

Supplementary Tables

Antiviral phytochemicals identified in *Rhododendron arboreum* petals exhibited strong binding to SARS-CoV-2 M^{Pro} and Human ACE2 receptor

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Supplementary	Description
Table S1	Secondary metabolites of <i>R. arboreum</i> petals. A) Hot water extracts analysed using GC-MS. The retention time (RT), m/z product ion, and their match scores against the NIST17 and Fiehn13 library are presented. The area of the peaks among all the replicates are listed below (n=4). B) Secondary metabolites reported by Shrestha et al., 2016 using LC-MS.
Table S2	Assessment of ADME (absorption, distribution, metabolism and elimination) properties of <i>R. arboreum</i> phytochemicals for drug discovery and development.
Table S3	Structural information of SARS-CoV-2 Main protease (6LU7) and ACE2 host cell receptor (1R4L) along with their native ligands. Active sites of the target proteins were analysed by using Computed Atlas for Surface Topography of Proteins (CASTp). The amino acids residues found in the active sites pockets of SARS-CoV-2 Main protease (6LU7) and ACE2 host cell receptor (1R4L) are tabulated.
Table S4	Blind docking of the secondary metabolites identified in <i>R. arboreum</i> petals against target protein-SARS-CoV-2 Mpro (6LU7) and Human Angiotensin Converting Enzyme (1R4L). Glycyrrhizic acid was taken as a positive control.
Table S5	<i>Rhododendron arboreum</i> petal phytochemicals with reported anti-viral studies

Table S1. Secondary metabolites of *R. arboreum* petals. A) Hot water extracts analysed using GC-MS. The retention time (RT), m/z product ion, and their match scores against the NIST17 and Fiehn13 library are presented. The area of the peaks among all the replicates are listed below (n=4). B) Secondary metabolites reported by Shrestha 2016 using LC-MS.

A								
Sl.No	Metabolite	RT	Product Ion (m/z)	Match Factor	% Peak Area Replicate 1	% Peak Area Replicate 2	% Peak Area Replicate 3	% Peak Area Replicate 4
1	3-Hydroxybenzoic acid	15.88	267	71.66	0.006	0.011	0.006	0.010
2	Shikimic acid	18.31	204	95.19	2.804	1.219	3.072	1.015
3	Protocatechuic acid	18.49	370	88.97	0.085	0.071	0.052	0.094
4	Quinic acid	19.5	345	91.04	12.028	6.517	8.291	5.559
5	Coumaric acid	20.86	293	88.69	0.056	0.058	0.062	0.048
6	Caffeic acid	27.00	396	70.41	0.013	0.018	0.016	0.016
7	Epicatechin	42.68	368	90.13	2.839	1.933	3.997	2.092
8	Catechin	42.98	368	79.33	2.830	1.930	3.997	2.097
9	5-O-Coumaroyl-D-quinic acid	44.40	345	97.41	1.137	0.513	0.764	0.778
10	3-Caffeoyl-quinic acid	44.70	345	93.87	1.113	10.790	0.764	0.783
11	2,4-Quinolinediamine	45.50	370	71.31	0.009	0.005	0.020	0.010
12	5-O-Feruloylquinic acid	46.35	345	75.82	14.215	10.869	14.348	10.107
	Others - Sugars, Organic acids, Aminoacids ...etc	variable	variable	variable	62.87	66.07	64.61	77.39

B		
Sl. No.	Metabolite	Mass
13	Epicatechin gallate	441.0827
14	Quercetin	301.0354
15	Quercetin-O-pentoside	433.0776
16	Quercetin-O-rhamnoside	447.0933
17	Kaempferol-O-pentoside	417.0827
18	Kaempferol	286.23

Reference:

Shrestha, A. (2016). Phytochemical Analysis of Rhododendron Species (Doctoral dissertation, IRC-Library, Information Resource Center der Jacobs University Bremen).

Table S2. Assessment of ADME (absorption, distribution, metabolism and elimination) properties of R. arboreum phytochemicals for drug discovery and development.

