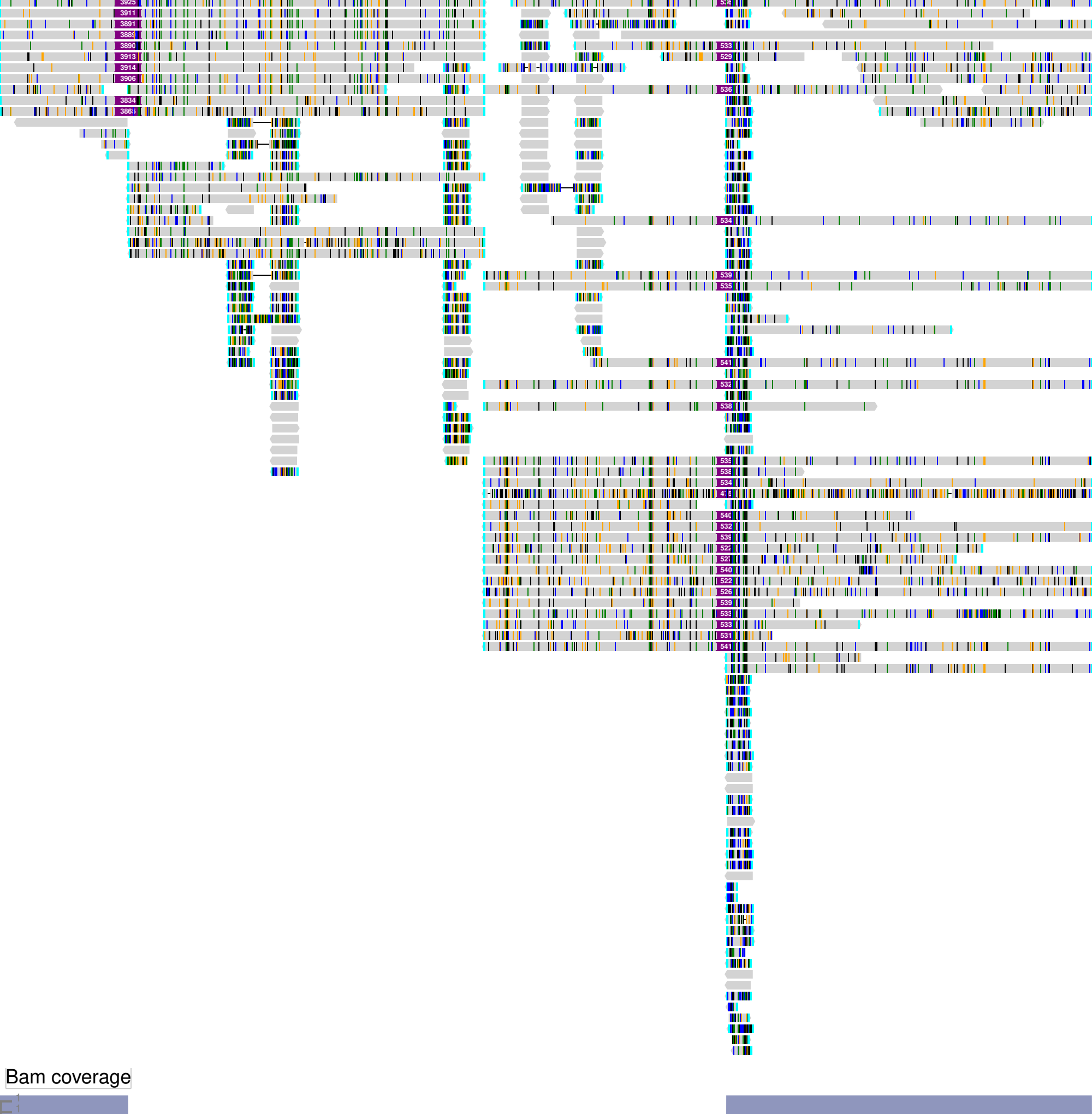


variants

Bam coverage

82
55

Child reads align to father allele1



Bam coverage

Child haplotypes align to father allele1



Bam coverage

