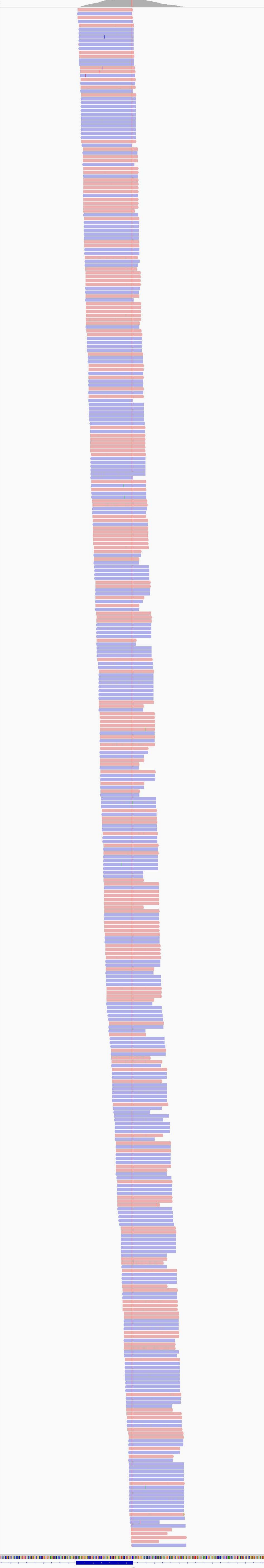


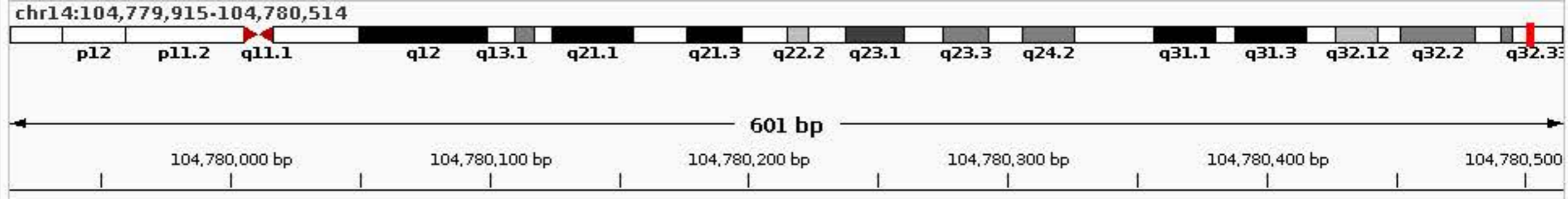
SL_0009-LT_3-B_mut.bam Coverage

SL_0009-LT_3-B_mut.bam



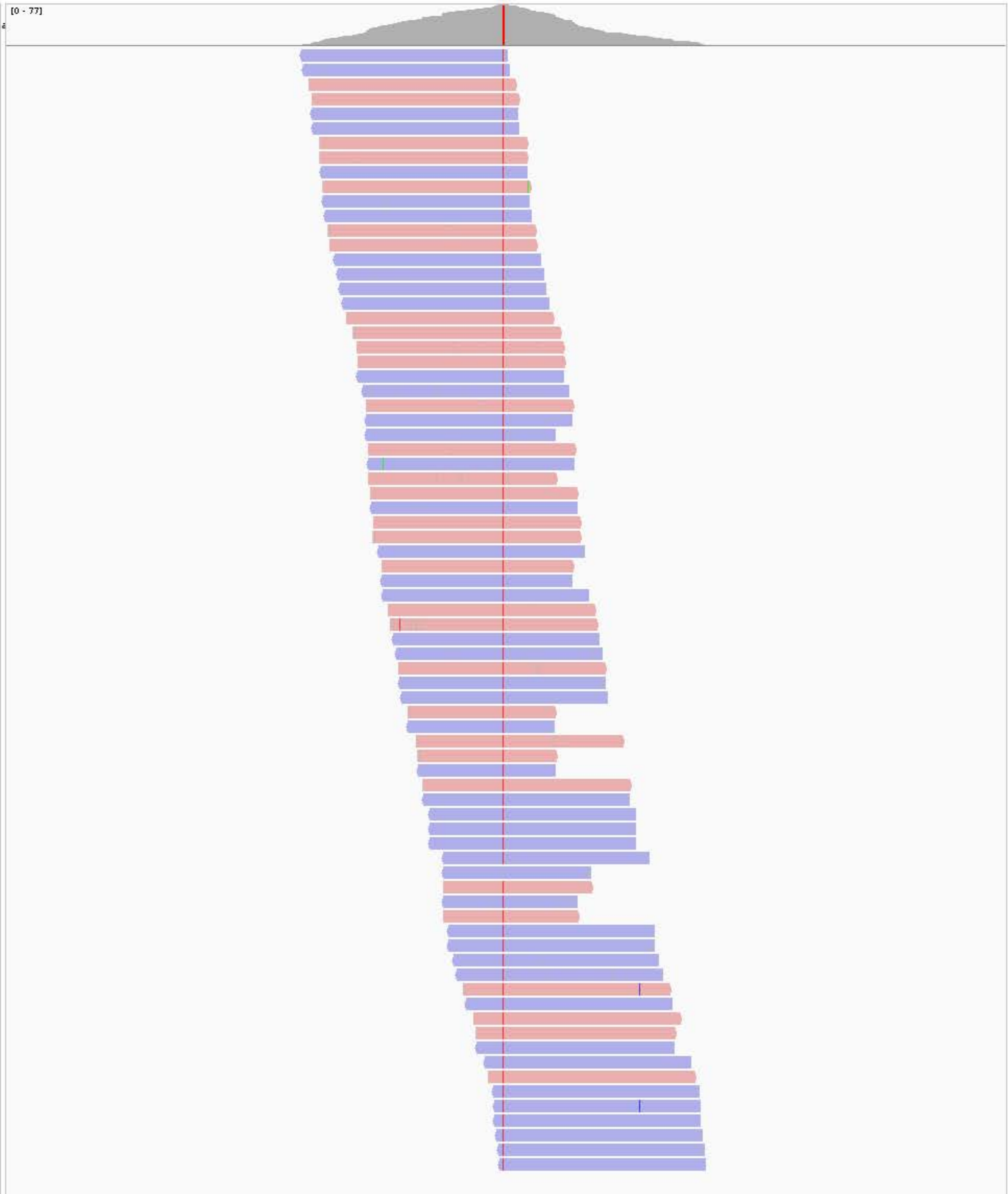
Sequence





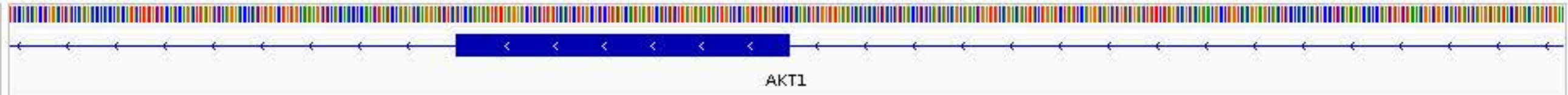
SL_0009-LT_9-B_mut.bam Coverage

SL_0009-LT_9-B_mut.bam

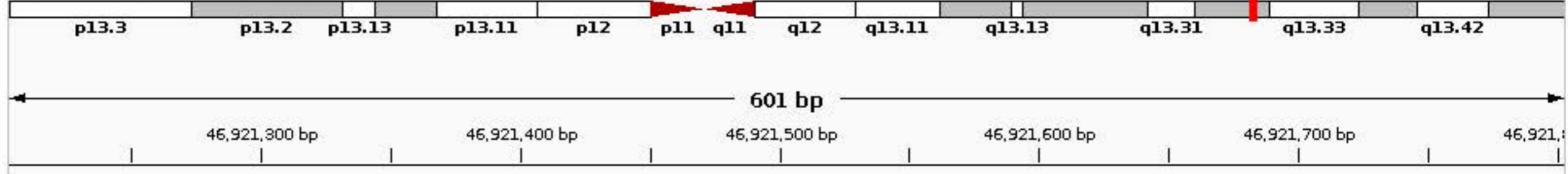


Sequence →

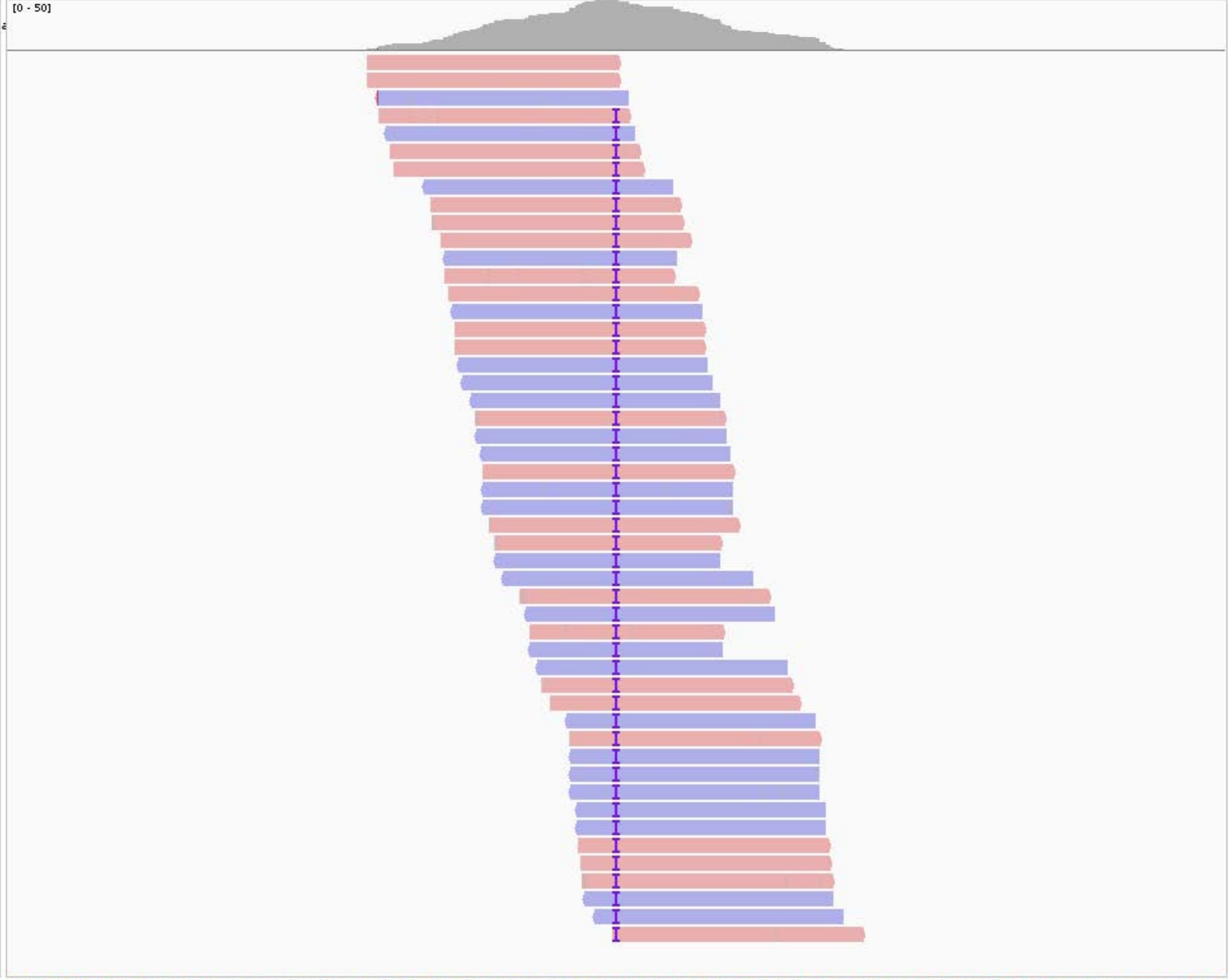
Refseq Genes



chr19:46,921,203-46,921,802



SL_0011-LT_6-B_mut.bam Coverage



SL_0011-LT_6-B_mut.bam

Sequence

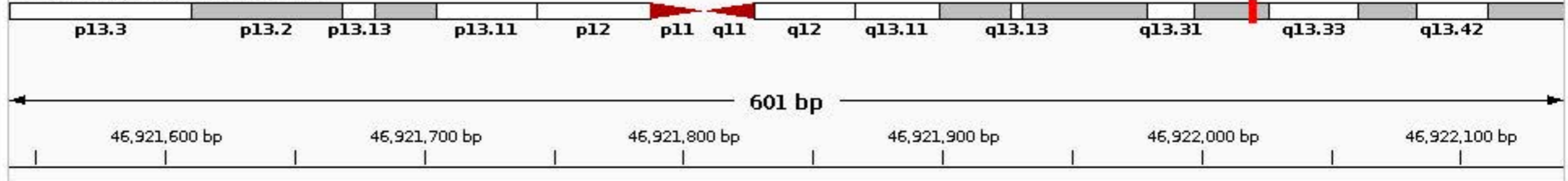


Refseq Genes

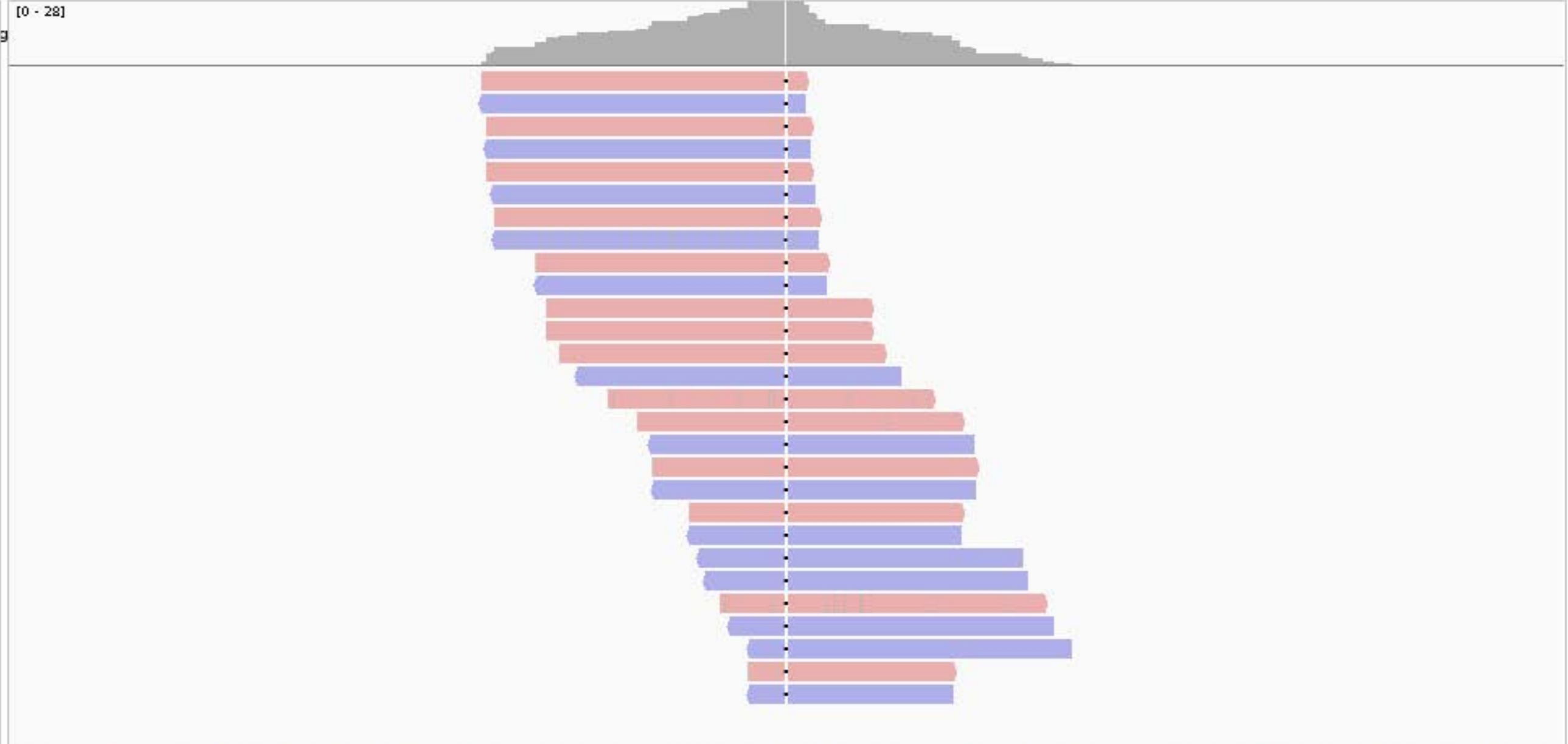


ARHGAP35

chr19:46,921,540-46,922,139



SL_1010-N6-B_mut.bam Coverage



SL_1010-N6-B_mut.bam

Sequence

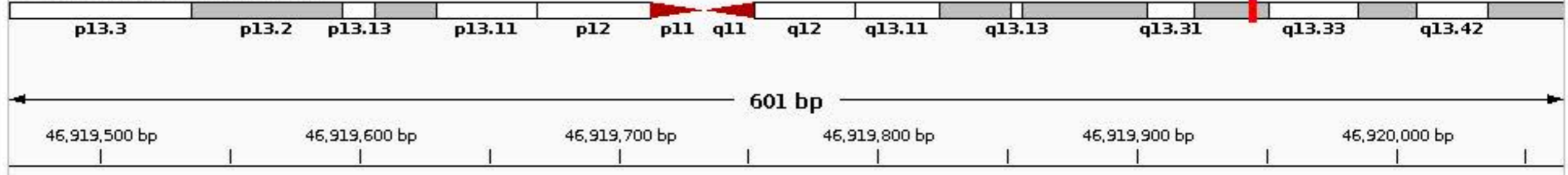


Refseq Genes



ARHGAP35

chr19:46,919,465-46,920,064



SL_0006_RN_4-B_mut.bam Cover



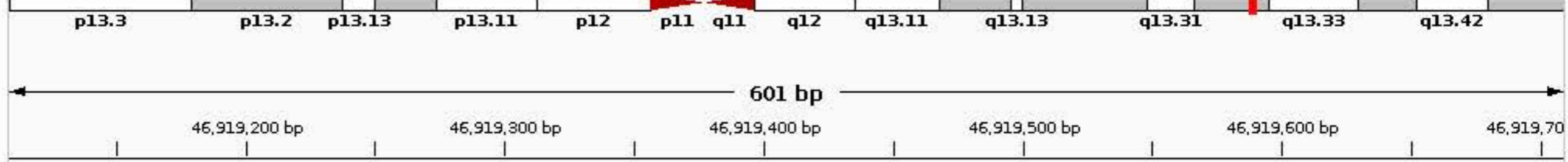
SL_0006_RN_4-B_mut.bam

Sequence →
Refseq Genes



ARHGAP35

chr19:46,919,109-46,919,708

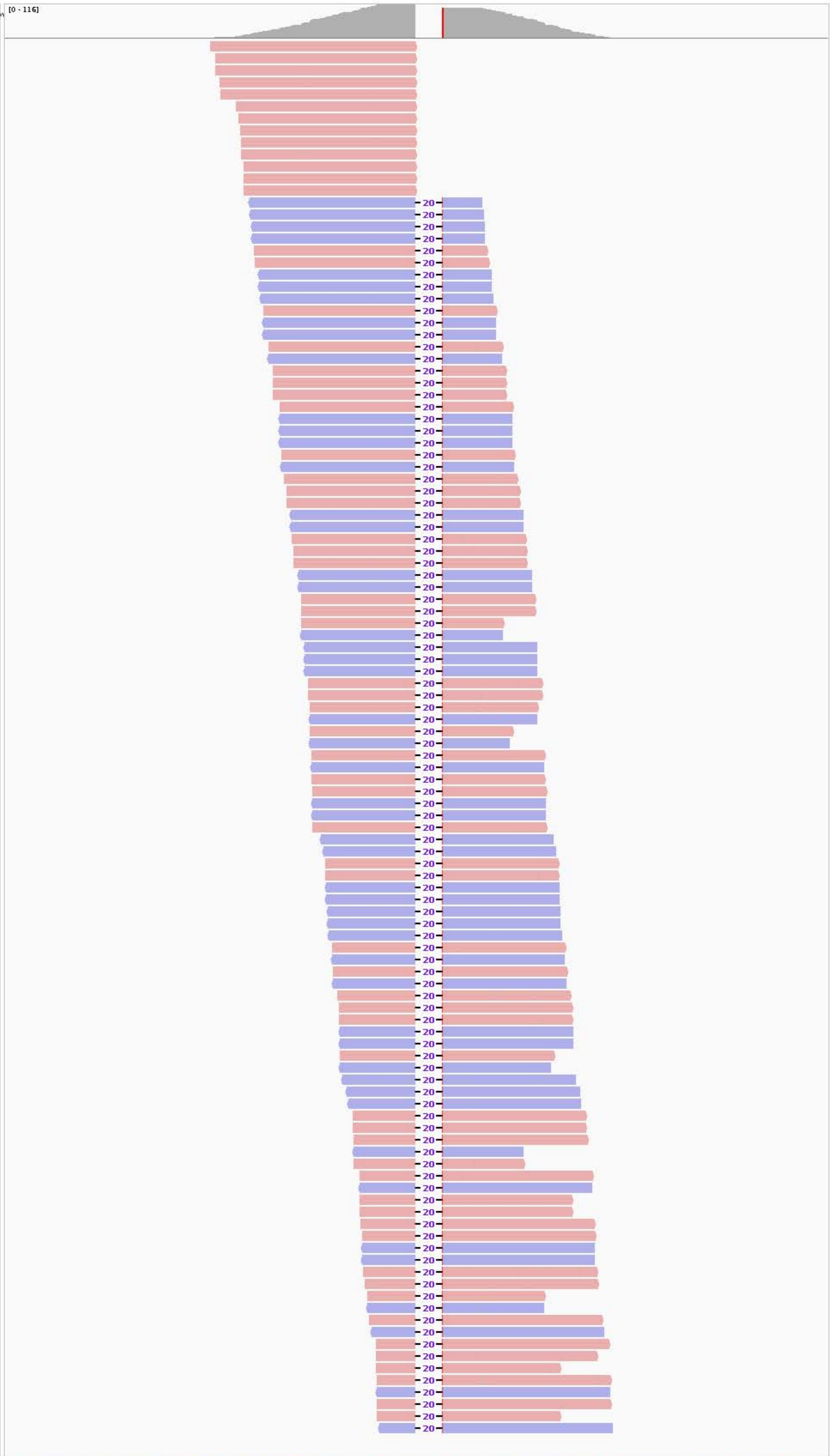


SL_0003_R_FFPE-B_mut.bam Coverage

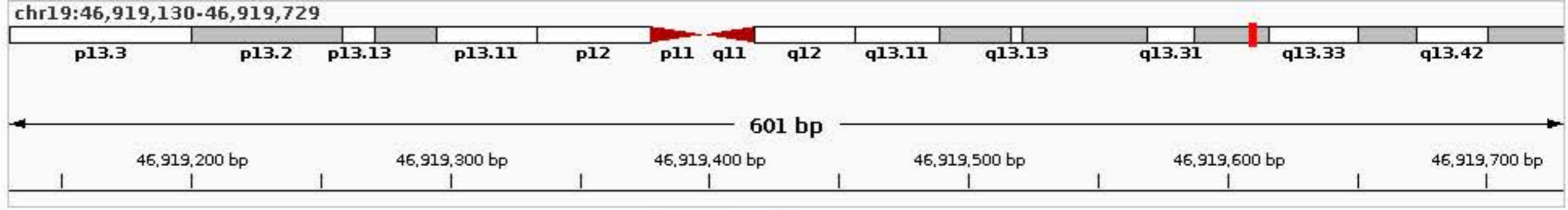
SL_0003_R_FFPE-B_mut.bam

Sequence →

Refseq Genes

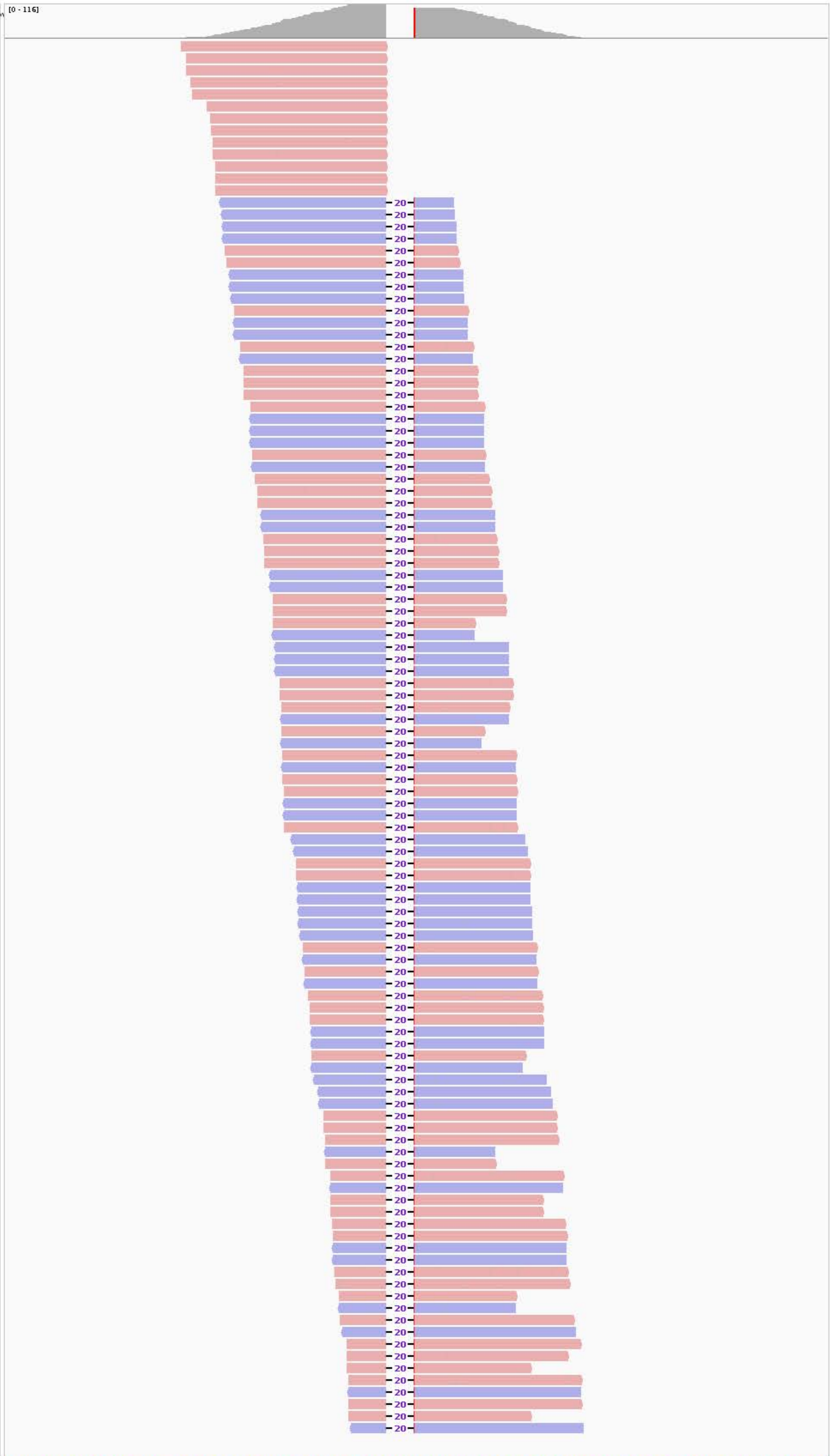


ARHGAP35



SL_0003_R_FFPE-B_mut.bam Coverage

SL_0003_R_FFPE-B_mut.bam

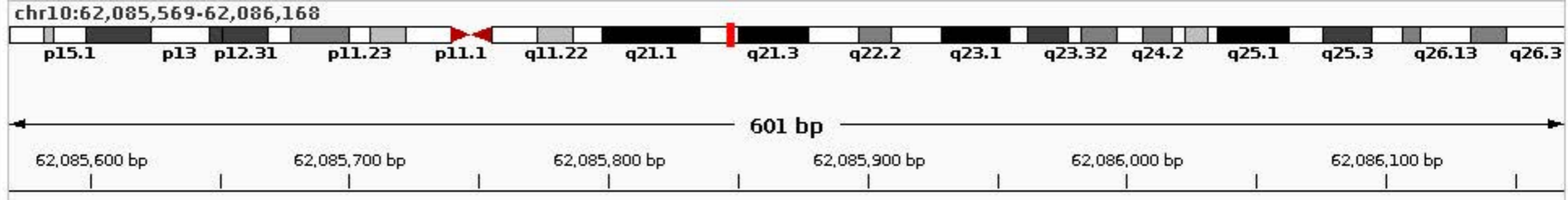


Sequence

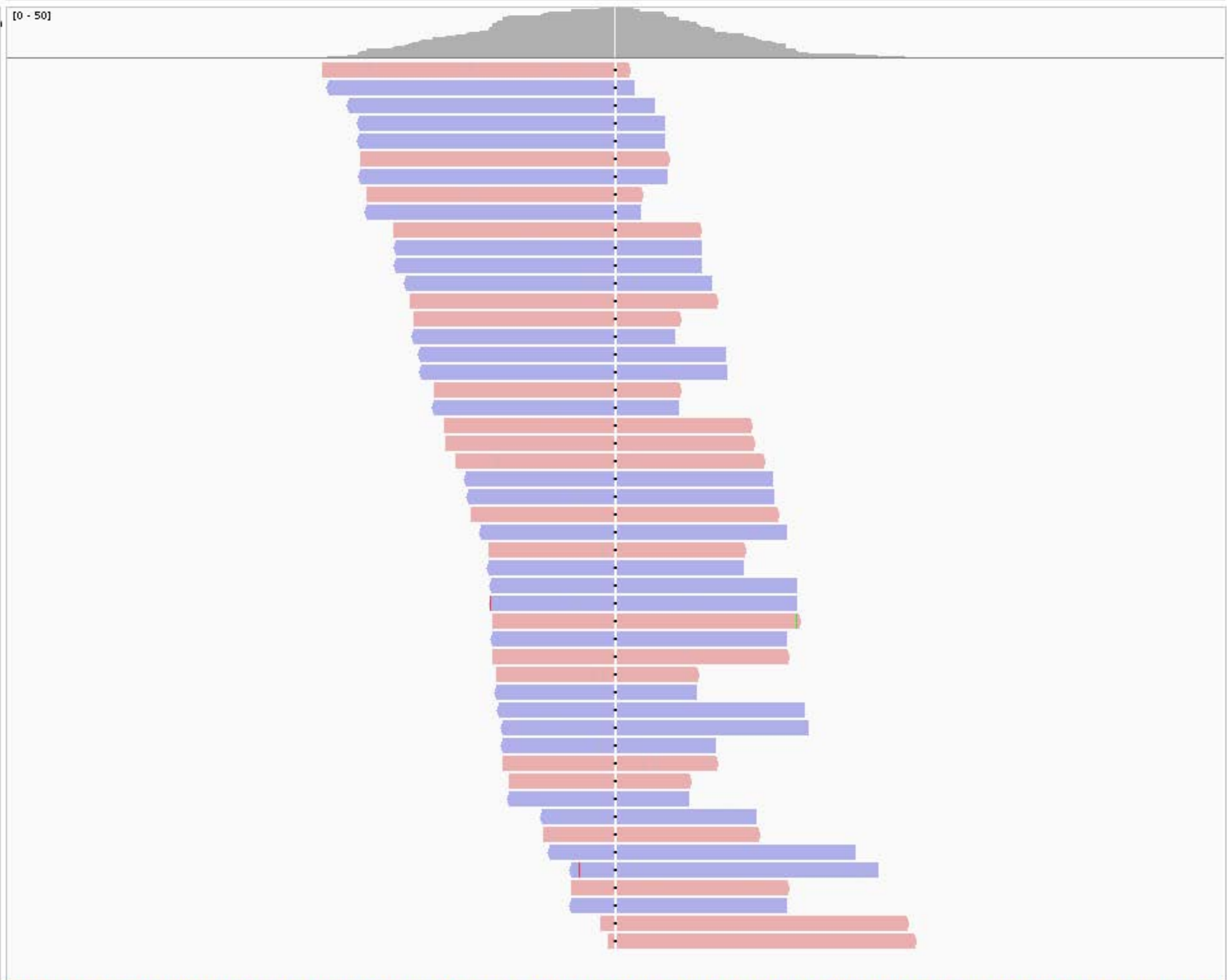
Refseq Genes



ARHGAP35



SL_0002_R_FFPE_57-B_mut.bam
range



SL_0002_R_FFPE_57-B_mut.ba
m

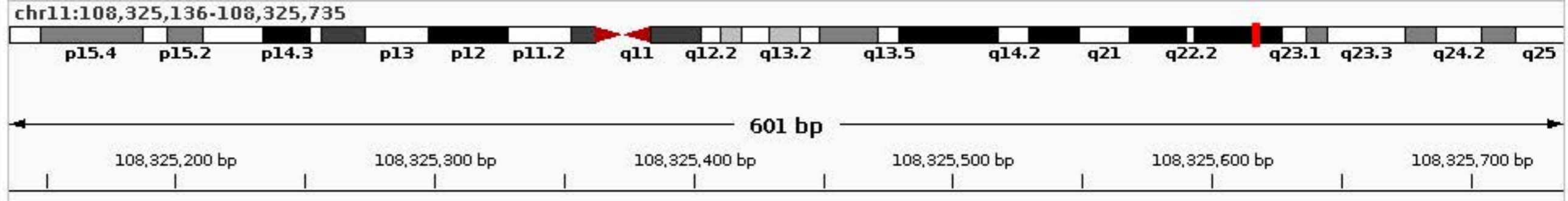
Sequence →



Refseq Genes

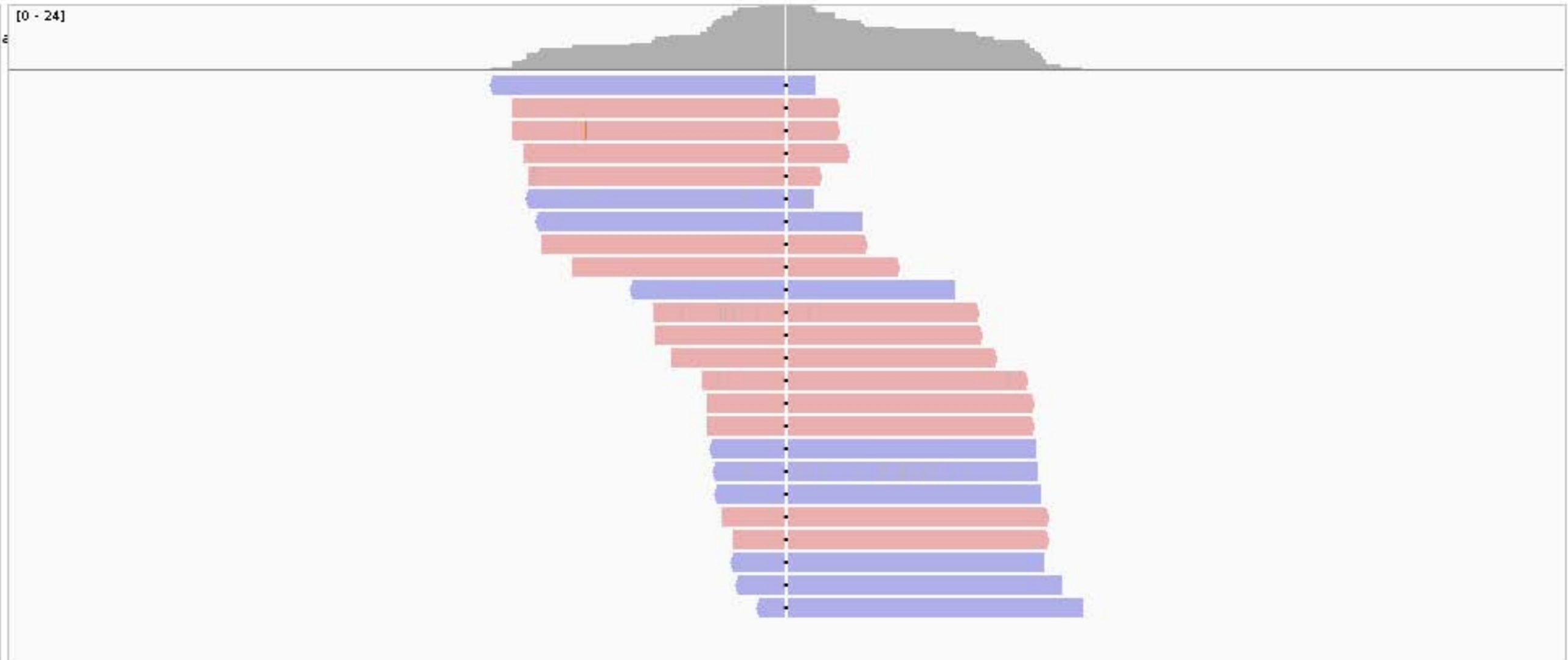


ARID5B



SL_0011-LT_2-B_mut.bam Coverage

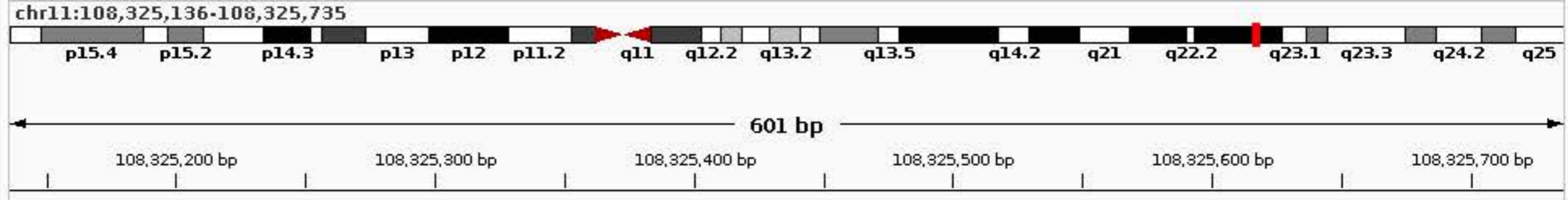
SL_0011-LT_2-B_mut.bam



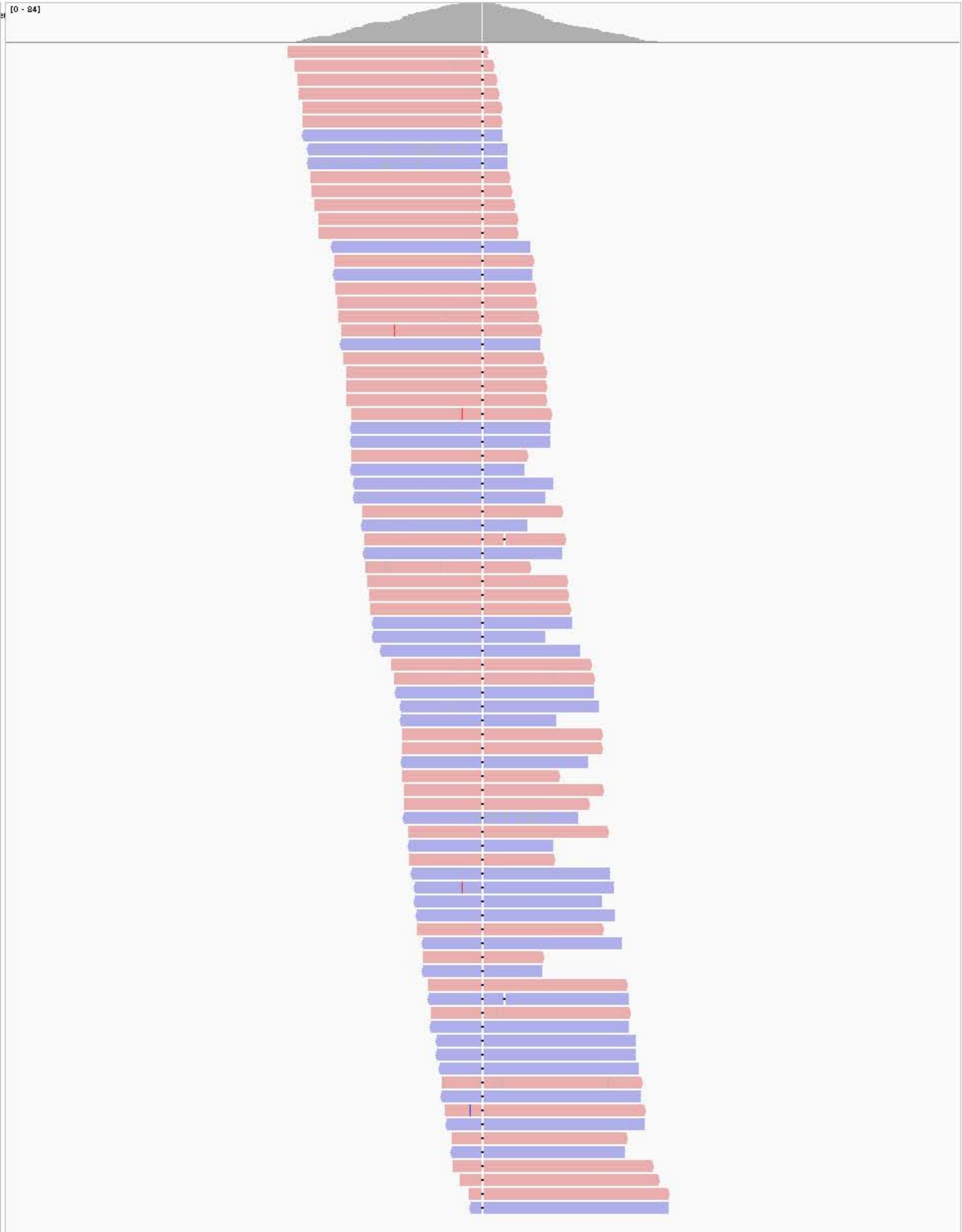
Sequence

Refseq Genes





SL_0011-LT_11-B_mut.bam Cov



SL_0011-LT_11-B_mut.bam

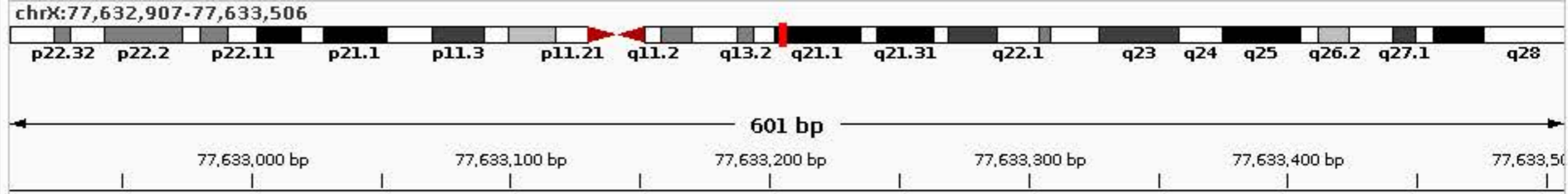
Sequence



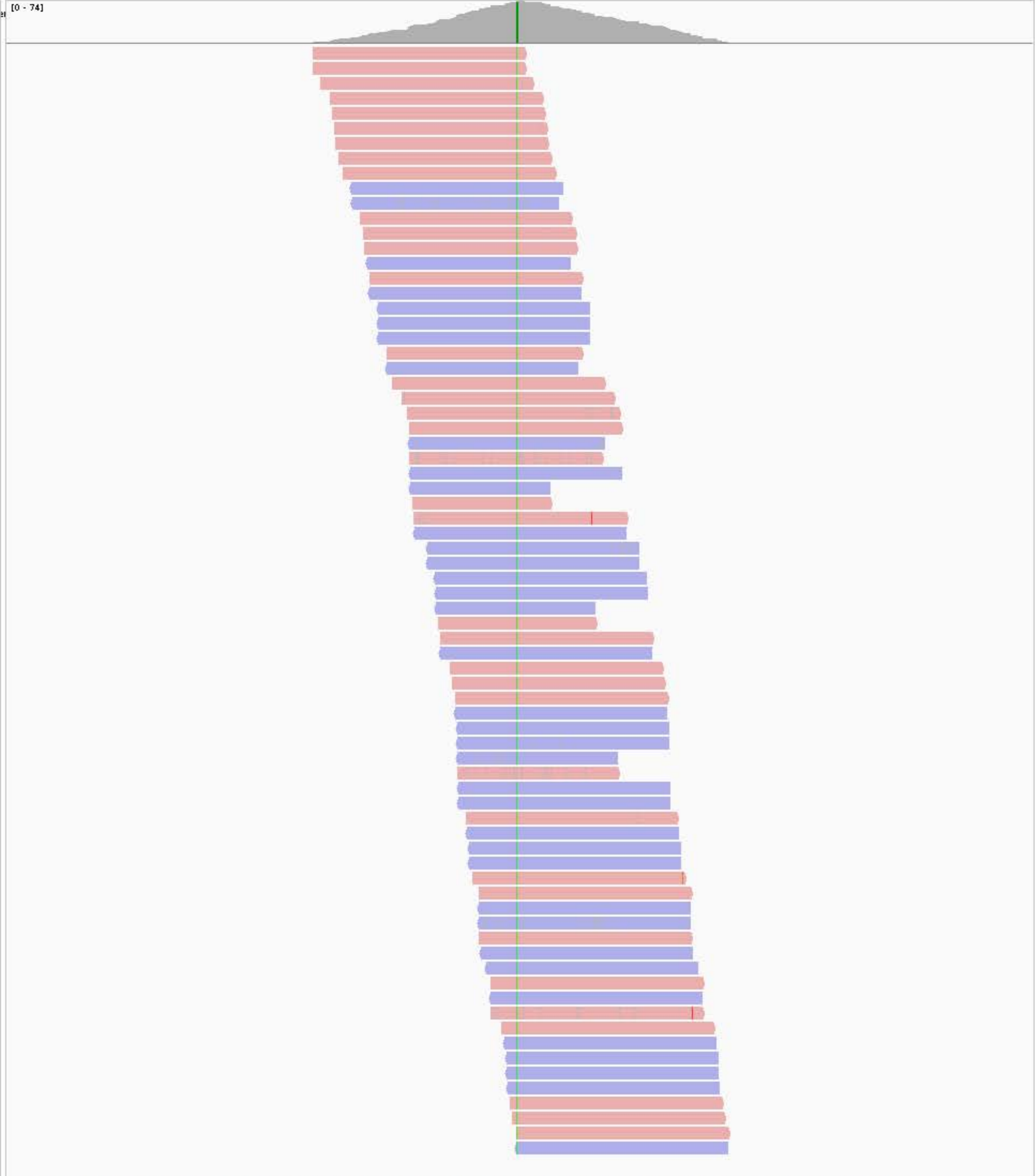
Refseq Genes



ATM



SL_0008-LT_15-B_mut.bam Coverage

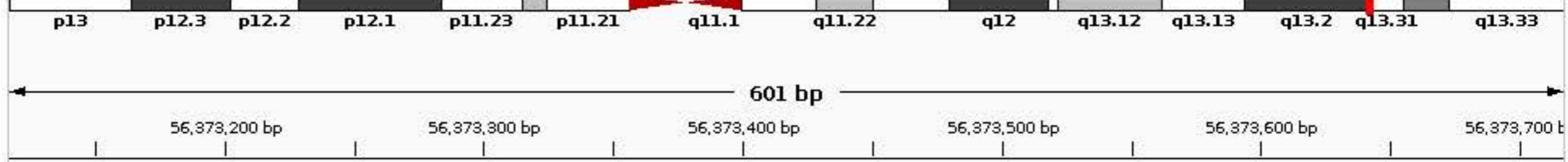


SL_0008-LT_15-B_mut.bam

Sequence →
Refseq Genes



chr20:56,373,117-56,373,716

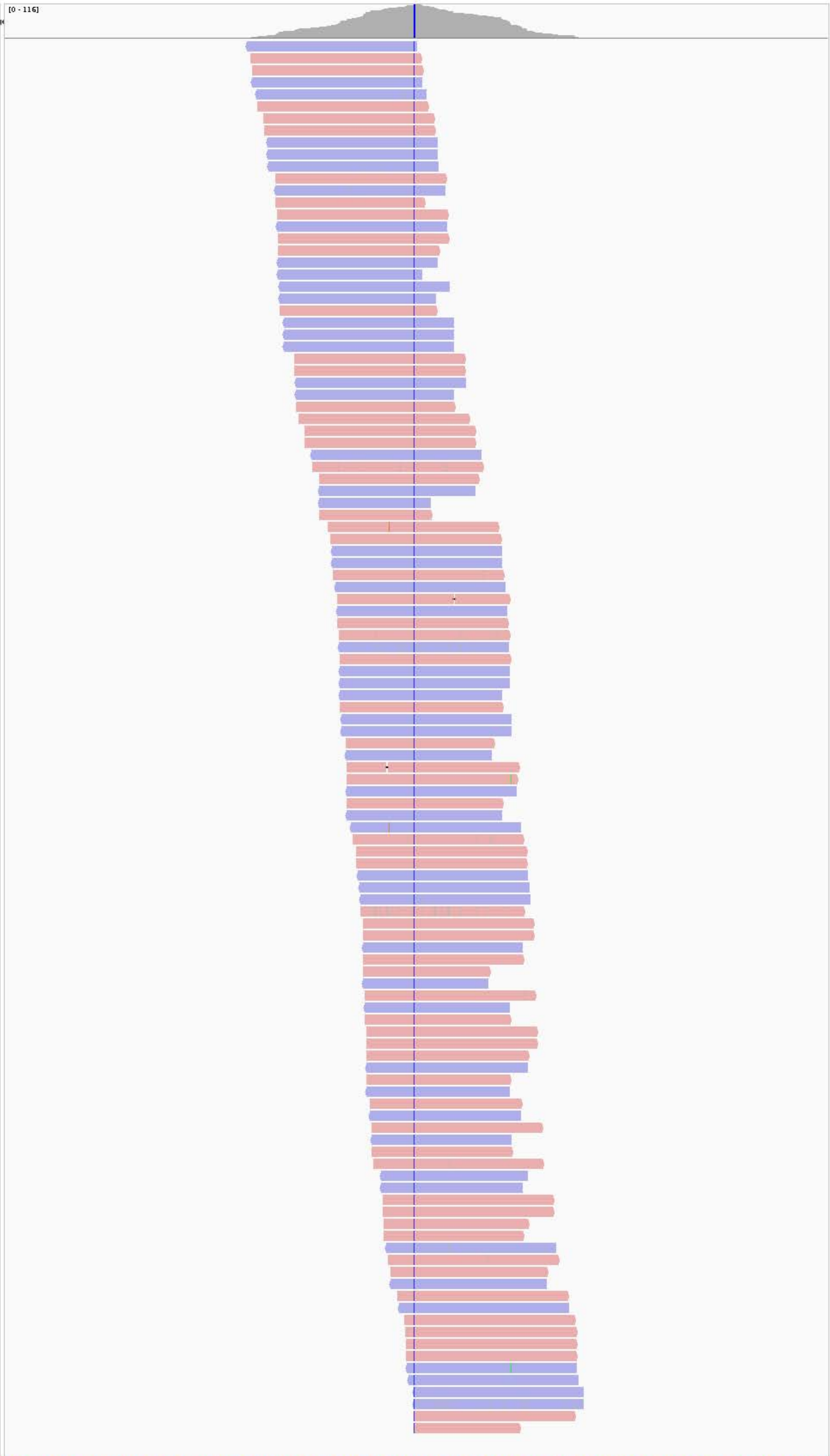


SL_1010-T1-B_mut.bam Coverage

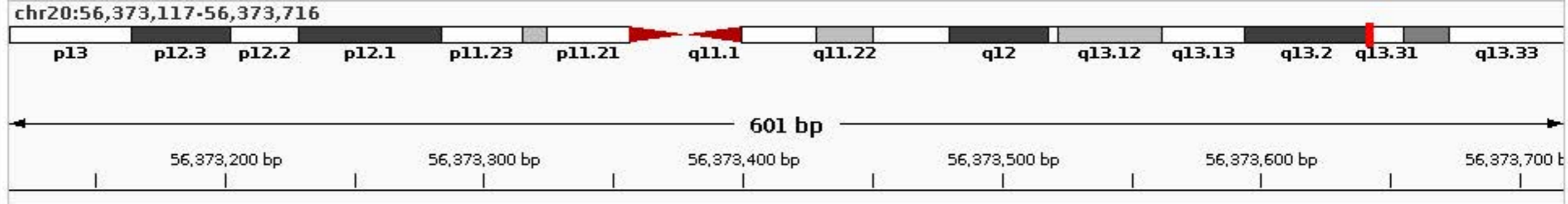
SL_1010-T1-B_mut.bam

Sequence →

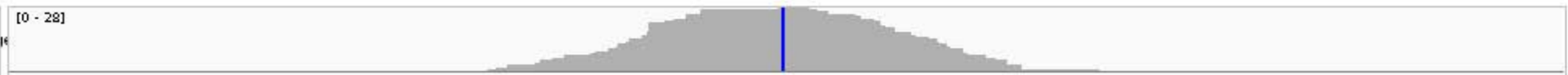
Refseq Genes



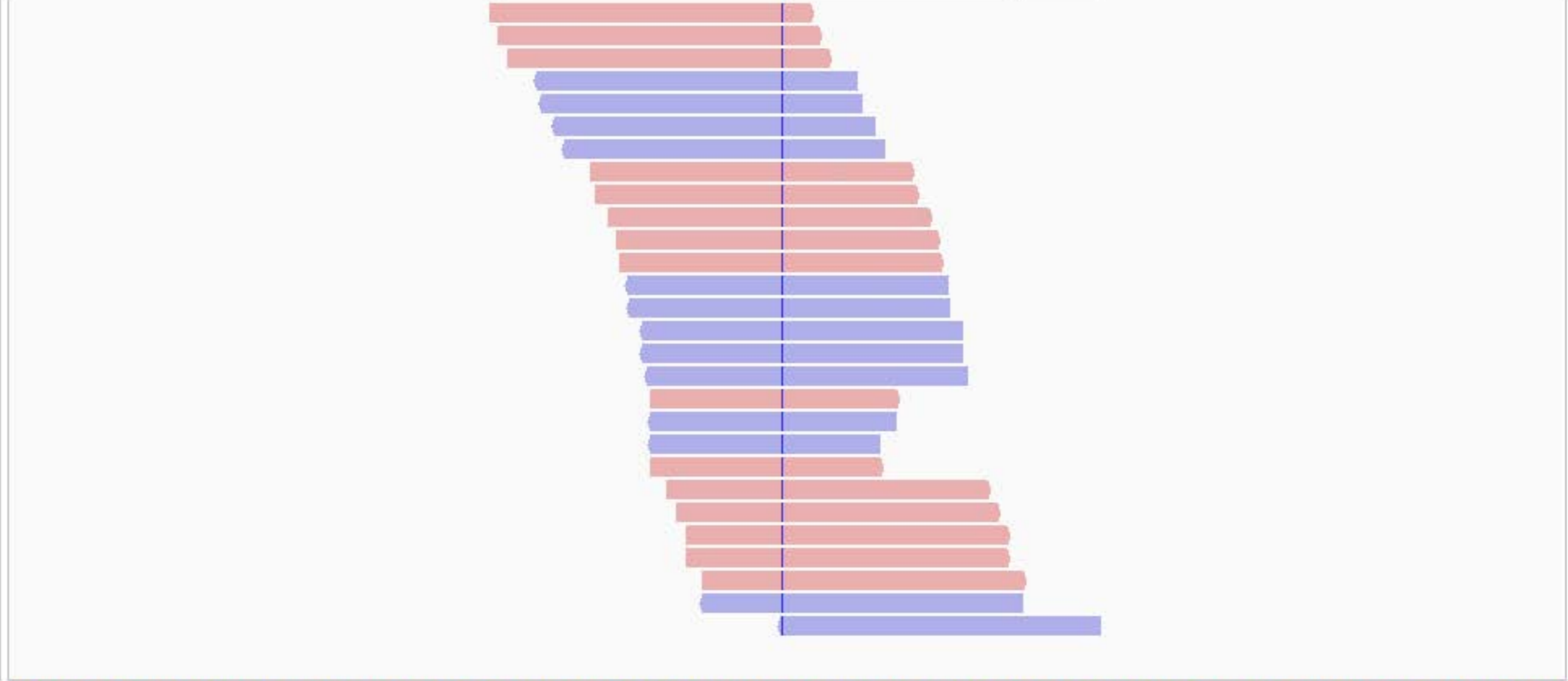
AURKA



SL_1010-T2-B_mut.bam Coverage



SL_1010-T2-B_mut.bam

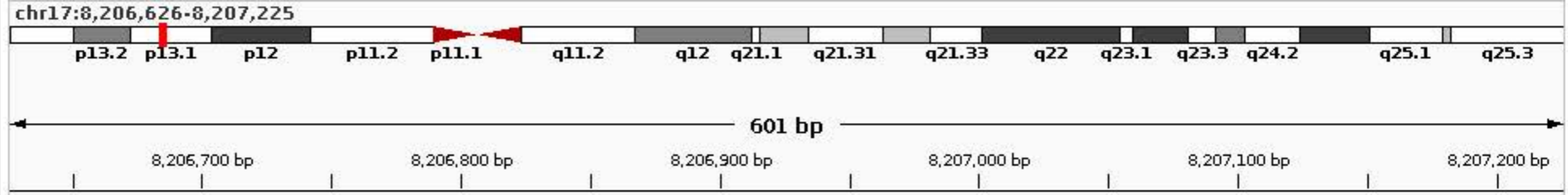


Sequence →

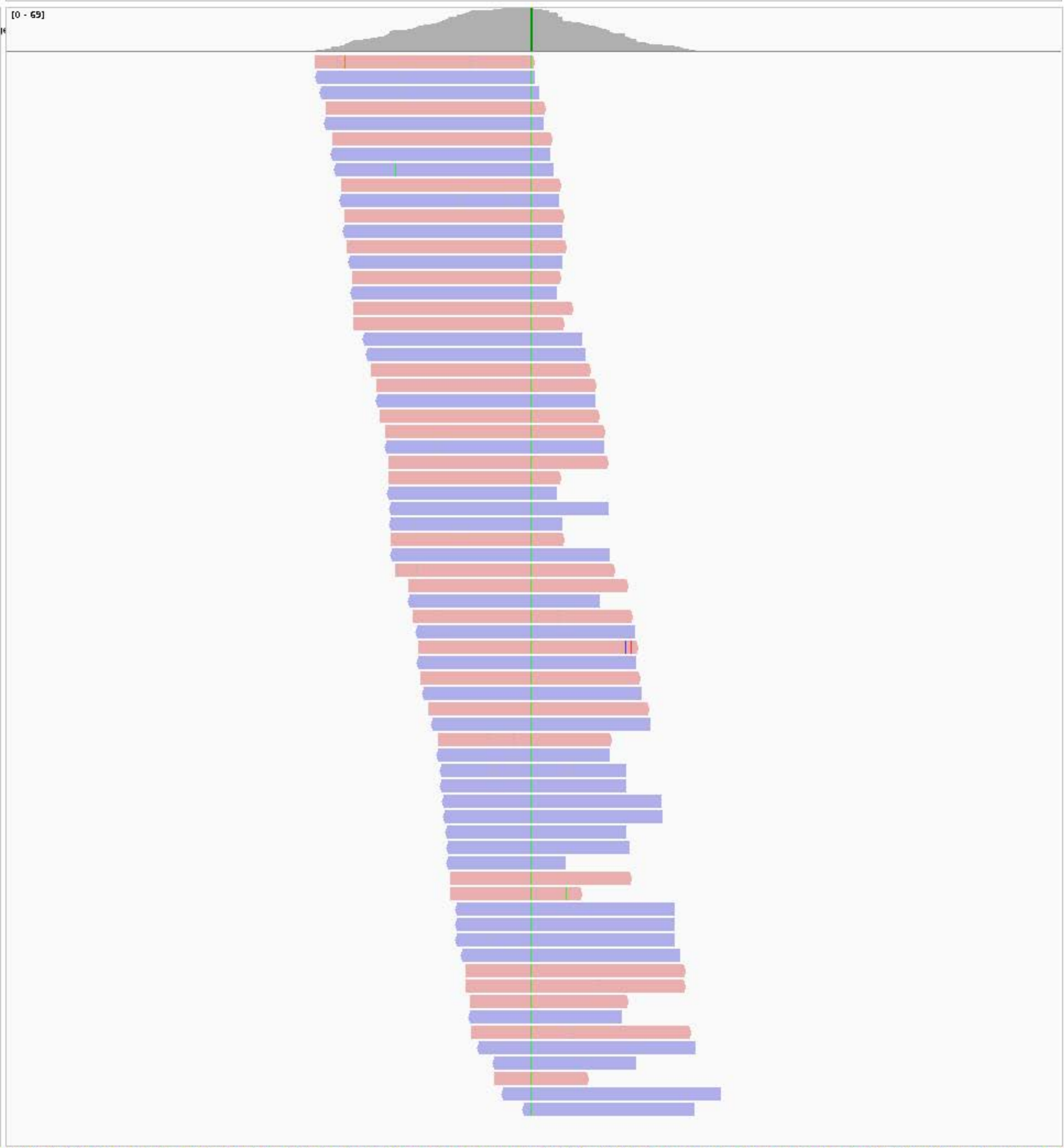


Refseq Genes





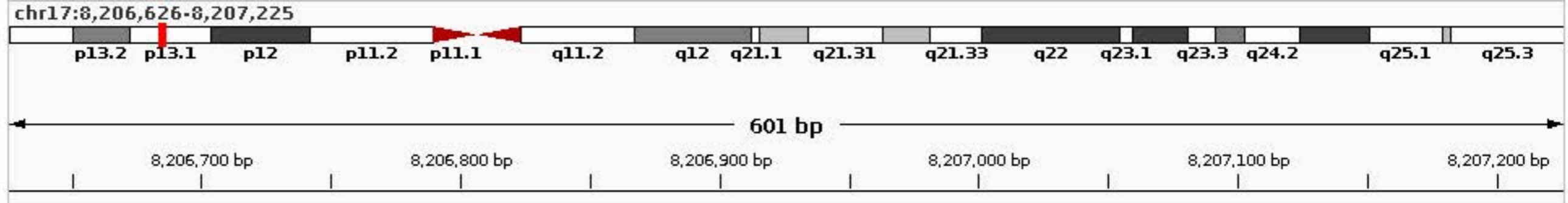
SL_1010-T1-B_mut.bam Coverage



SL_1010-T1-B_mut.bam

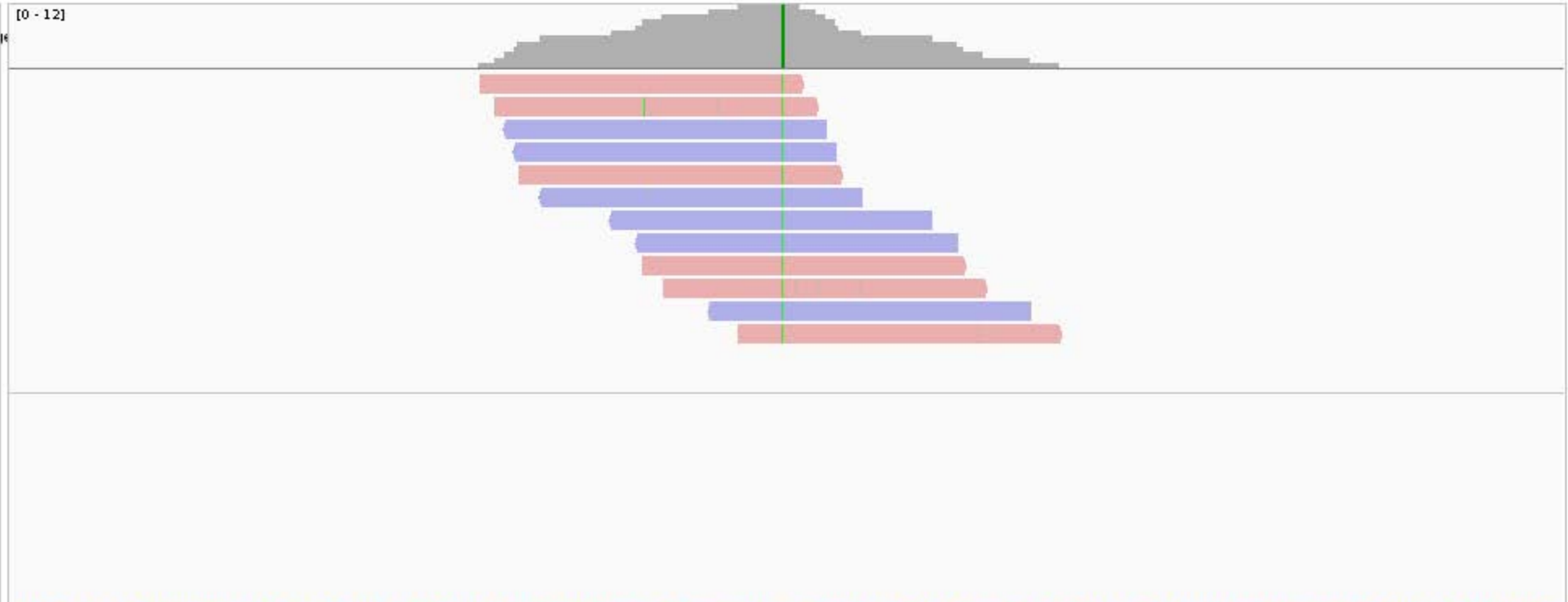
Sequence →
Refseq Genes





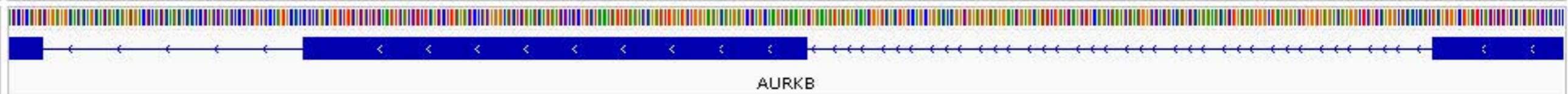
SL_1010-T2-B_mut.bam Coverage

SL_1010-T2-B_mut.bam

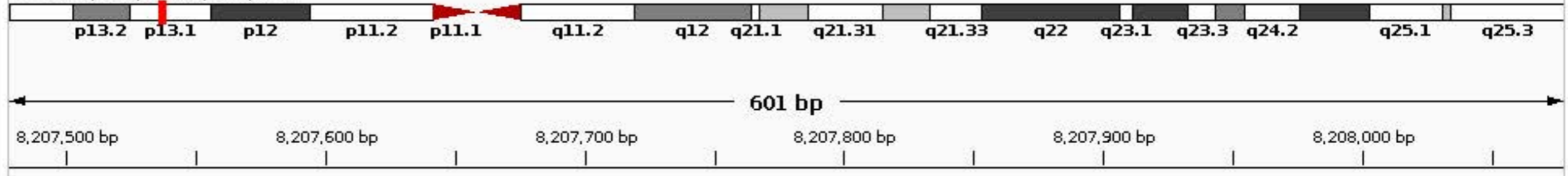


Sequence

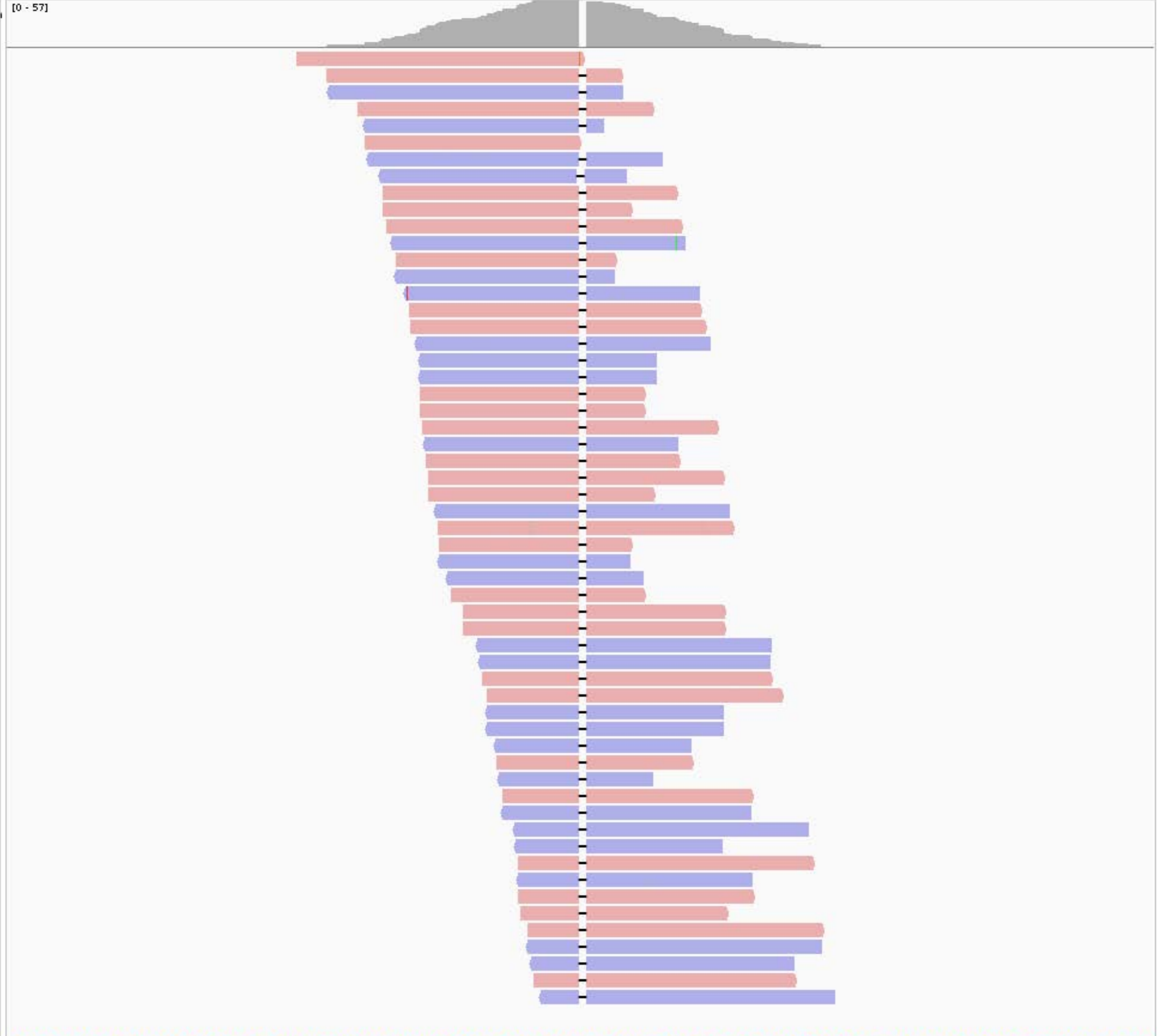
Refseq Genes



chr17:8,207,478-8,208,077



SL_1011_R_FFPE_31-B_mut.bam
rage



SL_1011_R_FFPE_31-B_mut.ba
m

Sequence →

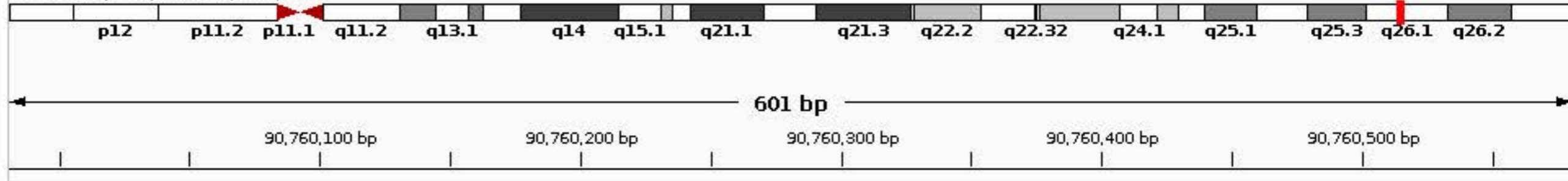


Refseq Genes

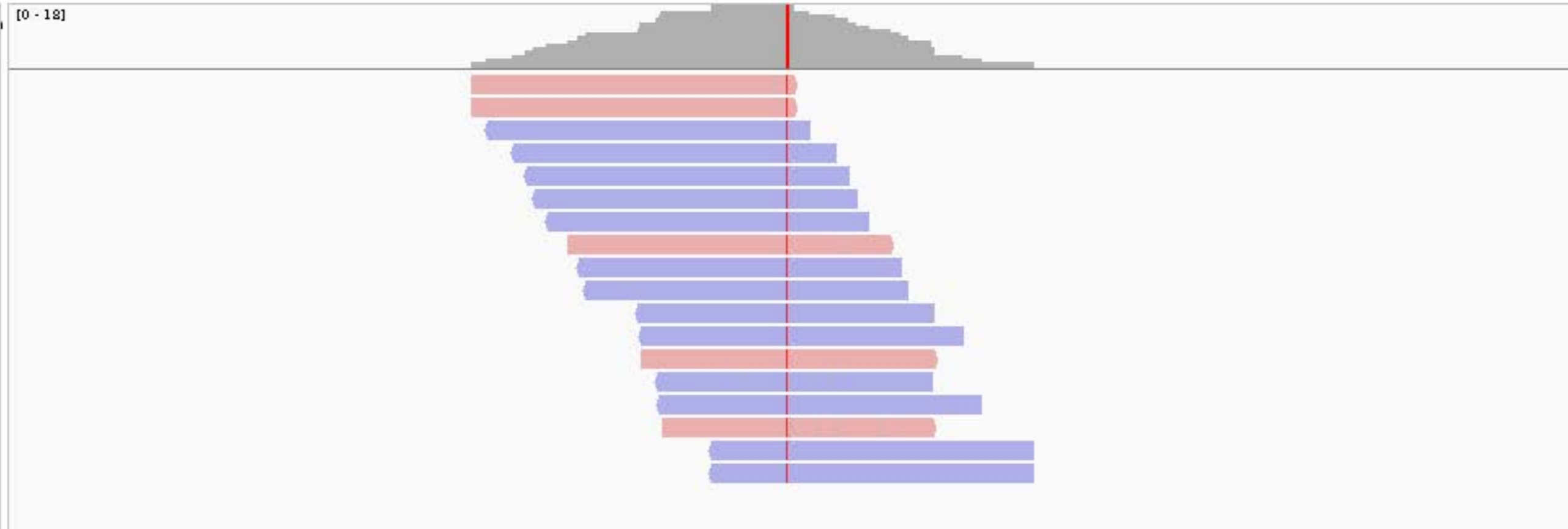


AURKB

chr15:90,759,981-90,760,580



SL_1008_R_51_FFPE-B_mut.bam
range



SL_1008_R_51_FFPE-B_mut.ba
m

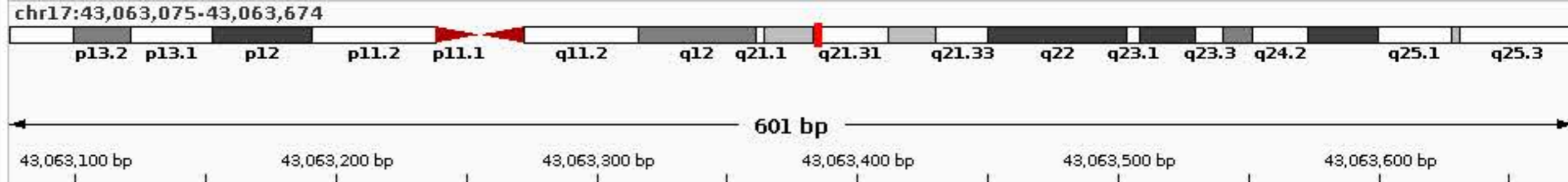
Sequence →



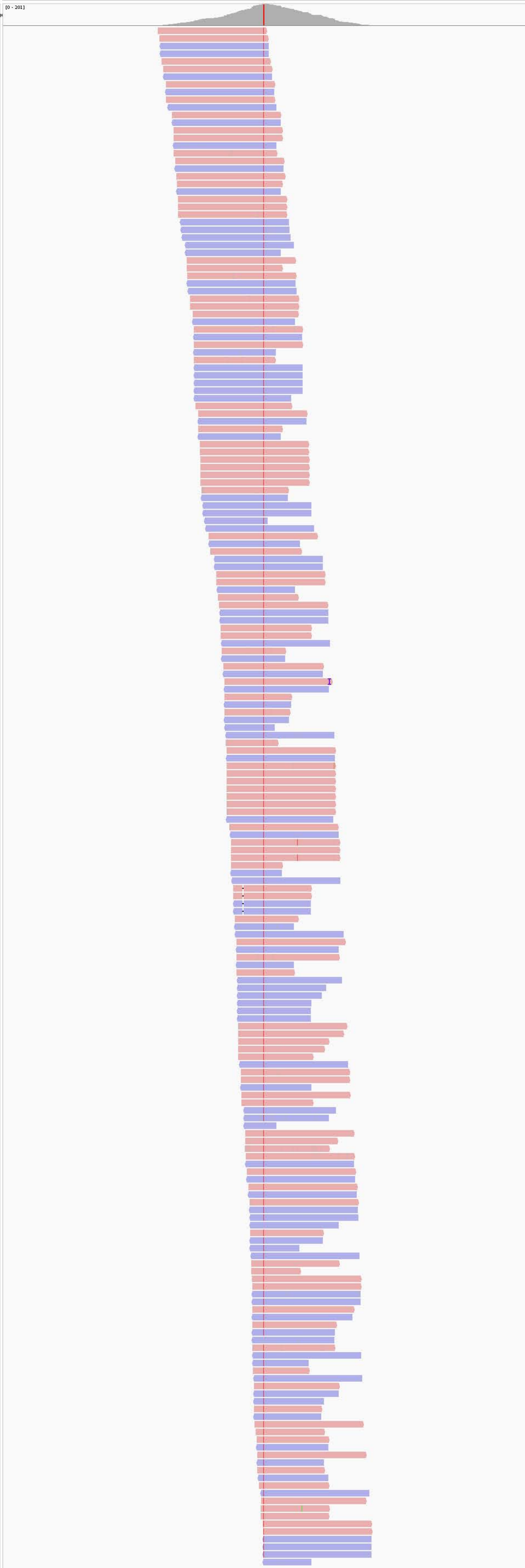
Refseq Genes



BLM



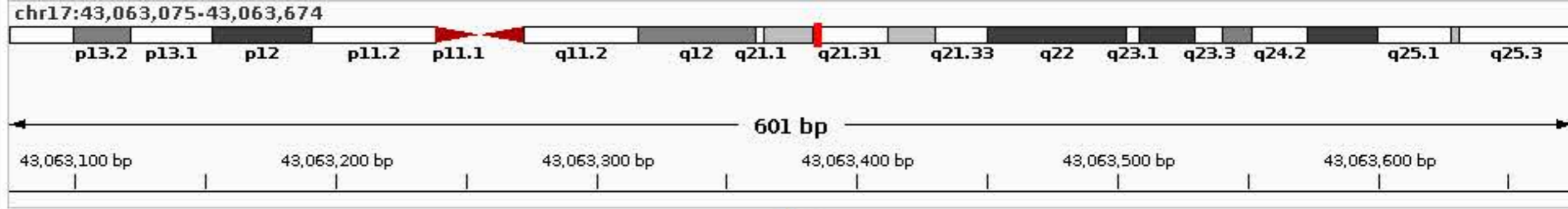
SL_1011-T1-B_mut.bam Coverage



SL_1011-T1-B_mut.bam

Sequence
Refseq Genes



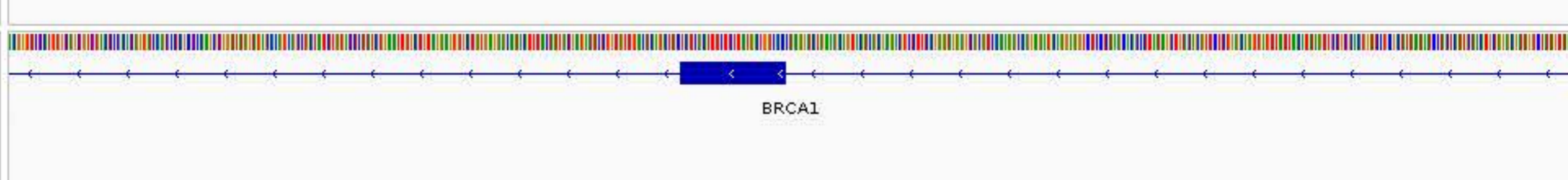
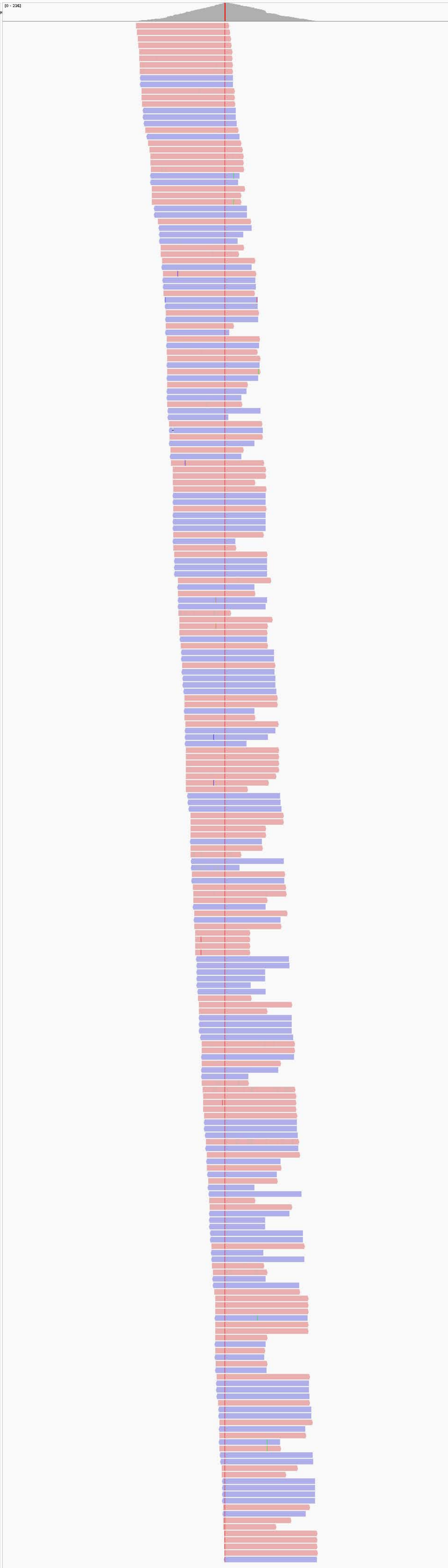


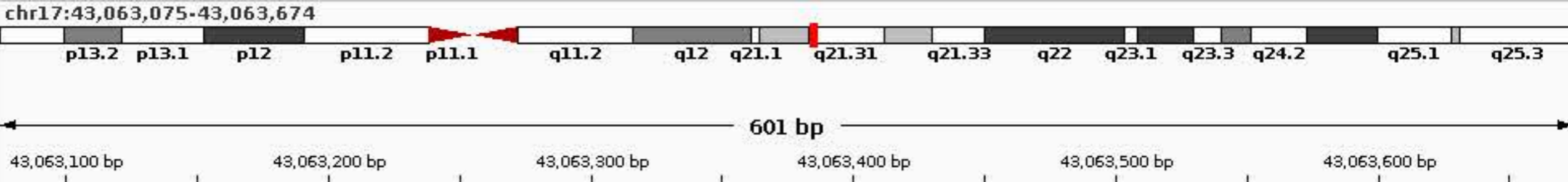
SL_1011-T2-B_mut.bam Coverage

SL_1011-T2-B_mut.bam

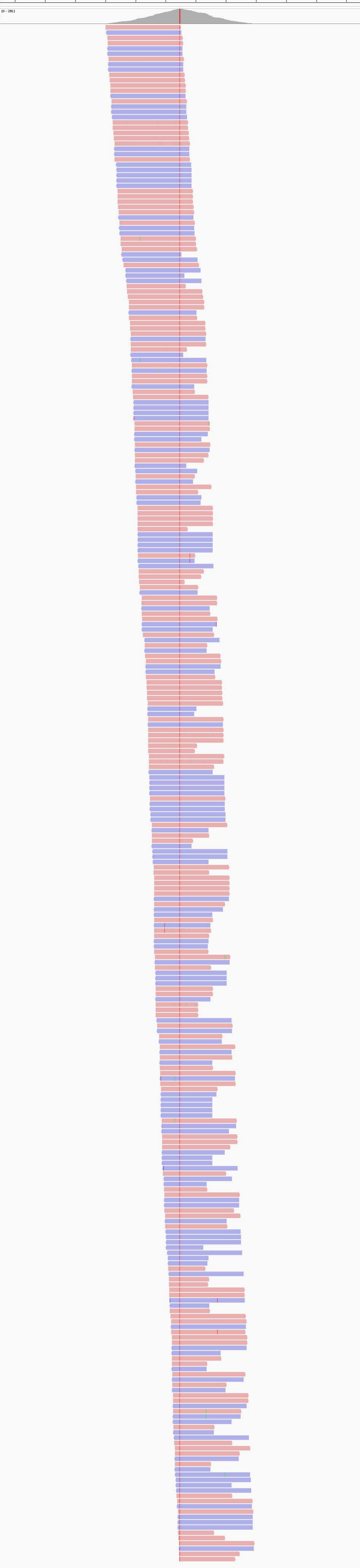
Sequence

Refseq Genes



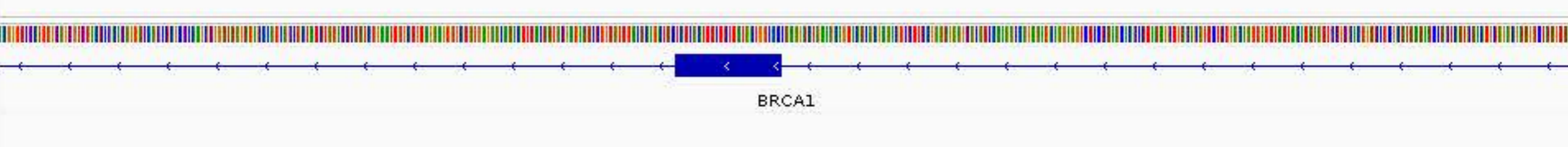


SL_1011-T9-B_mut.bam Coverage

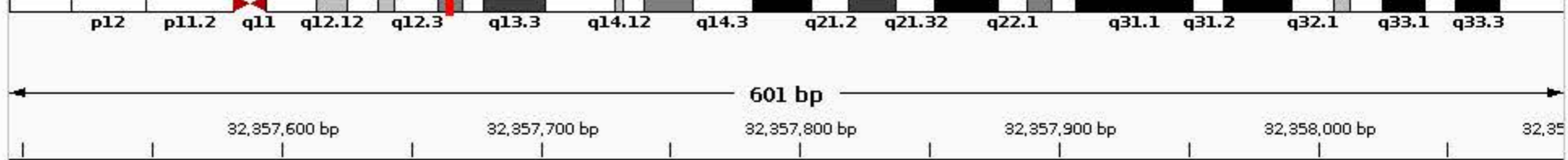


SL_1011-T9-B_mut.bam

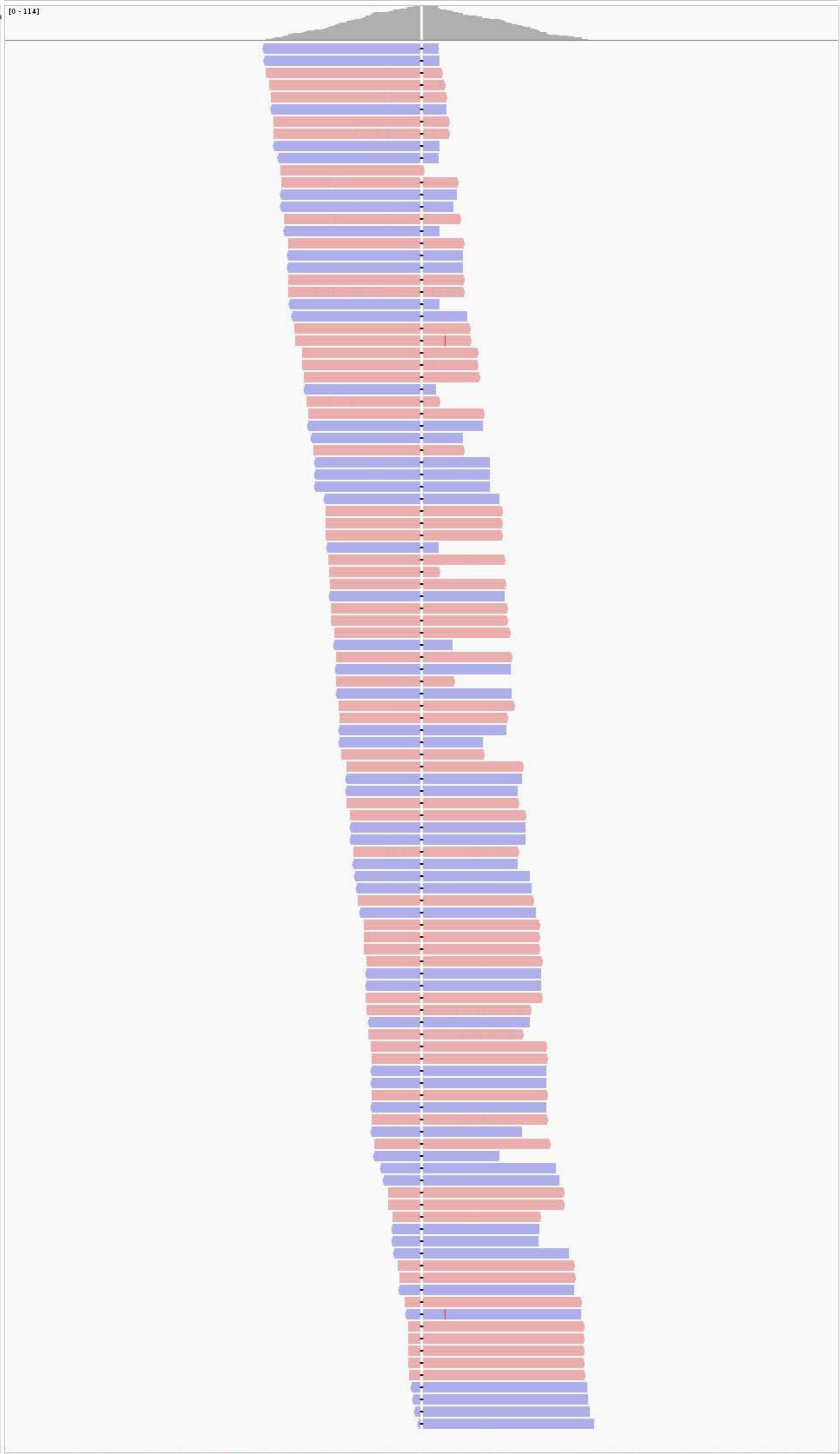
Sequence



chr13:32,357,495-32,358,094

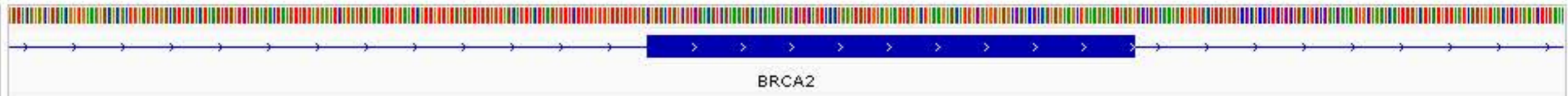


SL_1003_R_78_FFPE-B_mut.bam
rage

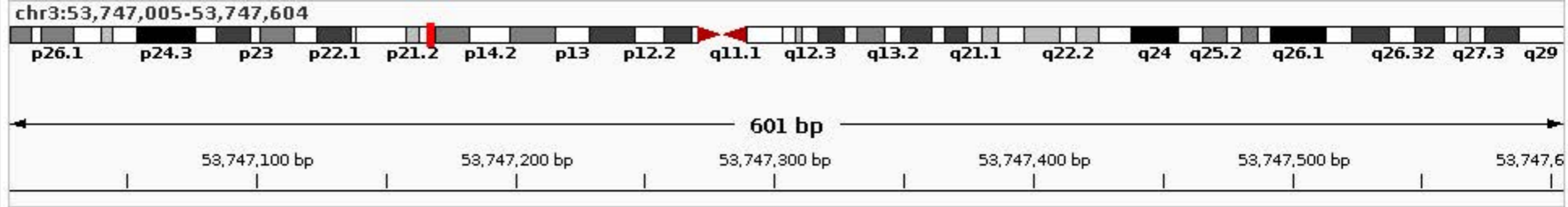


SL_1003_R_78_FFPE-B_mut.ba
m

Sequence →
Refseq Genes



BRCA2

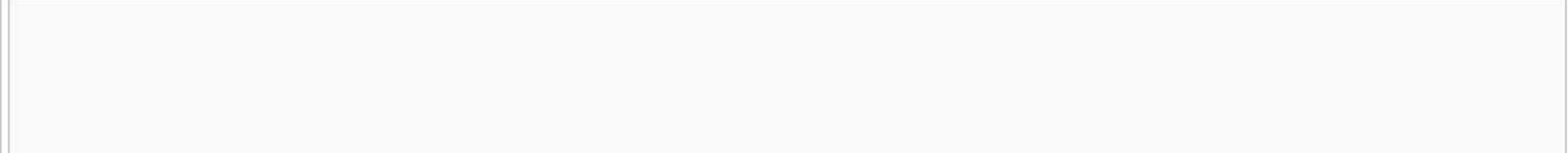
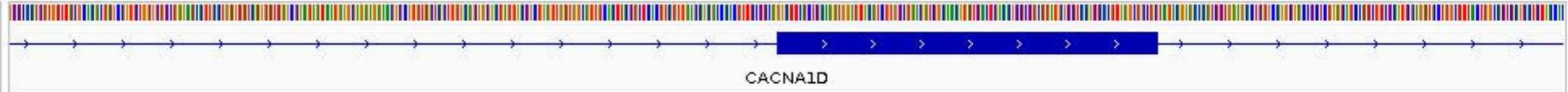
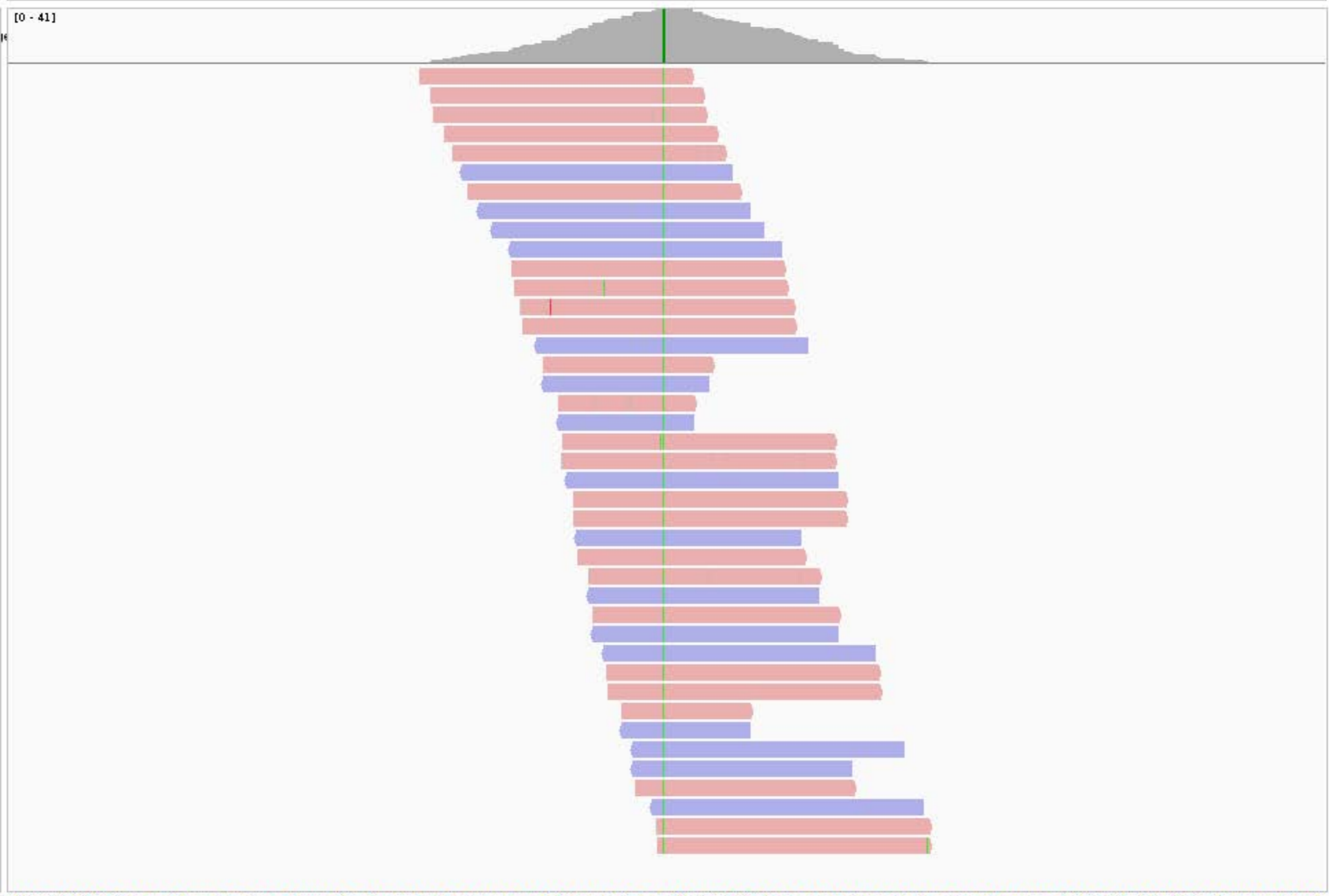


