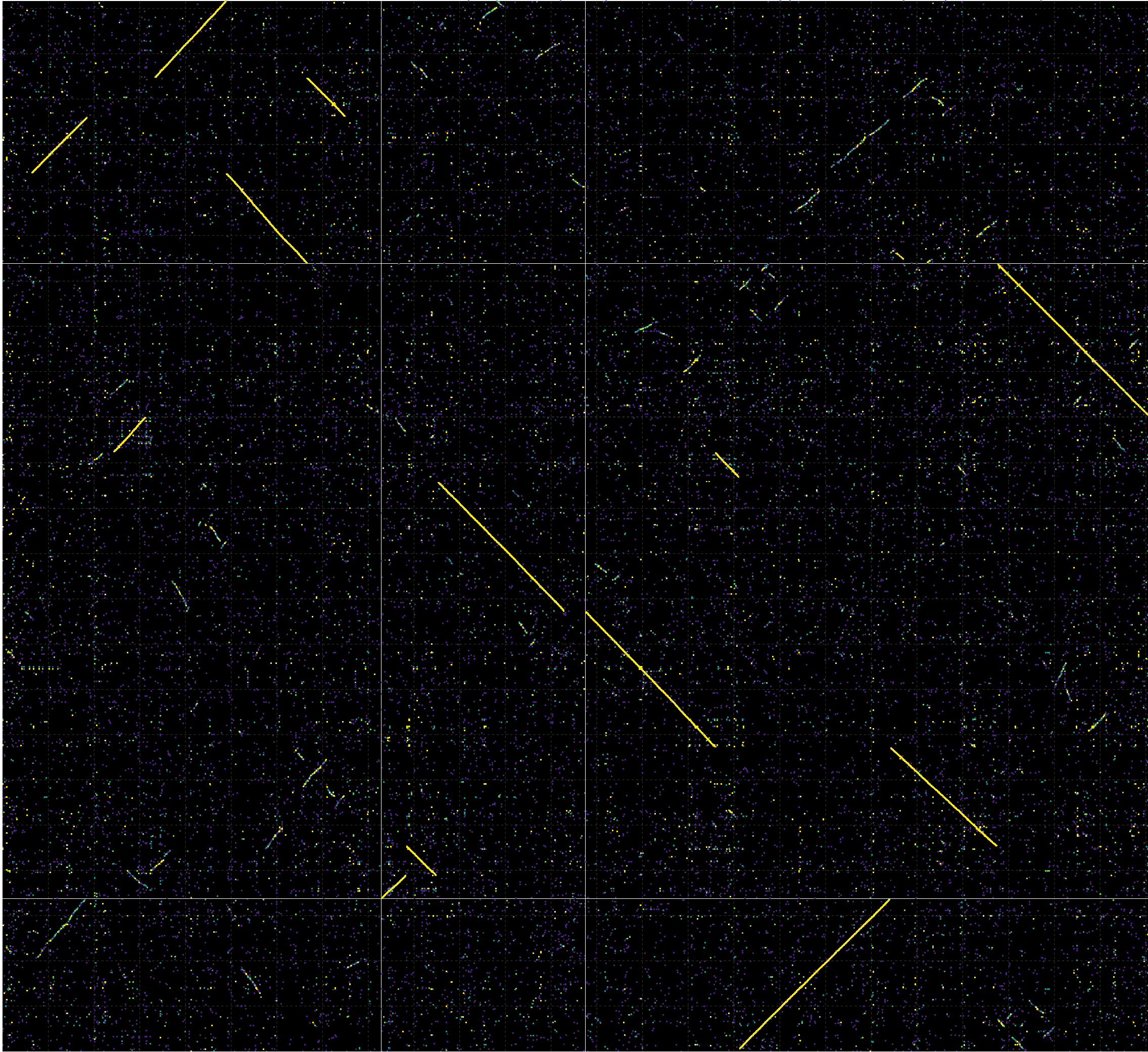


All blast hits with score > 50, 8/8-gene x/y windows (heatmap range: 2-5+ hits/window)

