	UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochndrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracellular region
0	O60732	MAGC1_HUMAN	Homo sapiens	Melanoma-associated antigen C1	22.636511	NaN	S63;S207;S382;S1063	34725712	MGDKDMPTAGMPSLLQSSSESPQS CPEGEDSQSPLQIPQSSPESDDTLYP LQSPQSRSEGEDSSDPLQRPPEGKD SQSPLQIPQSSPEGDDTQSPLQNSQ SSPEGKDSLSPLEISQSPPEGEDVQS PLQNPASSFFSSALLSIFQSSPESTQ SPFEGFPQSVLQIPVSAASSSTLVSIF QSSPESTQSPFEGFPQSPLQIPVSRS FSSTLLSIFQSSPERTQSTFEGFAQS PLQIPVSPSSSSTLLSLFQSFSERTQS TFEGFAQSSLQIPVSPSFSSTLVSLF QSSPERTQSTFEGFPQSPLQIPVSSS SSSTLLSLFQSSPERTHSTFEGFPQS LLQIPMTSSFSSTLLSIFQSSPESAQS TFEGFPQSPLQIPGSPSFSSTLLSLFQ SSPERTHSTFEGFPQSPLQIPMTSSF SSTLLSILQSSPESAQSAFEGFPQSP LQIPVSSSFSYTLLSLFQSSPERTHST FEGFPQSPLQIPVSSSSSSSTLLSLFQ SSPECTQSTFEGFPQSPLQIPQSPPE GENTHSPLQIVPSLPEWEDSLSPHY FPQSPPQGEDSLSPHYFPQSPPQGE DSLSPHYFPQSPQGEDSLSPHYFPQ SPPQGEDSMSPLYFPQSPLQGEEFQ SSLQSPVSICSSSTPSSLPQSFPESSQ SPPEGPVQSPLHSPQSPPEGMHSQS PLQSPESAPEGEDSLSPLQIPQSPLE GEDSLSSLHFPQSPPEWEDSLSPLH FPQFPPQGEDFQSSLQSPVSICSSST SLSLPQSFPESPQSPPEGPAQSPLQR PVSSFFSYTLASLLQSSHESPQSPPE GPAQSPLQSPVSSFPSSTSSSLSQSS PVSSFPSSTSSSLSKSSPESPLQSPVI SFSSSTSLSPFSEESSSPVDEYTSSS DTLLESDSLTDSESLIESEPLFTYTLD EKVDELARFLLLKYQVKQPITKAEML TNVISRYTGYFPVIFRKAREFIEILFGI SLREVDPDDSYVFVNTLDLTSEGCL SDEQGMSQNRLLILILSIIFIKGTYAS EEVIWDVLSGIGVRAGREHFAFGEP RELLTKVWVQEHYLEYREVPNSSPP RYEFLWGPRAHSEVIKRKVVEFLAM LKNTVPITFPSSYKDALKDVEERAQA IIDTTDDSTATESASSSVMSPSFSSE	None	None	None	None	None	None	None	None	None