

Resource Summary Report

Generated by ASWG on Sep 20, 2026

BUB/BnJ

RRID:IMSR_JAX:000653

Type: Organism

Proper Citation

RRID:IMSR_JAX:000653

Organism Information

URL: <https://www.jax.org/strain/000653>

Proper Citation: RRID:IMSR_JAX:000653

Description: Mus musculus with name BUB/BnJ from IMSR.

Species: Mus musculus

Synonyms: BUB/BnJ-Pde6b

Notes: gene symbol note: MX dynamin-like GTPase 1|cadherin related 23 (otocadherin)|adhesion G protein-coupled receptor V1|aryl-hydrocarbon receptor|interleukin 3 receptor; alpha chain|cytochrome c oxidase subunit 7A2 like|beta-2 microglobulin|phosphodiesterase 6B; cGMP; rod receptor; beta polypeptide|latent transforming growth factor beta binding protein 4|hemoglobin beta chain complex; inbred strain: Mx1|Cdh23|Adgrv1|Ahr|Il3ra|Cox7a2|B2m|Pde6b|Ltbp4|Hbb

Affected Gene: MX dynamin-like GTPase 1|cadherin related 23 (otocadherin)|adhesion G protein-coupled receptor V1|aryl-hydrocarbon receptor|interleukin 3 receptor; alpha chain|cytochrome c oxidase subunit 7A2 like|beta-2 microglobulin|phosphodiesterase 6B; cGMP; rod receptor; beta polypeptide|latent transforming growth factor beta binding protein 4|hemoglobin beta chain complex

Genomic Alteration: myxovirus susceptibility 1|age related hearing loss 1|frings|b-2 variant|mutation 1|long|a variant|retinal degeneration 1|Duchenne modifier 1 resistant|d

Catalog Number: JAX:000653

Database: JAX Mice and Services

Database Abbreviation: JAX

Availability: embryo

Organism Name: BUB/BnJ

Record Creation Time: 20250513T053620+0000

Record Last Update: 20260919T054809+0000

Ratings and Alerts

No rating or validation information has been found for BUB/BnJ.

No alerts have been found for BUB/BnJ.

Data and Source Information

Source: [Integrated Animals](#)

Source Database: JAX Mice and Services

Usage and Citation Metrics

We found 3 mentions in open access literature.

Listed below are recent publications. The full list is available at [ASWG](#) .

Ren W, et al. (2025) A tandem repeat atlas for the genome of inbred mouse strains: A genetic variation resource. iScience, 28(11), 113703.

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

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