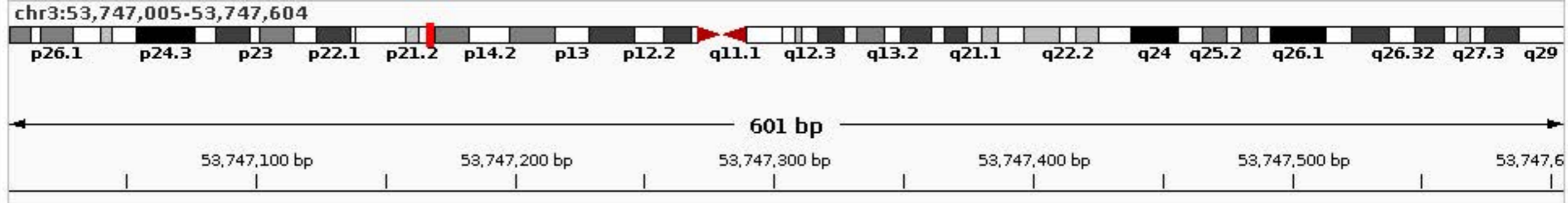
SL_1010-T1-B_mut.bam Coverage

SL_1010-T1-B_mut.bam

Sequence

Refseq Genes





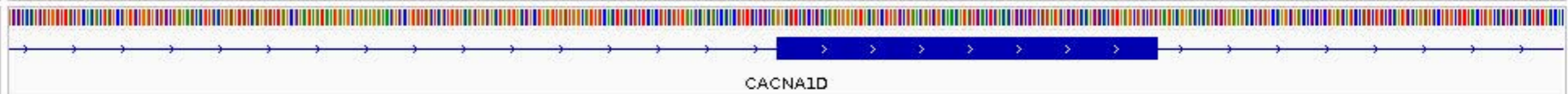
SL_1010-T2-B_mut.bam Coverage

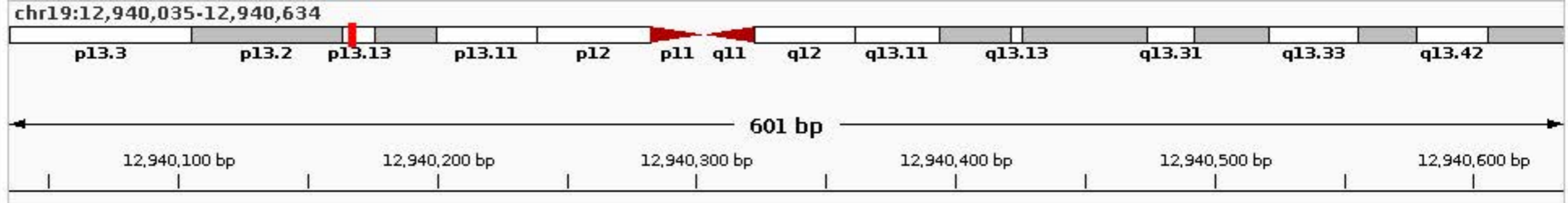
SL_1010-T2-B_mut.bam



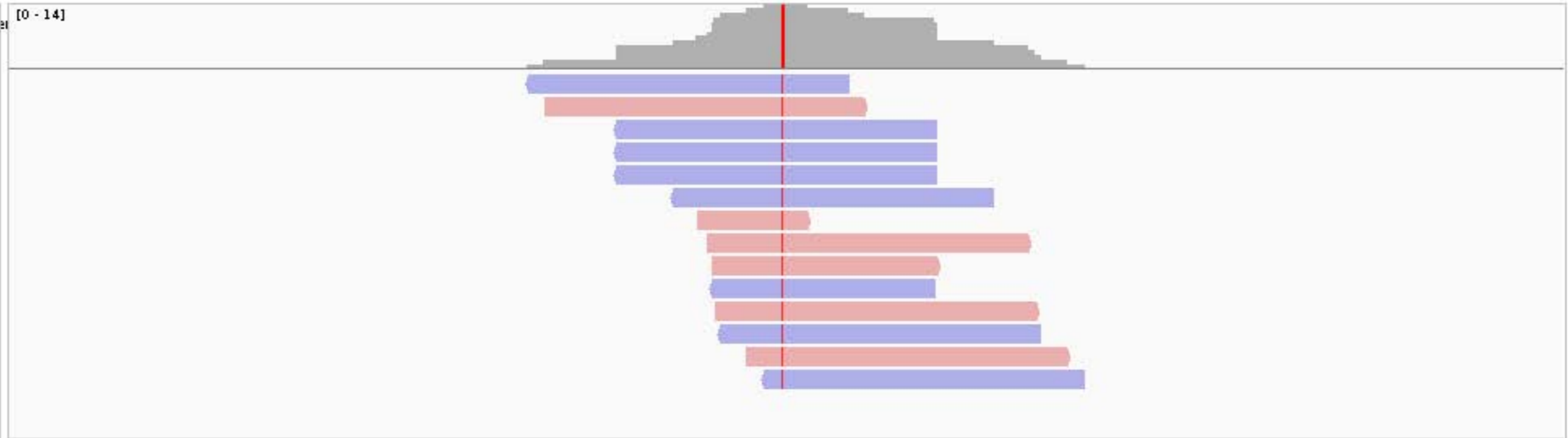
Sequence

Refseq Genes





SL_0005-LT_11-B_mut.bam Cover



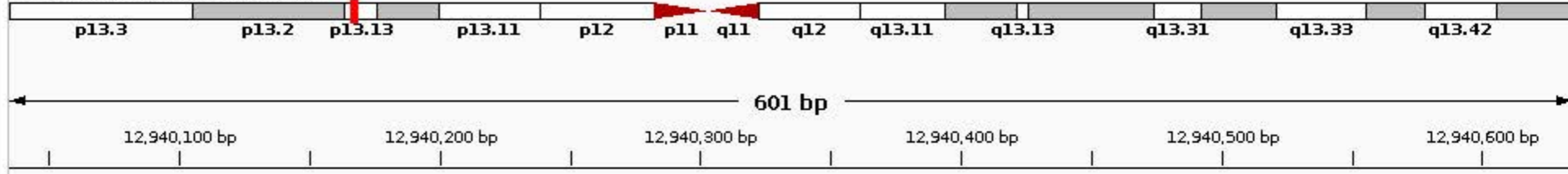
Sequence



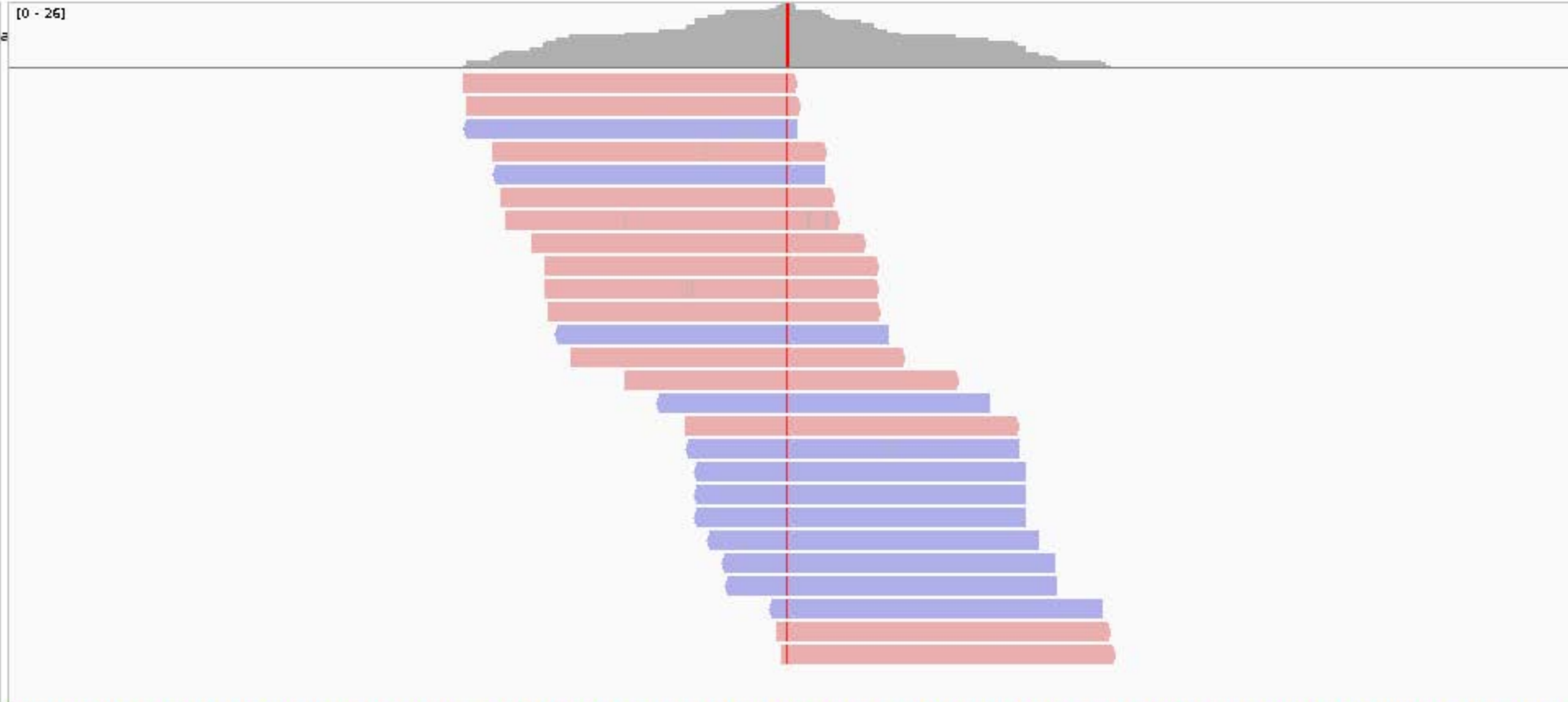
Refseq Genes



chr19:12,940,035-12,940,634



SL_0005-LT-1-B_mut.bam Coverage

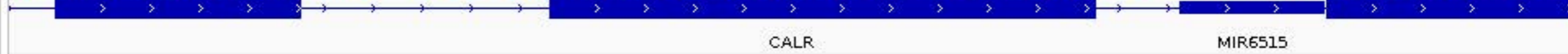


SL_0005-LT-1-B_mut.bam

Sequence

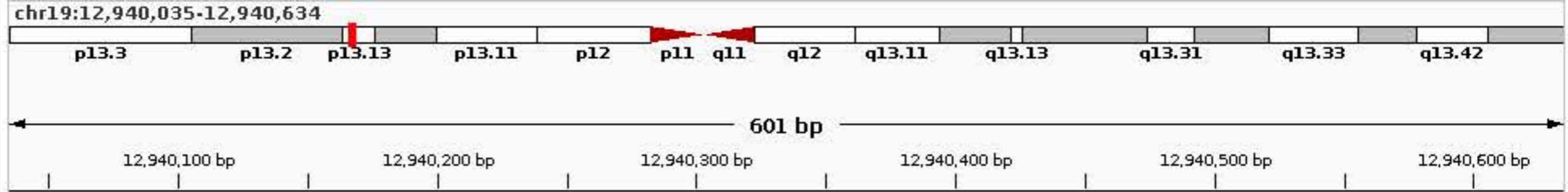


Refseq Genes

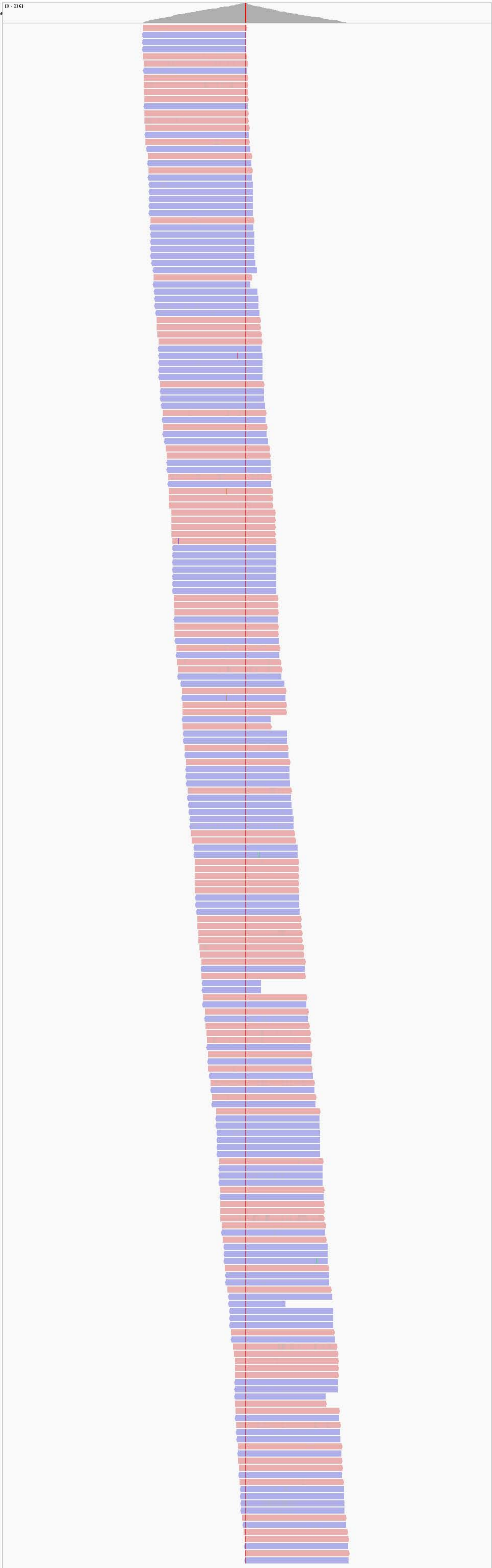


CALR

MIR6515



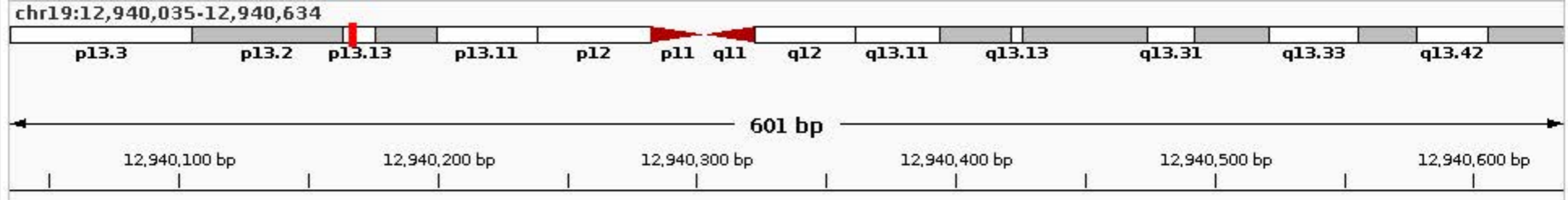
SL_0005-LT-4-B_mut.bam Coverage



SL_0005-LT-4-B_mut.bam

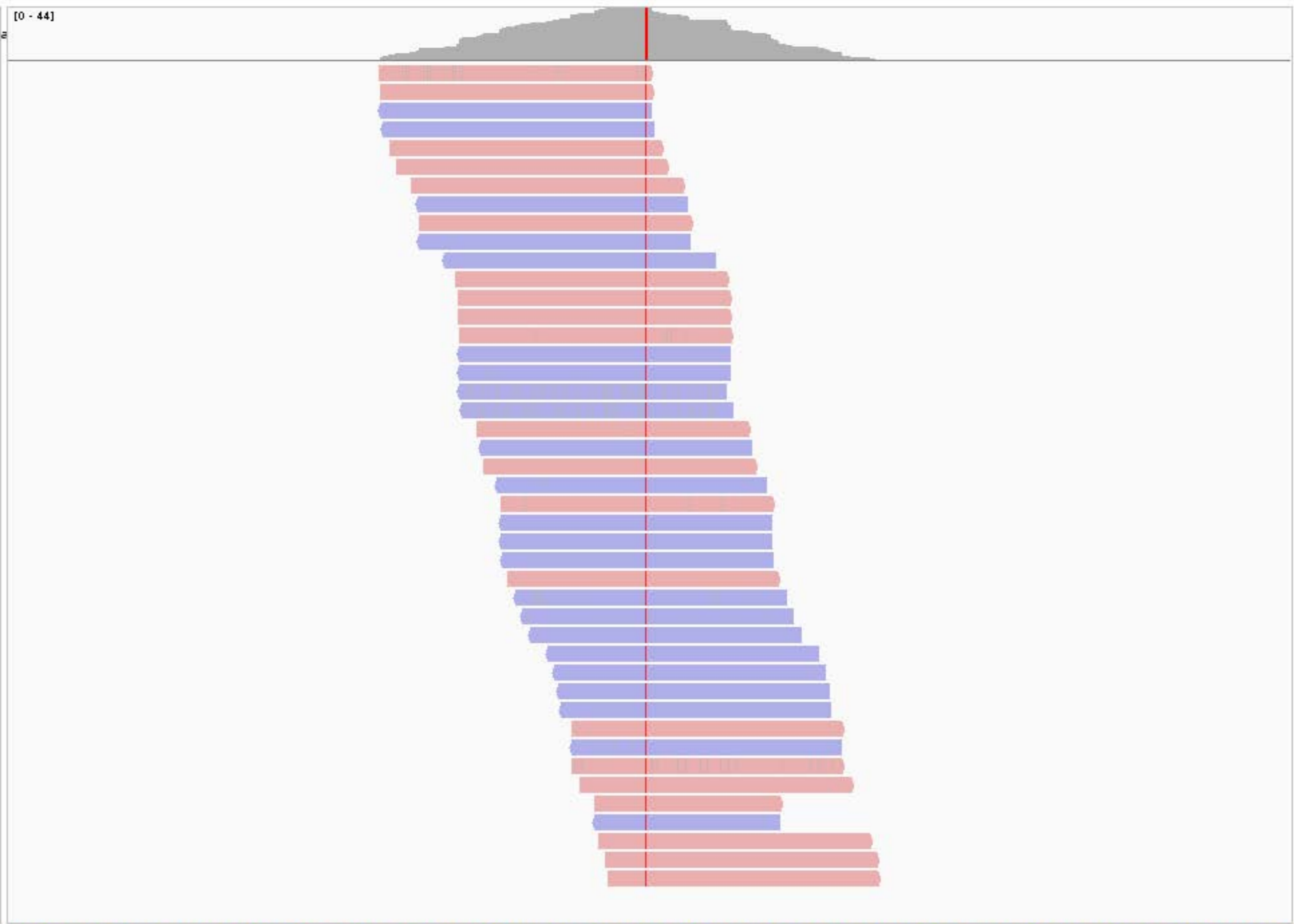
Sequence
Refseq Genes





SL_0005-LT-9-B_mut.bam Coverage

SL_0005-LT-9-B_mut.bam

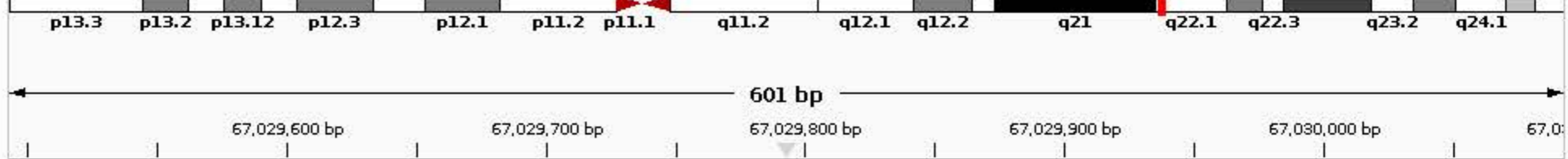


Sequence

Refseq Genes



chr16:67,029,493-67,030,092

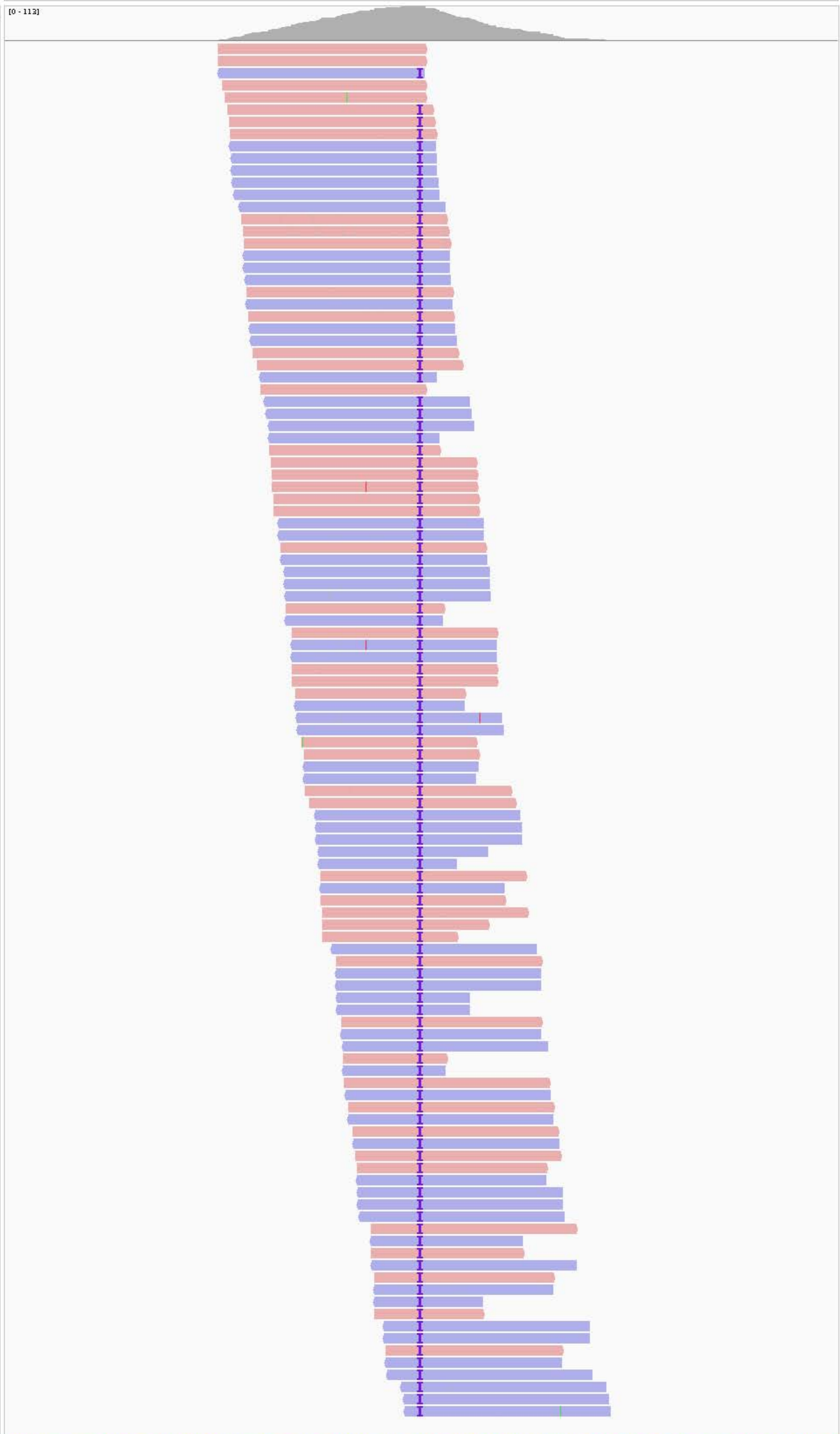


SL_1002_L_FFPE_24-B_mut.bam
rage

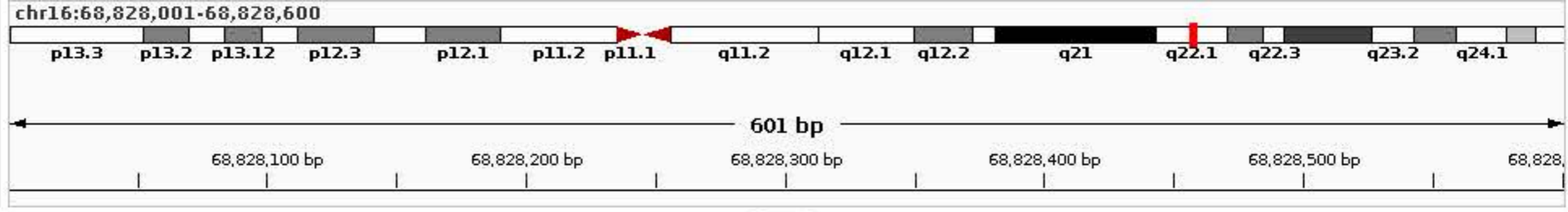
SL_1002_L_FFPE_24-B_mut.ba
m

Sequence →

Refseq Genes



CBFB

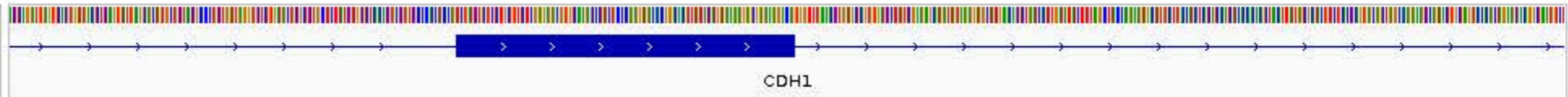
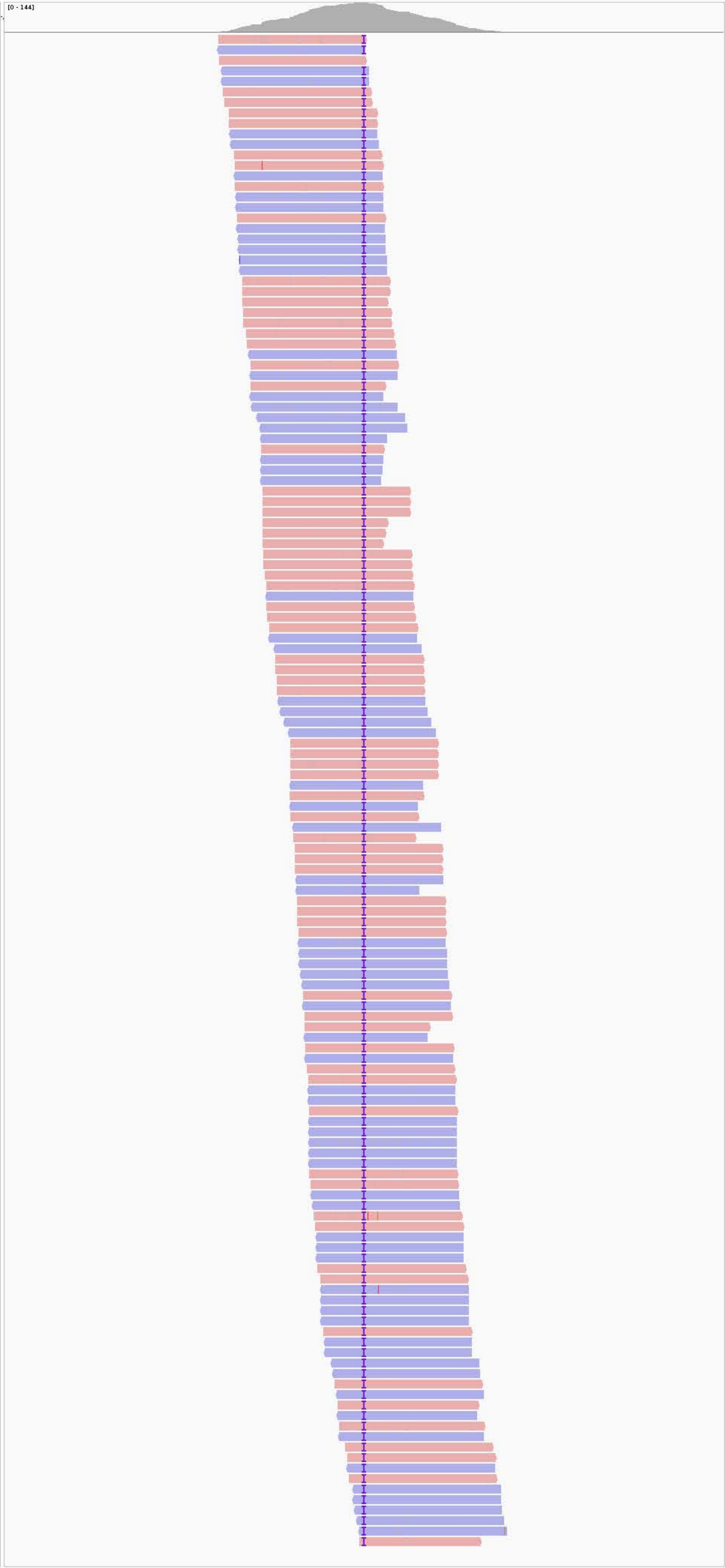


SL_0013_RT_4-B_mut.bam Coverage

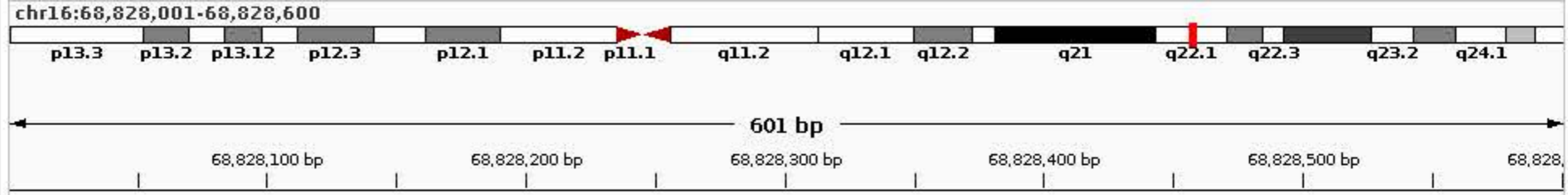
SL_0013_RT_4-B_mut.bam

Sequence

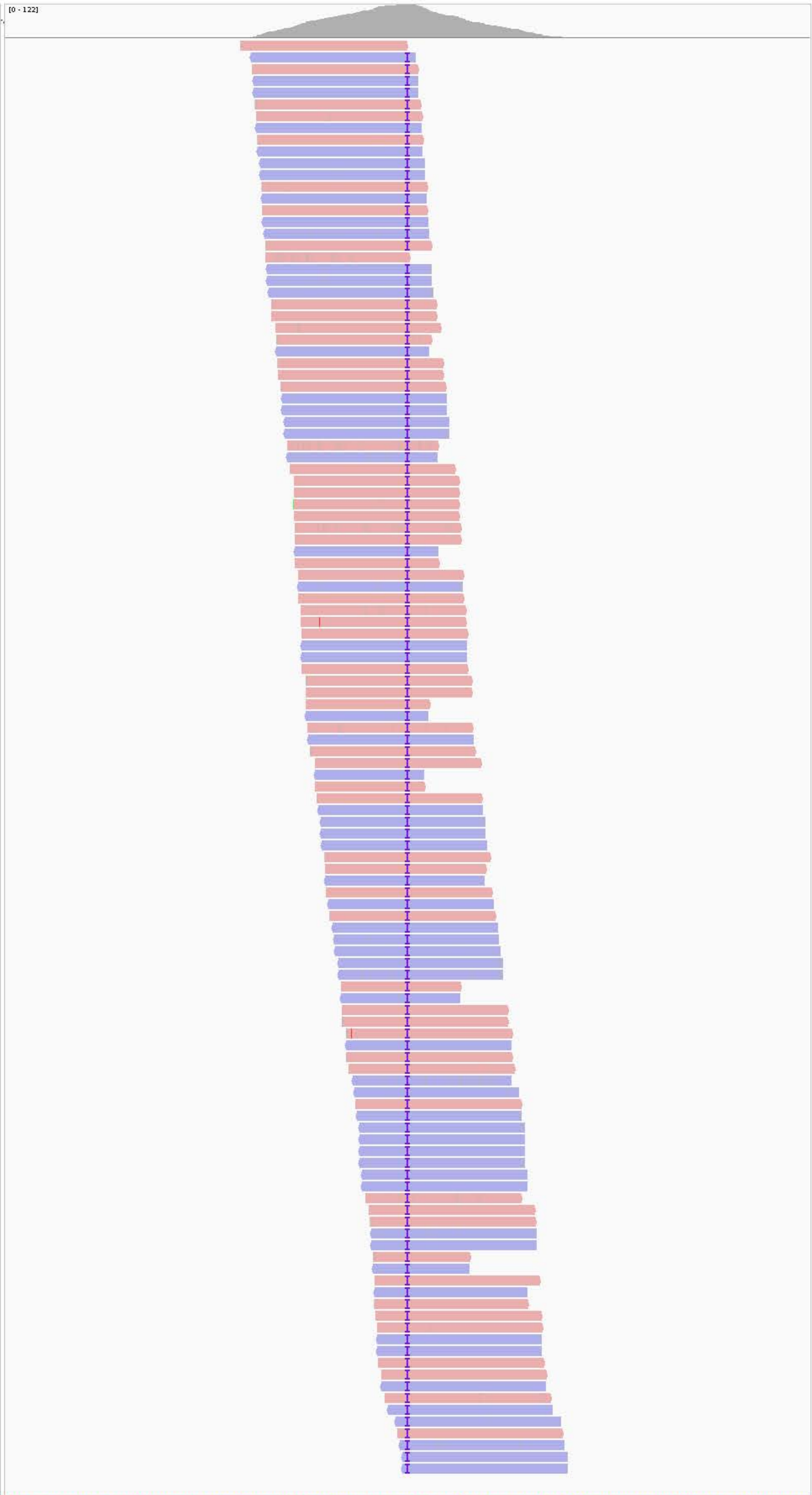
Refseq Genes



CDH1



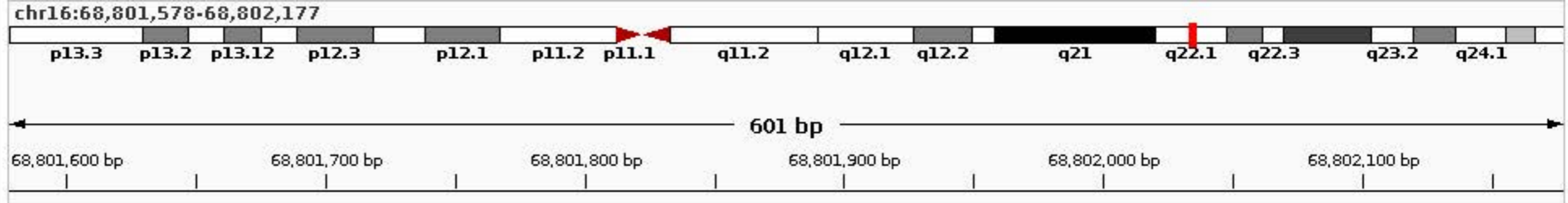
SL_0013_RT_5-B_mut.bam Cover



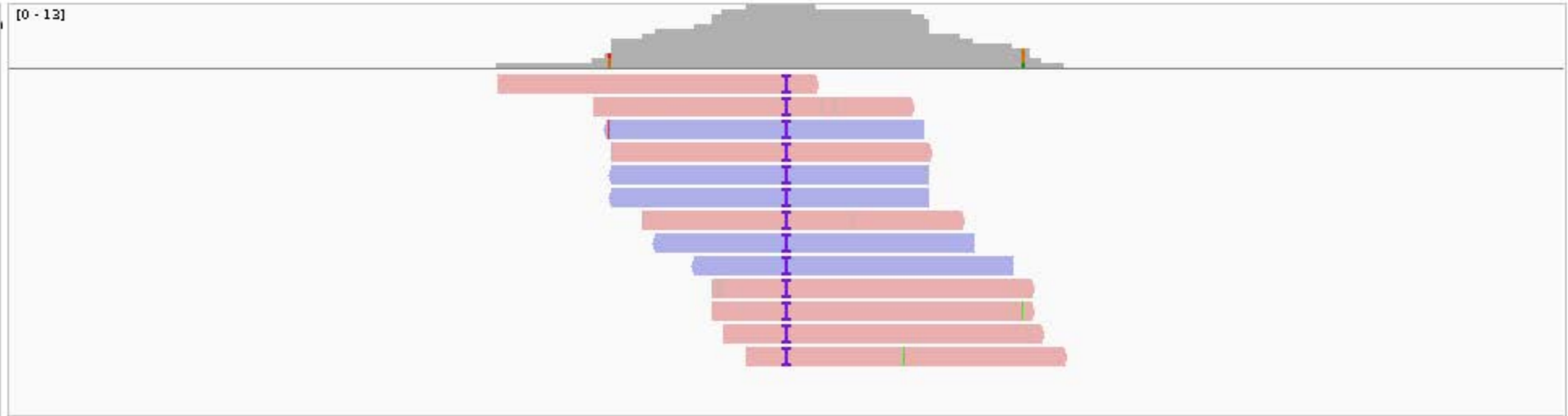
SL_0013_RT_5-B_mut.bam

Sequence →
Refseq Genes





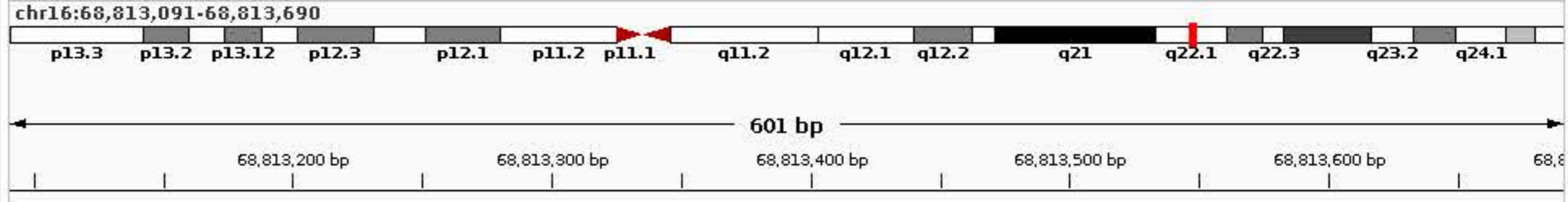
SL_1008_R_51_FFPE-B_mut.bam
 SL_1008_R_51_FFPE-B_mut.bam



Sequence →

Refseq Genes



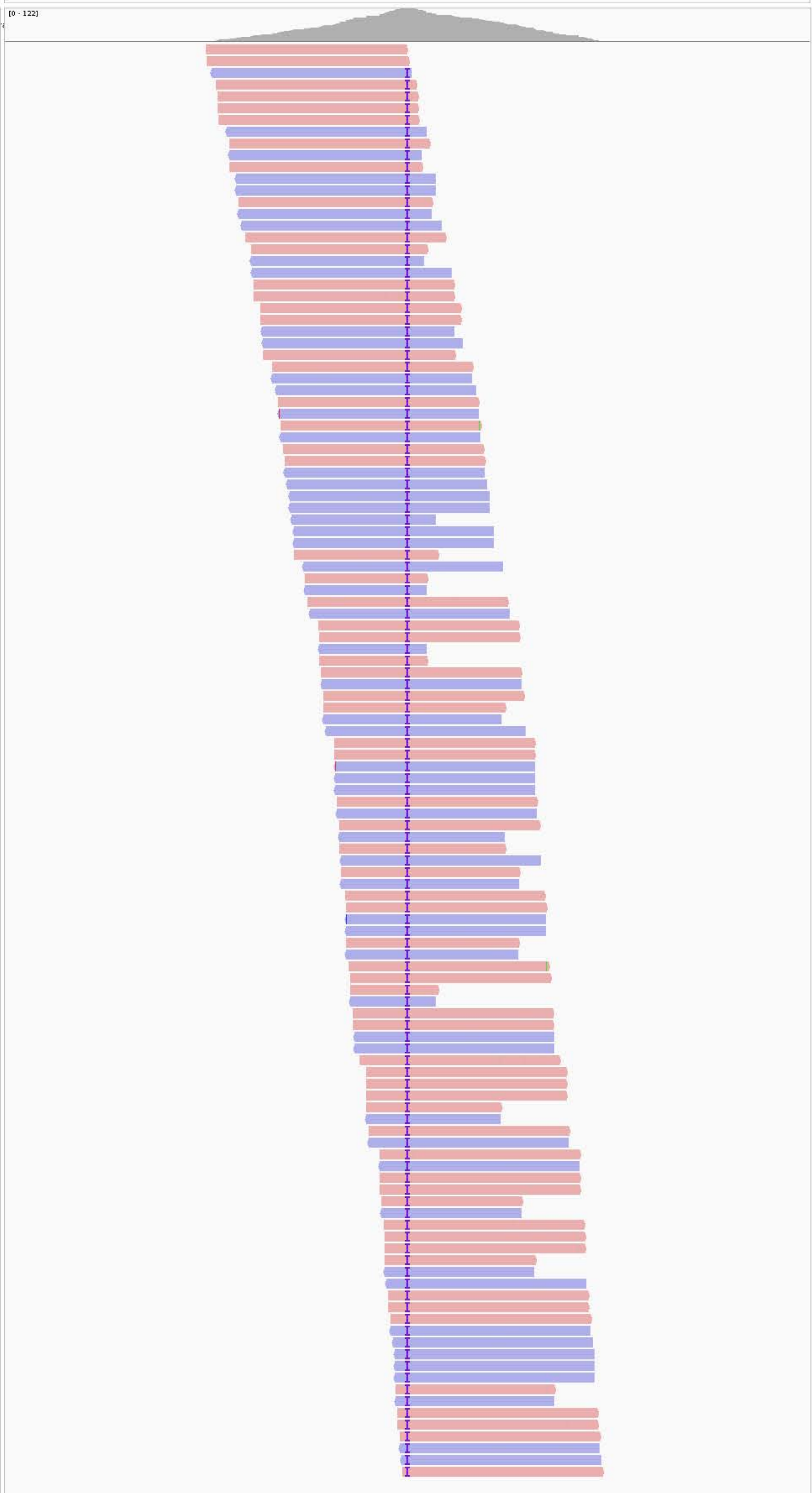


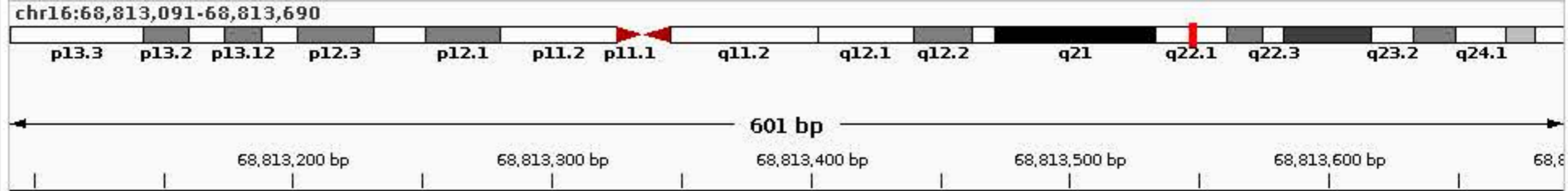
SL_0008_LT_8-B_mut.bam Coverage

SL_0008_LT_8-B_mut.bam

Sequence →

Refseq Genes



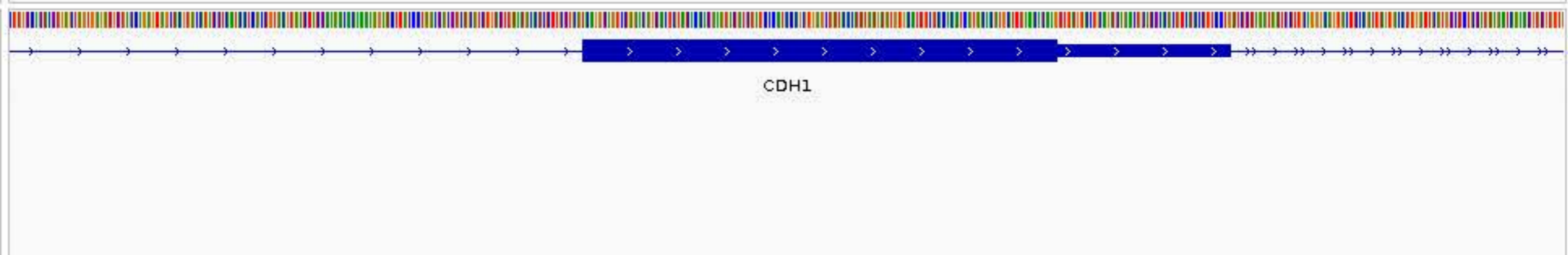
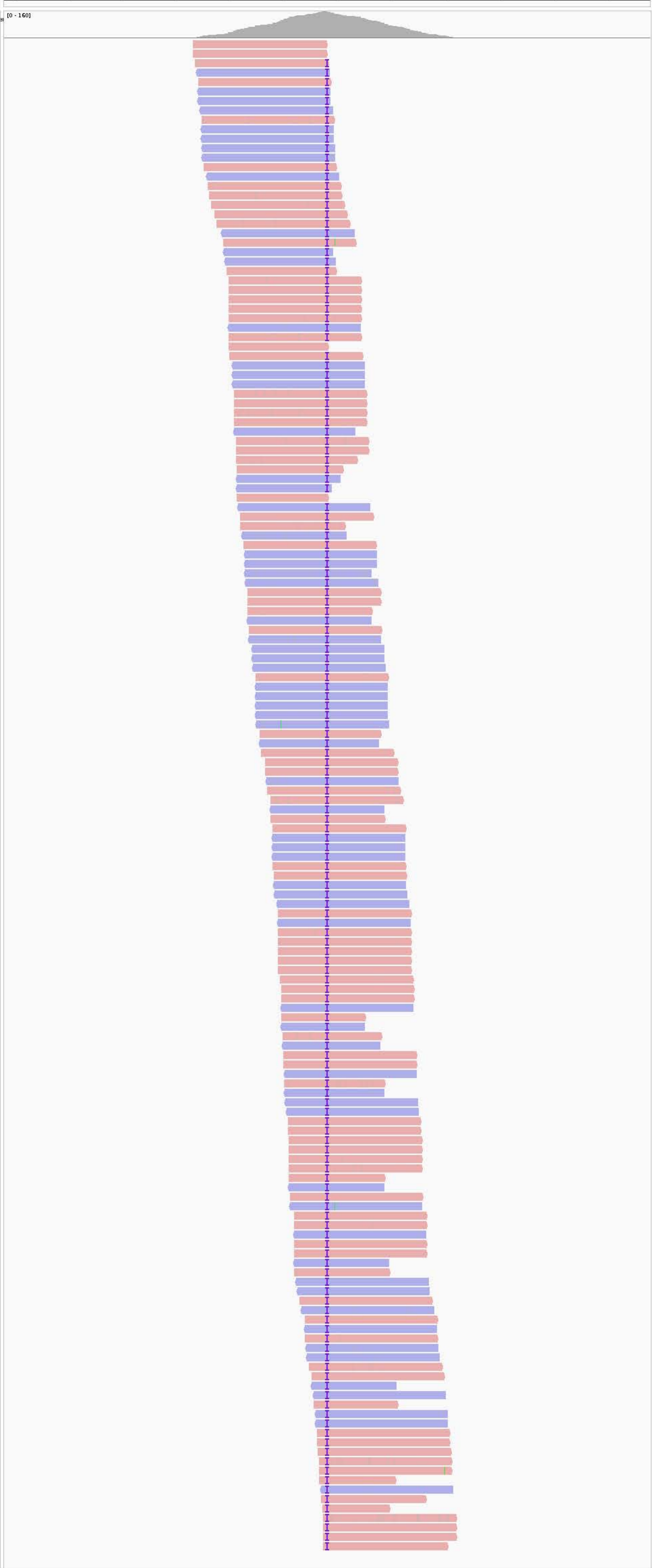


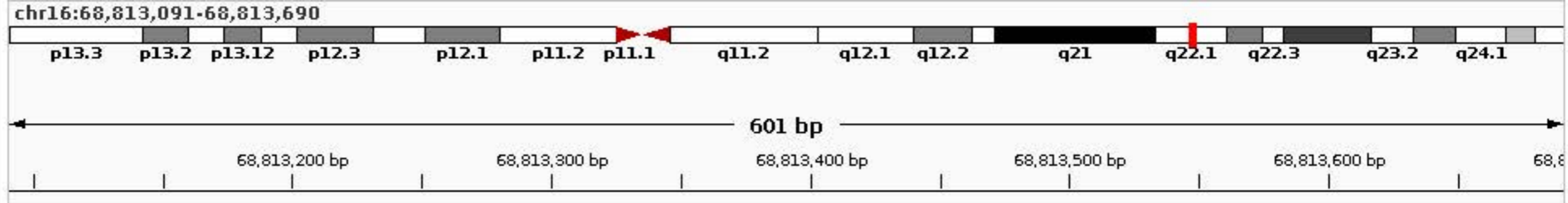
SL_0008-LT_15-B_mut.bam Cov

SL_0008-LT_15-B_mut.bam

Sequence

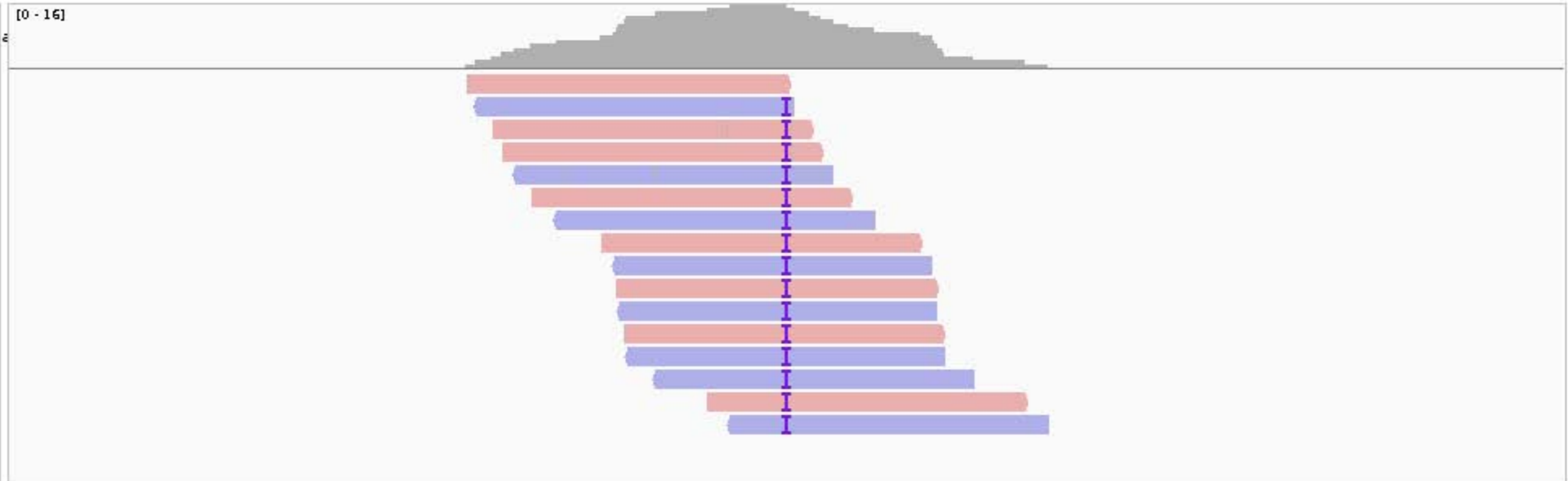
Refseq Genes





SL_0009-LT_9-B_mut.bam Coverage

SL_0009-LT_9-B_mut.bam



Sequence

Refseq Genes

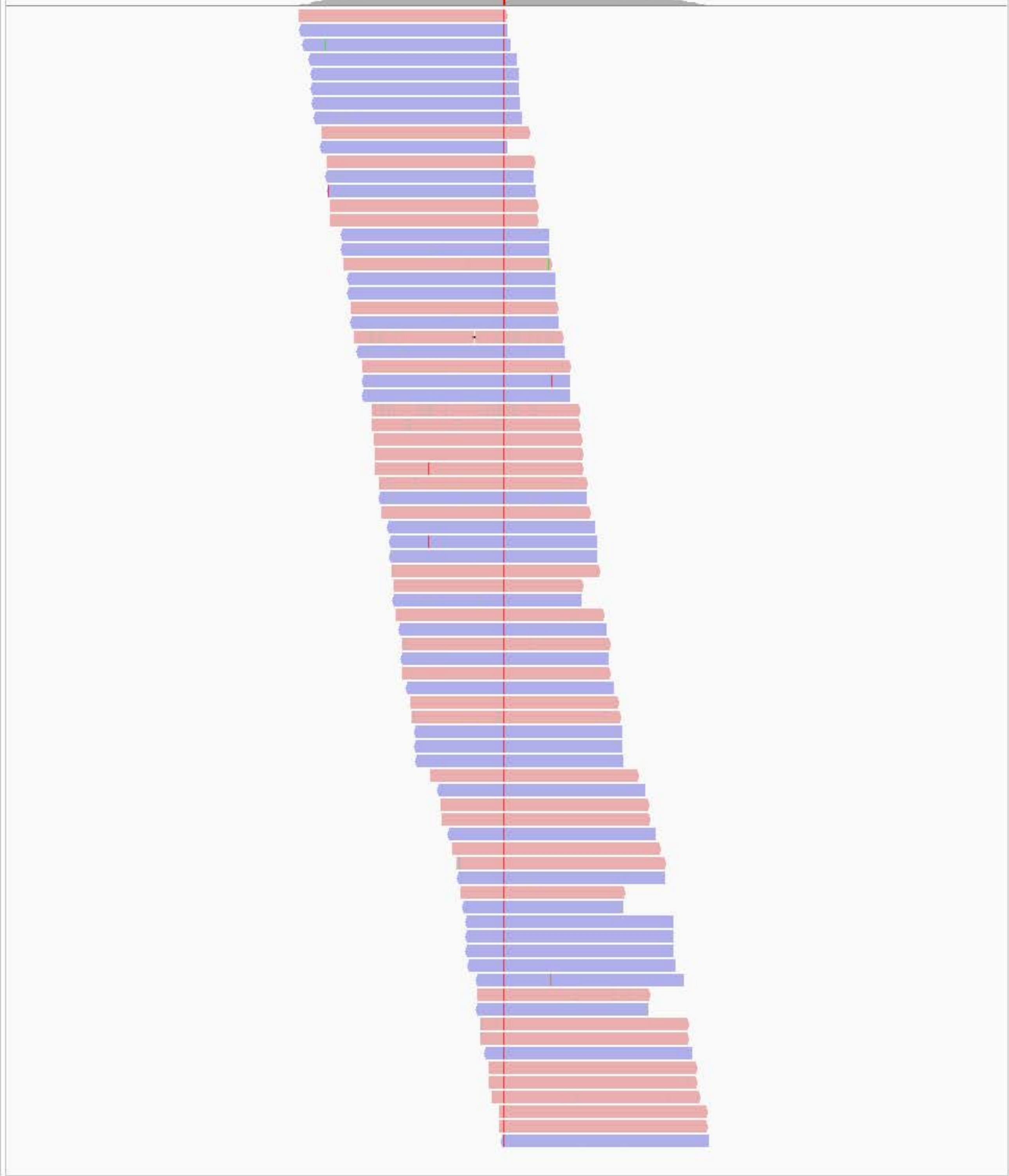


chr12:12,717,733-12,718,332
p13.32 p13.2 p12.2 p11.23 p11.1 q12 q13.12 q13.3 q14.2 q15 q21.2 q21.32 q22 q23.2 q24.11 q24.22 q24.32

601 bp
12,717,800 bp 12,717,900 bp 12,718,000 bp 12,718,100 bp 12,718,200 bp 12,718,300 bp

[0 - 78]

SL_0013_RT_4-B_mut.bam Cover

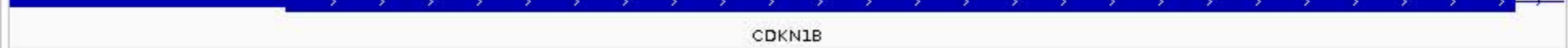


SL_0013_RT_4-B_mut.bam

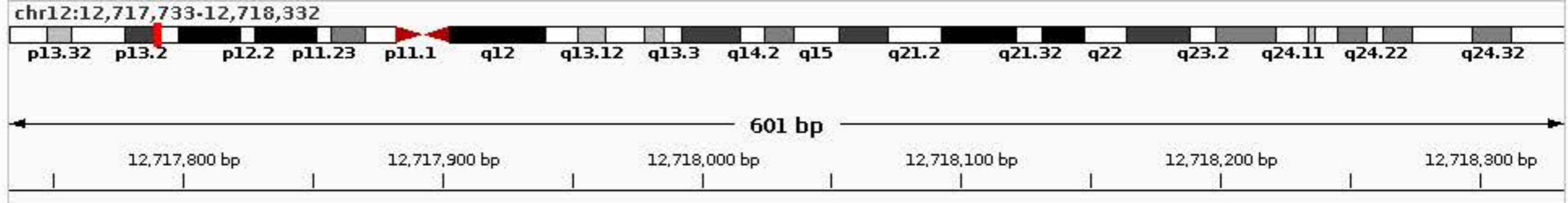
Sequence



Refseq Genes



CDKN1B

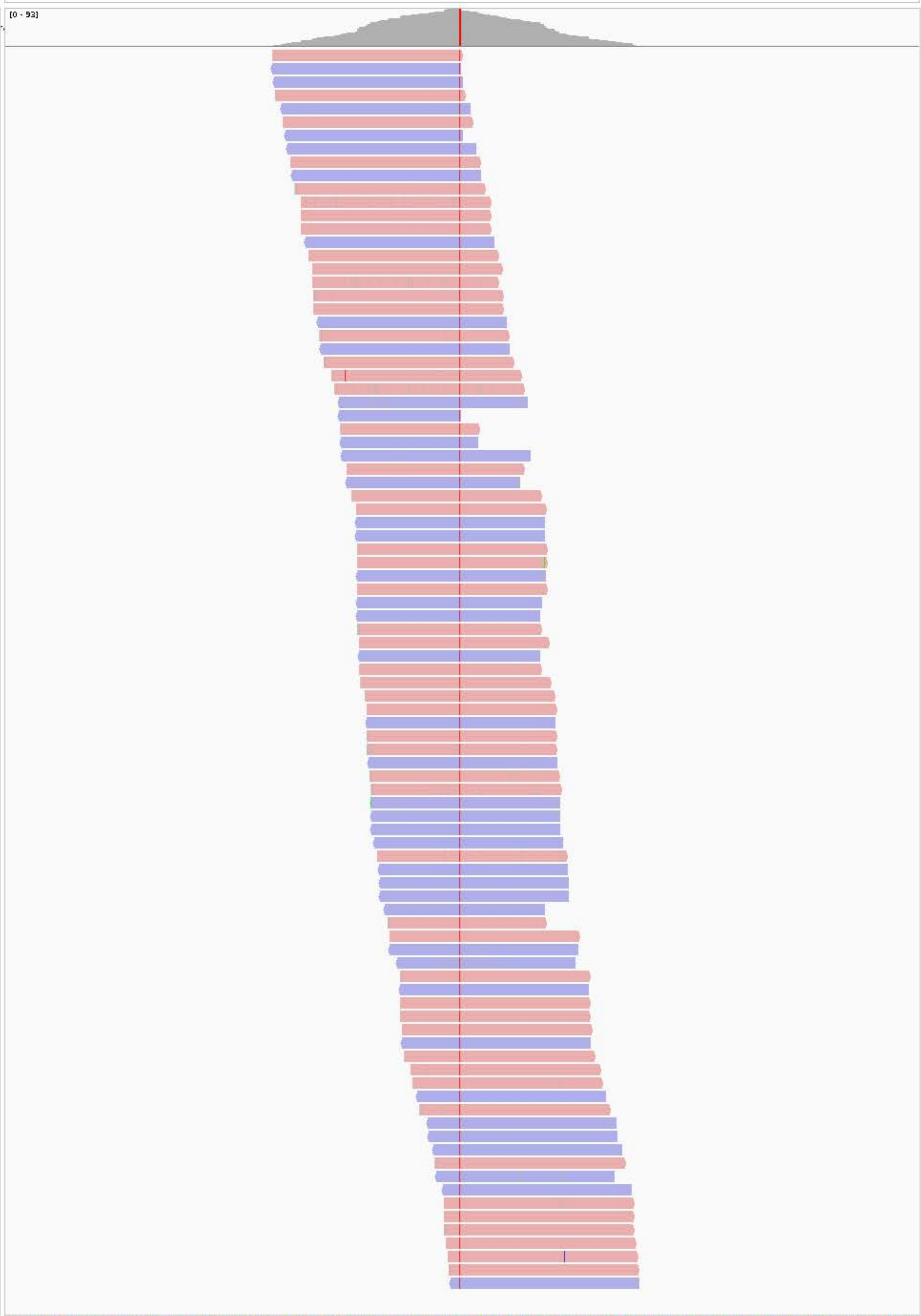


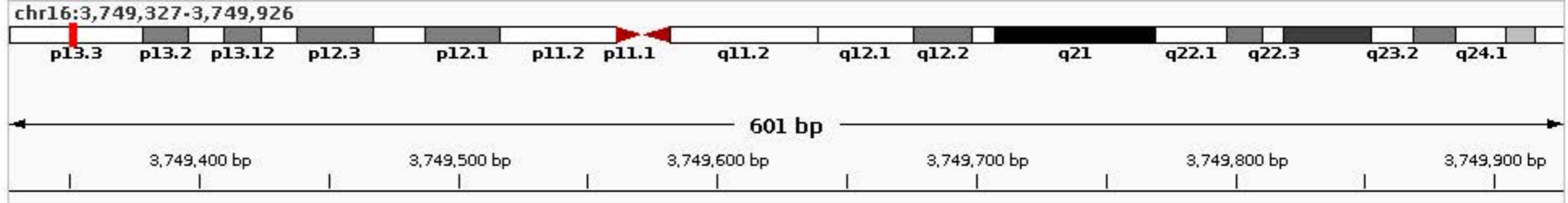
SL_0013_RT_5-B_mut.bam Cover.

SL_0013_RT_5-B_mut.bam

Sequence →

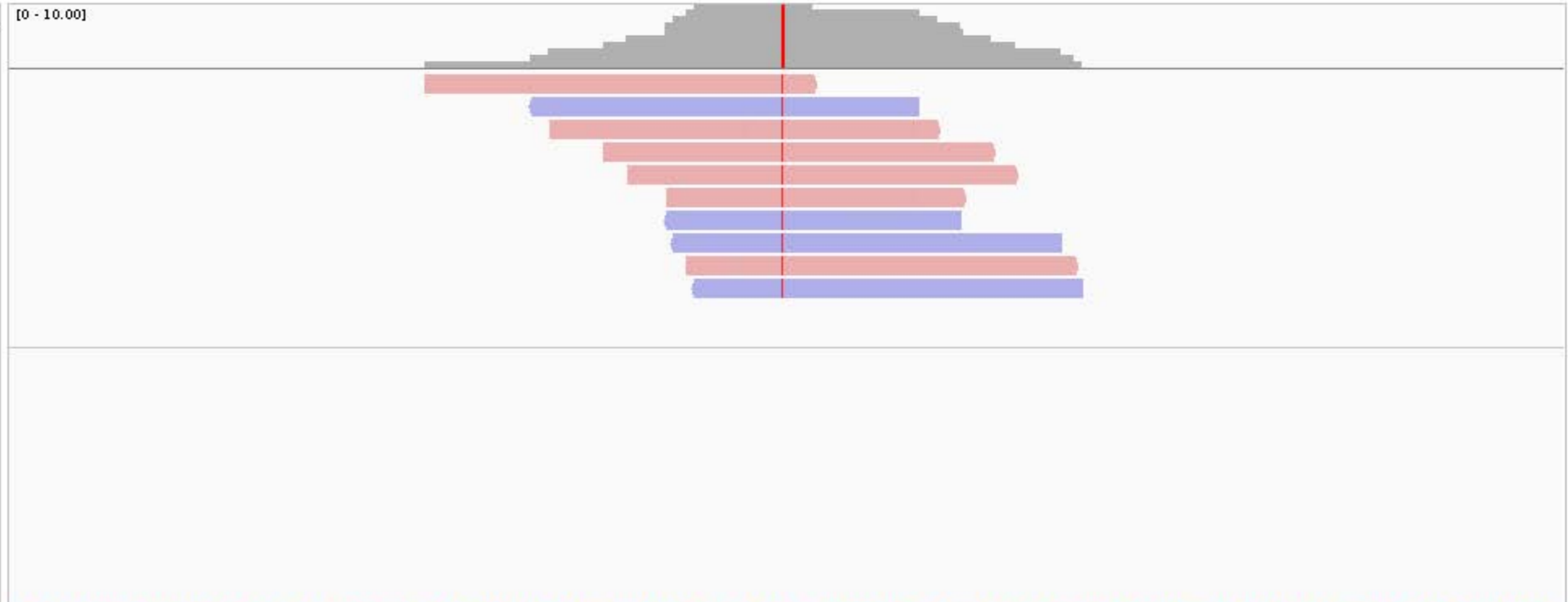
Refseq Genes





SL_0001_L_FFPE_10-B_mut.barr
rage

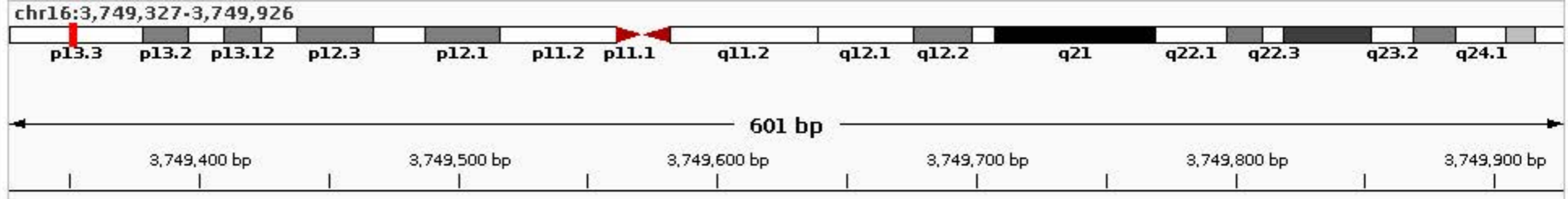
SL_0001_L_FFPE_10-B_mut.ba
m



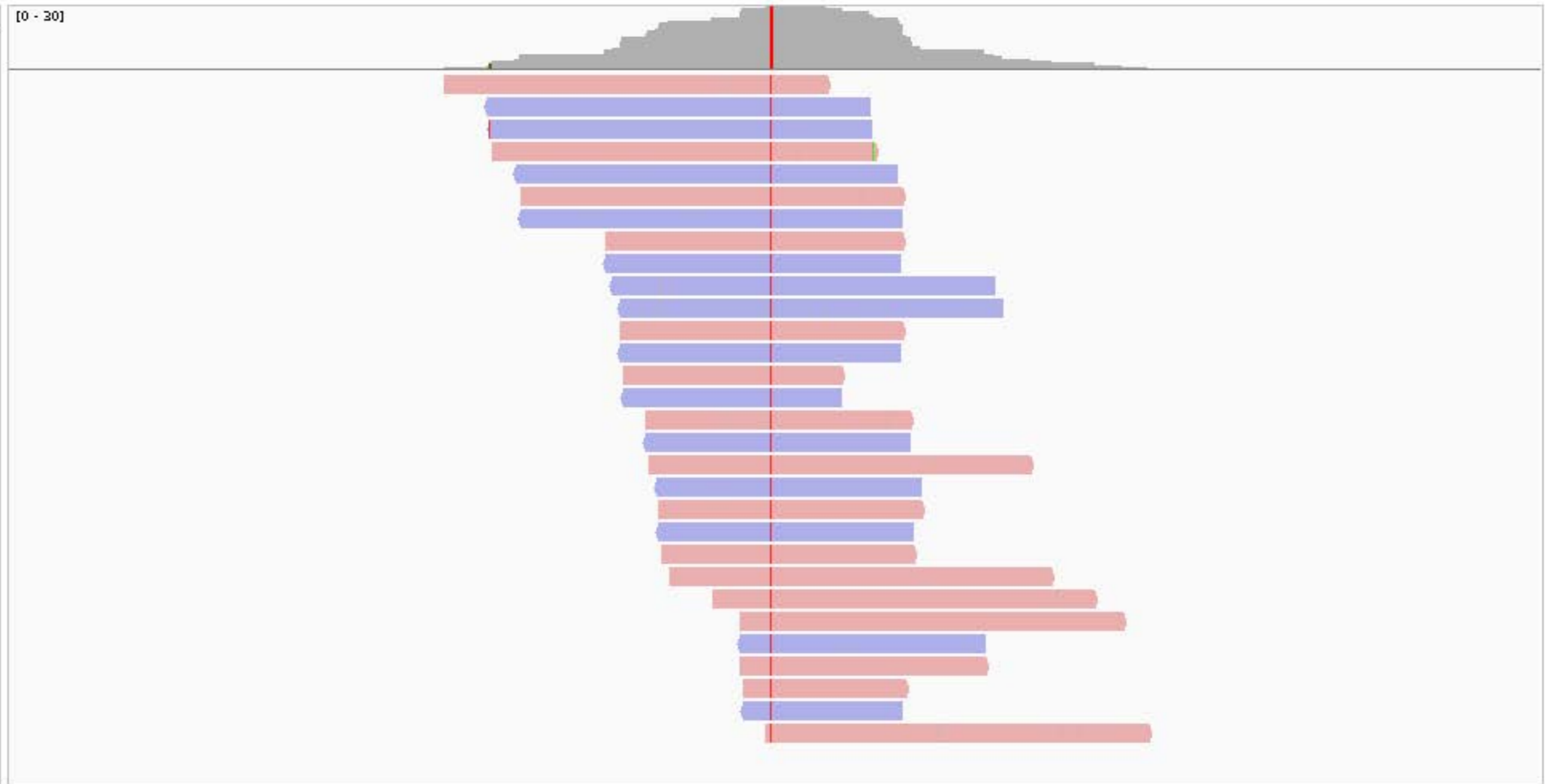
Sequence →

Refseq Genes





SL_1007_L_FFPE_21-B_mut.barrage



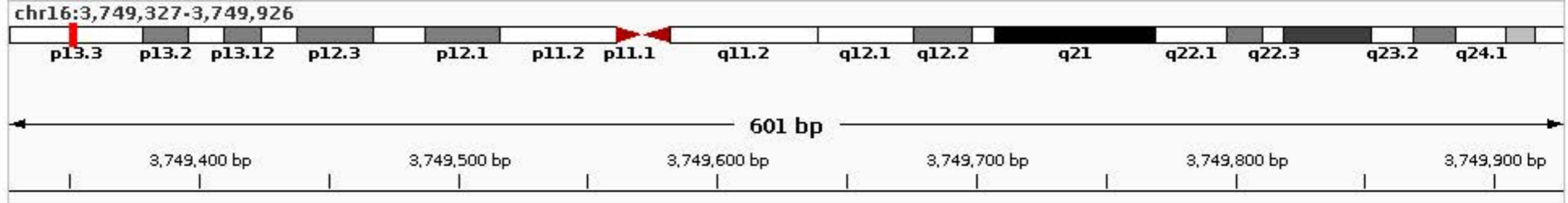
Sequence →



Refseq Genes

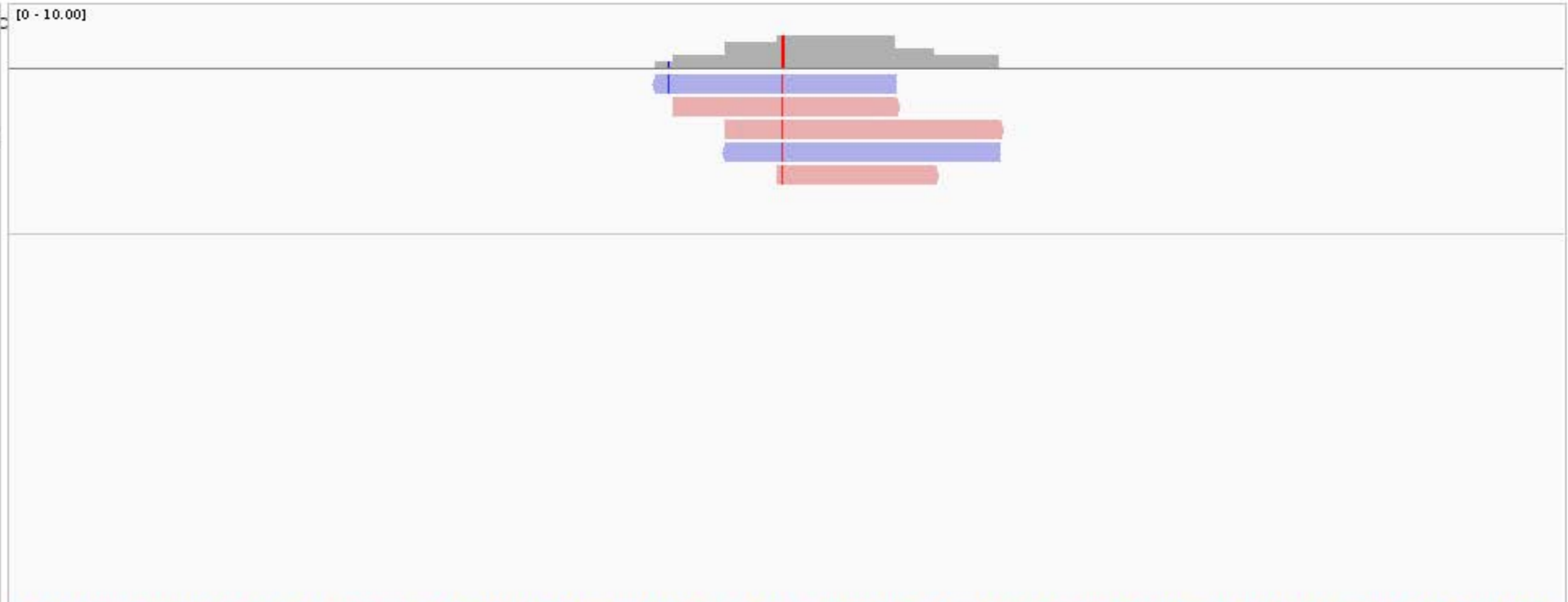


CREBBP



SL_1010_L_FFPE_3-B_mut.bam
age

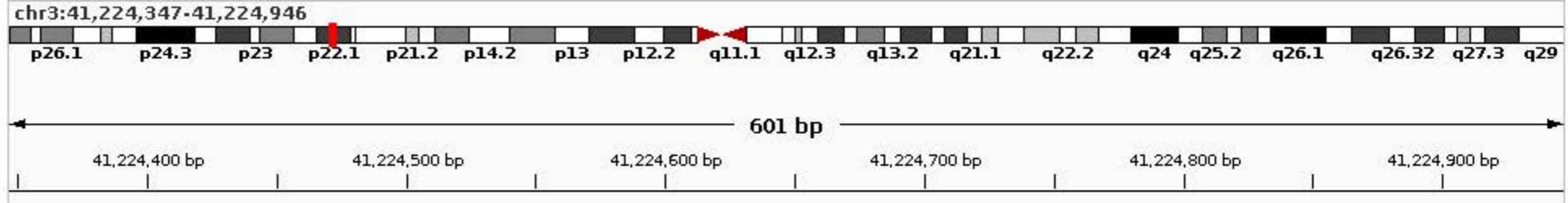
SL_1010_L_FFPE_3-B_mut.bam



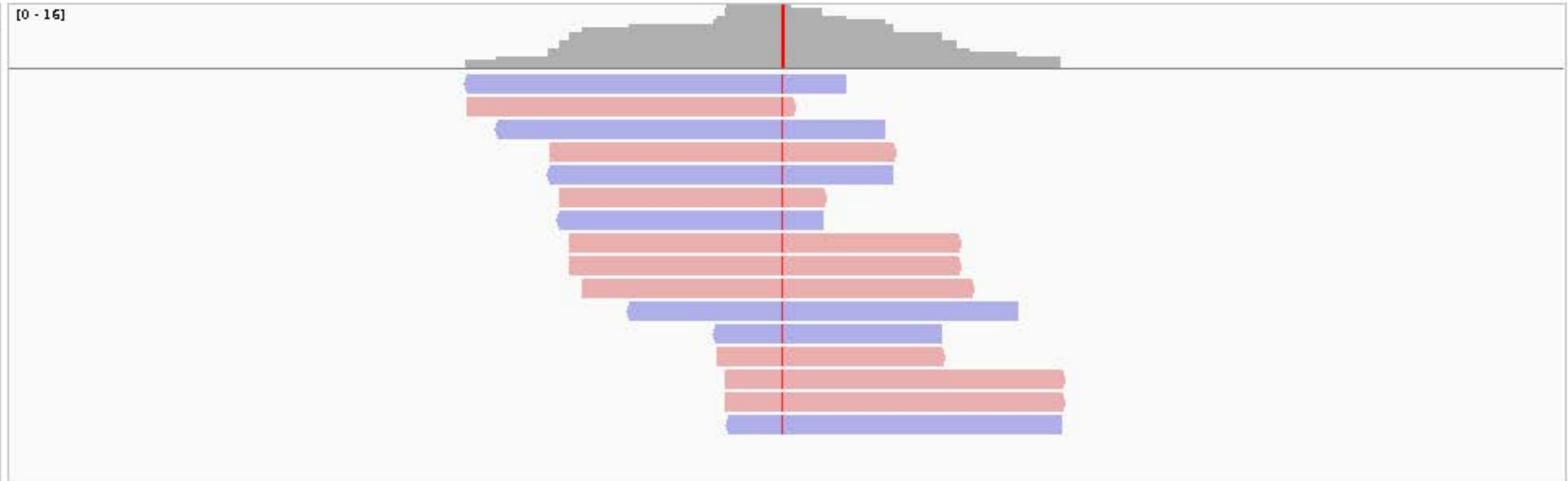
Sequence →

Refseq Genes





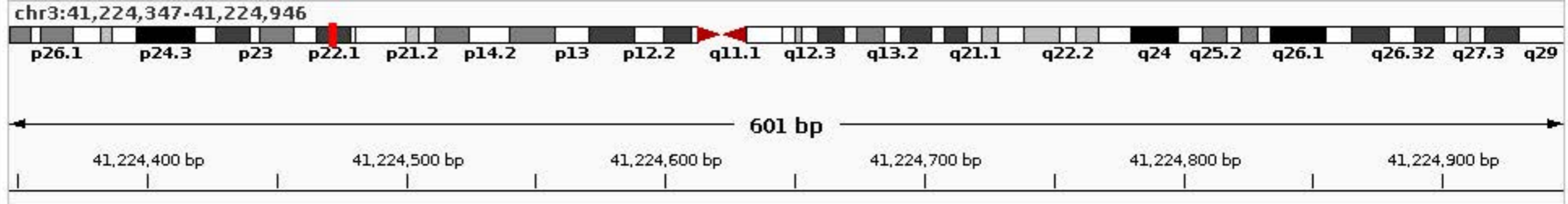
SL_0005_L_FFPE_21-B_mut.bam
range



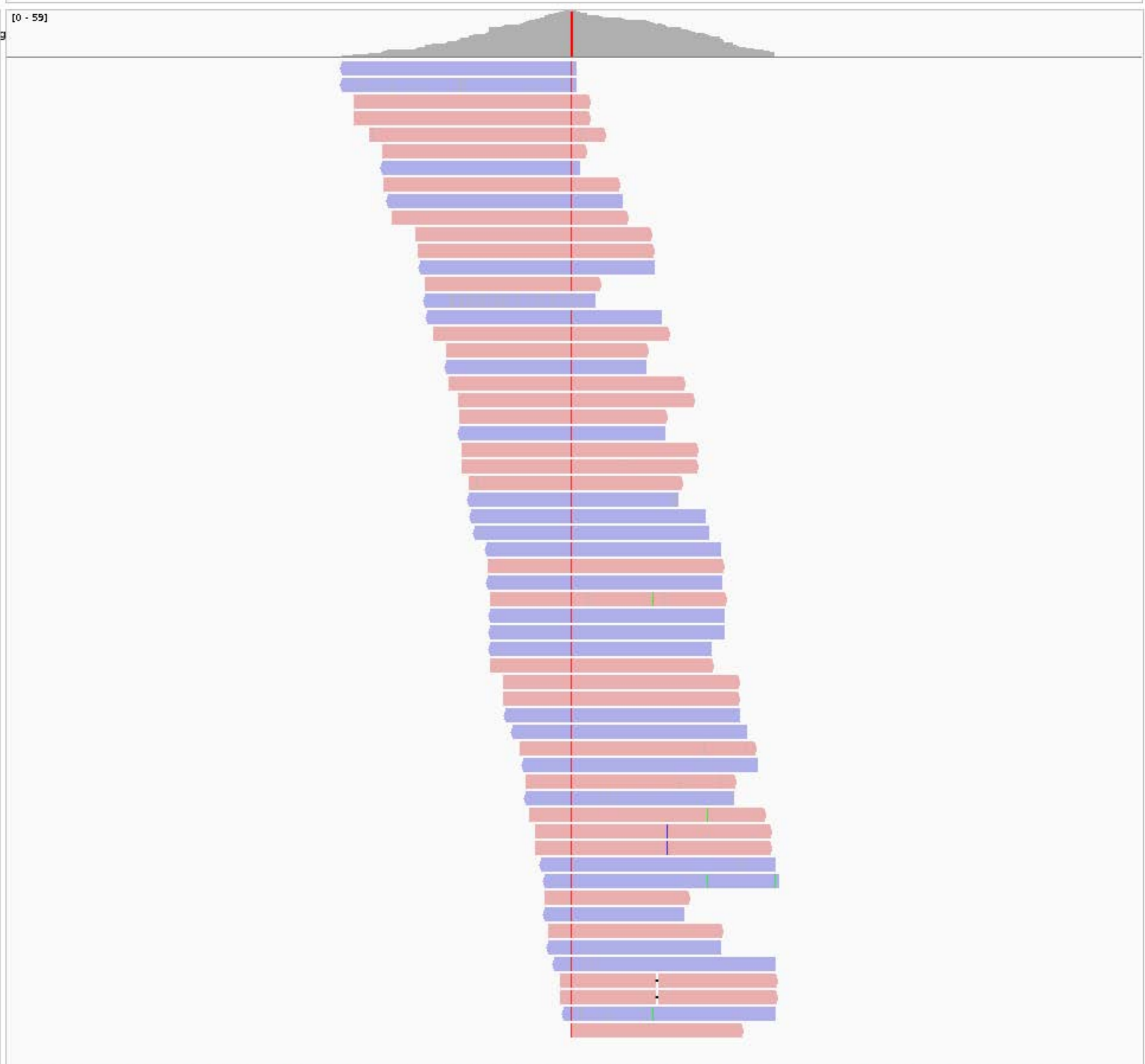
SL_0005_L_FFPE_21-B_mut.bam

Sequence →
Refseq Genes





SL_1010-N2-B_mut.bam Coverage

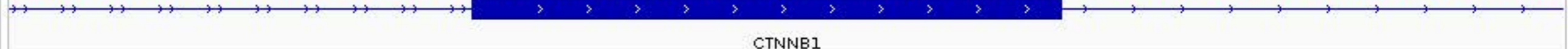


SL_1010-N2-B_mut.bam

Sequence

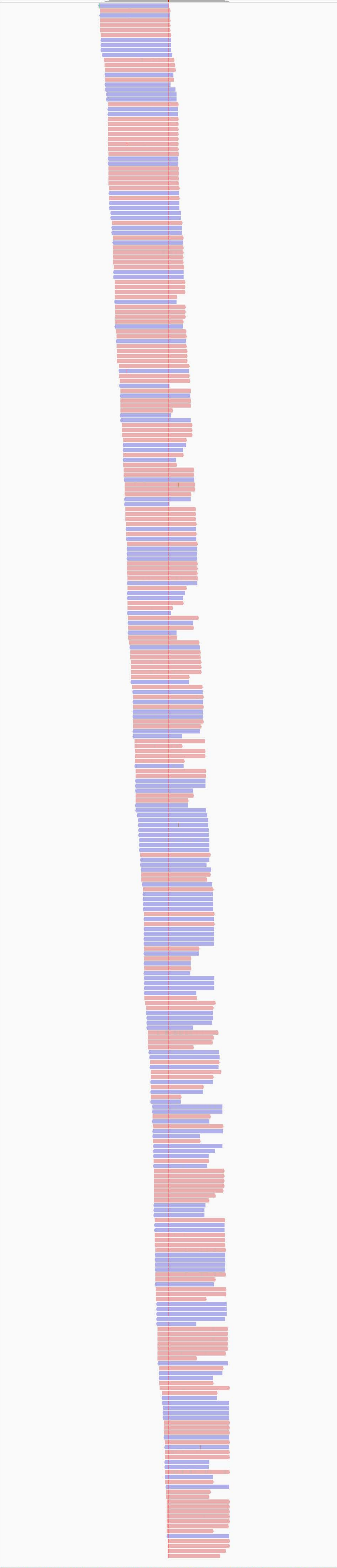


Refseq Genes



CTNNB1

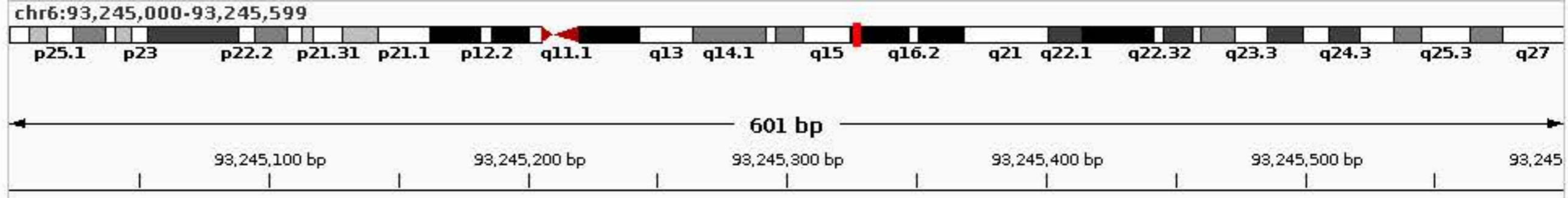
SL_1010-N3-B_mut.bam Coverage
[0 - 215]
SL_1010-N3-B_mut.bam



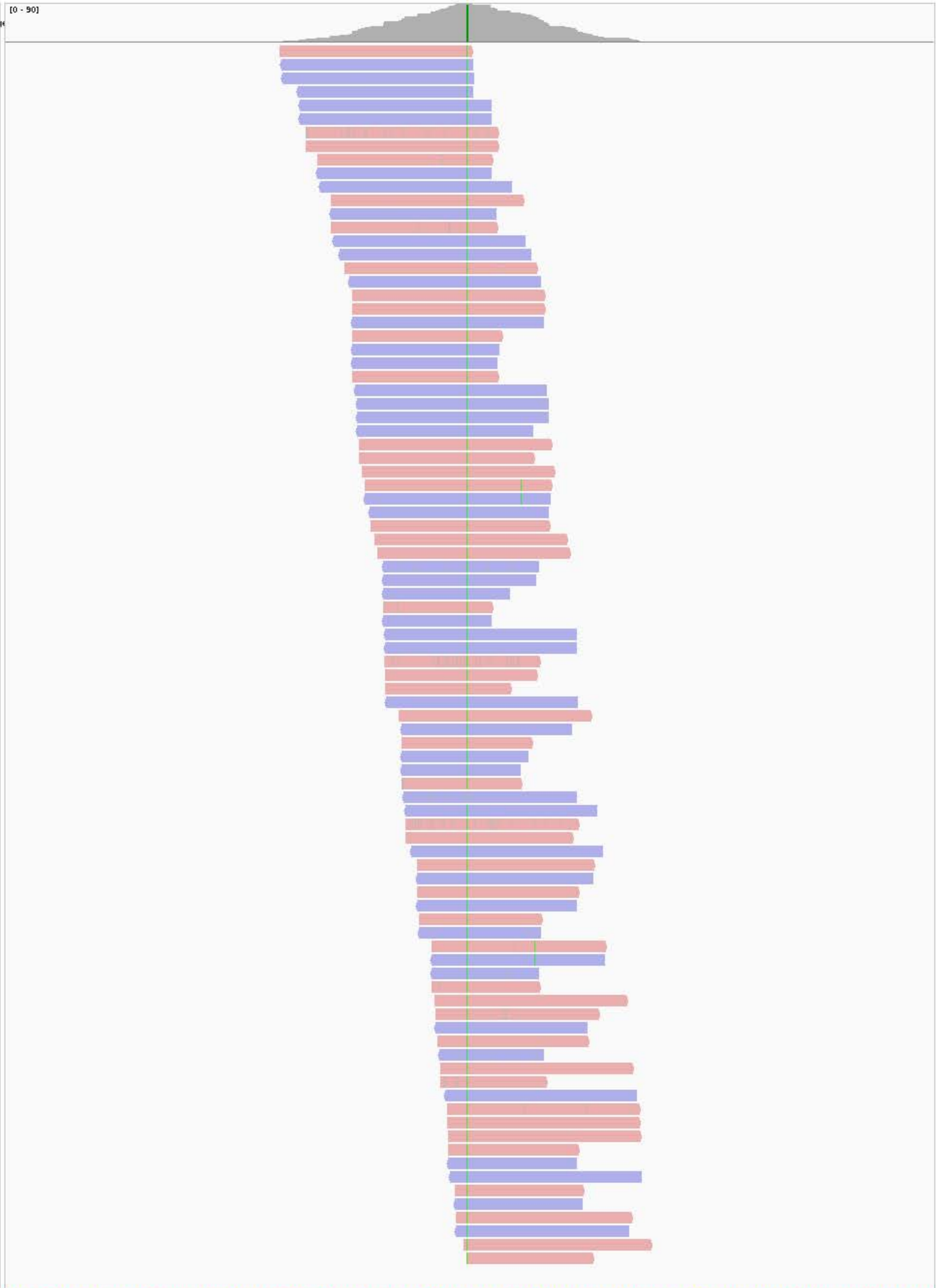
Sequence

Refseq Genes

CTNNE1



SL_1011-T1-B_mut.bam Coverage



SL_1011-T1-B_mut.bam

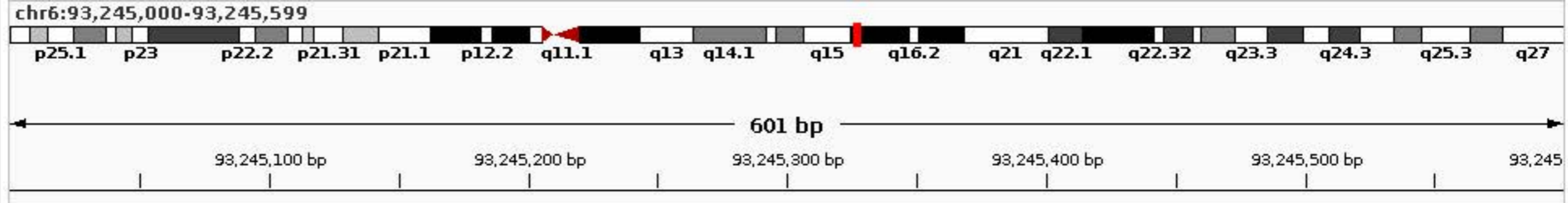
Sequence →



Refseq Genes

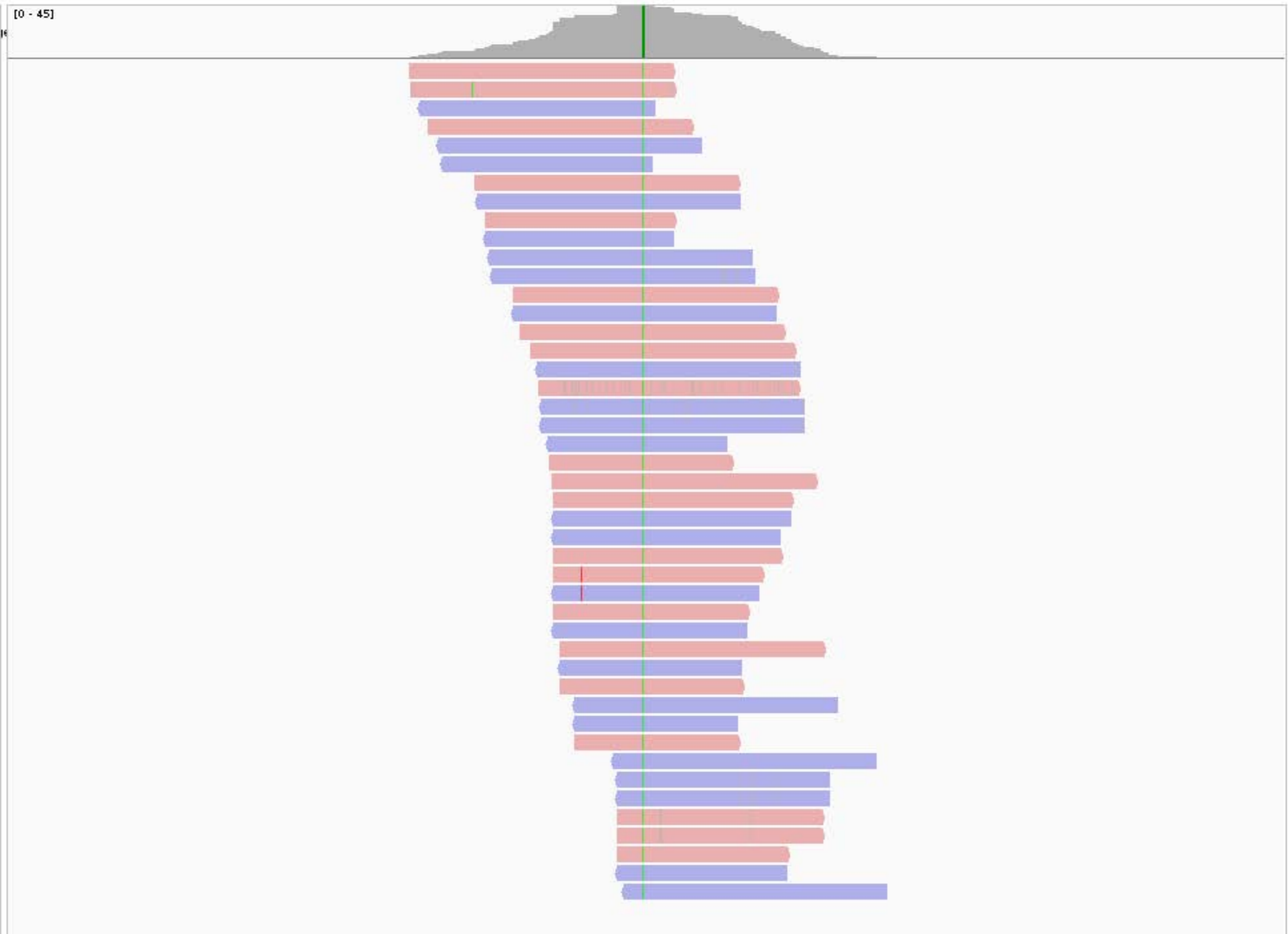


EPHA7



SL_1011-T2-B_mut.bam Coverage

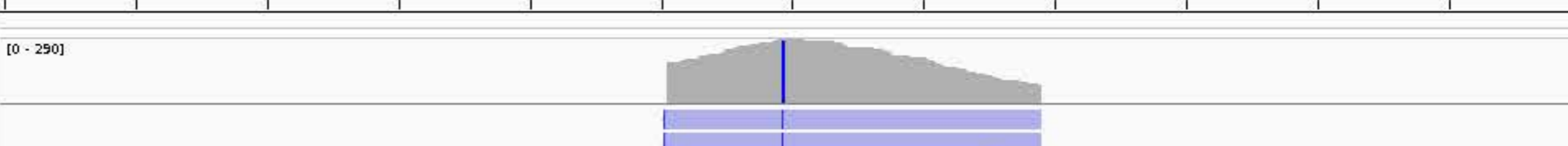
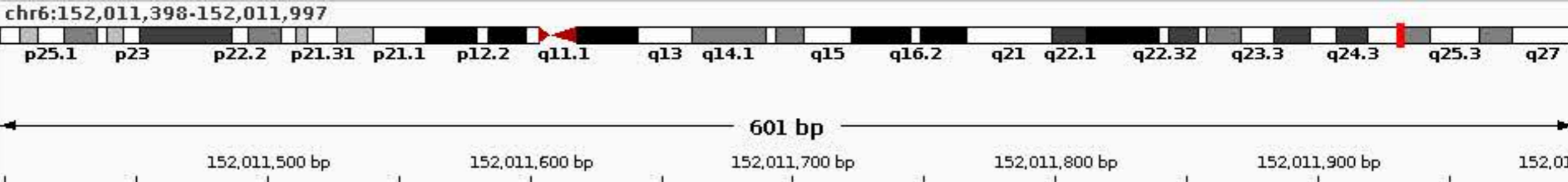
SL_1011-T2-B_mut.bam



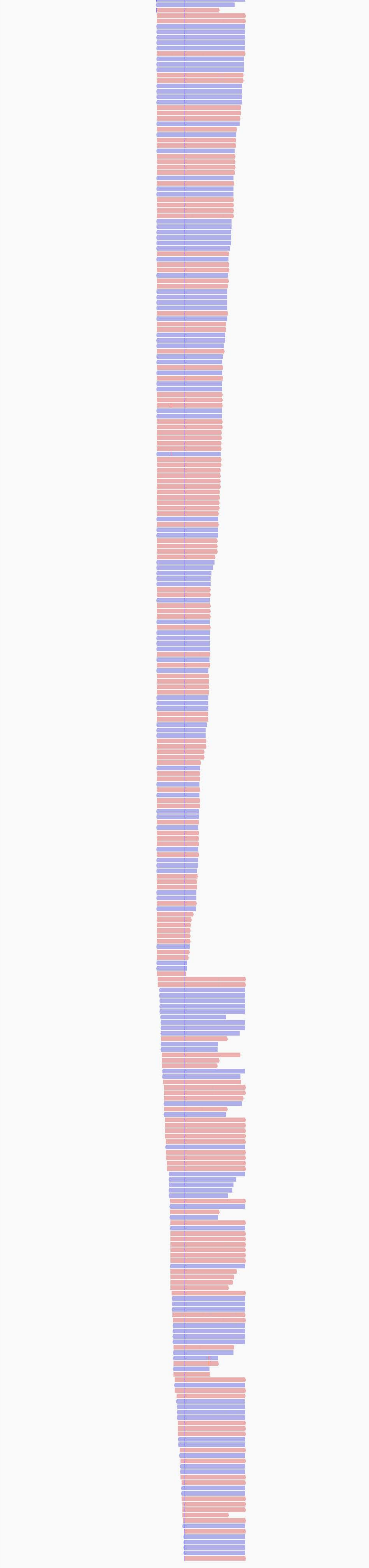
Sequence

Refseq Genes





SL_0003_L_FFPE_25-B_mut.barrage

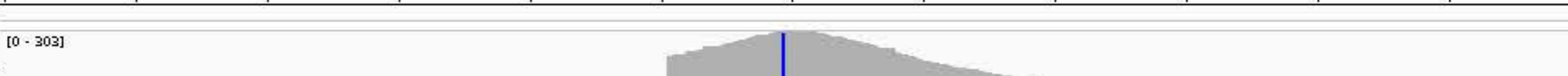
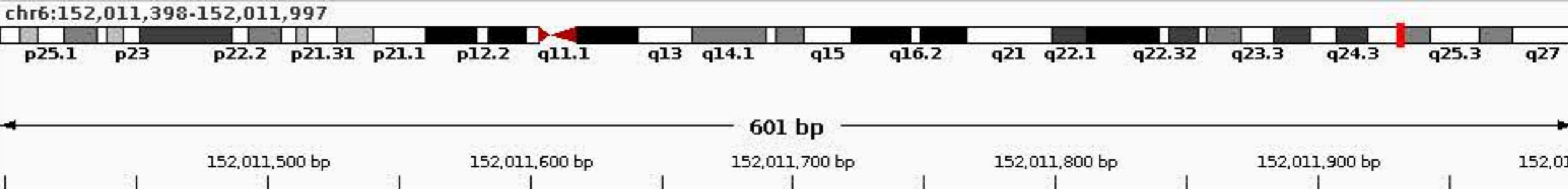


Sequence



Refseq Genes



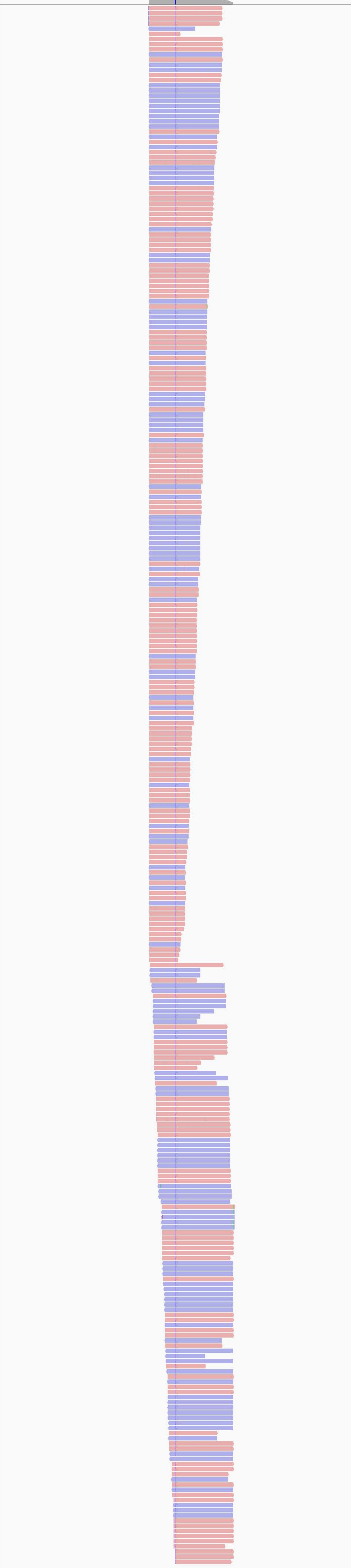


SL_1010-N2-B_mut.bam Coverage

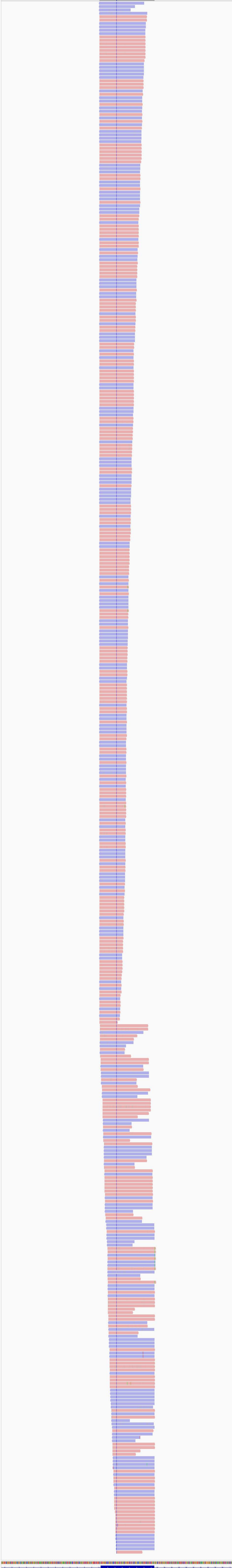
SL_1010-N2-B_mut.bam

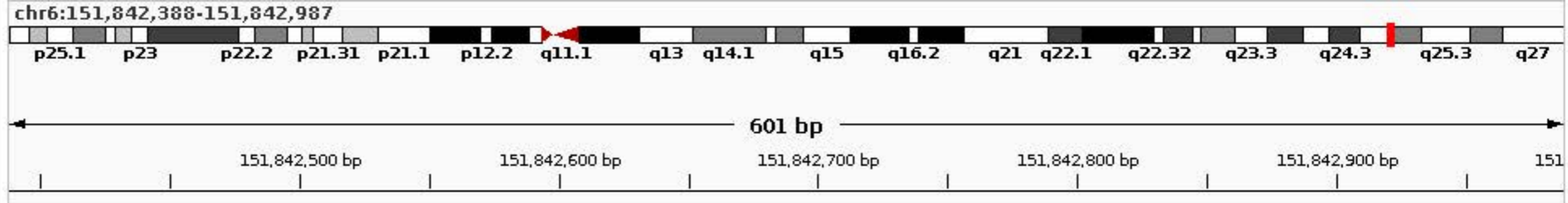
Sequence

Refseq Genes

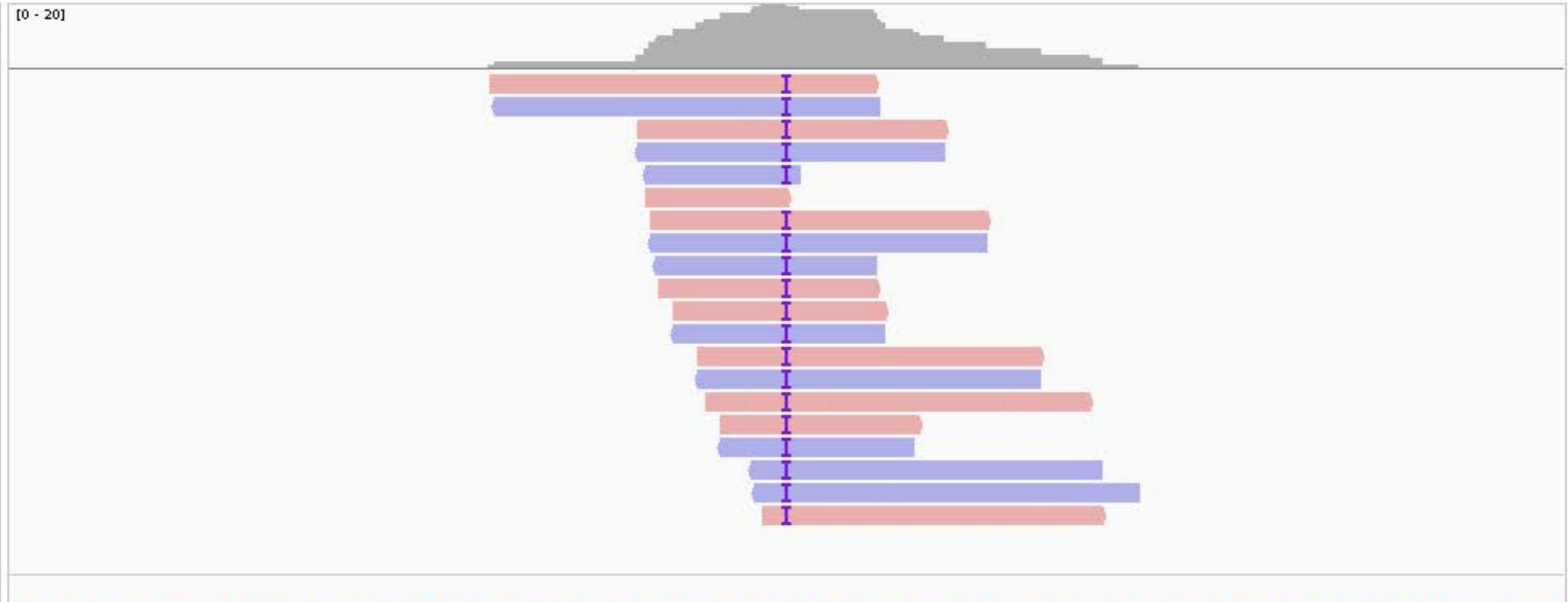


SL_1010-N3-8_mut.bam Coverage
SL_1010-N3-8_mut.bam
SL_1010-N3-8_mut.bam





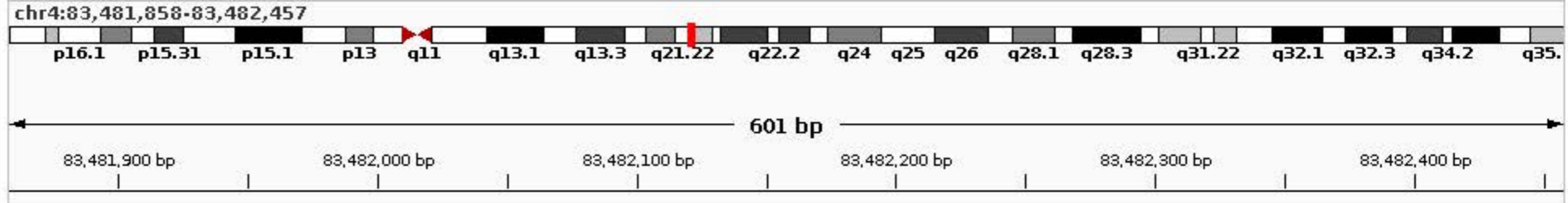
SL_1010_L_FFPE_17-B_mut.bam
range



SL_1010_L_FFPE_17-B_mut.bam

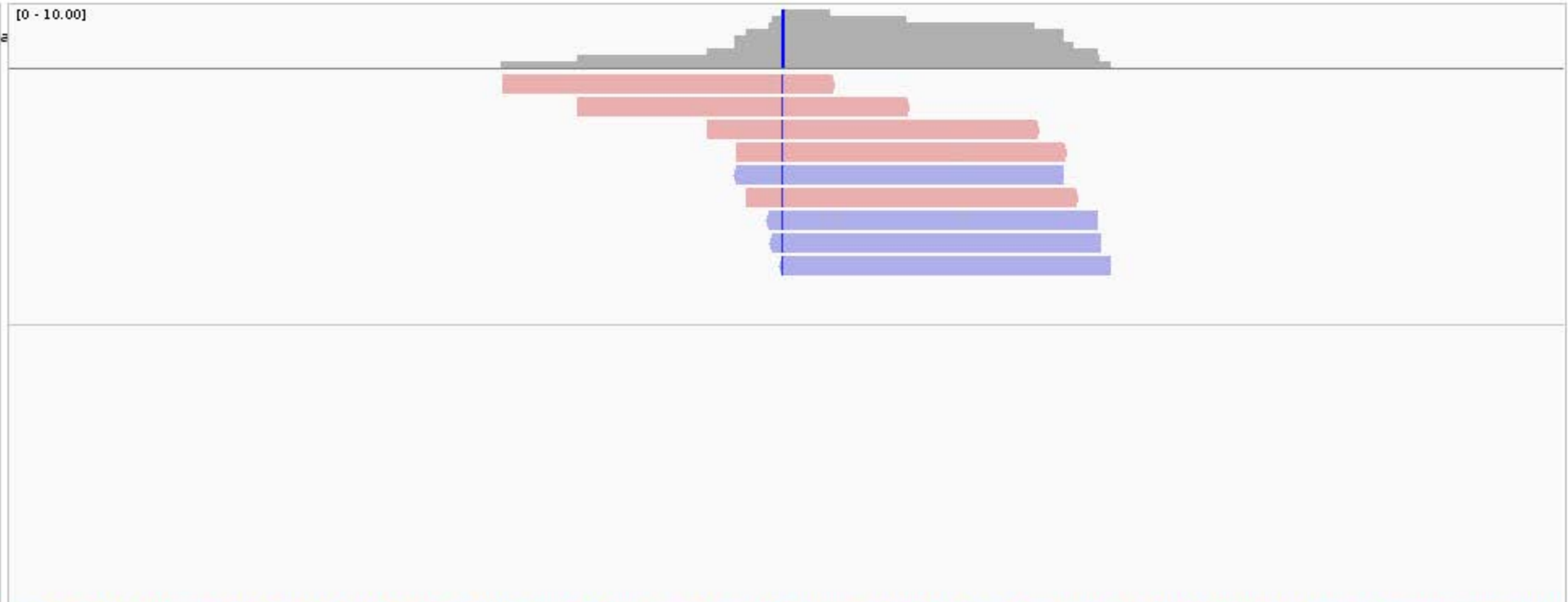
Sequence →
Refseq Genes





SL_0004-LT-6-B_mut.bam Coverage

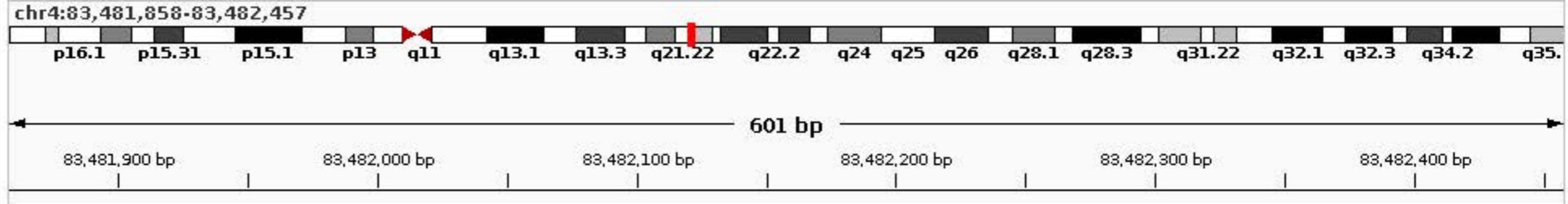
SL_0004-LT-6-B_mut.bam



Sequence

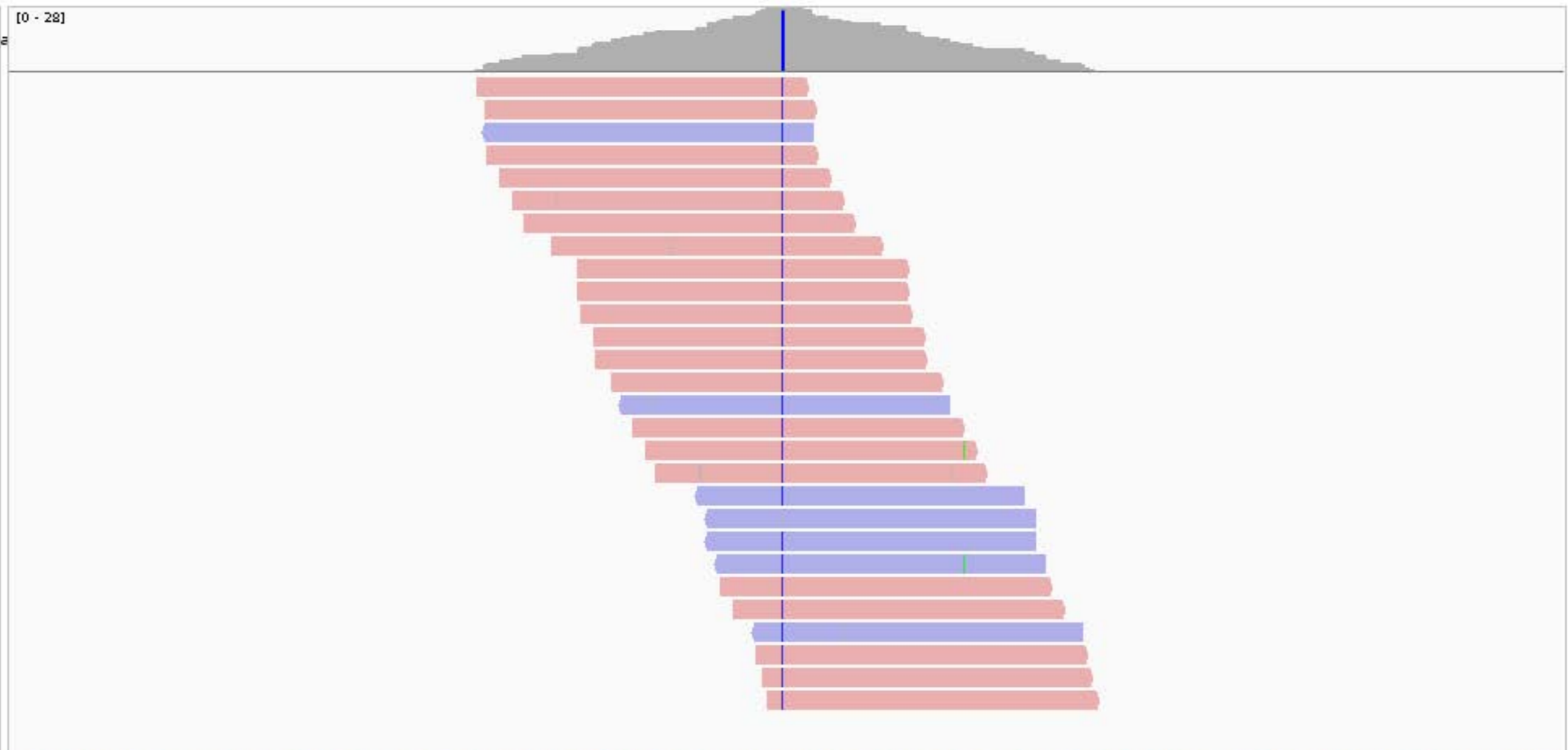
Refseq Genes





SL_0004-LT-8-B_mut.bam Coverage

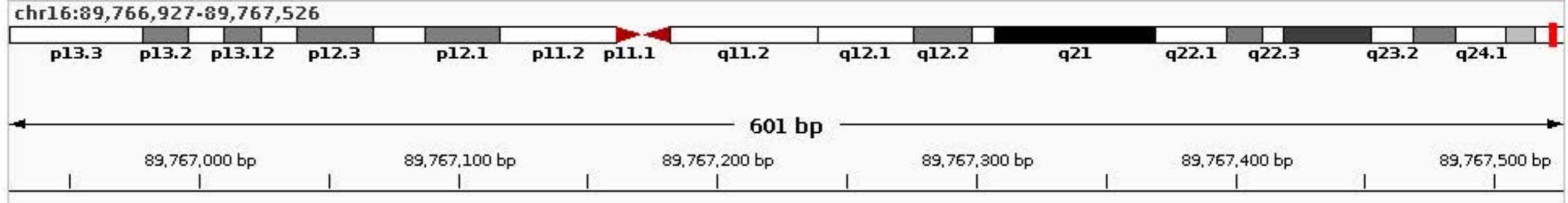
SL_0004-LT-8-B_mut.bam



Sequence →

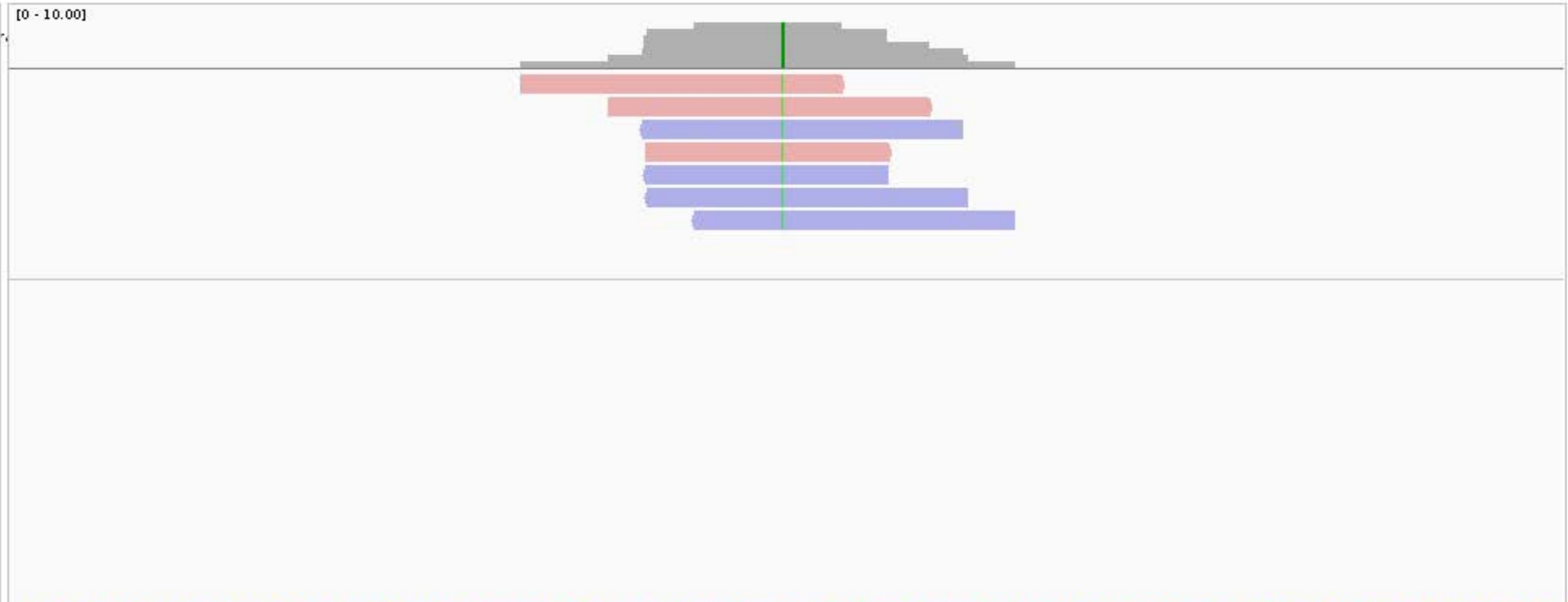
Refseq Genes





SL_0013_RT_2-B_mut.bam Cover

SL_0013_RT_2-B_mut.bam

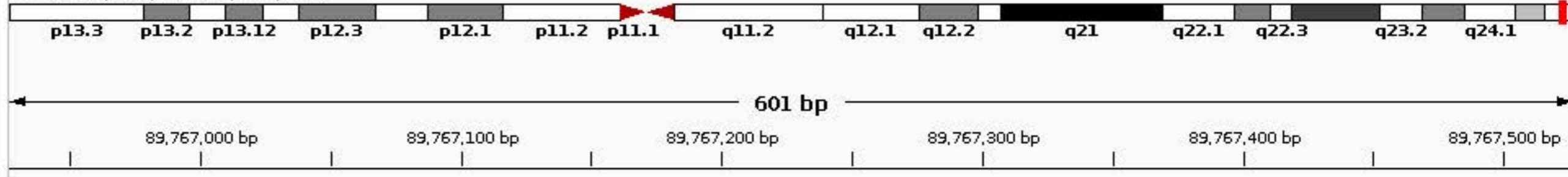


Sequence →

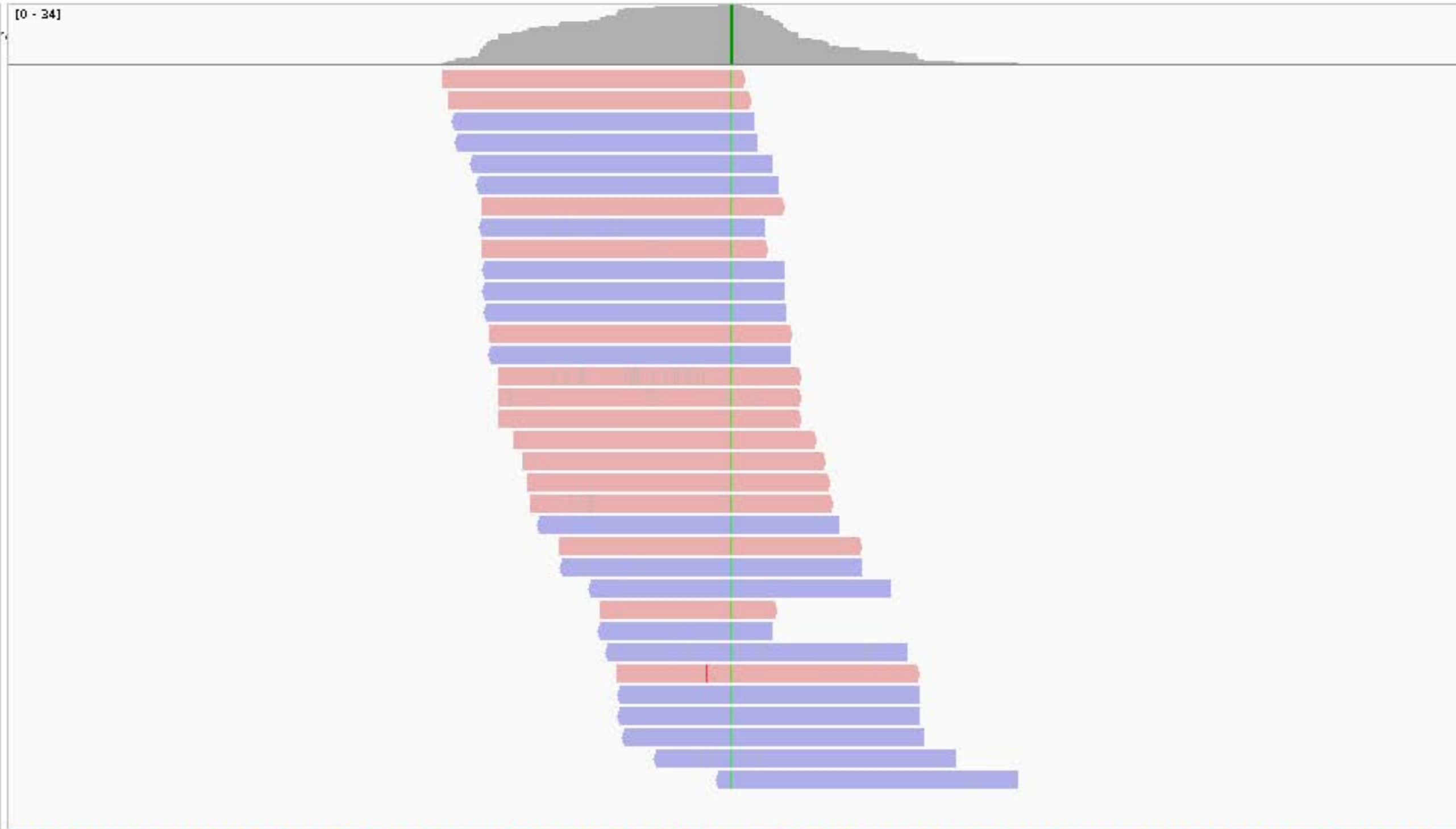
Refseq Genes

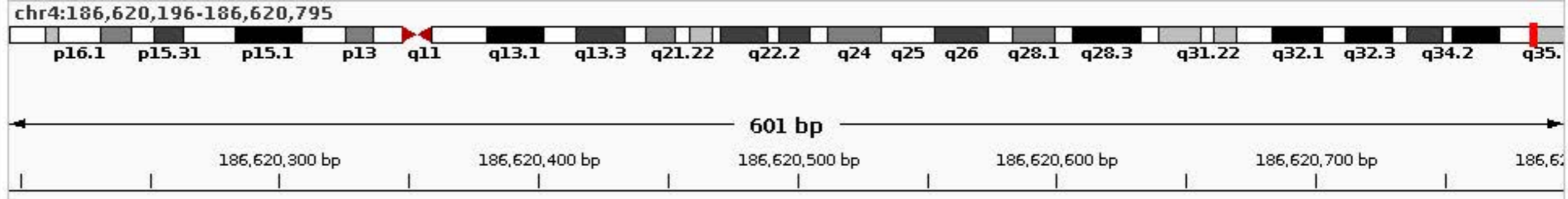


chr16:89,766,927-89,767,526



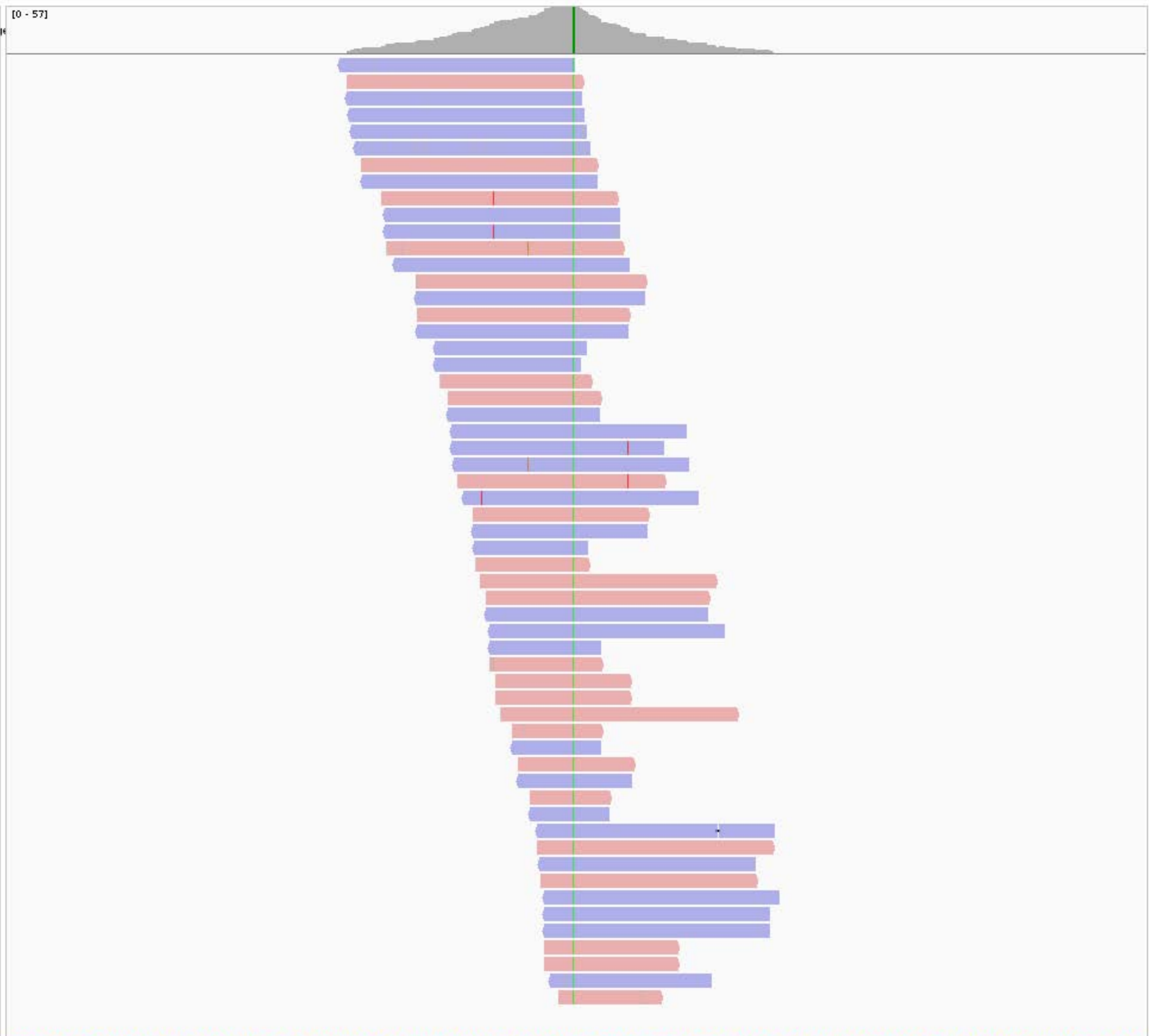
SL_0013_RT_6-B_mut.bam Cover.
SL_0013_RT_6-B_mut.bam
Sequence
Refseq Genes





