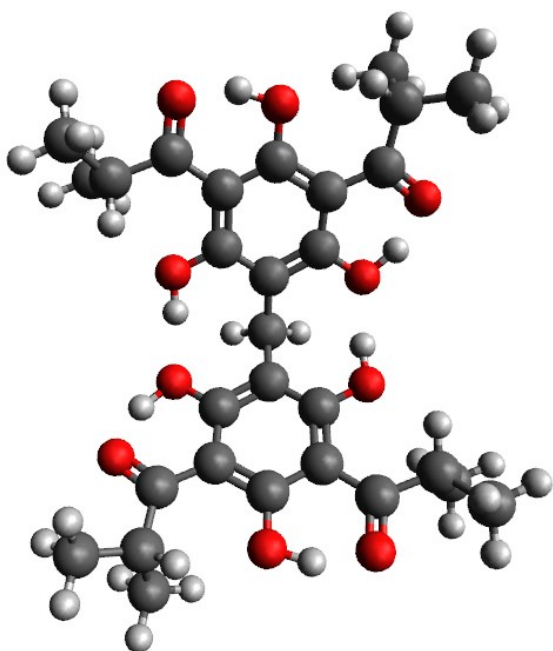
N3
N3_TD_chelpg B3LYP/6-31+G(d,p)
E=-2293.1680254 TD DFT
(-1438984.968767 kcal/mol)
E=-2293.1680945 DFT
(-1438985.01212796 kcal/mol)



Remdesivir
RDS_TD_chelpg B3LYP/6-31+G(d,p)
E=-1326.1005639 TD DFT
(-832140.845062037 kcal/mol)
E= -1326.1006115 DFT
(-832140.874931494 kcal/mol)

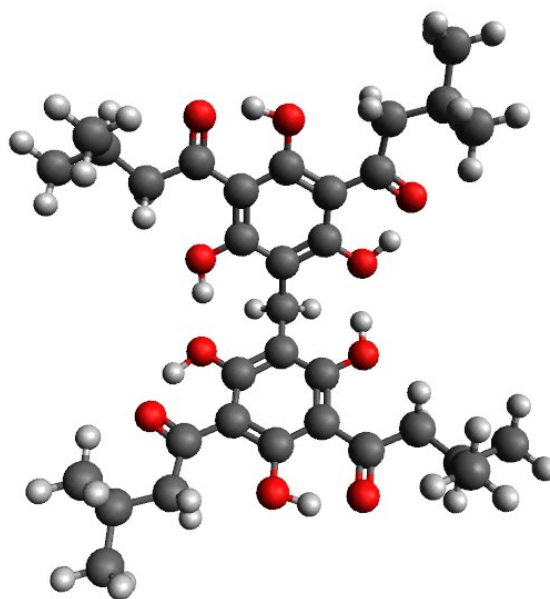


4a
4a_TD_chelpg B3LYP/6-31+G(d,p)
E= -1564.7022105 TD DFT
(-981865.670795457 kcal/mol)
E= -1564.7021775 DFT
(-981865.65008764 kcal/mol)



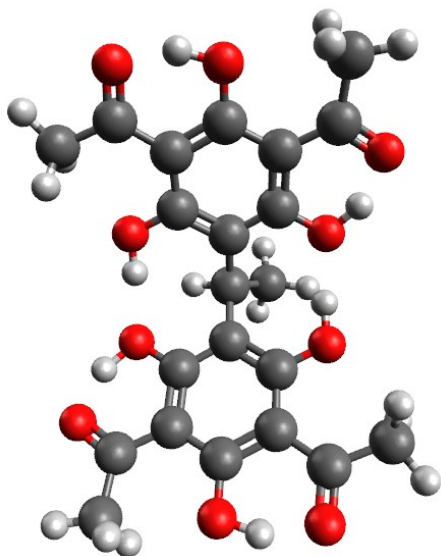
4b

4b_TD_chelpg B3LYP/6-31+G(d,p)
 E= -1879.2276558 TD DFT
 (-1179233.40969131 kcal/mol)
 E= -1879.2276558 DFT
 (-1179233.40969131 kcal/mol)



