

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
Q8BZ47	ZN609_MOUSE	Mus musculus	Zinc finger protein 609	10.914569	T332	S358;S361;S379;T381;S413;S433;S446;S452;S467;S470;S533;S575;S577;S742;T745;S757;S803;T822;S841;S845;S848;S1057	34418053;34678516	MSLSSGACGGKGV DANPVETYDSG DEWDIGVGNLIIDL DADLEKDQQKL EMSGSKEVGIPAPNAVATLPDNIKFV TPVPGPQKGEGKSKRSKSGKDS KPTPGTSLFSPSEGAASKKEVQGRA GDGASAGGLVA AVAPKGSEKAAKAS RSVAGSKKEKENS SSKGKKERSEGV GTCSEKDPGVLQPVPLGGRGSQYDG SAGMDTGTVEPLGSIAIEPGAALNPL GTKPEPEEGENE CRPLKVKSEKME SPVSTPAVLPLHLLVPVNNDISSPC EQIMVRTRSVGVNTCDVALATEPEC LGPCEPGTSVNLEGIVWQETEDGML VVNVTWRNKTYVGTLLDCTRHDWA PPRFCDSPTSDLEMRNGRGRGKRM RPSNTPVSEAAA ASDSKGTSSSSKT RAGANSKGRRGSONSSEHRPPASST SEVVKASPSSANKRKSPLSDMELN SSSEDSKGSKRVRTNSMGSATGPLP GTKVEPTLVDRNCPSVLIDCPHPN CNKKYKHINGLKYHQAHHTDDDS KPEADGDSEYGEEPALHADLSCNGA PVPQKGSLSPARSATPKVRLVEPHSP SPSSKFSTKGLCKKLSGEGDTDPG ALSNDGSDDGPSVMDETSNDAFDS LERKCMEMEKECKPSSLKSEKIPSKS LKSARPIAIPPPQIYTFQTATFTAA SPGSSSGLTTT VVQAMPNSPQLKPIQ PKPTVMGEPFTVNPALTPAKDKKKK DKKKKDSSELESPLTPGKVCRAEE GKSPFRDAAGDG I KVESLLNGSSES HQSRLASIKAEADKIYSFTDNAPSPSI GGSSRLDSTTPTQPLTPLHVVTQNG AEASSVKTNSPAYSDISDAGEDGEG KVDSAKSKDPEQLVKEGAKKTLFPP QPQSKDSPYYQGFESYYSPGYAQSSP GTLTSSSQAGMEGQPLKTKKDEEPE SVEGKVKNDVCEEKKPELSNSSQQP SVIQQRPNMYMQSLYNYAYVPPY GYS DQSYHSHLLSTNTAYRQYEEQ QKRQSLEQQQQQQRGLDKKTEMGL KEREASLKEEWKQKPSIPPTLTKAPS LTDLVKSGPGKAKEPGTDPKSVIIP KLDDSSKLPSQPPEGLKGLGEASH LGKEASEAKTGTECGRQAEVDPILW YRQETESRMWTVVYPAKYSDIKSED DRWKEERDRKLEDRSRKDSVPK EDGKESTSSDCKLPSEESRLGSKEP RPSVHVVPSSPLTQHQSYPYMHGY

