ConSurf Results for job:NETnat date:14/08/2023

1	11	21	31	41
MLLARMNPQV	Q P E N N G A D T G	P E Q P L R A R K T	AELLVVKERN	GVQCLLAPRD
51 GDAQPRETWG ebes	61 KKIDFLLSVV eeebbebbbb s	71 GFAVDLANVW bbbbbbbbbbbsssss	81 RFPYLCYKNG ebbebbeebe fssf s sf	91 GGAFLIPYTL bbbbeebbee ss sffss
101 FLIIAGMPLF bbeebbbbb	111 YMELALGQYN bbbbbbbbeb s ss	121 REGAATVWKI eebbbbbbee f	131 CPFFKGVGYA beebebbbeb f s s	141 VILIALYVGF bbebbebbb
151 YYNVIIAWSL bbbbbbbbbb	161 YYLFSSFTLN ebbebbeeee s s	171 LPWTDCGHTW bebeeeeee sfs f f	181 NSPNCTDPKL bbeebebeee	191 LNGSVLGNHT
201 KYSKYKFTPA ebbeeebbb	211 AEFYERGVLH bbbbbebbe	221 LHESSGIHDI beebebeee	231 GLPQWQLLLC eeebeebeeb f	241 LMVVVIVLYF beebbeebee
251 SLWKGVKTSG bbeebbeebb fs fss	261 KVVWITATLP eebeebbeeb ff f ss fs	271 Y F V L F V L L V H e e b b e e b e e e f f f	281 GVTLPGASNG beeeebeeb ff f s	291 INAYLHIDFY eeebeeeee f
301 RLKEATVWID eeeeebbebe f sf f	311 AATQIFFSLG bbbbbbbbbb s s s	321 AGFGVLIAFA bbbbbbbbbb	331 SYNKFDNNCY bebeeebebe sfs f f	341 RDALLTSSIN ebbeebbeeb s
351 CITSFVSGFA eebbeebbeb	361 IFSILGYMAH bbbbbbbbbee ss s s	371 EHKVNIEDVA eeeeebeebe	381 TEGAGLVFIL beeebbbbbb f s s	391 YPEAISTLSG bbbbbeebee
401 STFWAVVFFV eeebbeebbe ffss	411 MLLALGLDSS bbbbbbbbbbb s s s	421 MGGMEAVITG bbebbbbbbbb	431 LADDFQVLKR ebeeeeee	
451 FSTFLLALFC eebbeebeeb f	461 ITKGGIYVLT bbebbbebb	471 LLDTFAAGTS bbeebbebb f	481 ILFAVLMEAI bbebbebbe	491 GVSWFYGVDR bbbeebebee sf
501 FSNDIQQMMG bbebbeebee s f	511 FRPGLYWRLC eebeebeeb	521 WKFVSPAFLL beeebbebbe s	531 FVVVVSIINF ebbebeeee	541 KPLTYDDYIF eebeeeeb
551 PPWANWVGWG eeebbeebee f s	561 IALSSMVLVP ebeebbeebe	571 IYVIYKFLST eee	581 QGSLWERLAY	591 GITPENEHHL

601 611 VAQRDIRQFQ LQHWLAI

The conservation scale:



Variable

Average

Conserved

- 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.