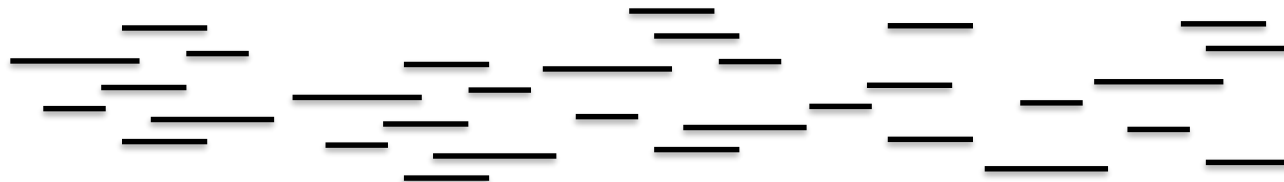




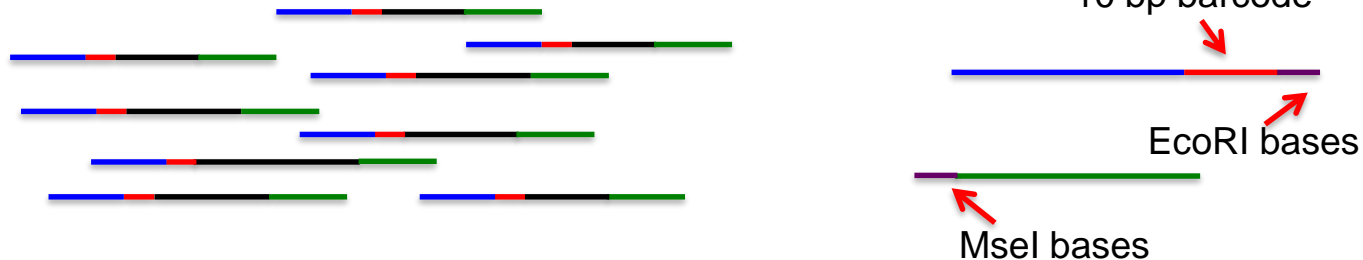
Genomic reduction and highly multiplexed library preparation

Parchman *et al.* 2012

1. Digest genomic DNA with EcoRI and MseI



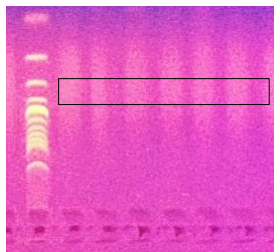
2. Ligate Illumina adaptors to fragments



3. Amplify with Illumina PCR primers

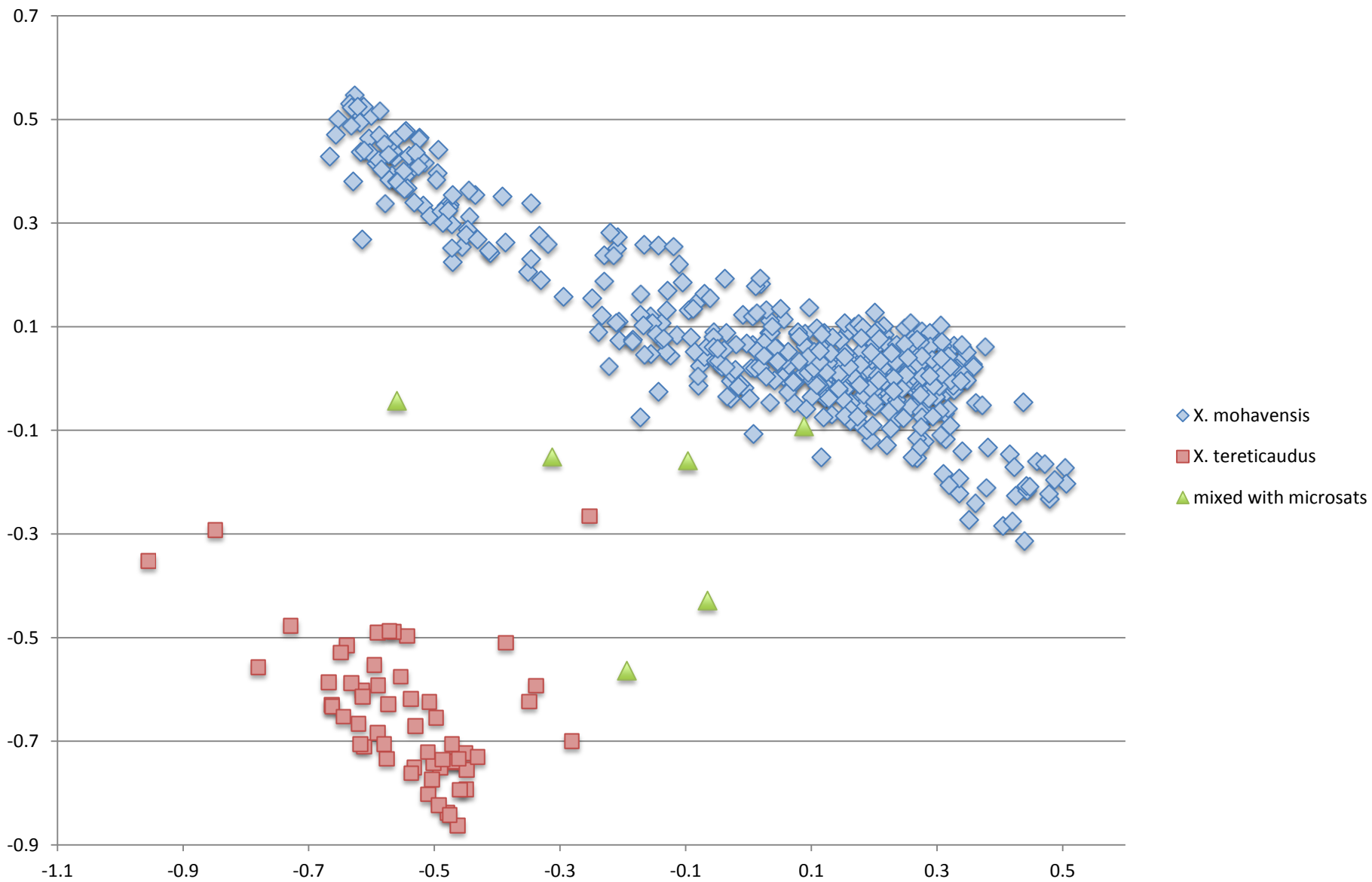


4. Gel purify product to desired size range (350-450bp)



Illumina sequencing

25,550 SNPs in 608 individuals



25,550 SNPs in 608 individuals

