

GeneMark

Version 2.5p (09.08.06)

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PROGRAM INFORMATION

Sequence : Microbacterium phage Hyperion complete squence, 61769 bp including 203 bp terminal repeat,
Analysis Date : 9/21/17 at 2:51:51
Pages : 31
Sequence Length : 61769 bp
GC Content : 67.01%

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,

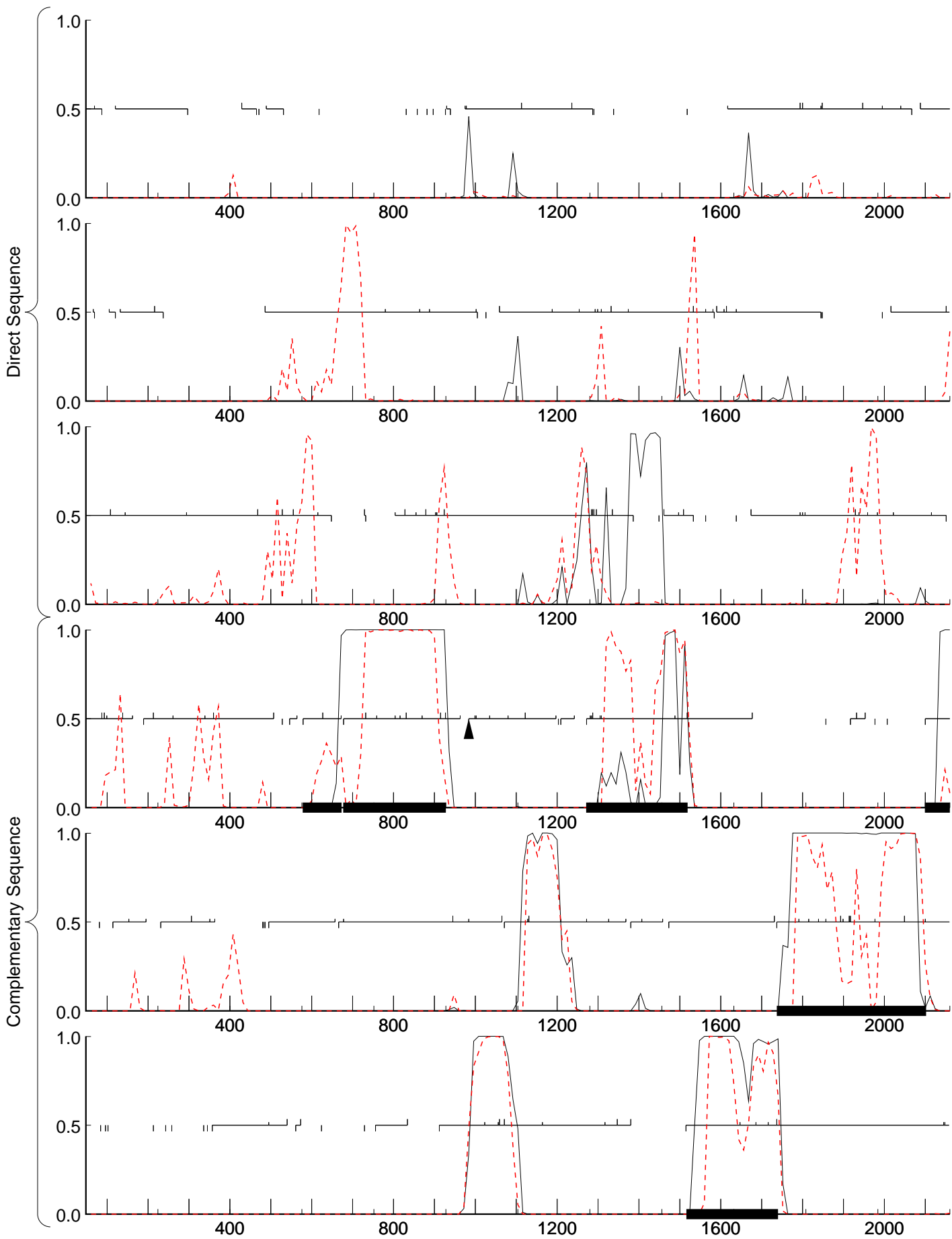
MATRIX INFORMATION

Matrix : -
Author : -
Order : 4

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

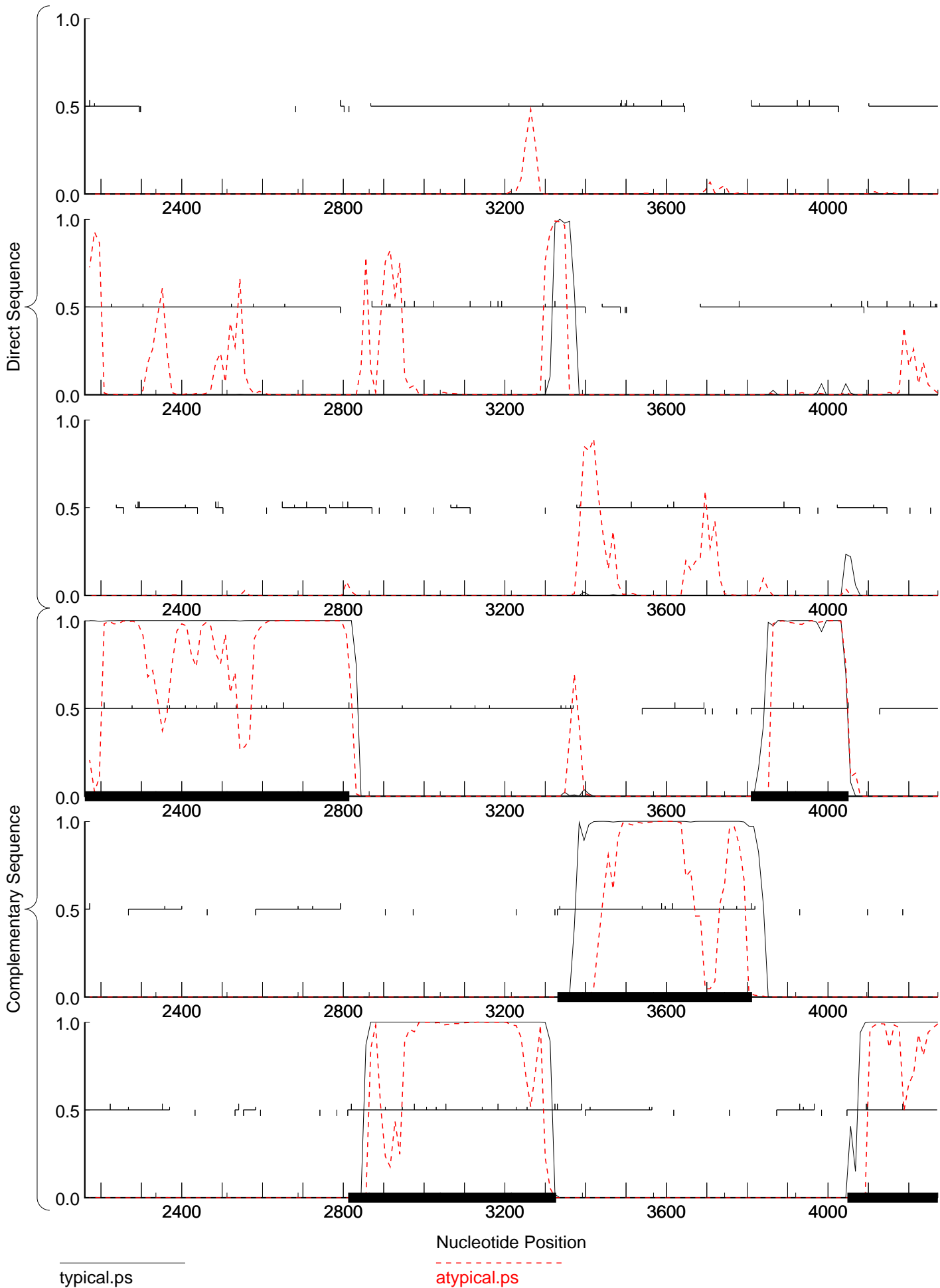
Matrix notes & comments

Training set derived by GeneMarkS, 4.30 October 2014
Thu Sep 21 02:51:51 2017



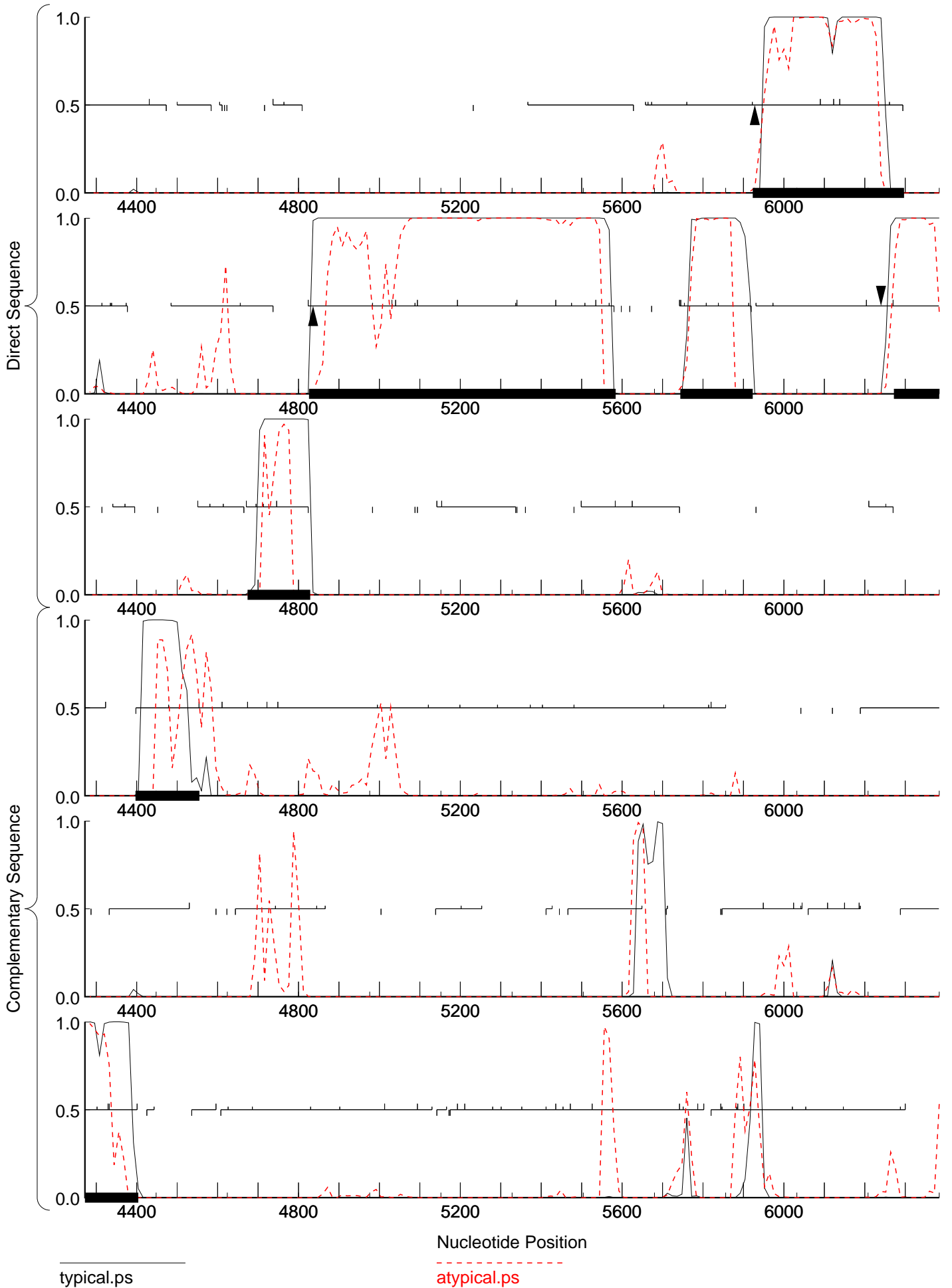
typical.ps

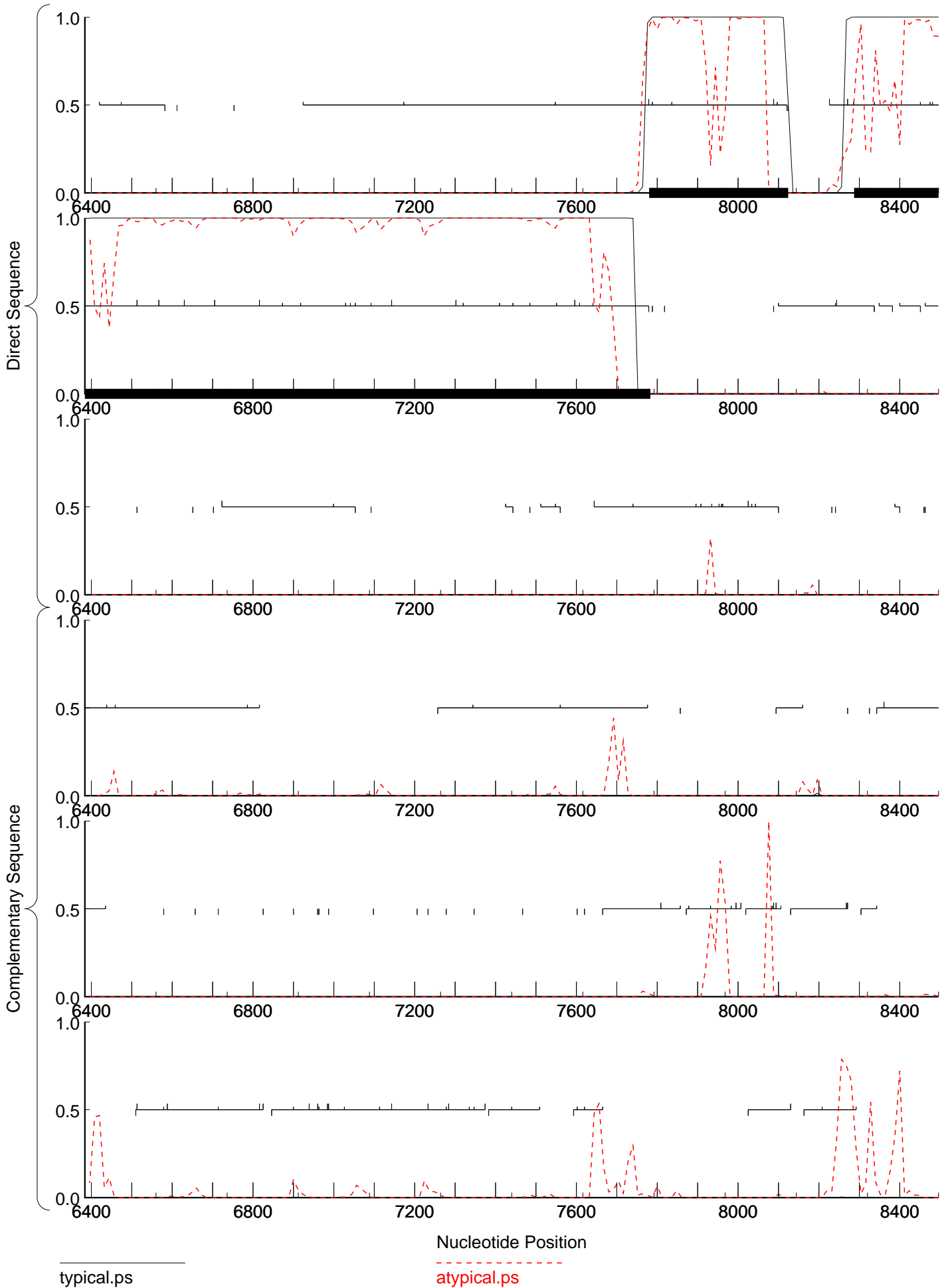
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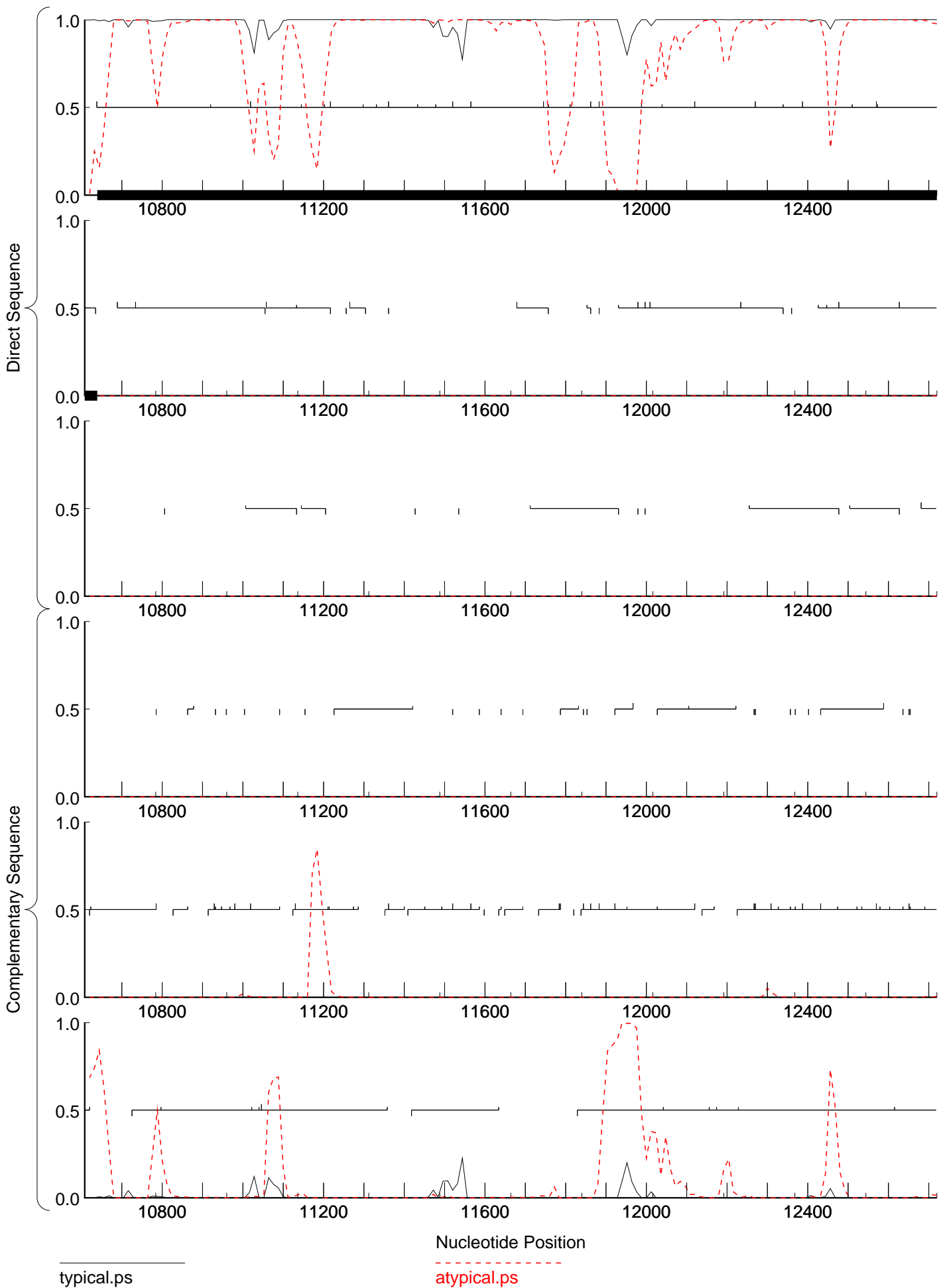


