

GeneMark

Version 2.5p (09.08.06)

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PROGRAM INFORMATION

Sequence : Mycobacterium phage Thumb complete sequence, 68408 bp, circularly permuted, Cluster H1
Analysis Date : 8/21/17 at 12:05:30
Pages : 34
Sequence Length : 68408 bp
GC Content : 57.50%

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,

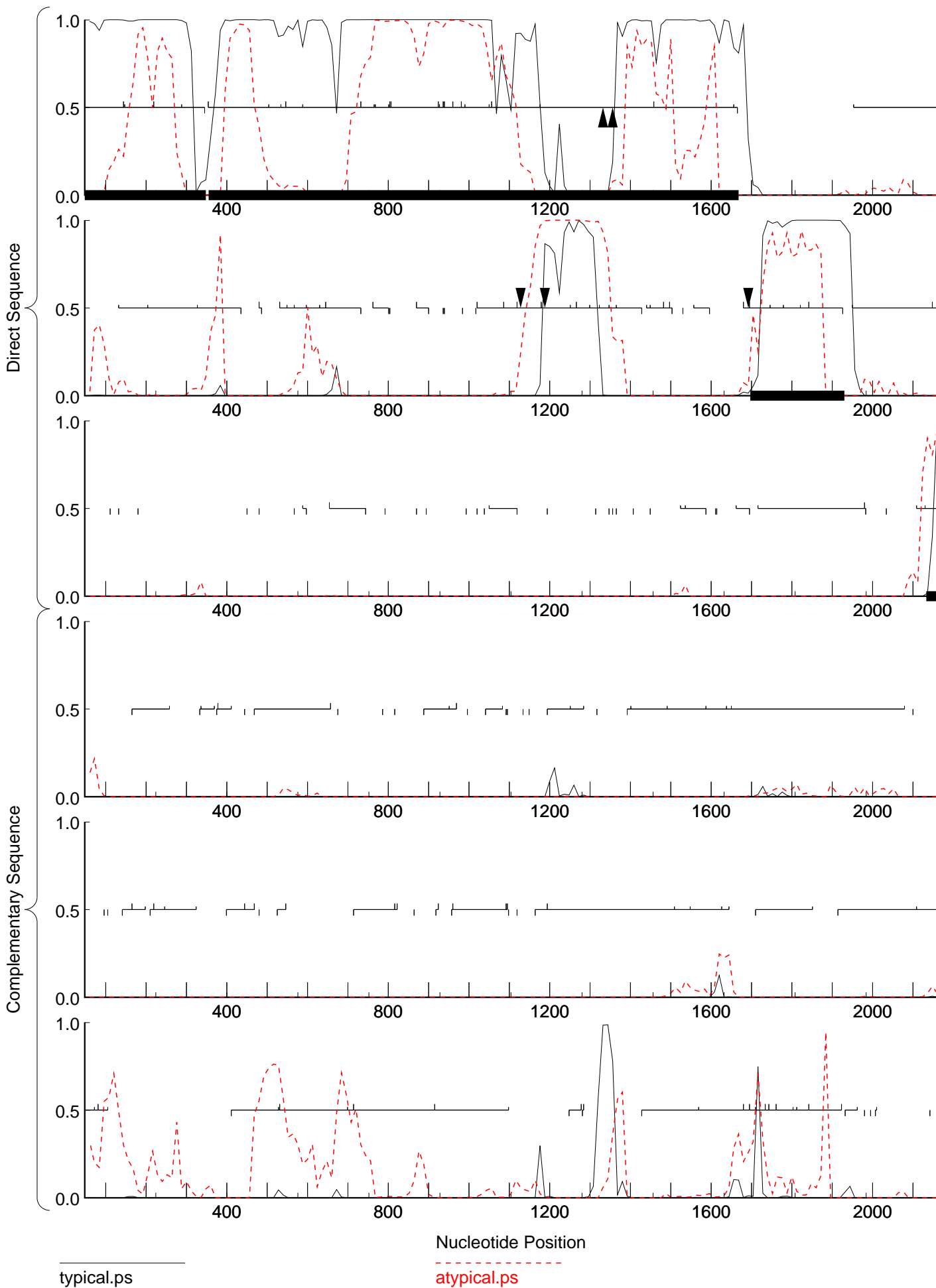
Matrix notes & comments

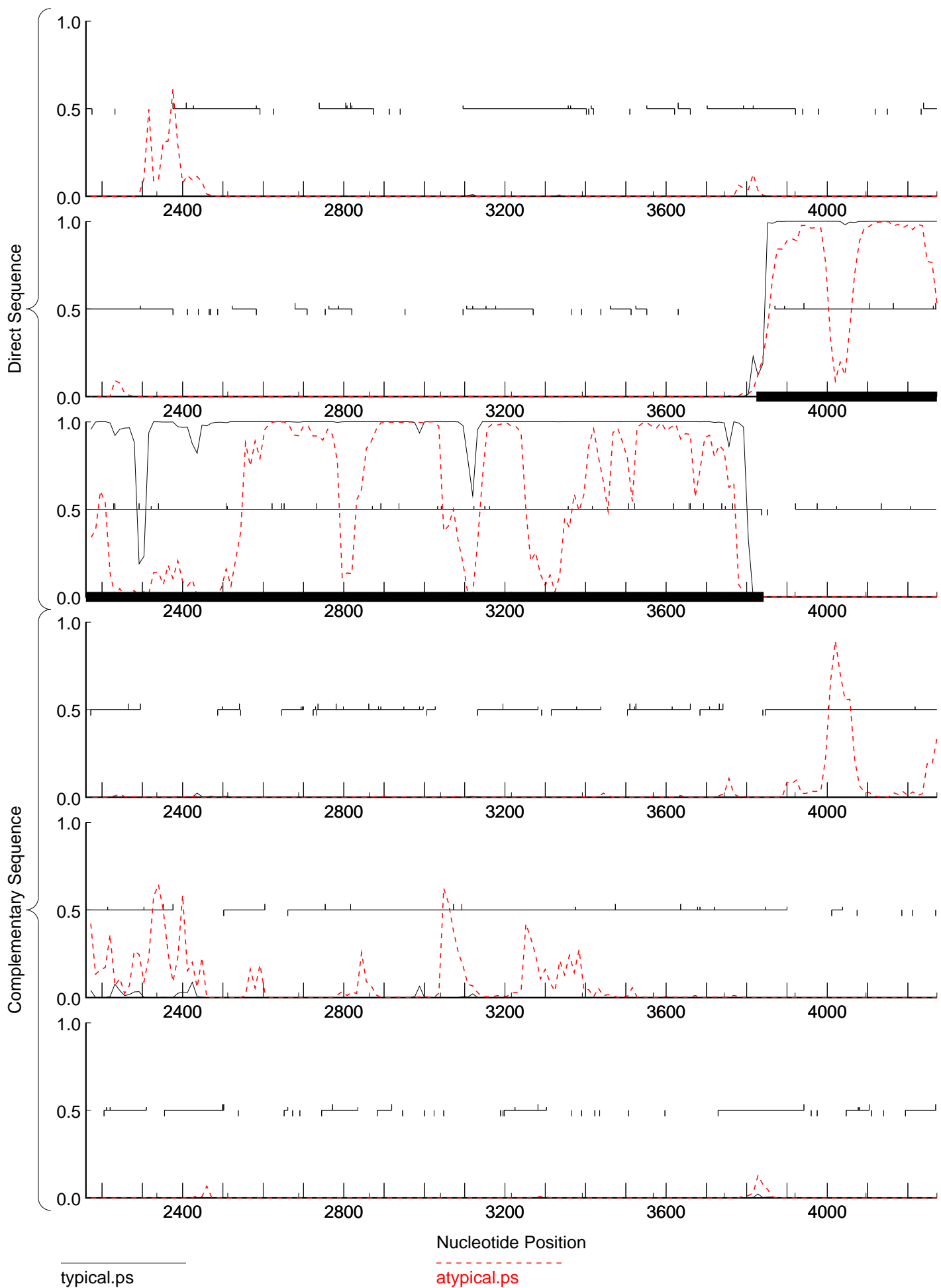
Training set derived by GeneMarkS, 4.30 October 2014
Mon Aug 21 12:05:30 2017

MATRIX INFORMATION

Matrix : -
Author : -
Order : 4

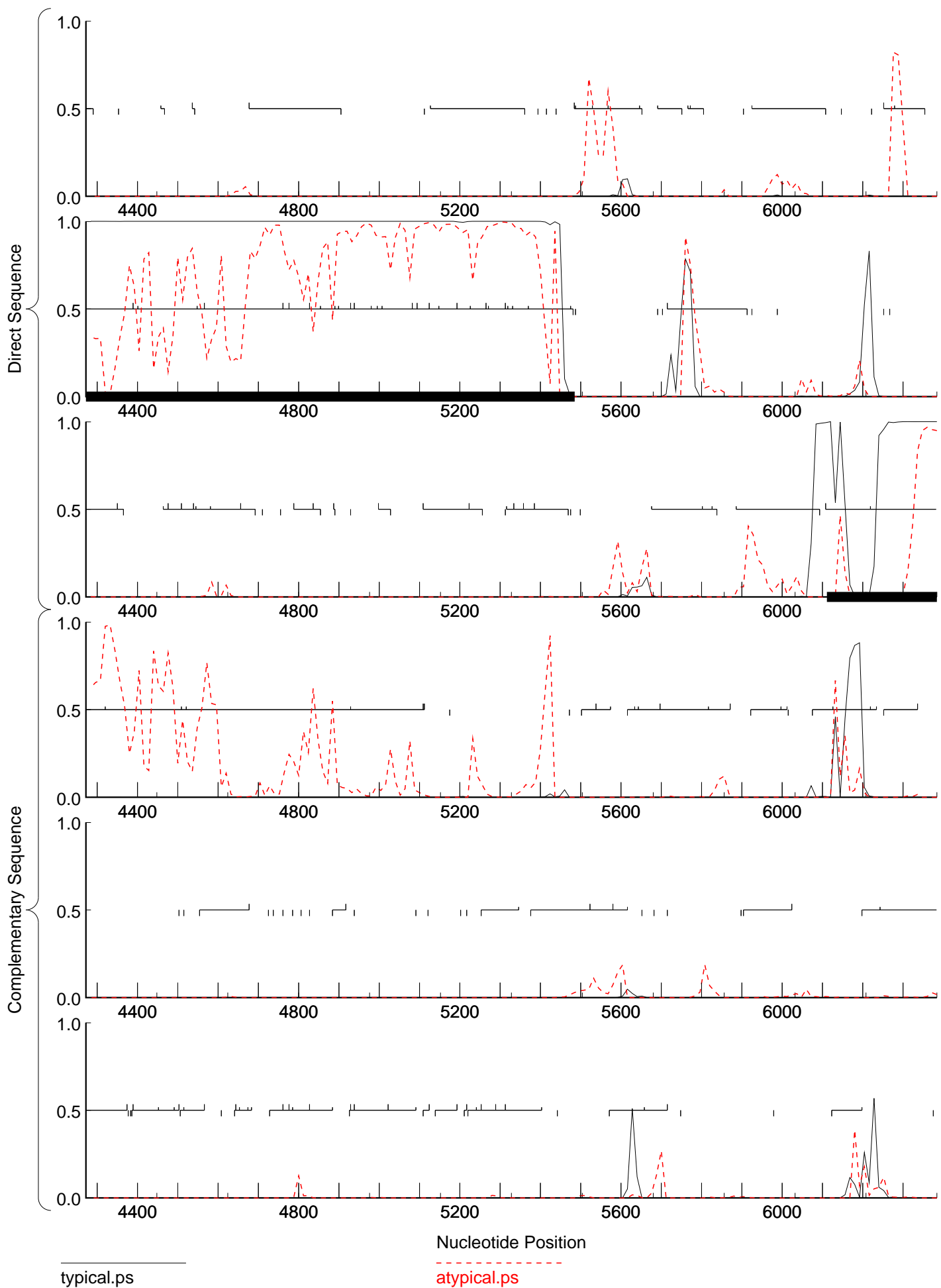
Send questions / comments to:
Dr. M. Borodovsky
Georgia Institute of Technology
School of Biology
Atlanta, GA 30332-0230

