

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	extr regi
Q80UM3	NAA15_MOUSE	Mus musculus	N-alpha-acetyltransferase 15, NatA auxiliary subunit	37.816643	NaN	S302;S537;S588;S854;S855	22645316;39627609	MPAVSLPPKENALFKRILRCYEHKQ YRNLKFKCKQLSNPKFAEHGETLA MKGLTLNLCGKKEEAYELVRRGLRN DLKSHVCWHVYGLLQRSDDKKYDEAI KCYRNALKWWDKDLNQLRDLSSLQI QMRDLEGYRETRYQLLQLRPAQRAS WIGYAIAYHLEDEYEMAAKILEEFK TQQTSPDKVDYSEYSELLLYONQVLR EAGLYREALEHLCTYEKQICDKLAVE ETKGELLQLCRLEDAADVYRGLQE RNPNGNWAYYKGLEKALKPANMLER LKIYEEAWTKYPRGLVPRRLPLNFLS GEKFKCECLDRFLRMNFSKGCPPVF NLTLSLYRDKKVAIVEELVVGYETS LKSCLFNPNDDGKEEPTLLWVQ YYLAQHYDKIGQPSIALEYINTAIEST PTLIELFLVKAKIYKHAGNIKEAARW MDEAQALDTADRFINSKCAKYVLKA NLIKEAEEMCSKFTREGTSAVENLN EMQCMWFQTECAQAYKAMNKFGE ALKKCHEIERHFIEITDDQDFHTYC MRKITLRSYVDLLKLEDVLRQHPFYF KAARIAEYIYKLDHNPPLTDENKEHE ADTANMSDKELKLRNKQRAQKK AQIEEEKNAEKEKQQRNQQKKKD DDDEEIGGPKELIPEKLAKVETPLE EAIKFLTPLKNLVKNKIETHLFAFEIY FRKEKFLMLQSVKRAFADSGHPW LHECMIRLFHVSVCESKDLPETVRTV LKQEMNRLFGATNPKNFNETFLKR NSDSLPHRLSAAKMVYYLDSSSQKR AIELATTLDGSLTNRNLQTCMEVLE ALCDGSLGDCKEAAEAYRASCHKLF PYALAFMPPGYEEDMKITVNGDSSA ETEELANEI	False	False	3.332	3.859	0.932	0.751	1.597	1.412	0.61