	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	M9PIF1	M9PIF1_DROME	Drosophila melanogaster	NaN	28.06426	T1069	NaN	40245499	MWIQPKELLLPSAFWIAEMHSRYFV LQKRRGHGESRGFGSMLVGTYDSV MNTKPAPYRILHQTPSSEVSYEIAIGI TQDEIVKDWEWLKANLFKVLDEME NEDEVTNFTICKIQSLYTQNNQDDT GESADFKVMKSKFRQIFKMPEEERL VISYSATYVKNKIPRQGQLYISLNHV CFYSYMLGQEIKRIIRFAELEDISRNA NTIYLKTTNNMTYNFTMLFNASDAH LLIEQLNKMAIQQLIHDPDSPVVDH DTSNFSRLGSKTSKKPVLLRDLTAR QKSEEFRIYFRLPQSEIIDGKIKANIW TPYSKRFNSGFIYLSPNFFCFRSDVK DLVSVVIPMKTIKSVEKKDDGQQRF ENQIVIITSENVPFMFAHIVDRAVLIS KITDLLARVHVPLSRERAKYDISWSK QTALMNTFKTQFSAEIIQKQEEKMV RWEAHFRDFGRGIGMFRTTDVINLI VEGIPDKLRQEIWLIFSGAIHDKEMN PGLYEDLVEKAACIKNCFAHDEIDRD LPRSLPEHPAFQSTDGIGALRRVLQA YALRNPQVGYCQAMNIVSSVFLLFC DEENAFWMLASLCENLLPDYYKDK VVGAQIDQGVLNELVETHLPDLHGH LEQLGVIKMISISWFLTIFMSVISYES SLHILDCFFYEGAKIIFMISLQIIEWN RDKLLICQDDGEAMLVLQNYLEGVY NPEYQVPPTTDKRKMERKVQTQTVQ TLIHEAYTKFGEDITQQRIEELRNKH RRLTMRQFDIDNEKTIVKAYVQNPY FNRSELHMLLTIIREEKHALKSLQQ QQQKVQCPLSETPQLVPQSTSRPIQ DAGASGGRYEAYSVSYEVFHTLFTEL TPWRKCVSVDIGEKLFRLTDKKGTG VLDFGQLINALGLVCSNKNMEKLKL LYVLHLPPLLSKAEIERSRRPRPRTK DDAEEAFEAEDFFDNDASESMEALP SPSDHNFDADDFALISATQHLHNLA GISGNTFMDLSRTPNLSVNSNTSSL AHRSSTFYVDLPGFQDLGAARITPPE AGVGADVVAGAASSAAADHMLLNV ETISNFSQISDLVAATRLERVDSNAT DLRSLGSLGYLLDQPDEGASNSQHS IPHMRKENFQLLWRSIIEIMGLVQD EEMQRAYENLLELGNSNIKKEPSLE SFTQLNMGGDEQPDSNGNPTTPAP EPIGTTRLFVALEEELRQASGKSRTT TTSSGDSSSTNISESWHISITQFIATV LTVNSIVRGFQTPIQISEQIEQLQKKR RKCLSTNY	NaN	NaN	NaN	NaN	NaN	NaN	NaN	NaN	None