







Phyre2

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|---------------|------------------------------|
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| Description | Q9F5L1 |
| Date | Wed Jan 25 15:21:15 GMT 2012 |
| Unique Job ID | 48c3c7f309e11310 |

Detailed template information

| # | Template | Alignment Coverage | 3D Model | Confidence | % i.d. | Template Information |
|---|-------------------------|--|---|------------|--------|---|
| 1 | d1l1ca_ |  Alignment |  | 96.4 | 45 | Fold: GroES-like Superfamily: SacY-like RNA-binding domain Family: BglG-like antiterminator proteins |
| 2 | d1auua_ |  Alignment |  | 96.2 | 47 | Fold: GroES-like Superfamily: SacY-like RNA-binding domain Family: BglG-like antiterminator proteins |
| 3 | d1pk8a1 |  Alignment |  | 12.0 | 44 | Fold: PreATP-grasp domain Superfamily: PreATP-grasp domain Family: Synapsin domain |