

sequence1 OR227935.1:1-157902

Alignment 1
sequence2
OR227937.1 (+)
1-157559
Criteria: 70%, 100 bp
Regions: 289

Alignment 2
sequence3
OR227936.1 (+)
1-157450
Criteria: 70%, 100 bp
Regions: 288

Alignment 3
sequence4
MH191388.1 (+)
1-157298
Criteria: 70%, 100 bp
Regions: 293

Alignment 4
sequence5
NC_062156.1 (+)
3-157848
Criteria: 70%, 100 bp
Regions: 287

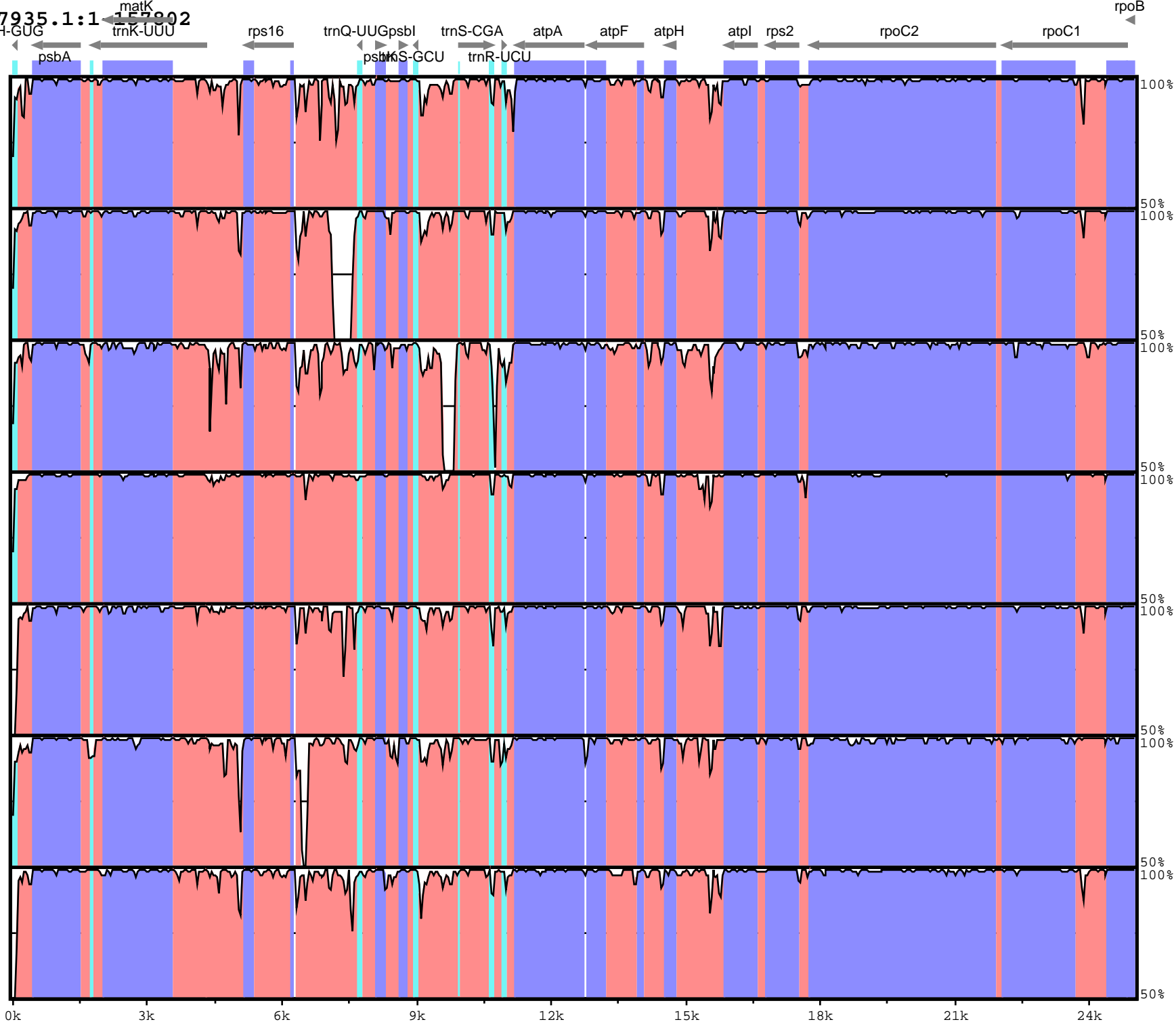
Alignment 5
sequence6
MN496075.1 (+)
1-157715
Criteria: 70%, 100 bp
Regions: 286

