

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus
O14617	AP3D1_HUMAN	Homo sapiens	AP-3 complex subunit delta-1	22.500897	NaN	S632;S634;S636;S658;S688;S758;S759;T762;S764;S788;S829;S785;S828;S931	40914422;34019948;37217939;28510447	MALKMVKGSIDRMFDKNLQDLVRG IRNHKDEAKYISQCIDEIKQELKQD NIAVKANAVCKLTYLQMLGYDISWA AFNIIIVMSASKFTFKRIGYLAASQS FHEGTDVIMLTTNQIRKDLSSPSQY DTGVALTGLSCFVTPDLARDLANDI MTLMSHTKPYIRKKAVALIMYKVFVKY PESLRPAFPRLKEKLEDPDPGVQSA VNVICELARRNPKNYLSLAPLFFKL MTSSTNNWVLIKIKLFGALTPLEPR LGKKLIEPLTNLIHSTSAMSLLYECV NTVIAVLISLSSGMPNHSASIQLCVQ KLRILIEDSDONLKYLGLLAMSKILK THPKSVQSHKDLILQCLDDKDESIRL RALDLLYGMVSKKNLMEIVKCLMT HVDKAEGTTYRDELLTKIIDCSQSN YQYITNFEWYISILVELTRLEGTRHG HLIAAQMLDVAIRVKAIRKFAVSQMS ALLDSAHLLASSTQRNGICEVLYAAA WICGEFSEHLQEPHHTLEAMLRPRV TTLPGHIQAVYVQNVVKLYASILQOK EQAGEAEGAQAVTQLMVDRLPOFV QSADLEVQERASCILQLVKHIQKLOA KDVPVAEEVSALFAGELNPVAPKAO KKVPVPEGLDLDAWINEPLSDSESE DERPRAVFHEEEQRRPKHRPSEADE EELARRREARKQEQANNPFYKSSP SPQKRYODTPGVEHIPVQIDL SVPL KVPGLPMSDQYVVKLEERRHRQKLE KDKRRKRKEKKGKRRHSSLPTE SDEIAPAQQVDIVTEEMPENALPS DEDDKDPNDPYRALDIDLKPLADS EKLPIQKHRNTEKSKSPEKDVPMVE KSKKPKKKEKKHKEKERDKEKKKE KEKKSPKPKKKKHRKEKEERTKGG KSKKQPPGSEEAAGEFPVQNGAPEE EQLPPESSYLLAENSIVKMTCDIRG SLQEDSQVTVAVLENRSSSILKGM LSVLDLNLNARMARPOGSSVHDGVPV PFQLPGVSNQAQYVFTIQSIVMAQK LKGTL SFIKNDGATHEKLDLFRH FSCSSYLITTPCYSDAFAKLLSGLS MSSIKVDGIRMSFQNLAKICFHHH FSVVERVDS CASMYSRSIQGHHVCL LVKKGENSVDGKCSDSLNSLL EEMKATLAKC	False	False	2.274	3.677	1.692	1.222	4.346