

Reference: *Laurus nobilis*

Criteria: 70%, 100 bp;
Regions: 275

- contig
- gene
- exon
- UTR
- CNS
- mRNA

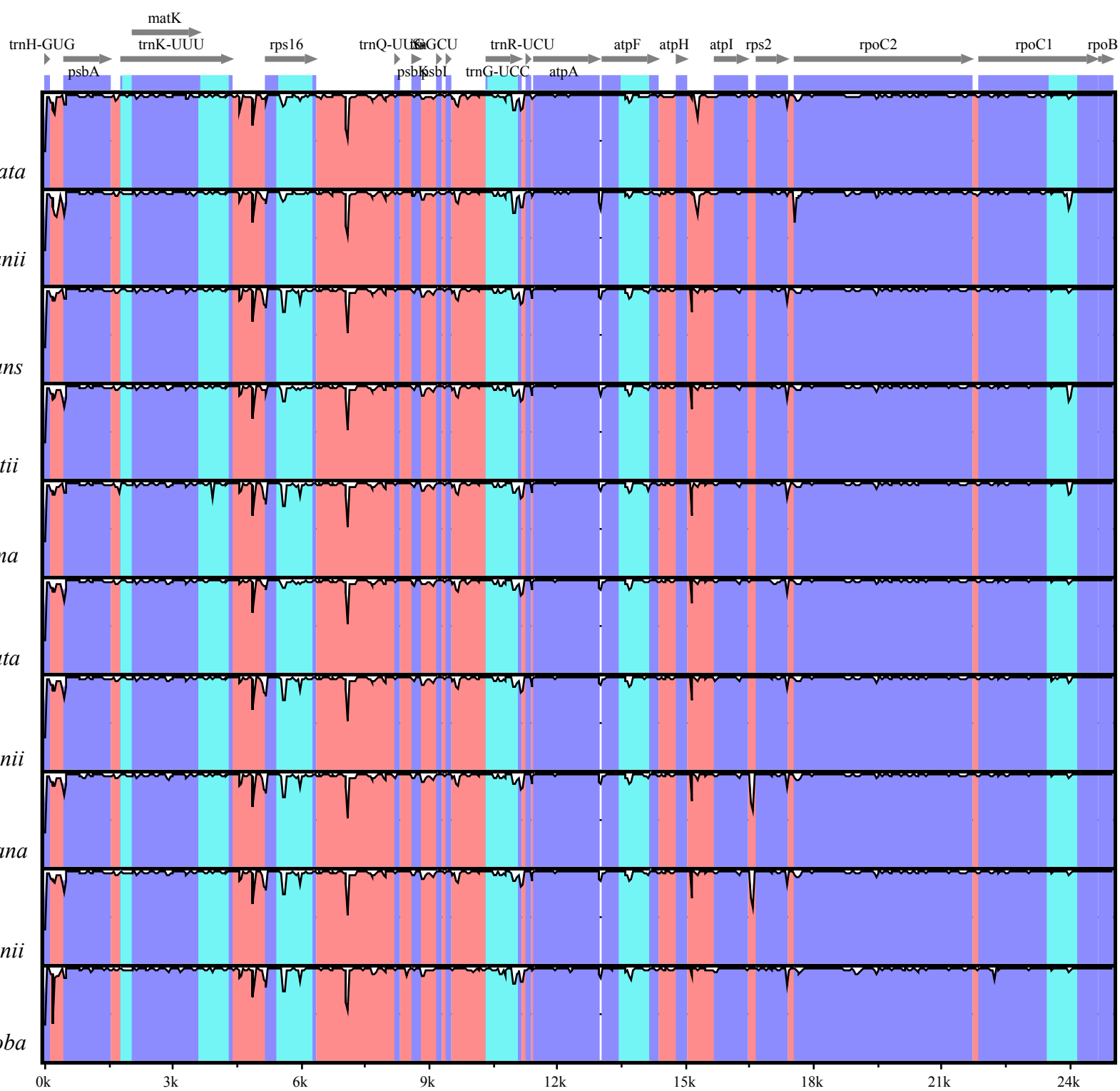
