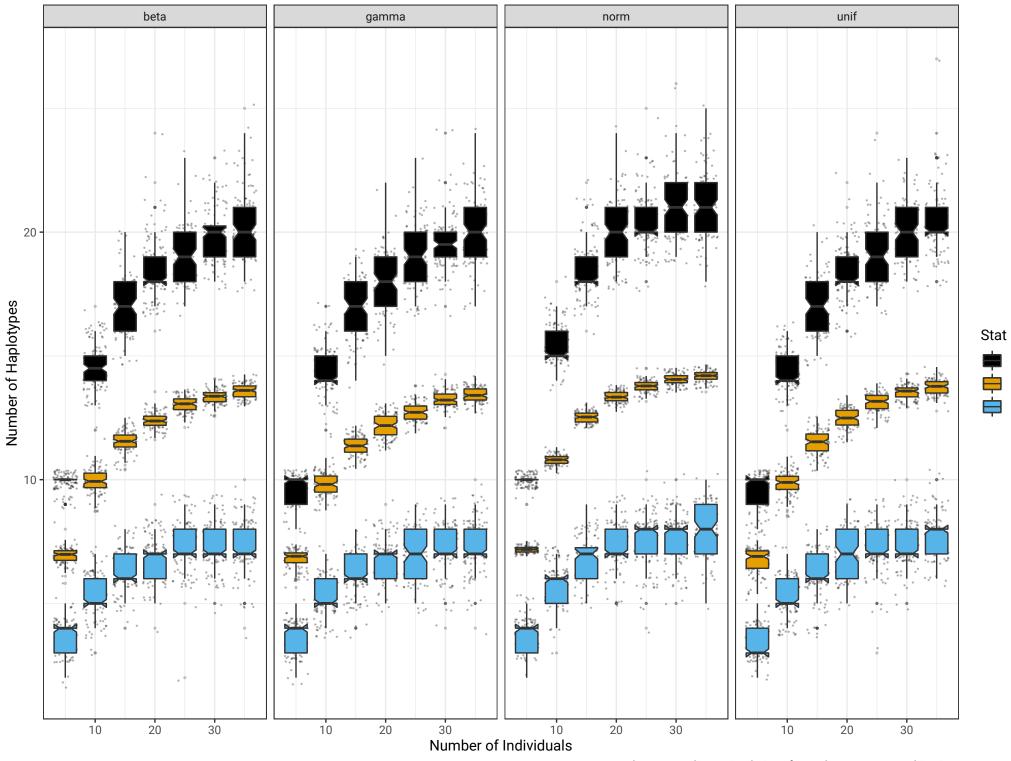
By Loci: Simulations of Haplotypes at 100x Coverage



Based on 100 replicate simulations for each parameter combination, performed in the program ms, using base parameters: **9=10**, **loci-length=10kb**, **100 loci**

max mean min