

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	extrac region
Q8C129	LCAP_MOUSE	Mus musculus	Leucyl-cystinyl aminopeptidase	25.875365	NaN	T70;S80;S91	39627609;33300544;37507081	MESFTNDRLQLPRNMIENSMFEEE PDVVDLAKEPCLHPLEPDEVEYEPR GSRLLVRLGEGHEMDEDEEYESSA KLLGMSFMNRSSGLRNSAAGYRQS PDGTCSLPSARTLVICVFVIVAVSVI MVIYLLPRCTFTKEGCHKTNQSAELI QPVATNGKVFPPWAQIRLPTAIPLCY ELSLHPNLTSMTRFGSVTISLQALQ DTRDIILHSTGHNISRVTFMSAVSSQ EKQVEILEYYPYHEQIAVVAPELLTG HNYTLKIEYSANISNSYGFYGYTYTD KSNEKKYFAATQFEPLAARSAFPCF DEPAFKATFIKITRNEHHTALSNMP KSSVPAEEGLIQDEFSESVKMSTYL VAFIVGEMRNLSQDVNGTLVSVYAV PEKIGQVHHALDTTIKLLEFYQTYFEI QYPLKKLDLVAIPDFEAGAMENWGL LTFREETLLYDNATSSVADRKLVTKII AHELAHQWFGNLTVMQWVNDLW LNEGFAFMFYFSVEKIFKELNSYE DFLDARFKTMRKDSLNSHPISSSV QSSEQIEEMFDSL SYFKGASLLML KSYLSEDFVRHAVILYLHNHSYAAIQ SDDLWDSFNEVTDKTLDVKKMMKT WTLQKGFPLVTVQRKGTLLQQRER FFLRMQPESQPSDTSHLWHIPISYVT DGRNYSEYRSVSLDDKSDVINLTE QVQWVKVNSNMTGYIVHYAHDDW TALINQKRNYPVLSDKDRANLINNI FELAGLGKVPRLRMAFDLIDYLNKNET HTAPITEALFQTNLIYNLLEKLGHMD LSSRLVARVHKLLQNIQQQTWTDE GTPSMRELRSALLEFACAHSLENCT TMATNLFDSWMASNGTQSLPTDVM VTVFKVGARTEKGWLFVLSMYSSM GSEAEKNKILEALASEDVHKLYWL MKSSLDGDIIRTQKLSLIHRTVGRHFP GHLLAWDFVKENWNKLVHKFHLGS YTIQSVAGSTHLFSTKTHLSEVQAF FENQSEATLKLRCVQEALEVIQLNIQ WMVRNLKTLTSQWL	False	True	2.706	2.593	1.923	2.168	2.081	5.0	2.294