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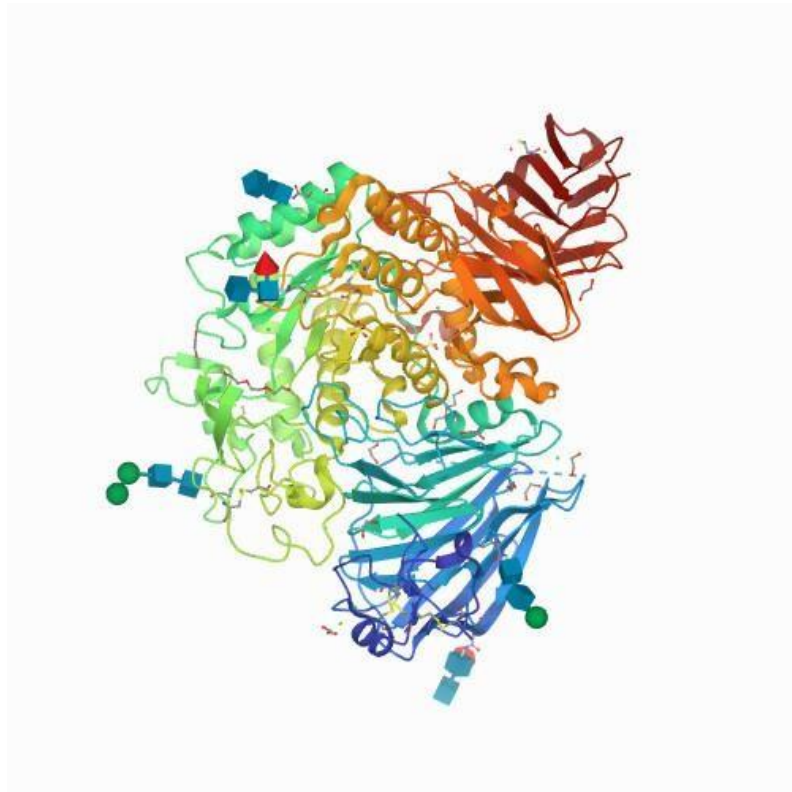
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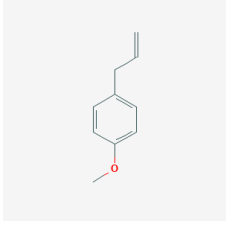
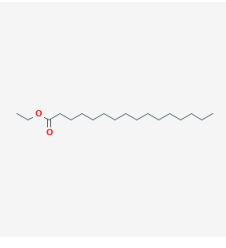
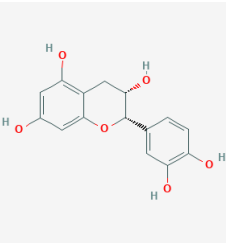
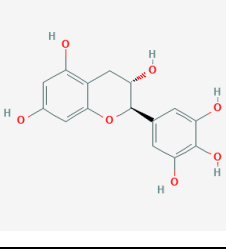
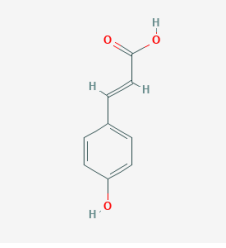
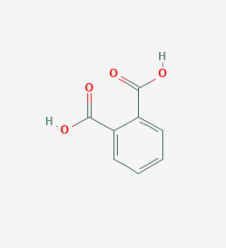
## **LAMPIRAN**

**Lampiran 1. Struktur Protein (Reseptor)**

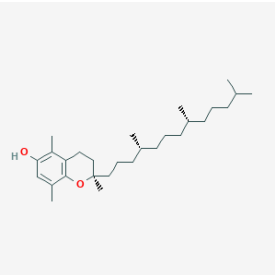
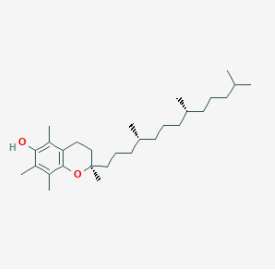
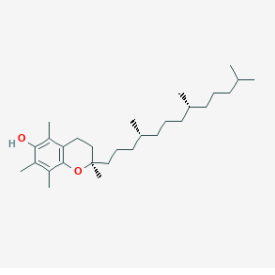
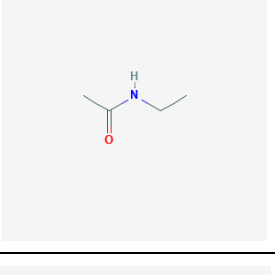
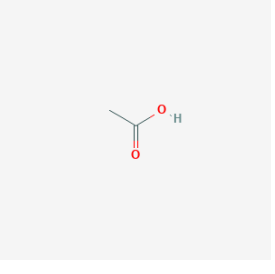
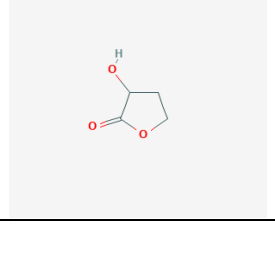


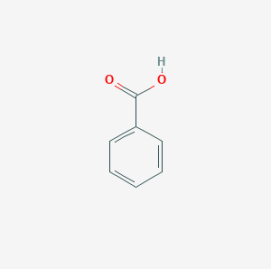
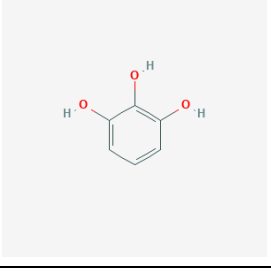
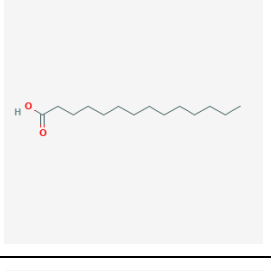
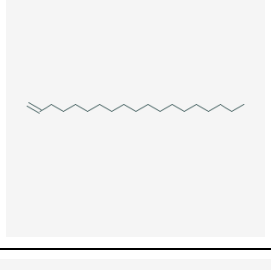
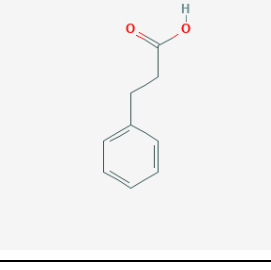
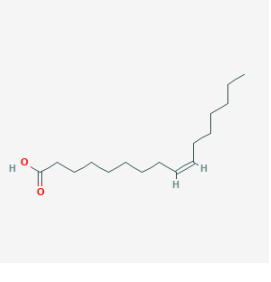
Struktur 3D dengan PDB ID 5NN3 (sumber : [www.rcsb.org](http://www.rcsb.org))

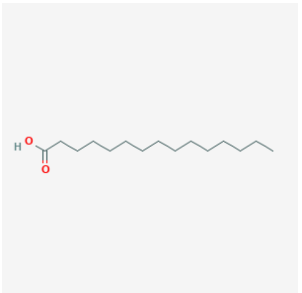
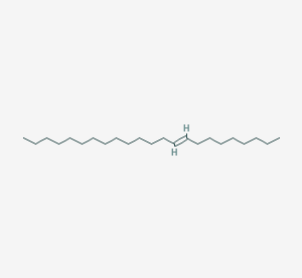
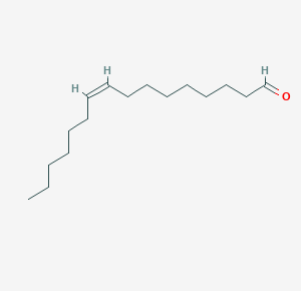
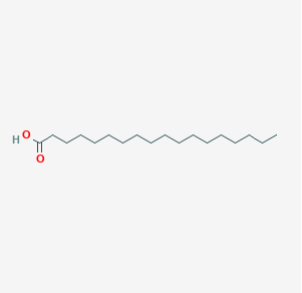

**Lampiran 2. Struktur Ligan Uji Ekstrak Kulit Pisang**

No .	Ligan	Struktur
1	<i>Estragole</i>	
2	<i>Hexadecanoic acid ethyl ester</i>	
3	<i>Epicatechin</i>	
4	<i>Gallocatechin</i>	
5	<i>p-coumaric acid methyl ester</i>	
6	<i>1,2-Benzenedicarboxylic acid mono (2-ethylhexylester)</i>	



7	<i>Beta-tocopherol</i>	
8	<i>Vitamin E</i>	
9	<i>Ethanimidic acid, ethyl ester</i>	
10	<i>Acetic acid</i>	
11	<i>2-hydroxy-gammbutyrolactone</i>	
12	<i>4H-pyran-4-one, 3 hidroxy</i>	

13	<i>Benzoic acid</i>	
14	<i>Pyrogallol</i>	
15	<i>Tetradecenoic acid</i>	
16	<i>1-Nonadecene</i>	
17	<i>Benzenepropanoic acid</i>	
18	<i>Cis-9-Hexadecenoic acid</i>	