Alignment 6
sequence7
MT081309.1 (+)
1-157152
Criteria: 70%, 100 bp
Regions: 291

Alignment 7
sequence8
MN496076.1 (+)
1-157781
Criteria: 70%, 100 bp
Regions: 286

X-axis: sequence1
Resolution: 39
Window size: 100 bp

- gene
- exon
- UTR
- CNS
- mRNA



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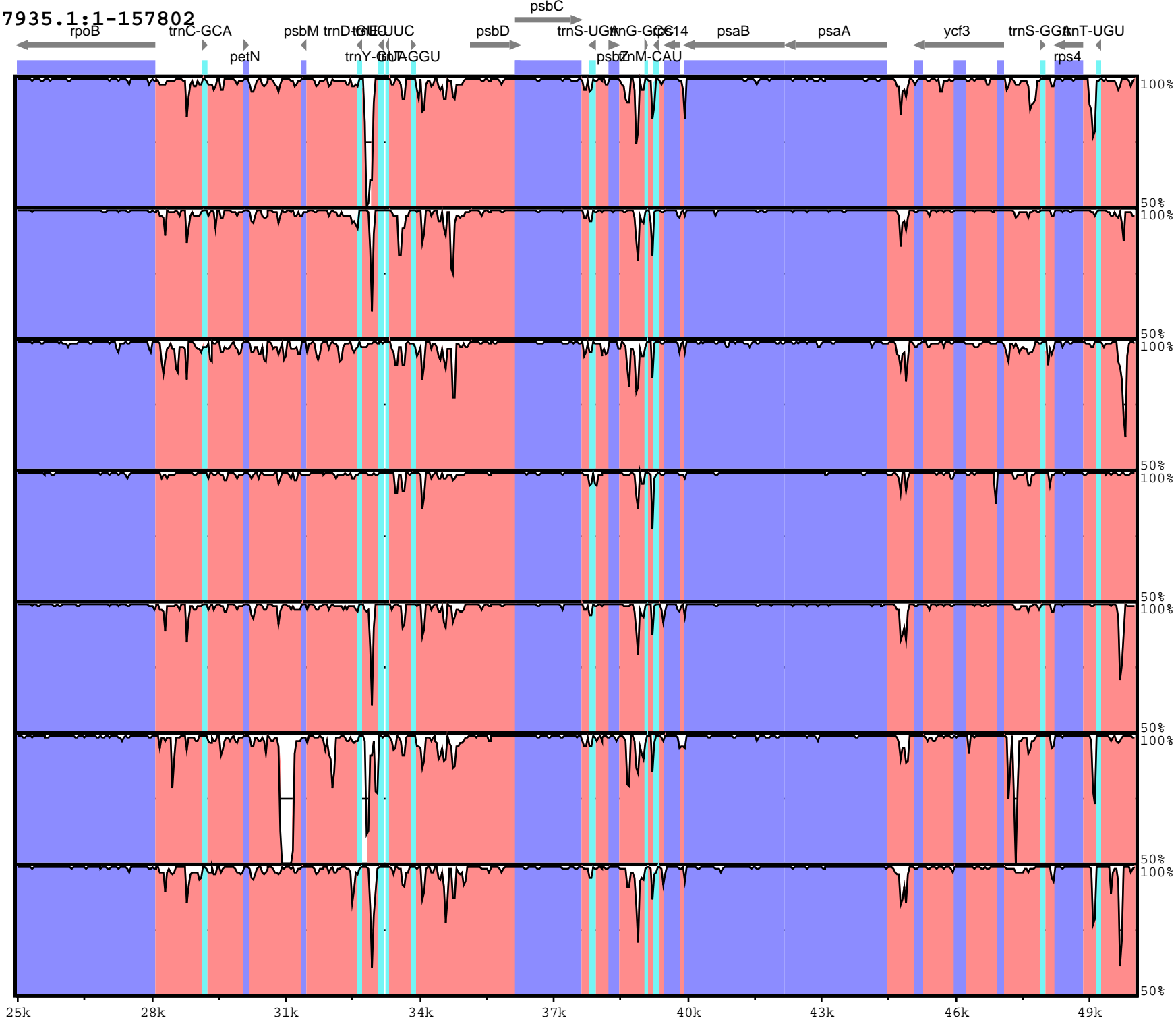
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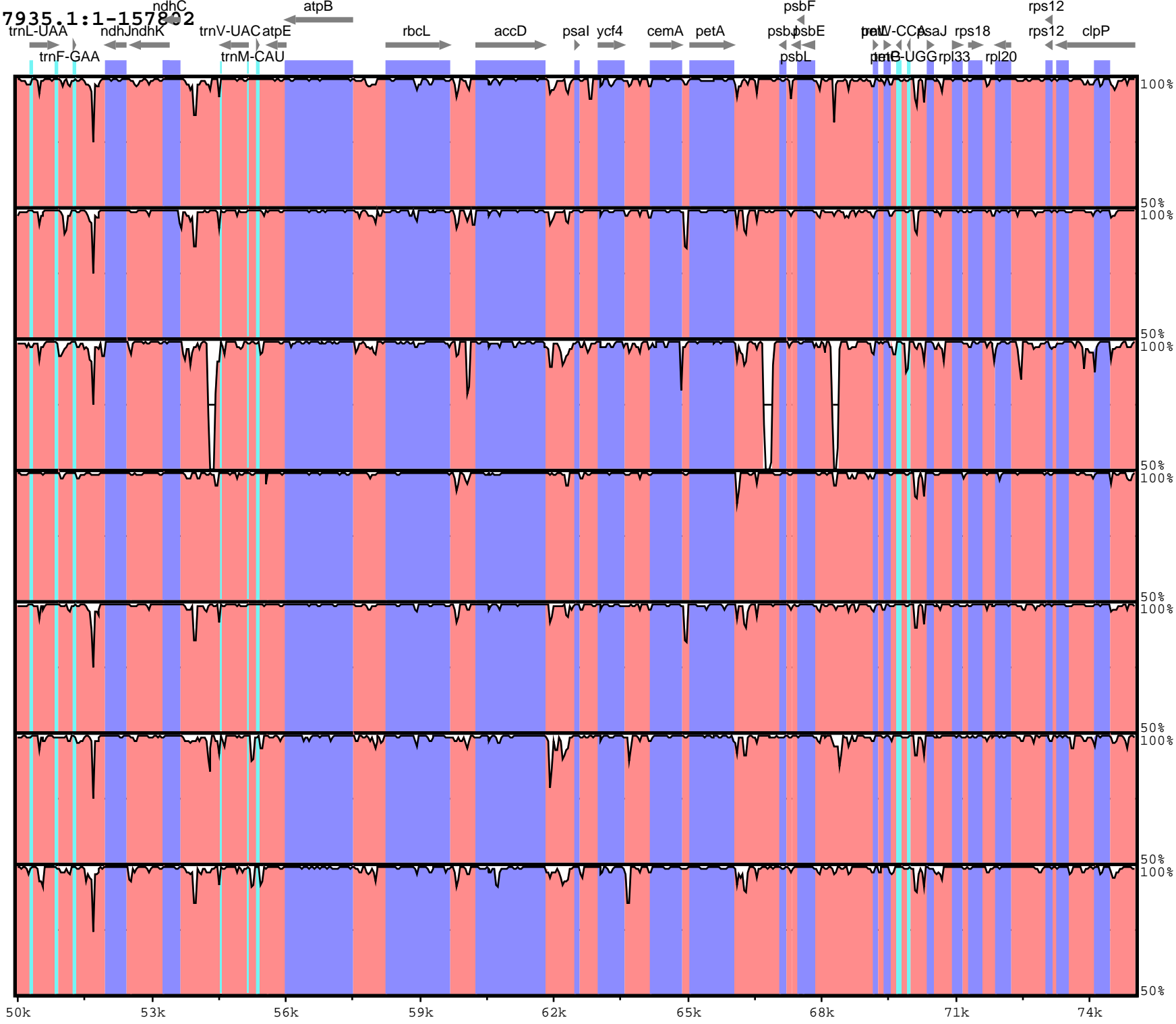
