

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	extracellular region
P32454	APE2_YEAST	Saccharomyces cerevisiae	Aminopeptidase 2, mitochondrial	24.475439	NaN	NaN	33229814	MPIVWRWLLKSAVRGSSLLIGKAHPCL RSIAAHPRYLNSVYSPAGVSRSLRI NVMWVKQSKLTPPRFVKIMNRRPLFT ETSACAKCQKTSQLLNKTNPNEIL PDNVVPLHYDLTVEPDFKTFKFEGS VKIELKINNPAIDTVTLNTVDTDIHSA KIGDVTSEIIEEEQVTTFAFPKGT MSSFKGNAPLDIKFTGILNDNMAGF YRAKYEDKLTGETKYMATTQMEPTD ARRAFPCFDEPNLKASFAITLVSDPS LTHLSNMDVKNEYVKDGKKVTLFN TTPKMSTYLVAFIVAELKYVESKNFR IPVRVYATPGNEKHGQFAADLTAKTL AFFEKTFGIQYPLPKMDNVAVHEFS AGAMENWGLVTYRVVLLLDKDNS TLDRIQRVAEVVQHELAHQWFGNLV TMDWWEGLWLNDFATWMSWYS CNEFQPEWKVWEQYVDTLQHALS LDSLRSSHPIEVVKKADEINQIFDAI SYSGASLLRMISKWLGEETFIKGVV QYLNKFKYGNAKTEDLWDALADAS GKDVRSVMNIWTKKVGFPVISVSED GNGKITFRQNYLSTADVDPDEDKTI YPVFLALKTKNGVDSVLSERSKTI ELEDPTFFKVNSEQSGIYTSYTDER WAKLGGQADLLSVEDRVGLVADVKT LSASGYTSTTNFLNLVSKWNNEKSF VVWDQIINSISSMKSTWLFEPKETO DALDNFTKQLISGMTHHLGWEFKS SDSFSTQRLKVTMFGAACAAARDADV EKAALKMFTDYCSGNKEAIPALIKPI VFNTVARVGGAEYKVKIYLDPI NDEKLAALRSLGRFKEPKLLERTLG YLFDGTVLNQDIYIPMQGMRAHQEG VEALWNWVKKNWDELVKRLPPGLS MLGSVVTLGTSGFTSMQKIDEIKKFF ATKSTKGFQSLAQSLDTITSKAQW VNRDRDVVNKYLKENGY	None	None	None	None	None	None	None	None	

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