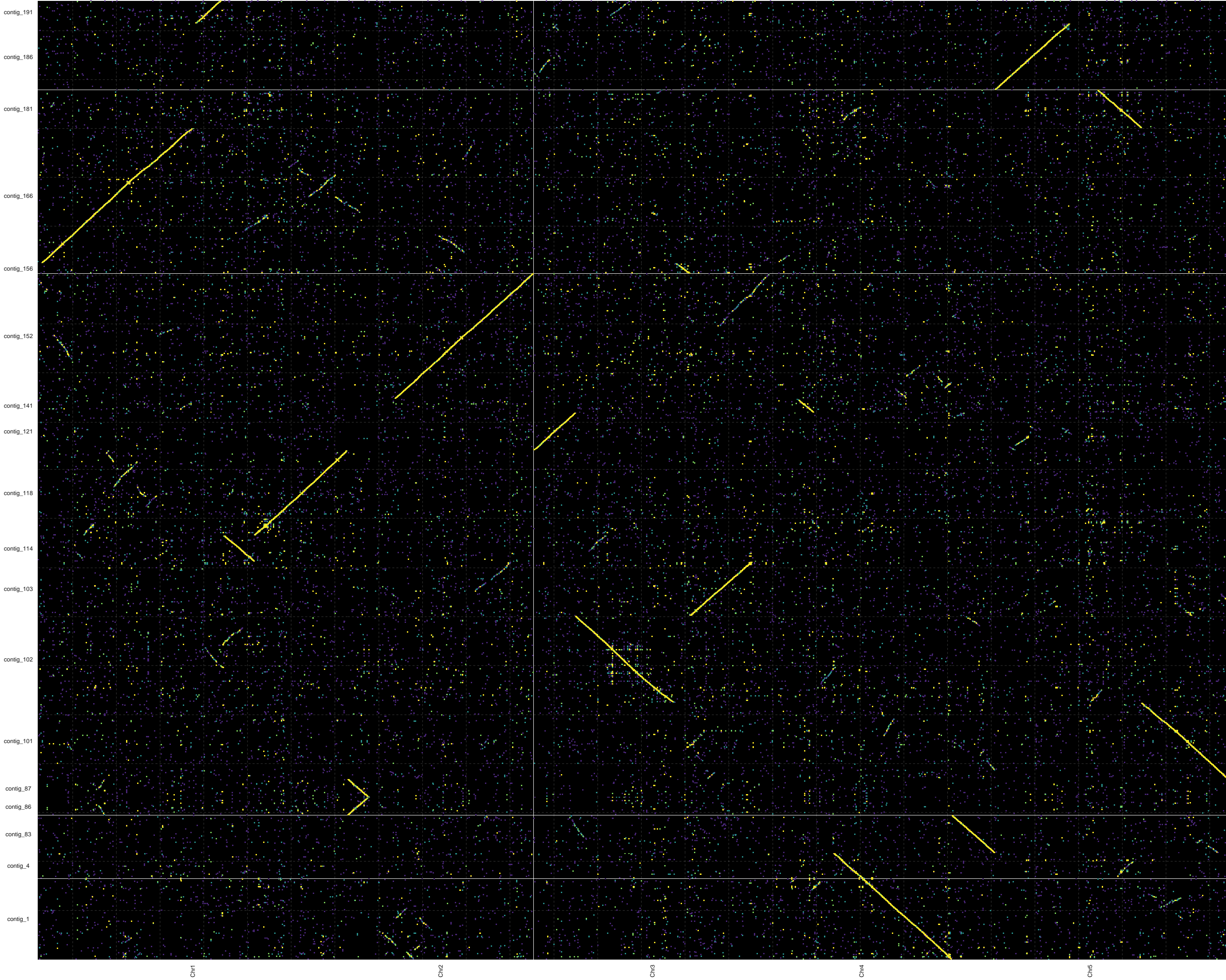


All blast hits with score > 50, 8/7-gene x/y windows (heatmap range: 2-5+ hits/window)

