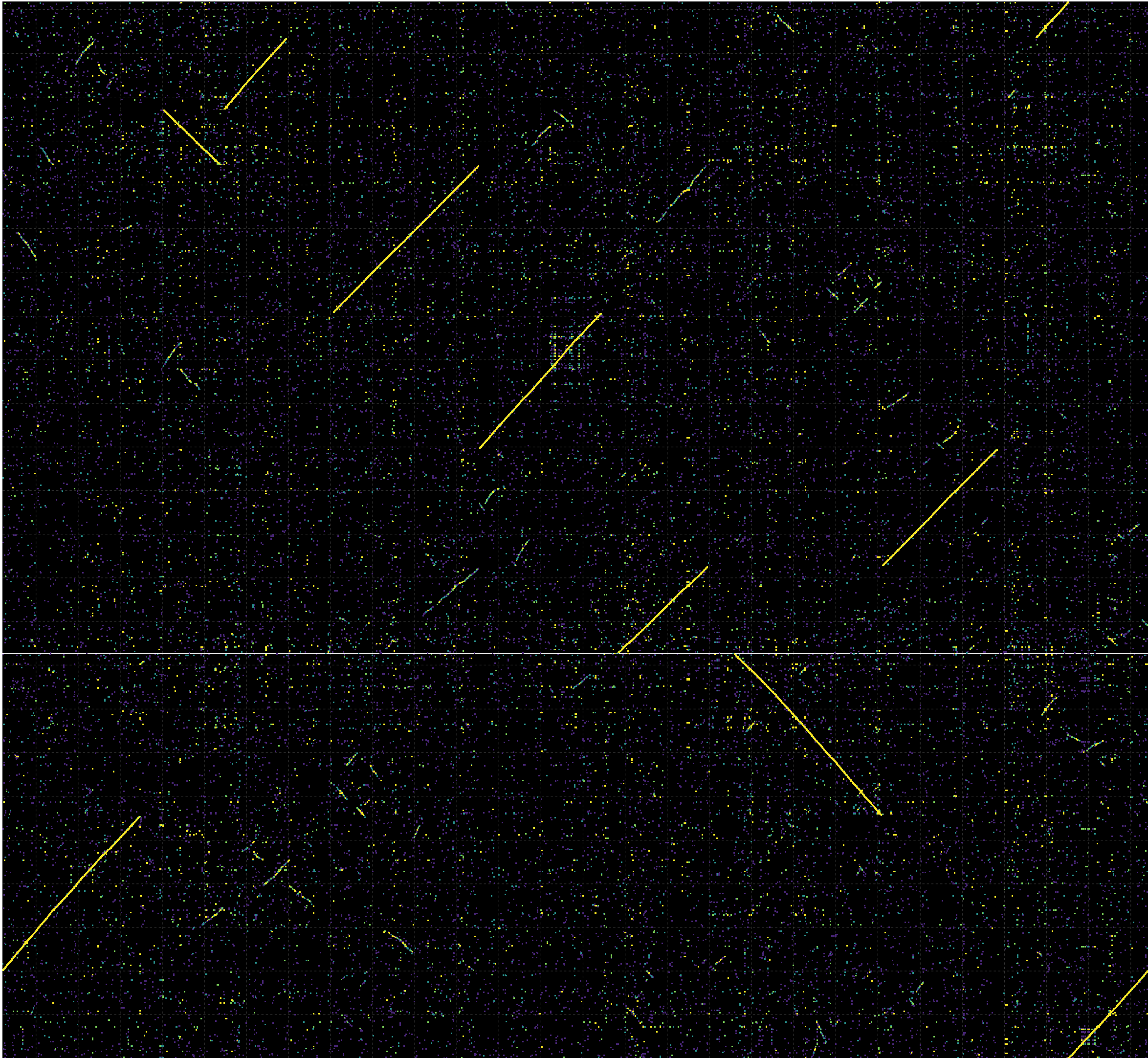


All blast hits with score > 50, 9/8-gene x/y windows (heatmap range: 2-5+ hits/window)

contig_629_pilon
contig_500_pilon
scaffold_497_pilon
scaffold_356_pilon
contig_349_pilon
contig_23_pilon
contig_10_pilon
contig_4_pilon
contig_3_pilon
contig_2_pilon



TAIR10: gene rank order position (25862 genes w/ blast hits), grids every 1000 genes

Blast hits where query and target are in the same orthogroup, 9/8-gene x/y windows (heatmap range: 1-5+ hits/window)

C24: gene rank order position (21567 genes w/ blast hits), grids every 1000 genes

contig_629_pilon
contig_500_pilon
scaffold_497_pilon
scaffold_356_pilon
contig_349_pilon
contig_23_pilon
contig_10_pilon
contig_4_pilon
contig_3_pilon
contig_2_pilon

Chr1

Chr2

Chr3

Chr4

Chr5

TAIR10: gene rank order position (22912 genes w/ blast hits), grids every 1000 genes

