

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
O43432	IF4G3_HUMAN	Homo sapiens	Eukaryotic translation initiation factor 4 gamma 3	20.917404	T168;T211;S232;S258;S277;T278;T283;S284;S285;S288;S289;T294;T295;S736;S806;S1050	T168;S230;S232;S267;S441;S495;S1156;S1218;S1409	33214551;28657654;22661428;31637018;35254053;30379171;38253038;29351928;30620550;31492838;32119511;35138101;32870666;34019948;23301498;37340703;35289036	MNSQPQTRSPFFQRPQIQPPRATIP NSSPSIRPGAQPTAVYQANQHIMM VNHLPMYPYVPVQGPQYCIQYRHS PPYVGGPPQYVPPGPGPFYVPGPGP GDFPNAYGTFYFYSQPVYQSAPIIVPT QQQPPAKREKKTIRIRDPNQGGKDI TEEIMSGGSRNPTPIGRPTSTPTP PQQLPSQVPEHSPVVYGTVESAHLA ASTPVTAASDQKQEEKPKDPVLKSP SPVLRVLVLSGEGKEQEGQTSSETTAIV SIAELPLPPSPTTVSSVARSTIAAPTS SALSSQPIFTTAIDDRCELSSPREDTI PIPSLTSCETSDPLPTNENDDDDICK KPCSVAPNDIPLVSSSTNLINEINGVS EKLSTATESIVEIVKQEVLPPLTELEIL ENPPEEMKLECIAPITPSTVPSFPPT PPTPPASPPHTPVIVPAAATTVSSPSA AITVQRVLEEDDESIRTCLSEDAKEIQ NKIEVEADGQTEEILDSQNLNSRRS PVPAQIAITVPKTWKKPKDRTRTTEE MLEAELELKAEEELSIDKVLSEEQD KMSQGFHPERDPSDLKVKAVEEN GEEAEPVRNGAESVSEGEIDANS STDSGSGDVTFFPKPESWKPDTDETEG KKQYDREFLLDFQFMPACIQKPEGL PPISDVVLDKINQPKLPMRTLDPRIL PRGPDFTPAFADFRQTPGGRGVPL LNVGSRRSQPGQRREPRKIITVSVKE DVHLKKAENAWKPSQKRDSQADDP ENIKTQELFRKVR SILNKLTPQM FN QLMKQVSGLTVDTTEERLKGVIDLVF EKAIDEPFSVAYANMCRCLVTLKVP MADKPGNTVNFRLKLLNRCQKEFE KDKADDDVFEKKQKELEAASAPEER TRLHDELEEA KDKARRRSIGNIKFIG ELFKLMLTEAIMHDCVVKLLKNHD EESLECLCRLLTTIGKDLDFEKAKPR MDQYFNQMEKIVKERKTSSRIRFML QDVIDLRLCNVWSRRADQGPKTIEQ IHKEAKIEEQEEQRKVQQLMTKEKR RPGVQRVDEGGWNTVQGAKNSRVL DPSKFLKITKPTIDEKIQLVPAQLGS WGKGSSGGAKASET DALRSSASSLN RFSALQPPAPSGSTPSTPVEFDSRRT LTSRGS MGREKNDKPLPSATARPNT FMRGGSKDLLDNQSQEEQRREML ETVKQLTGGVDVERNSTEAERNKTR ESAKPEISAMSAHDKAALSEEELER KSKSIIDFLHINDFKEAMQCVEELN AQGLLHVFRVGVVESTLERSQITRD HMGQLLYQLVQSEKLSKQDFFKGFS ETLELADDMAIDIPHIWLYLAELVTP MLKEGGISMRELTIEFSKPLLPVGRA GVLLSEILHLLCKQMSHKKVGALWR EADLSWKDFLPEGEDVHNFLLEQKL DFIESDPCSSEALSKELSAEELYK RLEKLIHEDKANDEQIFDWVEANLDE IQMSSPTFLRALMTAVCKAAIADSS TFRVDTAVIKQRPILLKYLDSDTEK ELQALYALQASIVKLDQPANLLRMF FDCLYDEEVISED AFYKWESSK DPA EQNGKGVALKSVTAFFTWLREAE SEDN

