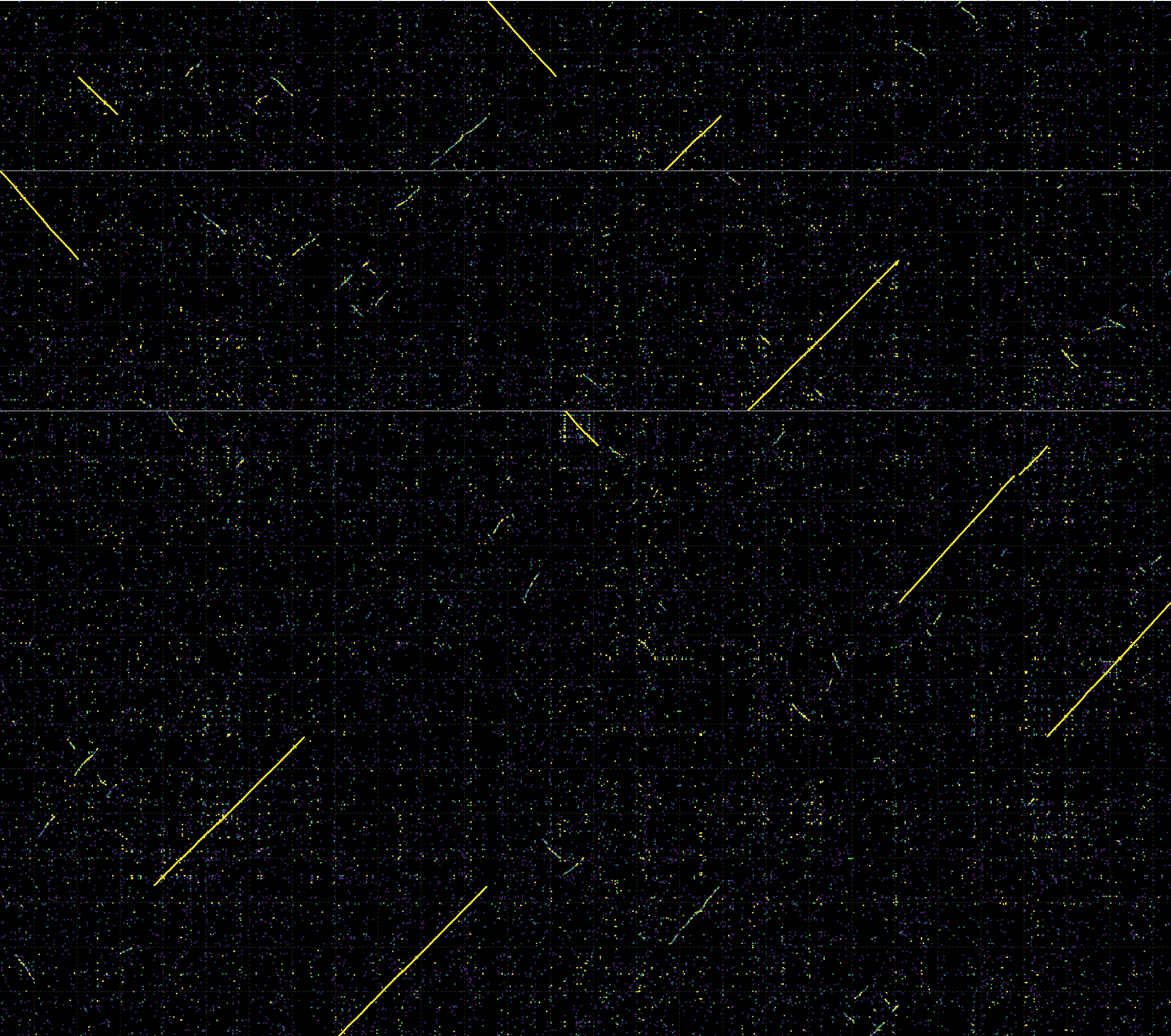


All blast hits with score > 50, 8/8-gene x/y windows (heatmap range: 2–5+ hits/window)

Ler_3: gene rank order position (22645 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

