

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
Q5VZ89	DEN4C_HUMAN	Homo sapiens	DENN domain-containing protein 4C	18.333285	S1634;T1636	S703;S737;S741;S965;S968;S973;T975;S989;S996;S1003;S1046;S1061;S1099;S1126;S1184;S1225;S1244;S1252;S1278;S1325;S1337;S1346;S1623;S1627;S1629;S1640;S1799;S953	38253038	<p> MIEDKGPRVTDYFVVAGLTDSTLLD QEINRLDTKSTGPKAPITDIAIIKSAG ETVPEGYTCVEATPSALQANLNYGS LKSPFLCYKRGTRDKPPLTDIGVLY EGKERLIPGCEVILATPYGRCANVNN SSTTSQRIFITYRRAPPVRPQNSLAVT DICVIVTSKGETPPHTFCKVDKNLNC GMWGSSVFLCYKKSVPASNAIAYKA GLIFRYPEEDYESFPLSESDVPLFCL PMGATIECWDPETKYPLPVFSTFVLT GSSAKKVYGAAIQFYEPYSRELLSEK QLMHLGLLTPVERKMVSKSINTNKC ICLLSHWPFEEAFRKFMLFIYKLSVS GPHPLPIEKHISHFMQNIPFPSPQRP RILVQLSVHDALILSQPVSTPLPLSGA NFSTLLMNLGPENCATLLLFVLLS KILLHSLRPAVLTGVAEAVVAMIFPF QWQCPYIPLCPLSLA AVL SAPLPFIV GVDSRYFDLHDPQDVVICDLDTNM LYVSDKKNMNWKQLPKKPKCNLL STLKKLYQLSSVHQKTQEGSAIDM TPIEADFSWQKKMTQLEMEIQEAF RFMASILKGYRTYLRPITEAPSNKAT AADSLFDRQGFLKSRDRAYAKFYTL LSKTQIFIRFIEECFVSDKDTGLAFF DDCIEKLFDPDKGTEKTDKVD FDSAE DTRLIELDDSQSEHTVFIMPPEPPP DDGKDLSPKYSYKYFPRDLKLFDRP QELKLCFSRHPTGNSITKSPPLMAK RTKQEIKTAHKLAKRCYTNPPQWAK CLFSHCYSLWFICLPAYVRVSHPKVR ALQQAYDVLIKMRKTDVDP LDEV CY RVVMQLCGLWGHPVLAVRVL FEMK TARIKPNAITYGYYNKVVLESPWPSS TRSGIFLWTKVRNVVRGLAQFRQPL KKT VQRSQVSSISGGQSDQGYGSKD ELIKDDAEIHVPEEQAARELITKTKM QTEEVCDASAI VAKHSQSPPEPHSPT EPPAWGSSIVKVP SGIFDVNSRKSST GSISNVLFSTQDPVEDAVFGEATNL KKN GDRGEK RQKHFPERSCSFSSES RAGMLLKSSLD SNSSEM AIMMGA DAKILTAALTCPKTSLLHIARTHSFE NVSCHLPDSRTC MSESTWNPEHRS SPVPEMLEESQELLEPVVDDVPKTT ATVDTYESLLSDS NSNQSRDLKTVS KDLRNKRSSLYGIKVVQREDVETG LDPLSLLATECTGGKTPDSEDKLFSP VIARNLADEIESYMN LKSPLGSKSSS </p>

MELHREENRESGMTTAFIHALERRS
SLPLDHGSPAQENPESEKSSPAVSR
SKTFTGRFKQQTPSRTHKERSTLSA
LVRSSPHGSLGSVVNSLSGLKLDNIL
SGPKIDVLKSGMKQAATVASKMWV
AVASAYSYSDDDEETNRDYSFPAGL
EDHILGENISPNTSISGLVPSELTS
NTSLGSSSSSGDVGKLHYPTGEVPF
PRGMKGQDFEKSDHGSSQNTSMSS
IYQNCAMEVLMSSCSQCACGALVY
DEEIMAGWTADDSNLNTACPFCKS
NFLPLLNIEFKDLRGSASFFLKPSTS
GDSLQSGSIPLANESLEHKPVSSLAE
PDLINFMDFPKHNIITEETGSAVEP
SDEIKRASGDVQTMKISSVPNSLSKR
NVSLTRSHSVGGPLQNIIDFTQRPFH
GISTVSLPNSLQEVVDPLGKRPNPPP
VSPYLSPLVLRKELESLENEGDQV
IHTSSFINQHPIIFWNLVWYFRRLDL
PSNLPGLILTSEHCNEGVQLPLSSLS
QDSKLVYIQLLWDNINLHQEPREPL
YVSWRNFNSEKKSSLLSEEQQTST
LVETIRQSIQHNNVLKPINLLSQQM
KPGMKRQRSLYREILFSLVSLGRE
NIDIEAFDNEYGIAYNSLSSEILERLQ
KIDAPPSASVEWCRKCFGAPLI