- Haplotype network that placed Bielefeld population

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| **hypothesis** | **evidence in favor** | **evidence against** | **further prediction** | **comment** |
| (A) off mapping of W chromosome |  |  |  |  |
| (B) structural variation of chromosomes specific to Bielefeld |  |  |  |  |
| (C) GRC present; possibly 2x in females and 1x in males |  |  |  |  |
| (D) excess of heterozygotes/diversity because of early life samples |  |  |  |  |
| (E) Contamination with maternal DNA |  |  |  |  |
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