

MAVID Alignment - HOMO vs. All

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

Alignment 2
Seqs: homo/cow
Criteria: 63%, 100 bp
Regions: 0

Alignment 3
Seqs: homo/mouse
Criteria: 87%, 100 bp
Regions: 2

Alignment 4
Seqs: homo/frog
Criteria: 80%, 100 bp
Regions: 1

Alignment 5
Seqs: homo/rat
Criteria: 86%, 100 bp
Regions: 1

Alignment 6
Seqs: homo/trout
Criteria: 82%, 100 bp
Regions: 2

Alignment 7
Seqs: homo/danio
Criteria: 83%, 100 bp
Regions: 2

Alignment 8
Seqs: homo/fly
Criteria: 74%, 100 bp
Regions: 1

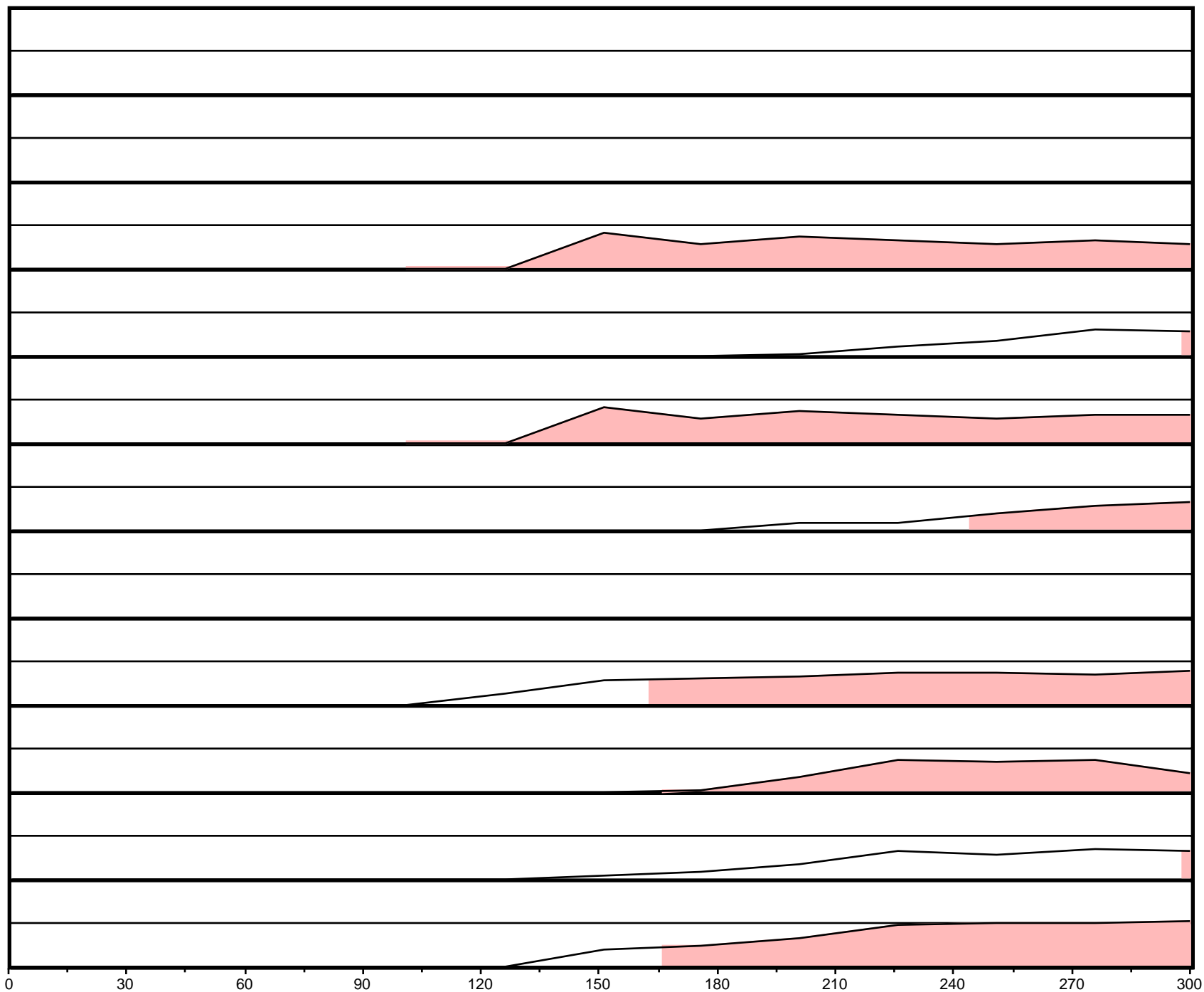
Alignment 9
Seqs: homo/celegans
Criteria: 72%, 100 bp
Regions: 2

