Locus: HLA-A

Resolution: 4th field

Full typing result:

HLA-A*03:397

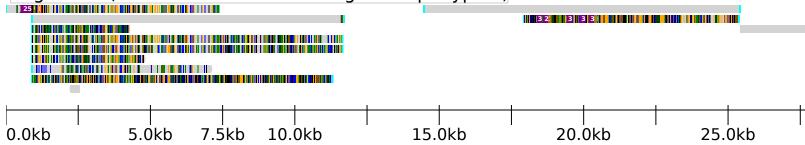
HLA-A*03:284N

bam coverage

variants



Alignment (reference: reconstructed gene haplotype 1)



HLA-A*03:397 (identity: 100.00%)

HLA-A*03:284N (identity: 100.00%)

Locus: HLA-B

Resolution: 4th field

Full typing result:

HLA-B*15:01:01:01

bam coverage

variants

HLA-B*15:01:01:01 (identity: 99.98%)

Locus: HLA-B

Resolution: 4th field

Full typing result:

HLA-B*44:02:01:01

bam coverage

variants

Alignment (reference: reconstructed gene haplotype 2) 0.0kb 1.0kb 2.0kb 3.0kb 4.0kb

HLA-B*44:02:01:01 (identity: 100.00%)

Locus: HLA-C

Resolution: 4th field

Full typing result:

HLA-C*03:04:01:01 HLA-C*03:04:01:02 HLA-C*03:04:01:86

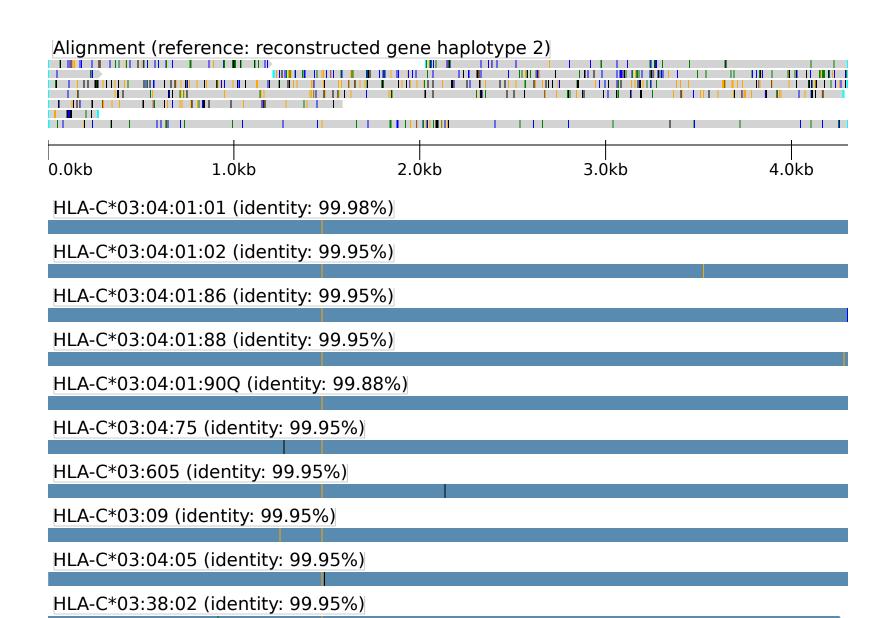
HLA-C*03:04:01:88 HLA-C*03:04:01:90Q HLA-C*03:04:75

HLA-C*03:605 HLA-C*03:09 HLA-C*03:04:05

HLA-C*03:38:02

bam coverage

variants



Locus: HLA-C

Resolution: 4th field

Full typing result:

bam coverage

variants

Alignment (reference: reconstructed gene haplotype 1)

0.0kb 1.0kb 2.0kb 3.0kb 4.0kb

HLA-C*05:01:01:83 (identity: 99.98%)

HLA-C*05:01:01:02 (identity: 99.93%)

HLA-C*05:287 (identity: 99.95%)

HLA-C*05:293 (identity: 99.95%)

HLA-C*05:01:01:47 (identity: 99.91%)

HLA-C*05:285 (identity: 99.95%)

HLA-C*05:32 (identity: 99.95%)

HLA-C*05:286 (identity: 99.95%)

HLA-C*05:255 (identity: 99.95%)

Locus: HLA-DMA

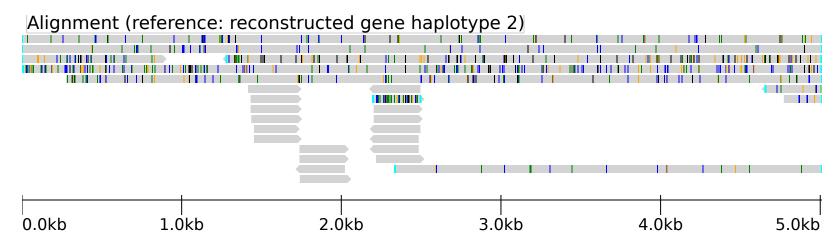
Resolution: 4th field

Full typing result:

HLA-DMA*01:01:01:01

bam coverage

variants



HLA-DMA*01:01:01 (identity: 99.94%)

Sample: NA12877 Locus: HLA-DMA Resolution: 4th field Full typing result: HLA-DMA*01:01:01:02 bam coverage **□** 27 18 variants Alignment (reference: reconstructed gene haplotype 1)

10.0kb

12.5kb

15.0kb

17.5kb

HLA-DMA*01:01:02 (identity: 99.94%)

