

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus
O88532	ZFR_MOUSE	Mus musculus	Zinc finger RNA-binding protein	57.348653	T102;T114;T135;T136;S148;T167;S180;S195;T202;T300;S422;T423;S429;S470	S1054	22517741;22645316;21606357;40885482;33300544;34887587;36852467;35822049;39927985;40021952;28528544;34678516;39627609;40997131;36757814	MIPICPVVSFTYVPSRLGEDAKMATG NYFGFTHSGAAAAAAQYSQQPAS GVAYSHPTTVASYTVHQAPVAAHTV TAAYAPAAATVAVARPAPVAVAAAAAT AAAYGGYPTAHTATDYGYTRQOQEA PPPPPPATTQNYQDSYSYVRSTAPAV AYDSKQYYQPTATAAAVAAAQPO PSVAETYYQTAPKAGYSOGATQYTOA QQARQVTAIKPATPSPATTTFSIYPVS STVQPVAATAAVVPSYTSQATYSTTA VTYSGTSYSGYEAAYVSAASSYYQQQ QQQQKQAAAAAAATAAWTGTTF TKKTPFQNKQLKPKQPPKPPQIHYC DVCKISCAGPQTYKEHLEGQKHKKK EAALKASQNTSSSNNSTRGTQNLRL CELCDVSCGTADAYAAHIRGAKHQK VVKLHHTKLGKPIPSTEPNVVQATSS TAASASKPTASPSIGASNCTLNTSSI ATSSVKGLSTTGNSSLNSTNTKVS IPTNMAAKTSTPKINFVGGNKLOST GNKTEDLKGIDCVKNTPAASAVQIPE VKODAGSEPVTPASLAALQSDVQPV GHDYVEEVRNDEGKIVRFHCKLCEC SFNDPNAKEMHLKGRRLQYKCK VNPDLQVEVKPSIRARKIQEEMRK QMQKEEYWRREERERWRMEIRRY EEDMYWRRMEEQHWWDRRRM PDGGYPHPPGLLGLVVRPMPPO PQGPAPLRRPDSDDRYVMTKHATI YPTTEELQAVQKIVSITERALKLVSDS LSEHEKSNKEGDDKKEGGKDRAL KGVLRVGVLAAGLLRGDRNVNLVL LCSEKPSKSLLSRIAENLPKQLAVISP EKYDIKCAVSEAAIILNSCPEPKMQV TITLTSPIIREENMREGDVTSGMVKD PPDVLDRQKCLDALALRHAKWFOA RANGLQSCVHILRLDLCQRVPTWS DFPSWAMELLVEKAISSASSPQSPG DALRRVFEICISSGILKGSPLLDPC KDPFDTLATMTDQOREDTSSAQFA LRLAFRQIHKVLMPLPQMNQRF NIHNNRKRDRSDGVDGFEAEGKK DKKDYDNF	True	False	2.527	4.471	1.138	0.715	0.814