

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracell region
Q61602	GLI3_MOUSE	Mus musculus	Transcriptional activator GLI3	24.992371	S78	S664;S849;S865;S877;S907;S980;S1006	36852467	MEAQAHSSSTATERKKAENSIGKCP RTDVSEKAVASSTTSNEDESPGQYH RERRNAITMQPQSVQGLNKISEEPS TSSDERASLIKKEIHGSLPHLAEP PYRGTVFAMDPRNGYMEPHYHPPH LFPAPFHPVPIDARHHEGRYHYDPS PIPPLHVPSALSSSPTYDLPFIRISPH RNPTAASESPFSPHPYINPYMDYIR SLHSSPSLSMISAARGLSPTDAPHAG VSPAEEYHQMALLTGQRSPYADILPS AATAGAGAIHMEYLHAMDSTRFPSP RLSARPSRKRTLISPLSDHSFDLQT MIRTS PNSLVTLNNSRSSSASGSY GHLASASIPALSFTYPSAPVSLMHM QQILSRQQLGSAFGHSPPLIHPAPT FPTQRPIPIPTVLPVQVSSGPESS QSKPTSEAVSSTGDPMHNKRKIK PDEDLPSPGSRGQEQPEGTTLVKE EADKDESKQEPVIYETNCHWEGCT REFDTQDQLVHHINNDHIHGEKKEF VCRWLDCSREKPFKAQYMLVVHM RRHTGEKPHKCTFEGCTKAYSRL LKTHLRSHTGEKPYVCEHEGCNKAF SNASDRAKHQNRTHSNEKPYVCKIP GCTKRYTDPSSLRKHVKTVHGPEAH VTKKQRGDMHPRPPPPRDSGSHSQ SRSPGRPTQGAFGGEKELSNITTSKR EECLQVKTVAEKPMTSQPSGGQS SCSSQSPISNYSNGLELPLTDGGS VADLSAIDETPIMDSTISTATTALALQ ARRNPAGTKWMEHIKLERLQVNG MFPRLNPLPSKAPAVSPLIGNGTQS NNNYSSGGPTLLPSRDLGVDFT VLNLTNRRDSNTSTISSAYLSSRRSS GISPCFSSRRSSEASQAEGRPQNVSV ADSYDPISTASRRSSEASQGDGLPS LLSLTPVQYRLKAKYAAATGGPPPT PLPHMERLSLKTMMALLGEGRDSGV TLPPVHPPRRCSGGGHYRGRHL MPHDALANSVRRASDPVRTVSEN SLARVQRFSSLNSFNPNLPPSVEK RSLVLQNYTROESSQPRYFOASPCP PSITENVALEALTMADANLNDEDL LPDDVVQYLN SQNTGYGQQLQSGI SEDSKVAHEPEDLDLAGLPDSHVQ EYPALEQPCSEGSKTDLPQWNEVS SGTDLSSSKLKCQQQRPSAQQPRG FGLYNNMVVHPHNLWKVGTGPAGG YQTLGENSSTYNGPEHFAHSGDGL GTNGNTFHEQPFTQYQYQSLNRQ PLTSSALDHACGTGIQGSKLKGNL QENGGLLDFSLVAPNELAGNTVNG MQTQDQMGQGYIAHQLLSGSMQH QGSPSRGQVQLGQV GATSHINIQG TESCLPQTQDNSSQPSSMAAIRGYQ PCASYGNNRRQAMPNGNLTLOQGO LSDMSQSSRVNSIKMEAQGSQQL CSTVQNYSGQFYDQTMGFSQDRK AGSFLSDANCLLQNGTENSSELLS PGANQVTSTVDSFESHDLGVQIDF DAIIDDGDHTSLMSGALSPSIQNL HSSSRLTTPRASLPFSLSMGTINM AIGDMSSLLTSLAEESKFLAVMQ	True	False	3.765	5.0	1.802	1.584	1.308	2.292	2.123