Locus: HFE

Resolution: 4th field

Full typing result:

HFE*001:01:01

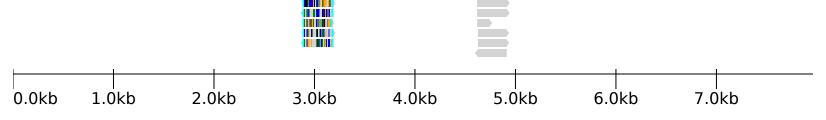
bam coverage

F 4

variants



Alignment (reference: reconstructed gene haplotype 1)



HFE*001:01:01 (identity: 100.00%)

Locus: HLA-A

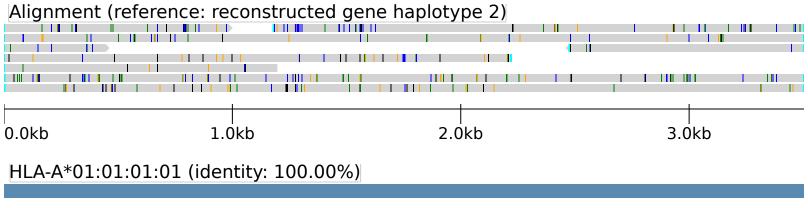
Resolution: 4th field

Full typing result:

HLA-A*01:01:01:01	HLA-A*01:01:01:21	HLA-A*01:452N
HLA-A*01:04:01:01N HLA-A*01:16N HLA-A*01:247N		
HLA-A*01:01:01:03	HLA-A*01:01:01:114	HLA-A*01:01:01:115
HLA-A*01:01:01:12	HLA-A*01:01:01:18	HLA-A*01:01:01:19
HLA-A*01:01:01:20	HLA-A*01:01:01:23	HLA-A*01:01:01:24

bam coverage

variants



HLA-A*01:01:01:21 (identity: 99.86%)

HLA-A*01:452N (identity: 99.86%)

HLA-A*01:04:01:01N (identity: 99.97%)

HLA-A*01:16N (identity: 99.97%)

HLA-A*01:247N (identity: 99.97%)

HLA-A*01:01:03 (identity: 99.97%)

HLA-A*01:01:01:114 (identity: 99.97%)

HLA-A*01:01:01:115 (identity: 99.97%)

HLA-A*01:01:01:12 (identity: 99.97%)

Locus: HLA-A

Resolution: 4th field

Full typing result:

 HLA-A*24:02:01:01
 HLA-A*24:586N
 HLA-A*24:02:01:09

 HLA-A*24:11N
 HLA-A*24:433N
 HLA-A*24:529N

 HLA-A*24:02:01:02L
 HLA-A*24:02:01:04
 HLA-A*24:02:01:05

 HLA-A*24:02:01:06
 HLA-A*24:02:01:08
 HLA-A*24:02:01:108

HLA-A*24:02:01:112 | HLA-A*24:02:01:140 | HLA-A*24:02:01:141

bam coverage

variants

Alignment (reference: reconstructed gene haplotype 1) - 11 HÌ Ш 1.0kb 0.0kb 2.0kb 3.0kb

HLA-A*24:02:01:01 (identity: 100.00%)

HLA-A*24:586N (identity: 99.35%)

23

HLA-A*24:02:01:09 (identity: 99.97%)

HLA-A*24:11N (identity: 99.97%)

HLA-A*24:433N (identity: 99.97%)

HLA-A*24:529N (identity: 99.97%)

HLA-A*24:02:01:02L (identity: 99.97%)

HLA-A*24:02:01:04 (identity: 99.97%)

HLA-A*24:02:01:05 (identity: 99.97%)

HLA-A*24:02:01:06 (identity: 99.97%)

Locus: HLA-B

Resolution: 4th field

Full typing result:

HLA-B*07:02:01:01

bam coverage

variants

Alignment (reference: reconstructed gene haplotype 1)

0.0kb 1.0kb 2.0kb 3.0kb 4.0kb

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

Locus: HLA-B

Resolution: 4th field

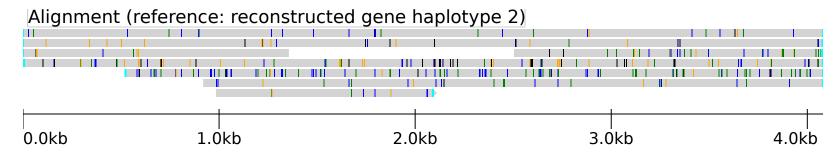
Full typing result:

HLA-B*08:01:01:01

bam coverage

— 7 5

variants



HLA-B*08:01:01:01 (identity: 100.00%)

Locus: HLA-C

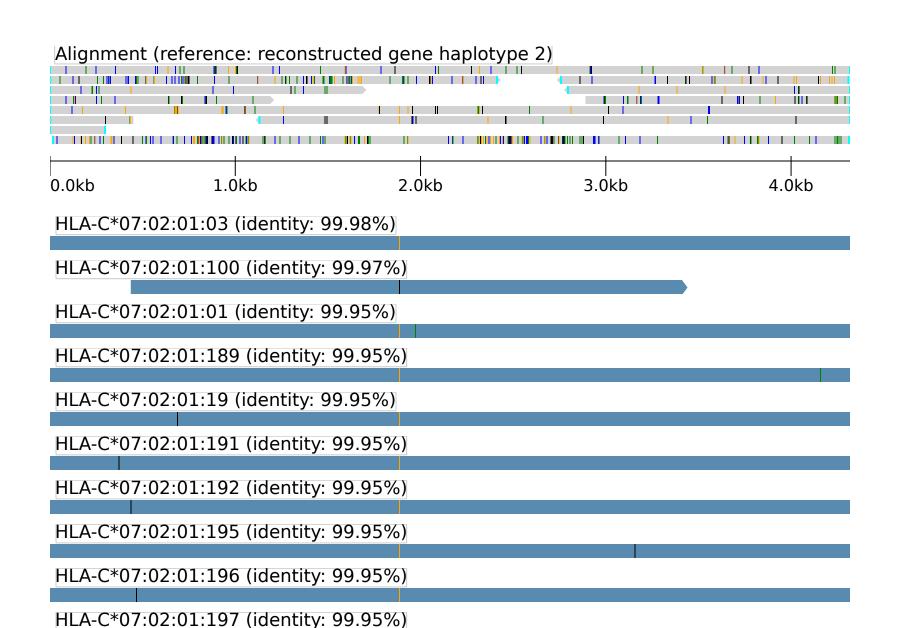
Resolution: 4th field

Full typing result:

HLA-C*07:02:01:03	HLA-C*07:02:01:100	HLA-C*07:02:01:01
HLA-C*07:02:01:189	HLA-C*07:02:01:19	HLA-C*07:02:01:191
HLA-C*07:02:01:192	HLA-C*07:02:01:195	HLA-C*07:02:01:196
HLA-C*07:02:01:197]	

bam coverage

variants



Locus: HLA-C

Resolution: 4th field

Full typing result:

```
HLA-C*07:01:01:01

HLA-C*07:01:01:132

HLA-C*07:01:01:131

HLA-C*07:01:01:16

HLA-C*07:01:05

HLA-C*07:01:08

HLA-C*07:01:31

HLA-C*07:01:76

HLA-C*07:21

HLA-C*07:24

HLA-C*07:250

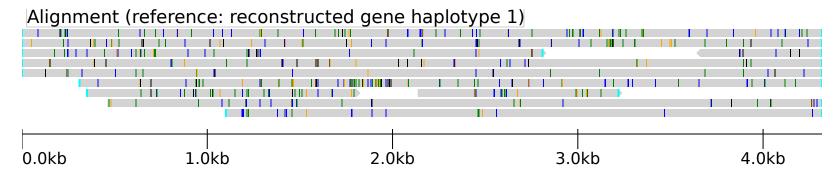
HLA-C*07:412

HLA-C*07:70

HLA-C*07:01:01:25
```

bam coverage

variants



HLA-C*07:01:01 (identity: 99.95%)

HLA-C*07:01:01:132 (identity: 99.95%)

