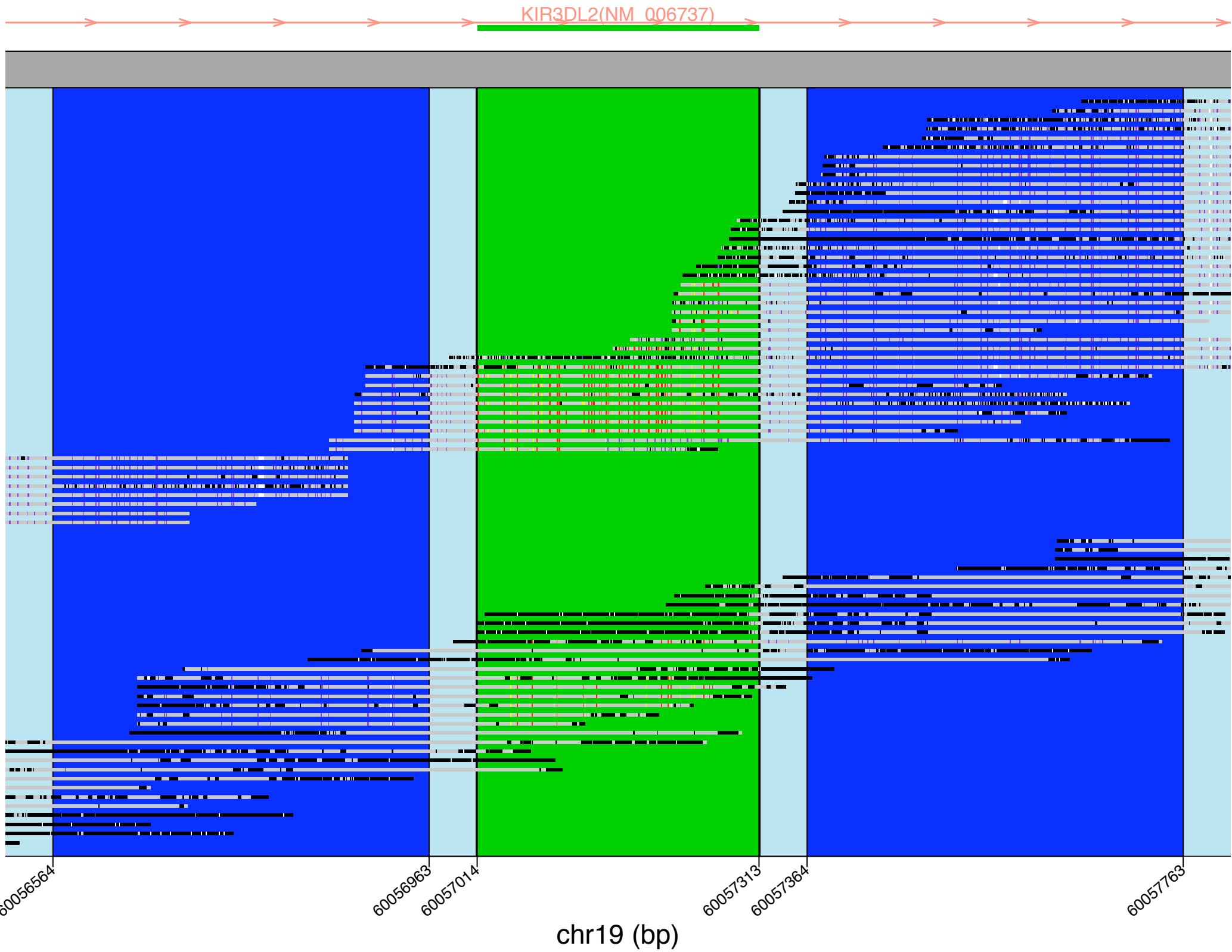
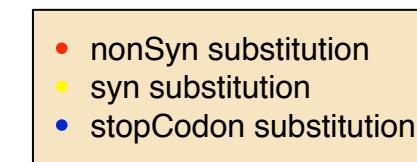
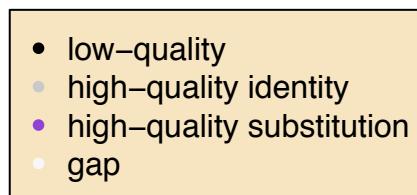
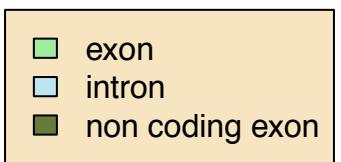
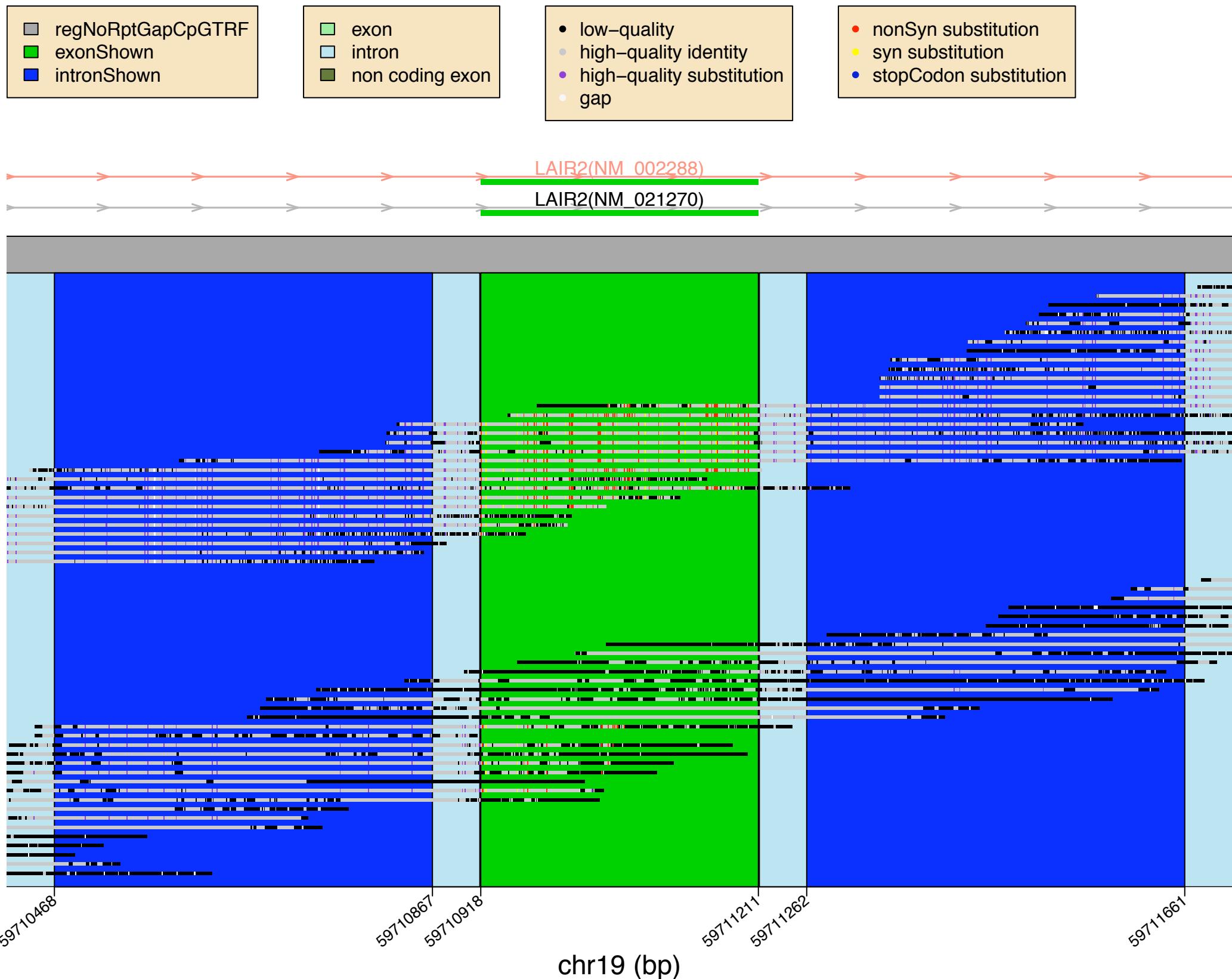


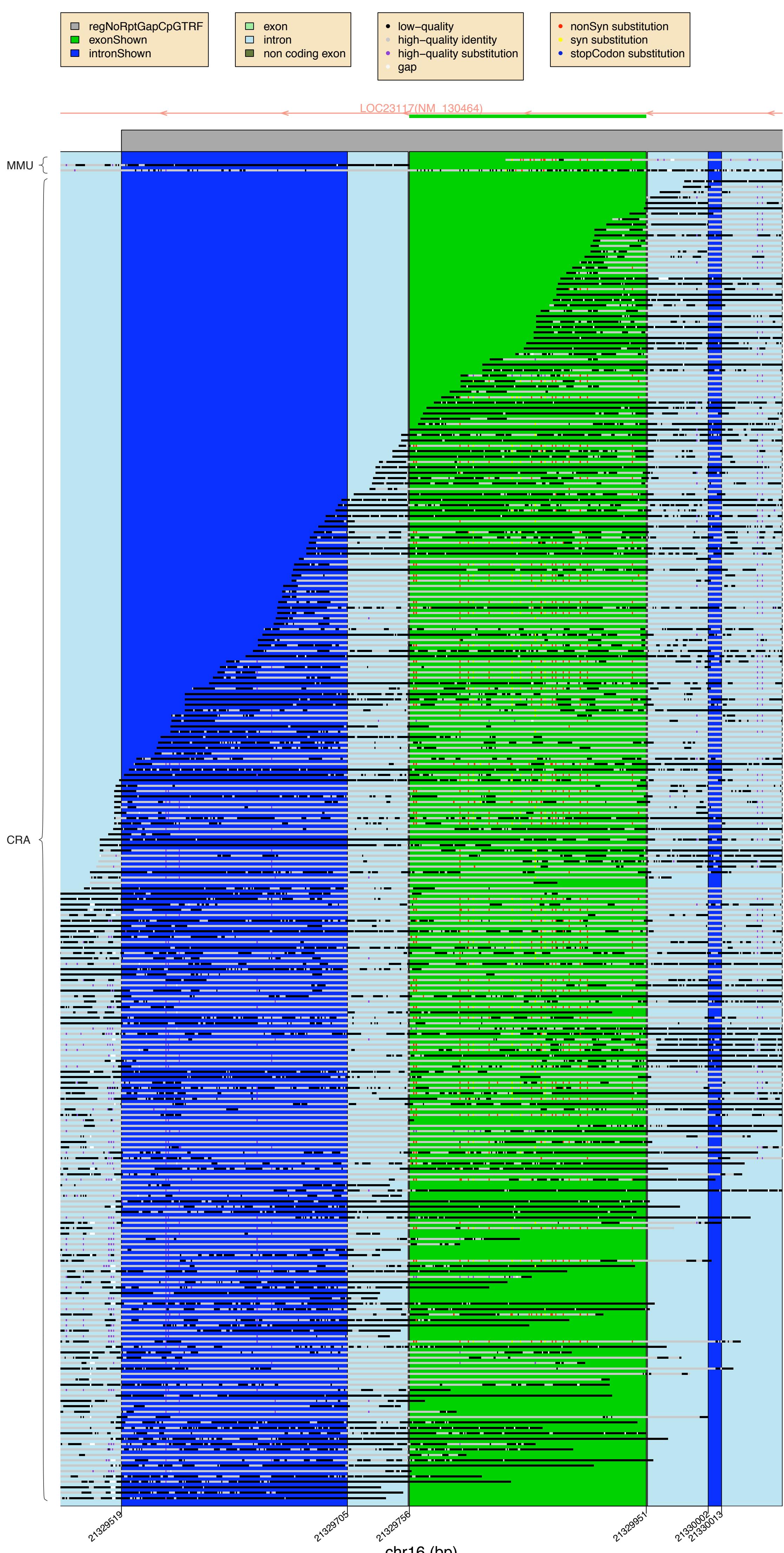
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# LOC23117\_NM\_130464\_21320950–21344159\_chr16\_exon5



# LOC23117\_NM\_130464\_21320950–21344159\_chr16\_exon3

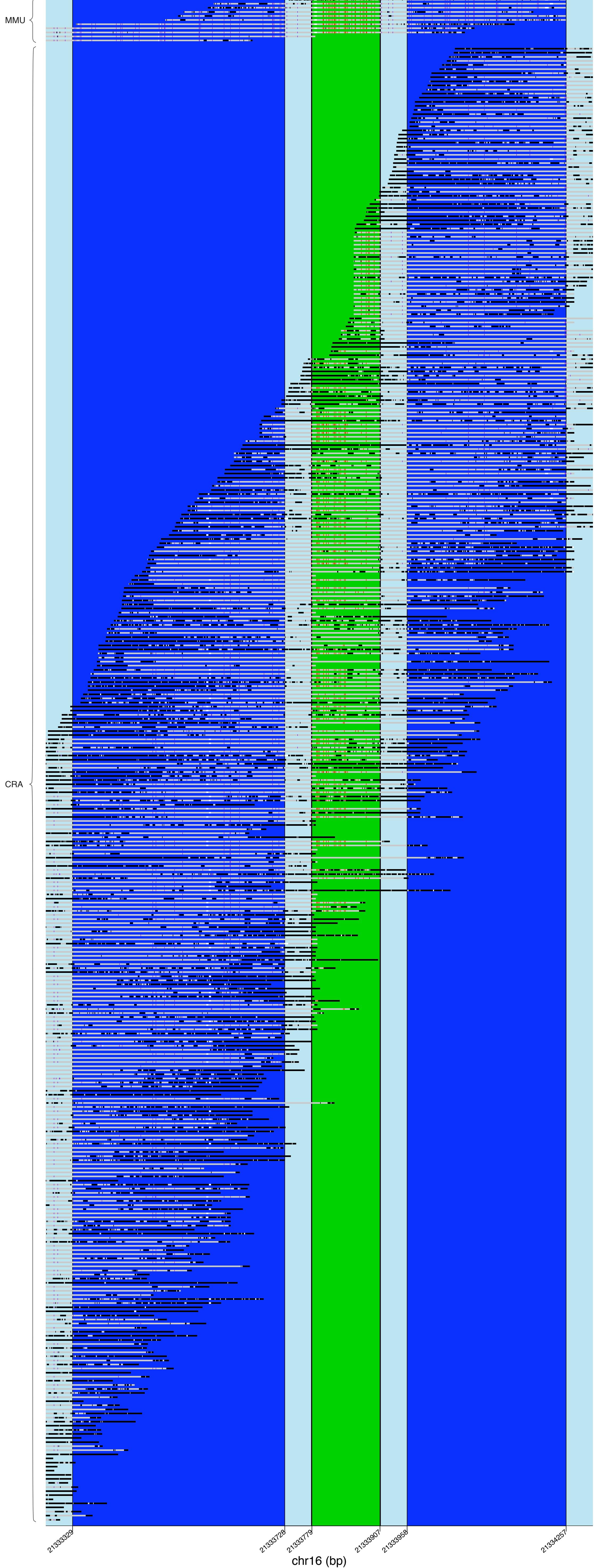
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exon  
intron  
non coding exon

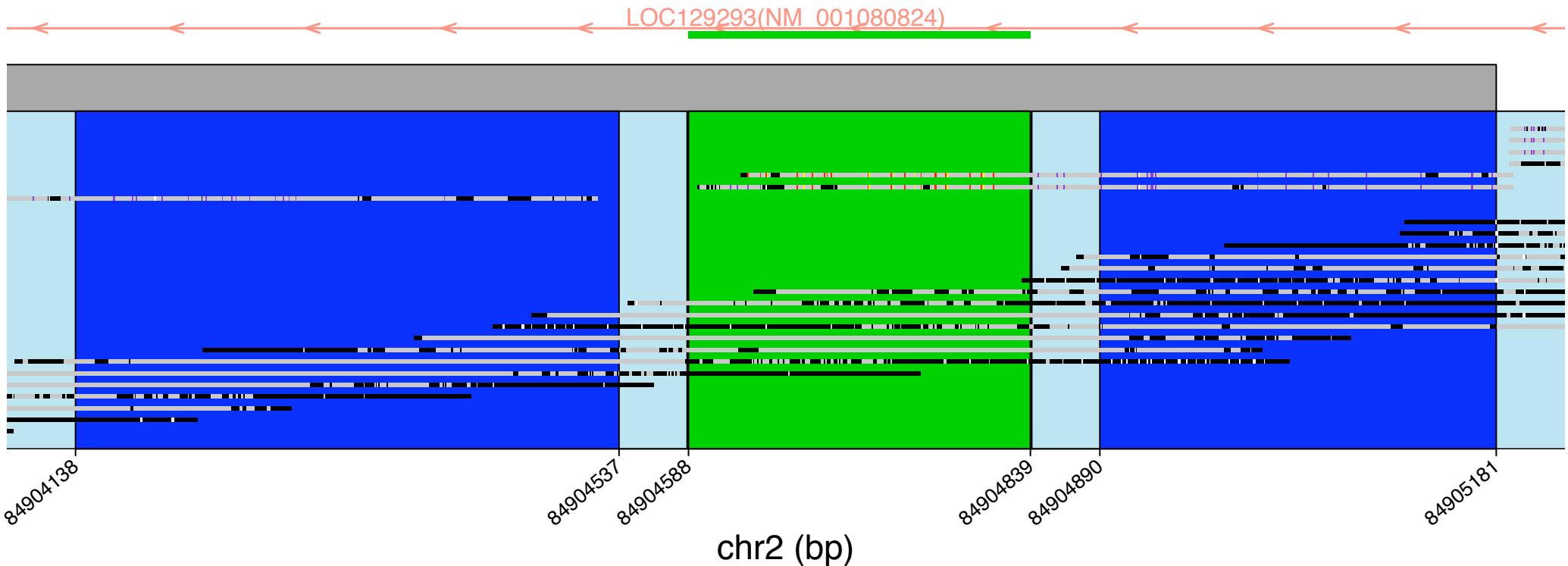
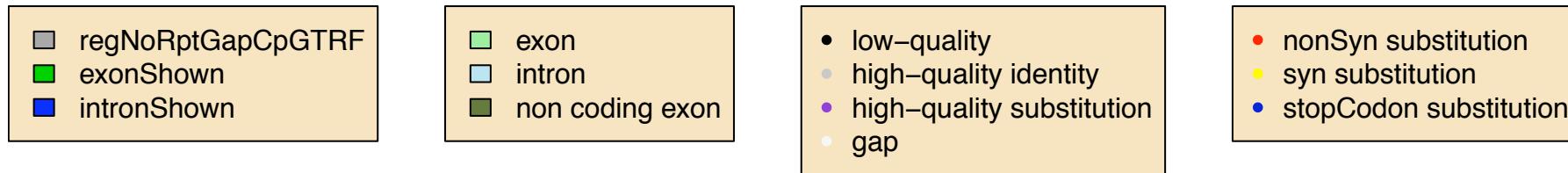
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• gap