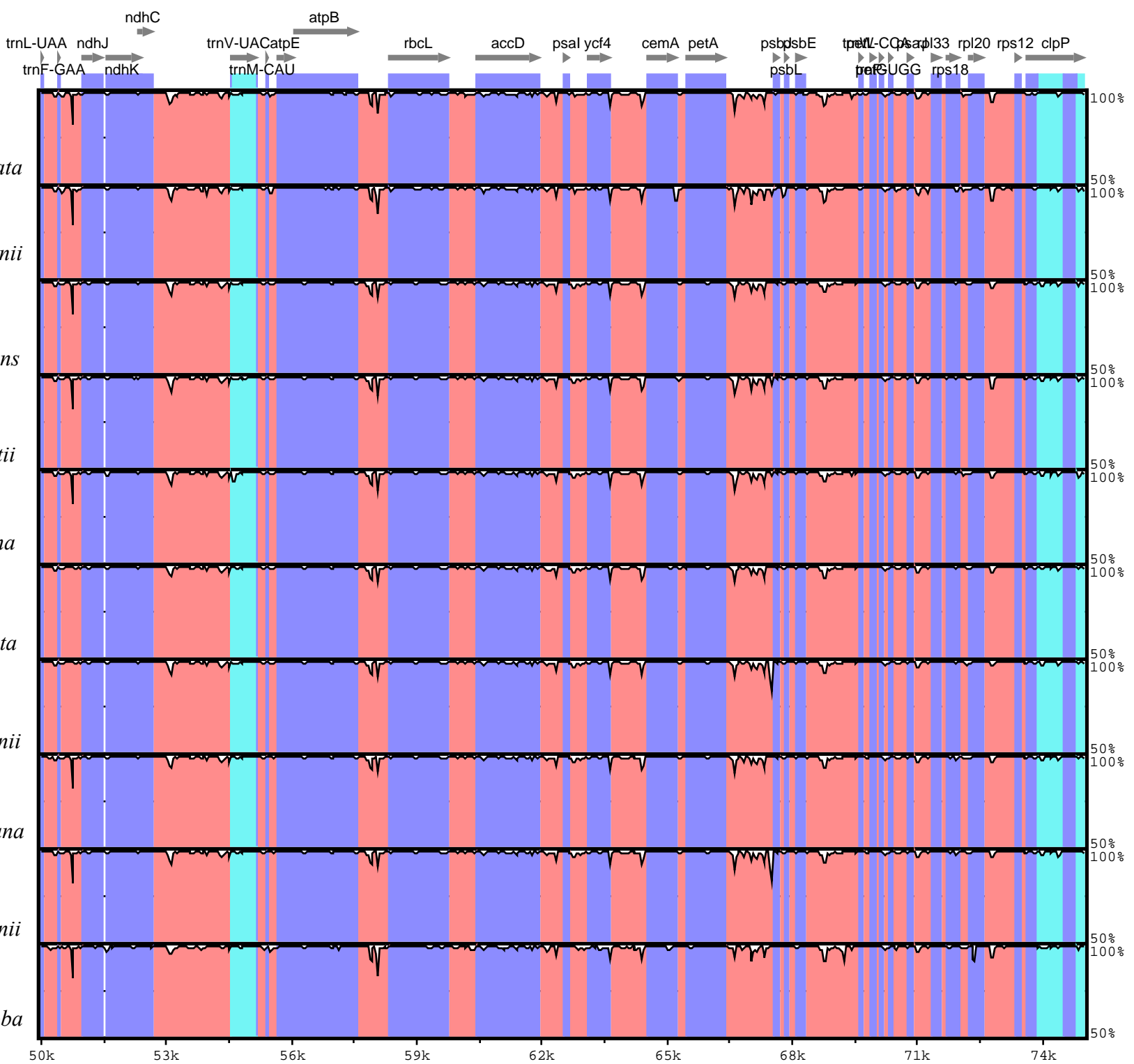


Fig S4. VISTA-based identity plots showed sequence identity of *Lindera* sequenced chloroplast genomes with *Laurus nobilis* as a reference.