5.0kb

7.5kb

0.0kb

2.5kb

Locus: HLA-DMB

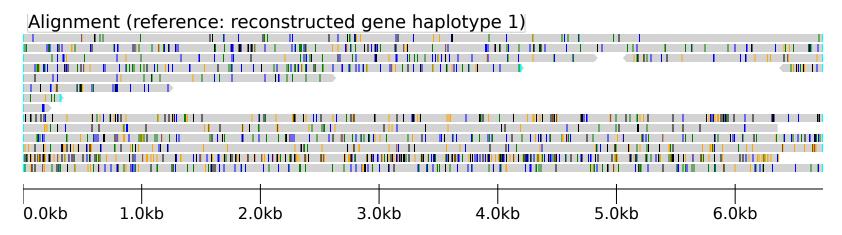
Resolution: 4th field

Full typing result:

HLA-DMB*01:01:01:01

bam coverage

variants



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

Locus: HLA-DOA

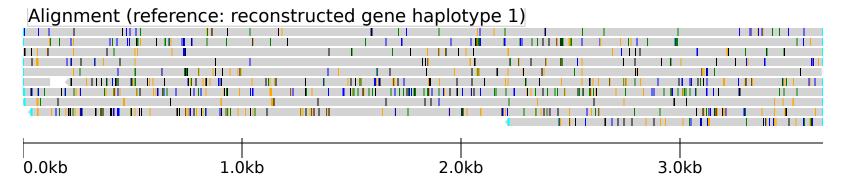
Resolution: 4th field

Full typing result:

HLA-DOA*01:01:02:45

bam coverage

variants



HLA-DOA*01:01:02:45 (identity: 99.97%)

Locus: HLA-DOA

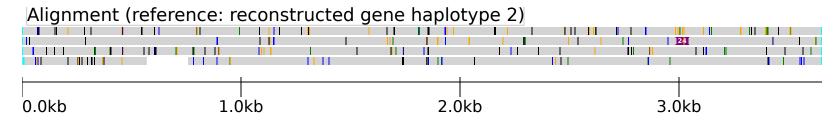
Resolution: 4th field

Full typing result:

HLA-DOA*01:01:02:02

bam coverage

variants



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

Locus: HLA-DOB

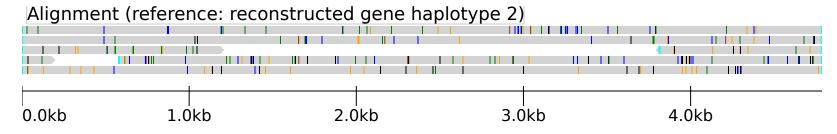
Resolution: 4th field

Full typing result:

HLA-DOB*01:01:01:03

bam coverage

variants



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

Locus: HLA-DOB

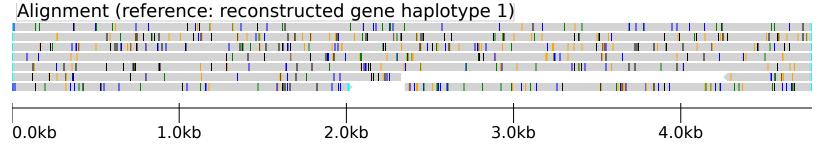
Resolution: 4th field

Full typing result:

HLA-DOB*01:02:01:01

bam coverage

variants



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

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 1)

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

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%)

Locus: HLA-DPA1

Resolution: 4th field

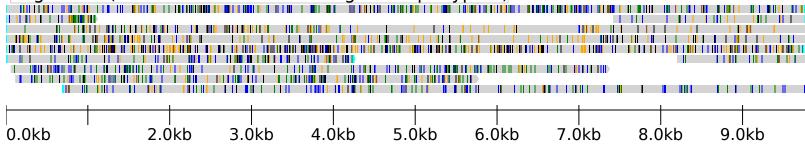
Full typing result:

HLA-DPA1*01:03:01:04

bam coverage

variants

Alignment (reference: reconstructed gene haplotype 2)



HLA-DPA1*01:03:01:04 (identity: 99.97%)

Locus: HLA-DPA2

Resolution: 4th field

Full typing result:

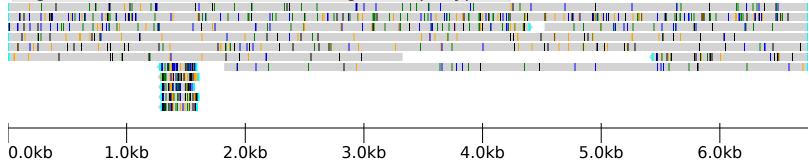
HLA-DPA2*01:01:01:02

bam coverage

 $=\frac{10}{7}$

variants

Alignment (reference: reconstructed gene haplotype 1)

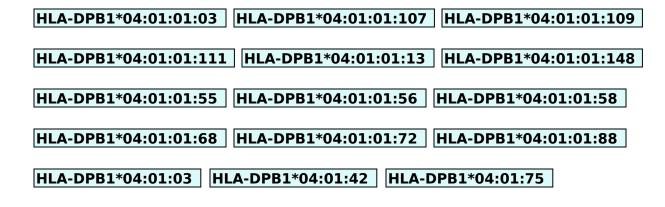


HLA-DPA2*01:01:01:02 (identity: 100.00%)

Locus: HLA-DPB1

Resolution: 4th field

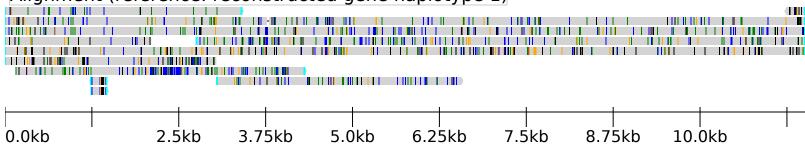
Full typing result:



bam coverage

variants

Alignment (reference: reconstructed gene haplotype 1)



HLA-DPB1*04:01:01:03 (identity: 99.98%)

HLA-DPB1*04:01:01:107 (identity: 99.97%)

HLA-DPB1*04:01:01:109 (identity: 99.97%)

HLA-DPB1*04:01:01:111 (identity: 99.97%)

HLA-DPB1*04:01:01:13 (identity: 99.97%)