HLA-C*07:01:01:131 (identity: 99.93%)

HLA-C*07:01:01:16 (identity: 99.93%)

HLA-C*07:01:05 (identity: 99.93%)

HLA-C*07:01:08 (identity: 99.93%)

HLA-C*07:01:31 (identity: 99.93%)

HLA-C*07:01:76 (identity: 99.93%)

HLA-C*07:1030 (identity: 99.93%)

HLA-C*07:21 (identity: 99.93%)

Locus: HLA-DMA

Resolution: 4th field

Full typing result:

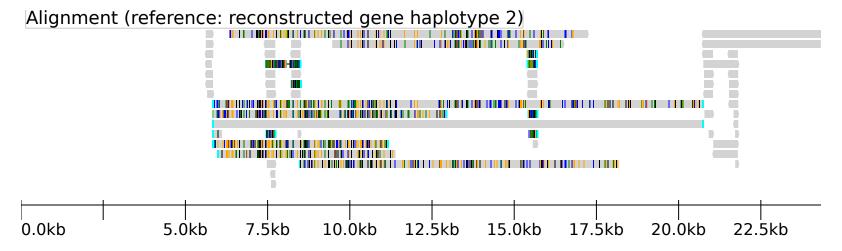
HLA-DMA*01:01:01:01

bam coverage

L 14 10

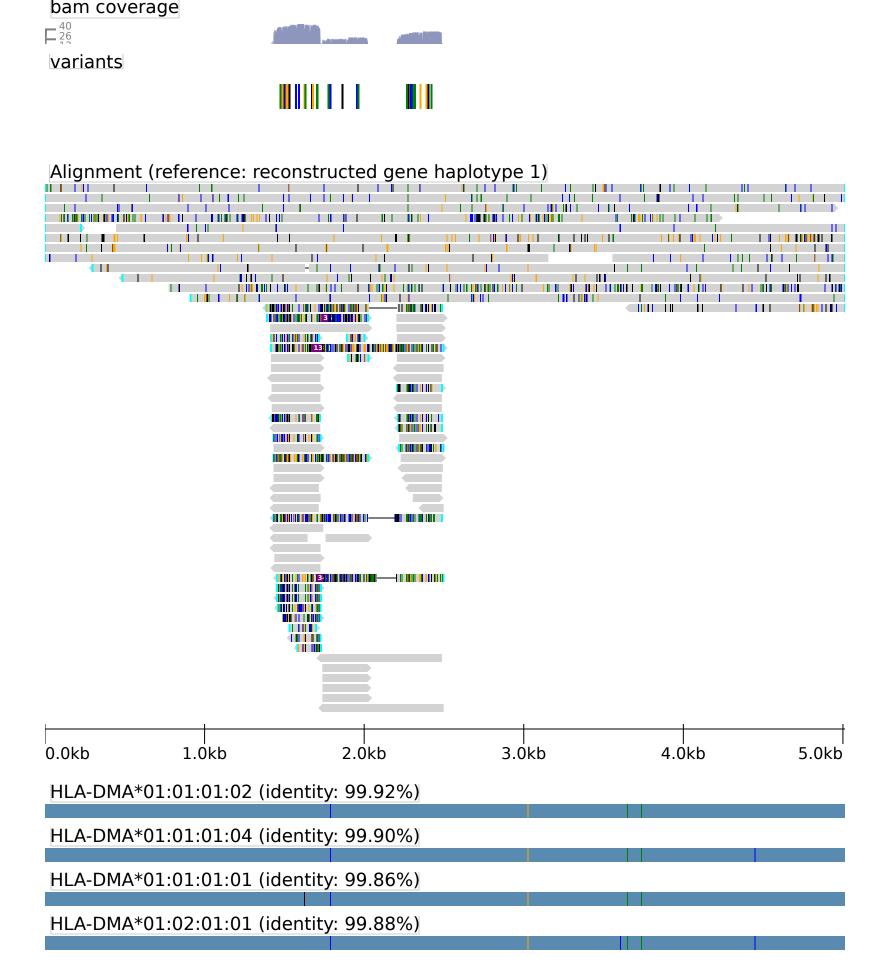
variants





HLA-DMA*01:01:01:01 (identity: 95.57%)

Sample: NA12891 Locus: HLA-DMA Resolution: 4th field Full typing result: HLA-DMA*01:01:01:02 HLA-DMA*01:01:04 HLA-DMA*01:01:01:01 HLA-DMA*01:02:01:01



Locus: HLA-DMB

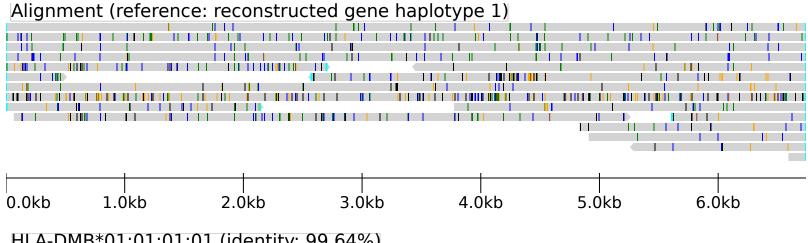
Resolution: 4th field

Full typing result:

HLA-DMB*01:01:01:01 HLA-DMB*01:01:03 HLA-DMB*01:01:01:46 HLA-DMB*01:01:01:48 HLA-DMB*01:03:01:01

bam coverage

variants



HLA-DMB*01:01:01:03 (identity: 99.63%)

HLA-DMB*01:01:01:46 (identity: 99.63%)

HLA-DMB*01:01:01:48 (identity: 99.63%)

HLA-DMB*01:03:01:01 (identity: 99.61%)

Locus: HLA-DMB

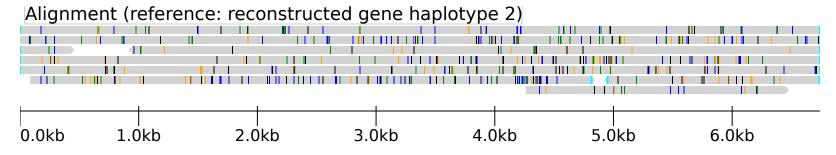
Resolution: 4th field

Full typing result:

HLA-DMB*01:01:01:03

bam coverage

variants



HLA-DMB*01:01:01:03 (identity: 100.00%)

Locus: HLA-DOA

Resolution: 4th field

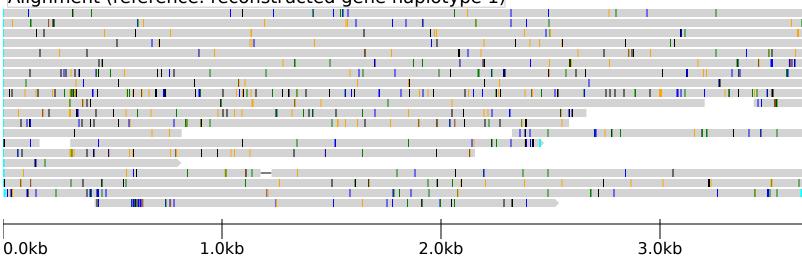
Full typing result:

HLA-DOA*01:01:01:01

bam coverage

variants

Alignment (reference: reconstructed gene haplotype 1)



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

Locus: HLA-DOB

Resolution: 4th field

Full typing result:

HLA-DOB*01:01:01:01

bam coverage

 $-\frac{7}{5}$

variants

Alignment (reference: reconstructed gene haplotype 2)

0.0kb 1.0kb 2.0kb 3.0kb 4.0kb

HLA-DOB*01:01:01 (identity: 100.00%)

Locus: HLA-DOB

Resolution: 4th field

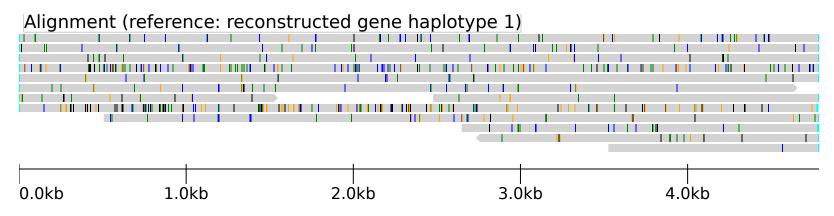
Full typing result:

HLA-DOB*01:01:03:02

bam coverage

__ 10 ___ 7

variants



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

Locus: HLA-DPA1

Resolution: 4th field

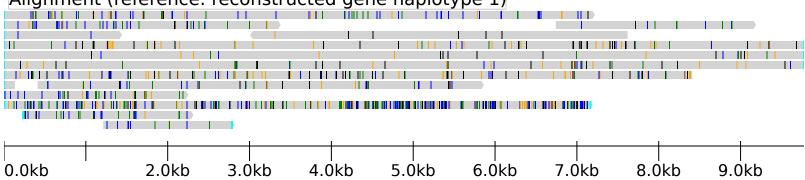
Full typing result:

HLA-DPA1*01:03:01:02



variants

Alignment (reference: reconstructed gene haplotype 1)



HLA-DPA1*01:03:01:02 (identity: 100.00%)

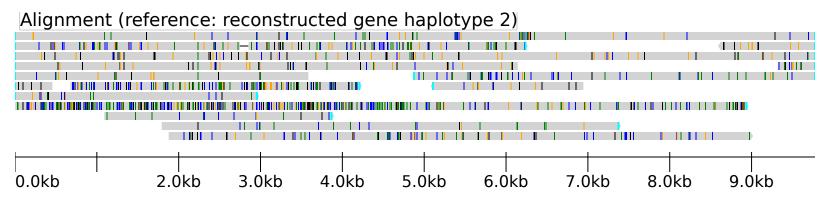
Locus: HLA-DPA1

Resolution: 4th field

Full typing result:

HLA-DPA1*01:03:01:03





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

Locus: HLA-DPA2

Resolution: 4th field

Full typing result:

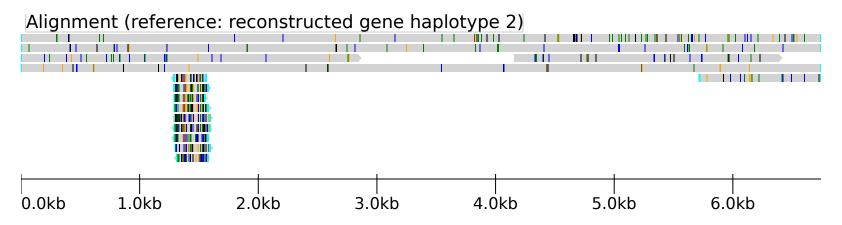
HLA-DPA2*01:01:01:01

bam coverage



variants





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

Locus: HLA-DPA2

Resolution: 4th field

Full typing result:

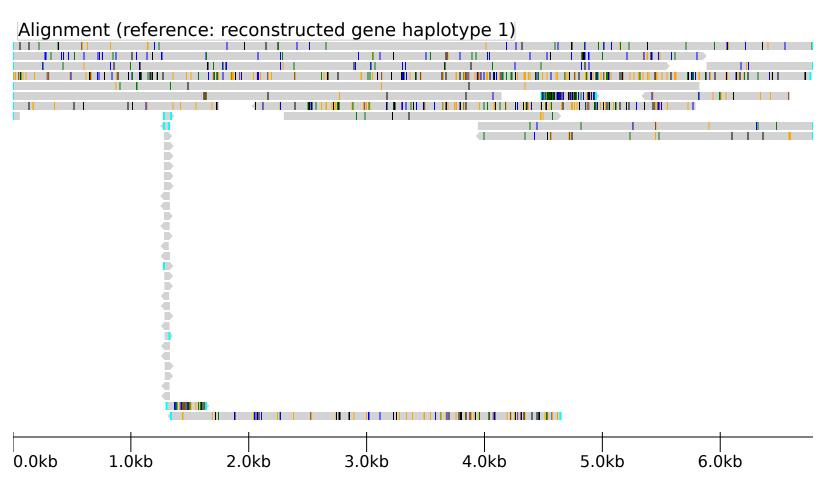
HLA-DPA2*01:01:02

bam coverage

□ 31 21

variants





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

Locus: HLA-DPB1

Resolution: 4th field

Full typing result:

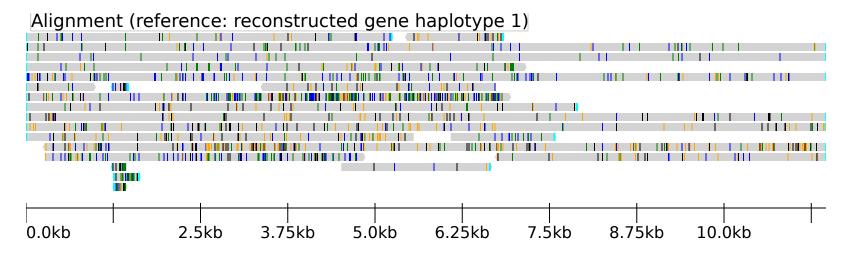
```
HLA-DPB1*03:01:01:01
HLA-DPB1*03:01:01:03
HLA-DPB1*03:01:01:04

HLA-DPB1*03:01:01:07
HLA-DPB1*03:01:01:14
HLA-DPB1*03:01:01:15

HLA-DPB1*03:01:01:16
HLA-DPB1*03:01:01:17
HLA-DPB1*03:01:01:18

HLA-DPB1*03:01:01:19
HLA-DPB1*03:01:01:21
HLA-DPB1*03:01:01:25
```

bam coverage variants



HLA-DPB1*03:01:01:01 (identity: 100.00%)

HLA-DPB1*03:01:01:03 (identity: 99.99%)

HLA-DPB1*03:01:01:04 (identity: 99.99%)

HLA-DPB1*03:01:01:07 (identity: 99.99%)

HLA-DPB1*03:01:01:14 (identity: 99.99%)

HLA-DPB1*03:01:01:15 (identity: 99.99%)

HLA-DPB1*03:01:01:16 (identity: 99.99%)

HLA-DPB1*03:01:01:17 (identity: 99.99%)

HLA-DPB1*03:01:01:18 (identity: 99.99%)

HLA-DPB1*03:01:01:19 (identity: 99.99%)

Locus: HLA-DPB1

Resolution: 4th field

Full typing result:

HLA-DPB1*04:01:01:01 | HLA-DPB1*04:01:01:35 | HLA-DPB1*04:01:01:06 |

HLA-DPB1*04:01:01:07 | HLA-DPB1*04:01:01:08 | HLA-DPB1*04:01:01:10 |

HLA-DPB1*04:01:01:101 | HLA-DPB1*04:01:01:104 | HLA-DPB1*04:01:01:108

| HLA-DPB1*04:01:01:113 | | HLA-DPB1*04:01:01:114 | | HLA-DPB1*04:01:01:115 |

bam coverage

variants



Alignment (reference: reconstructed gene haplotype 2) 11 11 111 1 0.0kb 2.5kb 5.0kb 6.25kb 7.5kb 8.75kb 10.0kb 3.75kb

HLA-DPB1*04:01:01:01 (identity: 100.00%)

HLA-DPB1*04:01:01:35 (identity: 99.93%)

HLA-DPB1*04:01:01:06 (identity: 99.97%)

HLA-DPB1*1423:01N (identity: 99.97%)

HLA-DPB1*04:01:01:02 (identity: 99.93%)

HLA-DPB1*04:01:01:04 (identity: 99.97%)

HLA-DPB1*04:01:01:07 (identity: 99.99%)

HLA-DPB1*04:01:01:08 (identity: 99.99%)

HLA-DPB1*04:01:01:10 (identity: 99.99%)

HLA-DPB1*04:01:01:101 (identity: 99.99%)

Sample: NA12891 **Locus: HLA-DPB2** Resolution: 4th field Full typing result: HLA-DPB2*03:01:01:01 bam coverage **■** 48 32 variants Alignment (reference: reconstructed gene haplotype 2) Alignment (reference: reconstructed gene napiotype 2) Ī 7.5kb 10.0kb 12.5kb 17.5kb 0.0kb 2.5kb 5.0kb 15.0kb

