MAVID Alignment - 2ENSMUSG00000027168proteincodi vs. All

Alignment 1
  Seqs: 2ensmusg00000027168proteincodi/11ensg00000007372proteincoding
  Criteria: 80%, 100 bp
  Regions: 38

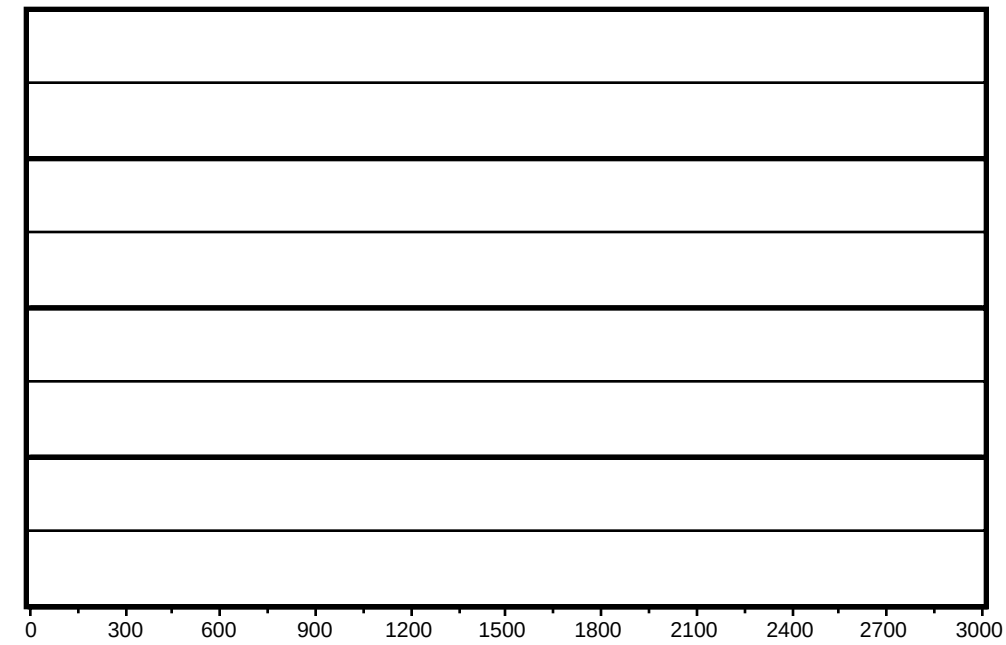
Alignment 2
  Seqs: 2ensmusg00000027168proteincodi/scaffold309newsinfrug000001215
  Criteria: 71%, 100 bp
  Regions: 10

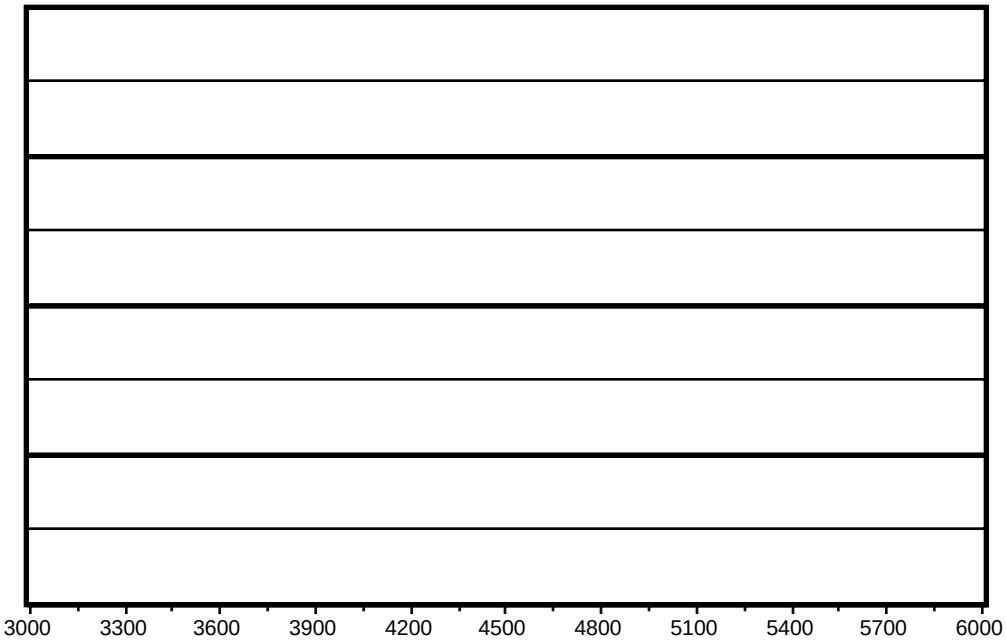
Alignment 3
  Seqs: 2ensmusg00000027168proteincodi/3ensgalg00000012123proteincodi
  Criteria: 74%, 100 bp
  Regions: 18

