

GeneMark

Version 2.5p (09.08.06)

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PROGRAM INFORMATION

Sequence : Streptomyces phage BayC complete sequence, 57243 bp, circularly permuted, Cluster BG
Analysis Date : 8/17/17 at 15:21:19
Pages : 29
Sequence Length : 57243 bp
GC Content : 69.23%

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,

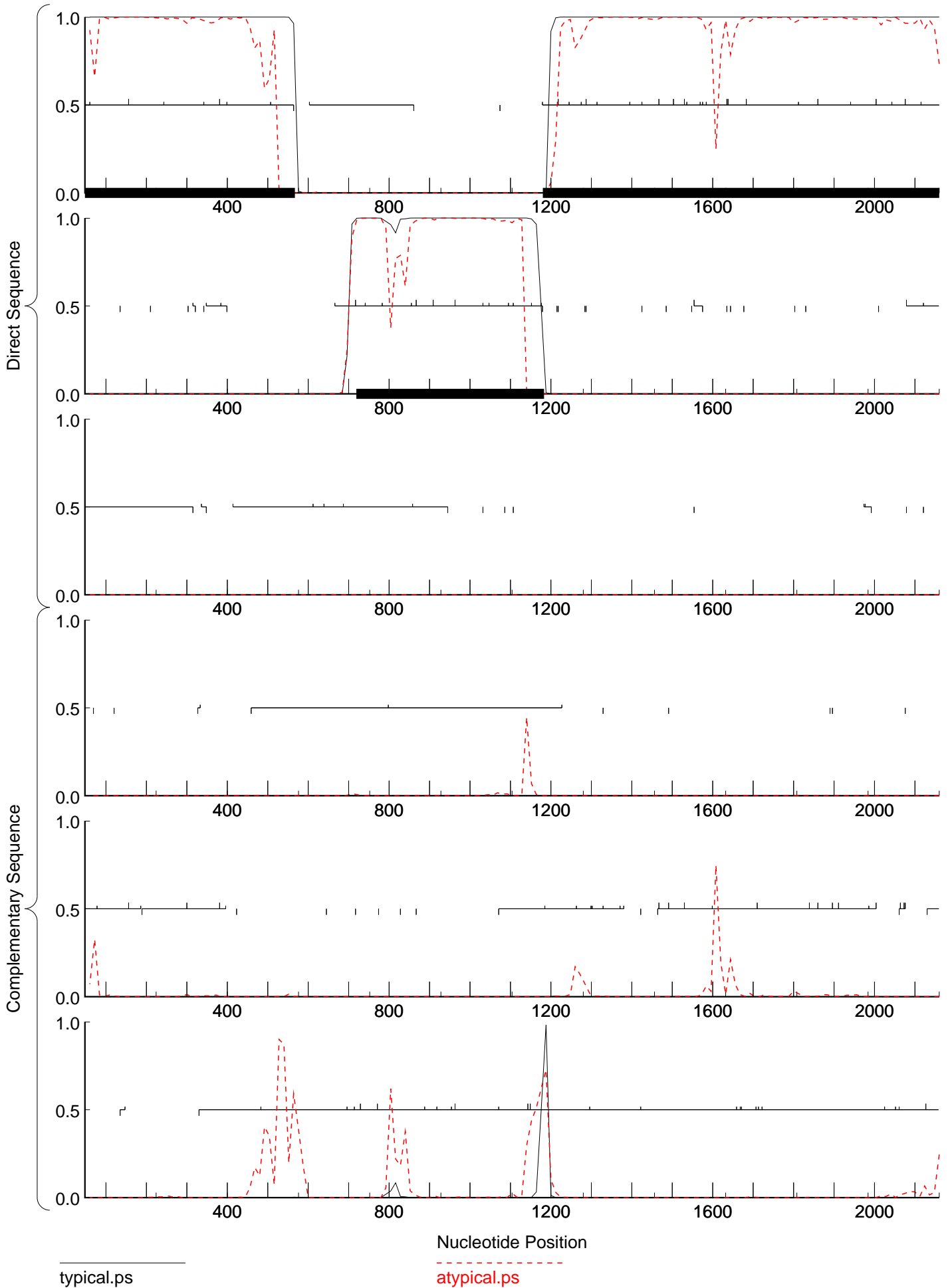
Matrix notes & comments

Training set derived by GeneMarkS, 4.30 October 2014
Thu Aug 17 15:21:19 2017

MATRIX INFORMATION

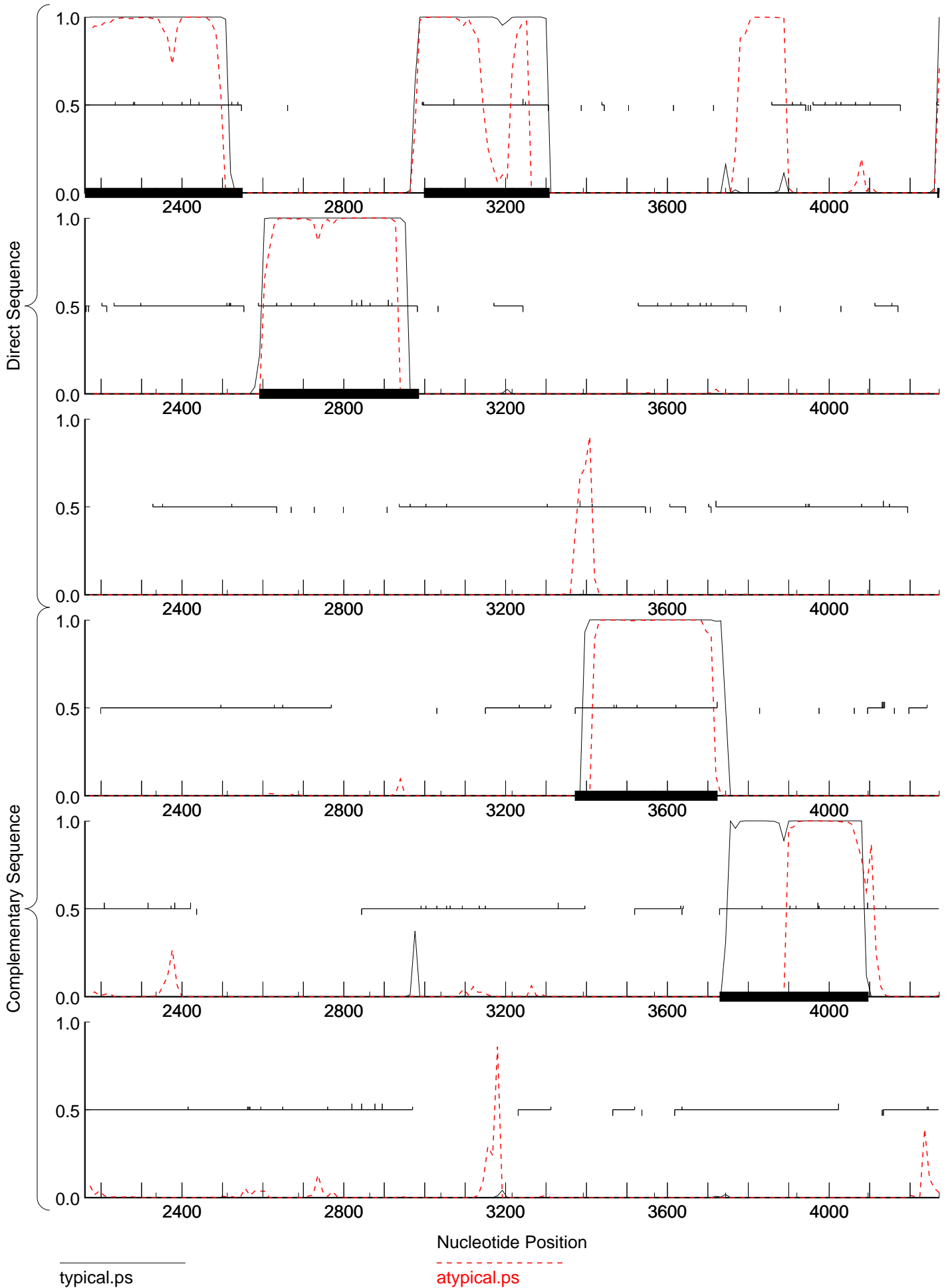
Matrix : -
Author : -
Order : 4

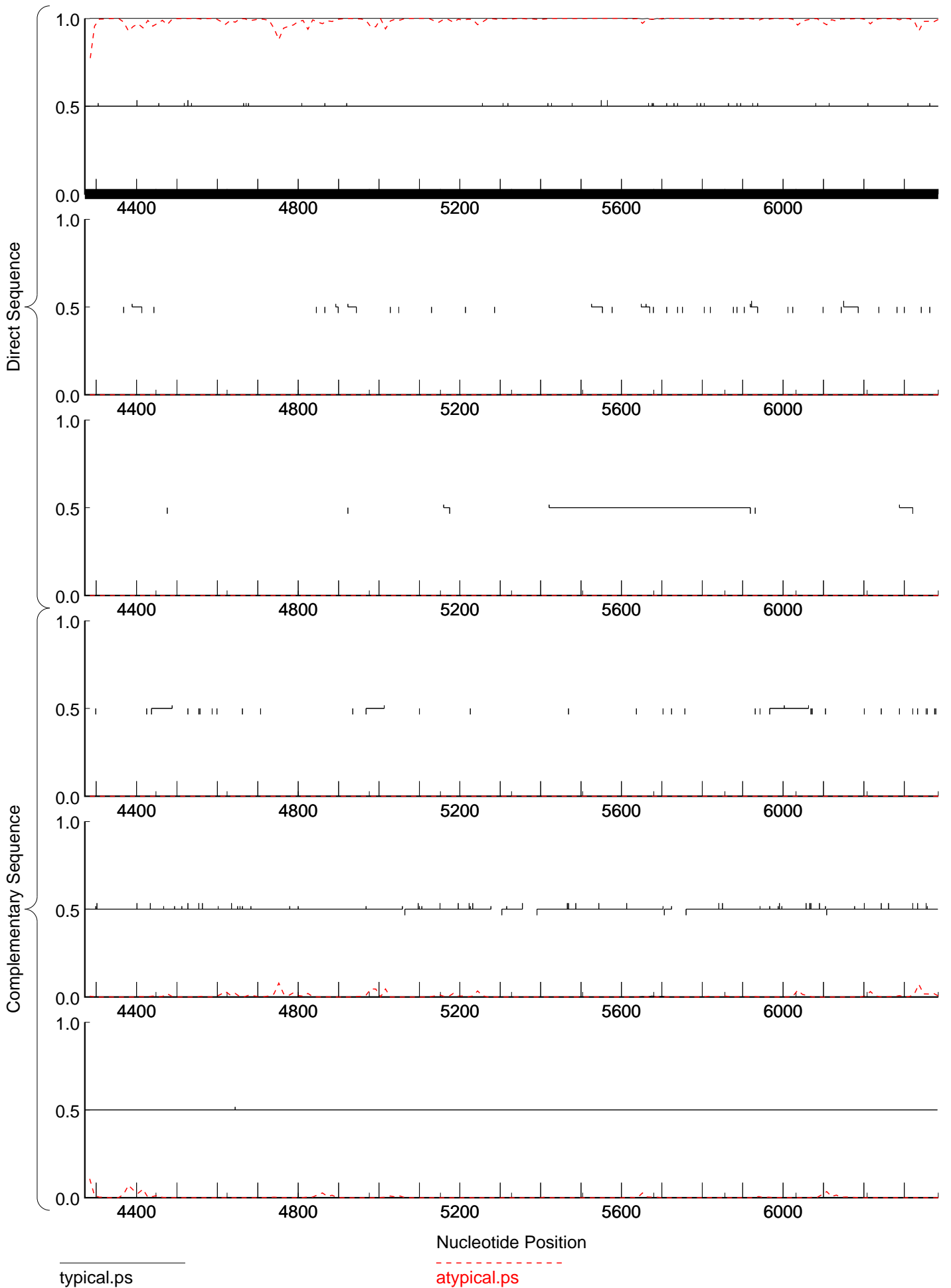
Send questions / comments to:
Dr. M. Borodovsky
Georgia Institute of Technology
School of Biology
Atlanta, GA 30332-0230

