

# GeneMark

Version 2.5p (09.08.06)

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## PROGRAM INFORMATION

Sequence : Streptomyces phage Kradal complete sequence, 186383 bp including ~1053 bp terminal repeat,  
Analysis Date : 7/16/18 at 12:30:27  
Pages : 90  
Sequence Length : 186383 bp  
GC Content : 66.72%

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,

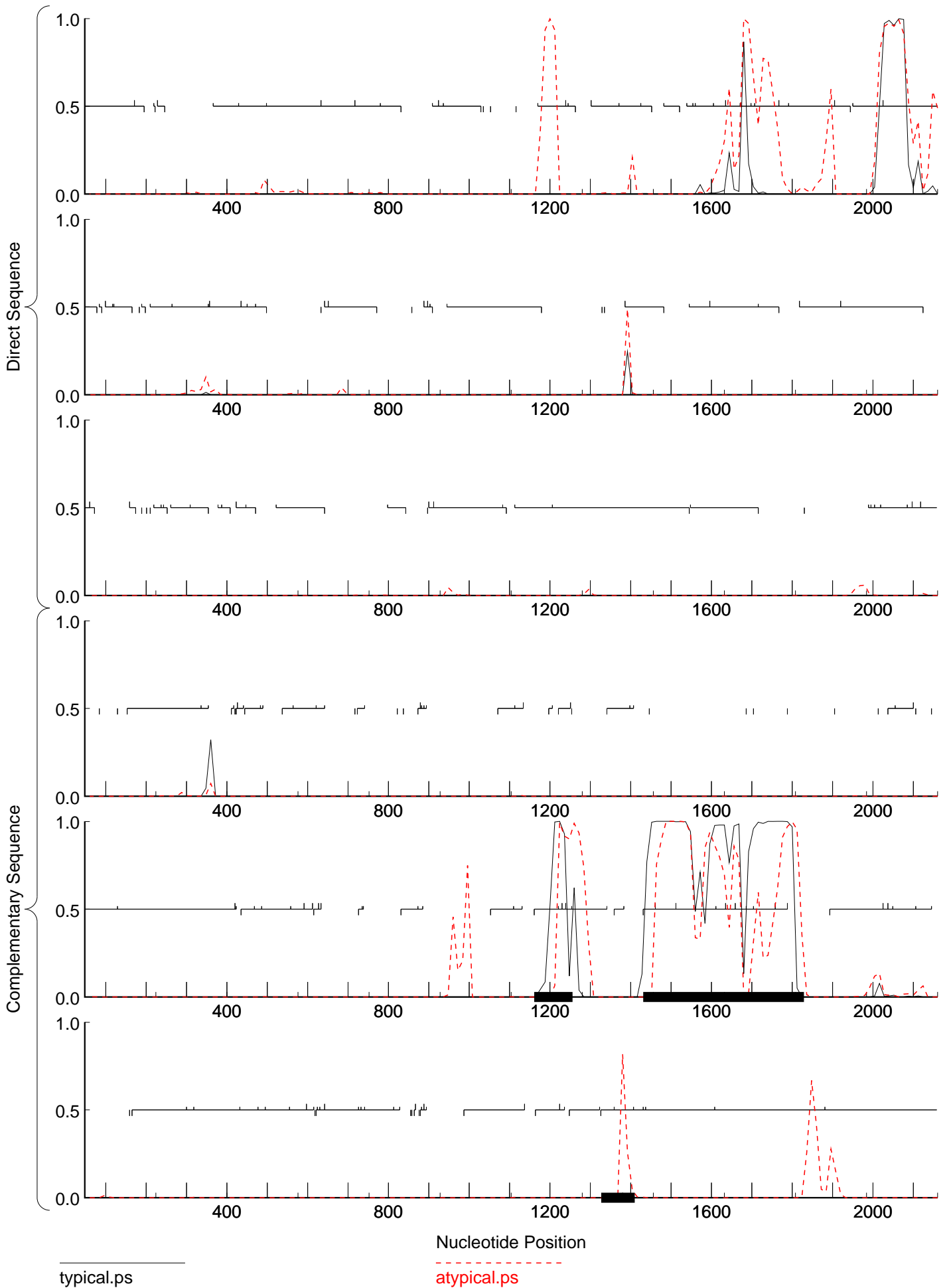
## MATRIX INFORMATION

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

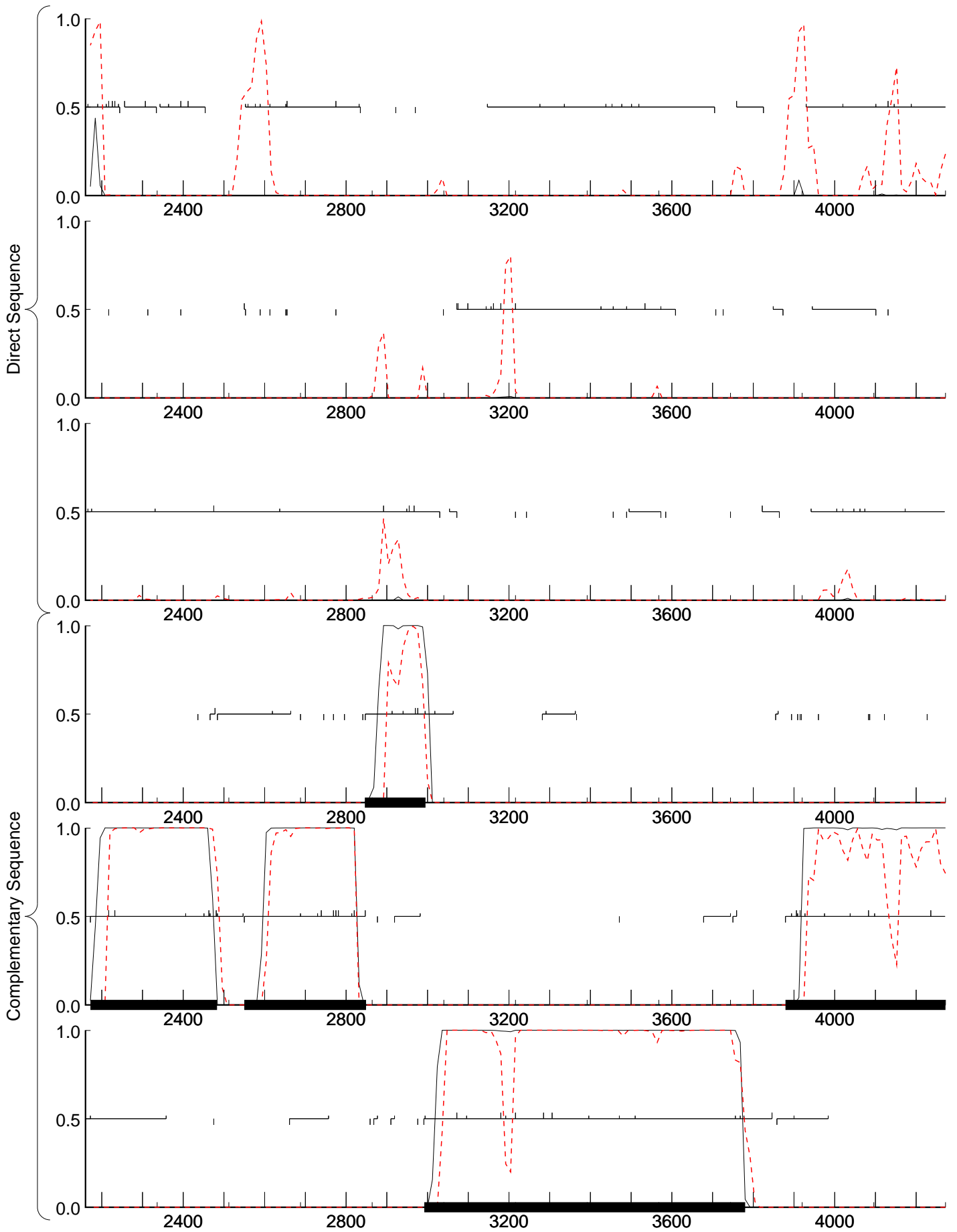
Training set derived by GeneMarkS, 4.30 October 2014  
Mon Jul 16 12:30:27 2018



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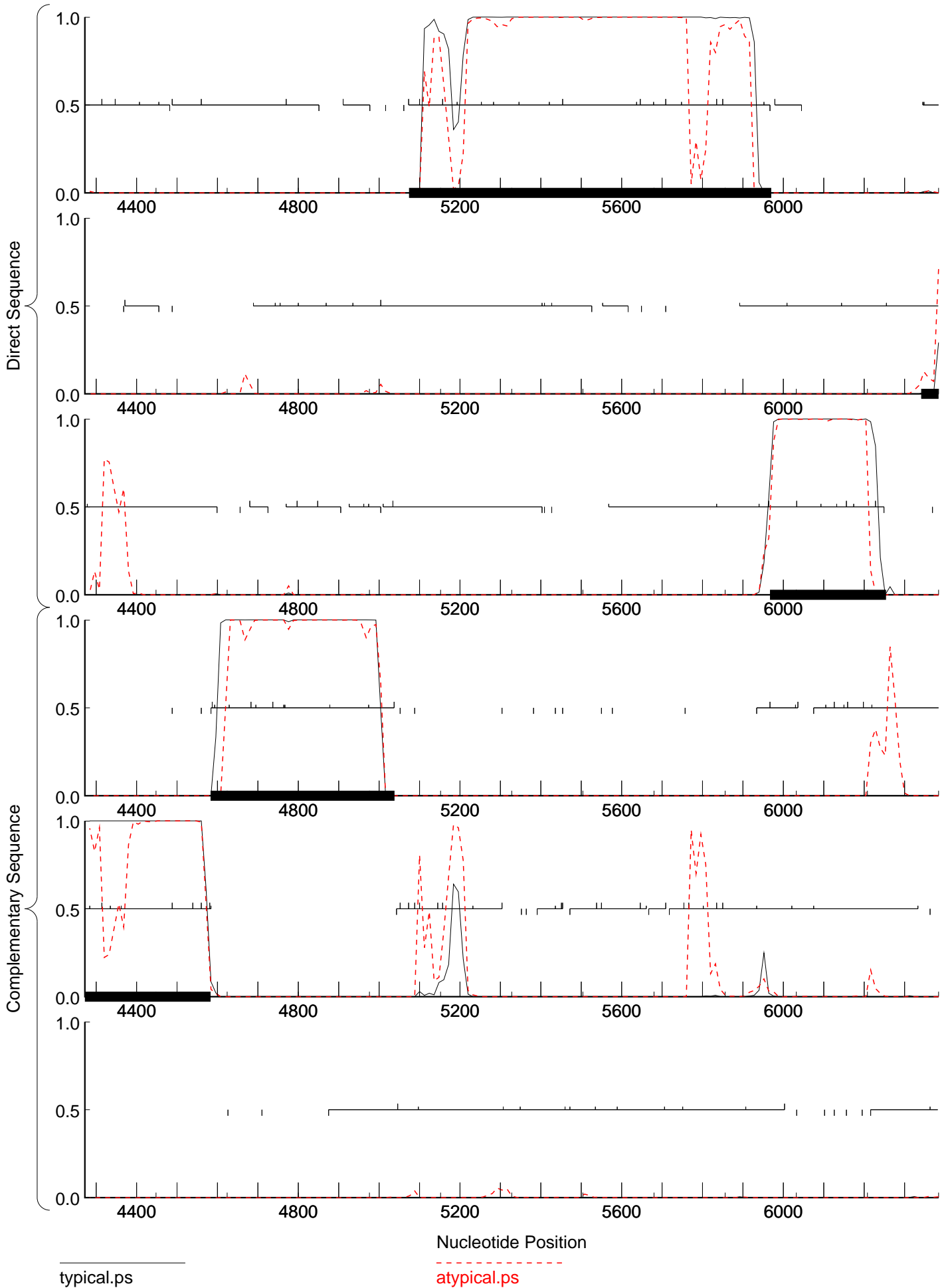
Nucleotide Position

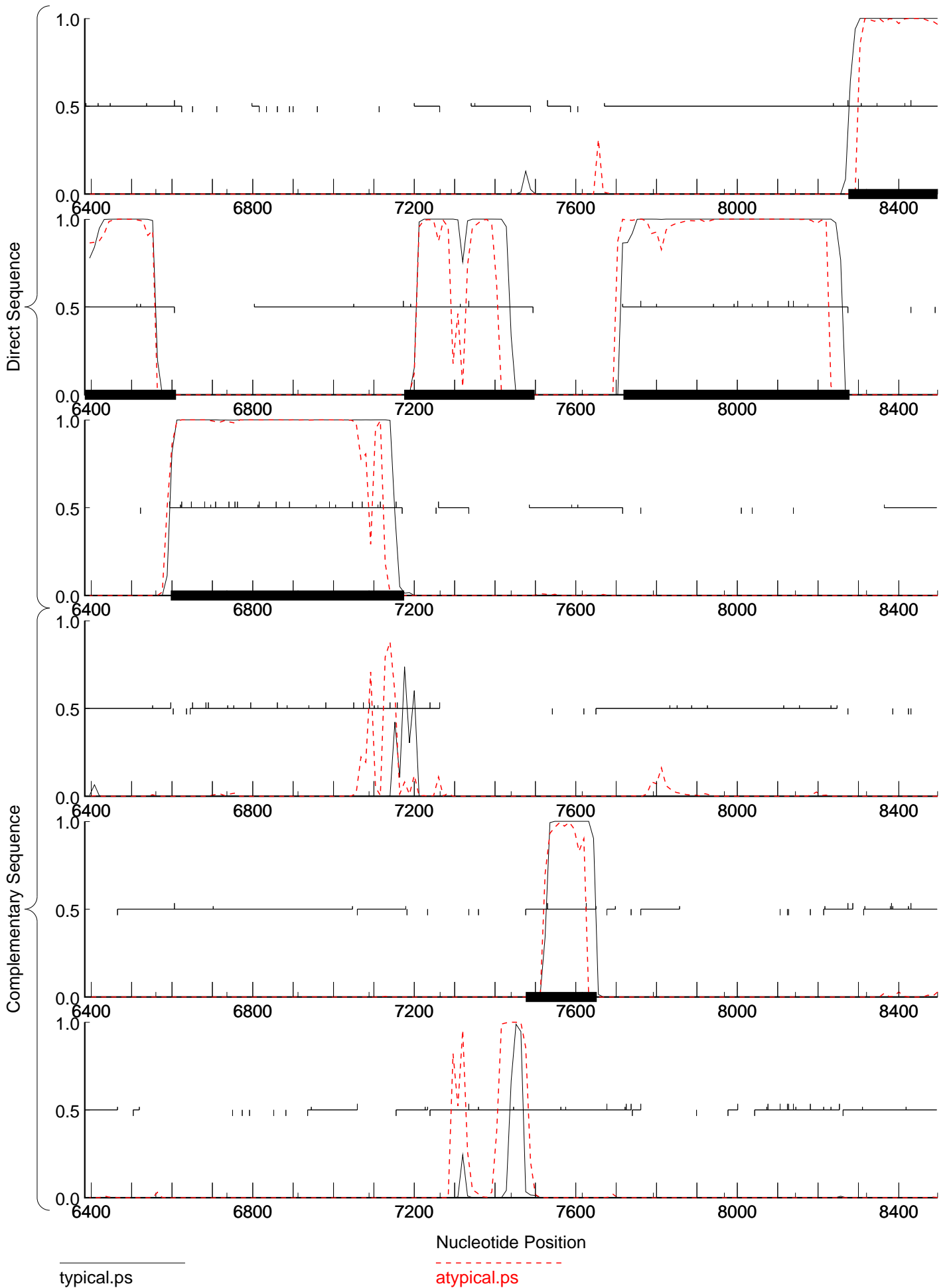
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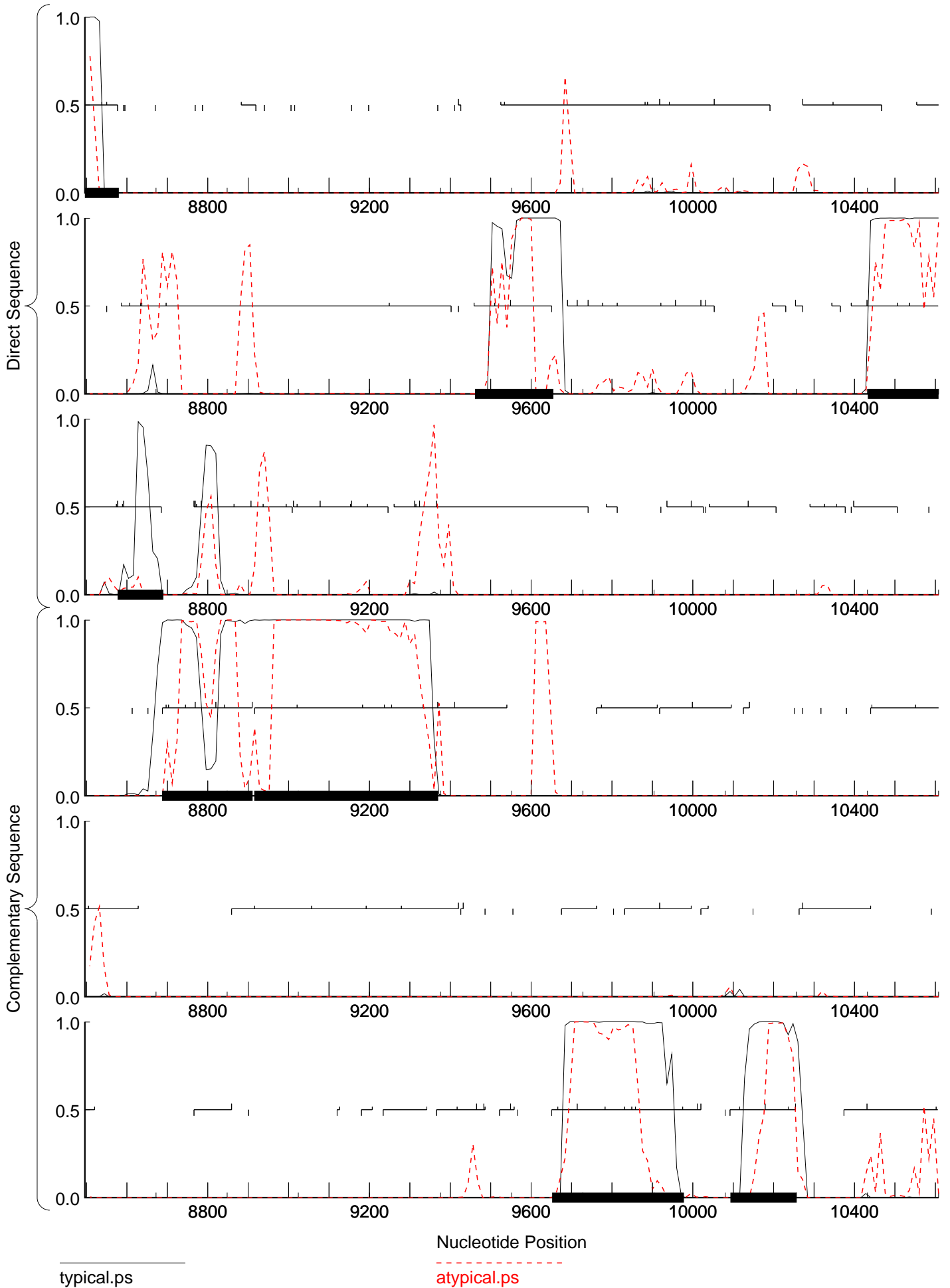
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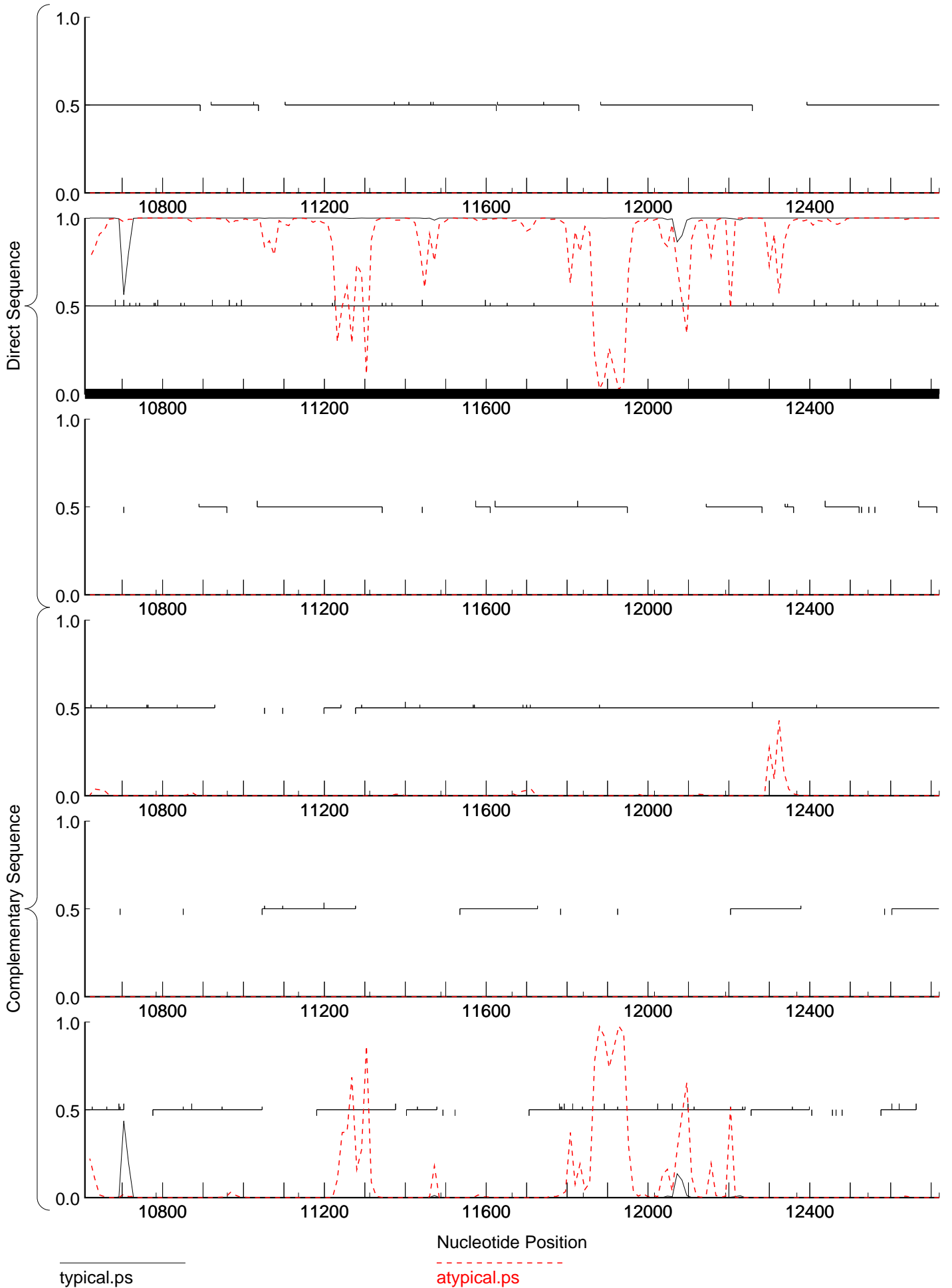


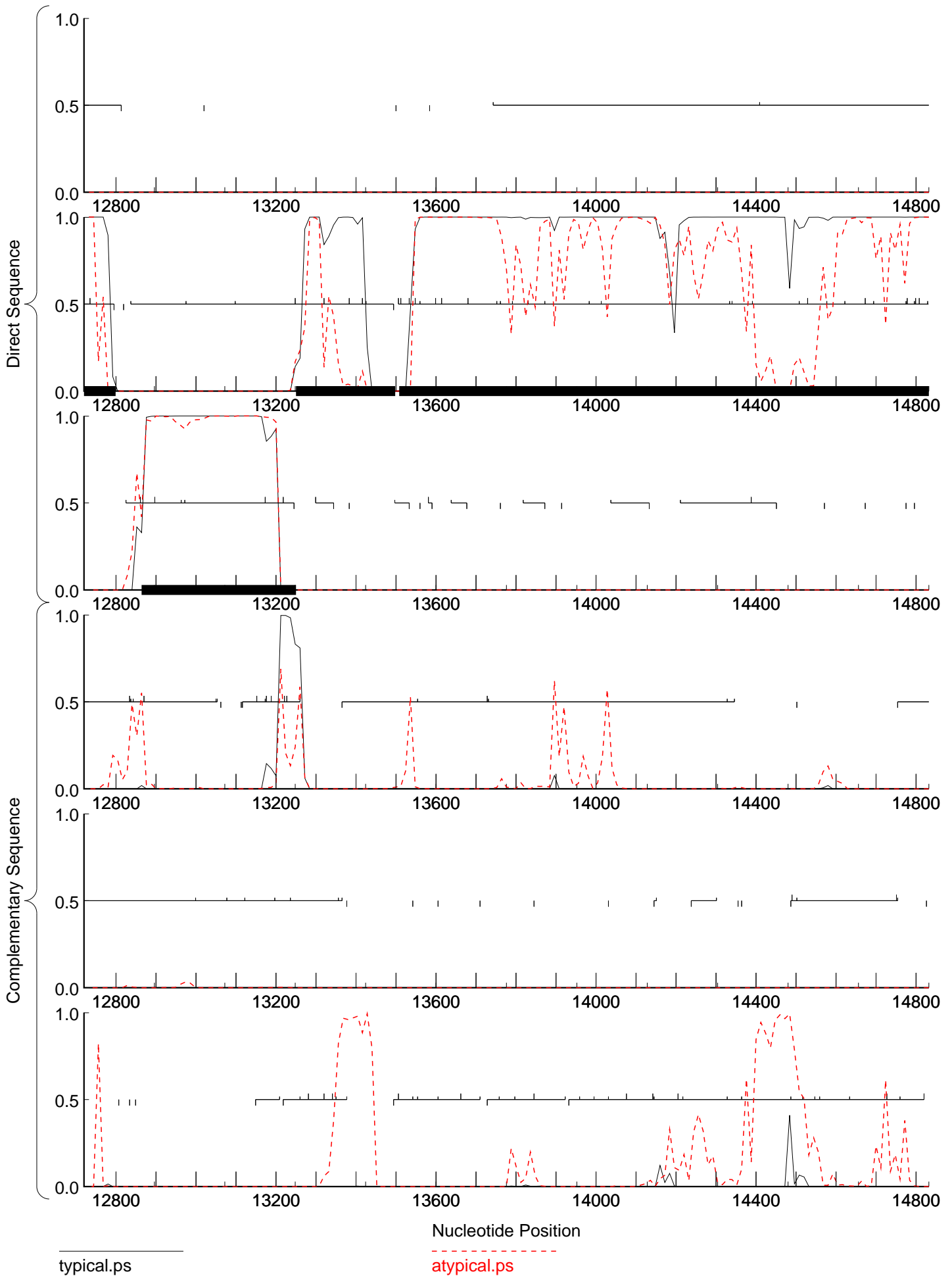


typical.ps

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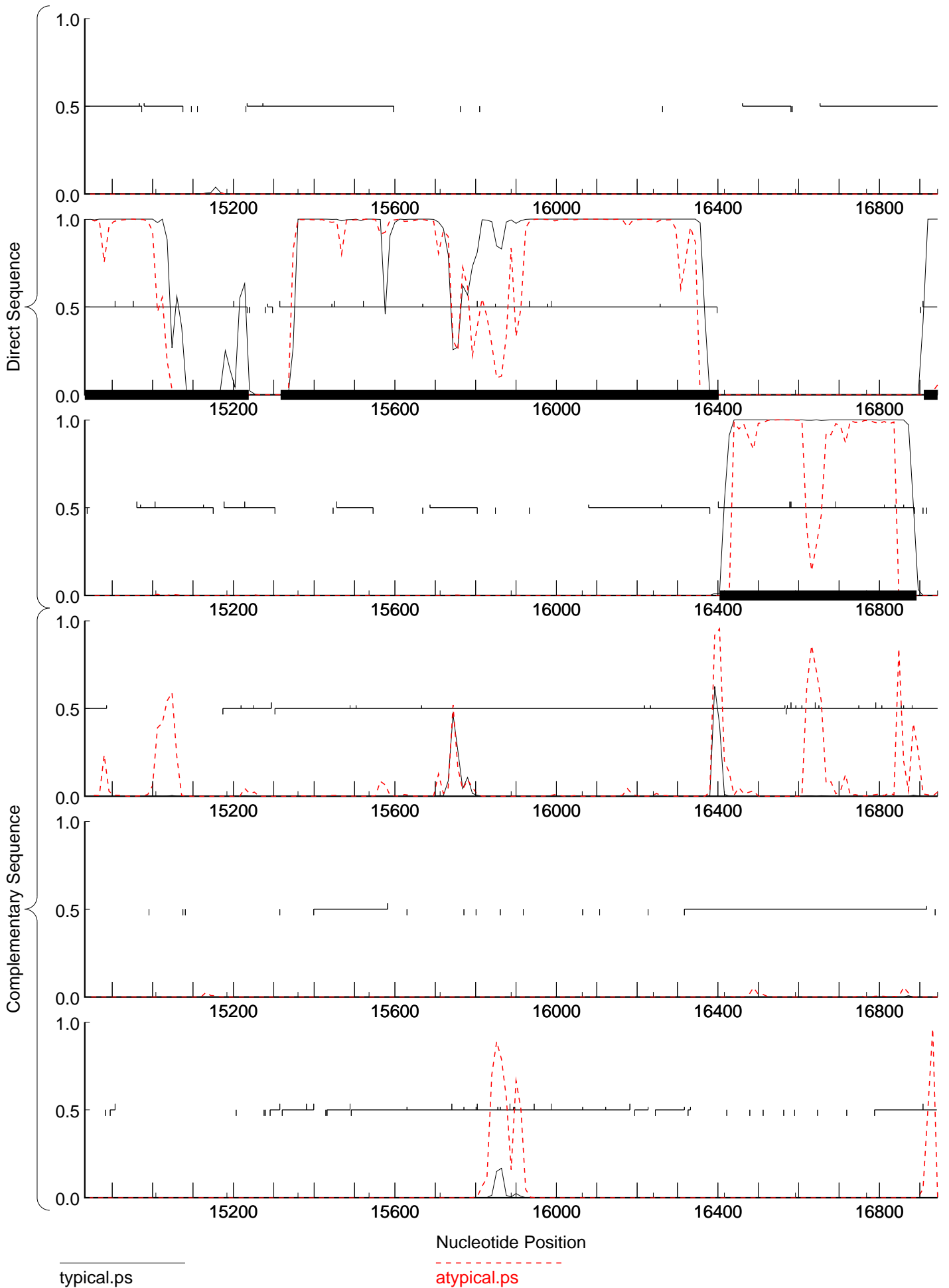


