Assignment -02

Task 01: Please conduct multiple Protein-Ligand Docking and provide the top 10 results in the following table.

	Ligands	Binding	Rmsd/ub	Rmsd/ib
		Affinity		
01	e4u9c_6436348_uff_E=402. 80	-6.2	0	0
02	e4u9c_10104370_uff_E=14 6.78	-5.5	0	0
03	e4u9c_196216_uff_E=215.0 6	-5.5	0	0
04	e4u9c_14529_uff_E=139.23	-5.5	0	0
05	e4u9c_79035_uff_E=1568.1 0	-5.4	0	0
06	e4u9c_5315469_uff_E=160. 58	-5.3	0	0
07	e4u9c_28930_uff_E=387.47	-5.2	0	0
08	e4u9c_7463_uff_E=91.91	-5	0	0
09	e4u9c_7461_uff_E=63.24	-5	0	0
10	e4u9c_31253_uff_E=90.99	-4.5	0	0

Task 02: Please perform ADME analysis on the 10 compounds obtained from the Protein-Ligand Docking and provide the results in the table below.

ADME Analysis						Pharmacokinetics		Drug likeness	Medicinal Chemistry		
Name	CID ID	Canonical SMILES	Molecular weight	Num . H-bo nd acce ptors	Num. H-bon d donors	Lipophilicity (iLOGP)	Water Solubility (Log S (SILICOS-I T))	GI absorption	BBB permeant	Lipinski	PAINS
beta- Bisab olen e		CC(=CCCC(=C)[C @H]1CCC(=CC1)C)C	204.35	0	0	3.67	-3.58	Low	No	1	0
2-(4- Meth ylphe nyl)p ropa n-2-o		Cc1ccc(cc1)C(O)(C)C	150.22	1	1	2.17	-3.01	High	Yes	0	0

Myrc	C=CC(=C)CCC=C(C)C	136.23	0	0	2.89	-2.42	Low	Yes	0	0
ene alpha -Fenc	Cc1ccc(cc1)C(O)(C)C		-		2.17			Yes	0	0
hene	6,6	150.22	1	1		-3.01	High			
gam ma-T erpin	CC1=CCC(=CC1)C (C)C				2.73			Yes	0	0
ene		136.23	0	0		-2.23	Low			
Bisac	CC(=CC(CC(c1ccc(2.87			Yes	0	0
umol	cc1)C)C)O)C	218.33	1	1		-3.97	High			
	CC(C1CCC(=C)C=				3.14			Yes	0	0
Curlo	C1)CC(=O)C=C(C)									
ne	C	218.33	1	0		-2.9	High			
р-Су	Cc1ccc(cc1)C(C)C				2.51			Yes	1	0
men e		134.22	0	0		-3.57	Low			
Ger	C/C/1=CCC(=C(C)				2.88			Yes	0	0
macr	C)C(=O)C/C(=C/C									
one	C1)/C	218.33	1	0		-3.64	High			
Tricy	CC12C3C1CC(C2(2.53			Yes	1	0
clene	C)C)C3	136.23	0	0		-2.23	Low			

Task 03: Perform Toxicity Prediction on the 10 compounds obtained from
the Protein-Ligand Docking and provide the results in the table below.

Toxicity Prediction Name CID ID Canonical SMILES Hepatotoxicity Carcinog **Immunotoxicity** Mutagenicity Cytotoxicity enicity beta-Bisab 1010437 CC(=CCCC(=C)[C@H]1C Inactive Active Inactive Inactive Inactive olene CC(=CC1)C)C 2-(4-Meth 14529 Cc1ccc(cc1)C(O)(C)C Inactive Active Inactive Inactive Inactive ylphenyl)p ropan-2-ol C=CC(=C)CCC=C(C)CInactive Inactive Inactive Inactive 31253 Inactive Myrcene alpha-Fen 28930 Cc1ccc(cc1)C(O)(C)C Inactive Active Inactive Inactive Inactive chene CC1=CCC(=CC1)C(C)C gamma-Te 7461 Inactive Inactive Inactive Inactive Inactive rpinene 5315469 CC(=CC(CC(c1ccc(cc1)C) Inactive Inactive Inactive Inactive Inactive Bisacumol C)O)C CC(C1CCC(=C)C=C1)CC(19216 Inactive Inactive Inactive Inactive Inactive Curlone =O)C=C(C)C 7463 Cc1ccc(cc1)C(C)C Inactive Inactive Inactive Inactive p-Cymene Inactive

Germacro	6436348	C/C/1=CCC(=C(C)C)C(=	Inactive	Inactive	Inactive	Inactive	Inactive
ne		O)C/C(=C/CC1)/C					
Tricyclene	79035	CC12C3C1CC(C2(C)C)C3	Inactive	Inactive	Inactive	Inactive	Inactive

Task 04: Identify the highest-ranking Protein – ligand complex and input the corresponding figures into the table below.

<u> </u>	onding figures in						
Figure Name	Sample Fig	Sample Figure		Input your Docking Figure			
Figure 01				A97			
Figure 02		·-		TO SEE TO			
Task 05: Identify	the highest-ranki	ng Prot	ein – liga		and input		
the Interaction d	etails into the tab	le belo	W.	_ (C) (C)			
Name		'	ategory		· !S		
N:UNK1:C FAgALA@103	C. C.	-	lydrophobic				
N:UNK1:C - A:ALA97	Jan Lung	H	lydrophobic	76117			
N:UNK1:C - A:VAL133	5.12820	H	lydrophobic	N. T.			
N:UNK1:C - A:LYS117	4.70026	H	lydrophobic				
N:UNK1:C - A:LYS117	4.5107	H	lydrophobic				
A:LYS117 - N:UNK1	<u>4 51099</u>	 	lydrophobic	 + ,	Alkyl		
N:UNK1:C - A:ALA97		H	lydrophobic	_ //>	Alkyl		
N:UNK1:C Figure 04		H	lydrophobic	- Gas	Alkyl		
N:UNK1:C - A:VAL133		H	lydrophobic		Alkyl		
N:UNK1:C - A:LYS117	4.70026	H	lydrophobic		Alkyl		
N:UNK1:C - A:LYS117	4.5107	Н	<u>lydrophobic</u>	`J	Alkyl		
A:LYS117 - N:UNK1	4.51099	I	lydrophobic	A:1954 A:19 7 9	:yl		
N:UNK1:C FACALA9705	3.7	 	lydrophobic	A:97	χl		
N:UNK1:C - A:ALA97	3.68		lydrophobic	Ā.Hrs	Ŋı		
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