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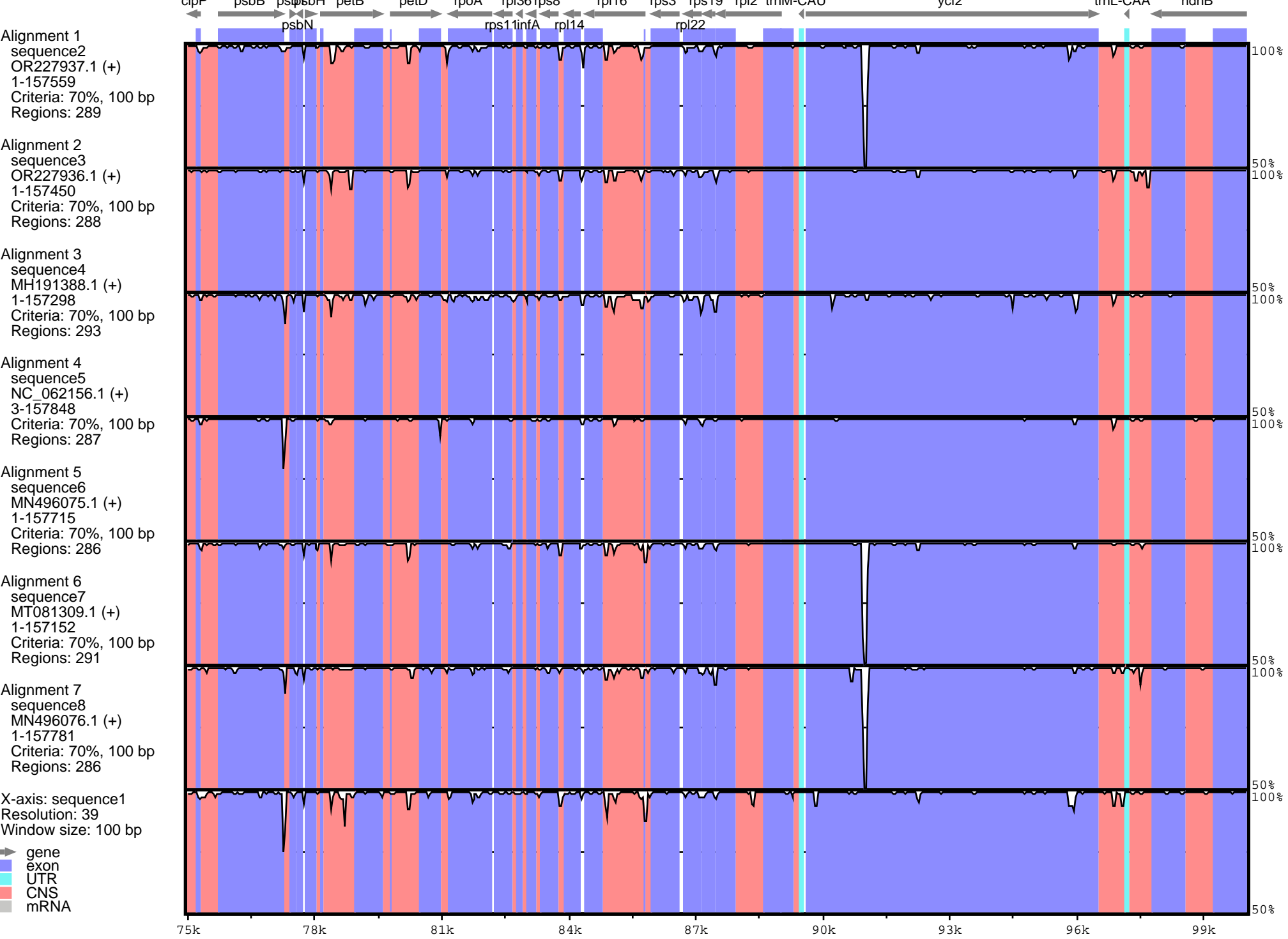
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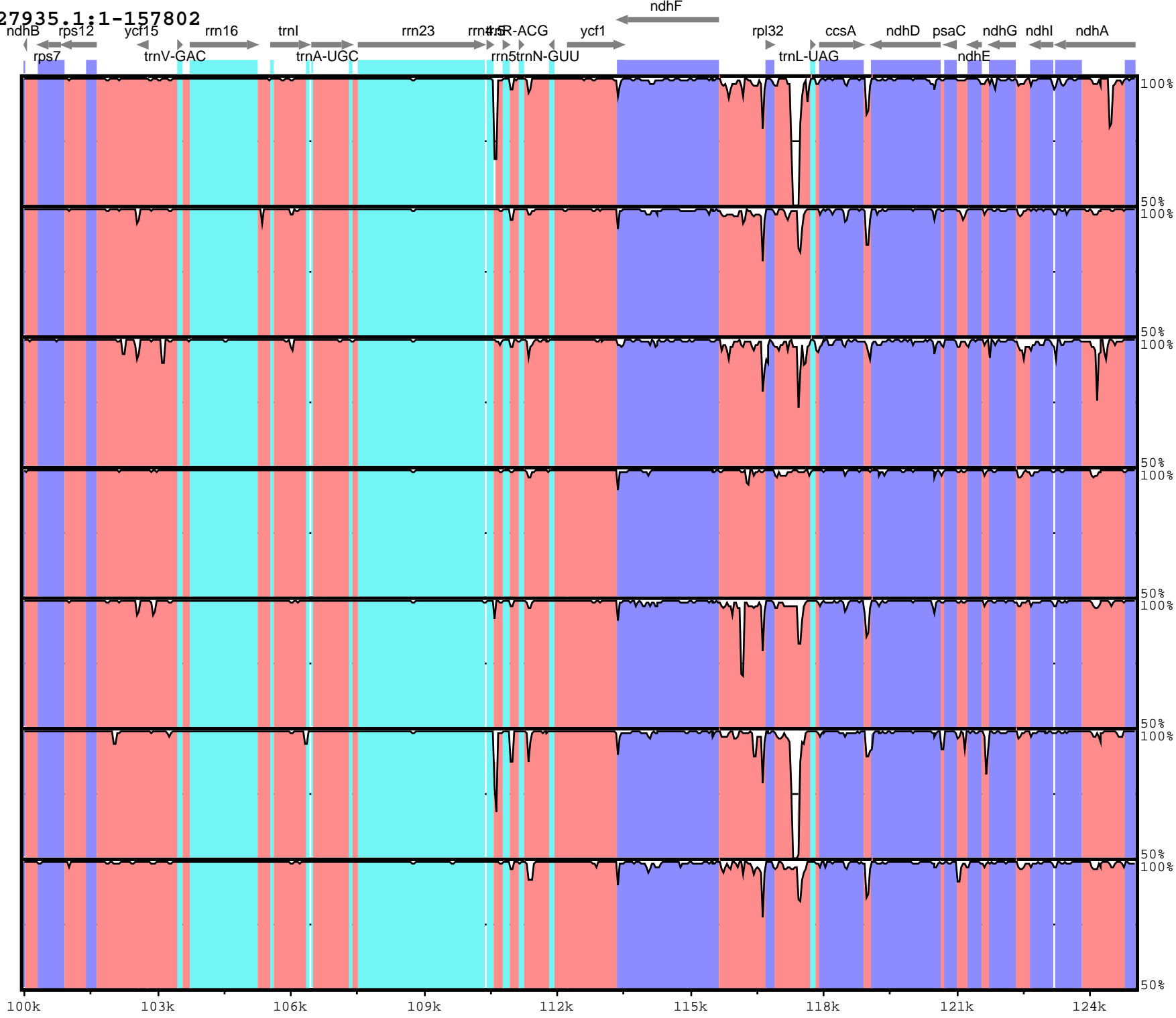
