

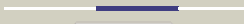





Phyre2

| | |
|---------------|------------------------------|
| Email | l.a.kelley@imperial.ac.uk |
| Description | C1P613 |
| Date | Wed Jan 25 15:20:07 GMT 2012 |
| Unique Job ID | a1328577bd5bd6b2 |

Detailed template information

| # | Template | Alignment Coverage | 3D Model | Confidence | % i.d. | Template Information |
|---|-------------------------|--|---|------------|--------|---|
| 1 | c2d9sA_ |  Alignment |  | 7.6 | 40 | PDB header: ligase Chain: A: PDB Molecule: cbl e3 ubiquitin protein ligase; PDBTitle: solution structure of rsgi ruh-049, a uba domain from mouse2 cdna |
| 2 | c2do6A_ |  Alignment |  | 6.8 | 40 | PDB header: ligase Chain: A: PDB Molecule: e3 ubiquitin-protein ligase cbl-b; PDBTitle: solution structure of rsgi ruh-065, a uba domain from human2 cdna |
| 3 | c2y35A_ |  Alignment |  | 5.7 | 53 | PDB header: hydrolase/dna Chain: A: PDB Molecule: ld22664p; PDBTitle: crystal structure of xrn1-substrate complex |