

# GeneMark

Version 2.5p (09.08.06)

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## PROGRAM INFORMATION

Sequence : Mycobacterium phage ScottMcG complete sequence, 154017 bp, circularly permuted, Cluster C1  
Analysis Date : 8/20/18 at 8:28:36  
Pages : 74  
Sequence Length : 154017 bp  
GC Content : 64.83%

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,

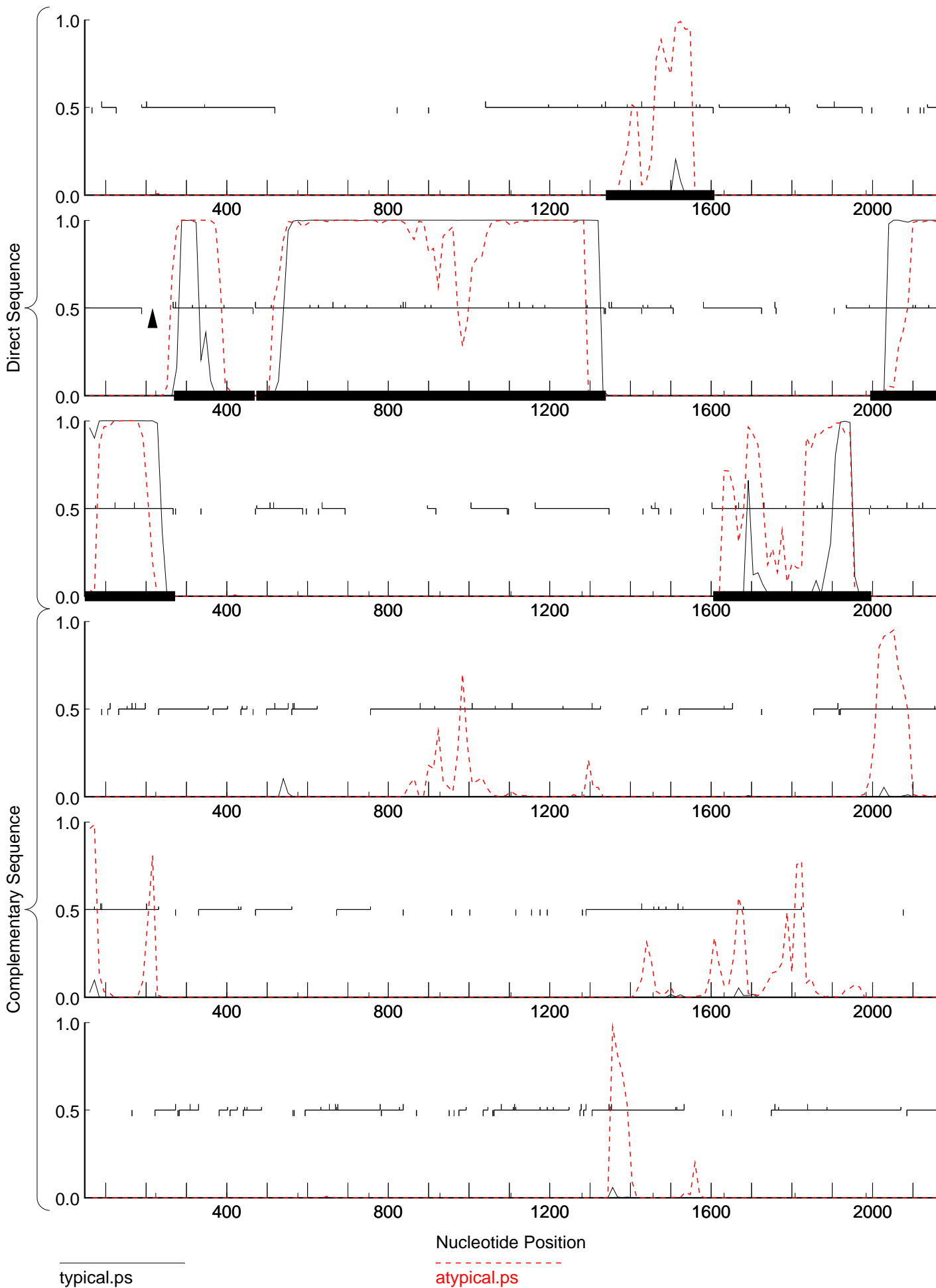
## MATRIX INFORMATION

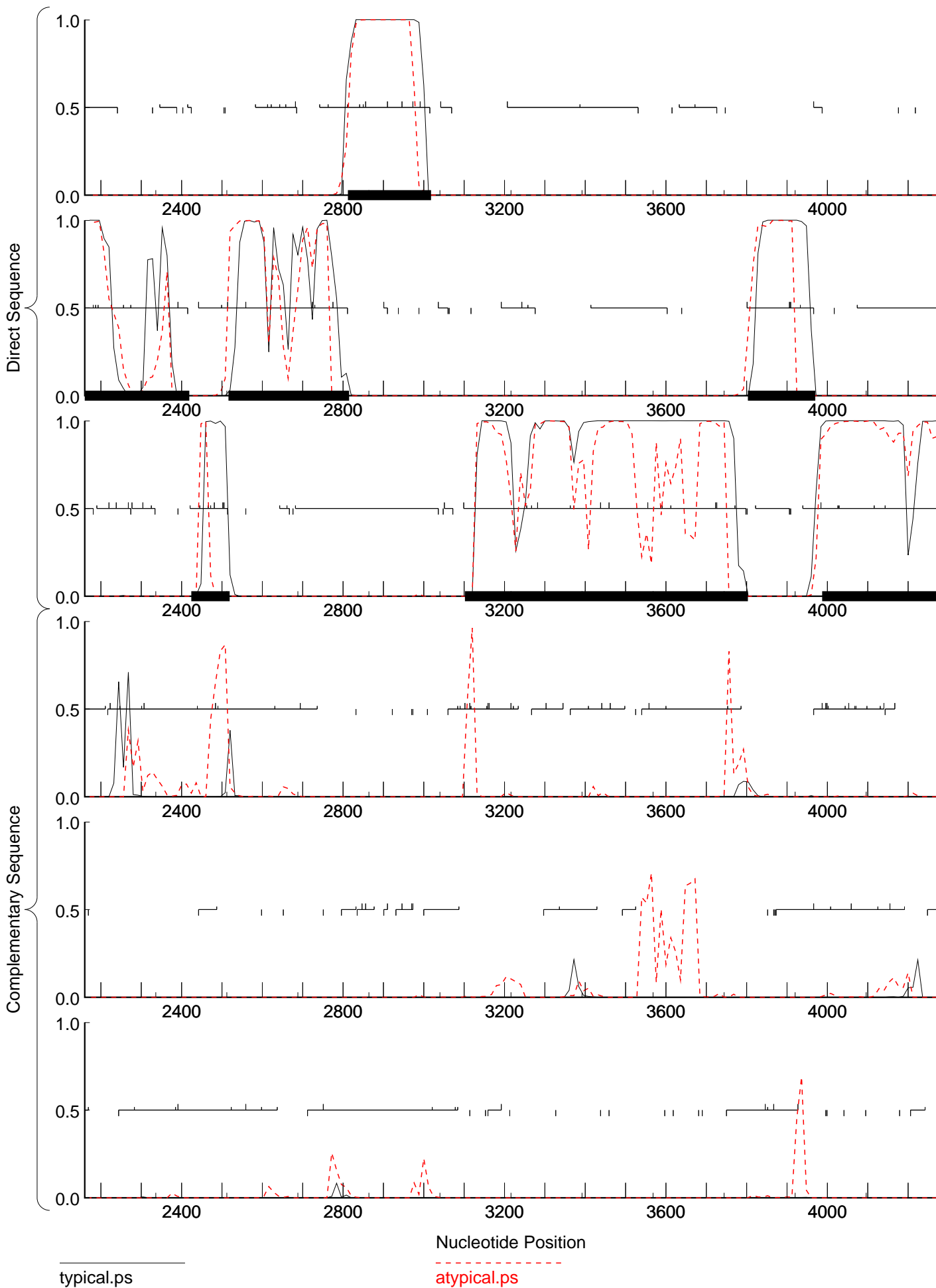
Matrix : -  
Author : -  
Order : 4

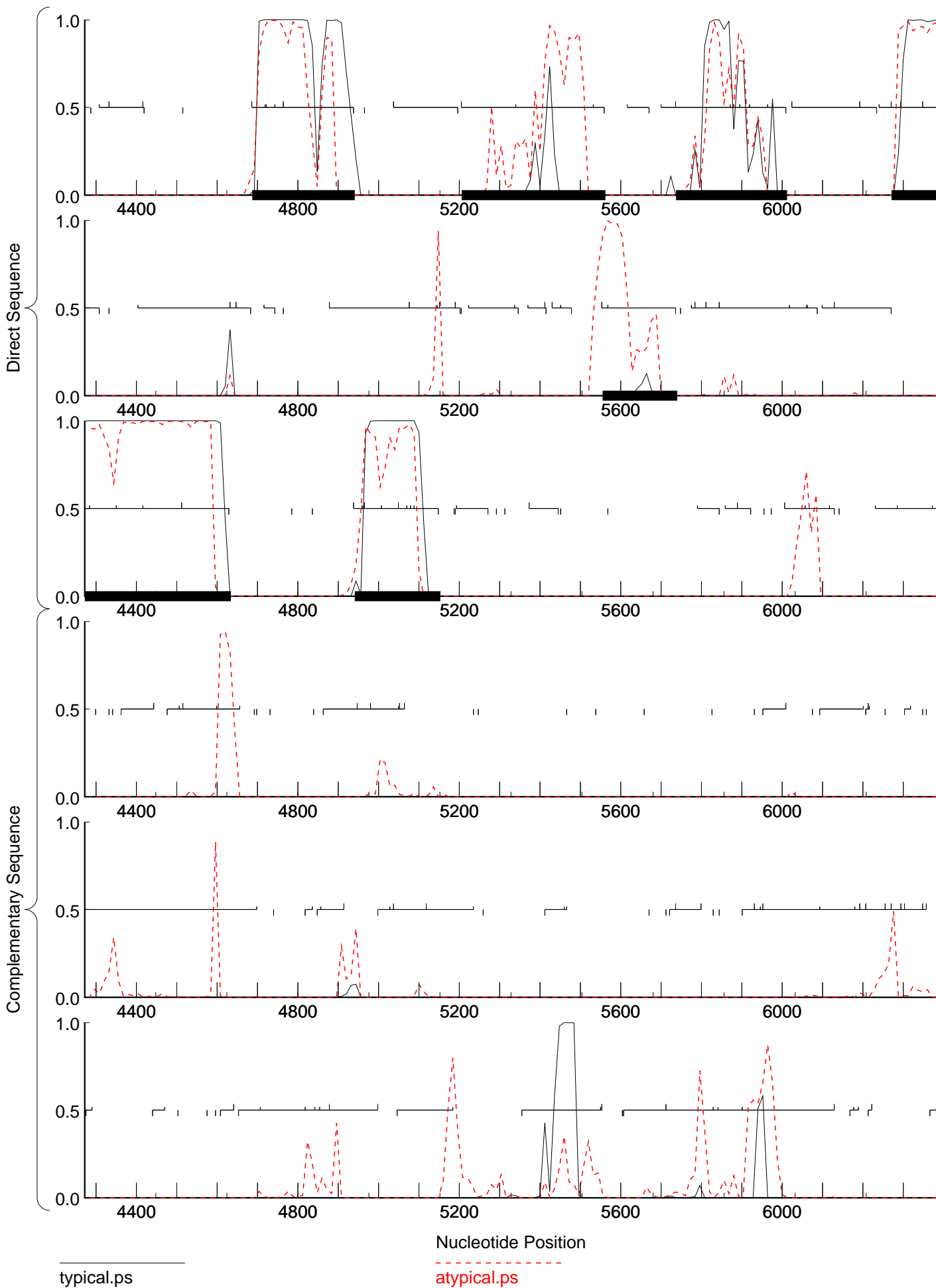
Send questions / comments to:  
Dr. M. Borodovsky  
Georgia Institute of Technology  
School of Biology  
Atlanta, GA 30332-0230