HLA-DPB1*04:01:01:148 (identity: 99.97%)

HLA-DPB1*04:01:01:55 (identity: 99.97%)

HLA-DPB1*04:01:01:56 (identity: 99.97%)

HLA-DPB1*04:01:01:58 (identity: 99.97%)

HLA-DPB1*04:01:01:68 (identity: 99.97%)

Locus: HLA-DPB1

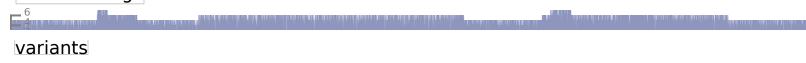
Resolution: 4th field

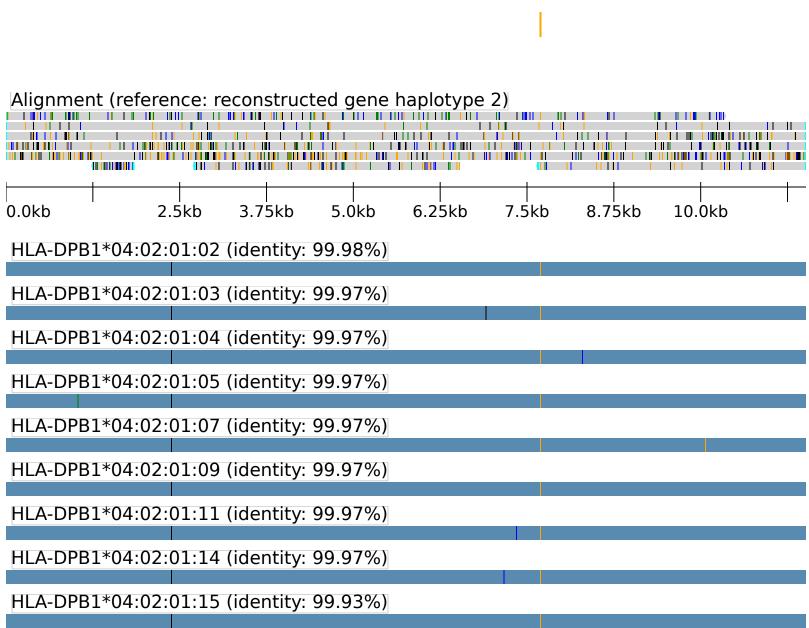
Full typing result:

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	HLA-DPB1*04:02:01:26

bam coverage

HLA-DPB1*04:02:01:18 (identity: 99.97%)





Sample: NA12877 Locus: HLA-DPB2 Resolution: 4th field Full typing result: HLA-DPB2*03:01:01:02 bam coverage **□** 35 23 13 variants Alignment (reference: reconstructed gene haplotype 2) III 10.0kb 0.0kb 2.5kb 5.0kb 7.5kb 12.5kb 15.0kb 17.5kb

HLA-DPB2*03:01:01:02 (identity: 99.94%)

Locus: HLA-DPB2 Resolution: 4th field Full typing result: HLA-DPB2*01:01:02 bam coverage **□** 57 38 variants \parallel Alignment (reference: reconstructed gene haplotype 1) Alignment (reference: reconstructed gene haplotype 1) Ш П Ш ш ш H 0.0kb 2.5kb 5.0kb 7.5kb 10.0kb 12.5kb 15.0kb HLA-DPB2*01:01:02 (identity: 99.70%)

Sample: NA12877

Sample: NA12877

Locus: HLA-DQA1

Resolution: 4th field

Full typing result:

HLA-DQA1*03:01:01:01 HLA-DQA1*03:03:01:01 HLA-DQA1*03:10

HLA-DQA1*03:01:01:04 HLA-DQA1*03:03:01:22 HLA-DQA1*03:71

bam coverage

Alignment (reference: reconstructed gene haplotype 2) 4 0.0kb 1.0kb 2.0kb 4.0kb 5.0kb 3.0kb 6.0kb

HLA-DQA1*03:01:01:01 (identity: 99.91%)

HLA-DQA1*03:03:01:01 (identity: 99.89%)

HLA-DQA1*03:10 (identity: 99.89%)

HLA-DQA1*03:01:01:04 (identity: 99.89%)

HLA-DQA1*03:03:01:22 (identity: 99.88%)

HLA-DQA1*03:71 (identity: 99.88%)

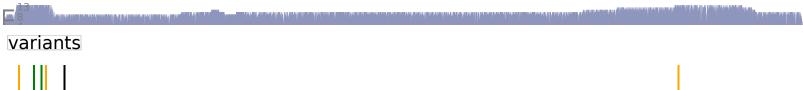
Locus: HLA-DQA1

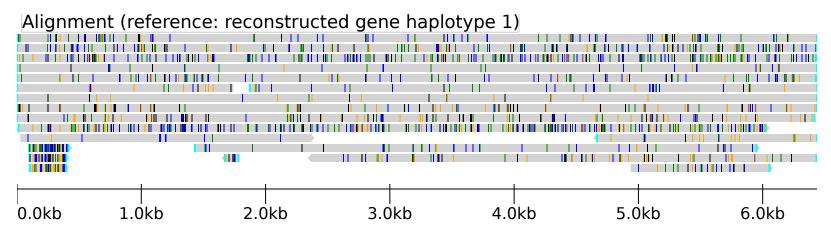
Resolution: 4th field

Full typing result:

HLA-DQA1*03:03:01:01







HLA-DQA1*03:03:01:01 (identity: 100.00%)

Sample: NA12877 **Locus: HLA-DQA2** Resolution: 4th field Full typing result: HLA-DQA2*01:08 bam coverage variants Alignment (reference: reconstructed gene haplotype 1) П Ш Ī Ш

20.0kb

30.0kb

HLA-DQA2*01:08 (identity: 11.74%)