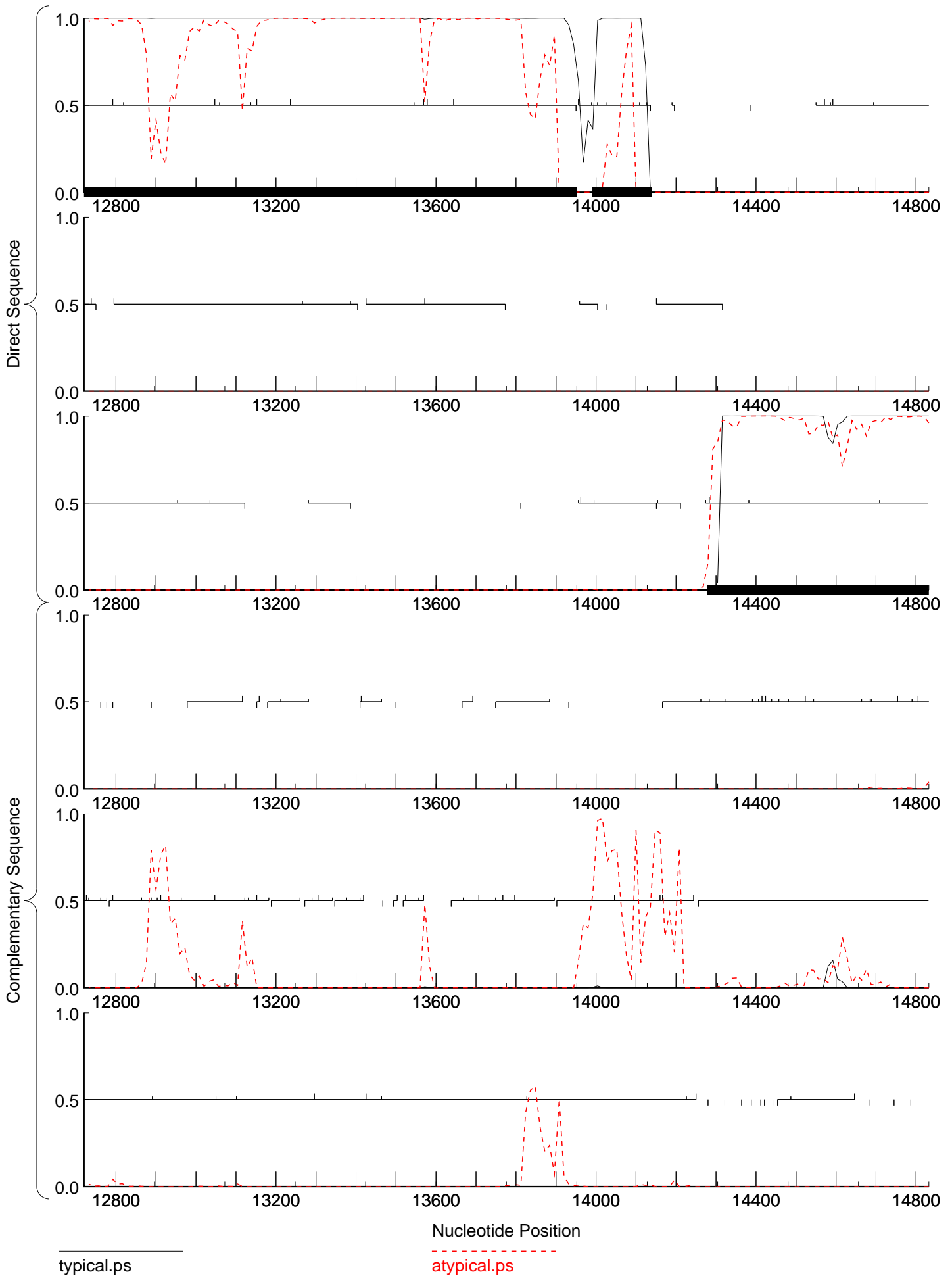
typical.ps

atypical.ps



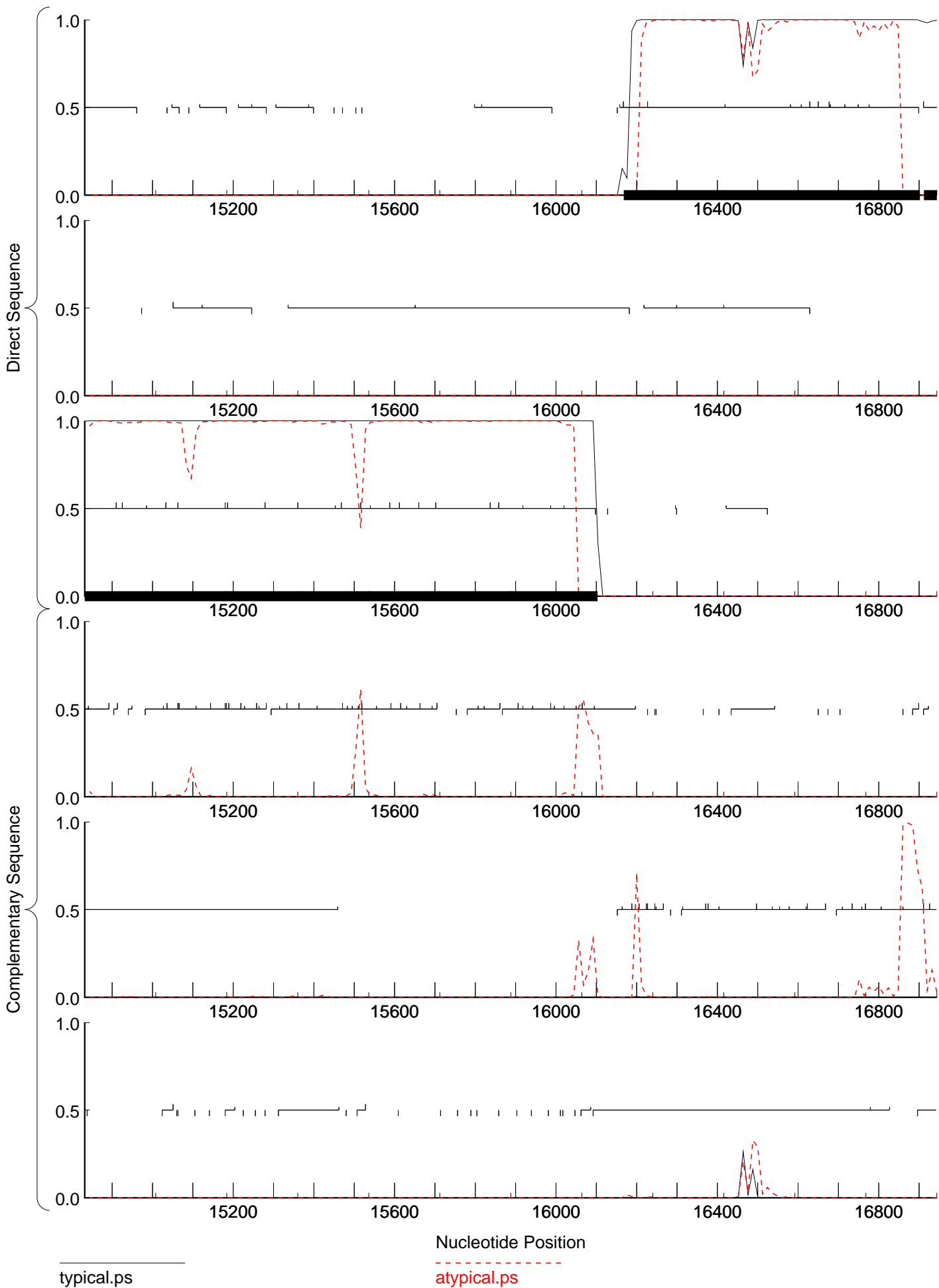


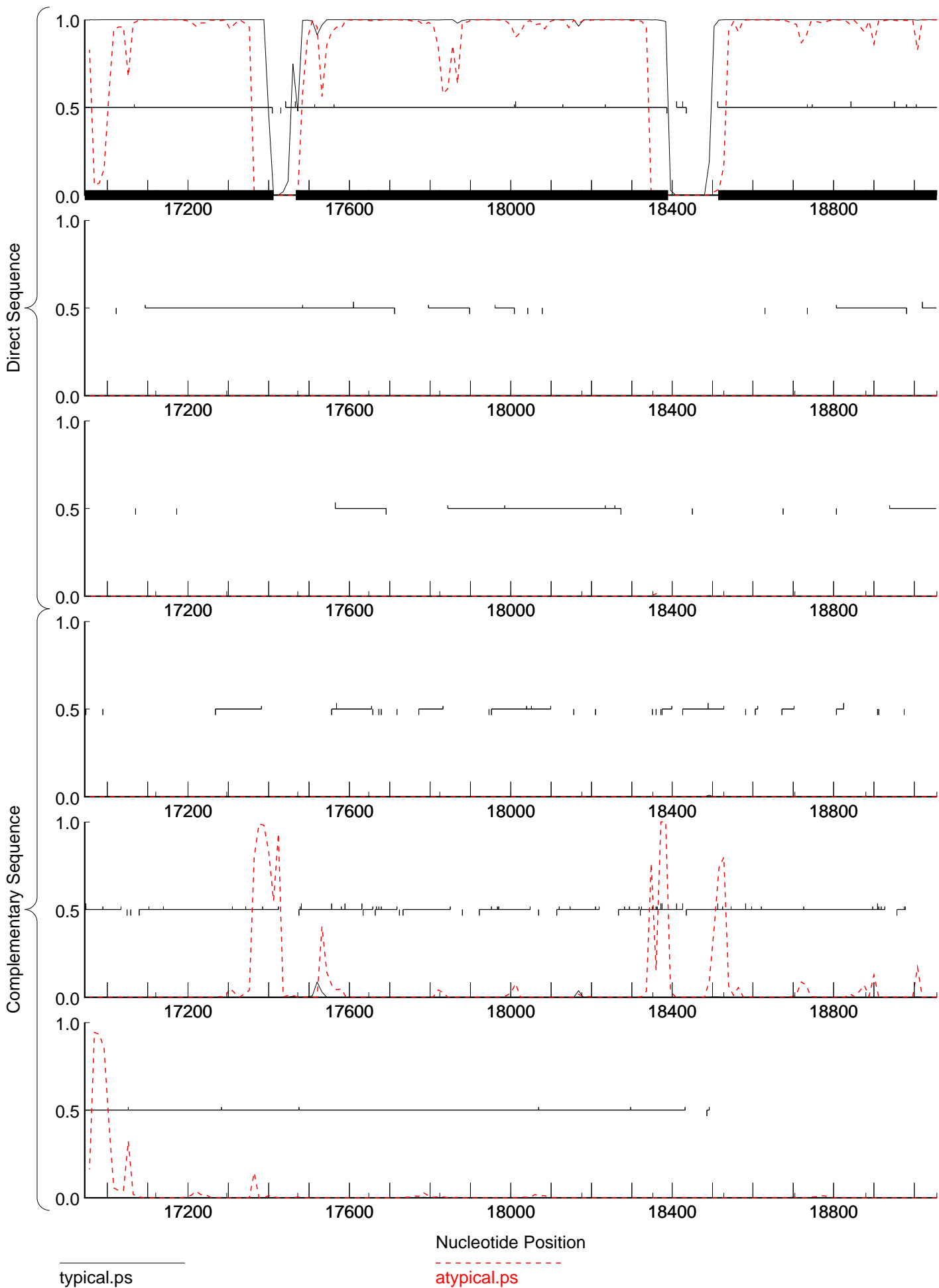


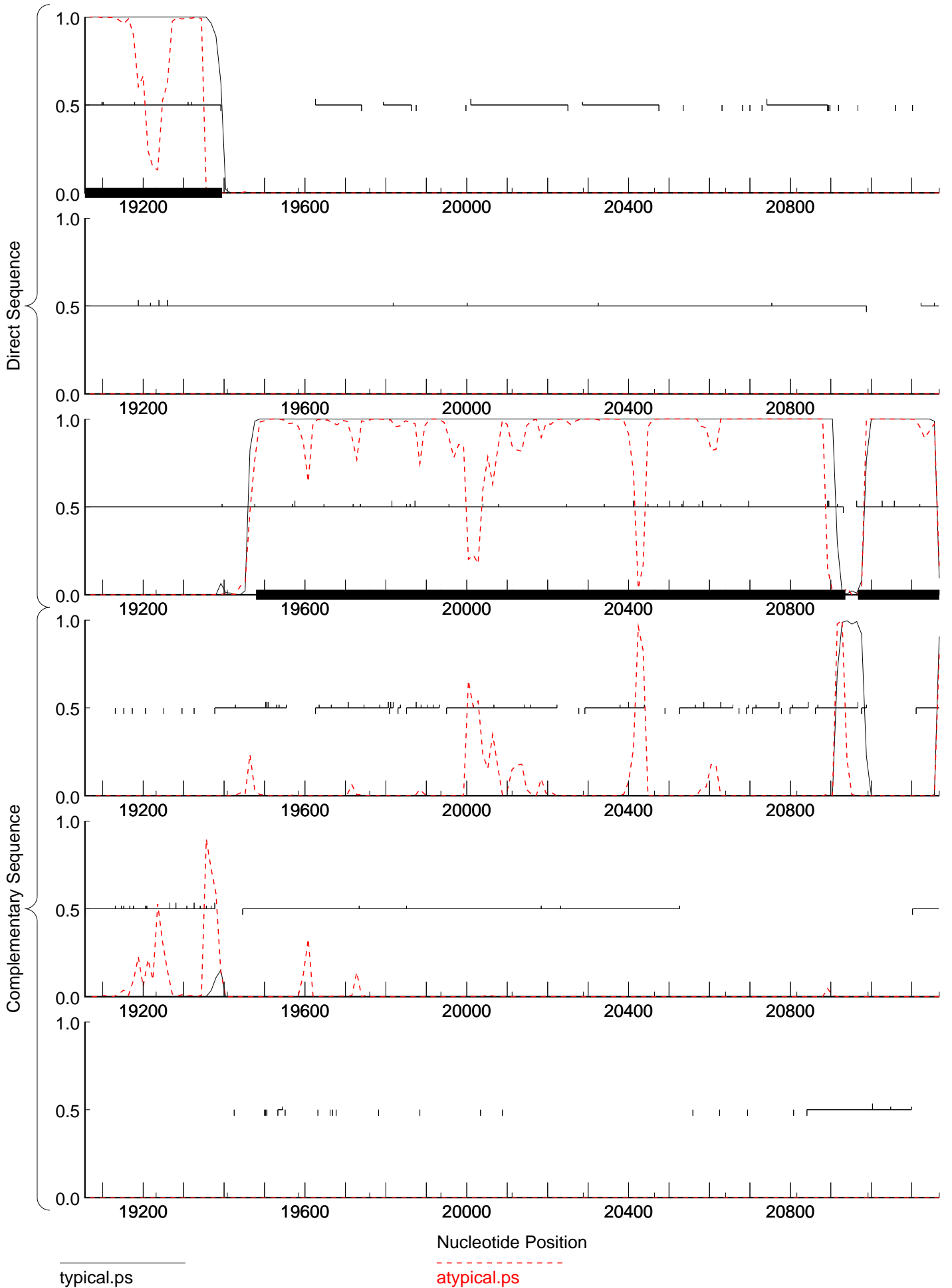