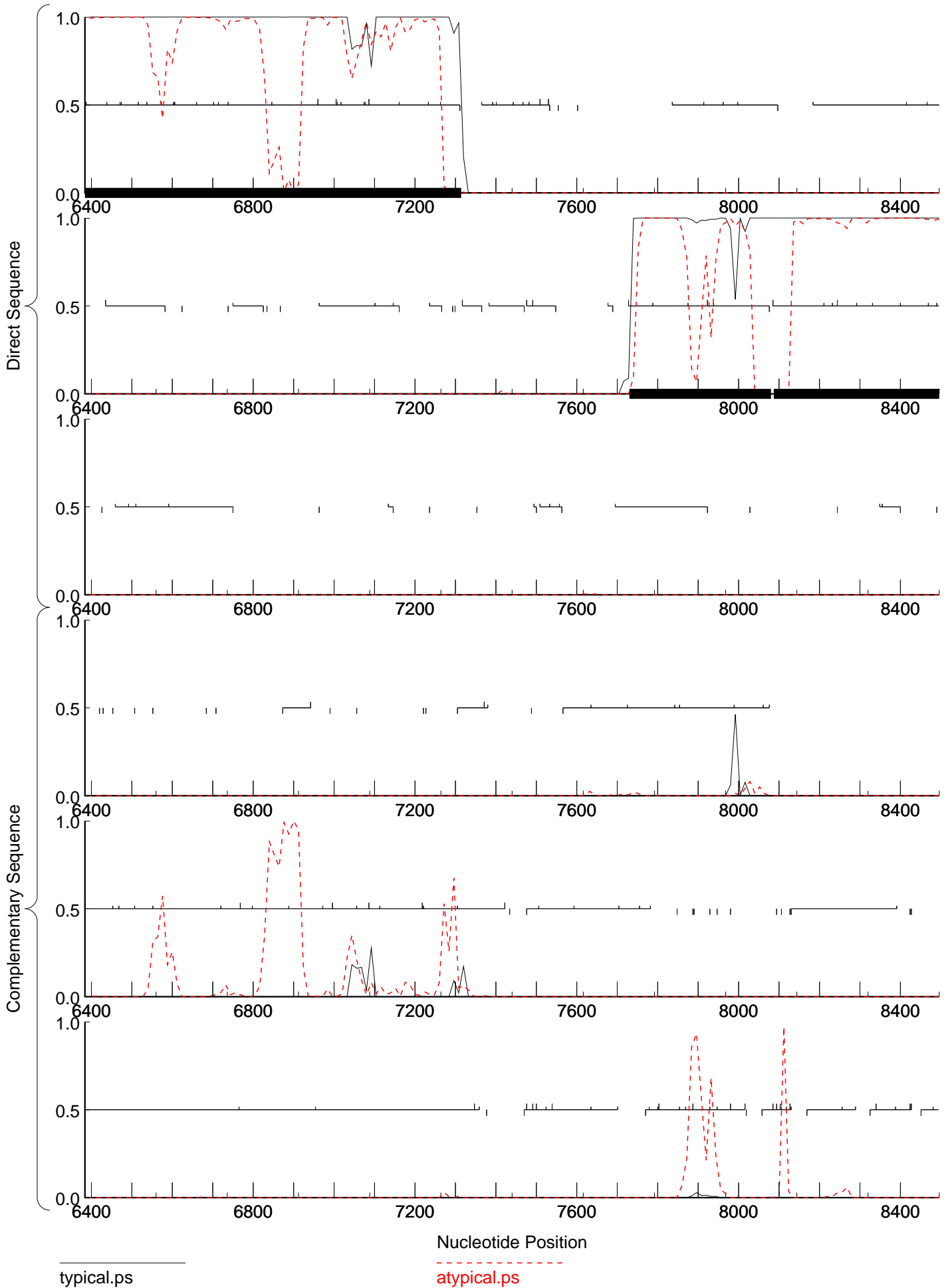


typical.ps

atypical.ps

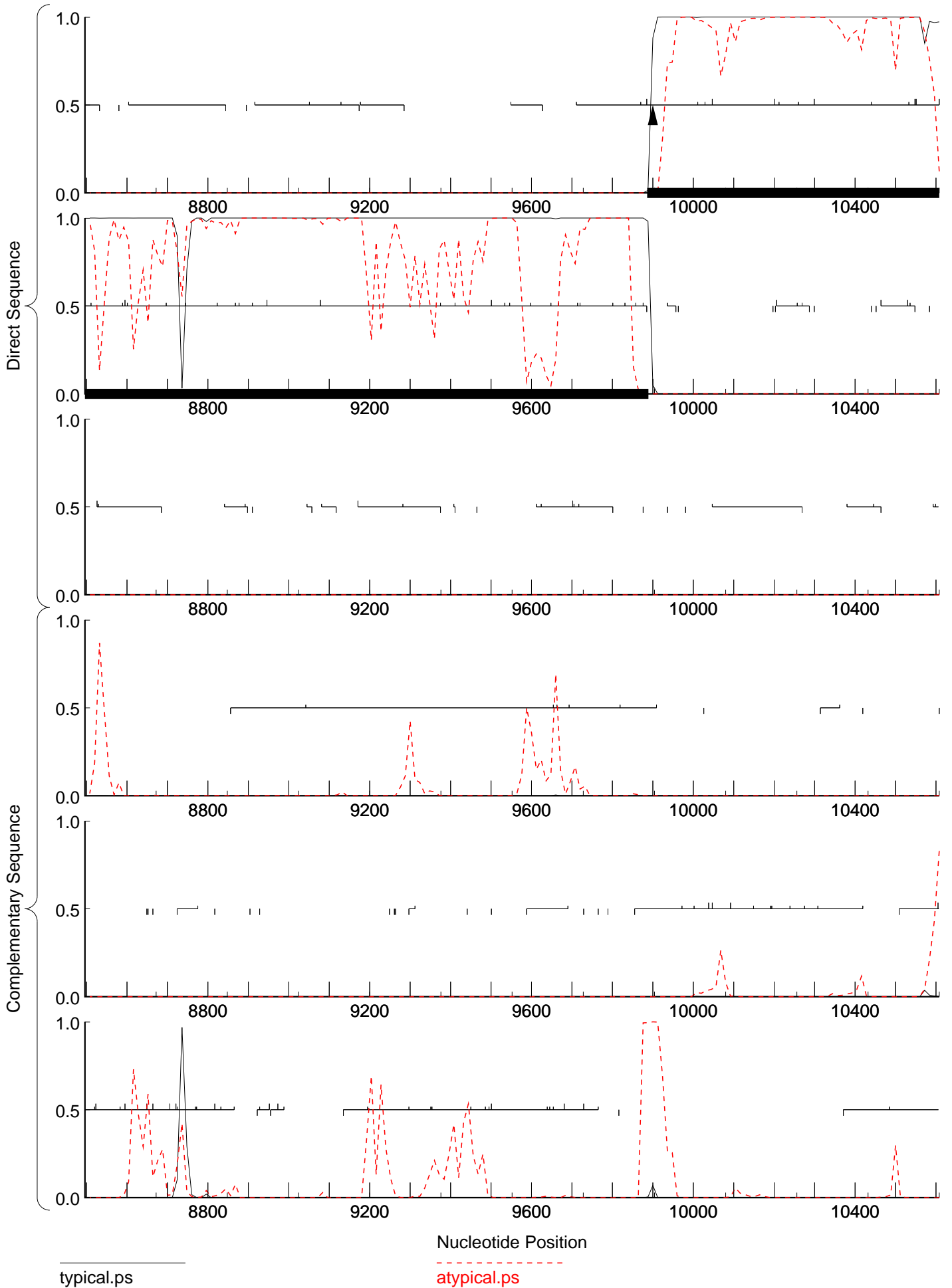


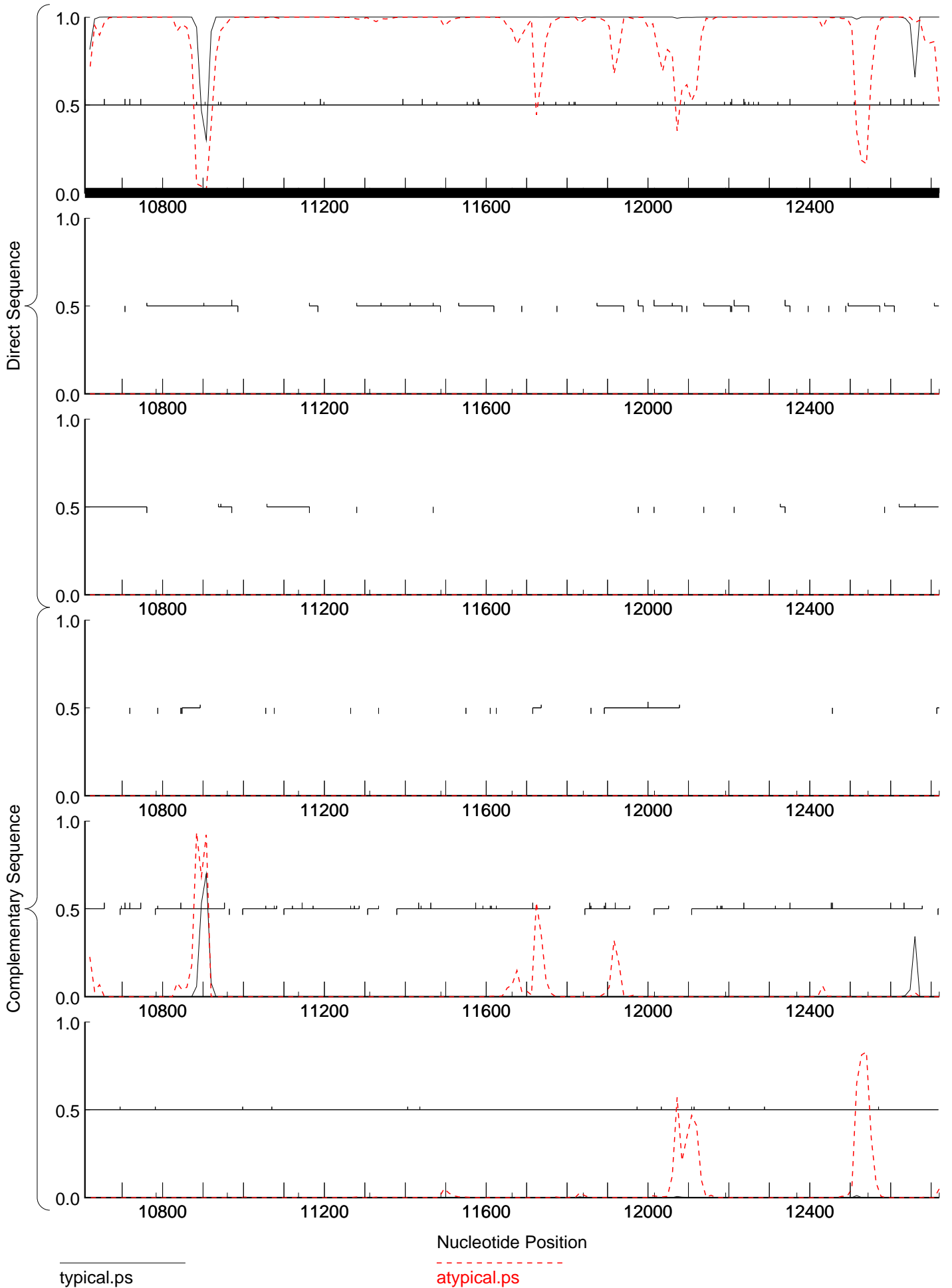


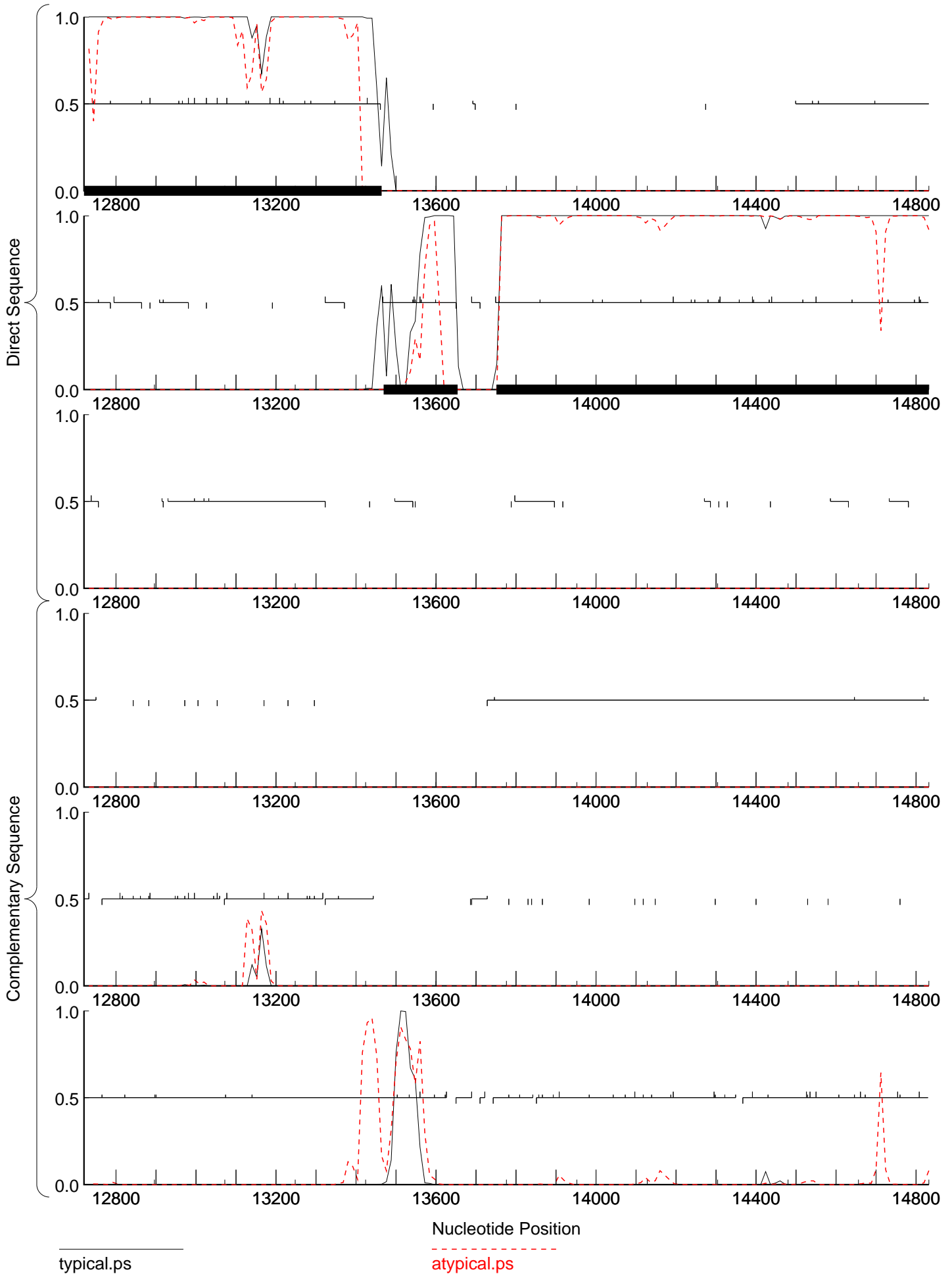


