

# GeneMark

Version 2.5p (09.08.06)

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## PROGRAM INFORMATION

Sequence : Gordonia phage Lennon complete sequence, 60237 bp, circularly permuted, Cluster DE1  
Analysis Date : 6/22/18 at 13:16:43  
Pages : 30  
Sequence Length : 60237 bp  
GC Content : 67.36%

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,

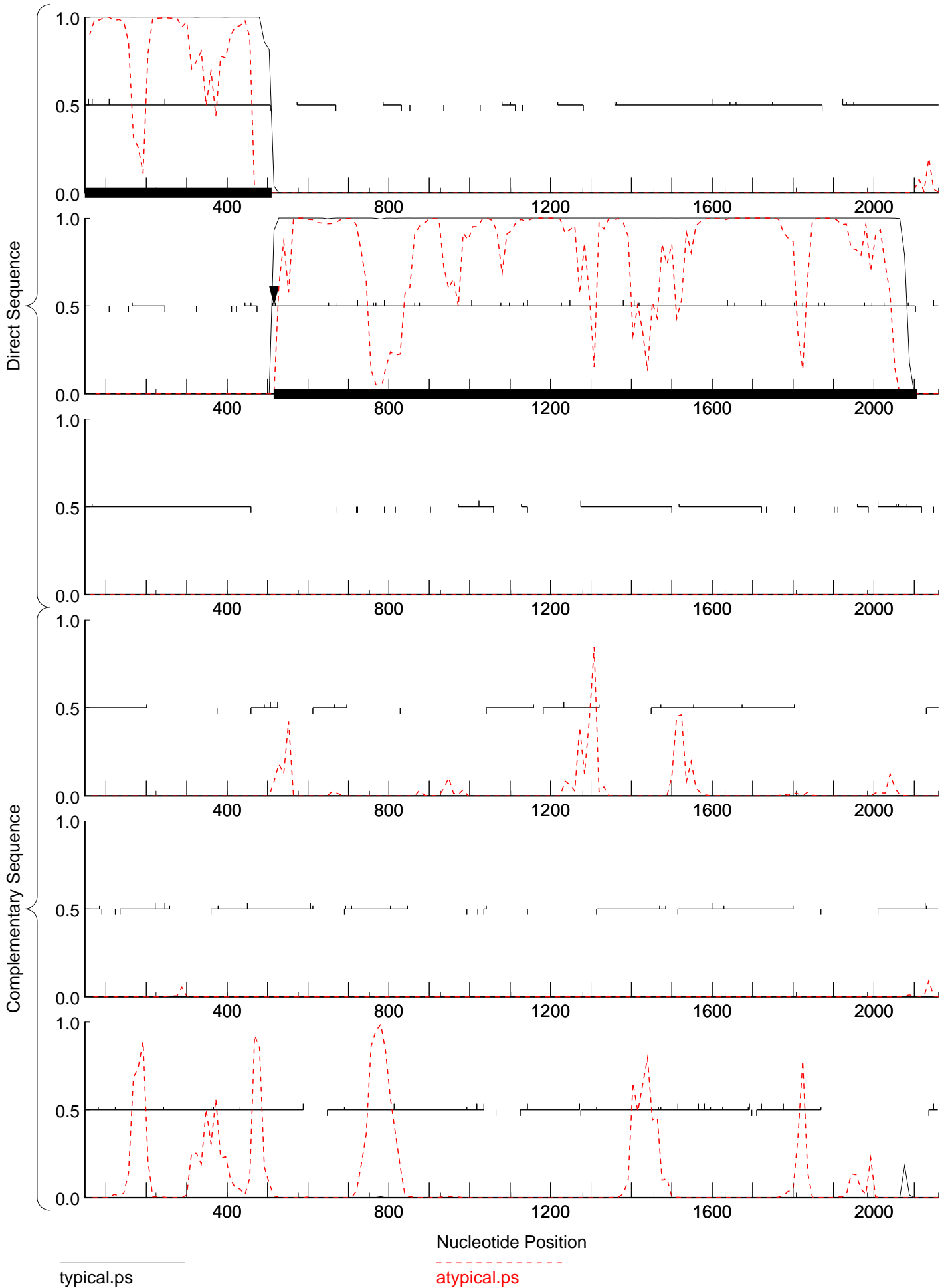
## Matrix notes & comments

Training set derived by GeneMarkS, 4.30 October 2014  
Fri Jun 22 13:16:43 2018