10.0kb

0.0kb

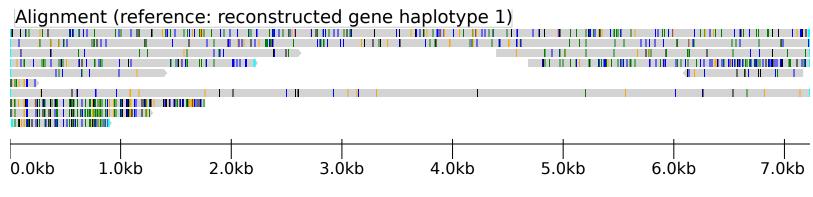
Locus: HLA-DQB1

Resolution: 4th field

Full typing result:

HLA-DQB1*03:01:01:01





HLA-DQB1*03:01:01:01 (identity: 99.99%)

Locus: HLA-DQB1

Resolution: 4th field

Full typing result:

 HLA-DQB1*03:04:01:01
 HLA-DQB1*03:01:01:20
 HLA-DQB1*03:55

 HLA-DQB1*03:01:01:31
 HLA-DQB1*03:01:01:34
 HLA-DQB1*03:01:01:51

 HLA-DQB1*03:01:01:52
 HLA-DQB1*03:01:52
 HLA-DQB1*03:01:03

 HLA-DQB1*03:01:01:43
 HLA-DQB1*03:01:01:01
 HLA-DQB1*03:421

HLA-DQB1*03:01:01:44 | HLA-DQB1*03:01:01:05 | HLA-DQB1*03:01:01:10 |

HLA-DQB1*03:01:01:51 (identity: 99.23%)

HLA-DQB1*03:01:01:52 (identity: 99.23%)

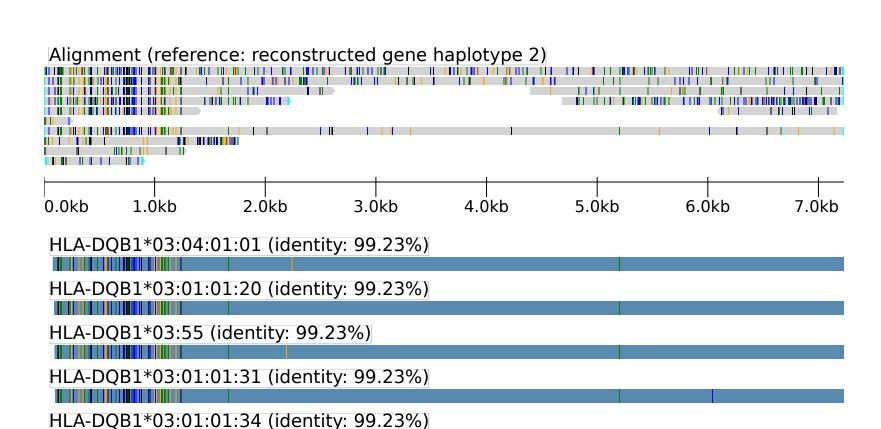
HLA-DQB1*03:01:52 (identity: 99.23%)

HLA-DQB1*03:01:03 (identity: 99.23%)

HLA-DQB1*03:01:01:43 (identity: 99.23%)

ham coverage





Locus: HLA-DQB2

Resolution: 4th field

Full typing result:

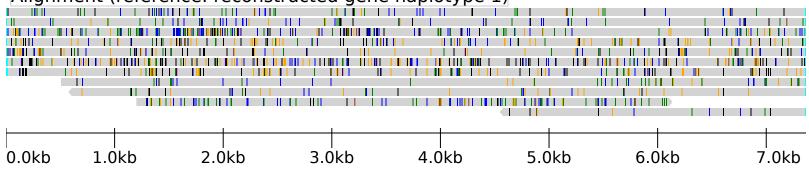
HLA-DQB2*01:01:01:02 HLA-DQB2*01:01:15

bam coverage

- 10

variants

Alignment (reference: reconstructed gene haplotype 1)



HLA-DQB2*01:01:01:02 (identity: 99.99%)

HLA-DQB2*01:01:01:15 (identity: 99.97%)

Locus: HLA-DQB2

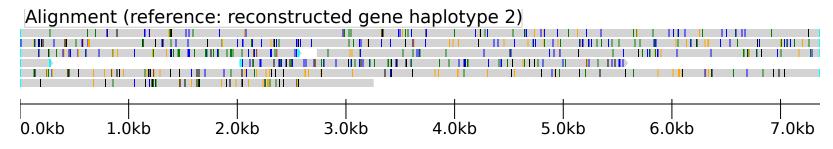
Resolution: 4th field

Full typing result:

