MAVID Alignment - 11ENSG00000007372proteincoding vs. All

Alignment 1
  Seqs: 11ensg00000007372proteincoding/scaffold309newsinfrug000001215
  Criteria: 71%, 100 bp
  Regions: 12

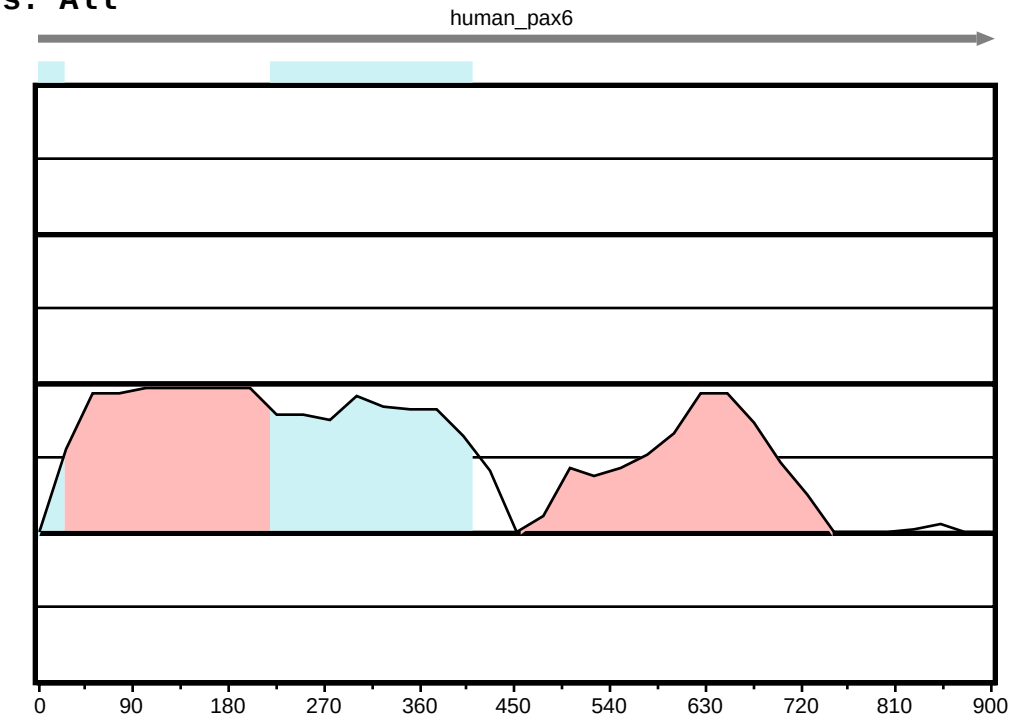
Alignment 2
  Seqs: 11ensg00000007372proteincoding/3ensgalg00000012123proteincodi
  Criteria: 73%, 100 bp
  Regions: 17

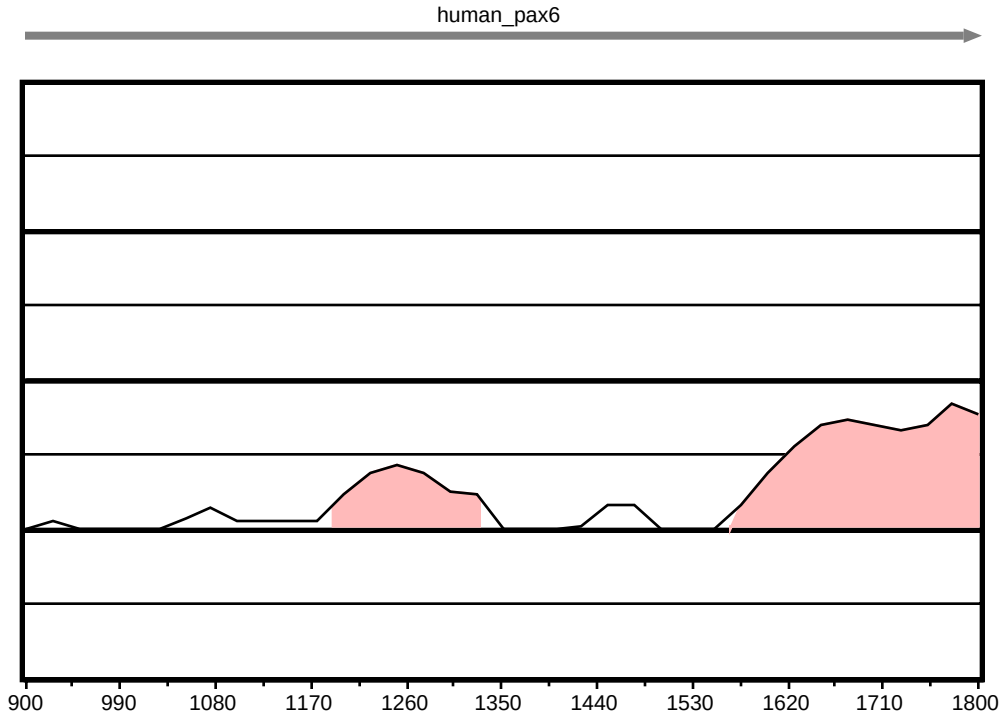
Alignment 3
  Seqs: 11ensg00000007372proteincoding/2ensmusg00000027168proteincodi
  Criteria: 80%, 100 bp
  Regions: 37

Alignment 4
  Seqs: 11ensg00000007372proteincoding/scaffold430newsinfrug000001506
  Criteria: 67%, 100 bp
  Regions: 5

