

GeneMark

Version 2.5p (09.08.06)

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PROGRAM INFORMATION

Sequence : Mycobacterium phage Citius complete sequence, 51393 bp including 10 bp 3' overhang (CGGCCGG
Analysis Date : 5/23/17 at 12:53:07
Pages : 26
Sequence Length : 51393 bp
GC Content : 63.89%

Window Length : 96 bp
Window Step : 12 bp
Threshold Value : 0.500

PS-Version : 1.2

GeneMark Options : PostScript graph,
Mark ORFs / splice sites,
List ORFs,
List regions and/or splice sites,

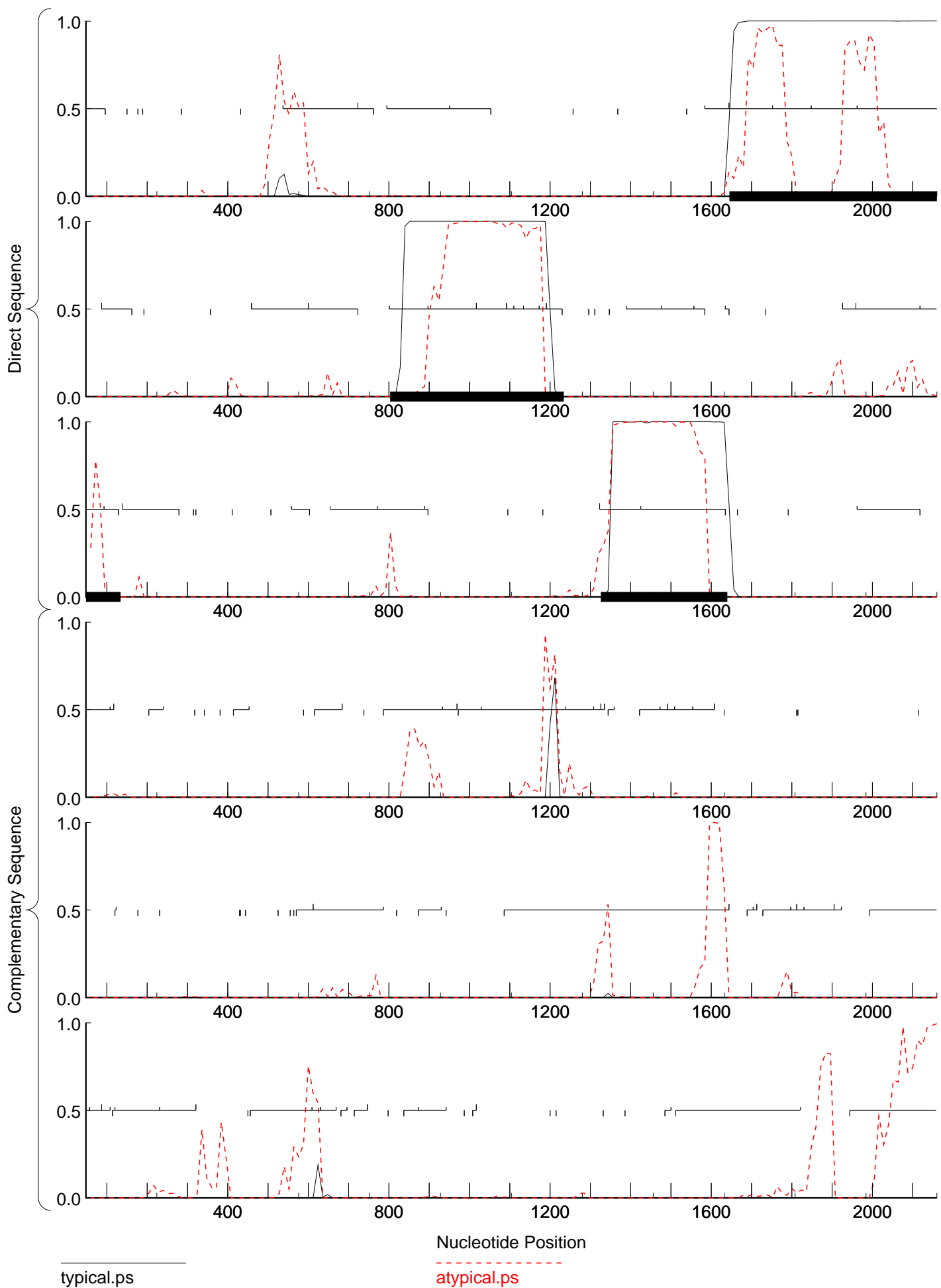
Matrix notes & comments

Training set derived by GeneMarkS, 4.30 October 2014
Tue May 23 12:53:07 2017

MATRIX INFORMATION

Matrix : -
Author : -
Order : 4

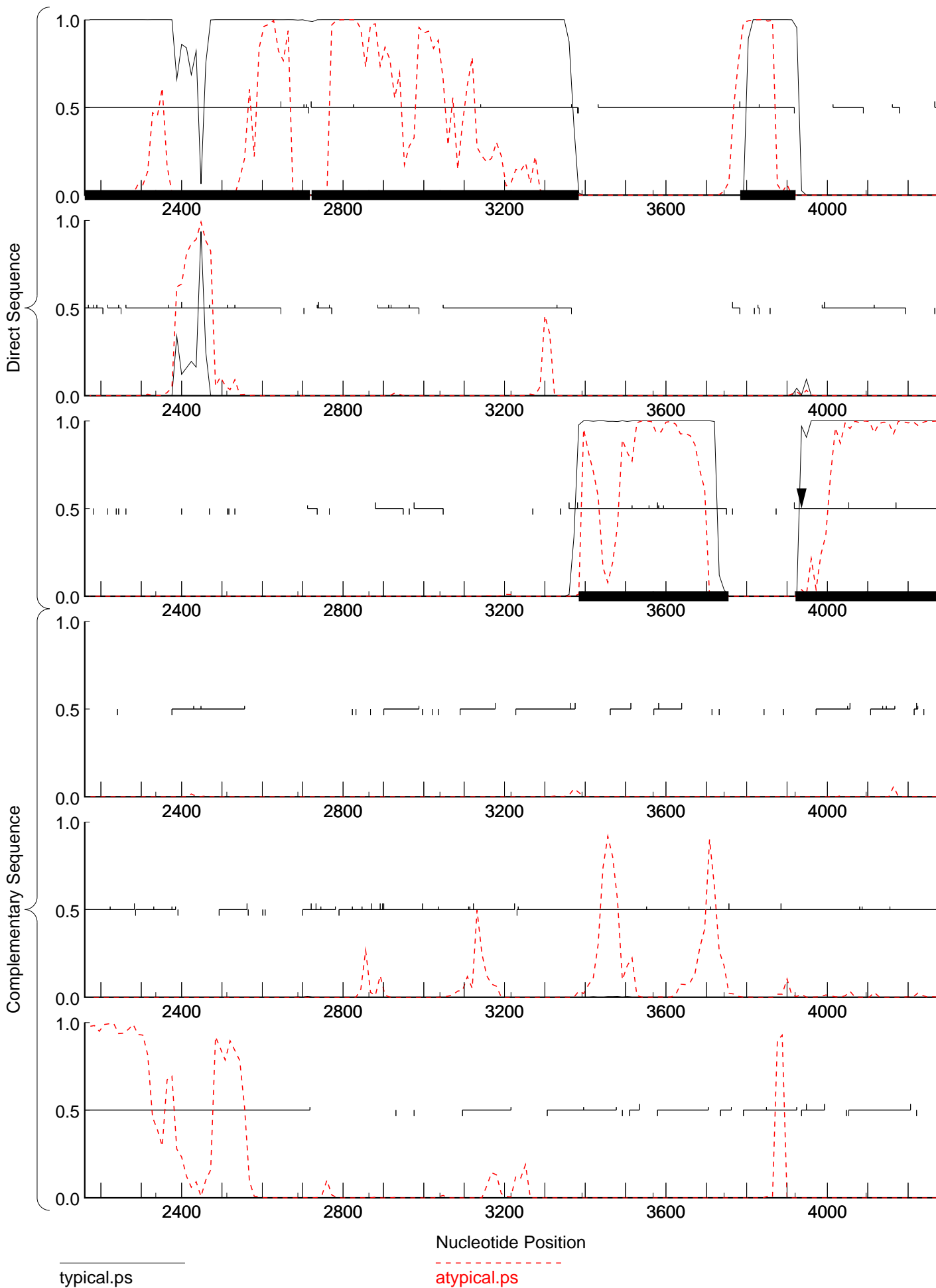
Send questions / comments to:
Dr. M. Borodovsky
Georgia Institute of Technology
School of Biology
Atlanta, GA 30332-0230

