

# GeneMark

Version 2.5p (09.08.06)

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## PROGRAM INFORMATION

Sequence : Streptomyces phage Yaboi complete sequence, 131250 bp including 12433 bp terminal repeat, C  
Analysis Date : 7/16/18 at 12:21:01  
Pages : 64  
Sequence Length : 131250 bp  
GC Content : 49.30%

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,

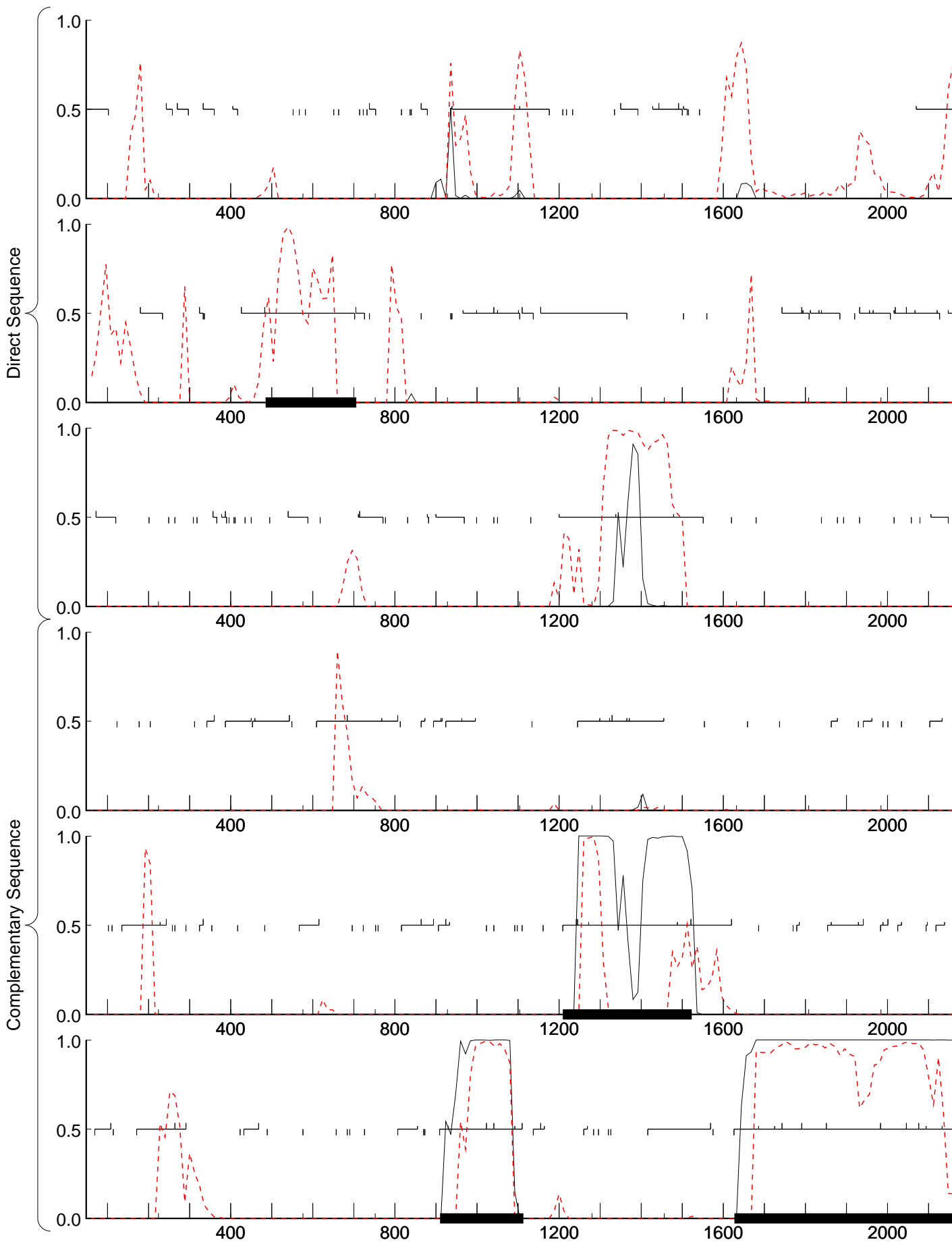
## MATRIX INFORMATION

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

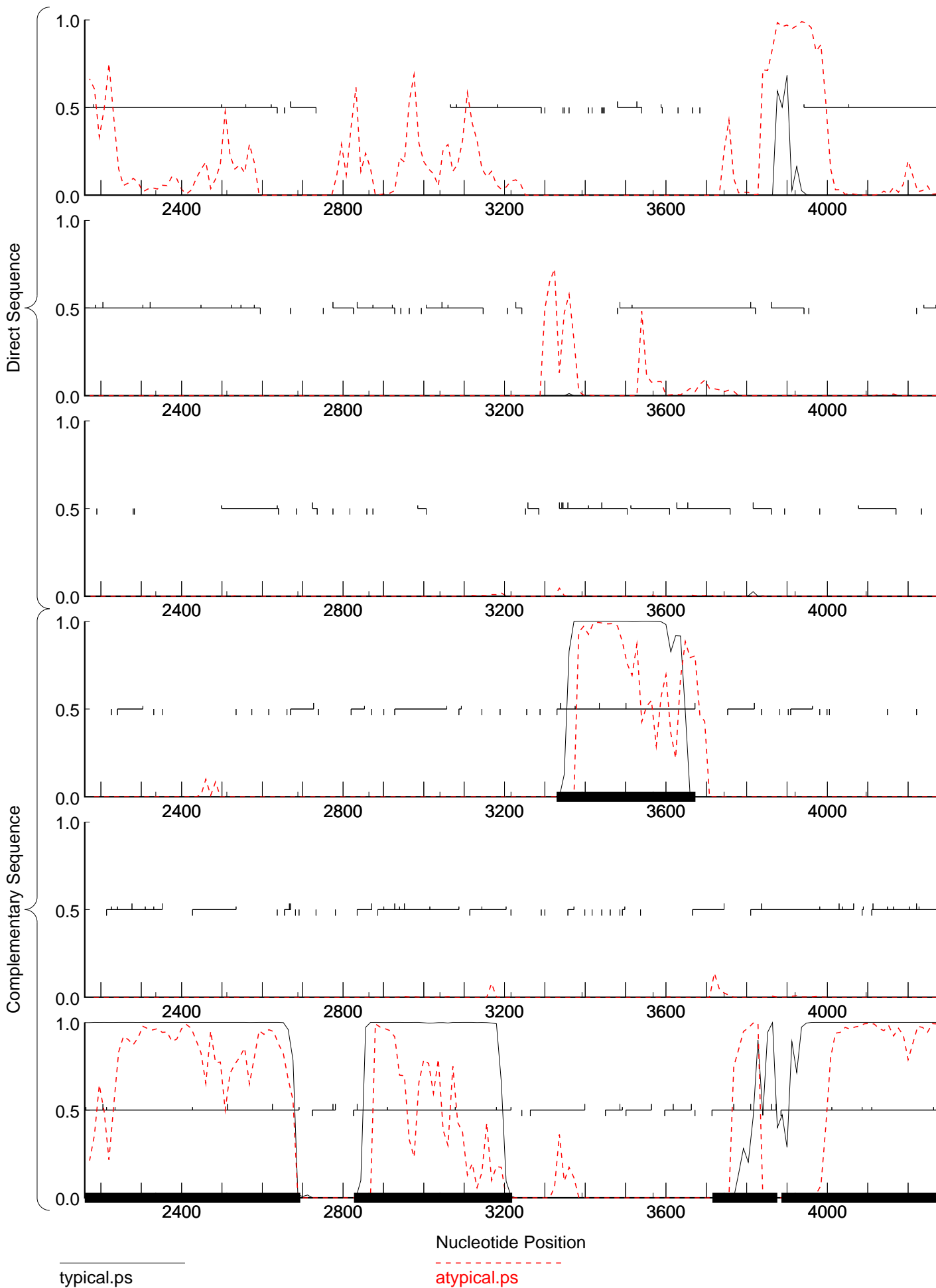
Training set derived by GeneMarkS, 4.30 October 2014  
Mon Jul 16 12:21:01 2018

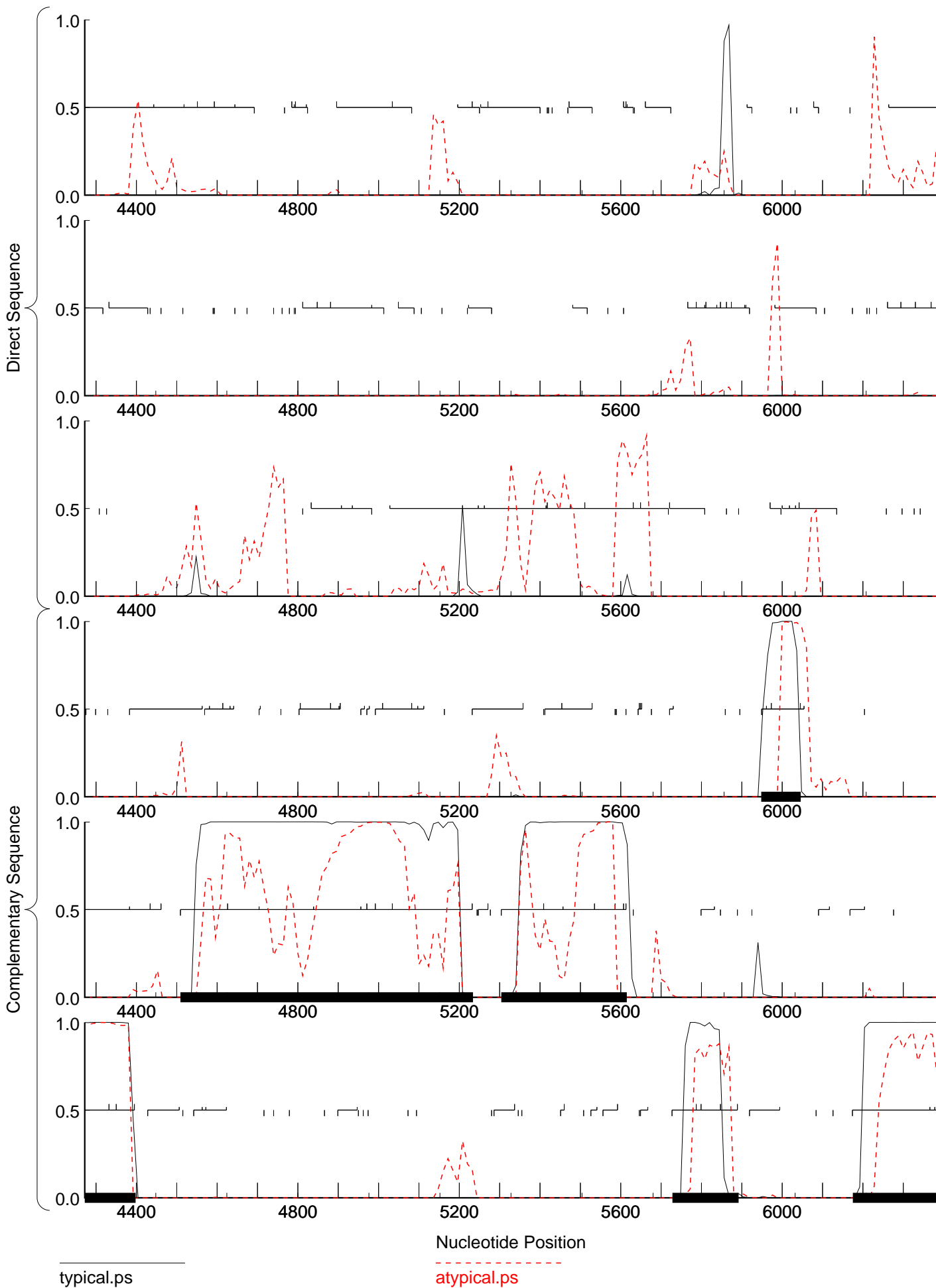


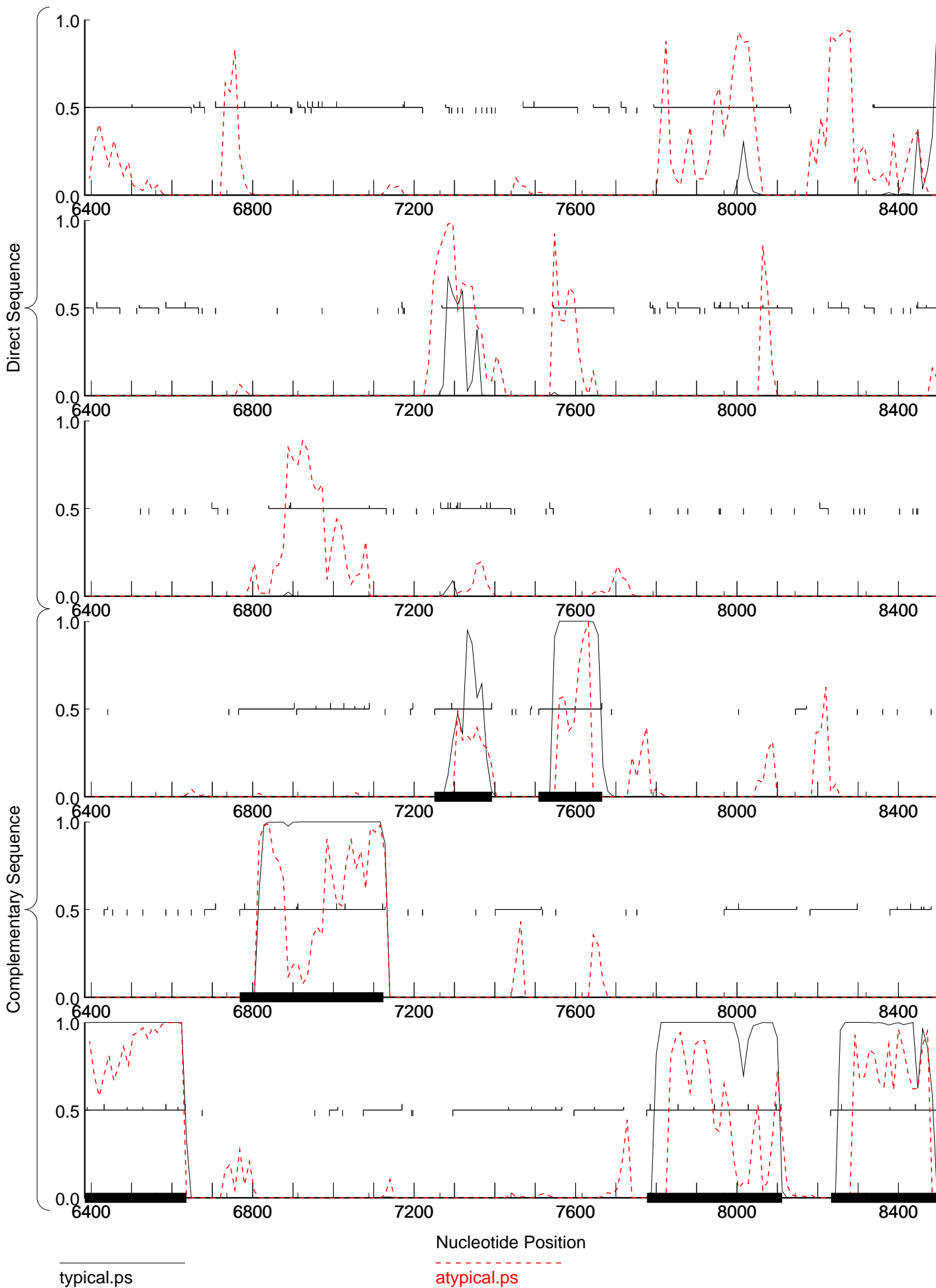
typical.ps

atypical.ps

phage Yaboi complete sequence, 131250 bp including 12433 bp terminal repeat, Cluster BE2, Order 4, Window 96, Step 12, 3/64  
phage Yaboi complete sequence, 131250 bp including 12433 bp terminal repeat, Cluster BE2, Order 2, Window 96, Step 12, 3/64

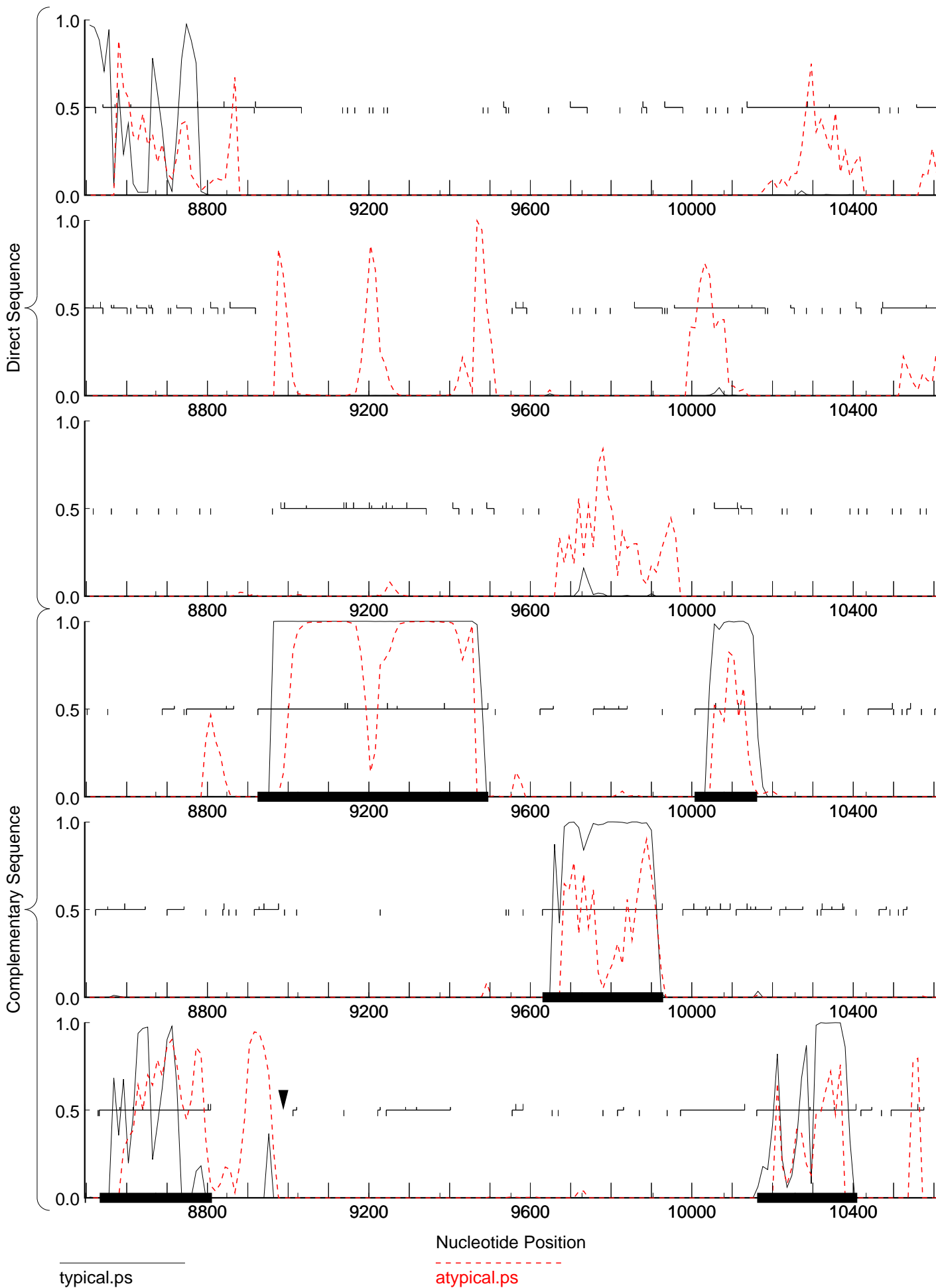






typical.ps

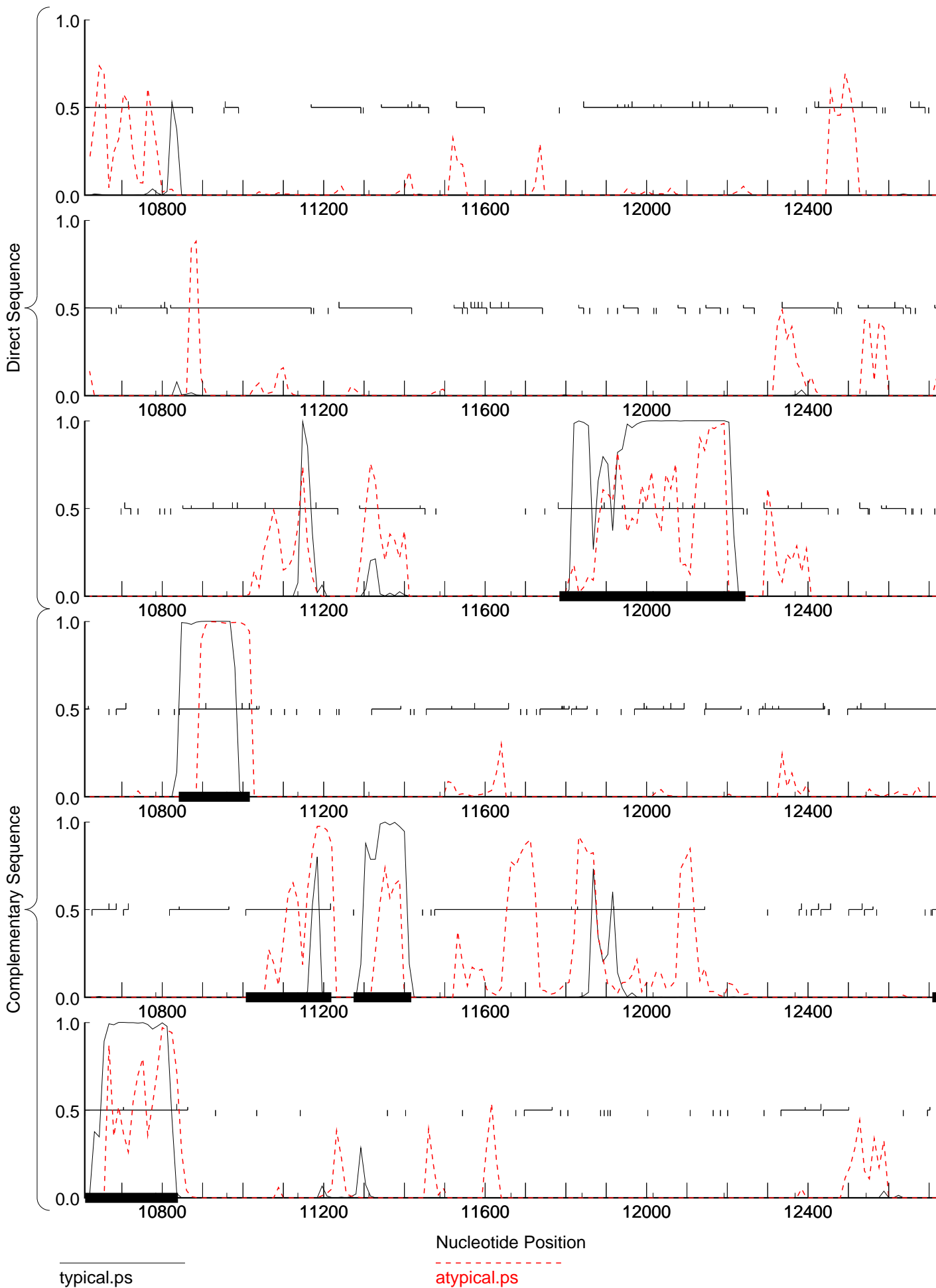
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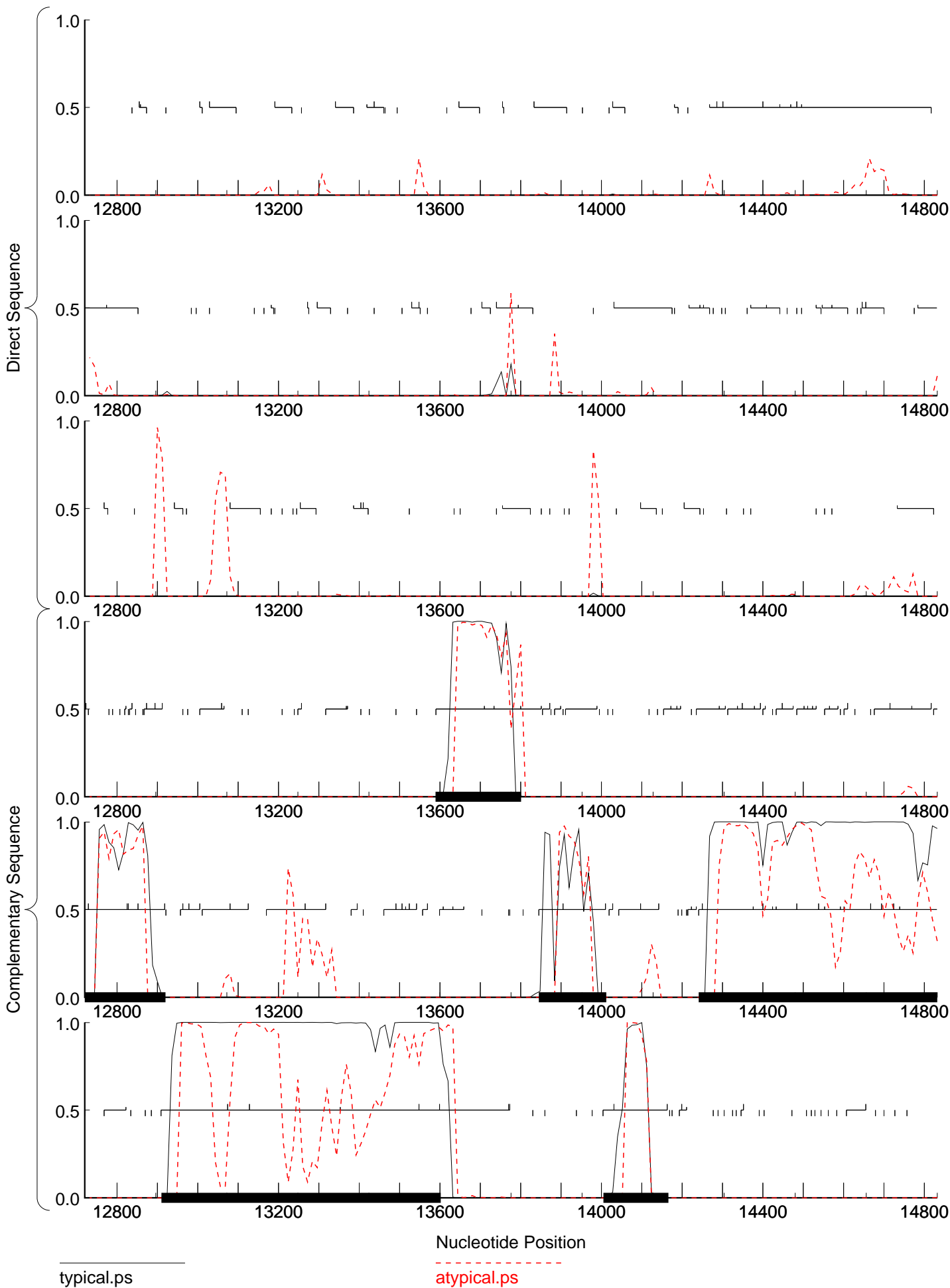


