

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
P53935	NST1_YEAST	Saccharomyces cerevisiae	Stress response protein NST1	9.365641	NaN	S266	33229814	MPPNSKSKRRKNKSKQHNNKNGNS DPEQSINPTQLVPRMEPELYHTESD YPTSRVIKRAPNGDVIVEPINTDDDK KERTANLTHNKDSMDSASSLAFTLD SHWESLSPEEKKTILRIEKEEVFVI RNYQDDHSCSCSVCGRRLAMDQE MERIYNTLYAMDKDKDPETNPIKFH LGIKELQISKNNQQNDLSSTKGEVV KNFLSSSTVGSLEEVLHFKQKQLS KQEQAHNETADNTSLLEENLNNIHI NKTSSSEISANFNVSVEELOQKYSN FTKTFISSHPKIAEEYVQKMMMYPNI RALTDDLMSNGQGFLNAIEDFVR DGQIQASKKDDSITEDEASSTDLDTP KEFTTMLHSGKPLTEDEYADLQRNI AERMTNAYDTASKKFKDVSQLEKEL FTRFMGRDKKSFRELIIQSFKNKFD GELGPSVLAATLSSCFSSQSKDTSLD TDSIYEDEDEEDYDDYSEYAEDSEEV SEYEGIEAVEKPEHDEKSNNGIRETLH LSYDHDHQRQNHPPHHYHSTSTHS EDELSEEEYISDIELPHDPHKHFHR DDDILDGDEDEPEEEDENEGDDEE DTYDGLDETDRLEEGRKLIIAITKL LQSRIMASYHEKQADNNRLKLLQEL EEEKRKKREKEEKQKKREKEKEK RLQQLAKEEEKRKEEEKERLKKEL EEREMRRREAQRKKVEEAKRKKDE ERKRRLEEQRREEMQEKQRKQKE ELKRKREEEKRIREQKRLEQEKLO KEKEEEERQLIAEDALRKQKLNEE QTSANILSAKPFTENGVGPNVSSQS HPNMTNYQEDNSCSINDEILKMVN SVAASKPVSPTGFNVHDL LLPSTNN QMPAMEQSHLPQPGNQNNHFGTT TIPNALDLATKSSLQ TENNYLMNSQ TLENTSLLMHNNSSPTKLLPNDFGL SSWGGTLNTMSINPTCKPPVIQTSE MESQAHKSSPQATMPSFGLPNGGT HRKSFTDELNTLTSMLSSSGFADTS LSSSGFPPSQRSVWNDQKSSFSGPS TAGNFNNSIQSGMLLAPTLGSVES FPNRTSIWDSSTTPMMNKSELGR NITSTAQDSPA FMASNIWSSNSQYN SPYLTSNVLQSPQISSGVDESHILDSI YNTYLAISPQDSLNPYIAIGTLFQNLV GLNLDYSTFINKLISMQ GAYNCEFFT DNNGSITHVRFARQTPAGHSGLLN QLFSGLNDPTATPFTSRPHTSTRASF

