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# 7BF4 mutated.fasta

Created: today at 12:42

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## Template Results ⓘ

<a href="#">Templates</a>	<a href="#">Quaternary Structure</a>	<a href="#">Sequence Similarity</a>	<a href="#">Alignment</a>	<a href="#">More ▾</a>			
↑↓ Sort	Coverage	GMQE	QSQE	Identity	Method	Oligo State	Ligands
<input type="checkbox"/> ✓	<a href="#">6z5t.1.A Replicase polyprotein 1ab</a> <i>SARS-CoV-2 Macrodomain in complex with ADP-ribose</i>		0.97	-	98.83	X-ray, 1.6Å	monomer ✓
▼							1 x APR ⓘ
<input type="checkbox"/>	<a href="#">6woj.1.A Non-structural protein 3</a> <i>Structure of the SARS-CoV-2 macrodomain (NSP3) in complex with ADP-ribose</i>		0.97	-	98.26	X-ray, 2.2Å	monomer ✓
▼							1 x APR ⓘ
<input type="checkbox"/>	<a href="#">6wey.1.A Non-structural protein 3</a> <i>High-resolution structure of the SARS-CoV-2 NSP3 Macro X domain</i>		0.97	-	99.41	X-ray, 1.0Å	monomer ✓
▼							None
<input type="checkbox"/>	<a href="#">6ywl.2.A NSP3 macrodomain</a> <i>Crystal structure of SARS-CoV-2 (Covid-19) NSP3 macrodomain in complex with ADP-ribose</i>		0.97	-	99.41	X-ray, 2.5Å	monomer ✓
▼							1 x APR ⓘ
<input type="checkbox"/>	<a href="#">6ywk.3.A NSP3 macrodomain</a> <i>Crystal structure of SARS-CoV-2 (Covid-19) NSP3 macrodomain in complex with HEPES</i>		0.97	-	99.41	X-ray, 2.2Å	monomer ✓
▼							1 x MG ⓘ
<input type="checkbox"/>	<a href="#">6ywm.2.A NSP3 macrodomain</a> <i>Crystal structure of SARS-CoV-2 (Covid-19) NSP3 macrodomain in complex with MES</i>		0.97	-	99.41	X-ray, 2.2Å	monomer ✓
▼							1 x MG ⓘ
<input type="checkbox"/>	<a href="#">7bf6.1.A Papain-like protease nsp3</a> <i>Crystal structure of SARS-CoV-2 macrodomain in complex with remdesivir metabolite GS-441524</i>		0.96	-	99.41	X-ray, 2.1Å	monomer ✓
▼							1 x U08 ⓘ
<input type="checkbox"/>	<a href="#">7c33.1.A Non-structural protein 3</a> <i>Macro domain of SARS-CoV-2 in complex with ADP-ribose</i>		0.96	-	99.41	X-ray, 3.8Å	monomer ✓
▼							1 x APR ⓘ
<input type="checkbox"/>	<a href="#">2acf.2.A Replicase polyprotein 1ab</a> <i>NMR STRUCTURE OF SARS-COV NON-STRUCTURAL PROTEIN NSP3A (SARS1) FROM SARS CORONAVIRUS</i>		0.92	-	73.21	X-ray, 1.4Å	monomer ✓
▼							None
<input type="checkbox"/>	<a href="#">8cb3.2.A Papain-like protease nsp3</a> <i>SARS-CoV Macro domain complexed with 3-(N-morpholino)propanesulfonic acid</i>						

<u>↓Sort</u>	Coverage	GMQE	QSQE	Identity	Method	Oligo State	Ligands
▼	<div style="width: 80%; background-color: #0070C0; height: 10px;"></div>	0.92	-	73.21	X-ray, 1.6Å	monomer ✓	1 x <u>MPO</u> 
<input type="checkbox"/> <u>8cb3.3.A</u> Papain-like protease nsp3 <i>SARS-CoV Macro domain complexed with 3-(N-morpholino)propanesulfonic acid</i>							
▼	<div style="width: 80%; background-color: #0070C0; height: 10px;"></div>	0.92	-	73.21	X-ray, 1.6Å	monomer ✓	None
<input type="checkbox"/> <u>2fav.2.A</u> Replicase polyprotein 1ab (pp1ab) (ORF1AB) <i>Crystal structure of SARS macro domain in complex with ADP-ribose at 1.8 Å resolution</i>							
▼	<div style="width: 80%; background-color: #0070C0; height: 10px;"></div>	0.92	-	73.21	X-ray, 1.8Å	monomer ✓	1 x <u>APR</u> 
<input type="checkbox"/> <u>8cb3.1.A</u> Papain-like protease nsp3 <i>SARS-CoV Macro domain complexed with 3-(N-morpholino)propanesulfonic acid</i>							
▼	<div style="width: 80%; background-color: #0070C0; height: 10px;"></div>	0.91	-	73.21	X-ray, 1.6Å	monomer ✓	None
<input type="checkbox"/> <u>2fav.3.A</u> Replicase polyprotein 1ab (pp1ab) (ORF1AB) <i>Crystal structure of SARS macro domain in complex with ADP-ribose at 1.8 Å resolution</i>							
▼	<div style="width: 80%; background-color: #0070C0; height: 10px;"></div>	0.91	-	73.21	X-ray, 1.8Å	monomer ✓	None
<input type="checkbox"/> <u>2acf.1.A</u> Replicase polyprotein 1ab <i>NMR STRUCTURE OF SARS-COV NON-STRUCTURAL PROTEIN NSP3A (SARS1) FROM SARS CORONAVIRUS</i>							
▼	<div style="width: 80%; background-color: #0070C0; height: 10px;"></div>	0.91	-	73.21	X-ray, 1.4Å	monomer ✓	None
<input type="checkbox"/> <u>2fav.1.A</u> Replicase polyprotein 1ab (pp1ab) (ORF1AB) <i>Crystal structure of SARS macro domain in complex with ADP-ribose at 1.8 Å resolution</i>							
▼	<div style="width: 80%; background-color: #0070C0; height: 10px;"></div>	0.91	-	73.21	X-ray, 1.8Å	monomer ✓	None

The full list of templates matching your target sequence includes the following templates which are not in the list above. The full template list is available in [text](#) or [html](#) format.

