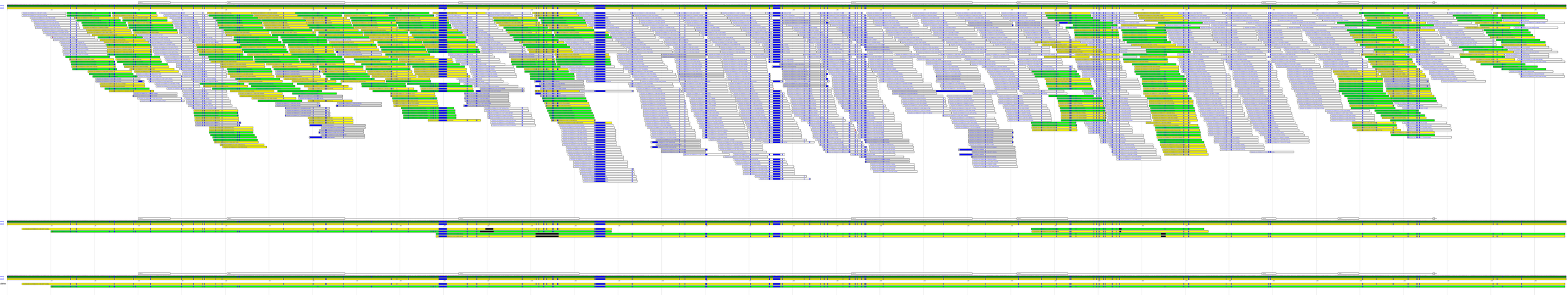
chr1:0-10,000,000 (reference)
chr1:0-10,000,000 (contigs)
a. Read alignment
chr1:0-10,000,000 (reference)
chr1:0-10,000,000 (contigs)
b. Assembly
chr1:0-10,000,000 (reference)
chr1:0-10,000,000 (contigs)
c. Assembly with known alleles