

# GeneMark

Version 2.5p (09.08.06)

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## PROGRAM INFORMATION

Sequence : Arthrobacter phage Racecar complete sequence, 173709 bp, circularly permuted, Cluster FC  
Analysis Date : 1/18/18 at 20:35:13  
Pages : 84  
Sequence Length : 173709 bp  
GC Content : 53.07%

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,

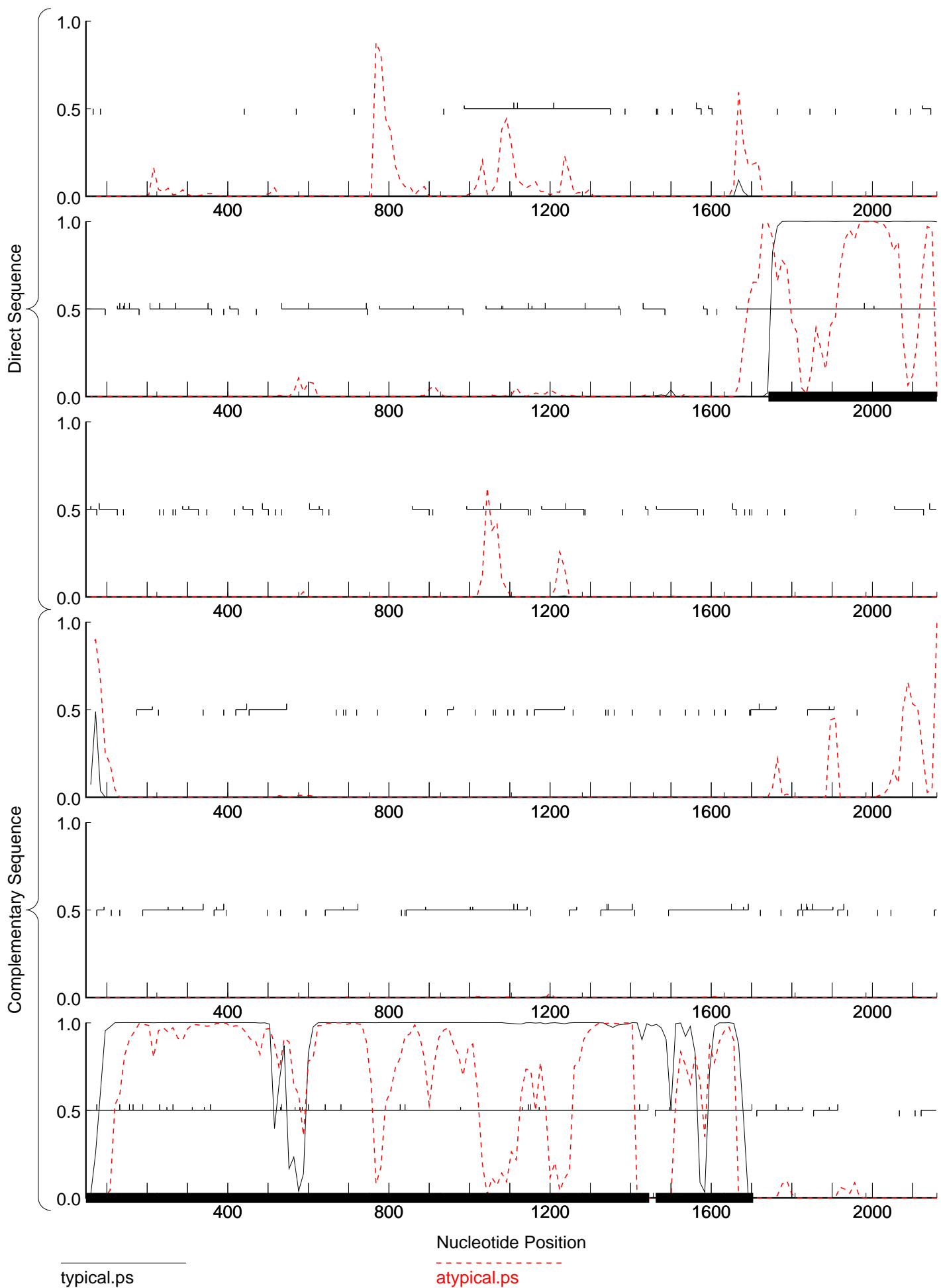
## MATRIX INFORMATION

Matrix : -  
Author : -  
Order : 4

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

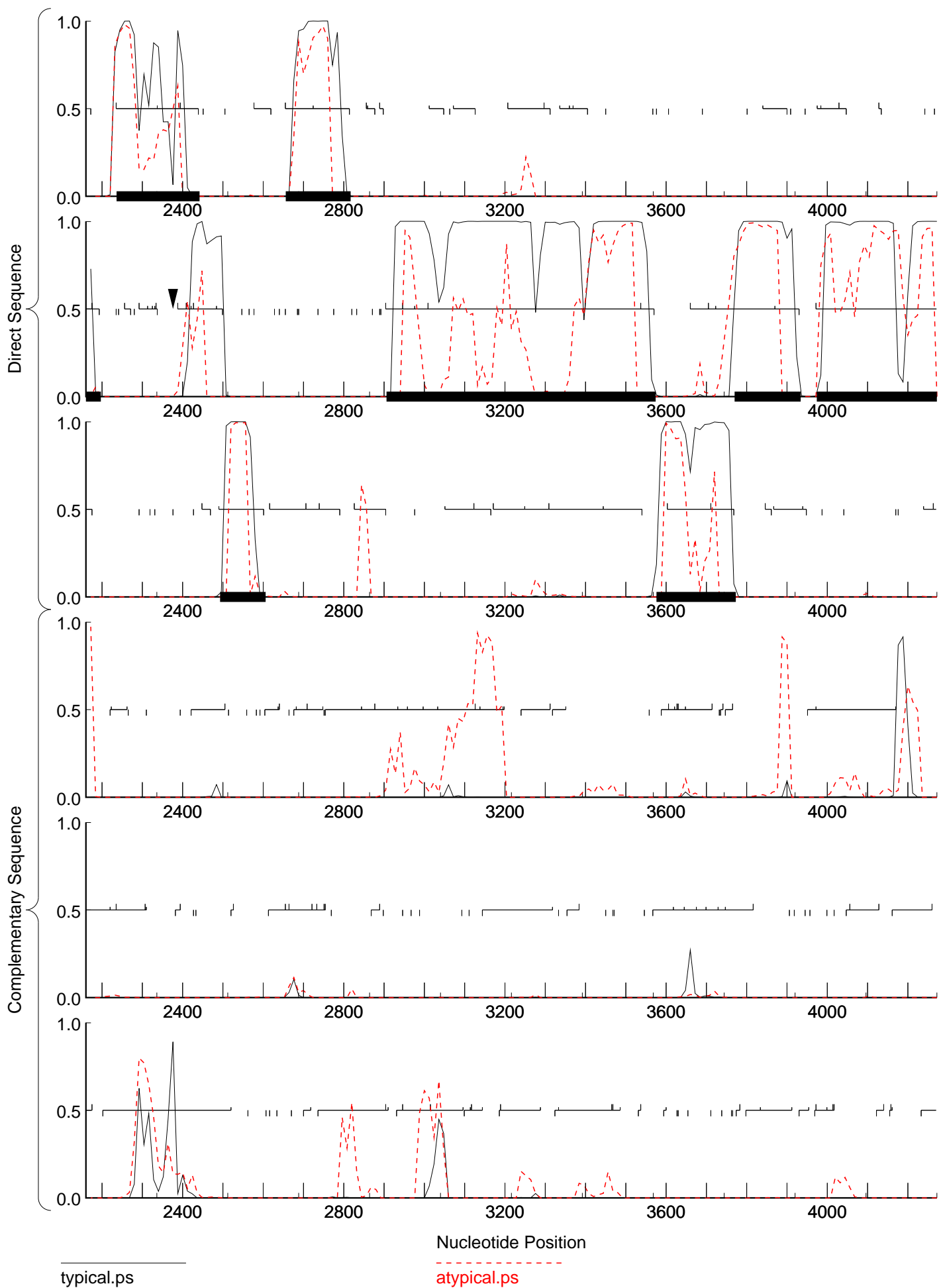
### Matrix notes & comments

Training set derived by GeneMarkS, 4.30 October 2014  
Thu Jan 18 20:35:13 2018



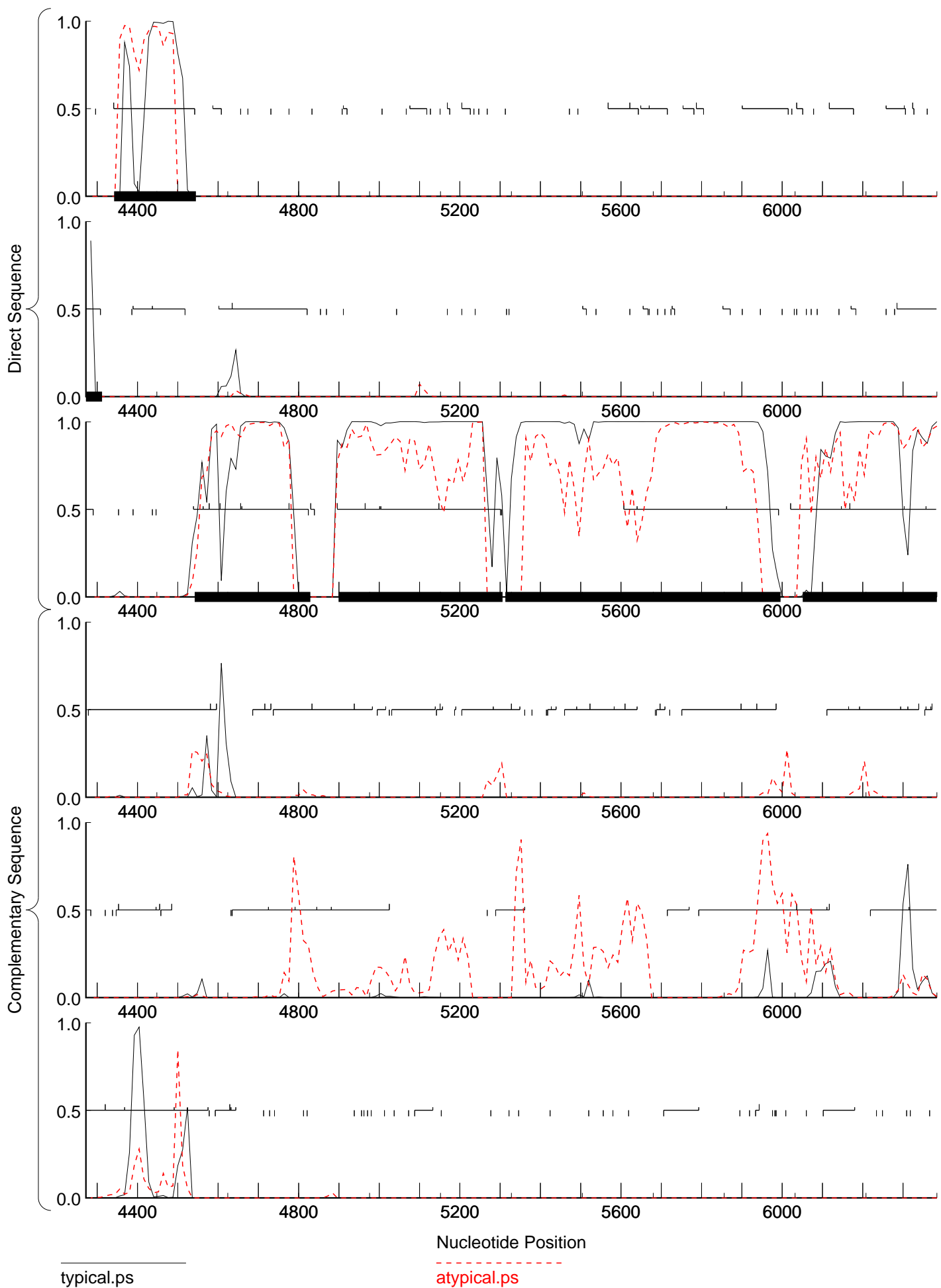
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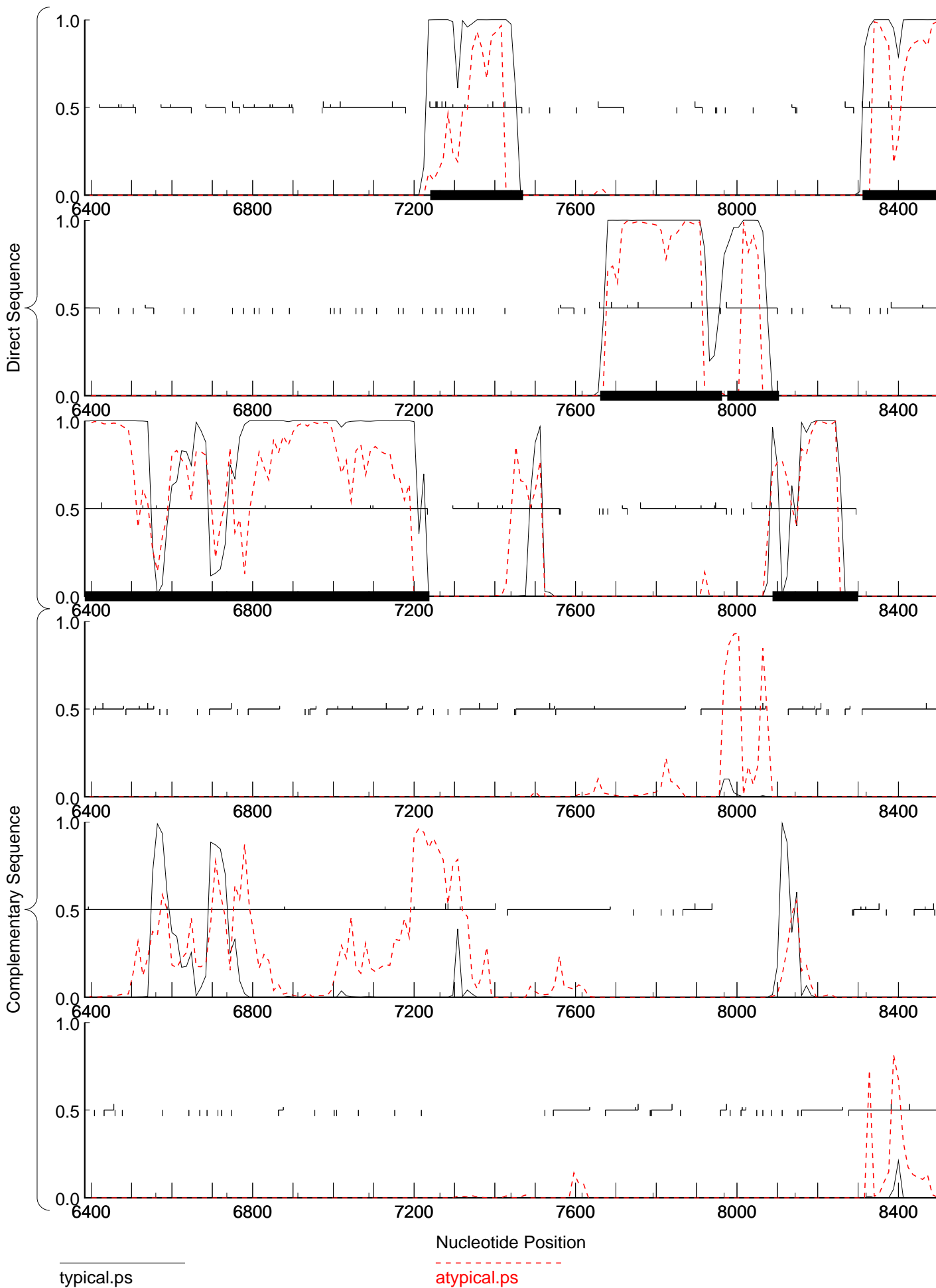
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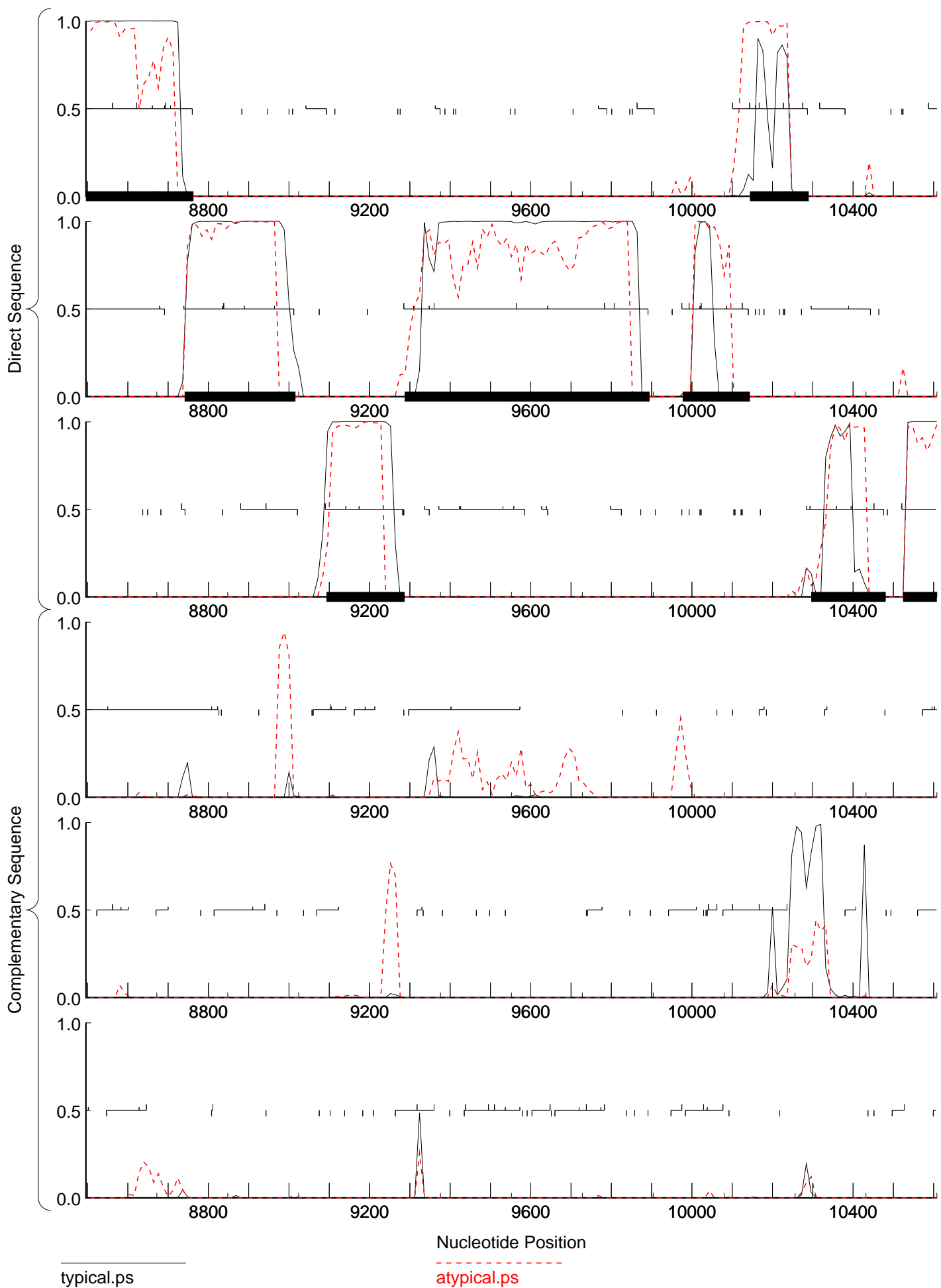


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atypical.ps

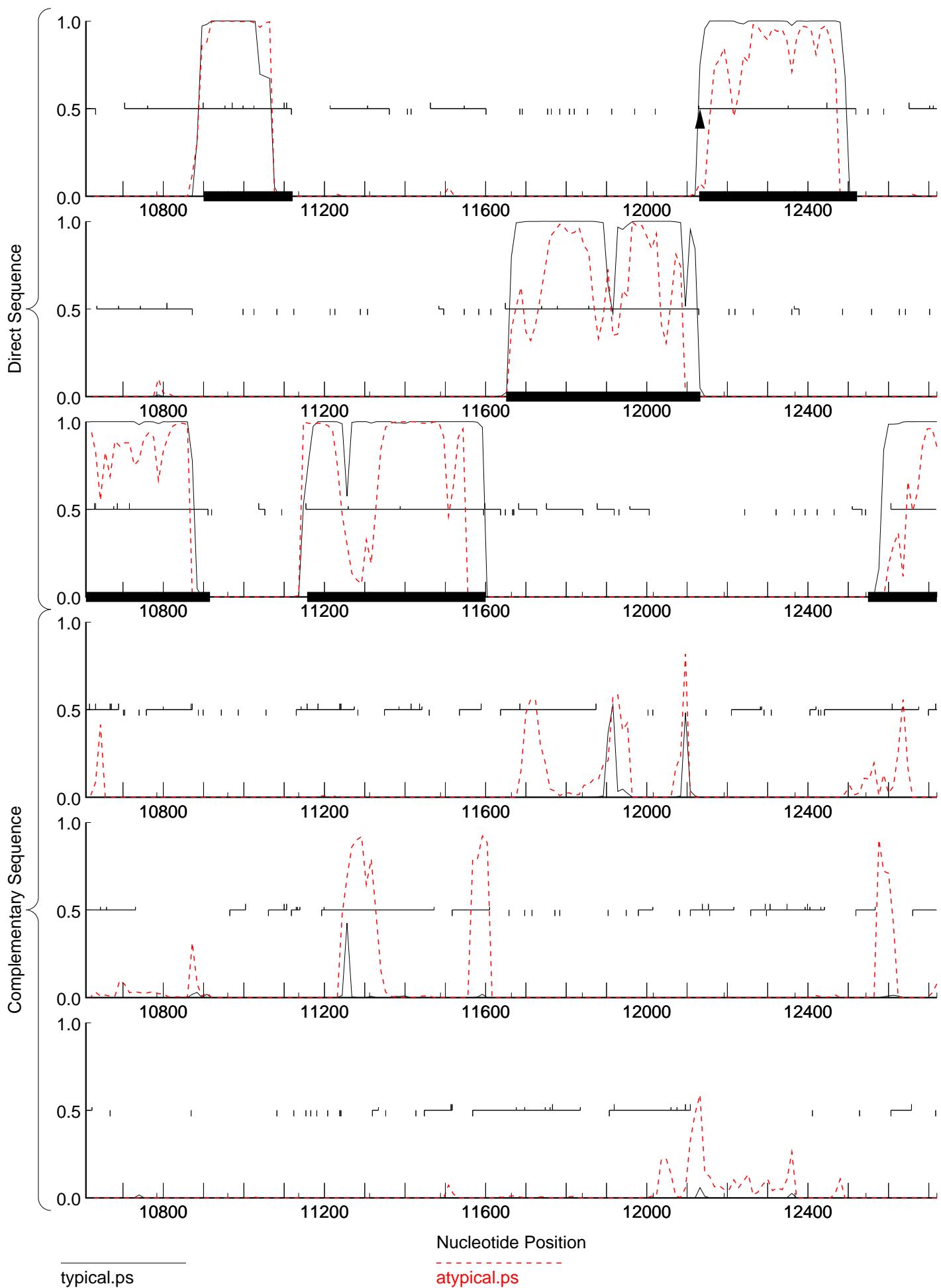


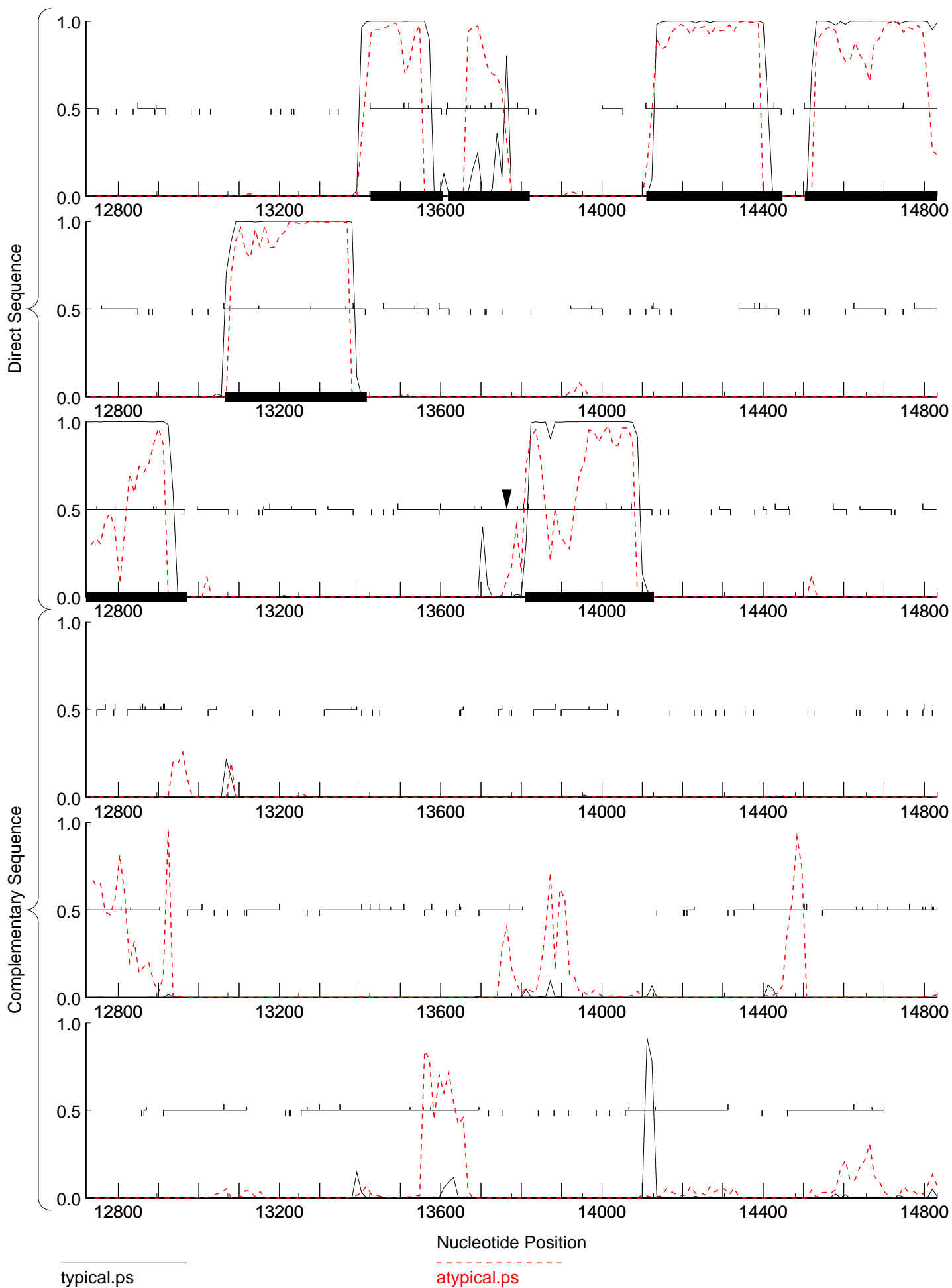




typical.ps

atypical.ps

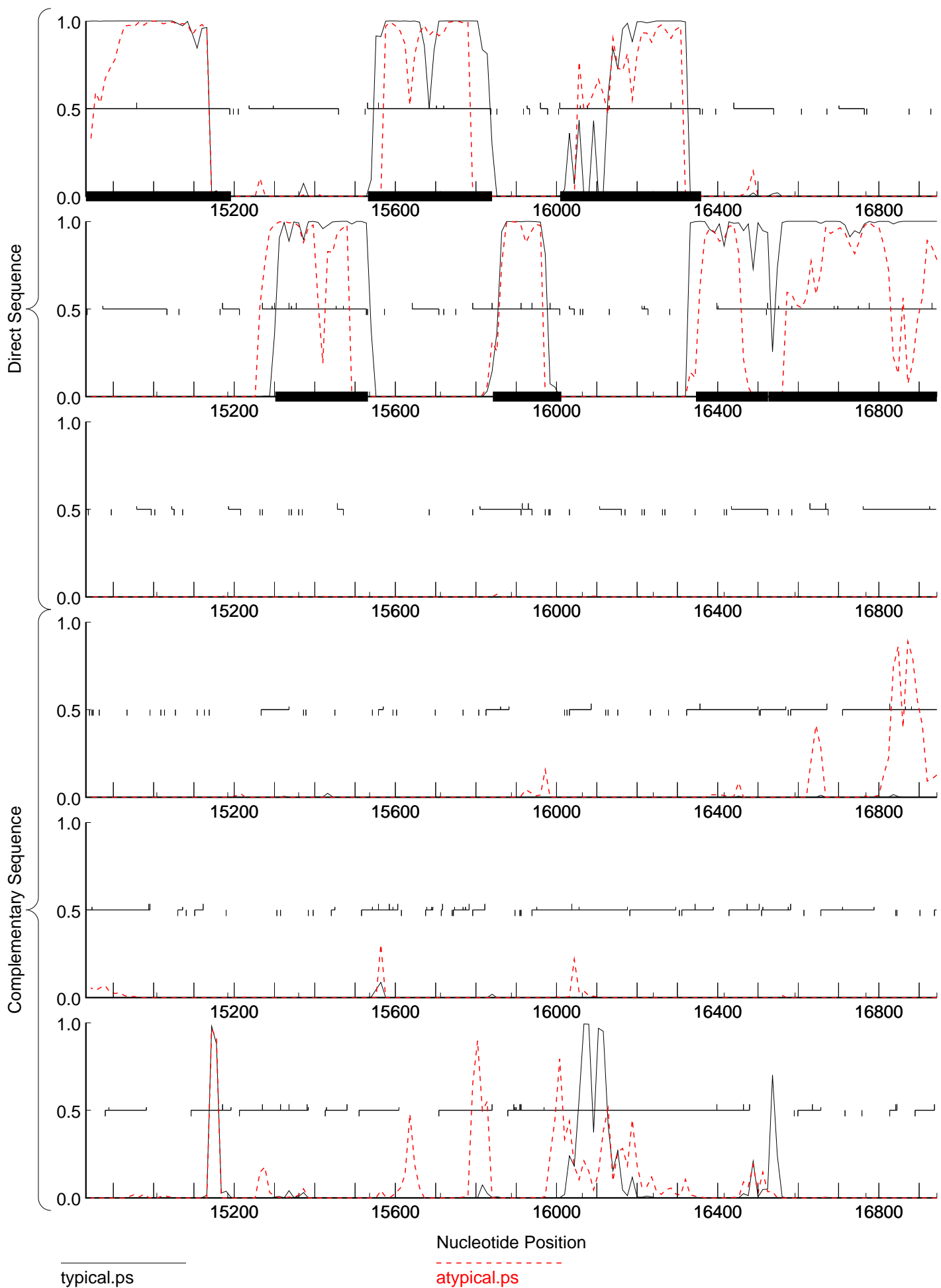




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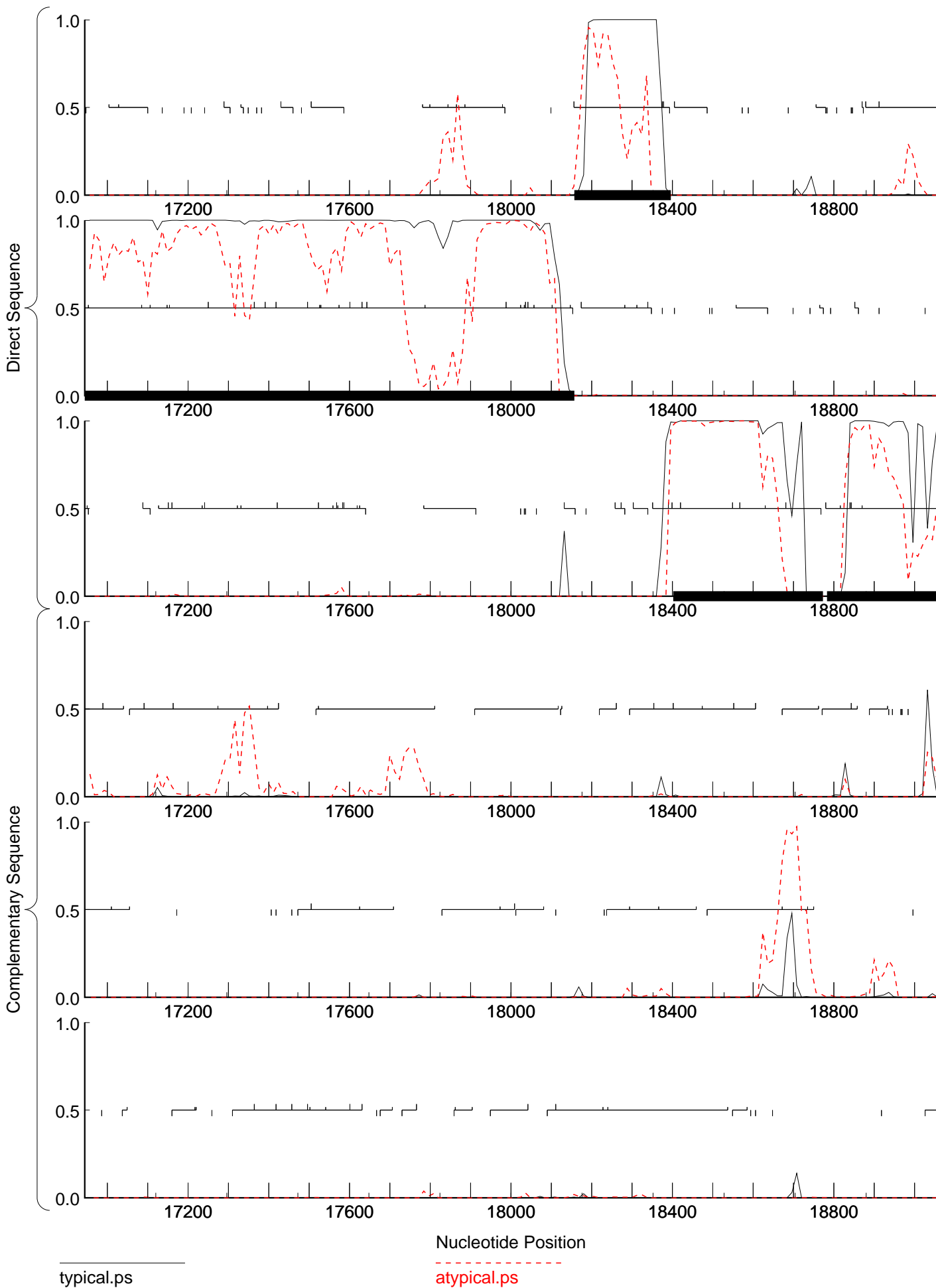
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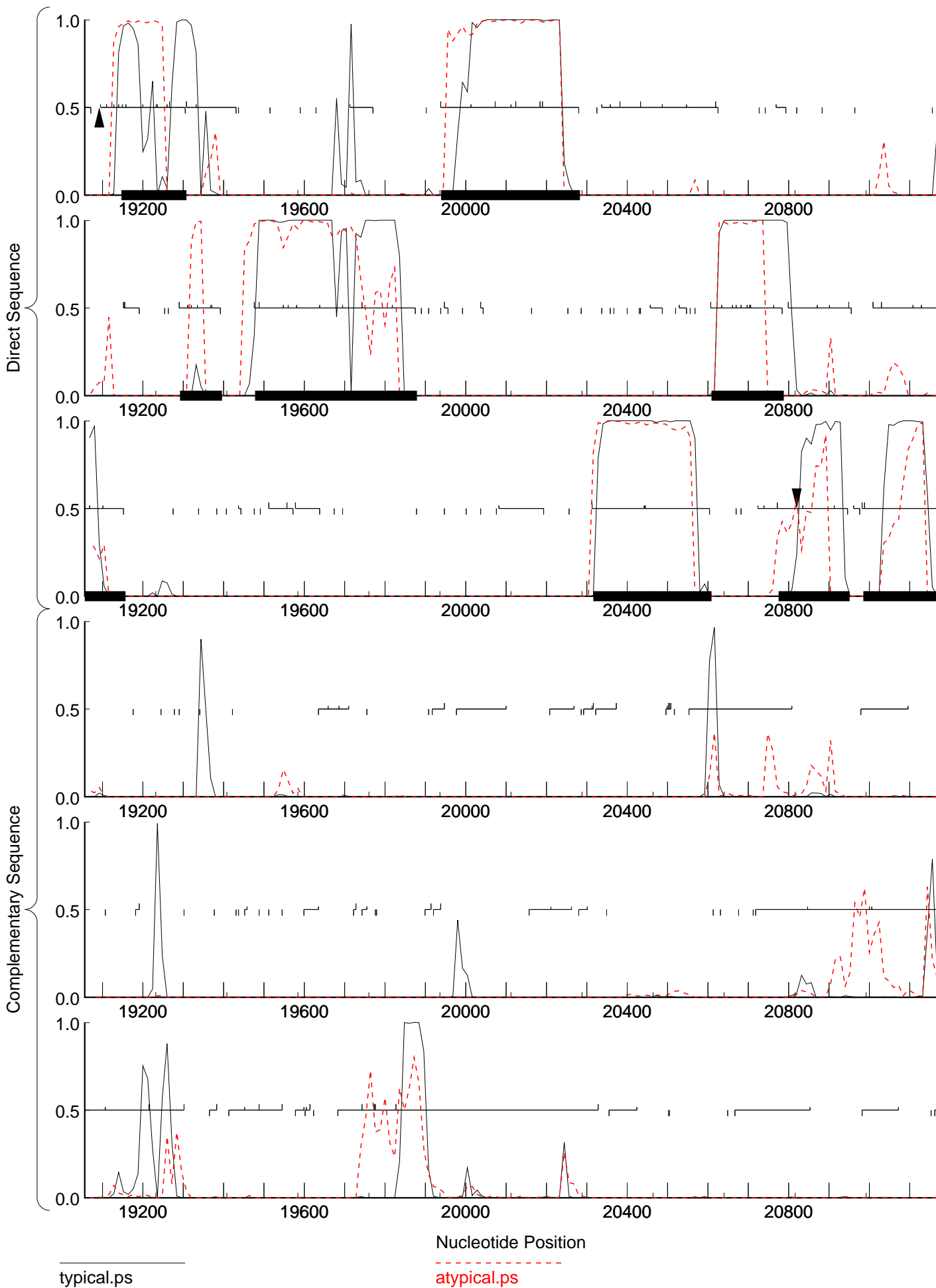


