

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	extract region
Q70EL4	UBP43_HUMAN	Homo sapiens	Ubiquitin carboxyl-terminal hydrolase 43	23.347461	T218;S1111	S969;S1041	39894887;30379171;38253038	MDLGPDAAGGGPLAPRRRRSLR RLFSRFLALGSRSRPGDSPRRPQPG HCDGDGEGGFACAPGPVPAAPGSPG EERPPGPQQLQPLAGDGARPPGAQ GLKNHGNTCFMNAVVCLSNTDLL AEFLALGRYRAAPGRAEVTEQLAALV RALWTRREYTPQLSAEFKNAVSKYGS QFQGN SQHDALEFLLWLLDRVHED LEGSSRGVSEKLPPEATKTSENCLS PSAQLPLGQSFVQSHFQAQYRSSLT CPHCLKQSNTFDPFLCVSLPIPLRQT RFLSVTLVFPKSRQFLRVGLAVPIL STVAALRKMVAEEGGVPADEVILVEL YPSGFQRSFFDEEDLNTIAEGDNVY AFQVPPSPSQGTL SAHPLGLSASPRL AAREGQRFSLSLHSESKVLILFCNLV GSGQQASRFPPFLIREDRAWVAQ LQQSILSKVRHLMKSEAPVQNLGSL FSIRVVGLSVACSYLSPKDSRPLCHW AVDRVLHLRRPGGPPHVKLAVEWDS SVKERLFGSLQEERAQDADSVWQQ QQAHQHSCITLDECFQFYTKEEQL AQDDAWKCPHCQVLQQGMVKLSL WTLDPDILIIHLKRFCQVGERRNKLST LVKFPPLSGLNMAPHVAQRSTSP EAG LGPWPSWKQPDCLPTSYPLDFLYDL YAVCNHHGNLQGGHYTAYCRNSLD GGWYSYDDSTVEPLREDEVNTRGAY ILFYQRNSIPPWSASSMRGSTSSS LSDHWLLRLGSHAGSTRGSLLSWSS APCPSLPQVDPSPIFTNSLCNQEKGK LEPRRLVRGVKGRSISMKAPTT SRAK QGPFKTMPLRWSFGSKEKPPGASVE LVEYLESRRRPRSTSQSIVSLLTGTA GEDEKSASPRSNVALPANSEDGGRA IERGPAGVPCPSAQP NHCLAPGNSD GPNTARKLKENAGQDIKLPKFDLP LTVMPVSEHEKPARPEGQKAMNWK ESFQMGSKSSPPSPYMGFSGNSKDS RRGTSELDRLPQGLTLLLRVFRKKE NRRNERAEVSPQVPPVSLVSGGLSP AMDGQAPGSPALRIPEGLARCLGS RLERDVWSAPSSLRLPRKASRAPRG SALGMSQRTVPGEQASYGTFQRVKY HTLSLGRKKTLPESFF	True	False	0.844	4.379	0.422	False	False	0.281	0.281

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