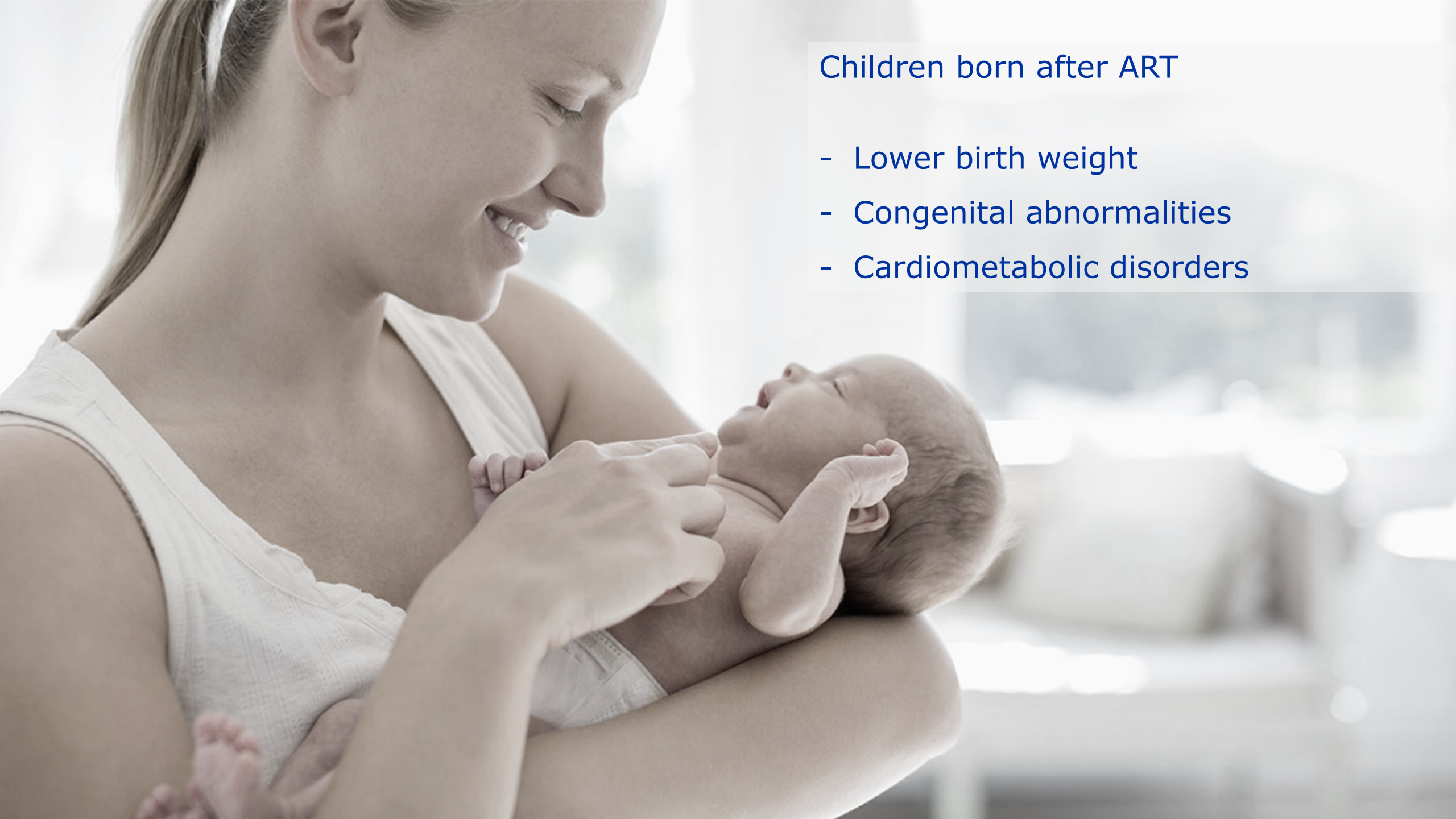


REPRODUCTION  
& GENETICS  
RESEARCH GROUP

# Mitochondrial DNA variants in individuals born after ART compared to spontaneously conceived peers

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## Children born after ART

- Lower birth weight
- Congenital abnormalities
- Cardiometabolic disorders

# Mitochondrial DNA mutations

**Lower birth weight**

Cancer

**β-cell failure**

Epilepsy

Ataxia

Neuropathy

**Ischemia**

Rapid ageing

**Atherogenesis**

Ophtalmoplegia

Myopathy

Increase in cholesterol synthesis

**Type 2 Diabetes**

Pancytopenia

**Infertility**

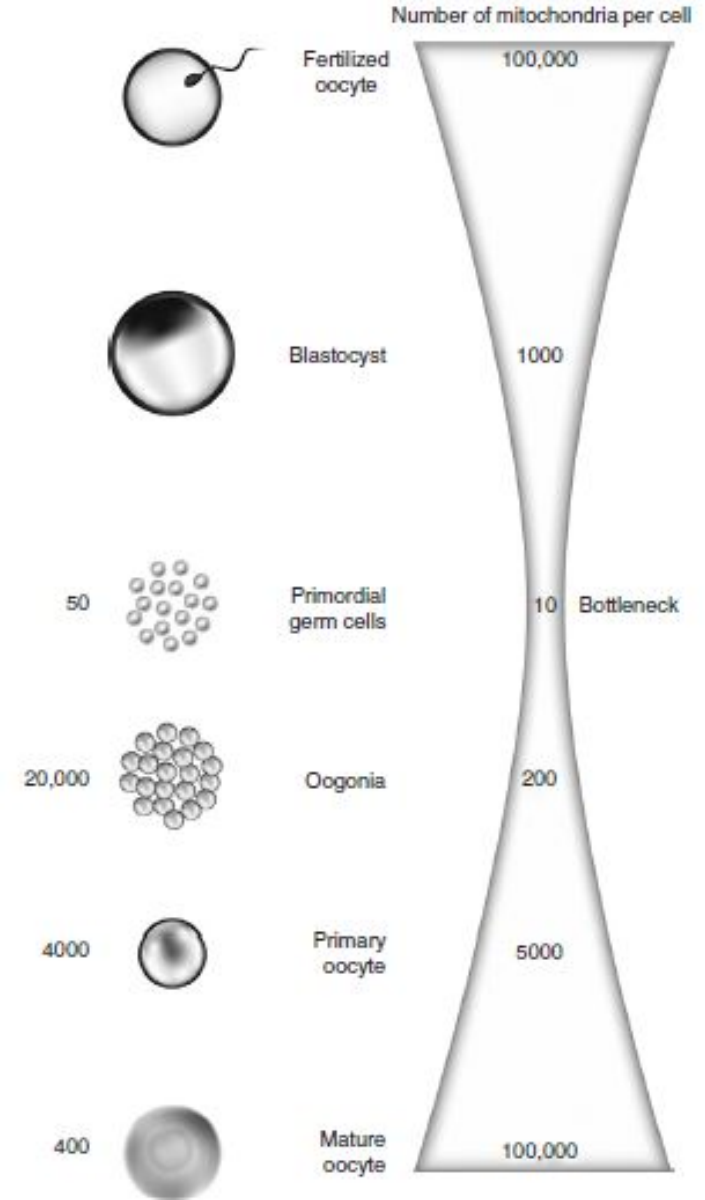
# Controlled ovarian stimulation during oogenesis

## Cellular stress

- Replication stress ↑
- mtDNA mutation rate ↑

## Recruitment of multiple follicles

- Prevents dominant follicle
- Includes recruitment of 'less fit' oocytes with mtDNA mutations?



