## ConSurf Results for job:VGLUT3 consurfseqres date:10/03/2023

| 1<br>MP <mark>FKAFD</mark> TFK           | 11<br>EKIL <mark>K</mark> PG <mark>KE</mark> G         | 21<br>VKNAVGDSL <mark>G</mark>                 | 31<br>ILQRKID <mark>G</mark> TT                     | 41<br>EEEDNIELNE                               |
|--|--|--|---|--|
| 51 EGR <mark>PVQTS</mark> RP             | 61<br>SPPLCDCH <mark>C</mark> C                        | 71<br>GLPK <mark>R</mark> YI <mark>I</mark> AI | 81<br>Msgl <mark>gfcis</mark> f                     | 91<br>GIRCNLGVAI                               |
| 101<br>VEMVNNSTVY                        | 111<br>VDGKPEIQTA                                      | 121<br>QFNWDPETVG                              | 131<br>LI <mark>HGSFFW</mark> GY                    | 141<br>IMTQIPGGFI                              |
| 151 SNKFAANRVF                           | 161 GAAIFLTSTL   | 171<br>NMFIPSAARV                              | 181<br>HY <mark>GCVMCV</mark> RI                    | 191<br><b>L Q <mark>G</mark> L V E G V T Y</b> |
| 201<br>PACHGMWSKW                        | 211 <b>APPLERSRLA</b>                                  | 221 TTSFCGSYAG eeeebee f                       |   |  |
| 251 FYIYGMFGII eeeeeeeee                 | 261  WYMFWLLQAY  e e e e e e e e e e e e e e e e e e e | 271 ECPAAHPTIS eeeeeeee f f f f f              | 281 NEEKTYIETS eeeeeeee                             | 291 IGEGANVVSL                                 |
| 301  SKFSTPWKRF  e e e e b e e e e b f f | 311 FTSLPVYAII eeeebbbb ff s                           |  |   | 341 <b>FEEVFGFAIS beeebeeee</b>                |
|  | 361 MVMTIVVPIG e e b e e b b e e b                     |  | 381  Q I L T T T A V R K  e e e e e e e b b e f s f |  |
|  | 411 HTKGVAISFL   | 421 V L A V G F S G F A b e b b e b b e b      |   |  |
| 451 GISNGVGTLS eebeeebeeb ff ff f        |  | 471  MTRHKTREEW  ebeeeebs s                    |   | 491 HYSGVIFYGV beebeeebee                      |
| 501  FASGEKQEWA  ebebee f                | 511  DPENLSEEKC  | 521<br>GIIDQDELAE                              | 531 <b>E I E L N H E S F A</b>                      | 541<br>SPKKKMSYGA                              |
| 551<br>T <mark>SQNCEVQK</mark> K         | 561<br>E <mark>WK</mark> GQRGATL                       | 571<br><b>DEEELTSYQN</b>                       | 581<br>E <mark>ERNFSTIS</mark>                      | 591  |

The conservation scale:

- e An exposed residue according to the NACSES algorithm.
- b A buried residue according to the NACSES algorithm.
- f A predicted functional residue (highly conserved and exposed).
- s A predicted structural residue (highly conserved and buried).
- X Insufficient data the calculation for this site was performed on less than 10% of the sequences.