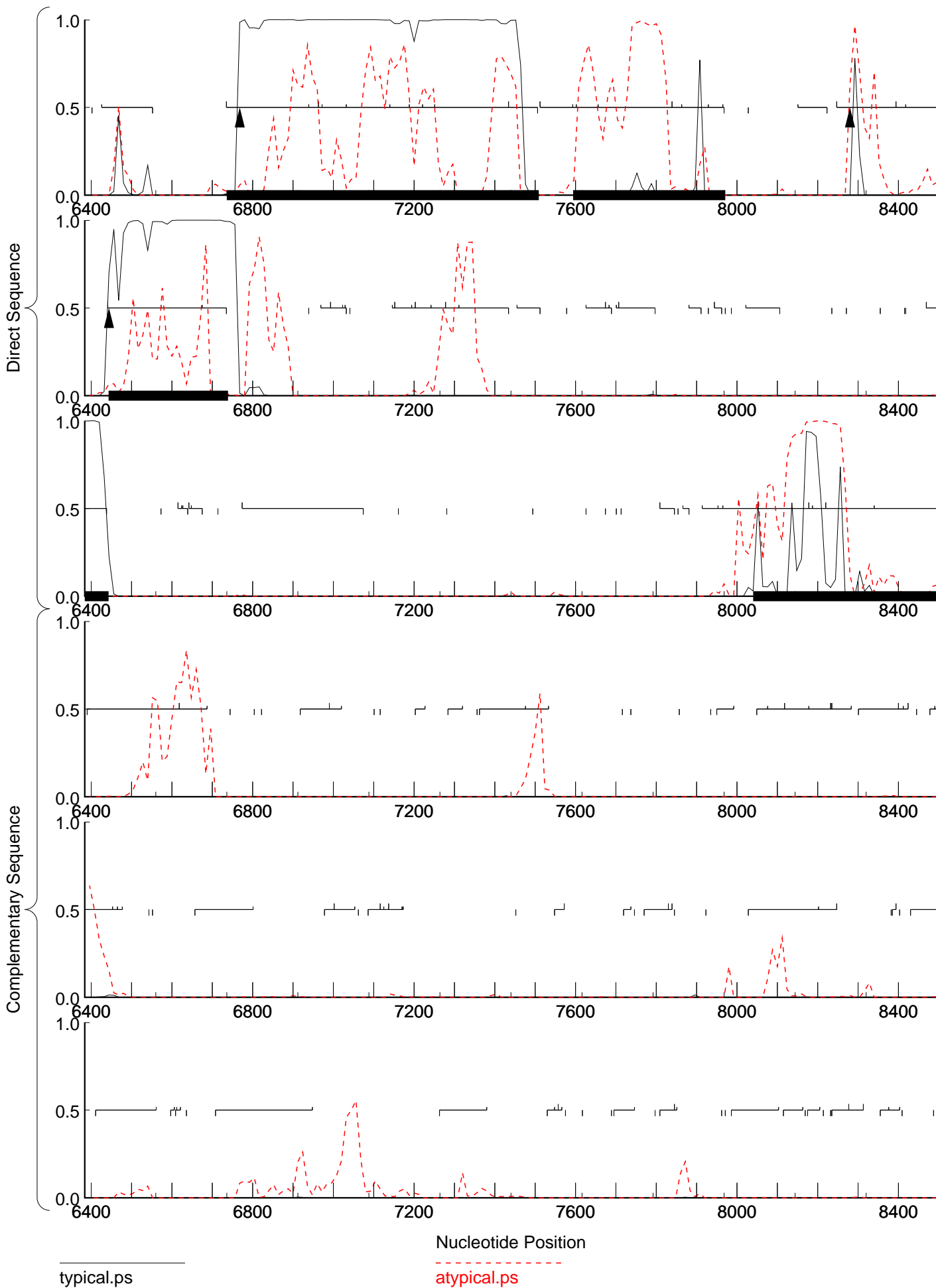


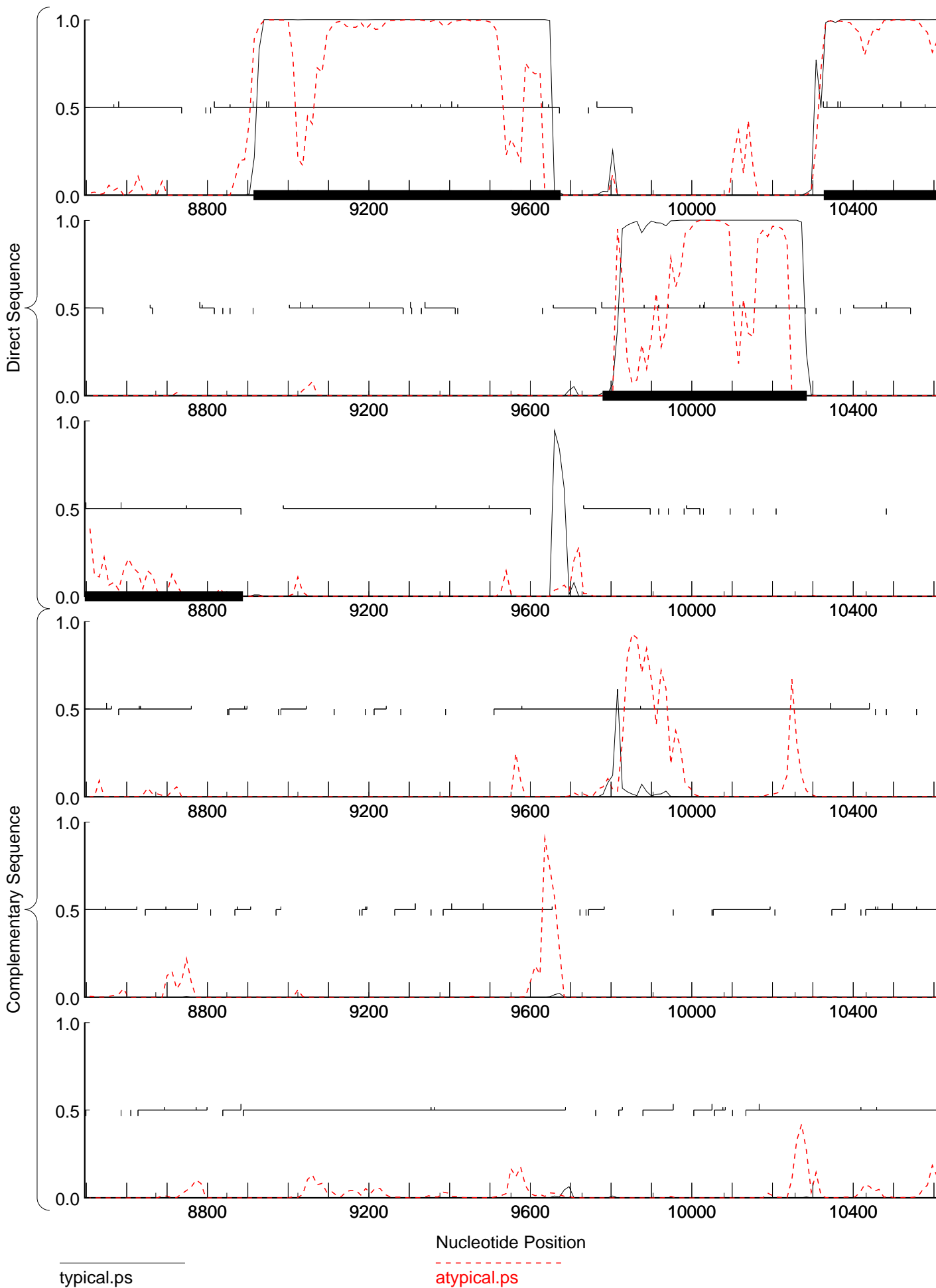


typical.ps

atypical.ps

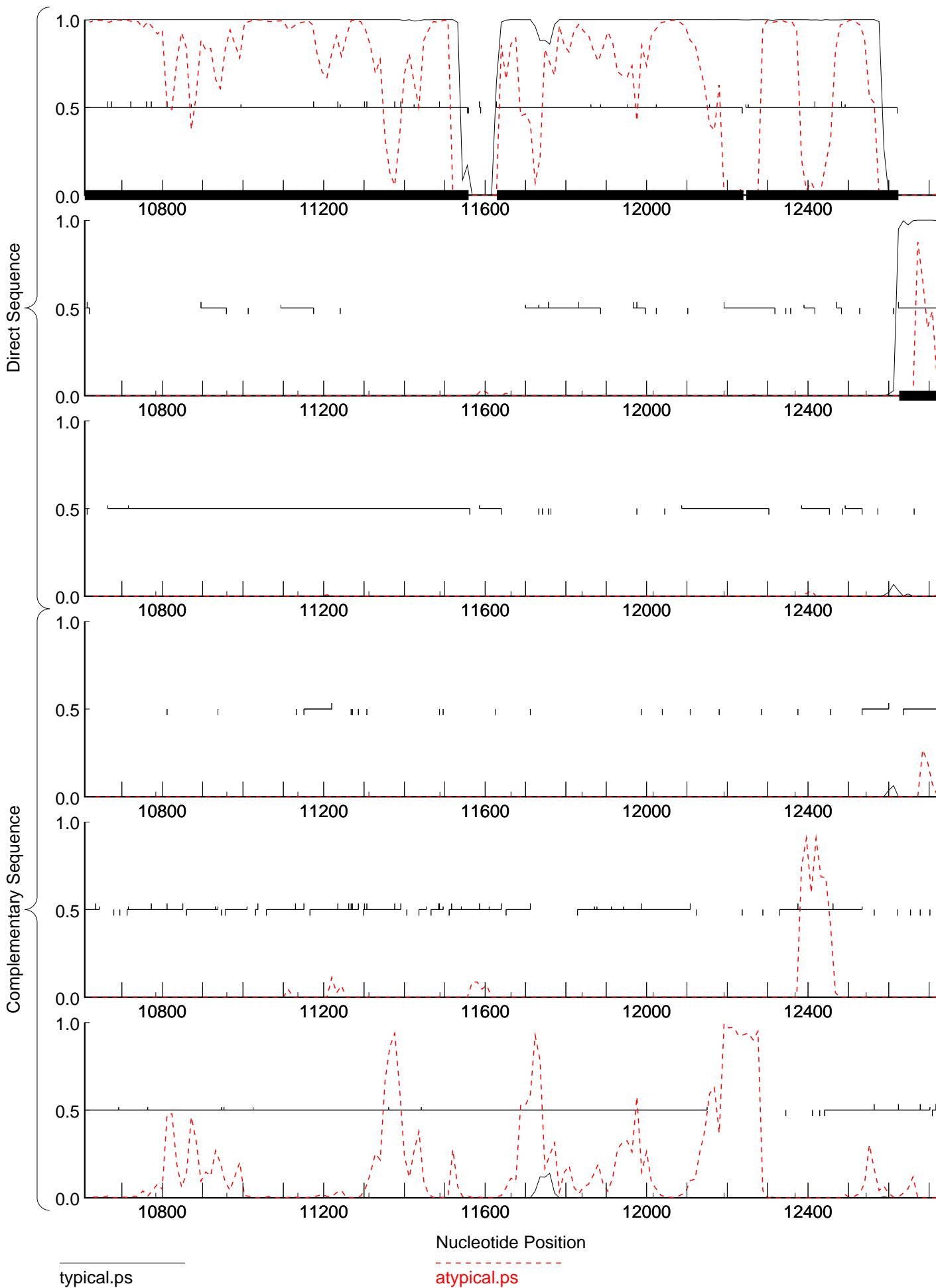


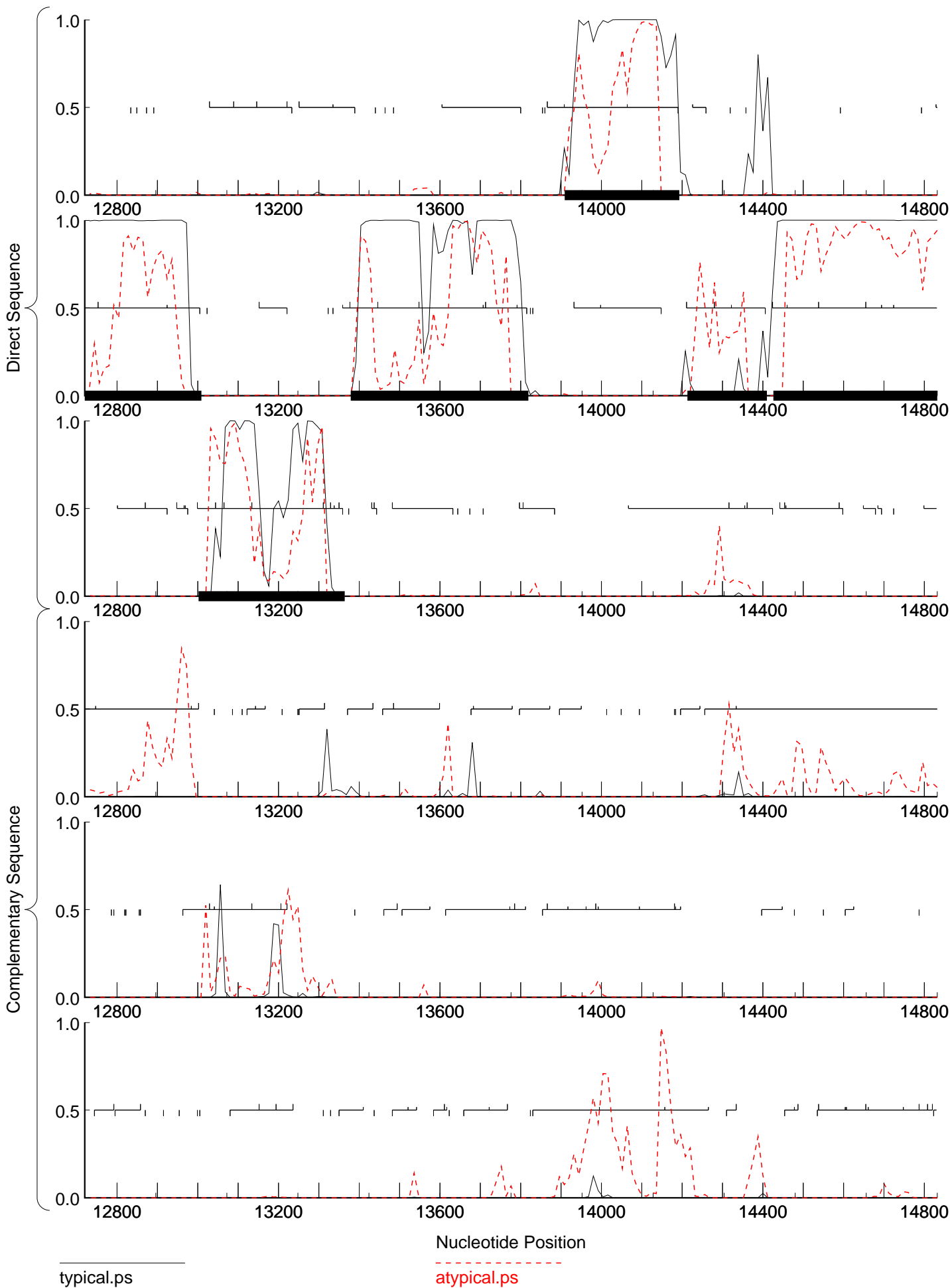


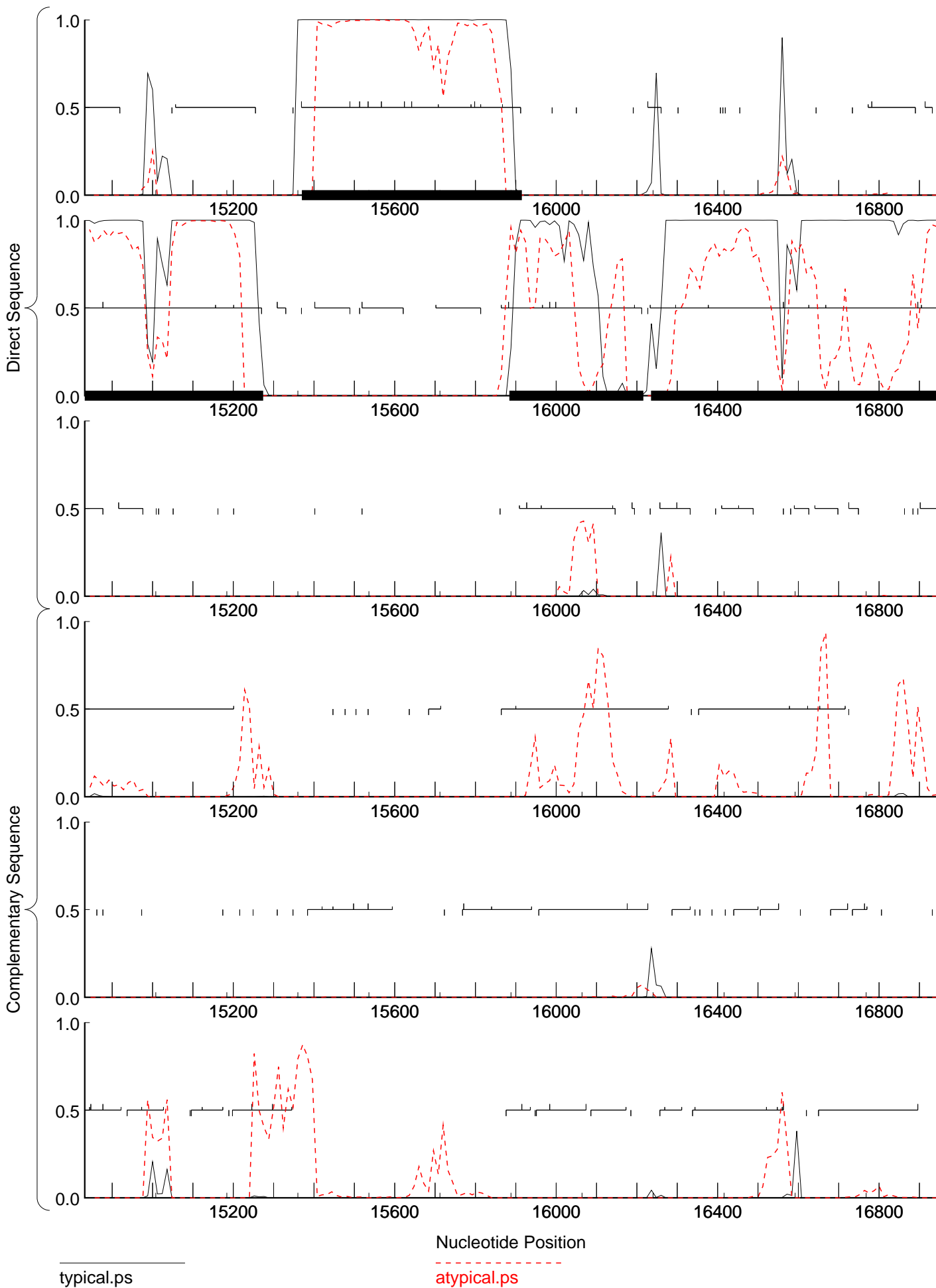