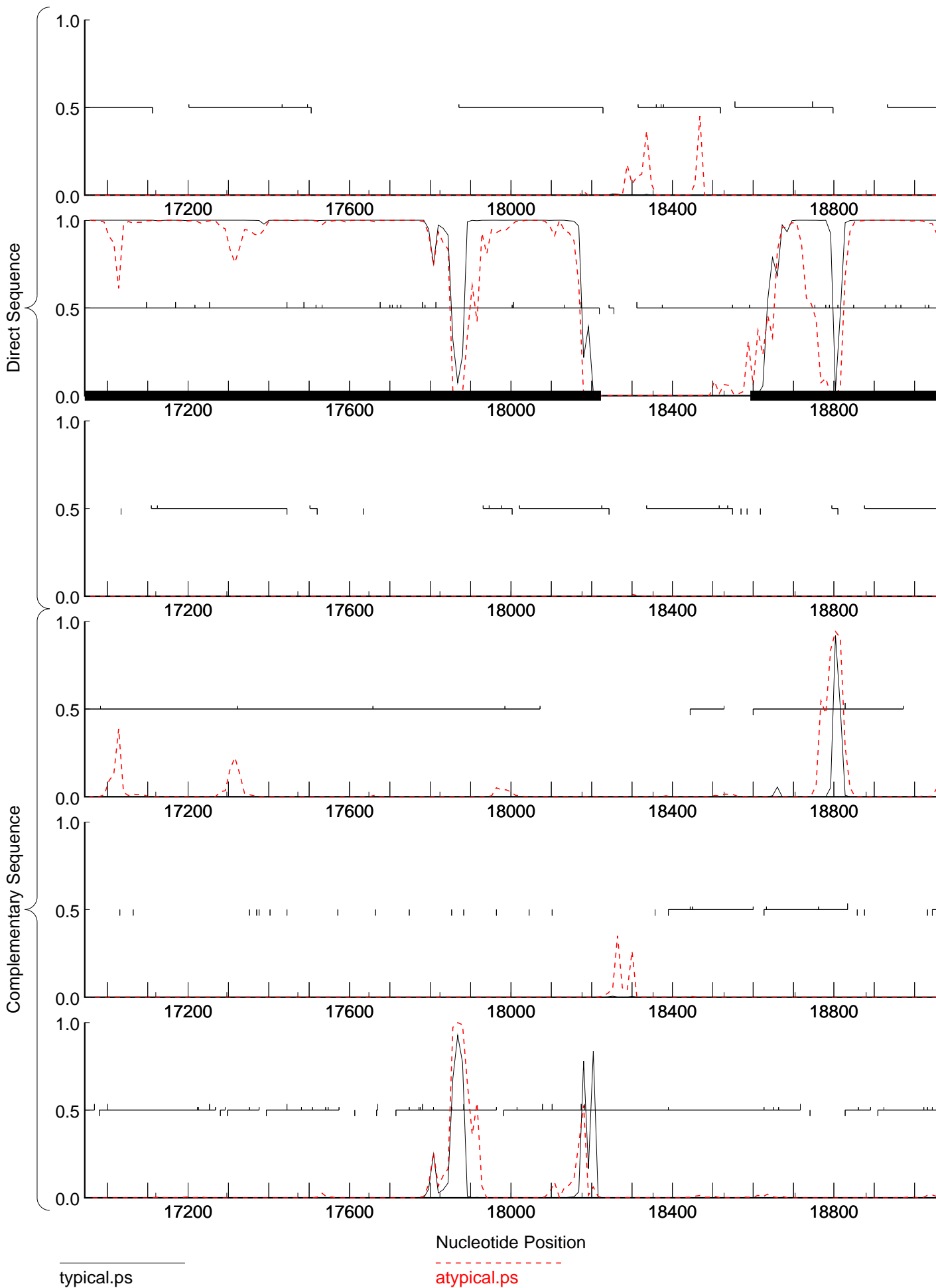


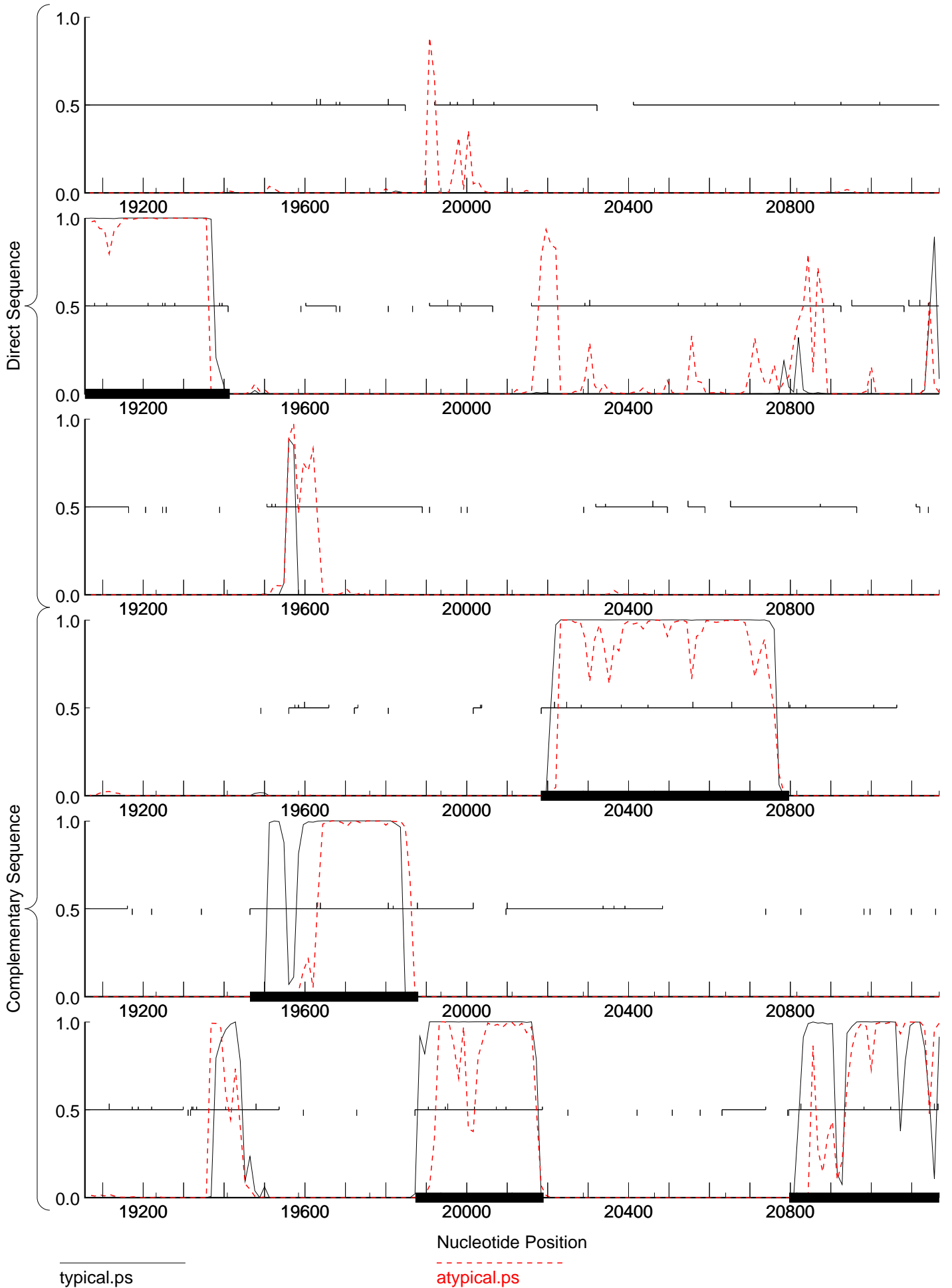
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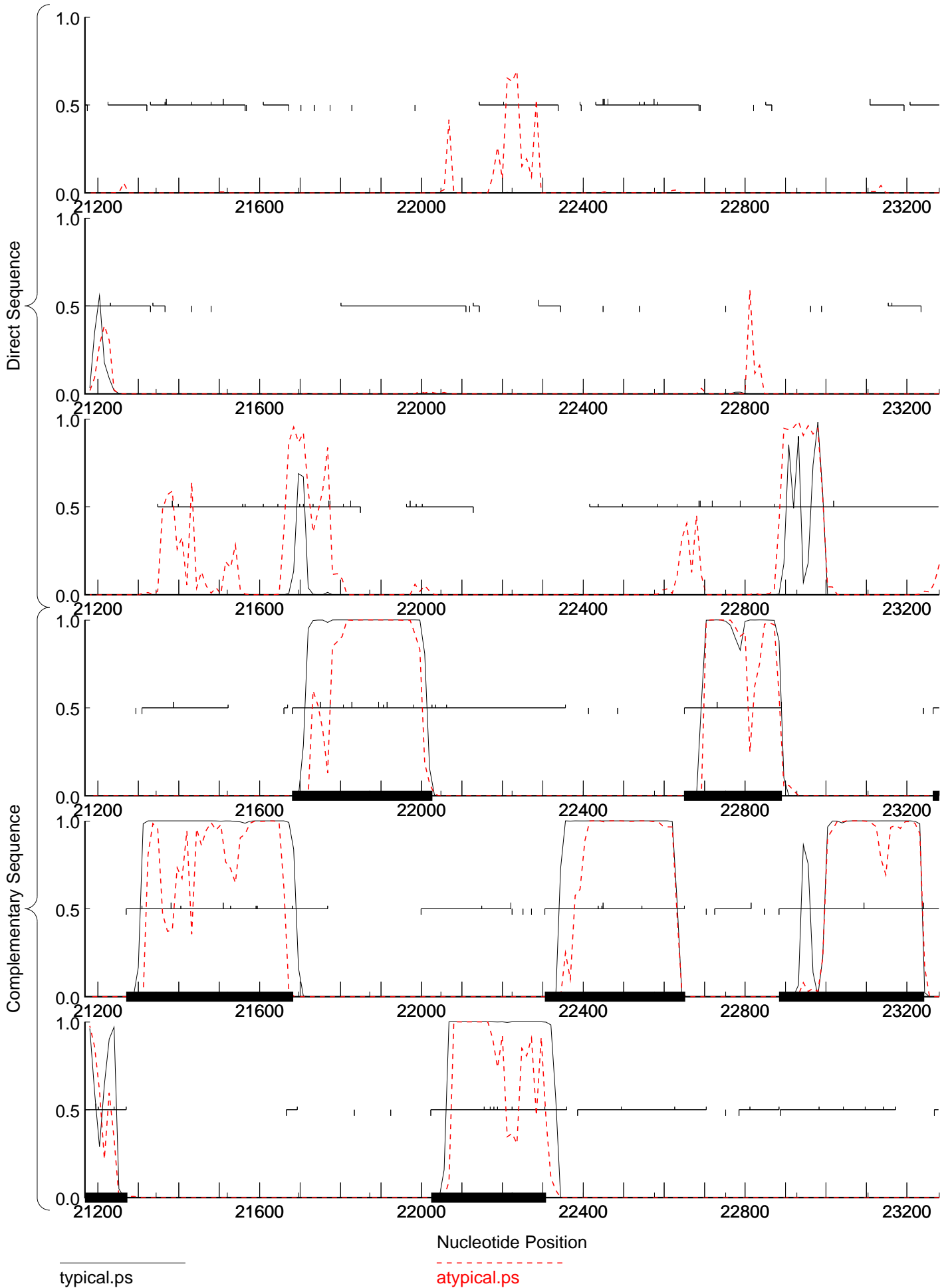
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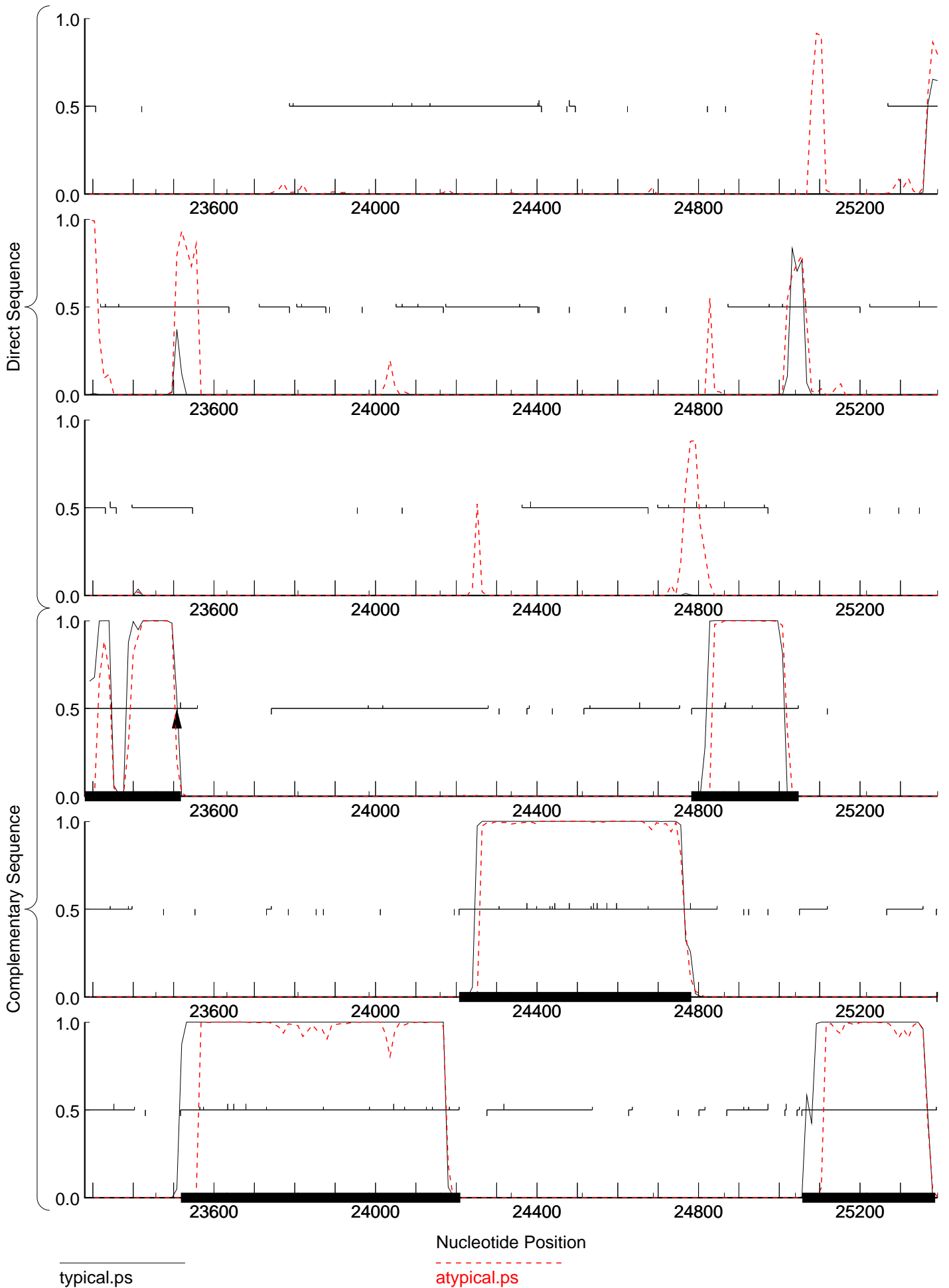


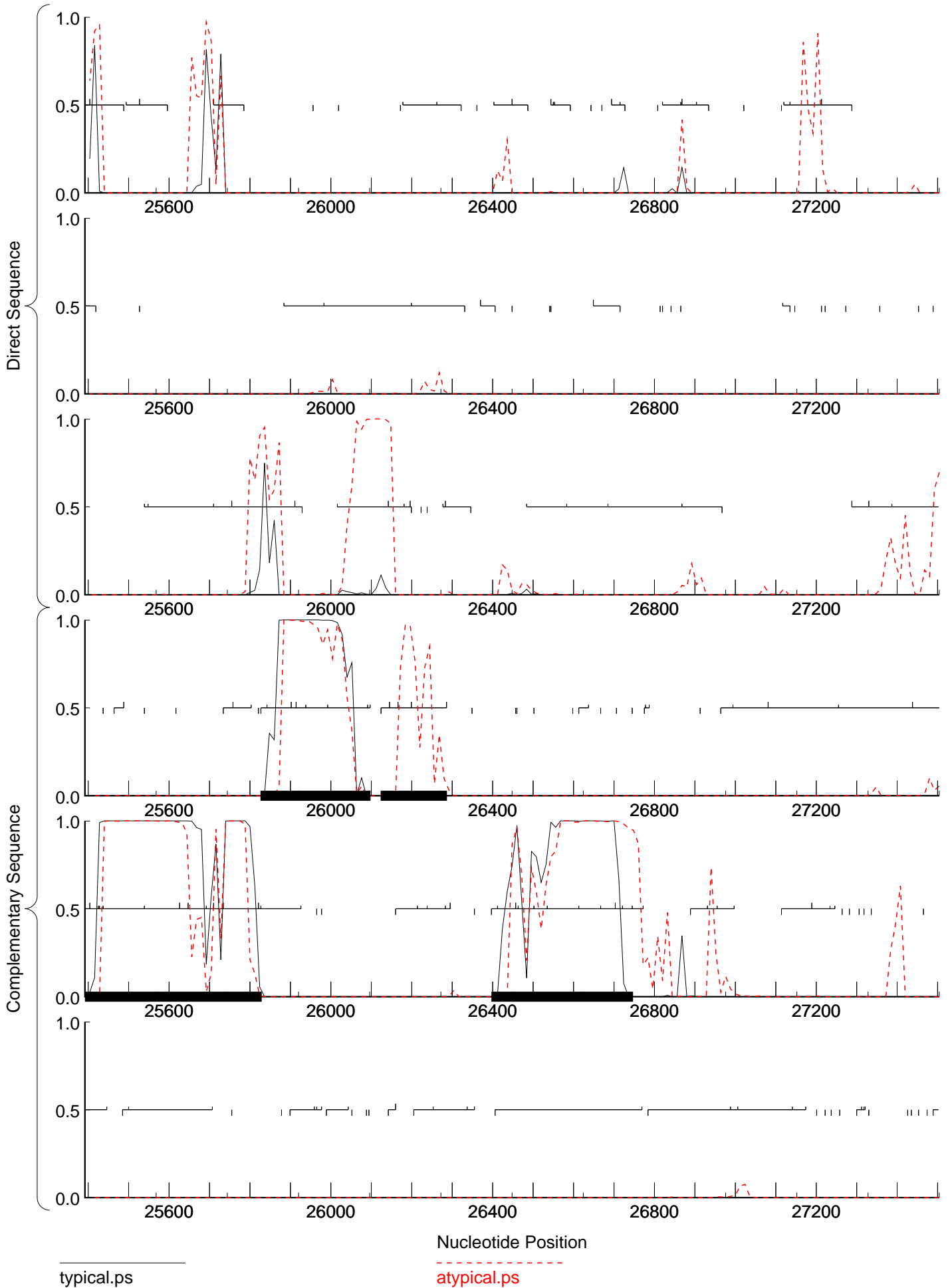






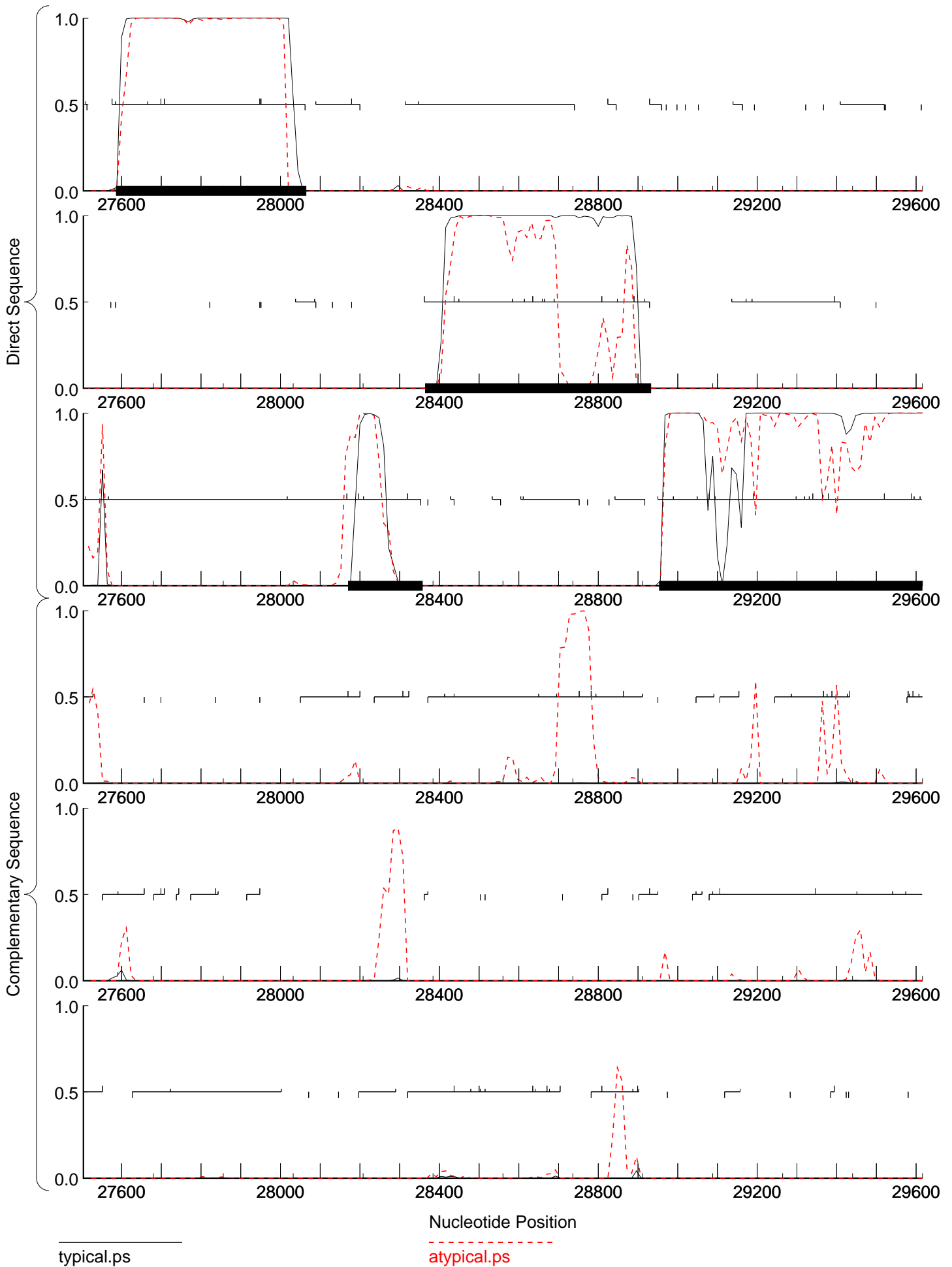






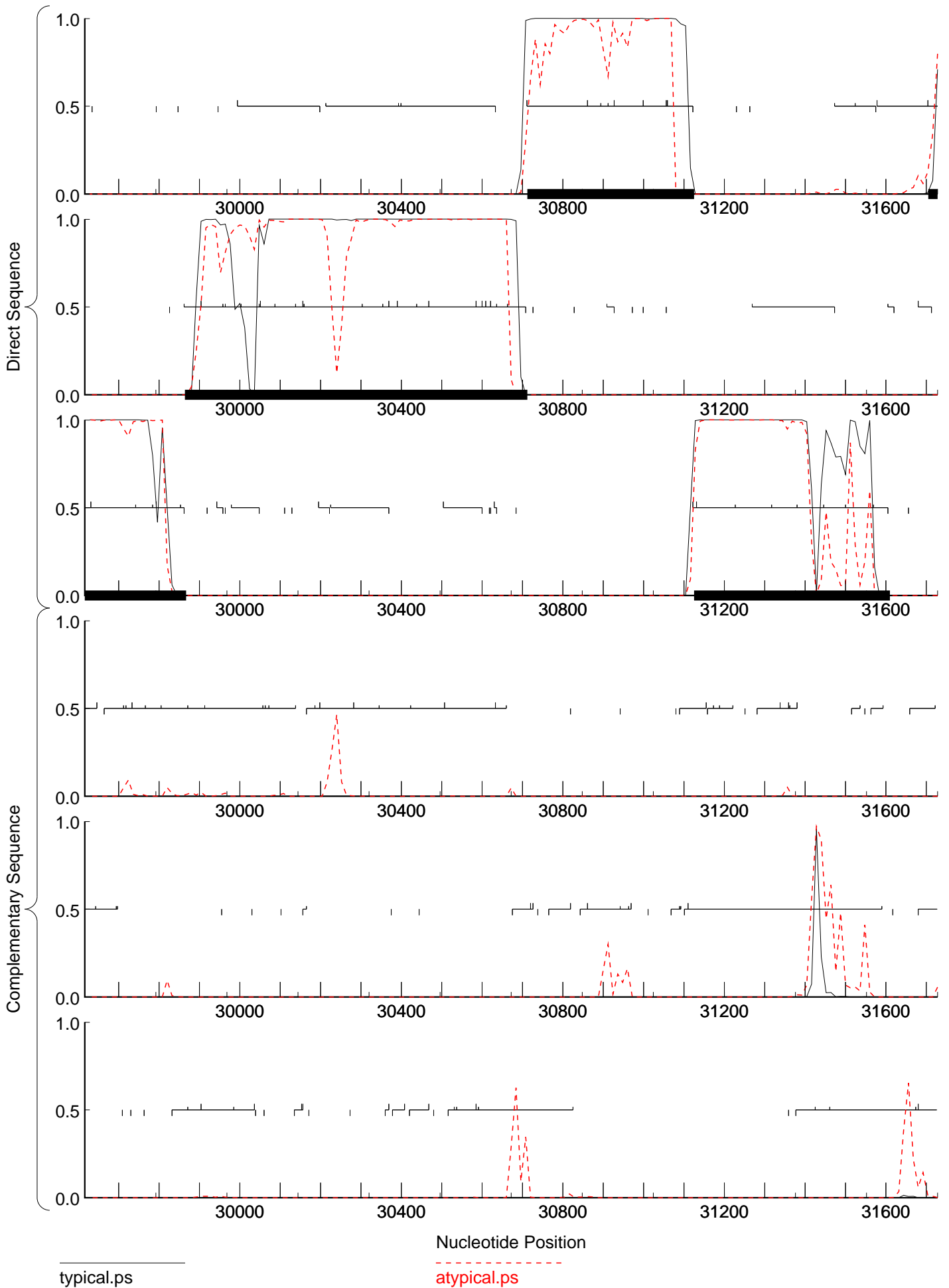
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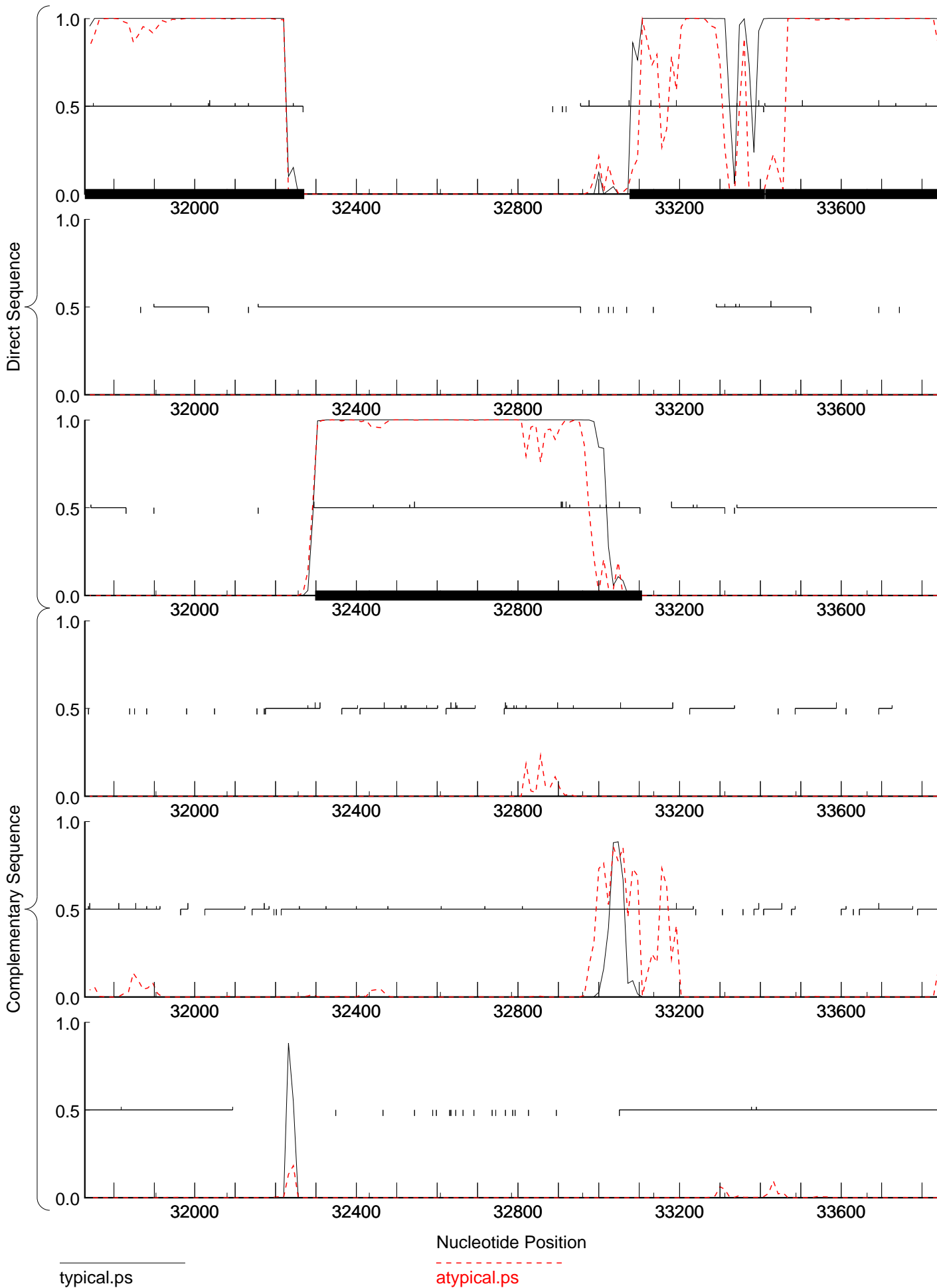


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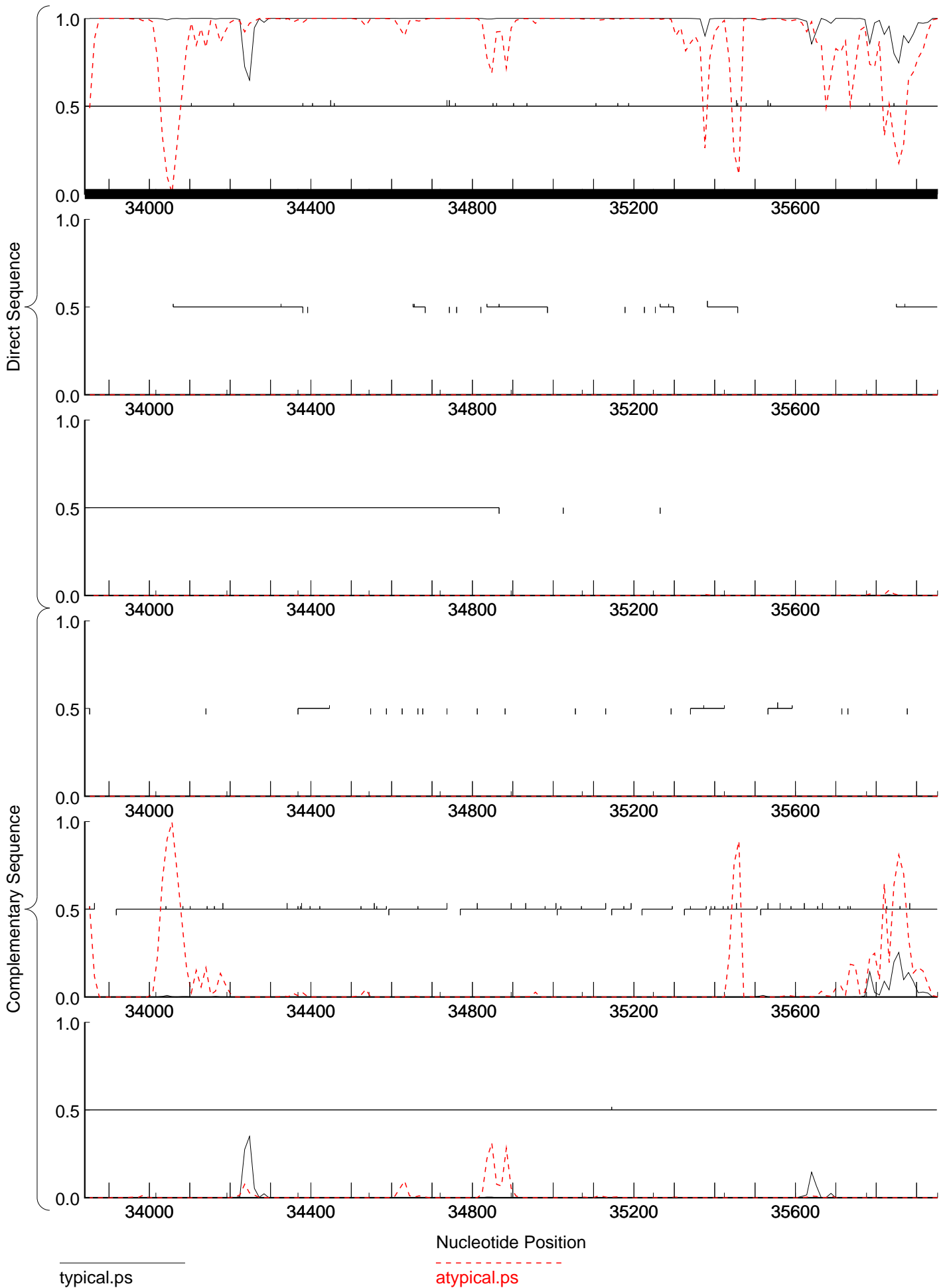


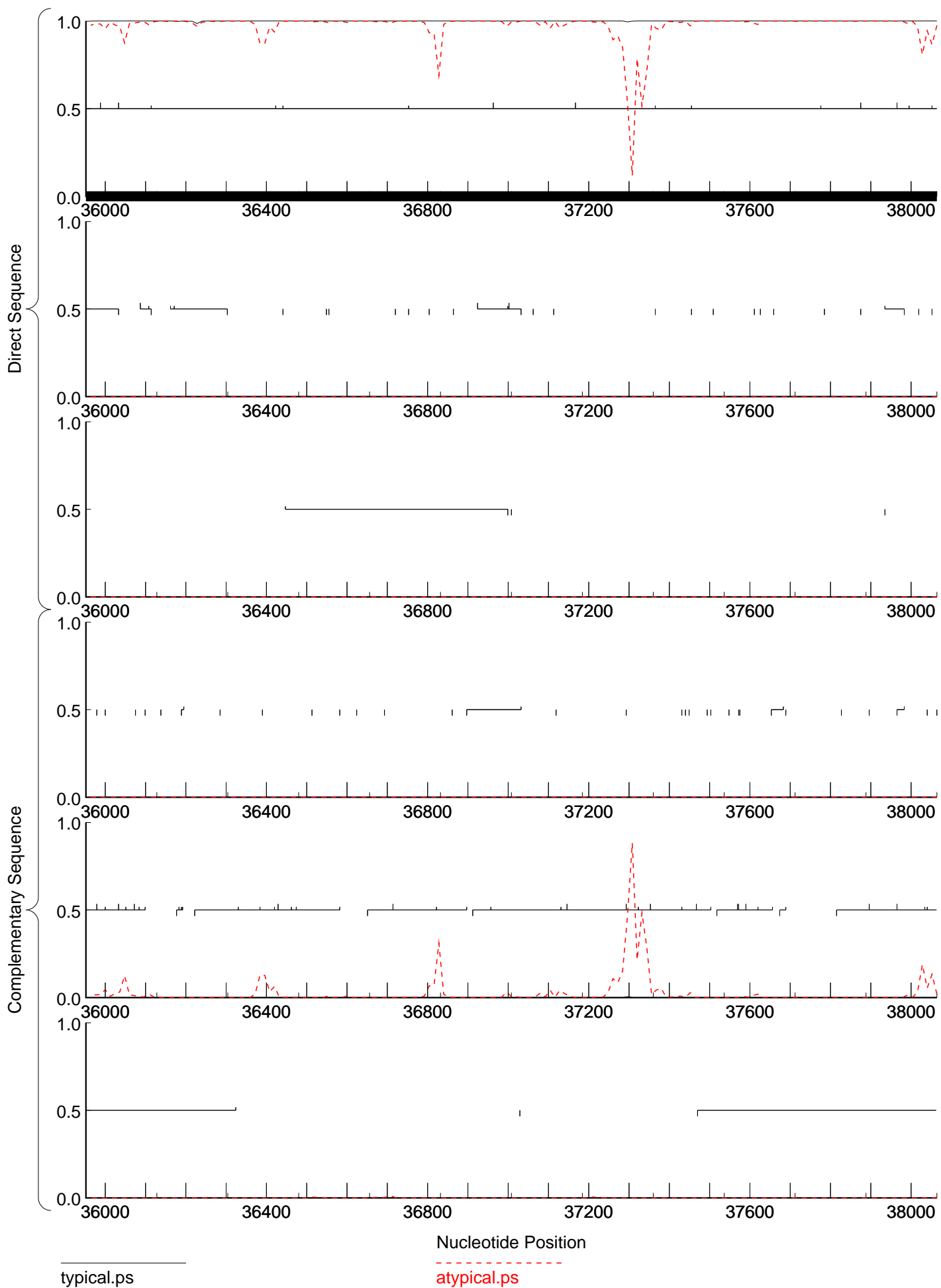




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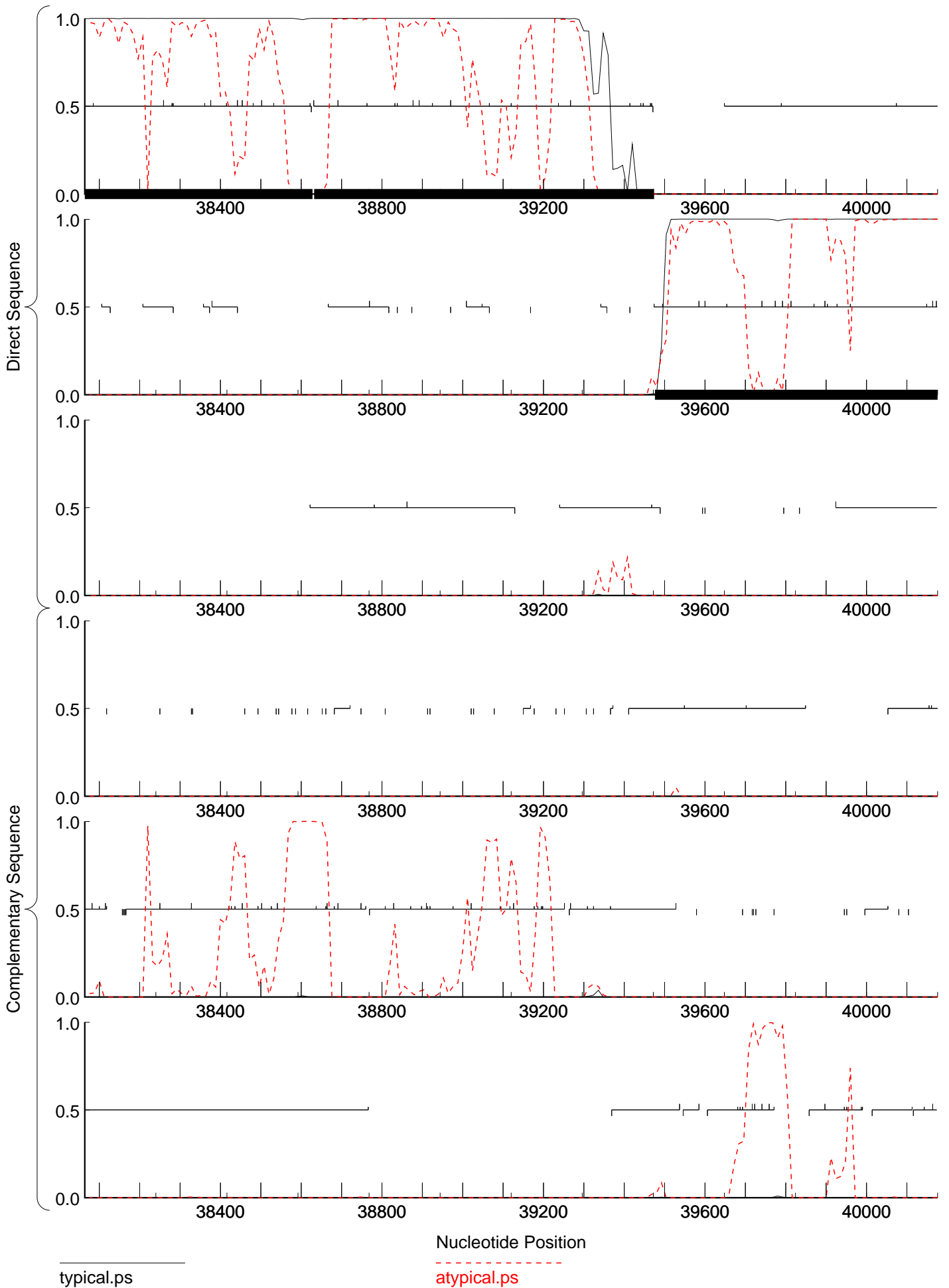
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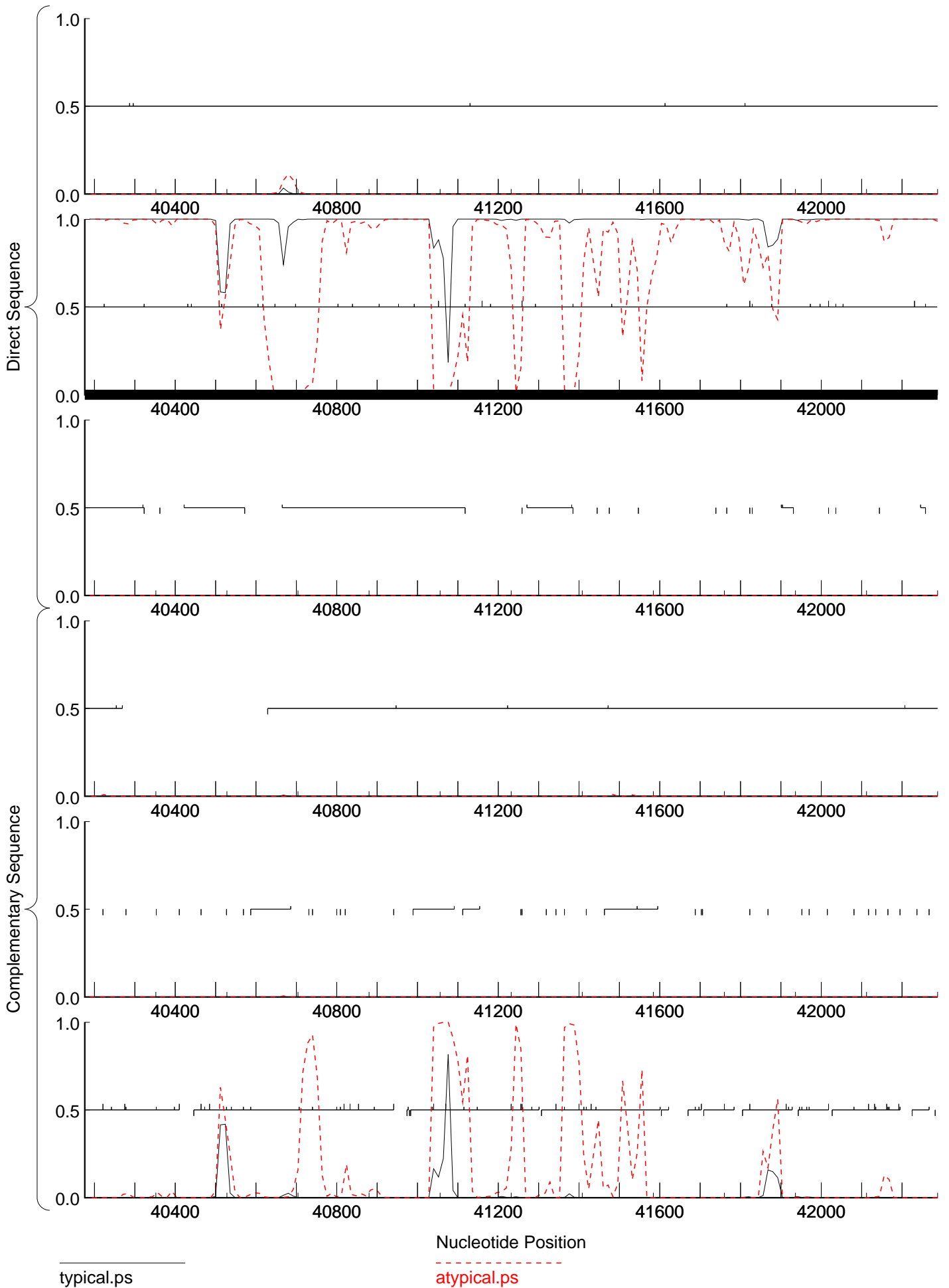