• nonSyn substitution  
• syn substitution  
• stopCodon substitution

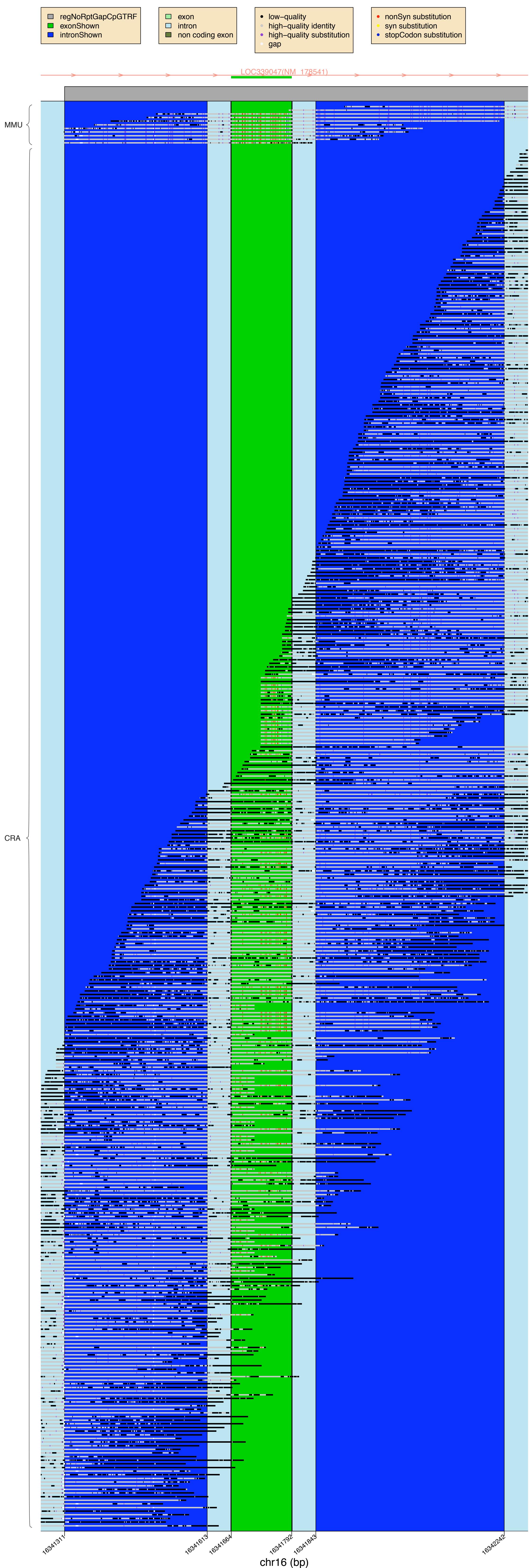
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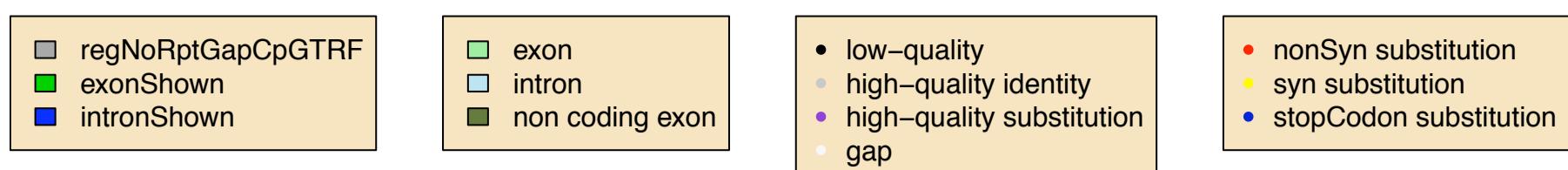
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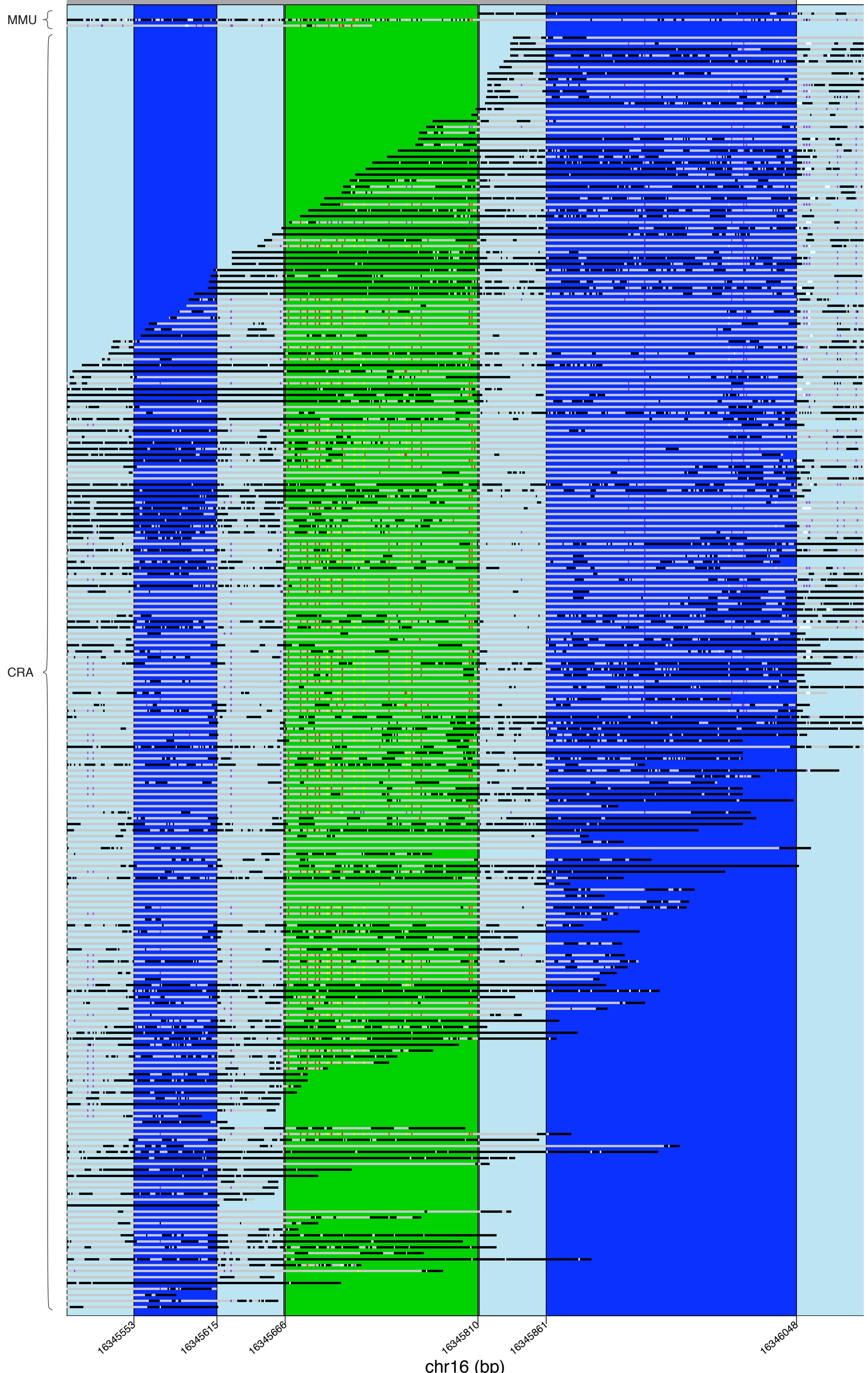
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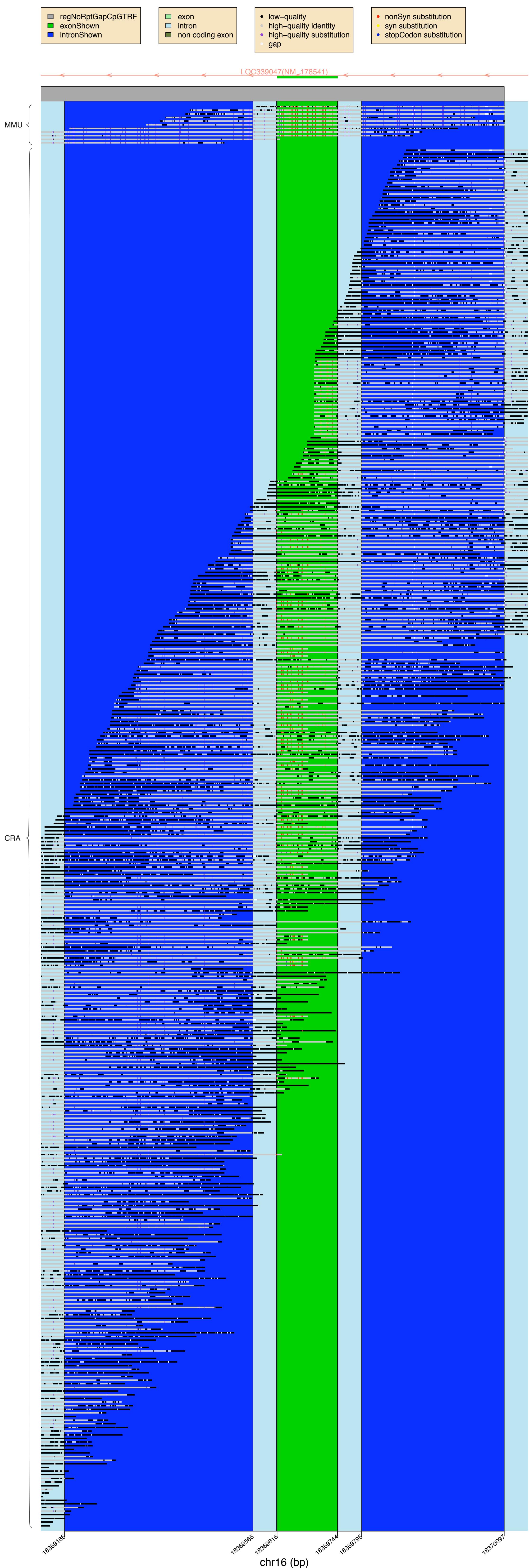
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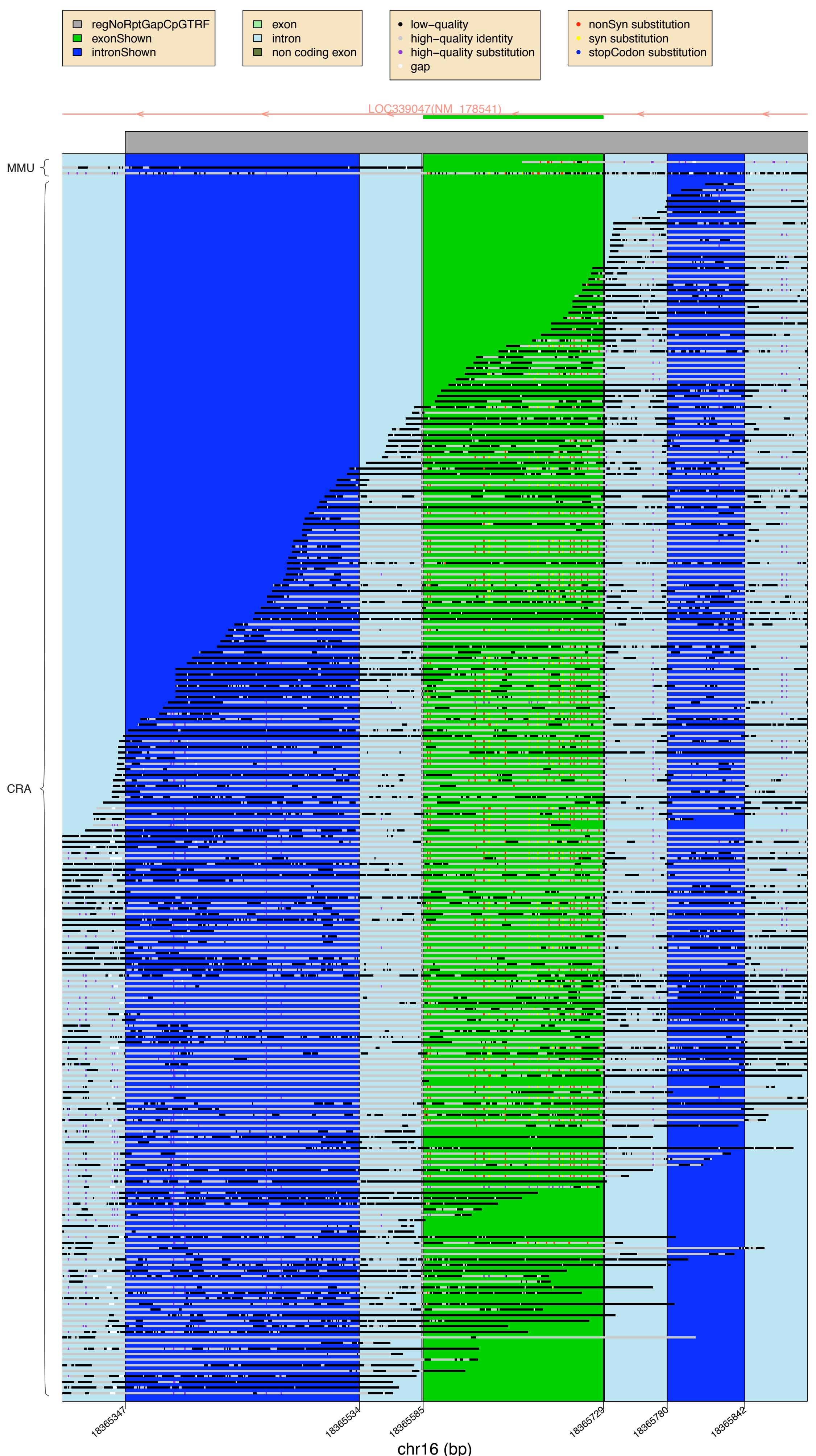
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# LOC339047\_NM\_178541\_18359455–18378197\_chr16\_exon4



# LOC339047\_NM\_178541\_18359455–18378197\_chr16\_exon6



# LOC100132247\_NM\_001135865\_2244292–22455329\_chr16\_exon2

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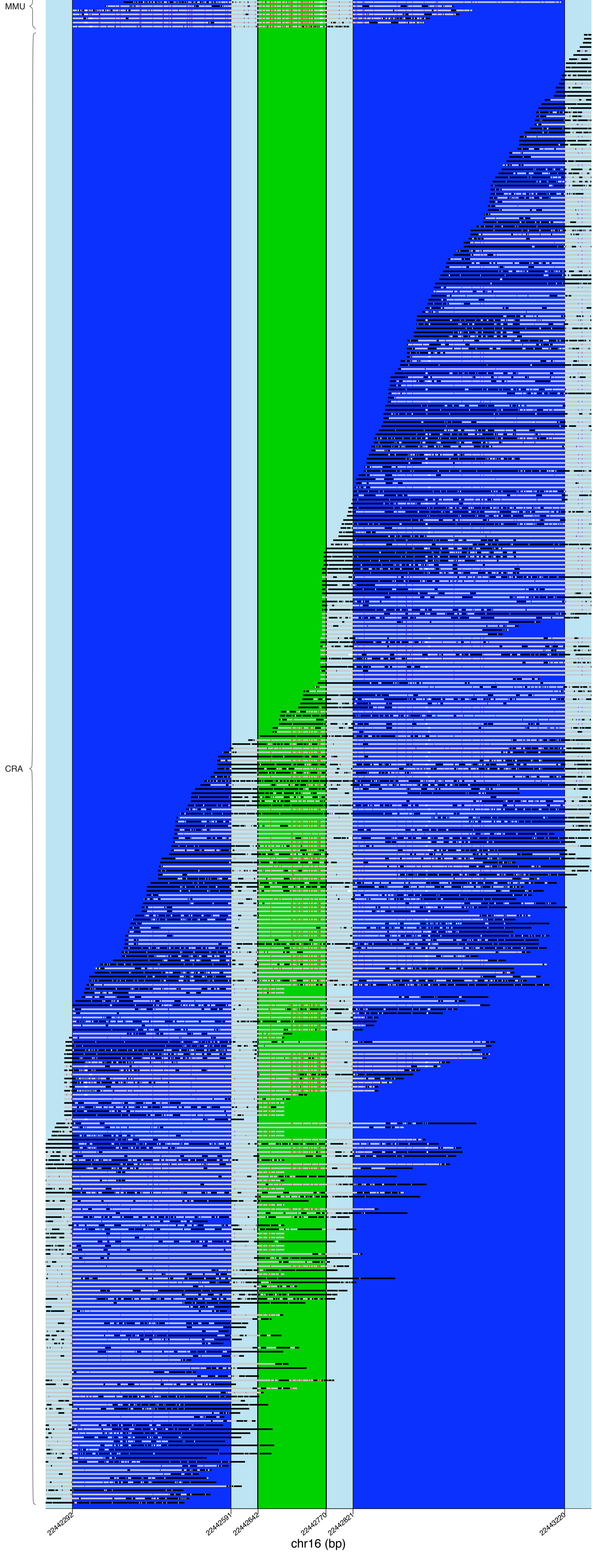
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- low-quality
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- high-quality substitution
- gap

- nonSyn substitution
- syn substitution
- stopCodon substitution

LOC100132247(NM\_001135865)

LOC23117(NM\_130464)