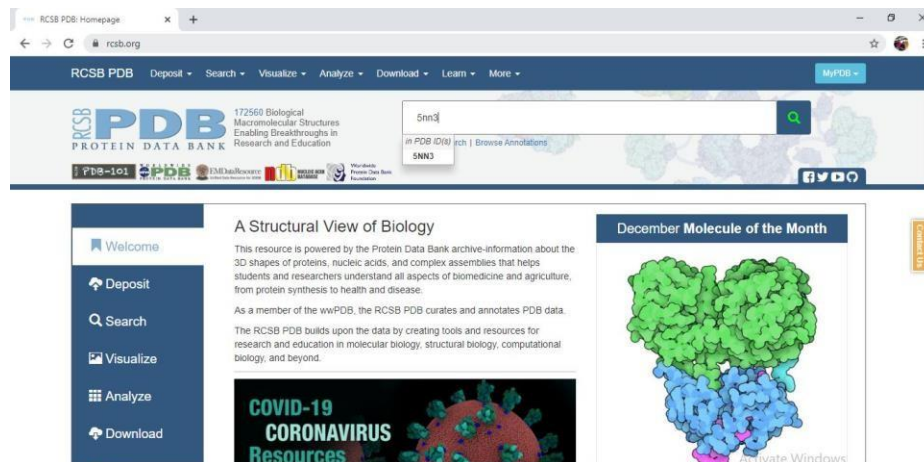
19	<i>Pentadecanoic acid</i>	
20	<i>9-Tricosene</i>	
21	<i>Cis-9-Hexadecenal</i>	
22	<i>Octadecanoic acid</i>	
23	<i>2,6,10,14,18,22 Tetracosahexaene</i>	

**Lampiran 3.** Hasil Pengaturan *grid box* reseptor

<i>Grid Box</i>						
	<i>Center</i>			<i>Size</i>		
<b>Makromolekul</b>	<b>X</b>	<b>Y</b>	<b>Z</b>	<b>X</b>	<b>Y</b>	<b>Z</b>
<i>α</i> -glukosidase (kode 5nn3)	4,116	-13,936	75,412	14	16	14

## Lampiran 4. Preparasi Struktur Reseptor

1. Struktur reseptor yang digunakan dapat diunduh dari <https://www.rcsb.org>



2. Reseptor  $\alpha$ -Glukosidase (kode : 5NN3) dicari pada *search engine* dan diunduh dalam format PDB

**5NN3**  
Crystal structure of human lysosomal acid-alpha-glucosidase

DOI: 10.2210/pdb5NN3/pdb

**Classification:** HYDROLASE  
**Organism(s):** Homo sapiens  
**Expression System:** Cricetulus griseus  
**Mutation(s):** Yes

**Deposited:** 2017-04-08 **Released:** 2017-10-25  
**Deposition Author(s):** Roig-Zamboni, V., Cobucci-Ponzano, B., Iacono, R., Bourne, Y., Moracci, M.

**Experimental Data Snapshot**  
**Method:** X-RAY DIFFRACTION  
**Resolution:** 1.90 Å  
**R-Value Free:** 0.189  
**R-Value Work:** 0.162  
**R-Value Observed:** 0.163

**wwPDB Validation**

Metric	Value
Rfree	0.189
Ciathscore	0.162
Ramachandran outliers	0.162
Sidechain outliers	0.162
RSRZ outliers	0.162

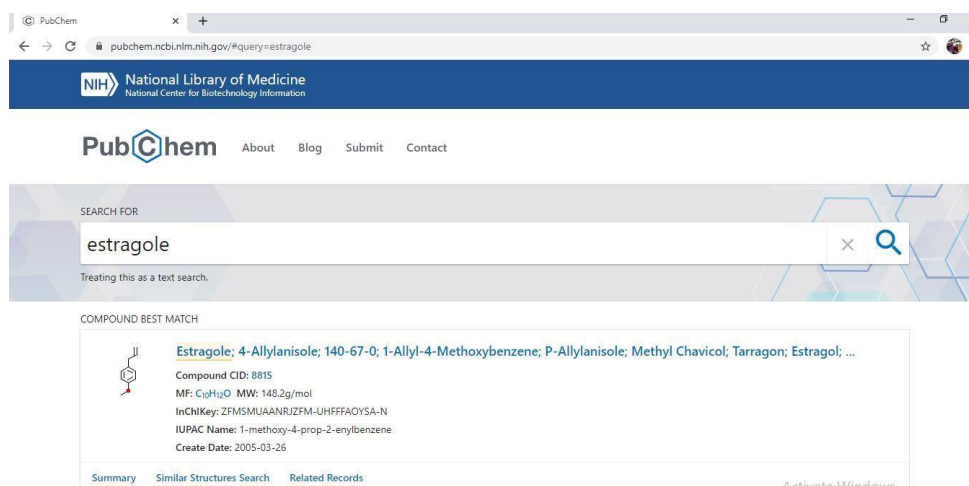
Display Files | Download Files

- FASTA Sequence
- PDB Format
- PDB Format (gz)
- PDBx/mmCIF Format
- PDBx/mmCIF Format (gz)
- PDBMLXML Format (gz)
- Biological Assembly 1
- Structure Factors (CIF)
- Structure Factors (CIF - gz)
- Validation Full PDF
- Validation XML
- f0-fc Map (DSN6)
- 2f0-fc Map (DSN6)

## Lampiran 5. Preparasi Struktur Ligan

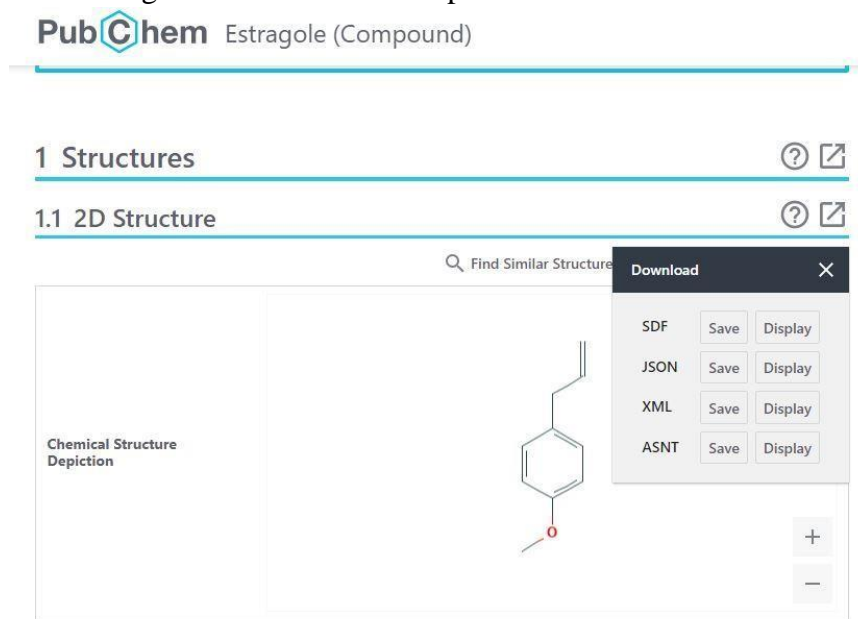
1. Ligan ekstrak kulit pisang yang digunakan disimpan dari situs

<http://pubchem.ncbi.nlm.nih.gov>



The screenshot shows the PubChem website interface. At the top, there is a search bar with the text "estragole" entered. Below the search bar, the results section is titled "COMPOUND BEST MATCH". It displays a small chemical structure icon and the following information: "Estragole; 4-Allylanisole; 140-67-0; 1-Allyl-4-Methoxybenzene; P-Allylanisole; Methyl Chavicol; Tarragon; Estragol; ...". Below this, it lists "Compound CID: 8815", "MF: C<sub>10</sub>H<sub>12</sub>O MW: 148.2g/mol", "InChIKey: ZFMSMUAANRIZFM-UHFFFAOYSA-N", "IUPAC Name: 1-methoxy-4-prop-2-enylbenzene", and "Create Date: 2005-03-26". At the bottom of the results box, there are links for "Summary", "Similar Structures Search", and "Related Records".

2. Struktur ligan 2D dicari dan disimpan dalam format SDF

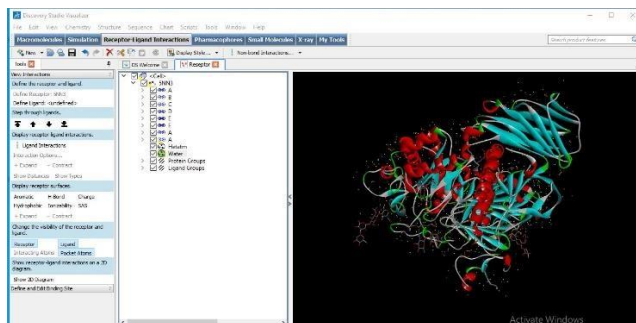


The screenshot shows the PubChem page for Estragole (Compound). The page title is "PubChem Estragole (Compound)". Below the title, there are navigation tabs for "1 Structures" and "1.1 2D Structure". The "1.1 2D Structure" tab is selected. The main content area shows the chemical structure of Estragole, which is 1-methoxy-4-prop-2-enylbenzene. To the right of the structure, there is a "Download" menu with options for SDF, JSON, XML, and ASNT. Each option has "Save" and "Display" buttons. The SDF option is highlighted. Below the structure, there are zoom in (+) and zoom out (-) buttons.

