

**Sample: HG01257**

**Locus: HLA-A**

**Resolution: 4th field**

**Full typing result:**

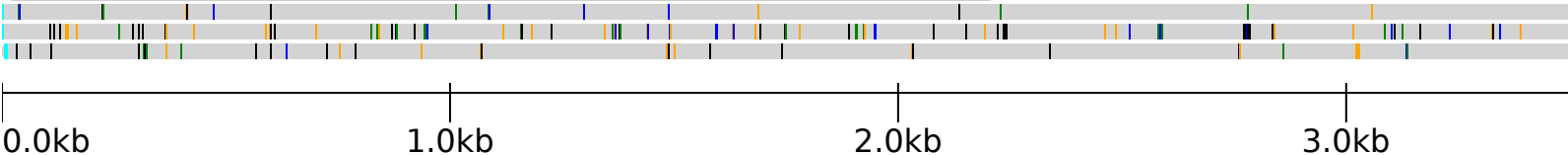
**HLA-A\*02:200**

bam coverage



variants

Alignment (reference: reconstructed gene haplotype 1)



HLA-A\*02:200 (identity: 99.97%)



**Sample: HG01257**

**Locus: HLA-B**

**Resolution: 4th field**

**Full typing result:**

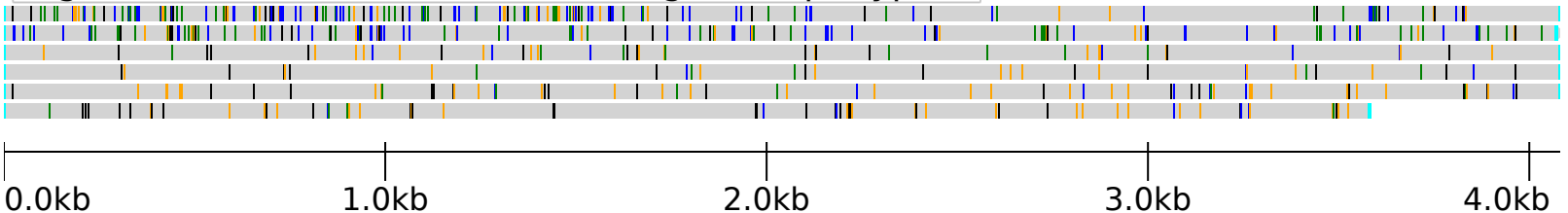
**HLA-B\*07:02:01:01**

bam coverage



variants

Alignment (reference: reconstructed gene haplotype 1)



HLA-B\*07:02:01:01 (identity: 100.00%)

**Sample: HG01257**

**Locus: HLA-B**

**Resolution: 4th field**

**Full typing result:**

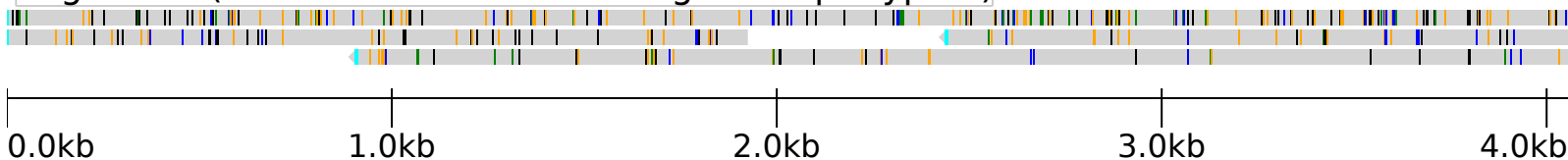
**HLA-B\*18:01:01:01**

bam coverage



variants

Alignment (reference: reconstructed gene haplotype 2)



HLA-B\*18:01:01:01 (identity: 99.98%)



Sample: HG01257

Locus: HLA-C

Resolution: 4th field

Full typing result:

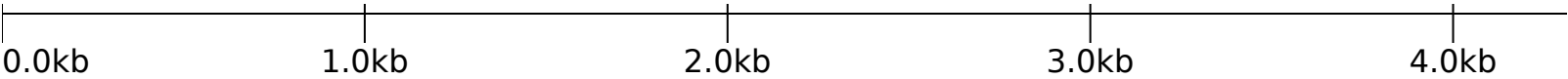
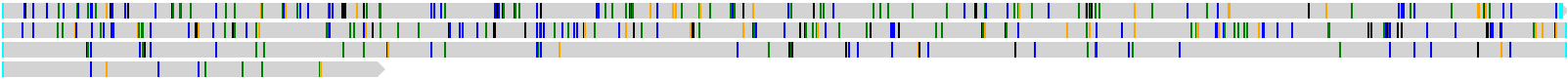
- HLA-C\*07:02:01:03
- HLA-C\*07:02:01:197
- HLA-C\*07:02:01:01
- HLA-C\*07:02:01:189
- HLA-C\*07:02:01:19
- HLA-C\*07:02:01:190
- HLA-C\*07:02:01:191
- HLA-C\*07:02:01:192
- HLA-C\*07:02:01:193
- HLA-C\*07:02:01:195

bam coverage



variants

Alignment (reference: reconstructed gene haplotype 2)



- HLA-C\*07:02:01:03 (identity: 100.00%)
- HLA-C\*07:02:01:197 (identity: 100.00%)
- HLA-C\*07:02:01:01 (identity: 99.98%)
- HLA-C\*07:02:01:189 (identity: 99.98%)
- HLA-C\*07:02:01:19 (identity: 99.98%)
- HLA-C\*07:02:01:190 (identity: 99.98%)
- HLA-C\*07:02:01:191 (identity: 99.98%)
- HLA-C\*07:02:01:192 (identity: 99.98%)
- HLA-C\*07:02:01:193 (identity: 99.98%)
- HLA-C\*07:02:01:195 (identity: 99.98%)

**Sample: HG01257**

**Locus: HLA-C**

**Resolution: 4th field**

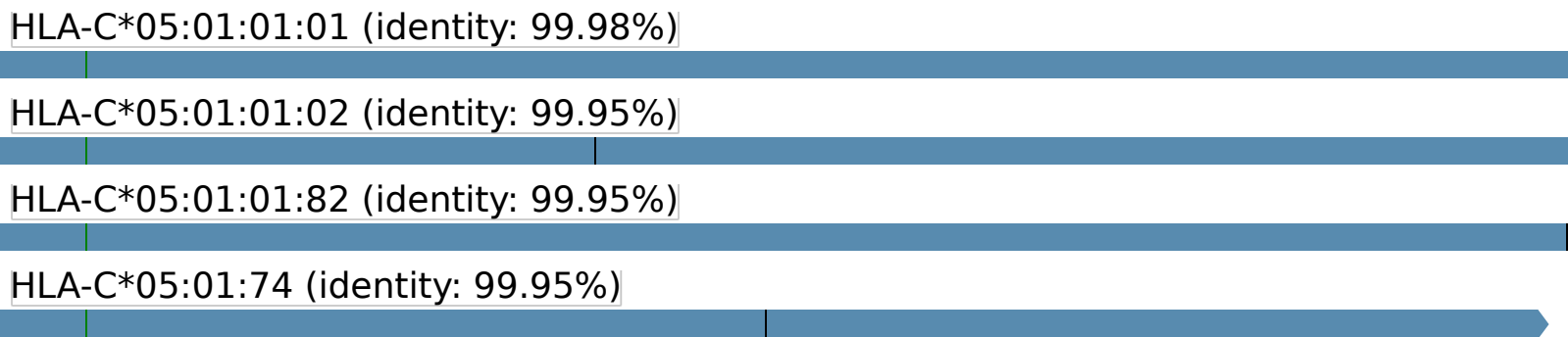
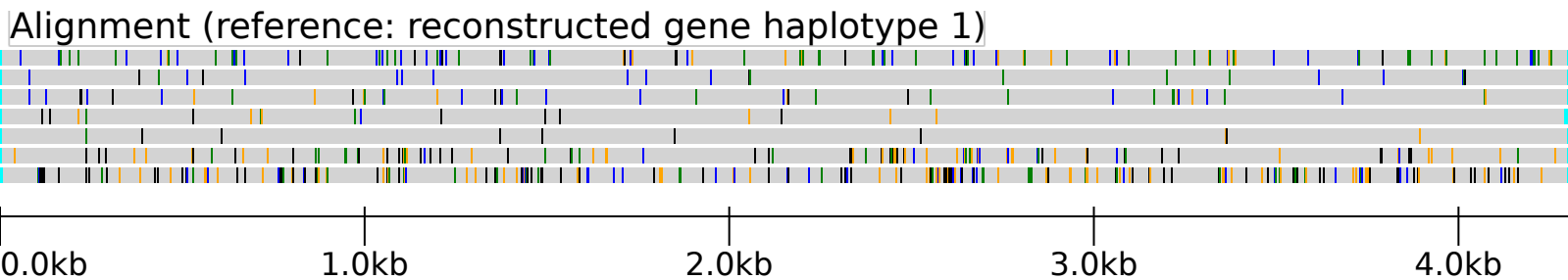
**Full typing result:**

HLA-C\*05:01:01:01

HLA-C\*05:01:01:02

HLA-C\*05:01:01:82

HLA-C\*05:01:74



Sample: HG01257

Locus: HLA-DMA

Resolution: 4th field

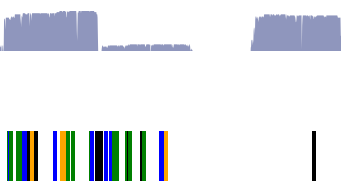
Full typing result:

HLA-DMA\*01:01:01:02

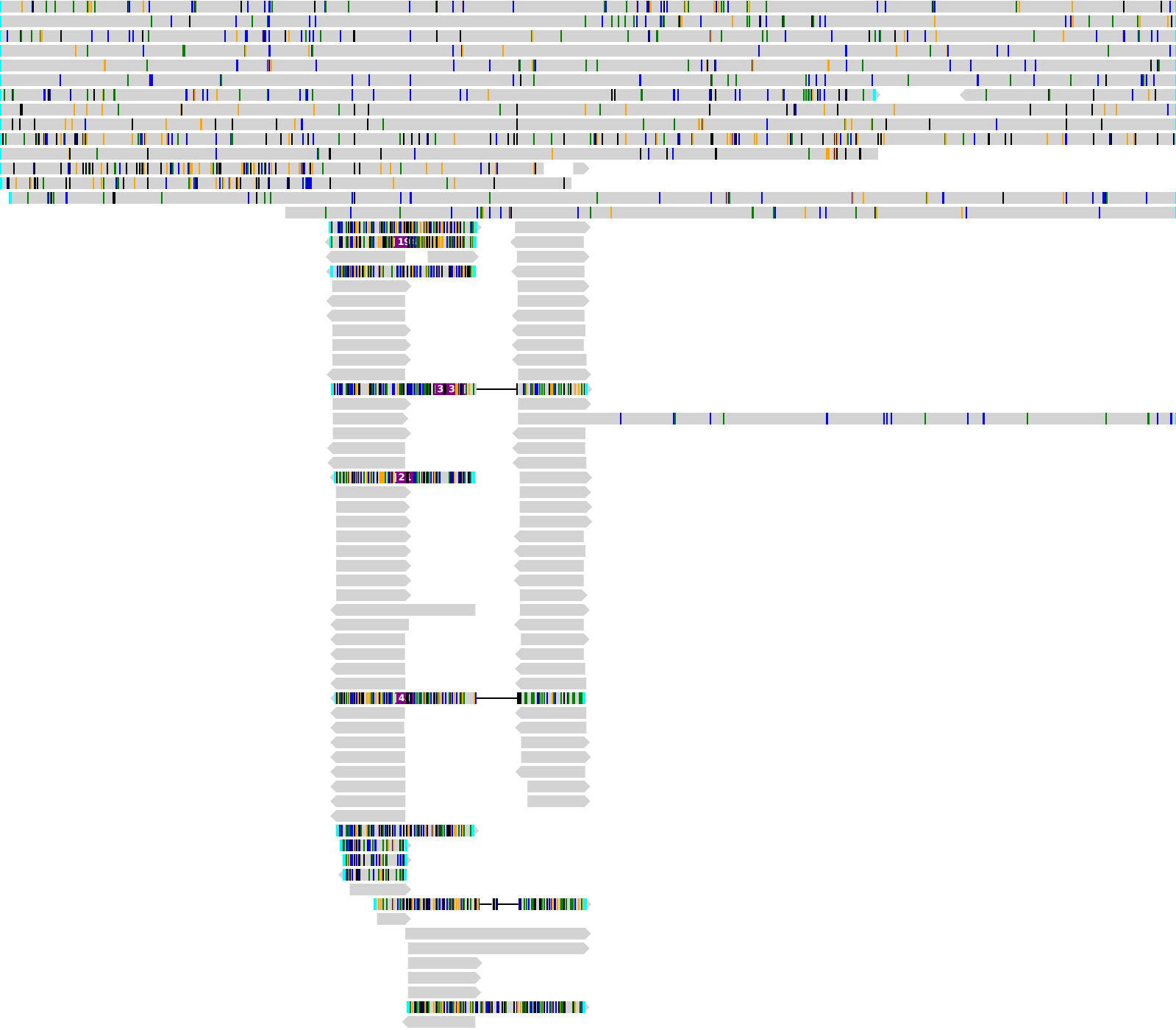
bam coverage

53  
35  
25

variants



Alignment (reference: reconstructed gene haplotype 1)



0.0kb 1.0kb 2.0kb 3.0kb 4.0kb 5.0kb

HLA-DMA\*01:01:01:02 (identity: 100.00%)



Sample: HG01257

Locus: HLA-DMB

Resolution: 4th field

Full typing result:

HLA-DMB\*01:03:01:17

HLA-DMB\*01:03:01:02

HLA-DMB\*01:03:01:18

HLA-DMB\*01:03:01:04

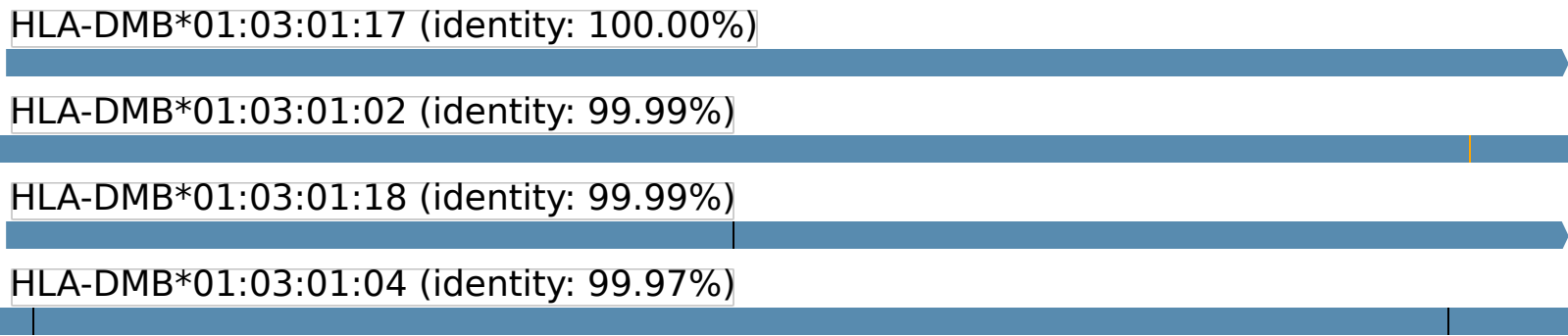
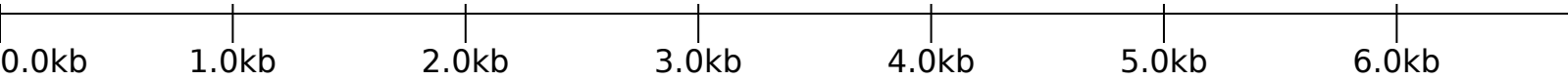
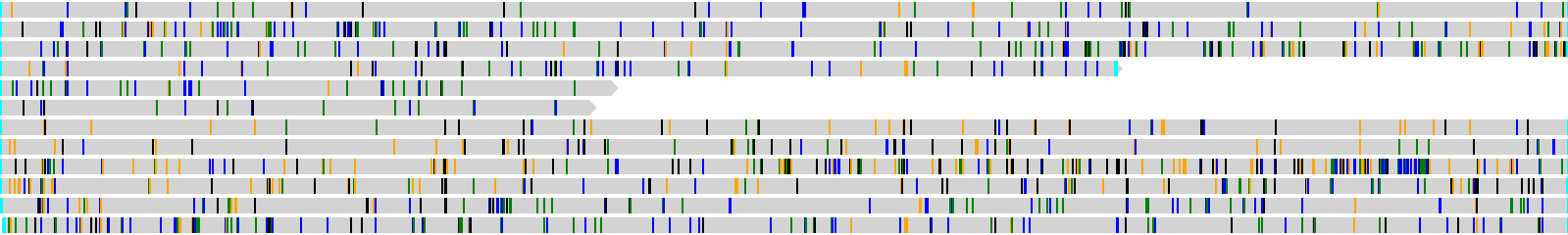
bam coverage



variants



Alignment (reference: reconstructed gene haplotype 1)



**Sample: HG01257**

**Locus: HLA-DOA**

**Resolution: 4th field**

**Full typing result:**

**HLA-DOA\*01:01:04:01**

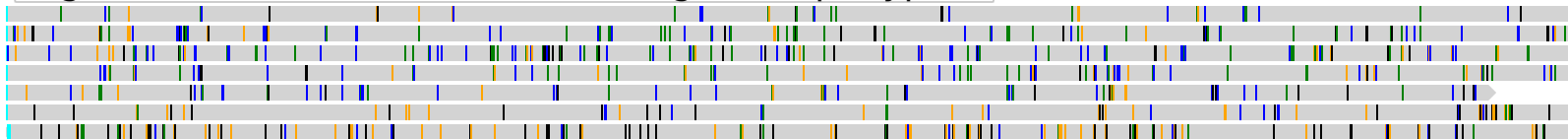
bam coverage



variants



Alignment (reference: reconstructed gene haplotype 1)



0.0kb

1.0kb

2.0kb

3.0kb

**HLA-DOA\*01:01:04:01 (identity: 100.00%)**



**Sample: HG01257**

**Locus: HLA-DOA**

**Resolution: 4th field**

**Full typing result:**

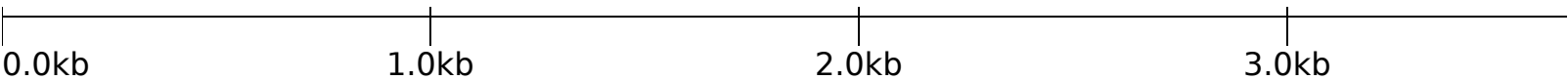
**HLA-DOA\*01:01:02:03**

bam coverage



variants

Alignment (reference: reconstructed gene haplotype 2)



**HLA-DOA\*01:01:02:03 (identity: 100.00%)**



**Sample: HG01257**

**Locus: HLA-DOB**

**Resolution: 4th field**

**Full typing result:**

**HLA-DOB\*01:03:01:02**

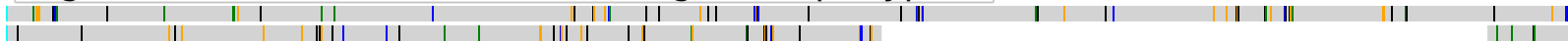
bam coverage



variants



Alignment (reference: reconstructed gene haplotype 2)



0.0kb

1.0kb

2.0kb

3.0kb

4.0kb

**HLA-DOB\*01:03:01:02 (identity: 100.00%)**



**Sample: HG01257**

**Locus: HLA-DOB**

**Resolution: 4th field**

**Full typing result:**

**HLA-DOB\*01:01:03:09**

**HLA-DOB\*01:01:03:02**

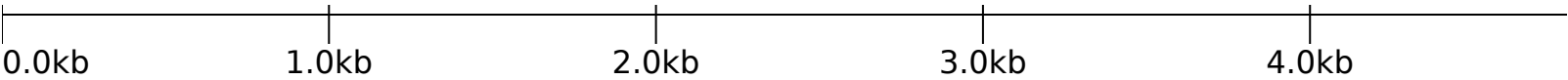
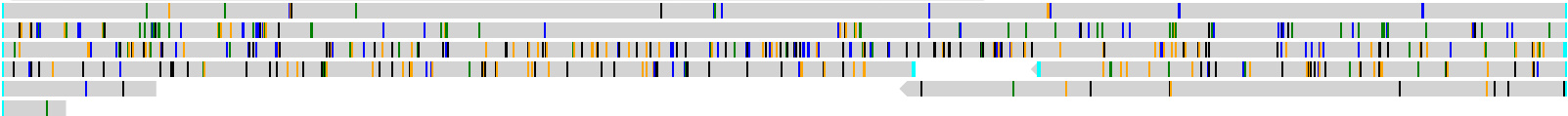
**HLA-DOB\*01:01:01:28**

bam coverage



variants

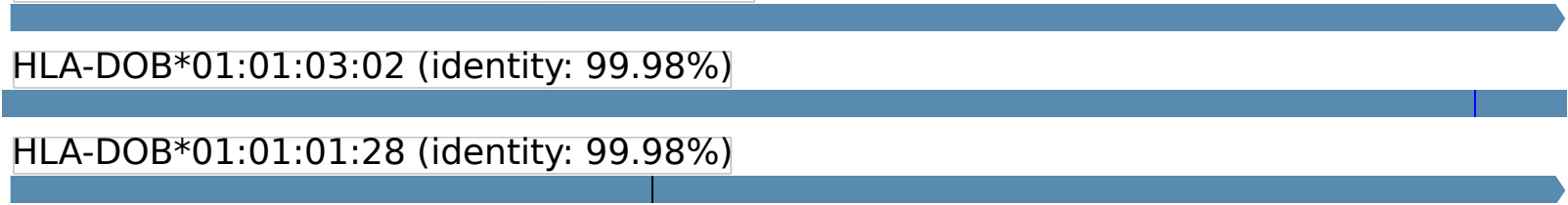
Alignment (reference: reconstructed gene haplotype 1)



**HLA-DOB\*01:01:03:09 (identity: 100.00%)**

**HLA-DOB\*01:01:03:02 (identity: 99.98%)**

**HLA-DOB\*01:01:01:28 (identity: 99.98%)**



**Sample: HG01257**

**Locus: HLA-DPA1**

**Resolution: 4th field**

**Full typing result:**

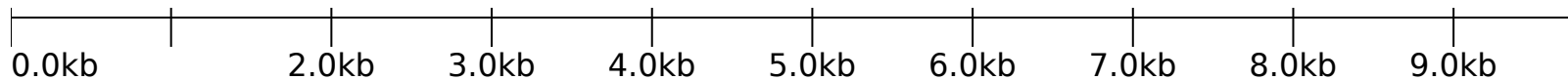
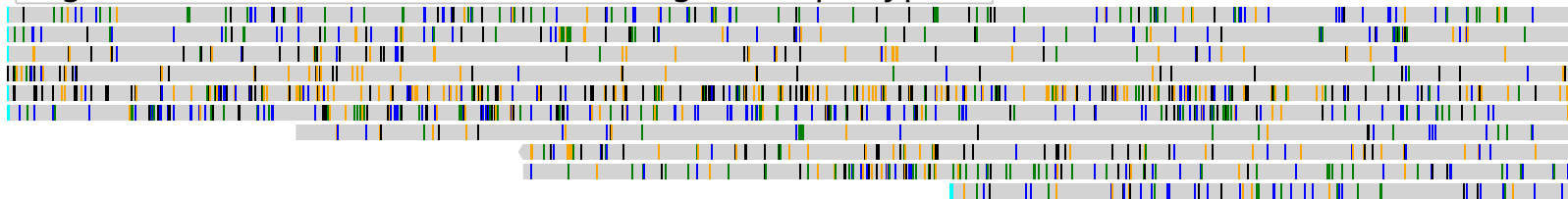
**HLA-DPA1\*01:03:01:01**

bam coverage



variants

Alignment (reference: reconstructed gene haplotype 1)



HLA-DPA1\*01:03:01:01 (identity: 100.00%)

**Sample: HG01257**

**Locus: HLA-DPA1**

**Resolution: 4th field**

**Full typing result:**

**HLA-DPA1\*01:03:01:05**

**HLA-DPA1\*01:03:01:15**

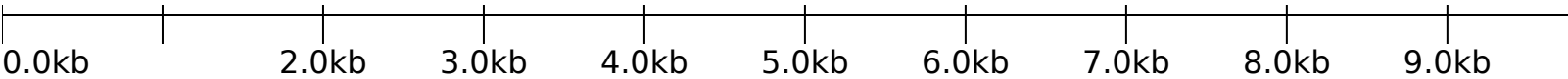
**HLA-DPA1\*01:03:01:74**

bam coverage



variants

Alignment (reference: reconstructed gene haplotype 2)



HLA-DPA1\*01:03:01:05 (identity: 99.99%)



HLA-DPA1\*01:03:01:15 (identity: 99.98%)



HLA-DPA1\*01:03:01:74 (identity: 99.98%)



**Sample: HG01257**

**Locus: HLA-DPA2**

**Resolution: 4th field**

**Full typing result:**

**HLA-DPA2\*01:01:01:02**

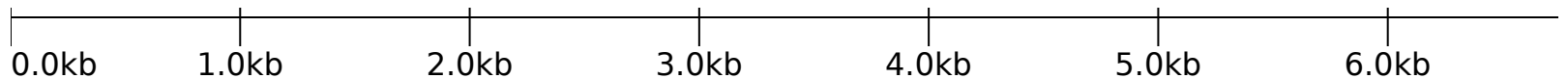
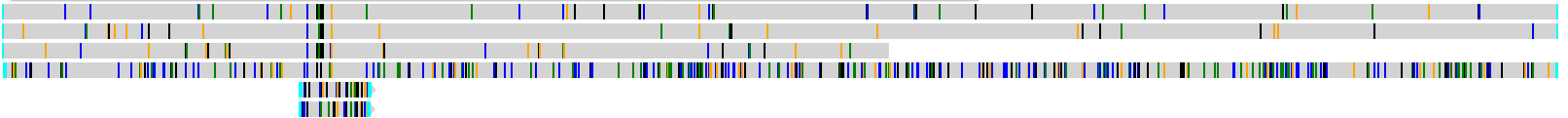
bam coverage



variants



Alignment (reference: reconstructed gene haplotype 2)



**HLA-DPA2\*01:01:01:02 (identity: 99.91%)**



**Sample: HG01257**

**Locus: HLA-DPA2**

**Resolution: 4th field**

**Full typing result:**

**HLA-DPA2\*02:01:01:01**

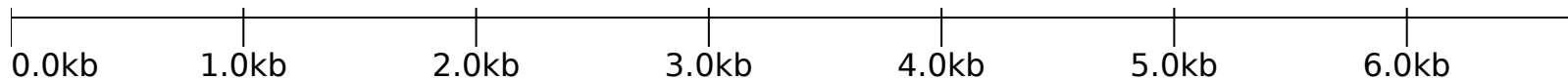
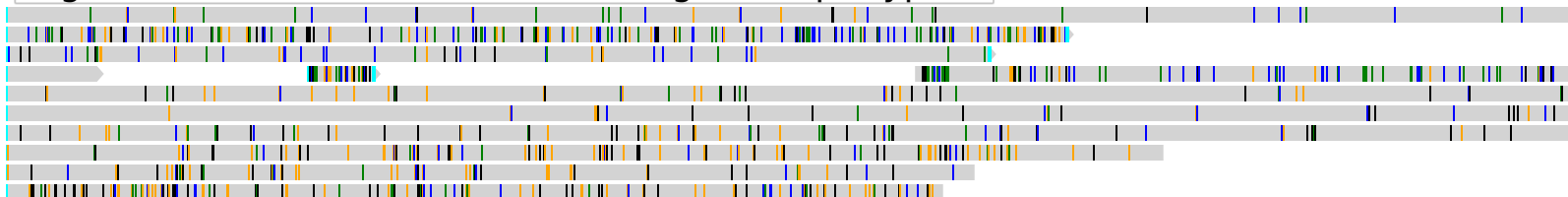
bam coverage



variants



Alignment (reference: reconstructed gene haplotype 1)