Ler_3: gene rank order position (22534 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_3_pilon

scaffold_88_pilon

contig_101_pilon

contig_102_pilon

contig_118_pilon

contig_449_pilon

contig_509_pilon

contig_589_pilon

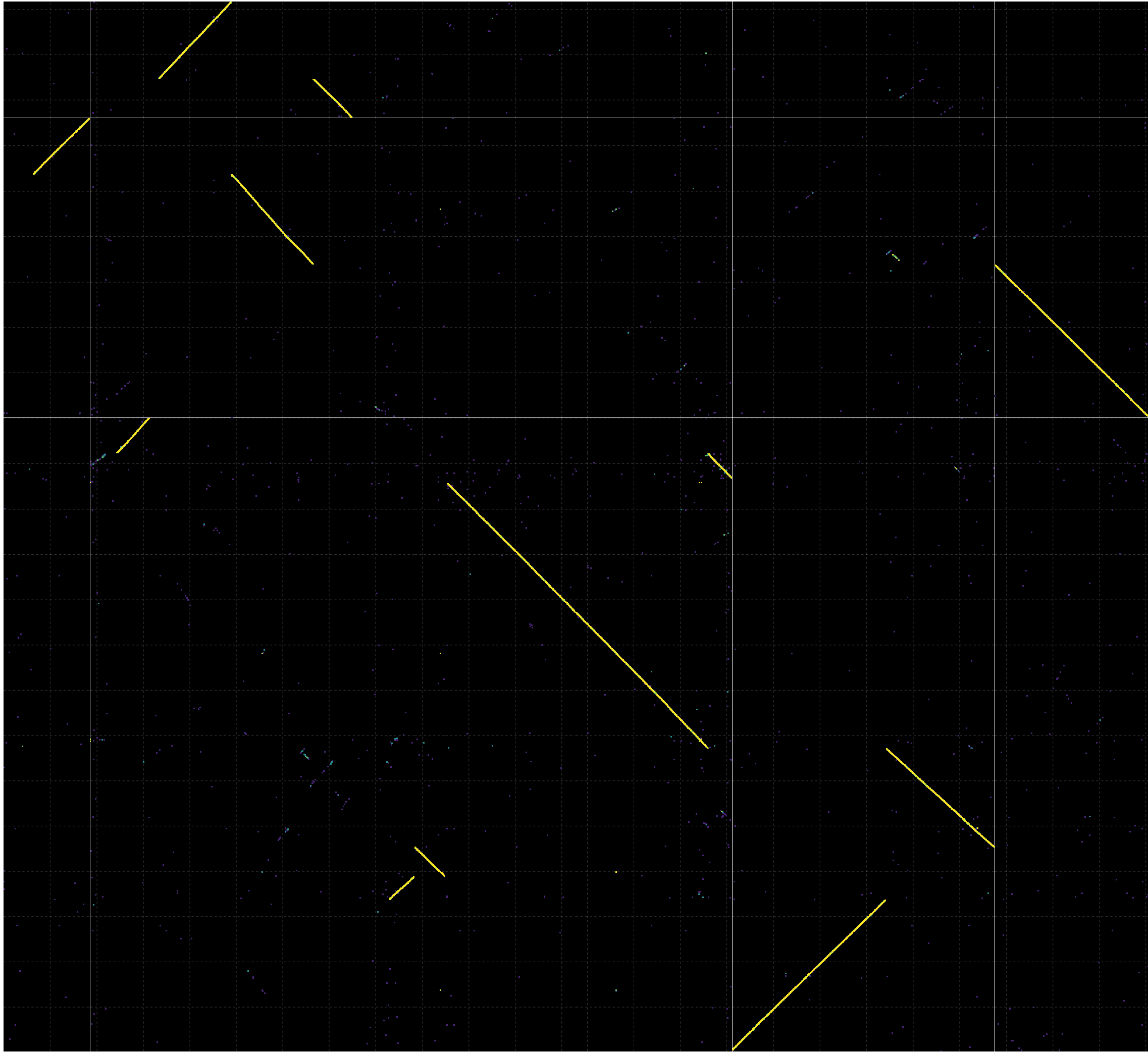
contig_591_pilon

scaffold_679_pilon

An_1: gene rank order position (24008 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 (20918 genes w/ blast hits), grids every 1000 genes



An_1: gene rank order position (21025 genes w/ blast hits), grids every 1000 genes