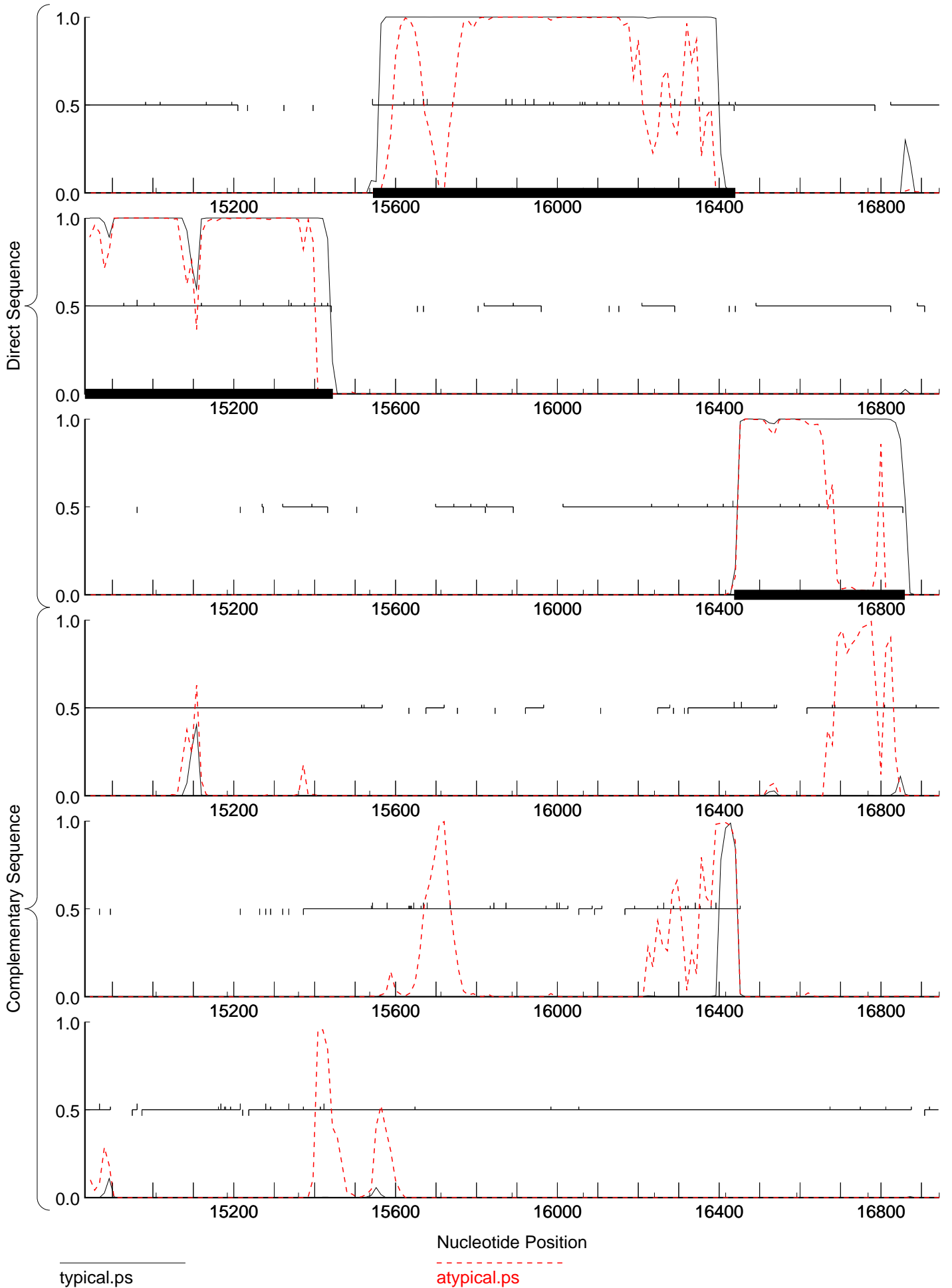
typical.ps

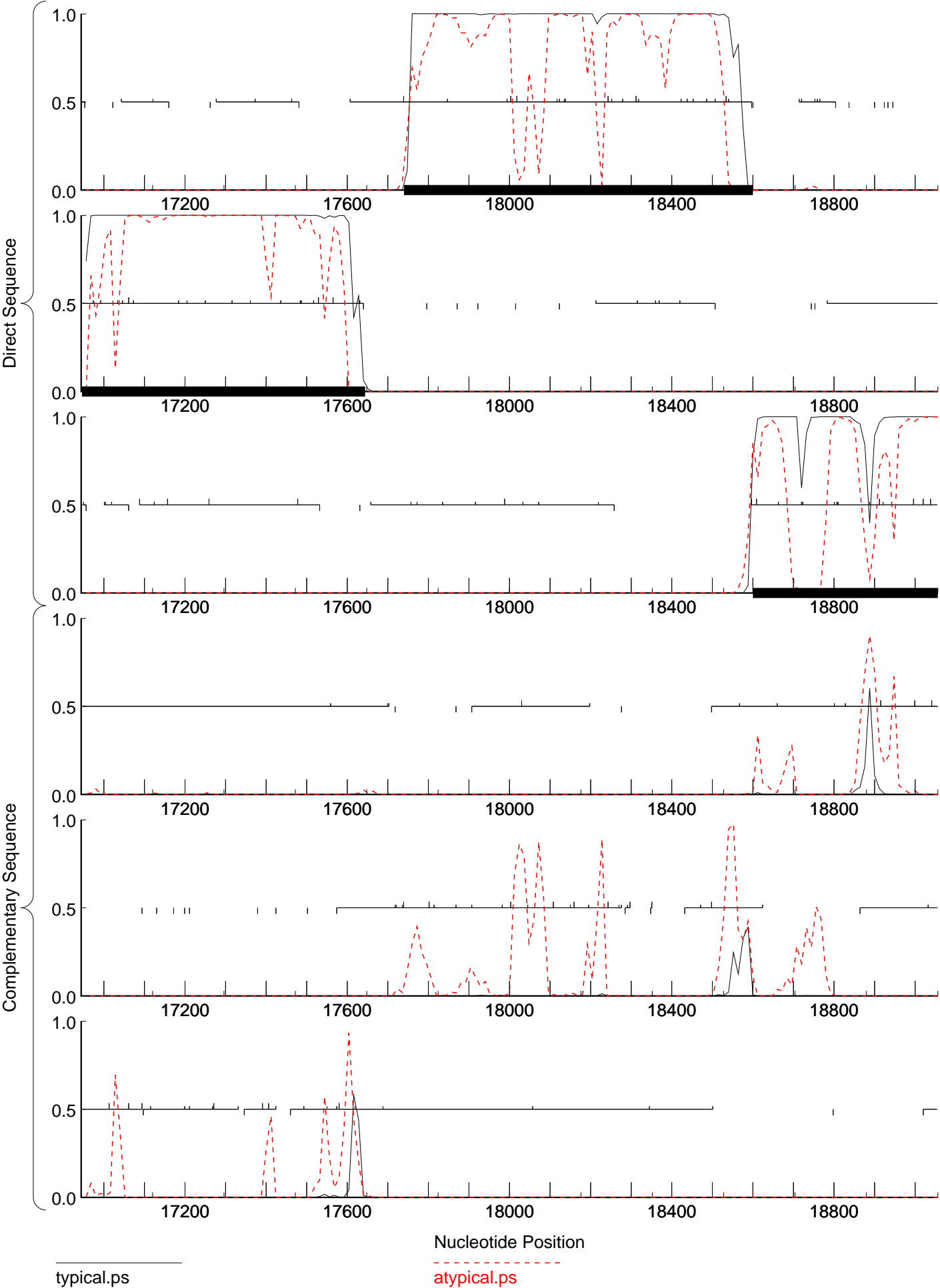
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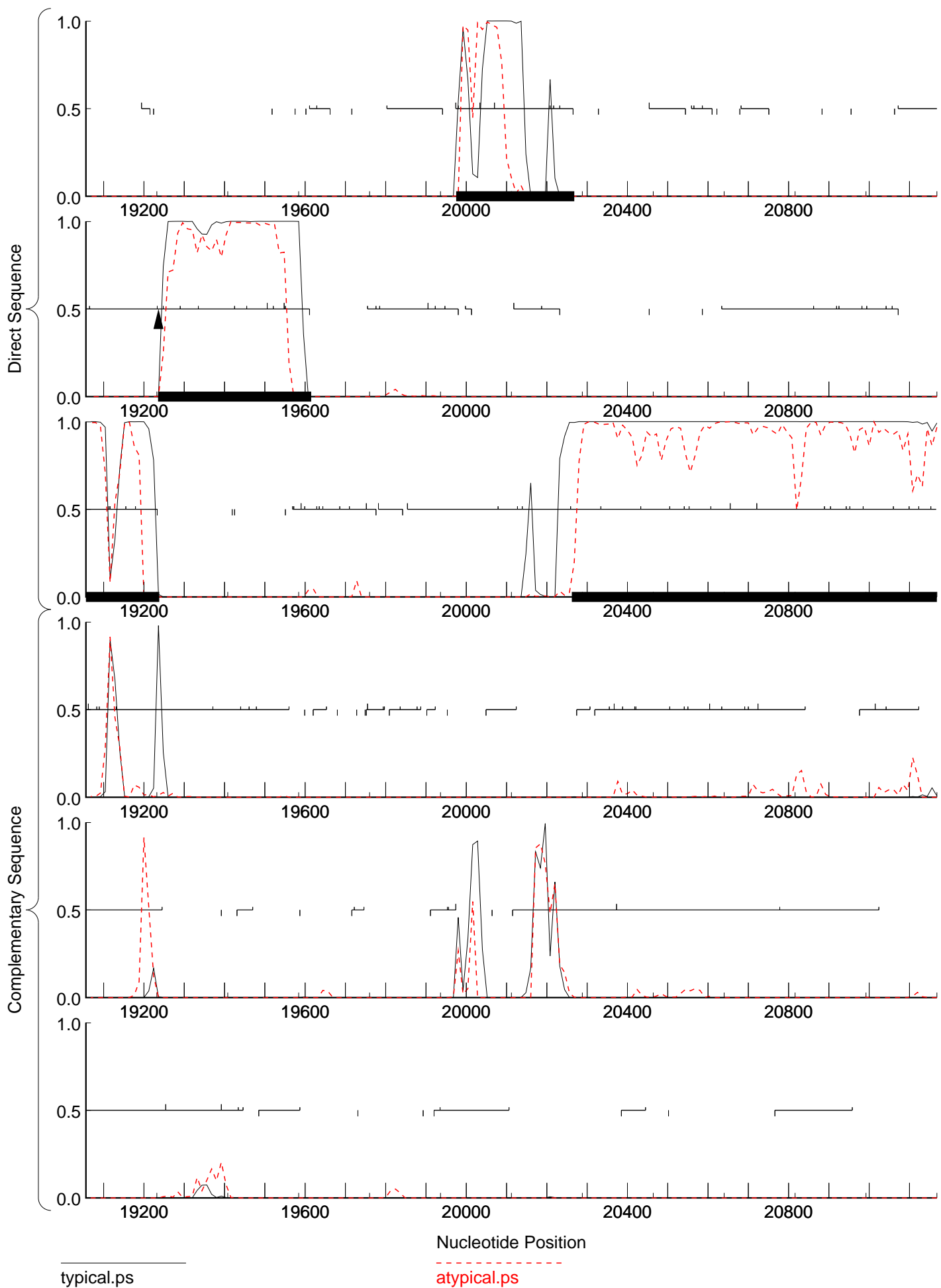