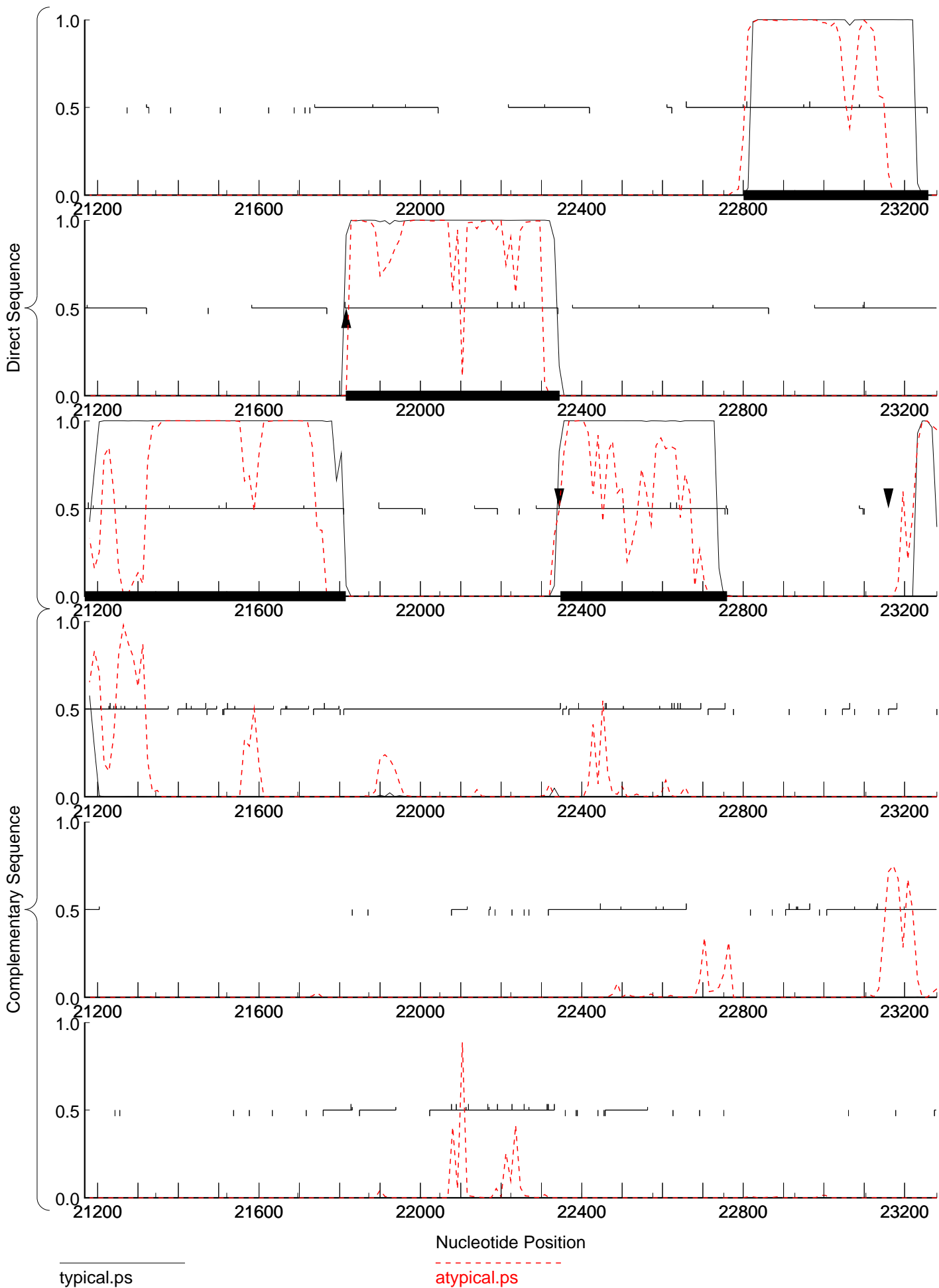
typical.ps

atypical.ps



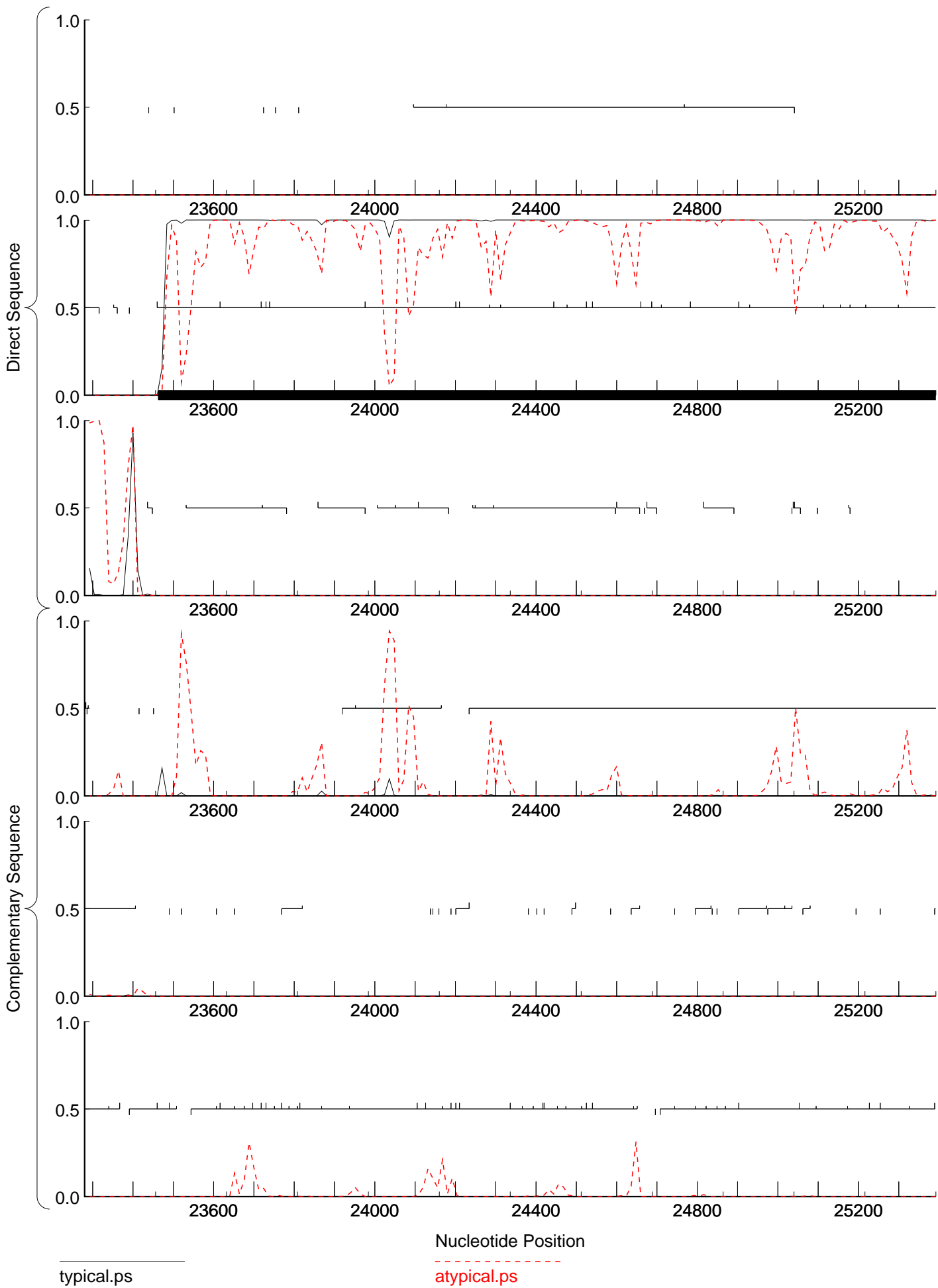


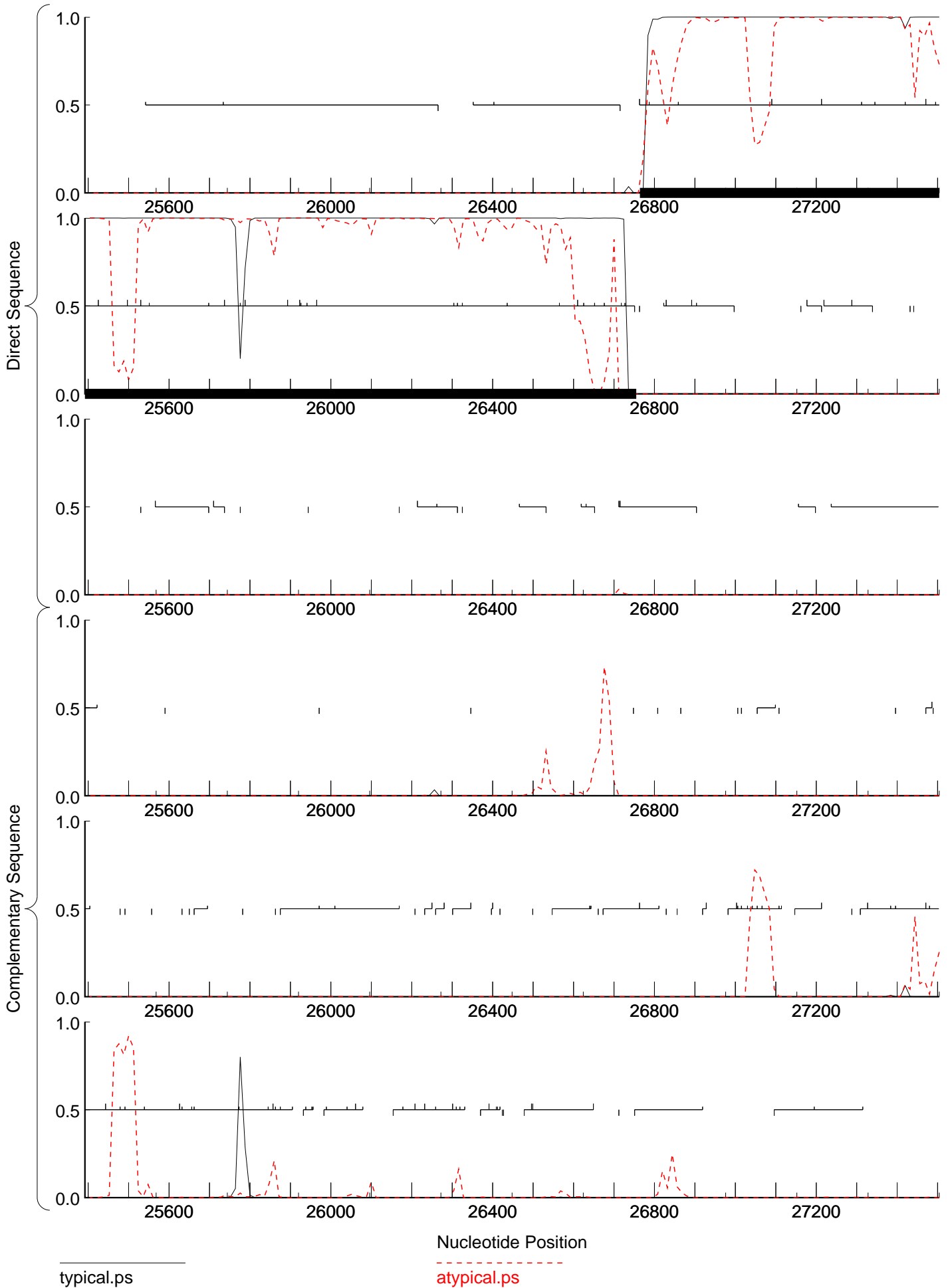


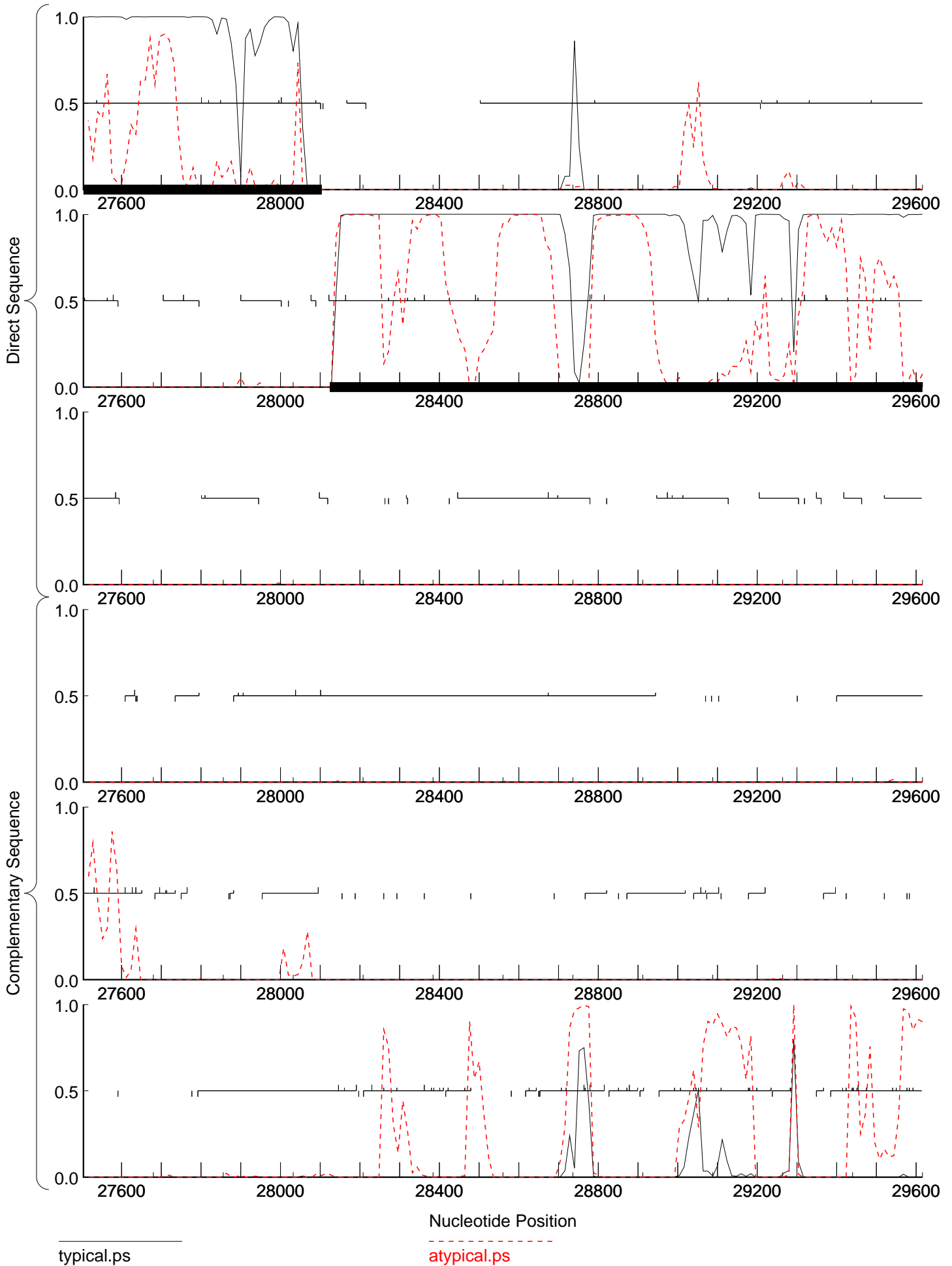


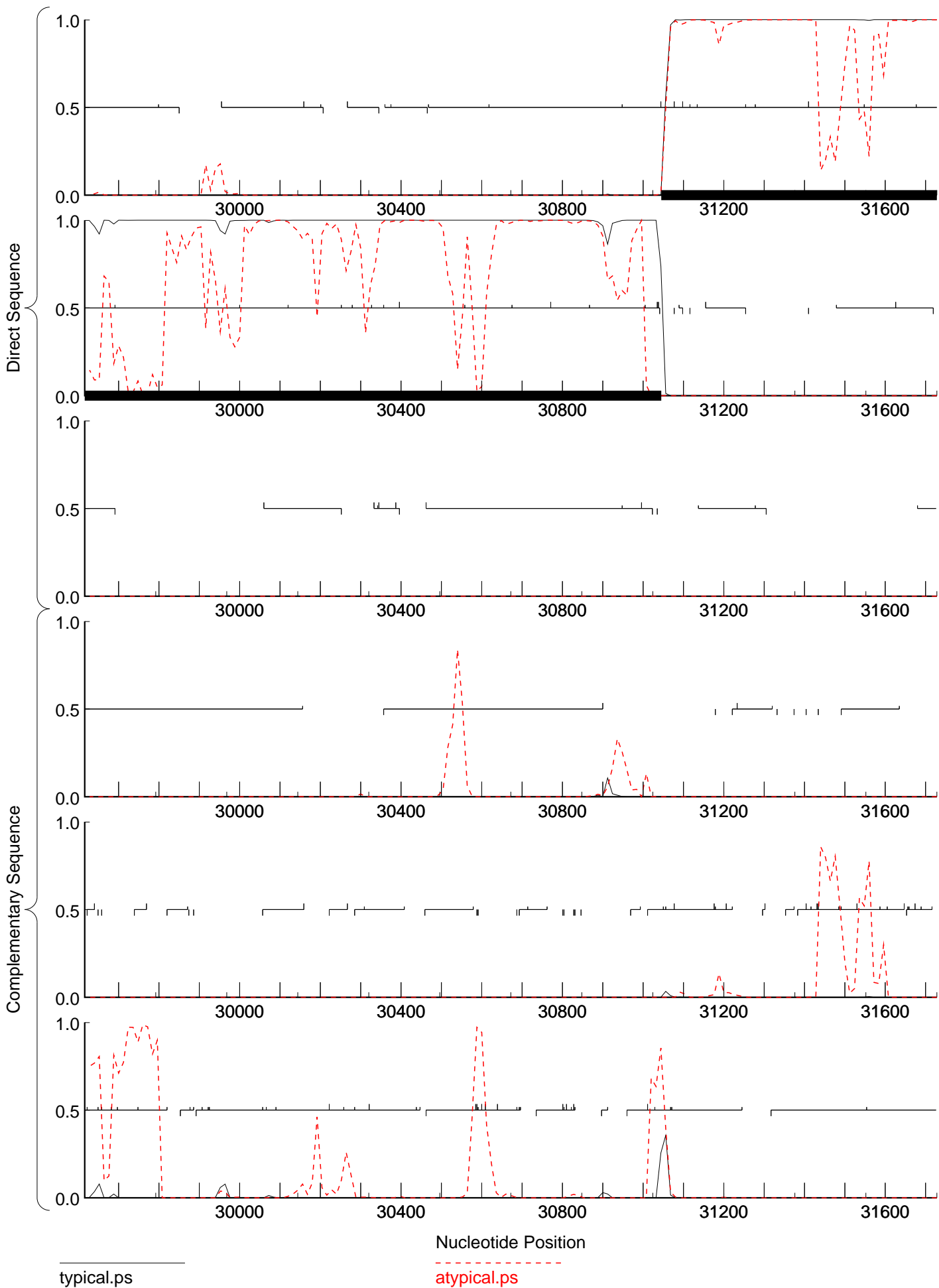
