

## LAMPIRAN

### Lampiran 1. Prediksi Aktivitas Biologis Senyawa Ikan Sidat dengan PASS Online

#### 1. Senyawa dengan aktivitas biologis antiviral influenza dan influenza A dengan Pa>0,6

Nama Senyawa	CID	Pa	Pi	Aktivitas Biologis	SMILE
4-Hydroxybenzenesulfonic acid	4765	0.792	0.002	Antiviral (Influenza A)	<chem>C1=CC(=CC=C1O)S(=O)(=O)O</chem>
15(R)-Prostaglandin D2	132282632	0.749	0.004	Antiviral (Influenza)	<chem>CCCCC(C=CC1C(C(CC1=O)O)CC=CCCC(=O)O)O</chem>
1beta-Hydroxycholic acid	5283893	0.728	0.004	Antiviral (Influenza)	<chem>CC(CCC(=O)O)C1CCC2C1(C(CC3C2C(CC4C3(C(CC(C4)O)O)C)O)O)C</chem>
4,4-dimethyl-5alpha-cholesta-8,14,24-trien-3beta-ol	443212	0.72	0.004	Antiviral (Influenza)	<chem>CC(CCC=C(C)C)C1CC=C2C1(CCC3=C2CCC4C3(CCC(C4(C)C)O)C)C</chem>
Carboprost	5281075	0.709	0.005	Antiviral (Influenza)	<chem>CCCCC(C)(C=CC1C(CC(C1CC=CCCC(=O)O)O)O)O</chem>
4-Dodecylbenzenesulfonic acid	8485	0.672	0.008	Antiviral (Influenza)	<chem>CCCCCCCCCCCCC1=CC=C(C=C1)S(=O)(=O)O</chem>
10-HDA	5312738	0.642	0.01	Antiviral (Influenza)	<chem>C(CCCC=CC(=O)O)CCCO</chem>
10,16-Dihydroxyhexadecanoic acid	441449	0.621	0.011	Antiviral (Influenza)	<chem>C(CCCCC(=O)O)CCCC(CCCCCCO)O</chem>
9,10-Dihydroxystearic acid	89377	0.599	0.013	Antiviral (Influenza)	<chem>CCCCCCCCC(C(CCCCCCCC(=O)O)O)O</chem>
(R)-2-hydroxystearic acid	439885	0.619	0.012	Antiviral (Influenza)	<chem>CCCCCCCCCCCCCCCCC(C(=O)O)O</chem>
2-Hydroxymyristic acid	1563	0.619	0.012	Antiviral (Influenza)	<chem>CCCCCCCCCCCCC(C(=O)O)O</chem>
2-Hydroxytetracosanoic acid	102430	0.619	0.012	Antiviral (Influenza)	<chem>CCCCCCCCCCCCCCCCCCCCCCCCC(C(=O)O)O</chem>
3-Hydroxysebacic acid	3017884	0.605	0.013	Antiviral (Influenza)	<chem>C(CCCC(=O)O)CCC(CC(=O)O)O</chem>
4-(Heptyloxy)benzoic acid	85154	0.625	0.011	Antiviral (Influenza)	<chem>CCCCCOC1=CC=C(C=C1)C(=O)O</chem>

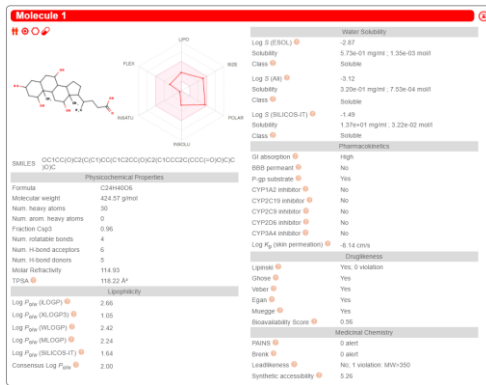
1,2,3-cyclopropanetricarboxylic acid	604010	0.609	0.012	Antiviral (Influenza)	CCOC(=O)C1C(C1C(=O)OCC)C(=O)OCC
(2R)-2,3-Dihydroxypropanoic acid	439194	0.6	0.013	Antiviral (Influenza)	C(C(C(=O)O)O)O
6-acetyloxy-7-ethenyl-1,4a,7-trimethyl-3,4,4b,5,6,8,10,10a-octahydro-2H-phenanthrene-1-carboxylic acid	51694232	0.674	0.007	Antiviral (Influenza)	CC(=O)OC1CC2C(=CCC3C2(CCCC3(C)C(=O)O)C)CC1(C)C=C
(±)13-HODE	6443013	0.646	0.009	Antiviral (Influenza)	CCCCC(C=CC=CCCCCCCC(=O)O)O
(R)-3-Hydroxy myristic acid	5288266	0.635	0.01	Antiviral (Influenza)	CCCCCCCCCCCC(CC(=O)O)O
3-Hydroxypentadecanoic acid	182089	0.635	0.01	Antiviral (Influenza)	CCCCCCCCCCCC(CC(=O)O)O
4-(hydroxymethyl)benzoic acid	76360	0.622	0.011	Antiviral (Influenza)	C1=CC(=CC=C1CO)C(=O)O
Androstanolone	10635	0.658	0.009	Antiviral (Influenza)	CC12CCC(=O)CC1CCC3C2CCC4(C3CCC4O)C
19(R)-hydroxy Prostaglandin A2	16061098	0.655	0.009	Antiviral (Influenza)	CC(CCCC(C=CC1C=CC(=O)C1CC=CCCC(=O)O)O)O
4-[(2,4-Dihydroxyphenyl)diazenyl] benzenesulfonic acid	11035	0.653	0.009	Antiviral (Influenza)	C1=CC(=CC=C1N=NC2=C(C=C(C=C2)O)O)S(=O)(=O)O
14(Z)-Eicosenoic acid	5282770	0.652	0.009	Antiviral (Influenza)	CCCCC=CCCCCCCCCCCCC(=O)O
Androst-4-en-3-one	250294	0.646	0.009	Antiviral (Influenza)	CC12CCCC1C3CCC4=CC(=O)CCC4(C3CC2)C
13Z,16Z-docosadienoic acid	5312554	0.599	0.013	Antiviral (Influenza)	CCCCC=CCC=CCCCCCCCCCCC(=O)O
8Z,11Z,14Z-Eicosatrienoic acid	5280581	0.599	0.013	Antiviral (Influenza)	CCCCC=CCC=CCC=CCCCCCCC(=O)O
Adrenic acid	5497181	0.599	0.013	Antiviral (Influenza)	CCCCC=CCC=CCC=CCC=CCCCCCC(=O)O
Arachidonic acid	444899	0.599	0.013	Antiviral (Influenza)	CCCCC=CCC=CCC=CCC=CCCC(=O)O
(±)11(12)-EET	53480479	0.609	0.012	Antiviral (Influenza)	CCCCC=CCC1C(O)CC=CCC=CCCC(=O)O
8-(4-Sulfophenyl) octanoic acid	53987276	0.668	0.008	Antiviral (Influenza)	C1=CC(=CC=C1CCCCCCCC(=O)O)S(=O)(=O)O