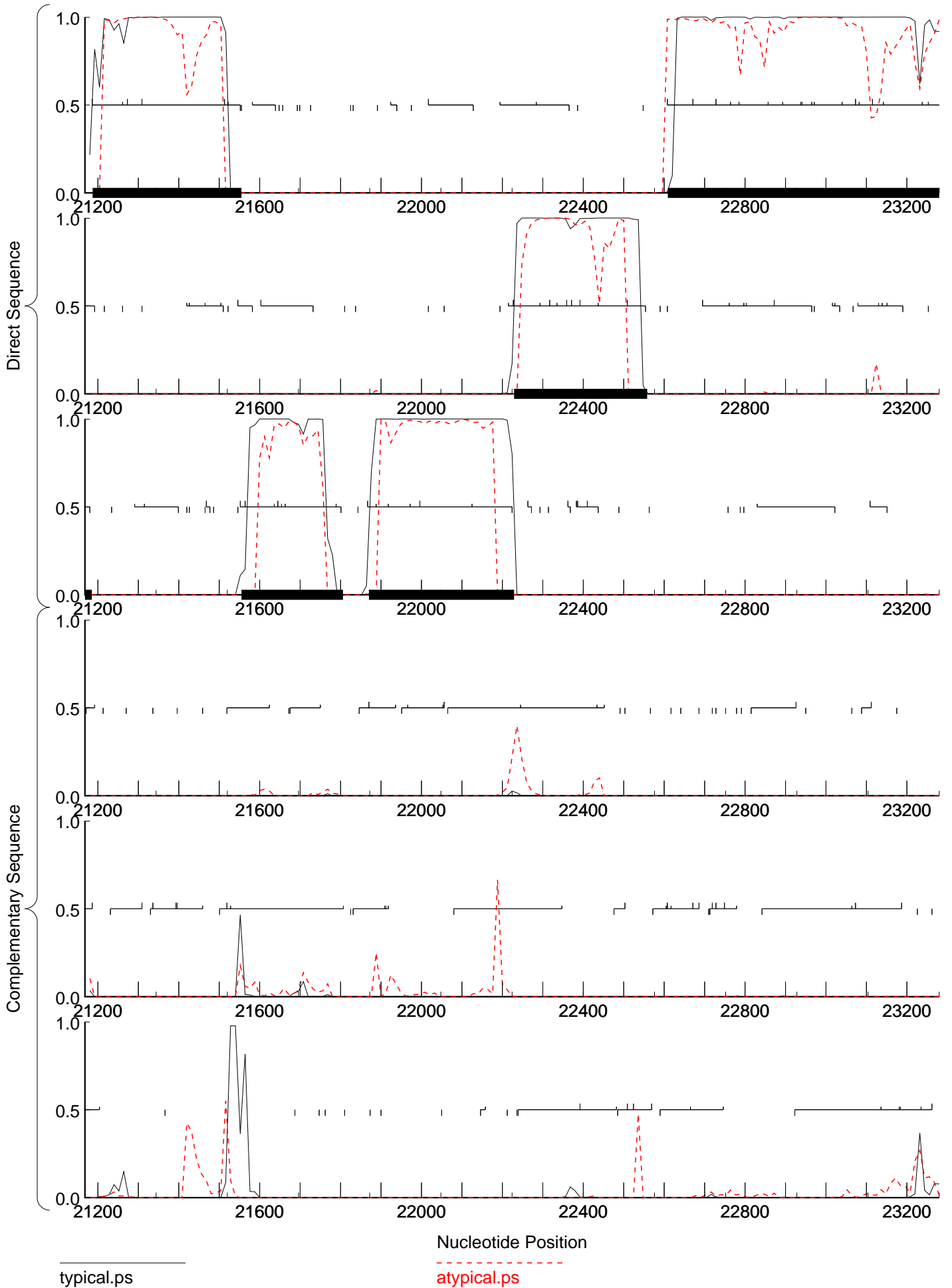


typical.ps

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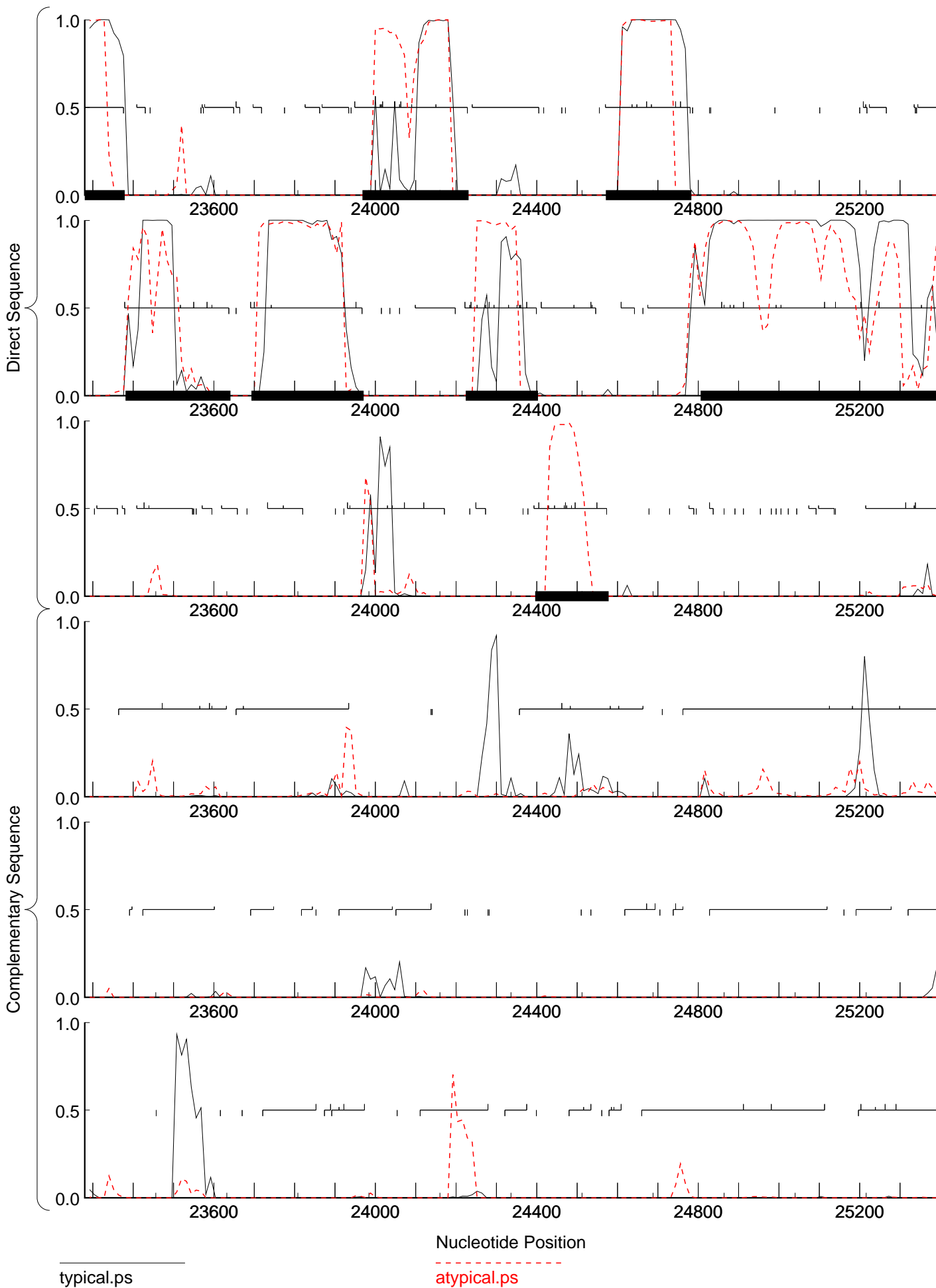


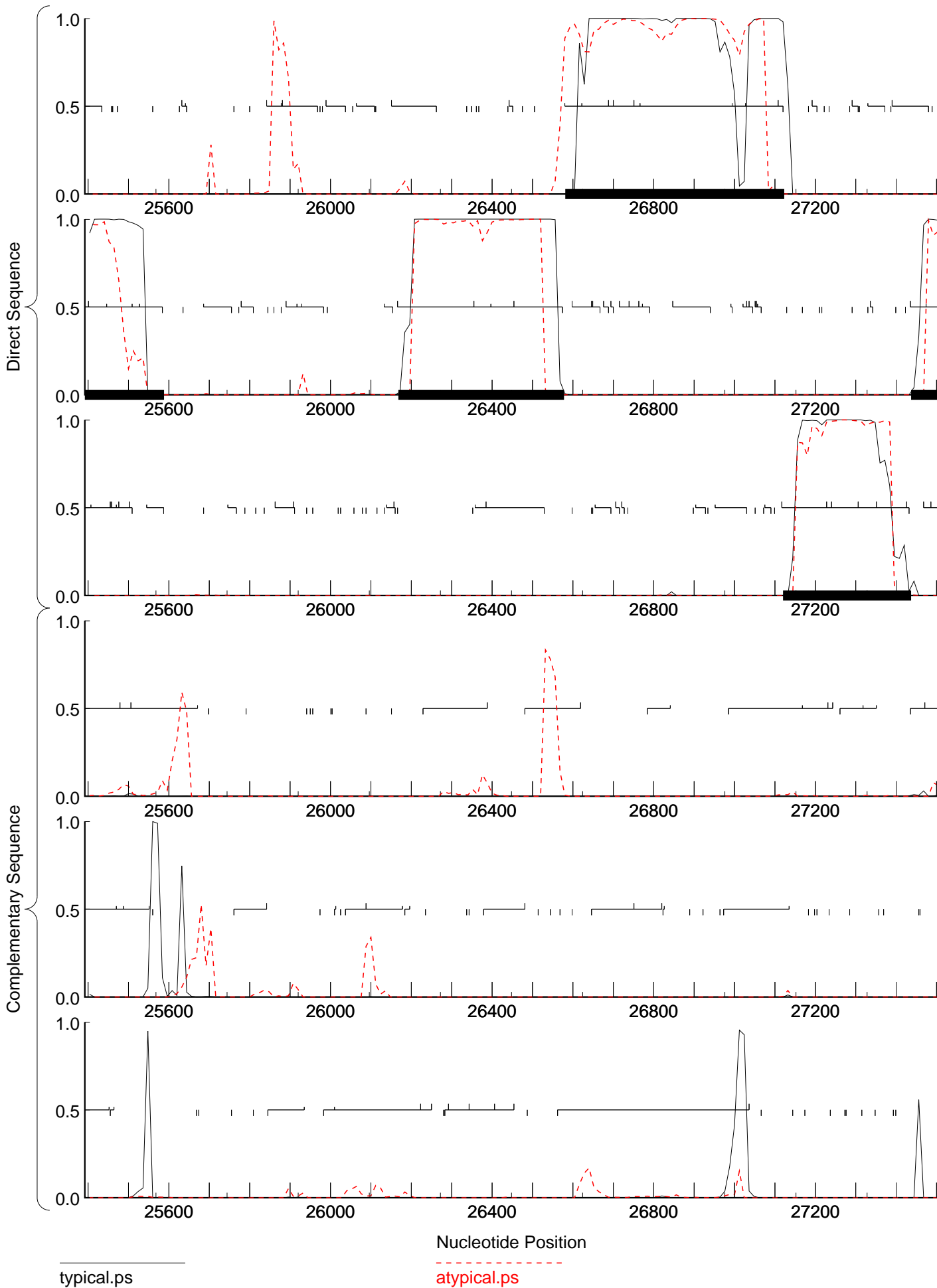




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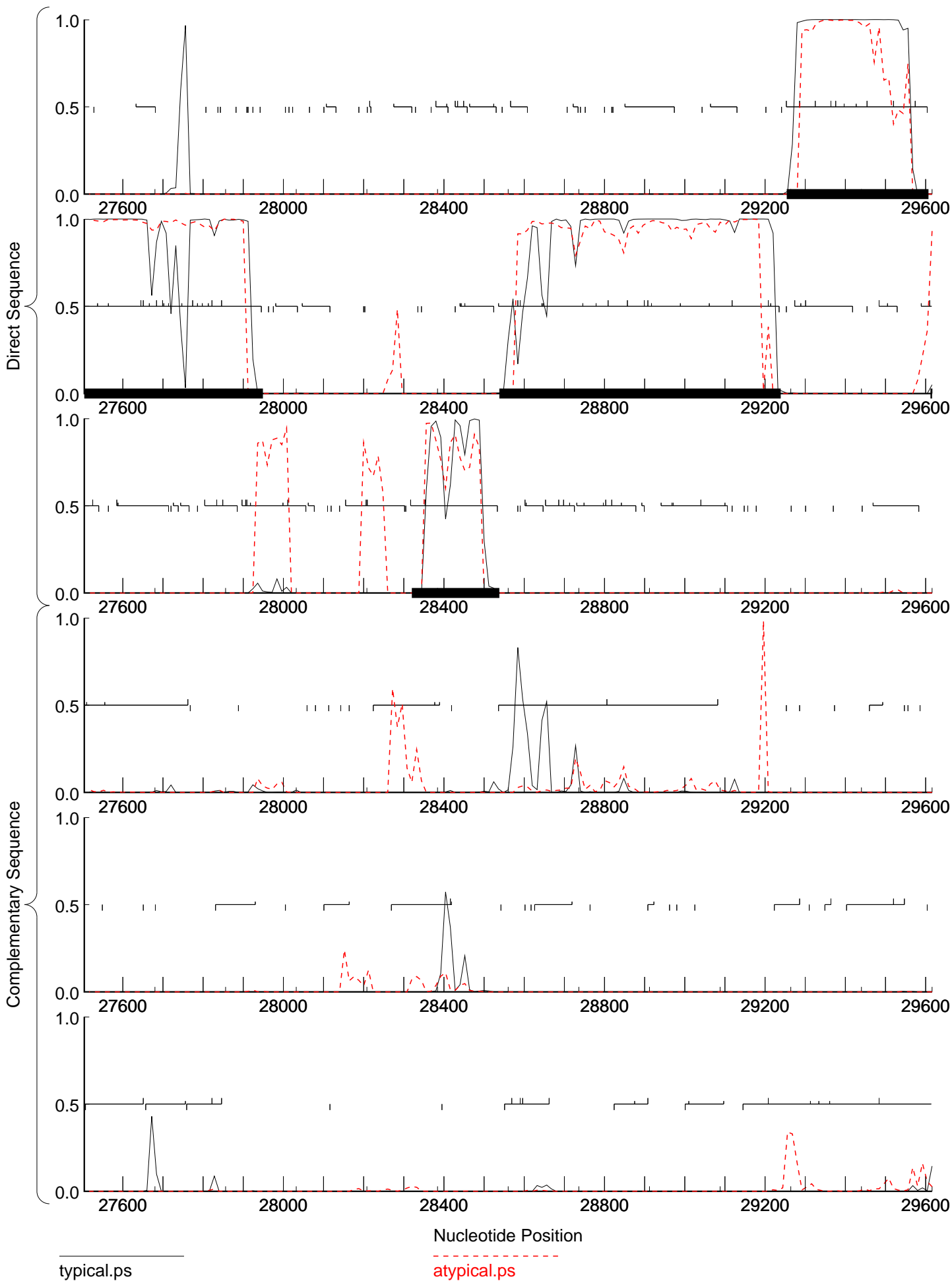
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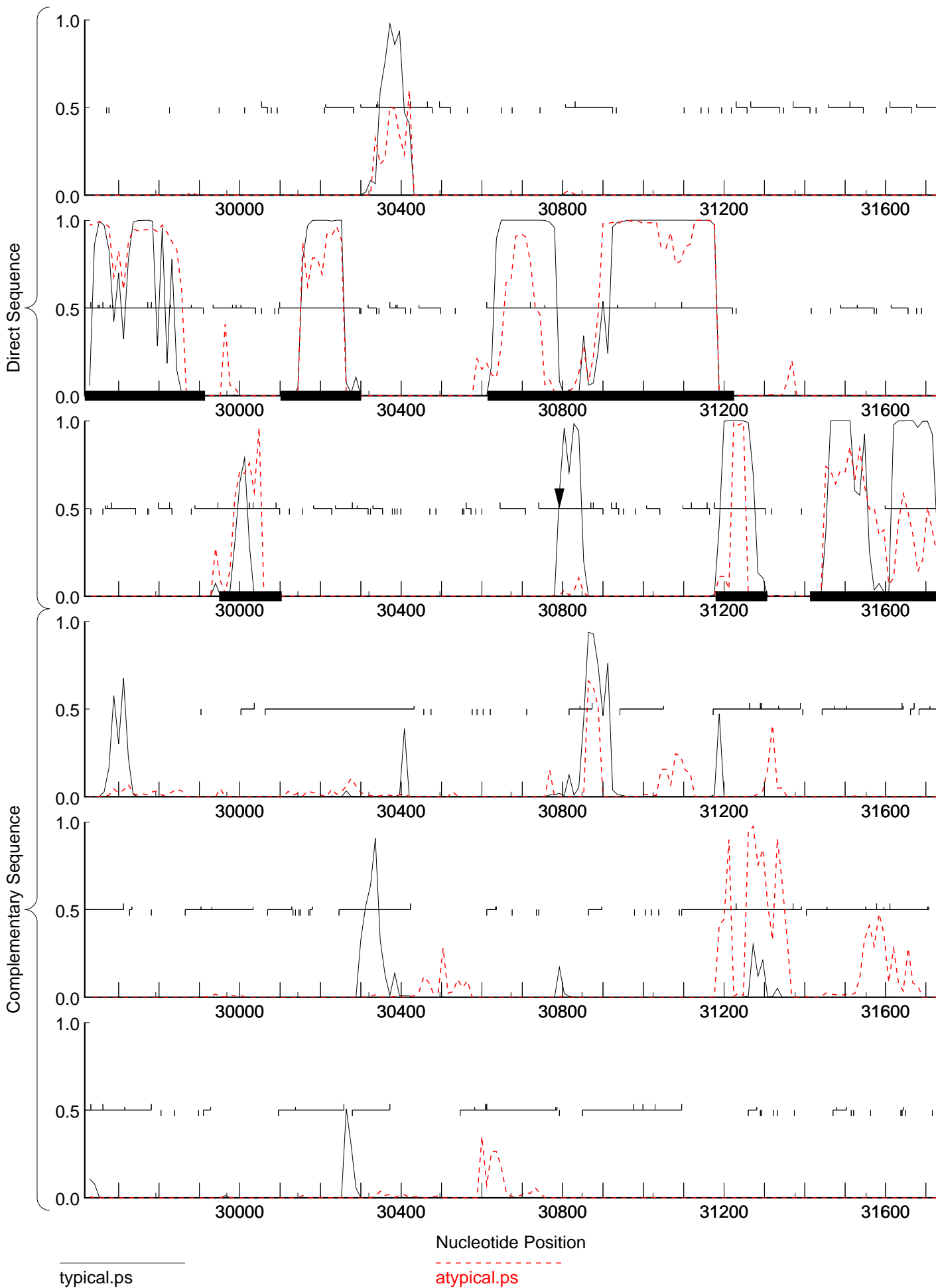
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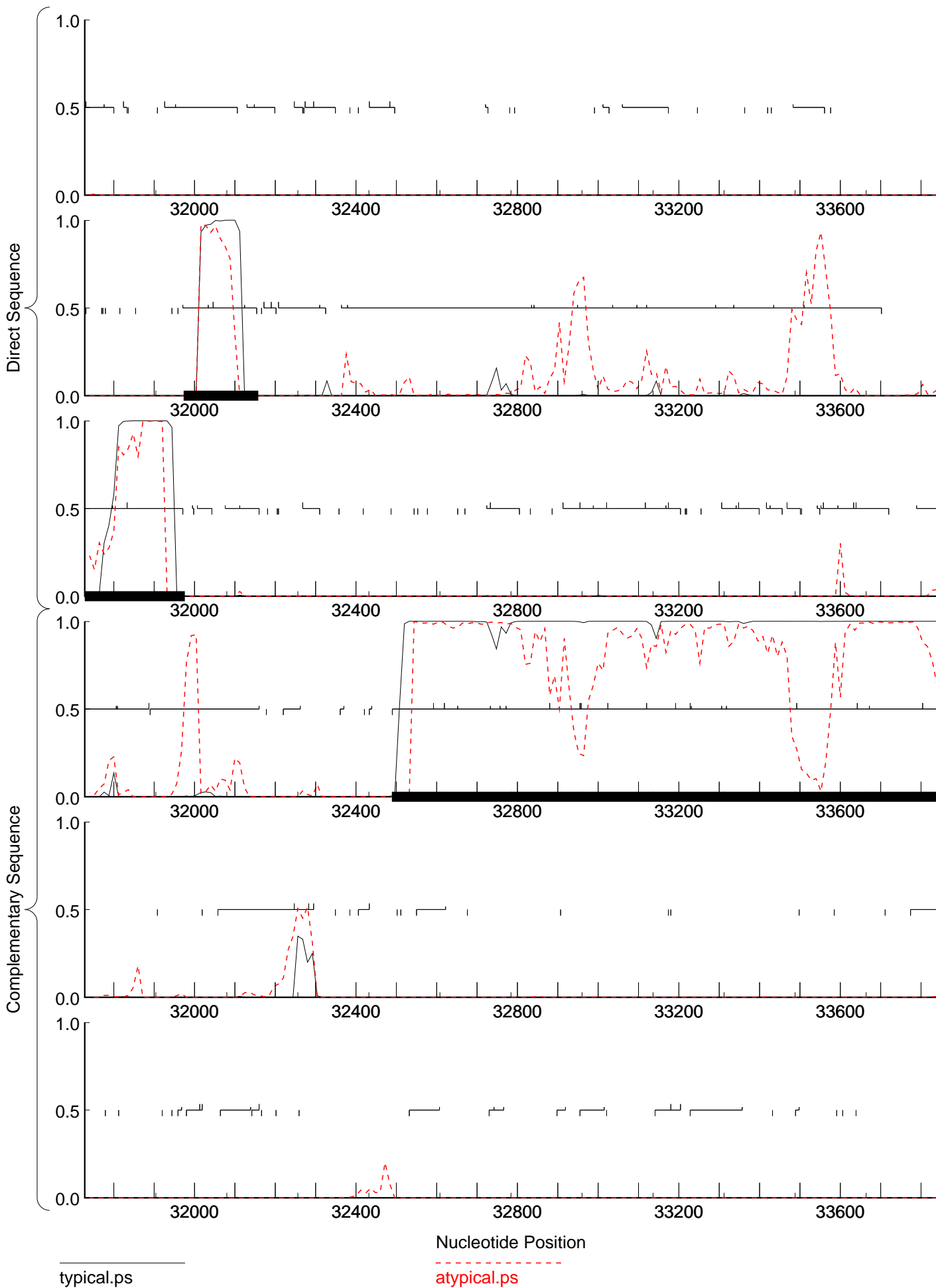


typical.ps

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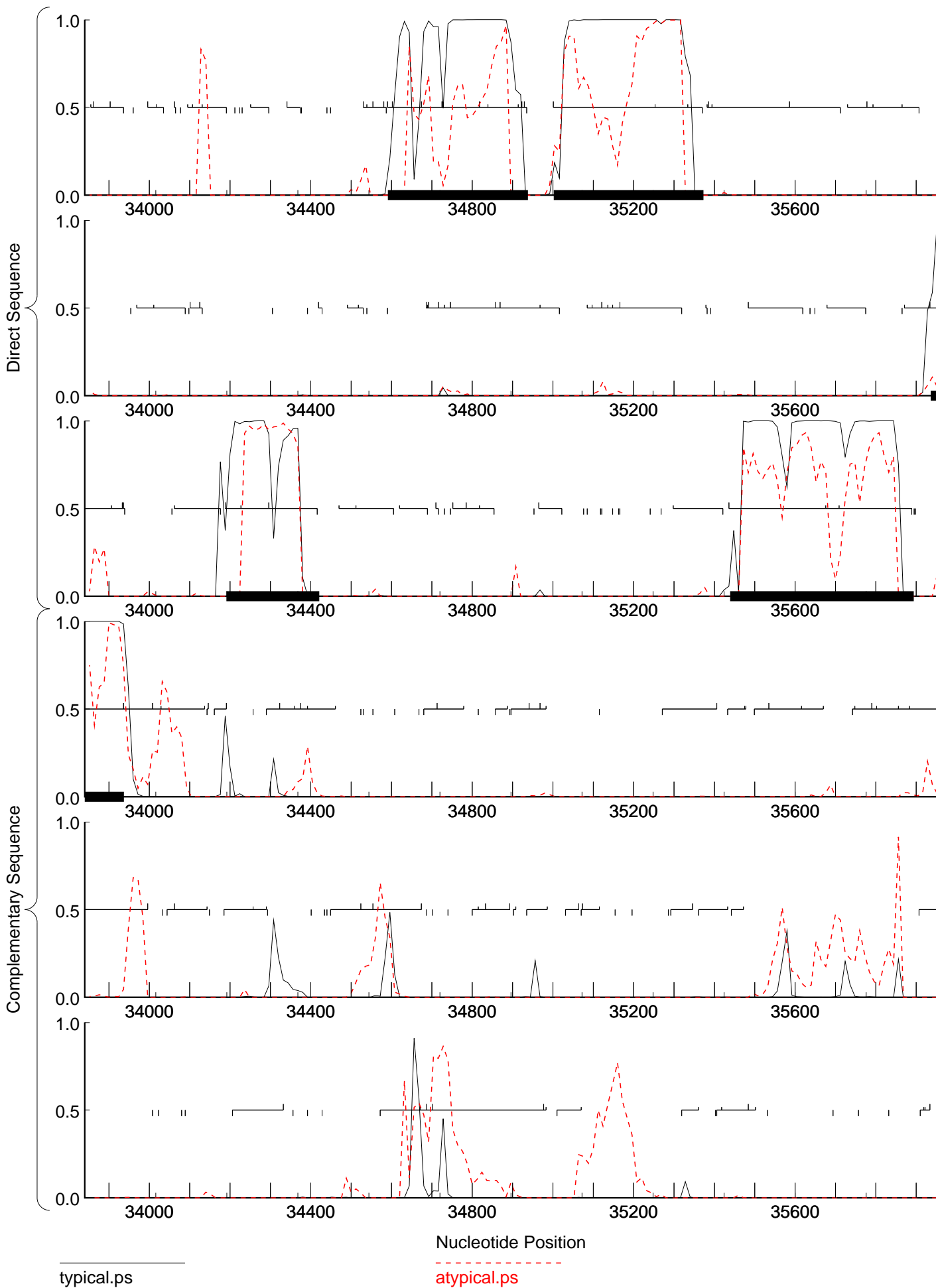






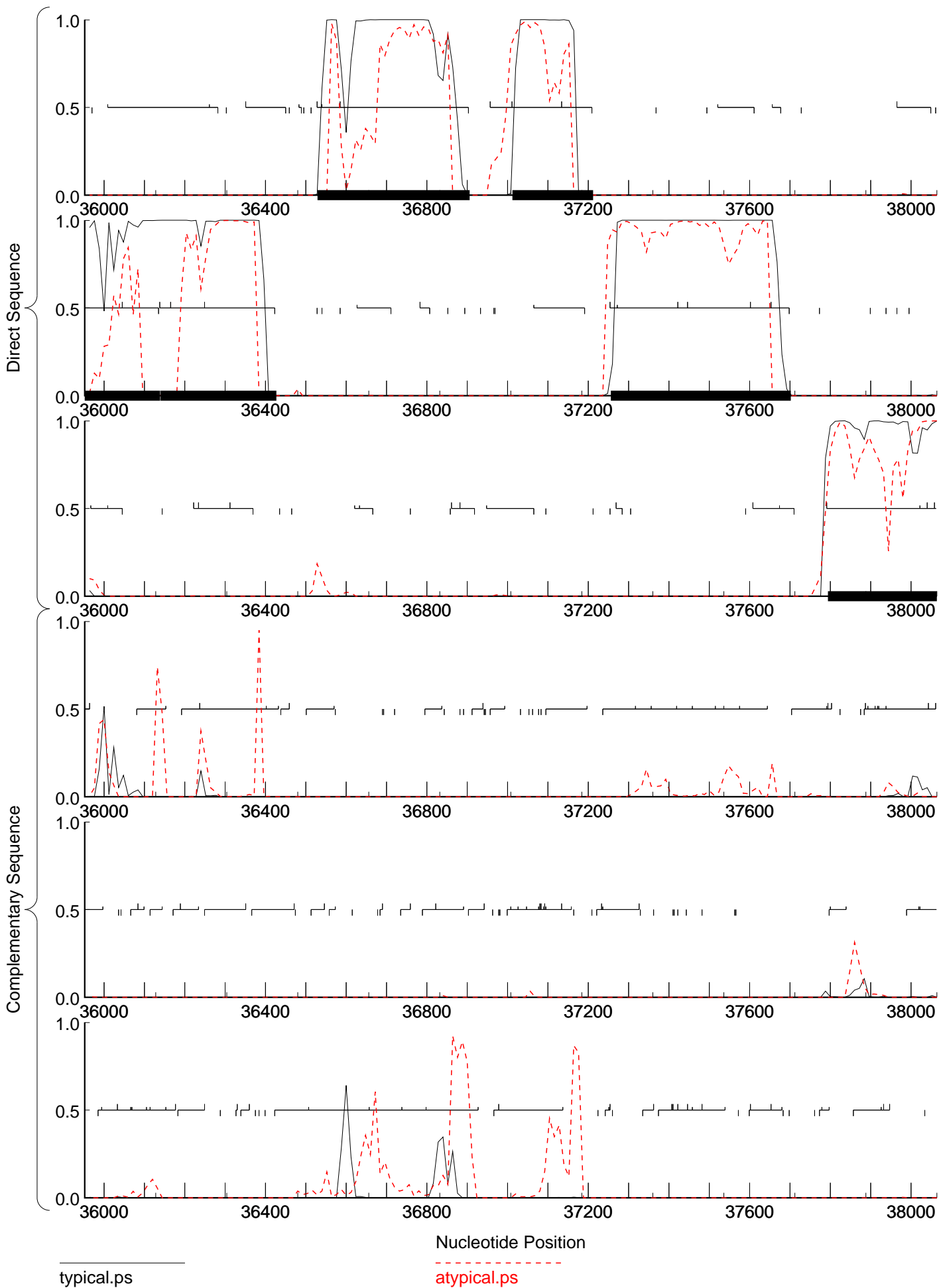
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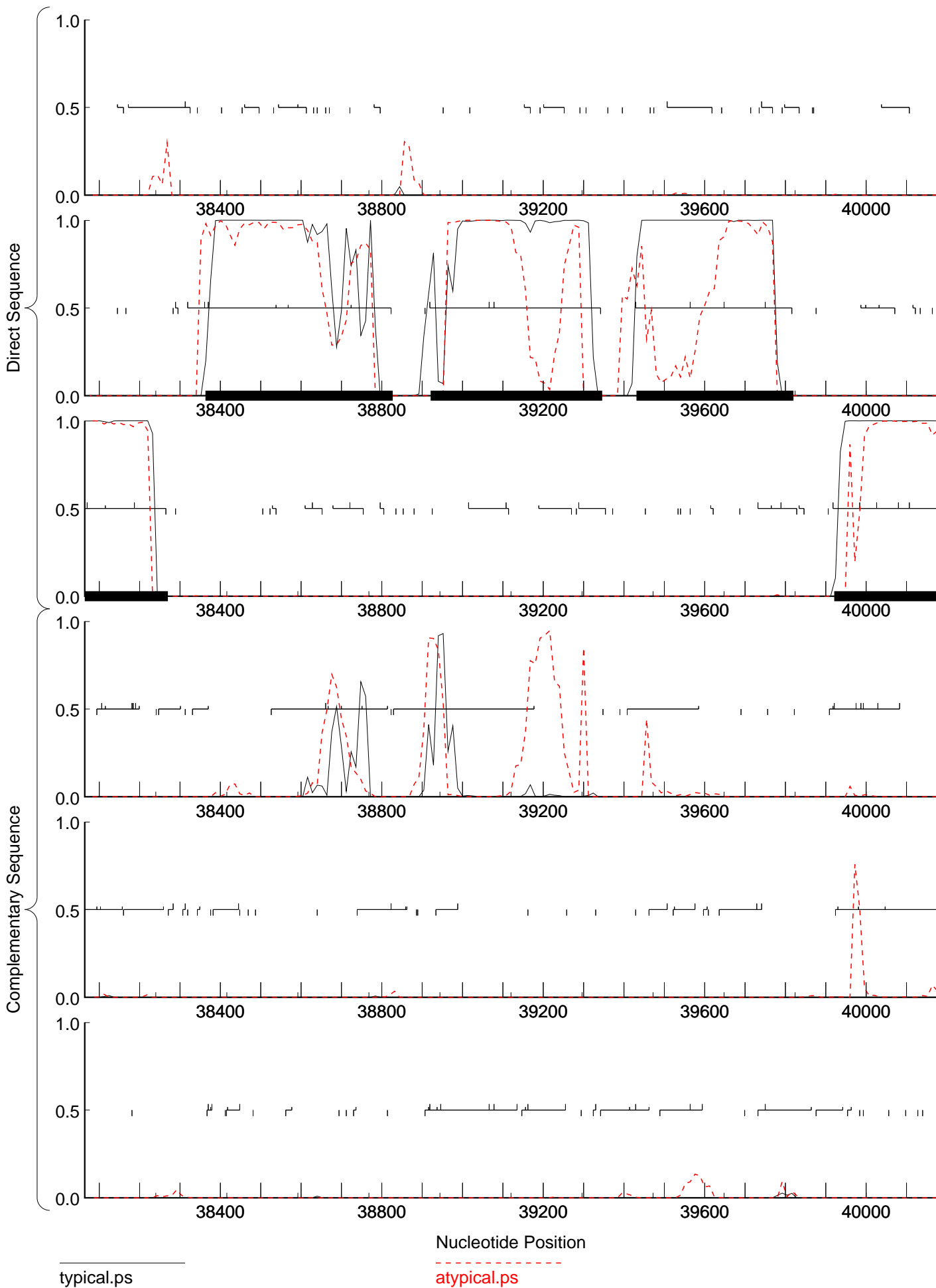
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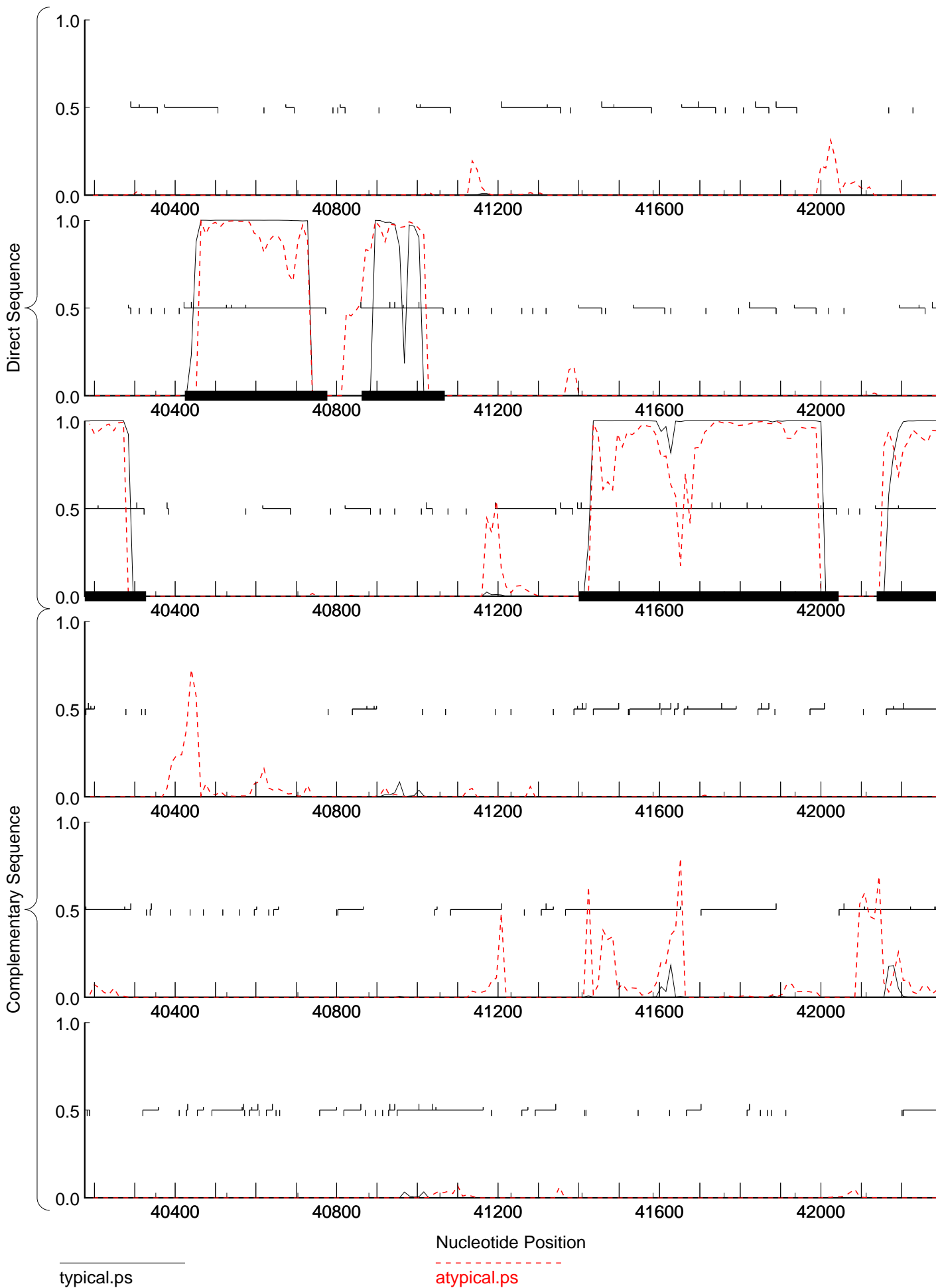
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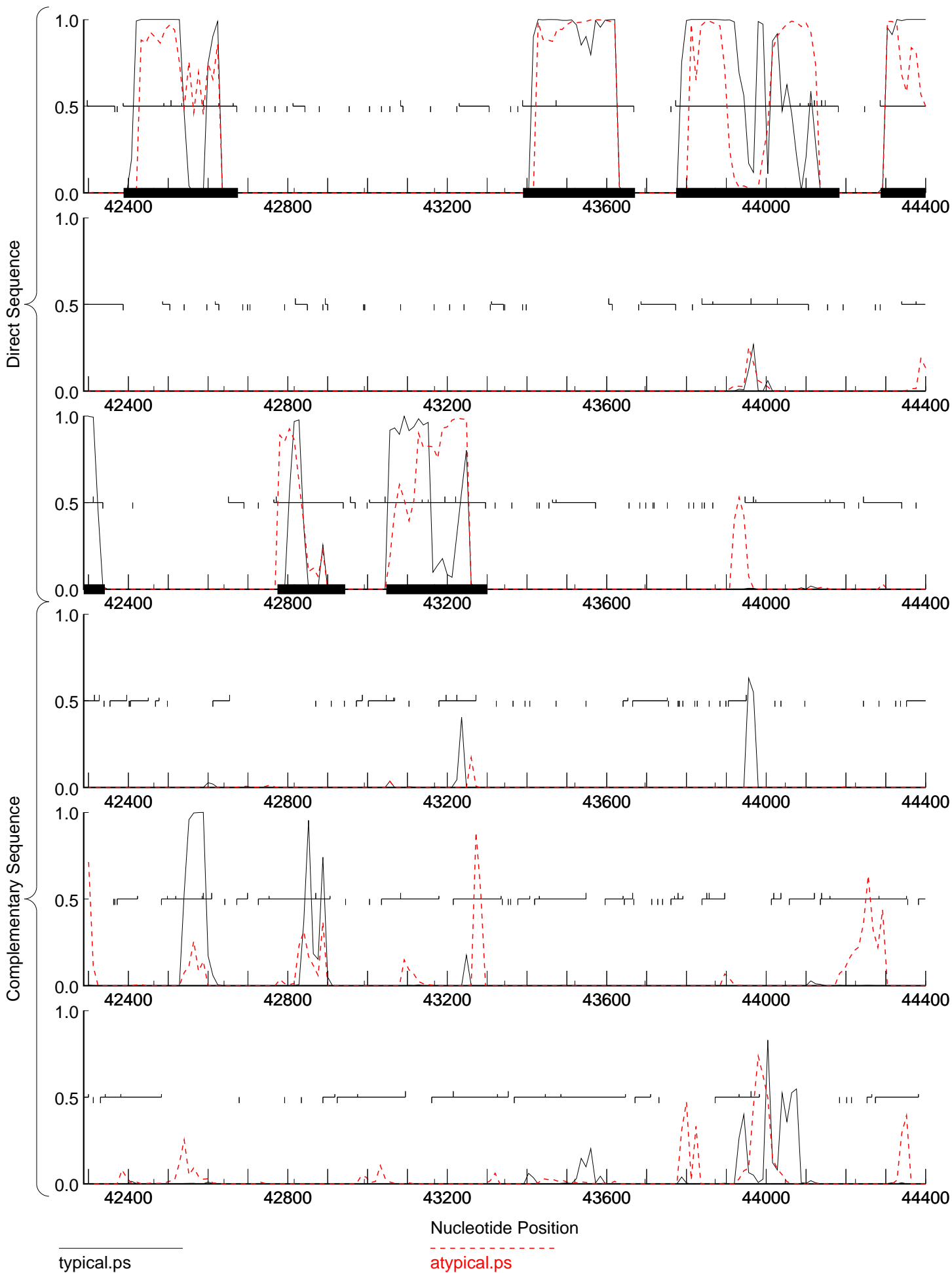