# LOC100132247\_NM\_001135865\_22432344–22455329\_chr16\_exon4

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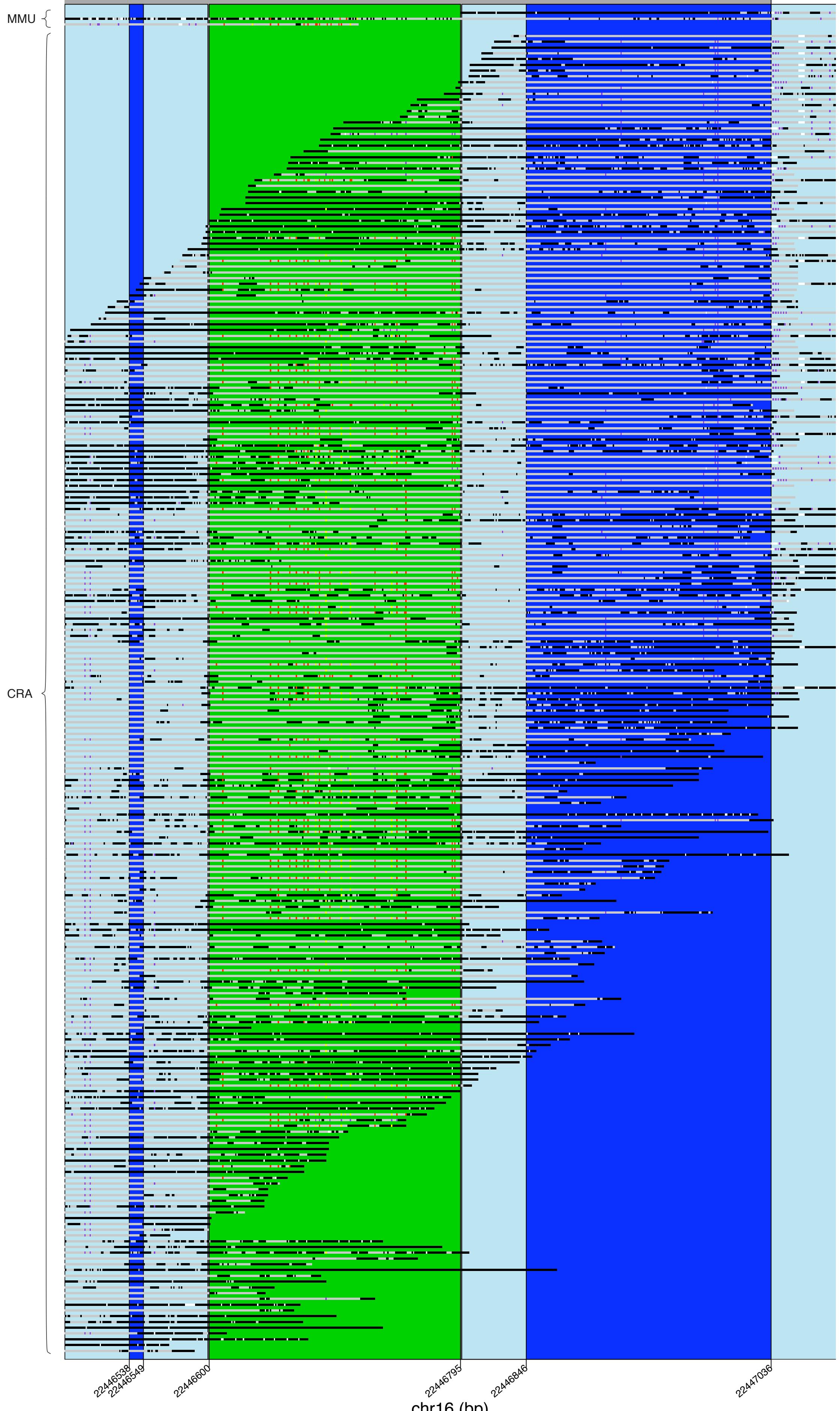
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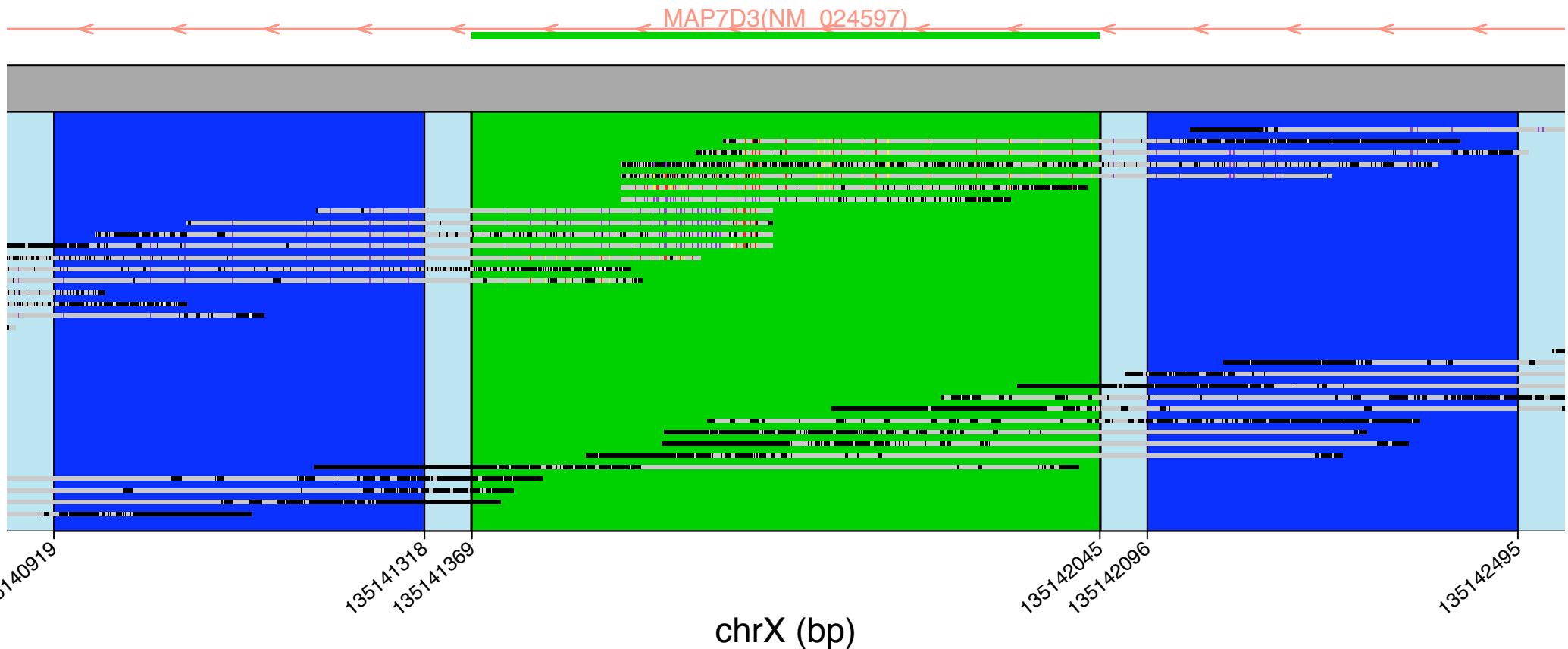
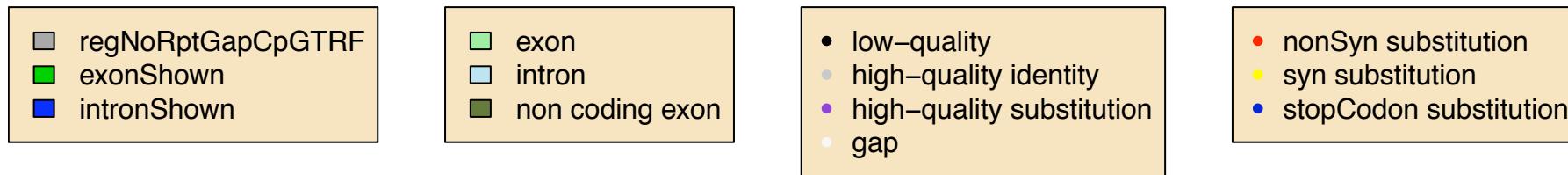
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LOC100132247(NM\_001135865)

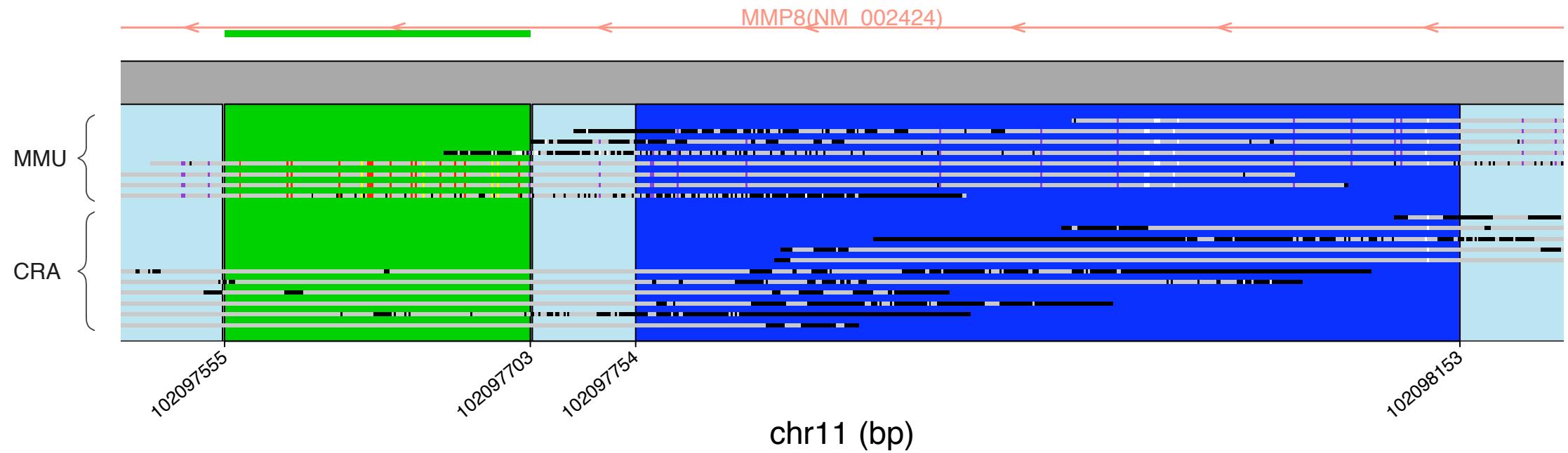
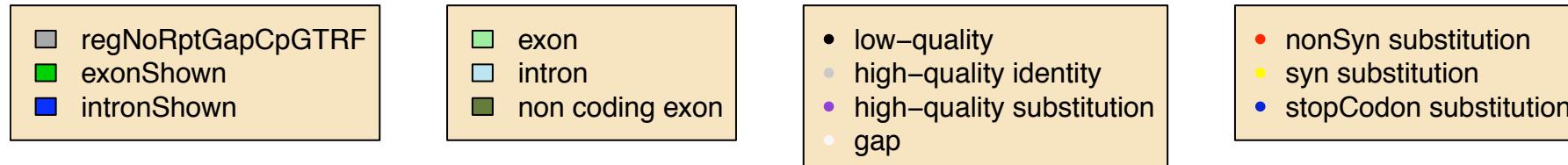
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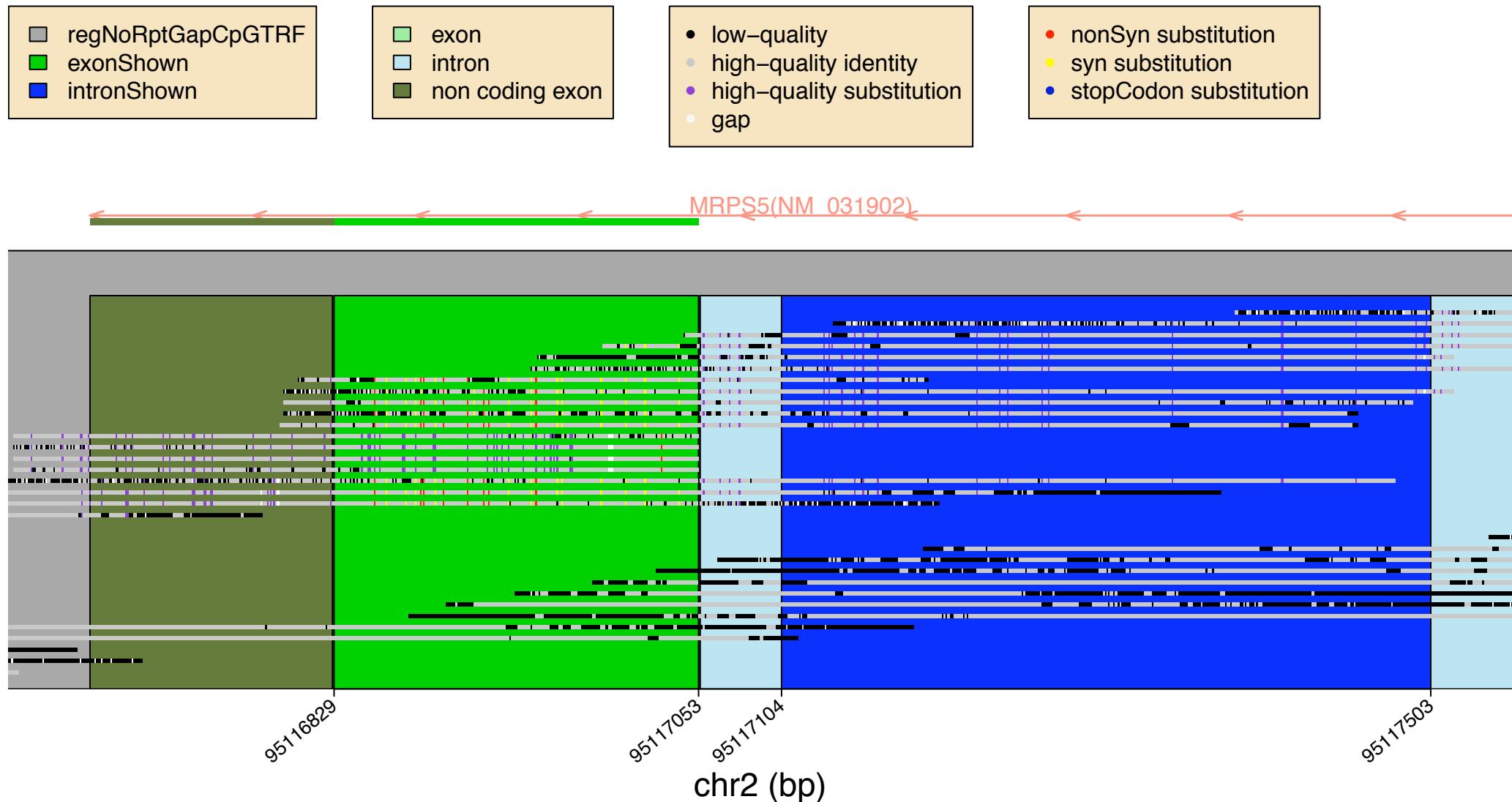
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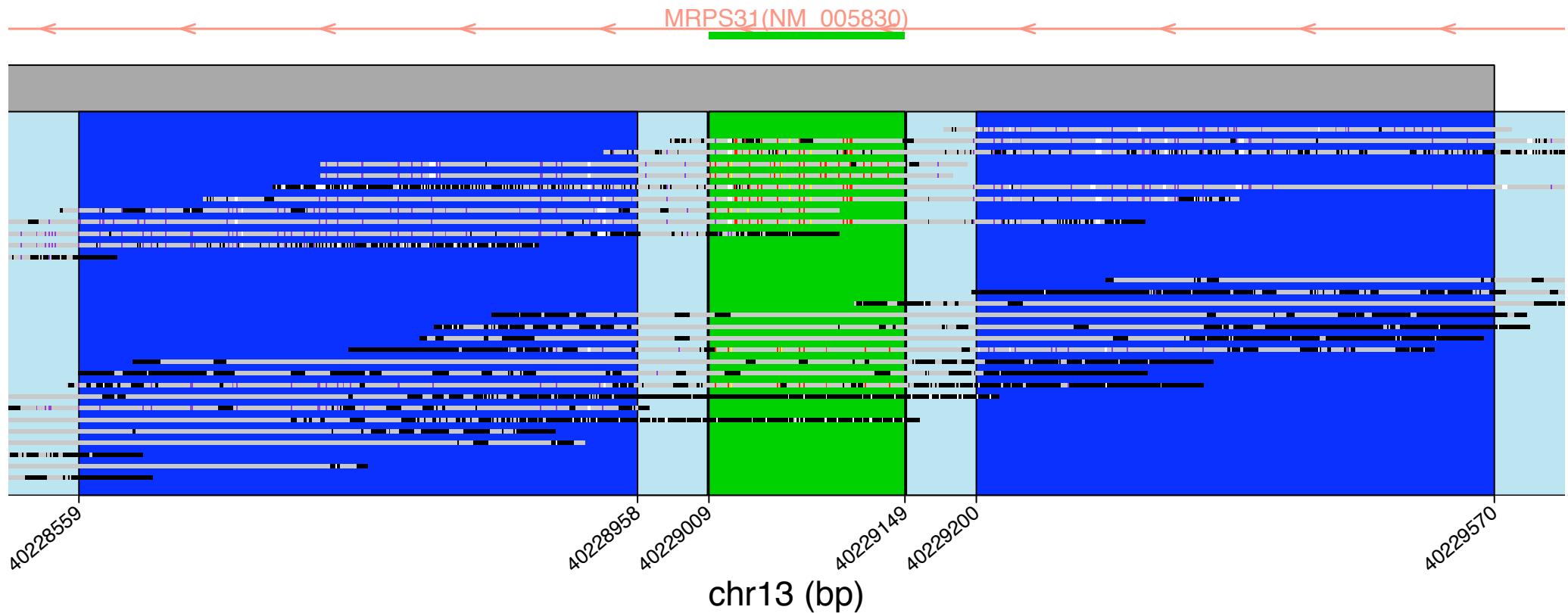
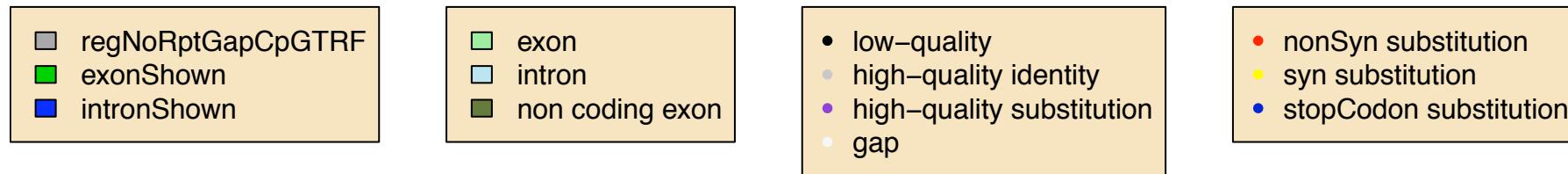
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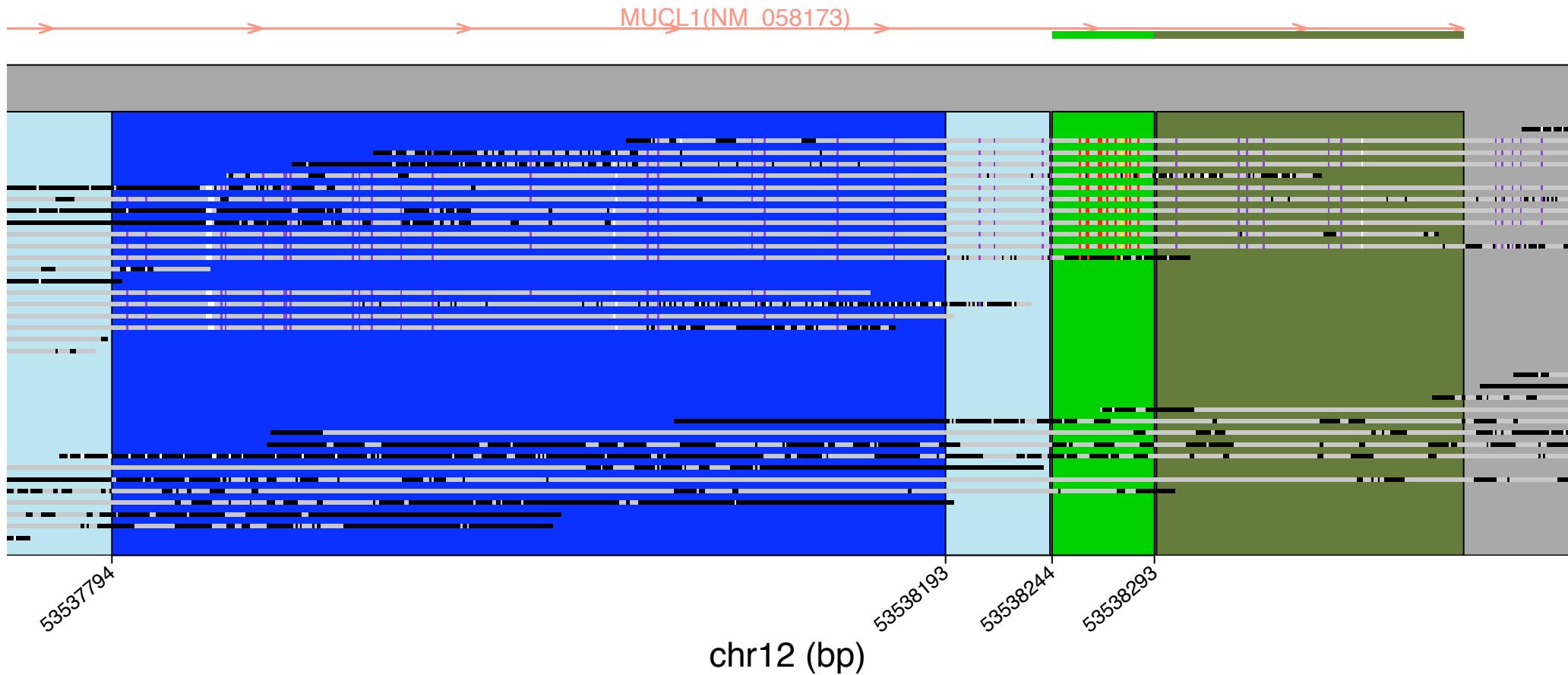
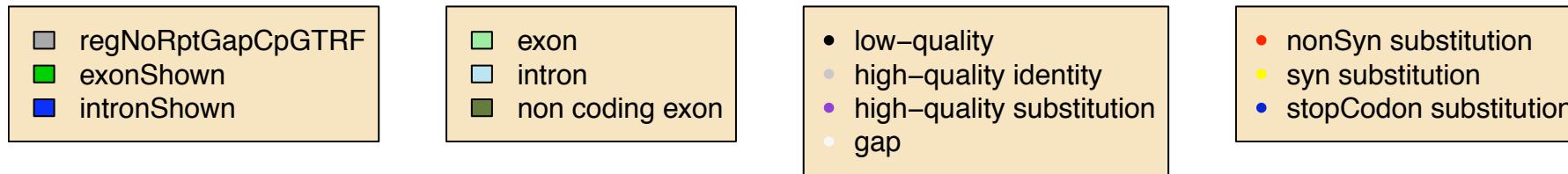
# MRPS5\_NM\_031902\_95116678–95151481\_chr2\_exon12



