

# MAVID Alignment - HOMO vs. All

Alignment 1  
Seqs: homo/pig  
Criteria: 94%, 100 bp  
Regions: 1

Alignment 2  
Seqs: homo/cow  
Criteria: 64%, 100 bp  
Regions: 0

Alignment 3  
Seqs: homo/mouse  
Criteria: 90%, 100 bp  
Regions: 1

Alignment 4  
Seqs: homo/frog  
Criteria: 78%, 100 bp  
Regions: 1

Alignment 5  
Seqs: homo/rat  
Criteria: 89%, 100 bp  
Regions: 1

Alignment 6  
Seqs: homo/trout  
Criteria: 80%, 100 bp  
Regions: 1

Alignment 7  
Seqs: homo/fly  
Criteria: 69%, 100 bp  
Regions: 1

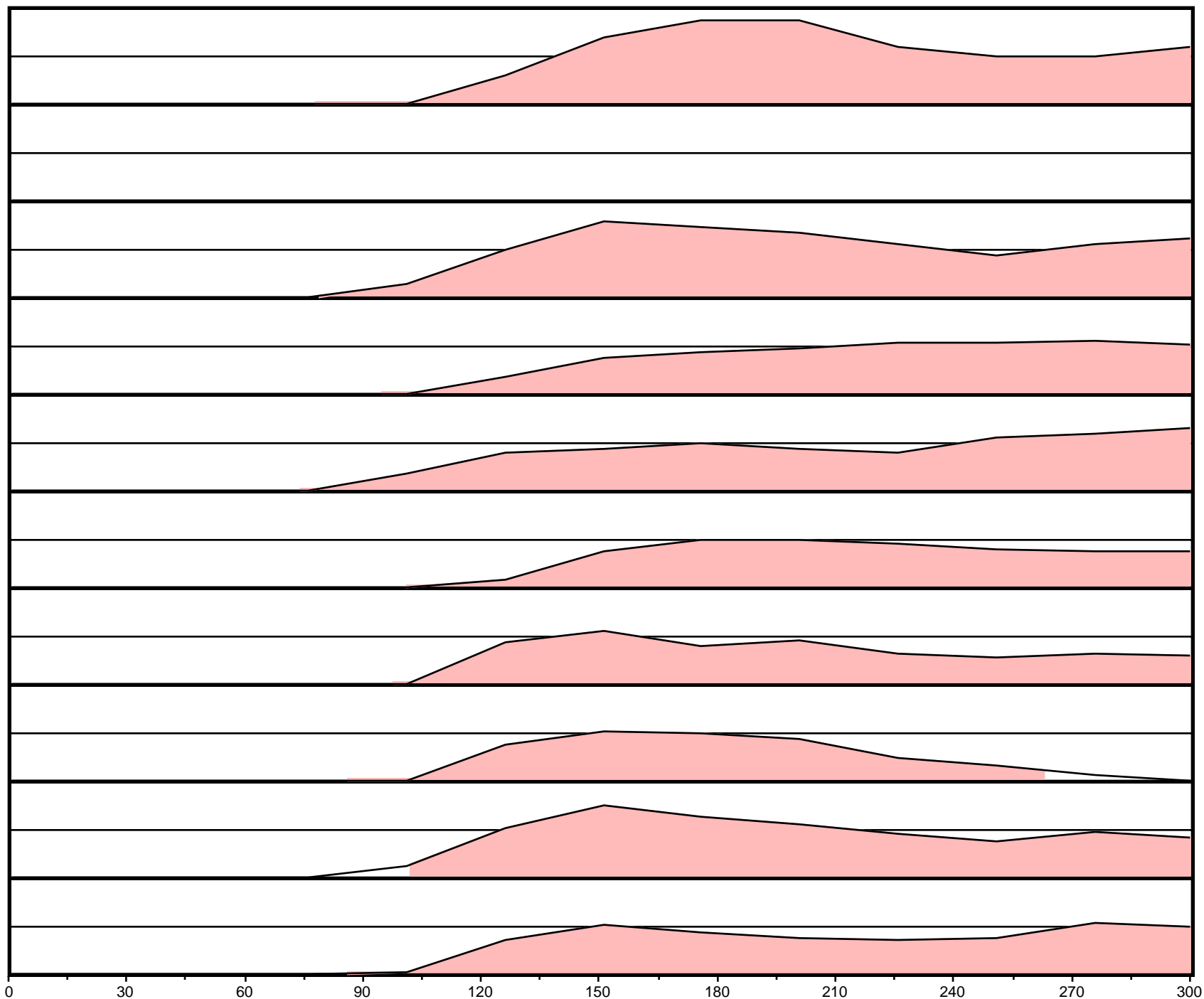
Alignment 8  
Seqs: homo/celegans  
Criteria: 67%, 100 bp  
Regions: 2