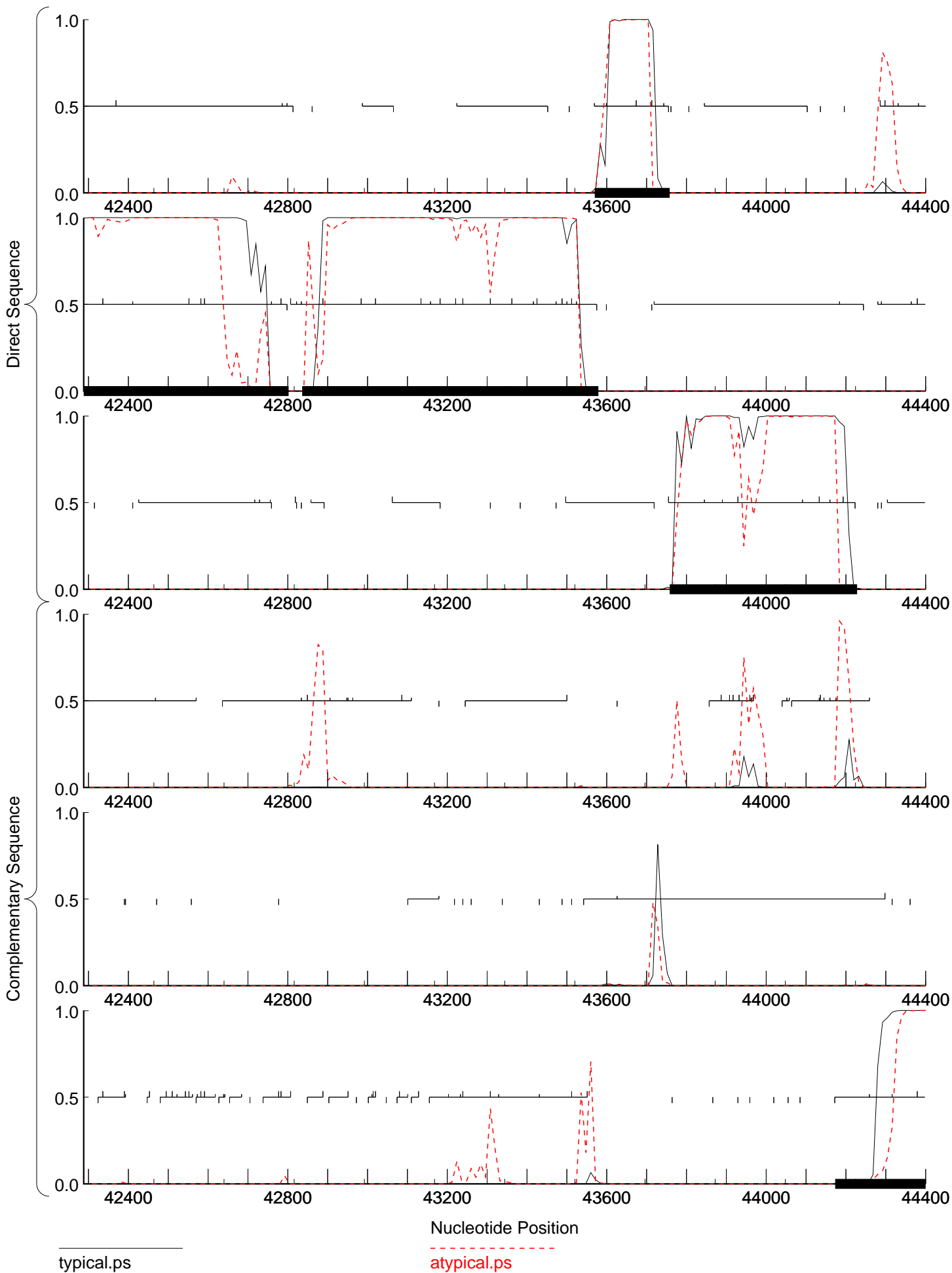


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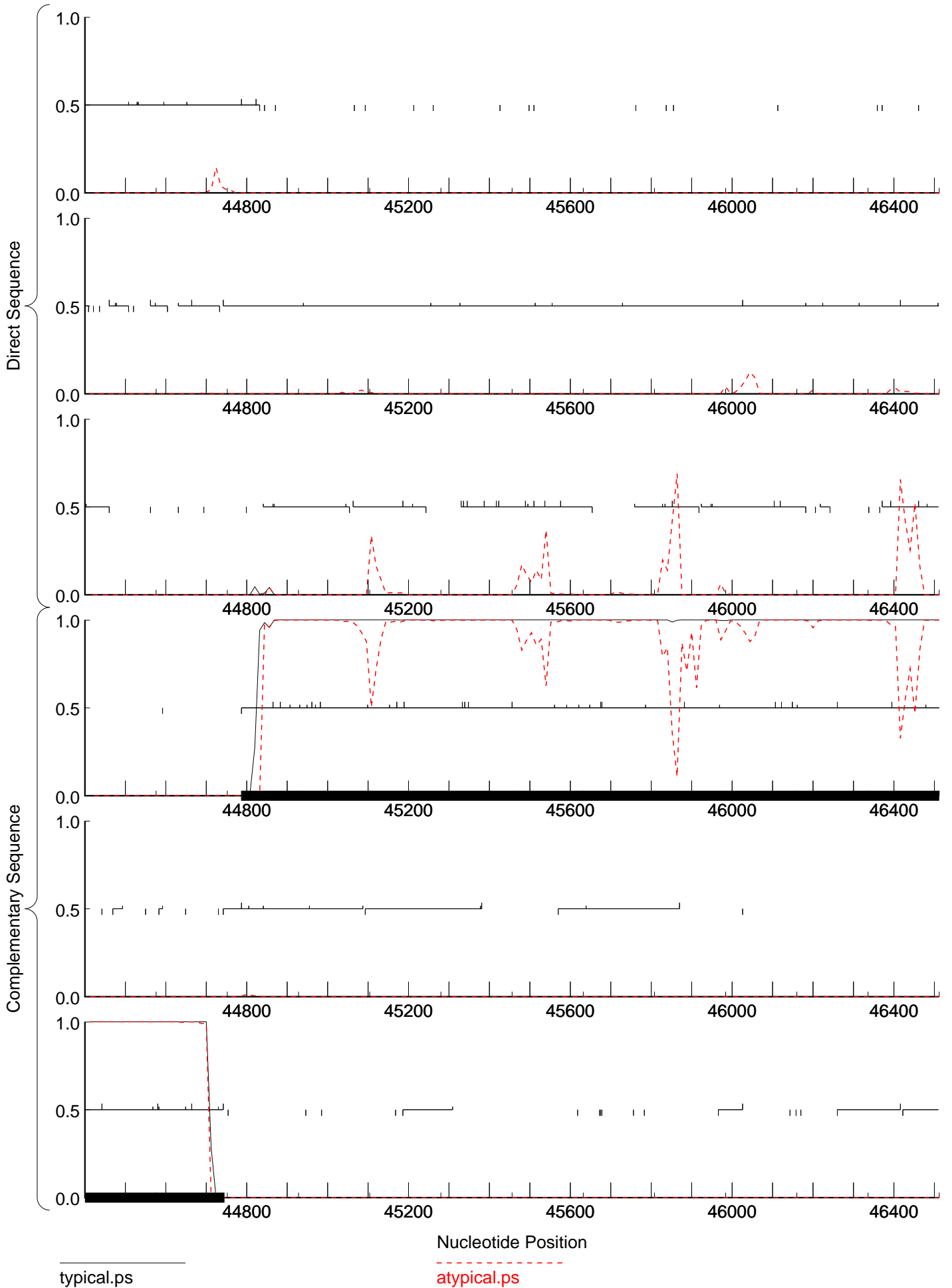


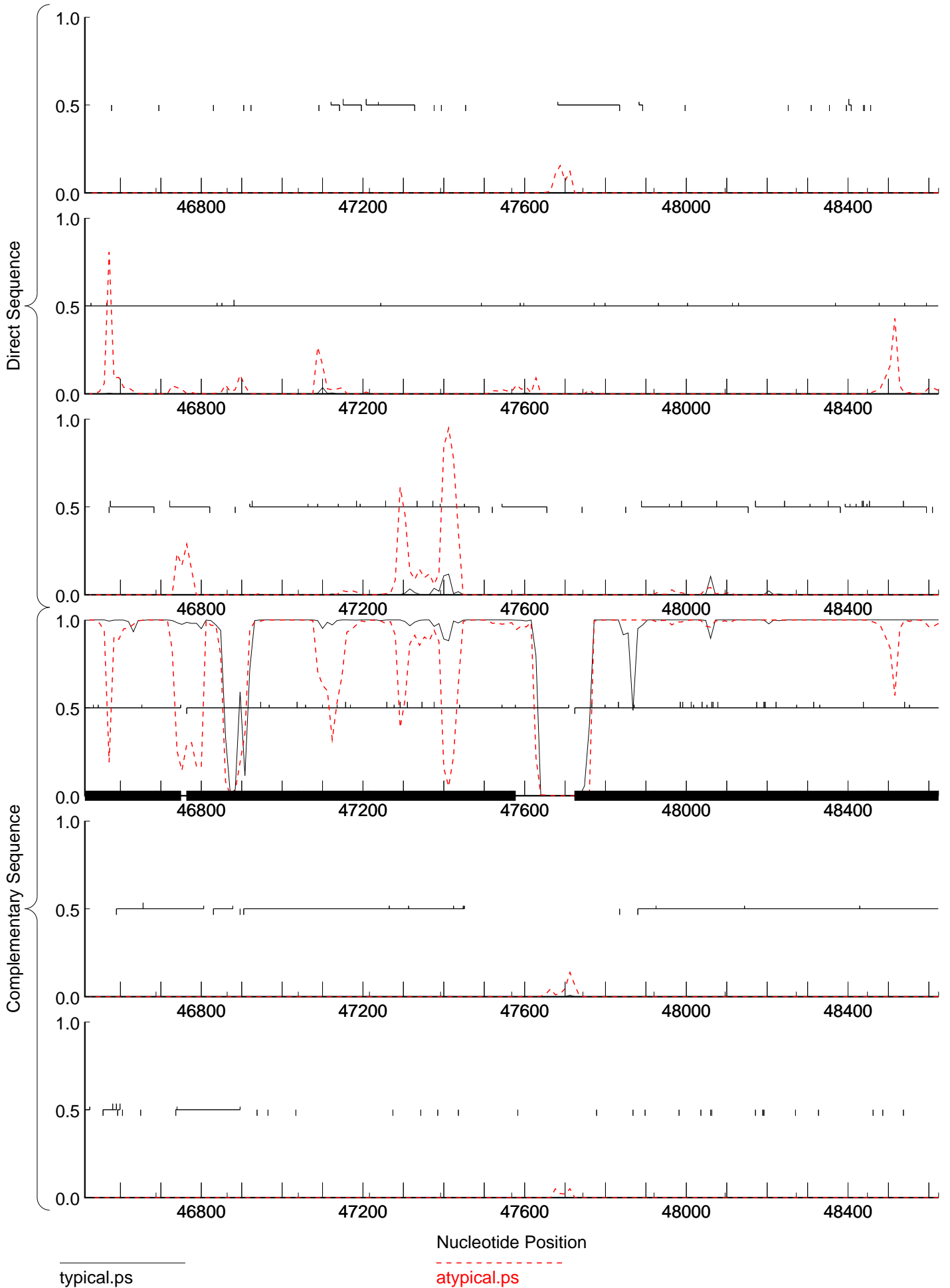




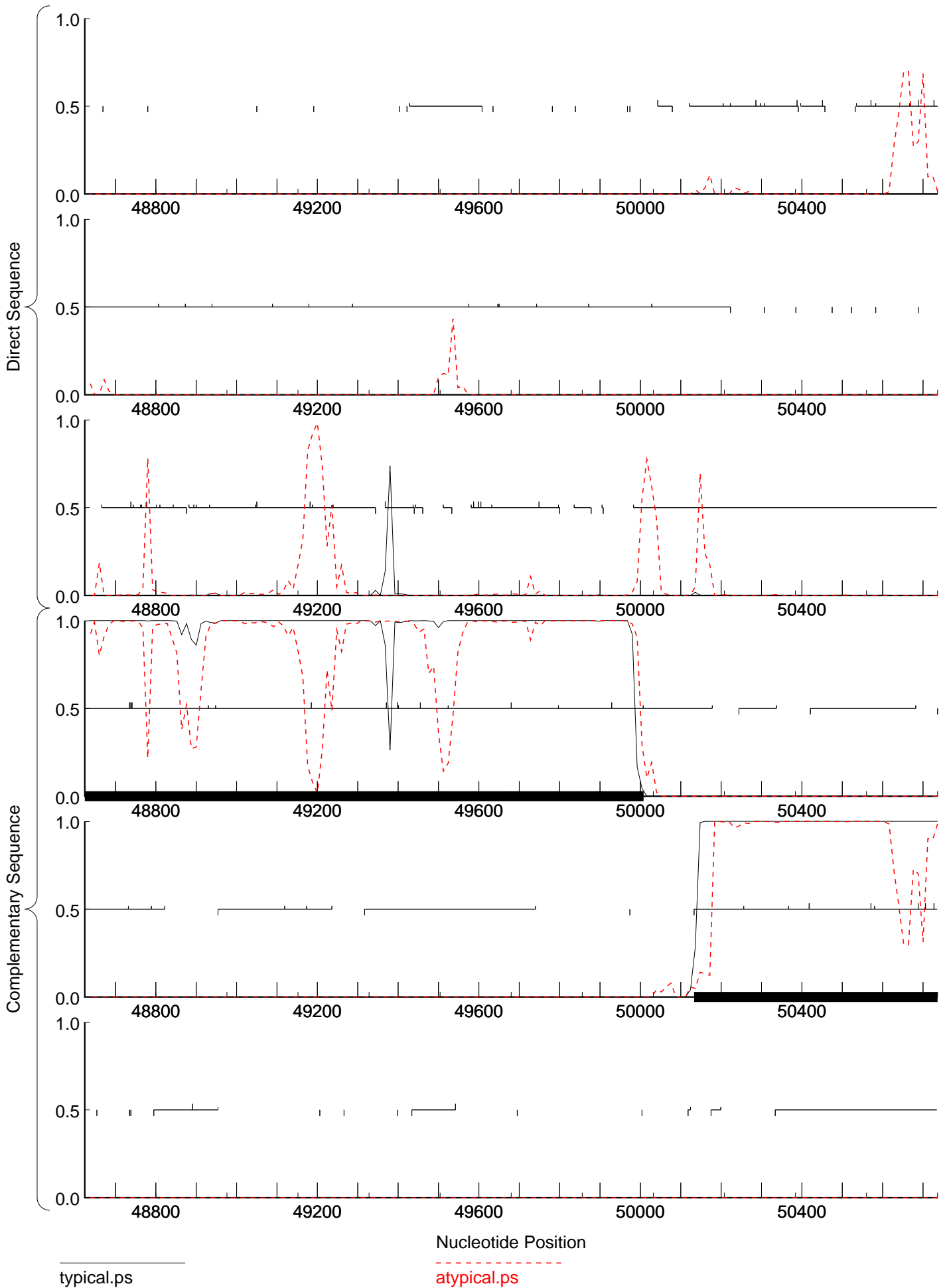
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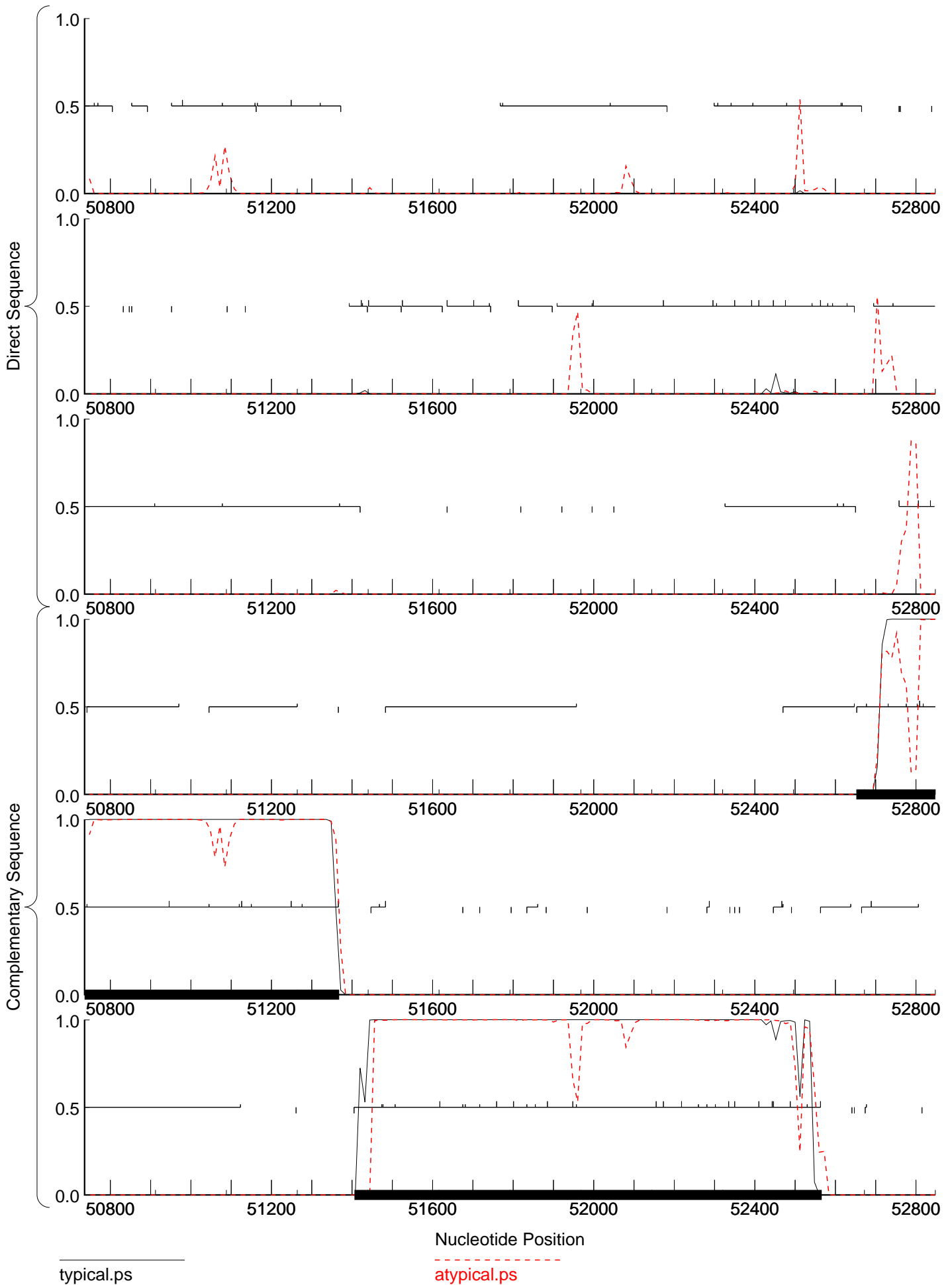


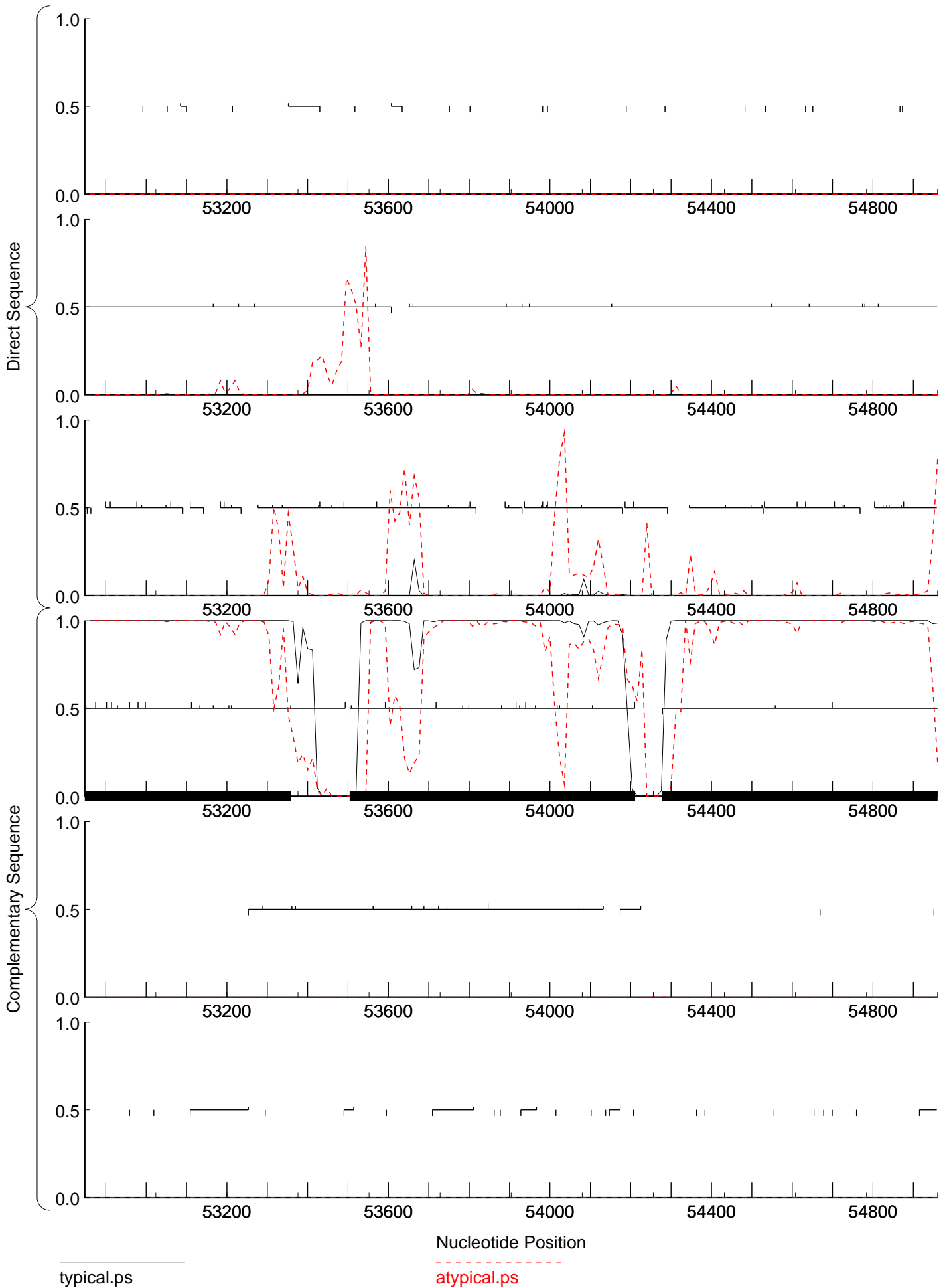


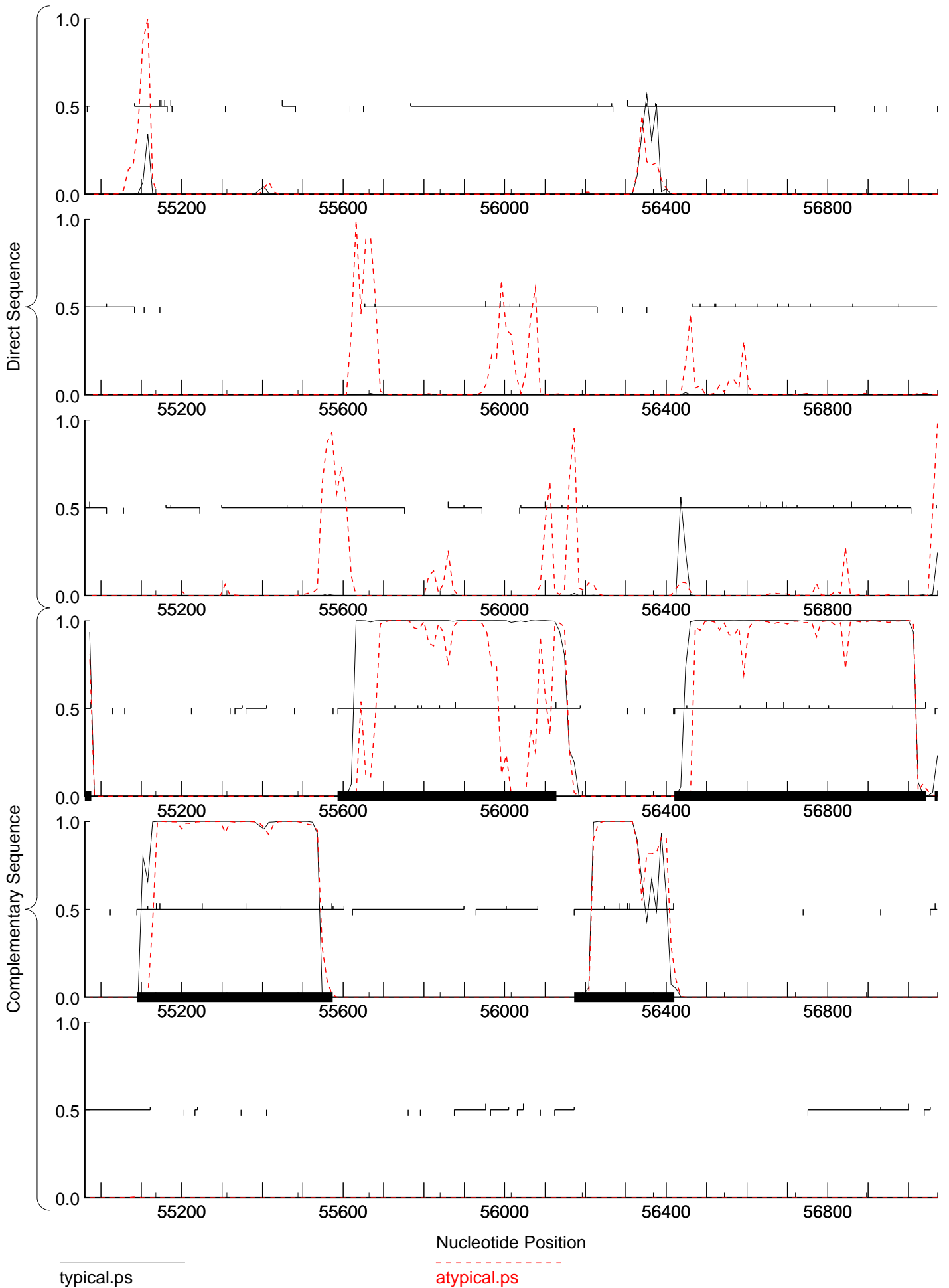


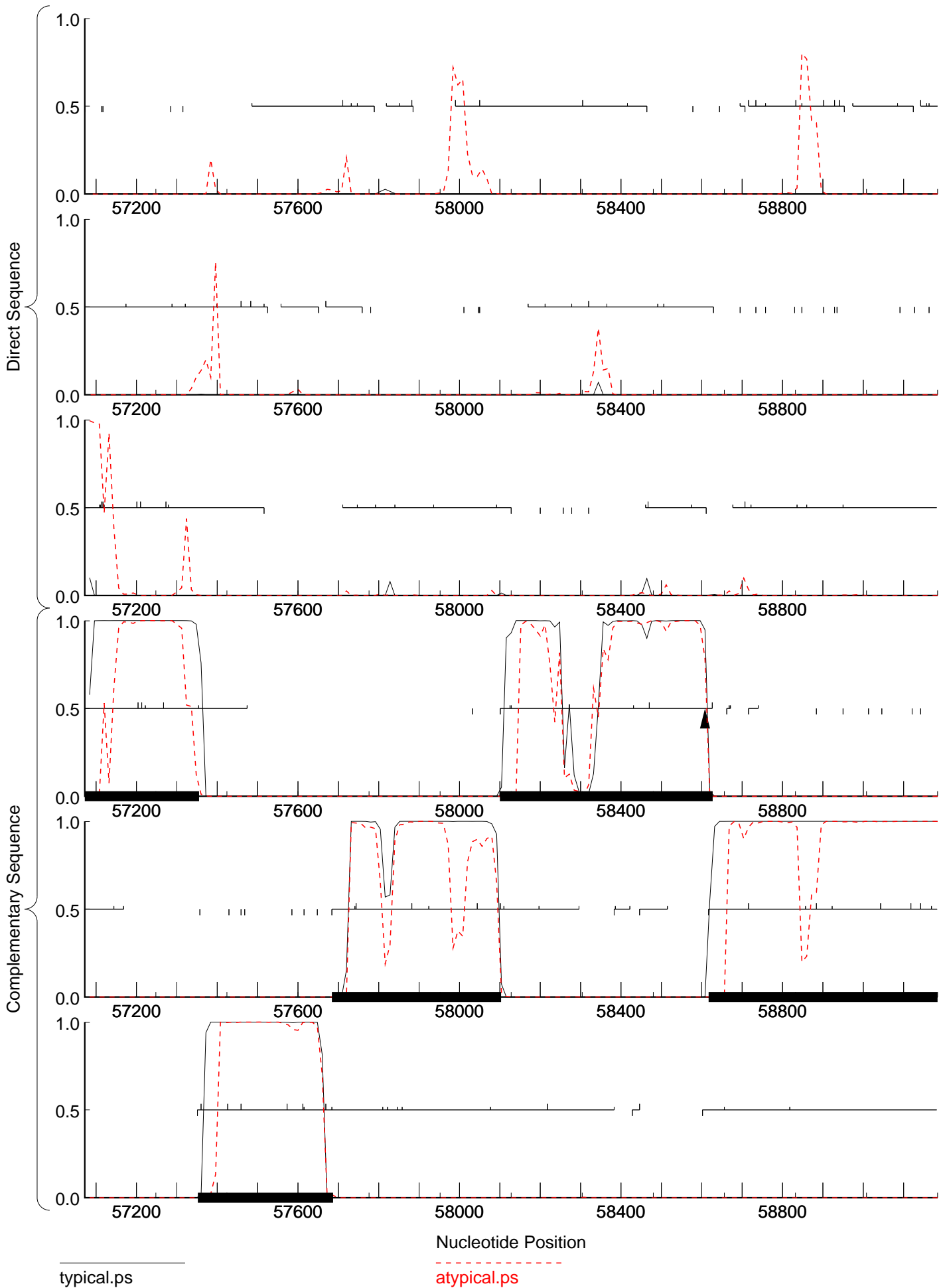
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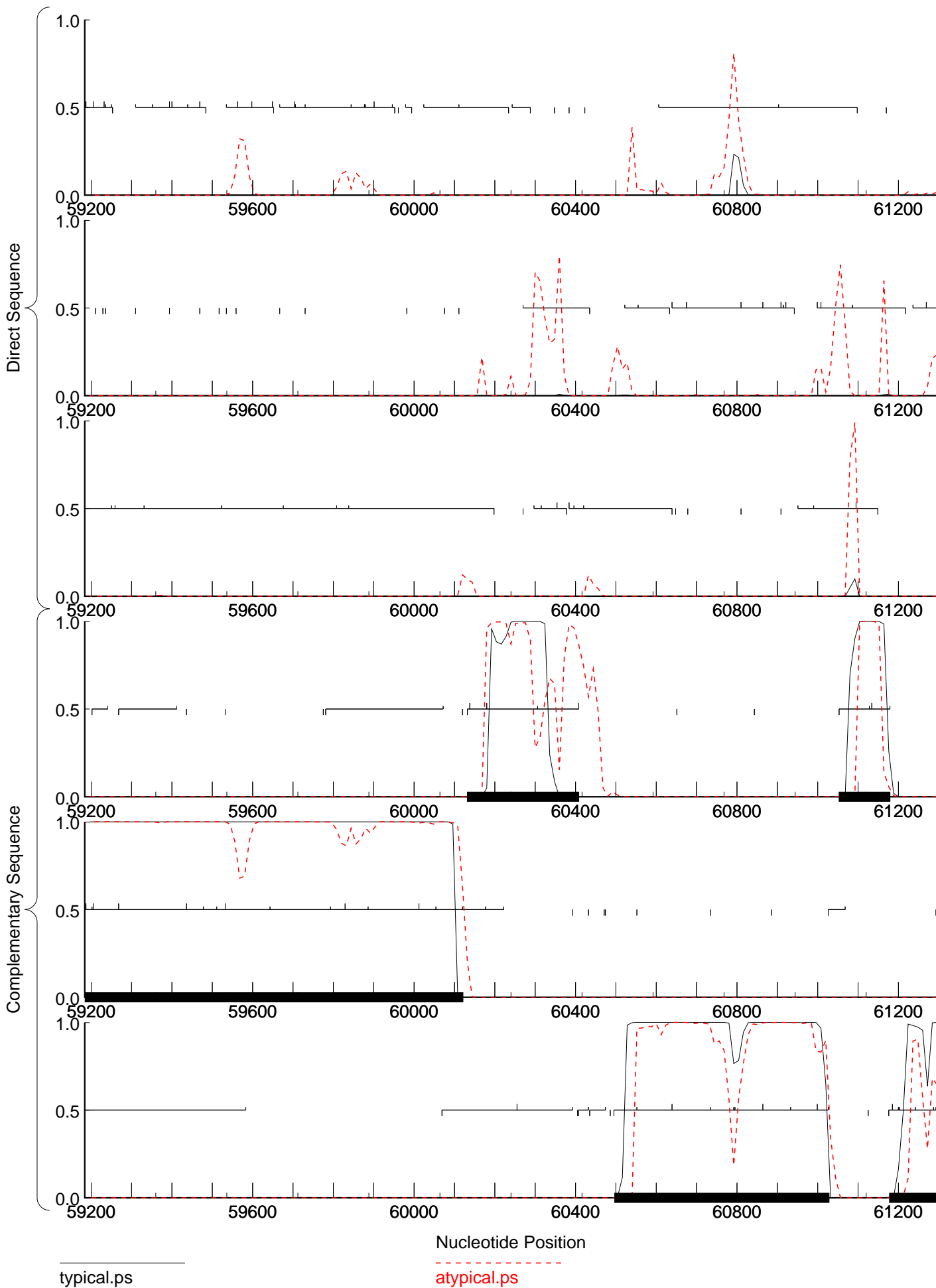
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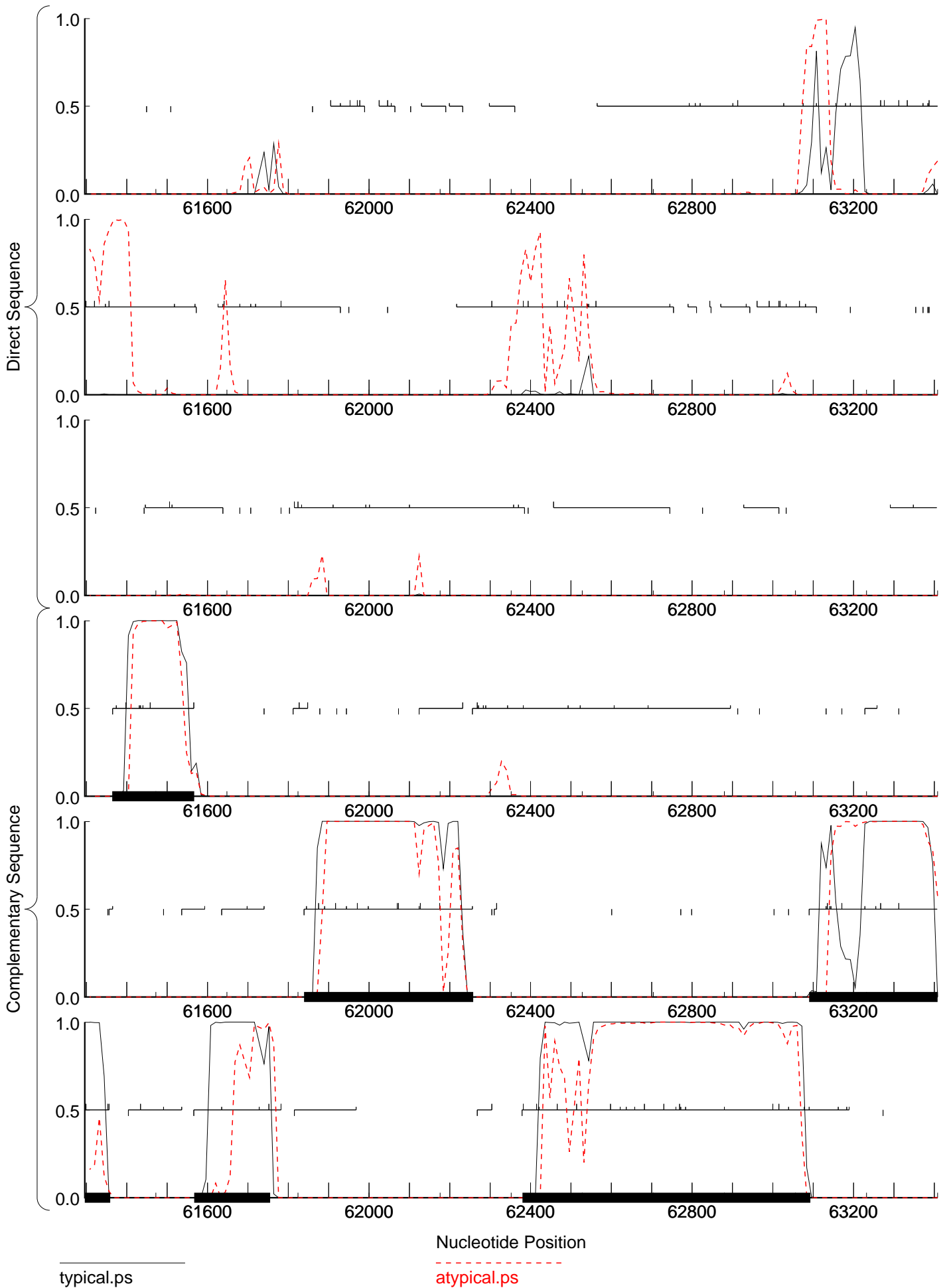


