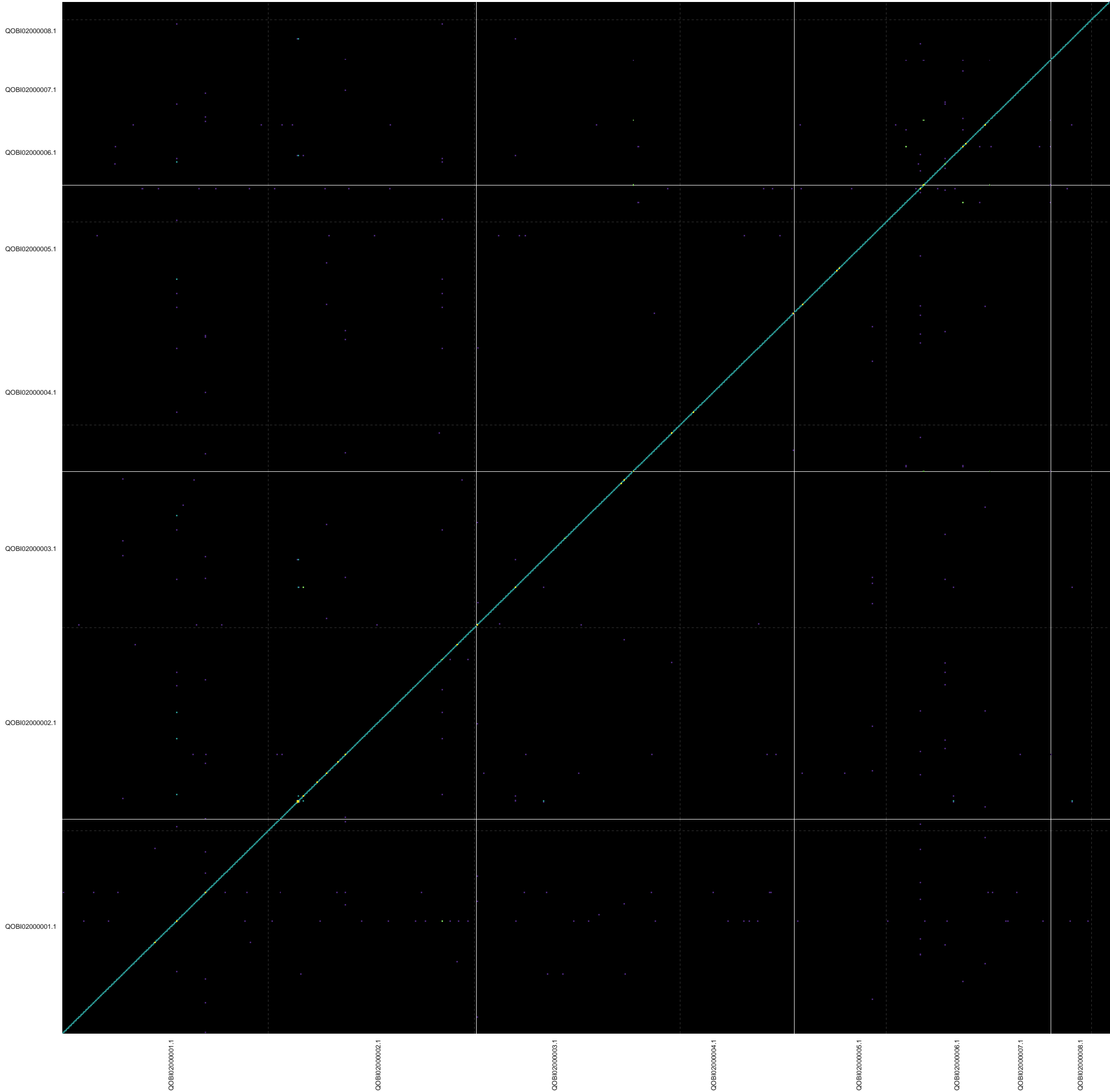


All blast hits with score > 50, 2/2-gene x/y windows (heatmap range: 2-5+ hits/window)

Clus_L17: gene rank order position (5088 genes w/ blast hits), grids every 1000 genes



Clus_L17: gene rank order position (5088 genes w/ blast hits), grids every 1000 genes

Blast hits where query and target are in the same orthogroup, 2/2-gene x/y windows (heatmap range: 1-5+ hits/window)

