|  |  |  |  |
| --- | --- | --- | --- |
| GPCR | 3D-Coordinates | Ramachandran Plot Summary from Procheck | Overall model qualityProSA-web |
| Most favouredregions | Additionallyallowed regions | Generouslyallowed regions | Disallowedregions | Z-Score |
| hADRB2  | PDBID 3P0G | 88,1% | 11,9% | 0,0% | 0,0% | -1,88 |
| PDBID 2RH1 | 94,2% | 5,4% | 0,0% | 0,0% | -2,52 |
| hTAAR6active-like  | Selected Model | 92,1% | 7,9% | 0,0% | 0,0% | -1,51 |
| SA-loop refinement | 93,5% | 6,5% | 0,0% | 0,0% | -1,77 |
| hTAAR6inactive-like  | Selected Model | 89,6% | 9,2% | 0,8% | 0,4% | -2,17 |
| SA-loop refinement | 92,9% | 6,3% | 0,8% | 0,0% | -2,34 |
| hTAAR8active-like  | Selected Model | 89,8% | 9,8% | 0,4% | 0,0% | -1,60 |
| SA-loop refinement | 91,2% | 8,8% | 0,0% | 0,0% | -1,84 |
| hTAAR8inactive-like  | Selected Model | 90,5% | 8,7% | 0,4% | 0,4% | -2,75 |
| SA-loop refinement | 93,3% | 6,7% | 0,0% | 0,0% | -2,93 |
| zTAAR13cactive-like | Selected Model | 94,1% | 5,5% | 0,4% | 0,0% | -2,57 |
| SA-loop refinement | 95,1% | 4,9% | 0,0% | 0,0% | -2,61 |
| zTAAR13cinactive-like | Selected Model | 95,3% | 4,3% | 0,4% | 0,0% | -2,66 |
| SA-loop refinement | 96,5% | 2,8% | 0,8% | 0,0% | -2.71 |