List of descriptors and labels for carbohydrate binding site clustering

General information on protein and carbohydrate:

pdb_id Model

Carbohydrate_name

Carbohydrate_label

Carbohydrate type: 1 = free ligand; 2 = glycosylation head; 3 = glycosylation body

Different properties of the binding site:

Polar_sasa

Apolar_sasa

Total_sasa

Residues_in_helices

Residues in strands

Residues_in_turns

Residues_in_bends

Residues_in_loops

Different functional groups present in ligands:

Function lipid

Function_acid

Function amide

Function_charged

Function_aromatic

Function_selenium

Function_halogen

Function_phosphorus

function_sulfurfunction_amino_acid

Carbohydrate binding site composition:

| count_A | count_C | count_D | count_E | count_F | count_G |
|---------|---------|---------|---------|---------|---------|
| count_H | count_I | count_K | count_L | count_M | count_N |
| count_P | count_Q | count_R | count_S | count_T | count_V |
| count_W | count_Y | | | | |