

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
P33894	STE13_YEAST	Saccharomyces cerevisiae	Dipeptidyl aminopeptidase A	24.475439	NaN	NaN	33229814	MSASTHSHKRKNSHLFPQRKSSNS SMDKPPFPNNDVANTDPQSNENG HTINEIRPTEATIDVTDVPPQTPFLQE QYSMRPRRESFQFNDIENQHHTHS FFSVNKFNRWGEWLSPEKRSYVL VFTLIALSVLVLVILIPSKLLPTKITR PKTSAGDSSLGKRFSIENVLNGDF AIPEDTFHFIDPPQRLGQSDPGLY FTTKEIDGHTNFIKQLFDETFEVNL GGNRFLYEGVEFTVSTVQINYKLDK LIFGTNLESEFRHSSKGFYWKDLNT GNIEPILPPEKSDDNYELGLSKLSYA HFSPAYNYIFVYENNLFLQQVNSG VAKKVTEDGSKDIFNAKPDWIYEEE VLASDQAIWWAPDDSKAVFARFNDT SVDDIRLNRYTNMNEAYLSDTKIKYP KPGFQNPQFDLFLVNLQNGIYSINT GGQKDSILYNGKWISPDTRFEITDR NSKILDVKVYDIPSSQMLTVRNTNS NLFNGWIEKTKDILSIPPKPELKRMD YGYIDIHADSRGFSHLFYPTVFAKE PIQLTKGNWEVTGNGIVGYEYETDTI FFTANEIGVMSQHLYSISLTDSTQN TFQSLQNPDKYDFYDFELSSSARYA ISKKLGPDTPIKVAGPLTRVLNVAEIH DDSILQTKDEKFKEKIKNYDLPITSY KTMVLDDGVEINYIEIKPANLNPKKK YPILVNIYGGPGSQTFTTKSSLAFEQ AVVSGLDVIVLQIEPRGTGGKGSWF RSWAREKLGWEPDITEVTKKFIQ RNSQHIDESKIAIWGWSYGGFTSLK TVELDNGDTFKYAMAVPVTNWTLY DSVYTERYMNQPSENHEGYFEVSTI QNFKSFESLKRFLFVHGTFDNDVHI QNTFRLVDQLNLLGLTNYDMHIFPD SDHSIRYHNAQRIVFQKLYYWL RDA FAERFDNTEVLHL	None	None	None	None	None	None	None	None	

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