MAVID Alignment - 3ENSGALG00000012123proteincodi vs. All

Alignment 1
  Seqs: 3ensgalg00000012123proteincodi/11ensg00000007372proteincoding
  Criteria: 73%, 100 bp
  Regions: 17

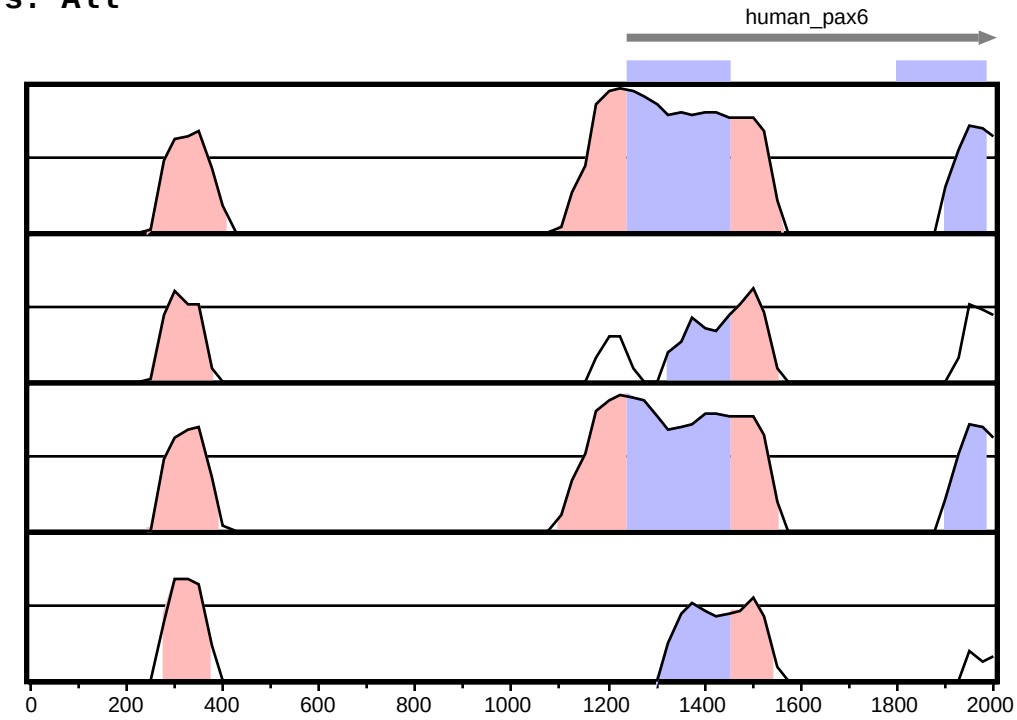
Alignment 2
  Seqs: 3ensgalg00000012123proteincodi/scaffold309newsinfrug000001215
  Criteria: 72%, 100 bp
  Regions: 8

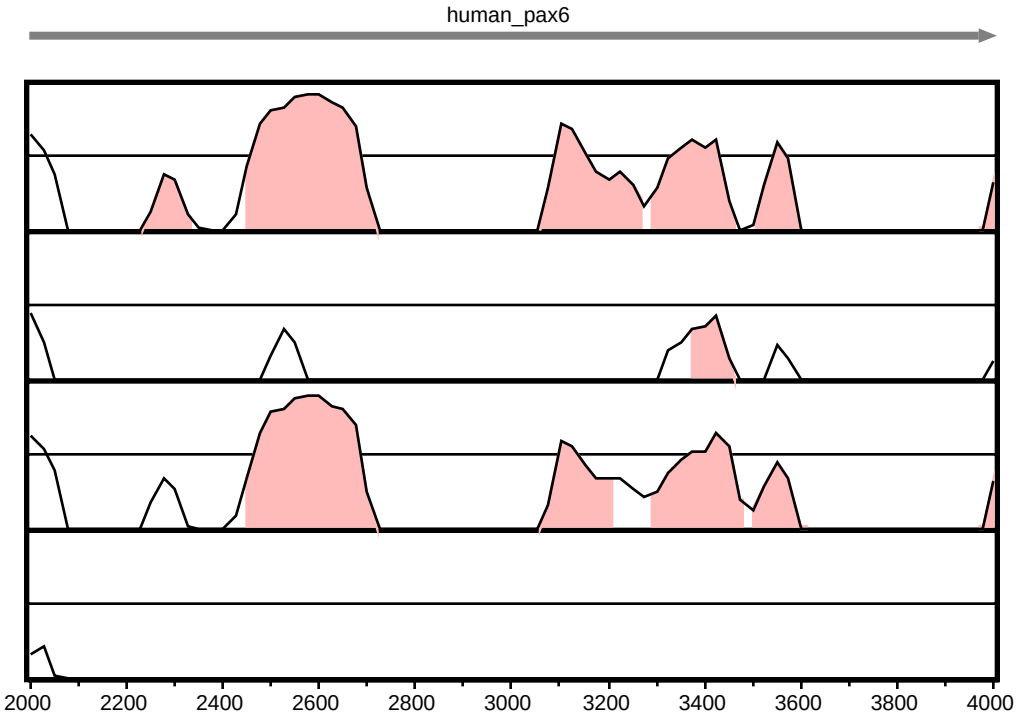
Alignment 3
  Seqs: 3ensgalg00000012123proteincodi/2ensmusg00000027168proteincodi
  Criteria: 74%, 100 bp
  Regions: 16

