Table S3 The root-mean-square distance (RMSD) matrix of the thiolases estimated by Swiss-PdbViewer

|  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- |
|  | BmorT1-1 | DmelCT | PxutAB-1 | BmorTFE | BmorSCP2 (type-1) |
| BmorT2 | 0.91 | 0.89 | 0.91 | 1.16 | 1.37 |
| BmorT1-1 |  | 0.25 | 0.91 | 1.19 | 2.01 |
| DmelCT |  |  | 0.89 | 1.16 | 1.95 |
| PxutAB-1 |  |  |  | 1.12 | 1.49 |
| BmorTFE |  |  |  |  | 2.03 |