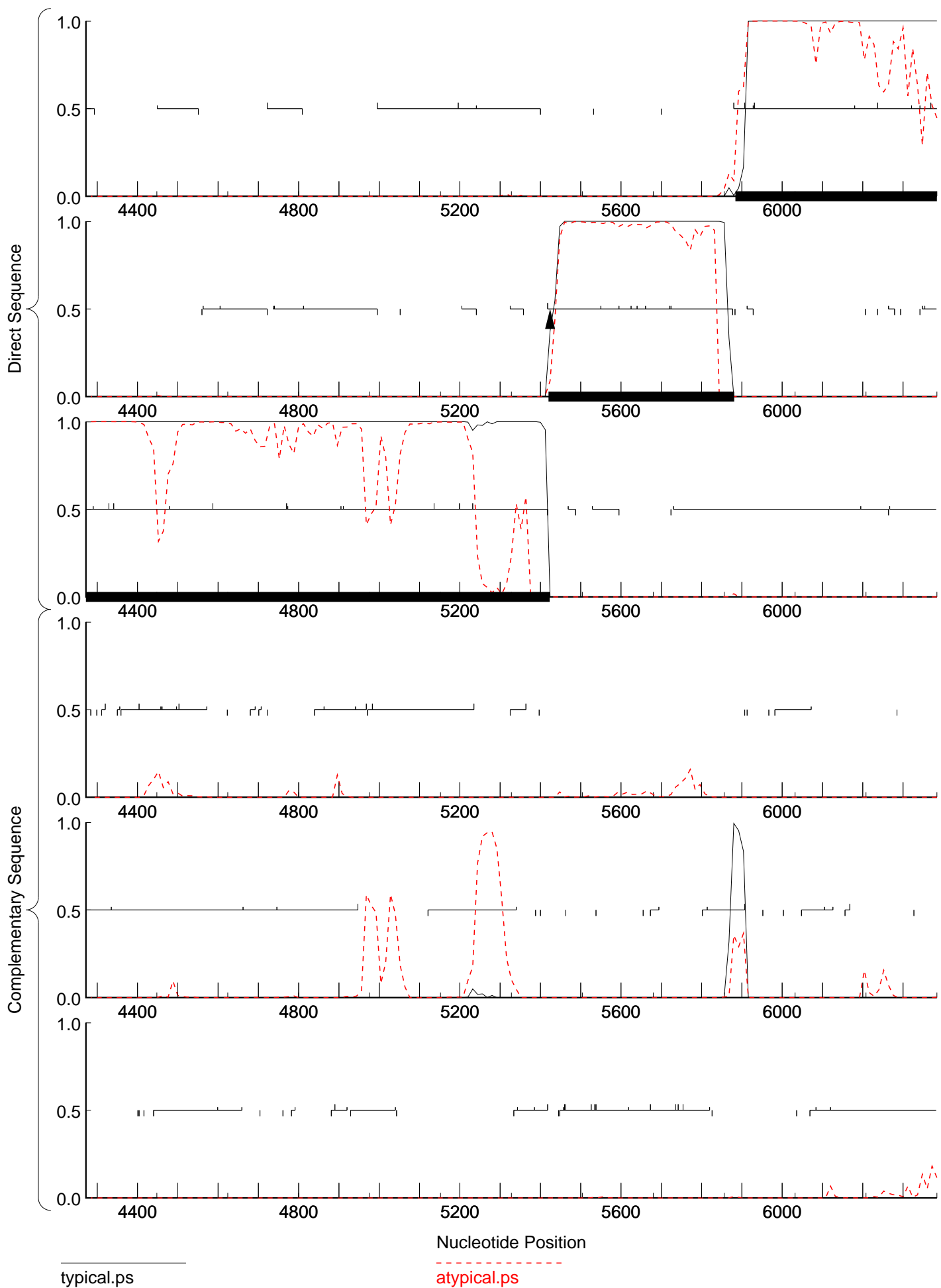


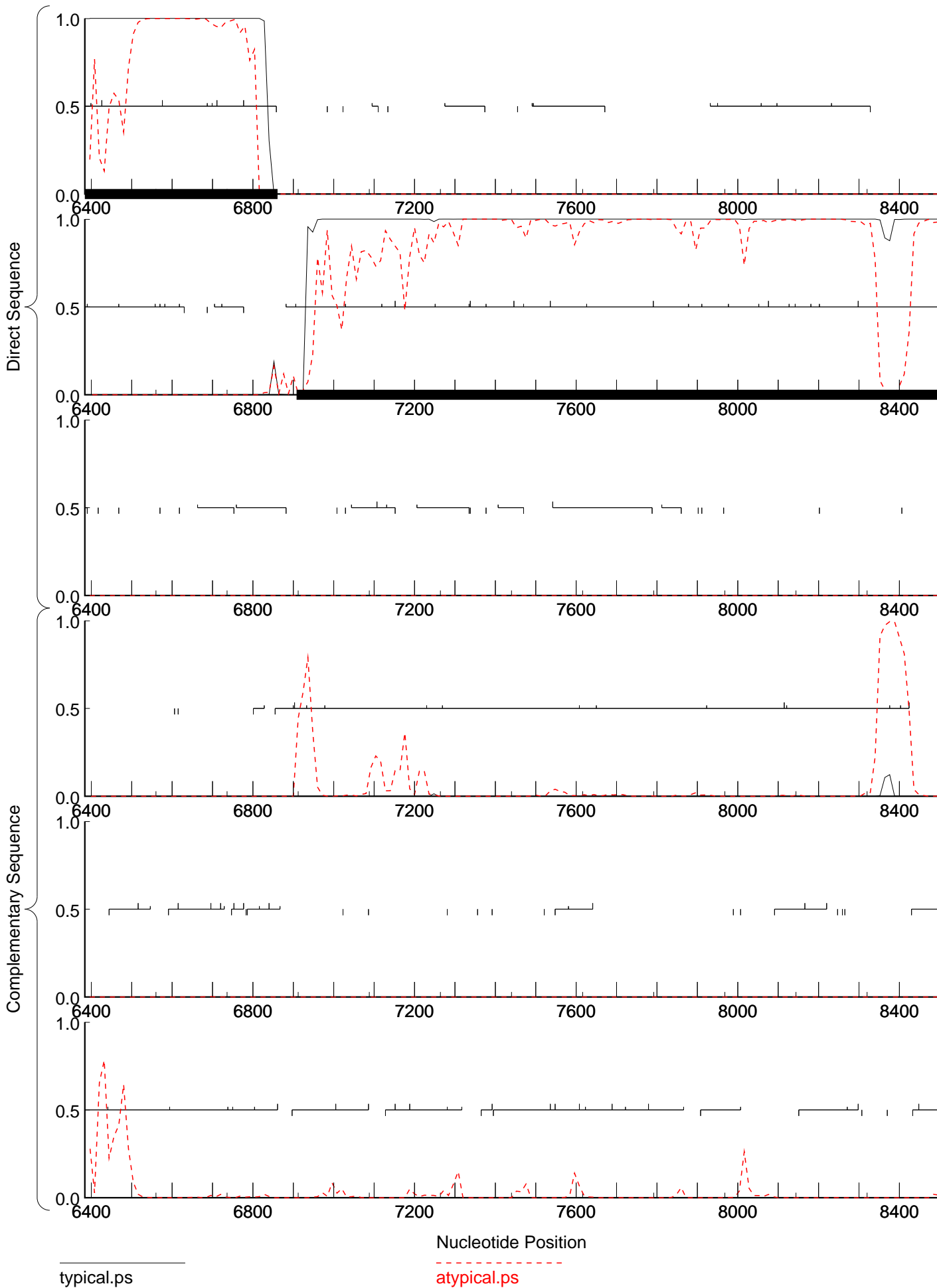
typical.ps

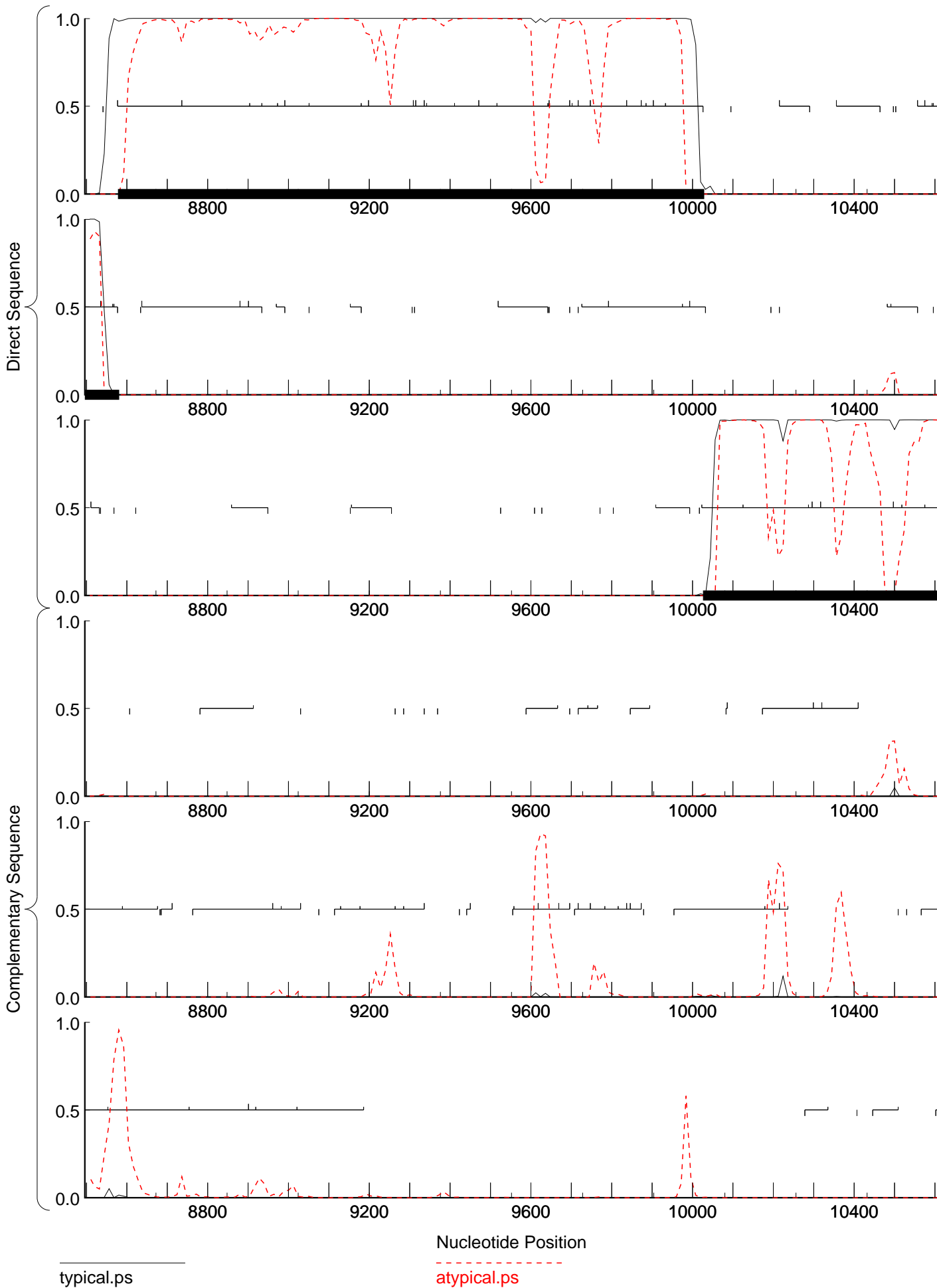
atypical.ps

... complete sequence, 51393 bp including 10 bp 3' overhang (CGGCCGGTAA), Cluster A4, Order 4, Window 96, Step 12, 3/26
... complete sequence, 51393 bp including 10 bp 3' overhang (CGGCCGGTAA), Cluster A4, Order 2, Window 96, Step 12, 3/26

