	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	A6NIR3	AGAP5_HUMAN	Homo sapiens	Arf-GAP with GTPase, ANK repeat and PH domain-containing protein 5	29.902547	S555	NaN	39894887;37340703	MGNILTCCVHPSVSLEFDQQQGSVC PSESEIYEAGAGDRMAGAPMAAAVQ PAEVTVEVGEDLHMHHIRDQEMPE ALEFSLSANPEASTIFQRNSQTDALE FNPSANPEASTIFQRNSQTDVVEIRR SNCTNHVSTERFSQQYSSCSTIFLD DSTASQHYLTMTIISVTLEIPHHITQR DADRSLSIPDEQLHSFAVSTVHITKN RNGGGSLNNYSSSIPSTPSTSQEDP QFSVPPTANTPTPVCKRSMRWSNLF TSEKGSHPDKERKAPENHADTIGSG RAIPIKQGMLLKRSGKWLKTWKKKY VTLCSNGVLTYYSSLGDYMKNIHKK EIDLRTSTIKVPGKWPSLATSACAPIS SSKSNGLSKDMDTGLGDSICFSPSIS STTSPKLNPPPSPHANKKKHLKKKS TNNFMIVSATGQTWHFEATTYEERD AWVQAIQSQILASLQSCESSKSKSQL TSQSEAMALQSIQNMRGNAHCVDY ETQNPKWASLNLGVLMCIECSGIHR SLGTRLSRVRSLELDDWPVELRKVM SSIGNDLANSIWEGSSQGQTKPSVK STREEKERWIRSKYEEKLFLAPLPCT ELSLGQHLLRATADEDLQTAILLLAH GSREEVNETCGEGDGCTALHLACRK GNVVLAQLLIWYGVDVMARDAHGN TALTYARQASSQECINVLLQYGCPDE CV	False	False	2.123	1.312	False	False	False	False	1.219