Pubchem CID	14207924	7420	637542	9799386	689043	9064	1794427	72276	72	8742	6441280	6508	107905	5280863	14749097	5280343	5878729	5280459
Molecules	2,4-Quinoline diamine	3-Hydroxy benzoic acid	4-Coumaric acid	5-O-Feruloyl quinic acid	Caffeic acid	Catechin	3-Caffeoyl-quinic acid	Epicatechin	Proto catechuic acid	Shikimic acid	5-O-Coumaroyl-D-quinic acid	Quinic acid	Epicatechin gallate	Kaempferol	Kaempferol-O-pentoside	Quercetin	Quercetin-O-pentoside	Quercetin-O-rhamnoside
Canonical SMILES	<chem>Nc1cc(N)c2c(n1)cccc2</chem>	<chem>Oc1cccc(c1)C(=O)O</chem>	<chem>OC(=O)C=Cc1cc(c(cc1)O</chem>	<chem>COc1cc(C=CC(=O)OC2CC(O)(CC(C2O)O)C(=O)O)ccc1O</chem>	<chem>OC(=O)C=Cc1cc(c(c(c1)O)O</chem>	<chem>Oc1cc2OC(C(c3ccc(c(c3)O)C(Cc2c(c1)O)O</chem>	<chem>O=C(OC1CC(O)(CC(C1O)O)C(=O)O)C(=O)O)C=Cc1ccc(c(c1)O)O</chem>	<chem>O=C(c1ccc(c(c1)O[Si](C)(C)C)O[Si](C)(C)C)O[Si](C)(C)C</chem>	<chem>O=C(OC1CC(O)(CC(C1O)O)C(=O)O)C=Cc1ccc(cc1)O</chem>	<chem>OC1CC(=CC(C1O)O)C(=O)O</chem>	<chem>O=C(OC1CC(O)(CC(C1O)O)C(=O)O)C=Cc1ccc(cc1)O</chem>	<chem>OC1C(O)CC(C1O)(CC1O)(C(=O)O)C(=O)O</chem>	<chem>Oc1cc(O)c2c(c1)OC(C(C2)OC(=O)c1cc(O)c(c(c1)O)O)c1ccc(c(c1)O)O</chem>	<chem>Oc1ccc(cc1)c1oc2cc(O)cc(c2c(=O)c1O)O</chem>	<chem>Oc1ccc(cc1)c1oc2cc(O)cc(c2c(=O)c1O)O</chem>	<chem>Oc1cc(O)c2c(c1)oc(c(c2=O)O)c1ccc(c(c1)O)O</chem>	<chem>Oc1cc(O)c2c(c1)oc(c(c2=O)OC1OCC(C(C1O)O)O)c1ccc(c(c1)O)O</chem>	<chem>Oc1cc(O)c2c(c1)oc(c(c2=O)OC1OC(C(C1O)O)O)c1ccc(c(c1)O)O</chem>
Formula	C9H9N3	C7H6O3	C9H8O3	C17H20O9	C9H8O4	C15H14O6	C16H18O9	C15H14O6	C16H30O4Si3	C7H10O5	C16H18O8	C7H12O6	C22H18O10	C15H10O6	C20H18O10	C15H10O7	C20H18O11	C21H20O11
MW	159.19	138.12	164.16	368.34	180.16	290.27	354.31	290.27	370.66	174.15	338.31	192.17	442.37	286.24	418.35	302.24	434.35	448.38
#Heavy atoms	12	10	12	26	13	21	25	21	23	12	24	13	32	21	30	22	31	32
#Aromatic heavy atoms	10	6	6	6	6	12	6	12	6	0	6	0	18	16	16	16	16	16
Fraction Csp3	0	0	0	0.41	0	0.2	0.38	0.2	0.56	0.57	0.38	0.86	0.14	0	0.25	0	0.25	0.29
#Rotatable bonds	0	1	2	6	2	1	5	1	7	1	5	1	4	1	3	1	3	3
#H-bond acceptors	1	3	3	9	4	6	9	6	4	5	8	6	10	6	10	7	11	11
#H-bond donors	2	2	2	5	3	5	6	5	0	4	5	5	7	4	6	5	7	7
MR	50.55	35.42	45.13	87.97	47.16	74.33	83.5	74.33	103.15	38.43	81.48	40.11	110.04	76.01	102.17	78.03	104.19	109
TPSA	64.93	57.53	57.53	153.75	77.76	110.38	164.75	110.38	44.76	97.99	144.52	118.22	177.14	111.13	170.05	131.36	190.28	190.28
iLOGP	1.1	0.86	0.95	1.47	0.97	1.47	0.96	1.47	4.72	0.62	1.08	-0.12	1.31	1.7	1.21	1.63	1.61	1.27
XLOGP3	1.16	1.5	1.46	-0.1	1.15	0.36	-0.42	0.36	5.99	-1.72	-0.07	-2.37	1.53	1.9	0.78	1.54	0.43	0.86
WLOGP	1.42	1.09	1.38	-0.45	1.09	1.22	-0.75	1.22	5.11	-1.52	-0.46	-2.32	2.2	2.28	0.39	1.99	0.1	0.49
MLOGP	0.98	0.99	1.28	-0.81	0.7	0.24	-1.05	0.24	2.97	-1.43	-0.54	-2.14	0.05	-0.03	-1.57	-0.56	-2.06	-1.84
Silicos-IT Log P	1	0.74	1.22	-0.07	0.75	0.98	-0.61	0.98	-0.22	-1.36	-0.15	-1.82	1.04	2.03	0.38	1.54	-0.1	0.01
Consensus Log P	1.13	1.03	1.26	0.01	0.93	0.85	-0.38	0.85	3.72	-1.08	-0.03	-1.75	1.23	1.58	0.24	1.23	0	0.16