SL_1011-T1-B_mut.bam Coverage

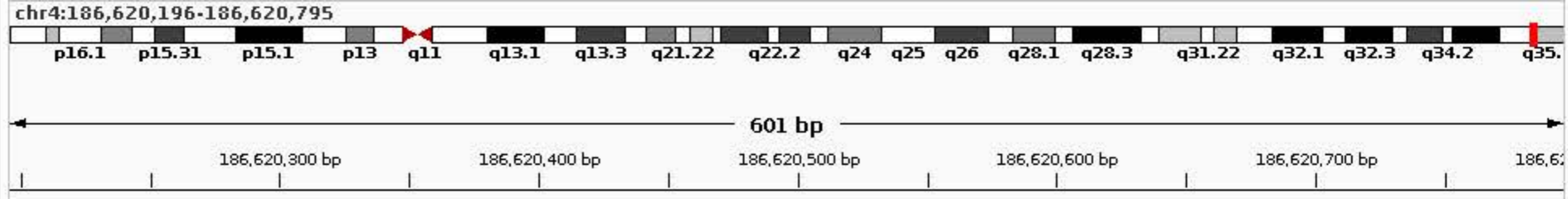
SL_1011-T1-B_mut.bam



Sequence

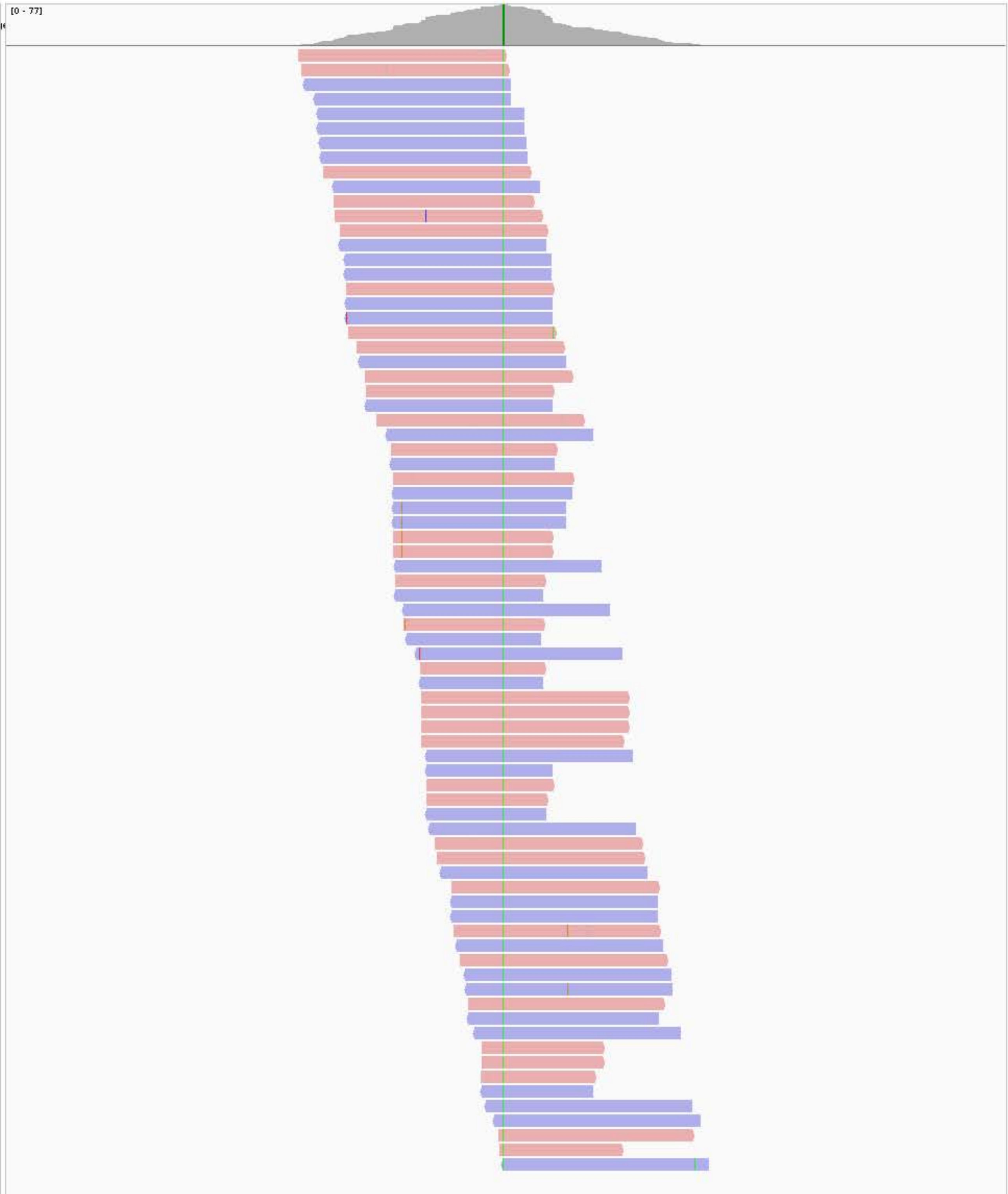
Refseq Genes





SL_1011-T2-B_mut.bam Coverage

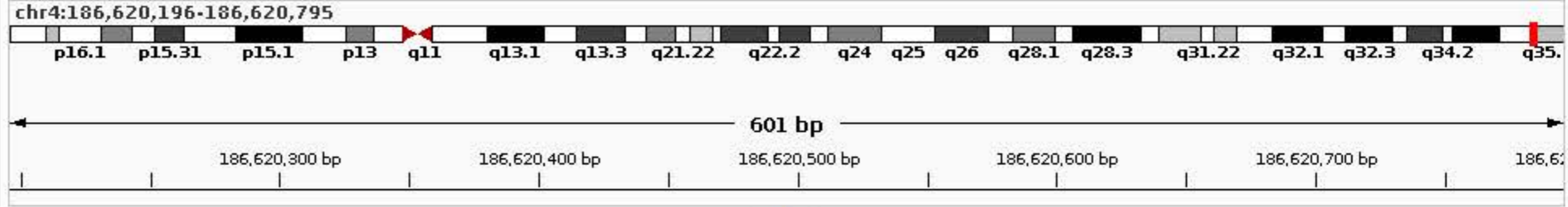
SL_1011-T2-B_mut.bam



Sequence →

Refseq Genes



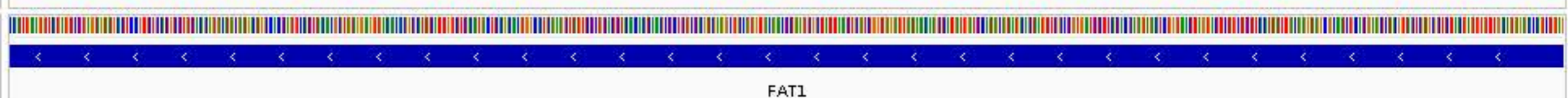
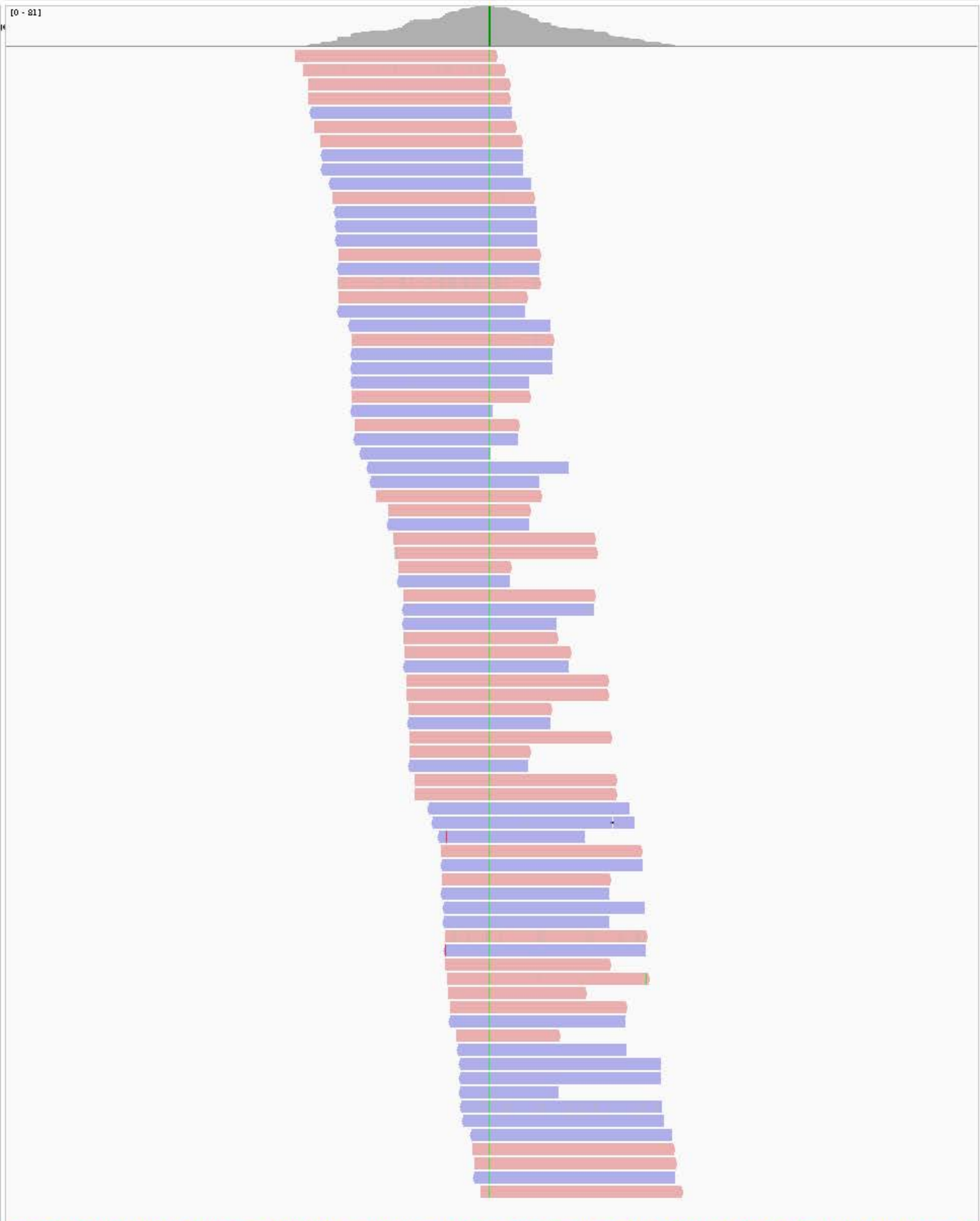


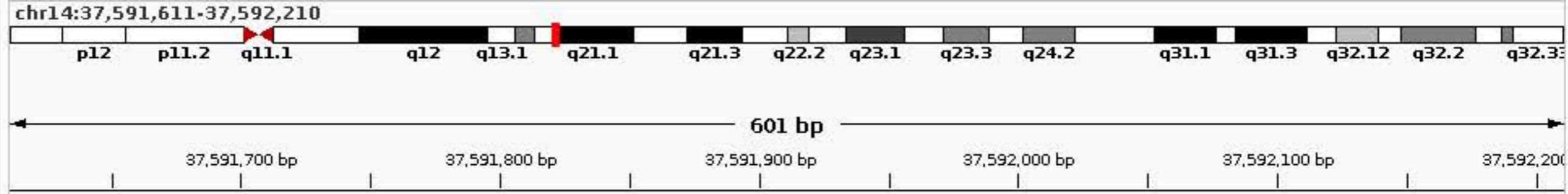
SL_1011-T3-B_mut.bam Coverage

SL_1011-T3-B_mut.bam

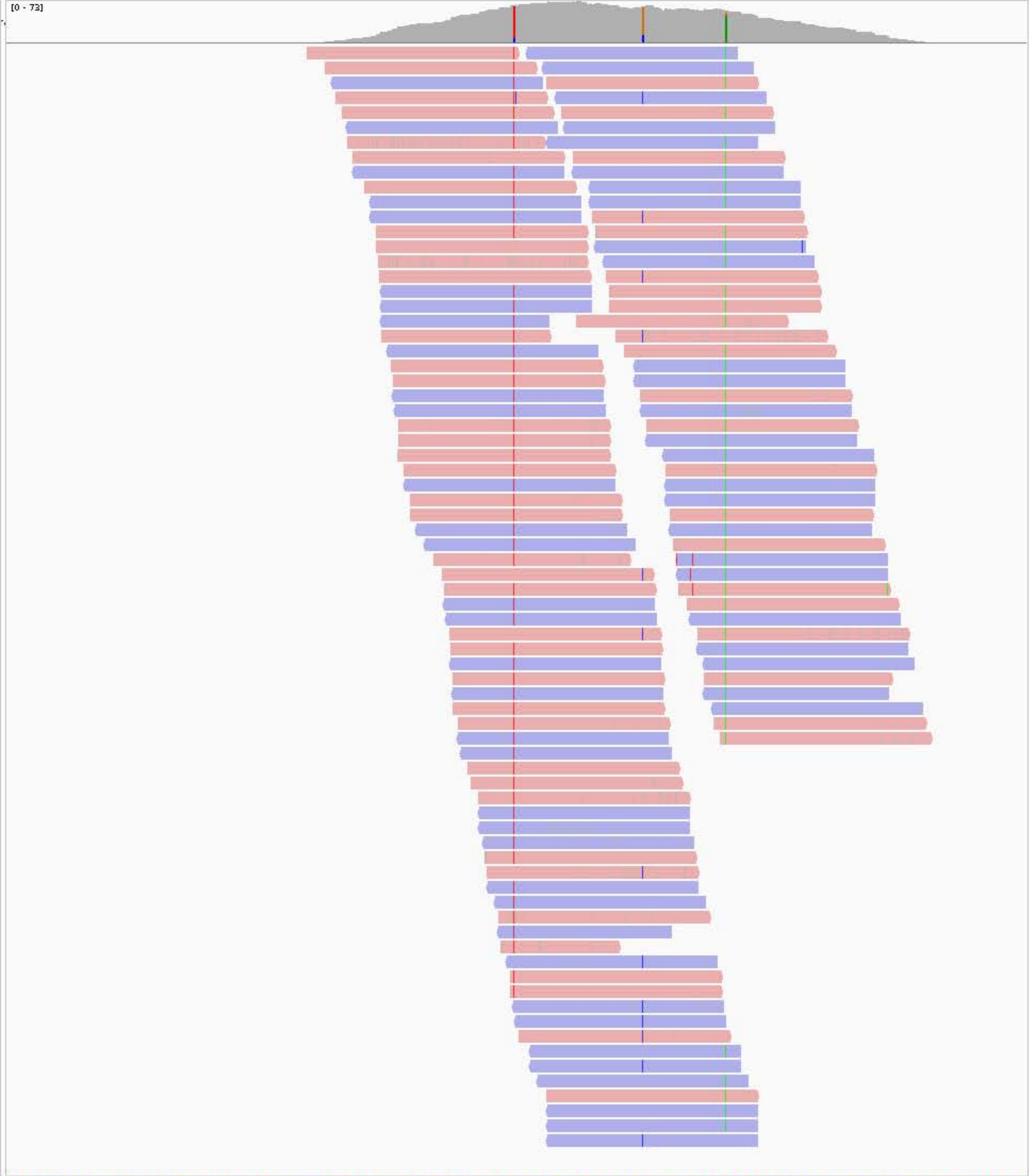
Sequence →

Refseq Genes





SL_0013_RT_4-B_mut.bam Cover



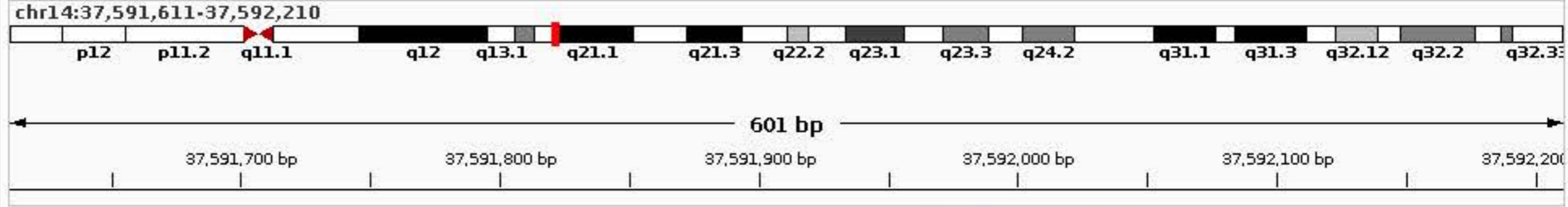
SL_0013_RT_4-B_mut.bam

Sequence →

Refseq Genes



FOXA1

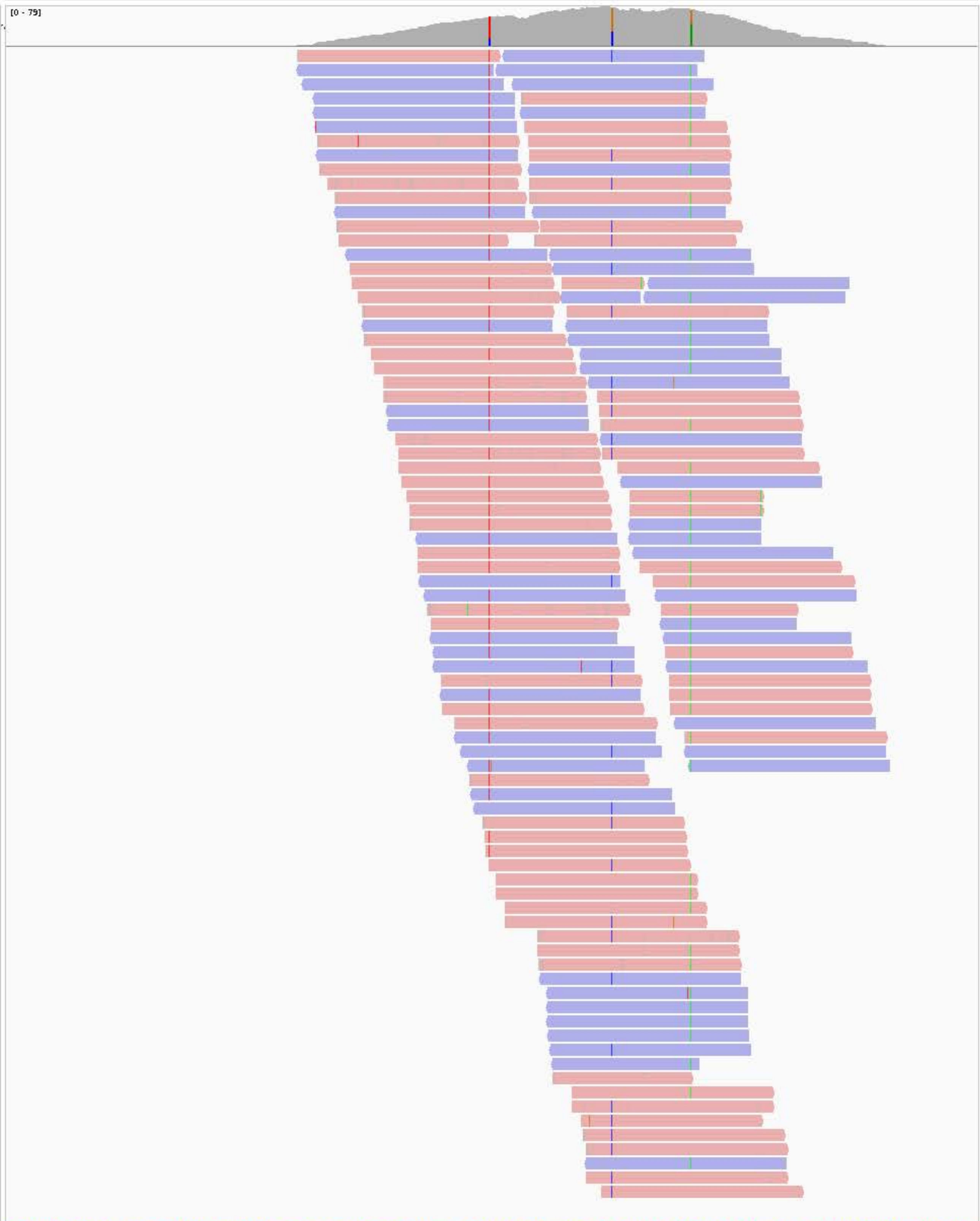


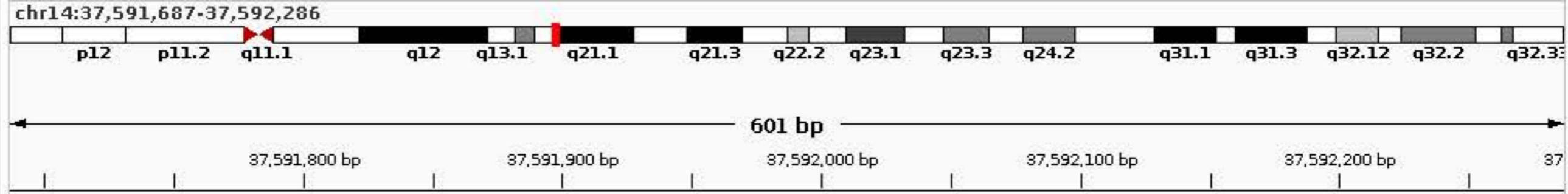
SL_0013_RT_5-B_mut.bam Cover

SL_0013_RT_5-B_mut.bam

Sequence →

Refseq Genes





SL_0013_RT_4-B_mut.bam Cover

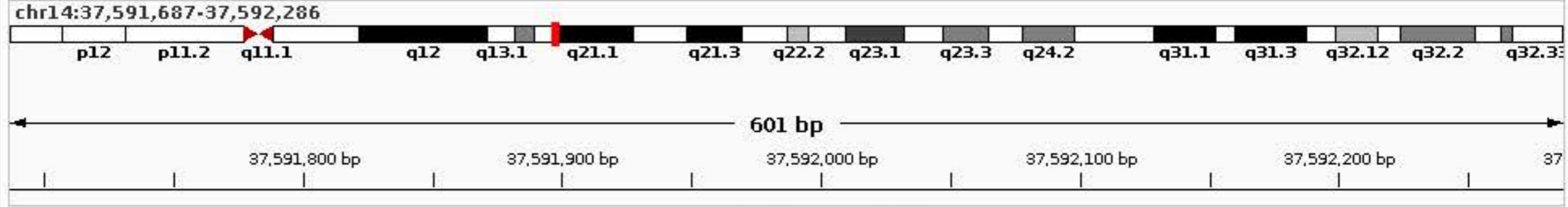


SL_0013_RT_4-B_mut.bam

Sequence →
Refseq Genes



FOXA1

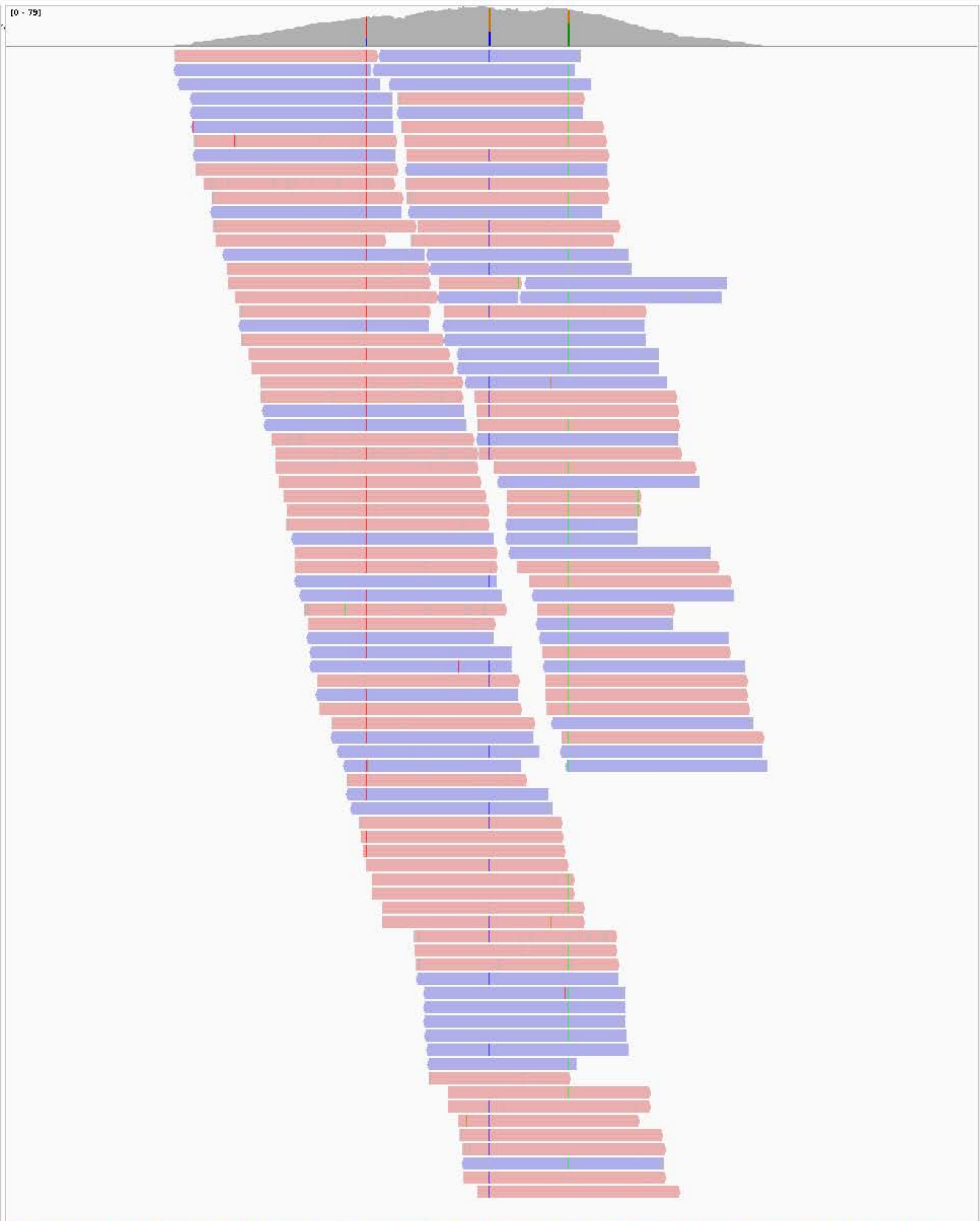


SL_0013_RT_5-B_mut.bam Cover

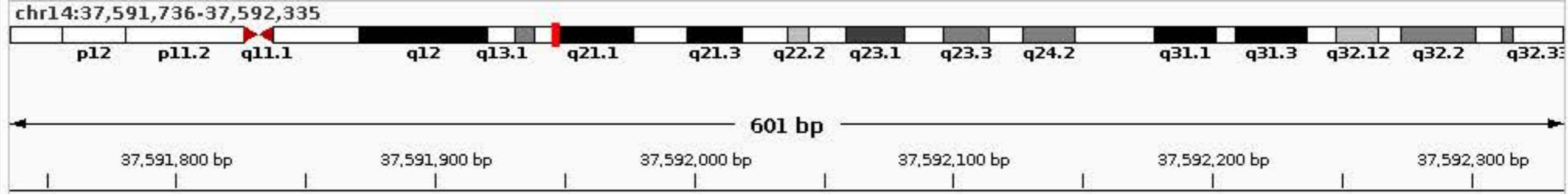
SL_0013_RT_5-B_mut.bam

Sequence →

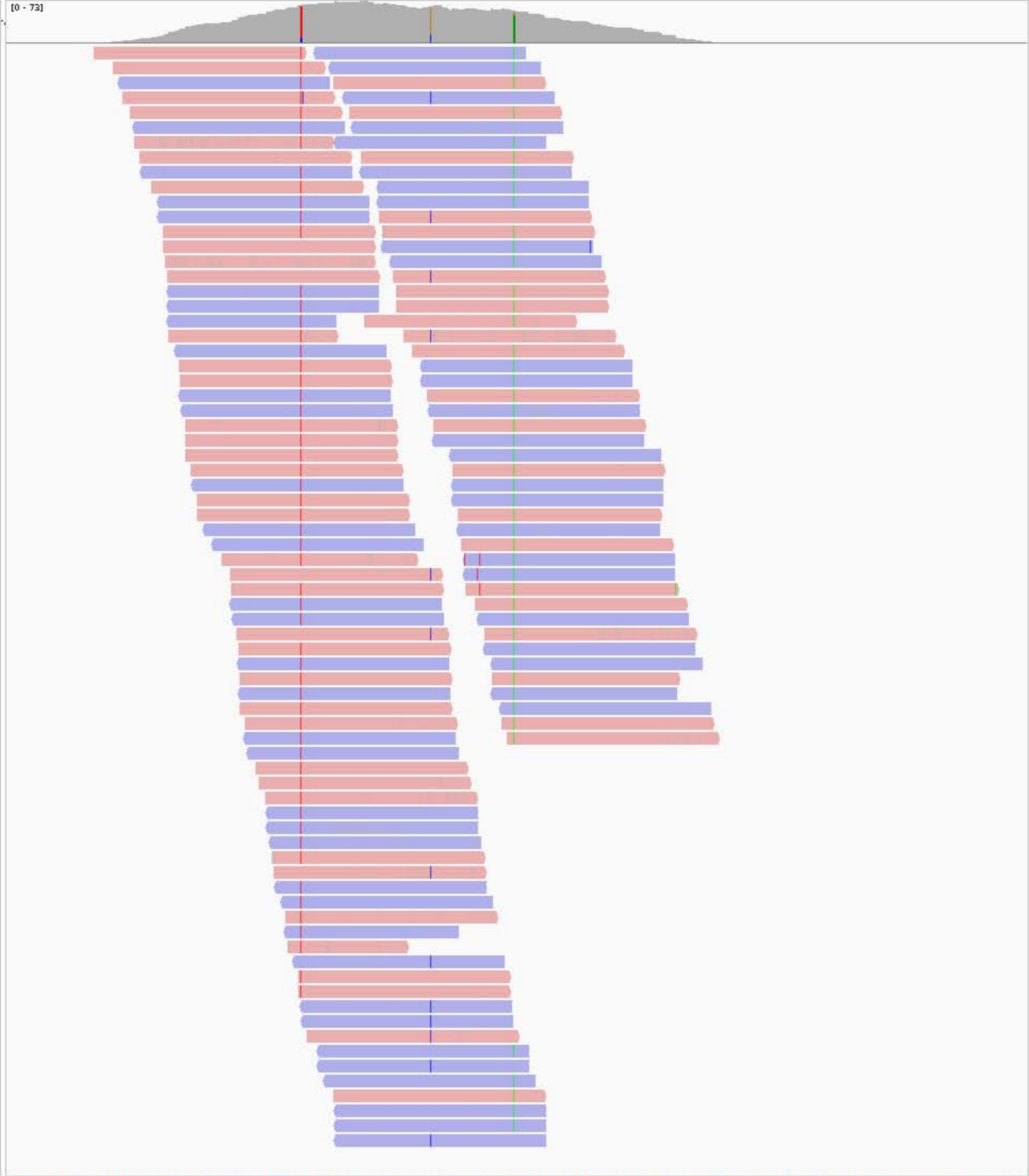
Refseq Genes



FOXA1



SL_0013_RT_4-B_mut.bam Cover

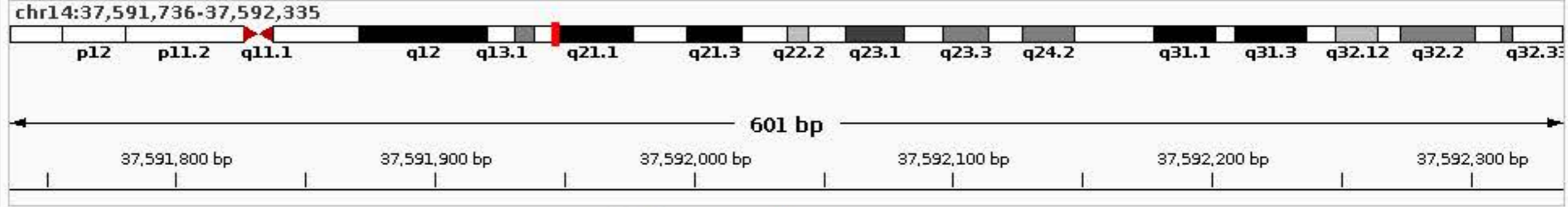


SL_0013_RT_4-B_mut.bam

Sequence →
Refseq Genes



FOXA1

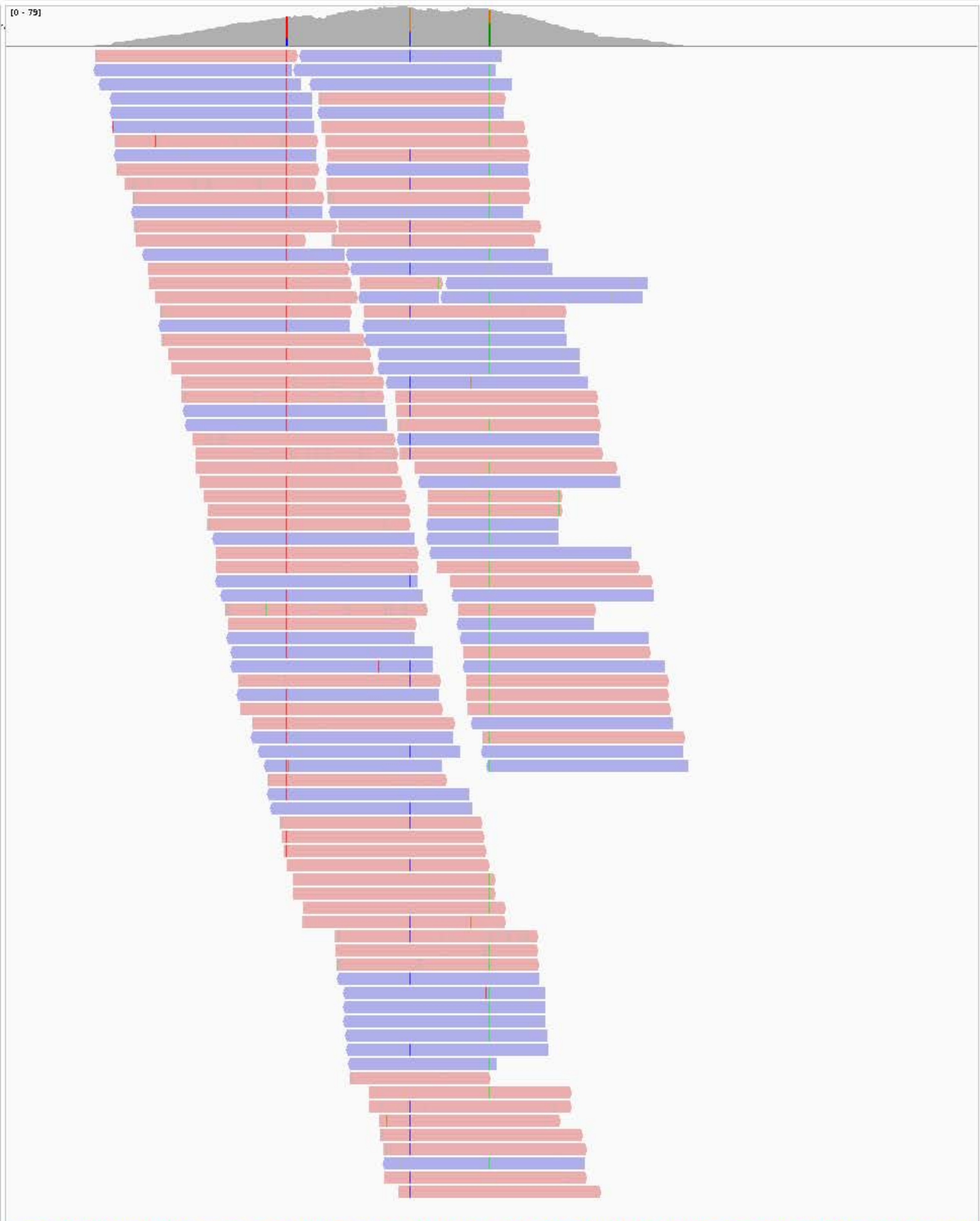


SL_0013_RT_5-B_mut.bam Cover

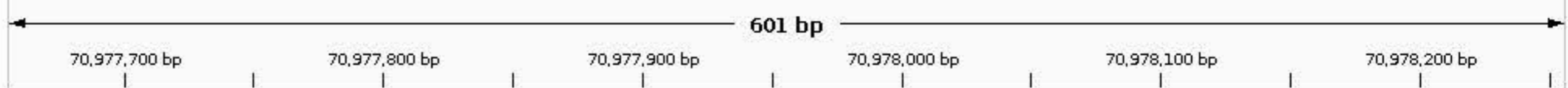
SL_0013_RT_5-B_mut.bam

Sequence →

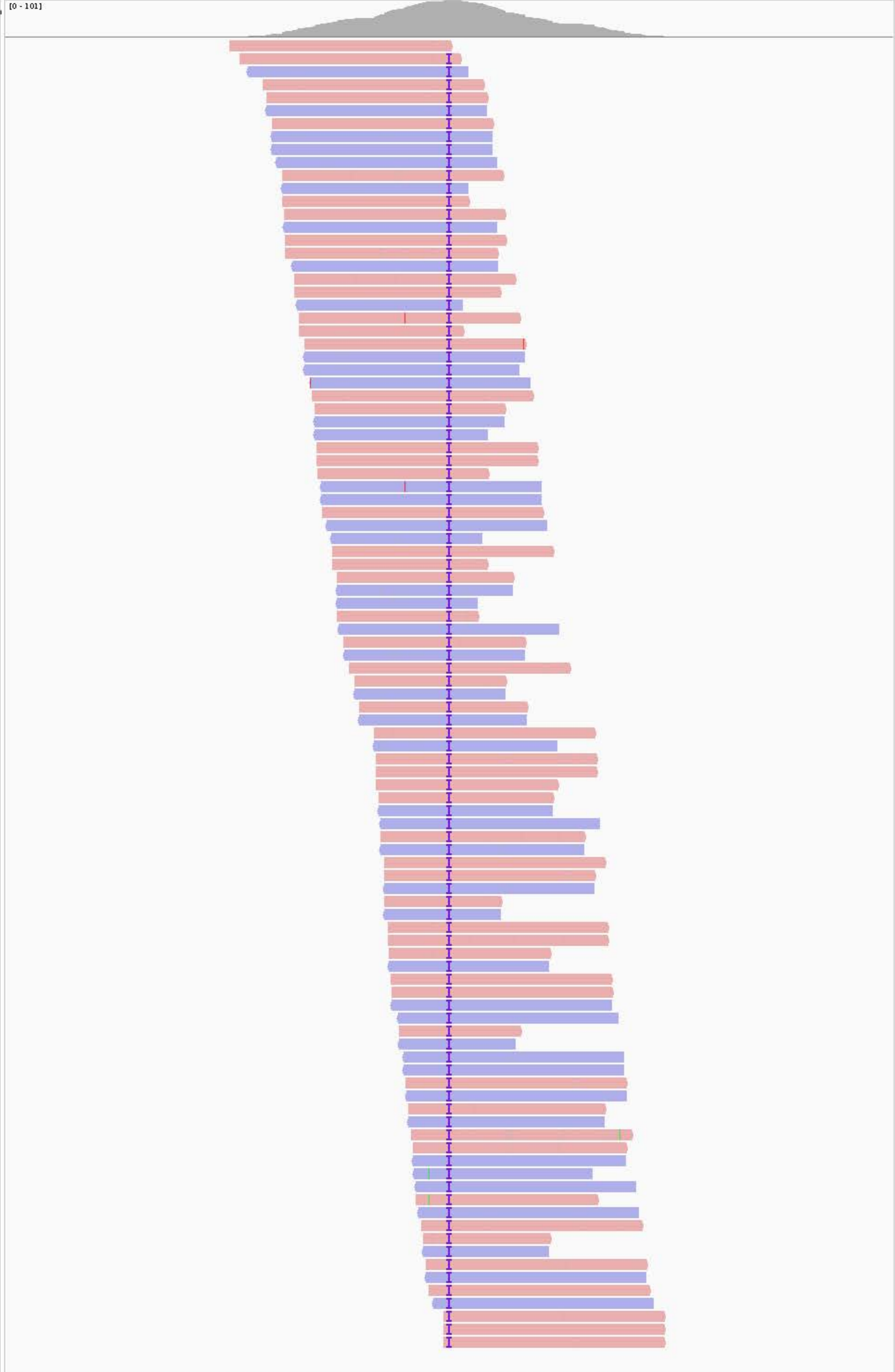
Refseq Genes



chr3:70,977,656-70,978,255
p26.1 p24.3 p23 p22.1 p21.2 p14.2 p13 p12.2 q11.1 q12.3 q13.2 q21.1 q22.2 q24 q25.2 q26.1 q26.32 q27.3 q29



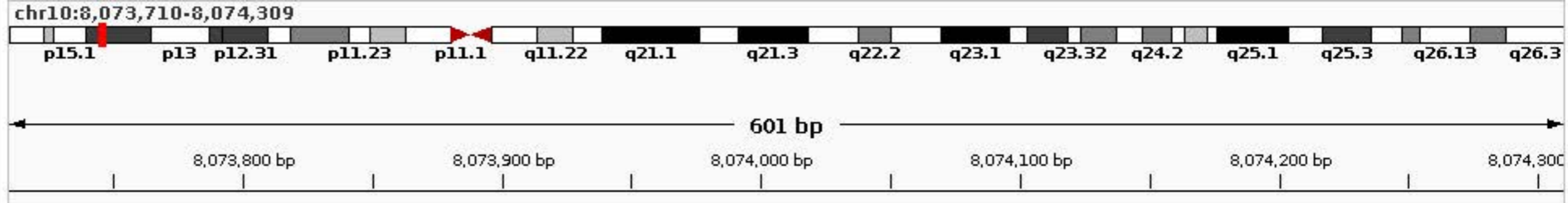
SL_0003_R_FFPE_52-B_mut.bam
rage



SL_0003_R_FFPE_52-B_mut.ba
m

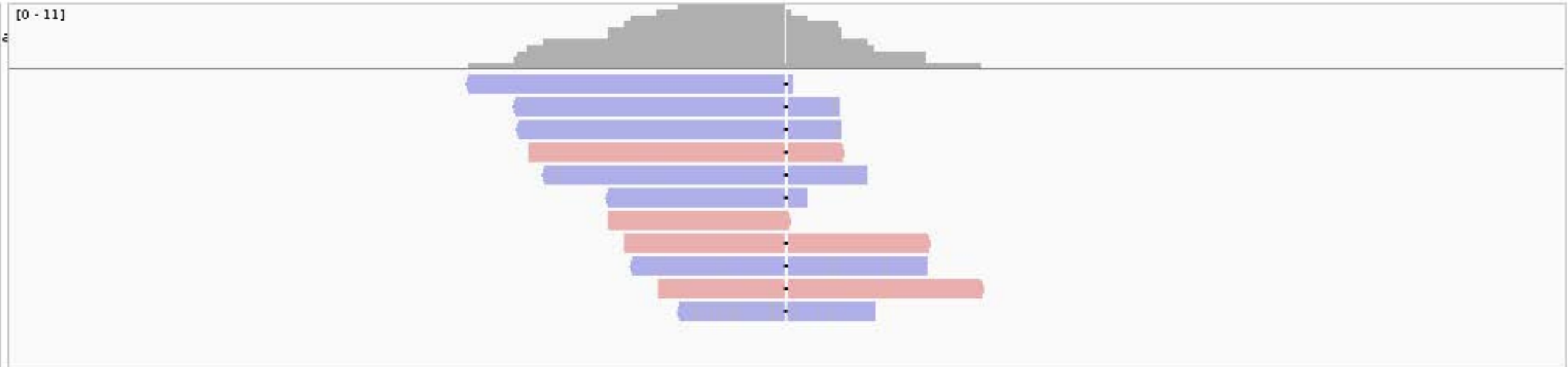
Sequence
Refseq Genes





SL_0009-LT_1-B_mut.bam Coverage

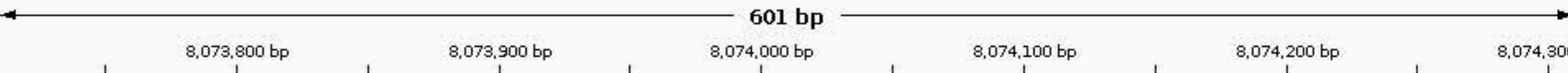
SL_0009-LT_1-B_mut.bam



Sequence

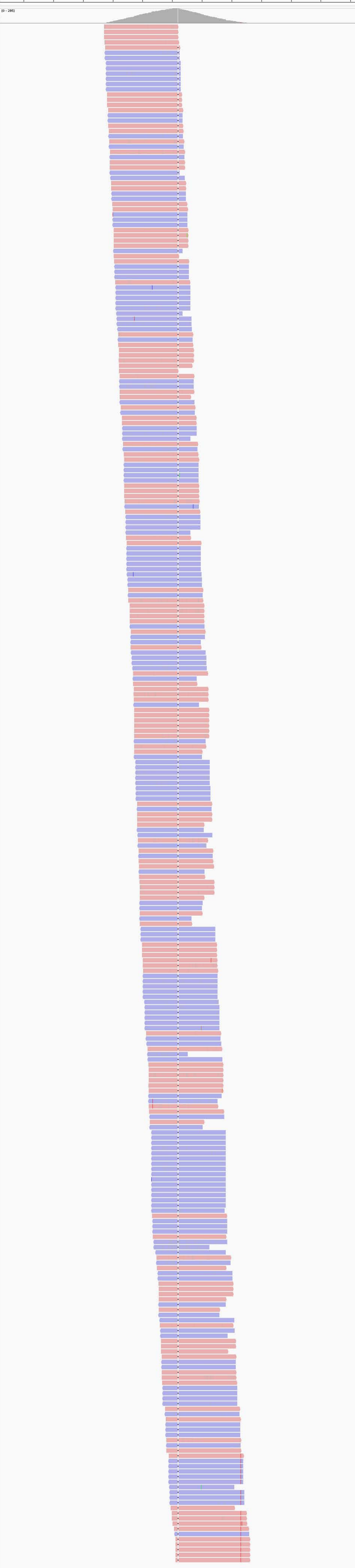
Refseq Genes





SL_0009-LT_3-B_mut.bam Coverage

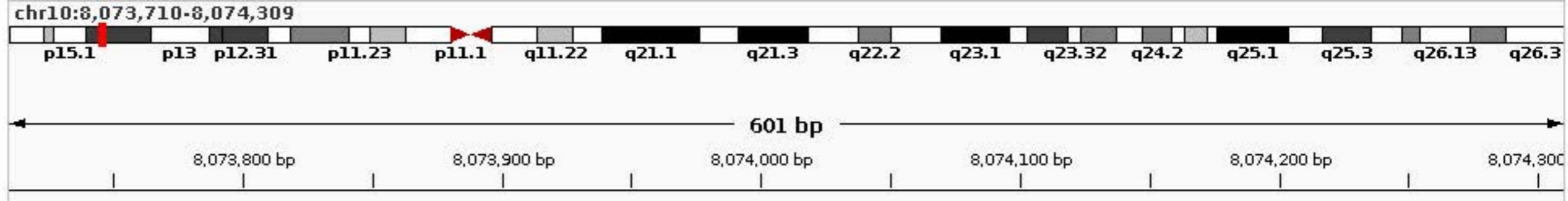
SL_0009-LT_3-B_mut.bam



Sequence
Refseq Genes

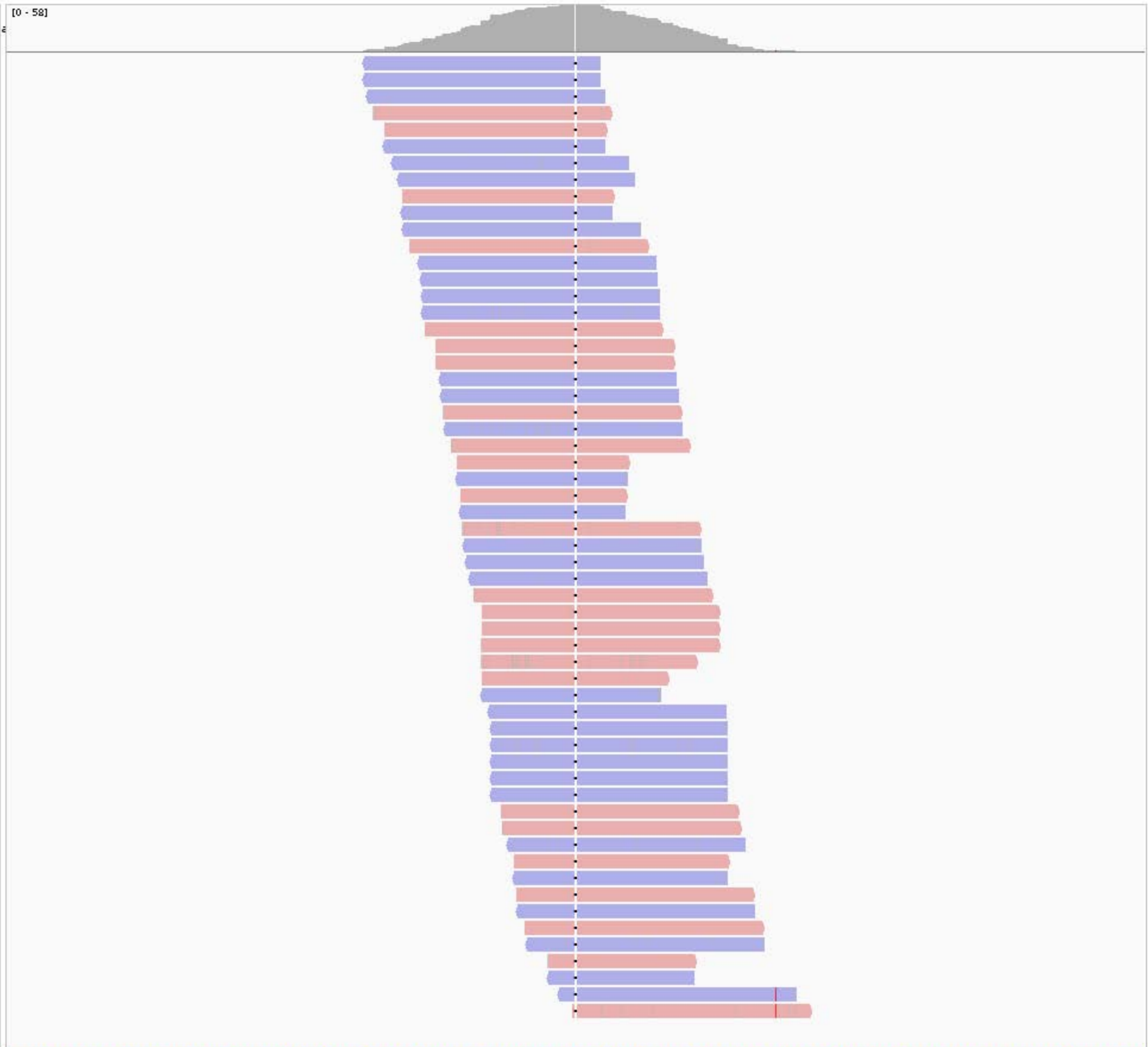


GAT3



SL_0009-LT_9-B_mut.bam Coverage

SL_0009-LT_9-B_mut.bam

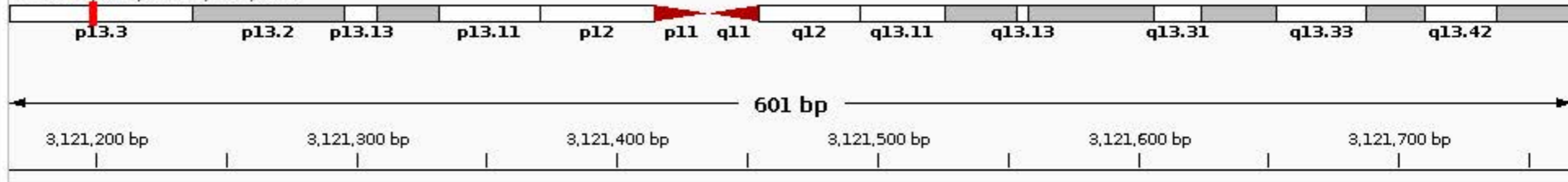


Sequence

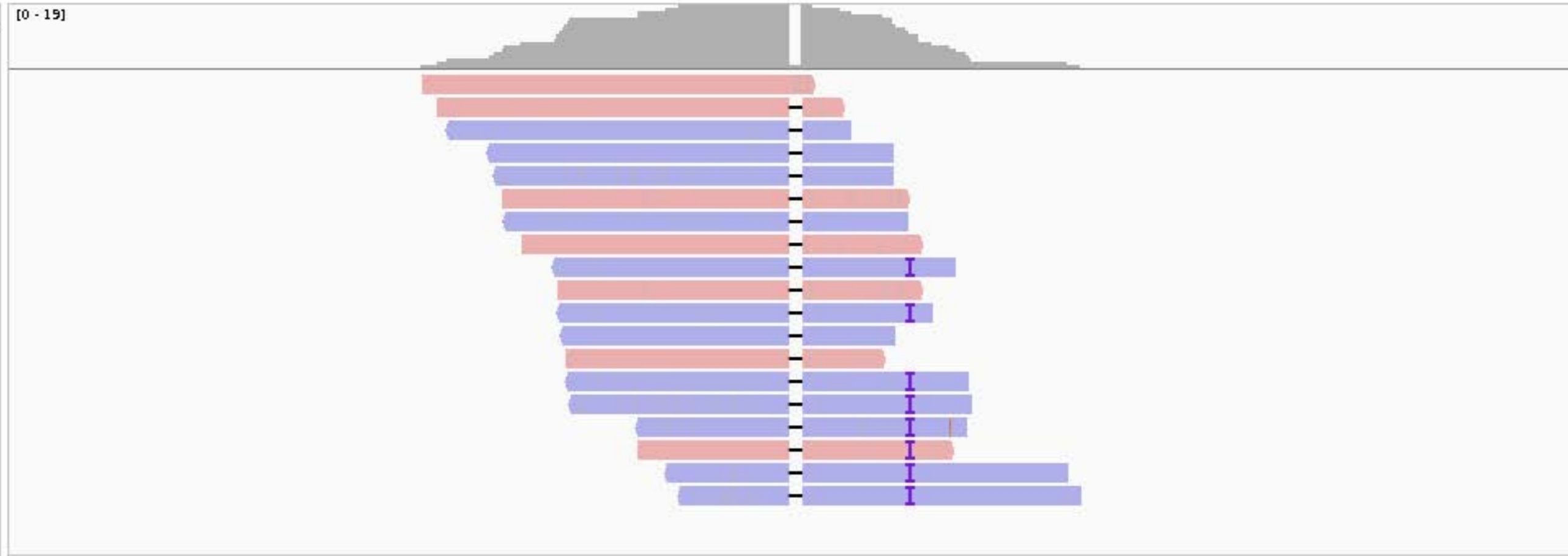
Refseq Genes



chr19:3,121,167-3,121,766



SL_0003_L_FFPE_11-B_mut.barrage



SL_0003_L_FFPE_11-B_mut.bam

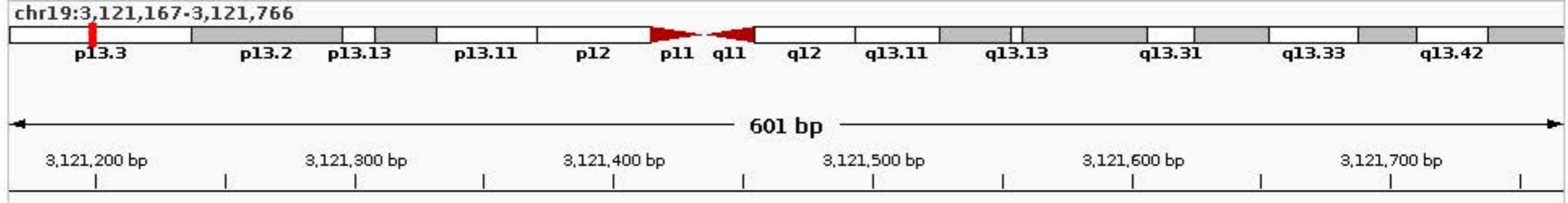
Sequence →



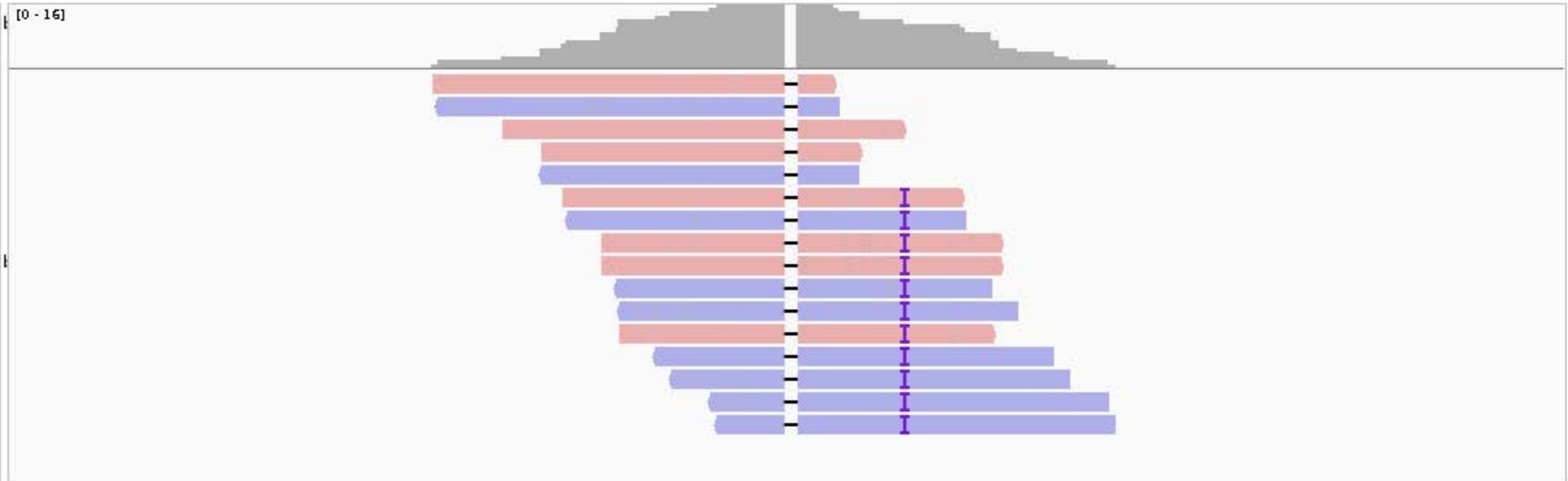
Refseq Genes



GNAT1



SL_0003_L_FFPE_40_skin-B_mut l
Coverage

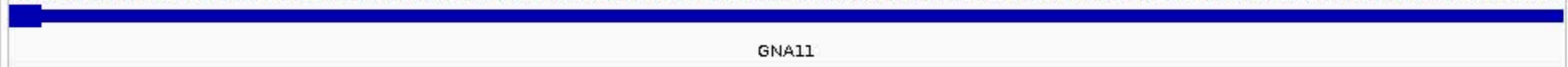


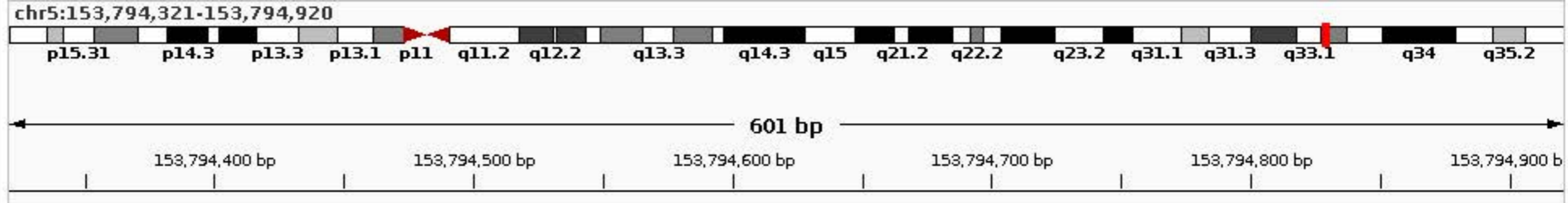
SL_0003_L_FFPE_40_skin-B_mut l

Sequence →



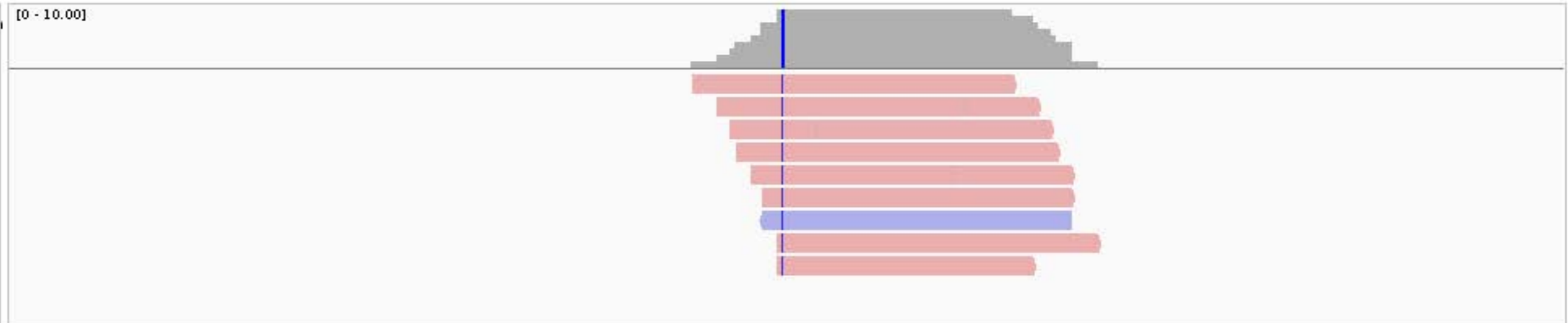
Refseq Genes





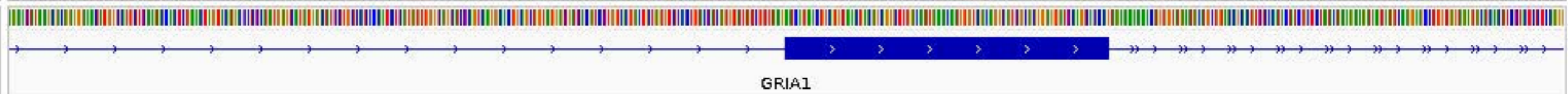
SL_1012_R_10_FFPE-B_mut.bam
rage

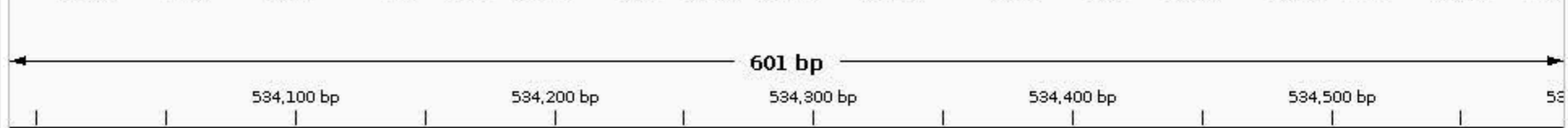
SL_1012_R_10_FFPE-B_mut.ba
m



Sequence →

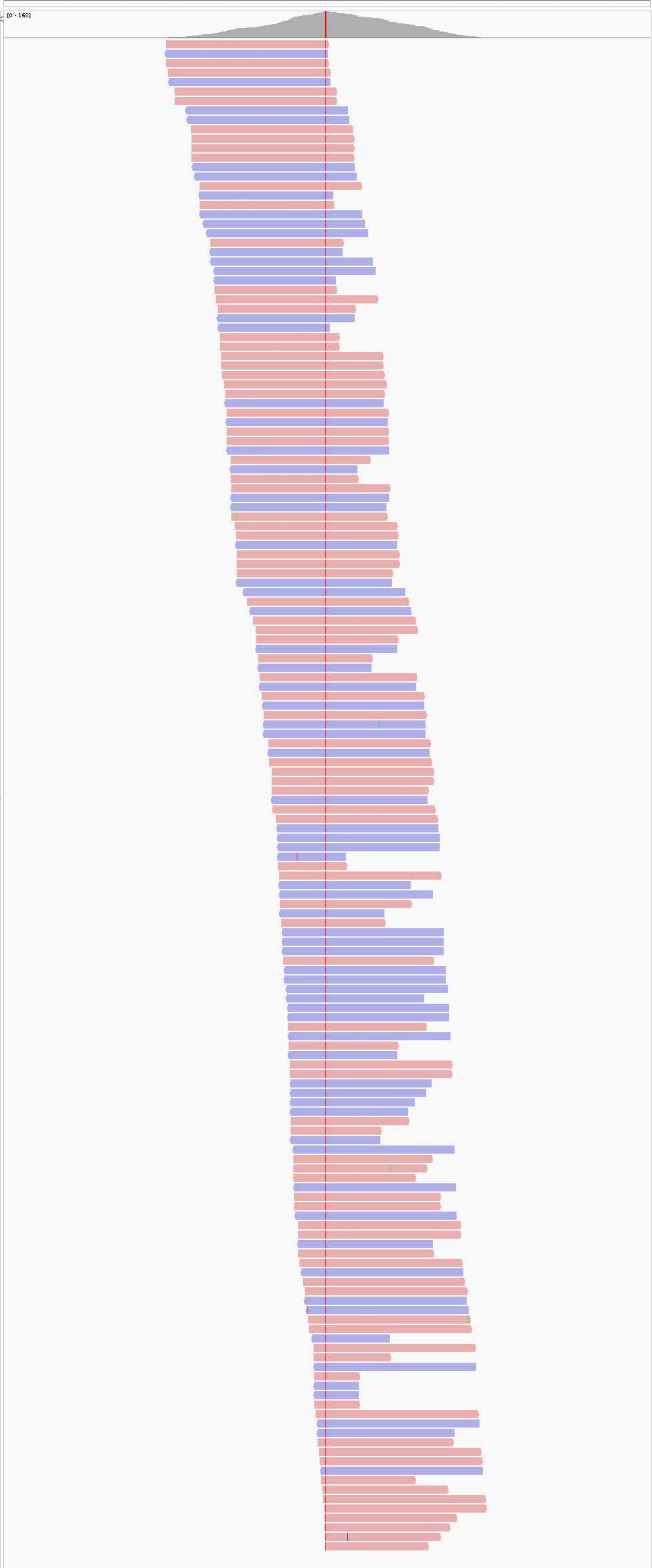
Refseq Genes





SL_1010_R_FFPE_2-B_mut.bam
age

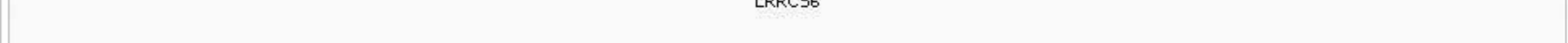
SL_1010_R_FFPE_2-B_mut.bam



Sequence

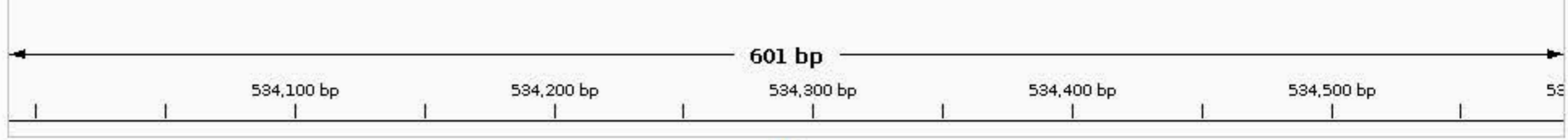


Refseq Genes



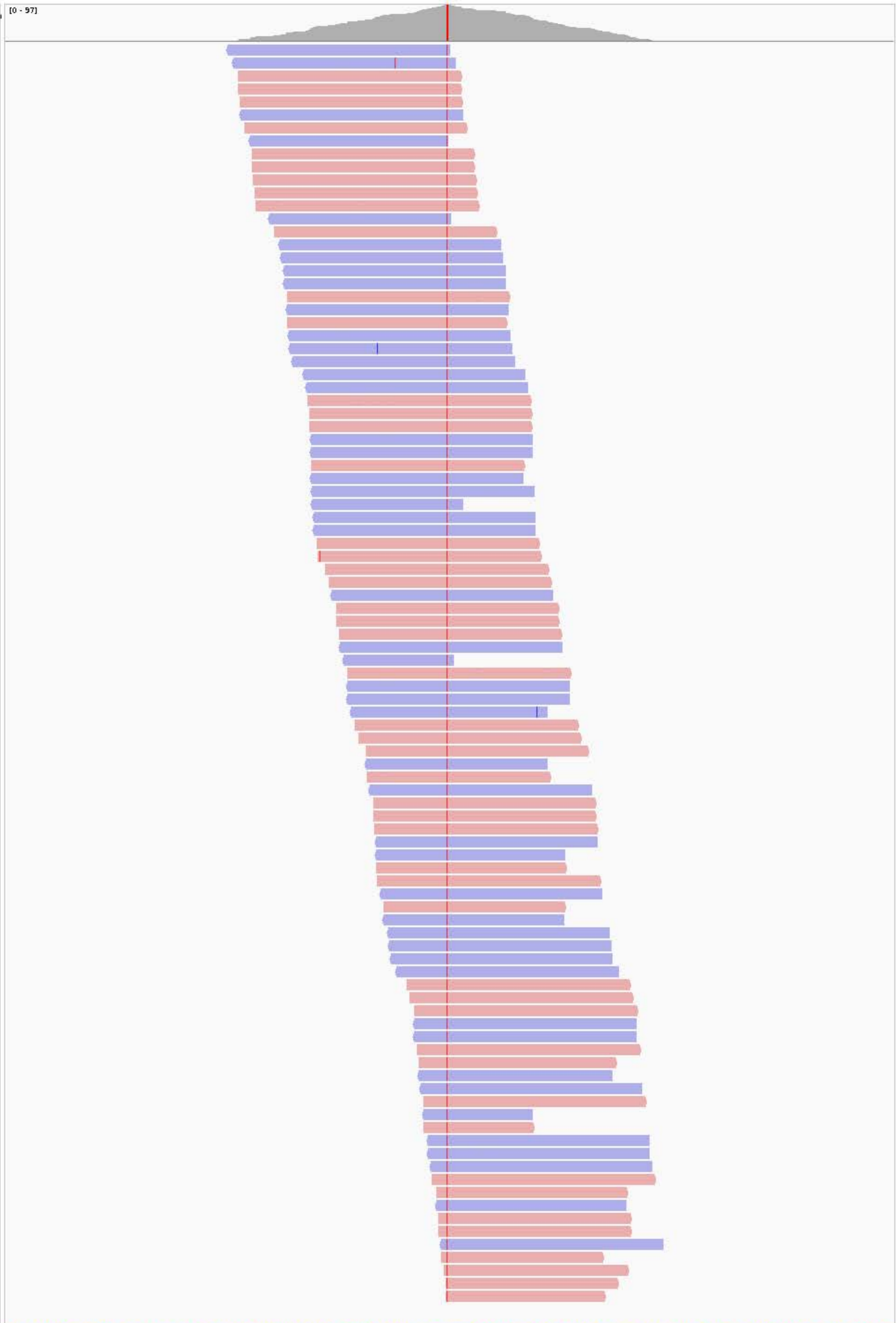
LRR56

chr11:533,990-534,589



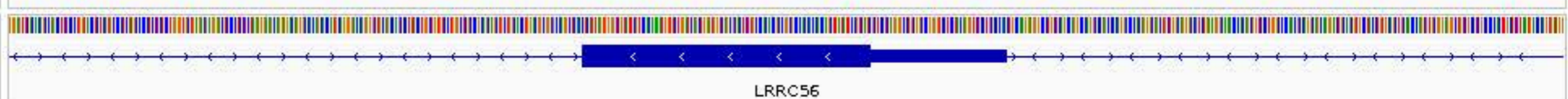
SL_1010_R_FFPE_11-B_mut.bam
rage

SL_1010_R_FFPE_11-B_mut.ba
m

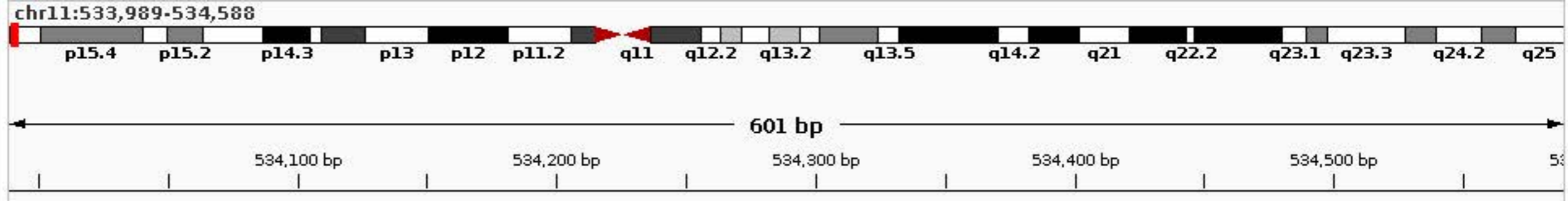


Sequence →

Refseq Genes

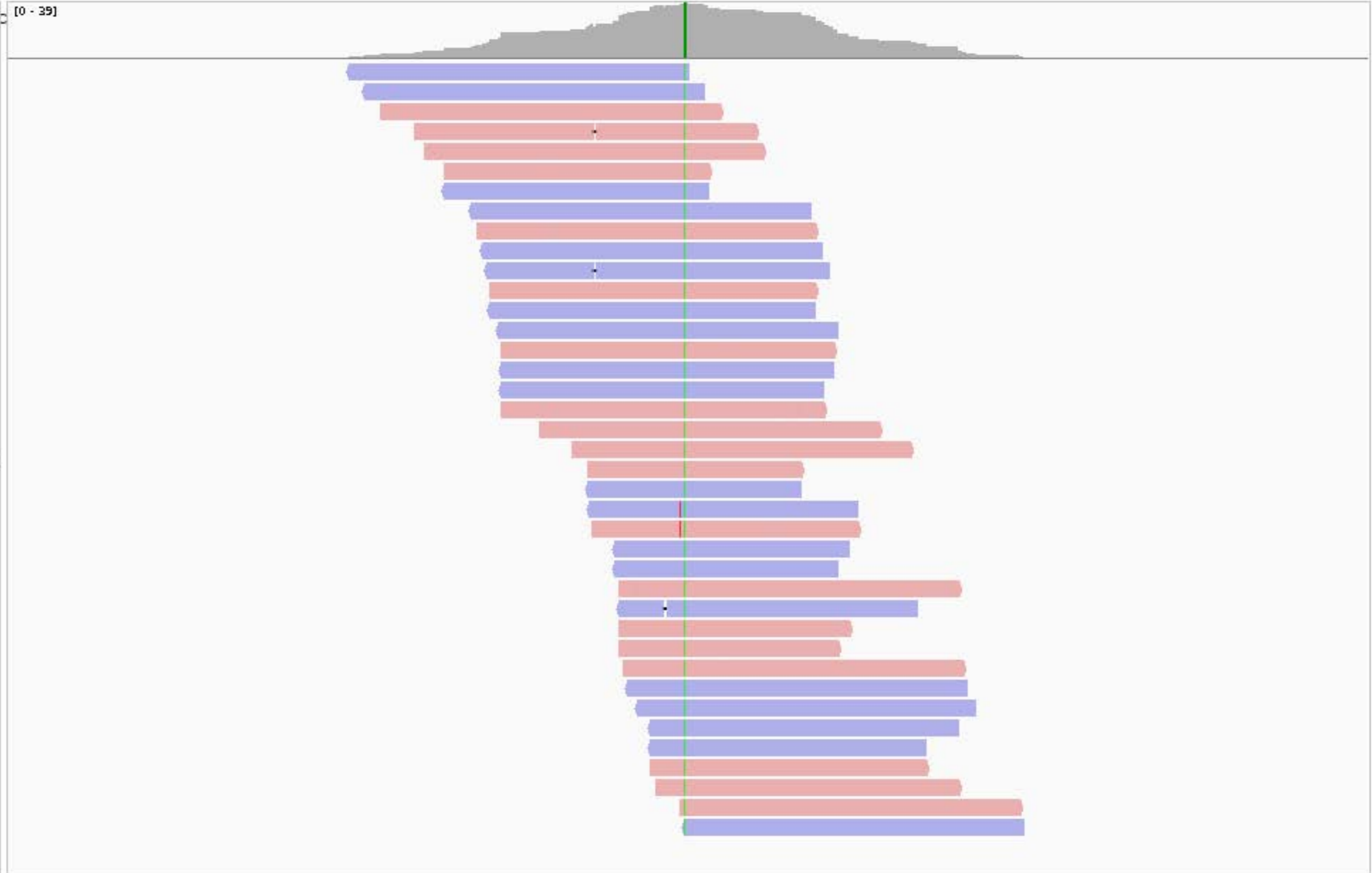


LRR56



SL_1006_L_FFPE_7-B_mut.bam
age

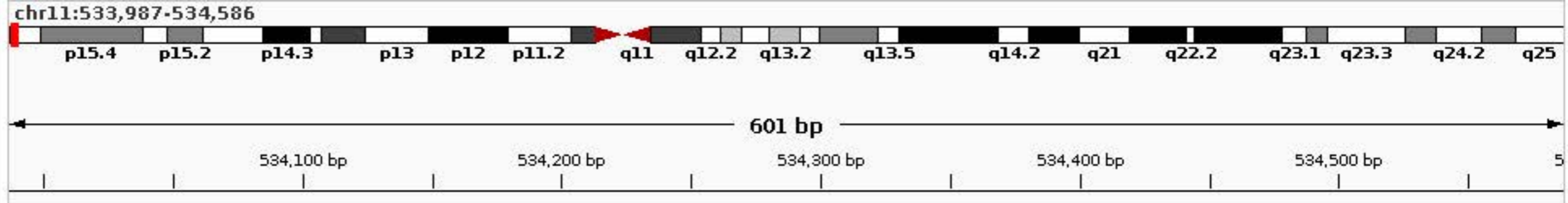
SL_1006_L_FFPE_7-B_mut.bam



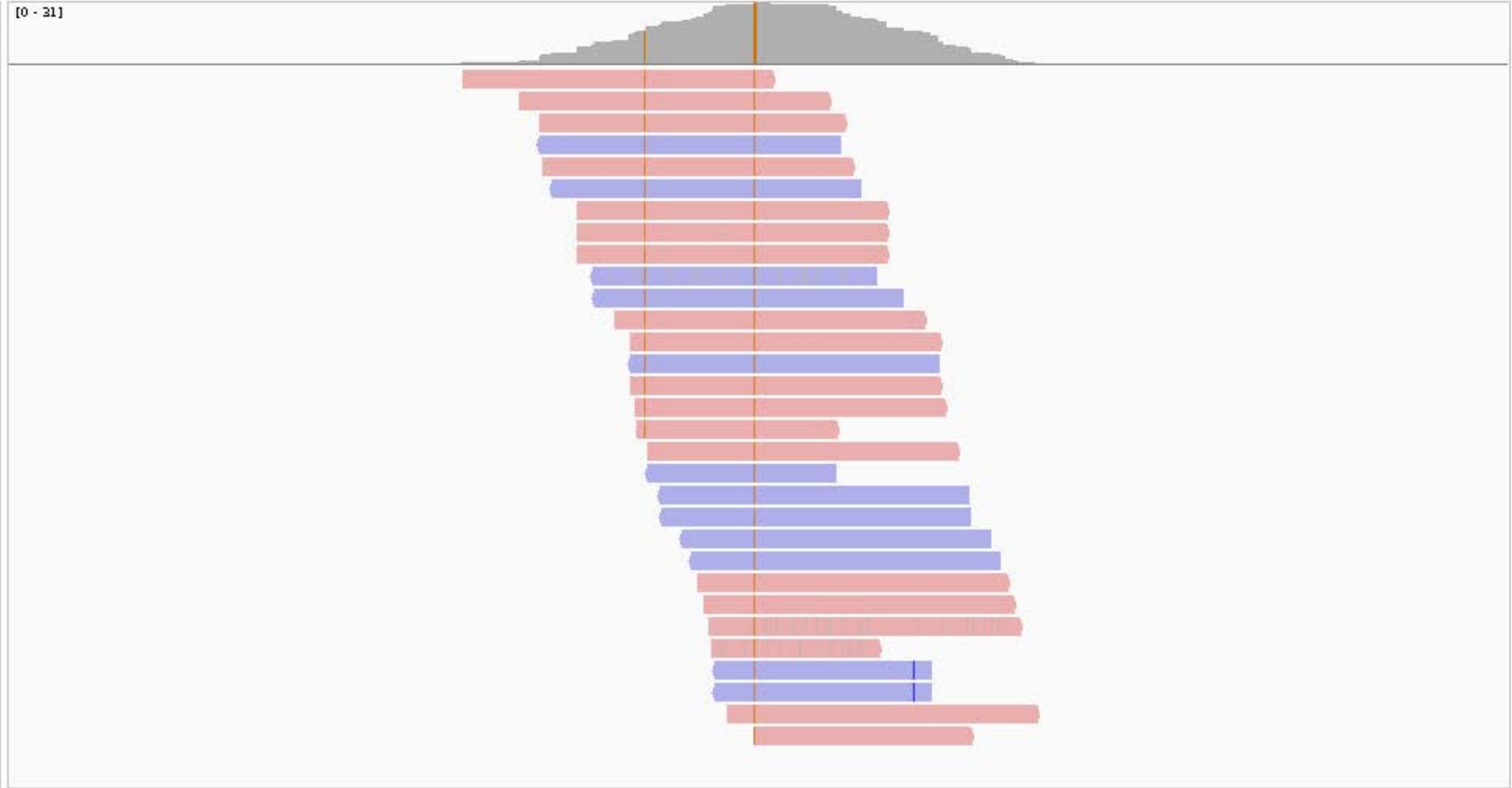
Sequence →

Refseq Genes





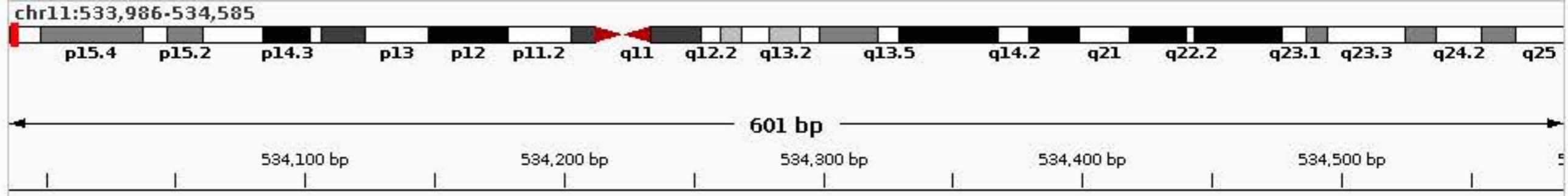
SL_1015_L_14_FFPE-B_mut.barr
rage



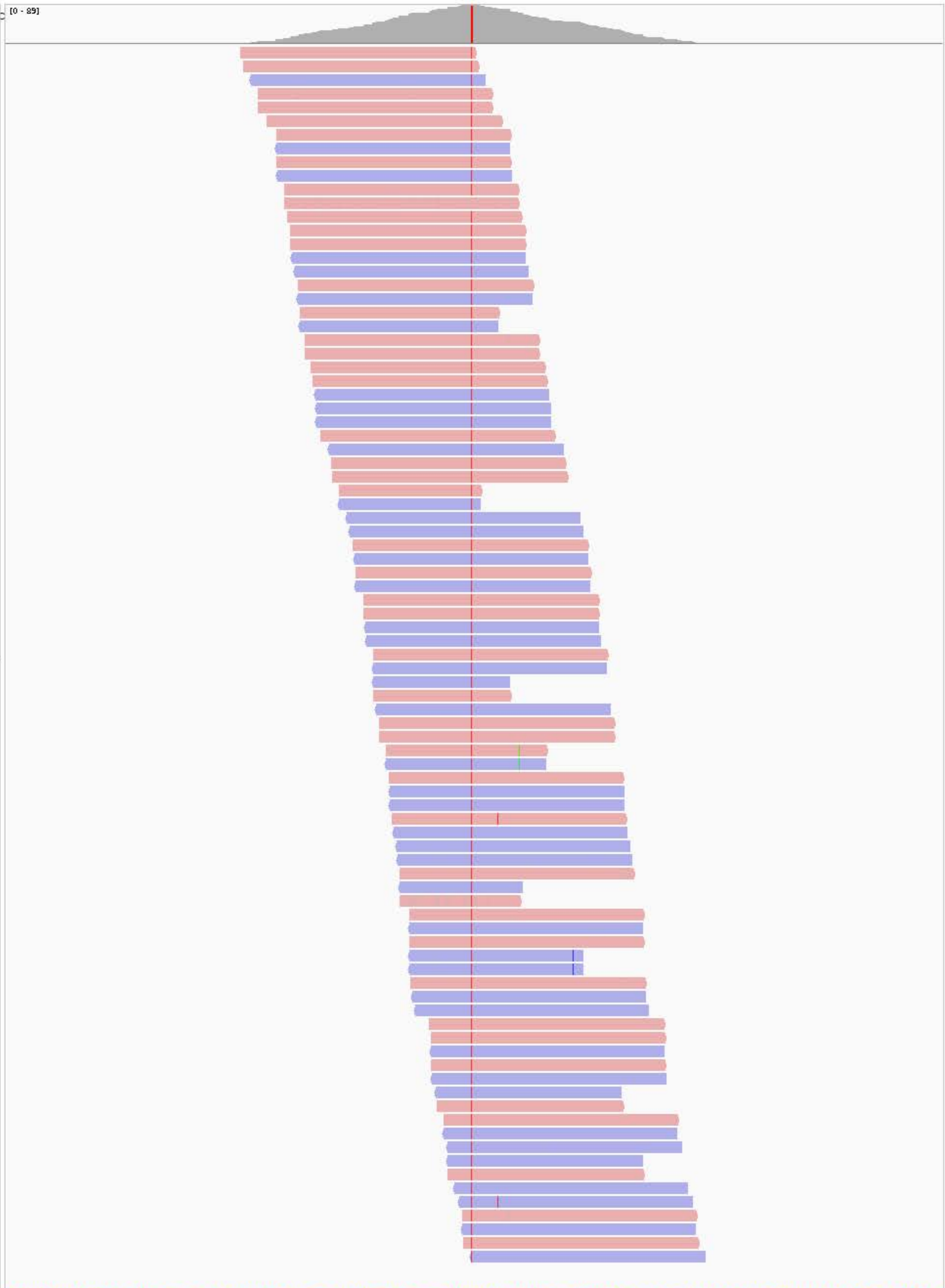
SL_1015_L_14_FFPE-B_mut.ba
m

Sequence →
Refseq Genes





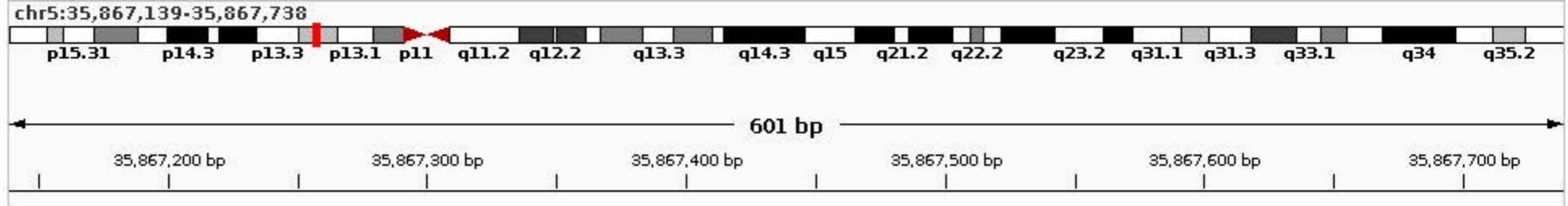
SL_0011_L_FFPE_9-B_mut.bam
age



SL_0011_L_FFPE_9-B_mut.bam

Sequence →
Refseq Genes





SL_1002_L_FFPE_24-B_mut.bam
rage



SL_1002_L_FFPE_24-B_mut.ba
m

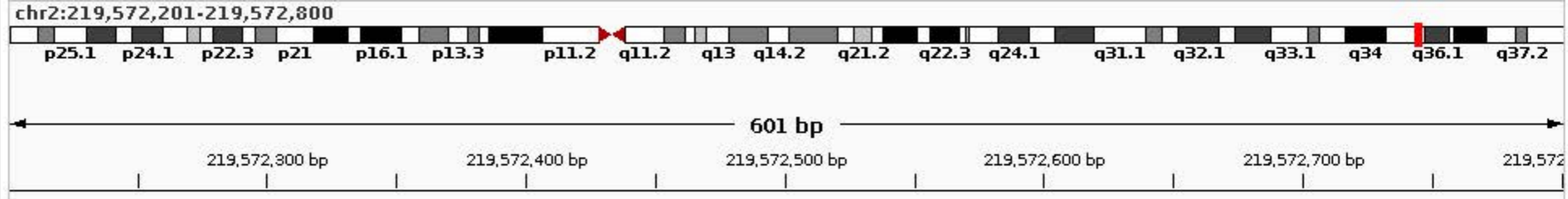
Sequence →



Refseq Genes

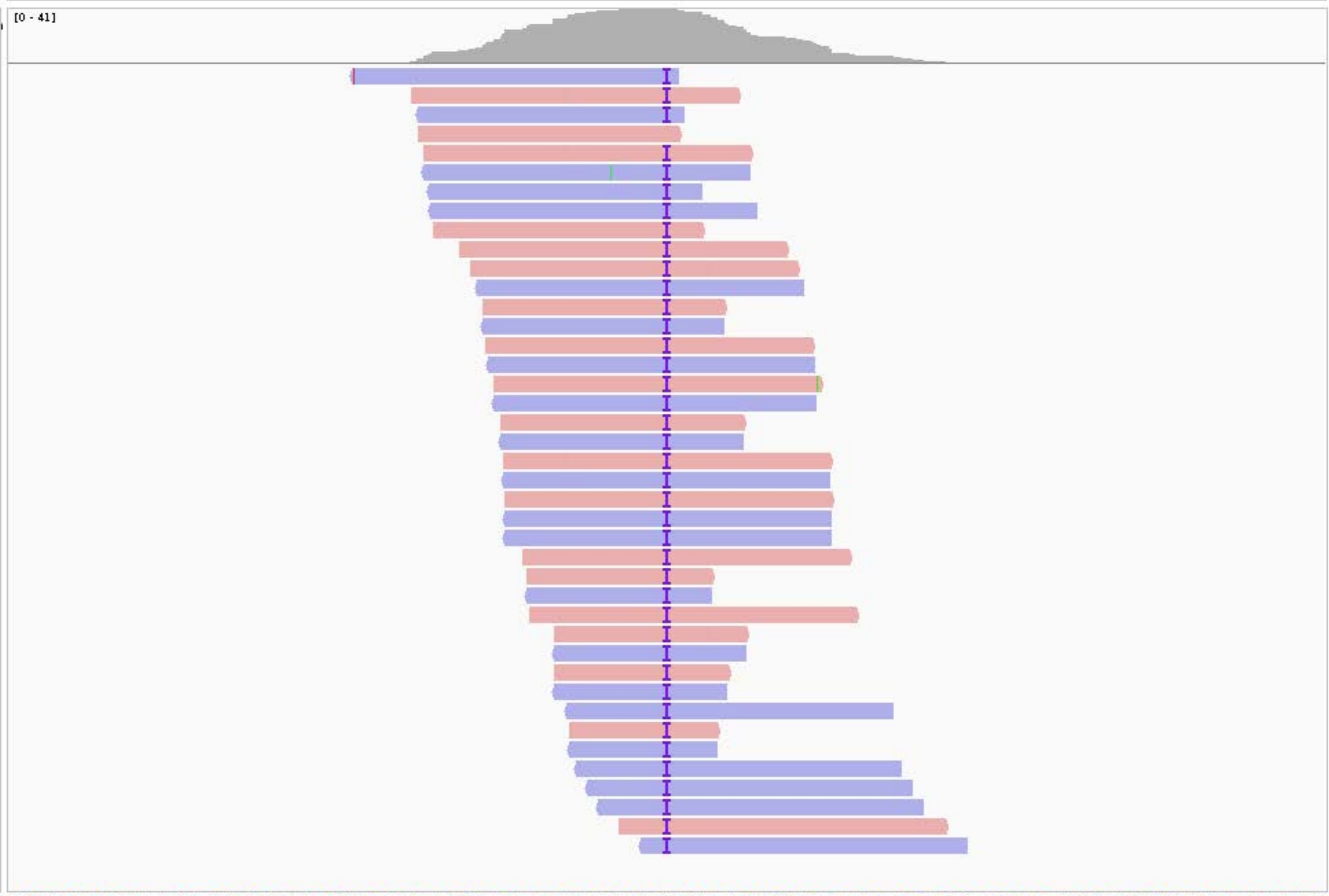


IL7R



SL_1002_R_FFPE_33-B_mut.bam
rage

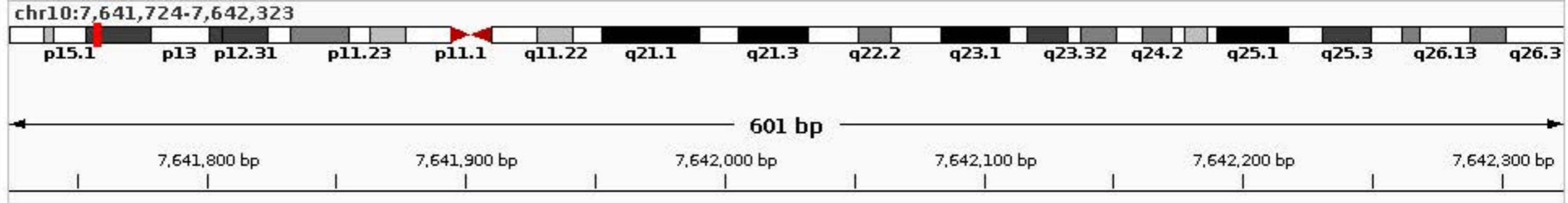
SL_1002_R_FFPE_33-B_mut.ba
m



Sequence →

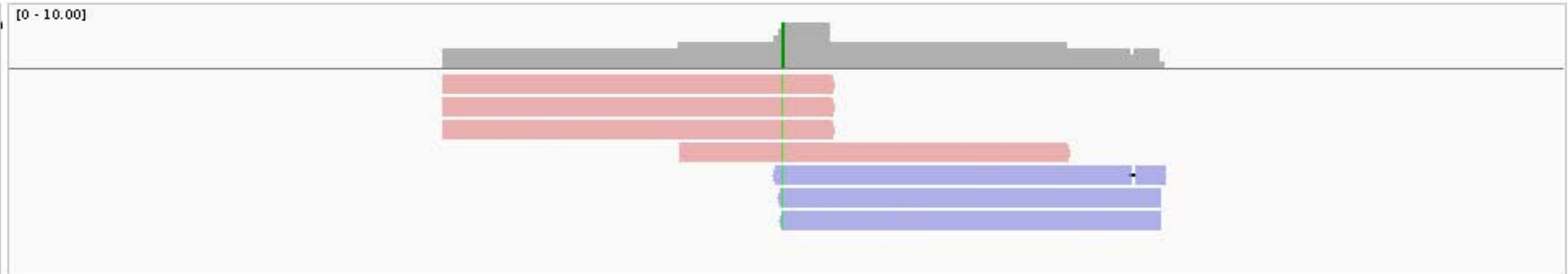
Refseq Genes





SL_0001_R_FFPE_55-B_mut.bam
rage

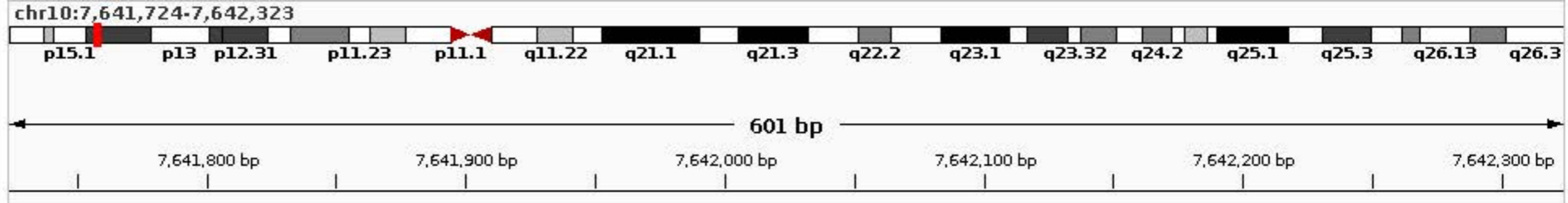
SL_0001_R_FFPE_55-B_mut.ba
m



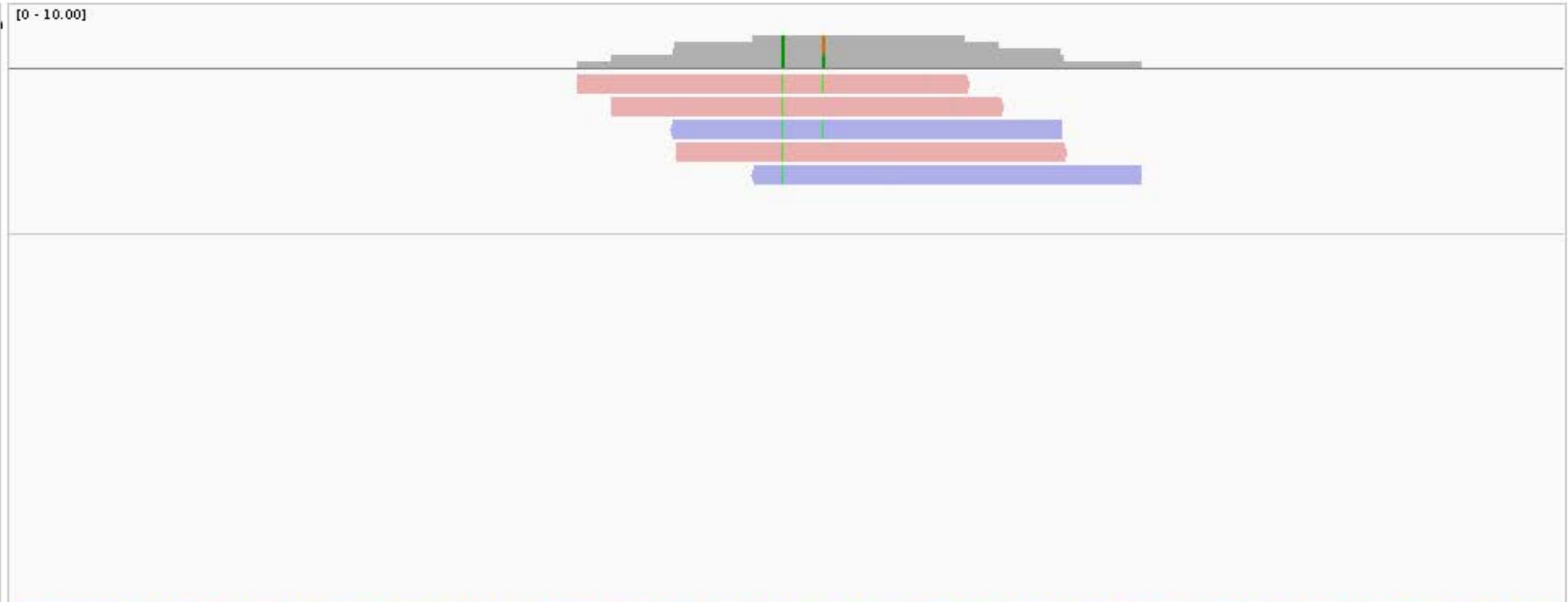
Sequence →

Refseq Genes





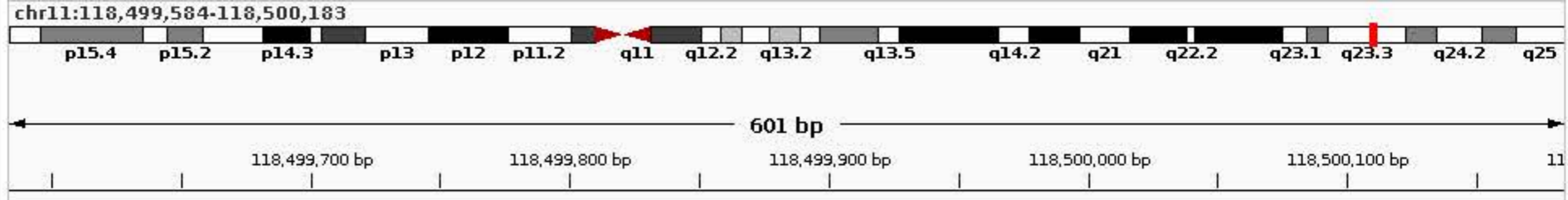
SL_0001_R_FFPE_56-B_mut.bam
 SL_0001_R_FFPE_56-B_mut.bam



