

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 Q6P1F6	2ABA_MOUSE	Mus musculus	Serine/threonine-protein phosphatase 2A 55 kDa regulatory subunit B alpha isoform	24.048659	NaN	NaN	33300544;37507081	MAGAGGGNDIQWCFSQVKGAVDDD VAEADIISTVEFNHSGELLATGDKGG RVVIFQQEQENKIQSHSRGEYNVYS TFQSHEPEFDYLSLEIEEKINKIRW LPQKNAAQFLLSTNDKTIKLWKISER DKRPEGYNLKEEDGRYRDPPTVTTL RVPVFRPMDLMVEASPRRIFANAHT YHINSISINSDYETYLSADDLRINLW HLEITDRSFNIVDIKIPANMEELTEVI TAAEFHPNSCNTFVYSSSGTIRLCD MRASALCDRHSKLFEEPEDPSNRSF FSEIISISDVKFSHSGRYMMTRDYL SVKIWDLNMENRPVETYQVHEYLRS KLCSLYENDCIFDKFECCWNGSDSV VMTGSYNFFRMFDRNTKRDITLEA SRENNKPRTVLKPVKCASGKRKCD EISVDSLDFNKKILHTAWHPKENIIA VATTNNLYIFQDKVN	True	False	4.184	2.311	2.357	0.584	0.802	1.154	1.254