

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	extracellular region
0 P11835	ITB2_MOUSE	Mus musculus	Integrin beta-2	28.201711	NaN	S747;S758;T760;T762	39627609;33300544	MLGLRPSLLLALAGLFFLGSAVSQE CTKYKVSSCRDCIQSGPGCSWCQKL NFTGPGEPDSLRCDDTRAQLLLKGCP ADDIMDPRSIANPEFDQRGQRKQLS PQKVTLYLRPGQAAAFNVTFRRAKG YPIDLYYLMDLSYSMLDDLNNVKKL GGDLLQALNEITESGRIGFGSFVDKT VLPFVNTHTPEKLRNPCPNKEKACQP PFAFRHVLKLTDNSNQFQTEVKGQL ISGNLDAPEGGLDAIMQVAACPEEIG WRNVTRLLVFATDDGFHFAGDGKL GAILTPNDGRCHLEDNMYKRSNEF DYPVSVGQLAHKLSNESNIQIFAVTKK MVKTYEKLTEIIPKSAVGELSDSSN VVQLIKNAYYKLSRVFLDHSTLPDT LKVTYDSFCNSGASSIGKSRGDCDG VQINNPFVTFQVKVMASECIEQSFVI RALGFTDTVTVQVRPQCECQCRDQS REQSLCGGKGVMECGICRCESGYIG KNCECQTQGRSSQELERNCRKDNS SIVCSGLGDCICGQCVCHTS DVPNK EIFGQYCECDNVNCEERYNSQVCGGS DRGSCNCGKCSCKPGYEGSACQCQ RSTTGCLNARLVECSGRGHCQCNR CICDEGYQPPMCEDCPSCGSHCRD NHTSCAECLKFKDFKGFPEKNCSVQCA GMTLQTIPLKKKPKERDSEGCWIT YTLQOKDGRNIYNIHVEDSLECVKG PNVAAIVGGTVVGVVLLIGVLLVIWK ALHTLTDLREYRRFEKEKLSQWNN DNPLFKSATTVMNPKFAES	False	True	2.419	2.656	2.25	2.475	1.613	5.0	3.627