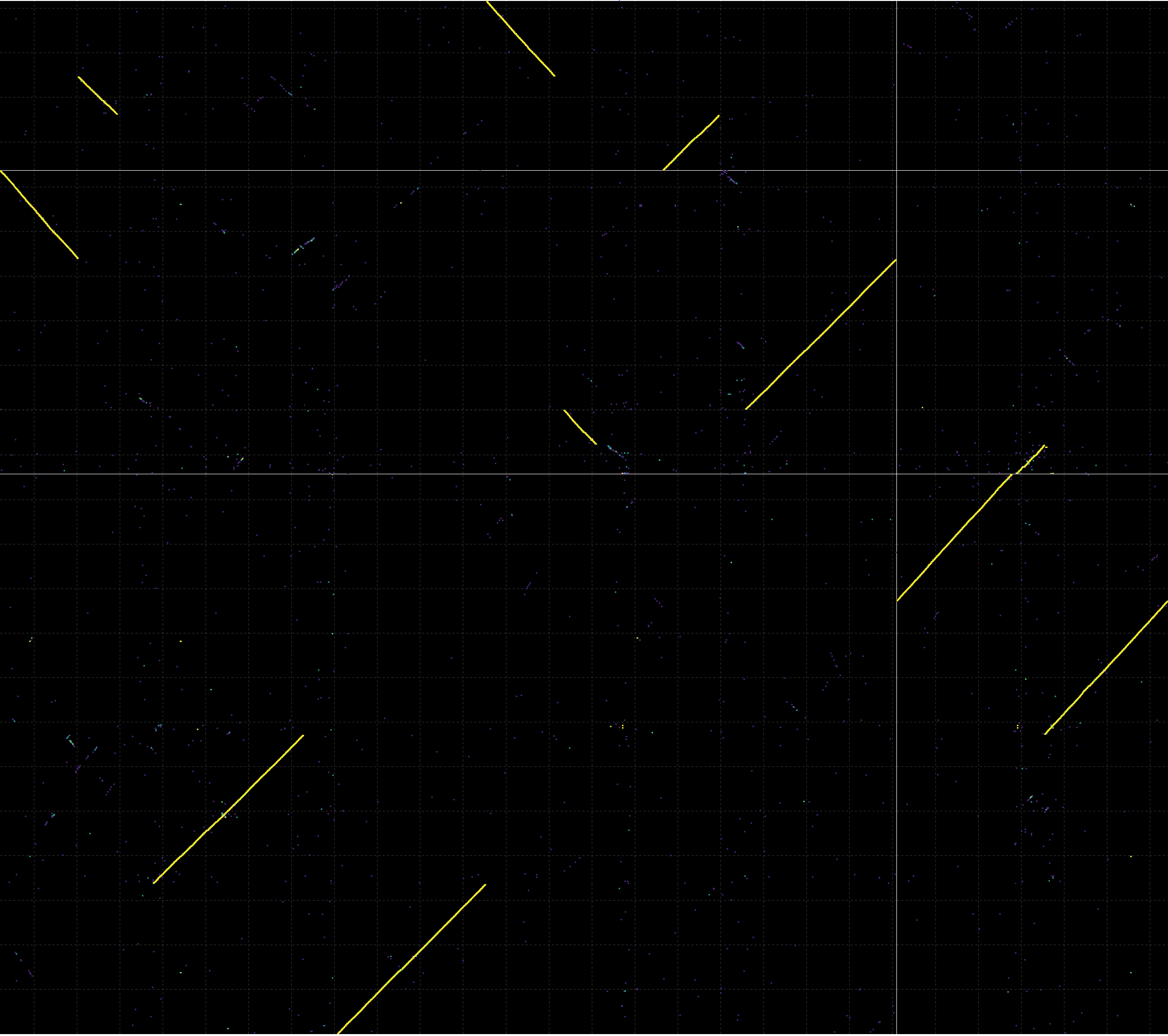


TAIR10: gene rank order position (25787 genes w/ blast hits), grids every 1000 genes

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 (21219 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



Chr1

Chr2

Chr3

Chr4

Chr5

TAIR10: gene rank order position (22045 genes w/ blast hits), grids every 1000 genes