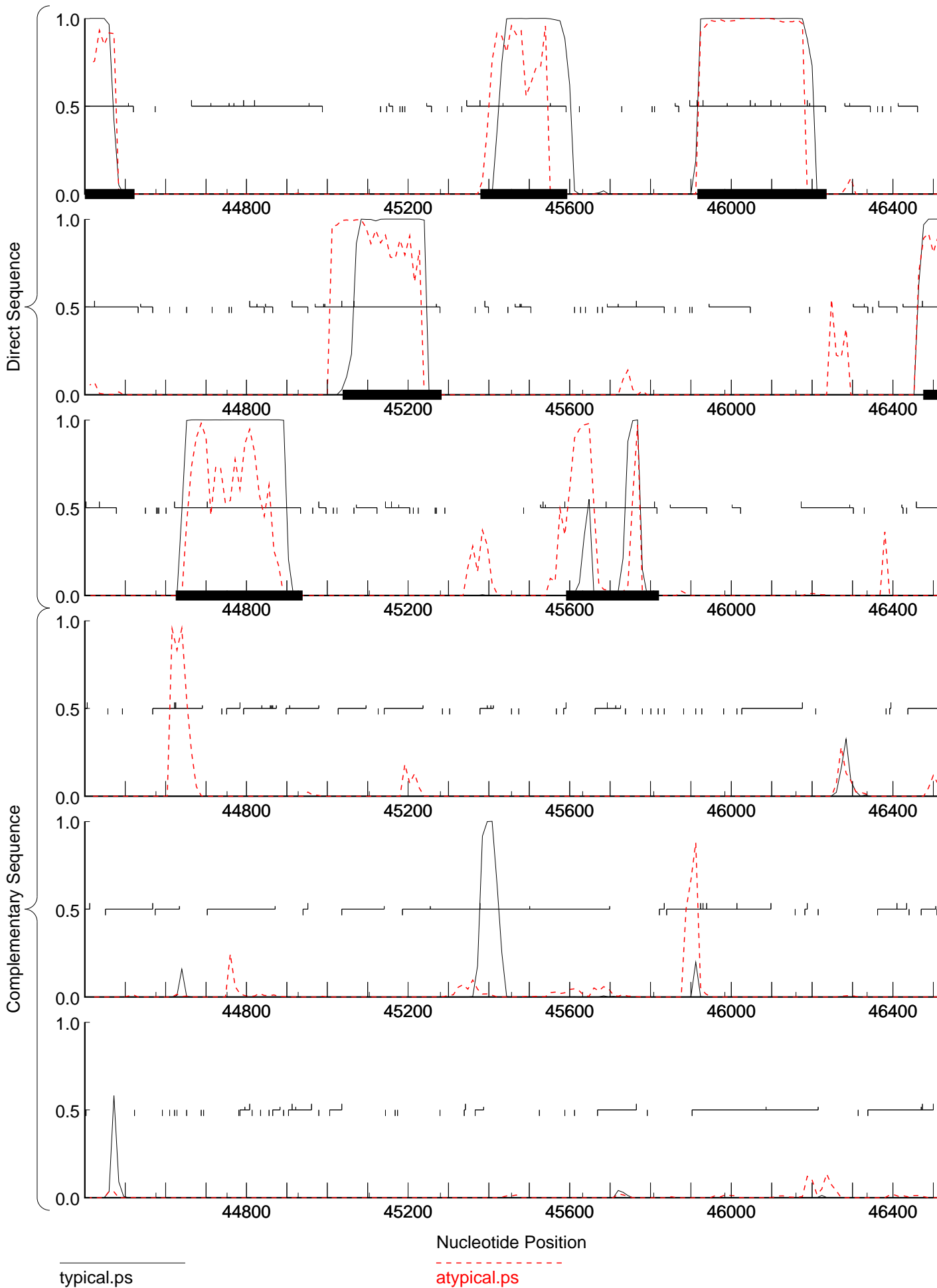


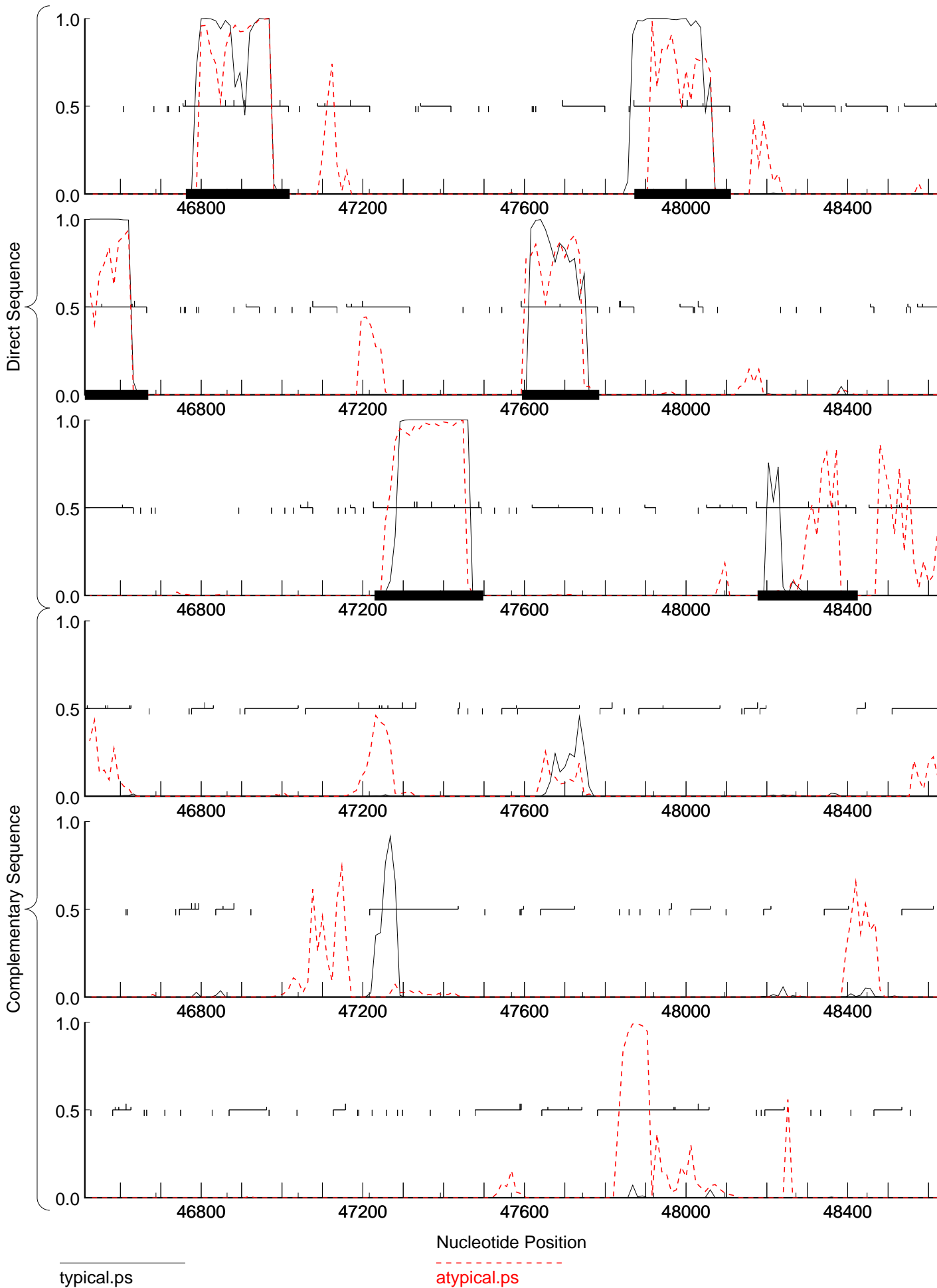
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atypical.ps





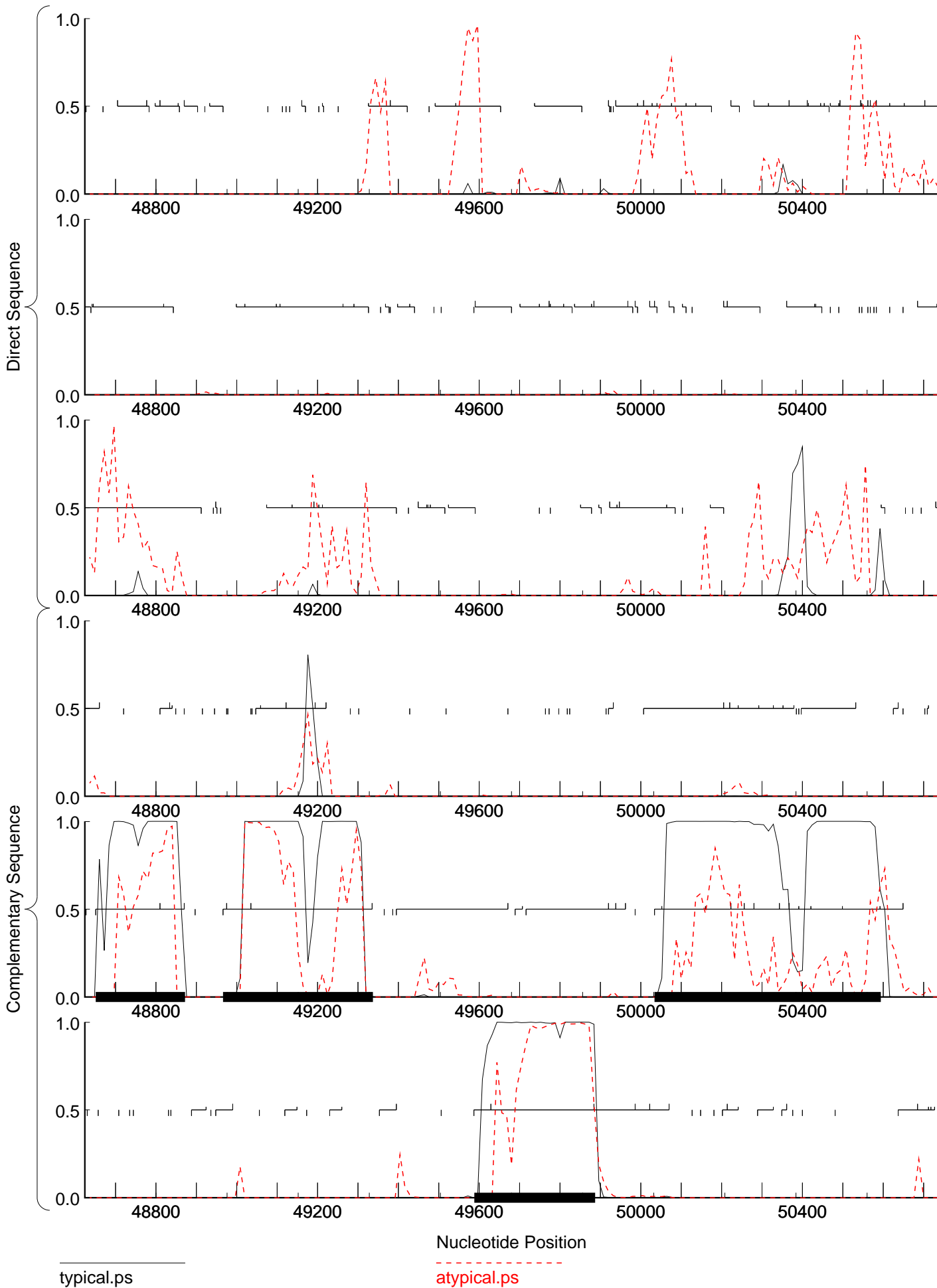


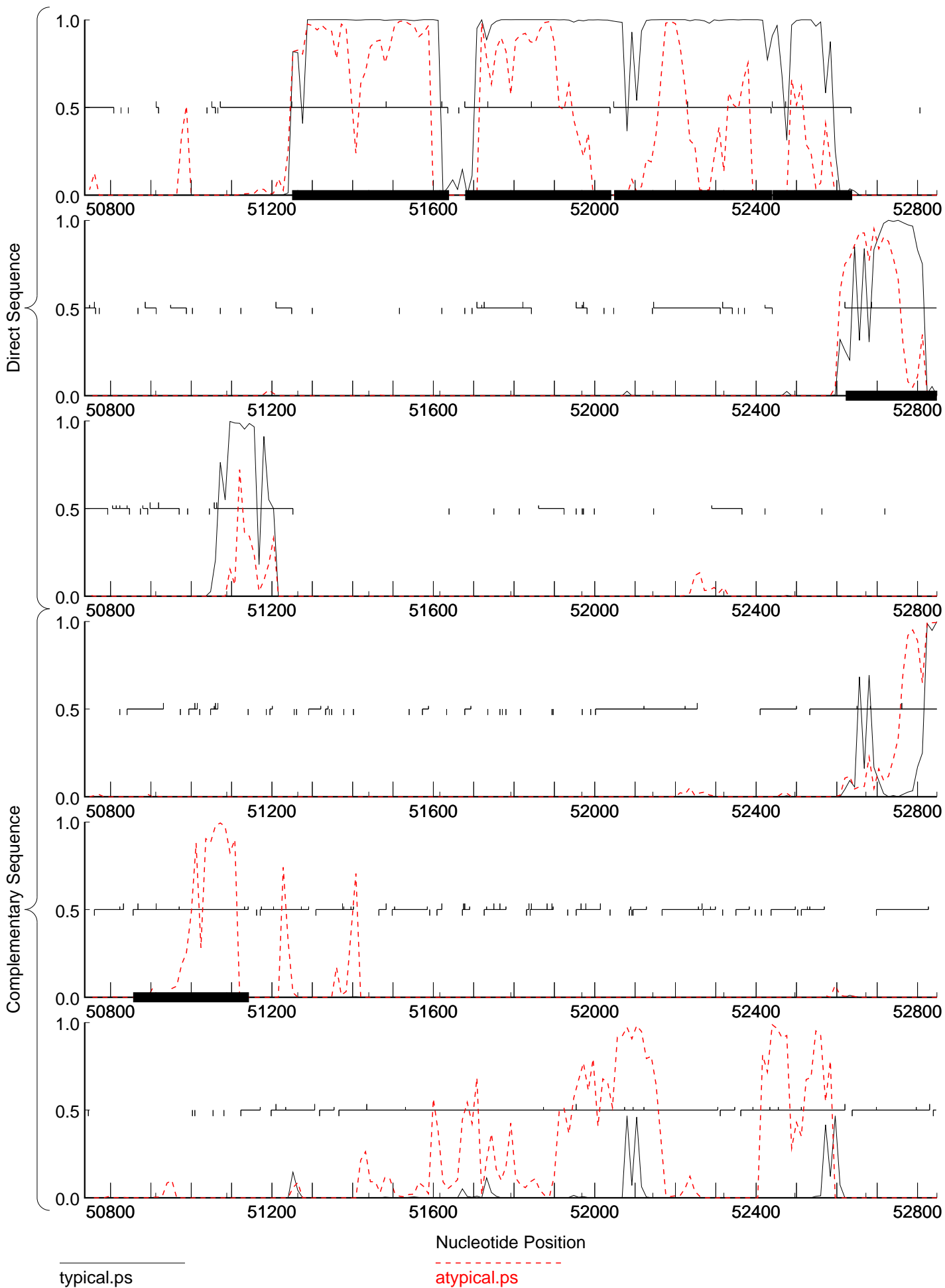


typical.ps

atypical.ps

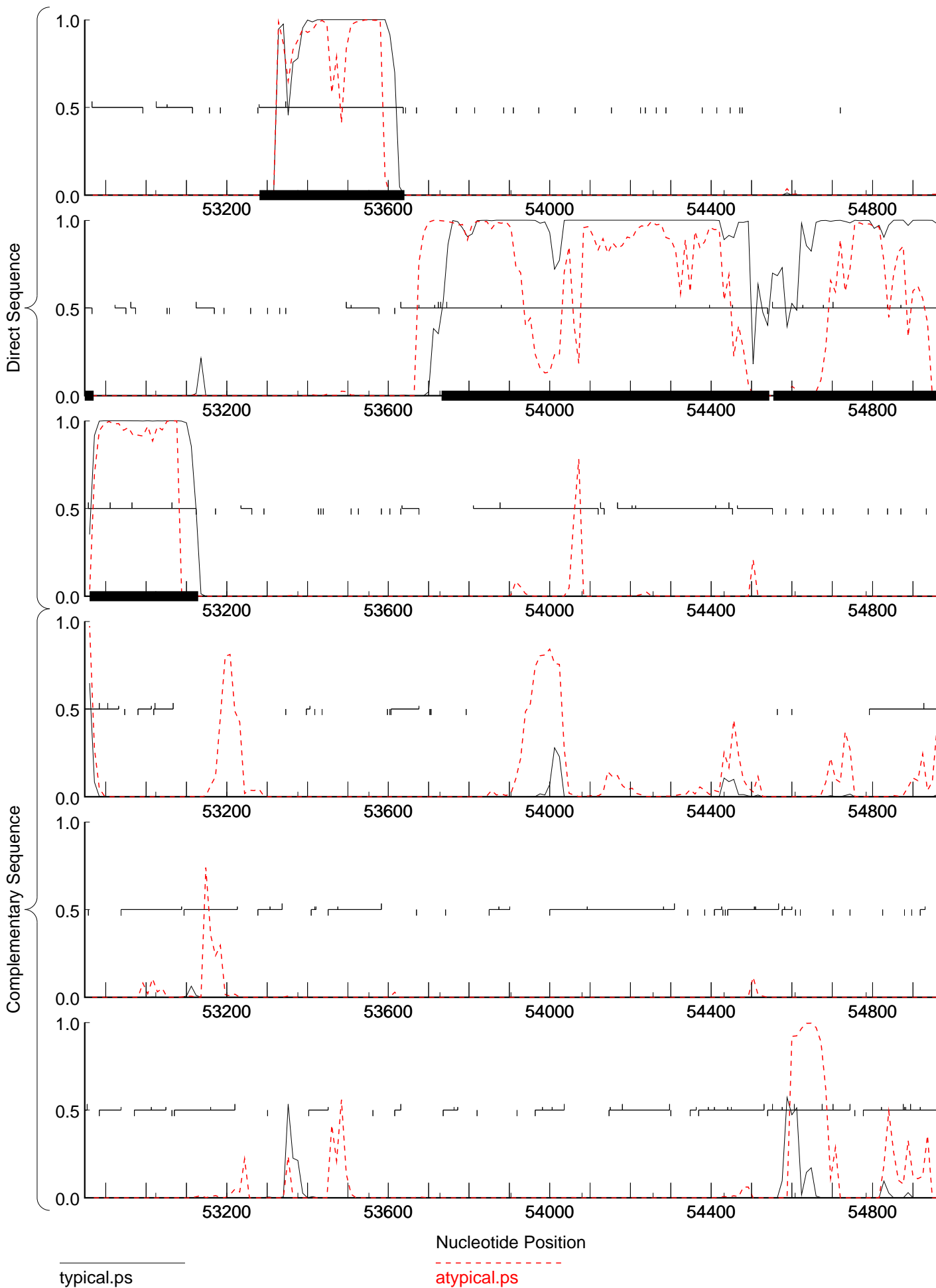






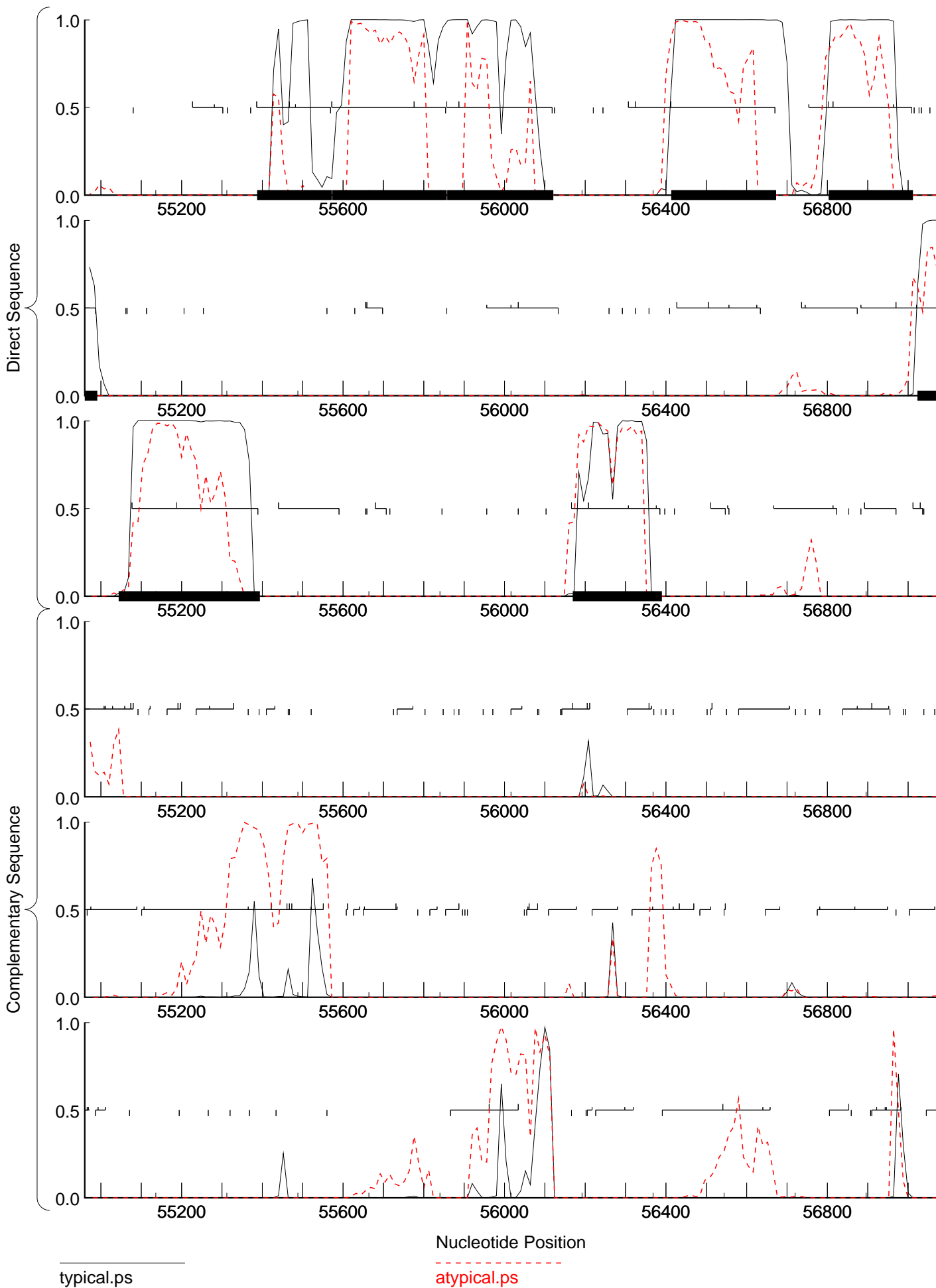
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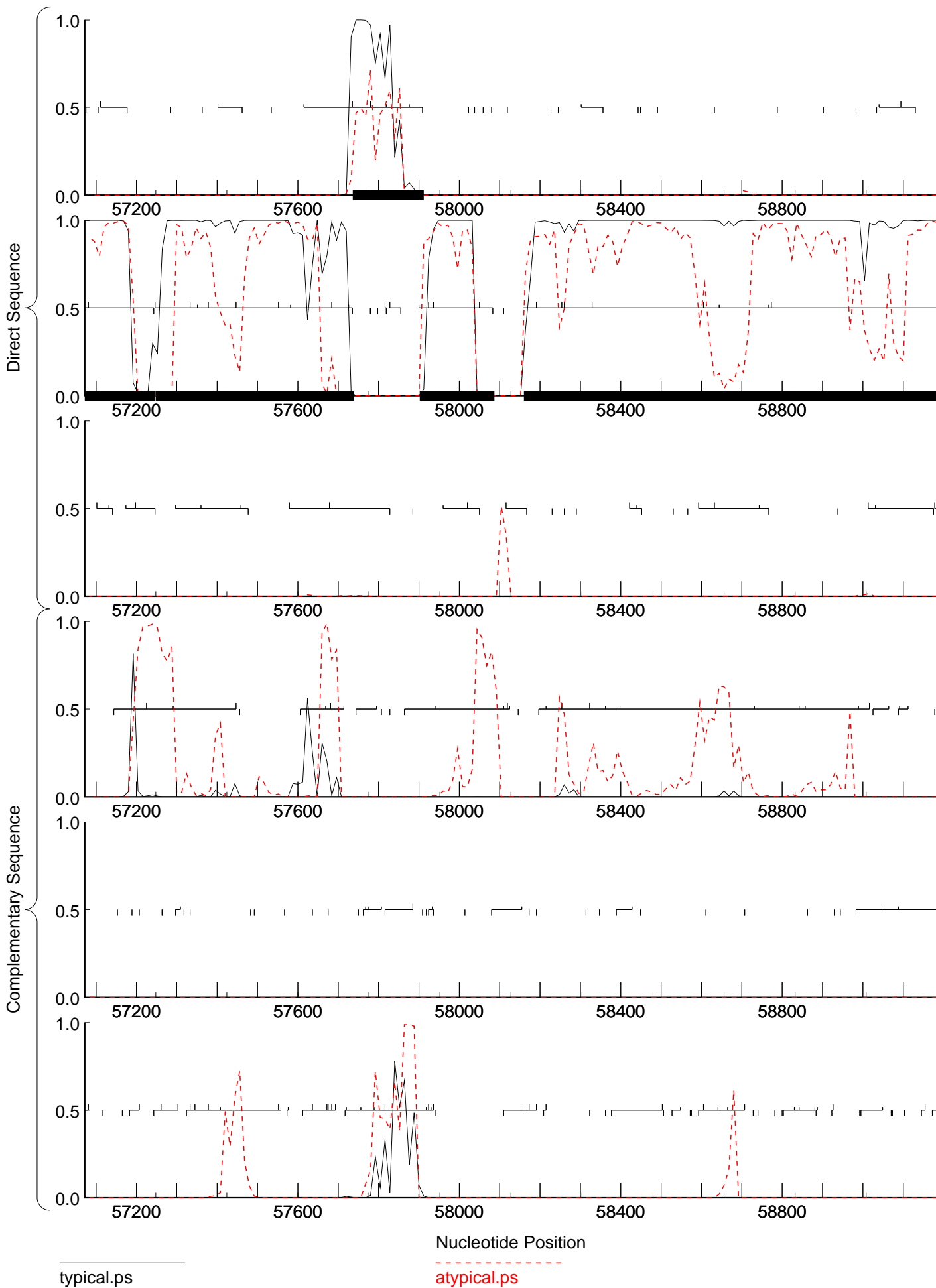
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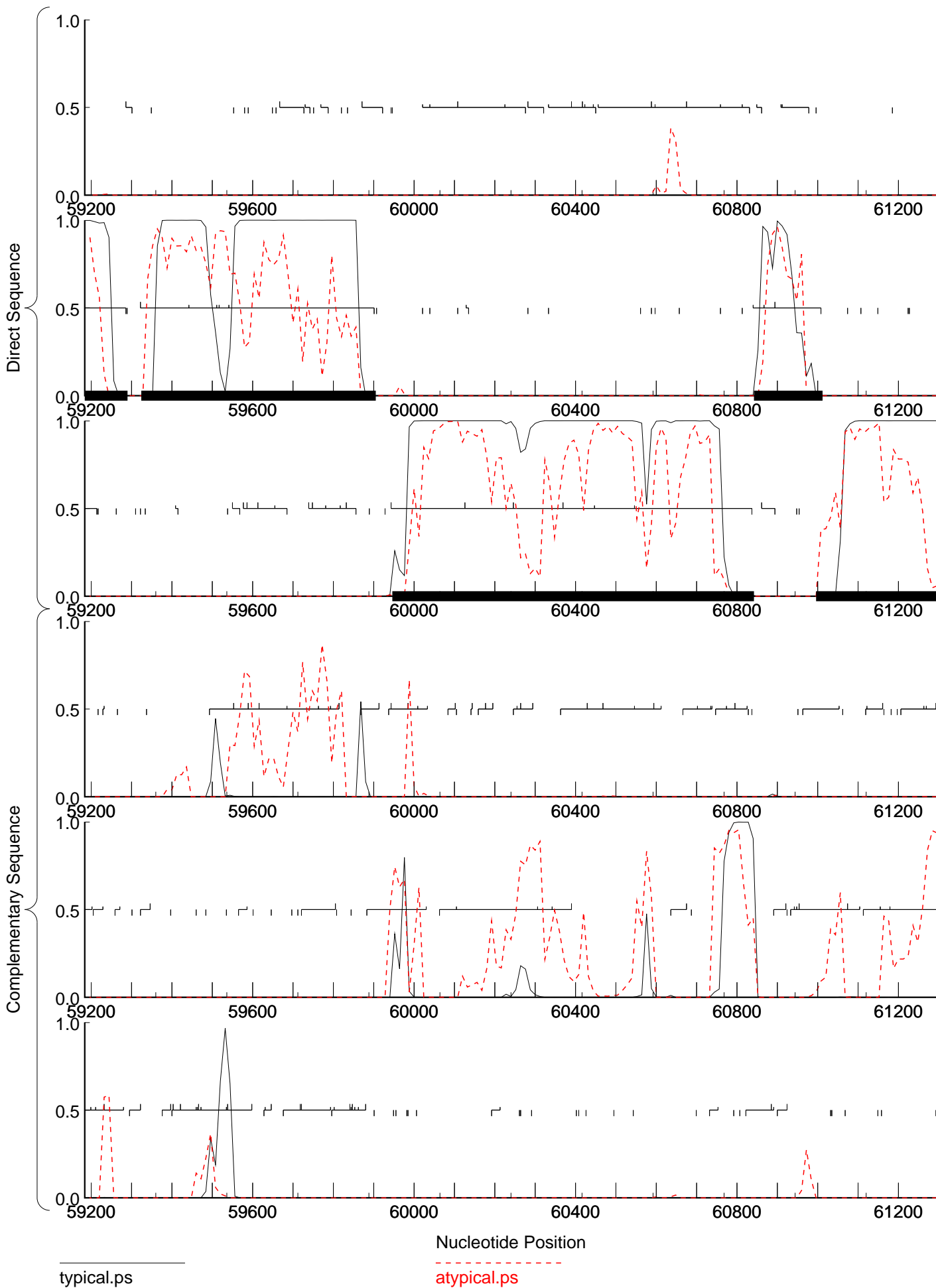
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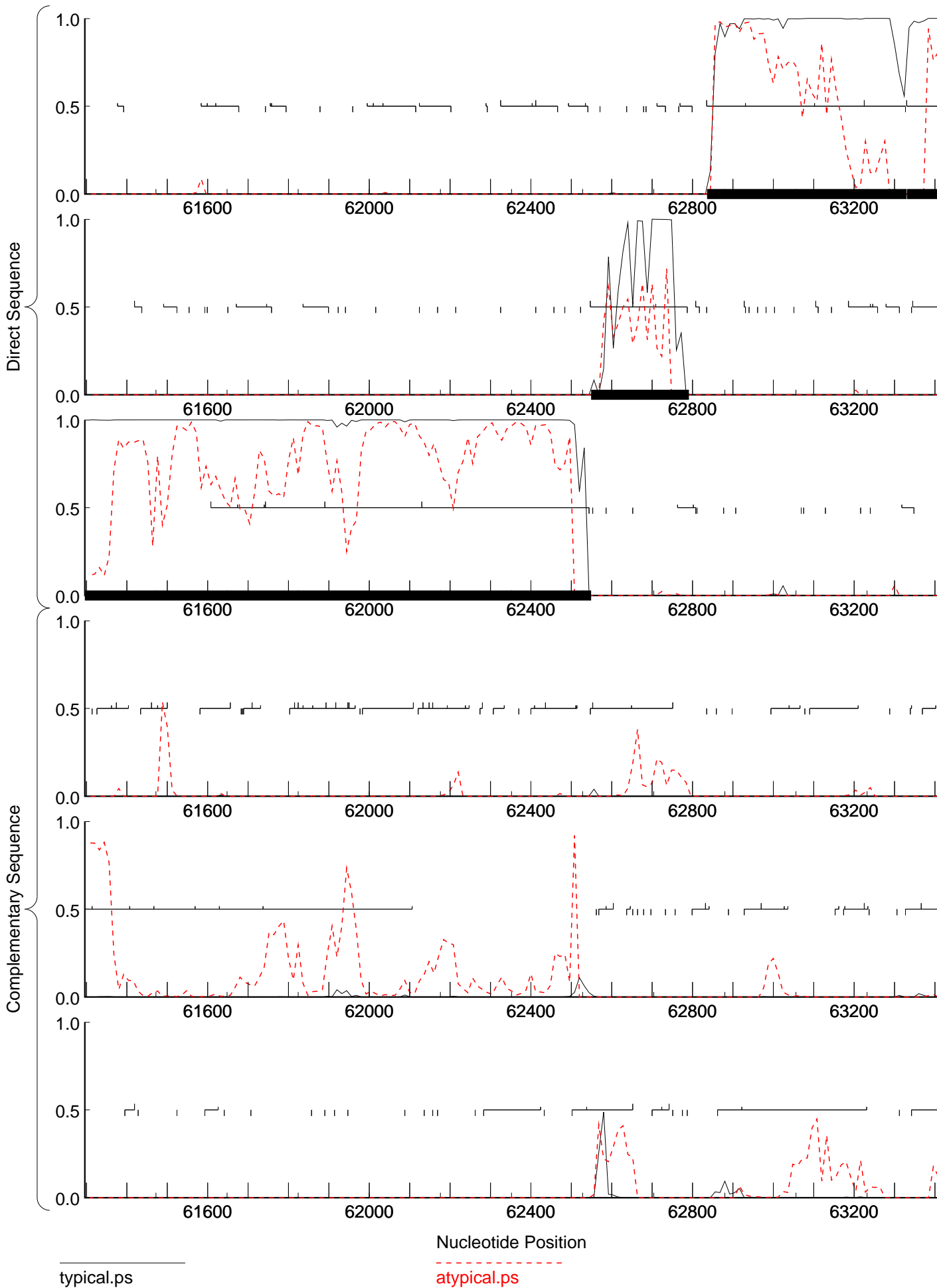


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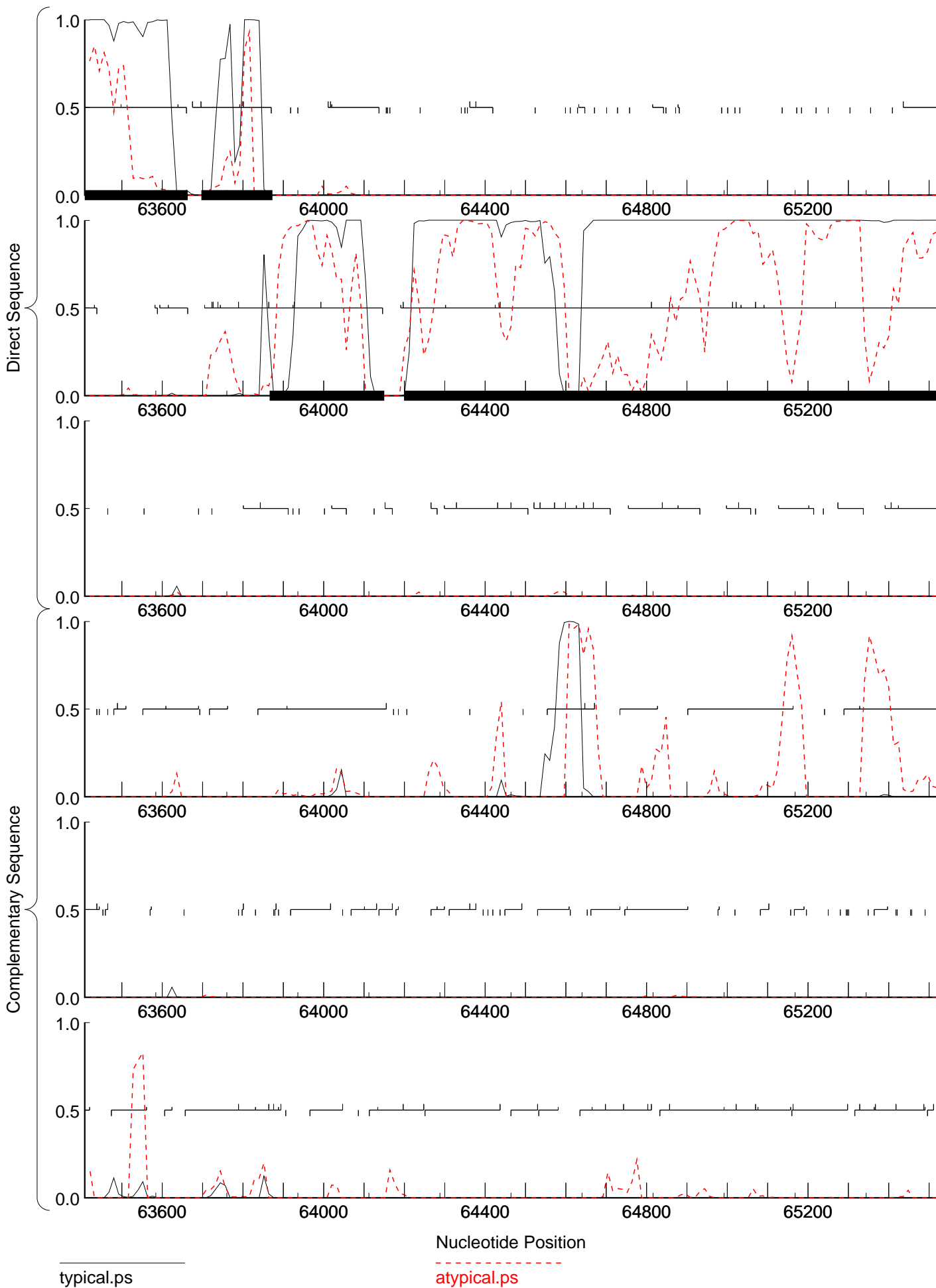




