

GeneMark

Version 2.5p (09.08.06)

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PROGRAM INFORMATION

Sequence : Mycobacterium phage Citius complete sequence, 51393 bp including 10-base 3' overhang (CGCC
Analysis Date : 7/24/17 at 22:32:05
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,

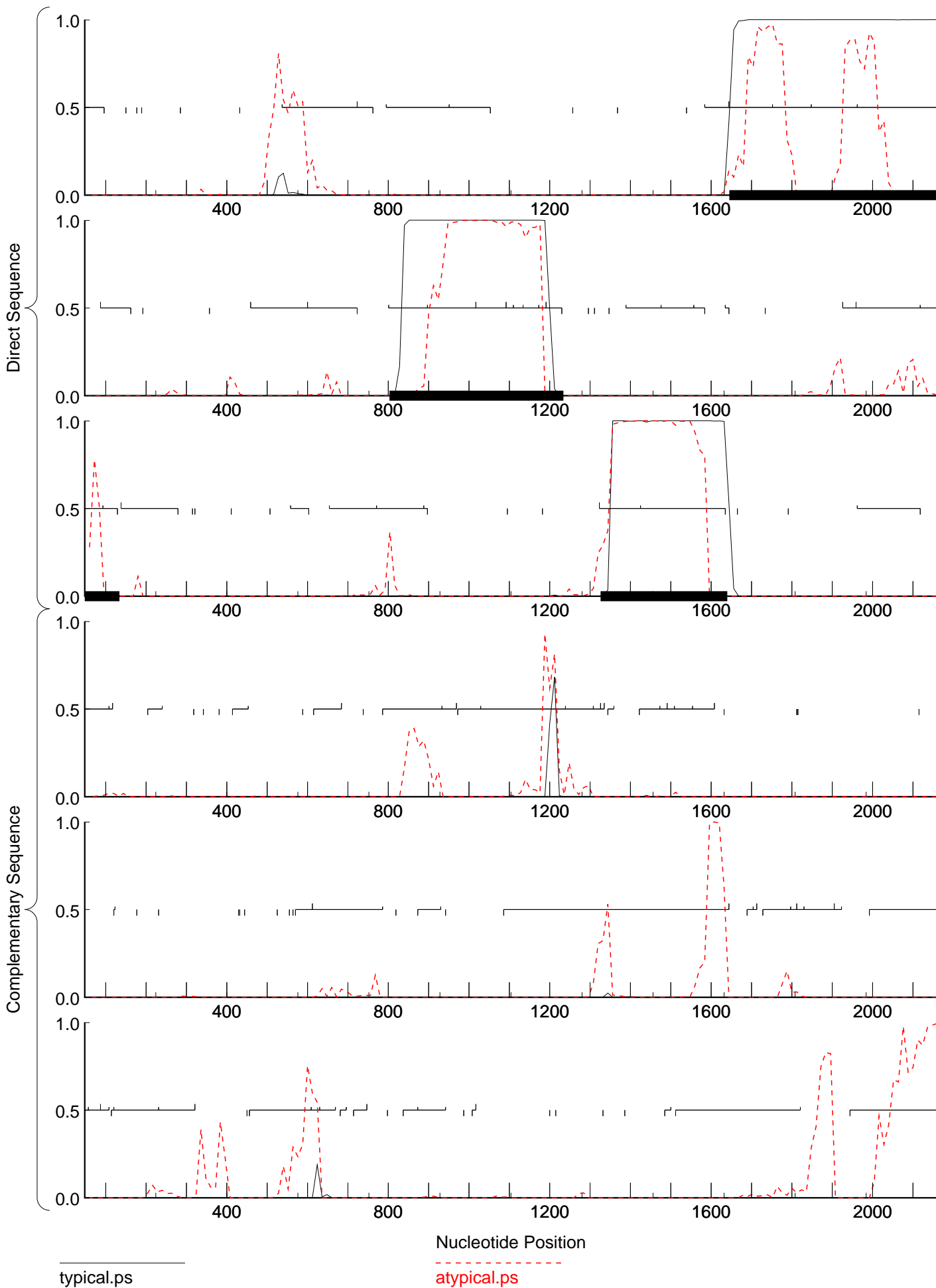
Matrix notes & comments

Training set derived by GeneMarkS, 4.30 October 2014
Mon Jul 24 22:32:05 2017

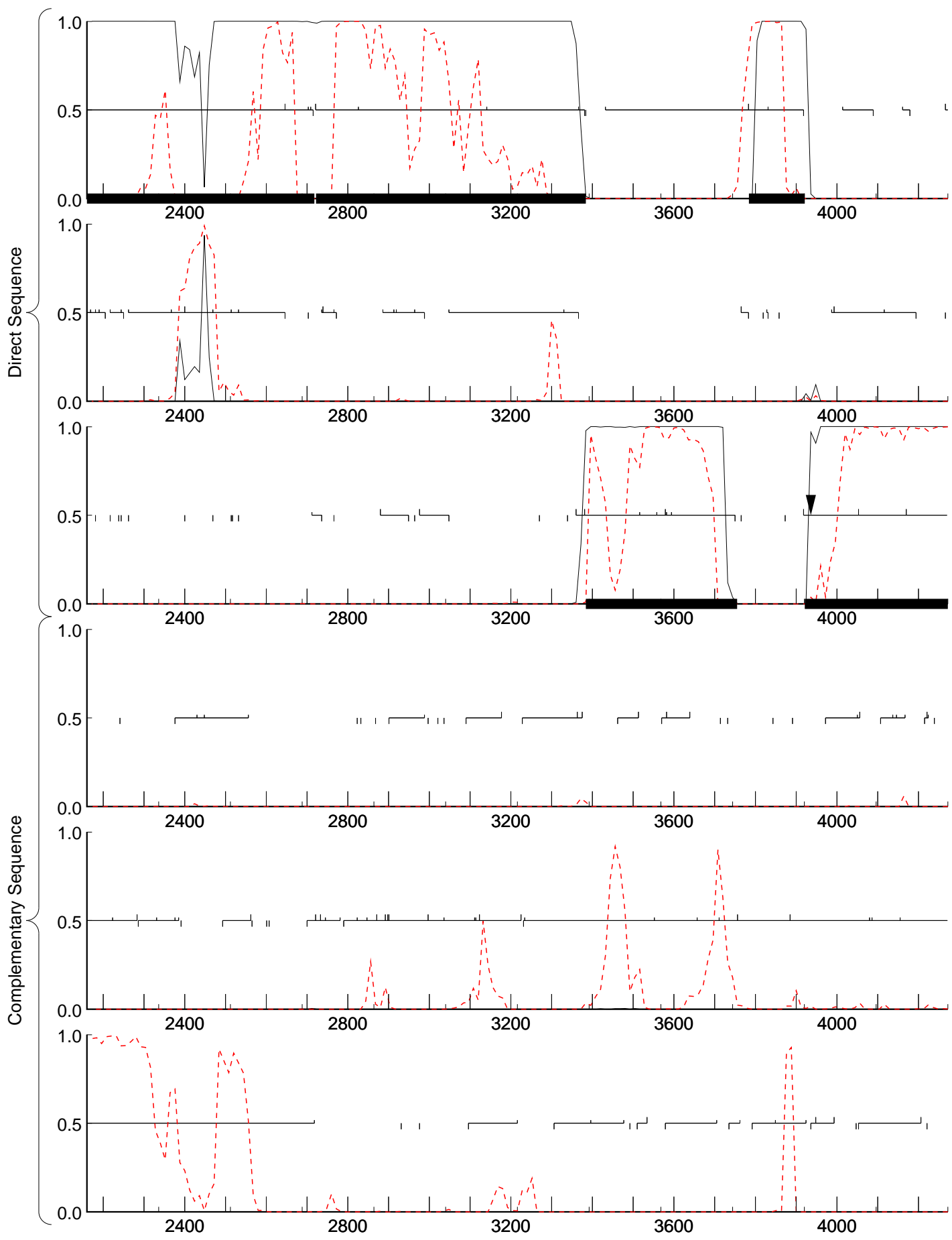
MATRIX INFORMATION

Matrix : -
Author : -
Order : 4

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

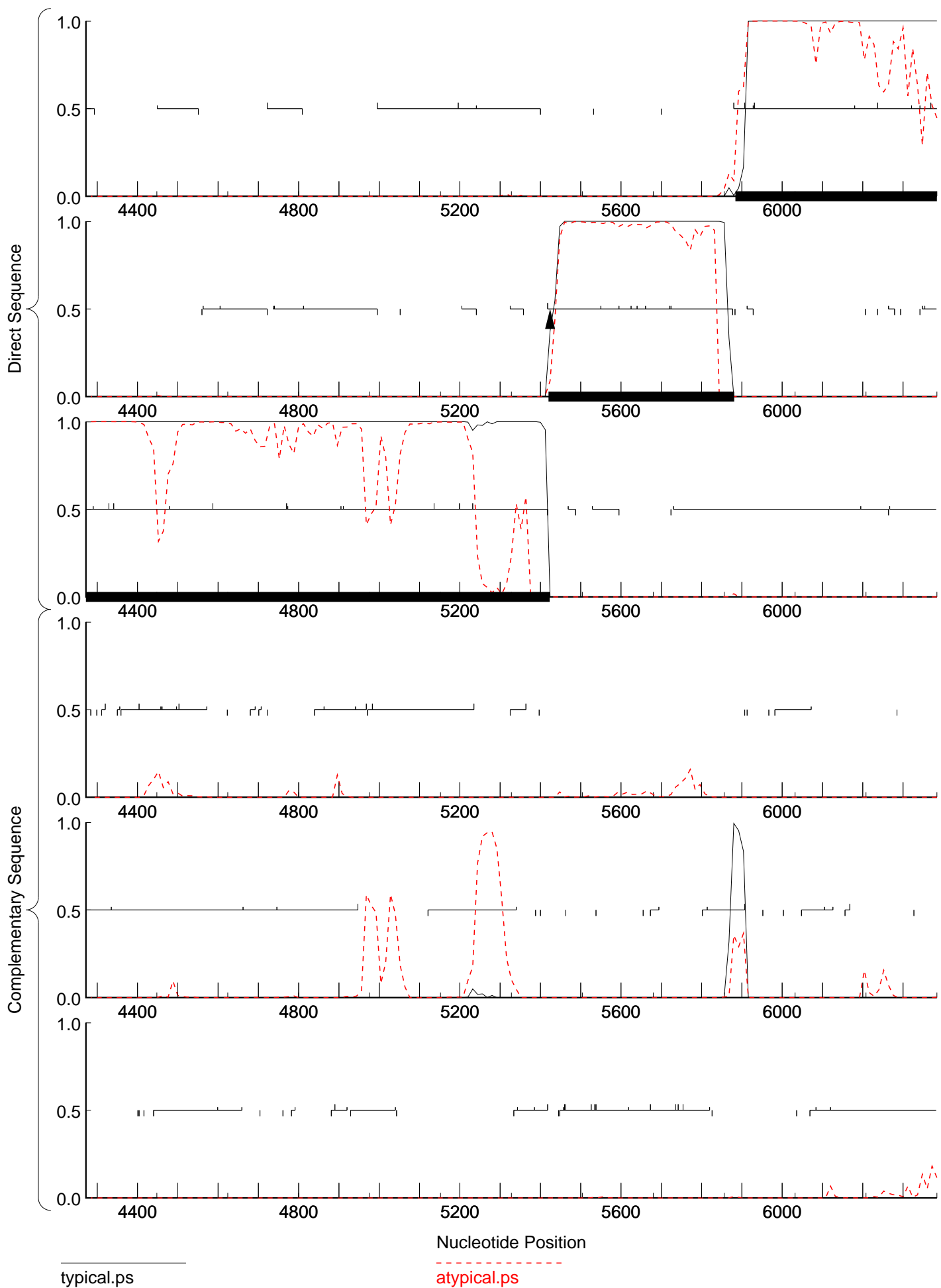


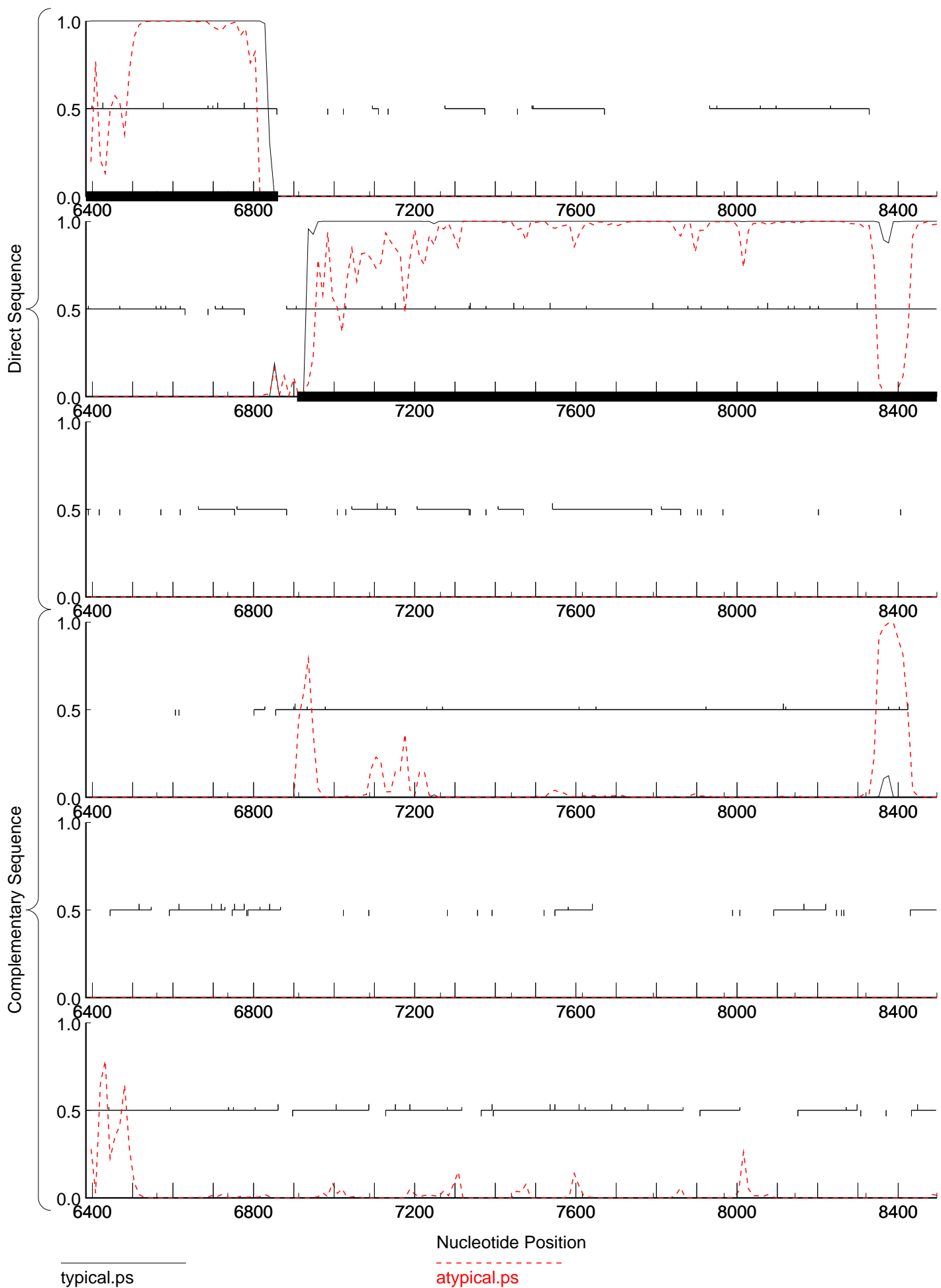
s complete sequence, 51393 bp including 10-base 3' overhang (CGGCCGGTAA), Cluster A4, Order 4, Window 96, Step 12, 3/26
s complete sequence, 51393 bp including 10-base 3' overhang (CGGCCGGTAA), Cluster A4, Order 2, Window 96, Step 12, 3/26



