

UniprotKB ID	Entry name	organism	full name	oglnacscore	oglnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracellular region
Q8K224	NAT10_MOUSE	Mus musculus	RNA cytidine acetyltransferase	25.161187	NaN	T716;S934;S984;S987	34887587	MNRKKVDNRIRILIENGVAERQSL FVVVGDRGKDQVVLHHMLSKATVK ARPSVLWCYKKEGFPSSHRKRRMR QLQKKIKSGTLNLIKQDDPFELFVAAT NIRYCYYNETHKILGNTFGMCVLQD FEALTPNLLARTVETVEGGGLVILL RTMNSLKQLYTMTMDVHSRYRTEA HQDVVGRFNERFILSLASCKKCLVID DQLDILPISSHVASIEALPPQAPDENL SPAALLELLEKESLQDTQPVGVLDVC CKTLDQAKAVLKFIEGISEKTLRSTV ALTAARGRGKSAALGLAIGAVAFGY SNIFVTSPPDNLHTLFEFVFKGFDA LQYQEHL DYEIVQSLNPEFNKAVIRV NVFREHRQTIQYIHPADAVKLGQAE LVVIDEAAAIPLPVKSLGPLYVFMMA STINGYEGTGRSLSLKLIQQLRQOSA QSQVSTTAENKTTTTARLASARTLH EVSLQESIRYAPGDAVEKWLNDLLC LDCLNITRIVSGCPLPEACELYVNR DTLFCYHKASEVFLQRLMALYVASH YKNSPNDLQMLSDAPAHHLFCLLP VPPTQNALPEVLAVVQVCLEGEISRO SILNSLSRGKKASGDLPVTVSEQFO DPDFGGLSGGRVRIAVHPDYOGMG YGSRALQLLQMYEKGKPCLEEKVL ETPQEIRTVSSEAVSLEEVITPRKDL PPLLLKLNERPAPERLDYLGVSGLTP RLKFWKRAQFVVPVLRQTPNDLTG EHSCIMLKTLADEDEAEQGAWLAAF WKDFRRRFLALLSYQFSTFSPALSL NIIQNRNVAKSALPALGREHLEALFL PYDLKRLEMYSRNMVDYHLIMDLIP AISRLYFLNQLGDLSSLAAQSALLGI GLQHKSVDQLEKEIELPSGQLMGLF NRIIRKVVKLFNDVQEKAIIEEQMVAV KDVVMEPTMKTLSDDLDEAAKEFQ EKHKKEVGKLDMDLSQYVIRGDDE EWNEVLSKAGQNASIVLSKSDKKRK LETKQEPKQSKLKKRDNNRKDMK LKRKK	False	False	2.443	3.774	2.119	1.209	0.792	1.043	0.914