

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	em ret
Q99L15	ZN281_MOUSE	Mus musculus	Zinc finger protein 281	47.724991	T60;S106;T682;S691;S693;T886;T888;S889	S392;S481;S648;S782;S805;T886	21606357;40885482;36852467;40128797;28528544;34678516;40997131	MKIGSGFLSGGGPSSSGSGSGGS SGSASGGSGGRRRAEMPTFPQSM VMFNHRLPPVTSFTRPAGTAAPPPQ CVLSSSTSAAPAAEPPPPAPDMTFK KEPAASAAAFPSQRTSWGFLQSLVSI KQEKPADPEEQPSHHHHHHHHYH GLFAGAEERSPLGGGGGGSHGVIQ DLSLLHQHAQQQPAQHHRDVLSS CSRTDEHGNQEPKQDANVKKAKRP KPESQGIKAKRKPSSASKPLVGELEG AVLSPSQPHICDHCSAAFSSYHL RRHVLHTGERPFQCSQCSMGFIQK YLLQRHEKHSREKPFQCDQCSMKF IQKYHMERHKRTHSGEKPKYKCDTC QQYFSRTDRLLKRRRTCGEAIKGA ASAEPGSSNHNSMGNLAVLSQGNT SSSRKSKSKSIAIENKEHKTGKTNE SQMSNNINMQSYVEMPTVSTSGSI IGTGIDELQKRVKLFKGSRKAD KSYLNFVSPDPVVGQKSLSGKPGG SLGIVSNNSVETISLLQSTSGKQGPIS SNYDDAMQFSKRRRYLPTASSNSAF SINVGHMVSQQSVIQSAGVSVLDNE APLSLIDSSALNAEIKSCHDKSGIPD EVLQSIDQYSGKSETQKEDPFNLTE PRVDLHTSGEHSSELVQEENLSPGTQ TPSNDKTSMLQEYSKYLQAFQEKST NAGFTLGHGFQFVLSLPLHNHTLF PEKQIYTTSPLECGFQSVTVLPS LPKPPFGMLFGSQPGLYLSALDATH QQLTPSQELDDLIDSQKNLETSSAF QSSSQKLTQKEQKNLESSTSFQIP SQELASQIDPKDIEPRTTYQIENFA QAFGSQFKSGSRVPMFITNSNGEV DHRVRTSVSDFSGYTNMMSDVSEP CSTRVKTPTSQSYR	True	False	0.921	4.762	1.182	0.5