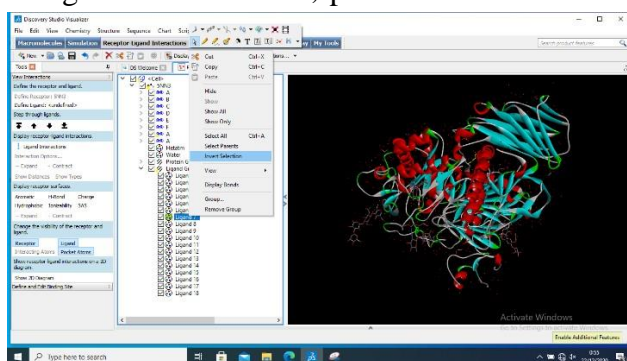
## Lampiran 6. Pemisahan Ligan dan Makromolekul

### 1. Pemilihan Ligan

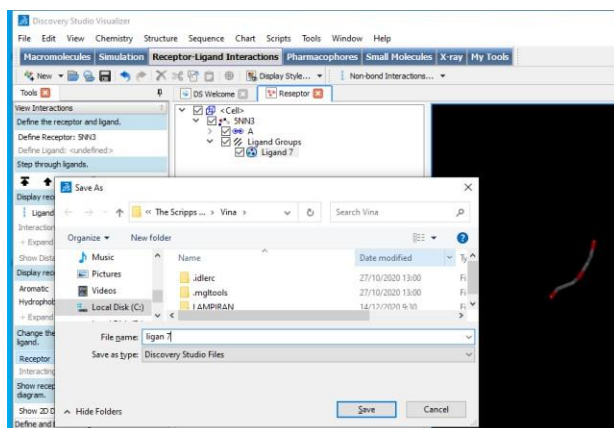
- a. Struktur 3D reseptor dalam format PDB yang diunduh dibuka dengan *discovery studio visualizer* kemudian munculkan *view hierarchy* dengan mengetik *ctrl+H* pada *keyboard*.



- b. Setelah muncul struktur reseptor lakukan penghilangan molekul air dengan meng-klik kanan *mouse* lalu pilih *delete*. Kemudian, lakukan pemilihan ligan dengan meng-klik kanan *mouse*, pilih *invert selection* dan tekan *delete*.

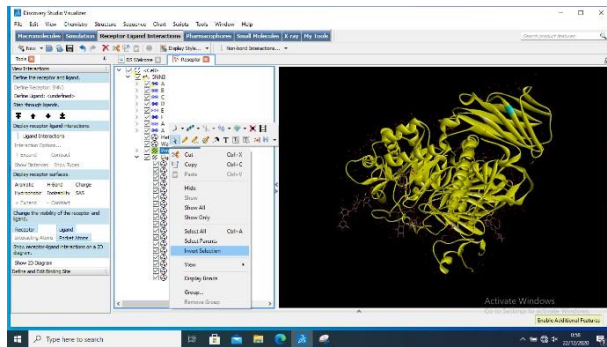


- c. Simpan file dengan format *\*.pdb*.

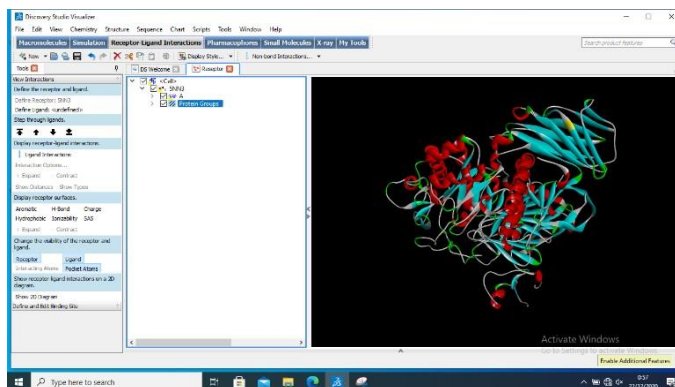


## 2. Pemilihan Makromolekul

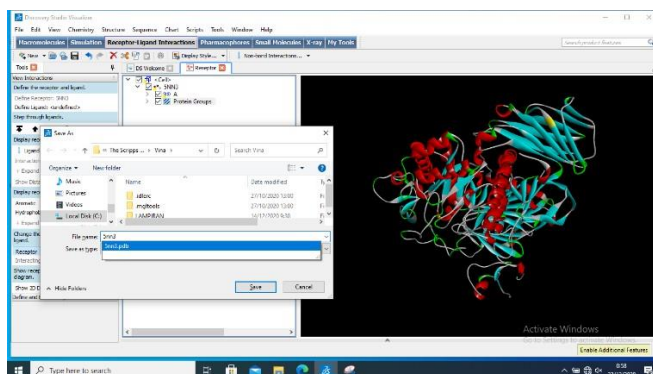
### a. Hal yang sama dilakukan hingga 1b



b. Langkah selanjutnya pilih unsur protein yang ada dengan meng-klik kanan, pilih *invert selection*, dan tekan tombol *delete*.



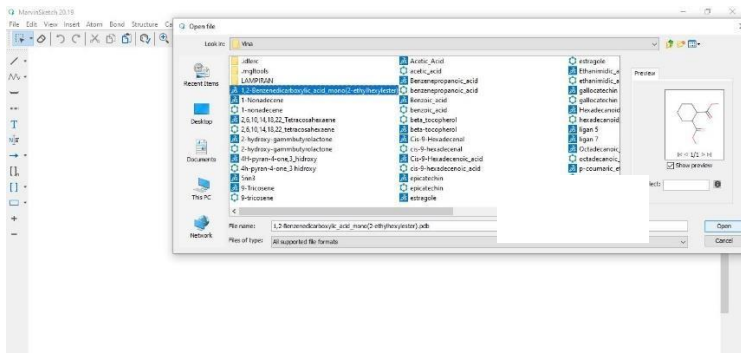
c. Kemudian, simpan *file* dengan format *\*.pdb*.



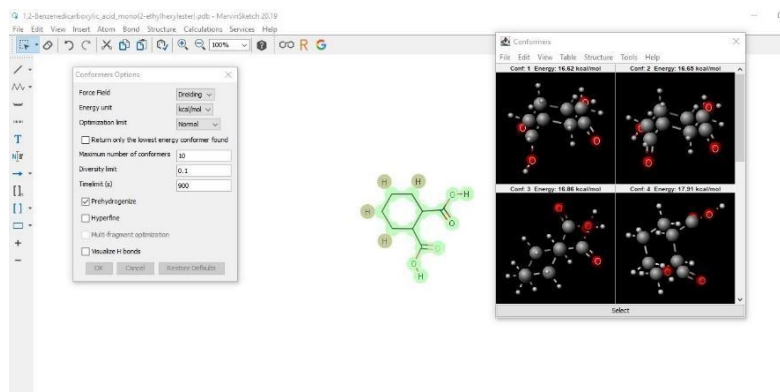


## Lampiran 7. Pengubahan Struktur Ligan 2D menjadi 3D

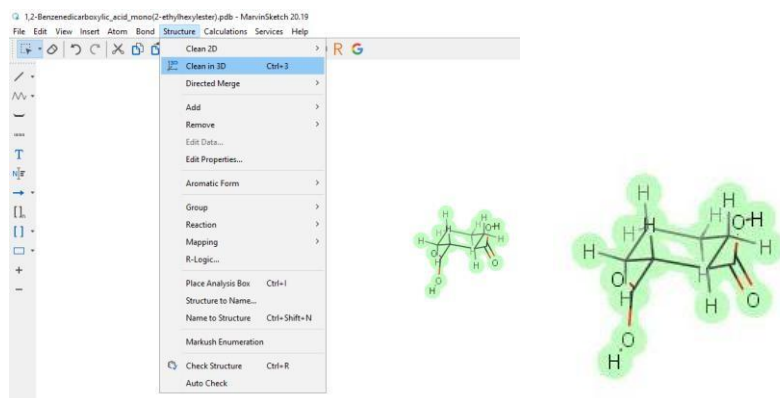
1. Pilih ligan yang telah diunduh dengan klik *open file* menggunakan *software* Marvin Sketch.



2. Blok struktur ligan, klik *calculation > conformation > conformers*, tekan ok. Setelah muncul pilih konformasi dengan energi yang paling rendah.

