

Product datasheet for **MR215666**

Loxl2 (NM_033325) Mouse Tagged ORF Clone

Product data:

Product Type:	Expression Plasmids
Product Name:	Loxl2 (NM_033325) Mouse Tagged ORF Clone
Tag:	Myc-DDK
Symbol:	Loxl2
Synonyms:	1110004B06Rik; 4930526G11Rik; 9430067E15Rik
Mammalian Cell Selection:	Neomycin
Vector:	pCMV6-Entry (PS100001)
E. coli Selection:	Kanamycin (25 ug/mL)



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ORF Nucleotide
Sequence:

>MR215666 representing NM_033325
 Red=Cloning site Blue=ORF Green=Tags(s)

TTTTGTAATACGACTCACTATAGGGCGGCCGGAATTCGTCGACTGGATCCGGTACCGAGGAGATCTGCC
 GCC**CGGATCGCC**

ATGGAGCTCCATTTGGCTCCTGCCTCTCCGGCTGTTTGGCTCTGCTTGTCTTGTCTCCCTGAGCC
 TAGCACAGTACGAGGGCTGGCCCTACCAGCTCCAGTACCCTGAGTACTTCCAGCAGCCCGCTCCTGAGCA
 CCATCAGCGGCAGGTGCCCTCCGATGTGGTCAAGATCCAGGTCCGCTGGCGGGCCAGAAGAGGAAGCAC
 AATGAGGGCCGCGTGGAGGTCTACTACGAAGGCCAGTGGGGCACGGTGTGCGACGATGACTTCTCGATCC
 ATGCCGCCATGTGGTCTGCCGGCAAGTGGGCTATGTAGAGGCCAAGTCTGGGTGCCAGCTCCTCCTA
 CGGTCCAGGGCAAGGCCCATCTGGTTGGACAATATCTACTGTACTGGCAAAGAGTCGACCTGGCATCT
 TGCTCCTCCAATGGCTGGGTGTACTGACTGCAAGCACACTGAAGACGTTGGAGTGGTGTGTAGTGA
 AAAGAATTCCTGGCTTCAAATTTGACAATTCGTTGATCAACAAATAGAGAGCCTAAATATACAGGTGGA
 AGACATCCGGATTCGGCCATCCTTTTCGCTTTTCGCCATCGCAAGCCTGTGACAGAGGGCTACGTGGAG
 GTGAAGGAGGGCAAGGCTTGAAGCAGATCTGCAACAAACTGGACAGCCAAGAATTCACAGTGGTCT
 GTGGCATGTTTCGGCTTCCCTGCAGAGAAGACTACAACCCCAAAGCCTATAAAACCTTTGCCTCGGGAG
 GAAGCTGCGTTACTGGAAGTTTTCTATGAACTGCACGGGCACTGAAGCGCATATCTCCAGCTGCAAGCTG
 GGCCCTTCCGTGACCCGGGACCTGTGAAGAAGCCACCTGTGAGAACGGGCAGCCAGCTGTGGTCAAGT
 GTGTGCCTAGCCAGATCTCAGCCCCGATGGACCTCAAGGTTCCGAAAGCCTACAAGCCAGAGCAACC
 CTTGGTGCCTGAGAGGTGGAGCCAGGTCCGGGAGGGCCGAGTGGAGGTGTGAAGAATGGAGAATGG
 GAAACCTCTGCGATGACAAGTGGACCTGGTATCTGCCAGTGTGGTCTGCCGAGAGCTGGGCTTTGGGA
 CGCTAAAGAGGCCATCACAGGCTCCAGACTAGGGCAAGGATTGGGCCATCCTCAATGAAGTCCA
 GTGCACAGGGACTGAGAAGTCCATCATAGACTGCAAAATCAACACAGAGTCTCAAGGCTGCAACCATGAA
 GAAGATGCCGGGTGCGATGCAACATCCCCATCATGGTTTTCCAGAAAAAGGTGCGCCTGAATGGAGGCC
 GCAATCCTTATGAGGGCCGAGTGGAGGTGCTAACAGAGAGAAATGGGTCCCTTGTGGGGGACTGTATG
 CGGCCAGAACTGGGGCATTGTGGAAGCCATGGTGGTCTGCCGGCAGCTAGGCCTGGGCTTTGCCAGCAAT
 GCCTTTCAGGAGACCTGGTACTGGCATGGAAATATCTTCGCCAACACGTGGTCAAGTGGAGTGAAGT
 GCTCAGGAACGGAGCTGTCCCTAGCACACTGCCCCATGACGAGGAGGTGGCCTGCCCGAGGGCGGGGT
 GCGGTTTTGGTGTGGAGTCGCTCGGAAACTGCACCTGACCTGGTGTAAATGCTGAGATTGCCAG
 CAGACTGCCTACCTGGAGACAGGCCATGTCTTGTGTCAGTGTCCATGGAGGAGAACTGCCTCTCCG
 CCTCCGCTGTGCACCCGACCCACCAGAGGCCACCGGCCCTTTTACGCTTCTCCTCCCAGATCCACAA
 CAATGGCCAGTCTGACTTCCGCCCAAGAATGGCCGCATGCGTGGATTTGGCACGACTGCCACAGGCAC
 TACCACAGCATGGAAGTCTTCACTTACTATGACCTGCTGAGCCTCAACGGCACCAAGGTGGCTGAGGGCC
 ACAAGGCCAGTCTGCCTGGAGGACACTGAGTGTGAGGGAGACATTGAGAAGAGTTACGAGTGTGCCAA
 CTTTGGAGAACAAGGCATCACCATGGGCTGTGGGACATGTACCGTCAAGATTGACTGCCAGTGGATA
 GACATACCGATGTGCCCCCTGGAGACTACCTGTTCCAGGTTGTCATTAACCCCAACTATGAAGTGCCAG
 AATCAGATTTCTTAACAACATCATGAAGTGCAGGAGCCGCTATGATGGCTACCGCATCTGGATGTACAA
 CTGTACAGTAGGTGGAGCCTTCAGTGGAGAGACAGAACAAGAAGTTCGAACACTTCAGTGGACTTCTAAAT
 AACCAGCTCTCTGTACAG

