

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
Q9P219	DAPLE_HUMAN	Homo sapiens	Protein Daple	15.734917	S1496;T1519;S1521;T1522;S1523;T1525;S1791	S227;S239;S486;S1444;S1601;S1806;T1954	34019948;35254053;30059200;29237092;37340703;31492838;30379171	MDVTVSELLELFLOSPLVTWVKTFG PFGSGSQDNLTMYMDLVDGIFLNQI MLQIDPRPTNQRINKHVNDVNLRI QNLTLVRNIKTYQEVLQQLIVMNL PNVLMIGRDPLSGKSMEEIKVLLL VLGCAVQCERKEEFIERIKQLDIETQ AGIVAHIQEVTHNQENVFDLQWLEL PDVAPEELEALSRSMLVHLRRLIDQ RDECTELIVDLTQERDYLQAQHPPSP IKSSADSTPSPTSSLSSSEDKQHLLAV ELADTKARLRVRQLEEDKTEQLVD TRHEVDQLVLELQKVKQENIQLAAD ARSARAYRDELDLREKANRVERLE LELTRCKEKLHDVDFYKARMEELRE DNIILJETKAMLEEQLTAAARAGDKV HELEKENLQLKSKLHDELEDRDTRDK KRIEELLEENMVLEIAQKQSMNESA HLGWELEQLSKNADLSDASRKSFVF ELNECASSRILKLEKENQSLQSTIQG LRDASLVLEESGLKCGELEKENHQL SKKIEKLQTQLEREKQSNQDLETLS EELIREKEQLQSDMETLKADKARQI KDLEQEKDHLNRAMWSLRERSQVS SEARMKDVEKENKALHQTVTEANG KLSQLEFEKRQLHRDLEQAKEKGER AEKLERELQRLQEENGLRARKVTSL ETATEKVEALEHESQGLQLENRTLRL KSLDTLQNVSLQLEGLERDNKQLDA ENLELRLVETMRFSTKLAQMERE NQQLEREKEELRKNVDLLKALGKKS ERLELSYQSVSAENLRLQSSLESSS HKTQTLESELGELEAERQALRRDLE ALRLANAQLEGAEKDRKALEQEVAQ LEKDKLLEKEAKRLWQVVELKDAV LDDSTAKLSAVEKESRALDKELARC RDAAGKLKELEKDNRDLTQVTVHA RTLTTTLREDLVLEKLSQQLSSELDK LSQELEKVGLENRELLQEDDSGSDT KYKILEGRNESALKTTLAMKEEKIVL LEAQMEEKASLNROLESELQMLKK ECETLRQNGEGQHLQNSFKHPAG KTAASHQGKEAWGPGHKEATMELL RVKDRAIELERNNAALQAEKQLLKE QLQHLETQNVTFSSQILTLQKQSAFL QEHNTTLQTQTAKLQVENSTLSSQS AALTAQYTLQNHHTAKETENESLQ RQQEQLTAAYEALLQDHEHLGTLHE RQSAEYEALIRQHSCLKTLHRNLEL EHKELGERHGDMLKRKAELEEREK VLTTEREALQQEQRTNALAMGENQ RLRGELDRVNFLHHQLKGEYEELH AHTKELKTSLNNAQLELNRWQARF DELKEQHQTMDISLTKLDNHCELLS RLKGNLEENHHLLSQIQLLSQQNQ MLLEQNMENTKQYHEEQKQYIDKL NALRRHKEKLEEKIMDQYKFDPPP KKKNHWIGAKALVKLIKPKKEGSRE RLKSTVDSPPWQLESSDPASPAASQ PLRSQAENPDTPALGNSNCAERDAH NGSVGKGPGLKPKRGSPhrgSLD RTDASTDLAMRWPSELGSRCTSTS ATTTAPSNSTPIARHPGRTRKGYNSDD

											NLCEPSLEFEVFNHRQYVSRPSSLE SSRNTSSNSSPLNLKGSSEQLHGRS ESFSSDLIPSRDLATLPREASTPGR NALGRHEYPLPRNGPLPQEGAQKRG TAPPYVGVRPCSASPSEMVTLEEFLL EESNRSSPHTDTPSCRDDLLSDYFR KASDPPAIGGQPGPPAKKEGAKMPT NFVAPTVMMAAPTSEGRPLKPGQYV KPNFRLTEAEAPPSVAPRQAQPPQSL SLGRPRQAPVPPASHAPASRSASLSR AFSLASADLLRASGPEACKQESPQK LGAPEALGGRETGSHTLQSPAPPSS HSLARERTPLVGKAGSSCQGPGRS RPLDTRRFSLAPPKEERLAPLHQSAT APAIATAGAGAAAAGSGSNSQLLHF SPAAAAPAARTKPKAPPRSGEVATITP VRAGLSLSEGDGVPGQGCSEGLPAK SPGRSPDLAPHLGRALEDCSRGSVS KSSPASPEPGGDPQTVWYEGCV
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