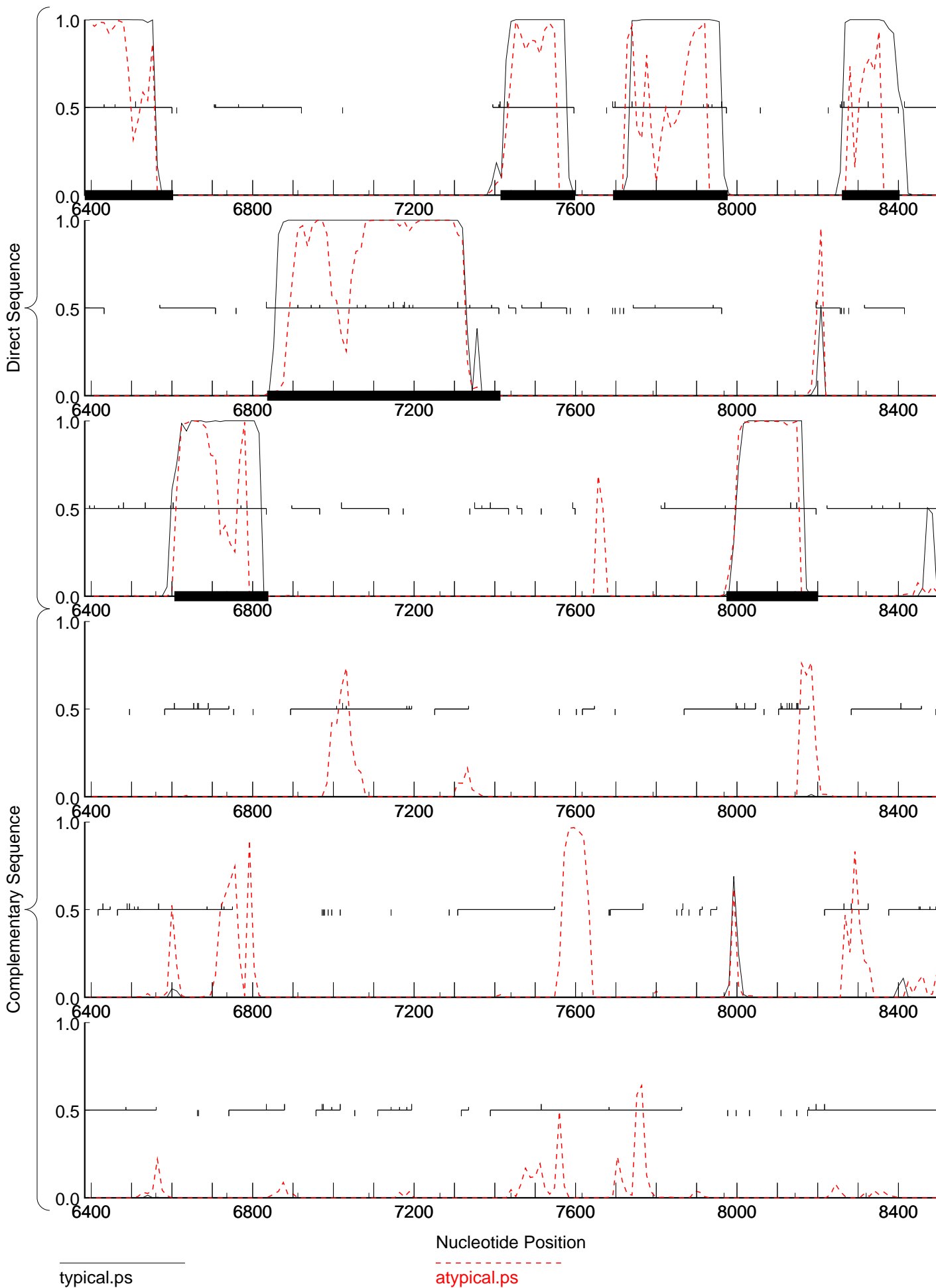
### *Matrix notes & comments*

Training set derived by GeneMarkS, 4.30 October 2014  
Mon Aug 20 08:28:35 2018



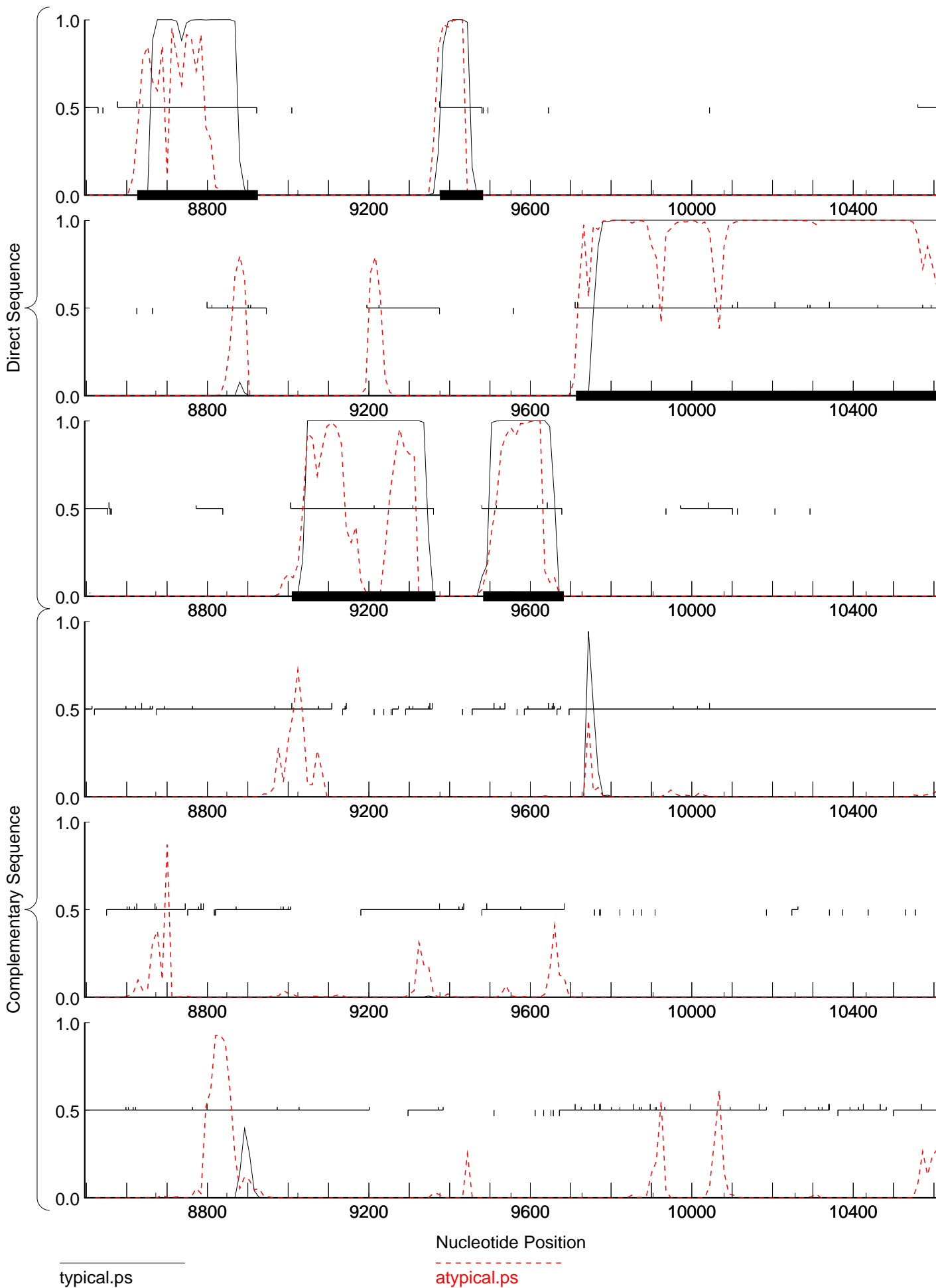


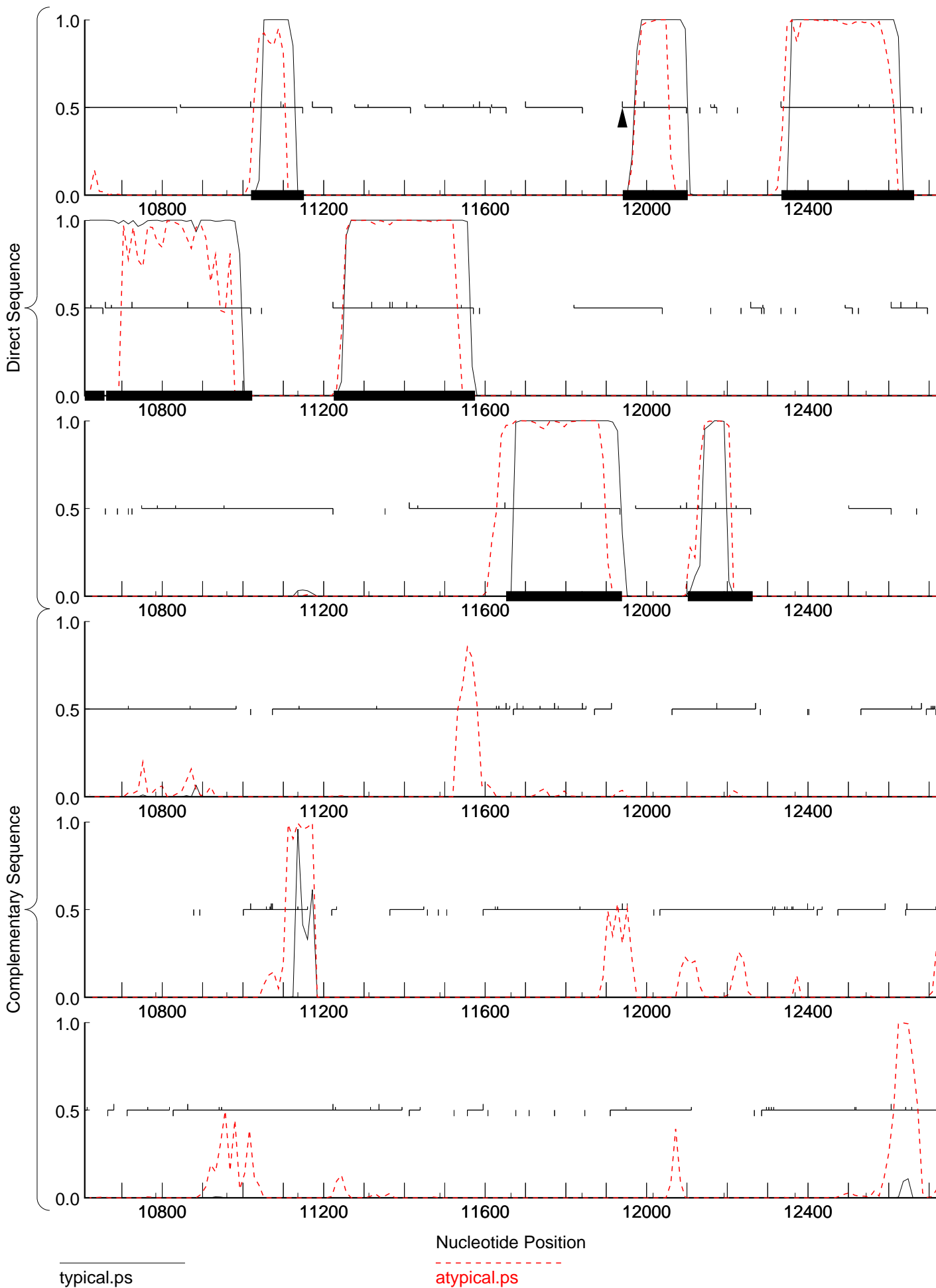


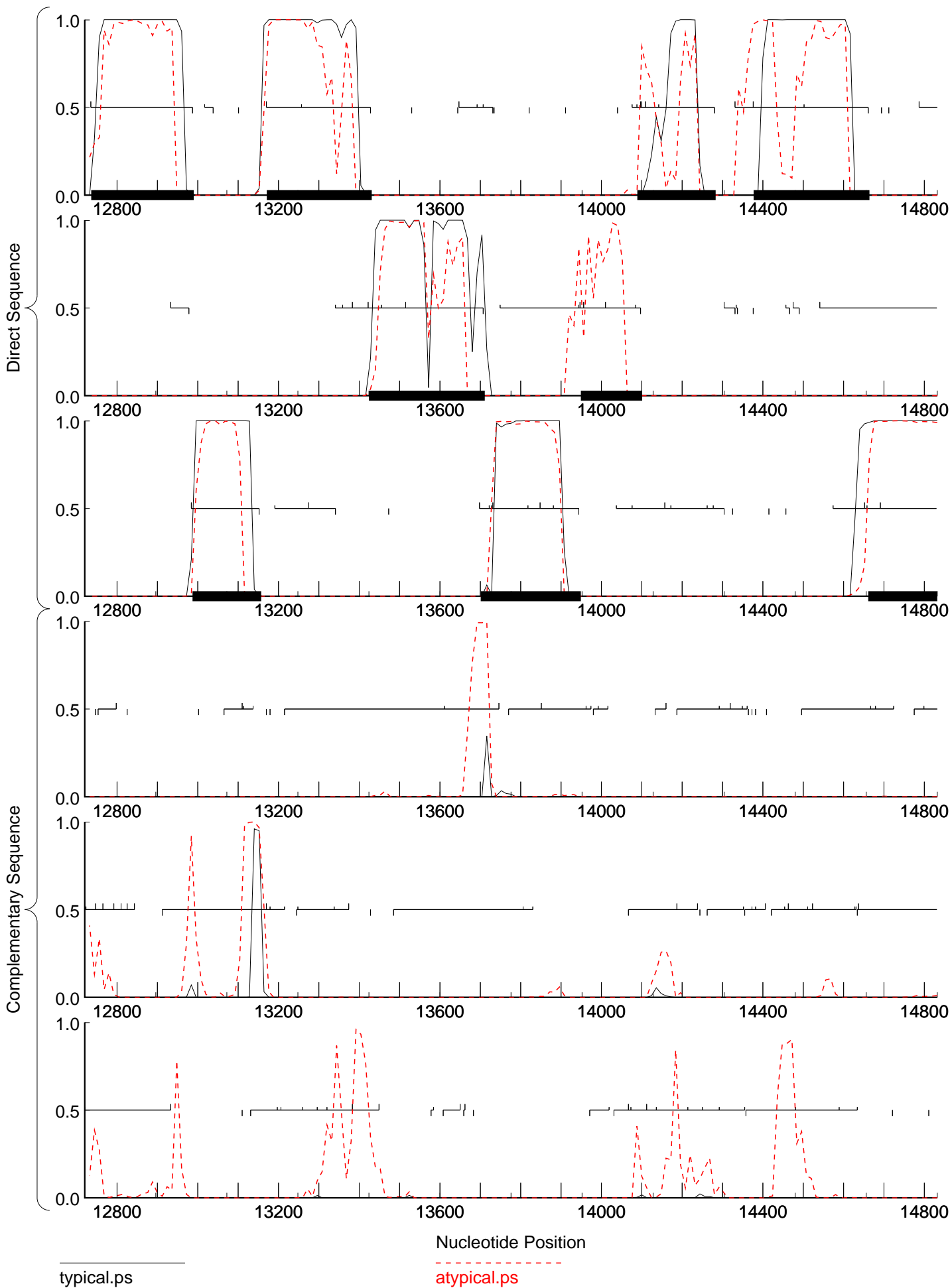


typical.ps

atypical.ps



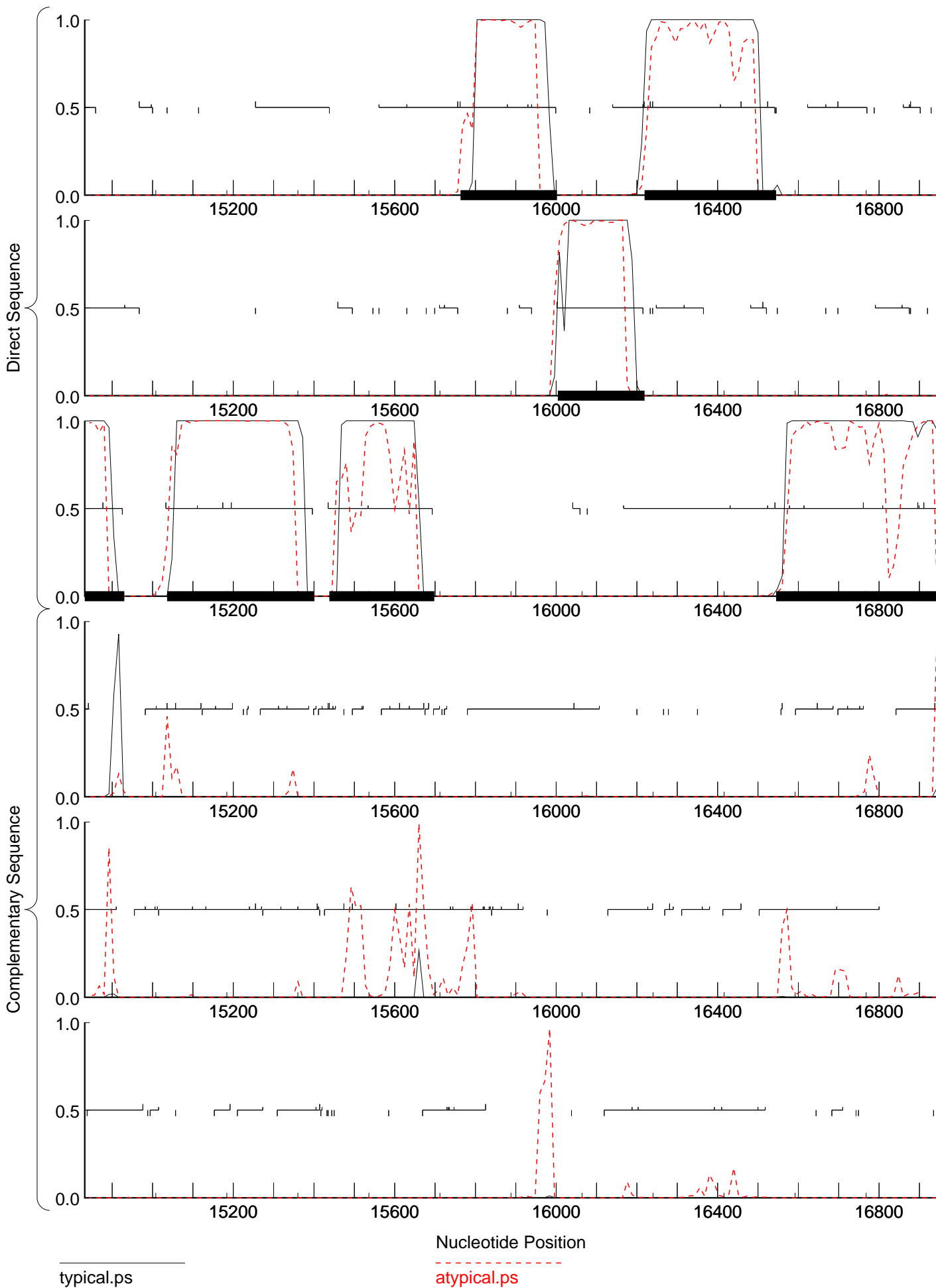




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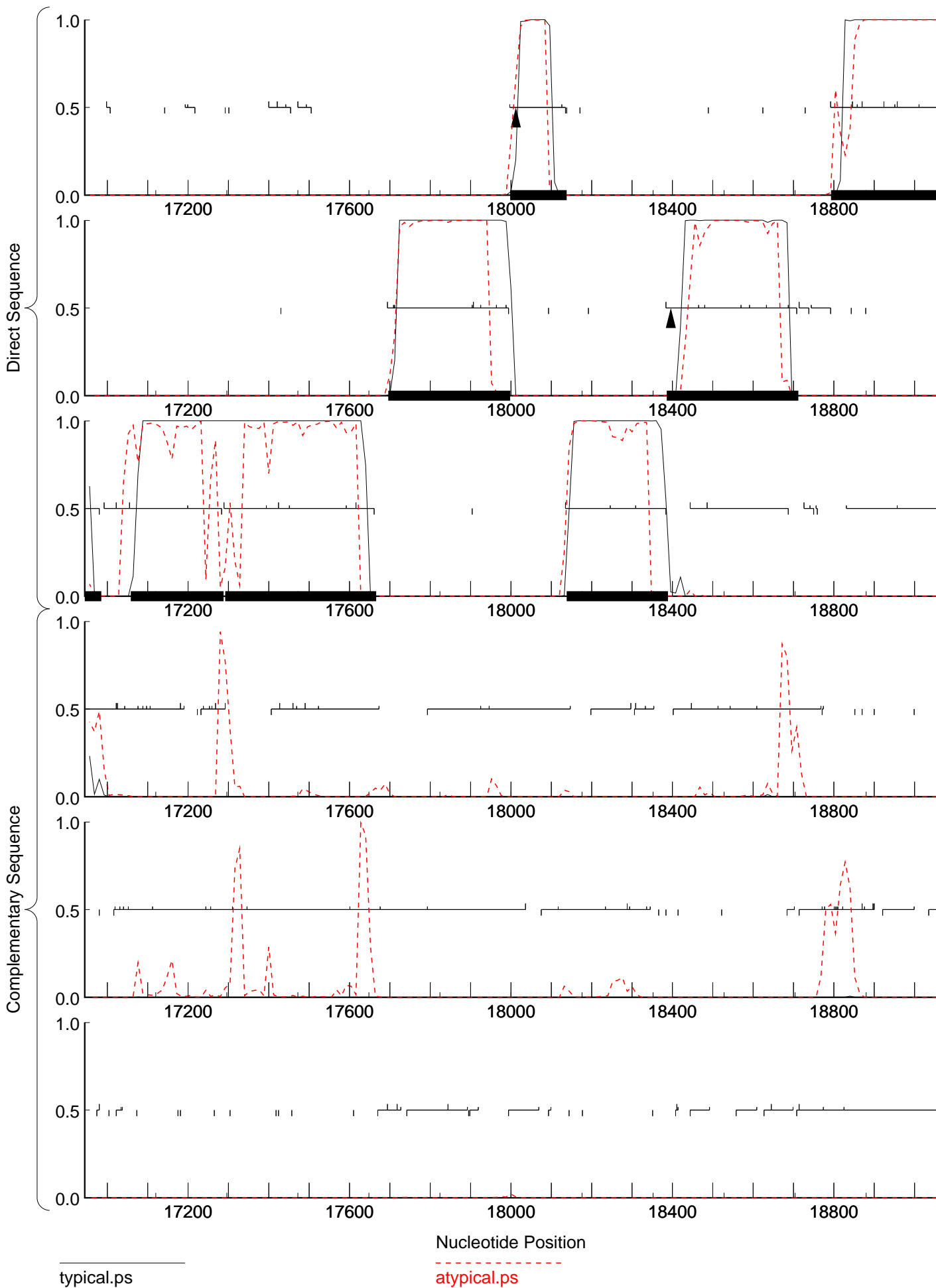
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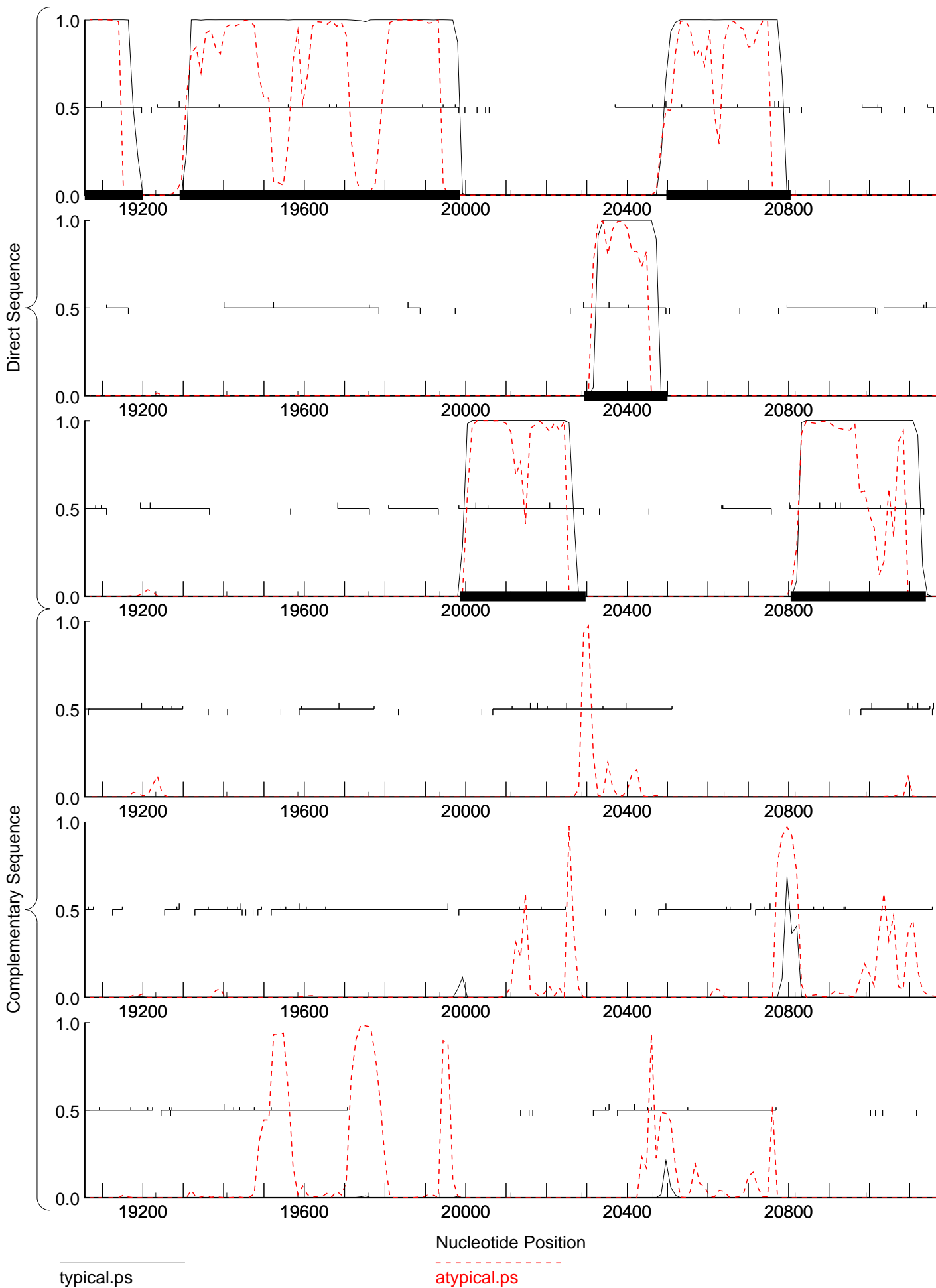
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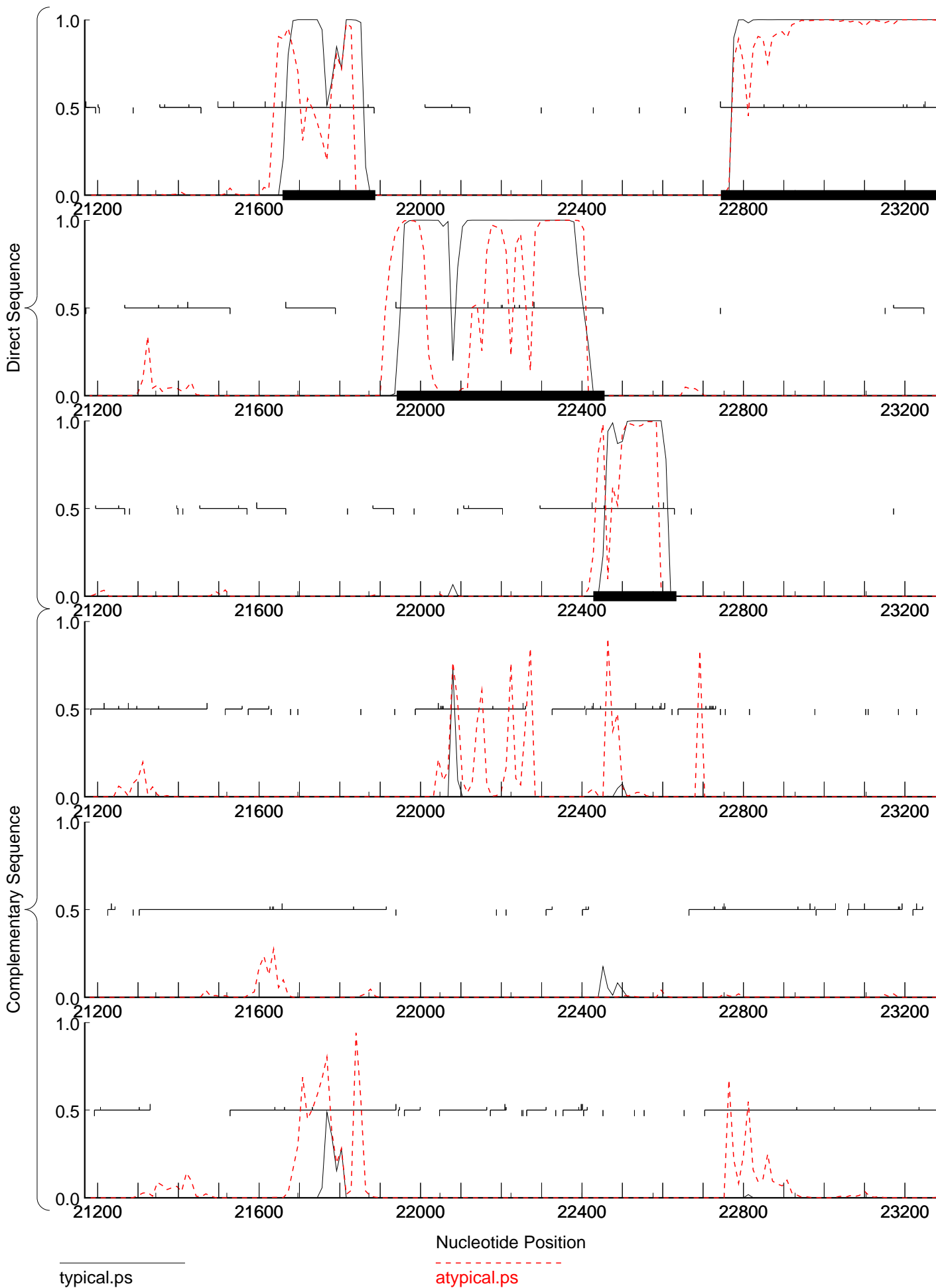
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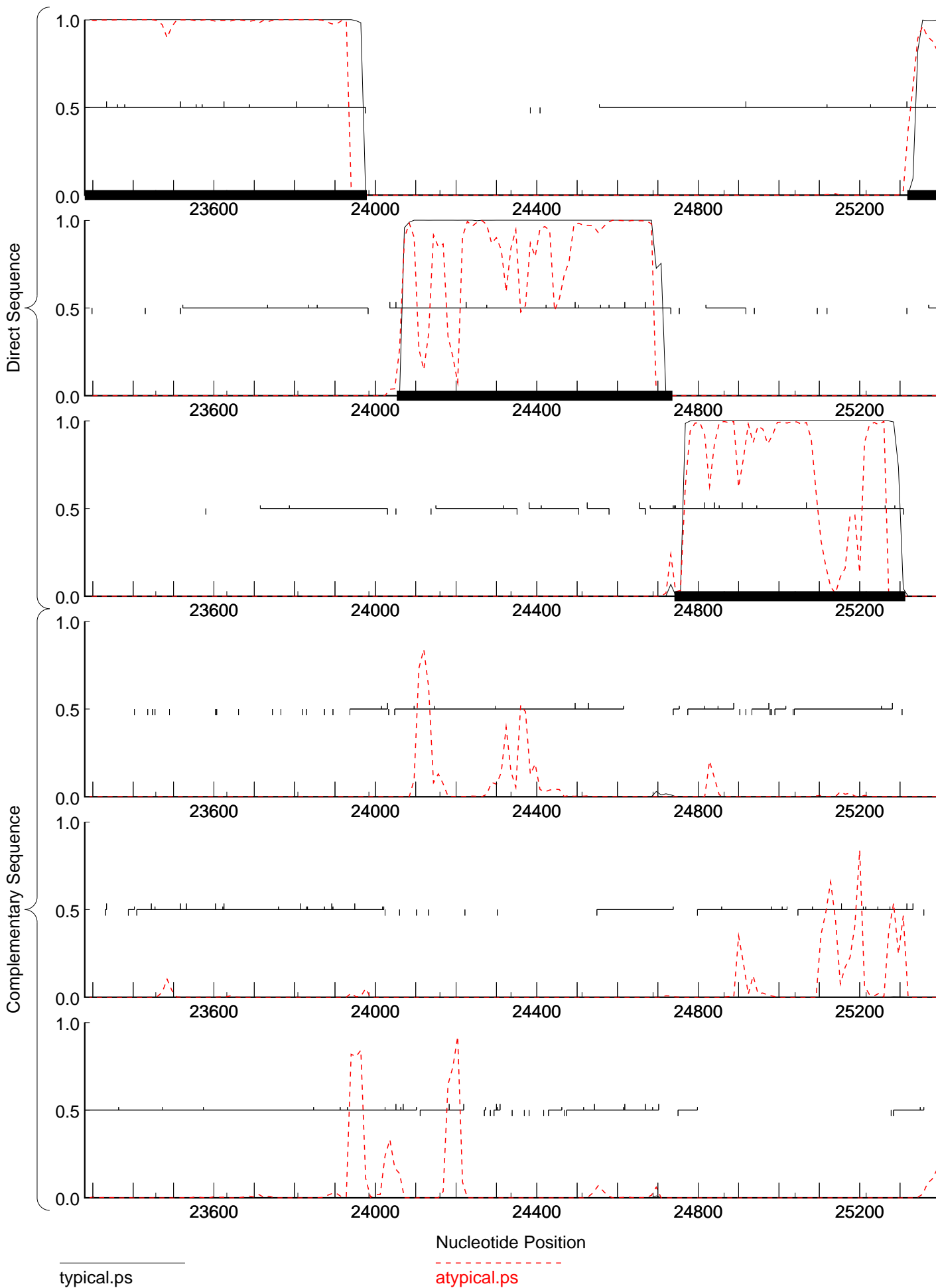


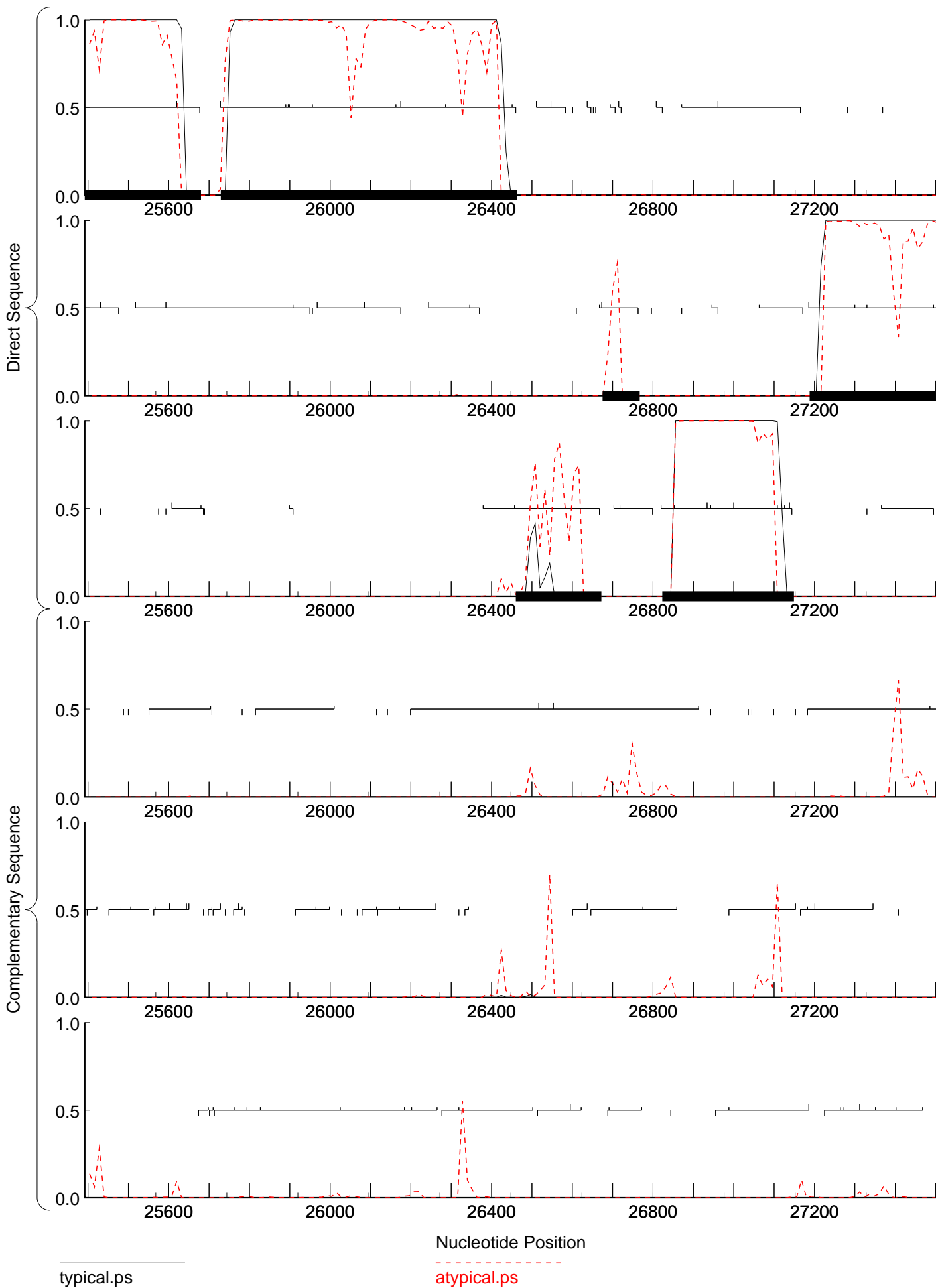
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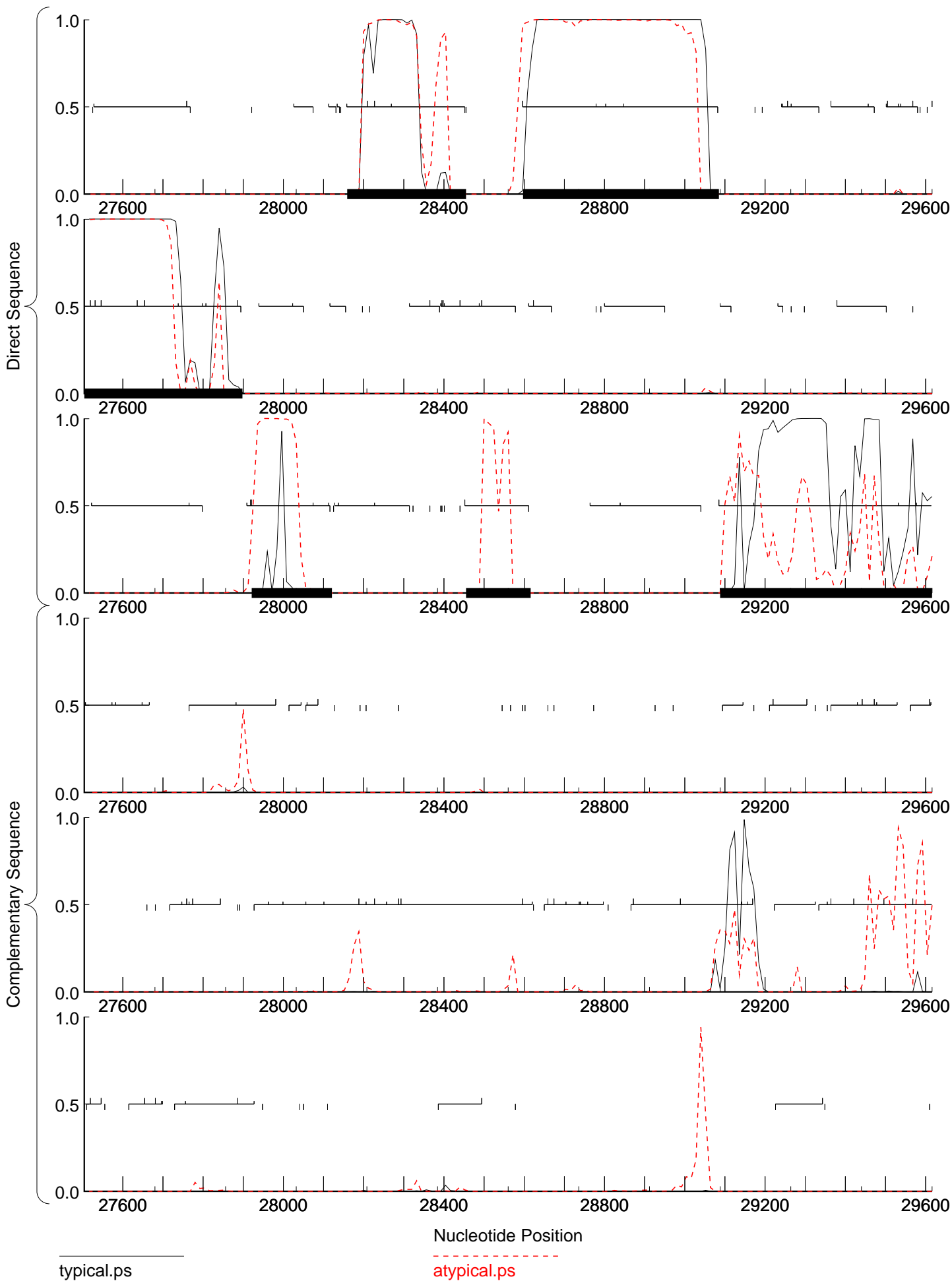
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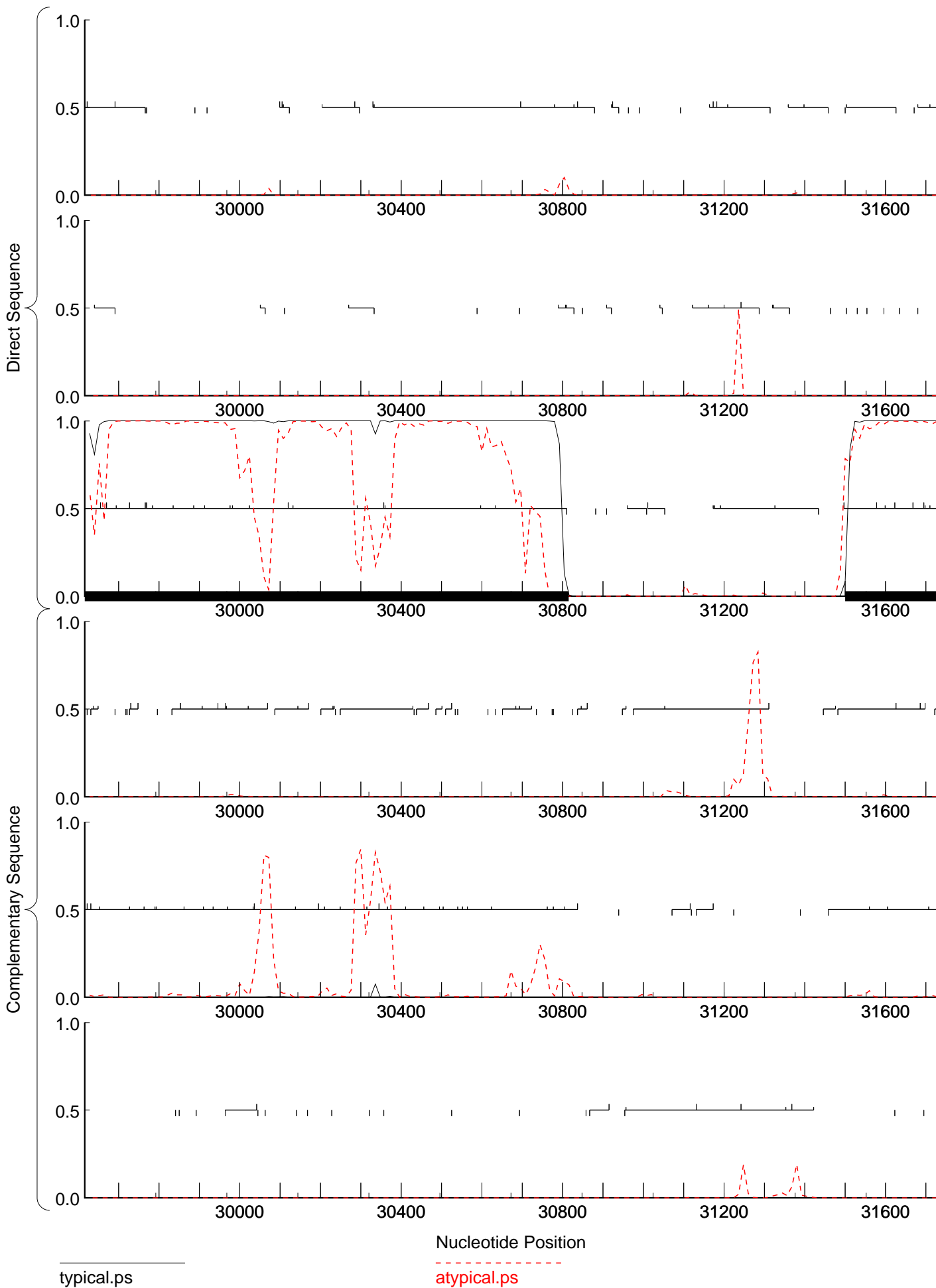




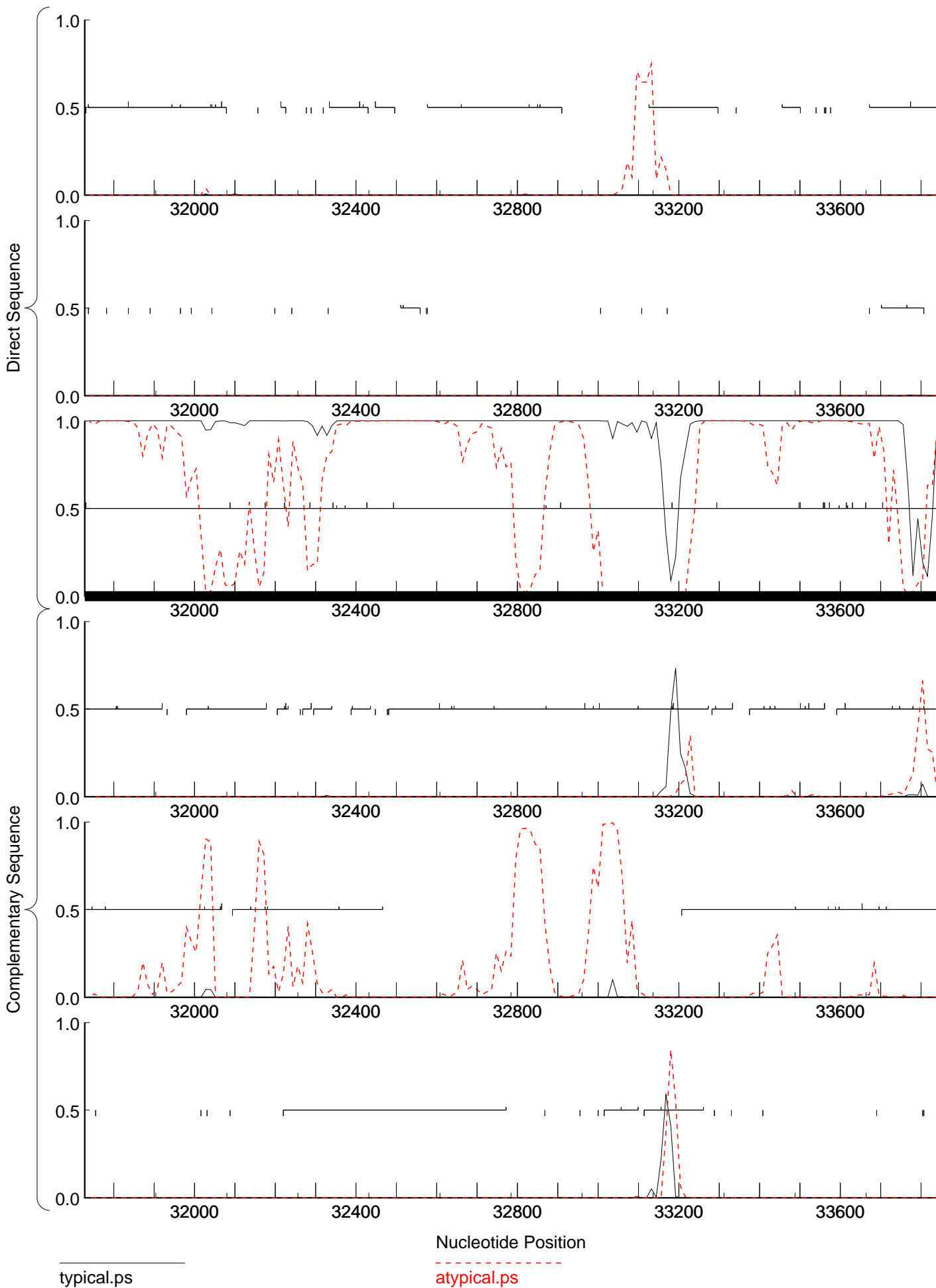


typical.ps

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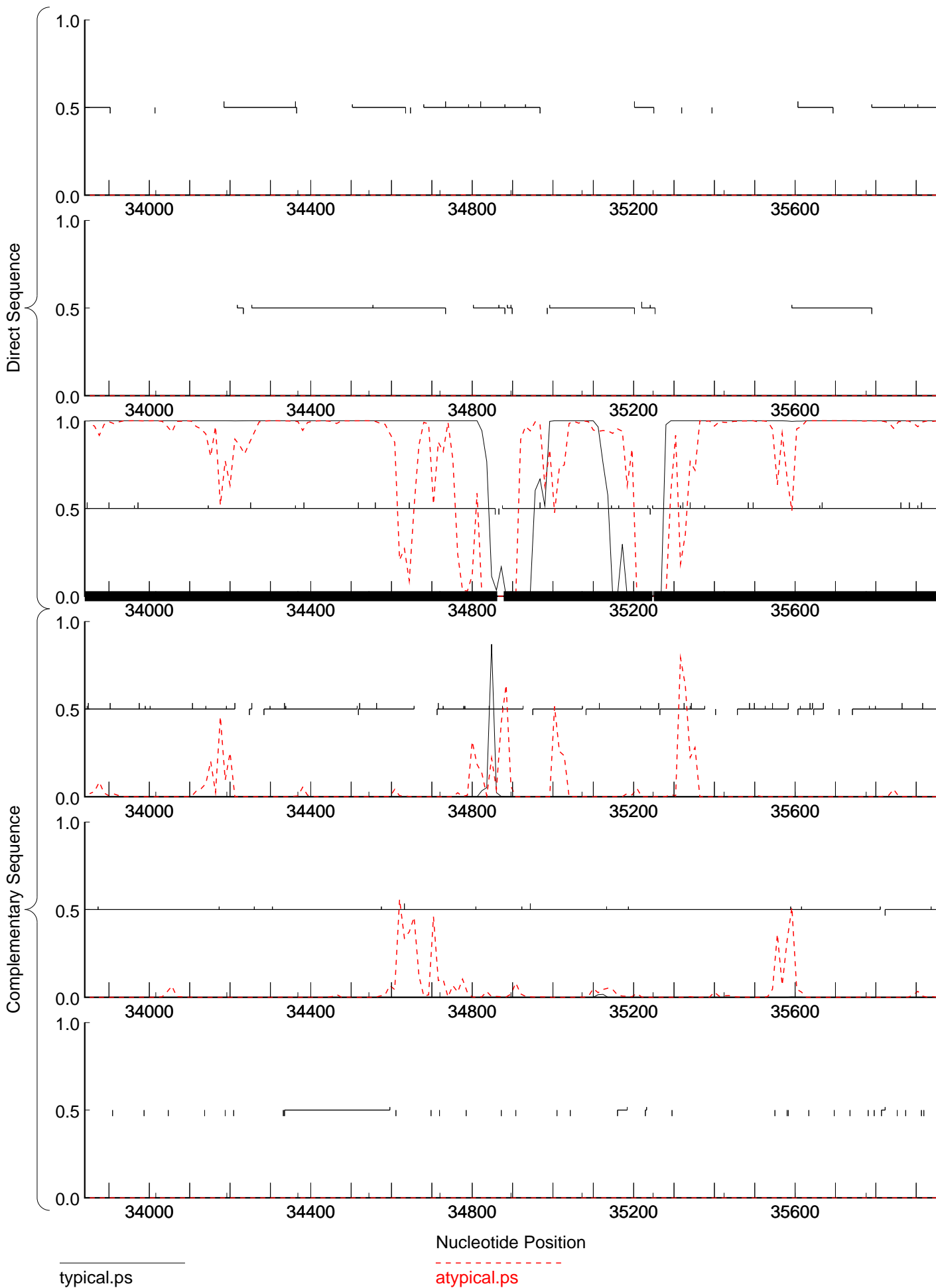


