	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	ex re
0	Q96P48	ARAP1_HUMAN	Homo sapiens	Arf-GAP with Rho-GAP domain, ANK repeat and PH domain-containing protein 1	26.040516	T722	S229;T354;S428;T431;T504;S738;S1428;S1435	34725712;29351928	MAEAGDAALSVAEWLRALHLEQYT GLFEQHGLVWATECQGLSDTRLMD MGMLLPGHRRRILAGLLRAHTSPAP APRPTPRPVPMKRHIFRSPPVPATPP EPLPTTTEDEGLPAAPPIPPRRSCLPP TCFTTPSTAAPDPVLPPLPAKRHLAE LSVPPVPPRTGPPRLLVSLPTKEEES LLPSLSSPPQPQSEEPLSTLPQGPPQ PPSPPPCPPEIPPKPVRLFPEFDDSDY DEVPEEGPGAPARVMTKKEEPPPSR VPRAVRVASLLSEGEELSGDDQGDE EEDDHAYEGVPNGGWHTSSLSLSLP STIAAPHPMDGPPGGSTPVTPVIKAG WLDKNPPQGSYIYQKRWVRLDTDH LRYFDSNKDAYSKRFISVACISHVAAI GDQKFEVITNNRTFAFRAESDVERK EWMQALQQAMAEQRARARLSSAYL LGVPGSEQPDRAGSLELRGFKNKLY VAVVGDKVQLYKNLEEYHLGIGITFI DMSVGNVKEVDRRSFDLTTPYRIFS FSADSELEKEQWLEAMQGAIAEALS TSEVAERIWAAAPNRFCADCGAPQP DWASINLCVVICKRCAGEHRGLGAG VSKVRSLKMDRKVWTETLIELFLQL GNGAGNRFWAANVPPSEALQPSSSP STRRCHLEAKYREGKYRRYHPLFGN QEELDKALCAAVTTTDLAETQALLG CGAGINCFSGDPEAPTPLALAEQAG QTLQMEFLRNNRTTEVPRLDSMKP LEKHYSVVLPTVSHSGFLYKTASAGK LLQDRRAREEFSRRWCVLGDGVLSY FENERAVTPNGEIRASEIVCLAVPPP DTHGFEHTFEVYTEGERLYLFGLES AEQAHEWVKCIAKAFVPPLAEDLLA RDFERLGRLPYKAGLSLQRAQEGWF SLSGSELRAVFPEGPCEEPLQLRKLQ ELSIQGDSENQVLVLVERRRTLYIQG ERRLDFMGWLGAIQKAAASMGDTL SEQQLGDSDIPVIVYRCVDYITQCGL TSEGIYRKCGQTSKTQRLLESLRQDA RSVHLKEGEQHVDDVSSALKRFLRD LPDGLFTRAQRLTWLEASEIEDEEE KVSRYRELLVRLPPVNRATVKALISH LYCVQCFSDTNQMNVHNLAIVFGPT LFQTDGQDYKAGRVVEDLINHYVVV FSVDEEELRKQREEITAIVKMRVAGT ASGTQHAGDFICTVYLEEKKAETEQ HIKVPASMTAEELTLEILDRRNVGIR EKDYWTCFEVNEREEAERPLHFAEK VLPILHGLGTDSHLVVKKHQAMEA MLLYLASRVGDTKHGMMKFREDRS LLGLGLPSGGFHDRYFILNSSCLRLY KEVRSQRPWSGAPETSHRPEKEWPI KSLKVYLGVKKKLRPPTCWGFTVVH ETEKHEKQQWYLCCDTQMELREWF ATFLFVQHDGLVWPSEPSRVSRAVP EVRLGSVSLIPLRGSENEMRRSVAA FTADPLSLLRNV	None	None	None	None	None	None	None	None	N