2. Senyawa dengan aktivitas biologis antiviral poxvirus, rhinovirus, dan picornavirus dengan  $P_a > 0,7$ 

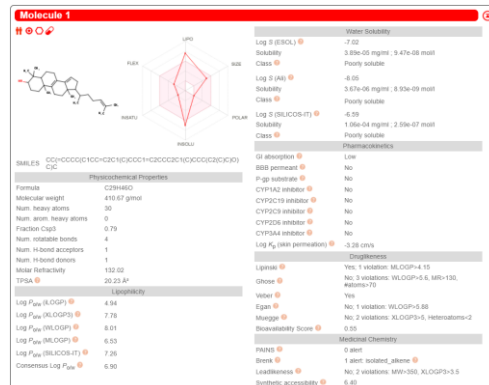
Nama Senyawa	CID	$P_a$	$P_i$	Aktivitas Biologis	SMILES
Adenosine-5'-diphosphate Di(monocyclohexylammonium)salt	197	0.956	0.001	Antiviral (Poxvirus)	<chem>C1=NC(=C2C(=N1)N(C=N2)C3C(C(C(O3)COP(=O)(O)OP(=O)(O)O)O)O)N</chem>
(10S)-Juvenile hormone III diol	10859043	0.891	0	Antiviral (Rhinovirus)	<chem>CC(=CCCC(=CC(=O)OC)C)CCC(C(C)(C)O)O</chem>
9-(2,3-dihydroxypropoxy)-9-oxononanoic acid	45783154	0.762	0.002	Antiviral (Rhinovirus)	<chem>C(CCCC(=O)O)CCCC(=O)OCC(CO)O</chem>
2,3-dihydroxypropyl 12-methyltridecanoate	10828227	0.749	0.002	Antiviral (Rhinovirus)	<chem>CC(C)CCCCCCCCCCC(=O)OCC(CO)O</chem>
4-methyl-2-oxopentanoic acid	70	0.739	0.004	Antiviral	<chem>CC(C)CC(=O)C(=O)O</chem>
4,4-dimethyl-5alpha-cholesta-8,14,24-trien-3beta-ol	443212	0.719	0.002	Antiviral (Rhinovirus)	<chem>CC(CCC=C(C)C)C1CC=C2C1(CCC3=C2CCC4C3(CCC(C4(C)C)O)C)C</chem>
16-Heptadecyne-1,2,4-triol	3015189	0.717	0.002	Antiviral (Rhinovirus)	<chem>C#CCCCCCCCCCCCC(CC(CO)O)O</chem>
Chenodeoxycholic acid 3-sulfate	21252312	0.707	0.003	Antiviral (Rhinovirus)	<chem>CC(CCC(=O)O)C1CCC2C1(CCC3C2C(CC4C3(CCC(C4)OS(=O)(=O)O)C)O)C</chem>
Cannabigerol	5315659	0.703	0.003	Antiviral (Rhinovirus)	<chem>CCCCC1=CC(=C(C(=C1)O)CC=C(C)CCC=C(C)C)O</chem>
2-Oxobutyric acid	58	0.7	0.004	Antiviral	<chem>CCC(=O)C(=O)O</chem>
Avocadyne Acetate	3952079	0.7	0.003	Antiviral (Rhinovirus)	<chem>CC(=O)OCC(CC(CCCCCCCCCCCC#C)O)O</chem>
C12-AE1S (TENTATIVE)	24761	0.701	0.003	Antiviral (Rhinovirus)	<chem>CCCCCCCCCCCCOCCOS(=O)(=O)O</chem>

## Lampiran 2. Prediksi Molekul Obat SwissADME

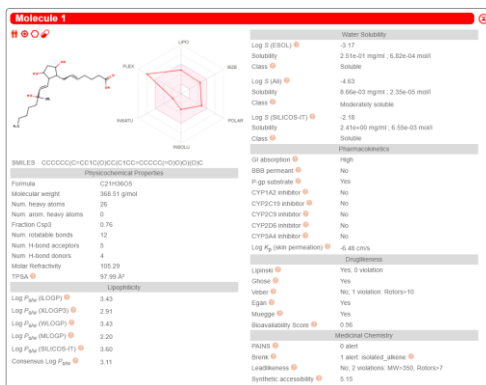
### 1. 1beta-Hydroxycholeic acid



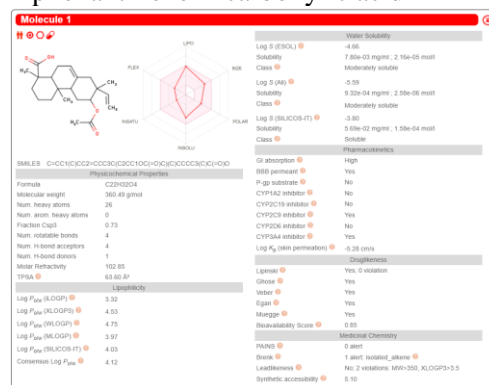
### 2. 4,4-dimethyl-5alpha-cholesta-8,14,24-trien-3beta-ol



