

| UniprotKB ID | Entry name | organism | full name | oglcnacscore | oglcnac sites | phosphorylation sites | PMIDS | sequence |
|--------------|------------|--------------|-----------------------------|--------------|---------------------|--|--|--|
| Q5T5P2 | SKT_HUMAN | Homo sapiens | Sickle tail protein homolog | 10.760686 | S169;S357;S743;S880 | S169;T244;S361;S365;T393;T470;S474;S526;S809;S1027;S1030;S1033;S1044;S1461;S1739;S1841;S1896;S1899;S1902 | 30379171;31492838;28657654;30620550;28510447 | MEENESQKCEPCLPYSADRRQMQE QGGKGNLHVTSPEDAECRRTKERLSN GNSRGSVSKSSRNIPRRHTLGGPRS SKEILGMQTSSEMDRKREAFLEHLKQ KYPHHAASAIMGHQERLRDQTRSPKL SHSPQPPSLGDPVEHLSETSADSLE AMSEGDAPTPFSRGSRTRASLPVVR STNQTKERSLGVLYLQYGDETKQLR MPNEITSADTIRALFVSAFPQQLTMK MLESPSVAIYIKDESRNVYELNDVR NIQDRSLLKVYNKPAHAFNHTPKT MNGDMRMQRELVYARGDGPVAPR PGSTAHPHAIIPNSPPSTVPVPHSMPP SPSRIPYGGTRSMVVPGNATIPRDRI SSLPVSRLPSPSAILERRDVKPDED MSGKNIAMYRNEGFYADPYLYHEG RMSIASSHGGHPLDVPDHIIAYHRTA IRSASAYCNPSMQAEMHMEQSLYR QKSRKYPDLSHLPTLGSKTPPASPHR VSDLRMIDMHAHYNAHGPPHTMQP DRASPSRQAFKKEPGLVYIEKPRSA AGLSSLDLGPPLMEKQVFAYSTATI PKDRETRERMQAMEKQIASLTGLVQ SALFKGPITSYSKDASSEKMMKTTA NRNHTDSAGTPHVSGGKMLSALLES TVPPSQPPPVGTSIAHMSLLEMRRS VAELRLQLQMRQLQLQNQELLRA MMKKAELISGKVMETMKRLEDPV QRQRVLVEQERQKYLHEEEKIVKKL CELEDFVEDLKKDSTAASRLVTLKD VEDGAFLLRQVGEAVATLKGEFPTL QNKMRAILRIEVAVRFLKEEPHKL DSLLKRVRSMTDVLTMLRRHVTDGL LKGTDAAQAAQYMAMEKATAAEVL KSQEEAAHTSGQPFHSTGAPGDAKS EVPPLSGMMVRHAQSSPVVIQPSQH SVALLNPAQNLPHVASSPAVPQEAT STLQMSQAPQSPQIPMNGSAMQSL FIEEIHVSVAKNRAVSIEKAEKKWEE KRQNLDHYNGKEFEKLEEAQANI MKSIPNLEMPPATGPLPRGDAPVDK VELSESPNSEQDLEKLGKSPPPP PPPPRRSYLPGSGLTTTRSGDVVYTG RKENITAKASSEDAGPSPQTRATKYP AEEPASAWTPSPPPVTTSSSKDEEEE EEEGDKIMAELQAFQKCSFMDVNS NSHAEPSRADSHVKDTRSGATVPPK EKKNLEFFHEDVRKSDVEYENGPQ MEFQKVTTGAVRPSDPPKWERGME NSISDASRTSEYKTEIIMKENSISNM SLLRDSRNYSQETVPKASFGFGISG LEDEINKGSKISGLQYSIPDTENQTL NYGKTKEMEKQNTDKCHVSSHTRL TESSVHDFKTEDQEVITDDFGQVVL RPKEARHANVNPNEDESSSSSPT ENAATDNIAFMITETTQVQLSSGEV HDIVSQKGEDIQTVNIDARKEMTPR |

