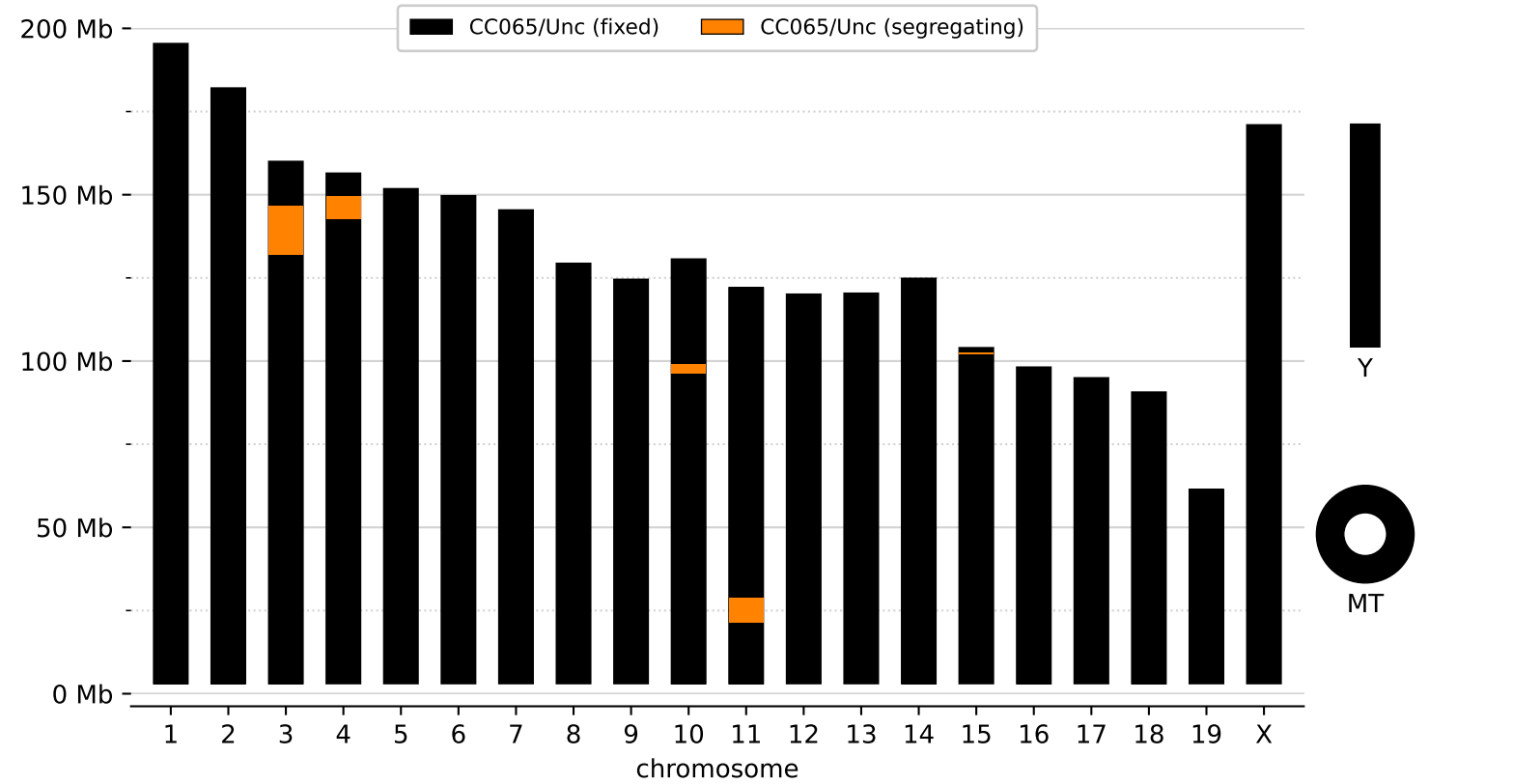


Summary: Incipient Recombinant Inbred strain from the Collaborative Cross population. This strain is derived from 8 inbred founders: A/J, C57BL/6J, 129S1/SvImJ, NOD/ShiLtJ, NZO/HILtJ, CAST/EiJ, PWK/PhJ, and WSB/EiJ. The genome of this strain is not fully replicable because 1.2% is segregating for the founder strains. Note that MiniMUGA may fail to detect additional small regions of residual heterozygosity. CC strains are maintained as live colonies at UNC and GQC is updated approximately every 18 months. This GQC report is based on the MiniMUGA G3 2024 pipeline (PMCID: PMC11457065) using 31 samples genotyped in 2025.



Genetic Background	Zygosity	Genome %	Estimated # Segregating SNPs
CC065/Unc (fixed)	Homozygous	98.8%	N/A
CC065/Unc (segregating)	Heterozygous	1.2%	147,000
Total		100.0%	147,000

Constructs Detected
None