[illegible]

CYP2D6 inhibitor	No	No	No	No	No	No	No	No	Yes	No	No	No	No	Yes	No	Yes	No	No
CYP3A4 inhibitor	No	No	No	No	No	No	No	No	No	No	No	No	No	Yes	No	Yes	No	No
log Kp (cm/s)	-6.45	-6.08	-6.26	-8.62	-6.58	-7.82	-8.76	-7.82	-4.31	-8.58	-8.41	-9.15	-7.91	-6.7	-8.3	-7.05	-8.64	-8.42
Lipinski #violations	0	0	0	0	0	0	1	0	0	0	0	0	1	0	1	0	2	2
Ghose #violations	1	3	0	1	0	0	1	0	0	2	1	1	0	0	0	0	0	0
Veber #violations	0	0	0	1	0	0	1	0	0	0	1	0	1	0	1	0	1	1
Egan #violations	0	0	0	1	0	0	1	0	0	0	1	0	1	0	1	0	1	1
Muegge #violations	1	1	1	1	1	0	2	0	1	1	0	2	2	0	2	0	3	3
Bioavailability Score	0.55	0.56	0.56	0.11	0.56	0.55	0.11	0.55	0.55	0.56	0.56	0.56	0.55	0.55	0.55	0.55	0.17	0.17
PAINS #alerts	0	0	0	0	1	1	1	1	0	0	0	0	1	0	0	1	1	1
Brenk #alerts	0	0	1	1	2	1	2	1	1	0	1	0	1	0	0	1	1	1
Leadlikeness #violations	1	1	1	1	1	0	1	0	2	1	0	1	1	0	1	0	1	1
Synthetic Accessibility	1.7	1	1.61	4.25	1.81	3.5	4.16	3.5	3.97	3.77	4.07	3.34	4.16	3.14	5.02	3.23	5.05	5.28

Remarks*		
A	Aqueous solubility is expressed on the basis of log S scale value ranges between -10 and 0 which is given in table mentioned below	
	Log S value	Aqueous solubility
	<-10	Insoluble
	-10 to -6	Poorly soluble
	-6 to -4	Moderately soluble

	-4 to -2	soluble	
	-2 to 0	Very soluble	
	>0	Highly soluble	
B	SwissADME have five methods to calculate the lipophilicity of molecules, and explained in		
1	Log Po/w (the partition coefficient between octanol and water which shows lipophilicity of molecules		
2	iLOGP is in-house physics-based method in which Generalized-Born and solvent accessible surface area model is used to calculate the free energy of solvation in n-octanol and water.		
3	XLOGP3 is an atomistic method comprising the corrective factors and knowledge-based library.		
4	WLOGP is a purely atomistic method based on the fragmental system.		
5	MLOGP is a prototype of topological method based on a linear relationship with 13 molecular descriptors		
	*SILICOS-IT is hybrid methods which rely on 7 topological descriptors and 27 fragments. Arithmetic mean of the values of the 5 methods is calculated in consensus log Po/w		

Table S3. Structural information of SARS-CoV-2 Main protease (6LU7) and ACE2 host cell receptor (1R4L) along with their native ligands. Active sites of the target proteins were analysed by using Computed Atlas for Surface Topography of Proteins (CASTp). The amino acids residues found in the active sites pockets of SARS CoV-2 Main protease (6LU7) and ACE2 host cell receptor (1R4L) are tabulated.