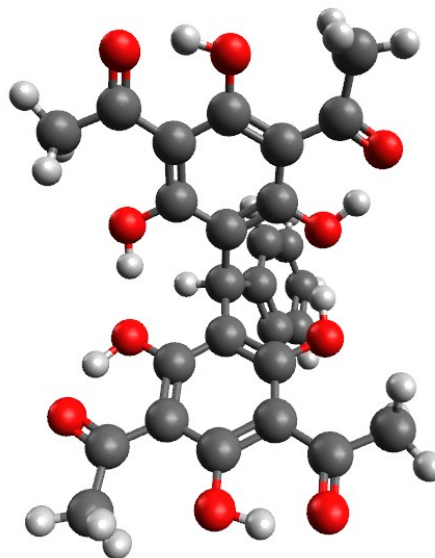
4c

4c_TD_chelpg B3LYP/6-31+G(d,p)
 E= -2036.5032204 TD DFT
 (-1277925.33758624 kcal/mol)
 E= -2036.5032204 DFT
 (-1277925.33758624 kcal/mol)



4d

4d_TD_chelpg B3LYP/6-31+G(d,p)
 E= -1604.007756 TD DFT
 (-1006530.2782456 kcal/mol)
 E= -1604.0077461 DFT
 (-1006530.27203325 kcal/mol)

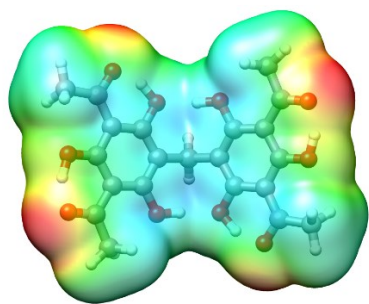


4e

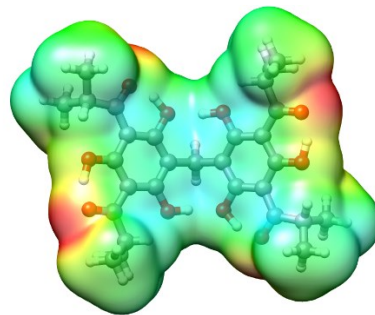
4e_TD_chelpg B3LYP/6-31+G(d,p)
 E= -1795.7512622 TD DFT
 (-1126851.1706636 kcal/mol)
 E= -1795.7512666 DFT
 (-1126851.17342461 kcal/mol)

Figure S7. DFT geometry optimisation of compound **4a-d**

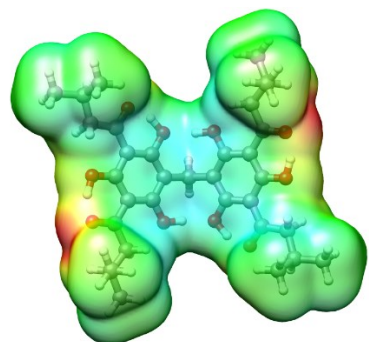
S12: MOLECULAR ELECTROSTATIC POTENTIAL (MEP) MAPS



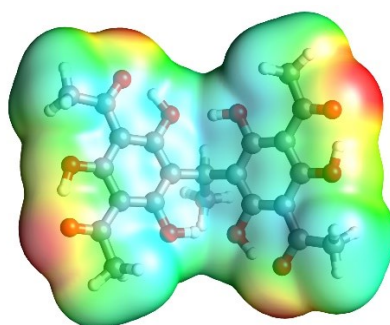
4a



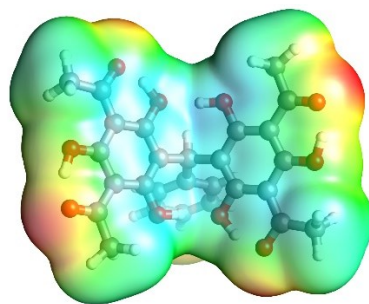
4b



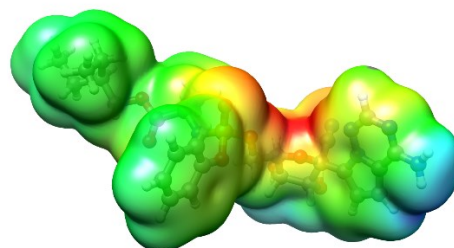
4c



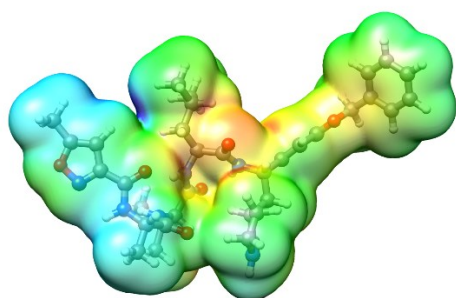
4d



4e



Chloroquine



Remdesivir

N3

Figure S8. MEP maps of compound **4a-d**

S13: CALCULATED QUANTUM CHEMICAL PARAMETERS

Table S3. Calculated quantum chemical parameters for the synthesised compounds **4a-e** along with N3, Chloroquine, and Remdesivir

