

Sr. No	Lecture Title	Description	Category	Durations
<b>Segment 6: Molecular Dynamics &amp; Simulation</b>				
1	Getting Started With Molecular Dynamics Simulation - Pre-processing of Protein Structure and Removal of Unnecessary Structural Features	<ul style="list-style-type: none"> <li>• Introduction to GROMACS and its installation.</li> <li>• Pre-processing of the protein structure and removal of unnecessary structural features.</li> </ul>	Protein Simulation	12:33
2	pdb2gmx - Construction of Topology File for Simulation	<ul style="list-style-type: none"> <li>• Introduction to pdb2gmx tool of GROMACS.</li> <li>• Construction of topology file for simulation.</li> </ul>	Protein Simulation	9:00
3	Defining a Solvant Box for Simulation	<ul style="list-style-type: none"> <li>• Introduction to GROMACS for defining solvant box.</li> <li>• Defining a solvant box to prepare the protein for simulation.</li> </ul>	Protein Simulation	4:14
4	Solvation - Adding Water Molecules in Solvant Box	<ul style="list-style-type: none"> <li>• Introduction to GROMACS for defining solvant box.</li> <li>• Adding water molecules in solvant box to prepare the protein for simulation.</li> </ul>	Protein Simulation	5:30
5	Generating Input Run File Replacement of Water Molecules With Ions	<ul style="list-style-type: none"> <li>• Generating input run file.</li> <li>• Replacing the water molecules with ions for protein simulation.</li> </ul>	Protein Simulation	6:55
6	genion - Replacement of Water Molecules With Ions	<ul style="list-style-type: none"> <li>• Introduction to genion tool of GROMACS.</li> <li>• Replacing the water molecules with ions for protein simulation.</li> </ul>	Protein Simulation	4:18
7	Energy Minimization - Relaxing and Fixing the Structure for Simulation	<ul style="list-style-type: none"> <li>• Introduction to gmx energy tool of GROMACS.</li> <li>• Relaxing and fixing the structure for protein simulation.</li> </ul>	Protein Simulation	11:25
8	GRACE - Visualization and Analysis of Minimized Structure	<ul style="list-style-type: none"> <li>• Introduction to gmgrace tool of GROMACS.</li> <li>• Visualization and analysis of minimized structure of the protein.</li> </ul>	Protein Simulation	4:11

9	Equilibration of Protein Structure NVT ENSEMBLE Phase 1	<ul style="list-style-type: none"> <li>• Introduction to nvt tool of GROMACS.</li> <li>• Phase 1 of NVT ENSEMBLE to equilibrate the protein structure.</li> </ul>	Protein Simulation	8:37
10	Equilibration of Protein Structure NPT ENSEMBLE Phase 2	<ul style="list-style-type: none"> <li>• Introduction to nvt tool of GROMACS.</li> <li>• Phase 2 of NVT ENSEMBLE to equilibrate the protein structure.</li> </ul>	Protein Simulation	8:09
11	mdrun - Executing Simulation Analysis	<ul style="list-style-type: none"> <li>• Introduction to mdrun tool of GROMACS.</li> <li>• Execution of MD simulation on the protein structure.</li> </ul>	Protein Simulation	3:46