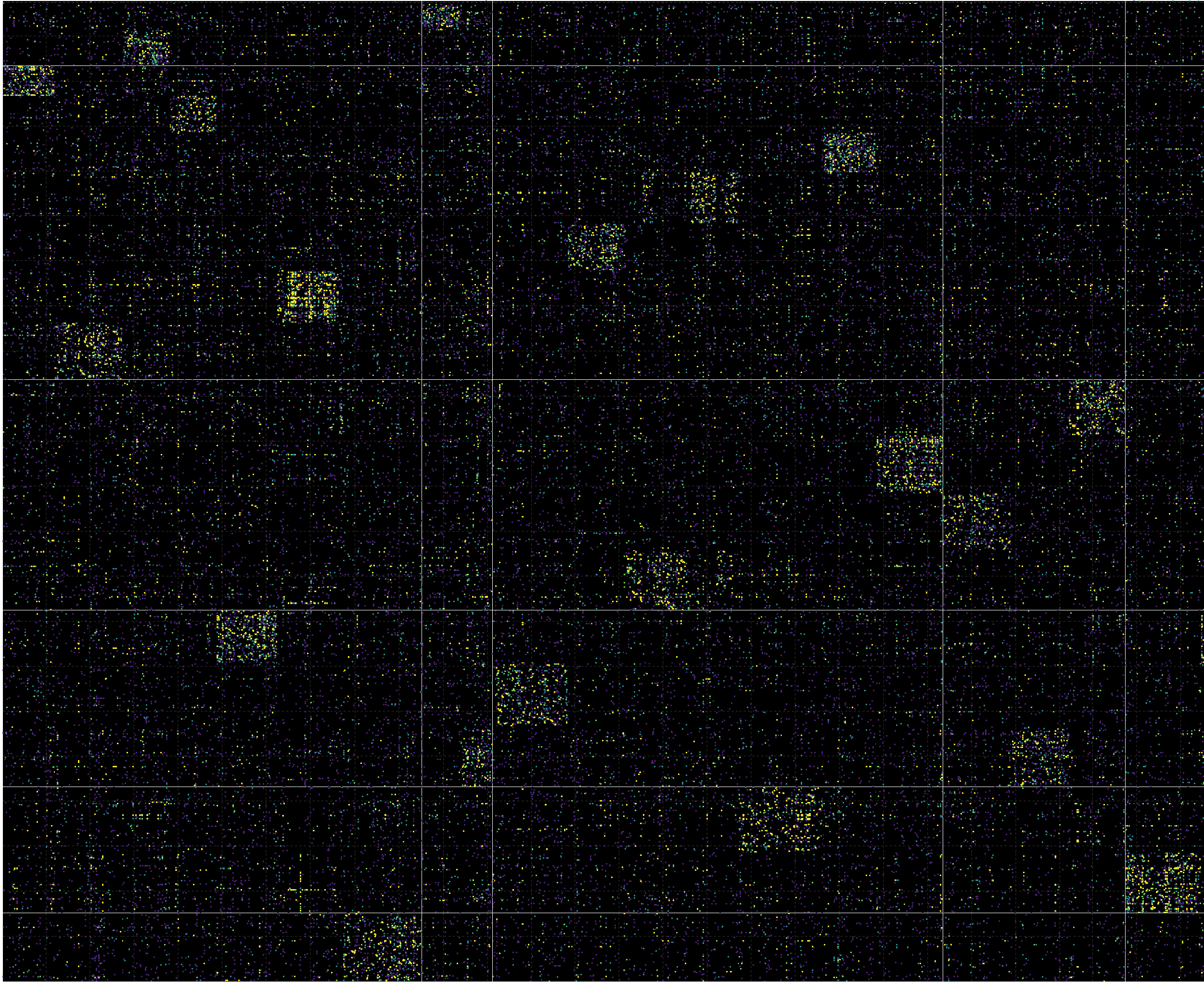


All blast hits with score > 50, 8/8-gene x/y windows (heatmap range: 2-5+ hits/window)

Sbro: gene rank order position (20050 genes w/ blast hits), grids every 1000 genes

Lachesis_group18
Lachesis_group17
Lachesis_group16
Lachesis_group15
Lachesis_group14
Lachesis_group13
Lachesis_group12
Lachesis_group11
Lachesis_group10
Lachesis_group9
Lachesis_group8
Lachesis_group7
Lachesis_group6
Lachesis_group5
Lachesis_group4
Lachesis_group3
Lachesis_group2
Lachesis_group1
Lachesis_group0