HLA-DQB2*01:02:01:01 HLA-DQB2*01:02:01:02

bam coverage

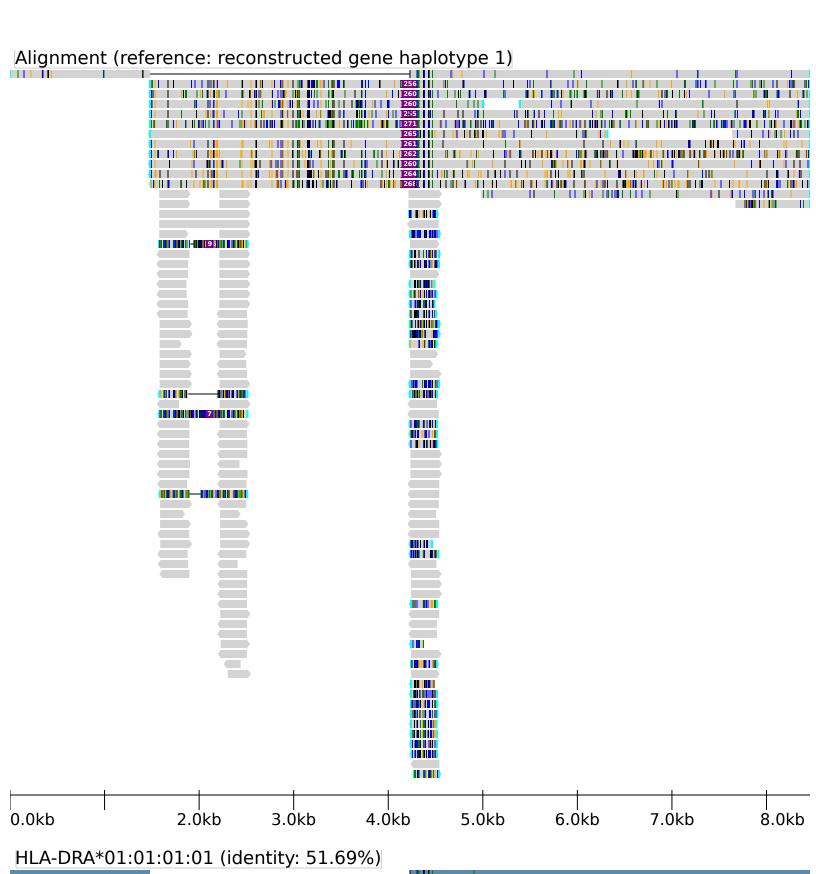
variants



HLA-DQB2*01:02:01:01 (identity: 99.96%)

HLA-DQB2*01:02:01:02 (identity: 99.93%)

Sample: NA12877 Locus: HLA-DRA Resolution: 4th field Full typing result: HLA-DRA*01:01:01 | HLA-DRA*01:01:03 bam coverage \$\begin{align*} \begin{align*} \begin{align

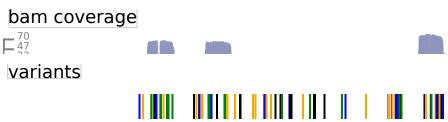


HLA-DRA*01:01:01:03 (identity: 51.69%)

Sample: NA12877 Locus: HLA-DRA Resolution: 4th field

Full typing result:

HLA-DRA*01:01:01:03





Sample: NA12877 Locus: HLA-DRB1 Resolution: 4th field Full typing result: HLA-DRB1*04:01:01:11 | HLA-DRB1*04:01:01:16 | HLA-DRB1*04:01:01:01 HLA-DRB1*04:01:01:04 | HLA-DRB1*04:01:01:21 | HLA-DRB1*04:01:01:25 HLA-DRB1*04:01:01:19 | HLA-DRB1*04:01:01:13 | HLA-DRB1*04:01:01:14 HLA-DRB1*04:01:01:23 | HLA-DRB1*04:01:01:09 | HLA-DRB1*04:01:01:10 HLA-DRB1*04:01:01:20 | HLA-DRB1*04:01:01:08 | HLA-DRB1*04:01:01:12 bam coverage **■** 58 39 variants Alignment (reference: reconstructed gene haplotype 2) 3: 30 - | | | | | | | | | | | -| | | | | | | | | | 91 27 5 2 0.0kb 10.0kb 20.0kb 30.0kb 40.0kb HLA-DRB1*04:01:01:11 (identity: -86.31%) HLA-DRB1*04:01:01:16 (identity: -86.27%) HLA-DRB1*04:01:01:01 (identity: -86.31%) HLA-DRB1*04:01:01:04 (identity: -86.30%) HLA-DRB1*04:01:01:21 (identity: -86.31%) HLA-DRB1*04:01:01:25 (identity: -86.31%) HLA-DRB1*04:01:01:19 (identity: -86.28%) HLA-DRB1*04:01:01:13 (identity: -86.28%) HLA-DRB1*04:01:01:14 (identity: -86.28%) HLA-DRB1*04:01:01:23 (identity: -86.29%)

Sample: NA12877 Locus: HLA-DRB1 Resolution: 4th field Full typing result: HLA-DRB1*04:01:01:20 | HLA-DRB1*04:01:01:04 | HLA-DRB1*04:01:01:25 HLA-DRB1*04:01:01:11 | HLA-DRB1*04:01:01:13 | HLA-DRB1*04:01:01:14 HLA-DRB1*04:01:01:09 | HLA-DRB1*04:01:01:10 | HLA-DRB1*04:01:01:01 HLA-DRB1*04:01:01:23 | HLA-DRB1*04:08:01:02 | HLA-DRB1*04:01:01:17 bam coverage **□** 57 38 variants Alignment (reference: reconstructed gene haplotype 1) Ш Ш 31 30 21 5 6: 8 31 **(** 2 0.0kb 5.0kb 17.5kb 22.5kb 7.5kb 12.5kb 27.5kb HLA-DRB1*04:01:01:20 (identity: -3.11%) HLA-DRB1*04:01:01:04 (identity: -3.11%) HLA-DRB1*04:01:01:25 (identity: -3.11%) HLA-DRB1*04:01:01:11 (identity: -3.11%) HLA-DRB1*04:01:01:13 (identity: -3.11%) HLA-DRB1*04:01:01:14 (identity: -3.11%) HLA-DRB1*04:01:01:09 (identity: -3.11%) HLA-DRB1*04:01:01:10 (identity: -3.11%) HLA-DRB1*04:01:01:01 (identity: -3.11%) HLA-DRB1*04:01:01:08 (identity: -3.11%)

Locus: HLA-DRB3

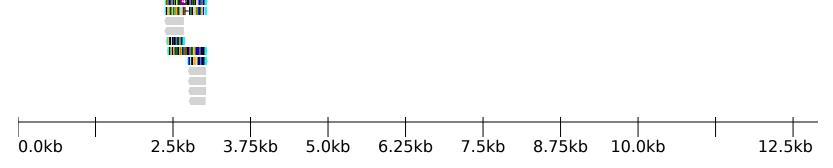
Resolution: 4th field

Full typing result:

