

UniprotKB ID	Entry name	organism	full name	oglcnaicscore	oglcnaicsites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracellular region
075094	SLIT3_HUMAN	Homo sapiens	Slit homolog 3 protein	27.086674	S1067	NaN	28657654	MAPGWAGVGAAVRARLALALASV LSGPPAVACPTKCTCSAASVDC HGL GLRAVPRGIPRNERLDLDRNNITRI TKMDFAGLKNLRVLHLEDNQVSVIE RGAFQDLKQLERLRLNKNKLOVLPE LLFQSTPKLTRLDLSENQIQGIPRKA FRGITDVKNLQLDNNHISCIEDGAF RALRDLEILTLNNTNISRILVTSFNH MPKIRTLRLHSNHLYCDCHLAWLSD WLRQRRTVGQFTLCMAPVHLRGFN VADVQKKEYVCPAPHSEPPSCNANS ISCPSPCTCSNNIVDCRGKGLMEIPA NLPEGIVEIRLEQNSIKAIPAGAFQY KCLKRIDISKNQISDIAPDAFQGLKSL TSLVLYGNKITEIVKGLFDGLVSLQL LLL NANKINCLRVNTFQDLQNLNLL SLYDNKLOQTSKGLFAPLQSIQTLHL AQNPFVCDCHLKW LADYLDQNPDIET SGARCSSPRLANKRISQIKSKKFRFC SGSEDYRSRFSSECFMDLVCPEKCR CEGTIVDCSNQKLVRIPSHLPEYVTD LRLNDNEVSVLEATGIFKKLPNLRKI NLSNNKIKEVREGAFDGAASVQELM LTGNQLETVHGRVFRGLSGLKTLML RSNLIGCVSNDTFAGLSSVRLSLYD NRITITPGAFTTLVSLSTINLLSNPF NCNCHLAWLGKWLKRKRIVSGNPR CQKPFFLKEIPIQDVAIQDFTCDGNE ESSCQLSPRCPEQCTC METVVRC SN KGLRALPRGMPKDVTLEYLEGNHLT AVPRELSALRHLLTLDLSNNSISMLT NYTFSNMSHLSTLILSYNRLRCIPVH AFNGLRSLRVLTLHGNDISSVPEGSF NDLTSLSHLALGTNPLHCDCSLRWL SEWVKAGYKEPGIARCSSPEPMADR LLLTTPTHRFQCKGPVDINIVAKCNA CLSSPCKNNGTCTQDPVELYRCACP YSYKGDCTVPINTCQNPCQHGGT CHLSDSHKDFSCSCPLGFEGQRCE INPDDECDNDCENNATCV DGINNY VICPPNYTGELCDEVIDHCVPELNL CQHEAKCIPLDKGFSCCEVPGYSGK LCETDNDDCVAHKCRHGAQCVDTI NGYTCTCPQGFSGPFCEHPPMVLL QTSPCDQYECQNGAQCI VVQEQEPTC RCPPGFAGPRCEKLITVNFVKGDSYV ELASAKVRPQANISLQVATDKDNGIL LYKGDNDPLALELYQGHVRLVYDSL SSPPTTVYSVETVNDGQFHSVELVTL NQTLNLVVDKGT PKSLGKLQKQPAV GINSPLYLGGIPTSTGLSALROGTD R PLGGFHGCIHEVRINNELQDFKALP PQSLGVSPGCKSCTVCKHGLCRSVE KDSVVCECRPGWTGPLCDQEARDP CLGHRCHHGKCVATGTSYMCKCAE GYGGDLCDNKNDSANACSAFKCHH GQCHISDQGEPYCLCPGFSGEHCQ QENPCLGQVVREVIRROKGYASCAT ASKVPIMECRGGCGPQCCQPTRSKR RKYVFQCTDGSSFVEEVERHLECCG LACS	True	True	1.724	2.399	5.0	1.539	1.266	3.716	5.0