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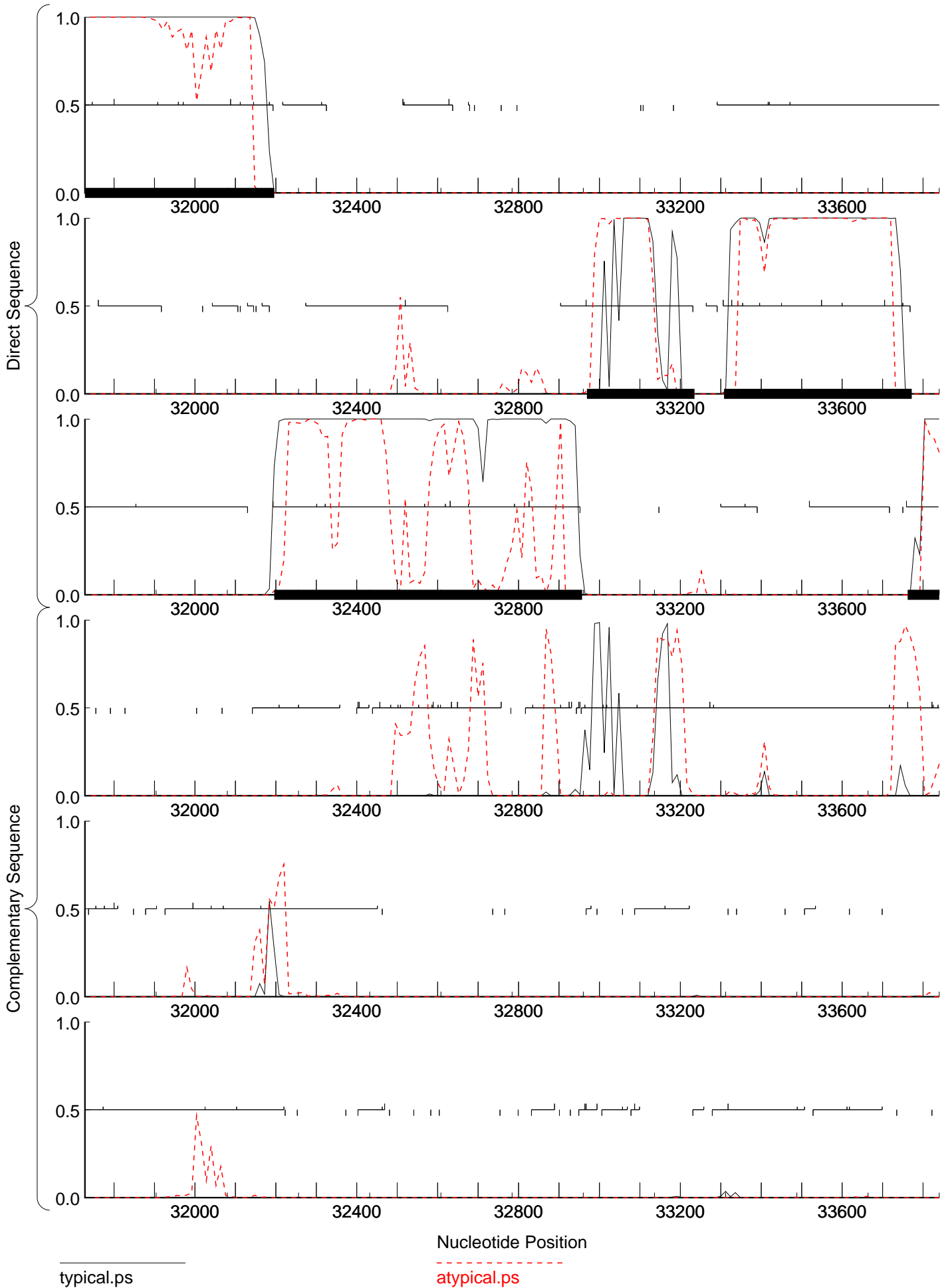
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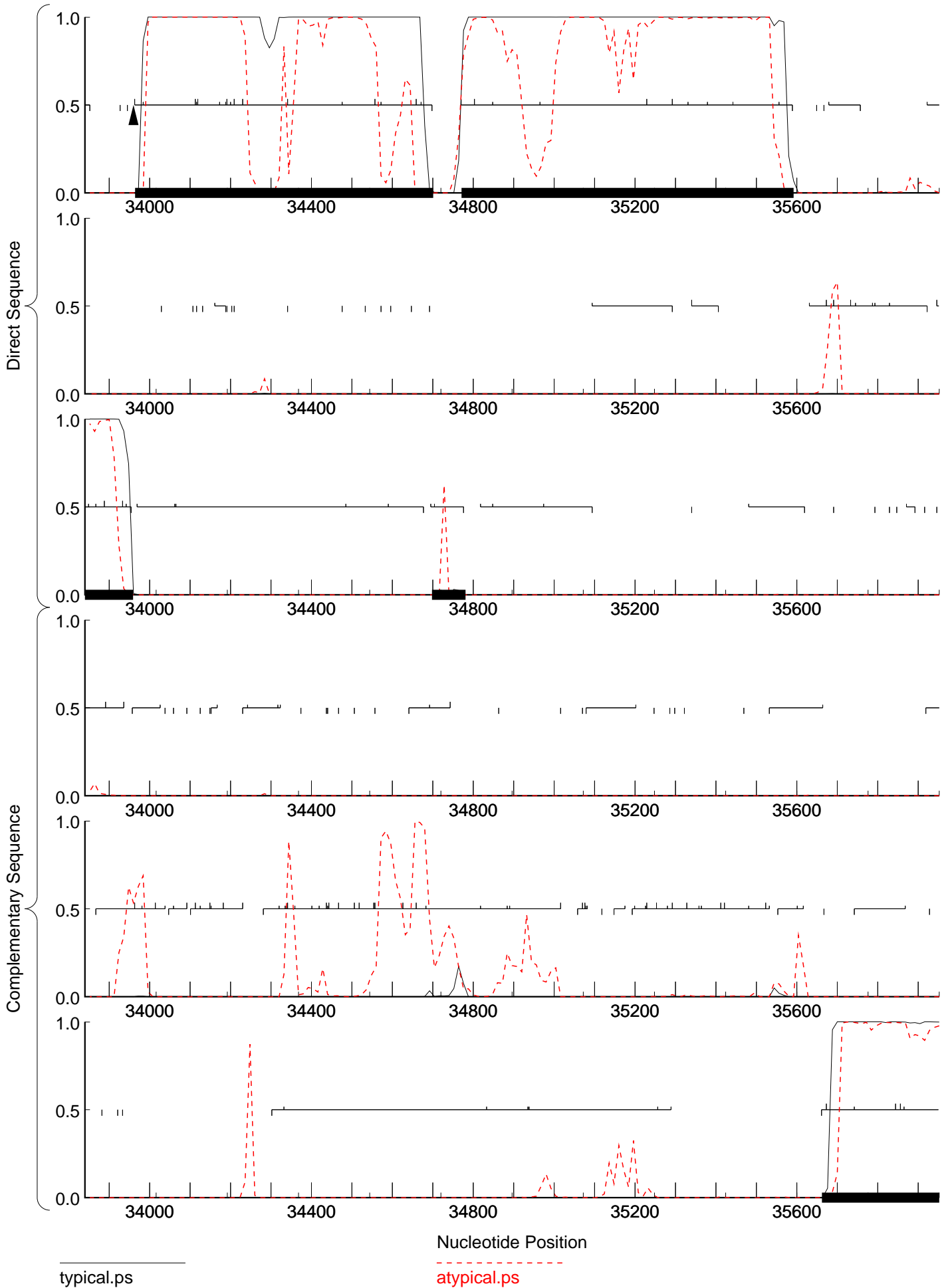