**HLA-DPA2\*02:01:01:01 (identity: 100.00%)**

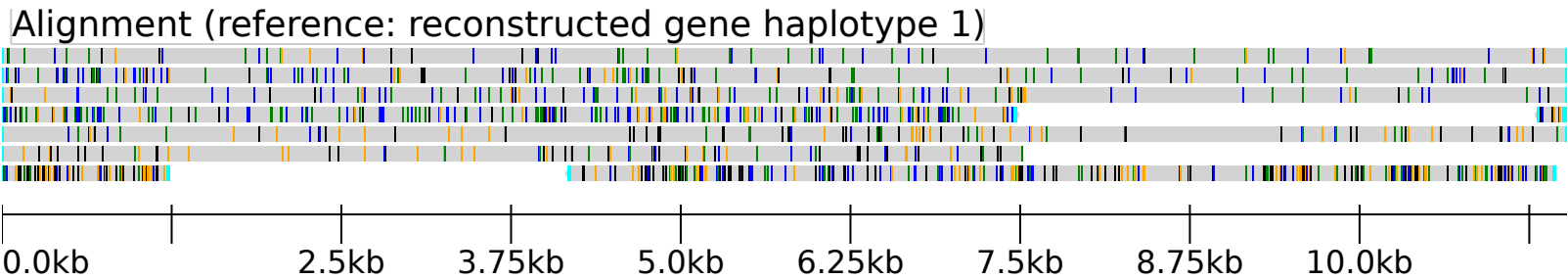
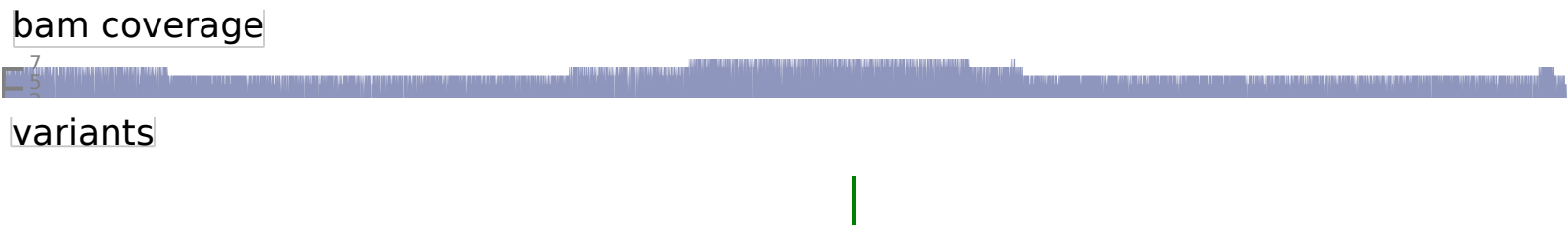
Sample: HG01257

Locus: HLA-DPB1

Resolution: 4th field

Full typing result:

- HLA-DPB1\*02:02:01:01
- HLA-DPB1\*02:02:01:05
- HLA-DPB1\*02:02:01:04
- HLA-DPB1\*02:02:03
- HLA-DPB1\*1188:01
- HLA-DPB1\*1376:01
- HLA-DPB1\*1403:01
- HLA-DPB1\*02:02:01:09
- HLA-DPB1\*47:01:01:01
- HLA-DPB1\*48:01
- HLA-DPB1\*02:01:02:04
- HLA-DPB1\*112:01:01:02
- HLA-DPB1\*02:01:02:05
- HLA-DPB1\*02:01:02:01
- HLA-DPB1\*71:01:01



- HLA-DPB1\*02:02:01:01 (identity: 100.00%)
- HLA-DPB1\*02:02:01:05 (identity: 99.97%)
- HLA-DPB1\*02:02:01:04 (identity: 99.97%)
- HLA-DPB1\*02:02:03 (identity: 99.99%)
- HLA-DPB1\*1188:01 (identity: 99.99%)
- HLA-DPB1\*1376:01 (identity: 99.99%)
- HLA-DPB1\*1403:01 (identity: 99.99%)
- HLA-DPB1\*02:02:01:09 (identity: 99.96%)
- HLA-DPB1\*47:01:01:01 (identity: 99.96%)
- HLA-DPB1\*48:01 (identity: 99.95%)



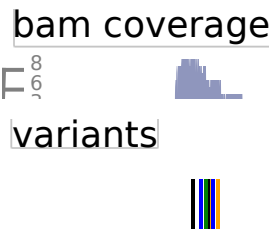
Sample: HG01257

Locus: HLA-DPB1

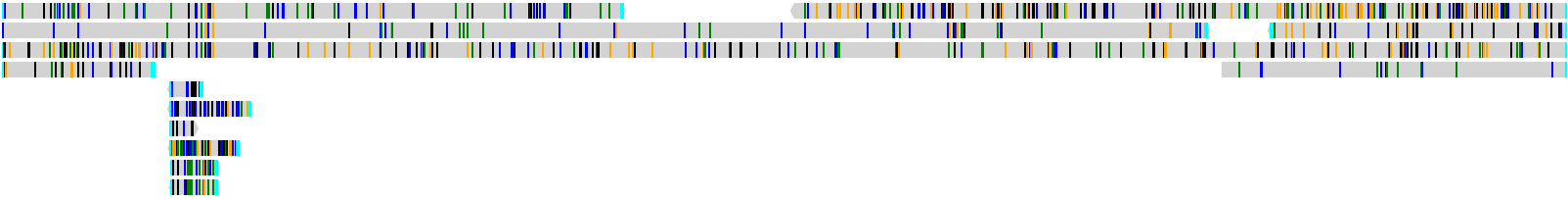
Resolution: 4th field

Full typing result:

HLA-DPB1*04:02:01:37	HLA-DPB1*04:02:01:02	HLA-DPB1*04:02:01:03
HLA-DPB1*04:02:01:04	HLA-DPB1*04:02:01:05	HLA-DPB1*04:02:01:07
HLA-DPB1*04:02:01:09	HLA-DPB1*04:02:01:11	HLA-DPB1*04:02:01:14
HLA-DPB1*04:02:01:15	HLA-DPB1*04:02:01:18	HLA-DPB1*04:02:01:21
HLA-DPB1*04:02:01:22	HLA-DPB1*04:02:01:24	HLA-DPB1*04:02:01:25



Alignment (reference: reconstructed gene haplotype 2)



HLA-DPB1*04:02:01:37 (identity: 99.93%)	
HLA-DPB1*04:02:01:02 (identity: 99.92%)	
HLA-DPB1*04:02:01:03 (identity: 99.91%)	
HLA-DPB1*04:02:01:04 (identity: 99.91%)	
HLA-DPB1*04:02:01:05 (identity: 99.91%)	
HLA-DPB1*04:02:01:07 (identity: 99.91%)	
HLA-DPB1*04:02:01:09 (identity: 99.91%)	
HLA-DPB1*04:02:01:11 (identity: 99.91%)	
HLA-DPB1*04:02:01:14 (identity: 99.91%)	
HLA-DPB1*04:02:01:15 (identity: 99.87%)	

**Sample: HG01257**

**Locus: HLA-DPB2**

**Resolution: 4th field**

**Full typing result:**

**HLA-DPB2\*03:01:01:02**

bam coverage

34  
23  
11

variants

Alignment (reference: reconstructed gene haplotype 2)



HLA-DPB2\*03:01:01:02 (identity: 99.96%)



Sample: HG01257

Locus: HLA-DPB2

Resolution: 4th field

Full typing result:

HLA-DPB2\*01:01:02

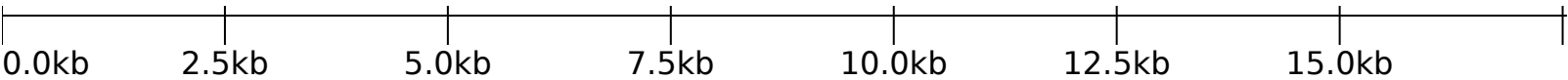
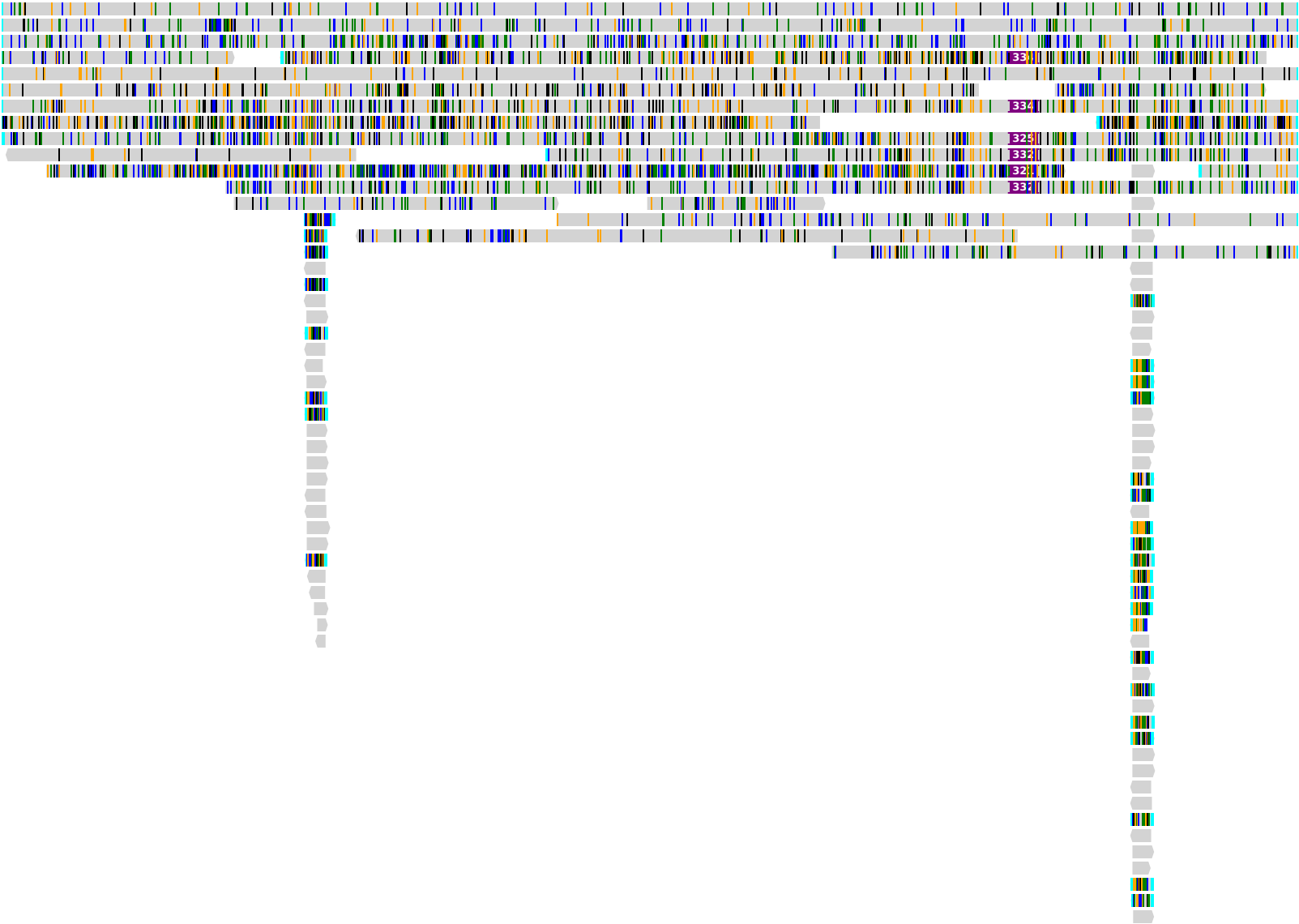
bam coverage



variants



Alignment (reference: reconstructed gene haplotype 1)



HLA-DPB2\*01:01:02 (identity: 99.99%)



**Sample: HG01257**

**Locus: HLA-DQA1**

**Resolution: 4th field**

**Full typing result:**

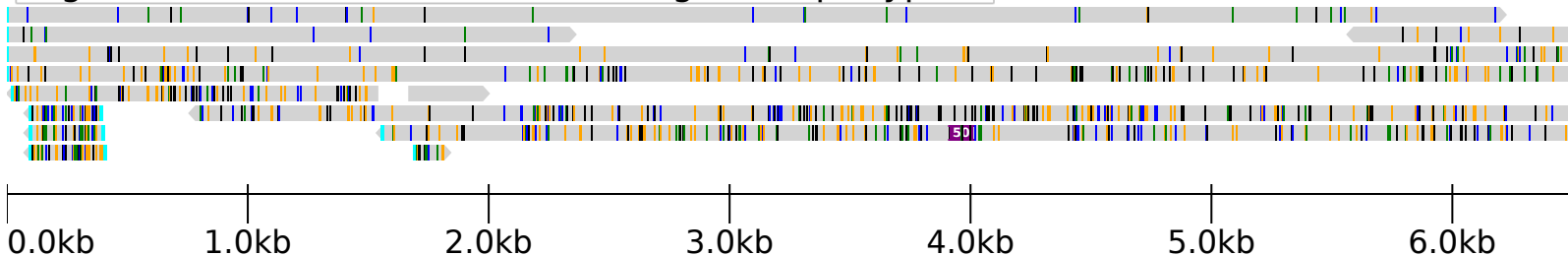
**HLA-DQA1\*05:01:01:01**

bam coverage



variants

Alignment (reference: reconstructed gene haplotype 2)



HLA-DQA1\*05:01:01:01 (identity: 99.98%)