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



Sample: NA12891 Locus: HLA-DQA1

Resolution: 4th field

Full typing result:

HLA-DQA1*01:02:01:03 | HLA-DQA1*01:02:01:01 | HLA-DQA1*01:02:01:02

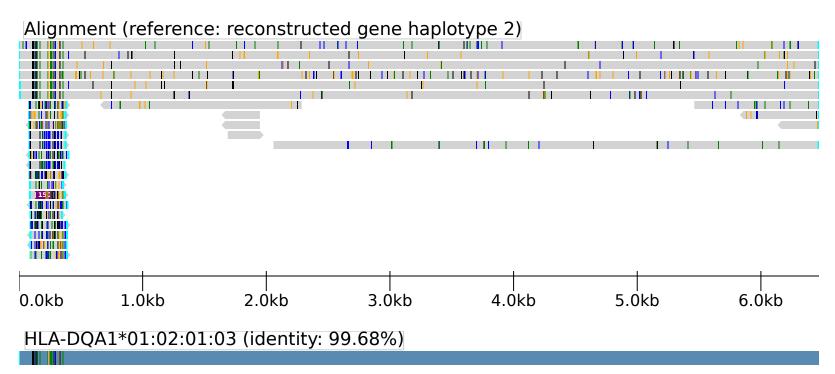
HLA-DQA1*01:02:01:21 | HLA-DQA1*01:02:01:05

bam coverage



variants





HLA-DQA1*01:02:01:01 (identity: 99.66%)

HLA-DQA1*01:02:01:02 (identity: 99.66%)

HLA-DQA1*01:02:01:21 (identity: 99.66%)

HLA-DQA1*01:02:01:05 (identity: 99.64%)

Locus: HLA-DQA1

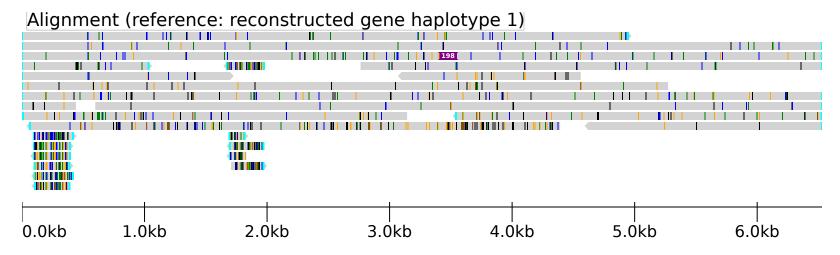
Resolution: 4th field

Full typing result:

HLA-DQA1*05:01:01:02



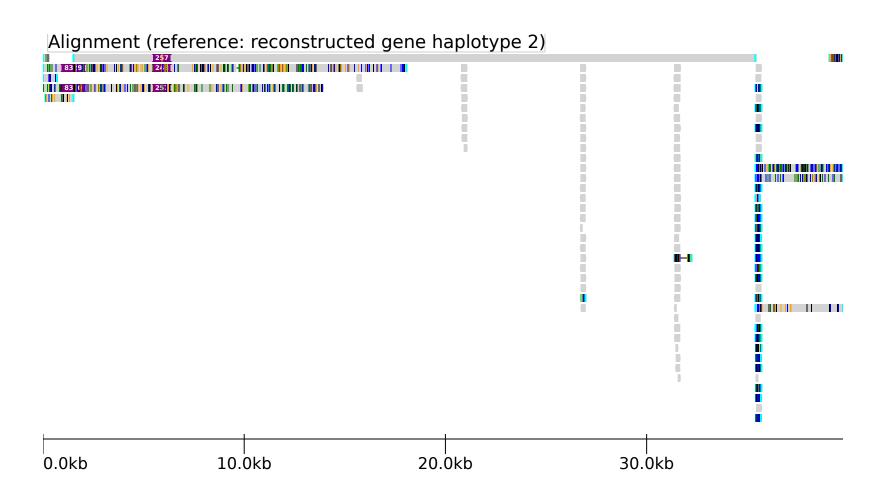
variants



HLA-DQA1*05:01:01:02 (identity: 100.00%)

Sample: NA12891 Locus: HLA-DQA2 Resolution: 4th field Full typing result: HLA-DQA2*01:01:03 bam coverage \[\frac{31}{21} \] variants

HLA-DQA2*01:01:01:03 (identity: 99.77%)

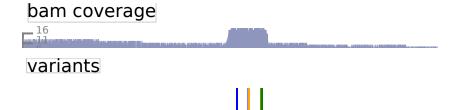


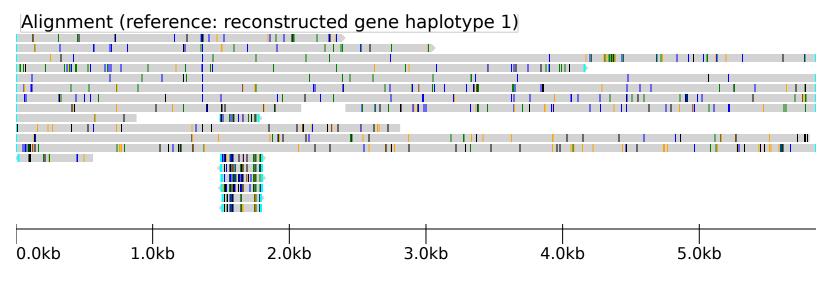
Locus: HLA-DQA2

Resolution: 4th field

Full typing result:

HLA-DQA2*01:01:01:01





HLA-DQA2*01:01:01:01 (identity: 100.00%)

Locus: HLA-DQB1

Resolution: 4th field

Full typing result:

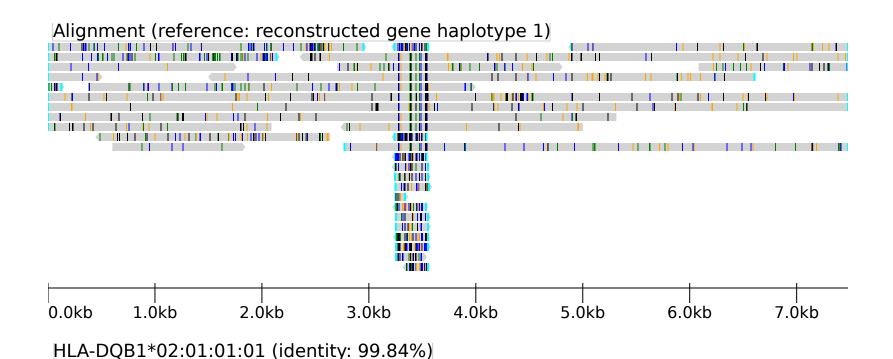
HLA-DQB1*02:01:01:01

bam coverage

□ 19 13

variants





Locus: HLA-DQB1

Resolution: 4th field

Full typing result:

HLA-DQB1*06:02:01:01

bam coverage

variants

Alignment (reference: reconstructed gene haplotype 2)

0.0kb 1.0kb 2.0kb 3.0kb 4.0kb 5.0kb 6.0kb 7.0kb

HLA-DQB1*06:02:01:01 (identity: 100.00%)

Locus: HLA-DQB2

Resolution: 4th field

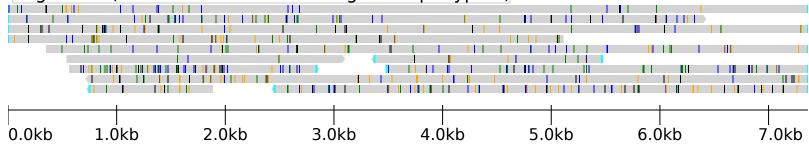
Full typing result:

HLA-DQB2*01:02:01:01

bam coverage

variants

Alignment (reference: reconstructed gene haplotype 1)



HLA-DQB2*01:02:01:01 (identity: 100.00%)

Locus: HLA-DQB2

Resolution: 4th field

Full typing result:

HLA-DQB2*01:02:02:01

bam coverage

variants

Alignment (reference: reconstructed gene haplotype 2)

