	UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochndrion	endoplasmic reticulum	gol app
0	Q14160	SCRIB_HUMAN	Homo sapiens	Protein scribble homolog	9.37818	T475;S764;S1140	S37;T378;T475;S504;S688;T689;S708;S764;T826;S835;S853;S875;S939;S1140;S1220;S1223;S1226;S1232;S1276;S1279;S1295;S1298;S1306;S1309;T1342;S1348;S1378;S1437;S1445;S1448;S1475;S1486;S1508;S1541;T1545;S1547;S1561;S1566	30379171;38665916;33214551;23301498;34019948;34725712;32119511;29351928	MLKCIPLWRCNRHVESVDKRHCSL QAVPEEIYRYSRSLEELLLDANQLRE LPKPFFRLLNLRKLGLSDNEIQRLPP EVANFMQLVELDVSRNDIPEIPESIK FCKALEIADFSGNPLSRLPDGFTQLR SLAHLALNDVSLQALPGDVGNLANL VTLELRENLLKSLPASLSFLVKLEQL DLGGNDLEVLPDTLGALPNLRELWL DRNQLSALPPELGNLRRLVCLDVSE NRLEELPAELGGLVLLTDLLLSQNLL RRLPDGIGQLKQLSILKVDQNRLCEV TEAIGDCENLSELILTENLLMALPRS LGKLTKLTNLNVDRNHLEALPPEIG GCVALSVLSLRDNRLAVLPPELAHTT ELHVLDVAGNRLQSLPFALTHLNLK ALWLAENQAQPMLRFQTEDDARTG EKVLTCYLLPQQPPPSLEDAGQQGS LSETWSDAPPSRVSVIQFLEAPIGDE DAEEAAAEKRGLQRRATPHPSELKV MKRSIEGRRSEACPCQPDSGSPLPA EEEKRLSAESGLSEDSRPSASTVSEA EPEGPSAEAQGGSQQEATTAGGEED AEEDYQEPTVHFAEDALLPGDDREI EEGQPEAPWTLPGGRQRLIRKDTPH YKKHFKISKLPQPEAVVALLQGMQP DGEGPVAPGGWHNGPHAPWAPRAQ KEEEEEEEGSPQEEEEEEEEENRAE EEEASTEEEDKEGAVVSAPSVKGVS FDQANNLLIEPARIEEEELTLTILRQT GGLGISIAGGKGSTPYKGDDEGIFISR VSEEGPAARAGVRVGDKLLEVNGVA LQGAEHHEAVEALRGAGTAVQMRV WRERMVEPENAVTITPLRPEDDYSP RERRGGGLRLPLLPPESPGPLRQRH VACLARSERGLGFSIAGGKGSTPYRA GDAGIFVSRIAEGGAAHRAGTLQVG DRVLSINGVDVTEARHDHAVSLLTA ASPTIALLLEREAGGPLPPSPLPHSSP PTAAVATTSITTATPGVPGLPSLAPSL LAAALEGPYPVEEIRLPRAGGPLGLSI VGGSDHSSHPFGVQEPGVFISKVLP RGLAARSGLRVGDRILAVNGQDVRD ATHQEAVSALLRPCLELSLLVRRDPA PPGLRELCIQKAPGERLGISIRGGAR GHAGNPRDPTDEGIFISKVSPTGAAG RDGRLRVGLRLLEVNQQSLLGLTHG EAVQLLRSVGDTLTVLVCDGFEAST DAALEVSPGVIANPFAAGIGHRNSLE SISSIDRELSPEGPGKEKELPGQTLH WGPEATEAAGRGLQPLKLDYRALAA VPSAGSVQRVPSGAAGGKMAESPCS PSGQQPPSPPSPDELPANVKQAYRA FAAVPTSHPPEDAPAQPPTPGPAASP EQLSFRERQKYFELEVRVPQAEGPP KRVSLVGADDLRKMQEEEARKLQQ KRAQMLREAAEAGAEARLALDGETL GEEEQEDEQPPWASPSPTSRQSPAS PPPLGGGAPVRTAKAERRHQERLRV QSPEPPAPERALSPAELRALEAEKRA LWRAARMKSLEQDALRAQMVLSRS QEGRGTRGPLERLAEAPSPAPTPSPT PVEDLGPQTSTSPGRLSPDFAEELRS LEPSPSPGPQEEDGEVALVLLGRPSP GAVGPEDVALCSSRRPVRPGRRGLG PVPS	None	None	None	None	None	None	No