

STRUCTURAL

BIOINFORMATICS IN PRACTICE

17.-18. November 2016

Universitetet på Færøerne

Day 1	Biomolecular visualisation and structure prediction	
09:15 - 10:00	Introduction to Structural Bioinformatics	
10:15 - 10:30	Hands-on: The Protein Data Bank	
10:30 - 12:00	Hands-on: Biomolecular visualisation with PyMOL	
12:00 - 12:30	Lunch	
12:30 - 13:00	Biomolecular visualisation continued	
13:15 - 15:00	Hands-on: <ul style="list-style-type: none">• Homology modelling with Swiss-Model• Secondary structure element prediction with PSI-Pred	

Day 2	Protein flexibility	
09:15-11:00	Hands-on: Comparative structural analysis with Bio3D-web	
11:15 - 12:00	Hands-on: Beginning protein structure analysis with Bio3D	
12:00 - 12:30	Lunch	
12:30 - 14:00	Structure analysis with Bio3D continued	
14:15 - 14:45	Other topics, e.g. protein-ligand docking, minimization and molecular dynamics simulations, other databases	
14:45 - 15:00	Course evaluation	

PyMOL

 **Bio3D**

 **R Studio**