CHILDREN BORN AFTER ART CARRY  
MORE VARIANTS IN THEIR MITOCHONDRIAL DNA  
AND AT A HIGHER LOAD



# Material & Methods

## Blood

- 116 ICSI
- 65 Control



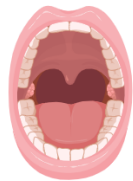
## Placentas

- 28 ICSI
- 27 Control



## Saliva

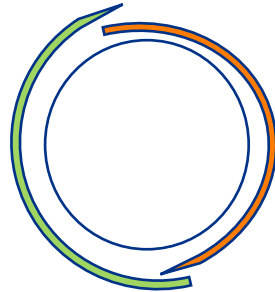
- 107 IVF
- 6 Control



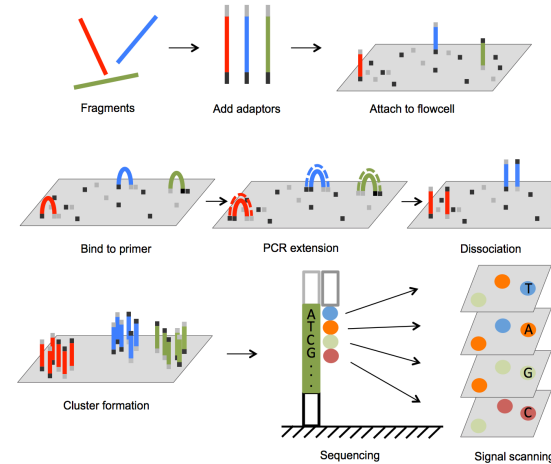
## Buccal swab

- 63 Control

## PCR amplification



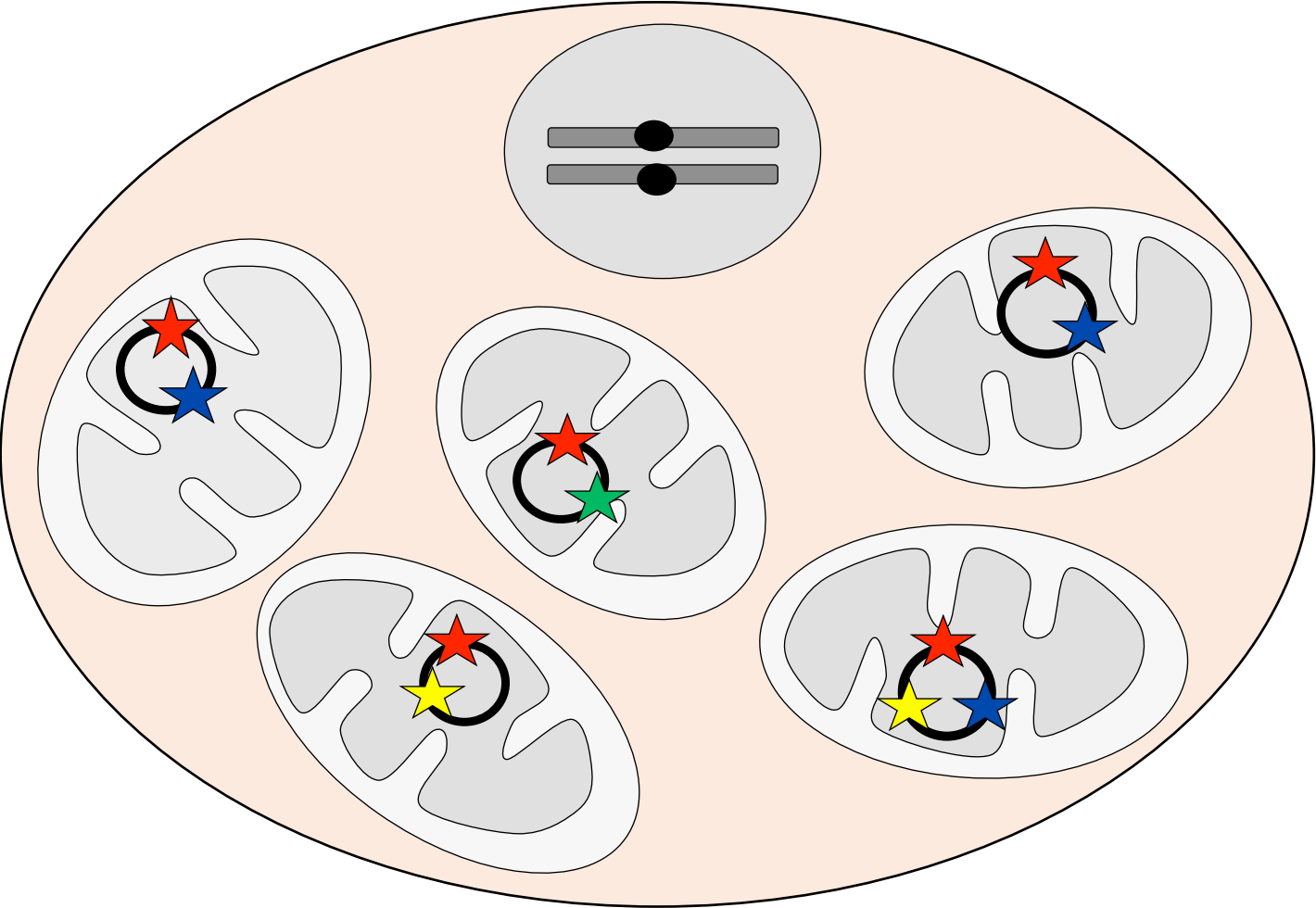
## Sequencing



## Data analysis

| T42            |         |               |             |         |                |           |           |        |        |
|----------------|---------|---------------|-------------|---------|----------------|-----------|-----------|--------|--------|
| Homoplasmies   |         |               |             |         |                |           |           |        |        |
| mDNA server    |         |               |             |         |                |           | CLCBio    | MuTect |        |
| Position       | Variant | Gene          | AAC         | Mutpred | Selection      | Haplotype |           |        |        |
| 1438           | A>G     | MT-RNR1       | .           | .       | yes (10.0)     |           | 99,958    | 1.000  |        |
| 15326          | A>G     | MT-CYB T194A  | 0.452       | 0.395   | yes (10.0)     |           | 99,854    | 0.998  |        |
| 4769           | A>G     | MT-ND2        | .           | .       | yes (8.8)      |           | 99,819    | 0.148  |        |
| 750            | A>G     | MT-RNR1       | .           | .       | yes (10.0)     |           | 99,881    | 0.999  |        |
| 8860           | A>G     | MT-ATP6 T112A | 0.369       | 0.287   | yes (10.0)     |           | 99,922    | 0.998  |        |
| 13865          | C>T     | MT-CYB        | .           | .       | no (n.d.)      |           | 99,485    | 0.996  |        |
| 16399          | A>G     | MT-DL01       | .           | .       | no (5.0)       |           | 99,772    | 0.998  |        |
| 3107           | N>T     | MT-RNR1       | .           | .       | no (n.d.)      |           |           |        |        |
| 9575           | G>A     | MT-CO3        | .           | .       | no (8.1)       |           | 99,574    | 0.997  |        |
| 9932           | G>A     | MT-CO3        | .           | .       | no (5.9)       |           | 99,354    | 0.995  |        |
| 16519          | T>C     | MT-DL01       | .           | .       | hotspot (n.d.) |           | 99,82     | 0.998  |        |
| Heteroplasmies |         |               |             |         |                |           |           |        |        |
|                |         |               |             |         |                |           | Frequency |        |        |
|                |         |               |             |         |                |           | mDNA seq  | CLCBio | MuTect |
| 263            | A>G     | SNV           | HV; MT-C HV |         |                |           | 58        | 58,962 | 0.598  |

