	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	extracellular region
0	Q62445	SP4_MOUSE	Mus musculus	Transcription factor Sp4	36.83566	S75;T92;T577;T587	S44	40885482	MSDQKKEEEEEAAAAMATEGGKTS EPENNNKKPKTSGSQDSHPSPLALL AATCSKIGTPGENQATGQQQIIIDPS QGLVQLQNQPQQLELVTTQLAGNA WQLVASTPPASKENNVSQPASSSSS SSSSNNGSSSPTKTKSGNPSTPNQF QVIQVQNPSGSVQYQVIPQLQTVEG QQIQINPTSSSSLQDLQGQIQLISAG NNQAILTAANRTASGNILAQNLANQ TVPVQIRPGVSIPLQLQTLPGTQAQV VTTLPINIGGVTLALPVINNVTAGGG TGQVGQPTTTTDSGTSNGNQLVSTP TTSTAPASTMPESPSSSTTCTTTAST TLTSSDTLVSSADTGQYASTSASSSE RTIEEPQTPAATESEAQSSSQLQSNG IQNAQDQSNSLQQVQIVGQPILQQIQ IQQPQQQIIQAIPPQSFQLQSGQTIQT IQQQPLQNVQLQAVNPTQVLIRAPTL TPSGQISWQTVQVQNIQSLSNLQVQ NAGLSQQLTITPVSSSGGTTLAQIAP VAVAGAPITLNTAQLASVPNLQTVSV ANLGAAGVQVQGVPVTITSVAGQQQ GQDGVKVQQATIAPVTVAVGGIANA TIGAVSPDQLTQVHLQQGQQTSDAE VQPGKRLRRVACSCPNCREGEGRGS SEPGKKKQHVCHIEGCGKVYGKTSH LRAHLRWHTGERPFICNWMFCGKR FTRSDELQRHRRTHTGEKRFECPEC SKRFMRSDHLSKHVKTHQNKKGGG TALAIVTSGELDSSVTEVLGSPRIVTV AAISQDSNPATPNVSTNMEEF	True	False	2.576	5.0	1.066	1.003	0.784	1.352	1.206