Alignment 4
  Seqs: 2ensmusg00000027168proteincodi/scaffold430newsinfrug000001506
  Criteria: 67%, 100 bp
  Regions: 6

X-axis: 2ensmusg00000027168proteincodi
Resolution: 25
Window size: 100 bp

gene
exon
UTR
CNS





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X-axis: 2ensmusg00000027168proteincodi
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Window size: 100 bp

gene
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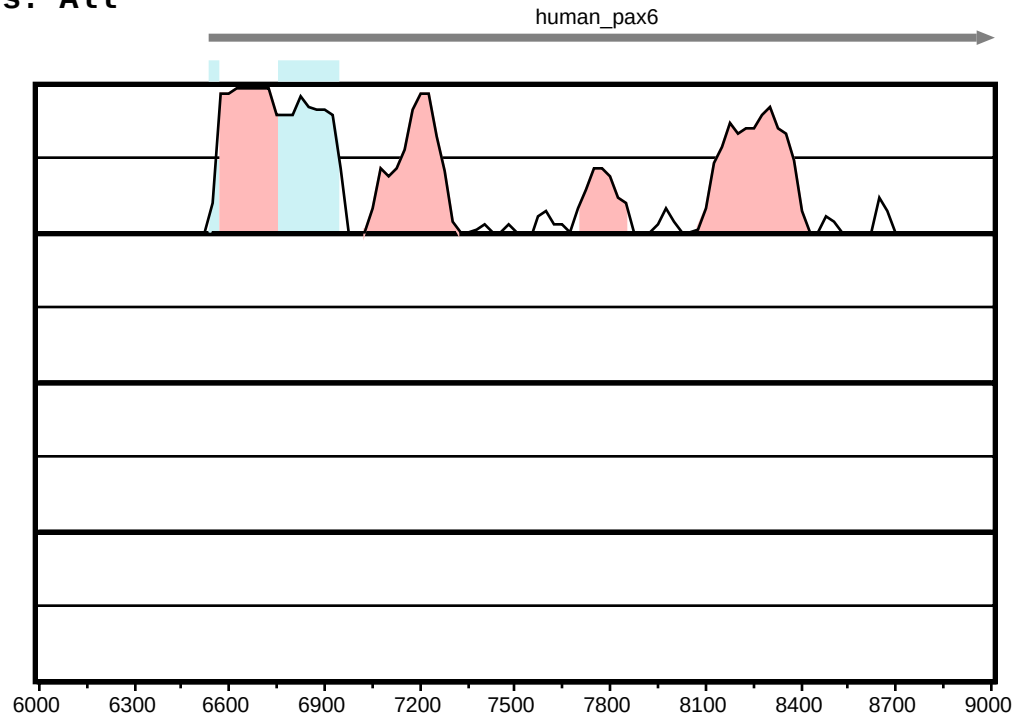