Sequence →

Refseq Genes



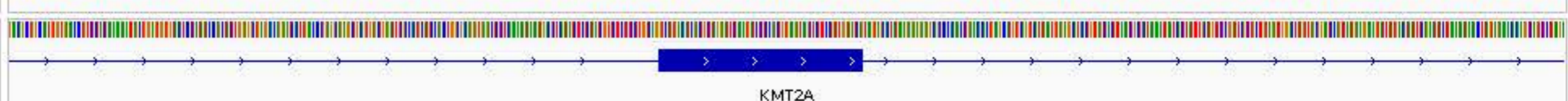
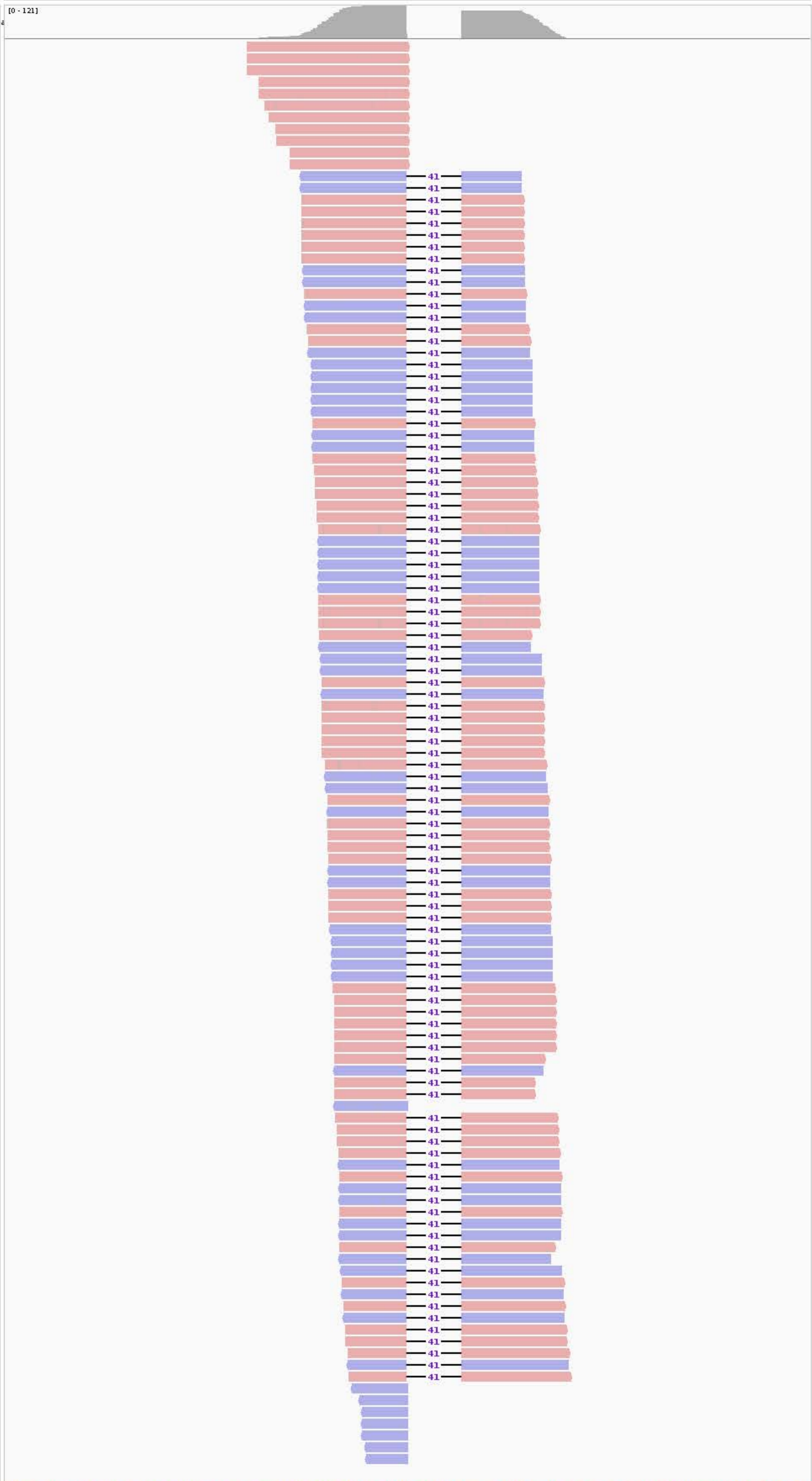


SL_0009-LT_3-B_mut.bam Coverage

SL_0009-LT_3-B_mut.bam

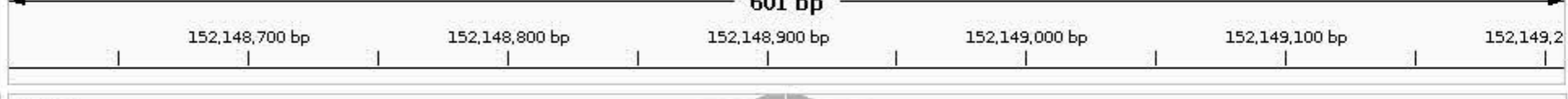
Sequence

Refseq Genes



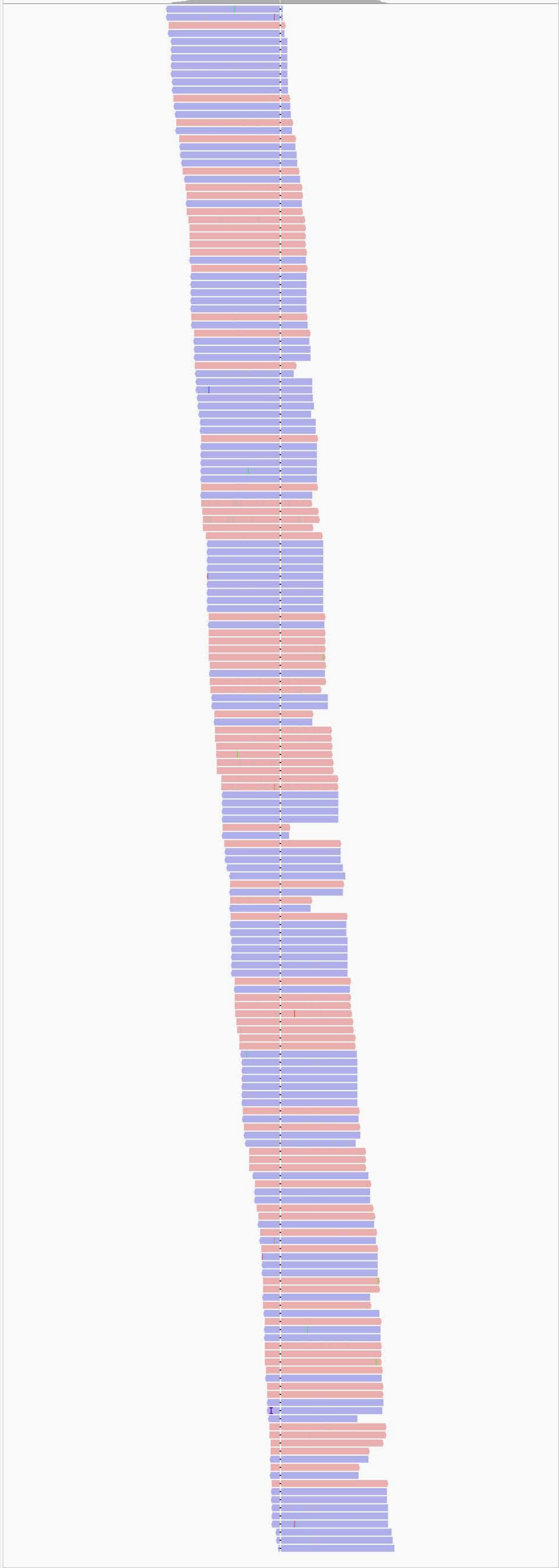
chr7:152,148,608-152,149,207

p22.1 p21.2 p15.3 p14.3 p14.1 p12.2 p11.1 q11.22 q21.11 q21.2 q22.1 q31.1 q31.31 q32.1 q33 q35 q36.2



[0 - 191]

