

9JP3	COMPLEX STRUCTURE OF ENDO-1,3-FUCANASE MUTANT D171A WITH FUCOTETRAOSE	HYDROLASE	Fucotetraose
9JOF	COMPLEX STRUCTURE OF ENDO-1,3-FUCANASE (FUN168D) FROM GH168 FAMILY WITH FUCOTRIOSE	HYDROLASE	
9JOO	COMPLEX STRUCTURE OF ENDO-1,3-FUCANASE (FUN168E) WITH FUCOTETRAOSES	HYDROLASE	Fucotetraose
9G3R	LecA from <i>Pseudomonas aeruginosa</i> in complex with a synthetic thiogalactoside	SUGAR BINDING PROTEIN	1-thio-bDGalp
9G3S	LecB from <i>Pseudomonas aeruginosa</i> in complex with a synthetic thiofucoside	SUGAR BINDING PROTEIN	aLFucp(1-1)-1-thio-bDGalp 1-thio-bDGalp(1-1)_aLFucp
9JOH	COMPLEX STRUCTURE OF ENDO-1,3-FUCANASE (FUN168D) WITH FUCOTETRAOSE FROM ISOSTICHOPUS BADIONOTUS	HYDROLASE	2OS-aLFucp(1-3)aLFucp(1-3) 2S4S-aLFucp(1-3)-2OS-aLFucp
8XZZ	Structure of a xylanase Xyl-1 M4 E175A in complex with xylobiose	HYDROLASE	
9JOG	COMPLEX STRUCTURE OF ENDO-1,3-FUCANASE (FUN168D) WITH FUCOTETRAOSE FROM HOLOTHURIA TUBULOSA	HYDROLASE	2OS-aLFucp(1-3)aLFucp(1-3) 2S4S-aLFucp(1-3)-2OS-aLFucp
9JP2	COMPLEX STRUCTURE OF ENDO-1,3-FUCANASE WITH FUCOTETRAOSE	HYDROLASE	aLFucp (1-3) 2OS4OS aLFucp(1-3)-2OS aLFucp(1-3)-2OSaLFucp
8Y00	Structure of a xylanase Xyl-1 M4 mutant E175A in complex with xylotriose	HYDROLASE	
8X05	X-Ray crystal structure of glycoside hydrolase family 6 cellobiohydrolase from <i>Phanerochaete chrysosporium</i> PcCel6A C240S/C393S soaked in cellobiose	HYDROLASE	
8X0Q	X-Ray crystal structure of glycoside hydrolase family 6 cellobiohydrolase from <i>Phanerochaete chrysosporium</i> PcCel6A C240S/C393S/D394C cocrystallized with cellobiose	HYDROLASE	aDManp (1-4) bDGlcNac (1-4) bDGlcNac
8X09	X-Ray crystal structure of glycoside hydrolase family 6 cellobiohydrolase from <i>Phanerochaete chrysosporium</i> PcCel6A C240S/C393S soaked in methyl-cellobiosyl-4-deoxy-4-thio-beta-D-cellobioside	HYDROLASE	bDGlcP (1-4) bDGlcP (1-4) bDGlcP (1-4) bDGlcP
9FNB	TMEM106B filaments from Biondi bodies (Biondi variant)	PROTEIN FIBRIL	bDGlcPNaC
8OXD	Structure of the human LYVE-1 (lymphatic vessel endothelial receptor-1) hyaluronan binding domain bound with decasaccharide hyaluronan.	CELL ADHESION	
8XZA	BA.2.86 Spike in complex with bovine ACE2 (Local refinement)	VIRAL PROTEIN	bDGlcPNaC
9CZF	Crystal structure of integrin avb6 headpiece in complex with compound MORF-627	CELL ADHESION/IMMUNE SYSTEM	
8Y32	BA.2.86-V483 RBD in complex with bACE2	VIRAL PROTEIN	bDGlcPNaC
9CZA	Crystal structure of integrin avb6 headpiece in complex with compound 18	CELL ADHESION/IMMUNE SYSTEM	
9CZD	Crystal structure of integrin avb6 headpiece in complex with compound 30	CELL ADHESION/IMMUNE SYSTEM	
9CZ7	Crystal structure of integrin avb6 headpiece in complex with compound 12	CELL ADHESION/IMMUNE SYSTEM	
8XZ8	BA.2.86 Spike in complex with bovine ACE2 (bound 1 ACE2)	VIRAL PROTEIN	
8Z2Y	High-resolution crystal structure of exo-beta-(1,3)-glucanase from <i>Aspergillus oryzae</i> (AoBgl) as a complex with glucose	HYDROLASE	bDGlcP
8RAL	CL3E peptide bound to the I-Ab murine MHC class II receptor	IMMUNE SYSTEM	
8Z2X	Crystal structure of exo-beta-(1,3)-glucanase from <i>Aspergillus oryzae</i> (AoBgl) as a complex with cellobiose	HYDROLASE	
9K7M	Coprinopsis cinerea GH131 protein CcGH131B E161A in complex with cellobiose	HYDROLASE	
8YWA	The structure of IgE receptor binding to IgE	MEMBRANE PROTEIN/IMMUNE SYSTEM	bDGlcPNaC
8QCM	ABC2 in complex with MZ82 and 5D3 Fab	TRANSPORT PROTEIN	
9FVE	Crystal structure of VcSiaP W73A mutant in complex with sialic acid and a VHH antibody (VHH_VcP#2)	TRANSPORT PROTEIN	Neu 5,9Ac
9JFF	<i>Nematostella vectensis</i> TRPM2 protomer in complex with ADPRP/Ca2+	MEMBRANE PROTEIN	
9GKU	Crystal Structure of Propanil hydrolase (PrpH) from <i>Sphingomonas</i> sp. Y57	HYDROLASE	bDGlcPNaC
9ARV	CryoEM structure of AMETA-A3	IMMUNE SYSTEM	bDGlcPNaC
8PXO	ABC2 in complex with AZ99 and 5D3 Fab	TRANSPORT PROTEIN	
9JJE	<i>Nematostella vectensis</i> TRPM2 tetramer in complex with ADPRP/Ca2+	MEMBRANE PROTEIN	
8X18	<i>Xanthomonas campestris</i> pv. <i>campestris</i> OpgD mutant-D379N with beta-1,2-glucan	TRANSFERASE	
8WEC	Crystal structure of <i>Arabidopsis thaliana</i> MIK2 ectodomain in complex with BAK1 ectodomain and SCOOP12	PLANT PROTEIN	