

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
P31374	PSK1_YEAST	Saccharomyces cerevisiae	Serine/threonine-protein kinase PSK1	9.365641	NaN	S10;S192;S202;S255;S286;S327;S926;S1018;S1023;S1035;S1055;T1079	33229814	MPYIGASNLSEHSFVNLKEKHAITH KGTSSSVASLQTPPSPDQENHIDNE LENYDTSLSDVSTPNKKEGDEFEQS LRDTFASFRKTKPPPLDFEQPRLPS TASSVDSTVSSPLTDEDIKELEFLP NESTHSYSYNPLSPNSLAVRLRILKR SLEIIIQNPSMLLEPTPDDLPLKEFA GRRSSLPRTSASANHLMNRNKSQI WNTTSATLNAFVNNTSSSSAASSAL SNKKPGTPVFPNLDPTHSTQTFHRAN SLAYLPSILPEQDPLLKHNNSLFRGD YGNNISPERPSFRQPFKDQTSNLRN SLLNERAYQEDETFLPHHGPSMDL LNEQRANLKSLLNLLNETLEKNTSE RASDLHMISLFLNLKMLGDPKKN NSERDKRTEKLLKILLDSLAEPFFEH YNFIGDNPIADTDELKEEIDEFTGSG DTTAITDIRPQQDYGRILRTFTSTKNS APQAIFTCSQEDPWQFRAANDLACL VFGISQNAIRALTLMDLIHTDSRNFV LHKLLSTEGQEMVFTGEIIGIVQPET LSSSKVWASFWAKRKNGLLVCVFE KVPCDYVDVLLNLDDFGAENIVDKC ELLSDGPTLSSSSTLSLPKMASSPTG SKLEYSLEKILEKSYTKPTSTENRN GDENQLDGDHSEPSLSSSPVRSKK SVKFANDIKDVKSISQSLAKLMDDV RNGVVFDPDDDLLPMPKVCNHINE TRYFTLNHLSYNIPCAVSSVLEDEL KLKIHS LPYQAGLFIVDSHTLDIVSS NKSILKNMFGYHFAELVGKSITEIIPS FPKFLQFINDKYPALDITLHKNKGLV LTEHFFRKIQAIEIMGDRKSFYTSVGI DGLHRDGCEIKIDFQLRVMNSKVILL WVTHSRDVVFEEYNTNPSQLKMLK ESELMLSSASSASSSKSSSRIST GTLKMSNLSTYEDLAHRTNKLKYE IGDDSRHSQSTLSEQEQVPLENDK DSGEMMLADPEMKHKLELARIYSR DKSQFVKEGNFKVDENLIISKISLSP STESLADSKSSGKGLSPLEEEKLIDE NATENGLAGSPKDEDGIIMTNKRGN QPVSTFLRTPEKNIGAQKHVKKFSD FVSLQKMGE GAYGKVNLCIHKKNRY IVVIKMIFKERILVDTWVRDRKLG TIP SEIQIMATLNK KPHENILRLLDFFED DDYYYIETPVHGETGCIDLFDLIEFKT NMTEFEAKLIFKQV VAGIKHLHDQG IVHRDIKDENVIVDSKGFVKIIDFGSA