HLA-DRB3*01:01:02:05 | HLA-DRB3*01:62:01:01 | HLA-DRB3*01:62:01:02

bam coverage F 5 5 variants

Alignment (reference: reconstructed gene haplotype 1)



HLA-DRB3*01:01:02:05 (identity: 97.68%)

HLA-DRB3*01:62:01:01 (identity: 97.68%)

HLA-DRB3*01:62:01:02 (identity: 97.68%)

Locus: HLA-DRB3

Resolution: 4th field

Full typing result:

HLA-DRB3*01:01:02:01

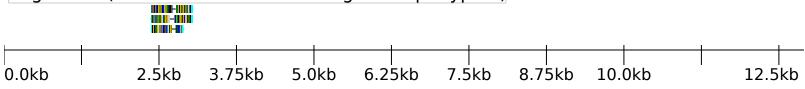
bam coverage

 $\frac{3}{2}$



variants

Alignment (reference: reconstructed gene haplotype 2)



HLA-DRB3*01:01:02:01 (identity: 86.75%)

Locus: HLA-DRB4

Resolution: 4th field

Full typing result:

HLA-DRB4*01:03:01:01 | HLA-DRB4*01:03:01:03 | HLA-DRB4*01:03:01:18

HLA-DRB4*01:152 | HLA-DRB4*01:03:01:02N

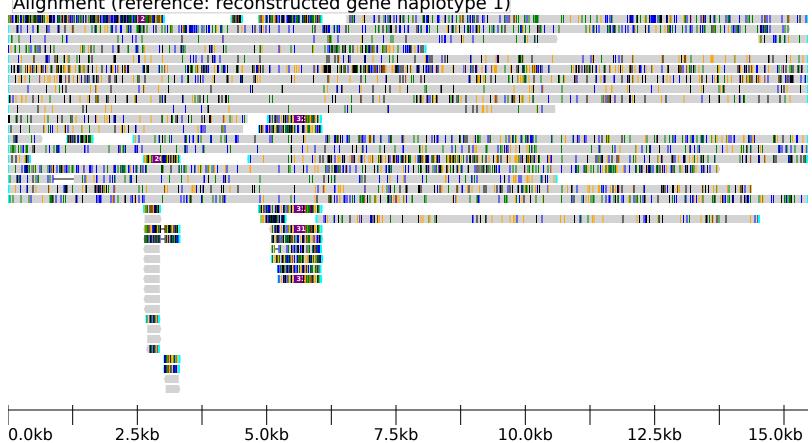
bam coverage

— 29

variants



Alignment (reference: reconstructed gene haplotype 1)



HLA-DRB4*01:03:01:01 (identity: 99.99%)

HLA-DRB4*01:03:01:03 (identity: 99.99%)

HLA-DRB4*01:03:01:18 (identity: 99.99%)

HLA-DRB4*01:152 (identity: 99.99%)

HLA-DRB4*01:03:01:02N (identity: 99.86%)

Resolution: 4th field Full typing result: HLA-DRB4*01:03:01:03 | HLA-DRB4*01:03:01:01 | HLA-DRB4*01:03:01:18 HLA-DRB4*01:152 bam coverage A a variants Alignment (reference: reconstructed gene haplotype 2) ııı. Ш 12 0.0kb 5.0kb 10.0kb 20.0kb 25.0kb 7.5kb 15.0kb HLA-DRB4*01:03:01:03 (identity: 22.61%) HLA-DRB4*01:03:01:01 (identity: 22.61%) ||||-|| HLA-DRB4*01:03:01:18 (identity: 22.61%) -|||||-||| HLA-DRB4*01:152 (identity: 22.61%) -|||||-|||

Sample: NA12877

Locus: HLA-DRB4

Sample: NA12877 Locus: HLA-DRB5 Resolution: 4th field Full typing result: HLA-DRB5*01:02:01 bam coverage **28** 19 variants Alignment (reference: reconstructed gene haplotype 1) 211 3) 24 11111-HILL IIII

0.0kb 2.5kb 3.75kb 5.0kb 6.25kb 7.5kb 8.75kb 10.0kb 12.5kb

HLA-DRB5*01:02:01 (identity: 68.72%)

Locus: HLA-E

Resolution: 4th field

Full typing result:

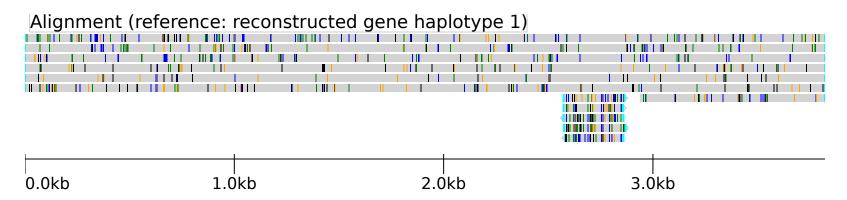
HLA-E*01:01:01:01

bam coverage

 \square_{7}^{10}

variants





HLA-E*01:01:01 (identity: 99.97%)

Locus: HLA-E

Resolution: 4th field

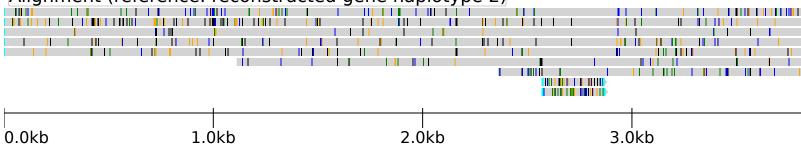
Full typing result:

HLA-E*01:03:02:02



variants

Alignment (reference: reconstructed gene haplotype 2)



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

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: 100.00%)

