

Syntenic anchor blast hits, colored by block ID

Ler_3: gene rank order position (20119 genes with blast hits), gridlines every 1000 genes

contig_538_pilon
scaffold_448_pilon
contig_446_pilon
contig_444_pilon
scaffold_322_pilon
contig_297_pilon
scaffold_290_pilon
contig_277_pilon
scaffold_250_pilon
contig_139_pilon

Chr1

Chr2

Chr3

Chr4

Chr5

TAIR10: gene rank order position (20119 genes with blast hits), gridlines every 1000 genes

