



INDIAN INSTITUTE OF TECHNOLOGY GUWAHATI

END SEMESTER EXAMINATION – BT 305: Computational Biology

April 29, 2024. Maximum Marks: 70 Time: 3 hours

Section A: 5 marks for each question

1. The SMILES representation of four molecules are as follows. Attempt to draw an approximate 2D structure of these molecules.

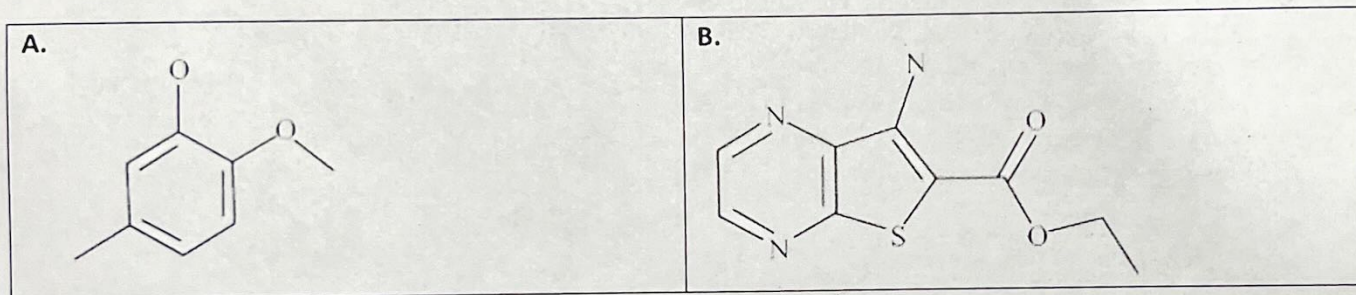
CCO

CC(=O)O

C1CCCCC1

c1cnccc1

2. Write the 1D string representation (SMILES) for the following molecules.



3. According to Hansch equation, how is the activity of a drug related to its hydrophobicity. How is the biological activity of a molecule related to its electronic characteristics and hydrophobicity?
4. Explain briefly the following concepts.
☒ A) Simulated Annealing ☒ B) Energy minimization ☒ C) RMSD based clustering ☒ D) Solvent accessible surface area.
5. ☒ A) Attempt to write an additive equation for free energy of binding in a molecular docking experiment.
☒ B) Free energy of binding can also be expressed based on simple molecular mechanics-based scoring functions. What are the most likely variables (or components) in such free energy estimations.
6. Explain briefly the following concepts.
☒ A) Pharmacophore ☒ B) Receptor ☒ C) Lipinski rule of five ☒ D) Molecular descriptor.

Section B: 20 marks for each question

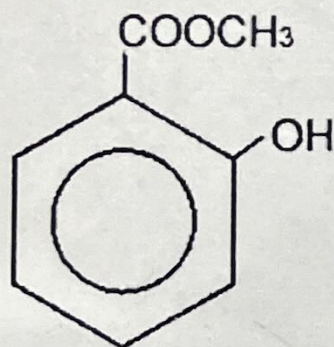
7. ☒ Bit strings for query and test molecules in a typical fingerprinting exercise is given below.

Mol 1	0	1	1	0	0	1	1	1	0
Mol 2	1	1	0	1	1	0	0	0	0
Mol 3	1	1	0	1	1	1	0	1	1
Mol 4	1	0	1	1	0	1	1	0	0

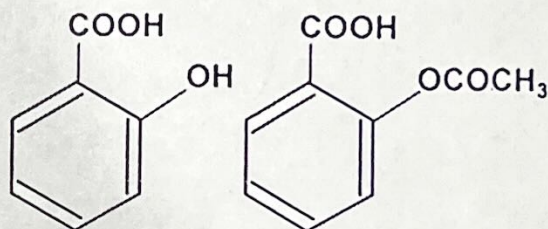
Calculate Tanimoto coefficients for structure comparison and attempt the following question:

If receptors of the four candidate structures Mol 1 Mol 2 Mol 3 and Mol 4 are respectively Rec 1, Rec 2, Rec 3 and Rec 4. Which molecule is more likely to bind with maximum number of Receptors? Rationalize your answer?

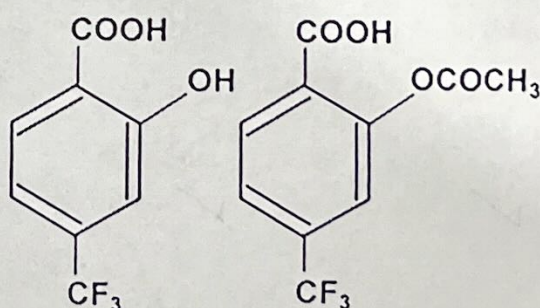
- ☒ Suppose we are performing an experiment of structure based molecular profiling for methyl salicylate (below structure) in a structure pool of four members on the right hand side, how do we determine the closest structure by calculating Tanimoto coefficient?



Salicylic acid Aspirin



HTB Triflusal



- ☒ 8. Using all three genetic operators (Selection, Crossover and Mutation), maximize the function $f(x) = x^3$, with 'x' lies in the interval 0 to 30. Generate initial population as 6, 9, 17 and 23.

End of Questions