typical.ps

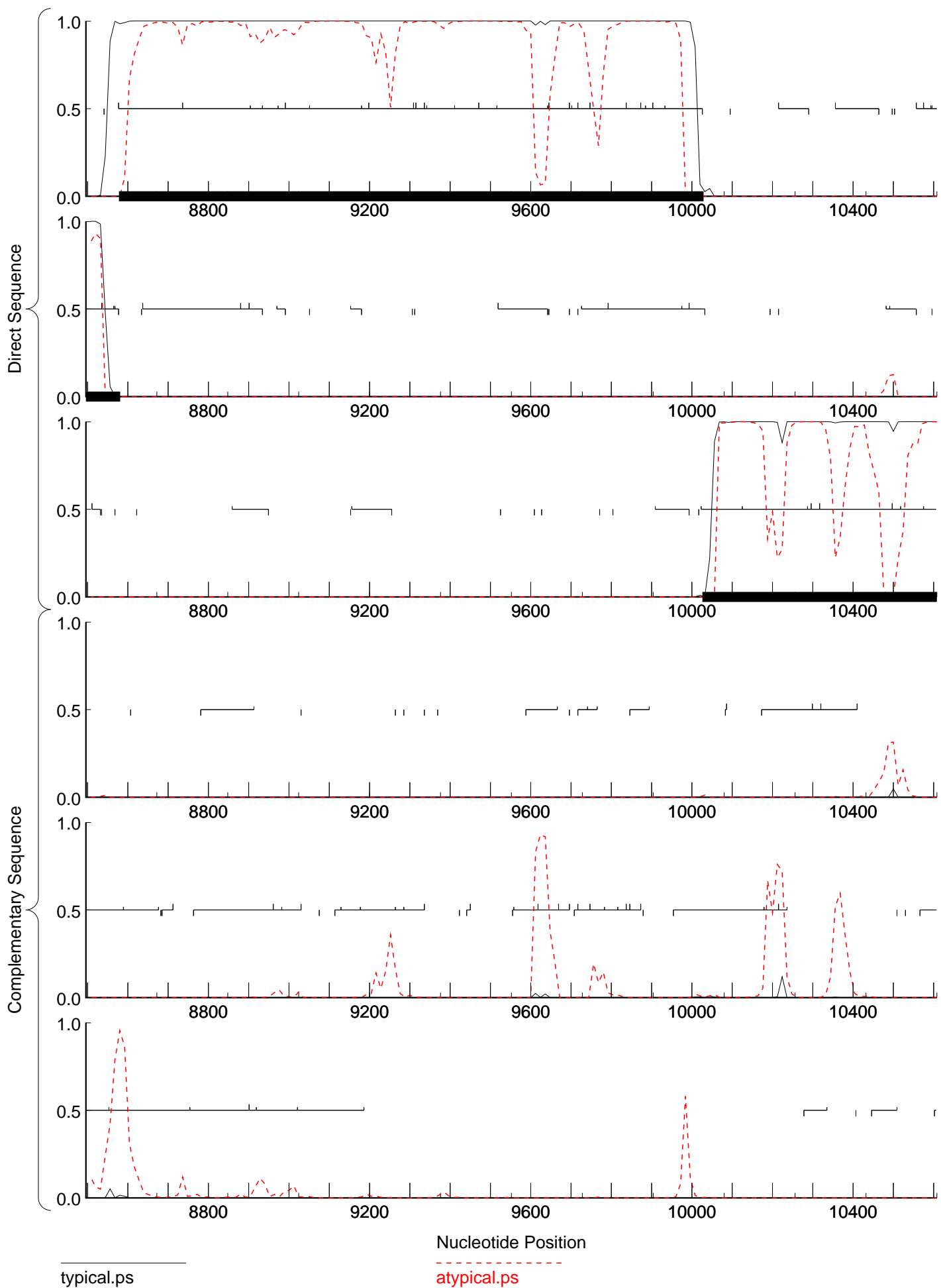
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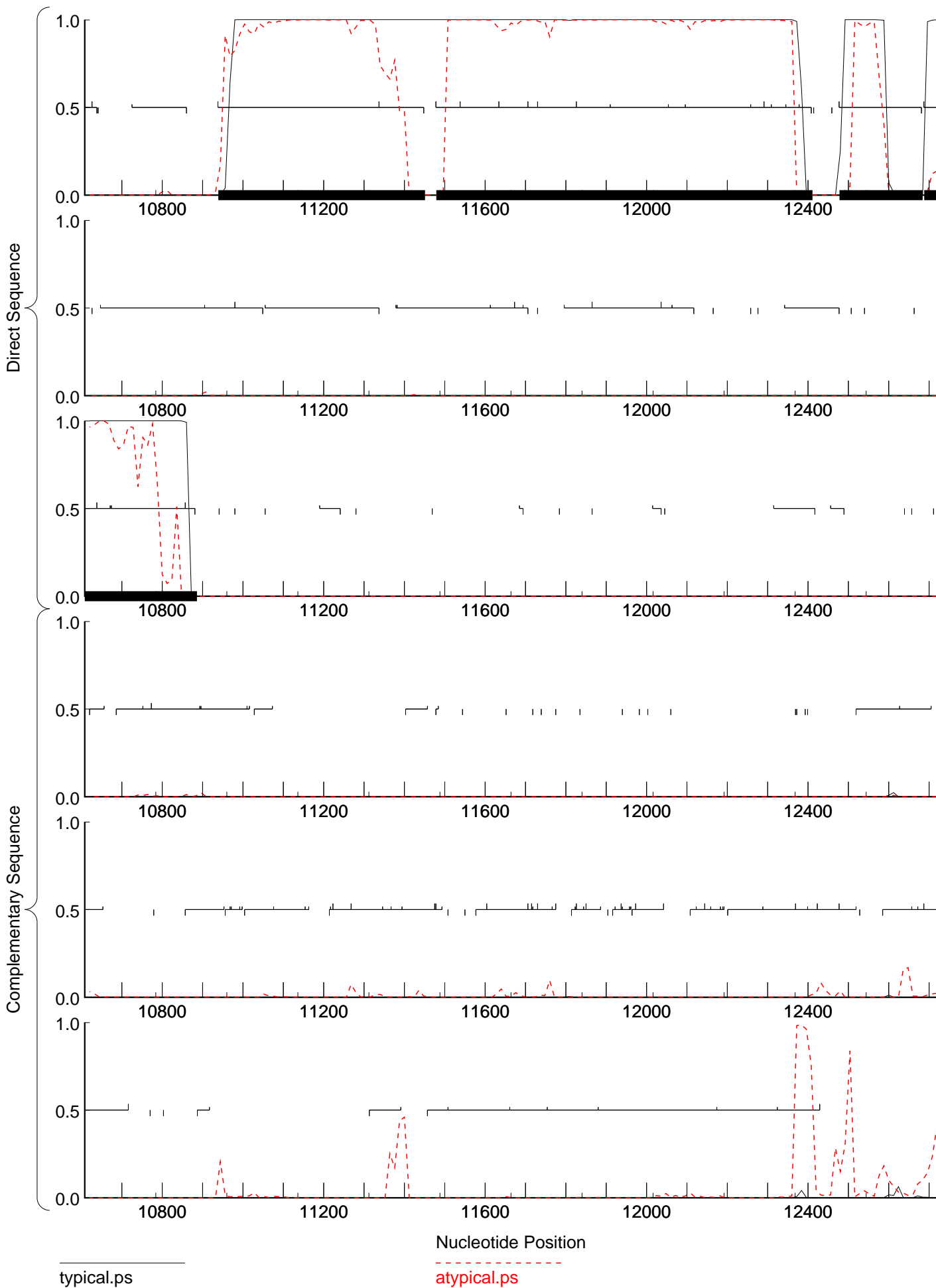
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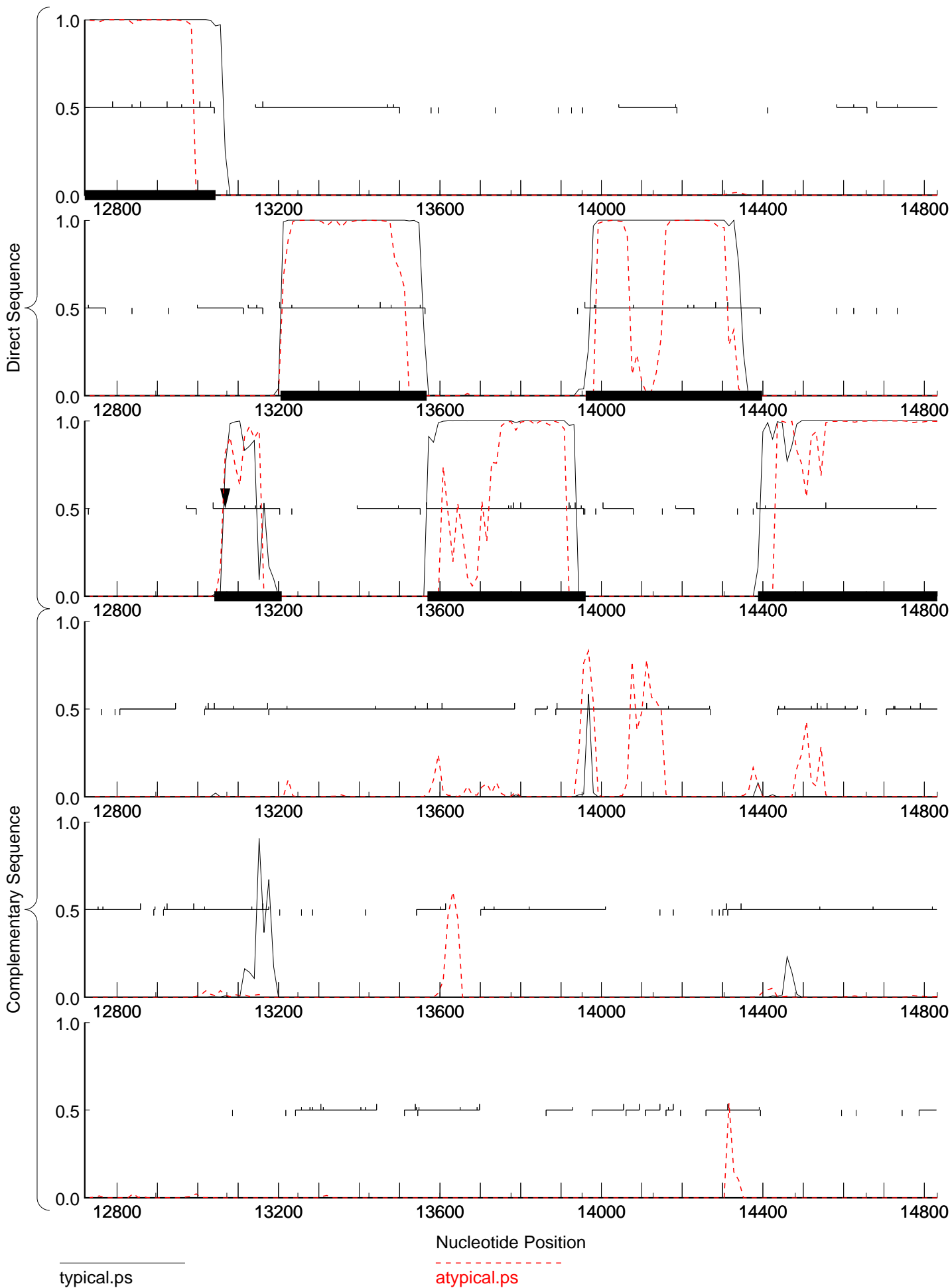
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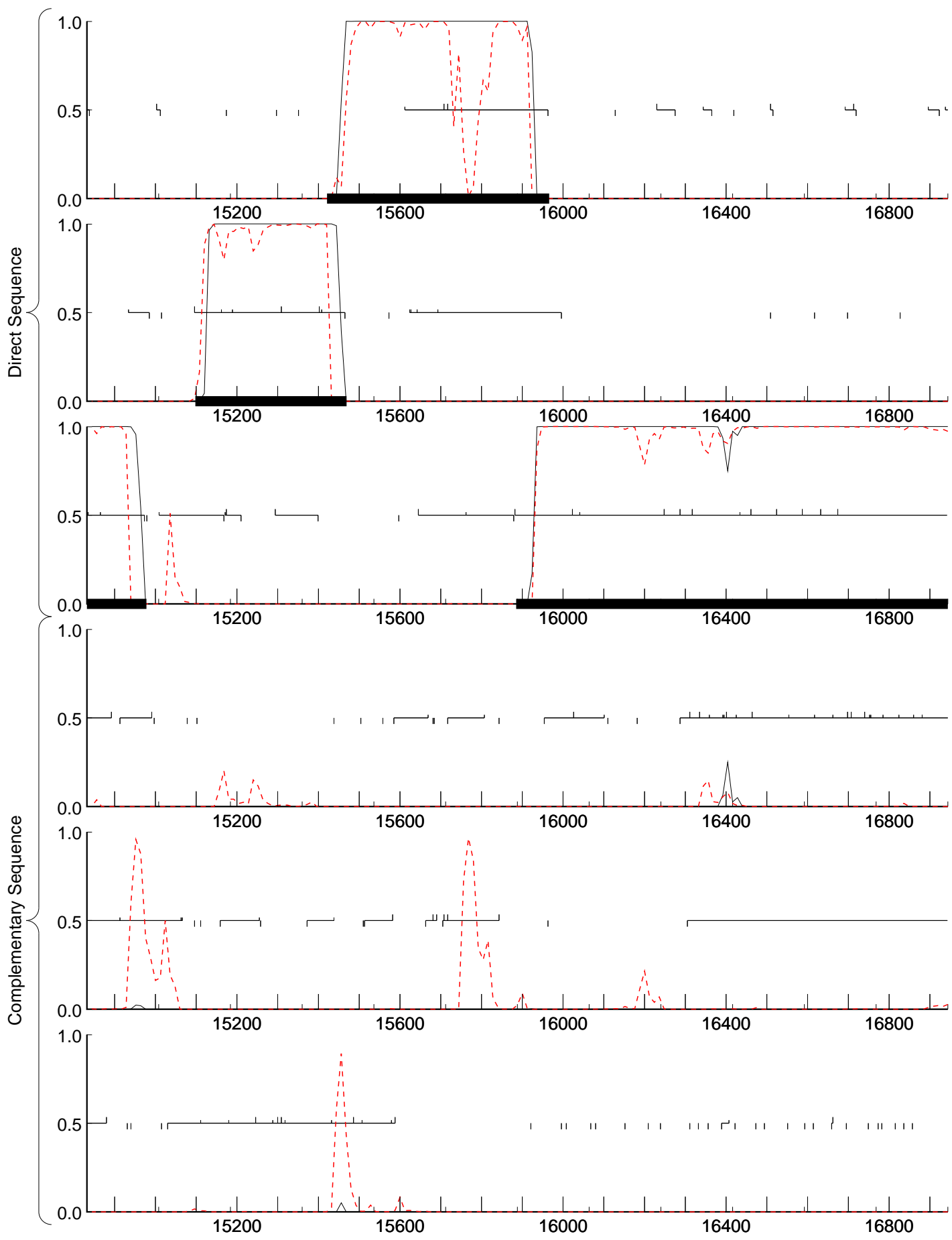