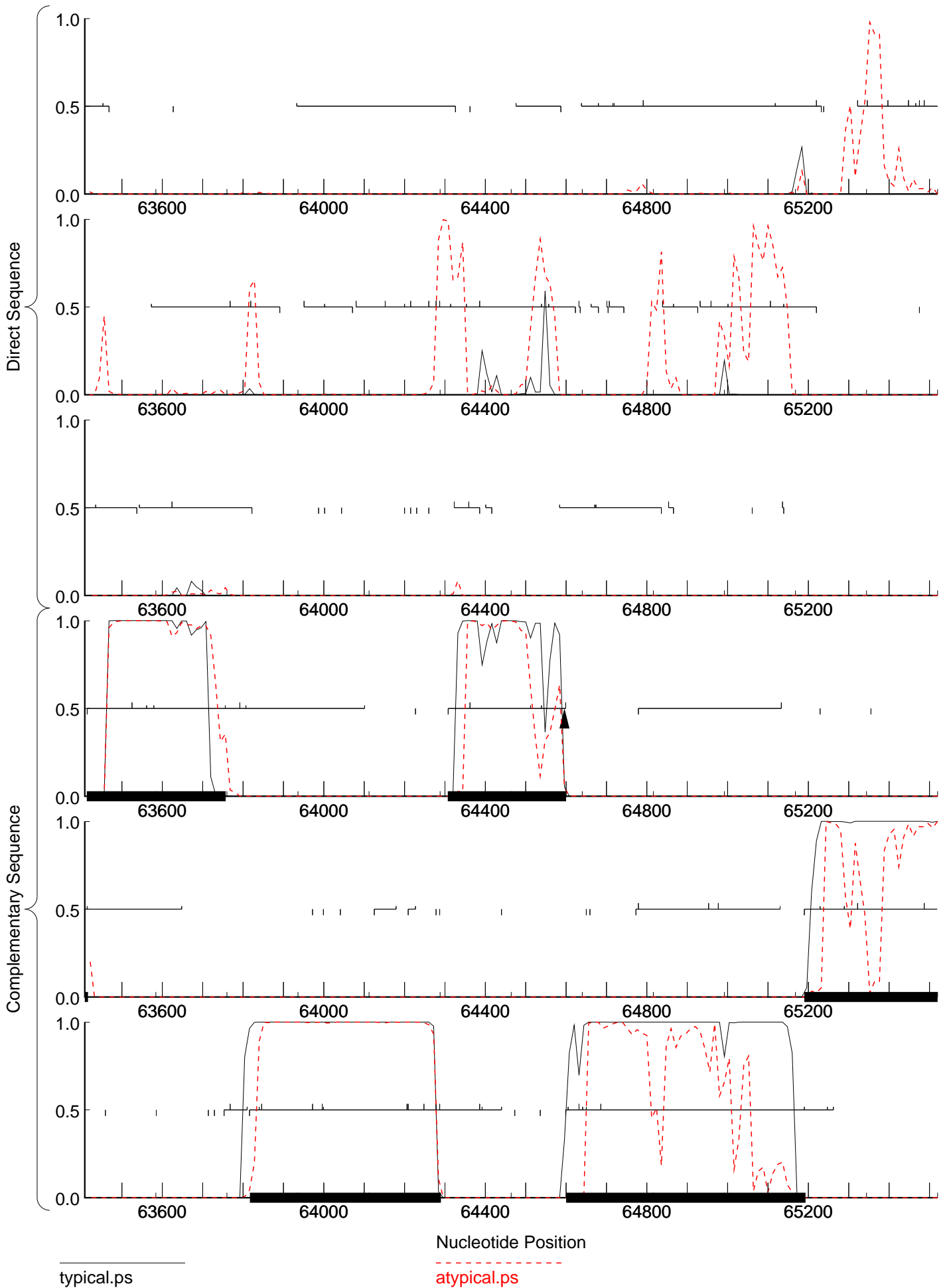




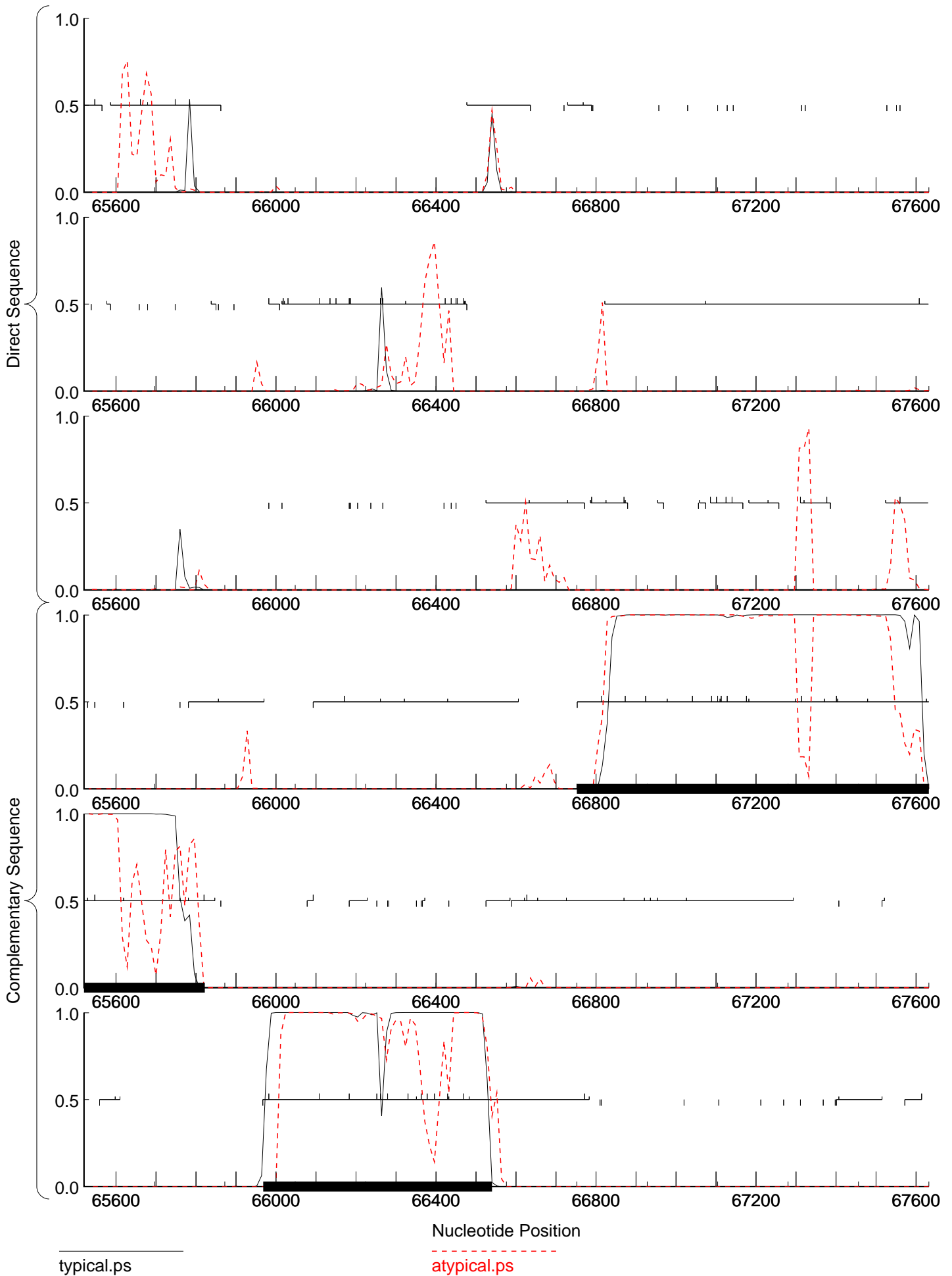


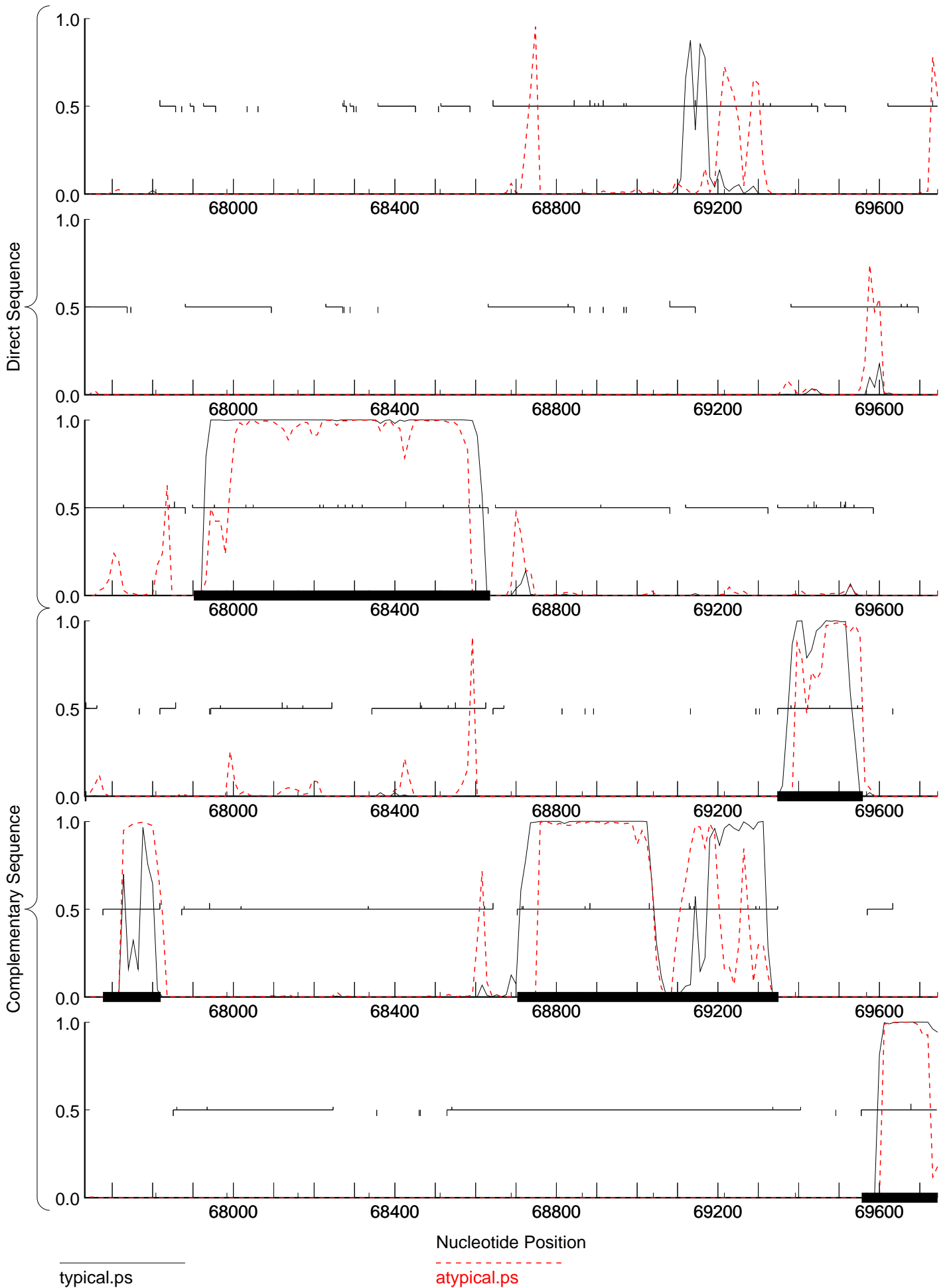


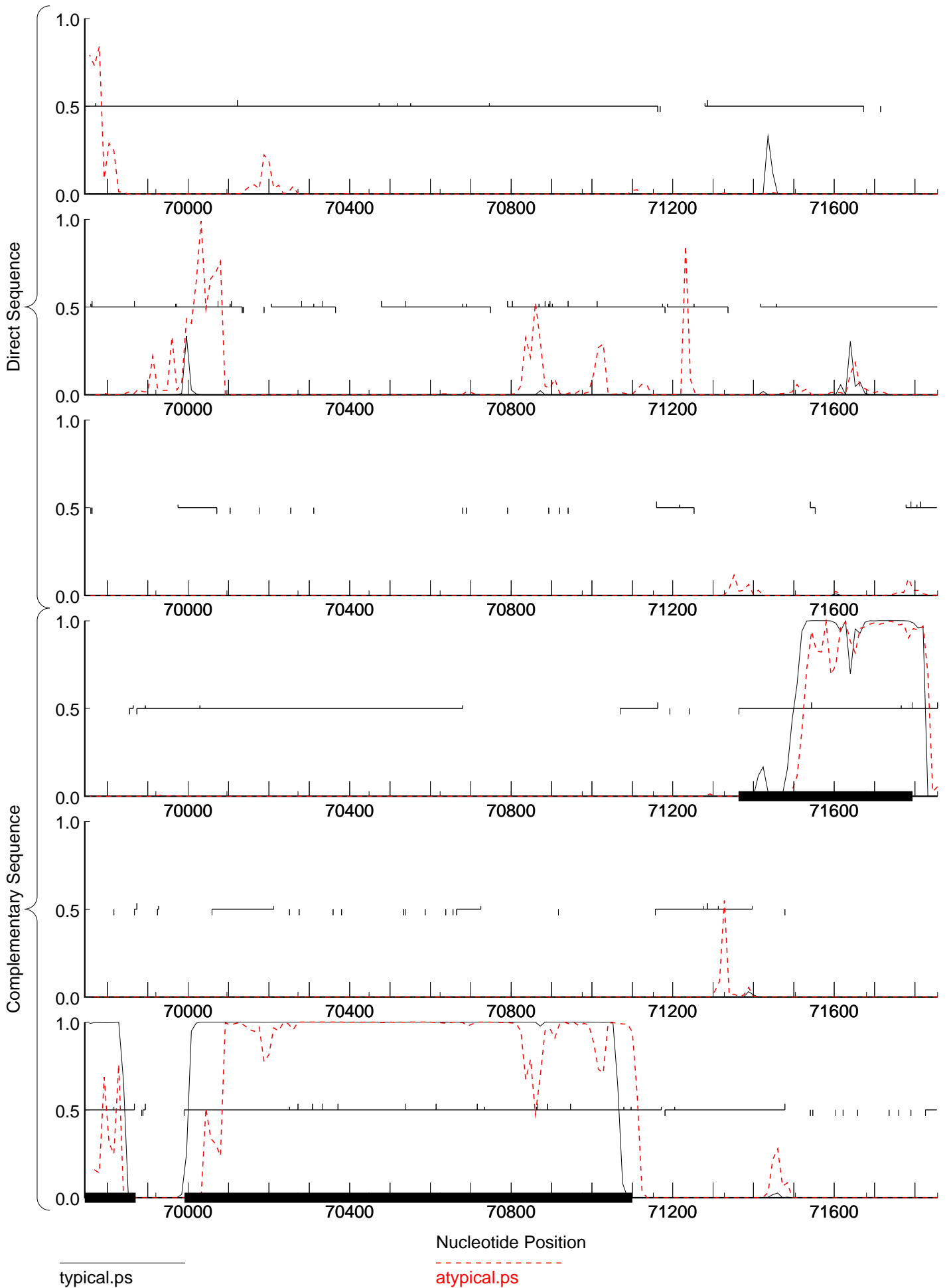


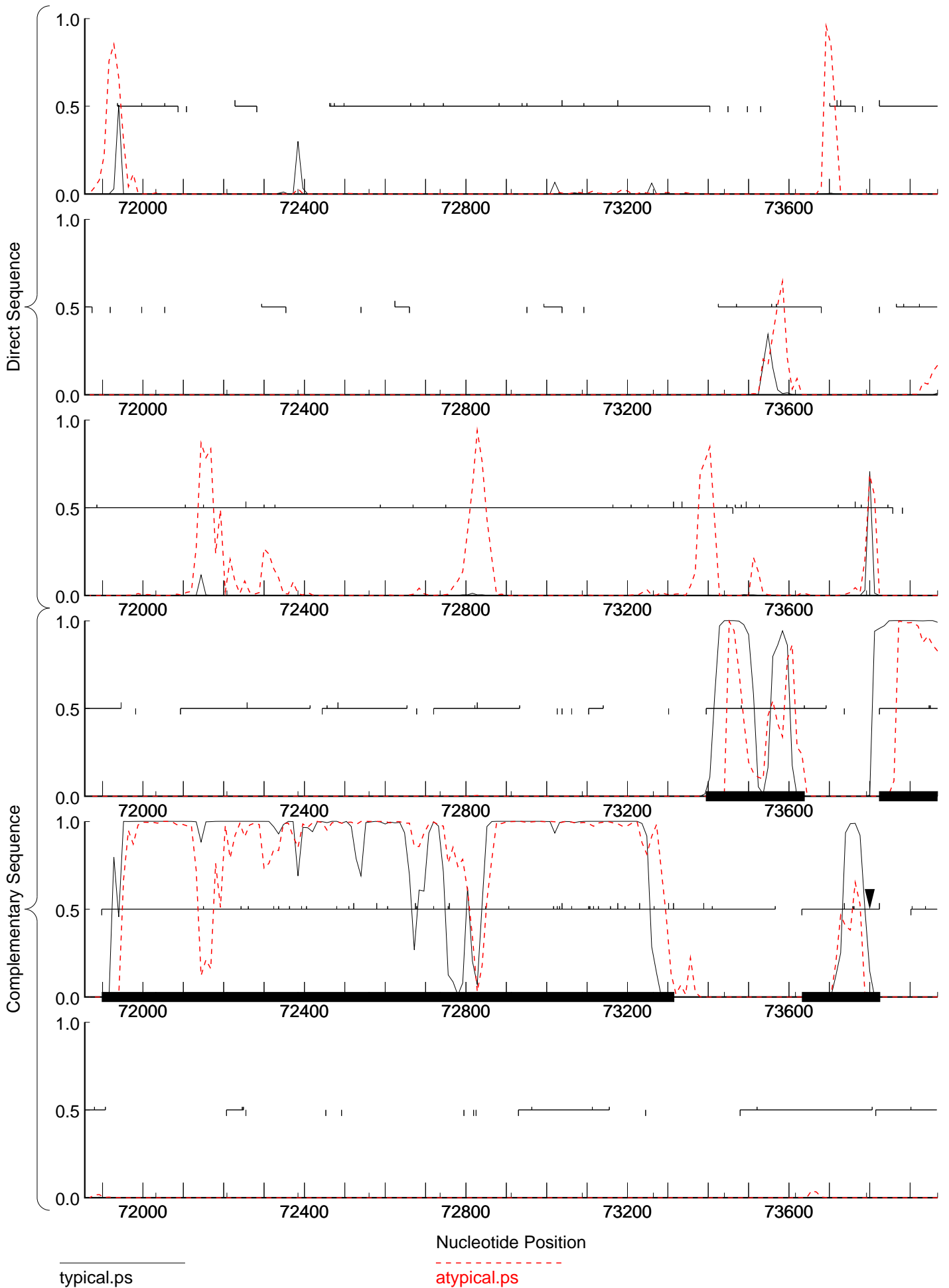






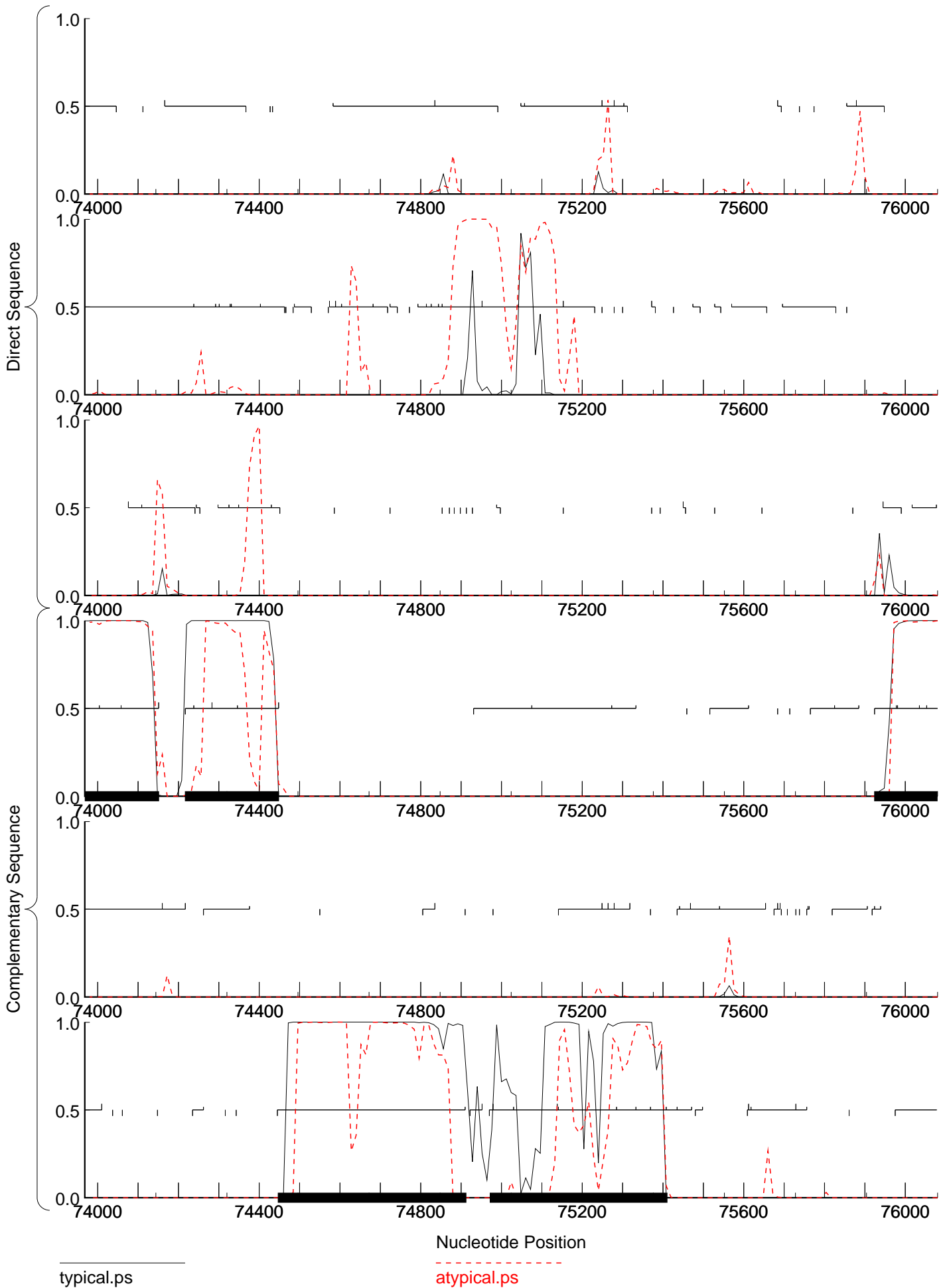


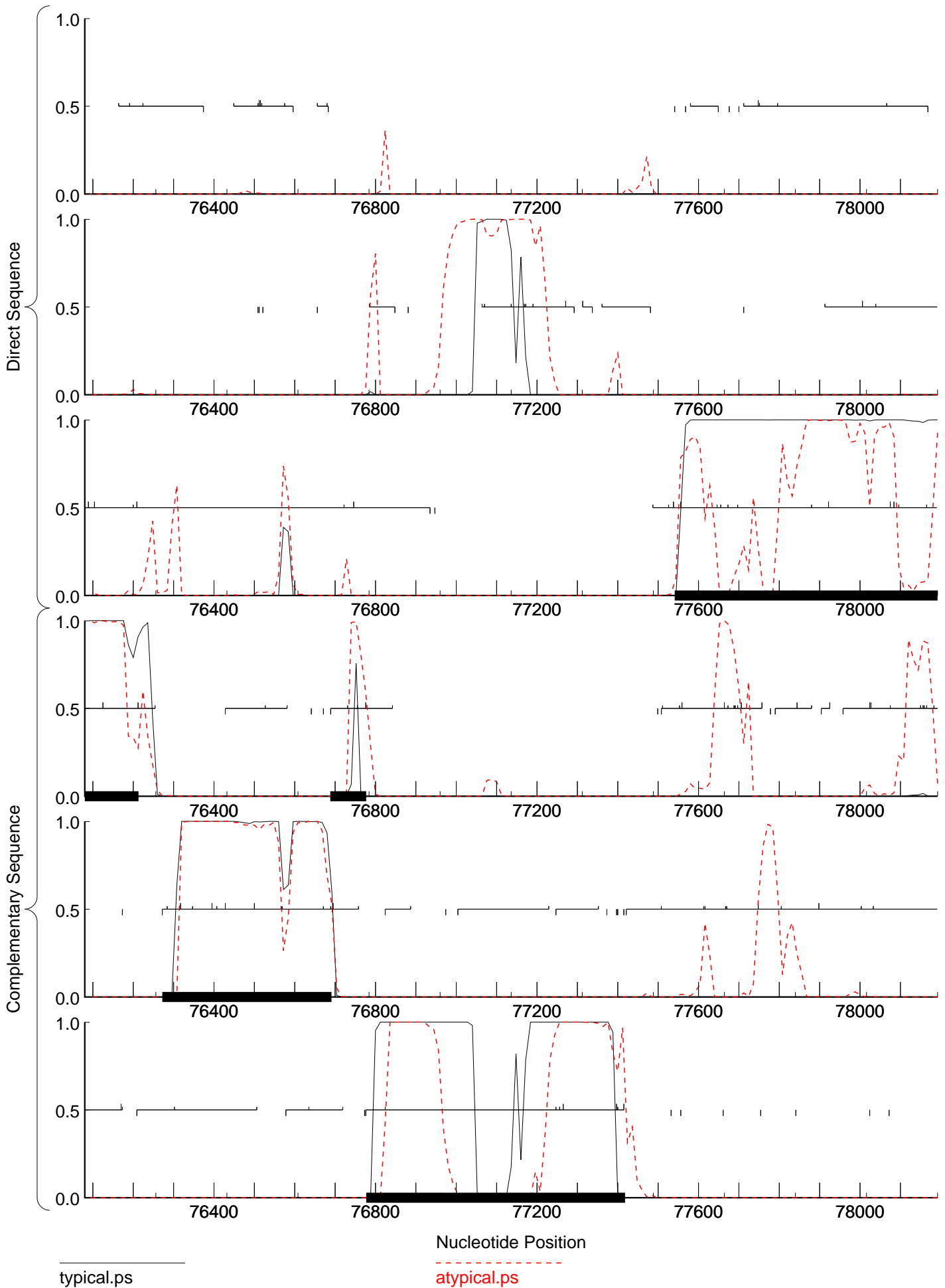


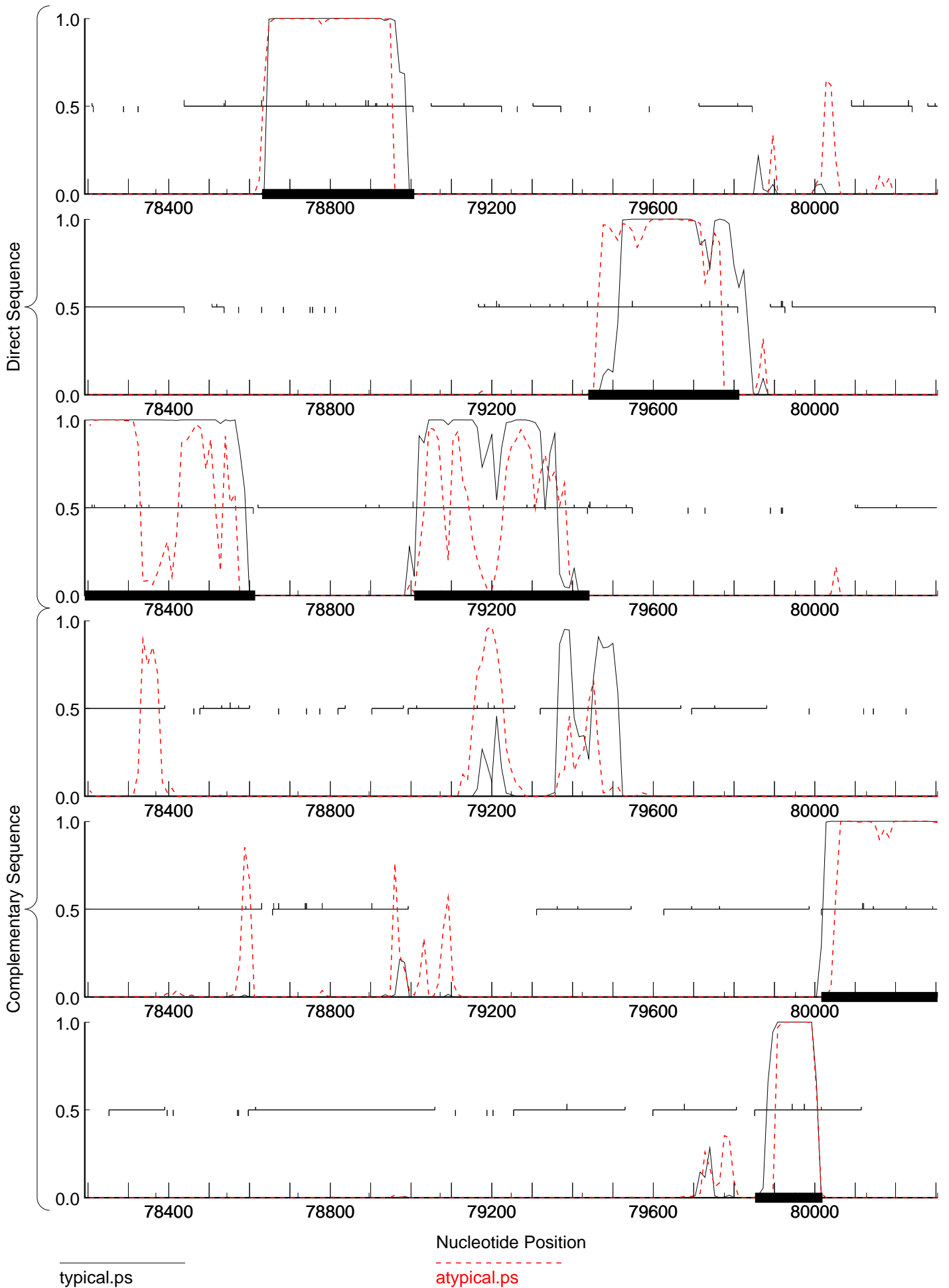


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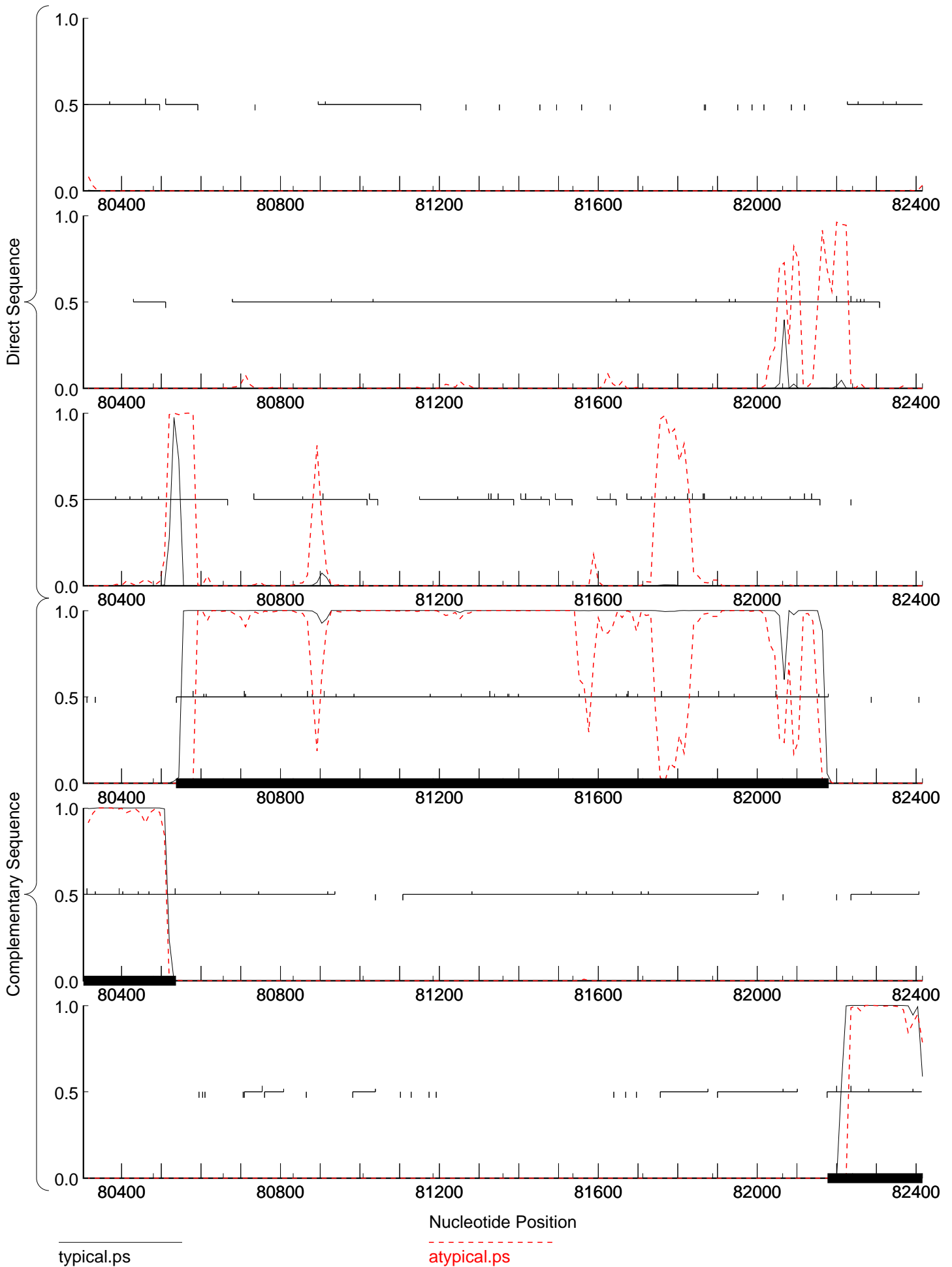




