	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	extracellular region
0	P53278	YG3A_YEAST	Saccharomyces cerevisiae	Uncharacterized protein YGR130C	24.475439	NaN	S286;S343;S347;T809	33229814	MLFNINRQEDDPFTQLINQSSANTQ NQQAHQQESPYQFLQKVVSNEPKG KEEWVSPFRQDALANRQNNRAYGE DAKNRKFPTVSATSAYSKQQPKDLG YKNIPKNAKRAKDIRFPTYLTQNEER QYQLLTELELKEKHLKYLKKCQKITD LTKDEKDDTDTTTSSSTSTSSSSSSS SSSSSSSSSDEGDVTSTTTSEATEAT ADTATTTTTTTSTSTTSTSTTNAVEN SADEATSVEEEHEDKVSESTSIGKGT ADSAQINVAEPISSENGVLEPRTTDQ SGGSKSGVVPTDEQKEEKSDVKKVN PPSGEEKKEVEAEGDAEEETEQSSA EESAERTSTPETSEPESEEDESPIDP SKAPKVPFQEPSRKERTGIFALWKSP TSSSTQKSKTAAPSNPVATPENPELI VKTKEHGYLSKAVYDKINYDEKIHQA WLADLRAKEKDKYDAKNKEYKEKL QDLQNQIDEIENSMKAMREETSEKI EVSKNRLVKKIIDVNAEHNNKKLMI LKDTENMKNQKLQEKNEVLDKQTN VKSEIDDLNNEKTNVQKEFNDWTT NLSNLSQQLDAQIFKINQINLKQGK VQNEIDNLEKKKEDLVTQTEENKKL HEKNVQVLESVENKEYLPQINDIDN QISSLLNEVTIIKQENANEKTQLSAIT KRLEDERRAHEEQLKLEAEERKRKE ENLLEKQRQELEEQAHQAQLDHEQ QITQVKQTYNDQLTELQDKLATEEK ELEAVKRERTRLQAEKAIEEQTRQK NADEALKQEILSRQHKQAEGIHAAE NHKIPNDRSQKNTSVLPKDDSLYEY HTEEDVMYA	None	None	None	None	None	None	None	None	None