X-axis: 11ensg00000007372proteincoding
Resolution: 25
Window size: 100 bp

gene
exon
UTR
CNS





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Window size: 100 bp

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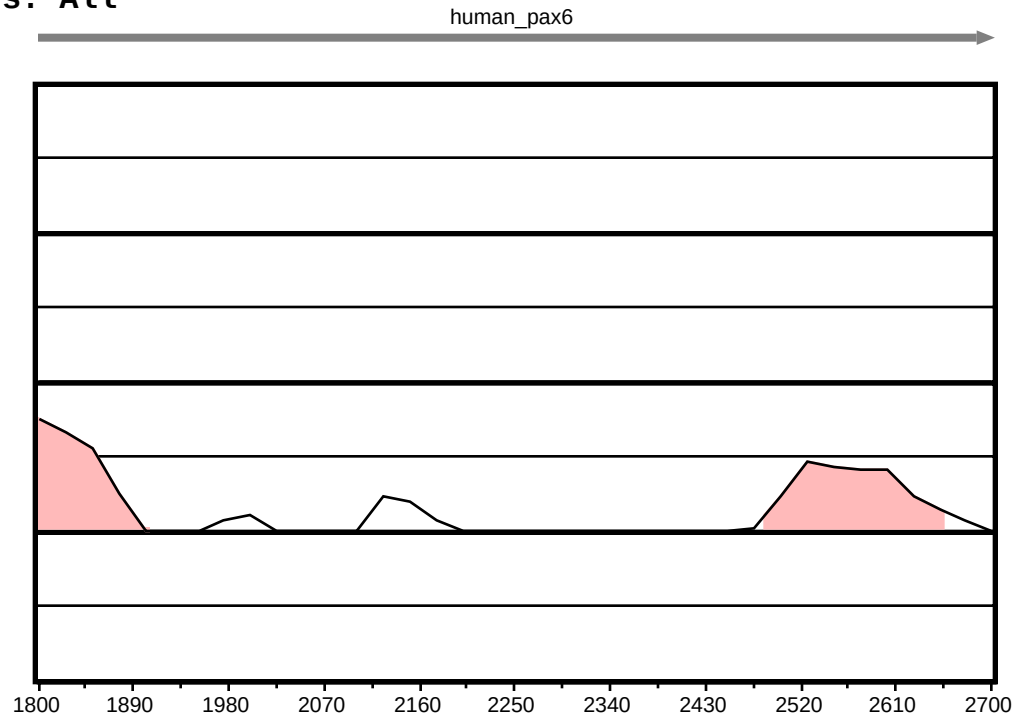

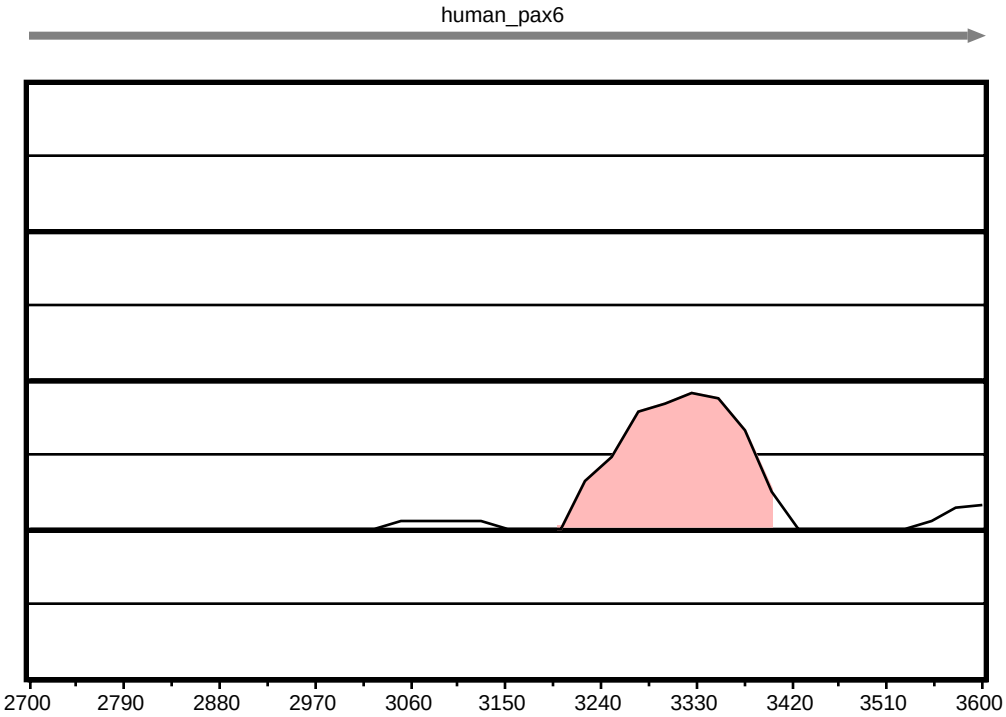

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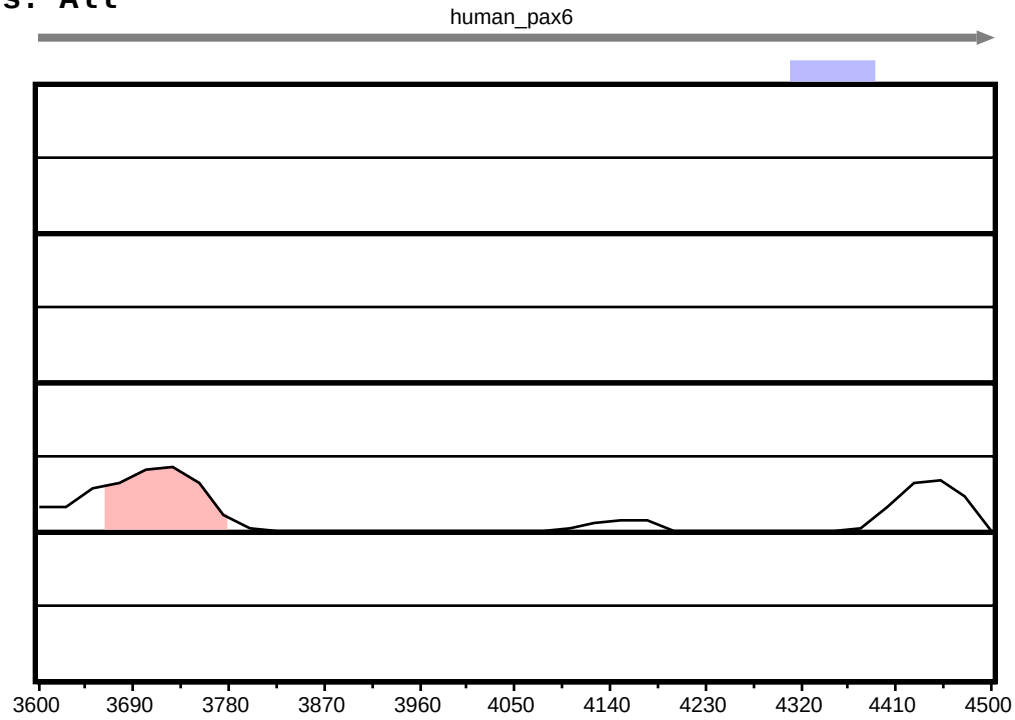