# Homoplasmy versus Heteroplasmy



Homoplasmic variant

★ 100%

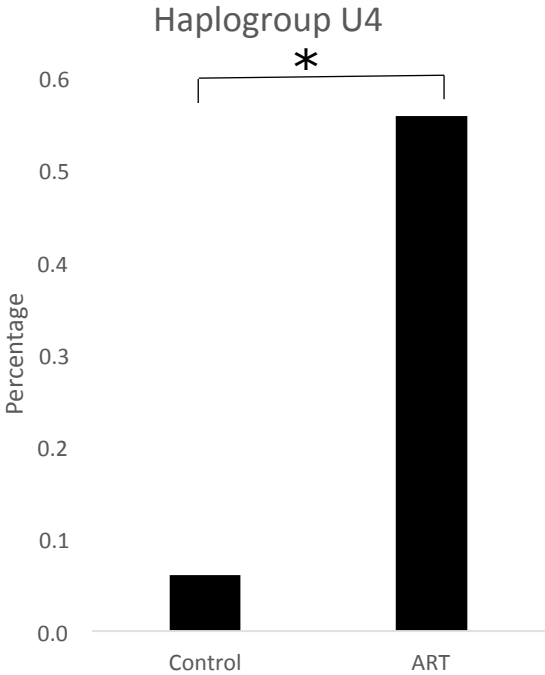
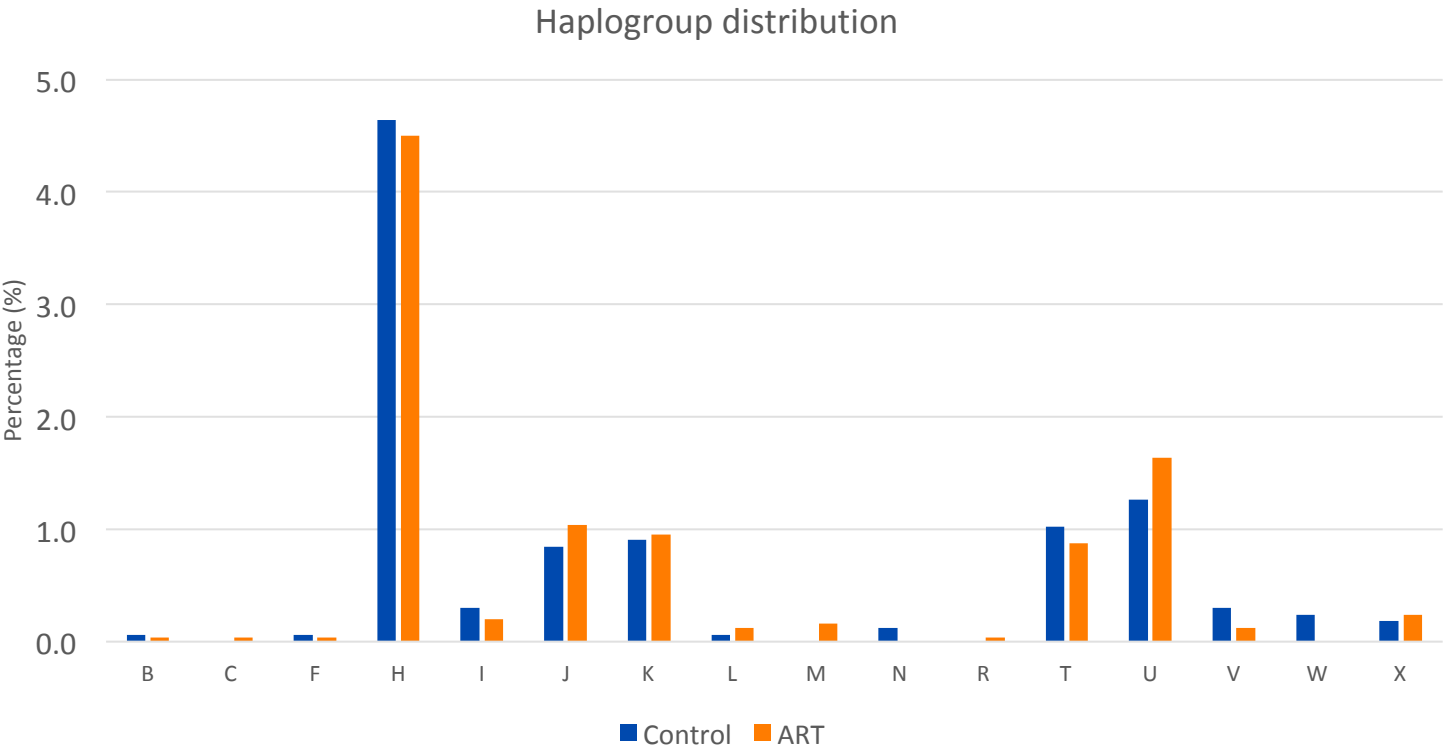