typical.ps

atypical.ps

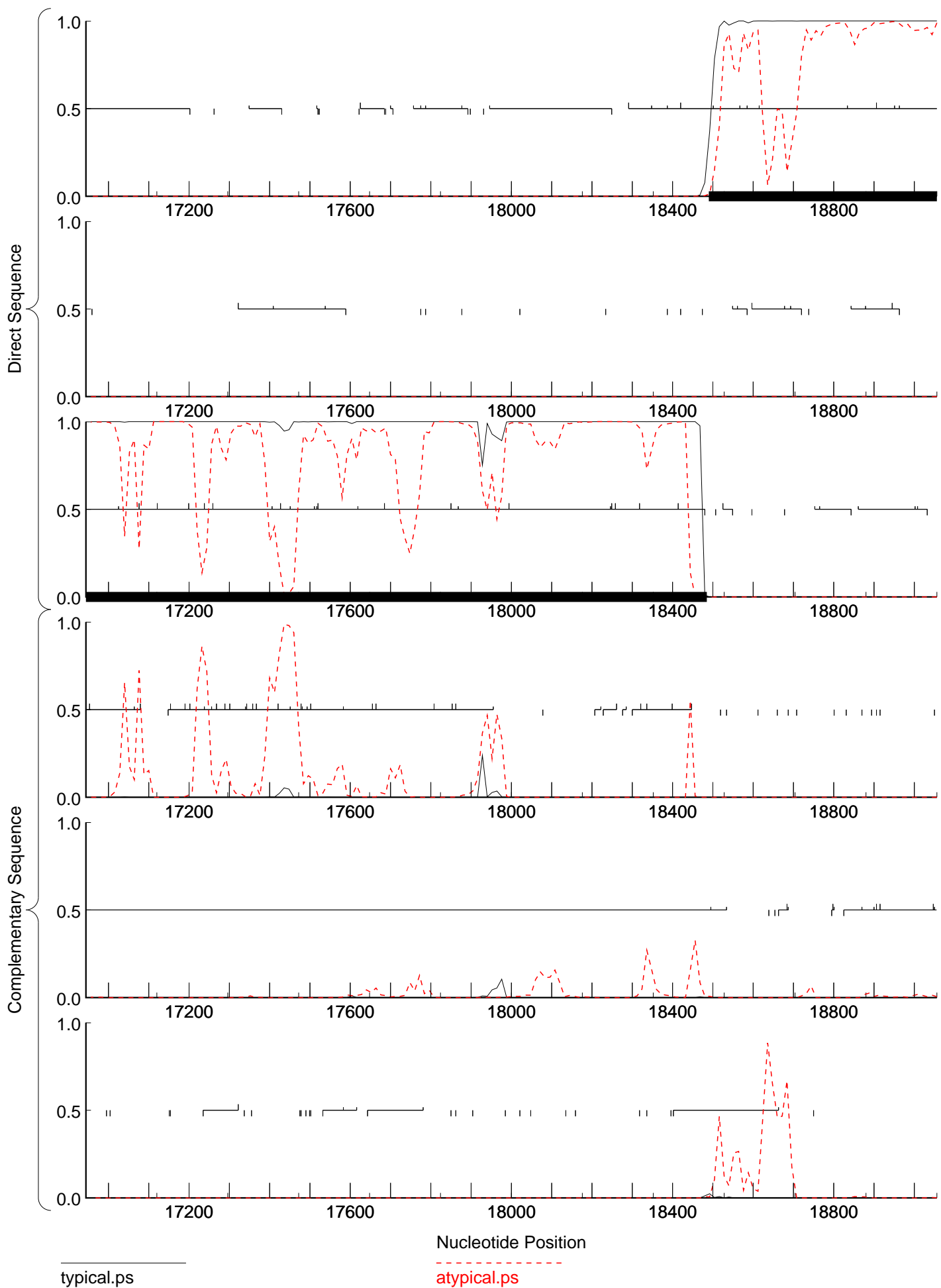


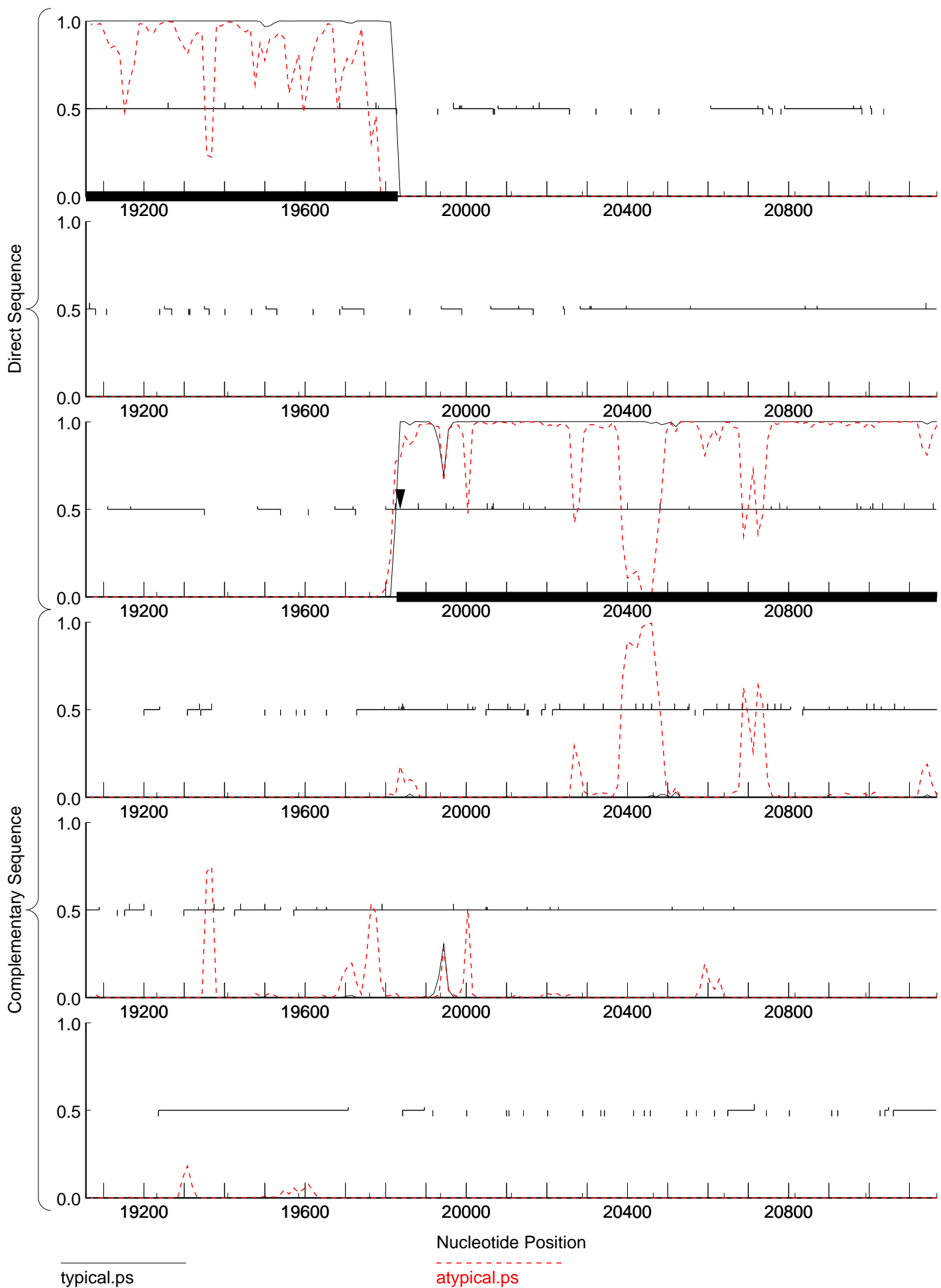




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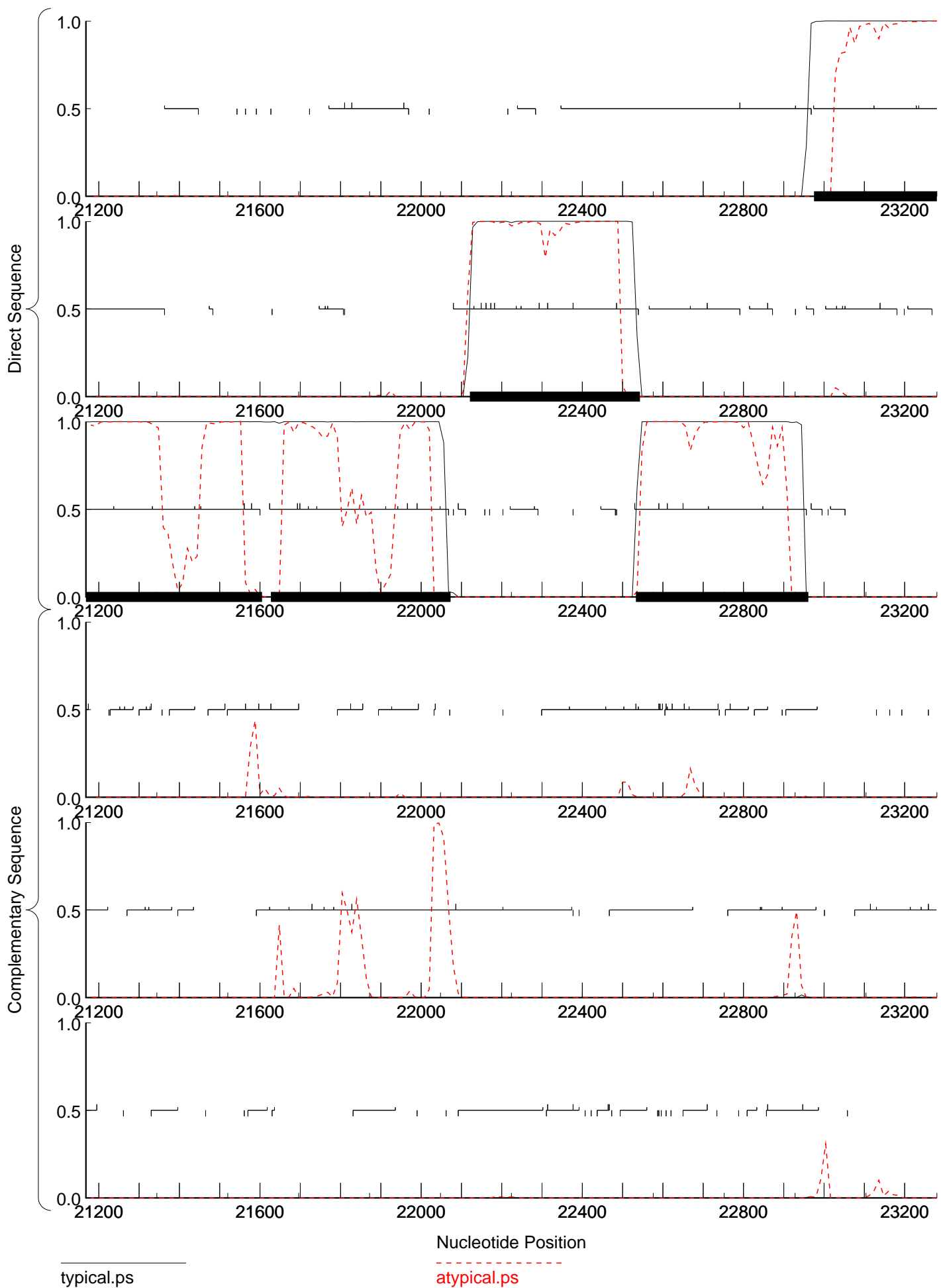
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