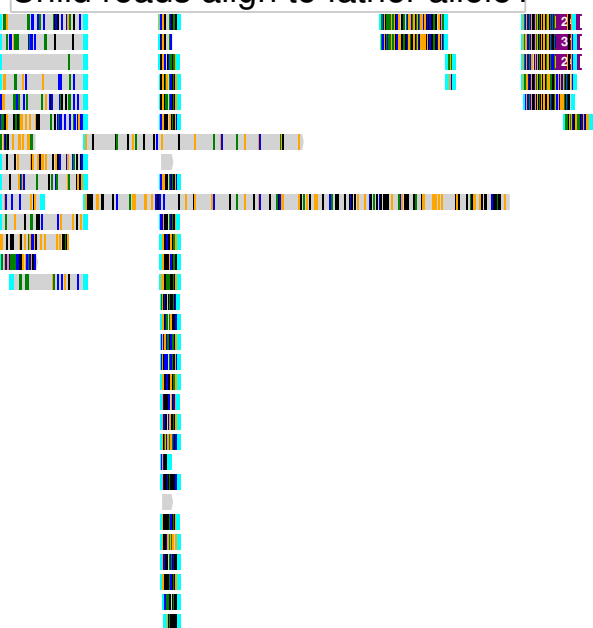


variants

Bam coverage

37
25
25

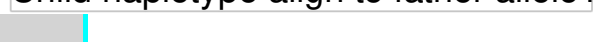
Child reads align to father allele1



Bam coverage

37
25
25

Child haplotype align to father allele1

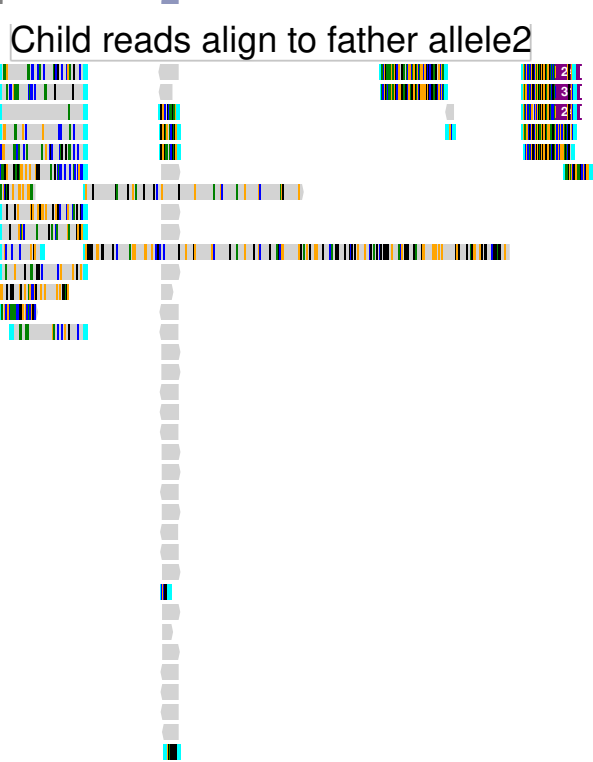


variants

Bam coverage

56
37
19

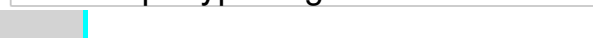
Child reads align to father allele2



Bam coverage

56
37
19

