

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
Q9H1B7	I2BPL_HUMAN	Homo sapiens	Probable E3 ubiquitin-protein ligase IRF2BPL	27.507895	S169;S179;S182;S183;S184;T186;T302;S306;T308;S314;S319	S69;S215;S519;S547;S639;S657;S658;S659;S662	3626323;30379171;33214551;34105348;30059200;31492838;32119511;32574038;29351928;29237092;34019948;23301498	MSAAQVSSRRQSCYLCDLPRMPW AMIWDFSEPVCRGCVNYEGADRIEF VIETARQLKRAHGCFQDGRSPGPPP PVGKTVALSAAKEAAAAAAAAAAAA AAQQQQQQQQQQQQQQQQQQQQ QQQLNHVDGSSKPAVLAAPSGLER YGLSAAAAAAAAAAAAAVEQRSFEYF PPPVSLGSSSHARLPNGLGGPNGF PKPTPEEGPPELNROSPNSSSAAASV ASRRGTHGGLVTGLPNPGGGGGPQL TVPPNLLPQTLLNGPASA AVLPPPPP HALGSRGPPTPAPPGAPGGPACLGG TPGVSATSSSASSSTSSSVAEVGVGA GGKRPGSVSSTDQERELKEKQRNAE ALAESESLRNRAEEWASKPKMVRD TLLTLGCTPYEVRFKDDHSLGRVF AFDAVSKPGMDYELKLFIEYPTGSG NVYSSASGVAKQMYQDCMKDFGRG LSSGFKYLEYEKKHSGDWRLLDL LPEAVRFFKEGVPADMLPQPYLDA SCPMLPTALVSLSRAPSAPPGTGALP PAAPSGRGAASLRKRKASPEPPDSA EGALKLGEEQQRQWMANQSEALK LTMSAGGFAAPGHAAGGPPPPPPPL GPHSNRTTPESAPQNGPSPMAAL MSVADTLGTAHSPKDGSSVHSTTAS ARRNSSPVSPASVPGQRRLASRNG DLNLQVAPPPSAHPGMDQVHPQNI PDSPMANSGLCCTICHERLEDTHF VQCPSPVSHKFCFPCSRESIKAQGAT GEVYCPSGEKCPLVGSNVPWAFMQ GEIATILAGDVKKKERDP

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