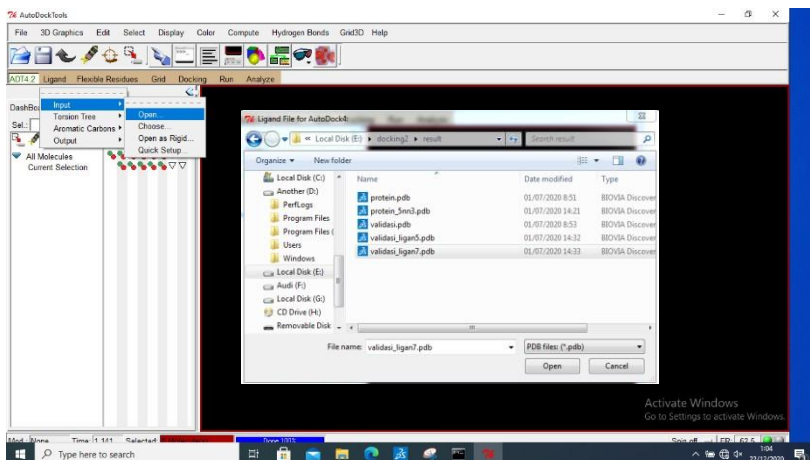


3. Setelah muncul blok kembali seluruh strukturnya, pilih *structure > clean in 3D* dan file disimpan dengan format \*.pdb.

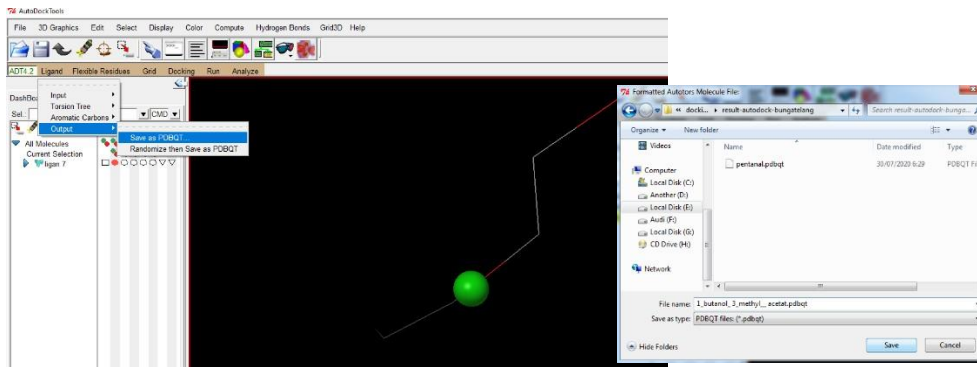


## Lampiran 8. Validasi *molecular docking*

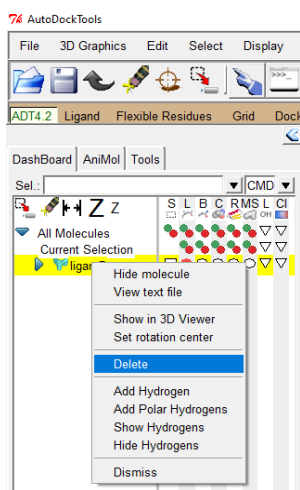
1. Buka struktur ligan dengan *software* Autodock Tools dengan klik *ligand* > *input* > *open*.



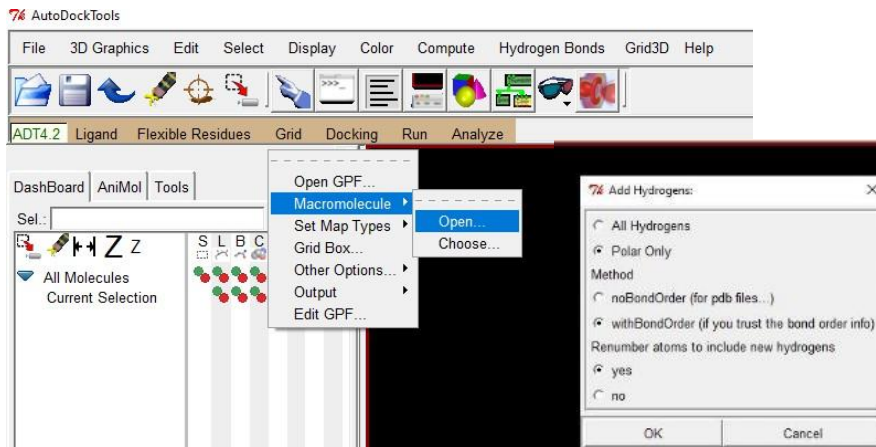
2. Dalam autodock tools struktur ligan otomatis diberi muatan *gasteiger* dan *choose torsio* sehingga dapat langsung disimpan dengan klik *Ligand* > *Output* > *Save as PDBQT*



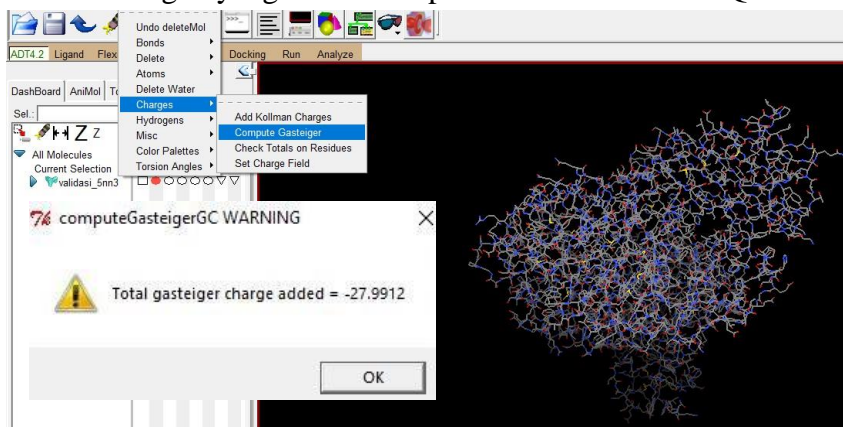
3. Setelah struktur ligan tersimpan, hapus struktur ligan pada *software* dengan klik kanan > *delete*.



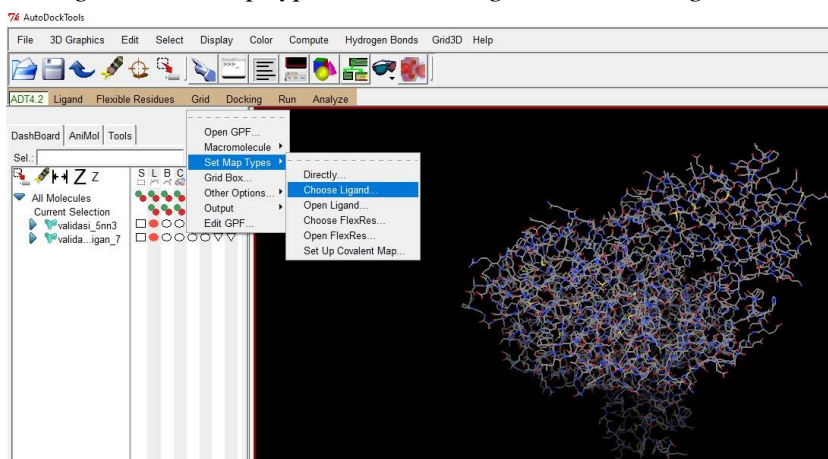
4. Buka struktur reseptor dengan meng-klik *grid* > *makromolekul* > *open*. Kemudian, tambahkan hidrogen polar dengan memilih menu *edit* > *Hydrogens* > *Add* > klik menu *Polar Only*.



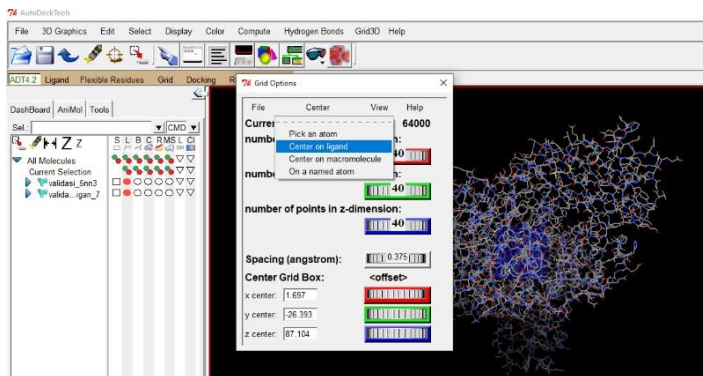
- Hitung muatan *gasteiger* dengan klik edit > charges > compute *gasteiger*. Kemudian masukkan ligan yang telah disimpan dalam format PDBQT.



