	UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochndric
0	Q14687	GSE1_HUMAN	Homo sapiens	Genetic suppressor element 1	12.720506	T46;S87;S116;T125;T129;T179;S212;S589;S917	S10;S84;S95;T433;S766;S826;S828;S857;T907;S909;S1101	38665916;34019948;33214551;30379171;31492838;39531497;40307207;35132862;35138101;32574038;39534244;35254053	MKGMSHEPKSPSLGMLSTATRTTAT VNPLTPSPLNGALVPSGSPATSSALS AQAAPSSSFAAALRKLAKQAEEPRG SSLSSESSPVSSPATNHSSPASTPKR VPMGPIIVPPGGHSVPSTPPVVTIAPT KTVNGVWRSESRQDAGSRSSSGGR ERLIVEPPLPQEKAGGPAIPSHLLSTP YPFGLSPSSVVQDSRFPPLNLQRPV HHVVPPSTVTEDYLRSFRPYHTTDD LRMSSLPPLGLDPATAAAYYHPSYLA PHPFPHPAFRMDDSYCLSALRSPFY PIPTPGSLPPLHPSAMHLHLSGVRYP PELSHSSLAALHSERMSGLSAERLQ MDEELRREREREREREREADRE REKEREREKEREQEKEREREKER ERELERQREQRAREKELLAAKALEP SFLPVAELHGLRGHATEERGKPSEQ LTPTRAEKLKDAGLQAPKPVQHPLH PVPTPHHTVPSLISNHGIFSLPSSSA ATALLIQRTNEEEKWLARQRRLRQE KEDRQSQVSEFRQQVLEQHLDMGR PPVPAEAEHRPESTTRPGPNRHEPG GRDPPQHFGGPPPLISPKPQLHAAPT ALWNPVSLMDNTLETRRAESHSLH SHPAAFEPSRQAAVPLVKVERVFCP EKAEEGPRKREPAPLDKYQPPPPPPR EGGSLEHQPFLPGPGPFLAELEKST QTILGQQRASLPQAATFGELSGPLKP GSPYRPPVPRAPDPAYIYDEFLQQRR RLVSKLDLEERRRREAQEKGYYYDL DDSYDESDEEEVRAHLRCVAEQPPL KLDTSSEKLEFLQLFGLTTQQQKEE LVAQKRRKRRRMLRERSPSPPTIQS KRQTPSPRLALSTRYSPDEMNNSPN FEEKKKFLTIFNLTHISAEKRKDKER LVEMLRAMKQKALSAAVADSLTNSP RDSPAVSLSEPATQQASLDVEKPVG VAASLSDIPKAAEPGKLEQVRPQELS RVQELAPASGEKARLSEAPGGKKSL SMLHYIRGAAPKDIPVPLSHSTNGKS KPWEPFVAEEFAHQFHESVLQSTQK ALQKHKGSVAVLSAEQNHKVDTSV HYNIPELQSSSRAPPPQHNGQQEPP TARKGPPTQELDRDSEEEEEEDDED GEDEEEVPKRKWQGIEAVFEAYQEH IEEQNLERQVLQTQCRRLEARHYSL SLTAEQLSHSVAELRSQKQKMVSER ERLQAELDHLRKCLALPAMHWPRG YLKGYPR	False	False	1.266	3.053	1.924