

All blast hits with score > 50, 8/8-gene x/y windows (heatmap range: 2-5+ hits/window)

Ler_3: gene rank order position (22275 genes w/ blast hits), grids every 1000 genes

contig_538_pilon

scaffold_448_pilon

contig_446_pilon

contig_444_pilon

scaffold_322_pilon

contig_297_pilon

scaffold_290_pilon

contig_277_pilon

scaffold_250_pilon

contig_139_pilon

contig_2_pilon

contig_3_pilon

contig_4_pilon

contig_10_pilon

contig_23_pilon

contig_349_pilon

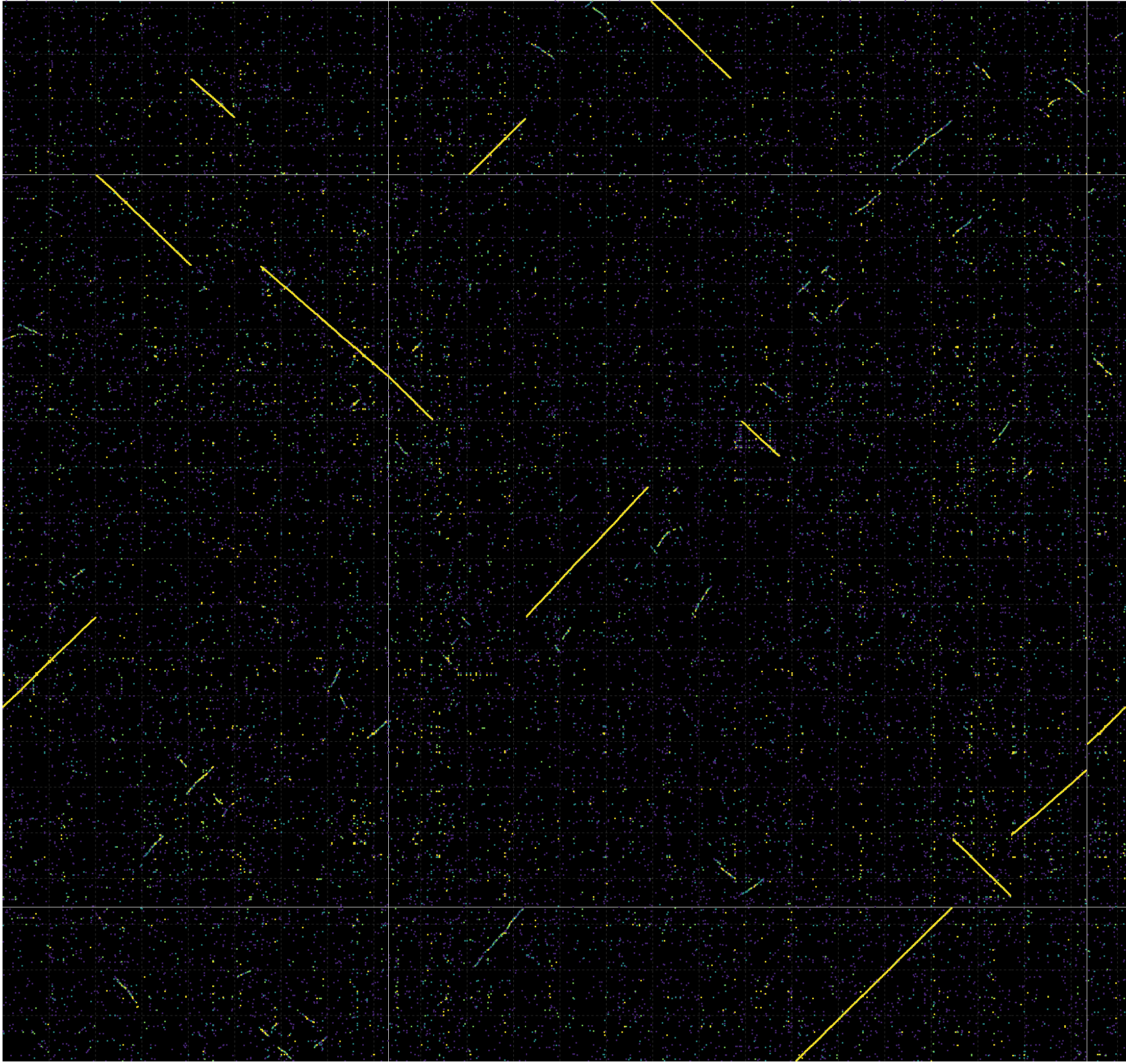
scaffold_356_pilon

scaffold_497_pilon

contig_500_pilon

contig_629_pilon

C24: gene rank order position (23139 genes w/ blast hits), grids every 1000 genes



Blast hits where query and target are in the same orthogroup, 8/8-gene x/y windows (heatmap range: 1-5+ hits/window)

Ler_3: gene rank order position (20453 genes w/ blast hits), grids every 1000 genes

contig_538_pilon

scaffold_448_pilon

contig_446_pilon

contig_444_pilon

scaffold_322_pilon

contig_297_pilon

contig_277_pilon

scaffold_250_pilon

contig_139_pilon

contig_2_pilon

contig_3_pilon

contig_4_pilon

contig_10_pilon

contig_23_pilon

contig_349_pilon

scaffold_356_pilon

scaffold_497_pilon

contig_500_pilon

contig_629_pilon

C24: gene rank order position (20659 genes w/ blast hits), grids every 1000 genes