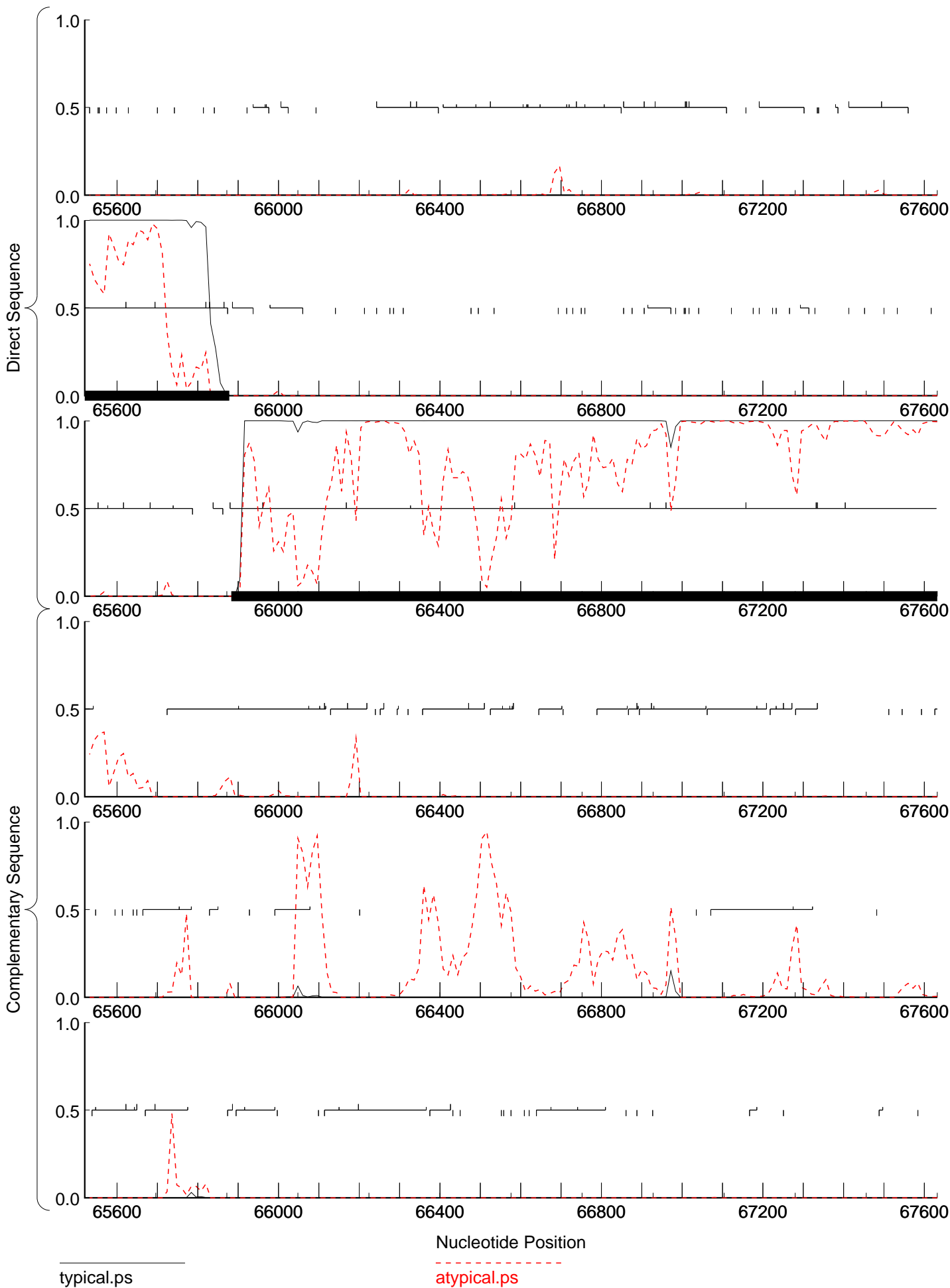


typical.ps

atypical.ps

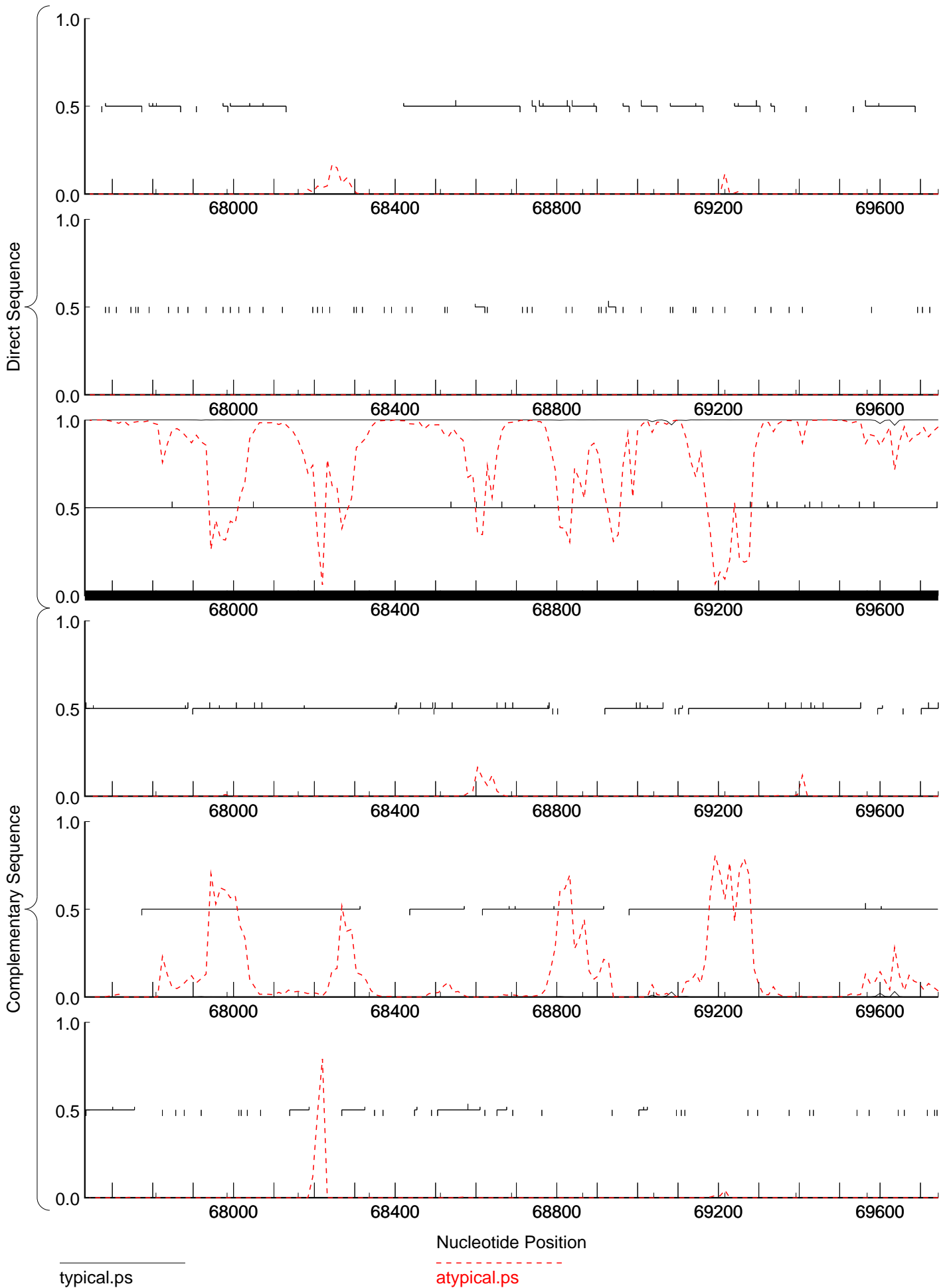


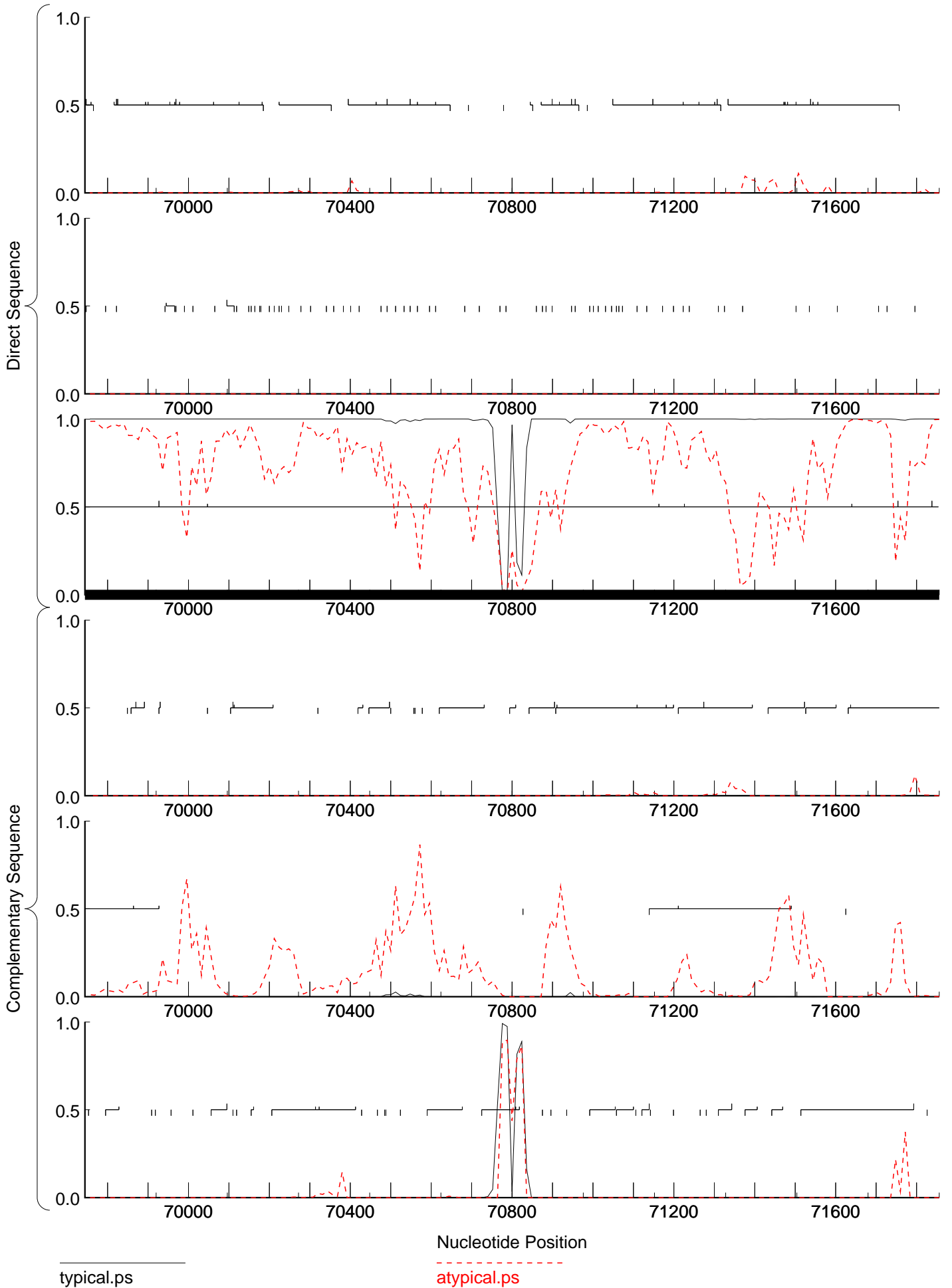


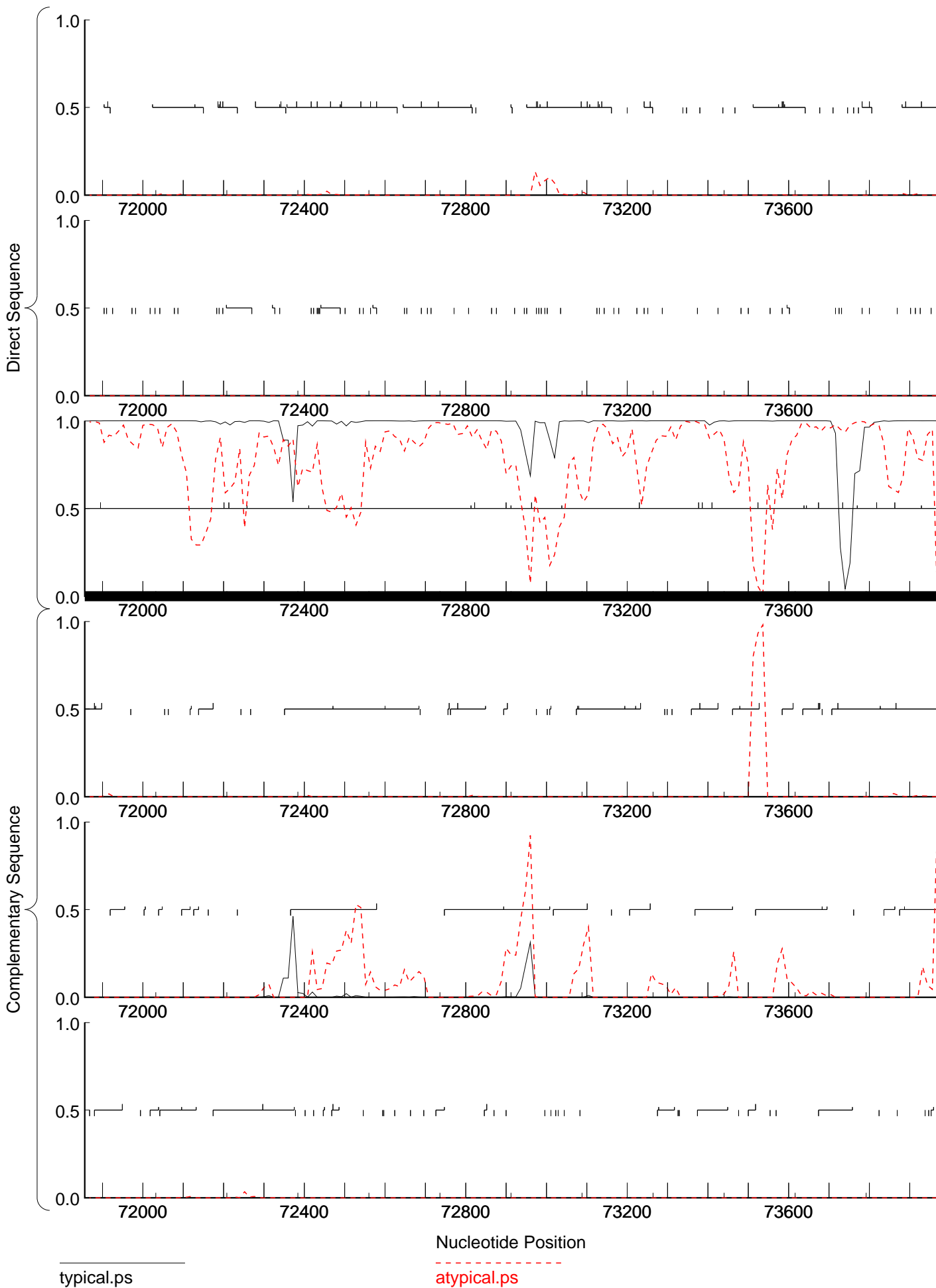


typical.ps

atypical.ps

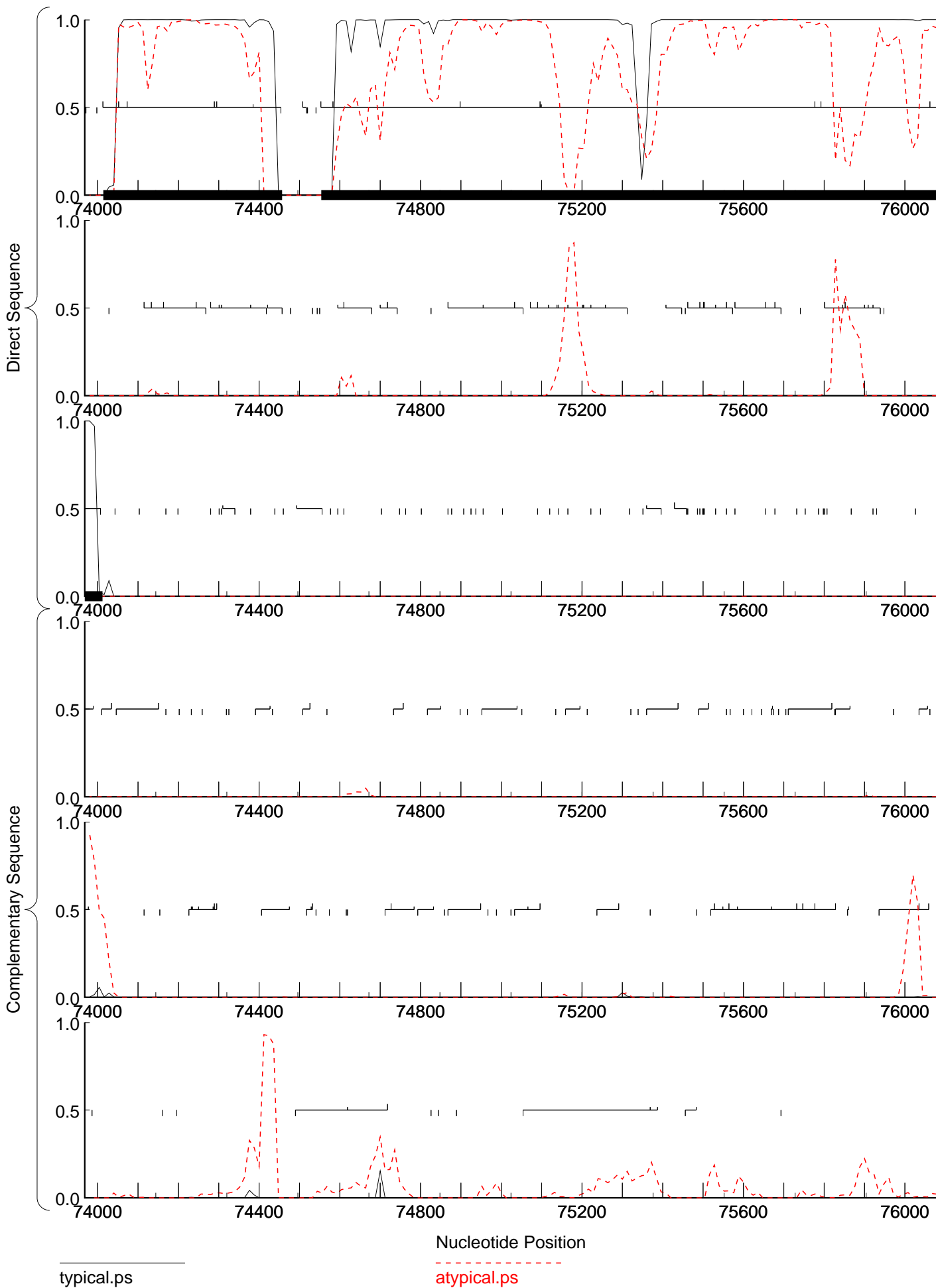






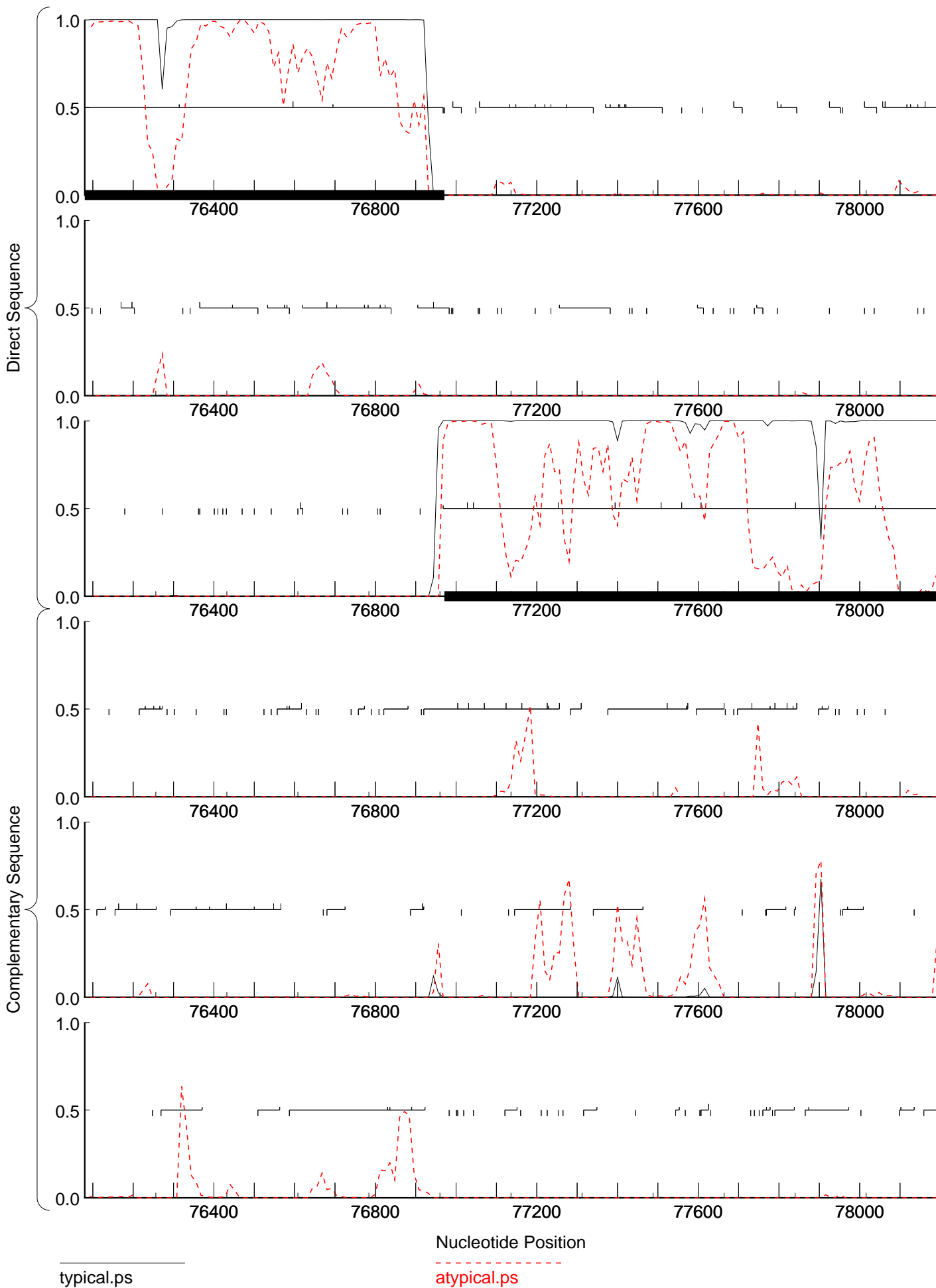
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atypical.ps



typical.ps

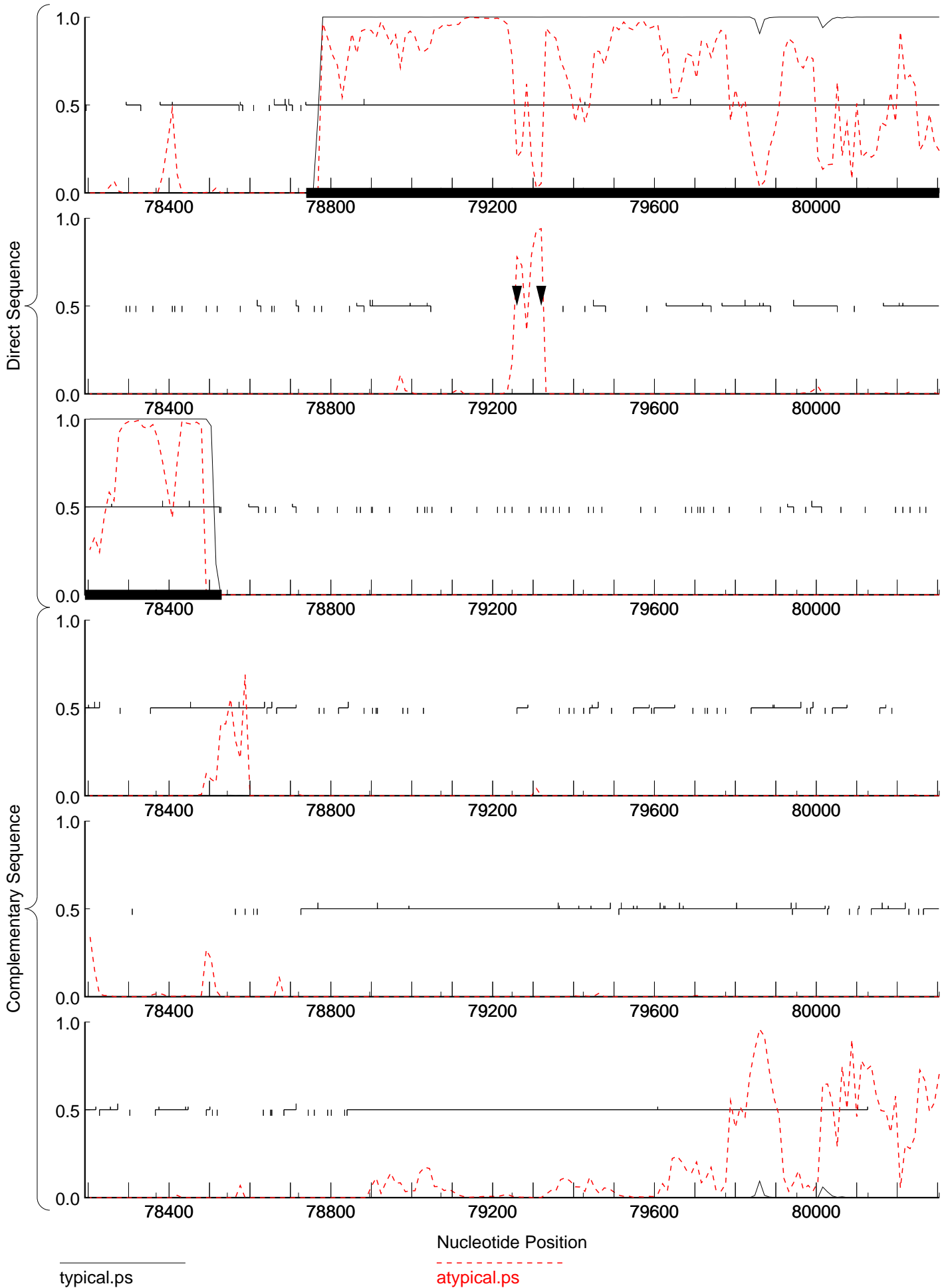
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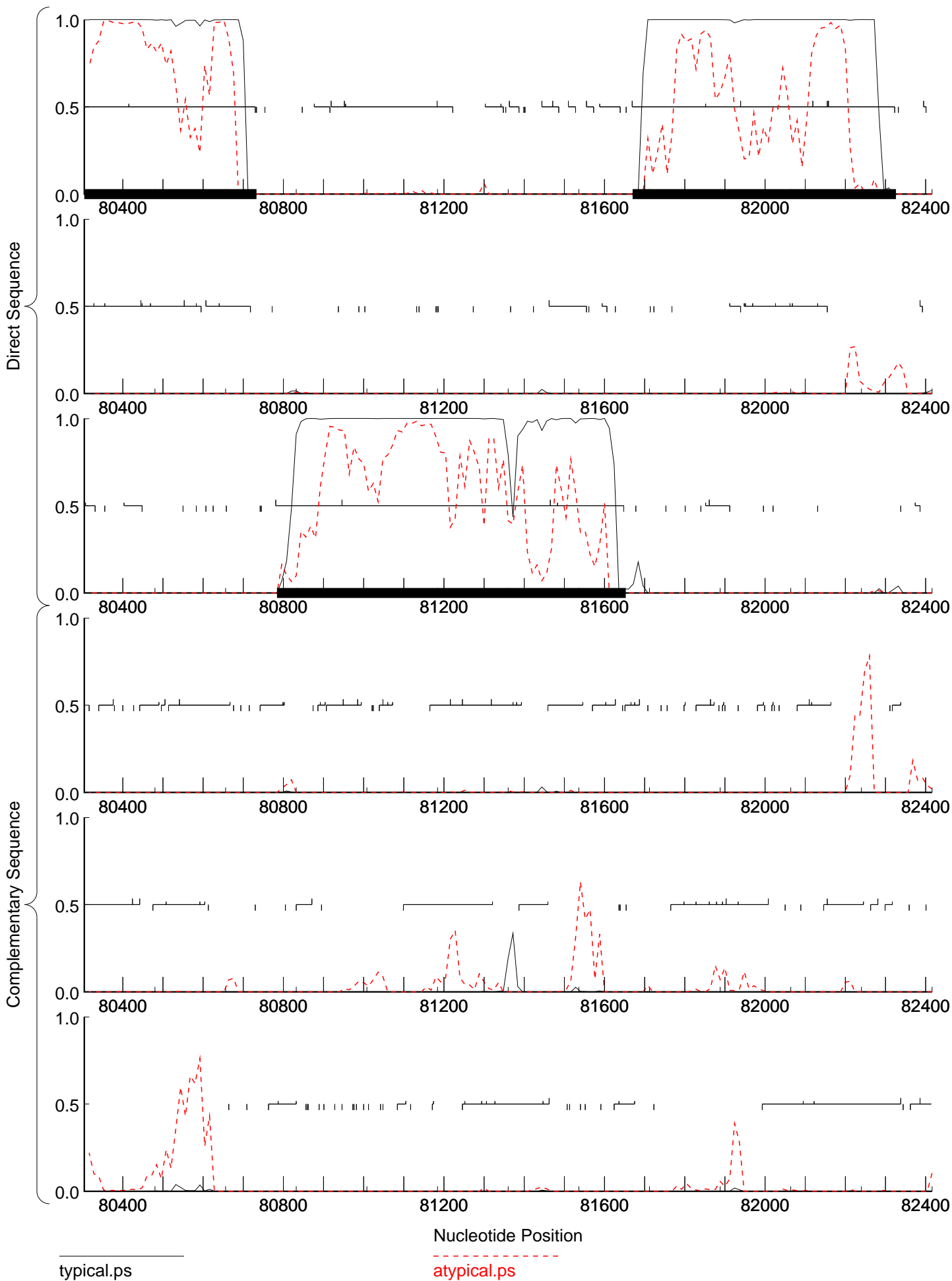