Heteroplasmic variants

★ 60%

★ 40%

★ 20%

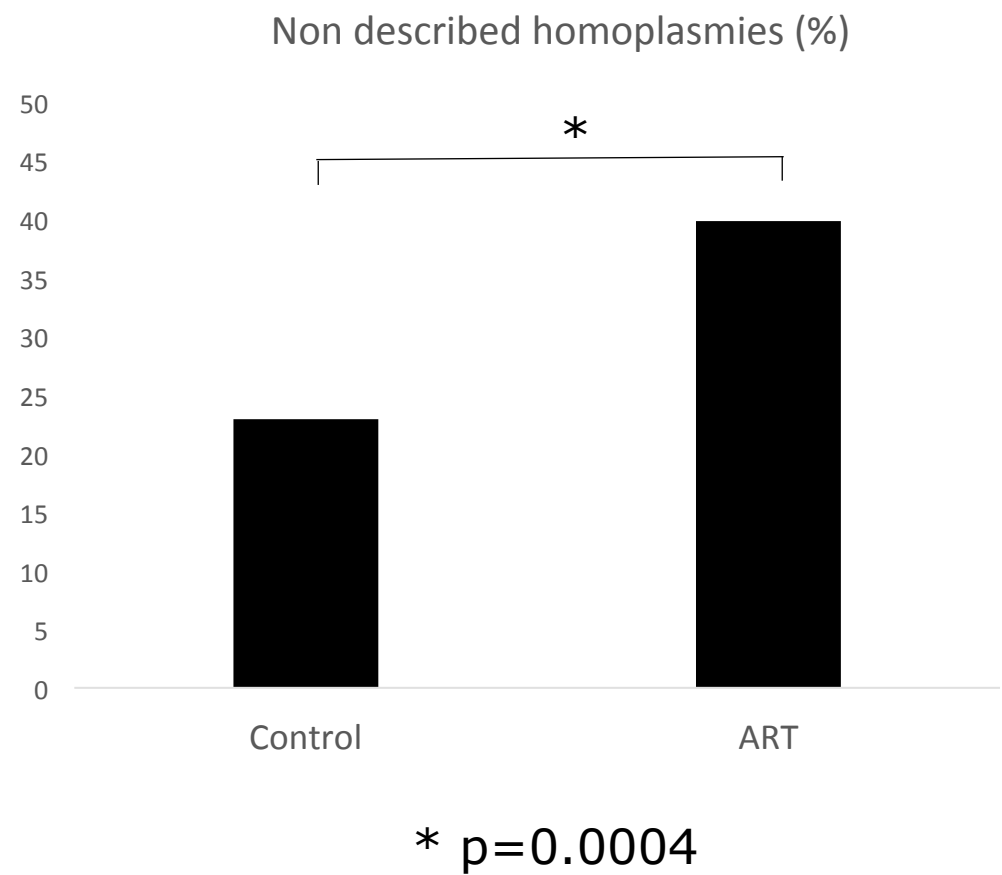
# Subhaplogroup U4 is over-represented in the ART group