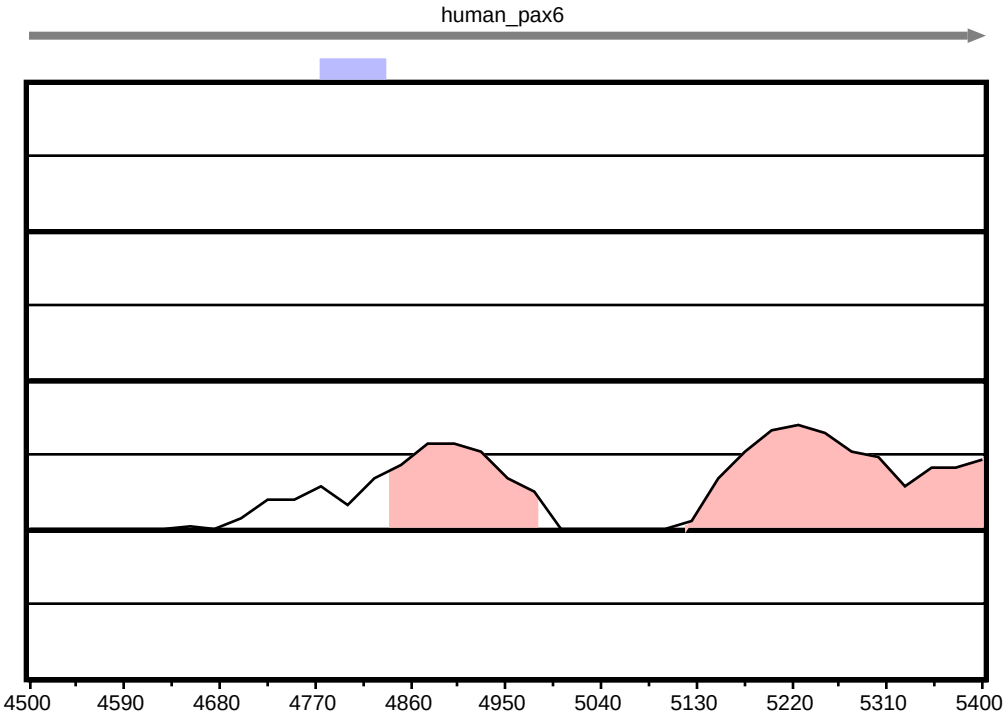

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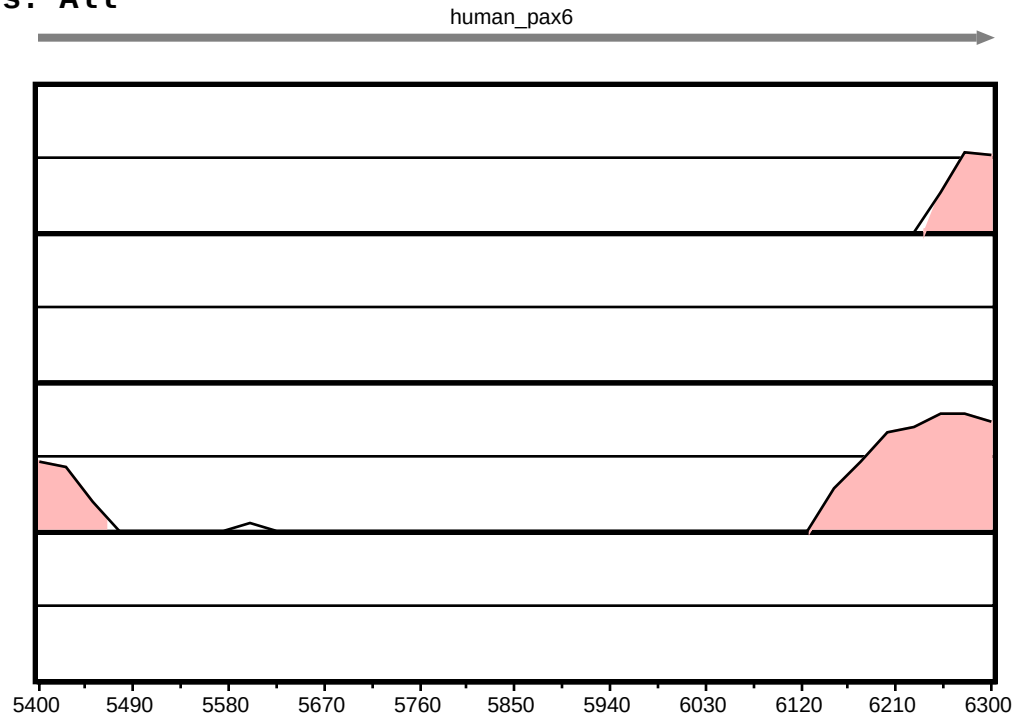

