	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	P05555	ITAM_MOUSE	Mus musculus	Integrin alpha-M	25.685855	NaN	NaN	39627609	MTLKALLVTALALCHGFNLDTEHPM TFQENAKGFGQNVVQLGGTSVVVAA PQEAKAVNQTGALYQCDYSTSRCHP IPLQVPPEAVNMSLGLSLAVSTVPQQ LLACGPTVHQNCKENTYVNGLCYLF GSNLLRPPQQFPEALRECPQQESDI VFLIDGSGSINNIDFQKMKEFVSTV MEQFKKSKTLFSLMQYSDEFRIHFT FNDFKRNPSPRSHVSPIKQLNGRTK TASGIRKVVRELFHKTNGARENAAKI LVVITDGEKFGDPLDYKDVIPEADRA GVIRYVIGVGNAFNKPQSRRELDTIA SKPAGEHVFQVDNFEALNTIQNQLQ EKIFAIEGTQTGSTSSFEHEMSQEGF SASITSNGPLLGSVGSFDWAGGAFL YTSKDKVTFINTTRVDSDMNDAYLG YASAVILRNRVQSLVLGAPRYQHIGL VVMFRENFGTWEPHTSIKGSQIGSY FGASLCSVDMDADGNTNLILIGAPH YYEKTRGGQVSVCPLPRGRARWQCE ALLHGDQGHPWGRFGAALTVLGDV NGDKLTDVAIGAPGEQENQGAVYIF YGASIASLSASHSHRIIGAHFSPGLQ YFGQSLSGGKDLTMDGLMDLAVGA QGHLLLLRAQPVLRLEATMEFSPKK VARSVFACQEQVLKNKDAGEVRVCL RVRKNTKDRLREGDIQSTVTYDLAL DPVRSRIRAFFDETKNNTRRRTQVF GLMQKCETLKLILPDCVDDSVSPIIL RLNYTLVGEPLRSFGNLRPVLAMDA QRFFTAMFPFEKNCGNDSICQDDLS ITMSAMGLDTLVVGGPQDFNMSVT LRNDGEDSYGTQVTVYYPSGLSYRK DSASQNPLTKKPWFVKPAESSSSE GHGALKSTTWNINHPIFPANSEVTF NVTFDVDSHASFGNKLLLKAIVASE NNMSRTHKTKFQLELPVKYAIYMIV TSDESSIRYLNFTASEMTSKVIQHQY QFNNLGQRSLPVSVVFWIPVQINNV TVWDHPQVIFSQNLSSACHTEQKSP PHSNFRDQLERTPVLNCSVAVCKRI QCDLPSFNTQEIFNVTLKGNLSFDW YIKTSHGHLLLVSSTEILFNDSAFAL LPGQESYVRSKTETKVEPYEVHNPV PLIVGSSIGGLVLLALITAGLYKLGFF KRQYKDMMNEAAPQDAPPQ	True	True	3.498	5.0	3.39	2.692	2.118	5.0	4.373