

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
Q61602	GLI3_MOUSE	Mus musculus	Transcriptional activator GLI3	15.818779	S78	S664;S849;S865;S877;S907;S980;S1006	36852467	MEAQAHSSSTATERRKKAENSIGKCPT RTDVSEKAVASSTTSNEDESPGQIYH RERRNAITMQPQSVQGLNKISEEPS TSSDERASLIKKEIHGSLPHLAEPSL PYRGTVFAMDPRNGYMEPHYHPPH LFPAPHPPVPIDARHHEGRYHYDPS PIPPLHVPSALSSSPTYDLPFIRISPH RNPTAASESPFSPHPYINPYMDYIR SLHSSPSLSMISAARGLSPTDAPHAG VSPA EYHQMALLTGQRSPYADILPS AATAGAGAIHMEYLHAMDSTRFPSP RLSARPSRKRTLSISPLSDHSFDLQT MIRTSPNSLVTILNNSRSSSSASGSY GHLSASAIPALSFTYPSAPVSLMHM QQILSRQQSLGSAFGHSPPLIHPAPT FPTQRPIPGIPTVLNPNVQVSSGPESS QSKPTSESAVSTGDPMHNKRSKIK PDEDLPSPGSRGQQEQPEGTTLVKE EADKDESKQEPEVIYETNCHWEGCT REFDTQDQLVHHINNDHIHGEKKEF VCRWLDCSREQKPFKAQYMLVVMH RRHTGEKPHKCTFEGCTKAYSRLN LKTHLRSHTGEKPYVCEHEGCNKAF SNASDRAKHQNRTHSNEKPYVCKIP GCTKRYTDPSSLRKHVKTVHGPEAH VTKKQRGDMHPRPPPPRDSGSHSQ SRSPGRPTQGAFGGEQKELSNNTTSKR EECLQVKTVKAEKPMTSQSPGGQS SCSSQQSPISNYSNSGLELPLTDGGS VADLSAIDETPIMDSTISTATTALALQ ARRNPA GTKWMEHIKLERLKQVNG MFPRLNPILPSKAPAVSPLIGNGTQS NNNYSSGGPGTLLPSRSDLGVDFT VLNTLNRRDSNTSTISSAYLSSRRSS GISPCFSSRRSSEASQAEGRPQNVSV ADSYDPISTASRRSSEASQGDGLPS LLSLTPVQYRLKAKYAAATGGPPPT PLPHMERLSLTKMALLGEGRDSGV TLPPVHPPRRCSDGGGHTYRGRHL MPHDALANSVRRASDPVRTVSENM SLARVQRFSSLNSFNPPNLPSSVEK RSLVLQNYTRQESSQPRYFQASPCP PSITENVALEALTMDADANLNDEDL LPDDVVQYLNSQNQTGYGQQLQSGI SEDSKVAHEPEDLDLAGLPDSHVQ EYPALEQPCSEGSKTDLPQWNEVS SGTSDLSSSKLKCGQQRPSAQPRG FGLYNNMVVHPHNLWKVGTGPAGG YQTLGENSSTYNGPEHFAIHSGDGL

