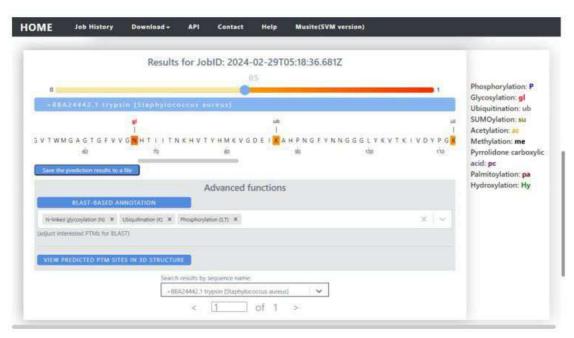
Experiment Result... Page No. 40 Result and Discussions Result and Discussion:

The output is visualized for each input sequence one by one which can be sufriened by its provided by its provided writer and the predicted top of sofresponding position multiple labules are shaw on top of prediction of that position is predicted to have multiple provided to their confidence level.

Protein Accession Identifier (ACCID) BLAST sequence identify and the known PTM annotation an homologous protein an presented for each input sequence. integrating 925 a tool to annotate genomic variouse on perotein structure & New Viewer a mel application for molecular vigualization. Study of post translational modification lesing relevant tools.

Alon: B study post translational modification using online tool musiledep.





trypsin [Staphylococcus aureus]

GenBank: BBA24442.1

GenPept Identical Proteins Graphics

>BBA24442.1 trypsin [Staphylococcus aureus]

MNKNIIIKSIAALTILTSVTGVGTTMVEGVQQTAKAENSVKQITNTNVAPYSGVTWMGAGTGFVVGNHTI ITNKHVTYHMKVGDEIKAHPNGFYNNGGGLYKVTKIVDYPGKEDIAVVQVEEKSTQPKGRKFKDFTSKFN IASETKENEPISVIGYPNPNGNKLQMYESTGKVLSVNGNIVSSDAIIQPGSSGSPILNSKYEAIGVIYAG NKPSGESTRAFAVYFSPEIKKFIADNLDK