Locus: HLA-G

Resolution: 4th field

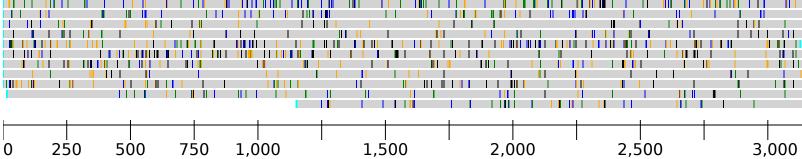
Full typing result:

HLA-G*01:01:01 | HLA-G*01:01:01:22

bam coverage

variants

Alignment (reference: reconstructed gene haplotype 1)



HLA-G*01:01:01 (identity: 99.94%)

HLA-G*01:01:01:22 (identity: 99.94%)

Locus: HLA-G

Resolution: 4th field

Full typing result:

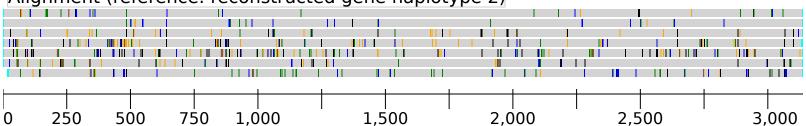
HLA-G*01:01:01:05

bam coverage

I gray

variants

Alignment (reference: reconstructed gene haplotype 2)



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

Locus: HLA-H

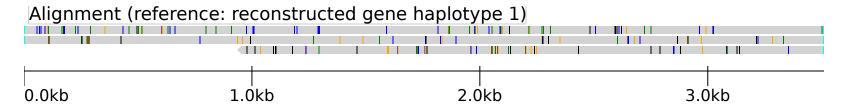
Resolution: 4th field

Full typing result:

HLA-H*02:04:01

bam coverage

variants



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

Locus: HLA-J

Resolution: 4th field

Full typing result:

HLA-J*01:01:01:04

bam coverage

variants

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

Locus: HLA-J

Resolution: 4th field

Full typing result:

HLA-J*01:01:05

bam coverage

variants

HLA-J*01:01:05 (identity: 100.00%)

Locus: HLA-K

Resolution: 4th field

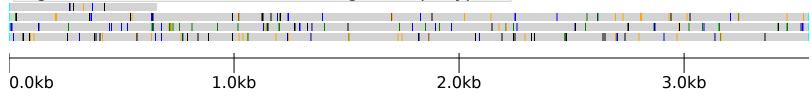
Full typing result:

HLA-K*01:02

bam coverage

variants

Alignment (reference: reconstructed gene haplotype 2)



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

Locus: HLA-K

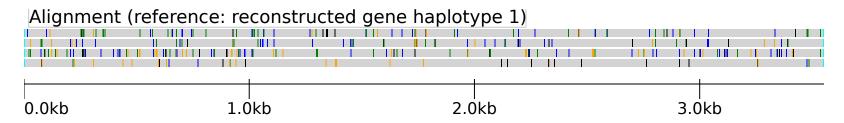
Resolution: 4th field

Full typing result:

HLA-K*01:01:01:01

bam coverage

variants



HLA-K*01:01:01:01 (identity: 99.97%)

Locus: HLA-L

Resolution: 4th field

Full typing result:

HLA-L*01:03

bam coverage

variants

Alignment (reference: reconstructed gene haplotype 1)

0.0kb 1.0kb 2.0kb 3.0kb

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

Locus: HLA-P

Resolution: 4th field

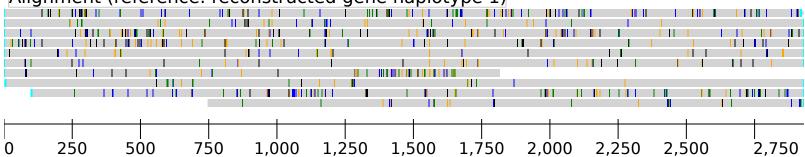
Full typing result:

HLA-P*02:01:01:01

bam coverage

variants

Alignment (reference: reconstructed gene haplotype 1)



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

Locus: HLA-P

Resolution: 4th field

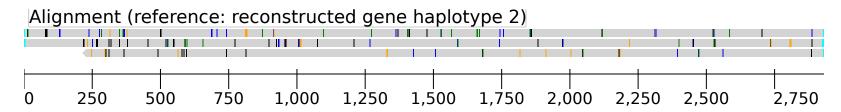
Full typing result:

HLA-P*02:01:01:02

bam coverage

 \square $\frac{3}{2}$

variants



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

Locus: HLA-T

Resolution: 4th field

Full typing result:

HLA-T*03:01:01:01

bam coverage



variants



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

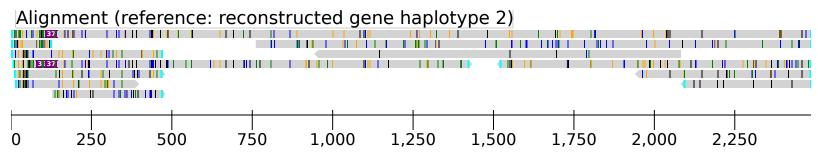
Locus: HLA-T

Resolution: 4th field

Full typing result:

HLA-T*03:01:01:02





HLA-T*03:01:01:02 (identity: 98.99%)

Locus: HLA-U

Resolution: 4th field

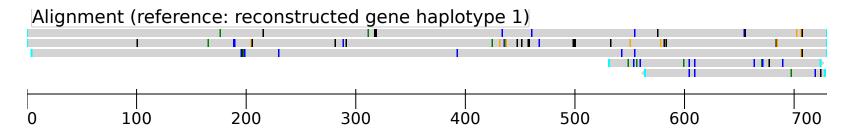
Full typing result:

HLA-U*01:04

bam coverage

−⁵₃

variants



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

Locus: HLA-U

Resolution: 4th field

Full typing result:

HLA-U*01:03

bam coverage

F 6

0

variants

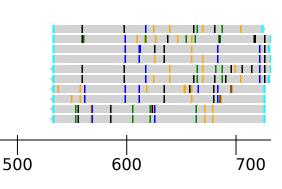


Alignment (reference: reconstructed gene haplotype 2)