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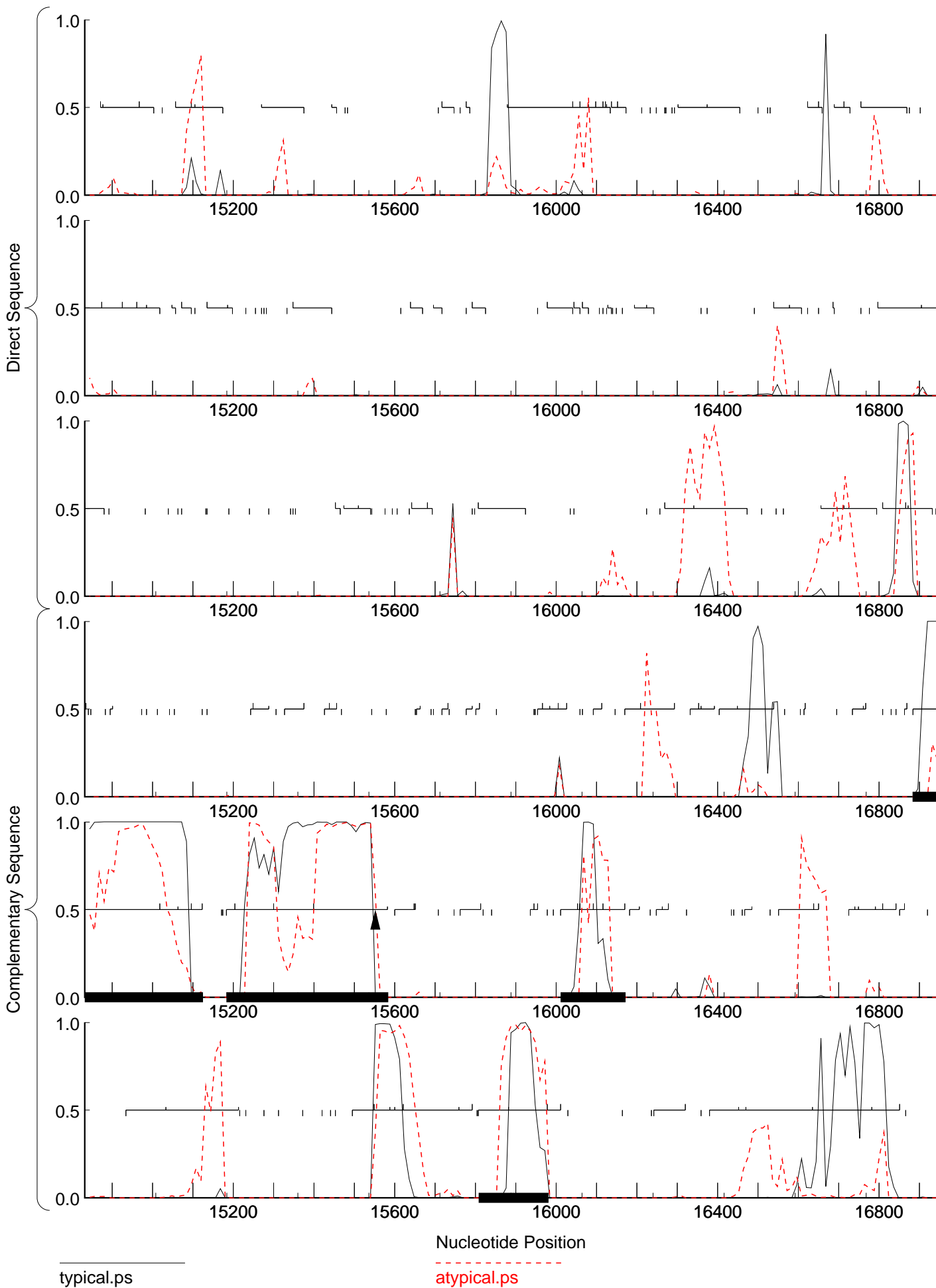
Nucleotide Position

atypical.ps



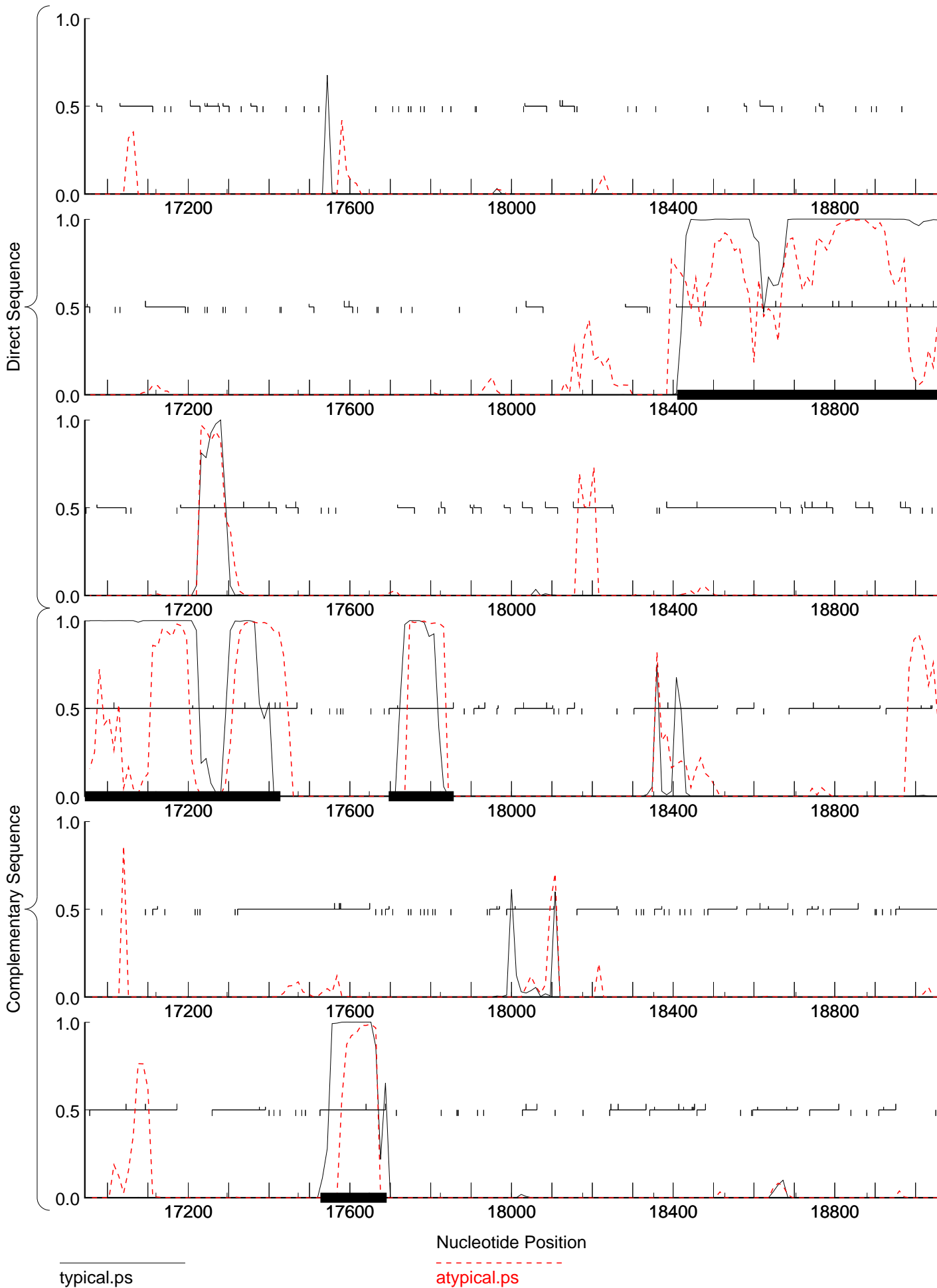






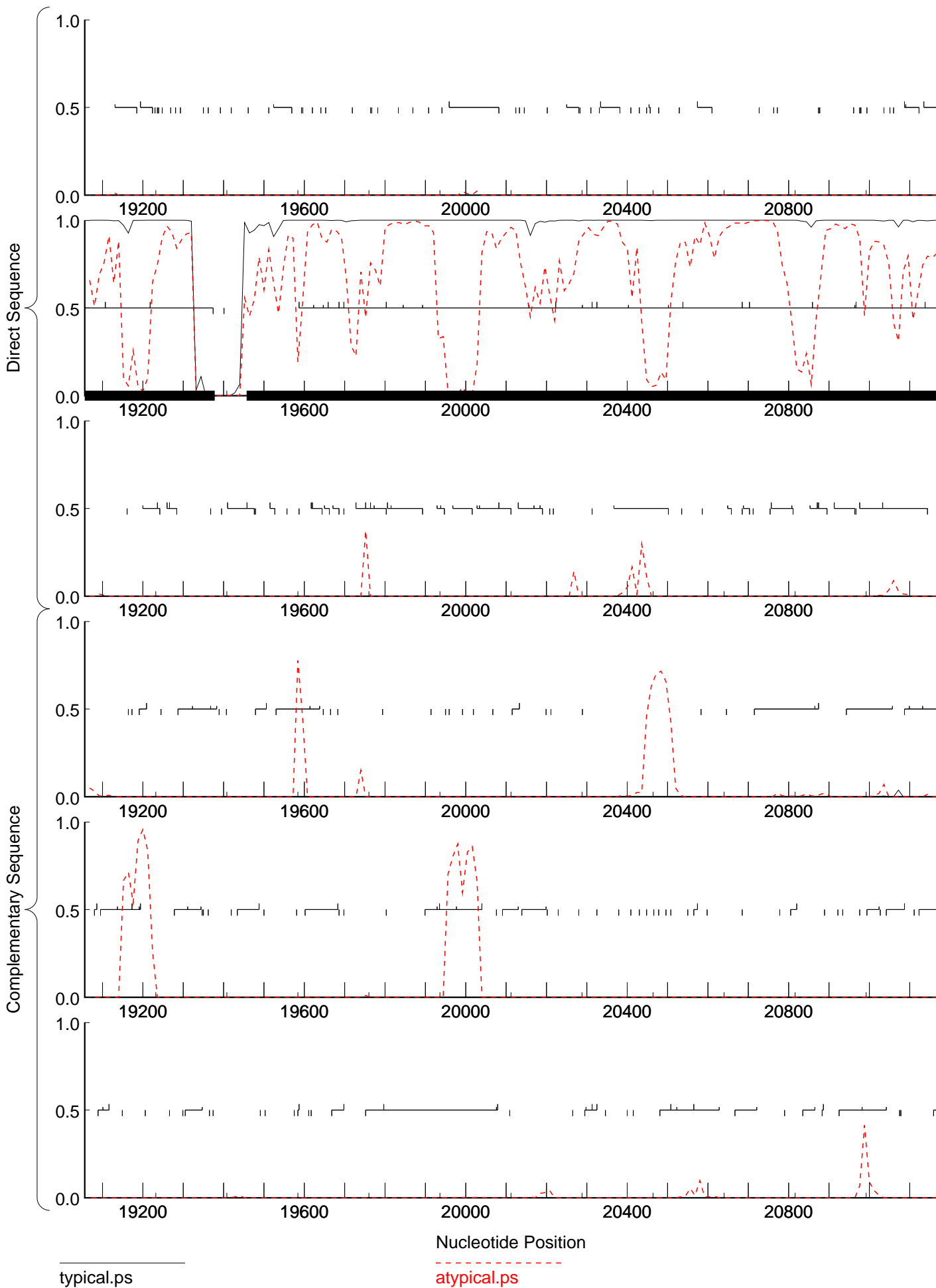
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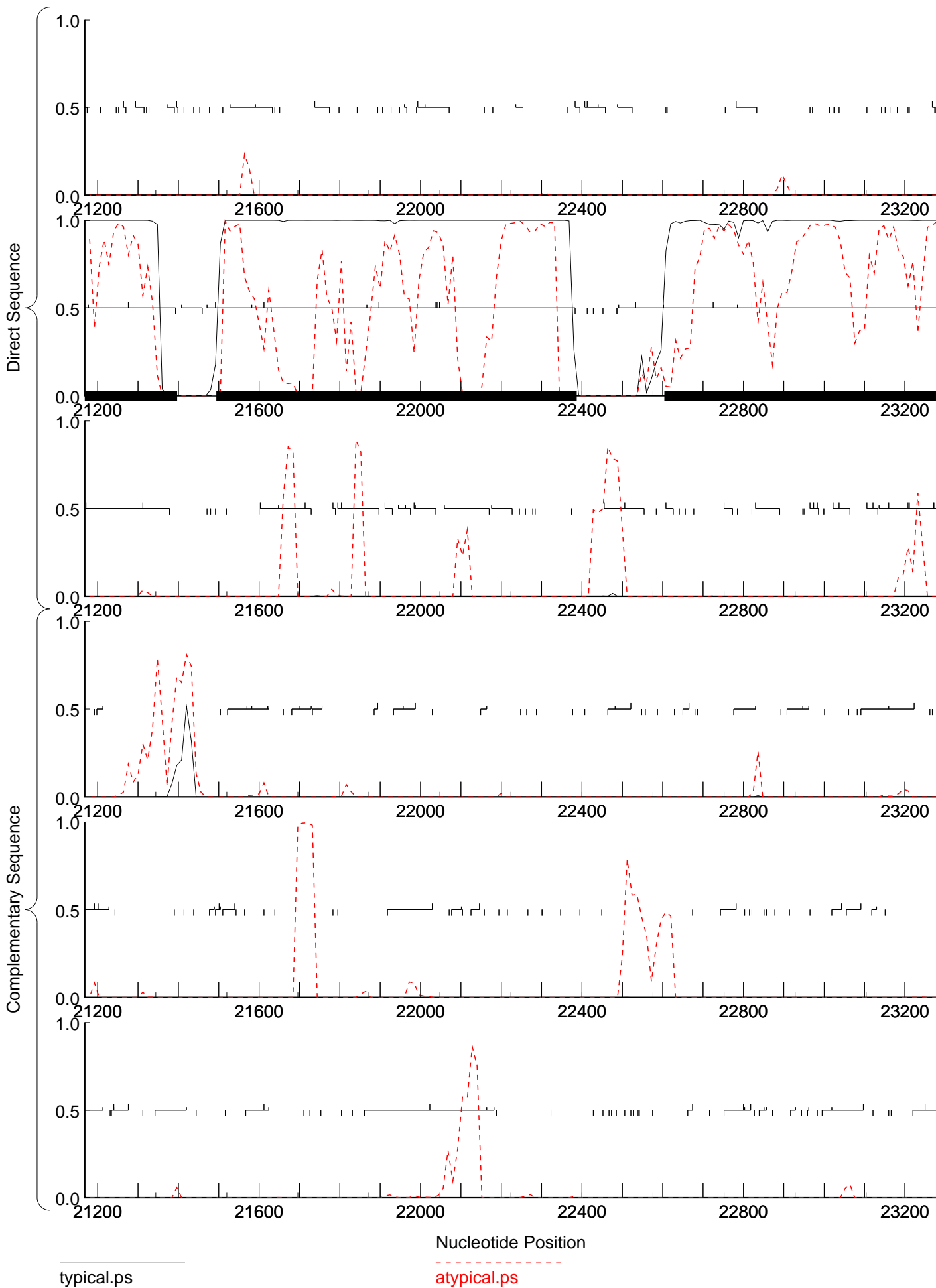
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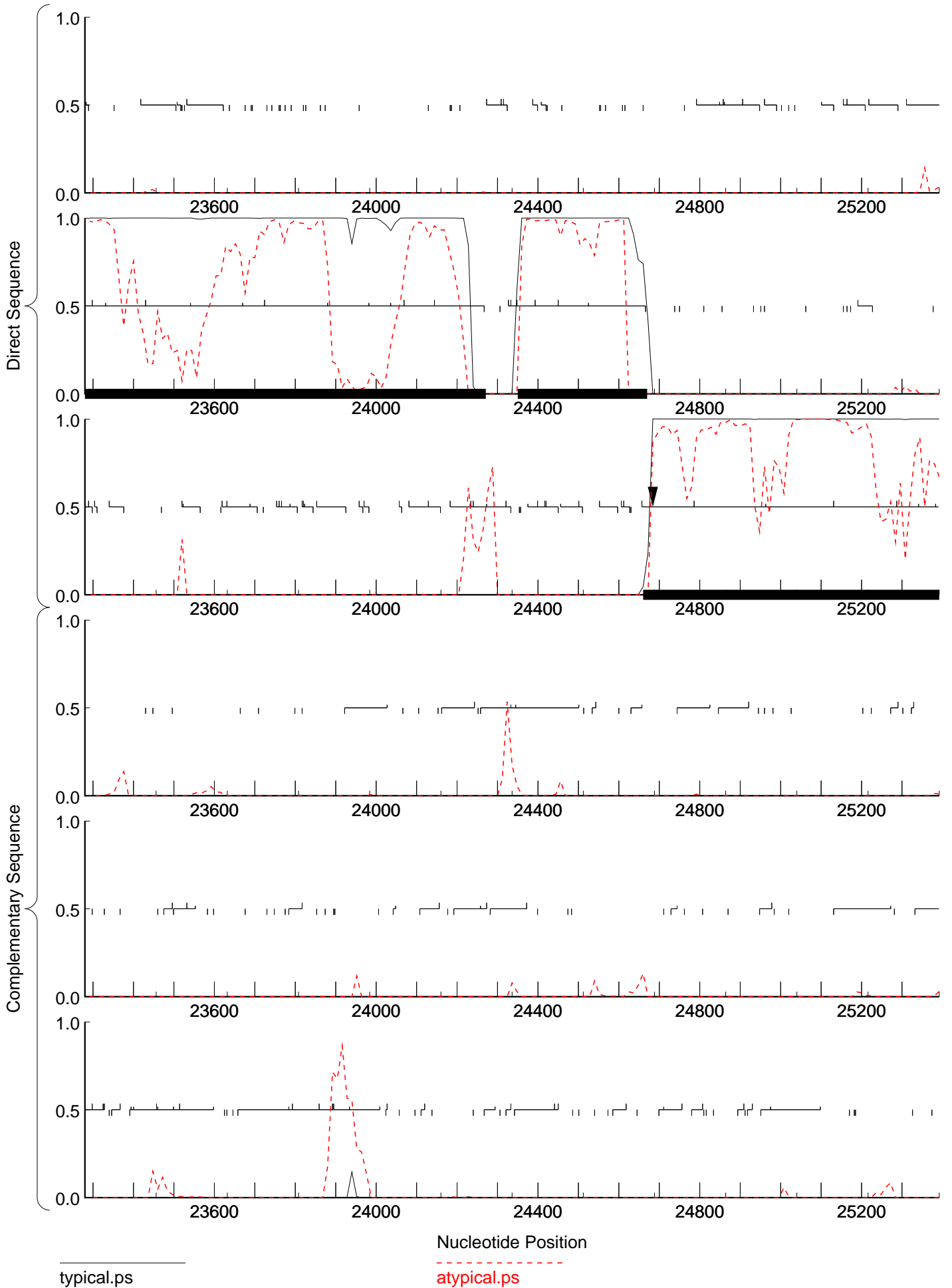


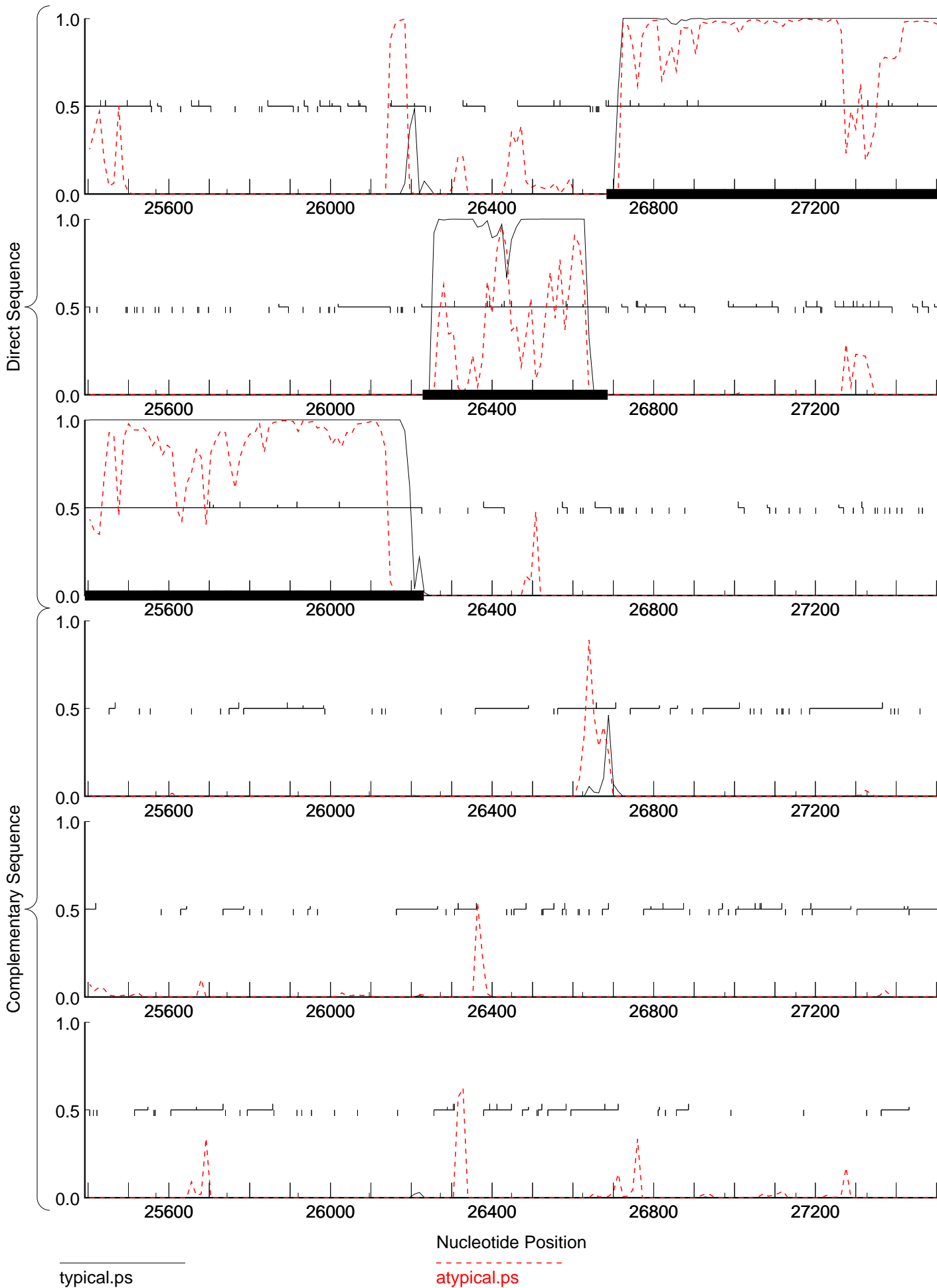
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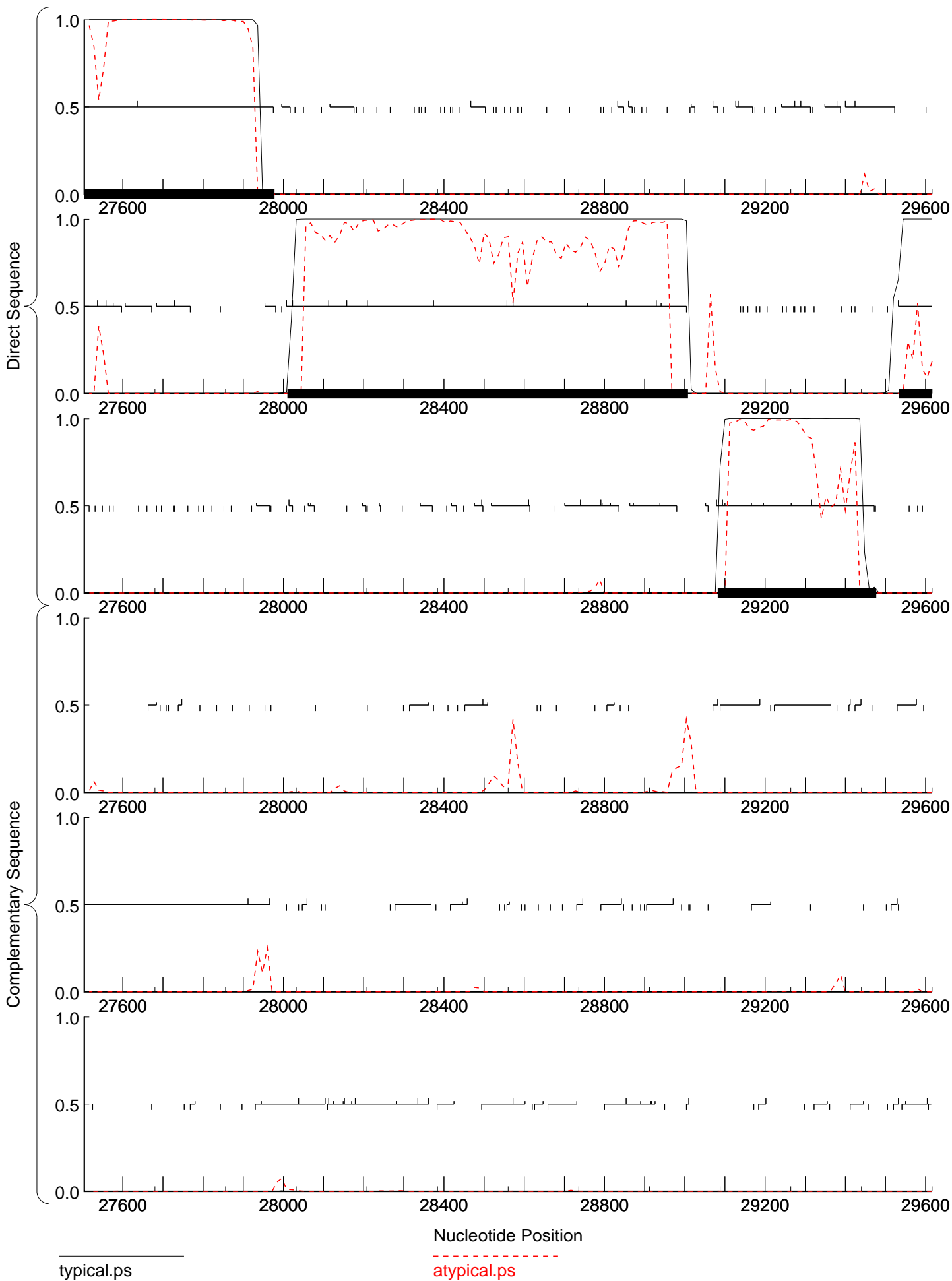
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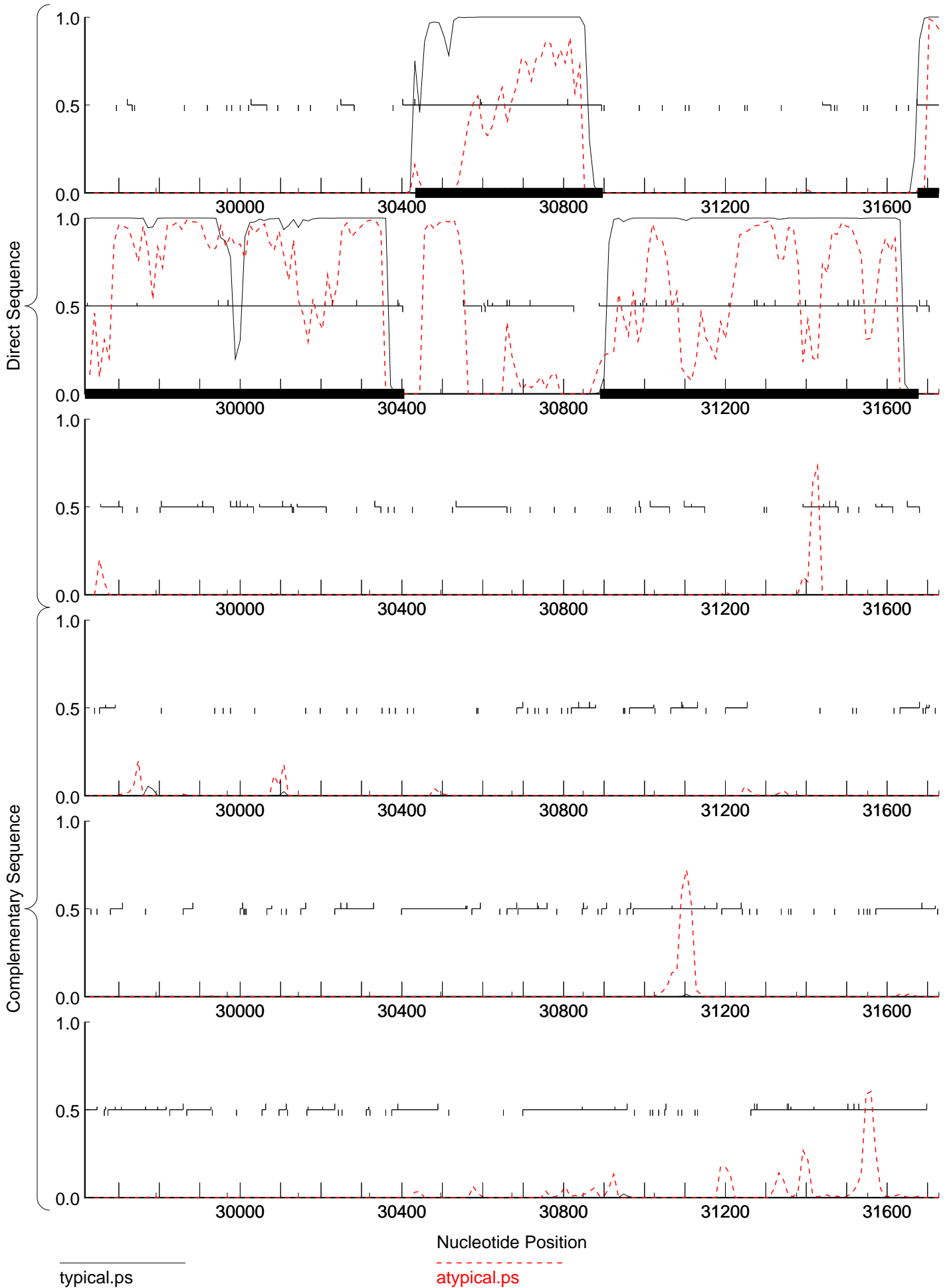