**Sample: HG01257**

**Locus: HLA-DQA1**

**Resolution: 4th field**

**Full typing result:**

**HLA-DQA1\*01:05:01:01**

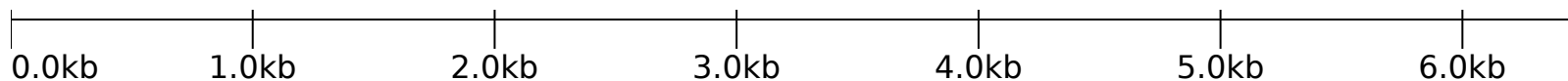
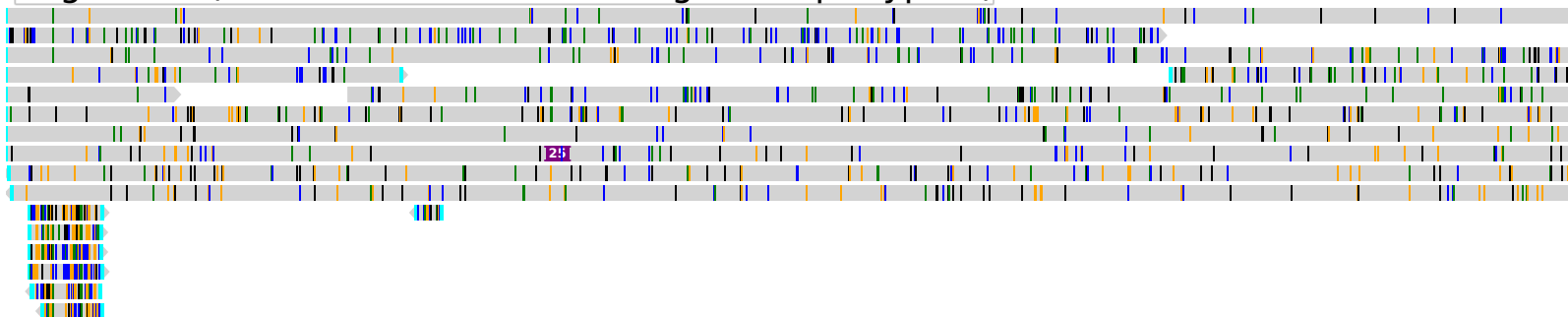
bam coverage



variants



Alignment (reference: reconstructed gene haplotype 1)



**HLA-DQA1\*01:05:01:01 (identity: 100.00%)**

**Sample: HG01257**

**Locus: HLA-DQA2**

**Resolution: 4th field**

**Full typing result:**

**HLA-DQA2\*01:04:01:02**

bam coverage

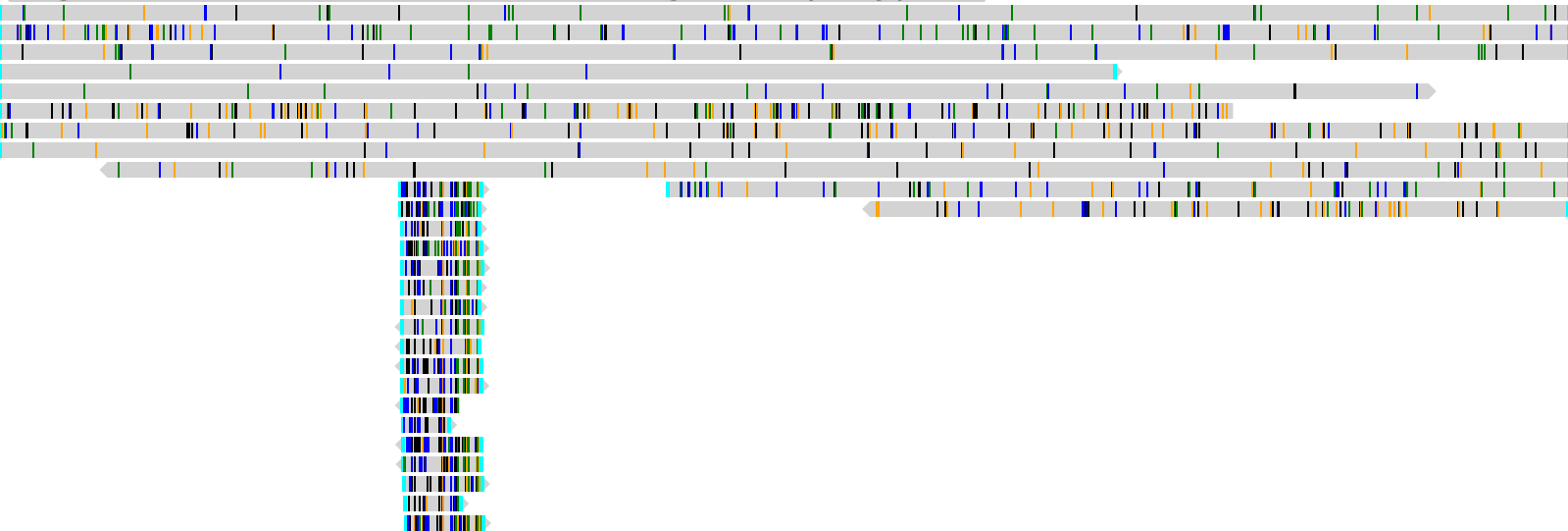
24  
16



variants



Alignment (reference: reconstructed gene haplotype 1)



0.0kb 1.0kb 2.0kb 3.0kb 4.0kb 5.0kb

HLA-DQA2\*01:04:01:02 (identity: 99.98%)



**Sample: HG01257**

**Locus: HLA-DQB1**

**Resolution: 4th field**

**Full typing result:**

**HLA-DQB1\*02:01:01:12**

bam coverage



variants



Alignment (reference: reconstructed gene haplotype 1)



0.0kb

10.0kb

20.0kb

30.0kb

HLA-DQB1\*02:01:01:12 (identity: 100.00%)



**Sample: HG01257**

**Locus: HLA-DQB1**

**Resolution: 4th field**

**Full typing result:**

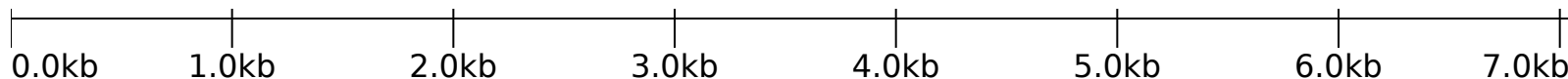
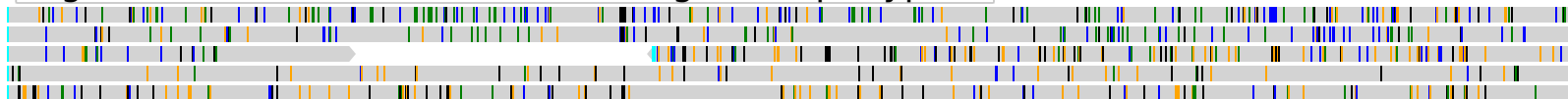
**HLA-DQB1\*05:01:01:05**

bam coverage



variants

Alignment (reference: reconstructed gene haplotype 2)



**HLA-DQB1\*05:01:01:05 (identity: 100.00%)**





Sample: HG01257

Locus: HLA-DRA

Resolution: 4th field

Full typing result:

HLA-DRA\*01:01:05

HLA-DRA\*01:01:01:01

HLA-DRA\*01:01:01:08

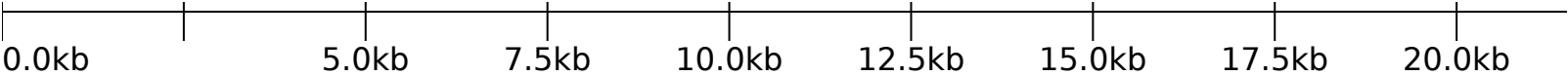
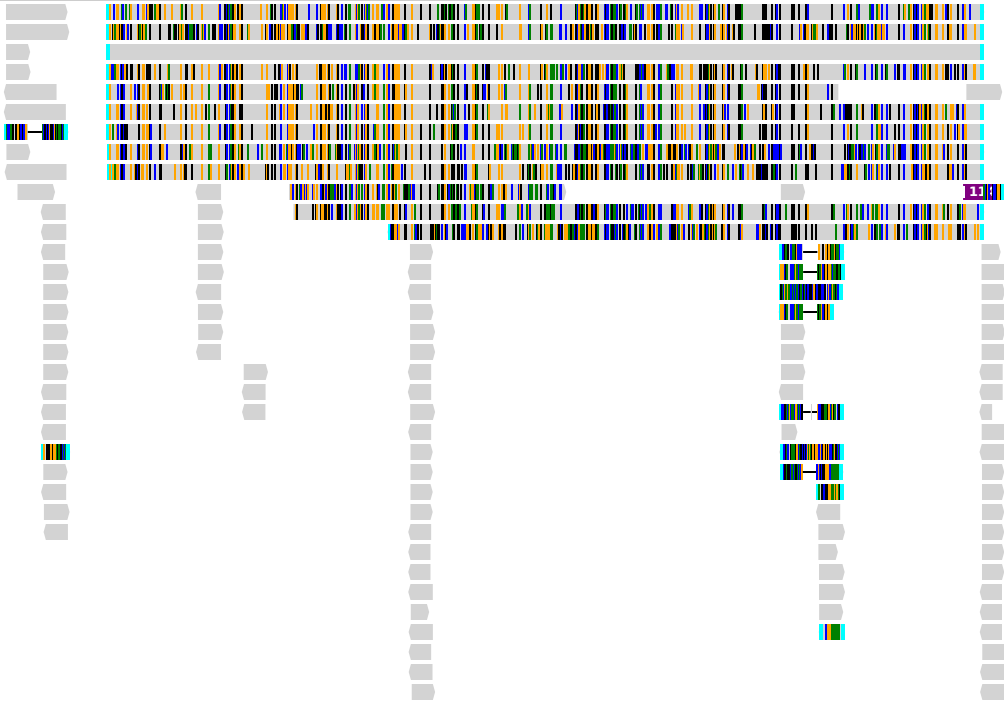
bam coverage



variants



Alignment (reference: reconstructed gene haplotype 1)



HLA-DRA\*01:01:05 (identity: 91.17%)



HLA-DRA\*01:01:01:01 (identity: 91.19%)



HLA-DRA\*01:01:01:08 (identity: 91.17%)



**Sample: HG01257**

**Locus: HLA-DRA**

**Resolution: 4th field**

**Full typing result:**

**HLA-DRA\*01:01:01:11**

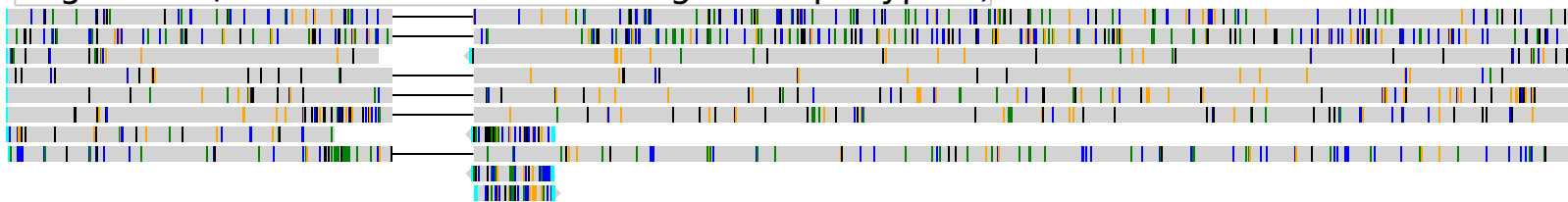
bam coverage



variants



Alignment (reference: reconstructed gene haplotype 2)



0.0kb 1.0kb 2.0kb 3.0kb 4.0kb 5.0kb 6.0kb

HLA-DRA\*01:01:01:11 (identity: 94.59%)



Sample: HG01257

Locus: HLA-DRB1

Resolution: 4th field

Full typing result:

- HLA-DRB1\*03:01:01:03
- HLA-DRB1\*03:01:01:02
- HLA-DRB1\*03:01:01:01
- HLA-DRB1\*03:01:01:06
- HLA-DRB1\*03:01:01:13
- HLA-DRB1\*03:01:01:21
- HLA-DRB1\*03:01:01:14
- HLA-DRB1\*03:01:01:17
- HLA-DRB1\*03:01:01:05
- HLA-DRB1\*14:06:01

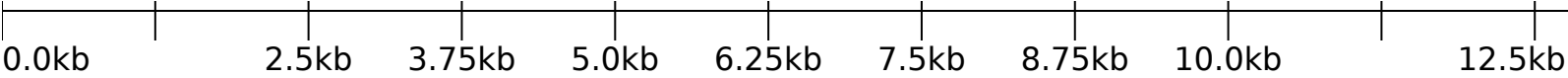
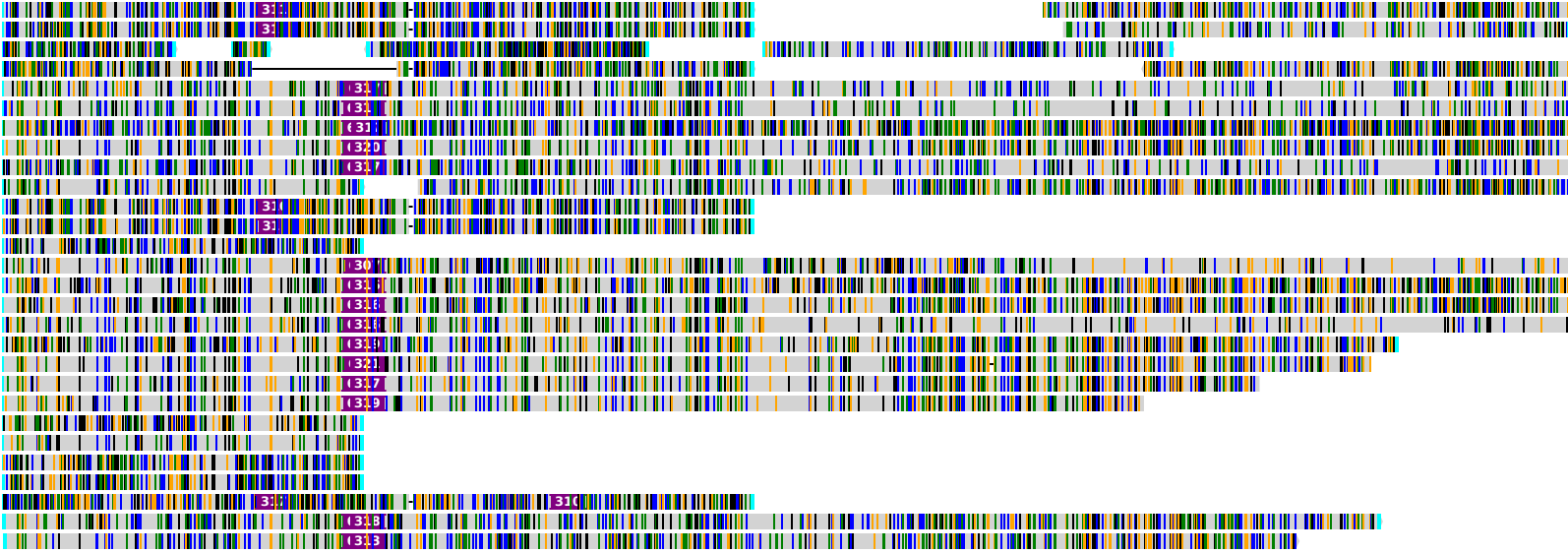
bam coverage



variants



Alignment (reference: reconstructed gene haplotype 2)



