

A sequencing we have choose the mtDNA lineages which are specific for populations of northern Asia but which are still underrepresented in the published data sets on complete mtDNA va
which are rarely found in populations of northern A
amples of northern and eastern Asians (including 24
a total of 55 samples representing haplogroups B (n
sequencing was performed using the methodology de

Application Usage
Purpose Analysis
were analyzed using SeqScape v.
Developer
software (Applied Biosystems) and compared with

indexes, the Tajima's D [70] and Fu's FS [71] neut

t (PC) analysis was performed using mtDNA haplog

dimensional scaling (MDS) analysis based on FST s
d other Asian populations around.

mtDNA diversity in western, eastern, central and north

s.
f the complete mtDNA phylogenies of haplogroups B
yloTree [18], were taken into account.

ich we hereby update, follows van Oven and Kayser [18], with several new modifications.

Open

Collection

/SoMeSci/PLoS_methods/

Collection Information

Document

PMC3283723

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| PMC2214211 | 3 weeks ago | 0 | 0 | 0 | 0 |

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regions of Asia.
r haplogroup-diagnostic RFLP markers and subje
d R11 (n=1) were selected (Table S4).

Developer_of
Version_of

Version
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version 3.01 [72].

Developer
tSoft, Inc., USA).

Developer_of
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6.0 software (StatSoft, Inc., USA)

ng now in eastern Europe but descended from we

published previously [4], [5], [20]–[28], [44]–[48],

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