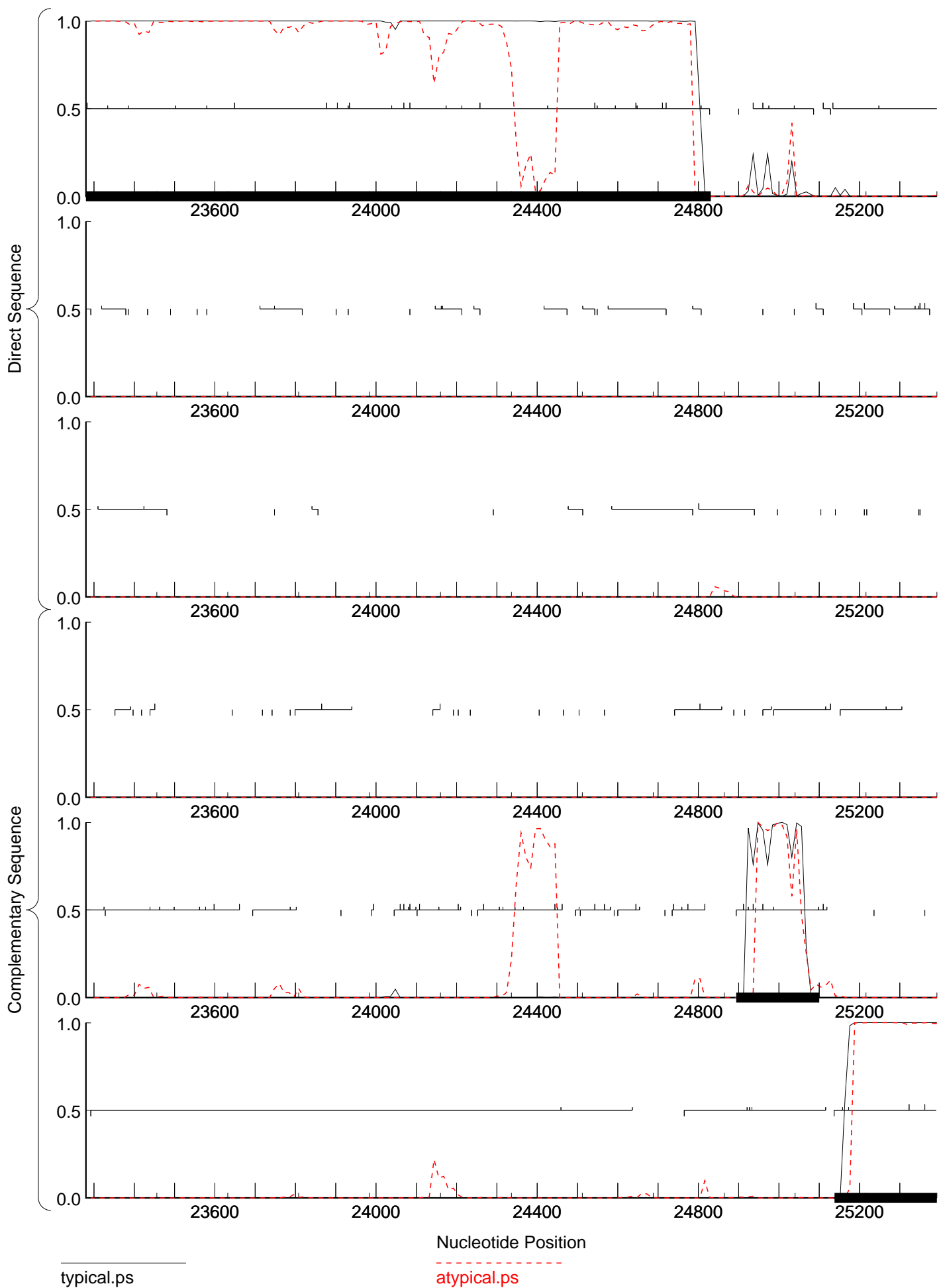
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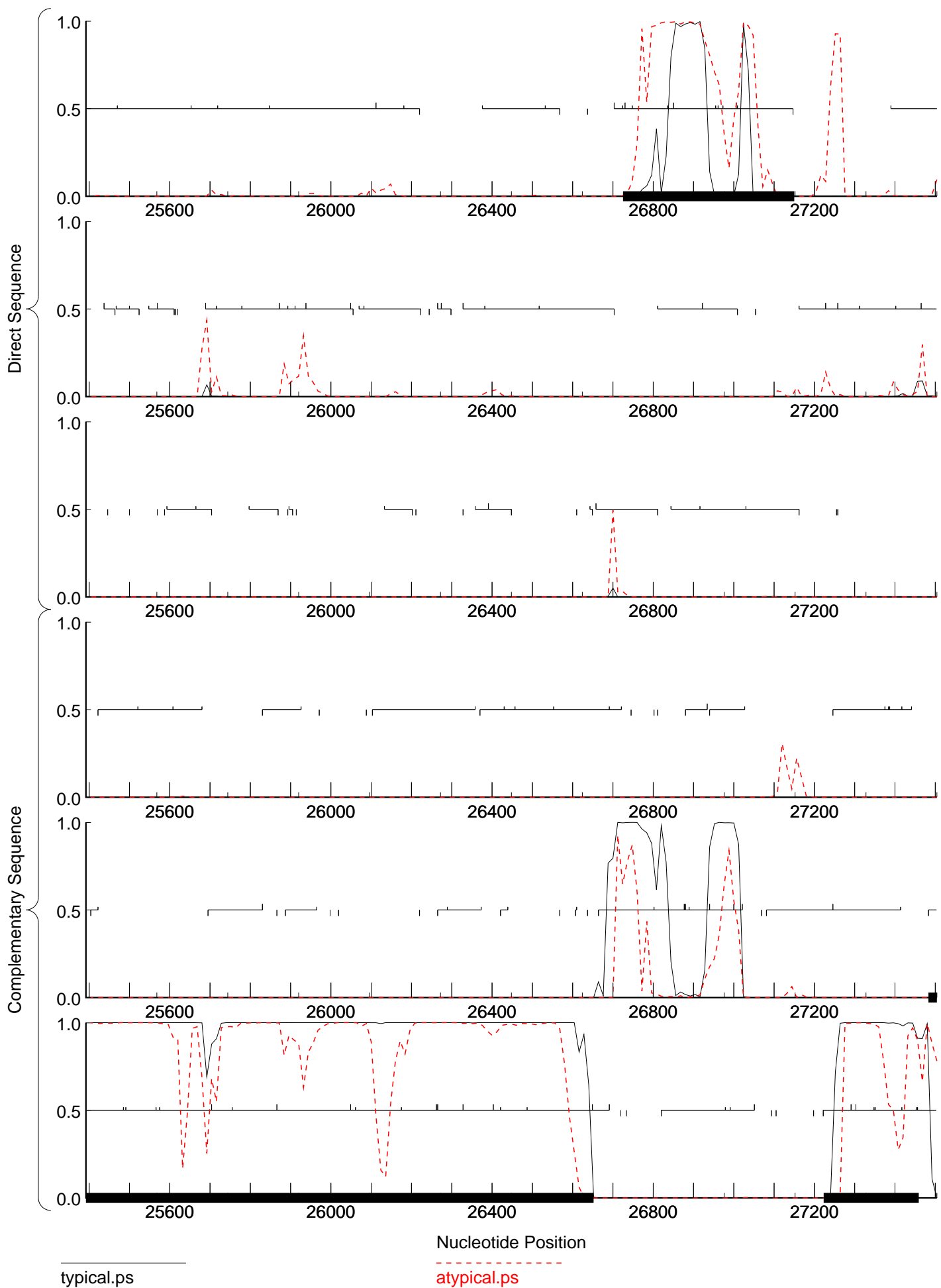
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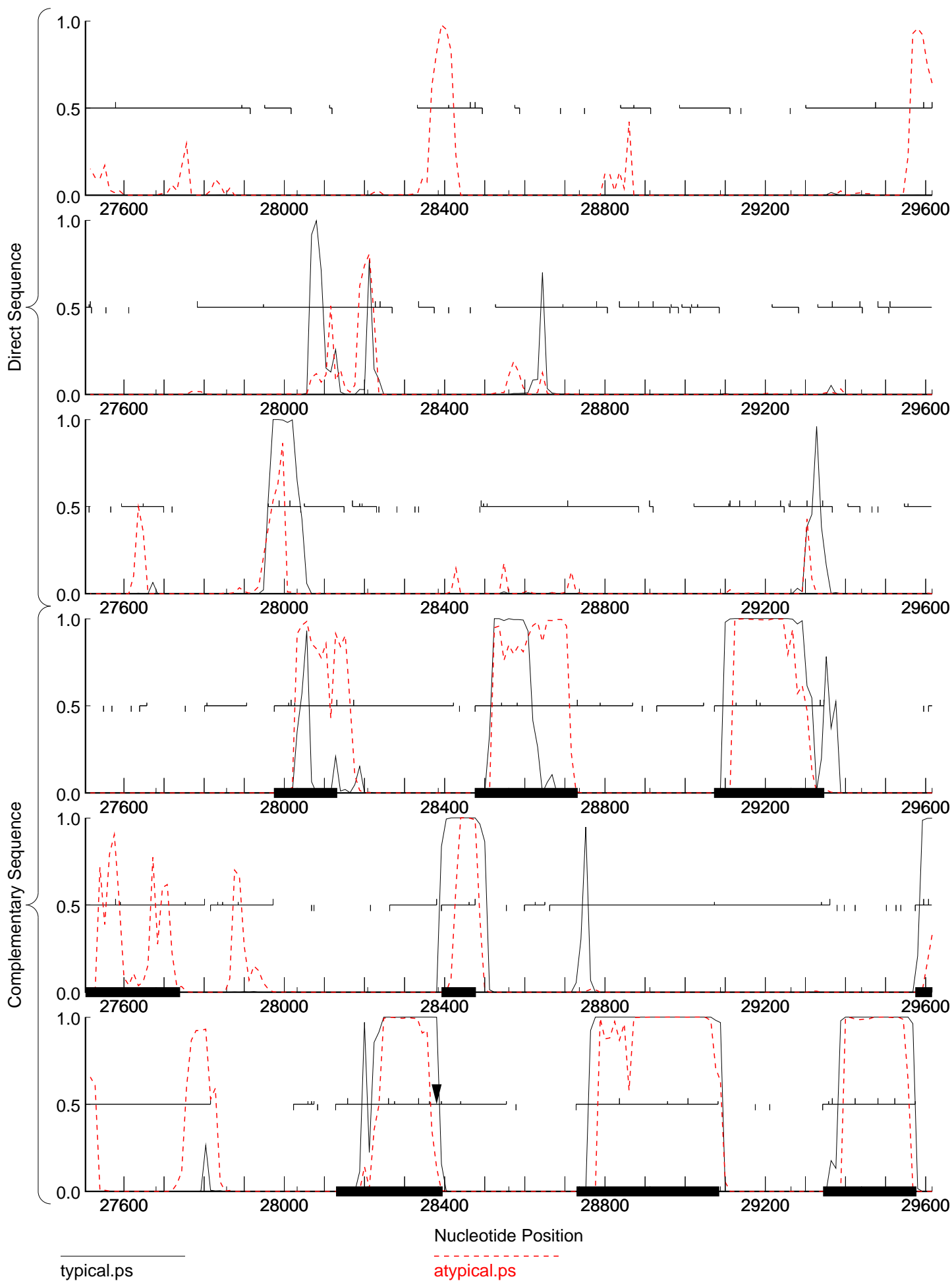


