

# Repurposing of Some Natural Product isolates as SARS-CoV-2 Main Protease Inhibitors via In vitro Cell Free and cell based anti-viral Assessments and Molecular Modeling Approaches

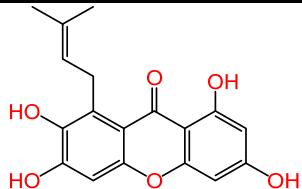
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## Supplementary Data

**Table S1.** The inactive natural isolates with high affinity to the viral protease (SARS-CoV-2 M<sup>PRO</sup>).

No.	Compound Name	Structure	Glide G-Score
9	1-Hydroxy-3,4-dihydronorharmane		-7.43
10	Aromadendrin-8-C- β-D-glucopyranoside		-7.394
11	Butyrolactone I		-7.316
12	Terrenolide S		-7.304
13	Ingenine C		-7.286

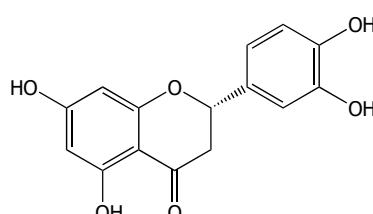
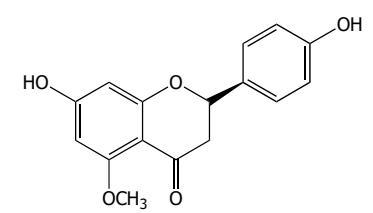
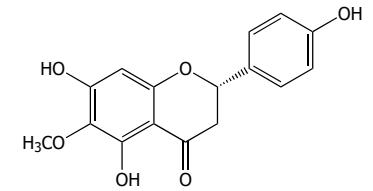
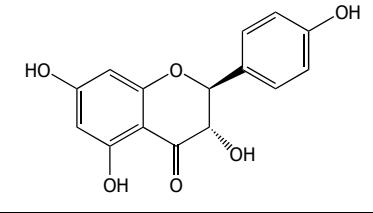
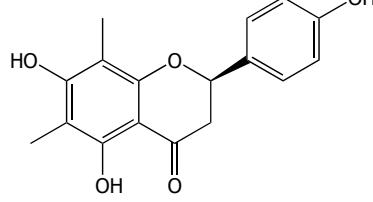
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14	1,3,6,7-Tetrahydroxy-8-prenylxanthone		-7.284
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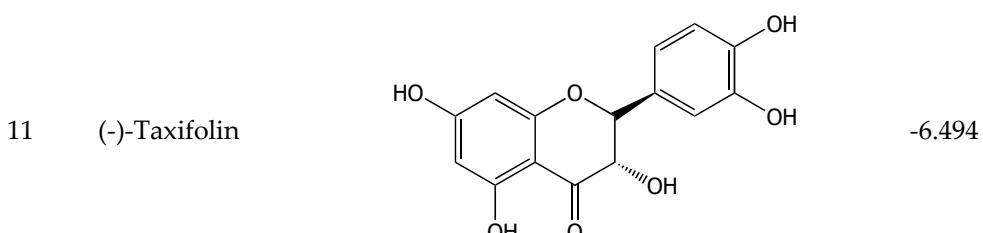
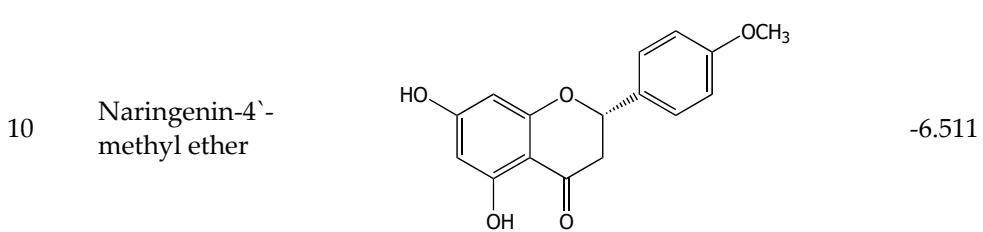
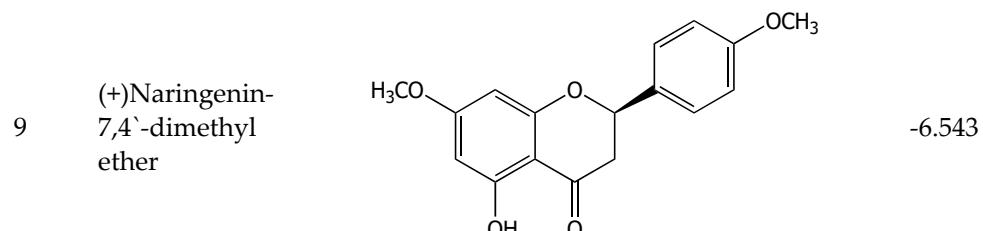
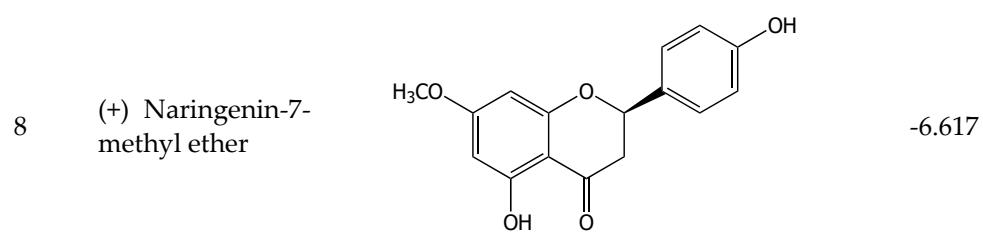
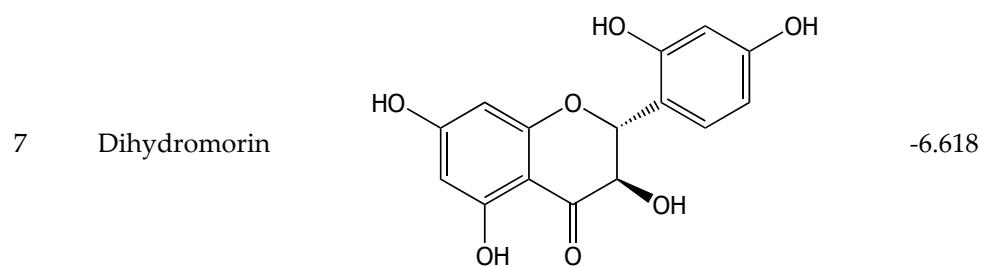
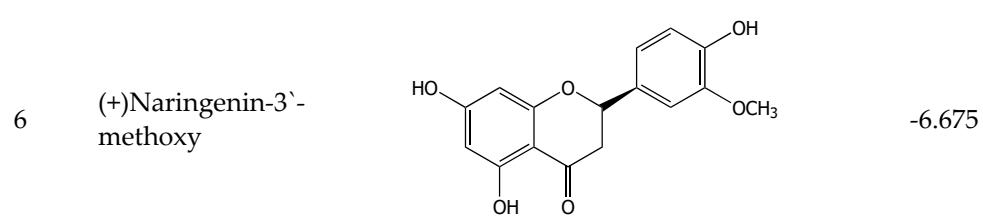
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**Table S2.** Similar compounds to naringenin that searched for their activity against the viral protease (SARS-CoV-2 M<sup>PRO</sup>).

