

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
089090	SP1_MOUSE	Mus musculus	Transcription factor Sp1	56.564599	S613;T641;S642;S699;S703	S2;S7;S61;S103;T280;T455;S613;T641;S642;T652;T669;S671;T682;S699;S703;T738	34260942;23315940;19070619;18596307;18513490;20138838;25657638;21493720;22874425;14749383;16879824;19598243;28528544;19285002;27010859;29125738	MSDQDHSMDDEVTAUVVKIEKDVGGN NGGSGNGGGAAFSQTRSSSTGSSSS SGGGGGQESQPSPLALLAATCSRIES PNENSNNNSQGSPSQSGGTGELDLTAT QLSQGANGWQIHISSSSGATPTSKEQS GNSTNGSNGSESSKNRTVSGGQYV VAATPNLQNNQVLTGLPGVMPNIQY QVIPQFQTVDDGQQLQFAATGAQVQQ DGSGQIQIIPGANQQIIPNRGSGGNII AAMPNLLQAVPLQGLANNVLSGQ TQYVTNVPVALNGNITLLPVNSVSA TLTPSSQAGTISSSGSQESSSQPVTS GTAISSASLVSSQASSSSFFTNANSY STTTTTSNMGIMNFTSSGSSGTSSQ GQTPQRVGGLQGSDSLNIQQNQTSG GSLQGSQQKEGEQSQQTQQQILIQ POLVQGGQALQALQAAPLSGOTFTT QAISQETLQNLQAVQNSGPIIIRTP TVGPNQVSWQTLQNLQVQNPQ AQITLAPMQVSLGQTSSTNTLTP IASAASIPAGTVTVNAAQLSSMPGLQ TINLSALGTSGIQVHQLPGLPLAIAN TPGDHGTQLGLHGSGGDDGIHDETA GGEGENSSDLQPQAGRRTREACTC PYCKDSEGRASGDPGKKKQHICHIQ GCGKVYGKTSHLRAHLRWHTGERP FMCNWSYCGKRFTRSDQLRHKRT HTGEKKFACPECPKRFMRSDHLSK HIKTHQNKKGPGVALSVGTLPDLS GAGSEGTATPSALITTNMVAMEAICP EGIARLANSGINVMQVTELQSNISG NGF