	UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochndrion	endoplasmic reticulum	golgi apparatus	plasma membrane	ext reg
0	Q03780	YD239_YEAST	Saccharomyces cerevisiae	Uncharacterized protein YDR239C	24.475439	NaN	S63;S254;S313;S342;S345;S390;S477;S492;S546;S683;S699	33229814	MFDGFSNNKGKRRSFFRFGSESKN NDSEKSVRKPSVPSTIKKSTNIARTS TAETSAPDIPPRSPNRNAHSRSHSIQ APLQKETLKNTNPFLNAEDTLGDSL ELTQSKEASGNDHKGIEYLQENNIIG QRTNPFTTSANSNAHFSKIKRSRPPP PPMDMKSITTSISNNTTKEEIESNN DSERDSIAISSTHNQHRRQRSEAEK LVDDIENYINEHKVSSGSSLSLDTSE NSDTKASQDKLPVDVMEAPILRNVS AESSLSYVKPLIVDNEEVNKASNGN LVQSDHLKEFSSNLDDGDDKFSFST SASGKSTKSLQQVSKDESSGFKAAH FDFAYKSNEHLGSDGSIASARKPLRI TNEIDSGSSNEDDDDGLQEKGFVDS ESKAFINYASDQGSSIKNDVSTQEPE LPSHRRIFRVVNEDRPSFYLNSVND TGSLTDKHSFDTASSGEYDAKSNFS SQSGLSISKGSKSTVLAALDSNGNTK SSNKTSELNSLNSISESLVPAAHSFN EHTVTIPATVDLPNPVHDAPSERSVK CSPLTSVVSNKSEKSVPLVSSYVEEL RLKYYKTSNFLQAPPNLPVALKQKN NLIQPKNIKVKLRTSSKQIGIKHGKV KQKLLALETRNEESDGTATGLKNKI NVDHTKEFHKLLGKENETGSISKKE GTDAEQAEDYLKDIPGDEAYNSDDI MAPLREKRGQNGSVDSVSRSNTVV SYYTRSQNRMRSGTLDNDYVNRQK LPTHISLQDYRDANARSNISRQDSVS TTNSDVVDLSYSLGHGLRVANPDSD PE	None	None	None	None	None	None	None	None	No