







# Phyre2

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Description	P08550
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Detailed template information

#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	<a href="#">c2qtyB_</a>	 Alignment		10.9	35	<b>PDB header:</b> hydrolase <b>Chain:</b> B: <b>PDB Molecule:</b> poly(adp-ribose) glycohydrolase arh3; <b>PDBTitle:</b> crystal structure of mouse adp-ribosylhydrolase 3 (marh3)
2	<a href="#">c3hfwA_</a>	 Alignment		8.2	7	<b>PDB header:</b> hydrolase <b>Chain:</b> A: <b>PDB Molecule:</b> protein adp-ribosylarginine hydrolase; <b>PDBTitle:</b> crystal structure of human adp-ribosylhydrolase 1 (harh1)
3	<a href="#">c3g9dB_</a>	 Alignment		5.9	38	<b>PDB header:</b> hydrolase <b>Chain:</b> B: <b>PDB Molecule:</b> dinitrogenase reductase activating <b>PDBTitle:</b> crystal structure glycohydrolase