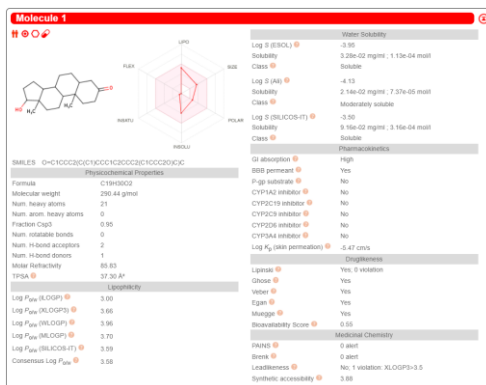
### 3. Carboprost



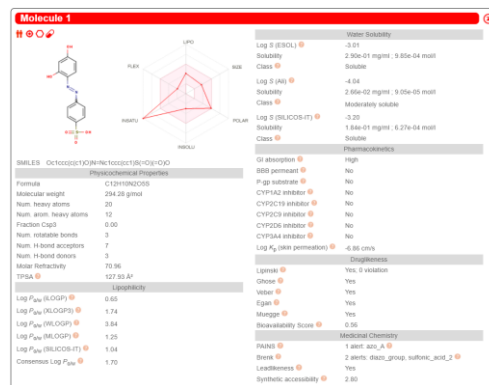
### 4. 6-acetyloxy-7-ethenyl-1,4a,7-trimethyl-3,4,4b,5,6,8,10,10a-octahydro-2H-phenanthrene-1-carboxylic acid



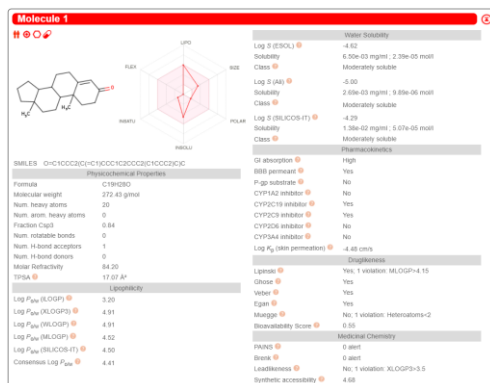
### 5. Androstanolone



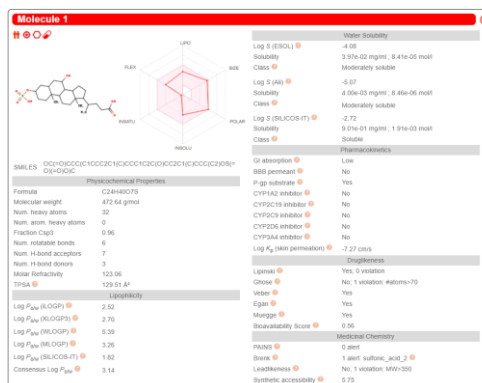
### 6. 4-[(2,4-Dihydroxyphenyl)diazenyl] benzenesulfonic acid



## 7. Androst-4-en-3-one



## 8. Chenodeoxycholic acid 3-sulfate



### Lampiran 3. Konfigurasi Penambatan Molekuler Autodock Vina

#### 1. Data input Main Protease (M<sup>pro</sup>) Sars-CoV-2

```
*Conf6w63_Ligand - Notepad
File Edit Format View Help
receptor = 6w63_Mpro.pdbqt
ligand = Ligan_ID.pdbqt

out = out1.pdbqt

center_x = -19.340
center_y = 18.376
center_z = -27.228

size_x = 18
size_y = 18
size_z = 18

log = log.txt
num_modes = 9
exhaustiveness = 32
```

#### 3. Data input Receptor Angiotensin-converting Enzyme (ACE2)

```
*Conf6m0jA_Ligand - Notepad
File Edit Format View Help
receptor = 6m0j_ACE2.pdbqt
ligand = Ligan_ID.pdbqt

out = out1.pdbqt

center_x = -34.607
center_y = 31.063
center_z = -3.383

size_x = 20
size_y = 40
size_z = 20

log = log.txt
num_modes = 9
exhaustiveness = 32
```

#### 2. Data input Receptor Binding Domain (RBD) Sars-CoV-2

```
*Conf6m0jE_Ligand - Notepad
File Edit Format View Help
receptor = 6m0jE_RBD.pdbqt
ligand = Ligan_ID.pdbqt

out = out1.pdbqt

center_x = -36.588
center_y = 28.757
center_z = 3.603

size_x = 24
size_y = 40
size_z = 24

log = log.txt
num_modes = 9
exhaustiveness = 32
```

## Lampiran 4. Log Penambatan M<sup>PRO</sup> Sars-CoV-2 Autodock Vina

### 1. Remdesivir

```
Log_6w63_Remdesivir - Notepad
File Edit Format View Help
Reading input ... done.
Setting up the scoring function ... done.
Analyzing the binding site ... done.
Using random seed: 234742180
Performing search ... done.
Refining results ... done.

mode | affinity | dist from best mode
      | (kcal/mol) | rmsd l.b. | rmsd u.b.
-----+-----+-----+-----
1      -8.2      0.000      0.000
2      -7.9      1.221      2.087
3      -7.8      2.914      7.437
4      -7.7      2.866      7.142
5      -7.6      2.632      7.390
6      -7.6      2.149      3.924
7      -7.6      2.154      4.884
8      -7.5      2.406      4.365
9      -7.5      2.689      7.549
Writing output ... done.
```

