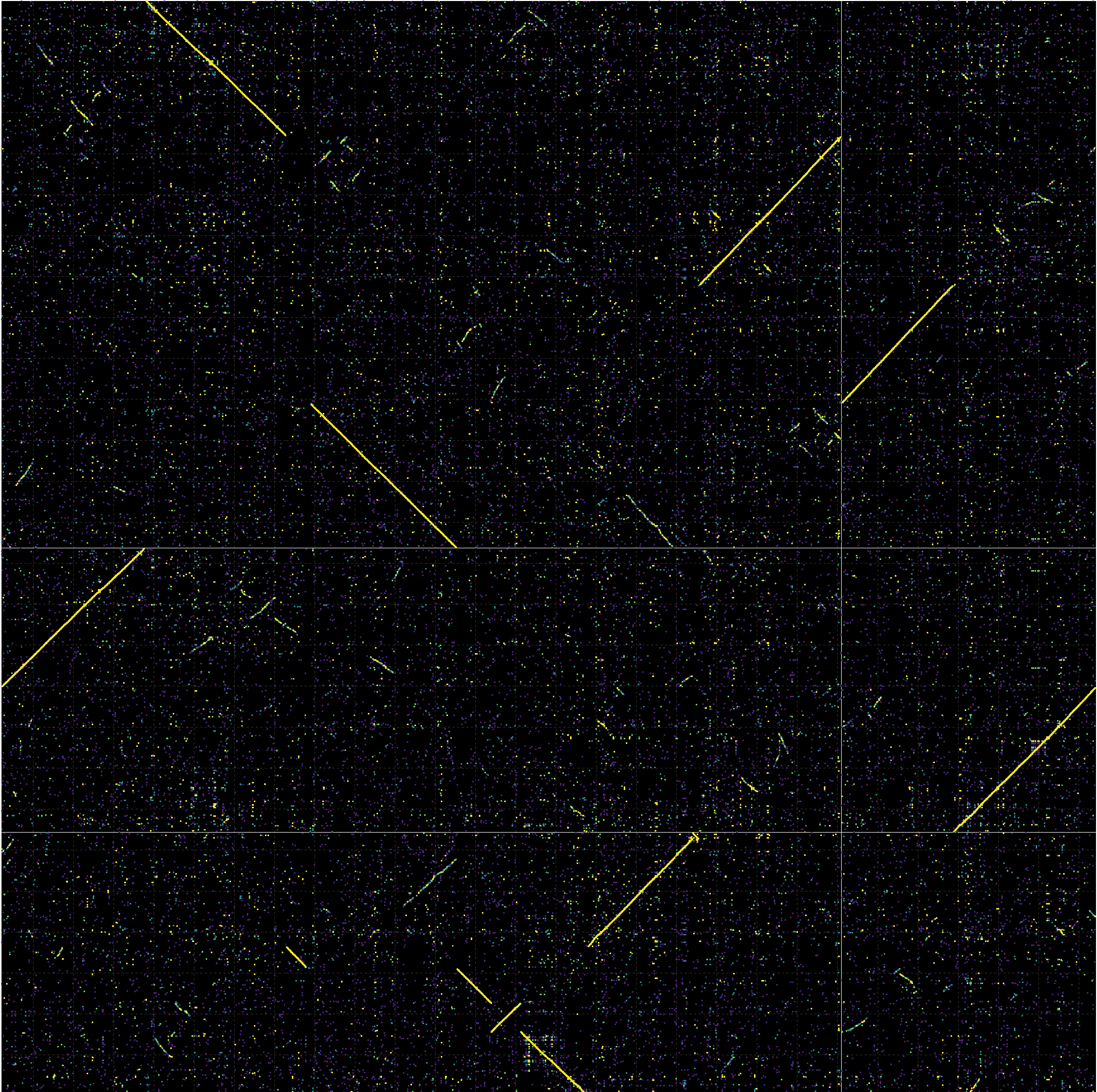


All blast hits with score > 50, 9/9-gene x/y windows (heatmap range: 2-5+ hits/window)

Sha: gene rank order position (26131 genes w/ blast hits), grids every 1000 genes

contig_71_pilon
scaffold_20_pilon
contig_19_pilon
scaffold_18_pilon
contig_17_pilon
contig_16_pilon
scaffold_15_pilon
scaffold_11_pilon
contig_10_pilon
contig_7_pilon



