

1 Linkage mapping of *Barley yellow dwarf virus* resistance
2 in connected populations of maize

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ADDITIONAL FILES

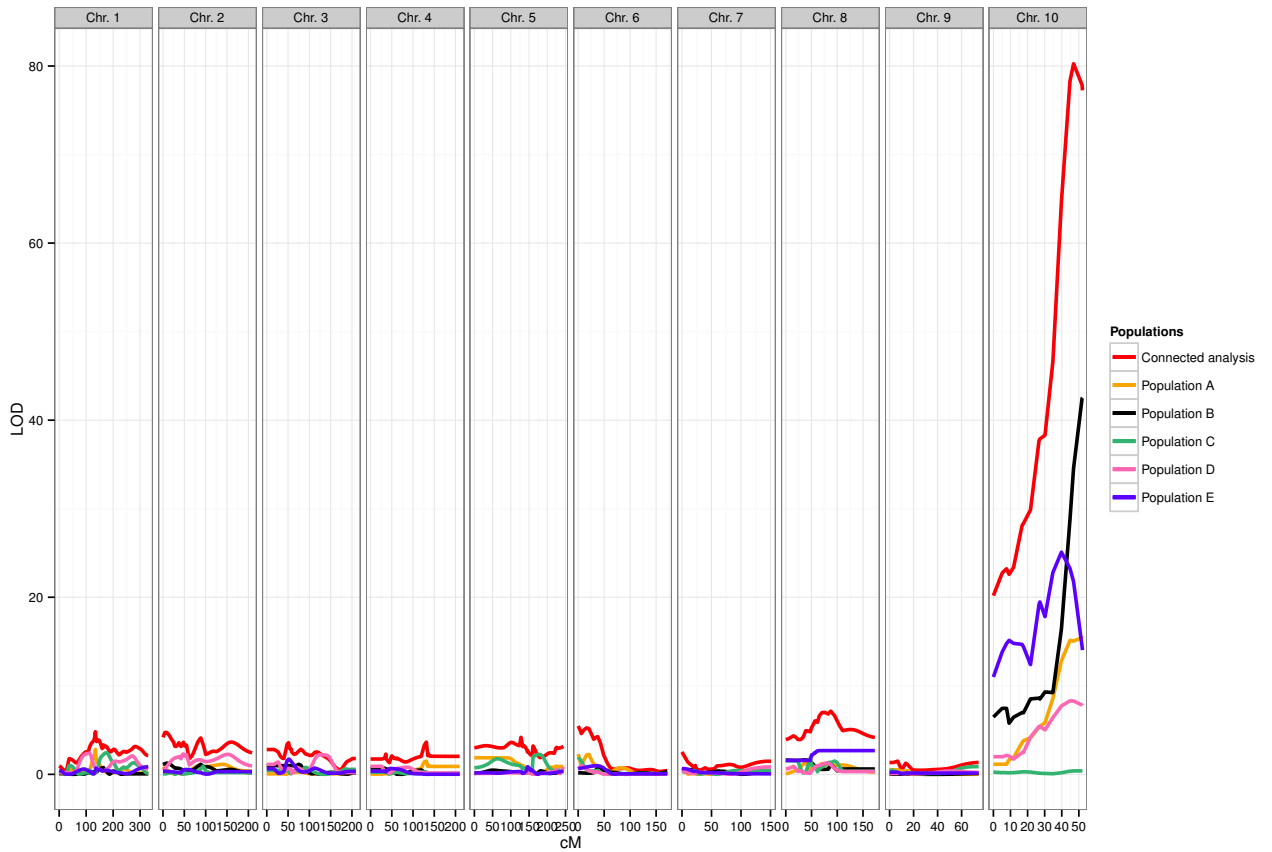


Figure S1 QTL for the trait infection rate (IR) across the genome in the inoculated field experiment.