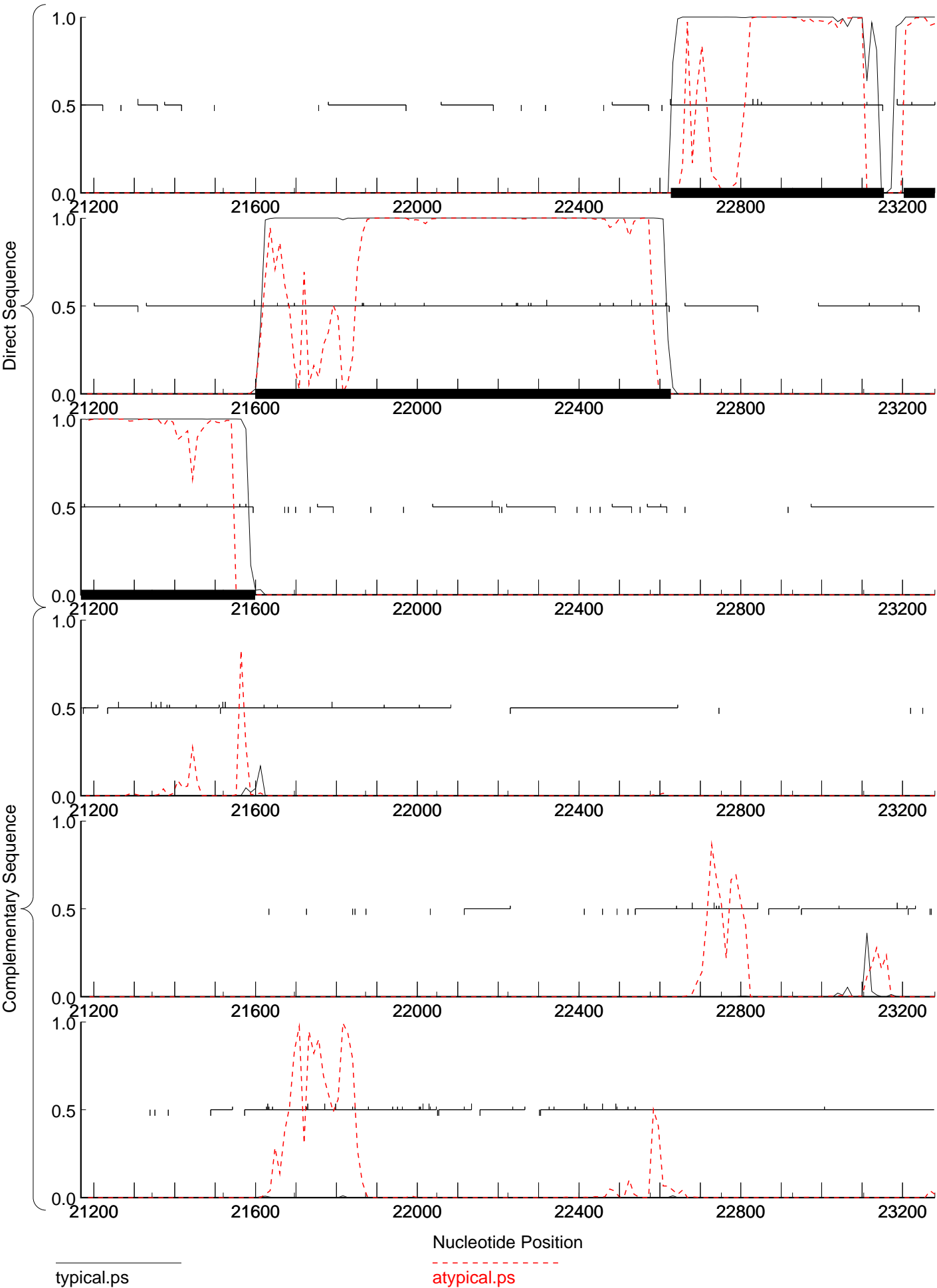






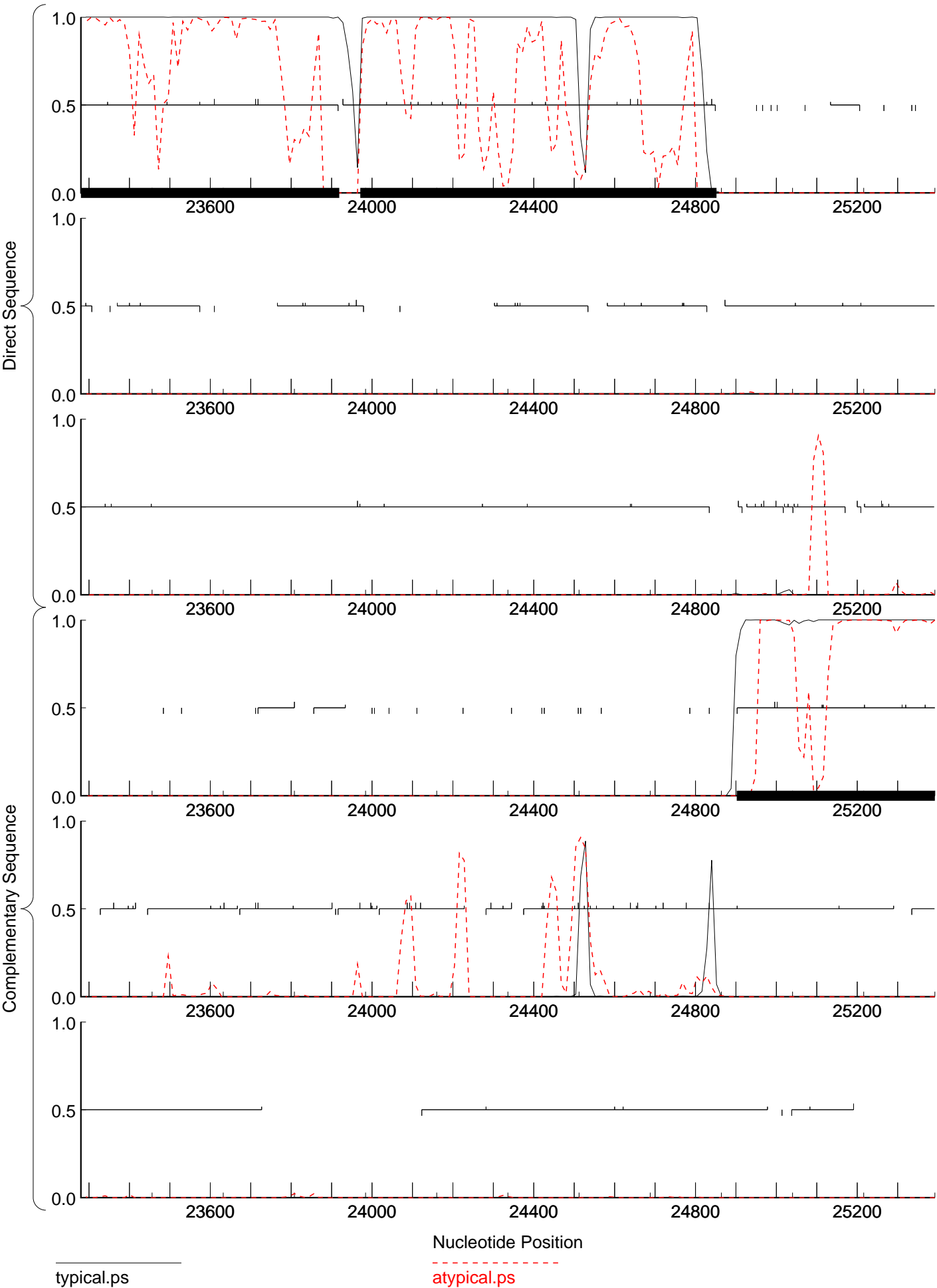


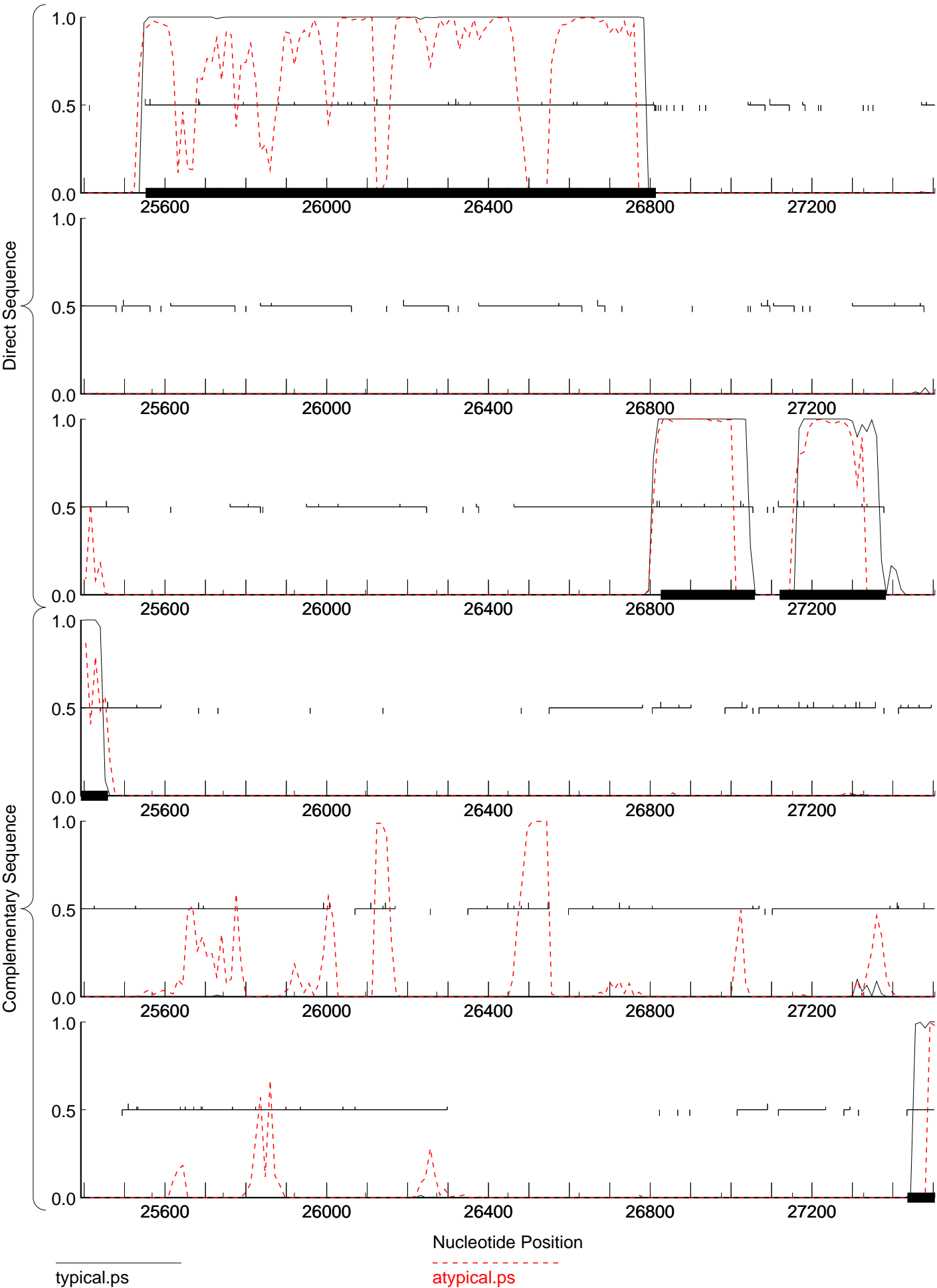




typical.ps

