

MAVID Alignment - HOMO vs. All

Alignment 1
Seqs: homo/pig
Criteria: 68%, 100 bp
Regions: 0

Alignment 2
Seqs: homo/cow
Criteria: 92%, 100 bp
Regions: 3

Alignment 3
Seqs: homo/mouse
Criteria: 88%, 100 bp
Regions: 3

Alignment 4
Seqs: homo/frog
Criteria: 80%, 100 bp
Regions: 2

Alignment 5
Seqs: homo/rat
Criteria: 88%, 100 bp
Regions: 1

Alignment 6
Seqs: homo/trout
Criteria: 85%, 100 bp
Regions: 3

Alignment 7
Seqs: homo/danio
Criteria: 81%, 100 bp
Regions: 1

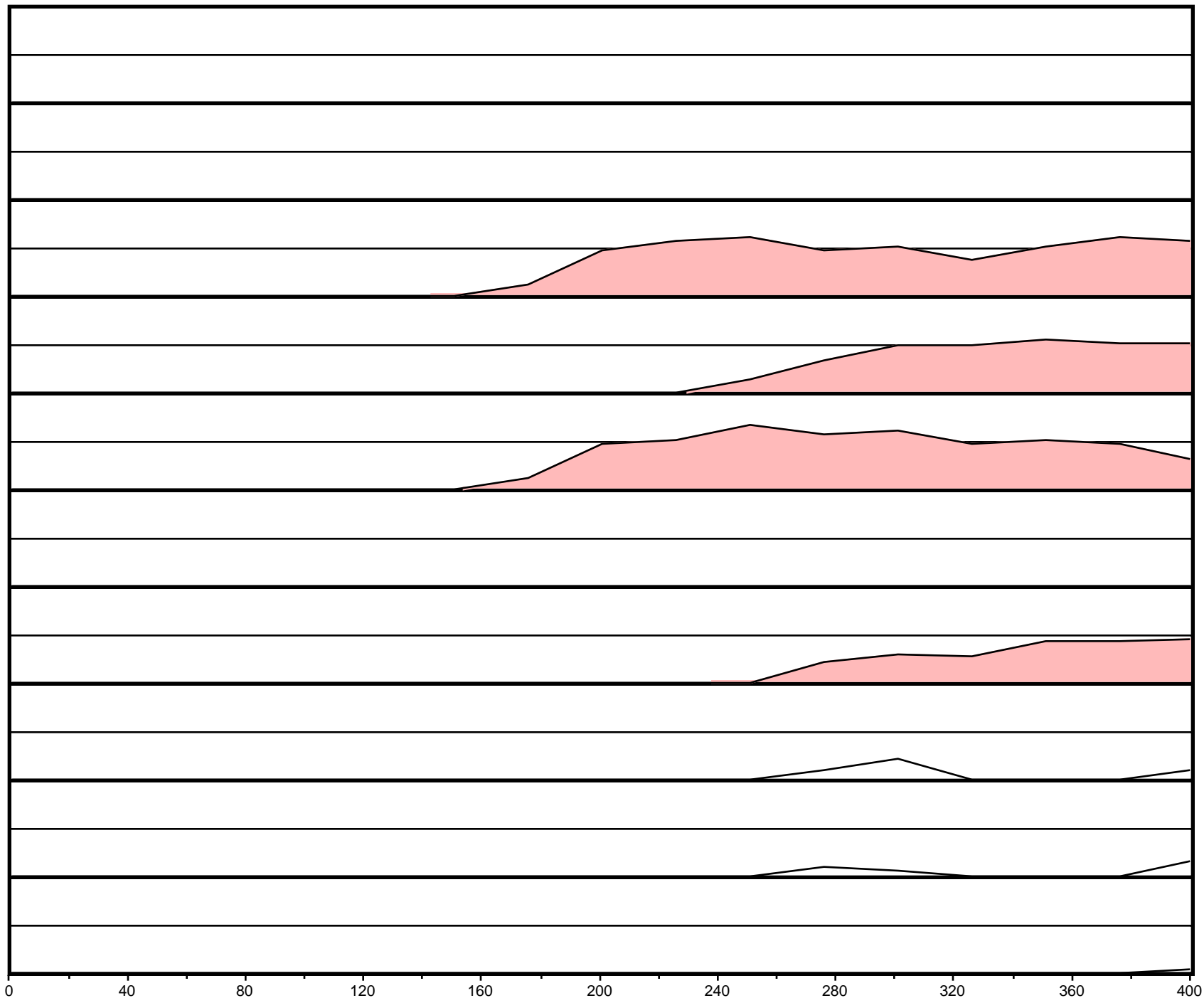
Alignment 8
Seqs: homo/fly
Criteria: 77%, 100 bp
Regions: 2