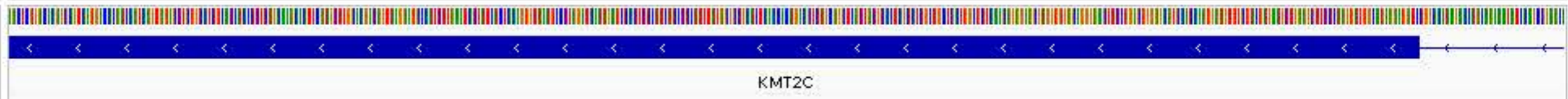
SL_0013_RT_4-B_mut.bam Cover



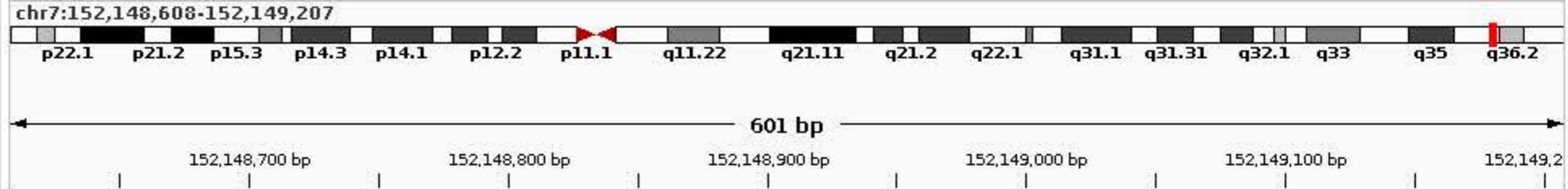
SL_0013_RT_4-B_mut.bam

Sequence

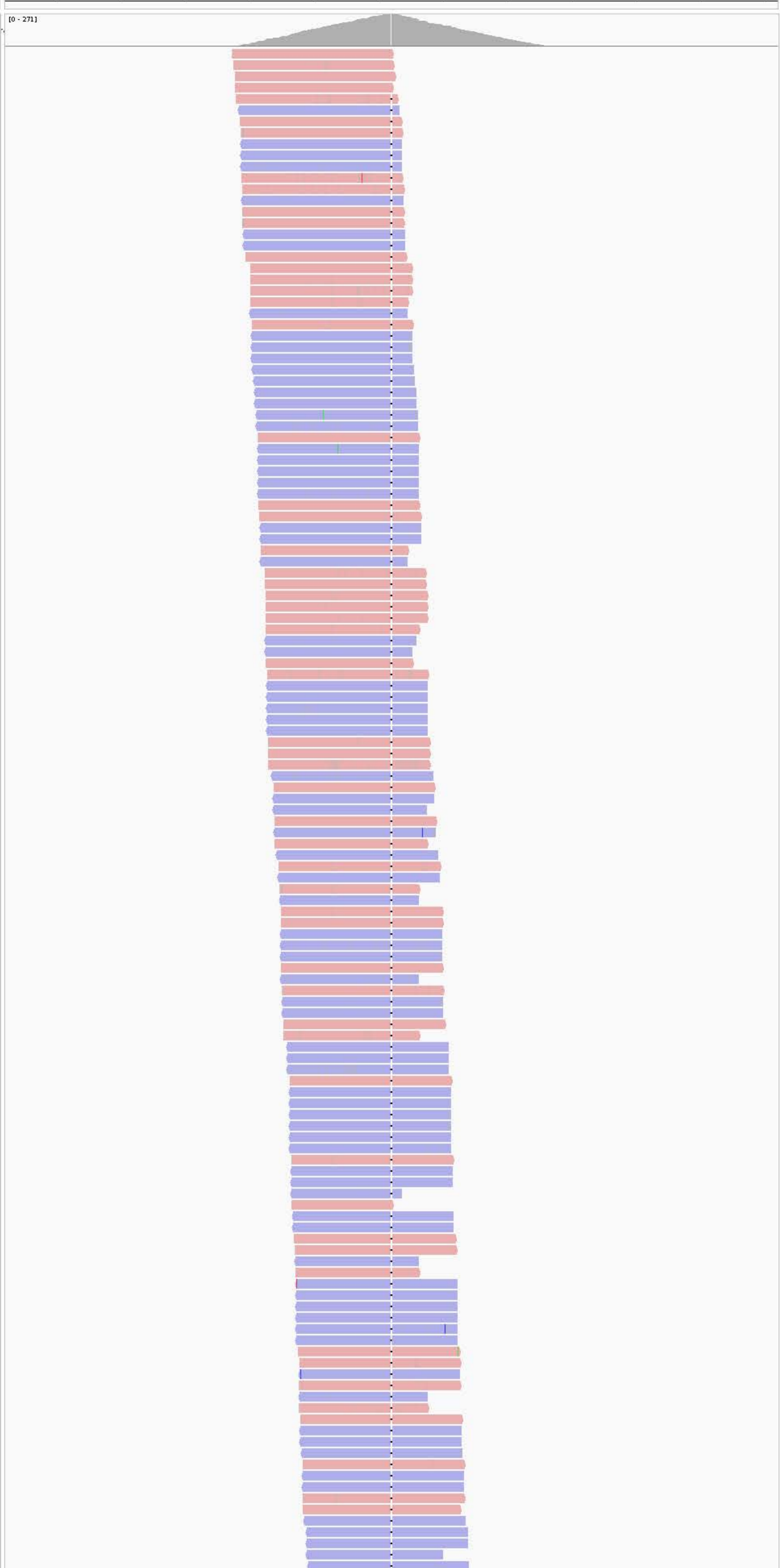
Refseq Genes



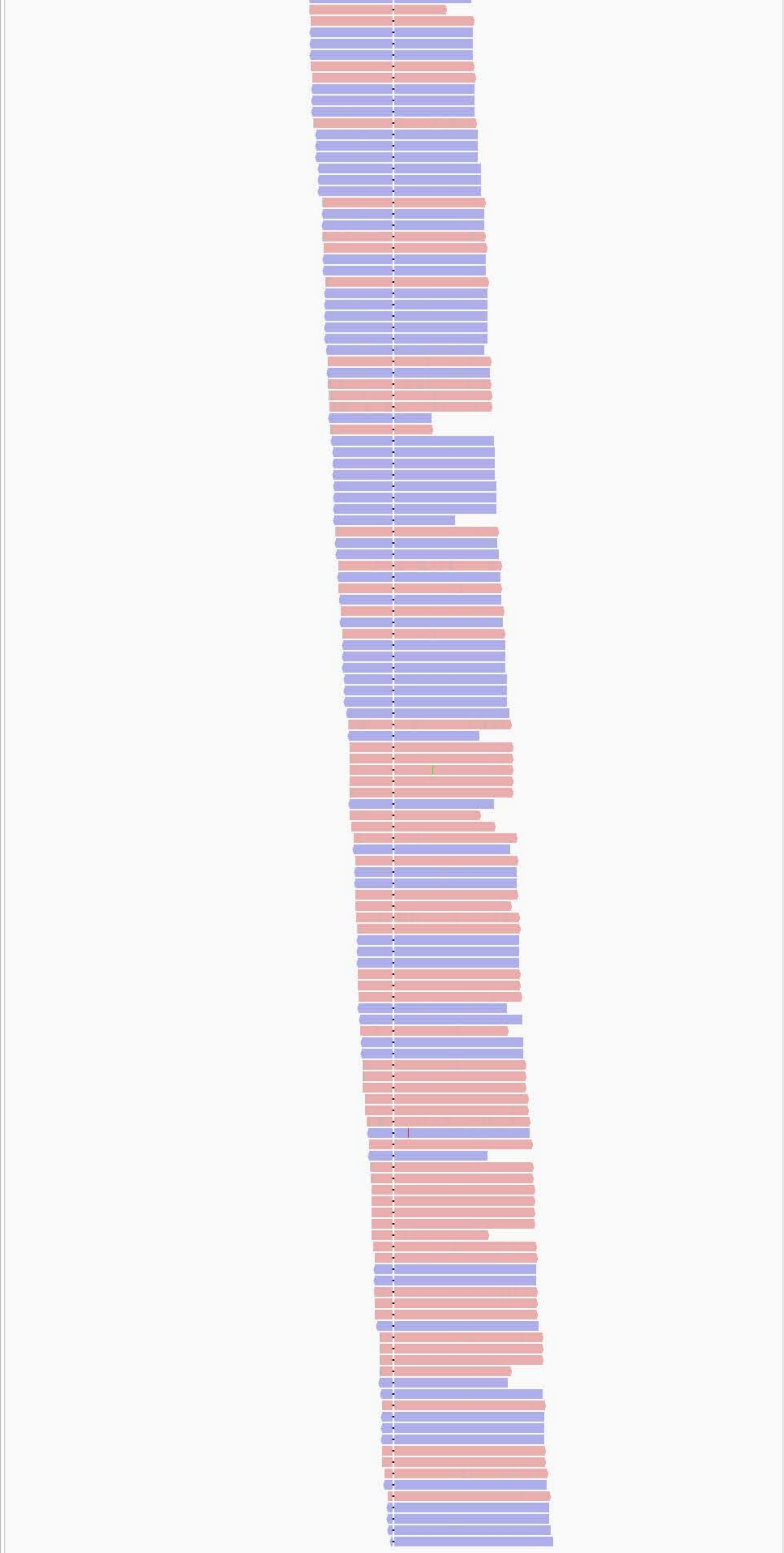
KMT2C



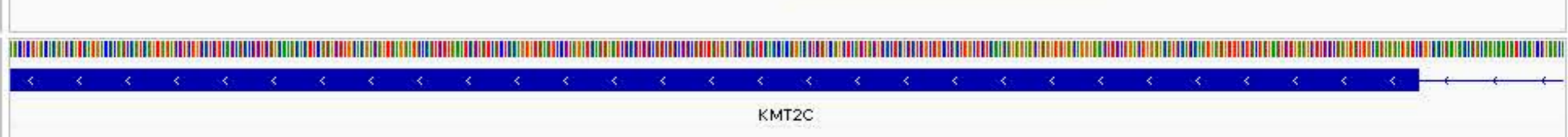
SL_0013_RT_5-B_mut.bam Cover



SL_0013_RT_5-B_mut.bam

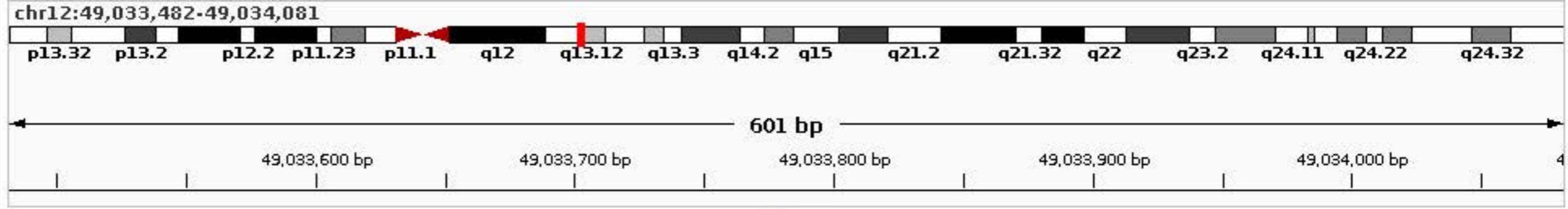


Sequence



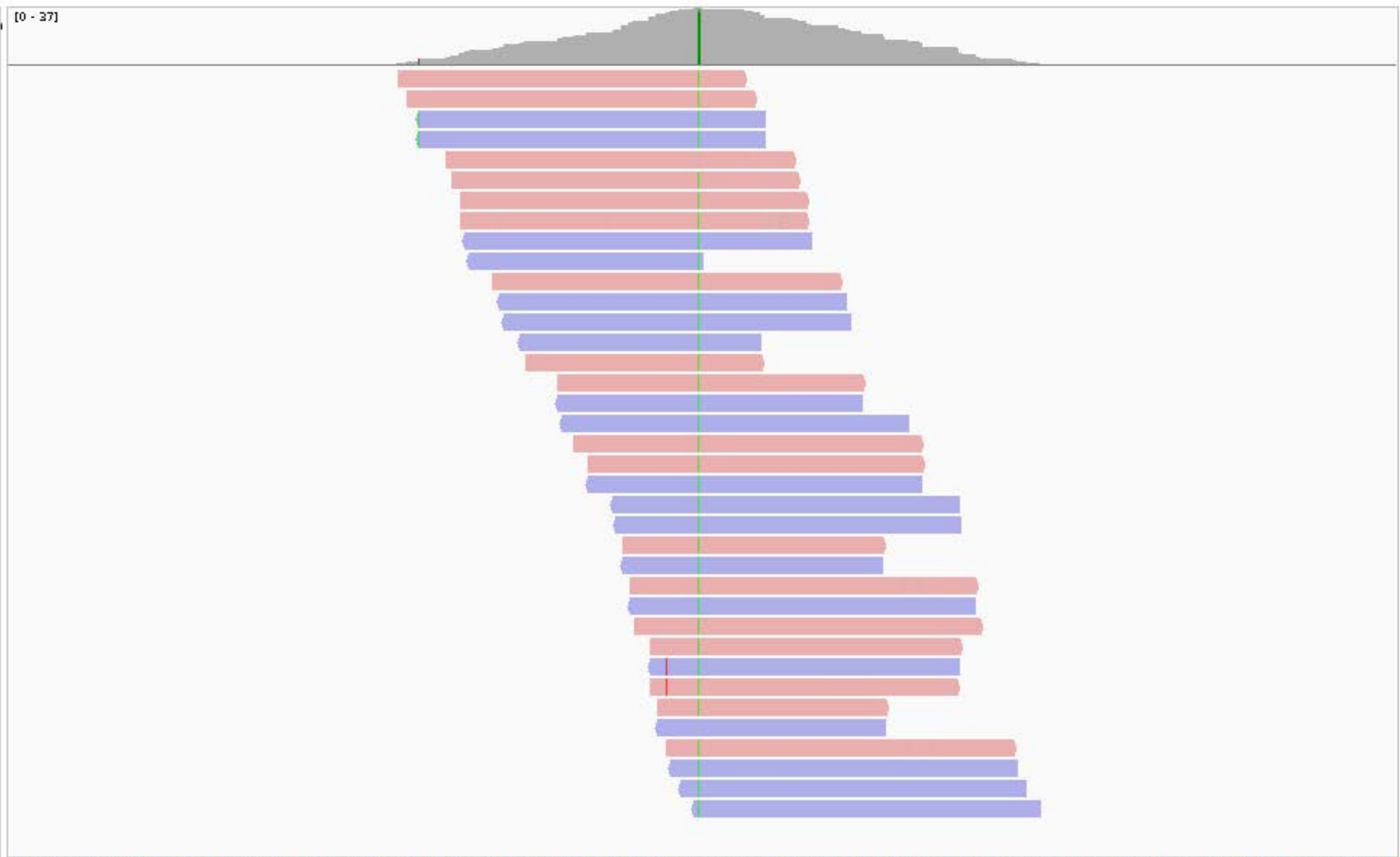
Refseq Genes





SL_0007_R_FFPE_58-B_mut.bam
rage

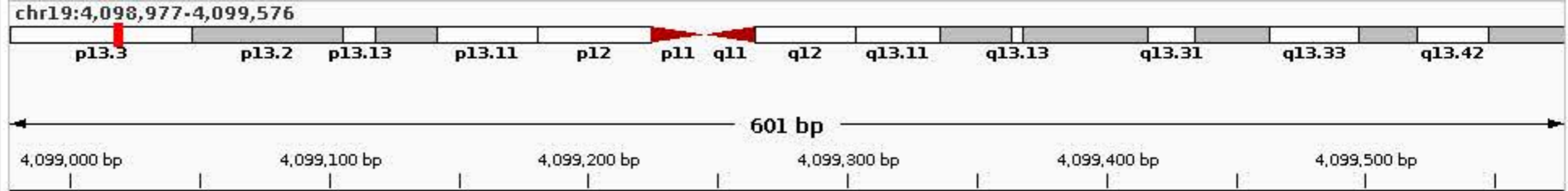
SL_0007_R_FFPE_58-B_mut.ba
m



Sequence →

Refseq Genes



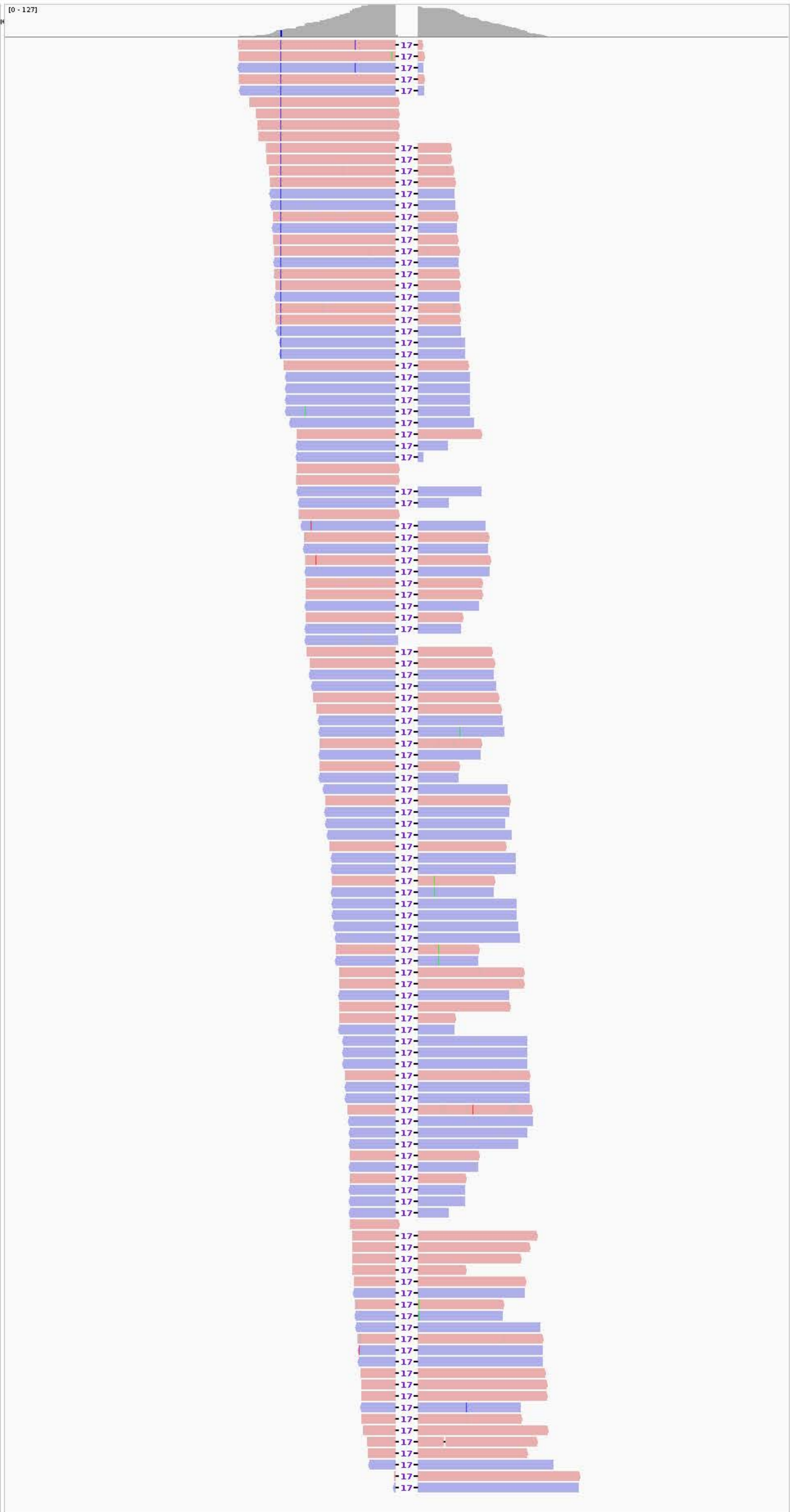


SL_1007-T1-B_mut.bam Coverage

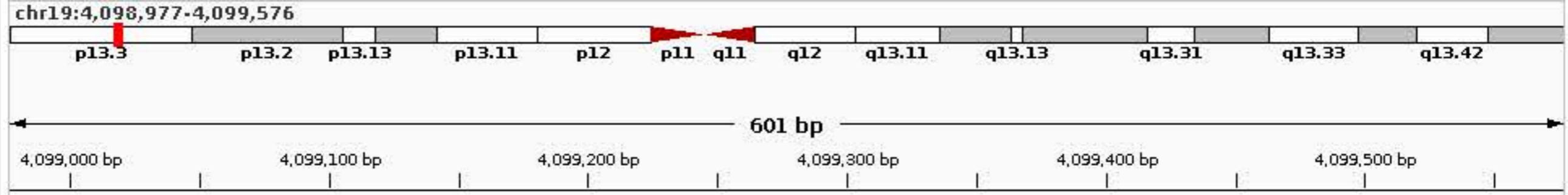
SL_1007-T1-B_mut.bam

Sequence →

Refseq Genes

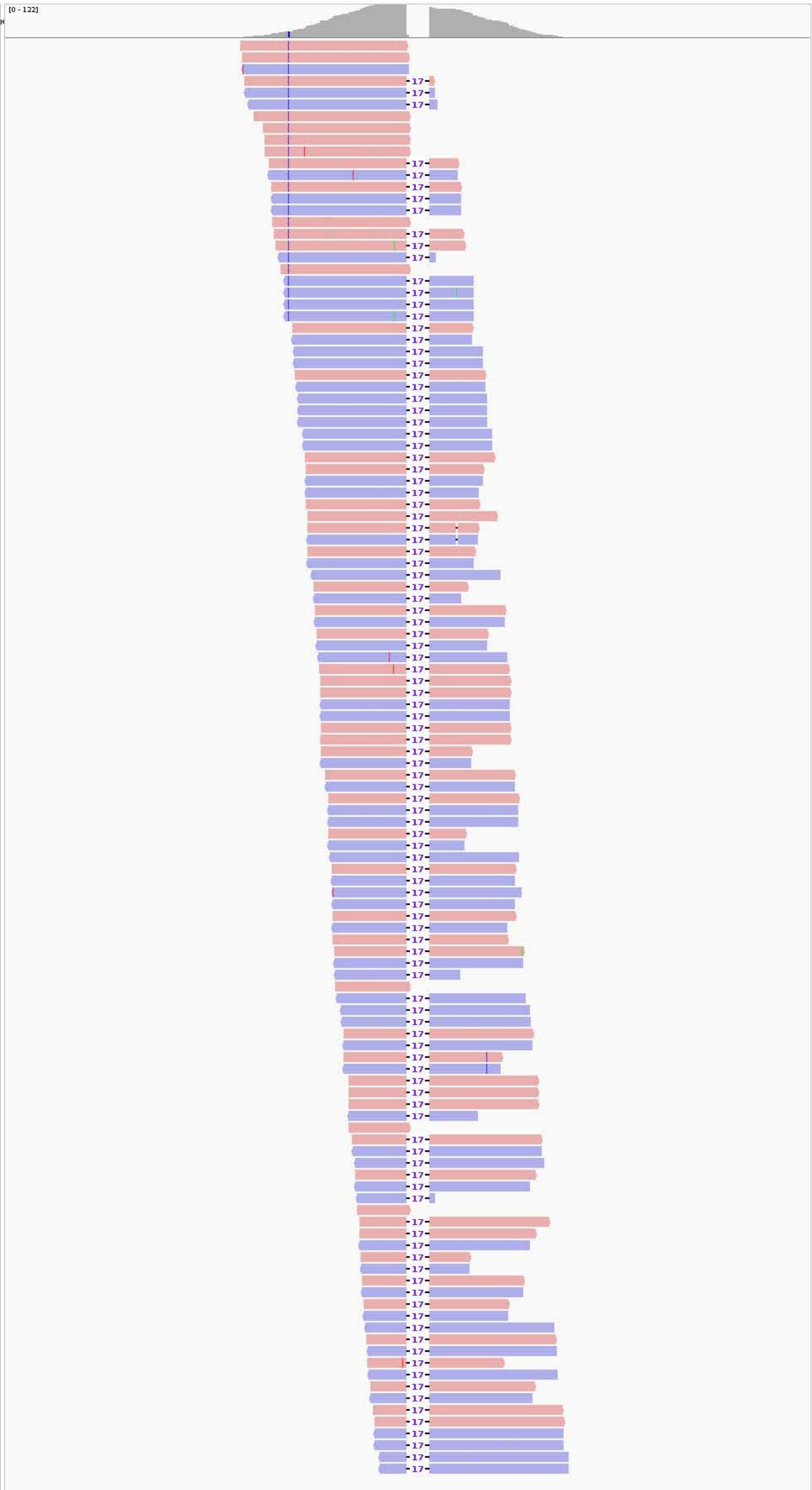


MAP2K2



SL_1007-T2-B_mut.bam Coverage

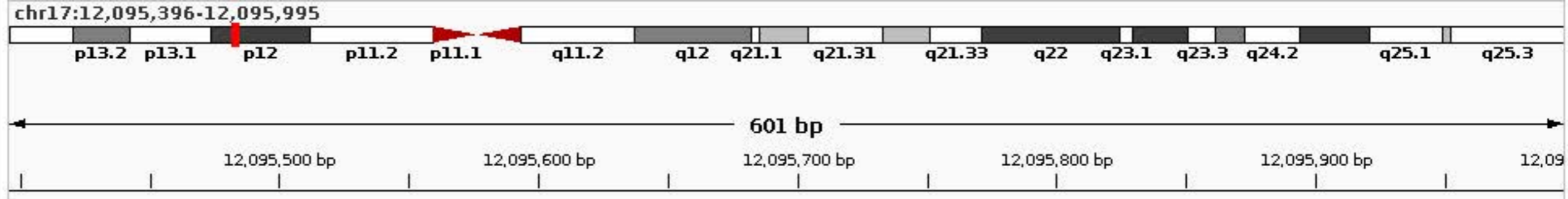
SL_1007-T2-B_mut.bam



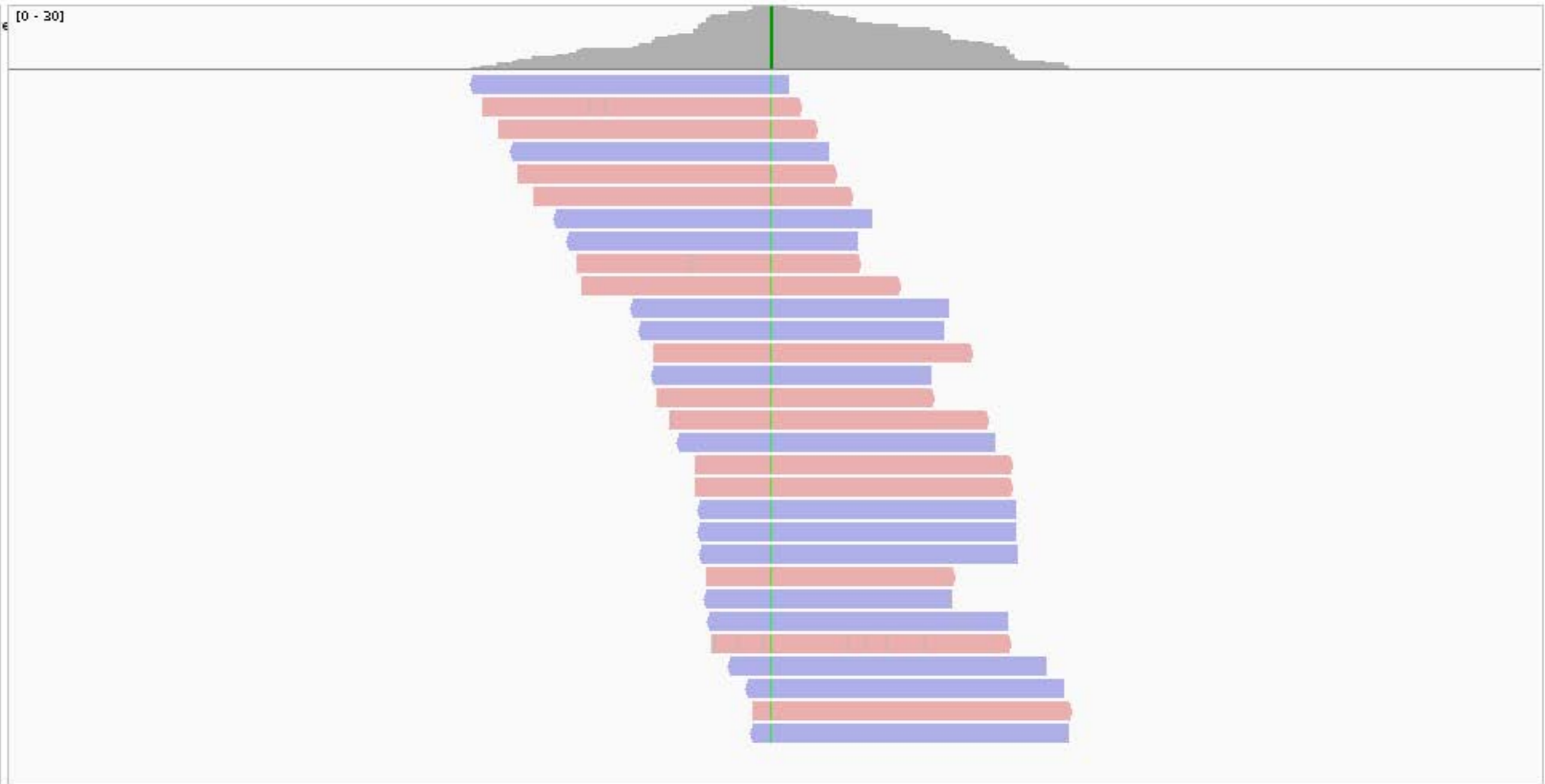
Sequence

Refseq Genes





SL_0005-RN_11-B_mut.bam Coverage



SL_0005-RN_11-B_mut.bam

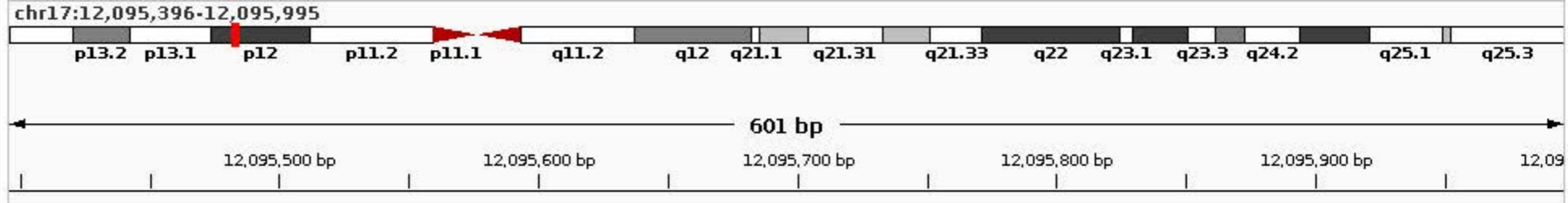
Sequence



Refseq Genes

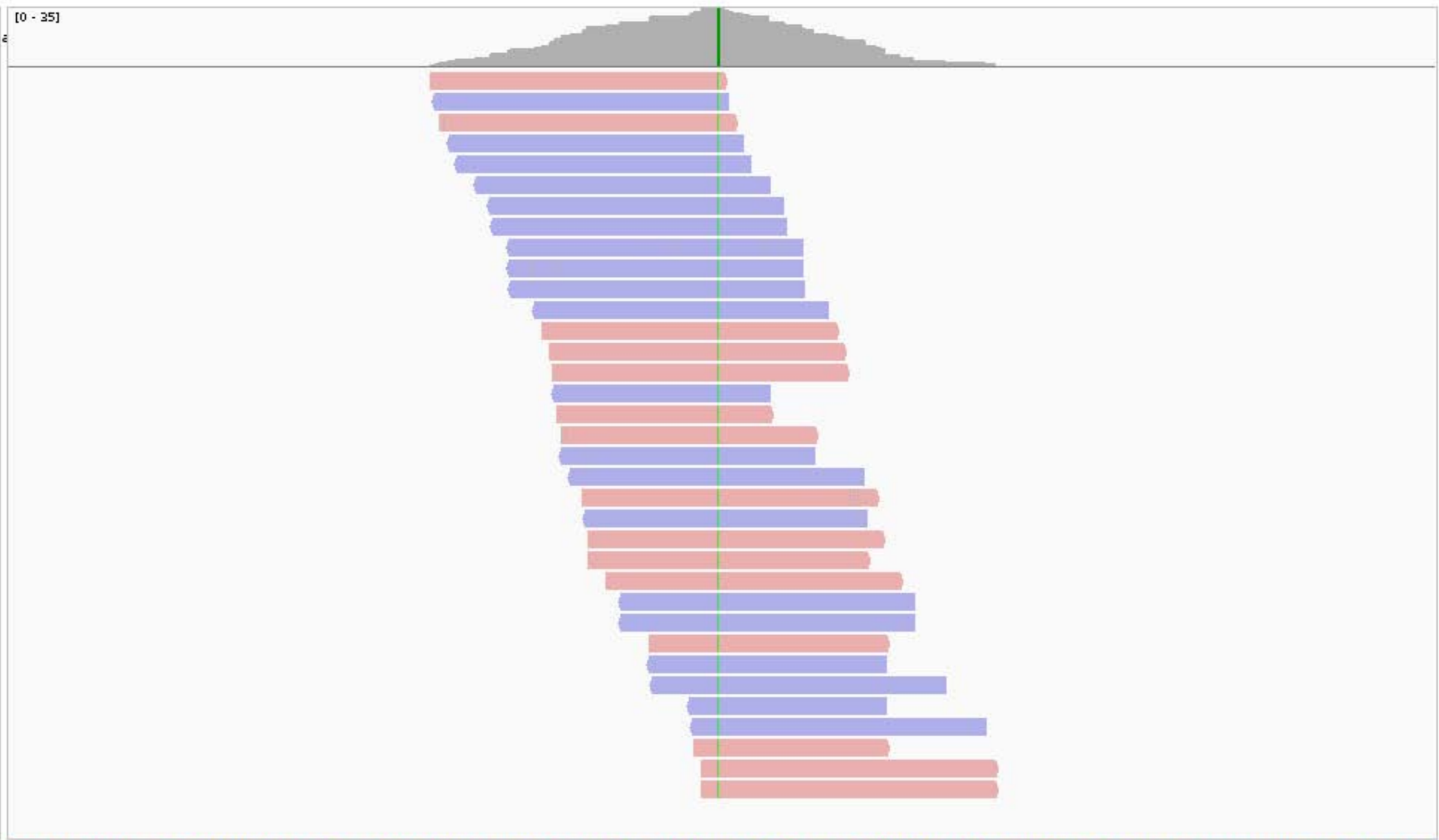


MAP2K4



SL_0009-LT_1-B_mut.bam Coverage

SL_0009-LT_1-B_mut.bam



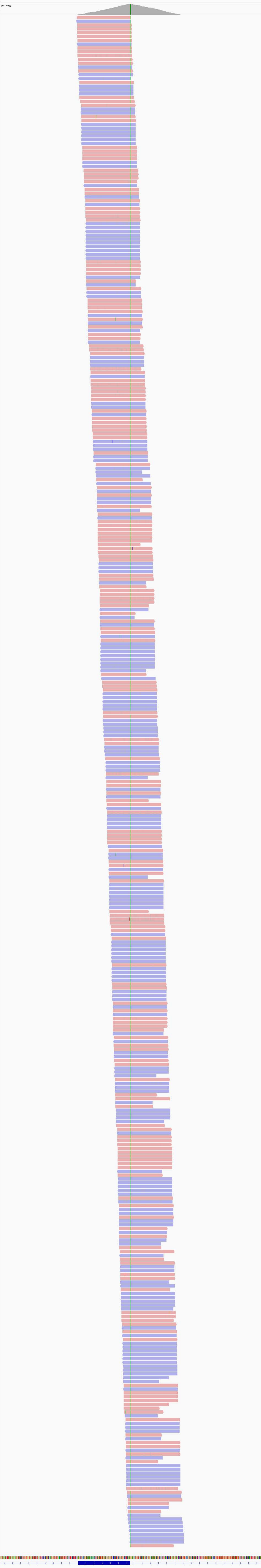
Sequence →

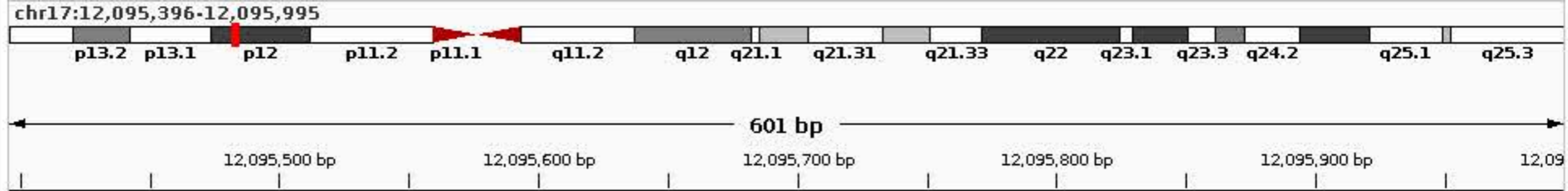
Refseq Genes



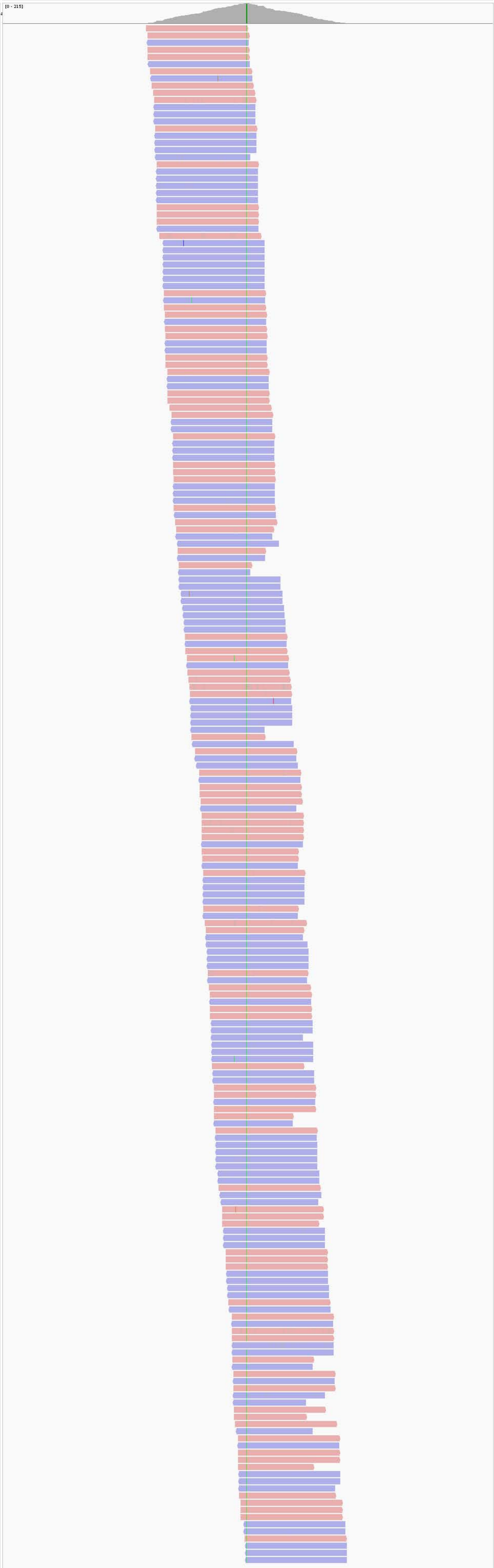
SL_0011-LT_2-B_mut.bam Coverage

SL_0011-LT_2-B_mut.bam





SL_0011-LT_5-B_mut.bam Coverage

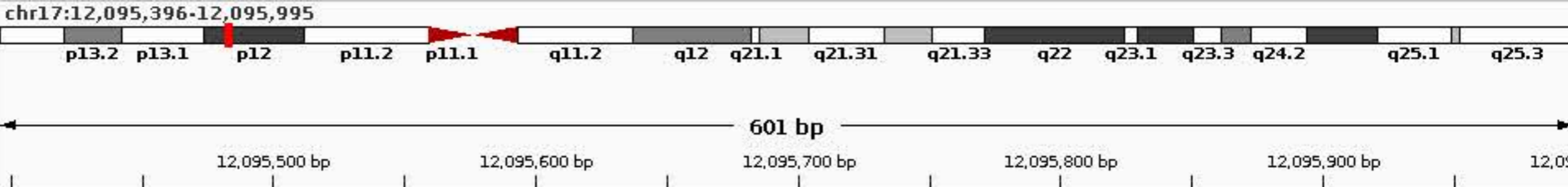


Sequence



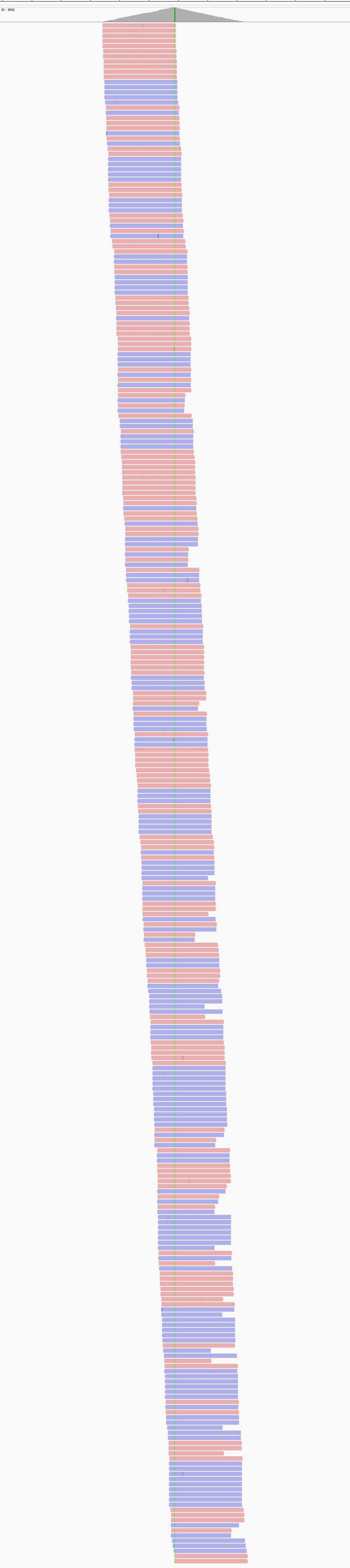
Refseq Genes





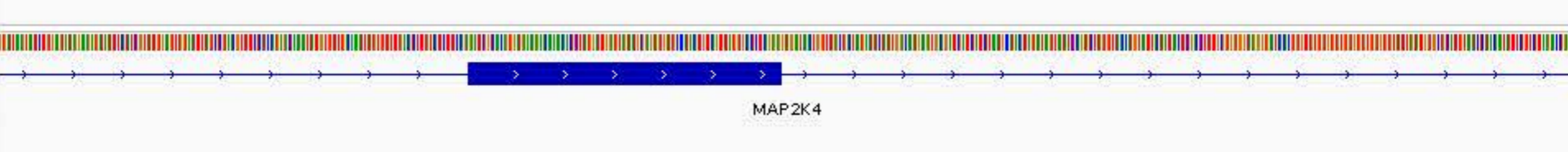
SL_0011-LT_6-B_mut.bam Coverage

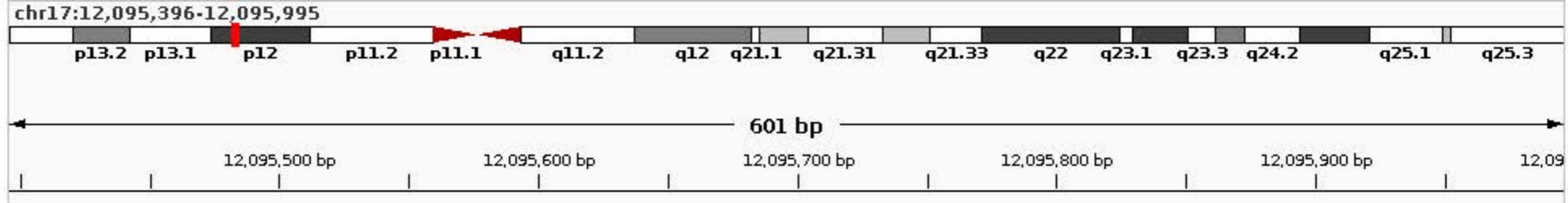
SL_0011-LT_6-B_mut.bam



Sequence

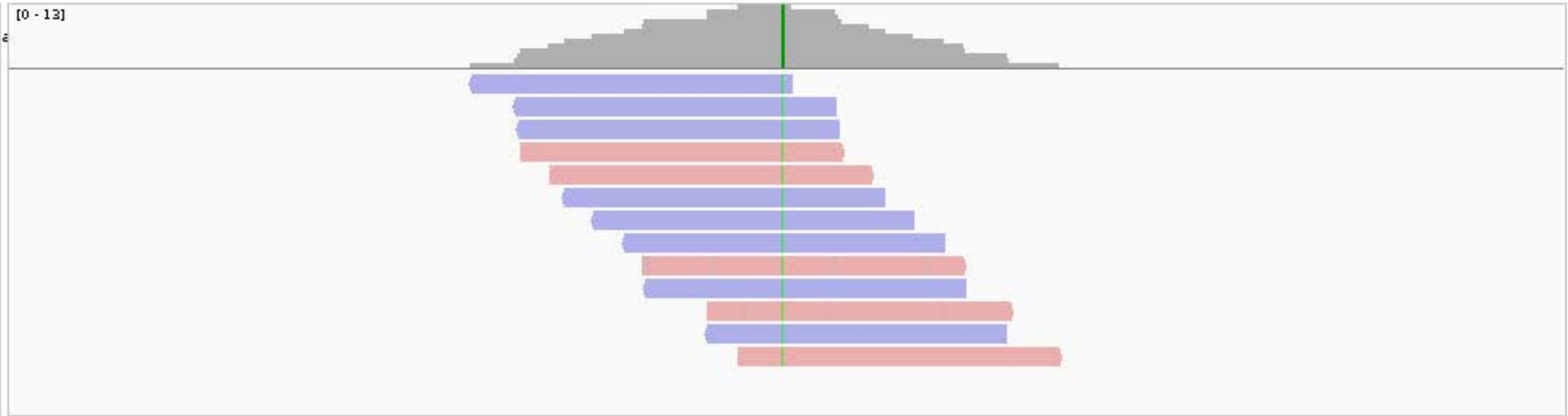
Refseq Genes





SL_0011-LT_7-B_mut.bam Coverage

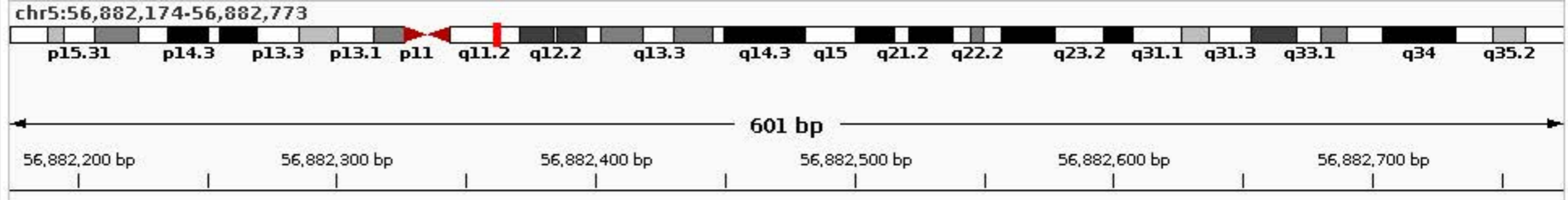
SL_0011-LT_7-B_mut.bam



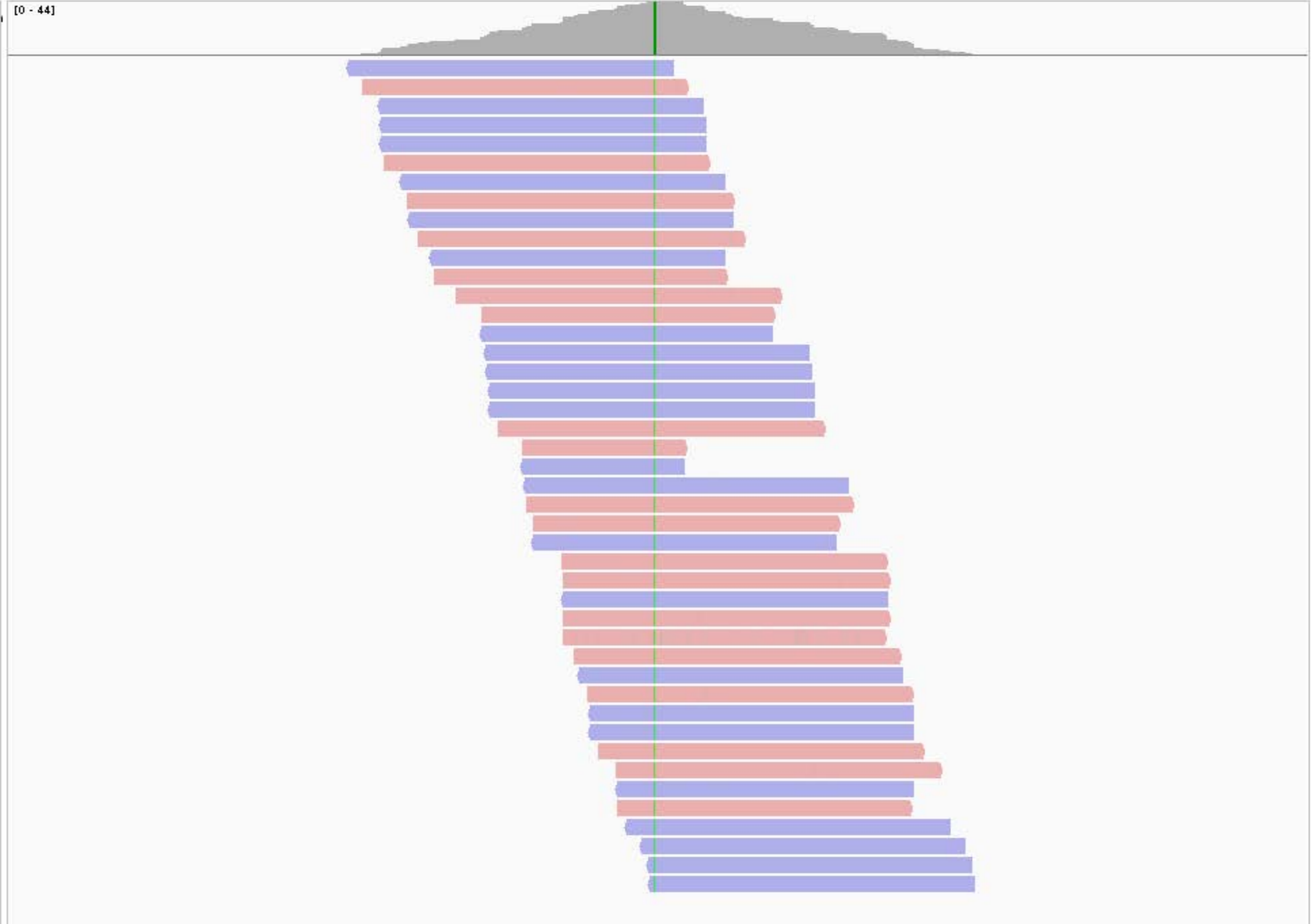
Sequence →

Refseq Genes





SL_0007_R_FFPE_58-B_mut.bam
rage



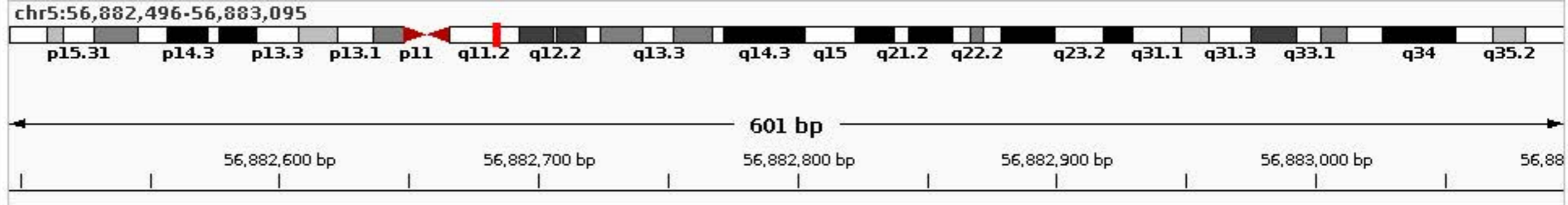
Sequence →



Refseq Genes

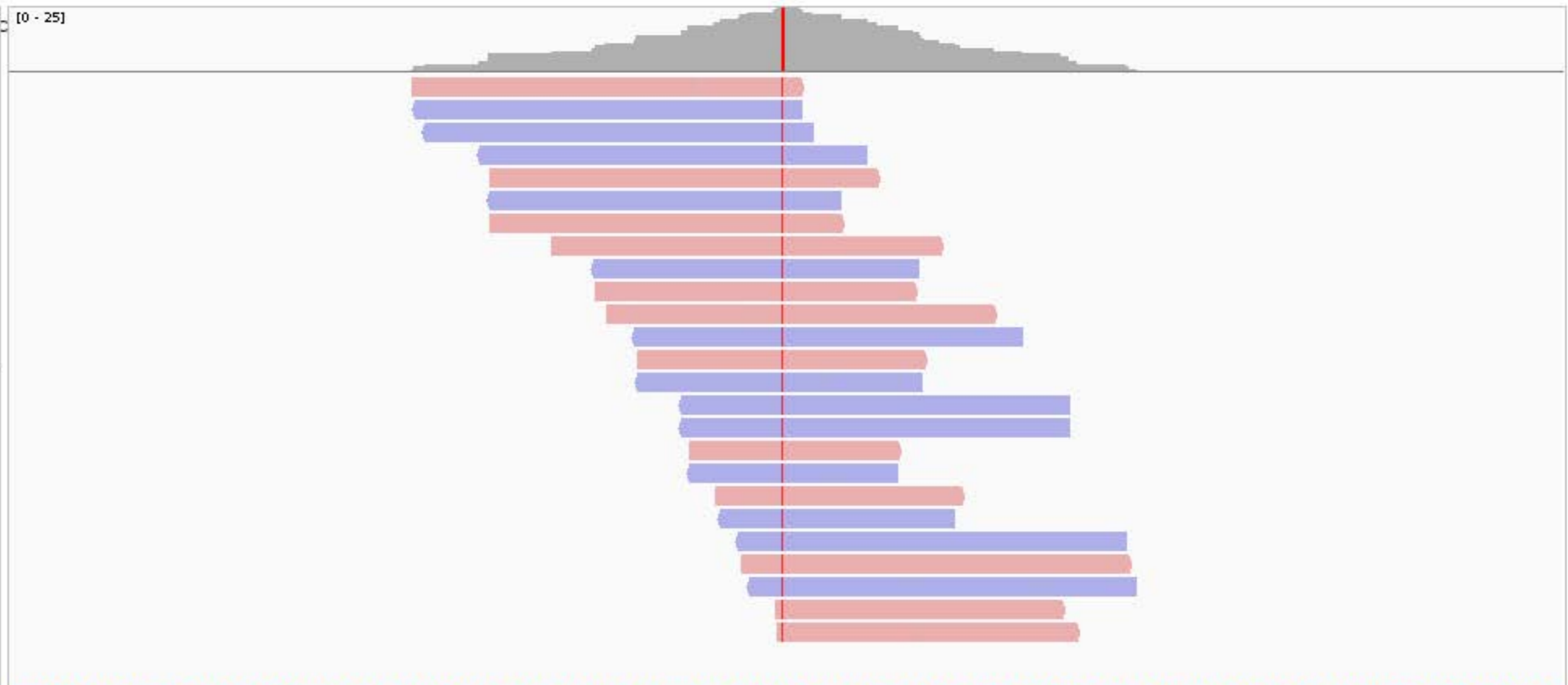


MAP3K1



SL_1010_L_FFPE_9-B_mut.bam
age

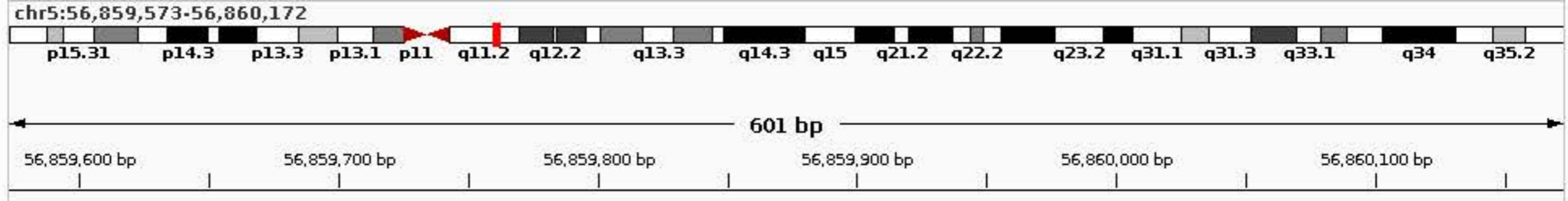
SL_1010_L_FFPE_9-B_mut.bam



Sequence →

Refseq Genes



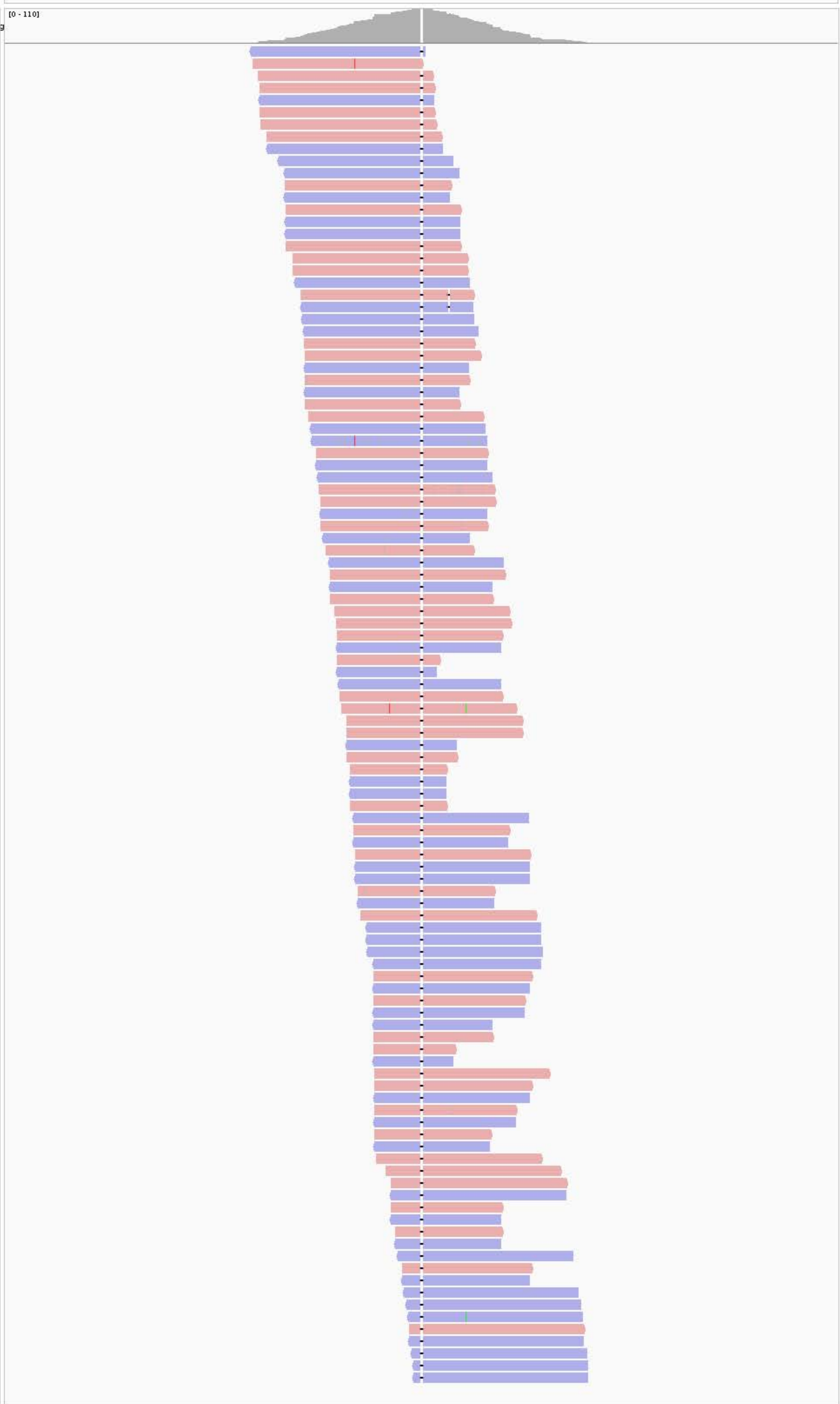


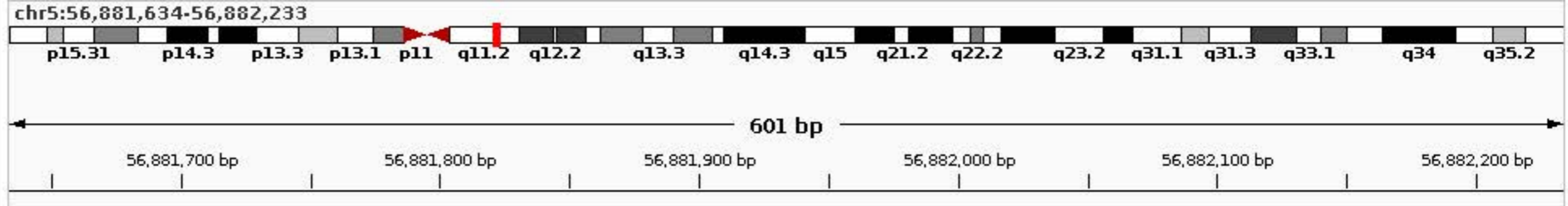
SL_1010-N5-B_mut.bam Coverage

SL_1010-N5-B_mut.bam

Sequence

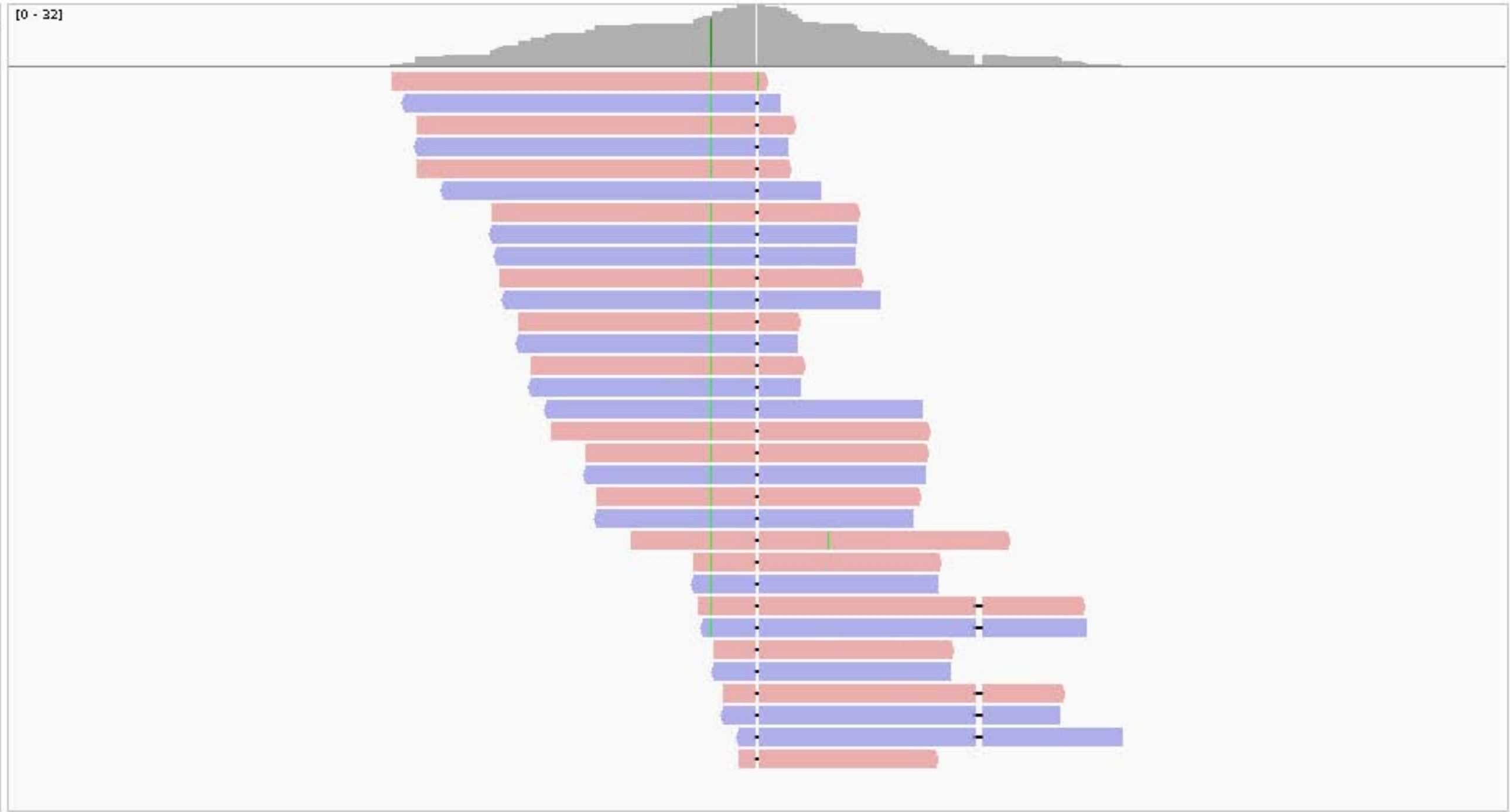
Refseq Genes





SL_1010_L_FFPE_13-B_mut.barrage

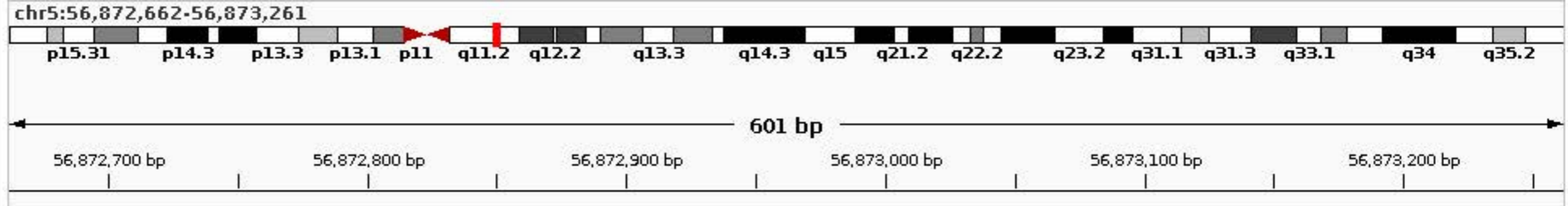
SL_1010_L_FFPE_13-B_mut.bam



Sequence

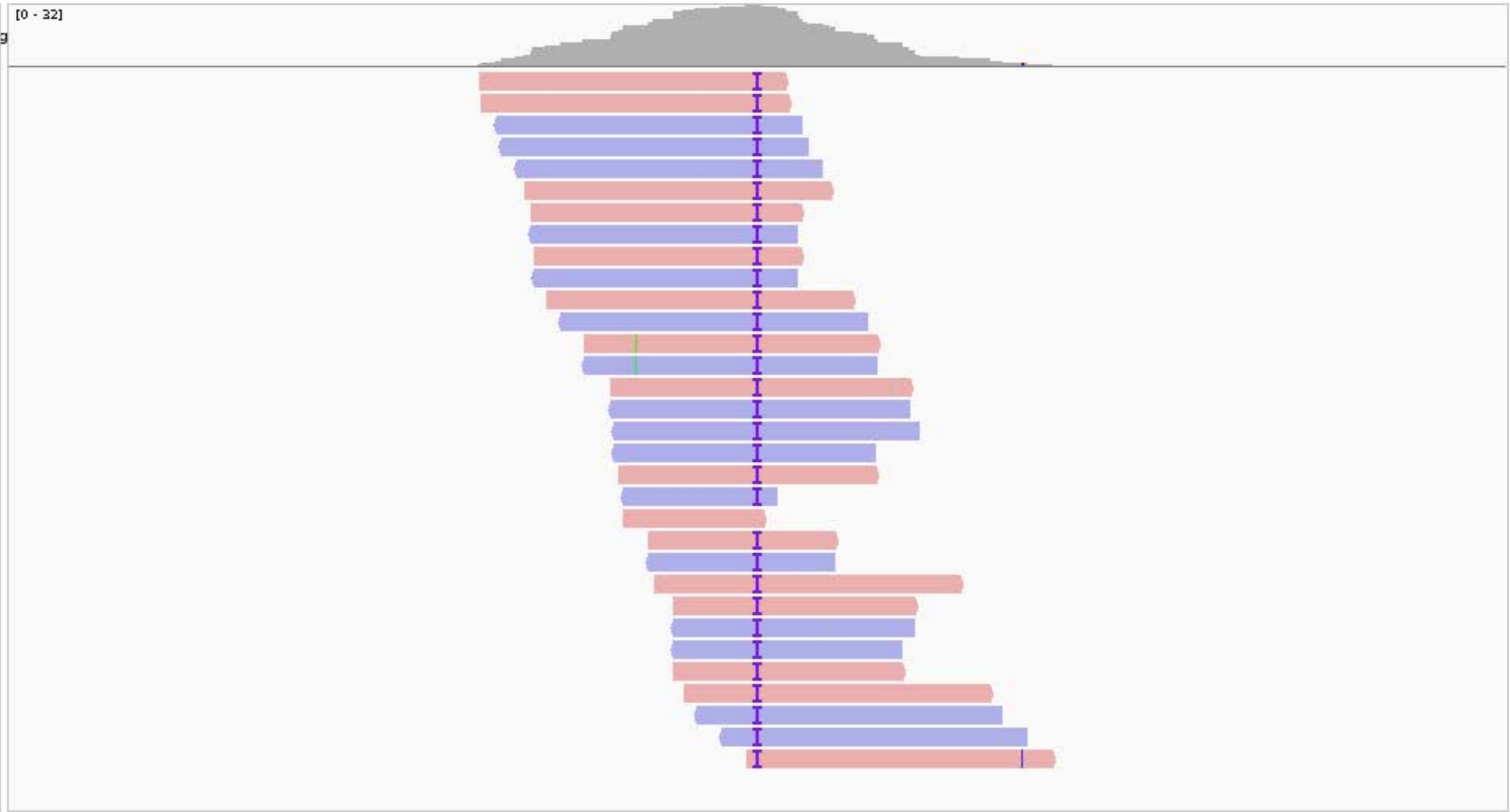
Refseq Genes





SL_1010-N5-B_mut.bam Coverage

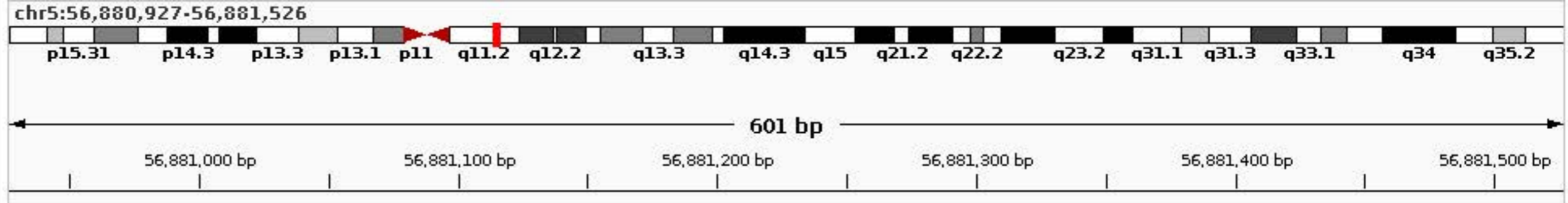
SL_1010-N5-B_mut.bam



Sequence

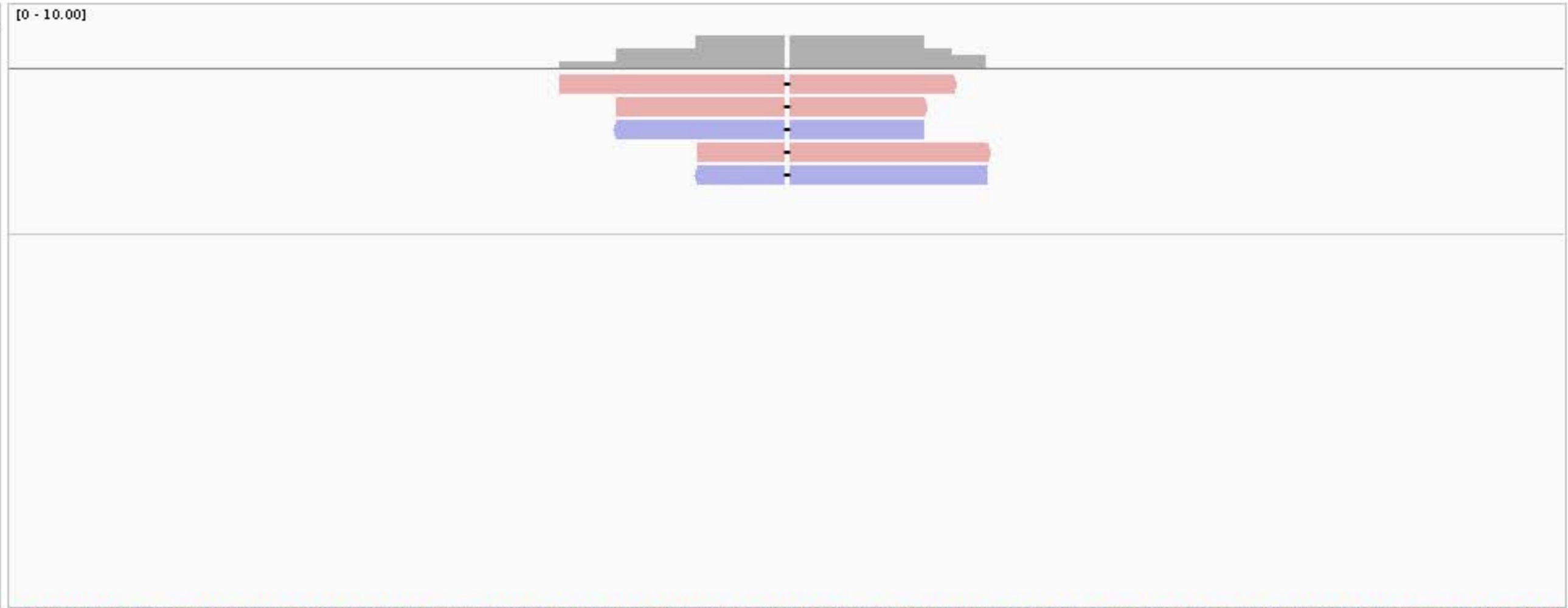
Refseq Genes





SL_1010_L_FFPE_14-B_mut.barr
rage

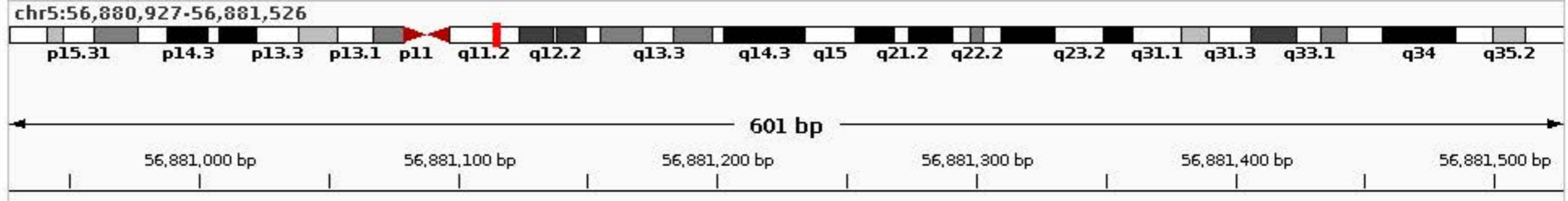
SL_1010_L_FFPE_14-B_mut.ba
m



Sequence →

Refseq Genes



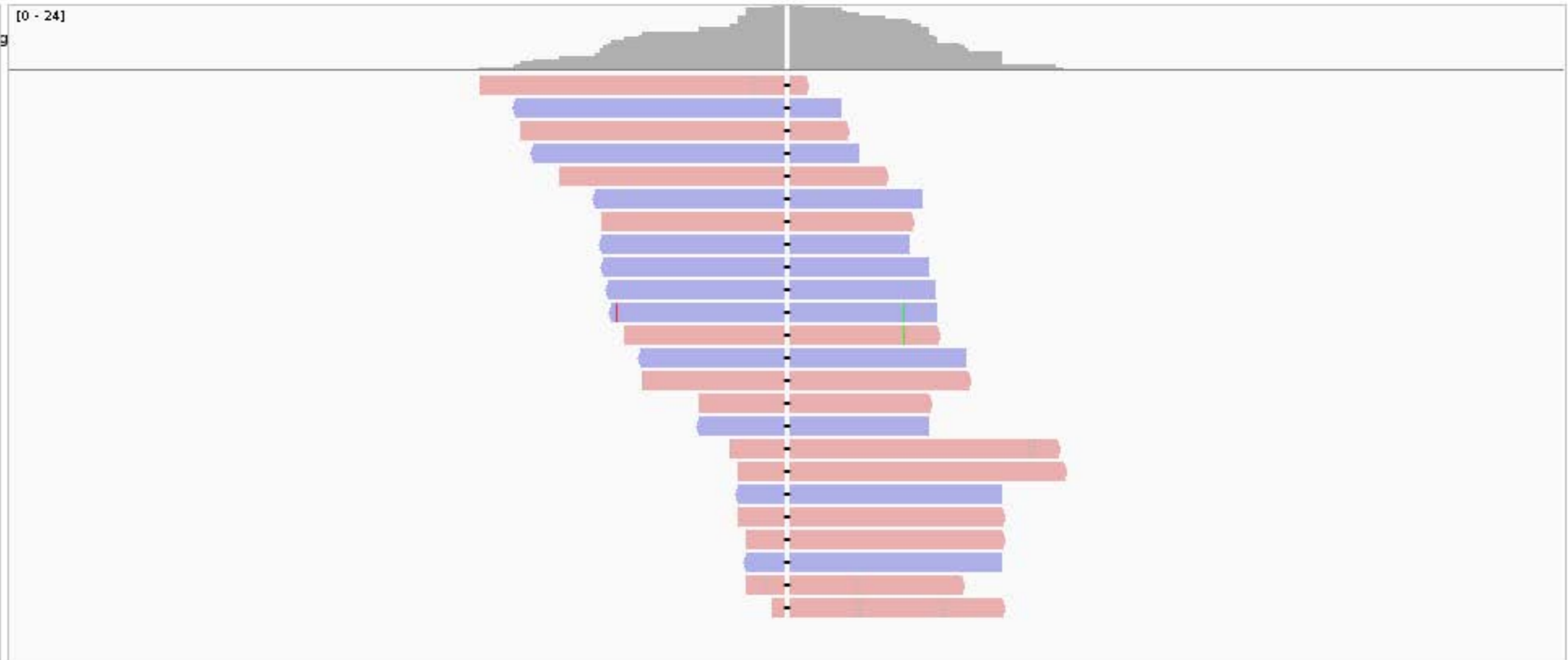


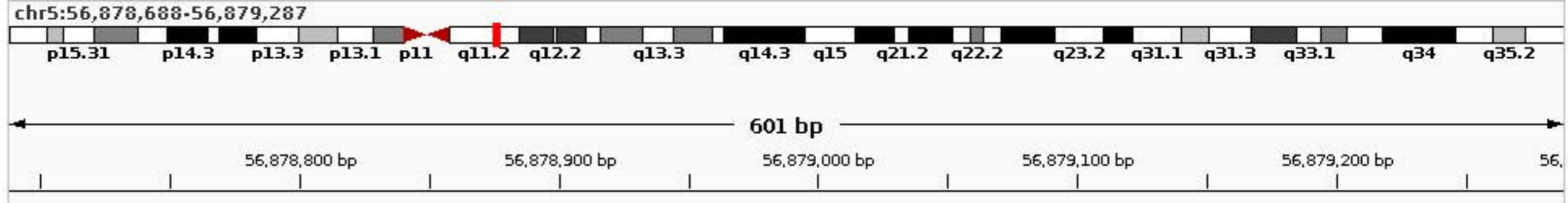
SL_1010-N4-B_mut.bam Coverage

SL_1010-N4-B_mut.bam

Sequence →

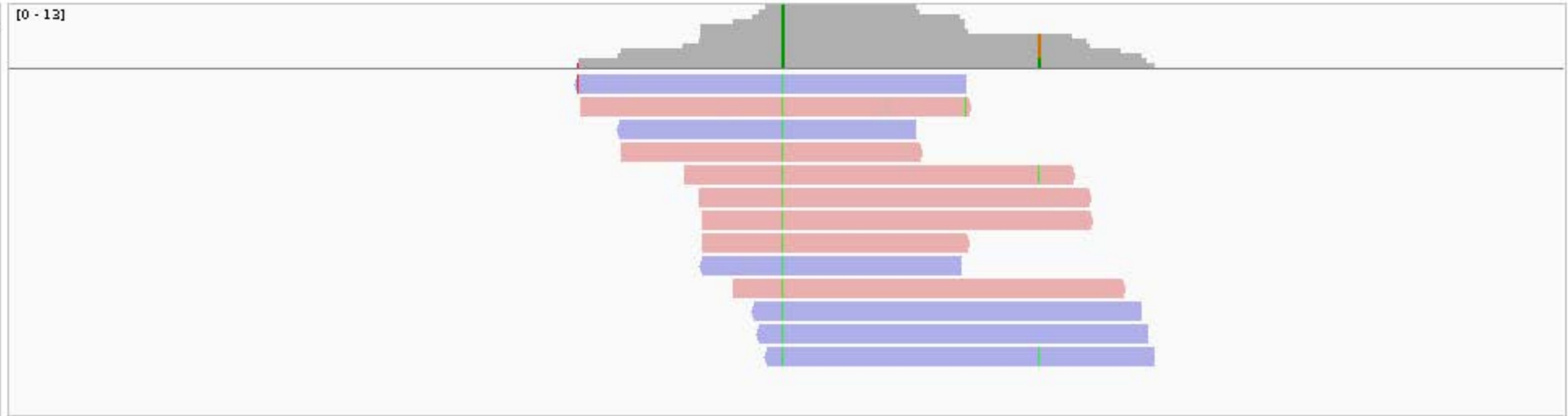
Refseq Genes





SL_1010_L_FFPE_12-B_mut.bam
range

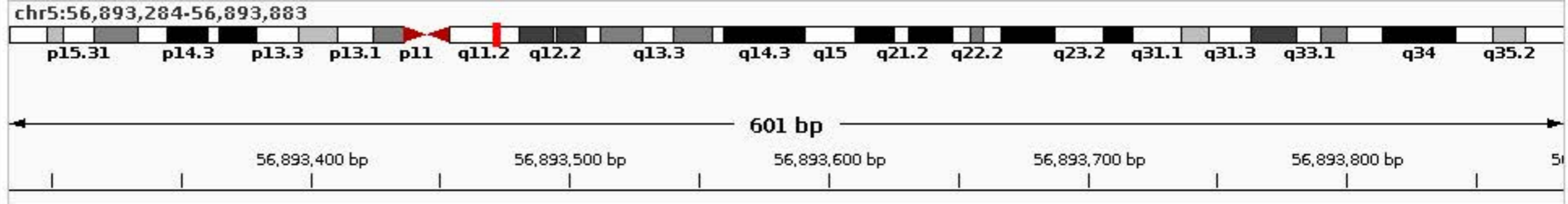
SL_1010_L_FFPE_12-B_mut.bam



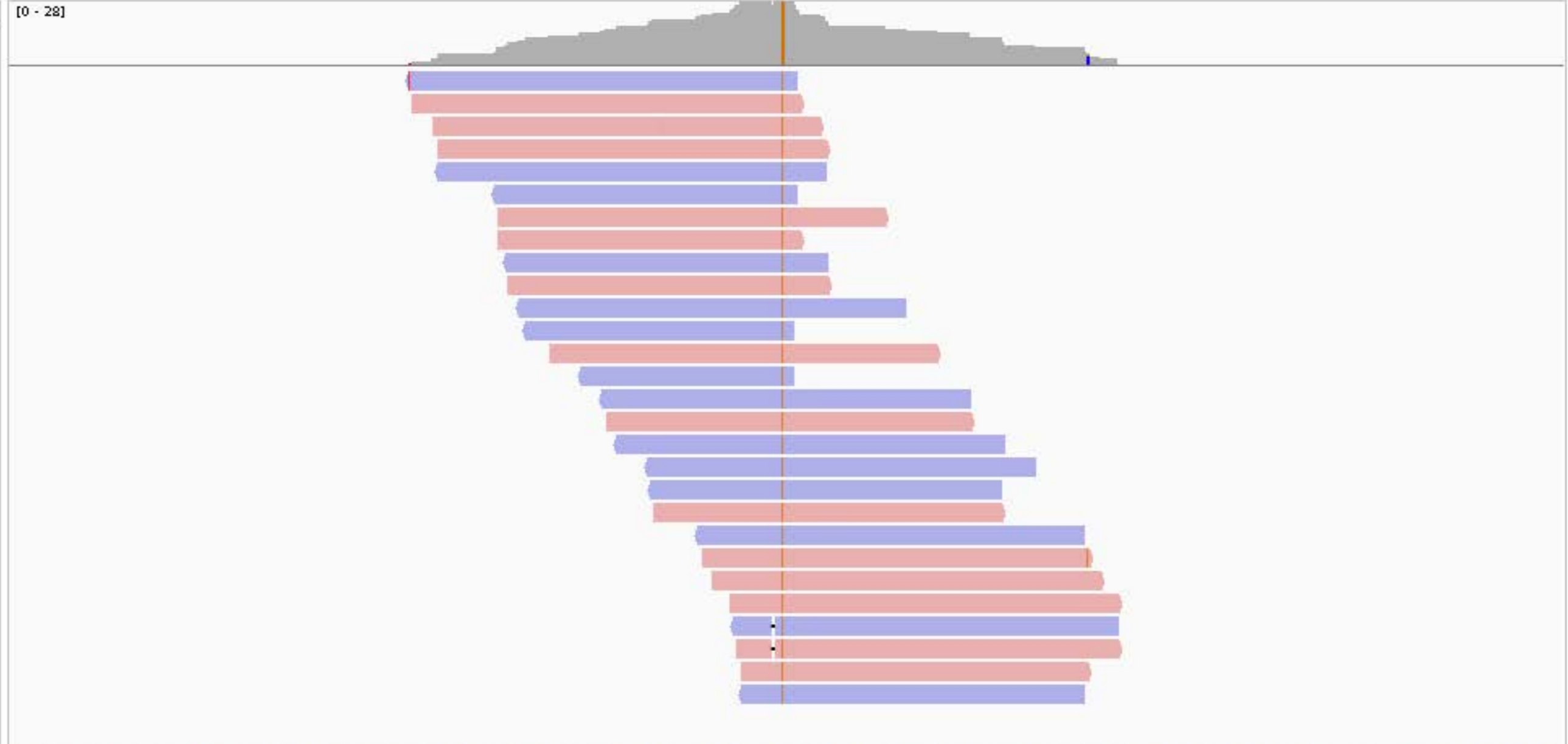
Sequence →

Refseq Genes





SL_1010_L_FFPE_10-B_mut.bam
rage



SL_1010_L_FFPE_10-B_mut.ba
m

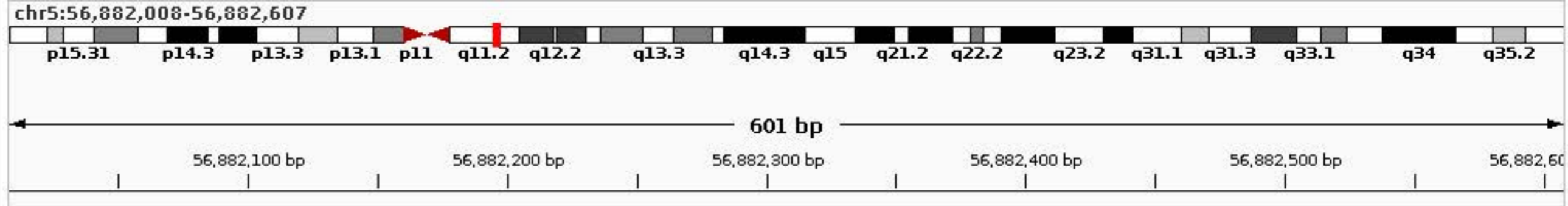
Sequence →



Refseq Genes

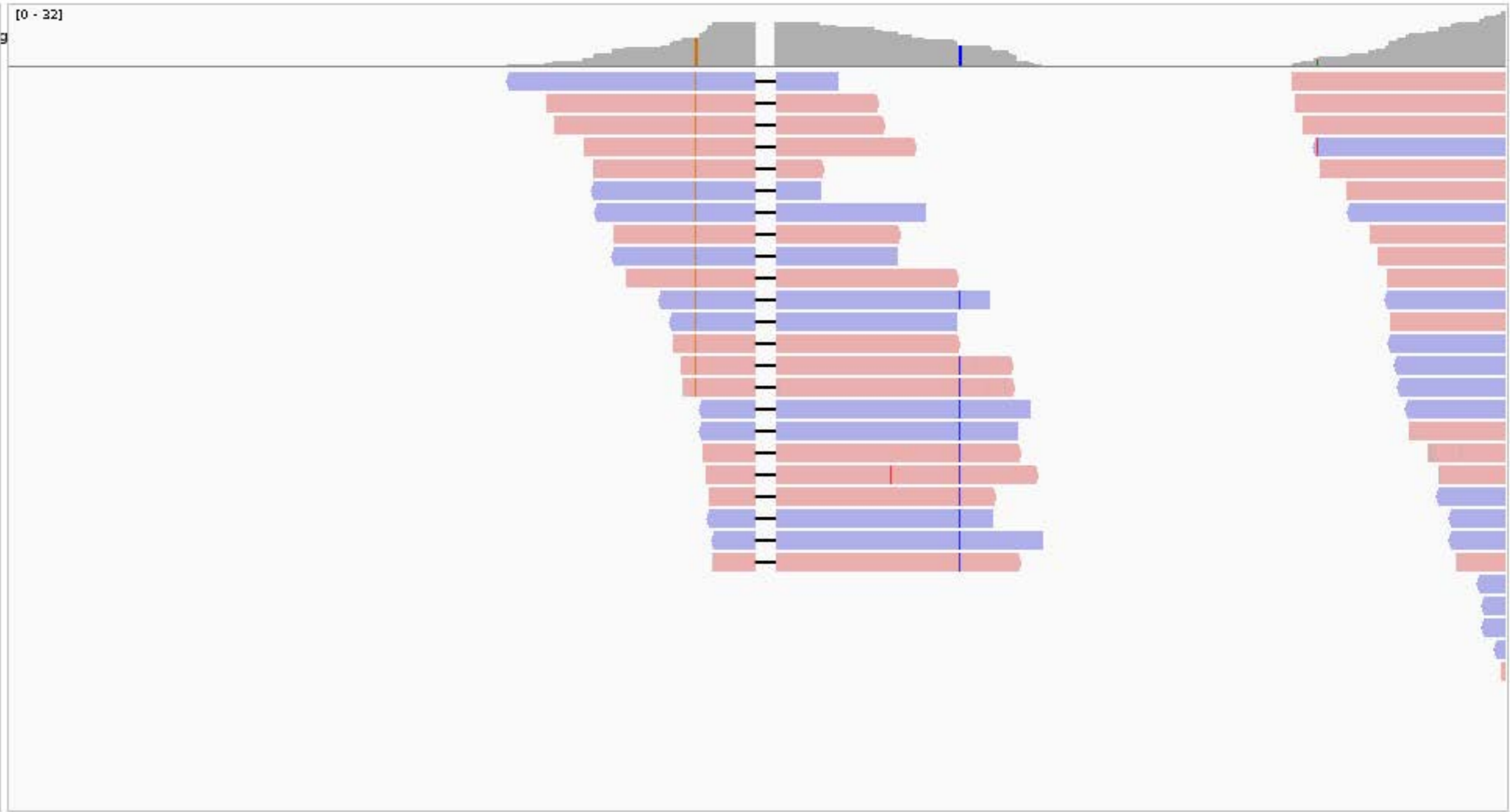


MAP3K1



SL_1010-N6-B_mut.bam Coverage

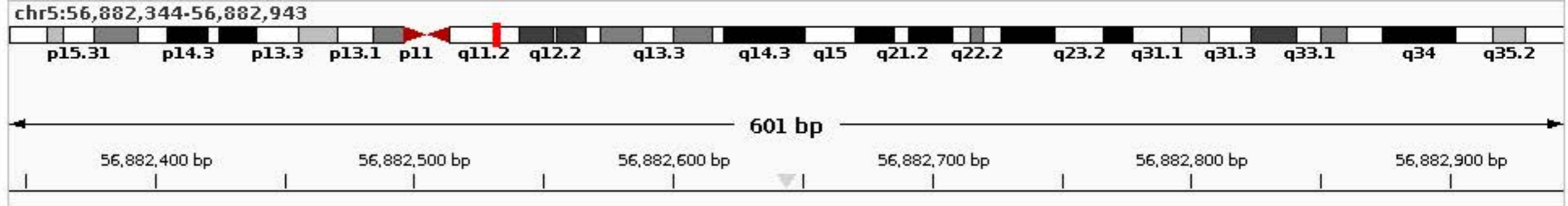
SL_1010-N6-B_mut.bam



Sequence

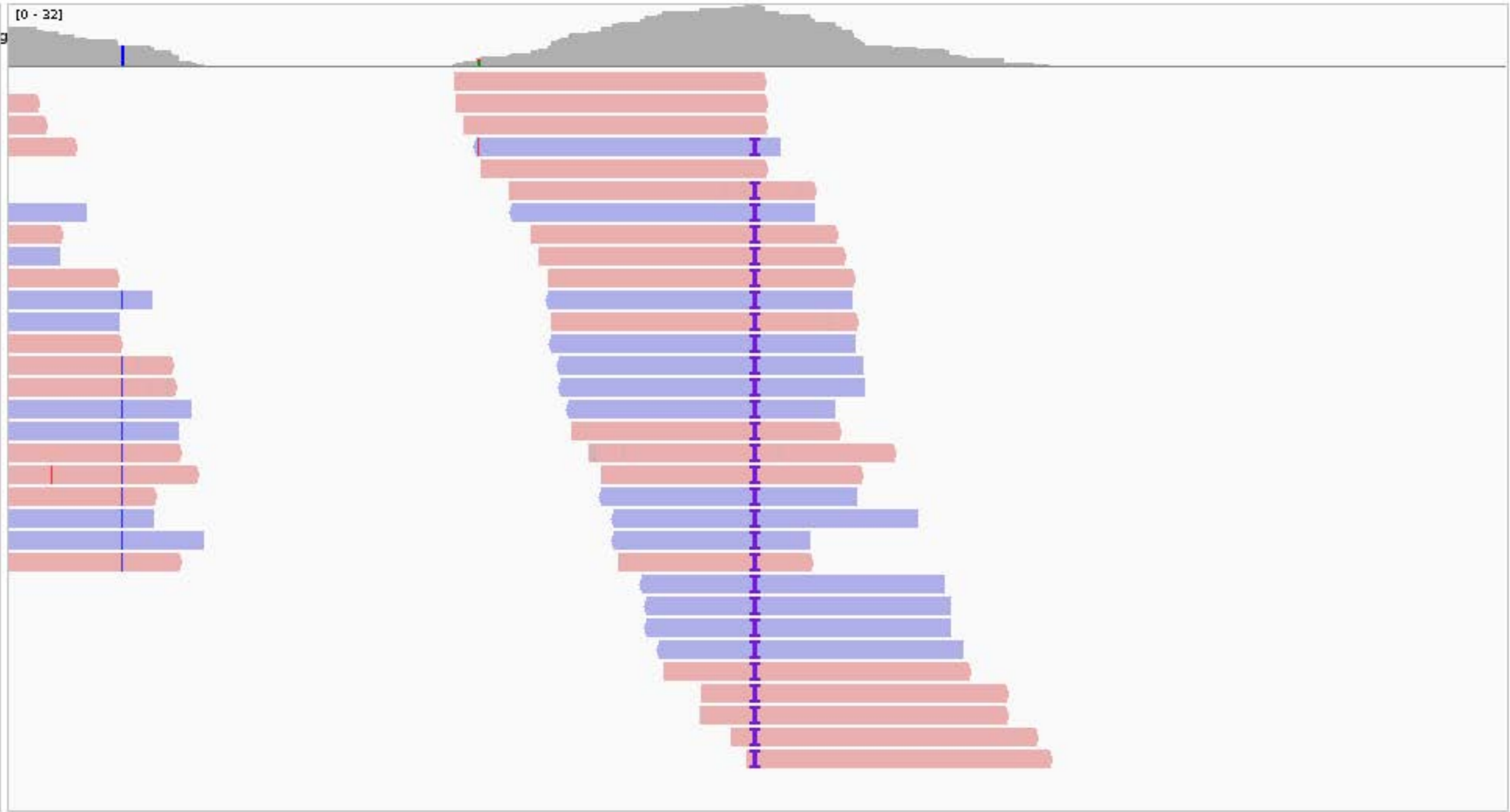
Refseq Genes





SL_1010-N6-B_mut.bam Coverage

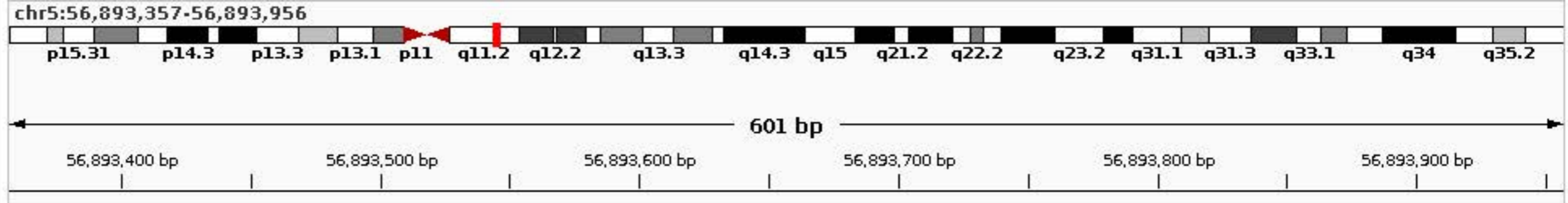
SL_1010-N6-B_mut.bam



Sequence

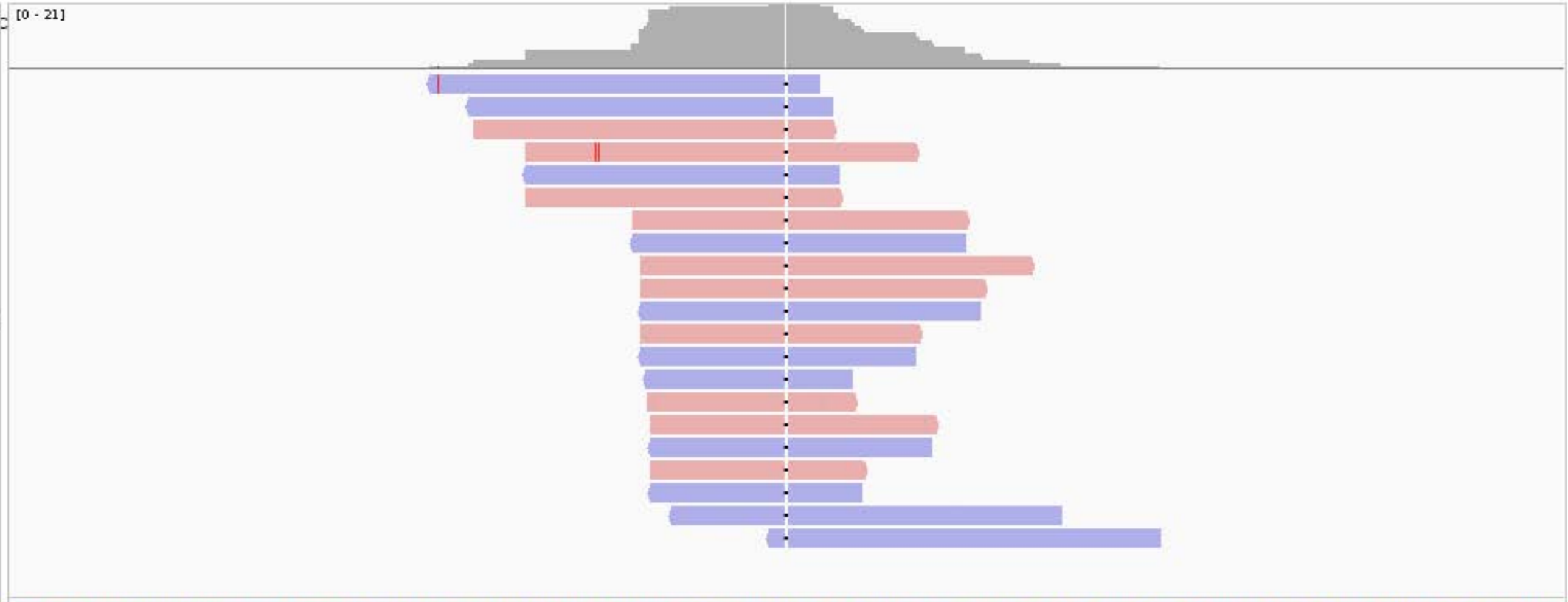
Refseq Genes





SL_1010_L_FFPE_9-B_mut.bam
age

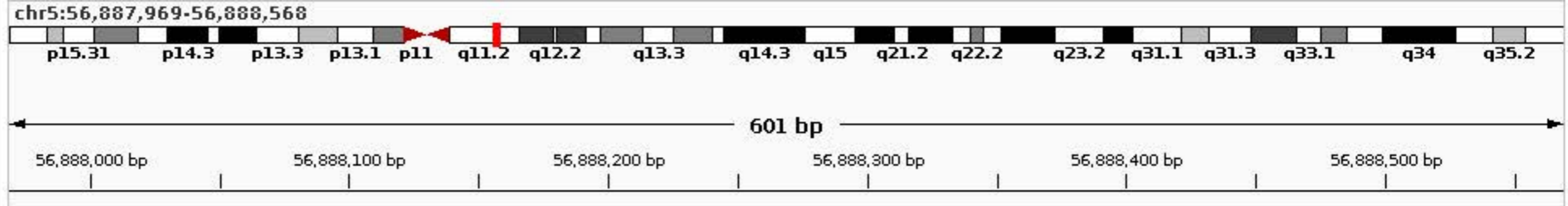
SL_1010_L_FFPE_9-B_mut.bam



Sequence →

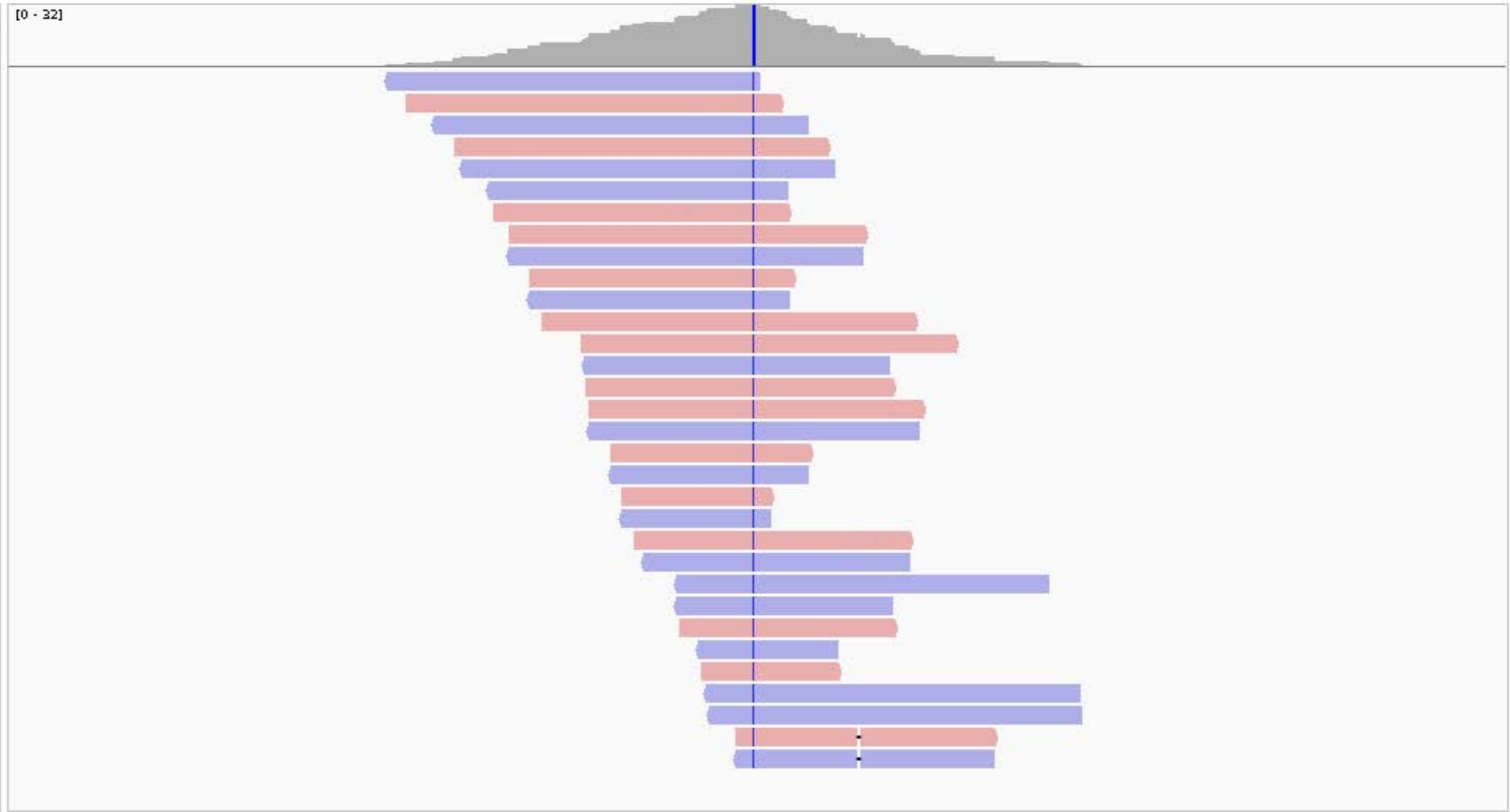
Refseq Genes





SL_1010_L_FFPE_13-B_mut.barr
rage

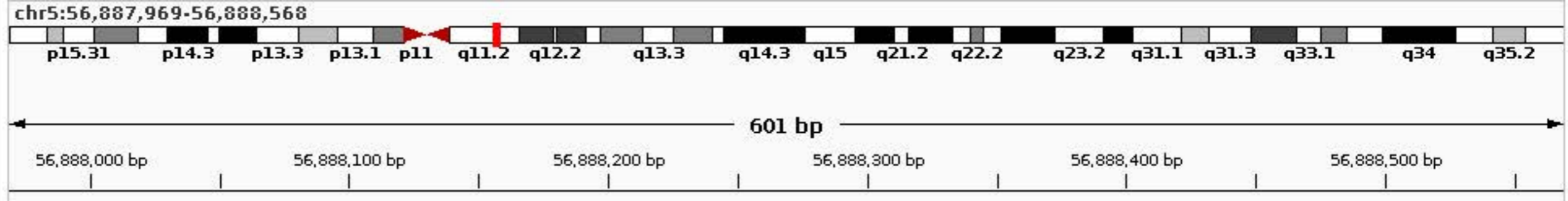
SL_1010_L_FFPE_13-B_mut.ba
m



Sequence →

Refseq Genes



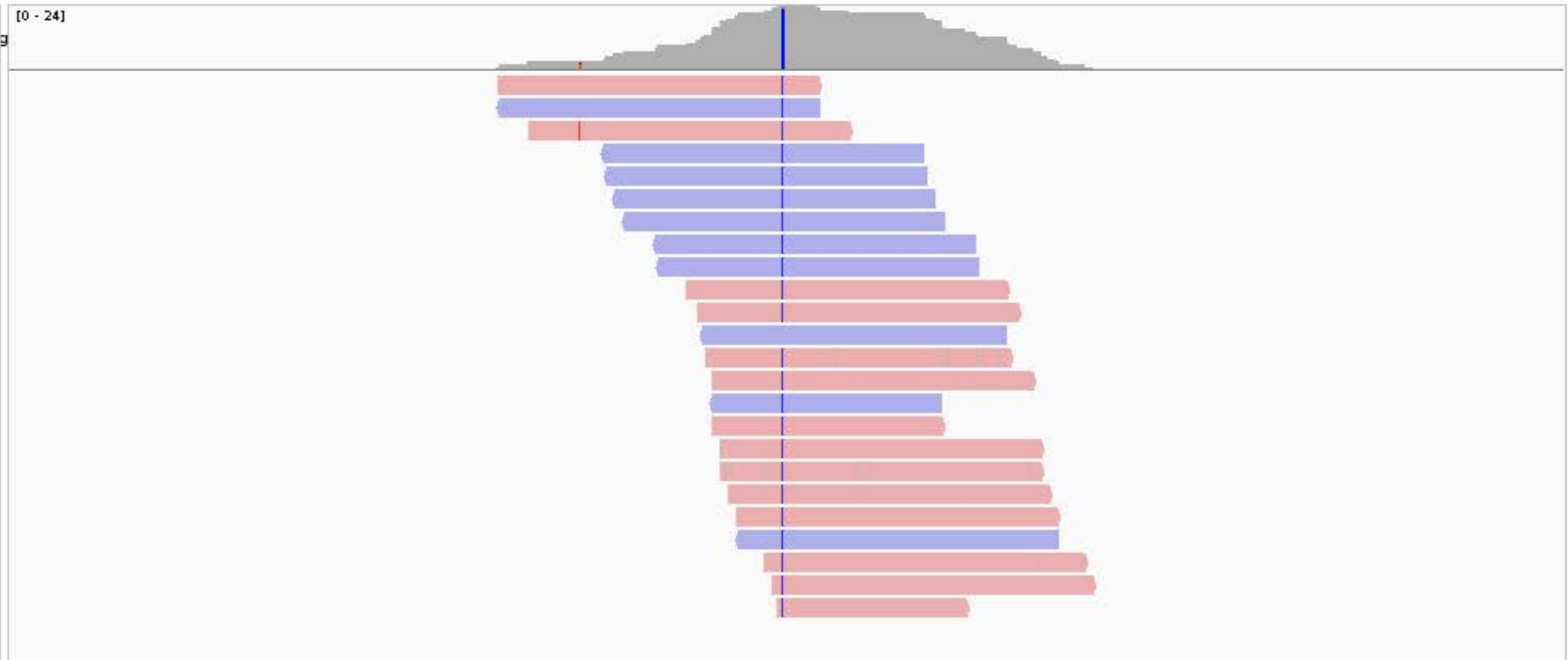


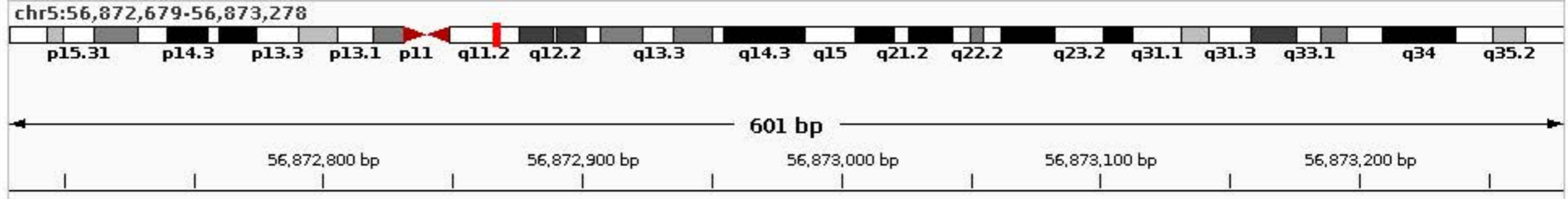
SL_1010-N6-B_mut.bam Coverage

SL_1010-N6-B_mut.bam

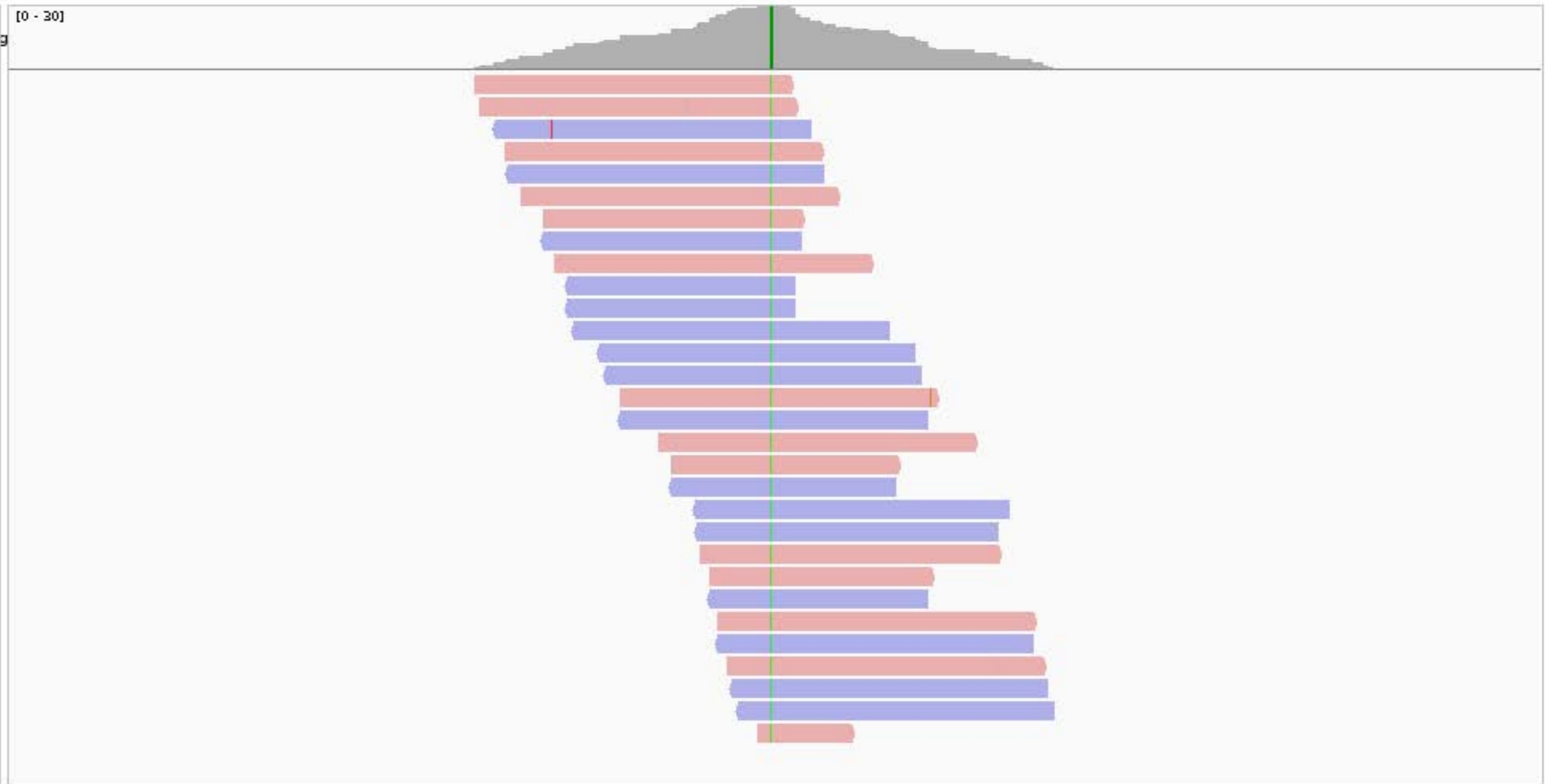
Sequence →

Refseq Genes





SL_1010-N4-B_mut.bam Coverage



SL_1010-N4-B_mut.bam

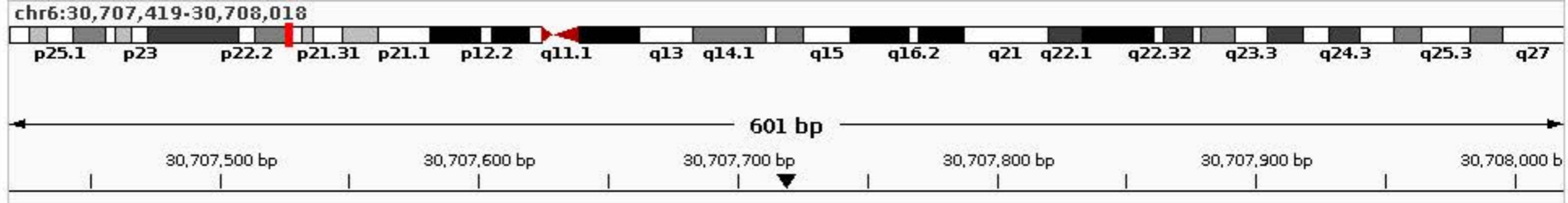
Sequence →



Refseq Genes



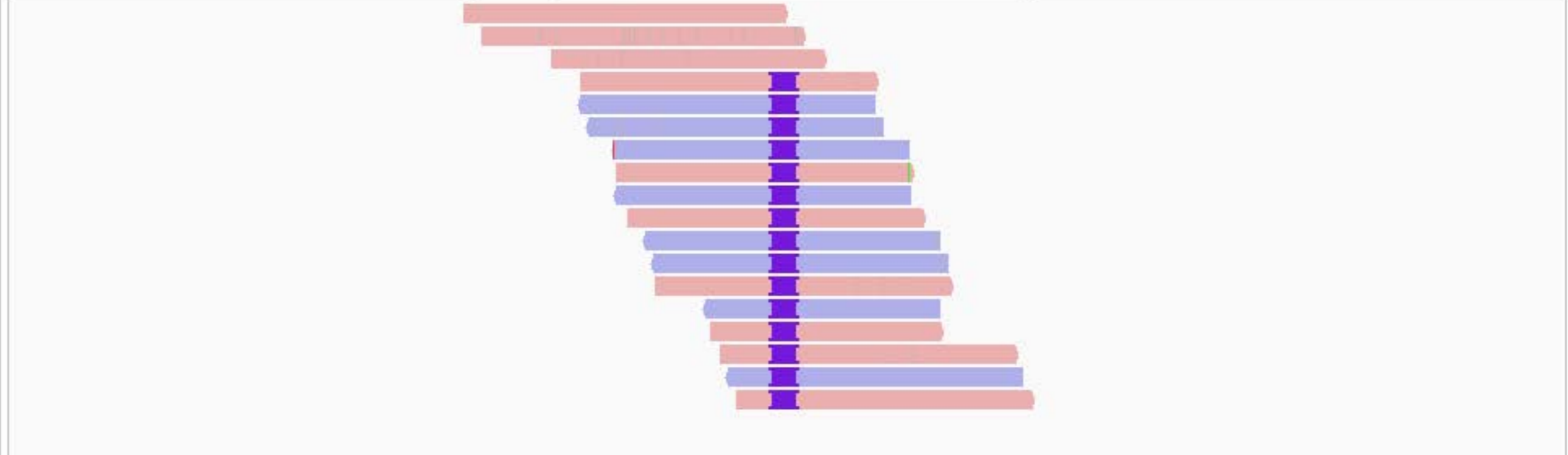
MAP3K1



SL_0005-LT_11-B_mut.bam Cover



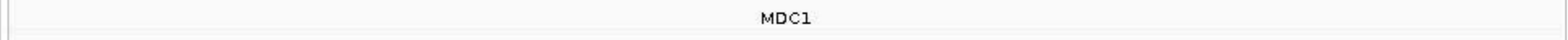
SL_0005-LT_11-B_mut.bam



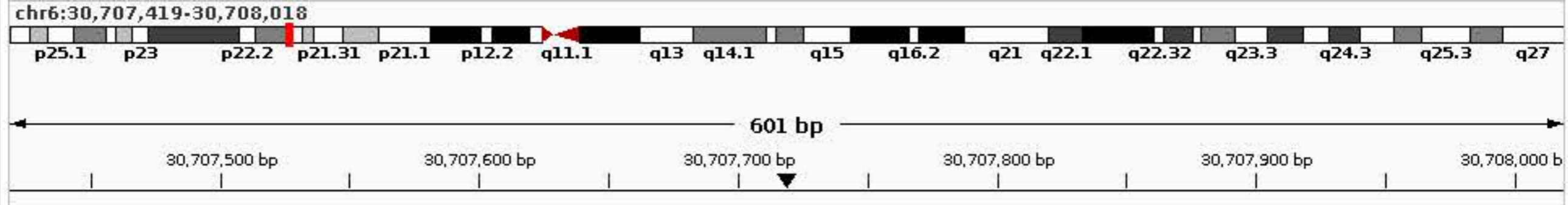
Sequence



Refseq Genes



MDC1

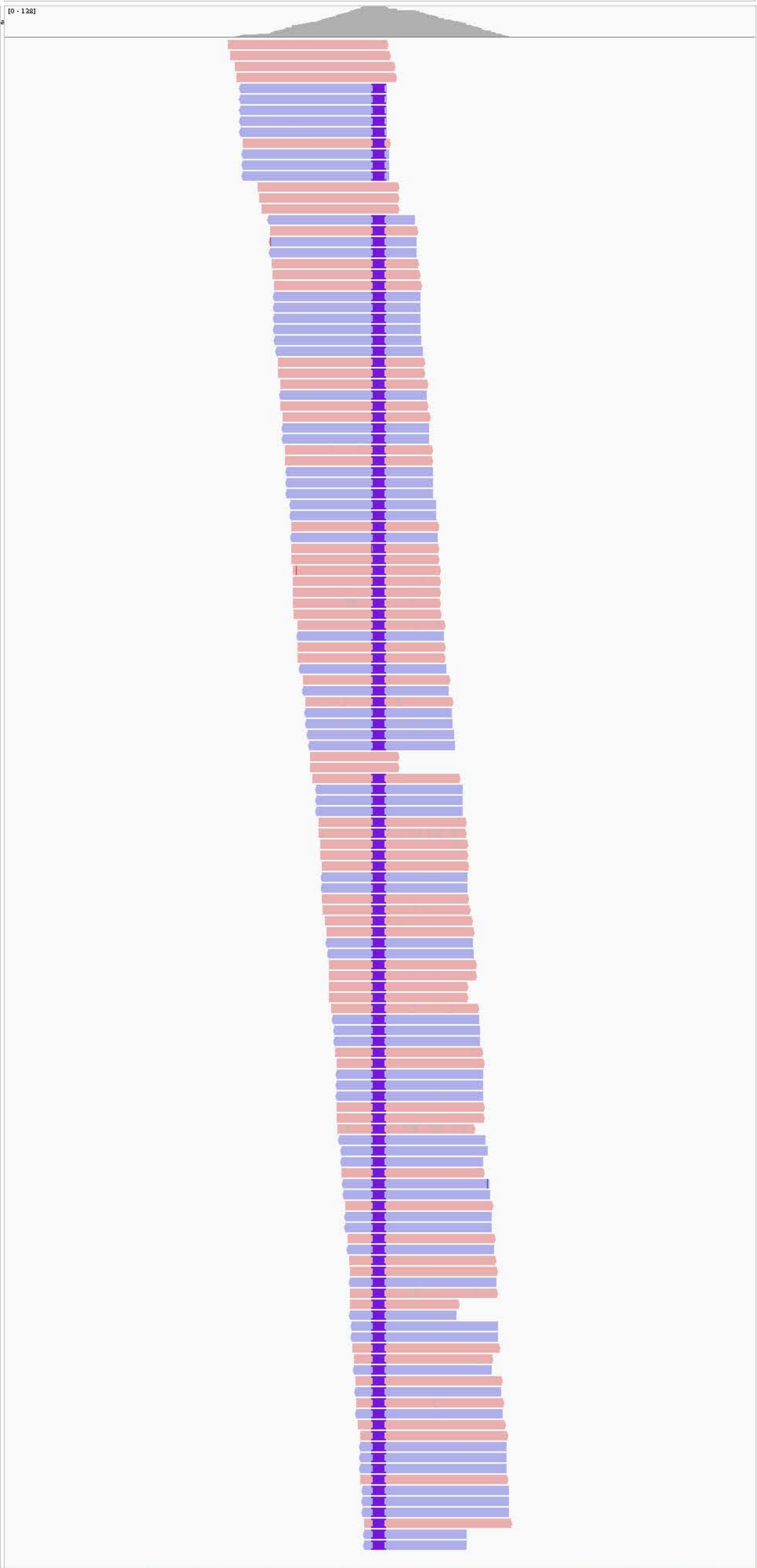


SL_0005-LT-4-B_mut.bam Coverage

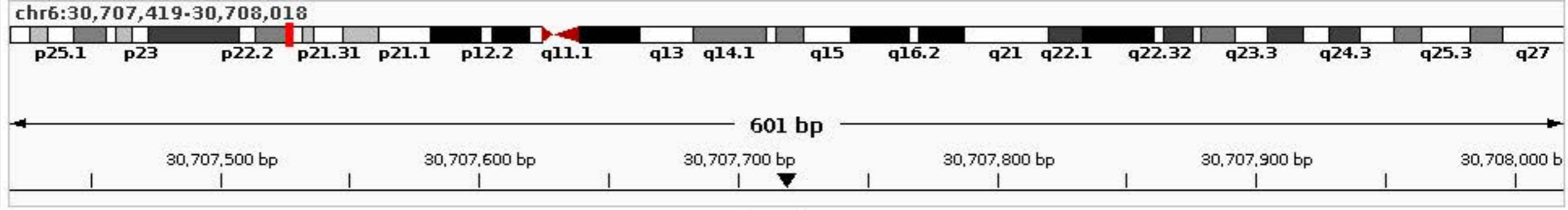
SL_0005-LT-4-B_mut.bam

Sequence

Refseq Genes

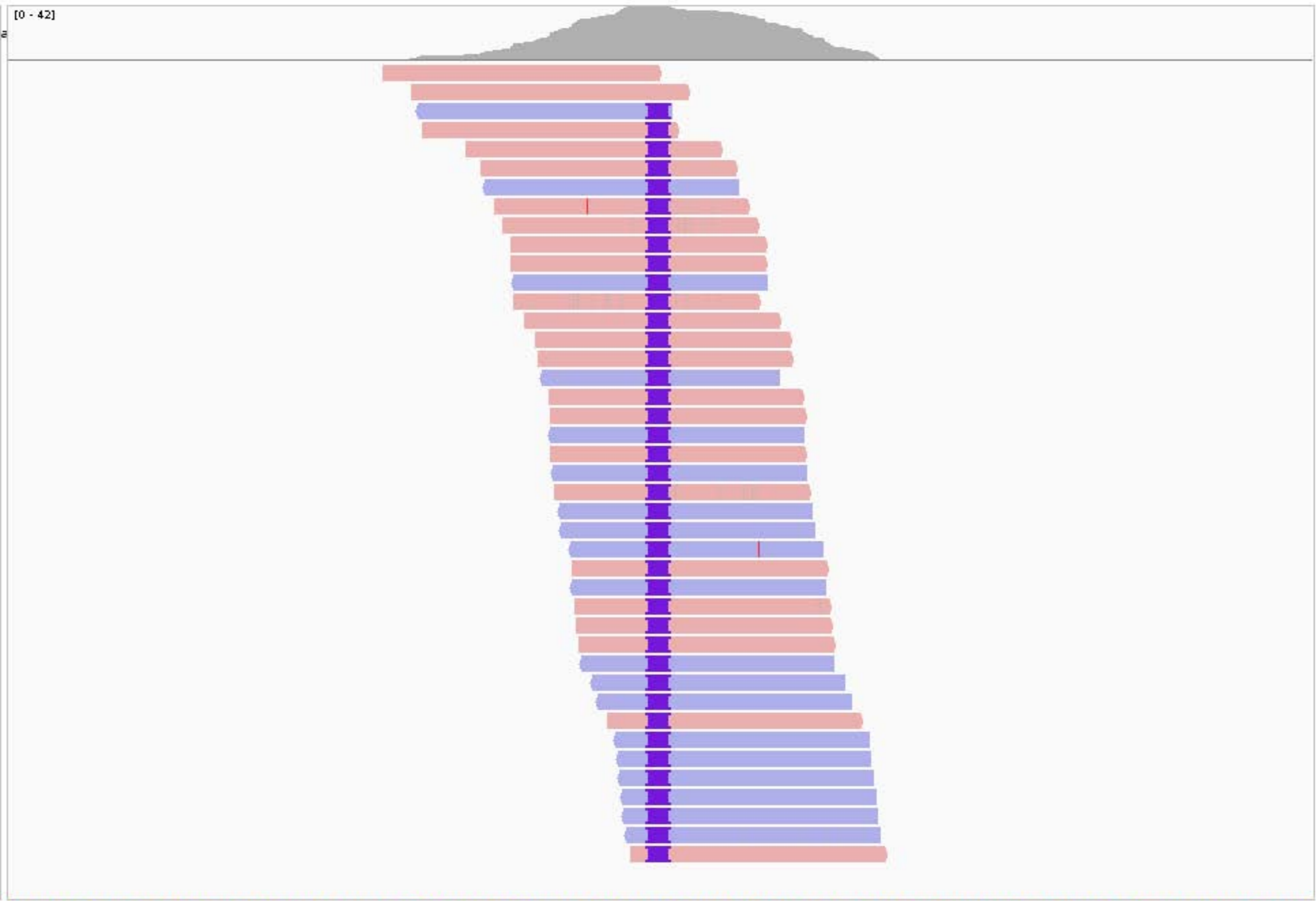


MDC1



SL_0005-LT-9-B_mut.bam Coverage

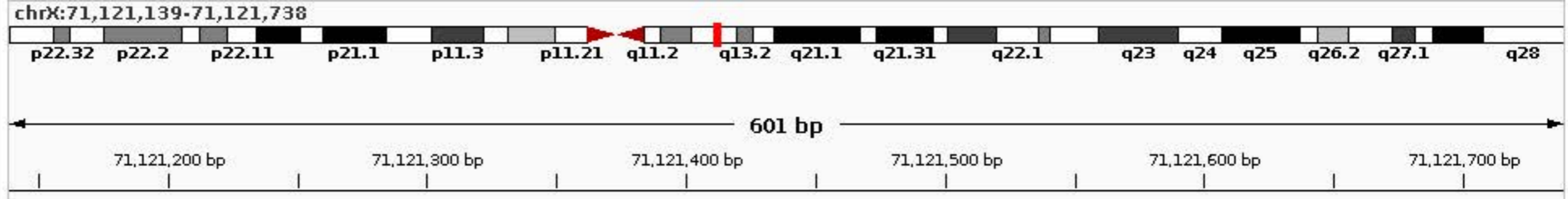
SL_0005-LT-9-B_mut.bam



Sequence →

Refseq Genes



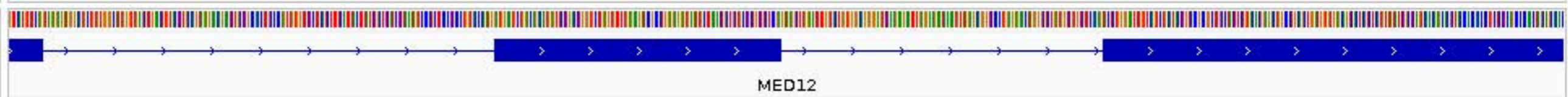
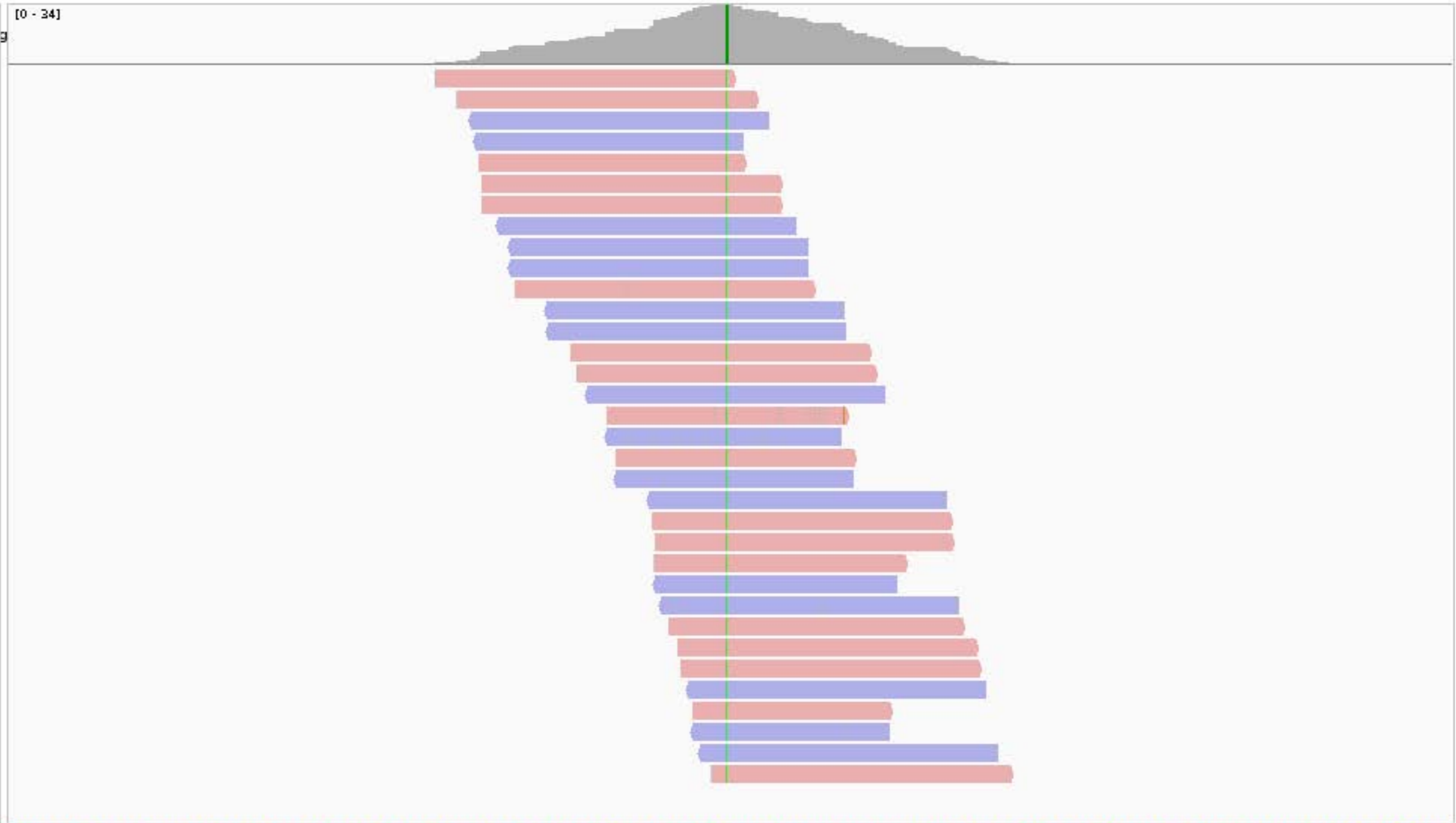


SL_1012-N2-B_mut.bam Coverage

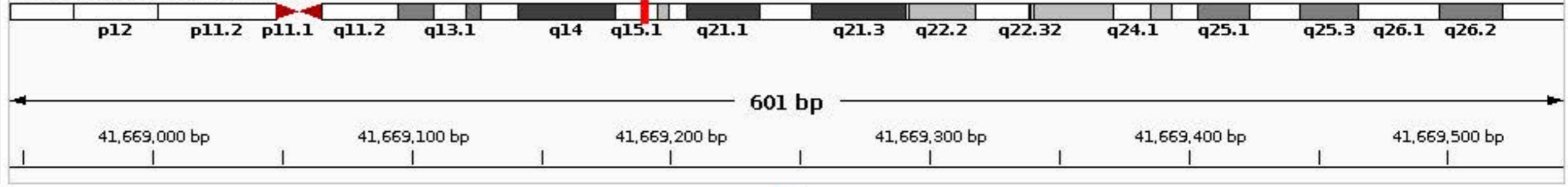
SL_1012-N2-B_mut.bam

Sequence

Refseq Genes

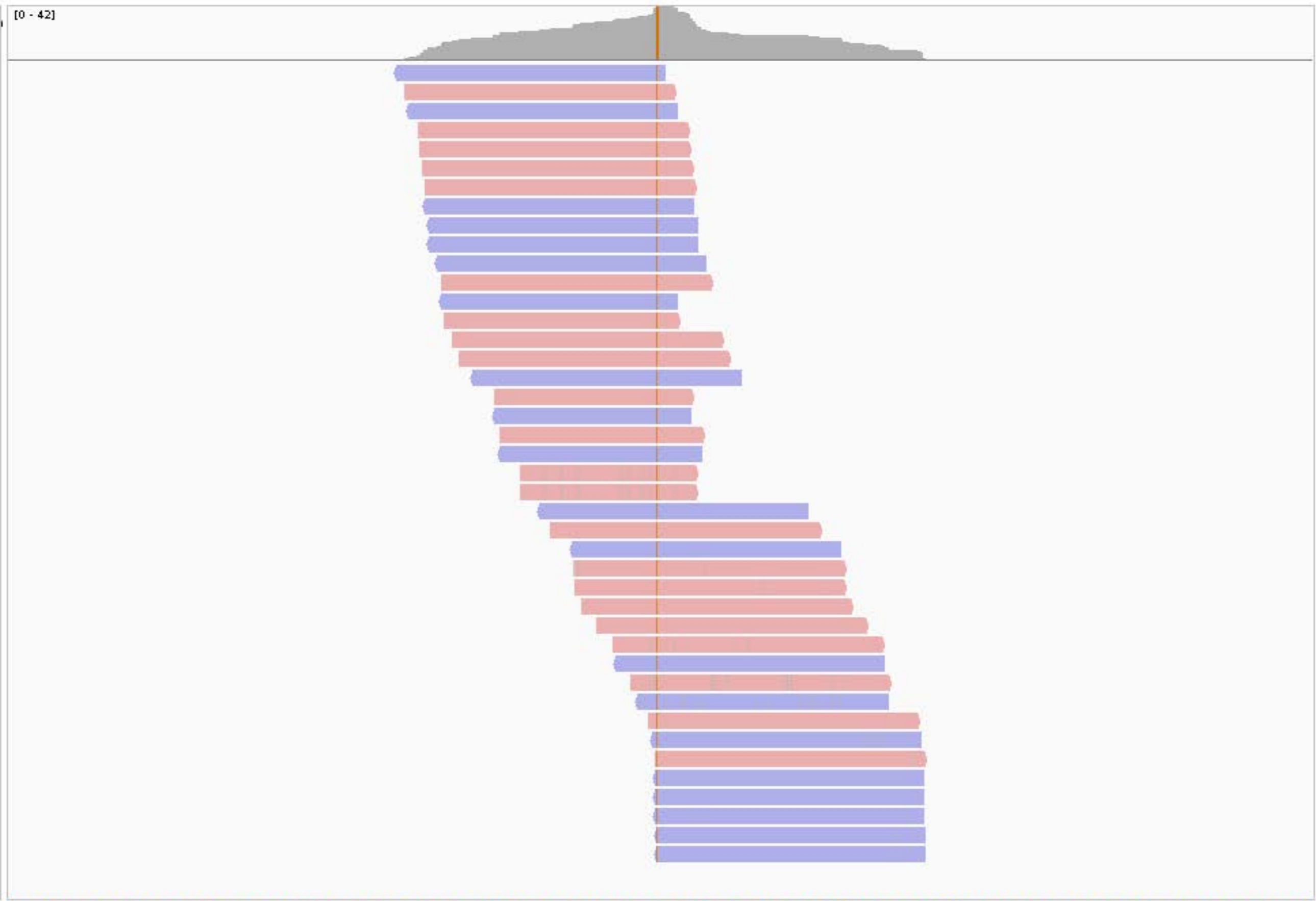


chr15:41,668,945-41,669,544



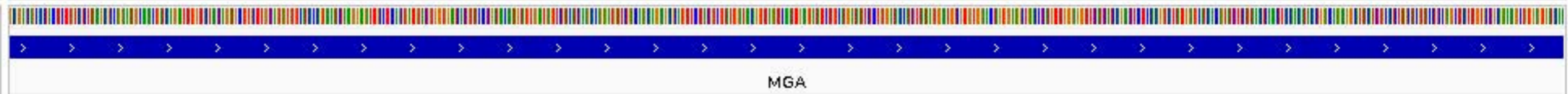
SL_1012_R_10_FFPE-B_mut.bam
rage

SL_1012_R_10_FFPE-B_mut.ba
m

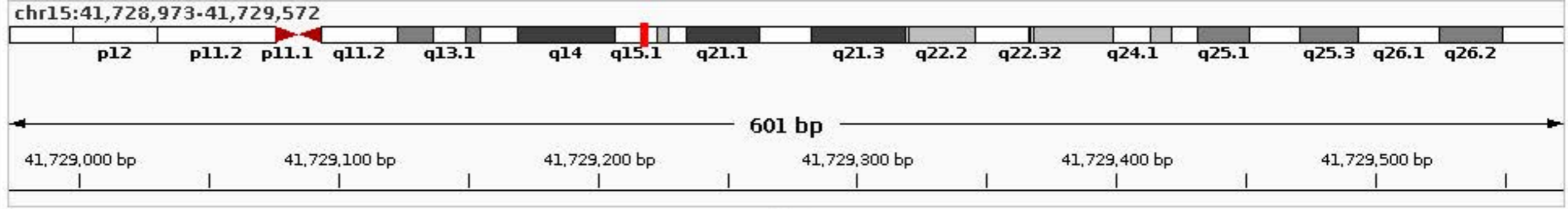


Sequence →

Refseq Genes

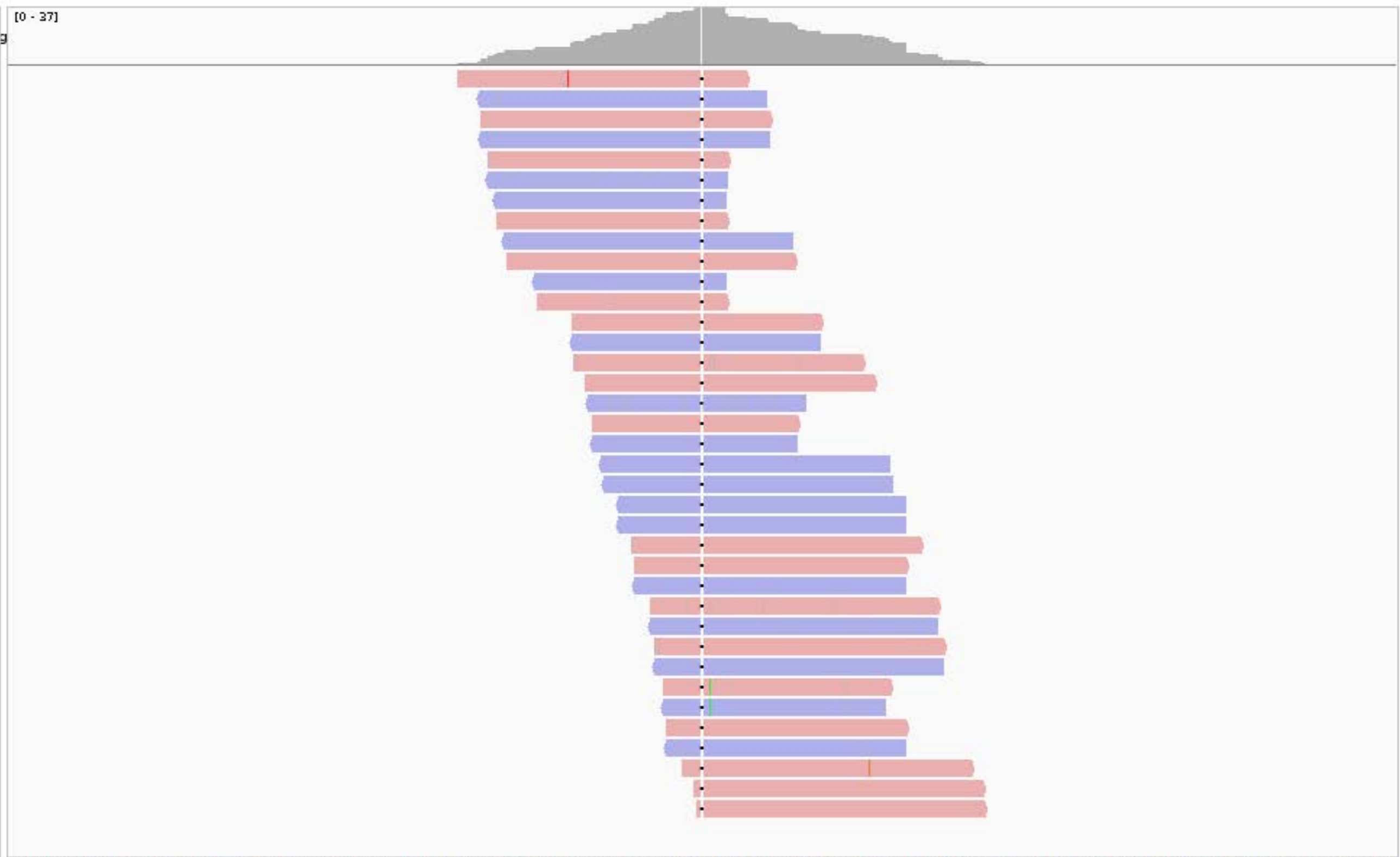


MGA



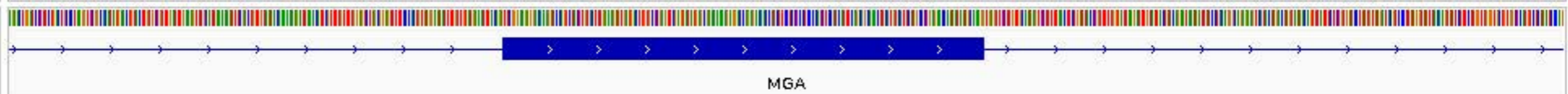
SL_1012-N1-B_mut.bam Coverage

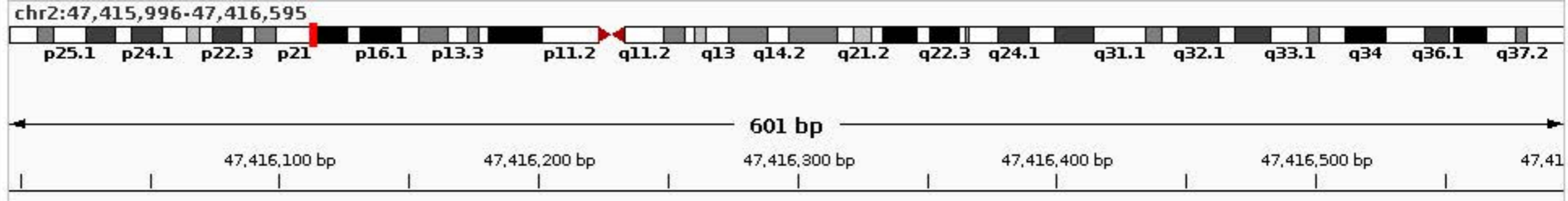
SL_1012-N1-B_mut.bam



Sequence →

Refseq Genes



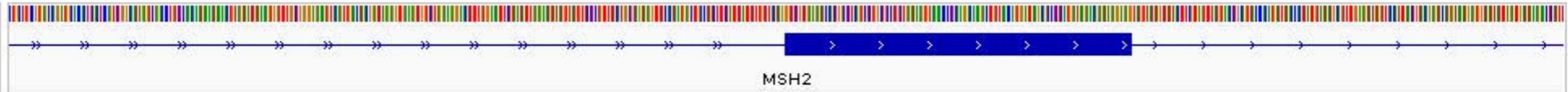
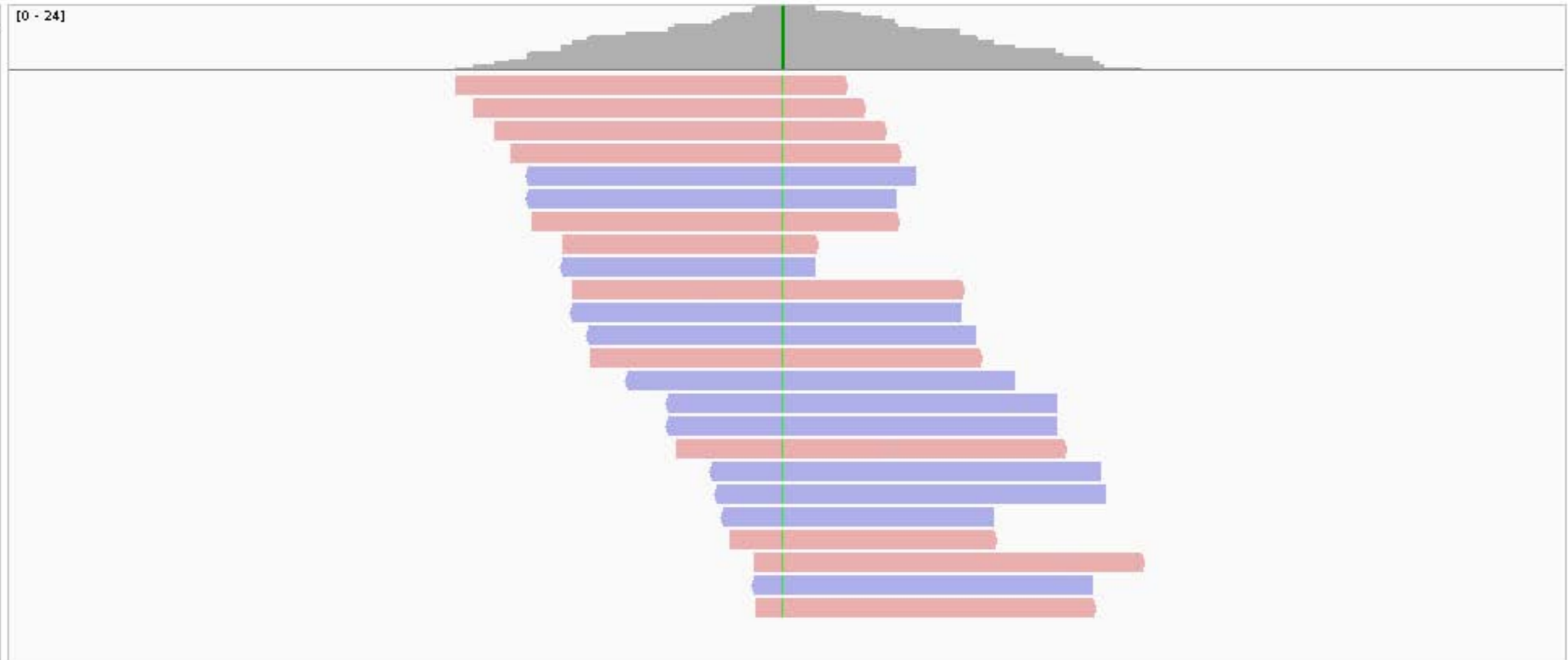