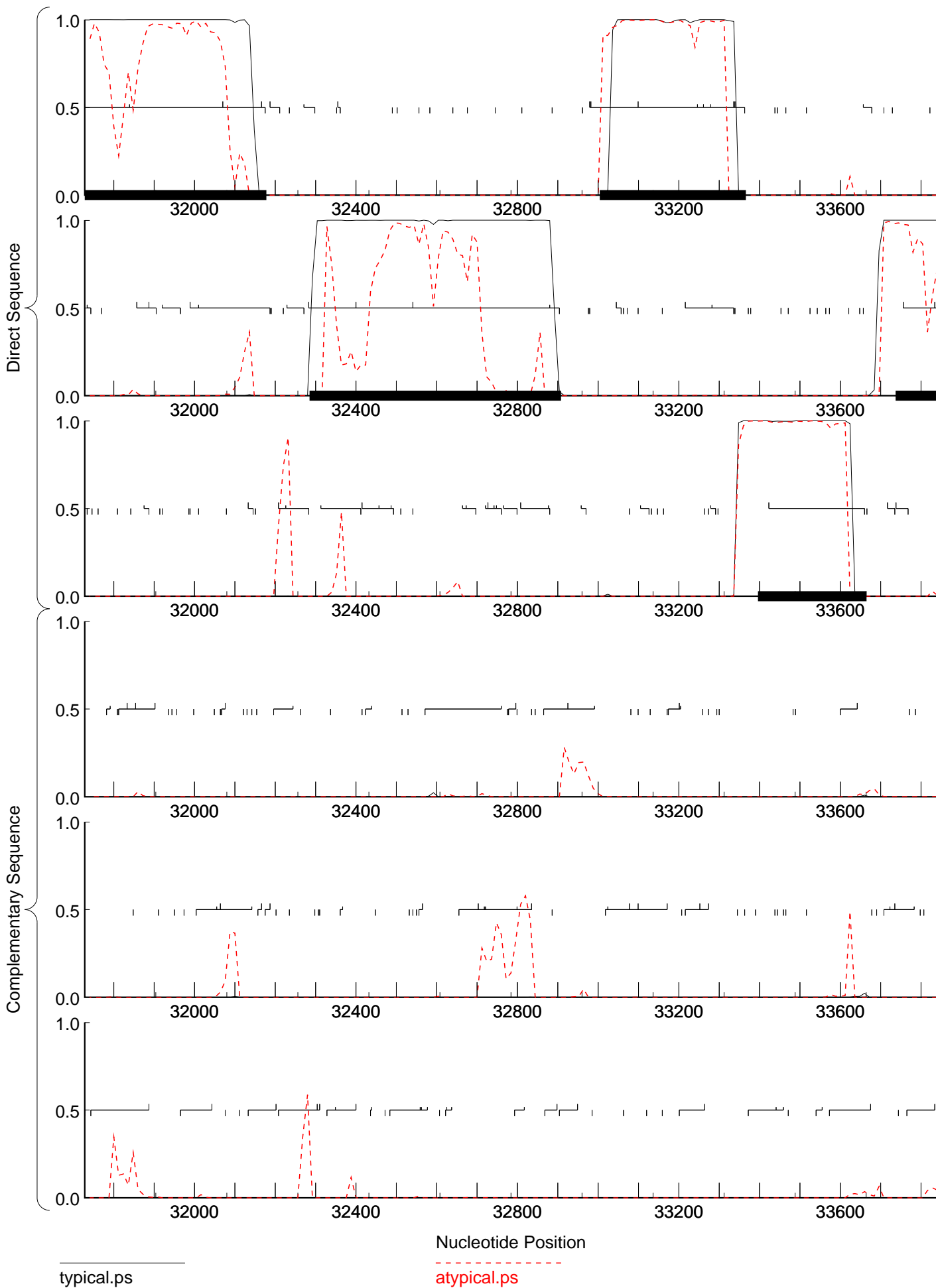


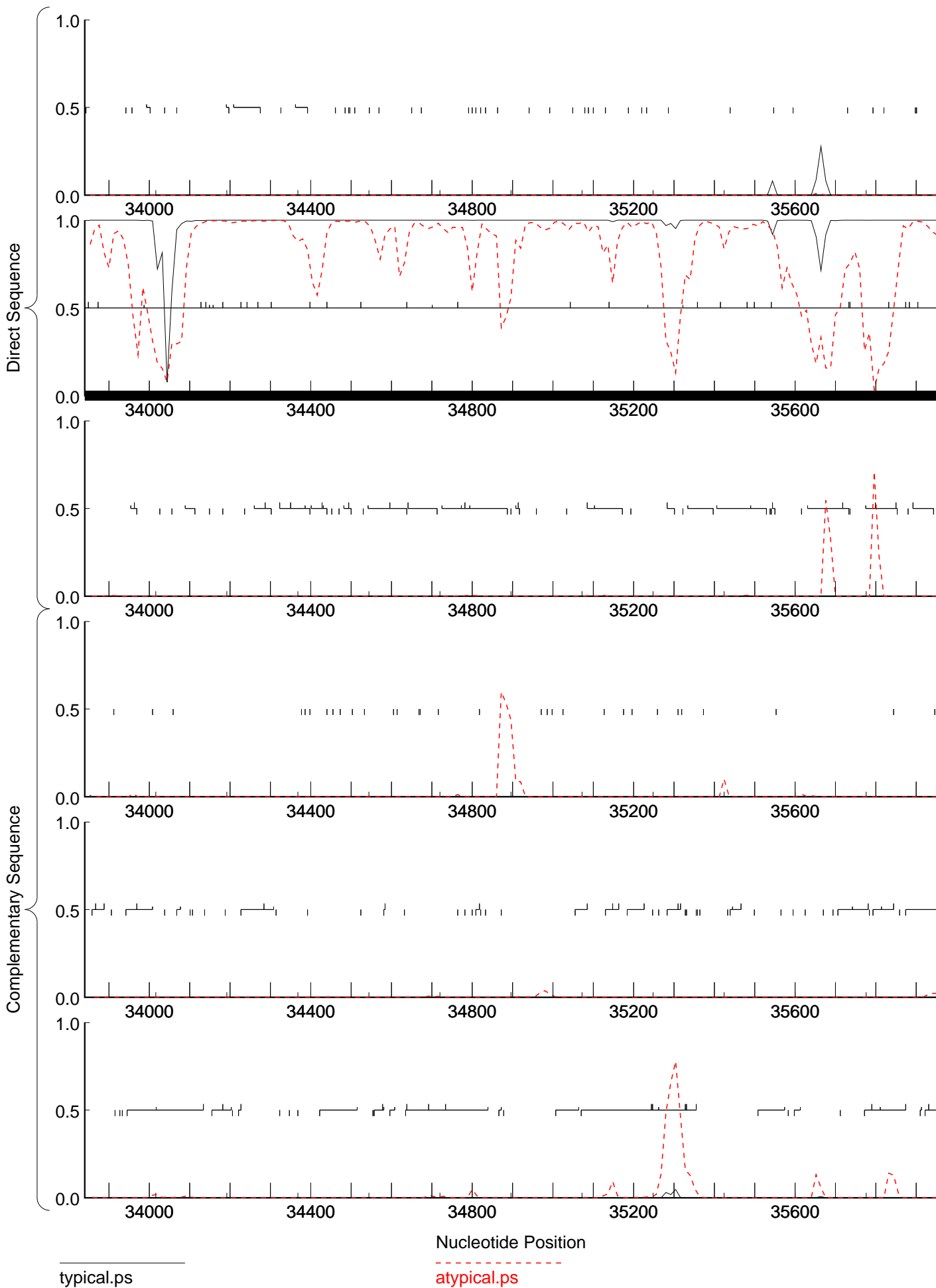
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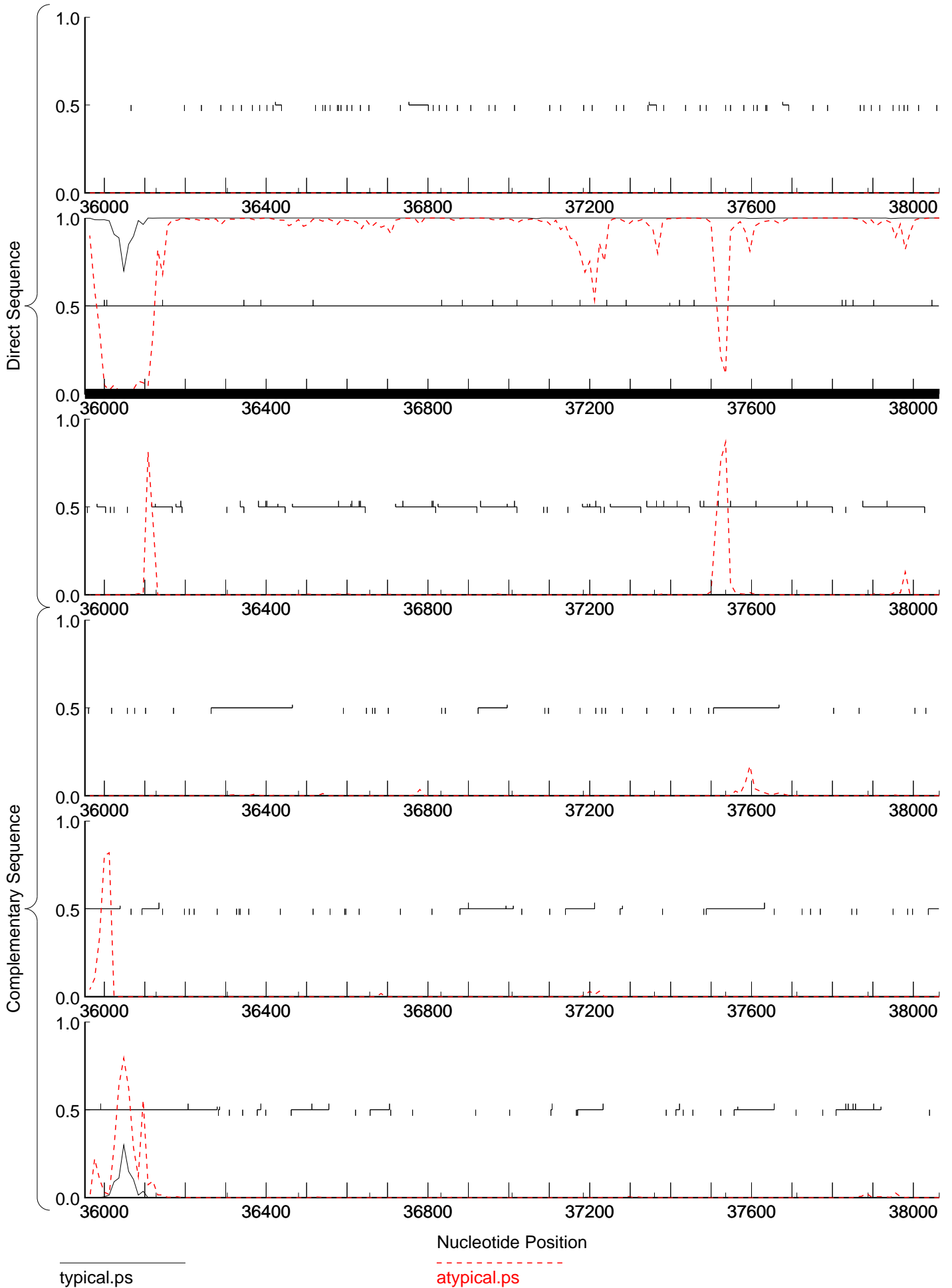
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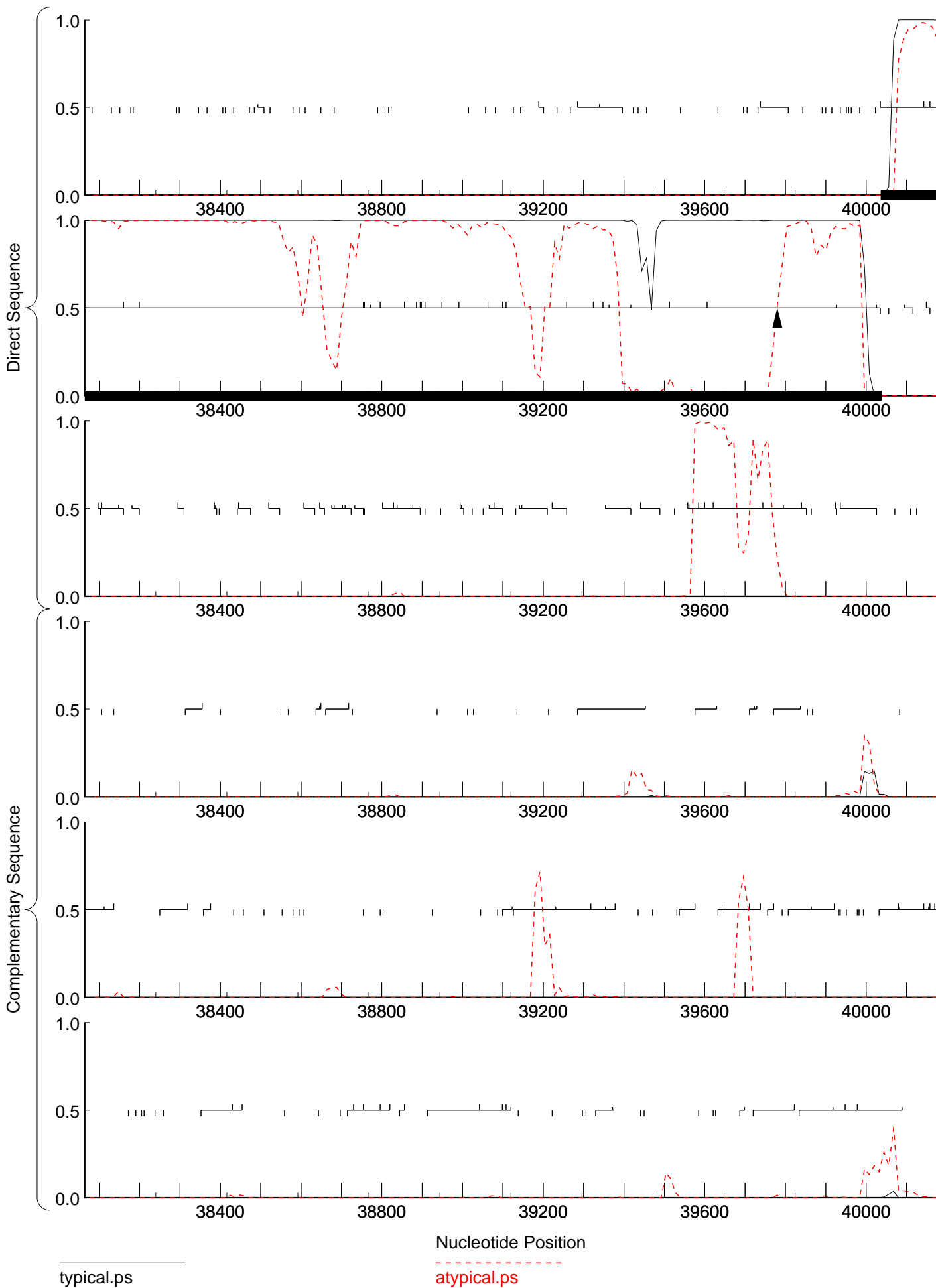