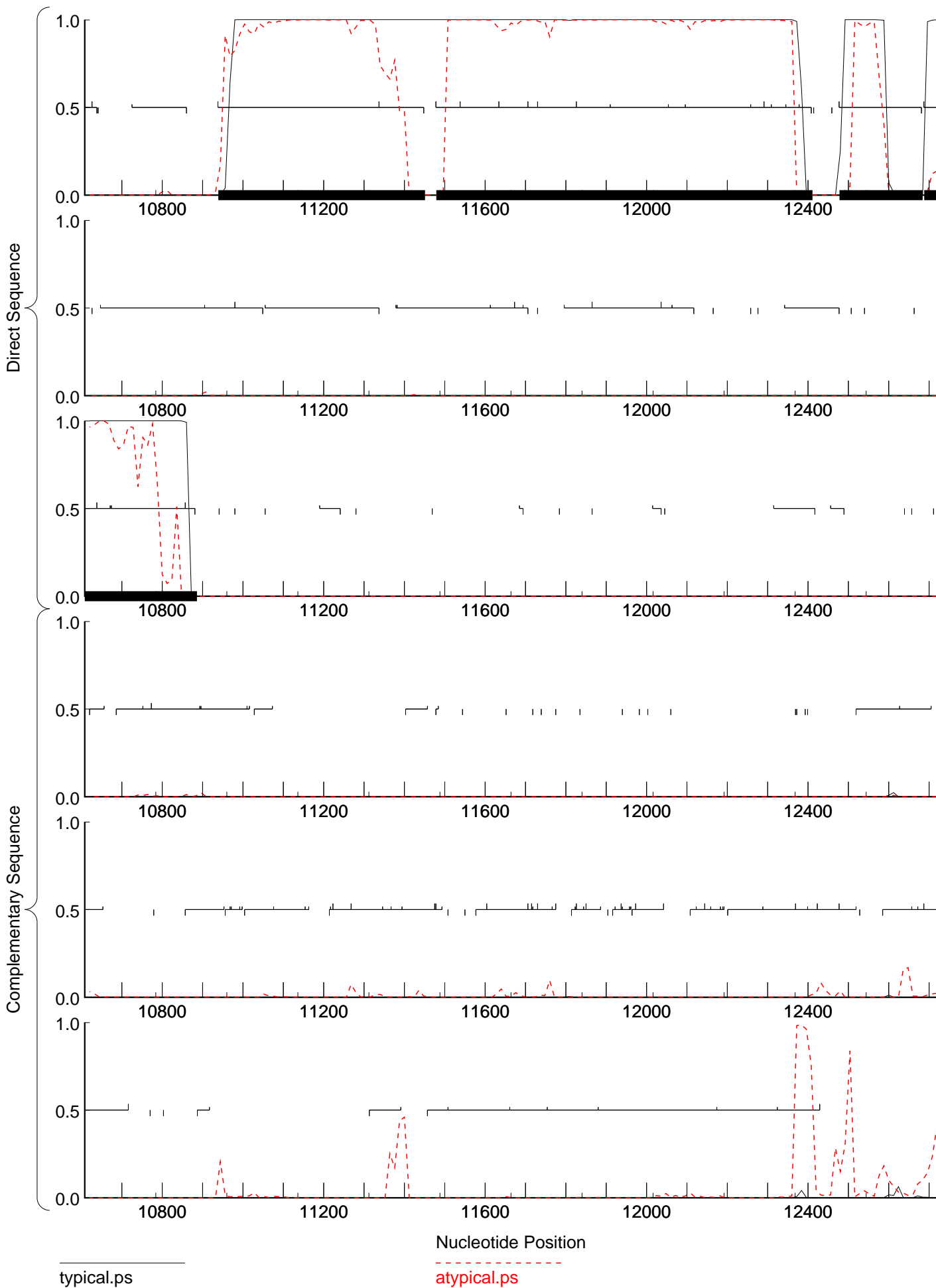


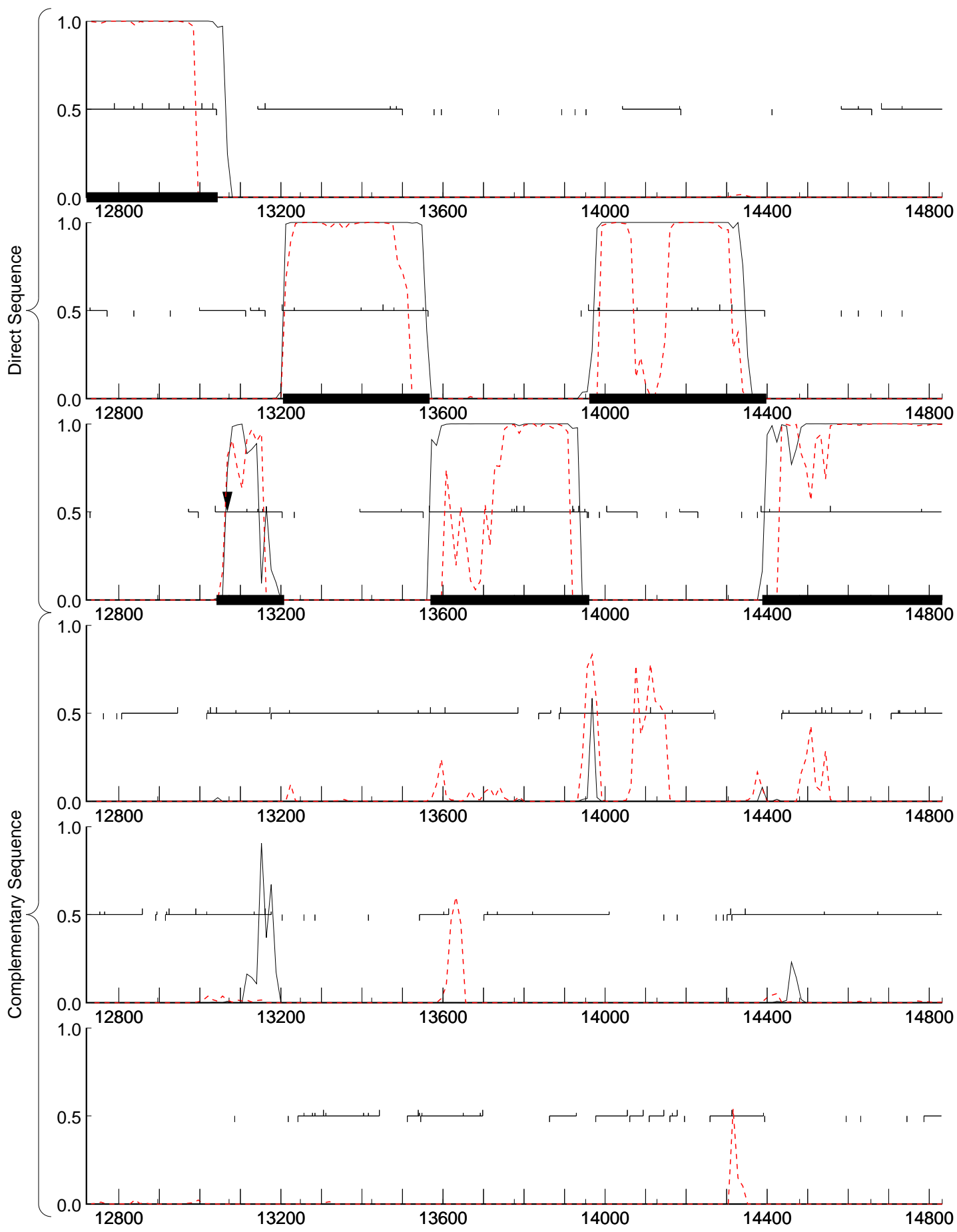






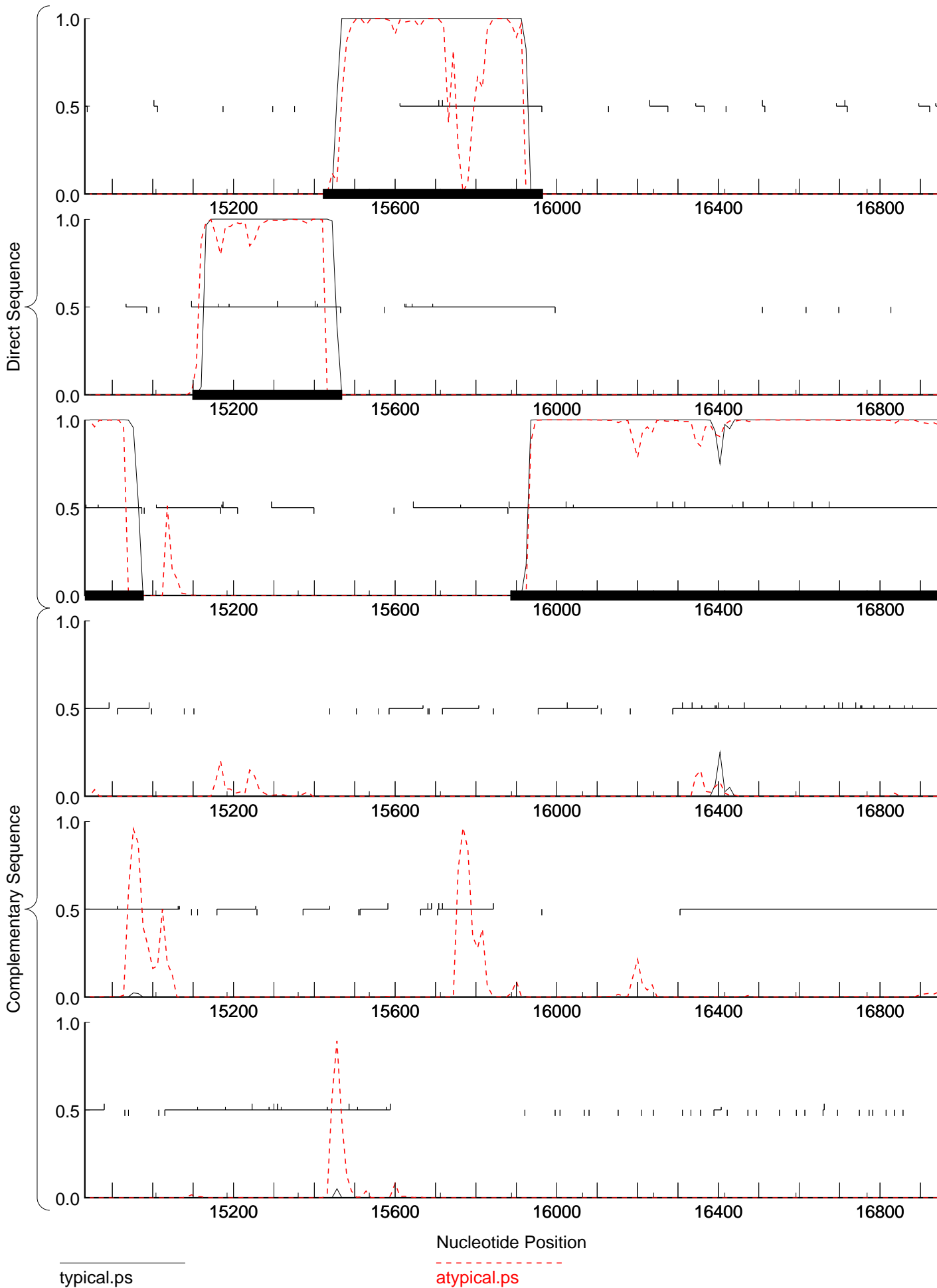
GeneMark.hmm, prediction
tius complete sequence, 51393 bp including 10 bp 3' overhang (CGGCCGGTAA), Cluster A4, Order 4, Window 96, Step 12, 7/26
tius complete sequence, 51393 bp including 10 bp 3' overhang (CGGCCGGTAA), Cluster A4, Order 2, Window 96, Step 12, 7/26

