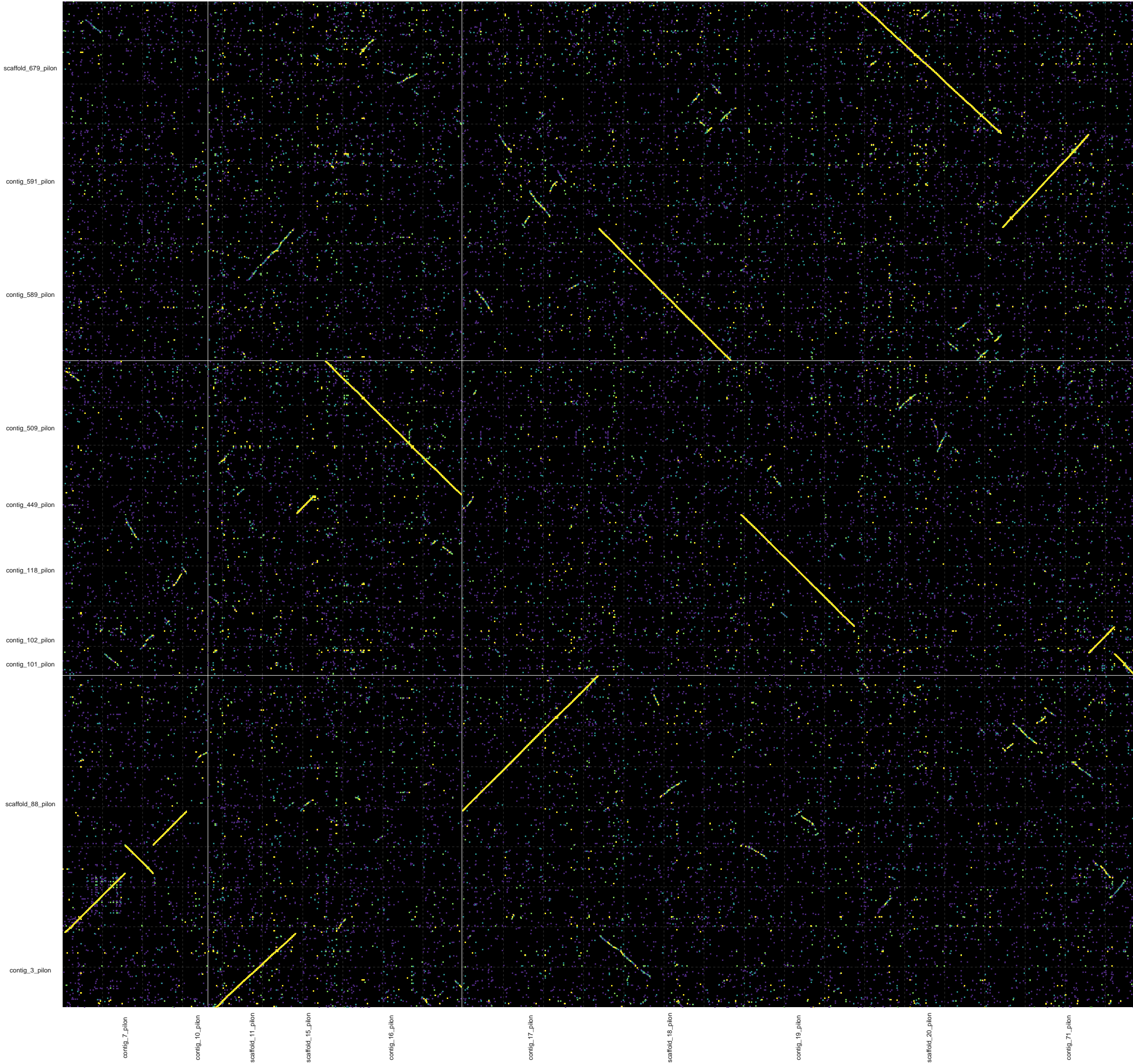


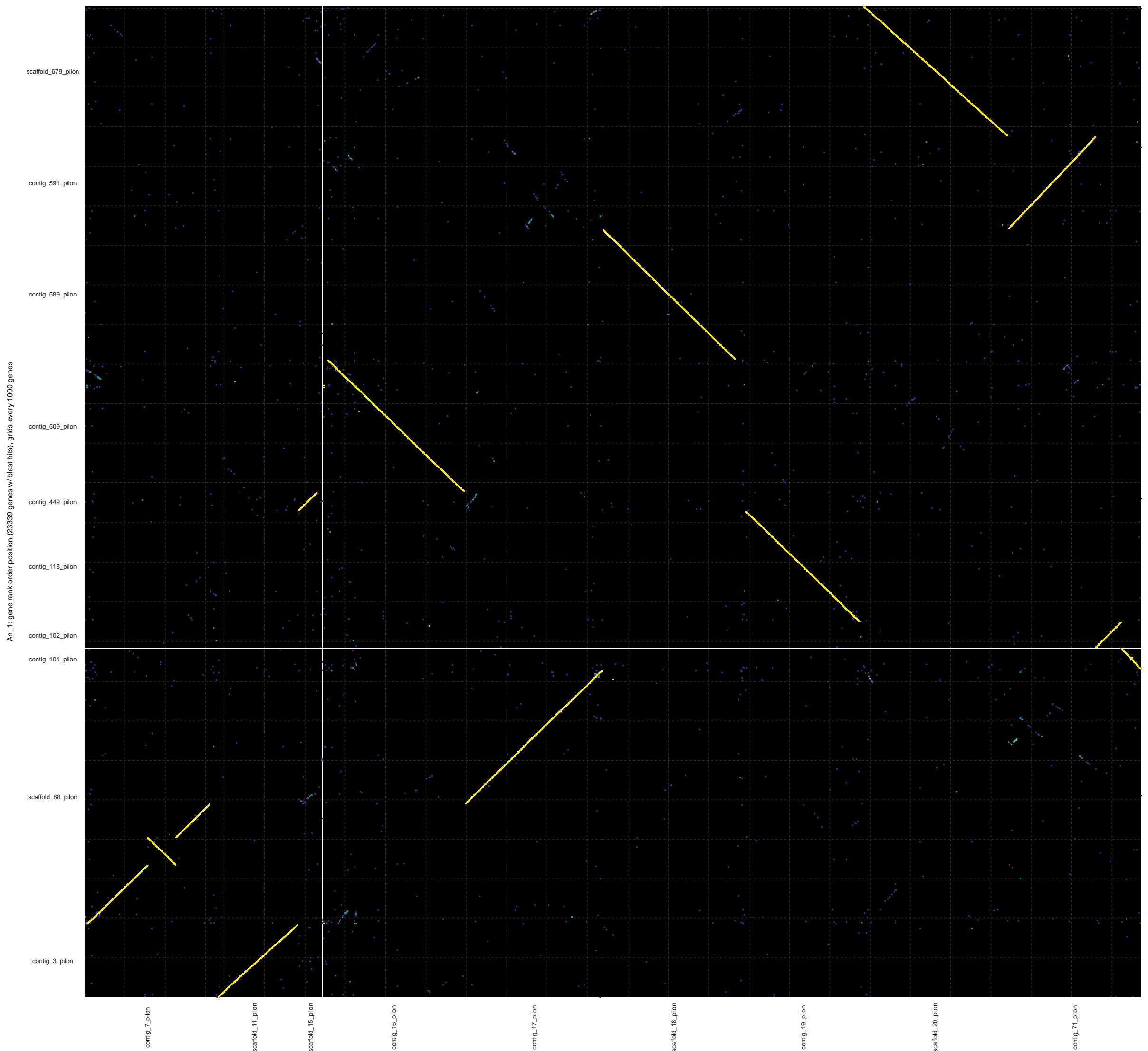
All blast hits with score > 50, 9/9-gene x/y windows (heatmap range: 2-5+ hits/window)

An_1: gene rank order position (24571 genes w/ blast hits), grids every 1000 genes



Sha: gene rank order position (25729 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 (23617 genes w/ blast hits), grids every 1000 genes