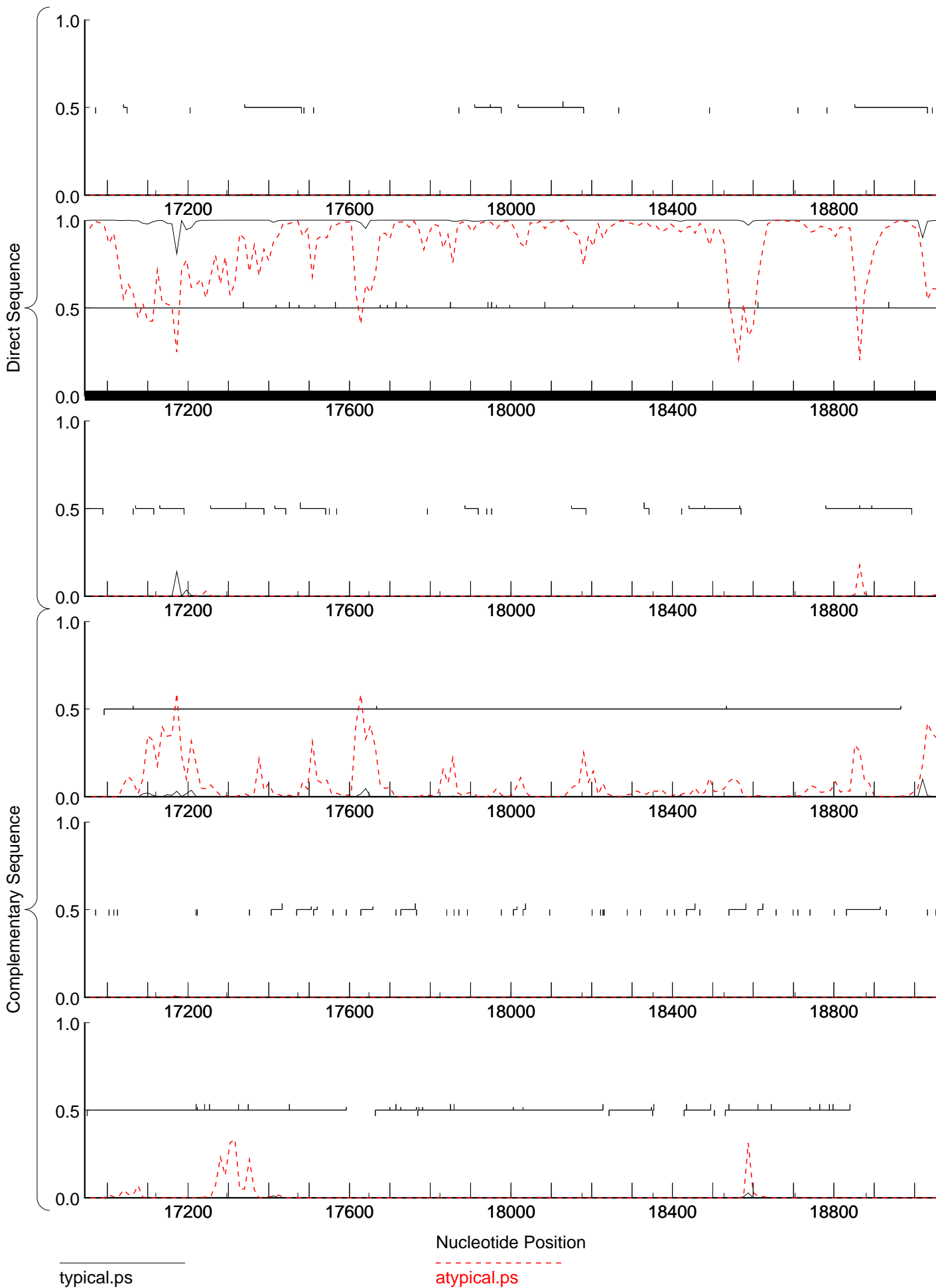
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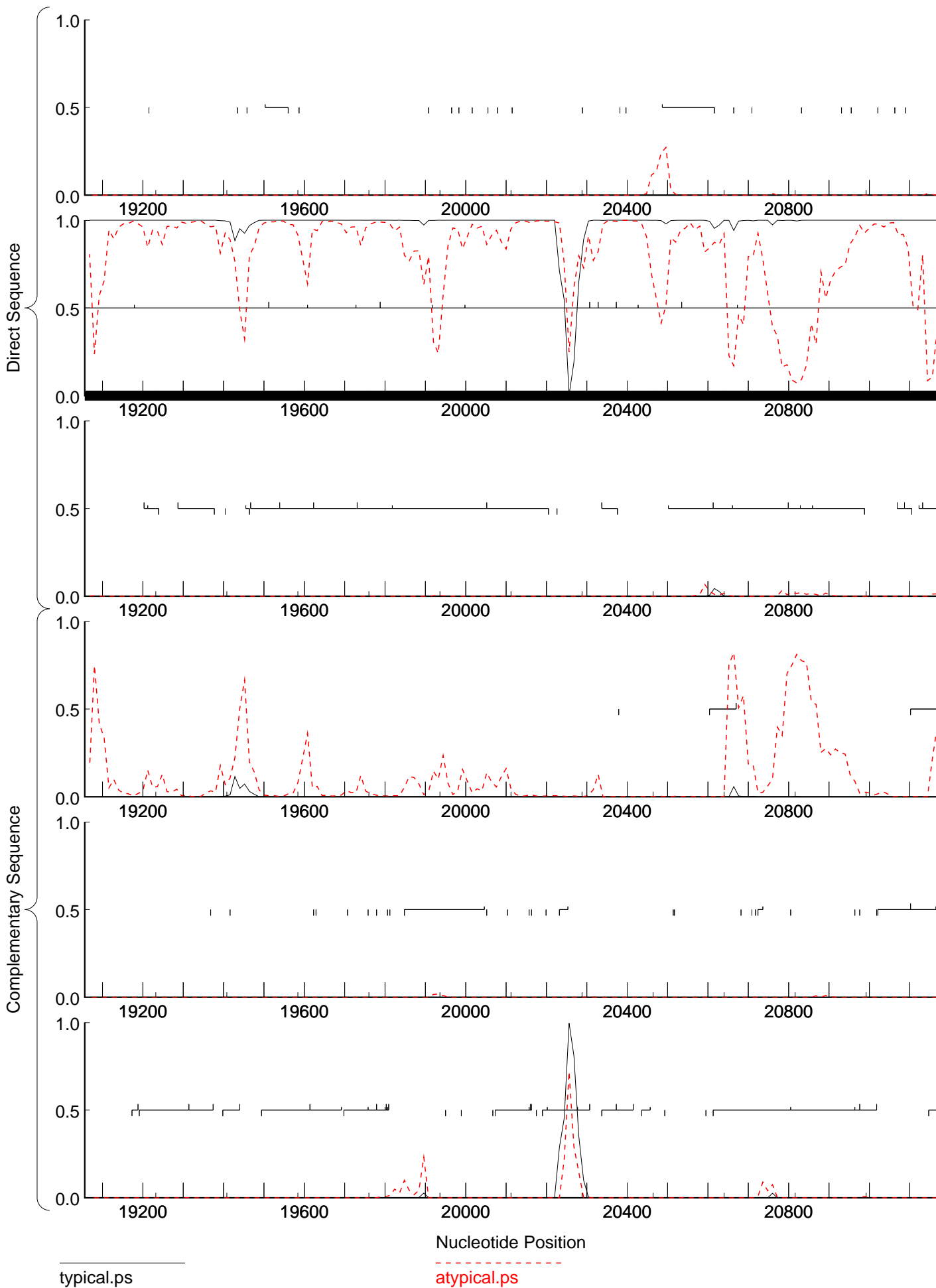
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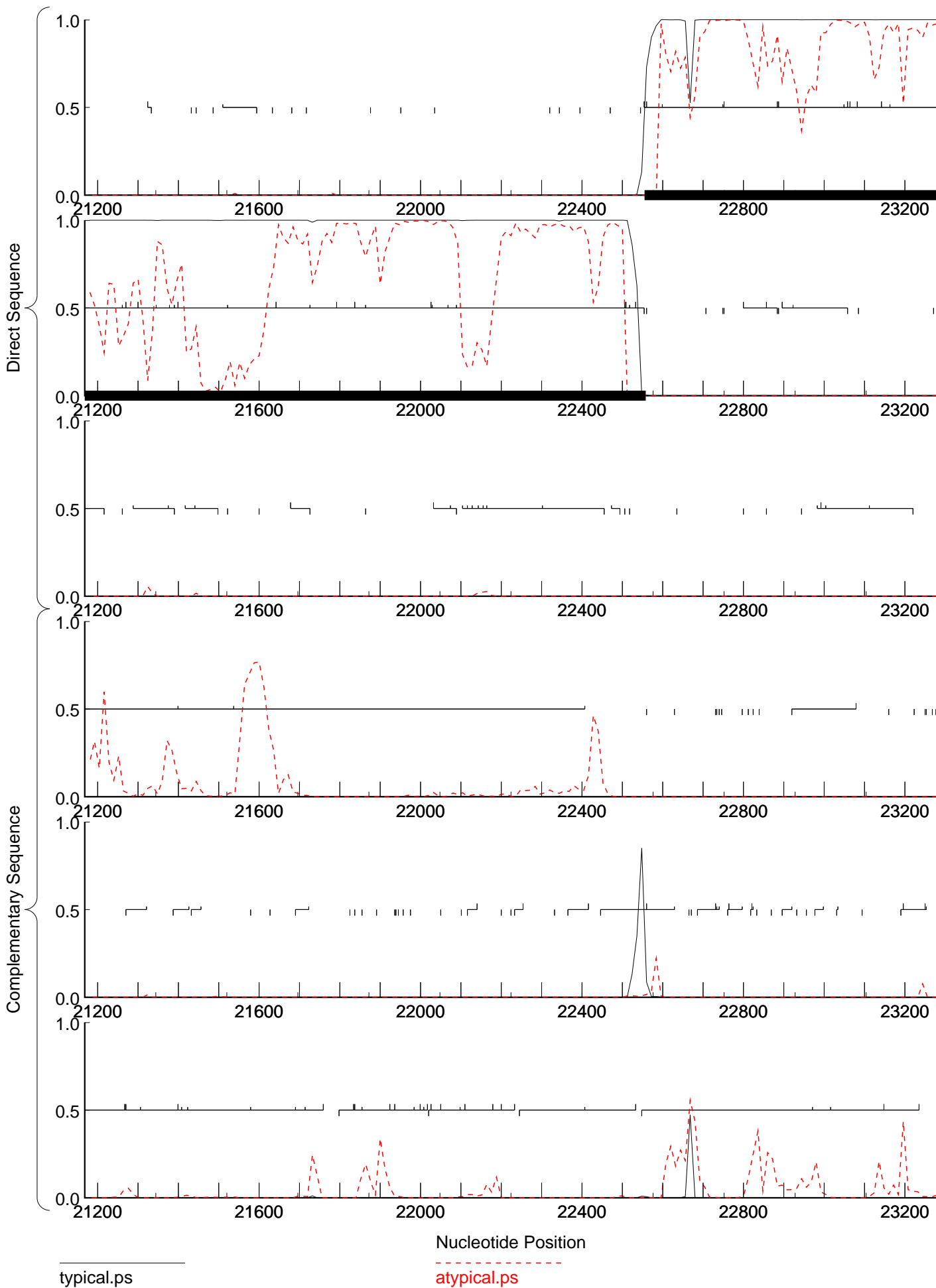