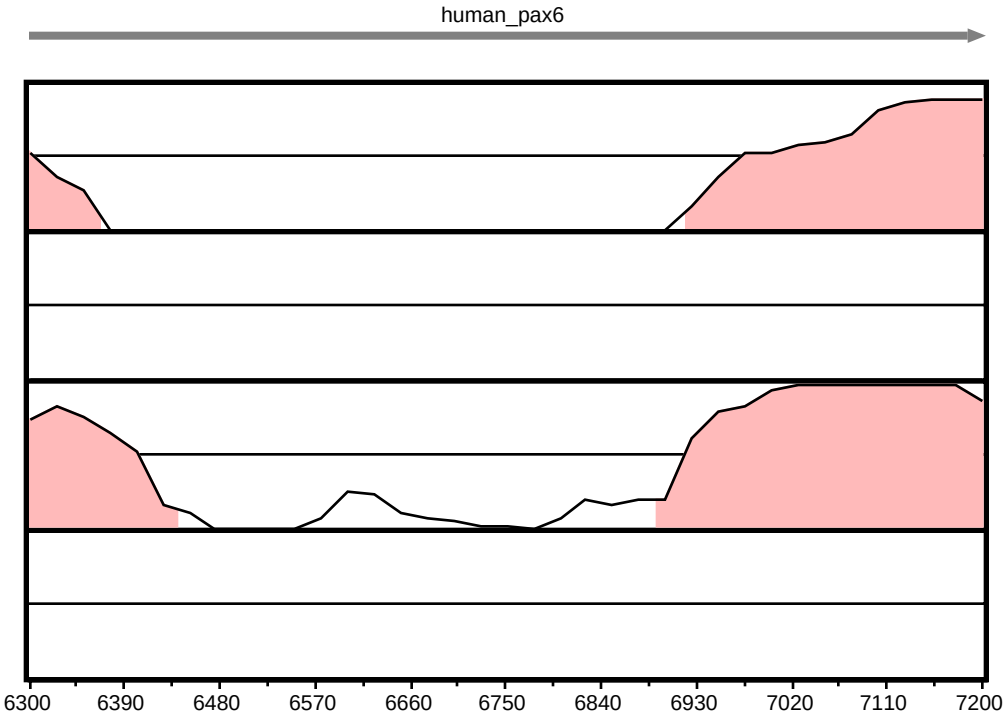

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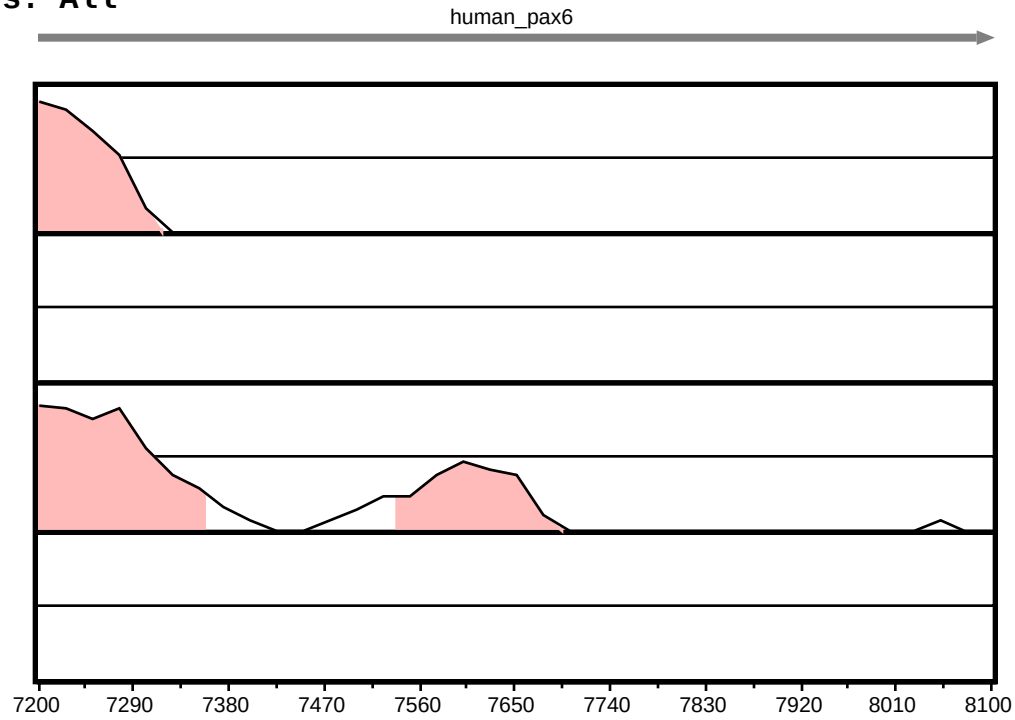

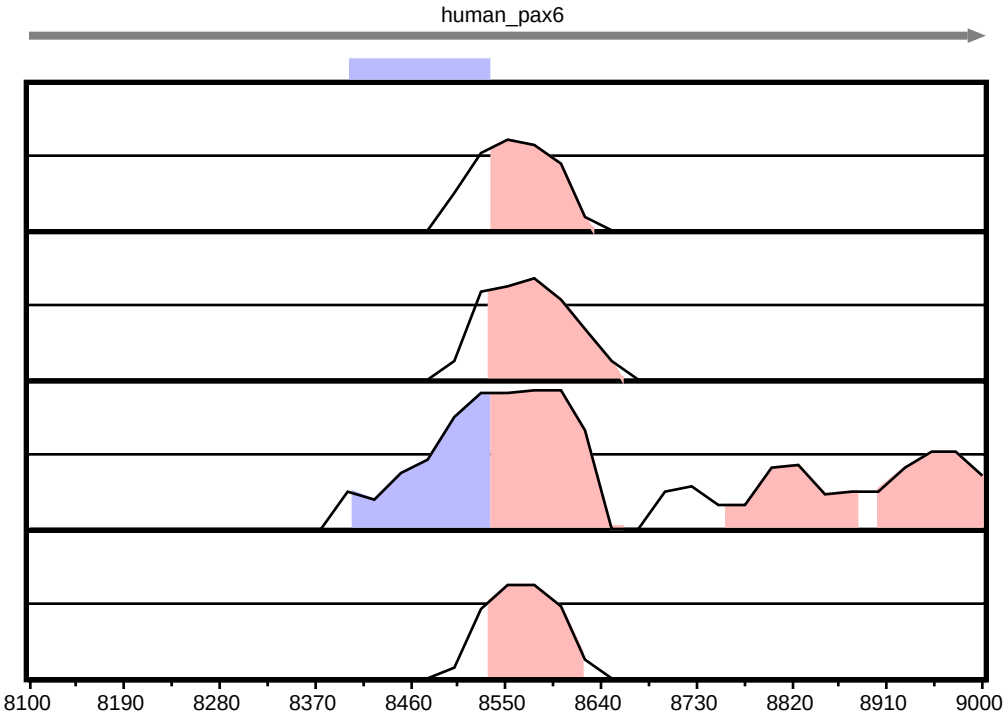

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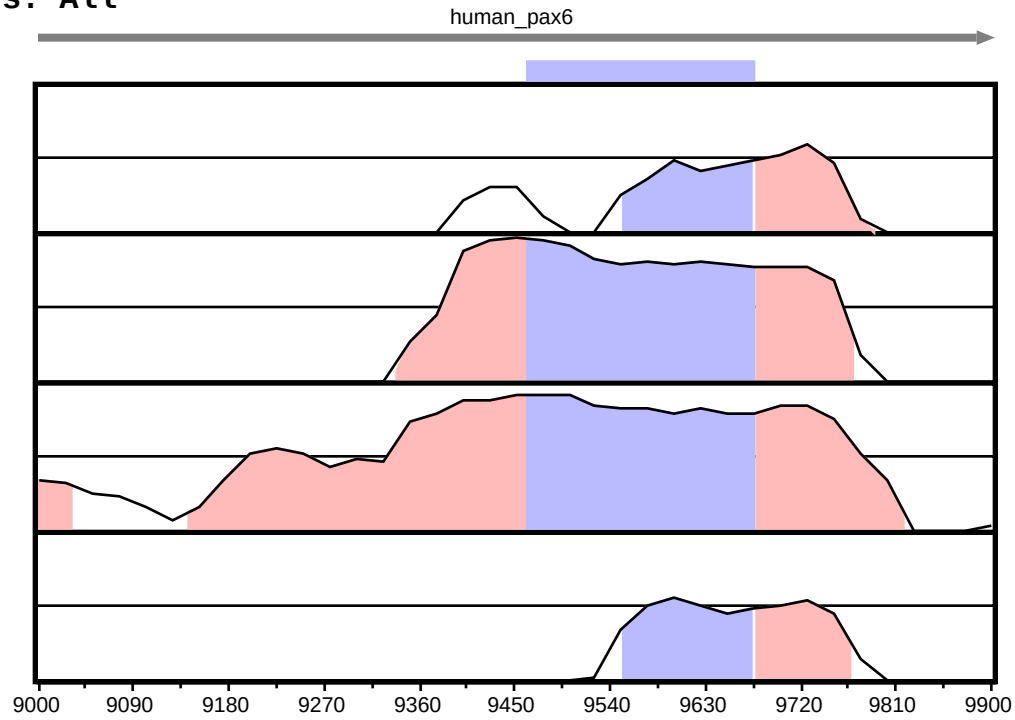

