

Figure 1. Habitancy map showing limited geographic extent of non-sanguinity, (absence of N trait). The darkest areas correspond to habitancies where the majority of the population is non-sanguine.

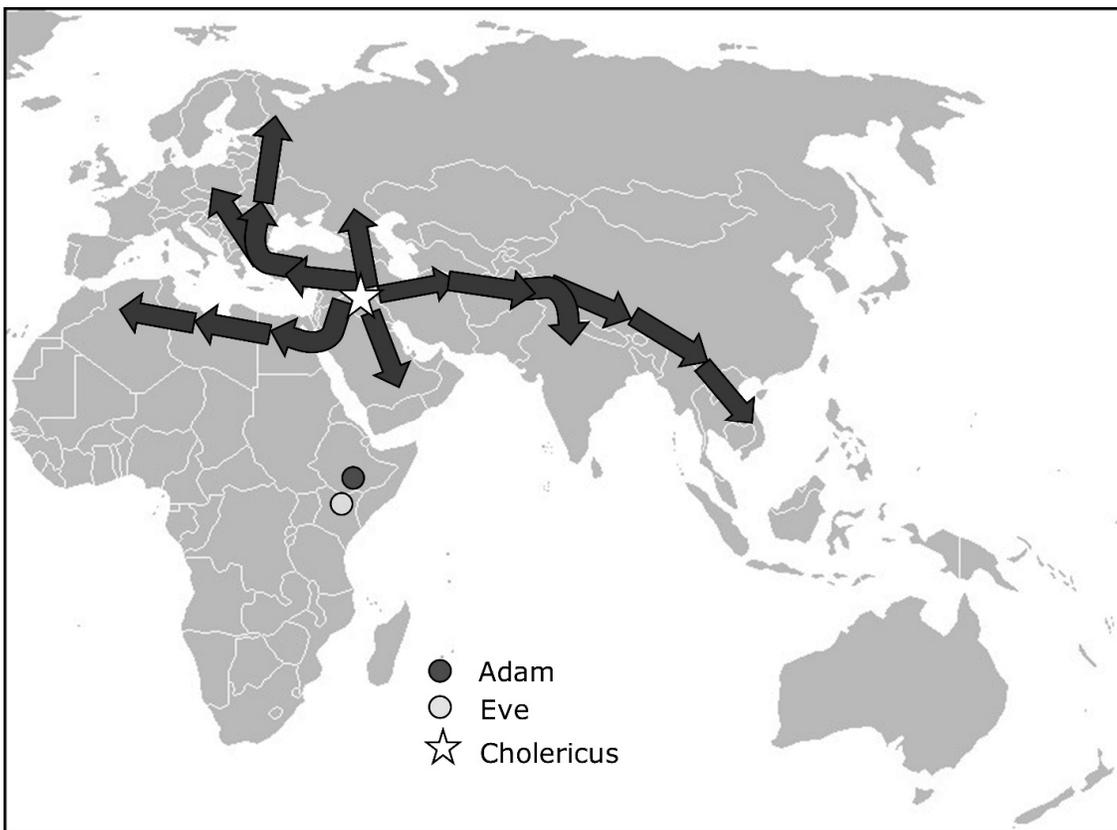


Figure 2. Possible non-sanguine trails emanating from a single founding individual, Cholericus, consistent with the "Out of Africa" ancestral migrations (Fig.3, below). The arrows show possible routes of dispersal of non-sanguinity consistent with 1) the limited present-day prevalence of non-sanguinity on our Habitancy map (above), and 2) the dispersal's having occurred relatively late during the epoch of ancestral migration. Non-sanguinity, or absence of the N trait, is a Mendelian dominant trait in the NPA model. The locations of ancestral Y-chromosomal "Adam" and m-DNA "Eve" are also shown.