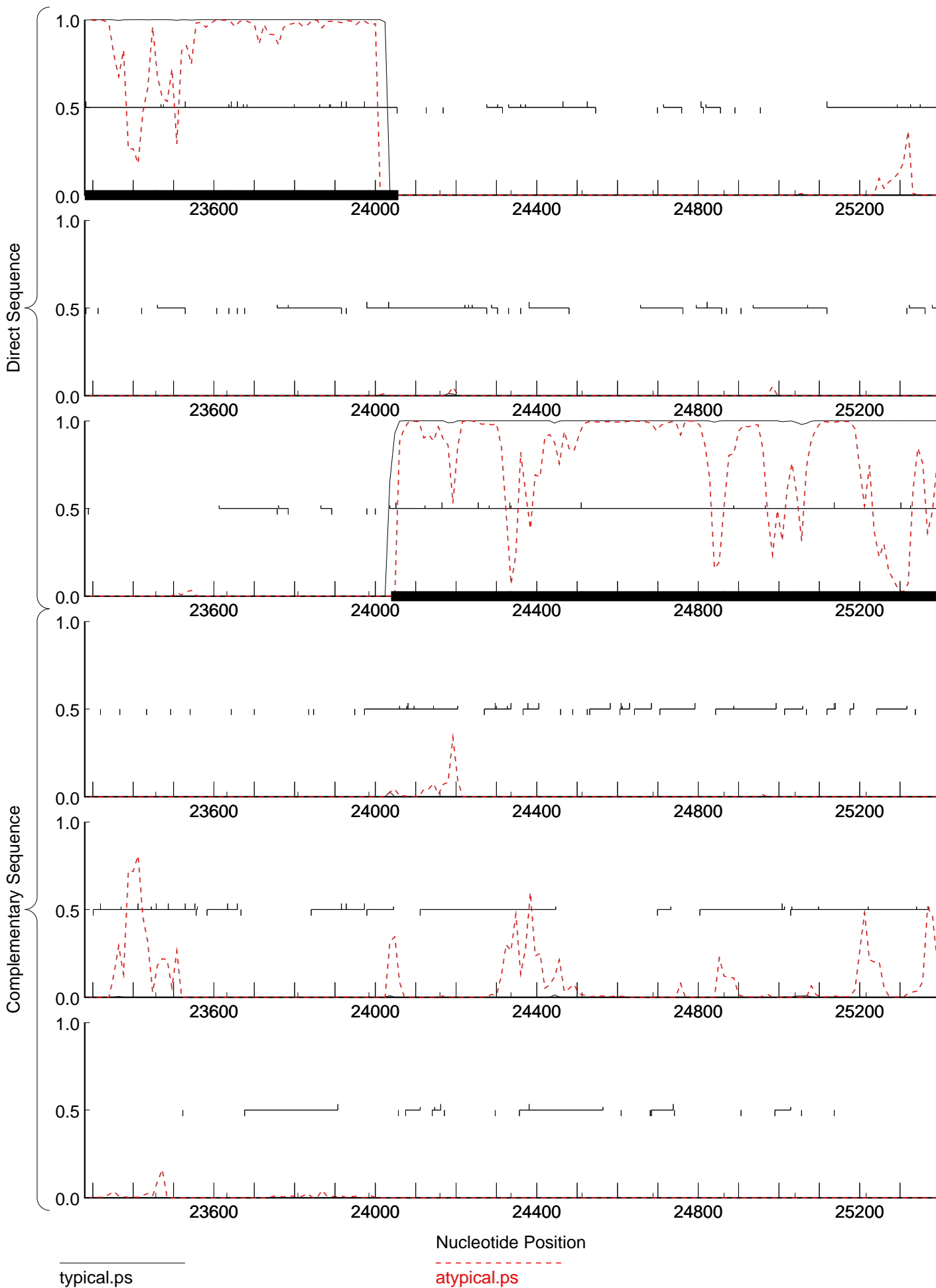






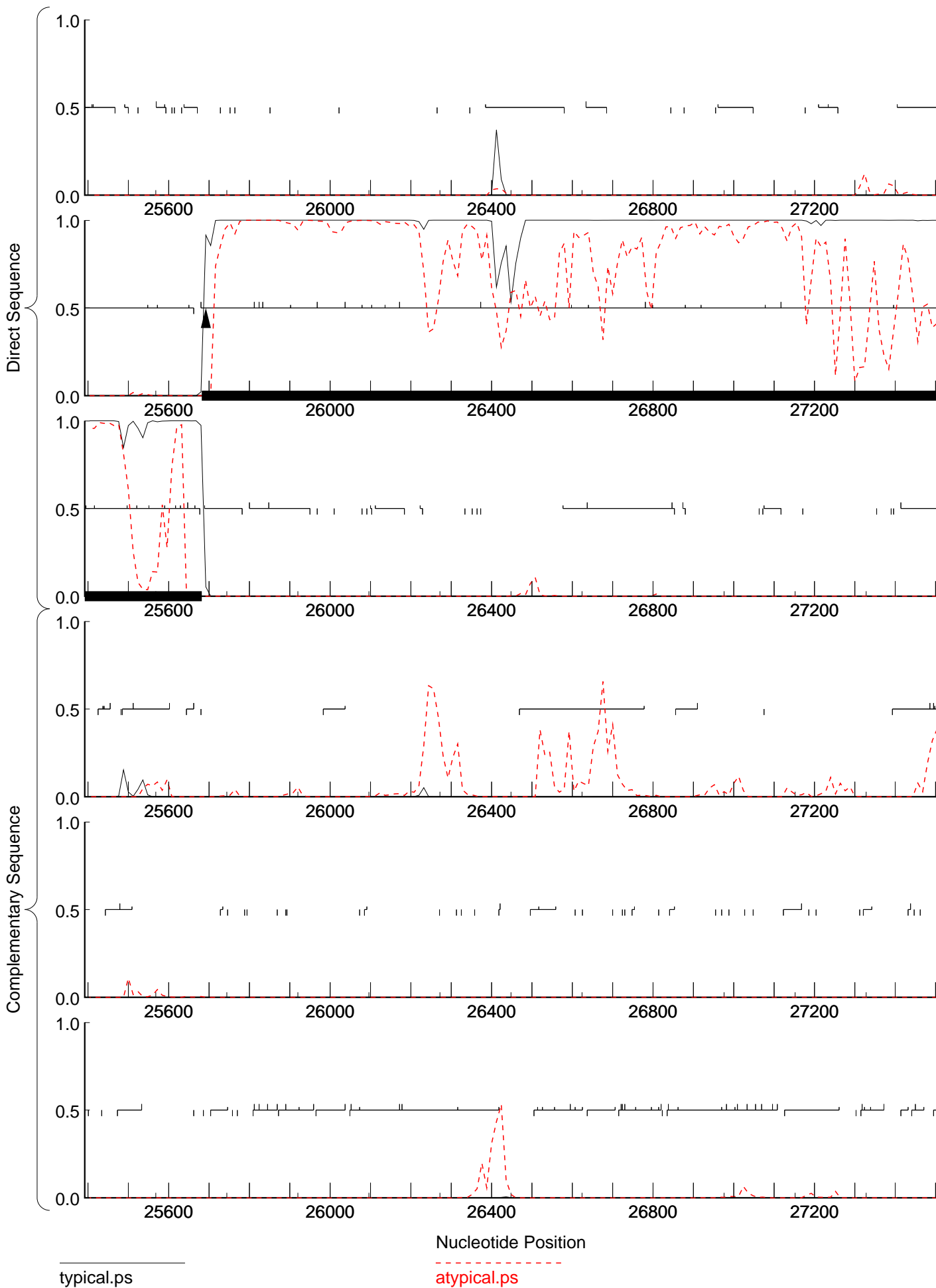


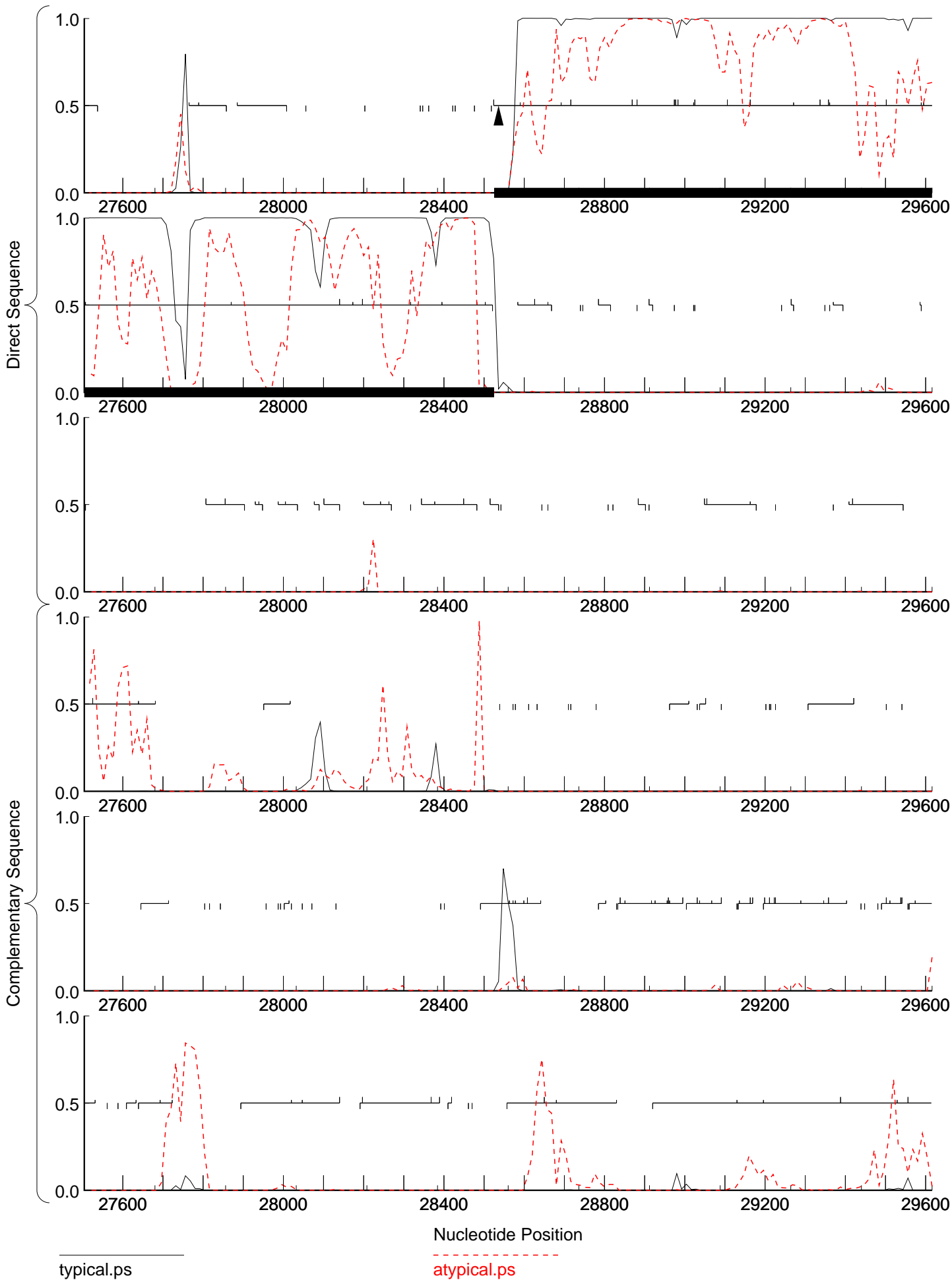


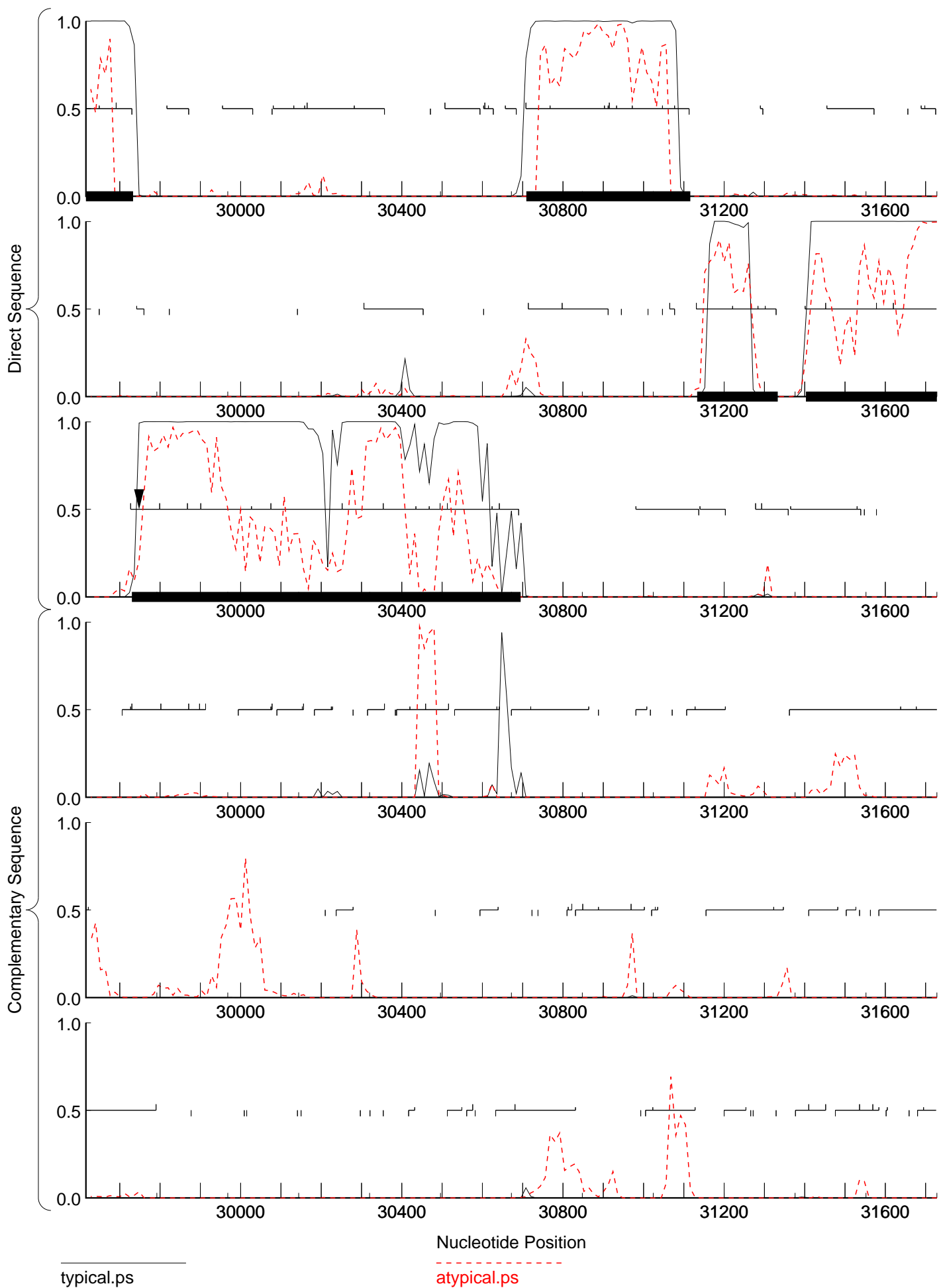


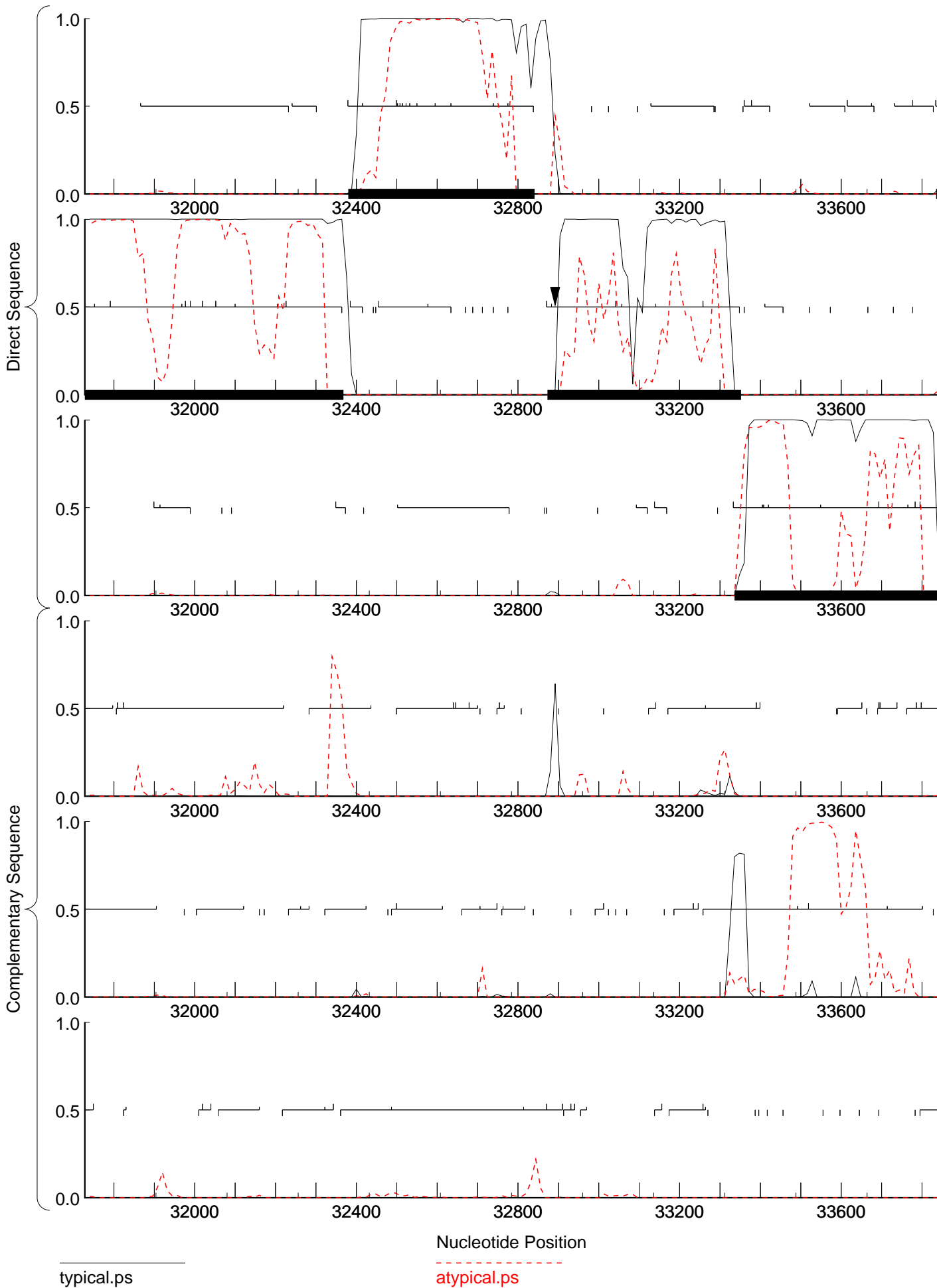
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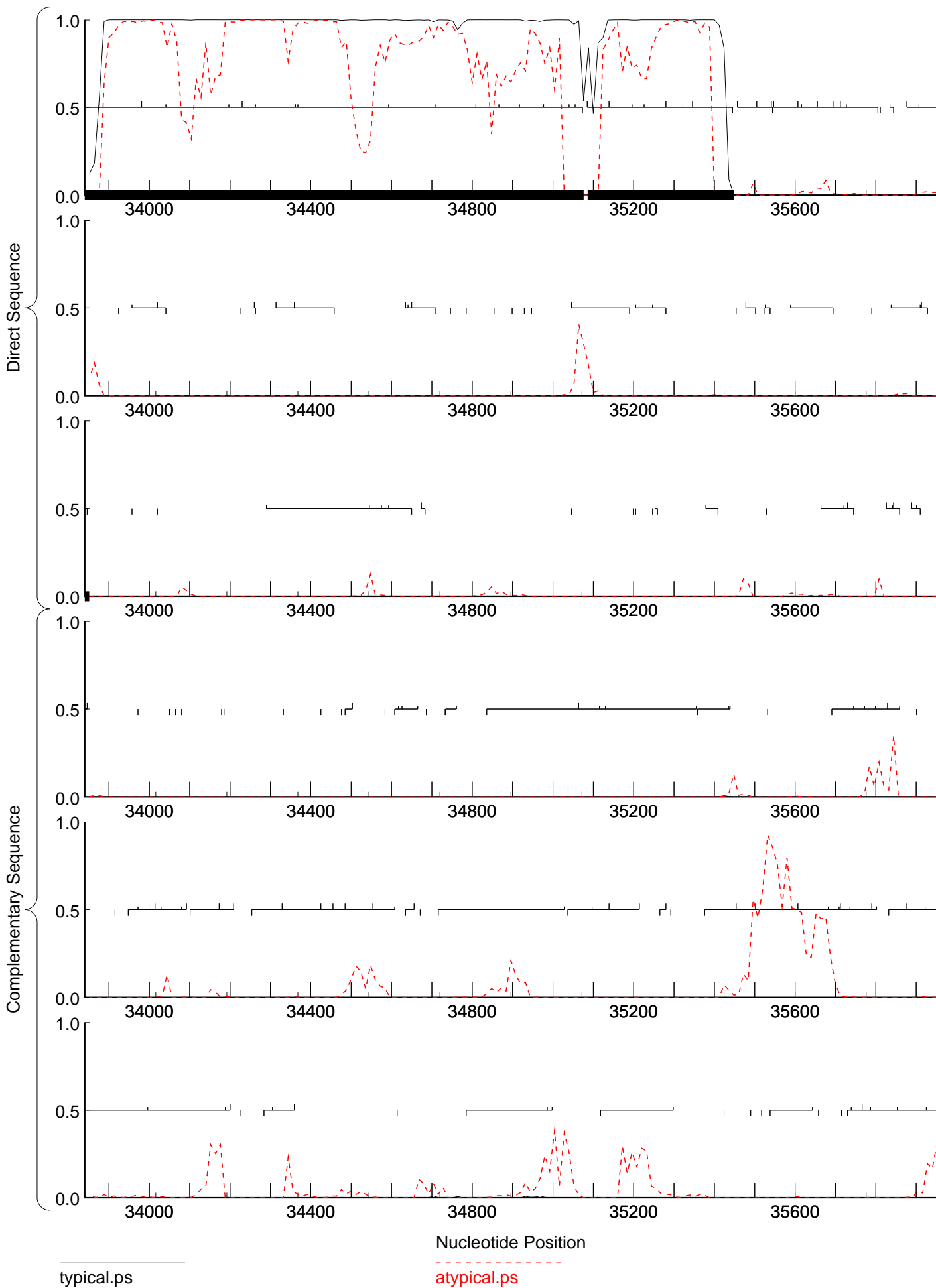
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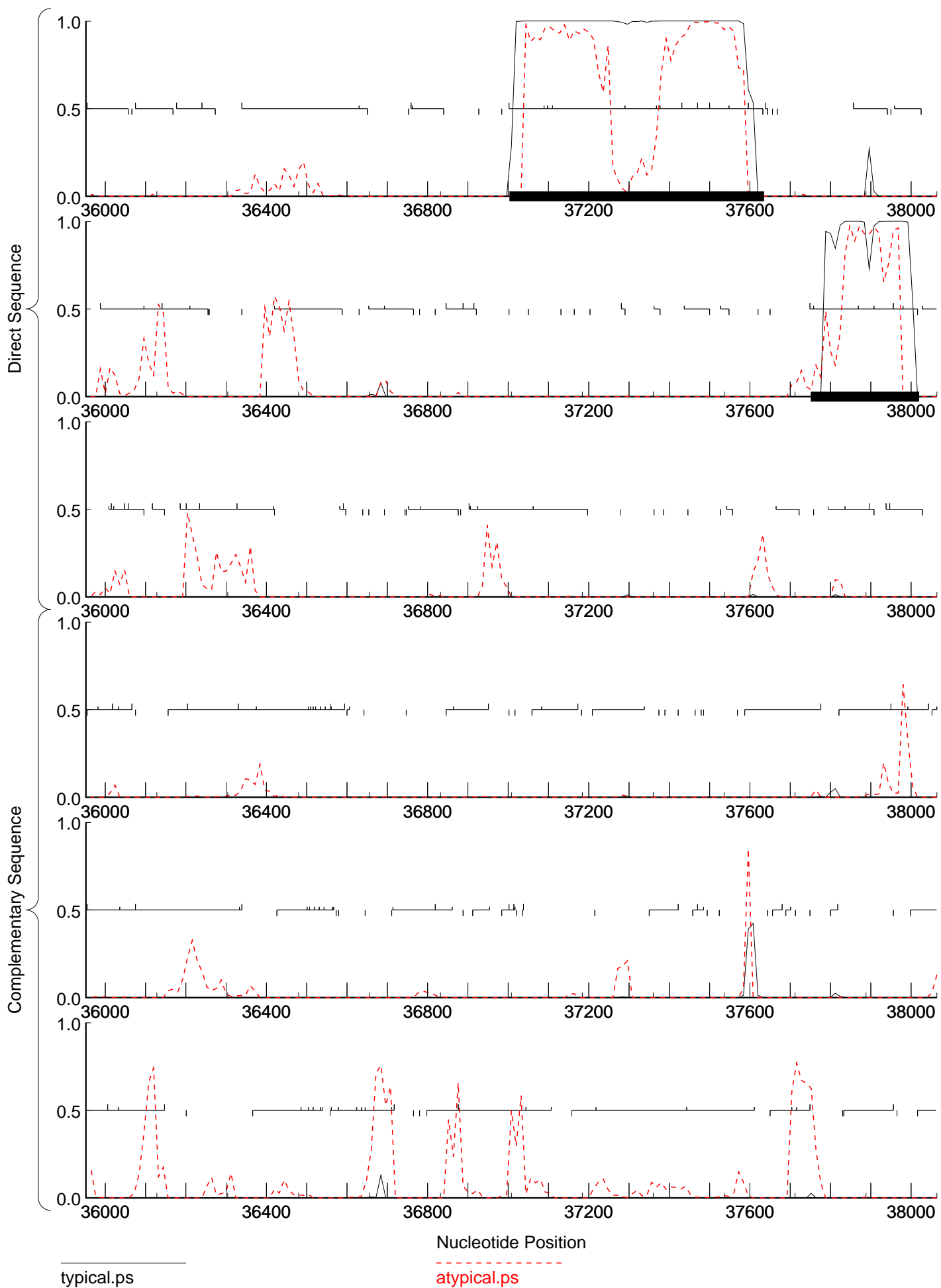


