

# Resource Summary Report

Generated by [FDI Lab - SciCrunch.org](https://fdi-lab.sci-crunch.org/) on Apr 12, 2025

## HmtPhenome

RRID:SCR\_017289

Type: Tool

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### Proper Citation

HmtPhenome (RRID:SCR\_017289)

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### Resource Information

**URL:** <https://www.hmtphenome.uniba.it>

**Proper Citation:** HmtPhenome (RRID:SCR\_017289)

**Description:** Collection of data about variants, genes, phenotypes and diseases involved in mitochondrial functionality. Users can search for variant position, gene, phenotype or disease and retrieve all related information through integrated network of biological entities.

**Resource Type:** data or information resource, data processing software, data visualization software, software resource, service resource, network graph visualization software, software application, database

**Defining Citation:** [DOI:10.1101/660282](https://doi.org/10.1101/660282)

**Keywords:** mitochondria, variant, gene, function, phenotype, data

**Funding:**

**Availability:** Free, Freely available

**Resource Name:** HmtPhenome

**Resource ID:** SCR\_017289

**Record Creation Time:** 20220129T080334+0000

**Record Last Update:** 20250412T060108+0000

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### Ratings and Alerts

No rating or validation information has been found for HmtPhenome.

No alerts have been found for HmtPhenome.

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## Data and Source Information

**Source:** [SciCrunch Registry](#)

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## Usage and Citation Metrics

We have not found any literature mentions for this resource.