

UniprotKB ID	Entry name	organism	full name	oglcnaicscore	oglcnaicsites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracellular region
P24063	ITAL_MOUSE	Mus musculus	Integrin alpha-L	31.413142	NaN	NaN	29187734;39627609	MSFRIAGPRLLLGLQLFAKAWSYN LDTRPTQSFLLAQAGRHFGYQVLQIE DGVVVGAPGEGDNTGGLYHCRSTSE FCQPVSLHGSNHTSKYLGMTLATDA AKGSLACDPGLSRTCDQNTYLSGL CYLFPQSLEGPMLQNRPAQECMK GKVDLVFLFDGSQSLDRKDFEKILEF MKDVMRKLNTSYQFAAVQFSTDC RTEFTFLDYVKQKNPDVLLGSVQP MFLLTNTFRAINYVVAHVFKKEESGA RPDATKVLVIITDGEASDKGNISAAH DITRYIIGIGKHFVSVQKQKTLHIFAS EPVEEFVKILDTFEKLKDLFTDLQRR IYAIEGTNRQDLTSFNMELSSSGISA DLKSGHAVVGAVGAKDWAGGFLDL REDLQGATFVGOEPLTSDVRGGYLG YTVAWMTSRSSRPLLAAGAPRYQHV GOVLLFQAEAGGRWNQTKIEGTQ IGSYFGGELCSVDLDQDGEAELLIG APLFFGEQRGGRVFTYQRRQSLFEM VSELQGDPGYPLGRFGAITALTDIN GDRLTDVAVGAPLEEQGAVYIFNGK PGGLSPQPSQRIQGAQVFPGRWFG RSIHGVKDLGGDRLADVVGAEGRV VVLSSRPVVDVVTLSFSPEEIPVHE VECSYSAREEQKHGVKLCACFRKPL TPQFQGRLLANLSYTLQLDGHMRMS RGLFPDGSHELSGNTSITPDKSCLDF HFHFPIQDLISPINVSLNFSLEEE GTPRDQKVGAMQPIRPSIHTVTKE IPFEKNCGEDKKCEANLTLSSPARS GPLRLMSSASLAVEWTLNSNGEDAY WVRLDLDFPRGLSFRKVEMLQPHS RMPVSCEELTEGSSLLTKLKNVS SPIFKAGQEVSLQVMFTLLNSSWE DFVELNGTVHCENENSSLQEDNSA ATHIPVLYPVNILTKEQENSTLYISFT PKGPKTQQVQHVYQVRIQPSAYDHN MPTLEALVGPWPHSEDPITYTWSV QTDPLVTCHESEDLKRPSSEAEQPCL PGVQFRCPVFRREILIQVTGTVELSK EIKASSTLSCSSLVSVFNSSKHFFHL YGSKASEAQVLVKVDLIHEKEMLHV YVLSGIGGLVLLFLIFLALYKVGFFKR NLKEKMEADGGVPNGSPPEDTDPL AVPGEETKDMGCLEPLRESKDK	False	True	2.41	2.428	1.991	1.744	1.343	5.0	3.075

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