typical.ps

Nucleotide Position

atypical.ps

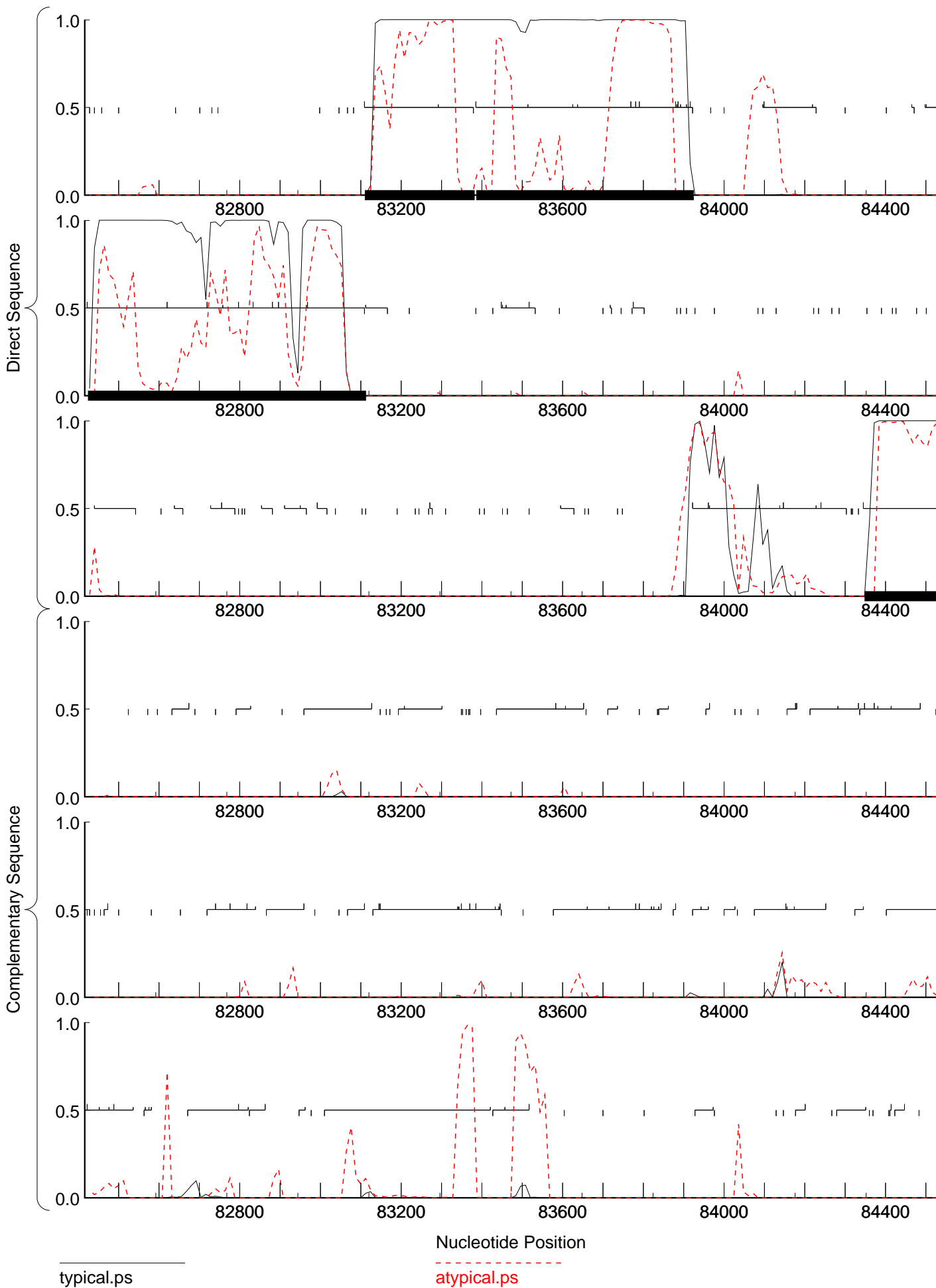




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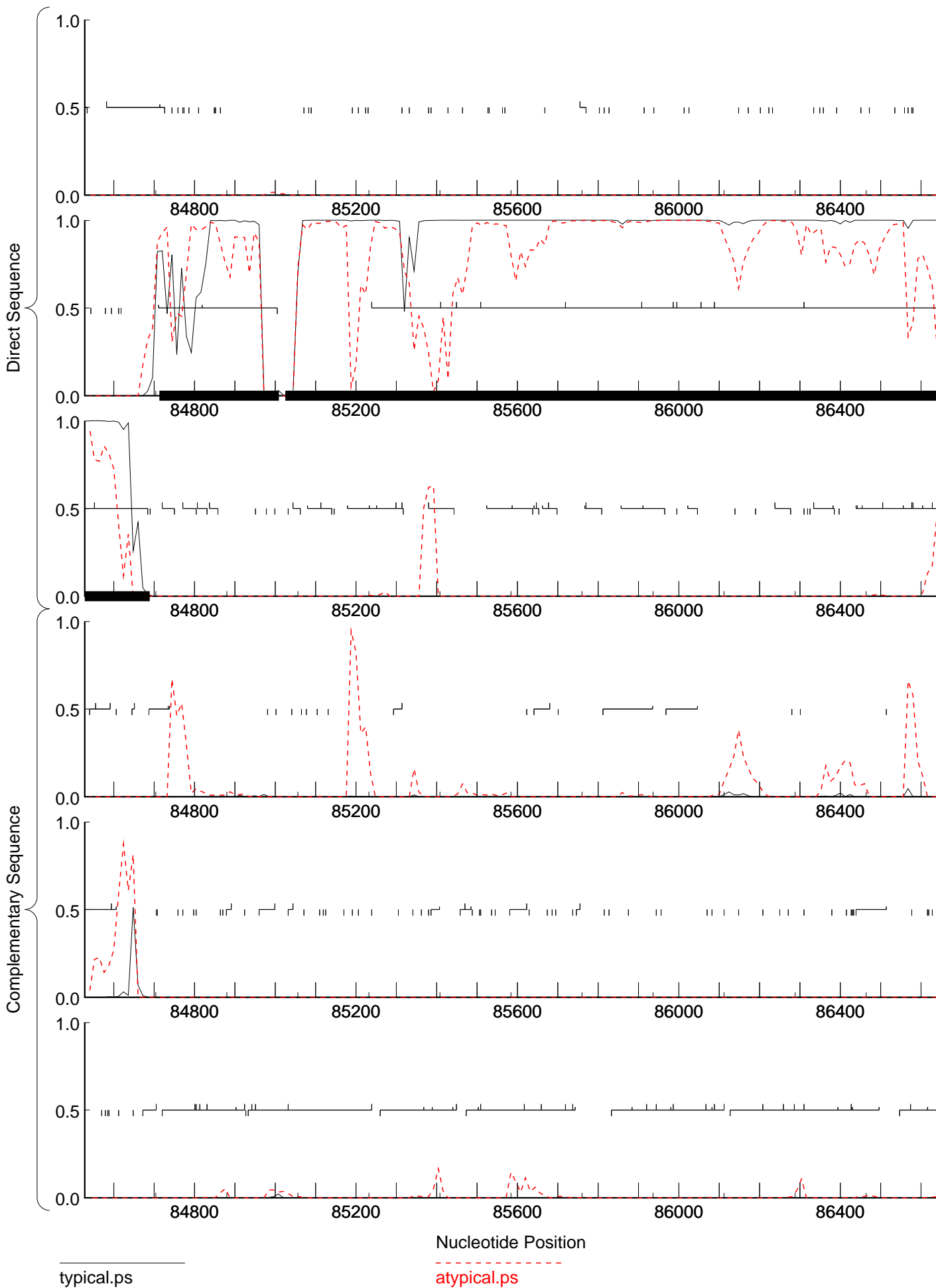
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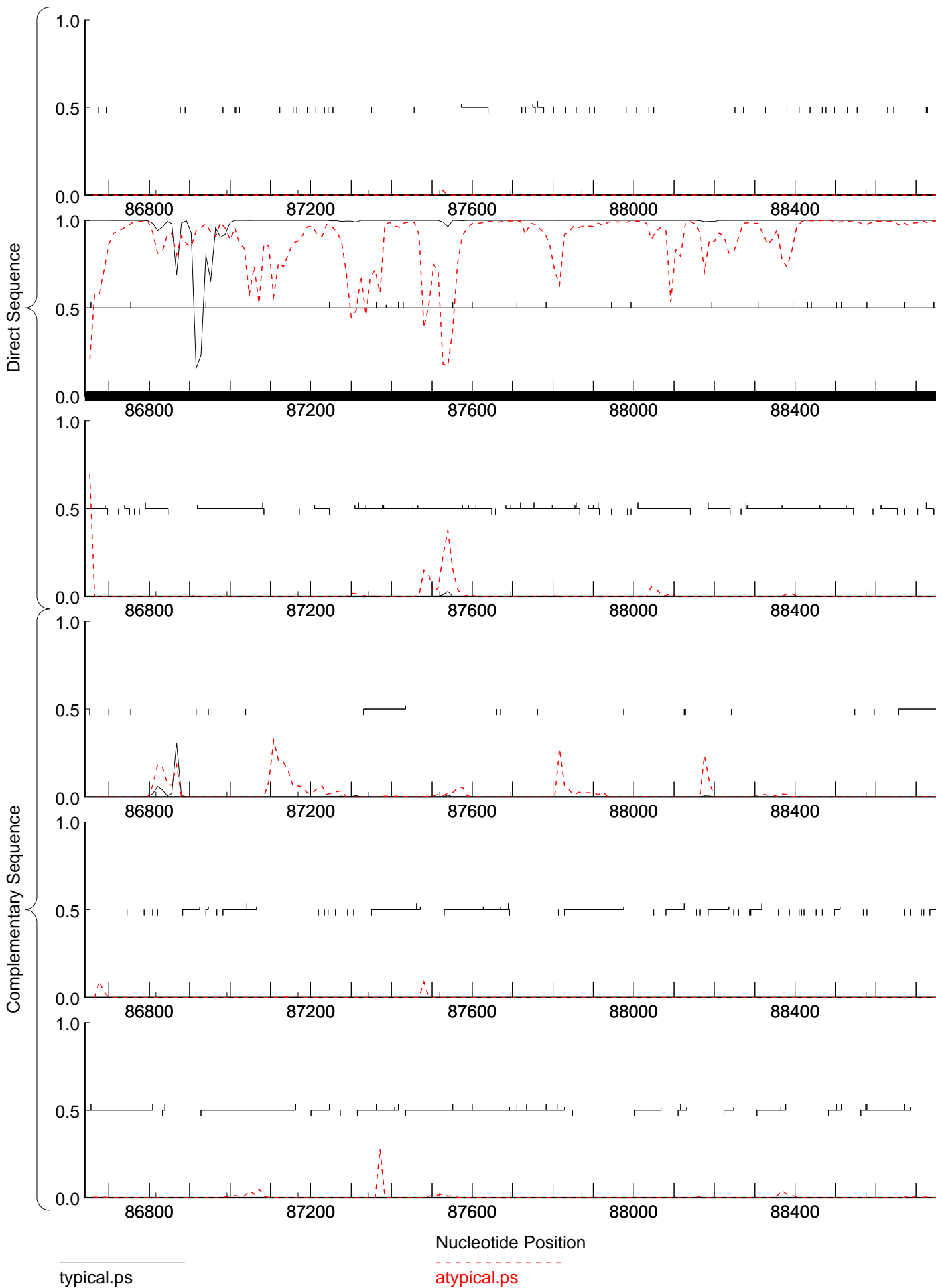
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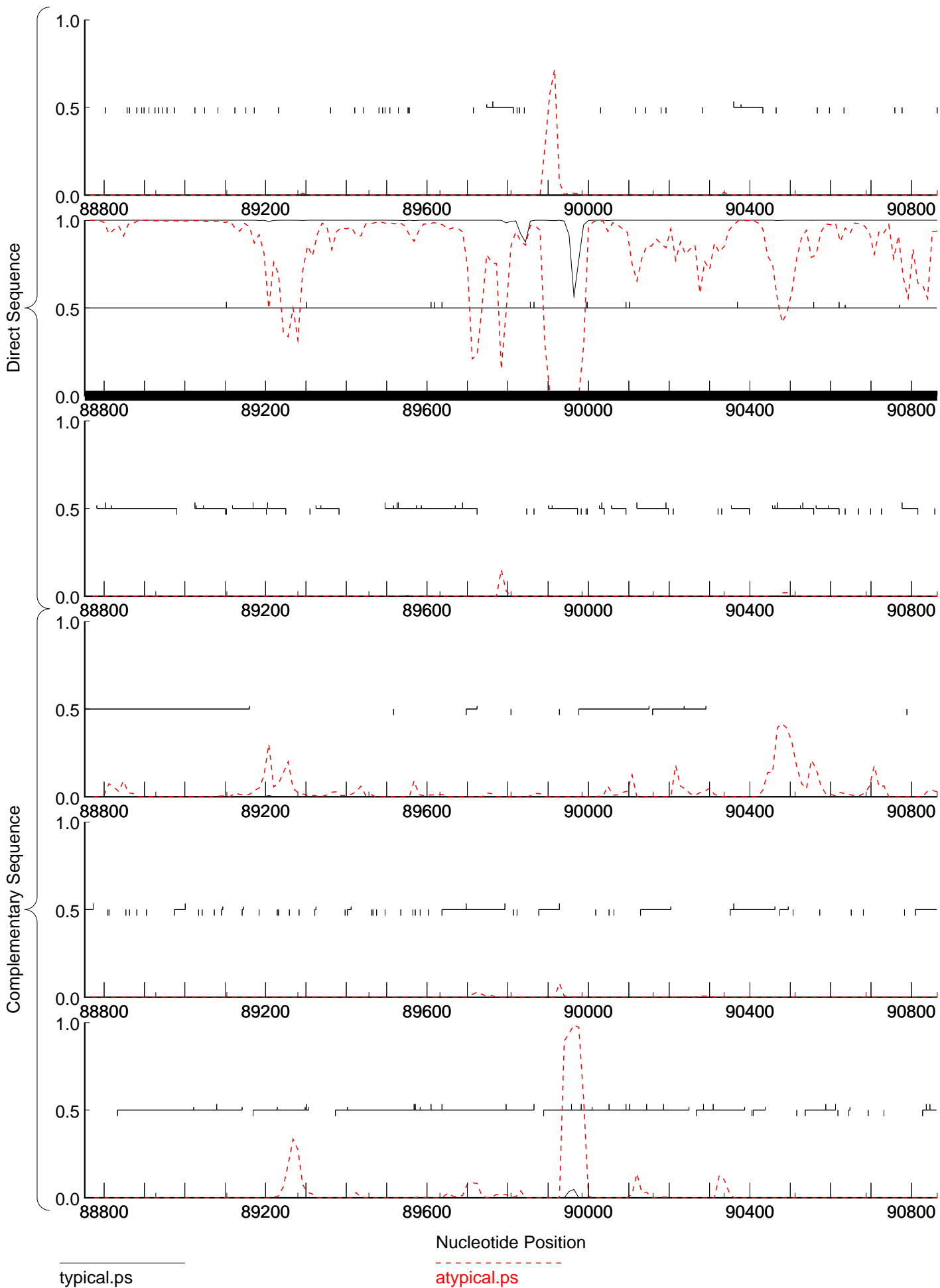
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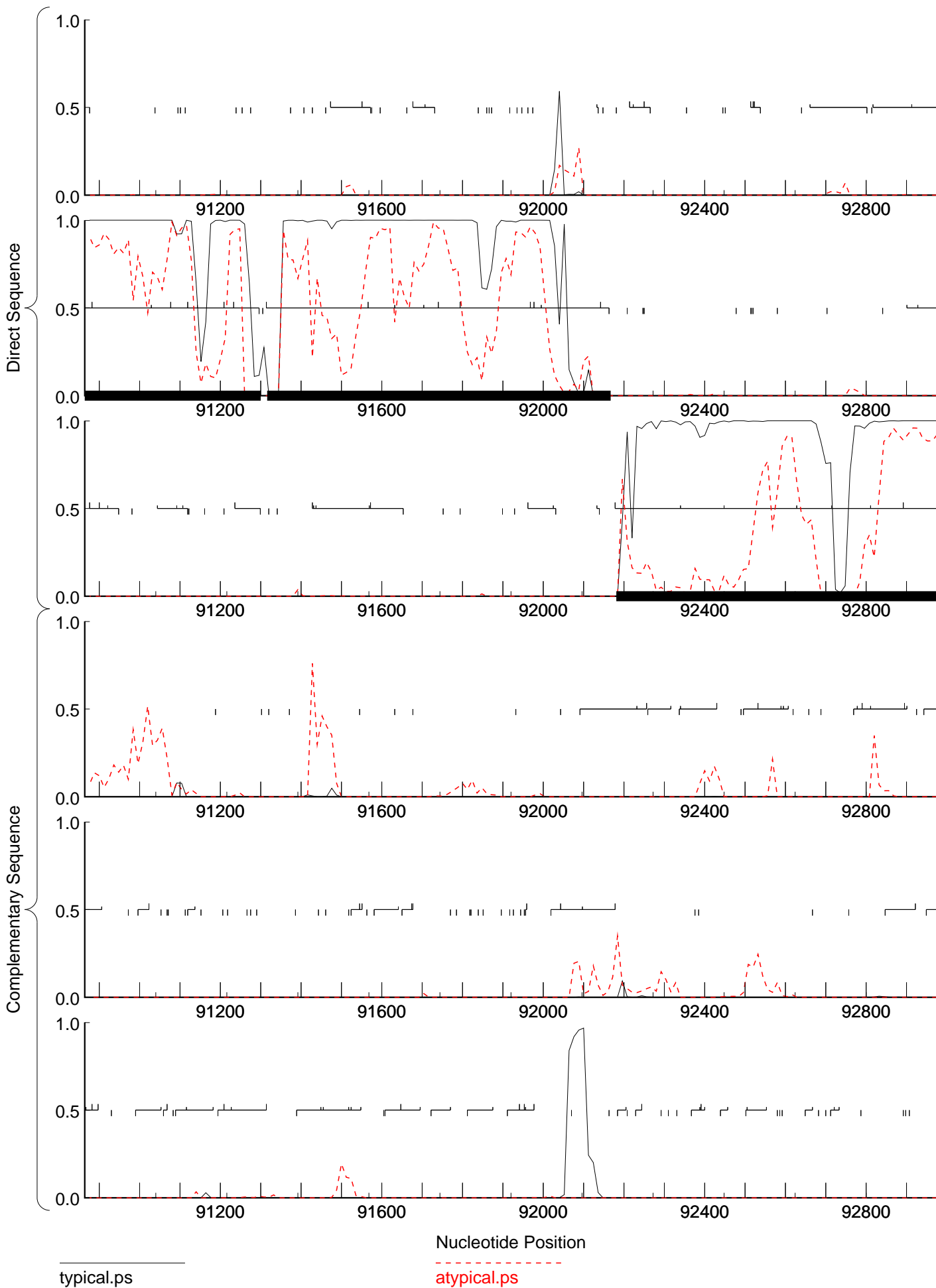


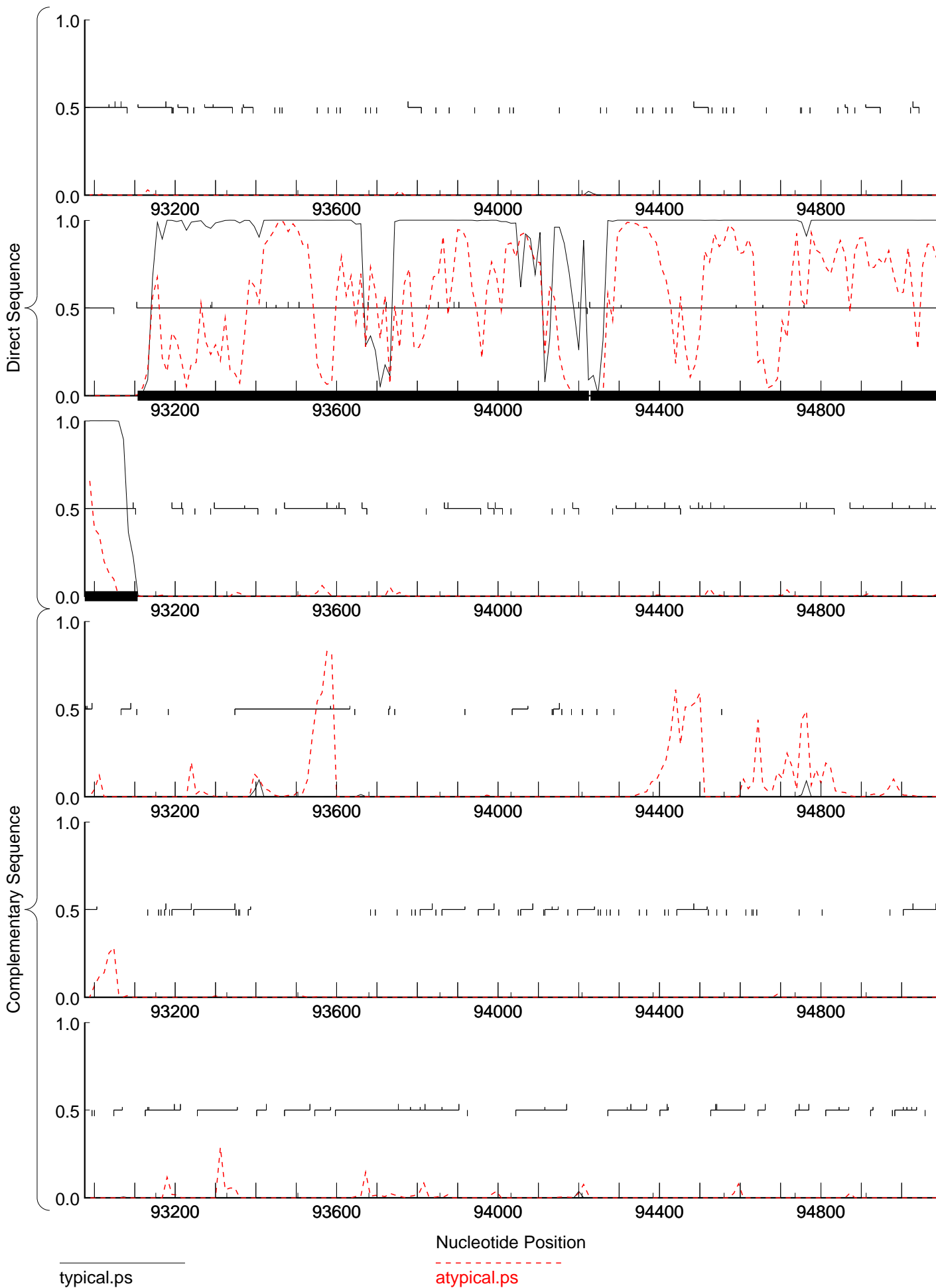
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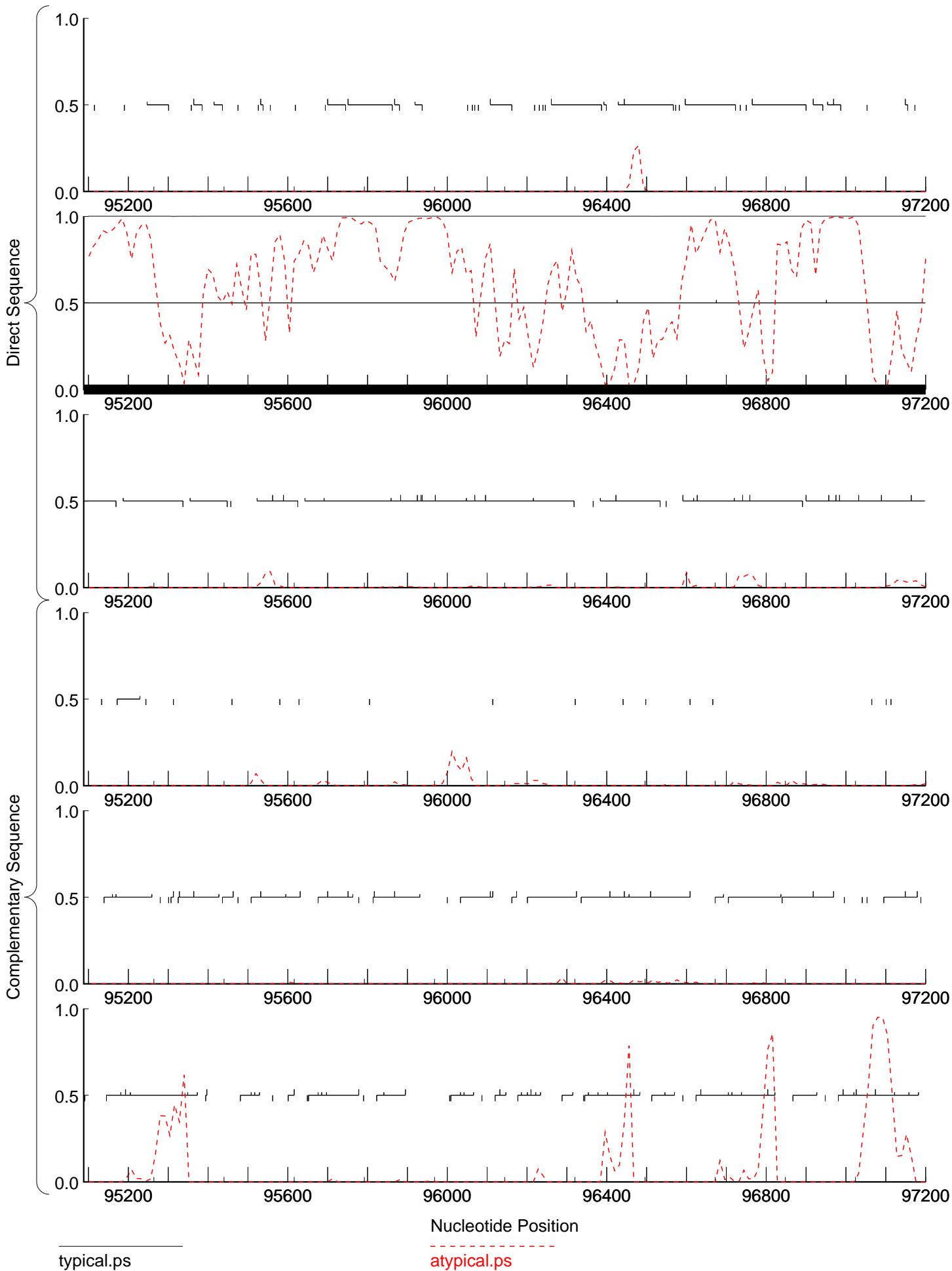
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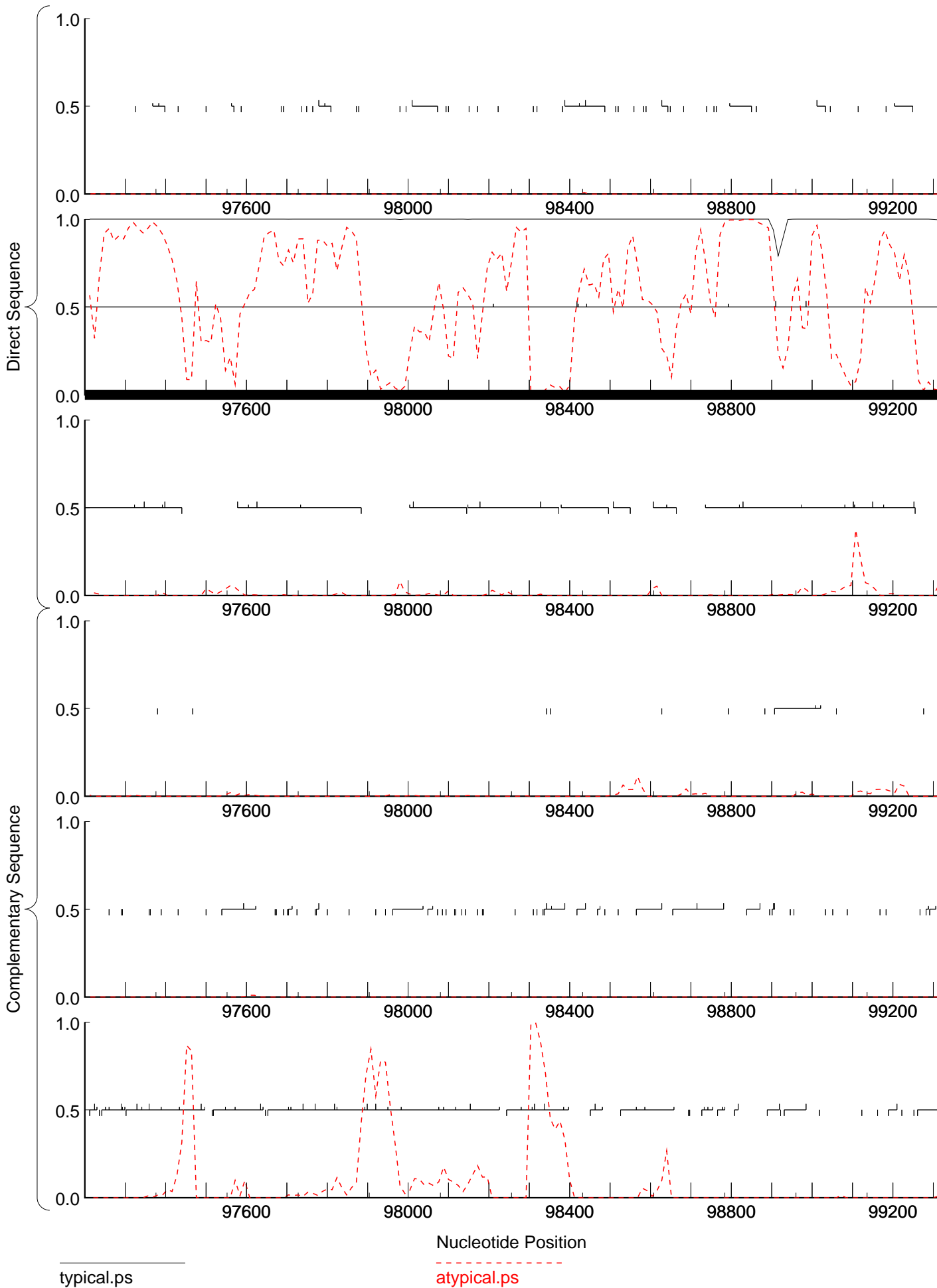