Alignment 10
Seqs: homo/flower
Criteria: 71%, 100 bp
Regions: 2

Alignment 11
Seqs: homo/soy
Criteria: 71%, 100 bp
Regions: 2

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

▶ gene
■ exon
■ UTR
■ CNS



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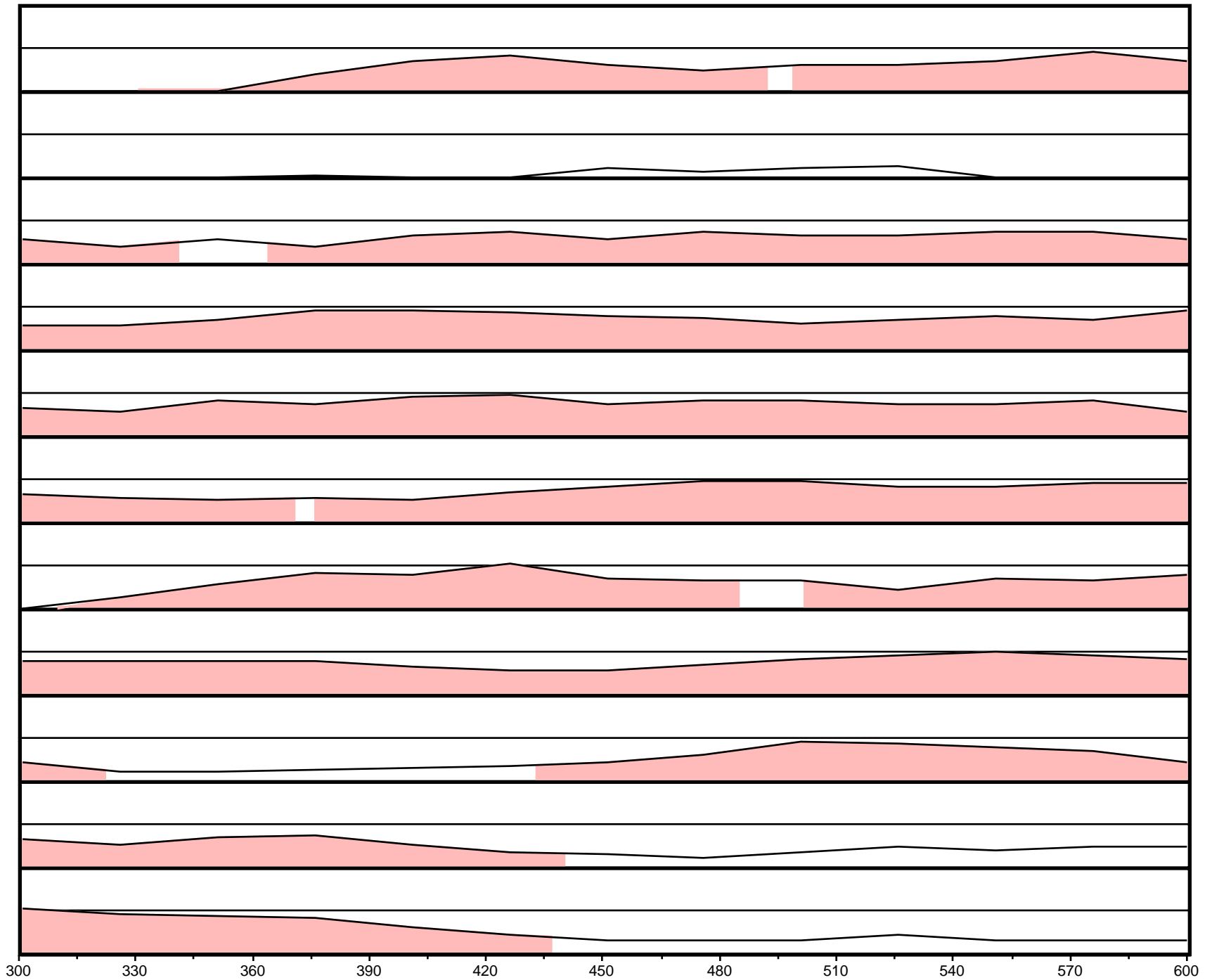
