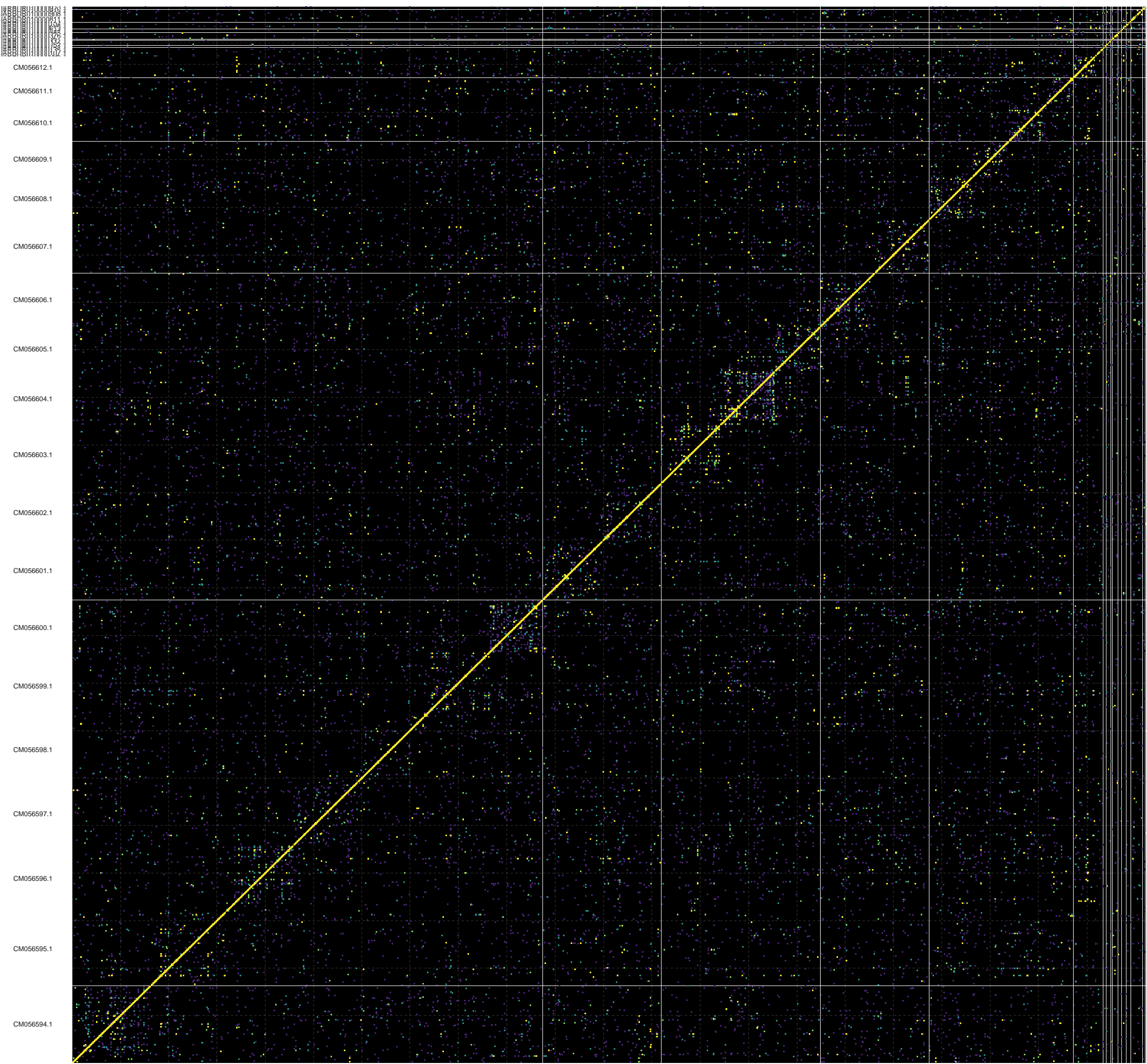All blast hits with score > 50, 8/8−gene x/y windows (heatmap range: 2−5+ hits/window)

Tgra: gene rank order position (23923 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)