

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	ext reg							
P38970	HAL5_YEAST	Saccharomyces cerevisiae	Serine/threonine-protein kinase HAL5	24.475439	NaN	S17;S19;S68;S72;S160;S273;S277;S324;S333;S336;S358;S391;S395	33229814	MGDEKLSRHTSLKRARSLSESIGL FKPSGISGSNNAAPSSRPGQDQAH SHQTAIRIITSNVSSPISIPVHSPVLQA APKHHKLGVPNIAKLSLSPSREPSLN SENEMFSQESFISEKDEDEANLLER EDLQNKKEEKARAKHVRSKEAYVPH HRYTVGSDEVERQPRERLKNFPONA GSSNPANSNANHVLVDQENNFSIDA MLDYDEESKLRRRNSLGVNRHNSNR TRSRKNSLSTPRSPPMKNGNGGMN SNATNNVNGGTGNRIYMRGRNHSD SISASSLPKFQEIIECKCILDGLGHFKVF ENGYHEHSLRVLPIITNNKNVDSGD EKDADASVNSGDDGDNDSEANMH KOKSVFSLSGFLFKSHKDGNOQQQQ QQQEEENGEQINLEKAFSIIQSQRFI KSQTLKKSRTSNLKNNGNDELKMN DGKNIPQIVNPNAAVGVVEELKLINAL SEKIRKGLKSENTKGNNGEGRSNSN KQEDSDDTEGKAGTTNDDTSHKPC SOKYGKSGVVGAGAYGVVKICARCK TAKDVLPYSTYSNGKCLFFAVKELKP KPGDQIDKFCTRLTSEFIIGHSLSHP HFEANAMIAGNVSRITPPKHVFNAP NILKILDLEYSNSFVEVMEFCASG DLYSLLTRNNISNESNNGSSRLIQTV KEGSGSPLHLEADCFMKQLLNGV QYMHDHGIAHCDLKPENILFQPNGL LKICDFGTSSVFQTAWEKHVHFQSG AMGSEPYVAPEEFIRDAEYDPRLVD CWSCGIVYCTMVMGQYLWKIAIPEK DSLFKSFLSEIKDDGQFYLFEELRHV SSELNRLRKIALYRTFQVDPKTRITIE QLLQSSWMRKTCCVVYRHLHTKV SK	None	None	None	None	None	None	None	None	None	None	None	None	None	None	None	None

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