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No	Compound Name	Structure	G-score
1	(+)-Eriodictyol		-6.933
2	(+)-Naringenin-5-methyl ether		-6.854
3	Naringenin-6'-methyl ether		-6.816
4	(-) Aromadendrin		-6.754
5	(+)-6,8-dimethyl Naringenin		-6.719

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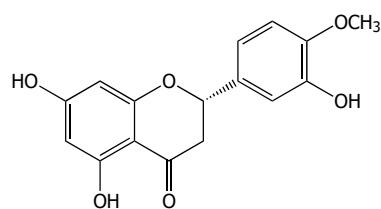


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12

(+)-Eriodictyol-4'-methyl ether



-5.783

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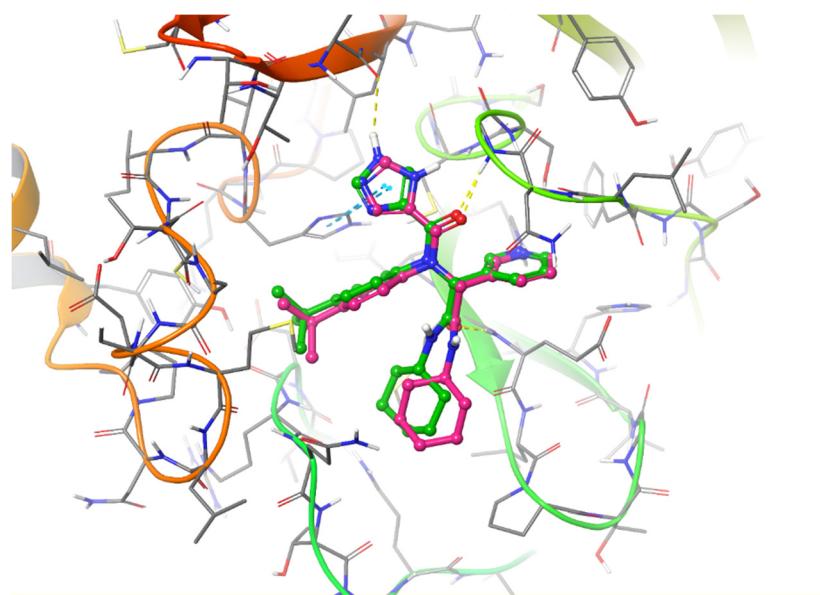
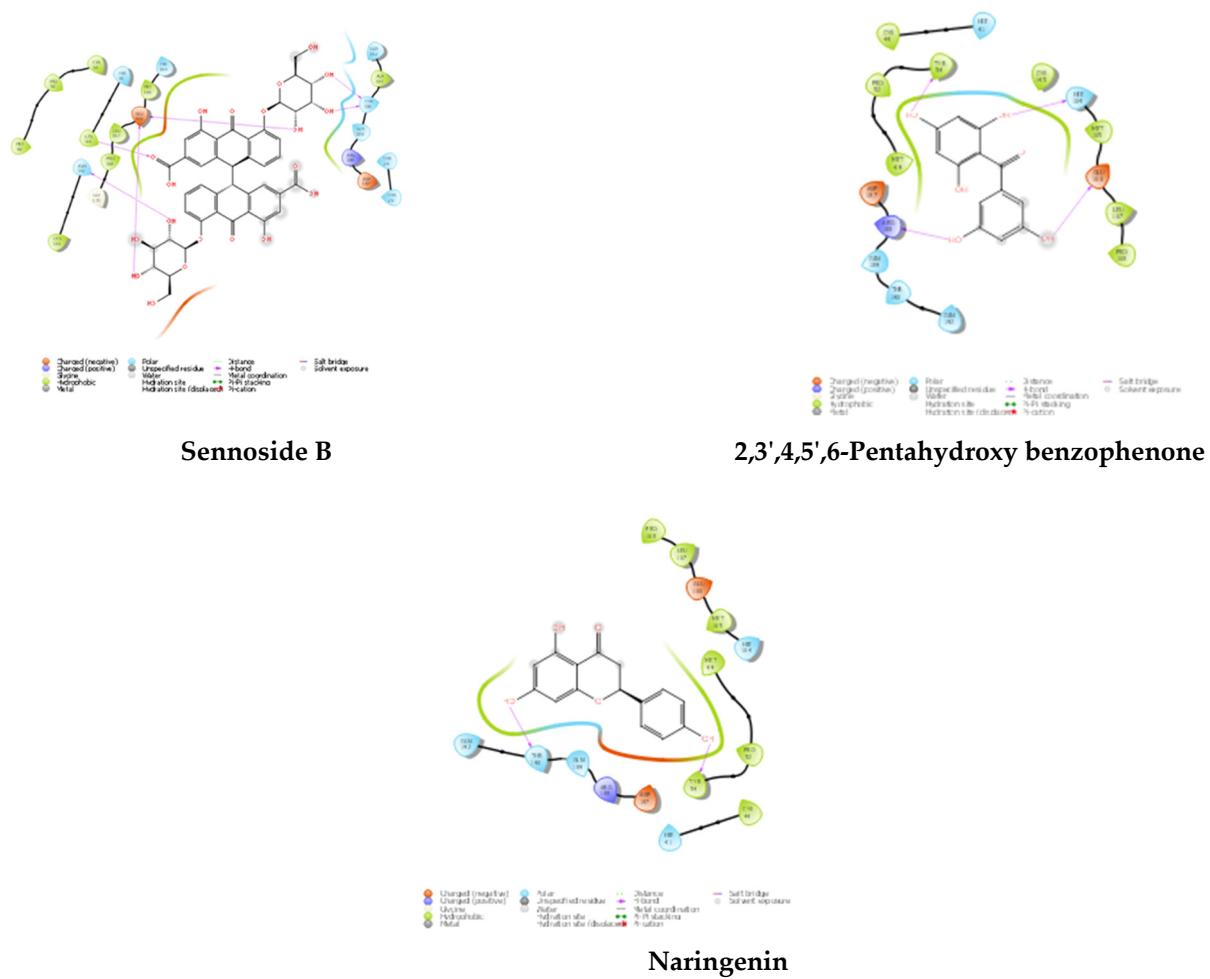
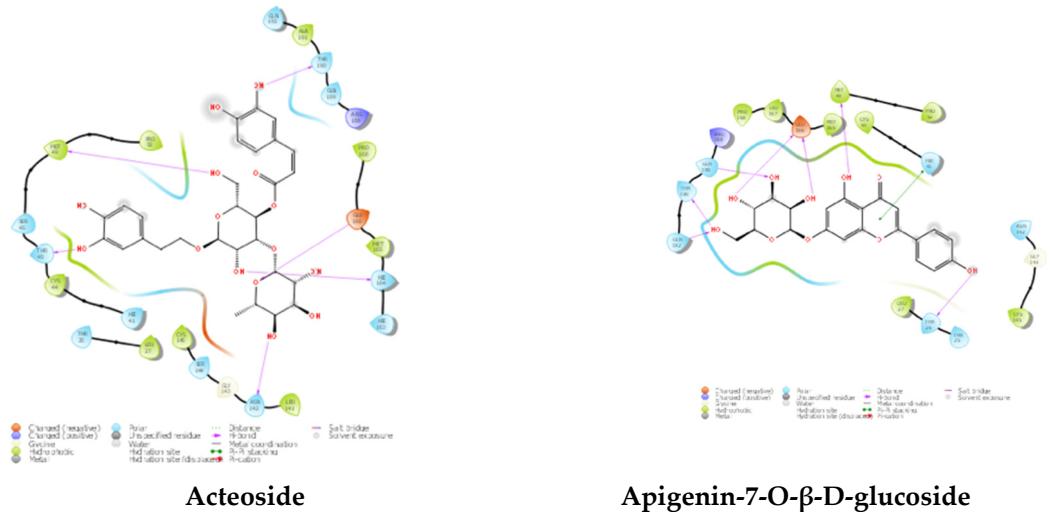
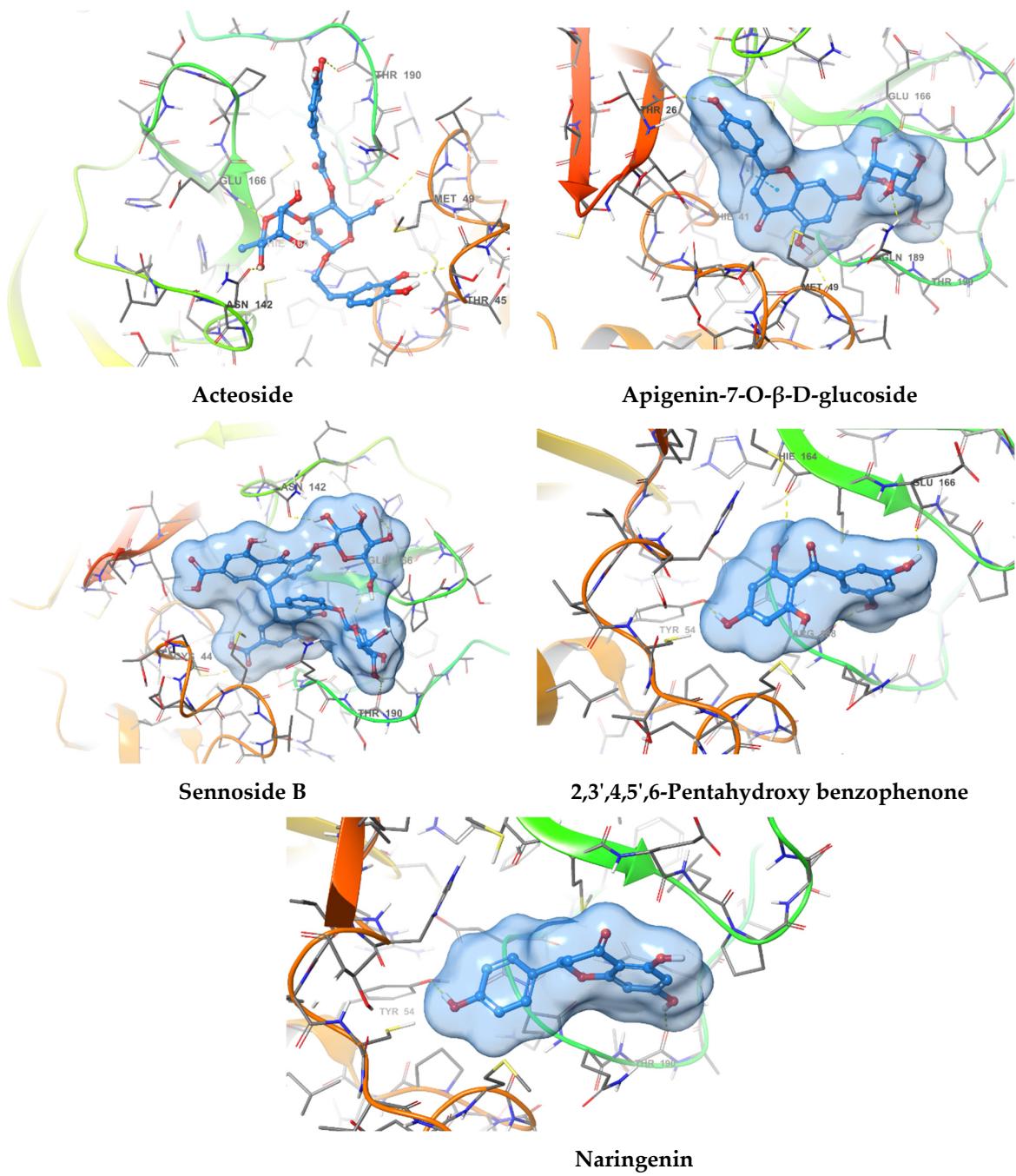


