	UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochndrion	endoplasmic reticulum	golgi apparatus
0	Q8C156	CND2_MOUSE	Mus musculus	Condensin complex subunit 2	35.186263	T93	S12;S25;T43;T46;S67;S75;S78;S83;S91;T182;S195;S227;S425;S489;T587	36852467;28528544;28135057;25153642;39627609	MRIPRSETMNSSFLKARGQQDVLSSPLERVPPASRPGKAPLGTPKTPVLEDFPQNDDEKERMQRRRSRVFDLQFSTDSIHLASPNRNIDVSTTISKFTNTQITEHYSTCIKLSSENKITTKNAFGLHLIDFMSEILKQKDAEPTNFKVAAGTLDASTKIYAVRVDAVHADVYRVLGGLGKDTPPQGEESHSGDGSTLETERTKKPAKPKKKQSCKTIEQNLSNINVSEADGKCAVDPMFQKTAASFDECSTTGVFLSTLHCQDYRSELLFPSDMQTLSSGEPLELPDLGFVDMTDLEASLQQCVEDRPLCPSLAGFQFTKWDSETHNESVSALVDKFKKNDQVFDINAEAEDDEEDVPDGPLVEDFVDNDEPDLSAAGDHEEFRSWKELCQVQSNQEEVISLEDRDIQVMCSFLSMKPGEYSYFSPRTMKMWAGPDHWRFRPRPKQDATSCTEHKKKSAKKDFEINFDDDIDFDAYFQKTKAATILTKSTLENQNWKATTLPTDFHYETDNLIQLHLKPGKRSLKMDQDQKAKTEHYEEIEDYDYNNPNDTSNYCPGLQAADSDYEEADDLFADPVGTLDLESDPKTTQENGHISPENQGVDITTYQELNLVAEPQKVNKIEIHYAKTAKKMDMKKLKQSMWSLLTKFSRKEADTEANHTESGQEGAPEEVADEKKLSGLTKDLQTRLPPLMAQNLSIPLAFACLLHLANEKNLKLEGTEDLSDVLVMQGD	True	False	2.901	5.0	0.919	0.72	False