	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	Q6ULP2	AFTIN_HUMAN	Homo sapiens	Aftiphilin	26.249992	T450;T452	S151;S395;S518;T617	23301498;29351928	MEPDIIRMYSSSPPPLDNGAEDDDD DEFGEFGGFSEVSPSGVGFVDFDTP DYTRPKEEFVPSNHFMPIHEFSENV DSLTSFKSIKNGNDKDITAELSAPVK GQSDVLLSTTSKEIISSEMLATSIDG MERPGNLNKVVEQRQNVGTLESFS PGDFRTNMNVVHQNKQLESCNGEK PPCLEILTNGFAVLETVNPQGTDDLD NVADSKGRKPLSTHSTEYNLDSVPS PAEEFADFATFSKKERIQLEEIECAV LNDREALTIRENNKINRVNELNSVK EVALGRSLDNKGDTDGEDQVCVSEI SIVTNRGFSVEKQGLPTLQQDEFLQ SGVQSKAWSLVDSADNSEAIRREQC KTEEKLDLLTSKCAHLCMDSVKTSD DEVGSPKEESRKFTNFQSPNIDPTE ENDLDDSLSVKNGDSSNDFVTCNDI NEDDFGDFGDFGSASGSTPPFVTGT QDSMSDATFEESSEHFPHFSEPGDD FGEFGDINAVSCQEETILTKSDLKQT SDNLSEECQLARKSSGTGTEPVAKL KNGQEGEIGHFDSVPNIQDDCNGF QDSDDFADFSSAGPSQVVDWNAFE DEQKDSCSWAAFGDQQATESHHRK EAWQSHRTDENIDTPGTPKTHSVPS ATSKGAVASGHLQESATSVQTALLN RLERIFEACFPSILVPDAEEEVTSLK HLLETSTLPIKTREALPESGELLDVW TELQDIHDAHGLRYQWGGSHSNKK LLSSLGIDTRNILFTGNKKQPVIVPM YAAGLGMLEPTKEPLKPLSAAEKIAS IGQTATMSPDMNTCTSDQFQESLPP VQFDWSSSGLTNPLDASGGSTLLNL DFFGPVDDSSSSSSTTIPGVDPELYE LTTSKLEISTSSLKVTDAFARLMSTV EKTSTSTRKPKREEHLSEEAIKVIAG LPDLTFMHAKVLMFPATLTPSTSSQ EKADG	None	None	None	None	None	None	None	None	None