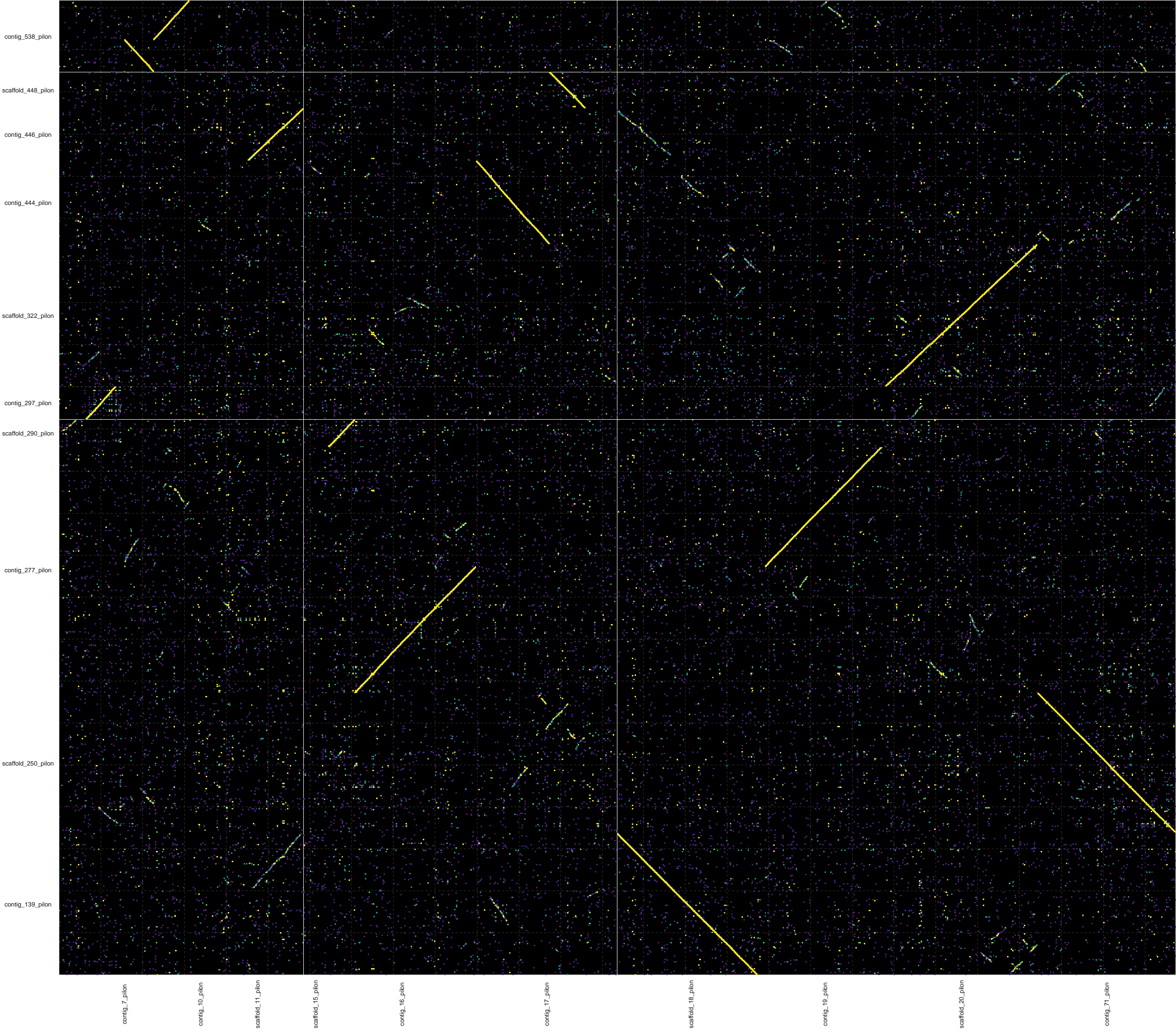


All blast hits with score > 50, 8/8-gene x/y windows (heatmap range: 2-5+ hits/window)