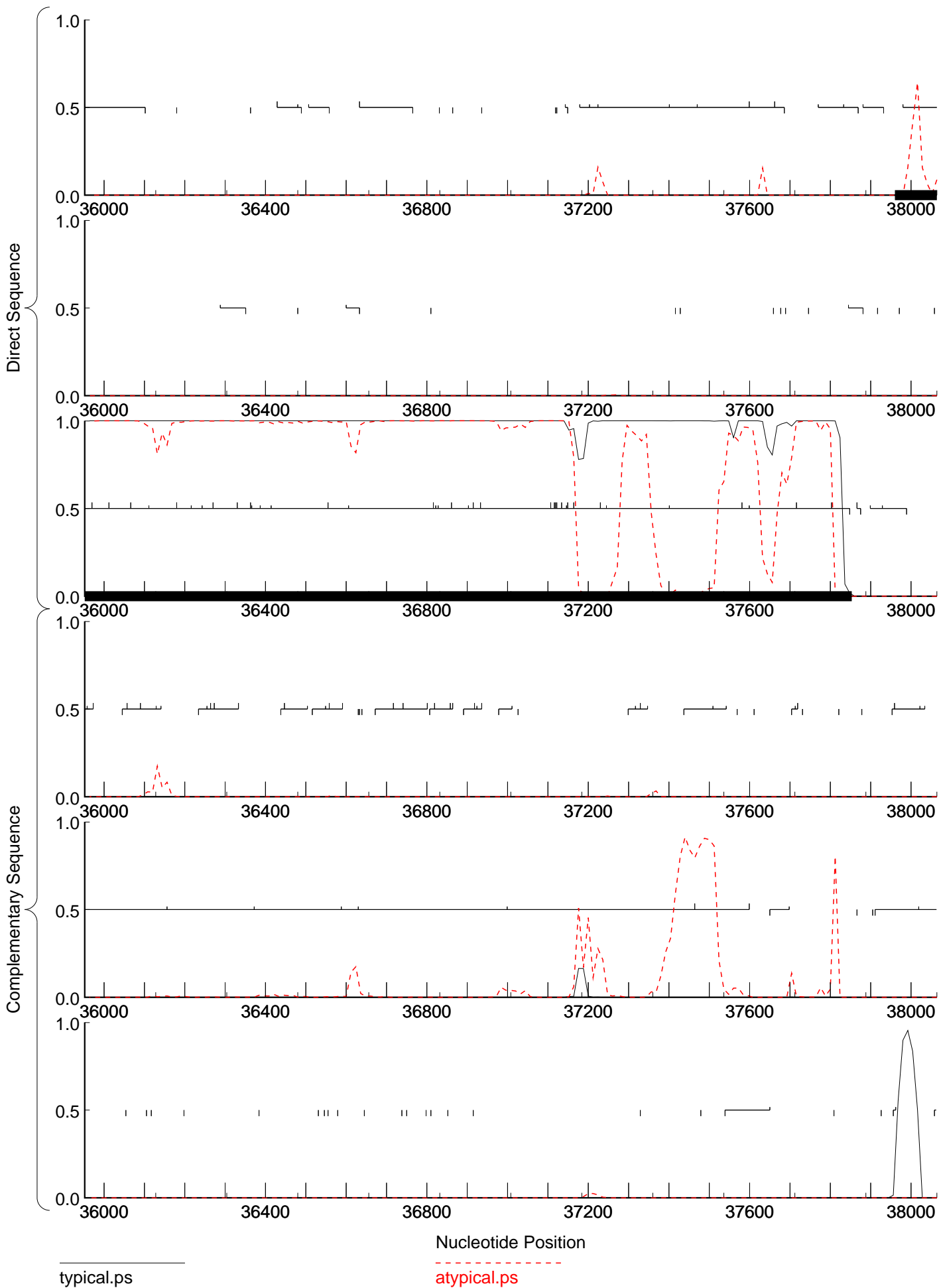


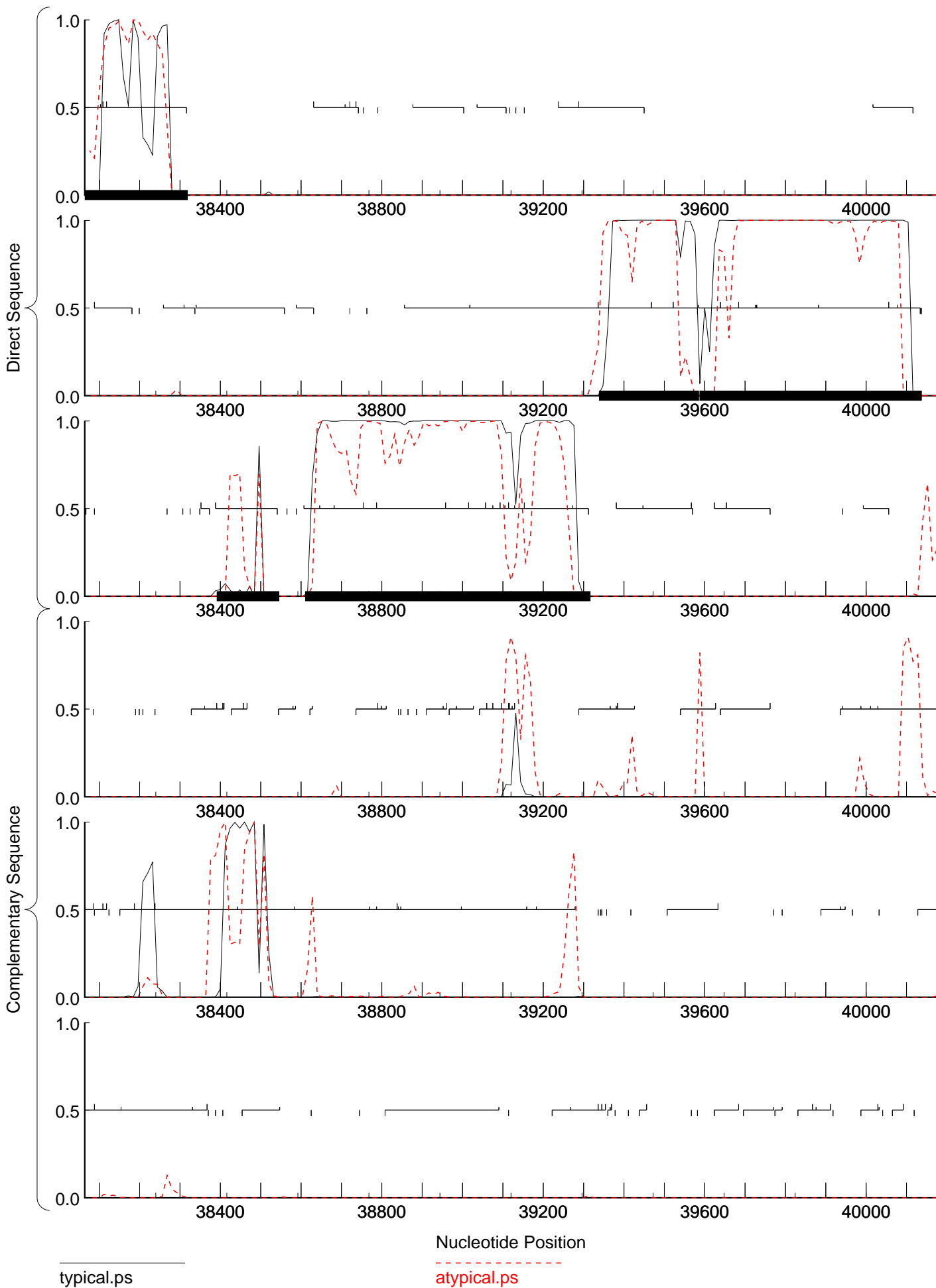


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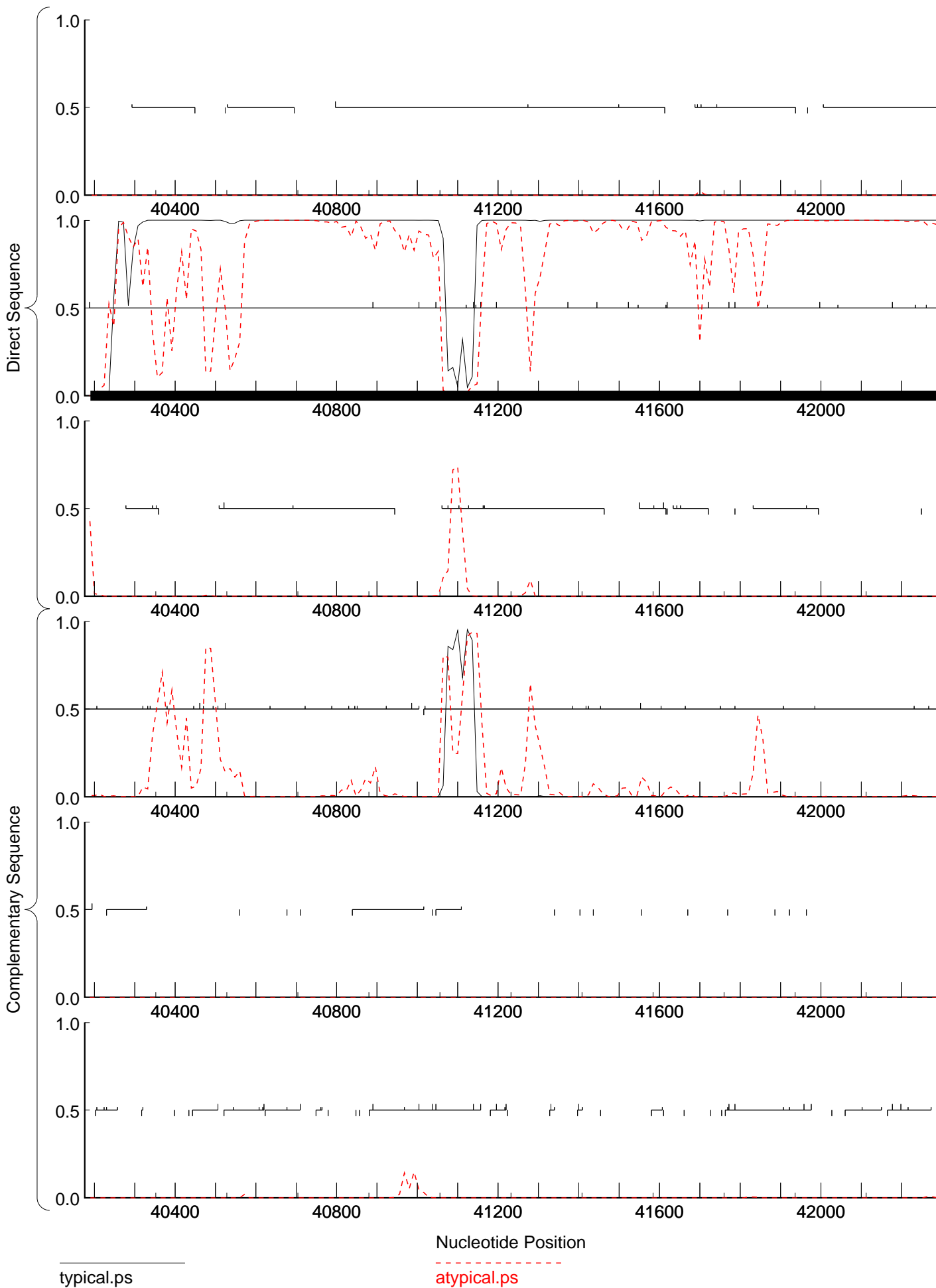


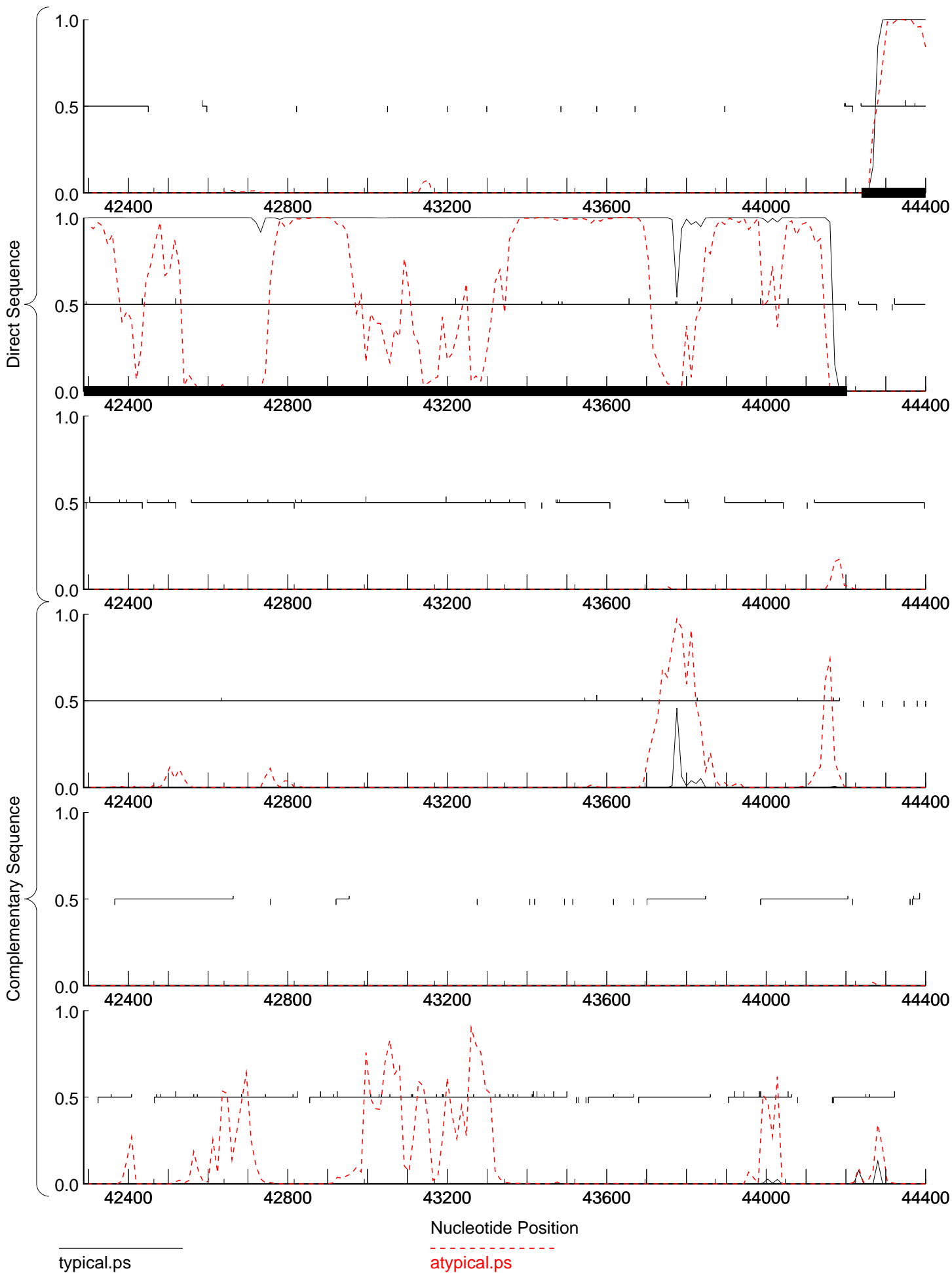


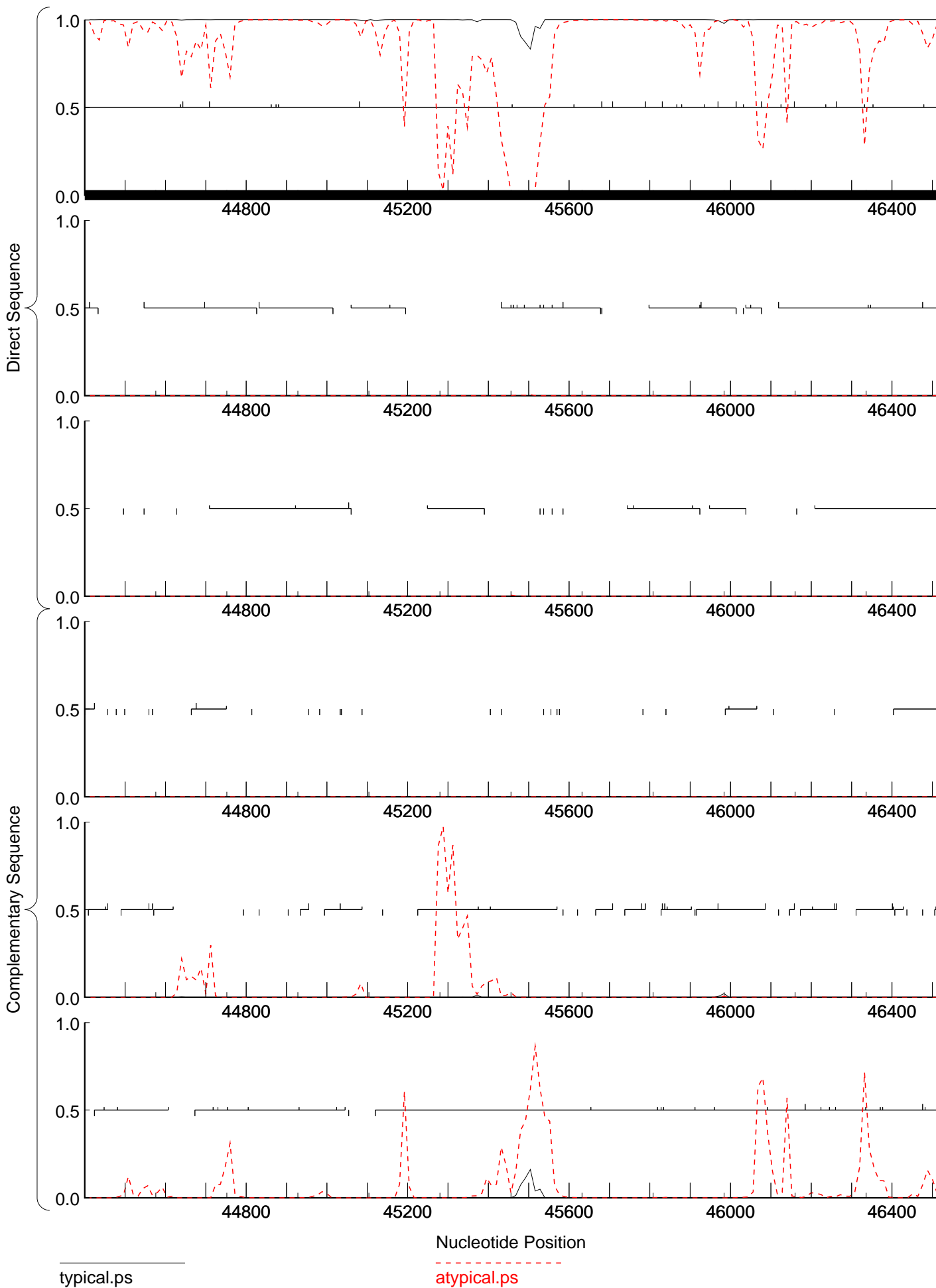


typical.ps

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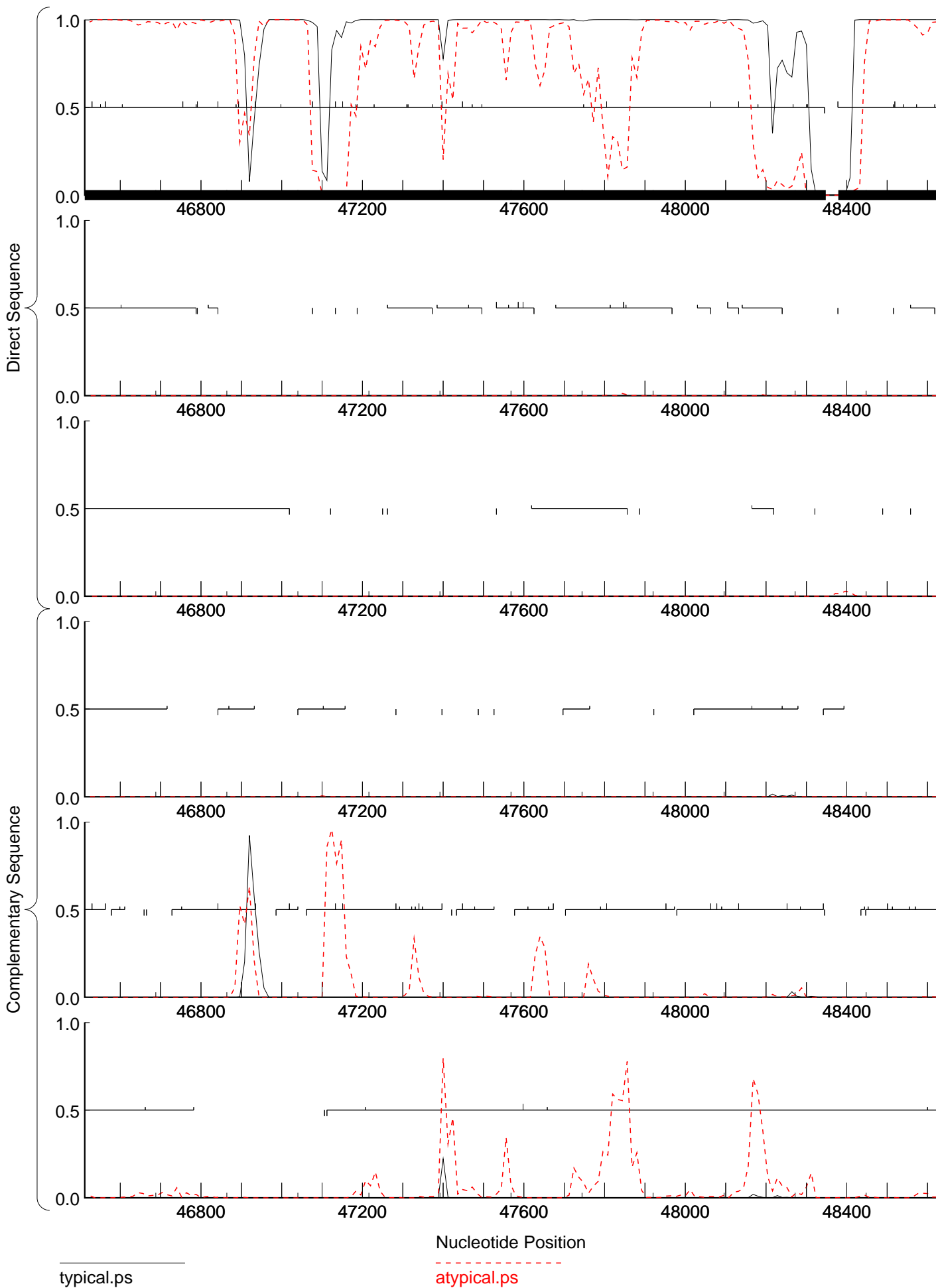






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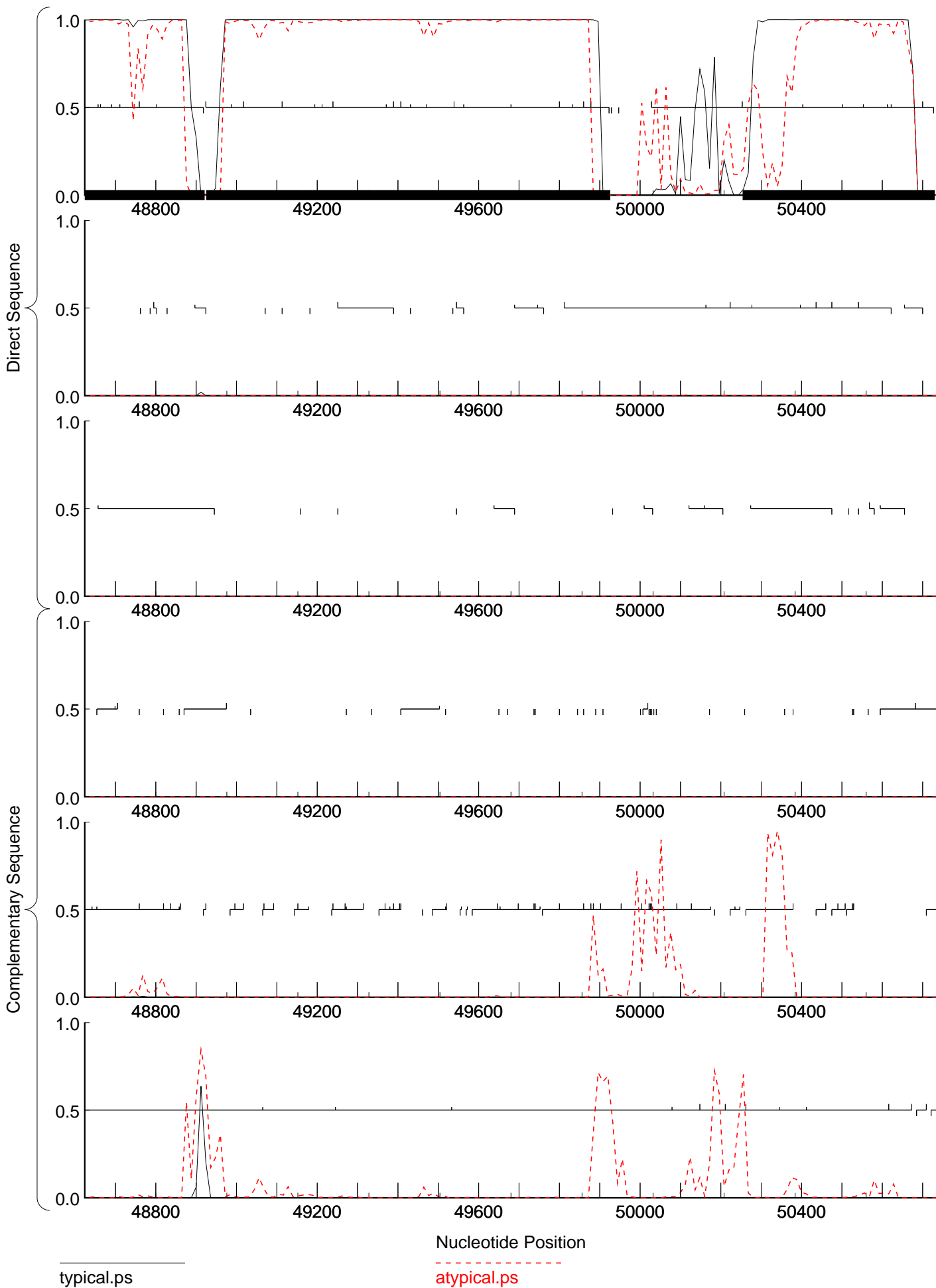
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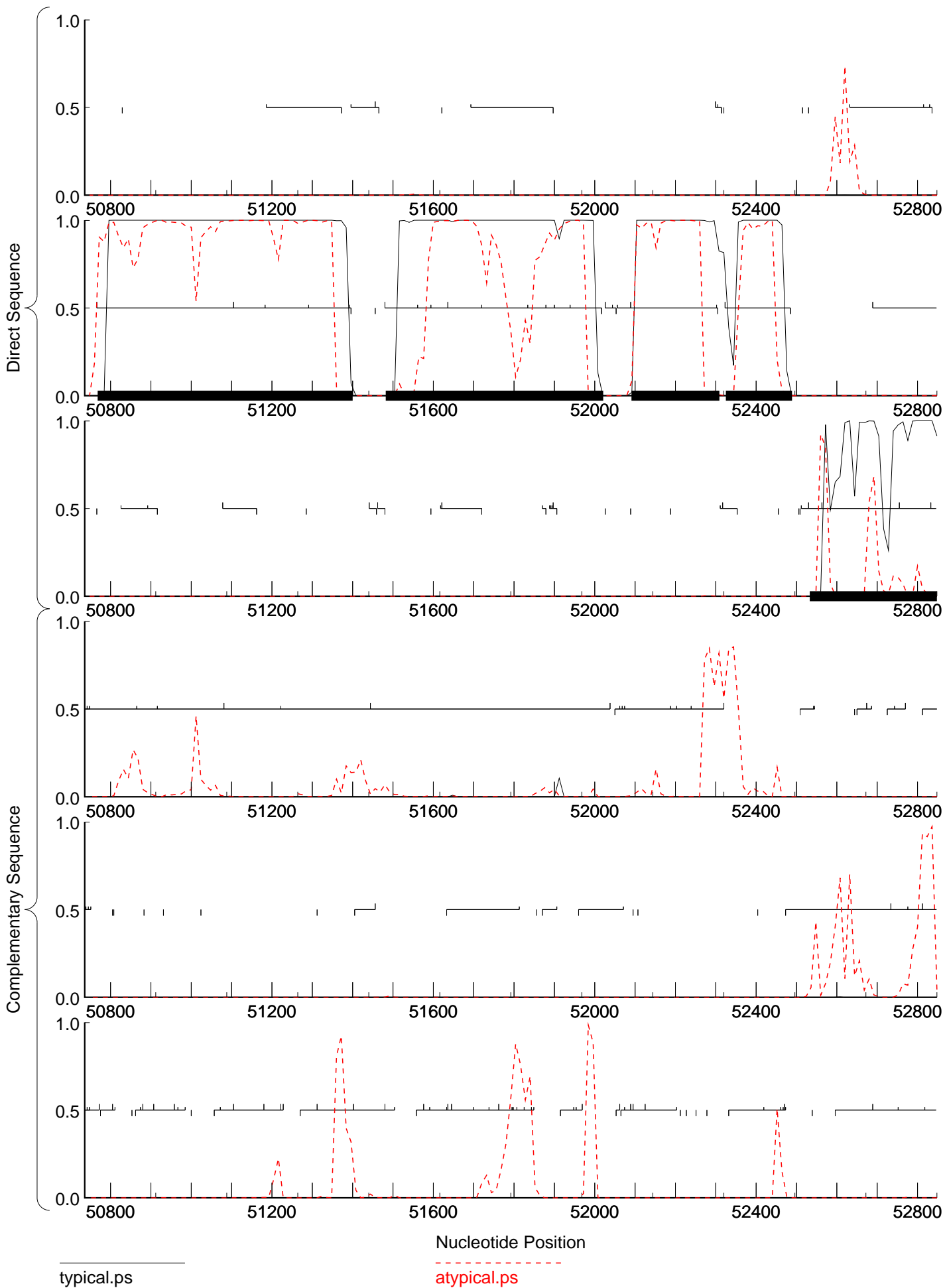


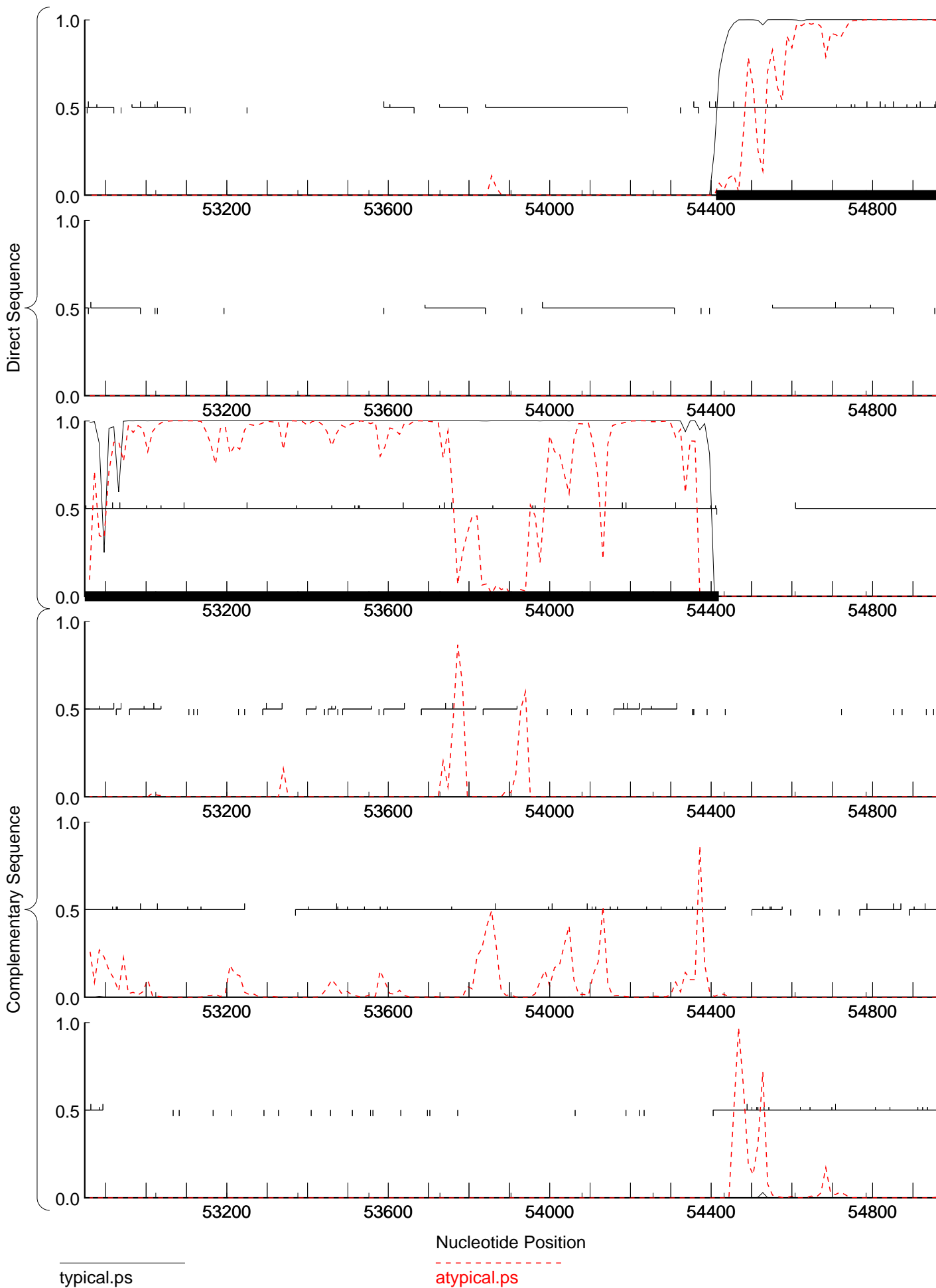
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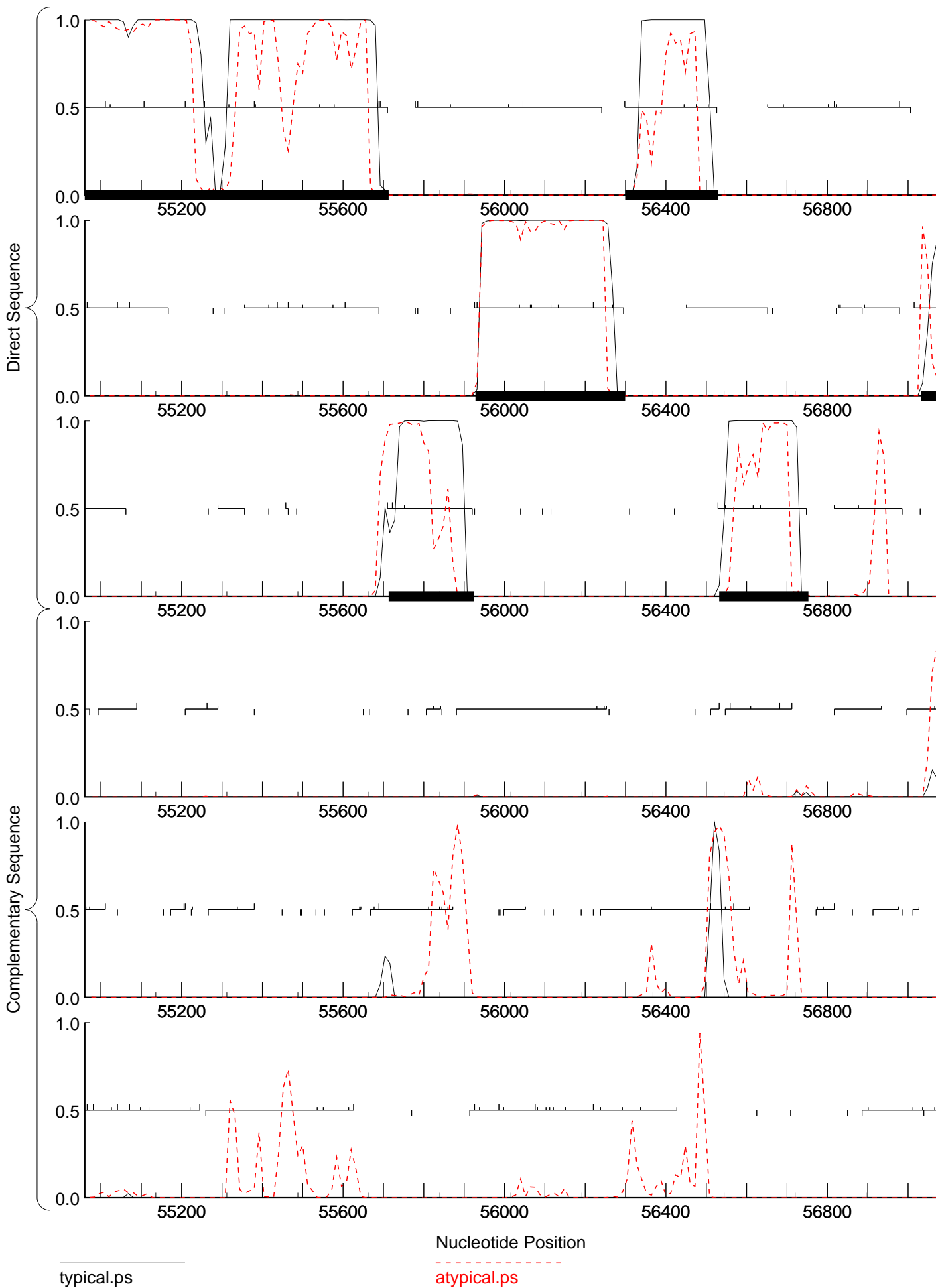
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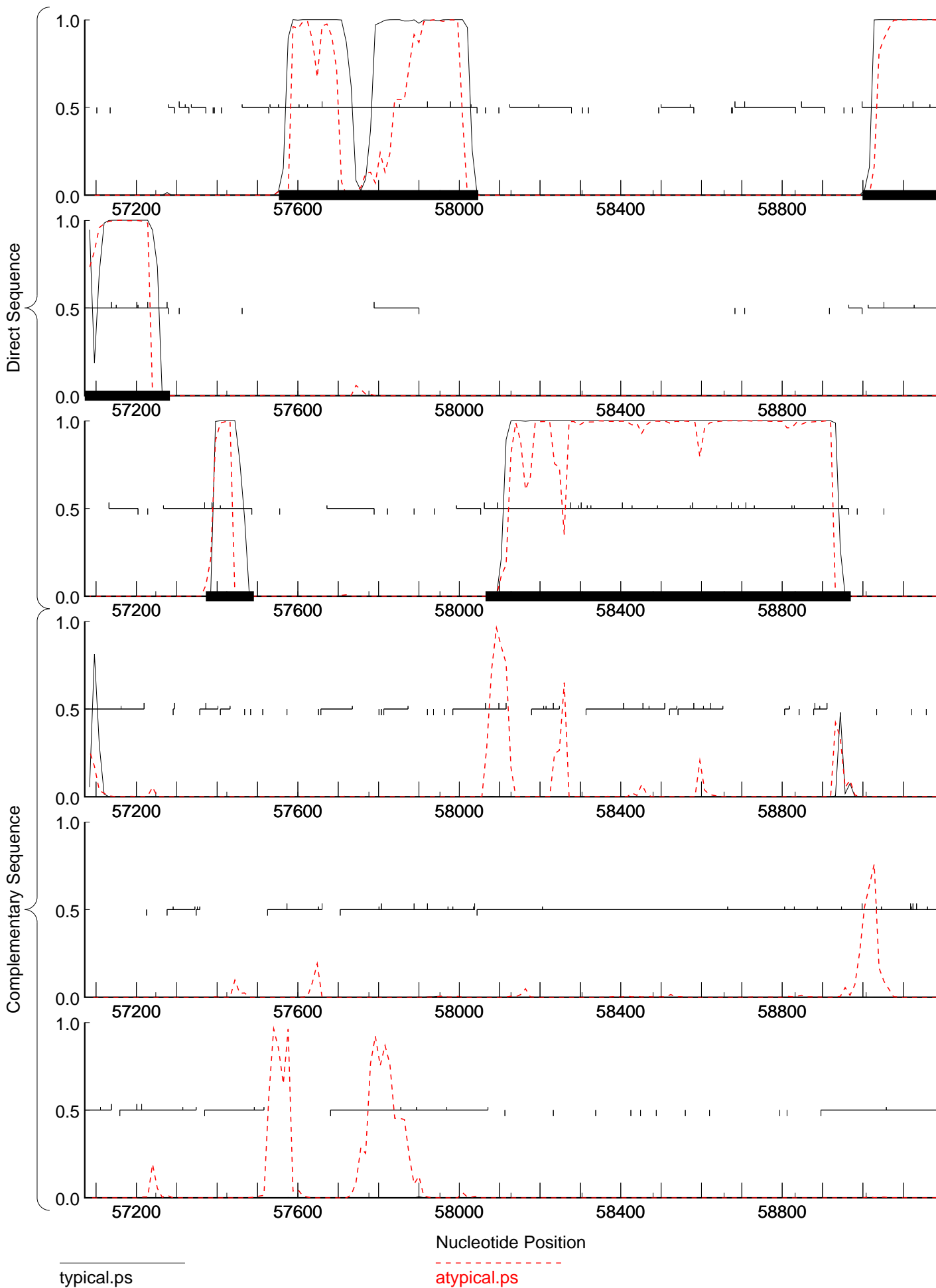