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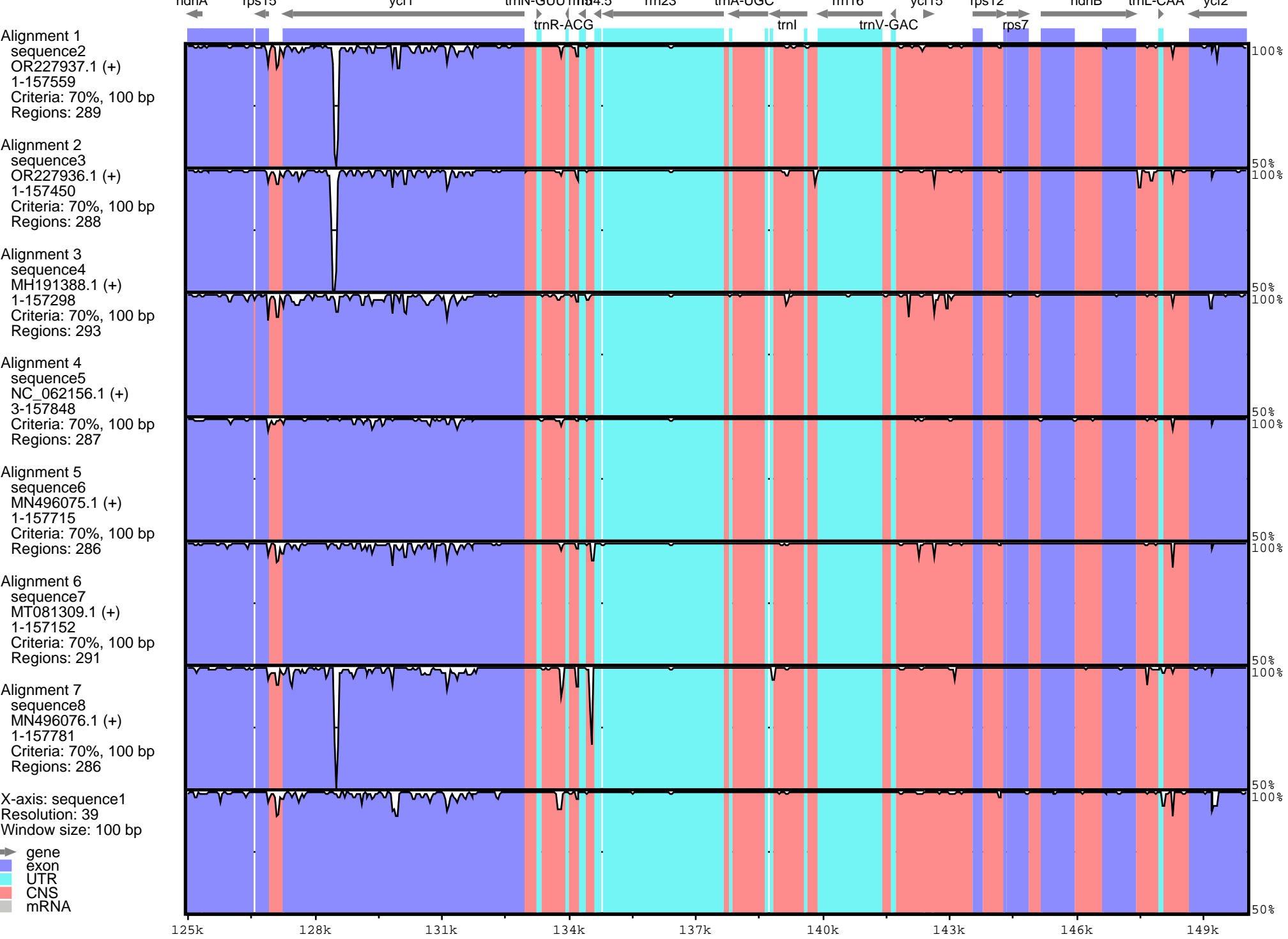
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