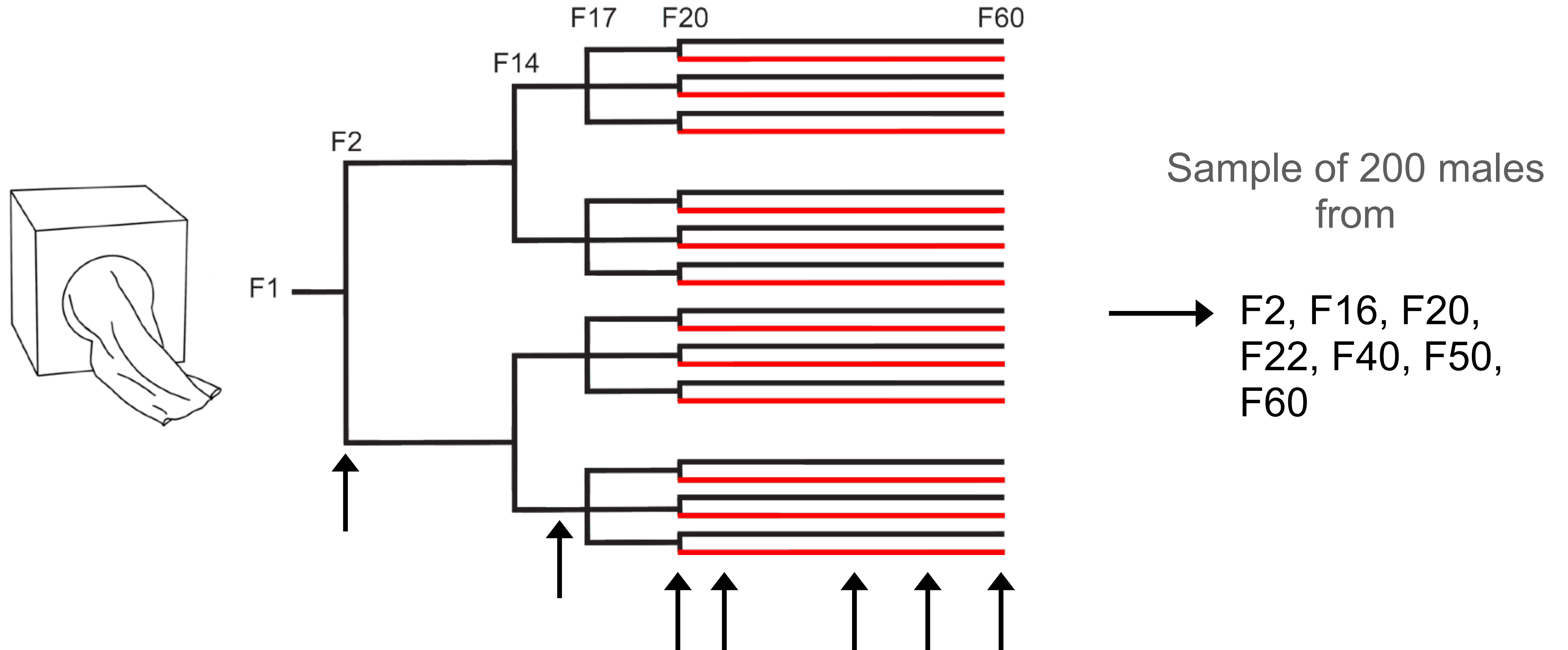


Pool-sequencing was used to quantify population allele and haplotype frequencies



Genome wide variation is shaped by **rotenone** treatment