0.0kb 1.0kb 2.0kb 3.0kb 4.0kb 5.0kb 6.0kb 7.0kb

HLA-DQB2*01:02:02:01 (identity: 100.00%)

Locus: HLA-DRA

Resolution: 4th field

Full typing result:

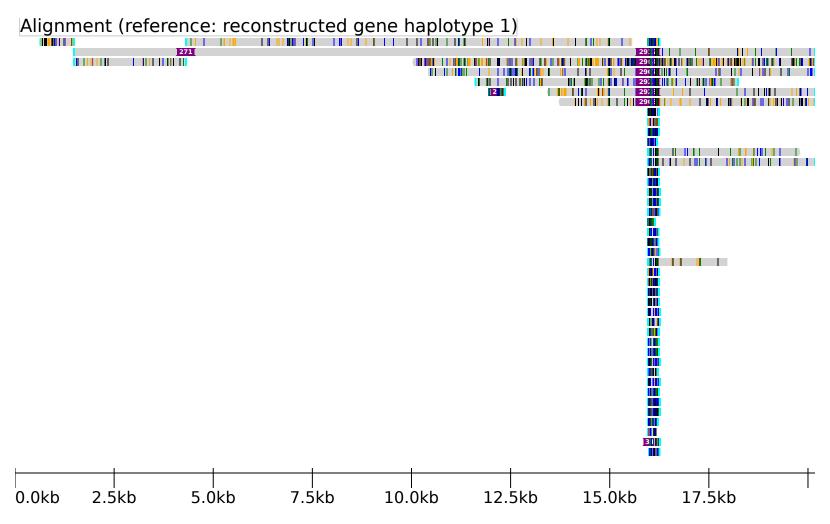
HLA-DRA*01:02:03:01 | HLA-DRA*01:02:03:02

bam coverage

L 24

variants





HLA-DRA*01:02:03:01 (identity: 99.69%)

HLA-DRA*01:02:03:02 (identity: 99.69%)

Locus: HLA-DRA

Resolution: 4th field

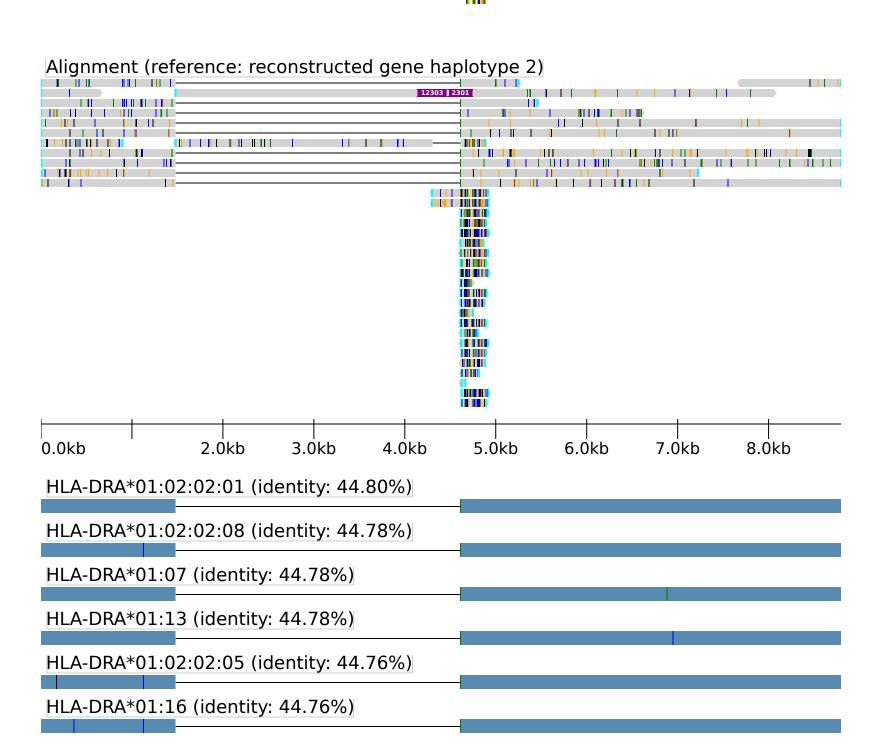
Full typing result:



bam coverage

Lightarrow

variants



Sample: NA12891 Locus: HLA-DRB1 Resolution: 4th field Full typing result: HLA-DRB1*15:01:01:05 | HLA-DRB1*15:14 | HLA-DRB1*15:03:01:01 bam coverage **□** 76 51 الوائلا variants Alignment (reference: reconstructed gene haplotype 2) 5 4 — | 3:.e 30: 2016 [31()] 313 21 21 8 12 7...-0.0kb 2.5kb 5.0kb 7.5kb 10.0kb 12.5kb 15.0kb HLA-DRB1*15:01:01:05 (identity: 45.06%) HLA-DRB1*15:14 (identity: 44.17%) HLA-DRB1*15:03:01:01 (identity: 45.05%)

Sample: NA12891 Locus: HLA-DRB1 Resolution: 4th field Full typing result: HLA-DRB1*03:01:01:01 | HLA-DRB1*03:01:01:08 | HLA-DRB1*03:01:01:09 HLA-DRB1*03:01:01:14 | HLA-DRB1*03:01:01:15 | HLA-DRB1*03:01:01:17 HLA-DRB1*03:01:01:22 | HLA-DRB1*03:01:01:27 | HLA-DRB1*03:01:01:21 HLA-DRB1*03:01:01:05 | HLA-DRB1*03:01:01:02 | HLA-DRB1*03:01:01:06 HLA-DRB1*03:01:01:13 | HLA-DRB1*03:01:01:03 | HLA-DRB1*03:01:01:18 bam coverage variants Alignment (reference: reconstructed gene haplotype 1) _____ - 30: 11 31 111111 3(4 - 301 32 22 34 ---21 22 21 - 30: 11 100101-10010 0.0kb 2.5kb 3.75kb 5.0kb 6.25kb 7.5kb 8.75kb 11.25kb HLA-DRB1*03:01:01:01 (identity: 100.00%) HLA-DRB1*03:01:01:08 (identity: 99.99%) HLA-DRB1*03:01:01:09 (identity: 99.99%) HLA-DRB1*03:01:01:14 (identity: 99.99%) HLA-DRB1*03:01:01:15 (identity: 99.99%) HLA-DRB1*03:01:01:17 (identity: 99.99%) HLA-DRB1*03:01:01:22 (identity: 99.99%) HLA-DRB1*03:01:01:27 (identity: 99.99%) HLA-DRB1*03:01:01:21 (identity: 99.98%) HLA-DRB1*03:01:01:05 (identity: 99.99%)

Locus: HLA-DRB3

Resolution: 4th field

Full typing result:

HLA-DRB3*01:62:01:02

bam coverage

□ 17 11

عيالا

variants

HLA-DRB3*01:62:01:02 (identity: 97.53%)

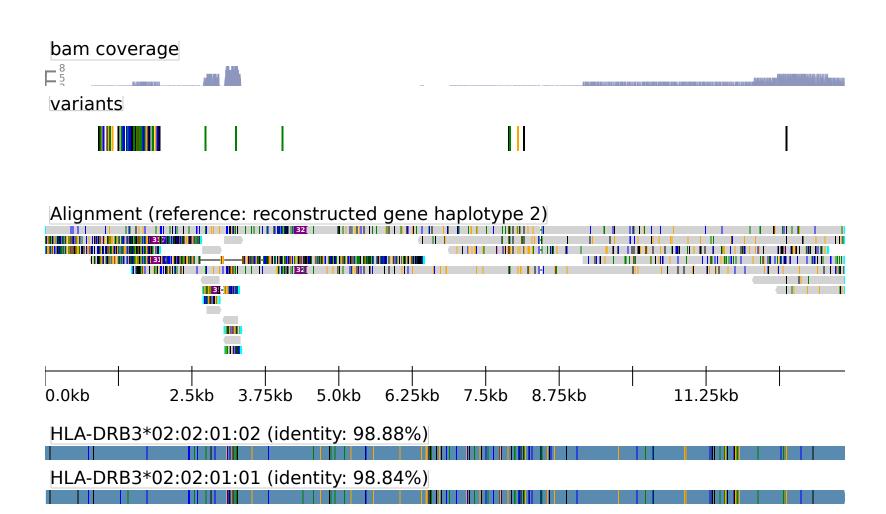
Alignment (reference: reconstructed gene haplotype 1) | 6 0.0kb 2.5kb 3.75kb 5.0kb 6.25kb 7.5kb 8.75kb 11.25kb