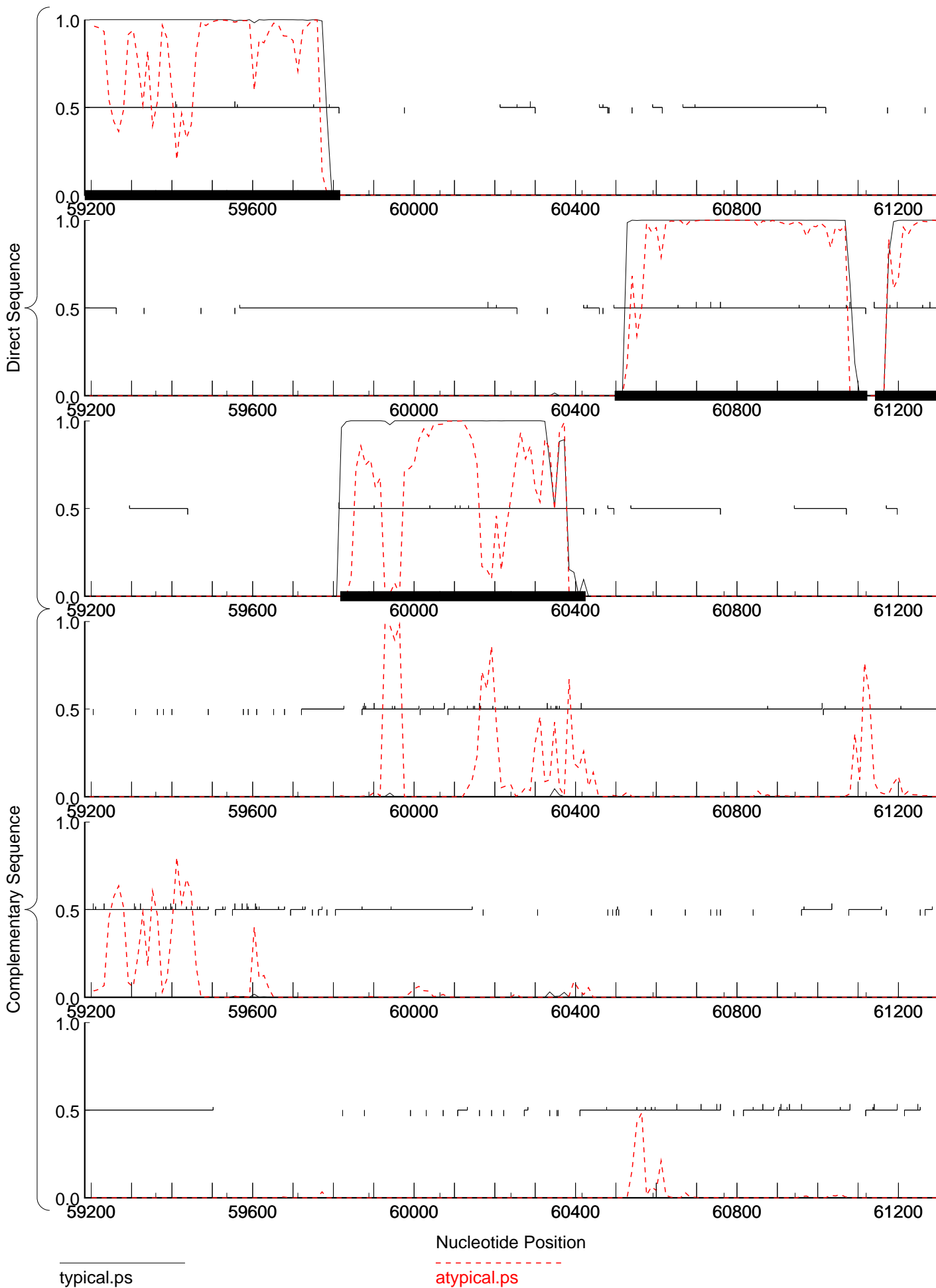
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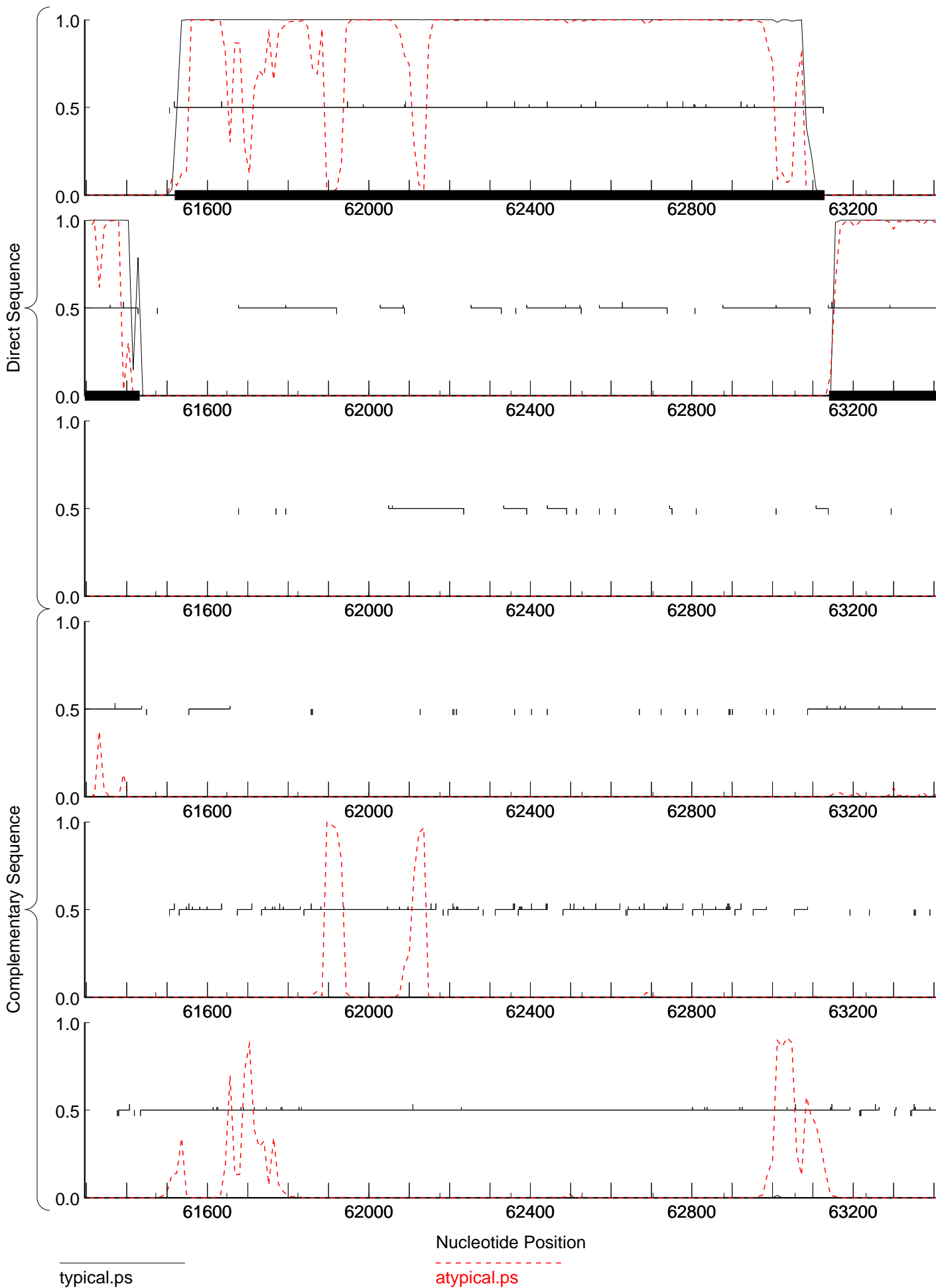
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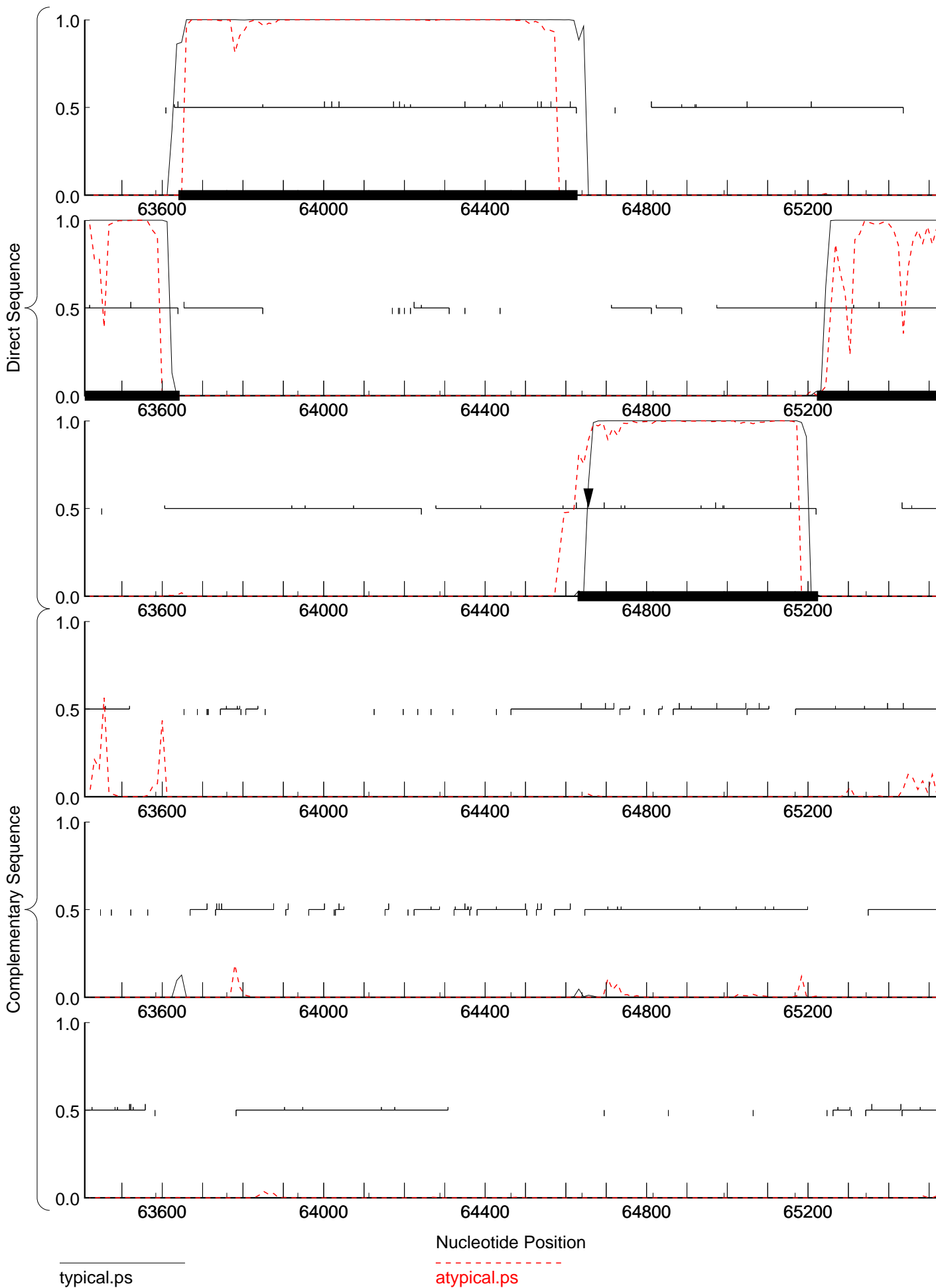


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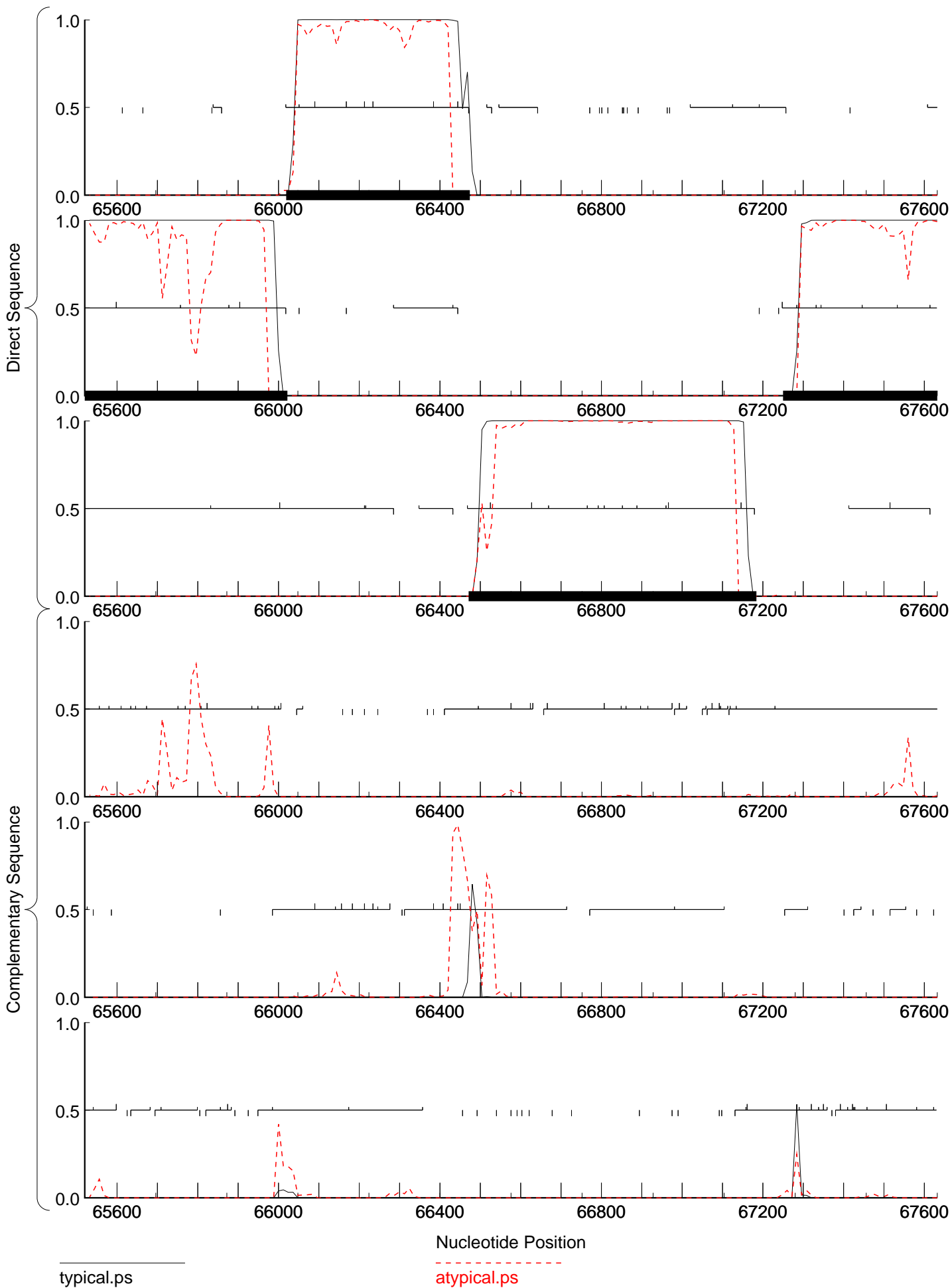
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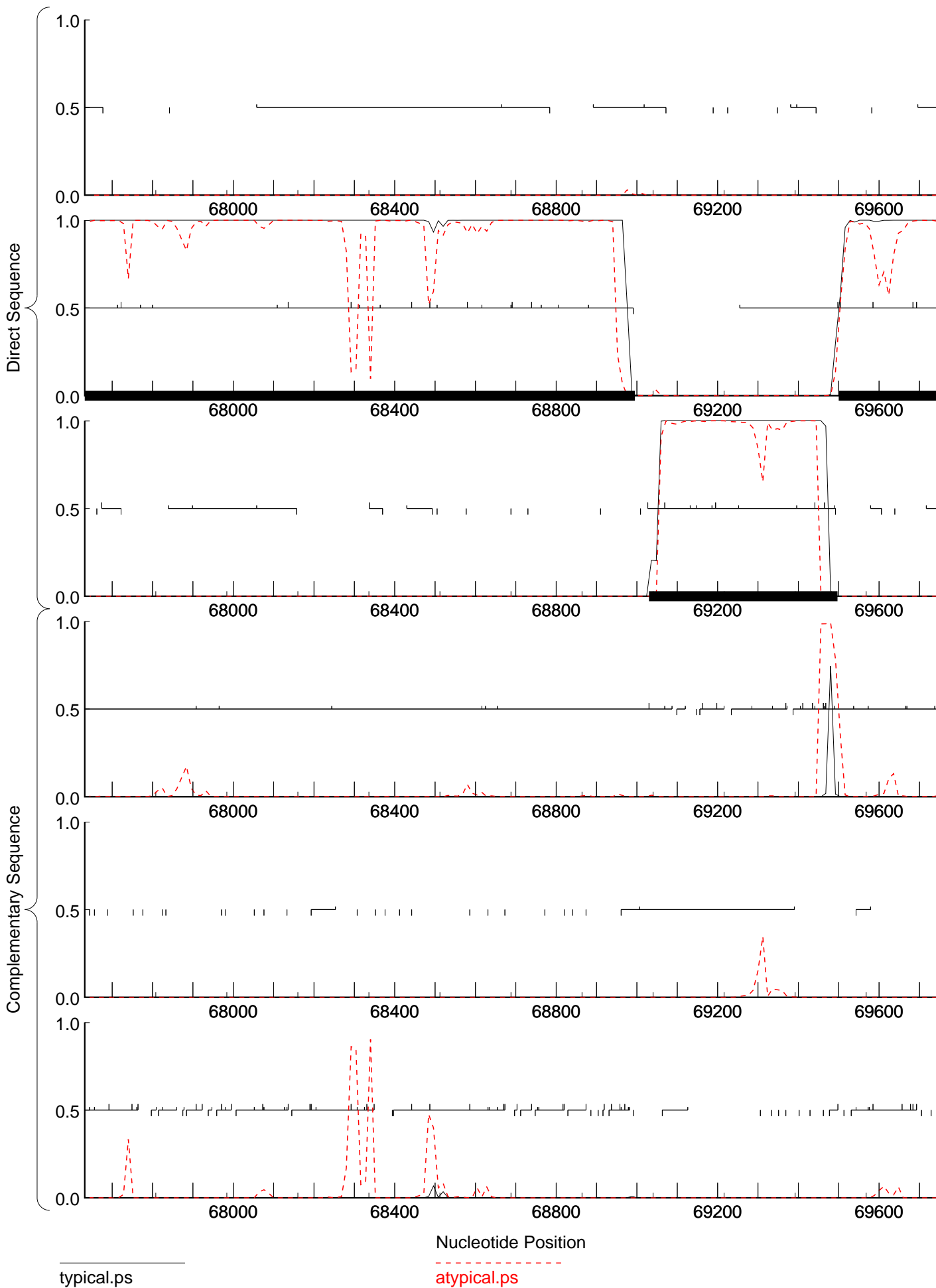