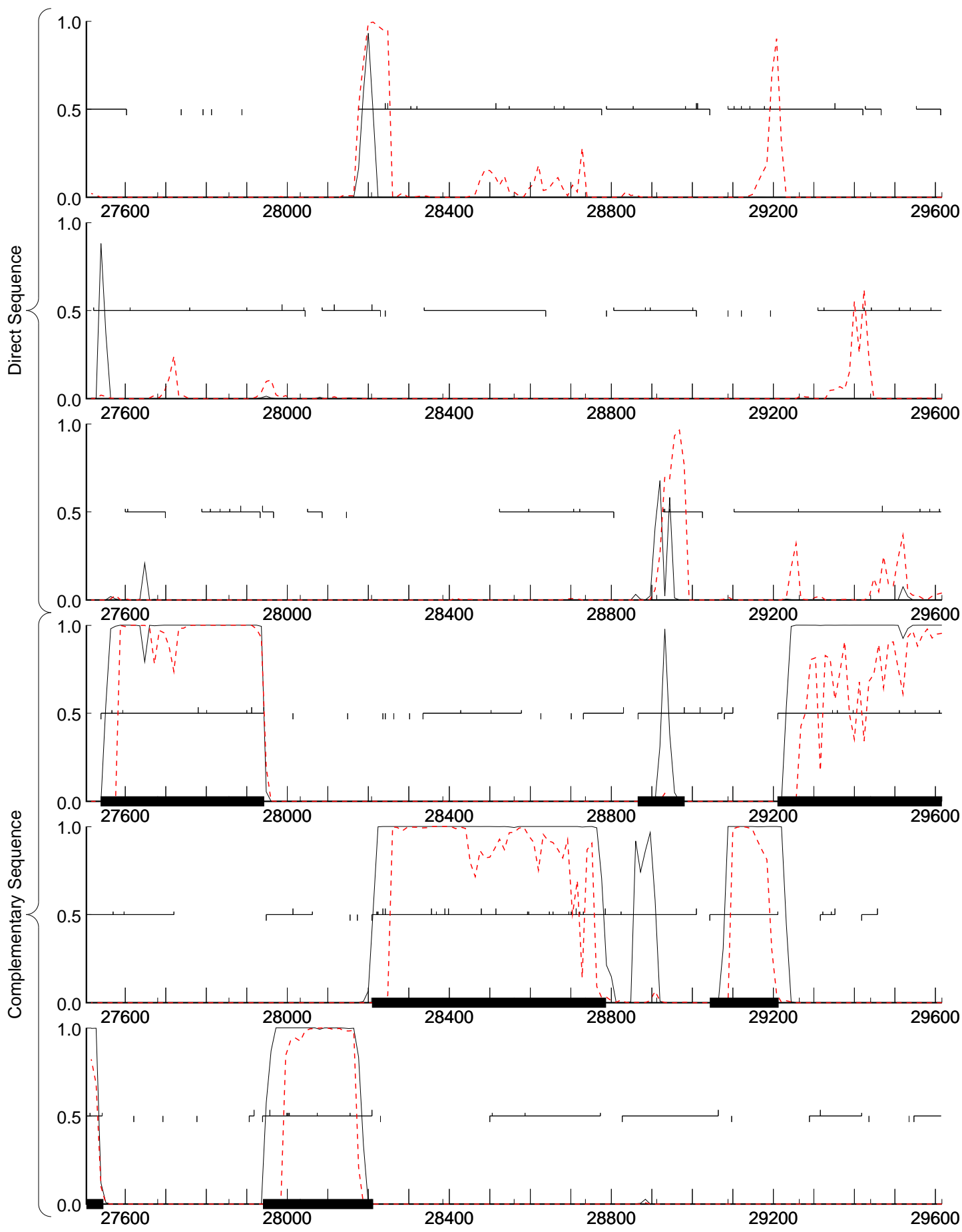
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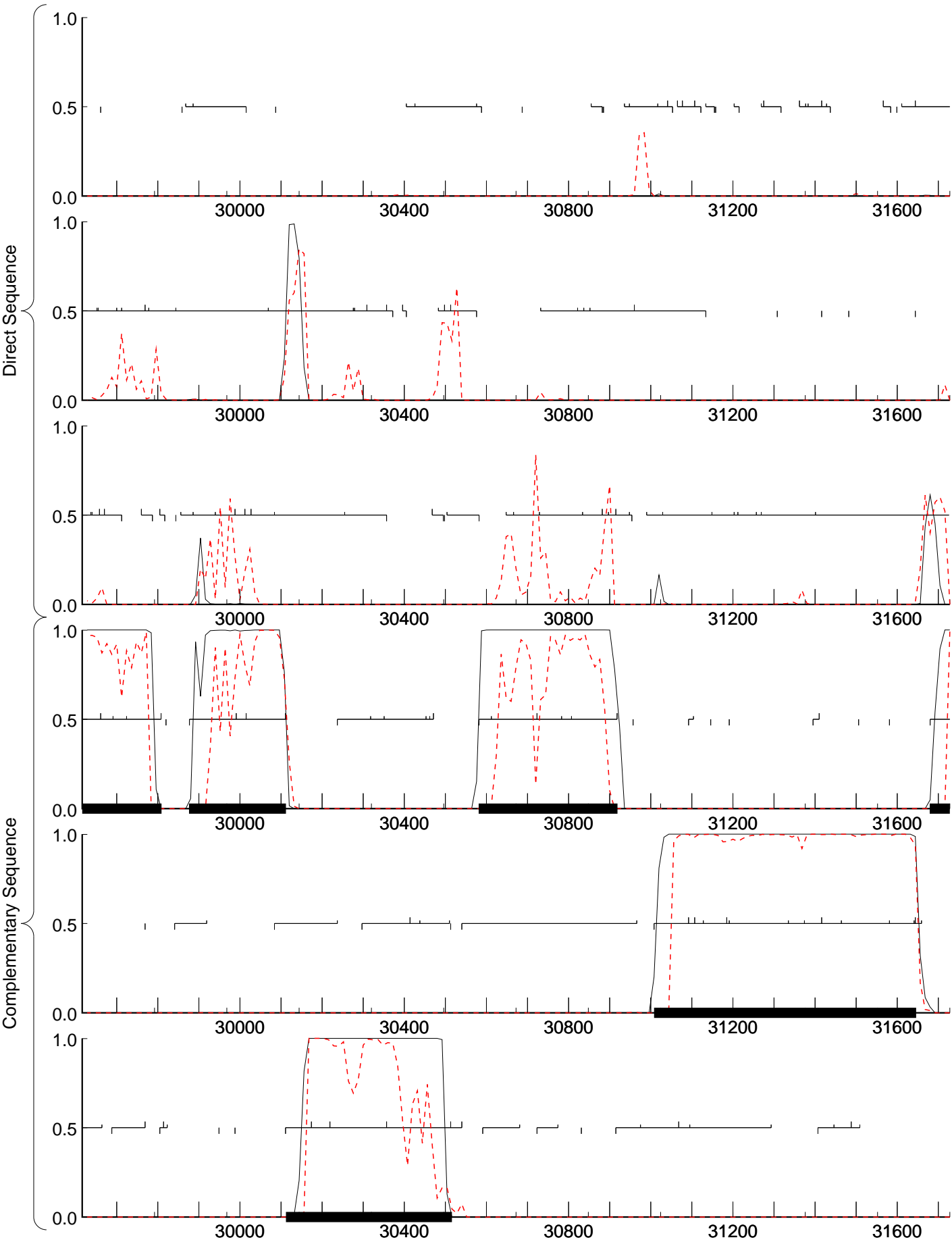
typical.ps