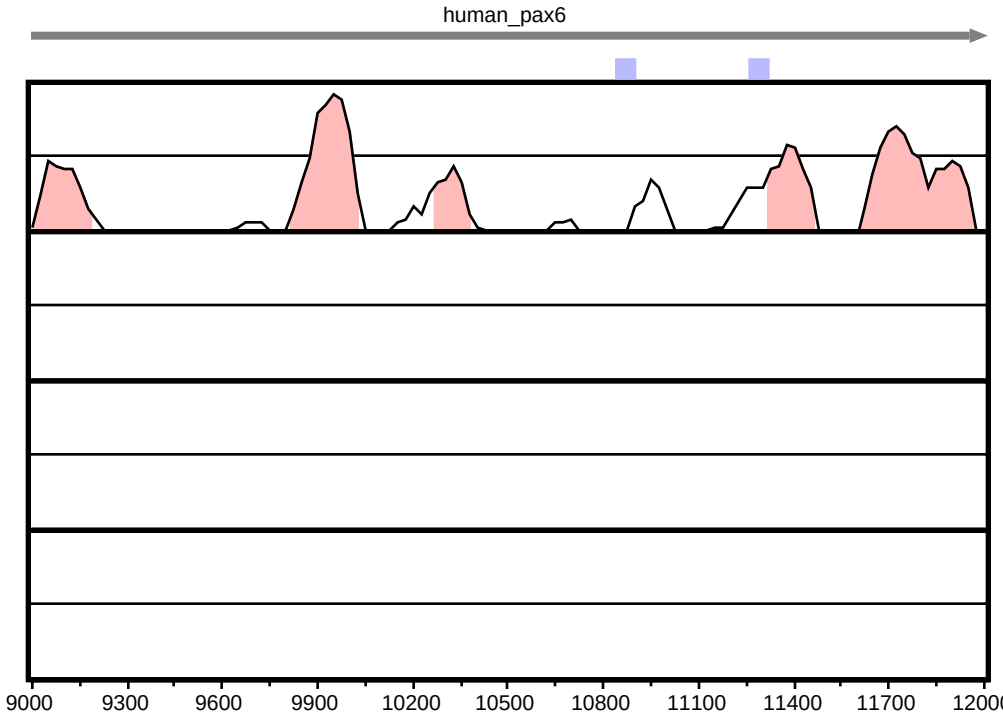

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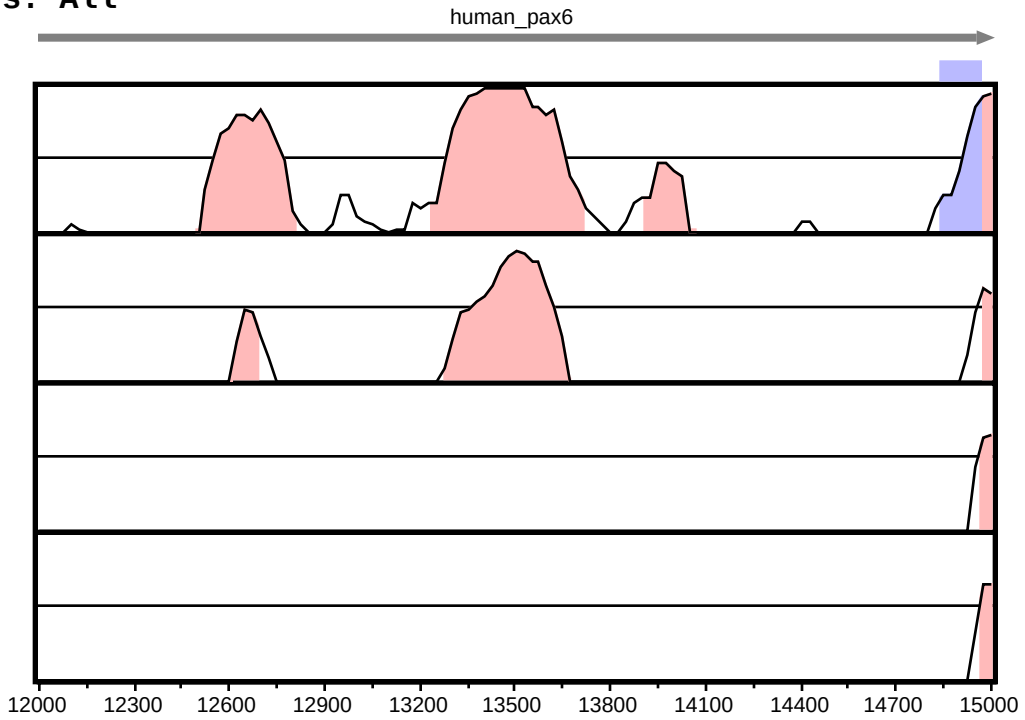