Chr1

Chr2

Chr3

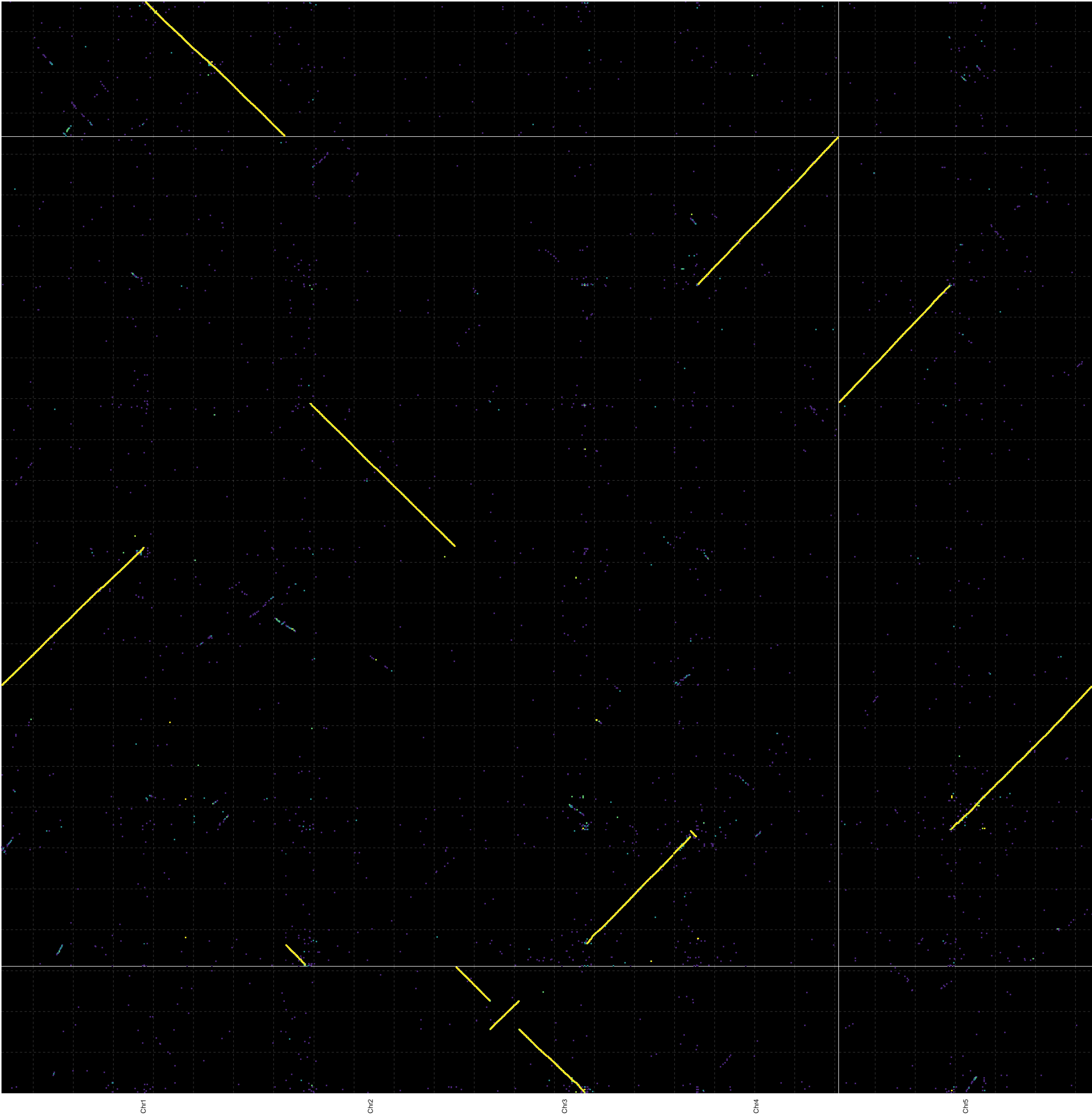
Chr4

Chr5

TAIR10: gene rank order position (26590 genes w/ blast hits), grids every 1000 genes

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

Sha: gene rank order position (24619 genes w/ blast hits), grids every 1000 genes



TAIR10: gene rank order position (25201 genes w/ blast hits), grids every 1000 genes