Child haplotype align to father allele2

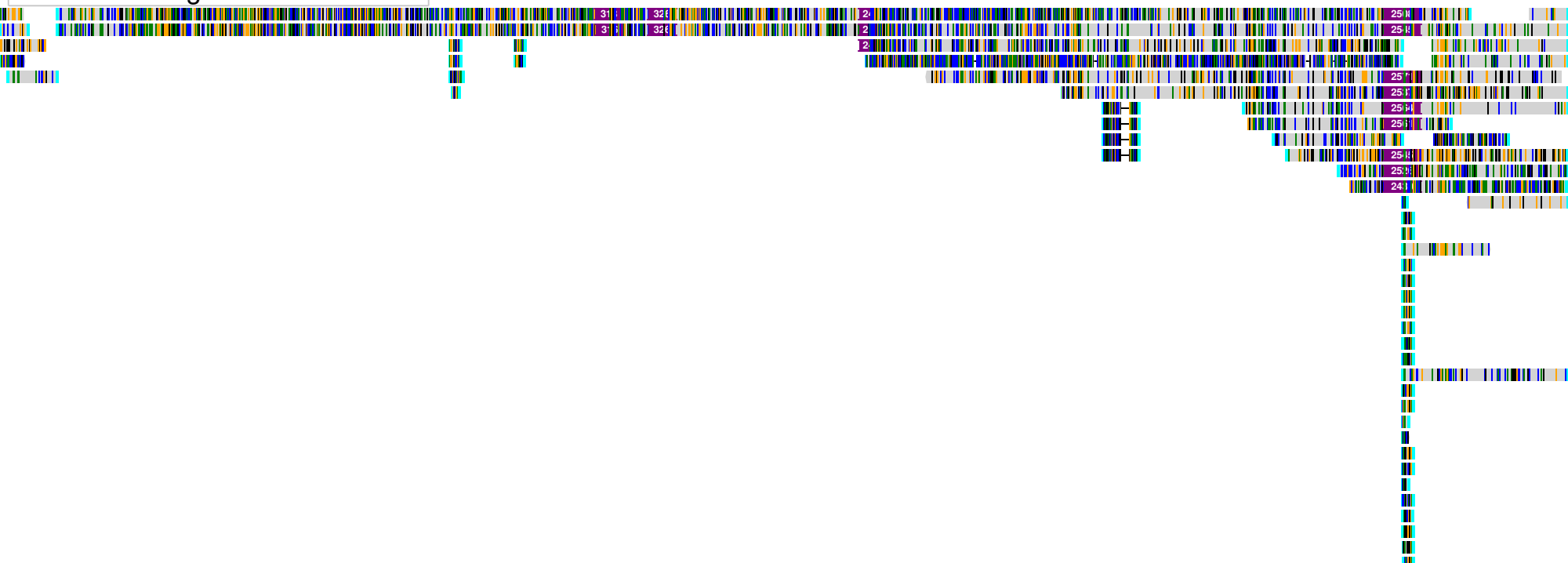


variants

Bam coverage

29
19
19

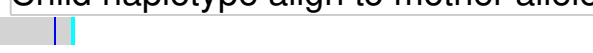
Child reads align to mother allele1



Bam coverage

29
19
19

Child haplotype align to mother allele1

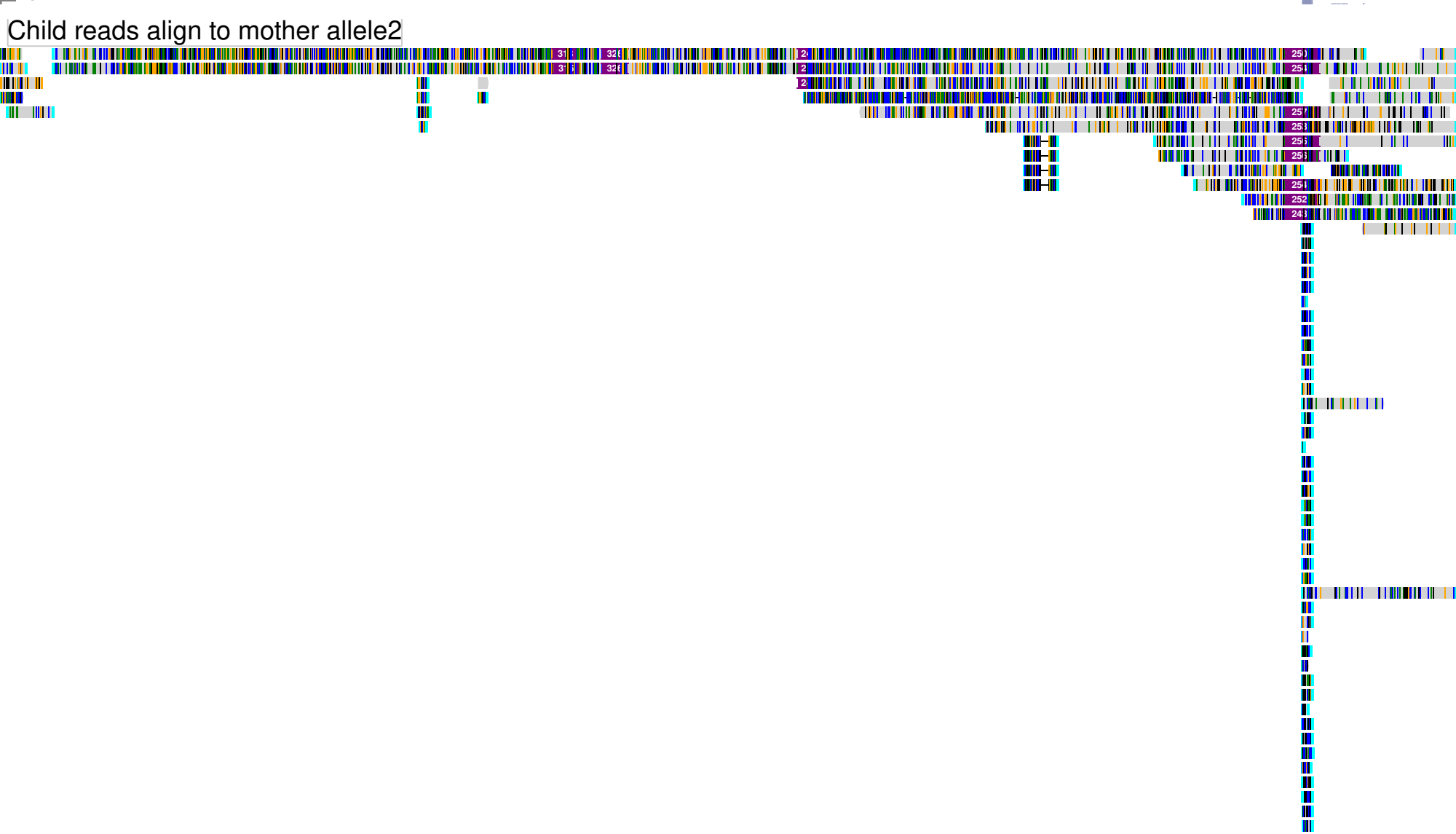


variants

Bam coverage

48
32
16

Child reads align to mother allele2



Bam coverage

48
32
16

Child haplotype align to mother allele2

