	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	Q7JRJ1	EAF_DROME	Drosophila melanogaster	Ell-associated factor Eaf	28.06426	T119	S194	40245499	MMMTKQKNSLAERLNIGEEVRELK LGATFNPKNTSTAFHTIKYDFKPASV DTSRMASVDVGSNNQVTVTVPNSE SSGVPHTVYKGNQREYAKECLMIYD KETGAITIEKLNHNIQVKKTRNEVTN KSVQLPGQNMGQPHNQGANGAAPV AVPVPGQGSGTAPKMENSTMRISTK TKVSTGSRRNNIIDFKPRNSPMQQN SPSRPVPVHRSPQSAPAWDANNAQ QTLPSIPLITDDDDFGLRAALHNSGH GNTSGTAAGQPDFGSTSSSTHIGKQ RQAPPHGHGKRQQMHQRLSPPMA QQQQPSNYGRGYNGGHNHAQQQQ HHQRNSPQQQRPSAYGHGNSMPID VDSSREHELTSQSVAQAAAALEQQI GGALSASSSSSESDSSDSDSGSDSD DSTEDDRSTQGQQQDHQQQQQQQ HQVYQNHKHTQQQVAQQHHNQLP NLGLGSISPAYGSSHQQQHQQQML PHQQKQKQQSGIYASNGGFPNDFL QNDLQLSSNSSDDDDD	True	False	1.484	4.594	0.538	False	0.516	False	False