| | UniprotKB ID | Entry name | organism | full name | oglcnacscore | oglcnac sites | phosphorylation sites | PMIDS | sequence | intracellular | extracellular | cytosol | nucleus | mitochndrion | endoplasmic reticulum | golgi apparatus | plasma membrane | extracellular region |
|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|
| **0** | K2NC82 | K2NC82_TRYCR | Trypanosoma cruzi marinkellei | NaN | 19.946178 | NaN | NaN | 30984116 | MSQTQETMASATLDALRAASASGH<br>MSALREVKADAVLLDEANALPELLR<br>LCGSIEAGHNLALMNAAWKLLLSVS<br>SLNTISPPVYSGILQFMLSQLHHHVF<br>SHDWSDAKLIKLASFVASHVVSAVR<br>AHPSIAARDERLVQSIVGLHAIVCLT<br>VATSGHNEAAQQLHQSIVVRLLAILE<br>AMGSSFSASSPLSAGEHLRLLLQCL<br>ADPPPAATAELATASVAAGSSVAALA<br>AMLAATRSFQVCDRAYVERILAAFLL<br>WLPDMQLVFSDVAEQEEKTEAAETA<br>RTTVLAASCDKDPDTVLCHRIVRHA<br>RNLPRAVFLPEQNGDGAAESSGEM<br>HLNAQLLLLLRSLACRSVDAIRGGE<br>WCDKEGSPGGGSLLRCRVFSLLMTA<br>LLVPNDTSAARVLLVWEAVLSRLSN<br>ADREGLGMELLQLGEEQLRALRELP<br>AESRGRATCLRRLLNALAVLLTEICR<br>QLQRPPKSLHEAFAETTATRISSSNP<br>TSFCTLVNVLYHQSSVYQWWNEKSI<br>PKEEVTGDTEEEAKAIHDELRAVLTE<br>WQDHSSSKTNSDMASRICMTIASVP<br>ERVHNILDRTNSPKSITAVLLIATKVL<br>QKLTEELNPSYAICECARFVAAELVA<br>VTGFVLSEDPTLLACIRRLVDISFAIT<br>DDGKEVQVDGMLRYHAAVCLRNFA<br>QRNAVVPLGQLRLSEAVTTALRTLL<br>QGGDVADTDVSMRDCEGFLAHRLE<br>EWKSLMSRSGHVAKSSQSADSLVEF<br>APQATSTFSQSVIDGHEGQFFLPLQR<br>GEETVRSVLQWLKAGRTLQPSEEET<br>LVRLEQLVSAAANMTRHALSSNGDV<br>FCSSADIGAGATQTTGLSPEVINVD | None | None | None | None | None | None | None | None | None |