

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
Q9UIQ6	LCAP_HUMAN	Homo sapiens	Leucyl-cystinyl aminopeptidase	30.529297	S147;T156	T70;S80;S91	40914422;35289036;34019948;37340703	MEPFTNDRLQLPRNMIENSMFEEE PDVVDLAKEPCLHPLEPDEVEYEPR GSRLLVRLGEGHEMEDEEYESSA KLLGMSFMNRSGLRNSATGYRQS PDGACSVPSARTMVVCAFVIVAVSV IMVIYLLPRCTFTKEGCHKKNQSIGL IQPFATNGKLPWAQIRLPTAVVPLR YELSLHPNLTSMTRGVSIVSQALQ VTWNILHSTGHNISRVTFMSAVSS QEKQAEILEYAYHGQIAIVAPEALLA GHNYTLKIEYSANISSYYGFYGFYSY TDESNEKKYFAATQFEPLAARSAFP CFDEPAFKATFIHKIRDEQYTALSNM PKKSSVLLDDGLVQDEFSESVKMST YLVAFIVGEMKNLSQDVNGTLVSIYA VPEKIGQVHYALETIVKLEFFQNYF EIQYPLKKLDLVAIPDFEAGAMENW GLLTFREETLLYDSNTSSMADRKL TKIIAHELAHQWFGNLVTMKWWND LWLNEGFAFMFYFSLEKIFKELSSY EDFLDARFKTMKKDSLNSHPISSS VQSSEQIEEMFDSLVSFKGSSLLLM LKTLYLSEDFVQHAVVLYLHNHSYASI QSDDLWDSFNEVTNQTLDVKRMM KTWTLQKGFPLVTVQKKGKELFIQQ ERFFLNMKPEIQPSDTSYLWHIPLSY VTEGRNYSKYQSVSLDKKSGVINLT EEVLWVKVINMNGYIVHYADDD WEALIHQLKINPYVLSKDRANLIN NIFELAGLGKVPKRAFDLINYLGN NHTAPITEALFQTDLIYNLLEKLG DLASRLVTRVFKLLQNQIQQTWTD EGTPSMRELRALLEFACTHNLGNC STTAMKLFDDWMASNGTQSLPTDV MTTVFKVGAKTDKGWSFLLGKYISIG SEAEKNKILEALASSEDVRKLYWLM KSSLNGDNFRITQKLSFIIRTVGRHFP GHLLAWDFVKENWNKLVQKFPPLGS YTIQNVAGSTYLFSTKTHLSEVQAF FENQSEATFRLRCVQEALEVIQLNIQ WMEKNLKSILTWWL	True	True	4.511	2.12	1.736	2.094	1.574	4.453