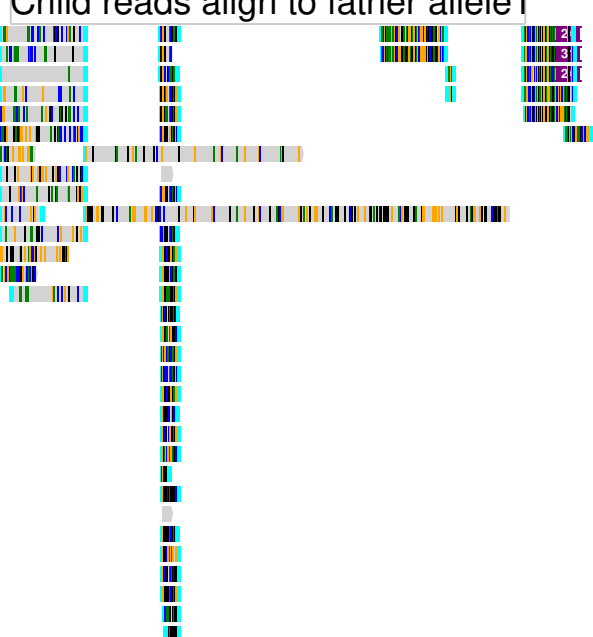


variants

Bam coverage

37
25
4

Child reads align to father allele1



Bam coverage

37
25
4

Child haplotype align to father allele1

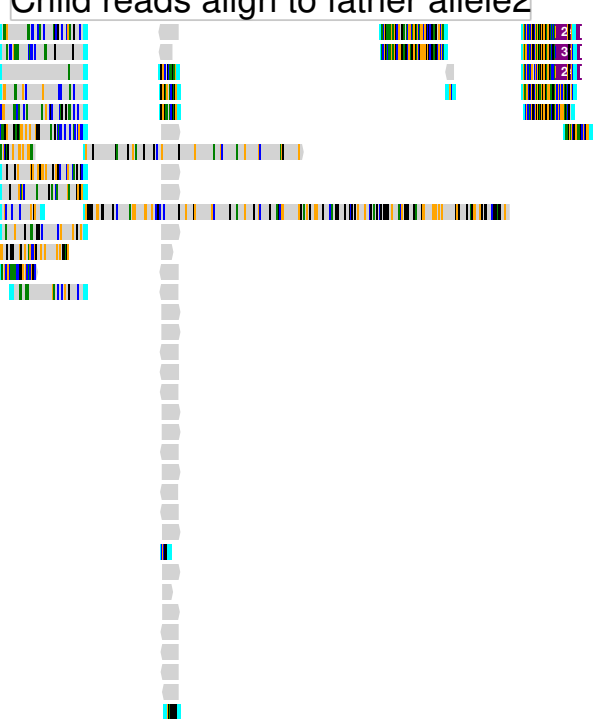


variants

Bam coverage

52
35
4

Child reads align to father allele2



Bam coverage

37
25
4

Child haplotype align to father allele2

