Amino acid positions corresponding to each UGGT domain taken as reference sequences.

|  |  |  |  |
| --- | --- | --- | --- |
|  | UDP-g\_GGTase  (Recognition domain) | GT8\_HUGT1\_C\_like  (Catalytic domain) | Reference |
| HUGT1 | 45-1202 | 1256-1503 | [link](https://www.ncbi.nlm.nih.gov/Structure/cdd/wrpsb.cgi?seqinput=NP_064505.1) |
| HUGT2 | 33-1189 | 1231-1478 | [link](https://www.ncbi.nlm.nih.gov/Structure/cdd/wrpsb.cgi?seqinput=NP_064506.3) |
| Ce-UGGT-a | 22-1170 | 1202-1449 | [link](https://www.ncbi.nlm.nih.gov/Structure/cdd/wrpsb.cgi?INPUT_TYPE=live&SEQUENCE=NP_509268.1) |
| Ce-UGGT-b | 20-1065 | 1095-1343 | Relative to CeUGGT-a |