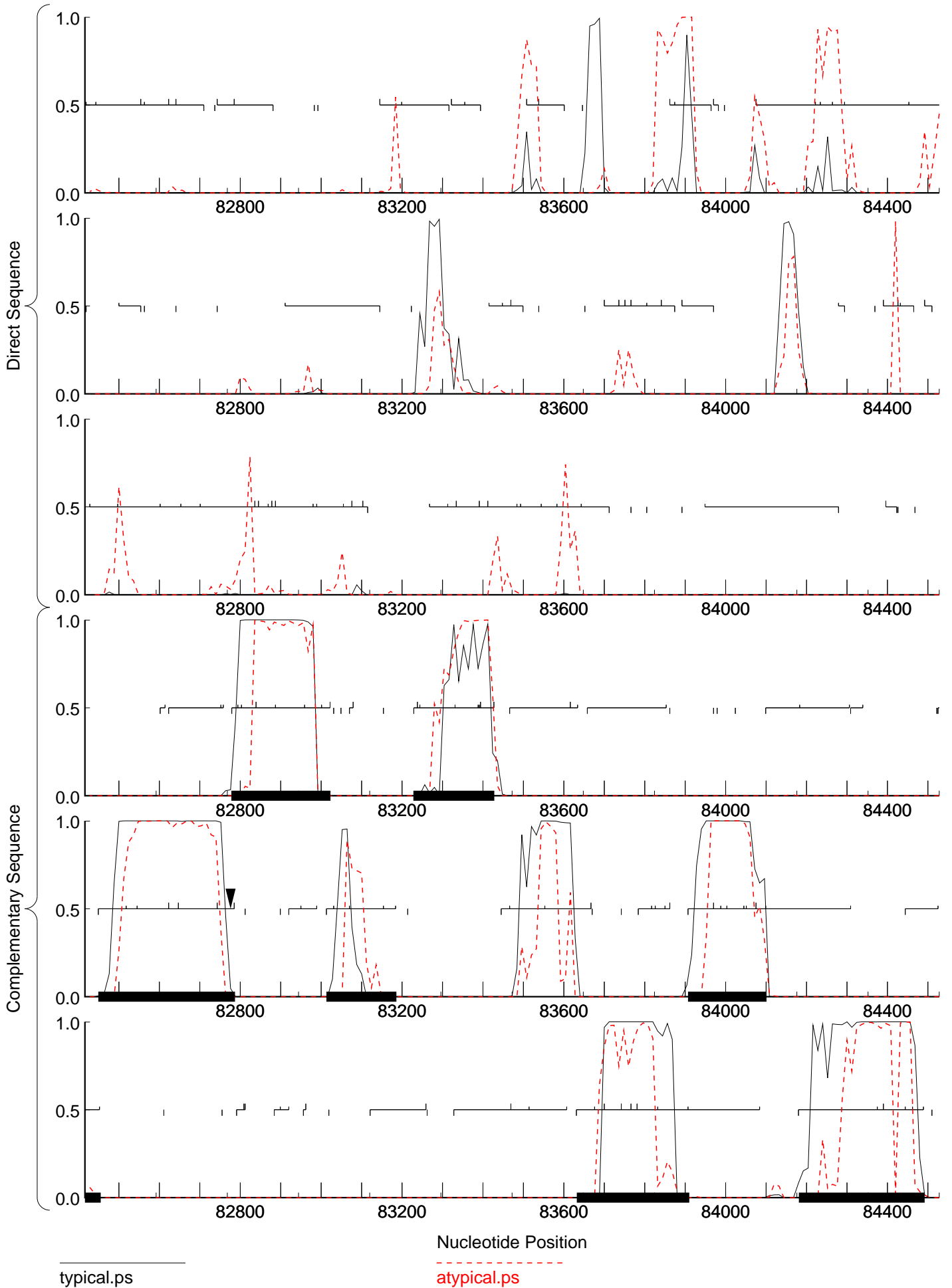


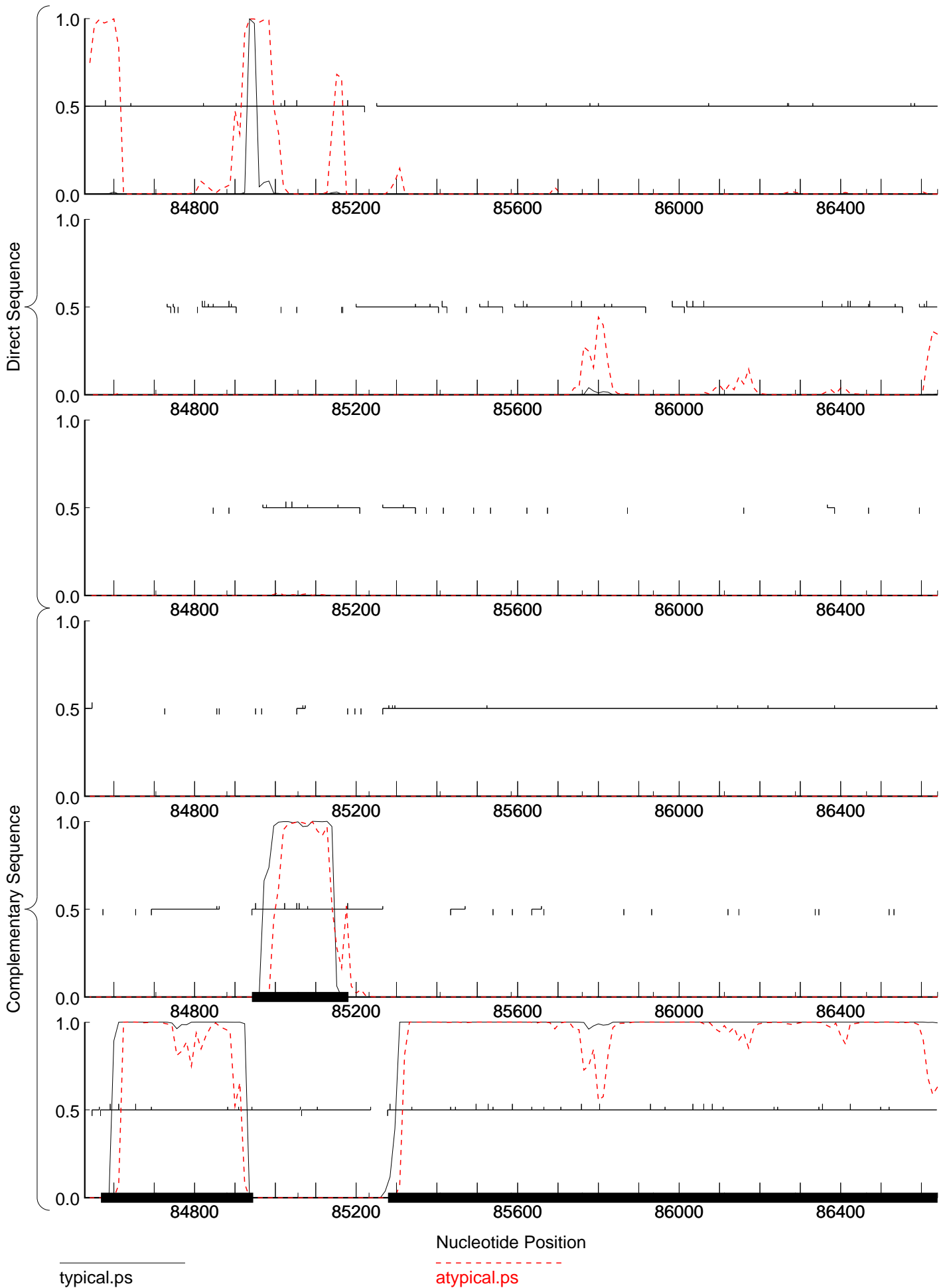
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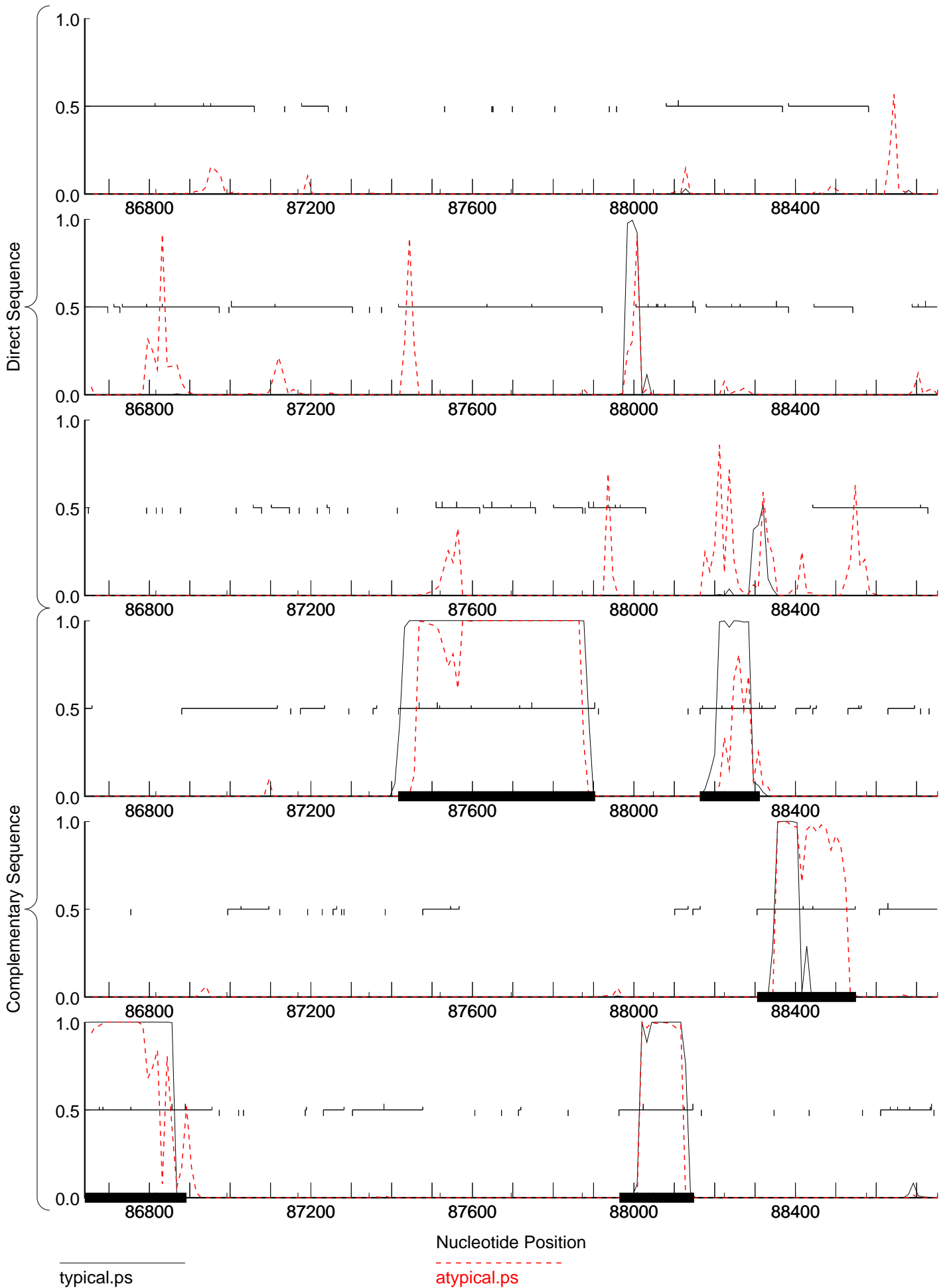


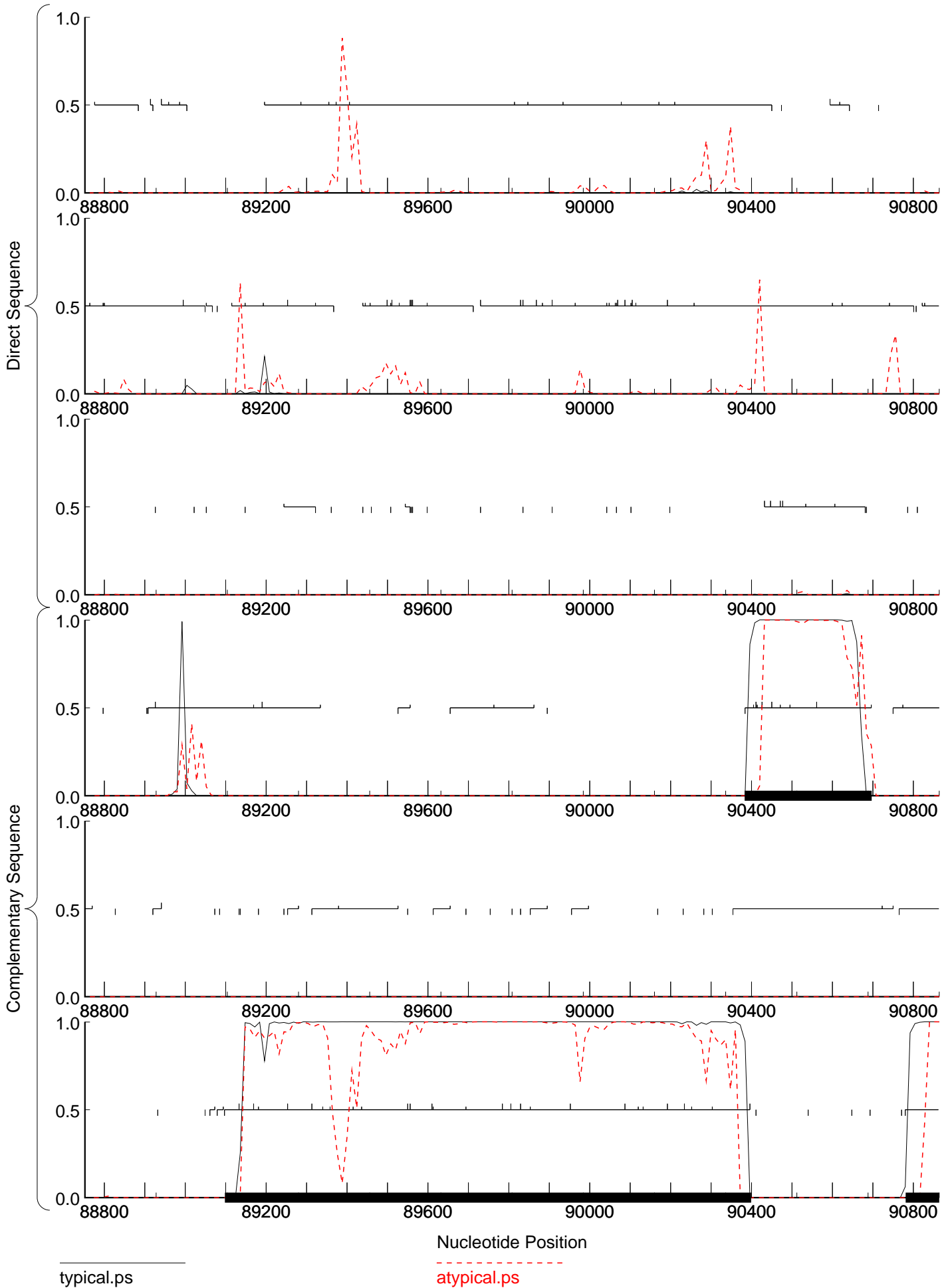




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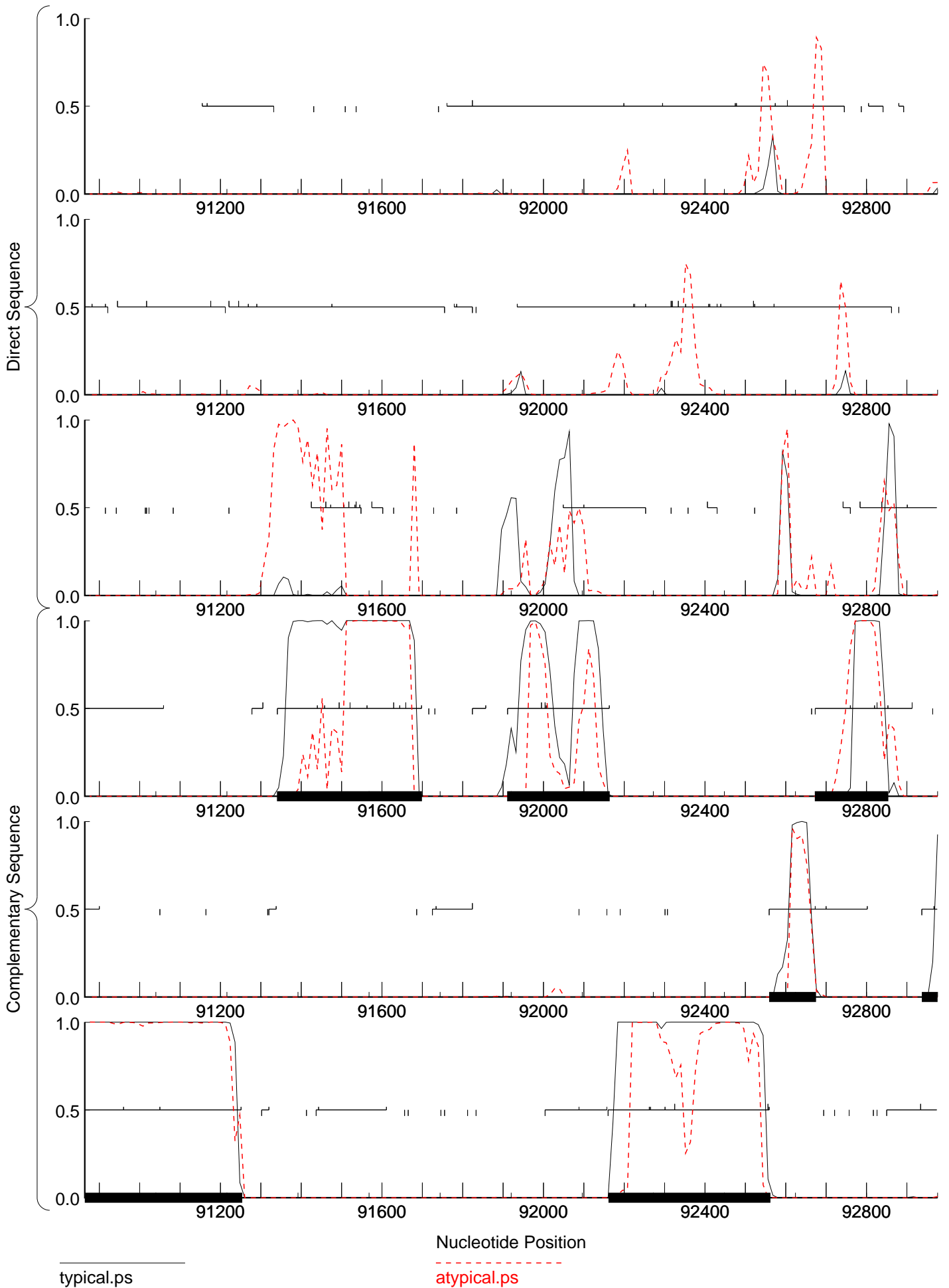
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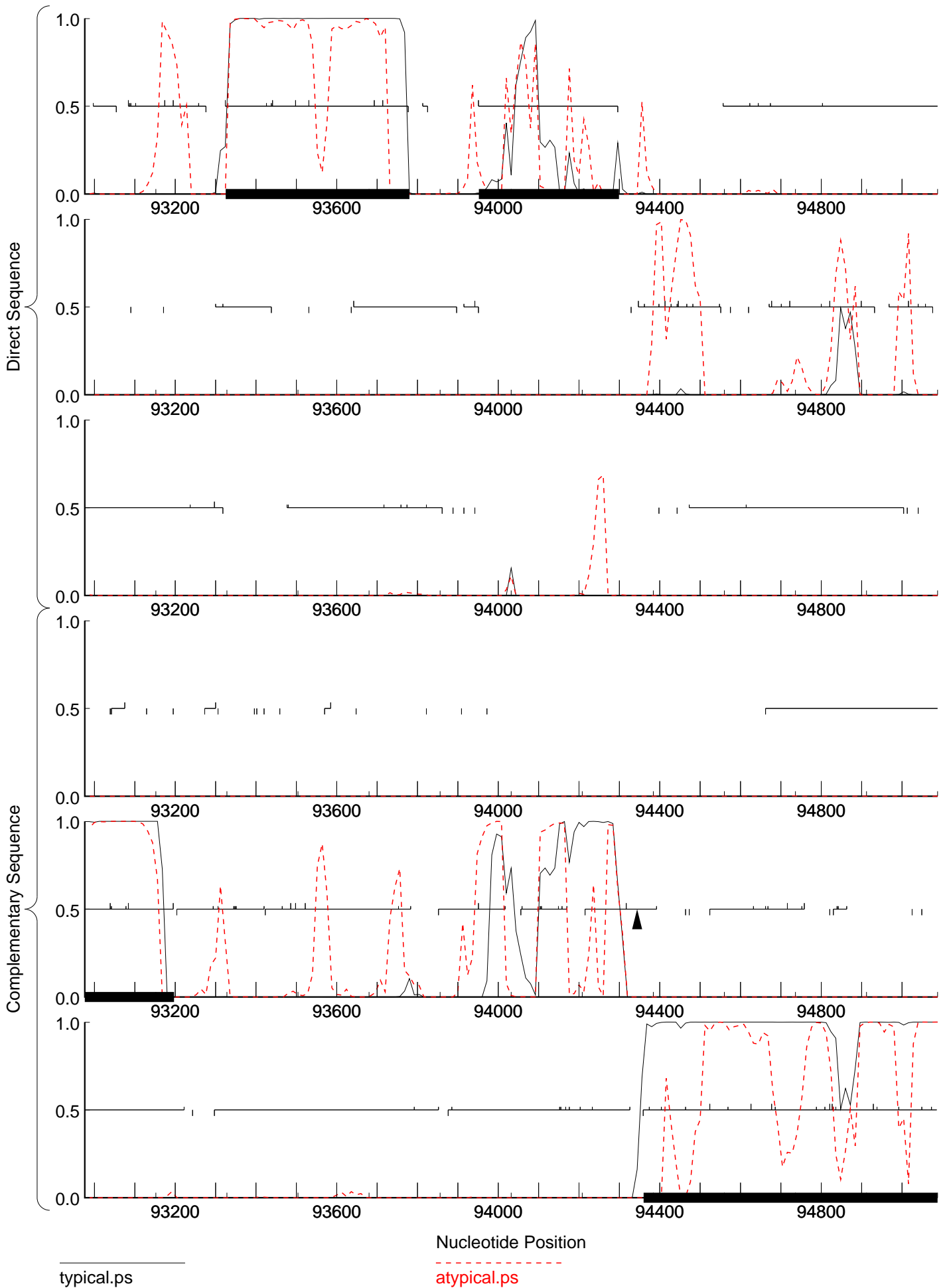
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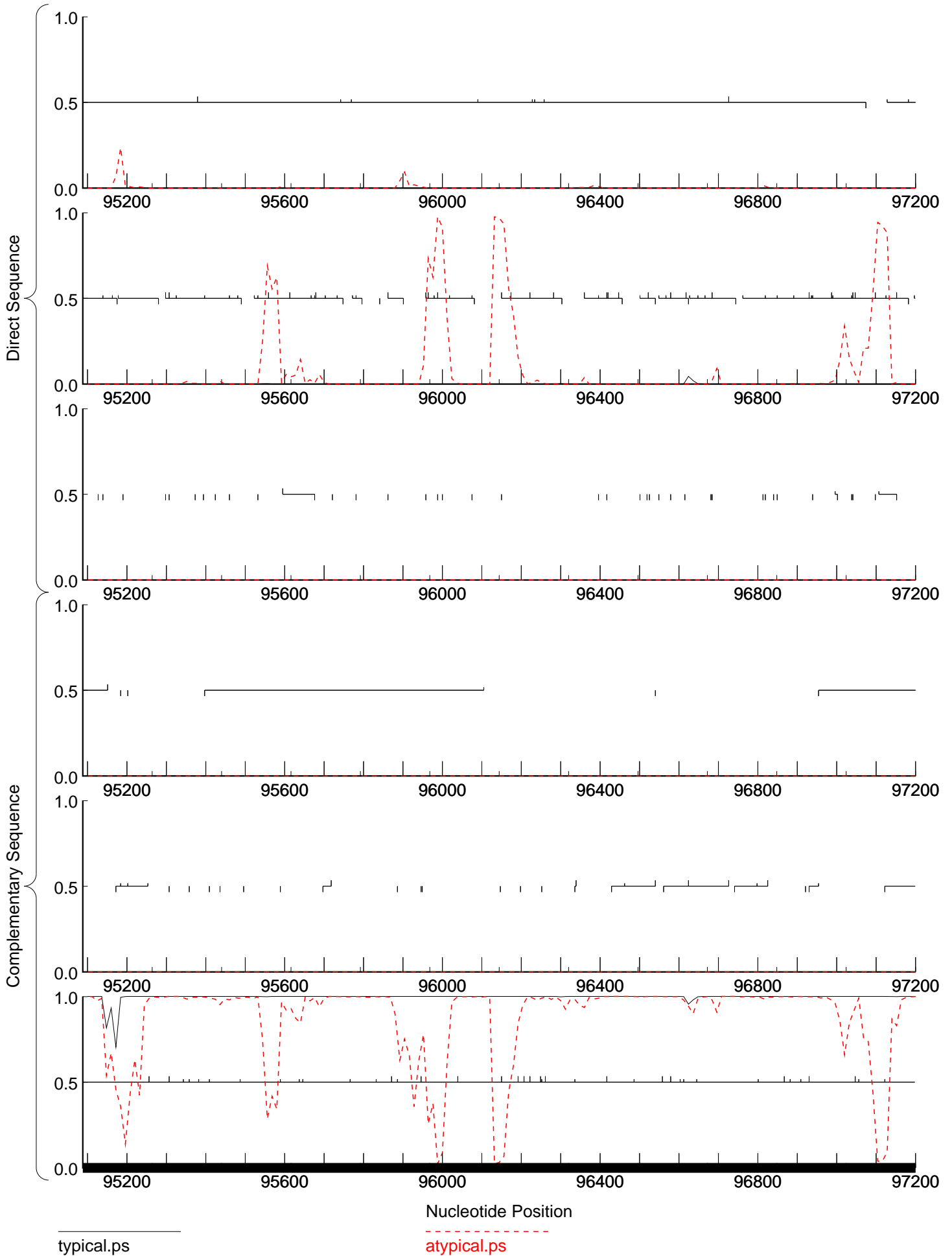
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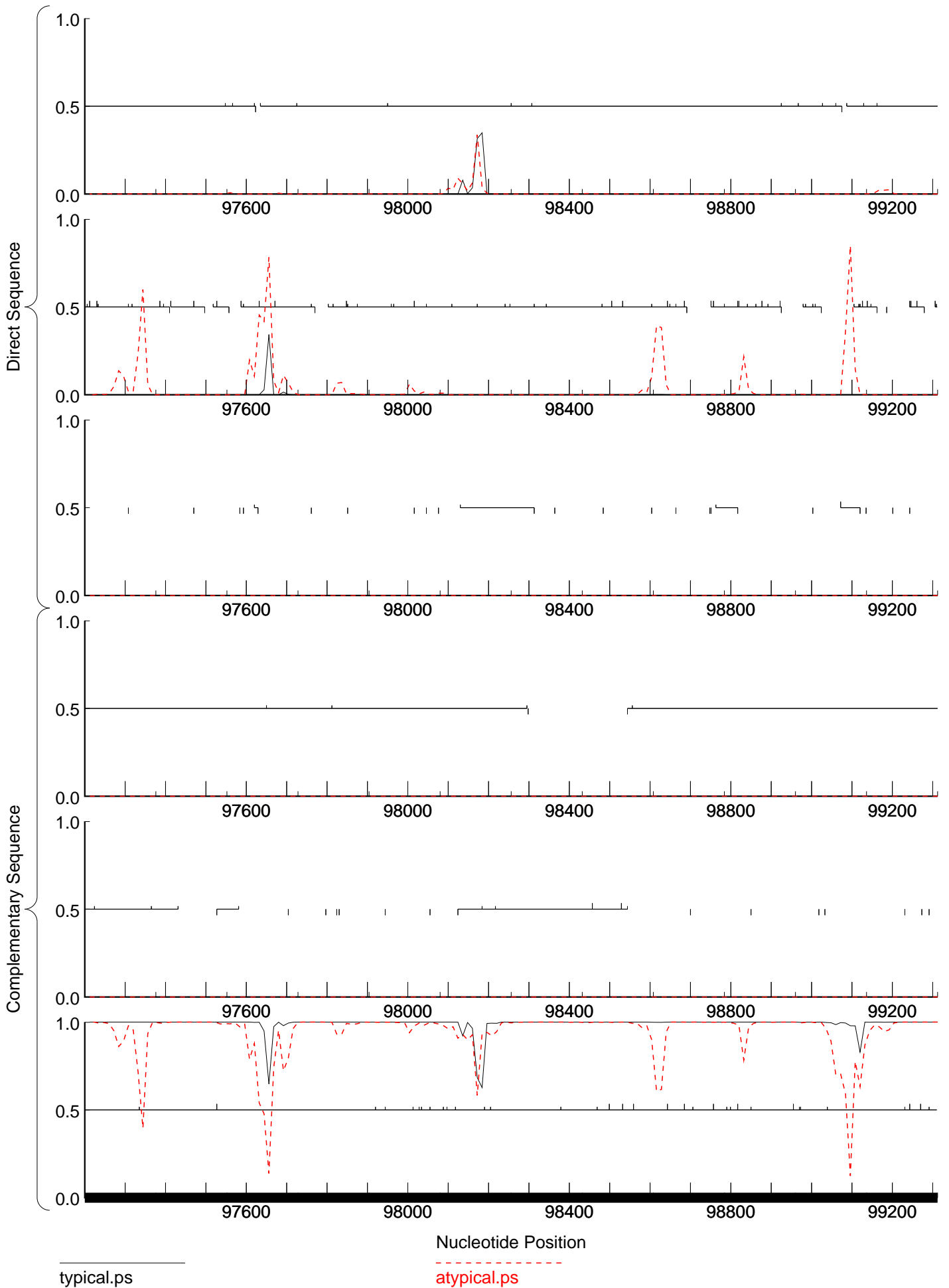
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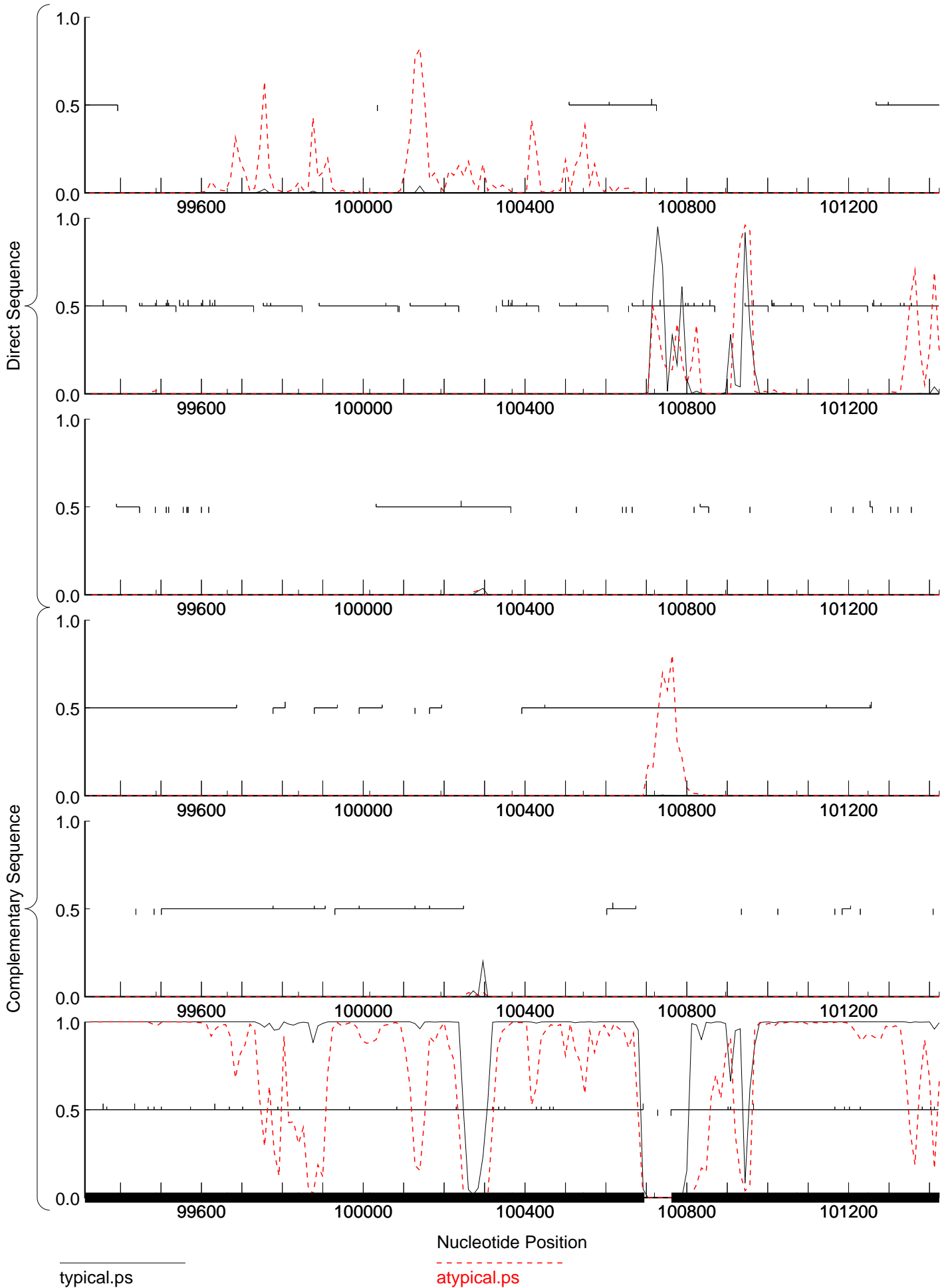
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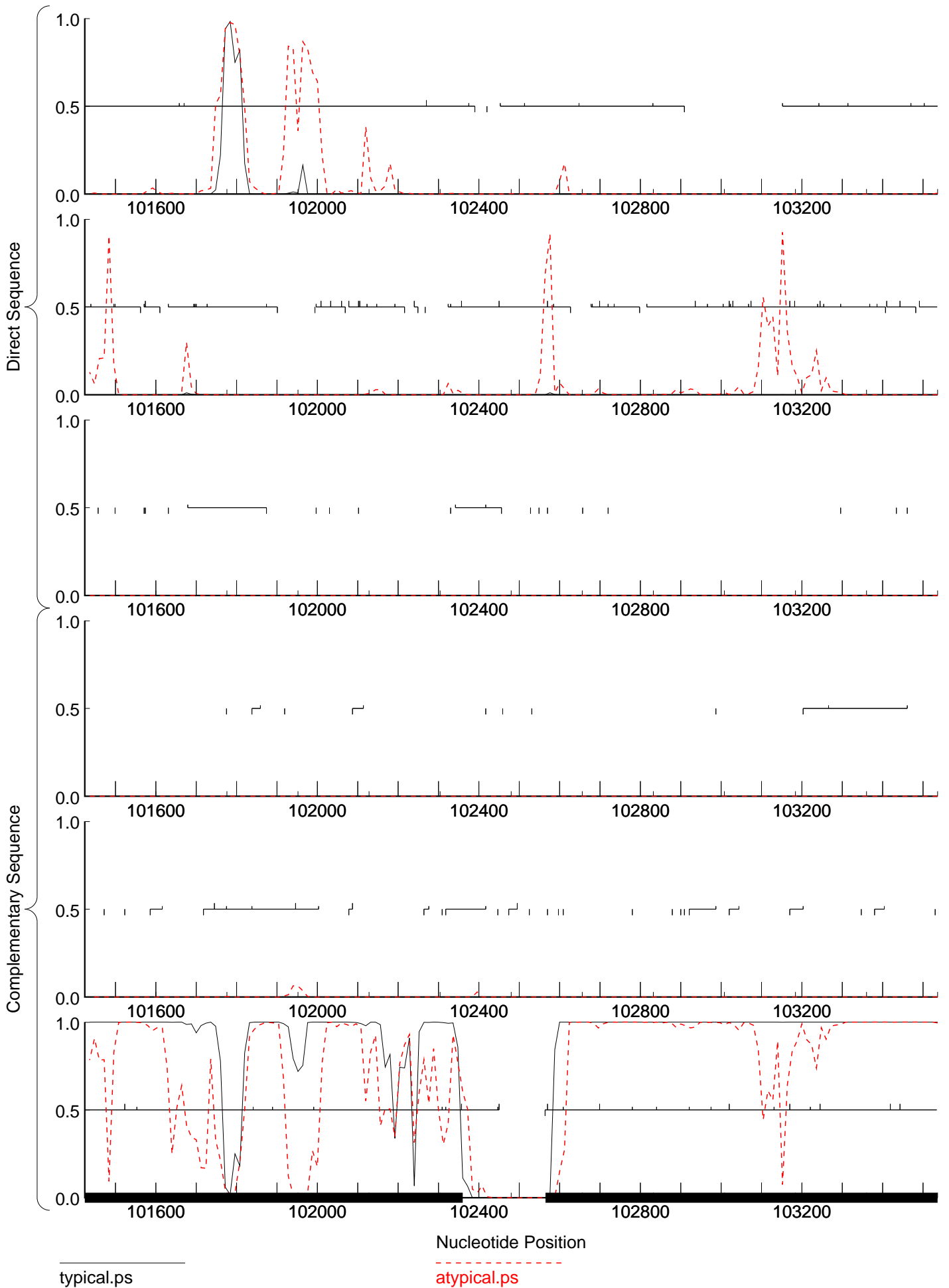
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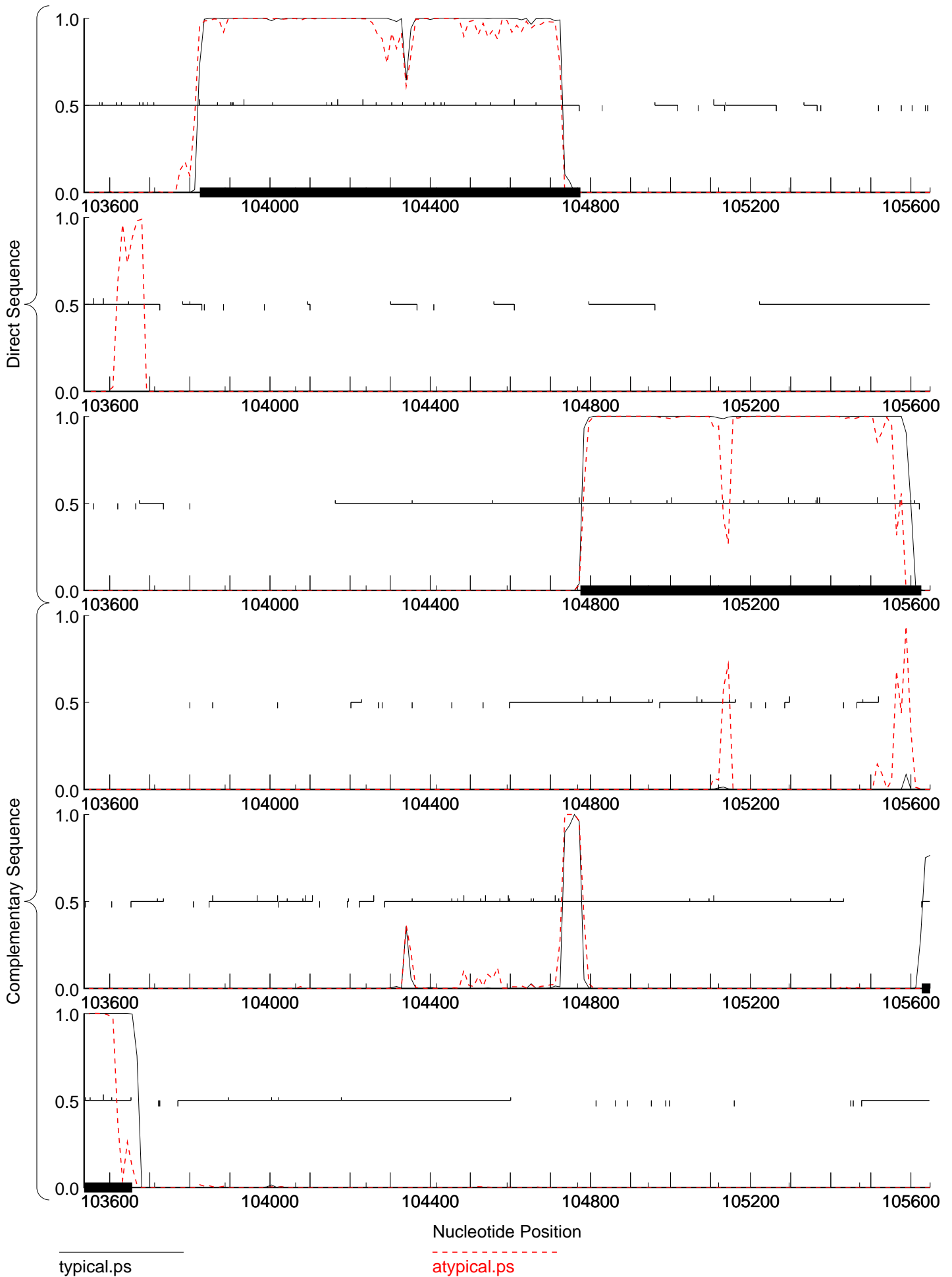


