



Paradendrynhiella salina PI 7C alginate lyase mutant H124 soaked with with tetra-mannuronic acid

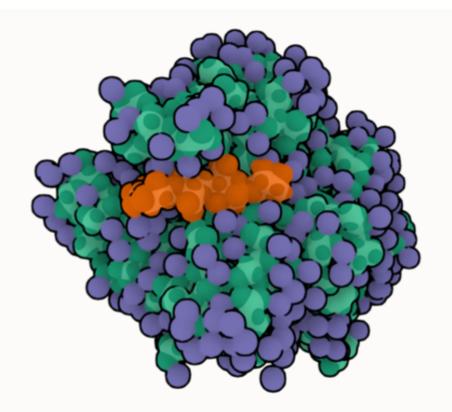
8ZDW The cryoEM structure of H5N1 HA split from symmetric filament in conformation

## 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 fileTéIécharger

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