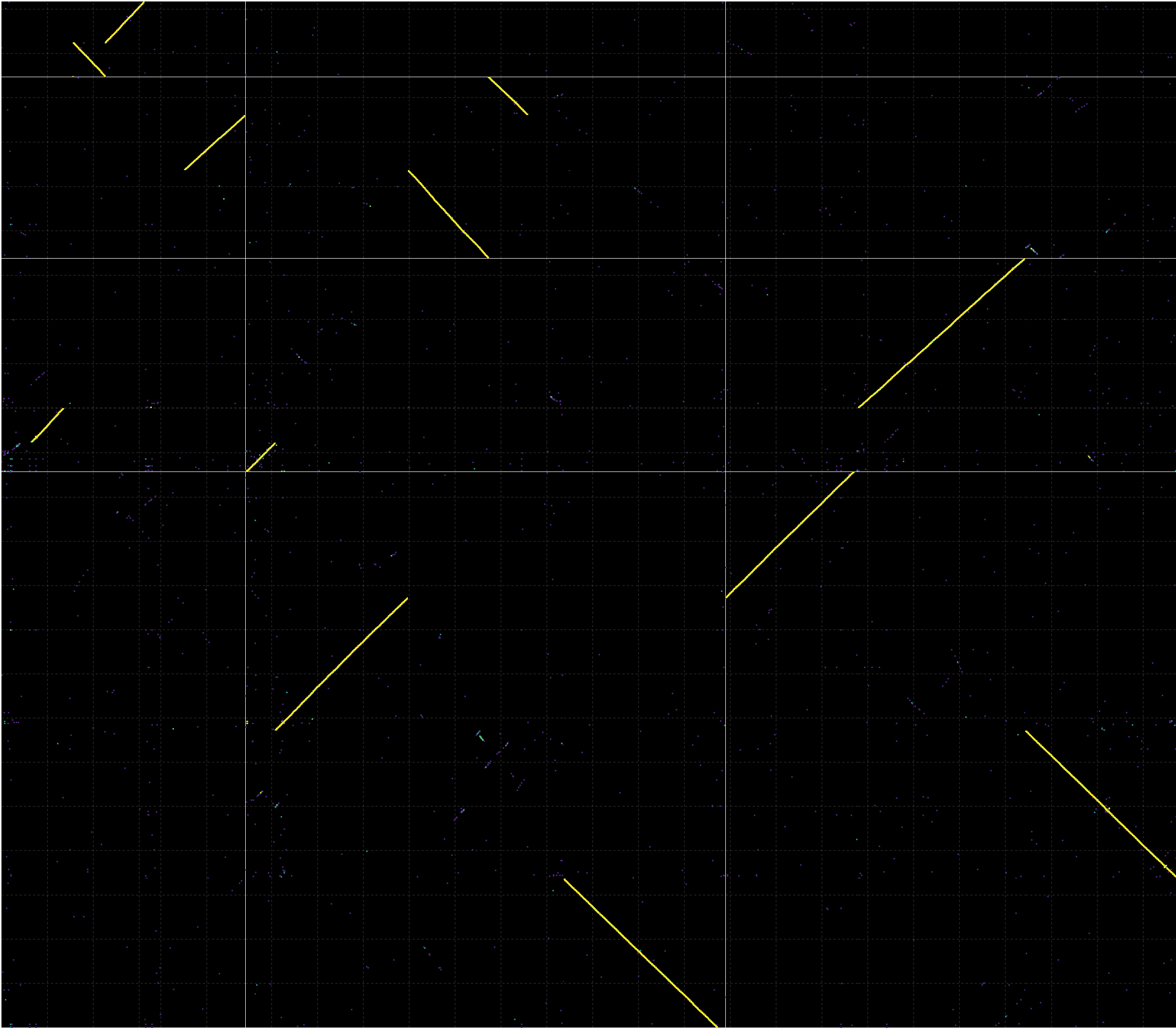
Ler_3: gene rank order position (22592 genes w/ blast hits), grids every 1000 genes



Sha: gene rank order position (25246 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 (21044 genes w/ blast hits), grids every 1000 genes



Sha: gene rank order position (21413 genes w/ blast hits), grids every 1000 genes