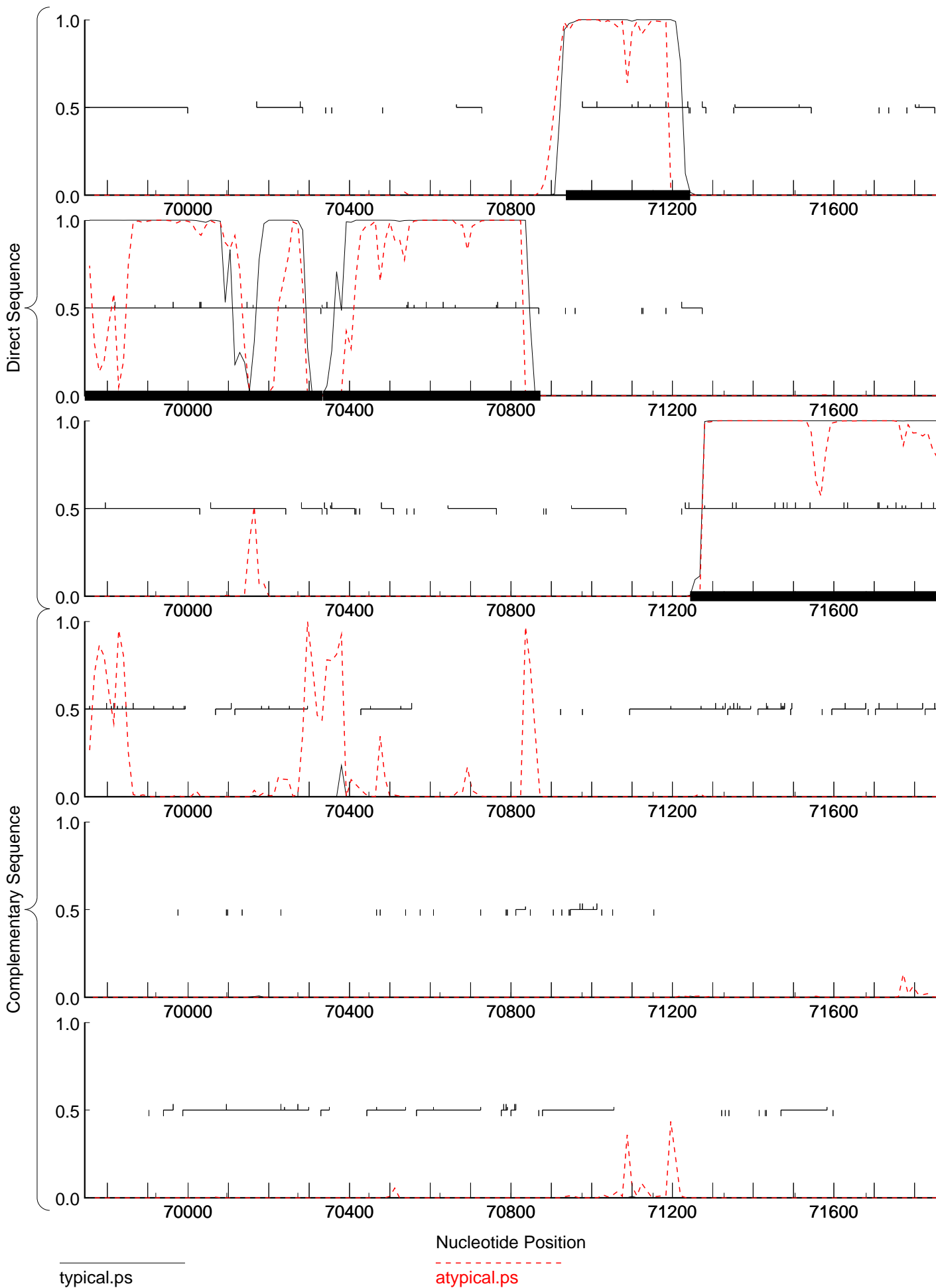


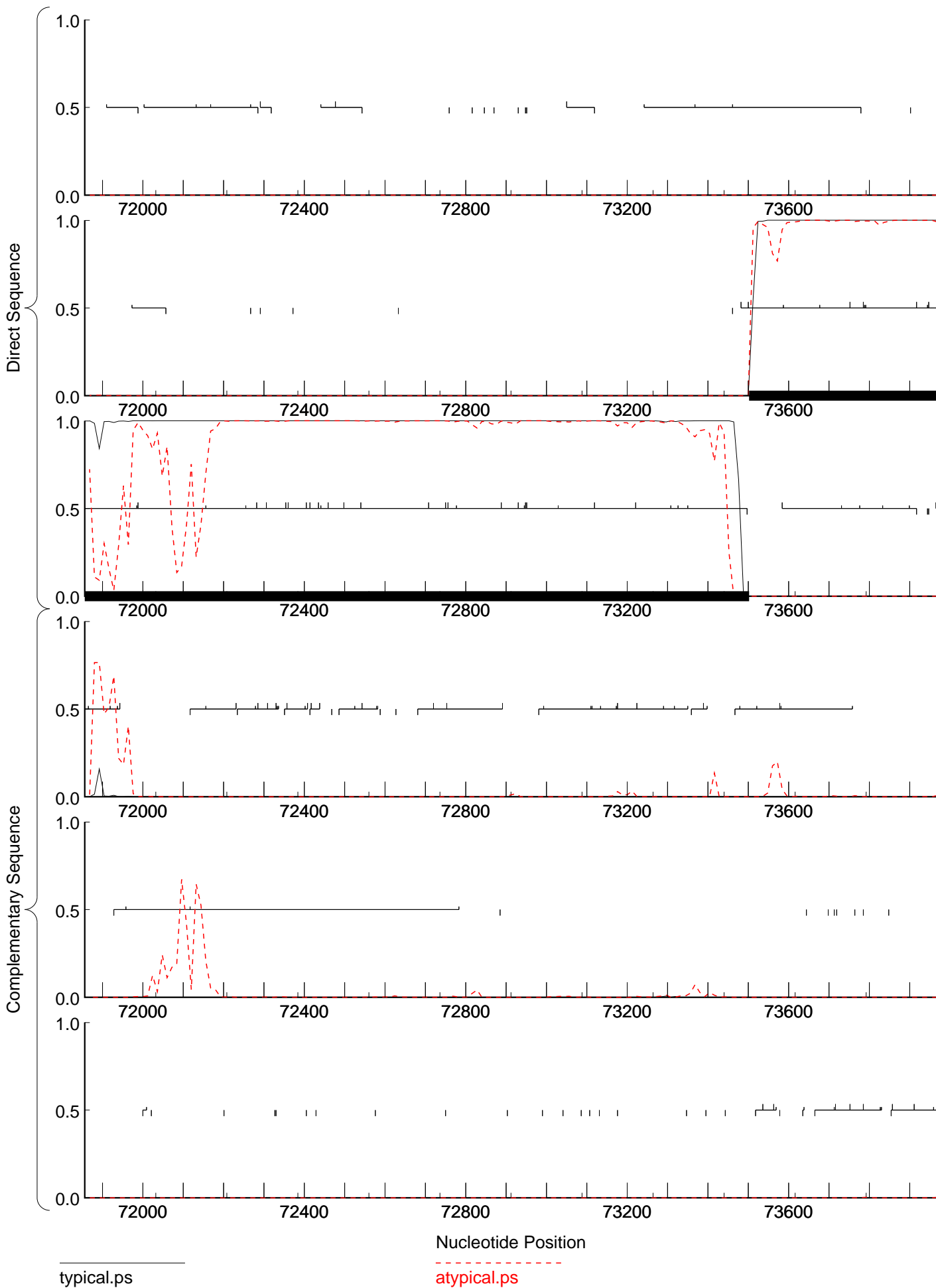






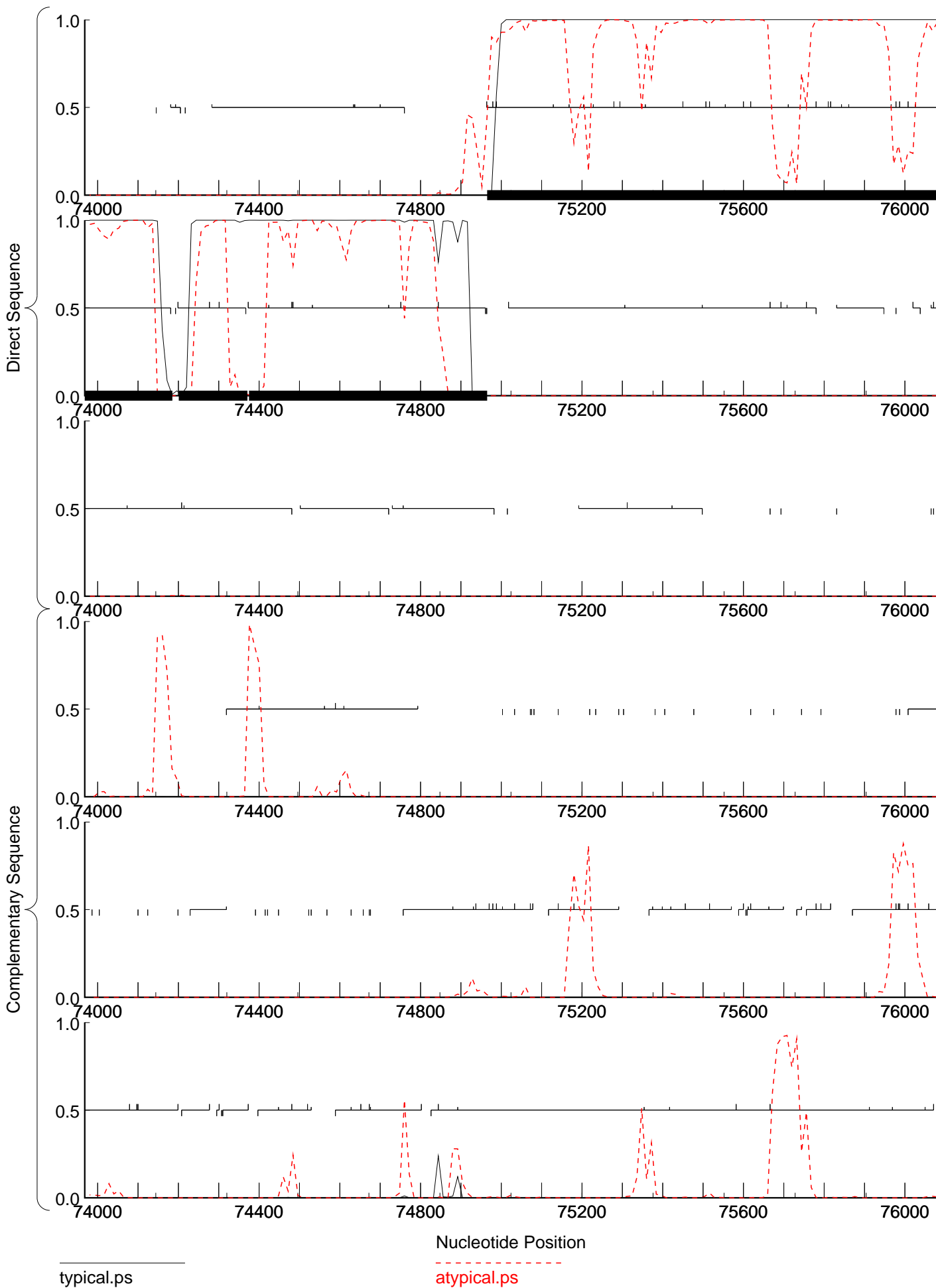


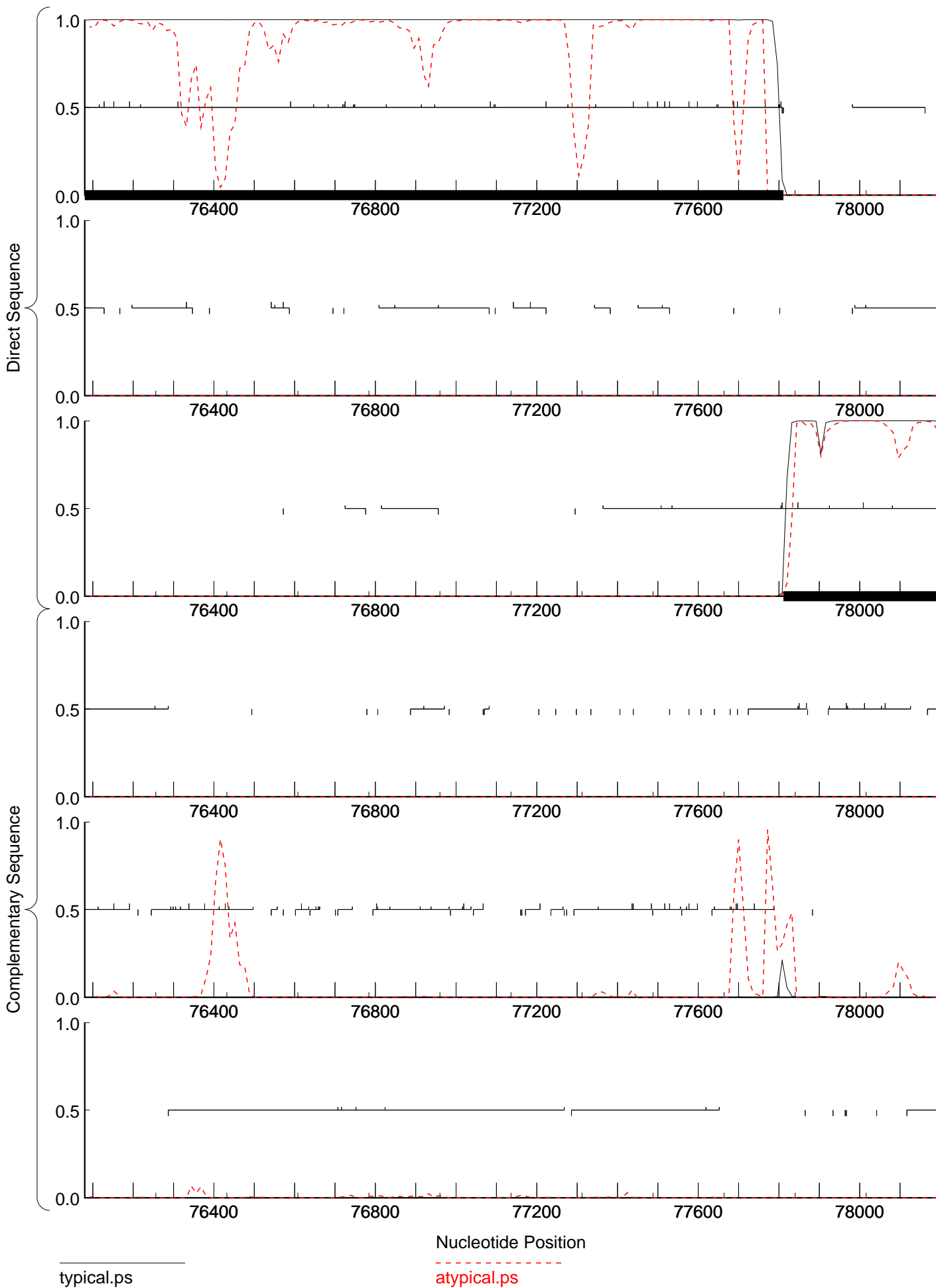




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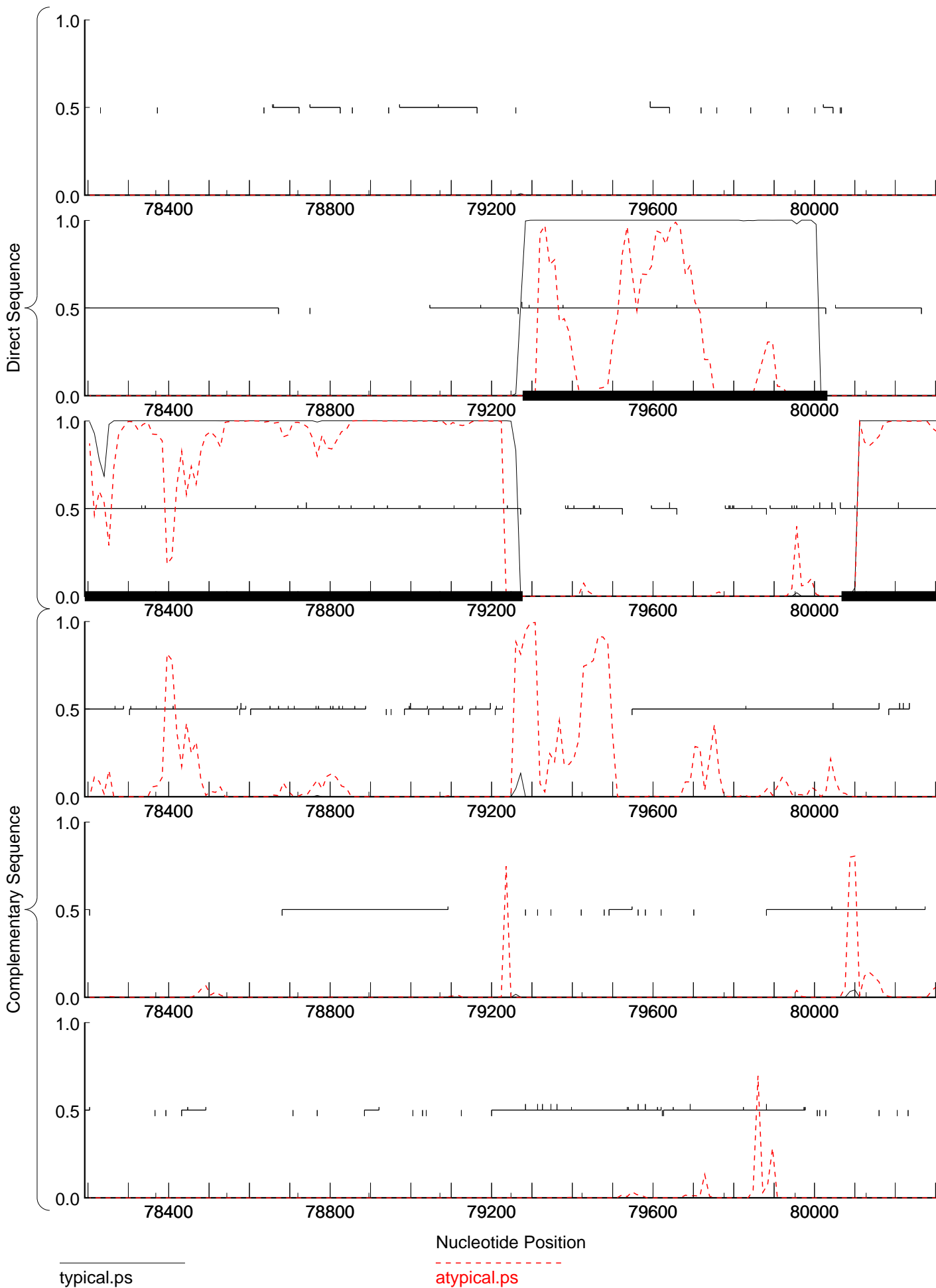
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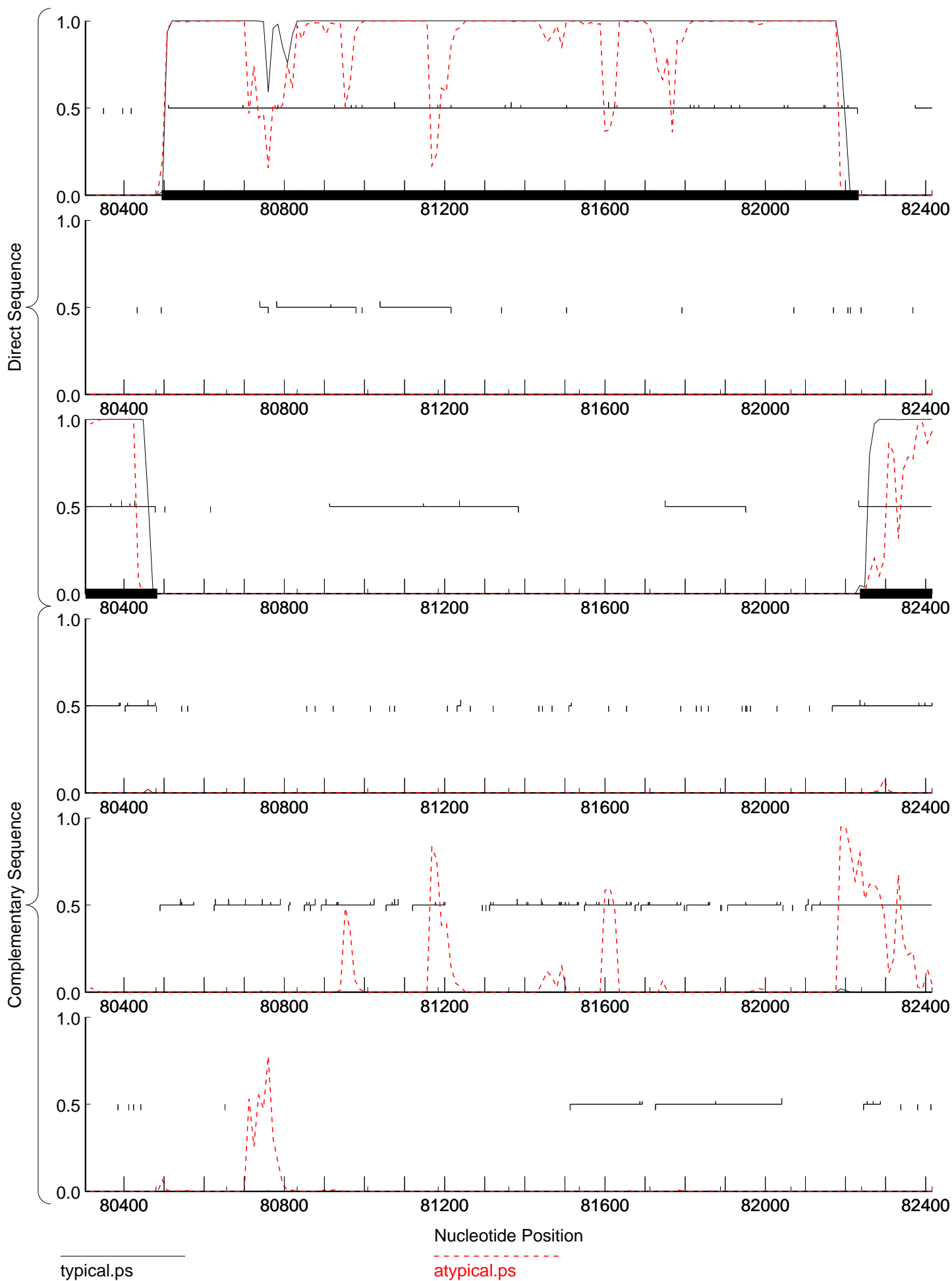




typical.ps

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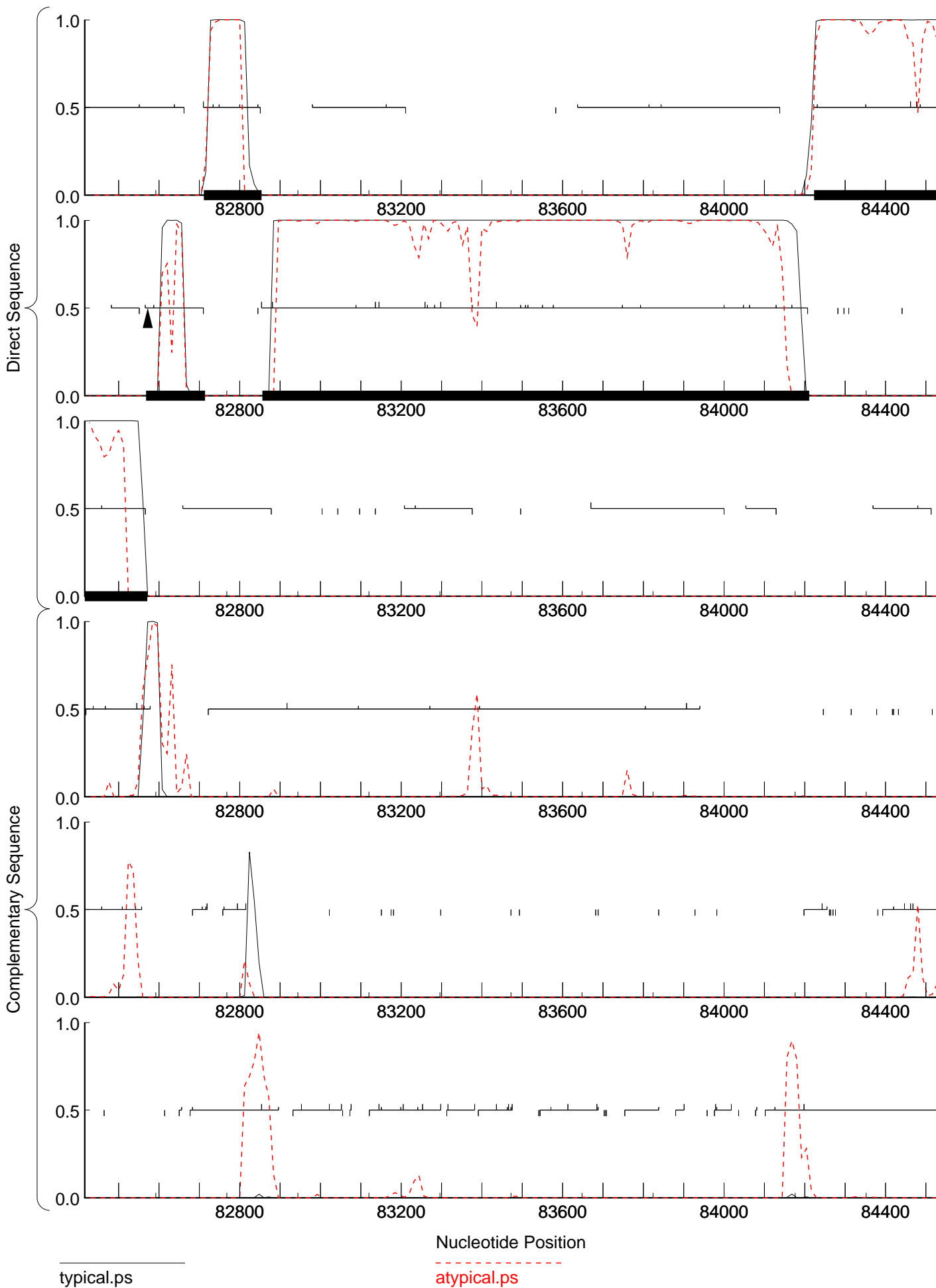




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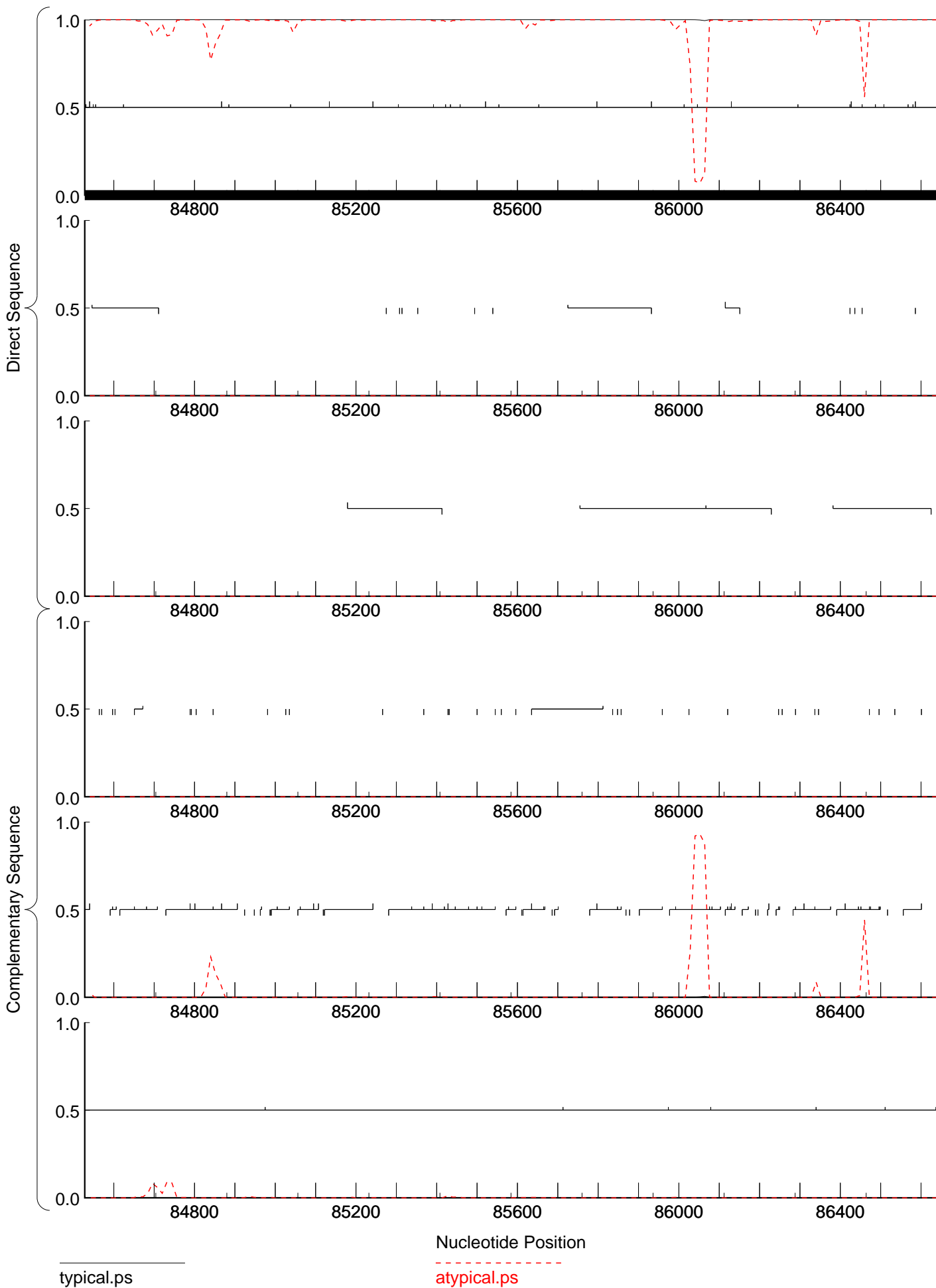
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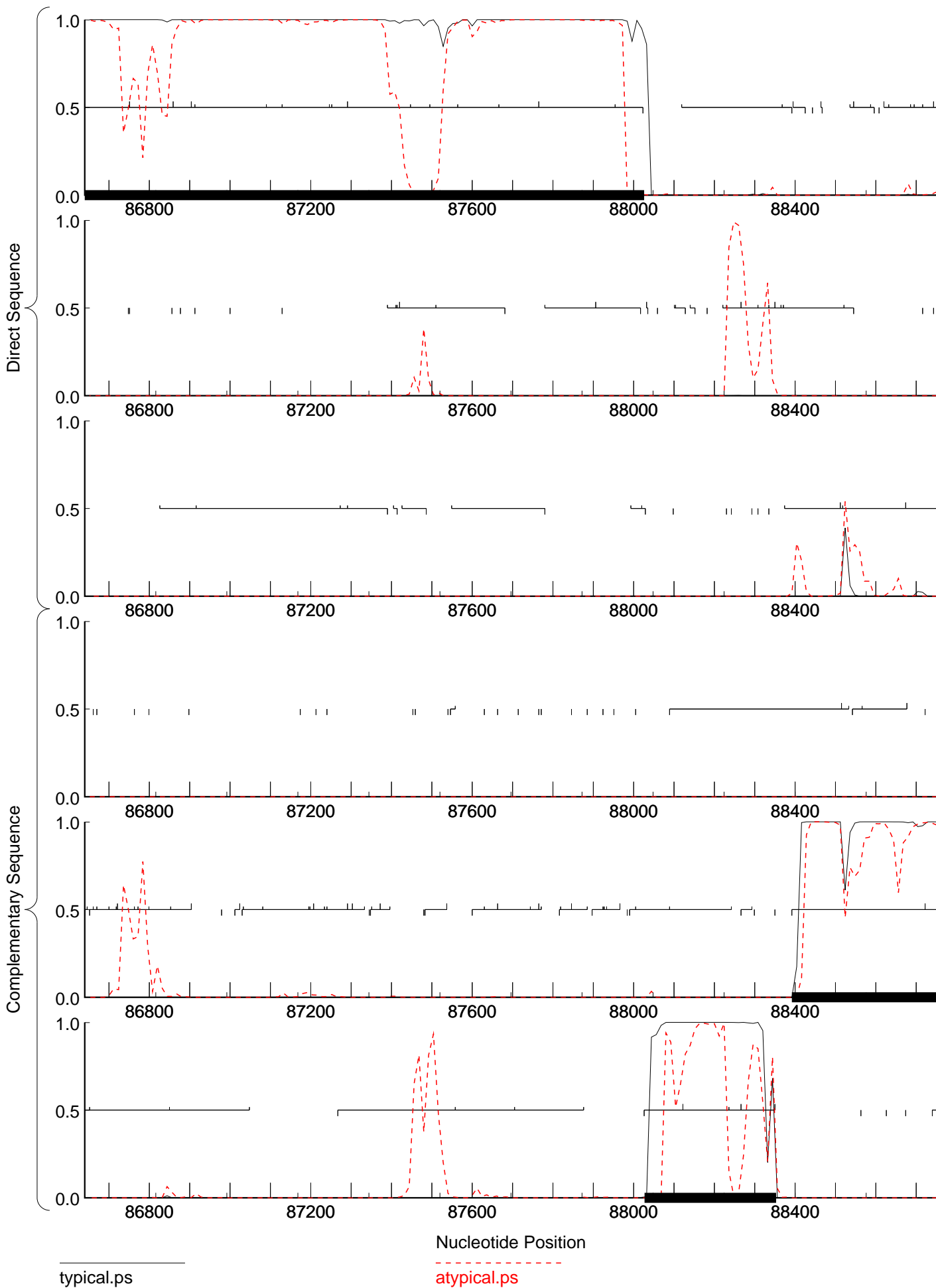


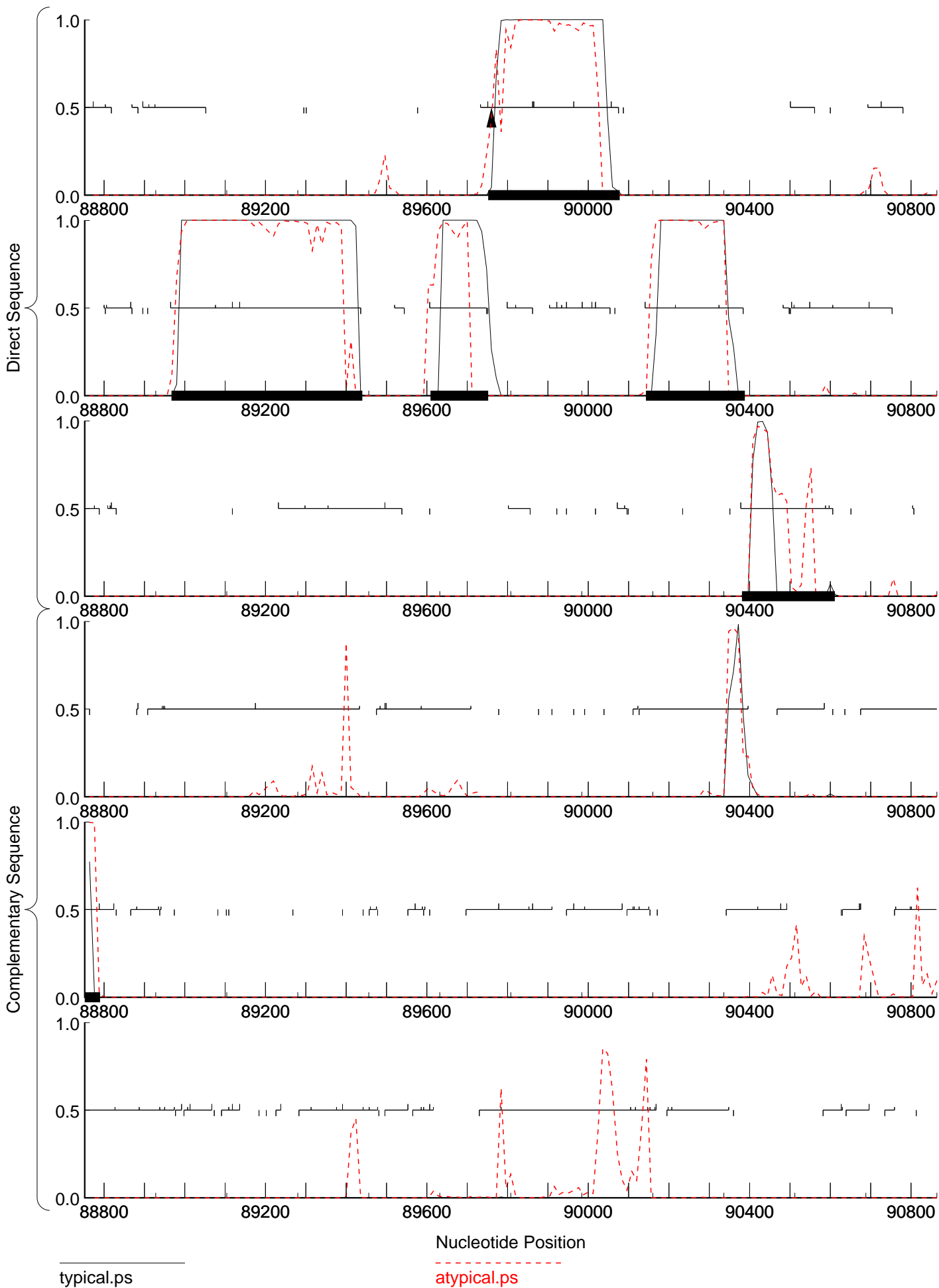


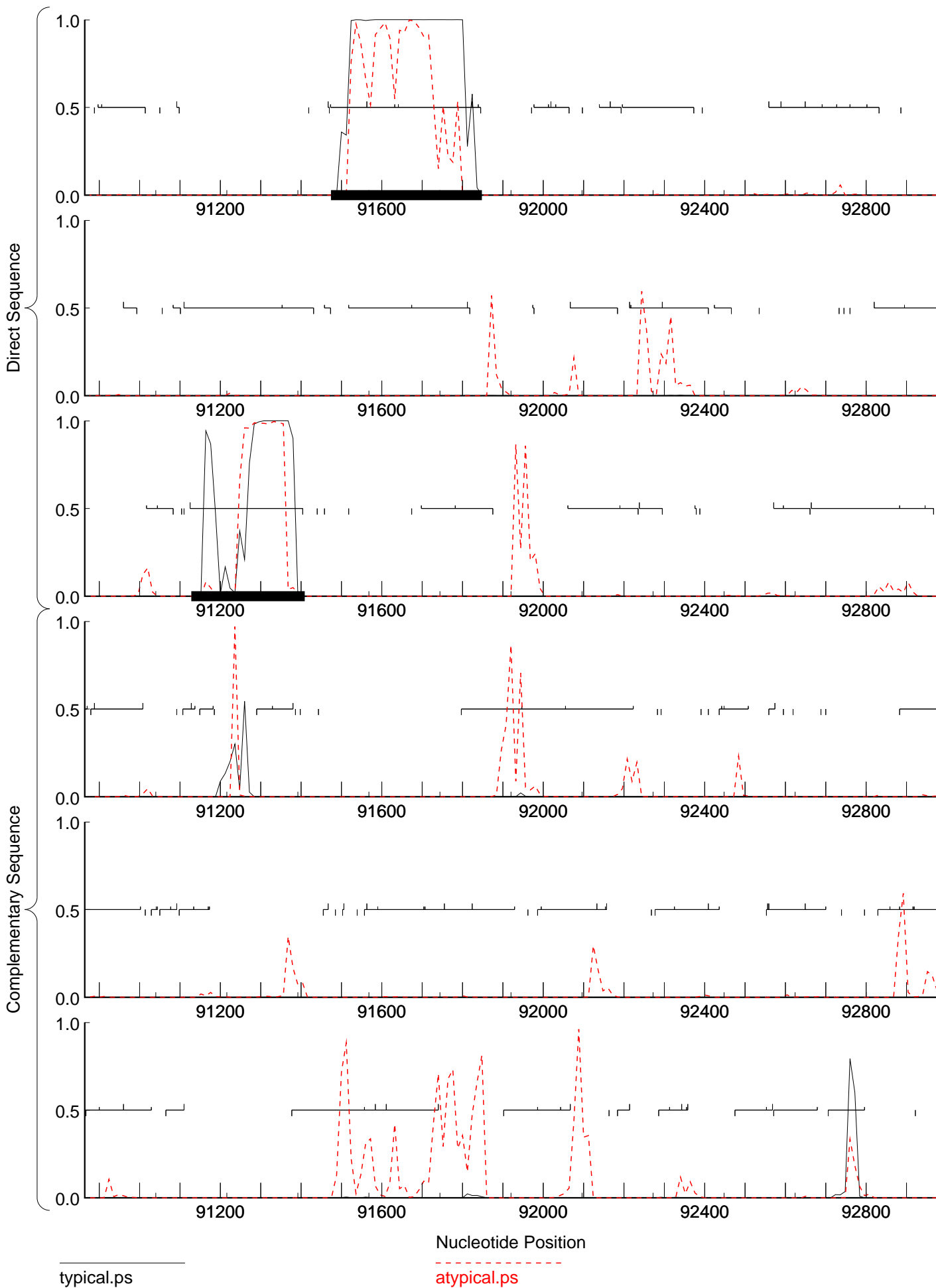
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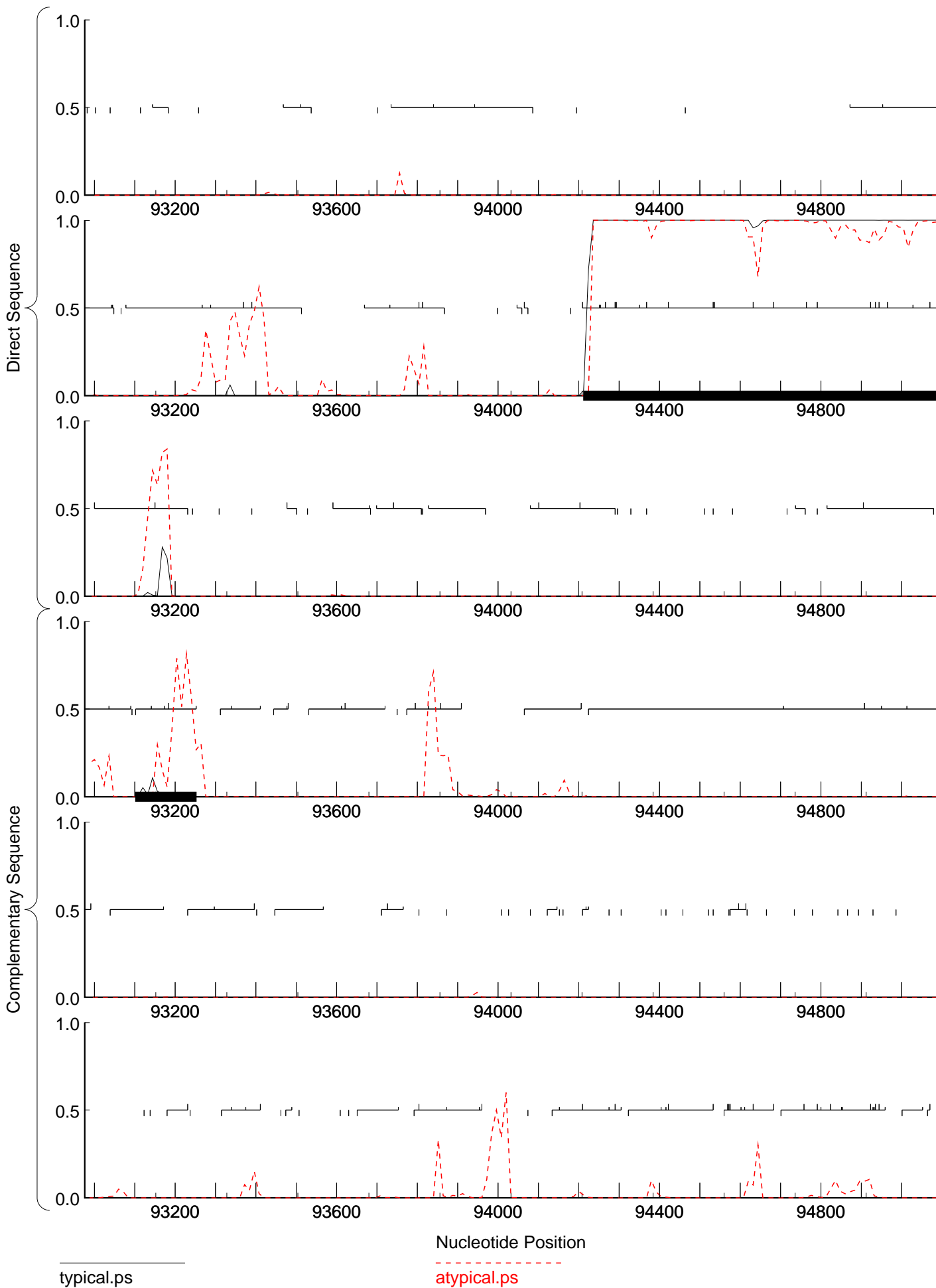


