	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	Q9VF20	Q9VF20_DROME	Drosophila melanogaster	ATP-binding cassette sub-family B member 6	28.06426	T55;T58;S63	NaN	40245499	MLYCPPNVTLSEVWTQHGISHCFM DTVGPAVYGGFLLLFGSIQLLMYRKY ATRITDPTQISKSRLFAMQLFLLLLLP VLALLRFLMNARIYPDSAVYGYMIFS TCVVCFSYPFSICLILKERYYQLPSMP TRGHGLVLLLFWTLAFINESLAFINL RHEDWWFHLKTNKDQIEMGLFVTR FLCSLLIFVLGLKAPGIMAPYNPHQR LDNDTANESQGNVQTGSAFRNGWR KLRTVFPYLWPKKNIALQIAVIVCIILL LAGRVIKLFLPIYRKKLVDSLTIAPIVF RWDFVLIYVALSFLQGGGTGSMGLF NNLRTFLWIRVQQYTTREIEIELFRH LHQLSLRWHLQRKTGEVLRVMDRG TDSINNLLNYIVFSIAPTILDLLVAVA YFVYAFNWWFGLIVFLTMFLYIASTI AITEWRTKYQRRMNLADNEQRARS VDSLLNFETVKYYGAEHYEVDCYRE AILKYQKEEFLSMLTLNMLNTAQNII LCLGLLAGSLLCVYLVVHHQTLTVG DFVLFSTYLMELYMPLNWFGTYYRA IQKNFVDMENMFDLLKEEEEIVDAP GCSPLLTAGGGIEFSNVTFGYSPEKI VLRNVSFTVPAGKTVAIVGPSGAGKS TIMRLLFRFYDVQTGAILIDGQNIKL VQQQSLRKAIGVVPQDTVLFNNTIFY NIEYAKLGASDEAVYEAARAADIHER ILGFPEKYETKVGERGLRLSGGEKQR VAIARTLLKAPIIVLLDEATSALDTHT ERNIQAALARVCANRTTIIVAHRLSTI IHADEILVLQQGSIAERGRHEELVLR EDGIYADMWQQQLKNLDAEQSGGS DNGDASAESGSEKRRAGGAGGPSGT GTGGAHFRAGHAHGGAR	False	False	1.529	1.059	1.699	0.82	0.712	3.551	1.108