- Pilih *grid* > set map types > choose ligand > select ligand



7. Tentukan *gridbox* dengan pilih *grid > grid box > center > center on ligand*



8. *Gridbox* diatur agar seluruh *box* menutupi ligan, lalu hasil validasi penambatan diketik dalam *notepad* dan disimpan dengan format \*.txt serta masukkan dalam folder vina

```

conf_rigid - Notepad
File Edit Format View Help
receptor =validasi_5nn3.pdbqt
ligand =vitamin_e.pdbqt

cpu =3

out =out_vitamin_e.pdbqt
log =log_vitamin_e.txt

center_x =4.116
center_y =-13.936
center_z =75.412

size_x =14
size_y =16
size_z =14

exhaustiveness =256
    
```

9. Parameter penambatan dilakukan *running* dengan aplikasi *command prompt (Run as Administrator)* lalu buka file konfigurasi yang ada di folder vina.

```

Administrator: C:\Windows\system32\cmd.exe
Microsoft Windows [Version 10.0.18363.1198]
(c) 2019 Microsoft Corporation. All rights reserved.

C:\Users\Amelya Putri Zamzami>cd C:\Vina\Program Files (x86)\The Scripps Research Institute\Vina
C:\Vina\Program Files (x86)\The Scripps Research Institute\Vina>vina.exe --config conf_rigid.txt
    
```

10. Tunggu proses *running* selesai. Contoh output dalam format \*.txt.

```

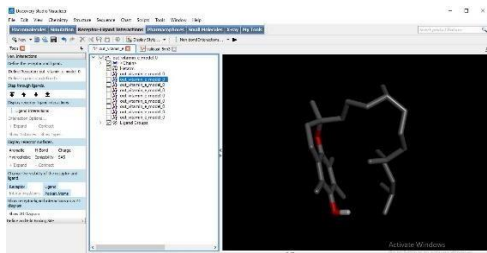
log_vitamin_e - Notepad
File Edit Format View Help
#####
# If you used AutoDock Vina in your work, please cite:
#
# O. Trott, A. J. Olson,
# AutoDock Vina: improving the speed and accuracy of docking
# with a new scoring function, efficient optimization and
# multithreading, Journal of Computational Chemistry 31 (2010)
# 455-461
# DOI 10.1002/jcc.21334
#
# Please see http://vina.scripps.edu for more information.
#####
Reading input ... done.
Setting up the scoring function ... done.
Analyzing the binding site ... done.
Using random seed: 1940121684
Performing search ... done.
Refining results ... done.

mode | affinity | dist from best mode
      | (kcal/mol) | rmsd l.b. | rmsd u.b.
-----|-----|-----|-----
1      | -5.5      | 0.000     | 0.000
2      | -5.5      | 2.086     | 5.098
3      | -5.5      | 2.765     | 6.136
4      | -5.4      | 1.441     | 2.615
5      | -5.4      | 1.850     | 4.952
6      | -5.4      | 2.700     | 5.967
7      | -5.4      | 2.105     | 5.339
8      | -5.4      | 2.691     | 6.952
9      | -5.4      | 1.814     | 6.261

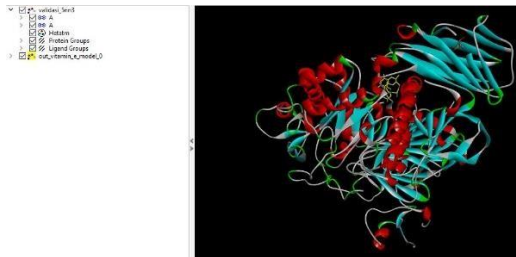
Writing output ... done.
    
```

## Lampiran 9. Visualisasi menggunakan *biovia discovery studio visualizer*

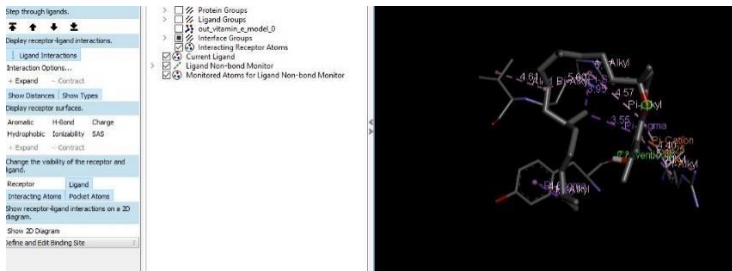
1. *File* reseptor dan hasil *out* ligan dalam format \*.pdbqt dibuka.



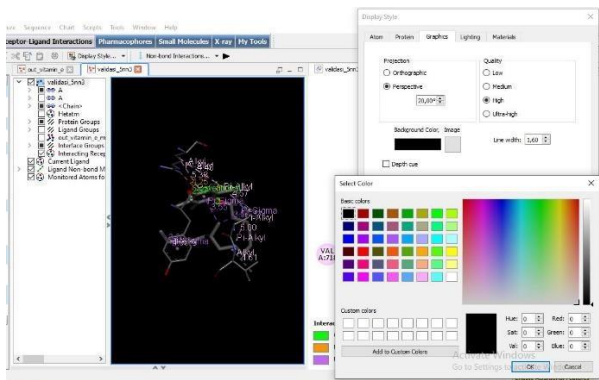
2. Hasil *out* ligan dipilih mode kedua lalu klik kanan > *Invert selection* > *delete*. *Out* ligan yang dipilih di *cut* dan disalin ke dalam struktur reseptor.



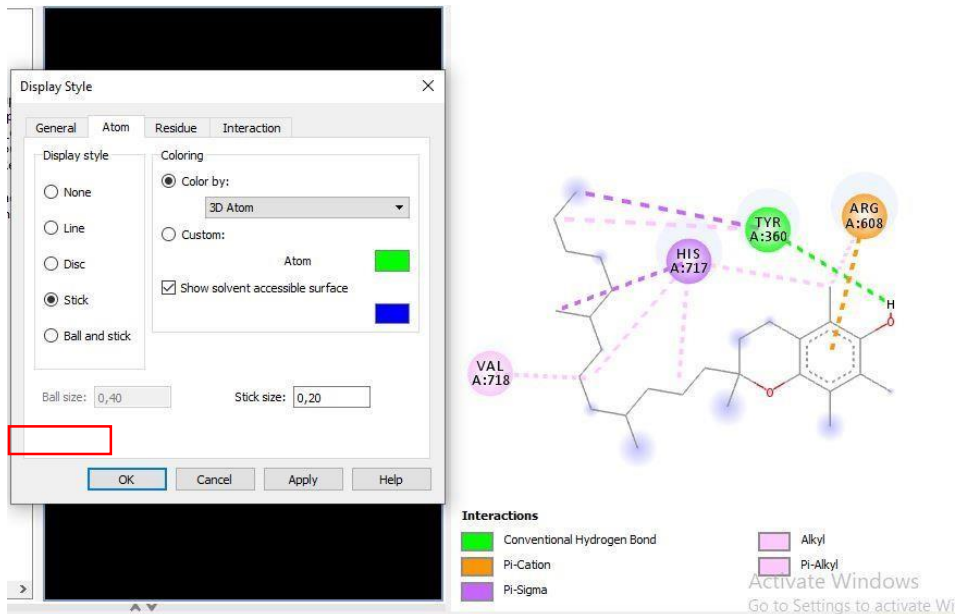
3. Untuk melihat interaksi ligan dengan reseptor pilih *Ligand Interaction* > *Show Distance* > *Show Types*.



4. Agar interaksi antara ligan dengan reseptor dapat terlihat ubah *background* menjadi putih dengan klik kanan > *display style* > *graphics* > *background*.

