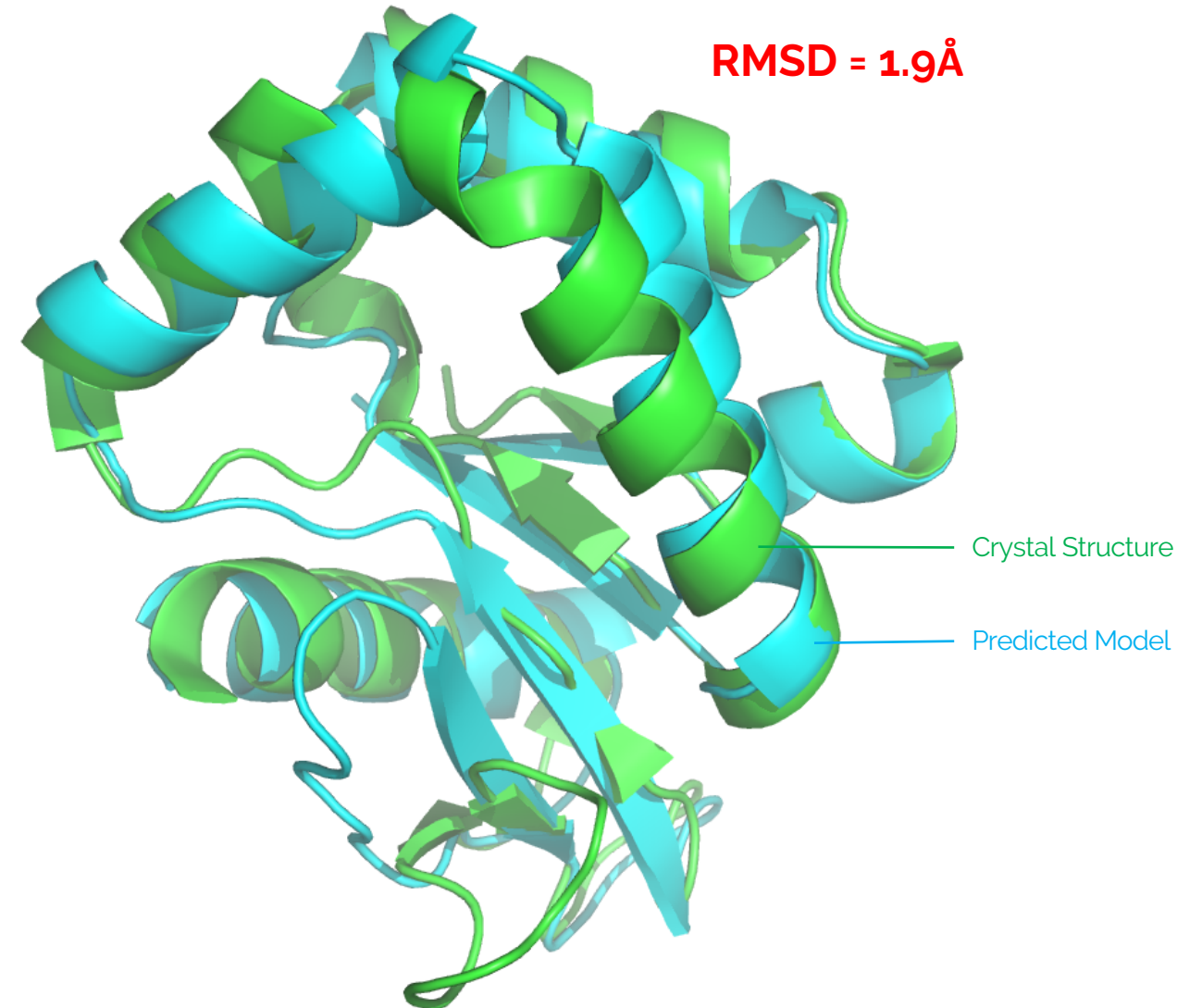


# Accurate Models of Very Low Homology Structures

**RMSD = 1.9Å**

24 different templates with sequence ID  $\leq$  18%



| Template PDB_ID | Coverage | Sequence Identity |
|-----------------|----------|-------------------|
| 4q6tA           | 67.24    | 15.38             |
| 2ddxA           | 62.93    | 16.44             |
| 3aquD           | 64.66    | 10.67             |
| 1ydoB           | 61.21    | 7.04              |
| 5jh8A           | 66.38    | 10.39             |
| 3wijA           | 70.69    | 10.98             |
| 4f9dA           | 57.76    | 17.91             |
| 4uriB           | 69.83    | 12.35             |
| 1hjxB           | 67.24    | 11.54             |
| 1v47A           | 56.03    | 12.31             |
| 2ftpA           | 62.07    | 8.33              |
| 3co4A           | 66.38    | 5.19              |
| 1hjbB           | 66.38    | 11.69             |
| 5hciD           | 62.93    | 1.37              |
| 3bg9C           | 62.93    | 10.96             |
| 4zquA           | 70.69    | 13.29             |
| 1ljbA           | 67.24    | 12.82             |
| 1m03A           | 62.93    | 8.22              |
| 1owqA           | 66.38    | 12.99             |
| 3cz8A           | 67.24    | 8.97              |
| 4s3jA           | 64.66    | 6.67              |
| 5jd3H           | 62.07    | 9.72              |
| 1hjbB           | 67.24    | 11.54             |
| 4g6uA           | 72.41    | 13.1              |