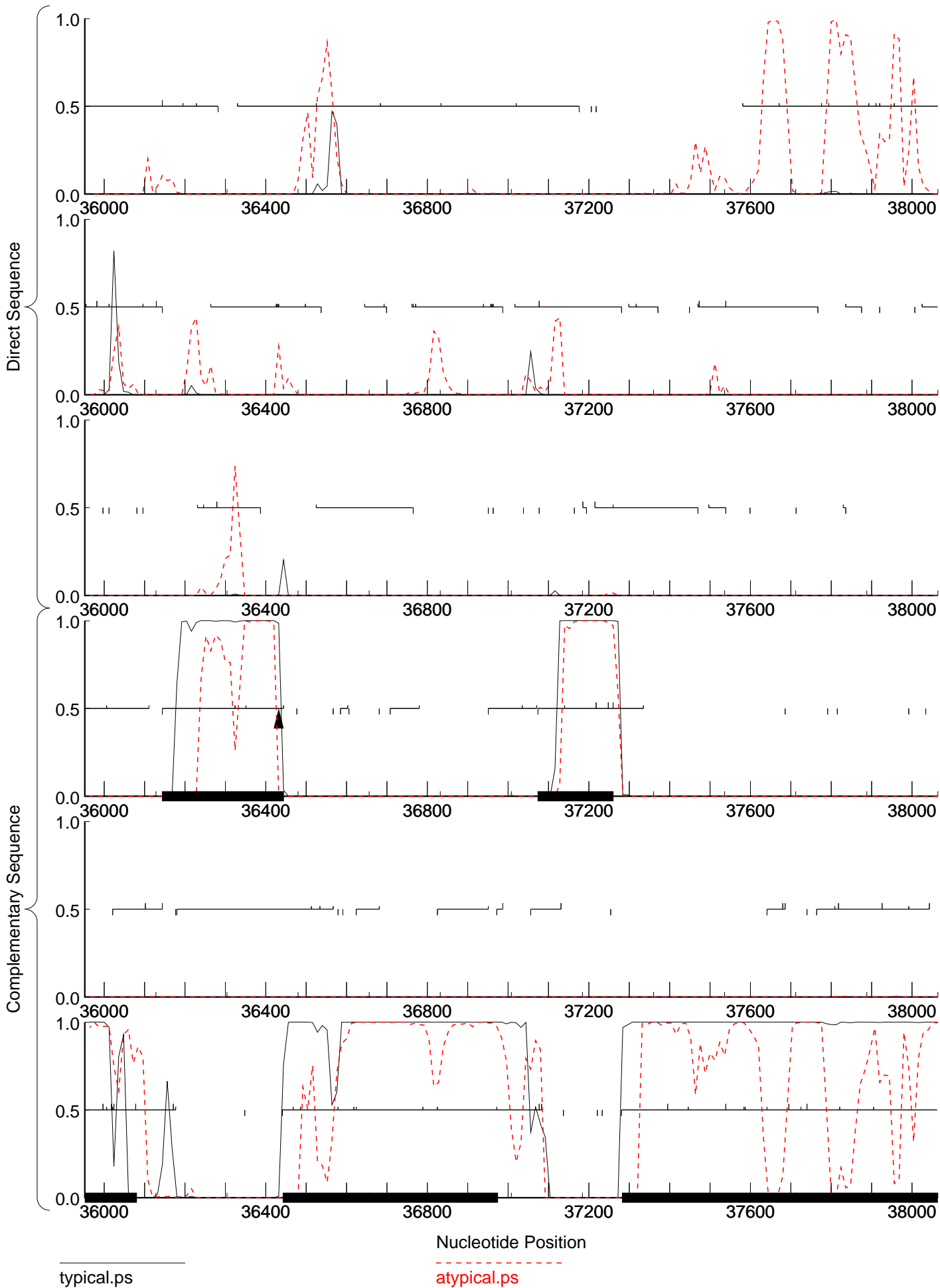


typical.ps

Nucleotide Position

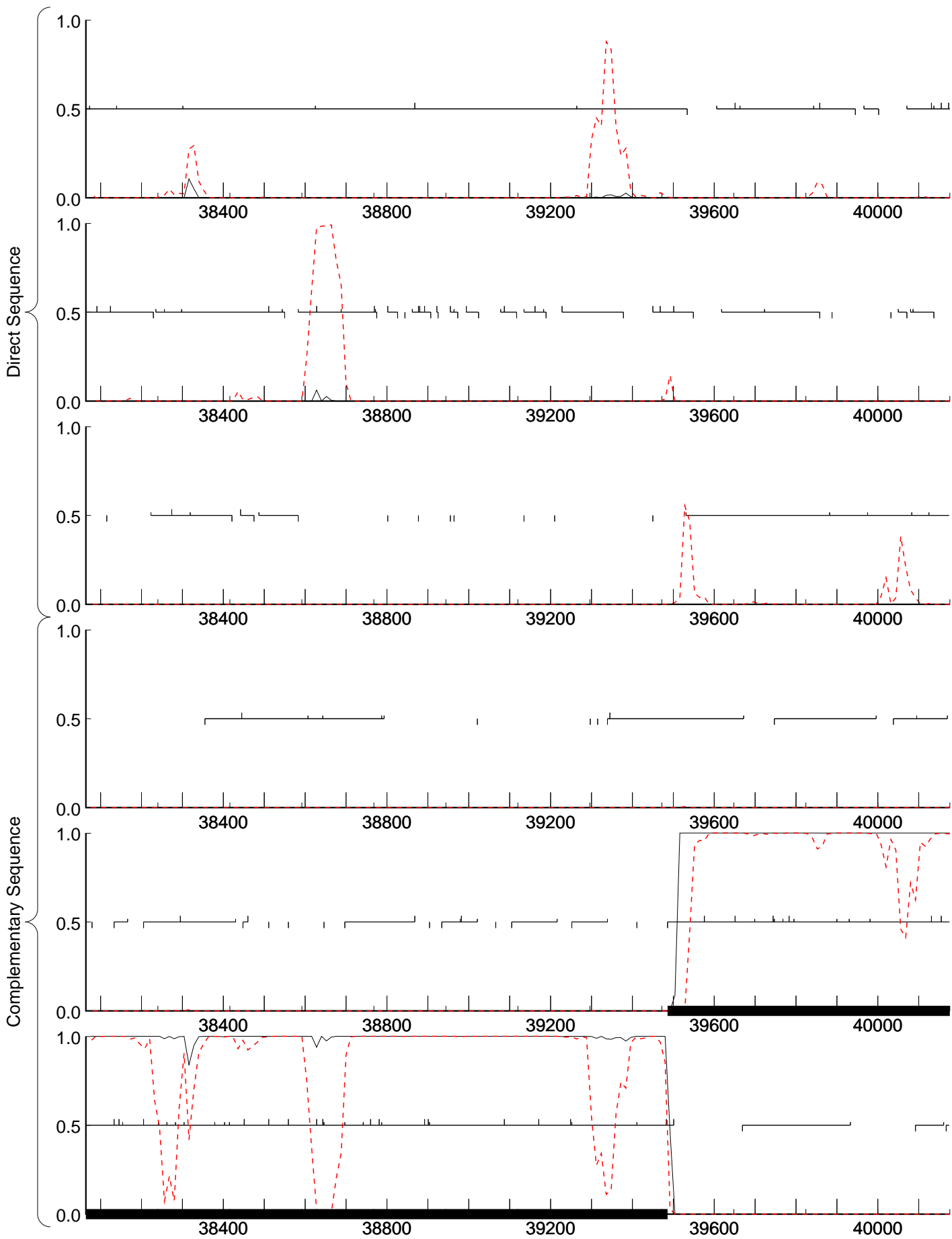
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typical.ps

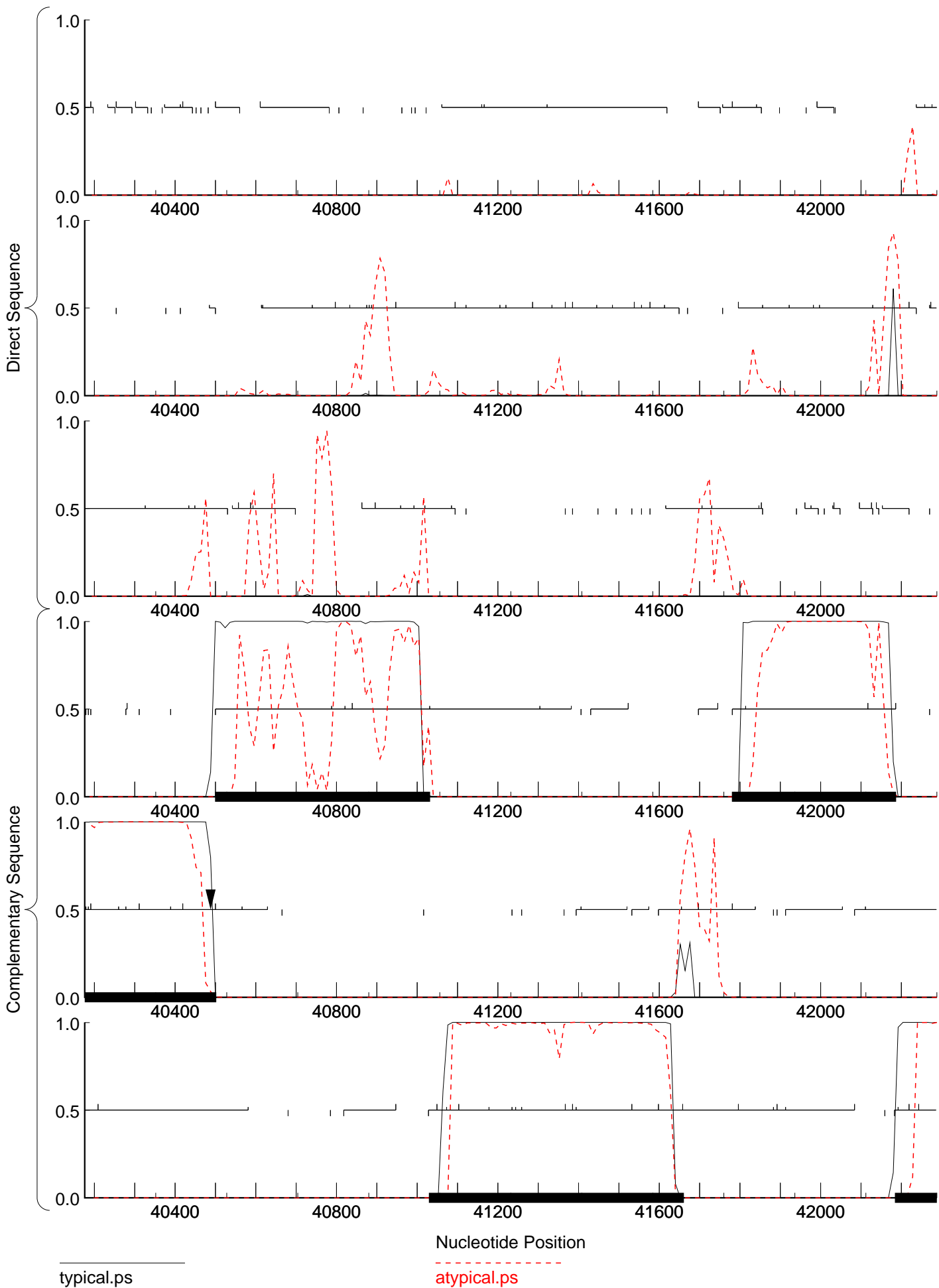
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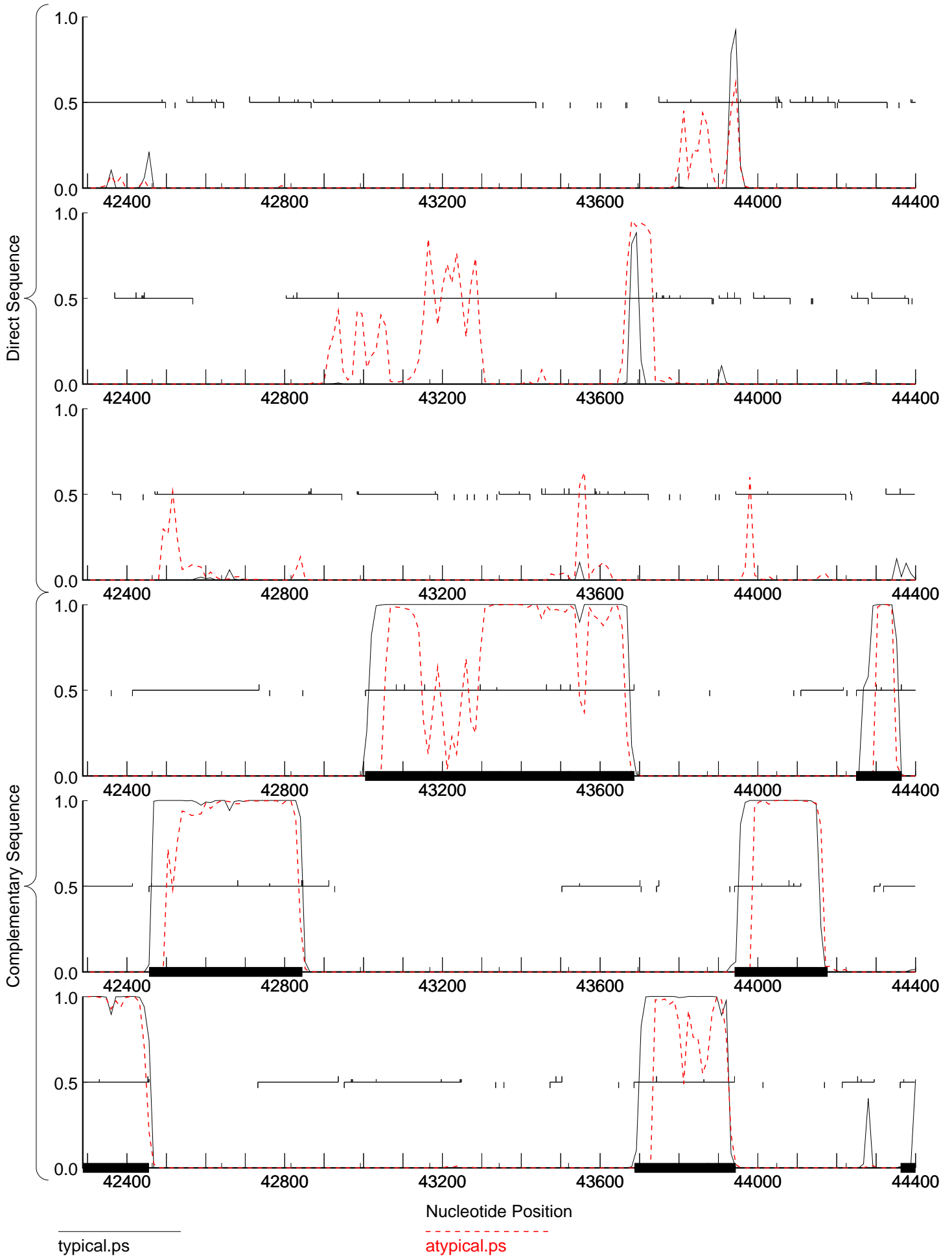