Alignment 9
Seqs: homo/celegans
Criteria: 75%, 100 bp
Regions: 2

Alignment 10
Seqs: homo/flower
Criteria: 74%, 100 bp
Regions: 3

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

► gene
■ exon
■ UTR
■ CNS



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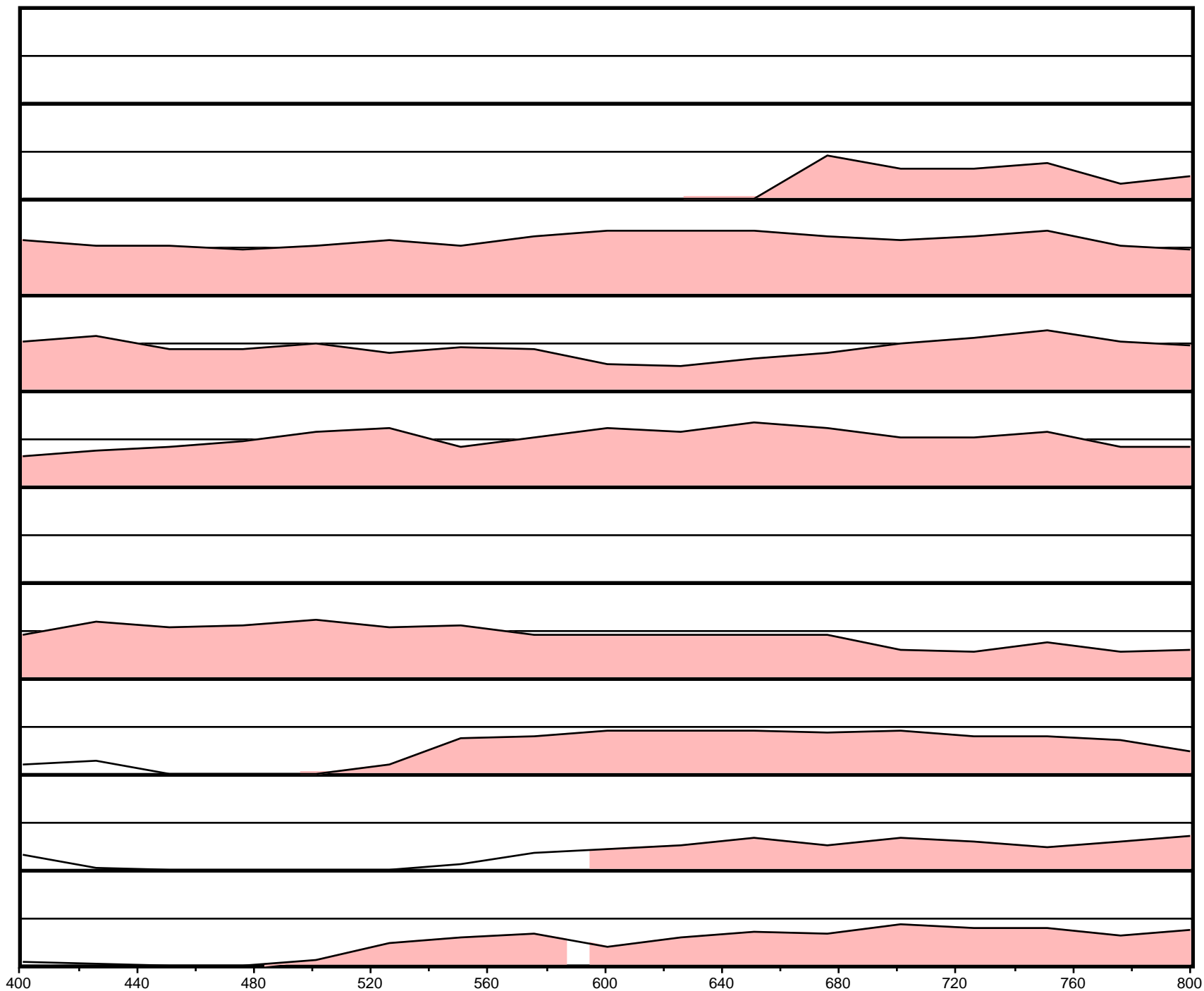
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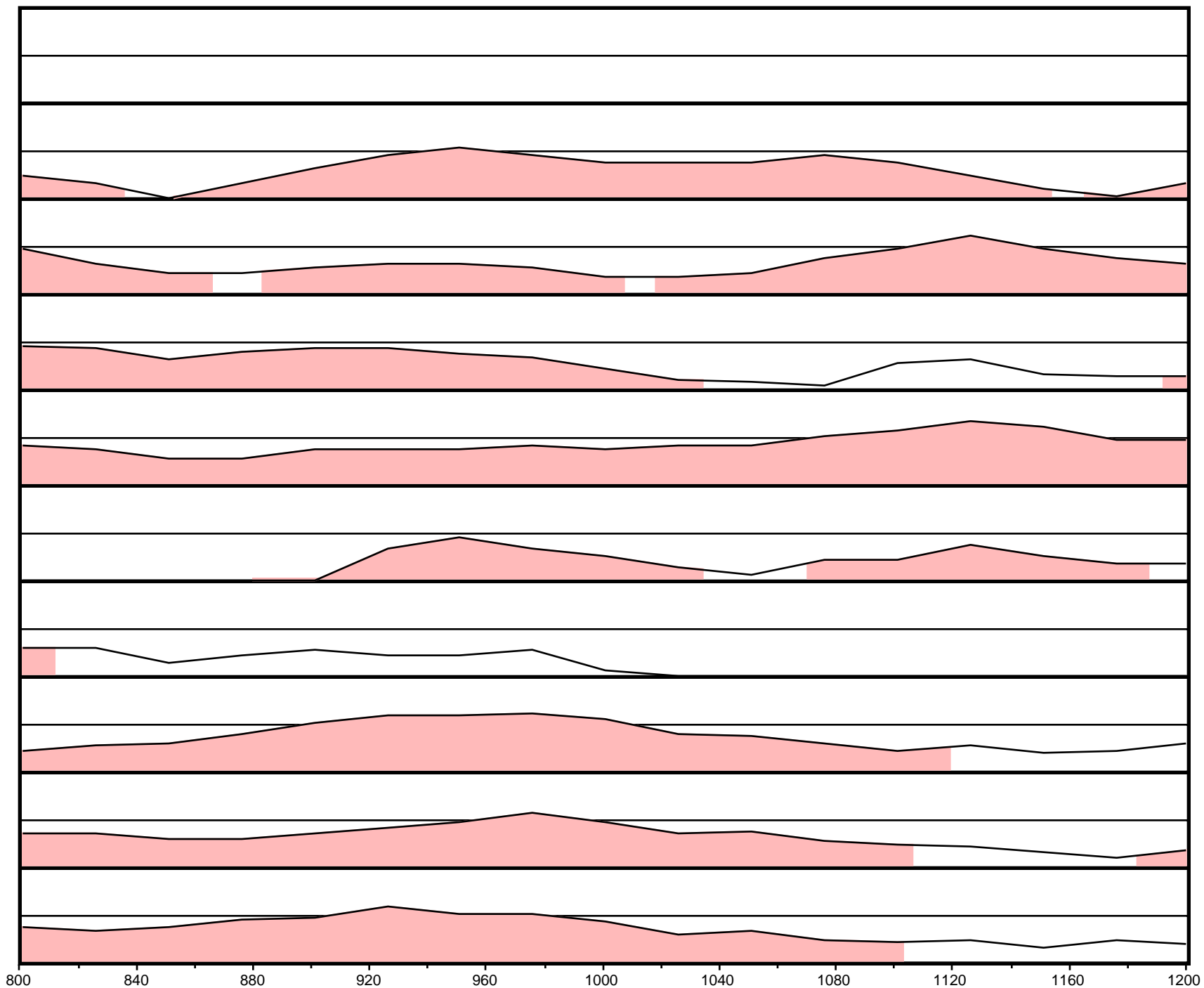
