	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	Q24318	TFDP_DROME	Drosophila melanogaster	Transcription factor Dp	28.06426	S90;T94;T98;T99	S398	40245499	MAHSTGGTVKTDEVNFFFRNEHGQ ISKMLKPAQNKSEMEGGKPVAVVYA TGSSARNSGSASGIGNVGRMGAFSQ MGSQGQFIRLQDNGLSIPKTEAGTT YTTVSAQKTSGAGSGHYDLPLKGDR YVKFTPNPIKMKSKLHAIQSNSLHS MSASSSSVQRKRKPDKAGKGLRHFS MKVCEKVEEKGKTTYNEVADDLVSE EMKNNAYDNNCDQKNIRRRVYDAL NVLMAINVISKDKKEIRWIGLPANST ETFLALEEENCQRRERIKQKNEMLR EMIMQHVAFKGLVERNKRNESQGV VPSPNASIQLPFIIVNTHKSTKINCSV TNDKSEYIFKFDKTFEMHDDIEVLK RMGFLLGLDKGECTPENIERVKSWV PPNLAKYVEAYGTGKTGENMYESDD EDNEFNGYLESANESQGFAQHSAQ HTTDGEFKLEMDDDELDDDID	True	False	1.406	5.0	False	False	False	False	0.798