	UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochndrion	endoplasmic reticulum	golgi apparatus	pla mem
0	Q86VM9	ZCH18_HUMAN	Homo sapiens	Zinc finger CCCH domain-containing protein 18	16.722134	NaN	S6;S34;S46;S53;S59;S67;S74;S78;S83;S95;T109;S110;S118;T162;S173;S179;S487;S532;S534;S536;S842;S852;S868;S893;S896	30379171;23301498;37217939	MDVAESPERDPHSPEDEEQPQGLS DDDILRDSGSDQDLDGAGVRASDLE DEESAARGPSQEEEDNHSDEEDRA SEPKSQDQDSEVNELSRGPTSSPCE EEGDEGEEDRTSDLRDEASSVTREL DEHELDYDEEVPEEPAPAVQEDEAE KAGAEDDEEKGEGTPREEGKAGVQS VGEKESLEAAKEKKKEDDDGEIDDG EIDDDDLEEGEVKDPSDRKVRPRPT CRFFMKGNCTWGMNCRFIHPGVN DKGNYSLITKADPFPPNGAPPLGPHP LMPANPWGGPVVDEILPPPPPEPPT ESAWERGLRHAKEVLKKATIRKEQE PDFEEKRFTVTIGEDEREFDKENEV FRDWNSRIPRDVRDTVLEPYADPYY DYEIERFWRGGQYENFRVQYTETEP YHNYRERERERERENRQRERERER ERDRERERRQRERERERERERDKE RQRRKEEWERERAKRDEKDRQHRD RDREKEREKEKGKPKPRSPQPPSRQ AEPPKKEAATTGPQVKRADEWKDP WRRSKSPKKKLGVSVSPSRARRRRK TSASSASASNSSRSSSRSSSYSGSGS SRSRSRSSSYSSYSSRSSRHSSFSGS RSRSRSFSSSPSPSPTPSPHRPSIRTK GEPAPPPGKAGEKSVKKPAPPPAPPQ ATKTTAPVPEPTKPGDPREARRKERP ARTPPRRRTLSGSGSGSGSSYSGSSS RSRSLSVSSVSSVSSATSSSSSAHSV DSEDMYADLASPVSSASSRSPAPAQ TRKEKGKSKKEDGVKEEKRKRDSST QPPKSAKPPAGGKSSQQPSTPQQAP PGQPQQGTFVAHKEIKLTLLNKAAD KGSRKRYEPSDKDRQSPPPAKRPNT SPDRGSRDRKSGGRLGSPKPERQRG QNSKAPAAPADRKRQLSPQSKSSSK VTSVPGKASDPGAASTKSGKASTLSR REELLKQLKAVEDAIARKRAKIPGKA	None	None	None	None	None	None	None	Nor