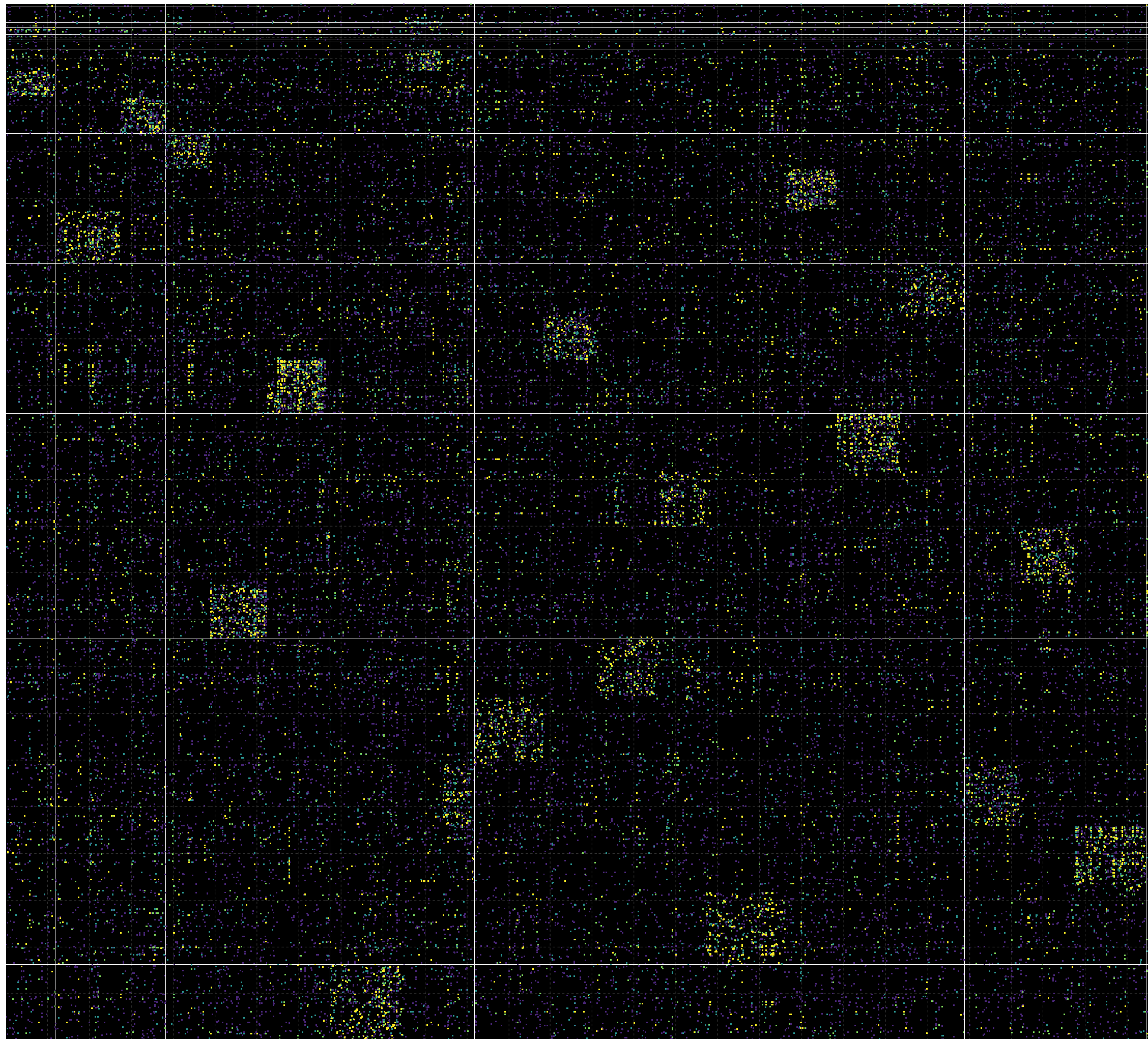
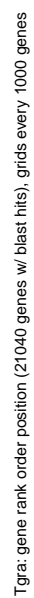