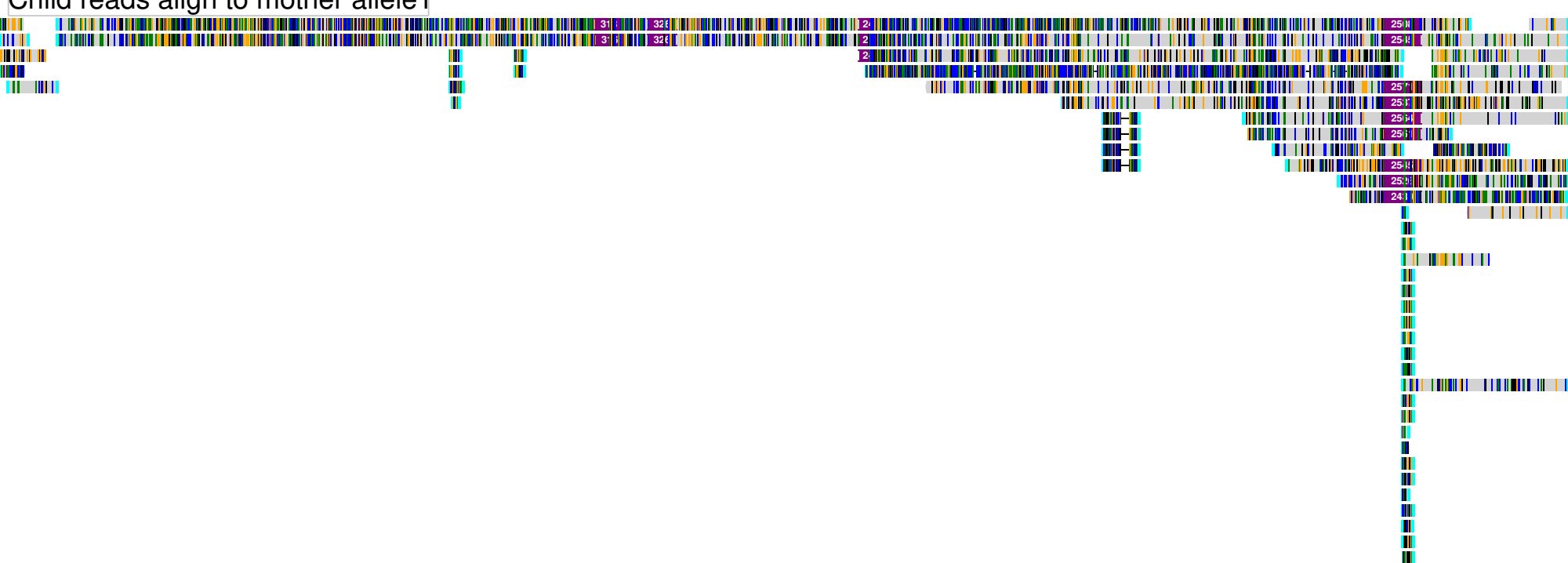


variants

Bam coverage

29
19
4

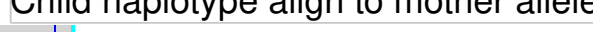
Child reads align to mother allele1



Bam coverage

37
25
4

Child haplotype align to mother allele1

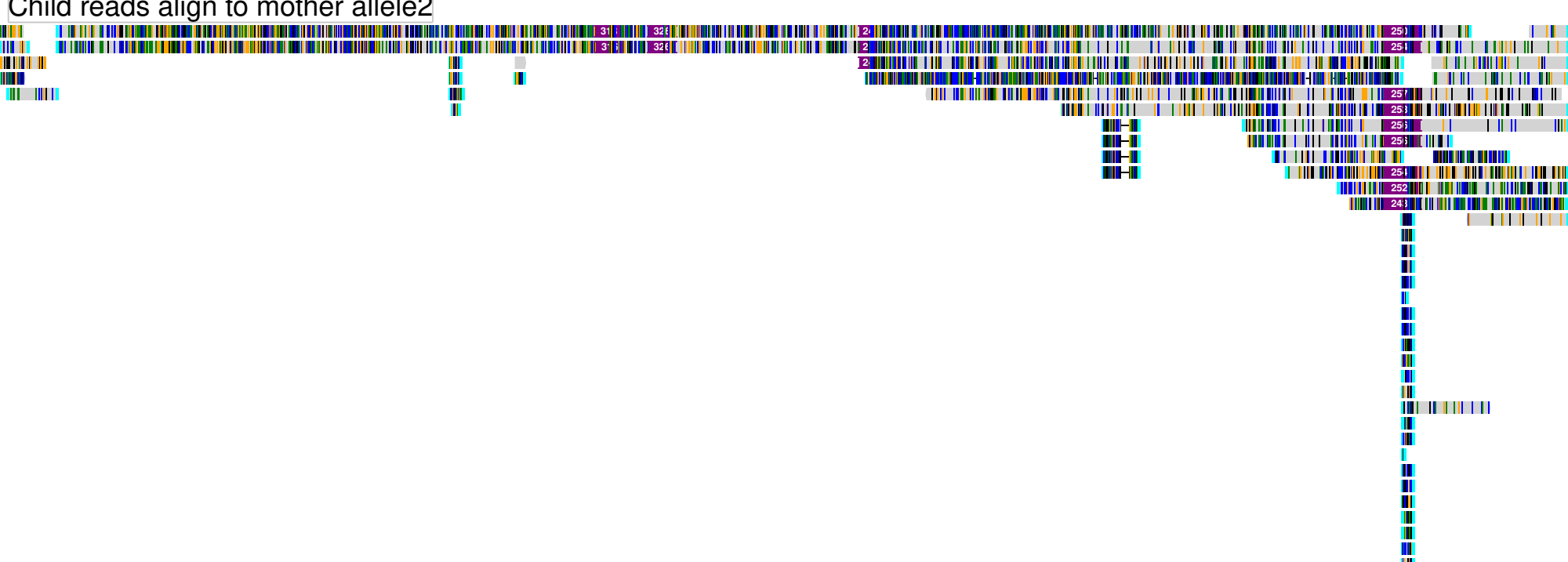


variants

Bam coverage

45
30
4

Child reads align to mother allele2



Bam coverage

37
25
4

Child haplotype align to mother allele2

