# Placental and blood mitochondrial DNA heteroplasmies in women living with HIV and their association with preterm birth



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Control

## OBJECTIVES

- To investigate the **mitochondrial DNA heteroplasmies** present in the placental tissue  $\bullet$ of a cohort of women living with and without HIV and determine any association with preterm birth status.
- To determine if previously-identified placental pathogenic mtDNA heteroplasmies are present within other fetal tissues and maternal blood.

#### HYPOTHESES

A greater number of pathogenic mtDNA heteroplasmies will be associated with HIV-

Position

**Fig 5a.** Gene-specific heteroplasmy frequency separated by preterm and term birth groups. COX: Cytochrome c oxidase subunits; CYTB: Cytochrome b; HVI: hypervariable region I ND: NADH Dehydrogenase; NC: non-coding region

Position 310 (HV2) is a **non-pathogenic** haplogroup marker. Levels of heteroplasmy vary from 10% to 95% with nearly every sample having some level of heteroplasmy.

- **positive** and **preterm** birth status.
- Pathogenic mtDNA heteroplasmies present in placental tissue will also be present in maternal blood and tissues.

#### METHODS

Sequenced a total of 69 samples with participants from two different study groups:

- Children and Women: Antiretrovirals and Markers of Aging (CARMA)-PREG
- **E**pigenetics of **P**lacenta in **C**omplications of Pregnancy (EPIC)



**Fig I.** Workflow for mitochondrial whole-genome sequencing (mtWGS): DNA extraction (QIAamp Mini Kit), mtDNA qPCR (LightCycler480), long PCR, gel extraction, mtDNA sequencing (Illumina MiSeq),



Position 2487 (rRNA) is an artifact of alignment. The aforementioned **new** alignment method removes heteroplasmy as this location, while significant pathogenic heteroplasmies remain.

Preterm and term groups differ the most in the HV3 and NADH Dehydrogenase subunit areas.

Living with HIV and not living with HIV groups do not visually differ in heteroplasmy location.

Fig 5b. Gene-specific heteroplasmy frequency separated by participants living with HIV and participants not living with HIV



#### and sequence processing (not pictured)

	<b>Term</b> (n = 29)	<b>Preterm</b> (n = 32)	p-value
Maternal HIV Status, n (%)			1.0000
Living with HIV	12 (41.4)	13 (40.6)	
Not living with HIV	17 (58.6)	19 (59.4)	
Maternal Ethnicity, n (%)			0.1171
African/Caribbean/Black	1 (3.45)	6 (18.6)	
Indigenous	7 (24.1)	5 (15.6)	
White/European	14 (48.3)	8 (25.0)	
Other	1 (3.45)	2 (6.25)	
Missing	6 (20.7)	11 (34.4)	
<b>Delivery Type</b> , n (%)			0.0096
Vaginal	23 (79.3)	13 (40.6)	
Elective C-section	4 (13.8)	14 (43.8)	
Emergency C-section	2 (6.90)	5 (15.6)	
Preeclampsia, n (%)			0.0091
Yes	3 (10.3)	13 (40.6)	
No	26 (89.7)	19 (59.4)	
Maternal Age (years), median [ <u>IQR</u> ] (range)	33.4 [7.20] (21.6, 40.4)	33.8 [7.55] (19.7, 42.9)	0.3515
Gestational Age at Birth (weeks), median [IOR] (range)	39.3 [2.10] (37.1, 42.1)	35.2 [2.55] (24.1, 36.9)	<0.0001

**Fig 2**. Demographics of study participants. The only significant variables are delivery type, preeclampsia, and gestational age at

## SIGNIFICANCE & FUTURE DIRECTIONS

This study will contribute vital information towards understanding of the role of mitochondrial health in pregnancy. It will also support the current body of research on the differences in mitochondrial health experienced by people living with HIV.

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