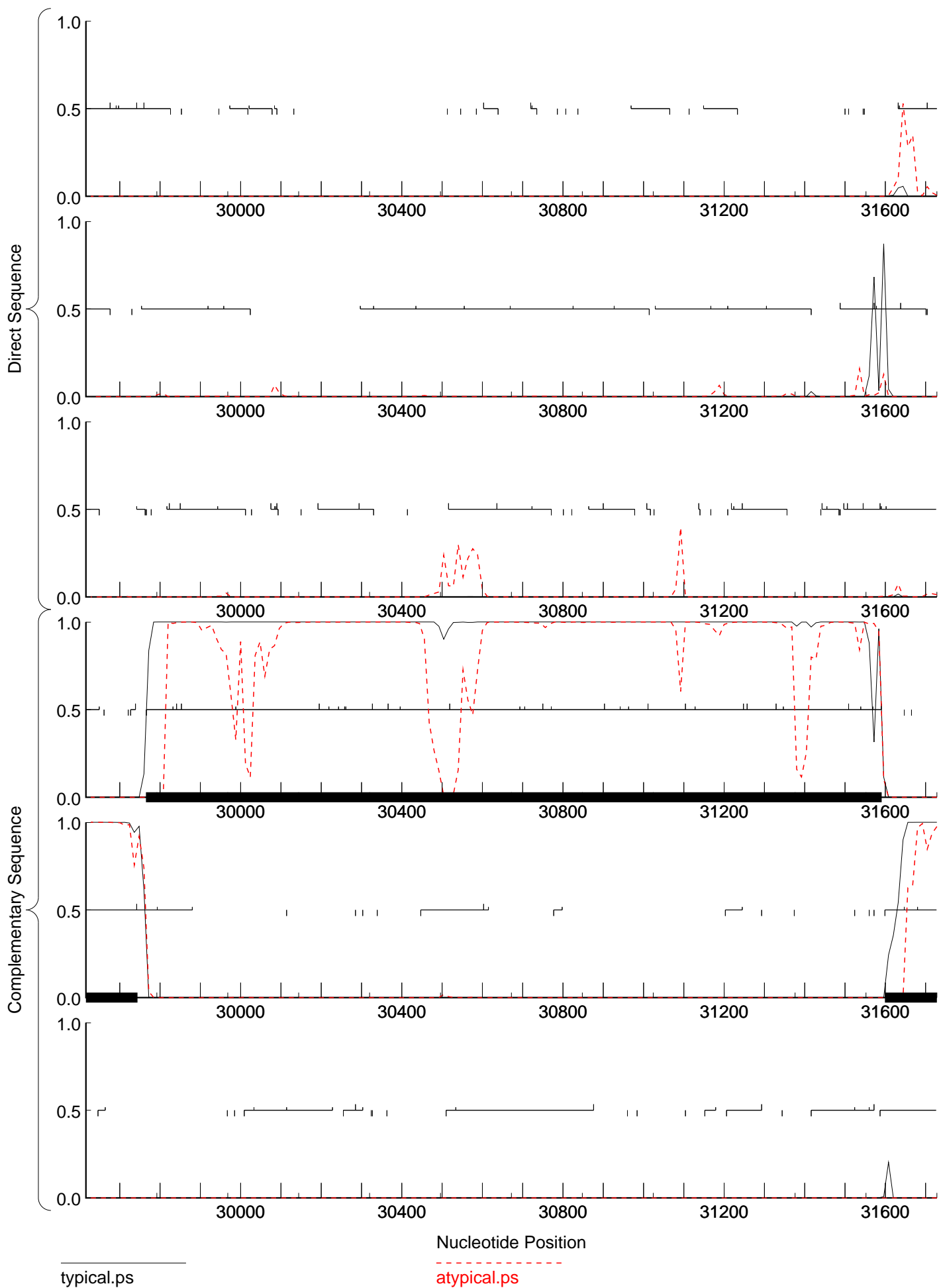
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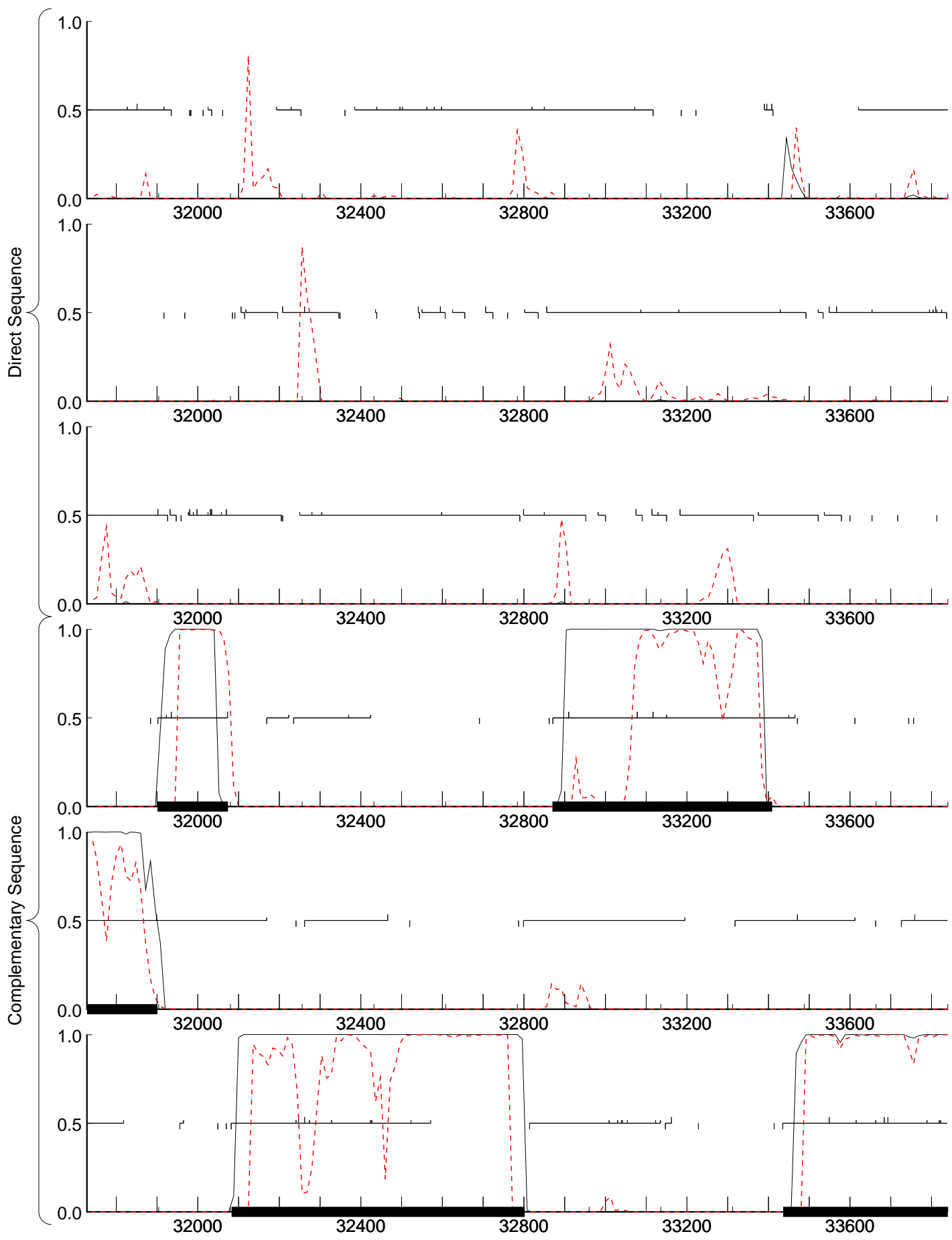
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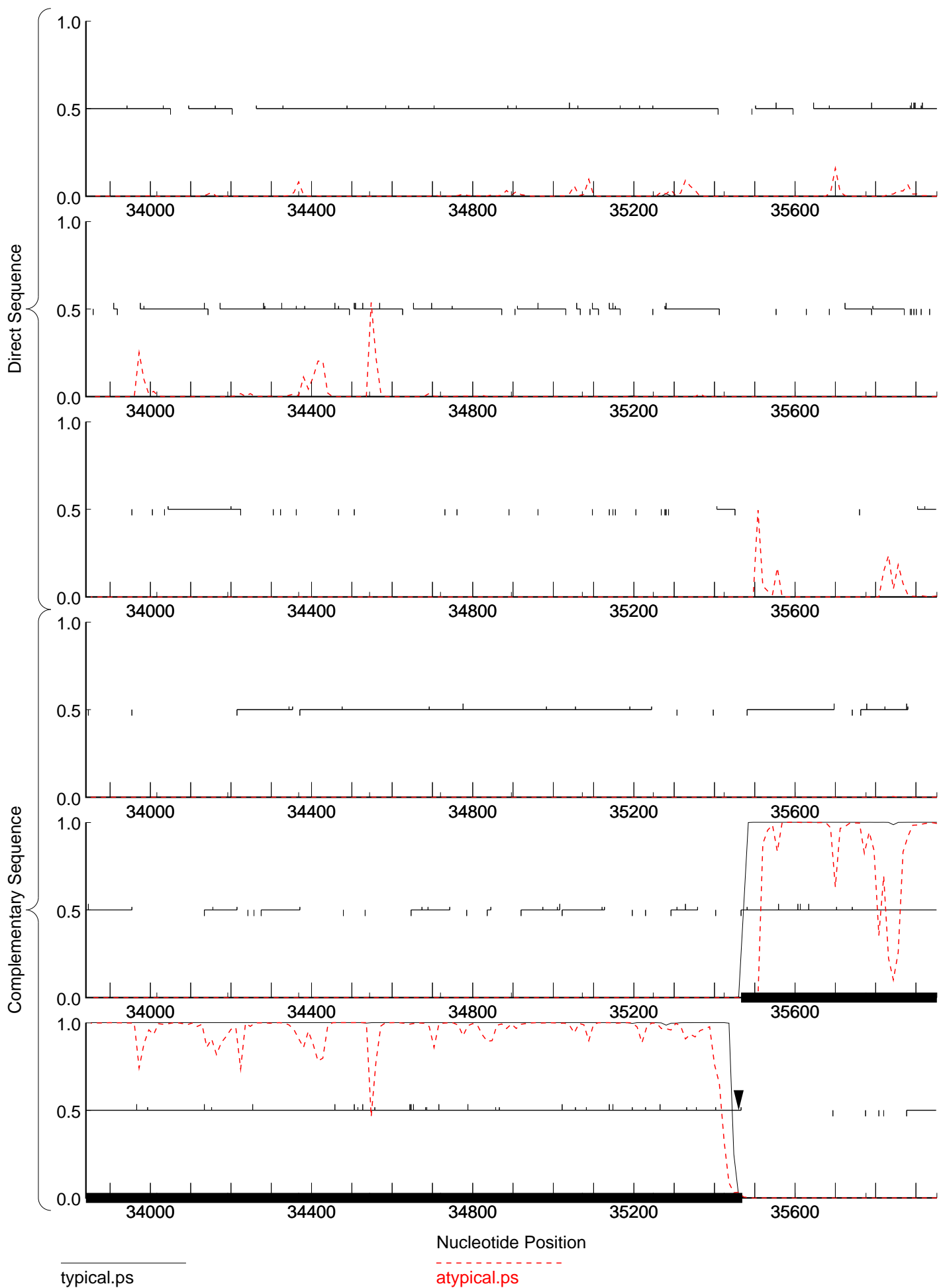