ACGCGTACGCGGCCGCTCGAGCAGAAACTCATCTCAGAAGAGGATCTGGCAGCAAATGATATCCTGGATT
 ACAAGGATGACGACGATAAGGTTTAA

Protein Sequence: >MR215666 representing NM_033325
Red=Cloning site Green=Tags(s)

MELHFGSCLSGCLALLVLLPSLSLAQYEGWPYQLQYPEYFQQPAPEHHQRQVPSDVVVKIQVRLAGQKRKH
NEGRVEVYYEGQWGTVCDDDFSIHAAHVCRQVGYVEAKSWAASSYGPGEPIWLDNIYCTGKESTLAS
CSSNGWGVTDCKHTEDVGVVCEKRIPGFKFDNSLINQIESLNIQVEDIRIRPILSAFRHRKPVTEGYVE
VKEGKAWKQICNKHWTAKNSHVVCGMFGFPAEKTYNPKAYKTFASRRKLRWKFMSNCTGTEAHISSCKL
GPSVTRDPVKNATCENGQPAVVSCVPSQIFSPDGPSRFRKAYKPEQPLVRLRGGAVGEGRVEVLKNGEW
GTICDDKWDLVASVVCRELGFGTAKEAITGSRLGQIGPIHLNEVQCTGTEKSIIDCKFNTESSQGCNHE
EDAGVRCNIPIMGFQKKVRLNGGRNPYEGRVEVLTERNGLVWGTVCGQNWGIVEAMVVCRLGLGFASN
AFQETWYWHGNIFANNVMSGVKCSGTELSLAHCRHDEEVACPEGGVRFGAGVACSETAPDLVLNAEIVQ
QTAYLEDRPMSLLQCAMEENCLSAVHTDPTRGHRLLRFSSQIHNNQSDFRPKNGRHAWIWHDCRRH
YHSMEVFYYDLLSLNGTKVAEGHKASFLEDTECEGDIQKSYECANFGEQGITMGCWDMYRHDIDCQWI
DITDVPPGDYLFQVVINPNYEVPESDFSNNIMKCRSRYDGYRIWYMNCHVGGAFSEETEQKFEHFSGLLN
NQLSVQ

TRTRPLEQKLISEEDLAANDILDYKDDDDKV

Restriction Sites: Sgfl-MluI

Cloning Scheme:



ACCN: NM_033325

ORF Size: 2328 bp

OTI Disclaimer: The molecular sequence of this clone aligns with the gene accession number as a point of reference only. However, individual transcript sequences of the same gene can differ through naturally occurring variations (e.g. polymorphisms), each with its own valid existence. This clone is substantially in agreement with the reference, but a complete review of all prevailing variants is recommended prior to use. [More info](#)

OTI Annotation: This clone was engineered to express the complete ORF with an expression tag. Expression varies depending on the nature of the gene.