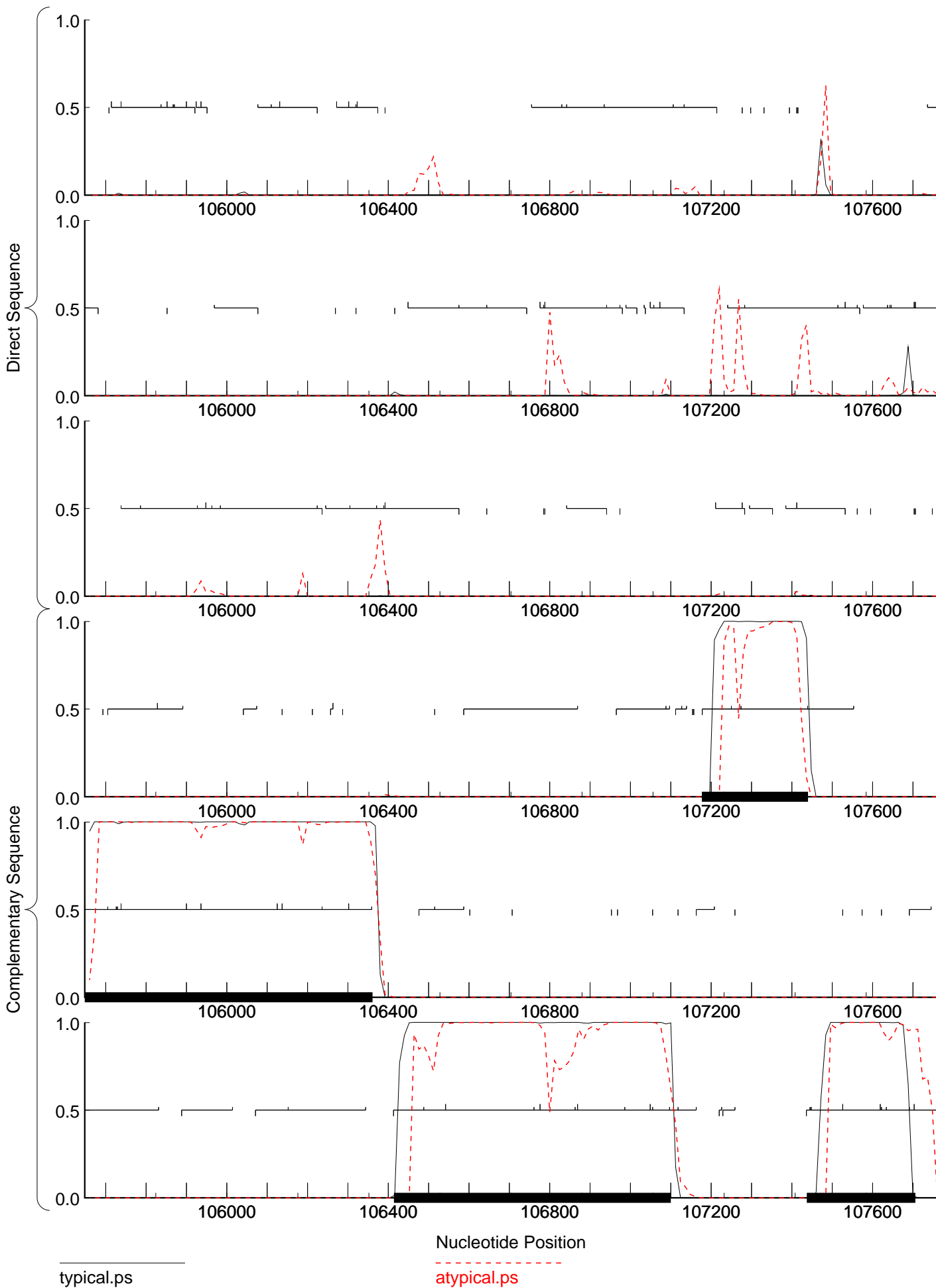






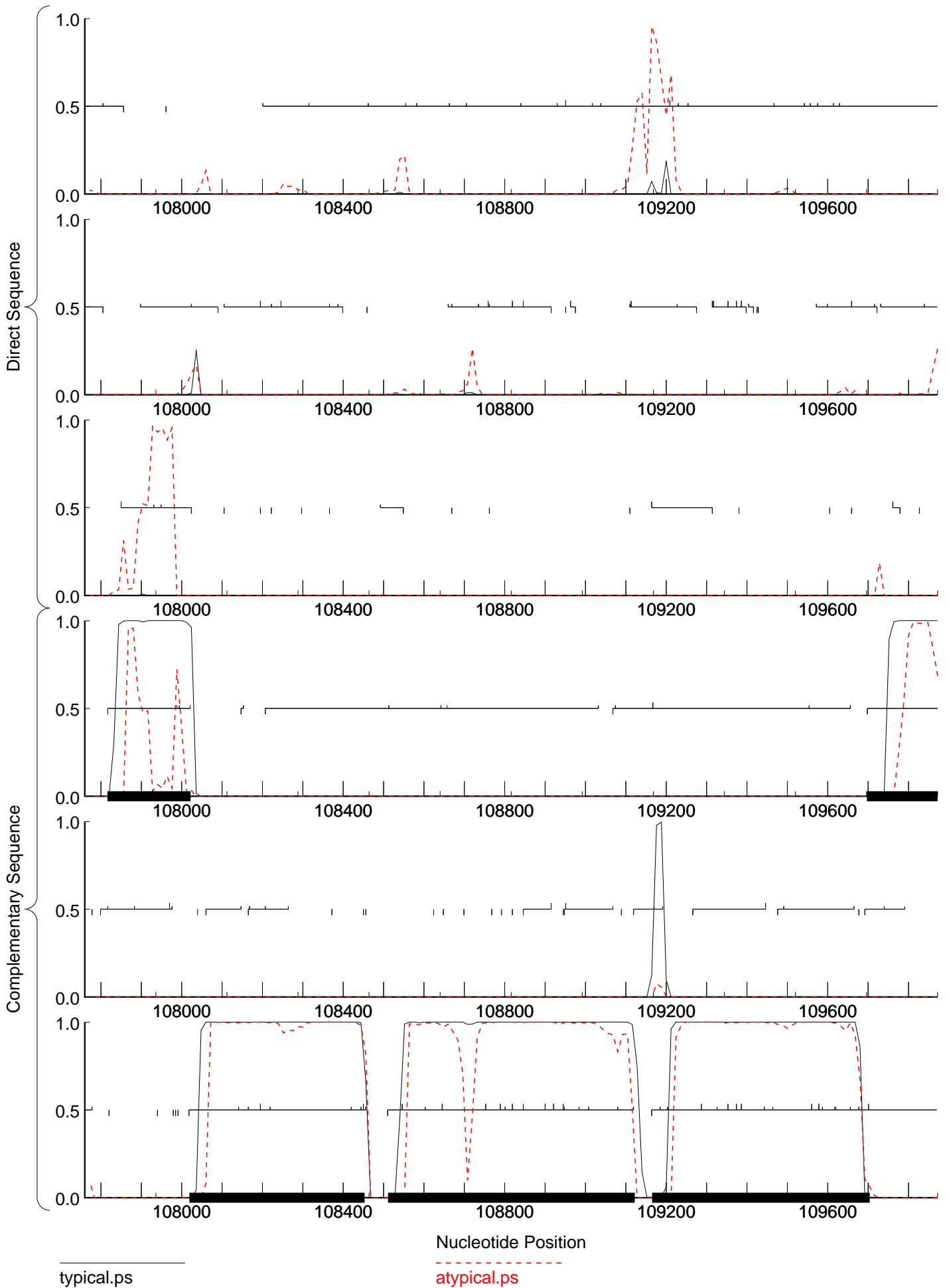


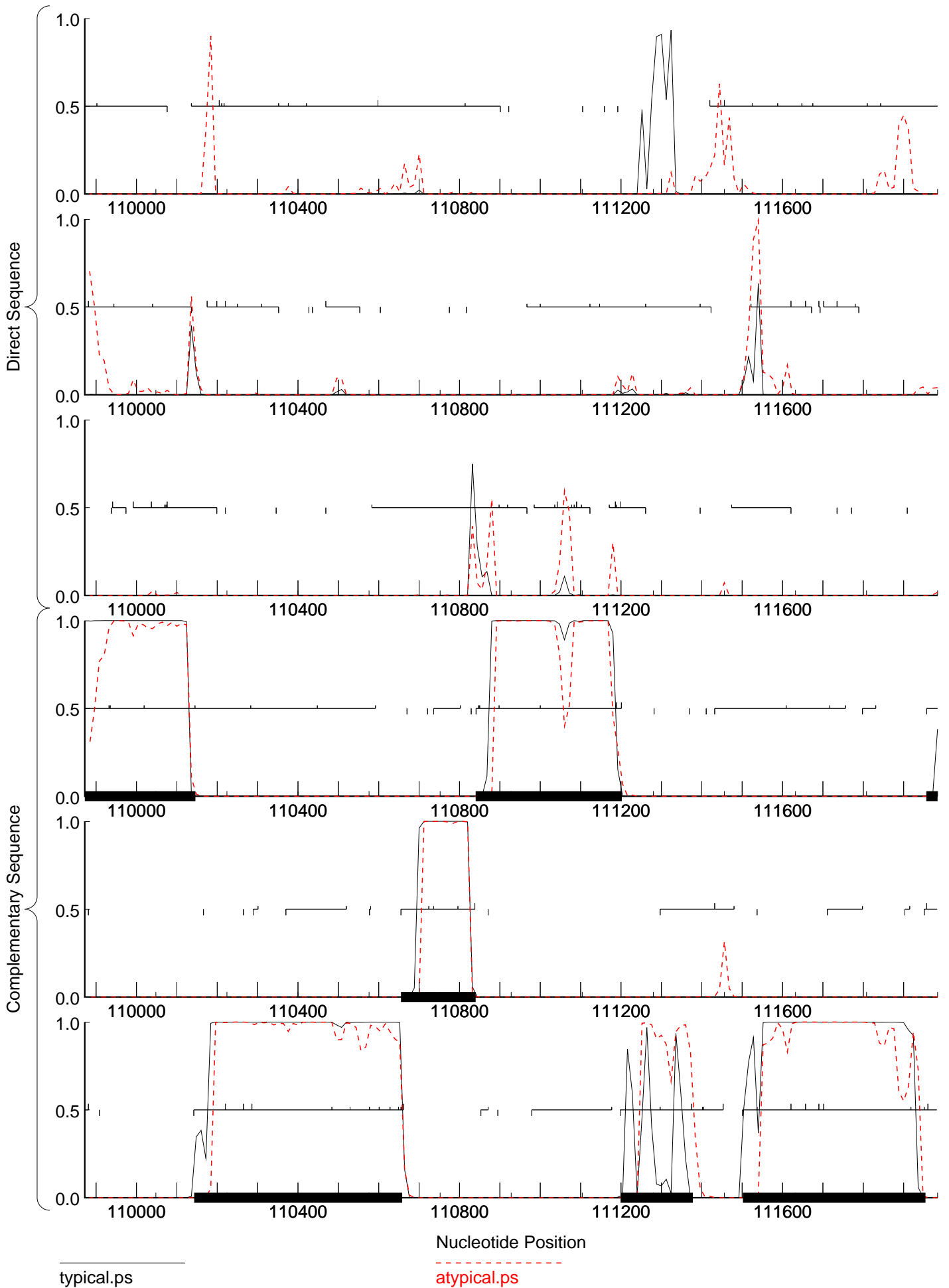


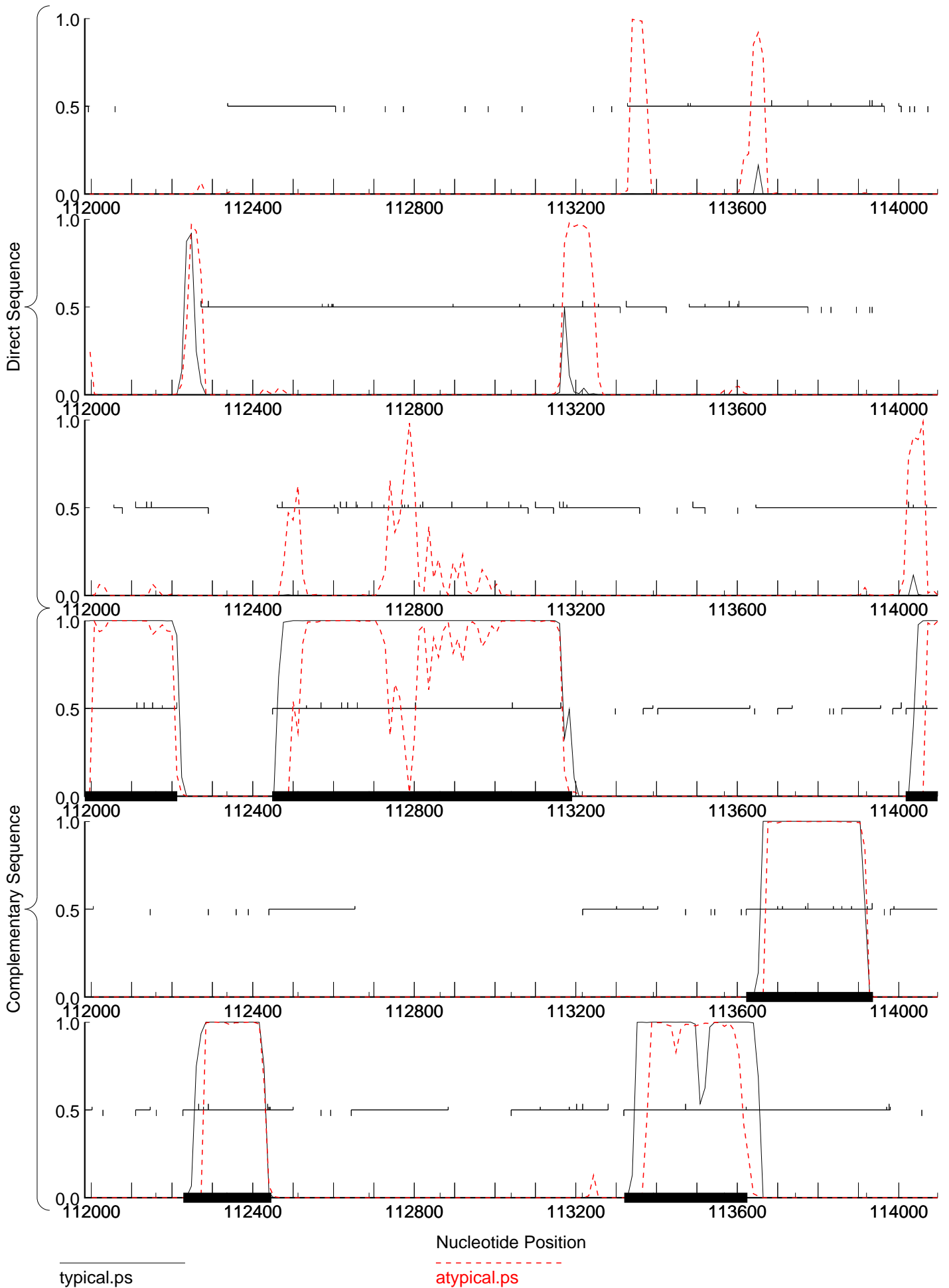


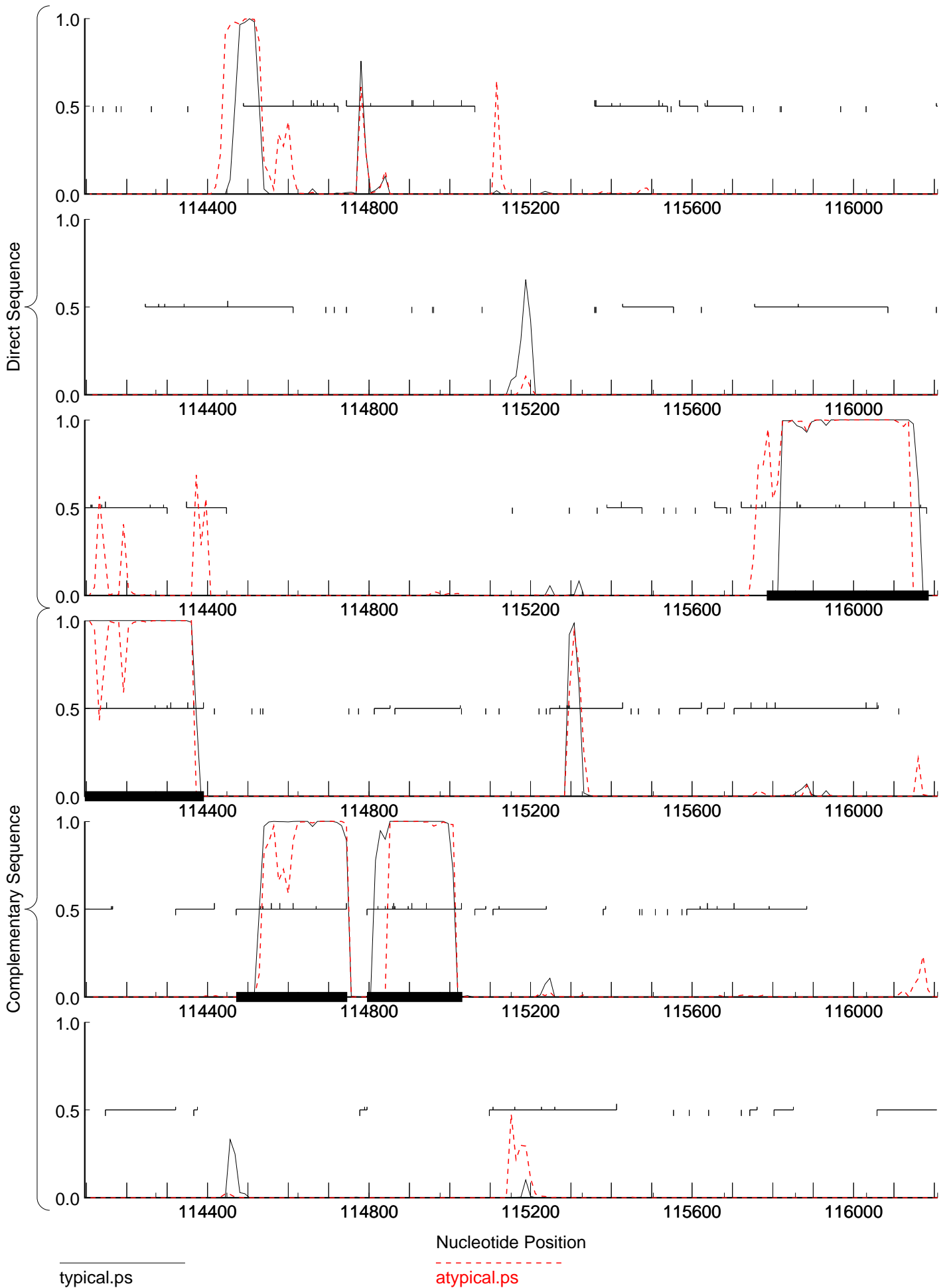
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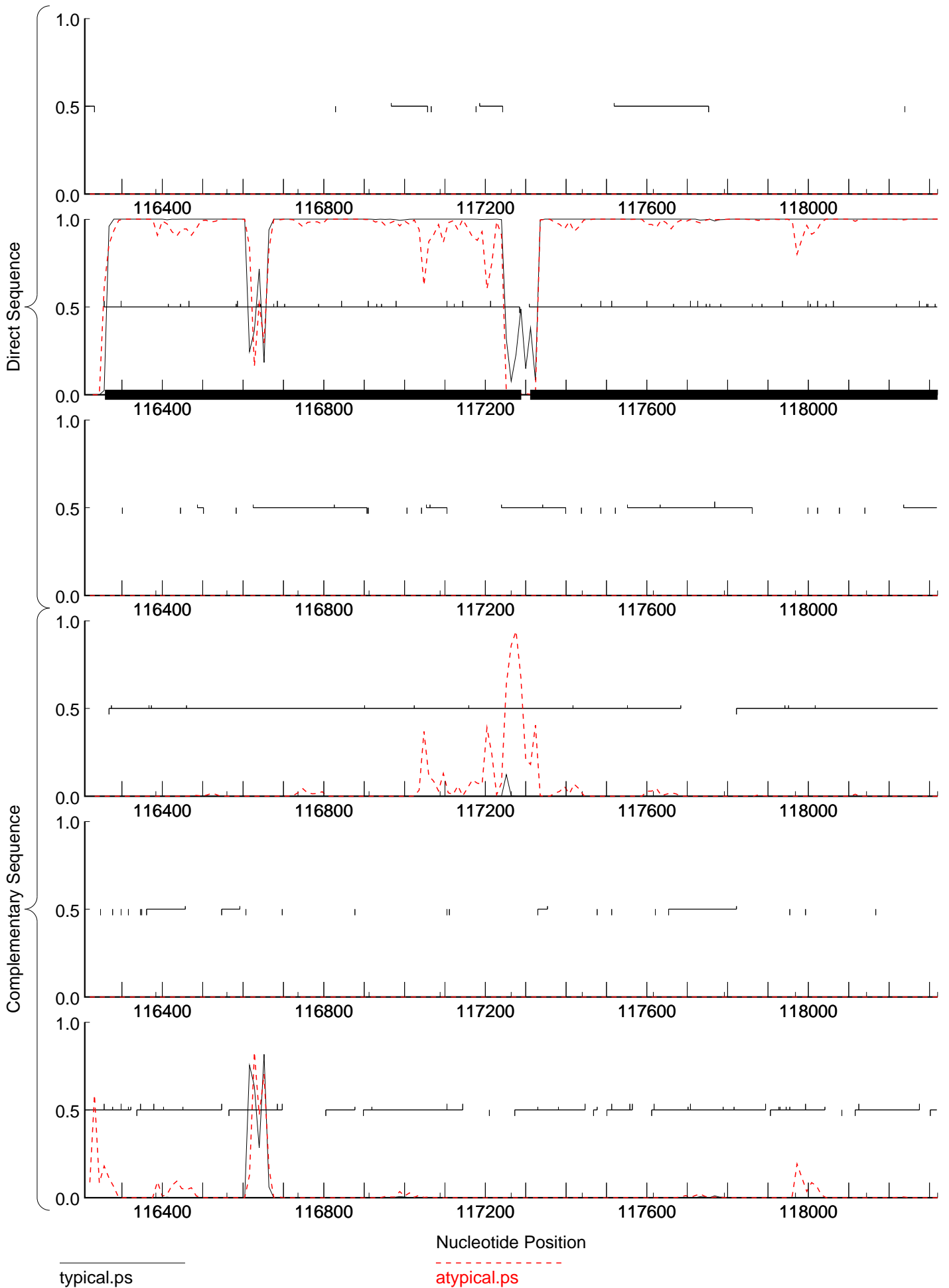


