



B6.Cg^{tm1.1(APOE*4)Aduj}App^{em2Aduj}Trem2^{em1Aduj}/J

Stock No: 031722 | App KO/APOE4/Trem2*R47H

◆ Congenic, Targeted Mutation, Endonuclease-Mediated Mutation



CRYORECOVERY

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Typically mice are recovered in 10-14 weeks. Contact Customer Service to place an order or for more information.

Overview



MODEL-AD

Model Organism Development & Evaluation for Late-Onset Alzheimer's Disease

Also Known As: App KO/APOE4/Trem2*R47H

This triple mutant strain carries a humanized ApoE knock-in mutation (sequence coding for isoform E4), a CRISPR/cas9-generated 94bp deletion in exon 14 of the *App* gene and a CRISPR/cas9-generated R47H point mutation of the *Trem2* gene. These mice may be suitable for use in studies related to Alzheimer's disease, lipoproteins, arteriosclerosis, and coronary heart disease.

Donating Investigator

Mike Sasner, The Jackson Laboratory

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GENETIC OVERVIEW

Genetic Background

Generation

Trem2^{em1A}diuj

Allele Type	Gene Symbol	Gene Name
Endonuclease-mediated (Humanized sequence)	<i>Trem2</i>	triggering receptor expressed on myeloid cells 2

App^{em2Adiuj}

Allele Type	Gene Symbol	Gene Name
Endonuclease-mediated (Not Specified)	<i>App</i>	amyloid beta (A4) precursor protein

*Apoe^{tm1.1(APOE*4)Adiuj}*

Allele Type	Gene Symbol	Gene Name
Targeted (Inserted expressed sequence, Humanized sequence)	<i>ApoE</i>	apolipoprotein E

[VIEW GENETICS](#)

RESEARCH APPLICATIONS

- Cardiovascular Research
- Research Tools
- Neurobiology Research
- Metabolism Research

[VIEW ALL RESEARCH APPLICATIONS](#)

BASE PRICE

Starting at:

\$2,854.50 Domestic price Cryo Recovery

[VIEW PRICE LIST](#)

➔ Details

➔ Detailed Description

This triple mutant strain carries a humanized ApoE knock-in allele, in which exons 2, 3 and most of exon 4 of the mouse *ApoE* gene were replaced by human *APOE4* gene sequence including exons 2, 3 and 4 (and some 3' UTR sequence); a mutant allele of the *App* gene containing a 94bp deletion in exon 14; and a knock-in of a point mutation into mouse *Trem2*, triggering receptor expressed on myeloid cells 2, gene containing a R47H point mutation, with two silent mutations. The targeted *ApoE* gene encodes apolipoprotein E, which is important in lipoprotein metabolism and cardiovascular disease as well as Alzheimer's disease, immunoregulation and cognition. The targeted *App* gene encodes amyloid beta precursor protein, a transmembrane cell surface receptor that is cleaved by secretases. Mutations in this gene have been associated with Alzheimer's disease. The targeted *Trem2* gene encodes a protein that is part of a receptor signaling complex with TYRO protein tyrosine kinase binding protein, and that activates macrophages and dendritic cells during immune responses. The *TREM2* R47H mutation is a missense mutation in exon 2 that is one of the strongest genetic risk factors for late-onset Alzheimer's disease. Mice that are homozygous for the *ApoE^{tm1.1(APOE*4)Adiuj}* and *Trem2^{em1Adiuj}*

alleles, and heterozygous for the *App*^{em2^AAdiuj} allele are viable and fertile. Homozygous viability/fertility has not been tested for the *App*^{em2^AAdiuj} allele (June 2018). As the mice are characterized, we will modify the strain description and add phenotype data. Of note, in brains of mice homozygous for the *Trem2*^{em1^AAdiuj} allele (and not carrying any other mutant alleles), expression of both transcripts of *Trem2* is decreased by about 50%. Mice expressing the *Trem2* R47H mutation also express a novel splice variant with a deletion of 119bp at the 5' end of exon 2, due to a cryptic splice acceptor site in exon 2 (see Stock No. [027918](#)).

Important note about App isoforms and exon numbering: The App-201 isoform (695 aa protein) is encoded by 16 exons, of which the Abeta sequence is encoded by exon 14. The App-206 isoform (770 aa protein) is encoded by 18 exons, of which the Abeta sequence is encoded by exon 16.

+ Development

+ Expression Data

+ Control Suggestions

- Genetics

+ *Trem2*^{em1^AAdiuj}

+ *App*^{em2^AAdiuj}

+ *ApoE*^{tm1.1(APOE*4)Adiuj}

- Disease/Phenotype

+ Disease Terms

+ Research Areas By Genotype

+ Mammalian Phenotype Terms by Genotype

+ References

- Technical Support

C H A T O  F L I N E

C O N T A C T T E C H N I C A L S U P P O R T

Genotyping Protocols

End Point Analysis: *Trem2*^{em1^AAdiuj}-EP

Standard PCR: *ApoE*^{tm1.1(APOE*4)Adiuj}

Probe: *App*^{em2^AAdiuj} (91bp del)

[Genotyping resources and troubleshooting](#)

Breeding Considerations

When maintaining a live colony, mice homozygous for the $Apoe^{tm1.1(APOE^*4)Adiuj}$ and $Trem2^{em1Adiuj}$ alleles, and heterozygous for the $App^{em2Adiuj}$ allele may be bred.

Homozygous viability/fertility has not been tested for the $App^{em2Adiuj}$ allele (June 2018).

As the mice are characterized, we will modify the strain description if necessary and add data related to viability and fertility.

[Additional Breeding and Husbandry Support](#)

Mating System

homozygous $Apoe^{tm1.1(APOE^*4)Adiuj}$, homozygous $Trem2^{em1Adiuj}$, heterozygous $App^{em2Adiuj}$ x homozygous $Apoe^{tm1.1(APOE^*4)Adiuj}$, homozygous $Trem2^{em1Adiuj}$, Wildtype $App^{em2Adiuj}$ and reciproca

Citation

When using the $App^{em2Adiuj}/Apoe^{tm1.1(APOE^*4)Adiuj}/Trem2^{em1Adiuj}$ mouse strain in a publication, please [cite the originating article\(s\)](#) and include JAX stock #031722 in your Materials and Methods section.

Animal Health Reports

[Facility Barrier Level Descriptions](#)

Production of mice from cryopreserved embryos or sperm occurs in a maximum barrier room, G200

➔ Pricing & Availability



Cryo Recovery

Typically mice are recovered in 10-14 weeks. Contact Customer Service to place an order or for more information.

Domestic International

Pricing effective for USA, Canada and Mexico shipping destinations

Cryorecovery - Domestic Pricing

SERVICE	GENOTYPE	PRICE
Cryo Recovery	Homozygous for $Apoe^{tm1.1(APOE^*4)Adiuj}$ and $Trem2^{em1Adiuj}$ Heterozygous or wildtype for $App^{em2Adiuj}$	\$2,854.50

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Terms of Use

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