

UniprotKB ID	Entry name	organism	full name	oglnacscore	oglnac sites	phosphorylation sites	PMIDS	sequence
Q9EST3	4ET_MOUSE	Mus musculus	Eukaryotic translation initiation factor 4E transporter	28.163225	S416;S576;S911;S919;T922	S4;S73;S77;S114;S119;S135;S137;S300;S344;S352;S373;S416;S563;S586;S692;S751;S919;S949	22517741;36757814;36852467	MEKSVAETENGDAFLELKKLPTS SKSPHRYTKEELLDIKERPYSKQRP SCLSKYDSDGVWDPEKWHASLYP ASGRSSPVESLKKESESDRPSL VRRRIADPRERVKEDDLDVVLS PQRRSFGGGCHVTAAVSSRRSG SPLEKDSGLRLLGGRRIGSGRII SARAFKDHRLSDKDLRLRDRDR RERDYKDKRFRREFGDSKR VFGERRRNDSTYEEEEPEWFS AGPTSQSETIELTGFDDKILEE DHKGRKRTRRRTASVKEGIVEC NNGGVAEEDEVEVILAQEPSAD QEVPRDVILPEQSPGEFD FNEFFNLDKVPCLASMIEDV LGEVSASARFSRWFSNPSRSG SRSSLSGTPHEELERLAGLEQ AVLSPGQNSGNYFAPISEDH AENKVDILEMLQKAKVDL KPLLSSLSANKEKLKESHSG VVLVVEEVEAGLKGLKVDQ QMKNSTPFMAEHLEETLSA ASSNRQLKKGDMTAFNKL VNTMKASGTLPTQPKVSRN VESHLLAPAEIPGPVSKN ILQELLSQPVPQRPASSN LLSGLMGSLEATASLS QRAPSPPMQVFRFTAAS ADYLHPRIPSPIGFPSG PQQLGDPFQGMRK PMSPVSAQMSQLELQQA ALEGLALPHDLAVQTAP FYQPGFSKPQVDRTRD GLRNRQQRMSKSPAPM HGGNSSSPPAPAASIT SMLSPSFTPTSVIRK MYESREKTKEEMAPGM VVPGDGKEDTQK TSEENLLSSNP IPNTDQDSSTTNP KLSTLQRSSCSTPL SQTSRYTKEQDYRP KTAGRKTPTL ASPVPGT PFLRPTHQV PLVPHVPIVR PAHQLHPGL VQRLIAQ GVHPQHLP SLLQAGVLP PGIDMAPL QGLSGPL LQPLYPLV SAASHPL LNPRPGT PLHLAVM QQQLQRS VLHPPGS SSQAAAIS VQTPQNV PSRSGM PHMHSQLEH RTSQRSS SPVGLAK WFGSD VLQQPL PSMPTK VISVDE LEYRQ

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