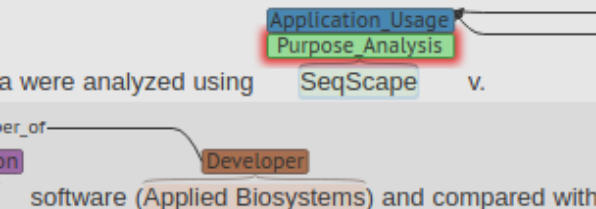


mtDNA sequencing we have chosen the mtDNA lineages which are specific for populations of northern Asia but which are still underrepresented in the published data sets on complete mtDNA variations which are rarely found in populations of northern Asia. For haplogroup-diagnostic RFLP markers and subject R11 (n=1) were selected (Table S4).



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

nt (PC) analysis was performed using mtDNA haplog

tidimensional scaling (MDS) analysis based on FST and other Asian populations around.

mtDNA diversity in western, eastern, central and north

of the complete mtDNA phylogenies of haplogroups E and H [18], were taken into account.

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

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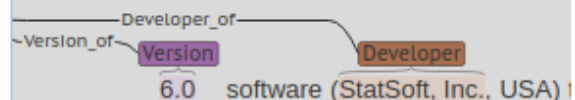
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