atypical.ps

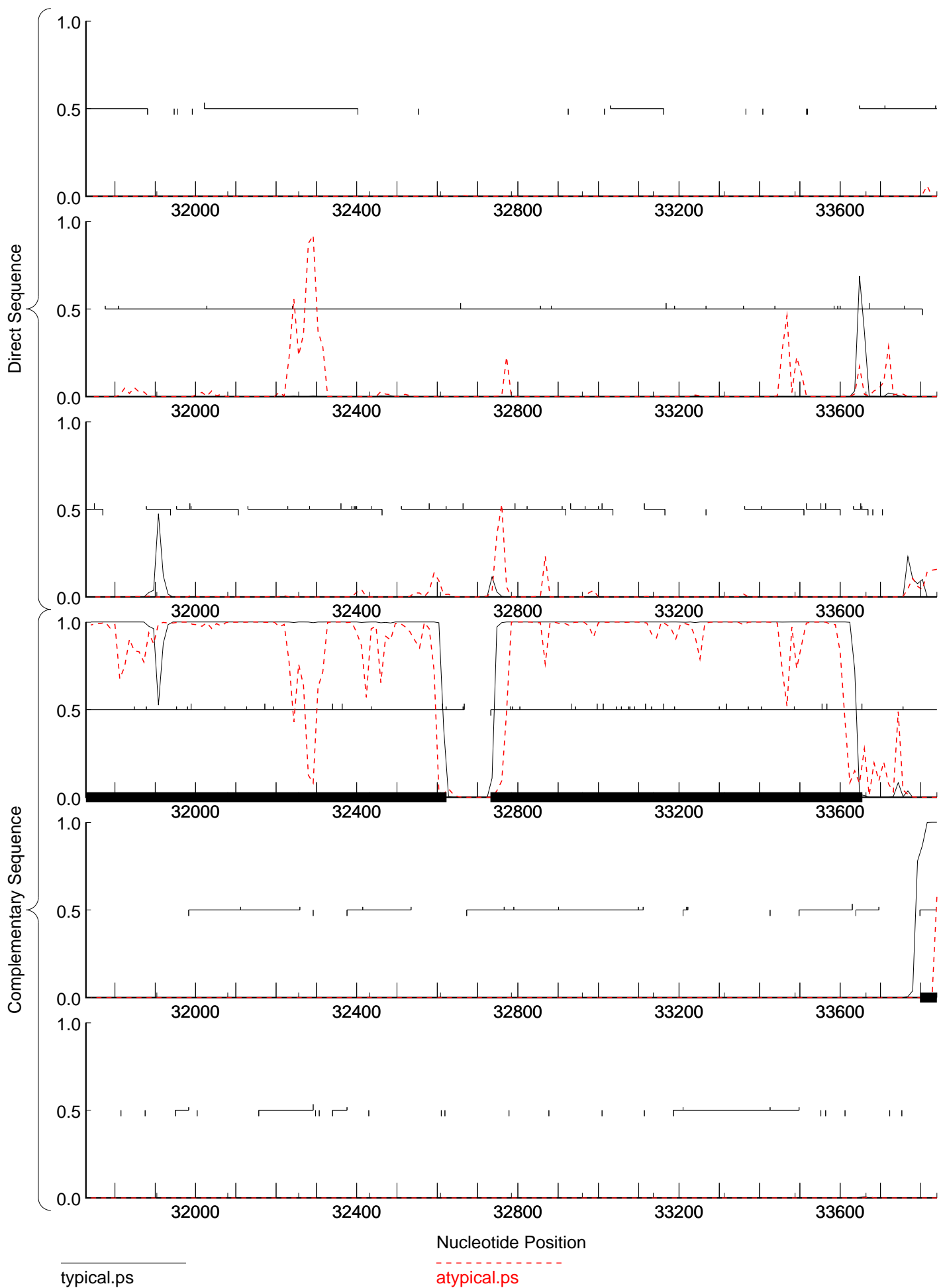


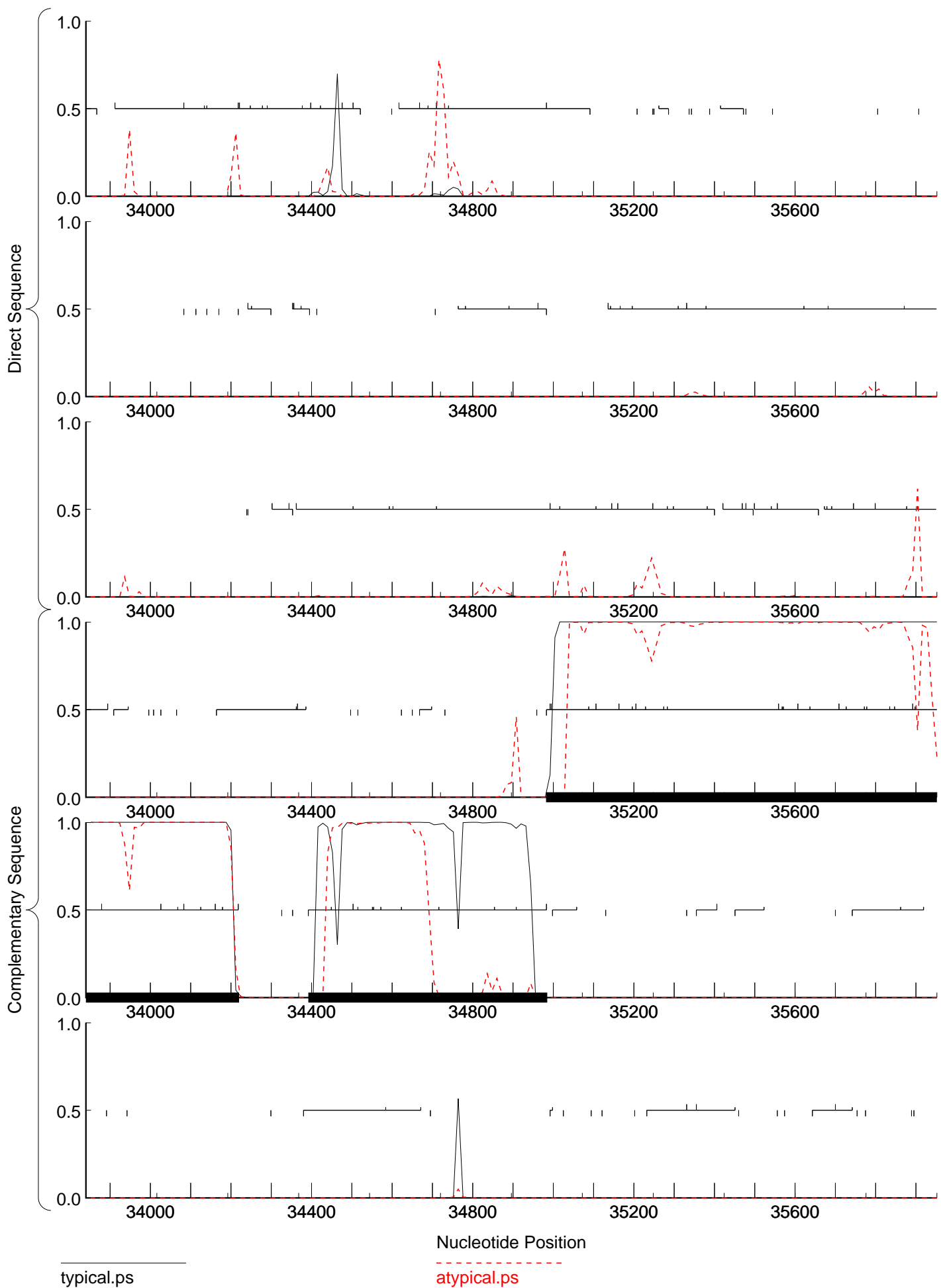
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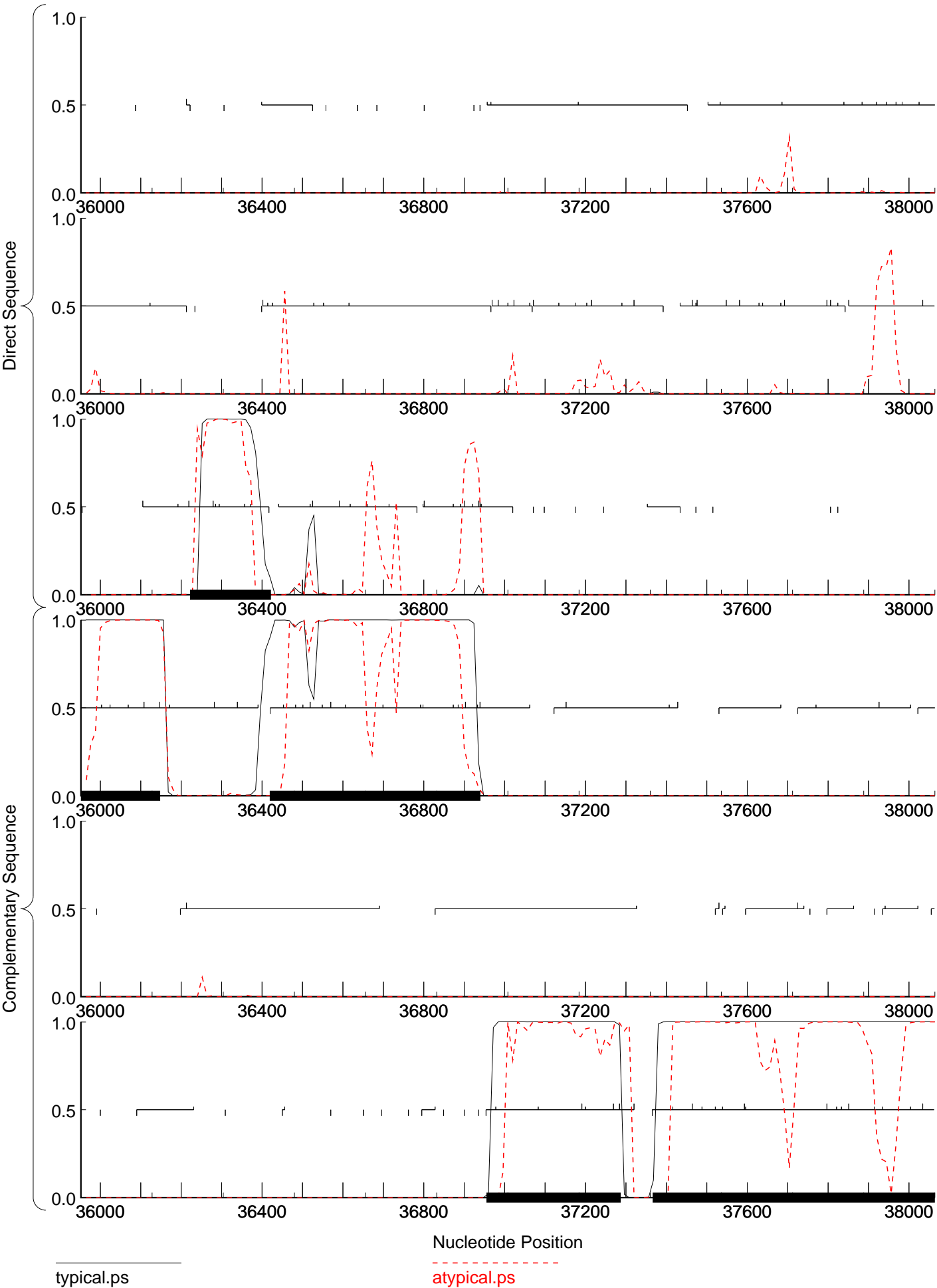
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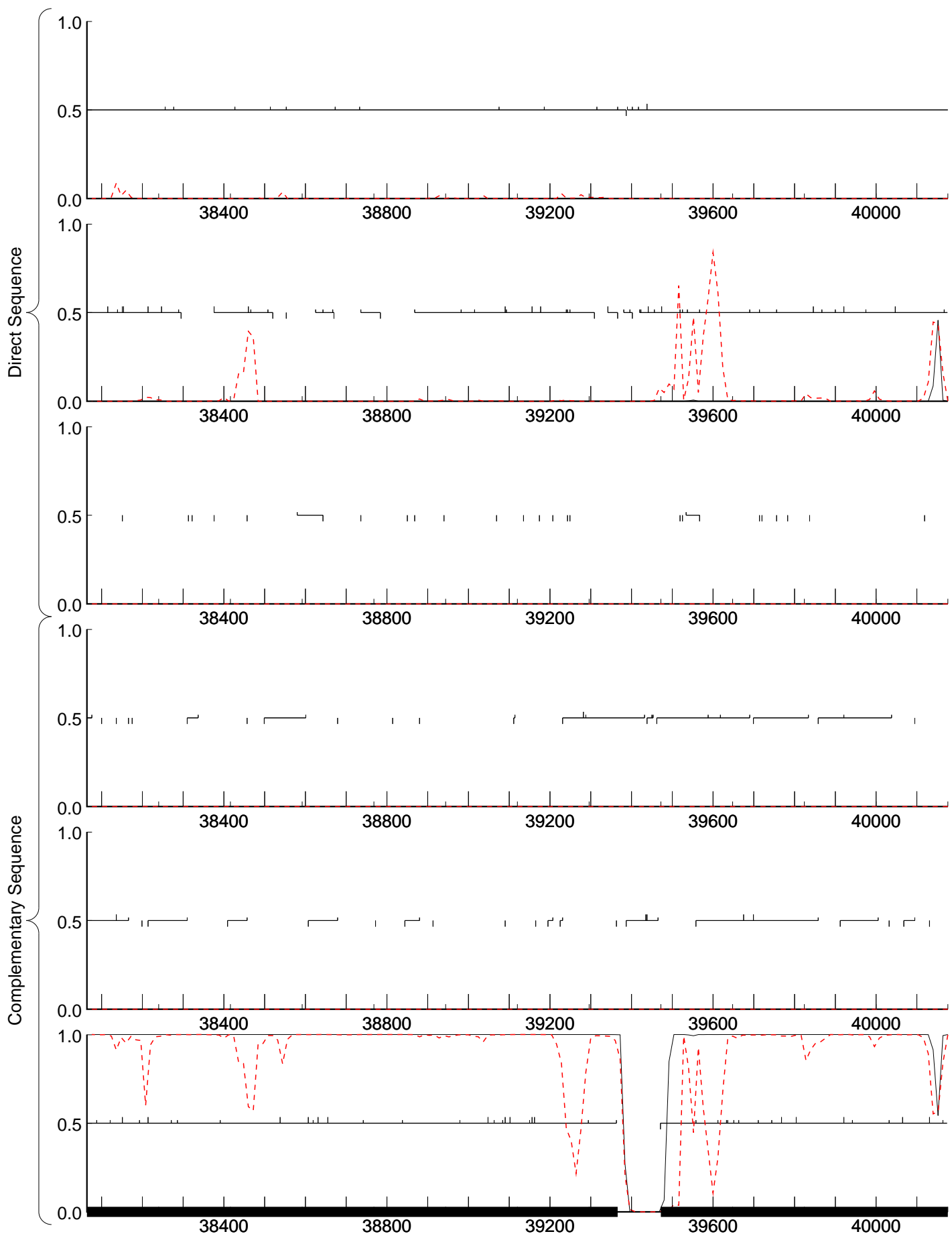


