



Supplemental Fig S3. Analysis of allele frequencies identified an excess of shared alleles between and private alleles within VNBI and VNBII, and an excess of rare alleles in VNI and VNBI. A) Two-way population comparisons of shared and fixed alleles between populations, and private alleles within each population of VNI, VNBI, and VNBII; B) Minor allele frequency in each of the VNI, VNBI and VNBII. VNI and VNBI had significantly more alleles than VNBII ($p < 2.2 \times 10^{-16}$ for both comparisons, Fisher's exact test).