300

400

200



HLA-U*01:03 (identity: 97.49%)

100

Locus: HLA-V

Resolution: 4th field

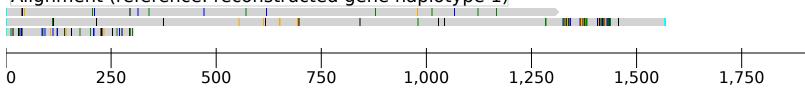
Full typing result:

HLA-V*01:01:01 HLA-V*01:01:04

bam coverage

variants

Alignment (reference: reconstructed gene haplotype 1)



HLA-V*01:01:01:01 (identity: 95.41%)

HLA-V*01:01:01:04 (identity: 95.41%)

Locus: HLA-W

Resolution: 4th field

Full typing result:

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

bam coverage

variants

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*03:01:01:02 (identity: 100.00%)

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

Locus: HLA-W

Resolution: 4th field

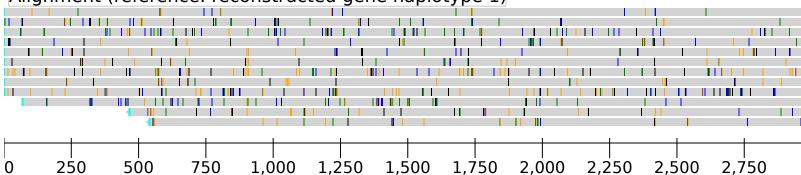
Full typing result:

HLA-W*01:01:01:05

bam coverage

variants

Alignment (reference: reconstructed gene haplotype 1)



HLA-W*01:01:01:05 (identity: 99.97%)

Sample: NA12877 Locus: MICA Resolution: 4th field Full typing result: MICA*008:01:02 bam coverage 52 34 variants Alignment (reference: reconstructed gene haplotype 1) Alignment (reference: reconstructed gene haplotype 1) 4 5 1(93 Ш Ш 0.0kb 7.5kb 2.5kb 5.0kb 10.0kb 12.5kb 15.0kb 17.5kb MICA*008:01:02 (identity: 99.98%)

Sample: NA12877 Locus: MICB Resolution: 4th field Full typing result: MICB*002:01:01 bam coverage 217 145 variants Alignment (reference: reconstructed gene haplotype 1) IIII 1 11111 0.0kb 2.5kb 5.0kb 3.75kb 6.25kb 7.5kb 8.75kb 10.0kb 12.5kb MICB*002:01:01 (identity: 99.80%)

Locus: MICB Resolution: 4th field Full typing result: MICB*005:02:01 bam coverage □ 134 90 variants Alignment (reference: reconstructed gene haplotype 2) 41111 11111 11111 IIIII IIIII 0.0kb 5.0kb 2.5kb 3.75kb 6.25kb 7.5kb 12.5kb 8.75kb 10.0kb MICB*005:02:01 (identity: 99.54%)

Sample: NA12877

Locus: TAP1

Resolution: 4th field

Full typing result:

TAP1*01:01:01:01

bam coverage



Alignment (reference: reconstructed gene haplotype 1)

O.0kb 2.0kb 3.0kb 4.0kb 5.0kb 6.0kb 7.0kb 8.0kb 9.0kb

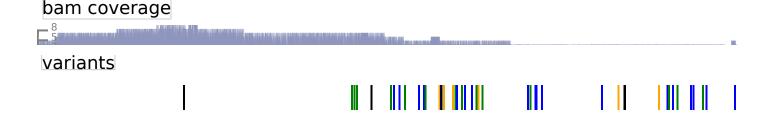
TAP1*01:01:01:01 (identity: 100.00%)

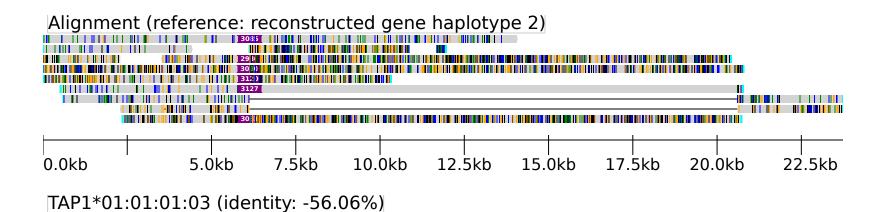
Locus: TAP1

Resolution: 4th field

Full typing result:

TAP1*01:01:01:03





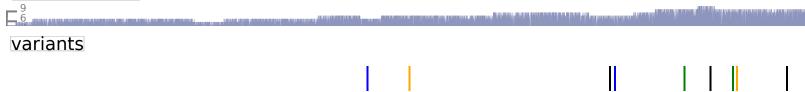
Locus: TAP2

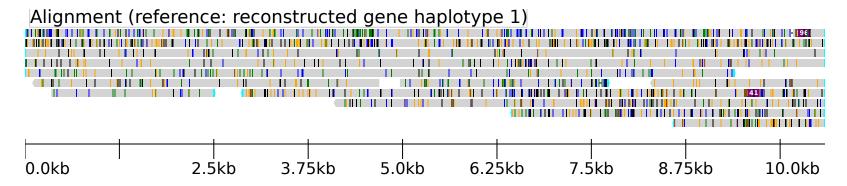
Resolution: 4th field

Full typing result:

TAP2*01:04:01:01







TAP2*01:04:01:01 (identity: 99.99%)

Locus: TAP2

Resolution: 4th field

Full typing result:

TAP2*02:01:02:01

bam coverage

TAP2*02:01:02:01 (identity: 99.89%)

variants

Alignment (reference: reconstructed gene haplotype 2)

0.0kb 2.5kb 3.75kb 5.0kb 6.25kb 7.5kb 8.75kb 10.0kb