typical.ps
atypical.ps





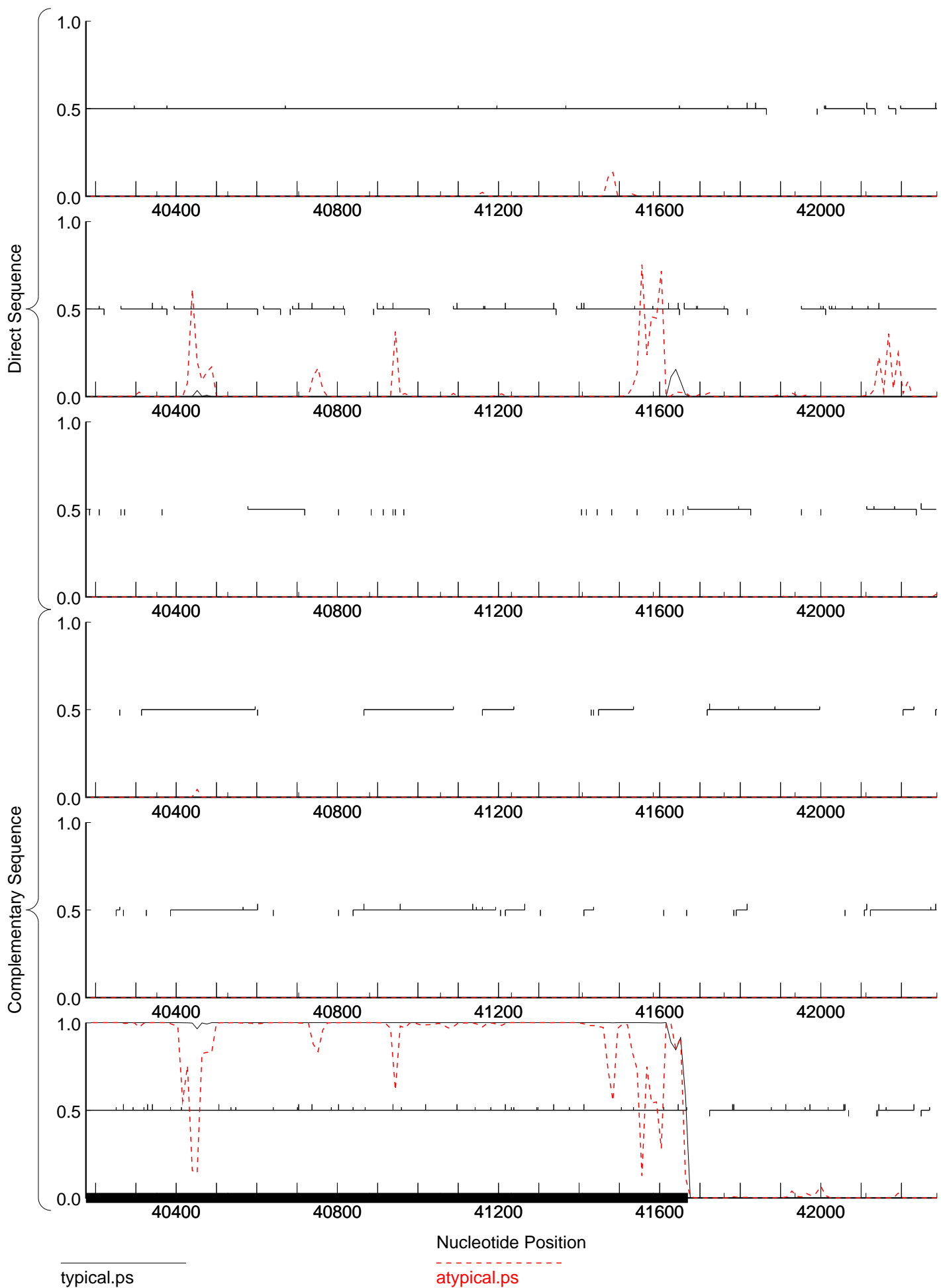


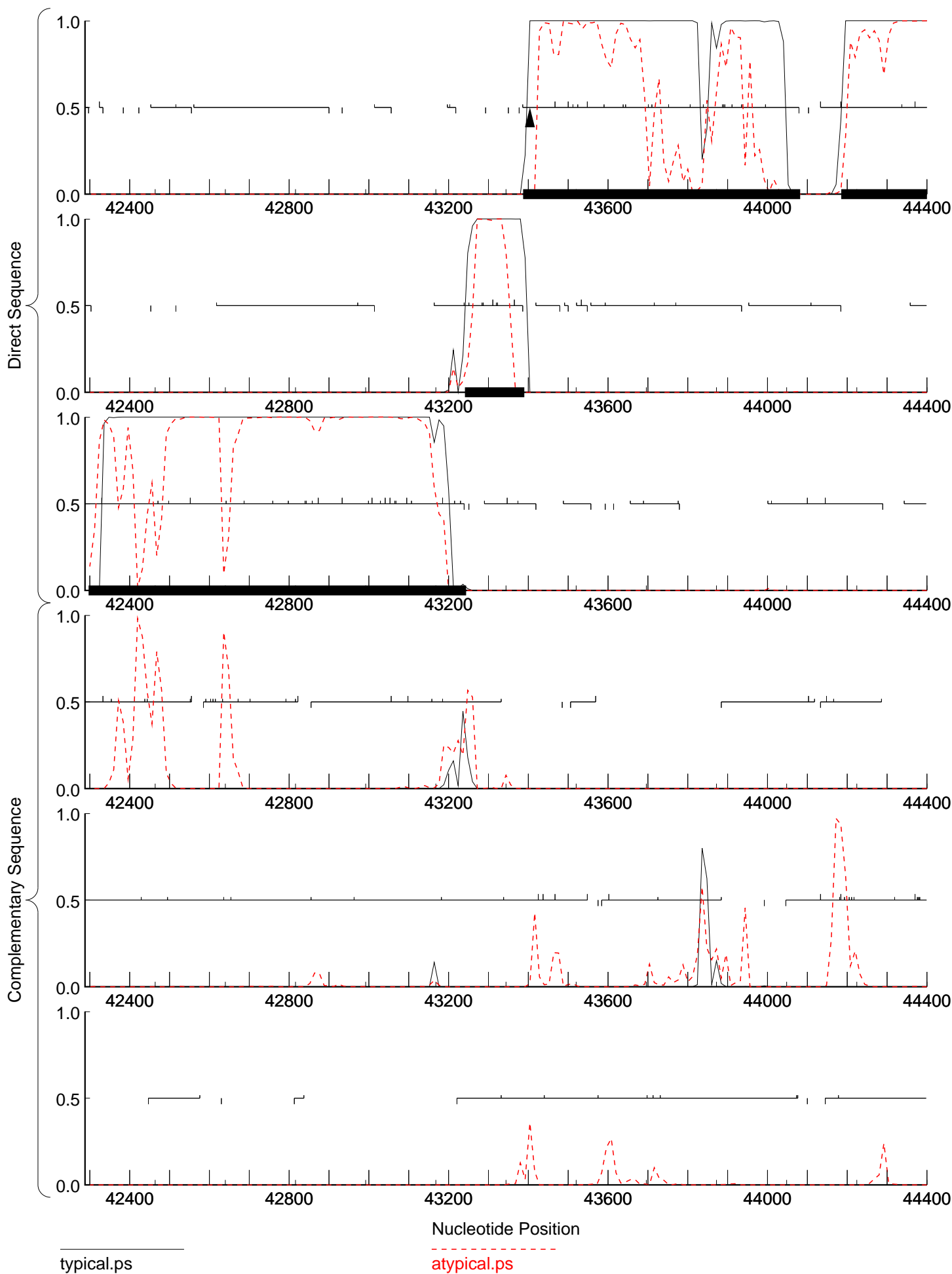


typical.ps

Nucleotide Position

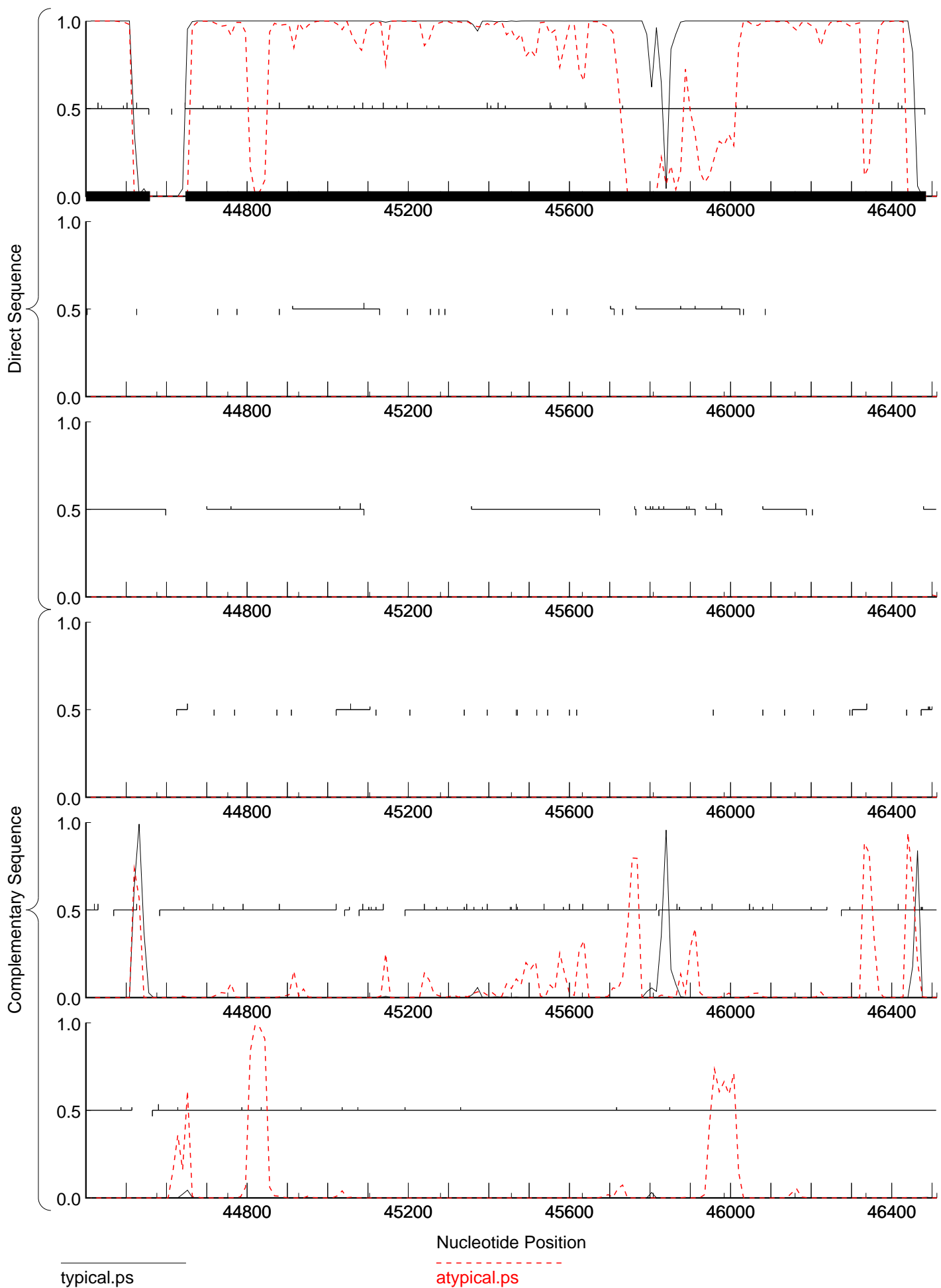
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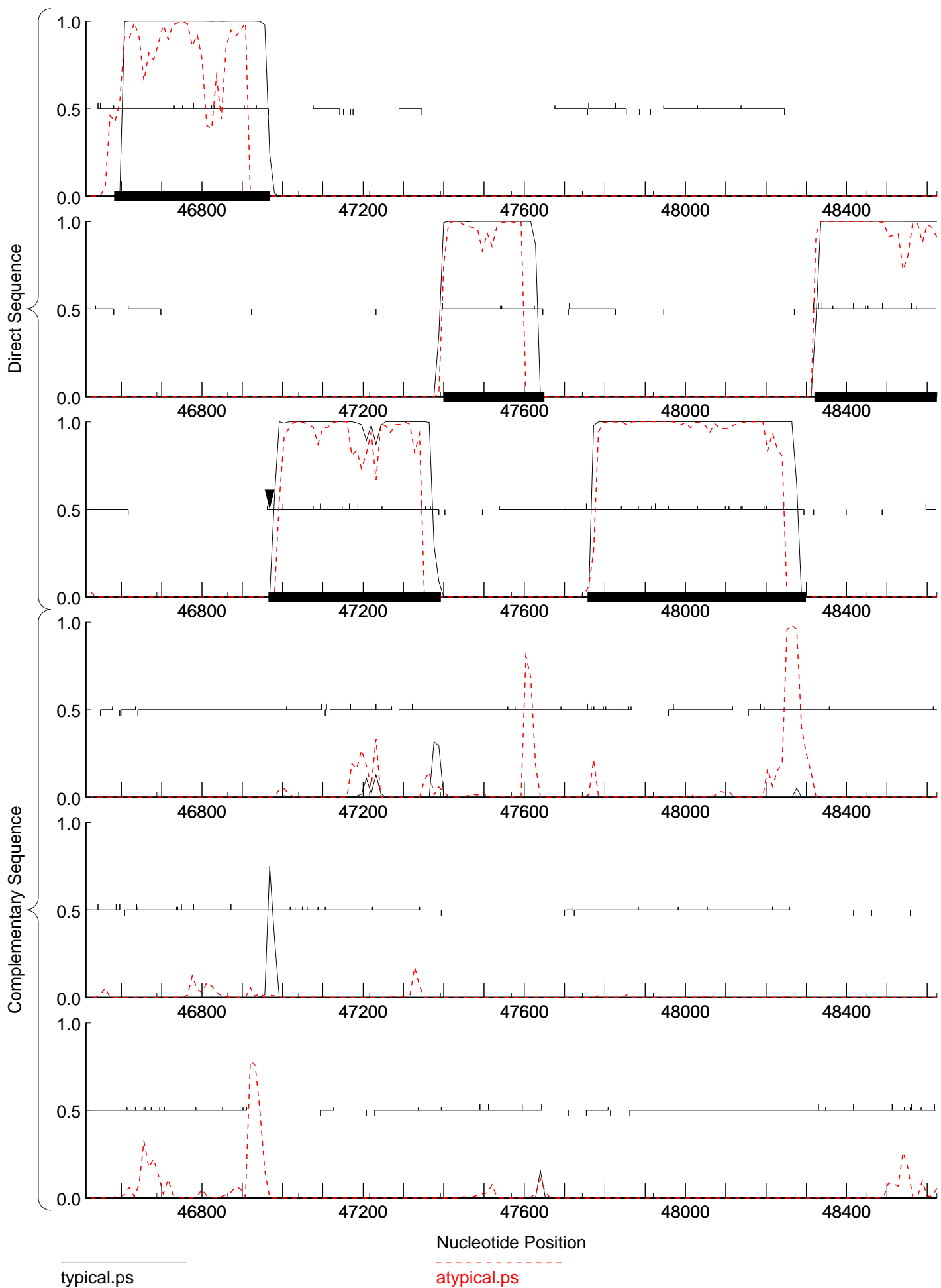