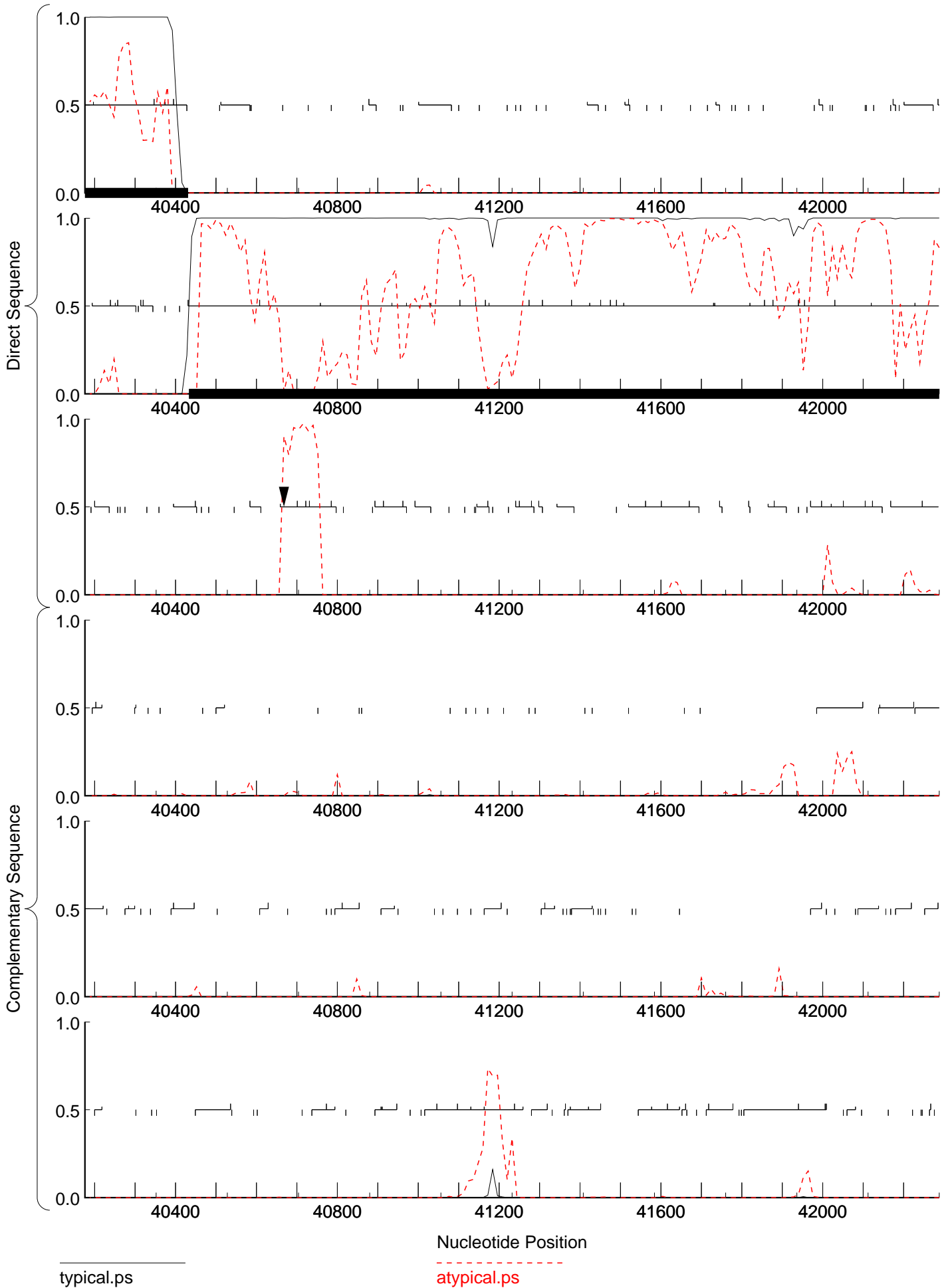


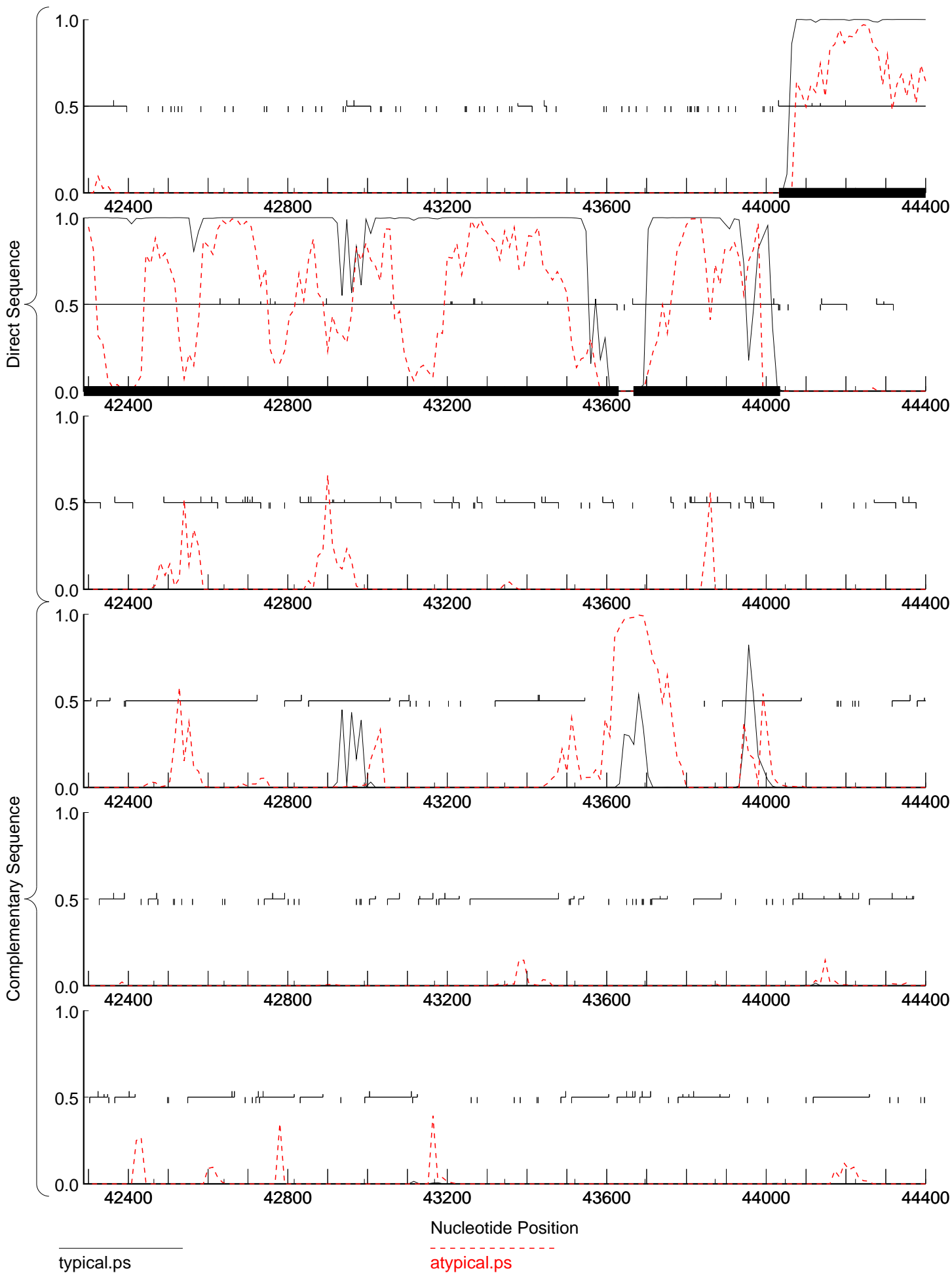


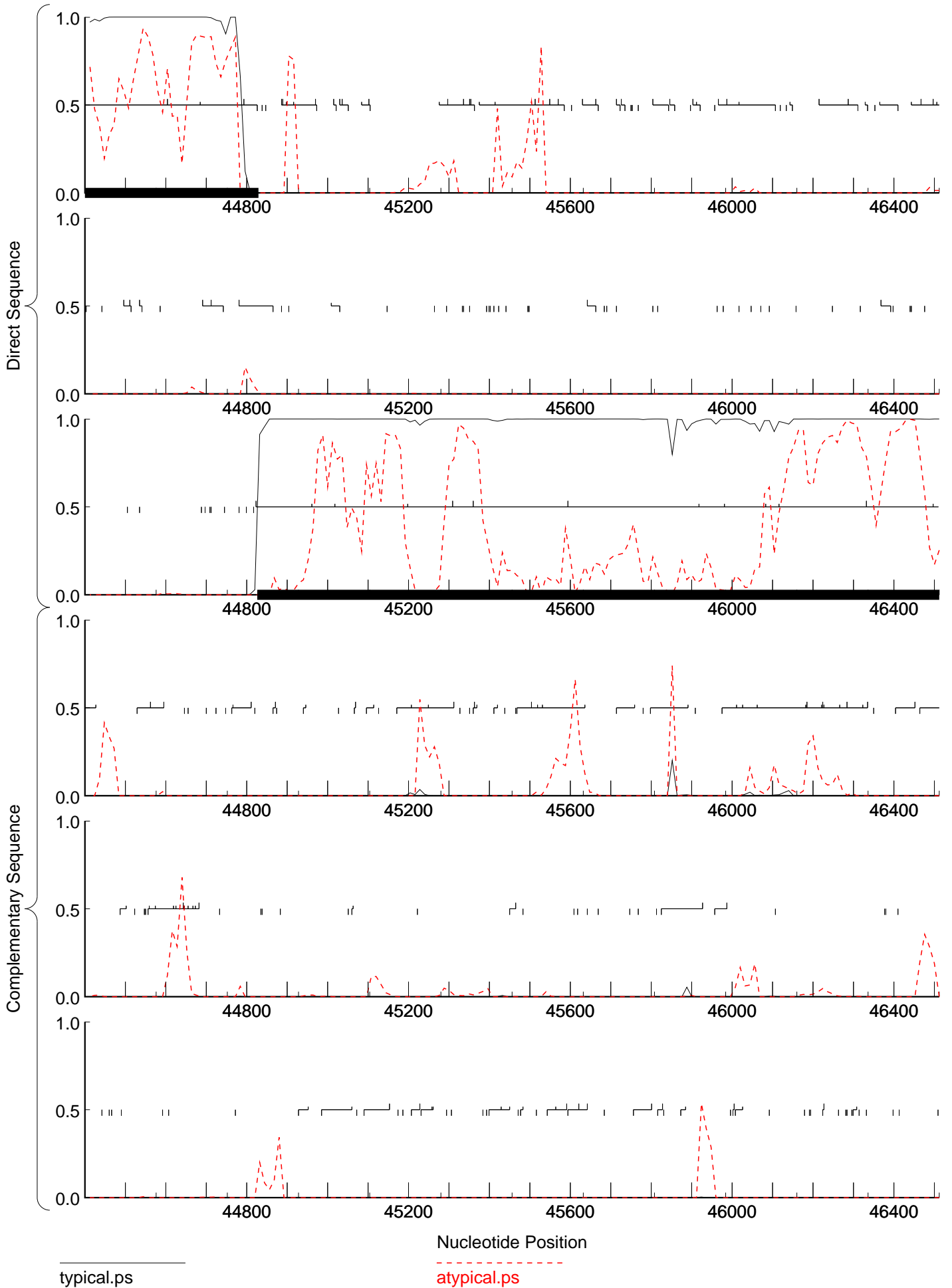






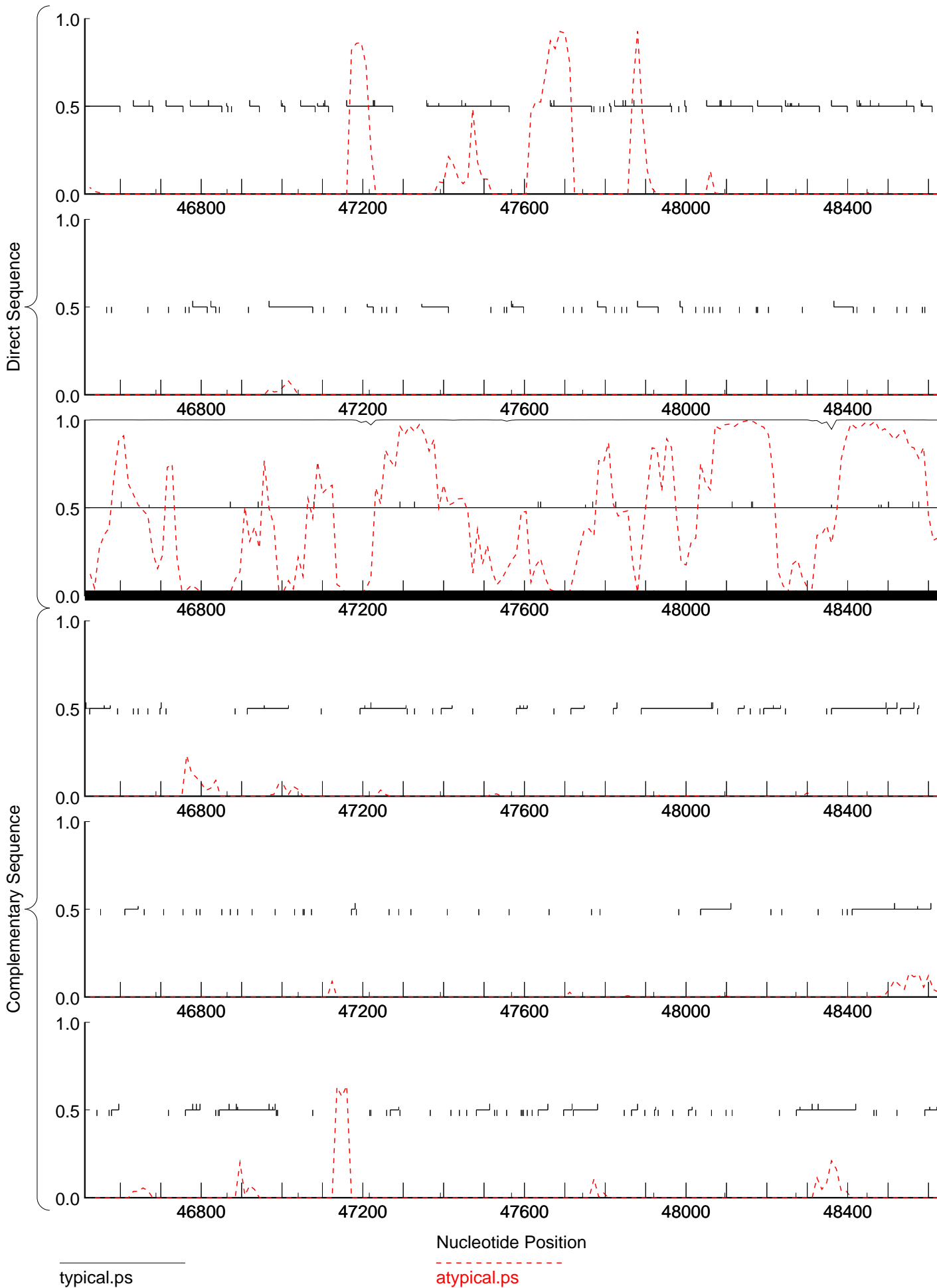




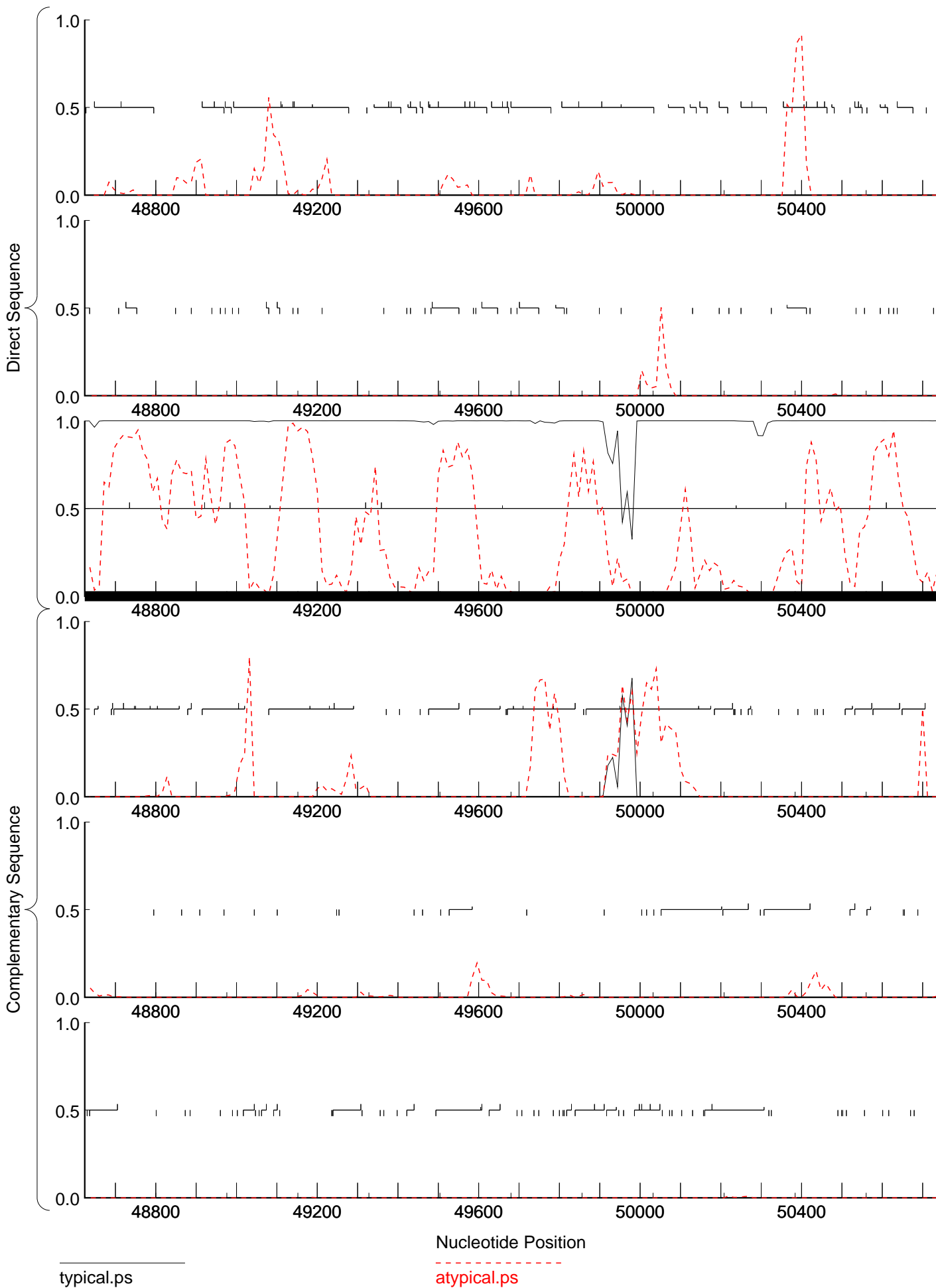


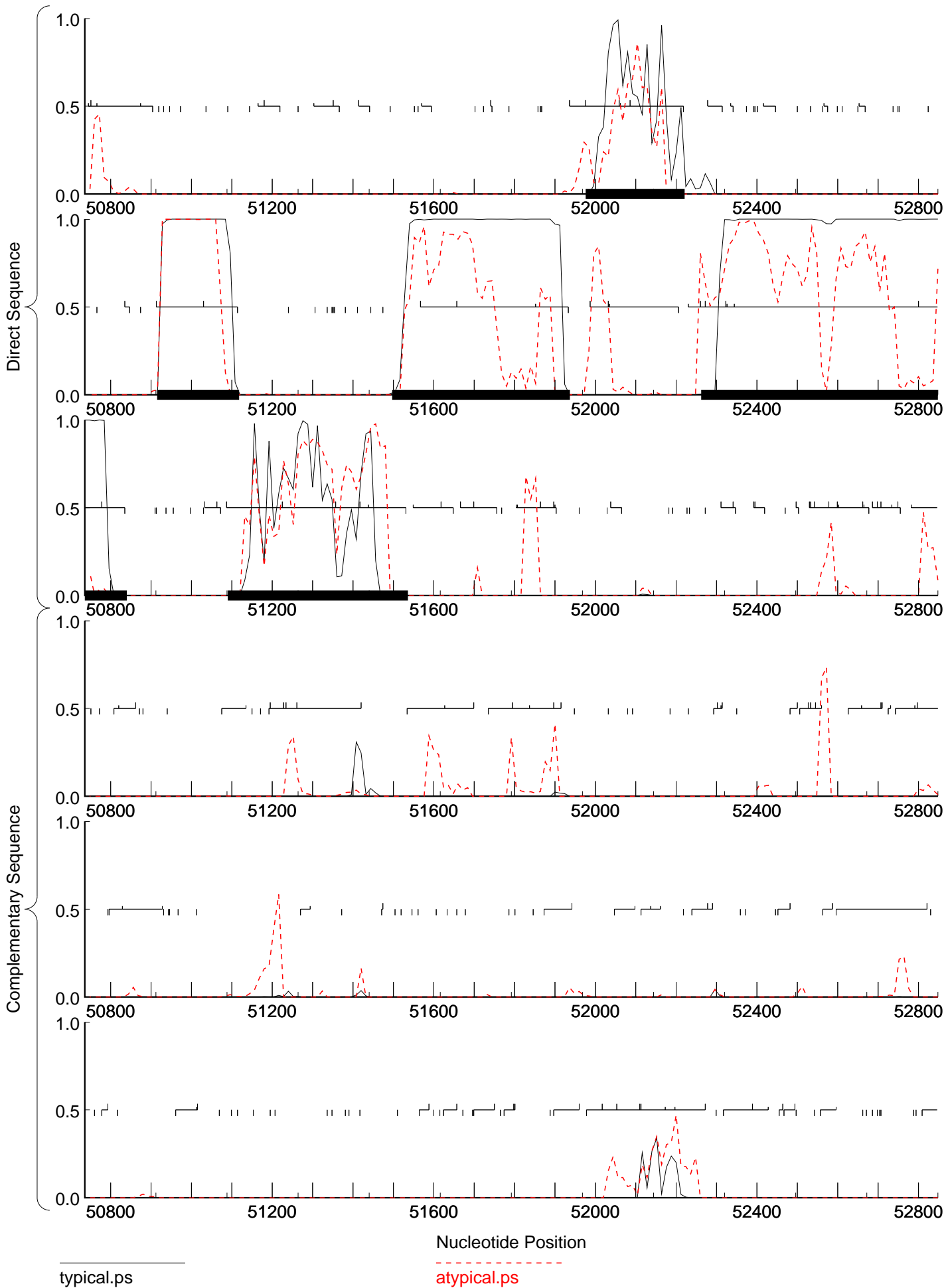
typical.ps

atypical.ps



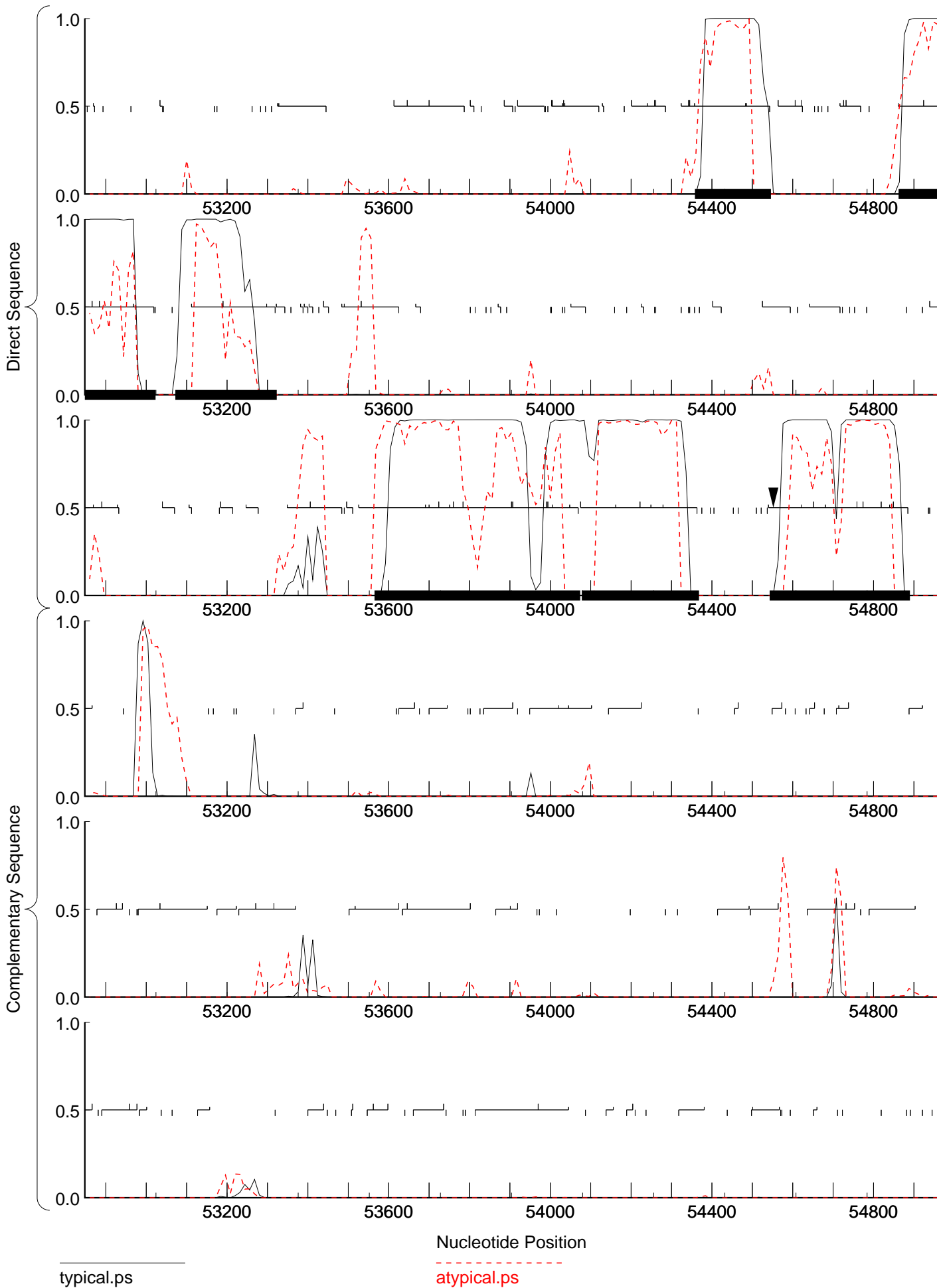






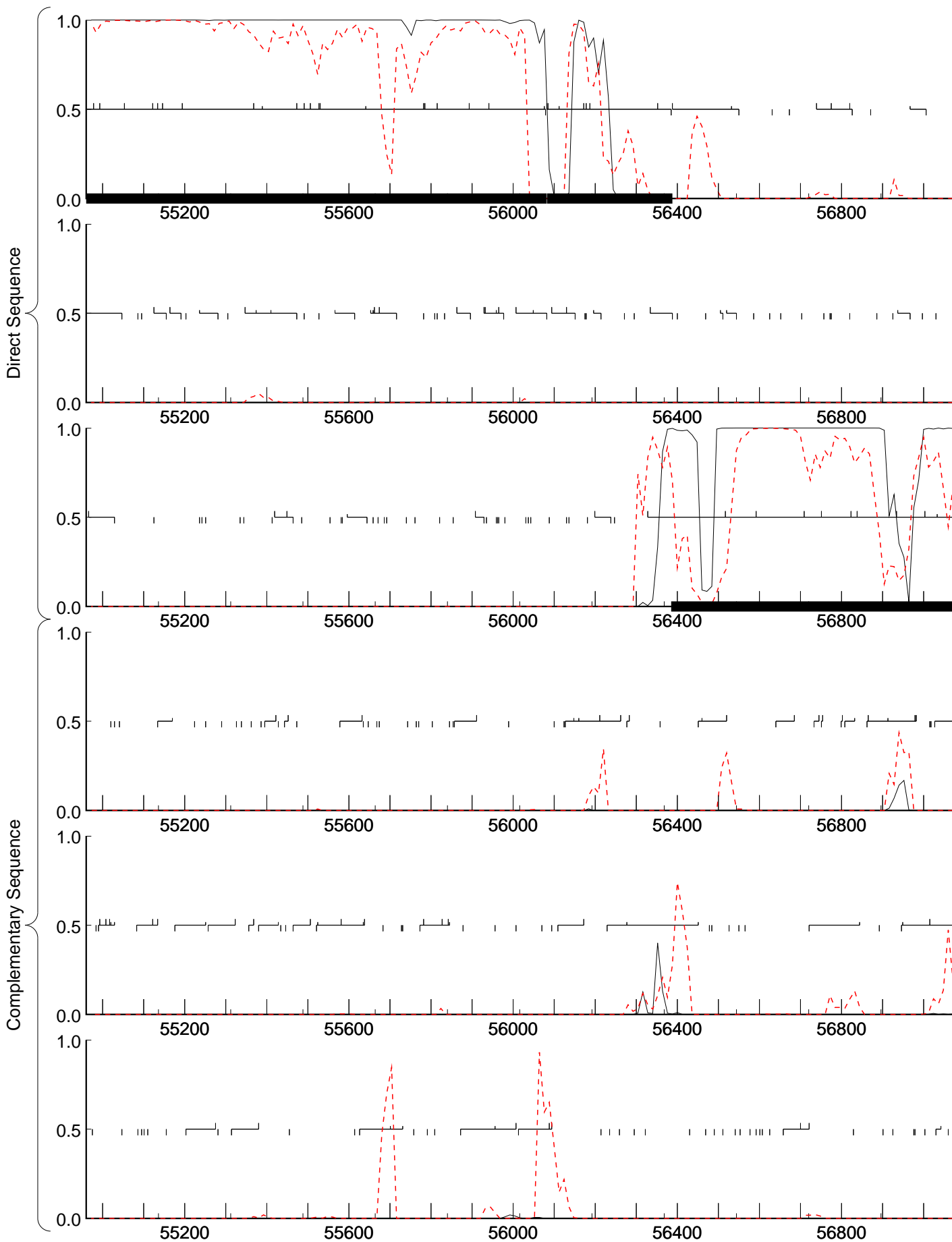
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atypical.ps



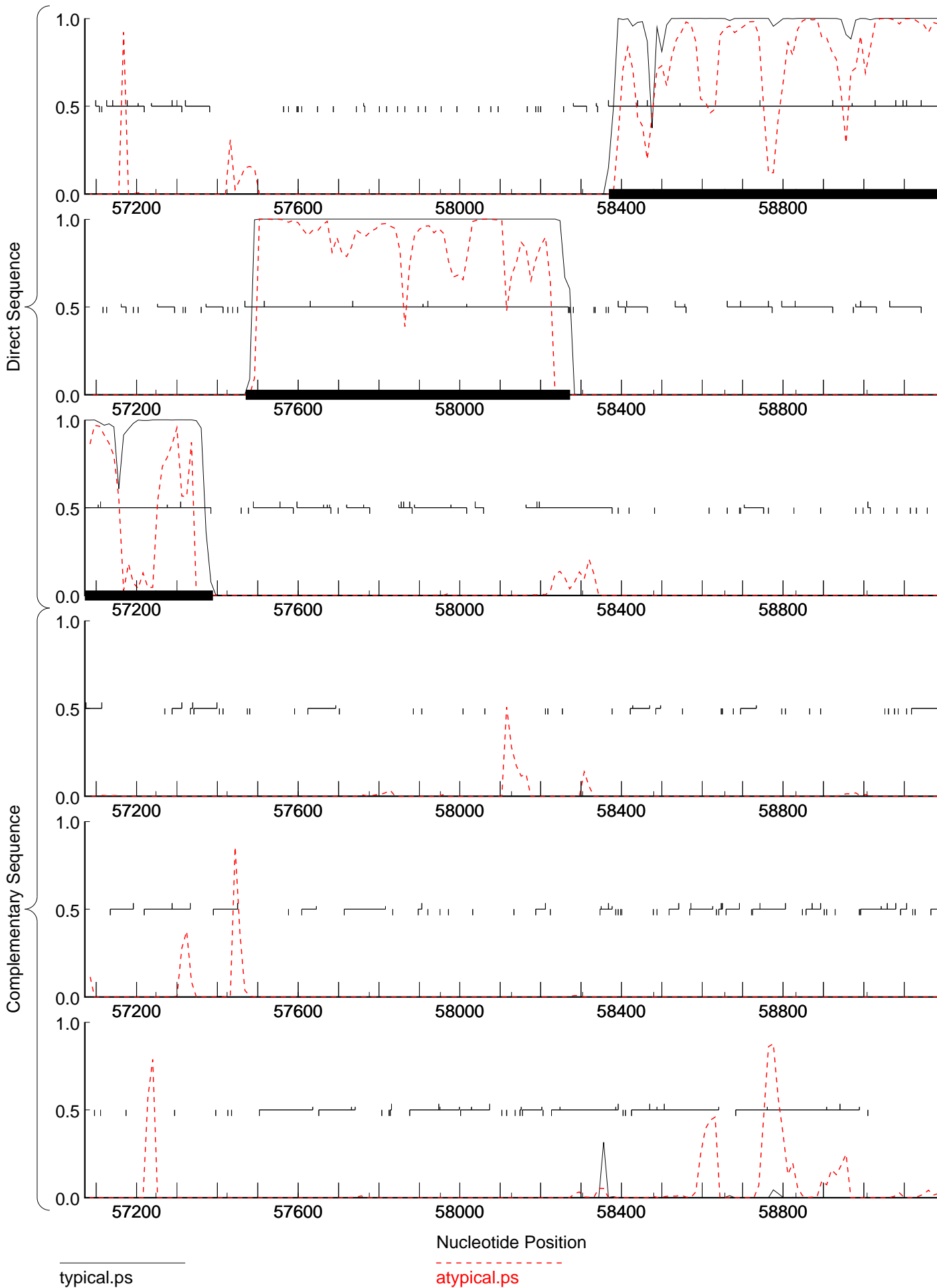
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atypical.ps



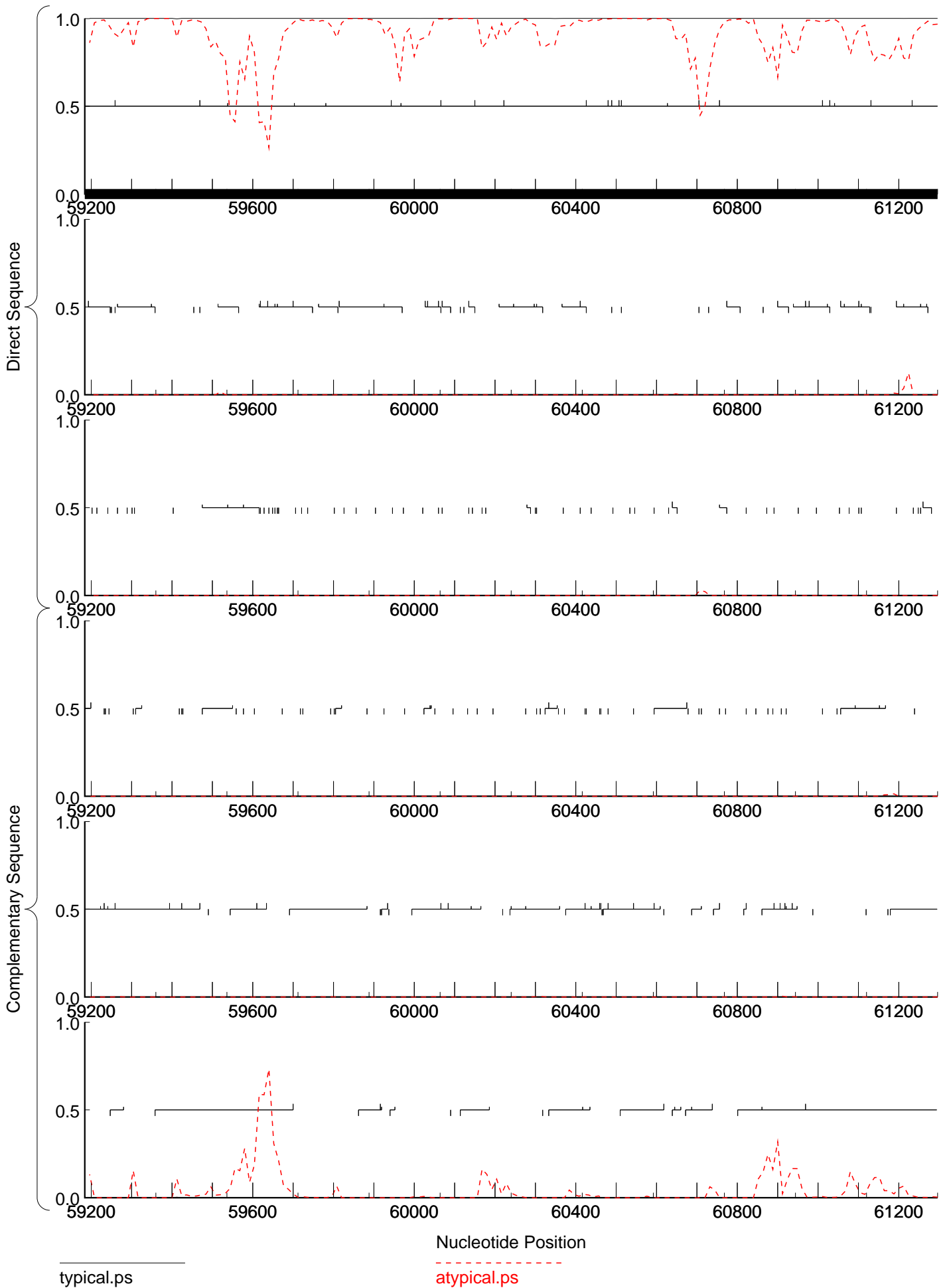
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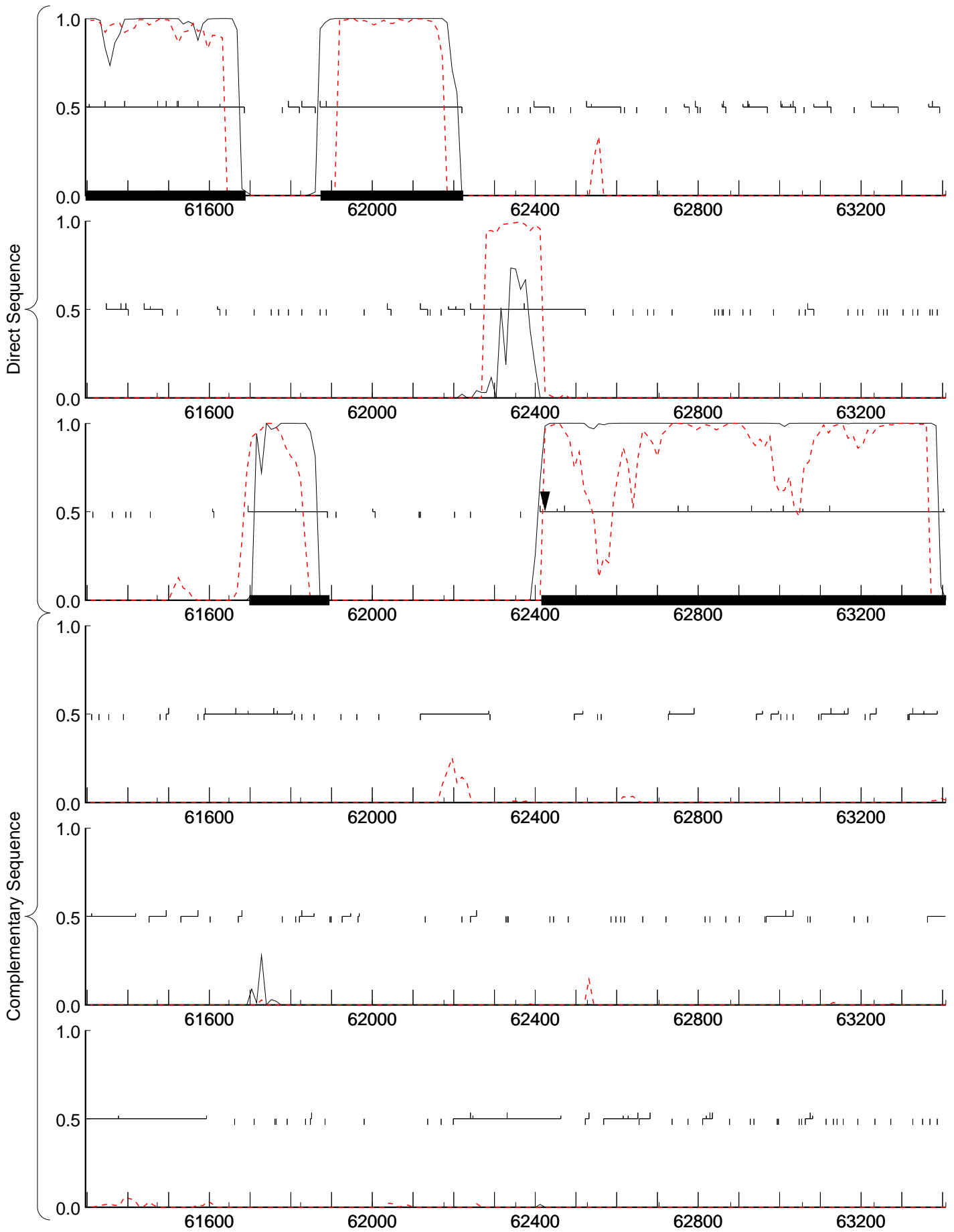
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typical.ps

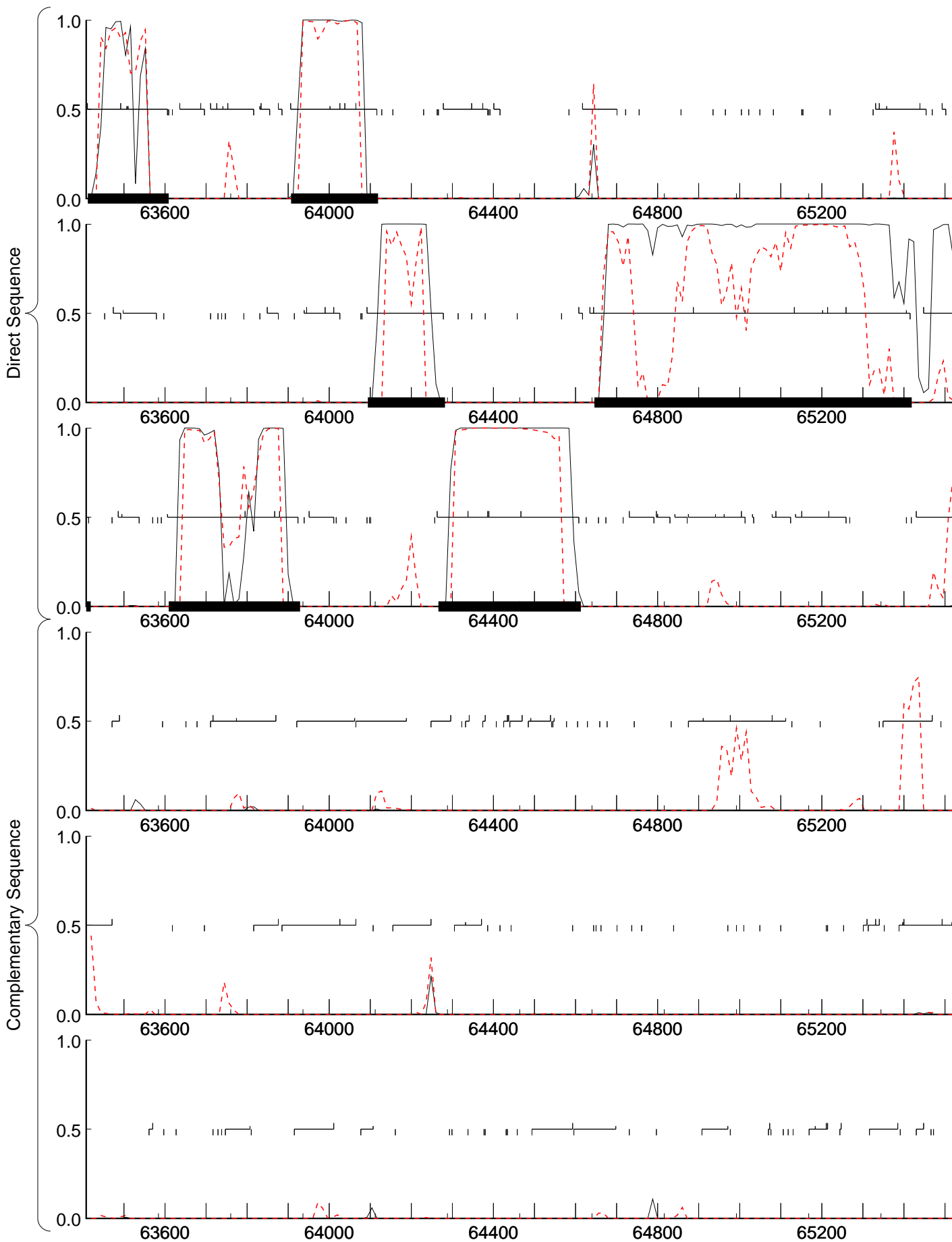
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typical.ps

