

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
P38903	2A5D_YEAST	Saccharomyces cerevisiae	Serine/threonine-protein phosphatase 2A 56 kDa regulatory subunit delta isoform	9.365641	NaN	T242;T257	33229814	MMRGFKQRLIKKTTGSSSSSSSKKK DKEKEKEKSSTTSSTSKKPASASSSS HGTTSSASSTGSKSTTEKKGQSGS VPSQGKHHSSSTSKTKTATTPSSSSS SSRSSVSRSGSSSTKKTSSRKGQEQ SKQSQQPSQSQKQSSSSSAAIMNP TPVLTVTKDDKSTSGEDHAHPTLLG AVSAVPSSPISNASGTAVSSDVENGN SNNNNMNINTSNTQDANHASSQSI DIPRSSHSFERLPTPTKLNPDTDLELI KTPQRHSSSRFEPSTRYPTLTKLPNFN EVSPEERIPFLIAKVDQCNTMFDNF DPSFDIQGKEIKRSTLDELIEFLVTNR FTYTNEMYAHVVNMFKINLFRPIPP PVNPVGDIDPDEDEPVNELAWPH MQAVYEFFLRFVESPDFNHQIAKQY IDQDFILKLELFDSEDIRERDCLKTT LHRIYGKFLSLRSFIRSMNNIFLQFI YETEFNGVAELLEILGSIINGFALPL KEEHKVFLVRILIPLHKVRCLSLYHP QLAYCIVQFLEKDLLTEEVVMGLLR YWPKINSTKEIMFLNEIEDIFEVIEPL EFIKVEVPLFVQLAKCISSPHFQVAE KVLSYWNNEYFLNLCIENAEVILPIIF PALYELTSQLELDTANGEDSISDPYM LVEQAINSGSWNRAIHAMAFKALKI FLETNPVLYENCNALYLSSVKETQQ RKVQREENWSKLEEYVKNLRINND KDQYTIKNPELRNSFNTASENNTLN EENENDCDSEIQ

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