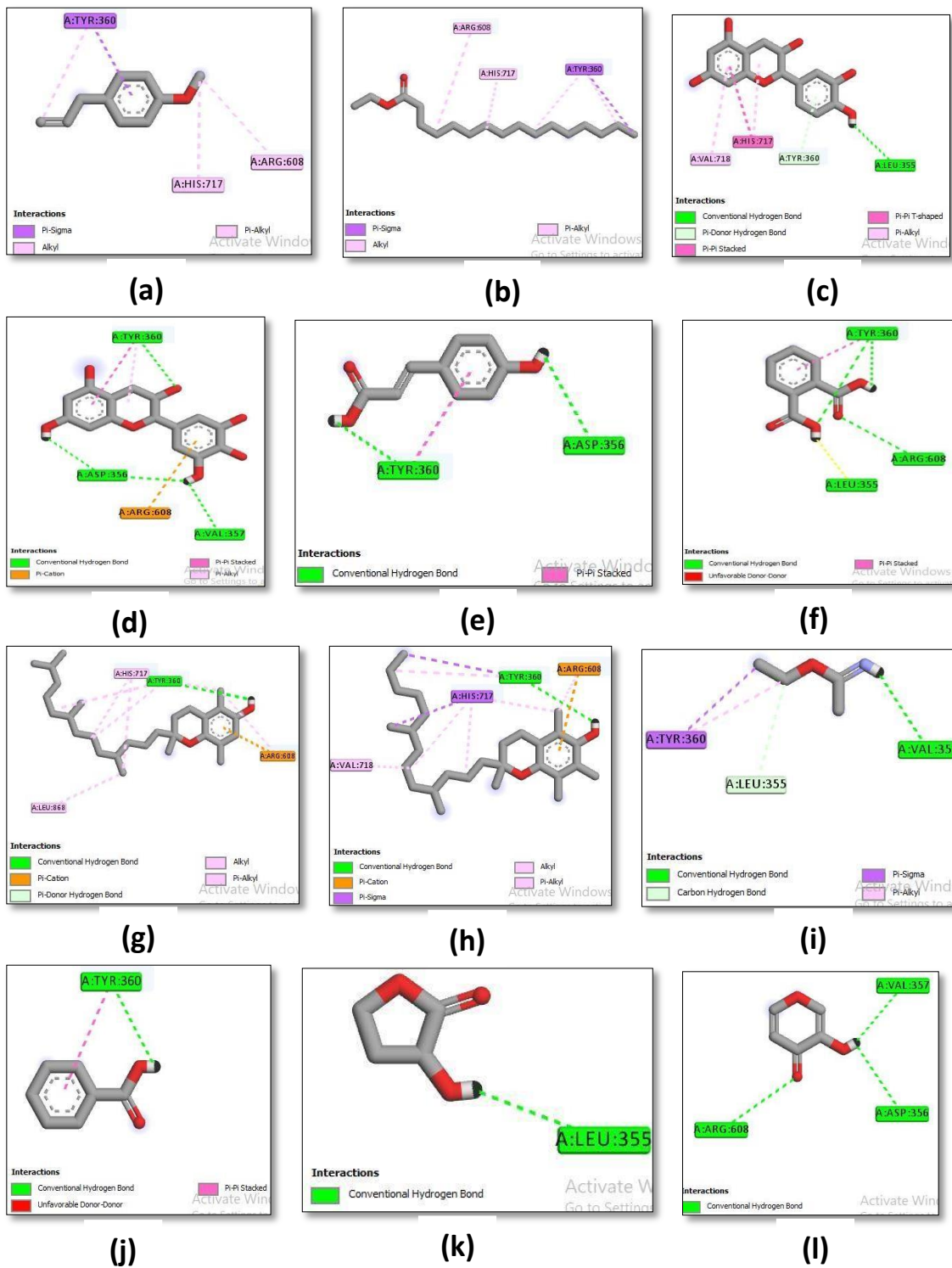


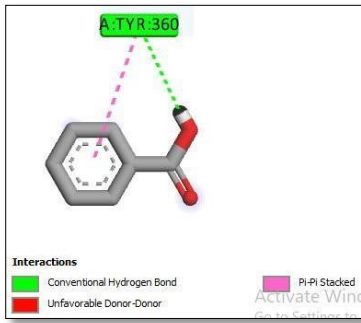
5. Agar visualisasi interaksi reseptor dan ligan lebih jelas maka ubah menjadi 2D dengan pilih *show 2D diagram*. Lalu agar visualisasi struktur dan ikatan dapat terlihat lebih jelas klik kanan > *display style* > atom > *stick* lalu tekan ok.



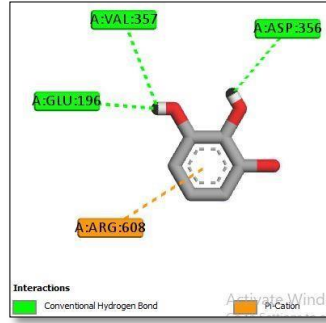


## Lampiran 10. Visualisasi 2D Interaksi Ligan dengan Reseptor

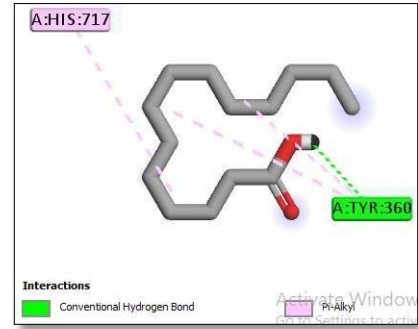




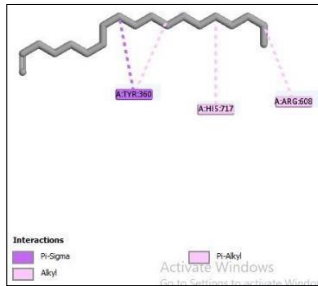
(m)



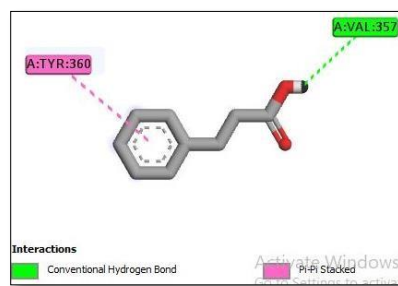
(n)



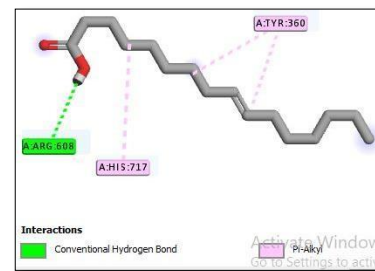
(o)



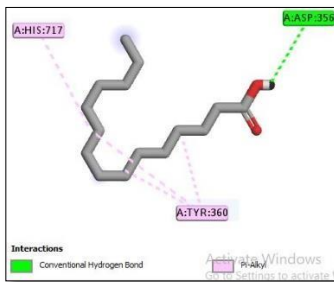
(p)



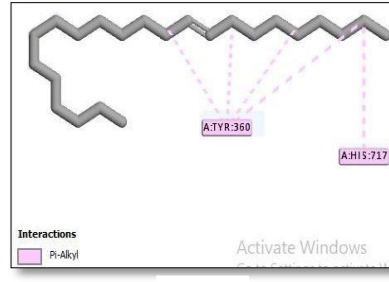
(q)



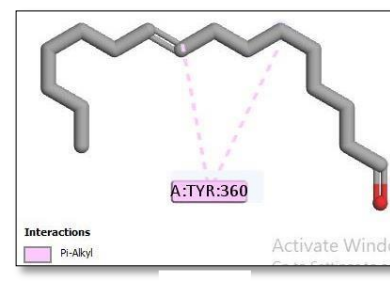
(r)



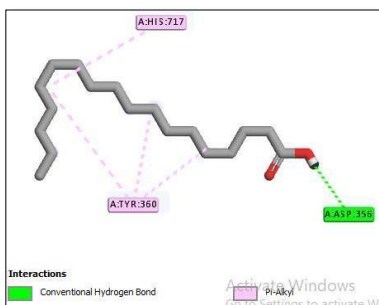
(s)



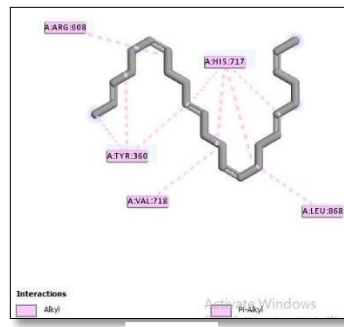
(t)



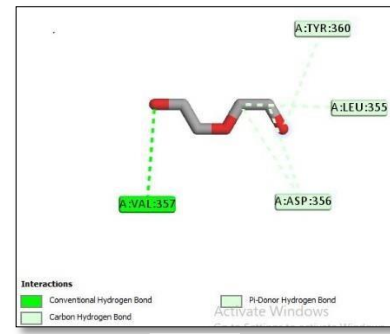
(u)



(v)



(w)



(x)



**Keterangan :**

<b>Kode</b>	<b>Ligan uji senyawa ekstrak kulit pisang</b>
a	<i>Estragole</i>
b	<i>Hexadecanoic acid ethyl ester</i>
c	<i>Epicatechin</i>
d	<i>Gallocatechin</i>
e	<i>p-coumaric acid methyl ester</i>
f	<i>1,2-Benzenedicarboxylic acid mono (2-ethylhexylester)</i>
g	<i>Beta-tocopherol</i>
h	<i>Vitamin E</i>
i	<i>Ethanimidic acid, ethyl ester</i>
j	<i>Acetic acid</i>
k	<i>2-hydroxy-gammbutyrolactone</i>
l	<i>4H-pyran-4-one, 3 hidroxy</i>
m	<i>Benzoic acid</i>
n	<i>Pyrogallol</i>
o	<i>Tetradecenoic acid</i>
p	<i>1-Nonadecene</i>
q	<i>Benzenepropanoic acid</i>
r	<i>Cis-9-Hexadecenoic acid</i>
s	<i>Pentadecanoic acid</i>
t	<i>9-Tricosene</i>
u	<i>Cis-9-Hexadecenal</i>
v	<i>Octadecanoic acid</i>
w	<i>2,6,10,14,18,22 Tetracosahexaene</i>
x	<i>Ligan 7 Validasi</i>

## Lampiran 11. Data Hasil *Docking Vina*

### a. Ligan 7 Validasi

Mode	Affinity (kcal/mol)	RMSD l b	RMSD u b
1	-2,4	0,000	0,000
2	-2,4	0,108	4,086
3	-2,4	1,582	4,170
4	-2,4	1,094	4,212
5	-2,4	6,641	7,089
6	-2,4	1,937	2,260
7	-2,4	1,583	1,711
8	-2,4	6,620	8,105
9	-2,4	1,620	3,754

### b. Estragole

Mode	Affinity (kcal/mol)	RMSD l b	RMSD u b
1	-4,5	0,000	0,000
2	-4,5	2,899	4,049
3	-4,4	2,875	4,678
4	-4,4	2,824	5,197
5	-4,4	2,790	5,329
6	-4,4	2,892	4,122
7	-4,4	2,270	3,629
8	-4,4	2,901	4,768
9	-4,4	3,029	4,932

