

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
P39535	PHO90_YEAST	Saccharomyces cerevisiae	Low-affinity phosphate transporter PHO90	9.365641	NaN	NaN	33229814	MRFSHFLKYNAVPEWQNHYMDYSE LKNLIYTLQTDELQVGDNEEGFGAG KSSNITDRFKNKFSFKNAKEDTSSG MNKDAGIVEETIELRELPTAQTVAAK PSPFRRMKEKIFYKRRSSASSVSST ANENLQLDITYDTFVGDLTAEKQKVD DFYKRTEAKFYDKFDALVKDLKKIGV IEYDIDDDTLFNEPIASTNDEVPLD LDDDEDDEFFYDDQSNIEDNTALLH HSQYNIKSQKKSLLKKSIVNLYIDLC QLKSFIELNRIGFAKITKKSVDKVLHL NTRTELIESEQFFKDTYAFQAETIEL LNSKISQLVTFYARITDRPHNISHSK QELKSYLHDHIVWERSNTWKDMLG LLSQADELTPKETEYNANKLVGKLD LEYRWRPLPRPINLKFTSINNVALPK LFFTKKAYKIYFIILVTGLLGIKTFND AAQHRCMALVECVAFLWASEAIPHLH ITAFVLVPLLVVLFKVLKTS DGAIMSA ASASSEILAAMWSSTIMILLAGFTLG EVLAQYNIKVLASWLLAFAGCKPR NVLLMAMCVVFFLSMWISNVAAPV LTYSLLSPLLDAMDADSPFAQALVLG VALAANIGGMSSPISPPQNIISMSYL KPYGIGWGQFFAVALPSGILAMLLV WILLFTTFKMNTKLEKFKPIKTKFT VKQYYIITVTVATILLWCVESQIEGAF GSSGQIAIPIVLFFGTGLLSTQDLNA FPWSIVILAMGGIALGKAVSSSGLLS TIAKALQKKIENDGVFAILCIFGILML VVGTFVSHTVSAIIIIPLVQEVGDKLG NPKAAPILVFGCALLSSCGMGLASS GFPNVTASKVDRKGDRYLSVMTFL TRGVPASILAFLCVITLGYGIMASVVK GNATSA