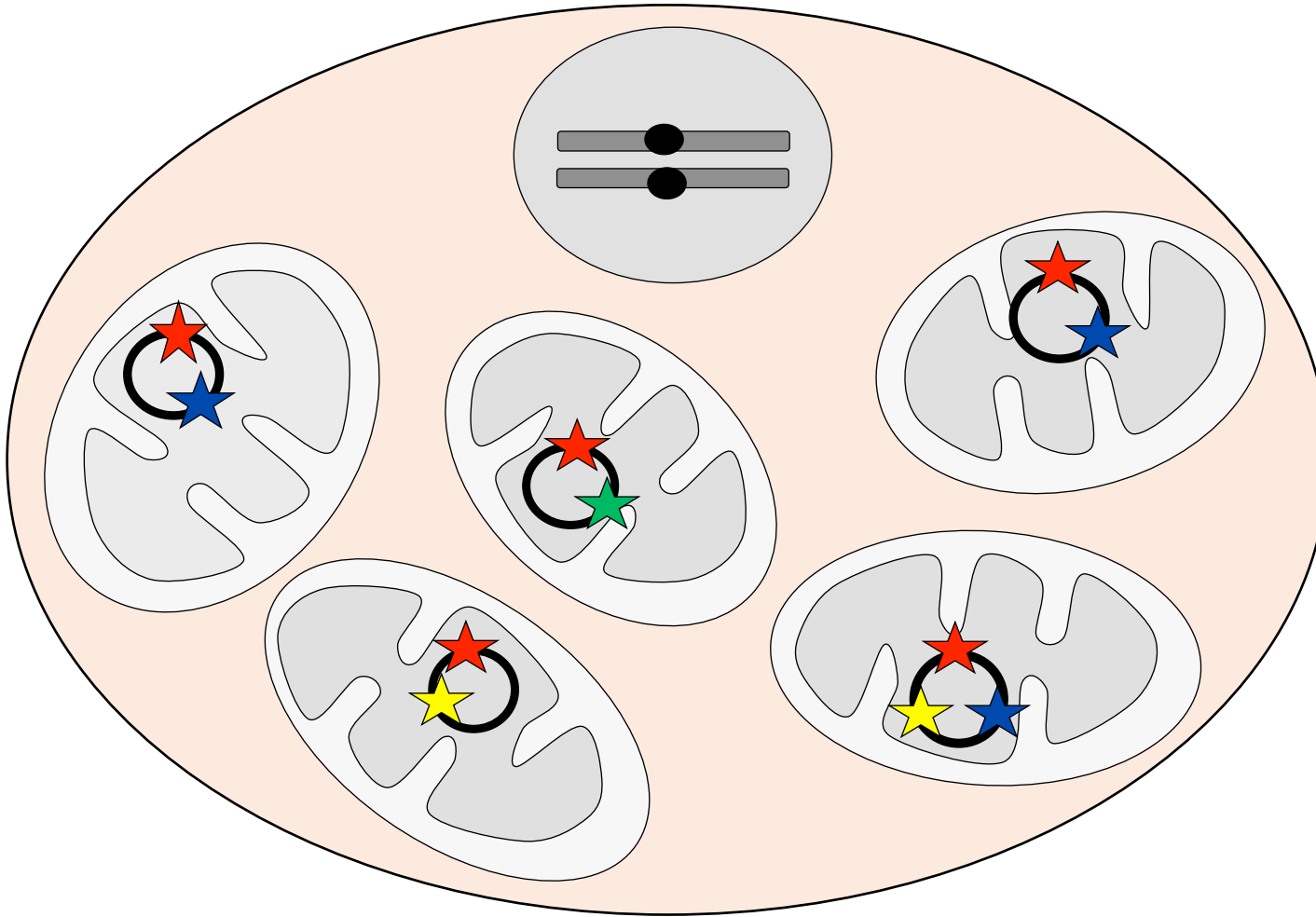
\* p=0.006



# More unique variants in the ART group



# Homoplasmy versus Heteroplasmy



Homoplasmic variant