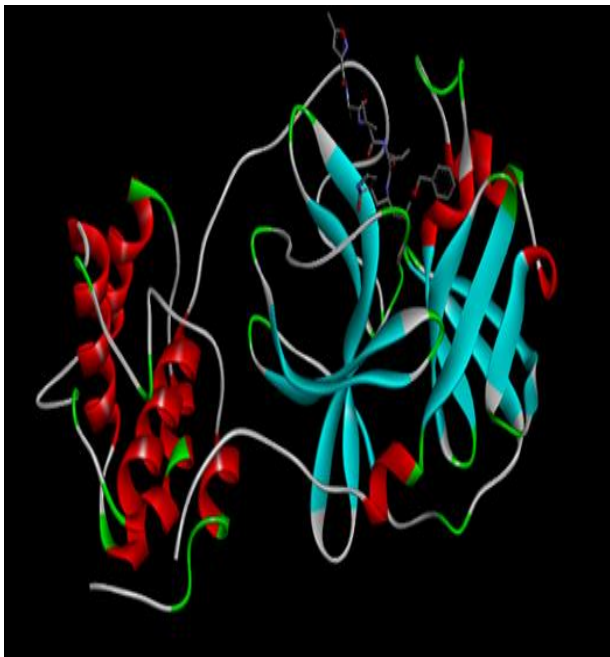
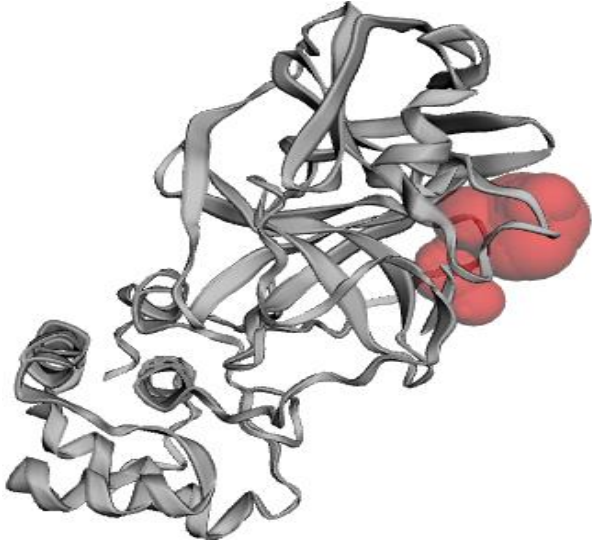
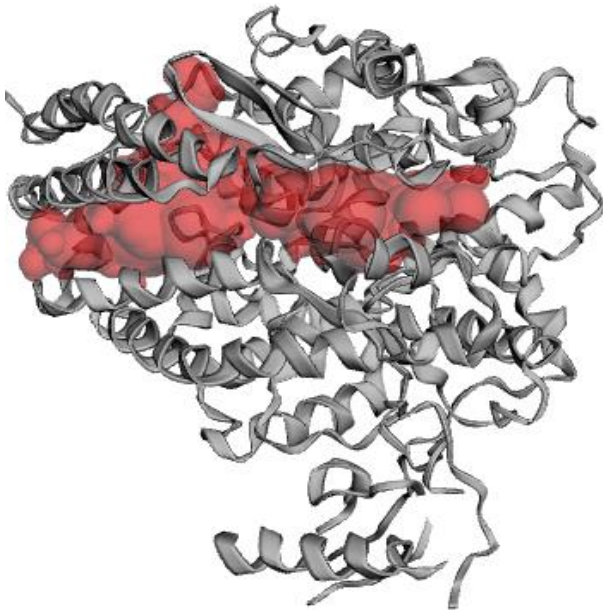
Target proteins	PDB ID	Structure	Native ligands	Active Sites	Active site amino acids
SARS-CoV-2 Main protease	6LU7		N-[(5-Methylisoxazol-3-Yl) Carbonyl] Alanyl-L-Valyl-N~1~-((1r,2z)-4-(Benzyloxy)-4-Oxo-1-[(3r)-2-Oxopyrrolidin-3-Yl]Methyl}But-2-Enyl)-L-Leucinamide		THRA:24, THRA:25, THRA:26, GLYA:143, LEUA:27, THRA:45, SERA:46, CYSA:145, HISA:164, ASPA:187, HISA:41, ARGA:188, META:49, GLNA:189, GLUA:166, THRA:190, ALAA:191, PROA:168, PHEA:140, LEUA:141, GLNA:192, HISA:172, HISA:163, ASNA:142, META:165, SERA:144, VALC:3, LEUC:4
Human Angiotensin Converting Enzyme	1R4L		N-Acetyl-D-Glucosamine Zinc Ion Chloride Ion (S,S)-2-{1-Carboxy-2-[3-(3,5-Dichloro-Benzyl)-3h-Imidazol-4-Yl]-Ethylamino}-4-Methyl-Pentanoic Acid		PHEA:32, GLUA:37, PHEA:40, SERA:43, SERA:44, SERA:47, TYRA:50, ASNA:52, ILEA:54, META:62, ASNA:63, GLYA:66, ASPA:67, TRPA:69, SERA:70, ALAA:71, LEUA:73, LYSA:74, SERA:77, LEUA:91, THRA:92, LYSA:94, LEUA:95, GLNA:96, GLNA:98, ALAA:99, LEUA:100, GLNA:102, ASNA:103, GLYA:104, SERA:105, SERA:106, LEUA:108, GLUA:110, SERA:113, LYSA:114, LEUA:116, ASNA:117, LEUA:120, ASNA:121, SERA:124, TYRA:127, SERA:128, THRA:129, GLYA:130, LYSA:131, CYSA:141, LEUA:142, LEUA:143, LEUA:144, GLUA:145, PROA:146, ASNA:149, GLUA:150, META:152, ALAA:153, ASNA:154, LYSA:187, META:190, ALAA:191, ASNA:194, TYRA:196, TYRA:199, TYRA:202, TRPA:203, GLYA:205, ASPA:206, GLUA:208, VALA:209, ASNA:210, GLYA:211, VALA:212, GLYA:268, ASPA:269, META:270, TRPA:271, ARGA:273, PHEA:274, THRA:276, ASNA:277, ILEA: 291, ASPA:292, THRA:294, ASPA:335, GLYA:337, ASNA:338, VALA:339, GLNA:340, LYSA:341, ALAA:342, VALA:343, CYSA:344, HISA:345, PROA:346, THRA:347, ALAA:348, TRPA:349, ASPA:350, GLYA:352, LYSA:353, GLYA:354, PHEA:356, LEUA:359, META:360, CYSA:361, LYSA:363, THRA:365, META:366, ASPA:367, ASPA:368, LEUA:370, THRA:371, HISA:374, GLUA:375, HISA:378, ASPA:382, TYRA:385, ALAA:386, PHEA:390, LEUA:391, LEUA:392, ARGA:393, ASNA:394, GLYA:395, ALAA:396, ASNA:397, GLUAA:398, HISA:401, GLUA:402, GLUA:406, SERA:409, LEUA:410, ALAA:413, PHEA:438, LYSA:441, GLNA:442, THRA:445, ILEA:446, THRA:449, LEUA:503, PHEA:504, HISA:505, SERA:507, ASNA:508, ASPA:509, TYRA:510, SERA:511, ARGA:514, TYRA:515, ARGA:518, THRA:519, GLNA:522, LYSA:562, SERA:563, GLUA:564, PROA:565, TRPA:566