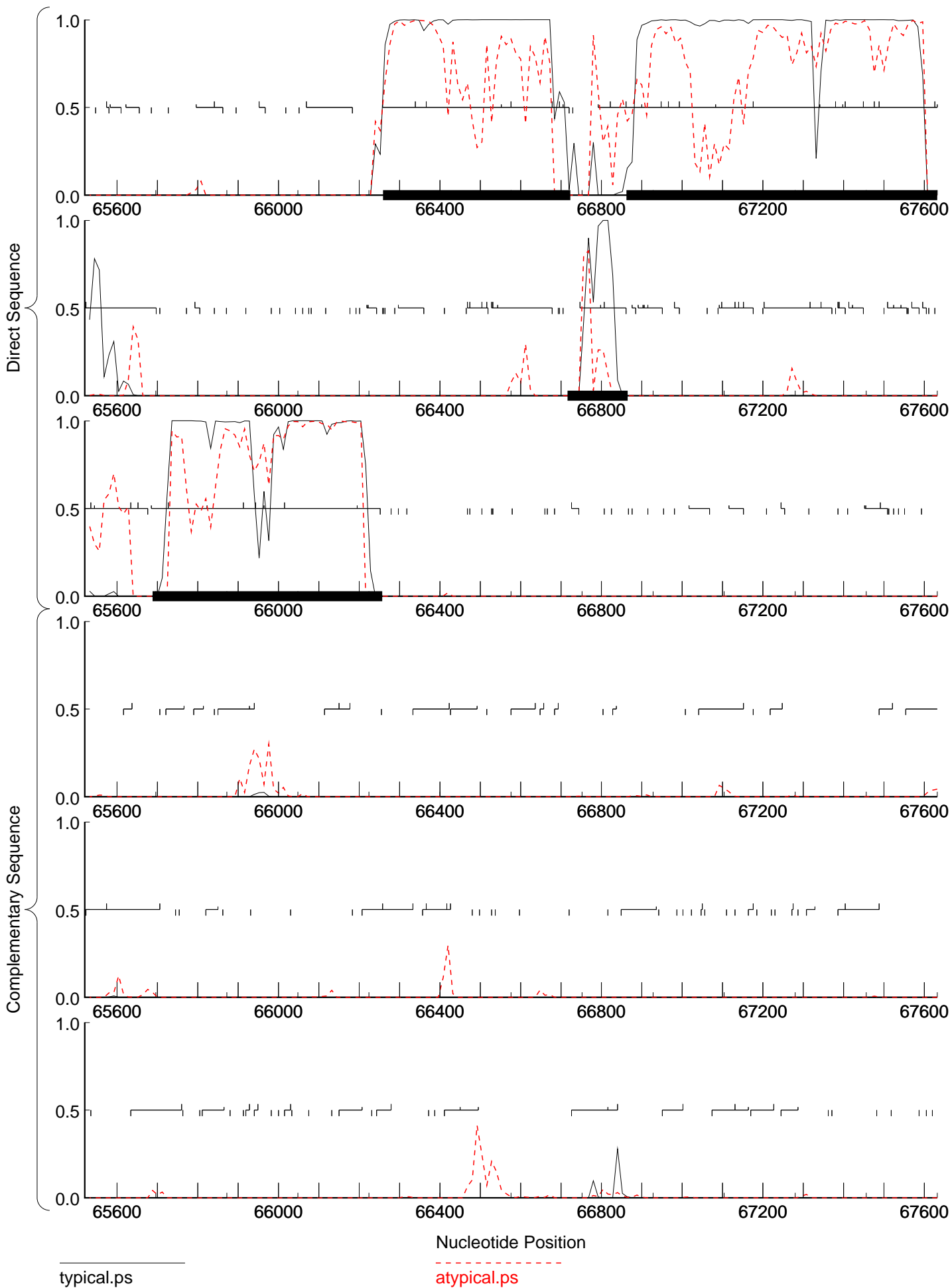
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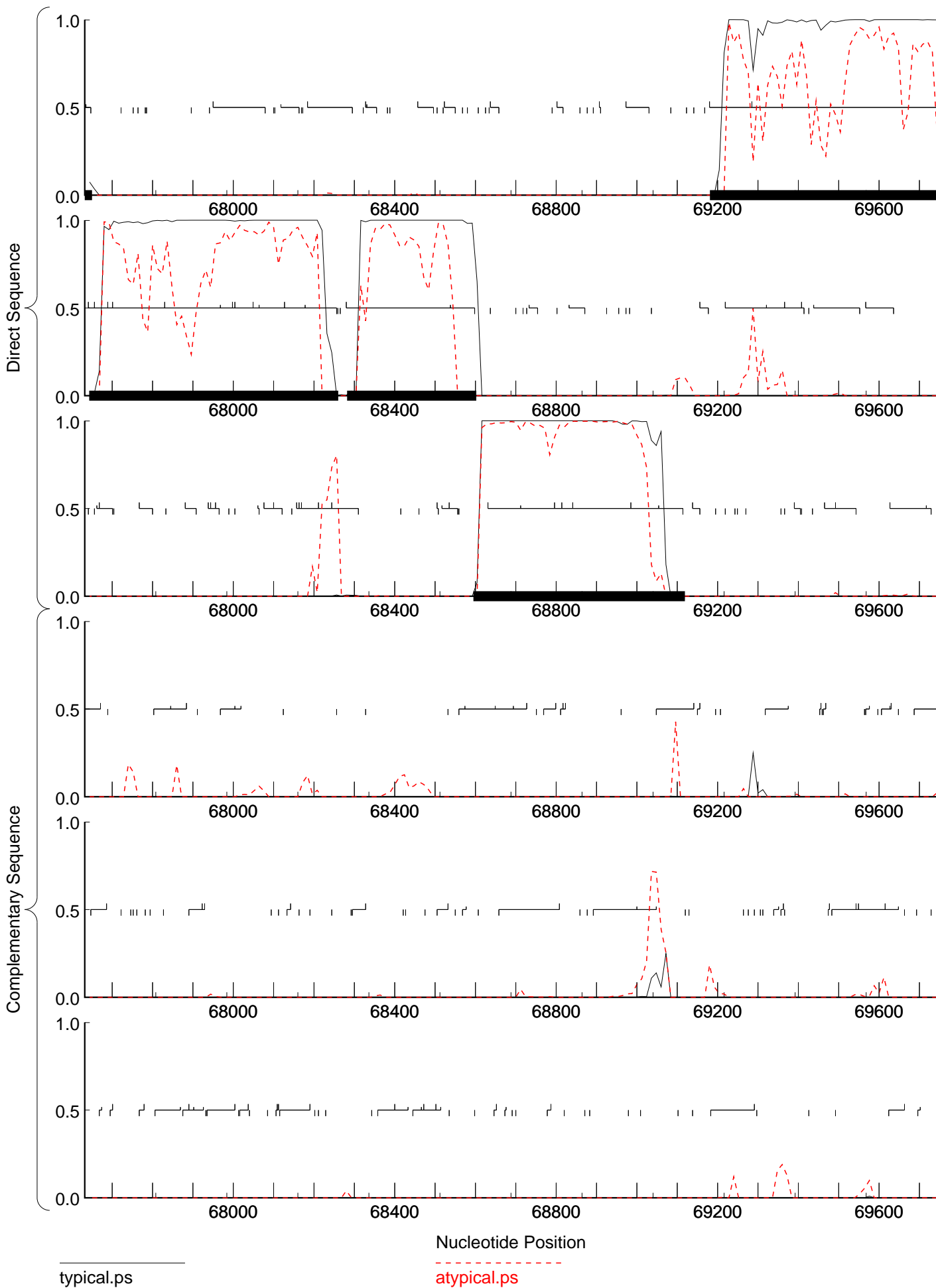


typical.ps

atypical.ps

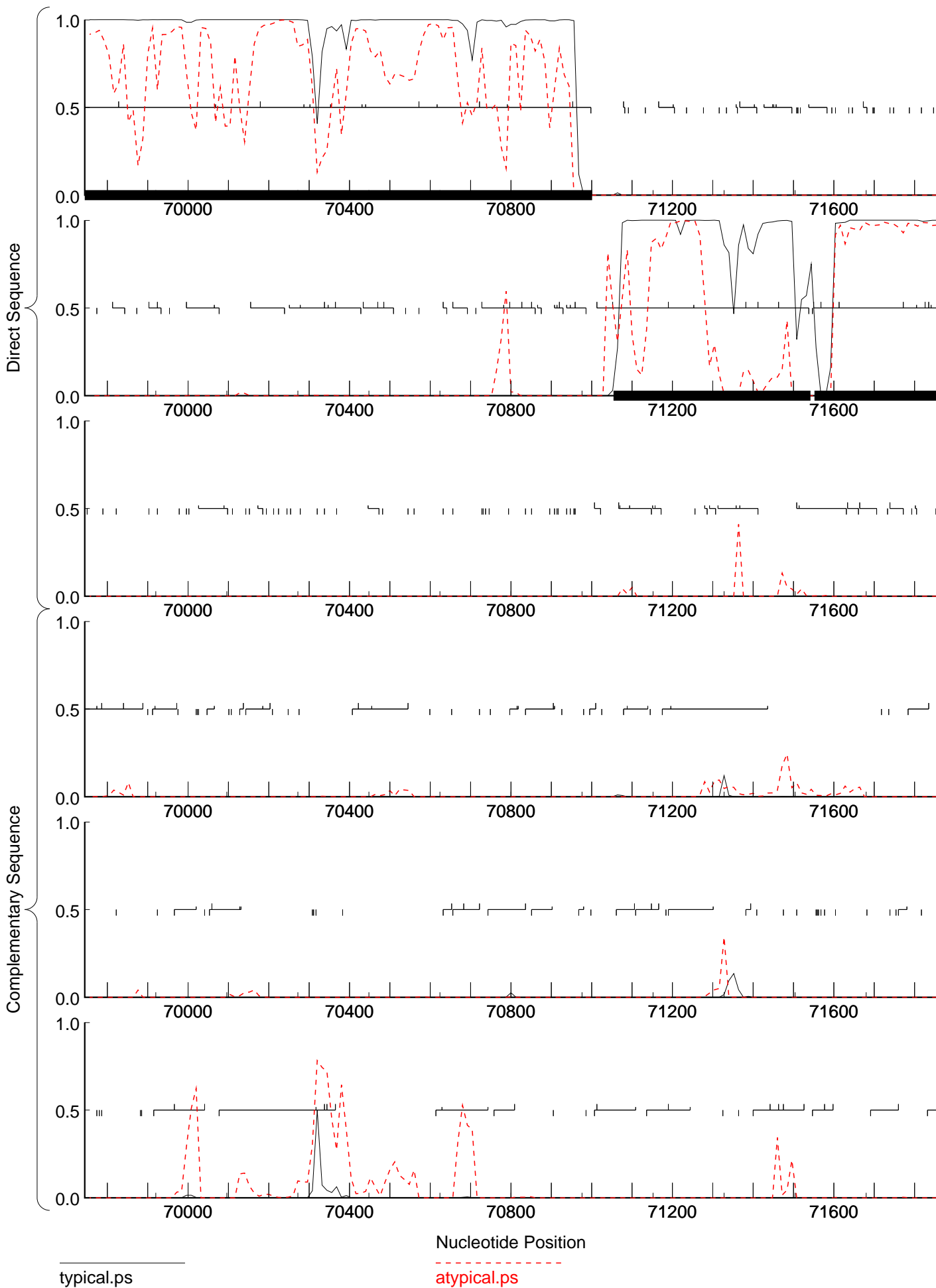


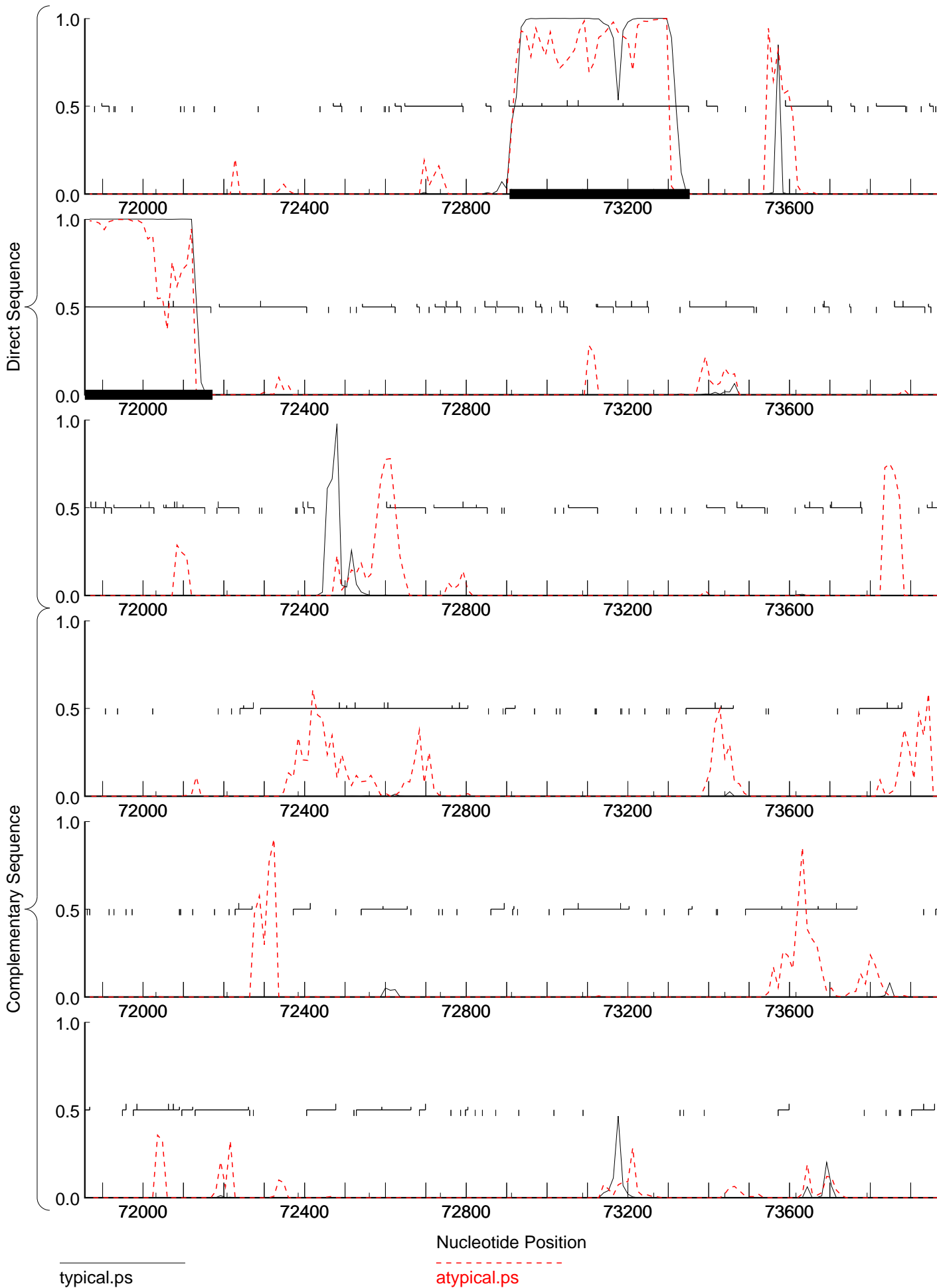


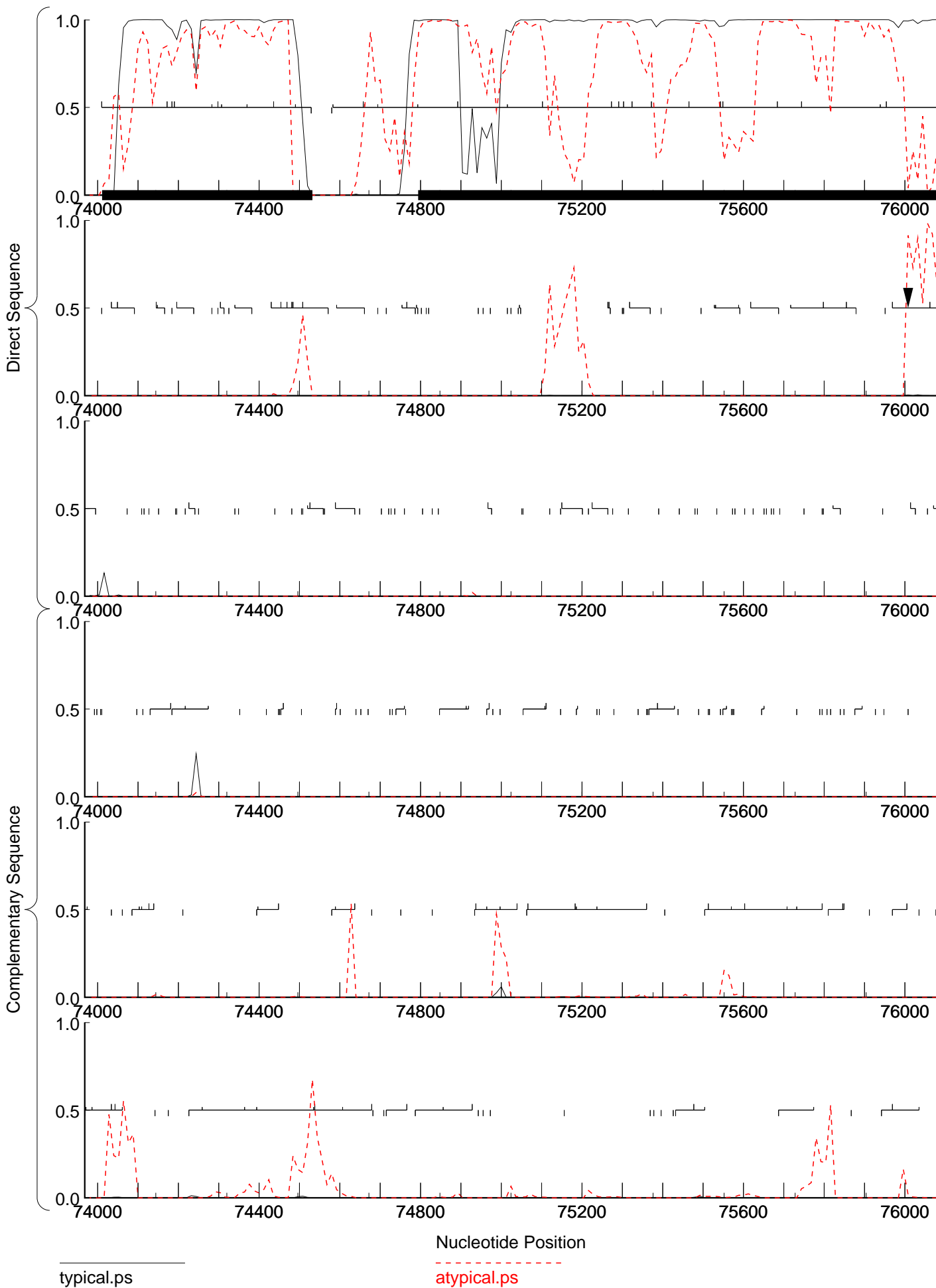


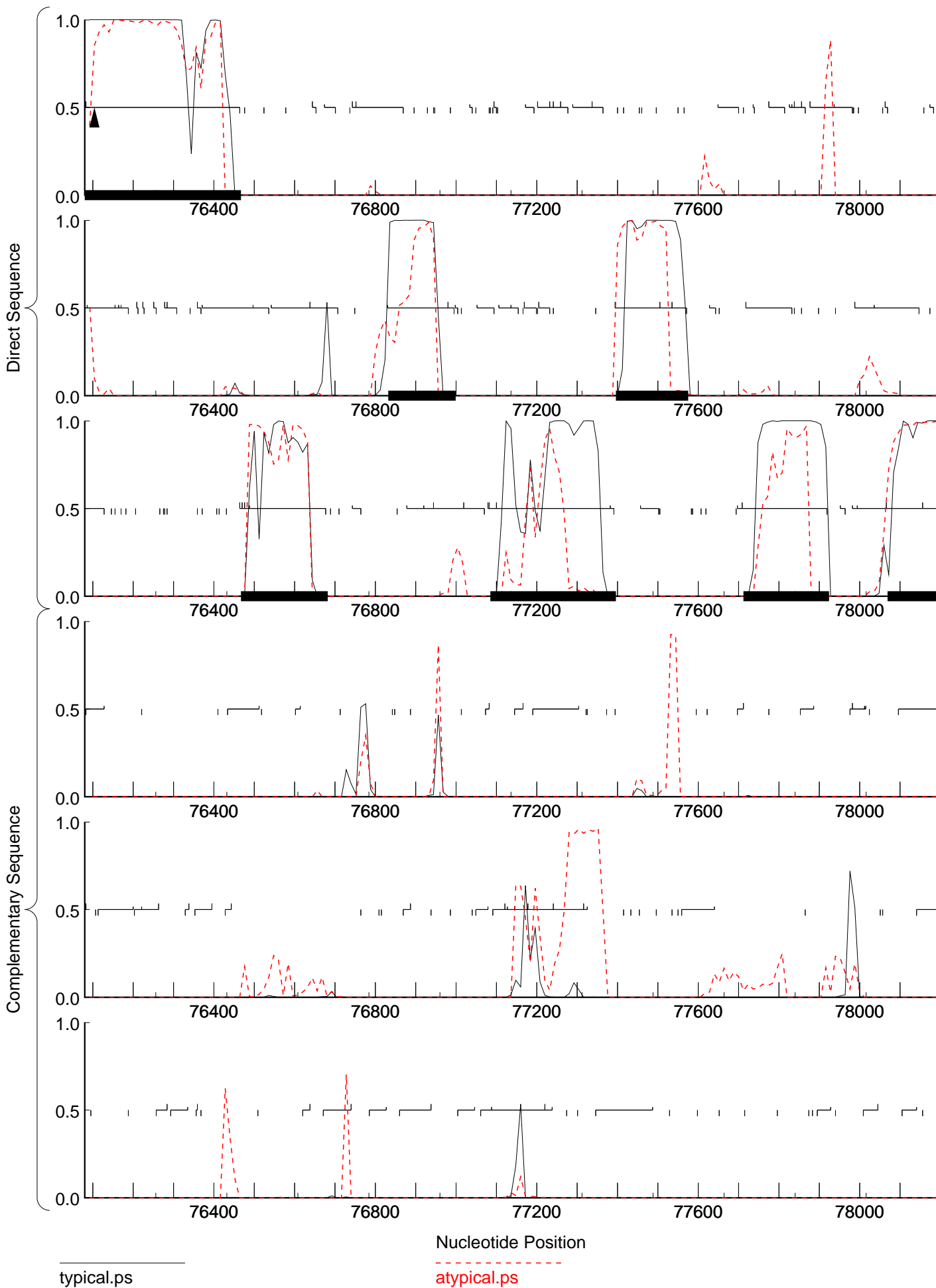
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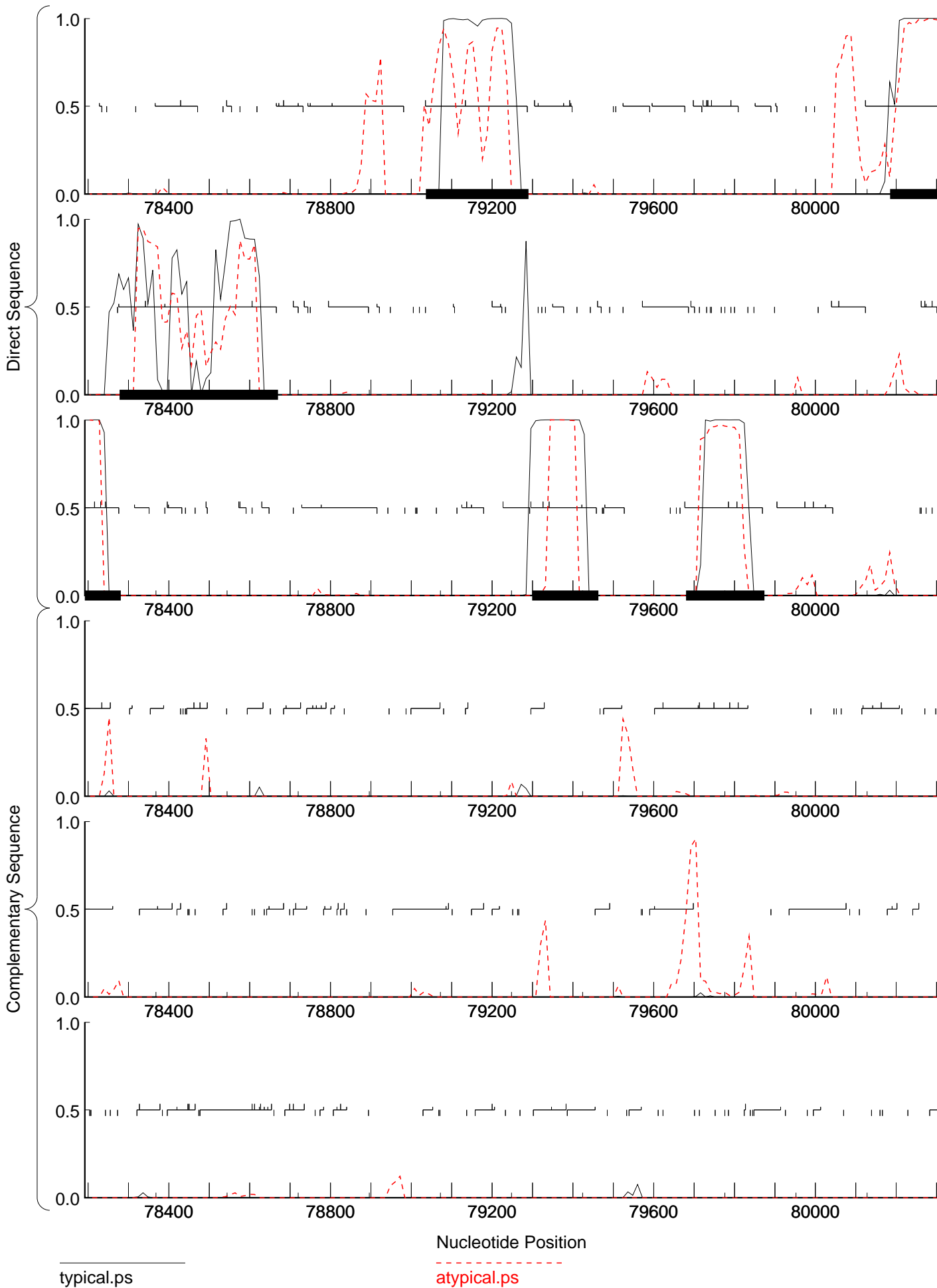
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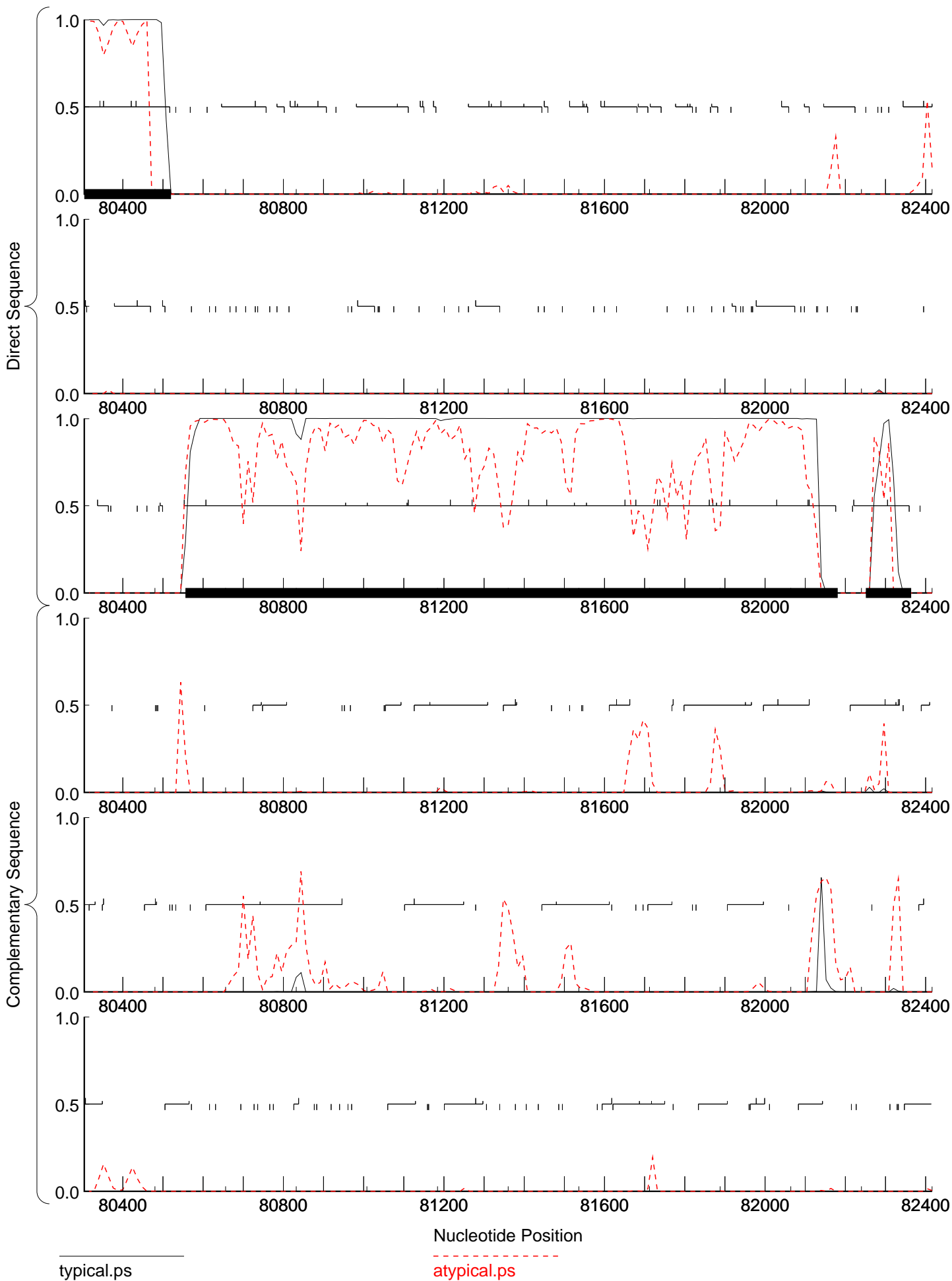