### 2. Nelfinavir

```
Log6w63_Nelfinavir - Notepad
File Edit Format View Help
Reading input ... done.
Setting up the scoring function ... done.
Analyzing the binding site ... done.
Using random seed: 826519980
Performing search ... done.
Refining results ... done.

mode | affinity | dist from best mode
      | (kcal/mol) | rmsd l.b. | rmsd u.b.
-----+-----+-----+-----
1      -8.5      0.000      0.000
2      -8.4      1.772      3.059
3      -8.4      1.476      2.224
4      -8.4      2.061      3.101
5      -8.3      1.536      1.994
6      -8.3      2.020      4.259
7      -8.2      2.442      7.604
8      -8.2      2.563      6.632
9      -8.1      2.380      3.922
Writing output ... done.
```

### 3. Hydroxychloroquine

```
Log6w63_Hydroxychloroquine - Notepad
File Edit Format View Help
Reading input ... done.
Setting up the scoring function ... done.
Analyzing the binding site ... done.
Using random seed: -870692872
Performing search ... done.
Refining results ... done.

mode | affinity | dist from best mode
      | (kcal/mol) | rmsd l.b. | rmsd u.b.
-----+-----+-----+-----
1      -6.2      0.000      0.000
2      -6.1      2.309      3.900
3      -6.1      2.472      3.984
4      -6.0      1.682      2.535
5      -5.9      2.777      4.391
6      -5.9      1.237      1.809
7      -5.9      2.798      4.445
8      -5.9      4.004      7.712
9      -5.8      3.575      6.212
Writing output ... done.
```

### 4. Molnupiravir

```
log6w63_Molnupiravir - Notepad
File Edit Format View Help
Reading input ... done.
Setting up the scoring function ... done.
Analyzing the binding site ... done.
Using random seed: 1646261760
Performing search ... done.
Refining results ... done.

mode | affinity | dist from best mode
      | (kcal/mol) | rmsd l.b. | rmsd u.b.
-----+-----+-----+-----
1      -6.8      0.000      0.000
2      -6.6      3.726      5.667
3      -6.4      2.918      5.839
4      -6.4      2.142      5.070
5      -6.4      3.030      6.572
6      -6.4      3.692      6.050
7      -6.4      2.234      5.149
8      -6.4      3.272      5.558
9      -6.3      3.359      6.916
Writing output ... done.
```

### 5. Ibeta-Hydroxycholic acid

```
Log6w63_5283893 - Notepad
File Edit Format View Help
Reading input ... done.
Setting up the scoring function ... done.
Analyzing the binding site ... done.
Using random seed: -1441435152
Performing search ... done.
Refining results ... done.

mode | affinity | dist from best mode
      | (kcal/mol) | rmsd l.b. | rmsd u.b.
-----+-----+-----+-----
1      -7.3      0.000      0.000
2      -7.3      3.670      7.738
3      -7.0      2.229      3.007
4      -6.9      3.130      7.796
5      -6.9      3.611      8.204
6      -6.8      2.191      3.663
7      -6.8      2.162      3.967
8      -6.8      1.862      8.210
9      -6.8      3.652      6.371
Writing output ... done.
```

### 6. 4,4-dimethyl-5alpha-cholesta-8,14,24-trien-3beta-ol

```
Log6w63_443212 - Notepad
File Edit Format View Help
Reading input ... done.
Setting up the scoring function ... done.
Analyzing the binding site ... done.
Using random seed: 1875534124
Performing search ... done.
Refining results ... done.

mode | affinity | dist from best mode
      | (kcal/mol) | rmsd l.b. | rmsd u.b.
-----+-----+-----+-----
1      -7.3      0.000      0.000
2      -7.3      2.605      5.072
3      -7.2      3.549      9.335
4      -7.0      1.539      3.543
5      -6.9      2.113      3.614
6      -6.9      2.282      3.941
7      -6.8      3.158      8.240
8      -6.8      2.232      4.945
9      -6.8      1.560      3.189
Writing output ... done.
```

## 7. Carboprost

```
Log6w63_5281075 - Notepad
File Edit Format View Help
Reading input ... done.
Setting up the scoring function ... done.
Analyzing the binding site ... done.
Using random seed: -2131044288
Performing search ... done.
Refining results ... done.
```

mode	affinity (kcal/mol)	dist from best mode rmsd l.b.	rmsd u.b.
1	-6.3	0.000	0.000
2	-5.9	2.042	3.632
3	-5.9	1.907	3.594
4	-5.8	2.169	3.365
5	-5.8	3.052	6.668
6	-5.8	3.330	5.728
7	-5.8	3.472	6.311
8	-5.7	1.520	2.296
9	-5.7	2.741	5.611

```
Writing output ... done.
```

## 9. Androstanolone

```
Log6w63_10635 - Notepad
File Edit Format View Help
Reading input ... done.
Setting up the scoring function ... done.
Analyzing the binding site ... done.
Using random seed: -195838696
Performing search ... done.
Refining results ... done.
```

mode	affinity (kcal/mol)	dist from best mode rmsd l.b.	rmsd u.b.
1	-7.5	0.000	0.000
2	-7.3	3.085	6.655
3	-6.8	2.762	3.796
4	-6.7	3.968	7.343
5	-6.7	4.801	7.050
6	-6.6	1.082	6.380
7	-6.5	1.519	2.965
8	-6.5	1.893	6.338
9	-6.5	1.267	6.556

```
Writing output ... done.
```