- HLA-DRB1\*03:01:01:03 (identity: 90.91%)
- HLA-DRB1\*03:01:01:02 (identity: 90.84%)
- HLA-DRB1\*03:01:01:01 (identity: 90.92%)
- HLA-DRB1\*03:01:01:06 (identity: 90.75%)
- HLA-DRB1\*03:01:01:13 (identity: 90.75%)
- HLA-DRB1\*03:01:01:21 (identity: 90.80%)
- HLA-DRB1\*03:01:01:14 (identity: 90.81%)
- HLA-DRB1\*03:01:01:17 (identity: 90.81%)
- HLA-DRB1\*03:01:01:05 (identity: 90.81%)
- HLA-DRB1\*14:06:01 (identity: 90.92%)

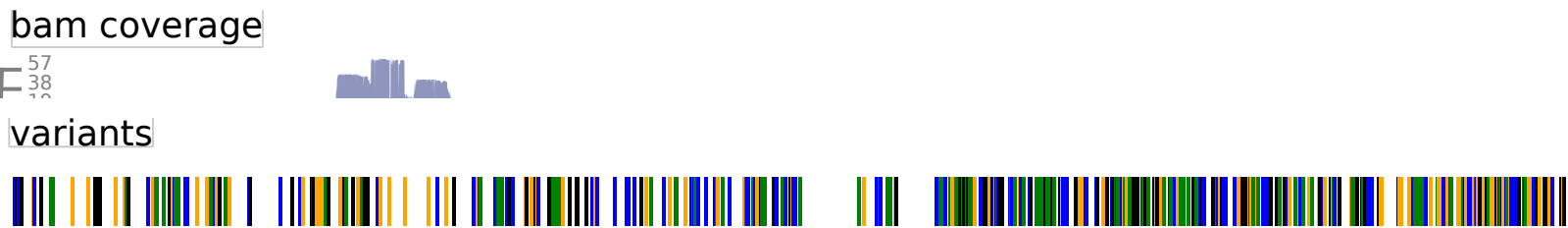
Sample: HG01257

Locus: HLA-DRB1

Resolution: 4th field

Full typing result:

HLA-DRB1\*10:01:01:03



Alignment (reference: reconstructed gene haplotype 1)



HLA-DRB1\*10:01:01:03 (identity: 99.99%)



**Sample: HG01257**

**Locus: HLA-DRB3**

**Resolution: 4th field**

**Full typing result:**

**HLA-DRB3\*02:02:01:02**

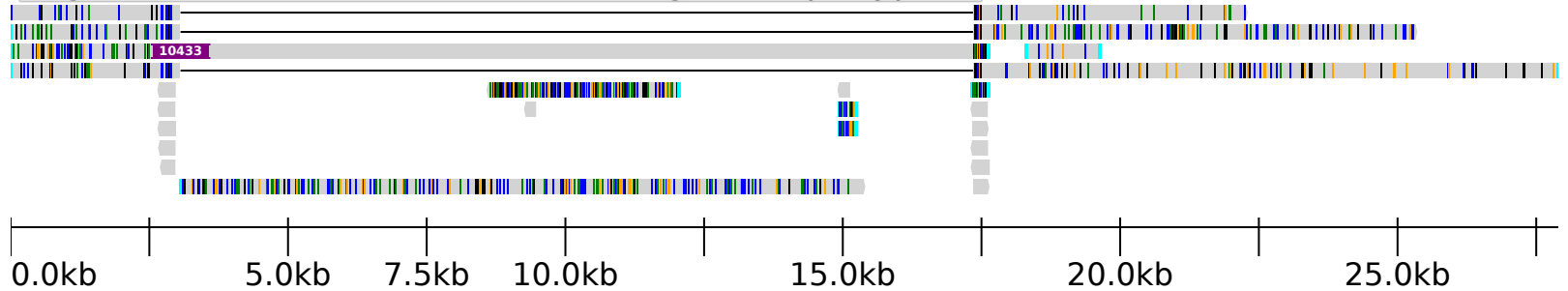
bam coverage

9  
6

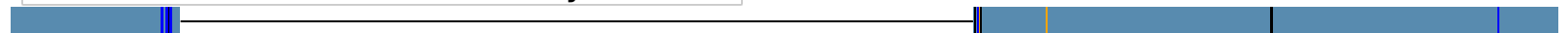
variants



Alignment (reference: reconstructed gene haplotype 1)



HLA-DRB3\*02:02:01:02 (identity: -5.07%)



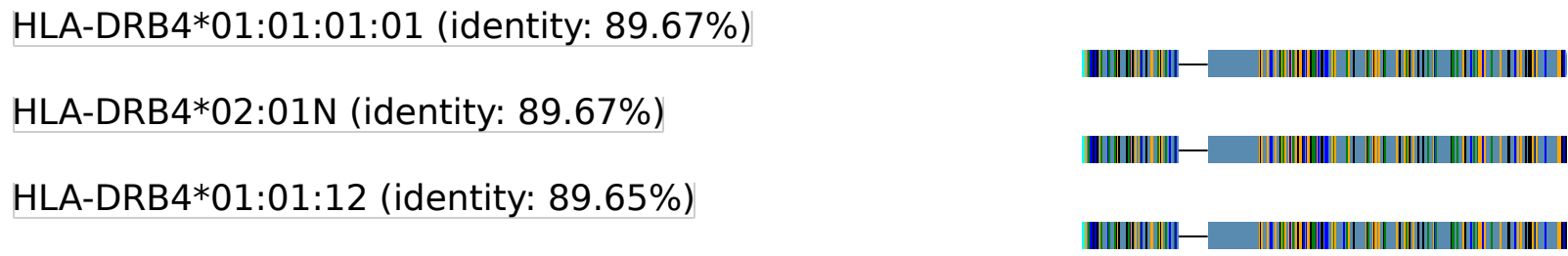
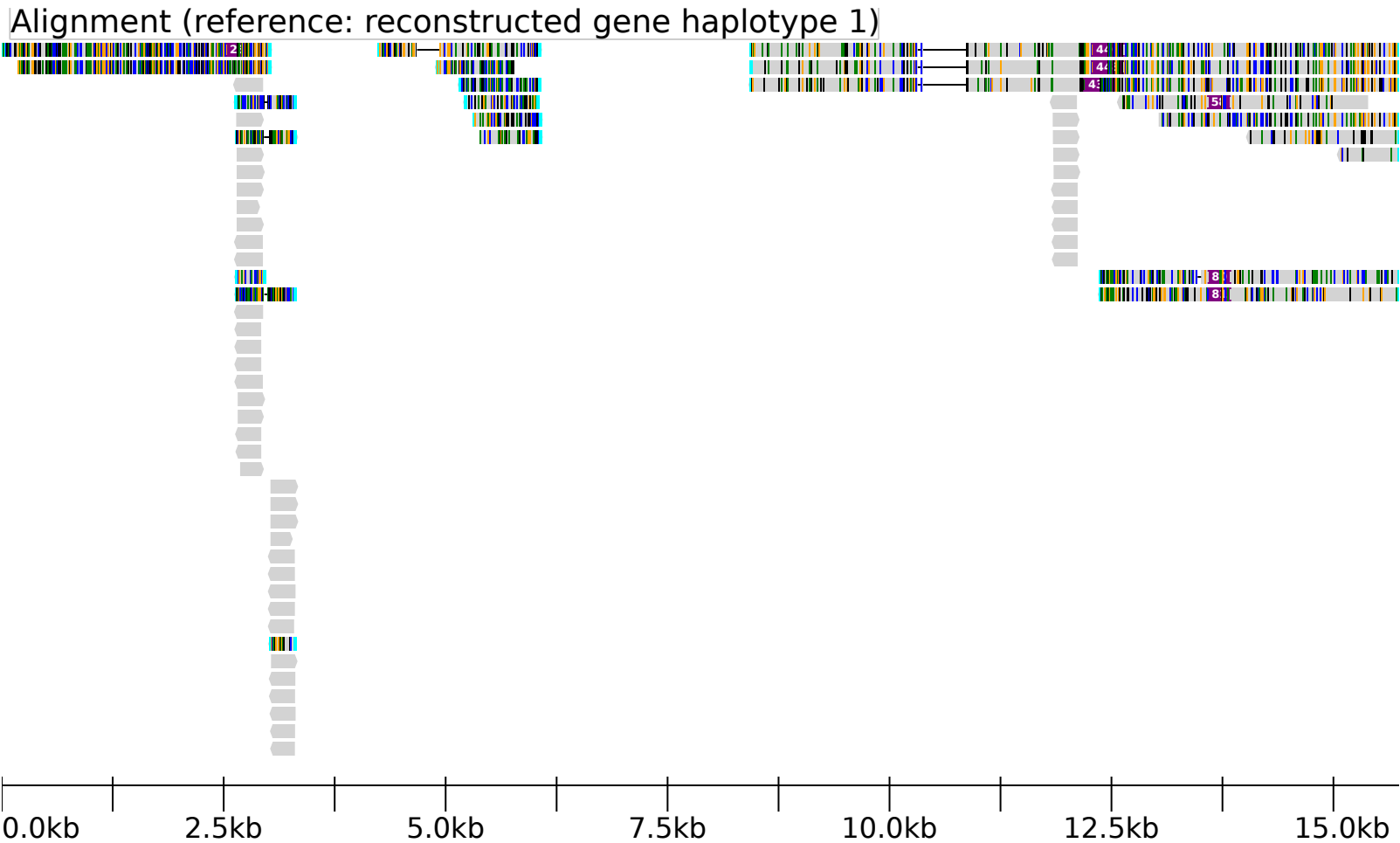
Sample: HG01257

Locus: HLA-DRB4

Resolution: 4th field

Full typing result:

HLA-DRB4\*01:01:01:01    HLA-DRB4\*02:01N    HLA-DRB4\*01:01:12



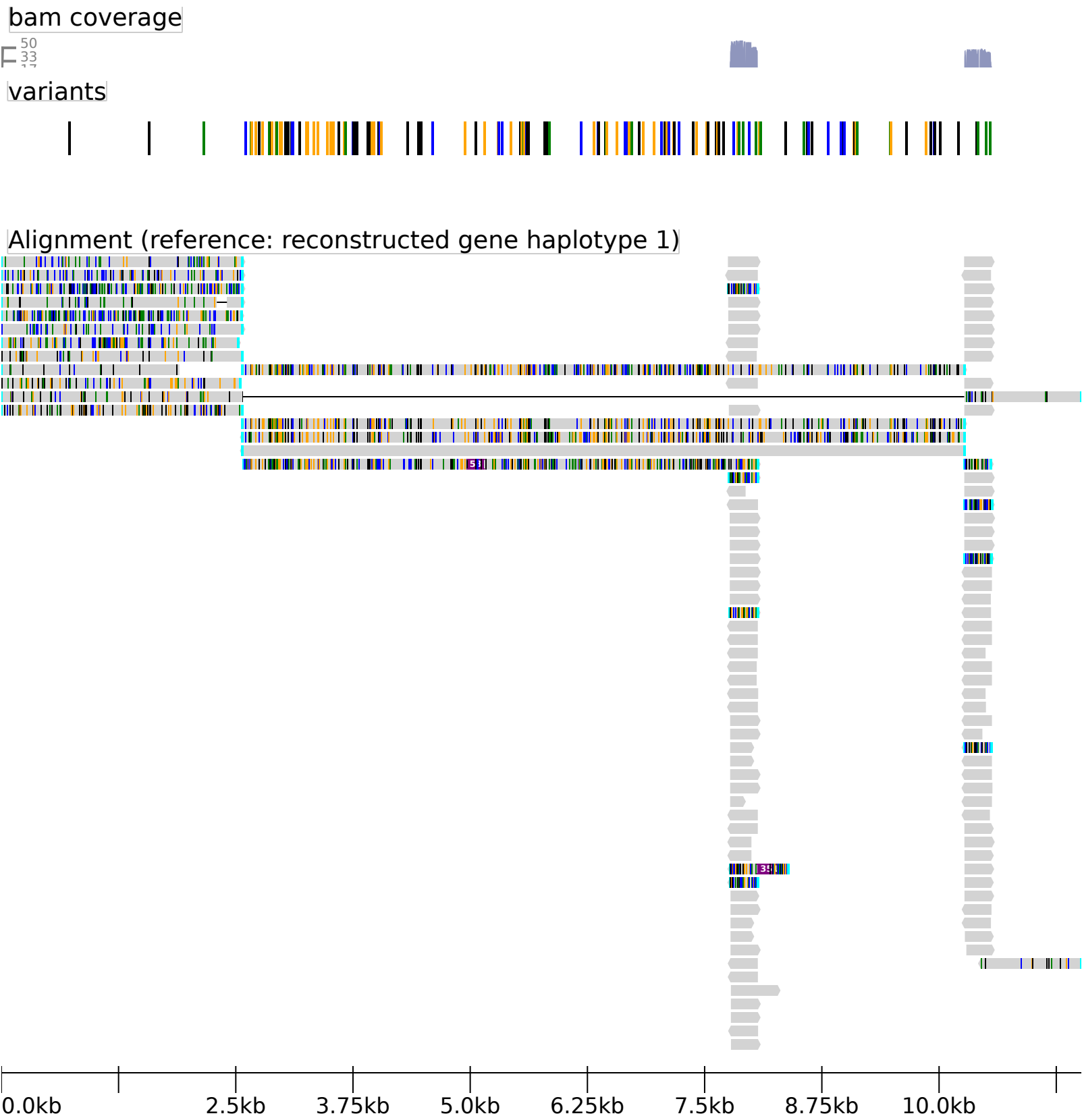
Sample: HG01257

Locus: HLA-E

Resolution: 4th field

Full typing result:

