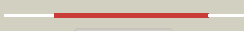
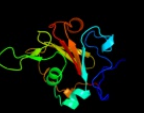
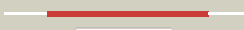
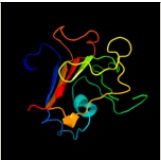


| #  | Template                | Alignment Coverage                                                                               | 3D Model                                                                            | Confidence | % i.d. | Template Information                                                                                                                                                                                                                                            |
|----|-------------------------|--------------------------------------------------------------------------------------------------|-------------------------------------------------------------------------------------|------------|--------|-----------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|
| 1  | <a href="#">c3fhhA_</a> | <br>Alignment   |    | 100.0      | 33     | <b>PDB header:</b> membrane protein<br><b>Chain:</b> A: <b>PDB Molecule:</b> outer membrane heme receptor shua;<br><b>PDBTitle:</b> crystal structure of the heme/hemoglobin outer membrane2 transporter shua from shigella dysenteriae                         |
| 2  | <a href="#">c2hdiA_</a> | <br>Alignment   |    | 100.0      | 45     | <b>PDB header:</b> protein transport,antimicrobial protein<br><b>Chain:</b> A: <b>PDB Molecule:</b> colicin i receptor;<br><b>PDBTitle:</b> crystal structure of the colicin i receptor cir from e.coli in complex2 with receptor binding domain of colicin ia. |
| 3  | <a href="#">c3qlbA_</a> | <br>Alignment   |    | 100.0      | 27     | <b>PDB header:</b> metal transport<br><b>Chain:</b> A: <b>PDB Molecule:</b> enantio-pyochelin receptor;<br><b>PDBTitle:</b> enantiopyochelin outer membrane tonb-dependent transporter from2 pseudomonas fluorescens bound to the ferri-enantiopyochelin        |
| 4  | <a href="#">c3efmA_</a> | <br>Alignment   |    | 100.0      | 18     | <b>PDB header:</b> membrane protein<br><b>Chain:</b> A: <b>PDB Molecule:</b> ferric alcaligin siderophore receptor;<br><b>PDBTitle:</b> structure of the alcaligin outer membrane recepteur fau from2 bordetella pertussis                                      |
| 5  | <a href="#">c1xkwA_</a> | <br>Alignment |  | 100.0      | 23     | <b>PDB header:</b> membrane protein<br><b>Chain:</b> A: <b>PDB Molecule:</b> fe(iii)-pyochelin receptor;<br><b>PDBTitle:</b> pyochelin outer membrane receptor fpta from pseudomonas2 aeruginosa                                                                |
| 6  | <a href="#">d1by5a_</a> | <br>Alignment |  | 100.0      | 25     | <b>Fold:</b> Transmembrane beta-barrels<br><b>Superfamily:</b> Porins<br><b>Family:</b> Ligand-gated protein channel                                                                                                                                            |
| 7  | <a href="#">d2gufa1</a> | <br>Alignment |  | 100.0      | 38     | <b>Fold:</b> Transmembrane beta-barrels<br><b>Superfamily:</b> Porins<br><b>Family:</b> Ligand-gated protein channel                                                                                                                                            |
| 8  | <a href="#">d1kmoa_</a> | <br>Alignment |  | 100.0      | 27     | <b>Fold:</b> Transmembrane beta-barrels<br><b>Superfamily:</b> Porins<br><b>Family:</b> Ligand-gated protein channel                                                                                                                                            |
| 9  | <a href="#">c1xkhC_</a> | <br>Alignment |  | 100.0      | 27     | <b>PDB header:</b> membrane protein<br><b>Chain:</b> C: <b>PDB Molecule:</b> ferripyoverdine receptor;<br><b>PDBTitle:</b> pyoverdine outer membrane receptor fpva from pseudomonas aeruginosa2 pao1 bound to pyoverdine                                        |
| 10 | <a href="#">c2grxB_</a> | <br>Alignment |  | 100.0      | 25     | <b>PDB header:</b> metal transport<br><b>Chain:</b> B: <b>PDB Molecule:</b> ferrichrome-iron receptor;<br><b>PDBTitle:</b> crystal structure of tonb in complex with fhua, e. coli2 outer membrane receptor for ferrichrome                                     |
| 11 | <a href="#">c2iahA_</a> | <br>Alignment |  | 100.0      | 27     | <b>PDB header:</b> membrane protein<br><b>Chain:</b> A: <b>PDB Molecule:</b> ferripyoverdine receptor;<br><b>PDBTitle:</b> crystal structure of the ferripyoverdine receptor of the outer2 membrane of pseudomonas aeruginosa bound to ferripyoverdine.         |

|    |                        |           |                                                                                  |       |    |                                                                                                                                                                                                                                  |
|----|------------------------|-----------|----------------------------------------------------------------------------------|-------|----|----------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|
| 12 | <a href="#">c1po3A</a> | Alignment |  | 100.0 | 28 | <b>PDB header:</b> membrane protein<br><b>Chain:</b> A: <b>PDB Molecule:</b> iron(iii) dicitrate transport protein feca<br><b>PDBTitle:</b> crystal structure of ferric citrate transporter feca in2 complex with ferric citrate |
|----|------------------------|-----------|----------------------------------------------------------------------------------|-------|----|----------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|

|    |                         |           |                                                                                  |       |    |                                                                                                                                                                                                                                                            |
|----|-------------------------|-----------|----------------------------------------------------------------------------------|-------|----|------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|
| 13 | <a href="#">c3cslB_</a> | Alignment |  | 100.0 | 24 | <p><b>PDB header:</b>membrane protein/heme binding protein</p> <p><b>Chain:</b> B: <b>PDB Molecule:</b>hasr protein;</p> <p><b>PDBTitle:</b> structure of the serratia marcescens hemophore receptor hasr in2 complex with its hemophore hasa and heme</p> |
|----|-------------------------|-----------|----------------------------------------------------------------------------------|-------|----|------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|