

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
Q8R4U7	LUZP1_MOUSE	Mus musculus	Leucine zipper protein 1	37.344396	T794	S256;S261;S395;S513;S571;S575;S612;S660;T680;S691;S746;S906;T952;S988	22645316;36852467;34887587;28528544;26192747	MAELTNYKDAASNRHLRFKLQSLSR RLDELEEATKNLQRAEDELDDLQDK VIQAEGSDSSTLAEIEVLRQRVLKIEG KDEEIKRAEDLCHTMKEKLEEEENL TRELKSEIERLQKRMVDLEKLEEALS RSKNECSQLCLSLNEERNLTKKISS ELEMRLRVKVKELSESSEDRDKTEQS LVSELEKCLKSLTSLFVNERKYLNEKE KENEKIIKELTQKLEQNKKMNRDH MRNASTFLERNDLRIEDGISSTLSSK ESKRKGLDYLKQVENETRDKSENE KNRNQEDNKVKDLNQEIEKLTQIK HFESLEEEELKKMRAKNNDLQDNYL TELNRRNSLASQLEEIKLQVRKQKE LGNGDIEGEDAFLLRGRHERTKLK GHGSEASVSKHTSRELSPOHKRERL RNREFALSNEHYSLSSKQASSPVFT NKRAAKASNMGMGTDSGTQETKRT EDRFAPGSSHSEKRGREQPSVLSR YPPAAQEHTKVWKGAPKPGTESGLK GKVEKTRTFSDSTHVSVPNDIVGK GDKTSDLSSSEAHCGKRGQVPGHAS QGTQAVESSCSKAIGALSSSQASSE GLSKGKKTANGLAADANFNSKAPI LSKYPYSSRSQENILQGFSLPNKEGV DQPVAVMEDSSQHEALRCRVIKSS GREKPDSDDDLIESFVTAKLVNTTI TPEPEPKPQPNREKVKSRGGTRTA LFENDKNAIENDSVKPTRPSSNAIE FPDANCAGVKNQRPFSREALRSRA IIPVIIDKDVKKIMGGSGTEVVLEKQ KSTSKSVTSKVTSSITIYPSDSSGPRA VPSEAPRERHTSTSNIQVGPPELTAI SNHVSSPLELSIHKHDITLQLTEAER VGDGSPKNRAEMVVSRSILIKPSES VEKNSHVPPAETIRWKSASASSDR HITVRNAWKSQRDLKCEDPPTGIG RNMEATNAYTQRPCDFLELEQPRS QPSEQGARRVGNSGDAPELSPRRTQ SSLTASEVLTRDRMGGAITAASCN HSSMEEGEDSTFVTSRRIHNPLEH SELPGKQGLPEPEPVWTEERLHPAK PYAEED