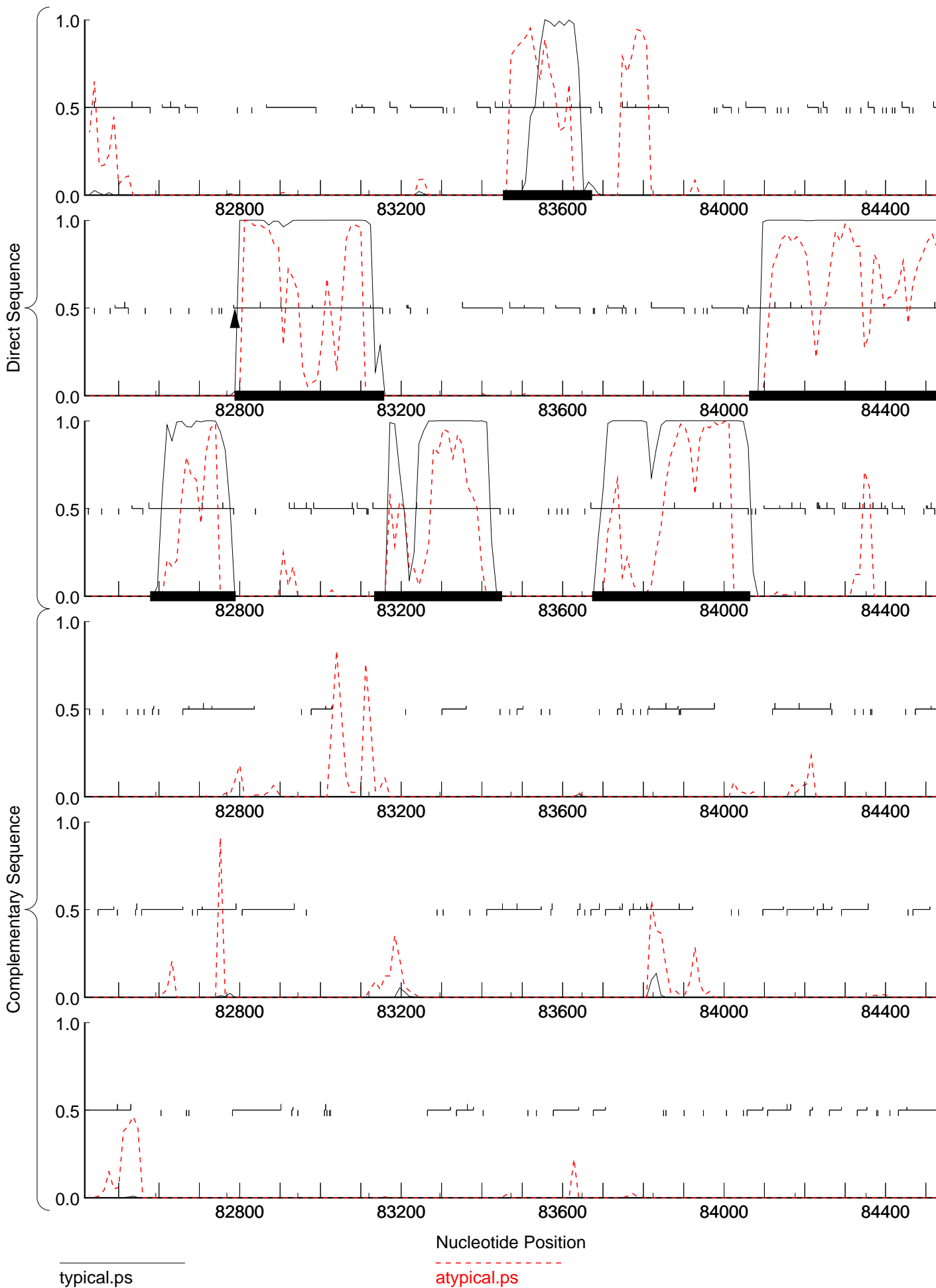


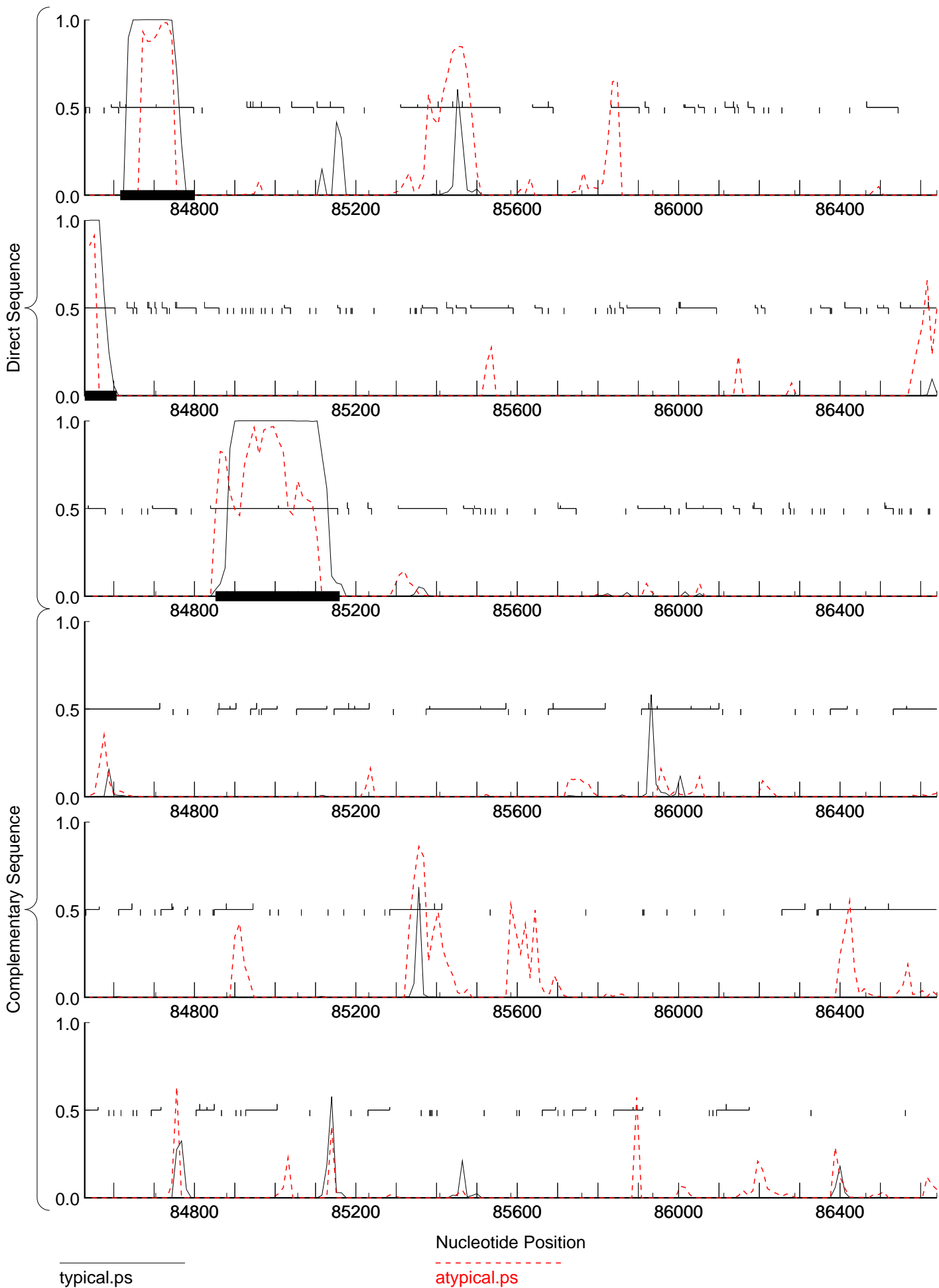


typical.ps

atypical.ps

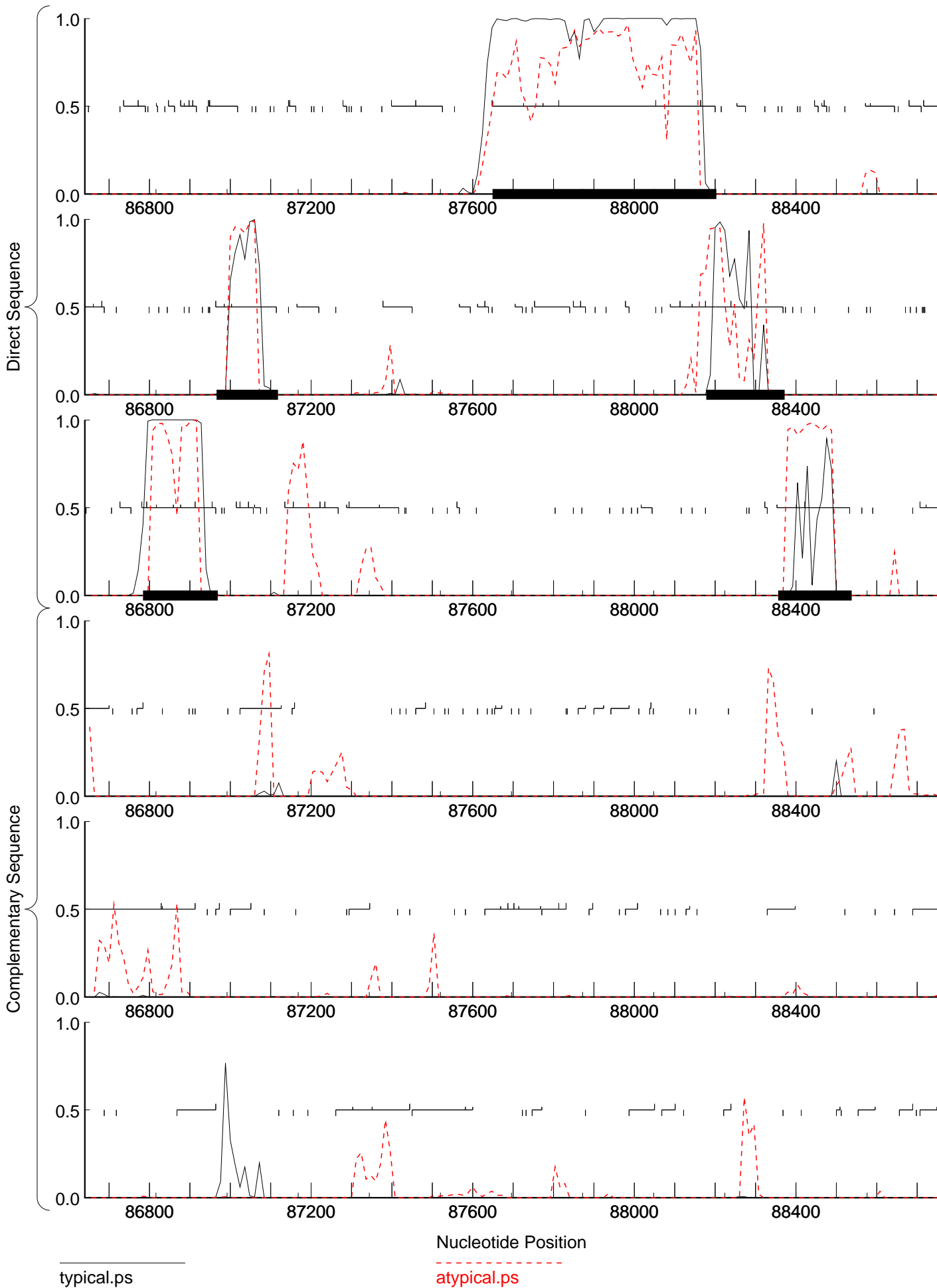






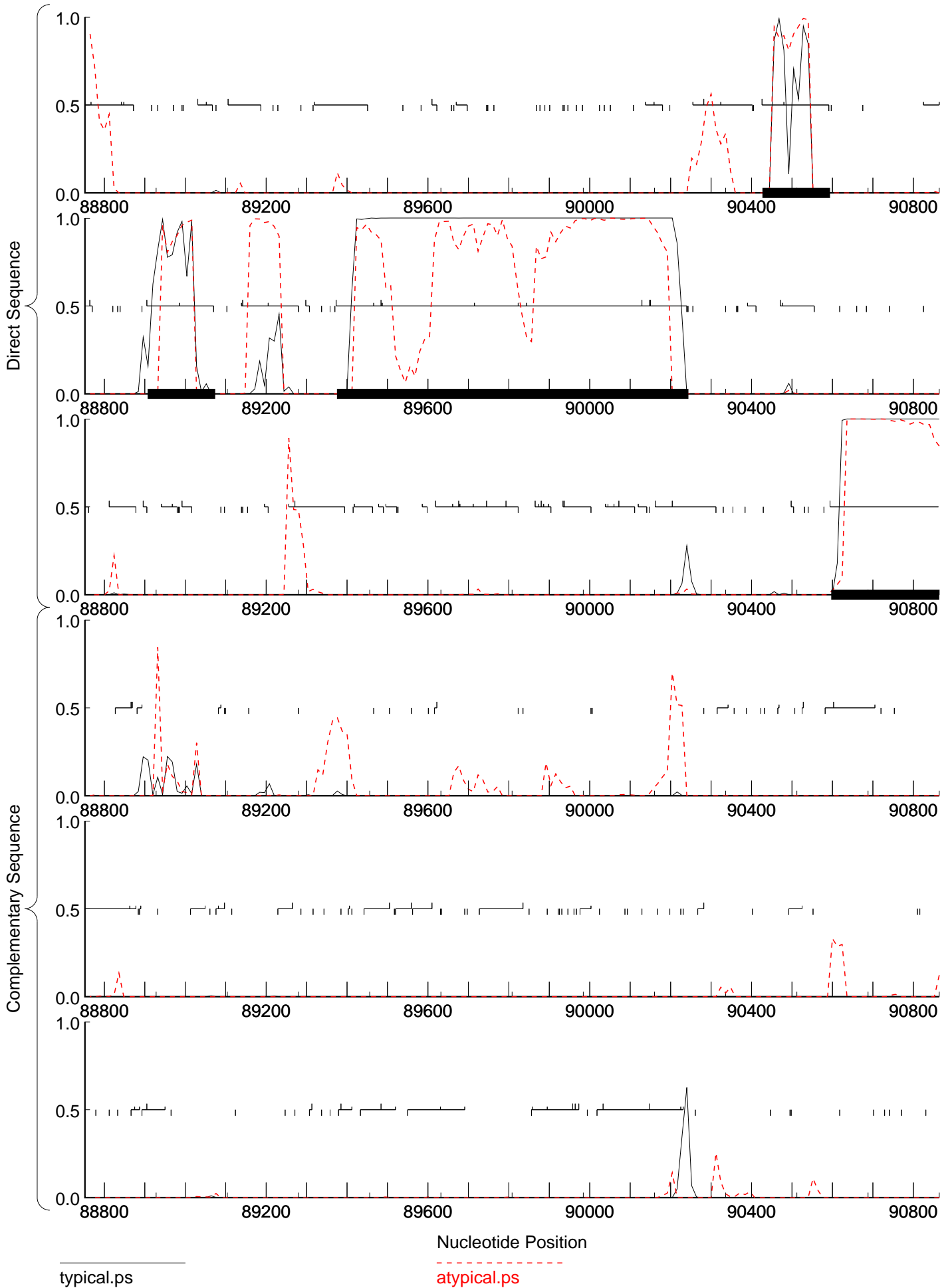
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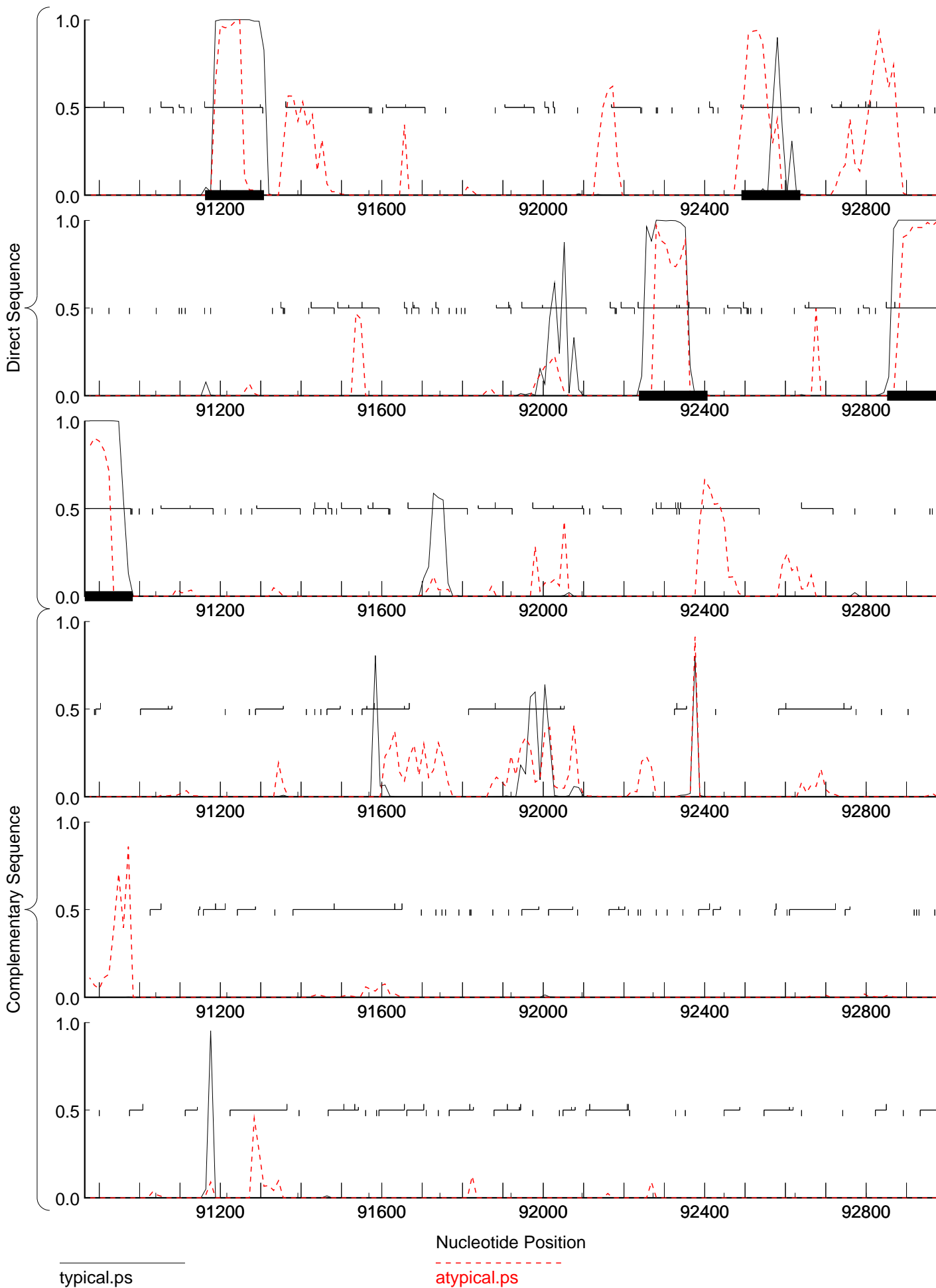
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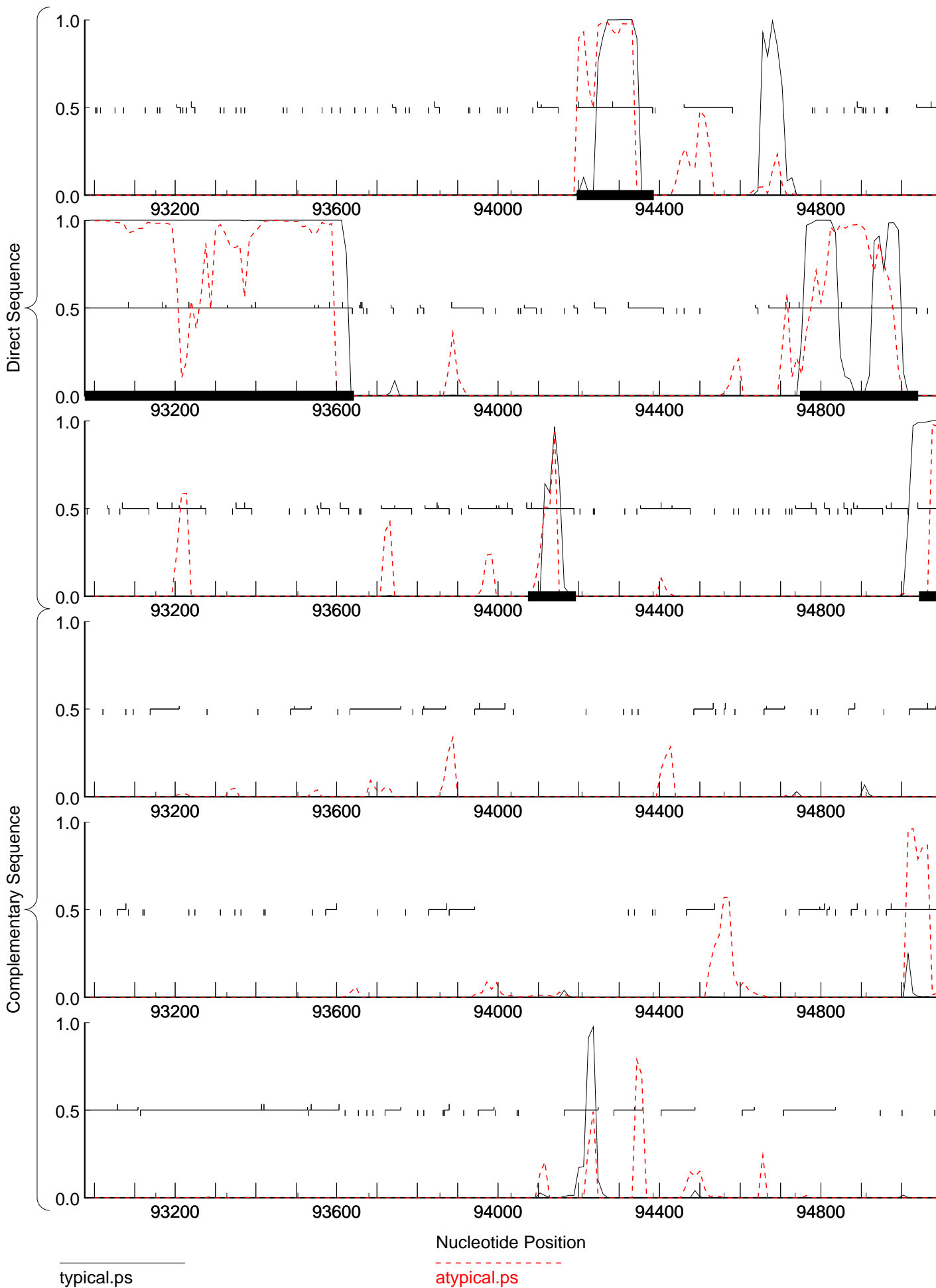