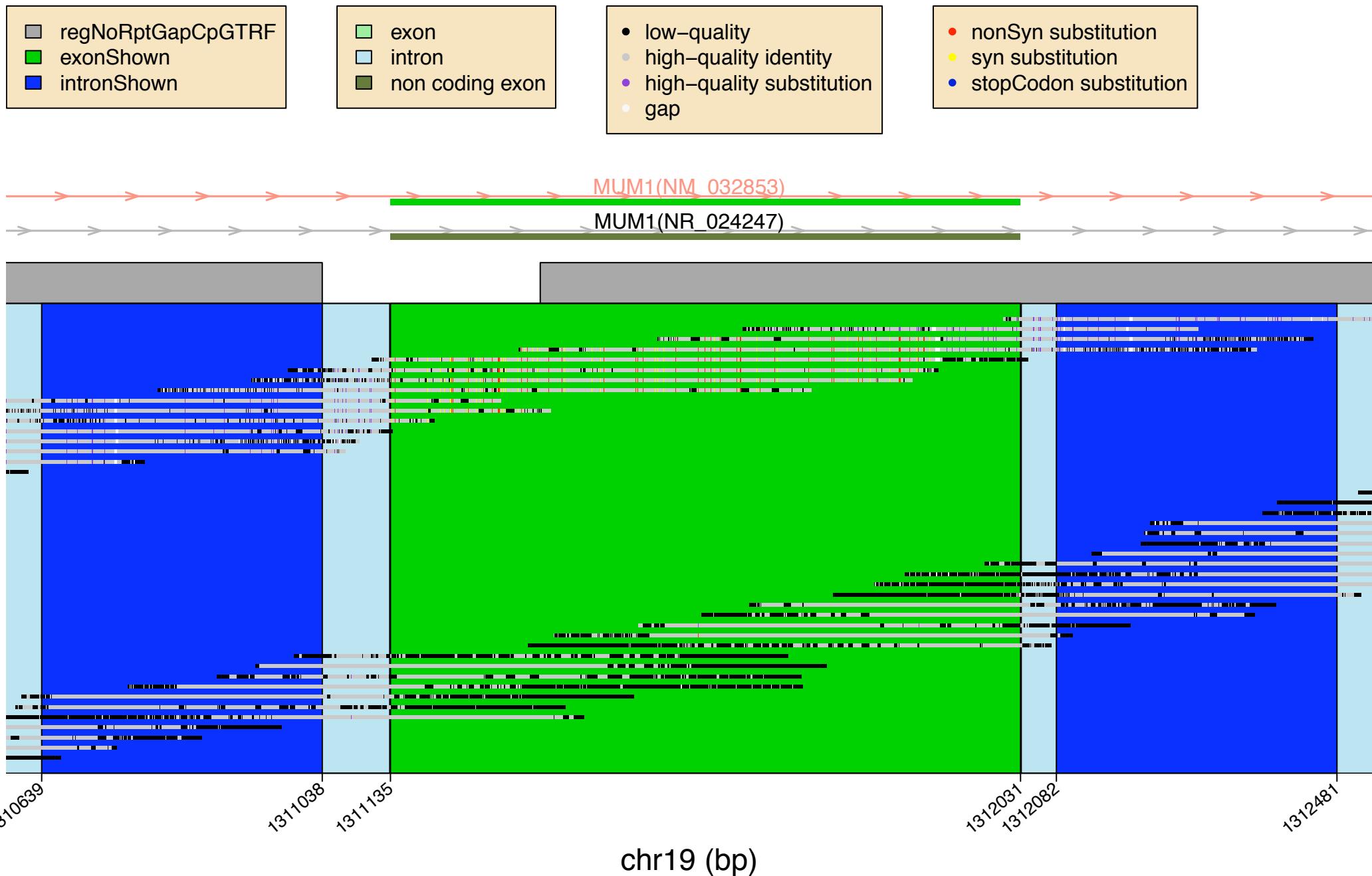
# MRPS31\_NM\_005830\_40201431–40243347\_chr13\_exon4



# MUCL1\_NM\_058173\_53534565–53538441\_chr12\_exon4



# MUM1\_NM\_032853\_1305975–1329430\_chr19\_exon5



# NPIP\_NM\_006985\_14938800–14953432\_chr16\_exon2

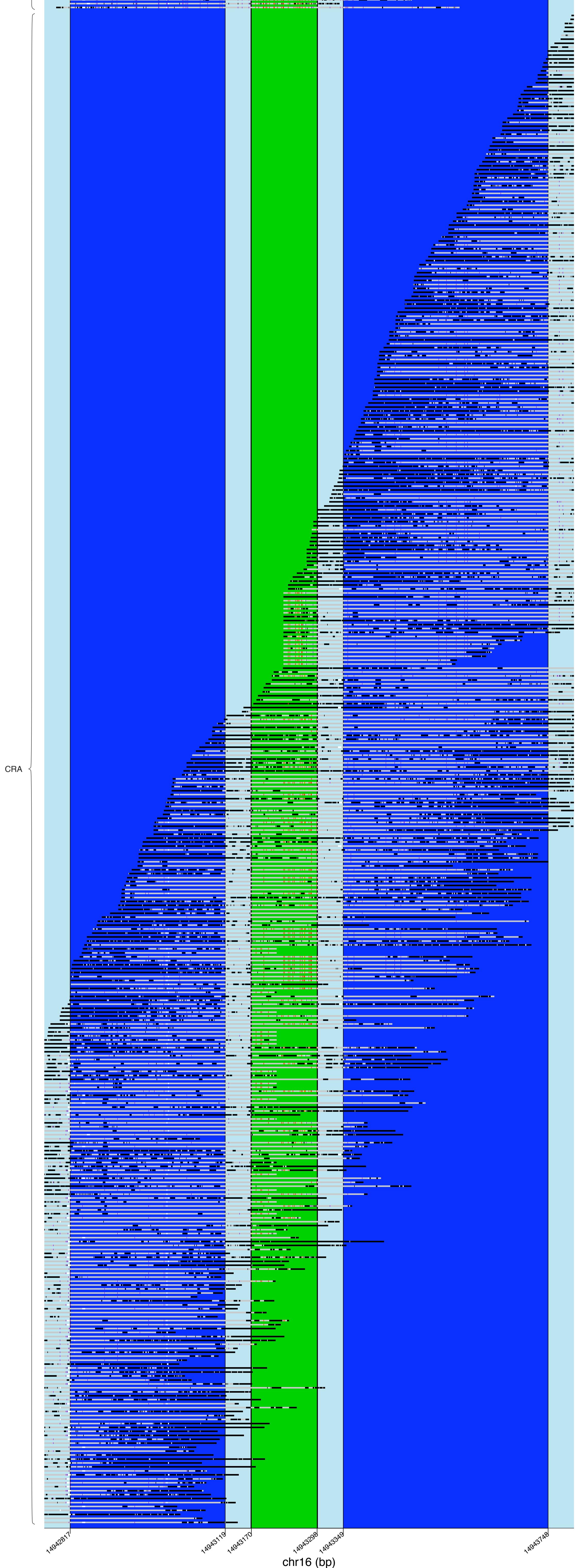
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intron  
non coding exon

low-quality  
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high-quality substitution  
gap

nonSyn substitution  
syn substitution  
stopCodon substitution

NPIP(NM\_006985)



