

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracellular region
043150	ASAP2_HUMAN	Homo sapiens	Arf-GAP with SH3 domain, ANK repeat and PH domain-containing protein 2	19.176706	S770	S701;S822;T903	39531497;38665916	MPDQISVSEFVAETHEDYKAPTASSF TTRTAQCRNTVAAIEEALDVDVDRMVL YKMKKSVKAINSSGLAHVENEQYT QALEKFGGNCVCRDDPDLGSAFLKF SVFTKELTALFKNLIQNMNNIISFPL DSSLKGDLDKGVKGDLDKPPDKAWK DYETKITKIEKEKKEHAKLHGMIRTE ISGAEIAEEMEKERRFFQLQMCEYL LKVNEIKIKKGVDLLQNLIKYFHAQC NFFQDGLKAVESLKPSETLSTDLHT IKQAQDEERRQLIQLRDILKSAQVE QKEDSQIRQSTAYSLHQPOGKKEHG TERNGLYKKSDBGIRKVVQKRKCSV KNGFLTISHGTANRPPAKLNLLTCQ VKTNPEEKCCFDLISHDRTYHFQAE DEQECQIWMMSVLQNSKEEALNNAF KGDDNTGENNIVQELTKEHSEVQR MTGNDVCCDCGAPDPTWLSTNLGI LTCIECSGIHRELGVHYSRMOQLTLD VLGTSELLAKNIGNAGFNEIMECC LPAEDSVKPNPGSDMNARKDYITAK YIERRYARKKHADNAAKLHSLCEAV KTRDIFGLLQAYADGVDLTEKIPLAN GHEPDETALHLAVRSVDRTSLHVD FLVQNSGNLDKQTKGKGSTALHYCCL TDNAECLKLLLRGKASIEIANESGET PLDIKRLKHEHCEELLTQALSGRF NSHVHVEYEWRLLEDLDESDDDM DEKLQSPNRRREDRPISFYQLGSNQ LQSNVAVSLARDAANLAKEKQRAFMP SILQNETYGALLSGSPPPAQPAAPST TSAPPLPPRNVGKVQTASSANTLWK TNSVSDGGSRQRSSDPPAVHPPL PPLRVTSTNPLTPTPPPVAKTSPVM EALSQPSKPAPPGISQIRPPPLPQPP SRLPQKKPAPGADKSTPLTNKGQPR GPVDLSATEALGPLSNAMVLQPPAP MPRKSQATKLPKRVKALYNADVADN PDELTFSEGDVIVDGEEDQEWVIG HIDGDPGRKGAFVSVVHFIAD	True	True	4.668	2.6	1.254	1.203	5.0	5.0	1.235