

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracellular region
P36123	SA190_YEAST	Saccharomyces cerevisiae	SIT4-associating protein SAP190	24.475439	NaN	S774;S857;S862;S892;T990;S991	33229814	MSGSFWKFGQDYSIESPVSKILNSA FKINKDQDDVPTGTCEENIADDE DNSSHDYAASEDNVNVNENEKEEEE NTLPITSEYENYRPNLDVLDLDDLLD DDELYTELMCSNFKLLIFLKYPDVLS KLIEYVTNEKILDEETDSAKKPEIIEG VNDHPILIERDRKDKKEDAEEGGDS EETTNSDHDSDGERSVDSEETSIT LPPESEEQVETRRARIAAEILSADVW PISAAIMQNKDLLGRLWSILDHPAPL PIPASTYFMKINERLLDMDITGMLEF ILSRDSLVARFLTHVDNPSLMDFFLL KVISTDKPDSPTGVIKILKSOELIPKL LDHLNPEYGISTQSAAGDFIKAFVTL STNSSNELASGIGPNELTRQLVSEE MIEKLIKIMLKGSTLSNGVGIIEELIR KNNSDYDFIQLVYTTLESHPPDRDP IHLIHLVKLFAKHMPDFADMLDKTK LPLMEMPFGNIEPLGFERFKICELIA ELLHCSNMTLLNEPNGEMIAQERDI ERAKELETSTEKENITAIVDNKSSYY DKDCVEKDITENLGAQINNQGSEE DELNDTGVSSVKLDVKSADAKVVEGL ENDASGVELYDETLSDTESVRECLR EKPLVGDRLKIALEDTKILISILDMFT EFPWNNFRLHNVIQFIAQIFNGPLK TGYNRFLKDYLVDAYLTKKIVDADK ACQDYKKTGLRYGYMGLHTLVAEE ISKFKEYIDEMKLTFCNTAVSDRLEE PFWKEYSETILADTREKYNTVLGDF GNDQESDDVIRNSDSEDHIGDTEG NENYNGENDELLENGHDSGNMMD LYYFNNEEENEEDYAEYSVDV NKNYNNVETNDDYDSDDGKSKS AESEFTDKISEHRDGNLYNEDNDE NGSDKWTSGTSLFPPDHFPSRSQPS DPKLQDQNFHHQDFEGVGDGDDDD YMDPNDDGQSYARPGNPLYTTPKTP PRPKTILFNSLSALDNNGEDEEVAL GTSVDDRMNEISSDEEDEDDEE NDMGNEEGYSLYRSRKEAF	None	None	None	None	None	None	None		

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