Table S4. Blind docking of the secondary metabolites identified in *R. arboreum* petals against Target protein- SARS-CoV-2 Mpro (6LU7) and Human Angiotensin Converting Enzyme (1R4L). Glycyrrhizic acid was taken as a positive control

Pubchem CID	Phytochemicals	Target protein-SARS-CoV-2 Mpro (6LU7)		Human Angiotensin Converting Enzyme (1R4L)	
		Binding affinity (Kcal/mol)	Amino acid Residues	Binding affinity (Kcal/mol)	Amino acid Residues
14982	Glycyrrhizic acid (Positive control) (Zhang et al., 2020)	-8	ASNA:151, SERA:158, VALA:104, ILEA:106, ARGA:105, GNA:107, THRA:111, ASPA:245, THRA:243, GLYA:108, THRA:292, HISA:246, VALA:202, ILEA:249, PHEA:294, PROA:293, GLNA:110	-10.5	ASPA:206, ARGA:514, GLUA:402, HISA:378, TYRA:202, LYSA:562, HISA:401, ASNA:394, GLUA:398, TYRA:510, ASPA:508, TRPA:206, ASNA:394, ASPA:206, TYRA:385, ASPA:350, PHEA:390, TRPA:69, PHEA:40, LEUA:391, LEUA:73, ALAA:99, PHEA:92
14207924	2,4-Quinolinediamine	-5.3	LEUA:286, ASPA:289, LEUA:287, THRA:199, TYRA:239, TYRA:237, LEUA:272, META:276	-6.3	ALAA:348, TRPA:349, THRA:347, HISA:378, GLUA:402, TYRA:515, HISA:505, HISA:345, PROA:346, PHEA:504, TYRA:510
7420	3-Hydroxybenzoic acid	-5.4	PHEA:294, THRA:111, THRA:292, GLNA:110, ASPA:295, ASNA:151, GLNA:127, PHEA:112, PHEA:8, ASA:153, ILEA:152	-5.8	ARGA:273, HISA:345, TYRA:515, HISA:505, HISA:378, GLUA:402, ARGA:514, PHEA:504, TYRA:510, THRA:347, PROA:346, THRA:371, GLUA:375, HISA:374
637542	4-Coumaric acid	-5.9	ASPA:153, LYSA:102, SERA:158, VALA:104, ASNA:151, GLNA:110, THRA:111, THRA:292, ASPA:295, PHEA:294	-6.2	PROA:346, HISA:345, TYRA:515, ARGA:273, HISA:505, THRA:371, PHEA:274, ASPA:367, THRA:276, THRA:445

