

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
Q06836	SYT1_YEAST	Saccharomyces cerevisiae	Arf guanine nucleotide exchange factor SYT1	11.278152	NaN	T277;S369	33229814	MNQSISSLIKLFQSHSNDKNGNK KGGSNVSTGIDKLRESESYRSPFLQL AEIQEHTNDDDKLDVKECEPTKKH SKLSRIRRKMGRLDLNFRSANEEKGS EDDEILVAQHLRNGQDPEEMPFKSE NNIDSIEKVPKPDGERVTLTSSGSDN VKRNSKHAPFIPVKPALEKFPSSNRL SRDYRKSQEPTLFNGDRLVPTLPTVS RISTSSSVGSSTAASRYFNPSKRAVV ASSSSSSSIKFNLSLHAIPDATPQIE LAKQQDEISKRRFGRRRSRTVDVFD YINKNNTAKNKPPLSPSSFIRTIDEK NTNSLMQDPMGSRGPLLPDDANIIS NTDGAEEASHPDHQVLSRSRSQSTS FVQGGKGRKSIEDEGYHNKLGPH GSGPTSVYNNKSNANSTITGMSRRS SSIVNALSSFVNLRSSLSRSSRQHL QQQQQLQKLDVSLDLPPVPAPPEF SDSCKDFLIKLPYKFIGIILTEKDD EFKKNCLNYLLTNCFEFKNPLDIA LRKLLMFLELPKETQIDRLIMAFSF AYYKAQKSYSKKKGIECPWSNADQV YFIAFSLMLHTDYFNPNNKSKMTK HDFVDLVHNDKYSGGNEIPMAVLTY FYENVTAKESPKFNYFLMSPMALDD SILDKDAFDTNFAITLSSNSMYSPID MIKRGSIIPKEASLPIFYPLTNSISAS GIAPSTAASCPPSTSGTINGANLGT NSNSNRPASNSISSYFSYNPSSSSSG NATLVQDDINVYSHIINDTLNEVNL PEVSKYWNKNALKANLLRNEEHKY EKYYSIMNDTKGGYLRFHKSQNLKL NLPNFEILNDNSRSGCKNSDYKYCK ILQMGAIMNLGMPSRKFSIVNSAKI HWKKEFAILTSLGLLICDKMDWINP QMMKDPKSGTTNYIIDFKSGFSFVP GSTIDVYNGLFADRERDSLKSHFA SLVLAYTEHHSTGSHTSNNTAASSA KHNEGVFEPSSDEEDSITNSTDGTS SVSNGESDNDVSSSDNQLSSNDS NEDYHSIKDEYPIFEDENADCLLYLH TCHRNFIWKANKYERDNWIDSINL FSAYDGCYVEIGSIANTICNKRKLTL QRMERLRSIKSAKWEKLLKFFESTLM LMGKCVPISTKTKTDMINRIRQLAVR MDWLIYEIKRSELFVSIIEVTRKQA EKNILEHGKGEEEGQGNNDDSDGI DDIEESFLFNENSLQVCVSDSSYDE YSNE

