

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
Q9QWY8	ASAP1_MOUSE	Mus musculus	Arf-GAP with SH3 domain, ANK repeat and PH domain-containing protein 1	21.849389	T823	T308;S732;S741;S854;S858;S1026;S1045;S1059;T1066;S1146	34678516;22517741	MRSSASRLSSFSSRDSLWNRMPDQI SVSEFIAETTEDYNSPTTSSFTTRLH NCRNTVTLLEEALDQDRALQKVKK SVKAIYNSGQDHSVQNEENYAQVLDK FGSNFLSRDNPDLGTAFVKFSTLTK ELSTLLKNLLQGLSHNVIFTLDSLLK GDLKGVKGDLLKPPDKAWKDYETKF TKIEKEKREHAKQHGMIRTEITGAEI AEEMEKERRLFQLQMCEYLKIVNEI KTKKGVDLLQNLIKYYHAQCFFQD GLKTADKLKQYIEKLAADLYNIKQTQ DEEKKQLTALRDLIKSSLQLDPKEVG GLYVASRANSSRRDSQSRQGGYSM HQLQGNKEYGSEKKGFLLKKS DGIR KVVQRRKCAVKN GILTISHATSNRQ PAKLNLLTCQVKPNAEDKKSFDLIS HNRTYHFQAEDEQDYIAWISVLTNS KEEALTMAFRGEQSTGENSLEDLTK AIIEDVQRLPGNDICDCGSSSEPTWL STNLGILTCIECSGIHREMGVHISRI QSLELDKLGTSSELLAKNVGNNSFN DIMEANLPSPSPKPTSSDMTVRKE YITAKYVDHRFSRKTCASSSAKLNEL LEAIKSRDLLALIQVYAEGVELMEPL LEPGQELGETALHLAVRTADQTS LH LVDFLVQNCGNLDKQTSVGN TVLH YCSMYGKPECLKLLLRSKPTVDIVN QNGETALDIKRLKATQCEDLLSQA KSGKFNPHVHVEYEWNL RQDEMDE SDDDLDDKPSPIKKERSPRPQS FCH SSSISPODKLALPGFSTPRDKQ RLSY GAFTNQIFASTSTDLPTSPTSEAPPLP PRNAGKGPTGPPSTLPLGTQTSSGSS TLSKKRPPPPPGHKRTLSDPPSPLP HGPPNKGAI PWGNDVGPLSSSKTAN KFEGLSQASTSSAKTALGPRVLPKL POKVALRKTETSHHLSLDR TNIPPET FOKSSQLTELPQKPPLGELPPKPV EL PKPQLSDLPPKQMKDLPPKQLGD LLAKSQAGDVSAKVQPPSEVTQRSH TGDLSPNVQSRDAIQKQASEDSNDL TPTLPETPVPLPRKINTGKNKVR RVK TIYDCQADNDELTFIEGEVIIVTGEE DQEW WIGHIEGQPERKGVFPVSVFH ILSD