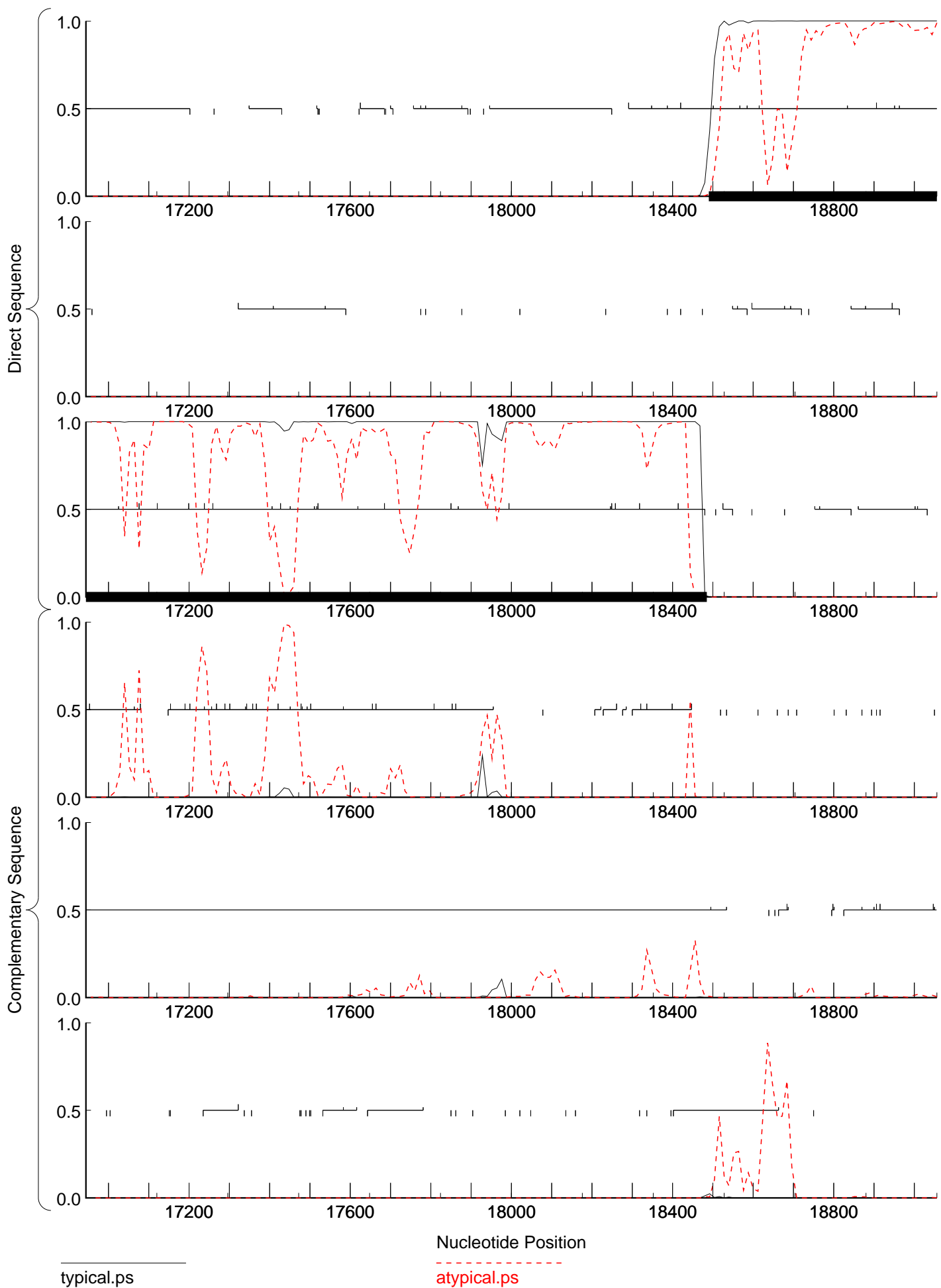


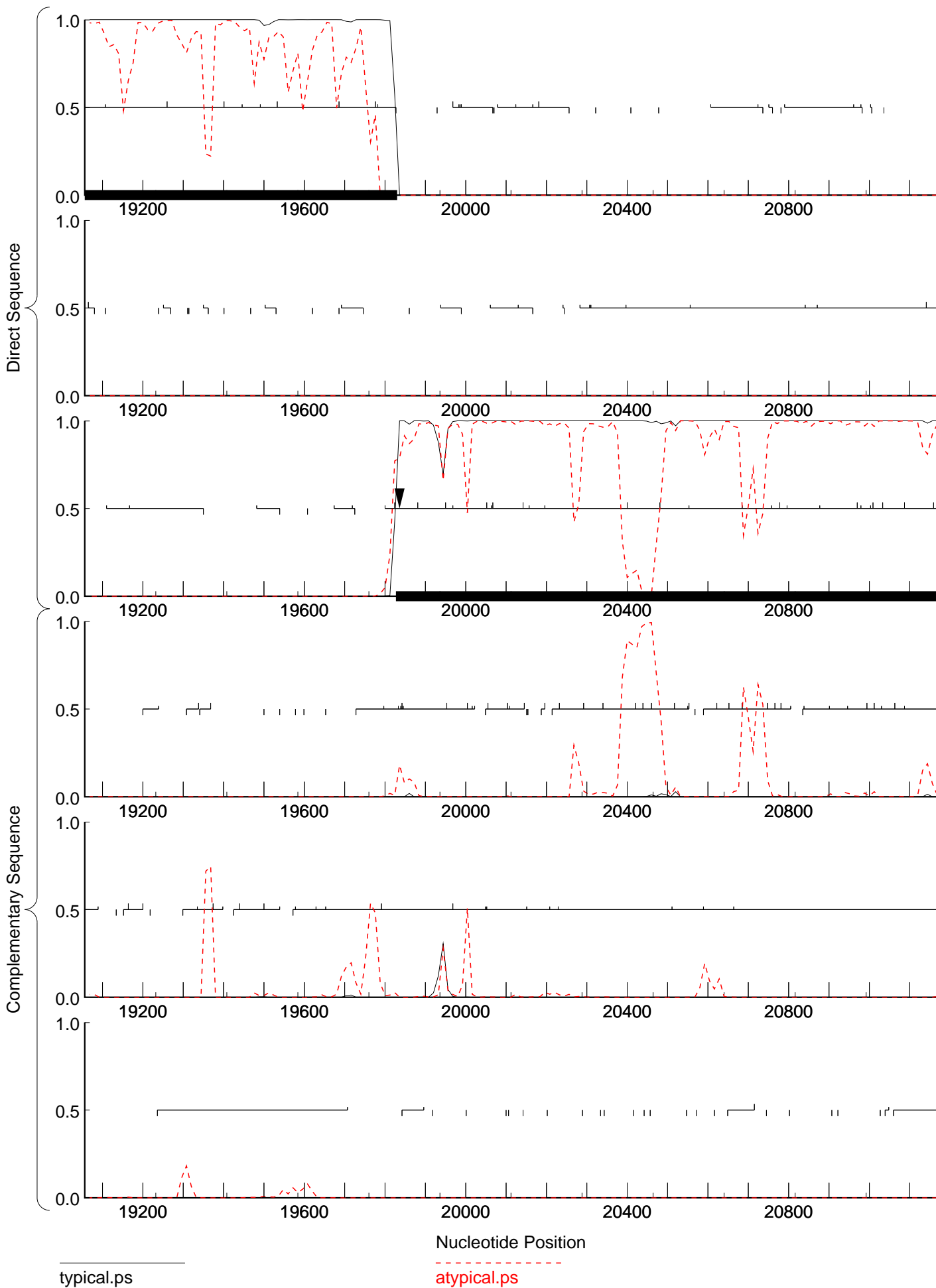


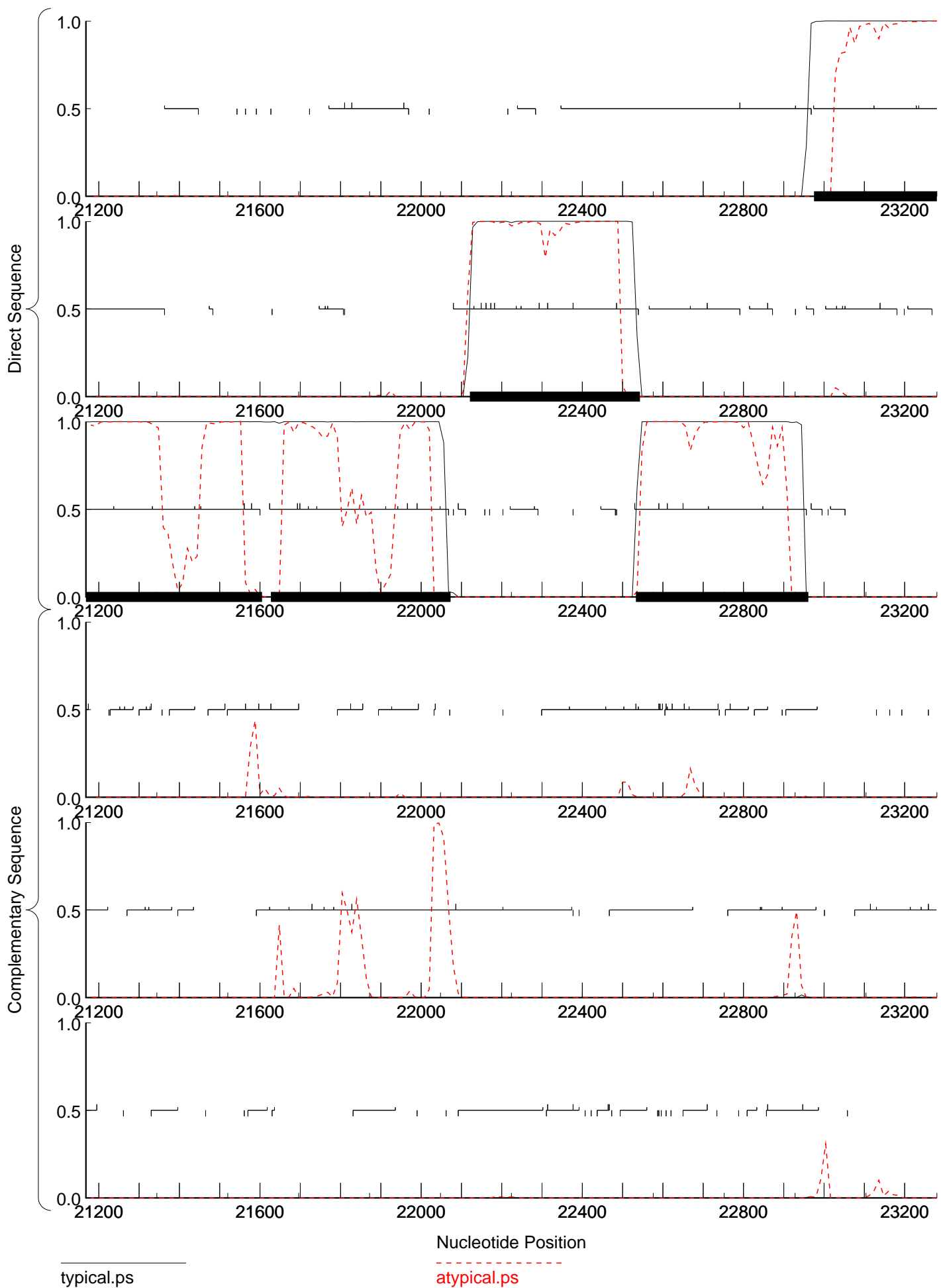
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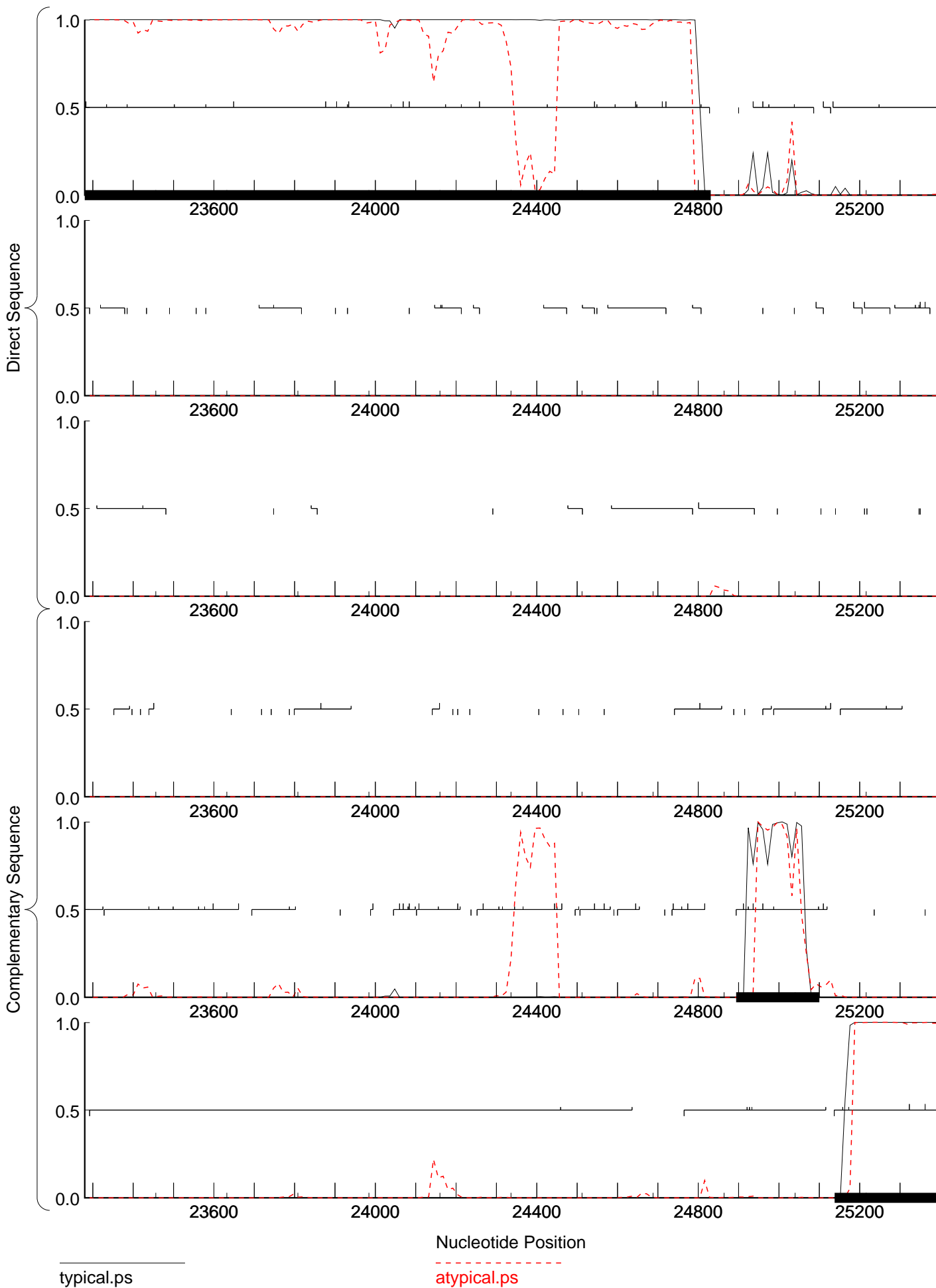
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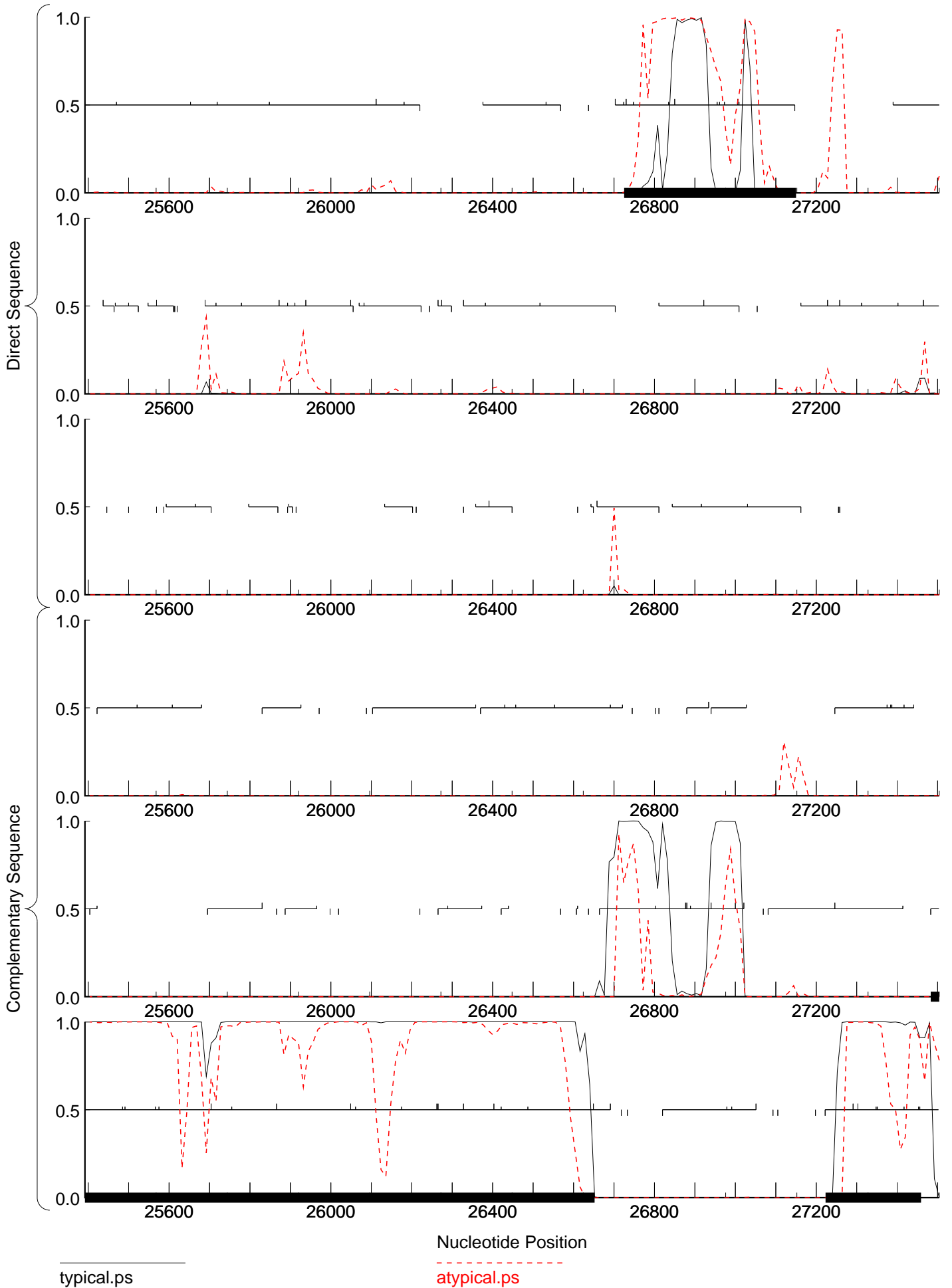