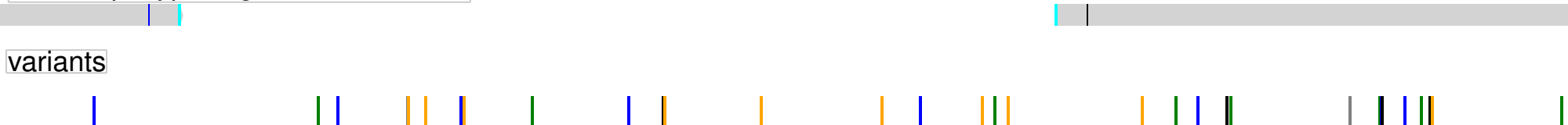
85
57

Child reads align to father allele2



Bam coverage

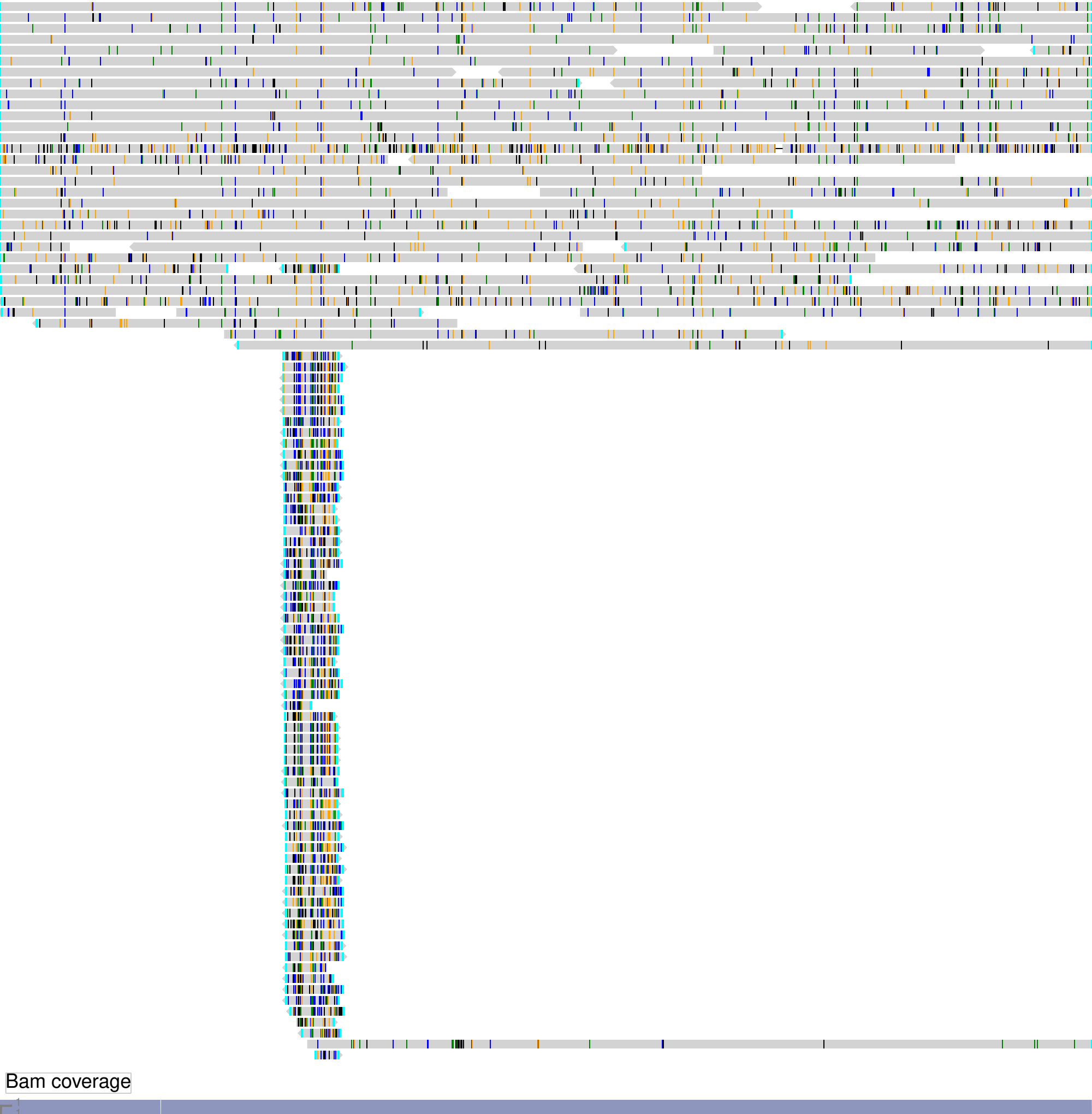
Child haplotypes align to father allele2