### c. Hexadecanoic acid ethyl ester

Mode	Affinity (kcal/mol)	RMSD l b	RMSD u b
1	-4,1	0,000	0,000
2	-4,1	2,198	5,437
3	-4,0	1,243	4,018
4	-4,0	2,617	5,182
5	-4,0	1,389	4,765
6	-4,0	2,611	5,932
7	-4,0	2,224	5,267
8	-4,0	1,754	4,080
9	-3,9	1,675	3,992

### d. Epicatechin

Mode	Affinity (kcal/mol)	RMSD l b	RMSD u b
1	-6,6	0,000	0,000
2	-6,4	2,628	6,420
3	-6,3	2,804	5,906
4	-6,3	1,846	2,619
5	-6,2	2,868	5,977
6	-6,2	2,803	5,926
7	-6,2	2,178	2,934
8	-6,2	3,039	4,609
9	-6,2	2,563	6,542

### e. Galocatechin

Mode	Affinity (kcal/mol)	RMSD l b	RMSD u b
1	-6,4	0,000	0,000
2	-6,4	0,913	1,956
3	-6,3	2,482	5,356
4	-6,2	1,116	1,549
5	-6,1	2,460	5,118
6	-6,0	2,105	4,843
7	-6,0	2,074	4,559
8	-6,0	4,184	6,499
9	-6,0	1,742	5,742

### f. *p*-coumaric acid methyl ester

Mode	Affinity (kcal/mol)	RMSD l b	RMSD u b
1	-5,2	0,000	0,000
2	-5,2	0,316	1,436
3	-5,1	3,099	3,660
4	-5,1	3,114	3,730
5	-5,0	1,192	1,836
6	-4,9	1,270	1,485
7	-4,8	2,915	5,051
8	-4,7	2,893	4,983
9	-4,7	2,860	4,901



*g. 1,2-Benzenedicarboxylic acid mono (2-ethylhexylester)*

Mode	Affinity (kcal/mol)	RMSD l b	RMSD u b
1	-5,1	0,000	0,000
2	-5,1	0,051	2,892
3	-4,9	1,958	3,120
4	-4,9	1,970	3,391
5	-4,8	2,588	3,512
6	-4,8	2,575	4,164
7	-4,7	1,995	3,182
8	-4,7	1,993	2,903
9	-4,6	1,954	3,757

*h. Beta-tocopherol*

Mode	Affinity (kcal/mol)	RMSD l b	RMSD u b
1	-5,5	0,000	0,000
2	-5,4	1,293	2,201
3	-5,3	2,897	7,442
4	-5,3	2,597	5,603
5	-5,3	1,950	4,458
6	-5,3	1,896	2,967
7	-5,3	2,081	5,165
8	-5,3	3,203	6,591
9	-5,3	2,794	5,573

*i. Vitamin E*

Mode	Affinity (kcal/mol)	RMSD l b	RMSD u b
1	-5,5	0,000	0,000
2	-5,5	2,086	5,098
3	-5,5	2,765	6,136
4	-5,4	1,441	2,615
5	-5,4	1,850	4,952
6	-5,4	2,700	5,967
7	-5,4	2,105	5,339
8	-5,4	2,691	6,952
9	-5,4	1,814	6,261

*j. Ethanimidic acid, ethyl ester*

Mode	Affinity (kcal/mol)	RMSD l b	RMSD u b
1	-3,1	0,000	0,000
2	-3,1	1,742	2,010
3	-2,9	3,309	3,624
4	-2,9	1,385	2,691
5	-2,9	7,066	7,284
6	-2,9	2,527	3,431
7	-2,9	2,359	3,293
8	-2,8	6,139	6,719
9	-2,8	6,239	6,650

*k. Acetic acid*

Mode	Affinity (kcal/mol)	RMSD l b	RMSD u b
1	-4,6	0,000	0,000
2	-4,6	0,057	1,623
3	-4,3	2,401	2,825
4	-4,3	3,494	4,558
5	-4,3	2,397	3,030
6	-4,2	2,308	3,187
7	-4,1	2,454	3,438
8	-4,1	1,884	2,573
9	-4,1	0,858	1,835

*l. 2-hydroxy-gammbutyrolactone*

Mode	Affinity (kcal/mol)	RMSD l b	RMSD u b
1	-3,3	0,000	0,000
2	-3,2	1,558	3,150
3	-3,2	3,381	3,779
4	-3,2	1,142	2,300
5	-3,1	1,944	3,465
6	-3,1	1,865	2,342
7	-3,1	1,478	2,472
8	-3,1	2,213	2,727
9	-3,0	1,767	2,642

m. *4H-pyran-4-one, 3 hidroxy*

Mode	Affinity (kcal/mol)	RMSD l b	RMSD u b
1	-3,5	0,000	0,000
2	-3,5	3,009	3,539
3	-3,5	1,578	1,906
4	-3,5	0,963	2,340
5	-3,4	2,342	3,359
6	-3,4	4,048	5,047
7	-3,4	2,732	3,850
8	-3,4	2,781	4,362
9	-3,4	1,882	2,685

n. *Benzoic acid*

Mode	Affinity (kcal/mol)	RMSD l b	RMSD u b
1	-4,7	0,000	0,000
2	-4,6	0,031	1,632
3	-4,3	3,510	4,291
4	-4,3	3,497	4,566
5	-4,3	2,412	2,845
6	-4,3	2,426	3,065
7	-4,2	0,904	1,907
8	-4,2	2,468	3,471
9	-4,1	2,466	3,415

o. *Pyrogallol*

Mode	Affinity (kcal/mol)	RMSD l b	RMSD u b
1	-4,2	0,000	0,000
2	-4,1	4,295	5,069
3	-4,1	4,293	5,198
4	-4,1	0,204	2,794
5	-4,1	4,245	5,099
6	-4,0	2,835	4,597
7	-4,0	4,080	5,556
8	-4,0	2,046	2,557
9	-4,0	1,982	2,592

p. *Tetradecenoic acid*

Mode	Affinity (kcal/mol)	RMSD l b	RMSD u b
1	-4,1	0,000	0,000
2	-4,1	2,926	2,926
3	-4,1	1,353	1,353
4	-4,0	1,717	1,717
5	-4,0	5,406	5,406
6	-4,0	2,645	2,645
7	-3,9	5,916	5,916
8	-3,9	5,157	5,157
9	-3,9	5,396	5,396

q. *1-Nonadecene*

Mode	Affinity (kcal/mol)	RMSD l b	RMSD u b
1	-3,8	0,000	0,000
2	-3,7	1,226	4,293
3	-3,7	1,609	5,701
4	-3,7	2,015	5,635
5	-3,7	2,231	4,076
6	-3,7	1,410	2,706
7	-3,7	2,212	5,827
8	-3,7	0,976	5,601
9	-3,6	1,596	4,891

r. *Benzenepropanoic acid*

Mode	Affinity (kcal/mol)	RMSD l b	RMSD u b
1	-4,9	0,000	0,000
2	-4,9	0,080	1,472
3	-4,8	3,065	4,494
4	-4,8	3,067	4,441
5	-4,8	0,917	1,749
6	-4,7	3,468	4,797
7	-4,7	3,497	4,836
8	-4,6	3,962	5,506
9	-4,6	1,396	2,087

*s. Cis-9-Hexadecenoic acid*

Mode	Affinity (kcal/mol)	RMSD l b	RMSD u b
1	-4,4	0,000	0,000
2	-4,4	1,153	1,789
3	-4,4	1,033	3,294
4	-4,3	1,094	3,713
5	-4,3	3,507	5,580
6	-4,3	3,532	5,980
7	-4,2	3,545	5,699
8	-4,2	1,545	2,809
9	-4,2	3,611	5,807

*t. Pentadecanoic acid*

Mode	Affinity (kcal/mol)	RMSD l b	RMSD u b
1	-4,2	0,000	0,000
2	-4,2	3,561	6,060
3	-4,1	3,613	5,560
4	-4,1	1,293	2,334
5	-4,0	1,528	3,749
6	-4,0	3,514	5,870
7	-4,0	3,569	6,048
8	-4,0	3,661	5,510
9	-4,0	4,102	6,648

*u. 9-Tricosene*

Mode	Affinity (kcal/mol)	RMSD l b	RMSD u b
1	-3,7	0,000	0,000
2	-3,7	0,912	2,733
3	-3,7	1,710	6,150
4	-3,7	1,330	3,980
5	-3,7	1,333	3,055
6	-3,7	2,070	6,067
7	-3,6	2,564	6,533
8	-3,6	1,733	3,922
9	-3,6	1,565	6,031

*v. Cis-9-Hexadecenal*

Mode	Affinity (kcal/mol)	RMSD l b	RMSD u b
1	-4,0	0,000	0,000
2	-4,0	1,491	2,192
3	-4,0	2,218	3,798
4	-3,9	2,511	5,615
5	-3,9	1,132	2,285
6	-3,9	2,853	5,838
7	-3,9	1,192	1,897
8	-3,9	1,306	4,192
9	-3,9	2,693	6,222