1hjz.1.A, 1spv.1.A, 1ty8.2.A, 1vhv.1.A, 1yd9.1.A, 1yd9.3.A, 1zr3.2.A, 1zr3.3.A, 1zr5.1.A, 2acf.1.A, 2acf.2.A, 2acf.4.A, 2afc.1.A, 2bfq.1.A, 2dx6.1.A, 2eee.1.A, 2fav.1.A, 2fav.2.A, 2fav.3.A, 2fg1.1.A, 2fxk.2.A, 2l8r.1.A, 2lgr.1.A, 2vri.1.A, 2x47.1.A, 2xd7.1.A, 3ejf.1.A, 3ejg.1.A, 3eke.1.A, 3eti.1.A, 3ew5.1.A, 3ew5.2.A, 3ewo.1.A, 3ewo.1.B, 3ewq.1.A, 3gpg.2.A, 3gpq.3.A, 3gpq.4.A, 3gqe.1.A, 3gqe.2.A, 3gqo.3.A, 3gqo.4.A, 3iid.1.A, 3iif.1.A, 3iif.2.A, 3iif.3.A, 3jzt.4.A, 3q6z.1.A, 3q71.1.A, 3sig.1.A, 3sij.1.A, 3v2b.1.A, 3v45.1.A, 3vfq.1.A, 4abl.1.A, 4ess.1.A, 4etj.1.A, 4etk.1.A, 4etk.2.A, 4gua.1.A, 4gvv.1.A, 4gvv.2.A, 4gvv.3.A, 4gvw.1.A, 4iqy.1.A, 4j4z.1.A, 4j5q.1.A, 4j5r.2.A, 4j5s.4.A, 4jca.1.A, 4jgk.1.A, 4jgk.3.A, 4jll.1.A, 4jvv.1.A, 4k0c.1.A, 4k0c.2.A, 4kyb.1.A, 4kyb.2.A, 4uml.1.A, 5ail.1.A, 5cb3.1.A, 5cb5.1.A, 5cms.1.A, 5dus.1.A, 5e3b.1.A, 5fsu.1.A, 5fsu.2.A, 5fsz.1.A, 5fud.1.A, 5hih.1.A, 5ii.1.A, 5iq5.1.A, 5isn.1.A, 5kiv.1.A, 5l9k.1.A, 5lau.1.A, 5lbp.1.A, 5lcc.1.A, 5lcc.2.A, 5lnc.1.A, 5lnc.2.A, 5lw0.1.A, 5lw6.1.A, 5m31.1.A, 5m3e.1.A, 5m3i.1.A, 5mqx.1.A, 5o2d.1.A, 5rts.2.A, 5s4c.1.A, 5zdg.1.A, 5zu7.1.A, 6fx7.1.A, 6fy5.1.A, 6fy5.2.A, 6lfq.1.A, 6lf.1.A, 6lf.1.A, 6lfu.1.A, 6lfu.2.A, 6lh4.1.A, 6lh4.4.A, 6ln0.1.A, 6mea.1.A, 6r0g.2.A, 6vxs.2.A, 6w02.1.A, 6w02.2.A, 6w0t.1.D, 6wey.1.A, 6woj.1.A, 6y4y.1.A, 6y4z.1.A, 6yw.3.A, 6yw.2.A, 6ywm.2.A, 6z5t.1.A, 7bf6.1.A, 7c33.1.A, 7d2c.1.A, 7d3y.1.A, 7d3y.1.B, 7epu.1.B, 7h0s.2.A, 7h6j.1.A, 7h78.2.A, 7h7f.3.A, 7h7x.4.A, 7jme.1.A, 7kg3.1.A, 7kqw.1.A, 7kr0.1.A, 7kr1.1.A, 7ny6.1.A, 7ny7.1.A, 7omu.1.A, 7p27.1.A, 7twt.1.A, 7tx3.2.A, 7tx4.1.A, 7tx5.1.A, 7wr6.1.B, 8azc.1.B, 8azd.1.A, 8cb3.1.A, 8cb3.2.A, 8cb3.3.A, 8gia.1.A, 8gia.2.A, 8p0c.1.A, 8p0c.2.A, 8rsi.1.A, 8rsj.1.A, 8rsk.2.A, 8rl.1.A, 8rsm.1.A, 8rsn.1.A, 8sh6.1.A, 8sh6.2.A, 8sh8.1.A, 8tv6.1.A, 8tv7.1.A, 9fia.1.Z, 9gub.2.A, 9hqv.44.A, 9i05.48.A, 9qyd.1.A, 9qye.1.A, 9qyf.1.A, 9qyg.1.B, 9qyh.2.A

Build Models 0

Clear Selection