

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum
Q02447	SP3_HUMAN	Homo sapiens	Transcription factor Sp3	19.197656	T118;S131;T134;S135;S136;T231;S334;T346;S375;S389;T390;T407;S408;T417;T420	S73;S563;S566;S646	38665916;34019948;33214551;35289036;39531497;26431879;35138101;30397120;29351928;40596516;35254053	MTAPEKPVKQEEAALDVIDSGGGG GGGGGHGEYLQQQQHNGAVAA AAAAQDTQPSPLALLAATCSKIGPPS PGDDEEEAAAAGAPAAAGATGDLA SAQLGGAPNRWEVLSATPTTIKDEA GNLVQIPSAATSSGQYVPLQLNLQN QQIFSVAPGSDSSNGTVSSVQYQVIP QIQSADGQVQIGFTGSSDNGGINQ ESSQIQIIPGSNQTLLASGTPSANIQN LIPQTGQVQVQVVAIGSSFPQTQV VANVPLGLPGNITFVPINSVDLDSL LSGSSQMTAGINADGHLINTGQAM DSSDNSERTGERVSPDINETNTD LFVPTSSSSQLPVTIDSTGILQNTN SLTTSSGQVHSSDLQGNVQSPVSEE TQAQNIQVSTAQPVVQHLQLQESQQ PTSQAQIVQGITPQTIHGVQASGQNI SQQALQNLQLQLNPGTFLIQAQIVT PSGQVTWQTFQVQVQNLQNLQIQ NTAAQQITLTPVQTLTLGQVAAGGAF TSTPVSLSGQLPNLQTVVNSIDSA GIQLHPGENADSPADIRIKEEPPDE EWQLSGDSTLNTNDLTHLRVQVVD EEGDQHQEGKRLRRVACTCPNCK EGGGRGTNLGKKKHICHIPGCGKV YGKTSHLRAHLRWHSGERPFCVNW MYCGKRFRSDELQRHRRHTHTGK KFVCEPKSRFMRSDHLAKHIKTHQ NKKGIHSSSTVLA SVEAARDTLITA GGTLLANIQQGSVSGIGTVNTSAT SNQDILTNTIPLQLVTVSGNETME	True	False	4.531	5.0	False	False