

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum
Q9NRA8	4ET_HUMAN	Homo sapiens	Eukaryotic translation initiation factor 4E transporter	23.218727	S331;S772;S905;S914	S5;S74;S78;S115;S120;S136;S138;S301;S345;S353;S374;S417;S513;S564;S587;S693;S752;S920;S951	38253038;38665916;31492838;39534244;39302247	MDRRSMGETESGDAFLDLKPPASKCPHRYTKSELLDIKELPHSKQRPSC LSEKYDSDGVWDPEKWHASLYPAS GRSSPVESLKKELDTDRPSLVRRIVD PRERVKEDDLDVVLSPPRRSFGGGC HVTAAVSSRRSGSPLEKSDGLRLL GGRRIGSGRIISARTFEKDHRLSDKD LRDLRDRDRERDFKDKRFRREFGDS KRVFGERRRNDSTEEPEWFSAGP TSQSETIELTGFDKILEEDHKGRKR TRRRRTASVKEGIVECNGGVAEEDEV EVILAQEPAADQEVPRDAVLPEQSPG DFDFNEFFNLDKVPCLASMIEDVLG EGSVSASRFSRWFSNPSRSGSRSSS LGSTPHEELERLAGLEQAILSPGQNS GNYFAPIPLEDHAENKVDILEMLQK AKVDLKPLSSLSANKEKLESSH GVVLSVEEVEAGLKGLKVDQQVKNS TPFMAEHLEETLSAVTNNRQLKKG DMTAFNKLVSMTKASGTLPSQPKVS RNLESHLMSPAEIPGQVPKNIQL LGQPVQRPASSNLLSGLMGSLEPTT SLLGQRAPSPPLSQVFQTRAASADYL RPRIPSPIGFTPGPQQLLGDPPFGMR KPMSPITAQMSQLELQQAALGLAL PHDLAVQAANFYQPGFGKQVDRTR DGFRNRQQRVTKSPAPVHRGNSSSP APAASITSMSPSFTPTSVIRKMYES KEKSEEPASGKAALGDSKEDTQKA SEENLSSSSVPSADRDSPTTNSKL SALQRSSCSTPLSQANRYTKEQDYR PKATGRKTPTLASPVPTTFLRPVHQ VPLVPHVPMVRPAHQLHPGLVORM LAQGVHPQHLPSSLQTVLPPGMDL SHLQGISGPILGQPFYPLAASHPLL NPRPGTPLHLAMVQQQLORSVLHP PGSGSHAAAVSVQTTQNVPSRSGL PHMHSQLEHRPSQRSSSPVGLAKW FGSDVLQQPLPSMPAKVISVDELEYR Q	True	False	3.481	5.0	1.379	1.208