

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus
P09405	NUCL_MOUSE	Mus musculus	Nucleolin	40.122178	NaN	S28;S34;S40;S41;S67;T69;T76;T84;T92;T99;T106;T121;S145;S157;S189;S212;T220;S303;S358;T369;S403;T407;S460;S462;S577;S588;S616	33300544;22645316;34887587;37453647;21540332;36288343;39627609;37507081	MVKLAKAGKTHGEAKKMAPPKKEVEEDSEDEEMSEDEDDSSGEEVVIPQKKGKATITPAKKVVVSQTKKAAVPTPAKKAAVTPGKKAVATPAKKNITPAKVIPTPGKKGAAQKALVPTPGKKGAAATPAKGAKNKNAKKEDSDEDED EDEDDSDDEDEDEEDEFEPPIVKGVKPAKAAAPASEDEDEDEDEDD EEDDDEEEDDSEEEVMEITAKGK KTPAKVVPMAKASVAEEEDDEEED EDEDEDDEEEDDEDDDEEEEEEP VKAAPGKRKKEMTKQKEAPEAKKQK VEGSEPTTFFNLFIGNLNPKNKSVNE LKFAISELFAKNDLAVVDVRTGTNR KFGYVDFESAEDLEKALELTGLKVF GNEIKLEKPKGRDSKKVRAARTLLA KNLSFNITEDELKEVFEADAMEIRLVS QDGGKSGIAYIEFKSEADAENLEEK QGAIEDGRSVSLYYTGEKGQRQERT GKTSTWSGESKTLVLSNLSYSATKET LEEVFEKATFIKVPONPHGKPKGYAF IEFASFEDAALNSCNKMEIEGRTI RLELQGSNSRSQPSKTLFVKGLSED TTEETLKESFEGSVRARIVTDRETGS SKGFVDFNSEEAKAAKEAMED GEIDGNKVTLDWAKPKGEGFGGRG GGGRGGFGGRGGGGGGGGFGGRG RGGFGGRGGFRGGGGDFKPKQG KKTKFE	True	True	3.178	5.0	2.217	1.998	1.636