

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
P16406	AMPE_MOUSE	Mus musculus	Glutamyl aminopeptidase	24.048659	NaN	NaN	33300544;37507081	MNFAEEEPSKKYCIKGGHVAICGVV VAVGLIVGLSVGLTRSCQDITPAPS QPPEASTALPPQDNVCPDSEDES GEWKNFRLPDFINPVHYDLEVKALM EEDRYTGIVTISVNLSKPTRDLWLHI RETKITKLPRLRRPSGEQVPIRRCFE YKKQEVVVIQAAEDLAATSGDSVYRL TMEFKGWLNGSLVGFYKTTYMEDG QIRSIATDHEPTDARKSFPCFDEPN KKSTYSISIIHPKEYSALSMPPEEKSE MVDDNWKKTTFVKSVPMSYLVCF AVHRFTAIERKSRSGKPLKVYVQPN QKETAEYAAANITQAVFDYFEDYFAM EYALPKLDKIAIPDFGTGAMENWGL VTYRETNLLYDPLLSASSNQORVAS VVAHELVHQWFGNTVTMDWDDDL WLNFGFASFFFLGVNHAEKDWQ MLSQVLEEDVFPVQEDDSLMSHPV VVTVSTPAEITSVFDGISYSKGASILR MLQDWITPEKFKQKGCQYLLKFFQFA NAKTSDFWDSLQEASNLPVKEVMD TWTSQMGYPVTVSQRONITQKRFL LDSKADPSQPPSELGYTWNIPVRWA DNDNSRITVYNNRLDKGGITLNANLS GDAFLKINPDHIGFYRVNIEGGTWD WIAEALSSNHTRFSAADRSSFIDDAF ALARAQLLNKYIALNLTYMKSEEDF LPWERVISSVSYIISMFEEDDRELYPM IETYFQGQVKPVADLLGWQDTGSHI TKLLRASILGFACKMGDREALGNAS QLFDSWLKGSASIPVNLRLLVYRYG MQNSGNEAAWNYTLEQYQKTSLAQ EKEKLLYGLASVKDVKLLARYLEML KDPNIIKTQDVFTVIRYISYNSYGKTM AWNWIQLNWDYLVSRFTINDRYLG RIVTIAEPFNTELQLWQMOSFFAKYP NAGAGAKPREQVLETVKNNIEWLN VNRQSIREWFASLP	False	True	1.601	2.006	1.64	2.861	0.892	5.0	3.888