9799386	5-O-Feruloylquinic acid	-7.1	THRA:24, THRA:25, LEUA:27, HISA:41, META:49, PHEA:140, LEUA:141, ASNA:142, GLYA:143, SERA:144, CYSA:145, GLUA:166, META:165, HISA:163, HISA:164, HISA:172, GLNA:189,	-8.4	ASPA:367, ASPA:368, THRA:371, ARGA:273, GLUA:375, ALAA:348, HISA:374, THRA:347, HISA:378, ARGA:514, GLUA:402, TYRA:510, HISA:505, PHEA:504, TYRA:515, HISA:345, PHEA:274 META:360, PROA:346,
689043	Caffeic acid	-5.8	ASPA:153, LYSA:102, SERA:158, VALA:104, ASNA:151, GLNA:110, THRA:111, THRA:292, ASPA:295, PHEA:294	-6.4	PHEA:274, THRA:449, ILEA:446, THRA:519, GLNA:522, ARGA:518, THRA:445, GLUA:406, LEUA:370, ASPA:367, THRA:371
9064	Catechin	-6.7	THRA:24, THRA:25, LEUA:27, HISA:41, META:49, PHEA:140, LEUA:141, ASNA:142, GLYA:143, SERA:144, CYSA:145, GLUA:166, META:165, HISA:163, HISA:172	-9.2	THRA:445, ARGA:518, GLUA:406, ARGA:273, TYRA:515, GLUA:402, HISA:378, HISA:505, THRA:347, PHEA:504, HISA:345, GLUA:375, HISA:374, THRA:371, PROA:346, ASPA:269, THRA:276, ASPA:367, PHEA:274,
1794427	3-Caffeoyl-quinic acid	-7	LEUA:271, GLYA:275, THRA:199, META:276, THRA:198, ASPA:197, LYSA:137, ASPA:289, GLUA:290, ARGA:131, LEUA:286, TYRA:239, VALA:204, LEUA:287, LEUA:272, TYRA:237	-8.5	SERA:44, PHEA:40, LEUA:351, ASPA:350, THRA:347, ALAA:348, GLUA:375, PROA:346, HISA:378, HISA:374, HISA:345, PHEA:504, GLUA:402, HISA:505, TYRA:515, SERA:511, TYRA:510, ARGA:514, TRPA:349, SERA:47

72276	Epicatechin	-7.1	TYRA:237, TYRA:239, THRA:199, THRA:198, ARG:131, ASPA:197, LYSA:137, ASPA:289, LEUA:286, VALA:204, LEUA:287, LEUA:272	-8.4	THRA:445, LEUA:370, ARG:518, GLUA:406, ARG:273, TYRA:515, GLUA:402, HISA:378, HISA:505, THRA:347, HISA:345, GLUA:375, THRA:371, PROA:346, THRA:276, PHEA:274, ASPA:367, HISA:374
72	Protocatechuic acid	-5.4	PHEA:140, LEUA:141, SERA:144, GLYA:143, ASNA:142, CYSA:145, GLNA:189, GLUA:166, META:165, HISA:163	-5.8	ALAA:348, TRPA:349, THRA:347, HISA:378, GLUA:402, TYRA:515, HISA:505, GLUA:375, HISA:374, HISA:345, PROA:346, PHEA:504, TYRA:510
8742	Shikimic acid	-5.3	PHEA:140, LEUA:141, SERA:144, GLYA:143, ASNA:142, CYSA:145, GLUA:166, META:165, HISA:163, HISA:164, HISA:172,	-6.1	ALAA:348, THRA:347, HISA:378, GLUA:402, TYRA:515, HISA:505, HISA:374, GLUA:375, HISA:345, PROA:346, PHEA:504, TYRA:510,
6441280	5-O-Coumaroyl-D-quinic acid	-7.2	THRA:25, THRA:26, LEUA:27, HISA:41, META:49, PHEA:140, LEUA:141, ASNA:142, GLYA:143, SERA:144, CYSA:145, GLUA:166, META:165, HISA:163, HISA:164 HISA:172, GLNA:189	-8.3	ASPA:367, THRA:371, GLUA:375, GLUA:402, HISA:378, ARG:514, ALAA:348, THRA:347, TYRA:510, PHEA:504, TYRA:515, HISA:505, HISA:345, ARG:273, HISA:374, PROA:346, PHEA:274