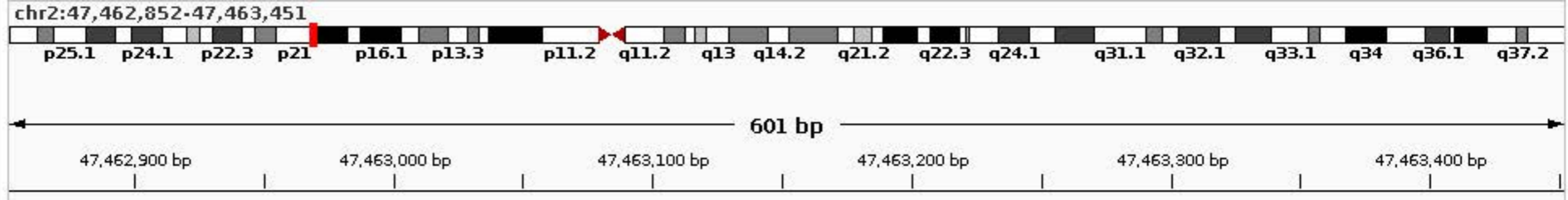
SL_0007_L_FFPE_13-B_mut.bam
rage

SL_0007_L_FFPE_13-B_mut.ba
m

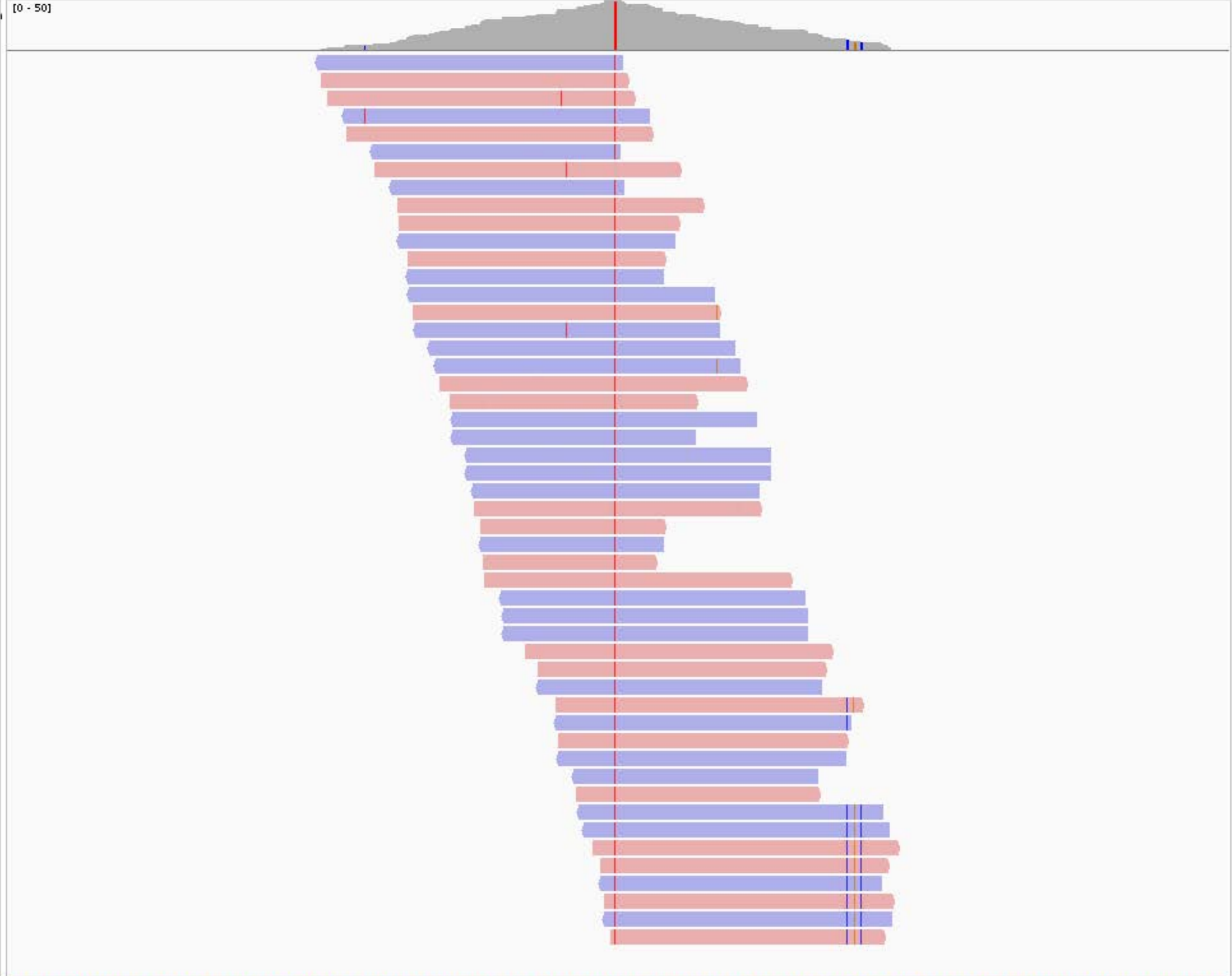
Sequence →

Refseq Genes





SL_1005_R_FFPE_59-B_mut.bam
 range



SL_1005_R_FFPE_59-B_mut.ba
 m

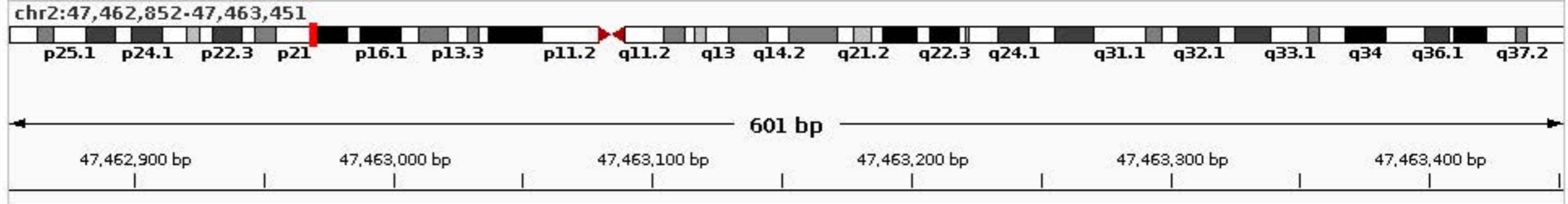
Sequence →



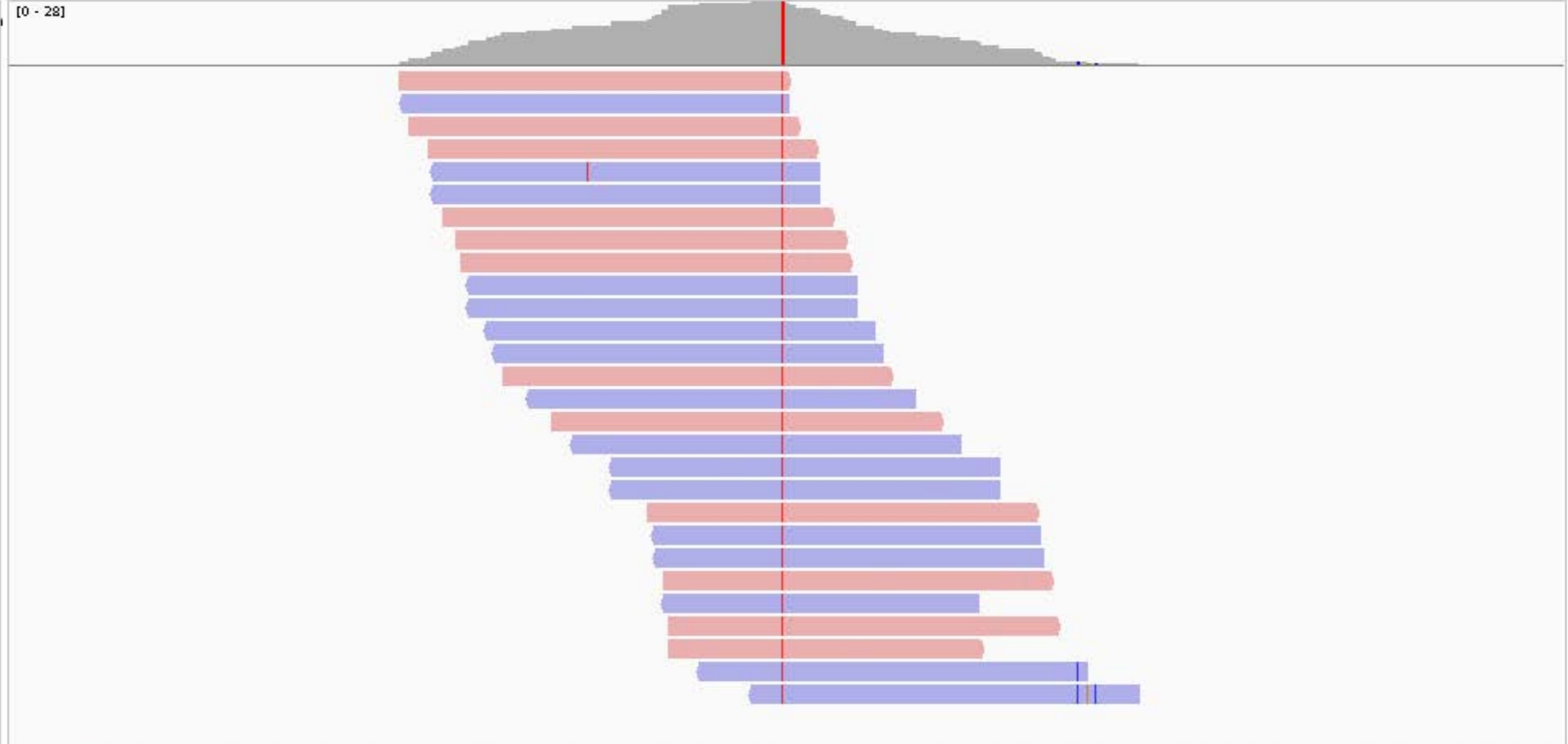
Refseq Genes



MSH2



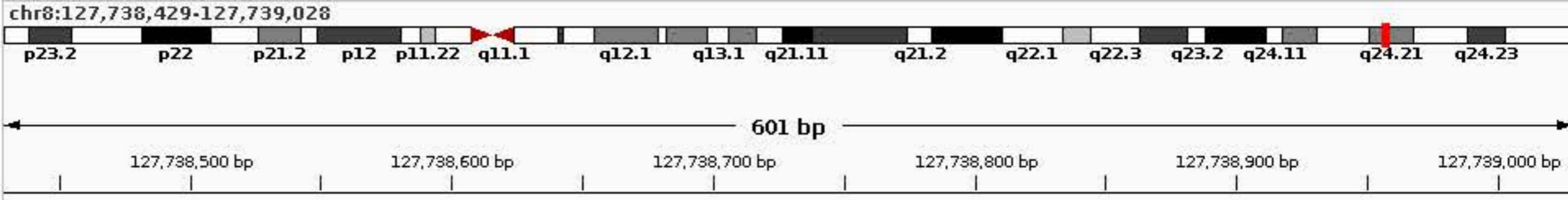
SL_1005_R_FFPE_66-B_mut.bam
rage



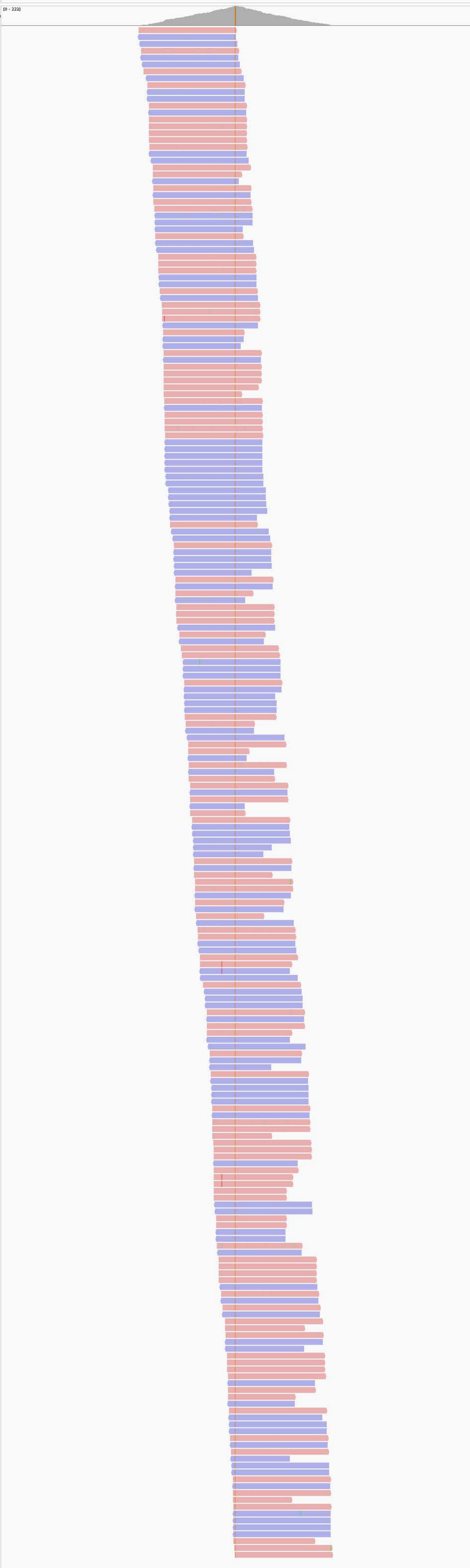
SL_1005_R_FFPE_66-B_mut.ba
m

Sequence →
Refseq Genes





SL_1010-T1-B_mut.bam Coverage



SL_1010-T1-B_mut.bam

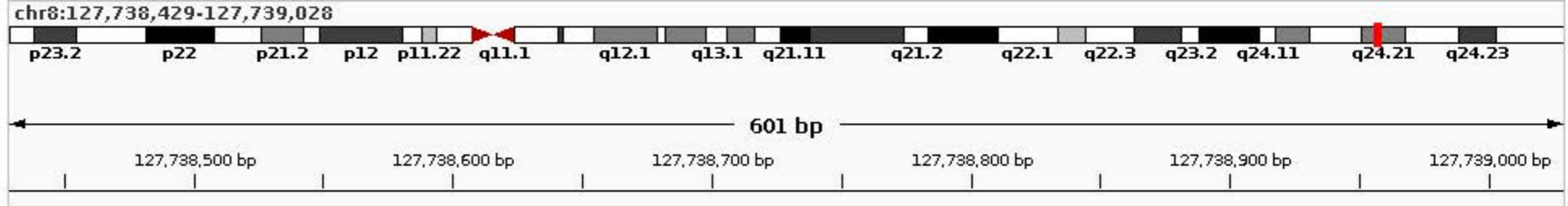
Sequence



Refseq Genes

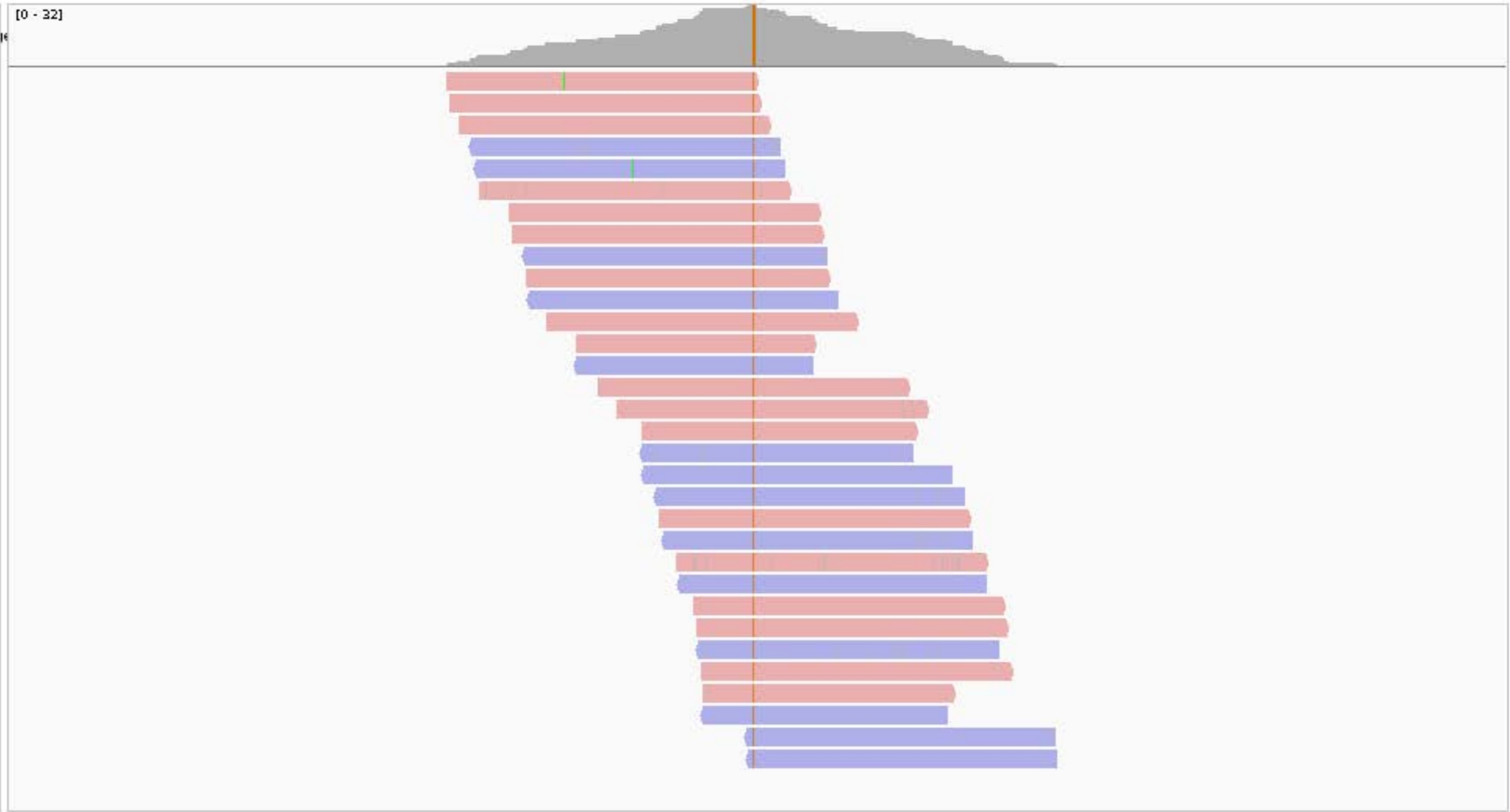


MYC



SL_1010-T2-B_mut.bam Coverage

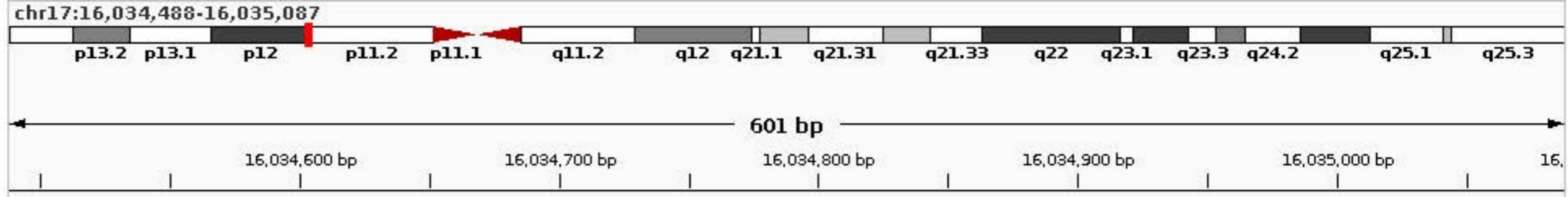
SL_1010-T2-B_mut.bam



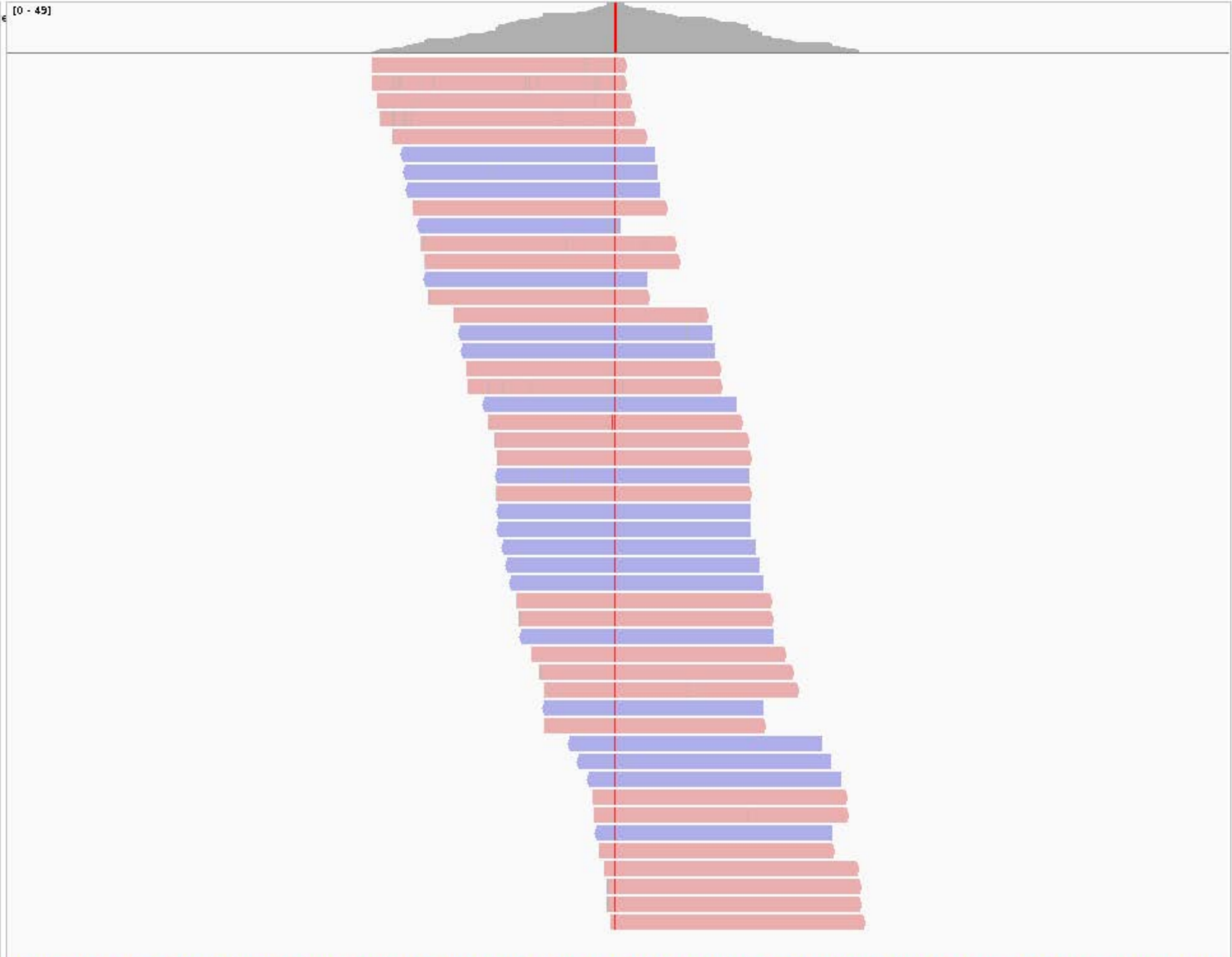
Sequence

Refseq Genes





SL_0005-RN_11-B_mut.bam Cov



SL_0005-RN_11-B_mut.bam

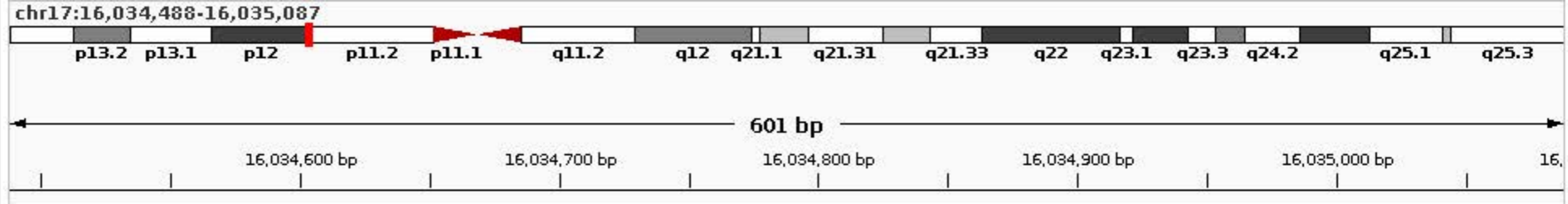
Sequence



Refseq Genes



TTC19

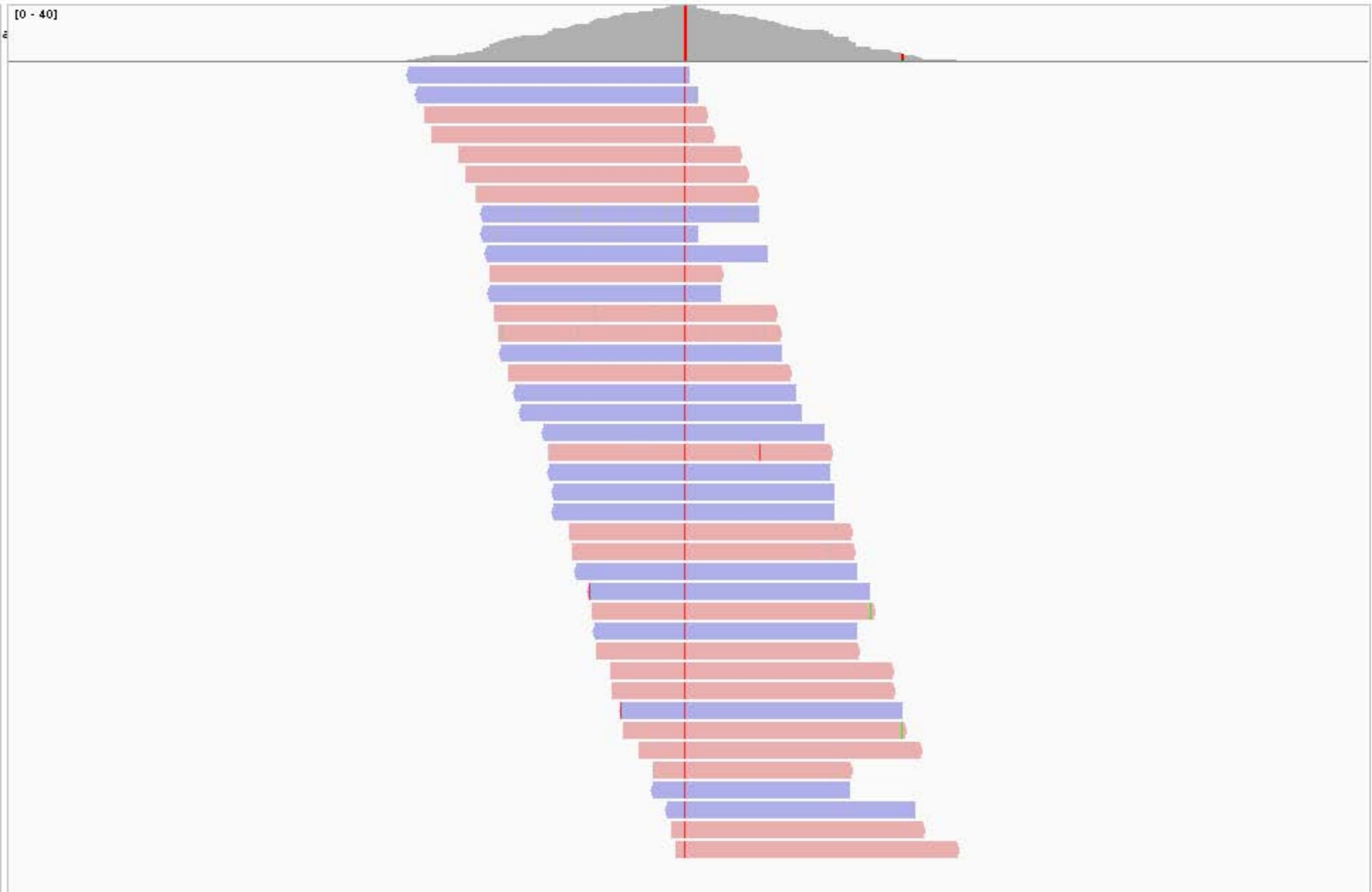


SL_0009-LT_1-B_mut.bam Coverage

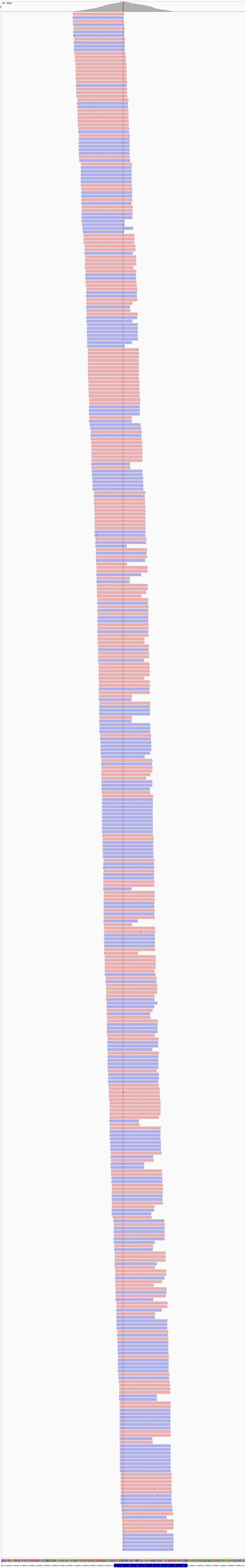
SL_0009-LT_1-B_mut.bam

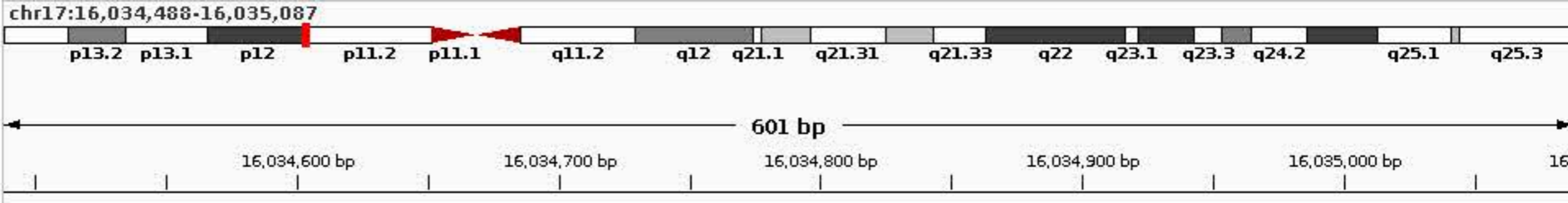
Sequence

Refseq Genes

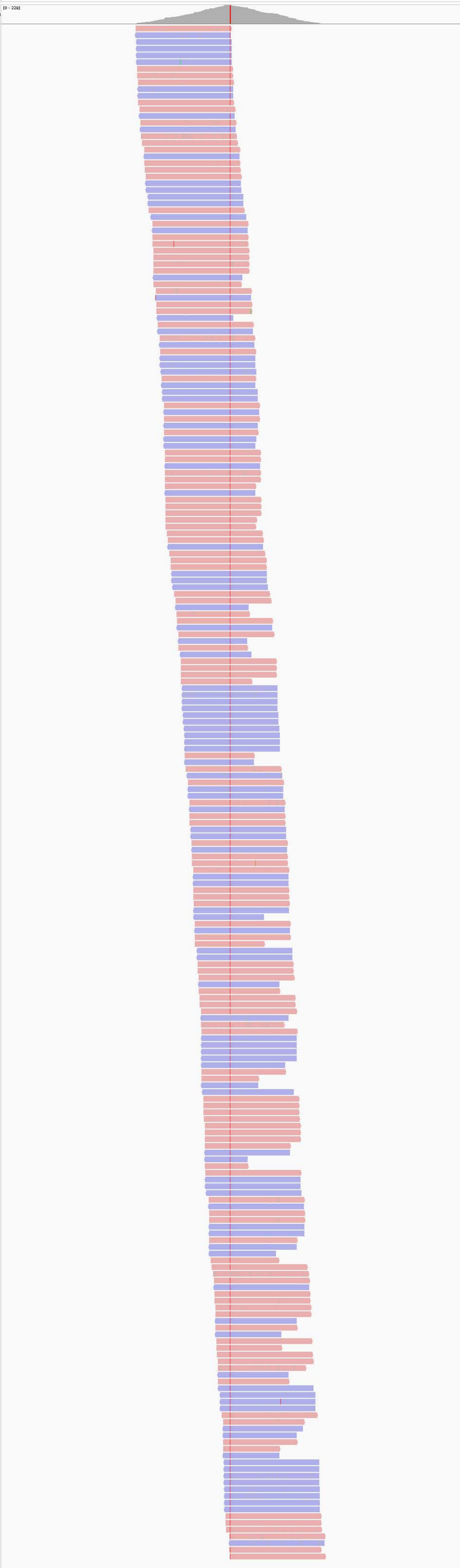


SL_0011-LT_2-8.bam Coverage
SL_0011-LT_2-8_rmt.bam



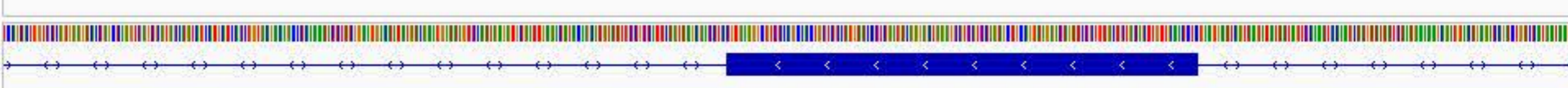


SL_0011-LT_5-B_mut.bam Coverage

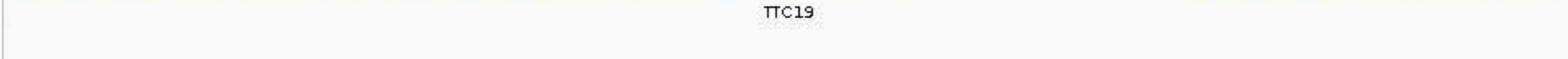


SL_0011-LT_5-B_mut.bam

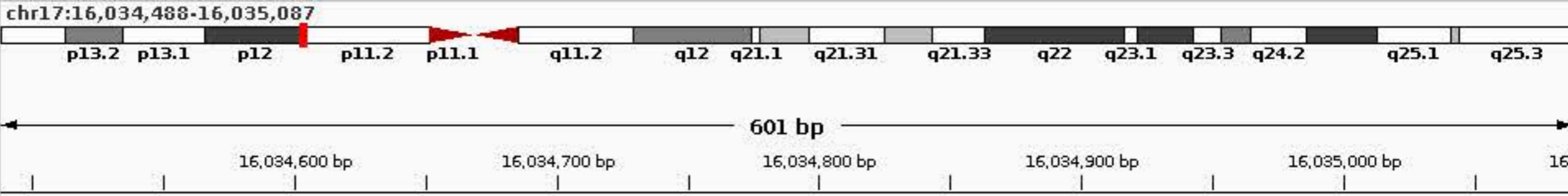
Sequence



Refseq Genes



TTC19

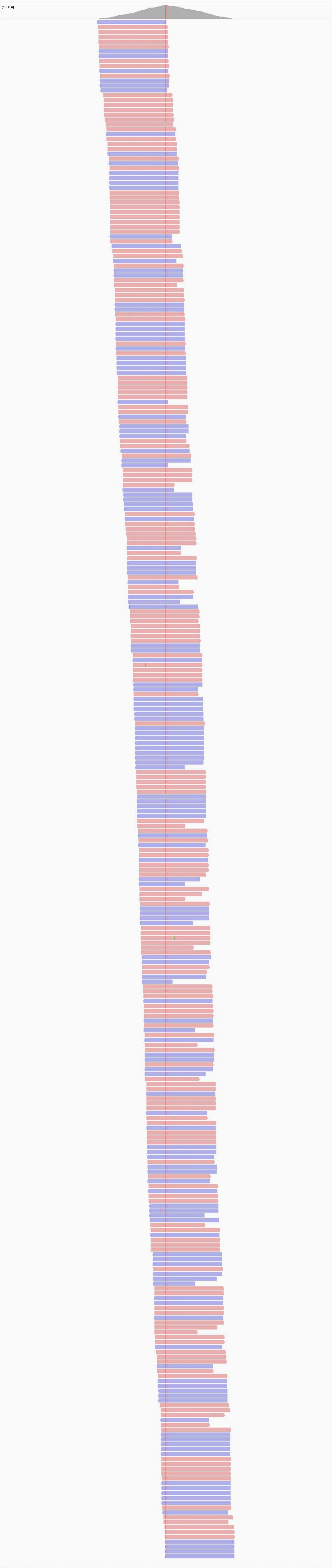


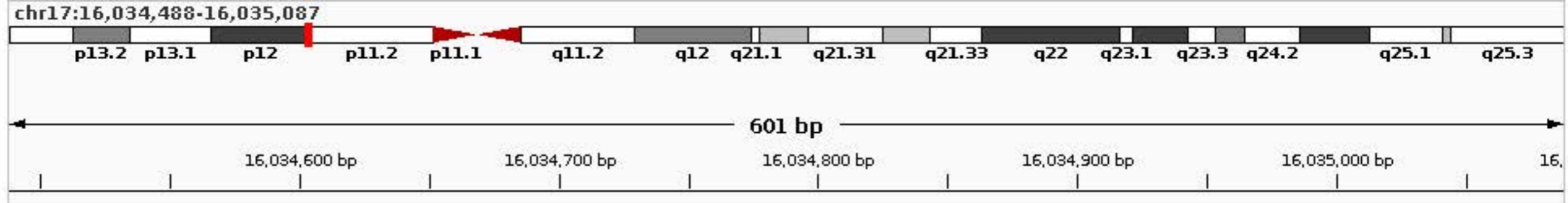
SL_0011-LT_6-B_mut.bam Coverage

SL_0011-LT_6-B_mut.bam

Sequence

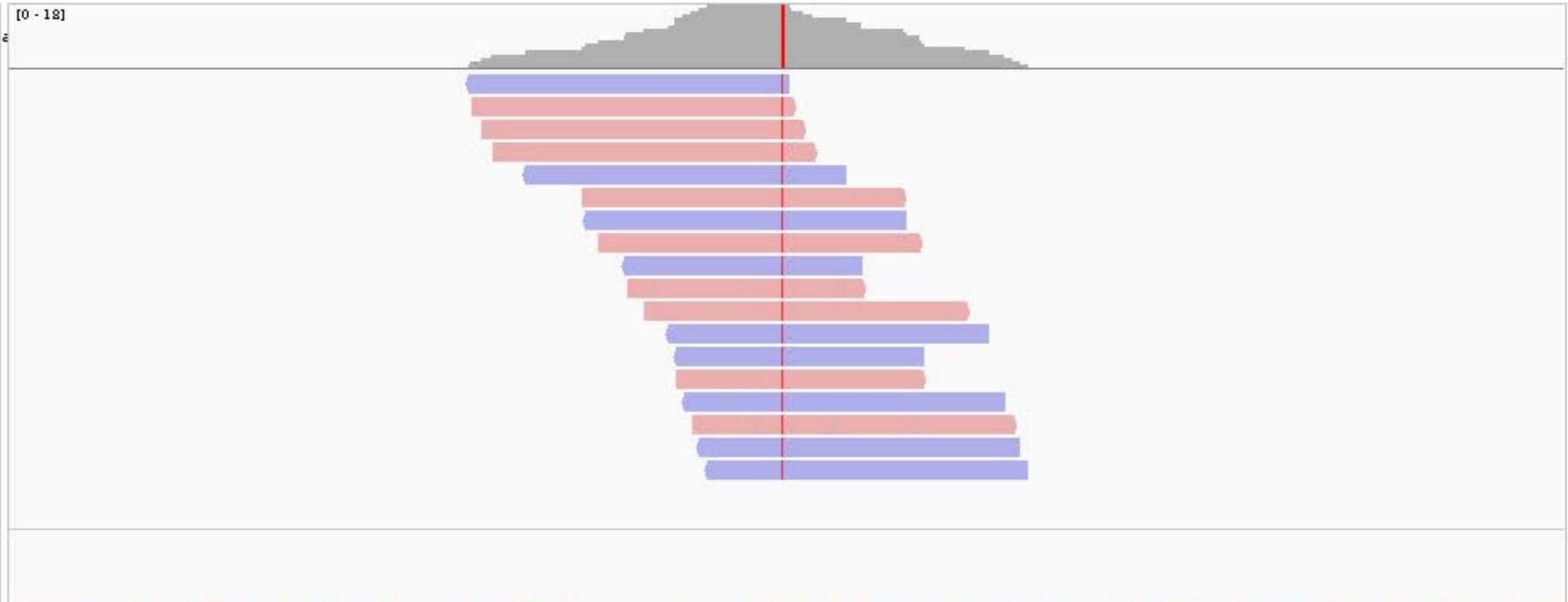
Refseq Genes





SL_0011-LT_7-B_mut.bam Coverage

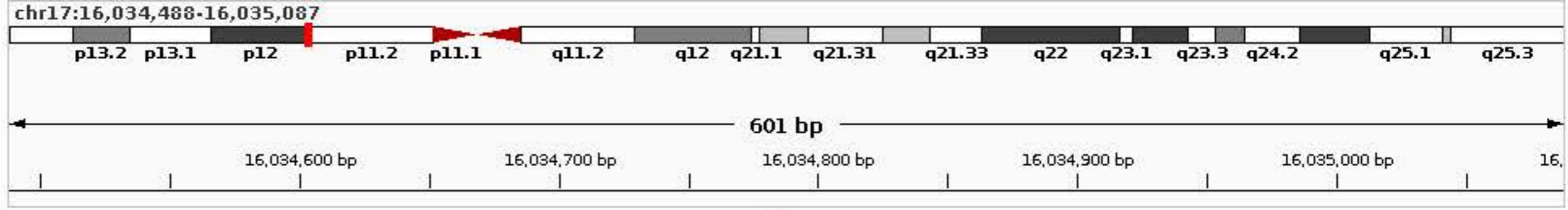
SL_0011-LT_7-B_mut.bam



Sequence →

Refseq Genes



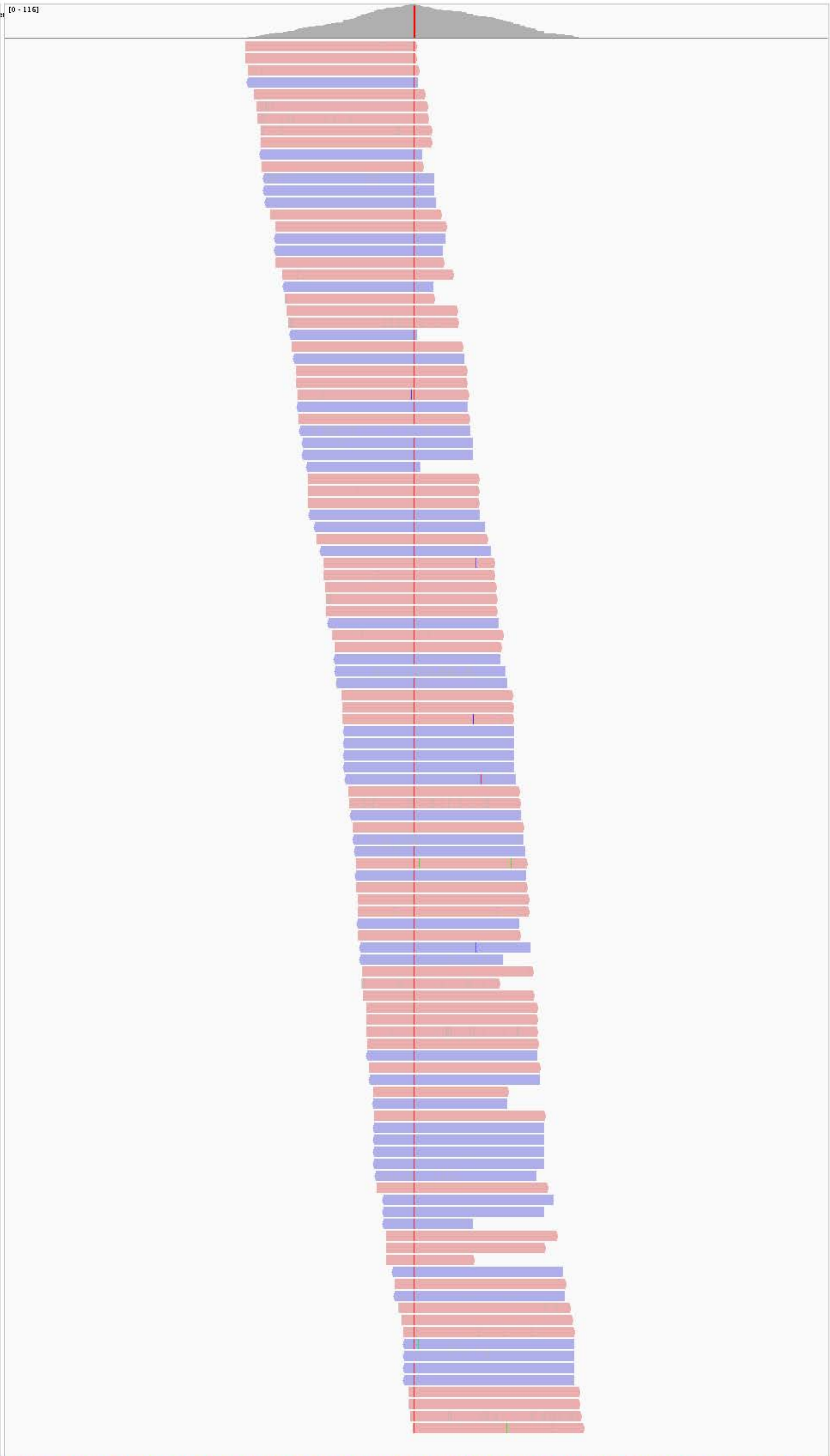


SL_0011-LT_11-B_mut.bam Cov

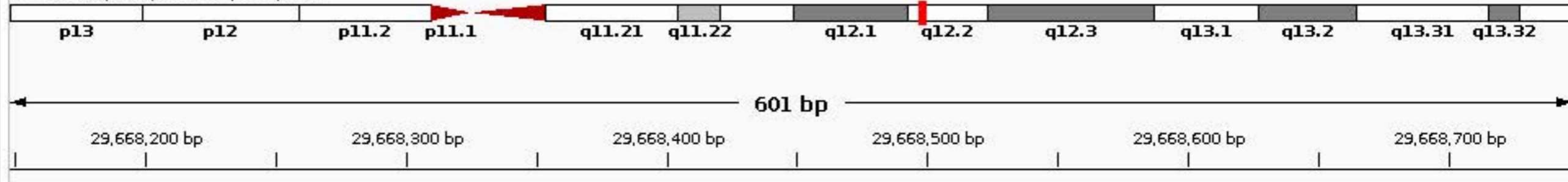
SL_0011-LT_11-B_mut.bam

Sequence

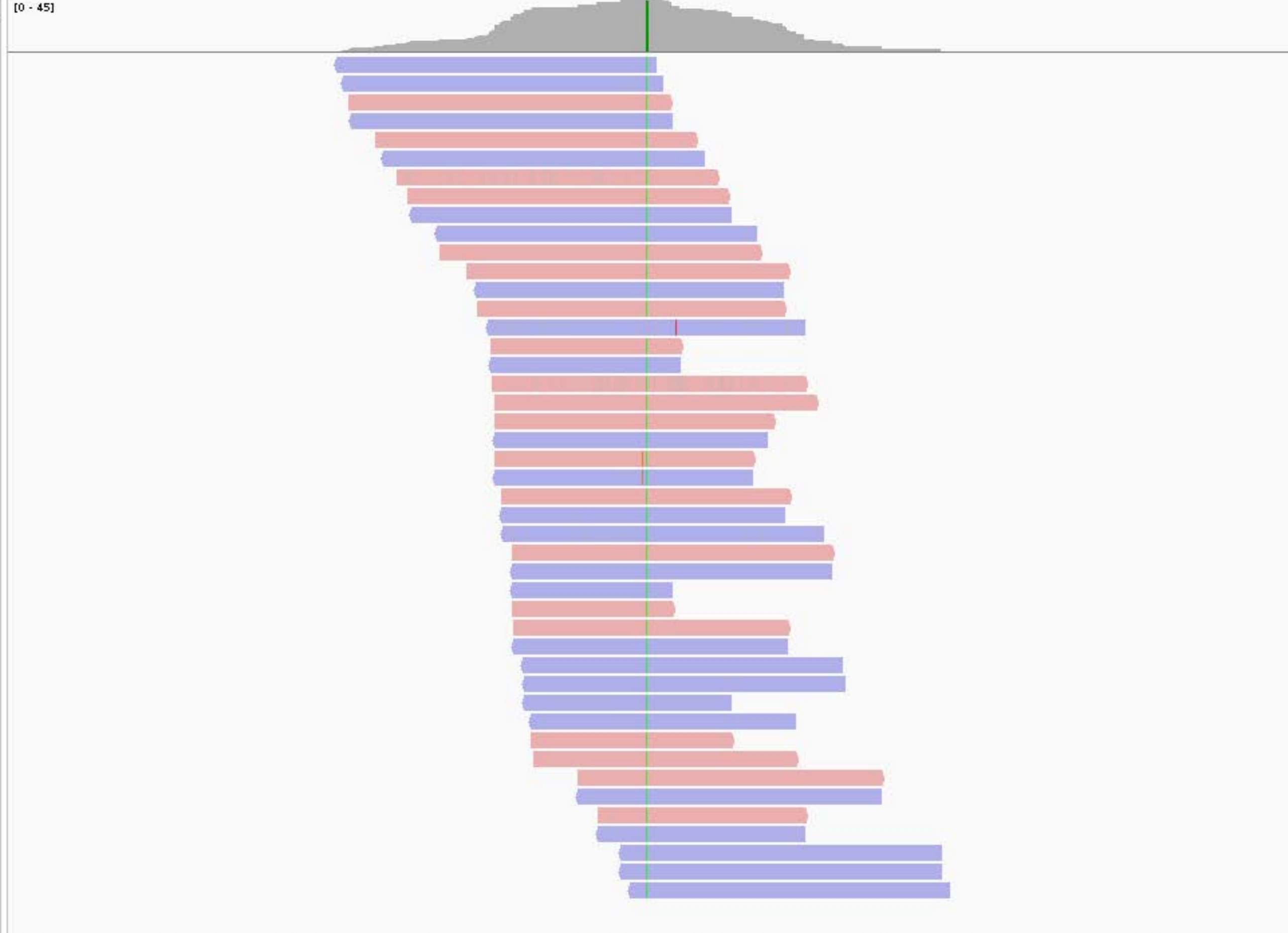
Refseq Genes



chr22:29,668,148-29,668,747



SL_1002_L_FFPE_24-B_mut.bam
rage



SL_1002_L_FFPE_24-B_mut.ba
m

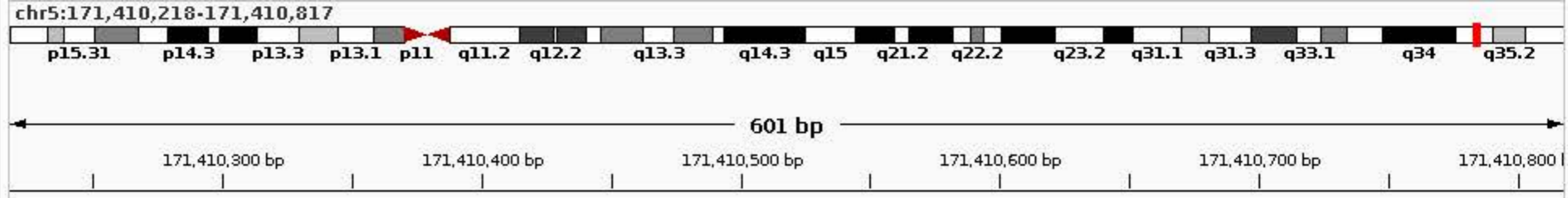
Sequence →



Refseq Genes



NF2

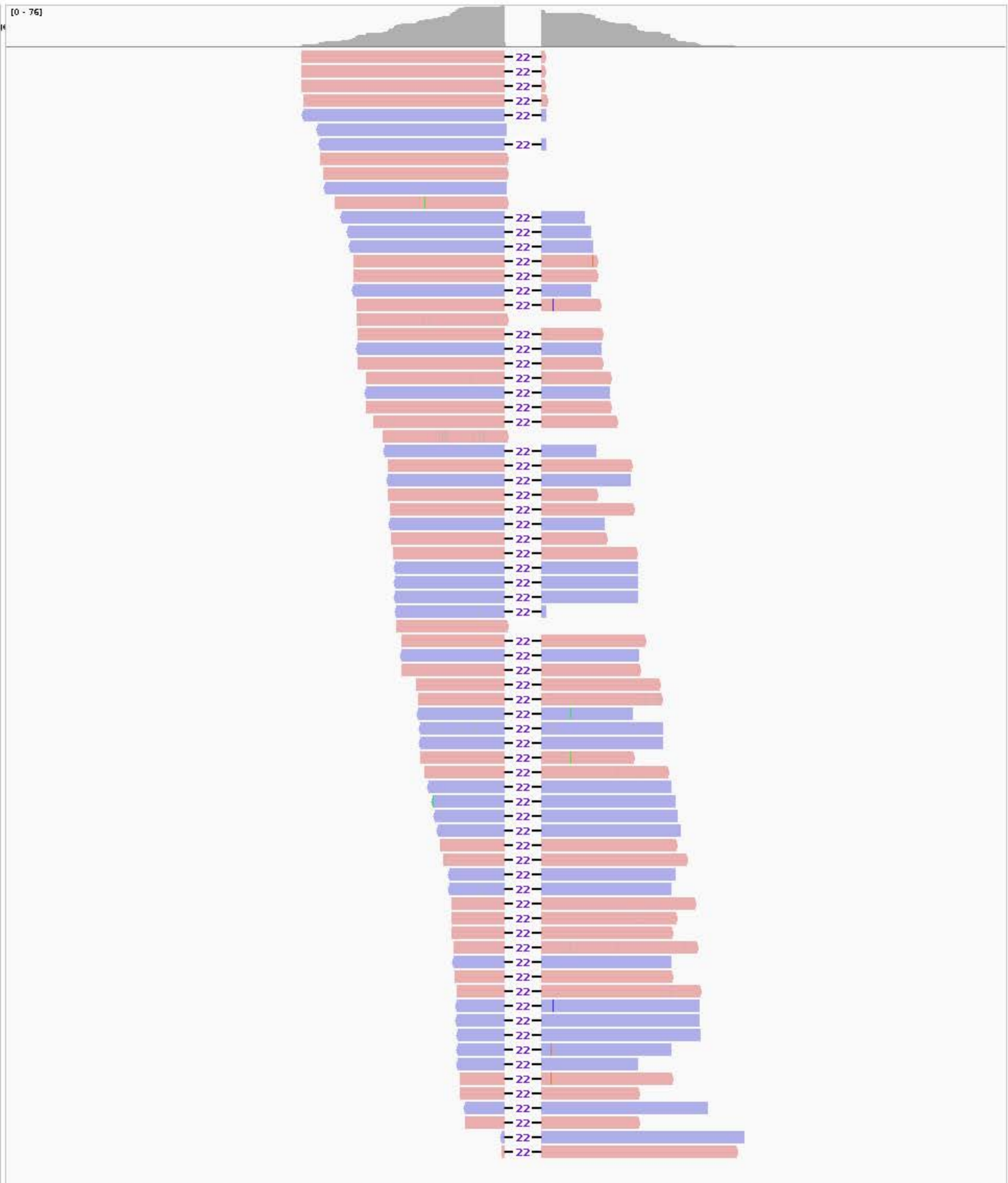


SL_1007-T1-B_mut.bam Coverage

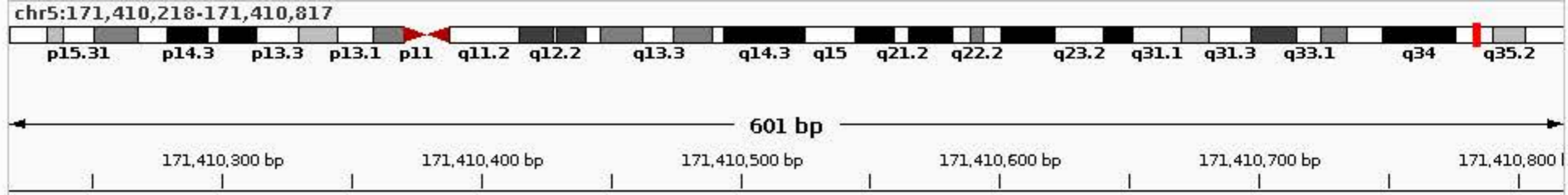
SL_1007-T1-B_mut.bam

Sequence

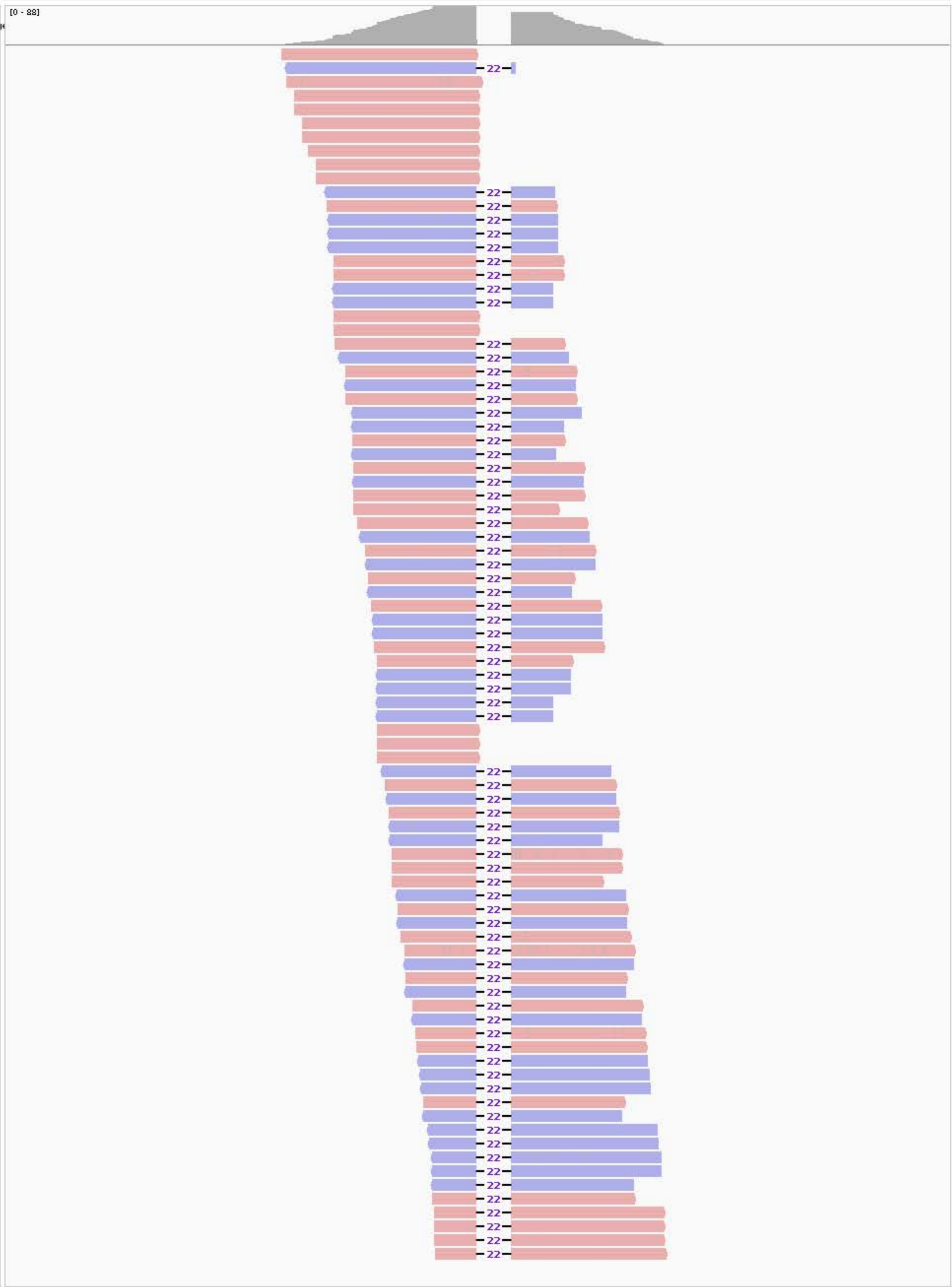
Refseq Genes



NPM1



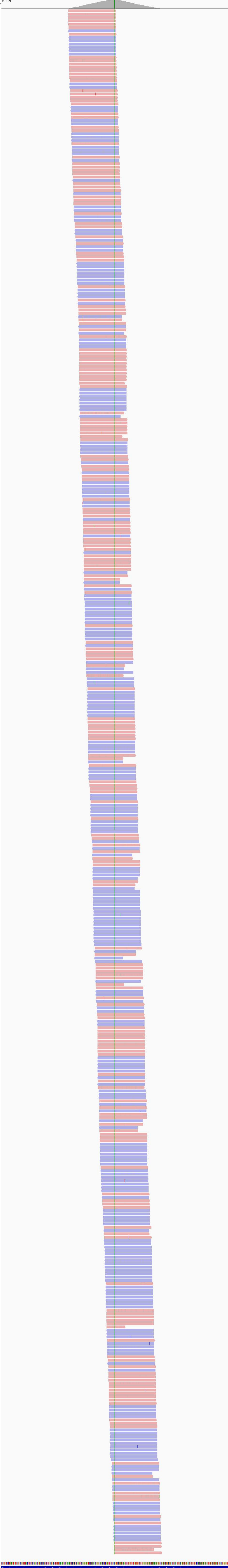
SL_1007-T2-B_mut.bam Coverage



Sequence →
Refseq Genes



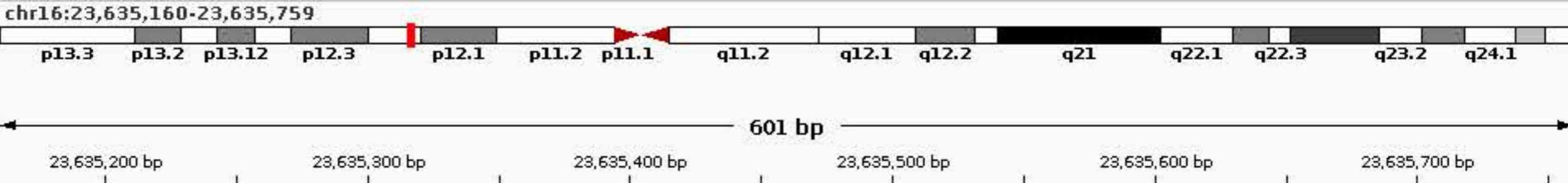
SL_0012_RT_3-B_mut.bam Cov



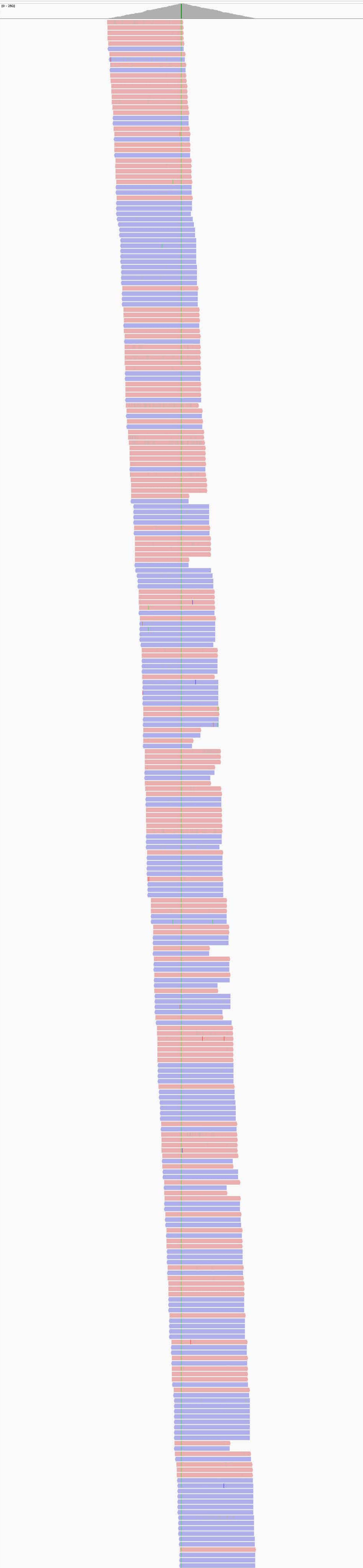
SL_0012_RT_3-B_mut.bam

Sequence
Refseq Genes





SL_0012_RT_4-B_mut.bam Coverage

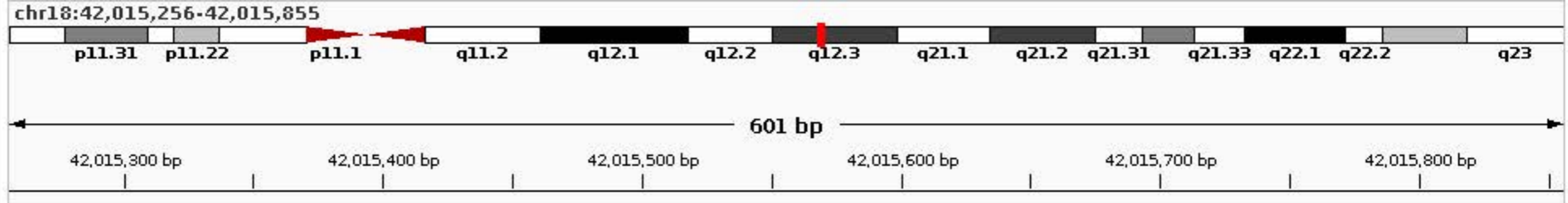


SL_0012_RT_4-B_mut.bam

Sequence



PALB2

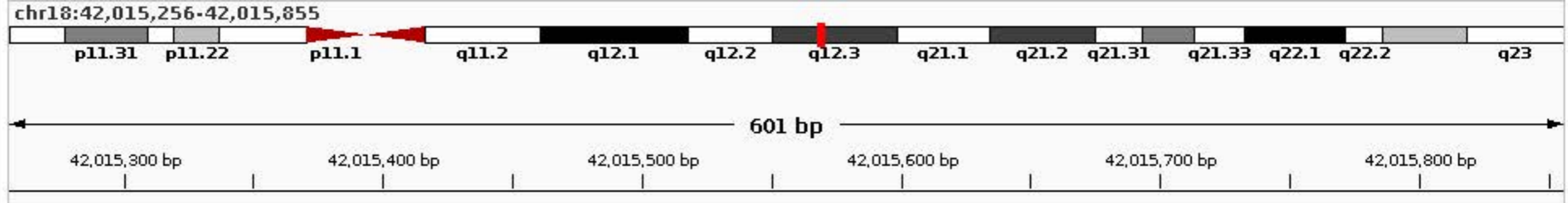


SL_1010_L_FFPE_3-B_mut.bam
age



Sequence →
Refseq Genes





SL_1010_L_FFPE_12-B_mut.barr
rage

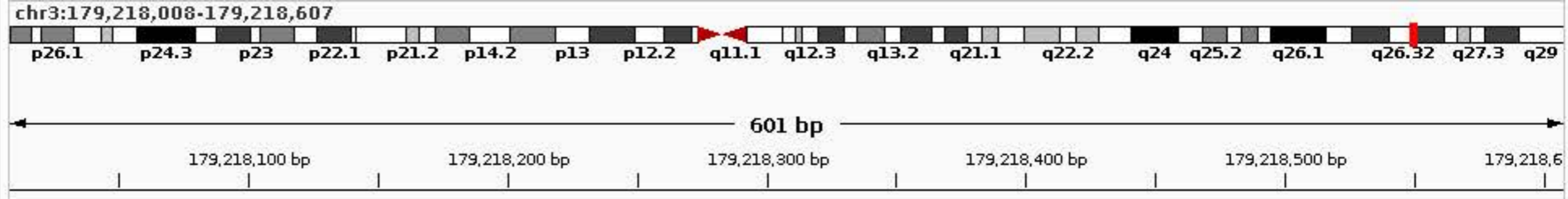
SL_1010_L_FFPE_12-B_mut.ba
m



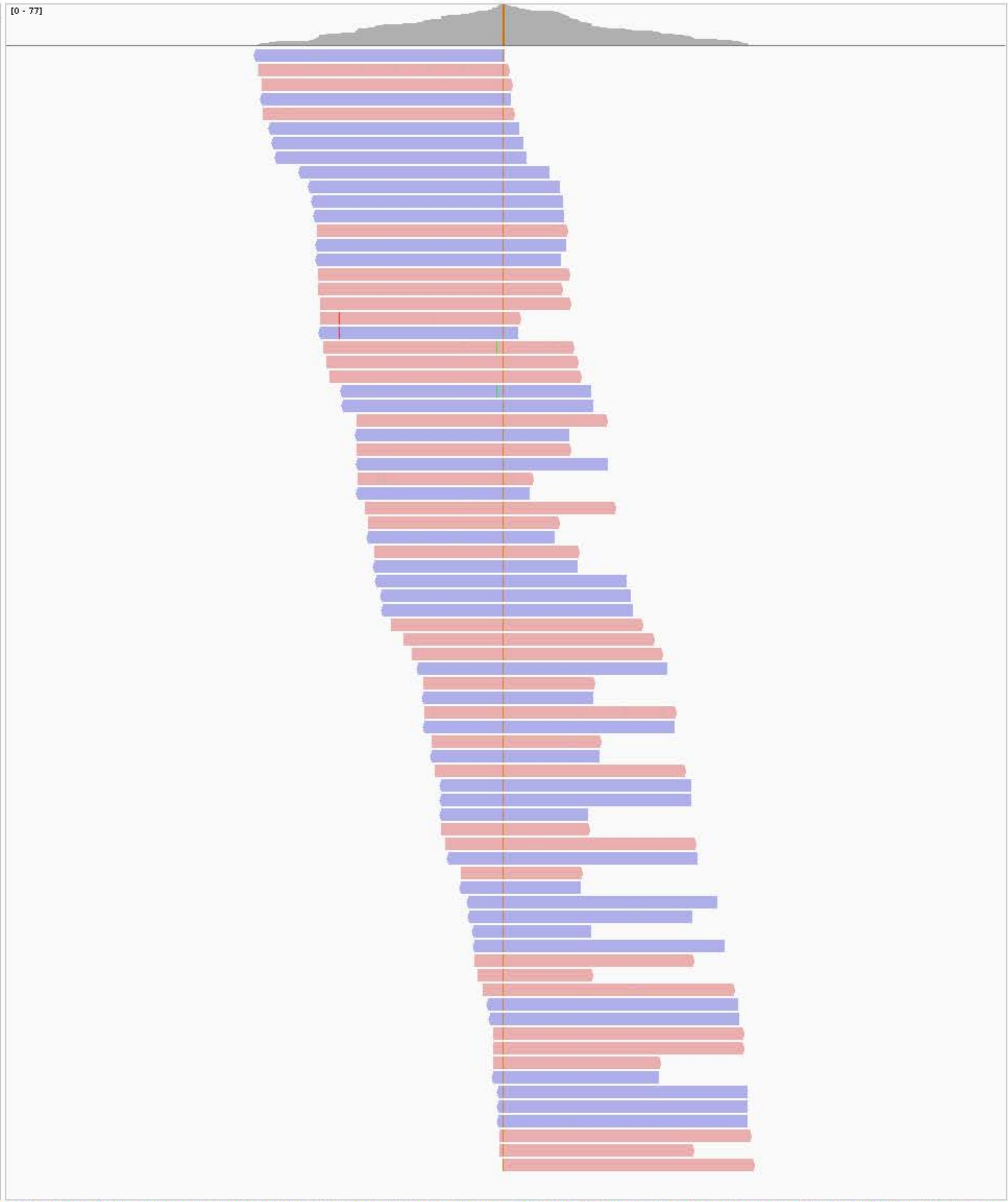
Sequence →

Refseq Genes





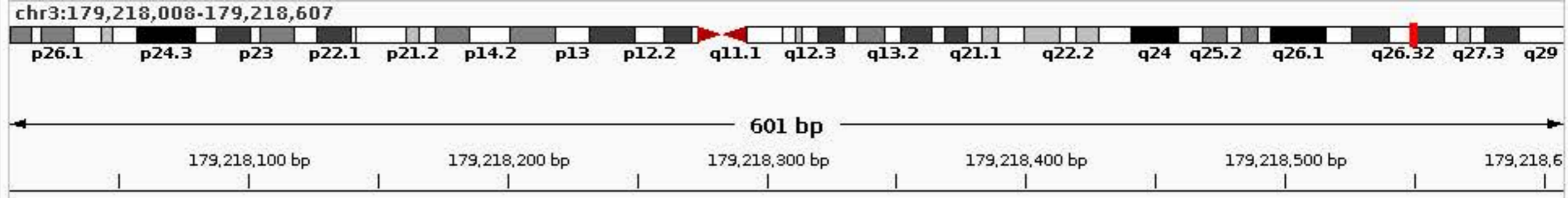
SL_0011_L_FFPE_20-B_mut.bam
rage



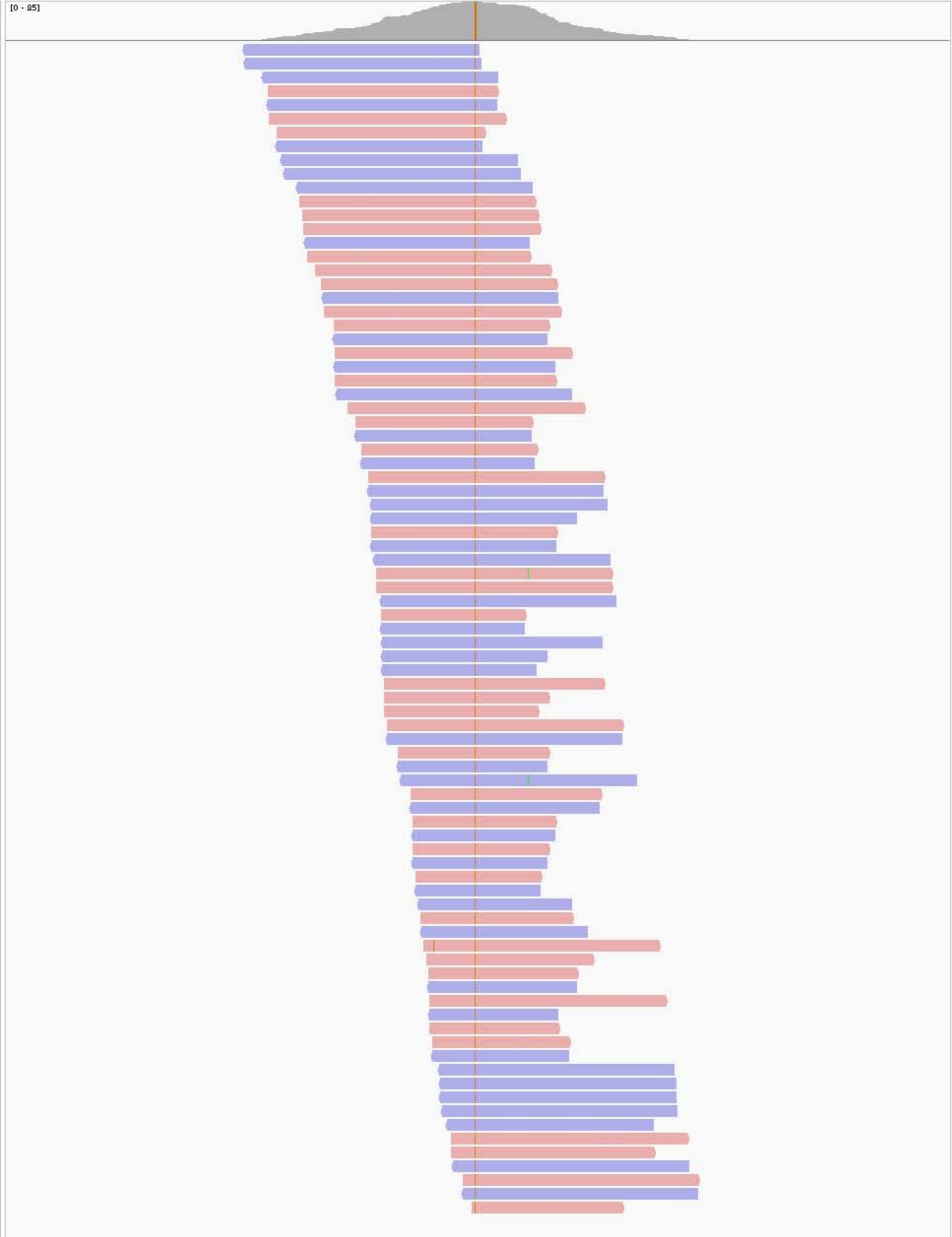
SL_0011_L_FFPE_20-B_mut.bam

Sequence →
Refseq Genes





SL_1002_L_FFPE_15-B_mut.bam
rage

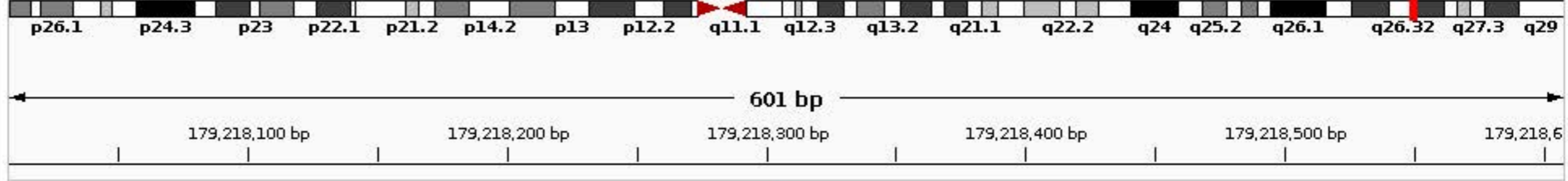


SL_1002_L_FFPE_15-B_mut.ba
m

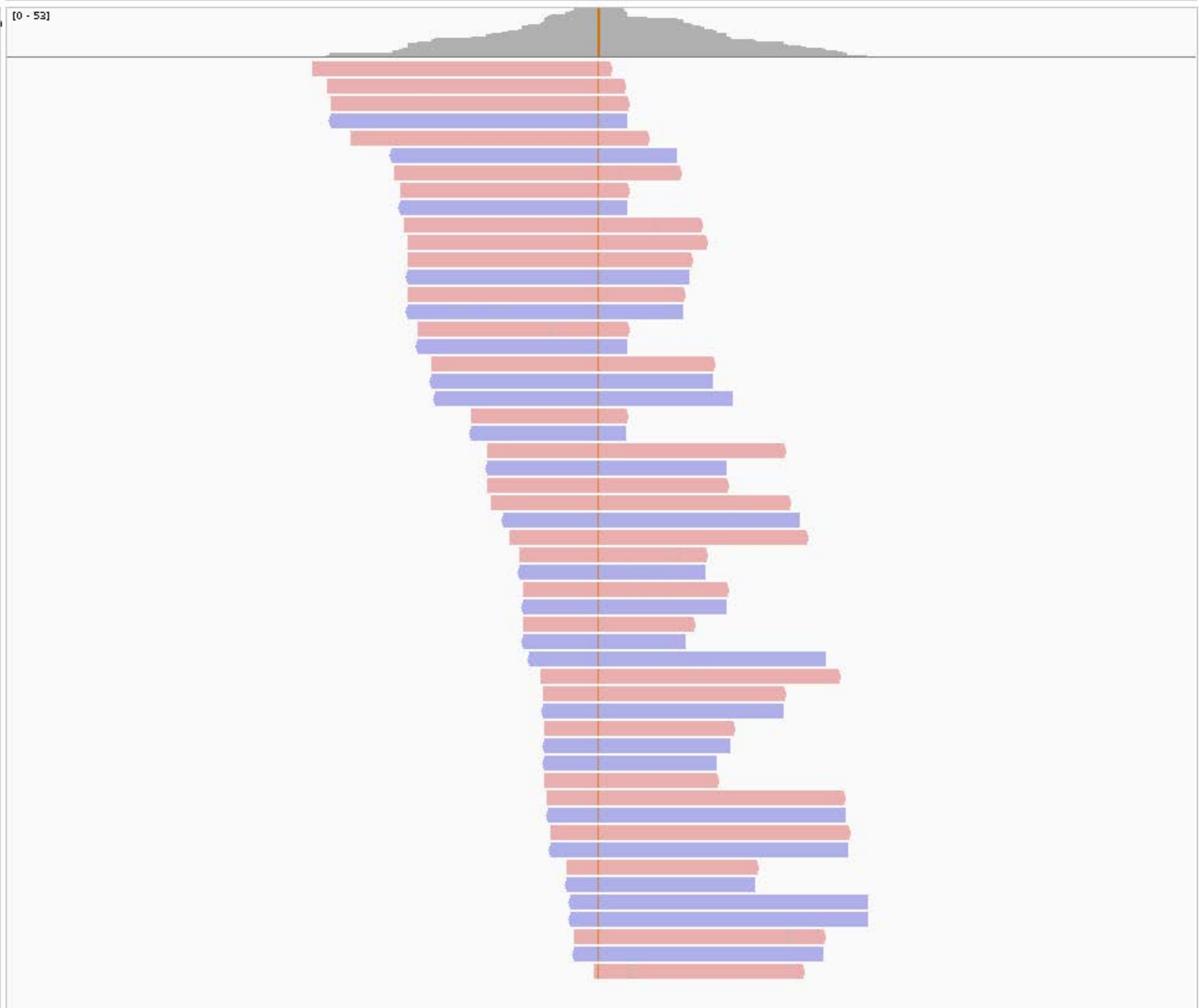
Sequence →
Refseq Genes



chr3:179,218,008-179,218,607



SL_1003_R_FFPE_93-B_mut.bam
rage



SL_1003_R_FFPE_93-B_mut.ba
m

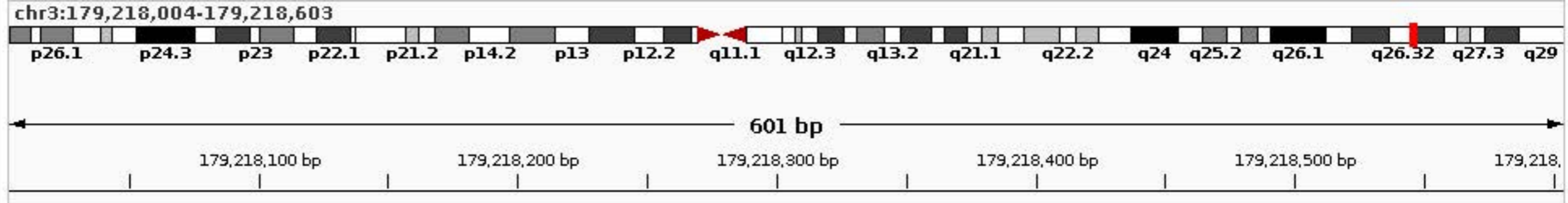
Sequence →



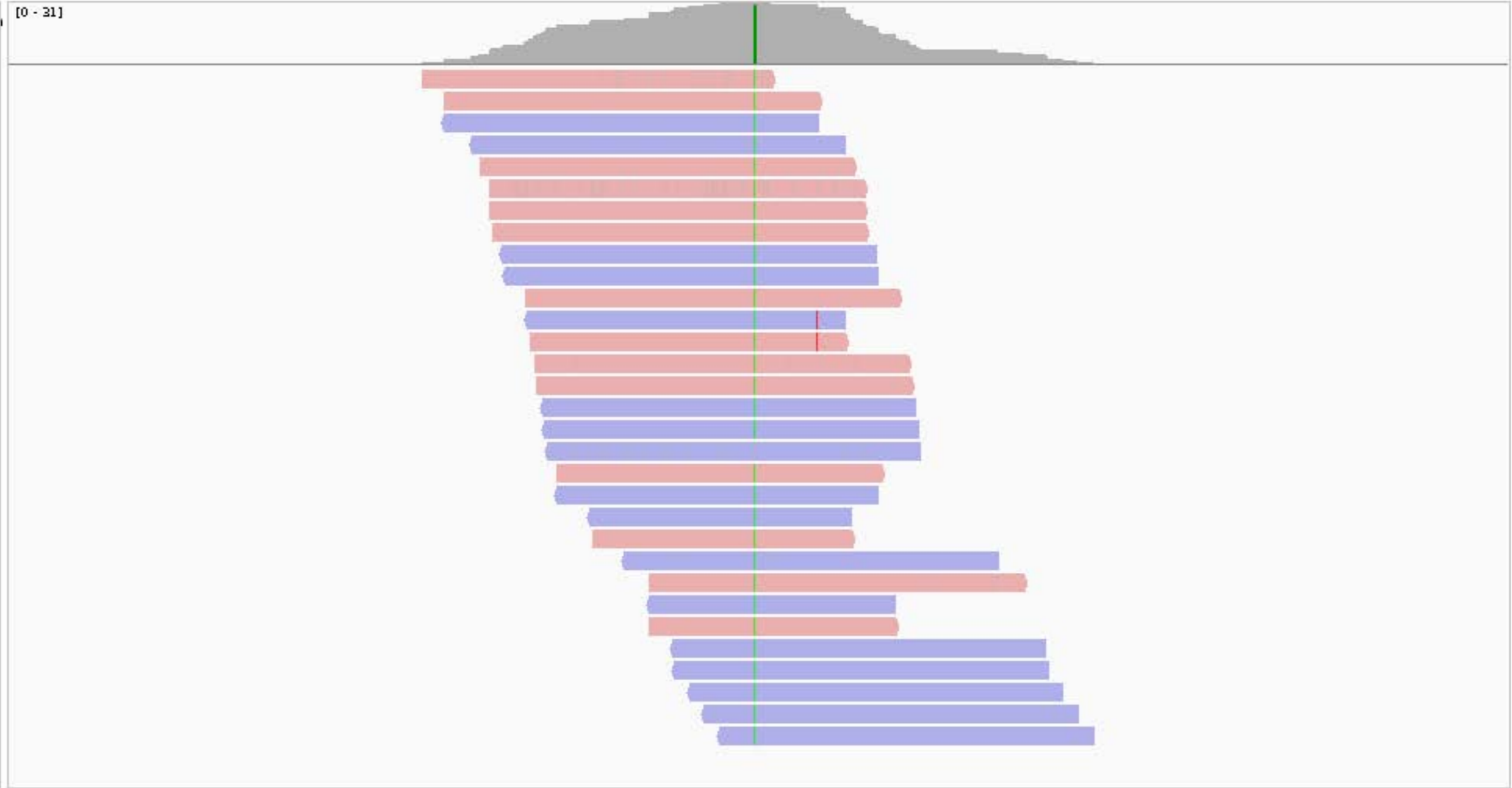
Refseq Genes



PIK3CA



SL_0005_R_FFPE_73-B_mut.bam
rage



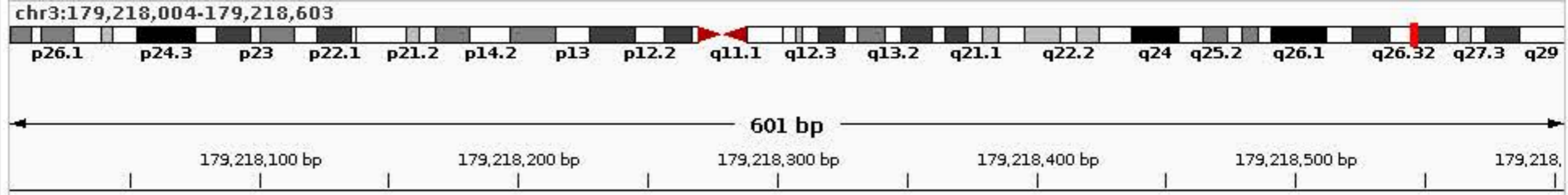
SL_0005_R_FFPE_73-B_mut.ba
m

Sequence →



Refseq Genes



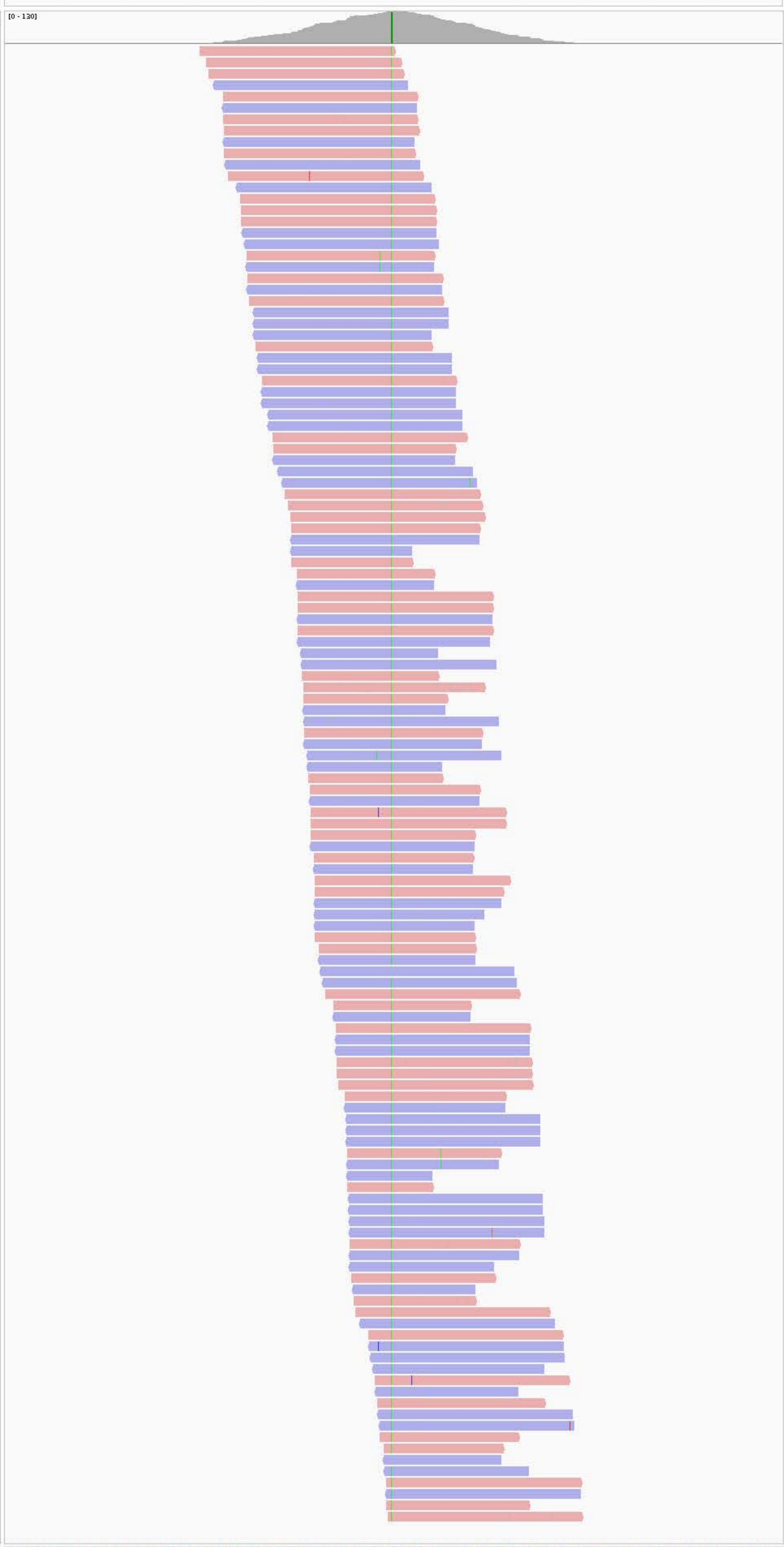


SL_1004_L_FFPE_38-B_mut.barrage

SL_1004_L_FFPE_38-B_mut.bam

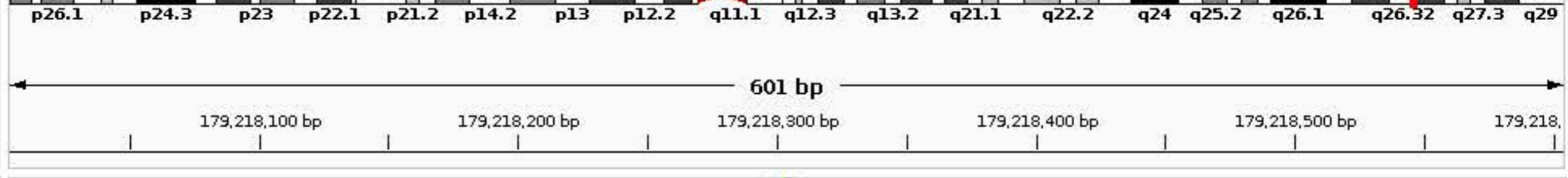
Sequence →

Refseq Genes



PIK3CA

chr3:179,218,004-179,218,603

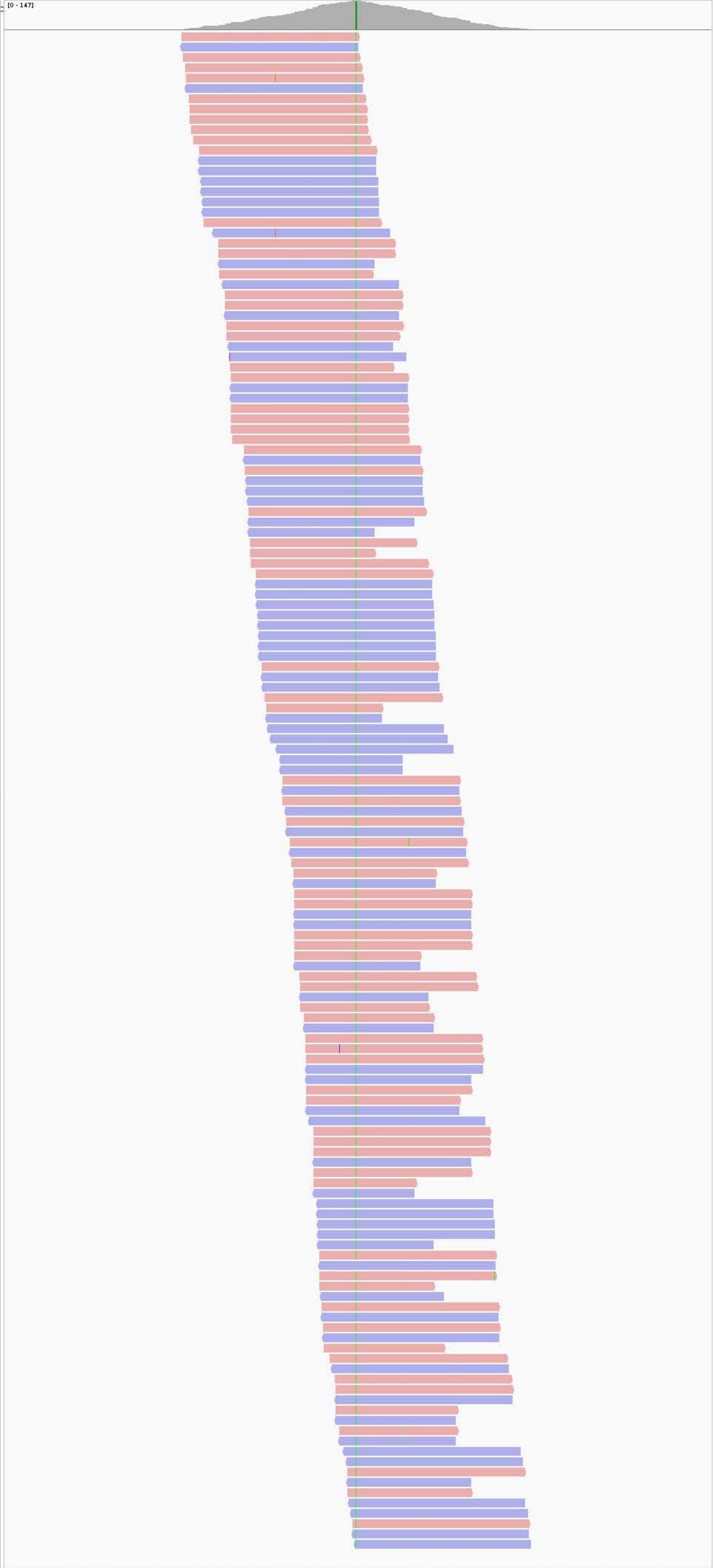


SL_1008_L_FFPE_3-B_mut.bam
age

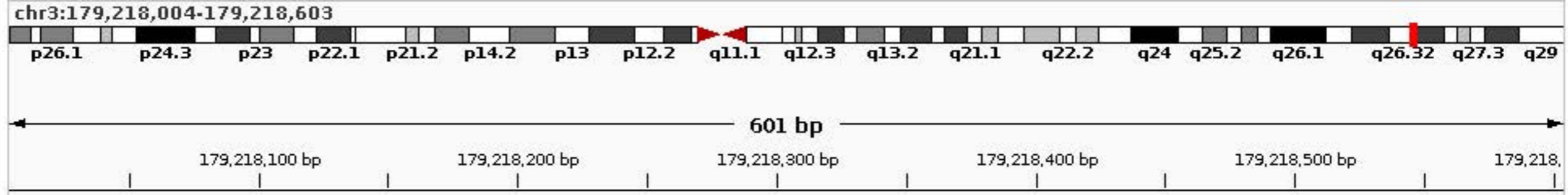
SL_1008_L_FFPE_3-B_mut.bam

Sequence →

Refseq Genes

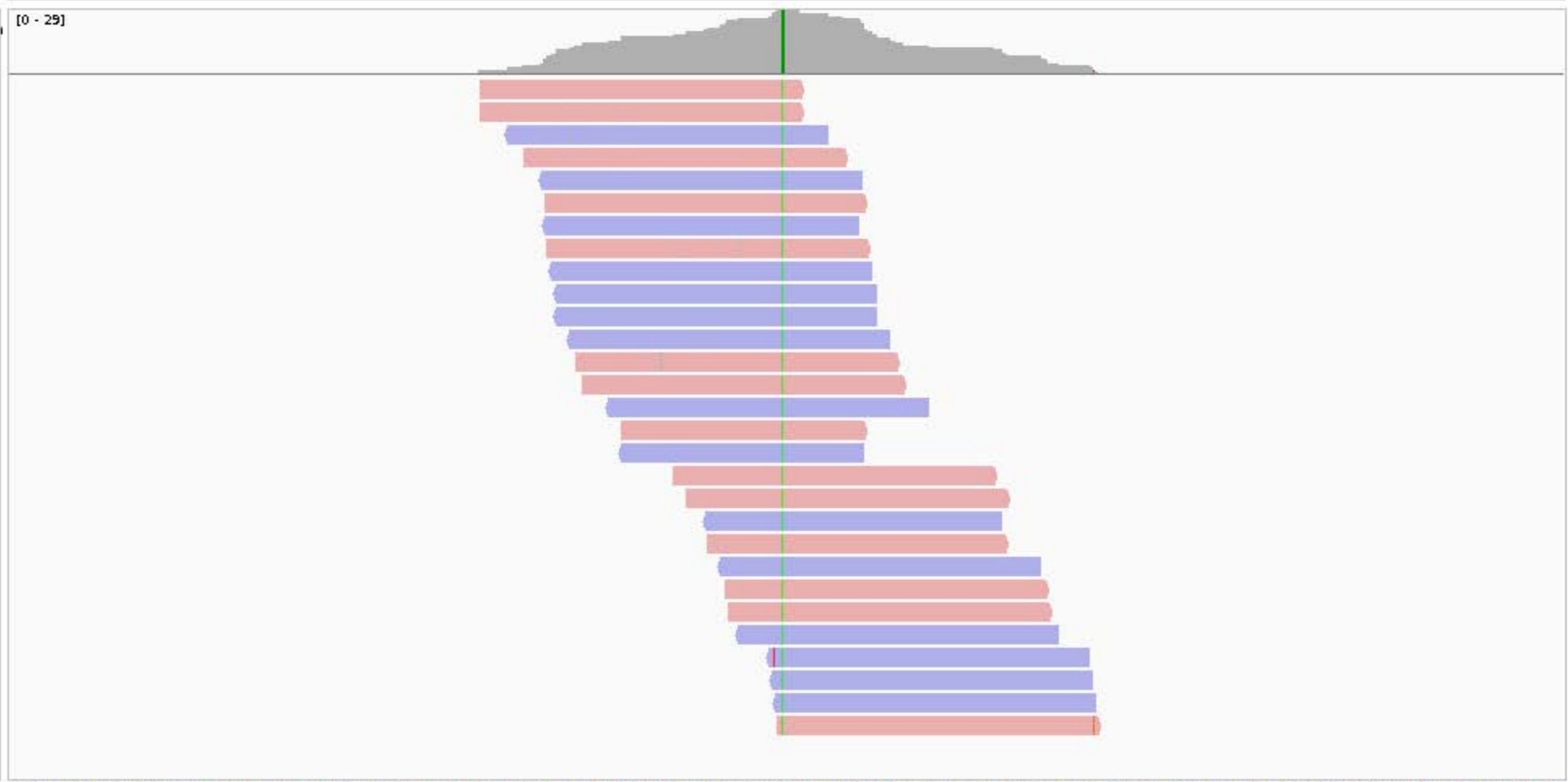


PIK3CA



SL_1012_R_10_FFPE-B_mut.bam
rage

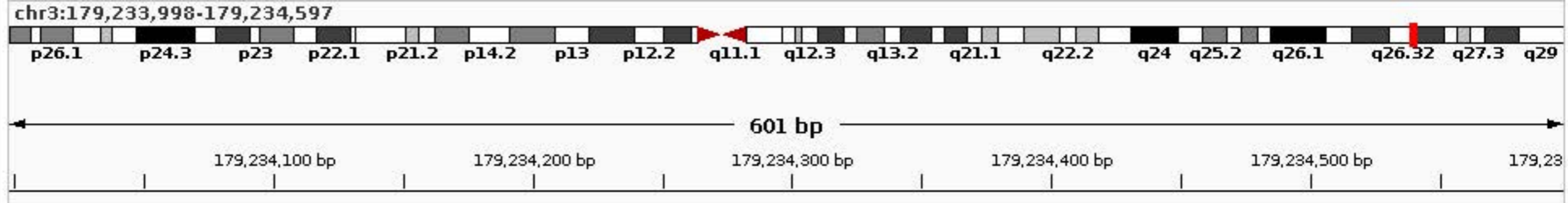
SL_1012_R_10_FFPE-B_mut.ba
m



Sequence →

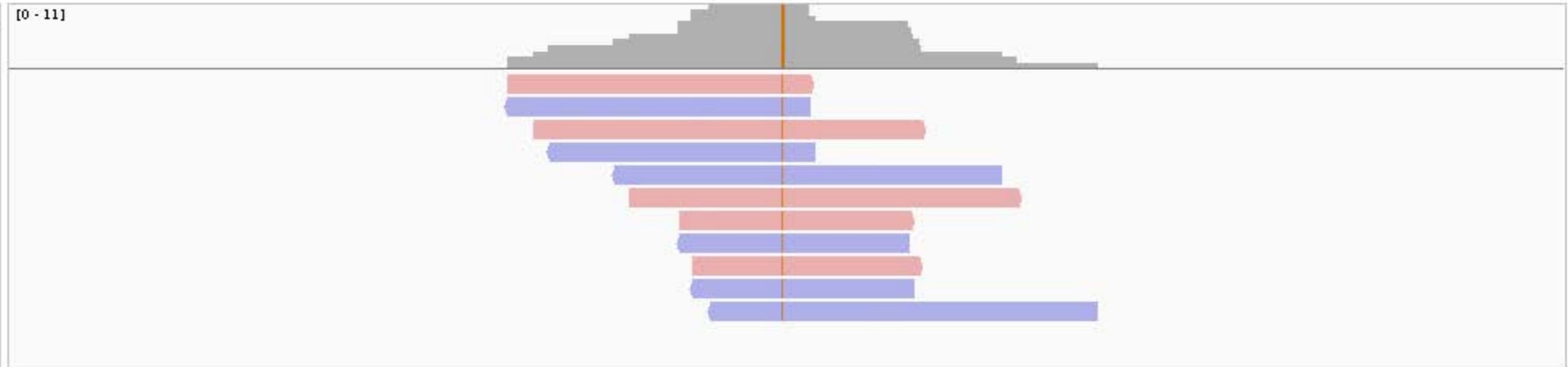
Refseq Genes





SL_0001_L_FFPE_10-B_mut.barr
rage

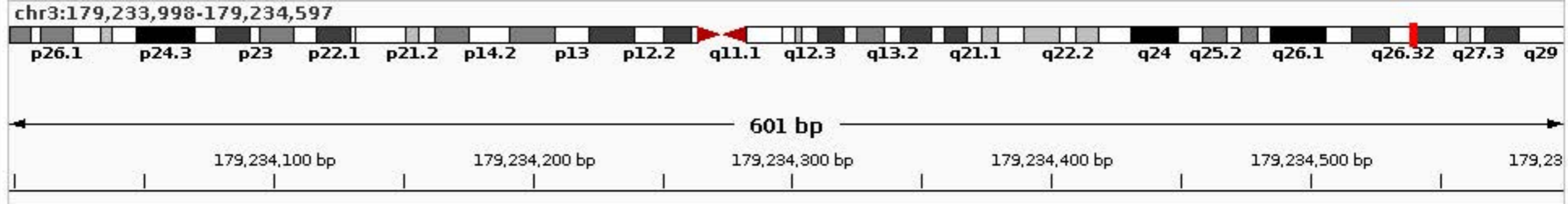
SL_0001_L_FFPE_10-B_mut.ba
m



Sequence →

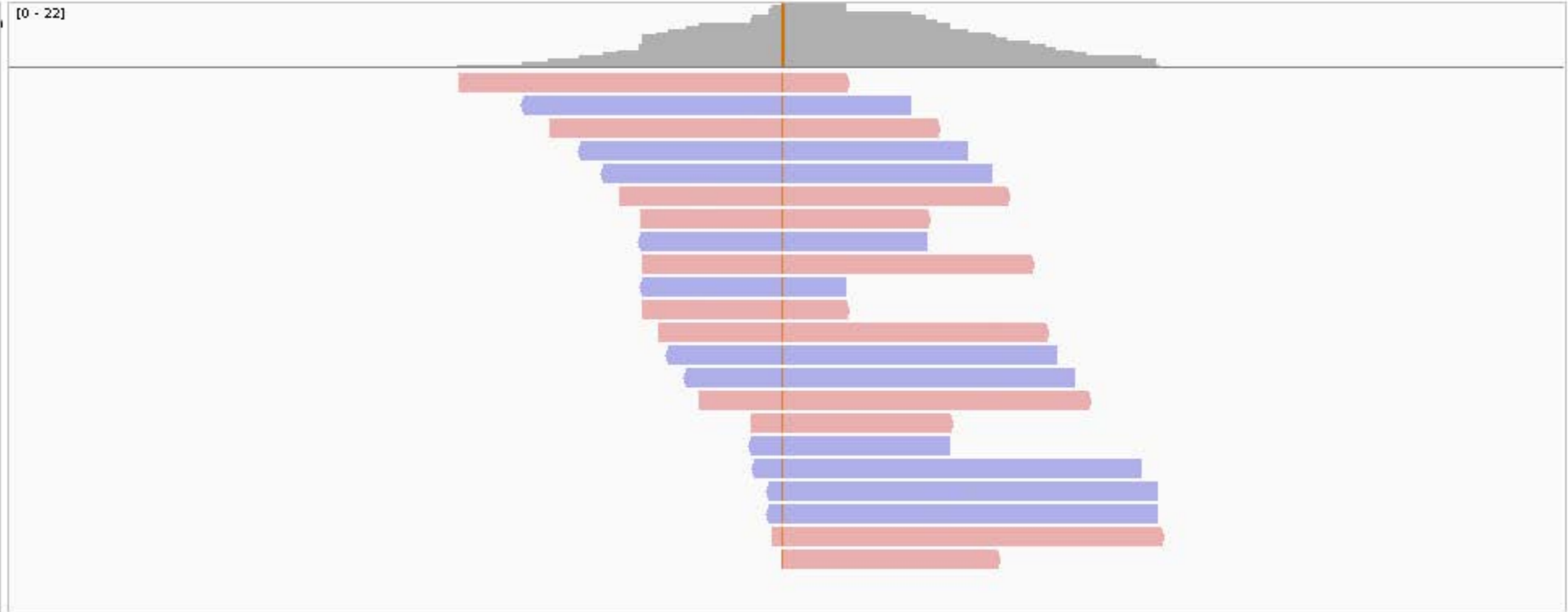
Refseq Genes





SL_0001_R_FFPE_58-B_mut.bam
rage

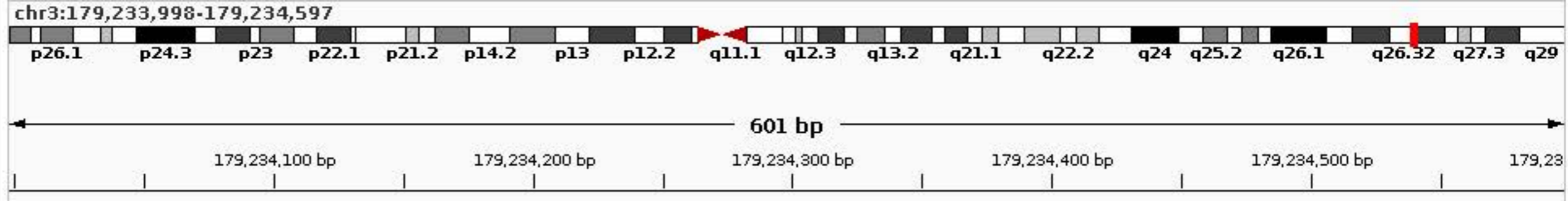
SL_0001_R_FFPE_58-B_mut.ba
m



Sequence →

Refseq Genes





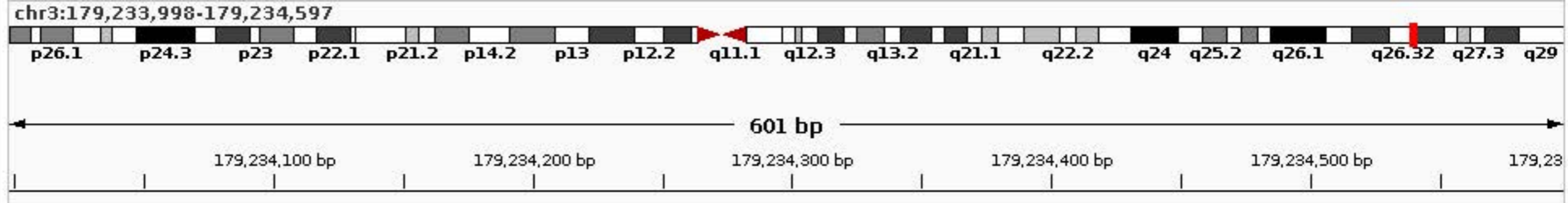
SL_0004_L_FFPE_14-B_mut.barrage



SL_0004_L_FFPE_14-B_mut.bam

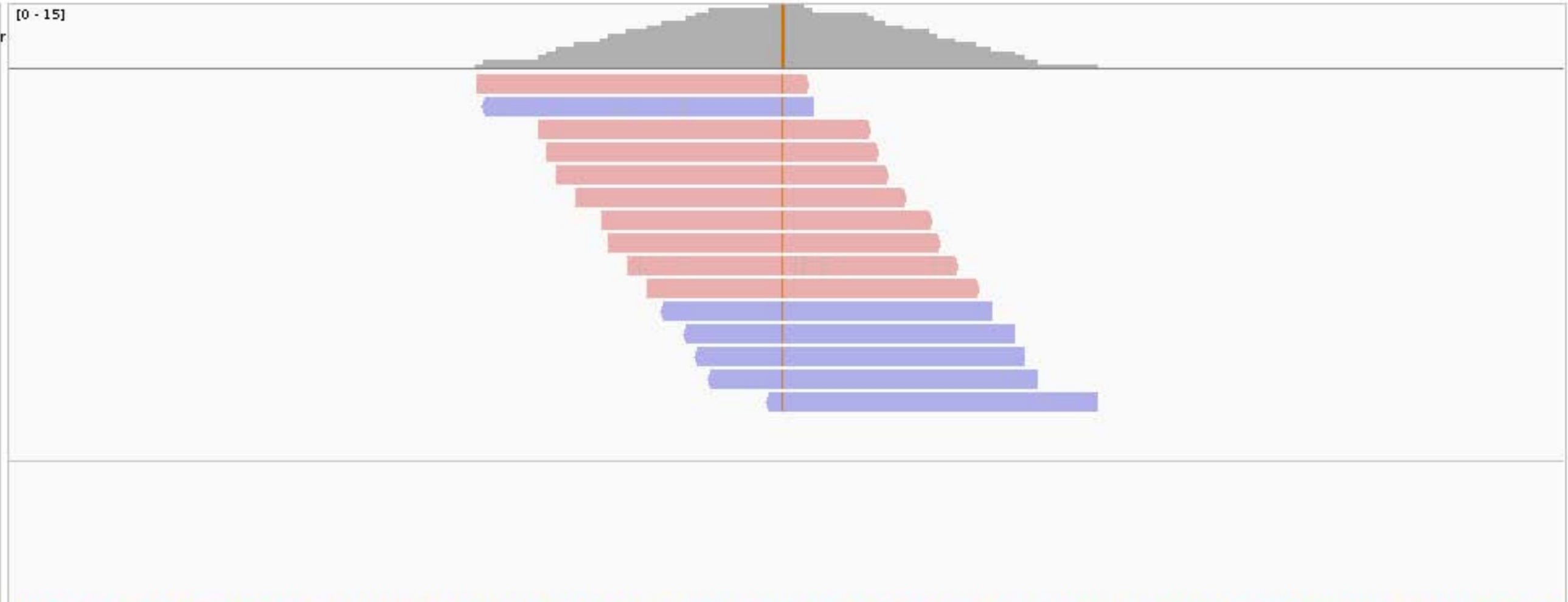
Sequence →
Refseq Genes





SL_0004-RN-8-B_mut.bam Cover

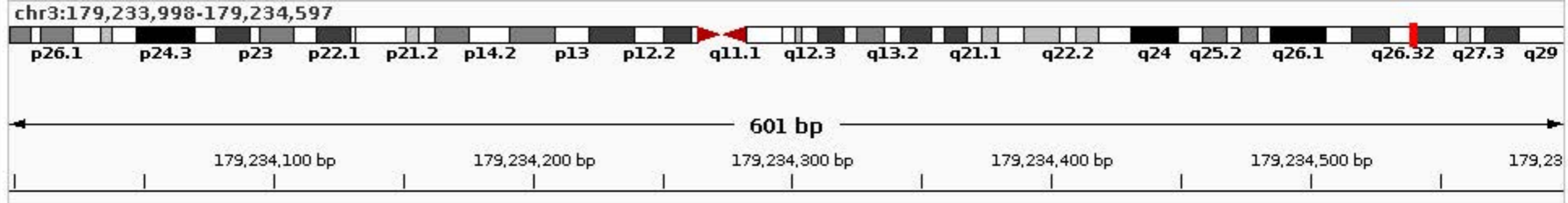
SL_0004-RN-8-B_mut.bam



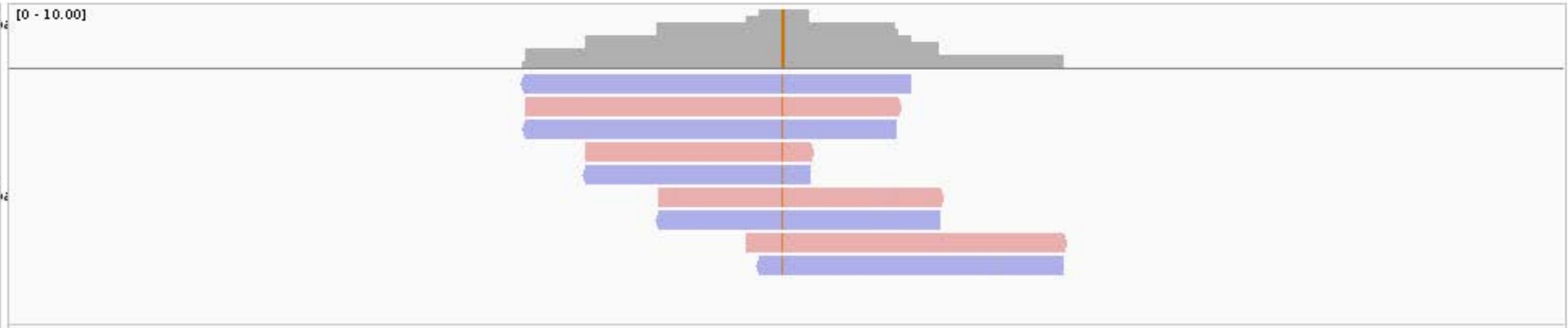
Sequence →

Refseq Genes



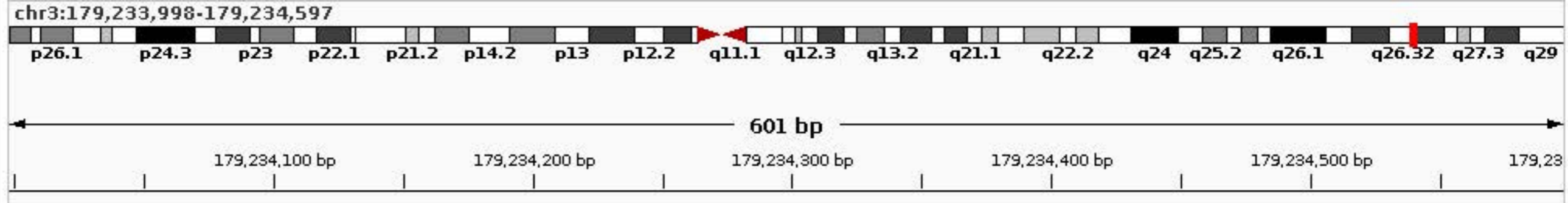


SL_0005_L_FFPE_6_skin-B_mut.hi
Coverage



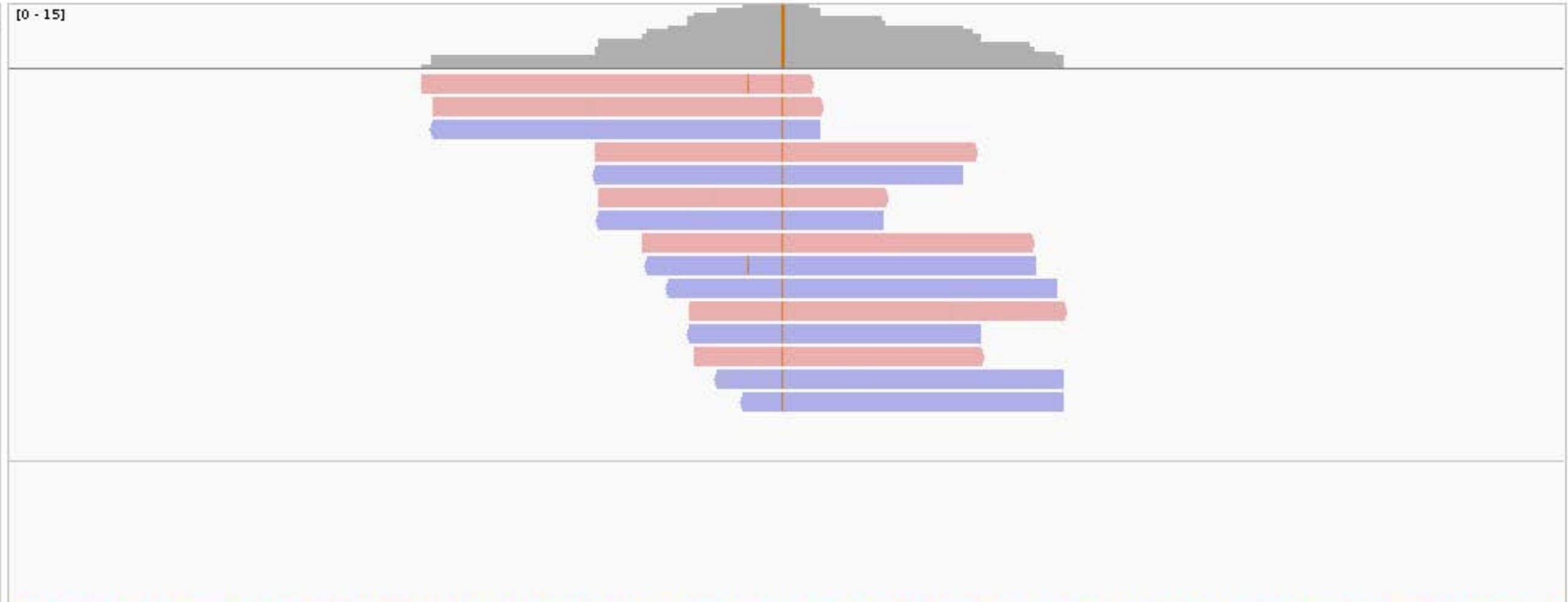
Sequence →
Refseq Genes





SL_0005_L_FFPE_38-B_mut.bam
rage

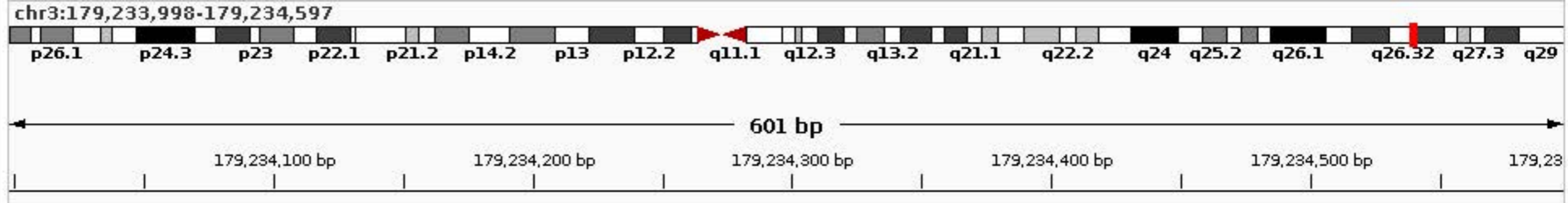
SL_0005_L_FFPE_38-B_mut.ba
m



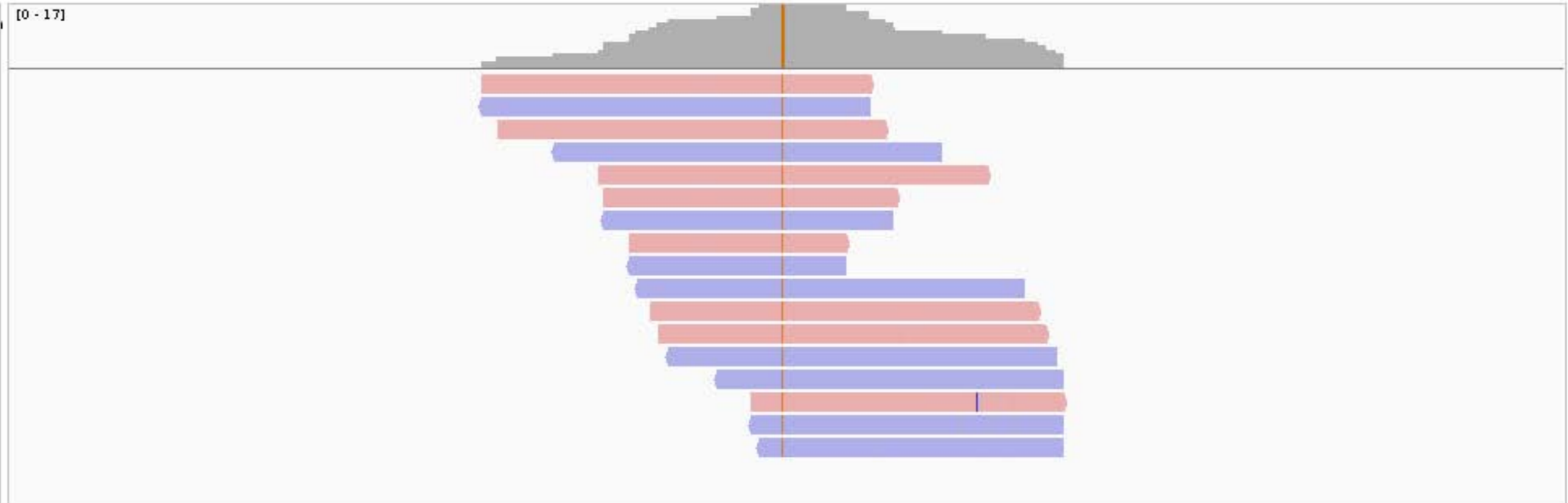
Sequence →

Refseq Genes





SL_0005_R_FFPE_78-B_mut.bam
range



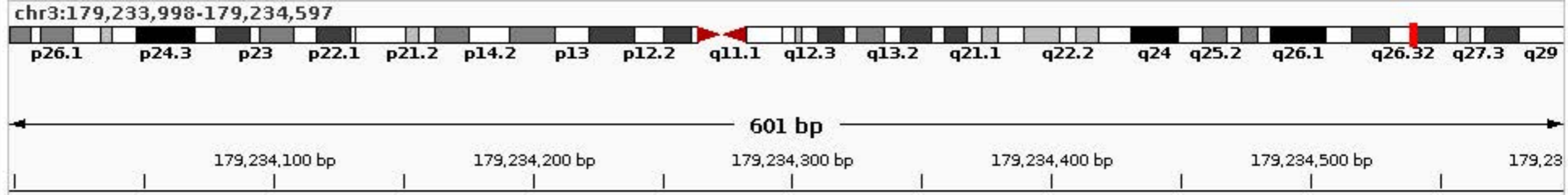
SL_0005_R_FFPE_78-B_mut.bam

Sequence →

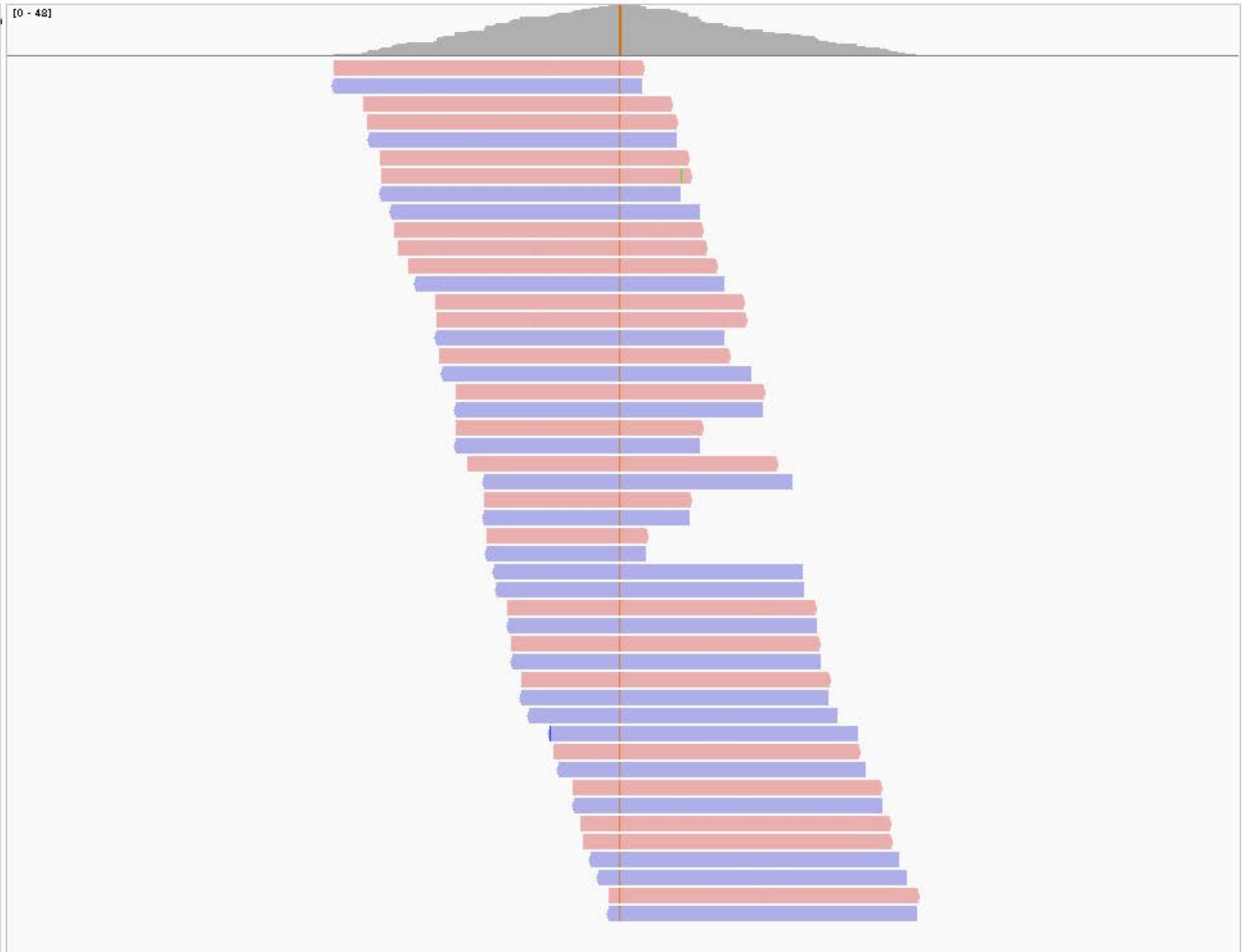


Refseq Genes



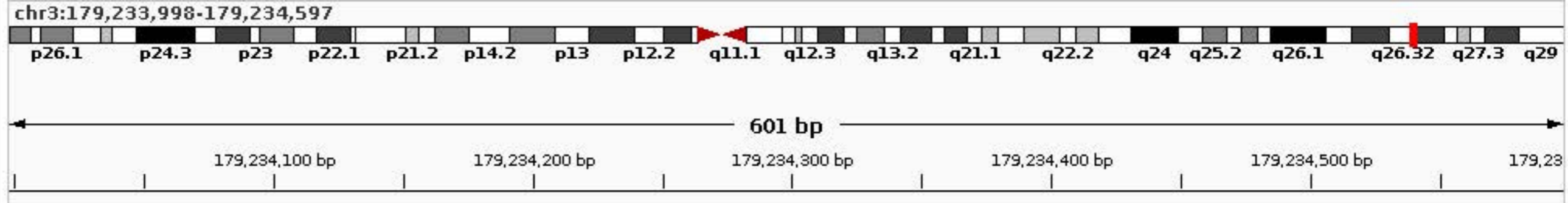


SL_0006_R_FFPE_21-B_mut.bam
rage



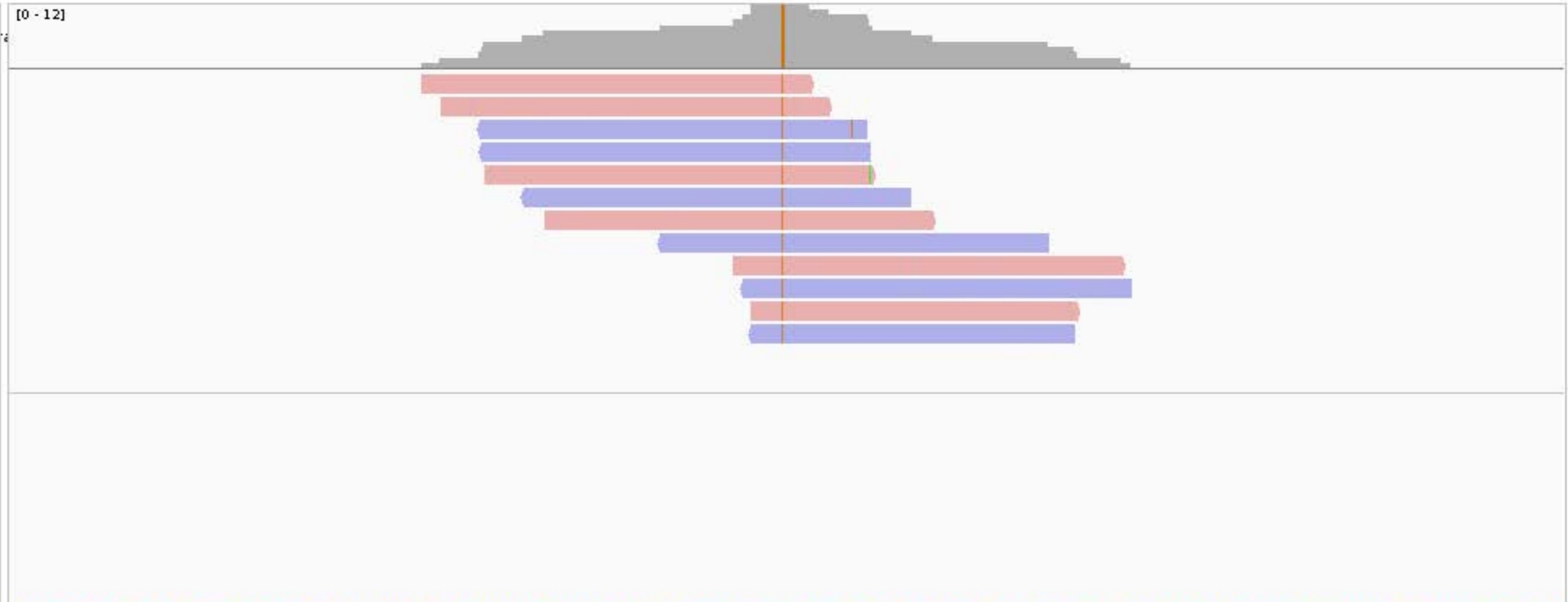
Sequence →
Refseq Genes





SL_0007_LT_9-B_mut.bam Coverage

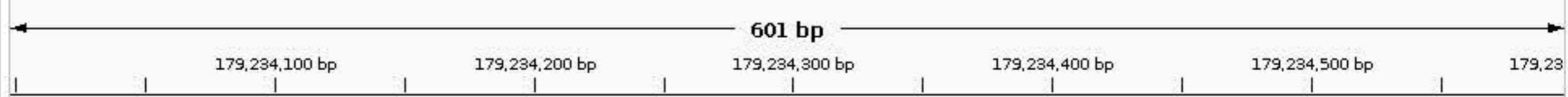
SL_0007_LT_9-B_mut.bam



Sequence →

Refseq Genes



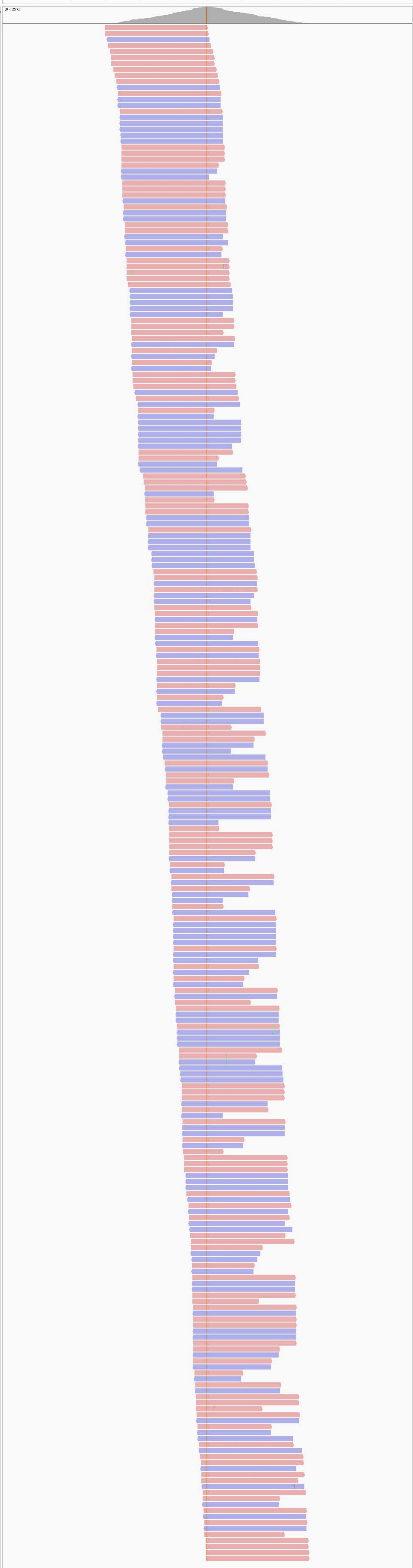


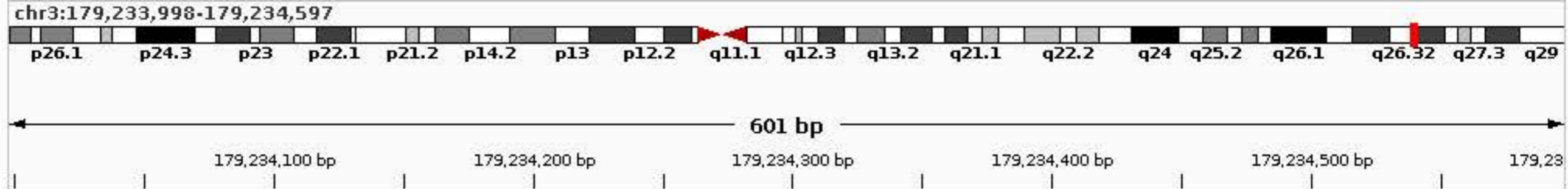
SL_0007_R_FFPE_67-B_mut.bam
rage

SL_0007_R_FFPE_67-B_mut.ba
m

Sequence →

Refseq Genes



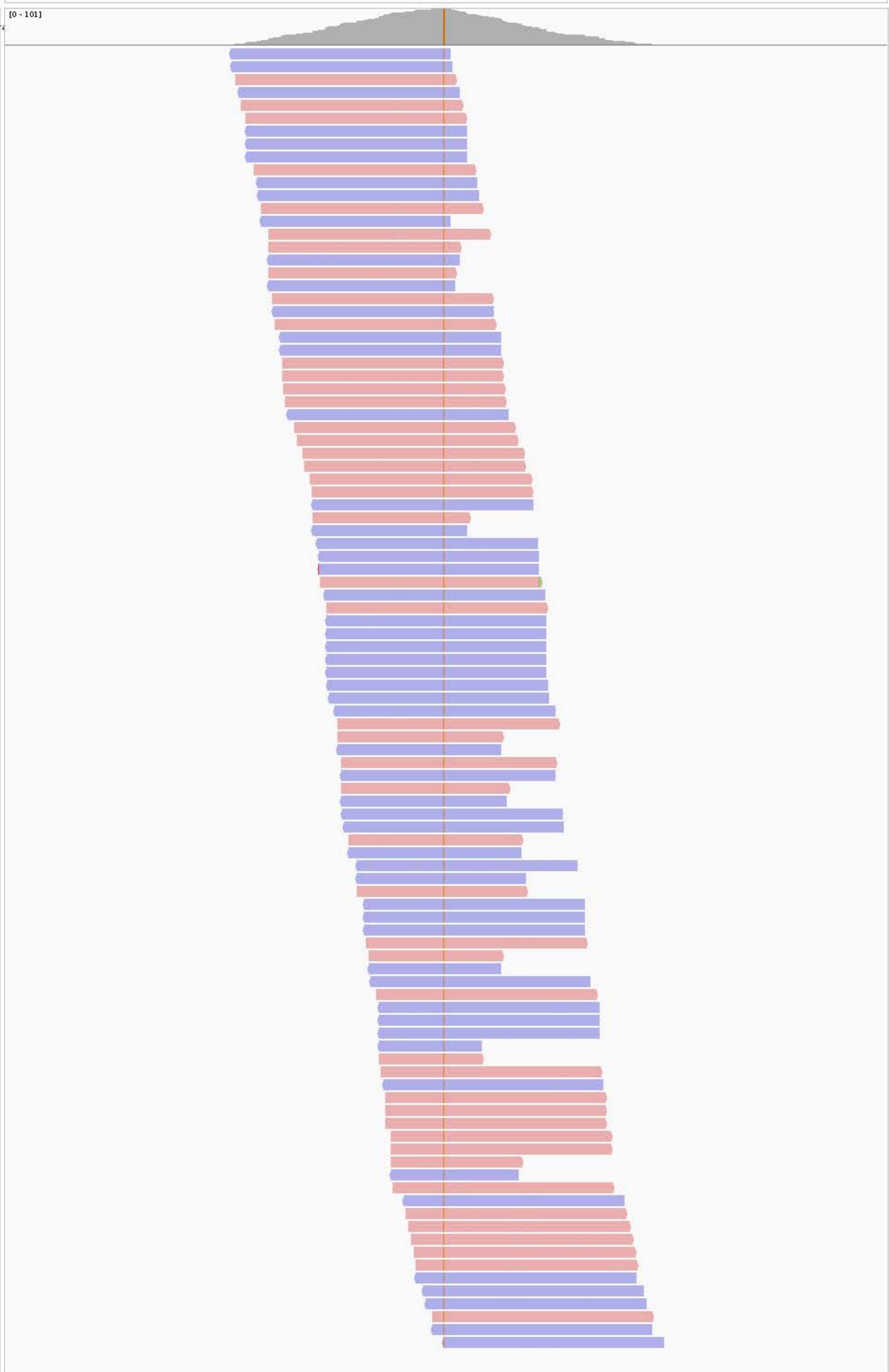


SL_0008_LT_8-B_mut.bam Coverage

SL_0008_LT_8-B_mut.bam

Sequence

Refseq Genes

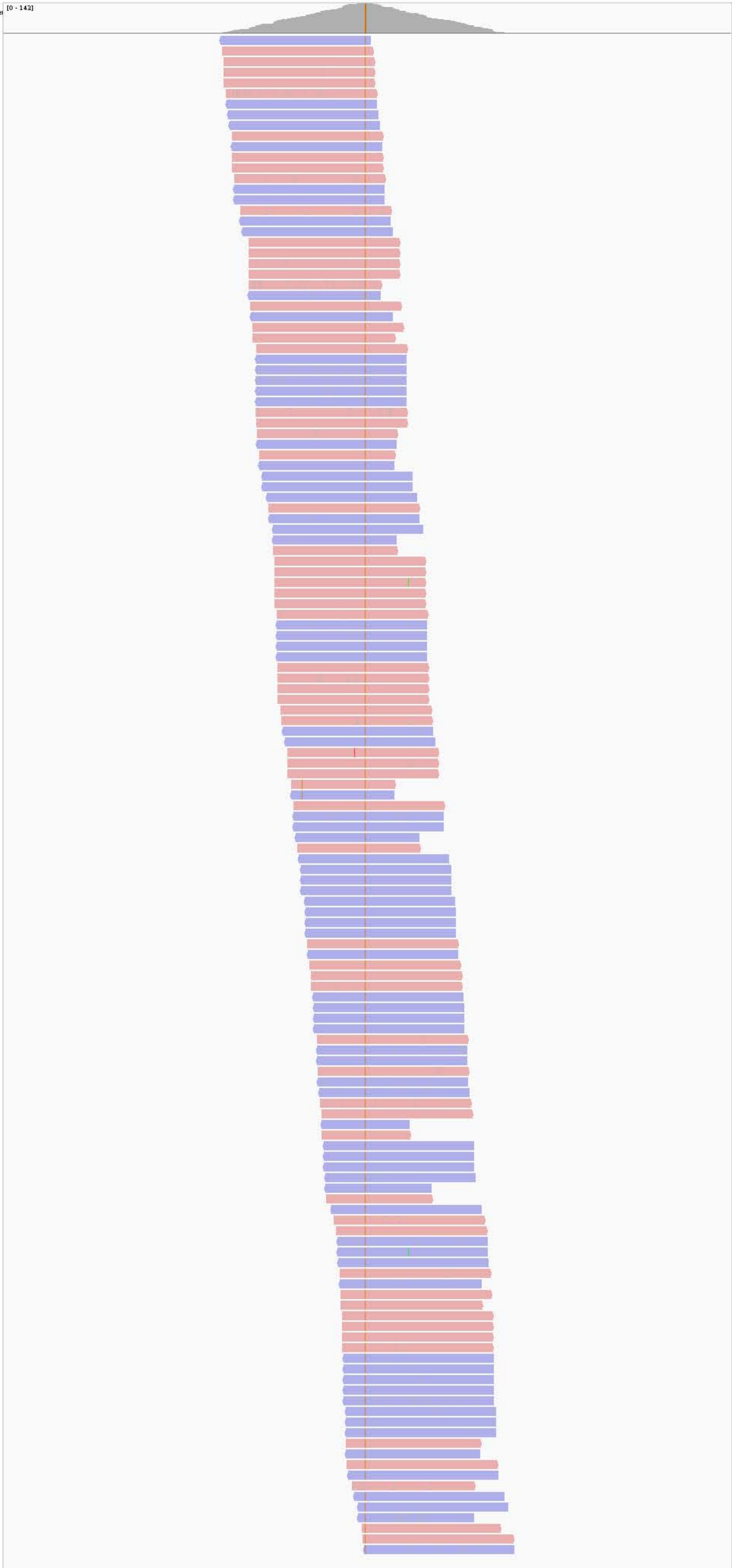


chr3:179,233,998-179,234,597

p26.1 p24.3 p23 p22.1 p21.2 p14.2 p13 p12.2 q11.1 q12.3 q13.2 q21.1 q22.2 q24 q25.2 q26.1 q26.32 q27.3 q29



