

UniprotKB ID	Entry name	organism	full name	oglnacscore	oglnac sites	phosphorylation sites	PMIDS	sequence
P53037	PSD2_YEAST	Saccharomyces cerevisiae	Phosphatidylserine decarboxylase proenzyme 2	9.365641	NaN	NaN	33229814	MRIIKGRKRGKKNKKPTLILKIHVIQAE NIEALKTFNCNPVCFVTTNTFYQSQT NKLKNSNTHWNQTLRIKLPRNPTSE WLRHIVYDALPTGAPPTTPSRPRTTTA NTSSSTLSNSGLSSHSHSRNLNVT SKGNQSTSTINSVSSSATPAPSHSSS SLSTTGPGSTHKNRINSYLVLGEAKI SLLDLFKRKDTTTSYKFSIEAQRVHL YDMKGGKDQDSLNCNFLVGDILLG FKLECNVKRTPTFQAFNAWRNELN TYLGRIDRNKARMRSSSSLPPLED MLSNSSAVSGNEIRREKPYSDTDLA HDEEVNAEDEIDAEEESIEDMNSSGS ICTERRYDIDNDTIFDSISEVVSLNDE ELDILNDFEADHPNVPDINVHDID EDTRISLSSMITALDEYDIVEPEDVA KLPVSENDITSVDDEESENQQESD EEFDIYNEDEREDSDFQSKEYIGSRL LHLQRGKHNSYANYLYRRAKSNFF ISKKEHAMGVVFMHIGAIKNLPALR NRLSKTNYEMDPFIVISFGRRVFKTS WRKHTLNPEFNEYAAFEVFPHETN FAFSIKVVDKDSFSFNDDVAKCELA WFDMLQQQHHENEWIPYEIPLDLT VEPAHAPKQPVLVYSSFKYVSYPFLLK SFWKEAVDTSVNLERLDIIQVMLYL ERLGSFTMADSFELFQHFNKSAWA GQSITRSQVLEGLQSWRKSTNFKRI WTCPRCMRSCKPTRNARRSKLVLE NDLITHFAICTFSKEHKTLPKPSYVSS AFASKRWFSKVLKLTYGKYALGSN NANILVQDRDTGIIIEEKISAHVKLG MRIIYNGKSPESKKFRSLLKTLIRQ GKKFDSTASAKQIEPFIKFHSLDLSQ CRDKDFKTFNEFFYRKLKPGSRLPE SNNKEILFSPADSRACTVFPTIQESKEI WVKGRKFSIKKLANNYNPETFNDN NCSIGIFRLAPQDYHRFHSPCNGTIG KPVYVDGEYYTVNPM AVRSELDFVG ENIRVIIPIDSPQFGKLLYPIGAMMV GSILLTCKENDVVEGQELGYFKFG GSTIIIIIPHNNFMFSDLVKNSSERI ETLVKVGMSIGHTSNVNELKRIRIKV DDPKKIERIKRTISVSDENAKSTGNV TWEYHTLREMMNKDFAGL