# NPIP\_NM\_006985\_14938800–14953432\_chr16\_exon4

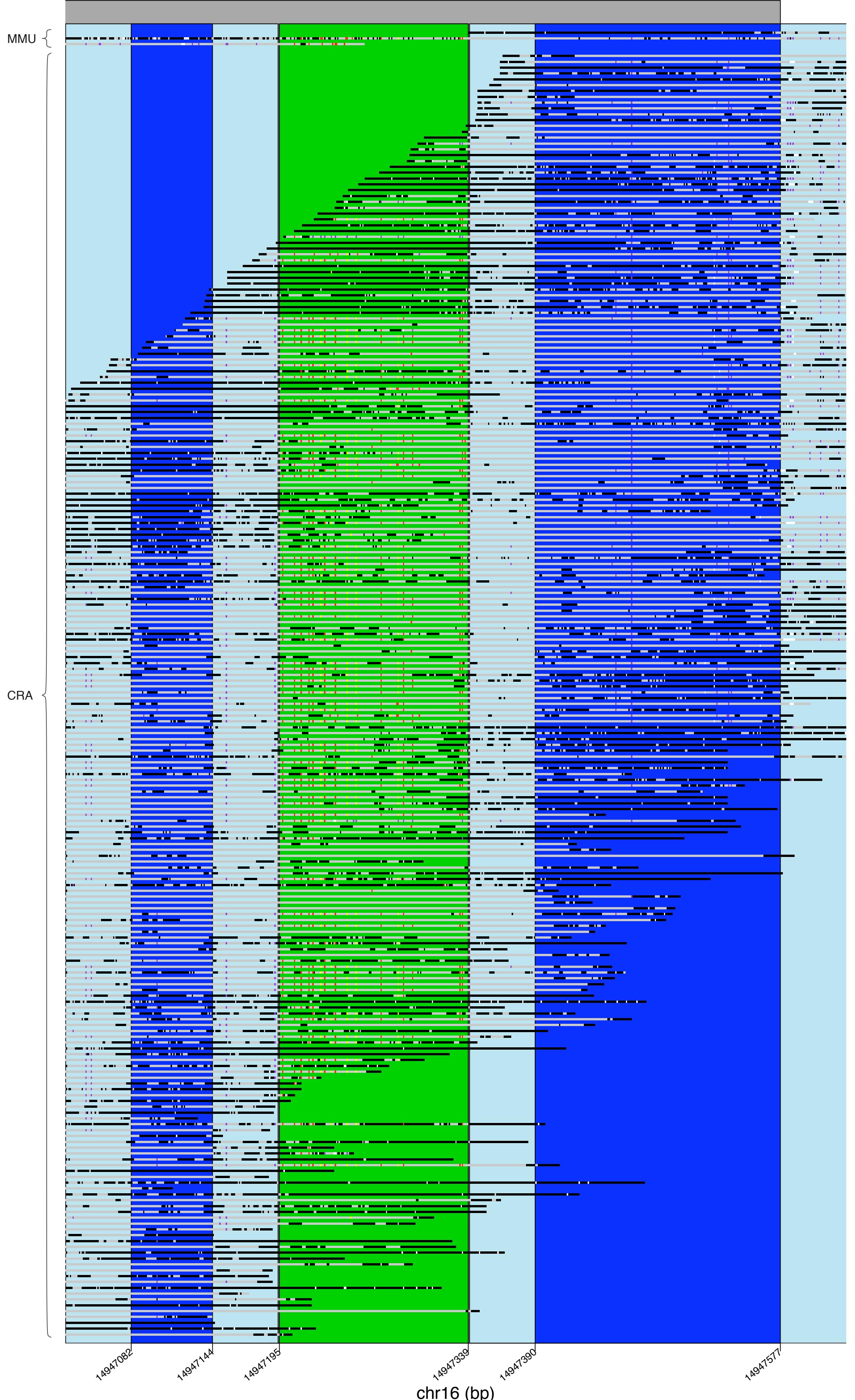
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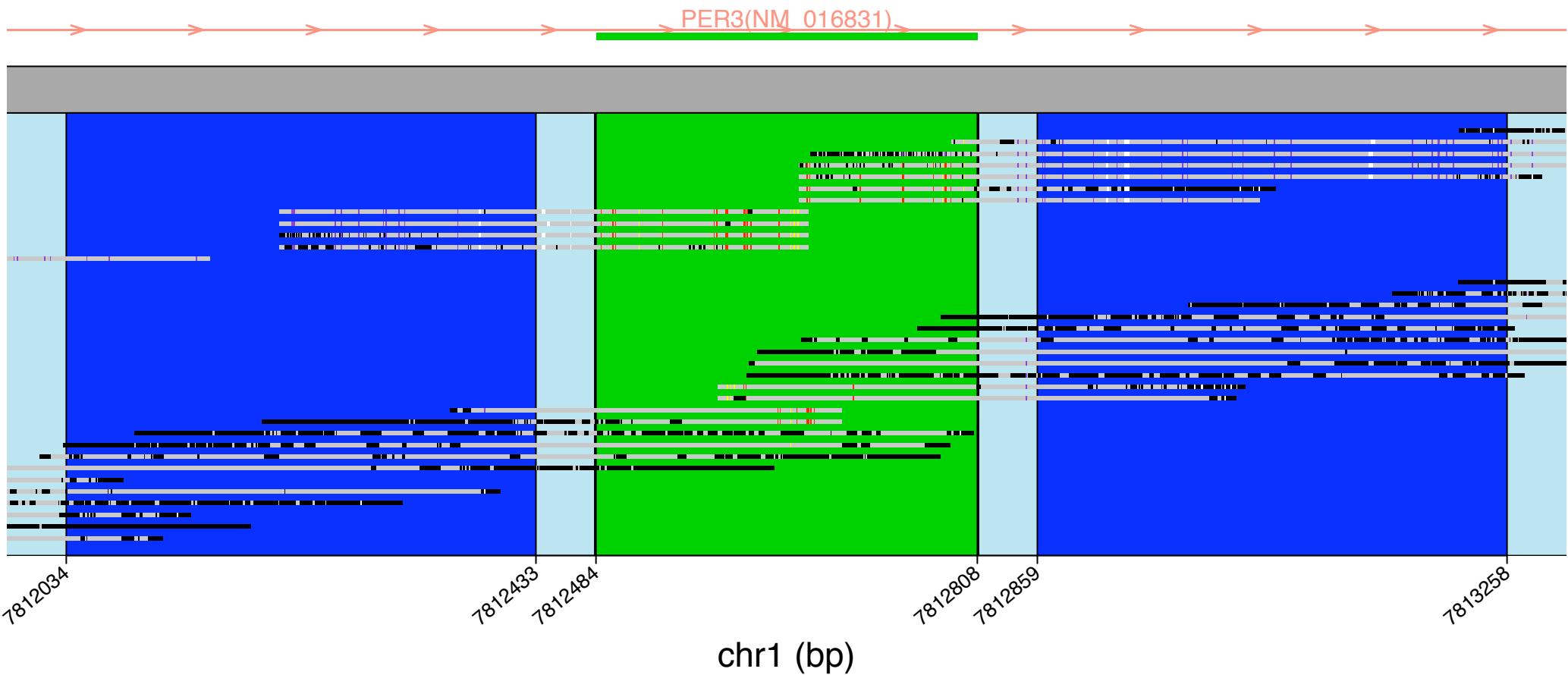
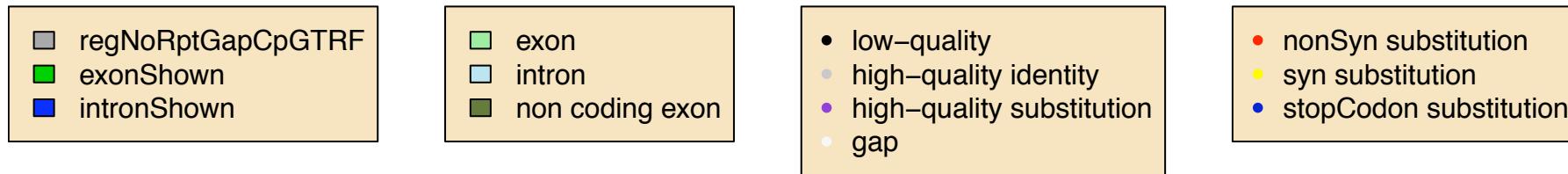
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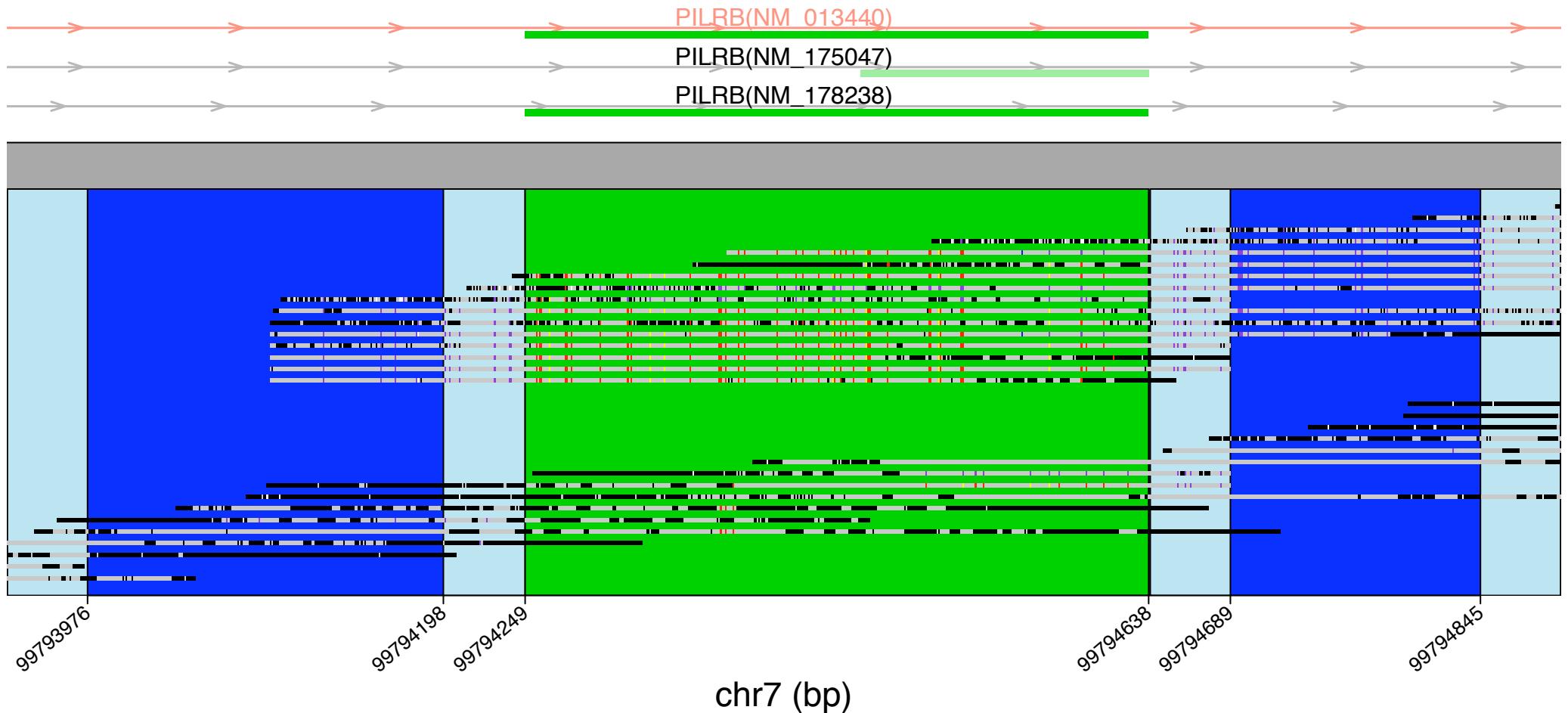
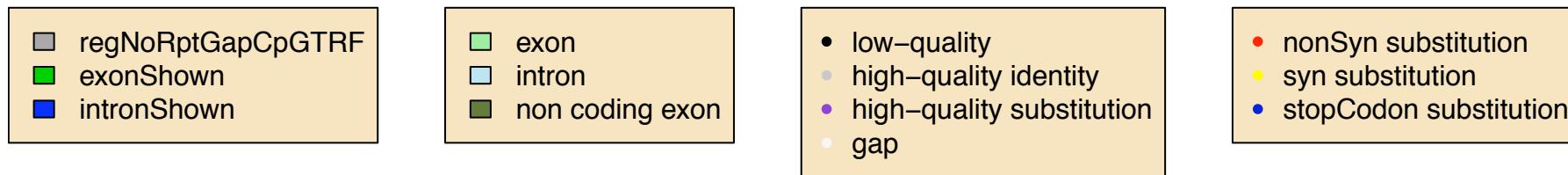
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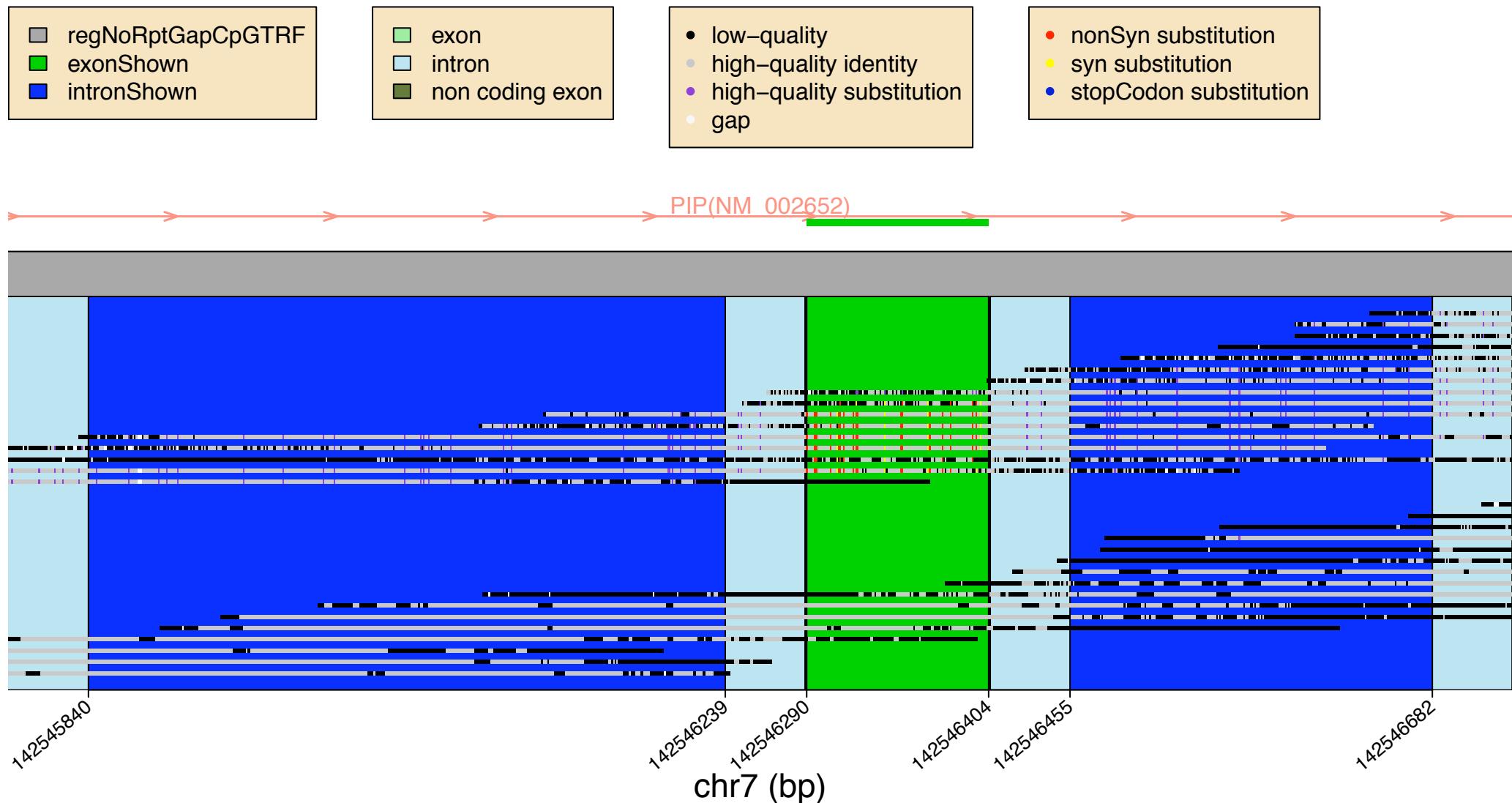
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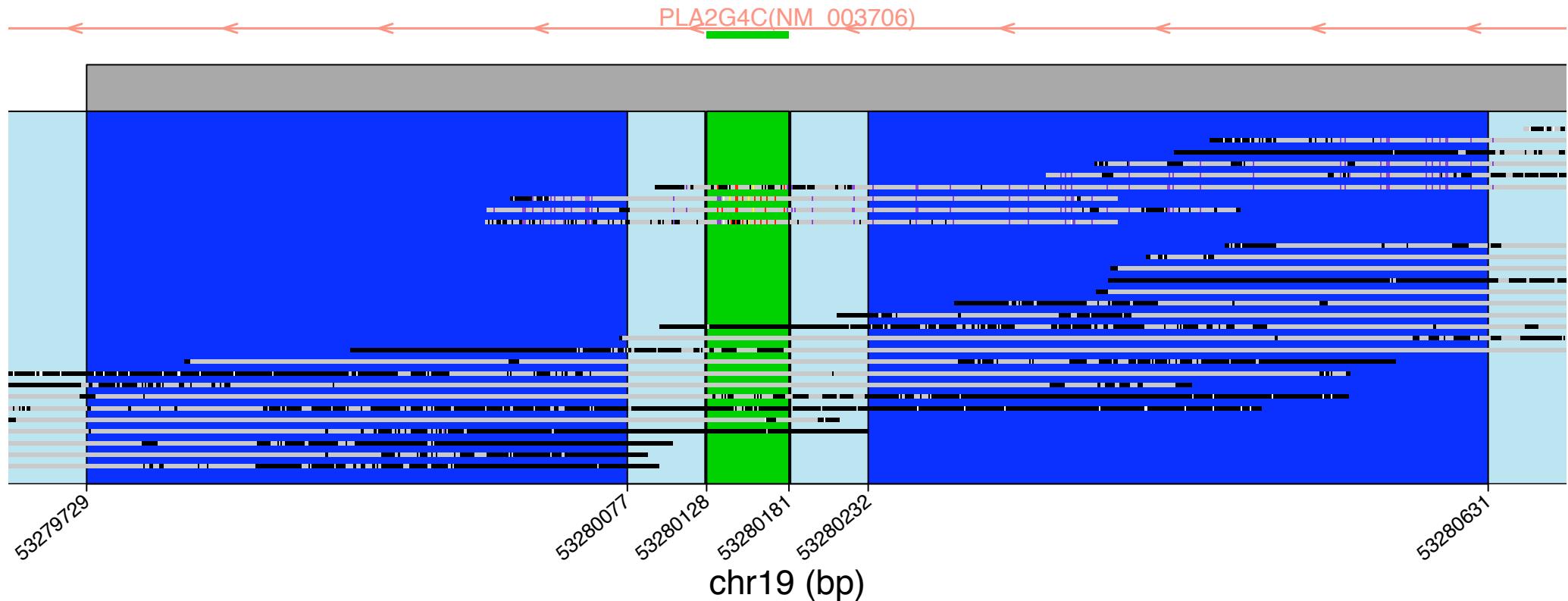
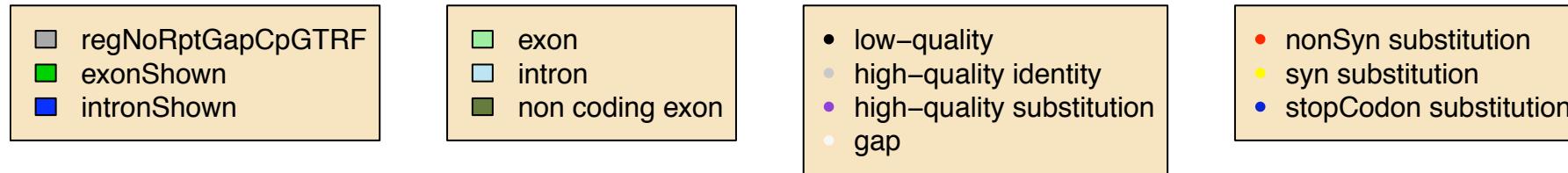
# PILRB\_NM\_013440\_99771672–99803388\_chr7\_exon16



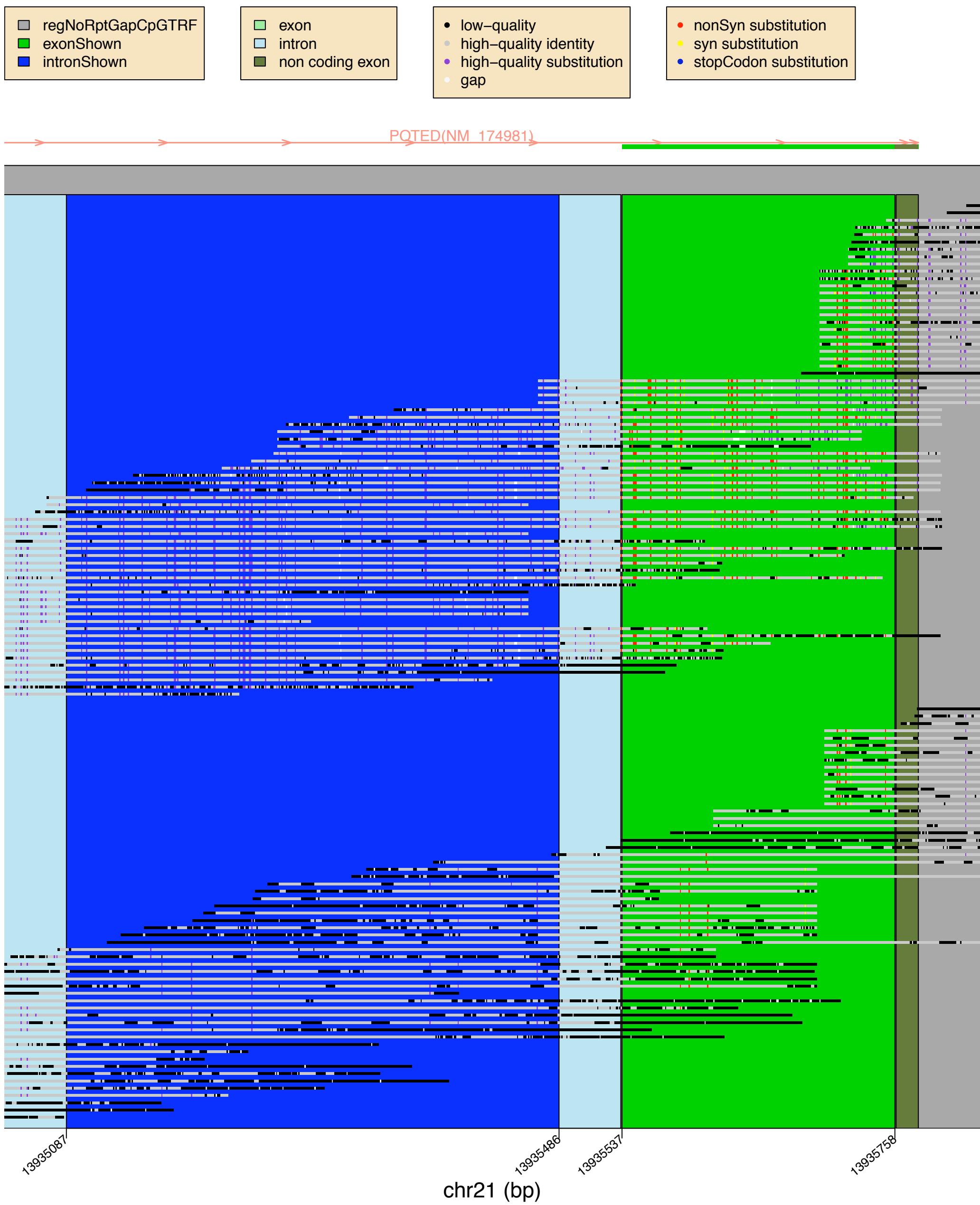
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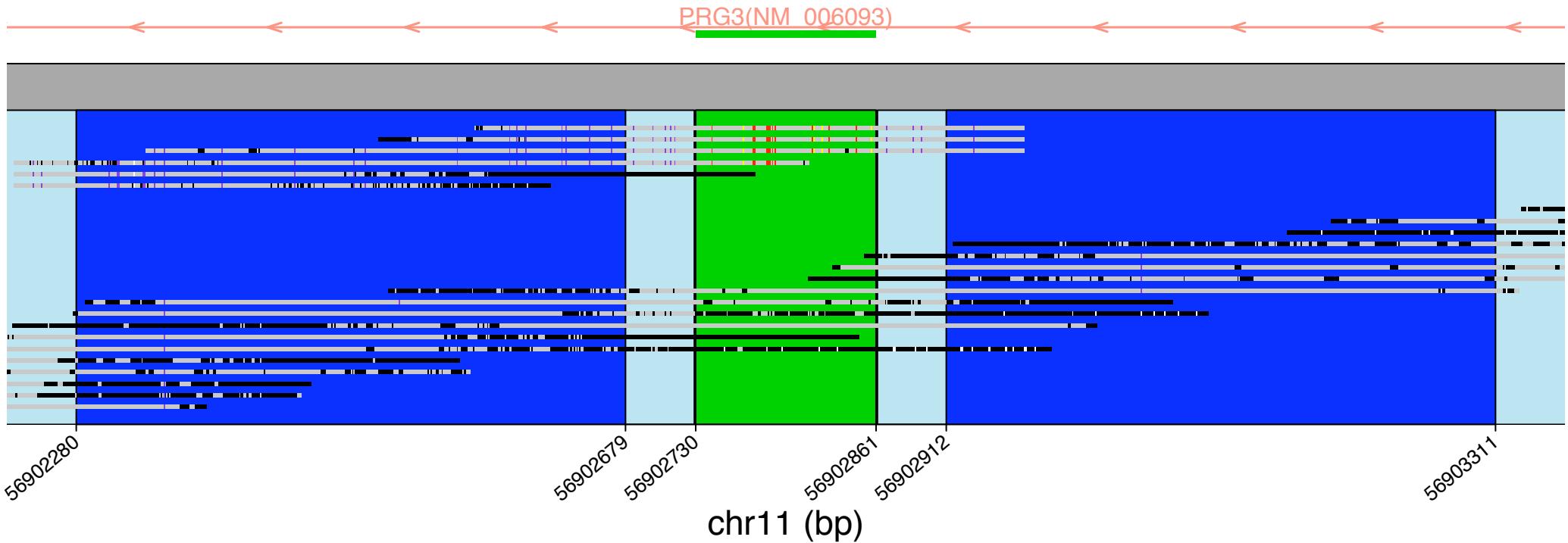
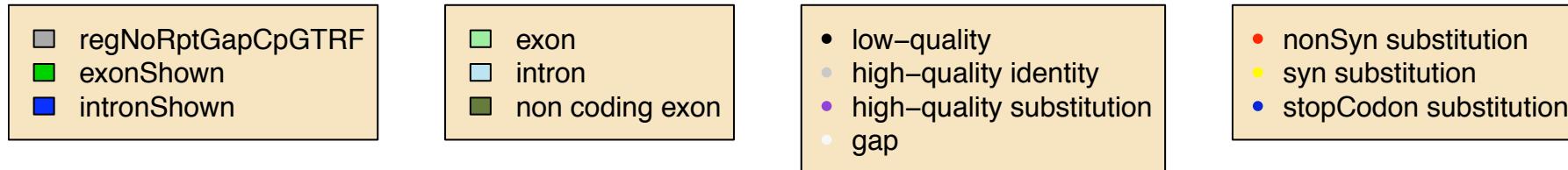
# PLA2G4C\_NM\_003706\_53242916–53305826\_chr19\_exon10