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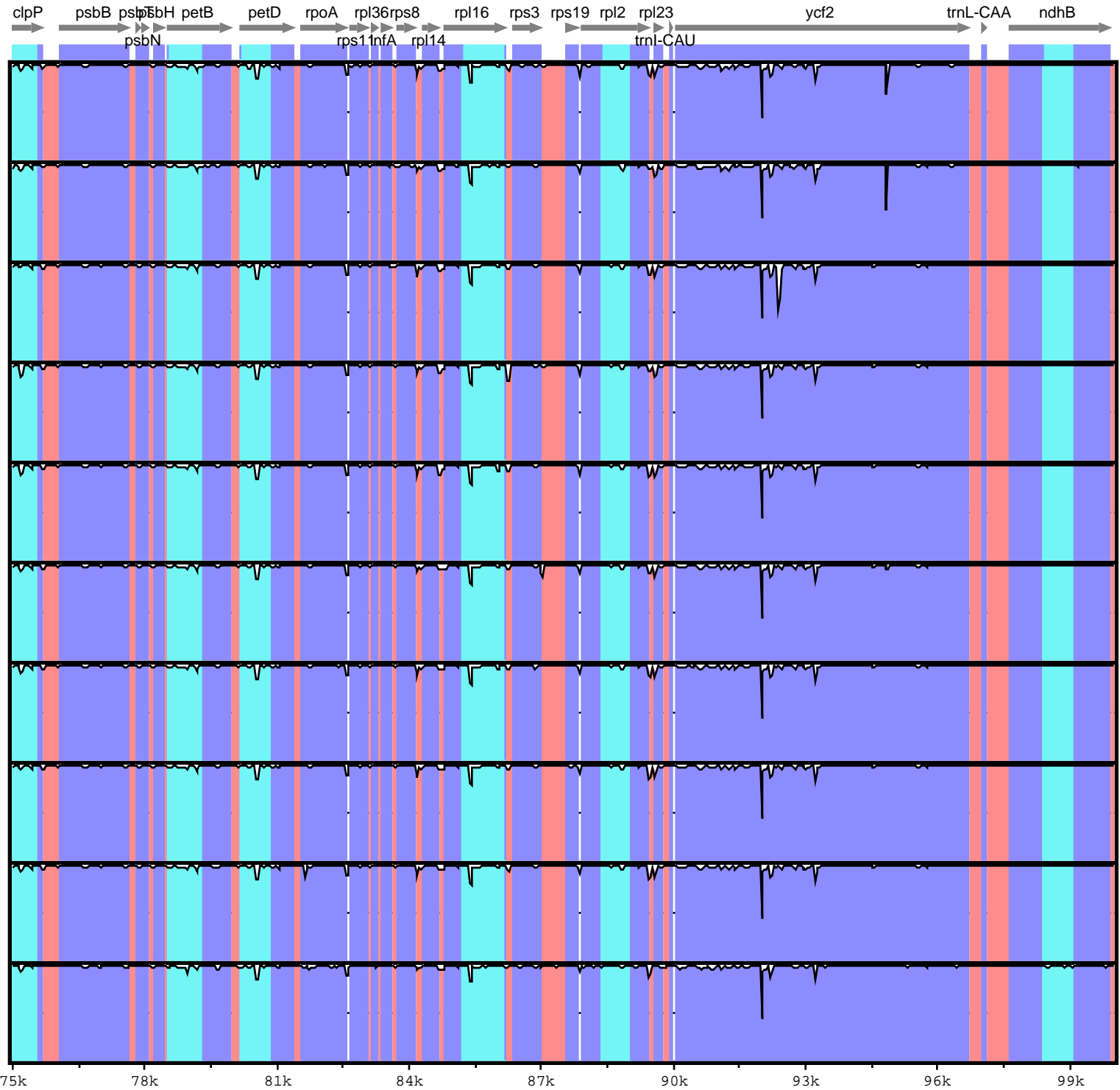
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Lindera aggregata

