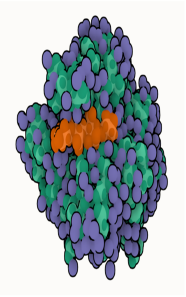


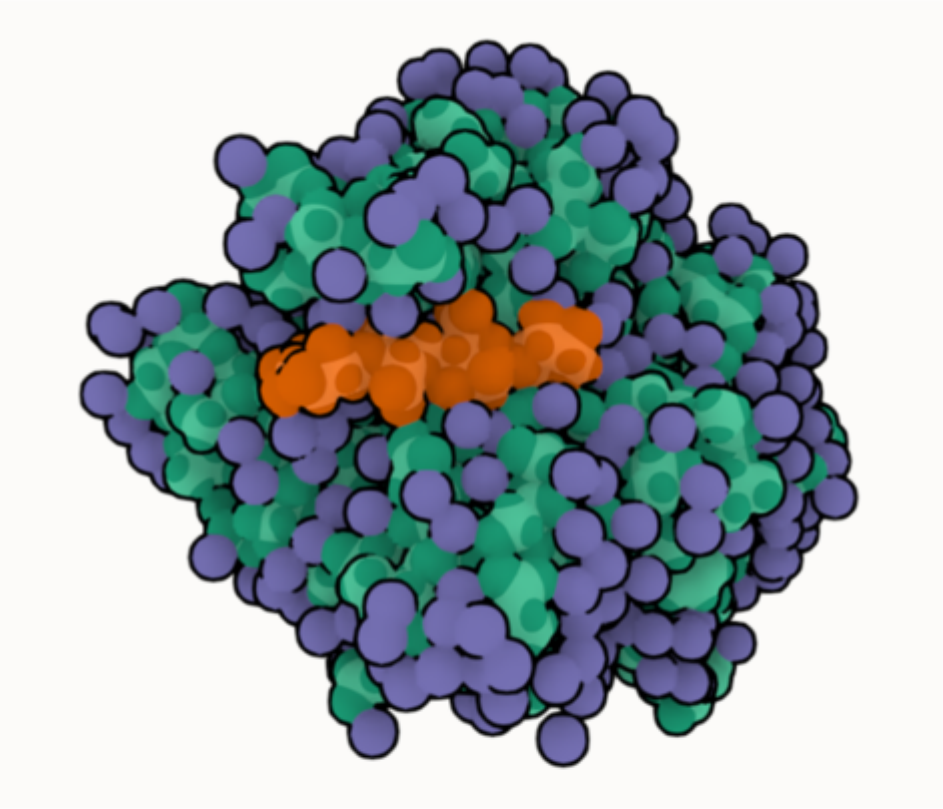
8QLI Crystal structure of Paradendryphiella salina PL7C alginate lyase mutant H124 soaked with with tetra-mannuronic acid

8ZDW The cryoEM structure of H5N1 HA split from symmetric filament in conformation

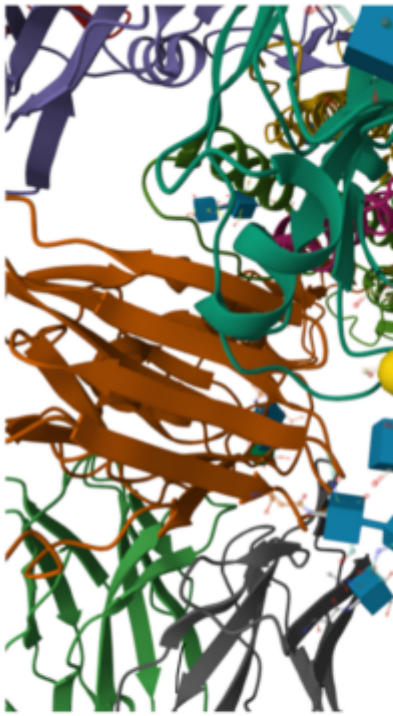
Protein-Carbohydrate Interfaces. PDB depositions: September_2024

Description

DIONYSUS is a database of protein-carbohydrate interfaces annotated according to proteins and carbohydratesâ?? structural, chemical and functional properties . https://www.dsimb.inserm.fr/DIONYSUS. It collects and analyzes protein-carbohydrate complexes of various types, including non-covalent ligand binding, glycosylated proteins, glycopeptides, and reaction intermediates in sugar enzymes. The update of DIONYSUS offers the possibility of creating a regular list of the latest structures added to the PDB, including PDB ID, title, protein function, and type of carbohydrate. Listed below are the latest structures added in **September 2024**



8QLI Crystal structure of Paradendryphiella salina PL7C alginate lyase mutant H124 soaked with with tetra-mannuronic acid



8ZDW The cryoEM s split from symmetri

Copie de September_PDB_Protein_CHO_Starting fileTÃ©lÃ©charger

Category

1. News