

Bioinformatics project

Submitted to: Dr. Samina Shakeel

Submitted by: Sayeda Zahra Batool

Department: Biochemistry Bs 7th Morning

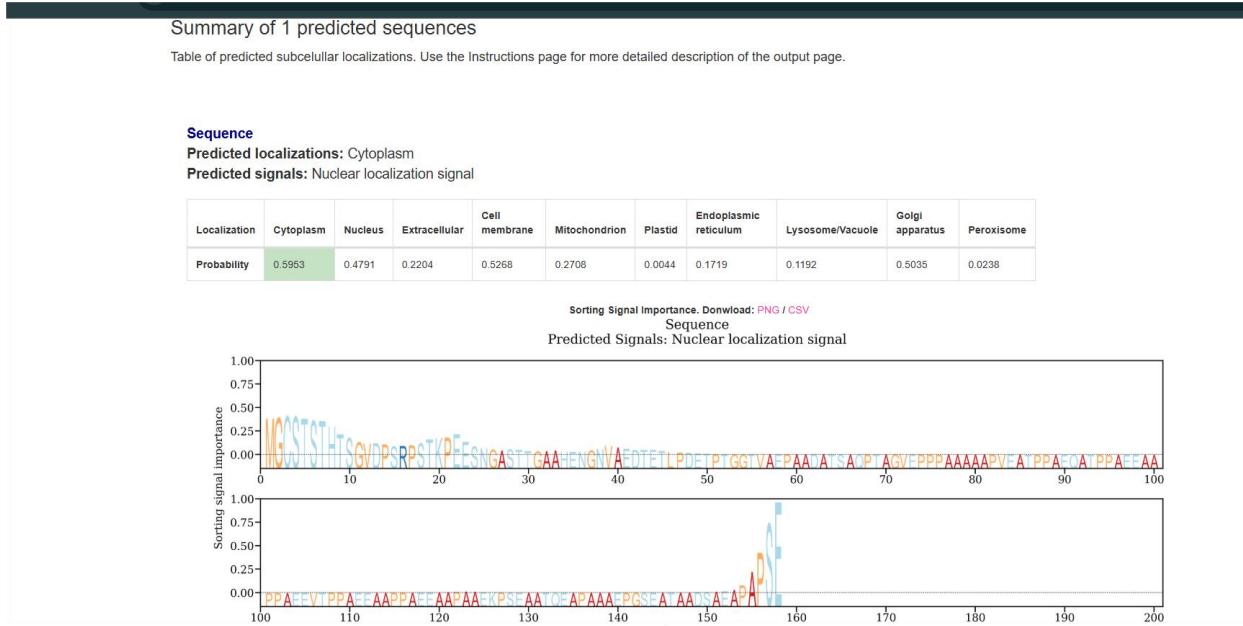
Task No: 4

Subcellular Localization Prediction:

1. PSORT:

The screenshot shows the PSORT web interface. At the top, there are tabs for 'Overview' and 'Quote/Order'. Below the tabs, it says 'CUE: Lupas's algorithm to detect coiled-coil regions' and 'total: 0 residues'. A blue header bar reads 'Results of the k-NN Prediction'. Underneath, it shows 'k = 9/23' and a list of localization probabilities: '73.9 %: nuclear', '13.0 %: cytoplasmic', '4.3 %: cytoskeletal', '4.3 %: peroxisomal', and '4.3 %: mitochondrial'. At the bottom, it says '>> prediction for 176670258622688 is nuc (k=23)'.

2. DeepLoc:



Subcellular Localization Justification:

Results Summary:

Tool 1 (PSORT):

Predicted Localization in Nucleus (Score: 72%).

Tool 2 (DeepLoc):

Predicted Localization in Cytoplasm (Probability: 0.5953) and Nucleus (Probability: 0.4791).

Scientific Justification:

The subcellular localization analysis of Tropomyosin 1 shows a dual distribution between the Nucleus and Cytoplasm. While PSORT strongly predicts a nuclear localization, DeepLoc indicates a primary presence in the cytoplasm with a significant secondary probability for the nucleus. This is scientifically consistent with the known behavior of Tropomyosin 1 isoforms, which are primarily structural components of the cytoskeleton but are also known to shuttle into the nucleus to participate in processes like RNA processing and chromatin remodeling. Therefore, the protein likely localizes in both compartments depending on the cellular requirements.