## 11. Androst-4-en-3-one

```
Log6w63_250294 - Notepad
File Edit Format View Help
Reading input ... done.
Setting up the scoring function ... done.
Analyzing the binding site ... done.
Using random seed: -1318792496
Performing search ... done.
Refining results ... done.
```

mode	affinity (kcal/mol)	dist from best mode rmsd l.b.	rmsd u.b.
1	-7.2	0.000	0.000
2	-7.0	2.877	3.872
3	-6.9	2.553	3.507
4	-6.9	1.391	1.712
5	-6.8	3.635	5.869
6	-6.7	3.031	6.252
7	-6.6	2.949	6.276
8	-6.5	3.340	5.166
9	-6.5	3.041	6.283

```
Writing output ... done.
```

## 8. 6-acetyloxy-7-ethenyl-1,4a,7-trimethyl-3,4,4b,5,6,8,10,10a-octahydro-2H-phenanthrene-1-carboxylic acid

```
Log6w63_51694232 - Notepad
File Edit Format View Help
Reading input ... done.
Setting up the scoring function ... done.
Analyzing the binding site ... done.
Using random seed: -1922542440
Performing search ... done.
Refining results ... done.
```

mode	affinity (kcal/mol)	dist from best mode rmsd l.b.	rmsd u.b.
1	-6.9	0.000	0.000
2	-6.6	1.775	6.897
3	-6.5	2.613	6.076
4	-6.4	3.382	7.052
5	-6.2	2.006	2.996
6	-6.2	2.193	6.727
7	-6.2	2.681	3.517
8	-6.1	2.737	5.203
9	-6.1	2.946	4.908

```
Writing output ... done.
```

## 10. 4-[(2,4-Dihydroxyphenyl)diazenyl]benzenesulfonic acid

```
Log6w63_11035 - Notepad
File Edit Format View Help
Reading input ... done.
Setting up the scoring function ... done.
Analyzing the binding site ... done.
Using random seed: -2117621460
Performing search ... done.
Refining results ... done.
```

mode	affinity (kcal/mol)	dist from best mode rmsd l.b.	rmsd u.b.
1	-6.9	0.000	0.000
2	-6.9	1.149	1.889
3	-6.5	5.434	7.313
4	-6.3	5.462	6.885
5	-6.3	5.155	5.951
6	-6.1	5.490	6.451
7	-6.1	3.455	8.469
8	-6.1	5.261	6.146
9	-6.1	5.714	8.513

```
Writing output ... done.
```

## 12. Chenodeoxycholic acid 3-sulfate

```
Log6w63_21252312 - Notepad
File Edit Format View Help
Reading input ... done.
Setting up the scoring function ... done.
Analyzing the binding site ... done.
Using random seed: 1217666832
Performing search ... done.
Refining results ... done.
```

mode	affinity (kcal/mol)	dist from best mode rmsd l.b.	rmsd u.b.
1	-8.1	0.000	0.000
2	-7.8	2.069	3.236
3	-7.4	2.852	4.230
4	-7.3	3.478	9.110
5	-7.2	1.866	2.506
6	-7.2	3.712	9.028
7	-7.1	3.422	8.824
8	-7.1	4.047	8.517
9	-7.0	3.426	8.918

```
Writing output ... done.
```



## Lampiran 5. Log Penambatan RBD SARS-CoV-2- Autodock Vina

### 1. Remdesivir

```
Log6m0jE_Remdesivir - Notepad
File Edit Format View Help
Reading input ... done.
Setting up the scoring function ... done.
Analyzing the binding site ... done.
Using random seed: 736359356
Performing search ... done.
Refining results ... done.

mode | affinity | dist from best mode
      | (kcal/mol) | rmsd l.b. | rmsd u.b.
-----+-----+-----+-----
1      -6.7      0.000      0.000
2      -6.5      1.554      1.955
3      -6.4      4.106      9.074
4      -6.4      1.748      2.281
5      -6.4      4.729      10.111
6      -6.3      1.741      2.265
7      -6.2      3.078      9.011
8      -6.2      3.360      8.983
9      -6.2      1.972      2.848
Writing output ... done.
```

### 2. Nelfinavir

```
Log6m0jE_Nelfinavir - Notepad
File Edit Format View Help
Reading input ... done.
Setting up the scoring function ... done.
Analyzing the binding site ... done.
Using random seed: 1757677536
Performing search ... done.
Refining results ... done.

mode | affinity | dist from best mode
      | (kcal/mol) | rmsd l.b. | rmsd u.b.
-----+-----+-----+-----
1      -7.0      0.000      0.000
2      -6.9      2.251      3.773
3      -6.8      5.017      11.608
4      -6.7      7.919      11.519
5      -6.7      3.581      6.305
6      -6.6      5.126      11.180
7      -6.6      2.674      5.485
8      -6.6      1.698      4.829
9      -6.6      1.141      1.444
Writing output ... done.
```

### 3. Hydroxychloroquine

```
Log6m0jE_Hydroxychloroquine - Notepad
File Edit Format View Help
Reading input ... done.
Setting up the scoring function ... done.
Analyzing the binding site ... done.
Using random seed: -1592583664
Performing search ... done.
Refining results ... done.

mode | affinity | dist from best mode
      | (kcal/mol) | rmsd l.b. | rmsd u.b.
-----+-----+-----+-----
1      -5.2      0.000      0.000
2      -5.1      2.156      3.160
3      -5.0      3.615      5.726
4      -5.0      3.818      6.098
5      -4.9      2.921      4.347
6      -4.9      2.846      4.604
7      -4.9      4.284      6.702
8      -4.9      2.683      4.188
9      -4.9      3.536      5.076
Writing output ... done.
```