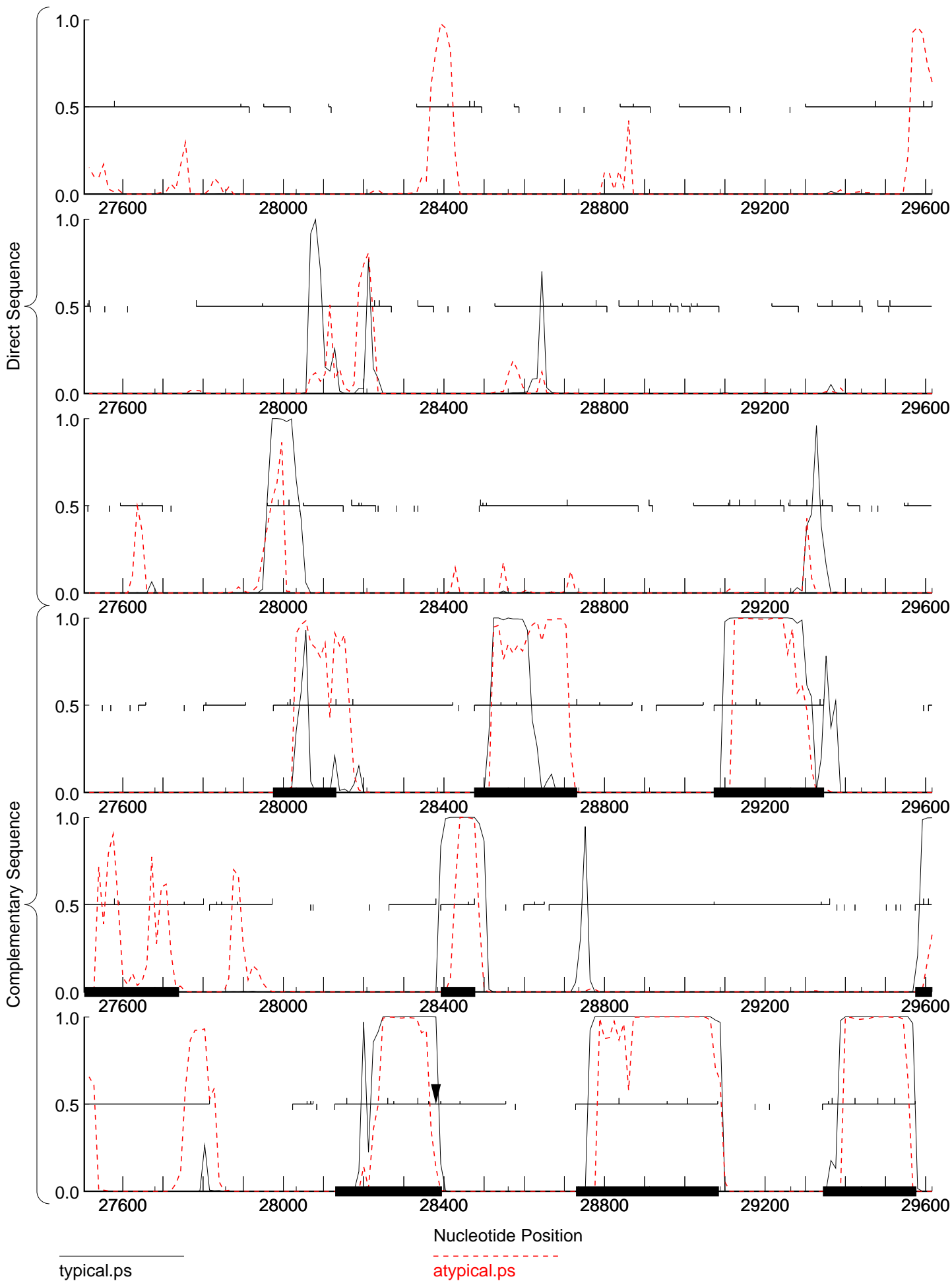


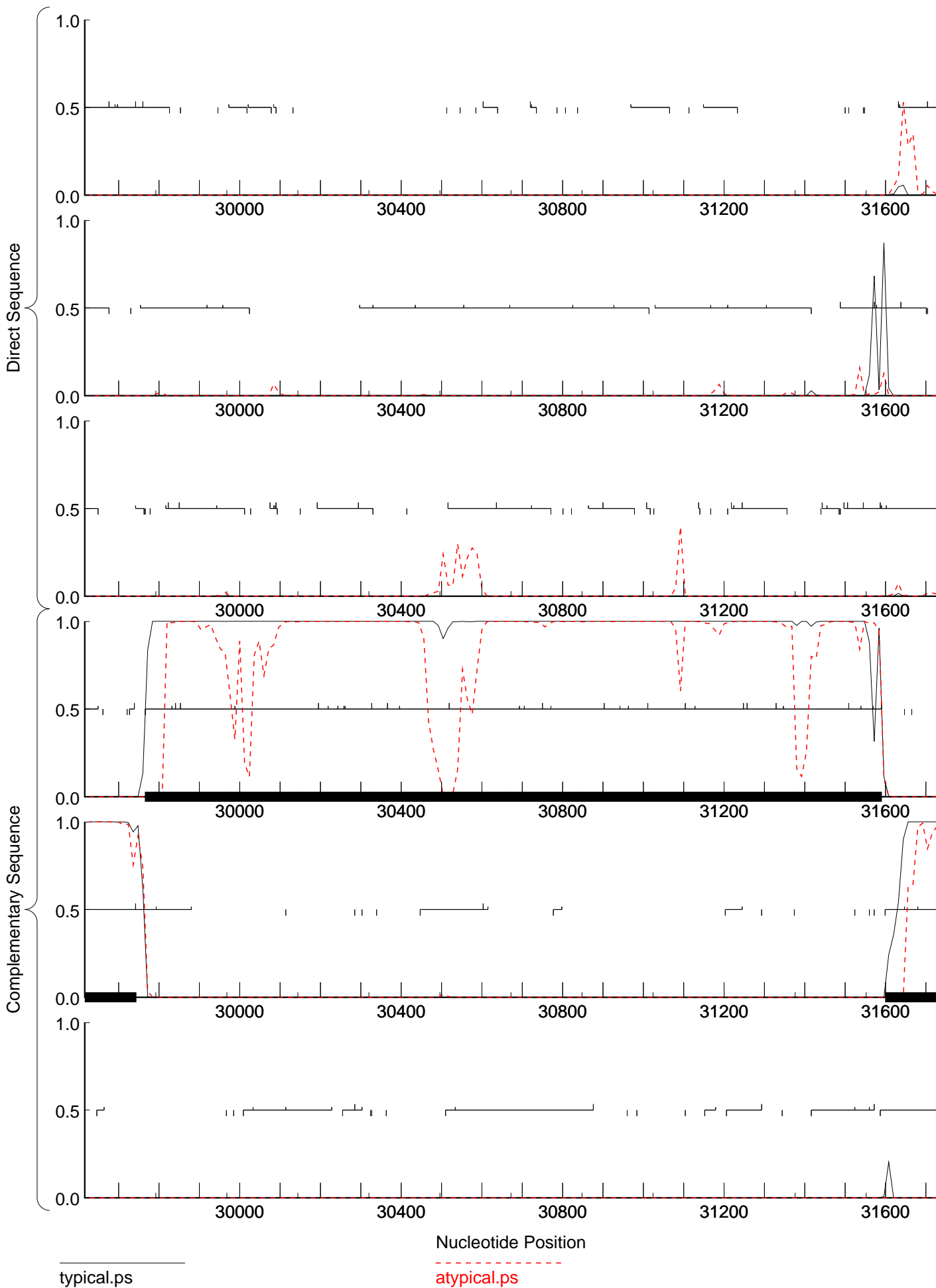






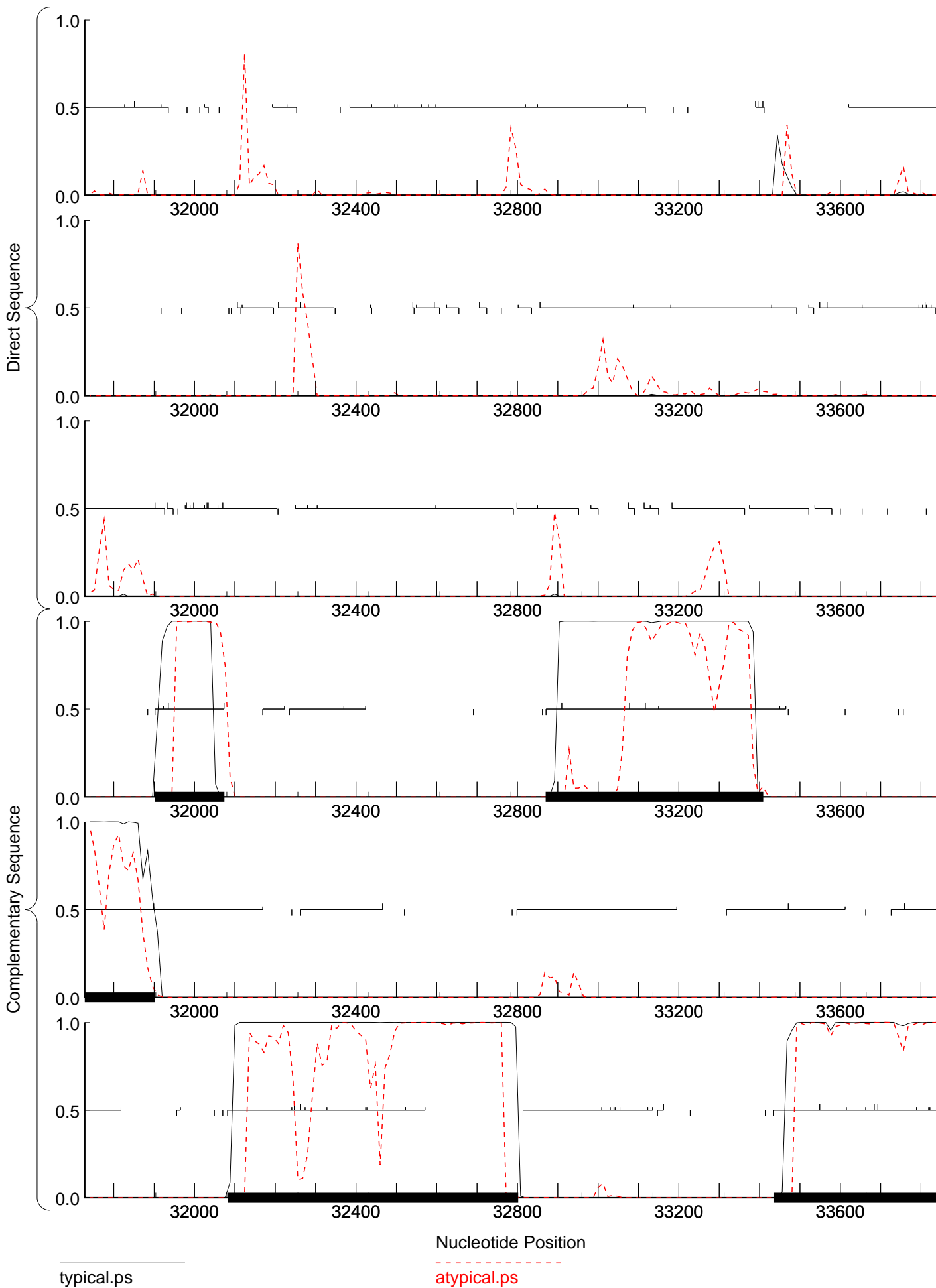


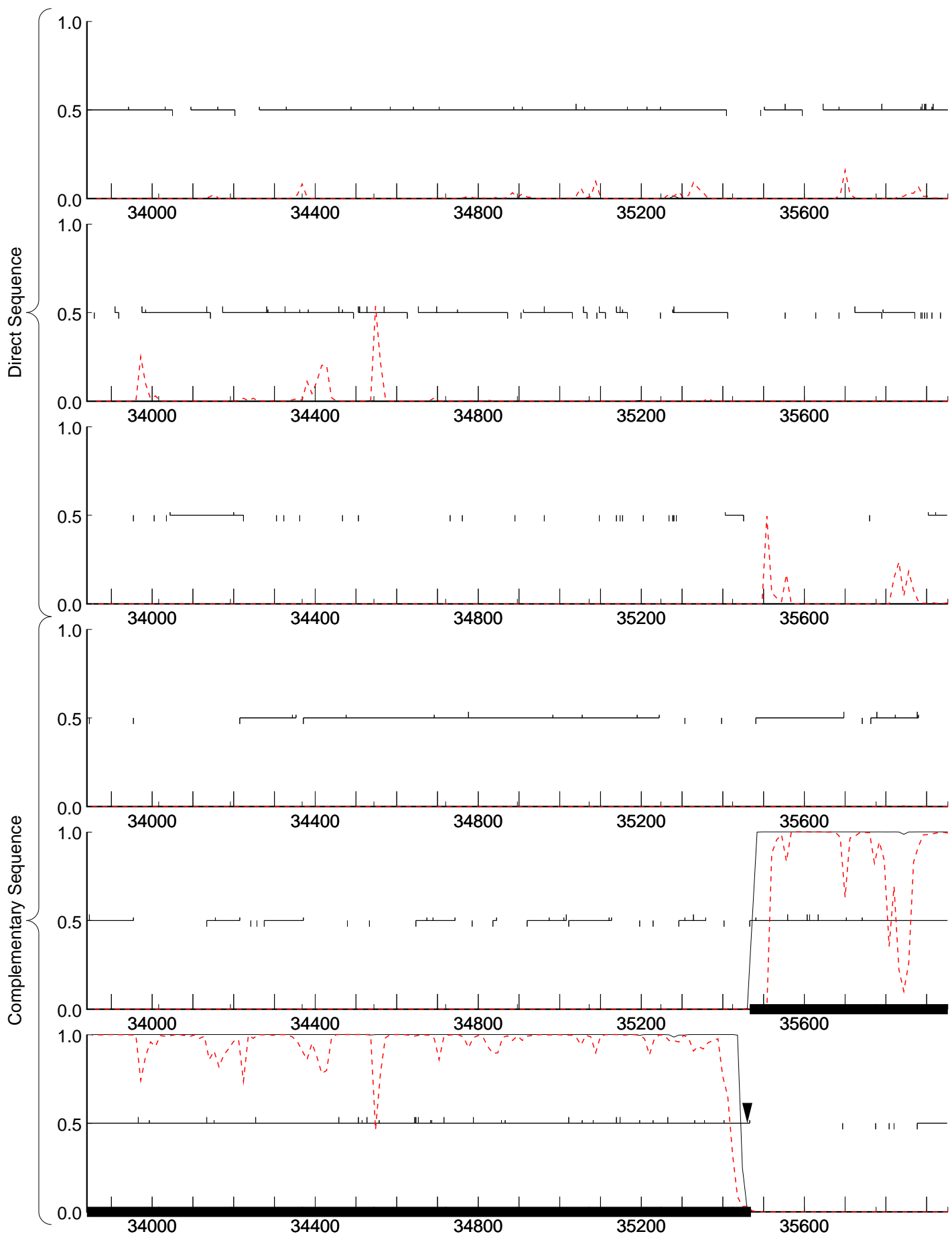




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