Locus: HLA-DRB3

Resolution: 4th field

Full typing result:

HLA-DRB3*02:02:01:02 | HLA-DRB3*02:02:01:01



Sample: NA12891 Locus: HLA-DRB4 Resolution: 4th field Full typing result: HLA-DRB4*02:01N HLA-DRB4*03:01N HLA-DRB4*01:01:01:01 bam coverage variants Alignment (reference: reconstructed gene haplotype 1) 441 31 45 491 485 4: ' 484 43



HLA-DRB4*03:01N (identity: 80.99%)

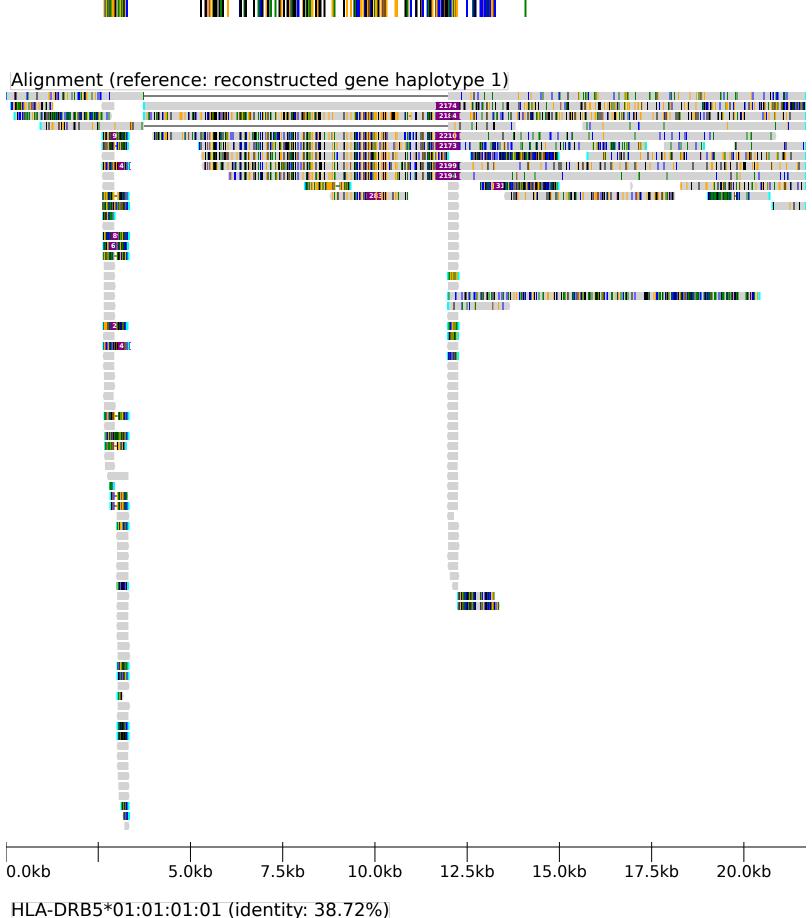
Locus: HLA-DRB5

Resolution: 4th field

Full typing result:

HLA-DRB5*01:01:01:01





Locus: HLA-E

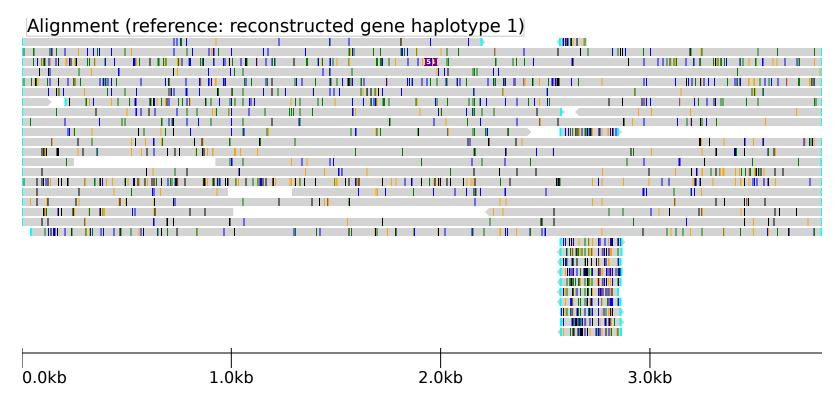
Resolution: 4th field

Full typing result:

HLA-E*01:01:01:01

bam coverage

variants



HLA-E*01:01:01 (identity: 100.00%)

Locus: HLA-F

Resolution: 4th field

Full typing result:

HLA-F*01:01:01:09	HLA-F*01:01:01:10	HLA-F*01:01:01:11
HLA-F*01:01:01:12	HLA-F*01:01:01:27	HLA-F*01:01:01:29
HLA-F*01:01:01:42	HLA-F*01:01:01:43	HLA-F*01:01:06
HLA-F*01:01:07 HLA-F*01:01:08 HLA-F*01:01:11		
HLA-F*01:02:01:02	HLA-F*01:10 HLA	-F*01:12

bam coverage

variants

Alignment (reference: reconstructed gene haplotype 2)

0.0kb 1.0kb 2.0kb 3.0kb

HLA-F*01:01:01:09 (identity: 100.00%)

HLA-F*01:01:01:10 (identity: 99.97%)

HLA-F*01:01:01:11 (identity: 99.97%)

HLA-F*01:01:01:12 (identity: 99.97%)

HLA-F*01:01:01:27 (identity: 99.97%)

HLA-F*01:01:01:29 (identity: 99.97%)

HLA-F*01:01:01:42 (identity: 99.97%)

HLA-F*01:01:01:43 (identity: 99.97%)

HLA-F*01:01:06 (identity: 99.97%)

HLA-F*01:01:07 (identity: 99.97%)

Locus: HLA-F

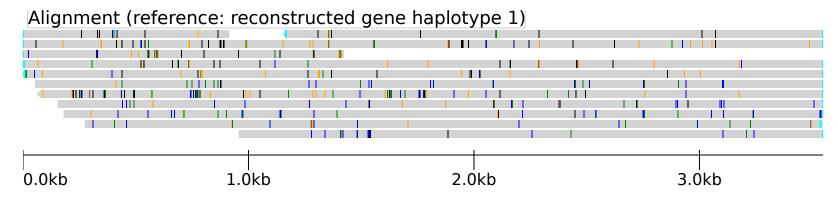
Resolution: 4th field

Full typing result:

HLA-F*01:03:01:01

bam coverage

variants



HLA-F*01:03:01:01 (identity: 100.00%)

Locus: HLA-G

Resolution: 4th field

Full typing result:

HLA-G*01:04:01:01

bam coverage

variants

Alignment (reference: reconstructed gene haplotype 1)

0 250 500 750 1,000 1,500 2,000 2,500 3,000

HLA-G*01:04:01:01 (identity: 100.00%)

Locus: HLA-G

Resolution: 4th field

Full typing result:

HLA-G*01:01:02:01 | HLA-G*01:01:14 | HLA-G*01:01:16

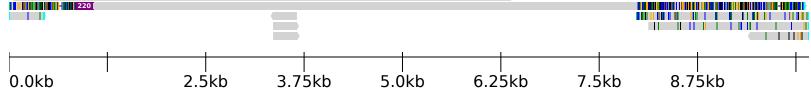
HLA-G*01:01:02:12 | HLA-G*01:01:02:11

bam coverage

L⁴₃

variants





HLA-G*01:01:02:01 (identity: -123.96%)

HLA-G*01:01:14 (identity: -124.00%)

HLA-G*01:01:16 (identity: 99.95%)