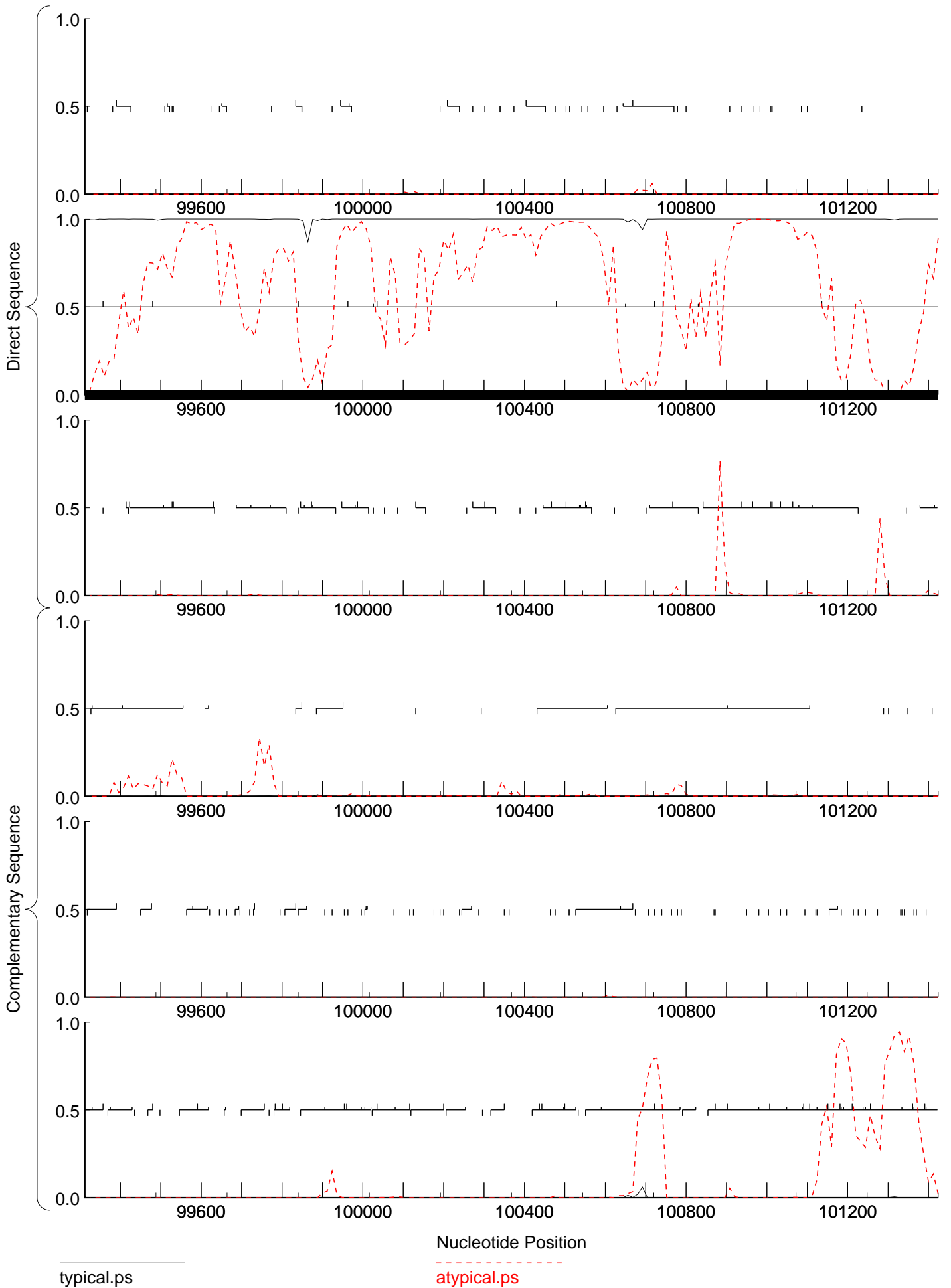


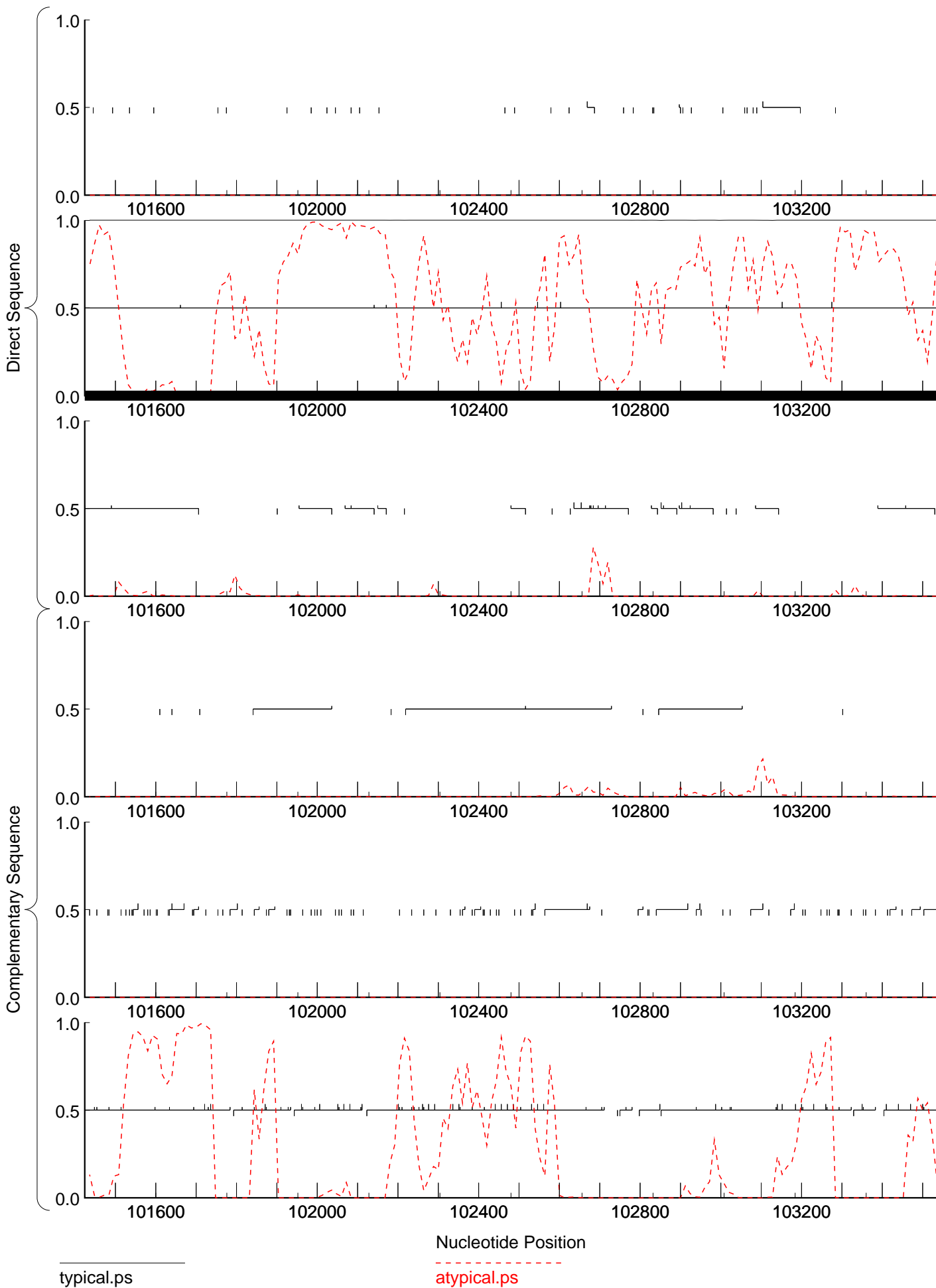






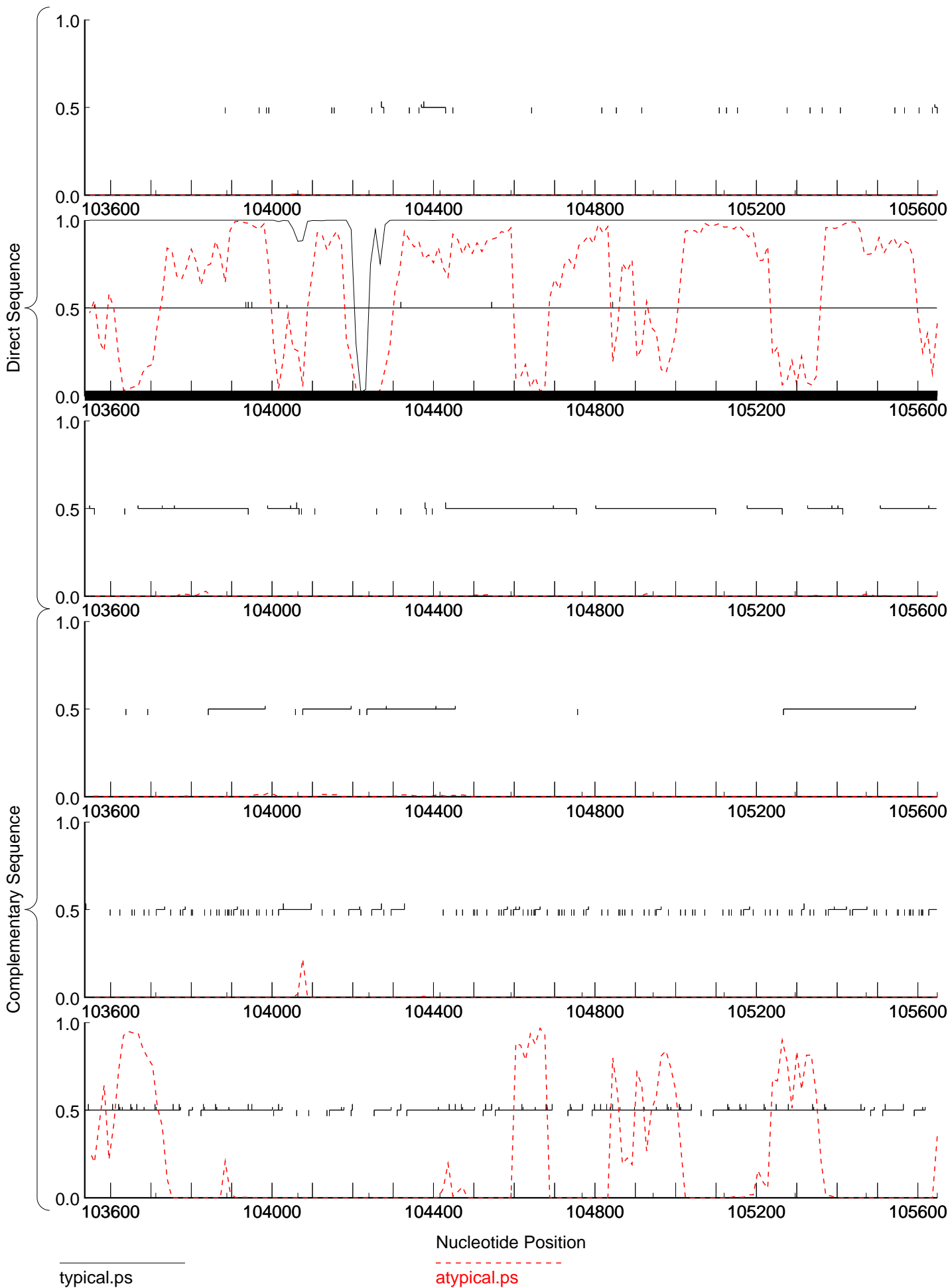






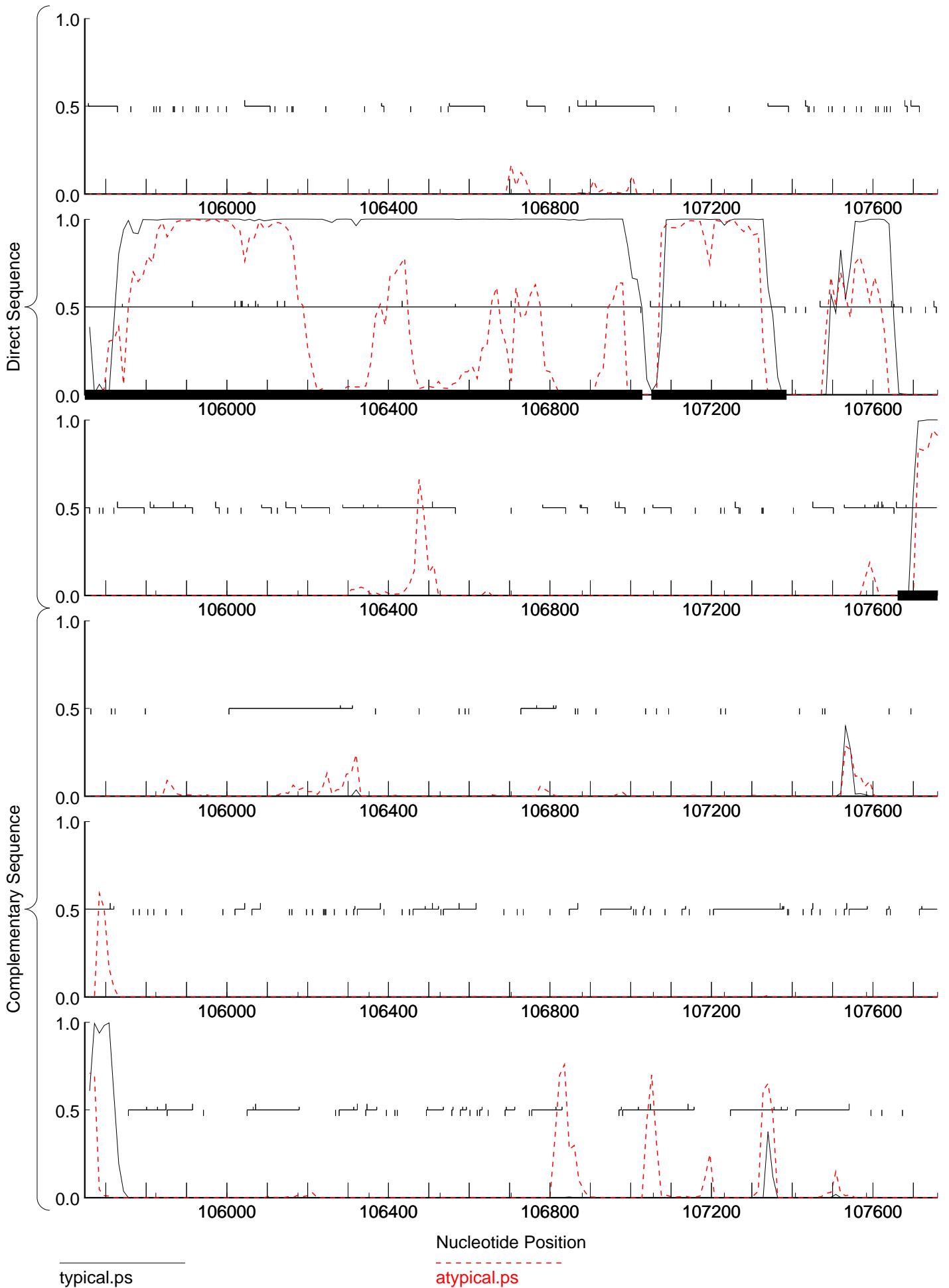
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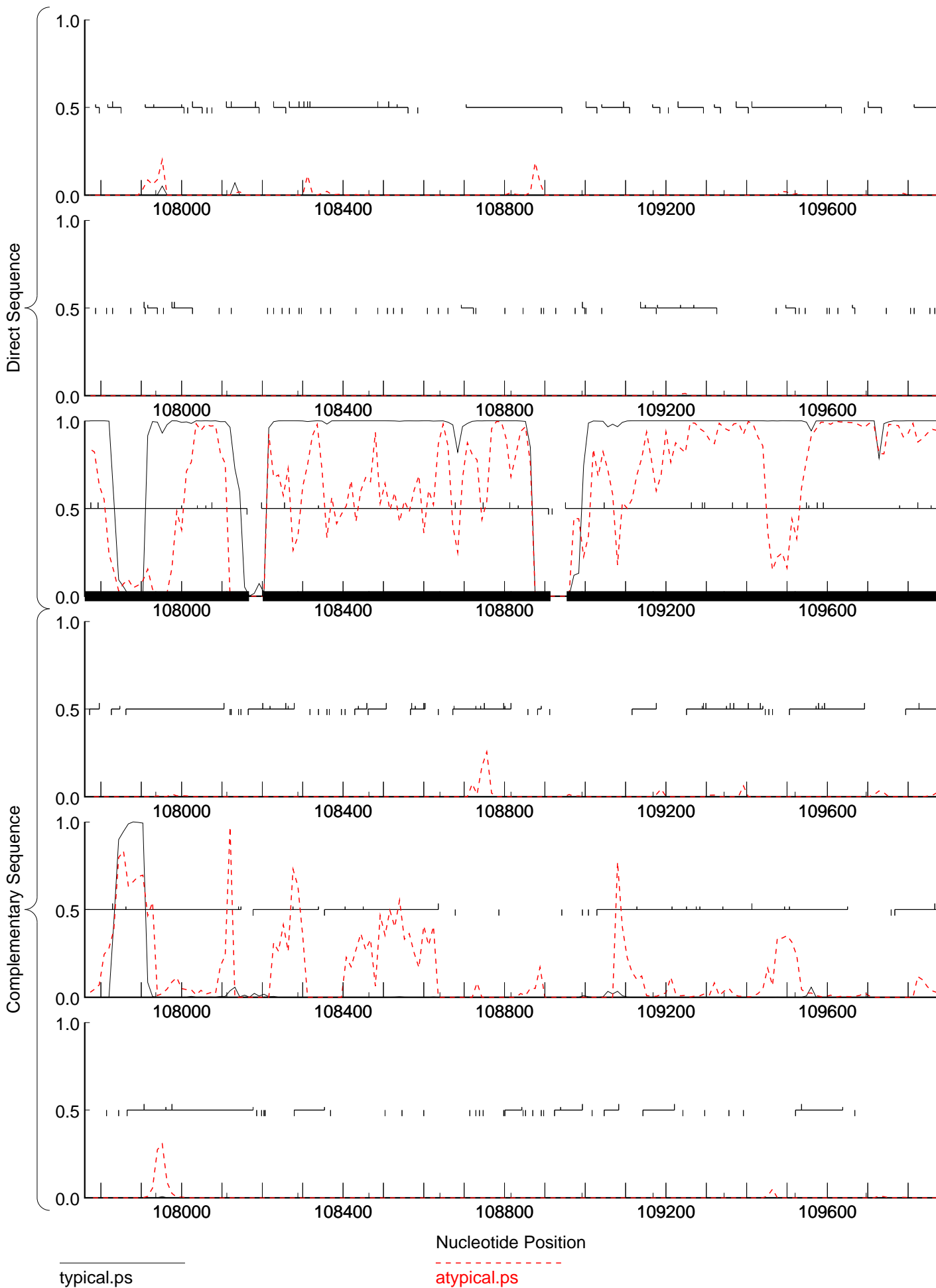
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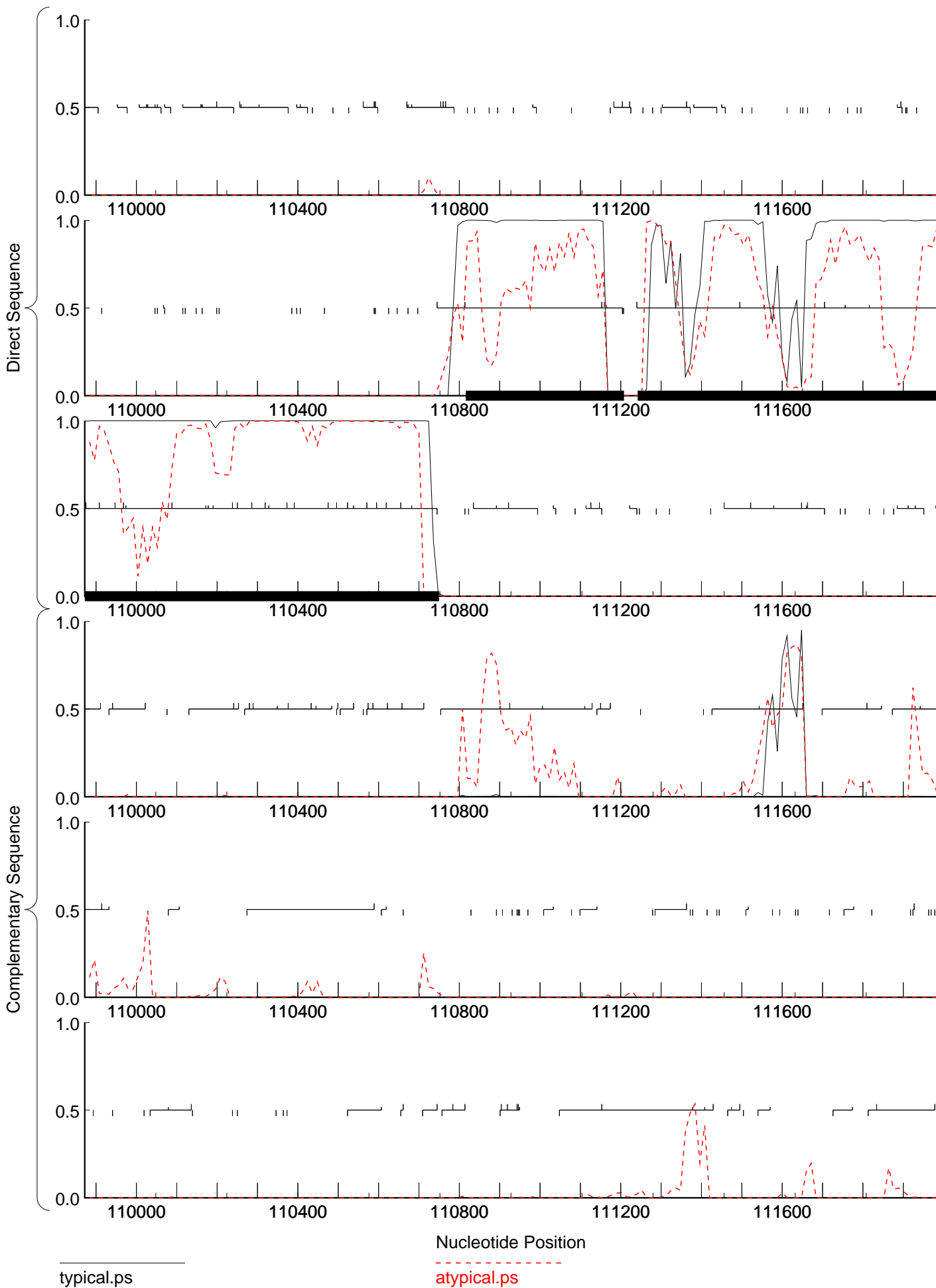


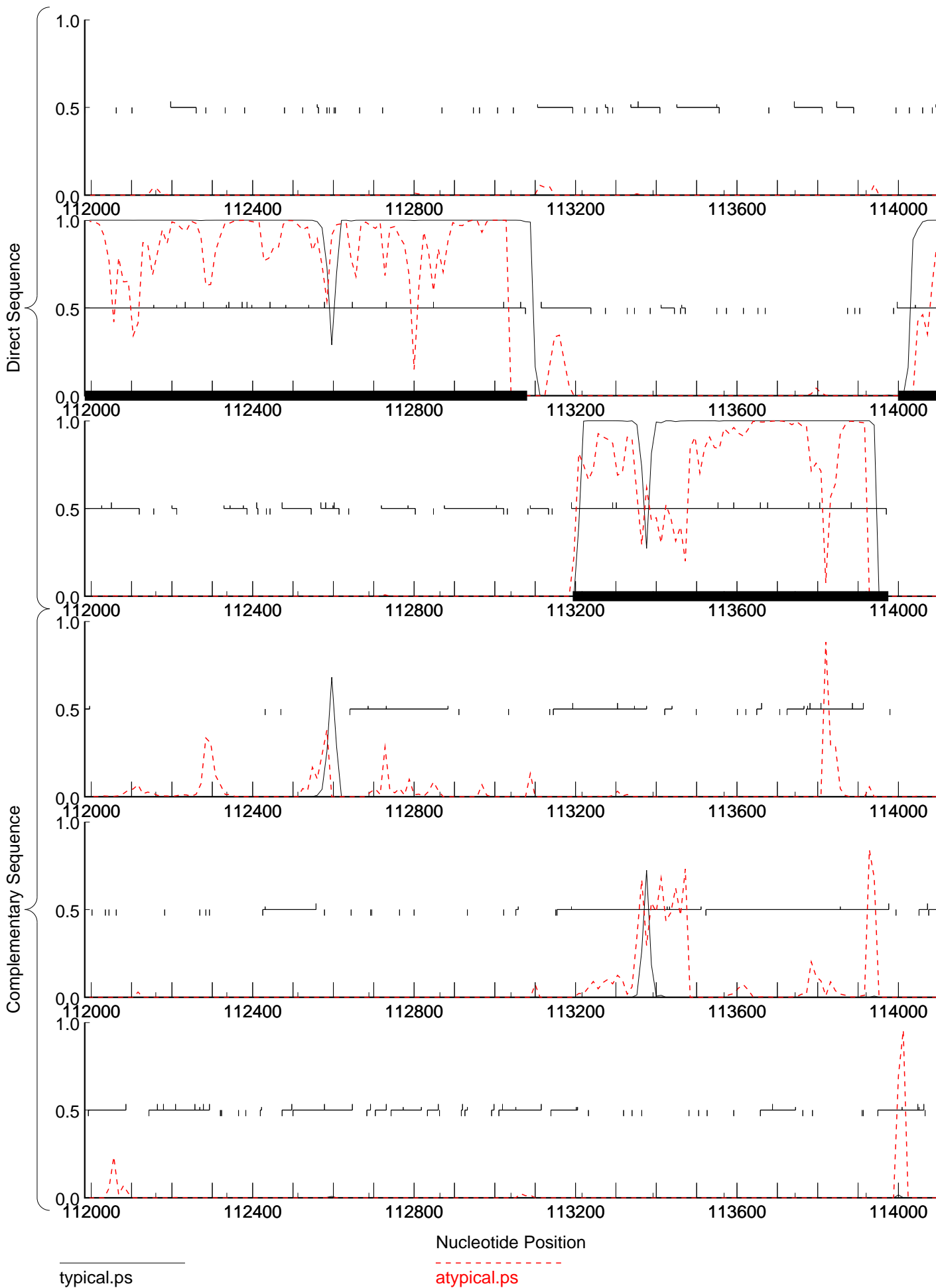
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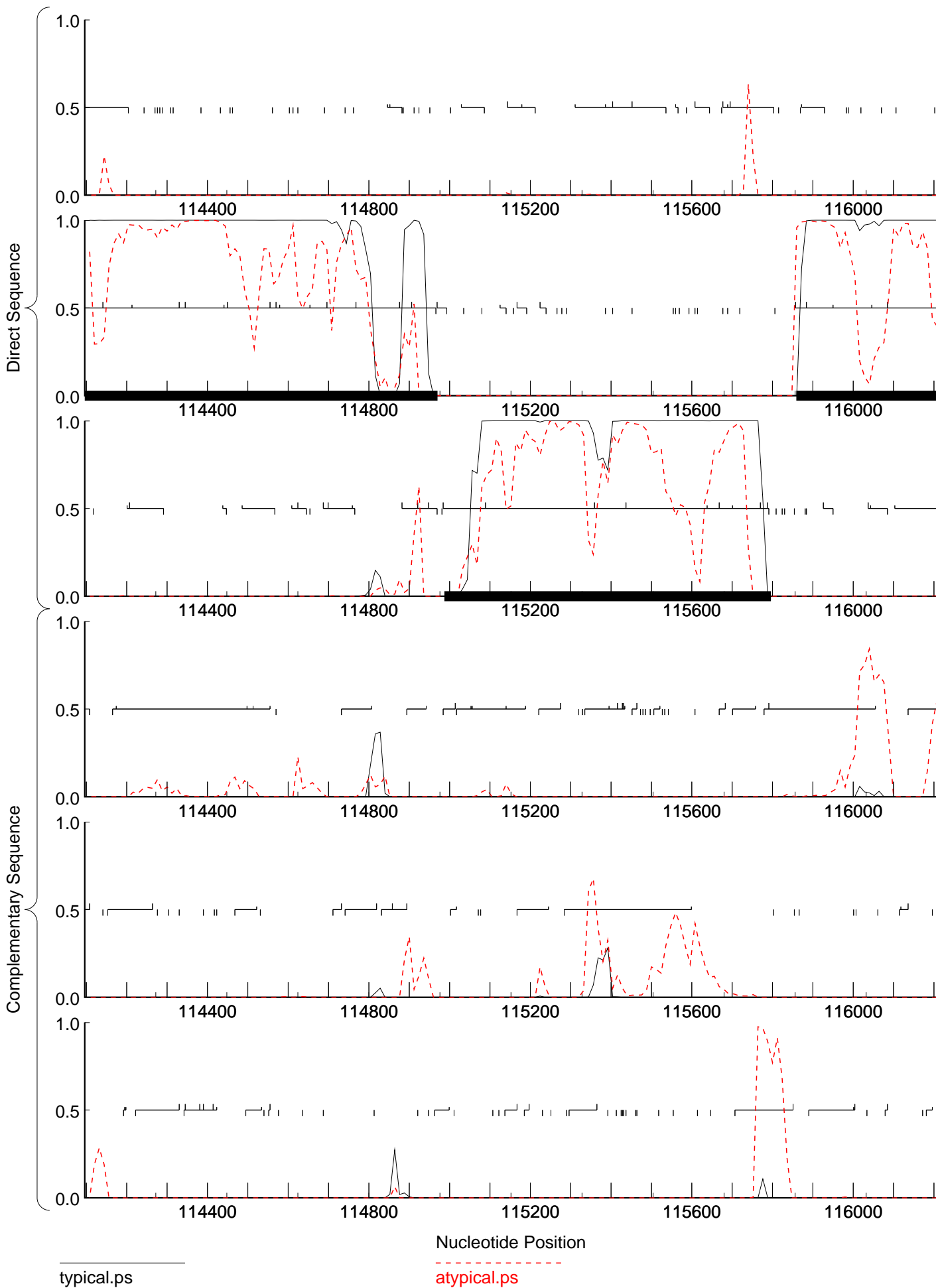
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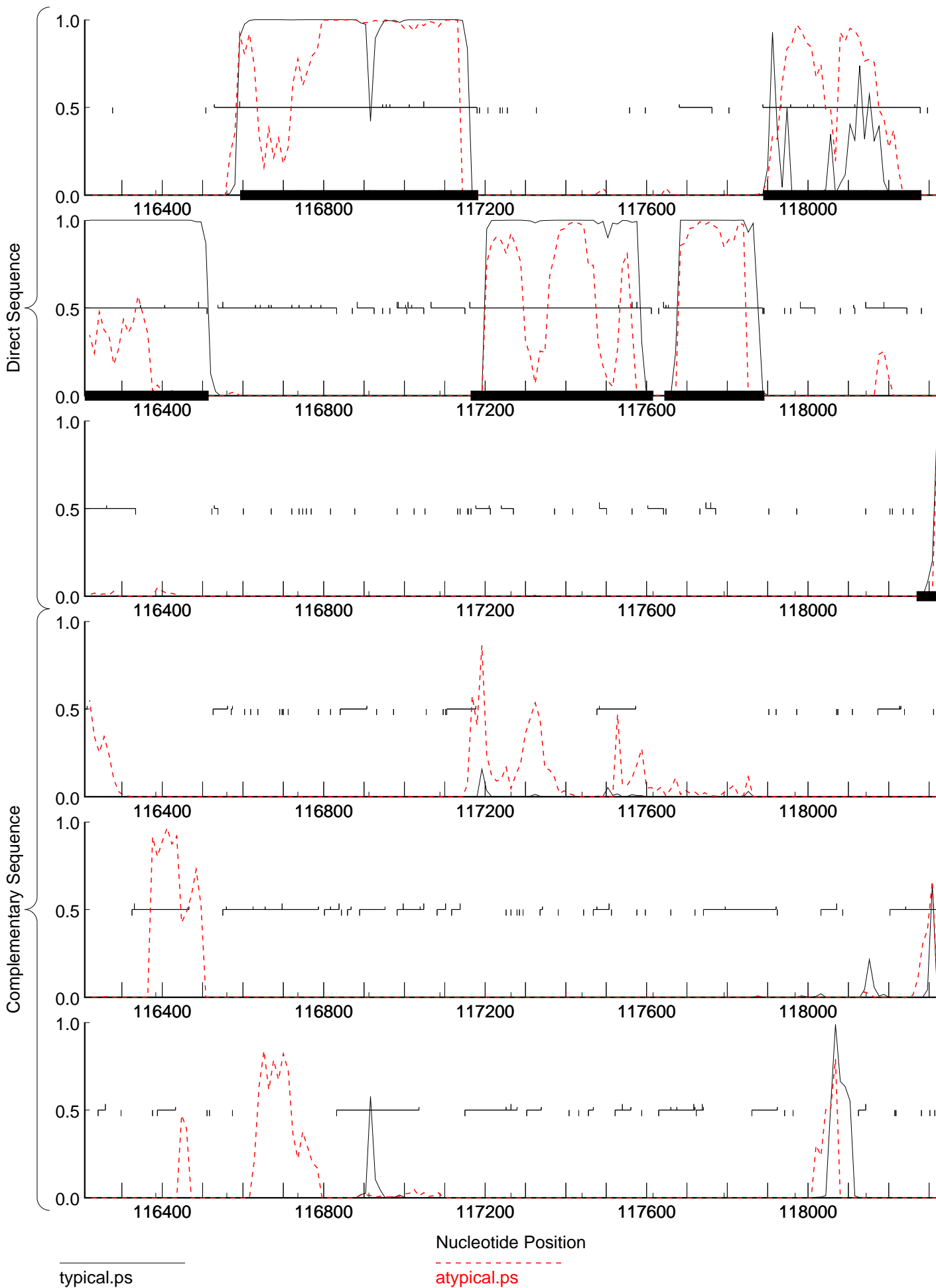




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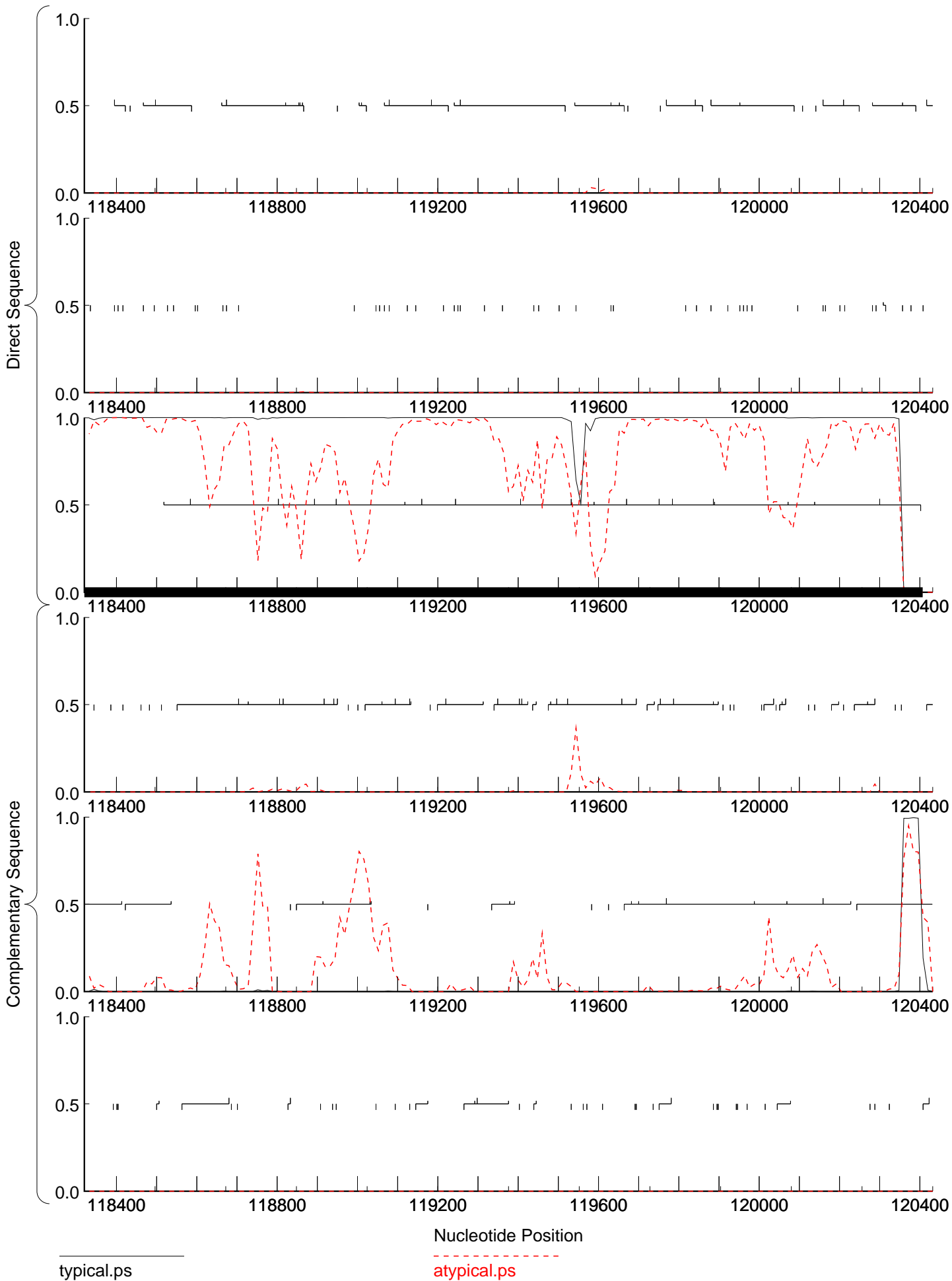
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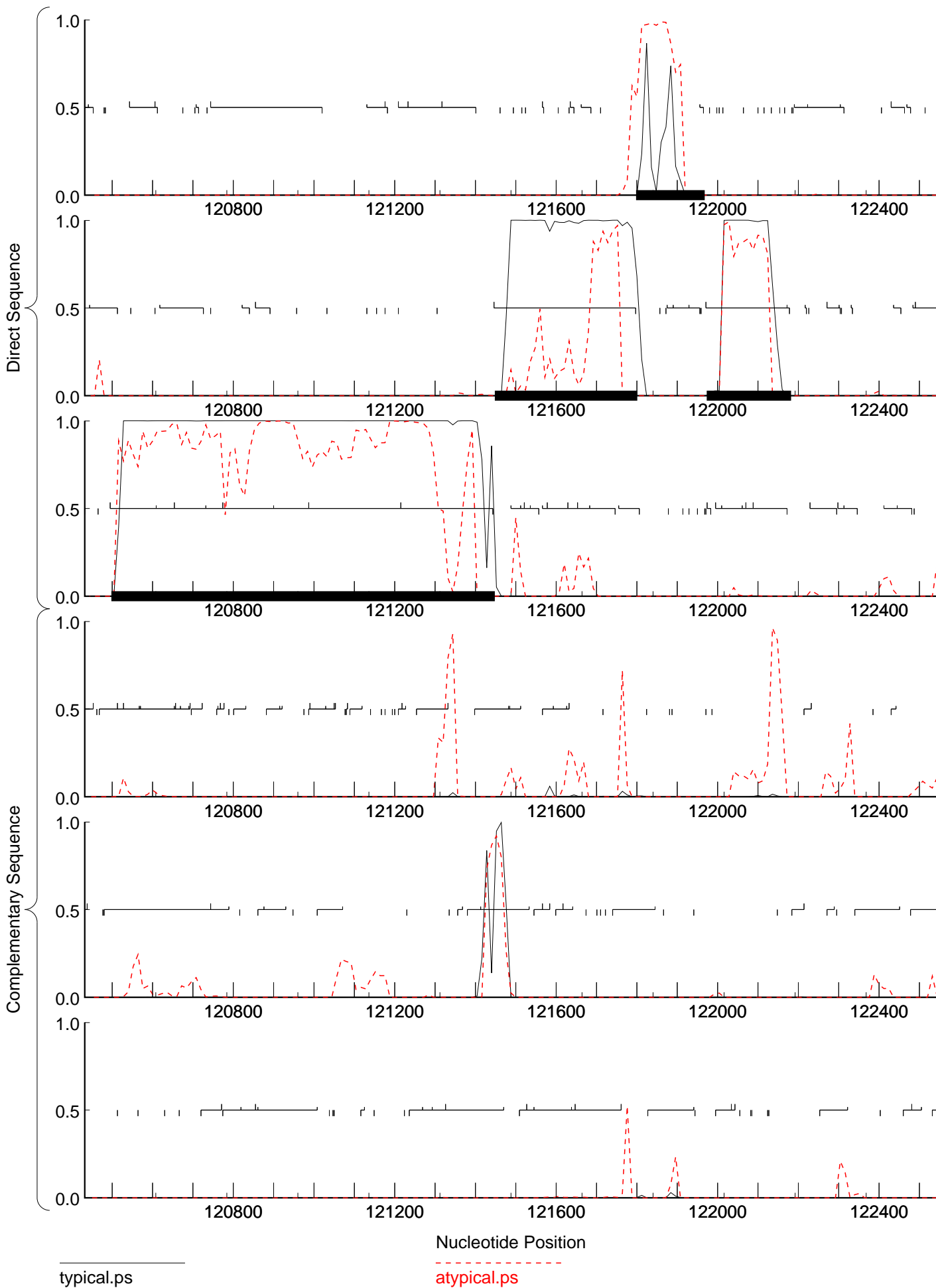


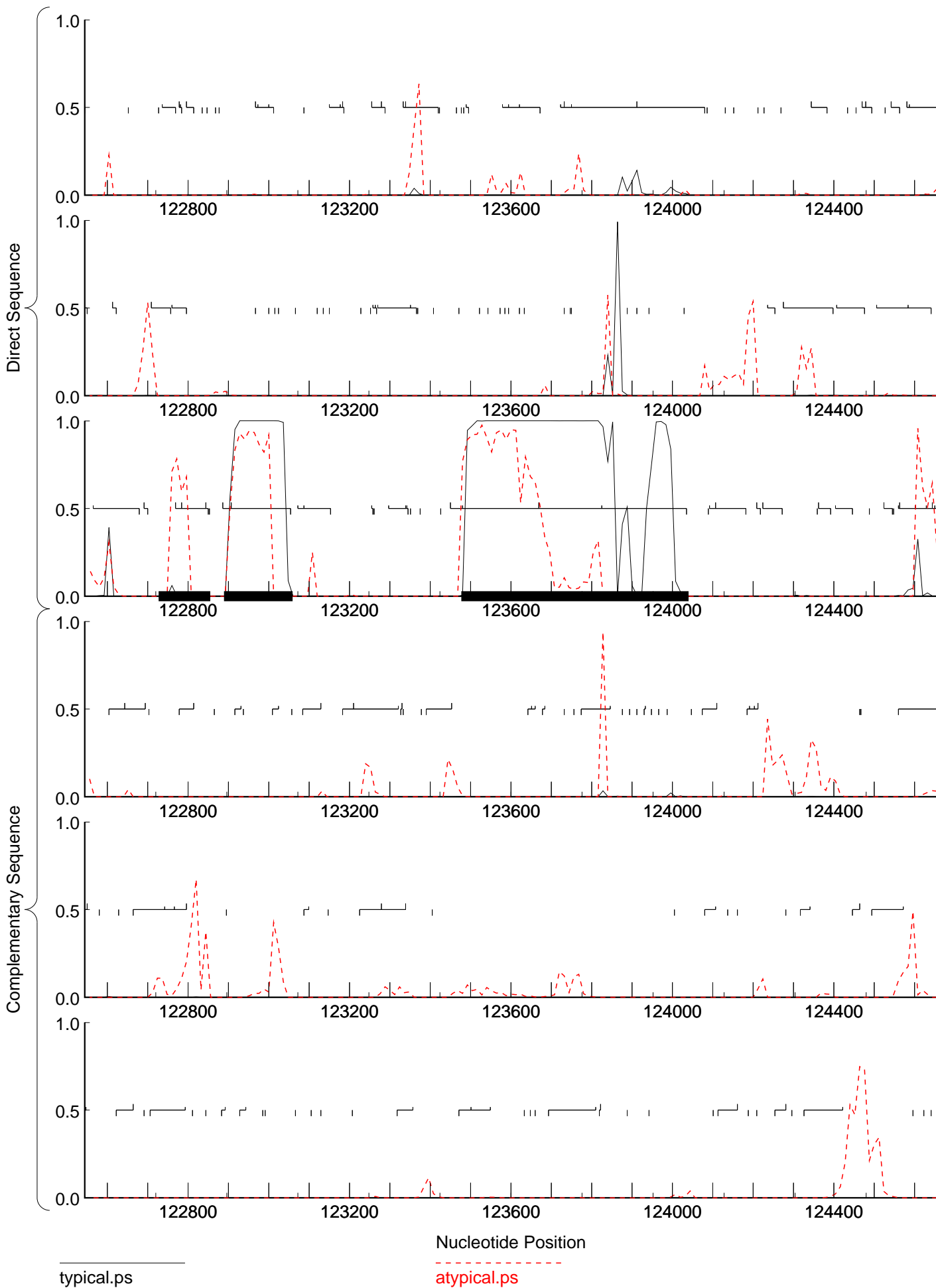


typical.ps

atypical.ps

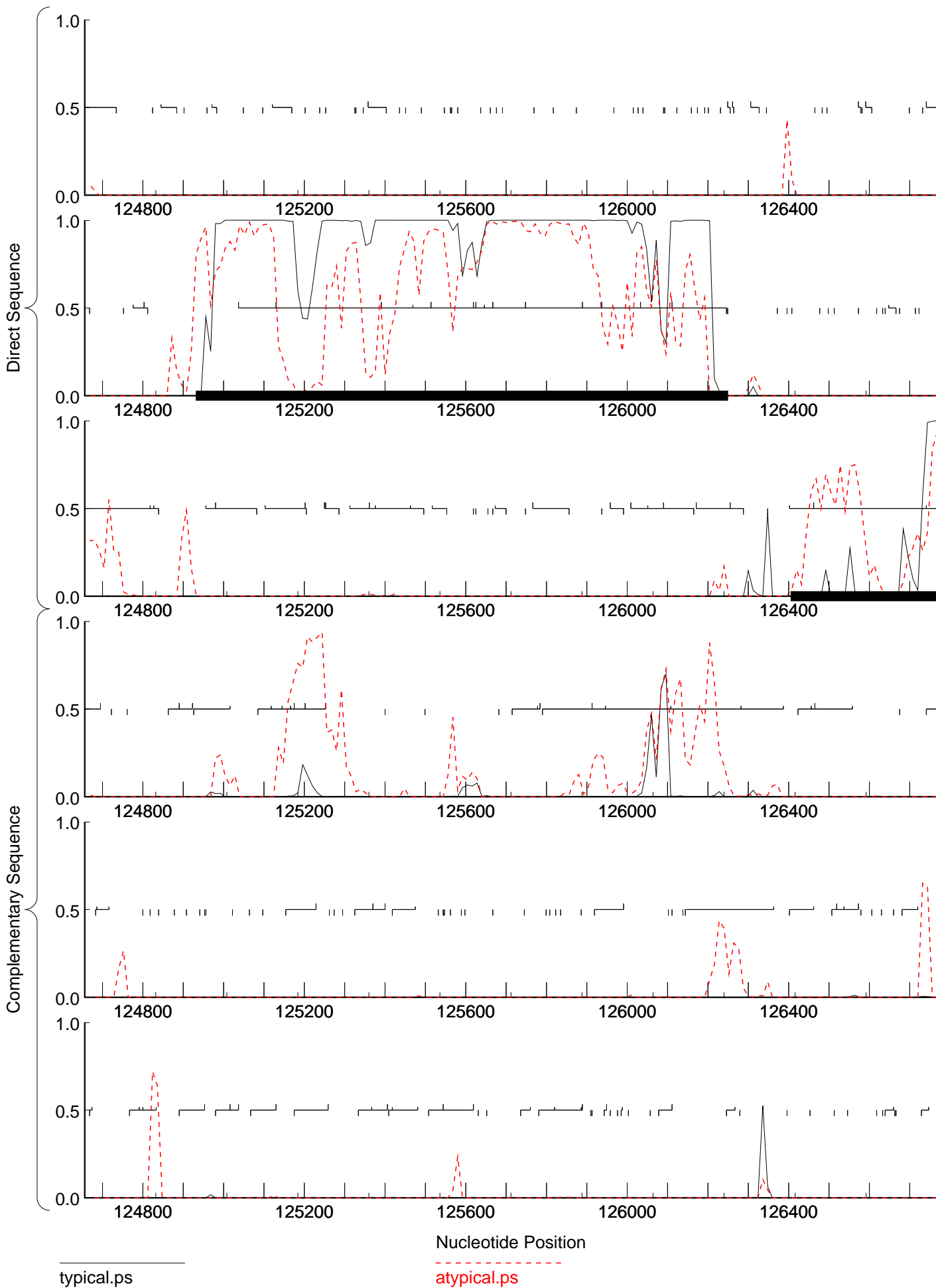


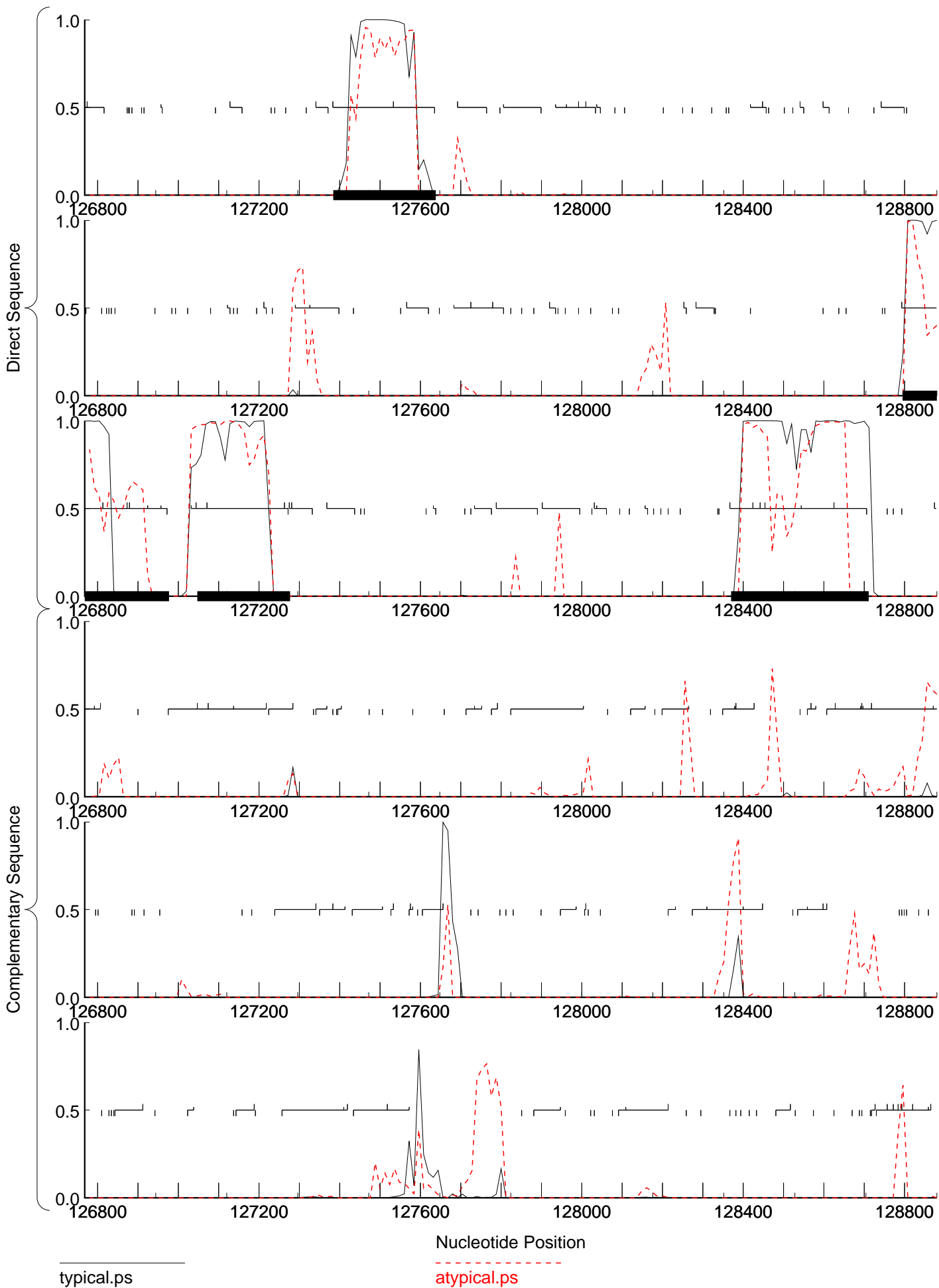


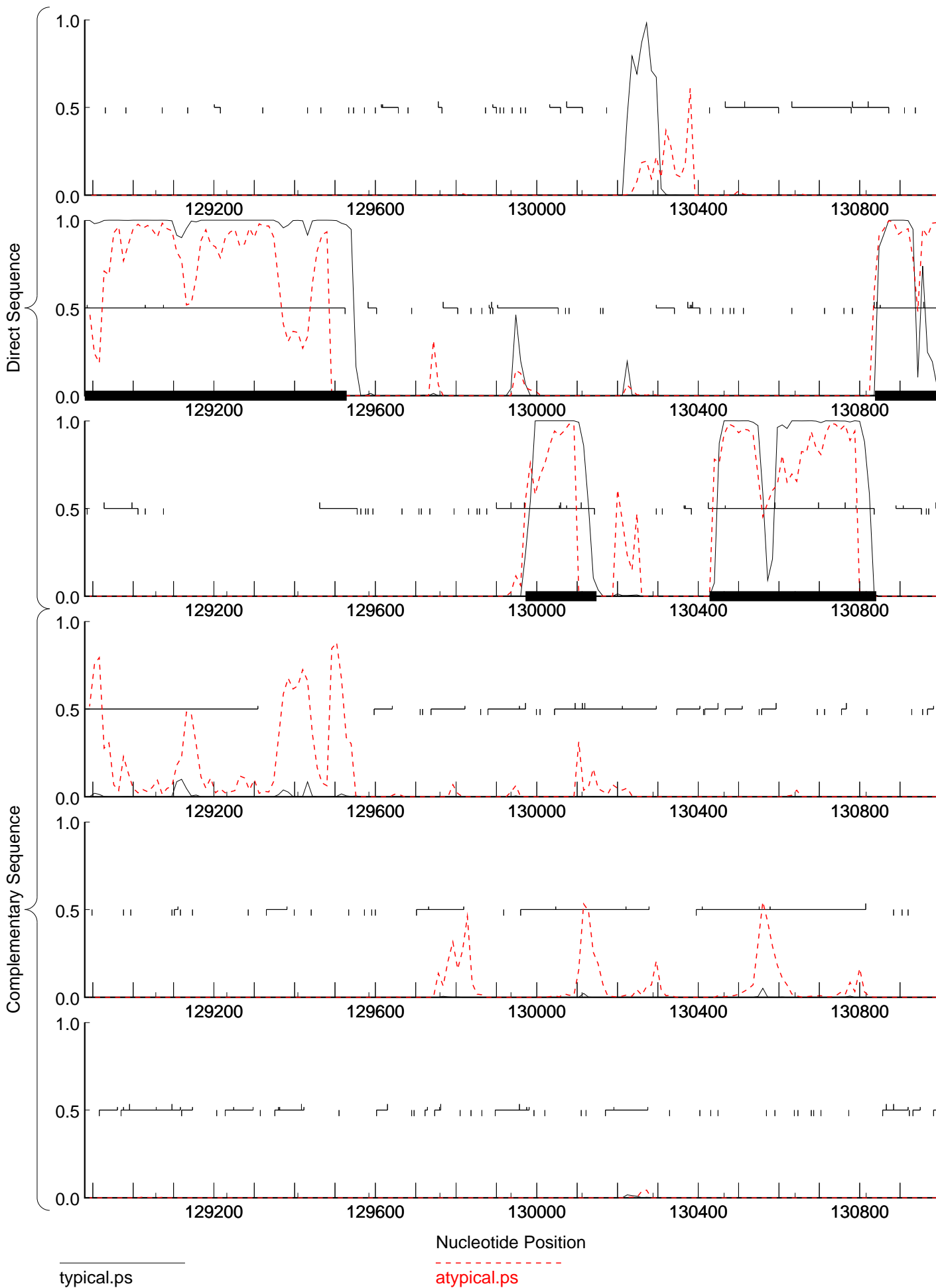


typical.ps

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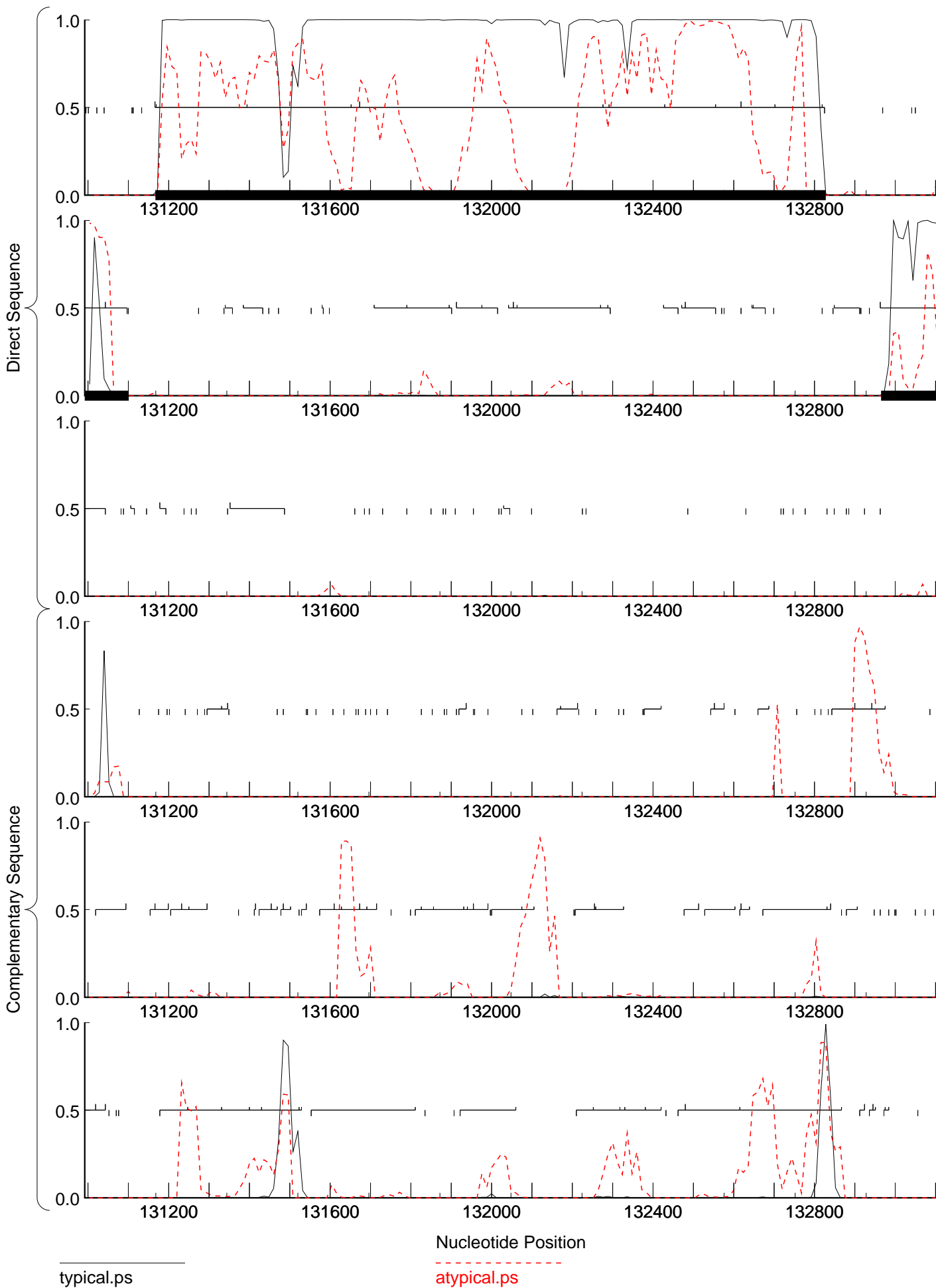