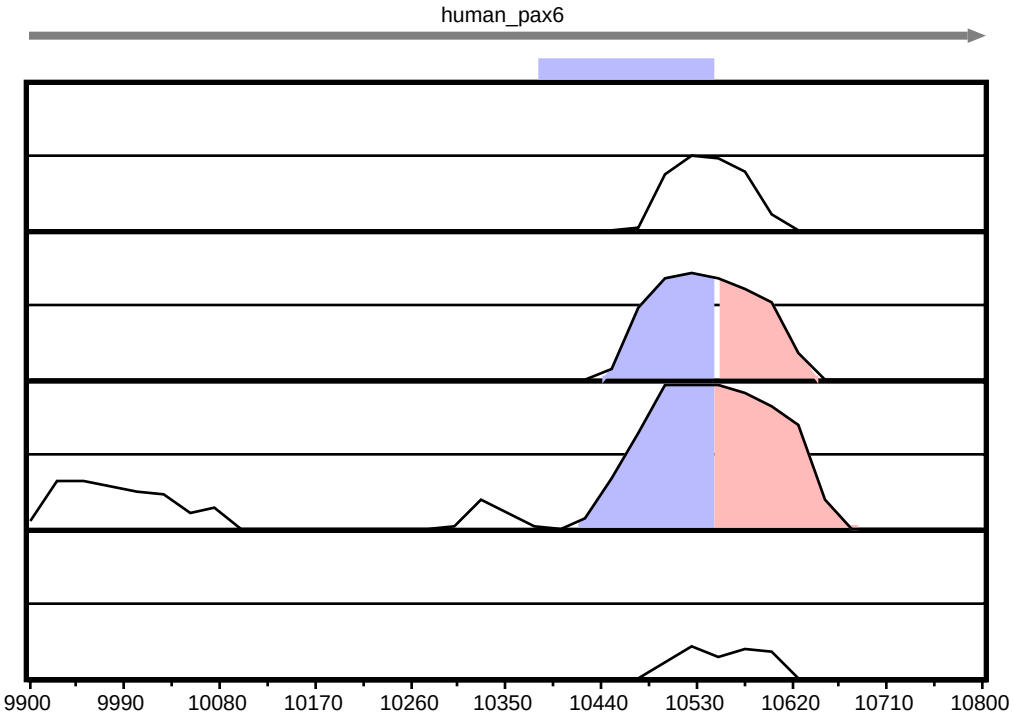

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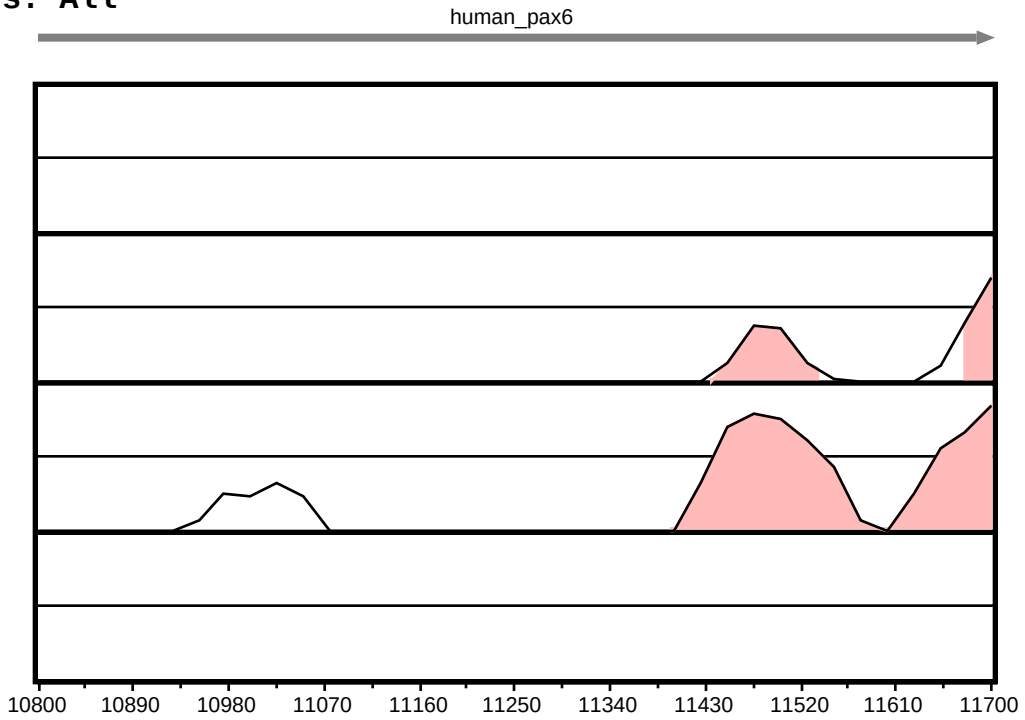

