	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	Q810B9	SLIK3_MOUSE	Mus musculus	SLIT and NTRK-like protein 3	24.474581	NaN	NaN	33300544	MMKPSIAEMLHRGRMLWIILLSTIA LGWTTPIPLIEDSEEIDEPCFDPCYCE VKESLFHIHCDSKGFTNISQITEFWS RPFKLYLQRNSMRRLYTNSFLHLNN AVSINLGNNALQDIQTGAFNGLKILK RLYLHENKLDVFRNDTFLGLESLEY LQADYNVIKRIESGAFRNLSKLRVLI LNDNLIPVLPTNLFKAVSLTHLDLRG NRLKVLFYRGMLDHIGRSLMELQLE ENPWNCTCEIVQLKSWLERIPYTAL VGDITCETPFHFHGKDLREIKKTELC PLLSDSEVEASLGIPHLSSSKENAWP TKPSSMLSSVHFTASSVEYKSSNKQ PKPTKQPRTPRPPSTSQALYPGPNQP PIAPYQTRPPIPIICPTGCTCNLHIND LGLTVNCKERGFNNISELLPRPLNA KKLYLSSNLIQKIYRSDFWNFSSLDL LHLGNNRISYVQDGAFINLPNLKSLF LNGNDIEKLTPGMFRGLQSLHYLYF EFNVIREIQPAAFSLMPNLKLLFLNN NLLRTLPTDAFAGTSLARLNLRKNY FLYLPVAGVLEHLNAIVQIDLNENP WDCTCDLVPFKQWIETISSVSVVGD VLCRTPENLTHRDVRTIELEVLCPE MLHIAQPGPSPPQPGDYHPNGGPTS ASPYEFSPPGGPVPLSVLILSLLVLFF SAVFVAAGLFAYVLRRRRKKLPFRSK RQEGVDLTGIQMQCHRLFEDSGGN SGGSGGGGRPTLSSPEKAPPVGHVY EYIPHPVTQMCNNPIYKPREEEEVAA SAAQDTGATDRGGPGTQPTGMAEVL LGSEQFAETPKENHSNYRTLLEKEK EWALAVSNSQLNTIVTVNHHHPHP HHSAVGGVSGVGGGTGGDLAGFRH HEKNGGVVLFPPGGGCGGGSLLLD RERPQPAPCTVGFVDCLYGTVPKLK ELHVHPPGMQYPDLQQDARLKETLL FSAGKGFTDHQTPKSDYLDLRAKLQ TKPDYLEVLEKTAYRF	None	None	None	None	None	None	None	None	None