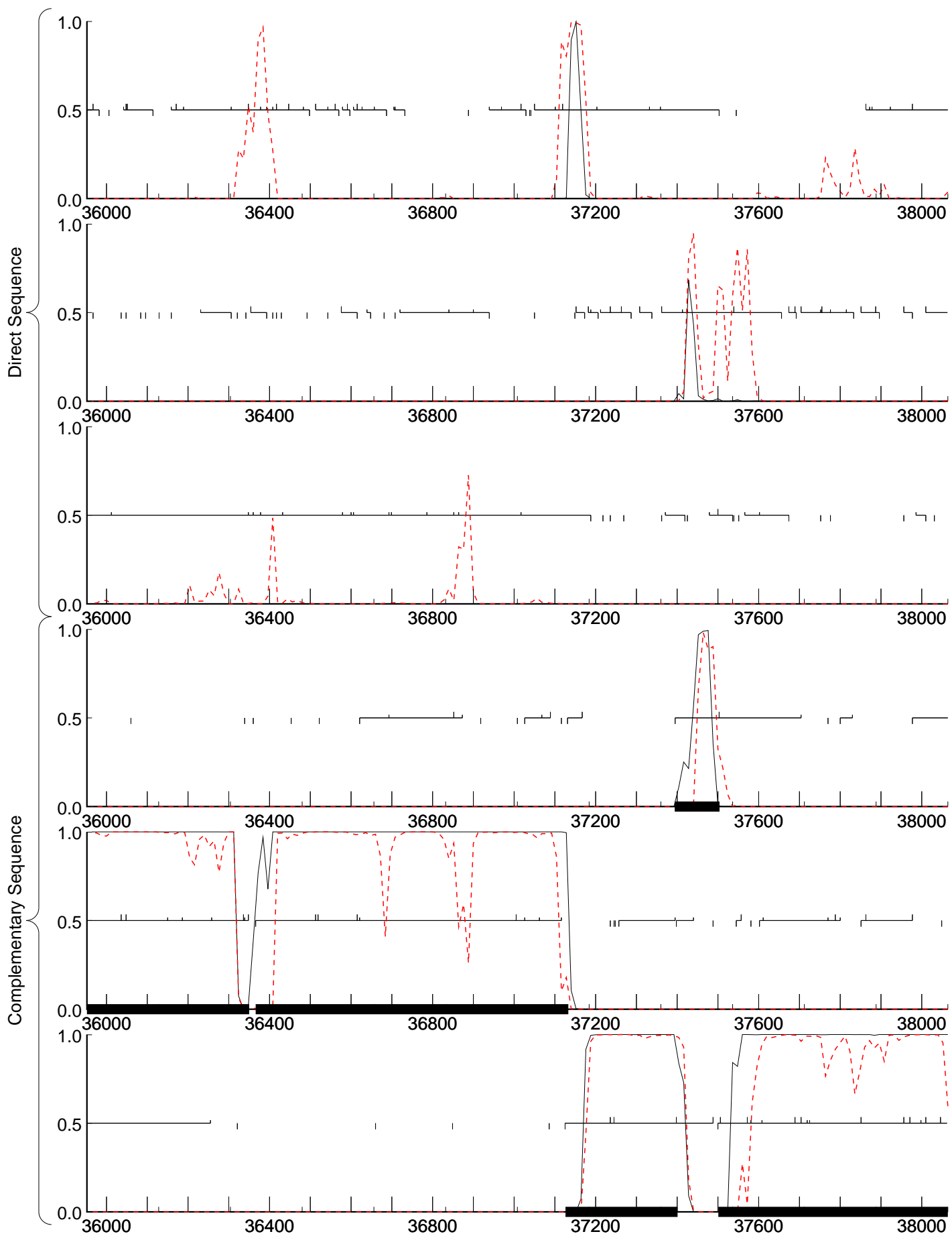
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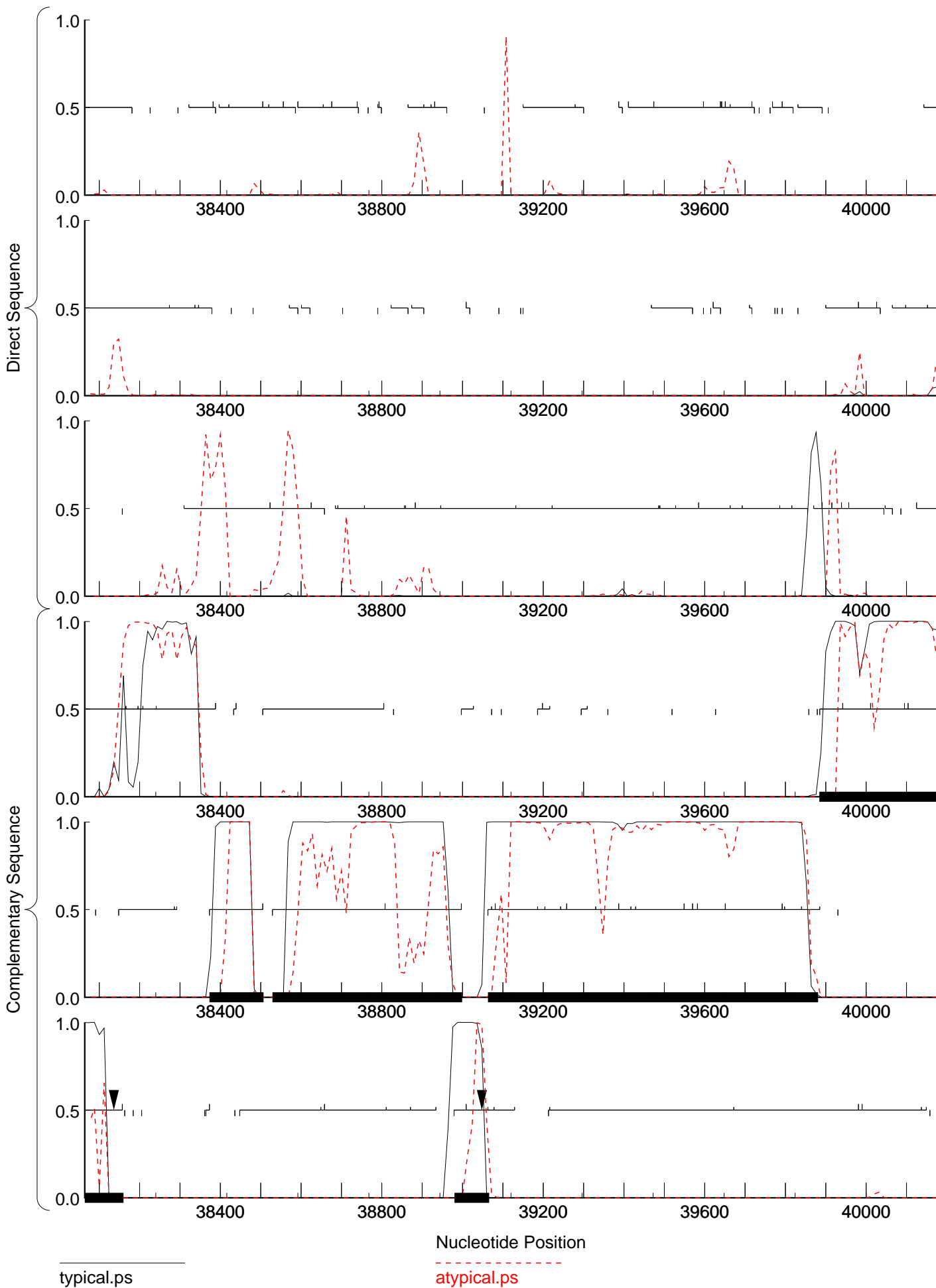
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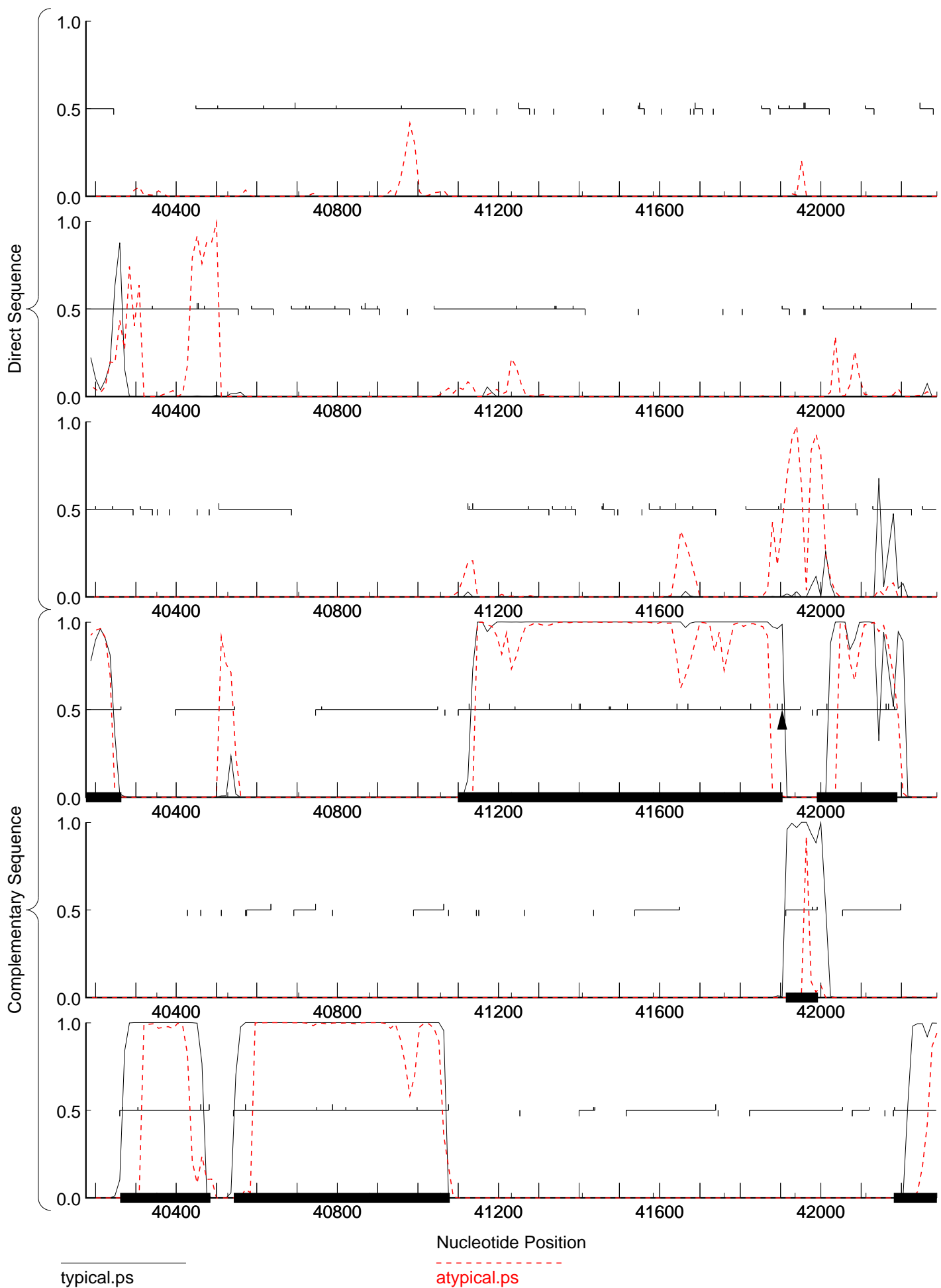