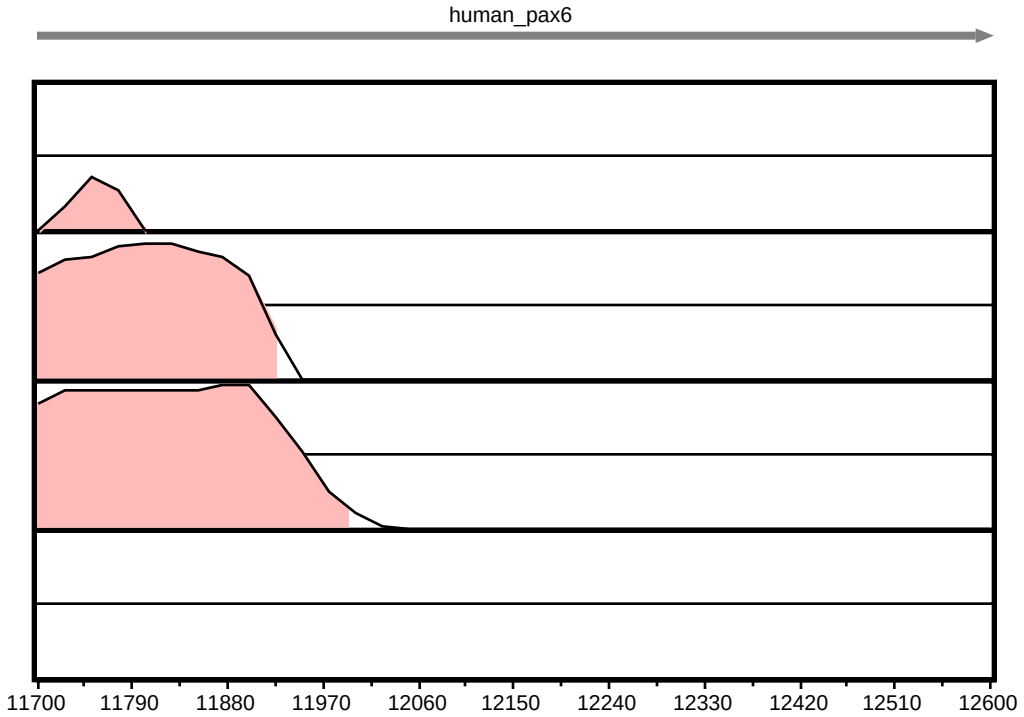

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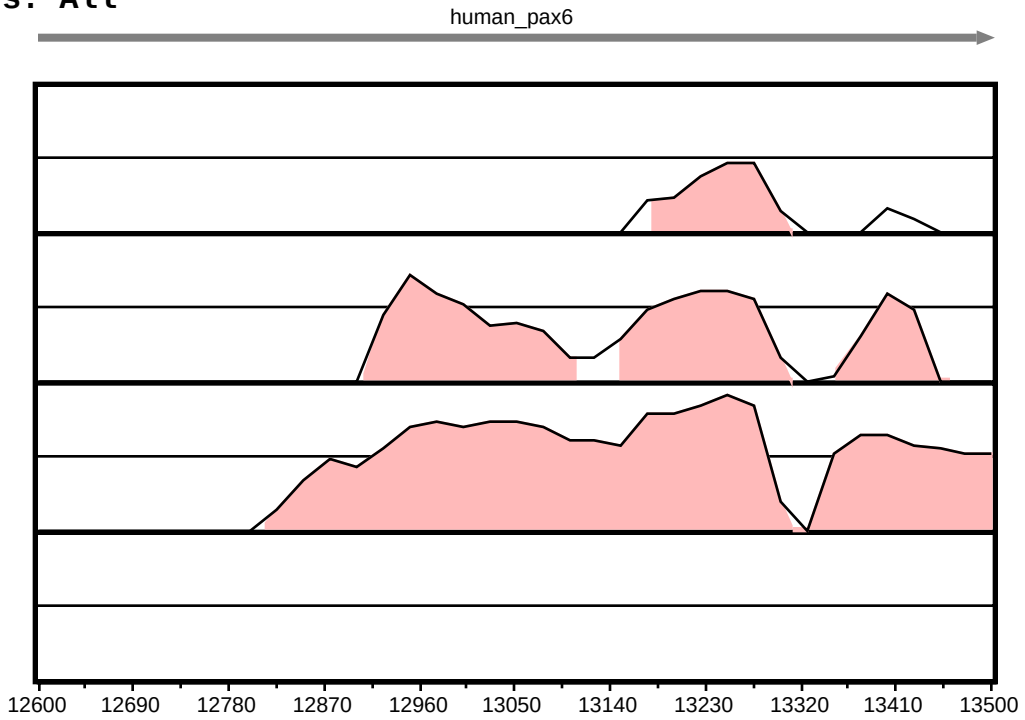