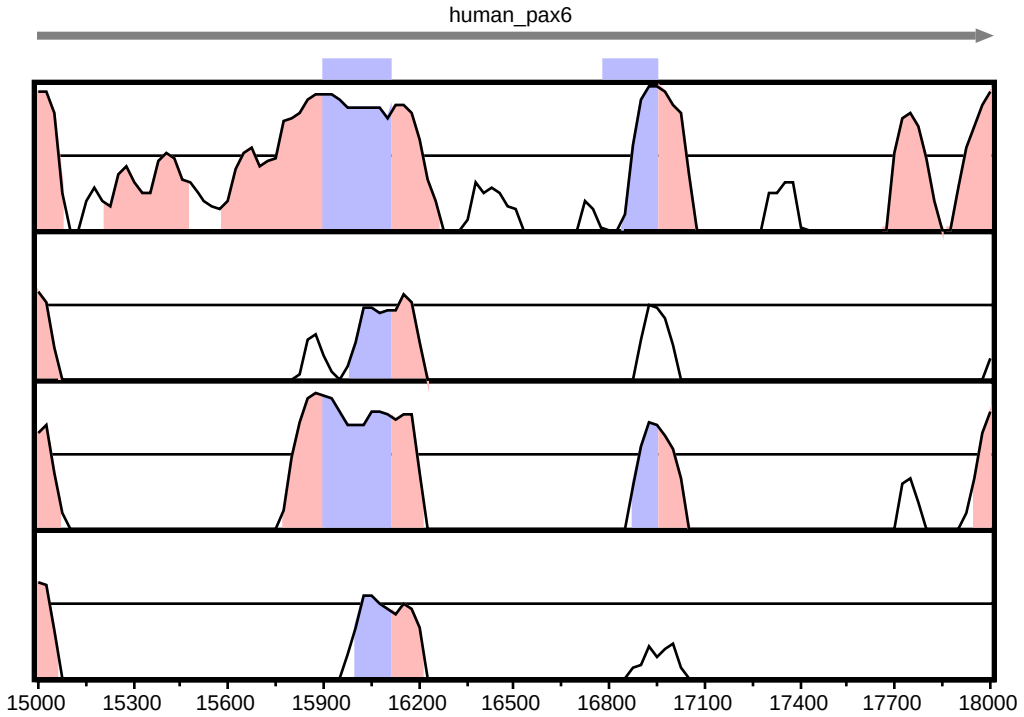

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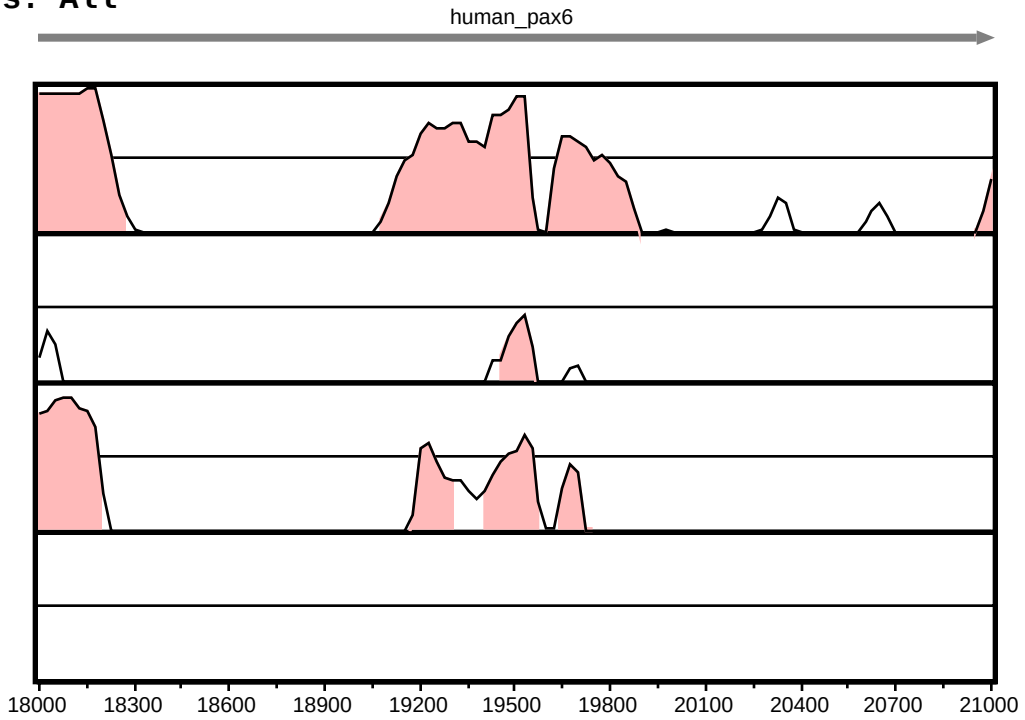

