

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

An_1: gene rank order position (24978 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

contig_3_pilon

scaffold_88_pilon

contig_101_pilon

contig_102_pilon

contig_118_pilon

contig_449_pilon

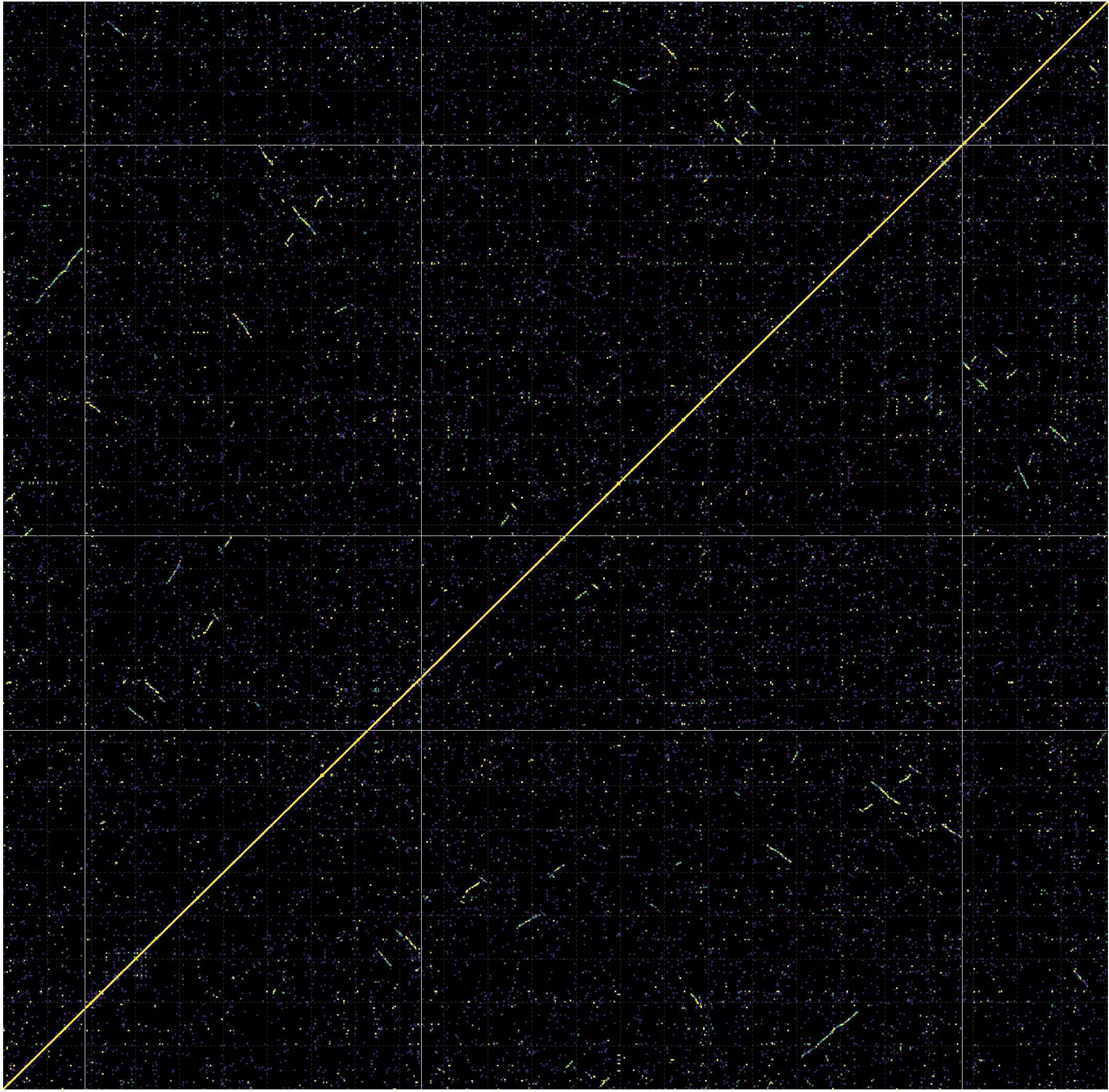
contig_509_pilon

contig_589_pilon

contig_591_pilon

scaffold_679_pilon

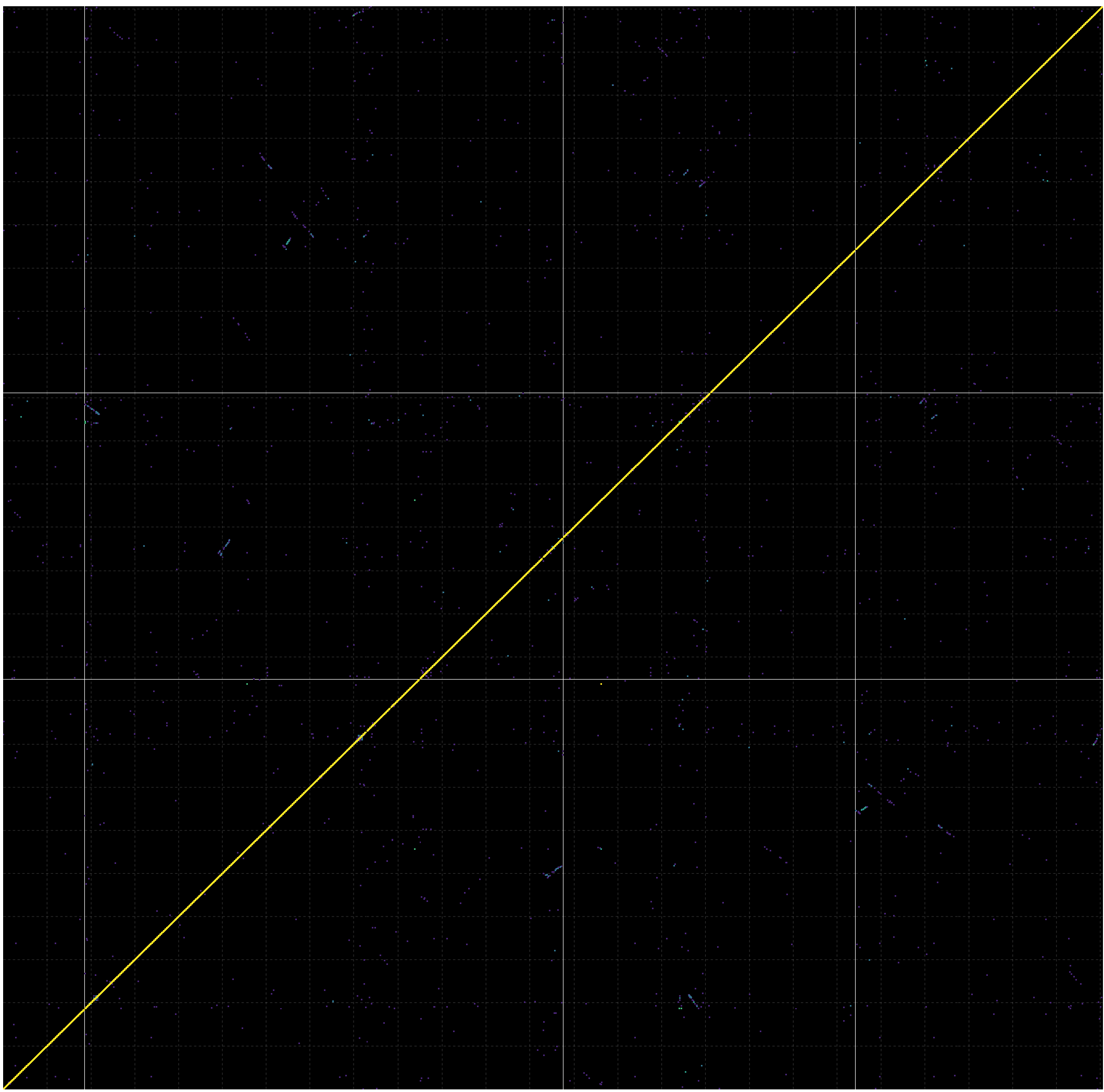
An_1: gene rank order position (24978 genes w/ blast hits), grids every 1000 genes



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

An_1: gene rank order position (25031 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



contig_3_pilon scaffold_88_pilon contig_101_pilon contig_102_pilon contig_118_pilon contig_449_pilon contig_509_pilon contig_589_pilon contig_591_pilon scaffold_679_pilon

An_1: gene rank order position (25031 genes w/ blast hits), grids every 1000 genes