
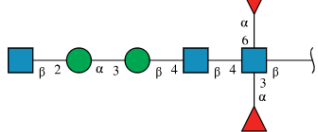

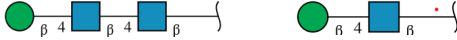

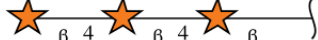

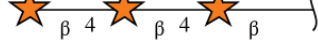
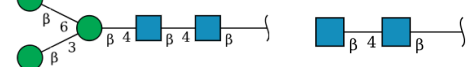

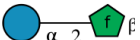
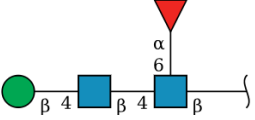
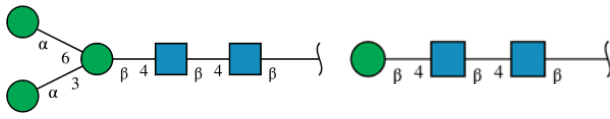
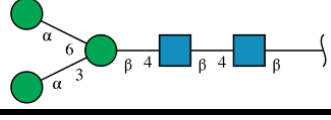
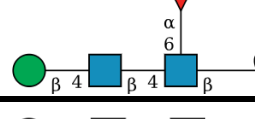


PDB ID	Title	Keywords	
<a href="#">8TFT</a>	Fab of O13-1 human IgG1 antibody bound to IgV domain of human TIM-3	IMMUNE SYSTEM	<b>NAG</b>
<a href="#">8XPK</a>	Marine bacterial laminarinase PtLam mutant E154A in complex with laminatriose	HYDROLASE	
<a href="#">8RHG</a>	CryoEM structure of human rho1 GABAA receptor in complex with GABA and pregnenolone sulfate	MEMBRANE PROTEIN	<b>NAG</b>
<a href="#">8YQQ</a>	Structure of HKU1B RBD with Tmprss2	VIRAL PROTEIN/HYDROLASE	<b>NAG</b>
<a href="#">8RH9</a>	CryoEM structure of human rho1 GABAA receptor in complex with pregnenolone sulfate	MEMBRANE PROTEIN	<b>NAG</b>
<a href="#">8RH4</a>	CryoEM structure of human rho1 GABAA receptor in complex with estradiol	MEMBRANE PROTEIN	<b>NAG</b>
<a href="#">8R7I</a>	X-ray structure of blue laccase BP76 from the termite Neocapritermes taracua	OXIDOREDUCTASE	
<a href="#">8RH7</a>	CryoEM structure of human rho1 GABAA receptor in complex with GABA and estradiol in primed state	MEMBRANE PROTEIN	<b>NAG</b>
<a href="#">8TGU</a>	Cryo-EM structure of BG505 SOSIP trimer purified via Galanthus nivalis lectin chromatography	VIRAL PROTEIN	
<a href="#">8TGW</a>	Cryo-EM structure of 1059 SOSIP trimer purified via Galanthus nivalis lectin chromatography	VIRAL PROTEIN	
<a href="#">8QY1</a>	Xylanase from Bacillus circulans mutant E78Q/Y69A bound to xylohexaose	HYDROLASE	
<a href="#">8QY0</a>	Xylanase from Bacillus circulans mutant E78Q/Y69A bound to xylotriose	HYDROLASE	
<a href="#">9B6G</a>	Cryo-EM structure of the mouse TRPM8 channel in complex with the antagonist AMTB	MEMBRANE PROTEIN	<b>NAG</b>
<a href="#">8QXY</a>	Xylanase from Bacillus circulans mutant E78Q bound to xylotriose	HYDROLASE	
<a href="#">9B6H</a>	Cryo-EM structure of the mouse TRPM8 channel in complex with the antagonist TC-I 2014 and the cooling agonist C3	MEMBRANE PROTEIN	<b>NAG</b>
<a href="#">8QY3</a>	Xylanase from Bacillus circulans mutant E78Q/F125A bound to xylotriose	HYDROLASE	
<a href="#">8KCS</a>	Cryo-EM structure of human gamma-secretase in complex with BMS906024	MEMBRANE PROTEIN	
<a href="#">8OXS</a>	Cholera holotoxin variant (chimera with E. coli heat-labile enterotoxin, 4 C-terminal substitutions)	TOXIN	<b>aGal; bGal</b>
<a href="#">9BCG</a>	Myeloid cell leukemia-1 (Mcl-1) complexed with compound	APOPTOSIS	
<a href="#">8KB7</a>	Crystal structure of UDP/mannose-bound AGO61/beta-1,4-N-Acetylglucosaminyltransferase 2 (POMGNT2)	TRANSFERASE	<b>NAG (5) aMan (4)</b>
<a href="#">8RUR</a>	High pH (8.0) nitrite-bound MSOX movie series dataset 2 of the copper nitrite reductase (NirK) from Bradyrhizobium japonicum USDA110 [1.38 MGy]	OXIDOREDUCTASE	
<a href="#">8X85</a>	Structure of leptin-LepR dimer	CYTOKINE	
<a href="#">8Q1F</a>	D10N,P146A variant of beta-phosphoglucomutase from Lactococcus lactis in complex with native beta-glucose 1,6-bisphosphate intermediate	ISOMERASE	<b>1,6-di-O-phosphono-beta-D-glucopyranose</b>
<a href="#">8RR2</a>	3-keto-glycoside eliminase/hydratase in complex with alpha-3-keto-glucose	LYASE	<b>Hydroxymethyl)-3,5,6-tris(oxidanyl)oxan-4-one</b>
<a href="#">8TNU</a>	Cryo-EM structure of TRNM-b*01 Fab in complex with HIV-1 Env trimer BG505.DS SOSIP	VIRAL PROTEIN/IMMUNE SYSTEM	
<a href="#">8Y3U</a>	Ebola virus glycoprotein in complex with a broadly neutralizing antibody 2G1	ANTIVIRAL PROTEIN	
<a href="#">8X80</a>	Structure of leptin-LepR trimer with a small gap	CYTOKINE	
<a href="#">8TO9</a>	Cryo-EM structure of TRNM-f*01 Fab in complex with HIV-1 Env trimer ConC SOSIP	VIRAL PROTEIN/IMMUNE SYSTEM	