

Product datasheet for **MC228201**

Eya2 (NM_001271963) Mouse Untagged Clone

Product data:

Product Type:	Expression Plasmids
Product Name:	Eya2 (NM_001271963) Mouse Untagged Clone
Tag:	Tag Free
Symbol:	Eya2
Synonyms:	Eab1
Vector:	pCMV6-Entry (PS100001)
E. coli Selection:	Kanamycin (25 ug/mL)
Cell Selection:	Neomycin



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Fully Sequenced ORF: >MC228201 representing NM_001271963
 Red=Cloning site Blue=ORF Orange=Stop codon

TTTTGTAATACGACTCACTATAGGGCGGCCGGGAATTCGTCGACTGGATCCGGTACCGAGGAGATCTGCC
 GCC**CGATCGCC**

ATGTTAGAAGTGGTGACCTCACCCAGCCTCGCAACAAGCAGTGACTGGAGCGAGCACGGTGTGCCGTGG
 GGACGCTGAGTGACAGGGAAGGCATCGCCAAATCAGCGGCTCTGAGTGTGCCTCAGCTCTTTGTGAAGTC
 TCATCCACGTGTCCCTCCTGGTCACTCCTCCACAGCCATGGCGGCCTATGGCCAGACACAGTACAGCACA
 GGCATTACAGCAGGCACCACCTATACAGCGTACCCAATCCGGCGCAAGCCTATGGAATCCCCCTTACA
 GCATCAAGACAGAAGACAGTTTGAATCACTCCCCAGCCAGAGCGGGTCTGAGCTATGGACCGAGCTT
 CAGCACCGCGCTGTGGACAGAGCCCCACACCTACCCCGTGCACAGCACCGCTGGGCTCTATCAAGGC
 GCCAACGGACTGACCAACACCGCTGGATTTGGGAGCGTGCACCAGGATTATCCGCTACCCAGCTTTT
 CACAGAACCAGTACCCCACTATTTAGCCCATCATAACAACCCGCTACGTCCTGCCAGCAGCCTCTG
 CTCTCGCCCTCTCCAGTCCACCTACGTCCTCCAGGAGGCTCCTCACAATGTCCCAGCCAGAGTTCT
 GAGTCCCTGGCCGGAGACTACAACACACAACCGACCCCTCCACACCAGCAAAGGAGGGTGACACAGAGA
 GGCCACATCGAGCCTCGGATGGGAAGCTACGGGGCCGGTCAAAGAGAAATAGTGACCCCTCCCCAGCAGG
 AGACAATGAAATCGAGCGGTGTTCTGCTGGGACCTGGACGAGACAATCATTATCTTCCACTCCCTGCTC
 ACAGGGACGTTTGCATCCAGATACGGGAAGGACACCACGACGCTGTGCGCATTGGCCTGATGATGGAGG
 AGATGATCTTCAACCTTGCTGACACACCTGTTCTTCAATGACCTGGAGGACTGTGACCAATCCACGT
 GGATGATGTCTCATCCGATGACAATGGTCAGGATTTAAGCACATACAACTTCTCCACTGATGGCTTCCAC
 AGCACGGCGCCAGGAGCCAGCTTGTGCCTGGGTACAGGTGTTTCATGGCGGTGGACTGGATGAGGAAAC
 TGGCCTCCGCTACCGTGTGTAAGGAGATGTACAACACCTACCGCAACAACGTGGGTGGCTTGTATAGG
 TGCTCCCAAAAGAGAGACCTGGCTGCAGCTGCGCGCCGAGCTGGAGGCCCTGACTGACCTCTGGCTCAC
 CACTCCCTGAAAGCCCTCAATCTCATCAACTCTGACCCAACTGTGTCAATGTGTTGGTACCACCACGC
 AACTGATCCCTGCATTGGCCAAGGTCTGCTGTACGGTCTGGGCTCCGTGTTCCCATCGAGAACATCTA
 CAGTGGCACAAGACAGGCAAGGAGAGCTGCTTCAAAGAATCATGCAGAGGTTTGGCCGCAAAGCTGTC
 TACATTGTGATAGGCGACGGGTAGAGGAAGAGCAAGGAGCCAAAAGCACAACATGCCTTTCTGGAGGA
 TATCTGTGATGCTGACCTGGAGGCTCTAAGGCATGCCCTGGAAGTGGAGTATCTA**TAG**

ACGGTACGGCGCCGCTCGAGCAGAACTCATCTCAGAAGAGGATCTGGCAGCAAATGATATCTGGATT
 ACAAGGATGACGACGATAAGGTTTAA

- Restriction Sites:** Sgfl-Mlul
- ACCN:** NM_001271963
- Insert Size:** 1599 bp
- OTI Disclaimer:** Our molecular clone sequence data has been matched to the reference identifier above as a point of reference. Note that the complete sequence of our molecular clones may differ from the sequence published for this corresponding reference, e.g., by representing an alternative RNA splicing form or single nucleotide polymorphism (SNP).
- OTI Annotation:** Clone contains native stop codon, and expresses the complete ORF without any c-terminal tag.
- 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_001271963.1](#), [NP_001258892.1](#)

RefSeq Size: 2459 bp

RefSeq ORF: 1599 bp

Locus ID: 14049

UniProt ID: [O08575](#)

Cytogenetics: 2 85.83 cM

Gene Summary: This gene encodes a member of the eyes absent protein family. The encoded protein is a tyrosine phosphatase which acts as a transcriptional activator during development. Alternative splicing results in multiple transcript variants. [provided by RefSeq, Dec 2012]
Transcript Variant: This variant (3) differs in the 5' UTR compared to variant 1. Variants 1, 2, and 3 encode the same protein. Sequence Note: This RefSeq record was created from transcript and genomic sequence data because no single transcript from the same strain was available for the full length of the gene. The extent of this transcript is supported by transcript alignments and orthologous data.