

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

An_1: gene rank order position (24615 genes w/ blast hits), grids every 1000 genes

scaffold_679_pilon

contig_591_pilon

contig_589_pilon

contig_509_pilon

contig_449_pilon

contig_118_pilon

contig_102_pilon

contig_101_pilon

scaffold_88_pilon

contig_3_pilon

Chr1

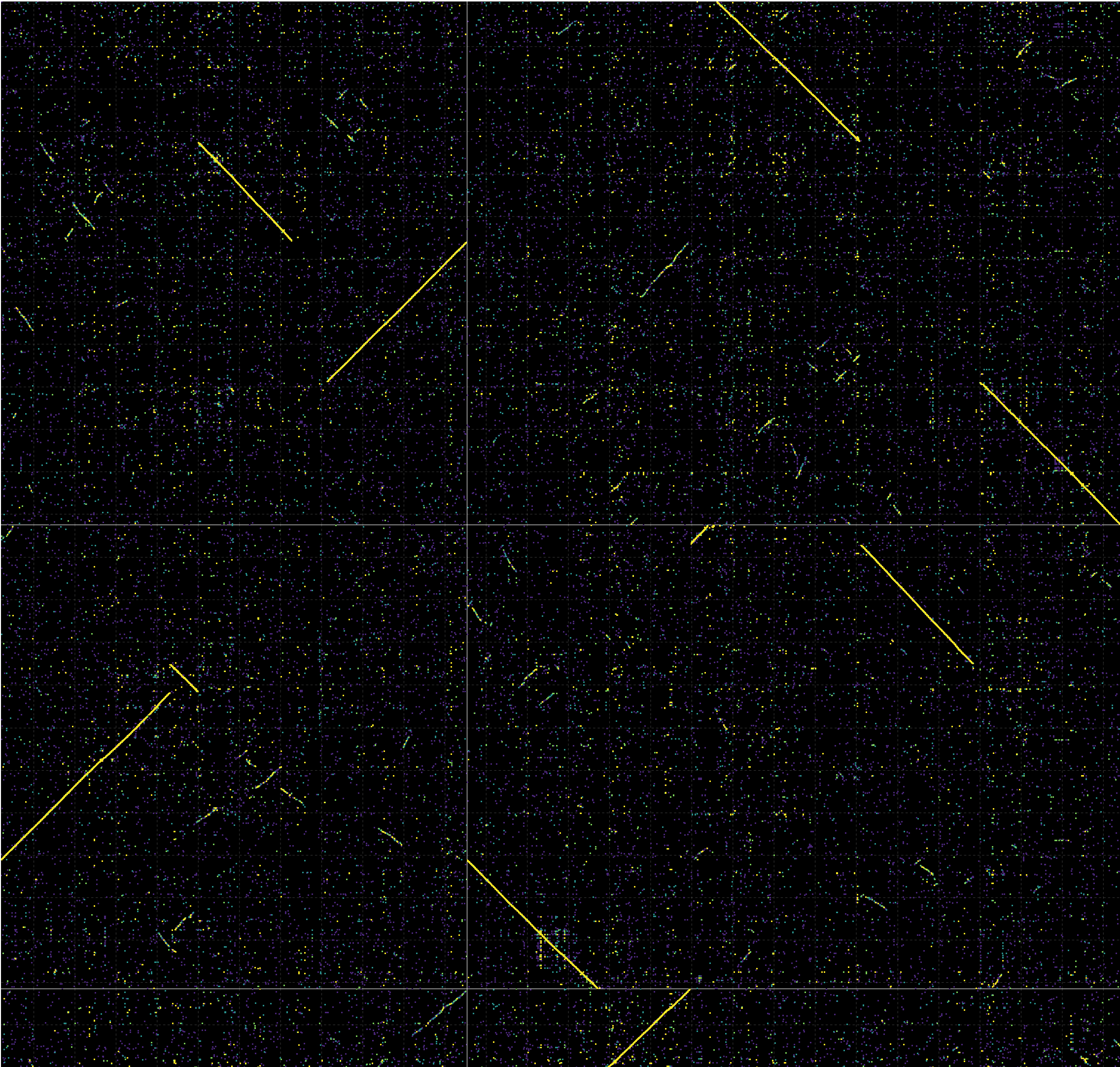
Chr2

Chr3

Chr4

Chr5

TAIR10: gene rank order position (26268 genes w/ blast hits), grids every 1000 genes



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

An_1: gene rank order position (23486 genes w/ blast hits), grids every 1000 genes

scaffold_679_pilon

contig_591_pilon

contig_589_pilon

contig_509_pilon

contig_449_pilon

contig_118_pilon

contig_102_pilon

contig_101_pilon

scaffold_88_pilon

contig_3_pilon

Chr1

Chr2

Chr3

Chr4

Chr5

TAIR10: gene rank order position (24309 genes w/ blast hits), grids every 1000 genes

