| Days | Name | Durations | Category | Main Category |
|------|--|-----------|-----------------------------------|--|
| 1 | Getting Started With Molecular Dyanmics Simulation - Pre-processing of Protein Structure and Removal of Unncessary Structural Features | 12:33 | Protein Structure Preparation | Molecular Dynamics Simulations: GROMACS |
| 2 | pdb2gmx - Construction of Topology File for Simulation | 9:00 | Protein Structure Preparation | Molecular Dynamics Simulations: GROMACS |
| 3 | Defining a Solvant Box for Simulation | 4:14 | Protein Structure Preparation | Molecular Dynamics Simulations: GROMACS |
| 4 | Solvation - Adding Water Molecules in Solvant Box | 5:30 | Protein Structure Preparation | Molecular Dynamics Simulations: GROMACS |
| 5 | Generating Input Run File Replacement of Water Molecues With Ions | 6:55 | Protein Structure Preparation | Molecular Dynamics Simulations: GROMACS |
| 6 | genion - Replacement of Water Molecules With Ions | 4:18 | Protein Structure Preparation | Molecular Dynamics Simulations: GROMACS |
| 7 | Energy Minimization - Relaxing and Fixing the Structure for Simulation | 11:25 | Energy Minimization | Molecular Dynamics Simulations: GROMACS |
| 8 | GRACE - Visualization and Analysis of Minimized Structure | 4:11 | Minimized Structure Visualization | Molecular Dynamics Simulations: GROMACS |
| 9 | Equibiliration of Protein Structure NVT ENSEMBLE Phase 1 | 8:37 | NVT Ensemble | Molecular Dynamics Simulations: GROMACS |
| 10 | Equibiliration of Protein Structure NPT ENSEMBLE Phase 2 | 8:09 | NPT Ensemble | Molecular Dynamics Simulations: GROMACS |
| 11 | mdrun - Executing Simulation Analysis | 3:46 | Simulation Execution | Molecular Dynamics Simulations: GROMACS |