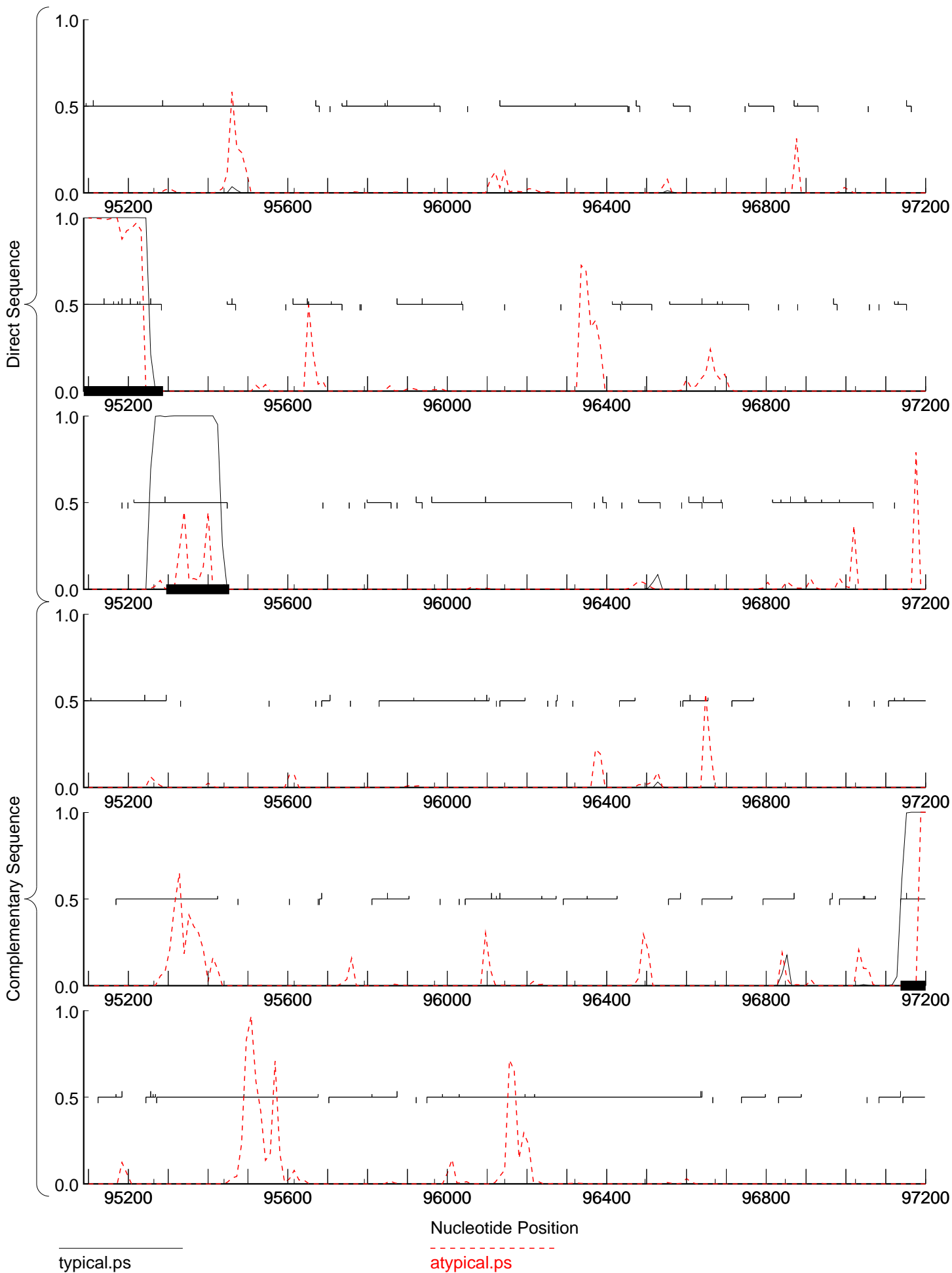


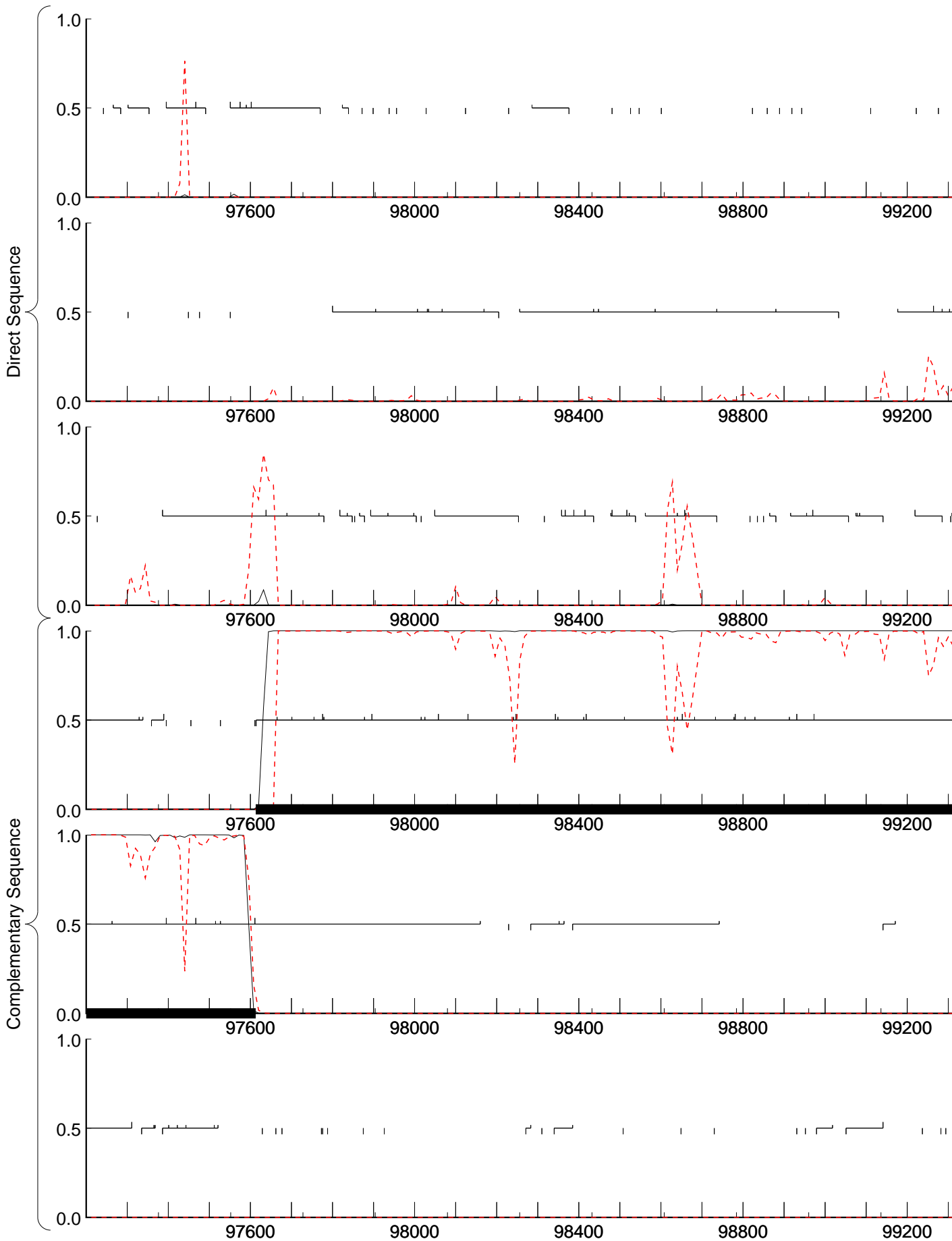


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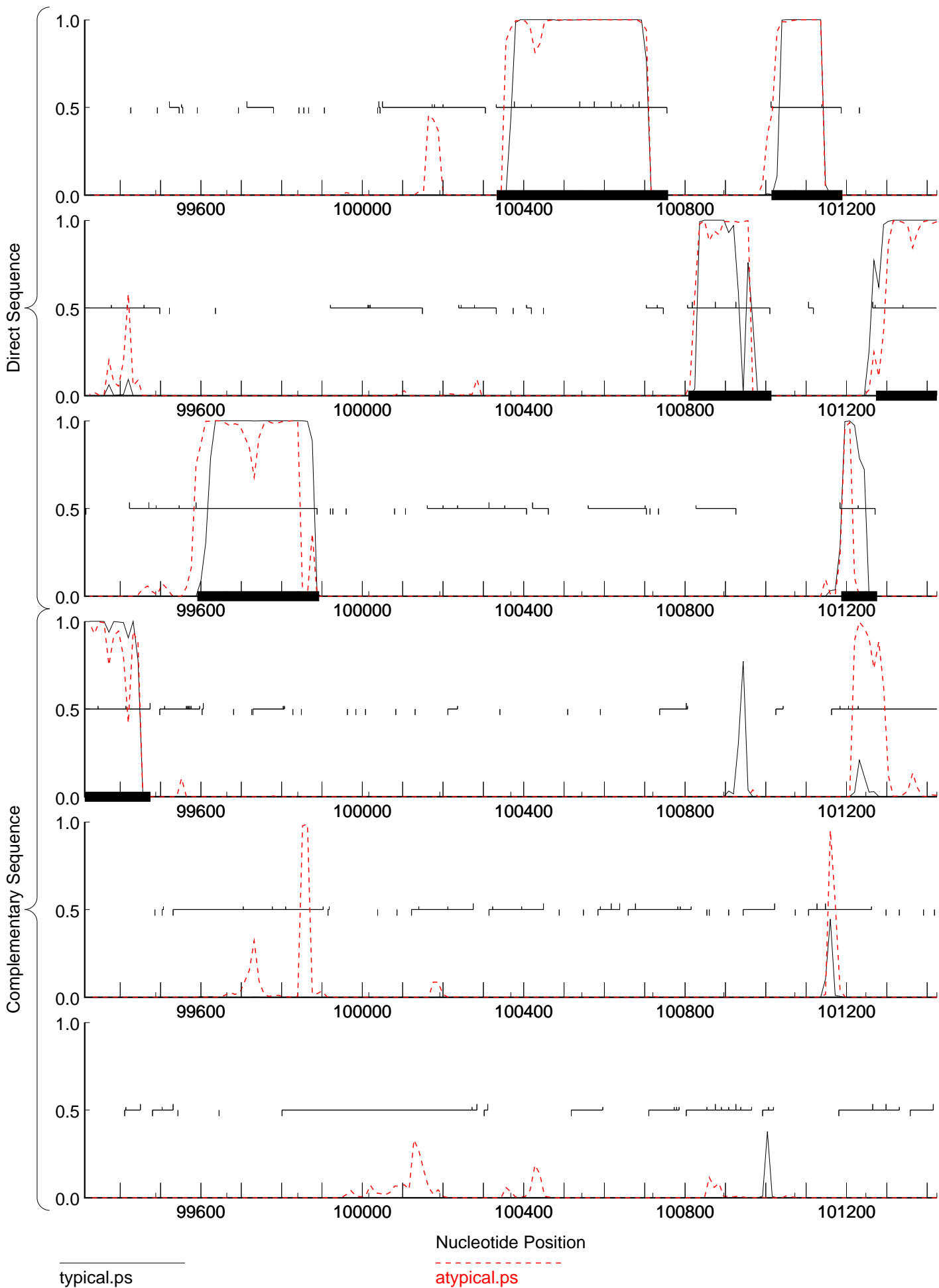
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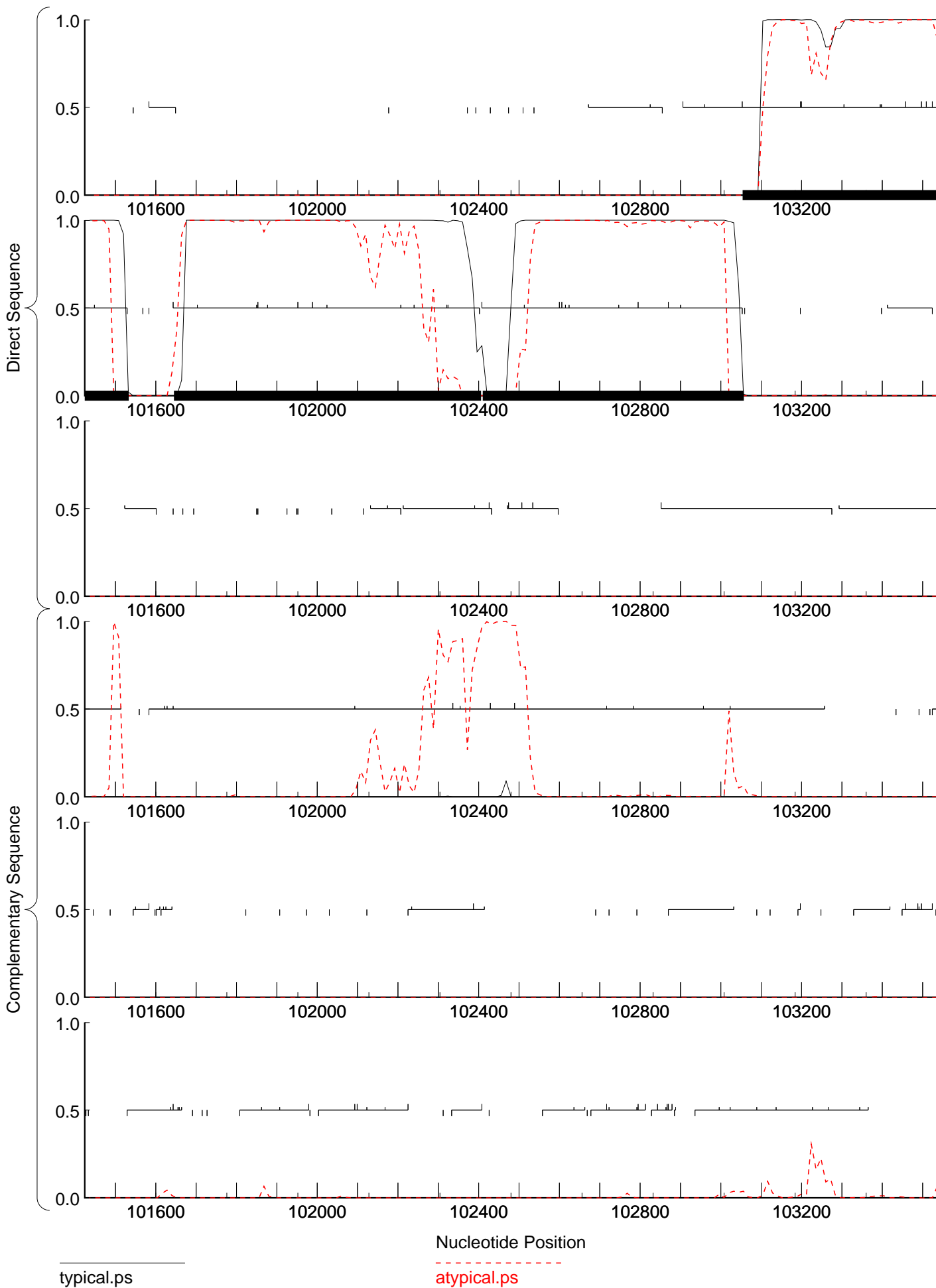


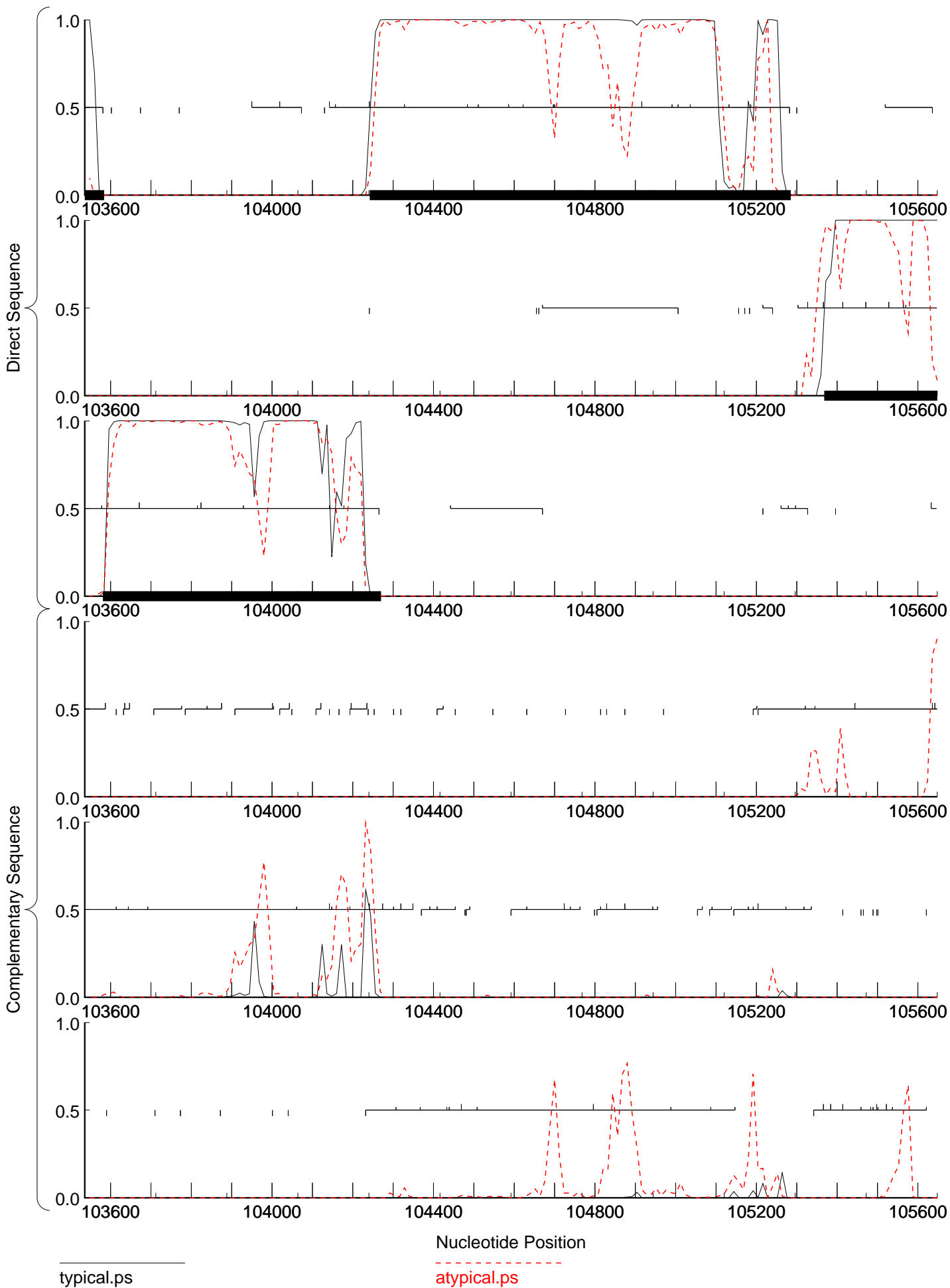






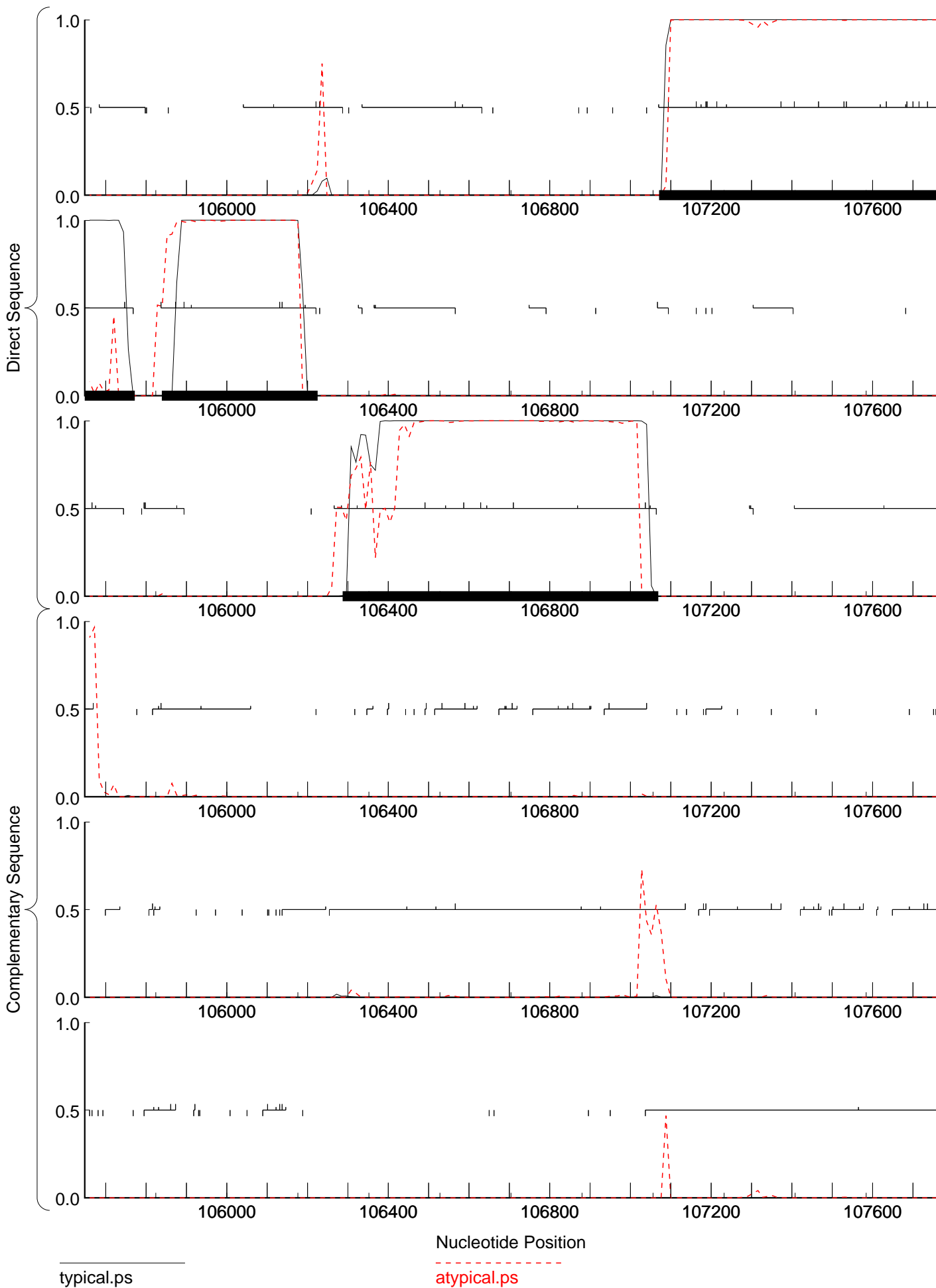


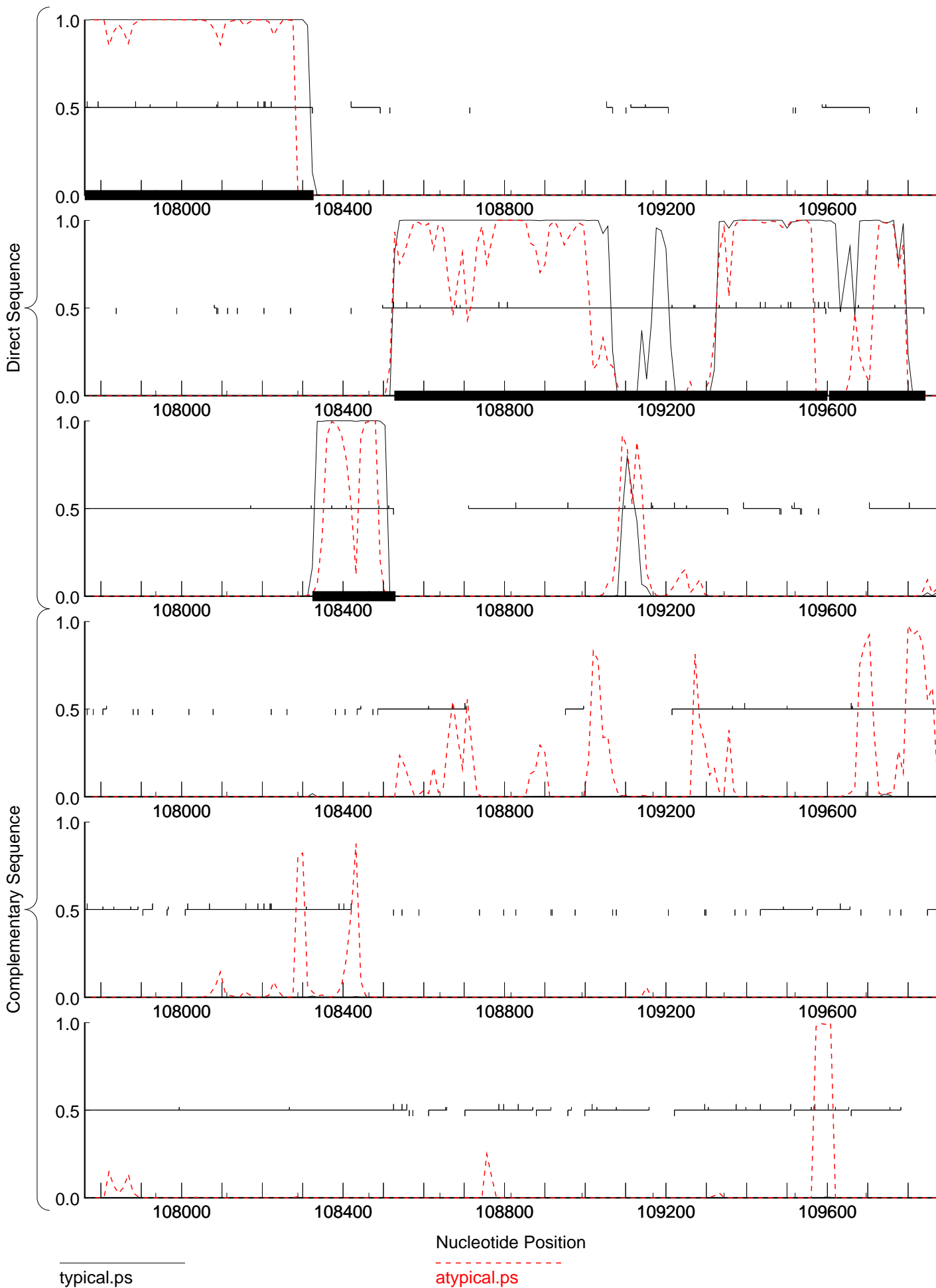




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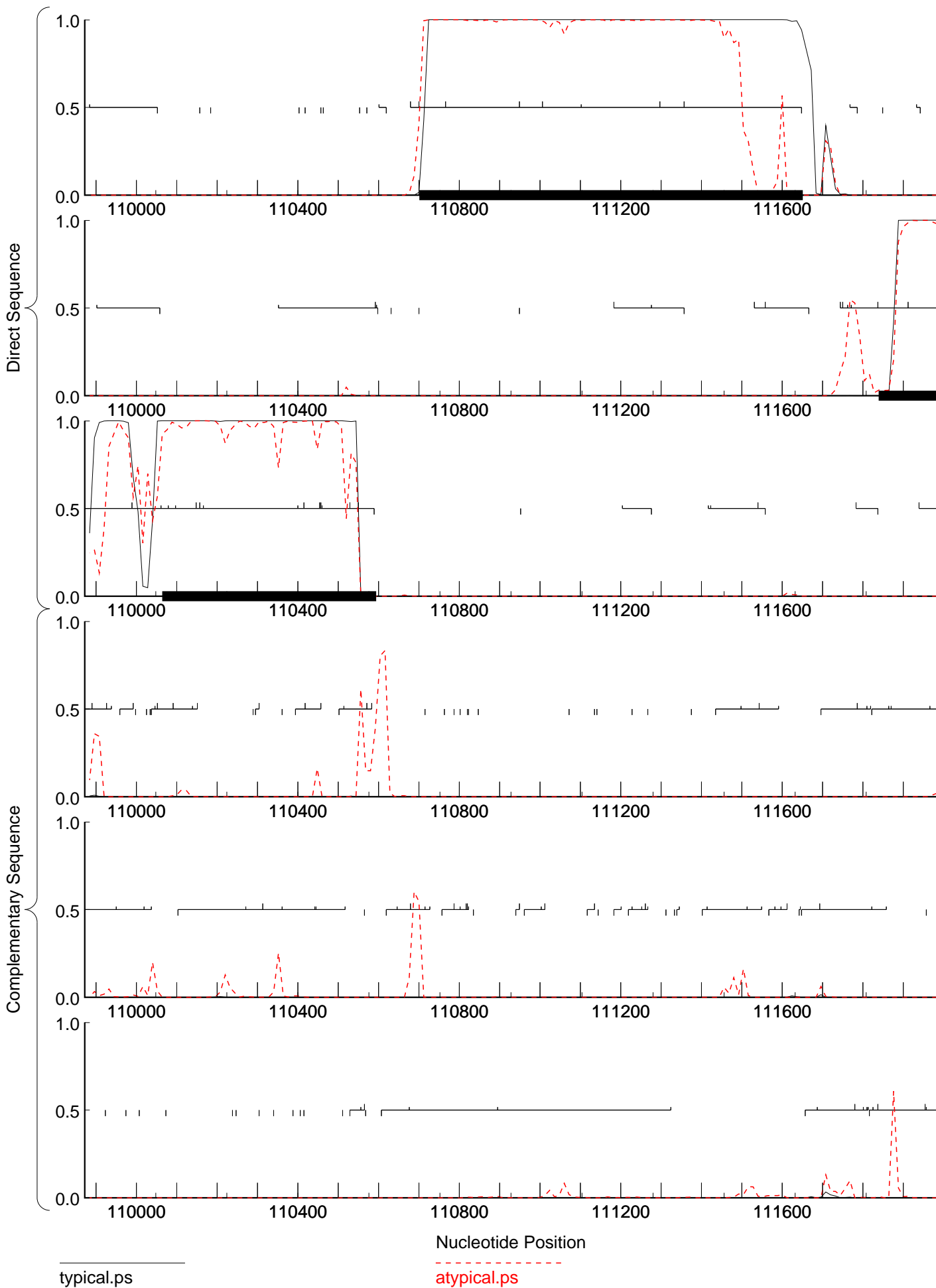
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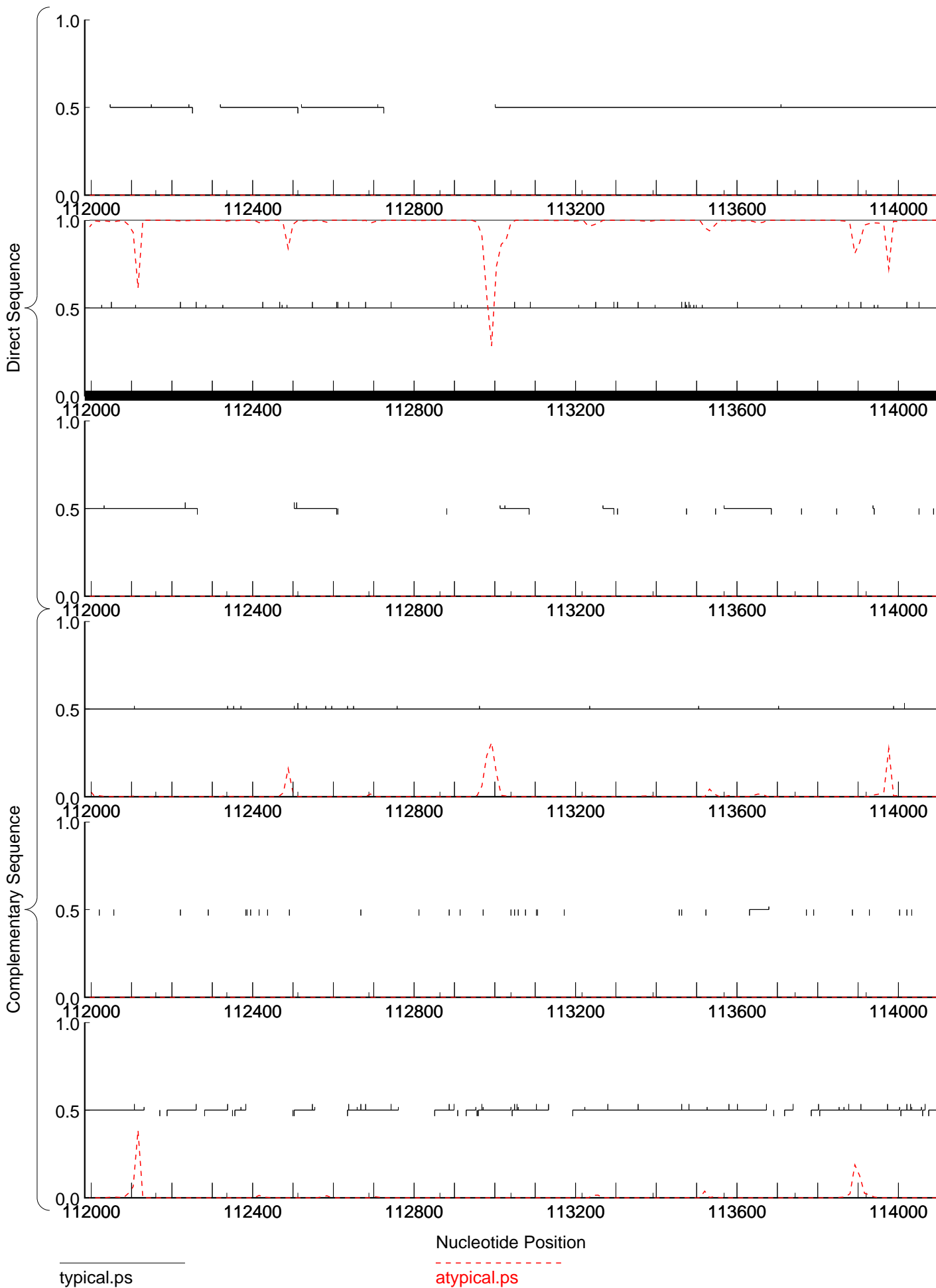