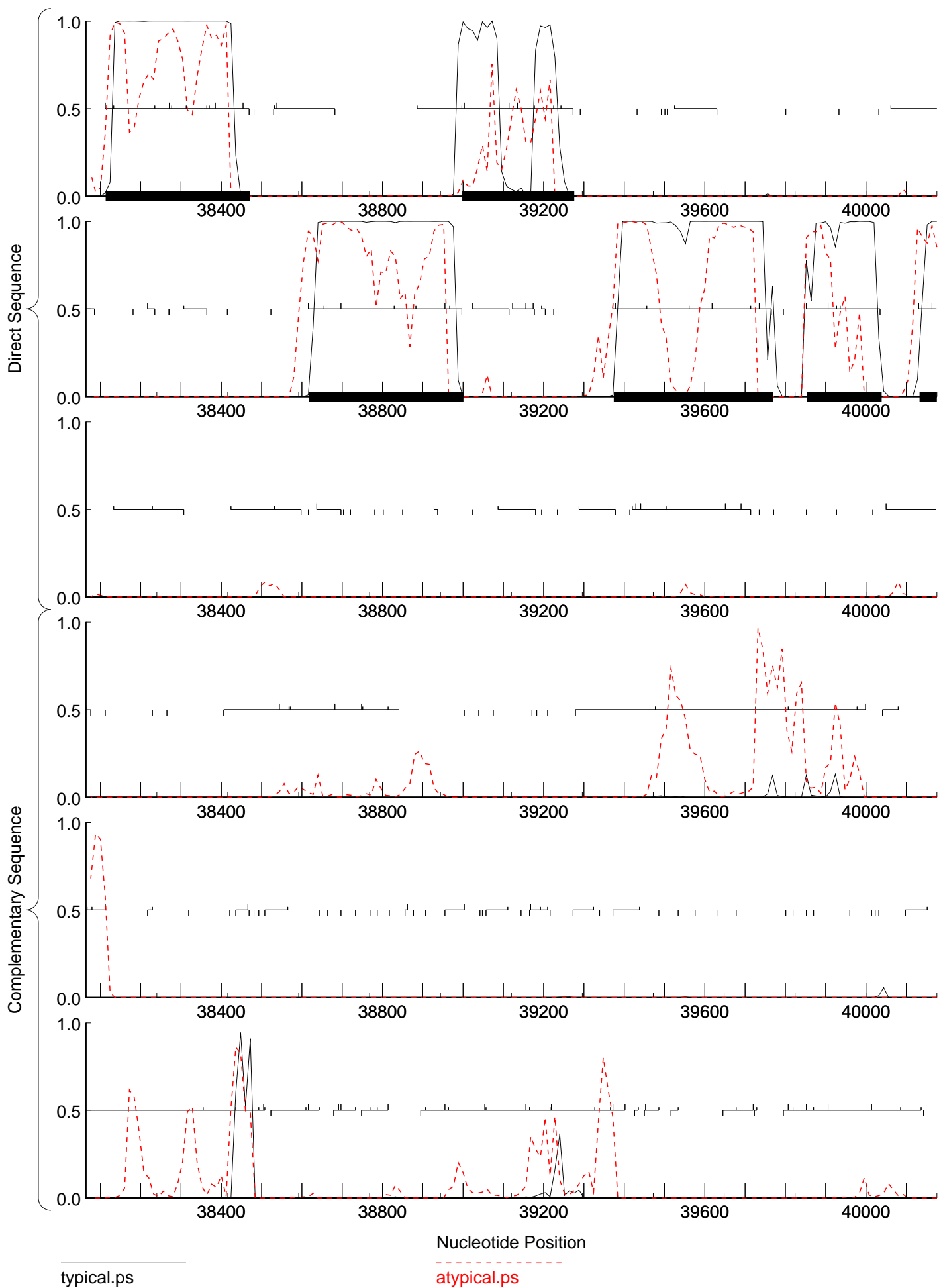






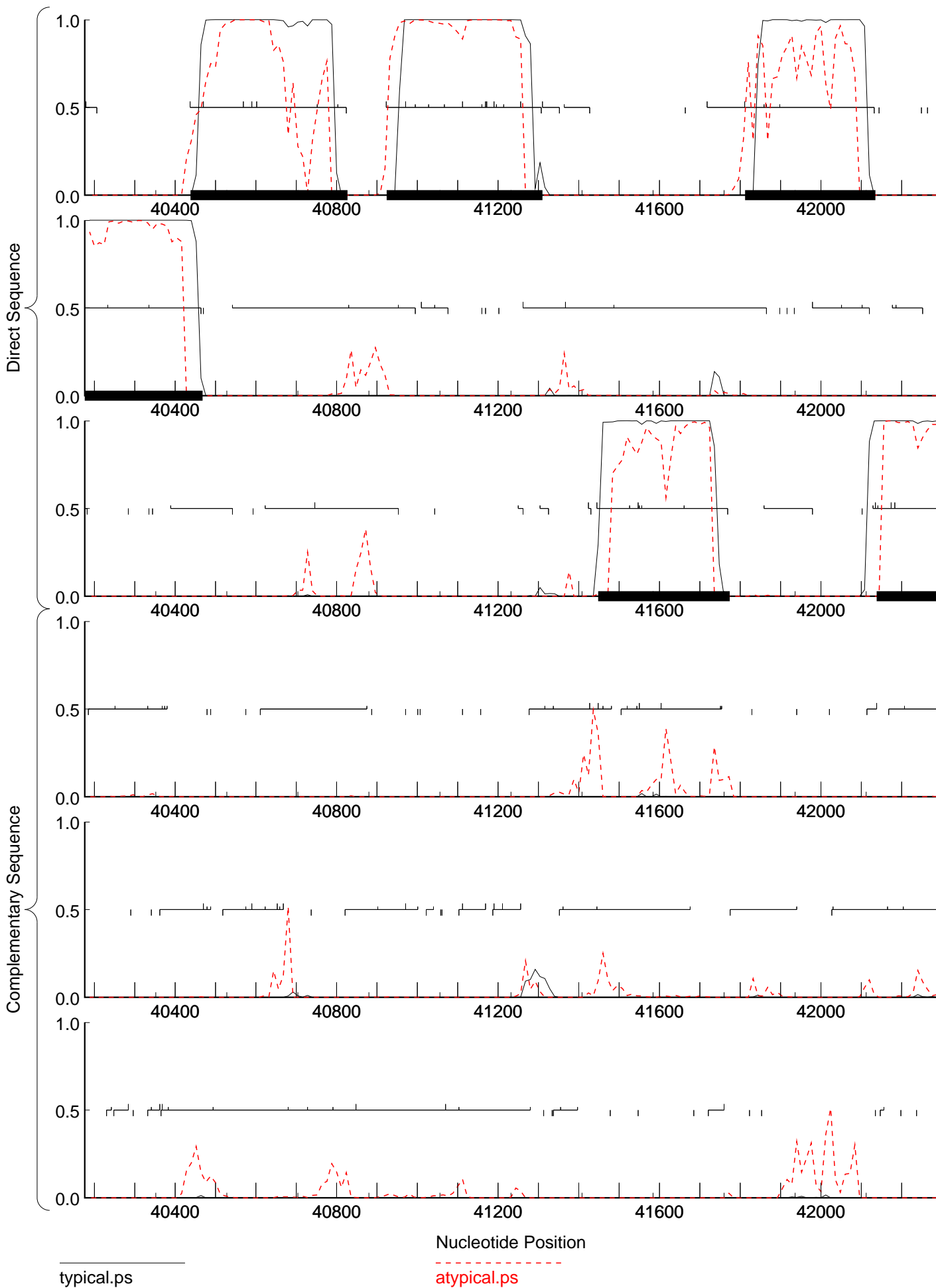






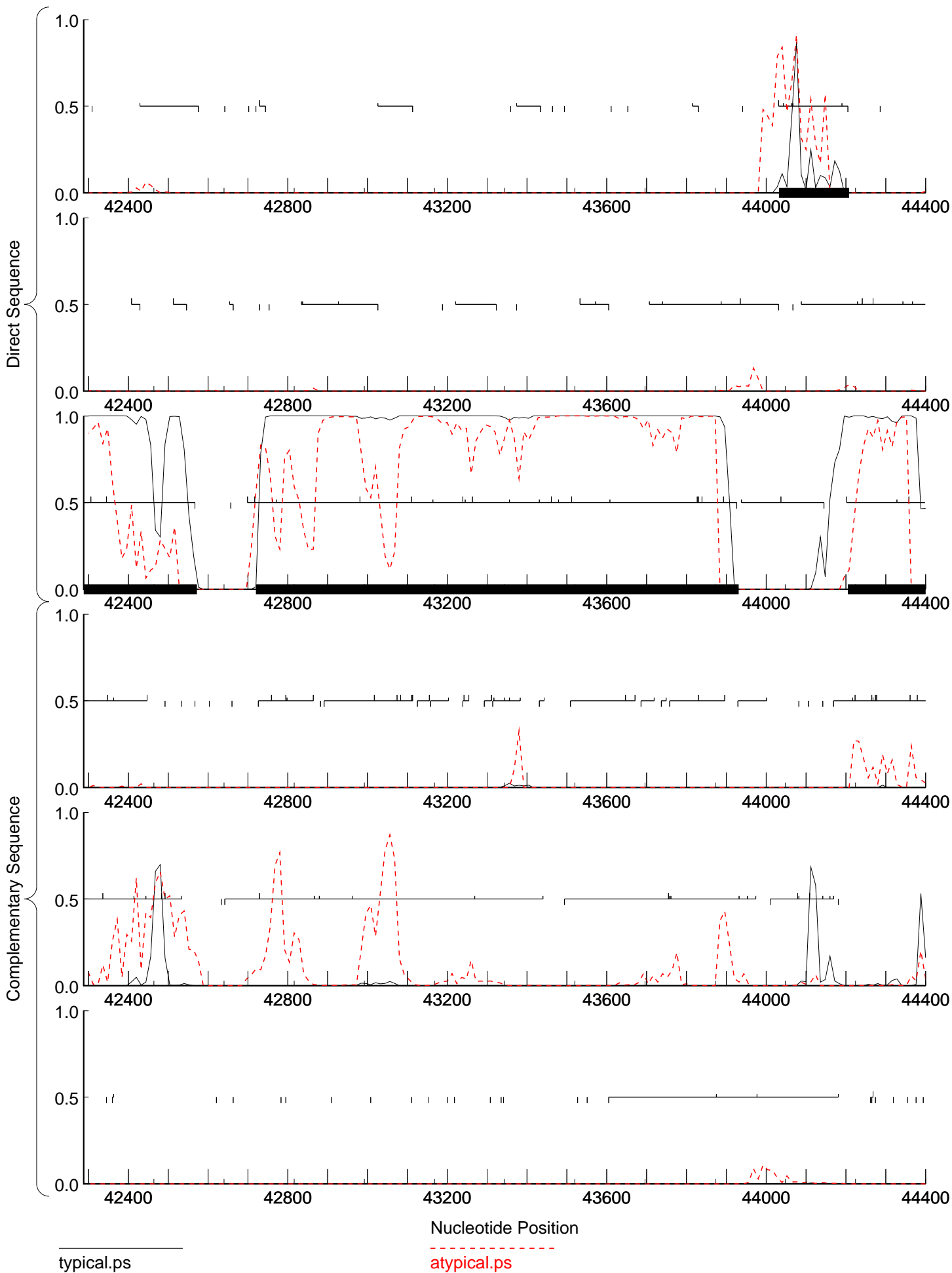
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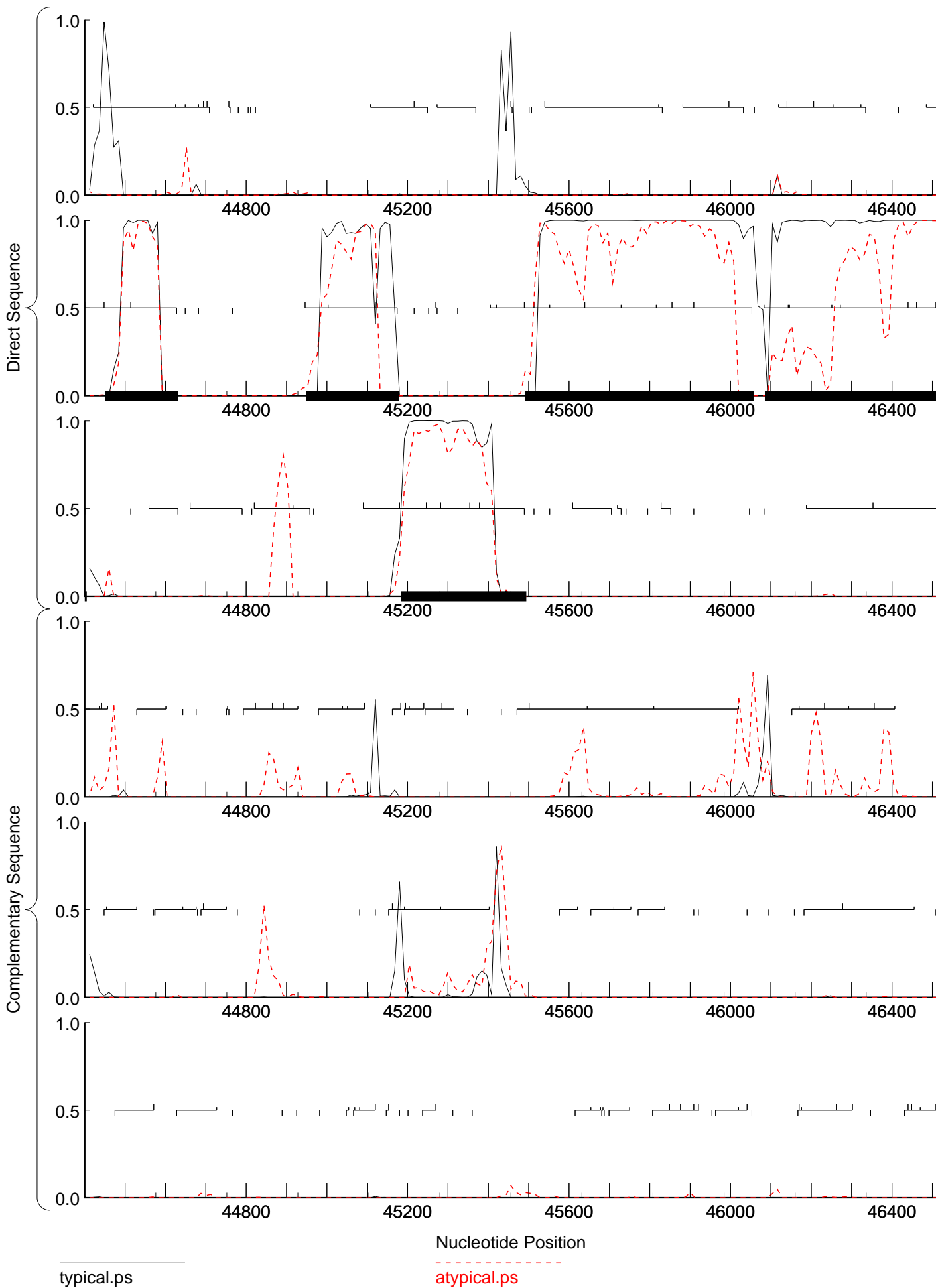
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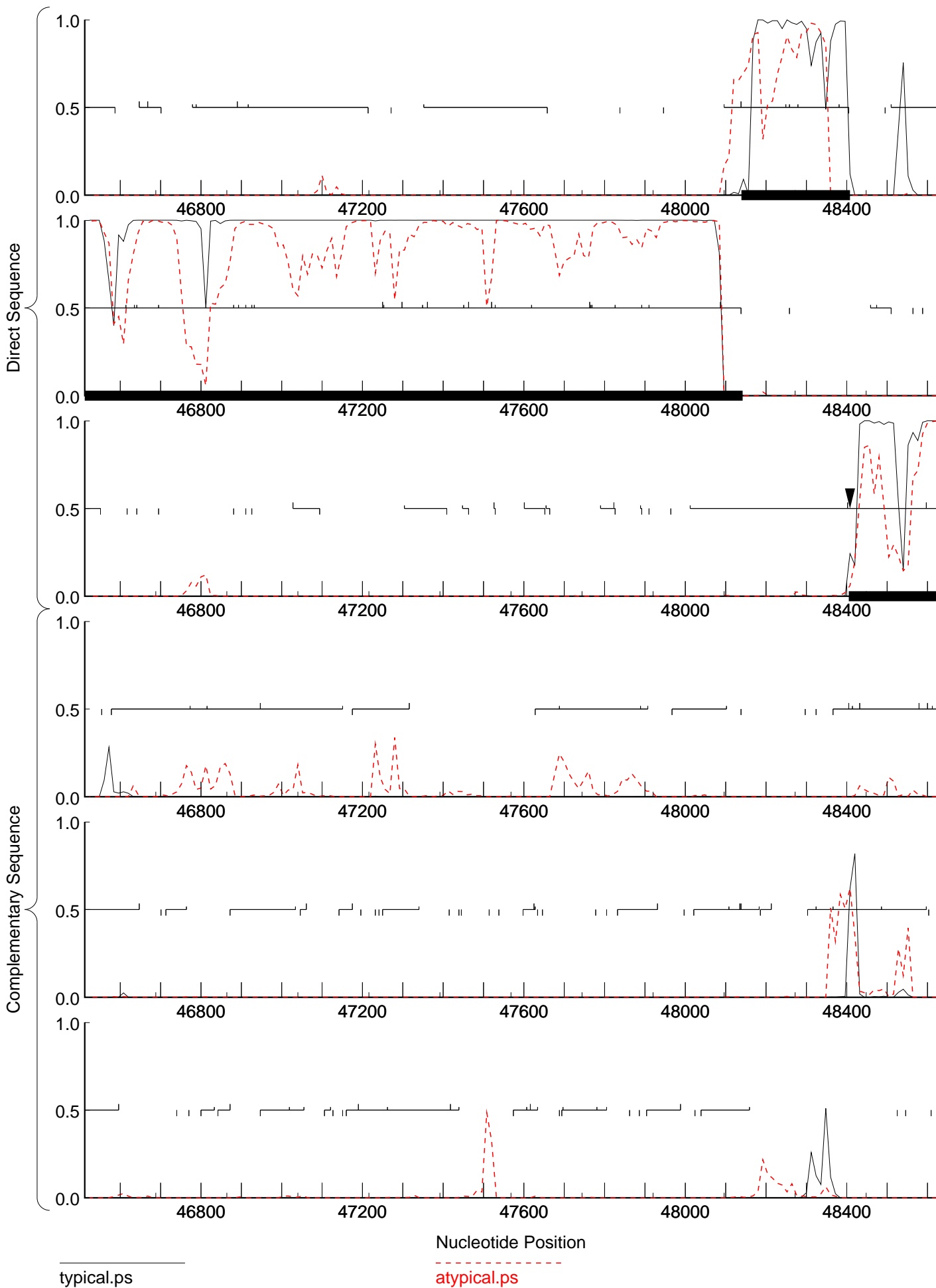