Bam coverage

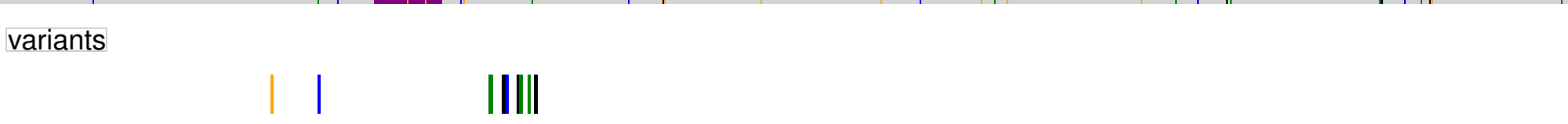
83
55

Child reads align to mother allele1



Bam coverage

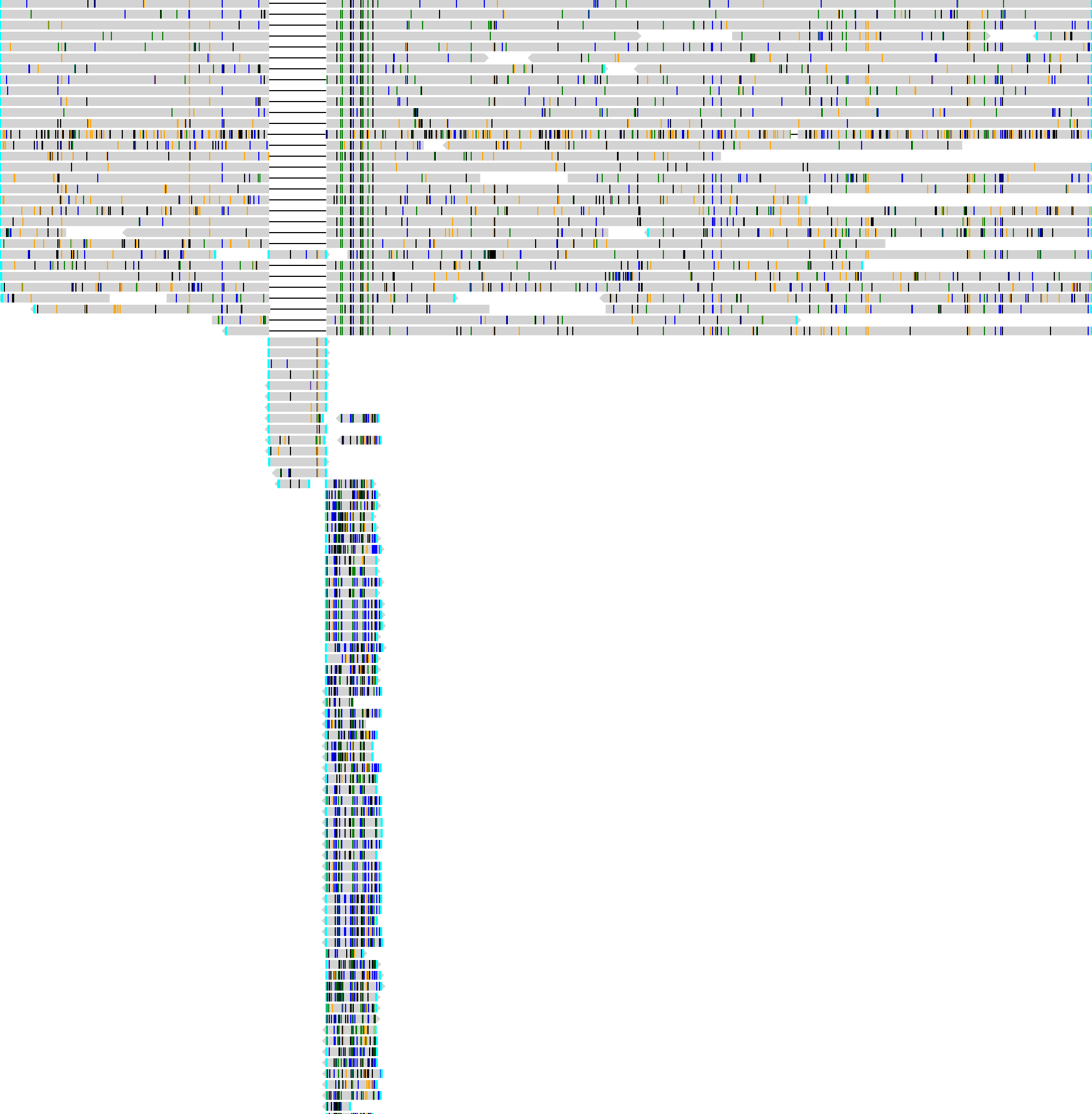
Child haplotypes align to mother allele1



Bam coverage

102
101

Child reads align to mother allele2



Bam coverage

Child haplotypes align to mother allele2