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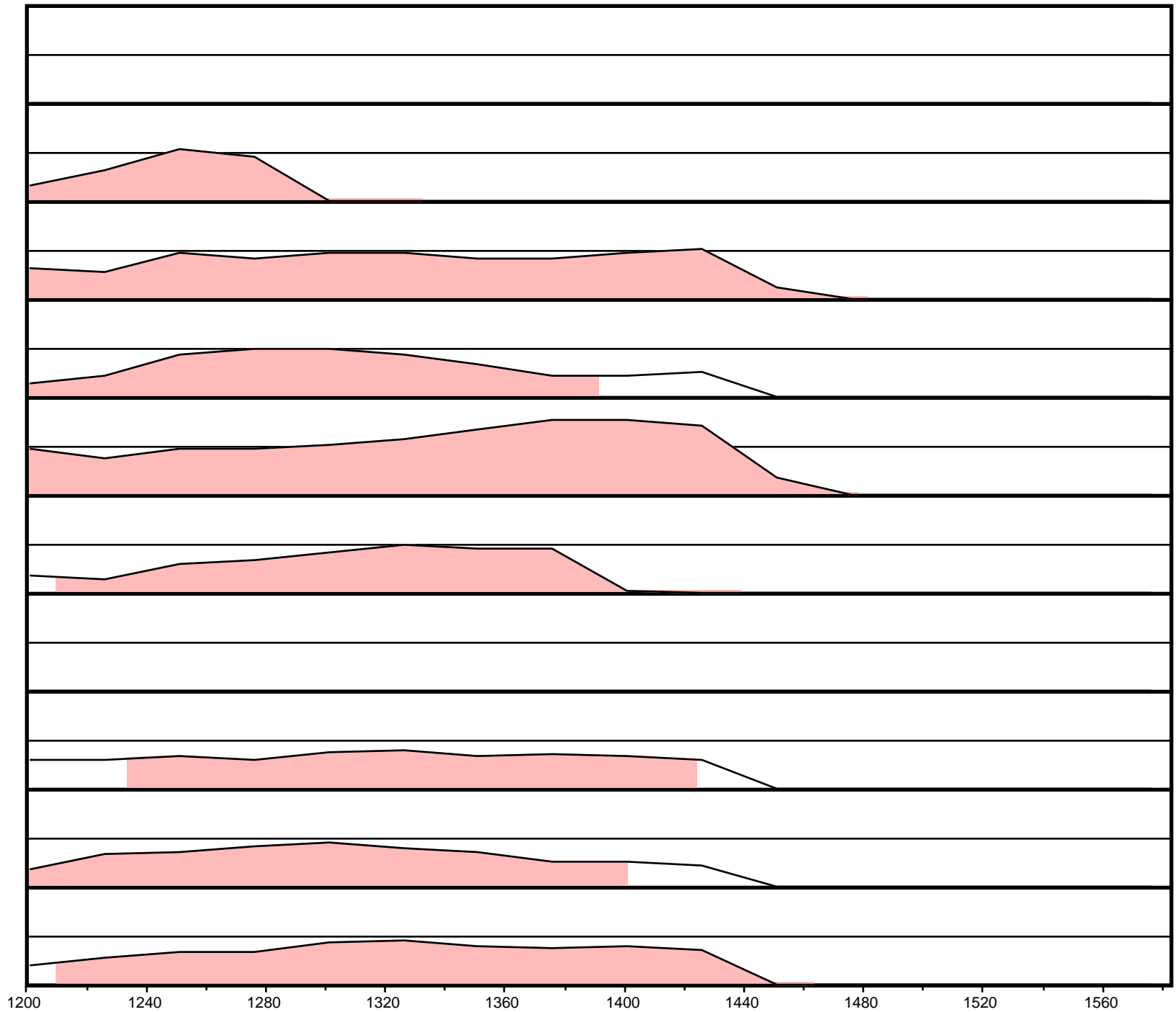
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1200 1240 1280 1320 1360 1400 1440 1480 1520 1560