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atypical.ps

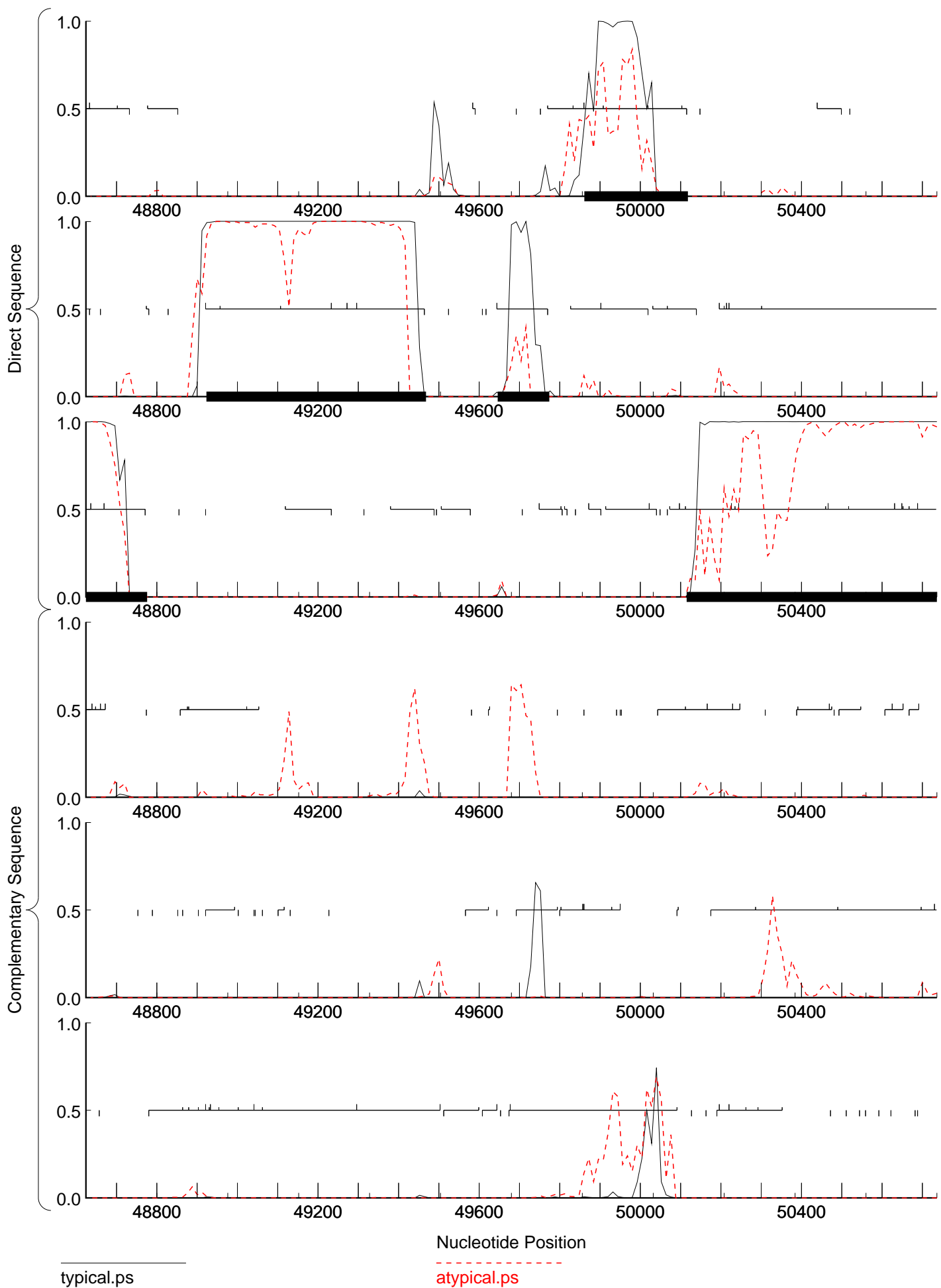






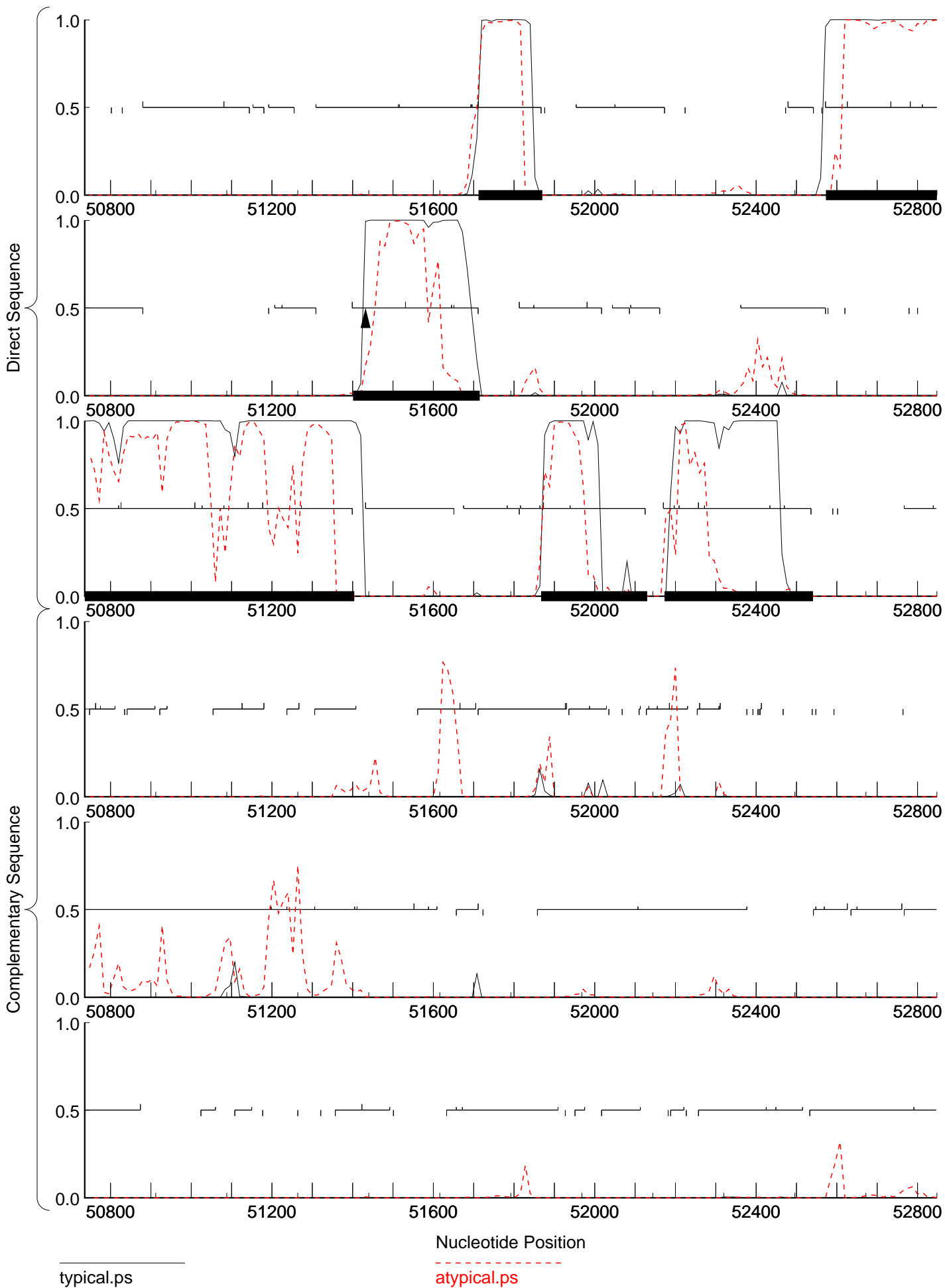
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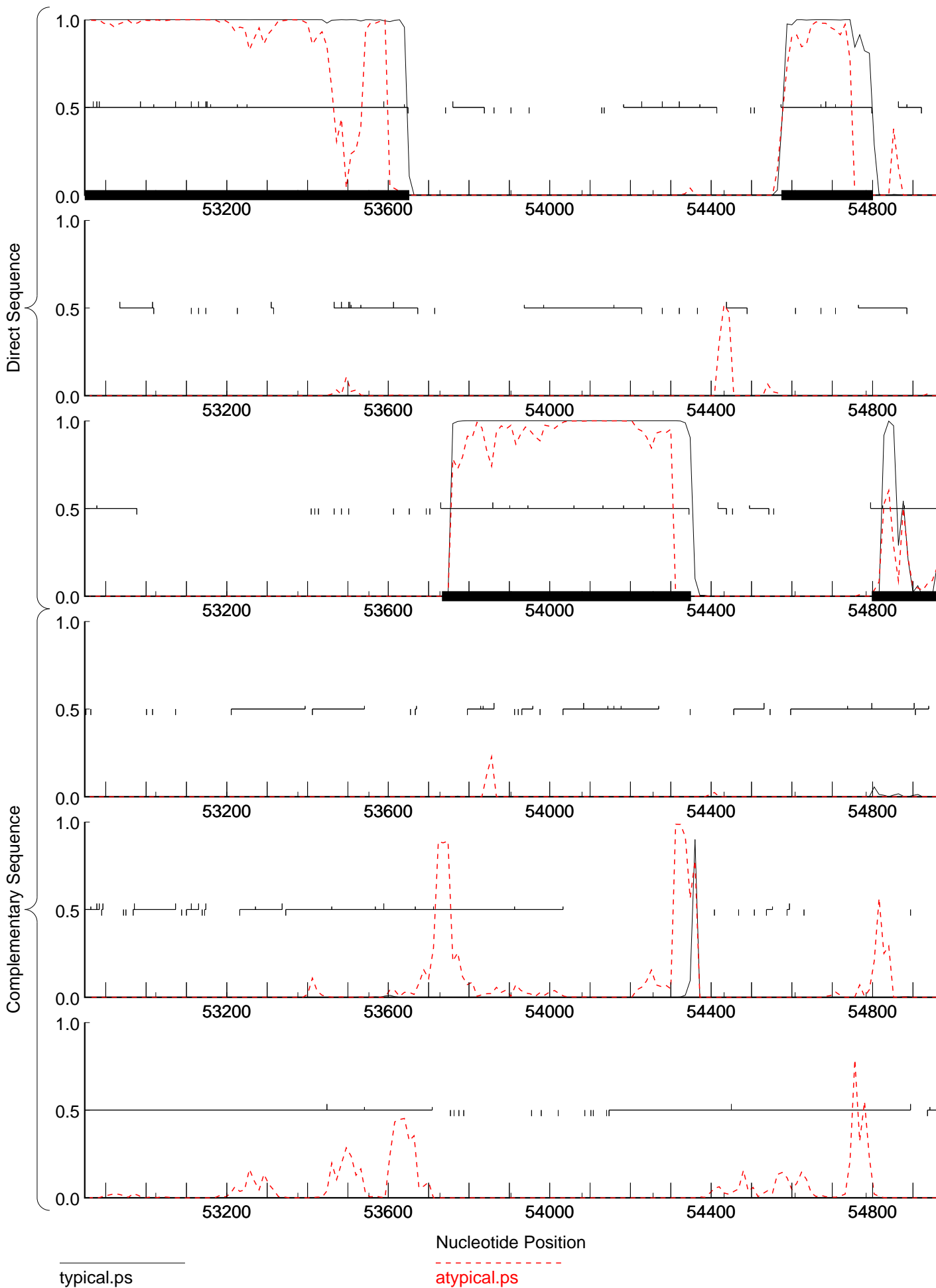
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typical.ps

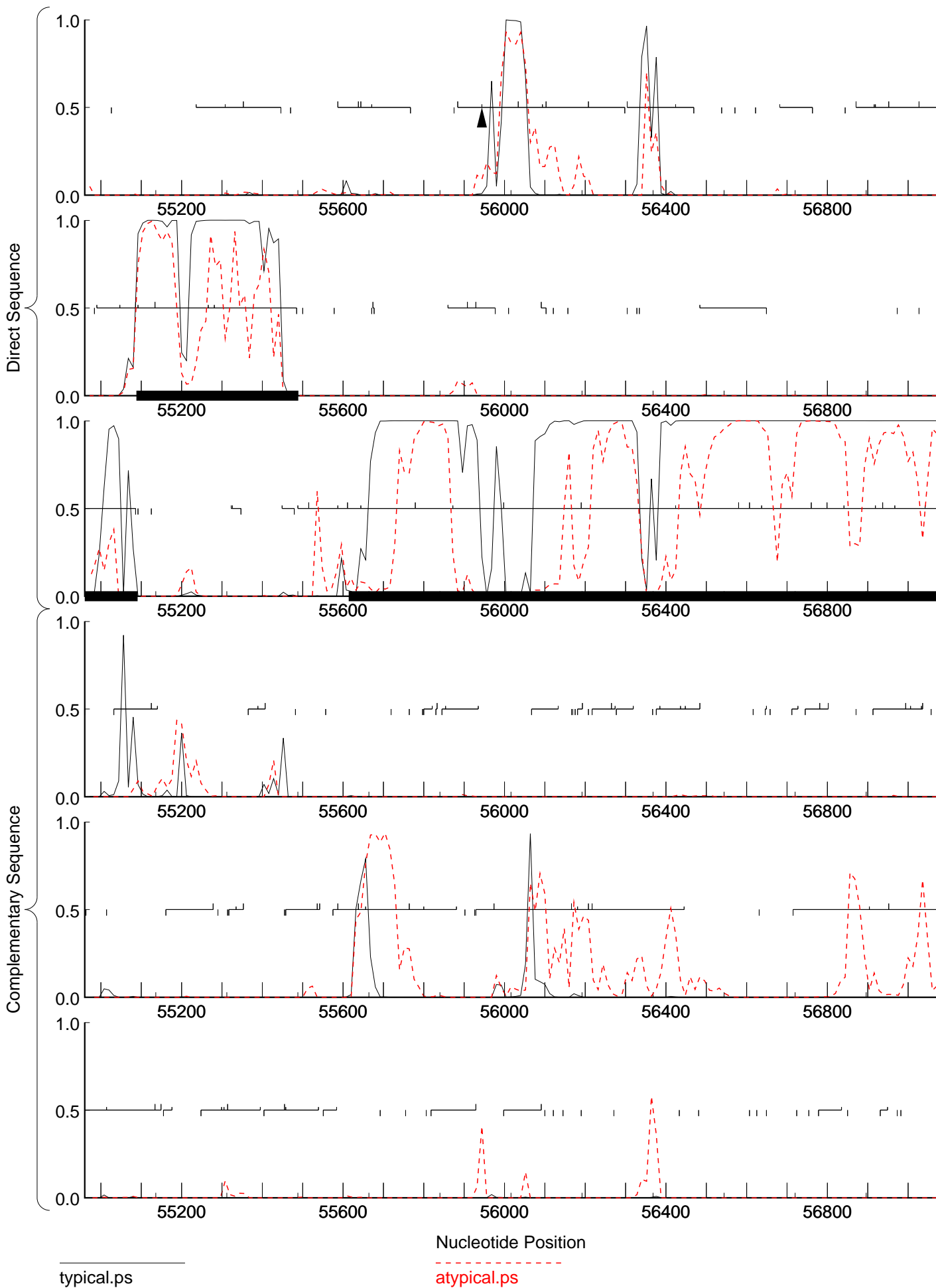