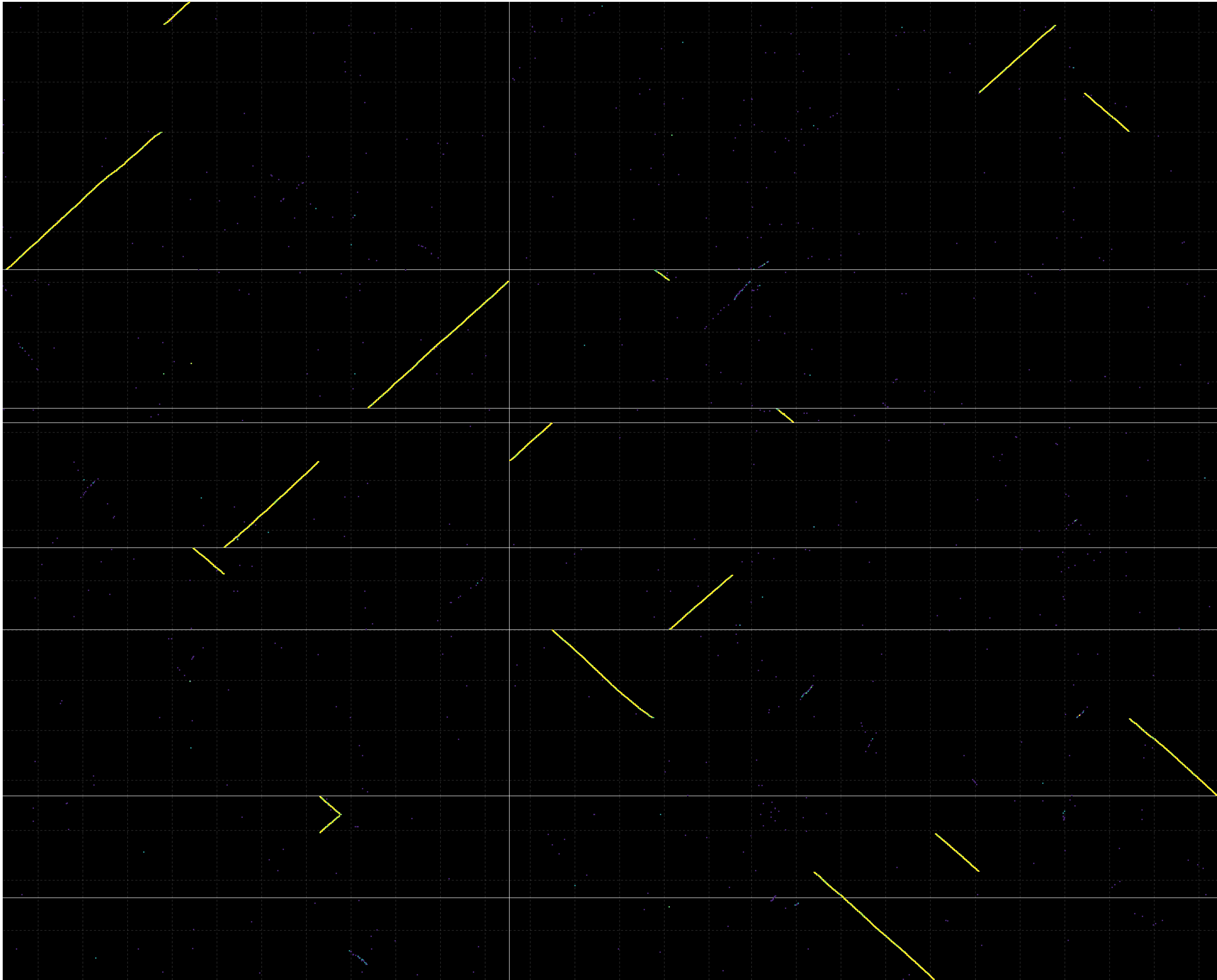
Pa_1: gene rank order position (19437 genes w/ blast hits), grids every 1000 genes



TAIR10: gene rank order position (23566 genes w/ blast hits), grids every 1000 genes

Blast hits where query and target are in the same orthogroup, 8/7-gene x/y windows (heatmap range: 1-5+ hits/window)

Pa_1: gene rank order position (18618 genes w/ blast hits), grids every 1000 genes



TAIR10: gene rank order position (19327 genes w/ blast hits), grids every 1000 genes