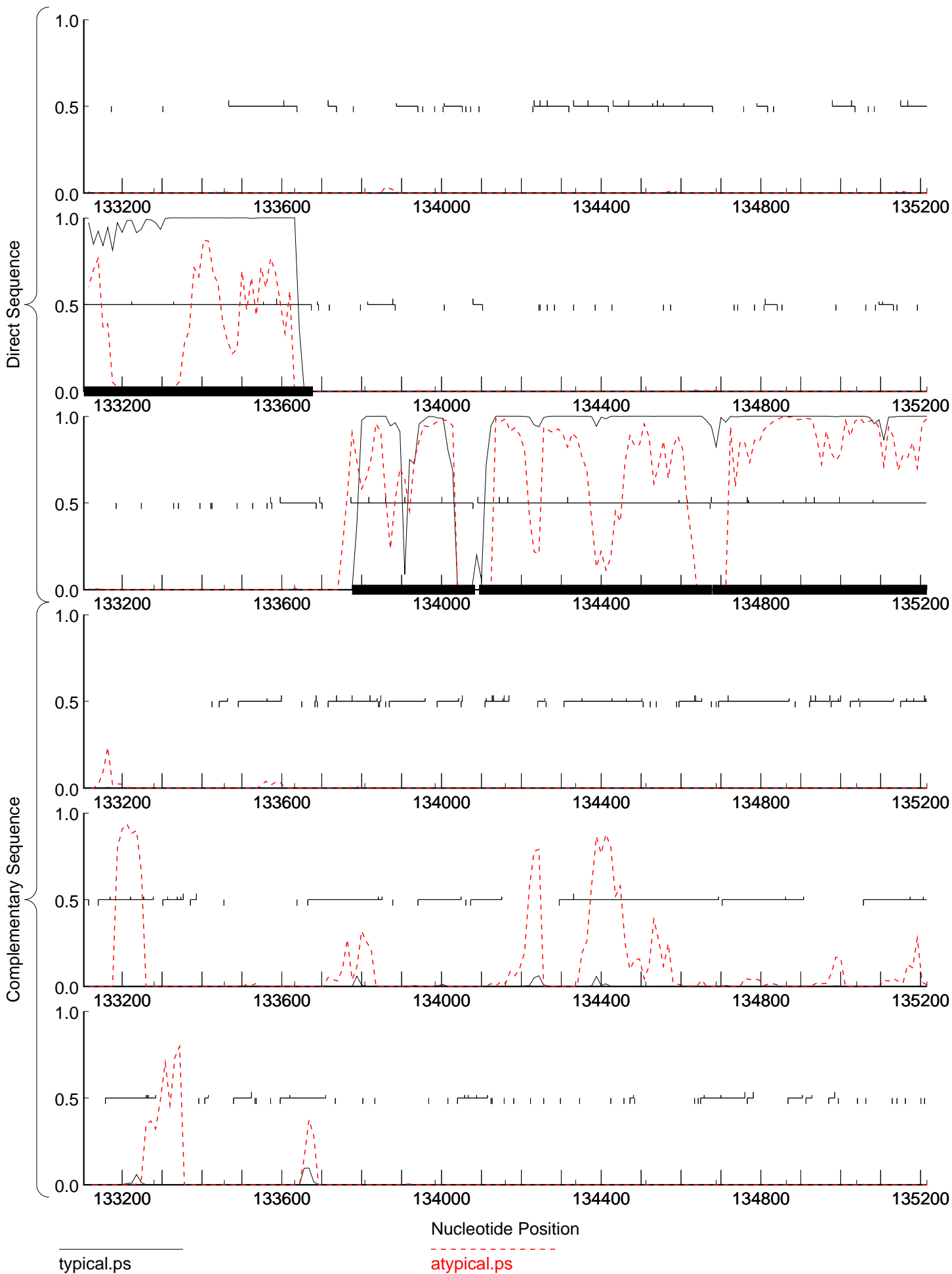


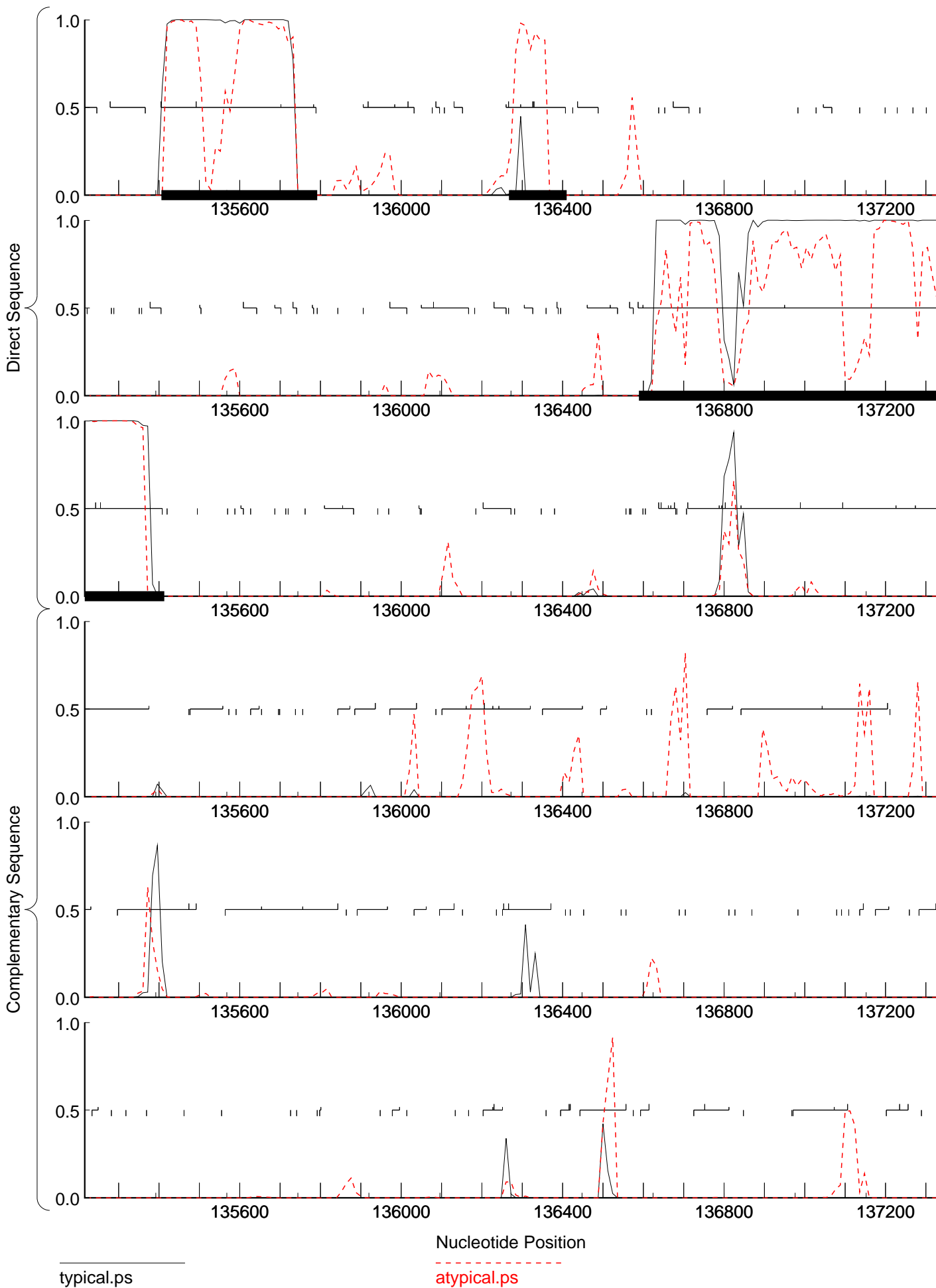
typical.ps

atypical.ps



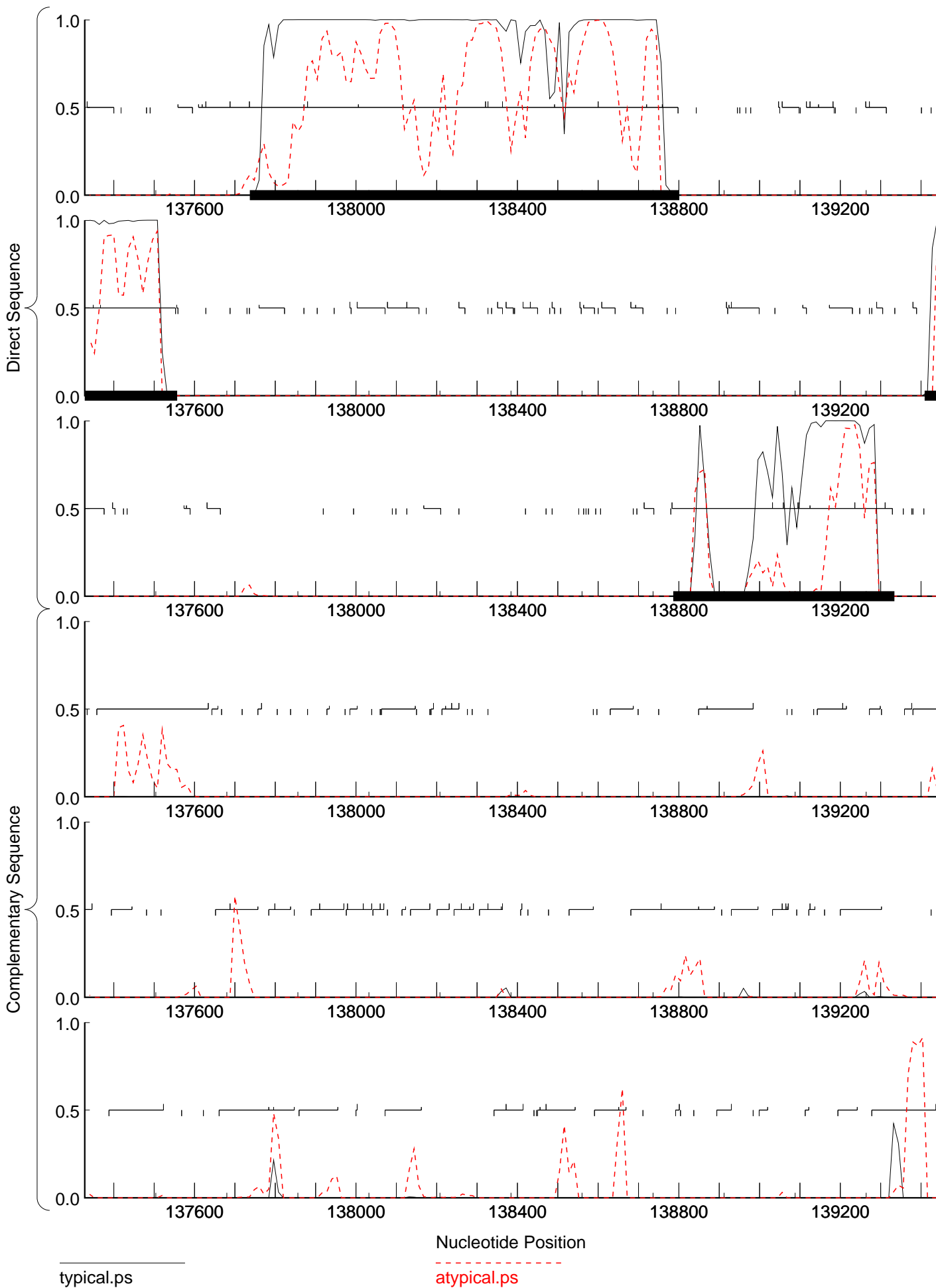






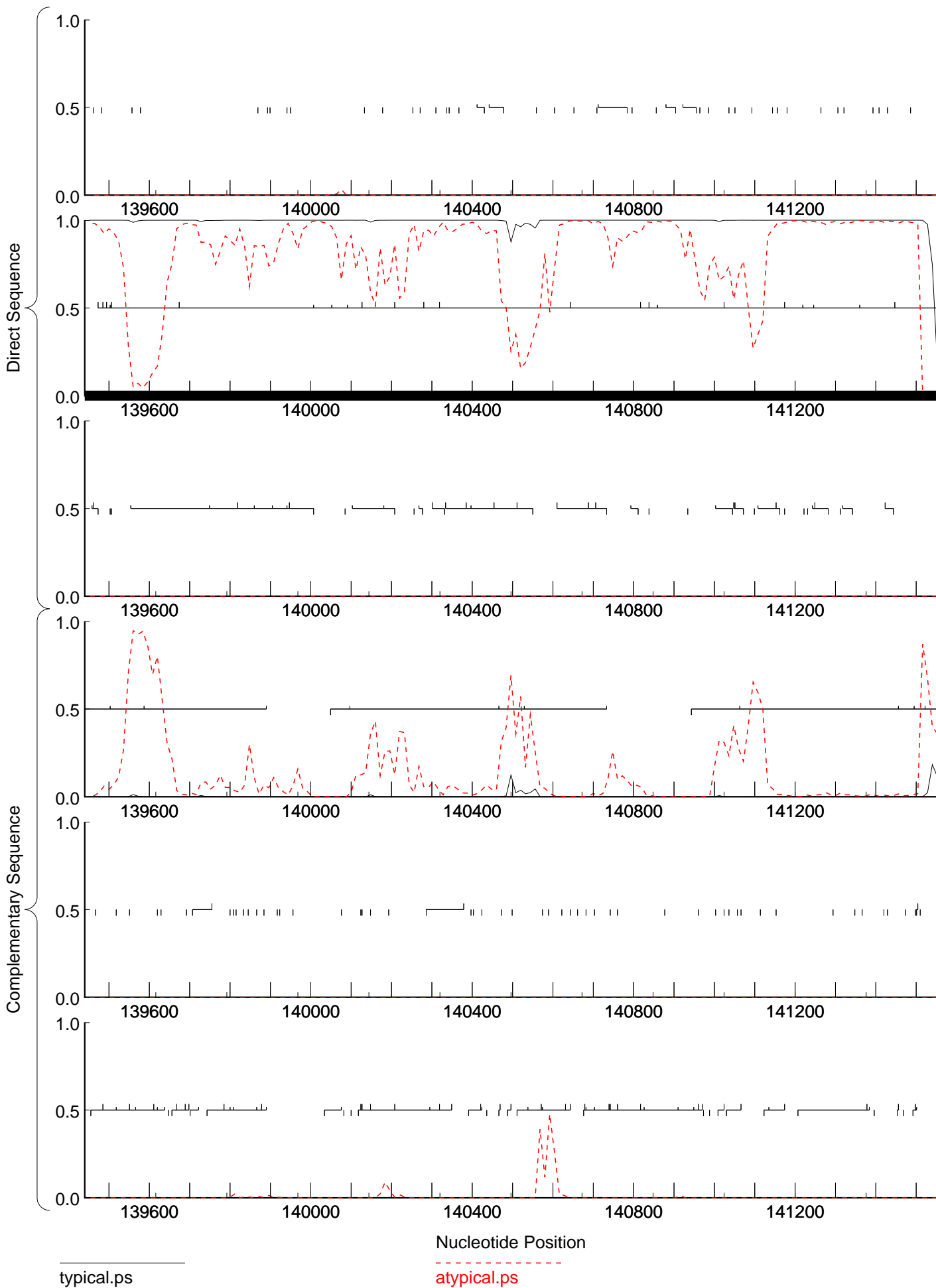
typical.ps

atypical.ps



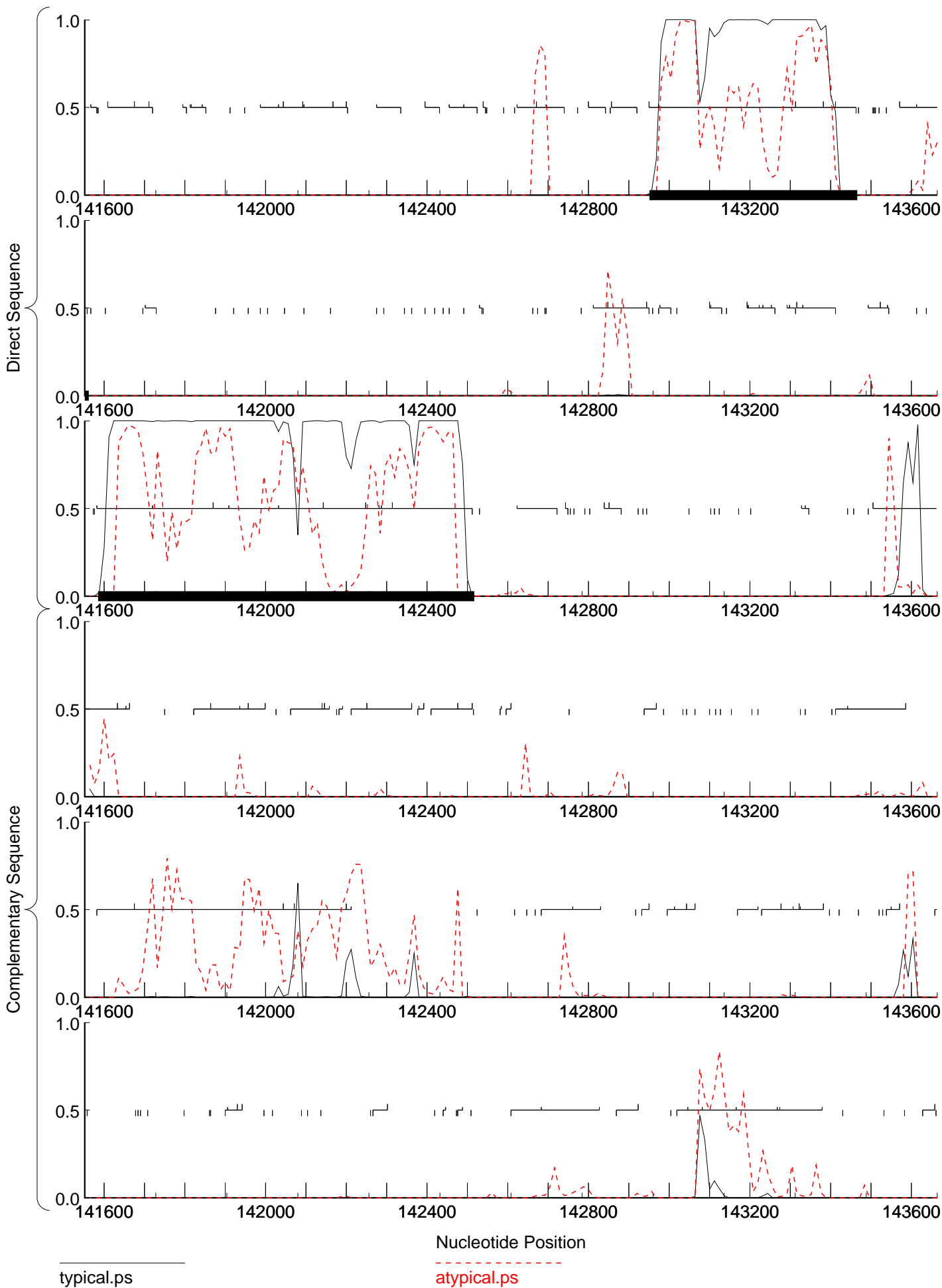
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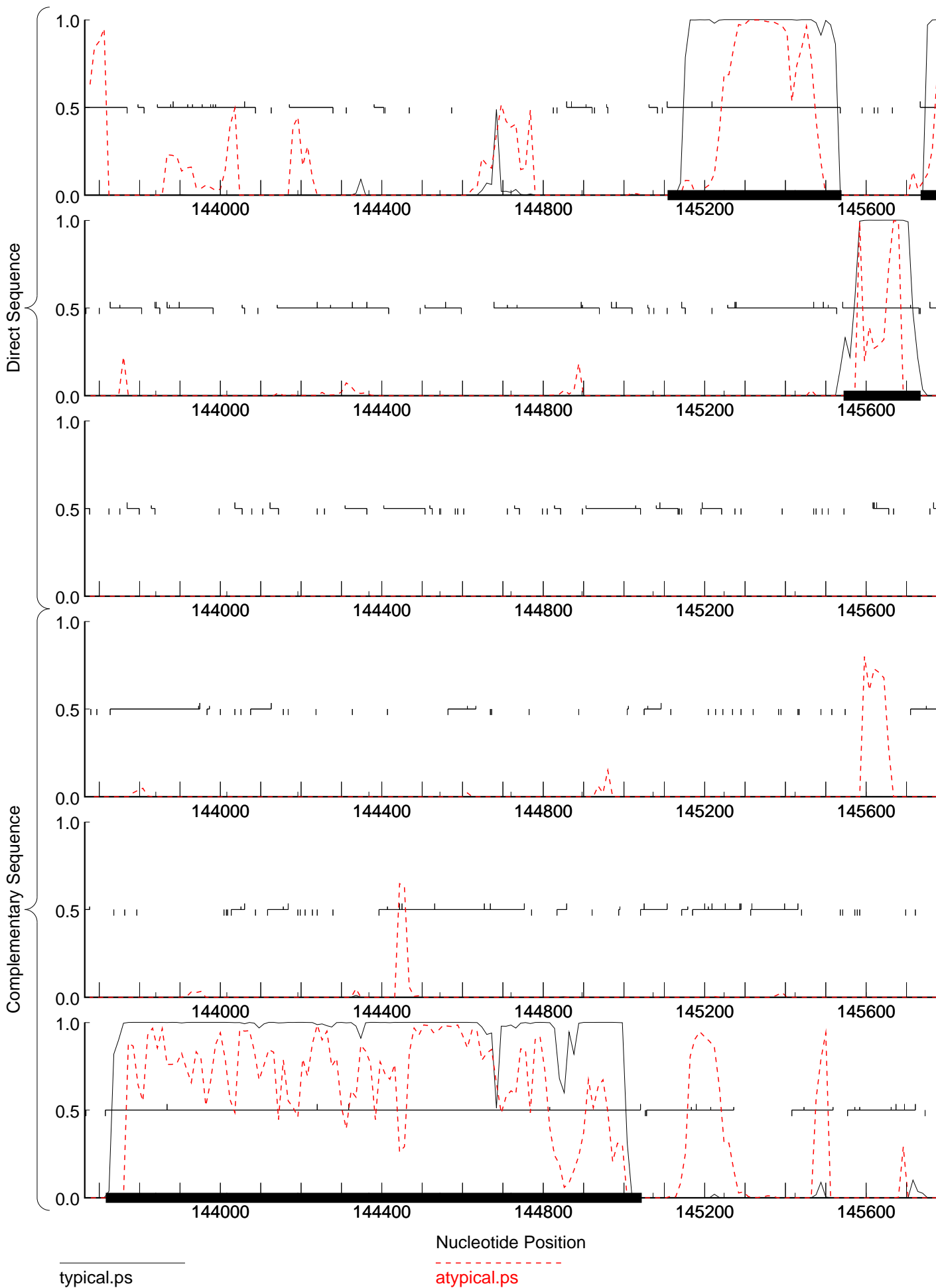
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typical.ps

atypical.ps

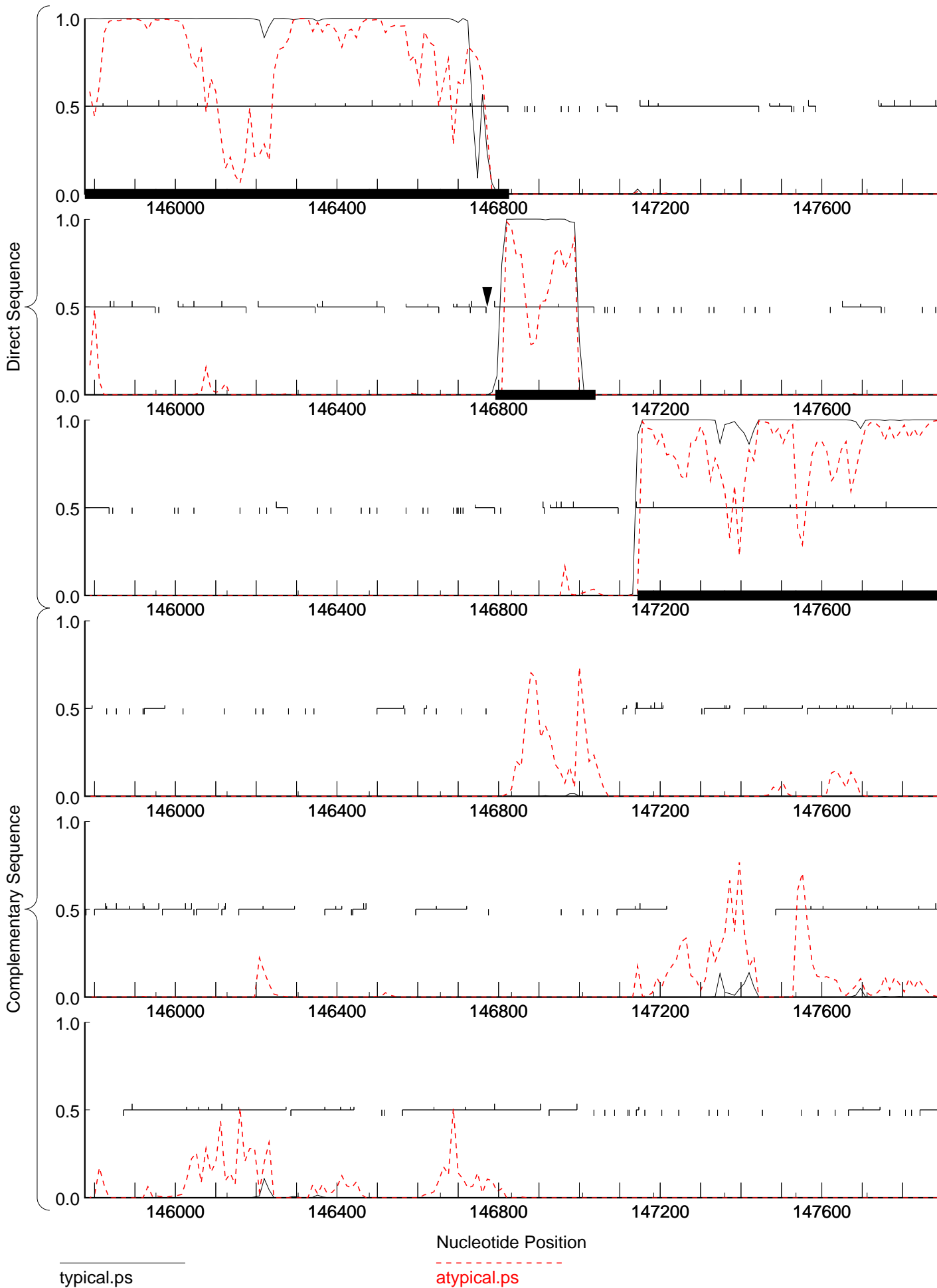




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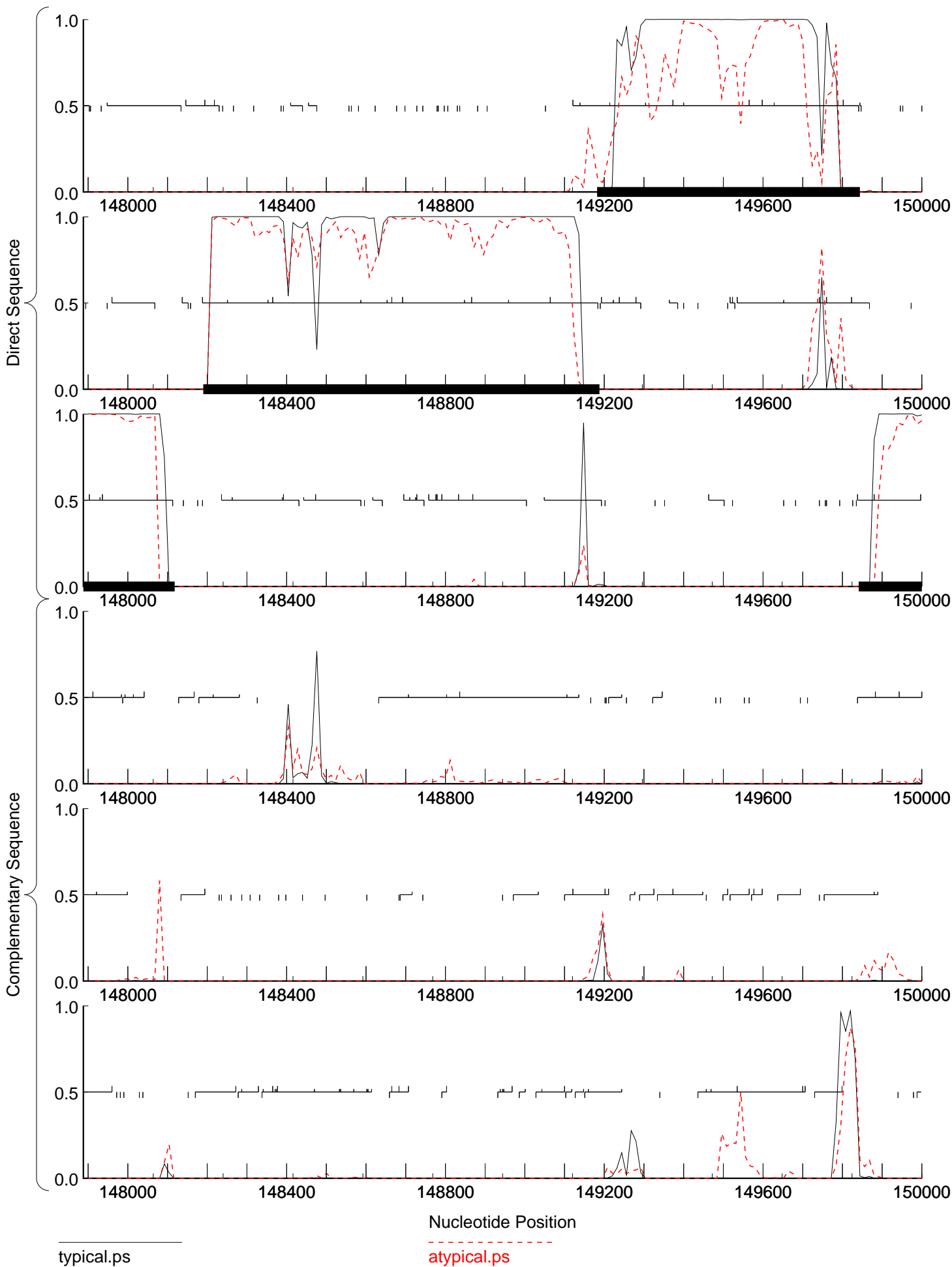
Nucleotide Position

