

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
Q80XI3	IF4G3_MOUSE	Mus musculus	Eukaryotic translation initiation factor 4 gamma 3	9.983052	NaN	T168;S230;S232;S267;S436;S470;S472;S490;S1150;S1212	35822049	MNSQPQARSPFFQRPQIQPPRAAIP NSSPSIRPGVQTPTAVYQANQHIMM VNHLPMYPYVPTQGHQYCIQYRHSG PPYVGGPPQQYYPVQPPGPGPFYPGGP GDFANAYGTPFYPSQPVYQSAPIIVPT QQQPPPAKREKKTIRIRDPNQGKDI TEEIMSGGSRNPTPIGRPASTPTP PQQLPSQVPEHSPVVYGTVESAHLA ASTPVTAASDQKQEEKPKPDPVFQS PSTVLRRLVLSGEKKEQAGQMPETAA GEPTPEPRTSSPTSLPPLARSSLPSP MSAALSSQPLFTAEDKCELPSSKEE DAPPVPSPTSCTAASGPSLTDNSDIC KKPCSVAPHDSQLISSTILINEMNGV GEKLSAKENTVGMLRQEVLPPLTELE EILEHPQEELKVECTPTPIAPSMPLA FSPAPPTPPTSPPCPPVLSAAIARSP AVATEVQRVADEGESLRTCLSKDAK EMQDKAESESDGQAEETADPQSLH SGRSPAPVQTATTAPKSWKKTKEQT RTPDEVLEAEAEPKAEELAVDSVLE PEQEKMSQGFPSERDPSALKRGKAE EGNGEEAEPVRNGAESASEGEGGD GNSGSADSSADGLTFFPKAESWKPA DTEGKKQYDREFLLDIQFMPACIQK PEGLPPISDVVLDKINQPRLSMRTL PRILPRGPDFTPAFADFPRTQTPGGRG VPLLNVGPRRSQPGQRREPRKIITVS VKEDVHLRKAENAWKPSQKRDSHA DDPESIKTQELFRKVRILNKLTPQM FNQLMKQVSALTVDTEERLKGVIDL VFEKAIDEPSFSVAYANMCRCLVTLK VPMADKPGNTVNFRKLLLNRCQKE FEKDKADDDVFEKKQKELEAASAPE ERTRLHDELEEAKDKARRRSIGNIKF IGELFKLMLTEAMHDCVVKLLKN HDEESLECLCRLTTIGKDLDFEKAK PRMDQYFNQMEKIVKERKTSSRIRF MLQDVIDLRLCNWVSRADQGPKTI EQIHKEAKIEEQEEQRKVQQLMTKE KRRPGVQRVDEGGWNTVQGAKNSR VLDPSKFLKITKPTIDEKIQLVPKAQL GSWKGSSGGAKASESDALRSSASS LNRFSPLQPPAPSGSPSATPLEFDSR RALTSRGSMDGREKSDKPIPAGTARP NTFLRGSSKDLLDNQSQEEQRREM LETVKQLTGGLDAERASTEADRSKT RELAKSEMCAVPAPDKPALSEEEVE RKSksiIDEFLHINDFKEATQCIEELS