Chemical reactivity indices	4a	4b	4c	4d	4e	N3	Chloroquine	Remdesivir
E_{HOMO} (eV)	-	-	-	-	-	-	-6.24	-5.87
E_{LUMO} (eV)	6.63	6.54	6.55	6.56	6.58	6.50	-1.41	-1.50
Energy gap (ΔE) = ($E_{LUMO} - E_{HOMO}$) (eV)	2.25	2.21	2.16	2.25	2.27	2.23	4.84	4.37
Ionisation potential (I) = $-E_{HOMO}$ (eV)	4.38	4.33	4.39	4.31	4.31	4.27	6.24	5.87
Electron affinity (A) = $-E_{LUMO}$ (eV)	6.63	6.54	6.55	6.56	6.58	6.50	6.24	5.87
Electronegativity (χ) = $-(E_{HOMO} + E_{LUMO})/2$ (eV)	2.25	2.21	2.16	2.25	2.27	2.23	1.41	1.50
Chemical potential (μ) = $1/2 (E_{LUMO} + E_{HOMO})$ (eV)	2.19	2.17	2.19	2.16	2.15	2.14	2.42	2.19
Global hardness (η) = $1/2 (E_{LUMO} - E_{HOMO})$ (eV)	-	-	-	-	-	-	-3.83	-3.68
Softness chemical softness (S) = $1/2 \eta$ (eV) ⁻¹	4.44	4.38	4.35	4.40	4.42	4.36	2.42	2.19
Electrophilicity index electrophilicity index (ω) = $\mu^2/2 \eta$ (ψ)	2.19	2.17	2.19	2.16	2.15	2.14	1.21	1.09
	1.10	1.08	1.10	1.08	1.08	1.07	3.02	3.10

S14: COORDINATES OF THE OPTIMISED STRUCTURES

Chloroquine DFT TD chelpg

P						
O	1	B1				
O	2	B2	1	A1		
O	3	B3	2	A2	1	D1 0
O	1	B4	2	A3	4	D2 0
O	1	B5	5	A4	2	D3 0
O	1	B6	5	A5	6	D4 0
O	1	B7	7	A6	5	D5 0
O	8	B8	1	A7	7	D6 0
N	2	B9	1	A8	7	D7 0
N	10	B10	2	A9	1	D8 0
N	1	B11	7	A10	5	D9 0
N	3	B12	2	A11	1	D10 0
N	11	B13	10	A12	2	D11 0
N	14	B14	11	A13	10	D12 0
C	2	B15	1	A14	7	D13 0
C	3	B16	2	A15	1	D14 0
C	4	B17	3	A16	2	D15 0
C	2	B18	1	A17	7	D16 0
C	10	B19	2	A18	1	D17 0
C	5	B20	1	A19	7	D18 0
C	13	B21	3	A20	2	D19 0
C	20	B22	10	A21	2	D20 0
C	10	B23	2	A22	1	D21 0
C	24	B24	10	A23	2	D22 0
C	14	B25	11	A24	10	D23 0
C	12	B26	1	A25	7	D24 0
C	11	B27	10	A26	2	D25 0
C	8	B28	1	A27	7	D26 0
C	8	B29	1	A28	7	D27 0
C	29	B30	8	A29	1	D28 0
C	29	B31	8	A30	1	D29 0
C	9	B32	8	A31	1	D30 0
C	27	B33	12	A32	1	D31 0
C	6	B34	1	A33	7	D32 0
C	31	B35	29	A34	8	D33 0
C	32	B36	29	A35	8	D34 0
C	35	B37	6	A36	1	D35 0
C	35	B38	6	A37	1	D36 0
C	38	B39	35	A38	6	D37 0
C	39	B40	35	A39	6	D38 0
C	40	B41	38	A40	35	D39 0
H	17	B42	3	A41	2	D40 0
H	18	B43	4	A42	3	D41 0
H	19	B44	2	A43	1	D42 0
H	21	B45	5	A44	1	D43 0
H	21	B46	5	A45	1	D44 0
H	3	B47	2	A46	1	D45 0