typical.ps

atypical.ps

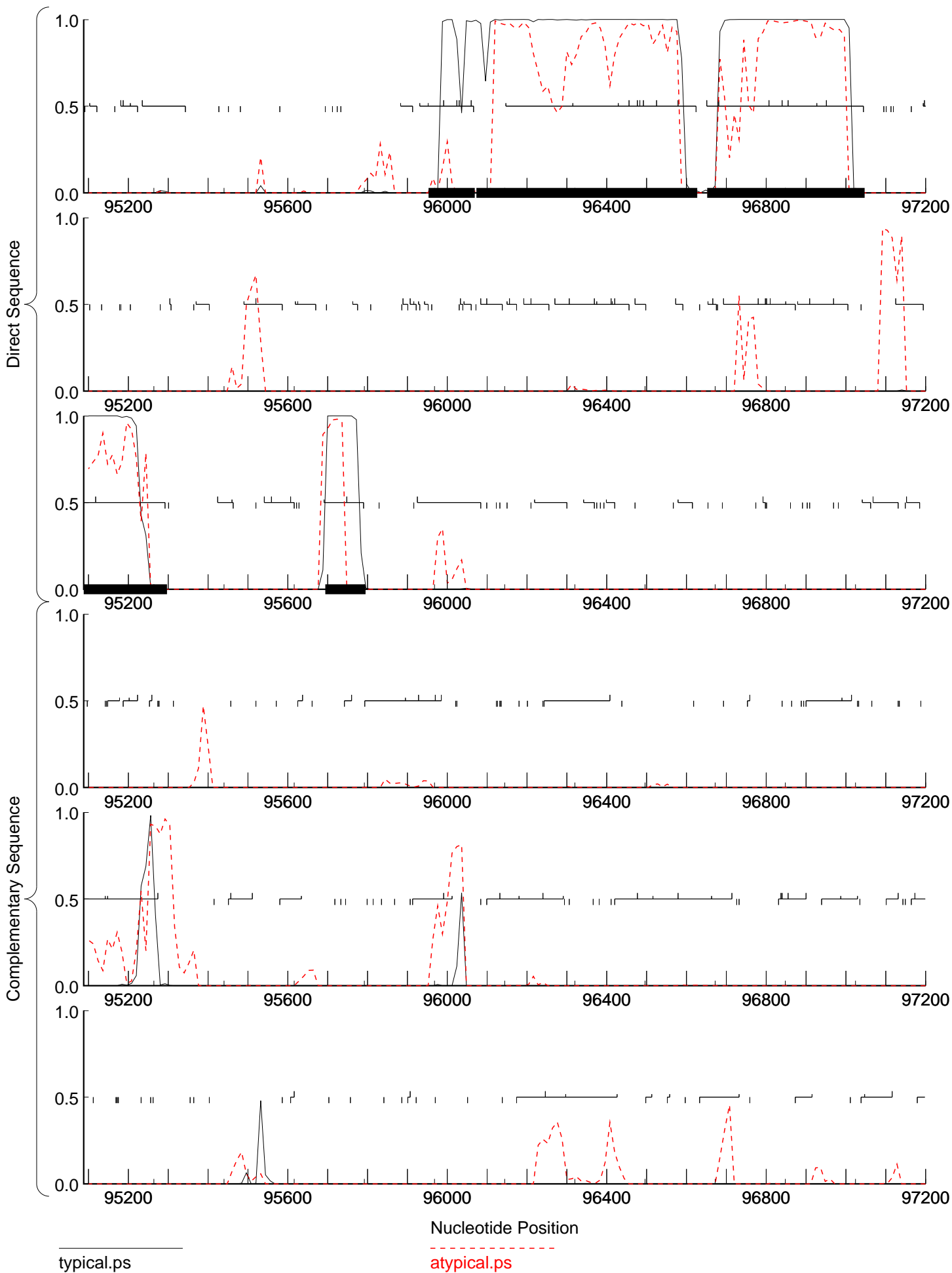


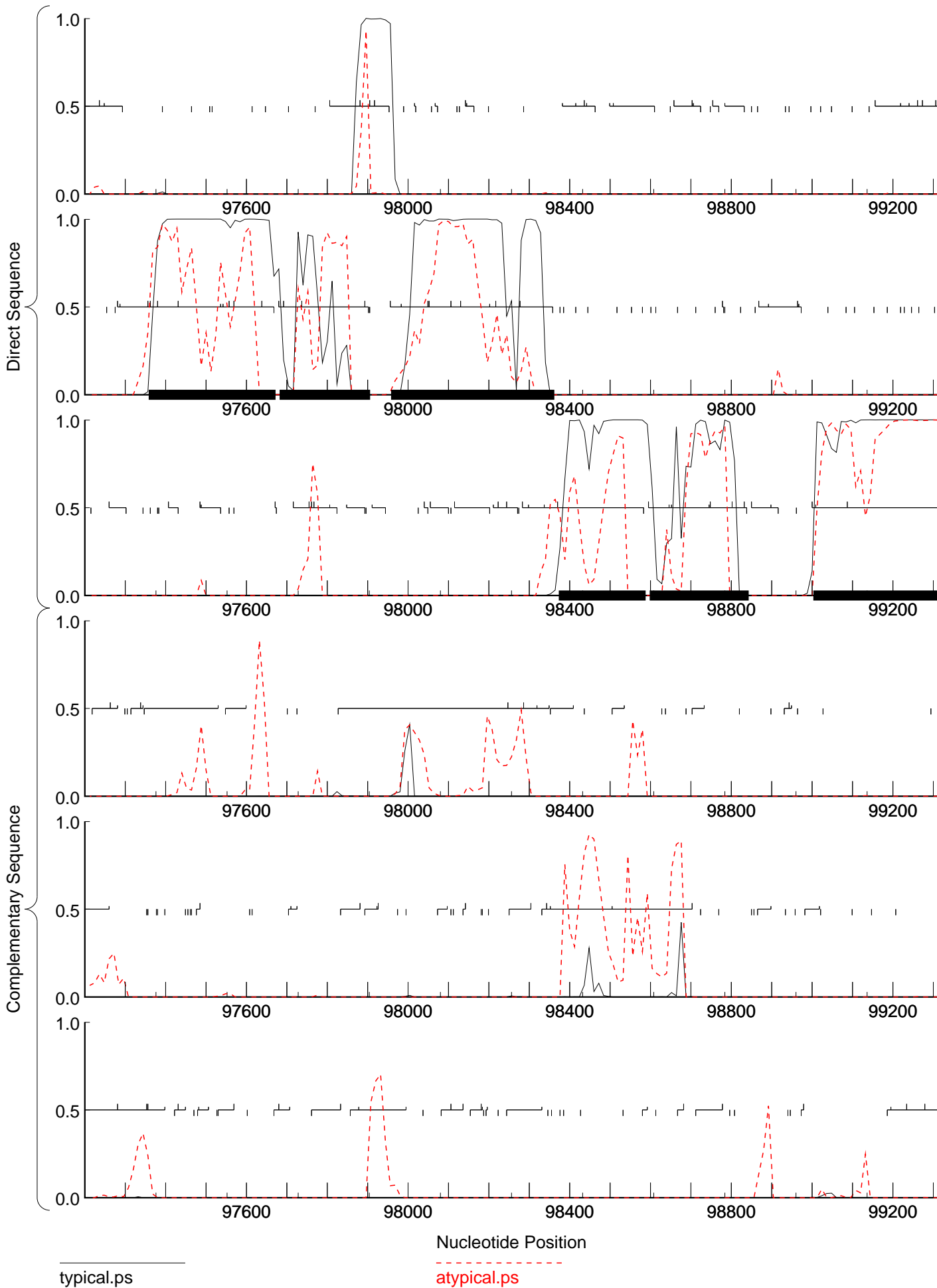




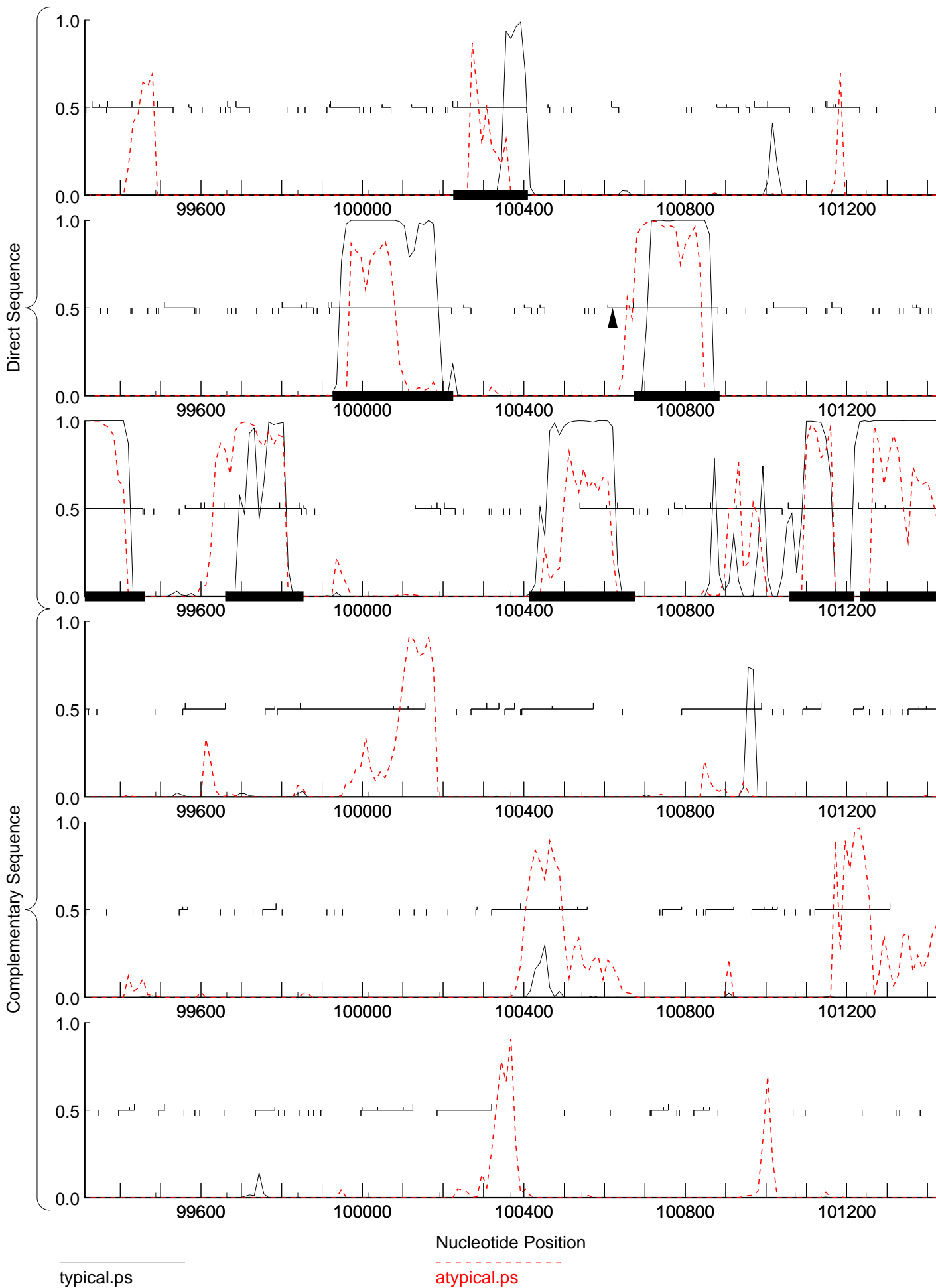
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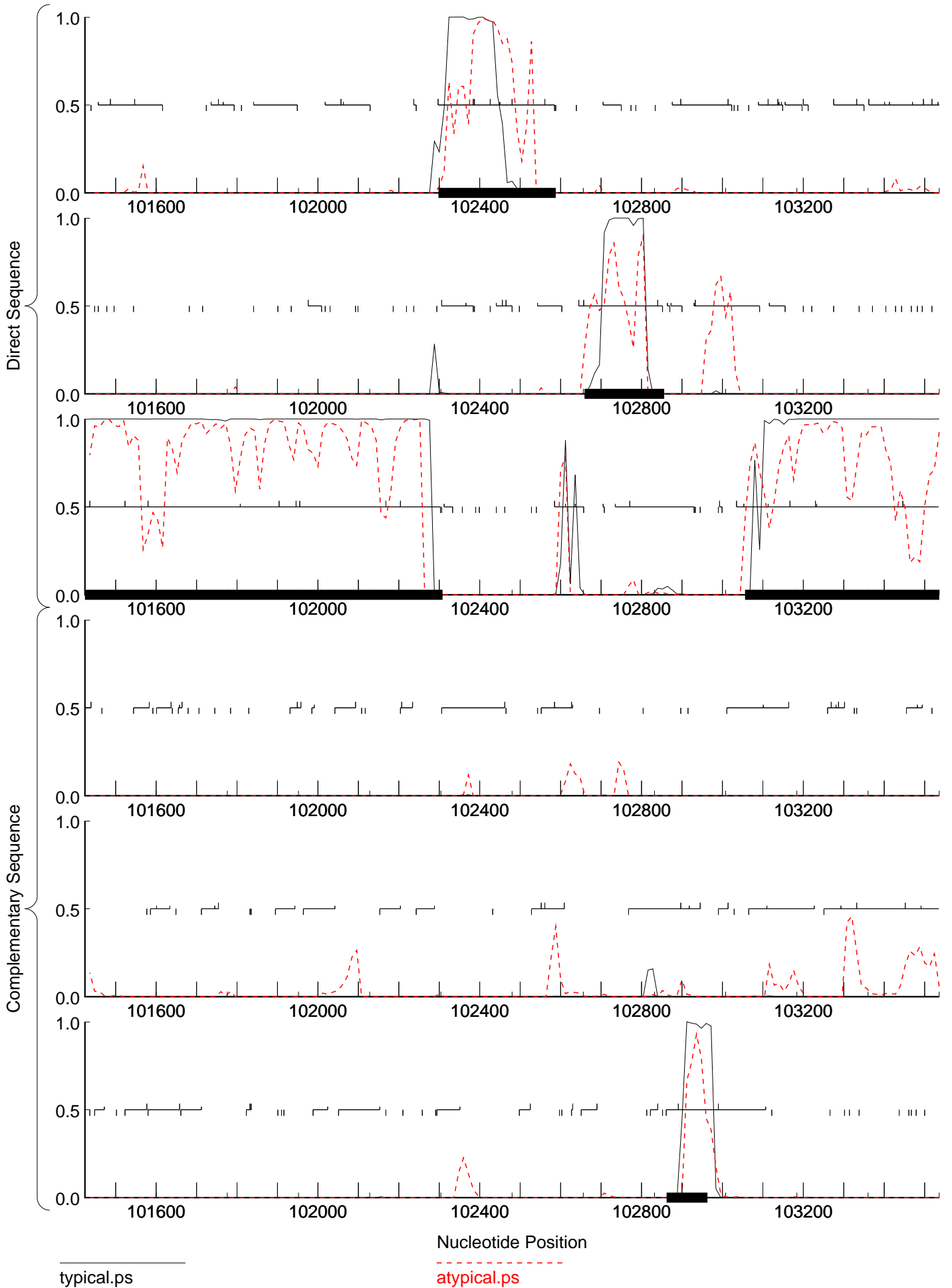






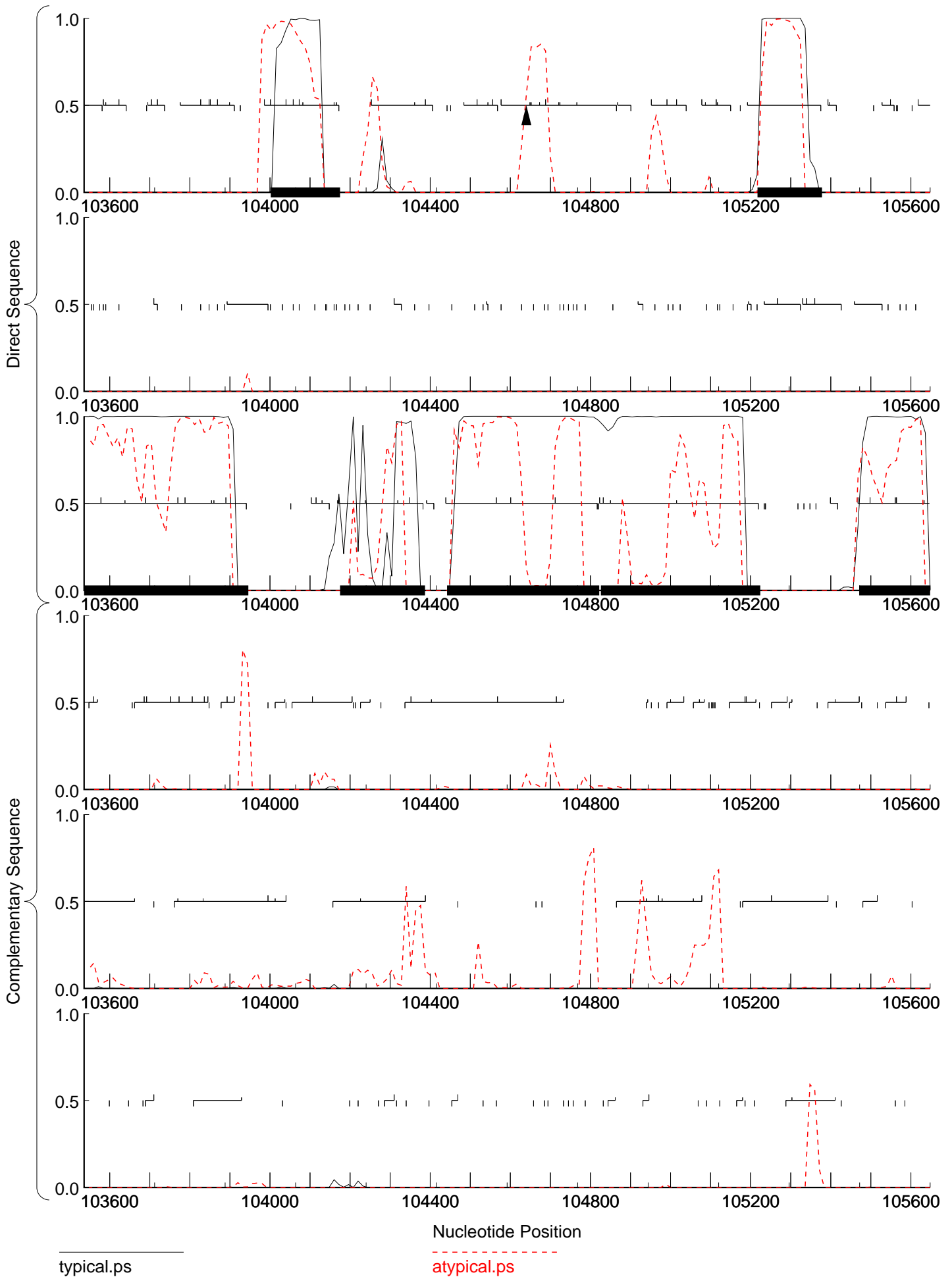
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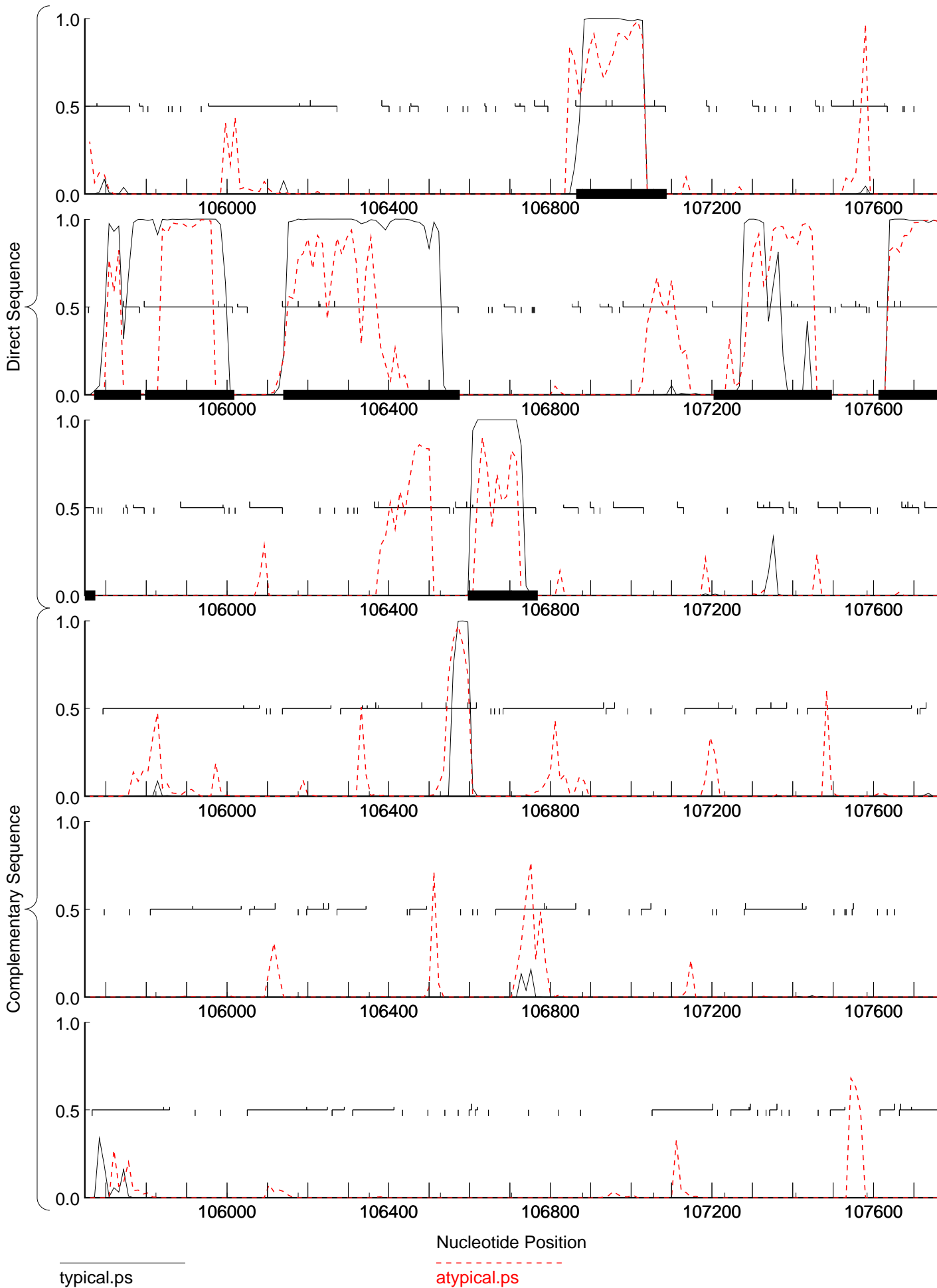
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typical.ps

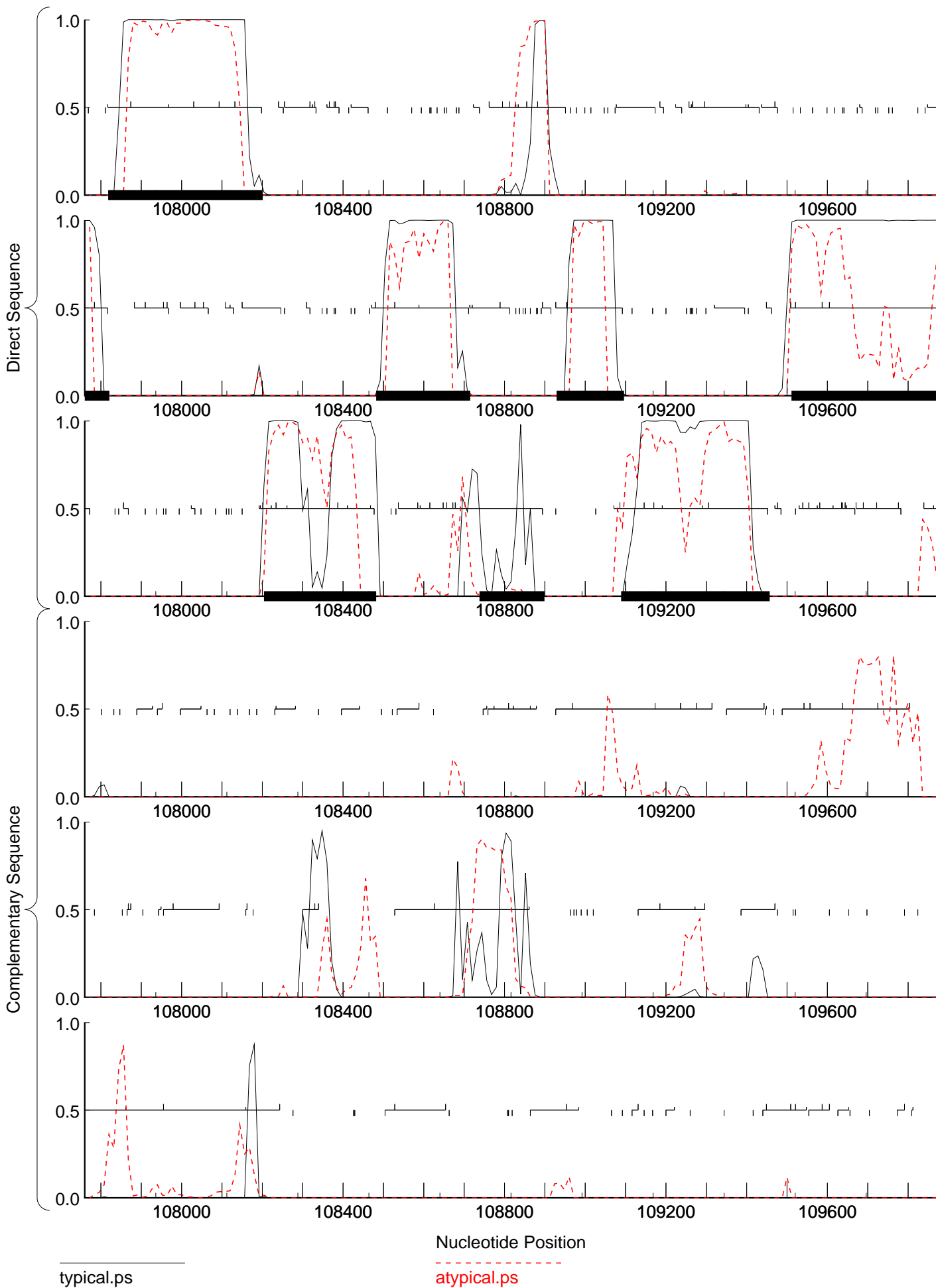
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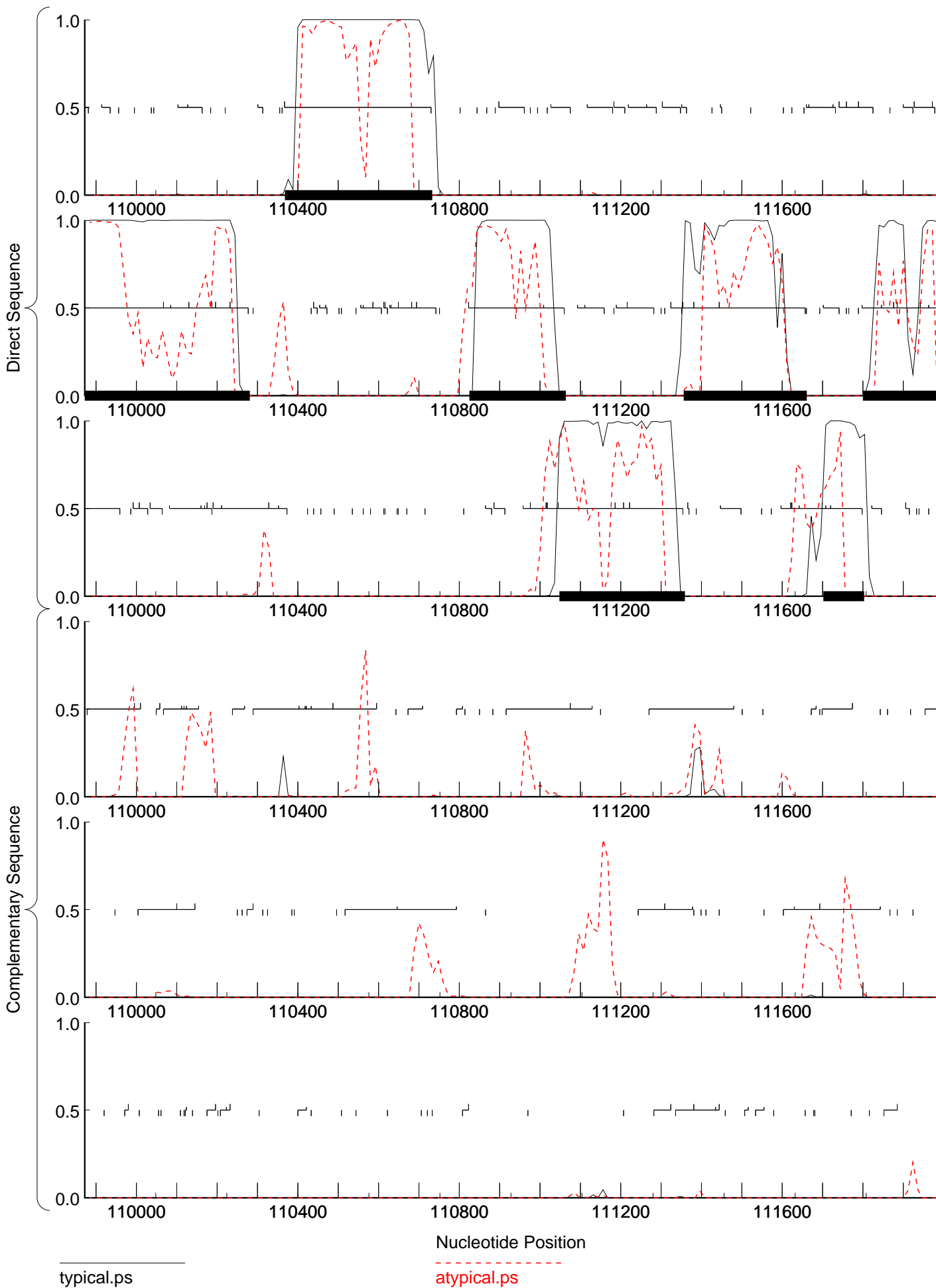
typical.ps