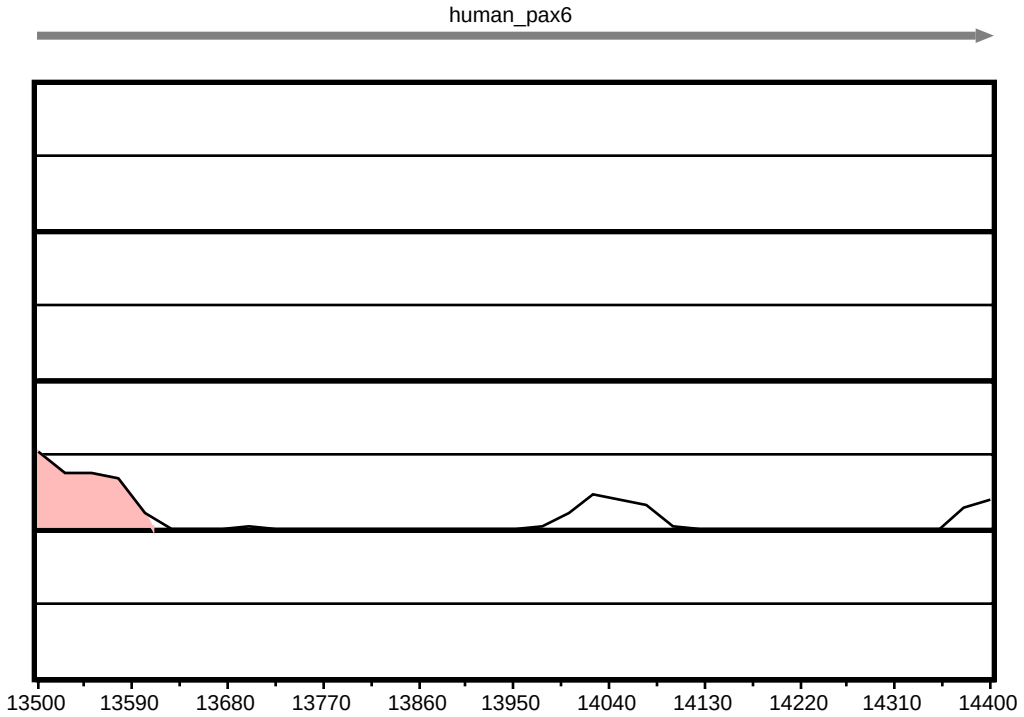

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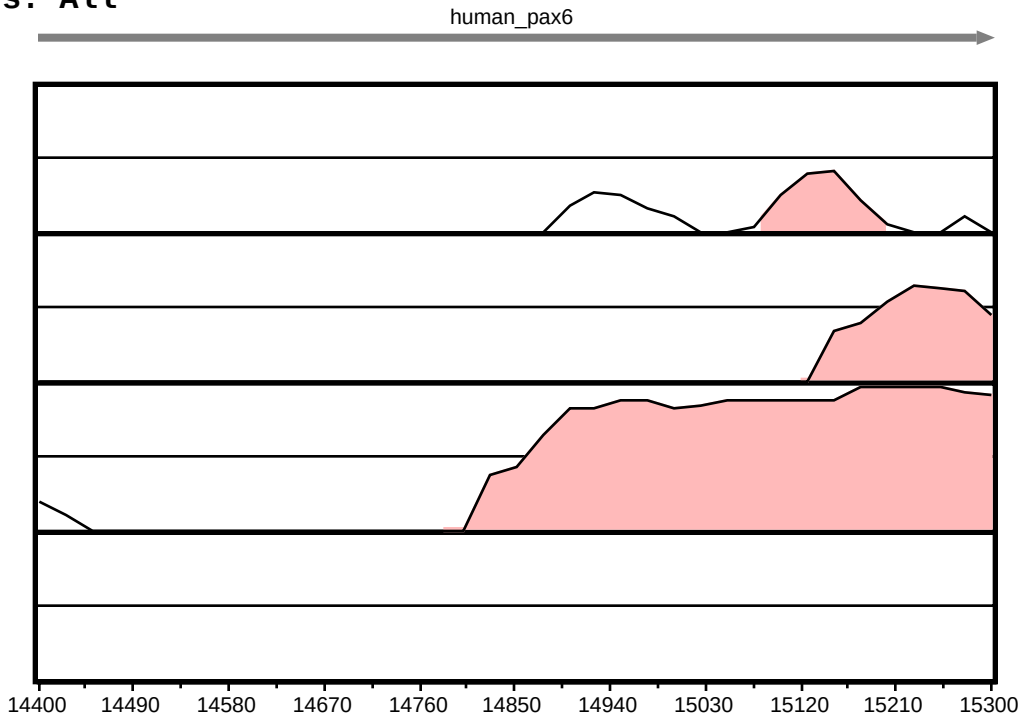

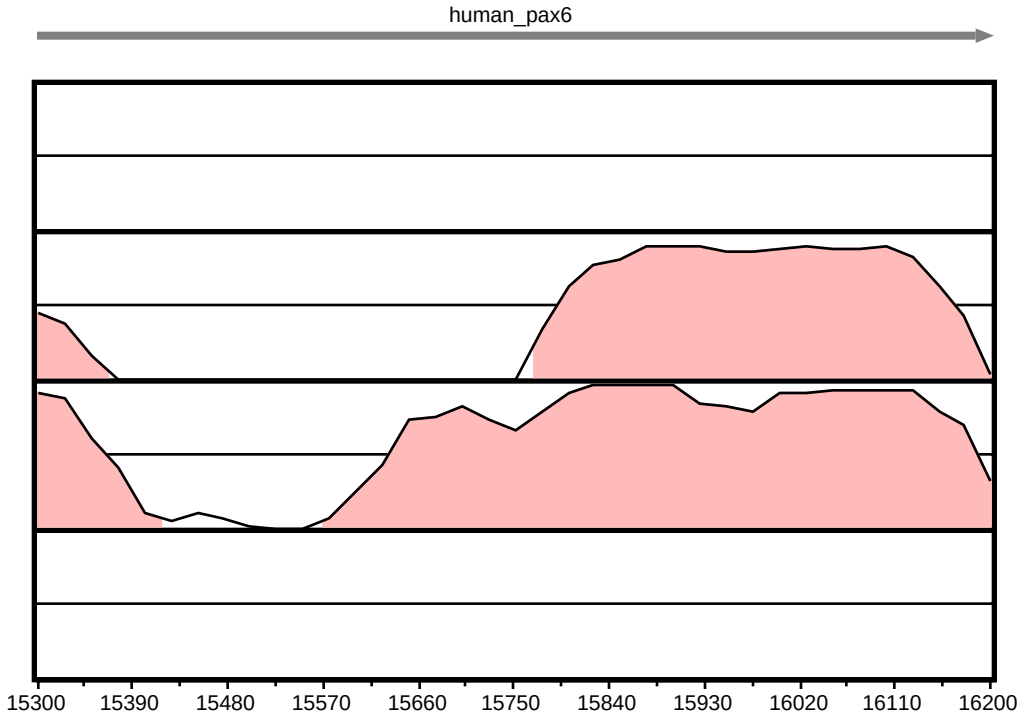

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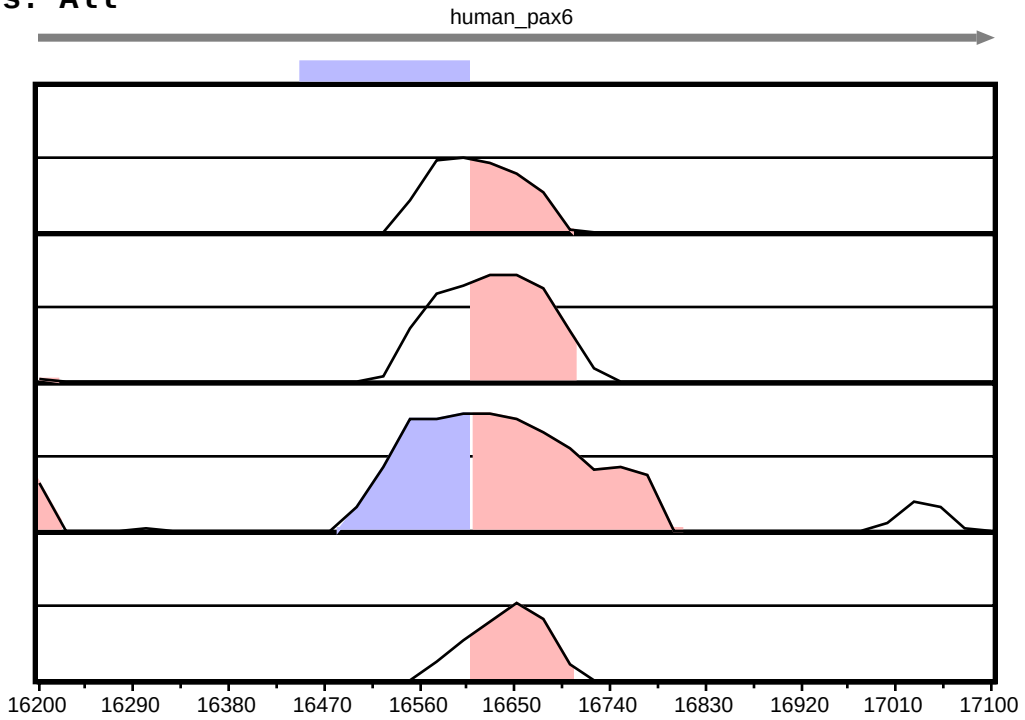