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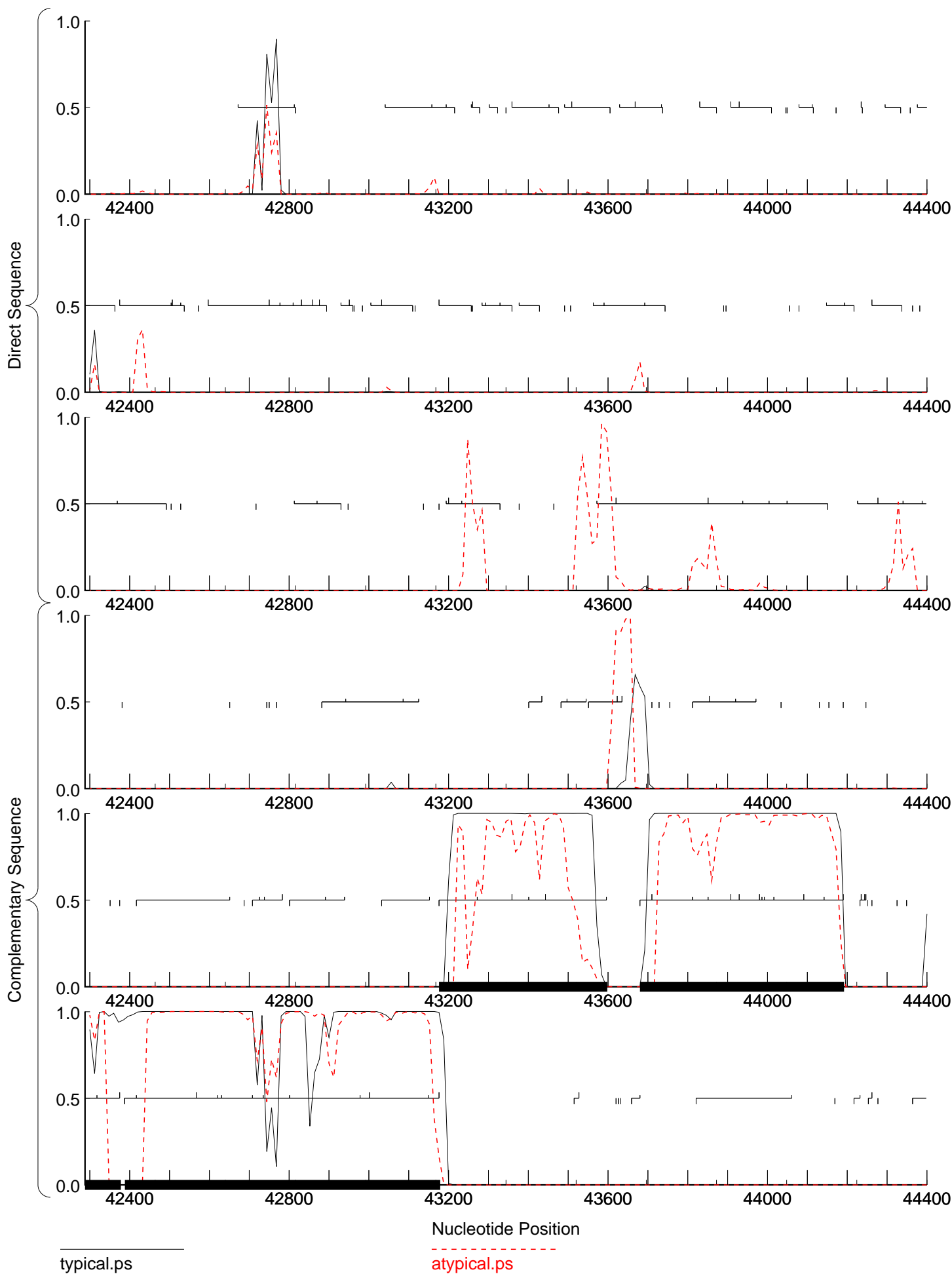
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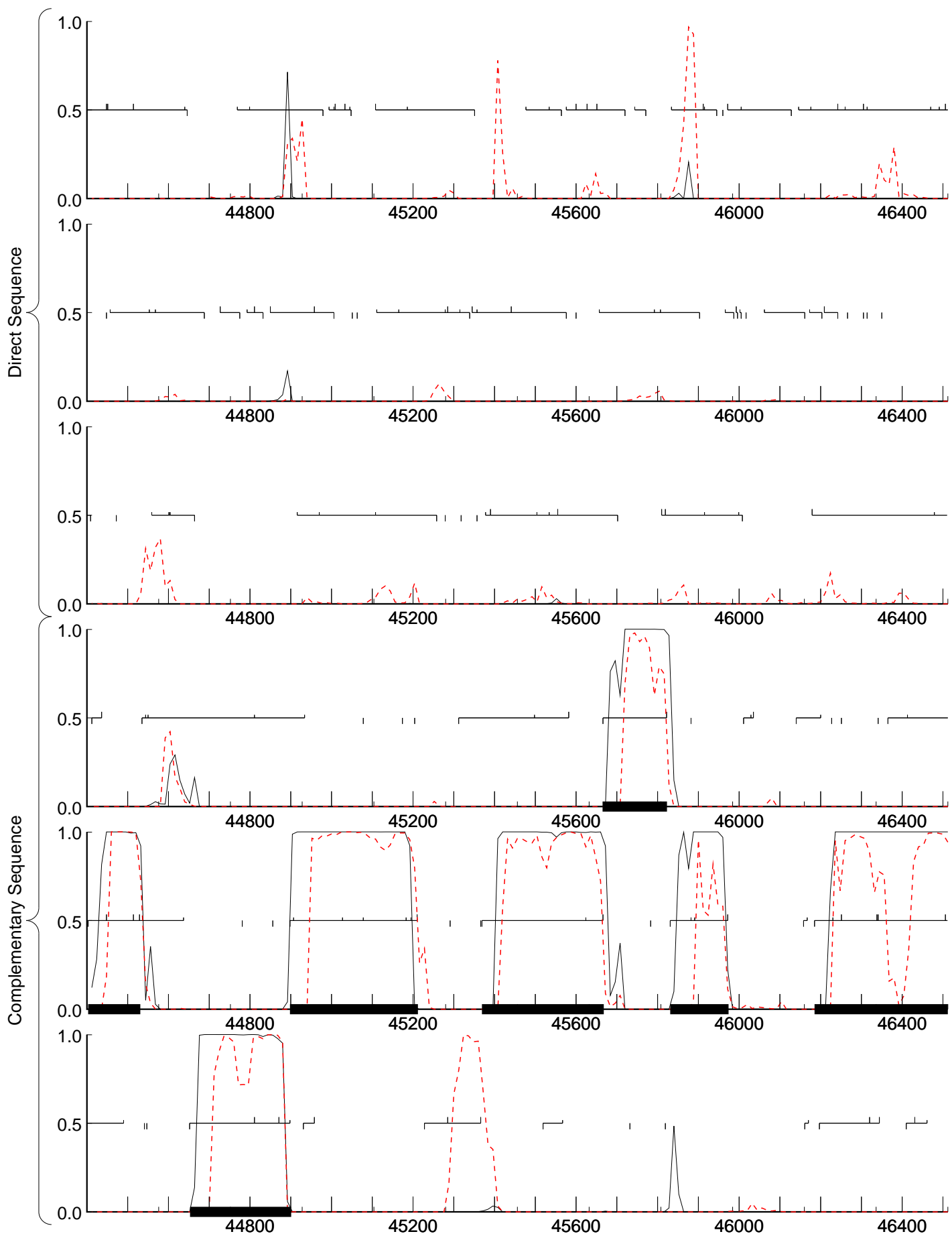




