# Pilot Screening of Cell-Free mtDNA in NIPT: Quality Control, Variant Calling and Haplogroup Determination

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Supplementary Materials
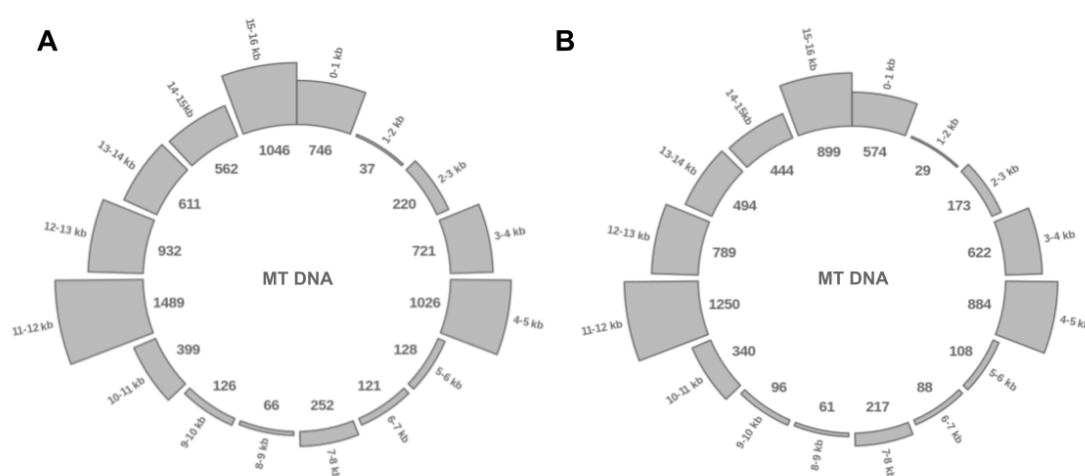
Figure S1. Distribution of variants throughout the mitochondrial genome before (A) and after (B) filtering of homopolymers. Figures outside the circle mark position in the mitochondrial genome (16kb mtDNA has been divided into 16 sections per 1kb), figures in the inner circle represent the number of variants in every section.