SL_0008-LT_15-B_mut.bam Coverage



SL_0008-LT_15-B_mut.bam

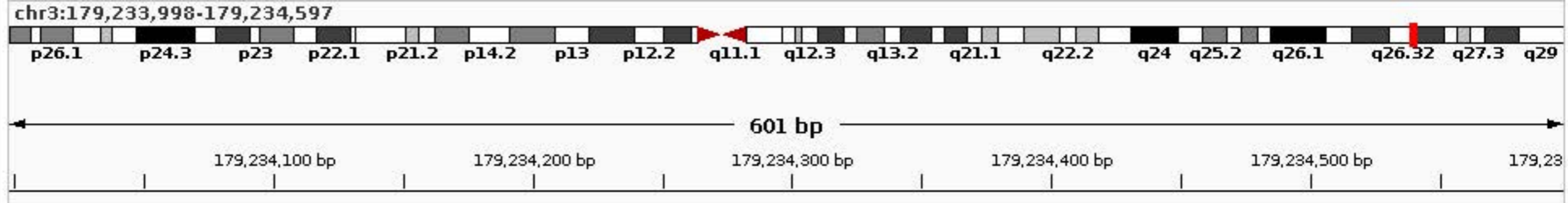
Sequence



Refseq Genes

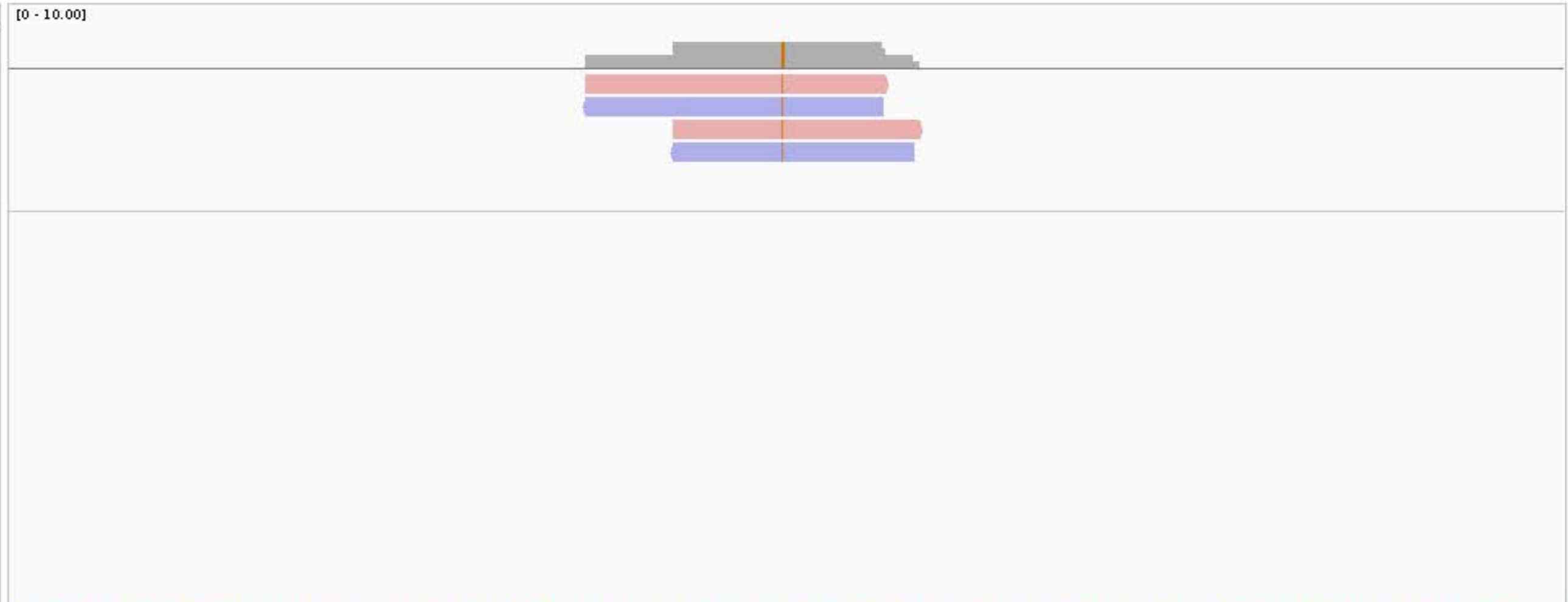


PIK3CA



SL_0009_L_FFPE_21-B_mut.bam
range

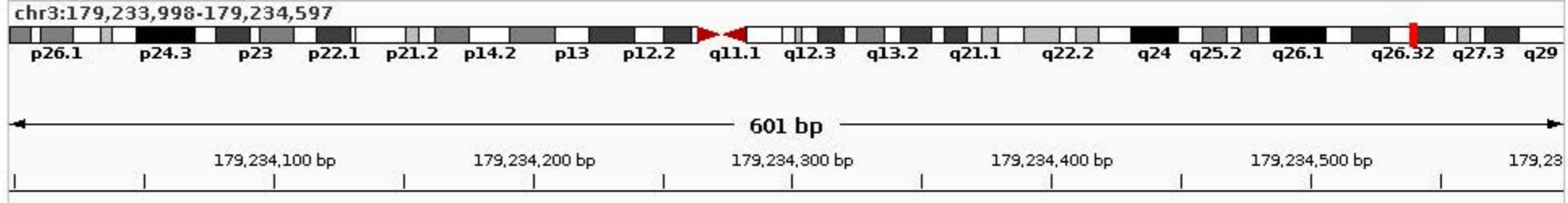
SL_0009_L_FFPE_21-B_mut.bam



Sequence →

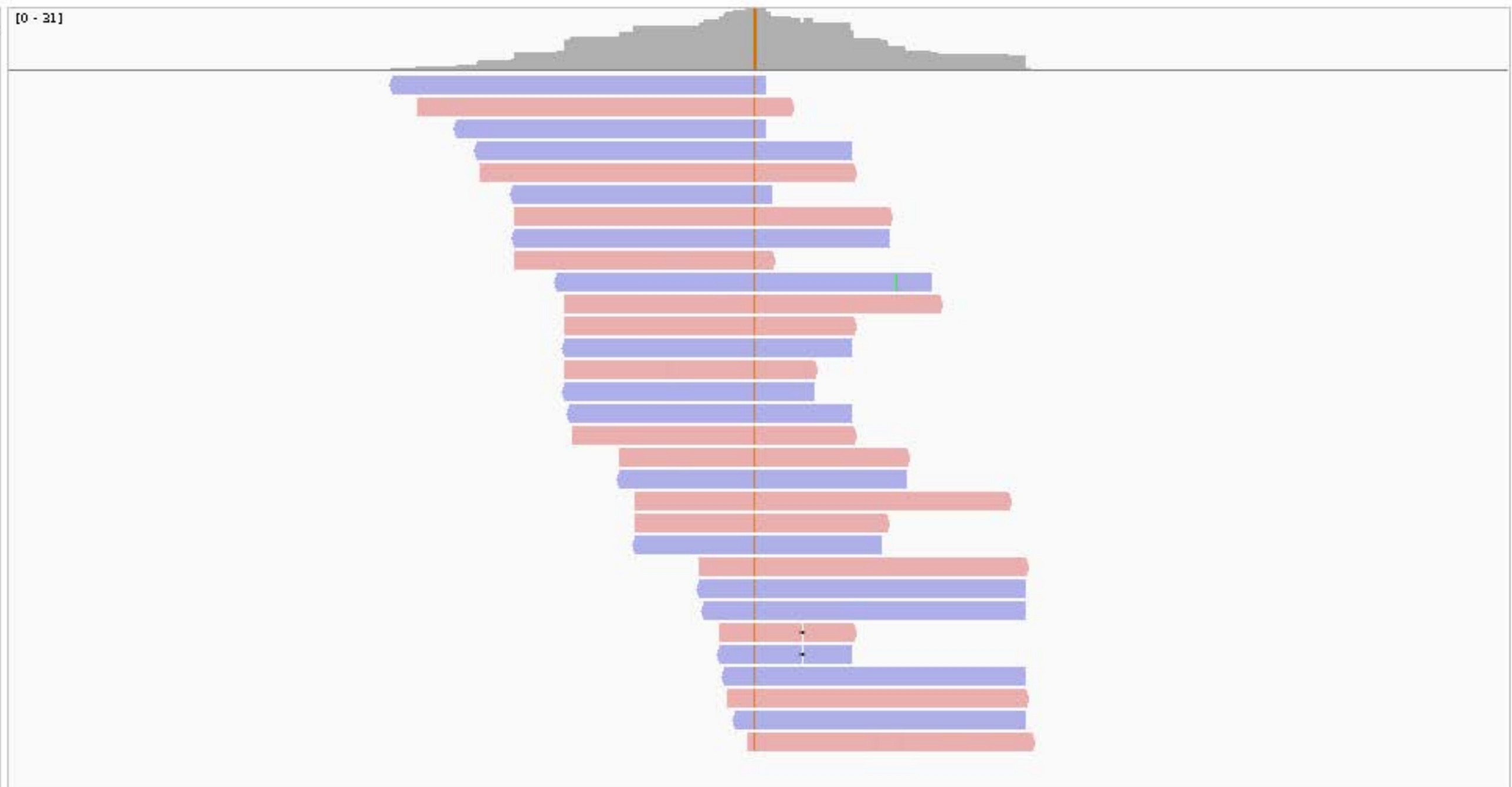
Refseq Genes





SL_0009_L_FFPE_26-B_mut.barrage

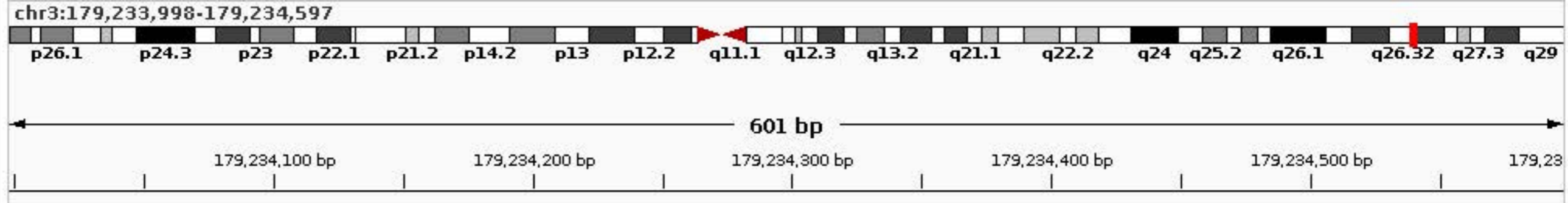
SL_0009_L_FFPE_26-B_mut.bam



Sequence

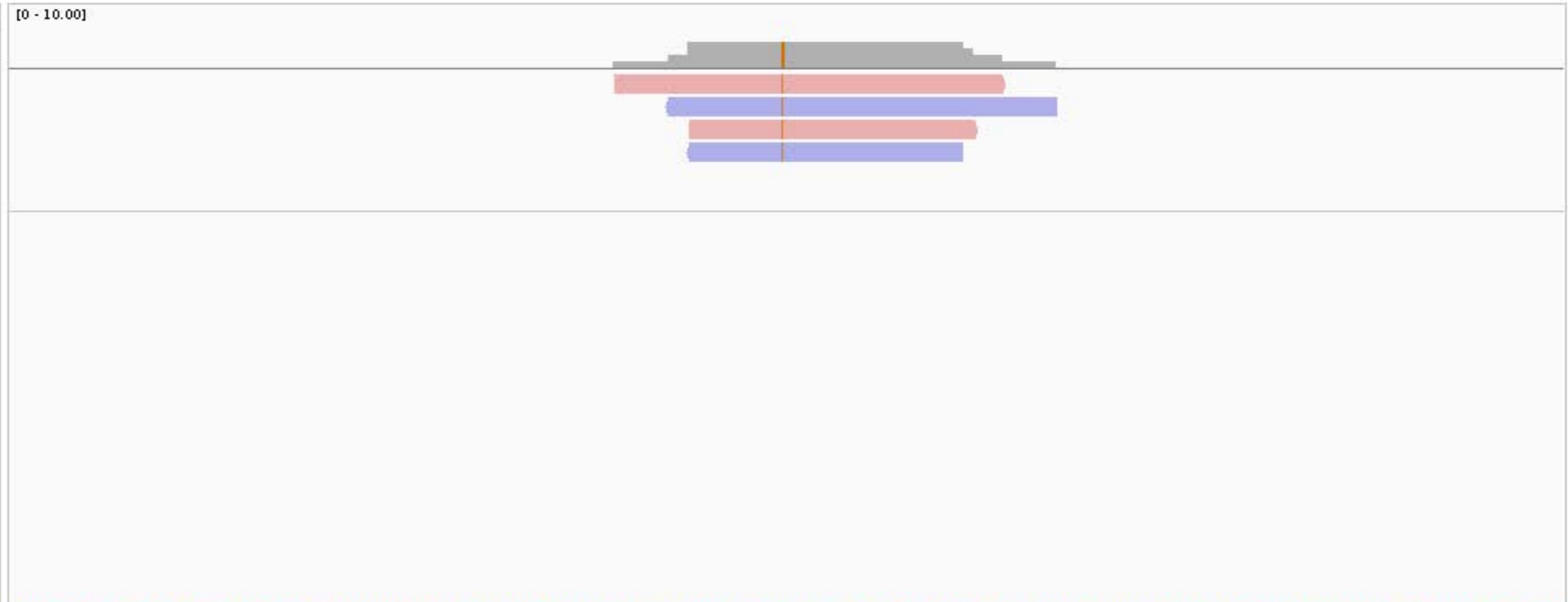
Refseq Genes





SL_0009_L_FFPE_30-B_mut.barr
age

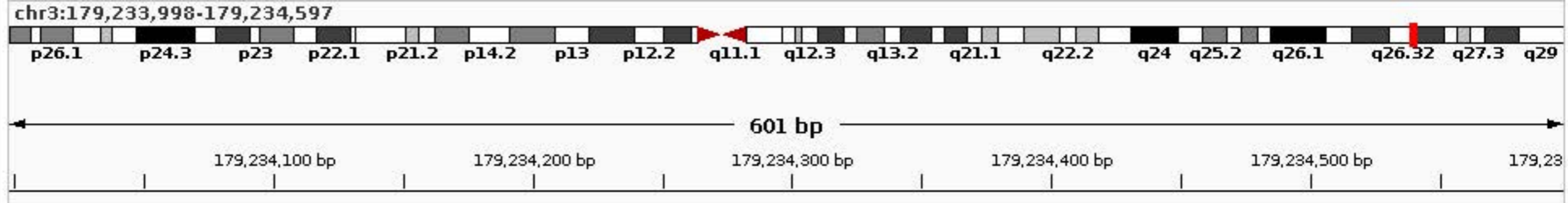
SL_0009_L_FFPE_30-B_mut.ba
m



Sequence →

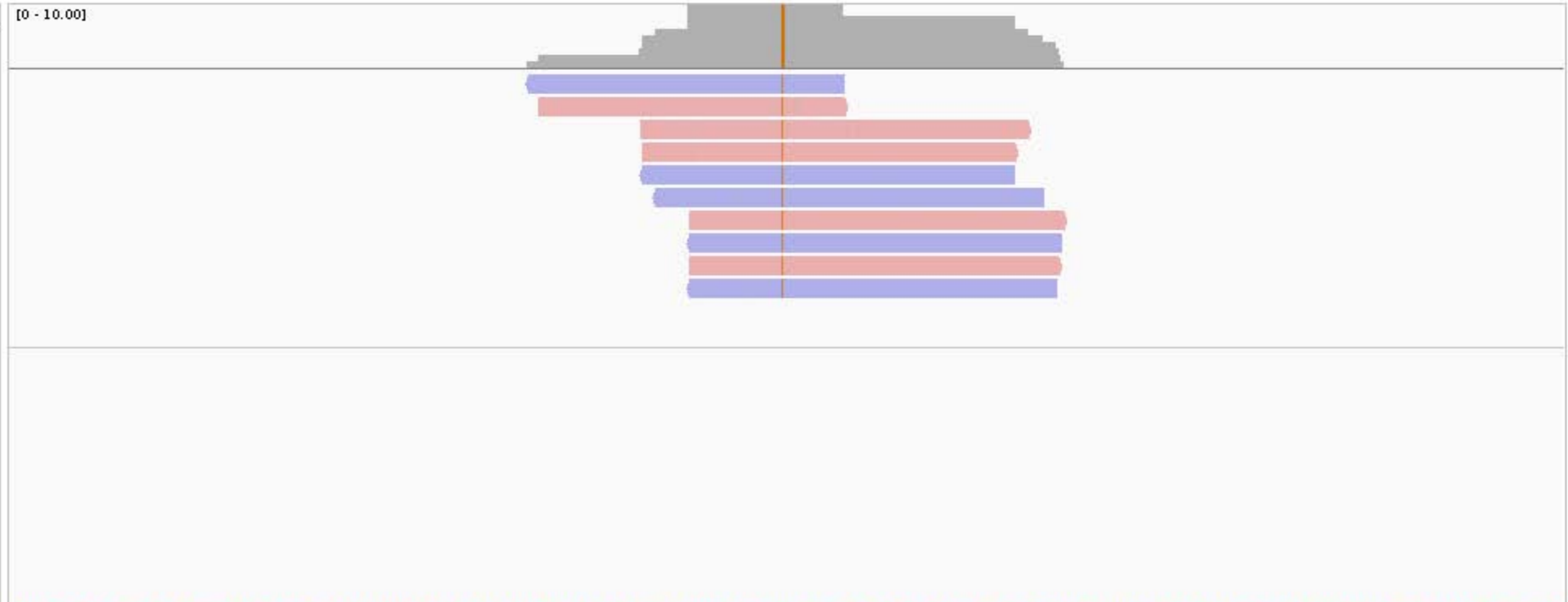
Refseq Genes





SL_0010_L_FFPE_18-B_mut.bam
range

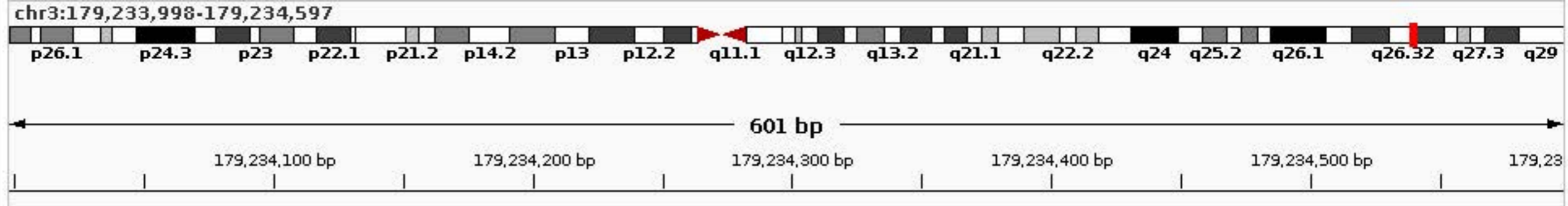
SL_0010_L_FFPE_18-B_mut.bam



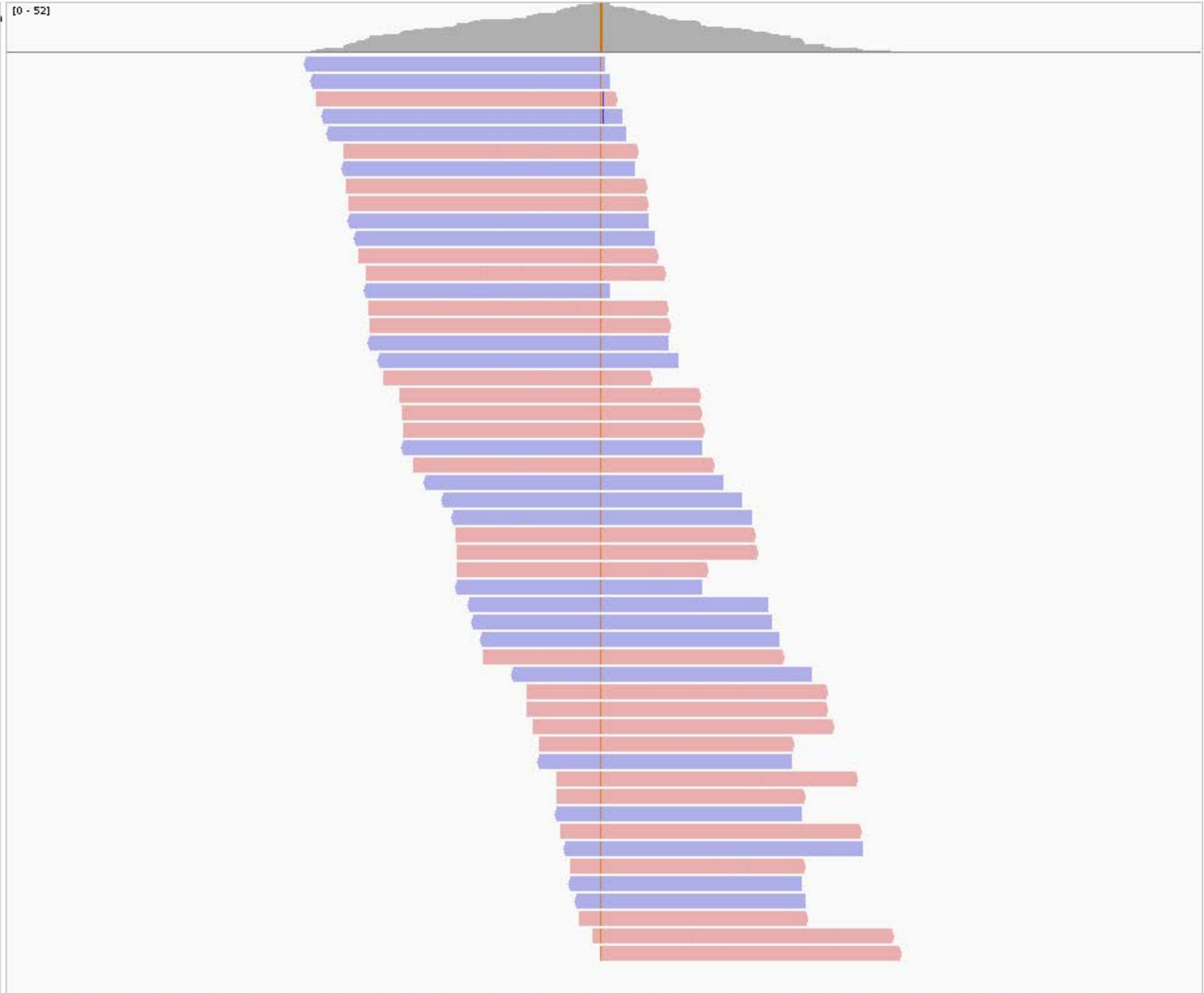
Sequence →

Refseq Genes





SL_0010_R_FFPE_42-B_mut.bam
range



SL_0010_R_FFPE_42-B_mut.bam

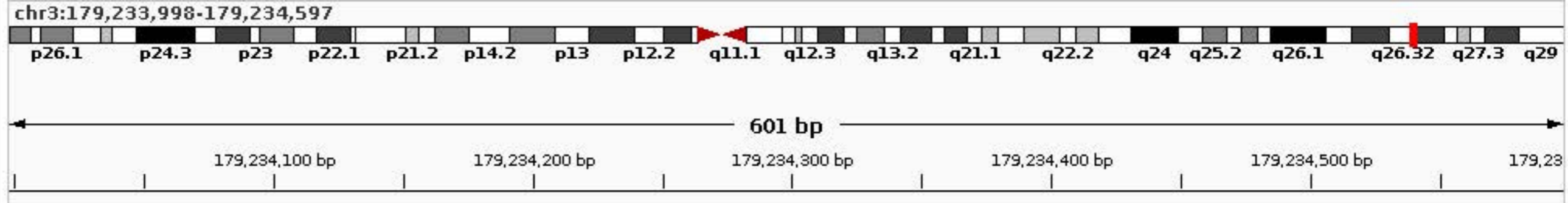
Sequence →



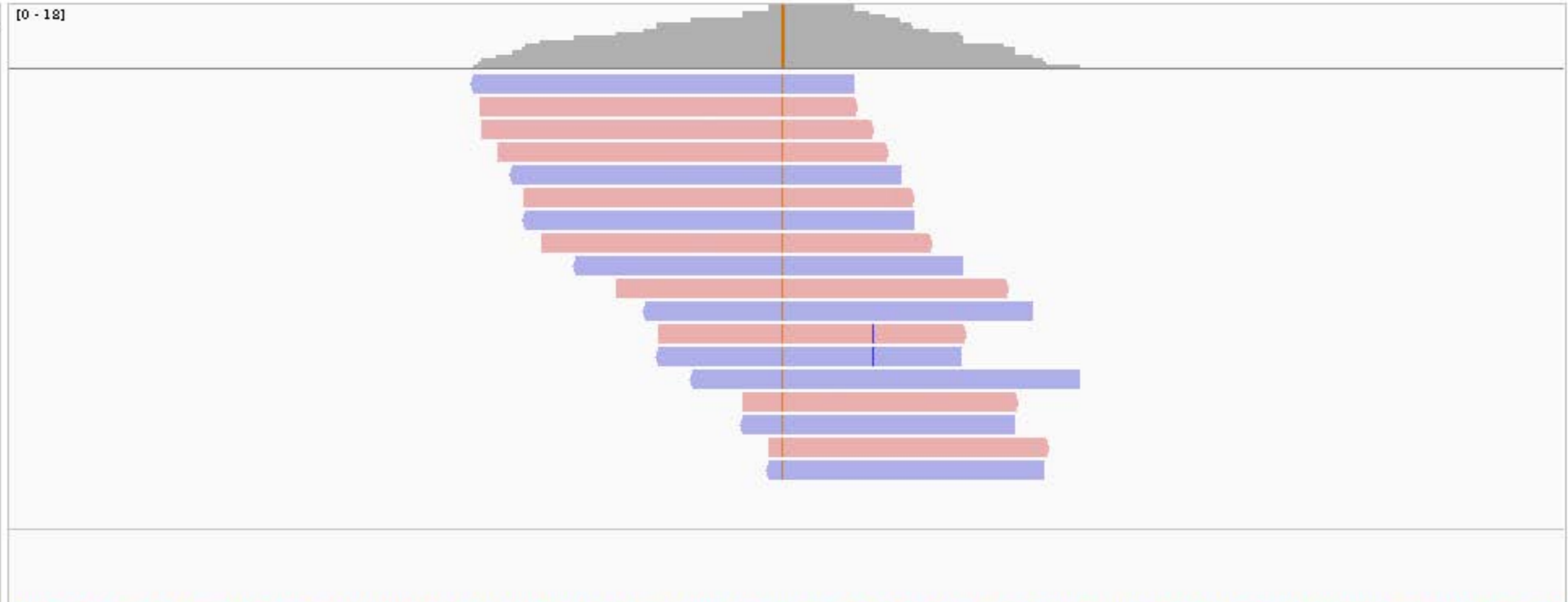
Refseq Genes



PIK3CA



SL_0011_L_FFPE_18-B_mut.bam
 SL_0011_L_FFPE_18-B_mut.bam

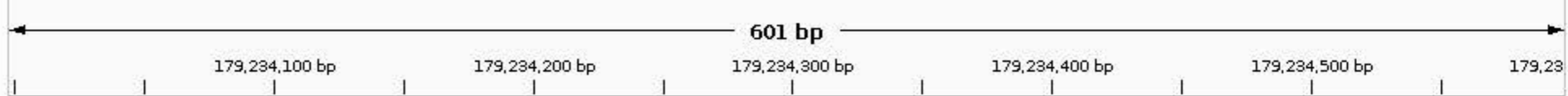


Sequence →

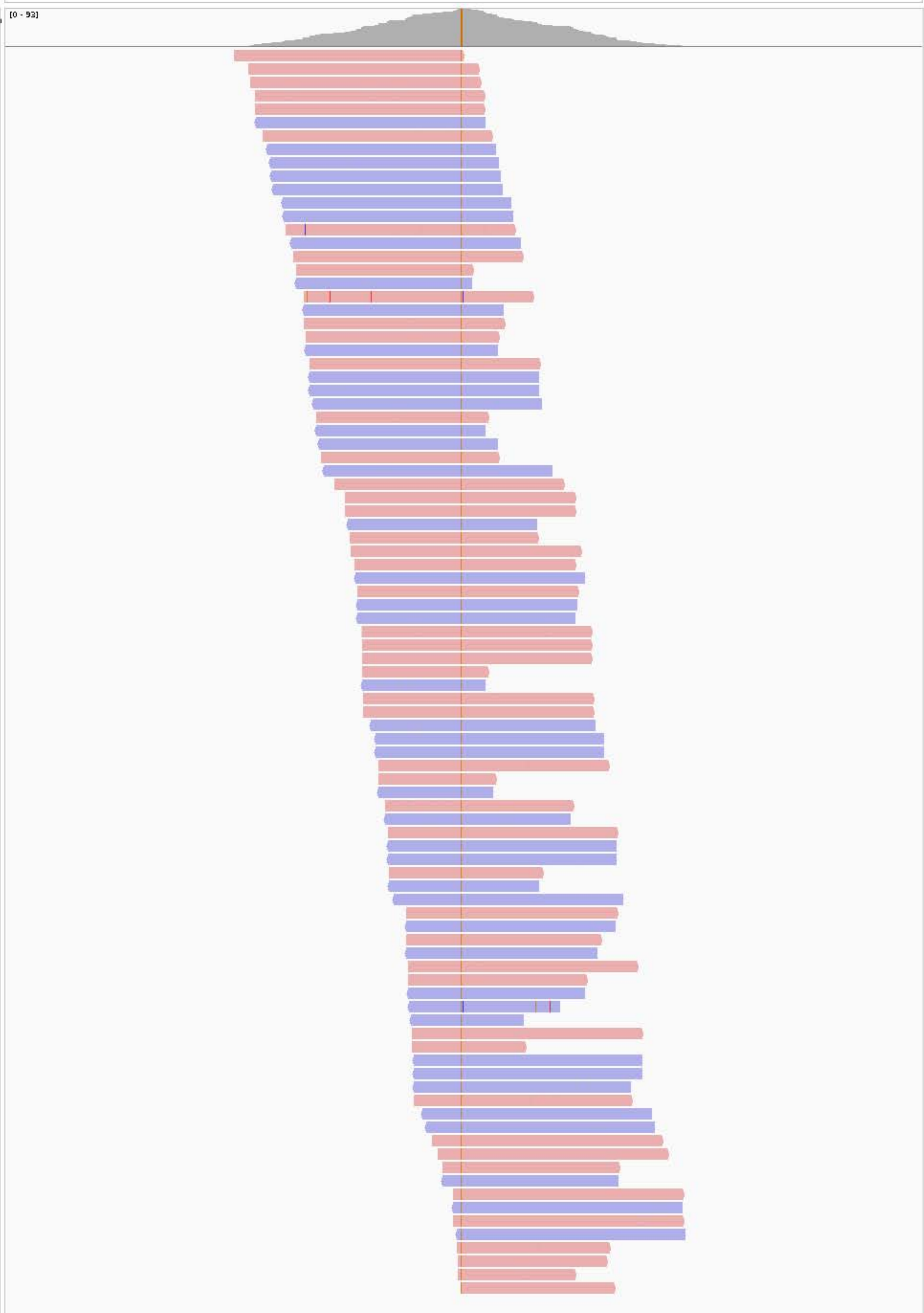
Refseq Genes



chr3:179,233,998-179,234,597
p26.1 p24.3 p23 p22.1 p21.2 p14.2 p13 p12.2 q11.1 q12.3 q13.2 q21.1 q22.2 q24 q25.2 q26.1 q26.32 q27.3 q29



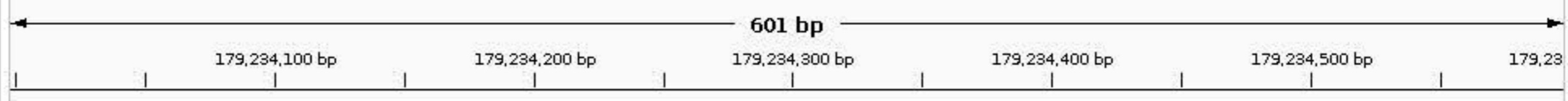
SL_1002_R_FFPE_33-B_mut.bam
rage



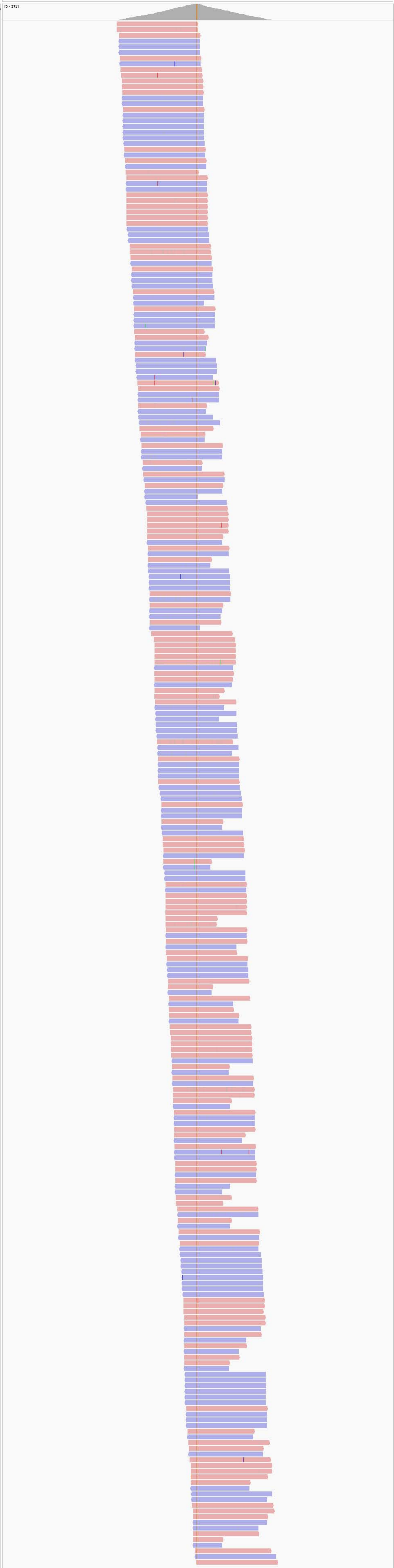
SL_1002_R_FFPE_33-B_mut.bam

Sequence →
Refseq Genes





SL_1004_R_49_FFPE-8_mut.bam
rage

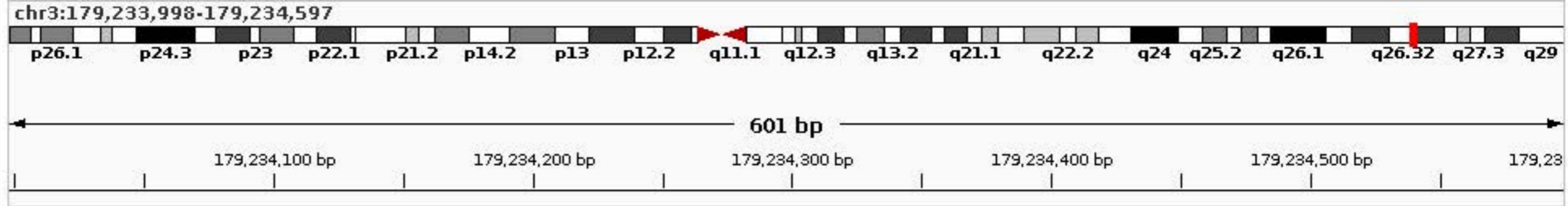


SL_1004_R_49_FFPE-8_mut.bam

Sequence
Refseq Genes

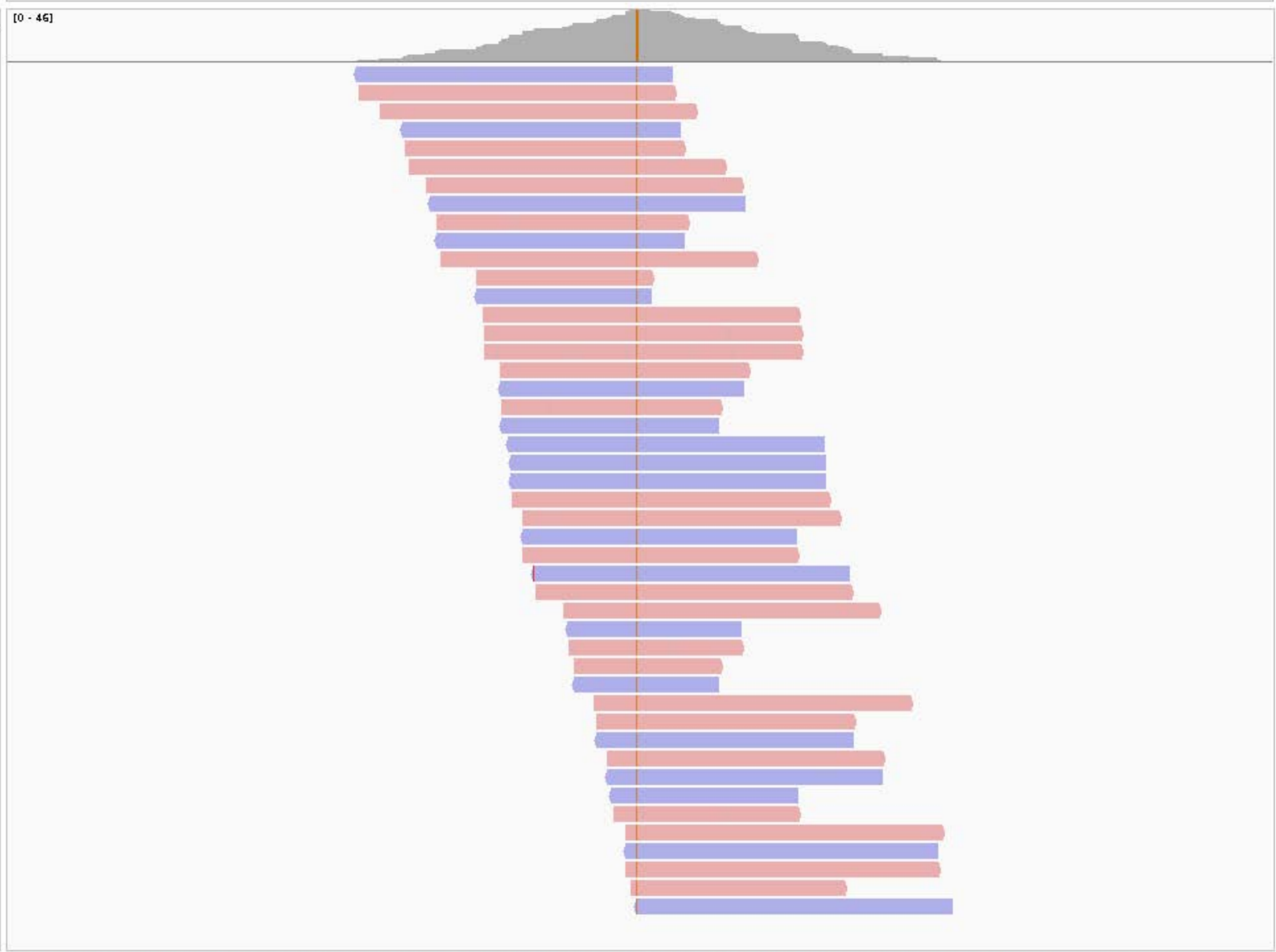


PIK3CA



SL_1007_L_FFPE_26-B_mut.bam
rage

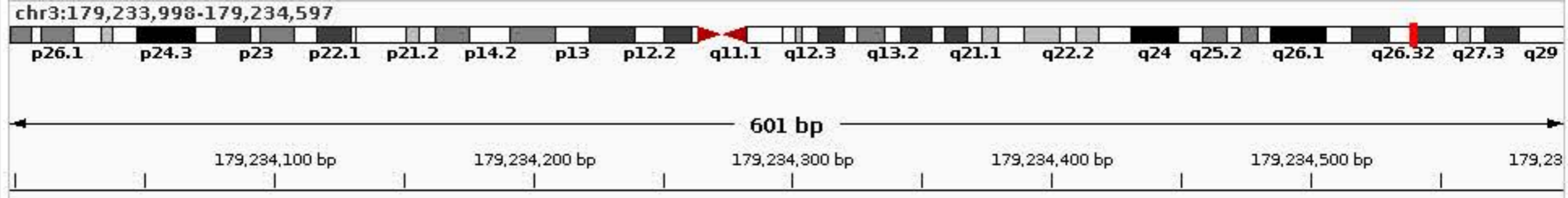
SL_1007_L_FFPE_26-B_mut.ba
m



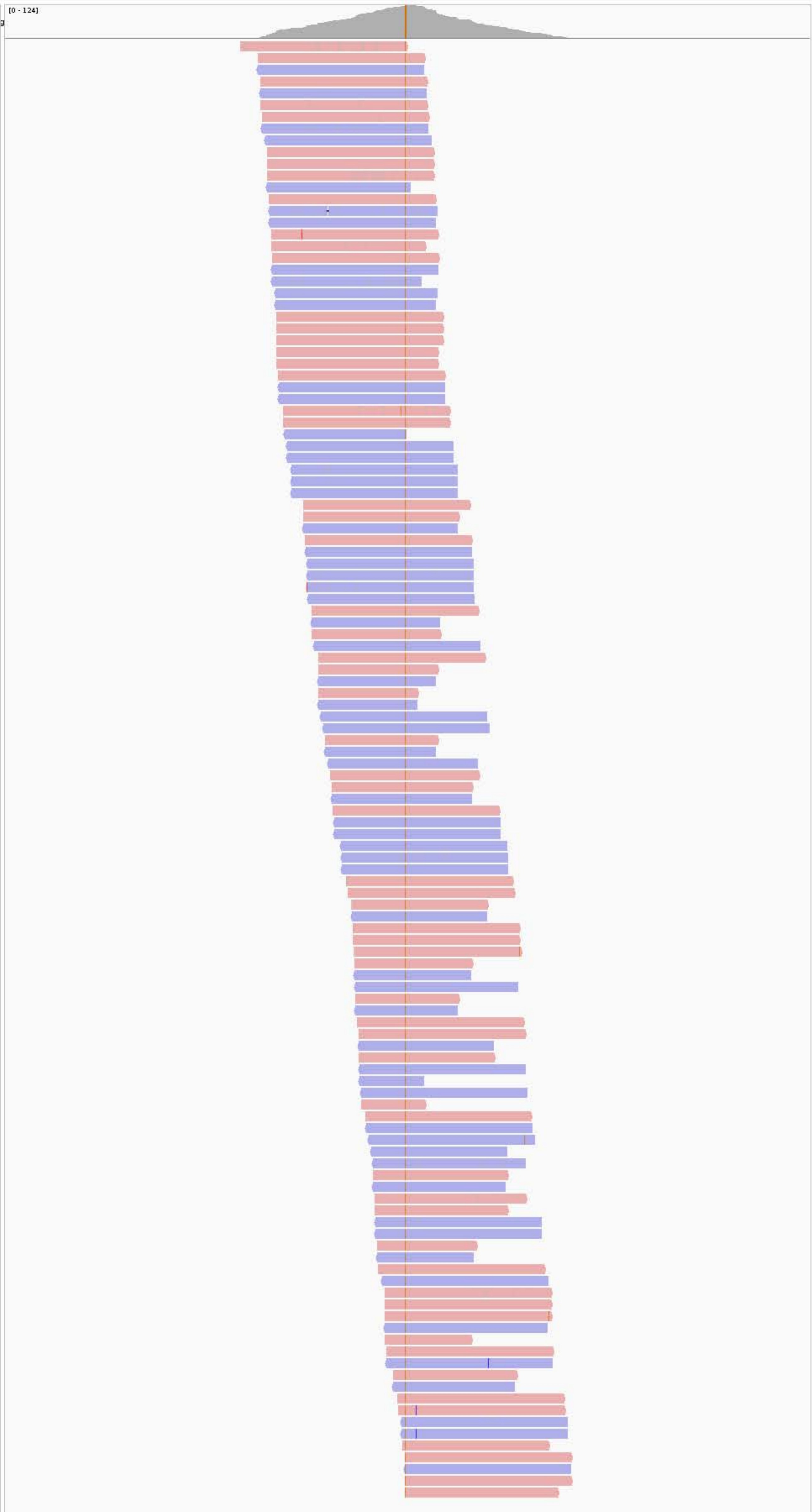
Sequence →

Refseq Genes





SL_1007-N1-B_mut.bam Coverage

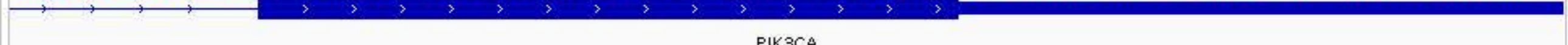


SL_1007-N1-B_mut.bam

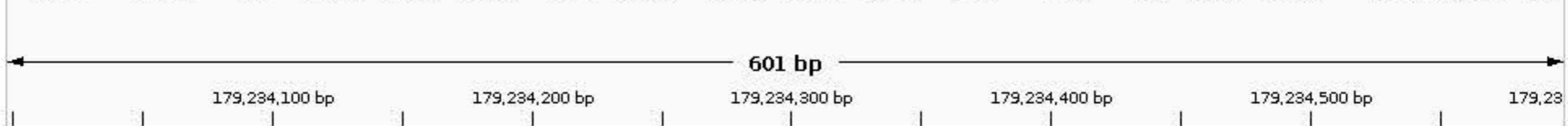
Sequence



Refseq Genes

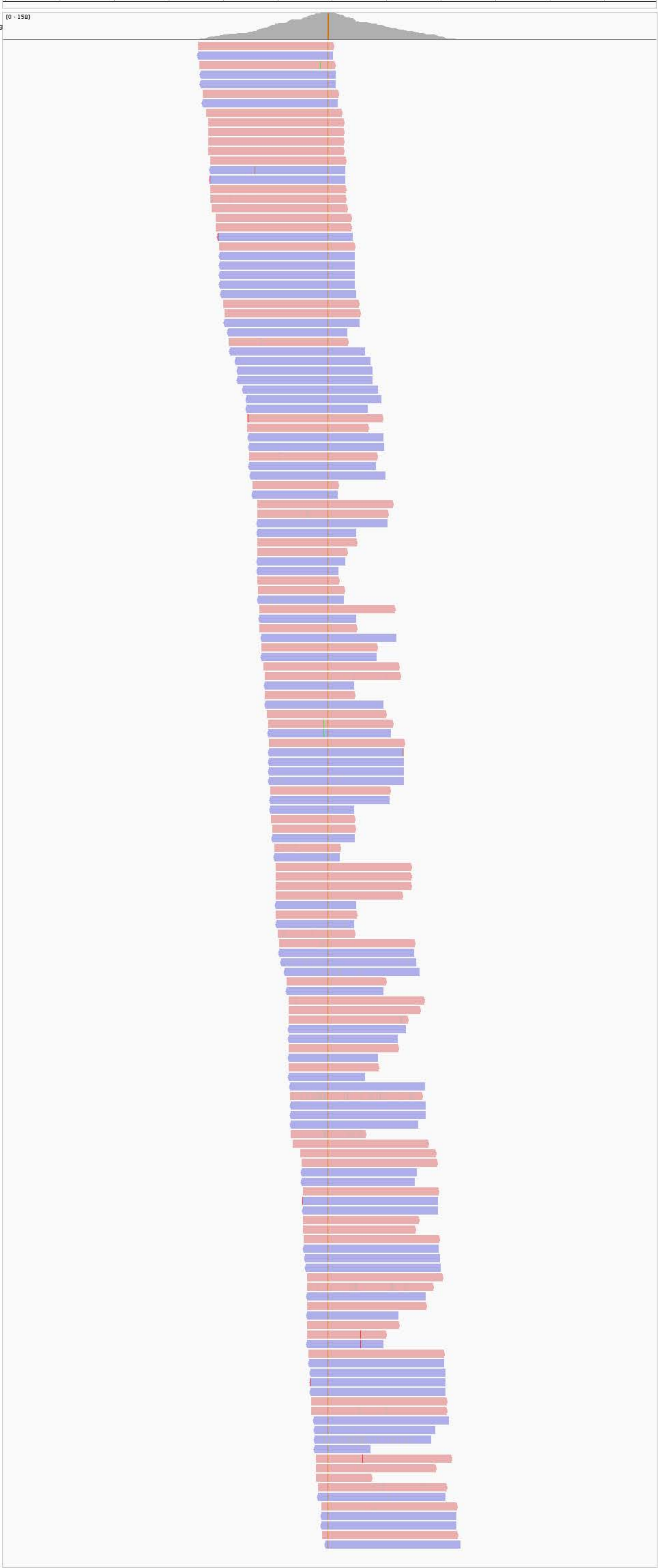


PIK3CA



SL_1007-N2-B_mut.bam Coverage

SL_1007-N2-B_mut.bam

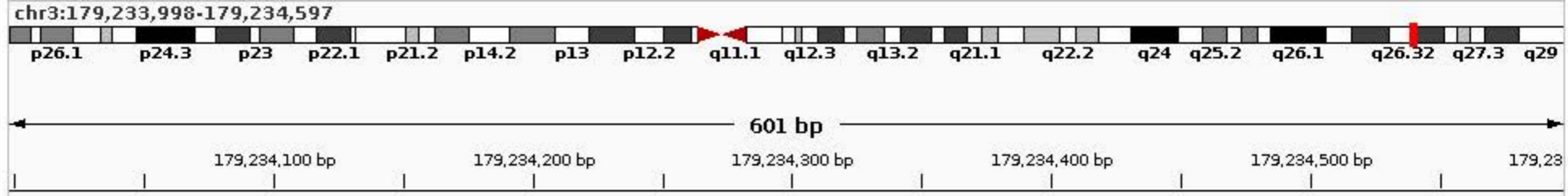


Sequence

Refseq Genes



PIK3CA



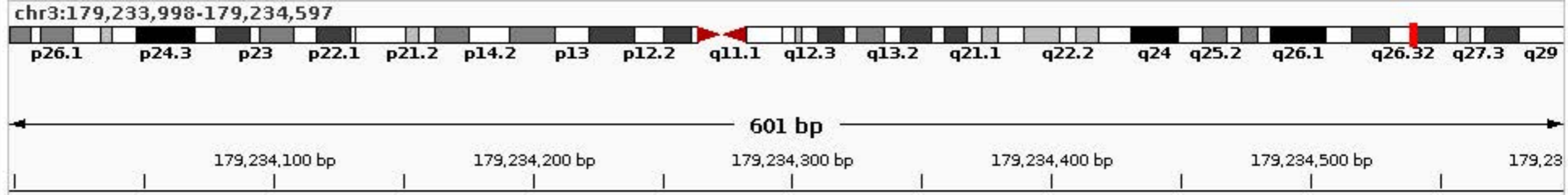
SL_1007-N3-B_mut.bam Coverage

SL_1007-N3-B_mut.bam

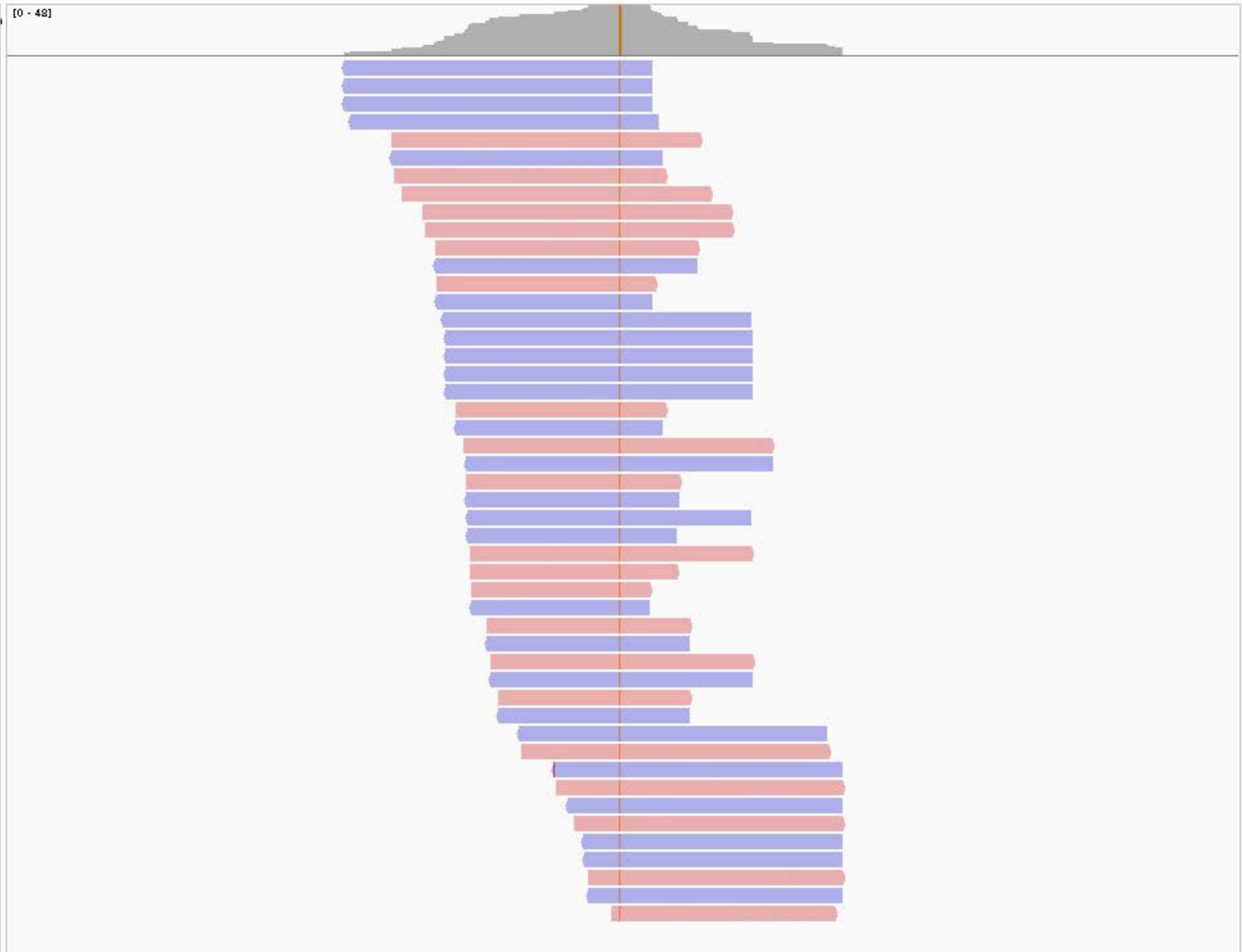
Sequence →

Refseq Genes





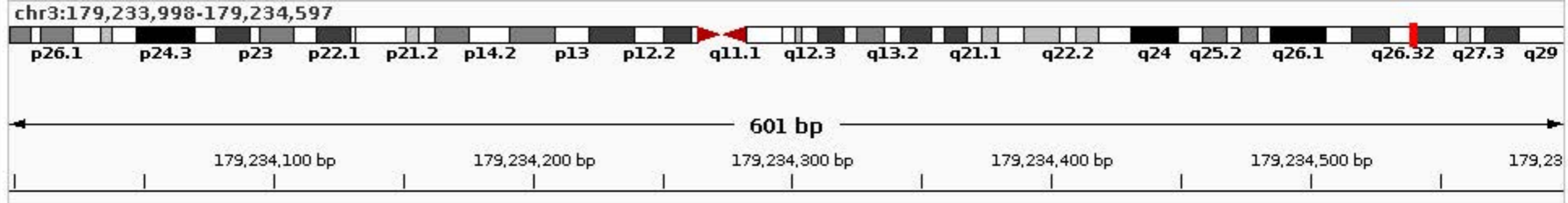
SL_1010_R_FFPE_15-B_mut.bam
rage



SL_1010_R_FFPE_15-B_mut.ba
m

Sequence →
Refseq Genes





SL_1011_R_FFPE_62-B_mut.bam
rage

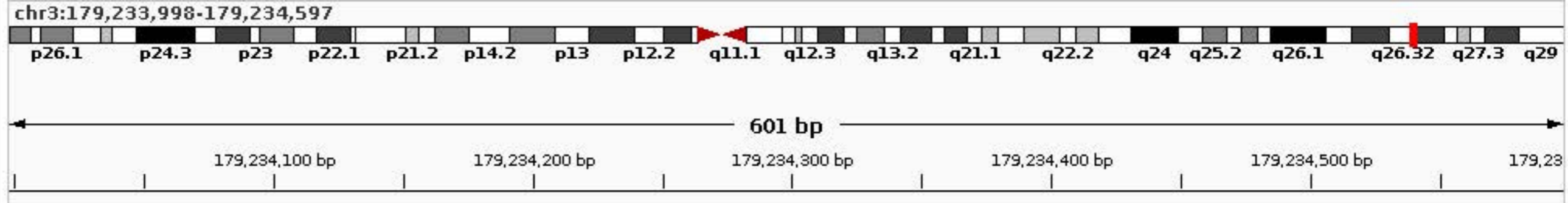
SL_1011_R_FFPE_62-B_mut.ba
m



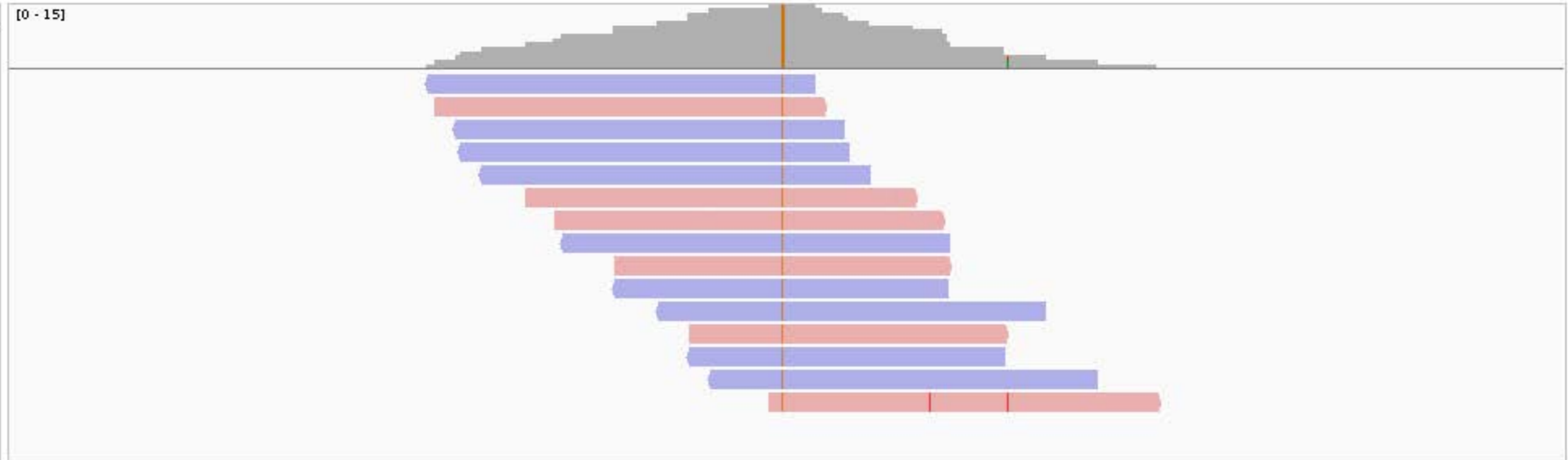
Sequence →

Refseq Genes





SL_1012_L_FFPE_47-B_mut.bam
range



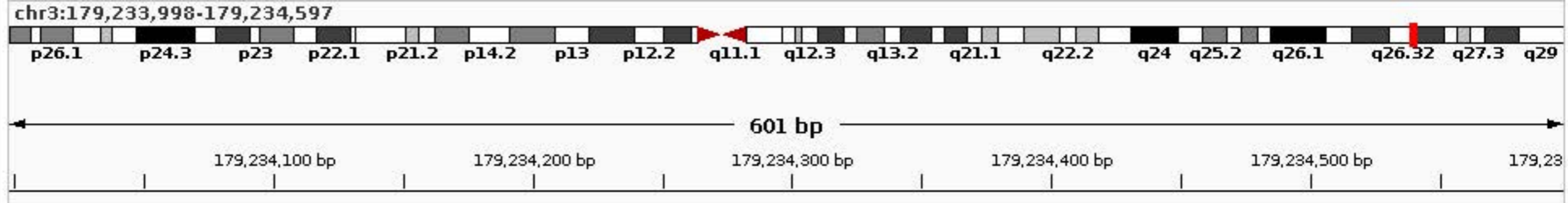
SL_1012_L_FFPE_47-B_mut.ba
m

Sequence →



Refseq Genes





SL_1013_R_12_FFPE-B_mut.bam
range

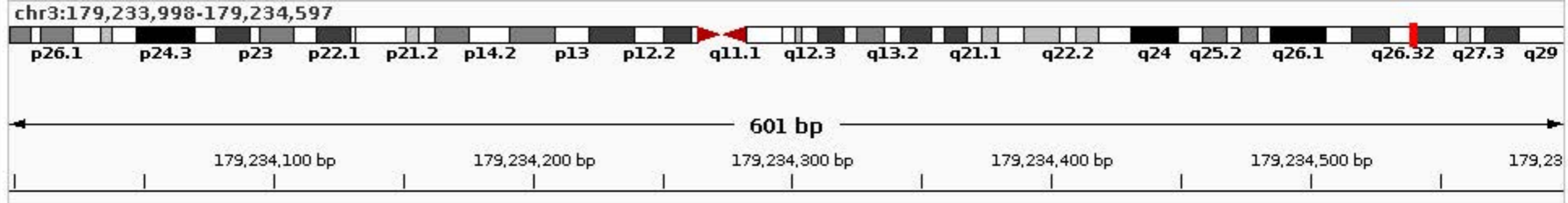
SL_1013_R_12_FFPE-B_mut.bam



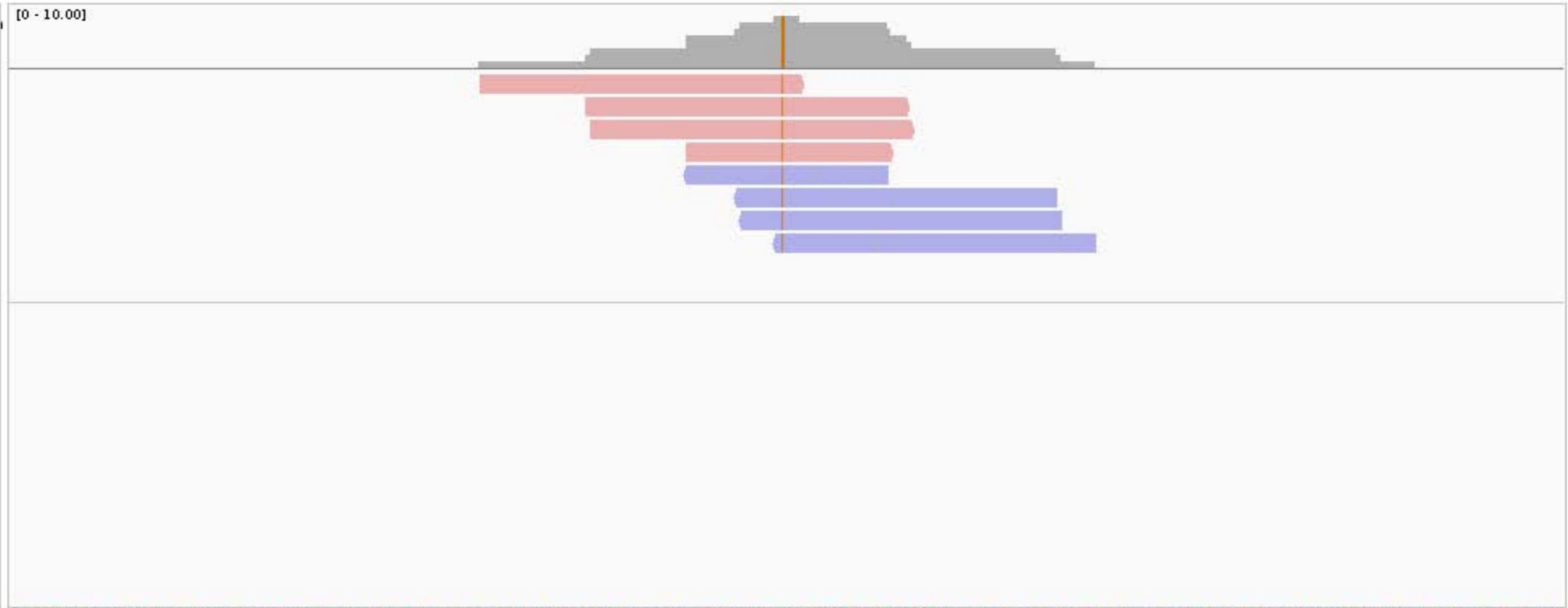
Sequence →

Refseq Genes



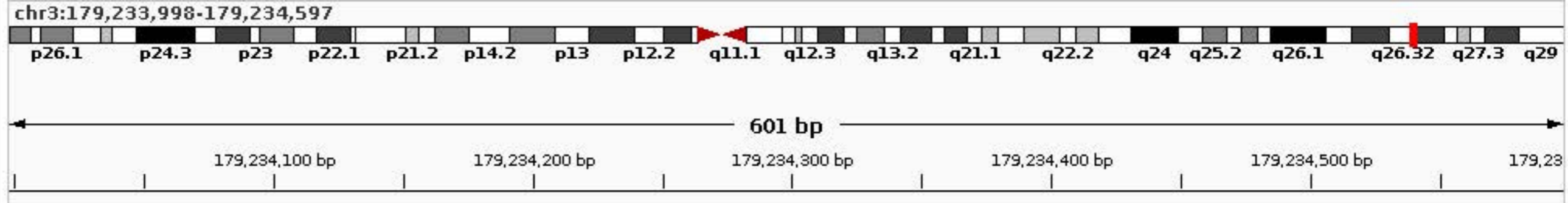


SL_1015_R_63_FFPE-B_mut.bam
SL_1015_R_63_FFPE-B_mut.bam



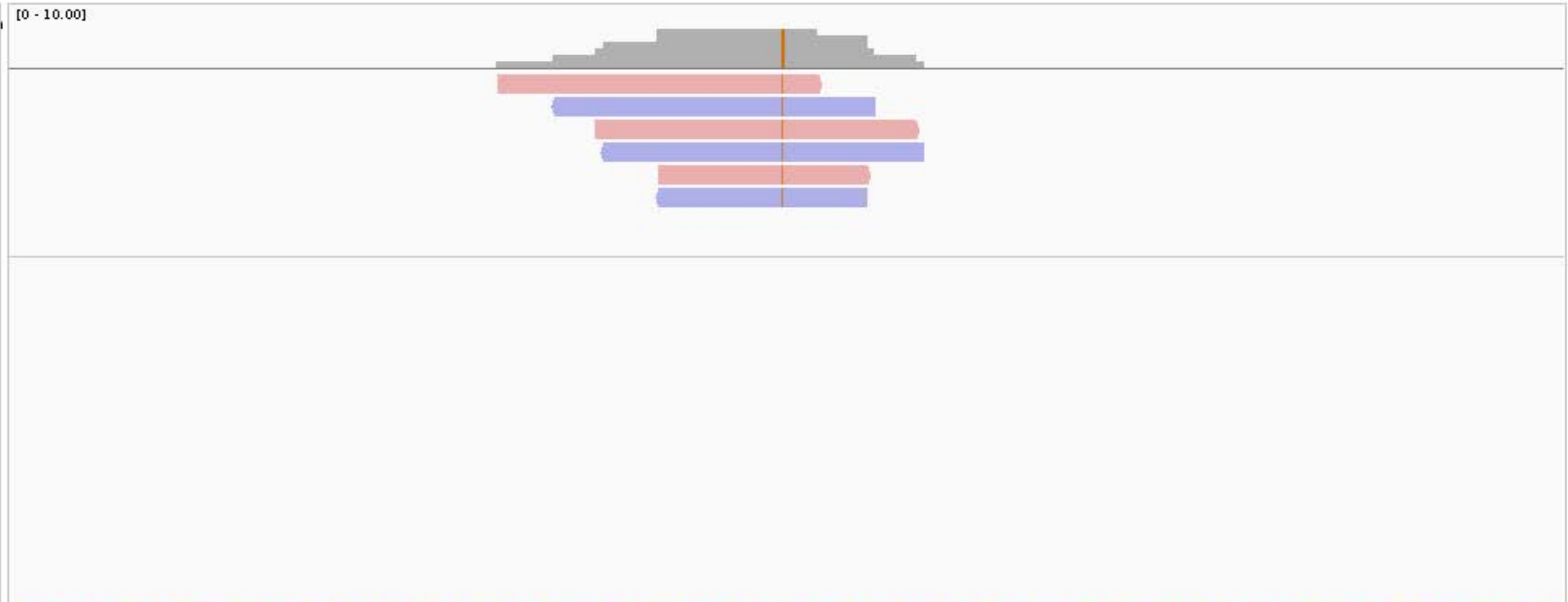
Sequence →
Refseq Genes





SL_1015_R_78_FFPE-B_mut.bam
range

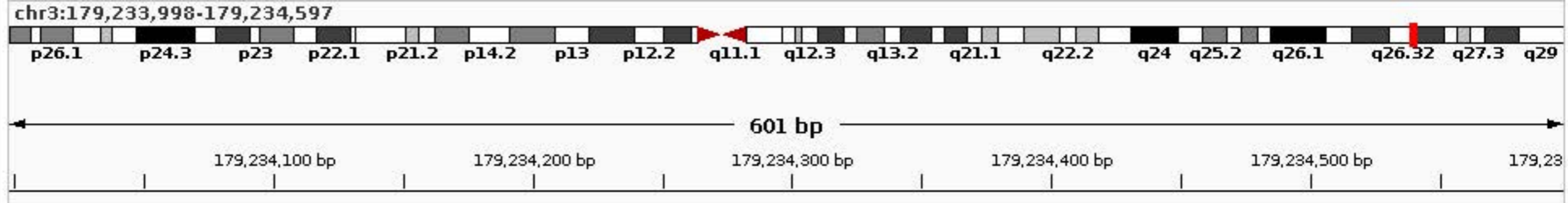
SL_1015_R_78_FFPE-B_mut.ba
m



Sequence →

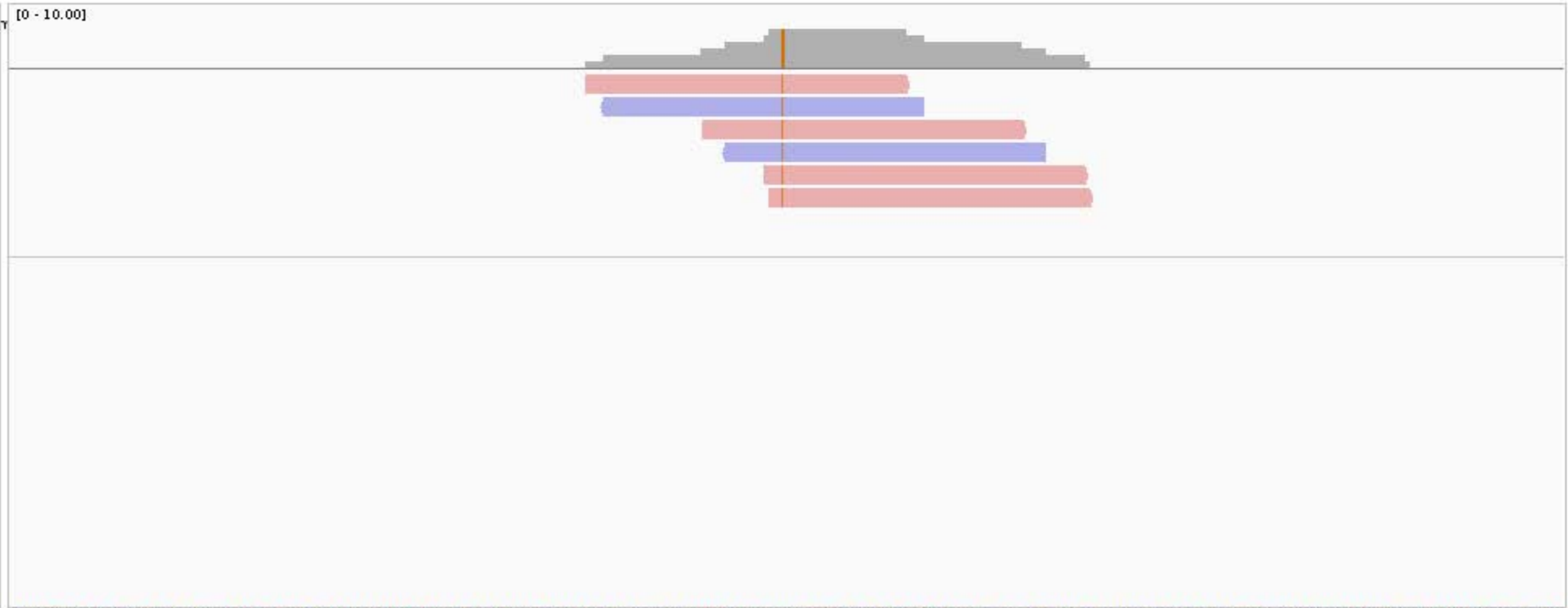
Refseq Genes





SL_1015_R_100_FFPE-B_mut.bam
verage

SL_1015_R_100_FFPE-B_mut.bam



Sequence →

Refseq Genes

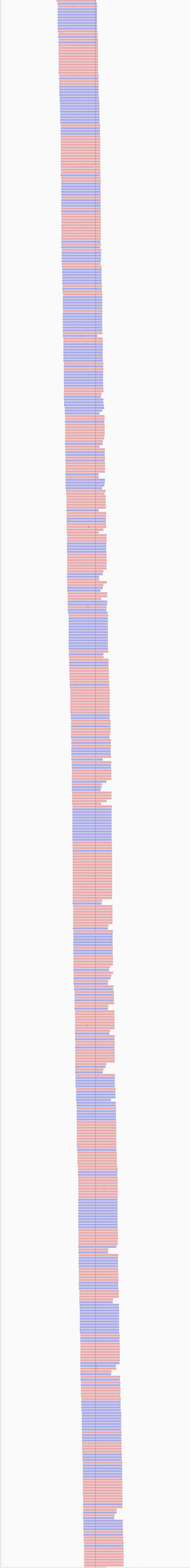


SL_0012_RT_3-8_mut.bam Cov

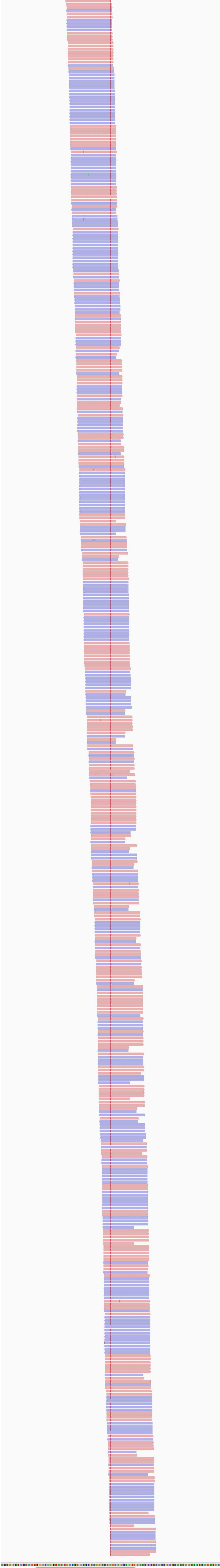
SL_0012_RT_3-8_mut.bam

Sequence

RefSeq Genes

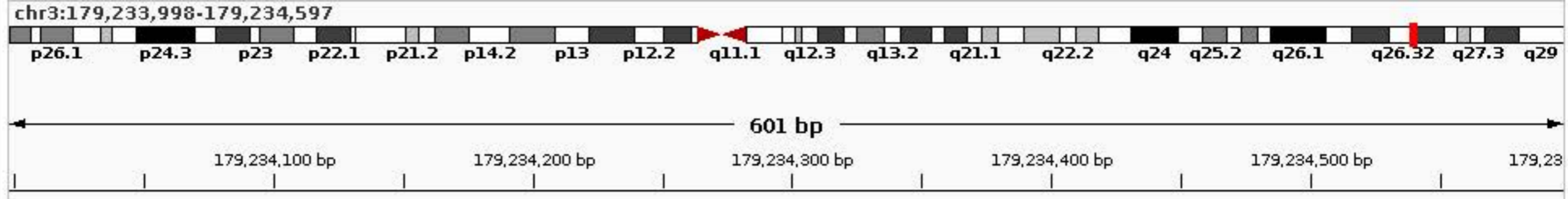


SL_0012_RT_4B_mut.bam Cov



Sequence
Refseq Genes





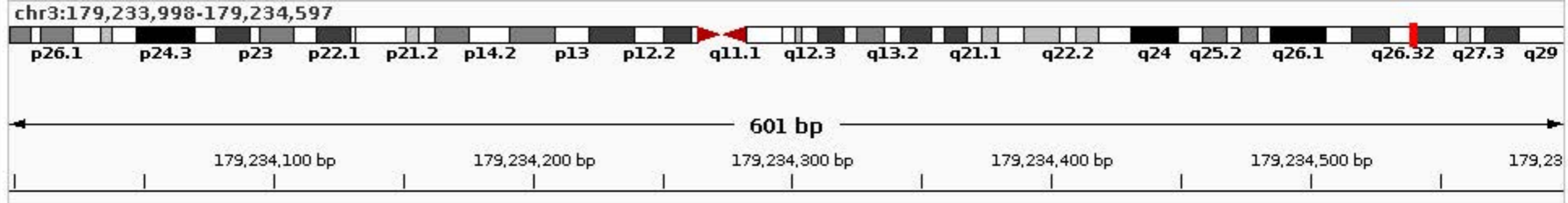
SL_1002_L_FFPE_15-B_mut.barrage



SL_1002_L_FFPE_15-B_mut.bam

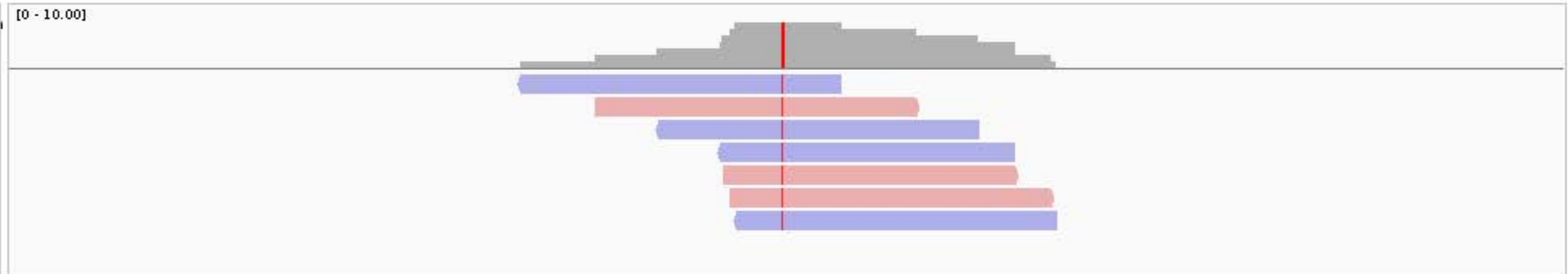
Sequence →
Refseq Genes





SL_1008_R_51_FFPE-B_mut.bam
range

SL_1008_R_51_FFPE-B_mut.bam

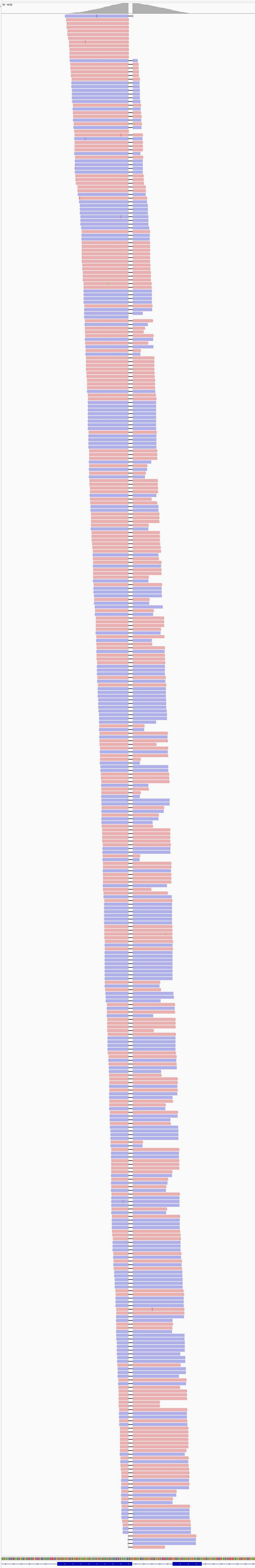


Sequence →

Refseq Genes

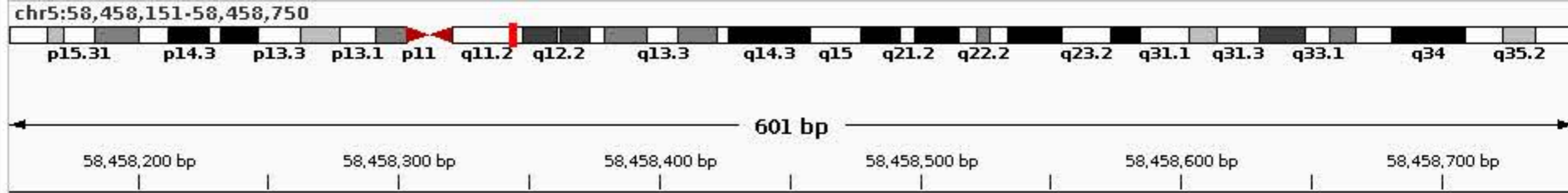


SL_0007_R_FFPE_39-B_mut.bam
rage

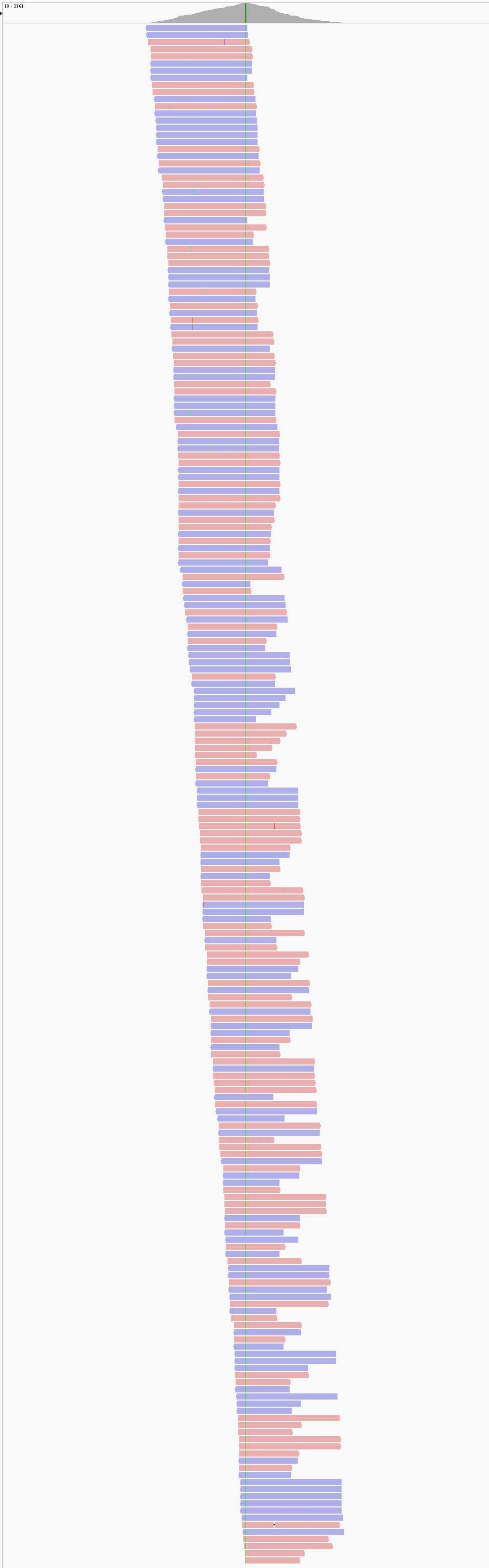


Sequence
Refseq Genes

PIK3R1



SL_1007-T1-B_mut.bam Coverage

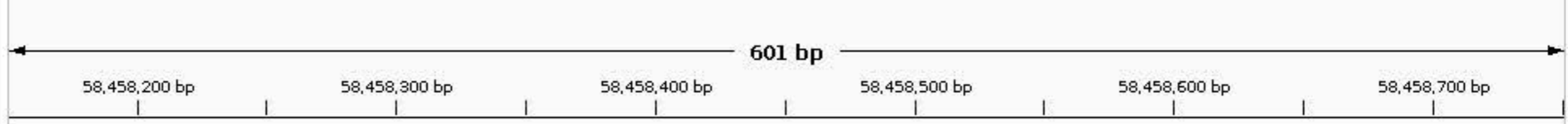


SL_1007-T1-B_mut.bam

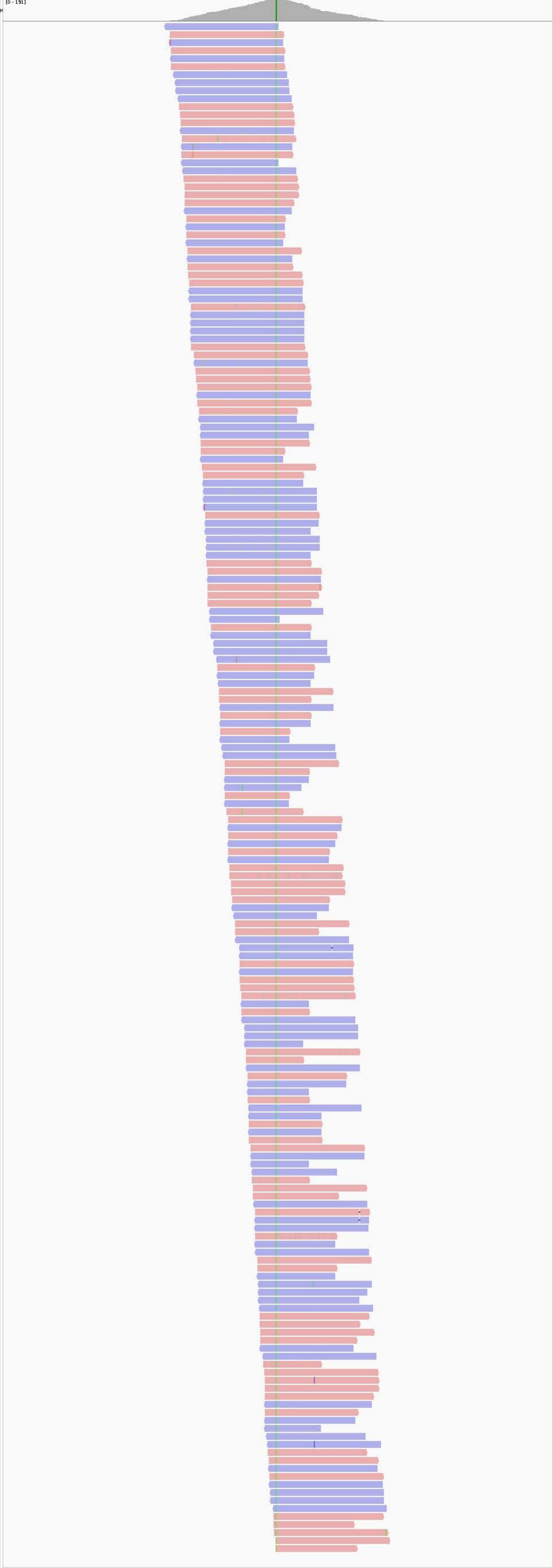
Sequence



PLK2



SL_1007-T2-B_mut.bam Coverage

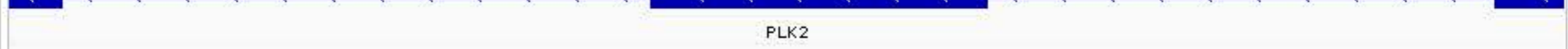


SL_1007-T2-B_mut.bam

Sequence

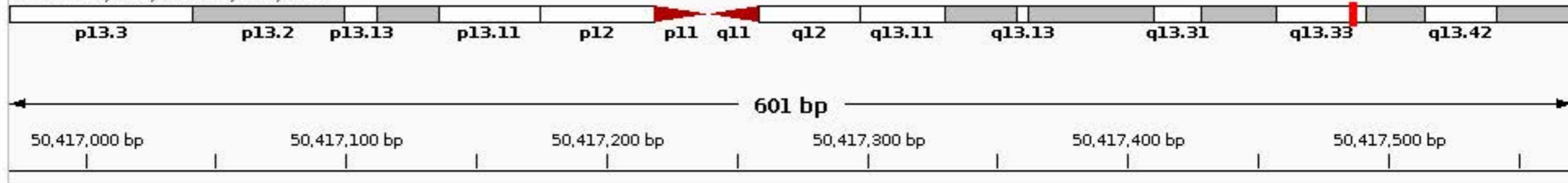


Refseq Genes

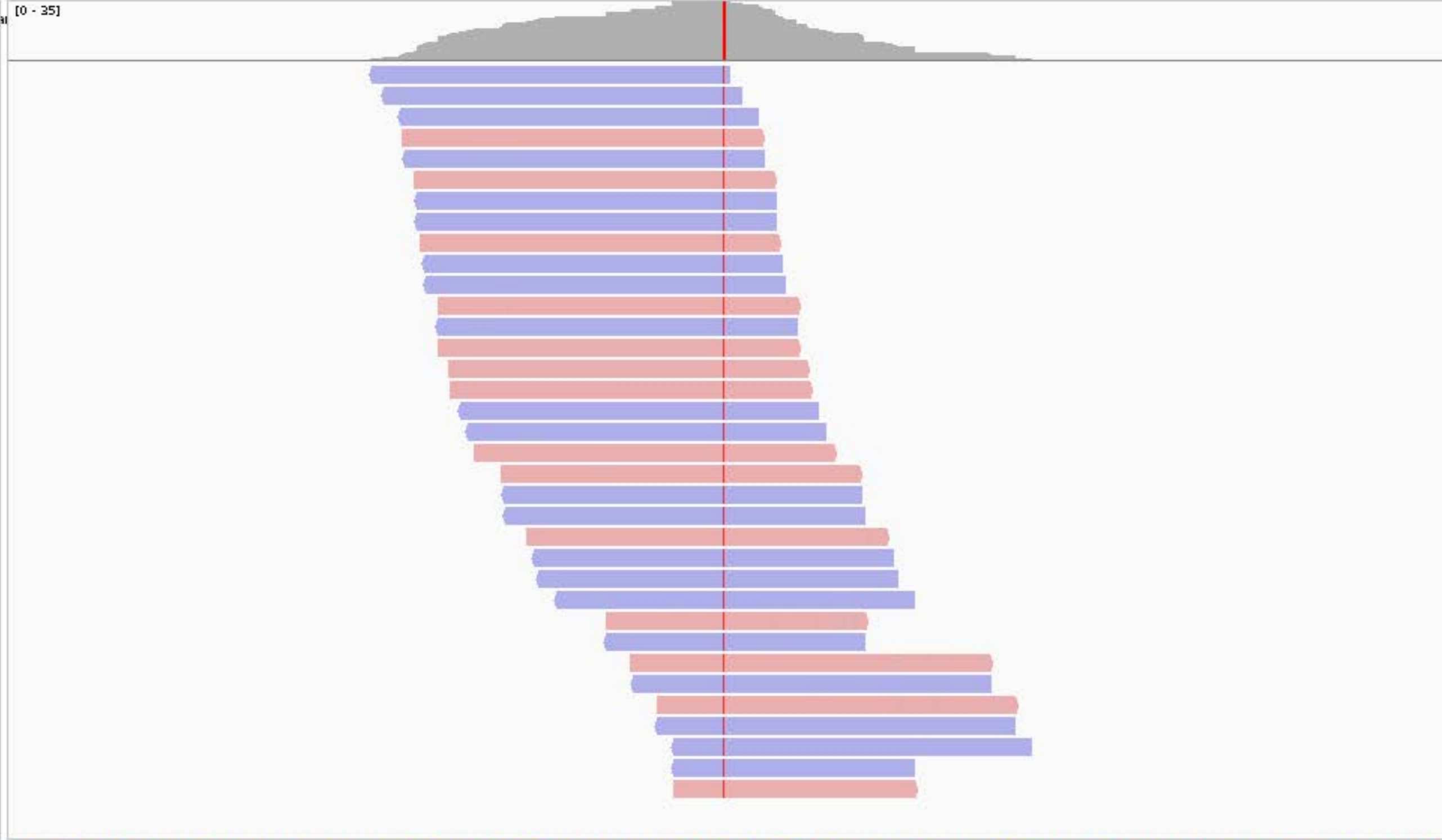


PLK2

chr19:50,416,971-50,417,570



SL_0007_R_FFPE_61_1-B_mut.bam
verage



SL_0007_R_FFPE_61_1-B_mut.
bam

Sequence →

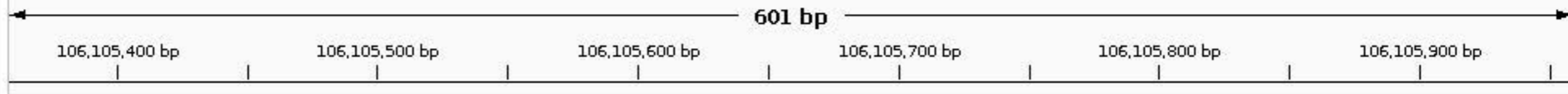


Refseq Genes



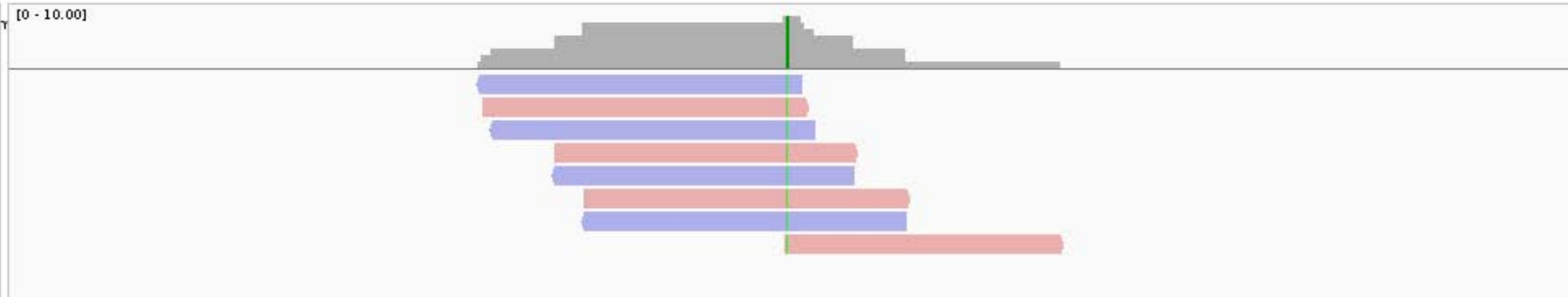
POLD1

chr6:106,105,359-106,105,958



SL_1015_R_100_FFPE-B_mut.bam
verage

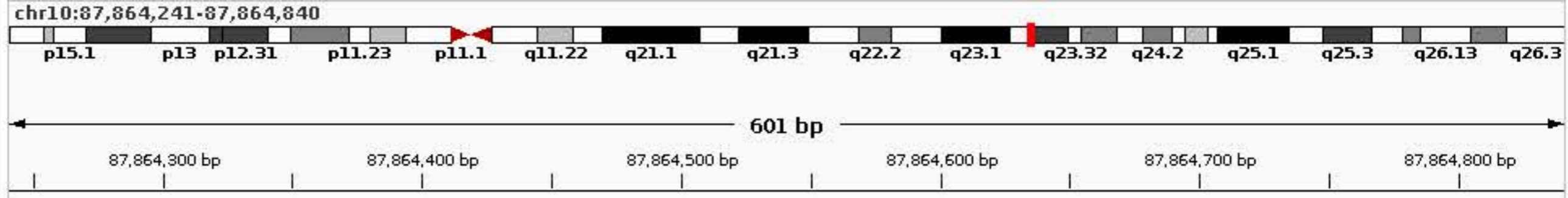
SL_1015_R_100_FFPE-B_mut.bam



Sequence →

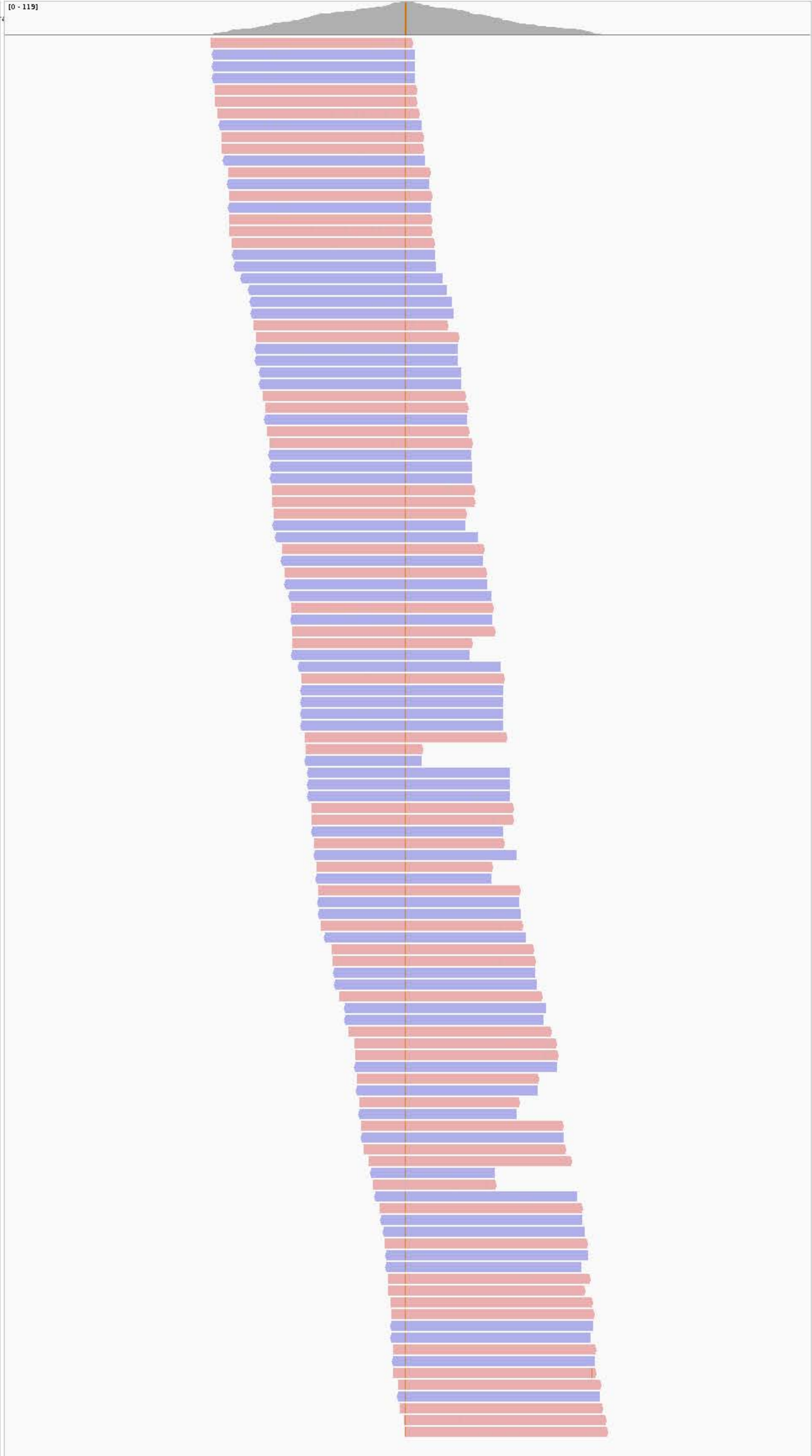
Refseq Genes





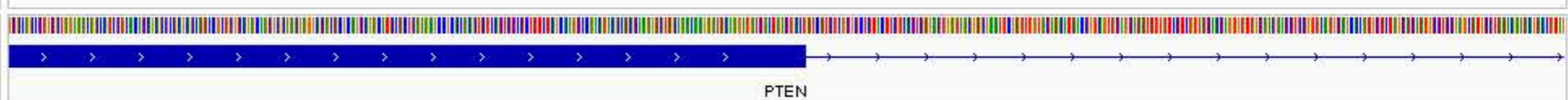
SL_0008_LT_8-B_mut.bam Coverage

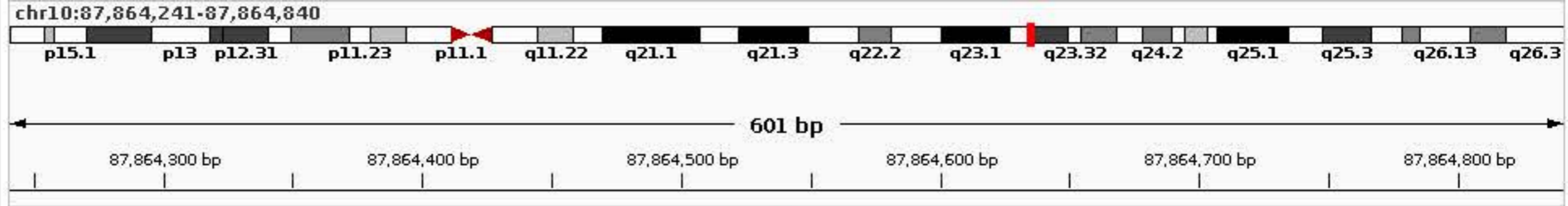
SL_0008_LT_8-B_mut.bam



Sequence

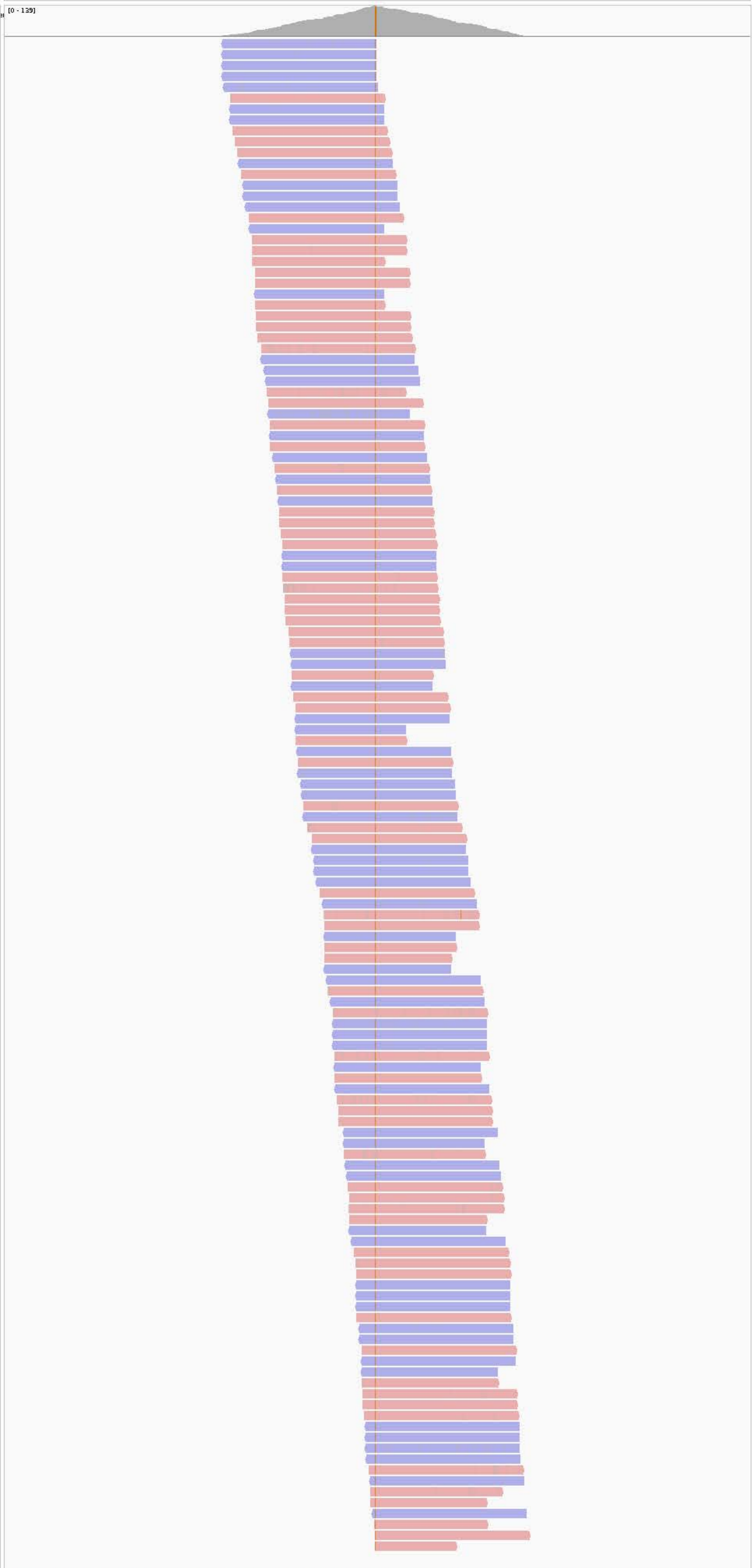
Refseq Genes





SL_0008-LT_15-B_mut.bam Coverage [0 - 129]

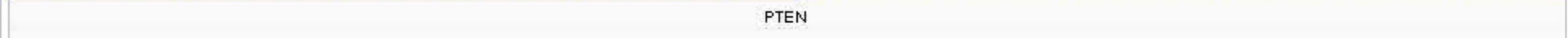
SL_0008-LT_15-B_mut.bam



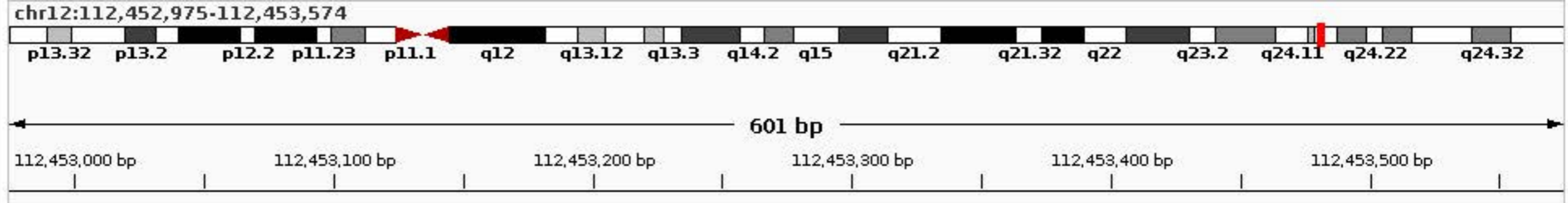
Sequence



Refseq Genes

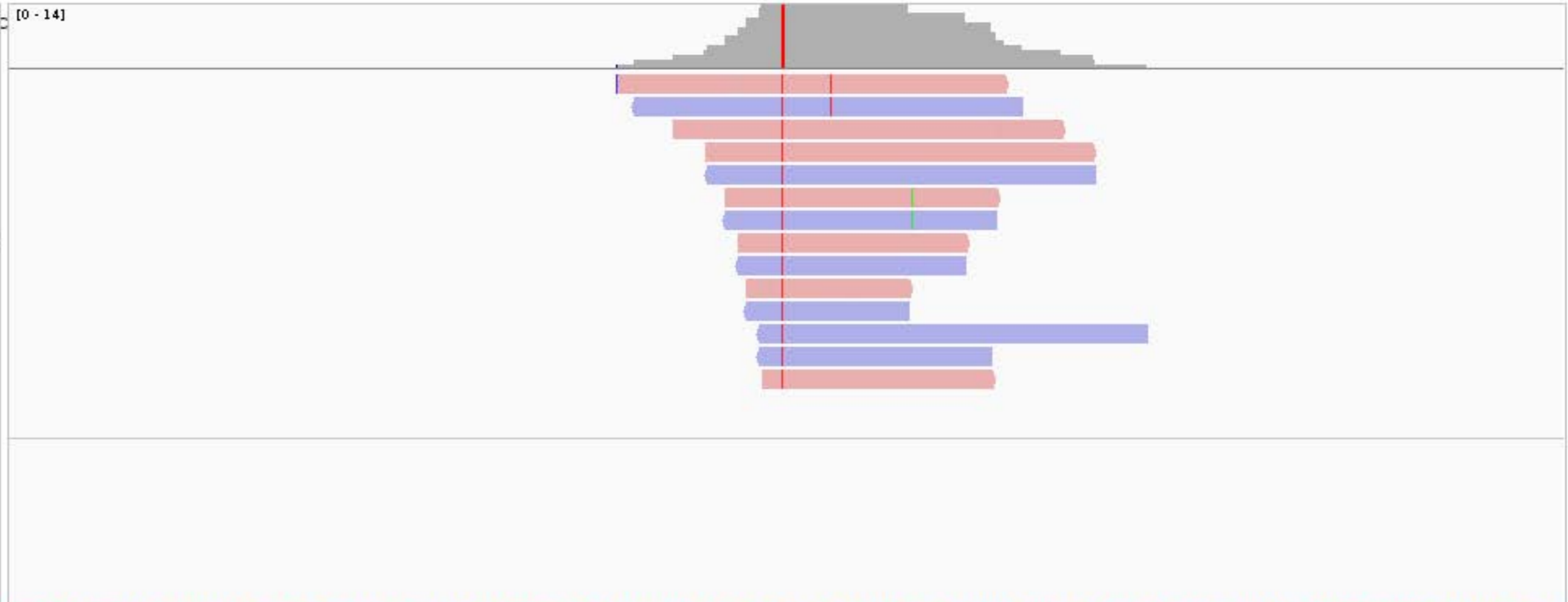


PTEN



SL_1010_L_FFPE_3-B_mut.bam
age

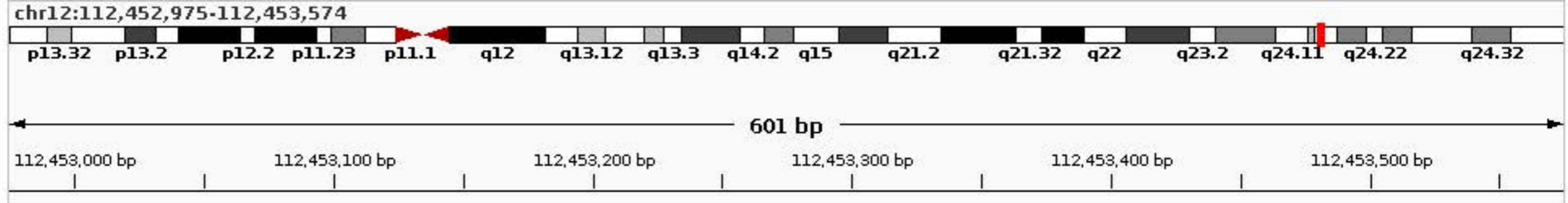
SL_1010_L_FFPE_3-B_mut.bam



Sequence →

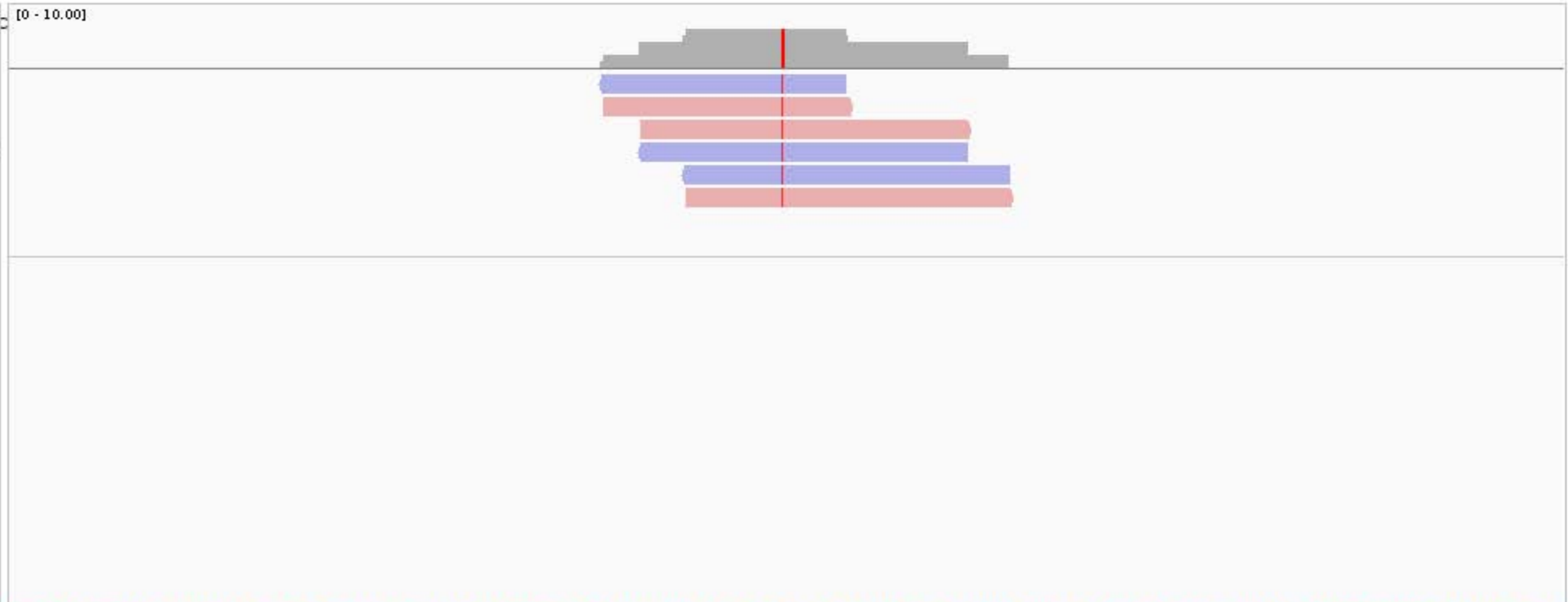
Refseq Genes





SL_1010_L_FFPE_8-B_mut.bam
age

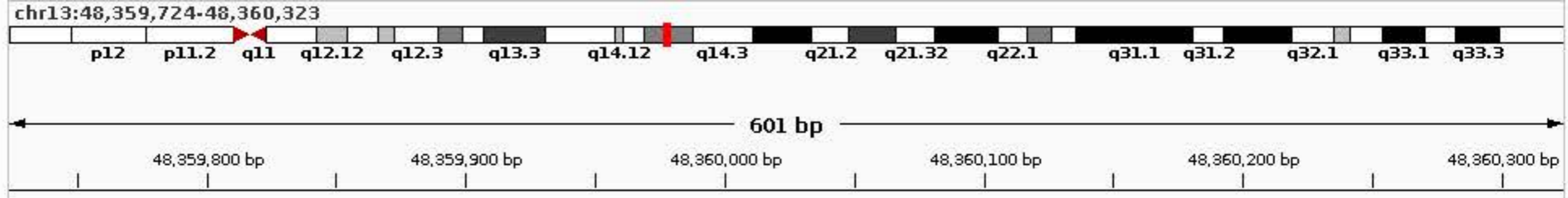
SL_1010_L_FFPE_8-B_mut.bam



Sequence →

Refseq Genes



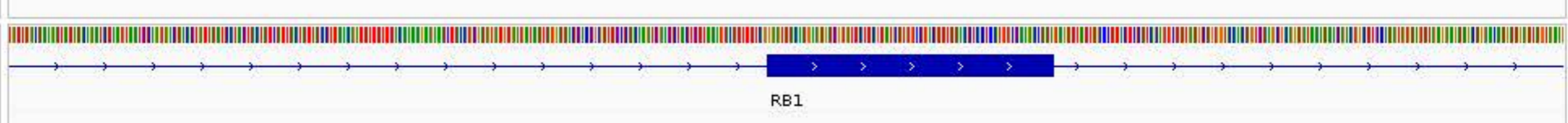
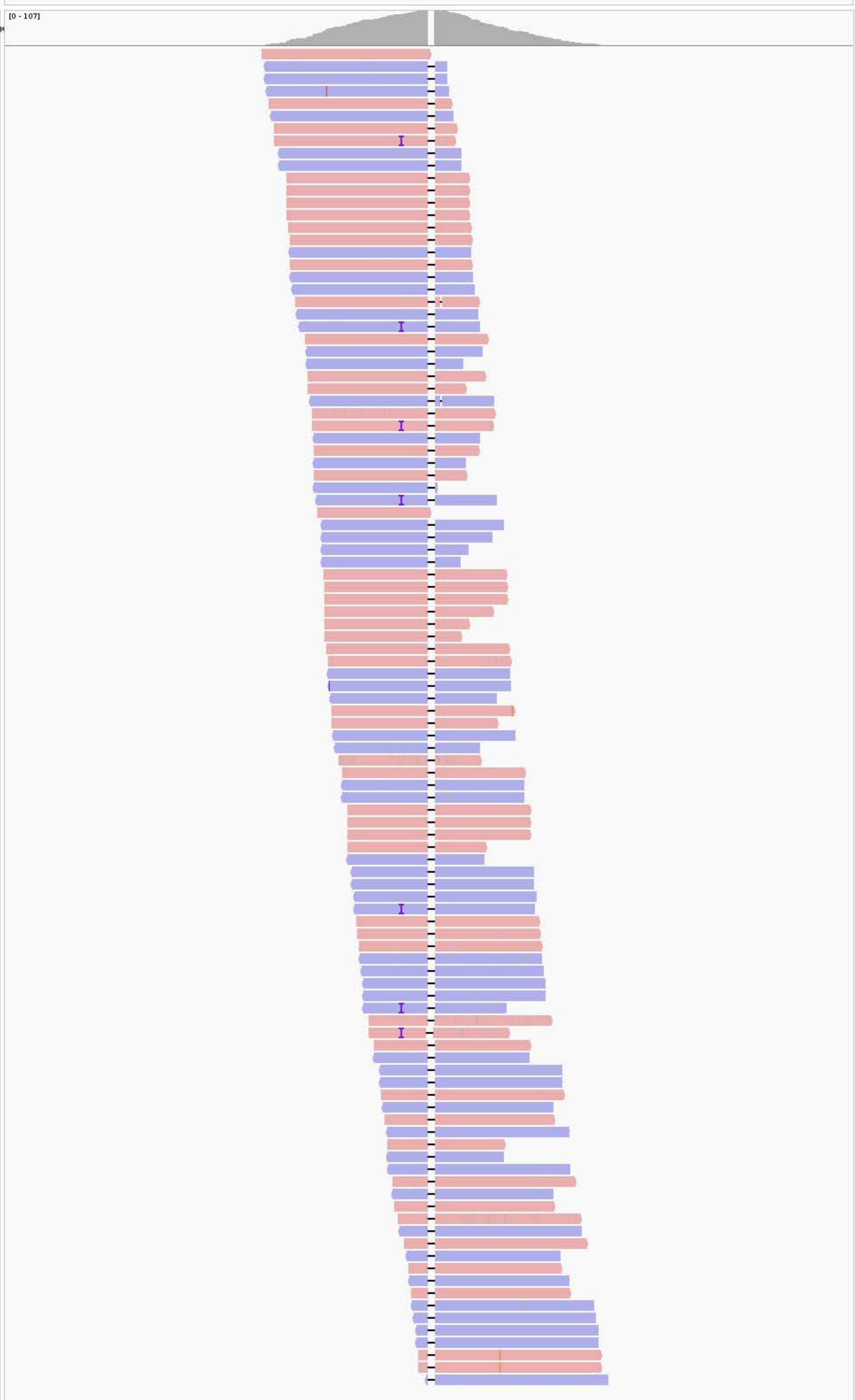


