

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	ext reg
B1AQJ2	UBP36_MOUSE	Mus musculus	Ubiquitin carboxyl-terminal hydrolase 36	28.801533	NaN	S429;S463;S547;S578;S663;S678;S709	28528544;24788674	MPIVDKLKEALKPGRKDSAEDGDLG RLLAASAKKVLQLRIEFEPASKSFSY QLESLSKSKYVLLSARAEGASRHRSG DELQARKPGTERVSGSGDGVPAPO KVLFPVERLSLRWERVFRVAGLHN LGNTCFLNSTIQCLTYTPPLANYLLS KEHARSCHQGGFCMLCLMQNHMV QAFANSGNAIKPVSFIRDKKIARHF RFGNQEDAHEFLRYTIDAMQKACL NGYAKLDRQTQATTLVHQIFGGYLR SRVKCSVCKSVSDTYDPYLDIALEIR QAANIVRALELFVKSDVLSGENAYM CAKCKKVPASKRFTIHRTSNVLTLS LKRFAFSGGKITKDVGYPEFLNIRP YMSQSSGDPVMYGLYAVLVHSGYSC HAGHYCYVKASNGQWYQMNDSLV HSSNVKVLNQAYVLFYLRIPGSK KSPEGPVSRVGATLPSRPKVPEHRSK KSPGNGVVPPLMAKRQDSVMMRK LPAPEEVGVPVSRNGSLPGLKLQNG CAPAKTPAGSPSPRLTPTPTHTMPTIL DEPGKKVKKSAPLQSLTTSPTTSQGS PGTGESRSQRPGSWASRDITFSTSPK LLARAITNGHRLKGEQGVLDLEKGD SSSSSPEHSASSDPAKAPQTAESRAA HACDSQGTNCPTAGHPKALLNGVD AKMVKLKSALSTTTEPTSLMSPPP AKKLALSAKASTLRRATGNDIGSPS PSAFCDLTSPMKATHPVVASTGPVS KTRTAAPAPRPSTHPHSASLSSSSAK PLGTSEPOQSCRPSAWTPLPQVNGHF TSHLHQLPEASEALHSPSKRKKKTP NGDPQRLGIDTLLPQCLRGAPAAAR RKRKKRCSEGEGATAPKQEGQFQD QSWSSGSQKEEGTQPQVNGHQVSH ILDSYHVSSRKRKRKRSEGLSQEAT PSQDLIQHSCSPVDHSEPEARTELQ KKKKKRKRKRKPEPQDEESKHPG DQRSPRPSVTPVPALSVNGHLPSDC LGLGQAPLVTVNRDQEPDVOALL QDSSDKAYGKVLTVWGEPSAISQD AIKDSRLARTQTVVDDWDEEFDRGK EKKIKFKREKRNFNFAFKLQSRRR NFWSVTHPAKVASLSYRR	None	None	None	None	None	None	None	None	

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