	UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochndrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extrace region
0	Q05140	AP180_RAT	Rattus norvegicus	Clathrin coat assembly protein AP180	31.640232	T310	S296;S300;S306;S313;T317;S594;S600;S640;S646;S775;S600;S627	18683930;21500857	MSGQTLTDRIAAAQYSVTGSAVARA VCKATTHEVMGPKKKHLDYLIQATN ETNVNIPQMADTLFERATNSSWVVV FKALVTTHHLMVHGNERFIQYLASR NTLFNLSNFLDKSGSHGYDMSTFIR RYSRYLNEKAFSYRQMAFDFARVKK GADGVMRTMVPEKLLKSMPILQGQI DALLEFDVHPNELTNGVINAAFMLL FKDLIKLFACYNDGVINLLEKFFEMK KGQCKDALEIYKRFLTRMTRVSEFL KVADEVGIDKGDIPDLTQAPSSLMET LEQHLNTLEGKKPGNNEGSGAPSPL SKSSPATTVTSPNSTPAKTIDTSPPVD IFATASAAAPVSSAKPSSDLLDLQPD FSGARAGAAAPVPPPTGGATAWGDL LGEDSLAALSSVPSEAPISDPFAPEP SPPTTTTEPASASASATTAVTAATTEV DLFGDAFAASPGEAPAASEGATAPAT PAPVAAALDACSGNDPFAPSEGSAE AAPELDLFAMKPPETSAPVVTPTAST APPVPATAPSPAPTAVAATAATTTAA AAATTTATTSAAAATTAAAPPALDIFG DLFDSAPEVAAASKPDVAPSIDLFGT DAFSSPPRGASPVPESSLTADLLSGS GFHCAEDDRHVPLFFTAVDAFAAPS PASTASPAKAESSGVIDLFGDAFGSS ASETQPAPQAVSSSSASADLLAGFG GSFMAPSTTPVTPAQNNLLQPNFEA AFGTTPSTSSSSSFDPSGDLLMPTM APSGQPAPVSMVPPSPAMSASKGLG SDLDSSLASLVGNLGISGTTSKKGDL QWNAGEKKLTGGANWQPKVTPATW SAGVPPQGTVPPTSSVPPGAGAPSVG QPGAGYGMPPAGTGMTMMPQQPV MFAQPMMRPPFGAAAVPGTQLSPS PTPATQSPKKPPAKDPLADLNIKDFL	None	None	None	None	None	None	None	None	None