### 4. Molnupiravir

```
log6m0jE_Molnupiravir1 - Notepad
File Edit Format View Help
Reading input ... done.
Setting up the scoring function ... done.
Analyzing the binding site ... done.
Using random seed: -1501864952
Performing search ... done.
Refining results ... done.

mode | affinity | dist from best mode
      | (kcal/mol) | rmsd l.b. | rmsd u.b.
-----+-----+-----+-----
1      -5.9      0.000      0.000
2      -5.8      1.807      1.886
3      -5.7      3.204      6.519
4      -5.6      1.898      2.155
5      -5.5      1.927      2.228
6      -5.4      4.017      7.682
7      -5.4      1.853      2.878
8      -5.4      3.985      6.935
9      -5.4      3.533      7.480
Writing output ... done.
```

### 5. Ibeta-Hydroxycholic acid

```
*Log6m0jE_5283893 - Notepad
File Edit Format View Help
Reading input ... done.
Setting up the scoring function ... done.
Analyzing the binding site ... done.
Using random seed: 1755523256
Performing search ... done.
Refining results ... done.

mode | affinity | dist from best mode
      | (kcal/mol) | rmsd l.b. | rmsd u.b.
-----+-----+-----+-----
1      -6.8      0.000      0.000
2      -6.6      3.804      6.760
3      -6.5      1.992      2.480
4      -6.3      2.906      4.625
5      -6.1      1.822      2.549
6      -6.1      3.527      7.135
7      -6.1      4.243      7.755
8      -6.1      3.476      6.701
9      -6.1      4.342      7.681
Writing output ... done.
```

### 6. 4,4-dimethyl-5alpha-cholesta-8,14,24-trien-3beta-ol

```
Log6m0jE_443212 - Notepad
File Edit Format View Help
Reading input ... done.
Setting up the scoring function ... done.
Analyzing the binding site ... done.
Using random seed: -2080540416
Performing search ... done.
Refining results ... done.

mode | affinity | dist from best mode
      | (kcal/mol) | rmsd l.b. | rmsd u.b.
-----+-----+-----+-----
1      -6.9      0.000      0.000
2      -6.4      2.772      8.344
3      -6.3      1.697      3.335
4      -6.2      1.800      3.363
5      -6.1      8.450      13.683
6      -6.0      4.060      5.820
7      -5.9      2.320      3.571
8      -5.9      4.351      6.039
9      -5.9      2.424      4.587
Writing output ... done.
```

## 7. Carboprost

```
Log6m0jE_5281075 - Notepad
File Edit Format View Help
Reading input ... done.
Setting up the scoring function ... done.
Analyzing the binding site ... done.
Using random seed: -1063172096
Performing search ... done.
Refining results ... done.

mode | affinity | dist from best mode
      | (kcal/mol) | rmsd l.b. | rmsd u.b.
-----+-----+-----+-----
1     | -5.3      | 0.000     | 0.000
2     | -5.2      | 1.783     | 2.092
3     | -5.0      | 2.839     | 7.069
4     | -5.0      | 3.052     | 5.618
5     | -4.9      | 3.456     | 7.279
6     | -4.9      | 2.496     | 5.486
7     | -4.8      | 2.178     | 7.859
8     | -4.8      | 2.412     | 5.427
9     | -4.8      | 2.536     | 4.689

Writing output ... done.
```

## 9. Androstanolone

```
Log6m0jE_10635 - Notepad
File Edit Format View Help
Reading input ... done.
Setting up the scoring function ... done.
Analyzing the binding site ... done.
Using random seed: -1391909756
Performing search ... done.
Refining results ... done.

mode | affinity | dist from best mode
      | (kcal/mol) | rmsd l.b. | rmsd u.b.
-----+-----+-----+-----
1     | -5.9      | 0.000     | 0.000
2     | -5.7      | 3.321     | 5.047
3     | -5.7      | 2.075     | 6.691
4     | -5.6      | 2.140     | 2.867
5     | -5.5      | 2.057     | 6.652
6     | -5.5      | 4.051     | 6.272
7     | -5.5      | 1.695     | 2.487
8     | -5.4      | 1.160     | 6.691
9     | -5.4      | 2.702     | 4.431

Writing output ... done.
```

## 11. Androst-4-en-3-one

```
Log6m0jE_250294 - Notepad
File Edit Format View Help
Reading input ... done.
Setting up the scoring function ... done.
Analyzing the binding site ... done.
Using random seed: 1059192832
Performing search ... done.
Refining results ... done.

mode | affinity | dist from best mode
      | (kcal/mol) | rmsd l.b. | rmsd u.b.
-----+-----+-----+-----
1     | -5.7      | 0.000     | 0.000
2     | -5.7      | 2.620     | 6.000
3     | -5.7      | 5.319     | 7.767
4     | -5.7      | 2.211     | 4.006
5     | -5.6      | 5.691     | 7.985
6     | -5.5      | 3.035     | 4.961
7     | -5.5      | 3.194     | 5.999
8     | -5.4      | 14.310    | 16.473
9     | -5.4      | 5.186     | 7.835

Writing output ... done.
```

## 8. 6-acetyloxy-7-ethenyl-1,4a,7-trimethyl-3,4,4b,5,6,8,10,10a-octahydro-2H-phenanthrene-1-carboxylic acid

```
Log6m0jE_51694232 - Notepad
File Edit Format View Help
Reading input ... done.
Setting up the scoring function ... done.
Analyzing the binding site ... done.
Using random seed: -287040336
Performing search ... done.
Refining results ... done.

mode | affinity | dist from best mode
      | (kcal/mol) | rmsd l.b. | rmsd u.b.
-----+-----+-----+-----
1     | -5.9      | 0.000     | 0.000
2     | -5.8      | 2.446     | 3.957
3     | -5.7      | 2.954     | 6.857
4     | -5.6      | 3.379     | 4.573
5     | -5.6      | 2.599     | 6.969
6     | -5.5      | 1.942     | 2.493
7     | -5.5      | 1.581     | 6.799
8     | -5.5      | 0.984     | 1.037
9     | -5.4      | 2.664     | 6.308

Writing output ... done.
```

