	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	extra regio
0	P27448	MARK3_HUMAN	Homo sapiens	MAP/microtubule affinity-regulating kinase 3	26.75988	NaN	S42;T211;S368;S374;S376;S380;S383;S400;S419;S469;S540;S543;T549;T564;S583;S598;S601;S643;S384;S407	29485866	MSTRTPLPTVNERDTENHTSHGDG RQEVTSRTSRSGARCRNSIASCADE QPHIGNYRLLKTIGKGNFAKVKLAR HILTGREVAIKIIDKTQLNPTSLQKLF REVRIMKILNHPNIVKLFEVIETEKTL YLIMEYASGGEVFDYLVAHGRMKEK EARSKFRQIVSAVQYCHQKRIVHRD LKAENLLLDADMNIKIADFGFSNEF TVGGKLDTFCGSPPYAAPELFQGKK YDGPEVDVWSLGVILYTLVSGSLPFD GQNLKELRERVLRGKYRIPFYMSTD CENLLKRFLVLNPIKRGTLEQIMKD RWINAGHEEDELKPFVEPELDISDQ KRIDIMVGMGYSQEEIQESLSKMKY DEITATYLLLGRKSSELDASDSSSSS NLSLAKVRPSSDLNNSTGQSPHHKV QRSVSSSQKQRRYSDHAGPAIPSVV AYPKRSQTSTADSDLKEDGISSRKSS GSAVGGKGIAPASPMLGNASNPNKA DIPERKKSSTVPSSNTASGGMTRRN TYVCSERTTADRHSVIQNGKENSTIP DQRTPVASTHSISSAATPDRIRFPRG TASRSTFHGQPRERRTATYNGPPAS PSLSHEATPLSQTRSRGSTNLFSKLT SKLTRRNMSFRFIKRLPTEYERNGR YEGSSRNVSAEQKDENKEAKPRSLR FTWSMKTTSSMDPGDMMREIRKVL DANNCDYEQRERFLLFCVHGDGHA ENLVQWEMEVCKLPRLSLNGVRFK RISGTSIAFKNIASKIANELKL	None	None	None	None	None	None	None	None	None