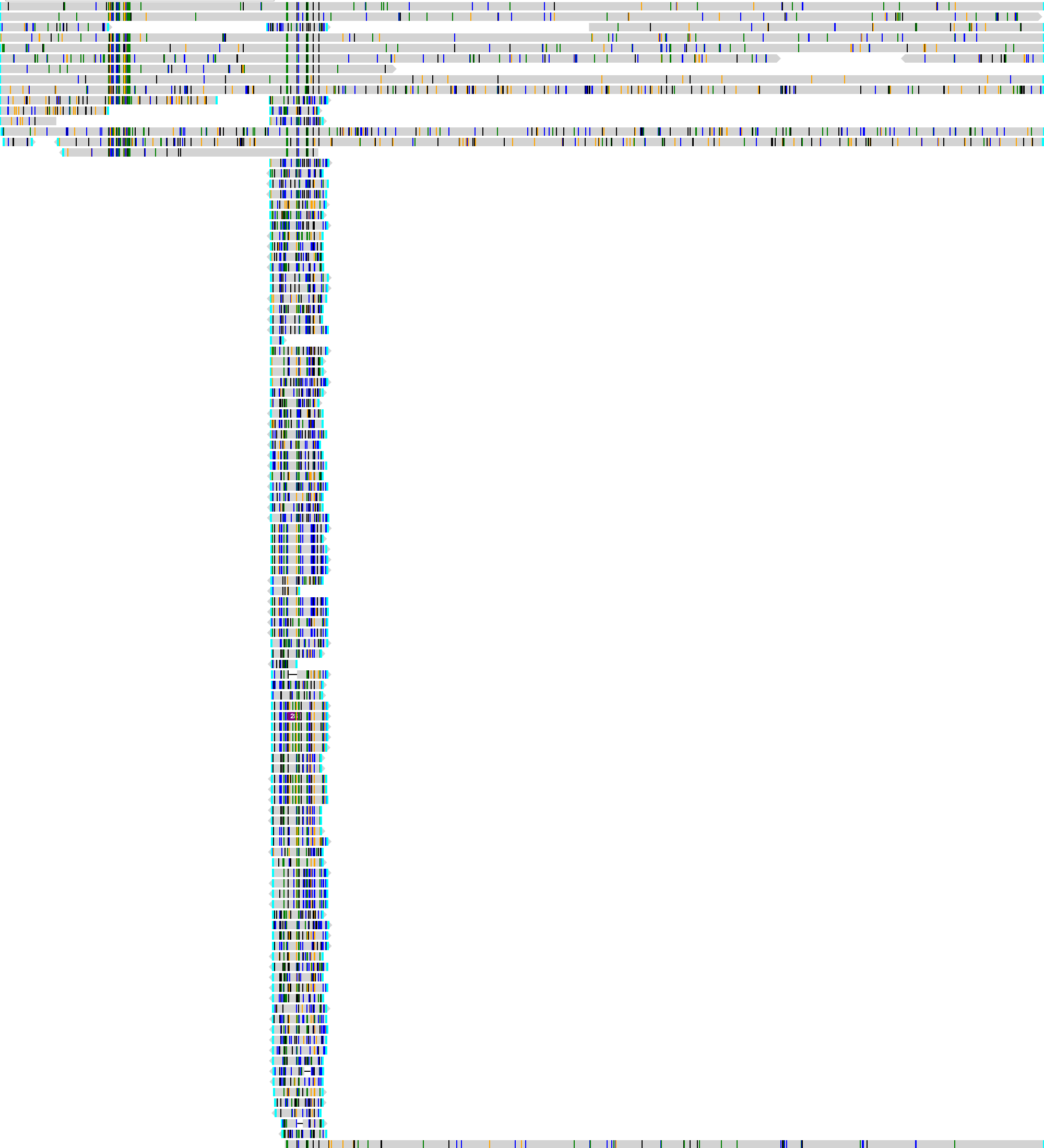


variants

Bam coverage

93
62

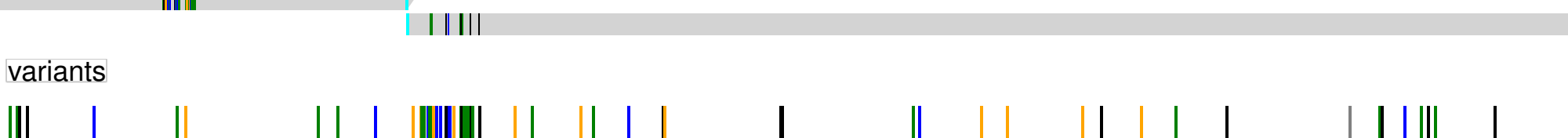
Child reads align to father allele1



Bam coverage

93
62

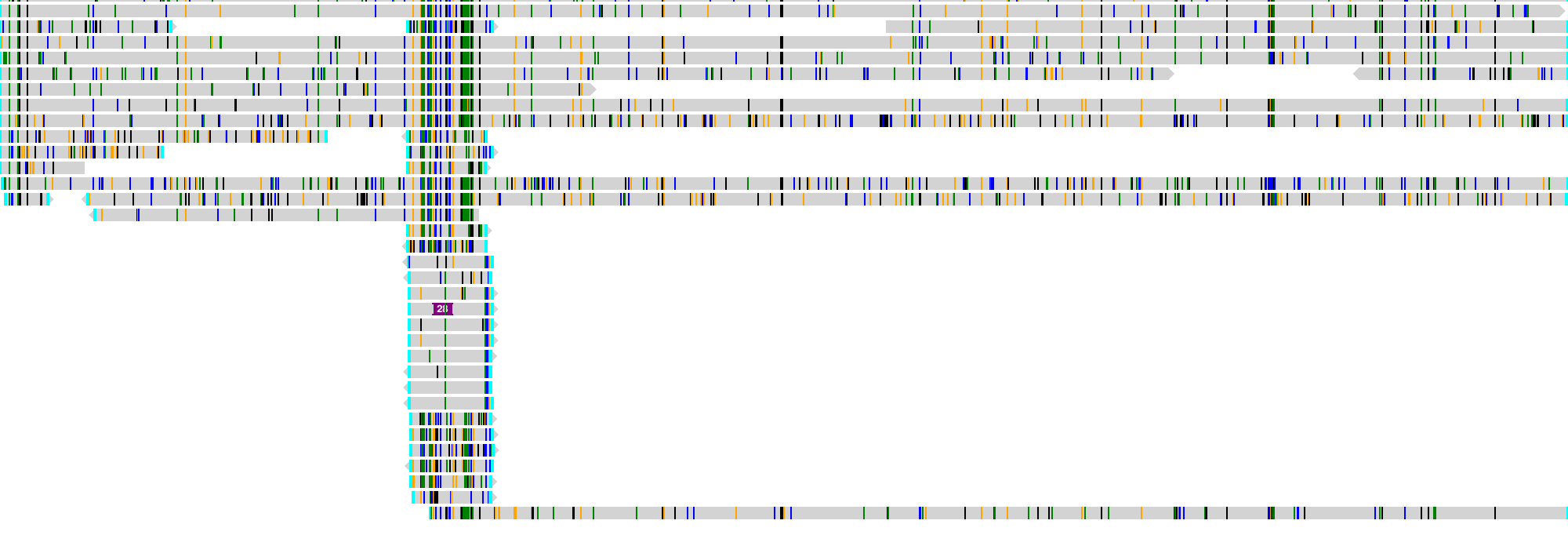
Child haplotype align to father allele1



Bam coverage

30
20

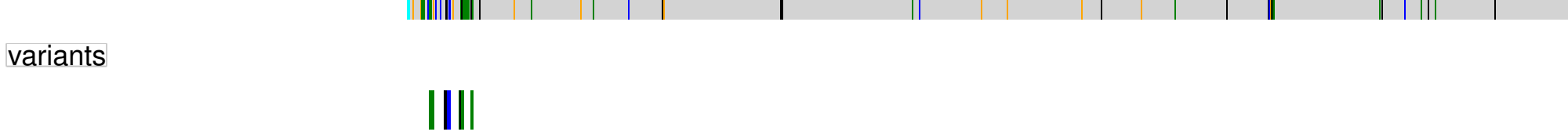
Child reads align to father allele2



Bam coverage

30
20

Child haplotype align to father allele2



Bam coverage

79
52

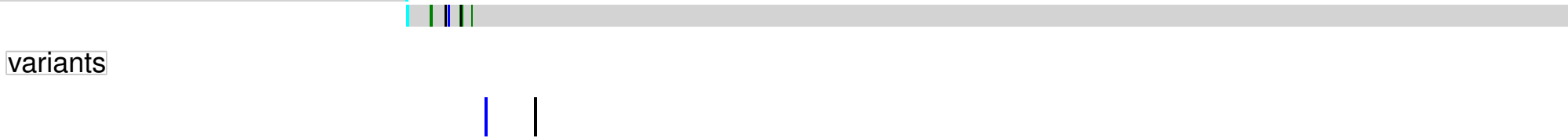
Child reads align to mother allele1



Bam coverage

79
52

Child haplotype align to mother allele1



Bam coverage

83
55

Child reads align to mother allele2



Bam coverage

83
55

Child haplotype align to mother allele2