HLA-G*01:01:02:12 (identity: -124.00%)

HLA-G*01:01:02:11 (identity: -124.00%)

Locus: HLA-H

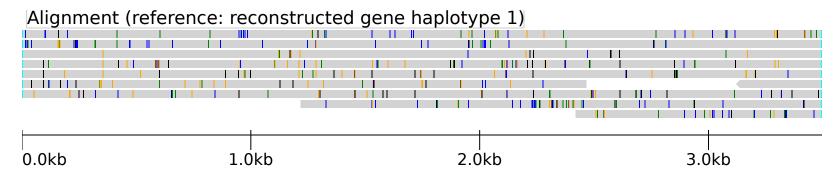
Resolution: 4th field

Full typing result:

HLA-H*02:01:01:01







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

Locus: HLA-J

Resolution: 4th field

Full typing result:

HLA-J*01:01:02

bam coverage

variants

Alignment (reference: reconstructed gene haplotype 1)

0.0kb 1.0kb 2.0kb 3.0kb

HLA-J*01:01:02 (identity: 100.00%)

Locus: HLA-K

Resolution: 4th field

Full typing result:

HLA-K*01:01:01:02

bam coverage

variants

Alignment (reference: reconstructed gene haplotype 1)

0.0kb 1.0kb 2.0kb 3.0kb

HLA-K*01:01:02 (identity: 100.00%)

Locus: HLA-L

Resolution: 4th field

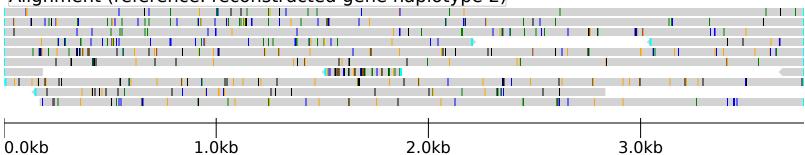
Full typing result:

HLA-L*01:01:01:01

bam coverage

variants

Alignment (reference: reconstructed gene haplotype 2)



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

Locus: HLA-L

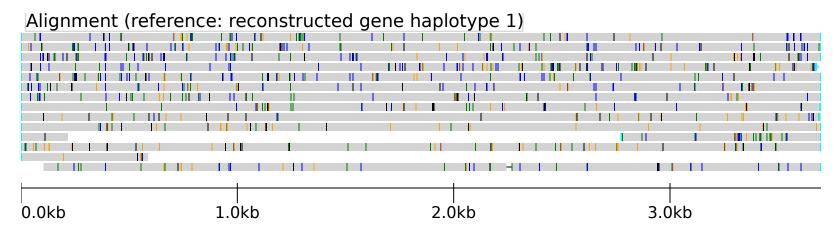
Resolution: 4th field

Full typing result:

HLA-L*01:01:02

bam coverage

variants



HLA-L*01:01:02 (identity: 100.00%)

Locus: HLA-N

Resolution: 4th field

Full typing result:

HLA-N*01:01:01:01

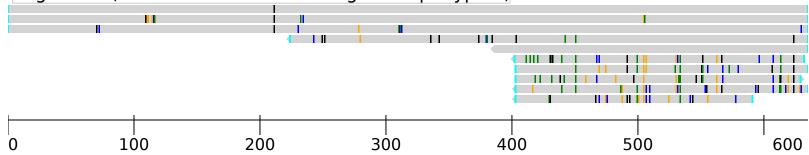
bam coverage

F 6

variants



Alignment (reference: reconstructed gene haplotype 1)



HLA-N*01:01:01:01 (identity: 100.00%)

Locus: HLA-P

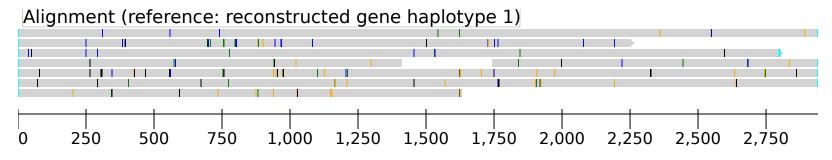
Resolution: 4th field

Full typing result:

HLA-P*01:01:01:01

bam coverage

variants



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

Locus: HLA-P

Resolution: 4th field

Full typing result:

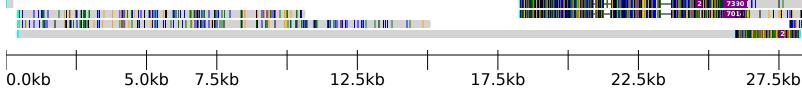
HLA-P*02:01:01:02



variants



Alignment (reference: reconstructed gene haplotype 2)



HLA-P*02:01:01:02 (identity: 99.84%)

Locus: HLA-T

Resolution: 4th field

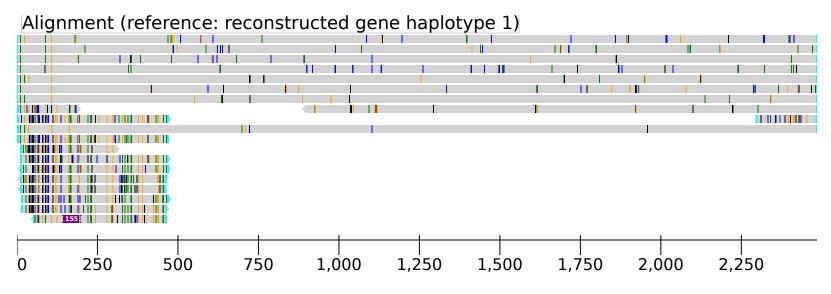
Full typing result:

HLA-T*01:01:01:01

bam coverage

variants





HLA-T*01:01:01:01 (identity: 99.84%)

Locus: HLA-U

Resolution: 4th field

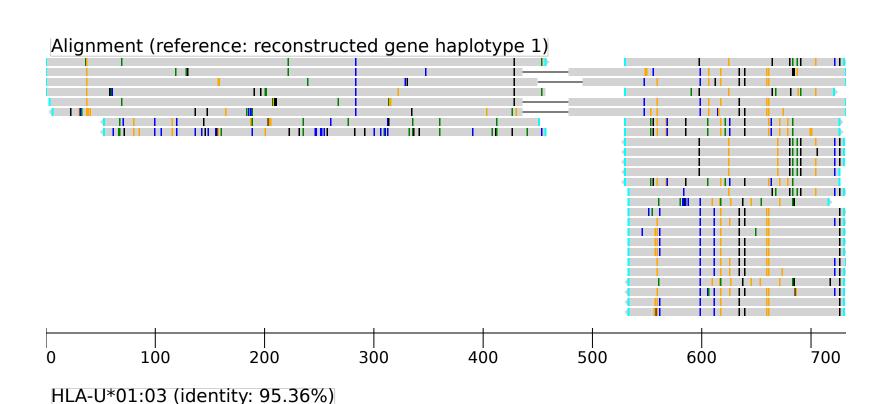
Full typing result:

HLA-U*01:03

bam coverage

23 15

variants



Locus: HLA-V

Resolution: 4th field

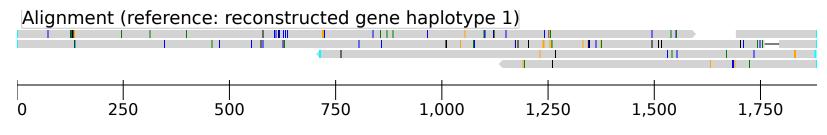
Full typing result:

HLA-V*01:01:01:02

bam coverage

3

variants



HLA-V*01:01:01:02 (identity: 99.42%)

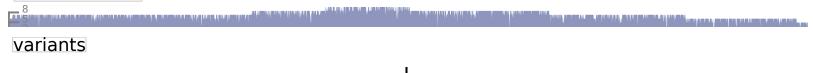
Locus: HLA-W

Resolution: 4th field

Full typing result:

HLA-W*01:01:01:07

bam coverage



Alignment (reference: reconstructed gene haplotype 2)

0 250 500 750 1,000 1,250 1,500 1,750 2,000 2,250 2,500 2,750

HLA-W*01:01:01:07 (identity: 99.83%)