- HLA-E\*01:01:01:01HLA-E\*01:01:01:03HLA-E\*01:01:01:49
- HLA-E\*01:01:01:52HLA-E\*01:01:01:53HLA-E\*01:01:02
- HLA-E\*01:03:01:01HLA-E\*01:118HLA-E\*01:125
- HLA-E\*01:01:01:14HLA-E\*01:01:36HLA-E\*01:01:37
- HLA-E\*01:01:39HLA-E\*01:01:41HLA-E\*01:01:42



HLA-E*01:01:01:01 (identity: -101.60%)	
HLA-E*01:01:01:03 (identity: -101.62%)	
HLA-E*01:01:01:49 (identity: -101.62%)	
HLA-E*01:01:01:52 (identity: -101.62%)	
HLA-E*01:01:01:53 (identity: -101.62%)	
HLA-E*01:01:02 (identity: -101.62%)	
HLA-E*01:03:01:01 (identity: -101.62%)	
HLA-E*01:118 (identity: -101.62%)	
HLA-E*01:125 (identity: -101.62%)	
HLA-E*01:01:01:14 (identity: -103.38%)	

**Sample: HG01257**

**Locus: HLA-F**

**Resolution: 4th field**

**Full typing result:**

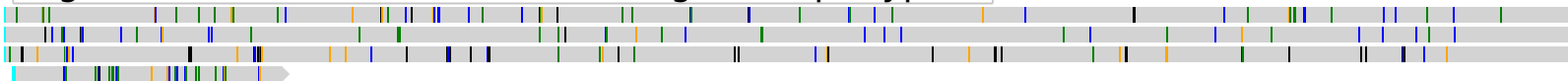
**HLA-F\*01:01:01:01**

bam coverage



variants

Alignment (reference: reconstructed gene haplotype 2)



0.0kb

1.0kb

2.0kb

3.0kb

**HLA-F\*01:01:01:01 (identity: 100.00%)**



**Sample: HG01257**

**Locus: HLA-F**

**Resolution: 4th field**

**Full typing result:**

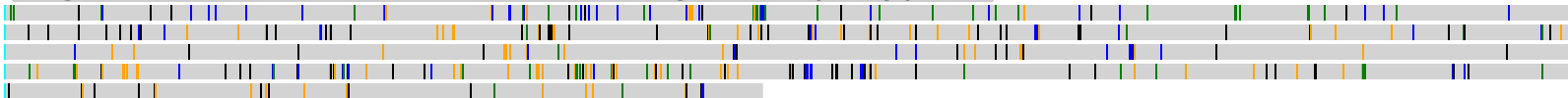
**HLA-F\*01:03:01:01**

bam coverage



variants

Alignment (reference: reconstructed gene haplotype 1)



0.0kb

1.0kb

2.0kb

3.0kb

**HLA-F\*01:03:01:01 (identity: 99.94%)**



**Sample: HG01257**

**Locus: HLA-G**

**Resolution: 4th field**

**Full typing result:**

**HLA-G\*01:01:01:05**

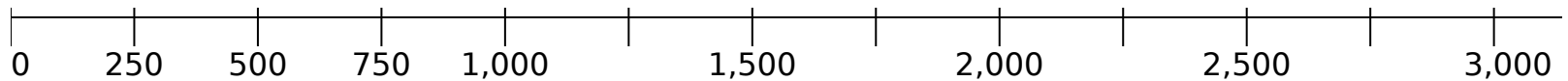
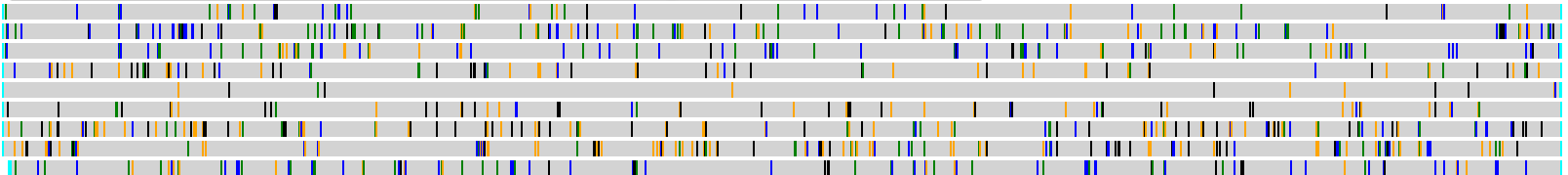
bam coverage



variants



Alignment (reference: reconstructed gene haplotype 1)



**HLA-G\*01:01:01:05 (identity: 100.00%)**



**Sample: HG01257**

**Locus: HLA-G**

**Resolution: 4th field**

**Full typing result:**

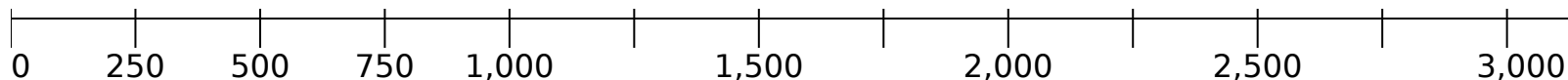
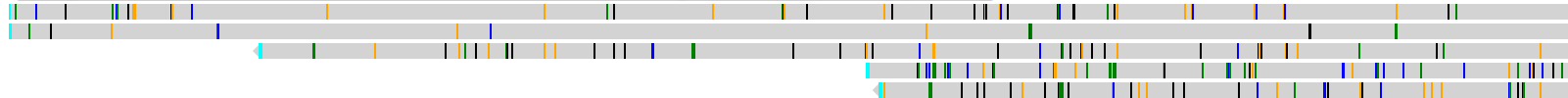
**HLA-G\*01:01:01:01**

bam coverage

5  
3

variants

Alignment (reference: reconstructed gene haplotype 2)



**HLA-G\*01:01:01:01 (identity: 100.00%)**

**Sample: HG01257**

**Locus: HLA-H**

**Resolution: 4th field**

**Full typing result:**

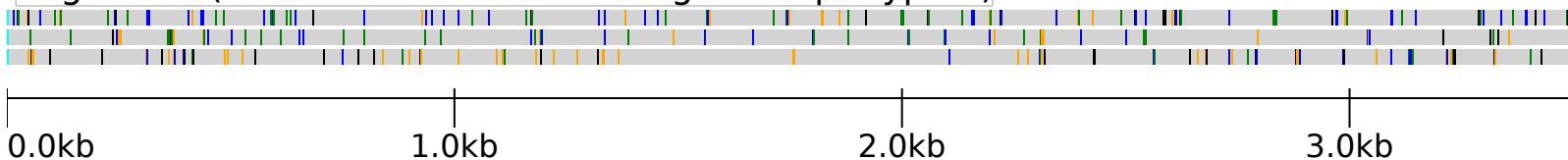
**HLA-H\*02:04:01**

bam coverage



variants

Alignment (reference: reconstructed gene haplotype 1)



HLA-H\*02:04:01 (identity: 100.00%)



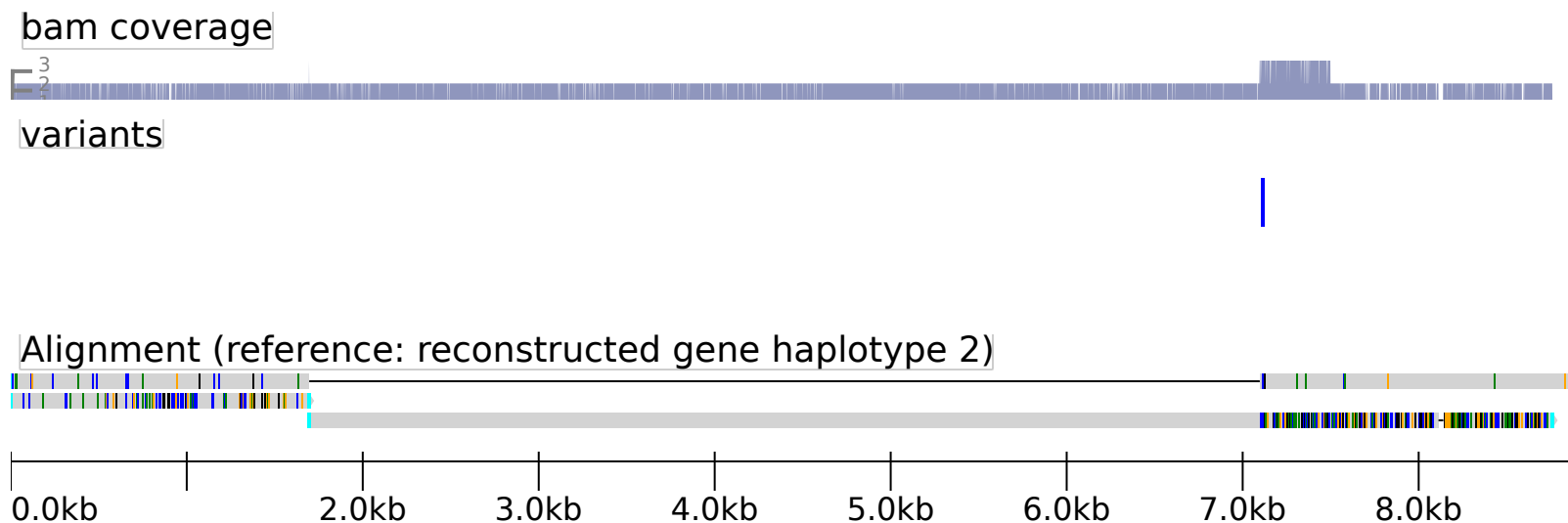
**Sample: HG01257**

**Locus: HLA-H**

**Resolution: 4th field**

**Full typing result:**

**HLA-H\*01:01:01:03**



**HLA-H\*01:01:01:03 (identity: -54.67%)**

**Sample: HG01257**

**Locus: HLA-J**

**Resolution: 4th field**

**Full typing result:**

**HLA-J\*01:01:01:06**

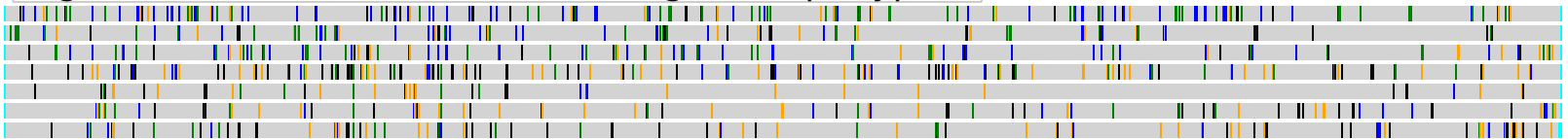
bam coverage



variants



Alignment (reference: reconstructed gene haplotype 2)



0.0kb

1.0kb

2.0kb

3.0kb

HLA-J\*01:01:01:06 (identity: 99.94%)



**Sample: HG01257**

**Locus: HLA-J**

**Resolution: 4th field**

**Full typing result:**

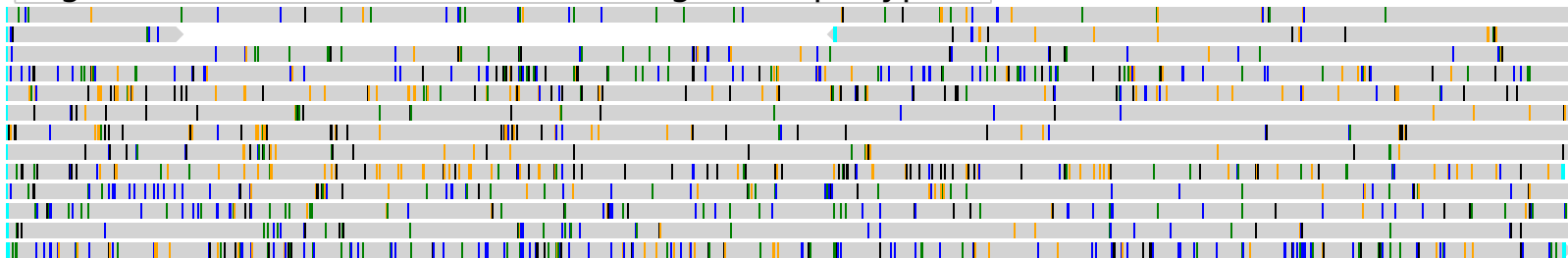
**HLA-J\*01:01:01:04**

bam coverage



variants

Alignment (reference: reconstructed gene haplotype 1)



0.0kb

1.0kb

2.0kb

3.0kb

HLA-J\*01:01:01:04 (identity: 100.00%)

**Sample: HG01257**

**Locus: HLA-K**

**Resolution: 4th field**

**Full typing result:**

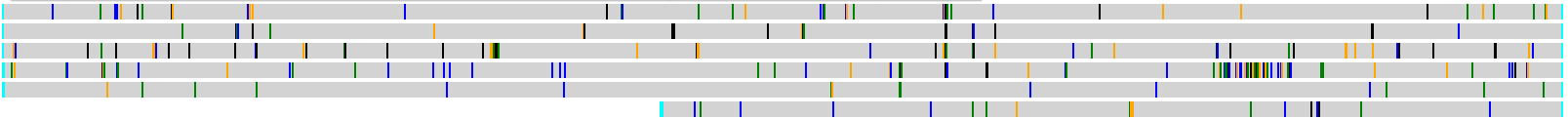
**HLA-K\*01:01:01:01**

bam coverage



variants

Alignment (reference: reconstructed gene haplotype 1)



0.0kb

1.0kb

2.0kb

3.0kb

HLA-K\*01:01:01:01 (identity: 100.00%)