atypical.ps

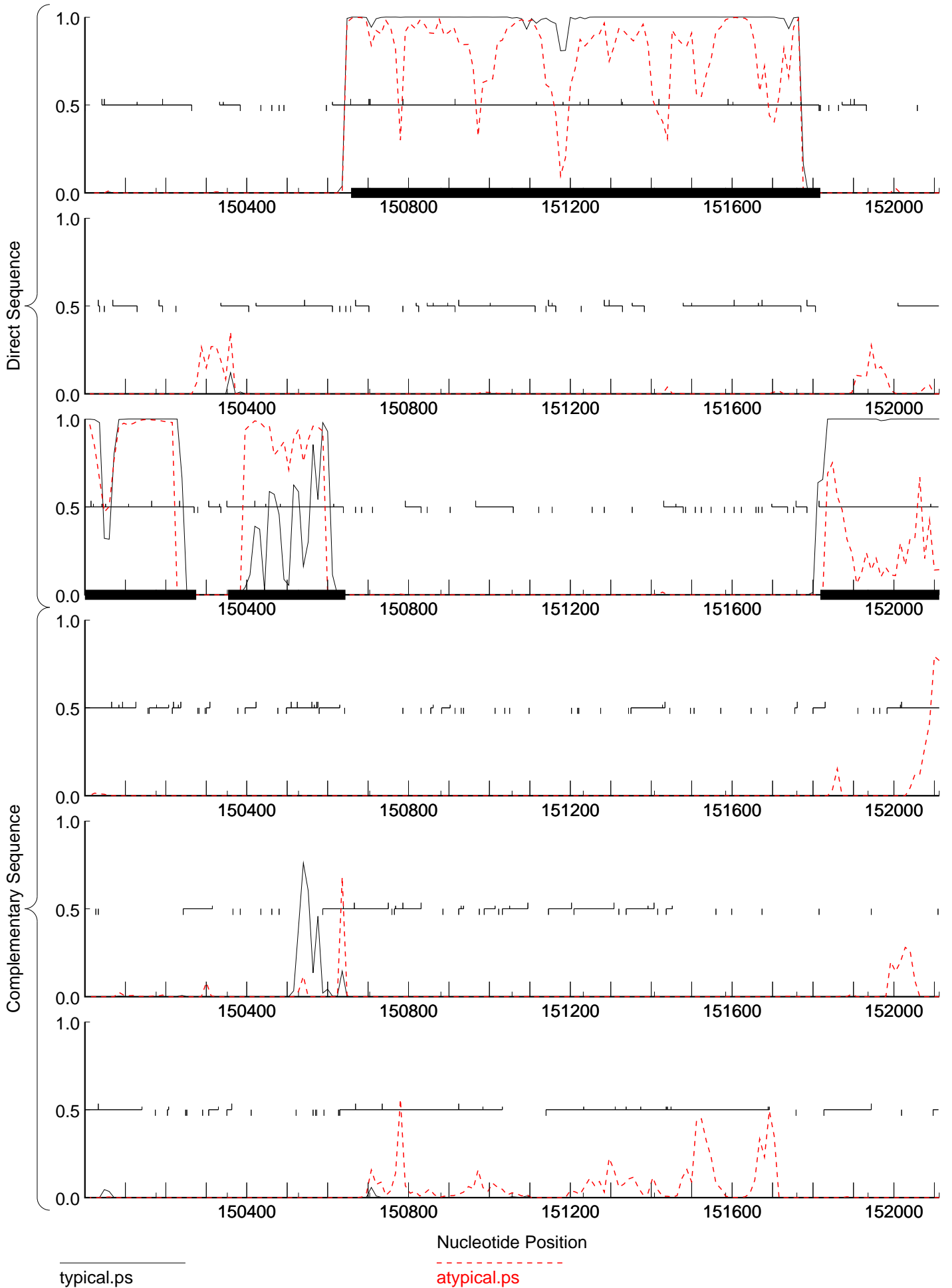


typical.ps

atypical.ps

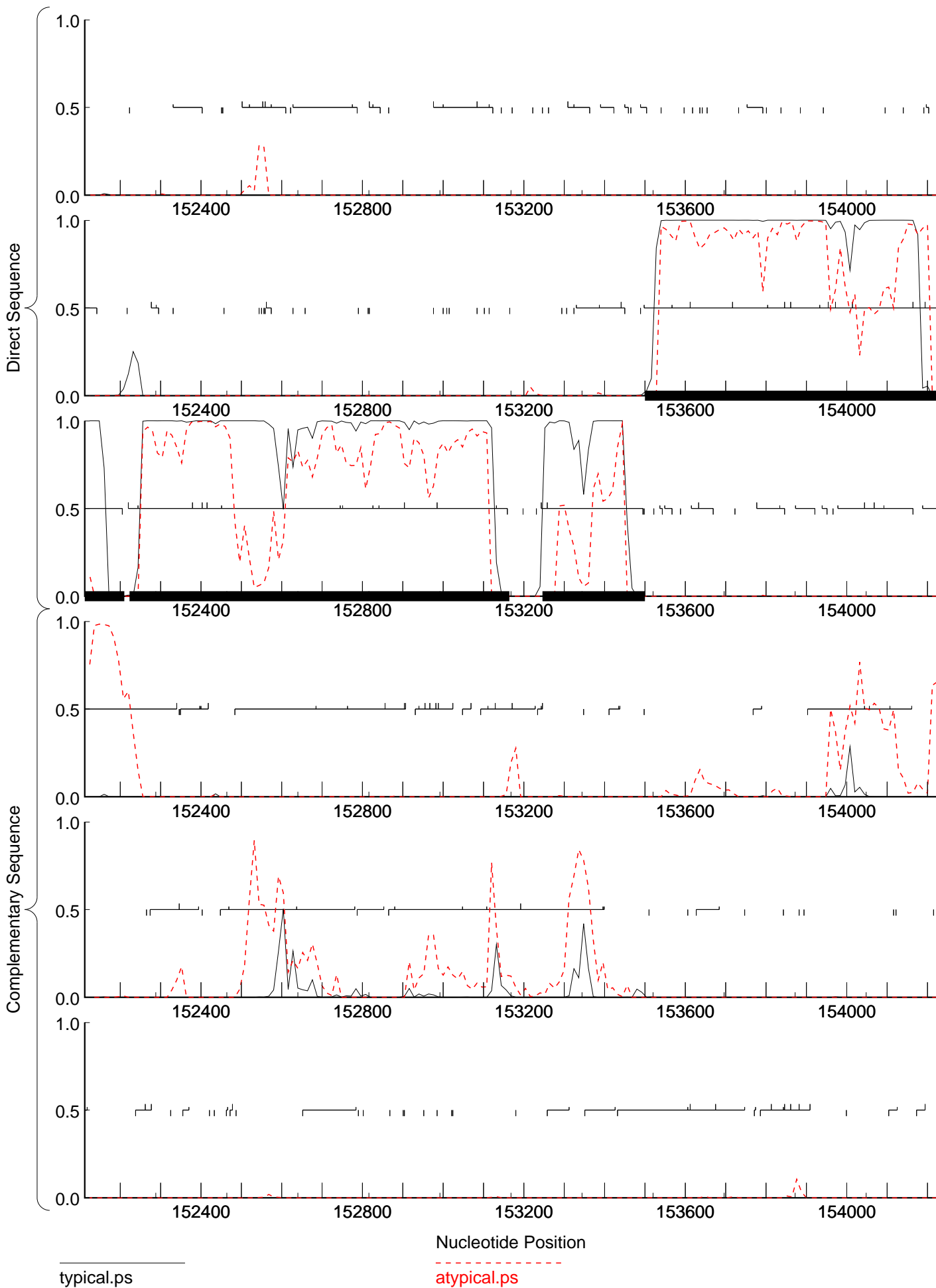


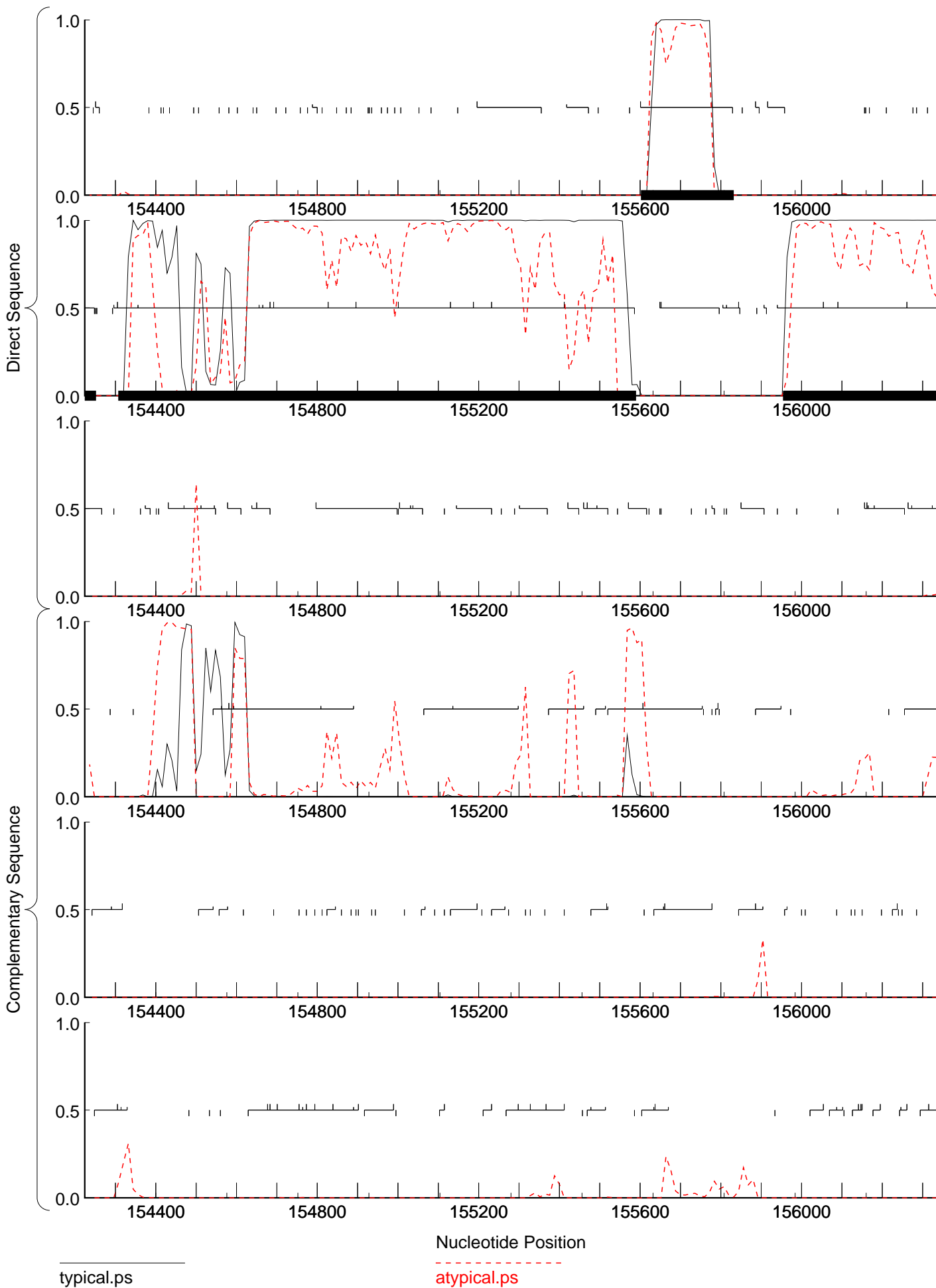


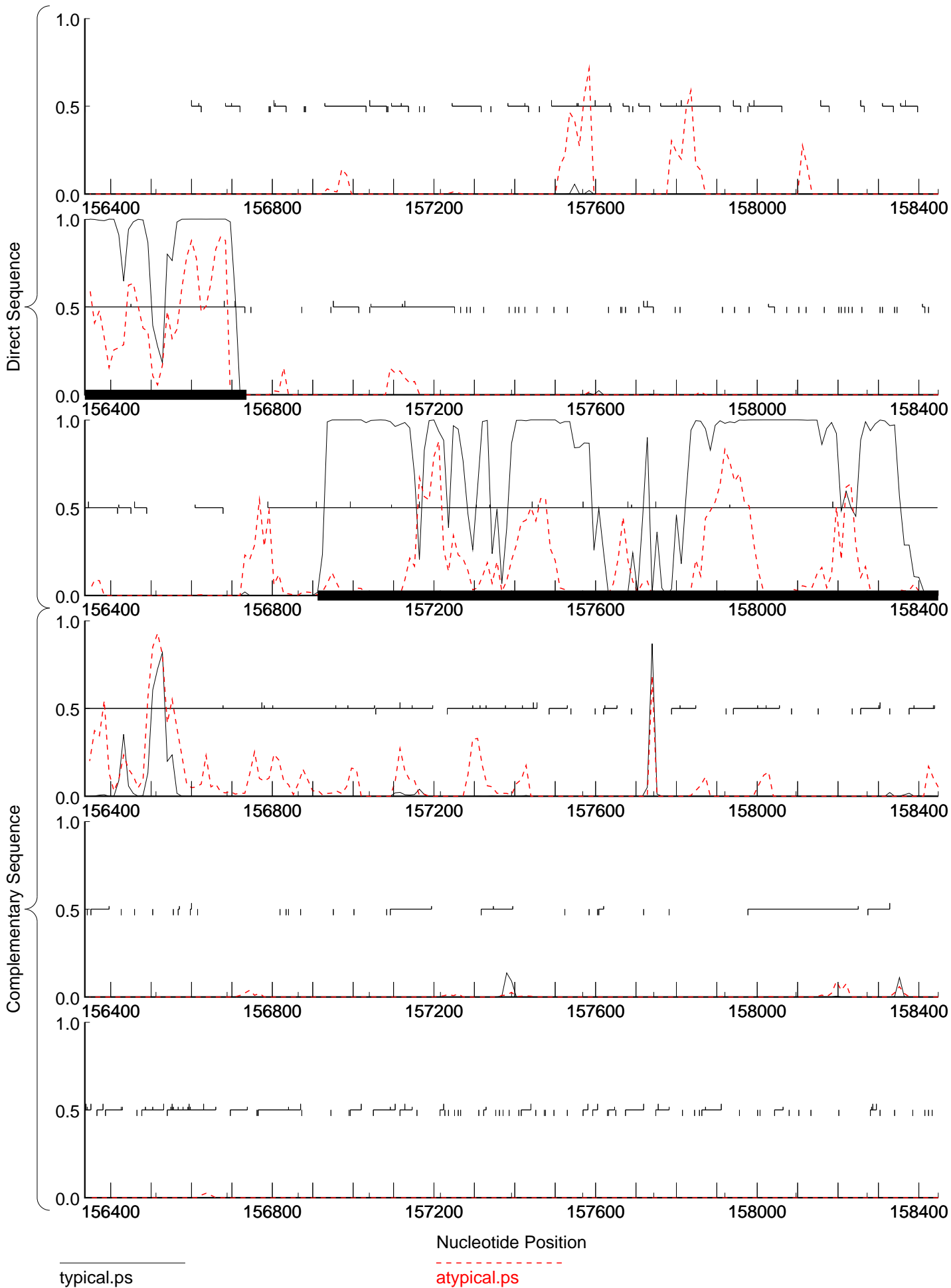


typical.ps

atypical.ps

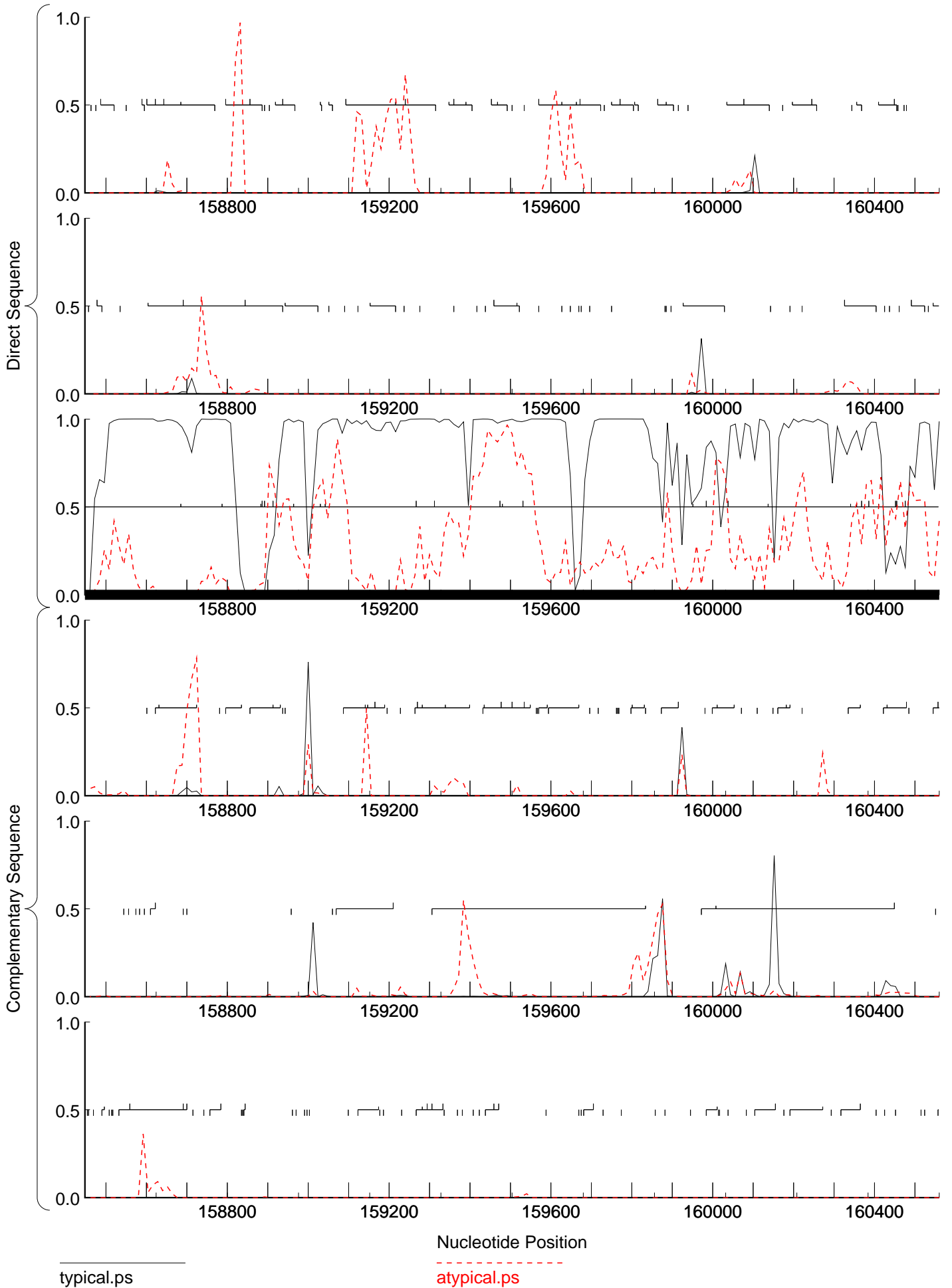






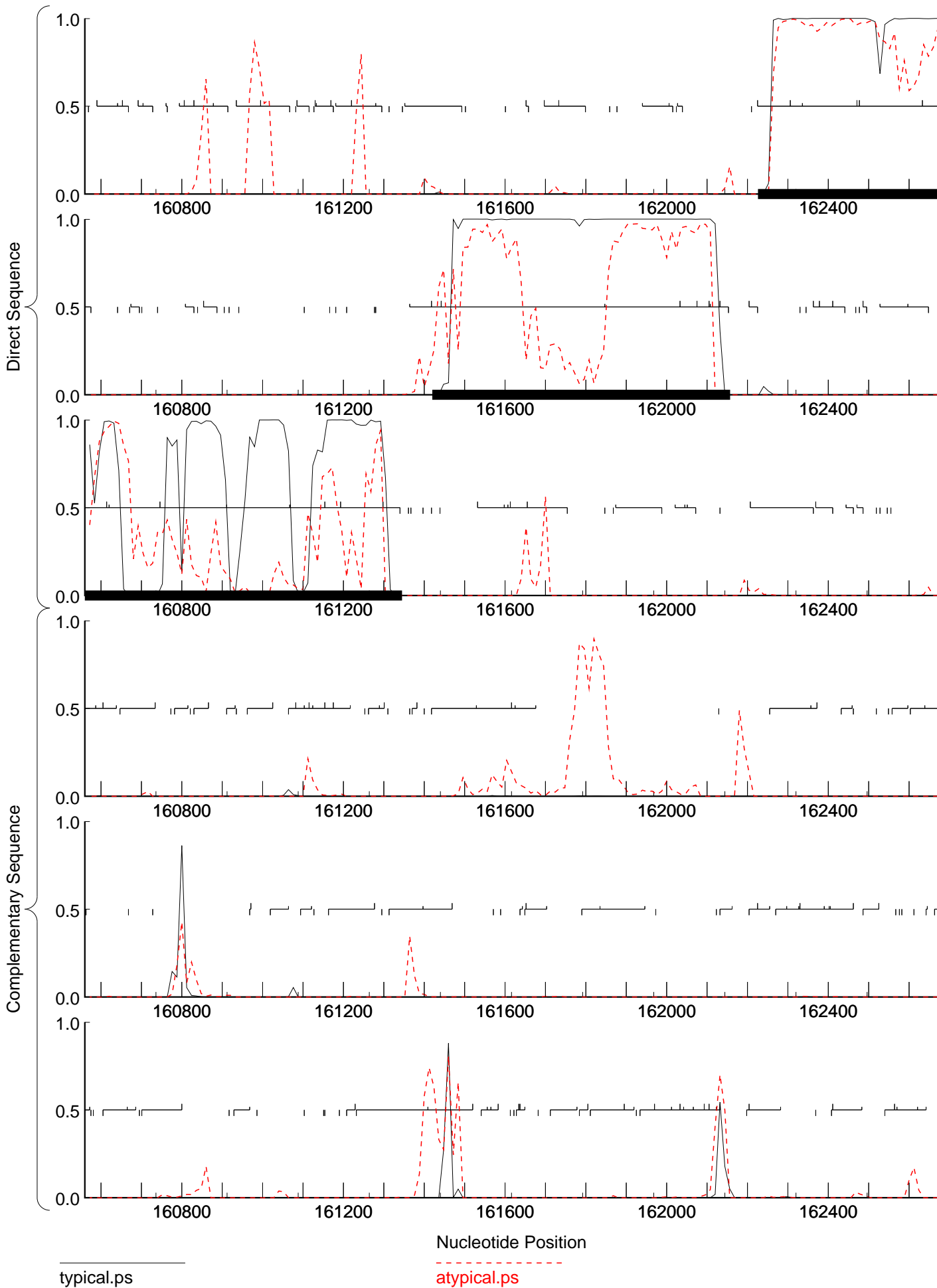
typical.ps

atypical.ps



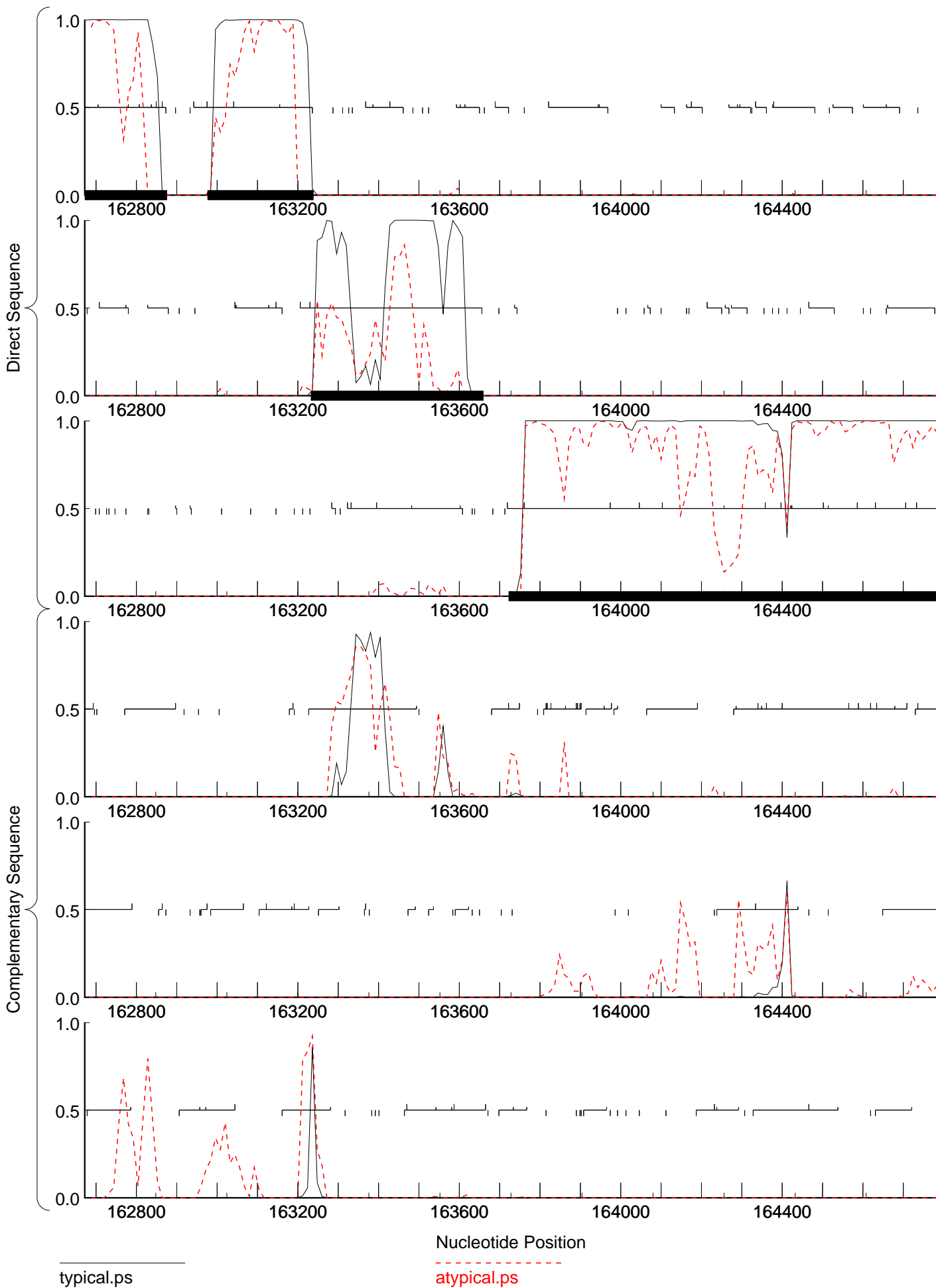
typical.ps

atypical.ps



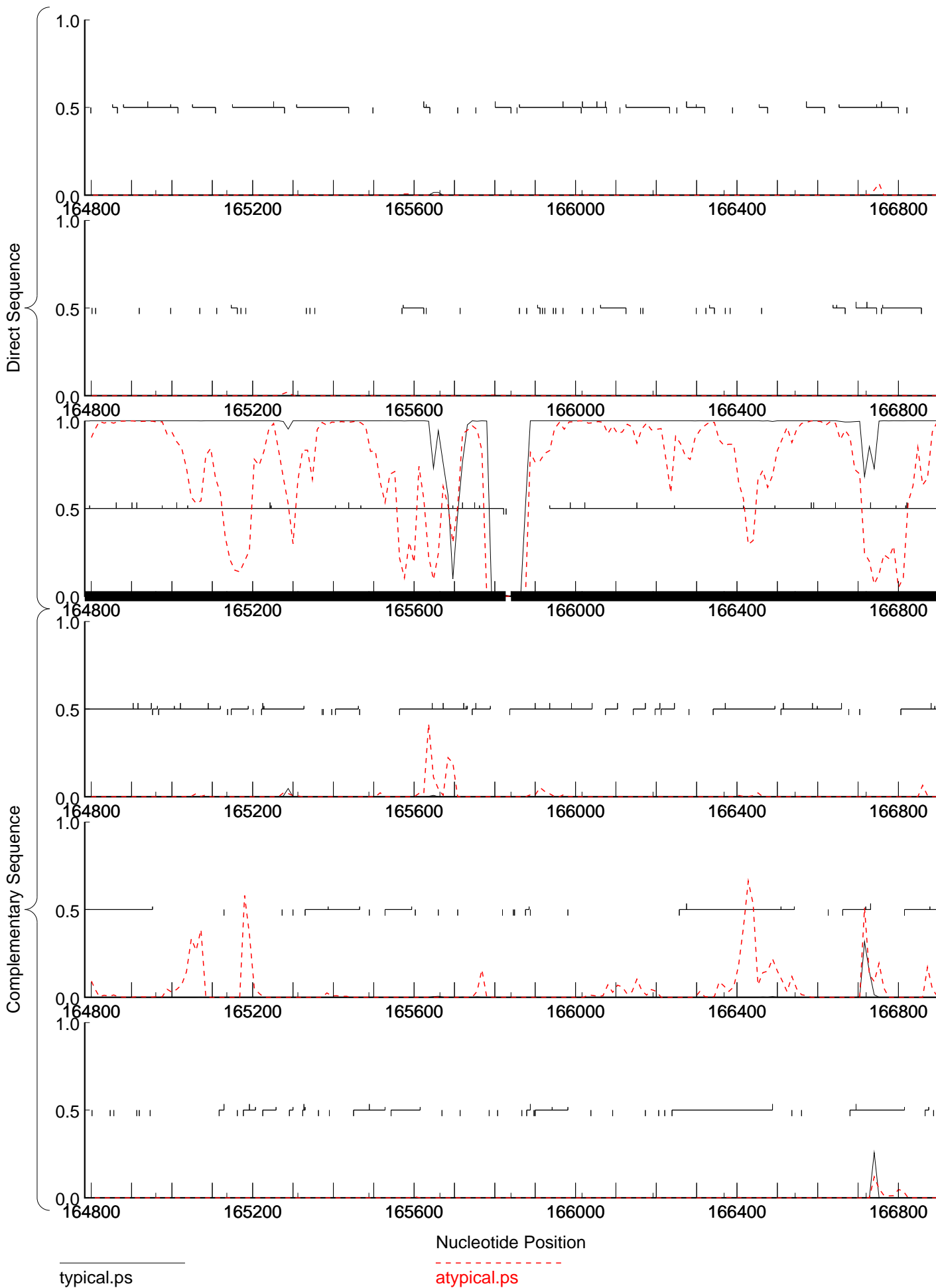
typical.ps

atypical.ps



typical.ps

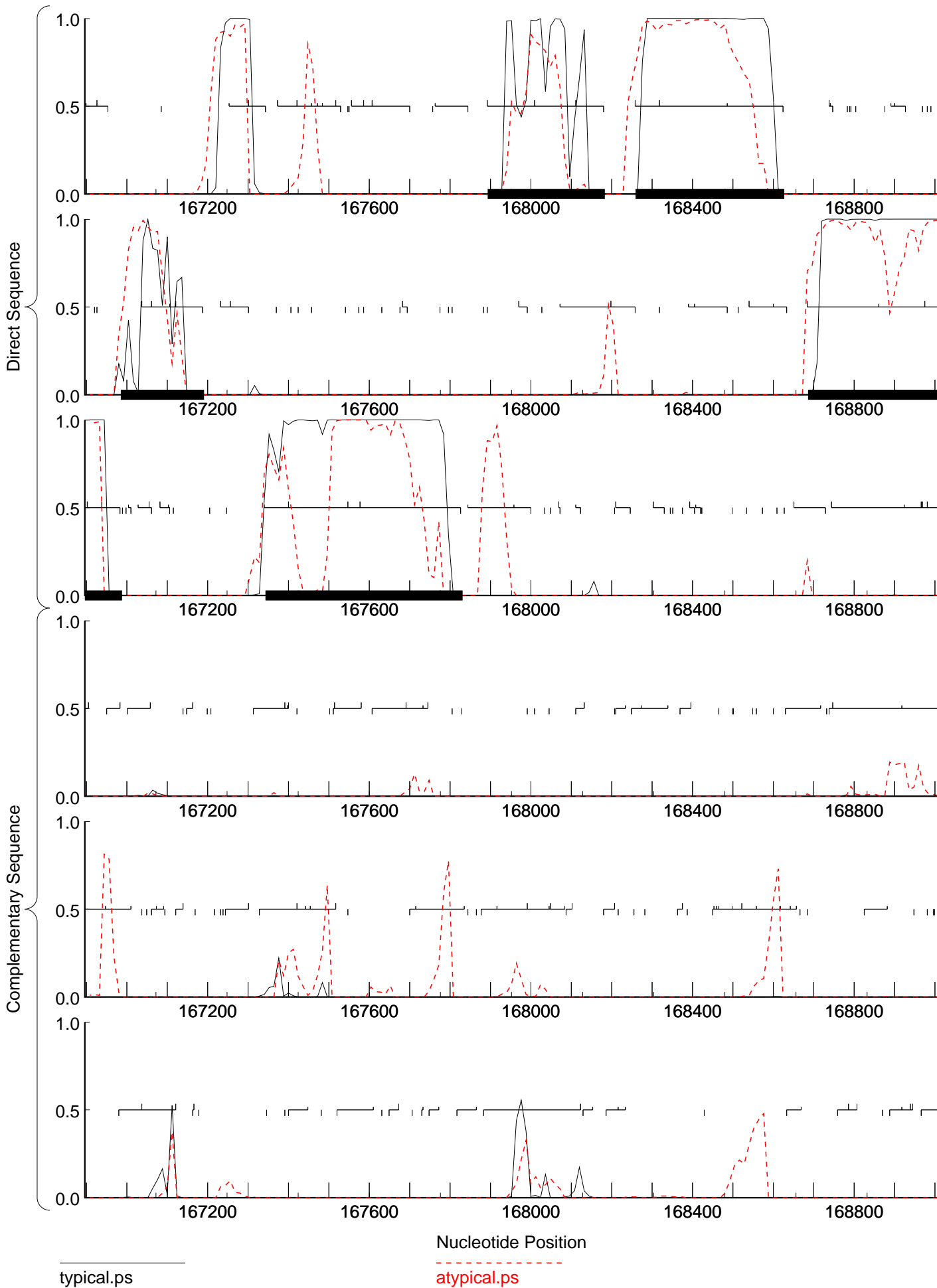
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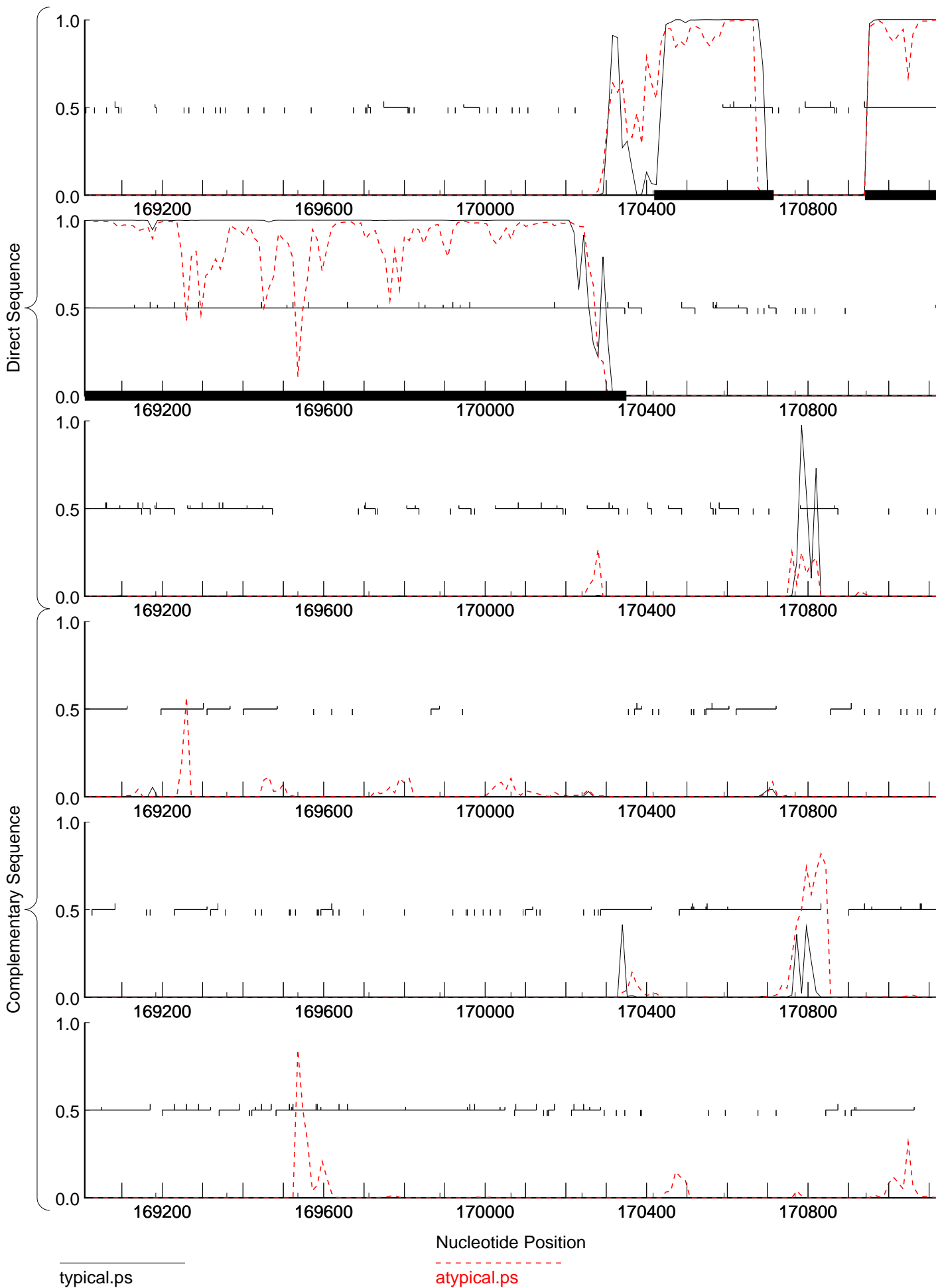


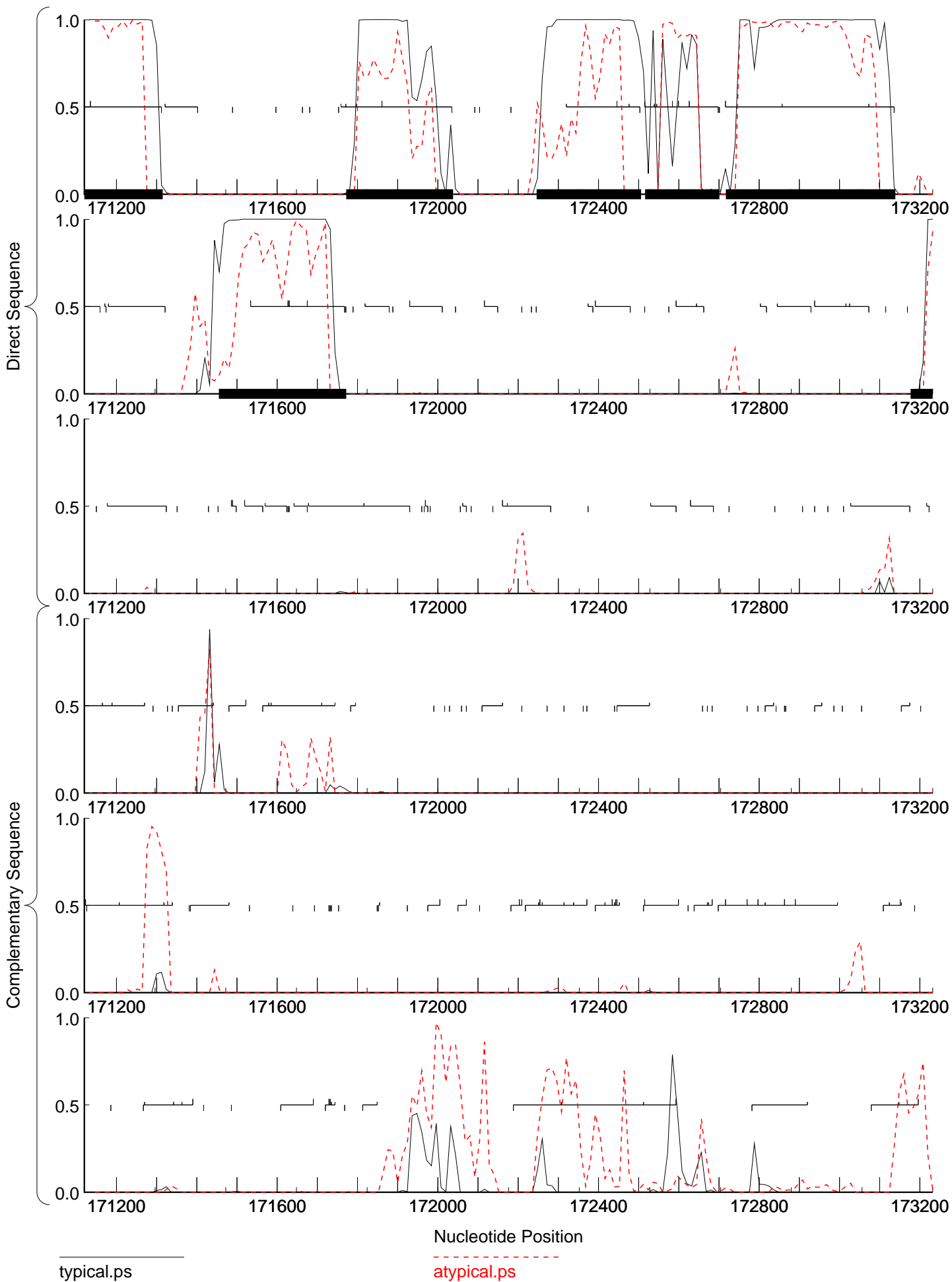
typical.ps

atypical.ps









typical.ps

atypical.ps

