

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
Q6PJT7	ZC3HE_HUMAN	Homo sapiens	Zinc finger CCH domain-containing protein 14	36.958015	S75;T77;T78;S132;T133;S134;S135;T202;T204;S211;S220;S240;T325;S329;S331;S360;S365;T367;T369;T370;S373;T374;T387;T389;S390;S658;S735	S85;S240;S281;S327;S343;S390;S409;S421;S498;S515;S527;S620	32119511;34725712;35132862;31637018;29249144;23301498;32574038;31373491;30059200;20068230;35289036;33214551;29237092;30379171;34019948;35254053;37340703;28657654;35138101;33465208;34846842;30620550;31492838;29351928	MEIGTEISRKIRSAIKGKQLQELGAYVD EELPDYIMVMVANKKSQDQMTEDL SLFLGNNTRFTVWLHGVLDKLRV TTEPSSLKSSDTNIFDSNVPSNKSNF SRGDERRHEAAVPLAIPSAPEKRD SRVSTSSQESKTTNVRQTYDDGAAT RLMSTVKPLREPAPSEDVIDIKPEPD DLIDEDLNFVQENPLSQKKPTVTLTY GSSRPSIEIYRPPASRNADSGVHLNR LQFQQQNSIHAAKQLDMQSSWVY ETGRLCEPEVLNSLEETYSPPFRNN SEKMSMEDENFRKRKLPVVSVVK VKKFNHGDGEEEEEDDYGSRTGSIS SSVSVPKPERRPSLPPSKQANKNLI LKAISEAQESVTKTTNYSTVPQKQTL PVAPRTRTSQEELLAEVVQGQSRT RISPPIKEEETKGDSVEKNQGTQQR QLLSRLQIDPVM AETLQMSQDYD MESMVHADTRSFILKKPKLSEEVVV APNQESGMKTADSLRVLSGHLMQT RDLVQPDKPASP KFIVTLDGVPSPPG YMSDQEEDMCFEGMKPVNQTAAS NKGLRGLLHPQQLHLLSRQLEDPN GSFSNAEMSELSVAQKPEKLLERCK YWPACKNGDECAYHHPISPC KAFP CKFAEKCLFVHPNCKYDAKCTKPDC PFTHVSRRI PVLSPKPAVAPPAPSSS QLCRYFPACKMECPFYHPKHCRF NTQCTRPDCTFYHPTINVPPRHALK WIRPQTSE