

# COMPUTATIONAL STRUCTURE-BASED SCREENING AND EXPLICIT MOLECULAR DYNAMICS

3 & 4 AUG  
2021

Two-days cloud-based hands-on workshop targeting structure-based drug designing. Participants will get practical experience and in-person guidance in using the Maestro GUI, covering the organic molecule sketching, protein selection, preparation, and screening for hit identification of molecules against therapeutic targets. The workshop will also include a brief recap of background theory for Molecular mechanics, Molecular Docking, and Molecular Dynamics via case studies on the real-time industrial projects.

## DAY 1

- 9.30**  
Technical set-up
- 9.50**  
Audio & Visual check
- 10.00**  
1. Opening -  
Agenda and split into breakout rooms
- 10.10**  
Breakout 1 -  
Introductions and open software
- 10.30**  
2. Maestro GUI:  
Building Molecules and Enumeration
- 11.00**  
Breakout 2 -  
Build Chalcones from SMILES
- 11.20**  
3. Ligand Preparation and ADME
- 11.35**  
Breakout 3 -  
Launch LigPrep
- 11.45**  
Intro to Molecular Docking, Protein  
and Ligand Preparations
- 12.10**  
4. Protein Preparation and  
Grid Generations
- 12.30**  
Breakout 4 -  
Prepare protein and grid  
generations  
and molecular docking
- 13.00**  
Break -  
Continue to use software  
during the break
- 14.00**  
Welcome Back
- 14.10**  
5. Molecular Docking
- 14.25**  
Breakout 5 -  
Analyzing 3D molecular  
docking analysis
- 14.40**  
6. Molecular Docking Analysis
- 14.55**  
Breakout 6 -  
2D molecular docking analysis
- 15.30**  
7. Full Revision of Day 1  
Breakout 7 -  
Full session practice
- 15.55**  
Wrap-up
- 16.00**  
Finish

## COMPUTATIONAL STRUCTURE-BASED SCREENING AND EXPLICIT MOLECULAR DYNAMICS

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### DAY 2

**9.30**

Technical set-up

**9.50**

Audio & Visual check

**10.00**

1. Opening -  
Agenda and molecular dynamics theory

**10.30**

Breakout 1 -  
Docking results and ligand sketching

**10.45**

2. Desmond Introduction

**11.00**

Breakout 2 -  
Protein ligand complex formation,  
system building

**11.15**

3. Molecular Dynamics Demo

**11.25**

Breakout 3 -  
MD submission

**11.35**

Desmond Trajectory Visualization

**11.50**

Breakout 4 -  
Participant analysis and  
molecular docking

**13.00**

Break -  
Continue to use software  
during the break

**14.00**

Welcome Back

**14.05**

5. Desmond Molecular Simulation  
Analysis

**14.20**

Breakout 5 -  
Analyzing 3D molecular  
docking analysis

**14.45**

6. Organic Molecules Enumeration

**15.00**

Breakout 6 -  
Reaction-based enumeration,  
library enumeration, ADME

**15.30**

7. Full Revision of Day 2

**15.55**

Wrap-up

**16.00**

Finish

#### Speakers:

Dr. Pritesh Bhat  
Dr. Prajwal Nandekar  
Dr. Koushik Kasavajhala  
Ms. Shelvia Malik  
Mr. Kishore V  
Mr. Vinod D  
Mr. Sivakumar K C



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#### Bioinformatics Facility

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[More Info](#)

[Register](#)

**FEES**

**500 INR** for Students and Academicians  
**2500 INR** for Non-Academics  
**100 \$** for Foreign Participants