★ 100%

Heteroplasmic variants

★ 60%

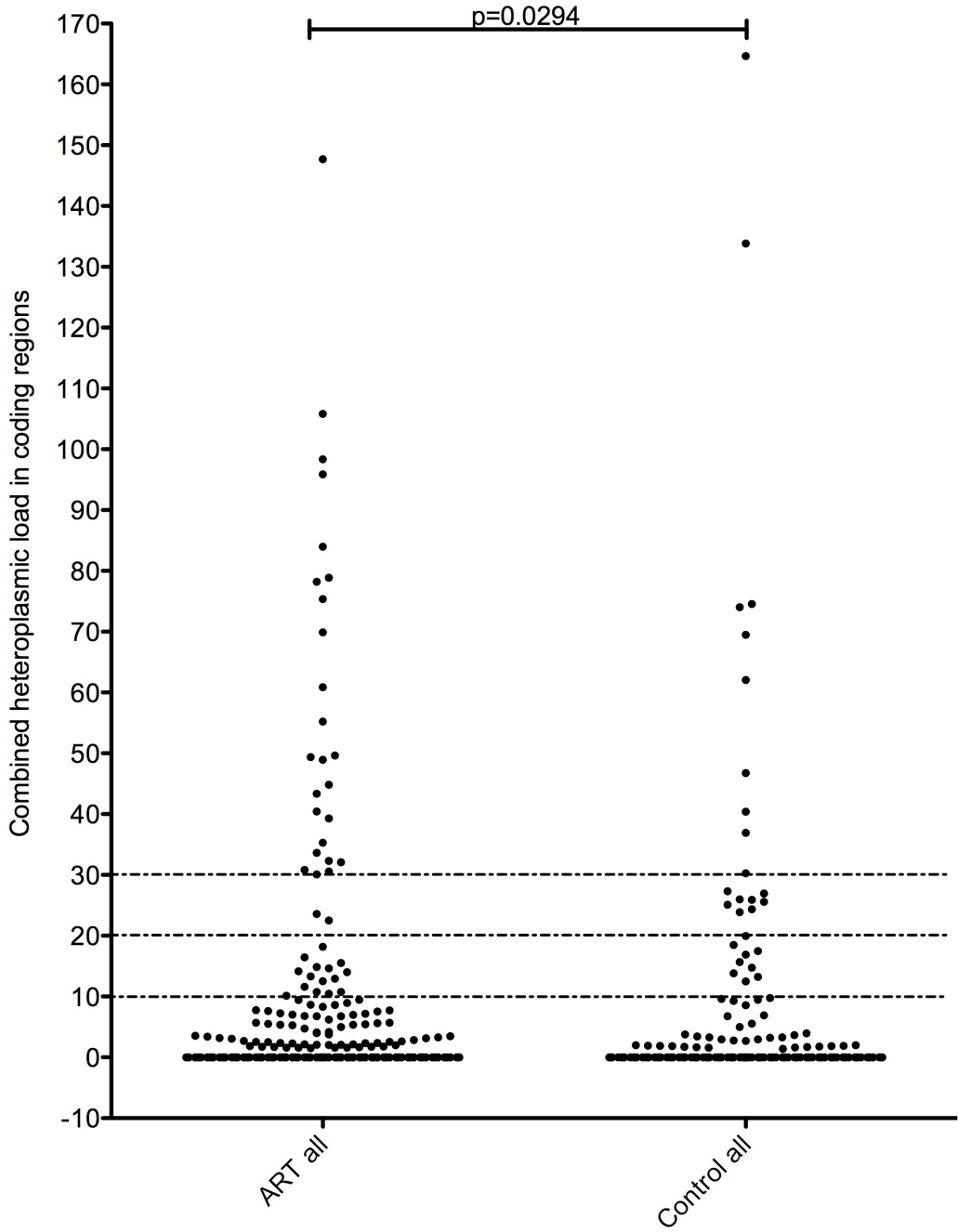
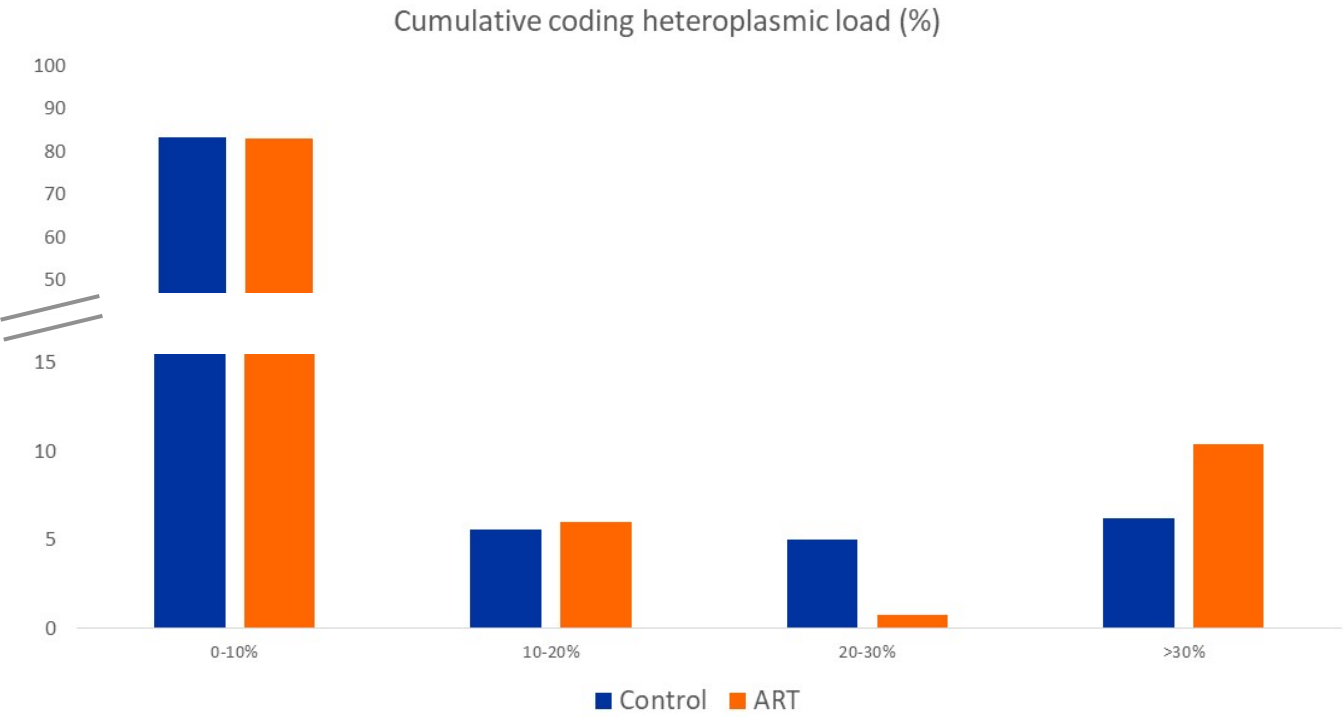
★ 40%