Components: The ORF clone is ion-exchange column purified and shipped in a 2D barcoded Matrix tube containing 10ug of transfection-ready, dried plasmid DNA (reconstitute with 100 ul of water).

Reconstitution Method:

1. Centrifuge at 5,000xg for 5min.
2. Carefully open the tube and add 100ul of sterile water to dissolve the DNA.
3. Close the tube and incubate for 10 minutes at room temperature.
4. Briefly vortex the tube and then do a quick spin (less than 5000xg) to concentrate the liquid at the bottom.
5. Store the suspended plasmid at -20°C. The DNA is stable for at least one year from date of shipping when stored at -20°C.

RefSeq: [NM_033325.2](#), [NP_201582.2](#)

RefSeq Size: 5186 bp

RefSeq ORF: 2331 bp

Locus ID: 94352

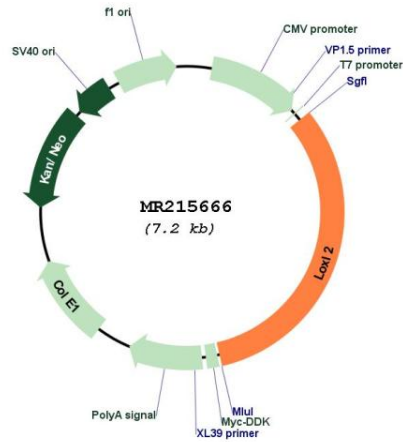
UniProt ID: [P58022](#)

Cytogenetics: 14 D2

MW: 87.5 kDa

Gene Summary: Mediates the post-translational oxidative deamination of lysine residues on target proteins leading to the formation of deaminated lysine (allysine) (By similarity). Acts as a transcription corepressor and specifically mediates deamination of trimethylated 'Lys-4' of histone H3 (H3K4me3), a specific tag for epigenetic transcriptional activation (By similarity). Shows no activity against histone H3 when it is trimethylated on 'Lys-9' (H3K9me3) or 'Lys-27' (H3K27me3) or when 'Lys-4' is monomethylated (H3K4me1) or dimethylated (H3K4me2) (By similarity). Also mediates deamination of methylated TAF10, a member of the transcription factor IID (TFIID) complex, which induces release of TAF10 from promoters, leading to inhibition of TFIID-dependent transcription (By similarity). LOXL2-mediated deamination of TAF10 results in transcriptional repression of genes required for embryonic stem cell pluripotency including POU5F1/OCT4, NANOG, KLF4 and SOX2 (PubMed:25959397). Involved in epithelial to mesenchymal transition (EMT) via interaction with SNAI1 and participates in repression of E-cadherin, probably by mediating deamination of histone H3 (By similarity). During EMT, involved with SNAI1 in negatively regulating pericentromeric heterochromatin transcription (By similarity). SNAI1 recruits LOXL2 to pericentromeric regions to oxidize histone H3 and repress transcription which leads to release of heterochromatin component CBX5/HP1A, enabling chromatin reorganization and acquisition of mesenchymal traits (By similarity). Interacts with the endoplasmic reticulum protein HSPA5 which activates the IRE1-XBP1 pathway of the unfolded protein response, leading to expression of several transcription factors involved in EMT and subsequent EMT induction (By similarity). When secreted into the extracellular matrix, promotes cross-linking of extracellular matrix proteins by mediating oxidative deamination of peptidyl lysine residues in precursors to fibrous collagen and elastin (By similarity). Acts as a regulator of sprouting angiogenesis, probably via collagen IV scaffolding (By similarity). Acts as a regulator of chondrocyte differentiation, probably by regulating expression of factors that control chondrocyte differentiation (PubMed:21071451).[UniProtKB/Swiss-Prot Function]

Product images:



Circular map for MR215666