

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
Q05D44	IF2P_MOUSE	Mus musculus	Eukaryotic translation initiation factor 5B	13.393436	T966	S66;T107;S108;S114;S137;S139;S145;S165;S172;S183;S184;S187;S191;S209;S215;S223;T300;S437;T497;S544;S553;S556;S584;S585;S587;S591;S1164	34418053;34887587	MGKKQKNKSEDSTKDDTDLGALAA EIEGAGAAKEQEPQKSKGKKKKEKK KQDFDENDILRELEELSLEAQGIRAD RDAAAVKPTENNEEESASKQDKKKK GQKGGKTSFDENDSELEDKDKSKSK KTARPNSEAPLSGSEDADDSNKLKSK KGKKAQKSTKKRDGSEEDEDNSKRS KERSRVNSSGESGGESDEFLOSRKG QKKNQKNKSVPTVDSGNEDDDSSF KIKTVAQKKAEEKKEREKKKRDEEKA KLRKMKEKEELEKGGKKEQSKQREP QKRPEEEVLTLRGTPDTGAASEEKG DTAAALEDDNEGDKKKKDKKKKKT EKDEKEKEKKGPSKSTVKAIQEALA KLKEEERQKREERIKRLEELEAK RKEEERLEQEKREKQKEKERKER LKKEGKLLTKSQREARARAEVTLRH LQAQGVVPSKDSLPPKRPVYEDKK KKKTPQQLSKEVSETLEISAPVEAV DQGGPEKEETPPSVEPEEEEDTEDA GLDDWEAMASDEEREKEGNMIHIE VEENPEEEEEEEEEEEEESEDEEE EGDSEGS DGEEDCKLSDEKDSGK AGDTKPSKDASSDSEYDSDDDRTKE ERAYDKARRIEKRRLEHGKNVNT KLRAPIICVLGHVDTGKTKILDKLRH THVQDGEAGGITQIGATNVPLEAIN EQTKMIKNFDRENVRIPGMLIIDTPG HESFSNLRNRGSSLCDIAILVVDIMH GLEPQTIESINILKSKKCPFIVALNKI DRLYDWKKSPSDVAVTLKKQKNT KDEFEEERAKAIIVEFAQQGLNAALFY ENKDPRTFVSLVPTSAHTGDGMGSL IYLLVELTQTMLSKRLAHCEELRAQV MEVKALPGMGTTIDVILINGRLKEG DTIIVPGVEGPIVTQIRGLLLPPMKE LRVKNQYEKHKEVEAAQGVKILGKD LEKTLAQLPLLVAYKDDEIPVLKDELI HELKQTLNAIKLEEKGVYVQASTLGS LEALLEFLKTSEVPYAGINIGPVHKK DVMKASVMLEHDPQYAVILAFDVRI ERDAQEMADSLGVRIFSAEIIYHLFD AFTKYRQDYKKQKQEEFKHIAVFC KMKILPQYIFNSRDPVIGVTVEAGQ VKQGTPMCPVPSKNFVDIGIVTSIEIN HKQVDVAKKGQEVCKIEPIGESPK MFGRHFEATDILVSKISRQSIDALKD WFRDEMOKSDWQLLIVELKKVFEII