Lindera chunii

Lindera fragrans

Lindera limprichtii

Lindera pulcherrima

Lindera supracostata

Lindera thomsonii

Lindera thomsonii var. *vernayana*

Lindera thomsonii

Lindera obtusiloba

75k

78k

81k

84k

87k

90k

93k

96k

99k

50%

50%

50%

50%

50%

50%

50%

50%

50%

50%

100%

100%

100%

100%

100%

100%

100%

100%

100%

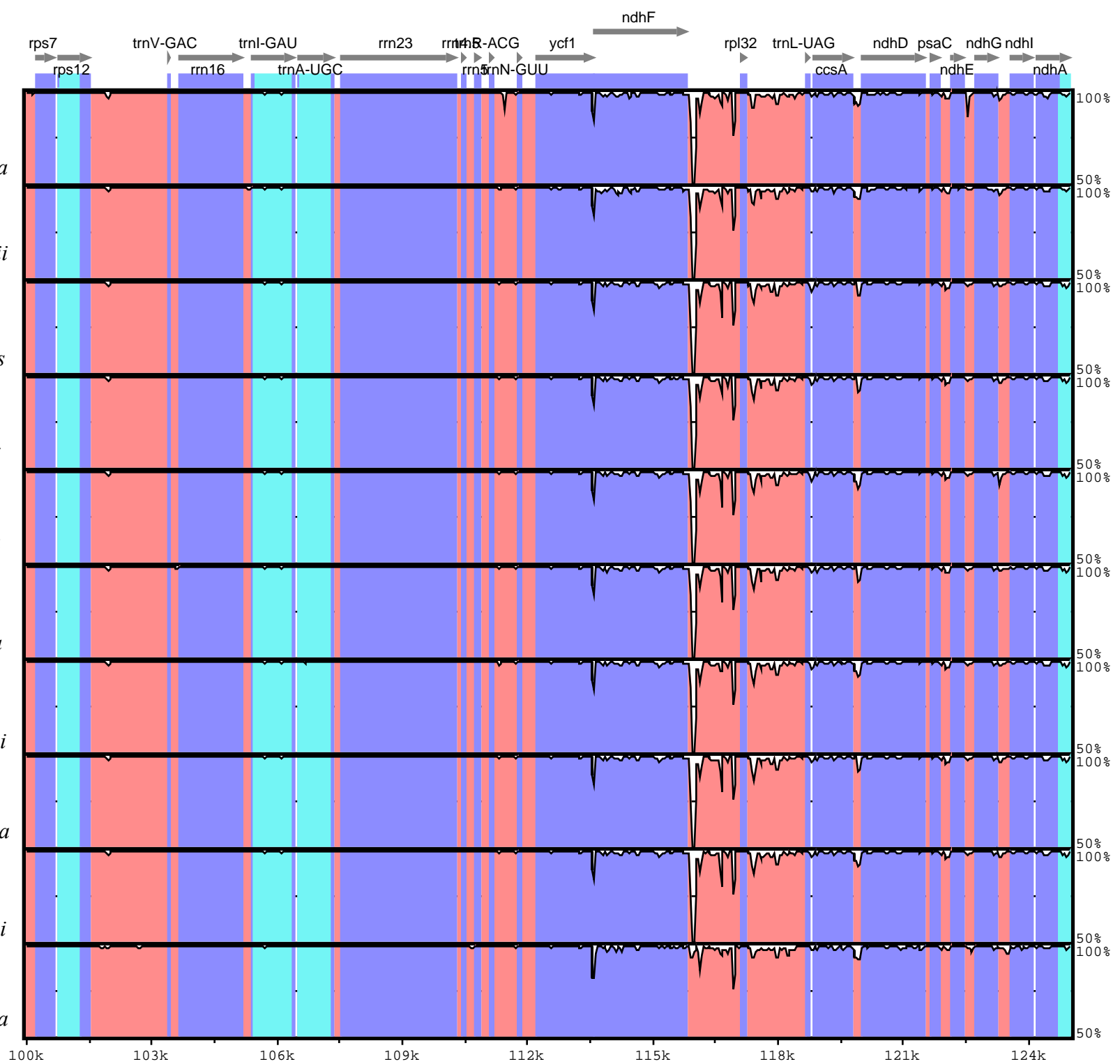
100%

50%

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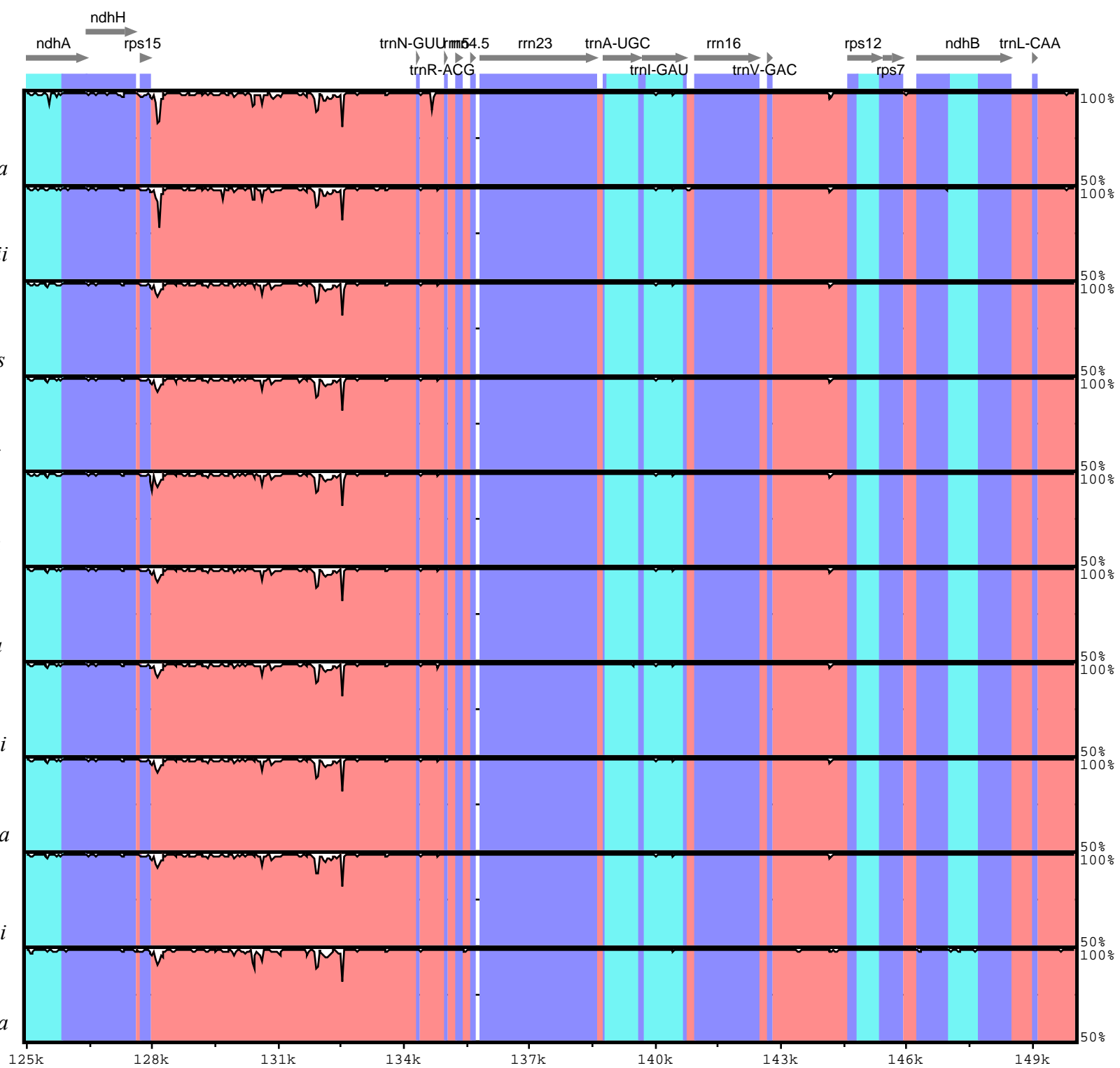
- ▶ contig
- ▶ gene
- exon
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