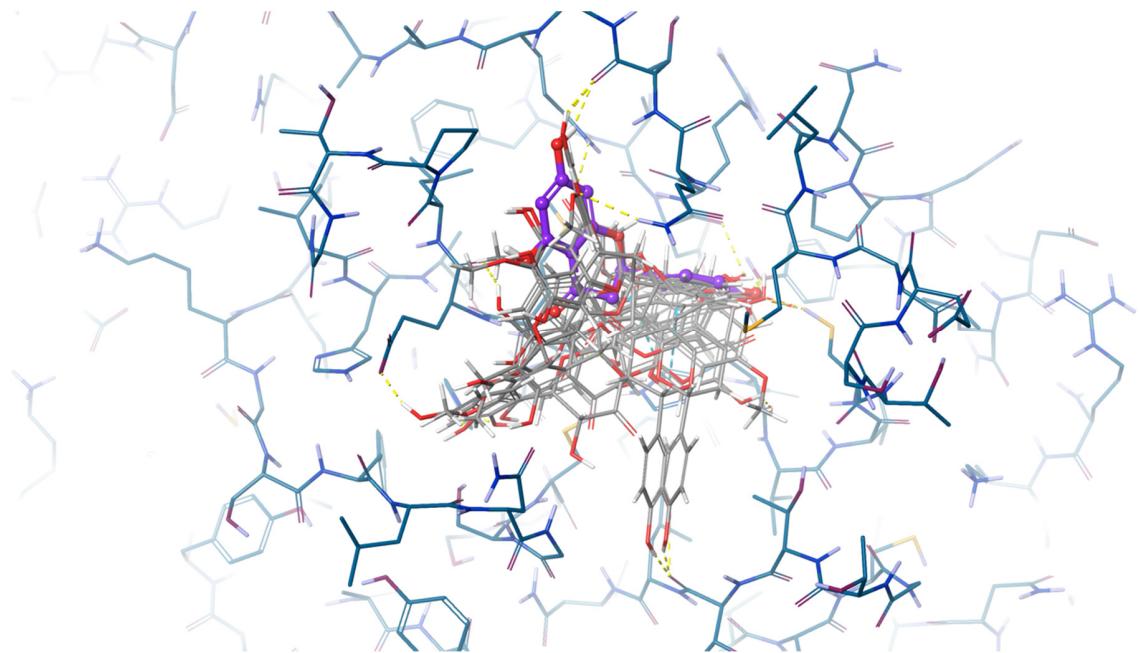
Figure S1. Superposition of the co-crystallized ligand (pink) and the redocked pose (green) with very small deviation and an RMSD value of 1.56 Å.



**Figure S2.** 2D Interaction diagrams of acteoside, apigenin-7-O-glucoside, sennoside B, 2,3',4,5',6-pentahydroxy benzophenone, and naringenin, respectively inside the Mpro active site in the crystal form.



**Figure S3.** 3D Interaction diagrams of acteoside, apigenin, sennoside B, 2,3',4,5',6-pentahydroxy benzophenone, and naringenin, respectively.



**Figure S4.** Overlay of naringenin analogues (grey) with naringenin (violet) in MPro active site