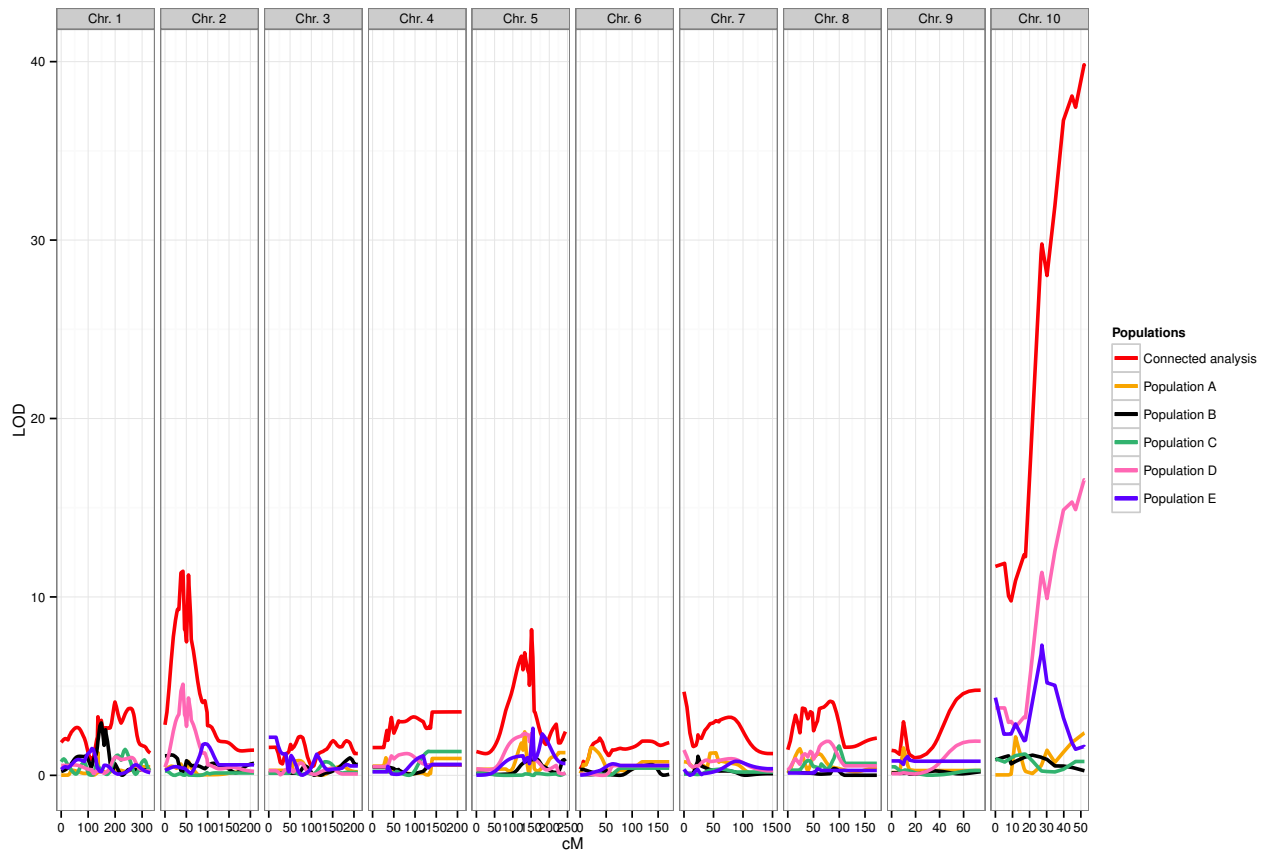


Figure S2 QTL for the trait red edges (RE) across the genome in the inoculated field experiment.

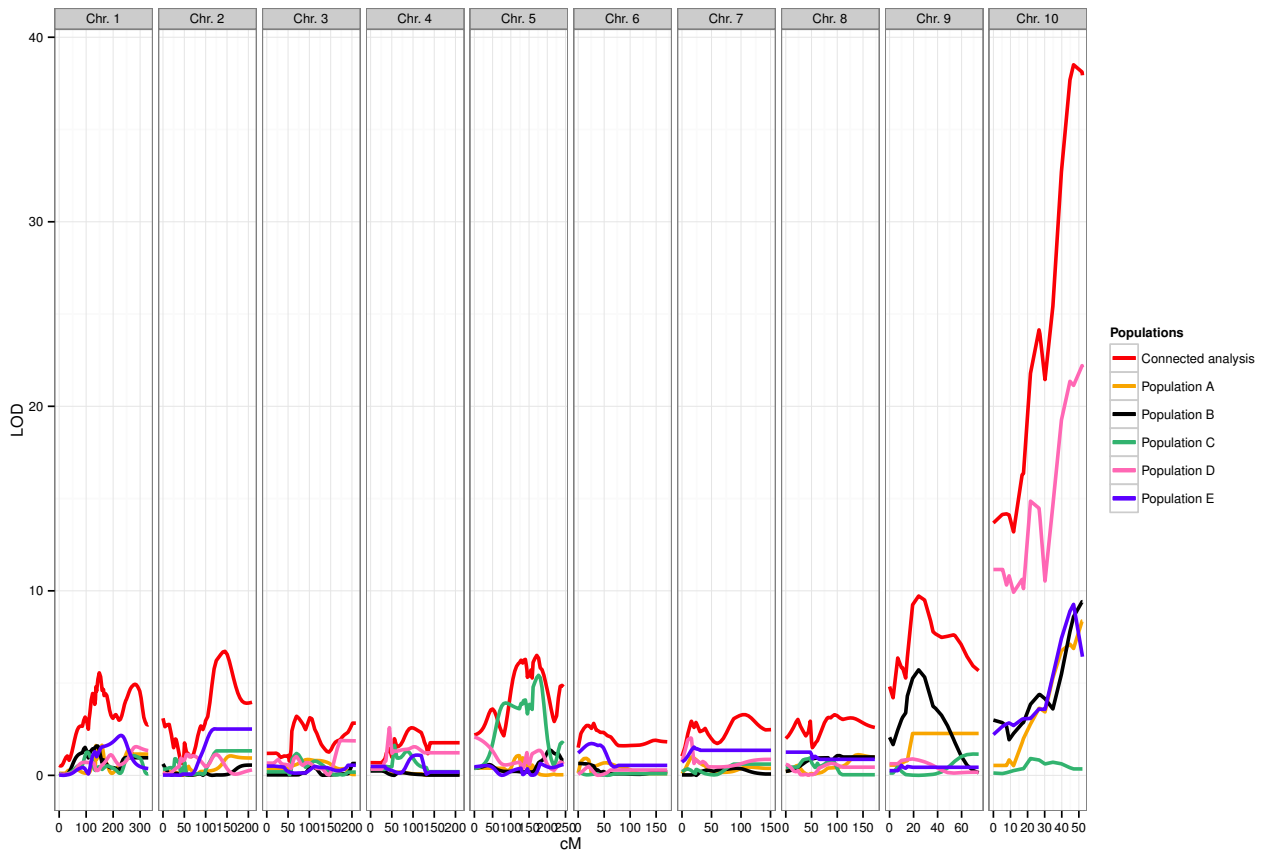


Figure S3 QTL for the trait infection rate (IR) across the genome in the greenhouse experiment.