# POTED\_NM\_174981\_13904368–13935777\_chr21\_exon11



# PRG3\_NM\_006093\_56900818–56905199\_chr11\_exon4



# PSG2\_NM\_031246\_48260201–48278654\_chr19\_exon5

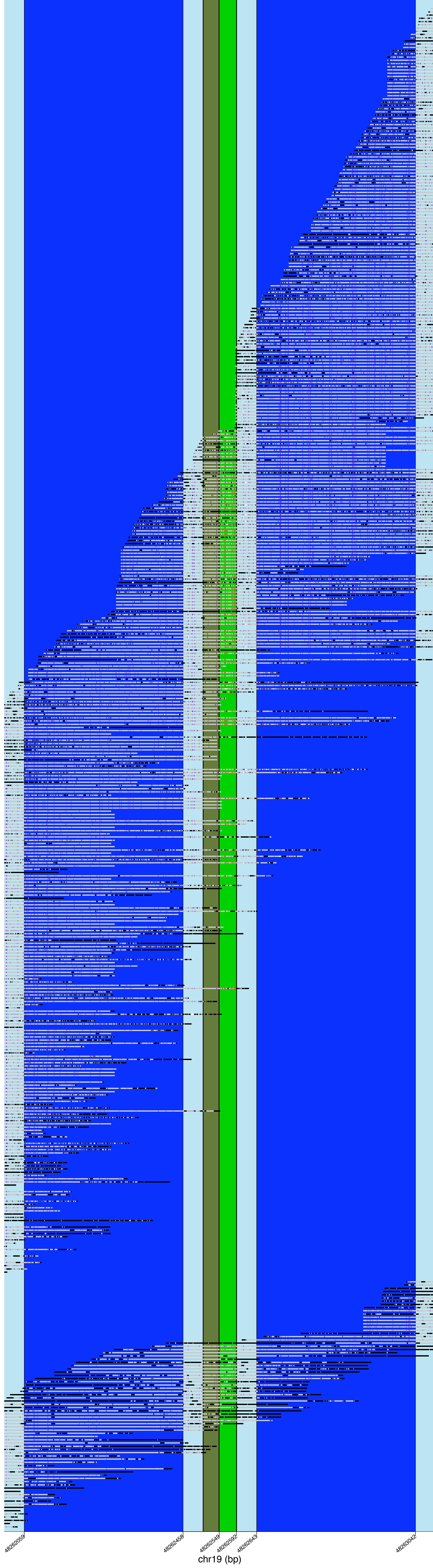
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exon  
intron  
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high-quality substitution  
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nonSyn substitution  
syn substitution  
stopCodon substitution

PSG2(NM\_031246)



# PSG7\_NM\_002783\_48120123–48133170\_chr19\_exon5

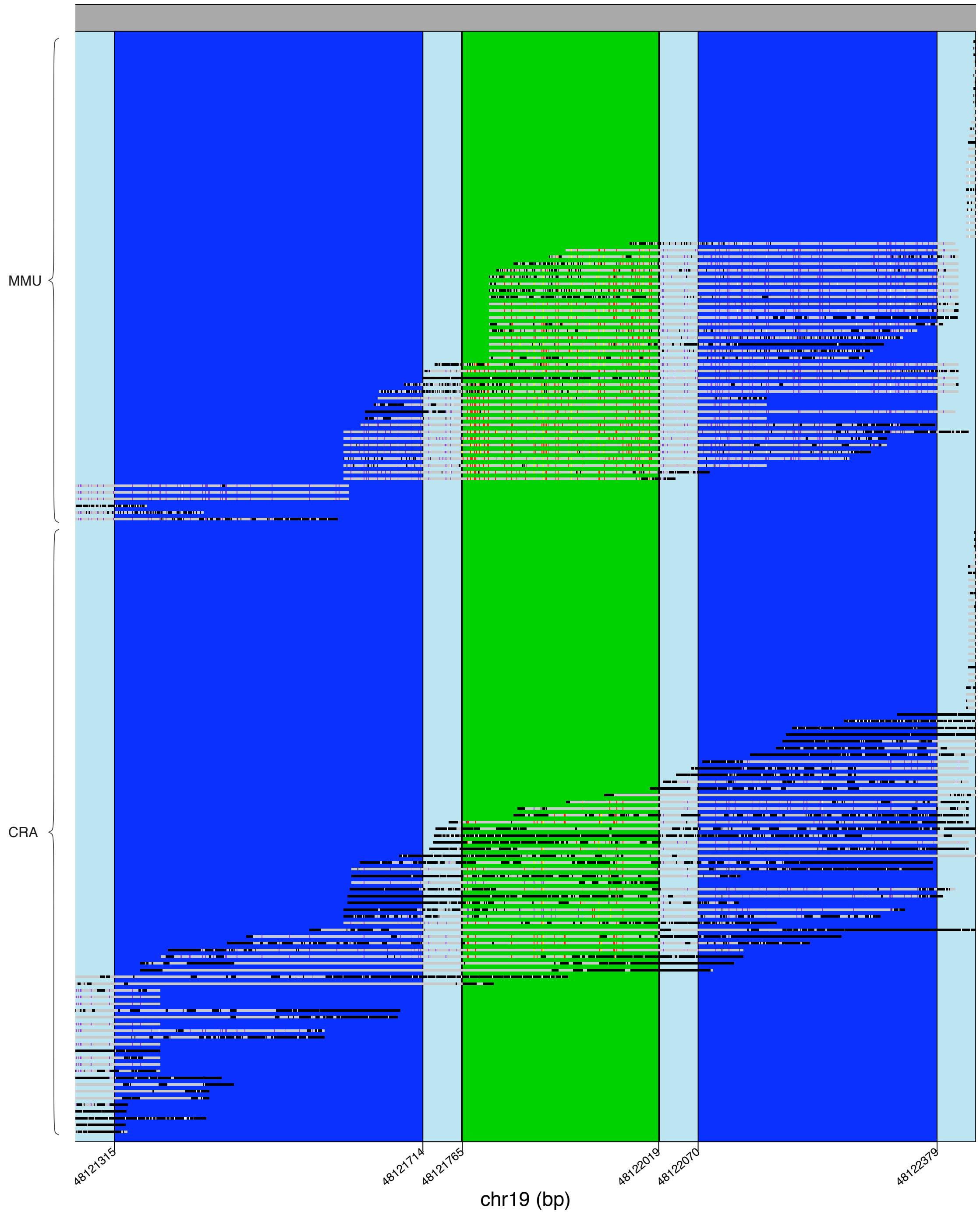
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exon  
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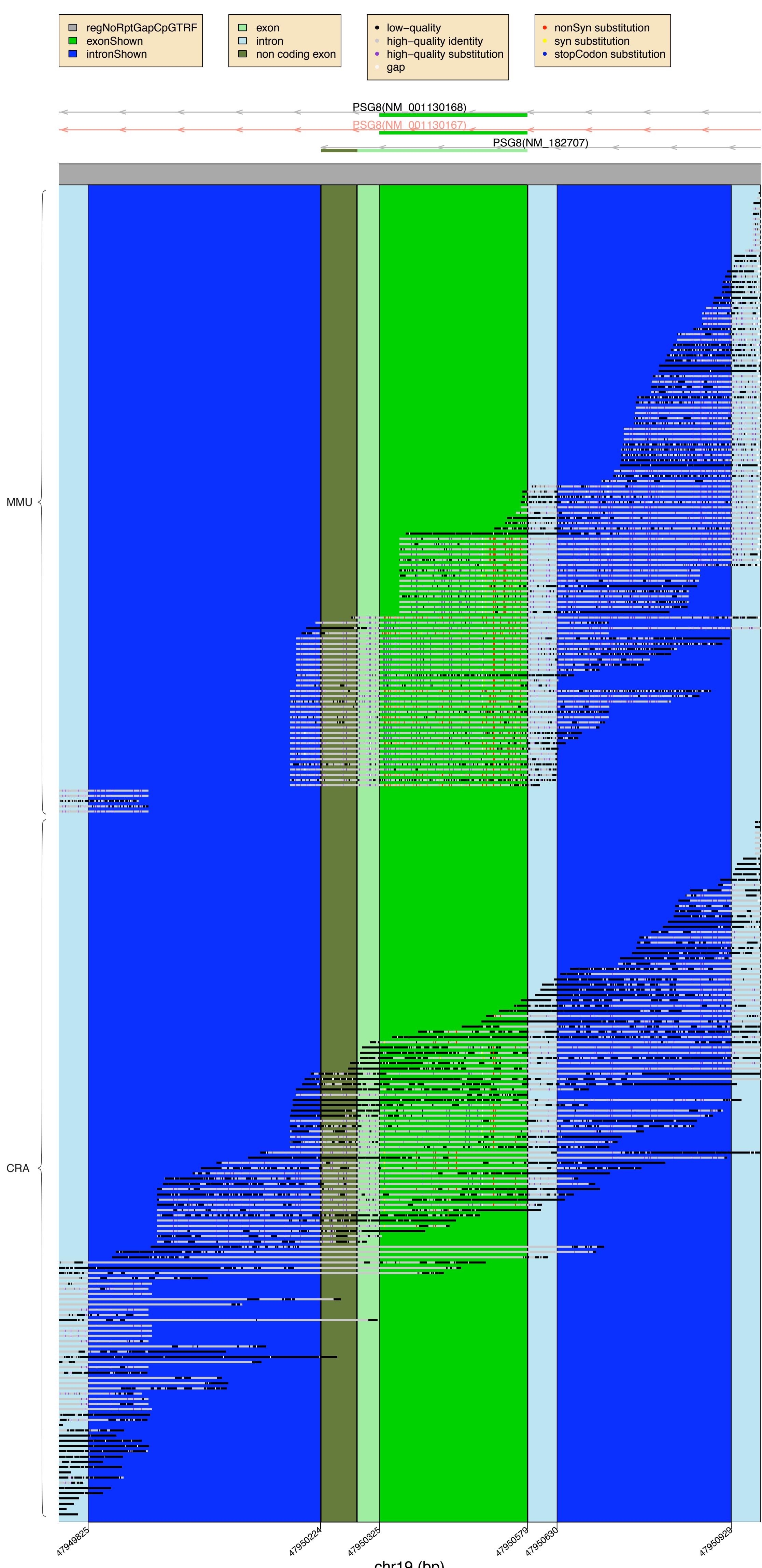
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- stopCodon substitution

