Resource Summary Report

Generated by FDI Lab - SciCrunch.org on Apr 15, 2025

LP/J

RRID:IMSR_JAX:000676 Type: Organism

Proper Citation

RRID:IMSR_JAX:000676

Organism Information

URL: https://www.jax.org/strain/000676

Proper Citation: RRID:IMSR_JAX:000676

Description: Mus musculus with name LP/J from IMSR.

Species: Mus musculus

Synonyms: LP. line 11 pied

Notes: gene symbol note: aryl-hydrocarbon receptor|cytochrome c oxidase subunit 7A2 like|MX dynamin-like GTPase 1|cadherin related 23 (otocadherin)|beta-2 microglobulin|disrupted in schizophrenia 1|endothelin receptor type B|hemoglobin beta chain complex; inbred strain: Ahr|Cox7a2l|Mx1|Cdh23|B2m|Disc1|Ednrb|Hbb

Affected Gene: aryl-hydrocarbon receptor|cytochrome c oxidase subunit 7A2 like|MX dynamin-like GTPase 1|cadherin related 23 (otocadherin)|beta-2 microglobulin|disrupted in schizophrenia 1|endothelin receptor type B|hemoglobin beta chain complex

Genomic Alteration: d variant|long|myxovirus susceptibility 1|age related hearing loss 1|a variant|deletion|piebald|d

Catalog Number: JAX:000676

Database: International Mouse Resource Center IMSR, JAX

Database Abbreviation: IMSR

Availability: live

Alternate IDs: IMSR_JAX:676

Organism Name: LP/J

Record Creation Time: 20230509T193230+0000

Record Last Update: 20250412T090207+0000

Ratings and Alerts

No rating or validation information has been found for LP/J.

No alerts have been found for LP/J.

Data and Source Information

Source: Integrated Animals

Source Database: International Mouse Resource Center IMSR, JAX

Usage and Citation Metrics

We found 59 mentions in open access literature.

Listed below are recent publications. The full list is available at FDI Lab - SciCrunch.org.

Durán A, et al. (2023) A Mouse Systems Genetics Approach Reveals Common and Uncommon Genetic Modifiers of Hepatic Lysosomal Enzyme Activities and Glycosphingolipids. International journal of molecular sciences, 24(5).

Oestereicher MA, et al. (2023) Comprehensive ECG reference intervals in C57BL/6N substrains provide a generalizable guide for cardiac electrophysiology studies in mice. Mammalian genome : official journal of the International Mammalian Genome Society, 34(2), 180.

Khan AH, et al. (2023) Genetic pathways regulating the longitudinal acquisition of cocaine self-administration in a panel of inbred and recombinant inbred mice. Cell reports, 42(8), 112856.

Sottoriva K, et al. (2022) A Notch/IL-21 signaling axis primes bone marrow T cell progenitor expansion. JCI insight, 7(9).

Zhou W, et al. (2022) DNA methylation dynamics and dysregulation delineated by high-throughput profiling in the mouse. Cell genomics, 2(7).

Shi LJ, et al. (2022) Genetic Evidence for a Causal Relationship between Hyperlipidemia and

Type 2 Diabetes in Mice. International journal of molecular sciences, 23(11).

Shi LJ, et al. (2022) Genetic Connection between Hyperglycemia and Carotid Atherosclerosis in Hyperlipidemic Mice. Genes, 13(3).

Molendijk J, et al. (2022) Proteome-wide systems genetics identifies UFMylation as a regulator of skeletal muscle function. eLife, 11.

Sheppard K, et al. (2022) Stride-level analysis of mouse open field behavior using deeplearning-based pose estimation. Cell reports, 38(2), 110231.

Munz M, et al. (2021) In silico candidate variant and gene identification using inbred mouse strains. PeerJ, 9, e11017.

Seemiller LR, et al. (2021) Genetic background determines behavioral responses during fear conditioning. Neurobiology of learning and memory, 184, 107501.

Mooney-Leber SM, et al. (2021) Genetic Differences in Dorsal Hippocampus Acetylcholinesterase Activity Predict Contextual Fear Learning Across Inbred Mouse Strains. Frontiers in psychiatry, 12, 737897.

Ren M, et al. (2020) Transcription factor p73 regulates Th1 differentiation. Nature communications, 11(1), 1475.

Strattan E, et al. (2019) Mast Cells Are Mediators of Fibrosis and Effector Cell Recruitment in Dermal Chronic Graft-vs.-Host Disease. Frontiers in immunology, 10, 2470.

Yang C, et al. (2019) Genome-wide association study using diversity outcross mice identified candidate genes of pancreatic cancer. Genomics, 111(6), 1882.

Zhou Y, et al. (2019) Integrative system genetic analysis reveals mRNA-IncRNA network associated with mouse spontaneous lung cancer susceptibility. Oncotarget, 10(3), 339.

Huang Q, et al. (2019) Delivering genes across the blood-brain barrier: LY6A, a novel cellular receptor for AAV-PHP.B capsids. PloS one, 14(11), e0225206.

Zimmerman H, et al. (2019) Interfrontal Bone Among Inbred Strains of Mice and QTL Mapping. Frontiers in genetics, 10, 291.

Yu T, et al. (2019) The piRNA Response to Retroviral Invasion of the Koala Genome. Cell, 179(3), 632.

Fu YY, et al. (2019) T Cell Recruitment to the Intestinal Stem Cell Compartment Drives Immune-Mediated Intestinal Damage after Allogeneic Transplantation. Immunity, 51(1), 90.