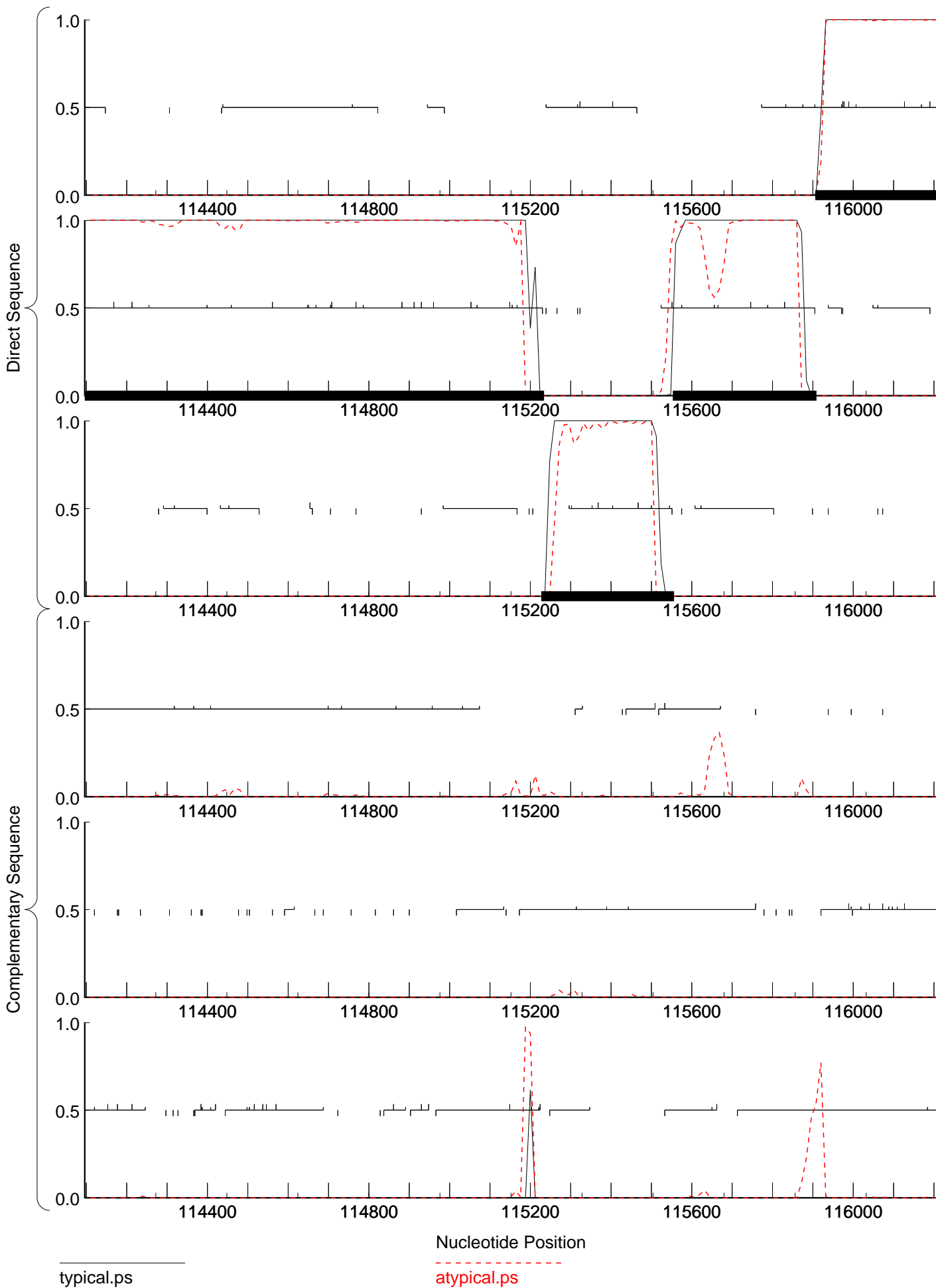


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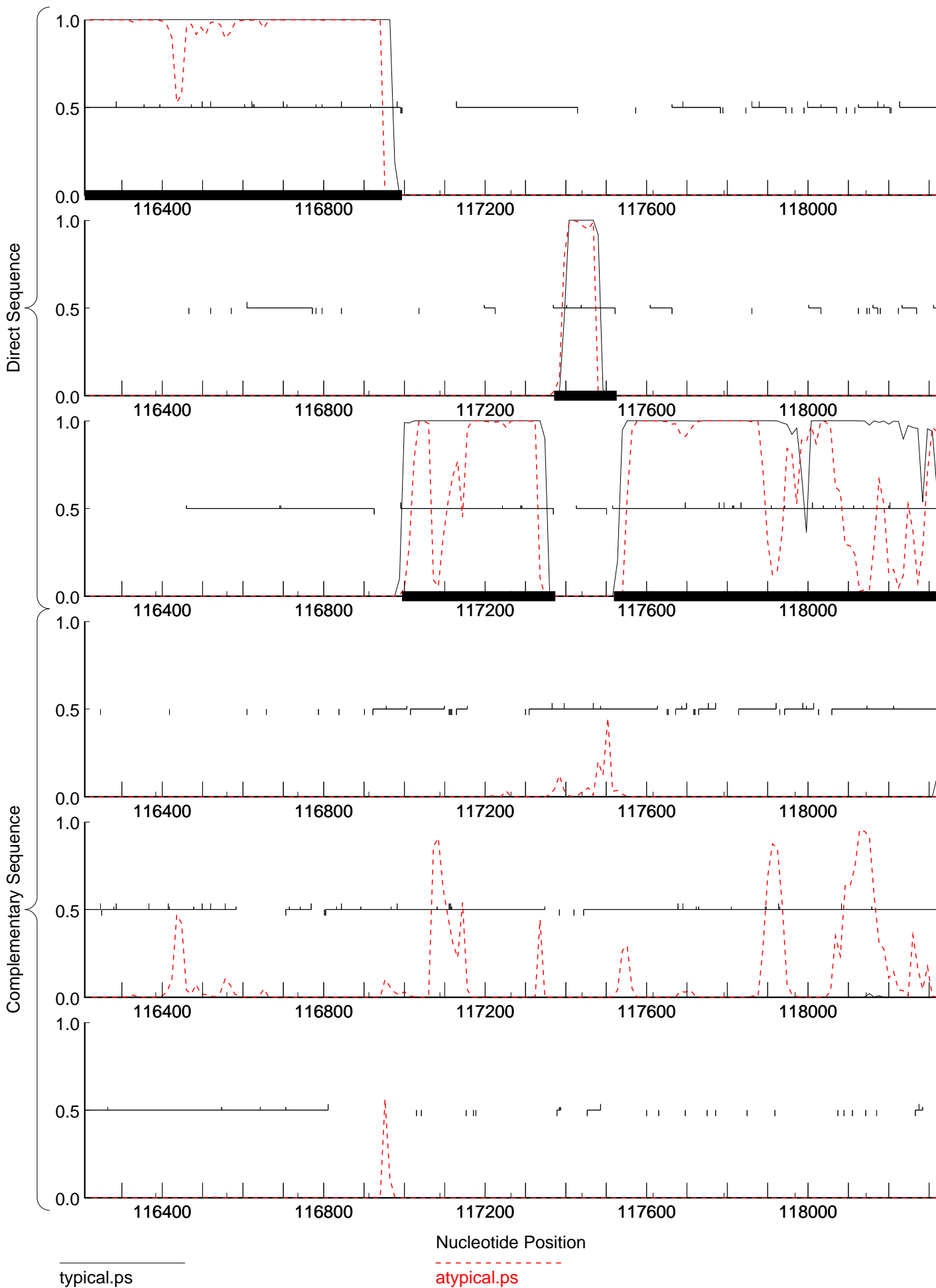
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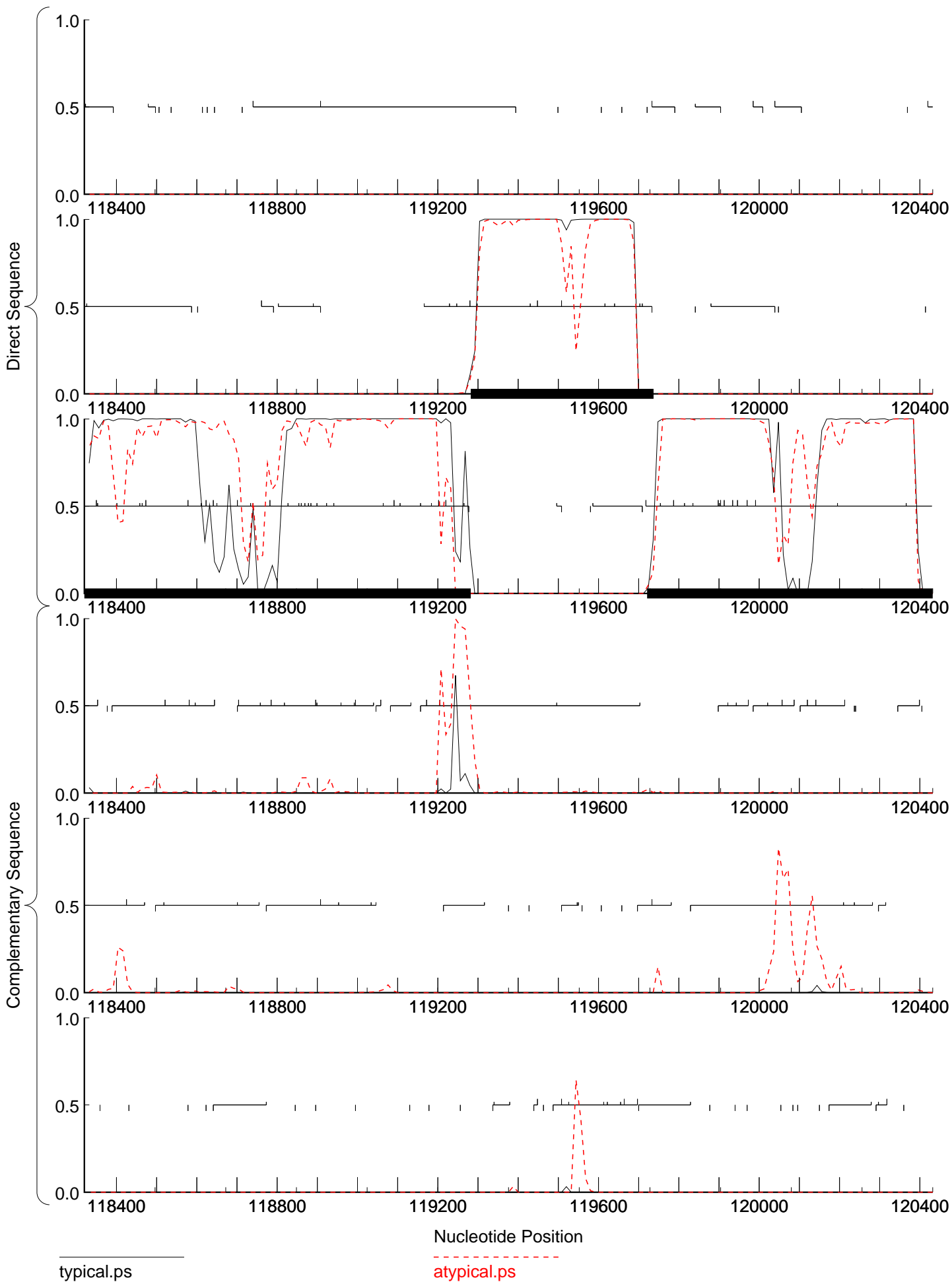


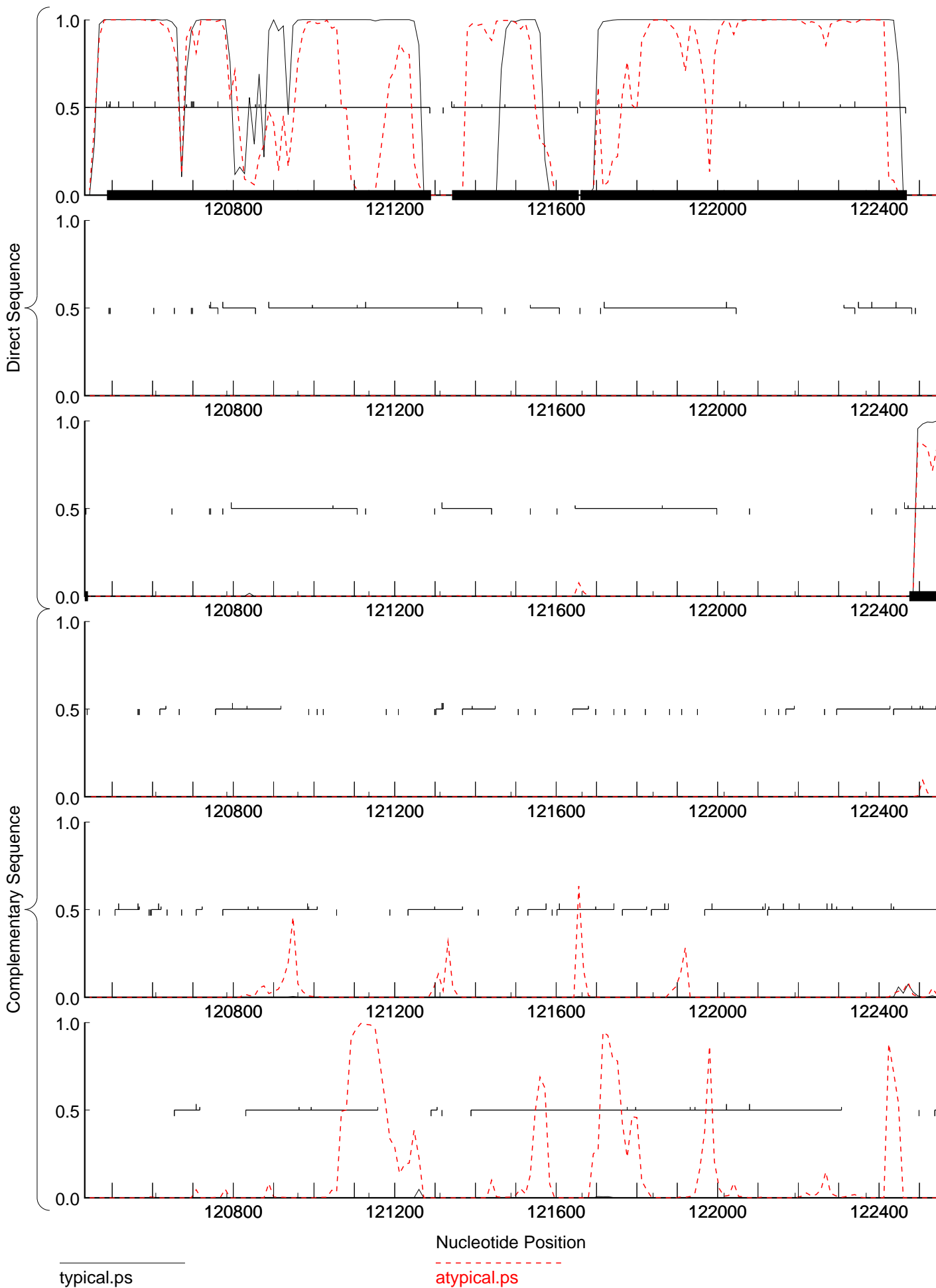


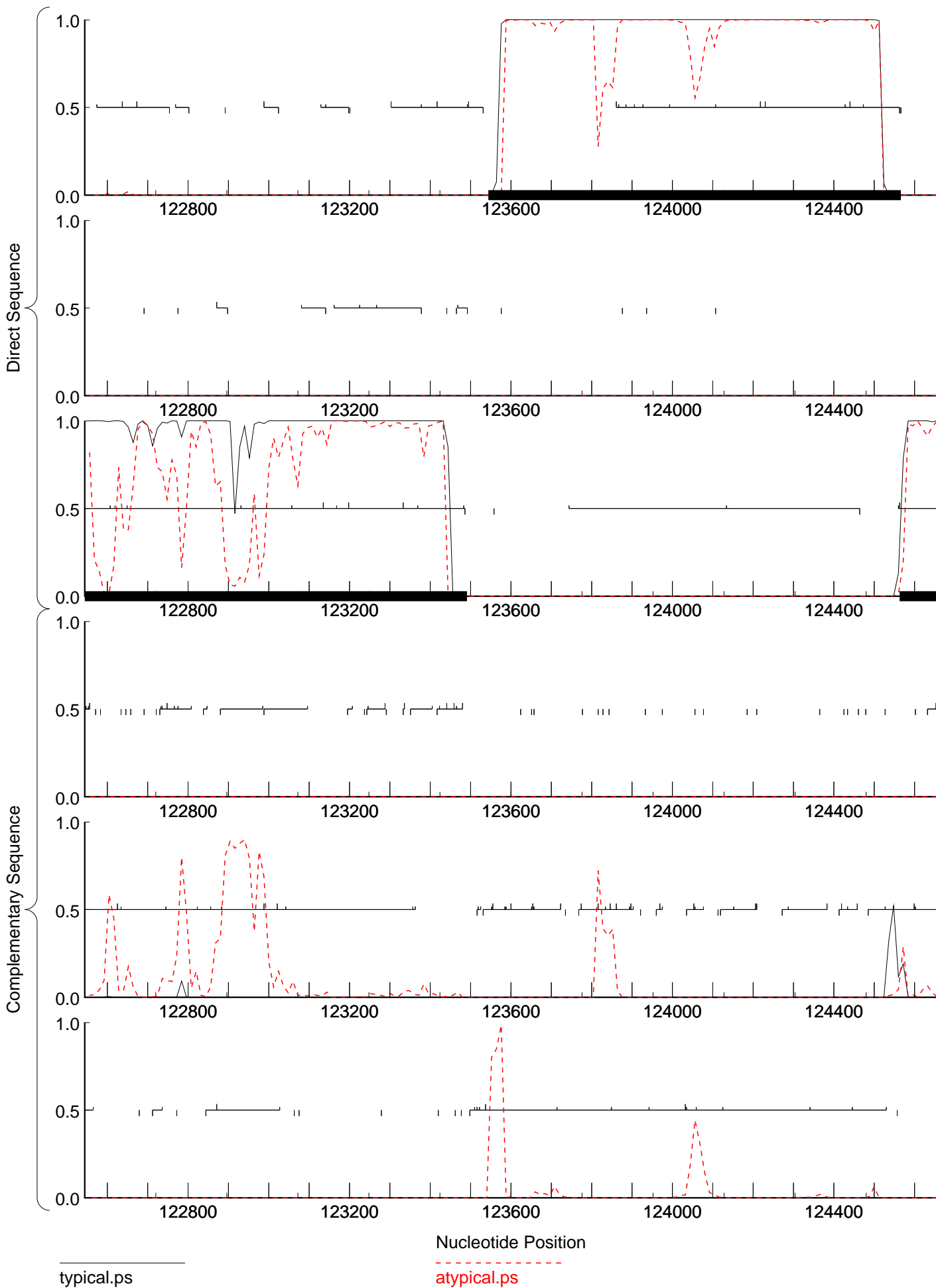


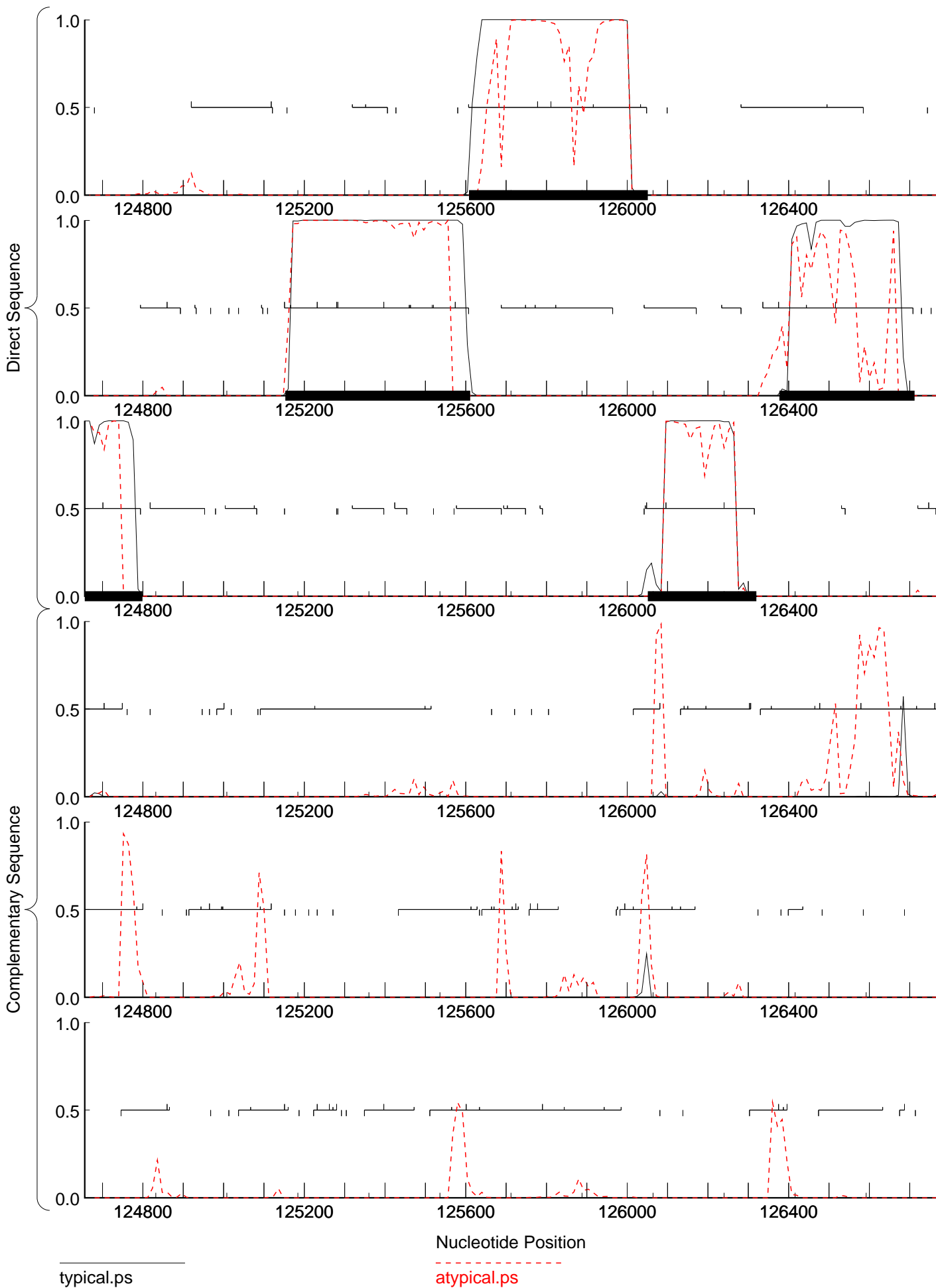


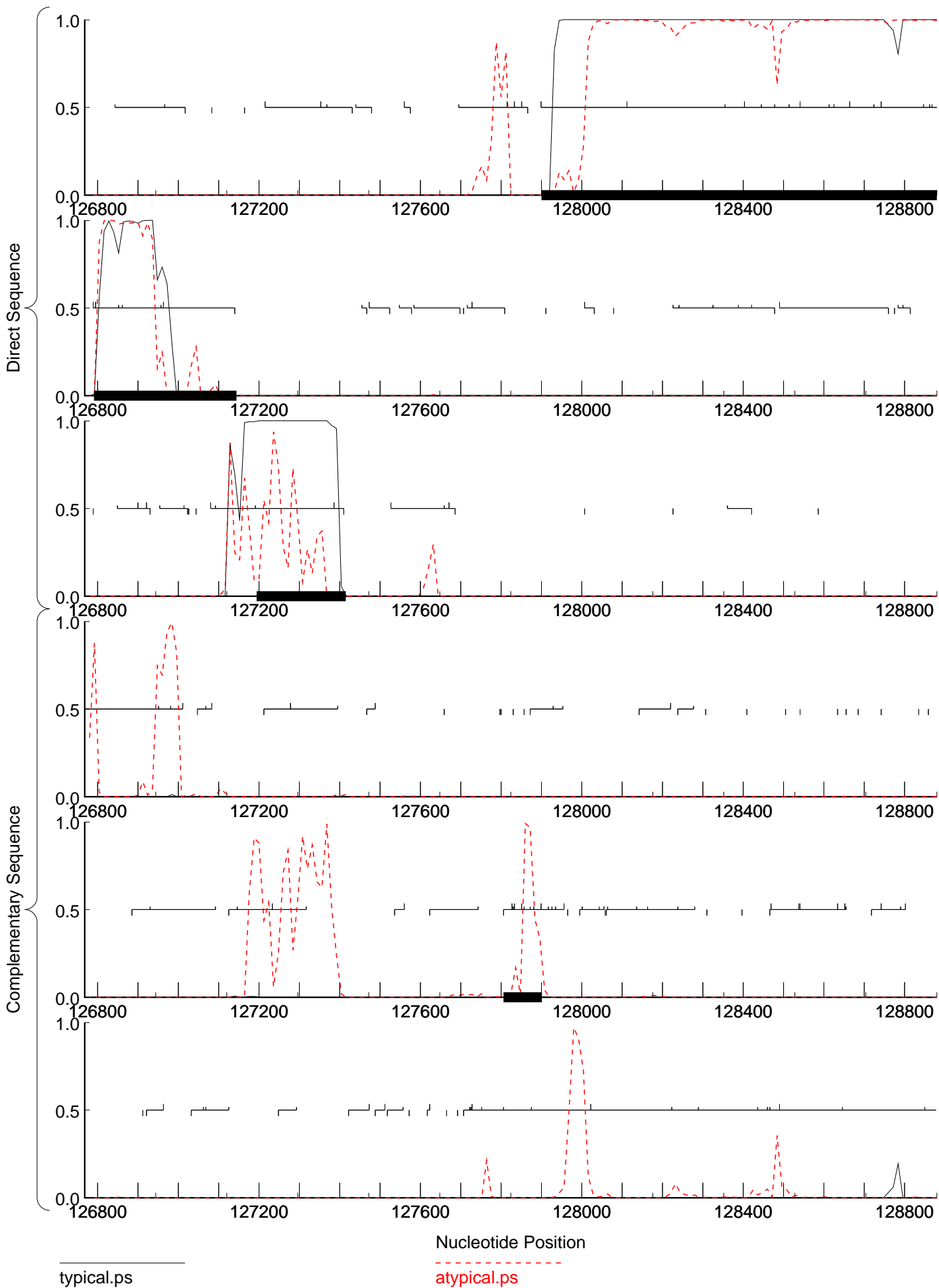


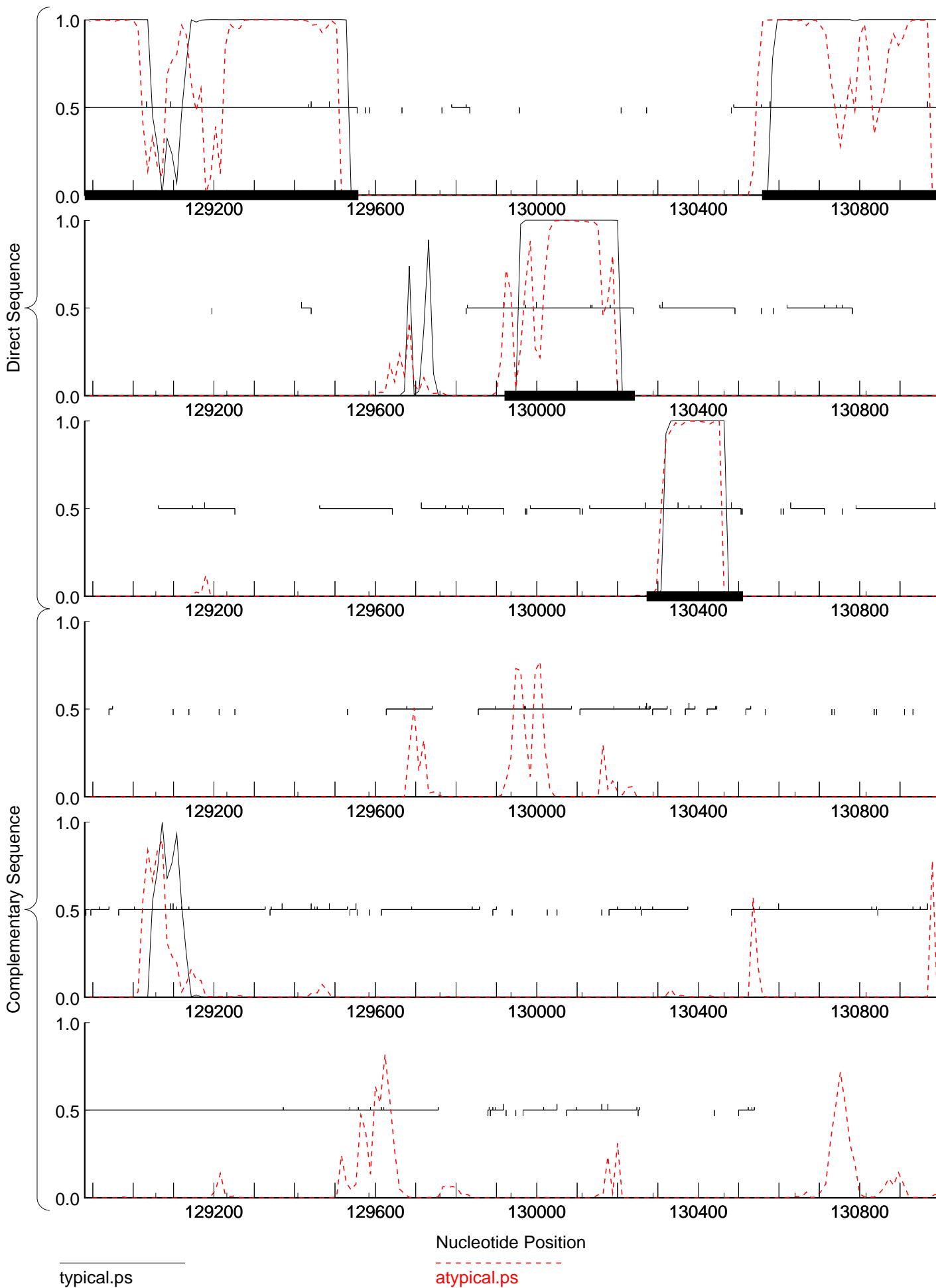






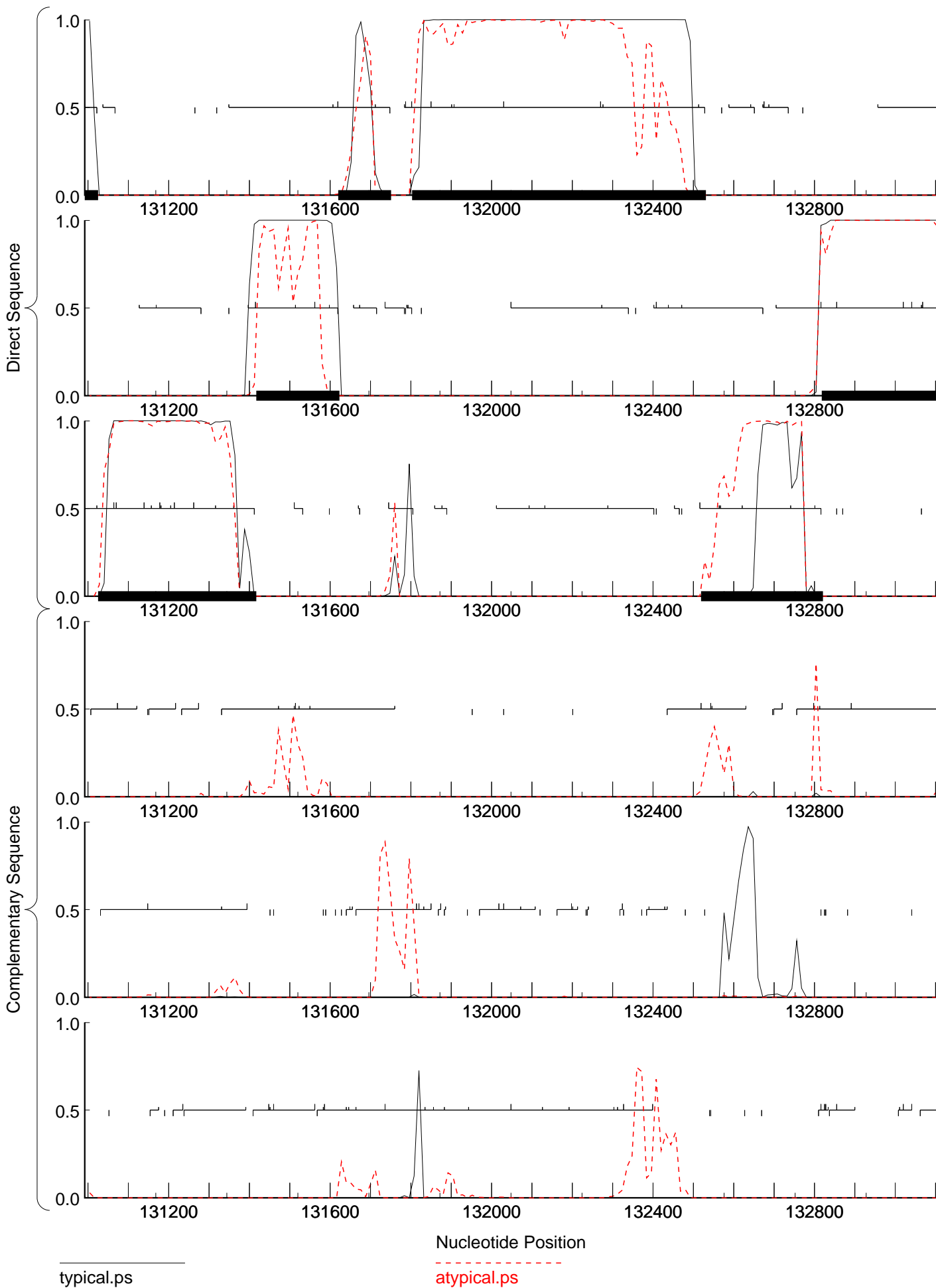






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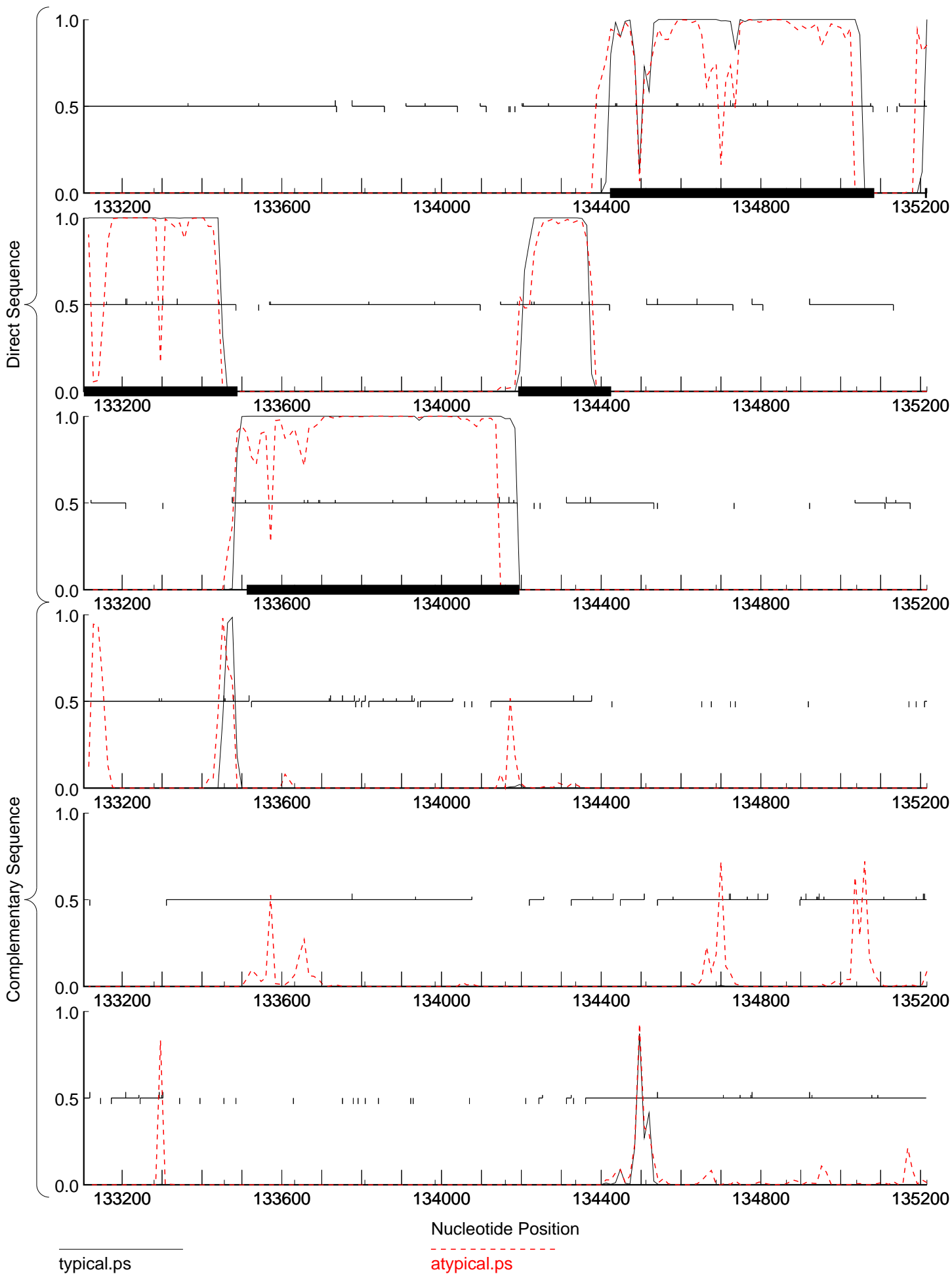
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typical.ps