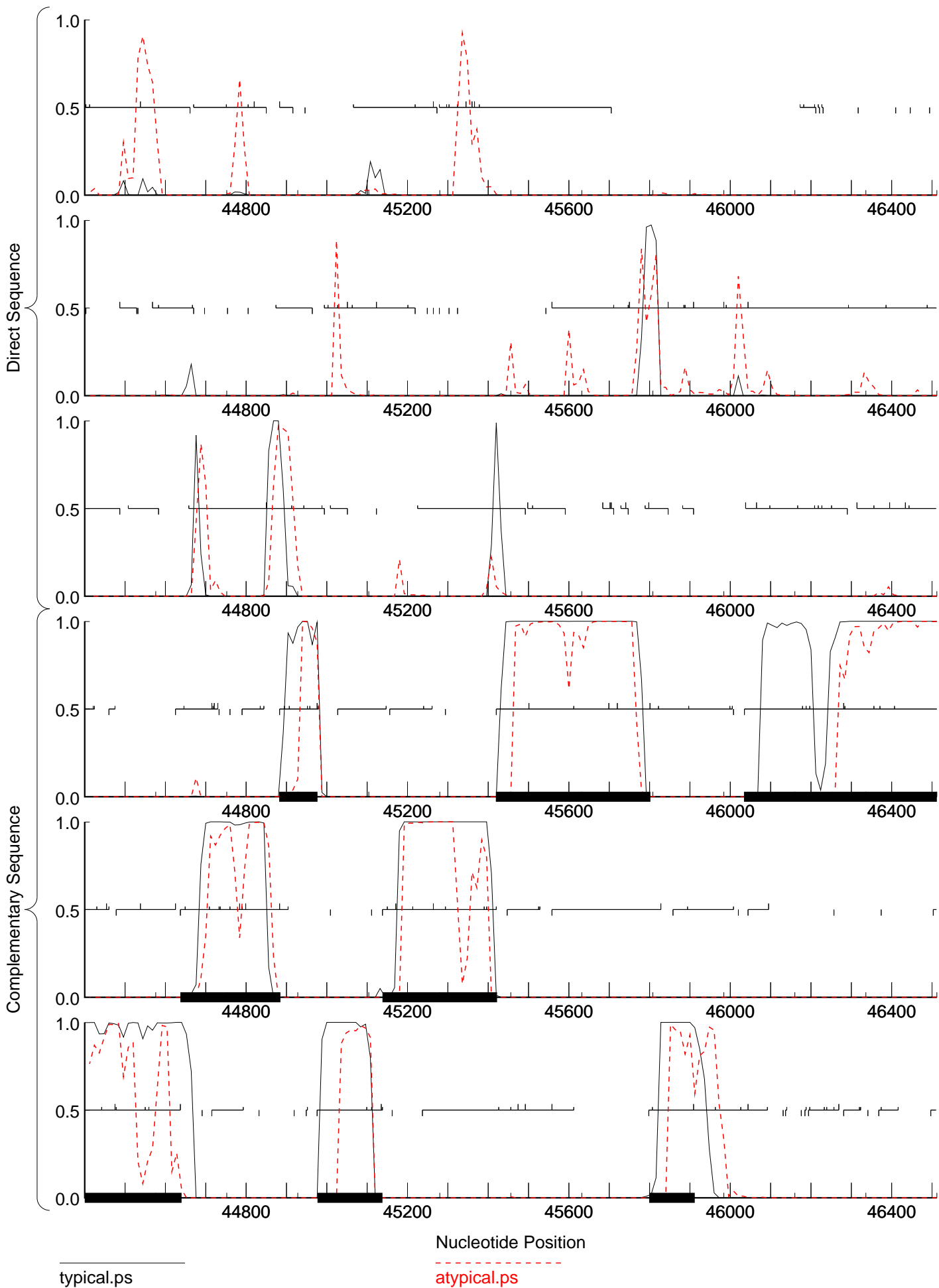
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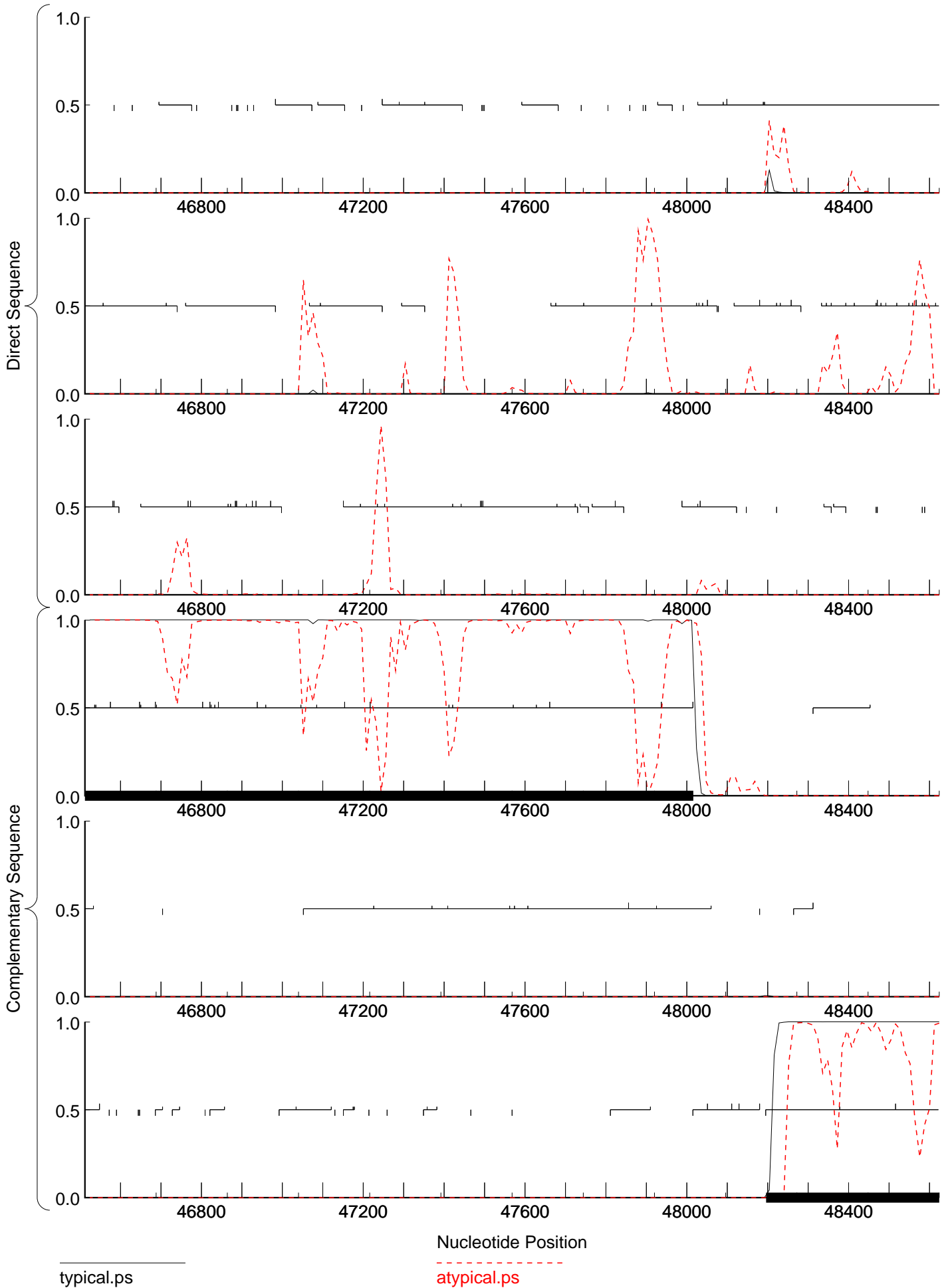
Nucleotide Position

atypical.ps



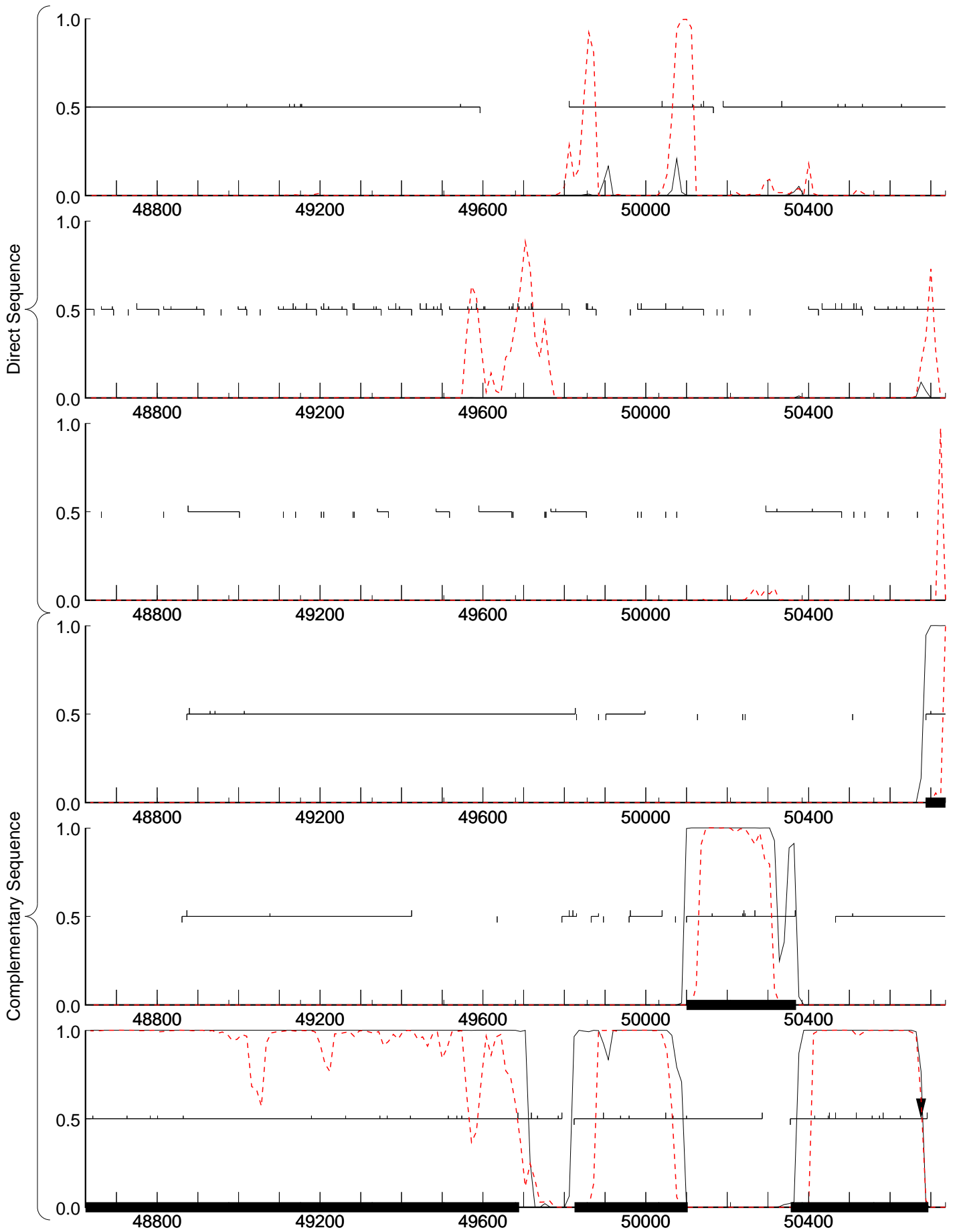






