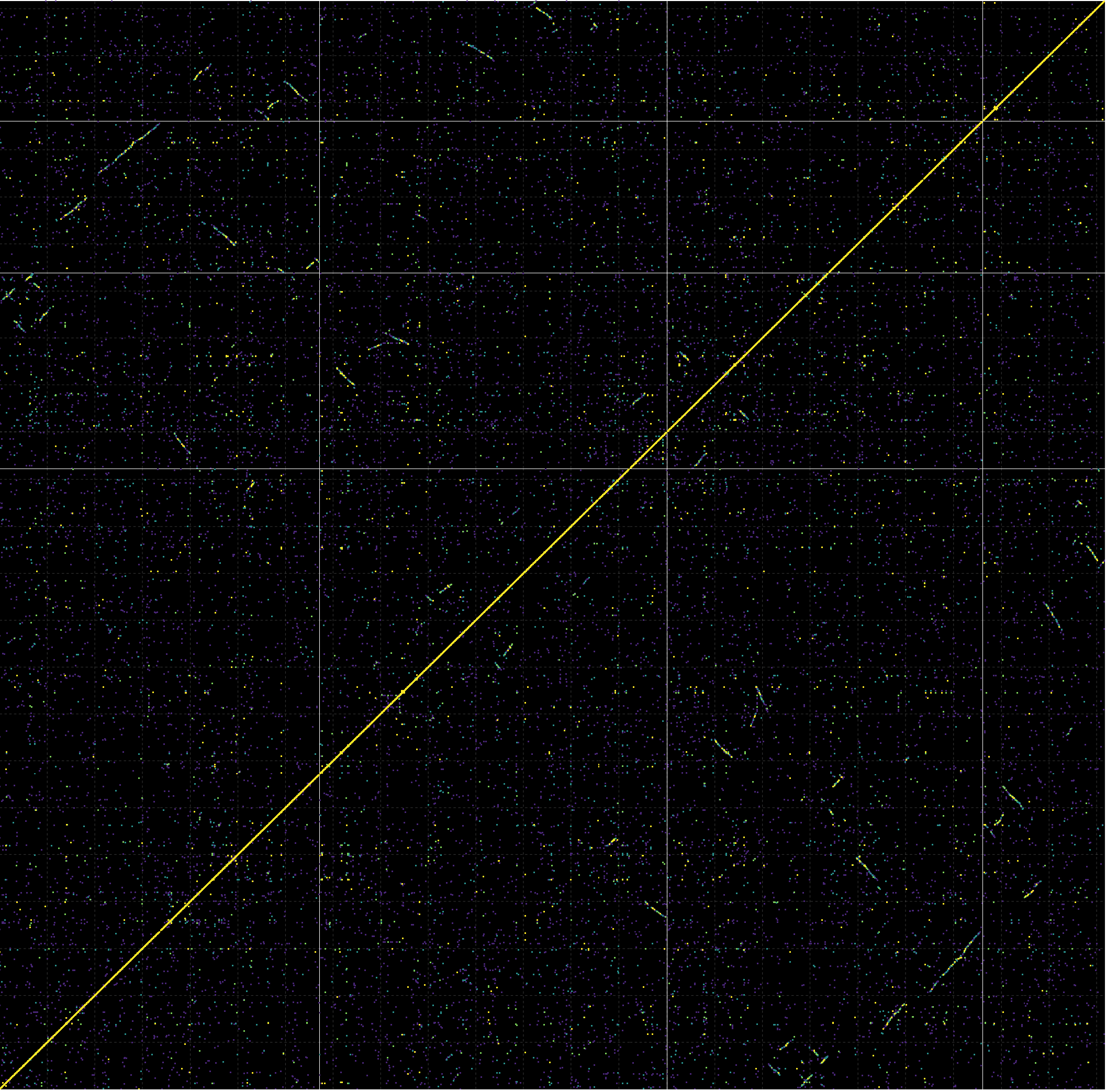


All blast hits with score > 50, 8/8-gene x/y windows (heatmap range: 2-5+ hits/window)