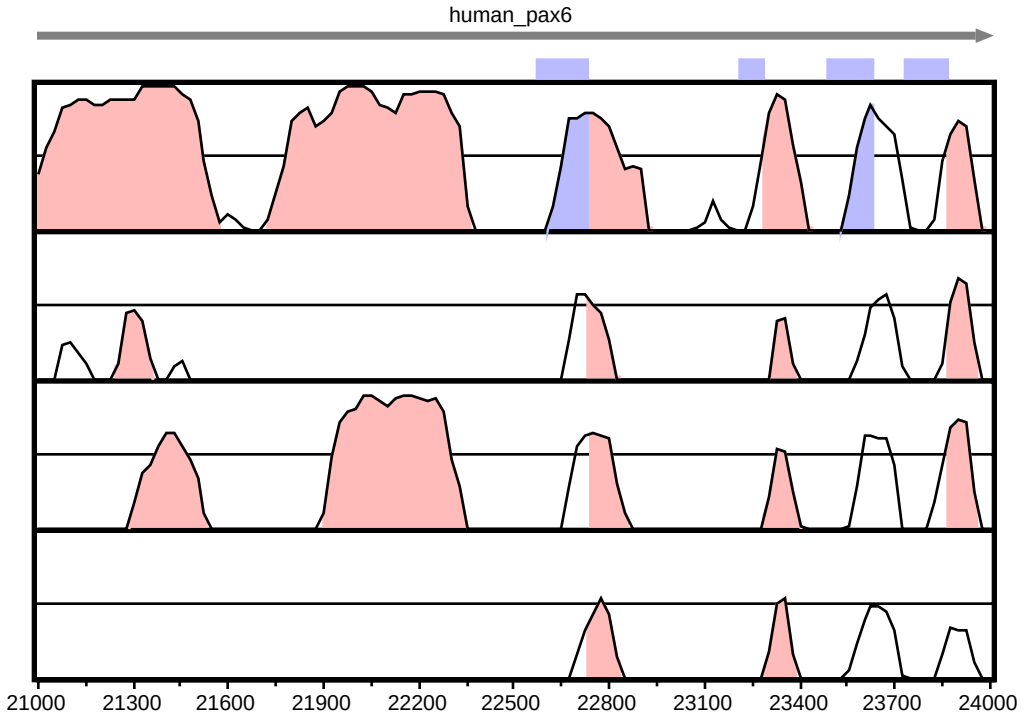

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