★ 20%

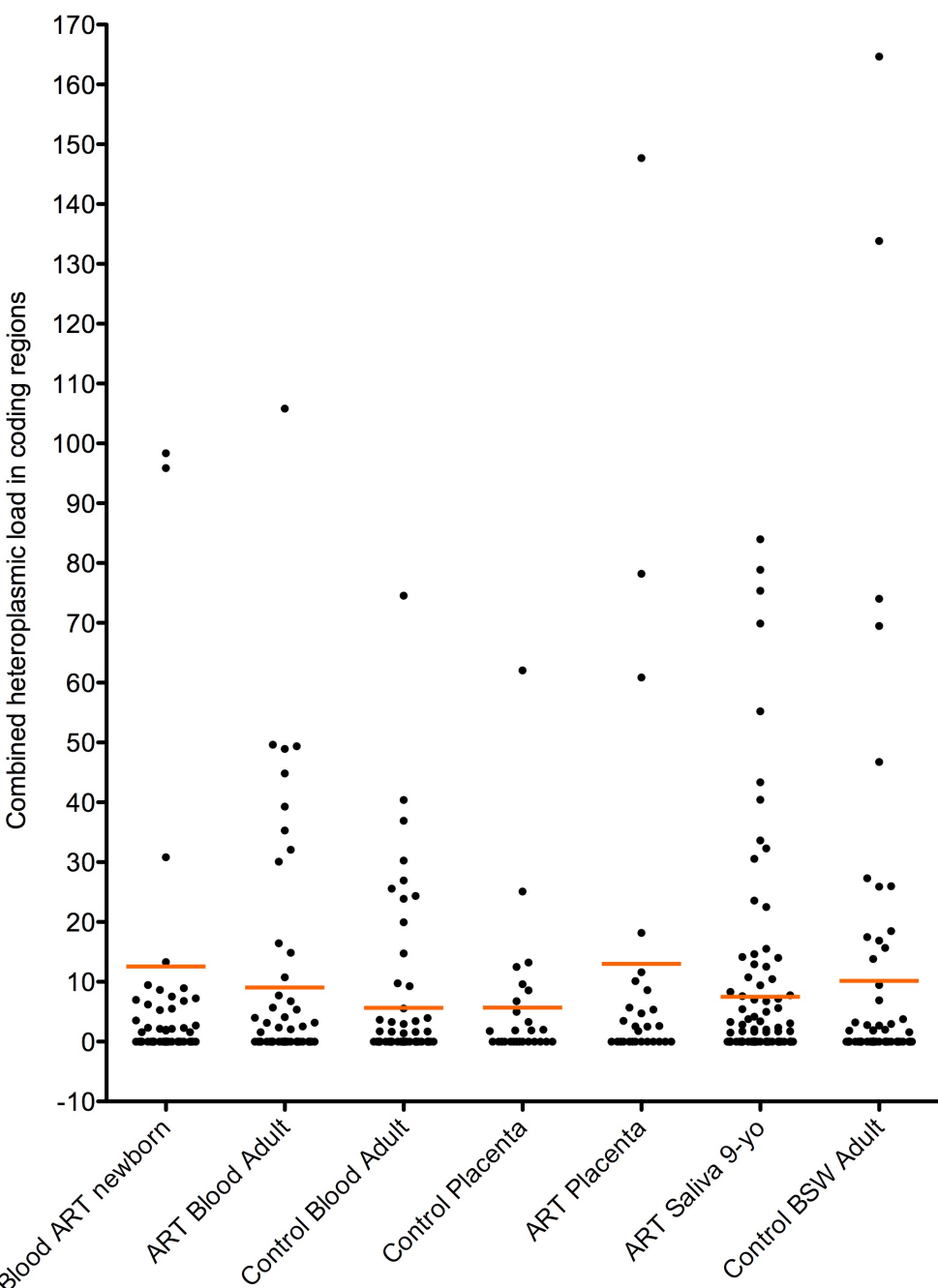
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= Cumulative load

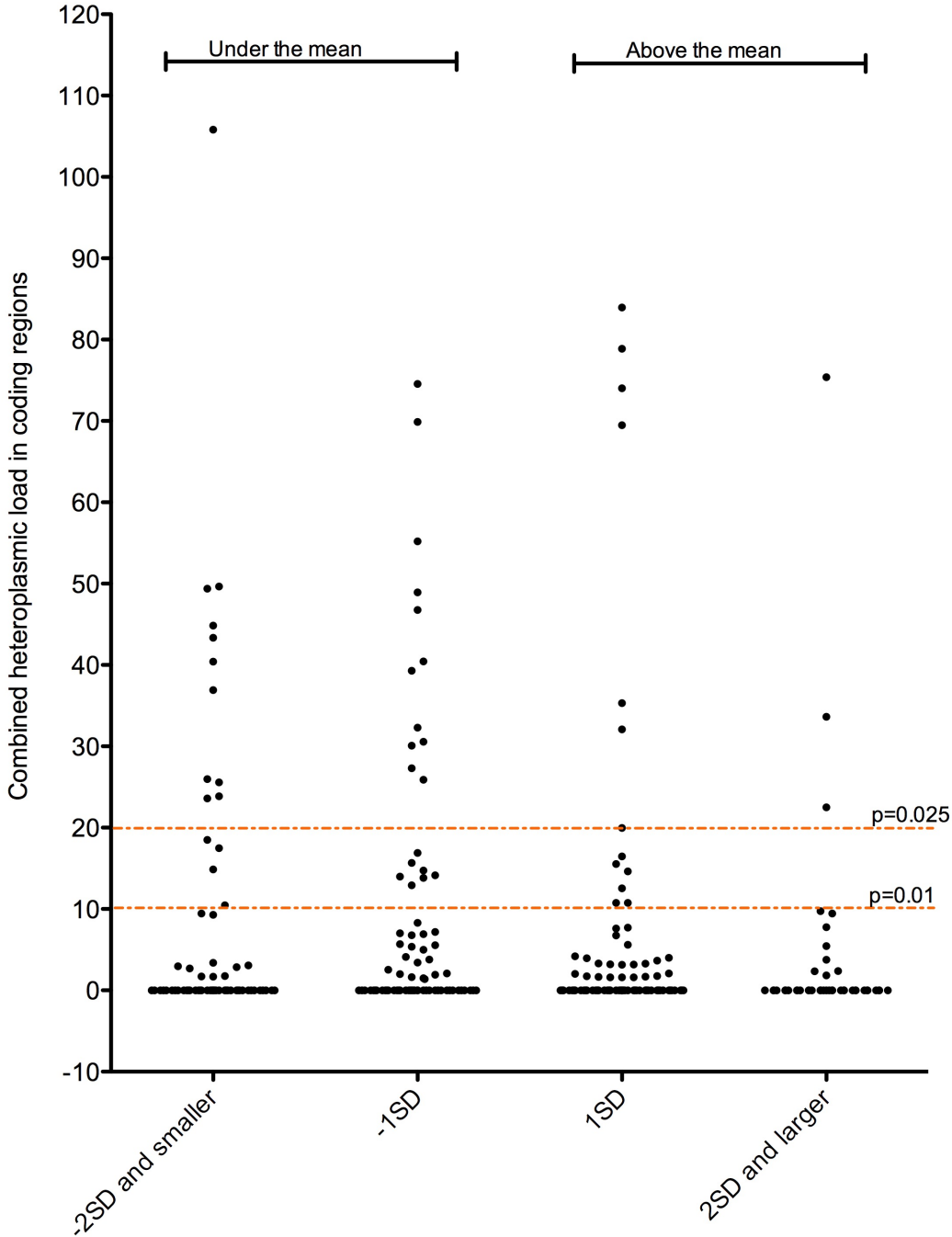
# Increased cumulative heteroplasmic load in the coding region in ART



