	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	extracellular region
0	Q9VD85	Q9VD85_DROME	Drosophila melanogaster	Aminopeptidase	28.06426	S2	NaN	40245499	MSSSIFRAFWLLCSLLICVTNAADYR LEGSVVPSHYNLTIGVLRNSVEPTIF DGEVSITLRVVGTLEVQQIILHADTL DITECWLLDAAGAQVEAIDISRLIYEA ATQQVRVPLTEAAQPGKNYTLGFKY TGHIRTDMAGFFSASYVERDTNVTR WLALTQMQRINARLVLPCFDEPALK AQFQLQIVRPNGYQSIANTKLKETKA LSQDRFVDHFKETPVMSTYLLAFMV ANYSARGNESEFAVLTRPEFYDNTE FSYHVGQQVVSAYGELFQSPYAELG NDVLQYASSPRFPHNGMENWGLIIY SDDVLIQEPGYSDDWSDKEFAIRIIA HETSHMWFGDSVTFSWWSYFWLN EAFARYYEYFMAHQLYPDYHLDEQF VVRQMQLIFGTDAQNGTQPMTSPES EIQTPSQIAYKFSGIAYAKGACIVRM WRNLMGAENFDTAIRSYLQQYHLT NTVPYNLFYHLFEHWPKNQDVDLV DFLTDYTEQVGYPMIIVKALQGNKVV TVEQSRFLLNKDDGSNATLKYTVPIT FTTNLERDFYNLTPYKYLHKTVNVL WLDFTSPIDWILFNLRQSNYQRVFY DDTLREGLLLAISATNHSGIPVENRA QIIDDLFNFAQAQYVDYADVFRFLEY LAKEVEYVPWYAVYENLNTVAKRLT PEQLPDFRIYLSSITAAVFEKLGVGW NAQDTPLDVANRNKLIKWLCRYQAS NCRPKVNVQFVTGHEVPSPDYRETF YCAASGADYNSYTMLWGQYYLENR TSEKNMLWAAISCTRDYETHYHKMI ISGTESVEQKKIGIARMYEENPDLVQ AIFEMITENIMELAYNFQSWSDTAE VINDMAEYFTTREQQQLLKEFYYSN HQFFGESEELLSKSLVTVEKNLNWA ETHLEGLVKYLADRNGSSCLGATSL VVVLLLAVGSMLAWP	False	False	1.737	1.536	1.757	1.822	1.369	2.086	3.733