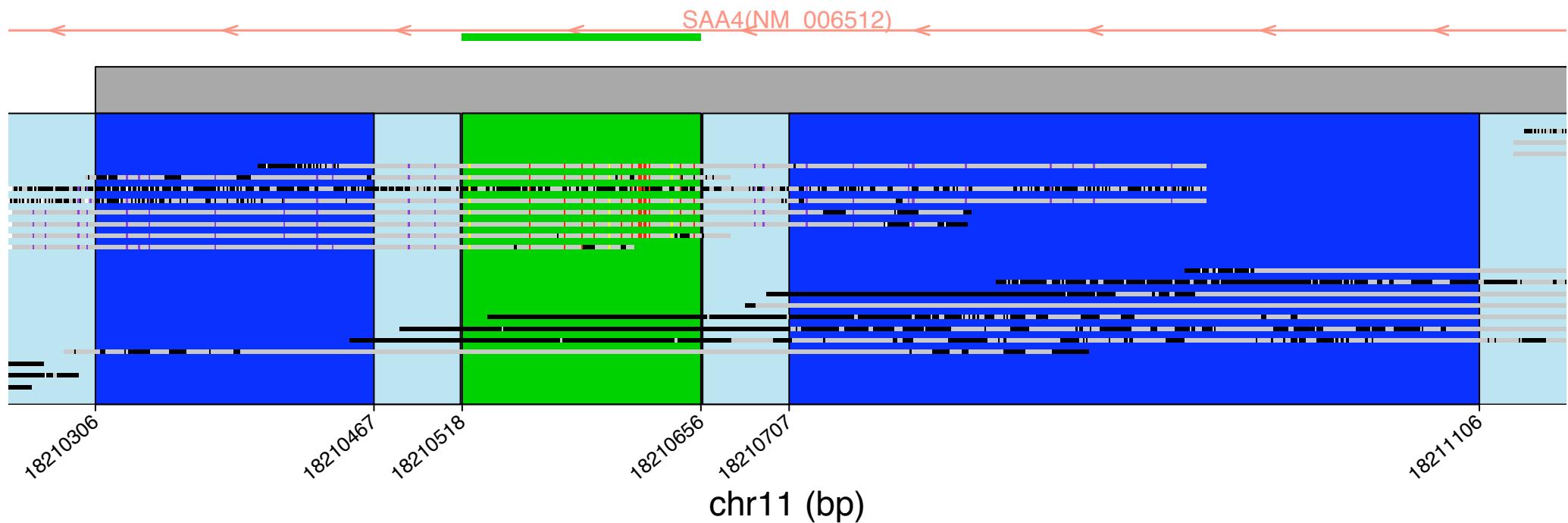
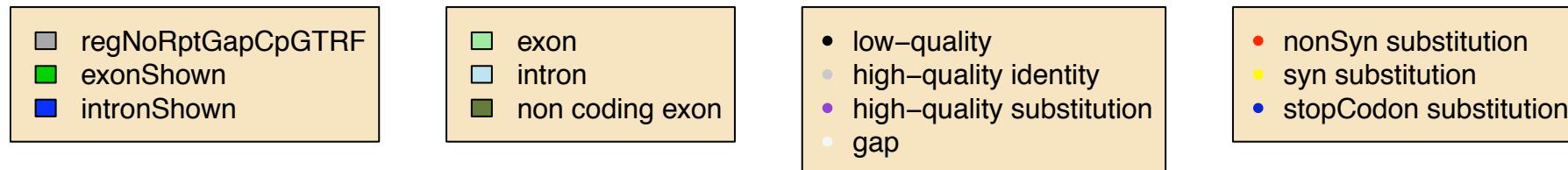
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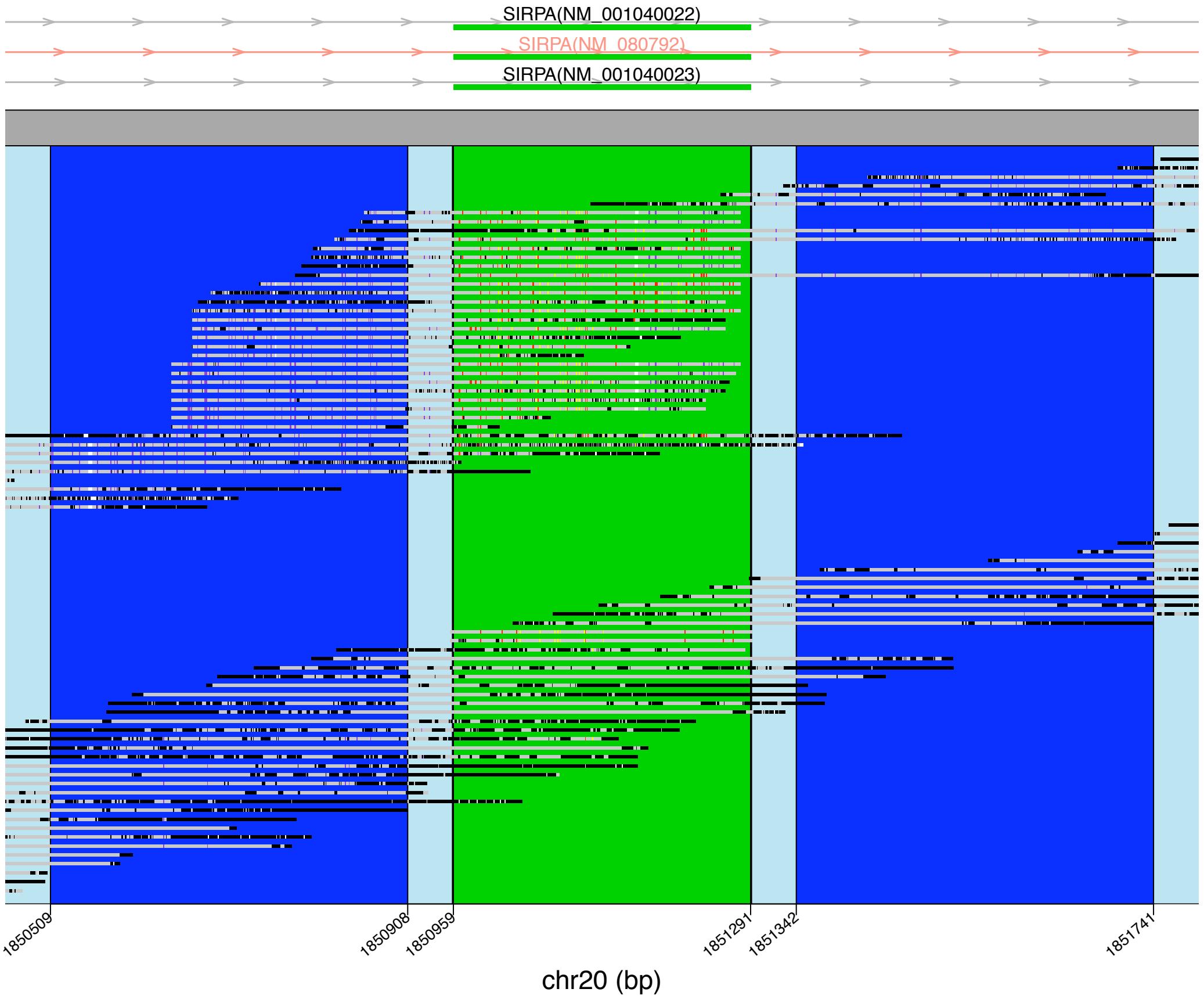
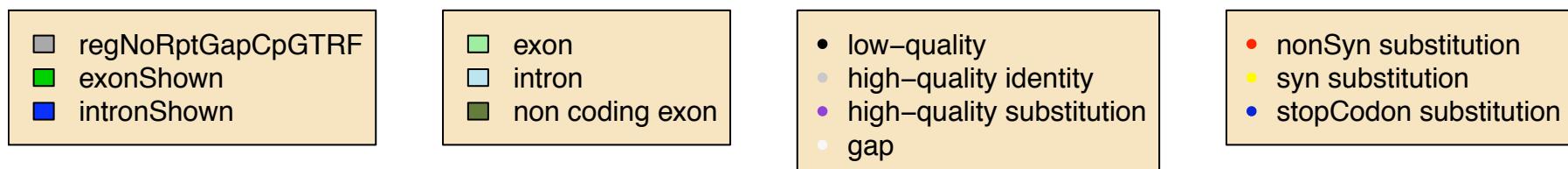
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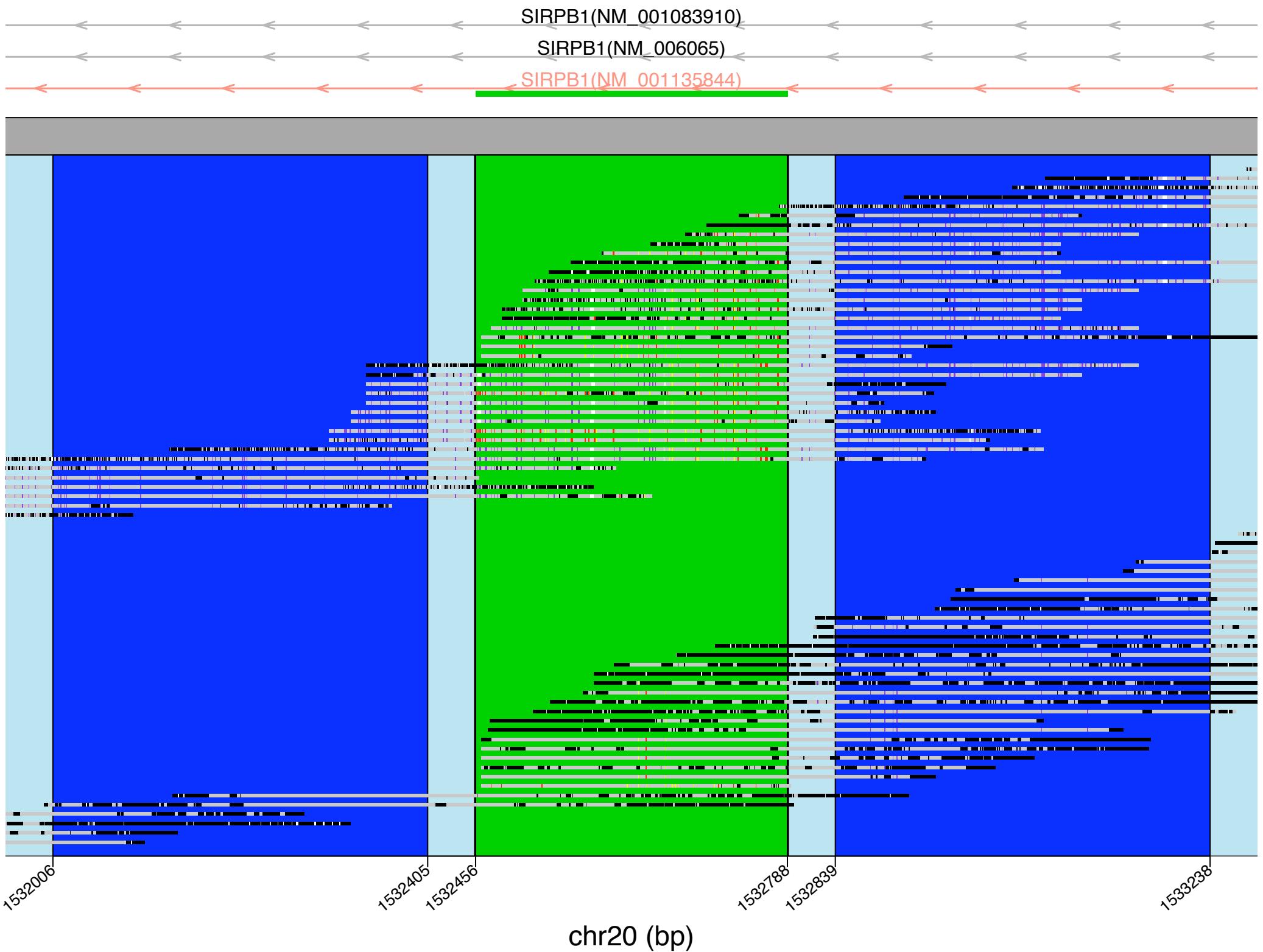
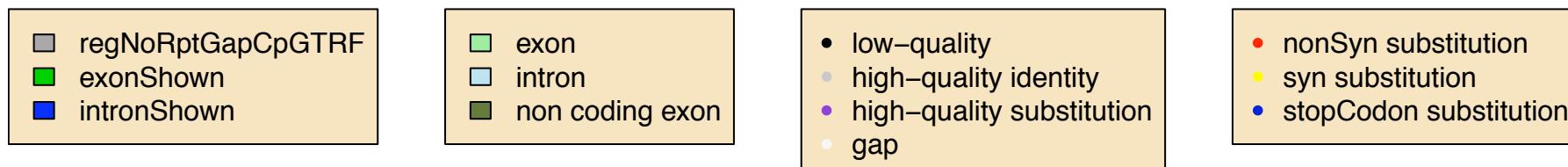
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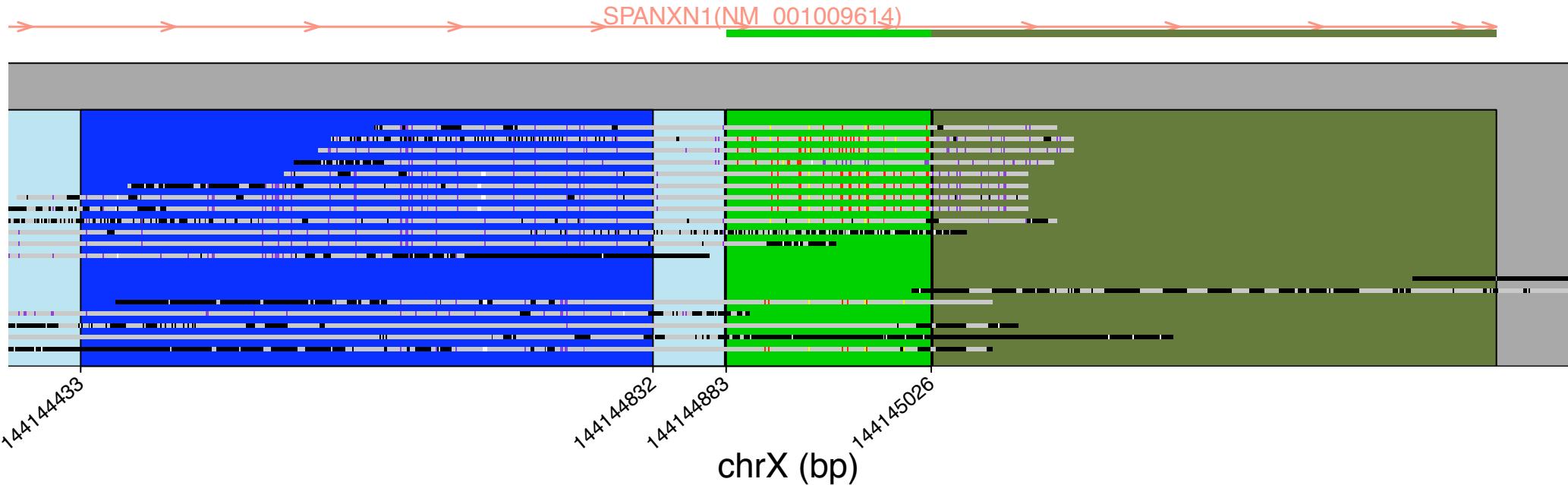
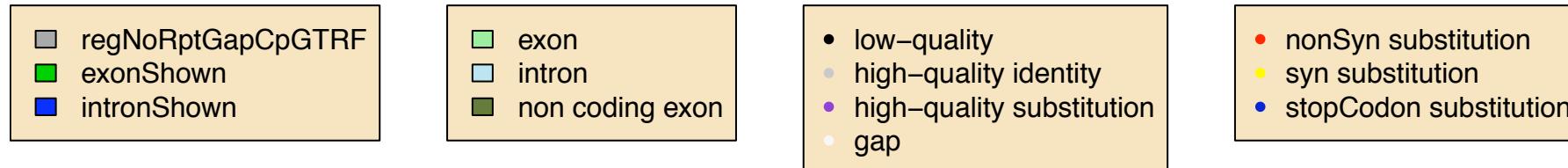
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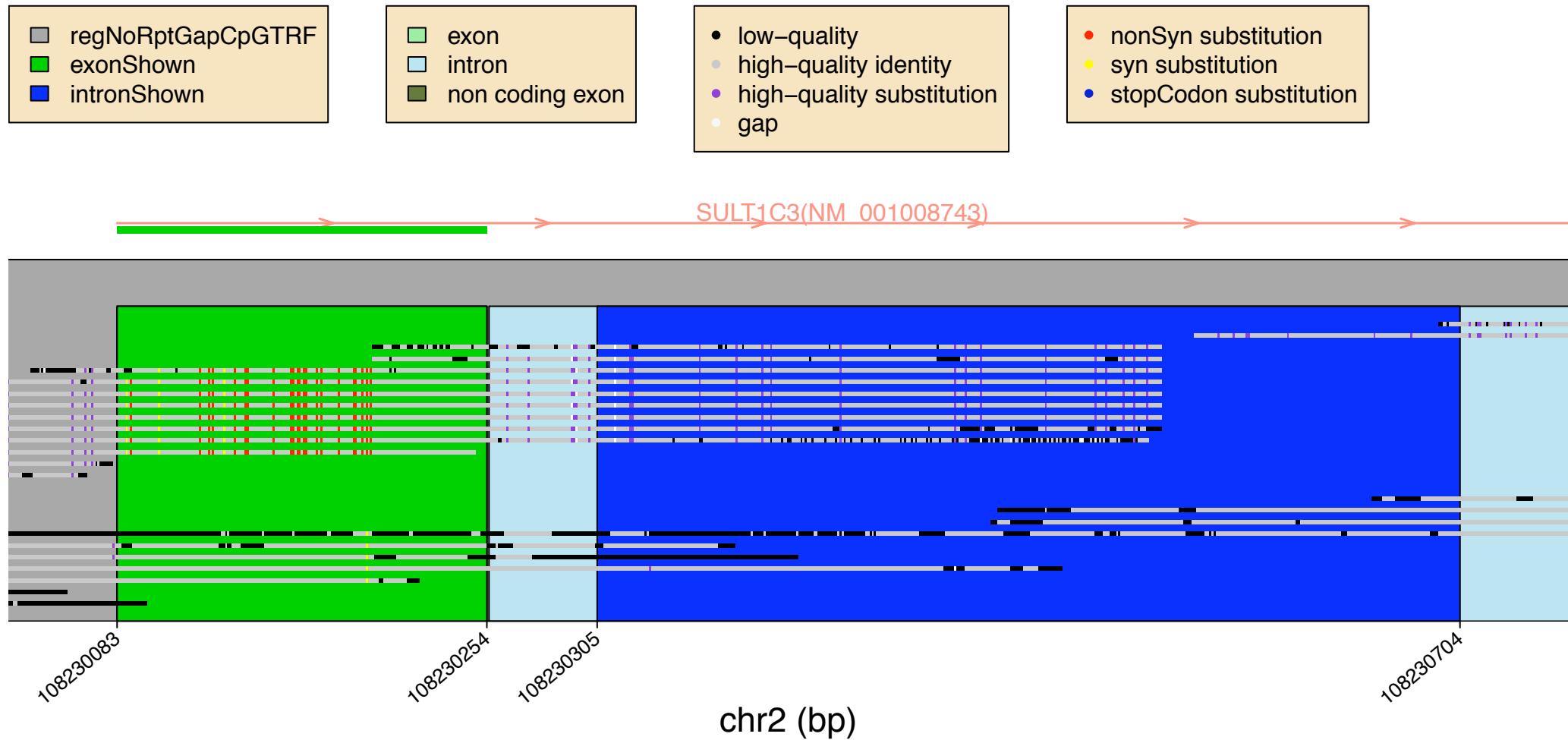
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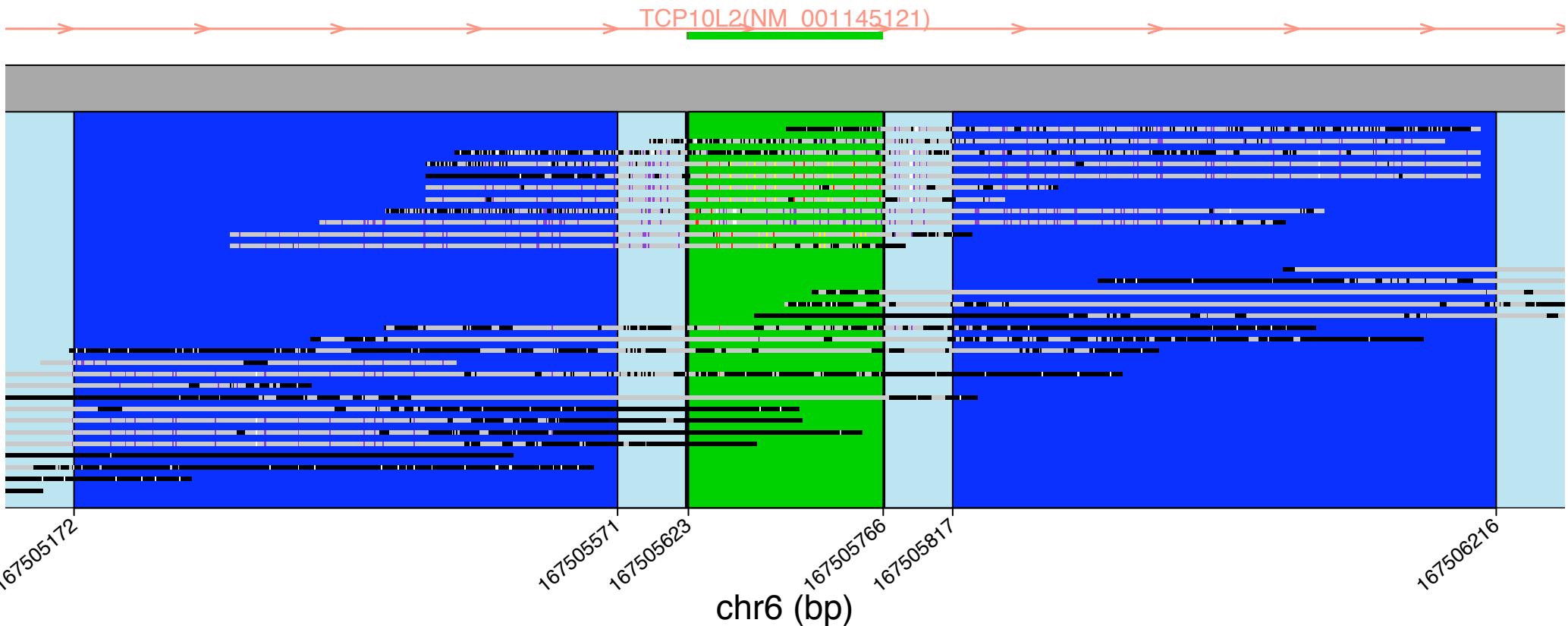
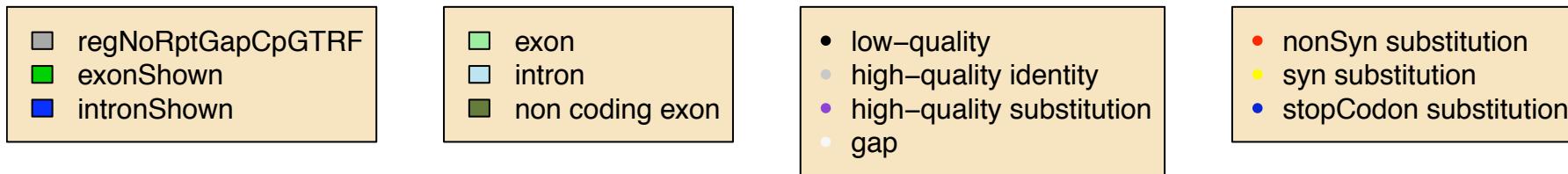
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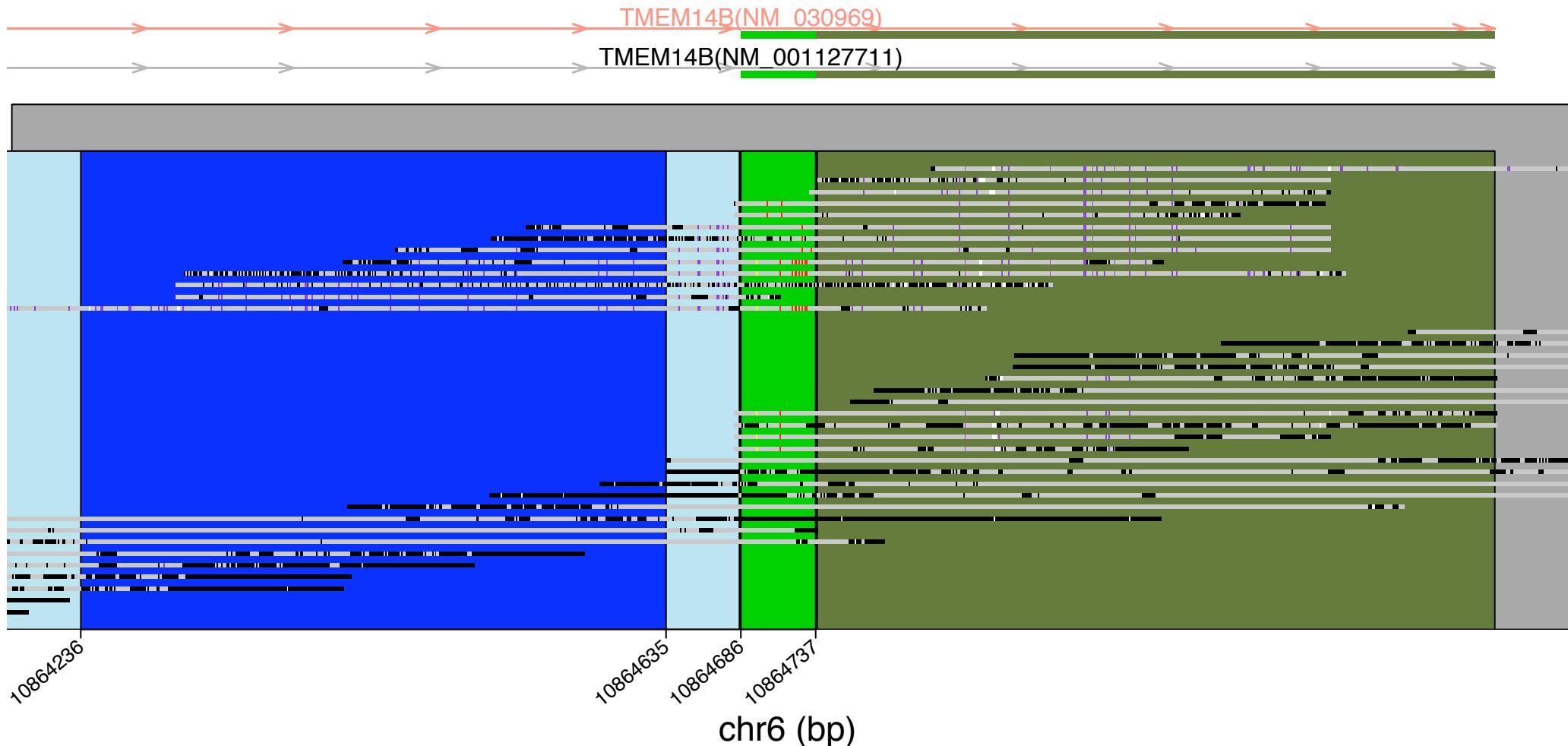