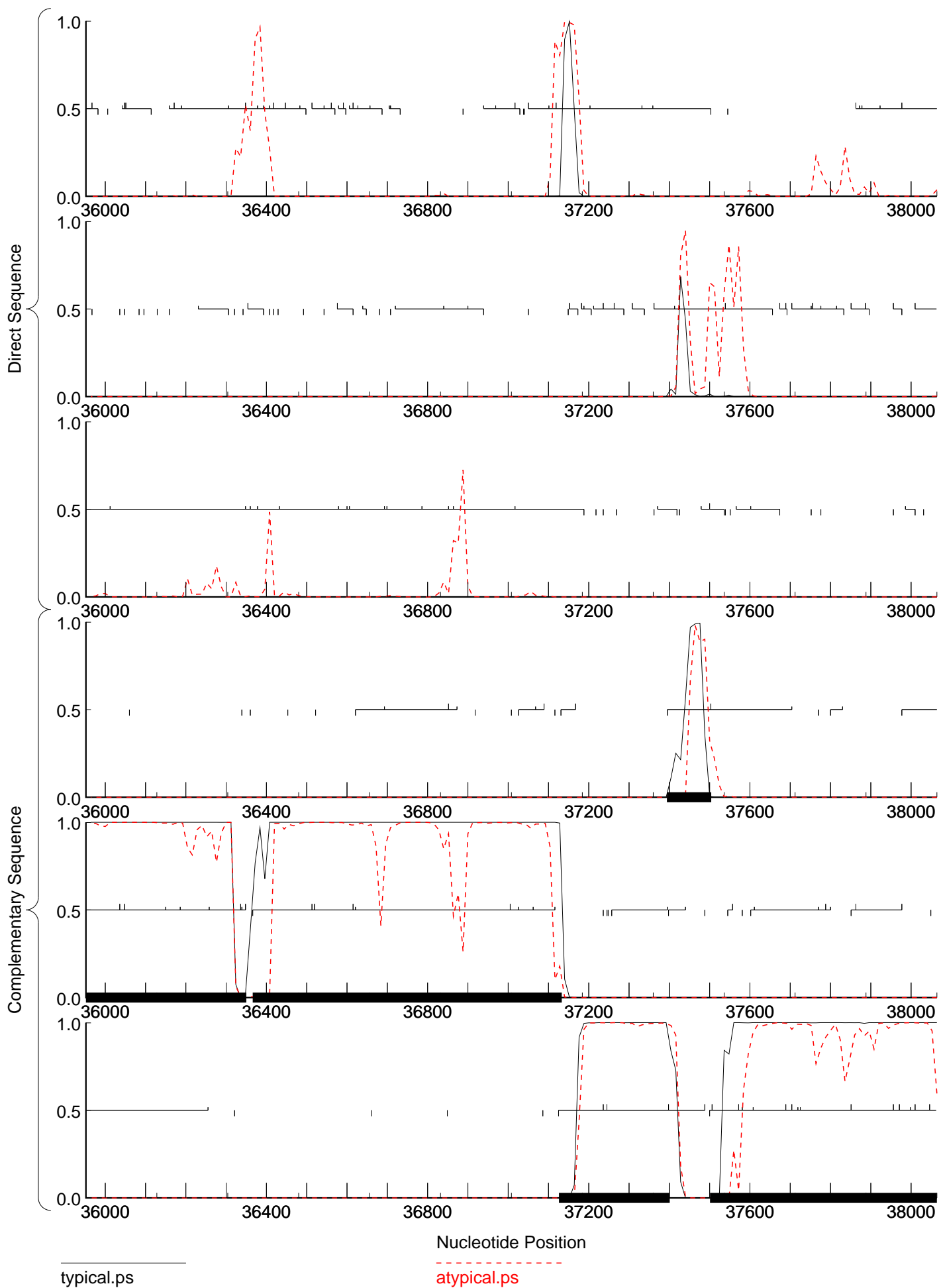
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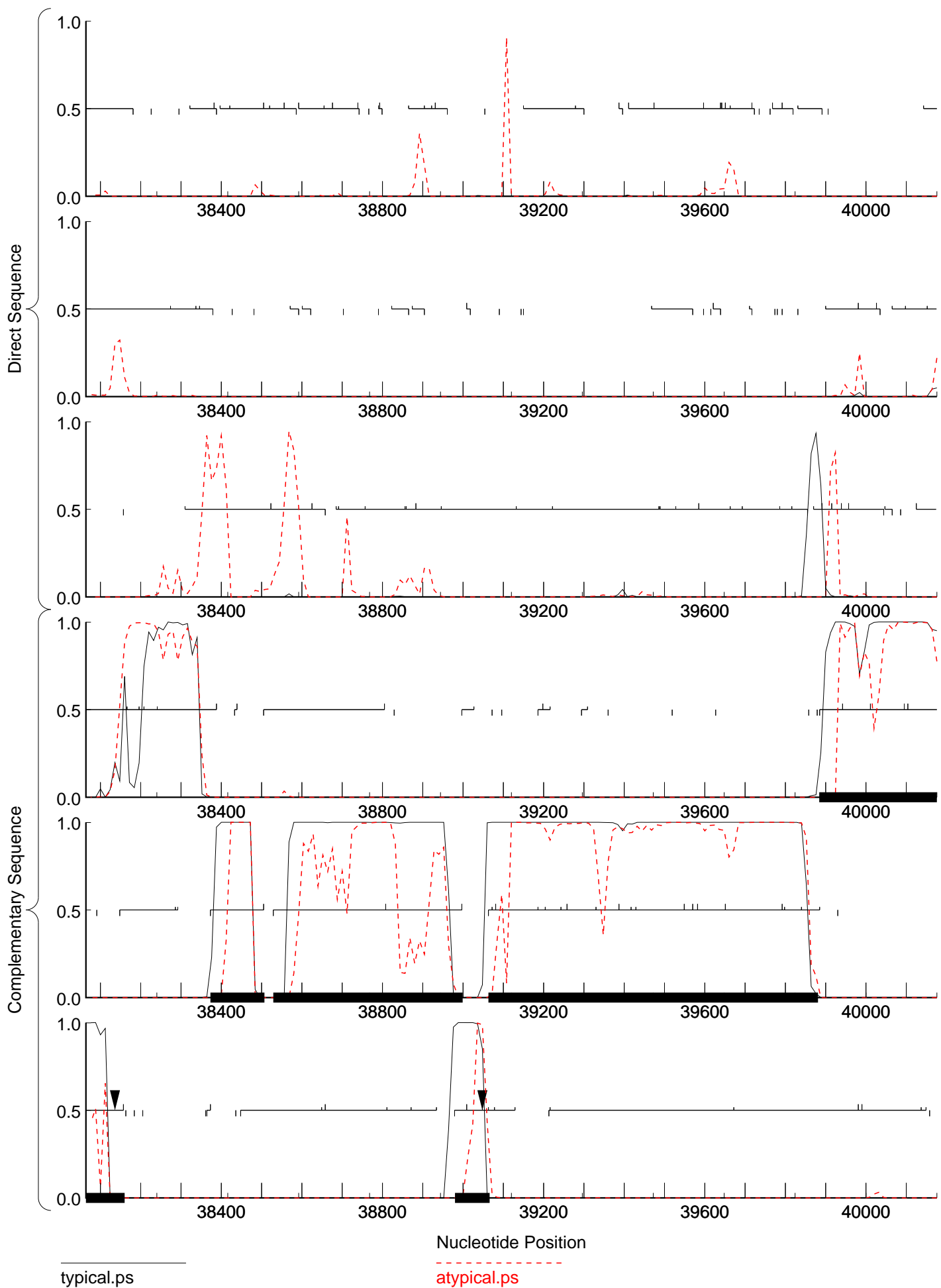
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typical.ps

