

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membra
Q6ZN30	BNC2_HUMAN	Homo sapiens	Zinc finger protein basonuclin-2	29.431777	S275;T514;S515;T525	S561	31492838;30620550;35254053	MAHLGPTPPPHSLNYKSEDRLSEQD WPAYFKVPCCGVDTSQIESEEAQVD VRERETQRDREPKRARDLTLRDSCT DNSMQFGTRTTTAEPPGFMGTWQNA DTNLLFRMSQQAIRCTLVNCTCECF QPGKINLRDTCQCKHGWVAHALDK LSTQHLYHPTQVEIVQSNVVFDISSL MLYGTQAVPVRLKILLDRLFSVLKQE EVLHILHGLGWLRLRDYVVRGILQDA AGKVLDRWAIMSREEEITLQOFLRF GETKSIVELMAIQEKEGQAVAVPSSK TDSDIRTFIESNNRTRSPSLLAHLEN SNPSSIHFFENIPNSLAFLLPFQYIN PVSAPLLGLPPNGLLLEQPLRLREP SLSTQNEYNESSESEVSPTPYKNDQ TPNRNALTSTNVEPKTEPACVSPIQ NSAPVSDLTKTEHPKSSFRHRMRR MGSASRKGRVFCNACGKTFYDKGT LKIHYNAVHLKIKHRCTIEGCNMVF SSLRSRNRHSANPNRHLHMPMLRN NRDKDLIRATSGAATPVIAS TKSNLA LTSRGRPPMGFTTTPPLDVLQNPPLPS QLVFSGLKTVQPVPPFYRSLTTPGE MVSPTSLPTSPIPTSGTIEQHPPPP SEPVVPAVMMATHEPSADLAPKKKP RKSSMPVKIEKEIITADEFDDEDDD PNDGGAVVNDMSHDNHCHSQEEM SPGMSVKDFSKHNRTRCISRTEIRR ADSMTSEDQEPERDYENESESEPK LGEESMEGDEHIHSEVSEKVLNNS ERPDENHSEPSHQDVIKVEEFTDP TYDMFYMSQYGLYNGGGASMAALH ESFTSSLNYGSPQKFSPEGDLCSSPD PKICYCKKSKFKSSYSVKLHYRNVHL KEMHVCTVAGCNAAFPSRRSRDRH SANINLHRKLLTKELDDMGLDSSQP SLSKDLRDEFVLKIYGAQHPMGLDV REDASSPAGTEDSHLNGYGRGMAE DYMVLDLSTTSSLQSSSIHSSRES AGSDEGILLDDIDGASDSESAHKA EAPALPGSLGAEVSGSLMFSLSLGS NGGIMCNICHKMYSNKGTLRVHYK TVHLREMHKCKVPGCNMMFSSVRS RNRHSQPNLHKNIPFTSVD	True	False	2.435	4.818	1.295	0.953	0.625	2.243