

# PROGRAMME SCHEDULE

## Structural Bioinformatics & Computer Aided Drug Designing (CADD)



### DAY - 1 (15/11/2024): Introduction to Structural Bioinformatics

08:30 - 9:00 am	<b>REGISTRATION</b>
	<i>Session-I: Protein Modeling and Structure Prediction</i>
09:00 - 10:00 am	<ul style="list-style-type: none"><li>✓ Sequence Structure Gap</li><li>✓ Protein Modeling: Homology Modeling and Threading</li><li>✓ Secondary structure prediction</li></ul>
10:00 - 10:30 am	<b>INAUGURATION</b>
10:30 - 11:00 am	<b>HIGH TEA</b>
	<i>Session-II: Protein Modeling and Structure Prediction</i>
11:00 - 12:15 pm	<ul style="list-style-type: none"><li>✓ Extended modelling approach using Python*</li><li>✓ Tertiary structure prediction through Ramachandran plot</li><li>✓ Visualisation of 3D protein structure*</li></ul>
12:15 - 12:30 pm	<b>Doubt Clearing Sessions</b>
12:30 - 01:45 pm	<b>LUNCH</b>
	<i>Session-III: Mutational Analysis and Thermodynamics</i>
01:45 - 03:00 pm	<ul style="list-style-type: none"><li>✓ Generation of mutants &amp; Thermodynamics*</li><li>✓ Energy minimisation of protein structures*</li><li>✓ Relative RMSD calculation*</li></ul>
03:00 - 03:30 pm	<b>HIGH TEA</b>
03:00 - 03:30 pm	<b>Doubt Clearing Sessions</b>
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### DAY - 2 (16/11/2024): Computer Aided Drug Designing

	<i>Session-I: Introduction to small compound databases</i>
09:00 - 10:30 am	<ul style="list-style-type: none"><li>✓ Data retrieval from PubChem and DrugBank</li><li>✓ Drug Designing using Avogadro/GaussView*</li><li>✓ Pharmacokinetics (ADMET) Analysis</li></ul>
10:30 - 11:00 am	<b>HIGH TEA</b>
	<i>Session-II: Molecular Docking and Analysis</i>
11:00 - 12:15 pm	<ul style="list-style-type: none"><li>✓ Introduction to Molecular Docking</li><li>✓ Molecular Docking using AutoDock*</li><li>✓ Intermolecular Interaction Analysis of Docked Poses</li></ul>
12:15 - 12:30 pm	<b>Doubt Clearing Sessions</b>
12:30 - 01:15 pm	<b>LUNCH</b>
	<i>Session-III: Molecular Dynamics Simulation</i>
01:15 - 03:00 pm	<ul style="list-style-type: none"><li>✓ Introduction to Linux OS</li><li>✓ Molecular Dynamics Simulation of Protein in GROMACS</li><li>✓ MD Visualization in VMD</li></ul>
03:00 - 03:15 pm	<b>HIGH TEA</b>
03:00 - 03:15 pm	<b>Doubt Clearing Sessions</b>
03:15 - 03:30 pm	<b>Valedictory Session and Vote of Thanks</b>
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\*Standalone softwares will be provided

## Two Days Workshop & Hands-on-Training

### Structural Bioinformatics & Computer Aided Drug Designing (CADD)

#### HIGHLIGHTS

- Introduction to Linux OS
- Computational Analysis using Python & ML
- Molecular Dynamics Simulation in GROMACS

#### Prerequisites

Personal Laptop (Windows / Linux OS)

Hurry !!  
Limited Seats



#### RESOURCE PERSON

Dr. Aniket Naha (M.Sc., Ph.D.)

Scientist & Student Coordinator  
Medical Biotechnology & Computational Drug Designing Laboratory,  
Pushpagiri Research Centre, PMS&RC, Tiruvalla

#### Registration Fees:

₹1500 (Ph.D. Scholars / Faculty)

₹1000 (Undergraduate / Postgraduate)

#### PAYMENT QR



Account Number: 024804176659190008

IFSC Code: CSBK0000248

MICR code: 689047003

Registration Link: <https://forms.gle/8sHfkn7STUzWqLUJ9>

15-16 November 2024

09:00 AM - 03:00 PM

Conference Hall, PRC

+91 - 8473912790

<http://prc.pushpagiri.in>

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