Ler_3: gene rank order position (23107 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_139_pilon

scaffold_250_pilon

contig_277_pilon

scaffold_290_pilon

contig_297_pilon

scaffold_322_pilon

contig_444_pilon

contig_446_pilon

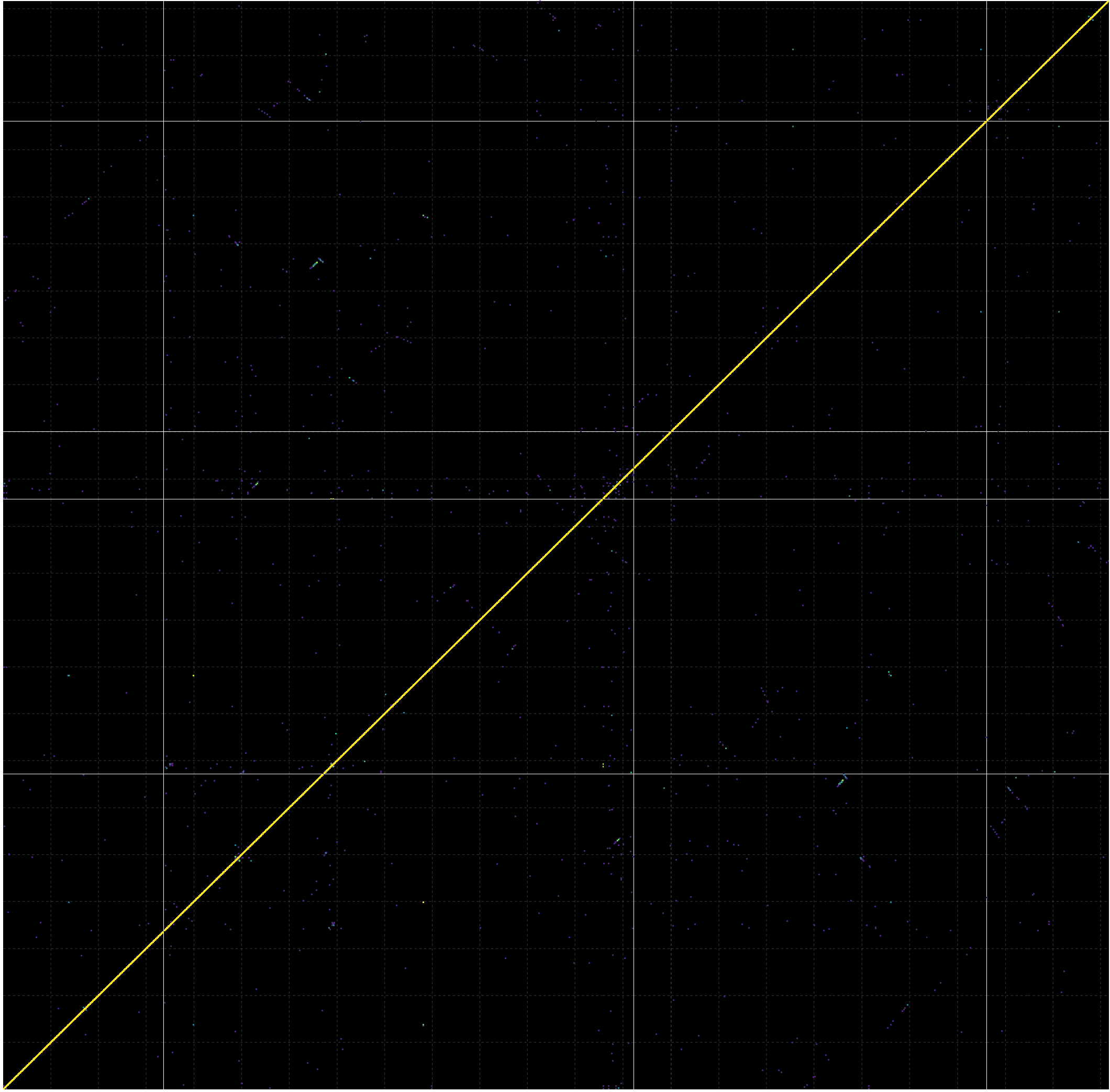
scaffold_448_pilon

contig_538_pilon

Ler_3: gene rank order position (23107 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: 1-7+ hits/window)

Ler_3: gene rank order position (23147 genes w/ blast hits), grids every 1000 genes



Ler_3: gene rank order position (23147 genes w/ blast hits), grids every 1000 genes