

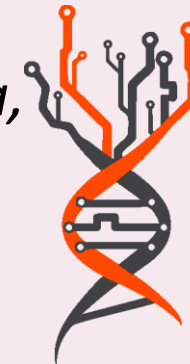
Exploration of Intrinsic Disorder Regions through Classification of Intrinsically Disordered Proteins Using PPI Network Structure and Sequence Attributes: A Case Study

Milana Grbić, Milan Predojević, Nenad Vilendečić, Dragan Matić

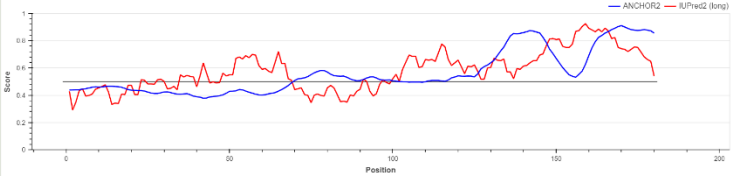
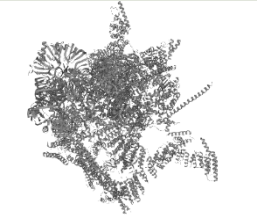
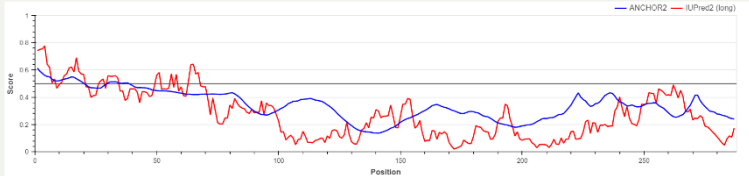
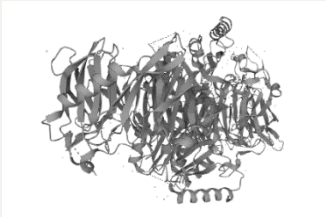
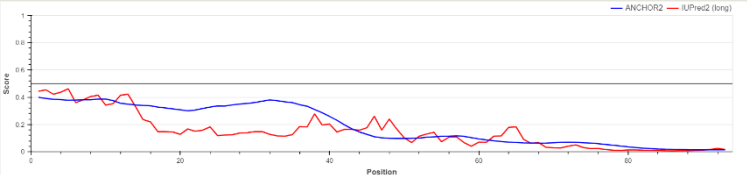
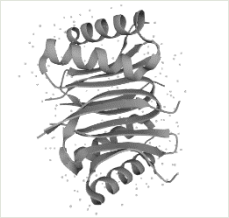
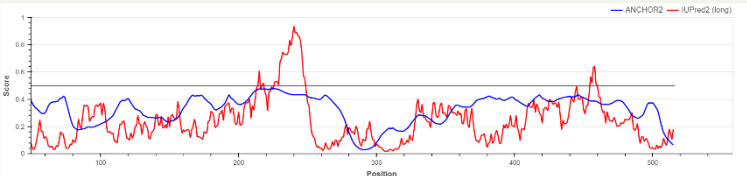
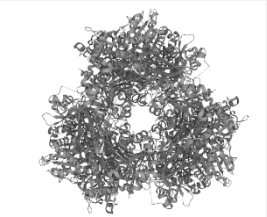
Faculty of Natural Science and Mathematics, University of Banja Luka

Belgrade Bioinformatics Conference 2024, Belgrade, Serbia,

June 17-21, 2024



**Belgrade
Bioinformatics
Conference**

Protein	IUPred2A	Function	Structure	Subcellular Location
YNL245C		Involved in pre-mRNA splicing.		Nucleus
YDL201W		Methyltransferase that catalyzes the		Nucleus
YDR424C		non-catalytic accessory components of the cytoplasmic dynein complex		Cytoplasm, cytoskeleton Nucleus, nuclear pore complex
YKL103C		Resident vacuolar enzyme that catalyzes the removal of amino acids from the N-terminus of peptides and proteins		Vacuole