

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
Q62383	SPT6H_MOUSE	Mus musculus	Transcription elongation factor SPT6	15.818779	S1526;T1532	S7;S12;S73;S78;S91;S125;S267;T1515;T1523;S1526;T1532;S1535;T1539;T1697;S1701;S1703;T1709;T1718	36852467	MSDFVESEAESEEEEEYNHEGEVVP VTKKFVEEEDDDDEEEEEENLDDQD ERGNLKDFFINDDDDEEEGEEDEGS DSGDEDDVGHKKRKRPSFDDRLE DDDFDLIEENLGVKVVRGQKYRRVK KMSDDDEDDEEYKKEEHEKEAIA GEIFQDEEGEGQEAVEAPMAPPDE EEEDDEESDIDDFIVDDDGQPLKKP KWRKKLPGYTDAALQEAQEIFGVDF DYDEFEKYNEYDEELEEDYEYEDDE TEGEIRVRPKTTTKRVSRRSIFEMY EPSELESSHLTDQDNEIRATDLPERF QLRSIPVKAEDDELEEEADWIYRN AFATPTISLQDSCDYLDRGQPTSSFS RKGPTVQKIKEALGFMRNQHFVFP FIAFYRKEYVEPELHINDLWRVWQW DEKWTQLRIRKENLTRLFEKMQAYQ YEQISADPKPLADGIRALDTTDMER LKDVQSMDELKDVYNHFLLYGRDI PKMQNAAKASRKKLKRIKEDGDEEG EGEEAEDEEQRGPELKQASRRDMY TICQSAGLDGLAKKFGLTPEQFGEN LRDSYQRHETEQFPAEPELELAKDYV CSQFPTPEAVLEGARYMVALQIAREP LVRQVLRQTFQERAKLNITPTKKGR KDVDEAHYAYSFKYLKNKPKVRELDRD DQFLKIGLAEDEGLLTIDISIDMKGV EGYGNQTYFEEIKQFYRDEFHQ VQEWNRQRTMAIERALQFLYVQM AKELKNKLLAEARESVMKACSRKLY NWLRVAPYRPDQQVEEDDDFMDEN QGKGIRVLGIAFSSARDHPVFCALVN GEGEVTDFLRLPHFTKRRTAWREEE REKKAQDIETLKKFLVNKKPHVVTIA GENRDAQMLTEDVKRIVHELDQGG QLSSIGVELVDNELAILYMNSKKSEA EFRDYPPVLRQAVSLARRIQDPLIEF AQVCSSEEDILCLKFHPLQEHVVKE ELLNALYCEFINRVNEVGVDVNRAIA HPYSQALIQYVCGLGPRKGTHLLKIL KQNNTRLESRTQLVTMCHMGPKVF MNCAGFLKIDTASLGDSTDSYIEVLD GSRVHPETYEWARKMAVDALEYDE SAEDANPAGALEEILENPERLKDLDL DAFAEELERQGYGDKHITLYDIRAEL SCRYKDLRTAYRSPNTEEIFNMLTK ETPETFYIGKLIICNVTGIAHRRPQGE SYDQAIRNDETGLWQCPFCQQDNF PELSEVWNHFDGSCPGQAIGVKTR

LDNGVTGFIPTKFLSDKVVKRPEERV
KVGMTVHCRIMKIDIEKFSADLTCRT
SDLMDRNNEWKLPKDTYYDFDAEA
ADHKQEEDMKRQQRTTYIKRVIAH
PSFHNINFKQAEKMMETMDQGDVII
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VREEGKENAFSLGATLWINSEEFED
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WFKDHYQDPVPGITPSSSNRTRTPA
SINATPANINLADLTRAVNALPQNM
TSQMFSIAAAVTGQGQNPATPAQ
WASSQYGYGGSGGSSAYHVFPTPA
QQPVATPLMTPSYSYTTSPQITTPQ
YHQLOASTTPQSTQAQPQPSSSRQ
RQQPKSNSHAAIDWGKMAEQWL
QEKEAERRKQKQRLTPRPSPPMIE
STPMSIAGDATPLLEMDR