

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
Q03088	SVL3_YEAST	Saccharomyces cerevisiae	Styryl dye vacuolar localization protein 3	9.365641	NaN	S471;T496;S551;S662;T739;T756;S757	33229814	MSSSSLRLVLAIGNNPILFYTSRFQL AKNIDLYHVNDKSKCQFEIETEYYGK DRFELENHFTSIEHLTEALSSKSSEA VFDIIIMSAPSLQELSSLASKLTSIIDS NTKIFLESSGFIQLEPFVKLSMESPH VNVFSILTDLDIRQIGPNHFKHFPST AKENTIYLGESKSSTEKYSSGVITLLT TFEKLFAKLFSNIKINLCNFSSIEFLS QQWKLAISSRICFDPLLIMFEQENPSD LDQQIIAKPLISGLVTEIITVAKTMGA RLNSSHDNENSLLSLWKNYSYHSTN KPPALVYHFHQTTPNLIDILLQITL LADDFGIKTPYLEFLYSVLSQFERLN SGKSKWFIRSDEKTQILQSLQKSQK NESALQTQITSLQGQISKLRQELLM QAKQHEMETNELKEKHQVALKAQA QAQAQAQSQAQTSIEALTPTEATNQ SDTNEYKATGTPNLRDIEDMALYSV NYGDSPPVRSPPPVVSSQPQMNSPLS SHSQTFGENNGTNDKLLQERELQL RKKELELQERELEFQKRALQQQRFN NSNNSIPRKPSFPQLQOSANVRSNS RGMHGTNGAMSQPASAGNFVDPIS SSIAAYDPQQPPSLPLQPPQQSVQVQ PFHSHSIKPTSRKNRNSNMPNIGNP SSINMSDFGRPPNNSQTRLNSMPT HSIVNQNRQRSQSKNKLNMPHAT NPNNTFNQVPAPSLNNHVPTQRQF SSSTMIEVTNNNNKVNNSSSNPDIS TNSVVHNAMQFTNTNNTSSTVDI NDPKNIAPPPTTSVSAPSTPTLSSSS QMANMASPSTDNGDNEEKNGGKK KRFGLFKKKKNSKK