6508	Quinic acid	-5.6	ASNA:72, SERA:121, ASNA:119, GLYA:120, VALA:18, ALAA:70, GLNA:69, GLYA:71, GLNA:19	-6.6	HISA:374, GLUA:375, HISA:345, HISA:505, PHEA:504, TYRA:515, TYRA:510, GLUA:402, TRPA:349, ALAA:348, ARG:514, THRA:347, HISA:378, PROA:346
5280343	Quercetin	-7	THRA:24, THRA:25, THRA:26, LEUA:27, HISA:41, PHEA:140, LEUA:141, ASNA:142, GLYA:143, SERA:144, CYSA:145, GLUA:166, META:165, HISA:163, HISA:172	-9.3	LEUA:370, GLUA:406, ARG:518, GLUA:402, HISA:505, TYRA:515, HISA:378, THRA:347, PROA:346, HISA:345, GLUA:375, HISA:374, THRA:371, PHEA:274, ARG:273, ASPA:269, THRA:276, ASPA:367, THRA:445
5878729	Quercetin-O-pentoside	-7.9	LYSA:137, ARG:131, ASNA:238, THRA:198, THRA:199, ASPA:197, LEUA:272, TYRA:239, LEUA:287, LEUA:286, ASPA:289, GLUA:288, LYSA:5, GLUA:290	-9.8	GLUA:406, ARG:518, THRA:347, GLUA:375, HISA:378, TYRA:515, HISA:505, GLUA:402, HISA:374, THRA:371, ARG:273, LEUA:503, TRPA:271, HISA:345, PHEA:274, PROA:346, THRA:276, ASPA:367, LEUA:370, THRA:445
14749097	Kaempferol-O-pentoside	-7.8	LEUA:272, TYRA:239, THRA:199, ASPA:197, THRA:198, ASNA:238, LYSA:137, ARG:131, GLUA:290, LYSA:5, GLUA:288, ASPA:289, LEUA:286, LEUA:287	-9.6	PHEA:274, ASPA:367, THRA:276, THRA:445, GLUA:406, PROA:346, ARG:518, GLUA:375, THRA:347, TYRA:515, HISA:505, GLUA:402, HISA:378, HISA:374, ARG:273, HISA:345, THRA:371

5280863	Kaempferol	-7	THRA:25, THRA:26, LEUA:27, HISA:41, PHEA:140, LEUA:141, ASNA:142, GLYA:143, SERA:144, CYSA:145, GLUA:166, META:165, HISA:163, HISA:172, GLNA:189	-8.6	HISA:505, HISA:345, ARGAS:273, TYRA:515, HISA:378, GLUA:375, GLUA:402, ARGAS:518, HISA:374, PHEA:274, GLUA:406, THRA:445, SERA:409, LEUA:370, THRA:371, PROA:346
107905	Epicatechin gallate	-7.4	ASPA:197, ASNA:238, TYRA:237, THRA:199, THRA:198, LYSA:137, ASPA:289, ARGAS:131, LEUA:286, LEUA:287, META:276, ALAA:285, GLYA:275, LEUA:271, LEUA:272, TYRA:239	-9.1	TYRA:515, HISA:345, HISA:505, PHEA:504, TYRA:510, THRA:347, TRPA:349, SERA:44, SERA: 44, ASPA:350, ALAA:348, ILEA:379, ASPA:382, TYRA:385, HISA:401, HISA:378, ARGAS:514, GLUA:375, HISA:374, GLUA:402, THRA:371, PROA:346,
5280459	Quercetin-O-rhamnoside	-7.7	LEUA:272, TYRA:239, THRA:199, ASPA:197, THRA:198, ASNA:238, LYSA:137, ARGAS:131, GLUA:290, LYSA:5, GLUA:288, ASPA:289, LEUA:286, LEUA:287	-10	THRA:445, ASPA:367, HISA:374, THRA:371, HISA:505, TYRA:515, GLUA:375, PROA:346, HISA:345, ARGAS:273, LEUA:144, ASNA:149, TRPA:271, ASPA:269, PHEA:274, GLUA:406, ARGAS:518