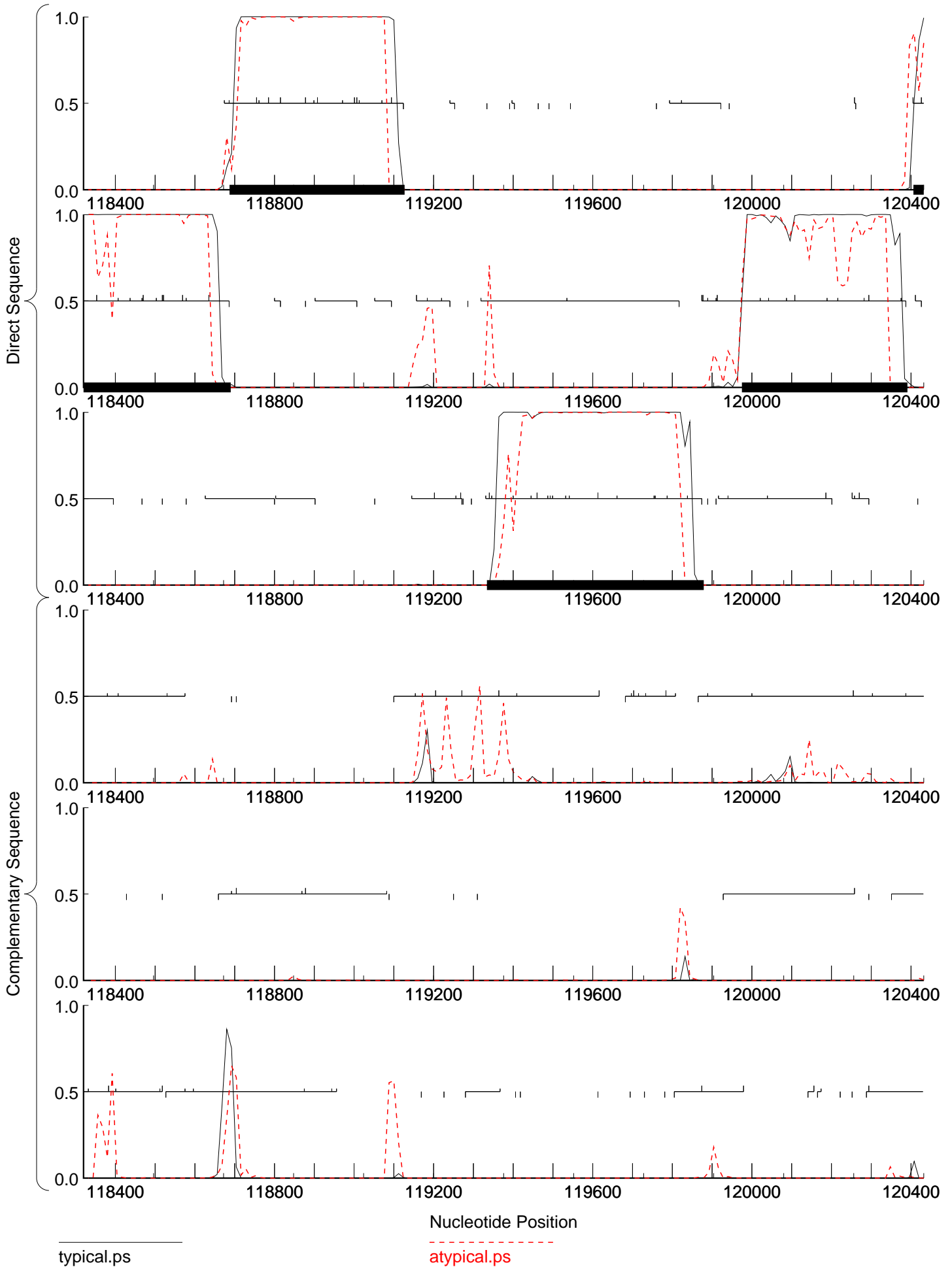


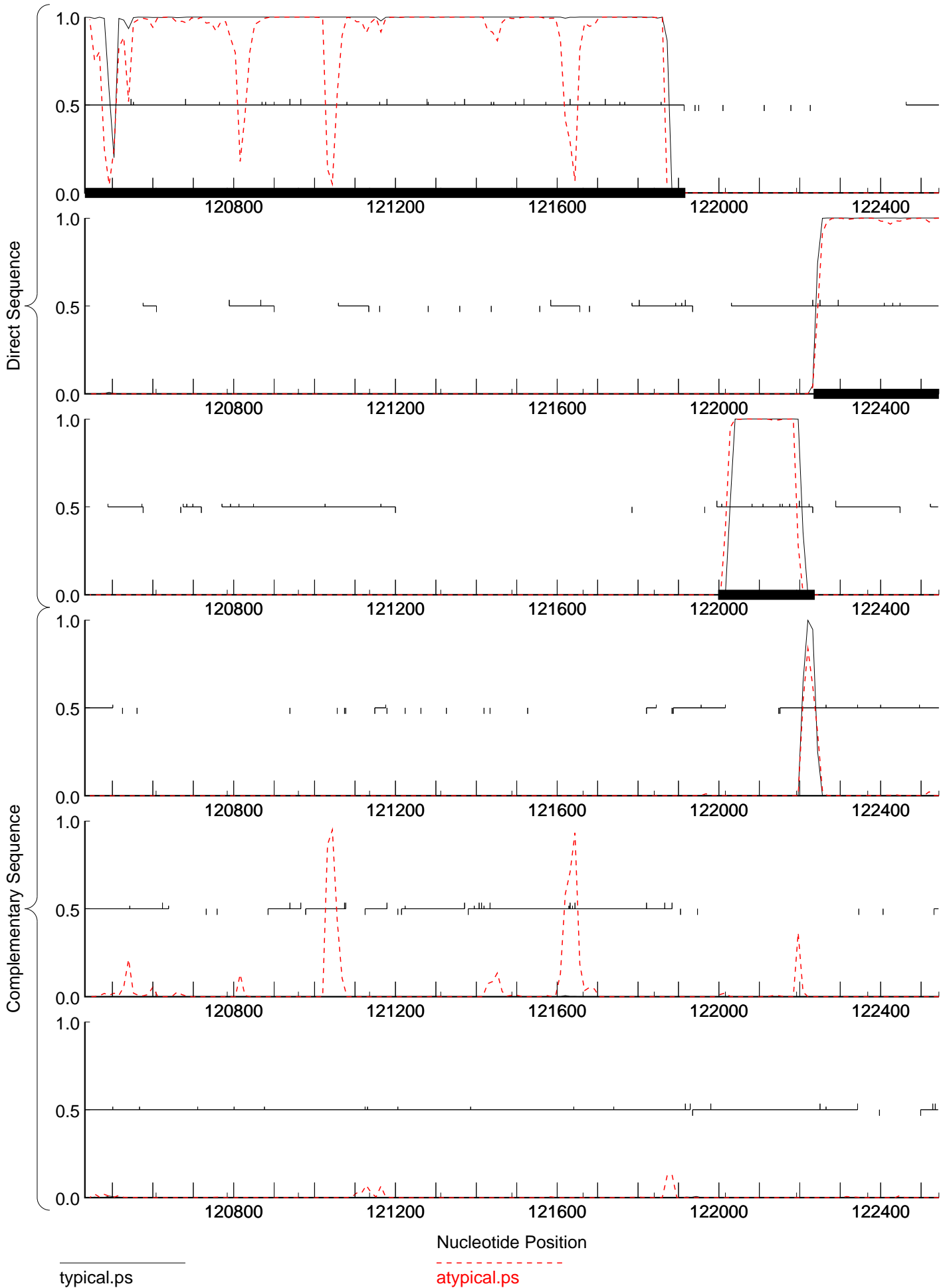


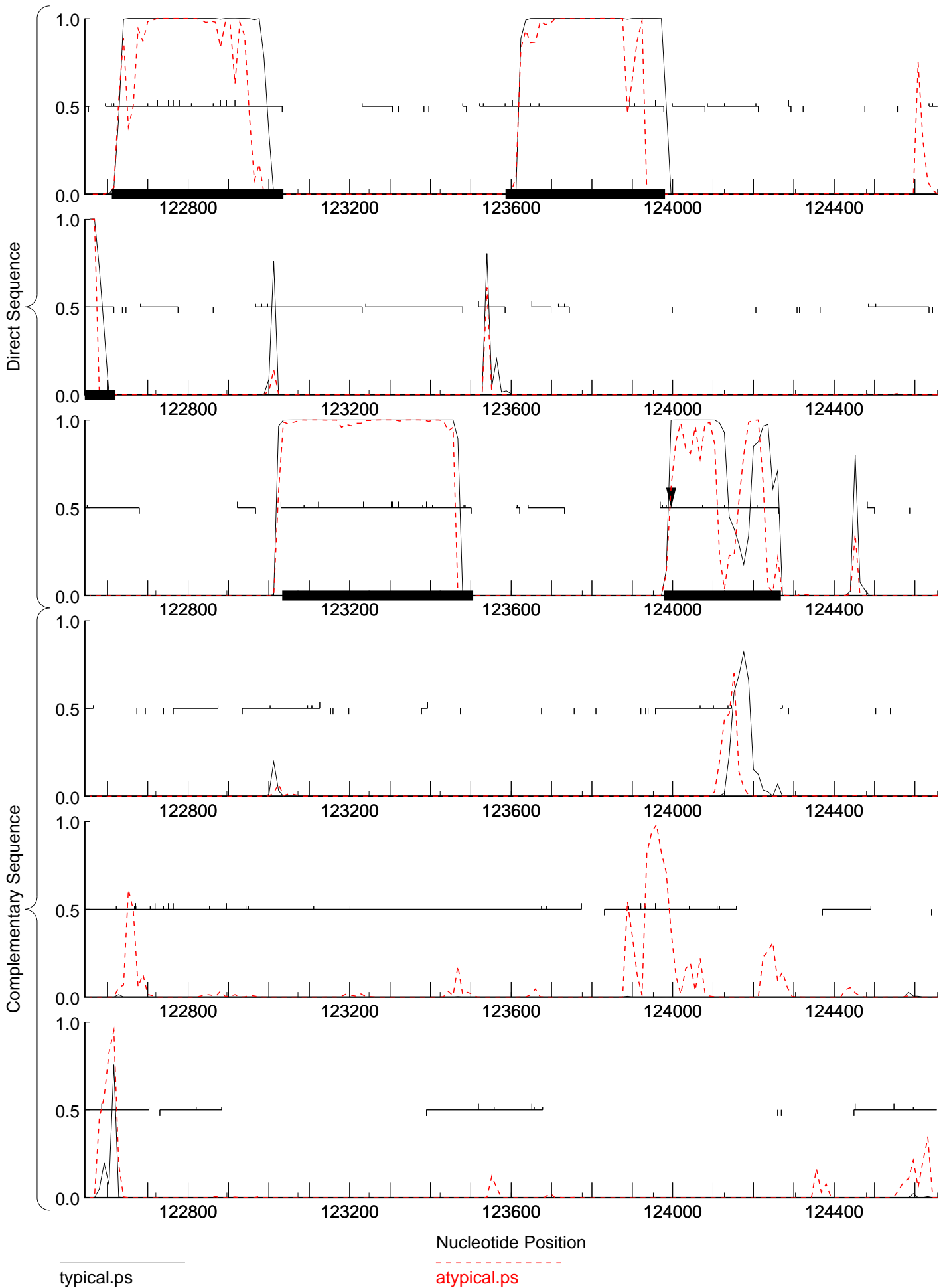


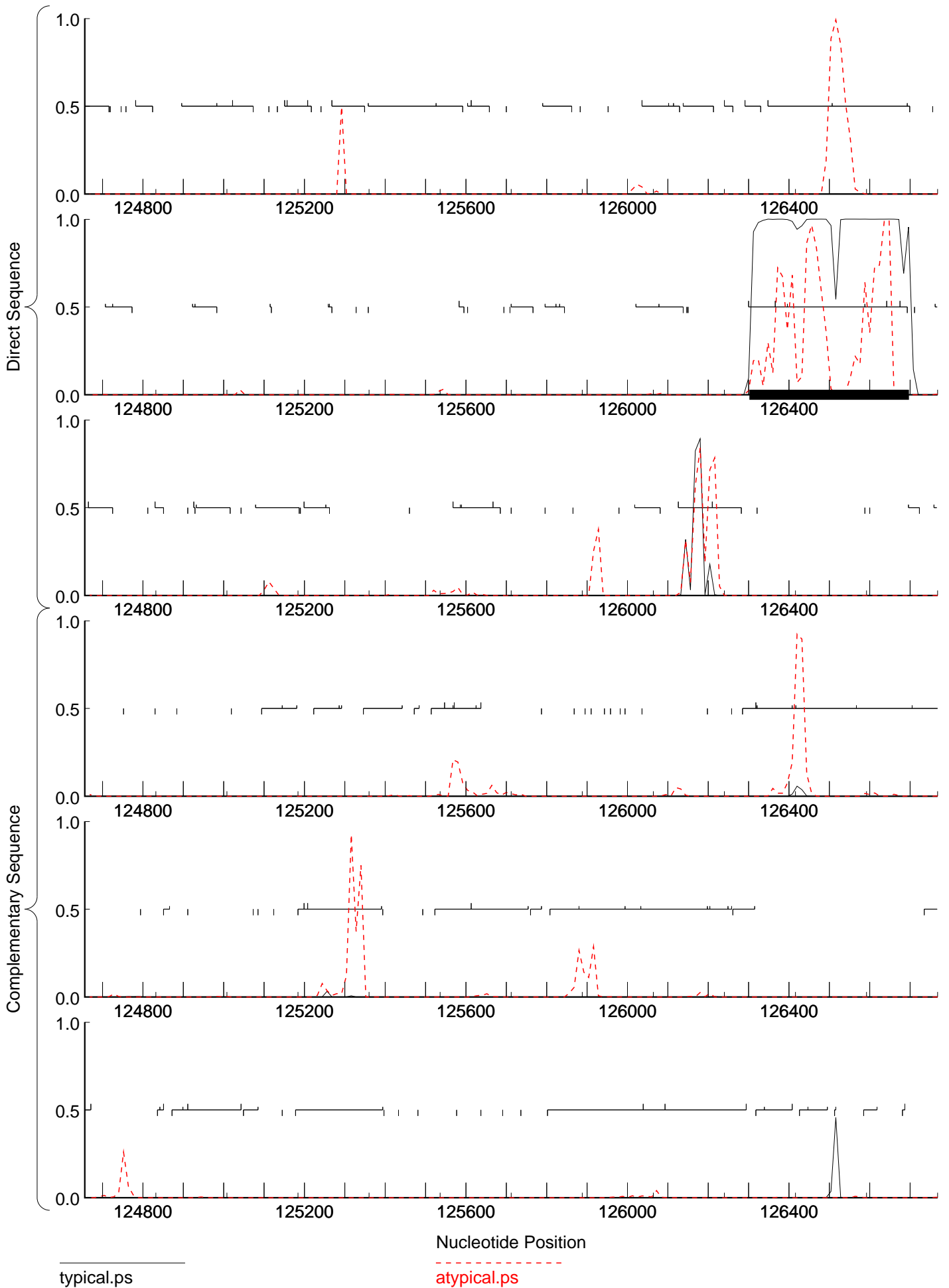


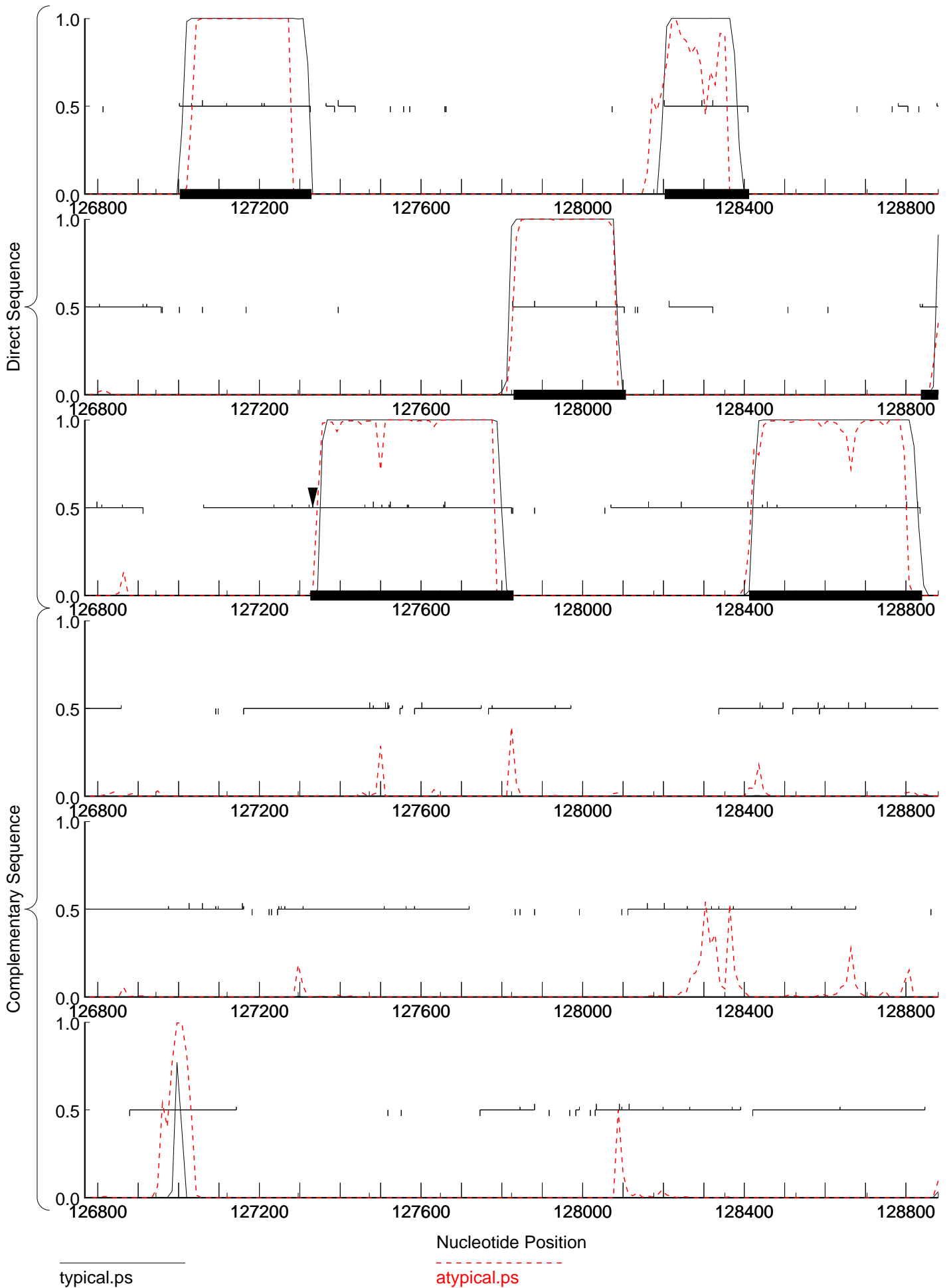


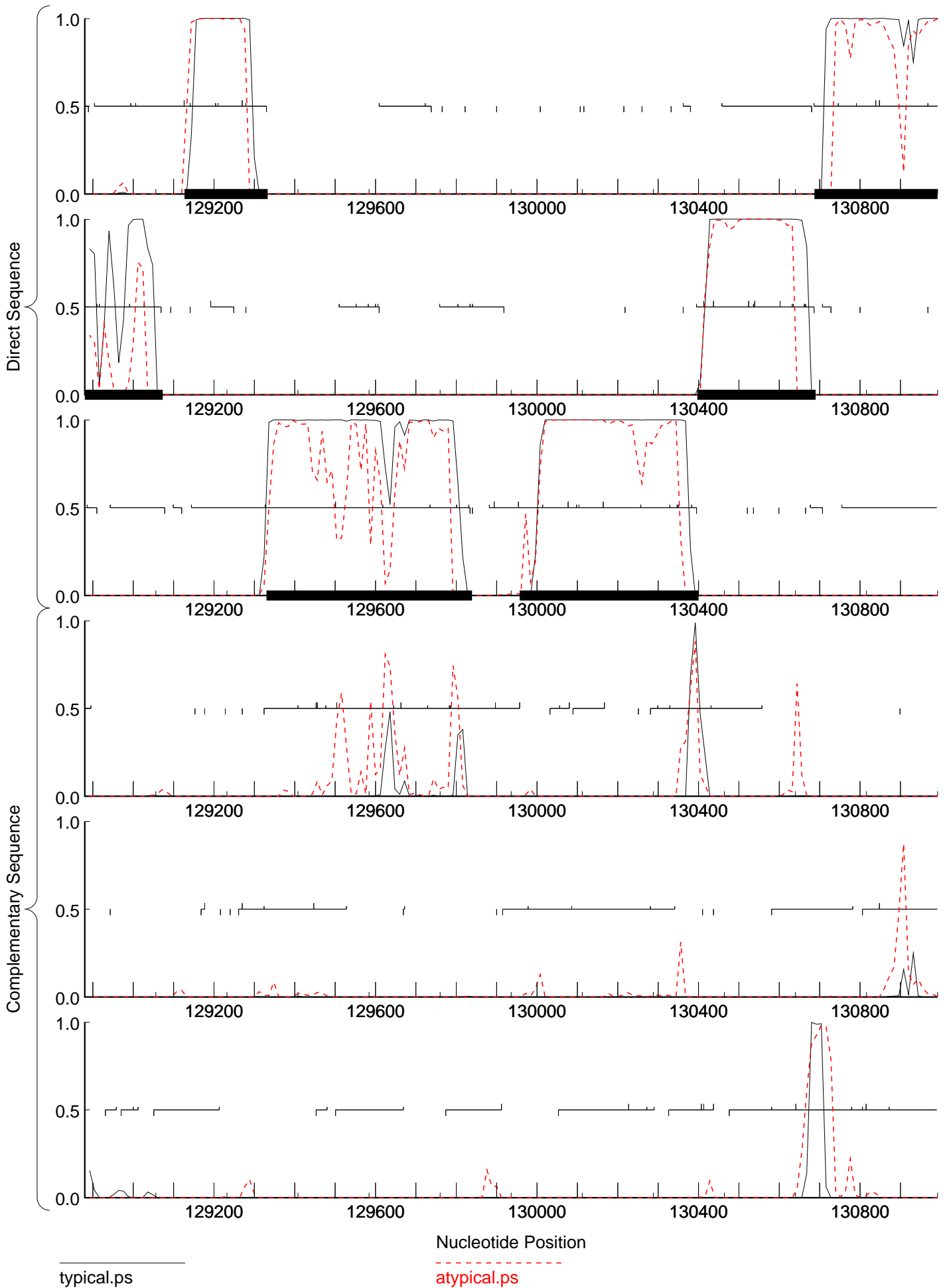


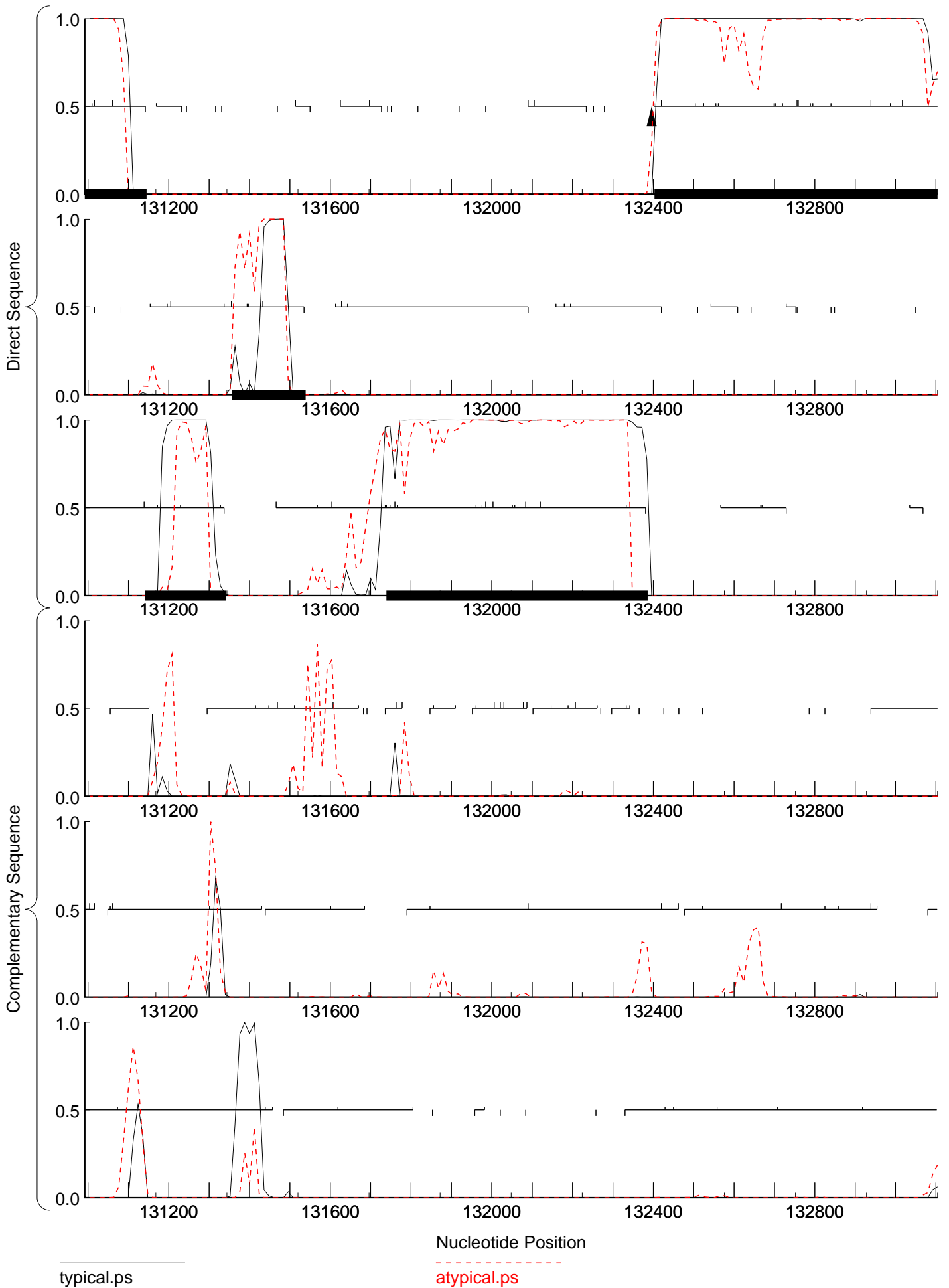




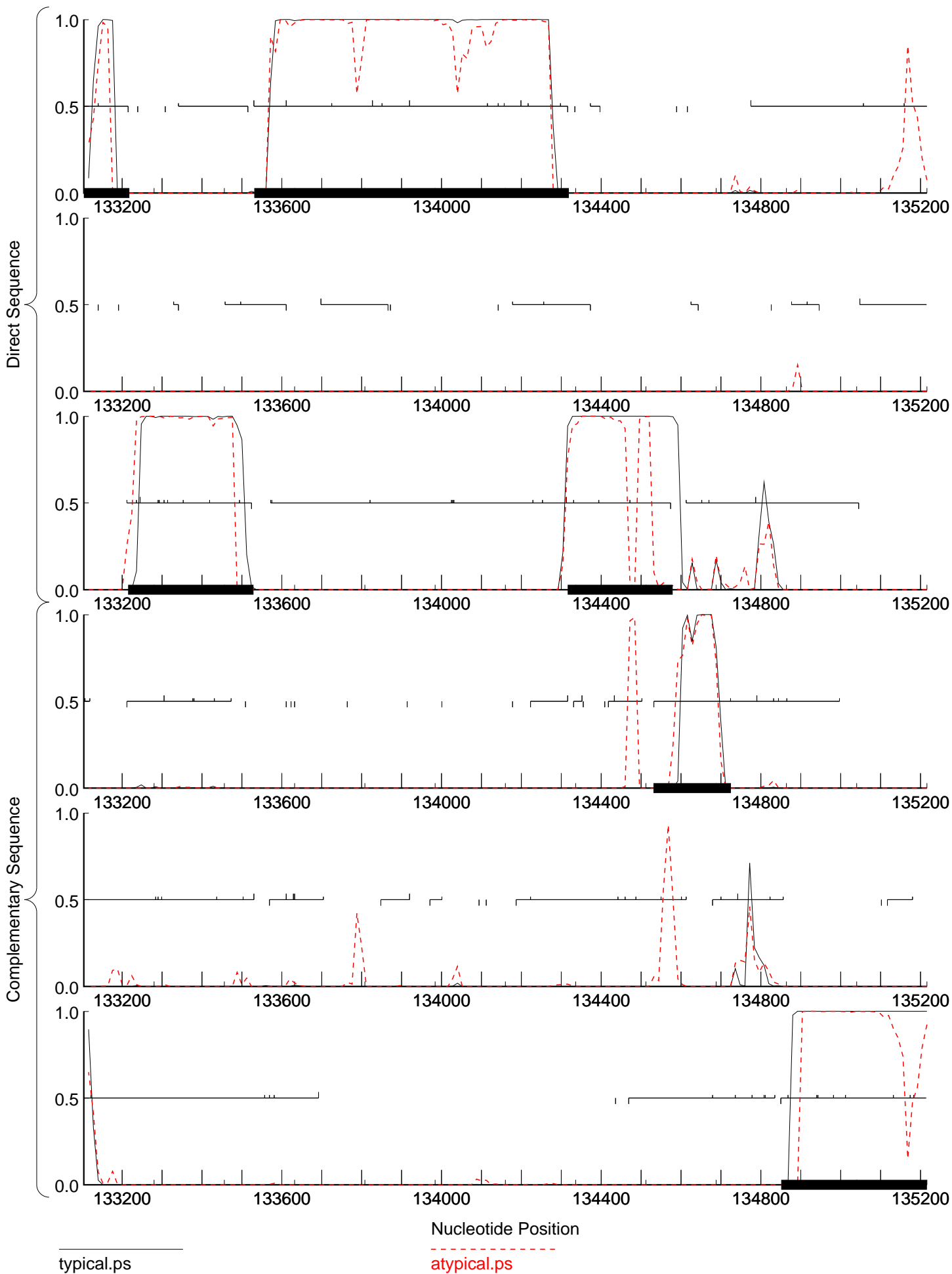


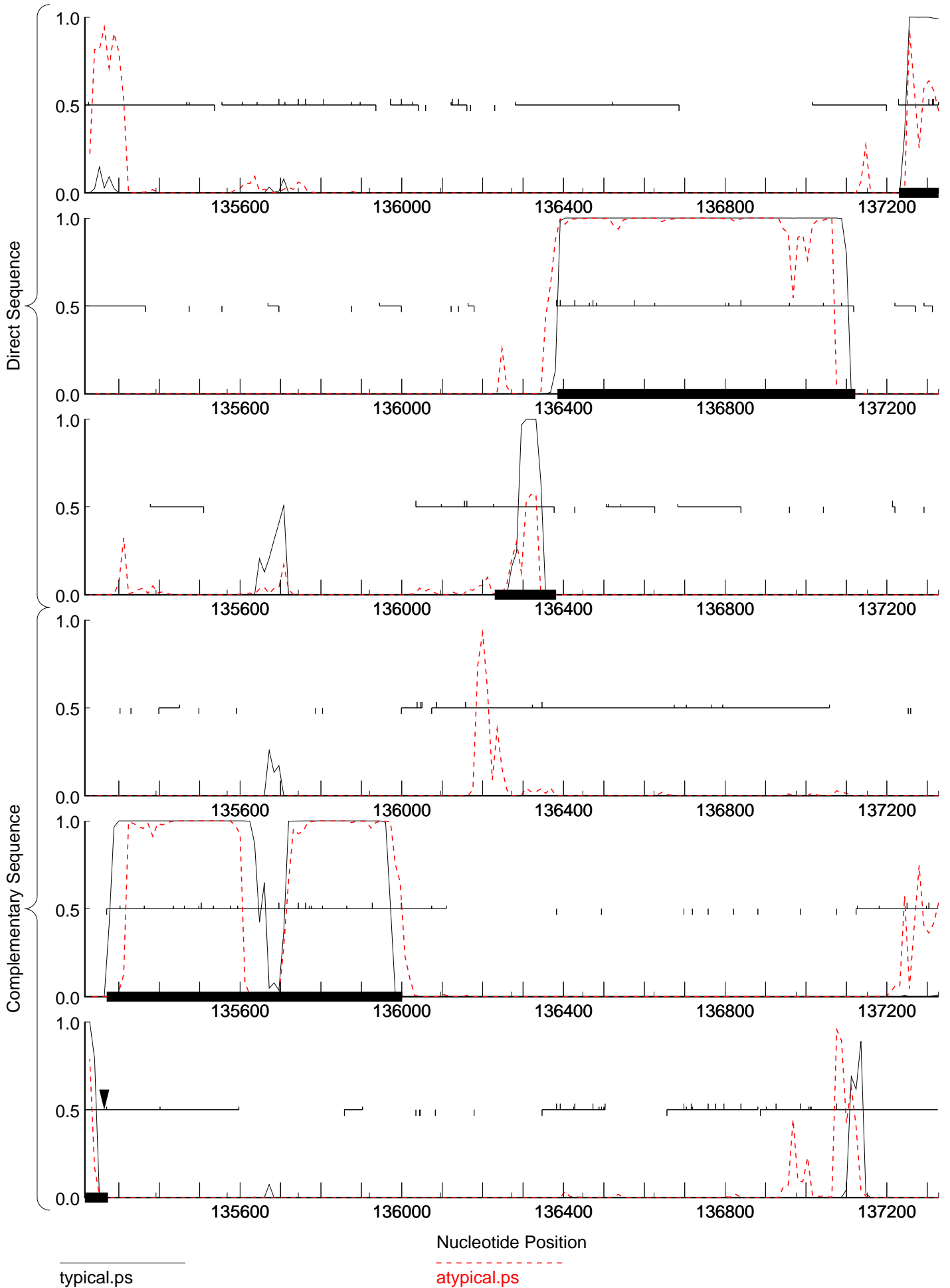






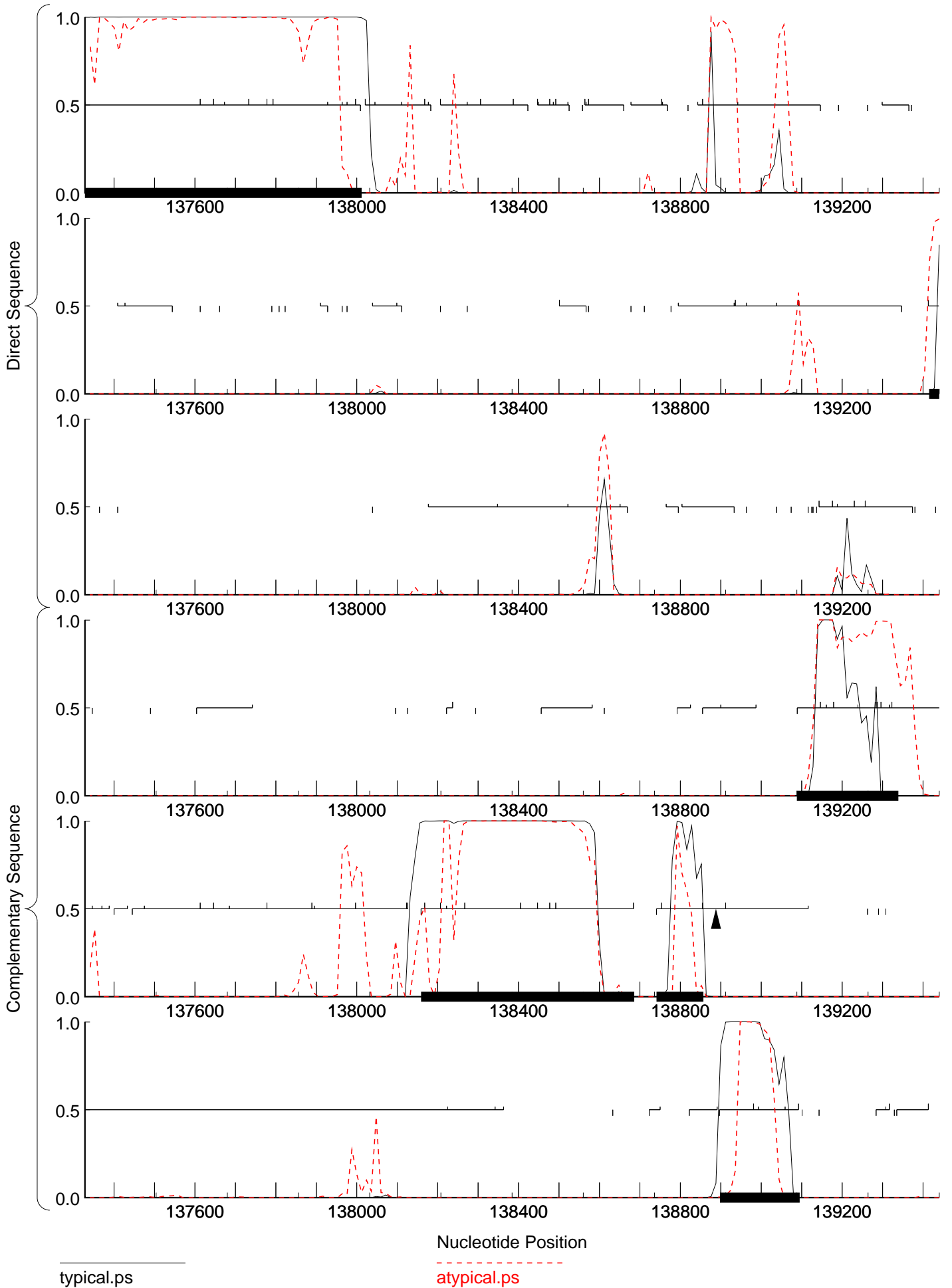


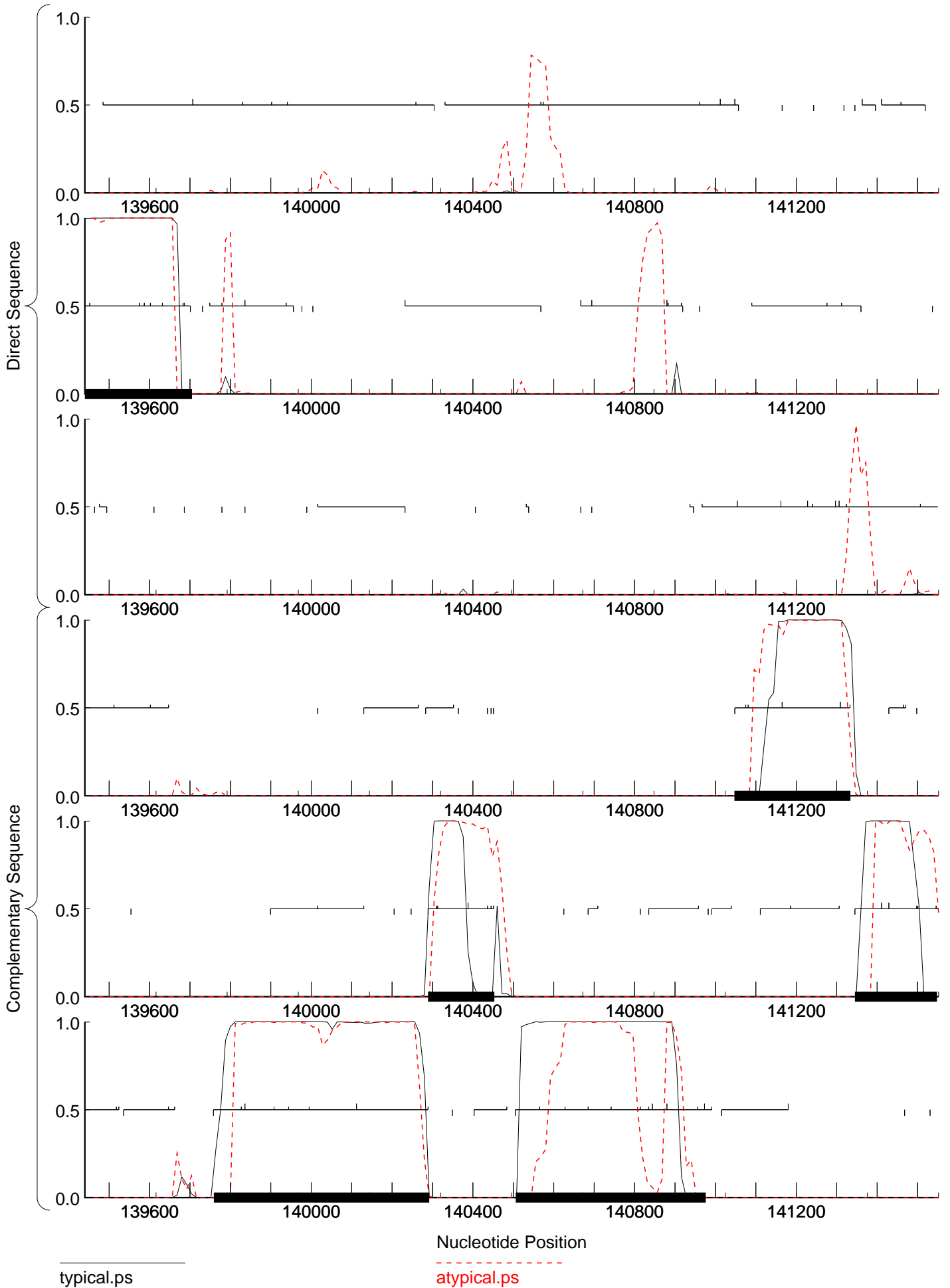


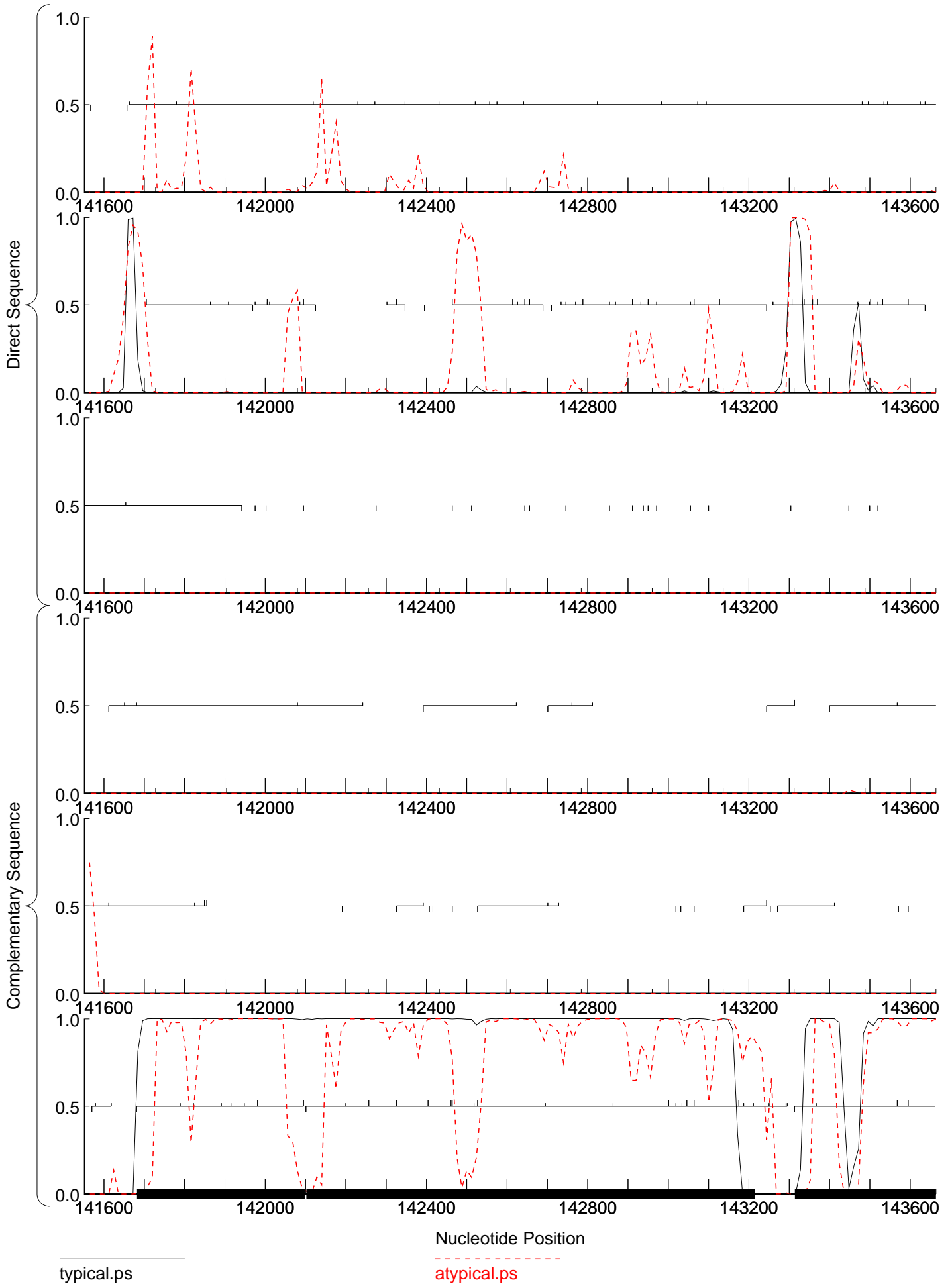


typical.ps

atypical.ps

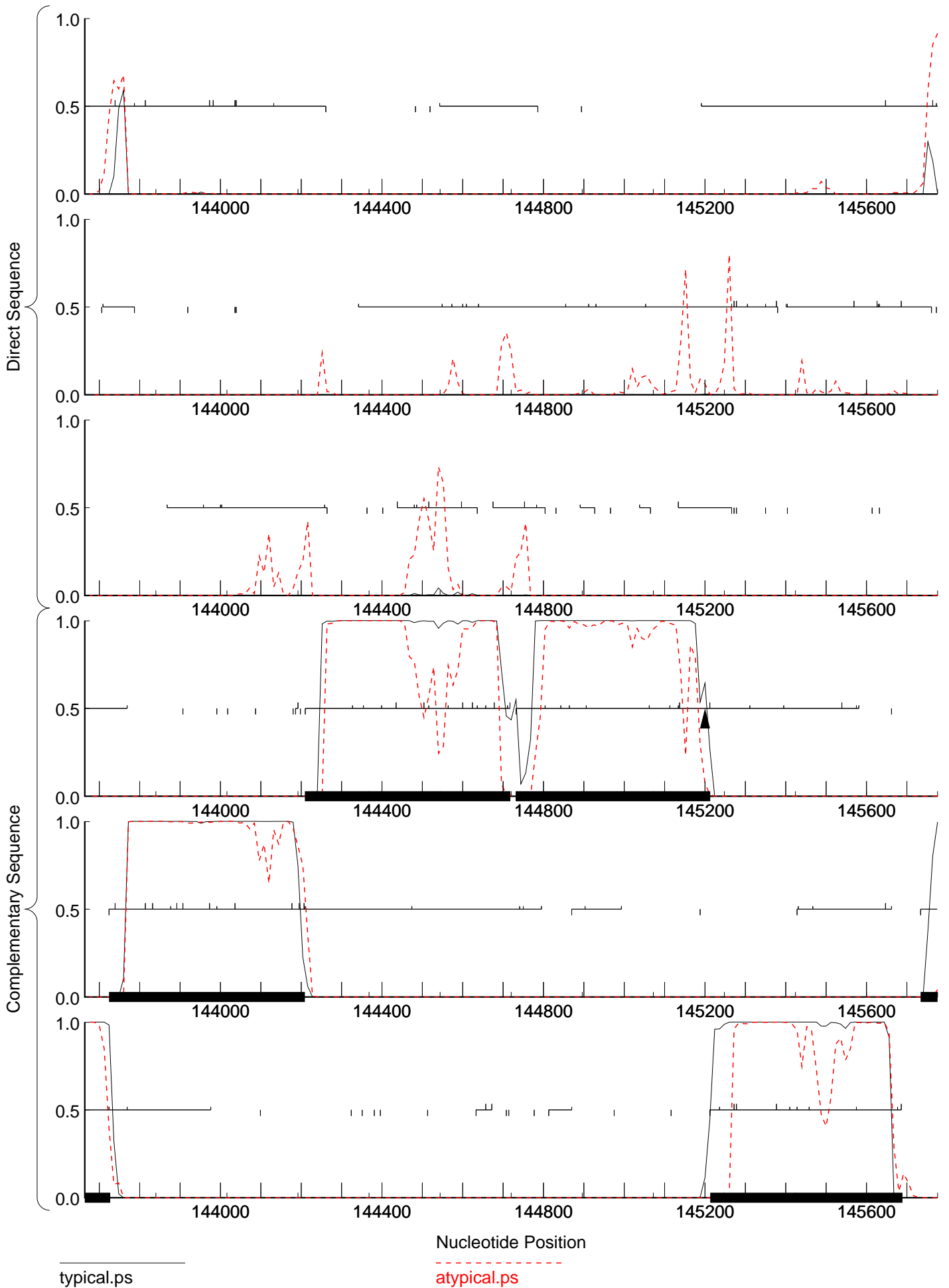


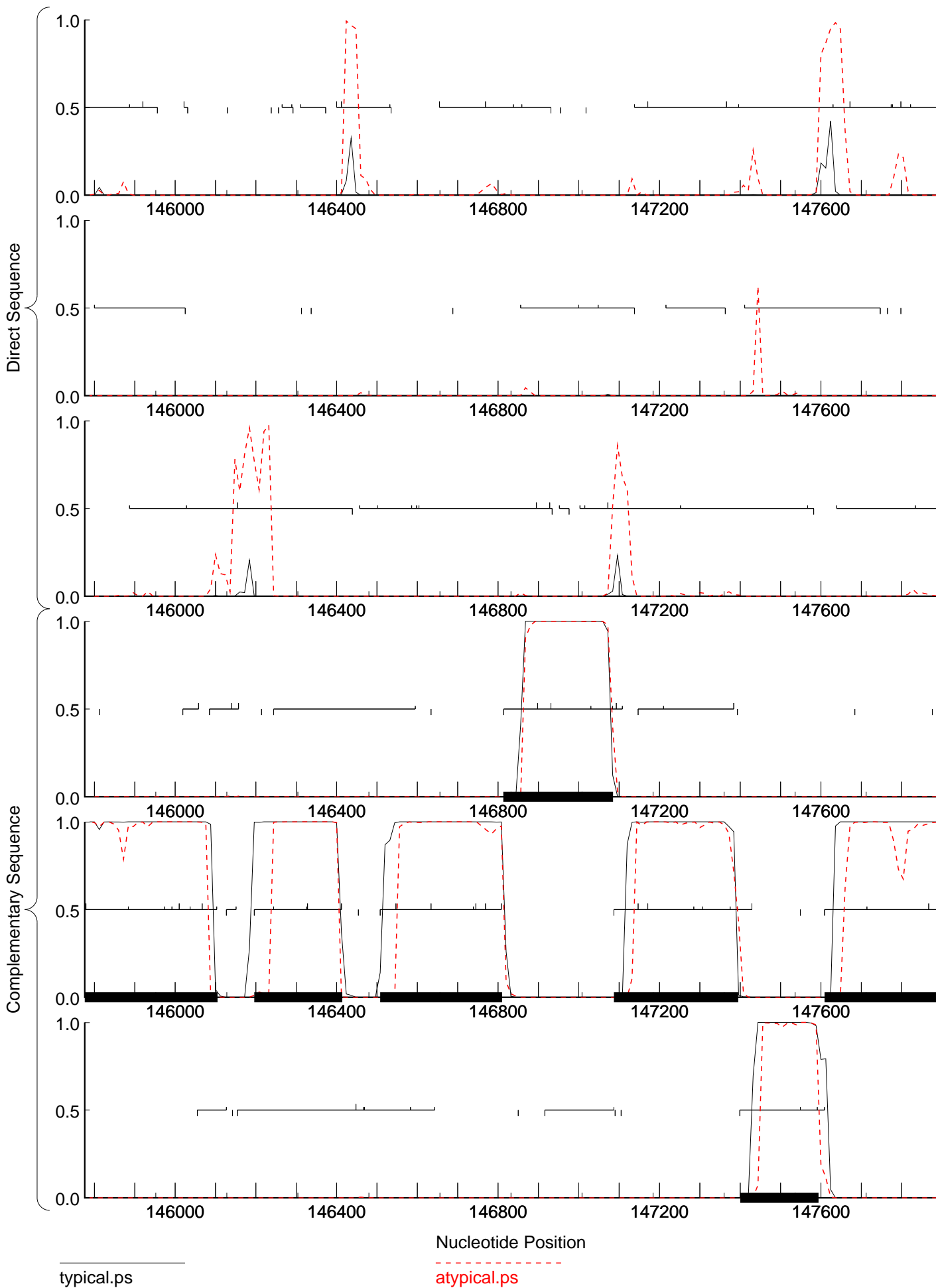




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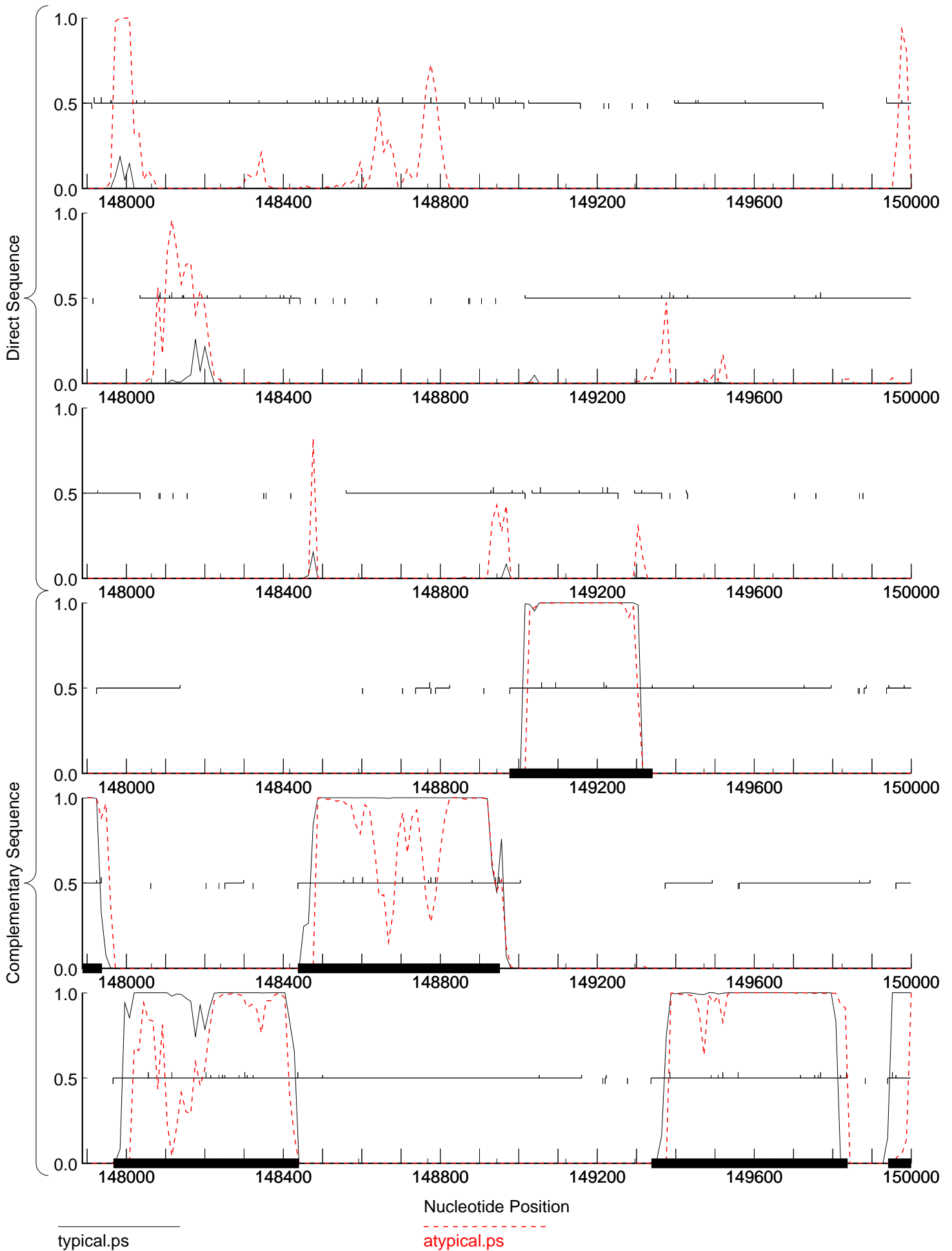
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typical.ps

