

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracellular region	
0	Q8N1L9	BATF2_HUMAN	Homo sapiens	Basic leucine zipper transcriptional factor ATF-like 2	26.75988	NaN	NaN	29485866	MHLCGGNLLTQDPKEQQRQLKK QKNRAAAQRSRQKHTDKADALHQQ HESLEKDNLALRKEIQSLQAEALWW SRTLHVHERLCPMDCASCAPGLLG CWDQAEGLLGPGPQGQHGCREQLE LFQTPGSCYPAQPLSPGPQPHDSPSL LQCPLPSLSLGPVVAEPPVQLSPSP LLFASHTGSSLQGSSSKLSALQPSLT AQTAPPQPLELEHPTRGKLGSSPDN PSSALGLARLQSRHHPALSAATWQ GLVVDPSHPHLLAFPLLSAQVHF	True	False	1.391	4.495	1.308	1.186	0.616	1.266	1.741