Locus: HLA-W

Resolution: 4th field

Full typing result:

HLA-W*01:01:01:02 | HLA-W*01:01:01:06

bam coverage

- 14

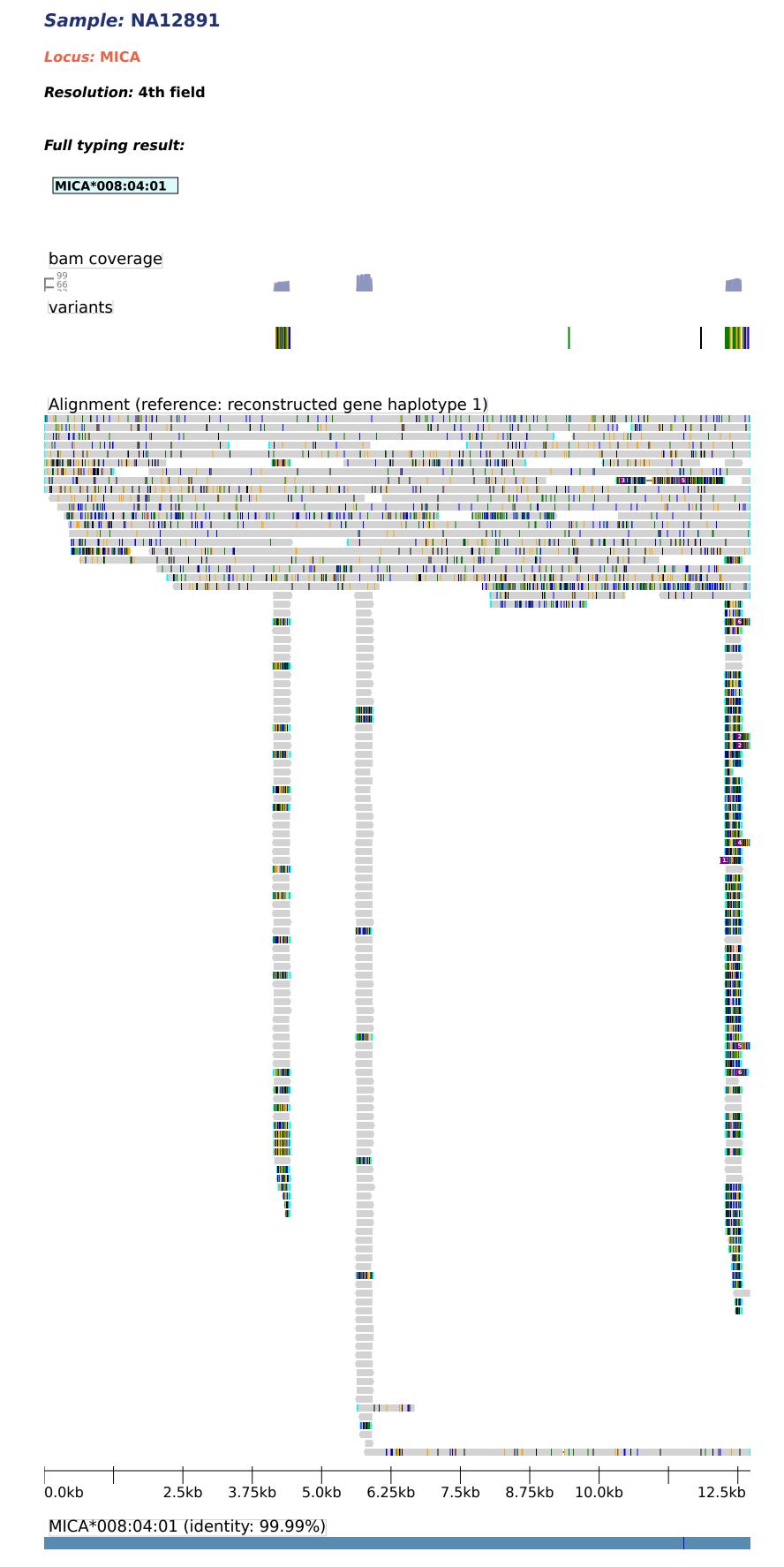
variants

Alignment (reference: reconstructed gene haplotype 1)

0 250 500 750 1,000 1,250 1,500 1,750 2,000 2,250 2,500 2,750

HLA-W*01:01:01:02 (identity: 100.00%)

HLA-W*01:01:01:06 (identity: 99.97%)



Sample: NA12891 Locus: MICB Resolution: 4th field Full typing result: MICB*052 bam coverage 166 111 variants Alignment (reference: reconstructed gene haplotype 1) Alignment (reference: reconstructed gene haplotype 1) -0.0kb 5.0kb 7.5kb 8.75kb 10.0kb 2.5kb 3.75kb 6.25kb 12.5kb MICB*052 (identity: 99.98%)

Locus: TAP1

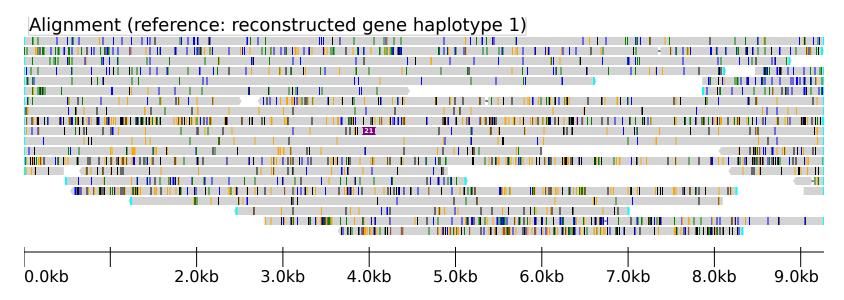
Resolution: 4th field

Full typing result:

TAP1*01:01:01:05

bam coverage

variants



TAP1*01:01:01:05 (identity: 100.00%)

Locus: TAP1

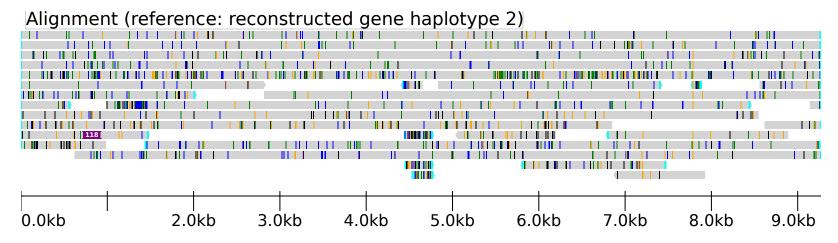
Resolution: 4th field

Full typing result:

TAP1*01:01:01:01

bam coverage





TAP1*01:01:01:01 (identity: 99.97%)

Locus: TAP2

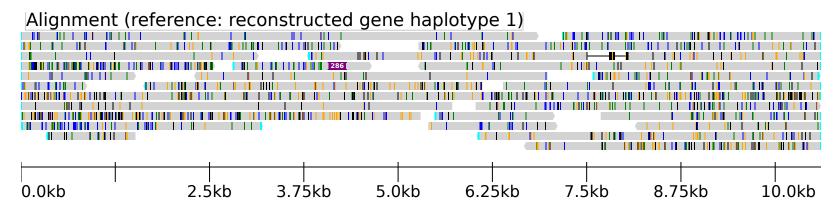
Resolution: 4th field

Full typing result:

TAP2*01:01:03:01







TAP2*01:01:03:01 (identity: 99.92%)

Locus: TAP2

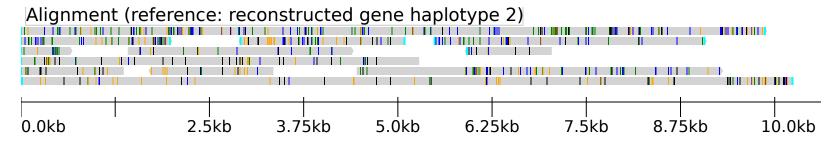
Resolution: 4th field

Full typing result:

TAP2*02:01:02:01



variants



TAP2*02:01:02:01 (identity: 99.11%)