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

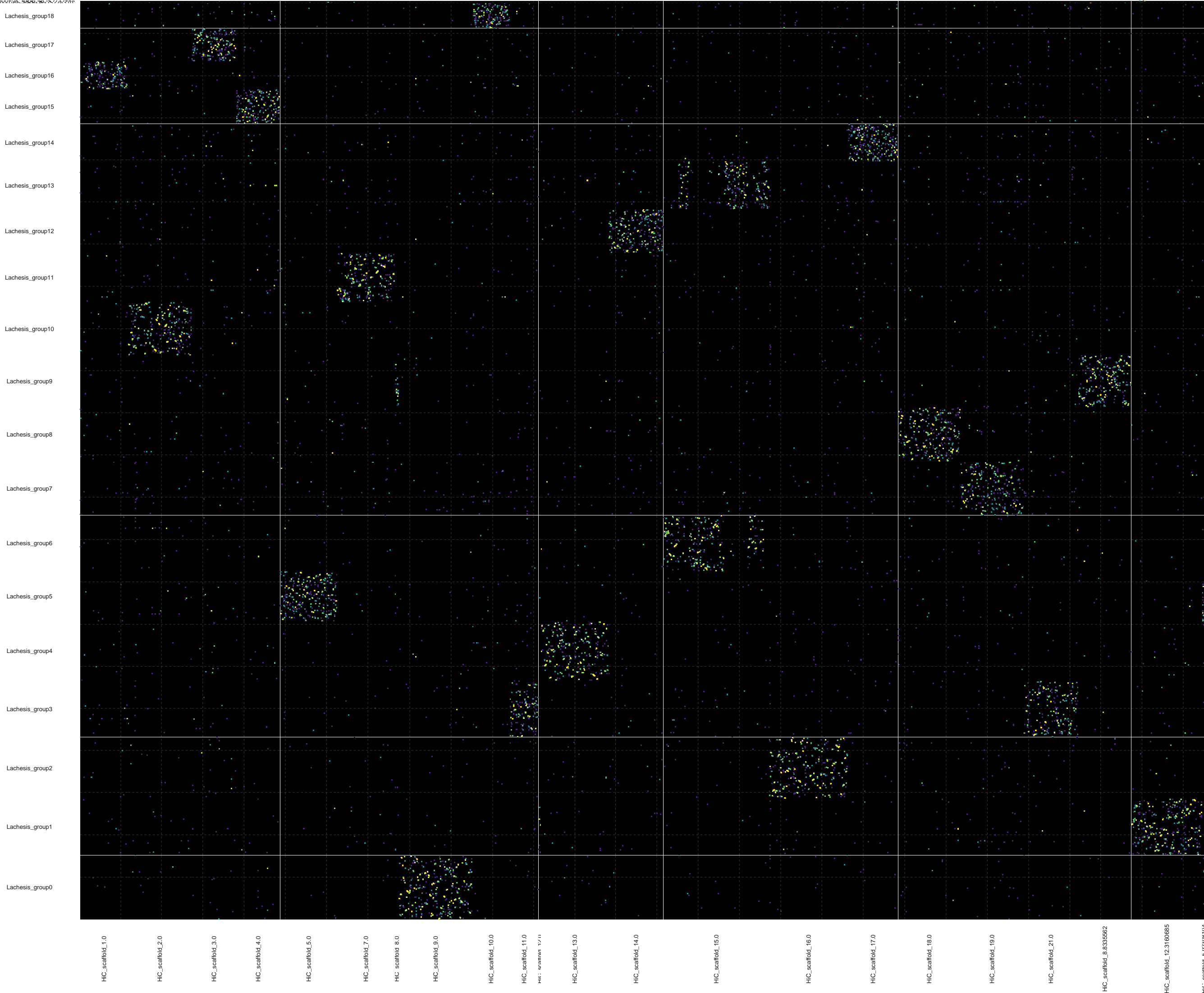
HIC_scaffold_12.3160685

HIC_scaffold_12.3160685

Azeb: gene rank order position (24111 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)

Sbro: gene rank order position (16414 genes w/ blast hits), grids every 1000 genes



Azeb: gene rank order position (17606 genes w/ blast hits), grids every 1000 genes

HIC_scaffold_5.9248114