

Structural Variant Effects, Stability & Function — Hands-on

Understand how amino acid substitutions, indels and domain rearrangements propagate from sequence to structure, stability and function. This module walks through mapping variants onto 3D models, using stability and impact predictors, combining docking and MD evidence, and communicating structural mechanisms of disease and protein engineering decisions.

Structural Variant Effects, Stability & Function

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Session Index

Session 1 — Structural Variants, Mapping & Annotation Session 2 — Stability, Packing &

Aggregation Risk Session 3 — Mechanisms, Dynamics & Functional Impact Session 4 — Interpretation, Reporting & Case Studies

Session 1

Fee: Rs 8800 Apply Now

Structural Variants, Mapping & Annotation

Classes of protein structural variants

missense, indels, truncations domain swaps and

fusions germline vs somatic contexts

Mapping variants from sequence to structure

reference sequences and numbering locating

residues on PDB or models isoforms, chains and

alignment issues

Basic annotation of variant context

secondary structure and environment buried vs

exposed positions proximity to sites and interfaces

Session 2

Fee: Rs 11800 Apply Now

Stability, Packing & Aggregation Risk

Stability and packing around variant sites

core vs surface substitutions hydrogen bonds and salt bridges cavities, clashes and packing defects

Computational stability and ddG style scores

concept of delta delta G of folding interpreting stabilising vs destabilising calls limits and calibration of predictors

Aggregation and misfolding risk indicators

exposed hydrophobics and patches disruption of disulfides and motifs relevance for expression and developability

Session 3

Fee: Rs 14800 Apply Now

Mechanisms, Dynamics & Functional Impact

Mechanistic classes of functional impact

active site and catalytic variants binding site and interface disruption allosteric, regulatory and conformational shifts

Integrating MD, docking and network views

local flexibility and RMSF changes contact maps and communication paths effect on ligand and partner

binding modes

Scoring and prioritising variants for follow up

combining structural and sequence evidence flagging likely loss vs gain of function shortlisting variants for experiments

Session 4
Fee: Rs 18800 Apply Now

Interpretation, Reporting & Case Studies

End to end structural variant case study

Theory + Practical

Structurally informed variant interpretation narratives

clear mechanism of impact statements figures and maps for reports and slides strength, caveats and alternatives

Linking to disease, diagnostics and design decisions

triaging clinical variants of interest guiding protein engineering hypotheses structural variant summaries for teams