

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	ex	
0	Q9CQ65	MTAP_MOUSE	Mus musculus	S-methyl-5'-thioadenosine phosphorylase	30.897006	NaN	NaN	24788674;25153642;37507081;37453647;28528544	MASGSACTAVKIGHGGTGLDDPEIL EGRTEKYVDTFFGKPSDALILGKIKN VDCVLLARHGRQHTIMPSKVNYQA NIWALKEEGCTHVIVTTACGSLREEI QPGDMVIIDQFIDRTSLRPQTFYDGS HCSARGVCHIPMAEPFCPKTREVLIIE TAKKLGRLRCHSKGTIVTIEGPRFSSR AESLIFRTWGADVNNMTTVPEVVLA KEAGICYASIAMATDYDCWKEHEEA VSVDGVLKTMKENANKAKSLLTTI PQIGSMEWSETLRNLKNMAQFSVL PPRH	True	False	4.411	3.248	1.724	1.156	0.589	1.461	1.