



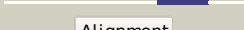

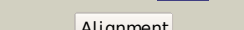





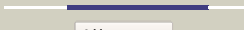

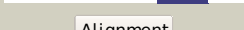









#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	c3e3vA_	 Alignment		14.6	17	PDB header: recombination Chain: A: PDB Molecule: regulatory protein recx; PDBTitle: crystal structure of recx from lactobacillus salivarius
2	c3se4B_	 Alignment		10.1	27	PDB header: immune system receptor Chain: B: PDB Molecule: interferon omega-1; PDBTitle: human ifnw-ifnar ternary complex
3	c3oq3A_	 Alignment		9.9	27	PDB header: cytokine/viral protein Chain: A: PDB Molecule: interferon alpha-5; PDBTitle: structural basis of type-I interferon sequestration by a poxvirus2 decoy receptor
4	d1au1a_	 Alignment		9.6	13	Fold: 4-helical cytokines Superfamily: 4-helical cytokines Family: Interferons/interleukin-10 (IL-10)
5	c3piwA_	 Alignment		9.6	7	PDB header: cytokine Chain: A: PDB Molecule: type i interferon 2; PDBTitle: zebrafish interferon 2
6	d1rh2a_	 Alignment		9.5	27	Fold: 4-helical cytokines Superfamily: 4-helical cytokines Family: Interferons/interleukin-10 (IL-10)
7	d1ogla_	 Alignment		9.2	18	Fold: all-alpha NTP pyrophosphatases Superfamily: all-alpha NTP pyrophosphatases Family: Type II deoxyuridine triphosphatase
8	d1b5la_	 Alignment		8.6	33	Fold: 4-helical cytokines Superfamily: 4-helical cytokines Family: Interferons/interleukin-10 (IL-10)
9	c3pivA_	 Alignment		8.3	20	PDB header: cytokine Chain: A: PDB Molecule: interferon; PDBTitle: zebrafish interferon 1
10	d1wu3i_	 Alignment		8.0	7	Fold: 4-helical cytokines Superfamily: 4-helical cytokines Family: Interferons/interleukin-10 (IL-10)
11	c3fwcO_	 Alignment		6.2	6	PDB header: cell cycle, transcription Chain: O: PDB Molecule: protein sus1; PDBTitle: sac3:sus1:cdc31 complex

12

[c2rrhA](#)

Alignment



5.7

56

PDB header:hormone
Chain: A: **PDB Molecule:**vip peptides;
PDBTitle: nmr structure of vasoactive intestinal peptide in methanol