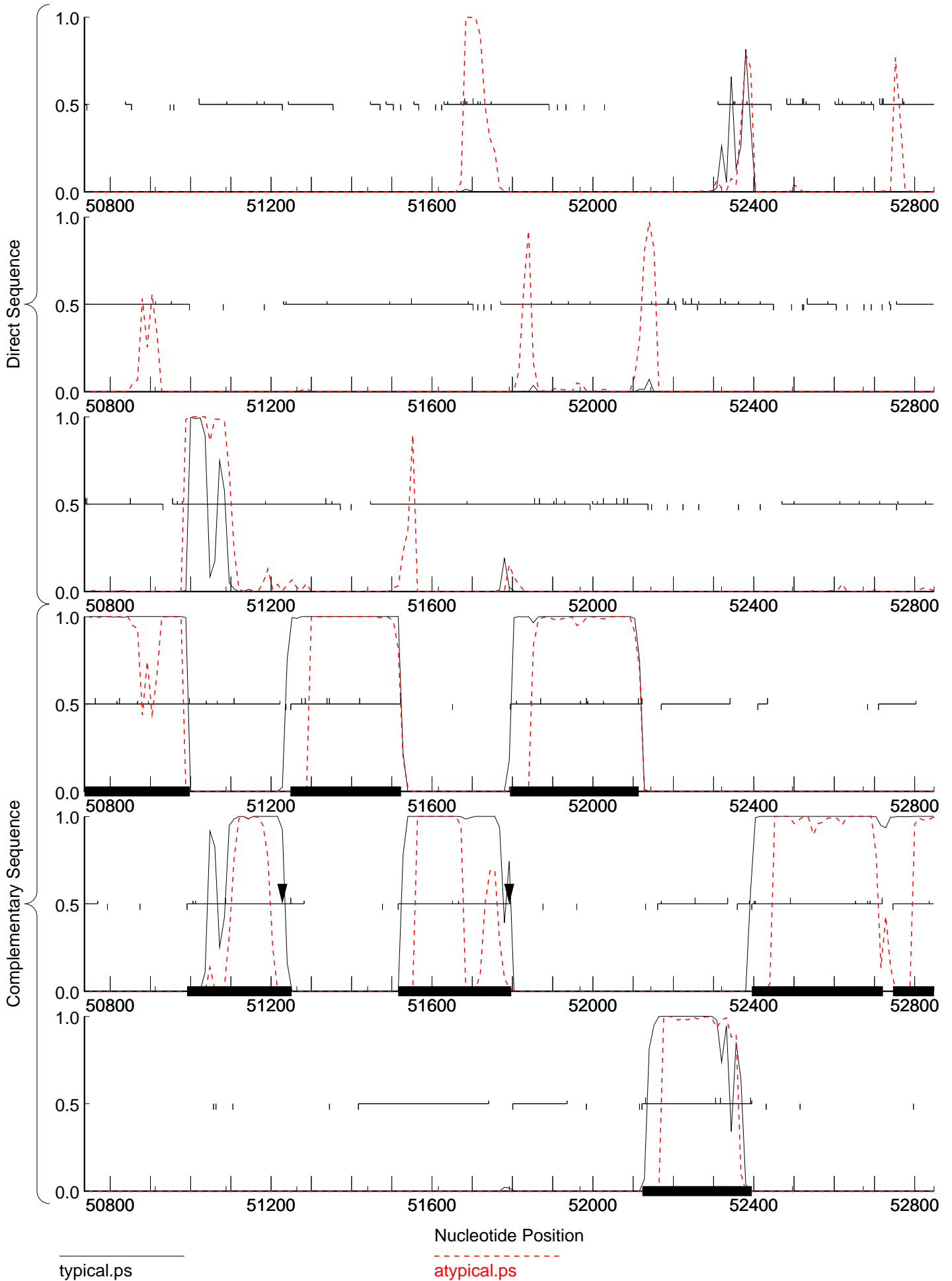
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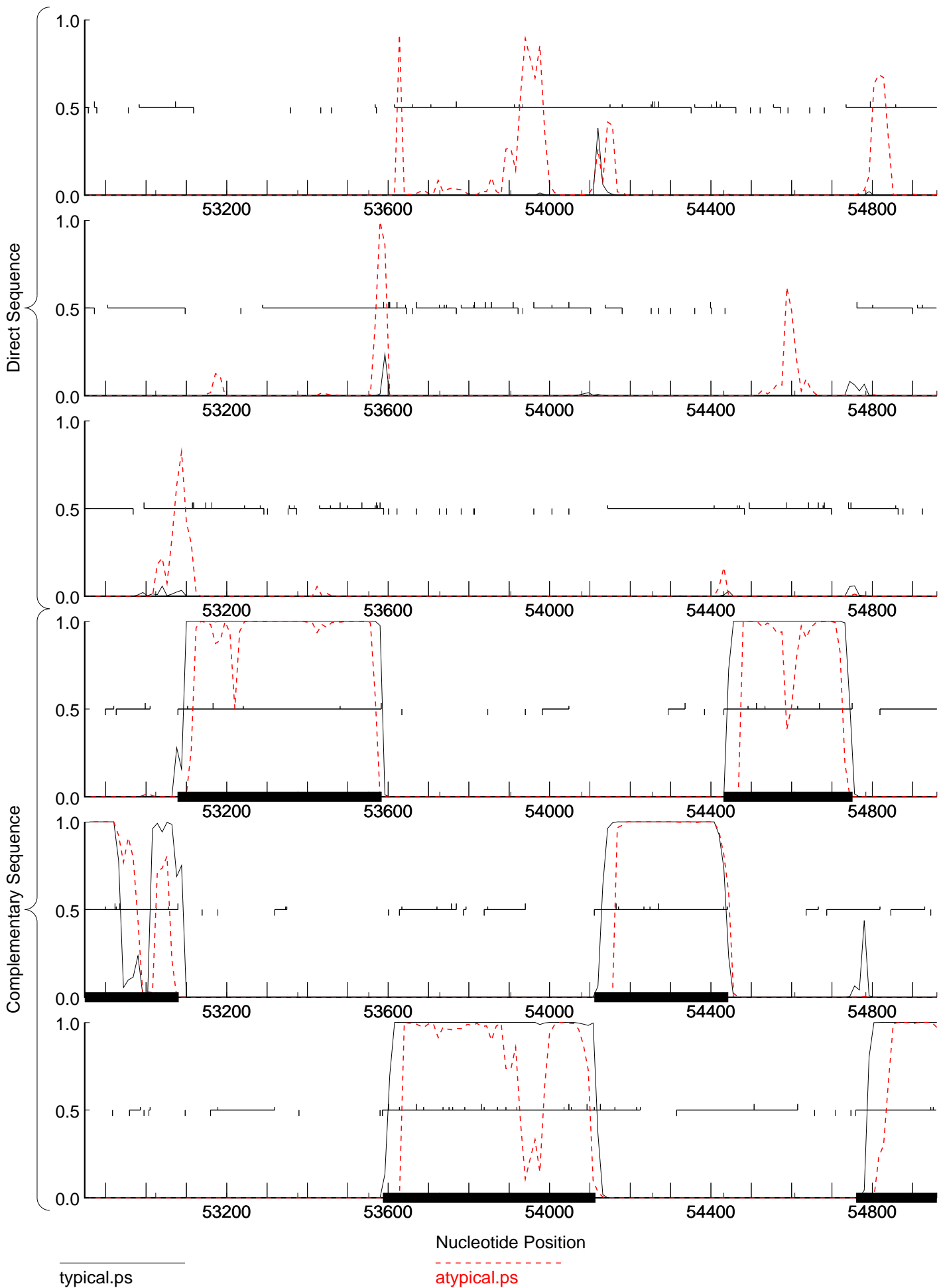
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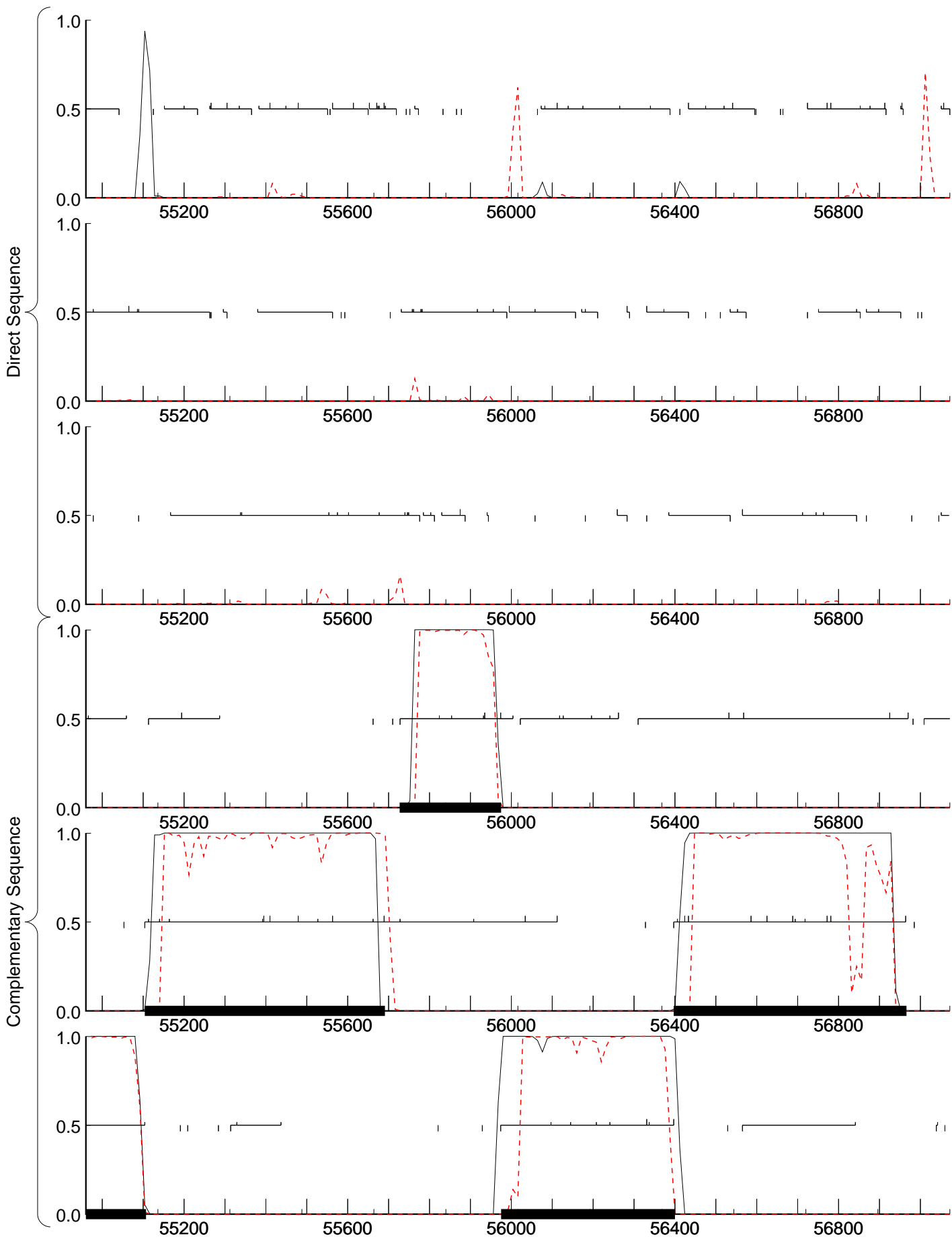