Alignment 9  
Seqs: homo/flower  
Criteria: 69%, 100 bp  
Regions: 1

Alignment 10  
Seqs: homo/soy  
Criteria: 68%, 100 bp  
Regions: 1

X-axis: homo  
Resolution: 25  
Window size: 100 bp

► gene  
■ exon  
■ UTR  
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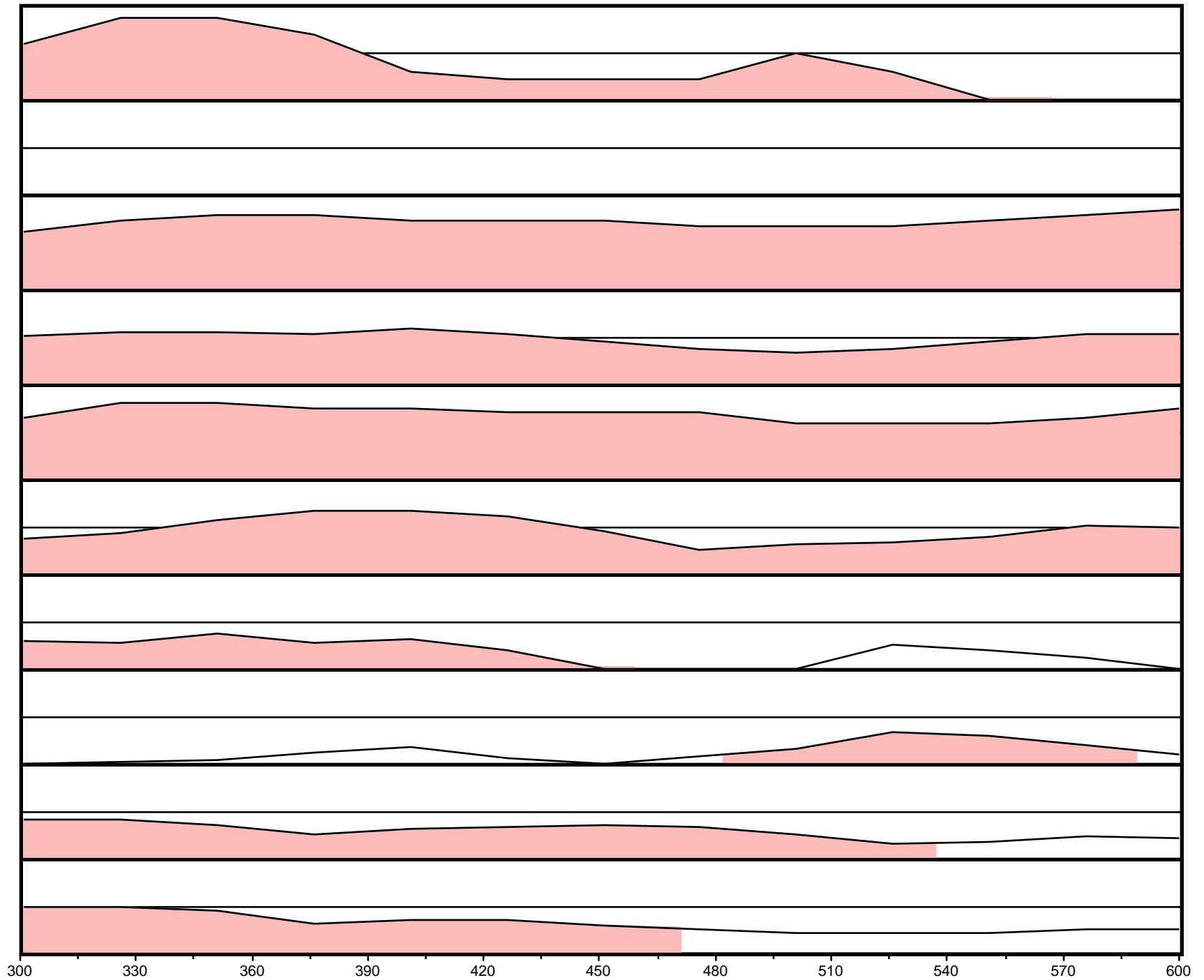
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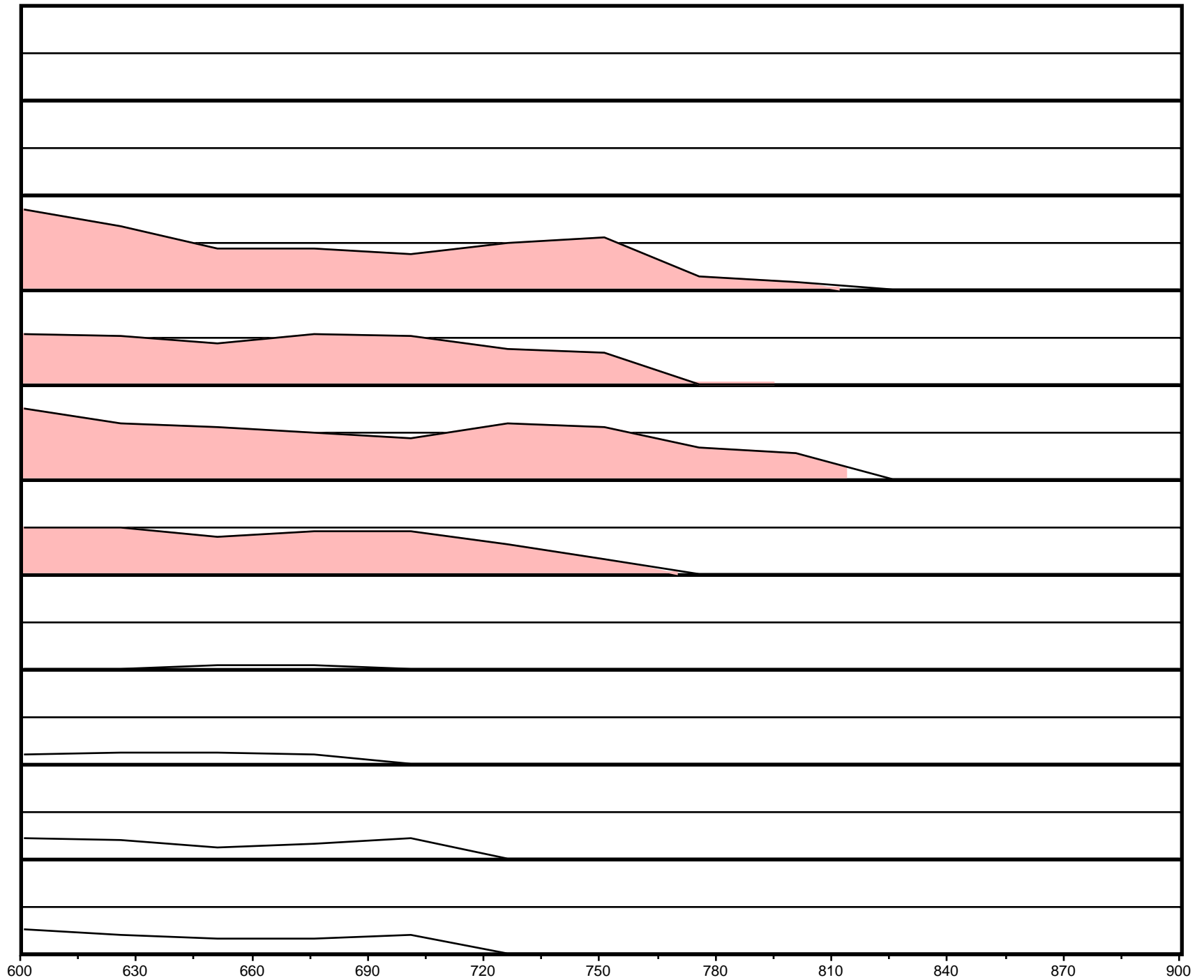
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