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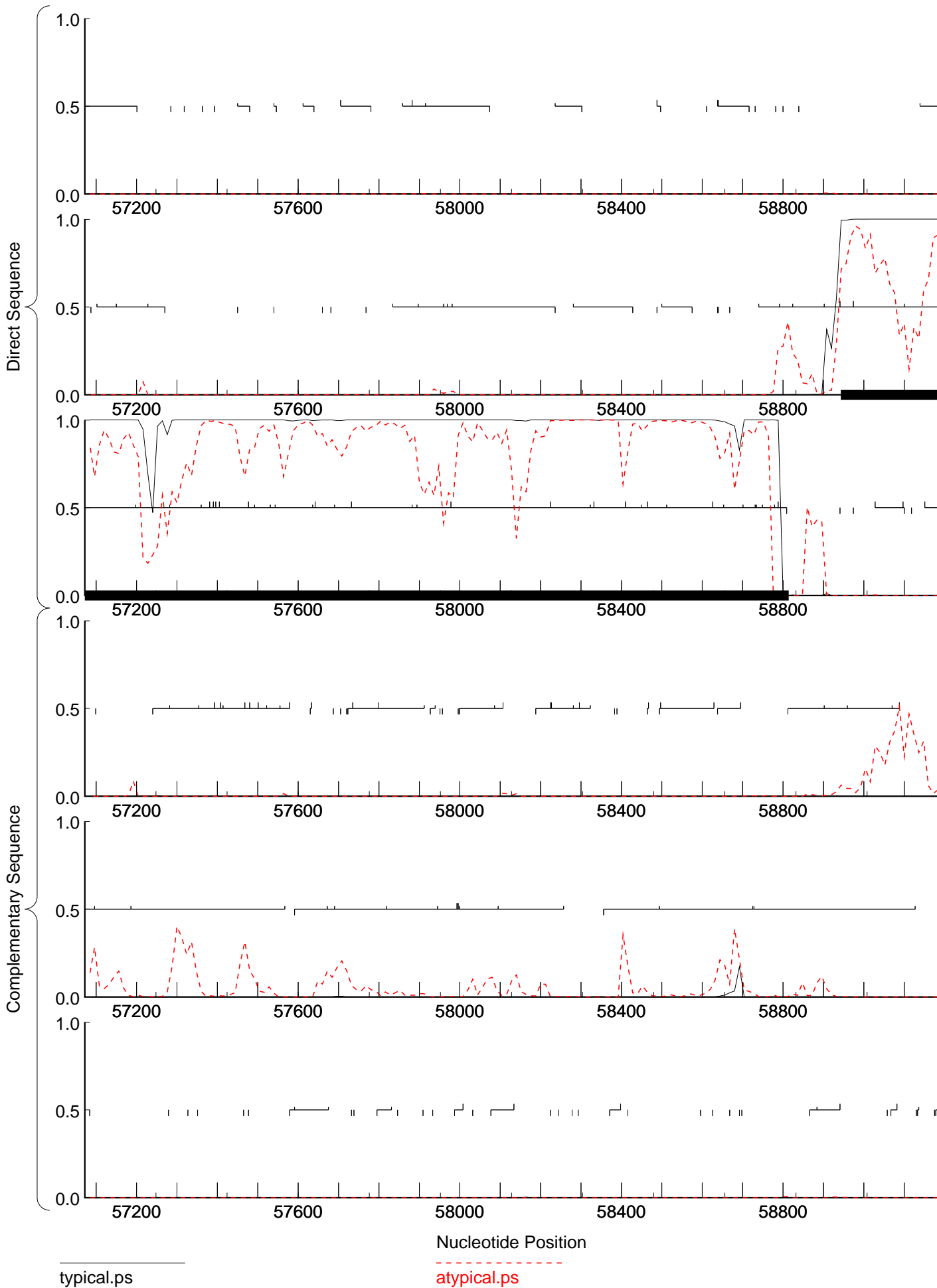
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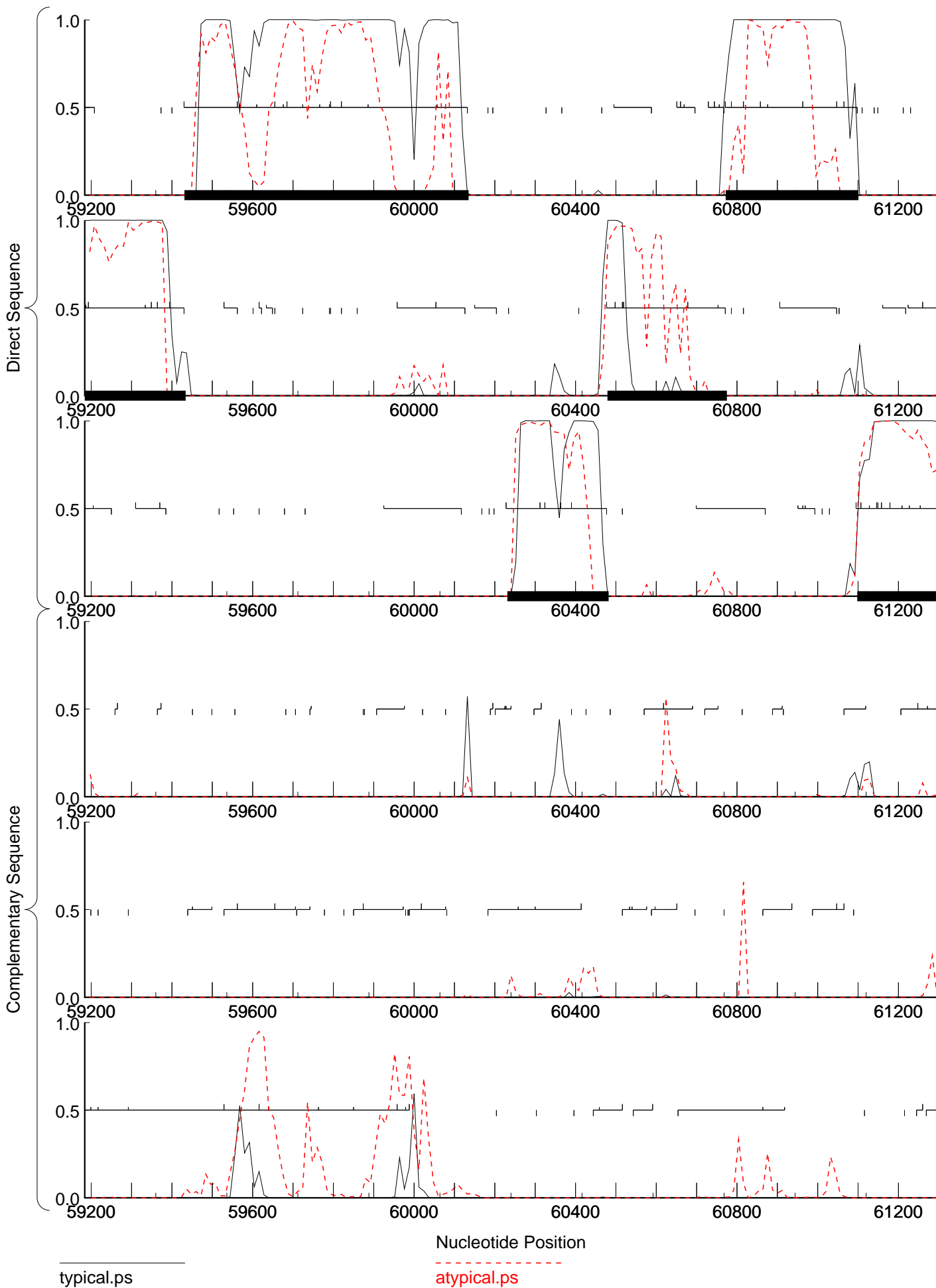
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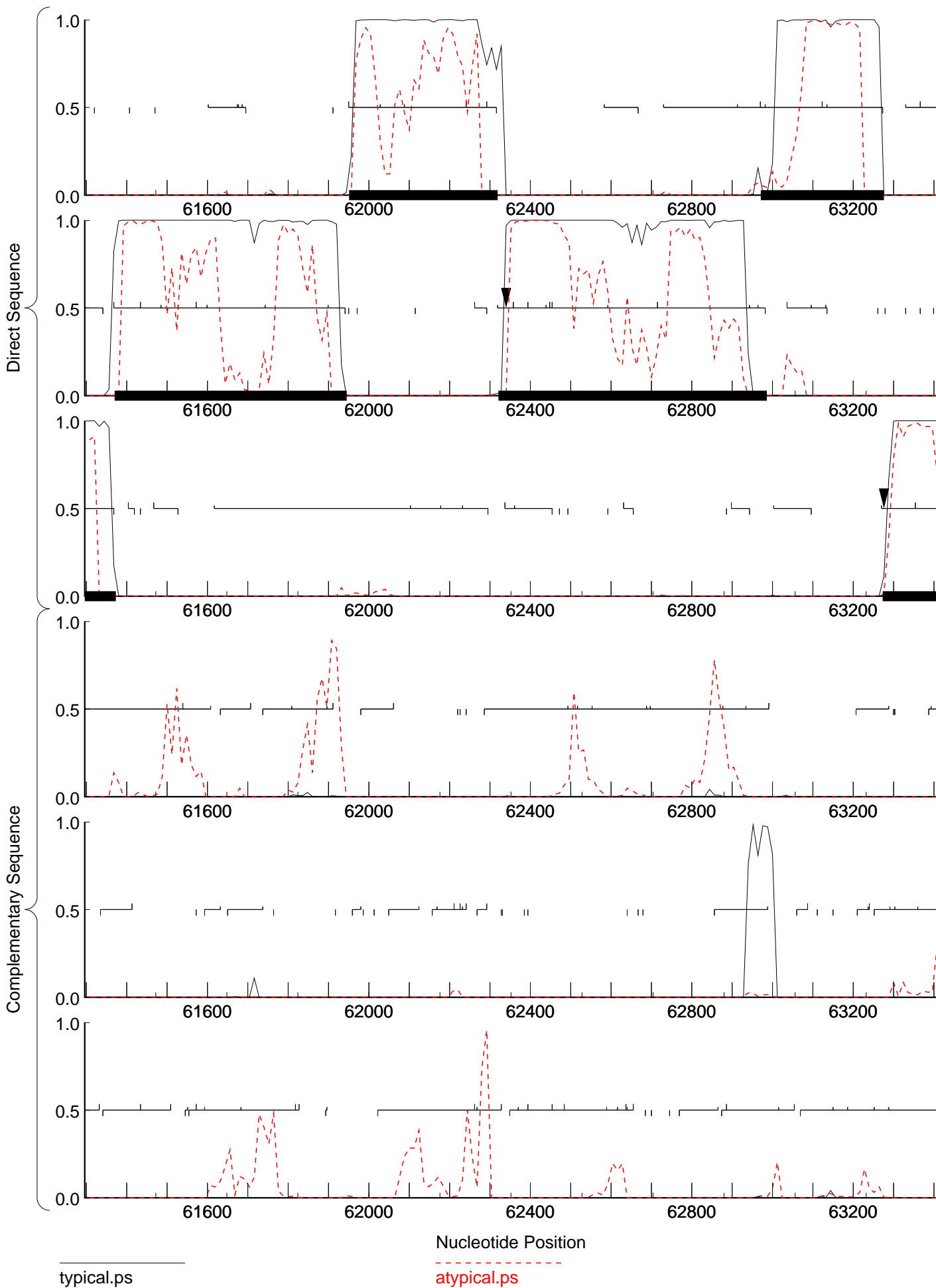
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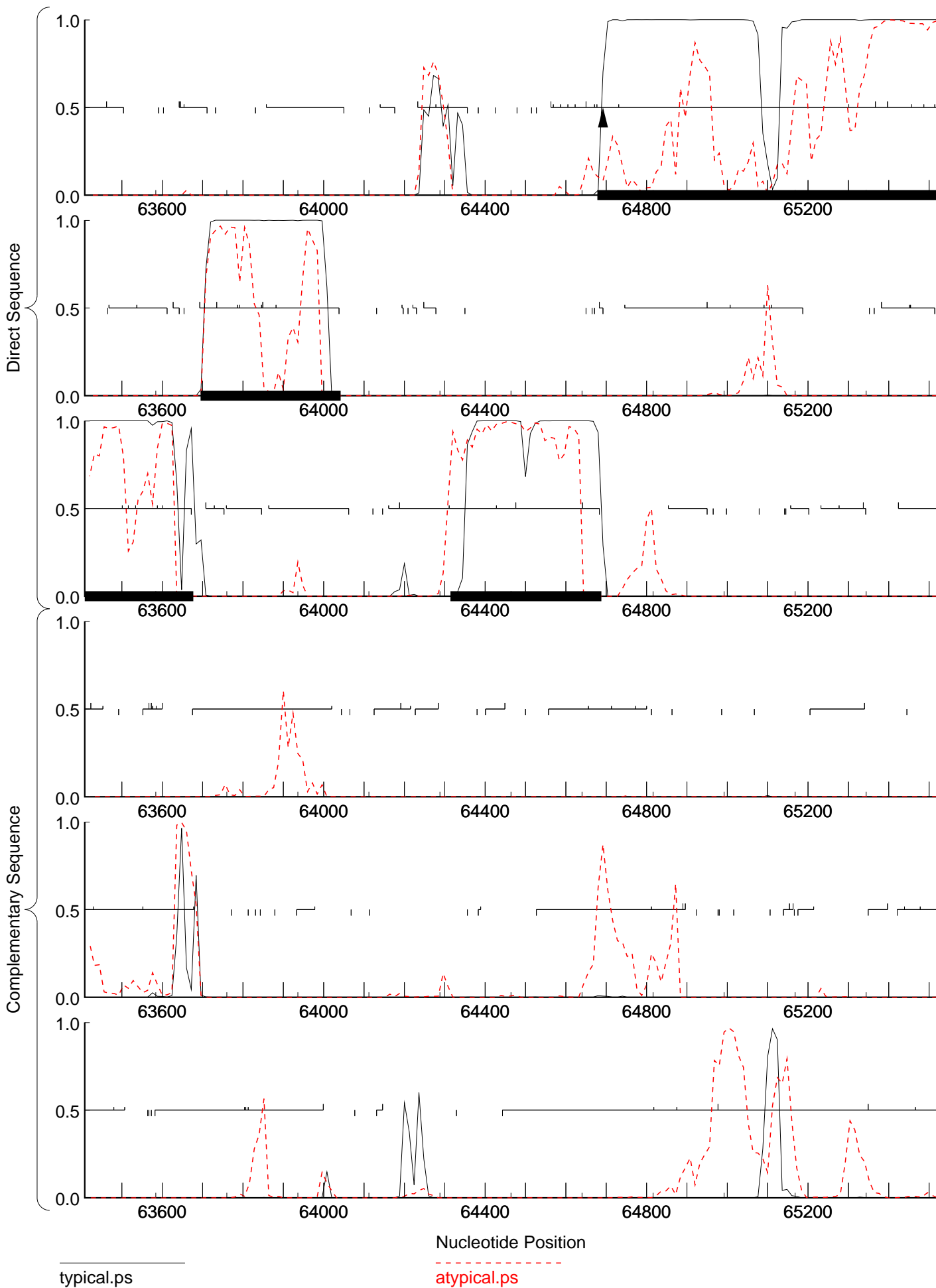