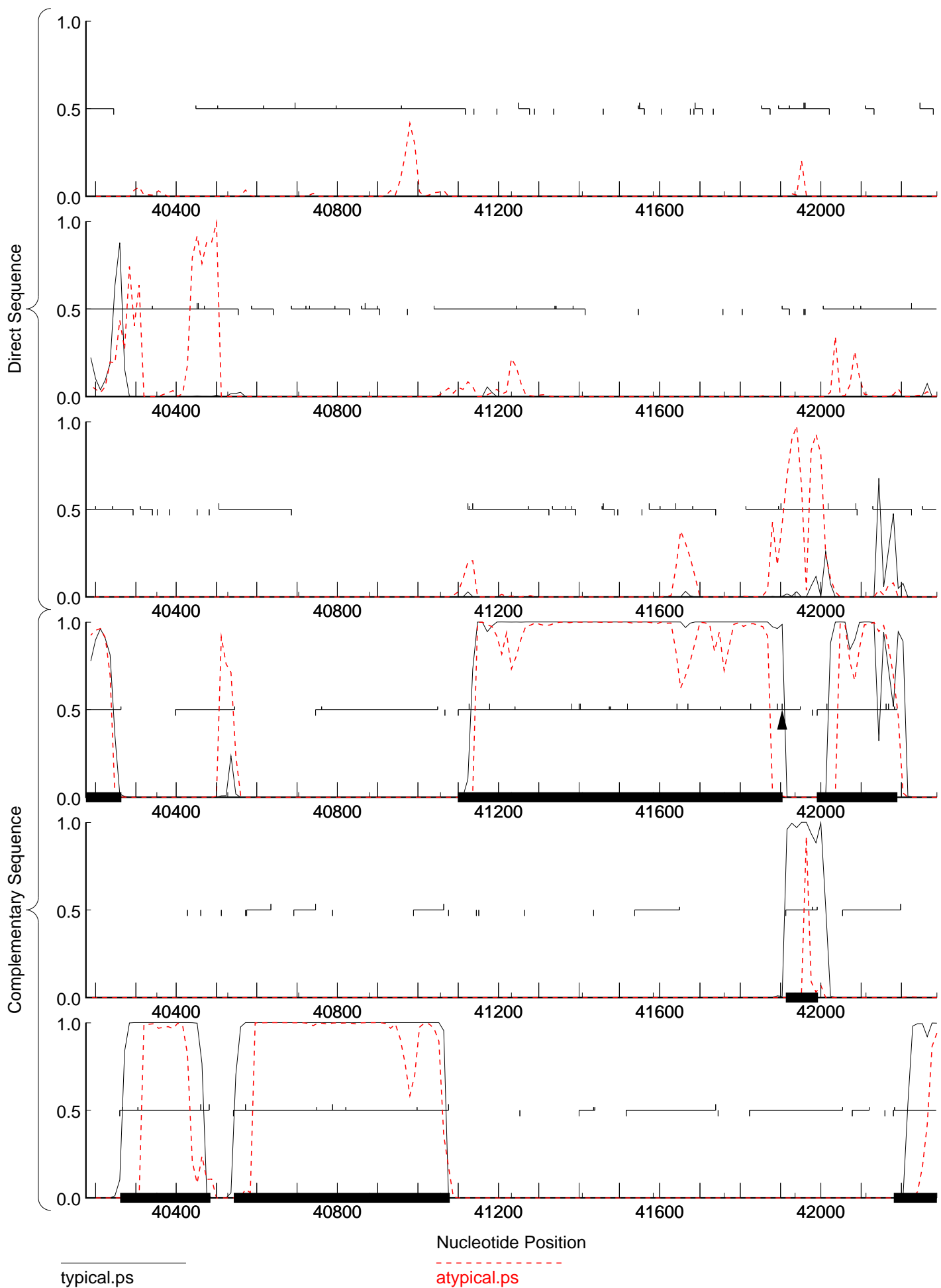
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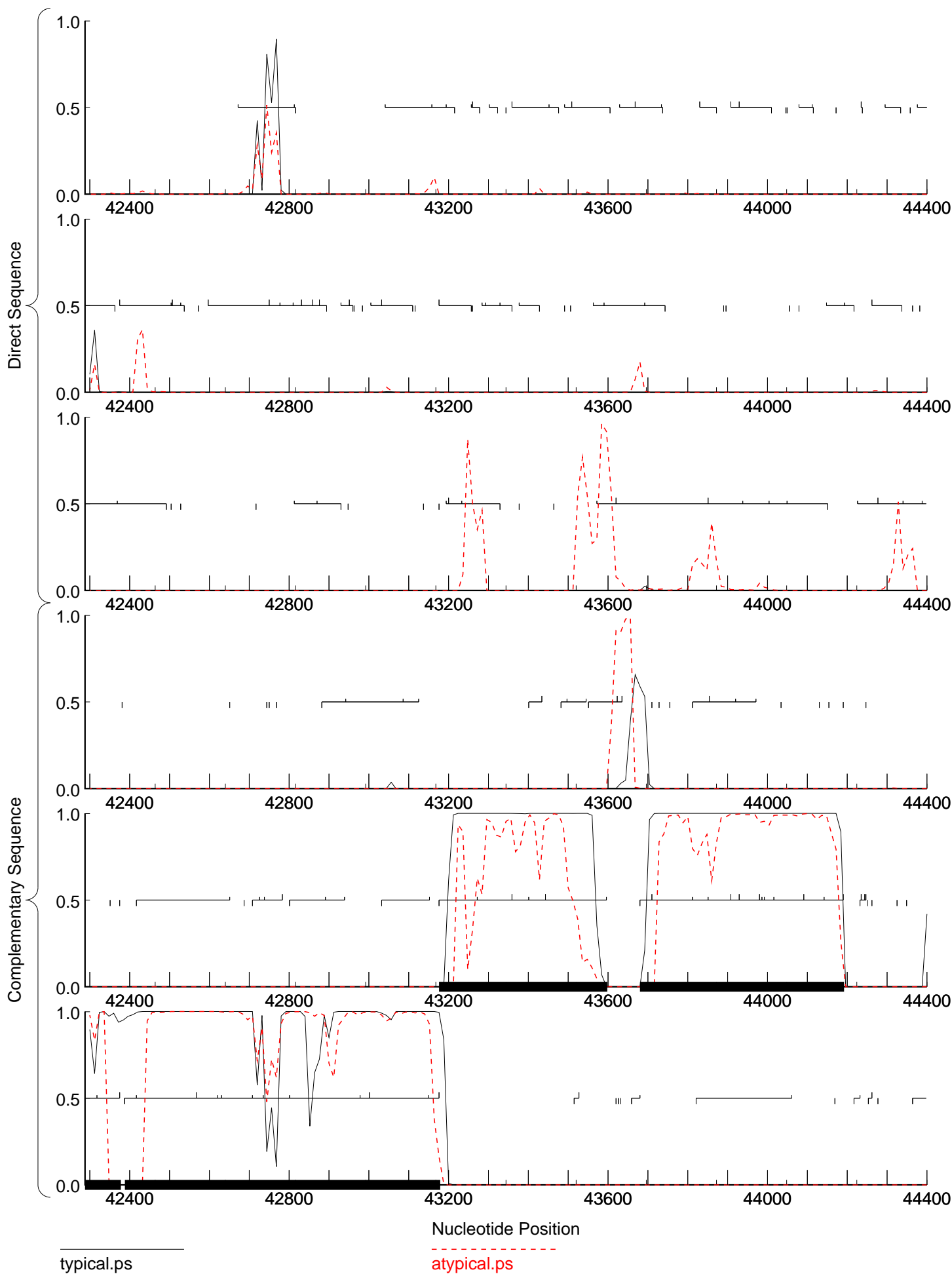
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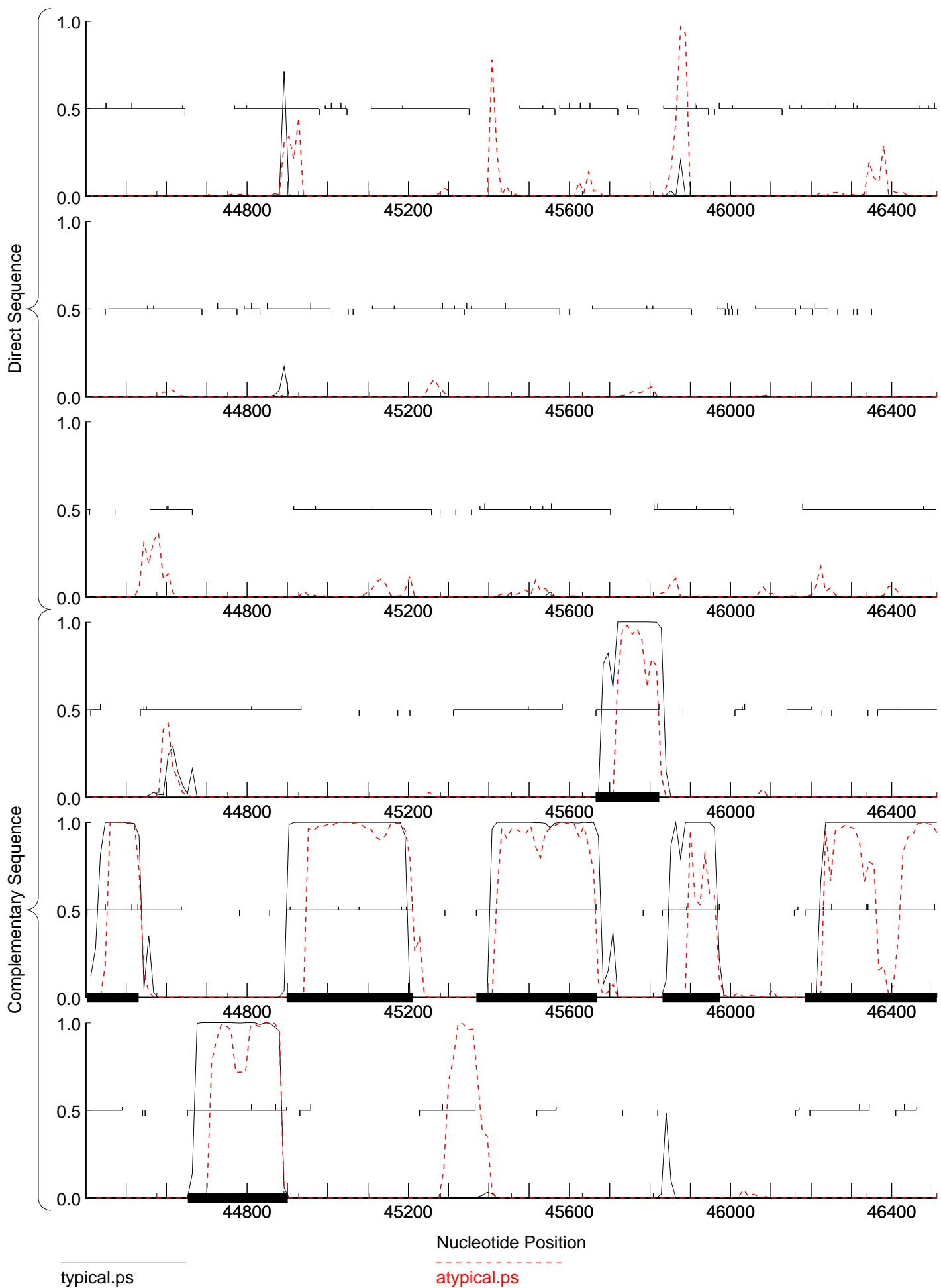
