

# SUPPLEMENTARY MATERIALS for Mitochondrial DNA genome variation in the Swedish population

**SND-ID:** 2024-357. **Version:** 1. **DOI:** <https://doi.org/10.57804/mfyp-ea25>

## Download data

Figure S1.pdf (348.21 KB)

Figure S2.pdf (289.52 KB)

Figure S3.pdf (402.76 KB)

Table S1.xlsx (50.11 KB)

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2024-357-1.zip (~1.07 MB)

## Citation

Sturk-Andreaggi, K. (2022) SUPPLEMENTARY MATERIALS for Mitochondrial DNA genome variation in the Swedish population (Version 1) [Data set]. Uppsala University. Available at: <https://doi.org/10.57804/mfyp-ea25>

## Creator/Principal investigator(s)

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## Research principal

[Uppsala University](#)

## Description

The data consists of:

- Table S1. The haplogroup breakdown for the 934 SweGen haplotypes included in the final mitochondrial genome dataset.
- Figure S1. Graphical description of the average read depths observed in the SweGen dataset.
- Figure S2. The distribution of read depth for the 16,569 positions of the mitochondrial genome based on the average observed in a subset of 100 representative SweGen haplotypes.
- Figure S3. The box-and-whisker plot presents the distribution of average variant frequency for each coverage classification group.

The dataset was originally published in DiVA and moved to SND in 2024.

## Data contains personal data

No

## Language

[English](#)

## Contributor(s)

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## Identifiers

URN: [urn:nbn:se:uu:diva-470698](https://nbn-resolving.org/urn:nbn:se:uu:diva-470698)

## Research area

[Genetics](#) (Standard för svensk indelning av forskningsämnen 2011)

[Forensic science](#) (Standard för svensk indelning av forskningsämnen 2011)

## Keywords

[Biology with specialisation in human evolution and genetics](#)

## Accessibility level

Access to data through SND  
Data are freely accessible

## Use of data

[Things to consider when using data shared through SND](#)

## Versions

Version 1. 2022-03-28

## This resource has the following relations

Is referenced by [Mitochondrial DNA genome variation in the Swedish population:](#)

## Download metadata

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[DDI 2.5](#)

[DDI 3.3](#)

[DCAT-AP-SE 2.0](#)

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