Alignment 4
  Seqs: 3ensgalg00000012123proteincodi/scaffold430newsinfrug000001506
  Criteria: 68%, 100 bp
  Regions: 6

X-axis: 3ensgalg00000012123proteincodi
Resolution: 25
Window size: 100 bp

gene
exon
UTR
CNS





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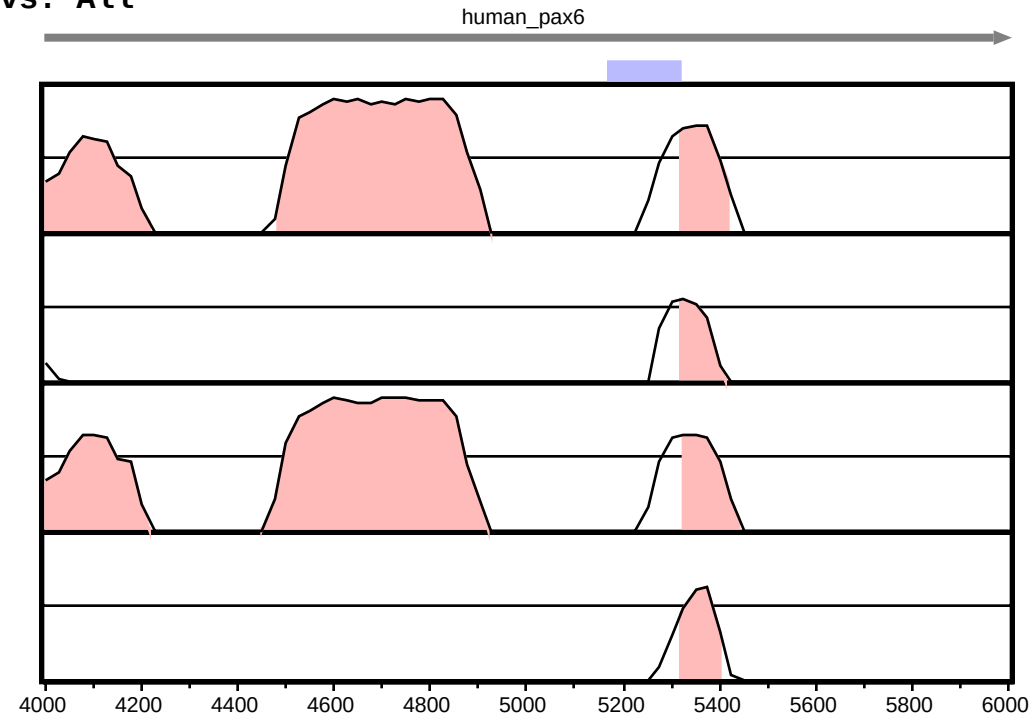
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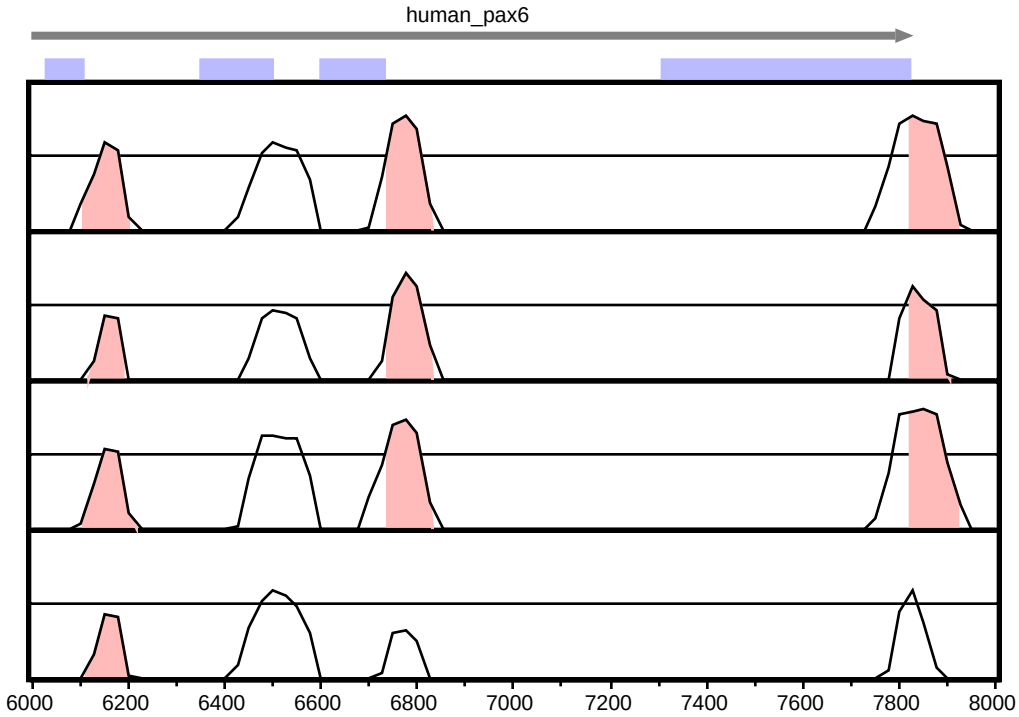
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