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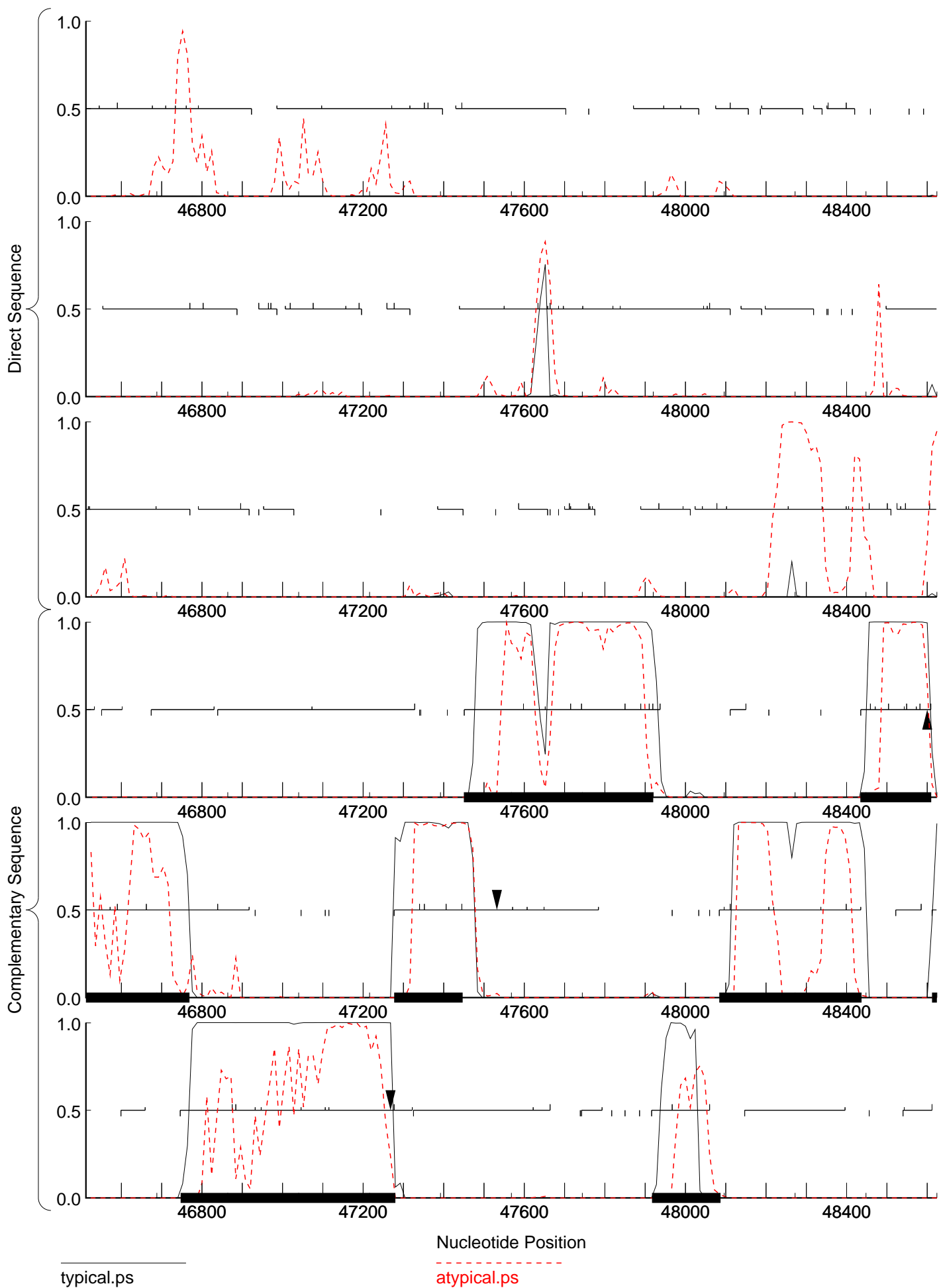


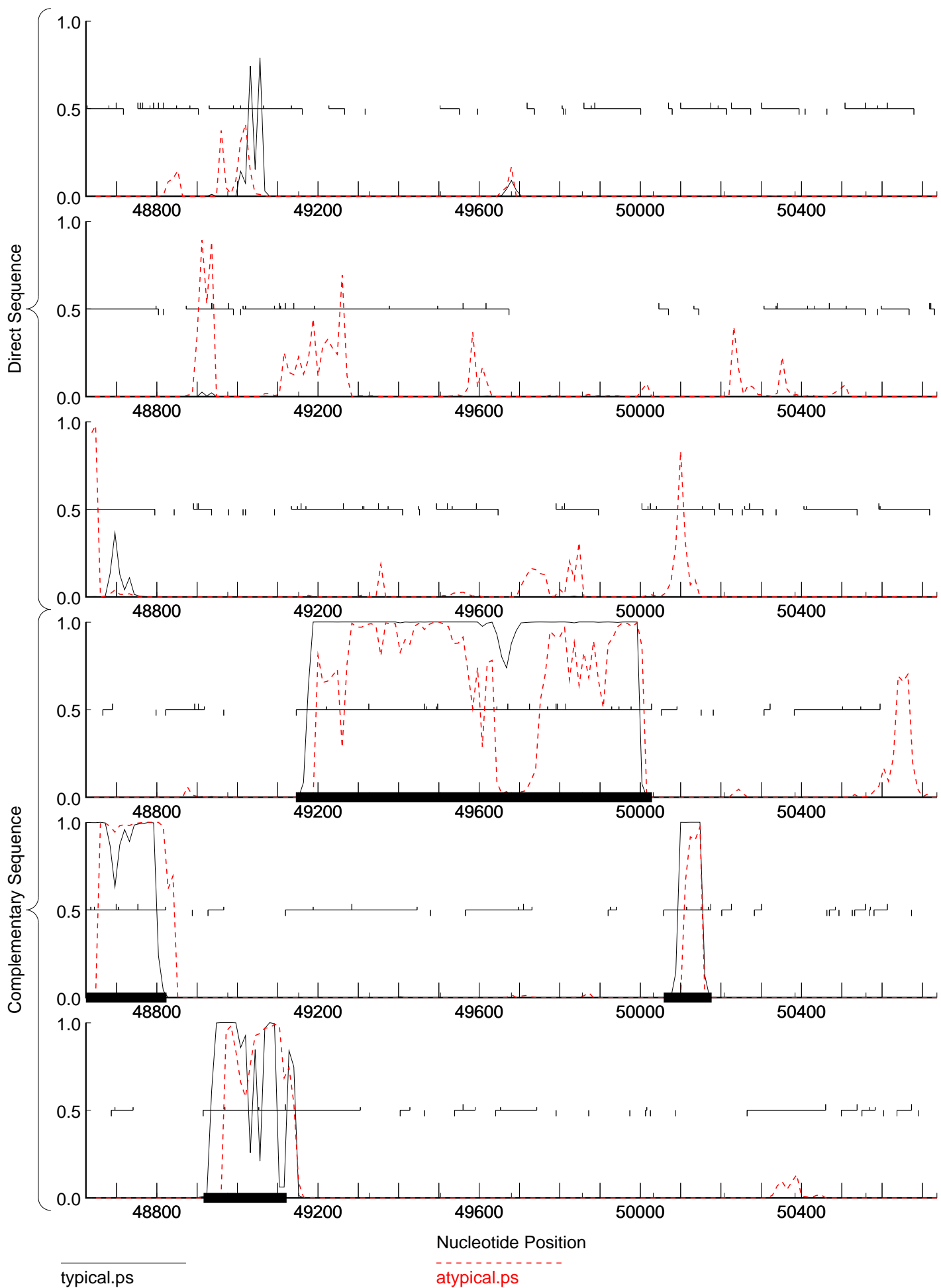
typical.ps

atypical.ps









typical.ps

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