## 10. 4-[(2,4-Dihydroxyphenyl)diazenyl]benzenesulfonic acid

```
Log6m0jE_11035 - Notepad
File Edit Format View Help
Reading input ... done.
Setting up the scoring function ... done.
Analyzing the binding site ... done.
Using random seed: 1667413348
Performing search ... done.
Refining results ... done.

mode | affinity | dist from best mode
      | (kcal/mol) | rmsd l.b. | rmsd u.b.
-----+-----+-----+-----
1     | -6.2      | 0.000     | 0.000
2     | -6.1      | 0.270     | 1.368
3     | -5.8      | 1.056     | 1.846
4     | -5.8      | 6.324     | 8.753
5     | -5.7      | 1.443     | 1.890
6     | -5.7      | 1.287     | 1.875
7     | -5.7      | 7.046     | 9.496
8     | -5.5      | 5.897     | 9.278
9     | -5.5      | 6.325     | 8.885

Writing output ... done.
```

## 12. Chenodeoxycholic acid 3-sulfate

```
Log6m0jE_21252312 - Notepad
File Edit Format View Help
Reading input ... done.
Setting up the scoring function ... done.
Analyzing the binding site ... done.
Using random seed: 256506792
Performing search ... done.
Refining results ... done.

mode | affinity | dist from best mode
      | (kcal/mol) | rmsd l.b. | rmsd u.b.
-----+-----+-----+-----
1     | -7.0      | 0.000     | 0.000
2     | -6.3      | 3.787     | 5.825
3     | -6.2      | 3.722     | 9.877
4     | -6.1      | 5.641     | 12.145
5     | -6.0      | 3.504     | 9.373
6     | -6.0      | 5.658     | 8.437
7     | -6.0      | 5.195     | 9.010
8     | -5.9      | 5.506     | 8.625
9     | -5.9      | 5.011     | 9.017

Writing output ... done.
```

## Lampiran 6. Log Penambatan ACE2 Autodock Vina

### 1. Remdesivir

```
Log6m0jA_Remdesivir - Notepad
File Edit Format View Help
Reading input ... done.
Setting up the scoring function ... done.
Analyzing the binding site ... done.
Using random seed: 639636848
Performing search ... done.
Refining results ... done.

mode | affinity | dist from best mode
      | (kcal/mol) | rmsd l.b. | rmsd u.b.
-----+-----+-----+-----
1      -6.5      0.000      0.000
2      -6.2      2.822      10.256
3      -5.9      4.432      8.643
4      -5.8      2.968      6.261
5      -5.8      1.582      2.160
6      -5.8      1.283      2.155
7      -5.7      2.538      5.589
8      -5.7      1.973      3.703
9      -5.7      13.307     15.449
Writing output ... done.
```

### 2. Nelfinavir

```
Log6m0jA_Nelfinavir - Notepad
File Edit Format View Help
Reading input ... done.
Setting up the scoring function ... done.
Analyzing the binding site ... done.
Using random seed: 303451112
Performing search ... done.
Refining results ... done.

mode | affinity | dist from best mode
      | (kcal/mol) | rmsd l.b. | rmsd u.b.
-----+-----+-----+-----
1      -6.4      0.000      0.000
2      -6.0      16.979     20.760
3      -6.0      3.122      6.130
4      -6.0      16.548     19.720
5      -5.9      16.031     19.723
6      -5.8      17.594     21.393
7      -5.8      3.713      8.309
8      -5.8      2.374      8.795
9      -5.8      2.915      8.480
Writing output ... done.
```

### 3. Hydroxychloroquine

```
Log6m0jA_Hydroxychloroquine - Notepad
File Edit Format View Help
Reading input ... done.
Setting up the scoring function ... done.
Analyzing the binding site ... done.
Using random seed: -1428733264
Performing search ... done.
Refining results ... done.

mode | affinity | dist from best mode
      | (kcal/mol) | rmsd l.b. | rmsd u.b.
-----+-----+-----+-----
1      -6.4      0.000      0.000
2      -6.2      1.753      2.305
3      -6.1      3.341      5.439
4      -5.8      5.226      8.125
5      -5.8      3.398      7.807
6      -5.8      2.994      4.193
7      -5.7      4.320      8.678
8      -5.6      3.316      4.992
9      -5.6      3.418      7.636
Writing output ... done.
```

### 4. Molnupiravir

```
log6m0jA_Molnupiravir - Notepad
File Edit Format View Help
Reading input ... done.
Setting up the scoring function ... done.
Analyzing the binding site ... done.
Using random seed: -1633839400
Performing search ... done.
Refining results ... done.

mode | affinity | dist from best mode
      | (kcal/mol) | rmsd l.b. | rmsd u.b.
-----+-----+-----+-----
1      -6.8      0.000      0.000
2      -6.6      2.684      5.982
3      -6.5      2.890      6.116
4      -6.3      3.780      6.372
5      -6.1      3.696      6.804
6      -5.7      4.019      7.112
7      -5.7      15.849     19.000
8      -5.7      15.092     18.124
9      -5.6      15.440     16.917
Writing output ... done.
```

### 5. 1beta-Hydroxycholeic acid

```
Log6m0jA_5283893 - Notepad
File Edit Format View Help
Reading input ... done.
Setting up the scoring function ... done.
Analyzing the binding site ... done.
Using random seed: -458512792
Performing search ... done.
Refining results ... done.

mode | affinity | dist from best mode
      | (kcal/mol) | rmsd l.b. | rmsd u.b.
-----+-----+-----+-----
1      -6.2      0.000      0.000
2      -6.2      14.698     19.419
3      -6.0      15.395     19.248
4      -5.9      12.473     15.326
5      -5.9      15.655     18.900
6      -5.8      15.010     17.626
7      -5.7      14.587     17.946
8      -5.6      15.075     19.078
9      -5.5      14.306     17.841
Writing output ... done.
```

