	UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochndrion	endoplasmic reticulum	golgi appara
0	Q92667	AKAP1_HUMAN	Homo sapiens	A-kinase anchor protein 1, mitochondrial	23.348165	S580	T70;S105;S107;S151;S169;S429;S445;T447;T533;S573;T590;S592	35083852;30379171;32119511;38253038	MAIQFRSLFPLALPGMLALLGWWW FFSRKKGHVSSHDEQQVEAGAVQL RADPAIKEPLPVEDVCPKVVSTPPSV TEPPEKELSTVSKLPAEPPALLQTHP PCRRSESSGILPNTTDMRLRPGTRR DDSTKLELALTGGEAKSIPLECPLSS PKGVLFSSKSAEVCKQDSPFSRVPR KVQPGYPVVPAEKRSSGERARETGG AEGTGDAVLGEKVLEEALLSREHVL ELENSKGPSLASLEGEEDKGKSSSS QVVGPVQEEEYVAEKLPSRFIESAHT ELAKDDAAPAPPVADAKAQDRGVEG ELGNEESLDRNEEGLDRNEEGLDR NEESLDRNEEGLDRNEEIKRAAFQII SQVISEATEQVLATTVGKVAGRVCQA SQLQGQKEESCVPVHQKTVLGPDTA EPATAEAAVAPPDAGLPLPGLPAEGS PPPKTYVSCLKSLLSSPTKDSKPNISA HHISLASCLALTTPSEELPDRAGILV EDATCVTCMSDSSQSVPLVASPGHC SDSFSTSGLEDSCTETSSSPRDKAIT PPLPESTVPFSNGVLKGELSDLGAED GWTMDAEADHSGGSDRNSMDSVD SCCSLKKTESFQNAQAGSNPKKVDL IIWEIEVPKHLVGRLIGKQGRYVSFL KQTSGAKIYISTLPYTQSVQICHIEGS QHHVDKALNLIGKKFKELNLTNIYA PPLPSLALPSLPMTSWLMLPDGITVE VIVVNQVNAGHLFVQQHTHPTFHAL RSLDQQMYLCYSQPGIPTLPTPVEIT VICAAPGADGAWWRAQVVASYEETN EVEIRYVDYGGYKRVKVDVLRQIRSD FVTLPFQGAEVLLDSVMPLSDDDQF SPEADAAMSEMTGNTALLAQVTSYS PTGLPLIQLWSVVGDEVVLINRSLVE RGLAQWVDSYYTSL	None	None	None	None	None	None	None