	UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochndrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extra regio
0	Q9ULD9	ZN608_HUMAN	Homo sapiens	Zinc finger protein 608	16.682301	NaN	S421;S424;T481;S493;S627;S782;S895;S964;S1098	30379171;37217939	MSVNISTAGKGVDPNTVDTYDSGDD WEIGVGNLIIDLDADLEKDRQKFEM NNSTTTTSSSNSKDCGGPASSGAGA TAALADGLKFASVQASAPQGNSHKE TSKSKVKRSKTSKDANKSLPSAALYG IPEISSTGKRQEVQGRPGEATGMNS ALGQSVSSGGSGNPNSNSTSTSTSA ATAGAGSCGKSKEEKPGKSQSSRGA KRDKDAGKSRKDKHDLLQGHQNGS GSQAPSGGHLYGFGAKSNGGGASPF HCGGTGSGSVAAAGEVSKSAPDSGL MGNSMLVKKEEEEEESHRRIKKLKT EKVDPLFTVPAPPPPISSSLTPQILPS YFSPSSSNIAAPVEQLLVRTRSVGVN TCEVGVVTEPECLGPCEPGTSVNLE GIVWHETEEGVLVVNVTWRNKTYV GTLLDCTKHDWAPPRFCESPTSDLE MRGGRGRGKRARSAAAAPGSEASFT ESRGLQNKNRGGANGKGRRGSLNA SGRRTPPNCAAEDIKASPSSTNKRK NKPPMELDLNSSSEDNKPGKRVRT NSRSTPTTPQGKPETTFLDQGCSSP VLIDCPHPNCNKKYKHINGLRYHQA HAHLDPENKLEFEPDSEDKISDCEE GLSNVALECSEPSTSVSAYDQLKAPA SPGAGNPPGTPKGKRELMSNGPGSI IGAKAGKNSGKKKGLNNELNNLPVI SNMTAALDSCSAADGSLAAEMPKLE AEGLIDKKNLGDKEKGKKATNCKTD KNLSKLKSARPIAPAPAPTPPQLIAIP TATFTTTTTGTIPGLPSLTTTVVQATP KSPPLKPIQPKPTIMGEPITVNPALVS LKDKKKKEKRKLKDKEGKETGSPKM DAKLGKLEDSKGASKDLPGHFLKDH LNKNEGLANGLSESQESRMASIKAE ADKVYTFTDNAPSPSIGSASRLECST LVNGQAPMAPLHVLTQNGAESSAA KTSSPAYSDISDAADDGGSDSRSEG MRSKASSPSDIISSKDSVVKGHSSTT AQSSQLKESHSPYYHSYDPYYSPSY MHPGQVGAPAAGNSGSTQGMKIKK ESEEDAEKKDKAEQLDSKKVDHNS ASLQPQHQSVITQRHPALAQSLYYG QYAYGLYMDQKSLMATSPAYRQQYE KYYEDQRLAEQKMAQTGRGDCERK SELPLKELGKEETKQKNMPSATISKA PSTPEPNKNHSKLGPSVPNKTEETG KSQLLSNHQQQLQADSFKAKQMEN HQLIKEAVEMKSVMDSMKQTGVDP TSRFKQDPDSRTWHHYVYQPKYLDQ QKSEELDREKKLKEDSPRKTPNKES GVPSLPVSLTSIKEEPKEAKHPDSQS MEESKLKNDDRKTPVNWKDSRGTR VAVSSPMSQHQSYIQYLHAYPYPQM YDPSHPAYRAVSPVLMHSYPGAYLS PGFHYPVYGKMSGREETEKVNTSPS VNTKTTTESKALDLLQQHANQYRSK SPAPVEKATAEREREAERERDRHSP FGQRHLHTHHHTHVGMGYPLIPGQ YDPFQGLTSAALVASQQVAAQASAS GMFPGQRRE	None	None	None	None	None	None	None	None	None