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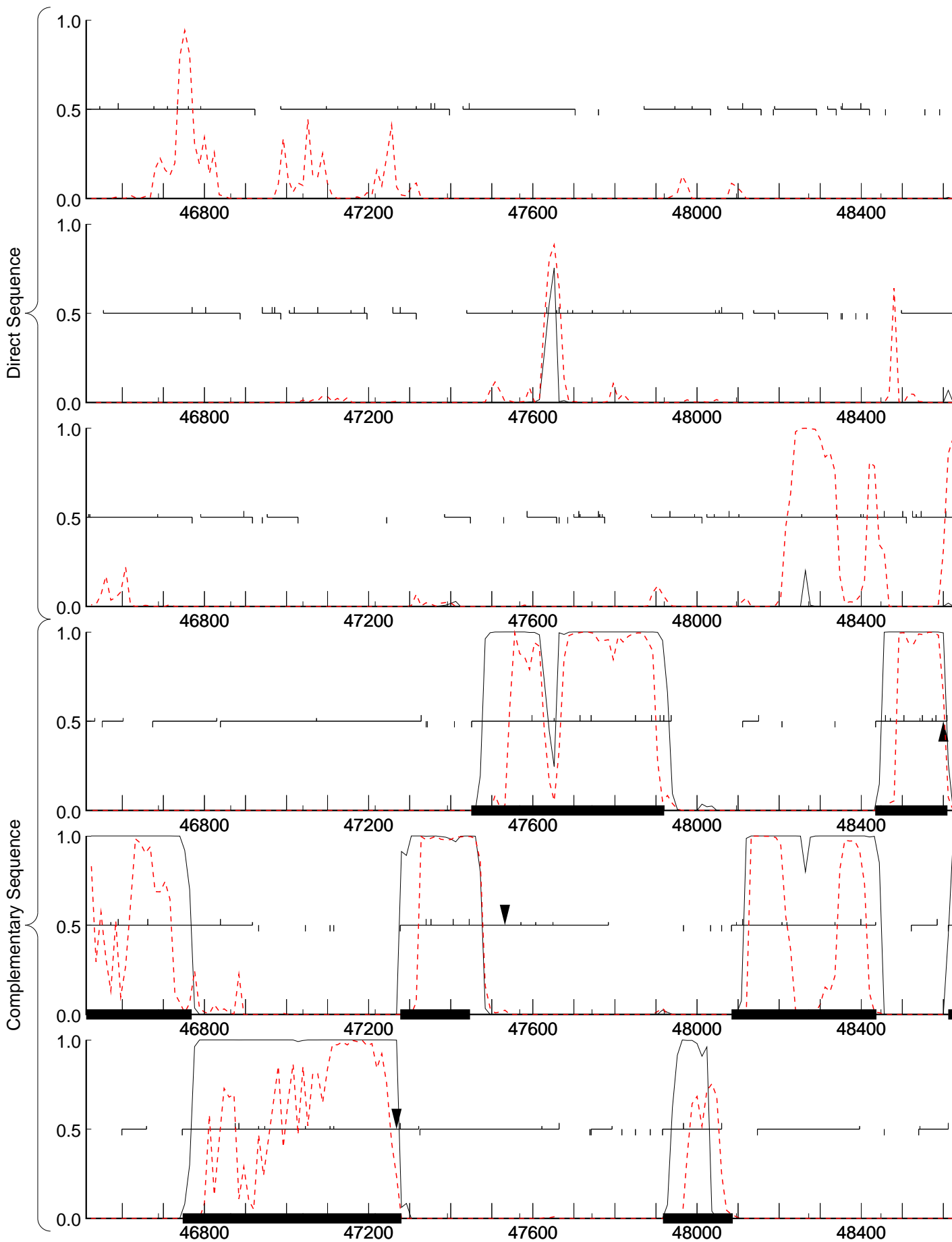
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