typical.ps

atypical.ps

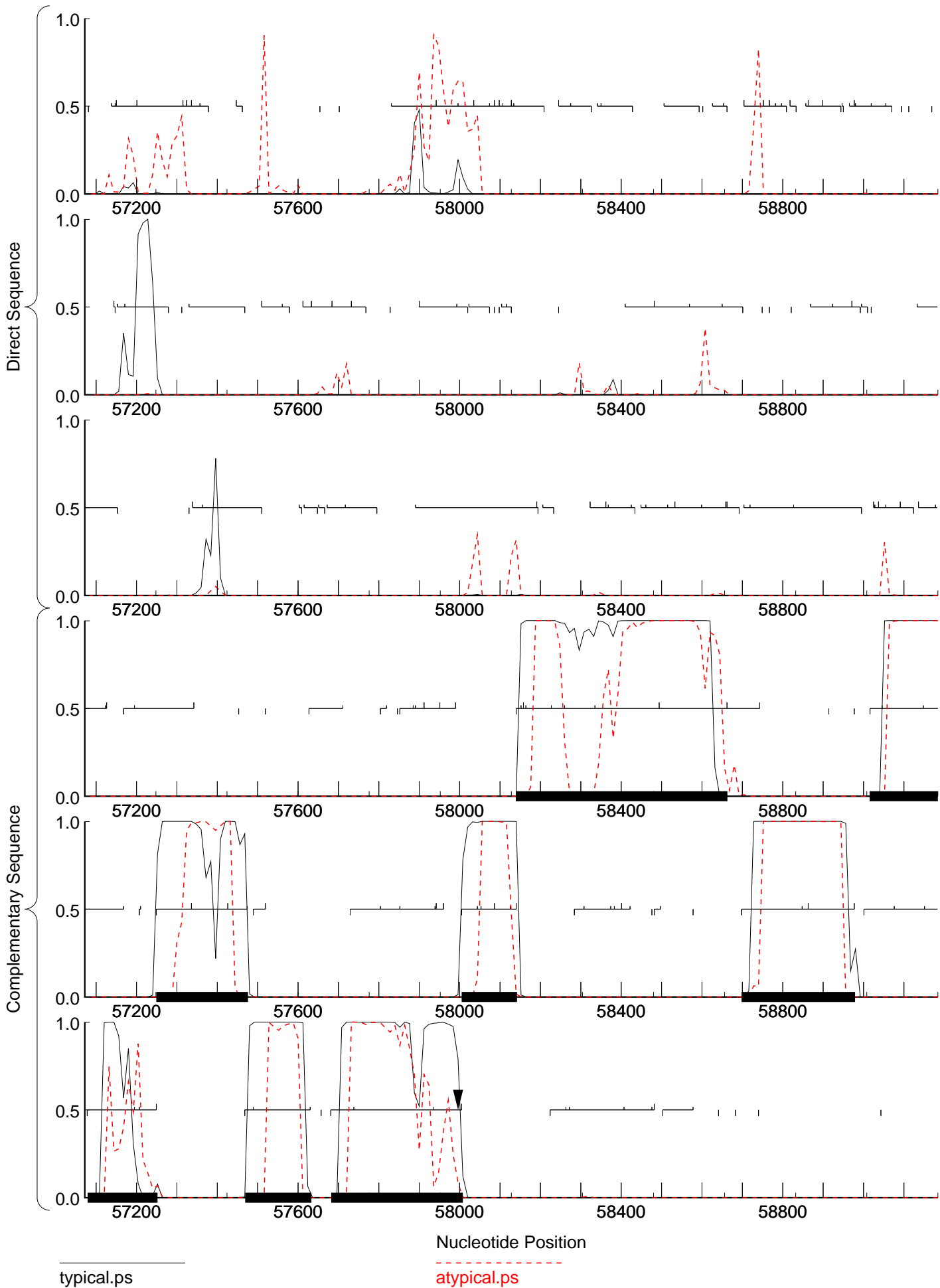


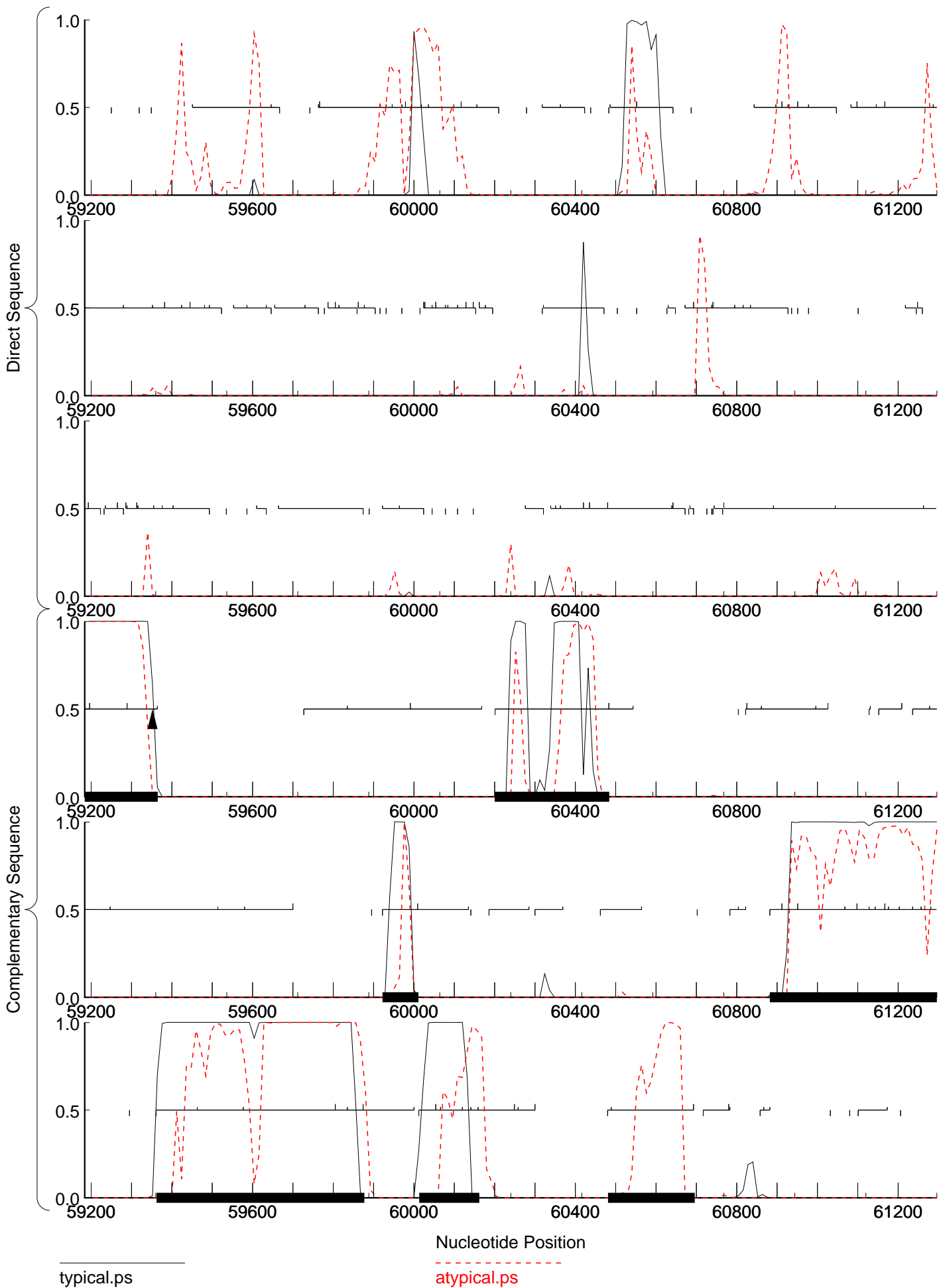




typical.ps

atypical.ps





typical.ps

atypical.ps

