

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
Q92667	AKAP1_HUMAN	Homo sapiens	A-kinase anchor protein 1, mitochondrial	17.153812	S580	T70;S105;S107;S151;S169;S429;S445;T447;T533;S573;T590;S592	38253038;32119511;30379171;35083852	MAIQFRSLFPLALPGMLALLGWWW FFSRKKGHVSSHDEQQVEAGAVQL RADPAIKEPLPVEDVCPKVVSTPPSV TEPPEKELSTVSKLPAEPPALLQTHP PCRRSESSGILPNTTDMRLRPGTRR DDSTKLELALTGGEAKSIPLECLSS PKGVLFSKSAEVCKQDSPFSRVPR KVQPGYPVVPAEKRSSGERARETGG AEGTGDAVLGEKVL EEALLSREHVL ELENKGP SLASLEGEEDK GKSSSS QVVG PVQ EEEYVAEKLPSRFIESAHT ELAKDDAAPAPPVADAKAQDRGVEG ELGNEESLDRNEEGLDRNEEGLDR NEESLDRNEEGLDRNEEIKRAAFQII SQVISEATEQVLATTVGKVAGRVCQA SQLQGQKEESCVPVHQKTVLGPDTA EPATAEAAVAPPDAGLPLPGLPAEGS PPPPTYVSCLKSLSSPTKDSKPNISA HHISLASCLALTPSEELPDRAGILV EDATCVTCMSDSSQSVPLVASPGHC SDSFSTSGLEDSC TETSSSPRDKAIT PPLPESTVPF SNGVLKGELSDLGAED GWTMDAEADHSGGSDRNSMDSVD SCCSLKKTESFQNAQAGSNPKKVDL IIWEIEVPKHLVGR LIGKQGRYVSFL KQTSGAKIYISTLPYTQSVQICHIEGS QHHVDKALNLIGKKFKELNLNIYA PPLPSLALPSLPMTSWLMLPDGITVE VIVVNQVNAGHLFVQQHTHPTFHAL RSLDQQMYLCYSQPGIPTLPTPVEIT VICAAPGADGAWWRAQVVASYEETN EVEIRYVDYGGYKRVKVDVLRQIRSD FVTLPFQGAEVLLDSVMPLSDDDQF SPEADAAMSEMTGNTALLAQVTSYS PTGLPLIQLWSVVGDEVVLLINRSLVE RGLAQWVDSYYTSL

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