

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
O08788	DCTN1_MOUSE	Mus musculus	Dynactin subunit 1	25.964899	T1252	T108;T145;T146;T147;S179;S212	22645316;36852467;33300544	MAQSRRHMSRTPSGSRMSTEASARPLRVGSRVEVIGKGRGTVAAYVGA TLFATGKWVGVILDEAKGKNDGTVOGRKYFTCDEGHGIFVRQSQIQVFED GADTTSPETPDSSASKVLKREGADA AAKTSKLRGLKPKKAPTARKTTTRRP KPTRPASTGVAGPSSSLGPSGSASAG ELSSSEPSTPAQTPLAAPIIPTALTSP GAAPPLSPSKEEEGLRAQVRDLEE KLETLRLKRSEDKAKLKELEKHKIQL EQVQEWKSKMQEQQADLQRRLKEA RKEAKEALEAKERYMEEMADTADAI EMATLDKEMAEERAESLQQEVEAL KERVDELTTDLEILKAEIEEKGSDDGA ASSYQLKQLEEQNARLKDALVRMRD LSSSEKQEHVKLQKLMEKKNQELE VVRQQRERLQEELSQAESTIDELKE QVDAALGAEEMVEMLTDRNLNLEE KVRELRETVGDLEAMNEMNDELQENARETELELREQLDMAGARVREAQ KRVEAAQETVADYQQTIKKYRQLTA HLQDVNRELTNQQEASVERQQQPP PETFDKIKFAETKAHAKAIEMELRQ MEVAQANRHMSLLTAFMPDSFLRP GGDHDCVLVLLMPRLICKAELIRK QAQEKFDLSENCSEKPLRGAAGEQ LSFAAGLVYSLSLQATLHRYEHALS QCSVDVYKKVGSLYPEMSAHERSLD FLIELLHKDQLDQETVNVPLTKAIKY YQHLYSIHLAEQPEDSTMQLADHIK FTQSALDCMGVEVGRRLRAFLOGGQ EATDIALLLRDLETSCSDTRQFCKKI RRRMPGTDAPGIPAALAFGSQVSDT LLDCRKHLLTWVAVLQEVAAAAAQL IAPLAENGLPVAALEELAFKASEQI YGSPPSSPYECLRQSCSTILISTMNKL ATAMQEGEYDAERPPSKPPVELRA AALRAEITDAEGLGLKLEDRETVIKE LKKSLKIKGEELSEANVRLSLEKKL DSAAKDADERIEKVQTRLDETQTLLR KKEKDFEETMDALQADIDQLEAEKA ELKQRLNSQSKRTIEGLRGPPPSGIA TLVSGIAGEEPQRGGAPGQAPGALP GPGLVKDSPLLLQOISAMRLHISQLQ HENSILRGAQMKASLAALPPLHVAK LSLPPHEGPGGNLVAGALYRKTSQL LEKLNQLSTHVVVDITRSPAAKSP SAQLMEQVAQLKSLSDTIEKLKDEV LKETVTQRPGATVPTDFATFPSSAFL RAKEEQDDTVYMGKVTFSACAAGL GQRHRLVLTQEQLHQLHSRLIS