**Sample: HG01257**

**Locus: HLA-L**

**Resolution: 4th field**

**Full typing result:**

**HLA-L\*01:01:02**

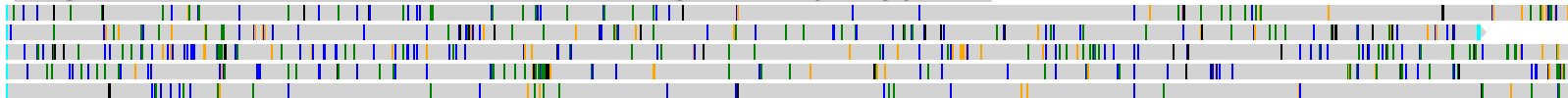
bam coverage



variants



Alignment (reference: reconstructed gene haplotype 2)



0.0kb

1.0kb

2.0kb

3.0kb

**HLA-L\*01:01:02 (identity: 99.95%)**



**Sample: HG01257**

**Locus: HLA-L**

**Resolution: 4th field**

**Full typing result:**

**HLA-L\*01:01:01:03**

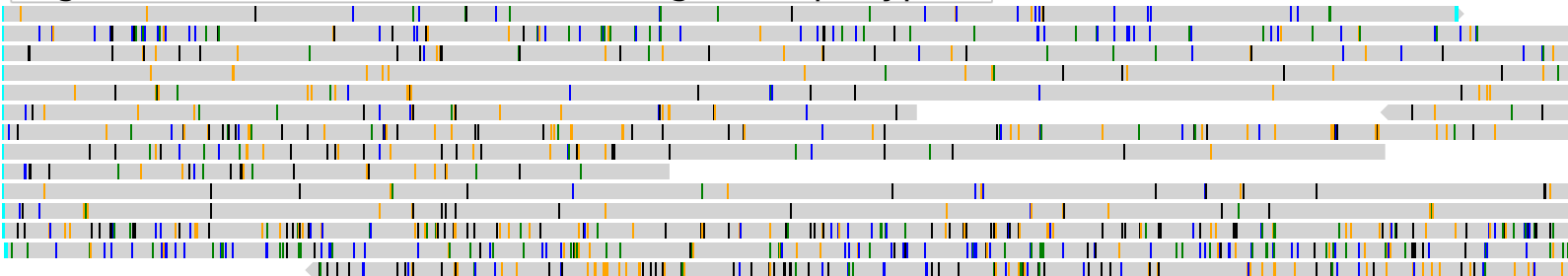
bam coverage



variants



Alignment (reference: reconstructed gene haplotype 1)



0.0kb

1.0kb

2.0kb

3.0kb

HLA-L\*01:01:01:03 (identity: 100.00%)

**Sample: HG01257**

**Locus: HLA-P**

**Resolution: 4th field**

**Full typing result:**

**HLA-P\*02:01:01:02**

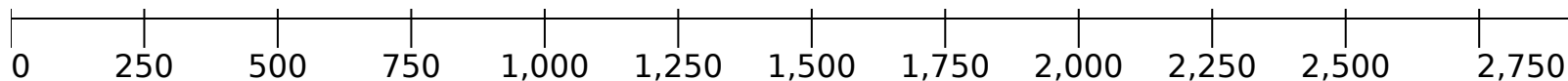
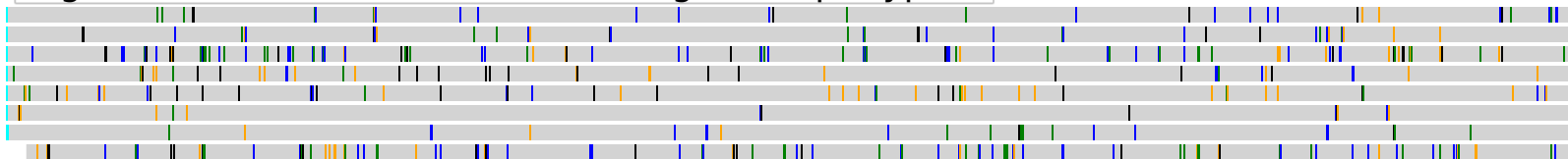
bam coverage



variants



Alignment (reference: reconstructed gene haplotype 1)



HLA-P\*02:01:01:02 (identity: 100.00%)



**Sample: HG01257**

**Locus: HLA-U**

**Resolution: 4th field**

**Full typing result:**

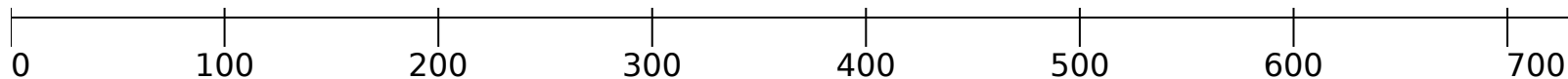
**HLA-U\*01:03**

bam coverage

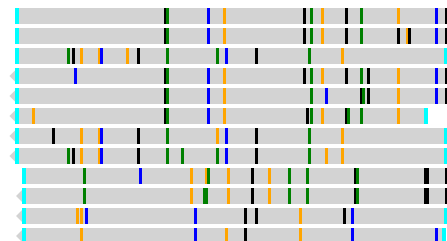
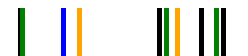
$\frac{11}{7}$

variants

Alignment (reference: reconstructed gene haplotype 1)



HLA-U\*01:03 (identity: 69.26%)



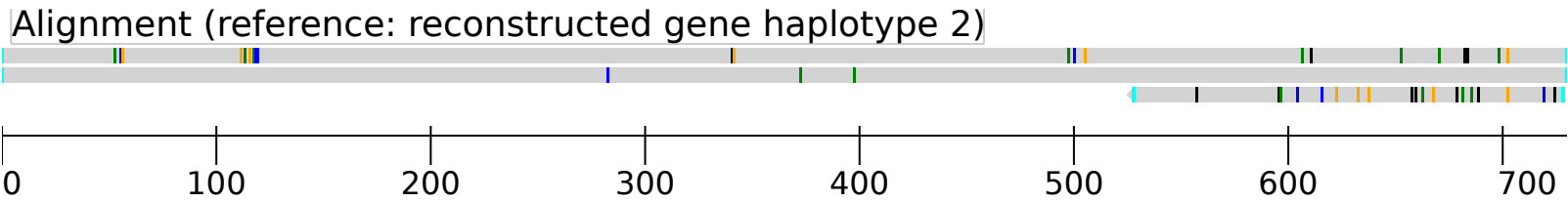
**Sample: HG01257**

**Locus: HLA-U**

**Resolution: 4th field**

**Full typing result:**

**HLA-U\*01:04**



HLA-U\*01:04 (identity: 100.00%)



A solid blue horizontal bar representing the HLA-U\*01:04 allele with 100.00% identity.

**Sample: HG01257**

**Locus: HLA-W**

**Resolution: 4th field**

**Full typing result:**

**HLA-W\*03:01:01:01**

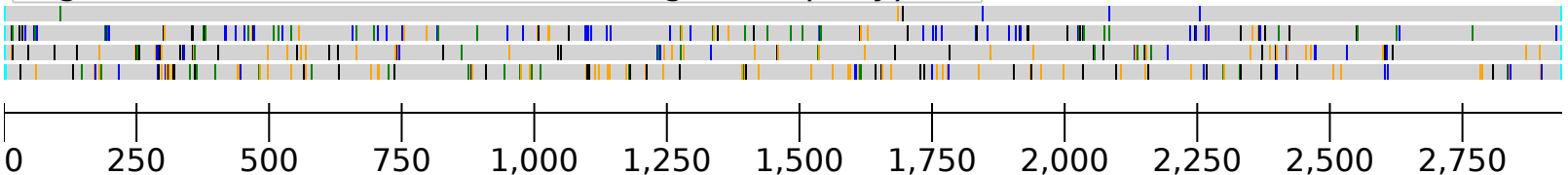
**HLA-W\*03:01:01:02**

bam coverage



variants

Alignment (reference: reconstructed gene haplotype 1)



HLA-W\*03:01:01:01 (identity: 100.00%)

HLA-W\*03:01:01:02 (identity: 99.97%)

Sample: HG01257

Locus: MICA

Resolution: 4th field

Full typing result:

MICA\*008:04:01

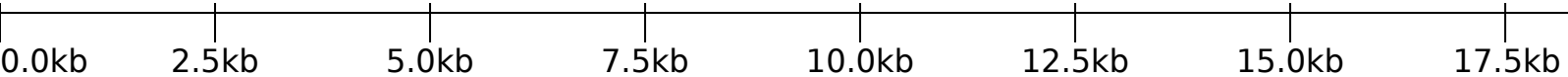
bam coverage



variants



Alignment (reference: reconstructed gene haplotype 1)



MICA\*008:04:01 (identity: 100.00%)



Sample: HG01257

Locus: MICA

Resolution: 4th field

Full typing result:

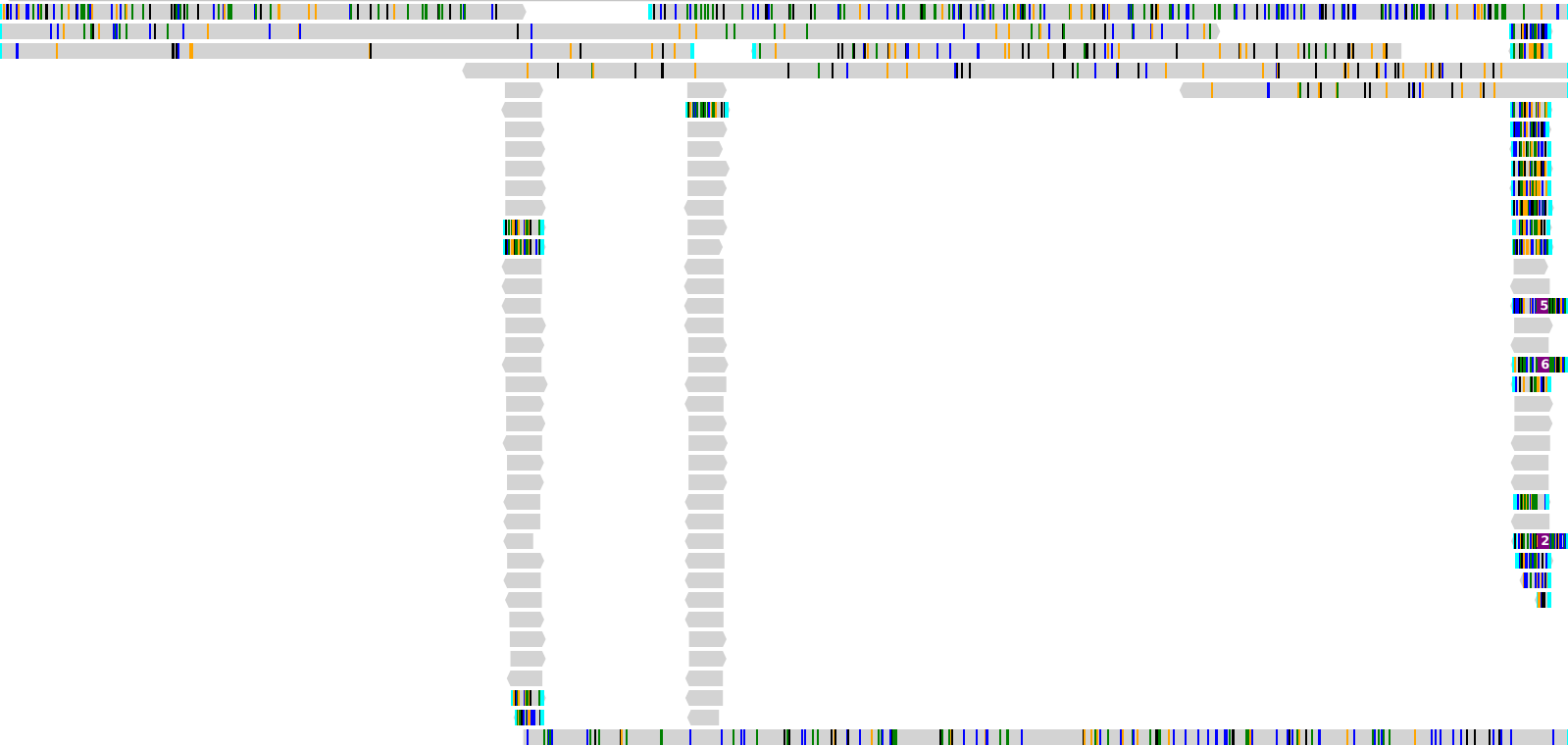
MICA\*001:01

bam coverage

33  
22  
11

variants

Alignment (reference: reconstructed gene haplotype 2)



MICA\*001:01 (identity: 99.99%)



Sample: HG01257

Locus: MICB

Resolution: 4th field

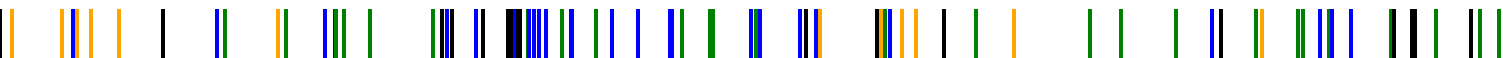
Full typing result:

MICB\*052

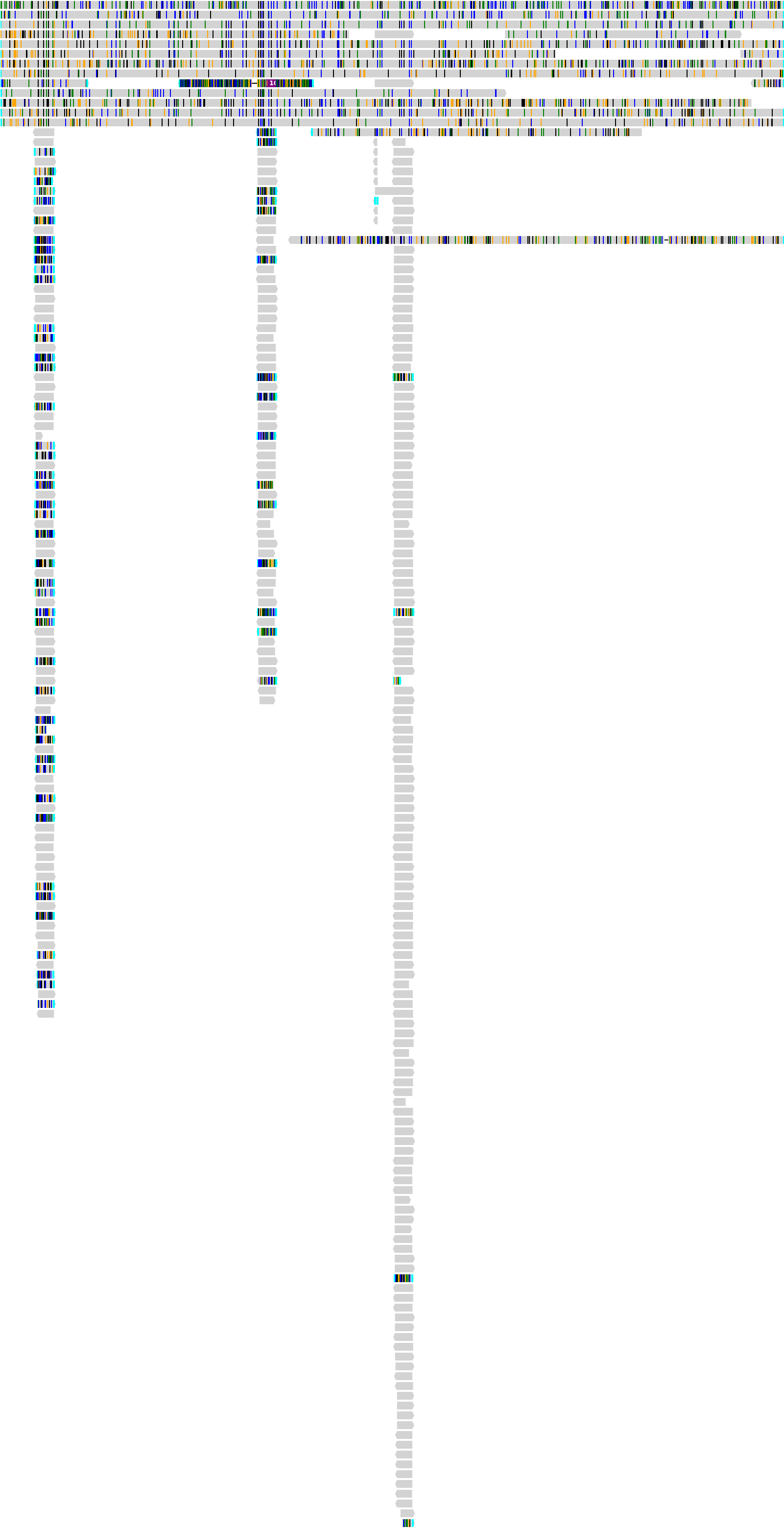
bam coverage



variants



Alignment (reference: reconstructed gene haplotype 1)



MICB\*052 (identity: 99.85%)



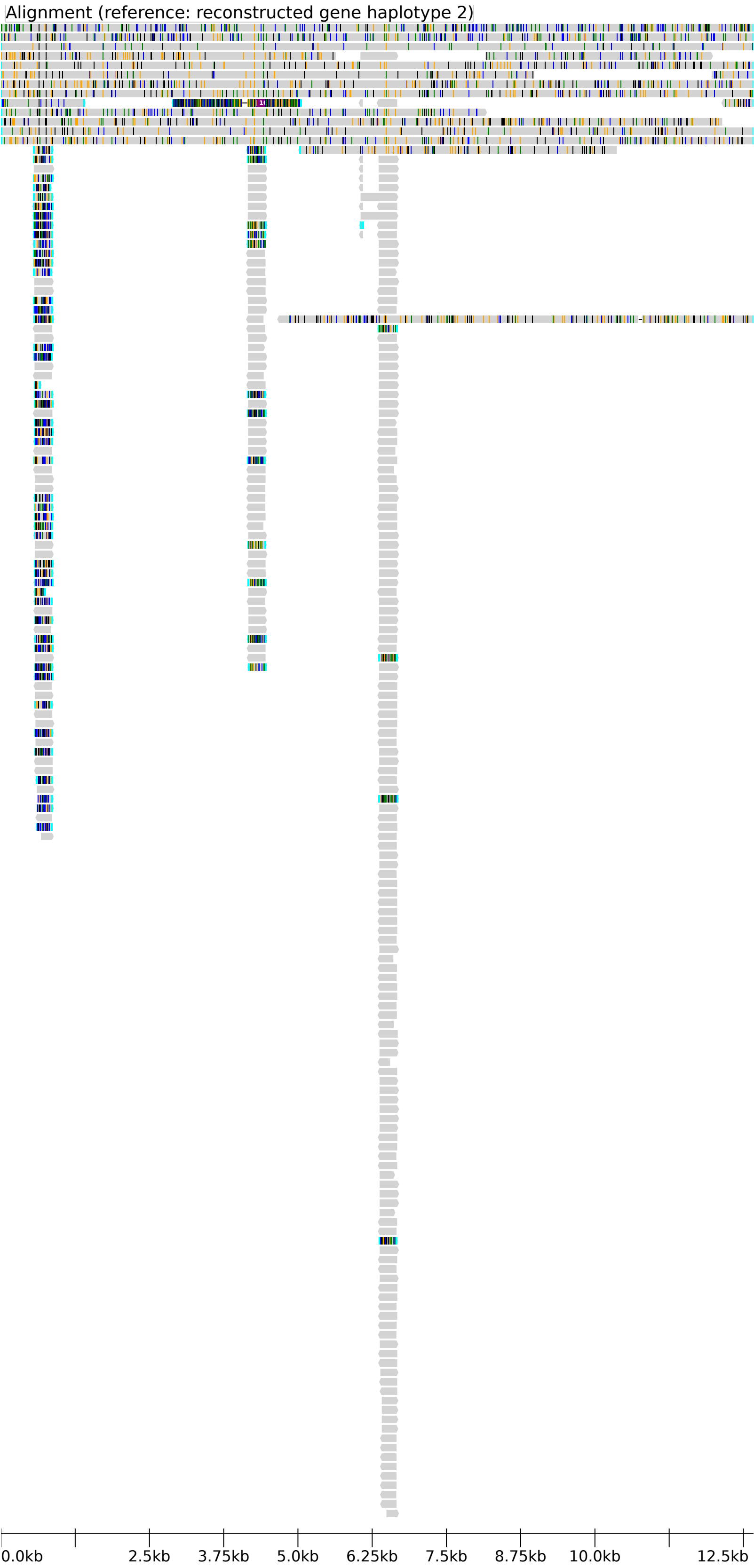
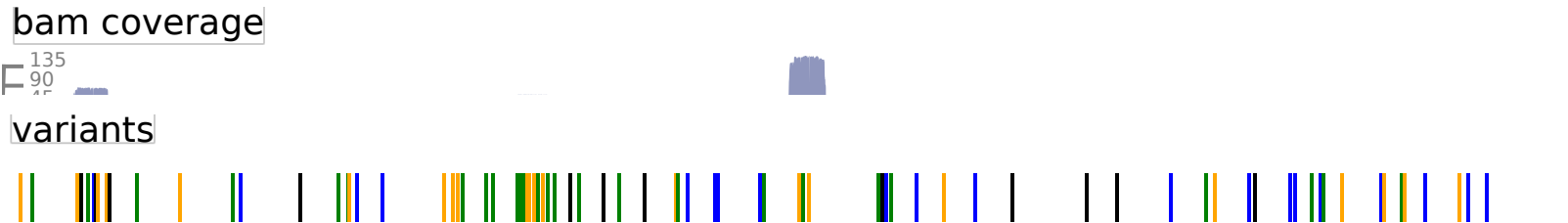
Sample: HG01257

Locus: MICB

Resolution: 4th field

Full typing result:

MICB\*005:02:01



MICB\*005:02:01 (identity: 99.74%)



**Sample: HG01257**

**Locus: TAP1**

**Resolution: 4th field**

**Full typing result:**

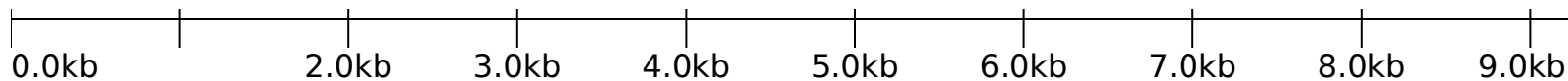
**TAP1\*01:01:01:02**

bam coverage



variants

Alignment (reference: reconstructed gene haplotype 1)



**TAP1\*01:01:01:02 (identity: 99.99%)**



**Sample: HG01257**

**Locus: TAP1**

**Resolution: 4th field**

**Full typing result:**

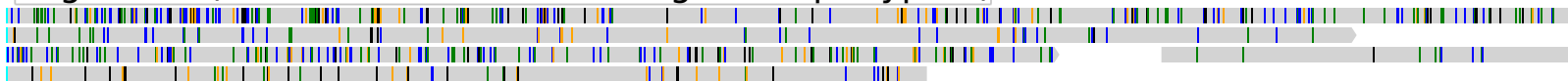
**TAP1\*01:01:01:04**

bam coverage



variants

Alignment (reference: reconstructed gene haplotype 2)



0.0kb 2.0kb 3.0kb 4.0kb 5.0kb 6.0kb 7.0kb 8.0kb 9.0kb

TAP1\*01:01:01:04 (identity: 100.00%)



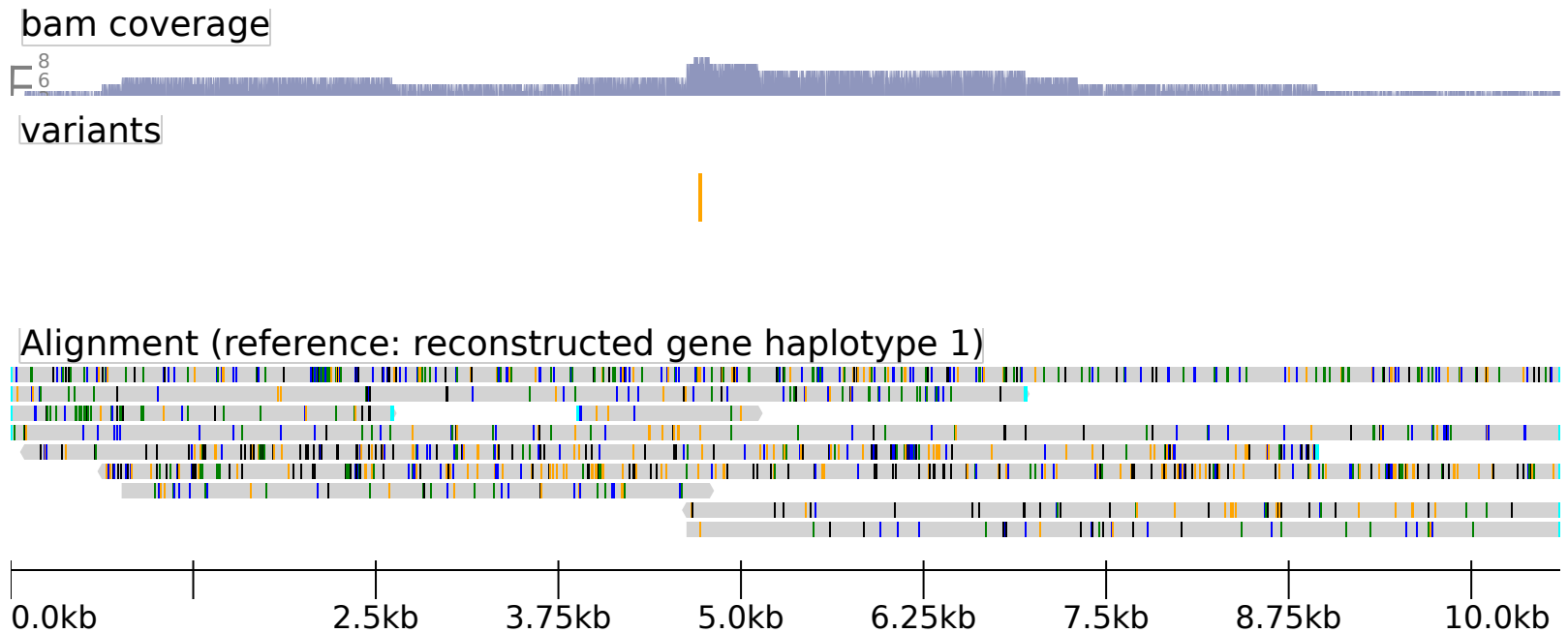
**Sample: HG01257**

**Locus: TAP2**

**Resolution: 4th field**

**Full typing result:**

**TAP2\*01:01:03:03**



**TAP2\*01:01:03:03 (identity: 99.99%)**

**Sample: HG01257**

**Locus: TAP2**

**Resolution: 4th field**

**Full typing result:**

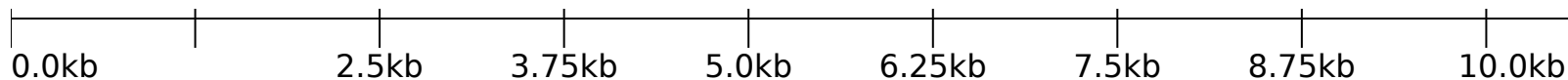
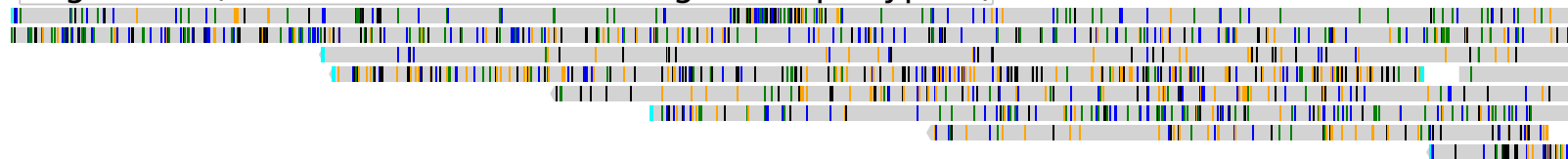
**TAP2\*01:01:07:05**

bam coverage

7  
5

variants

Alignment (reference: reconstructed gene haplotype 2)



TAP2\*01:01:07:05 (identity: 99.98%)

