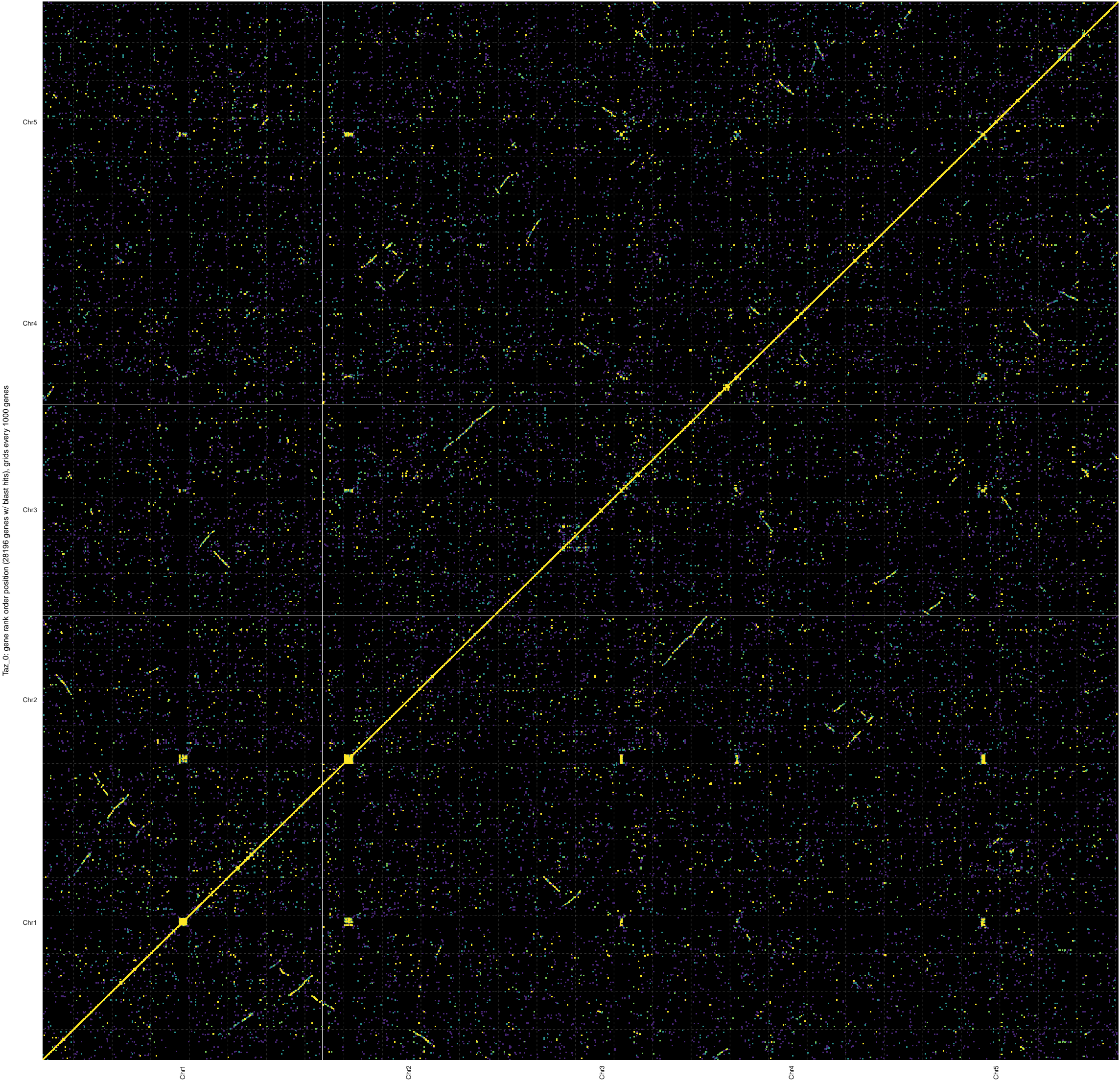


All blast hits with score > 50, 10/10–gene x/y windows (heatmap range: 2–5+ hits/window)

Taz_0: gene rank order position (28196 genes w/ blast hits), grids every 1000 genes



Taz_0: gene rank order position (28240 genes w/ blast hits), grids every 1000 genes

Blast hits where query and target are in the same orthogroup, 10/10–gene x/y windows (heatmap range: 1–5+ hits/window)

Taz_0: gene rank order position (28205 genes w/ blast hits), grids every 1000 genes

Taz_0: gene rank order position (28249 genes w/ blast hits), grids every 1000 genes

