

## **Descriptions of Additional Supplementary Files**

**Supplementary Data 1.** Sample collection metadata and inclusion in RADseq, COI sequencing, and 16S sequencing analyses (Y=sample included, N=sample not included).

**Supplementary Data 2.** Genomic positions and  $p$  values for SNPs identified as  $F_{ST}$  outliers for comparisons between agricultural plots in Hermiston, Oregon.

**Supplementary Data 3.** Maximum genetic distance along scaffolds (“Max distance”) for  $F_{ST}$  outlier SNPs identified from comparisons between agricultural plots in Hermiston, Oregon. Boxes surround SNPs occurring on the same scaffold. This table only includes outlier SNPs occurring on scaffolds that had more than one outlier SNP. SNPs less than ~800bp apart are associated with the same restriction cut site.