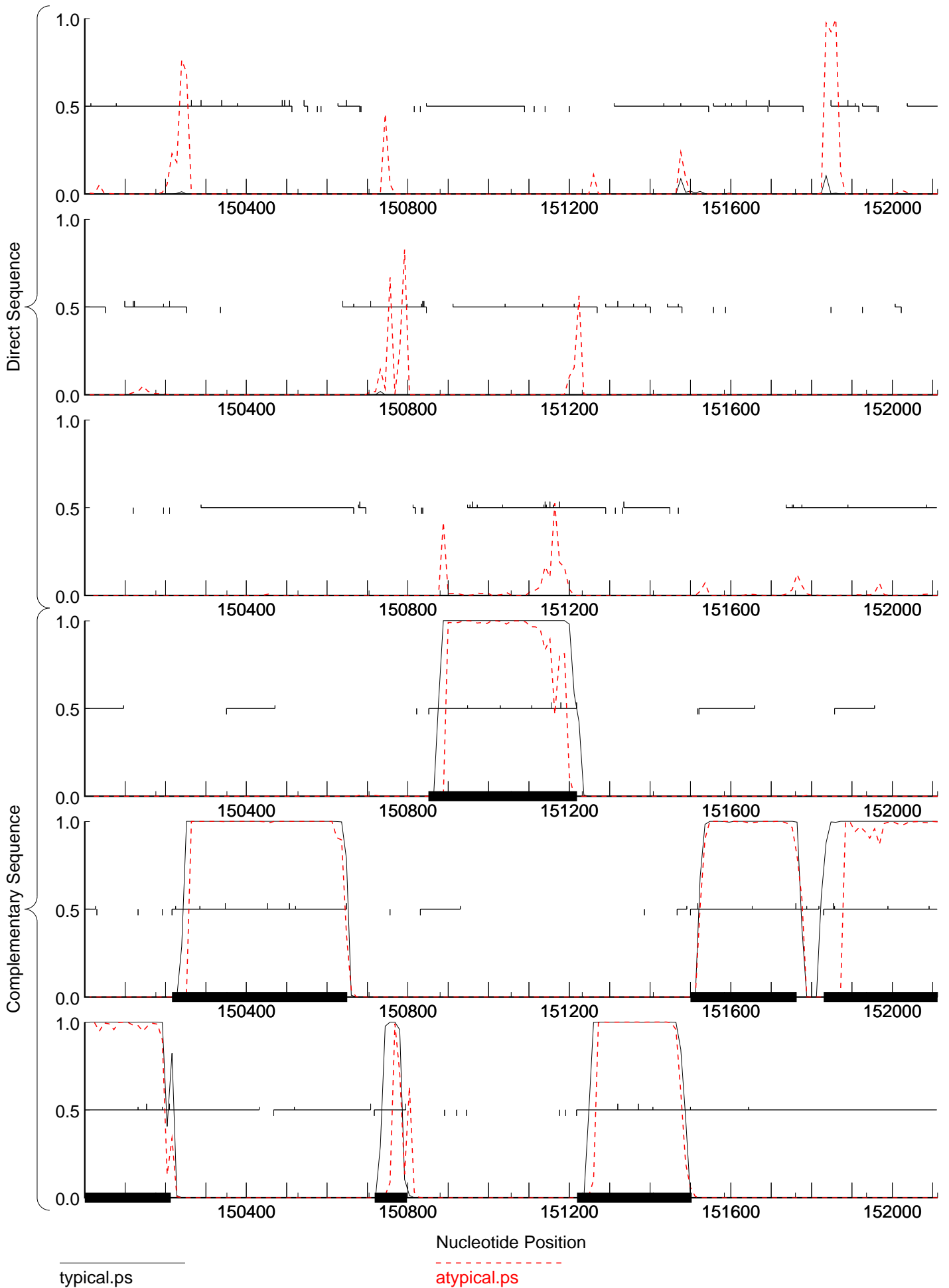
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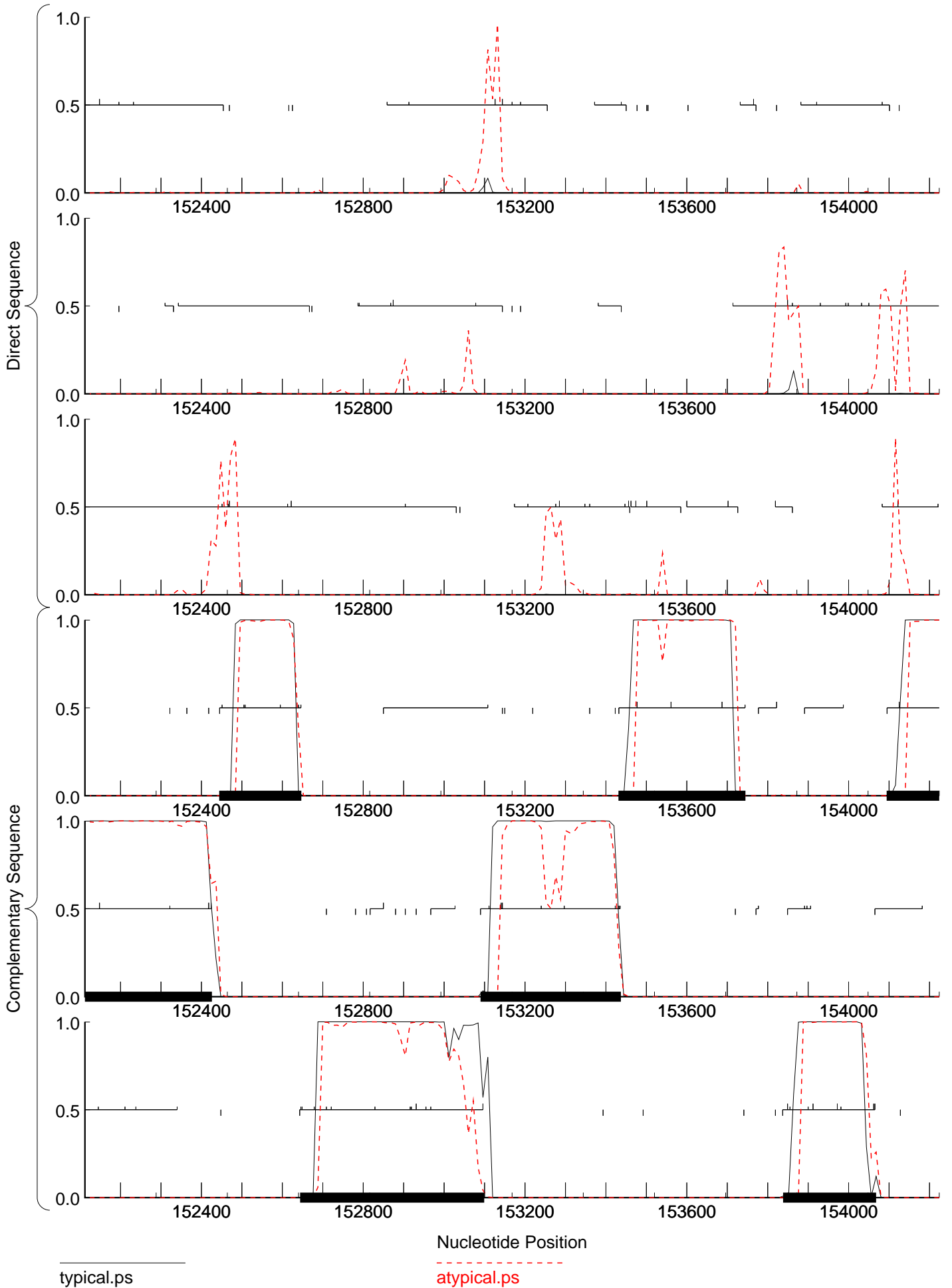


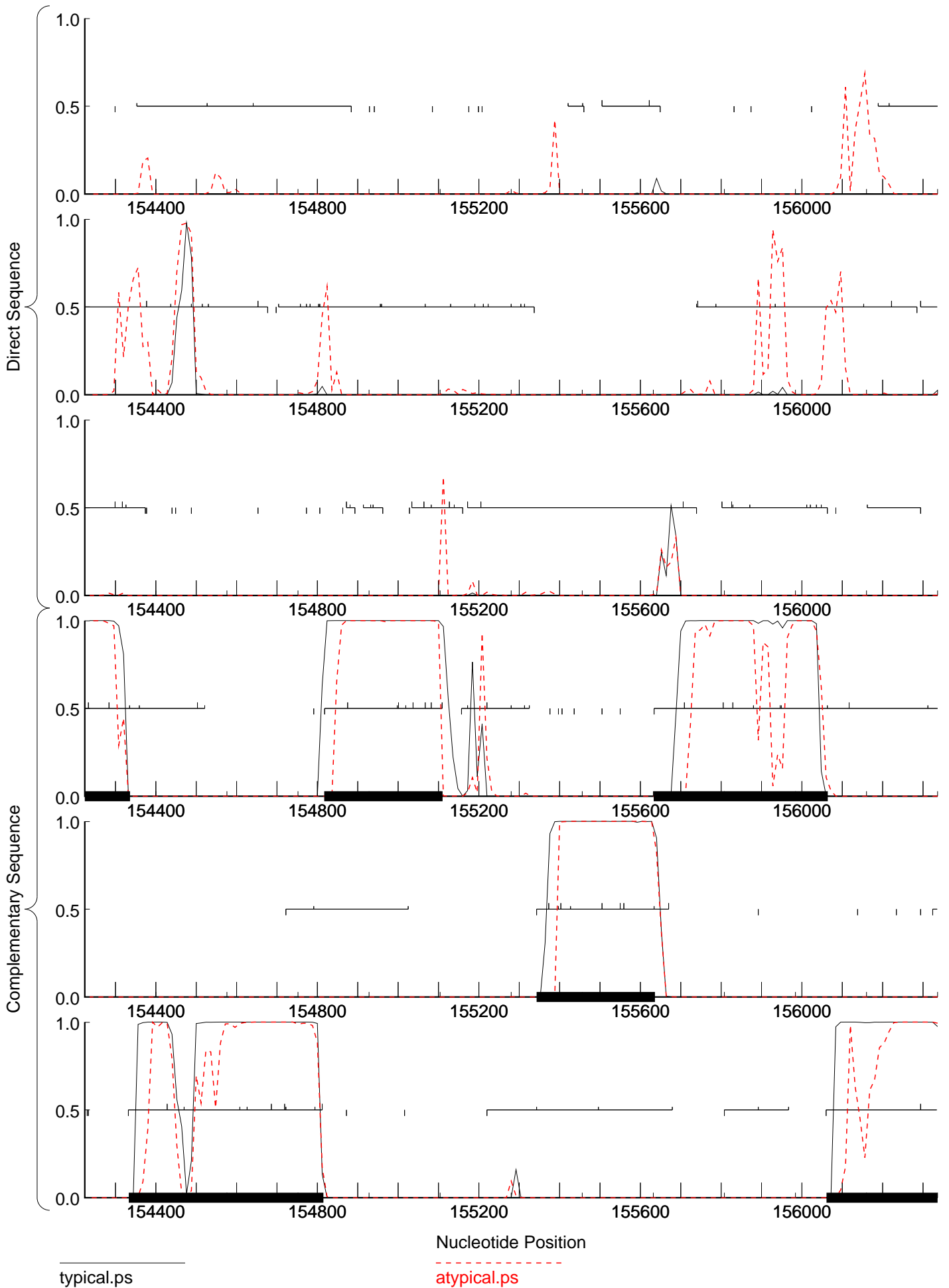
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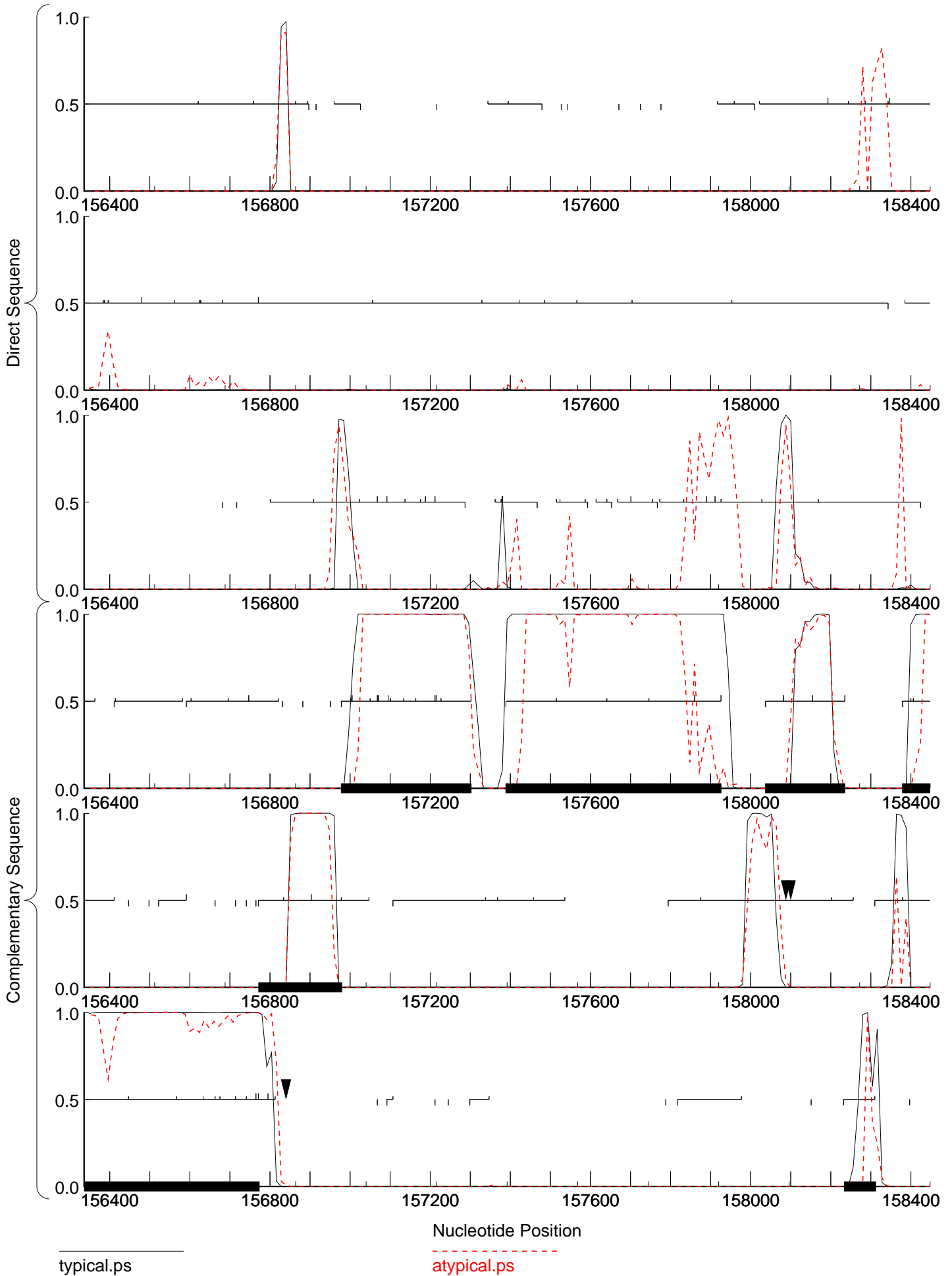
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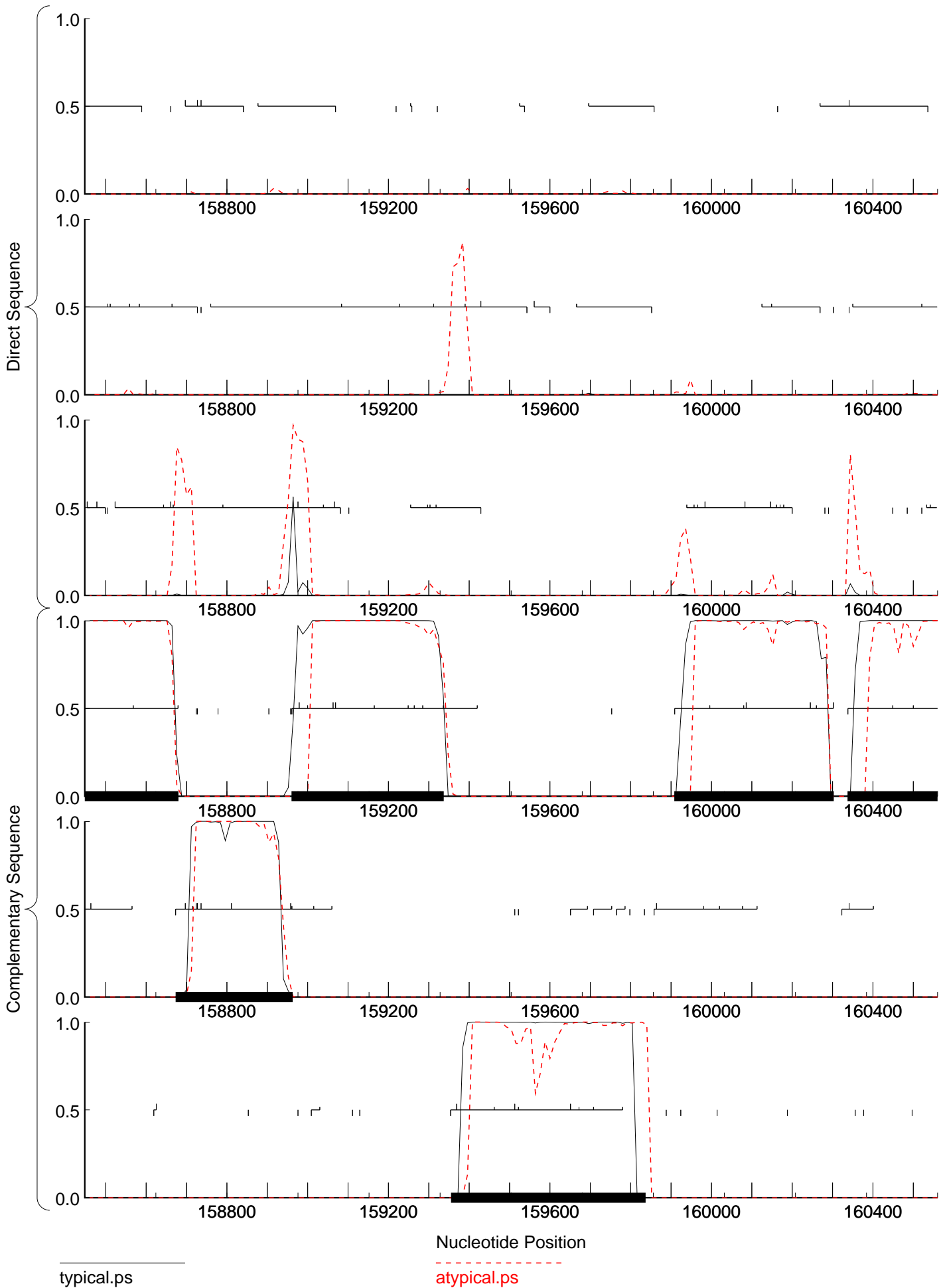


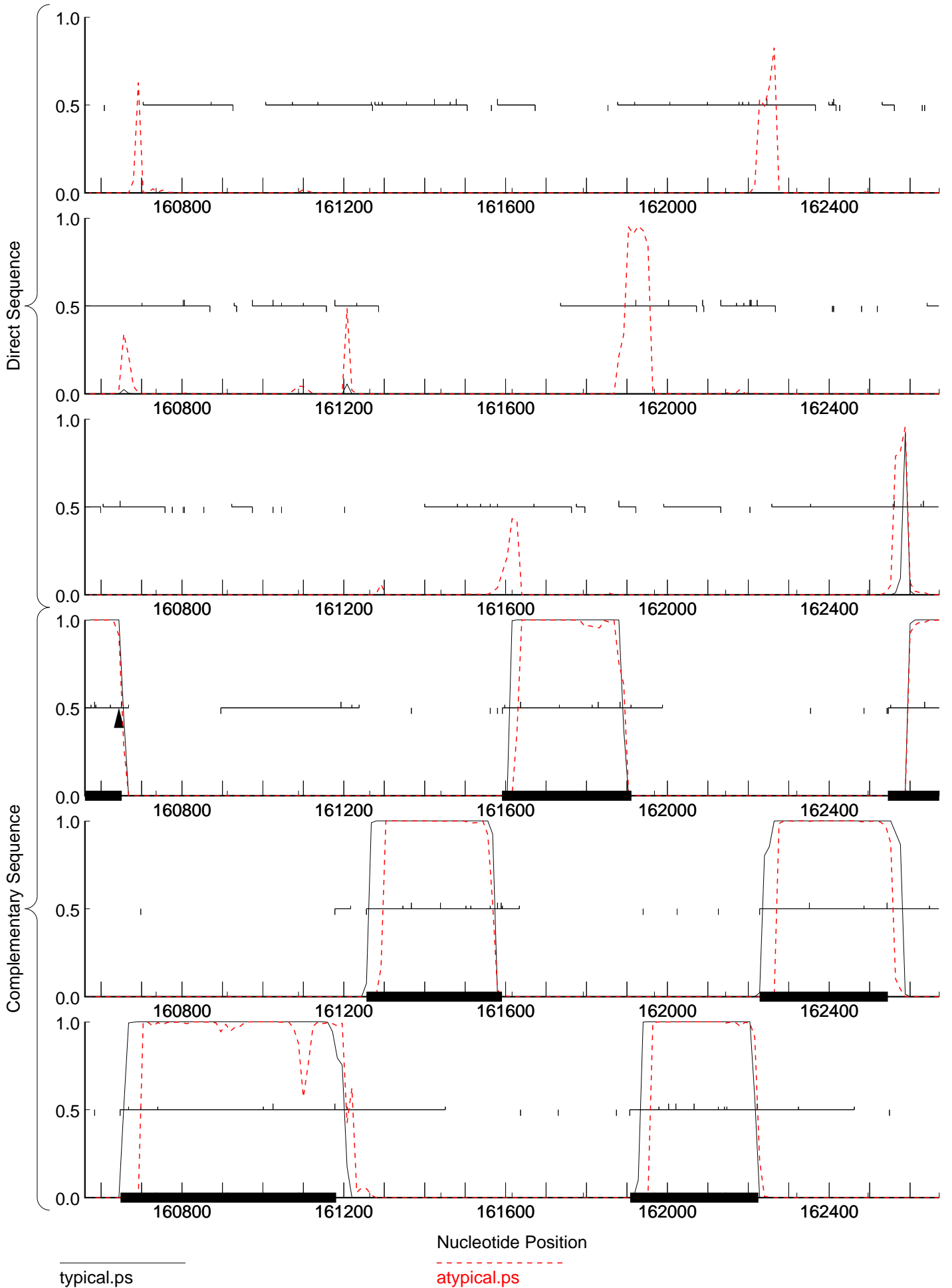


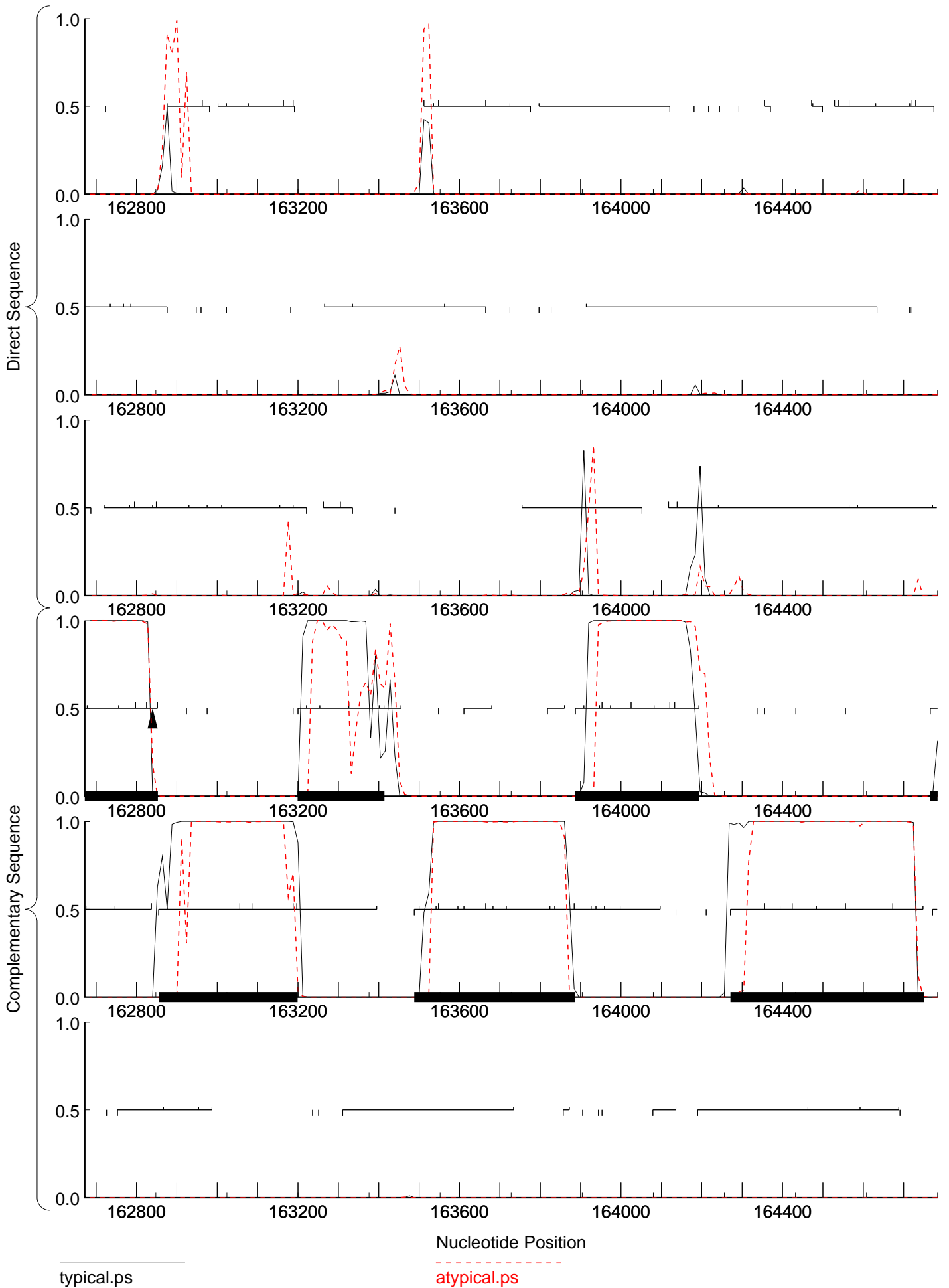


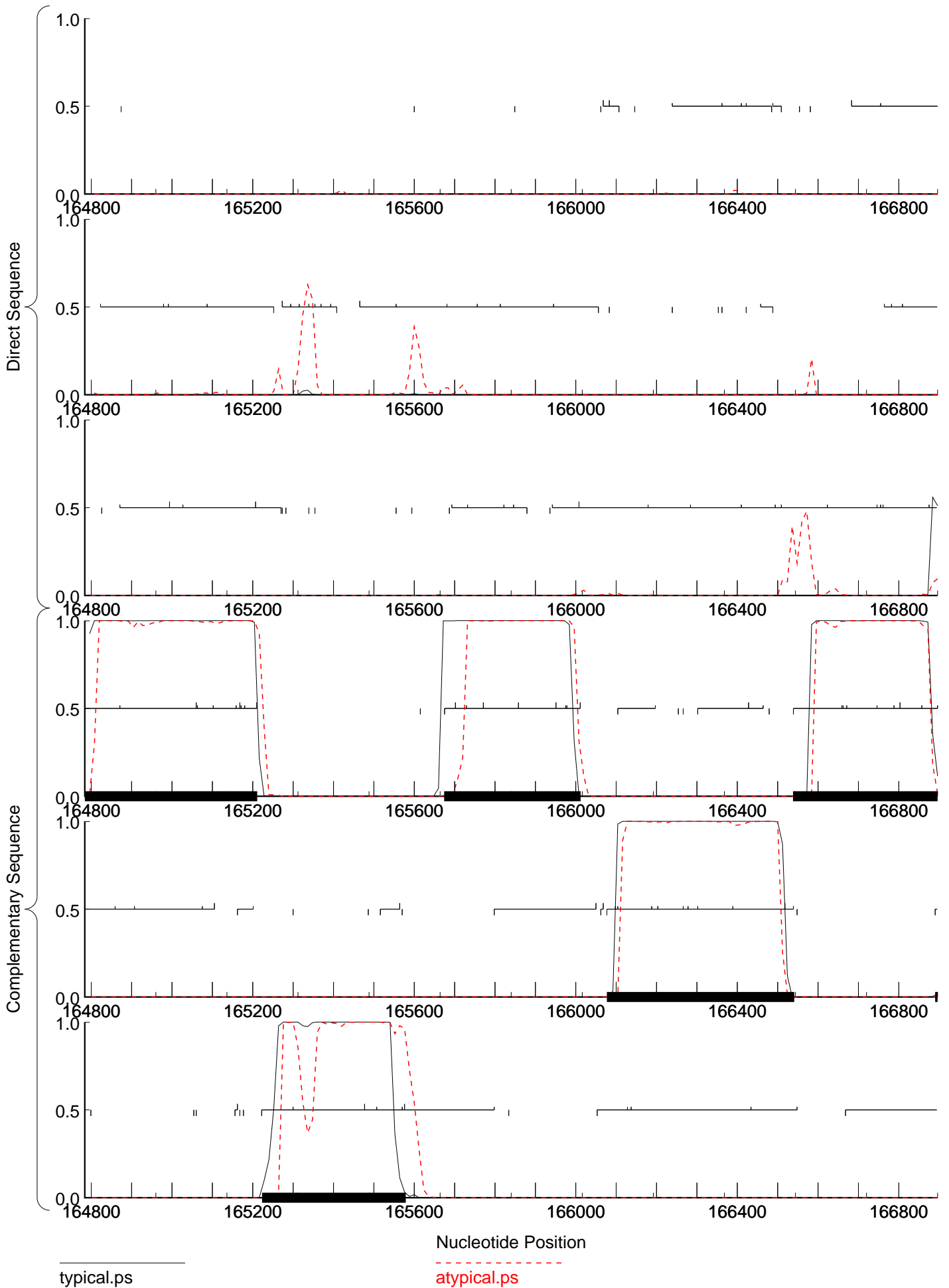
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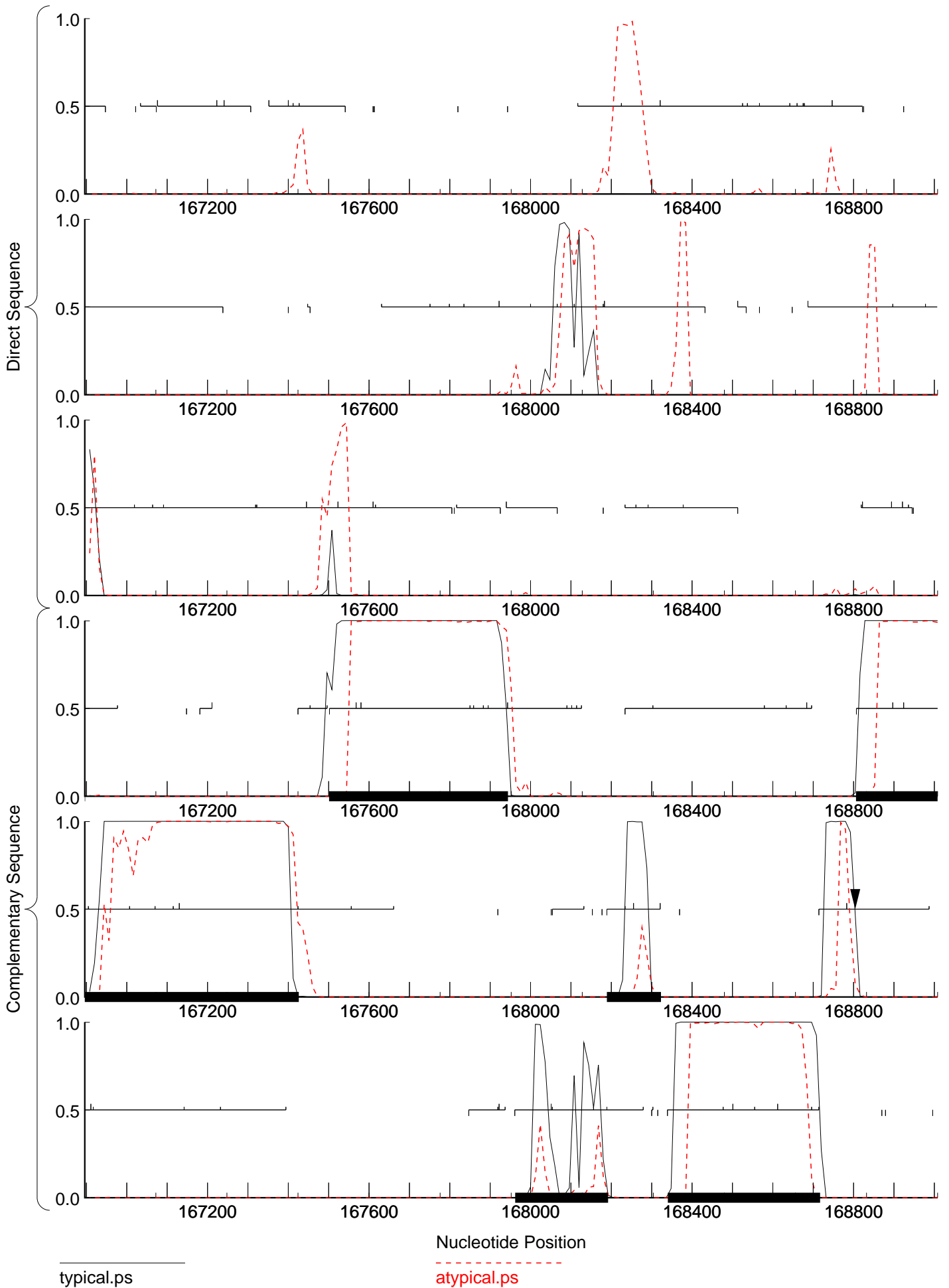












typical.ps

atypical.ps

