

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	extra regio
P09055	ITB1_MOUSE	Mus musculus	Integrin beta-1	31.948243	NaN	T777;T783;S785;T789	36288343;33300544;34678516;29187734;28528544	MNLQLVSWIGLISLICSVFGQTDKN RCLKANAKSCGECIQAGPNCGWCT NTTFLQEGMPTSARCDLLEALKKKG CQPSDIENPRGSQTIKKNKNTNRS KGMAEKLRPEDITQIQPQQLLLKLR GEPQKFTLKFKAEDYPIDLYLMDL SYSMKDDLENVKS LGTDLNMEMRR ITSDFRIGFGSFVEKTVMPYISTTPAK LRNPCTSEQNCTSPFSYKNVLSLTD RGEFFNELVGQQRISGNLDSPEGGF DAIMQVAVCGSLIGWRNVTRLLVFS TDAGFH FAGDGKLGIVLPNDGQC HLENNVYTM SHYYDYP SIAHLVQKL SENNIQTIFAVTEEFQPVYKELKNLI PKSAVGTLSGNSSNVIQLIADAYNSL SSEVILENSKLPDGVTINYKSYCKNG VNGTGENGRKCSNISGDEVQFEISI TANKCPNKESETIKIKPLGFTEEEV VLQFICKCNCQSHGIPASPKCHEGN GTFECGACRCNEGRVGRHCECSTD EVNSEMDAYCRKENSSEICSNNG ECVCGOCVCRKRDNTNEIYSGKFCE CDNFNCDRSNGLICGGGVCRCRV CECYPNYTGSA CDCSLDTGPCLASN GQICN GRGICEGACKCTDPKFQGP TCETCQTCLGVCAEHKECVQCRAFN KGEEKDTCAQECSHFNLTKVESREK LPQPVQVDPVTHCKEKDIDDCWFYF TYSVNGNNEAIVHV VETPDCPTGPDI IPIVAGVVAGIVLIGLALLLIWKLLMII HDRREFAKFEKEKMNAKWDGTGENP IYKSAVTTVVNPKYEKG	False	True	2.828	3.189	2.717	2.728	2.252	5.0	3.713