

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
Q08951	AP3D_YEAST	Saccharomyces cerevisiae	AP-3 complex subunit delta	9.365641	NaN	S700;S727;T767;S770;S773;S798;S888;S918	33229814	<p>MTSLYAPGAEDIRQLRPFQFFFEKS LKDLIKGIRSHNETPEKLDQFFKQVL SECREEVNSPDLNSKTNVAVLKLTYL EMYGFDMAWCNFHILEVMSSNKLQ QKRVGYLAASQSFKDSDILMLATN LLKKDLKYDGNNDVVKVGIASGLS TIITPSLARDIADDLFTMLNSTRPYIR KKAITALFKVFLQYPEALRDNFDKFV SKLDDDDISVVSAAVVICELSKKNP QPFIQLSPLLYEILVTIDNNWIIIRLLK LFTNLSQVEPKLRKLLPKILELMES TVATSVIYESVNCIVKGNMLEEDDFE TAMACLERLHTFCDSQDPNLRYSICI LFYKIGKINTDFISRFDQLIIRLLSDV DVSIRSKAIELVEGIVDEDNLKAIVQT LMKQFVDEDVVILQGTGSIVYEKSKRI PIIPENYKIKMVNVIISICSADNYSSV NDFEWYNAVIMDLAMLCQDISDKS LGSKIGEQRNLMIKVPSMREVTIA NIIKLISNDNINKQLPTVLRECIWCL GEFSTLVENGNDLIKIMTENISYYSH SVQEVLLILALVKVFSNWCNNFQEDK RFEIKMVLKELIEFFENLSYSSTFEV QERSVEVLEFLRLSLEALEEDTEGLP MLLSEVLPSFFNAYELAPIARGTQLK LAVDENLDLETPFLTKEAADELLEDE QKSDAISDLMSDISMDEQVELKFVD DSDTSYEEKEKLDDFENPFEIEREKE RMSNPYYLGEDEEERTKNSKDLLDL NEEESDCKPETIRLNRTDNSLNSL SLSTTEISRKKKKGKKNRVQVLSDE PVIEAAPKRKDAFQKPHDNHSTQNP LKKDKINLRMHSQLENFDFSNFGQ SSNAGRGSQEEGNLRKEDELELSRL EANLIVKDEKDNLSDTEEVIVIKKKK KGKKSksknklktkakspepnefl RDQSTDI</p>