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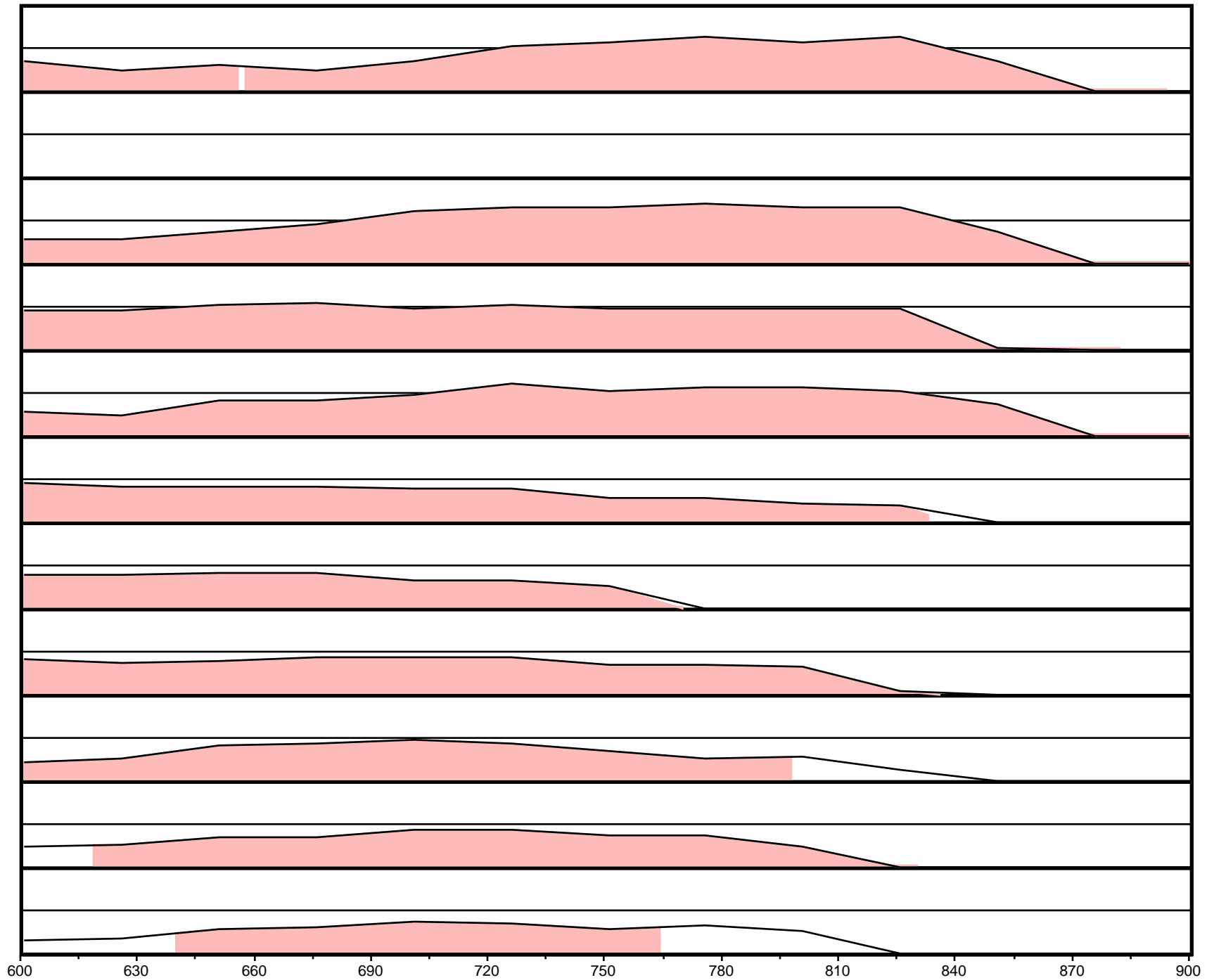
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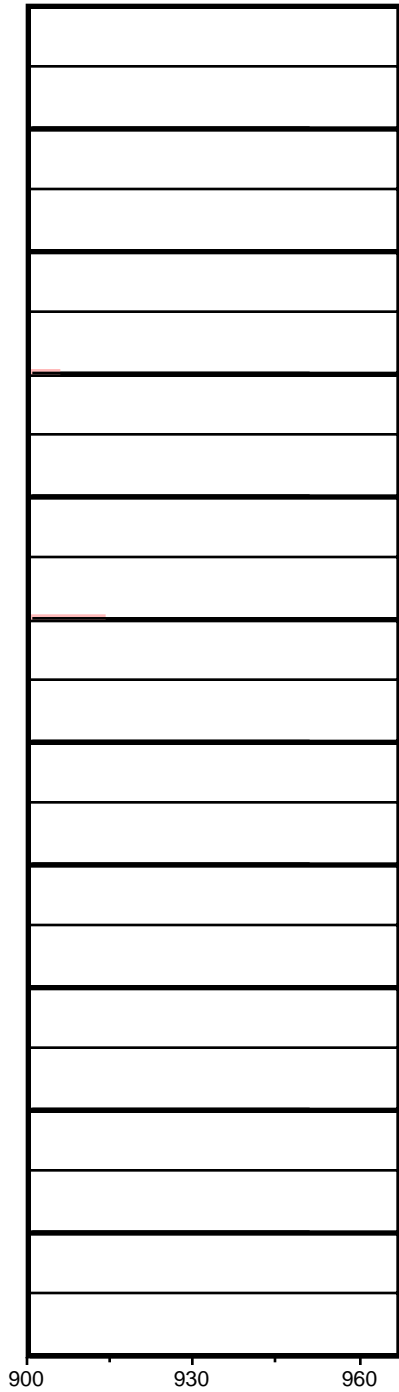
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900 930 960