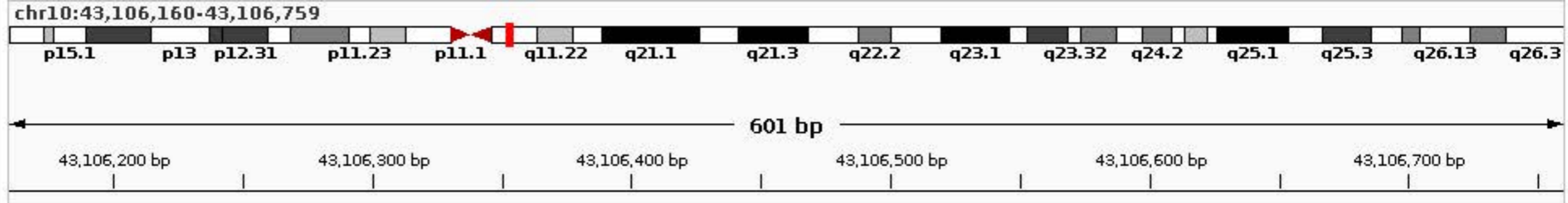
SL_1007-T2-B_mut.bam Coverage

SL_1007-T2-B_mut.bam

Sequence →

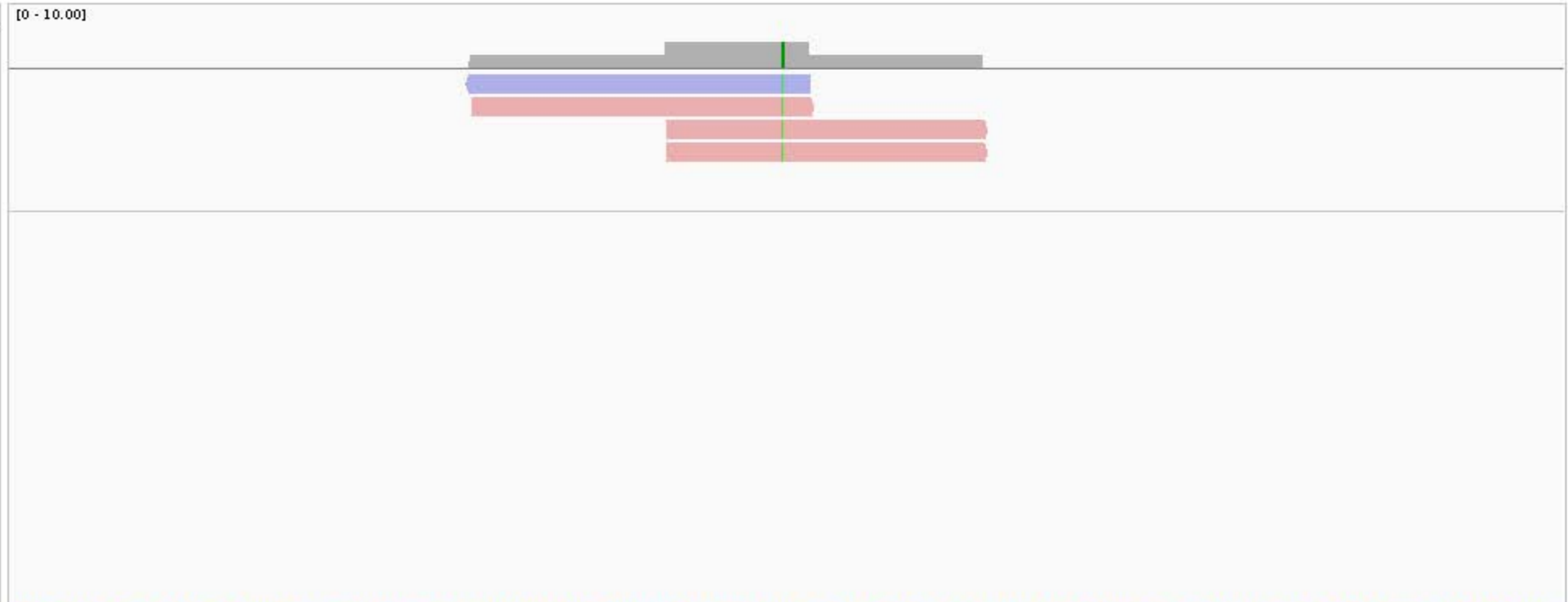
Refseq Genes





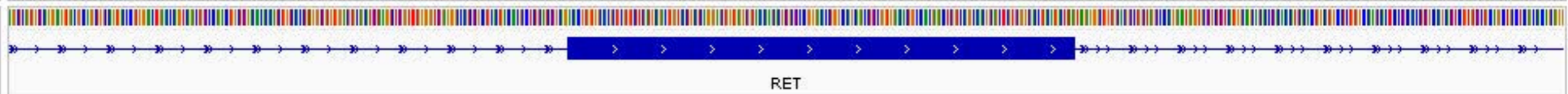
SL_0001_L_FFPE_10-B_mut.barr
rage

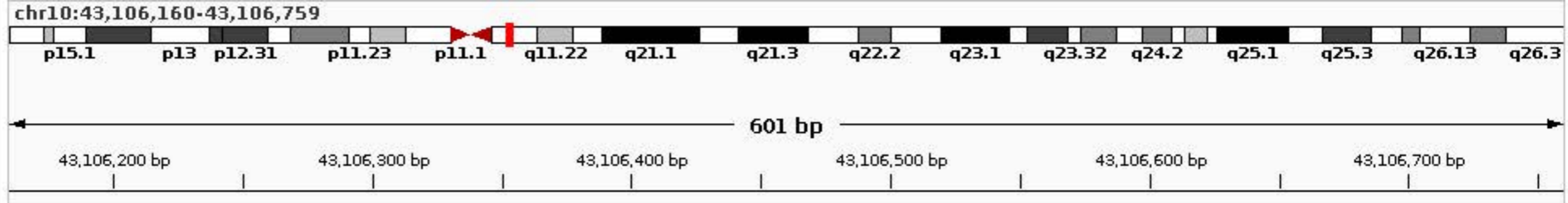
SL_0001_L_FFPE_10-B_mut.ba
m



Sequence →

Refseq Genes

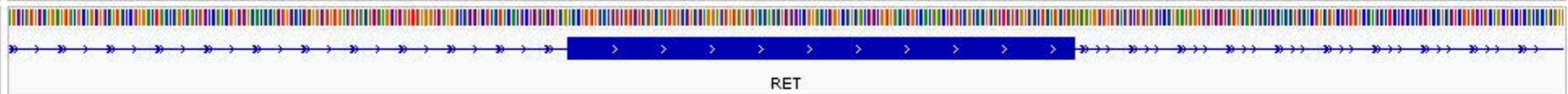




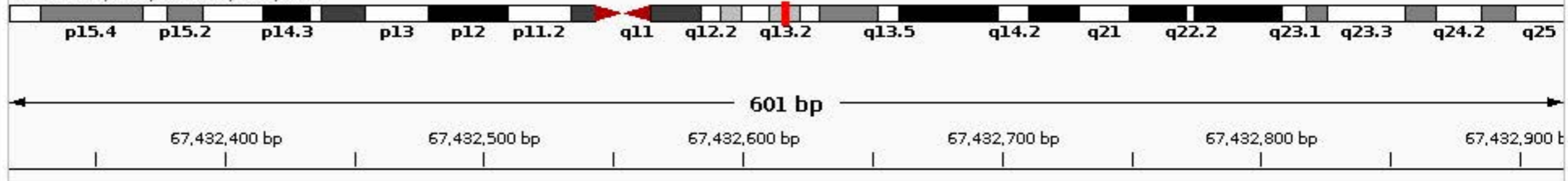
SL_0001_L_FFPE_29-B_mut.bam
SL_0001_L_FFPE_29-B_mut.bam



Sequence
Refseq Genes



chr11:67,432,317-67,432,916



SL_1006_L_FFPE_7-B_mut.bam
age

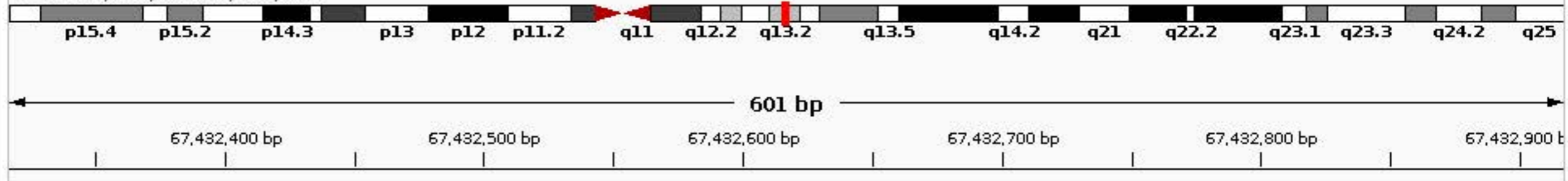


SL_1006_L_FFPE_7-B_mut.bam

Sequence →
Refseq Genes

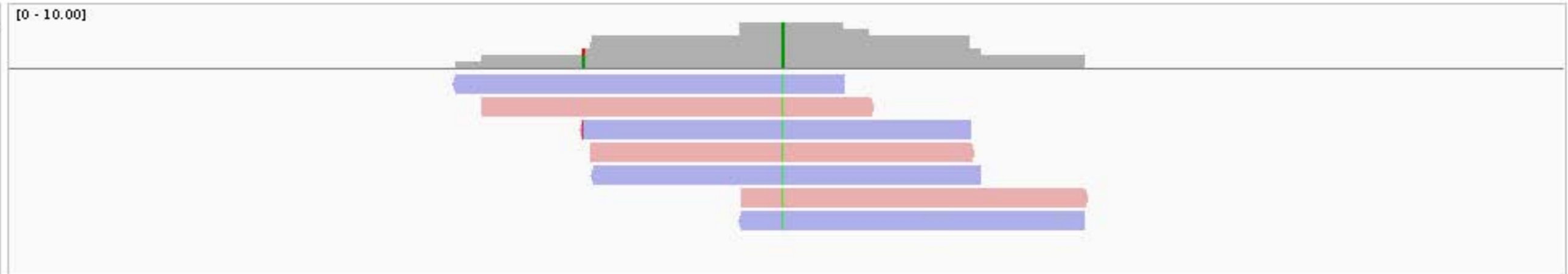


chr11:67,432,317-67,432,916



SL_1006_L_FFPE_28-B_mut.barr
rage

SL_1006_L_FFPE_28-B_mut.ba
m

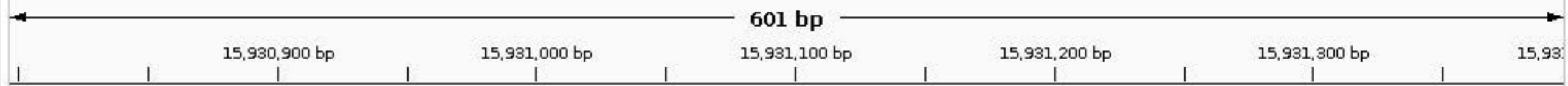


Sequence →

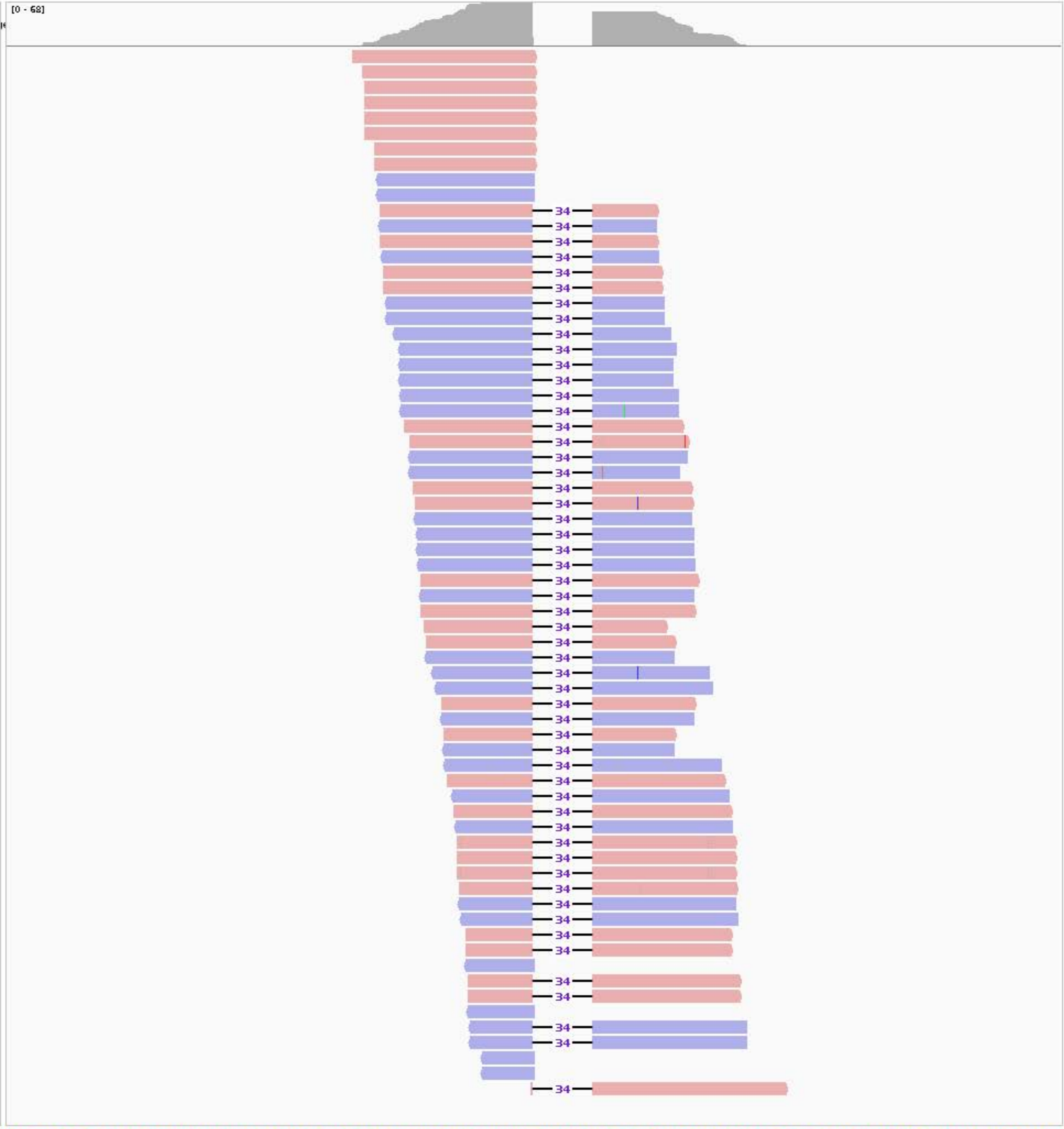
Refseq Genes



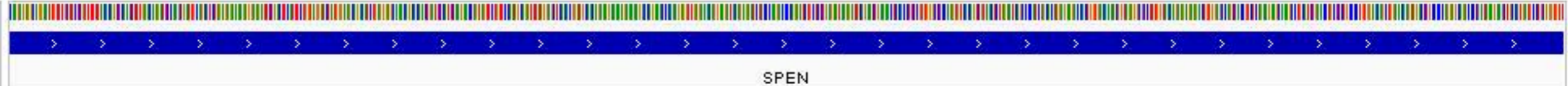
chr1:15,930,797-15,931,396
p36.23 p36.12 p34.3 p33 p32.1 p31.1 p22.2 p21.1 p13.1 q12 q21.1 q22 q24.1 q25.2 q31.2 q32.2 q41 q42.2 q44

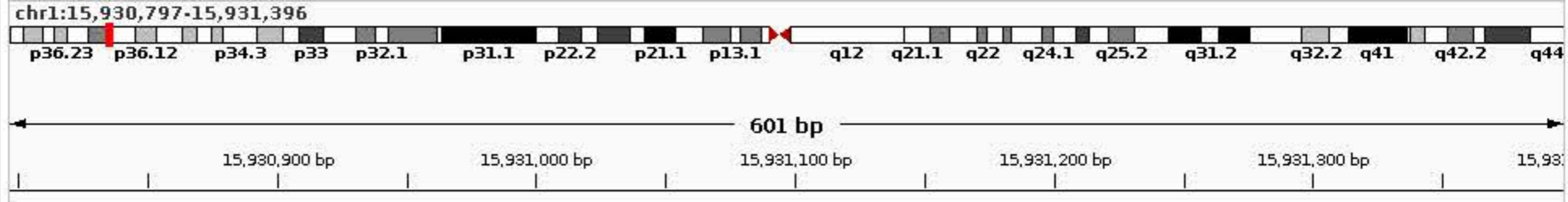


SL_1011-T2-B_mut.bam Coverage



Sequence →
Refseq Genes



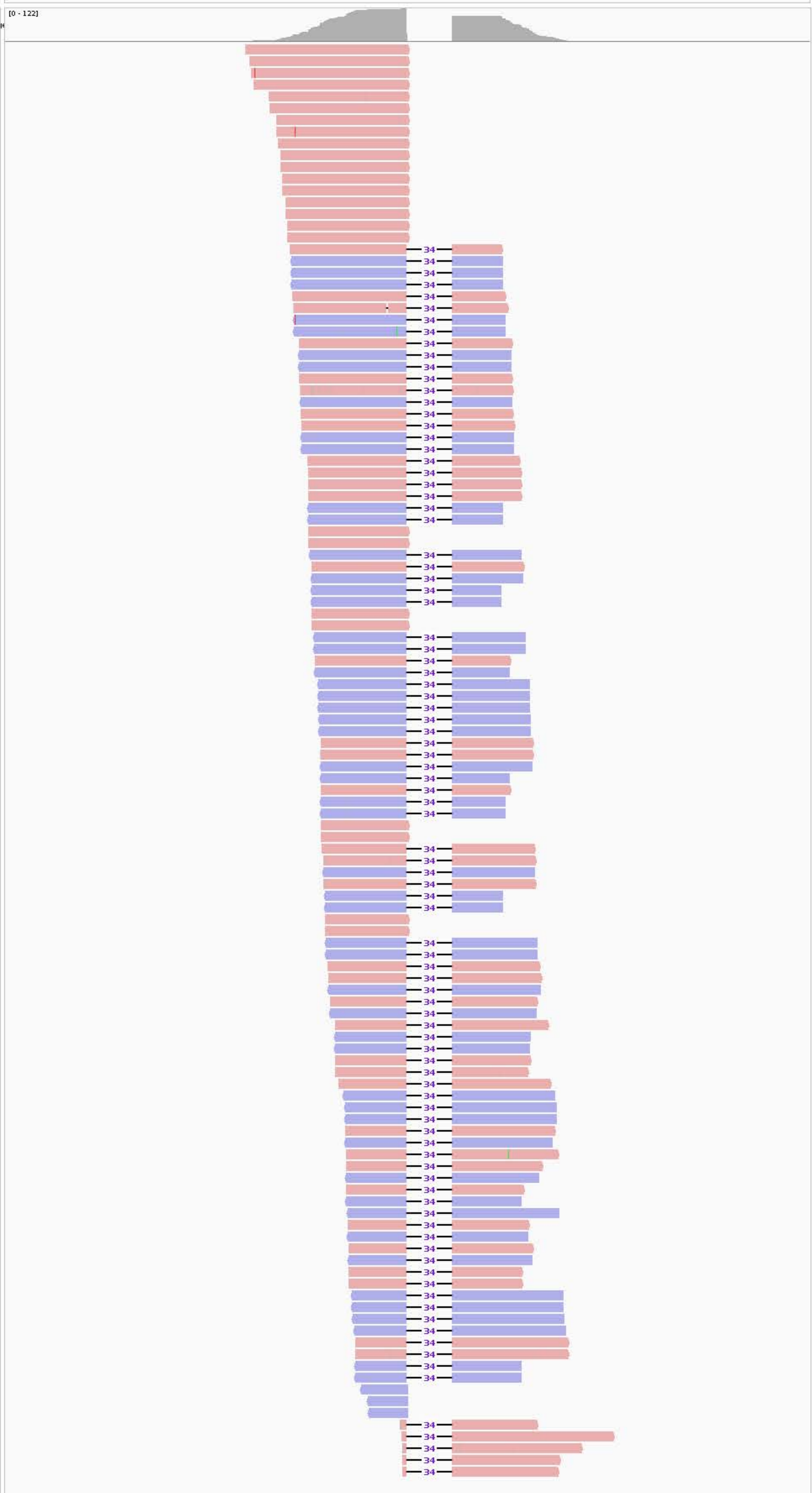


SL_1011-T3-B_mut.bam Coverage

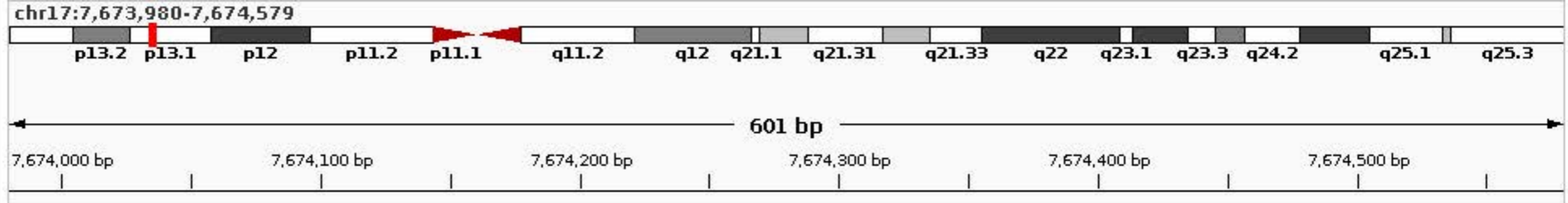
SL_1011-T3-B_mut.bam

Sequence

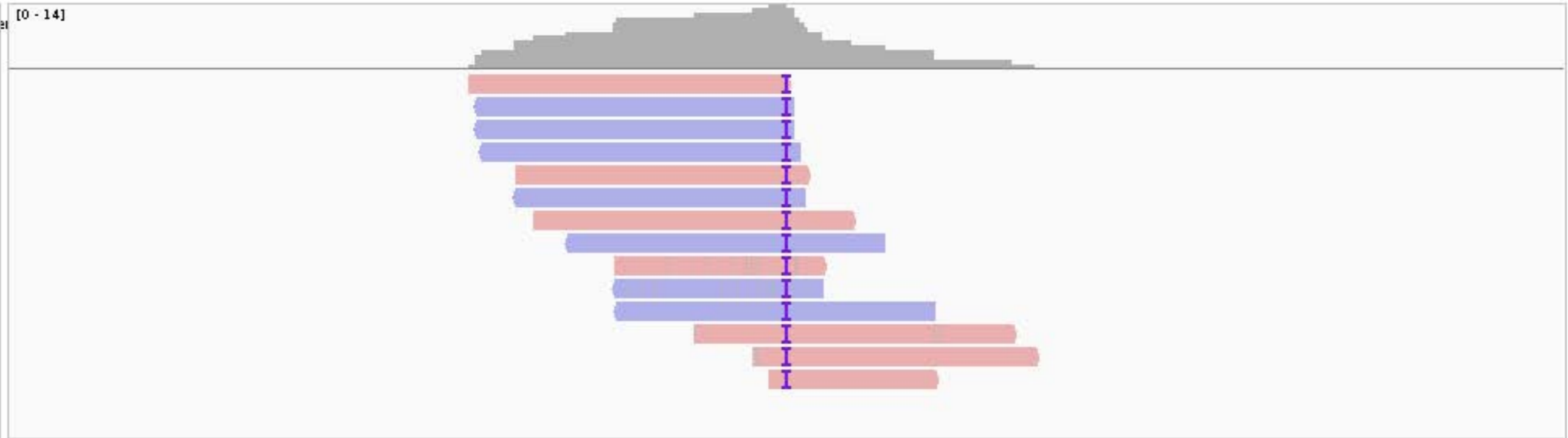
Refseq Genes



SPEN



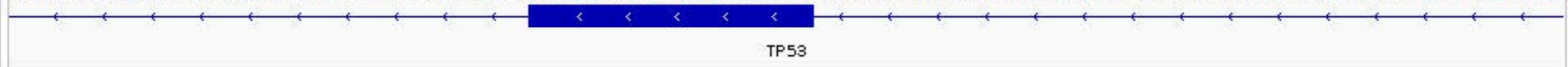
SL_0005-LT_11-B_mut.bam Cover



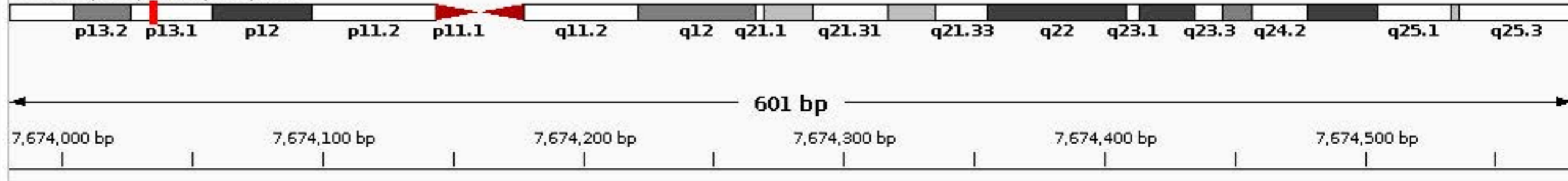
Sequence →



Refseq Genes



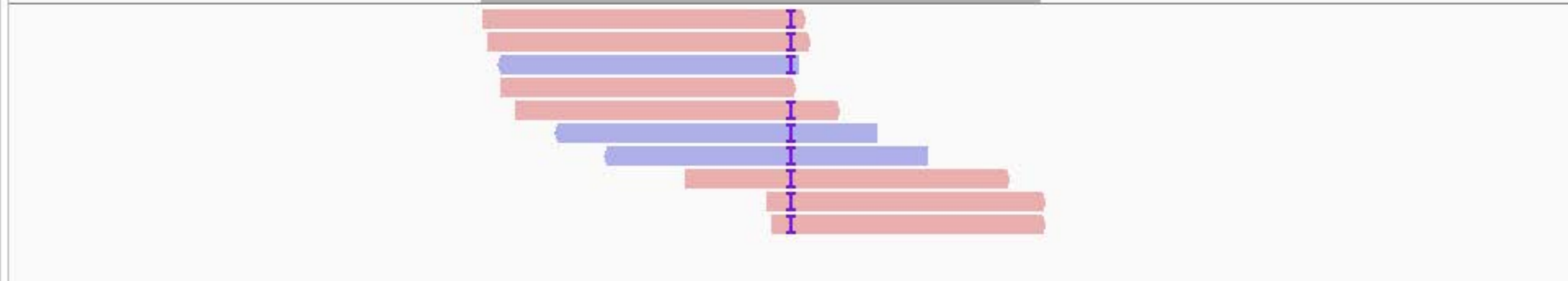
chr17:7,673,980-7,674,579



SL_0005-LT_12-B_mut.bam Cover



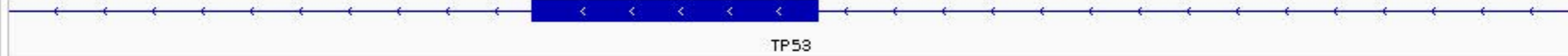
SL_0005-LT_12-B_mut.bam

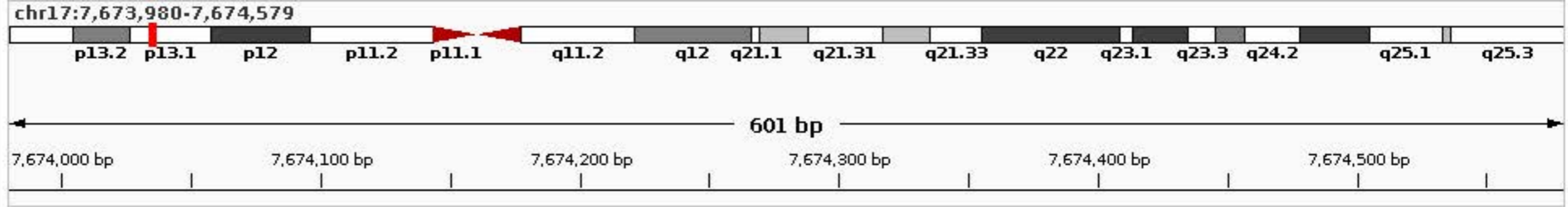


Sequence



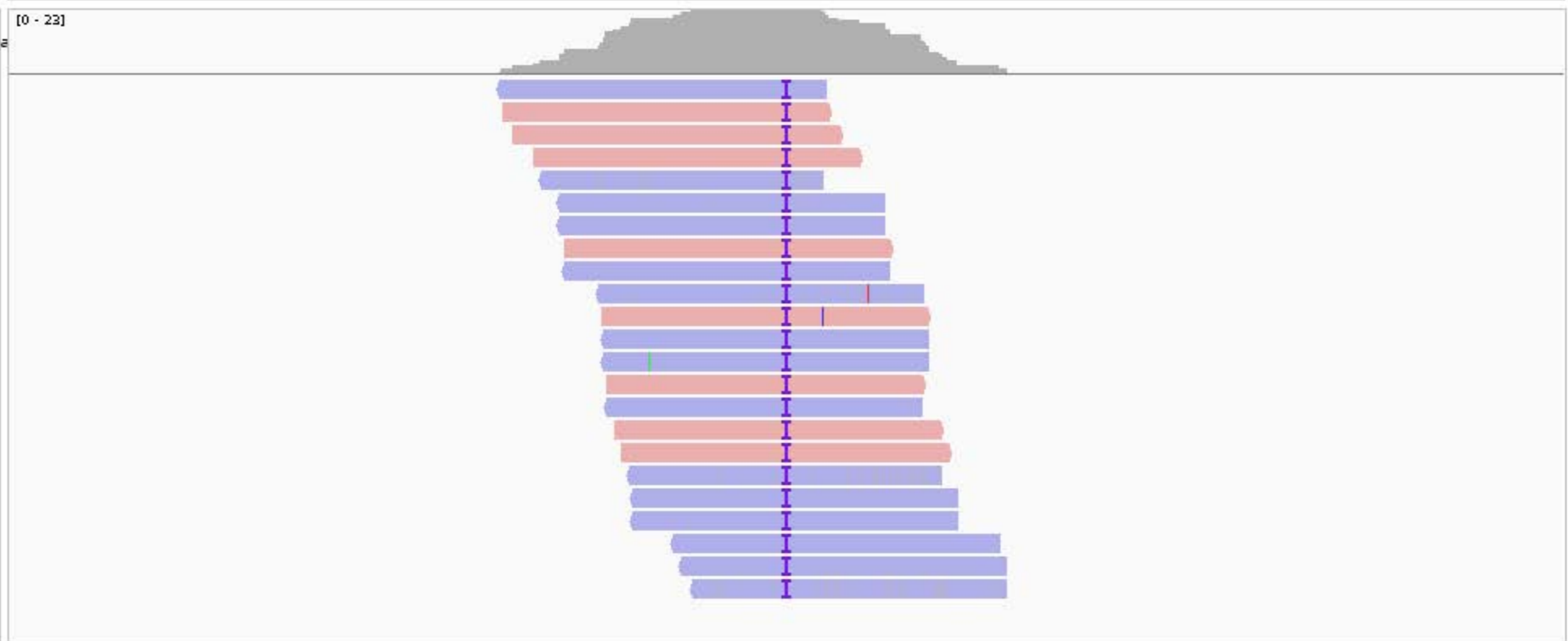
Refseq Genes





SL_0005-LT-1-B_mut.bam Coverage

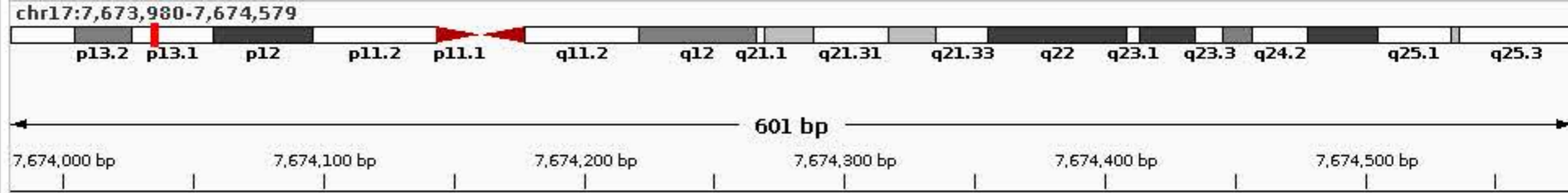
SL_0005-LT-1-B_mut.bam



Sequence →

Refseq Genes



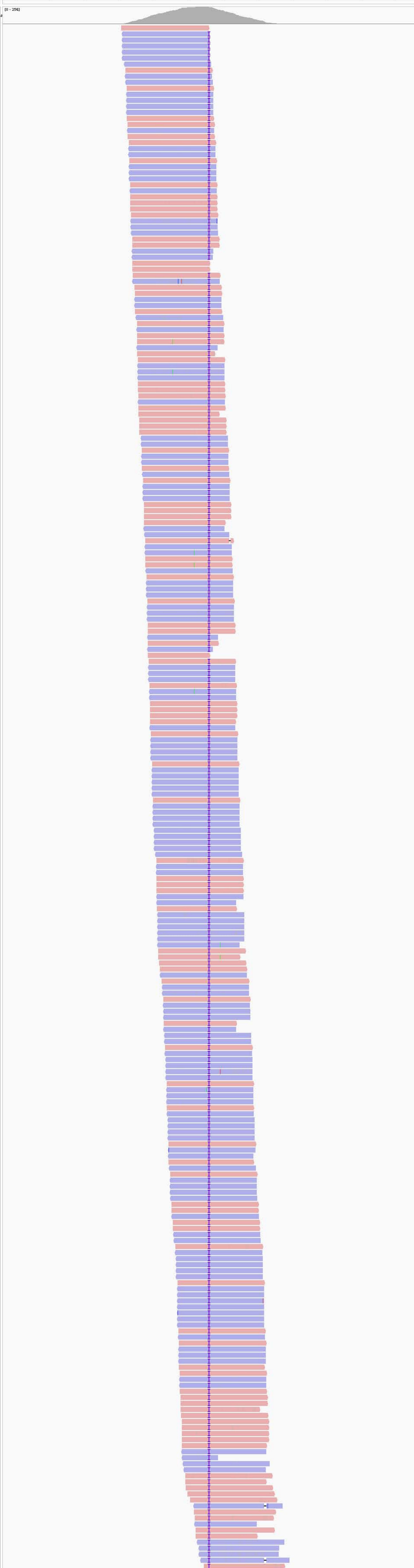


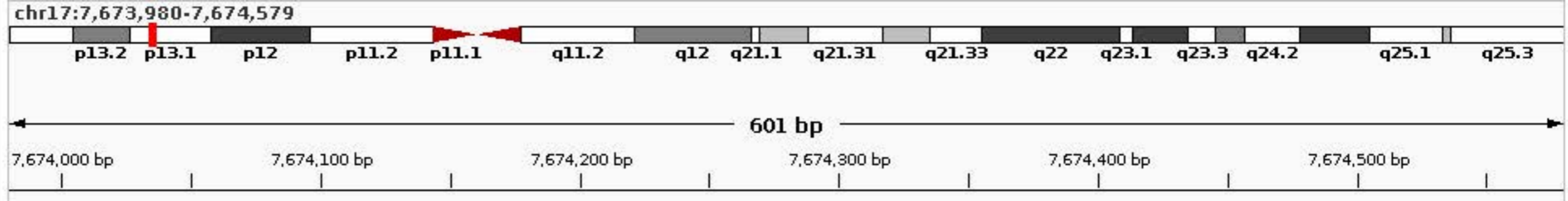
SL_0005-LT-4-B_mut.bam Coverage

SL_0005-LT-4-B_mut.bam

Sequence

Refseq Genes





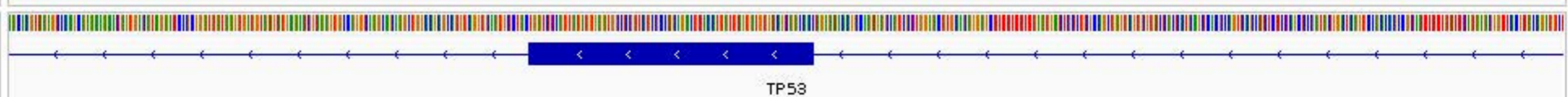
SL_0005-LT-9-B_mut.bam Coverage

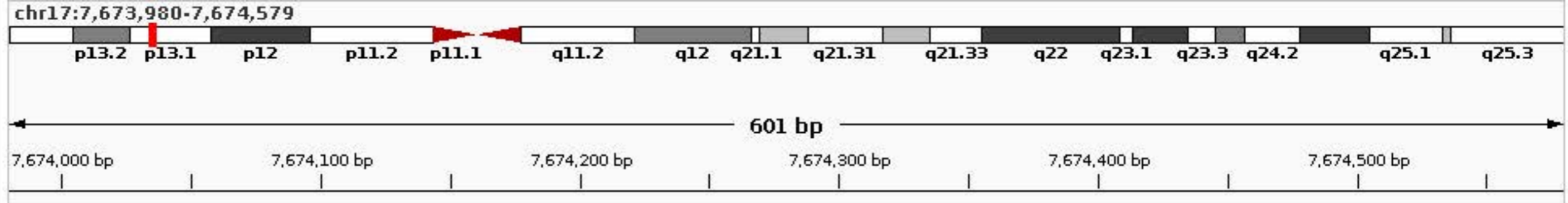
SL_0005-LT-9-B_mut.bam



Sequence

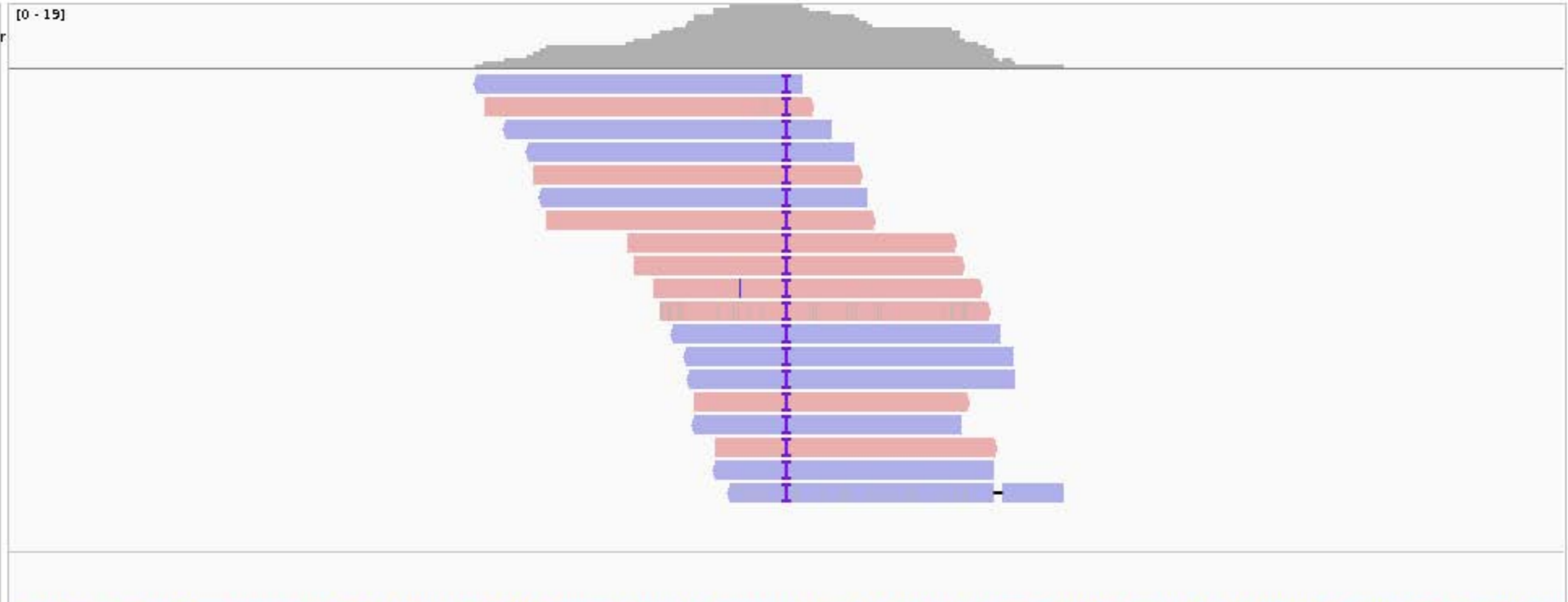
Refseq Genes





SL_0005-RN-1-B_mut.bam Cover

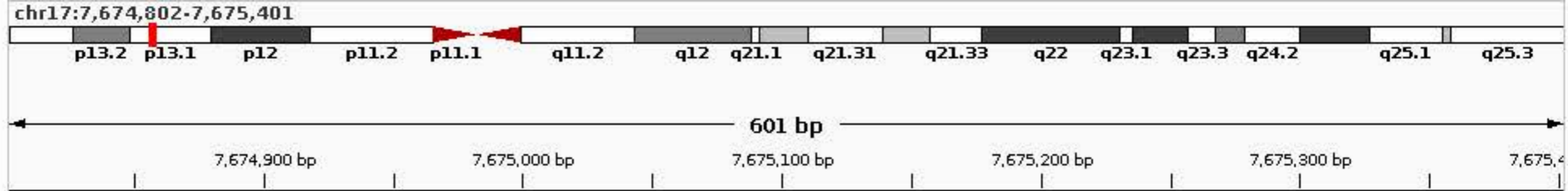
SL_0005-RN-1-B_mut.bam



Sequence →

Refseq Genes



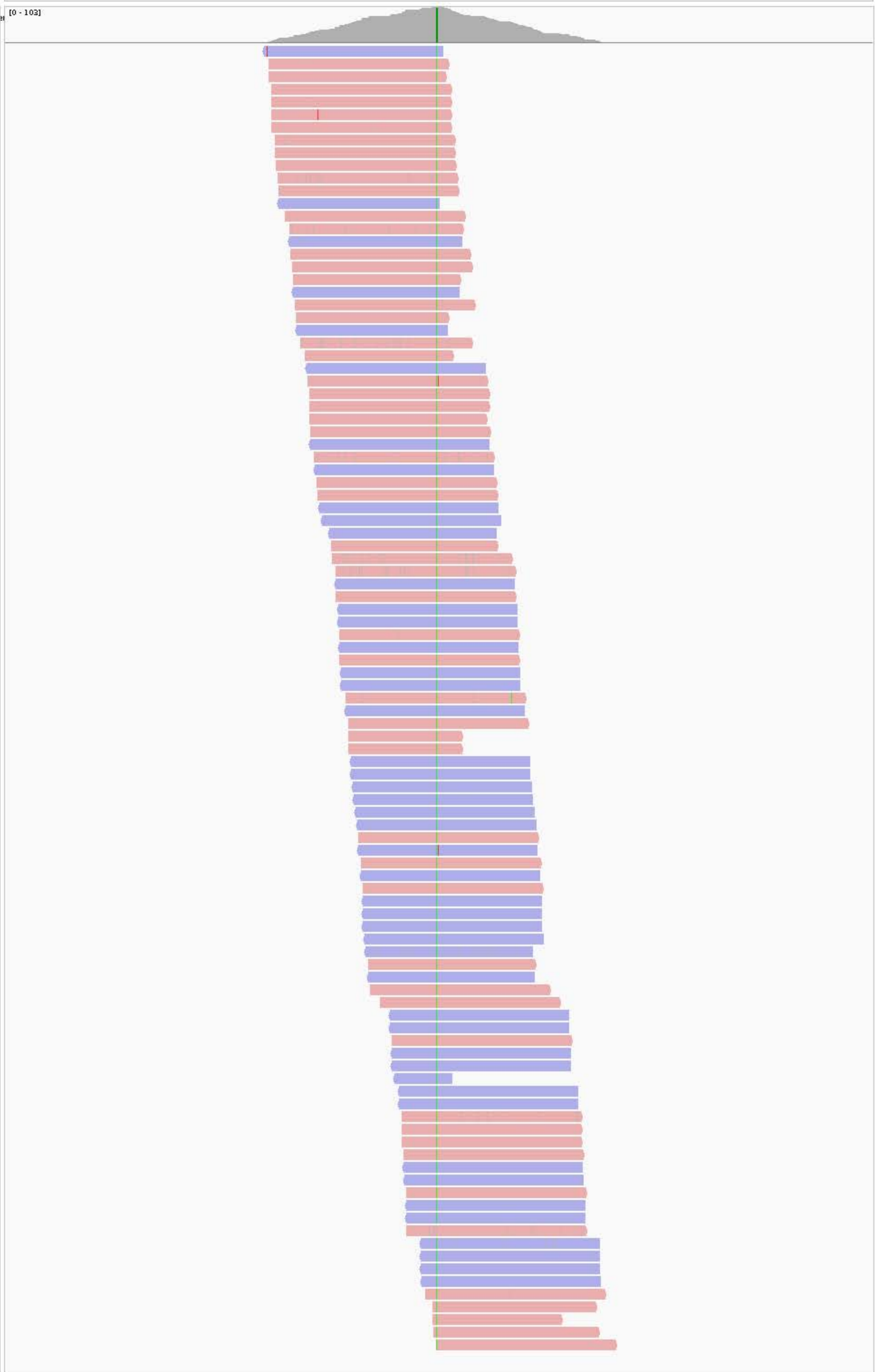


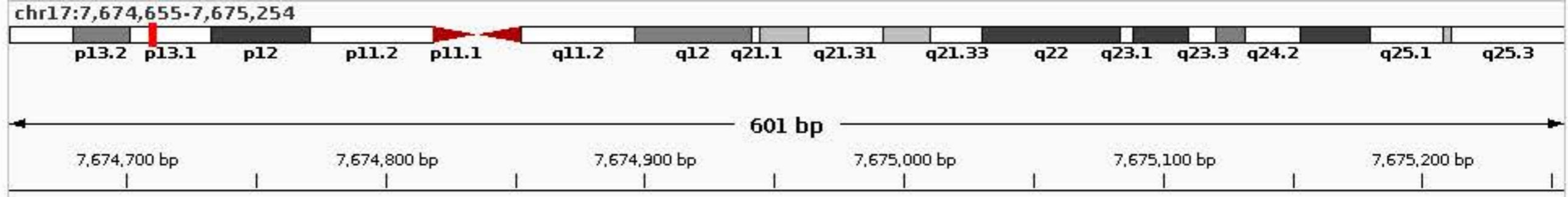
SL_0008-LT_15-B_mut.bam Coverage [0 - 103]

SL_0008-LT_15-B_mut.bam

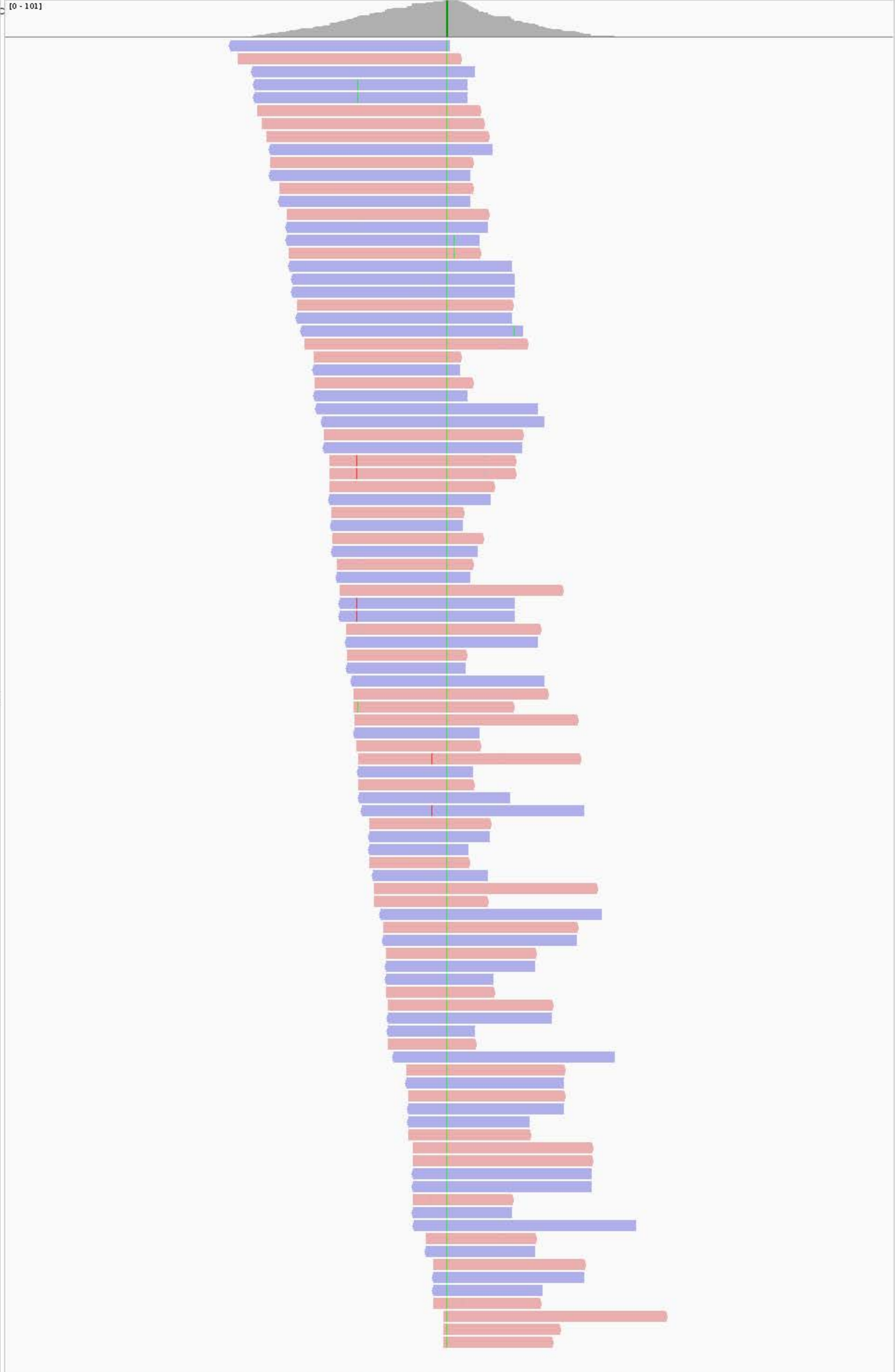
Sequence →

Refseq Genes





SL_1010_L_FFPE_3-B_mut.bam
age

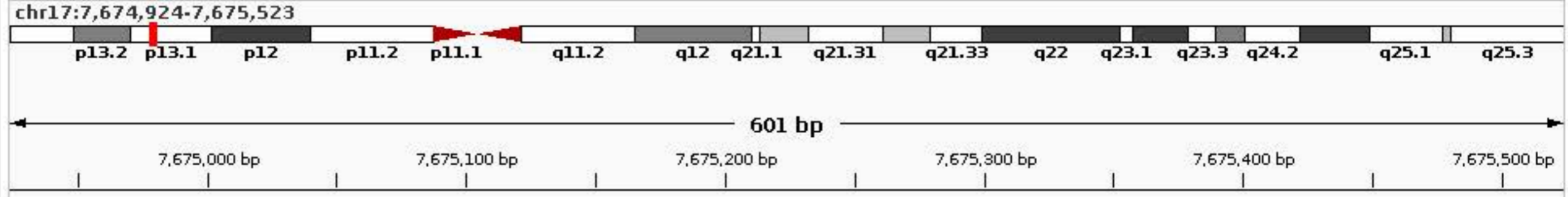


SL_1010_L_FFPE_3-B_mut.bam

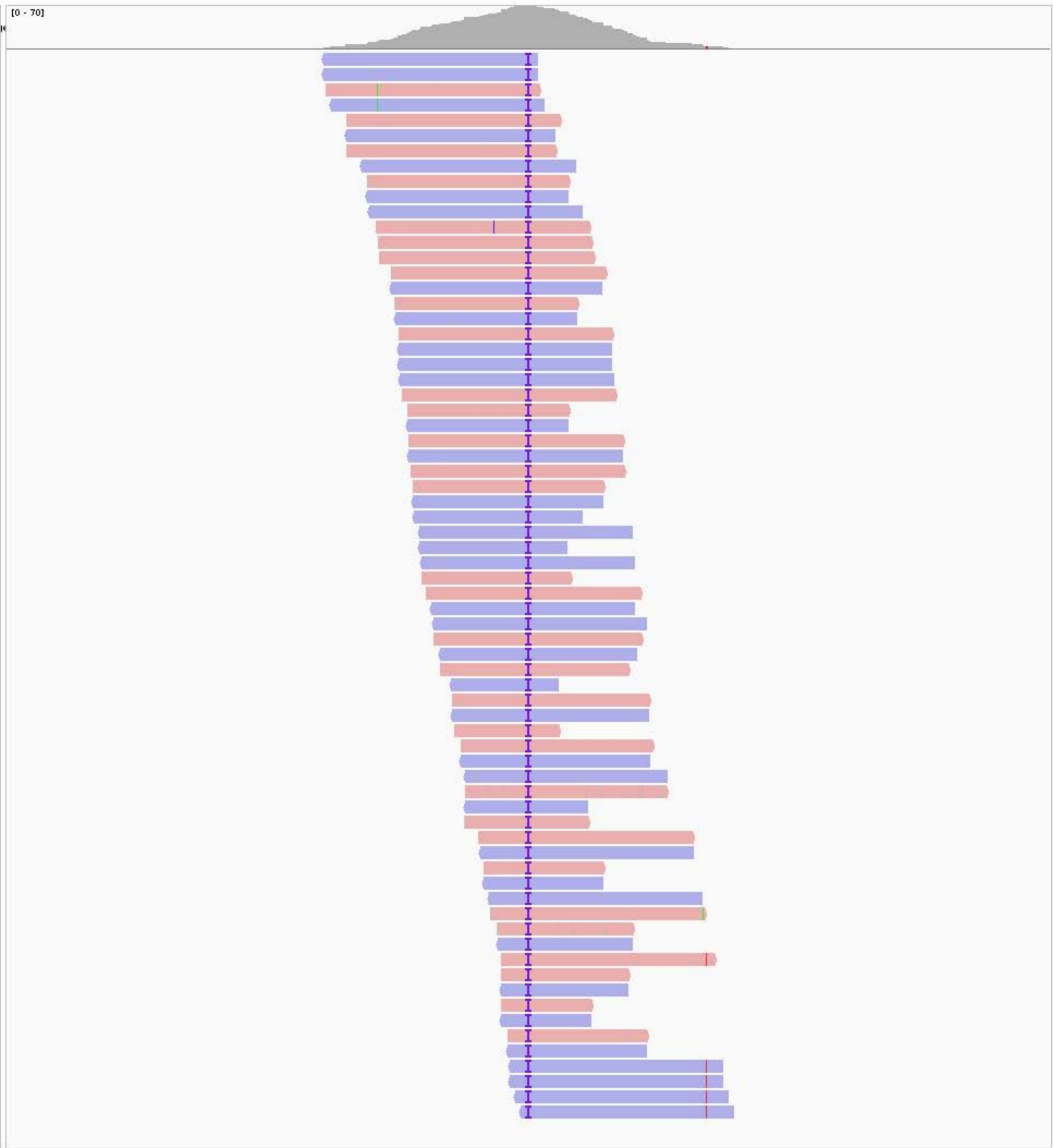
Sequence →
Refseq Genes



TP53



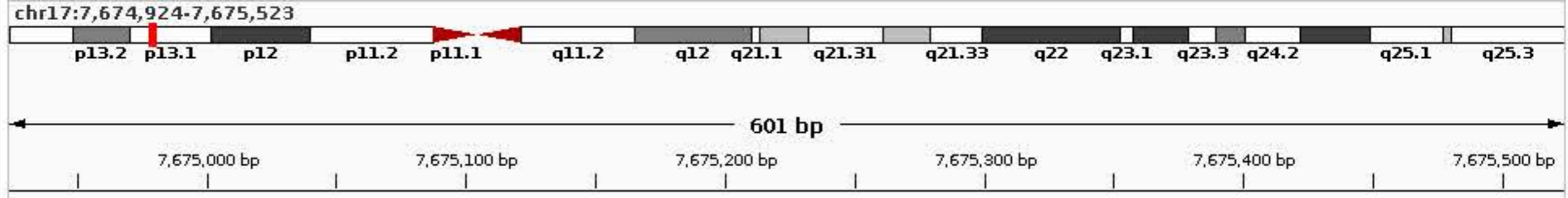
SL_1007-T1-B_mut.bam Coverage



SL_1007-T1-B_mut.bam

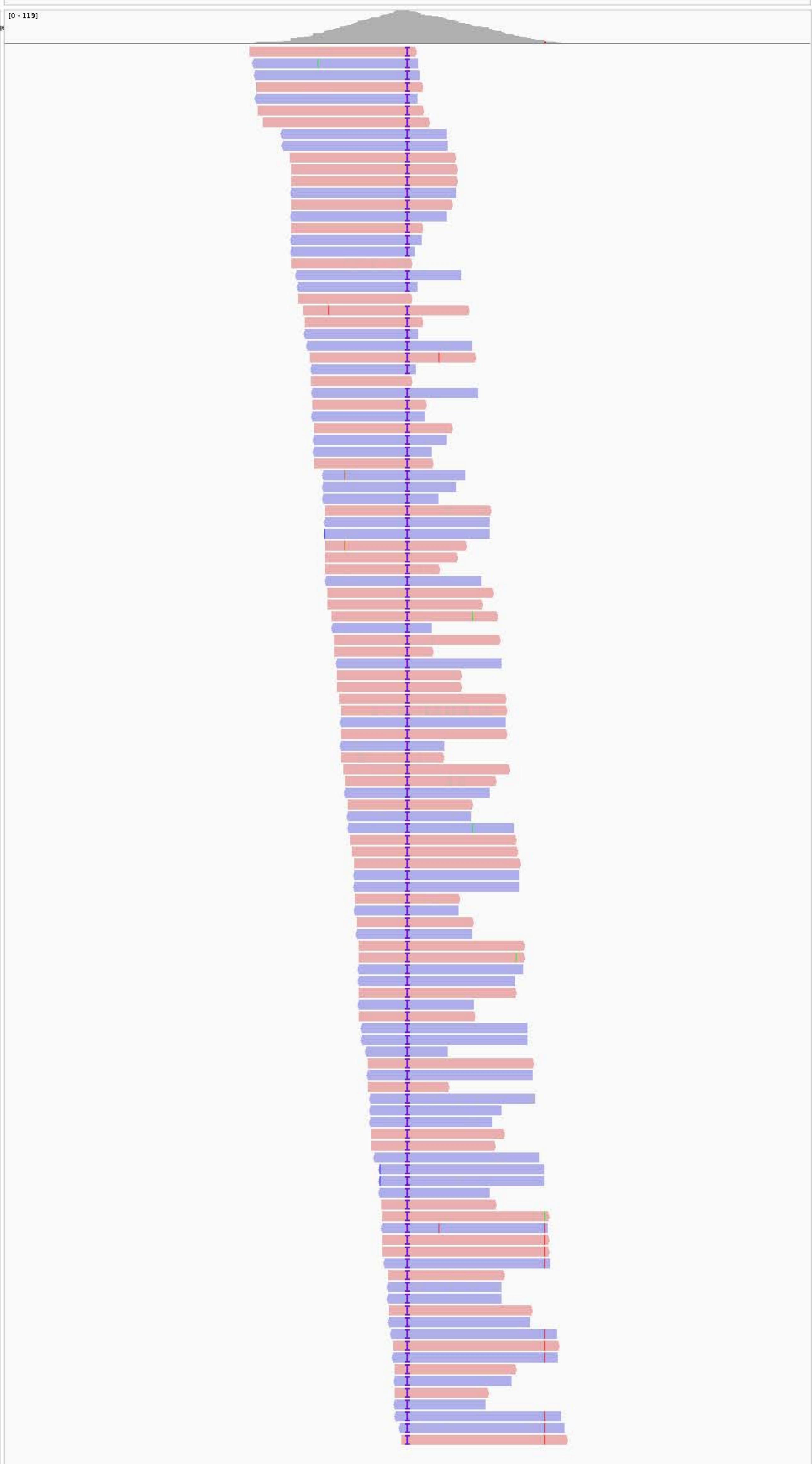
Sequence →
Refseq Genes





SL_1007-T2-B_mut.bam Coverage

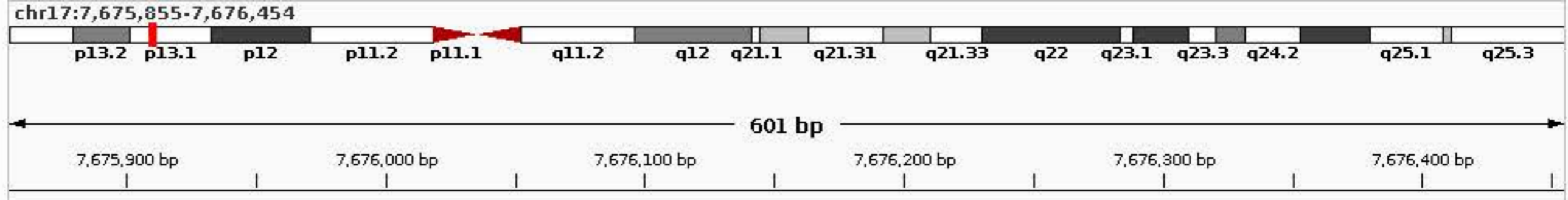
SL_1007-T2-B_mut.bam



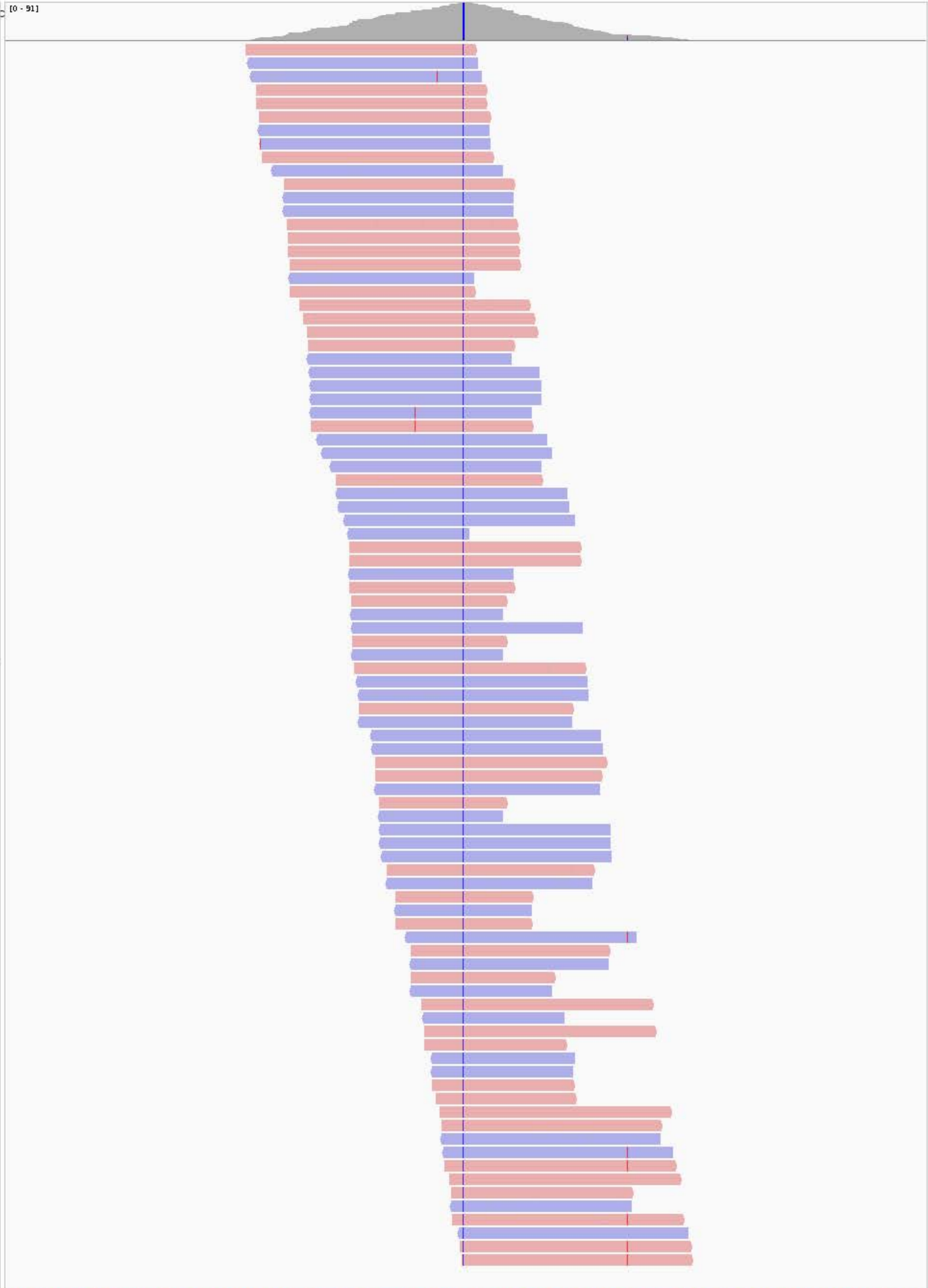
Sequence

Refseq Genes





SL_0011_L_FFPE_4-B_mut.bam
age

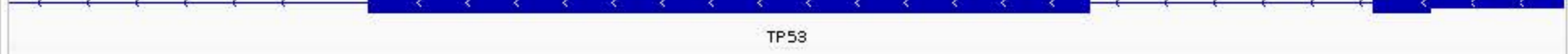


SL_0011_L_FFPE_4-B_mut.bam

Sequence →



Refseq Genes

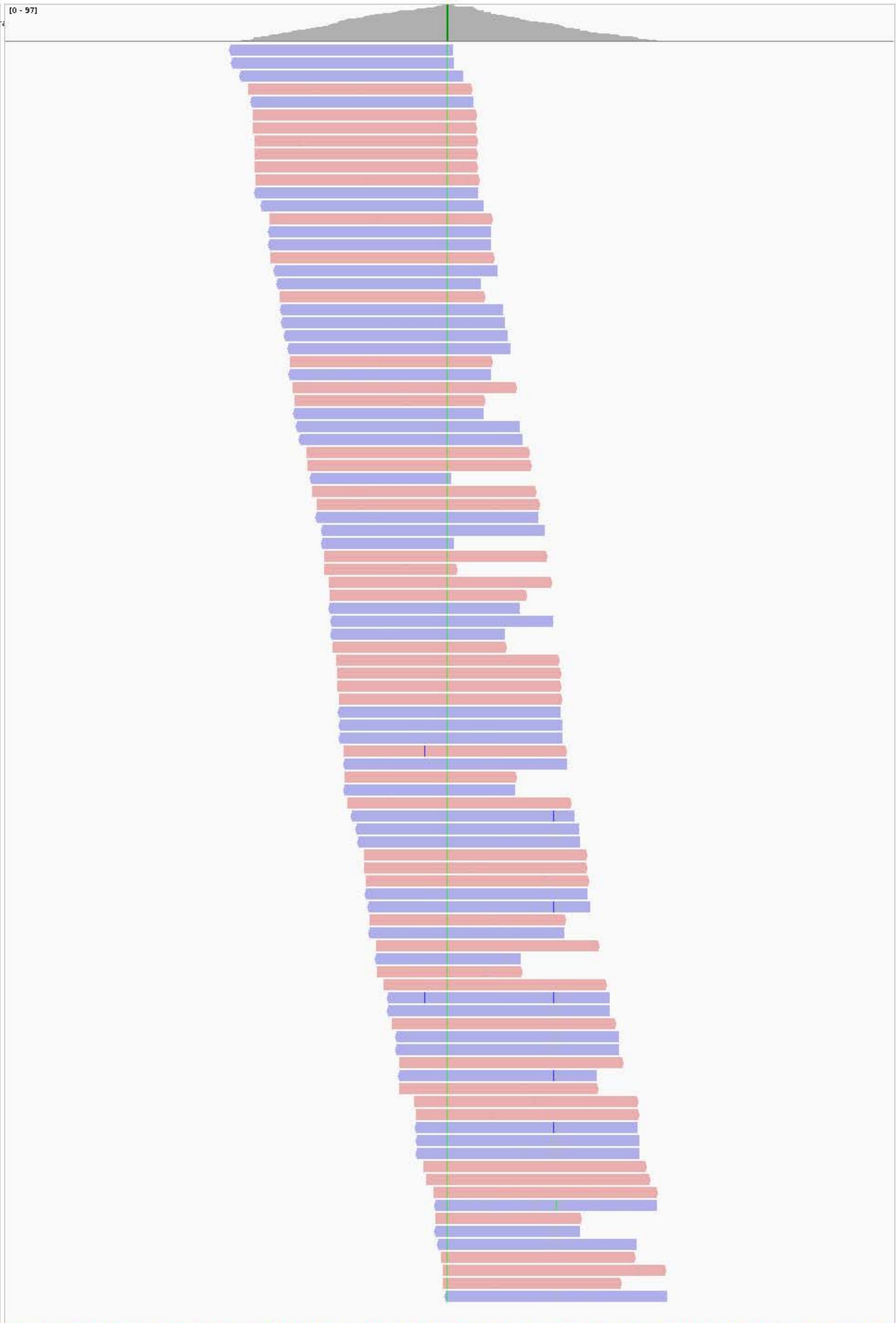


TP53

chr6:80,007,621-80,008,220
p25.1 p23 p22.2 p21.31 p21.1 p12.2 q11.1 q13 q14.1 q15 q16.2 q21 q22.1 q22.32 q23.3 q24.3 q25.3 q27



SL_0008_LT_8-B_mut.bam Coverage



SL_0008_LT_8-B_mut.bam

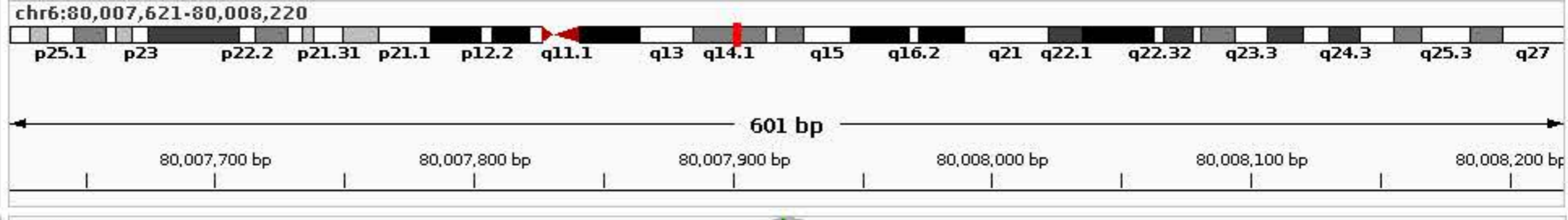
Sequence



Refseq Genes



TTK

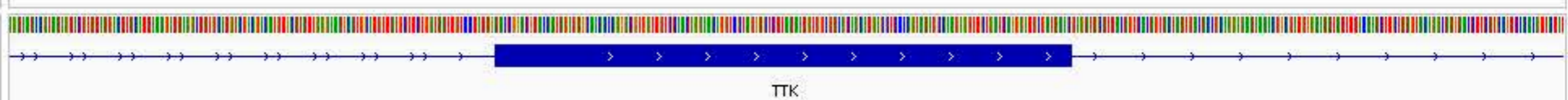
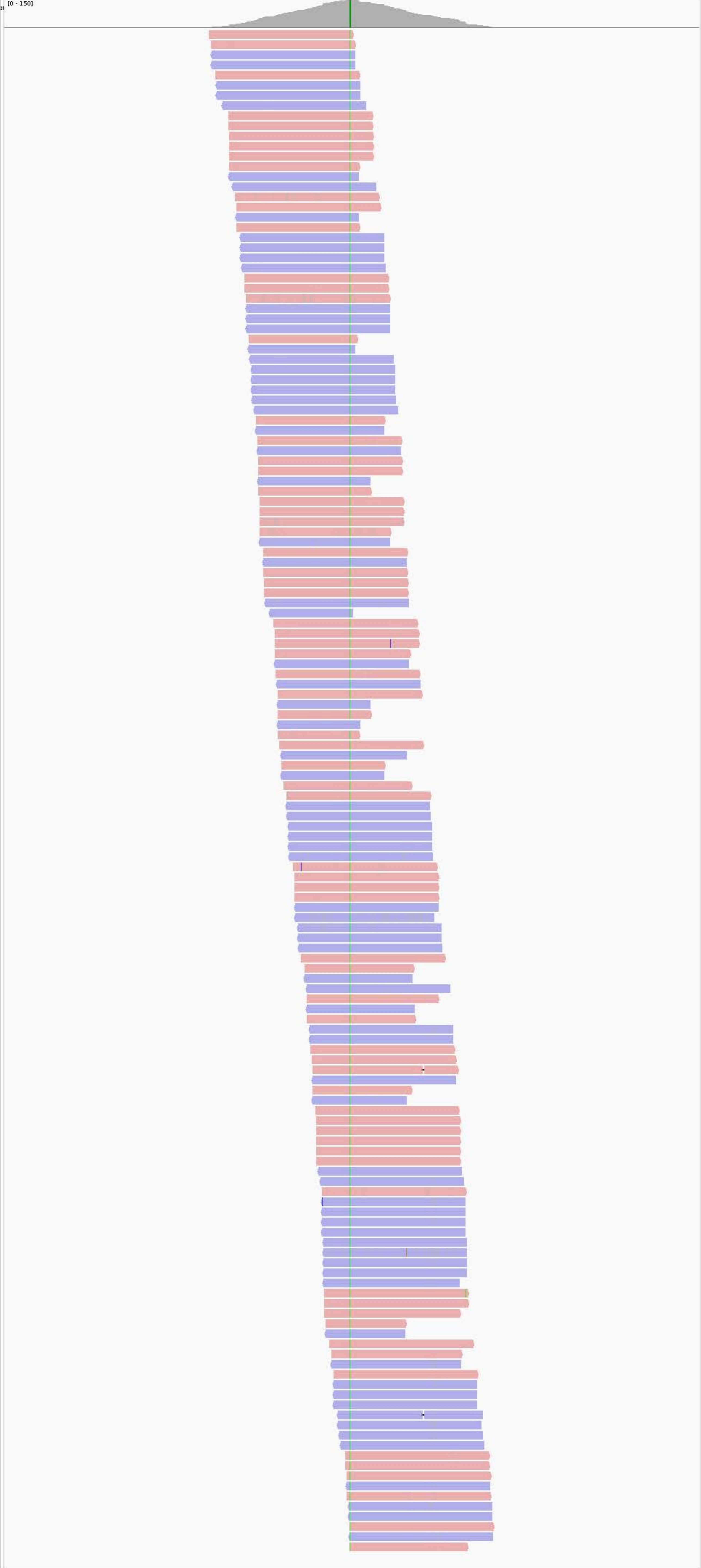


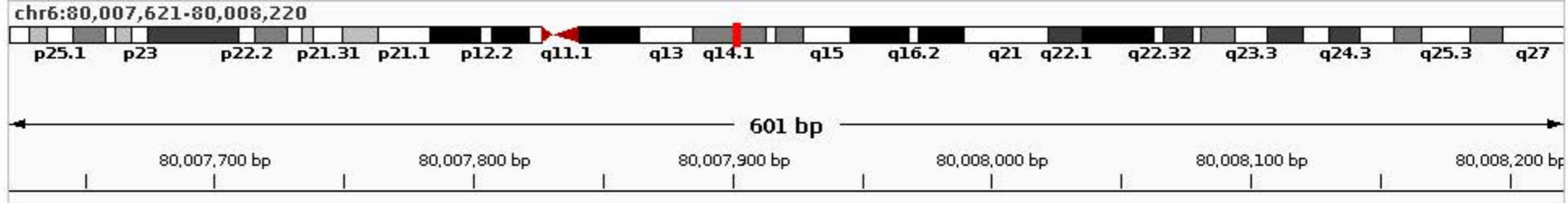
SL_0008-LT_15-8_mut.bam Cov

SL_0008-LT_15-8_mut.bam

Sequence

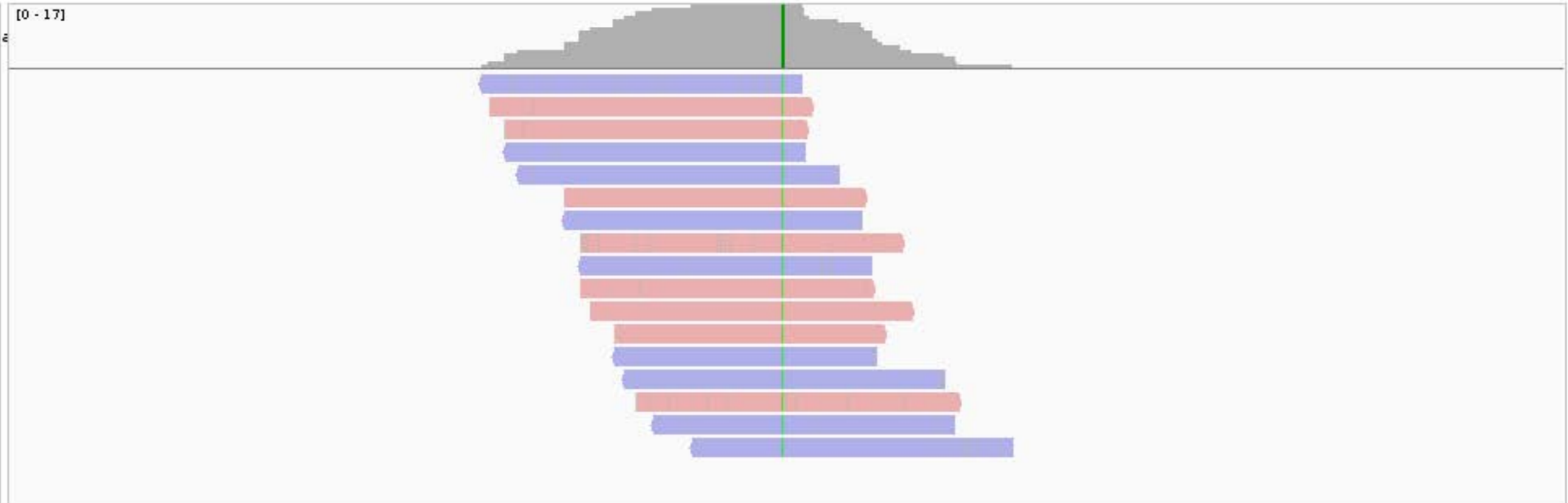
Refseq Genes





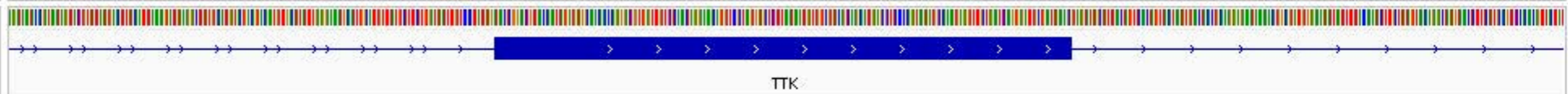
SL_0009-LT_9-B_mut.bam Coverage

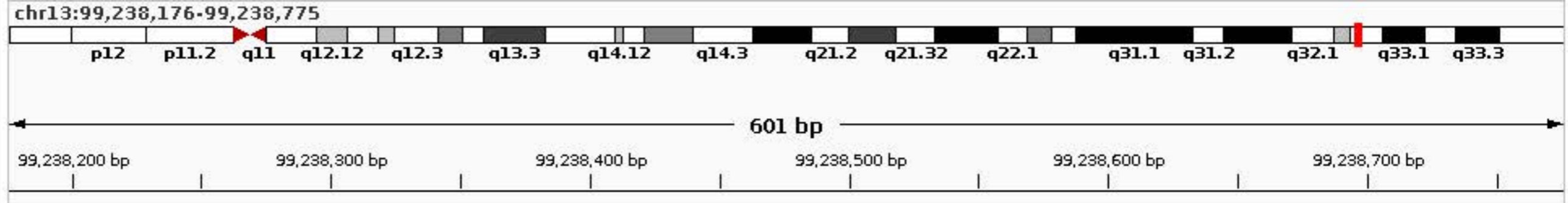
SL_0009-LT_9-B_mut.bam



Sequence →

Refseq Genes





SL_0004_R_FFPE_45-B_mut.bam
rage

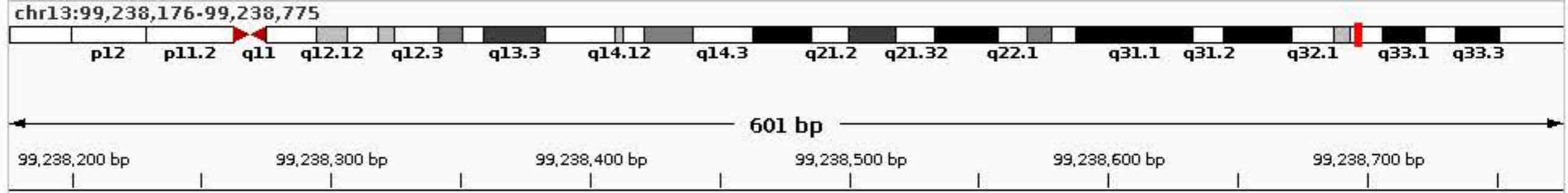
SL_0004_R_FFPE_45-B_mut.ba
m



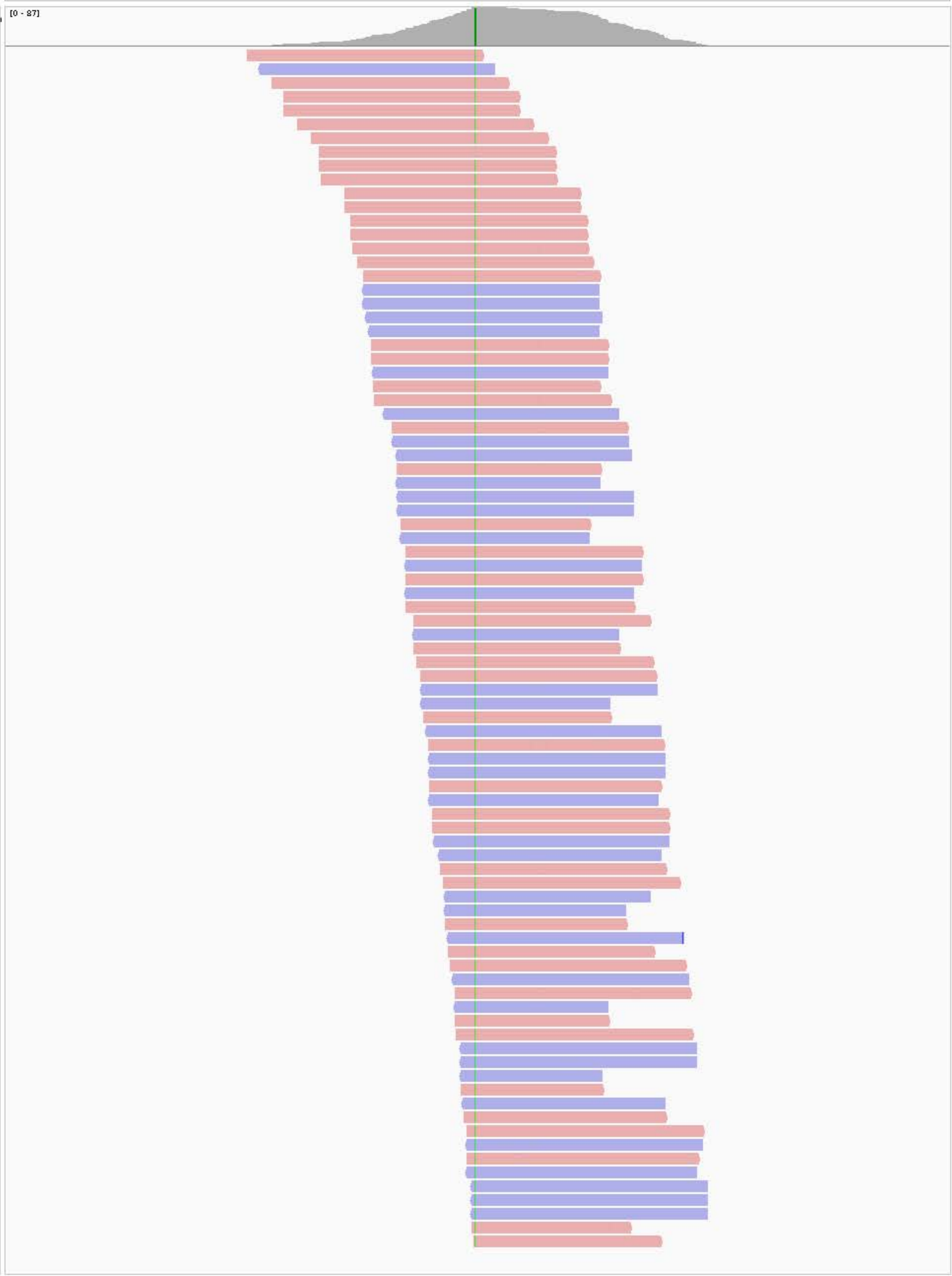
Sequence →

Refseq Genes





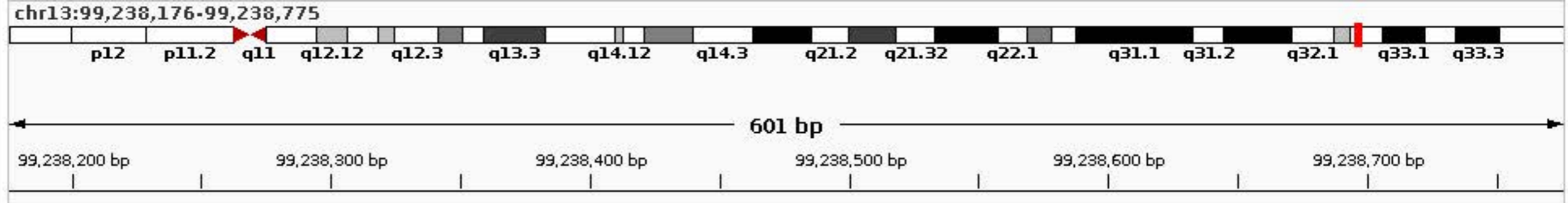
SL_0004_R_FFPE_50-B_mut.bam
 range



SL_0004_R_FFPE_50-B_mut.ba
 m

Sequence →
 Refseq Genes





SL_0004-RN-3-B_mut.bam Cover

SL_0004-RN-3-B_mut.bam

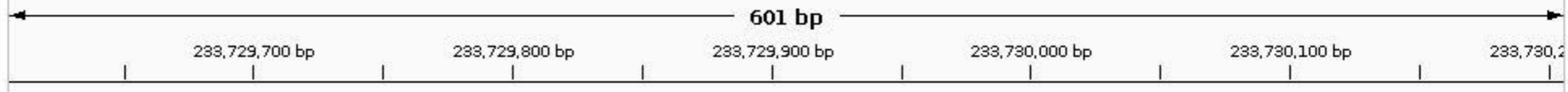
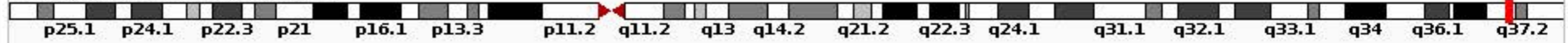


Sequence →

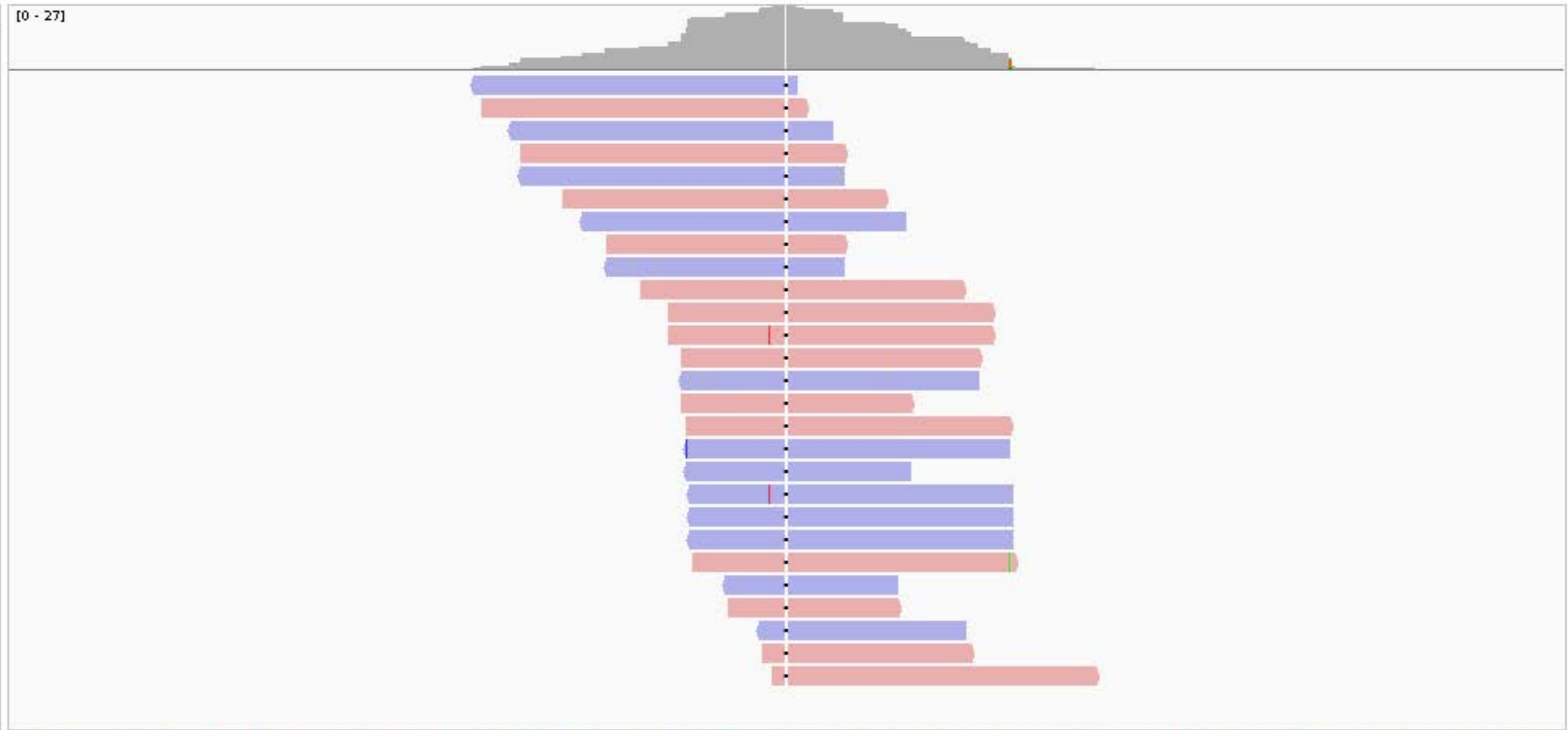
Refseq Genes



chr2:233,729,606-233,730,205



SL_1006_L_35_FFPE-B_mut.bam
range



SL_1006_L_35_FFPE-B_mut.bam

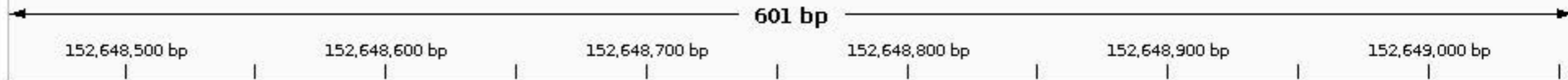
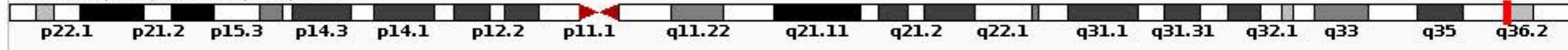
Sequence →



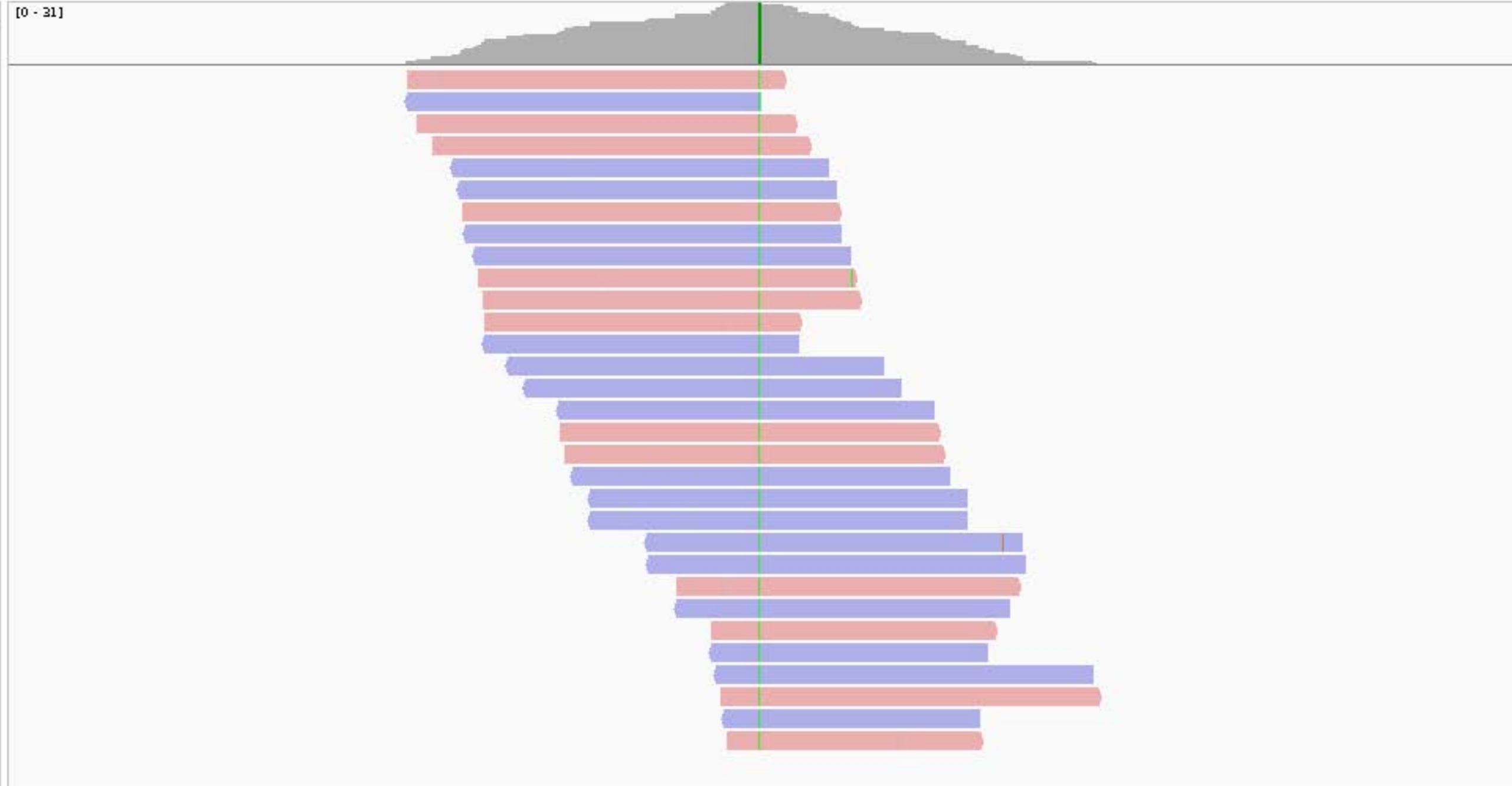
Refseq Genes

UGT1A8

chr7:152,648,456-152,649,055



SL_1012_L_FFPE_36-B_mut.barrage



SL_1012_L_FFPE_36-B_mut.bam

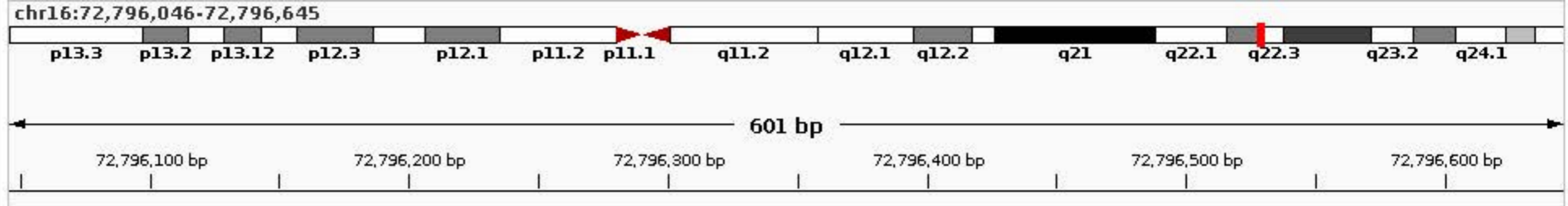
Sequence



Refseq Genes

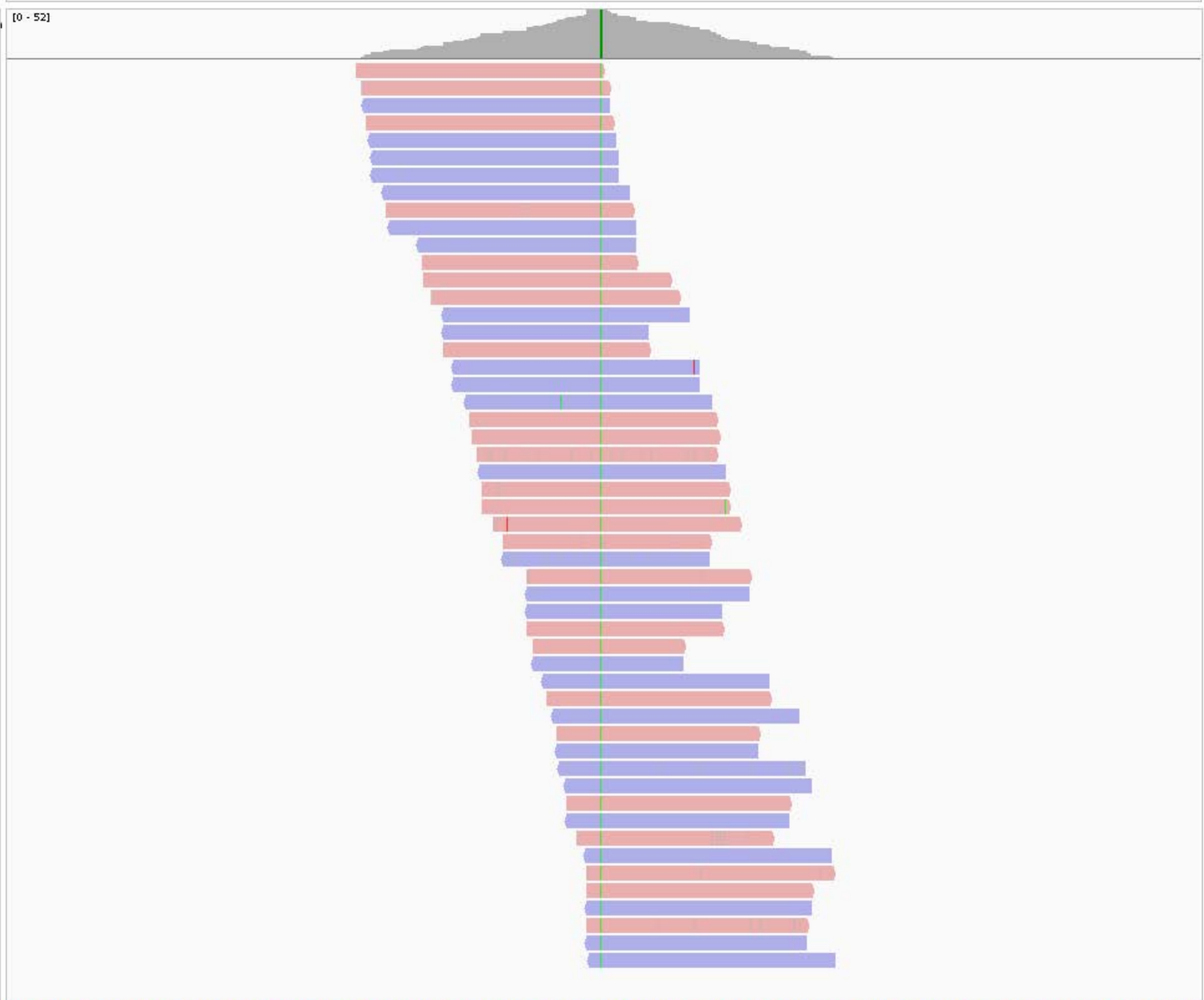


XRC2



SL_1012_R_10_FFPE-B_mut.bam
rage

SL_1012_R_10_FFPE-B_mut.ba
m



Sequence →

Refseq Genes