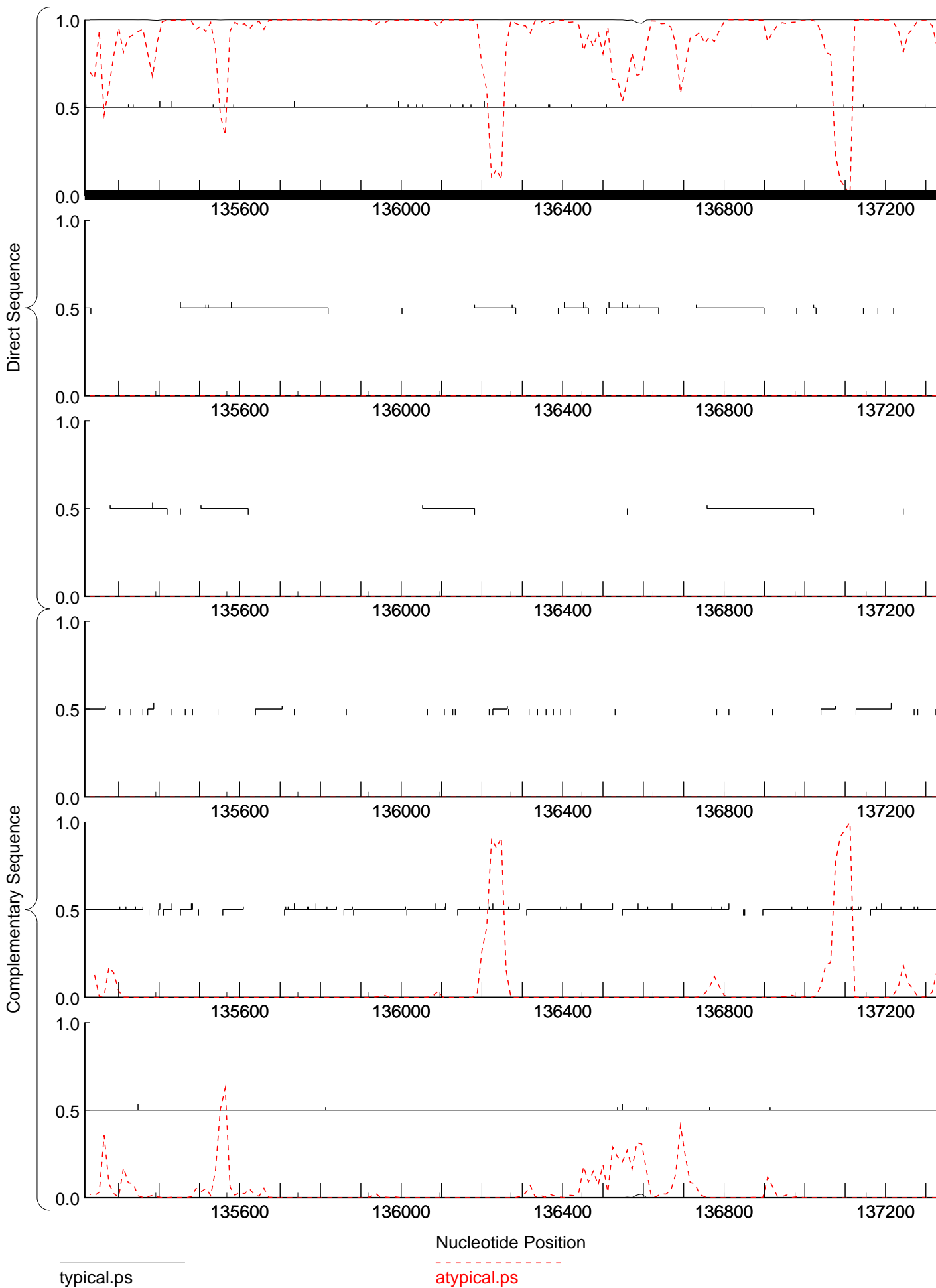
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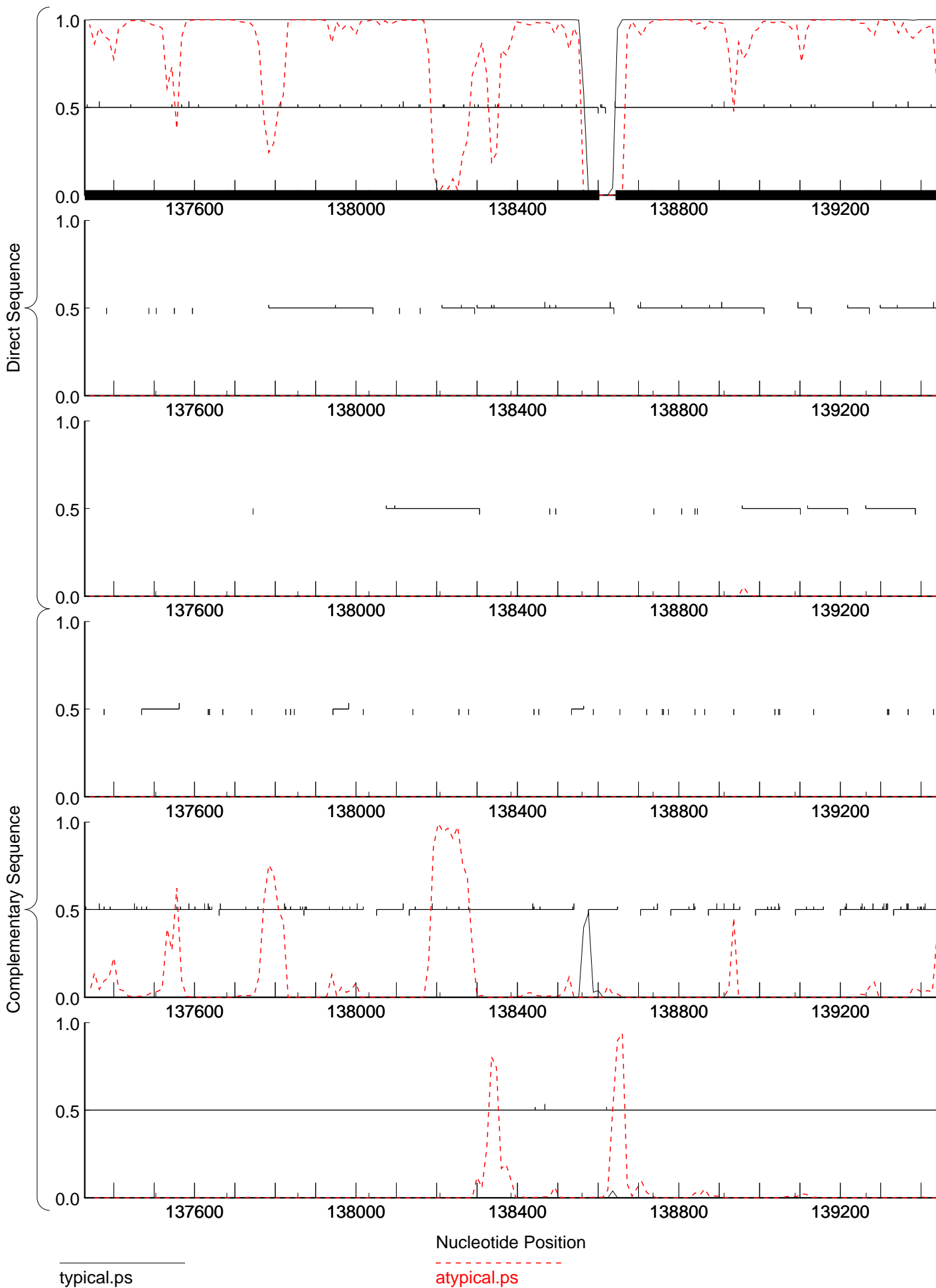




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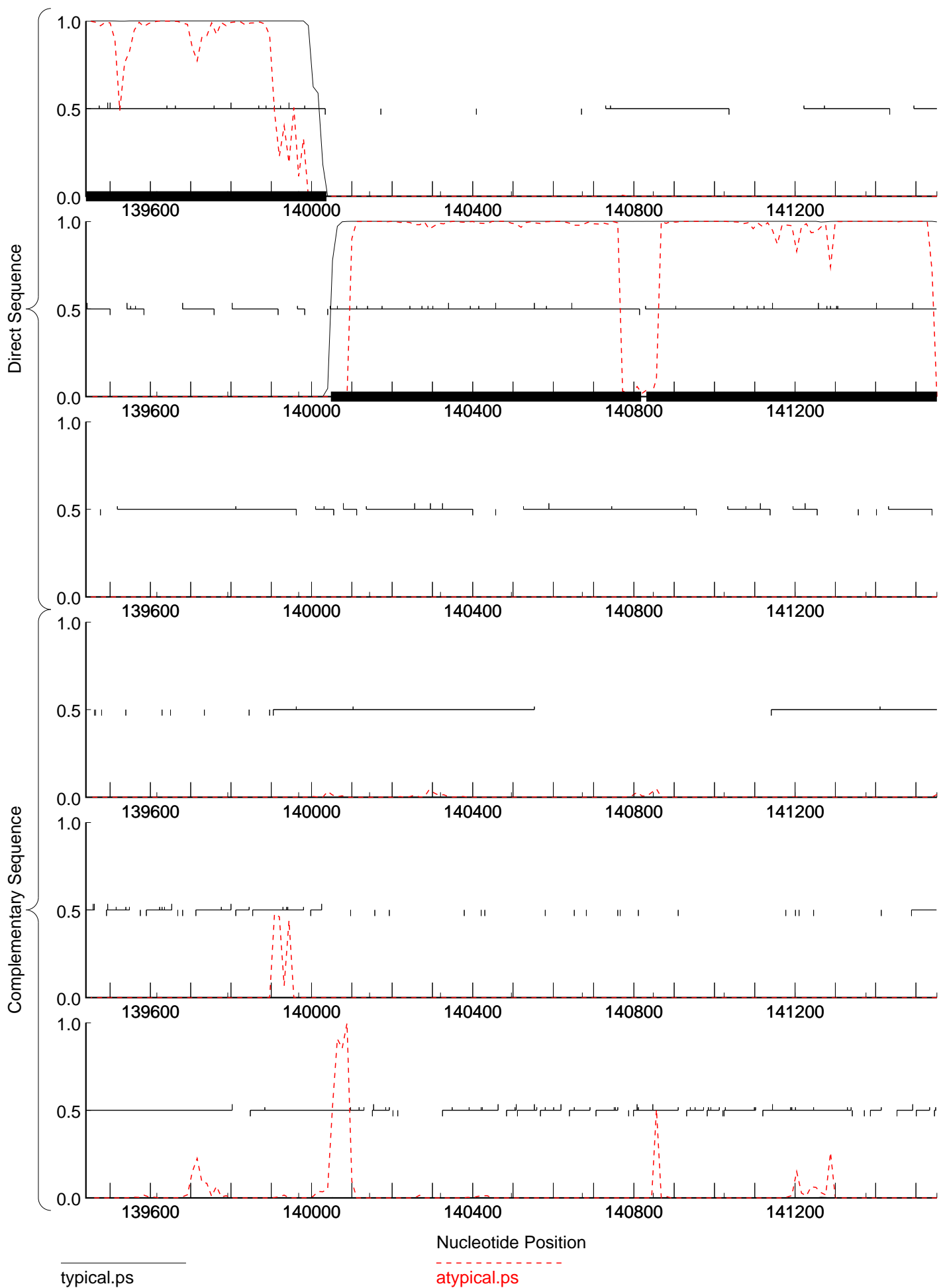
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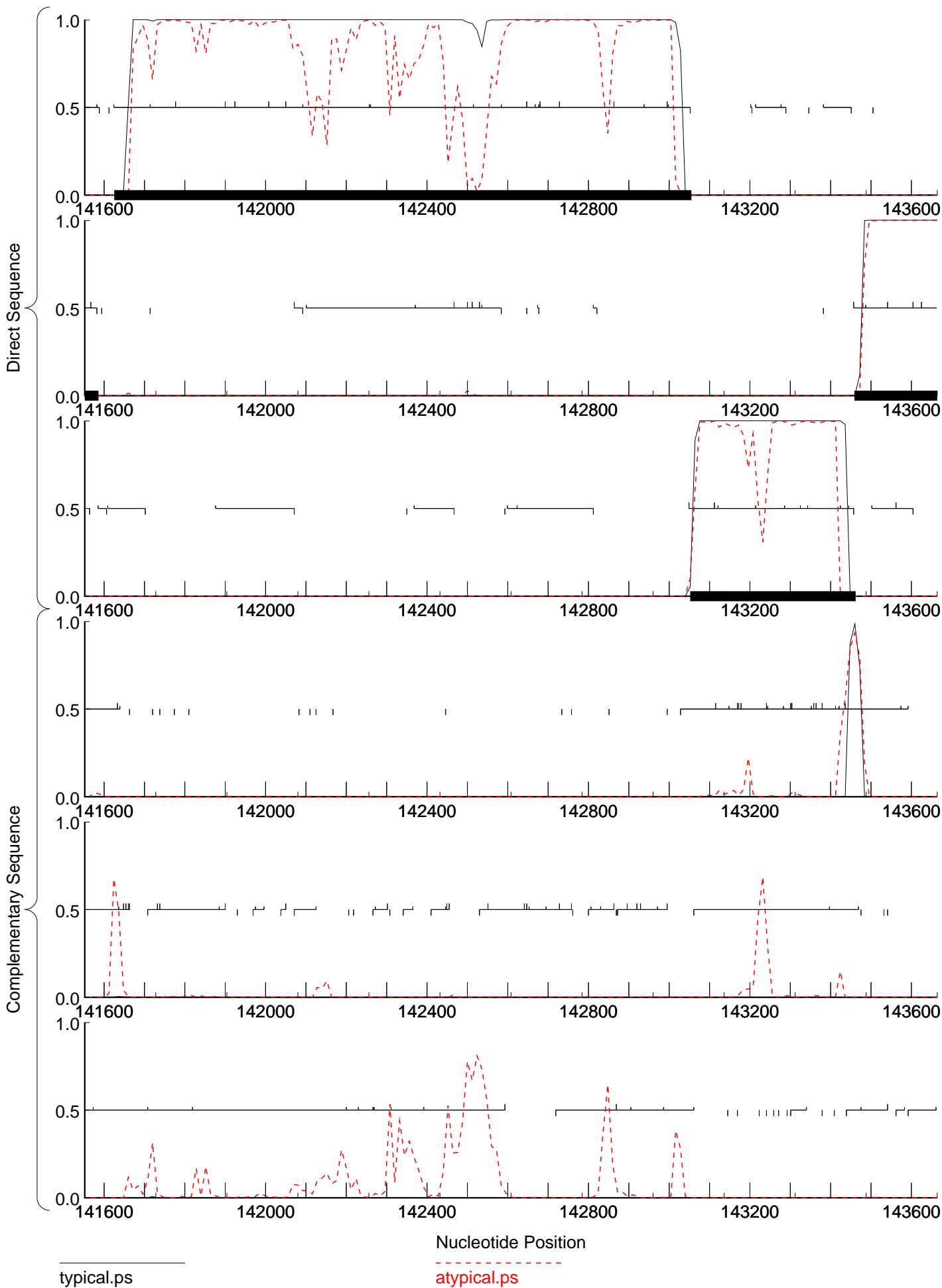


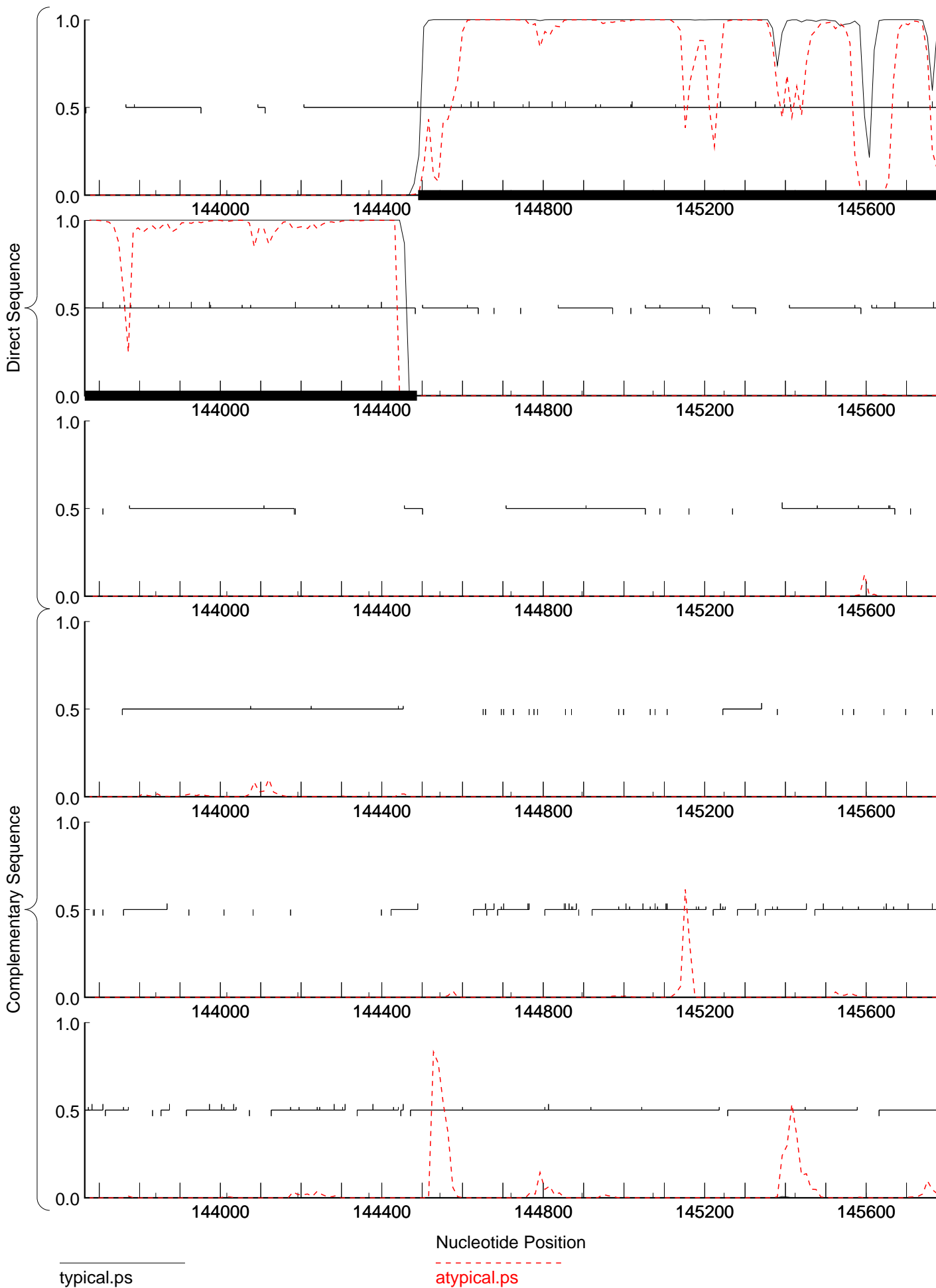


typical.ps

atypical.ps

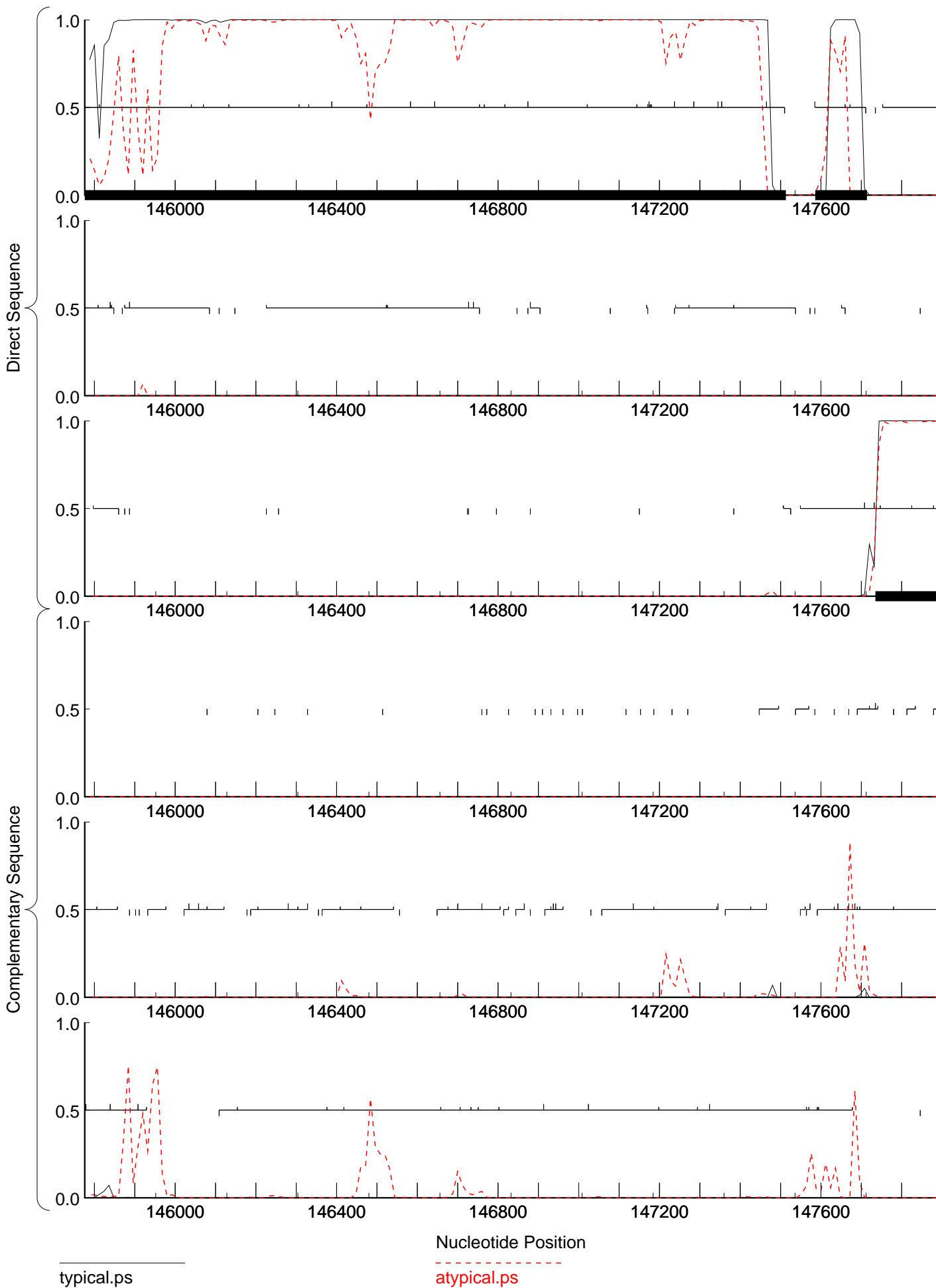


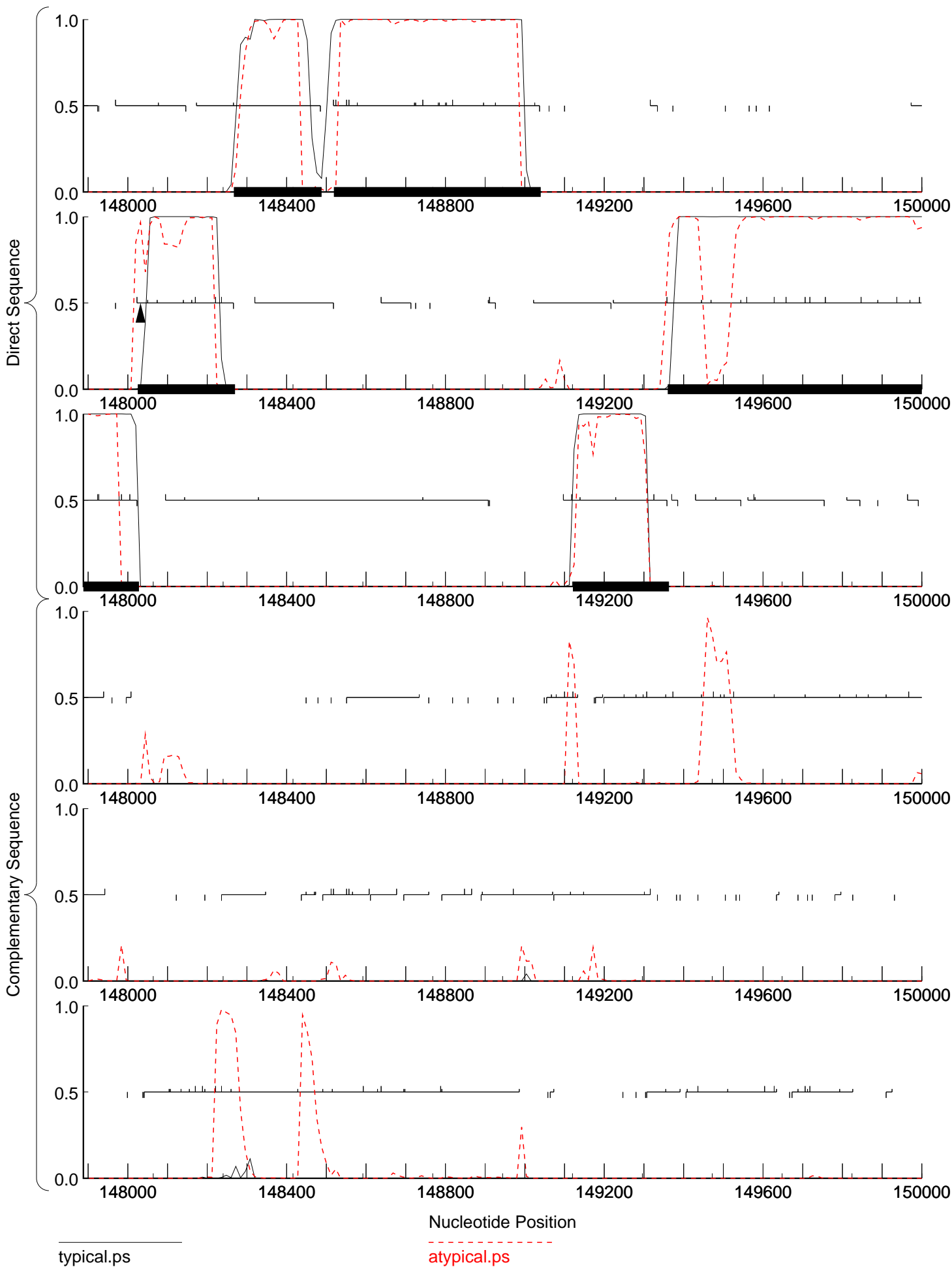




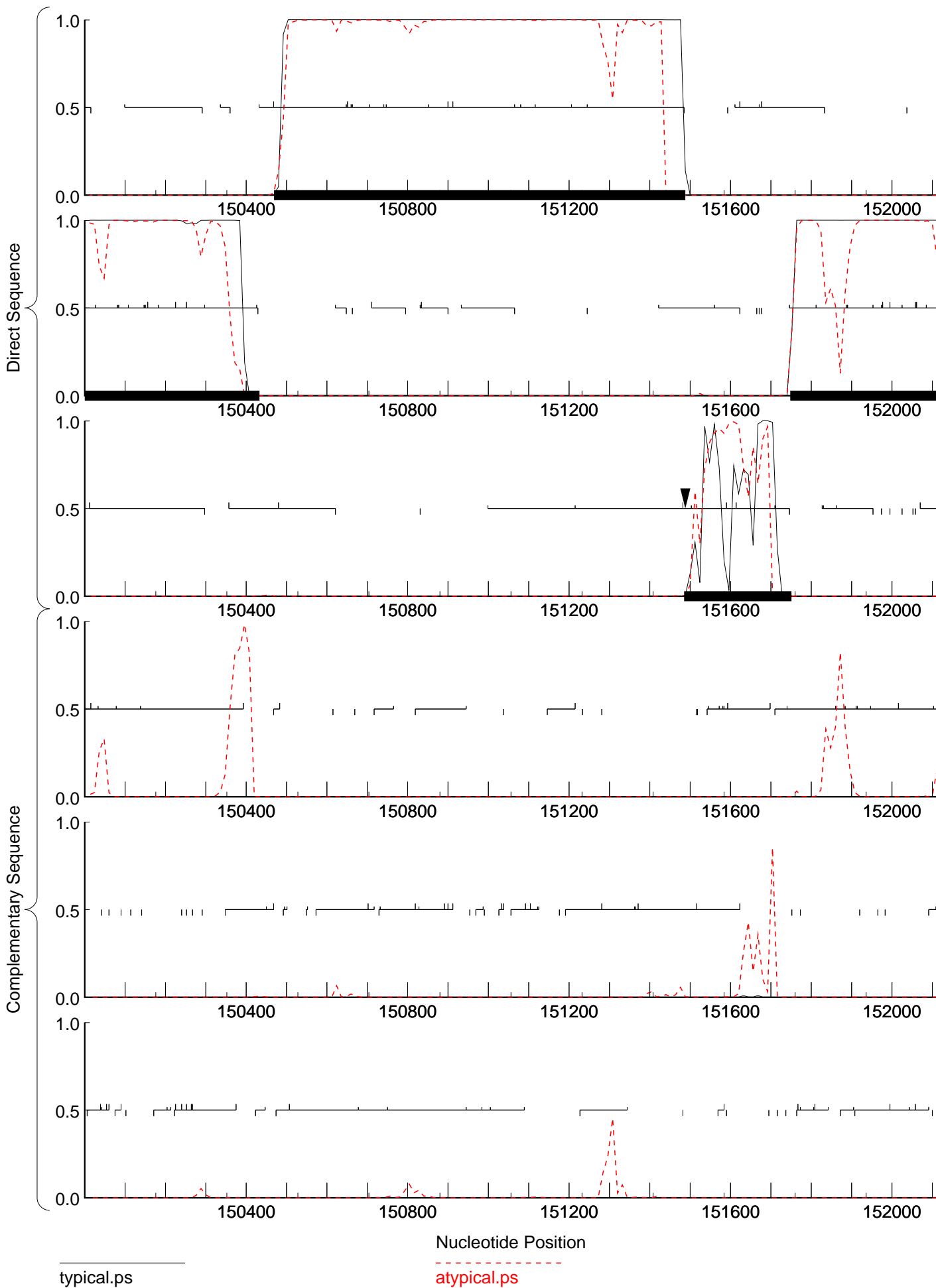
typical.ps

atypical.ps









typical.ps

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