

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	extracellular region
A6H6E2	MMRN2_MOUSE	Mus musculus	Multimerin-2	25.019509	T903	NaN	34418053	MIPTLLLGFGVYLSWGLLGSWAQDP GTKFSHLNRPGMPEGWRLGAEDTS RDPIRRNWCPYQKSRLVTFVAACKT EKFLVHSQQPCPQGAPDCQGVVVM YRVAQKPVYVQVQKVLISVDWRCCP GFQGPDCQDHNPTANPEPTEPSGKL QETWDSMDGFELGHPVPEFNEIKVP QEQQENLLQNLQNDAAQSVEDGFPG SWEAPPSNLTDEMTEANLTFEFPG RTSEHLLQPHIDAFKKAHFSPIWKN FNDSLHLSQAIRNLSLDVEANHQA IKMIQEGTVARADFQELGAKFEAKV QQNSORLGQLWQDVEDQLHAQRRS VHHALSDVQAEVSTKLKQLVKAQEL PGANGSLVMASAAAAARPEPESLQA RLGQLQRNLSALHMVTSQREELQ STLKNMDSVLKQHAEEIKELYSED ETFDQISKVERQVEELLVNHTGLRE LRVILMEKSLIMEENKEIEIRQLLEL NLTLOHLHAGHADLIKVKDCNCOR VNSDQVVAPEGHRDVMHTLEETQV SLDEQHQLDGSSLQALQSTVDAMSS AMDAYRGEGERARAERARIRSQLRA LDHAVEALKTAANGTRKEIRLLHGS FTALLEDALRHQAVLAALFGEEMID EMSEEAPRPLPDYEQIRLALQDAAS GLQEQAIGWEDLATRVEALEKAAGG FVEQHPQLAEGLEPSHDSGREEEA MTLAELEQEIRRLSSDVKQIGOCCE ASWAASLNSSLEDLHSMMLDTQHG LRQHRQLFHNLFQNFQGLVASNISL DLGKLQAMLKDKKKQPRGPGESR KRDKKQVVMSTDAHAKGLELWETG SPVAFYAGSSEGATALQMKFNNTSI NVGSSYFPEHGYFRAPKRGVYLFV SITFGPGPMGQLVFEHHRVPVYS TEQRGGSTATTFAMVELQKGERAW FELIQGSATKGSQPGTAFGGFLMFK T	None	None	None	None	None	None	None	None	

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