

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
Q03868	SPO71_YEAST	Saccharomyces cerevisiae	Prosopore membrane adapter protein SPO71	24.475439	NaN	NaN	33229814	MDSIVNVVEDDVKYAQRVTSFSSPQ NANVKVFTIPRHSFTAFLRSYVSPTE LSACSQVTLLGGIPKQWYADQNNQV WKLTKISLRKVRKQSDMLRRYGYG TIYKKRVGKIPTALYLRRKHFTWSYED NTSIHNGHRLKEAEMEMKRTRSSP VQKSEYKLSLPRRCRSSSDQNFMRQ ELLKEKKSELNRNNSLPLIDTAQAVD IHPVLHEEDQENTNKRKNSLLSNLK RKDLGESKISIRKDYSHFDRIPSPSS ARSVGETDFNYNREPS EDTLRYPD SI IEVTNRTSPAPNSILSRSGQFVNSND LSDGFSTNTINNVLNANЕКIFFN ALQSMЕKENLMLWKT SQKYGT YLD ERRKSADAFQKREGSCVDIGKLSH SHLPFINLPPWPTELTEEERIIHDRL ASKHSHHIRKHVHNARNKTSCKIKD SVGTF LGMTNSLTNKATVKKRTGQI LKKEKMLVMVKEAIAQNKVPLPNFSE NECFDTRVSRWKEYVIARSTGRFD PPILLQFYRHRHIPEIEDISSIATKYH RNPLDFFLSRNCIVKFYSLLDKTISIQ KPKRLGGFIDESIEKKDELKHYSPI KIFILRCSSIRSSGRWYKFLES LDR QLFTPAINLKIPLTEISIKINLNEIIFQ KLIDLGKQEKDRLKICFLQRGYKIFQ HPILRYFTVAILEKLLAHYDYLIRKW DTENPVLGCALKRYDRLEWIPCED SLVTGIFAFQCQSHLIQYRPIANRLRET KSLEGKCLKEPTPIEGFLIRLTDKYGS ARTNFGKYSISTAYFFTCENLLFSMK AYRANPPLPIDS MIDD TSTEIEKEEI WKQWKKIPEVYEQOPYPLDTNDHIE WMNCQTTQSEYDSRDFYAFHCFHR RIDQILKTDNVIDL TEVKDIYQGTRTD YEADKIKYGVYKEASEIFWHRNYEID DVSRSVINIETSNGLLLKLLATSATV AEEWVVKLQMSIYWK NKQREDTER LLKIRRSNAGLLMLNGEETKIGEN TLRWIVEHGRADEQTFNANGISLSR PLIQKGPLYQPKHKS VFSKYVVVLI SGFIVLFHCFHRSTTGFAKEVLEYAH YVTIPIDDCYLYSGTTTEL DLLORDRT FDEINYGSHALPRVYGDGWR SVEDE SSRCFTLWFGTRRALSSNRLQKKGN EKQYTQDYGRQDNNIDPPSAPEADL NNSNVPSNTDKIHFTK KLGVS GKSM VFMARSRQERDLWVMSIYYELERLR RTASTSNSRNQTM	None	None	None	None	None	None	None		