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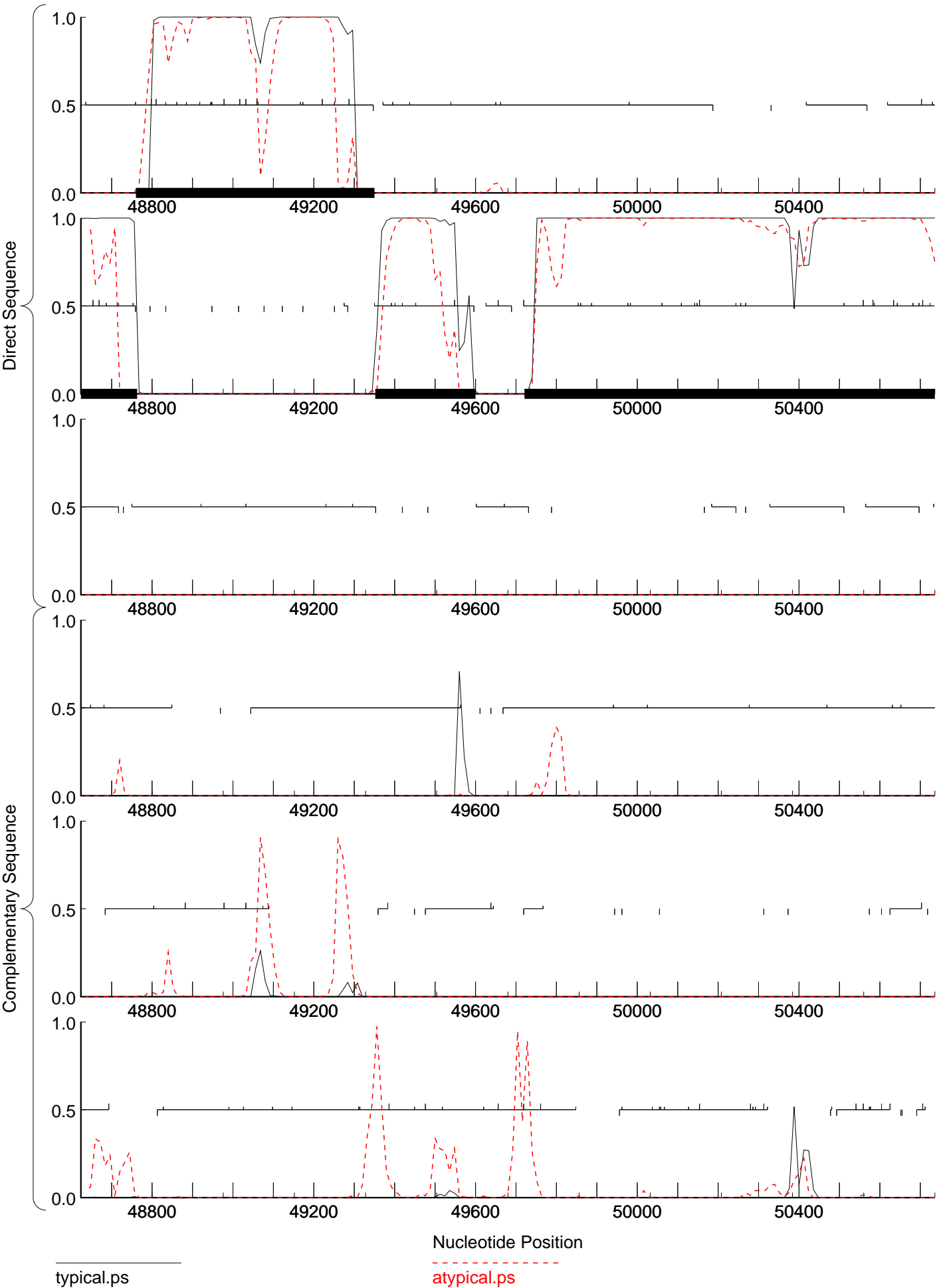
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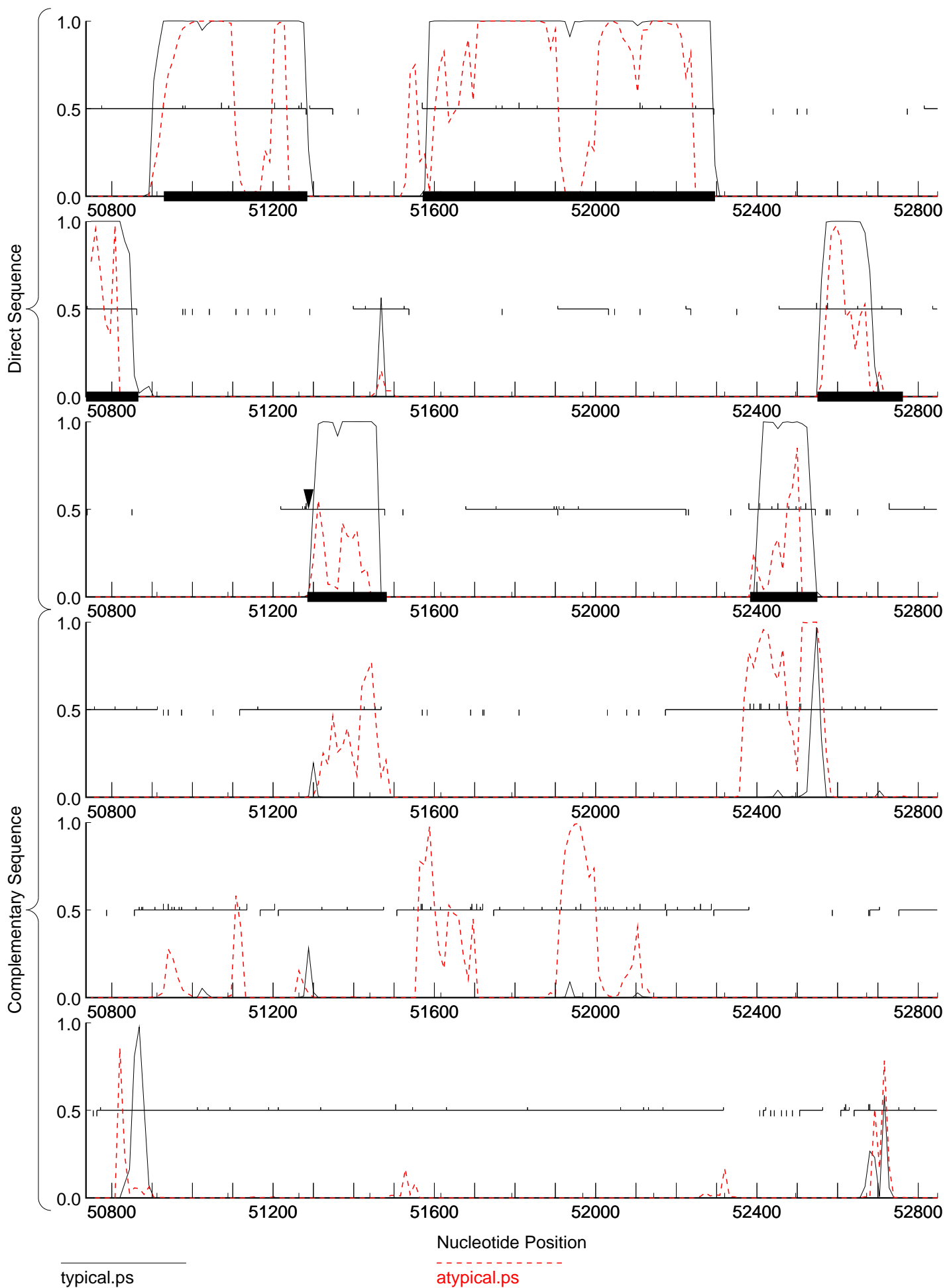


typical.ps

atypical.ps







typical.ps

atypical.ps