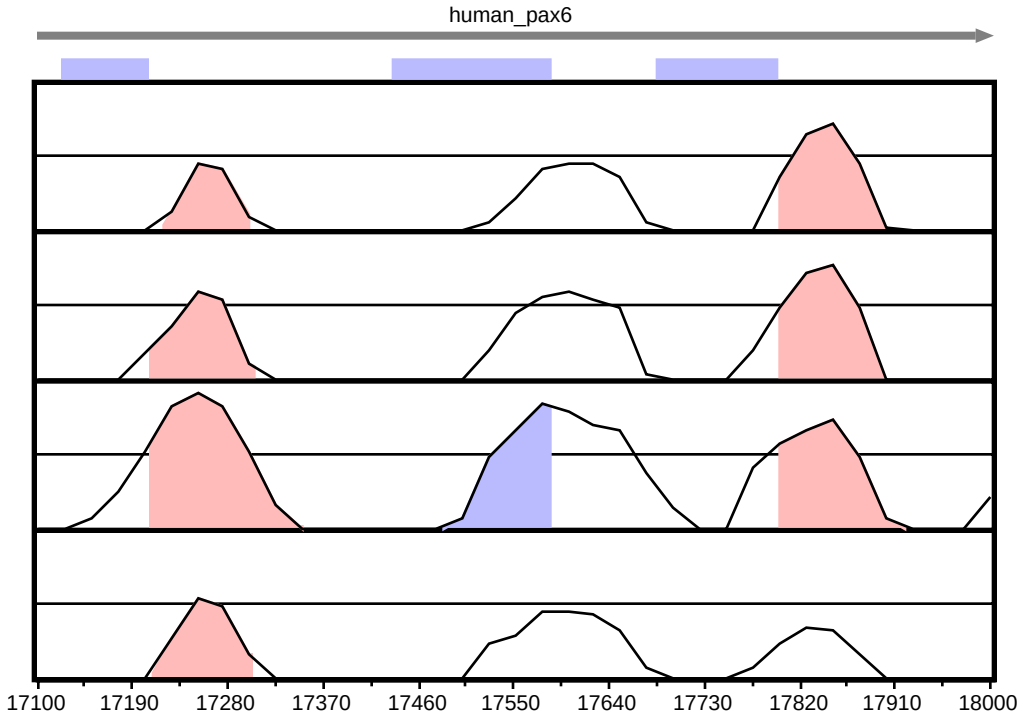

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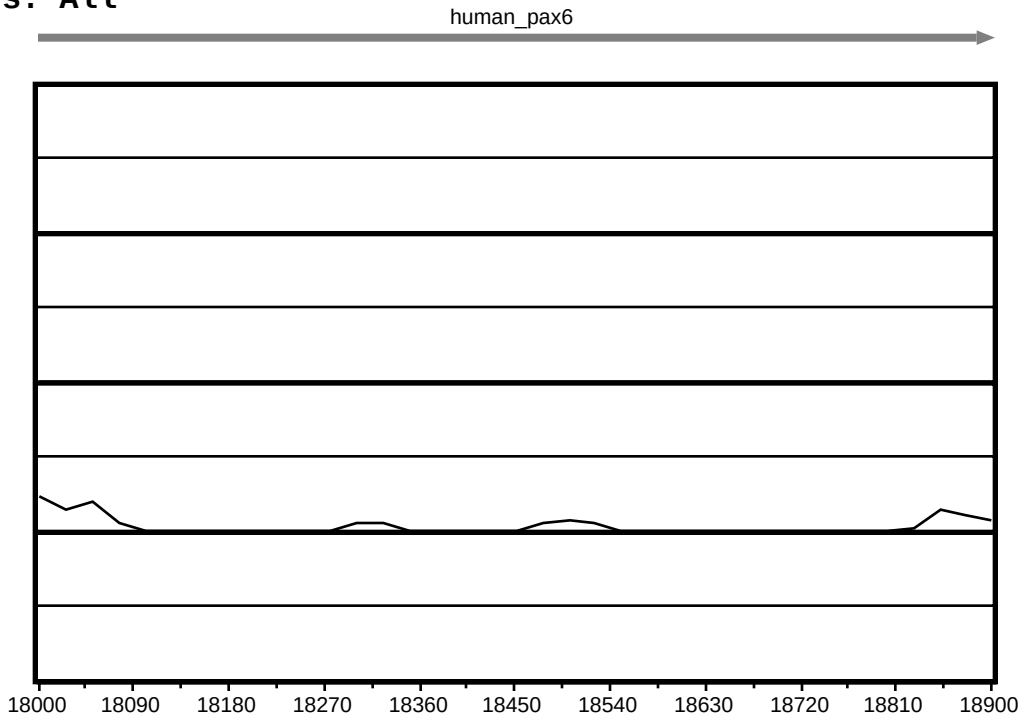

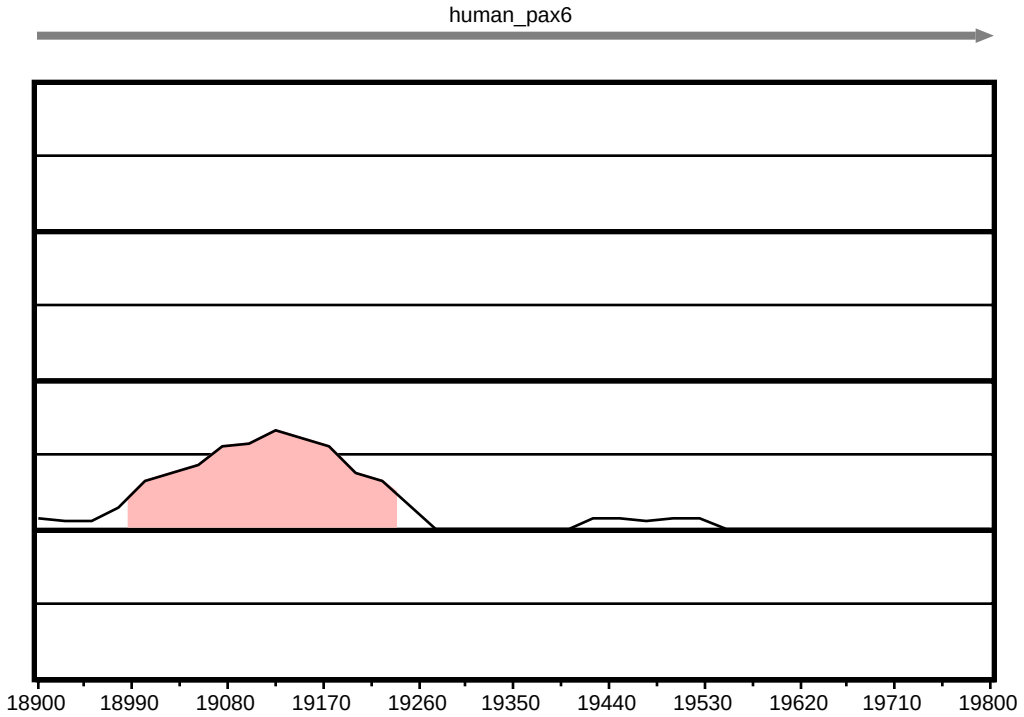

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