	UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochndrion	endoplasmic reticulum	golgi apparatus	plasma membran
0	Q9HCK1	ZDBF2_HUMAN	Homo sapiens	DBF4-type zinc finger-containing protein 2	22.971194	T40;S41;S266;S1870	S539	30379171;23301498;29237092	MQKRQGYCSYCRVQYNNLEQHLFS AQHRSLTRQSRRQICTSSLMERFLQ DVLQHHPYHCQESSSTQDETHVNT GSSSEVVHLDDAFSEEEEEDEDKVE DEDATEERPSEVSEPIEELHSRPHKS QEGTQEVSVRPSVIQKLEKGQQQPL EFVHKIGASVRKCNLVDIGQATNNR SNLVRPPVICNAPASCLPESSNDRPV TANTTSLPPAAHLDSVSKCDPNKVE KYLEQPDGASRNPVPSSHVETTSFSY QKHKESNRKSLRMNSDKLVLWKDV KSQGKTLSAGLKFHERMGTKGSLRV KSPSKLAVNPNKTDMPSNKGIFEDT IAKNHEEFFSNMDCTQEEKHLVFN KTAFWEQKCSVSSEMKFDCISLQSA SDQPQETAQDLSLWKEEQIDQEDNY ESRGSEMSFDCSSSFHSLTDQSKVS AKEVNLSKEVRTDVQYKNNKSYVSK ISSDCDDILHLVTNQSQMIVKEISLQ NARHISLVDQSYESSSSETNFDCDA SPQSTSDYPQQSVTEVNLPKEVHIGL VDKNYGSSSSEVSADSVFPLQSVVD RPPVAVTETKLRKKAHTSLVDNYGS SCSETSFDCDVSLESVVDHPQLTVK GRNLKGRQVHLKHKKRKPSSAKAH LDCDVSLGTVADESQRAVEKINLLK EKNADLMDMNCESHGPEMGFQAD AQLADQSQVAEIERQKVDVDLENKS VQSSRSSLSSDSPASLYHSAHDEPQ EALDEVNLKELNIDMEVRSYDCSSS ELTFDSDPPLLSVTEQSHLDAEGKE RHIDLEDESCESDSSEITFDSDIPLYS VIDQPEVAVYEEETVDLESKSNESCV SEITFDSDIPLHSGNDHPEVAVKEVI QKEEYIHLERKNDEPSGSEISSDSHA PLHSVTNSPEVAVKKLNPQKEEQVH LENKENEPIDSEVSLDYNIIFHSVTG RSEDPIKEISLHTKEHMYLENKSVFE TSLDSDVPLQAATHKPEVIVKETWL QREKHAEFQGRSTEFSGSKTSLDSG VPHYSVTEPQVAVNKINRKKQYVLE NKNDKCSGSEIILDSNVPPQSMTDQ PQLAFLKEKHVNLKDKNSKSGDSKI TFDSEQLQEAVKKIDQWKEEVIGLK NKINEPSTYKLIHHPDVSVQSVADQP KVAIKHVNLGNENHMYLEVKNSQY SCSEMNLDSGFLGQSIVNRPQITILE QEHIELEGKHNQCCGSEVSFDSDDP LQSVADRLRETVKEISLWKDEEVDT EDRRNEAKGFEIMYDSDVLQPVAGQ PEEVVKEVSLWKEHVDLENKIVKPT DSRINFDSHEPLQSVTNKIPGANKEI NLLREEHVCLDDKGYVPSDSEIIYVS NIPLQSVIKQPHILEEEHASLEDKSS NSYSPEESSDSNDSFQAAADELQKP VKEINLWKEDHIYLEDKSYKLGDFD VSYASHIPVQFVTDQSSVPVKEINLQ KKDHNDLENKNCEVCGSEIKCHSC VHLQSEVDQPQVSYKEADLQKEEHV VMEEKTDQPSDSEMMYDSDVPFQI VVNQFPGSVKETHLPKVVLVDLVPG DSDYEVISDDIPLQLVTDPPQLTVKDI SCINTECIDIEDKSCDFFGSEVRCNC KASTPSMTNQCKETFKIINRKKDYIIL GEPSCQSCGSEMNFNVDASDQSMT YESQGPDEKMVKYIDSEDKSCGYNG SKGKFNLEDTSHRTTHRLQKAHKEA SLRKDPRNAGLKGKSCQSSASAVDF GASSKSALHRRADKKKRSKLKHRDL EVSCEPDGFEMNFQCAPPLPSDTDQ PQETVKKRHPCKKVSSDLKEKNHDS QSSSVLKVDSVRNLKKAKDVIEDNP DEPVLEALPHVPPSFVGKTWSQIMR EDDIKINALVKEFREGRFHCYFDDD CETKKVSSKGKKKVTWADLQGKEDT APTQAVSESDDIVCGISDIDDLSVAL DKPCHRHPPAERPPKQKGRVASQC QTAKISHSTQTSCKNYPVMKRKIIRQ EEDPPKSKCSRLQDDRKTKKKVKIG TVEFPASCTKVLKPMQPKALVCVLS	None	None	None	None	None	None	None	None

									SLNIKLKEGEGLPFPKMRHHSWDN DIRFICKYKRNIFDYYEPLIKQIVISPP LSVIVPEFERRNWVKIHFNRSNQNS SAGDNDADGQGSASAPLMAVPARY GFNSHQGTSDSSLFLEESKVLHARE LPKKRNFQLTFLNHDVVKISPKSVR NKLLESQSKKKIHGKRVTTSSNKLG FPKKVYKPIILQQKPRKASEKQSIWIR TKPSDIIRKYISKYSVFLRHRYQSRSA FLGRYLKKKKSVVSRLKKAKRTAKVL LNSSVPPAGAEELSSAMANPPPKRP VRASCRVARRRKKTDESYHGRQKGP STPVRAYDLRSSSCLQQRERMMTRL ANKLRGNEVK								