

Supplemental Figure S5: Whole-genome alignment of Trayshed (Query, y-axis) on Cabernet Sauvignon (Reference, x-axis); **A)** Hap2 vs Hap1, **B)** Hap1 vs Hap2, and **C)** Hap2 vs Hap2. The percent identity (%id) between the alignments is displayed as a color gradient.

