

UniprotKB ID	Entry name	organism	full name	oglnacscore	oglnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracellular region
Q9NS15	LTBP3_HUMAN	Homo sapiens	Latent-transforming growth factor beta-binding protein 3	22.327667	NaN	NaN	37217939;39302247	MPGPRGAAGGLAPEMRGAGAAGLL ALLLLLLLLLLGLGGRVEGGPAGERG AGGGGALARERFKVVFAPVICRKTCL KGQCRDSCQGSNMTLIGENGHST DTLTGSGFRVVCPLPCMNGGQCSS RNQCLCPPDFTRFCQVPAGGAGG GTGGSGPGLSRTGALSTGALPPLAPE GDSVASKHAIYAVQVIADPPGPEGP PAQHAAFLVPLGPGQISAEVQAPPPV VNVRVHHPPEASVQVHRIESNAES AAPSQHLLPHPKPSHPRPPTQKPLG RCFQDTLPKQPCGSNPLPGLTKQED CCGSIGTAWGQSKCHKCQPLQYTG QKPGPVRGEVGADCPQGYKRLNSTH CQDINECAMPVGCVRHGDCLNPNPS YRCVCPGHSGLGSRTOCIADKPEEK SLCFRLVSPHQHPLTTRLRQL CCCSVGKAWGARCQRCPDGTAAAF KEICPAGKGYHILTSHTLTIQGESD FSLFLHPDGPQPLPESPSQAPPP EDTEERGVTTDSPVSEERSVQQSH PTATTTTPARPYELISRPSPPTMRWF LPDLPPSRSAVEIAPTQVTETDECRL NQNICGHGECVPGPPDYSCHCNPG YRSHPQHRYCVDVNECEAEPCGPG RGICMNTGGSYNCHCNRYRLHVG AGGRSCVDLNECAKPHLCGDGGFCI NFPGHYKCNCYPGYRLKASRPVCE DIDECRDPSSCPDGKCKENKPGSFKCI ACQPGYRSQGGGACRDVNECAEGS PCSPGWENLPGSFRCTCAQGYAPA PDGRSCLDVDECEAGDVCNNGICSN TPGSFQCQLSGYHLSRDRSHCEDI DECDPAAACIGGDCINTNGSYRCLC PQGHRLVGGRCQDIDECSDPPLC LPHGACKNLQGSYVVCDEGFTPTQ DQHGCEEVEQPHHKKCYLNFDDT VFCDSVLATNVVQEQCCSLGAGW GDHCEIYCPVYSSAEFFSLCPDGK GYTQDNNIVNYGIPAHRDIDECMLF GSEICKEGKCVNTQPGYECYCKQGF YYDGNLLECVDVDECLDESNCRNG VCENTRGGYRCACTPPAEYSPAQRQ CLSPPEMDVDECODPAACRPGRCV NLPGSYRCECRPPWVPGPSGRDCQL PESPAERAPERDVCWSQRGEDGM CAGPLAGPALTFDCCCRQGRGWG AQCRCPPRGAGSHCPTSQSESNSF WDTSPLLGKPPREDSSEEDSDEC RCVSGRCVPRPGGAVCECPGGFOLD ASRARCVDIDECRELNQRGLLCKSE RCVNTSGSFRVCCKAGFARSRPHGA CVPQRRR	False	True	1.317	2.03	1.625	1.704	1.155	3.573	4.59