Table S5. *Rhododendron arboreum* petal phytochemicals with reported anti-viral studies

Phytochemicals	Antiviral Activity		References
4-Coumaric acid	Anti-human rhinoviruses	In vitro study	Kwon et al., 2019,
	Anti-influenza virus A (H3N2) and human herpes virus 1 and 2 (HSV-1 and HSV-2)	In vitro study	Stankova et al., 2009
Caffeic acid	Anti-hepatitis B virus	In vivo and in vitro study	Wang et al., 2009
	Anti-herpes simplex virus type 1 (HSV-1)	In vitro study	Ikeda et al, 2011
	Anti-severe fever with thrombocytopenia syndrome virus (SFTSV)	In vitro study	Ogawa et al., 2018
Catechin	Anti-influenza	In vitro study	Song et al., 2005
	Anti-hepatitis B virus	In vivo study	Li et al., 2001
	Anti-herpes simplex virus type 1 (HSV-1) and type 2 (HSV-2)	In vitro study	Lyu et al., 2005
3-Caffeoyl-quinic acid	Anti-hepatitis B virus	In vivo and in vitro study	Wang et al., 2009
	Anti-influenza virus (H1N1)	In vitro study	Urushisaki et al., 2011
	Anti-HIV activity	Molecular docking study	Serina et al., 2017
	Anti-avian influenza virus (H5N1)	Molecular docking study	Ren et al., 2019
	Anti-influenza A (H1N1/H3N2)	In vivo and in vitro study	Ding et al., 2017
	Anti-enterovirus 71	In vitro study	Li et al., 2013
	Anti-herpes virus	In vitro study	Xiehuang et al., 2008
Epicatechin	Anti-mayaro virus	In vitro and molecular docking study	Ferreira et al., 2018
	Anti-herpes simplex virus type 1 (HSV-1) and type 2 (HSV-2)	In vitro study	Lyu et al., 2005
Protocatechuic acid	Anti-avian influenza virus H9N2	In vivo study	Ou et al., 2014
	Anti-hepatitis B virus	In vitro study	Dai et al., 2017
	Anti-influenza virus A	In vitro study	Lu et al., 2002

	Anti-herpes simplex virus	In vitro study	Hassan et a., 2017
Shikimic acid	Anti-avian influenza virus	In vitro study	Singh et al., 2020
	Precursor molecule for the drug oseltamivir phosphate used against influenza viruses types A and B, avian influenza virus H5N1, and human influenza virus H1N1	In vitro study	Quiroz et al., 2014
Quinic acid	Anti-dengue virus	In vitro study	Zanello et al., 2015
	Anti-hepatitis B virus	In vivo and in vitro study	Wang et al., 2009
	Anti-HIV-1-RT activity	Molecular docking study	Yazdi et al., 2019
Quercetin	Anti-dengue virus 2 (DENV2)	Molecular docking study	Ismail et al., 2017
	Anti-herpes simplex virus type 1 (HSV-1) and type 2 (HSV-2)	In vitro study	Lyu et al., 2005
	Anti-influenza A virus	In vitro study	Wu et al., 2016
	Anti-hepatitis C virus	In vivo study	Rojas et al., 2016
	Anti-feline calicivirus (FCV) and murine norovirus (MNV)	In vivo study	Seo et al., 2016
	Anti-SARS coronavirus	In vivo study	Schwarz et al., 2014
Kaempferol	Anti-influenza B virus	In vitro study	Yang et al., 2014
	Anti-feline calicivirus (FCV)	In vivo study	Seo et al., 2016
	Anti-human cytomegalovirus	In vivo study	Mitrocotsa et al., 2000
	Anti-dengue virus and japanese encephalitis virus	In vivo study	Care et al., 2020
	Anti-japanese-encephalitis-virus	In vivo study	Zhang et al., 2012

	Anti-SARS coronavirus	In vivo study	Schwarz et al., 2014
	Anti-human rhinoviruses	In vitro study	Kwon et al., 2019
Epicatechin gallate	Anti-grass carp reovirus (GCRV)	In vitro study	Zhang et al., 2020
	Anti-feline calicivirus (FCV) and murine norovirus (MNV)	In vivo study	Seo et al., 2016
Quercetin-O-rhamnoside	Anti-influenza A virus	In vitro study	Choi et al., 2009
	Anti-influenza virus	In vitro and Molecular docking study	Gansukh et al., 2020

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