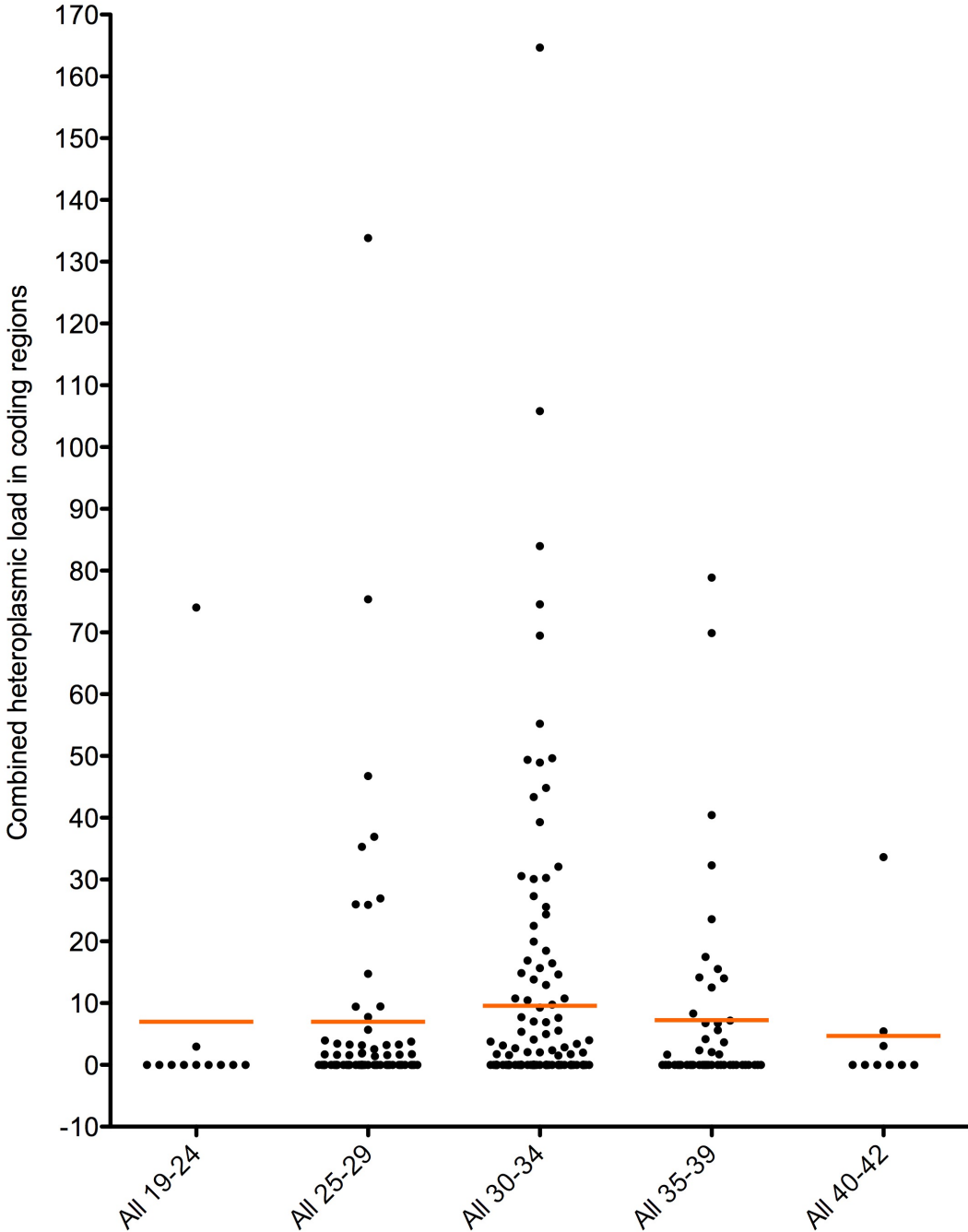
Increased cumulative heteroplasmic load in the coding region in ART



Increased cumulative loads seem to correlate with a lower birth weight



Maternal age doesn't seem to correlate with a higher cumulative load



# Conclusion

- Subhaplogroup U4 is over-represented in the ART group: a link to maternal infertility?
- Increase in non-described homoplasmies:  
→ mostly synonymous: origin?
- Increased cumulative coding heteroplasmic variant load
- Higher cumulative load appears to correlate with lower birth weight
- Maternal age doesn't seem to influence mtDNA variant load

# Acknowledgements



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