

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
P40091	PEA2_YEAST	Saccharomyces cerevisiae	Protein PEA2	24.475439	NaN	S230	33229814	MHKFDLELSRRANPLLFSAERYEY PLKYDELKQYLLSQNPSPHNSRP YTSIDYFDYLLYRSKNDESEIDLDDK LVSEFALYYVQKEHLNSDDLNP ELLLKLPKSADWYEMMLRILESINT TGIDQLTKENNSFPNSKRARSSTN MGGTDKFNGAYHTDKADDDKNEI LQELTSFLMSNSIQKIDIKPIDDP VKFLKNGINSILDTCVSLEKNTSSPI SPNAAAIQEEDSSKKLELETAFSDL QLAHNFLTQFENDRAEYVQDIEKL TRTNRELQDKLLNNHNSLSKTEKKL HELEQENKELEKANNKLNSSRHNF GMSSPASSPVTWDPSSPSVGSPTS GSGSRSLSIMTSEFKKVLSTQRKYE KELSDEREHRFKLERELALLKNAEA NTSLALNRDDPPDML	None	None	None	None	None	None	None	None	None

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