### 6. 4,4-dimethyl-5alpha-cholesta-8,14,24-trien-3beta-ol

```
Log6m0jA_443212 - Notepad
File Edit Format View Help
Reading input ... done.
Setting up the scoring function ... done.
Analyzing the binding site ... done.
Using random seed: 512043268
Performing search ... done.
Refining results ... done.

mode | affinity | dist from best mode
      | (kcal/mol) | rmsd l.b. | rmsd u.b.
-----+-----+-----+-----
1      -7.3      0.000      0.000
2      -6.2      2.718      8.389
3      -6.2      14.890     19.759
4      -6.2      3.668      7.068
5      -5.9      15.221     17.428
6      -5.9      14.241     16.778
7      -5.9      13.291     16.198
8      -5.8      14.740     16.258
9      -5.8      19.534     25.177
Writing output ... done.
```

## 7. Carboprost

```
Log6m0jA_5281075 - Notepad
File Edit Format View Help
Reading input ... done.
Setting up the scoring function ... done.
Analyzing the binding site ... done.
Using random seed: -1214675844
Performing search ... done.
Refining results ... done.
```

mode	affinity (kcal/mol)	dist from best mode rmsd l.b.   rmsd u.b.
1	-7.6	0.000   0.000
2	-6.4	2.274   2.910
3	-6.3	2.847   4.816
4	-5.6	3.225   5.296
5	-5.5	4.502   8.002
6	-5.4	15.774   21.483
7	-5.3	14.760   20.514
8	-5.3	13.764   19.233
9	-5.2	3.476   4.809

Writing output ... done.

## 9. Androstanolone

```
Log6m0jA_10635 - Notepad
File Edit Format View Help
Reading input ... done.
Setting up the scoring function ... done.
Analyzing the binding site ... done.
Using random seed: 1468768272
Performing search ... done.
Refining results ... done.
```

mode	affinity (kcal/mol)	dist from best mode rmsd l.b.   rmsd u.b.
1	-6.9	0.000   0.000
2	-6.6	6.409   8.877
3	-6.5	0.908   6.647
4	-6.4	6.844   9.990
5	-6.1	3.867   8.317
6	-5.9	2.957   7.602
7	-5.7	6.269   9.002
8	-5.6	2.181   6.951
9	-5.6	13.776   15.233

Writing output ... done.

## 11. Androst-4-en-3-one

```
Log6m0jA_250294 - Notepad
File Edit Format View Help
Reading input ... done.
Setting up the scoring function ... done.
Analyzing the binding site ... done.
Using random seed: 1233320584
Performing search ... done.
Refining results ... done.
```

mode	affinity (kcal/mol)	dist from best mode rmsd l.b.   rmsd u.b.
1	-6.9	0.000   0.000
2	-6.7	8.026   10.270
3	-6.4	4.039   7.416
4	-6.3	2.885   6.044
5	-6.2	6.932   9.367
6	-6.2	13.343   15.368
7	-6.2	2.738   5.110
8	-5.9	4.487   7.627
9	-5.9	18.010   20.528

Writing output ... done.

## 8. 6-acetyloxy-7-ethenyl-1,4a,7-trimethyl-3,4,4b,5,6,8,10,10a-octahydro-2H-phenanthrene-1-carboxylic acid

```
Log6m0jA_51694232 - Notepad
File Edit Format View Help
Reading input ... done.
Setting up the scoring function ... done.
Analyzing the binding site ... done.
Using random seed: 424532720
Performing search ... done.
Refining results ... done.
```

mode	affinity (kcal/mol)	dist from best mode rmsd l.b.   rmsd u.b.
1	-6.1	0.000   0.000
2	-5.7	2.356   4.524
3	-5.2	13.958   17.437
4	-4.9	6.680   11.205
5	-4.9	6.172   9.570
6	-4.9	13.322   15.824
7	-4.9	5.395   9.104
8	-4.8	12.286   15.067
9	-4.8	12.076   14.318

Writing output ... done.

## 10. 4-[(2,4-Dihydroxyphenyl)diazenyl]benzenesulfonic acid

```
Log6m0jA_11035 - Notepad
File Edit Format View Help
Reading input ... done.
Setting up the scoring function ... done.
Analyzing the binding site ... done.
Using random seed: -298365312
Performing search ... done.
Refining results ... done.
```

mode	affinity (kcal/mol)	dist from best mode rmsd l.b.   rmsd u.b.
1	-6.0	0.000   0.000
2	-5.7	3.478   8.120
3	-5.7	14.229   15.175
4	-5.5	14.139   15.390
5	-5.5	14.327   16.840
6	-5.4	16.854   18.423
7	-5.4	14.493   17.813
8	-5.4	14.203   15.157
9	-5.3	1.820   2.182

Writing output ... done.

## 12. Chenodeoxycholic acid 3-sulfate

```
Log6m0jA_21252312 - Notepad
File Edit Format View Help
Reading input ... done.
Setting up the scoring function ... done.
Analyzing the binding site ... done.
Using random seed: -1817535500
Performing search ... done.
Refining results ... done.
```

mode	affinity (kcal/mol)	dist from best mode rmsd l.b.   rmsd u.b.
1	-6.2	0.000   0.000
2	-6.1	4.771   10.879
3	-6.0	3.165   8.752
4	-6.0	16.756   18.076
5	-5.9	3.712   9.294
6	-5.9	2.739   4.023
7	-5.7	4.142   9.042
8	-5.7	2.903   8.956
9	-5.7	3.382   9.433

Writing output ... done.