

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membrane	ext req
000203	AP3B1_HUMAN	Homo sapiens	AP-3 complex subunit beta-1	18.152543	S671	S276;S609;S750;S752	30379171;23301498;9502801;34019948;32119511;37217939;38665916	MSSNSFPYNEQSGGGEATELGQEATSTISPSGAFGLFSSDLKKNEIDLKQMLESNKDSAKLDAMKRIVGMIAKGKNASELFPVAVKKNVASKNIEIKLVVYLVRYAEEQDLALLSISTFQRALKDPNQLIRASALRVLSSIRVPIIPIIMMLAIKEASADLSPYVRKNAAHAIQKLYSLDPEQKEMLIEVIEIKLLKDKSTLVAGSVVMAFEEVCPDRIDLHKNYRKLCLNLLVDVEEWGQVVIHMLTRYARTQFVSPWKEGDELEDNGKNFYESDDDQKEKTDKKKKPYTMDPDHRLLRINTKPLLQSRNAAVVMAVAQLYWHISPKSEAGIISKSLVRLRSNREVQYIVLQNIATMSIQRKGMFEPYKSFYVRSTDP TMIKTLKLEILTNLANEANISTLLREFQTYVKSQDKQFAAATIQTIGRCATNI LEVTDTCNLGLVCLLSNRDEIVVAESVVIKLLQMQPAQHGEIHKHMAKLLDSITVVARASILWIGENCERVPKI APDVLKMAKSFTSEDDLVLKQILNLGAKLYLTNSKQTKLLTQYILNLGKYDQNYDIRDRTRFIRQLIVPNVKS GAL SKYAKKIFLAQKPAPLLES PFKDRDH FQLGTLSHTLNIKATGYLELSNWPE VAPDPSVRNVEIELAKEWTPAGKA KQENS AKKFYSESEEEEDSSDSSD SESESGESGEQGESGEGDSNEDS SEDSSEQDSESGRESGLENKRTAK RNSKAKGKSDSEEDGEKENEKSTSD SSNDESSSIEDSSDSESESEPESES ESRRVTKEKEKTKQDRTP LTKDVS LLDLDDFNVPSTPVALPTALSPSLM ADLEGLHLSTSSSVISVSTPAFVPTK THVLLHRMSGKGLAAHYFFPRQPCI FGDKMVS IQITLNNTDRKIENIHIG EKKLPIGMKMHVFNPIDSLPEGSIT VSMGIDFCDSTQTASFQLCTKDDCF NVNIQPPVGELLLPVAMSEKDFKKE QGVLTMNETSAVIAAPONFTPSVI FQKVVNVANVGAVPSGQDNHRFAA KTVHSGSLMLVTVELKEGSTAQLIIN TEKTIVGSVLLRELKPVLSQG	False	True	2.373	3.173	3.073	1.192	4.486	2.077	2.9