

Sr. No	Lecture Title	Description	Category	Duration
Segment 3: 3D Structure Prediction				
1	PEPFOLD 3 Peptide Structure Modeling	<ul style="list-style-type: none"> • Introduction to PEPFOLD_3 server and its purpose. • Generating 3D models of a peptide using the query sequence. • Defines parameters and analysis of results. 	Peptide Structure prediction	13:14
2	MODELLER	<ul style="list-style-type: none"> • Introduction to Modeller tool and its uses. • Procedure to predict a protein structure through Modeller. • Evaluation method of MODELLER to find out the most optimal and good protein structure predicted. 	3D Structure Prediction	36:13:00
3	SwissModel	<ul style="list-style-type: none"> • Introduction to homology modeling and SwissModel, a homology modeling server. • Prediction of protein structure for a target sequence using SwissModel. • Defines parameters for a good protein structure and analysis of resulting protein. 	3D Structure Prediction	12:52
4	HHPRED	<ul style="list-style-type: none"> • Basic description of HHPred tool and its purpose. • Procedure to predict the protein structure from target sequence through HHPred tool. • Selecting a particular template structure for homology modeling of target protein using HHPred tool. 	3D Structure Prediction	14:09
5	M4T	<ul style="list-style-type: none"> • Introduction to M4T, a protein structure prediction tool. • Procedure to predict the protein structure from a Target protein sequence, using the M4T server. 	3D Structure Prediction	9:26

6	IntFOLD	<ul style="list-style-type: none"> • Introduction to IntFOLD and its purpose. • Procedure to predict the protein structure from target protein sequence, using the IntFOLD server. • Interpretation of results. 	3D Structure Prediction	8:41
7	ROBETTA	<ul style="list-style-type: none"> • Introduction to Robetta, a protein structure prediction tool. • Procedure to predict and analyse protein structure. • Prediction of protein structure for a query sequence using ab-initio techniques. 	3D Structure Prediction	14:39
8	Homology Modeling Using MOE	<ul style="list-style-type: none"> • Introduction to homology modeling and Molecular Operating Environment (MOE) tool. • Generating a homology 3D model of a target protein using MOE. • Analysis of results and how to align the two structures of the protein using the MOE tool. 	3D Structure Prediction	12:34