H	23	B48	20	A47	10	D46	0
H	4	B49	3	A48	2	D47	0
H	25	B50	24	A49	10	D48	0
H	12	B51	1	A50	7	D49	0
H	27	B52	12	A51	1	D50	0
H	29	B53	8	A52	1	D51	0
H	28	B54	11	A53	10	D52	0
H	30	B55	8	A54	1	D53	0
H	30	B56	8	A55	1	D54	0
H	32	B57	29	A56	8	D55	0
H	31	B58	29	A57	8	D56	0
H	32	B59	29	A58	8	D57	0
H	31	B60	29	A59	8	D58	0
H	15	B61	14	A60	11	D59	0
H	15	B62	14	A61	11	D60	0
H	34	B63	27	A62	12	D61	0
H	34	B64	27	A63	12	D62	0
H	34	B65	27	A64	12	D63	0
H	37	B66	32	A65	29	D64	0
H	36	B67	31	A66	29	D65	0
H	37	B68	32	A67	29	D66	0
H	37	B69	32	A68	29	D67	0
H	36	B70	31	A69	29	D68	0
H	36	B71	31	A70	29	D69	0
H	38	B72	35	A71	6	D70	0
H	39	B73	35	A72	6	D71	0
H	40	B74	38	A73	35	D72	0
H	41	B75	39	A74	35	D73	0
H	42	B76	40	A75	38	D74	0

B1	4.46157339
B2	3.61097956
B3	2.73586475
B4	1.60906964
B5	1.63495435
B6	1.48112331
B7	4.99001303
B8	2.26148093
B9	3.02755247
B10	1.36190702
B11	1.66603993
B12	3.44101461
B13	2.40697271
B14	2.30624533
B15	1.42674806
B16	1.40193278
B17	1.41891439
B18	1.43853854
B19	1.36901011
B20	1.43422509
B21	1.16160140
B22	1.39738935
B23	1.39422091
B24	1.39624714
B25	1.32568118

B26	1.46545151
B27	1.31050921
B28	2.41965901
B29	1.45477877
B30	1.54675022
B31	1.54504098
B32	1.21783630
B33	1.54013769
B34	1.39755477
B35	1.53533564
B36	1.53358628
B37	1.39648113
B38	1.39516713
B39	1.39772155
B40	1.39667350
B41	1.39767079
B42	1.09134448
B43	1.09837694
B44	1.09816061
B45	1.09566614
B46	1.09728520
B47	0.97089589
B48	1.08011855
B49	0.96717367
B50	1.08072375
B51	1.01624882
B52	1.09649108
B53	1.10022816
B54	1.08512619
B55	1.09451677
B56	1.09372322
B57	1.09828958
B58	1.09624537
B59	1.09809866
B60	1.09564383
B61	1.00688174
B62	1.00953104
B63	1.09330627
B64	1.09317757
B65	1.09472690
B66	1.09454927
B67	1.09554713
B68	1.09497142
B69	1.09643879
B70	1.09551956
B71	1.09501955
B72	1.08414797
B73	1.08525118
B74	1.08610217
B75	1.08601914
B76	1.08571822
A1	137.23086642
A2	64.27641946
A3	10.72155051
A4	99.07527313

A5	111.27422239
A6	86.65952948
A7	57.99019077
A8	136.07453735
A9	80.40796965
A10	114.37559394
A11	58.64986320
A12	87.13286290
A13	123.01987182
A14	166.66344859
A15	19.46938664
A16	65.01302021
A17	59.63276026
A18	50.53645318
A19	122.47069913
A20	50.09954737
A21	107.16729078
A22	146.93665276
A23	107.58466657
A24	91.81997787
A25	121.66961911
A26	113.27877173
A27	164.68555627
A28	146.13645240
A29	92.57311214
A30	143.15191666
A31	29.25899489
A32	112.52702495
A33	124.08663988
A34	116.07964017
A35	114.32932552
A36	121.78542339
A37	116.77907517
A38	118.65582828
A39	119.25809358
A40	120.80632844
A41	106.38410012
A42	110.15810827
A43	109.86569236
A44	109.38517859
A45	109.73346315
A46	92.49206439
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N3 TD chelpg

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Remdesivir TD chelpg

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D26	93.81314374
D27	72.48000363
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4a TD chelpg

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O	2	1.34482067	1	117.39704942	6	178.96158090 0
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4b TD chelpg

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C	1	B5	2	A4	3	D3	0
O	4	B6	3	A5	2	D4	0
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C	5	B10	4	A9	3	D8	0
C	3	B11	2	A10	1	D9	0
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H	68	B69	27	A68	26	D67	0
H	68	B70	27	A69	26	D68	0
C	27	B71	26	A70	23	D69	0
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4c TD chelpg

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O	4	1.32586431	3	118.41611506	2	-177.20712410 0
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O	6	1.34168243	1	120.62325717	2	-179.55318083 0
O	2	1.34580618	1	117.17543829	6	179.12289730 0
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C	24	1.42943772	22	123.52911525	21	174.81267622	0
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C	25	1.47449472	23	124.38626278	22	179.83608815	0
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H	33	1.09548161	28	109.03908177	25	48.42270559	0
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4d TD chelpg

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4e TD chelpg

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O	2	B9	1	A8	6
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