	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	extracel region
0	Q8TDZ2	MICA1_HUMAN	Homo sapiens	[F-actin]-monooxygenase MICAL1	26.21016	T498	T475;S617;S872;S875;S876;S1057	29351928	MASPTSTNPAHAHFESFLQAQLCQD VLSSFQELCGALGLEPGGGLPQYHKI KDQLNYWSAKSLWTKLDKRAGQPV YQQGRACTSTKCLVVGAGPCGLRVA VELALLGARVVLVEKRTKFSRHNVL HLWPFTIHDLRALGAKKFYGRFCTG TLDHISIRQLQLLLLKVALLLGVEIH WGVTFTGLQPPPRKGSGWRAQLQP NPPAQLANYEFDVLISAAGGKFVPE GFKVREMRGKLAIGITANFVNGRTV EETQVPEISGVARIYNQSFFQSLLKA TGIDLENIVYYKDDTHYFVMTAKKQ CLLRLGVLRQDWPDTNRLLGSANVV PEALQRFTRAAADFATHGKLGKLEF AQDAHGQPDVSAFDFTSMMRAESS ARVQEKHGARLLLGLVGDCLVEPFW PLGTGVARGFLAAFDAAWMVKRWA EGAESLEVLAERESLYQLLSQTSPEN MHRNVAQYGLDPATRYPNLNLRAV TPNQVRDLYDVLAKEPVQRNNDKT DTGMPATGSAGTQEELLRWCQEQT AGYPGVHVSDLSSSWADGLALCALV YRLQPGLLEPSELQGLGALEATAWA LKVAENELGITPVVSAQAVVAGSDPL GLIAYLSHFHSAFKSMAHSPGPVSQ ASPGTSSAVLFLSKLQRTLQRSRAKE NAEDAGGKKLRLEMEAETPSTEVPP DPEPGVPLTPPSQHQEAGAGDLCAL CGEHLYVLERLCVNGHFFHRSCFRC HTCEATLWPGGYEQHPGDGHFYCL QHLPQTDHKAEGSDRGPESPELPTP SENSMPPGLSTPTASQEGAGPVPDP SQPTRRQIRLSSPERQRLSSLNLTPD PEMEPPPKPPRSCSALARHALESSF VGWGLPVQSPQALVAMEKEEKESPF SSEEEEEDVPLDSDVEQALQTFAKT SGTMNNYPTWRRTLLRRAKEEEMK RFCKAQTIQRRLNEIEAALRELEAEG VKLELALRRQSSSPEQQKKLWVGQL LQLVDKKNSLVAEEAELMITVQELN LEEKQWQLDQELRGYMNREENLKT AADRQAEDQVLRKLVDLVNQRDALI RFQEERRLSELALGTGAQG	None	None	None	None	None	None	None	None	None