

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondr
Q7Z417	NUFP2_HUMAN	Homo sapiens	FMR1-interacting protein NUFIP2	24.510444	S374;S376;S379;S380;S381;S382;S384;S385;T387;T437;S546	T87;S112;S113;S212;S214;T218;T219;T220;S304;S376;T571;S572;S592;S608;S629;T633;S637;S652;S655;S692	20305658;36240223;30379171;22661428;35132862;29351928;33214551;35138101;34019948;21740066;35254053;30620550;32119511;37217939;27114449;38665916	MEEKPGQPQPHHHSHHHPHHHP QQQQQPHHHHHYFYNHSHNHH HHHHHQPHOYLQHGAEQSPKAQP KPLKHEQKHTLQQHQETPKKKTGY GELNGNAGEREISLKNLSSDEATNPI SRVLNGNQVVDTSLKQTVKANTFG KAGIKTKNFIQKNSMDKKNGKSYEN KSGENQSVDKSDTIPIPNGVVTNNS GYITNGYMGKADNDGSGSESGYTT PKKRKARRNSAKGCENLNIVQDKIM QQETSVPTLKQGLETFKPDYSEQKG NRVDGSKPIWKYETGPGGTSRQKPA VGDMLRKSSDSKPGVSSKKFDDRPK GKHASAVASKEDSWTLFKPPVFPV DNSSAKIVPKISYASKVKENLNKTIQ NSSVSPTSSSSSSSTGETQTQSSSR LSQVPMASALKSVTSANFNGPVLG TDGNVYPPGGQPLLTTAANTLTPISS GTDSVLQDMSLTSAAVEQIKTSLFIY PSNMOTMLLSTAQVDLPSQTDQQN LGDIFQNWGLSFINEPSAGPETVT GKSSEHKVMEVTFQGEYPATLVSQG AEIIPSGTEHPVFPKAYELEKRTSPQV LGSILKSGTTSSESGALSLEPSHIGDL QKADTSSQGALVFLSKDYEIESQNPL ASPTNTLLGSAKEQRYQRGLERNDS WGSFDLRAAIVYHTKEMESIWNLQK QDPKRUIYNEAMDSPDQ	True	False	4.617	5.0	0.951