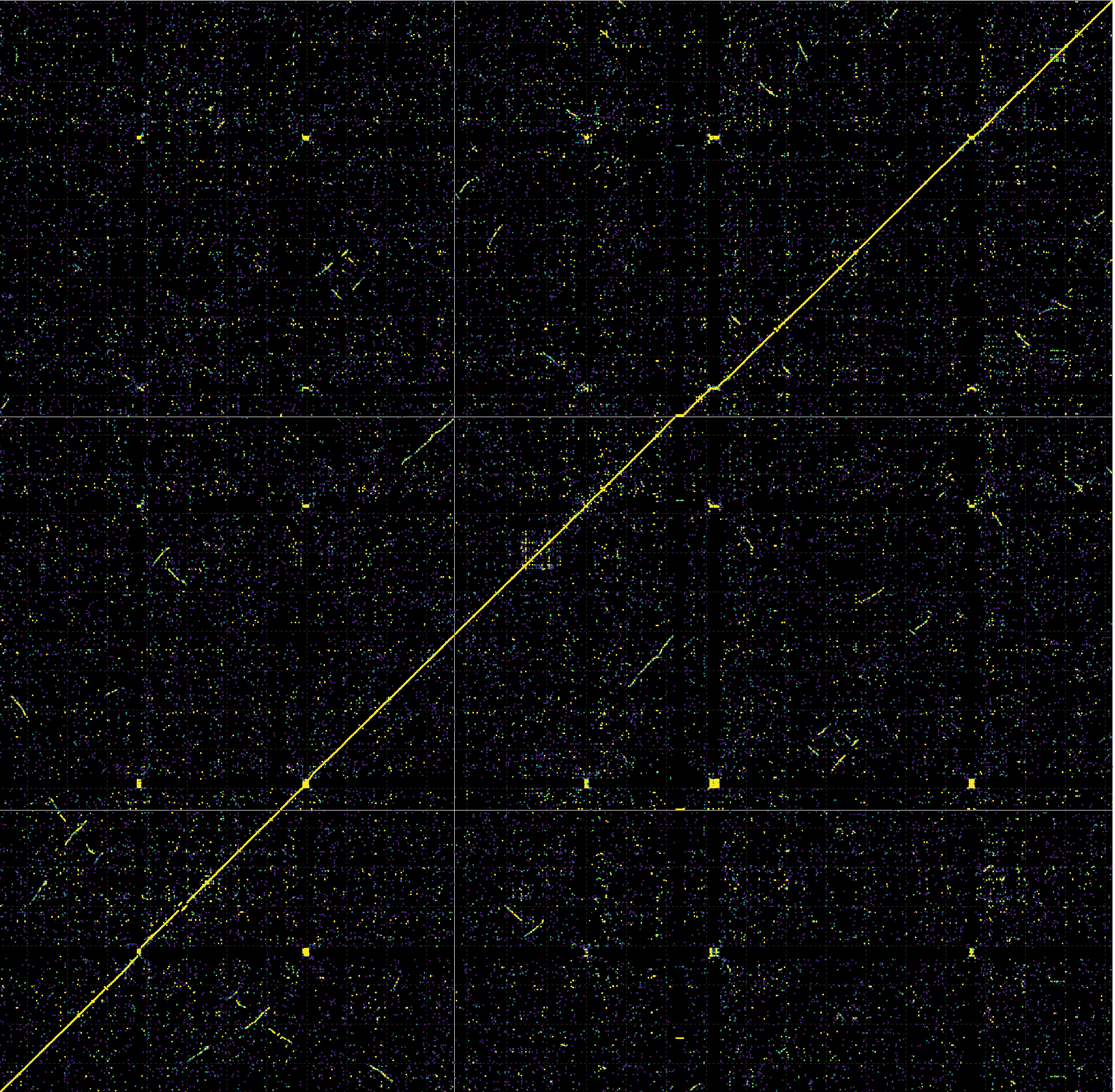


All blast hits with score > 50, 10/10–gene x/y windows (heatmap range: 2–5+ hits/window)

Taz_0: gene rank order position (27732 genes w/ blast hits), grids every 1000 genes



Altai_5: gene rank order position (27862 genes w/ blast hits), grids every 1000 genes

Blast hits where query and target are in the same orthogroup, 10/10-gene x/y windows (heatmap range: 1–5+ hits/window)

Taz_0: gene rank order position (26435 genes w/ blast hits), grids every 1000 genes

Altai_5: gene rank order position (26330 genes w/ blast hits), grids every 1000 genes

