

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
P39993	GEA2_YEAST	Saccharomyces cerevisiae	ARF guanine-nucleotide exchange factor 2	9.365641	NaN	S46;S284	33229814	MSDREFVTVDPVTIIIKECINLSTAM RKYSKFTSQSGVAALLGGGSEIFSNQ DDYLAHTFNNLNTNKHNDPFLSGFI QLRLMLNKLKLNLDNIDSLTILQPFL IVSTSSISGYITSLALDSLQFFTLNII NESSQNYIGAHRAVNALTHCRFEG SQQLSDDSVLLKVVFLRSIVDSPYG DLLSNSIYDVLQTILSLACNNRRSE VLRNAAQSTMIAVTVKIFSKLKTIEPV NVNQIYINDESYTNDVLKADTIGTNV ESKEEGSQEDPIGMKVNNEEAISED DGIEEHHSEKSTNGAEQLDIVQKT TRSNSRIQAYADDNYGLPVVRQYLN LLLSLIAPENELKHSYSTRIFGLELIQ TALEISGDRQLYPRFLTISDPFKSI LFIIQNTTKLSLLQATLQLFTTLVVIL GNNLQLQIELTLTRIFSILLDDGTAN NSSSENKNKPSIIKELLIEQISILWTR SPSFFTSTFINFDCNLDRAVDSINFL KALTKLALPESALTTTESVPPICLEGL VSLVDDMFDHMKDIDREEFGRQKN EMEILKKRDRKTEFIECTNAFNEKP KKGIPMLIEKGFASDSKDIAEFLF NNNNRMNKKTIGLLLCHPKVSLL NEYIRLFDGSLRVDEAIRILLTKFRL PGESQQIERIIEAFSSAYCENQDYDP SKISDNAEDDISTVQPDADSVFILSYS IIMLNTDLHNPQVKEHMSFEDYSGN LKGCCNHKDFPFWYLDRIYCSIRDK EIVMPEEHHGNEKWFEDAWNLLIS STTVITEIKKDTQSVMDKLTPLELLN FDRAIFKQVGPSIVSTLFNIYVVASDD HISTRMITSLDKCSYISAFFDFKDLF NDILNSIAKGTTLINSSHDELSTLA FEYGPMPVQIKFEDTNTTEIPVSTDA VRFGRSFKGQLNTVVFFRIIRNKDP KIFSKELWLNIVNIILTYEDLILSPDI FPDLQKRLKLSNLPKPSPEISINKSK ESKGLLSTFASYLKGDEEPTEEEIKS SKKAMECIKSSNIAASVFGNESNITA DLIKTLLDSAKTEKNADNSRYFEAEL LFIIELTIALFLFCKEEKELGKFIQKV FQLSHTKGLTKRTVRRMLTYKILLIS LCADQTEYLSKLINDELKKGDIQFTQ KFFATNQGKEFLKRLFSLTESEFYR GFLLGNENFWKFLRKVTAMKEQSE SIFEYLNESIKTDSNILTNNENFMWVL GLLDEISSMGAVGNHWEIEYKKLTE SGHKIDKENPYKKSIELSLKSIQLTS

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