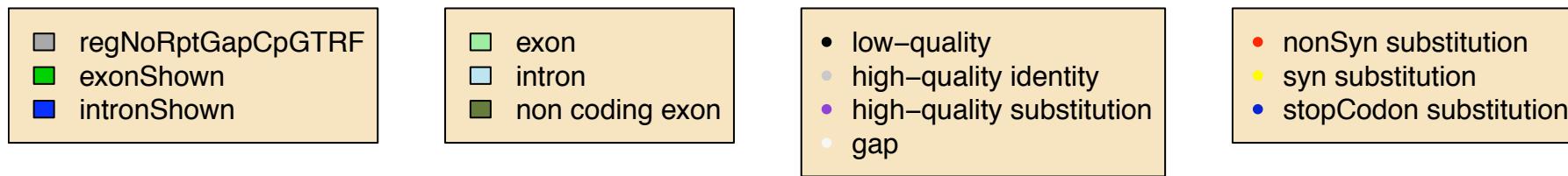
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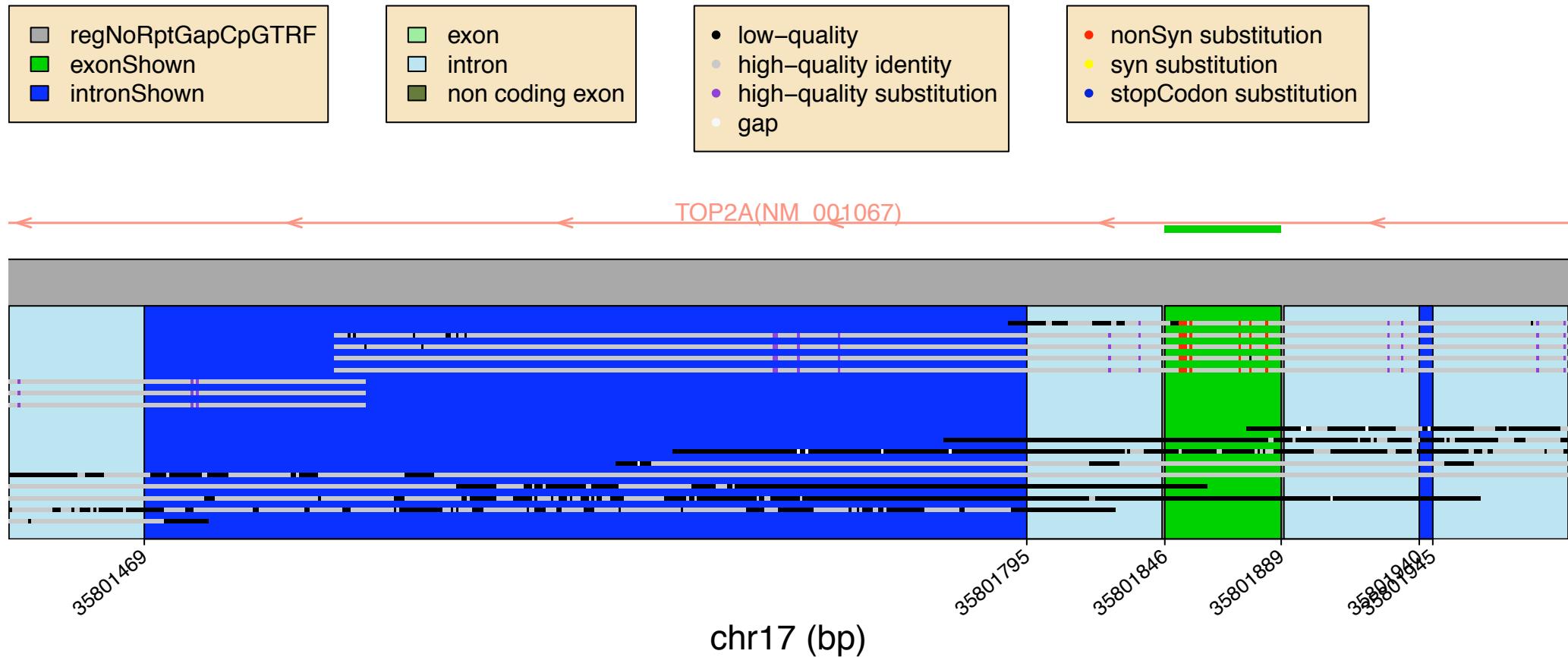
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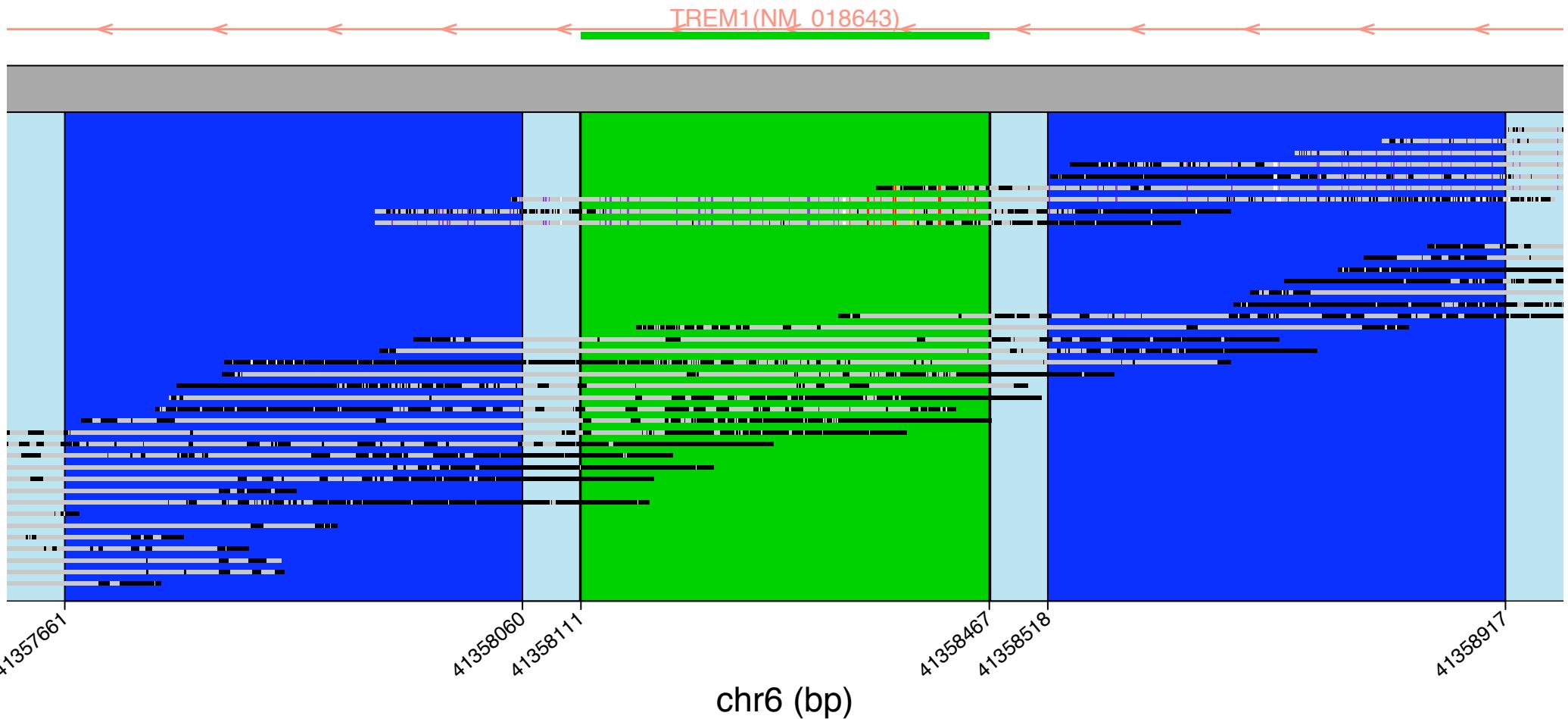
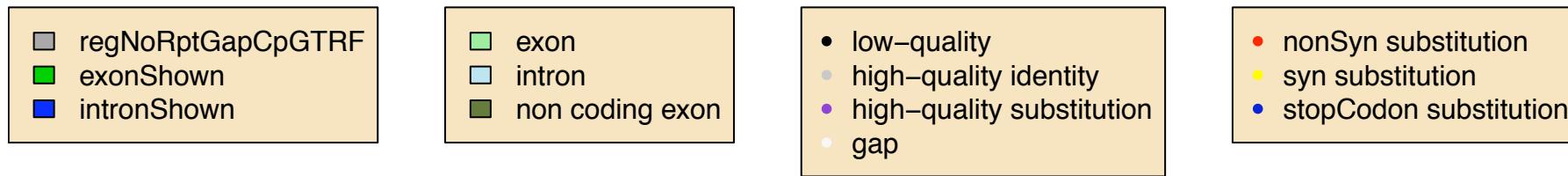
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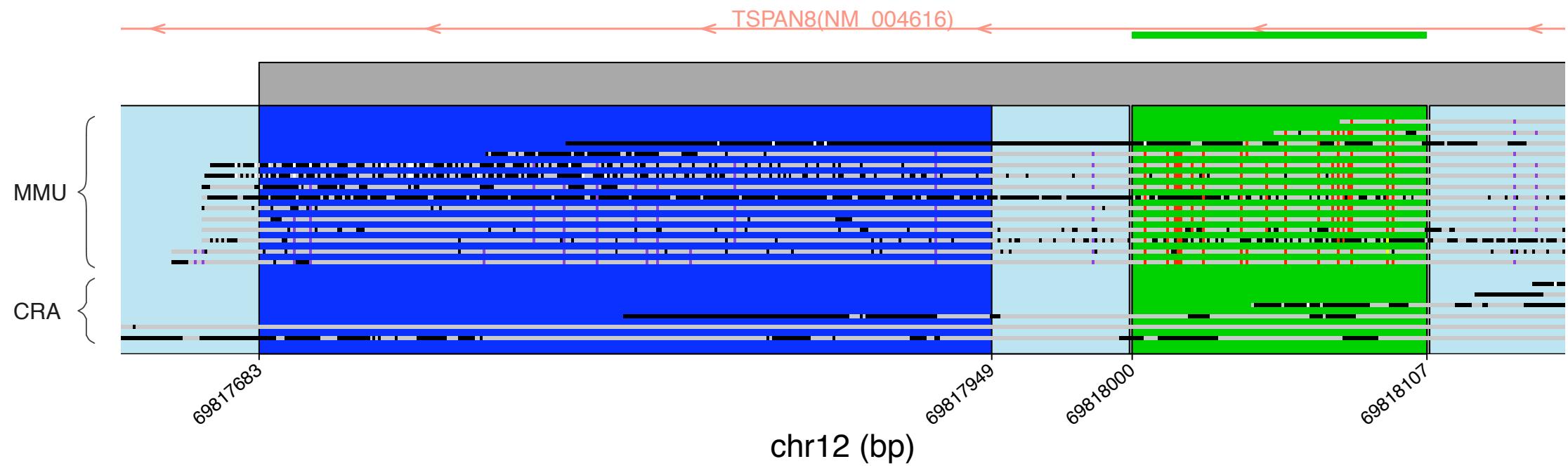
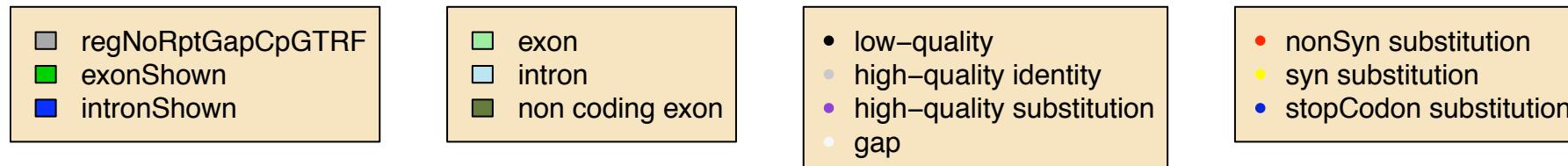
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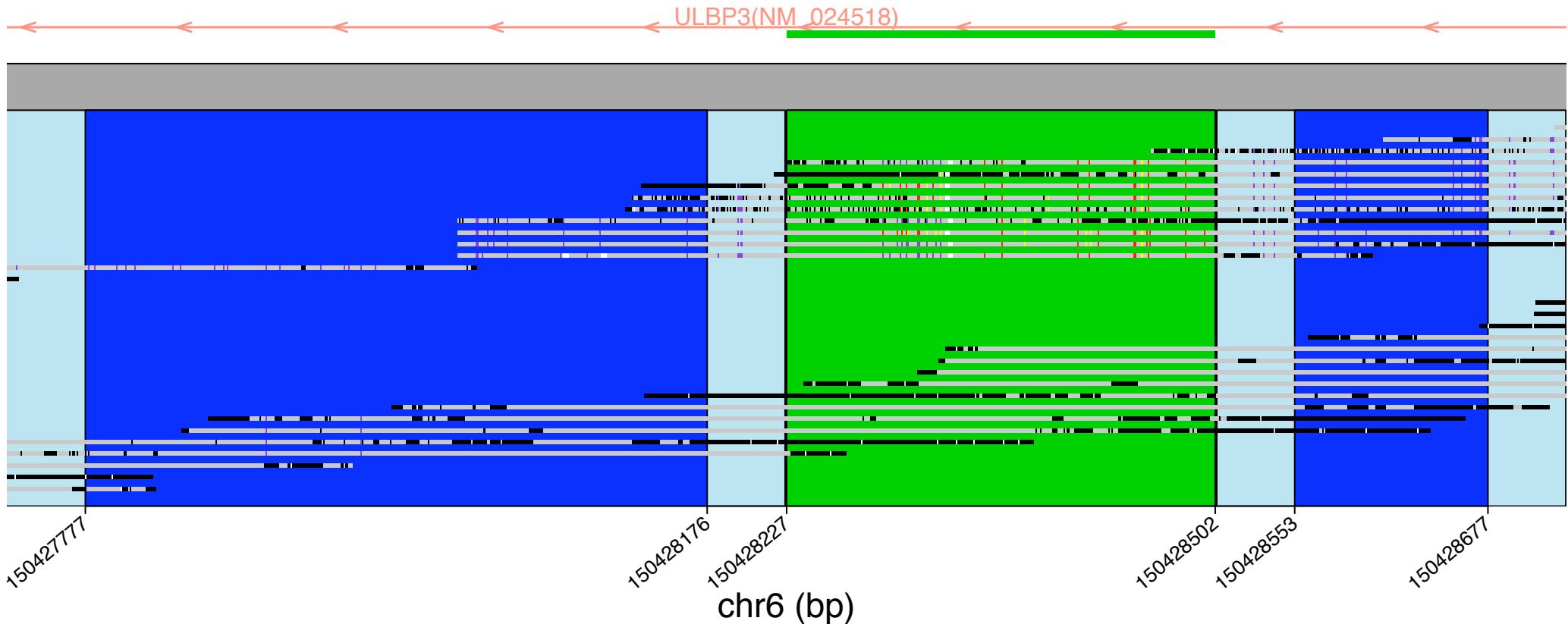
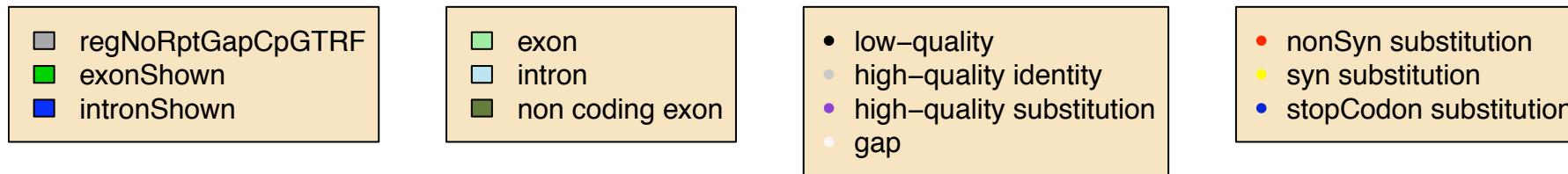
# TREM1\_NM\_018643\_41351689–41362435\_chr6\_exon2



# TSPAN8\_NM\_004616\_69805143–69838046\_chr12\_exon6



# ULBP3\_NM\_024518\_150427435-150431895\_chr6\_exon3



# VWA5A\_NM\_014622\_123491320–123522828\_chr11\_exon15

