

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
Q86V48	LUZP1_HUMAN	Homo sapiens	Leucine zipper protein 1	8.237967	S785;T793;T797;S875	S255;S394;S425;S440;S512;S570;S574;S611;S659;T679;S690;S745;S905;S932;T958;S995;S1042	23301498;30379171;35254053;28657654;28510447;30059200;34725712;34019948	MAEFTSYKETASSRHLRFKLQSLSR RLDELEEATKNLQKADELDDLQDK VIQAEGSNSSMLAEIEVLRQRVLRIE GKDEEIKRAEDLCRLMKEKLEEEEN LTRELKSEIERLQKRMAEIEKLEEF SRSKNDCTQLCLSLNEERNLTKKIS SELEMLRVKVKESSEDRLDKTEQ SLASELEKLSLTLFSVSRKYLNEK EKENEKLIKELTQKLEQNKMMNRDY TRNASNLERNDLRIEDGISSTLPSKE SRRKGGLDYLGQVENETRNXSENE KNRNQEDNKVKDLNQEIEKLTQIK HFESLEEEELKMKSKNNDLQDNYL SEQNKNKLLASQLEEIKLQIKKQKEL ENGEVEGEDAFLSSKGRHERTKFRG HGSEASVSKHTARELSPQHKRERLR NREFALNNENYSLSNRQVSSPSFTN RRAAKASHMGVSTDSGTQETKKTED RFVPGSSQSEGKKSREQPSVLSRYPP AAQEHKAWKGTSKPGTESGLKGV EKTRTFSDTTHGVSVPDPLGRADK ASDTSSETVFGKRGHVLGNGSQVTQ AANSGCSCAIGALASSRRSSSEGLSK GKKAANGLEADNSCPNSKAPVLSKY PYSCRSQENILQGFSTSHKEGVNQP AAVVMEDSSPHEALRCRVIKSSGRE KPDSDDDLDIASLVTAKLVTITPE PEPKQPNSREKAKTRGAPRTSLFE NDKDAGMENESVKSVRASNTMEL PDTNGAGVKSQRPFSPREALRSRAII KPVIVDKDVKIMGGSGTETTLEKQ KPVSKPGPNKVTSSITIYPSDSSSPRA APGEALRERHTSTSNIQVGLAELTSV SNHVSPFELSIHKHDITLQLAEAEER MADGPKDRPETVVSRSIIKPSDP VERNSHAPPAETIRWKSHPSEVVG FSDARHVTVRNAWKSRRDLKSLEDP PTRIGKNVESTNSNAYTQRSSTDFSE LEQPRSCLEFQGTRRVGPGSSGDAPE PSSRRTQSSLTVSEVLTRRNRVGDIT TVAAWNHSASMEEEGEDCTLSVYR QLHNSLDPSELPGKQGLPESGRVRA EERLRPTRPCAEEN