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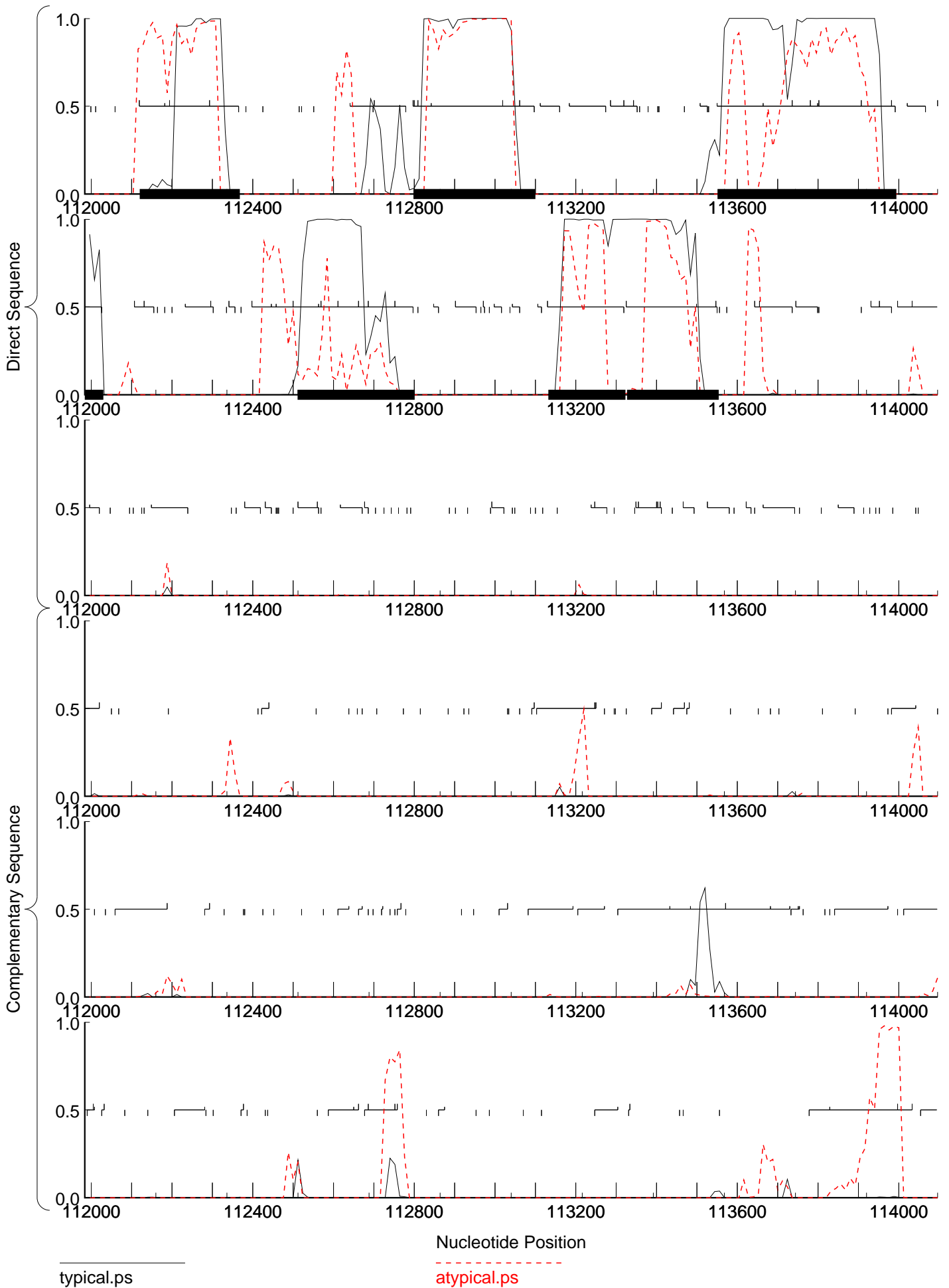
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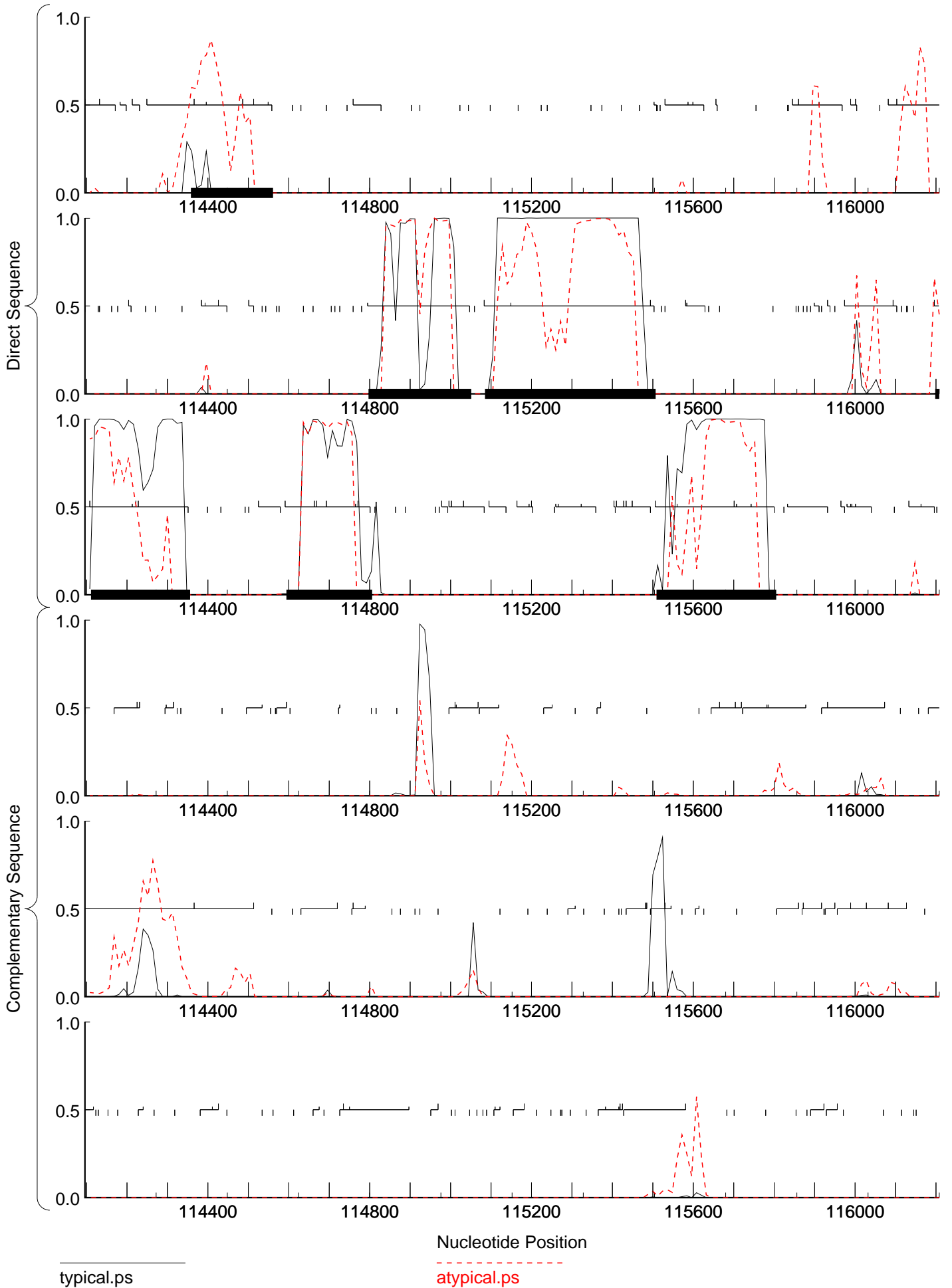
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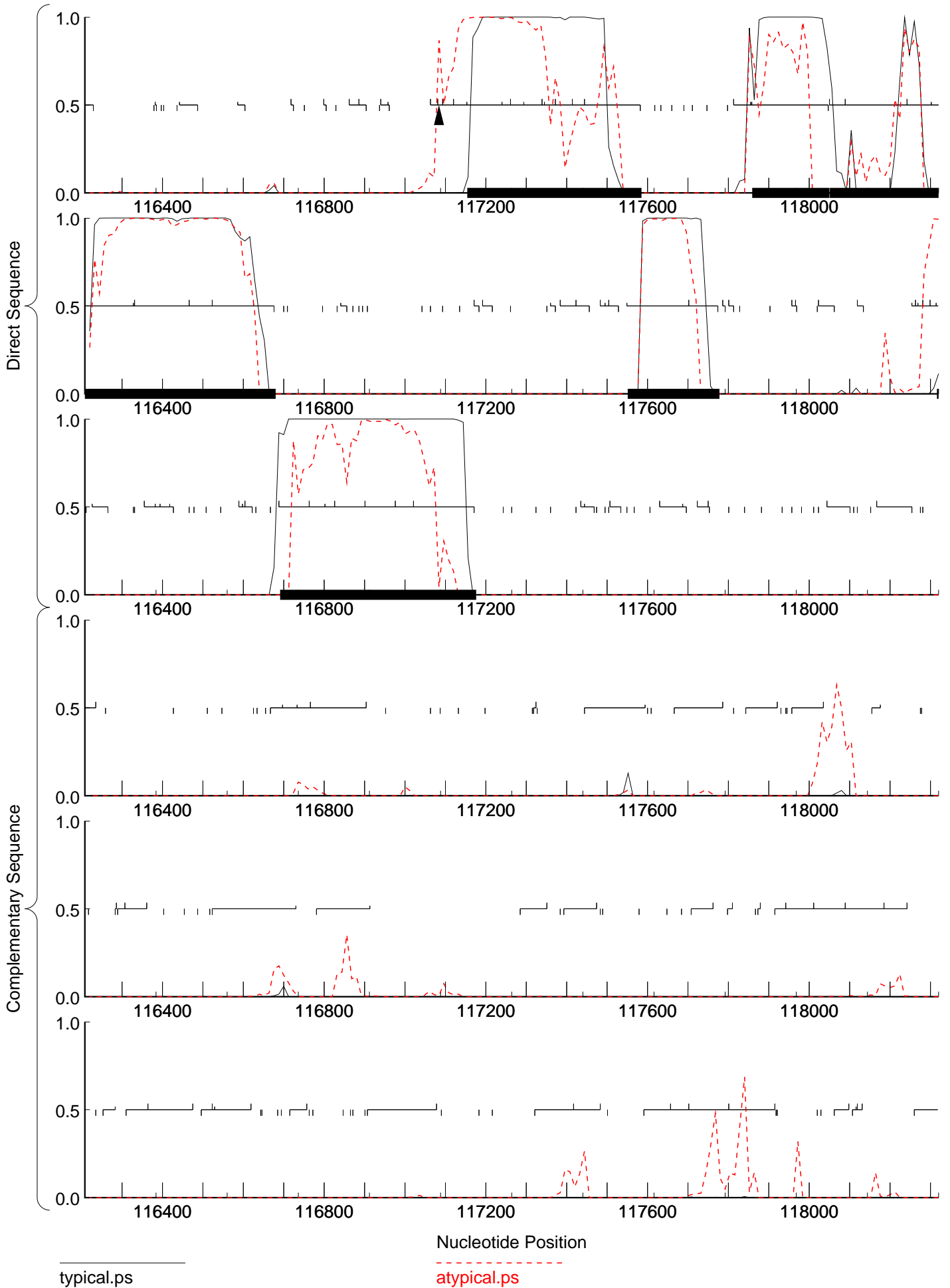
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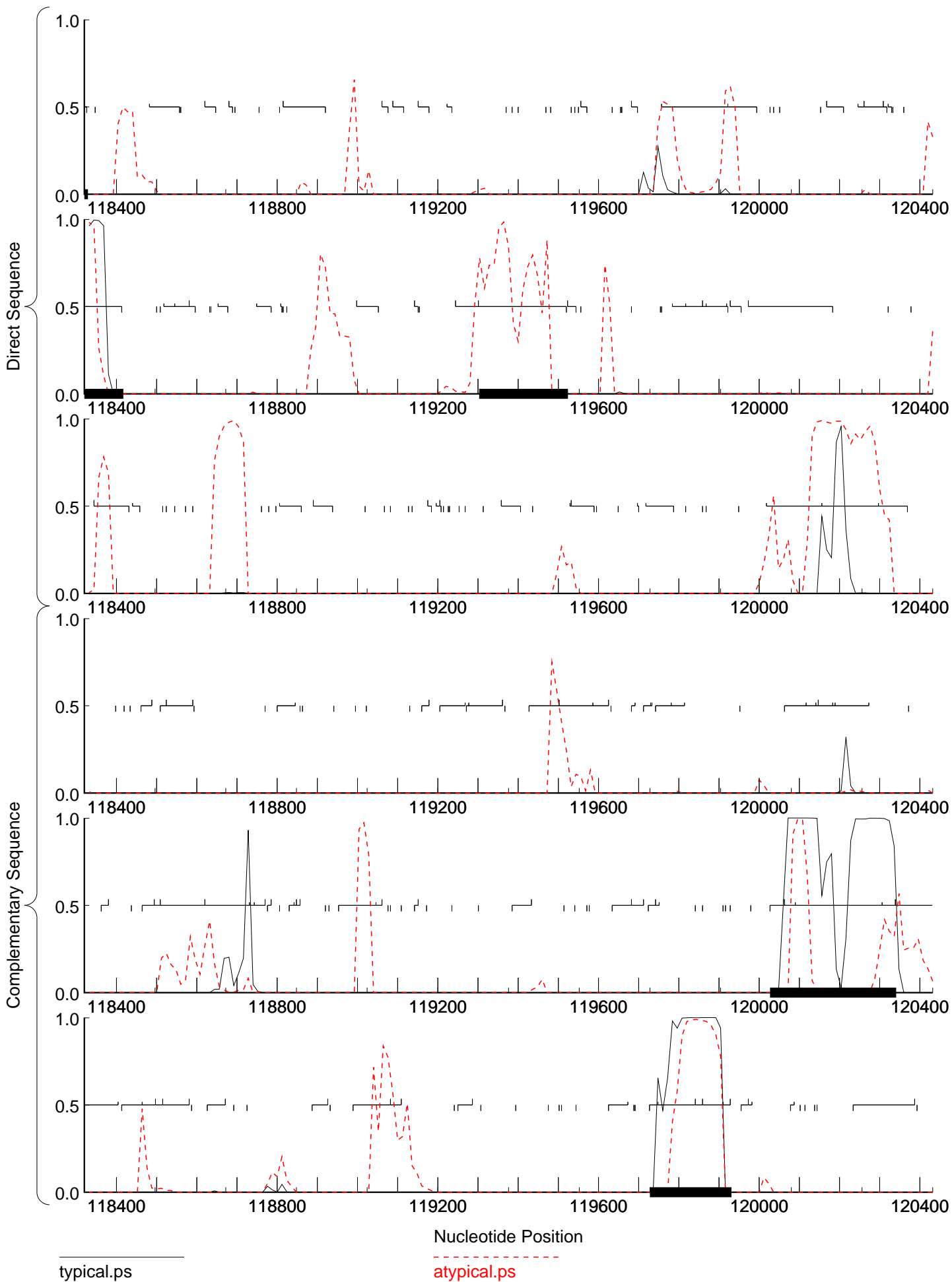
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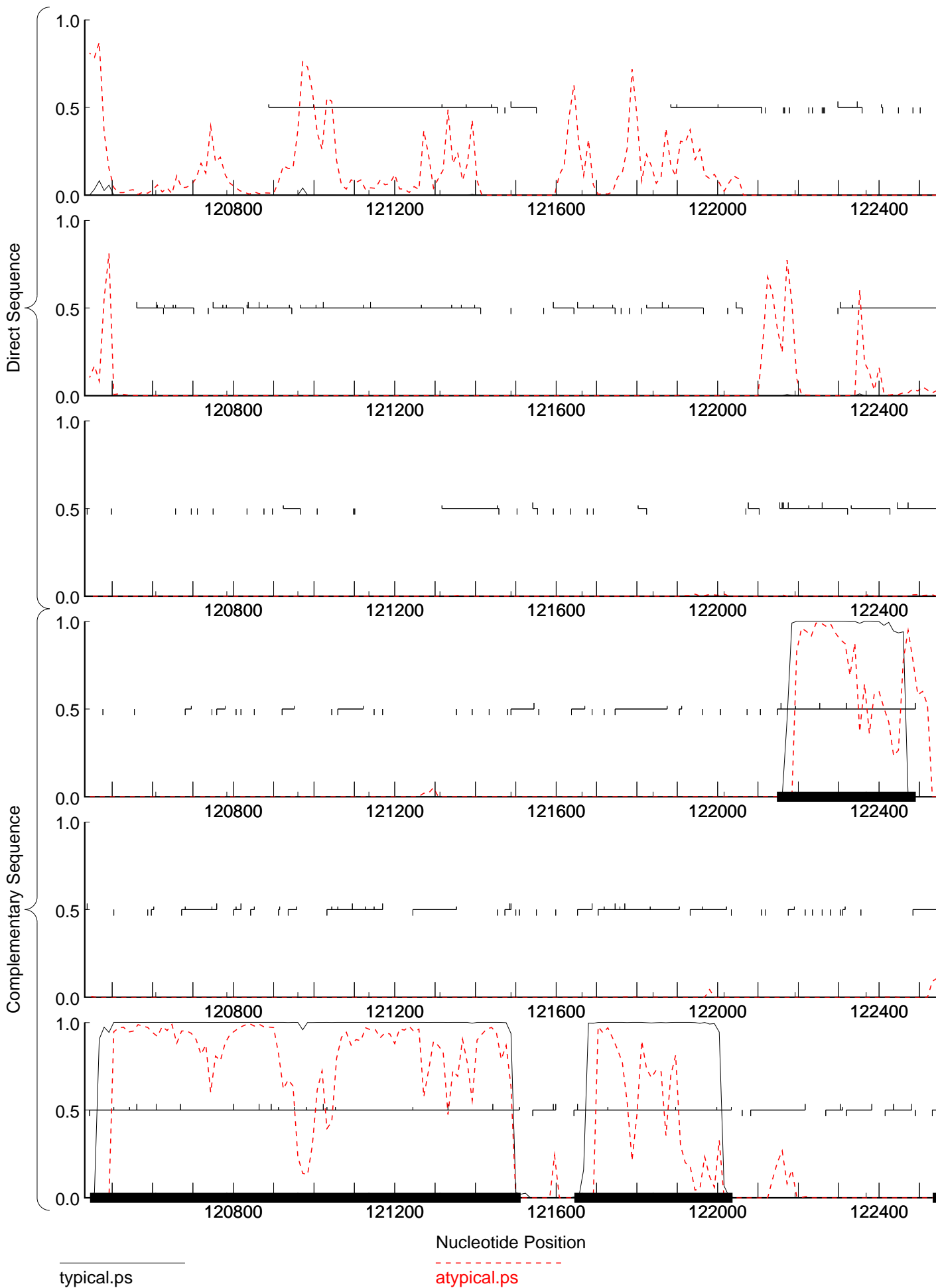






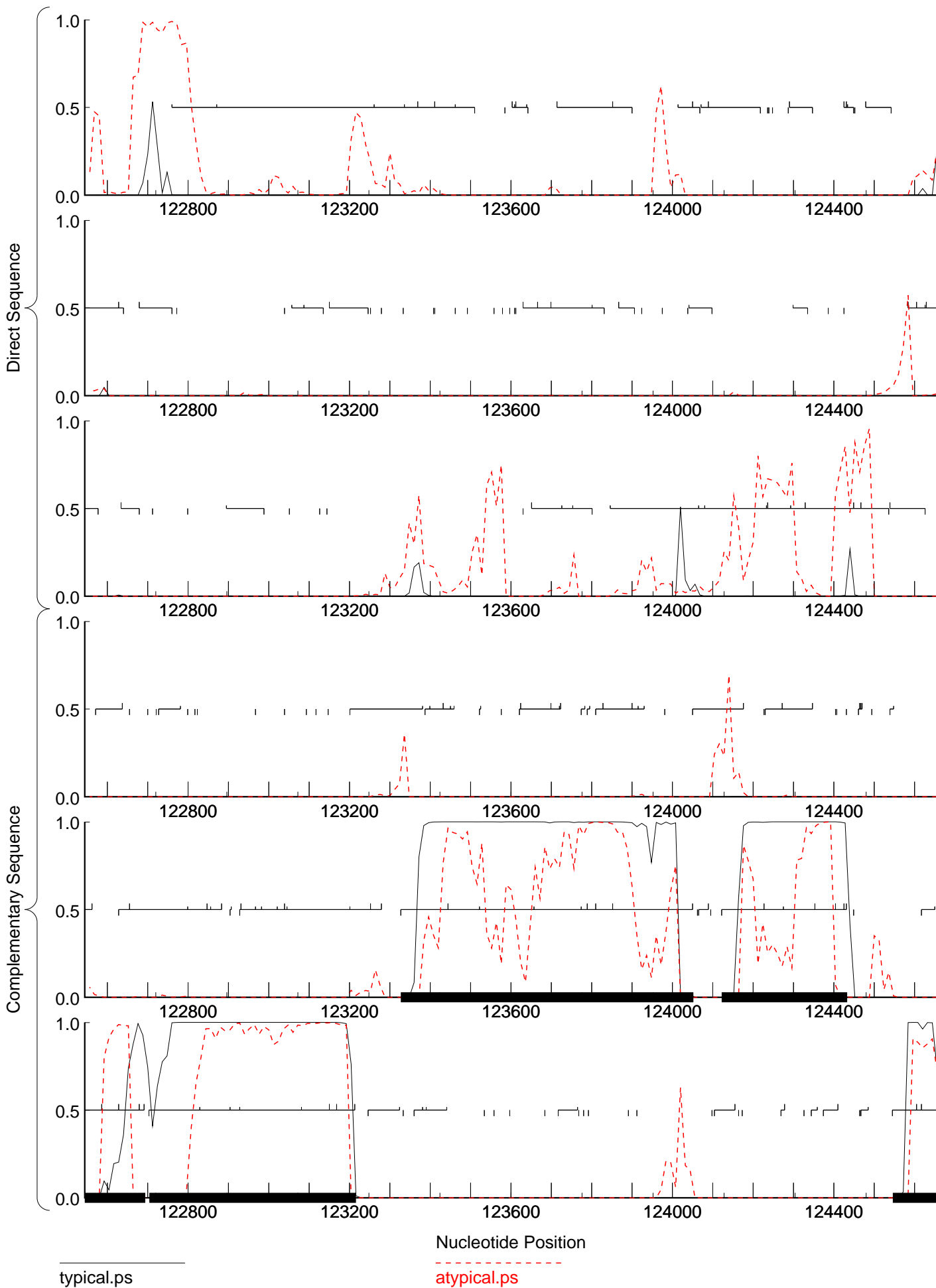


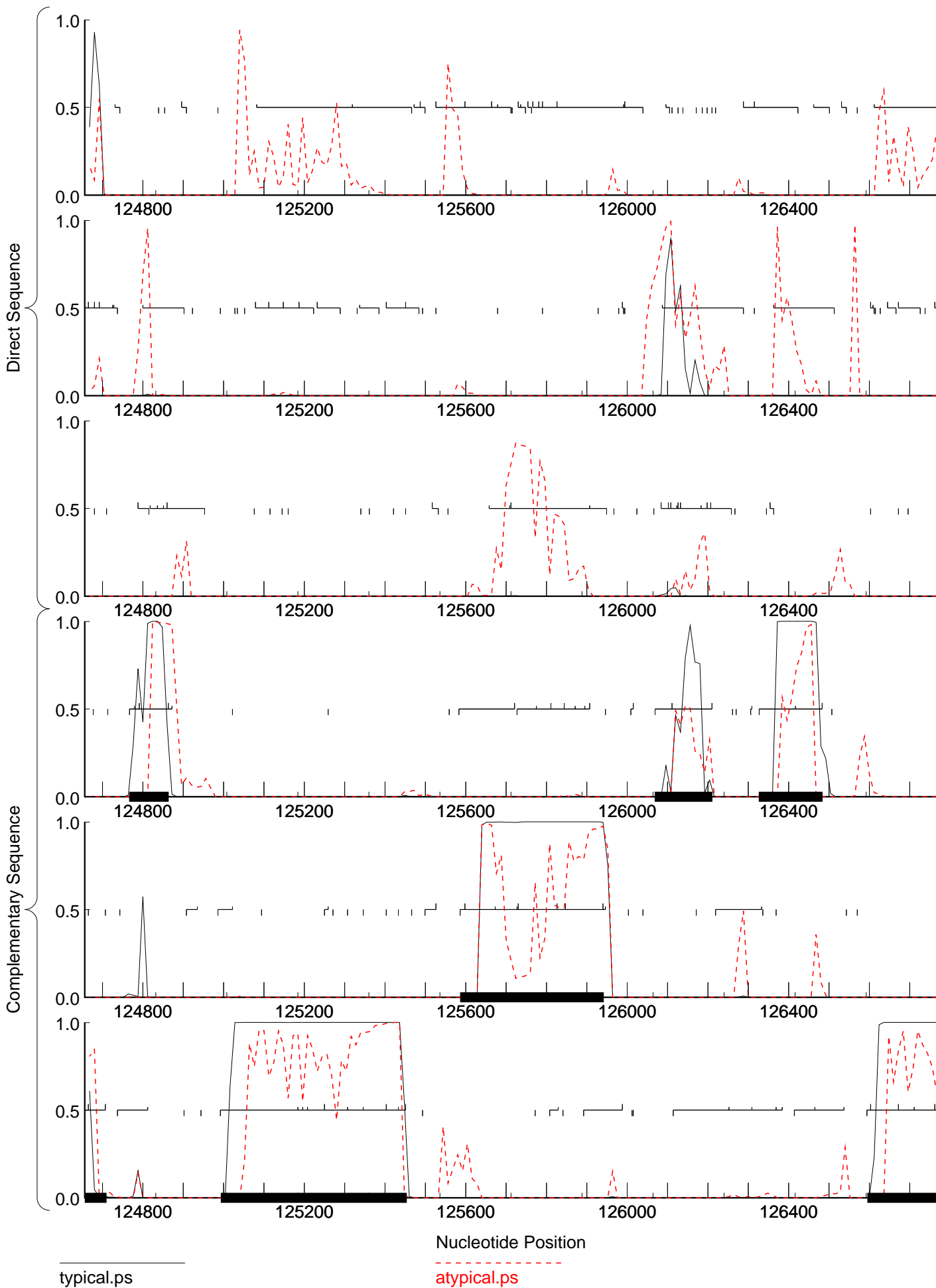


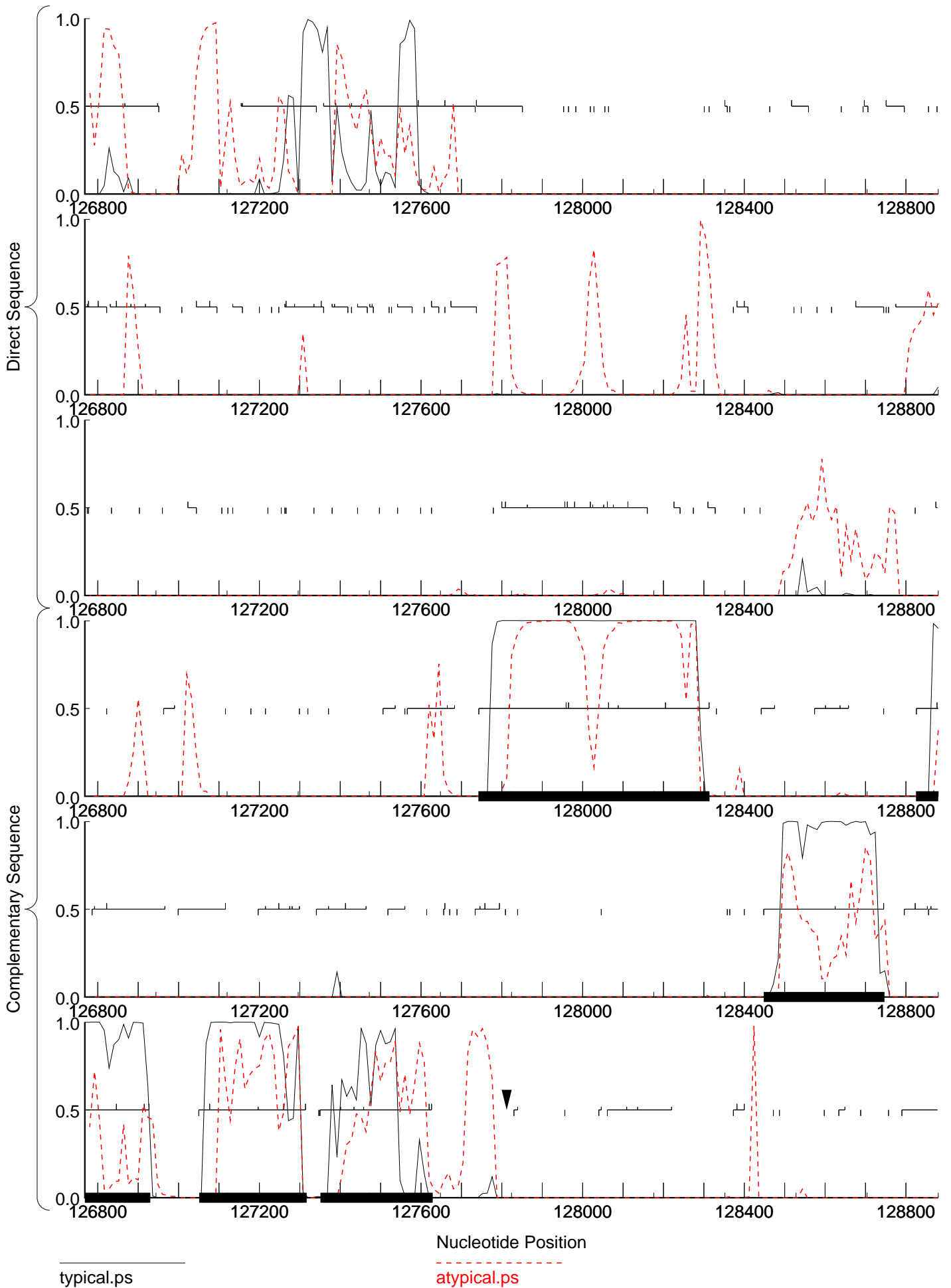


typical.ps

atypical.ps

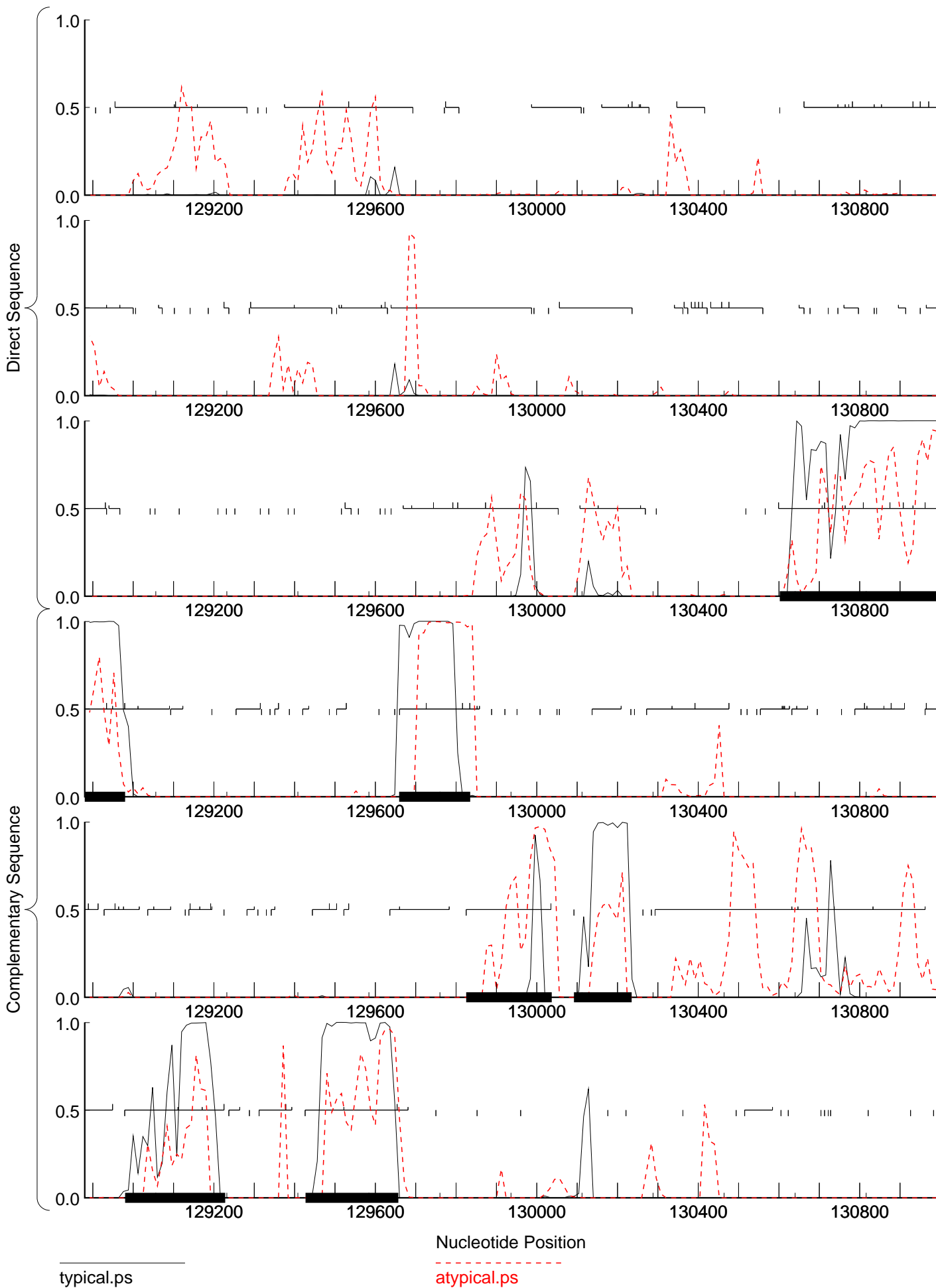






typical.ps

atypical.ps



typical.ps

atypical.ps

