

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus
Q3U0V1	FUBP2_MOUSE	Mus musculus	Far upstream element-binding protein 2	47.67849	T95	T101;S126;S130;S182;S185;S194;S275;S481	21540332;36852467;34887587;24788674;22645316;25153642;36288343;28135057;28528544;35822049	MSDYNTGGPPGPPPPAGGGGAA GAGGGPPGPPGAGDRGGGGPGGG GPGGGGASGGPSQPPGGGGPGIRKD AFADAVQRARQIAAKIGGDAATTVN NNTPDFGFGGQKRQLEDGDQPSK KLASQGDSIGSQLGPIHPPRTSMTE EYRVPDGMVGLIIGRGGEQINKIQDD SGCKVQISPDSSGGLPERSVSLTGAPE SVQKAKMMLDDIVSRGRGPPGQF HDNANGGQNGTVQEIMIPAGKAGL VIGKGGETIKQLQERAGVKMILIQDG SQNTNVDKPLRIIGDPYKVVQACEM VMDILRERDQGGFGDRNEYGSRVG GGIDVPVPRHSVGVVIGRSGEMIKKI QNDAGVRIQFKQDDGTGPEKIAHIM GPPDRCEHAARIINDLLQSLRSGPPG PPGAPGMPPGGRGRGRQGNWGP GGEMTFSIPTHKCGLVIGRGGENVK AINQQTGAFVEISRQLPPNGDPNFK LFVIRGSPQQIDHAKQLIEEKIEGPLC PVGPGGGPGPAGPMGPFNPGPFN QGPPGAPPHAGGPPPHQYPPQGWG NTYPQWQPPAPHPNKAATAATDP NAAWAAYSHYYQPPGPVPGPAPA PAAPPAQGEPPQPPPTGQSDYTKAW EEYYKIQQQPQQPGAPPQDYTKA WEEYYKQAQVATGGGPGAPPGSQP DYSAAWAEYRQQAAYGQTPGPGG PQPPPTQQGQQQASGNCHPPPPFFS FQPPATVHPALVGSAGNPFPCGVCP	True	False	4.382	4.068	1.626	1.531	1.042