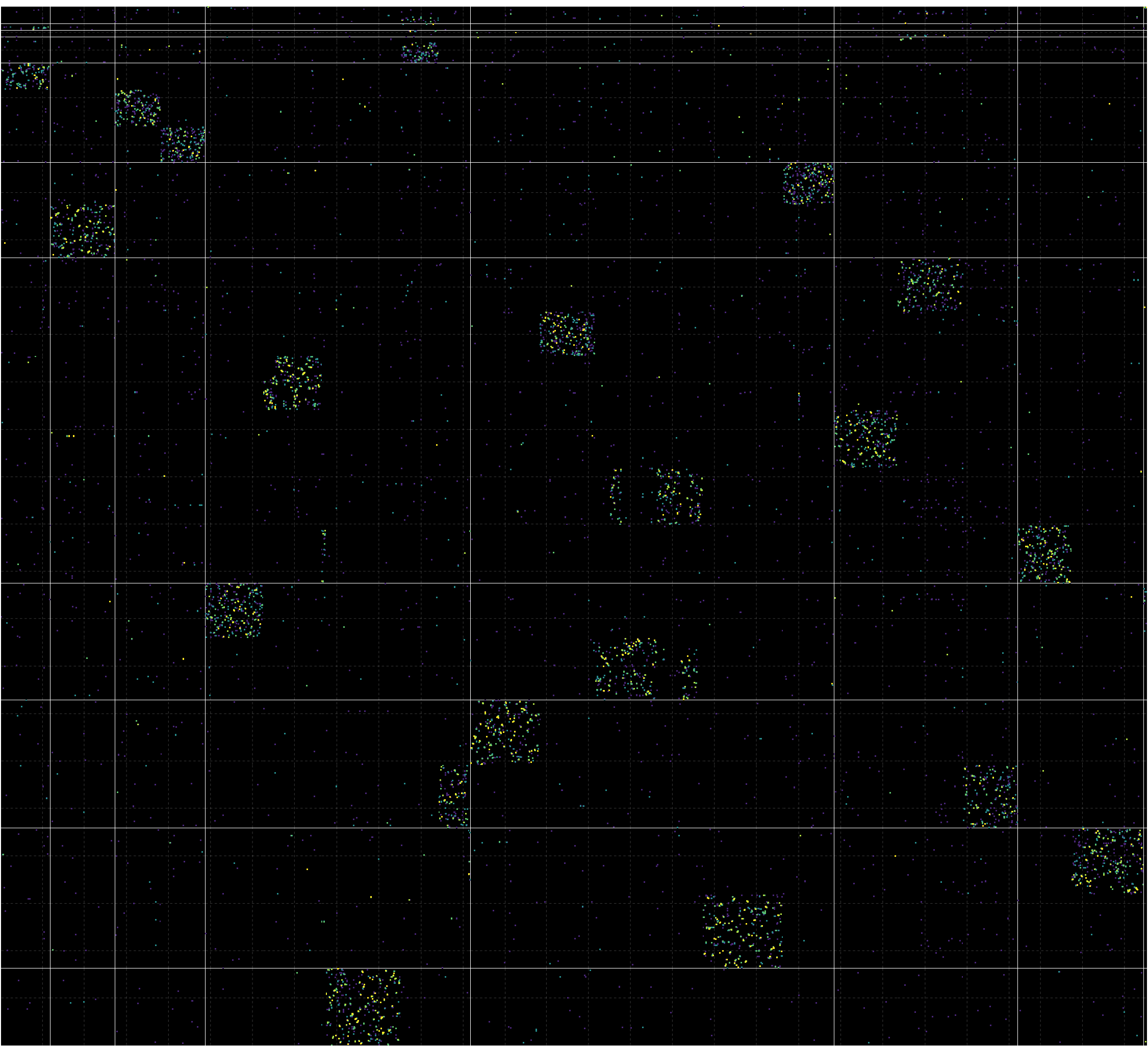
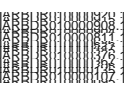
All blast hits with score > 50, 9/8-gene x/y windows (heatmap range: 2–5+ hits/window)



Azeb: gene rank order position (22377 genes w/ blast hits), grids every 1000 genes

Blast hits where query and target are in the same orthogroup, 9/8–gene x/y windows (heatmap range: 1–5+ hits/window)

Tgra: gene rank order position (15518 genes w/ blast hits), grids every 1000 genes



HIC_scaffold_1.0
HIC_scaffold_2.0
HIC_scaffold_3.0
HIC_scaffold_4.0
HIC_scaffold_5.0
HIC_scaffold_7.0
HIC_scaffold_8.0
HIC_scaffold_9.0
HIC_scaffold_10.0
HIC_scaffold_11.0
HIC_scaffold_12.0
HIC_scaffold_13.0
HIC_scaffold_14.0
HIC_scaffold_15.0
HIC_scaffold_16.0
HIC_scaffold_17.0
HIC_scaffold_18.0
HIC_scaffold_19.0
HIC_scaffold_21.0
HIC_scaffold_8.8335562
HIC_scaffold_12.310085
HIC_scaffold_5.9238104

Azeb: gene rank order position (15351 genes w/ blast hits), grids every 1000 genes