

# Resource Summary Report

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## HmtPhenome

RRID:SCR\_017289

Type: Tool

### Proper Citation

HmtPhenome (RRID:SCR\_017289)

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

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

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

**Availability:** Free, Freely available

**Resource Name:** HmtPhenome

**Resource ID:** SCR\_017289

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

**Record Last Update:** 20260728T043531+0000

### Ratings and Alerts

No rating or validation information has been found for HmtPhenome.

No alerts have been found for HmtPhenome.

### 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.