*w. Octadecanoic acid*

Mode	Affinity (kcal/mol)	RMSD l b	RMSD u b
1	-4,0	0,000	0,000
2	-3,9	3,440	6,083
3	-3,9	3,391	5,588
4	-3,9	3,401	5,963
5	-3,9	1,019	1,999
6	-3,8	1,981	3,509
7	-3,8	3,366	5,303
8	-3,8	1,409	2,470
9	-3,8	3,305	6,525

*x. 2,6,10,14,18,22 Tetracosahexaene*

Mode	Affinity (kcal/mol)	RMSD l b	RMSD u b
1	-4,4	0,000	0,000
2	-4,3	0,881	5,825
3	-4,3	2,165	5,347
4	-4,3	1,576	5,166
5	-4,2	0,825	5,636
6	-4,2	1,834	4,680
7	-4,2	2,020	6,618
8	-4,2	1,586	4,120
9	-4,2	0,424	5,590

## Lampiran 12. Jarak Residu Asam Amino Reseptor dengan Ligan

### a. Ikatan Hidrogen

No.	Ligan	Ikatan Hidrogen	Jarak Interaksi (Å)
1	Ligan Validasi	VAL <sup>357</sup>	3,34
		ASP <sup>356</sup>	3,53 ; 3,48
		LEU <sup>355</sup>	3,58
		TYR <sup>360</sup>	3,67
2	<i>Estragole</i>		
3	<i>Hexadecanoic acid ethyl ester</i>		
4	<i>Epicatechin</i>	LEU <sup>355</sup>	2,64
		TYR <sup>360</sup>	3,82
5	<i>Gallocatechin</i>	ASP <sup>356</sup>	2,36 ; 2,68
		VAL <sup>357</sup>	2,68
		TYR <sup>360</sup>	3,13
6	<i>p-coumaric acid methyl ester</i>	ASP <sup>356</sup>	2,96
		TYR <sup>360</sup>	2,05
7	<i>1,2-Benzenedicarboxylic acid mono (2-ethylhexylester)</i>	LEU <sup>355</sup>	2,18
		ARG <sup>608</sup>	3,29
		TYR <sup>360</sup>	3,22 ; 2,73
8	<i>Beta-tocopherol</i>	TYR <sup>360</sup>	2,54
9	<i>Vitamin E</i>	TYR <sup>360</sup>	2,37
10	<i>Ethanimidic acid, ethyl ester</i>	VAL <sup>357</sup>	2,7
		LEU <sup>355</sup>	3,48
11	<i>Acetic acid</i>	TYR <sup>360</sup>	2,87
12	<i>2-hydroxy-gammbutyrolactone</i>	LEU <sup>355</sup>	2,68
13	<i>4H-pyran-4-one, 3 hidroxy</i>	ARG <sup>608</sup>	3,39
		ASP <sup>356</sup>	2,92
		VAL <sup>357</sup>	2,41
14	<i>Benzoic acid</i>	TYR <sup>360</sup>	2,4
15	<i>Pyrogallol</i>	GLU <sup>196</sup>	2,35
		VAL <sup>357</sup>	2,45
		ASP <sup>356</sup>	2,1
16	<i>Tetradecenoic acid</i>	TYR <sup>360</sup>	2
17	<i>1-Nonadecene</i>		
18	<i>Benzenepropanoic acid</i>	VAL <sup>357</sup>	2,39
19	<i>Cis-9-Hexadecenoic acid</i>	ARG <sup>608</sup>	2,28
20	<i>Pentadecanoic acid</i>	ASP <sup>356</sup>	2,93
21	<i>9-Tricosene</i>		
22	<i>Cis-9-Hexadecenal</i>		
23	<i>Octadecanoic acid</i>	ASP <sup>356</sup>	1,97
24	<i>2,6,10,14,18,22 Tetracosahexaene</i>		

b. interaksi elektrostatik

No.	Ligan	Elektrostatik	Jarak Interaksi (Å)
1	Ligan Validasi		
2	<i>Estragole</i>		
3	<i>Hexadecanoic acid ethyl ester</i>		
4	<i>Epicatechin</i>		
5	<i>Gallocatechin</i>	ARG <sup>608</sup>	4,22
6	<i>p-coumaric acid methyl ester</i>		
7	<i>1,2-Benzenedicarboxylic acid mono (2-ethylhexylester)</i>		
8	<i>Beta-tocopherol</i>	ARG <sup>608</sup>	4,33
9	<i>Vitamin E</i>	ARG <sup>608</sup>	4,25
10	<i>Ethanimidic acid, ethyl ester</i>		
11	<i>Acetic acid</i>		
12	<i>2-hydroxy-gammbutyrolactone</i>		
13	<i>4H-pyran-4-one, 3 hidroxy</i>		
14	<i>Benzoic acid</i>		
15	<i>Pyrogallol</i>	ARG <sup>608</sup>	4,01
16	<i>Tetradecenoic acid</i>		
17	<i>1-Nonadecene</i>		
18	<i>Benzenepropanoic acid</i>		
19	<i>Cis-9-Hexadecenoic acid</i>		
20	<i>Pentadecanoic acid</i>		
21	<i>9-Tricosene</i>		
22	<i>Cis-9-Hexadecenal</i>		
23	<i>Octadecanoic acid</i>		
24	<i>2,6,10,14,18,22 Tetracosahexaene</i>		



### A. Ikatan Hidrofobik

No.	Ligan	Ikatan Hidrofobik	Jarak Interaksi (Å)
1	Ligan Validasi		
2	<i>Estragole</i>	HIS <sup>717</sup> ARG <sup>608</sup> , TYR <sup>360</sup>	5,19 4,29 3,98 ; 3,61
3	<i>Hexadecanoic acid ethyl ester</i>	ARG <sup>608</sup> HIS <sup>717</sup> TYR <sup>360</sup>	5,17 4,05 3,82 ; 3,73 ; 4,91
4	<i>Epicatechin</i>	VAL <sup>718</sup> HIS <sup>717</sup>	4,78 5,04 ; 5,05
5	<i>Galocatechin</i>	TYR <sup>360</sup>	4,08 ; 4,7
6	<i>p-coumaric acid methyl ester</i>	TYR <sup>360</sup>	3,82
7	<i>1,2-Benzenedicarboxylic acid mono (2-ethylhexylester)</i>	TYR <sup>360</sup>	4,17
8	<i>Beta-tocopherol</i>	LEU <sup>868</sup> HIS <sup>717</sup> ARG <sup>608</sup> TYR <sup>360</sup>	4,59 4,42 ; 4,66 ; 5,21 4,15 5,39 ; 3,84
9	<i>Vitamin E</i>	TYR <sup>360</sup> ARG <sup>608</sup> VAL <sup>718</sup> HIS <sup>717</sup>	3,8 ; 4,3 4,4 4,61 3,95 ; 5 ; 5,01
10	<i>Ethanimidic acid, ethyl ester</i>	TYR <sup>360</sup>	3,53 ; 4,1
11	<i>Acetic acid</i>	TYR <sup>360</sup>	3,89
12	<i>2-hydroxy-gammbutyrolactone</i>		
13	<i>4H-pyran-4-one, 3 hidroxy</i>		
14	<i>Benzoic acid</i>	TYR <sup>360</sup>	3,89
15	<i>Pyrogallol</i>		
16	<i>Tetradecenoic acid</i>	TYR <sup>360</sup> HIS <sup>717</sup>	4,86 3,92 ; 5,14
17	<i>1-Nonadecene</i>	TYR <sup>360</sup> HIS <sup>717</sup> ARG <sup>608</sup>	4,8 4,71 4,23
18	<i>Benzenepropanoic acid</i>	TYR <sup>360</sup>	3,97
19	<i>Cis-9-Hexadecenoic acid</i>	HIS <sup>717</sup> TYR <sup>360</sup>	5,04 3,71 ; 4,55
20	<i>Pentadecanoic acid</i>	HIS <sup>717</sup> TYR <sup>360</sup>	4,56 4,21 ; 4,94 ; 5,36
21	<i>9-Tricosene</i>	TYR <sup>360</sup> HIS <sup>717</sup>	3,94 ; 3,98 ; 4,86 ; 4,92 4,64

<b>22</b>	<i>Cis-9-Hexadecenal</i>	<b>TYR<sup>360</sup></b>	4,03 ; 4,54
<b>23</b>	<i>Octadecanoic acid</i>	<b>TYR<sup>360</sup></b> <b>HIS<sup>717</sup></b>	5,01 5,48 ; 5,35 ; 4,24
<b>24</b>	<i>2,6,10,14,18,22 Tetracosahexaene</i>	<b>ARG<sup>608</sup></b> <b>HIS<sup>717</sup></b> <b>TYR<sup>360</sup></b> <b>VAL<sup>718</sup></b> <b>LEU<sup>868</sup></b>	4,03 4,27 ; 4,6 ; 5,01 ; 5,08 4,14 ; 4,46 ; 5,43 5,06 4,49

**Lampiran 13.** Hasil Pemeriksaan *Similarity* Skripsi



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