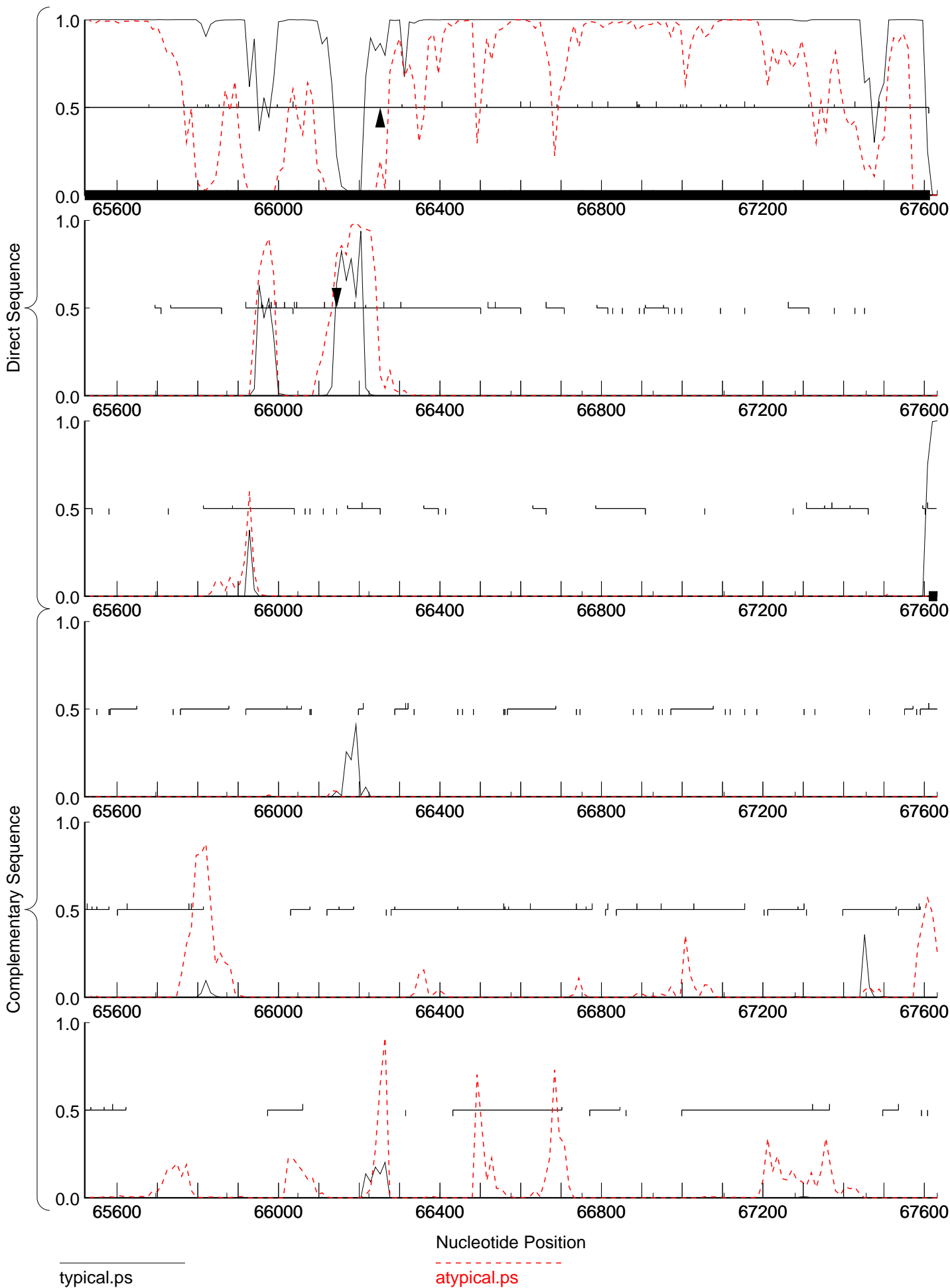


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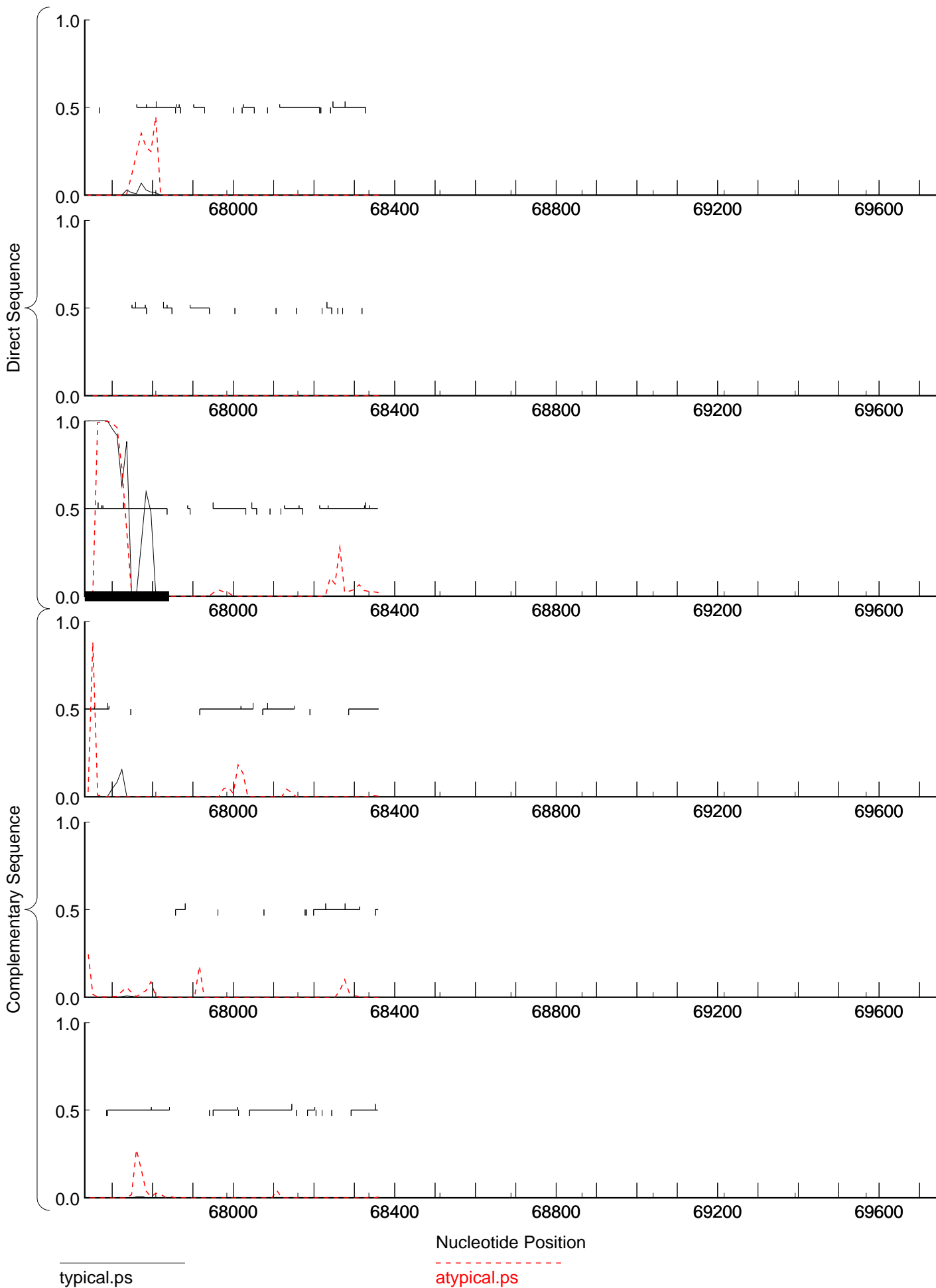






typical.ps

atypical.ps



typical.ps

Nucleotide Position

atypical.ps