## MATRIX INFORMATION

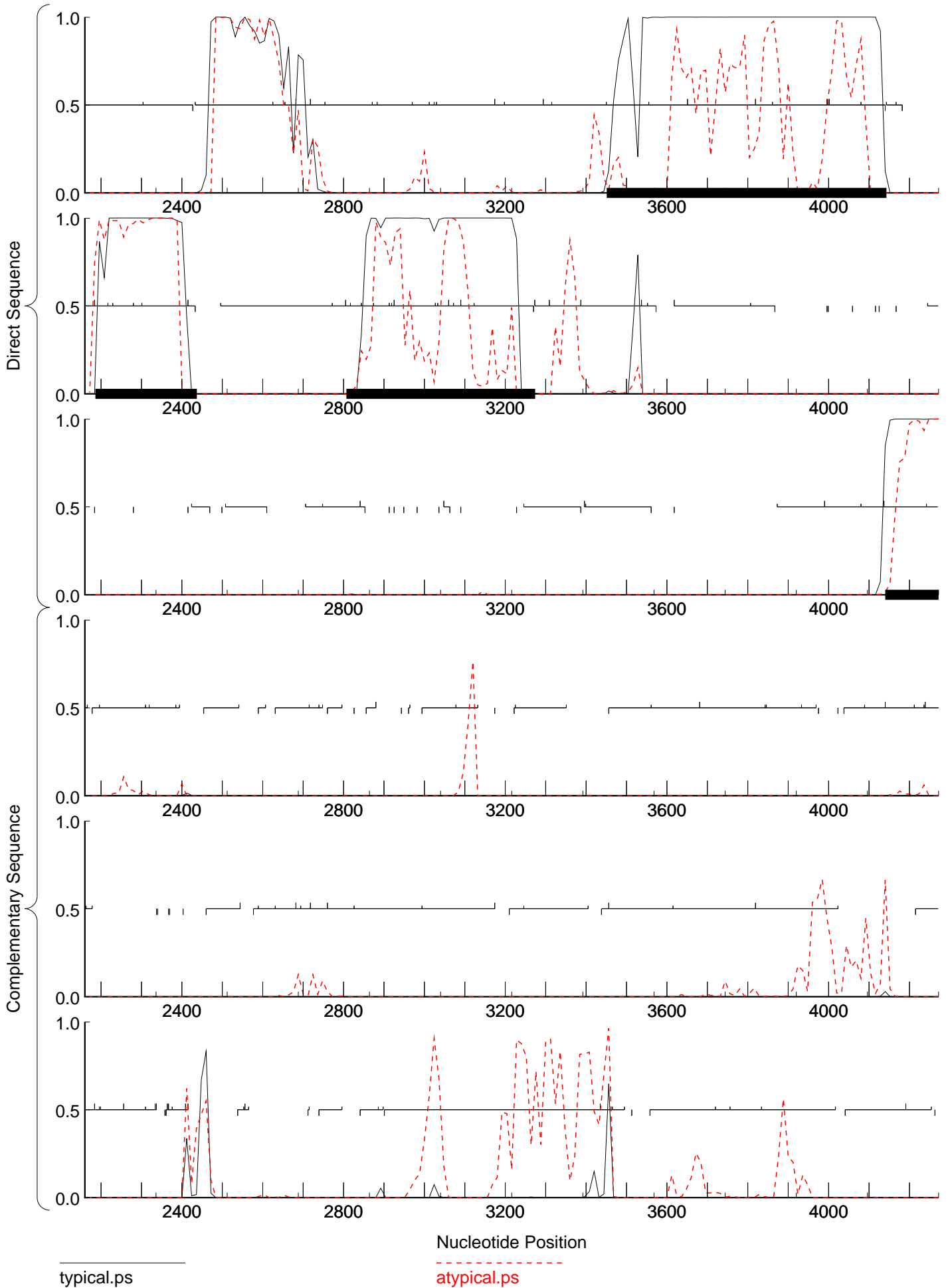
Matrix : -  
Author : -  
Order : 4

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



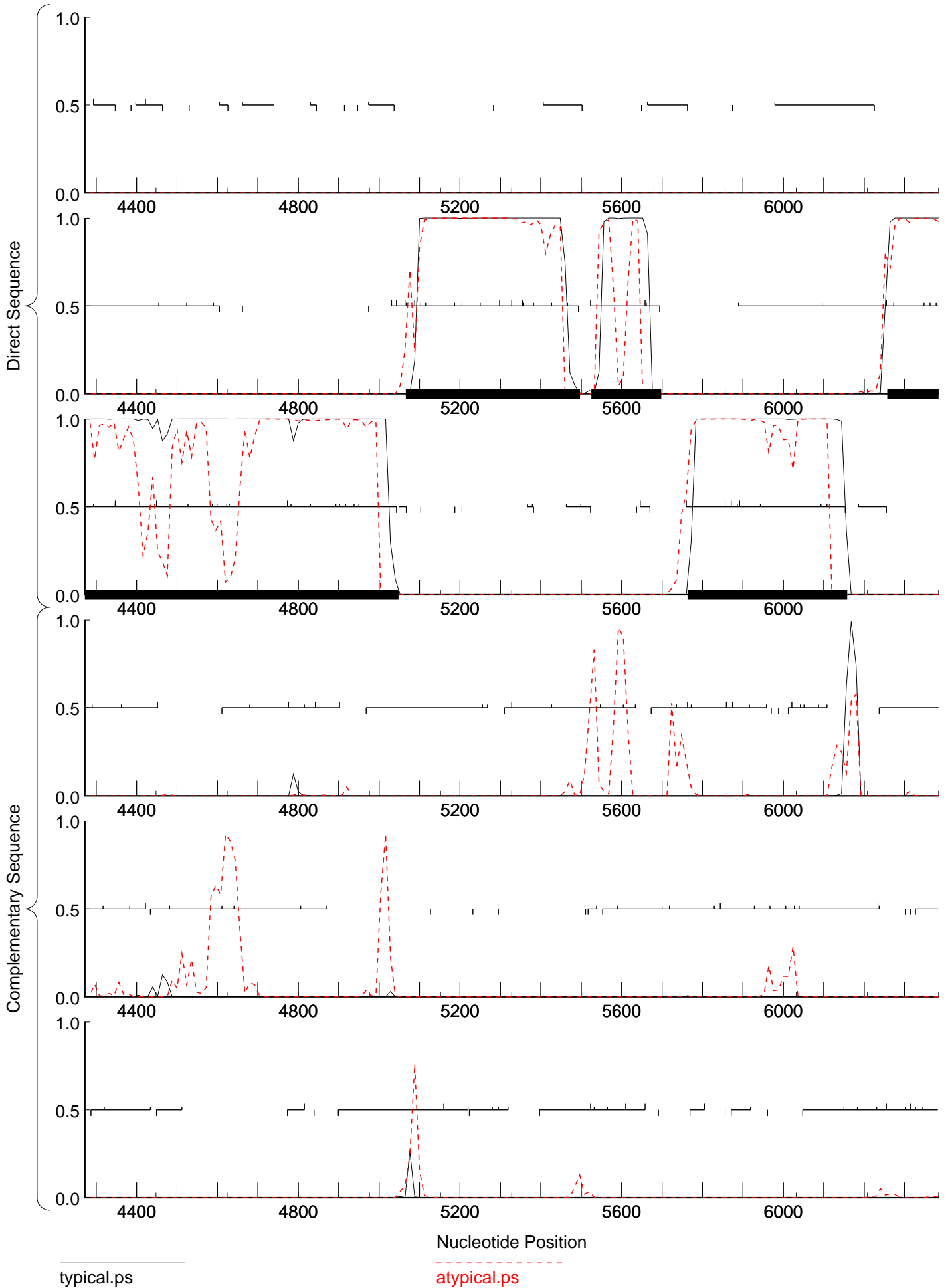
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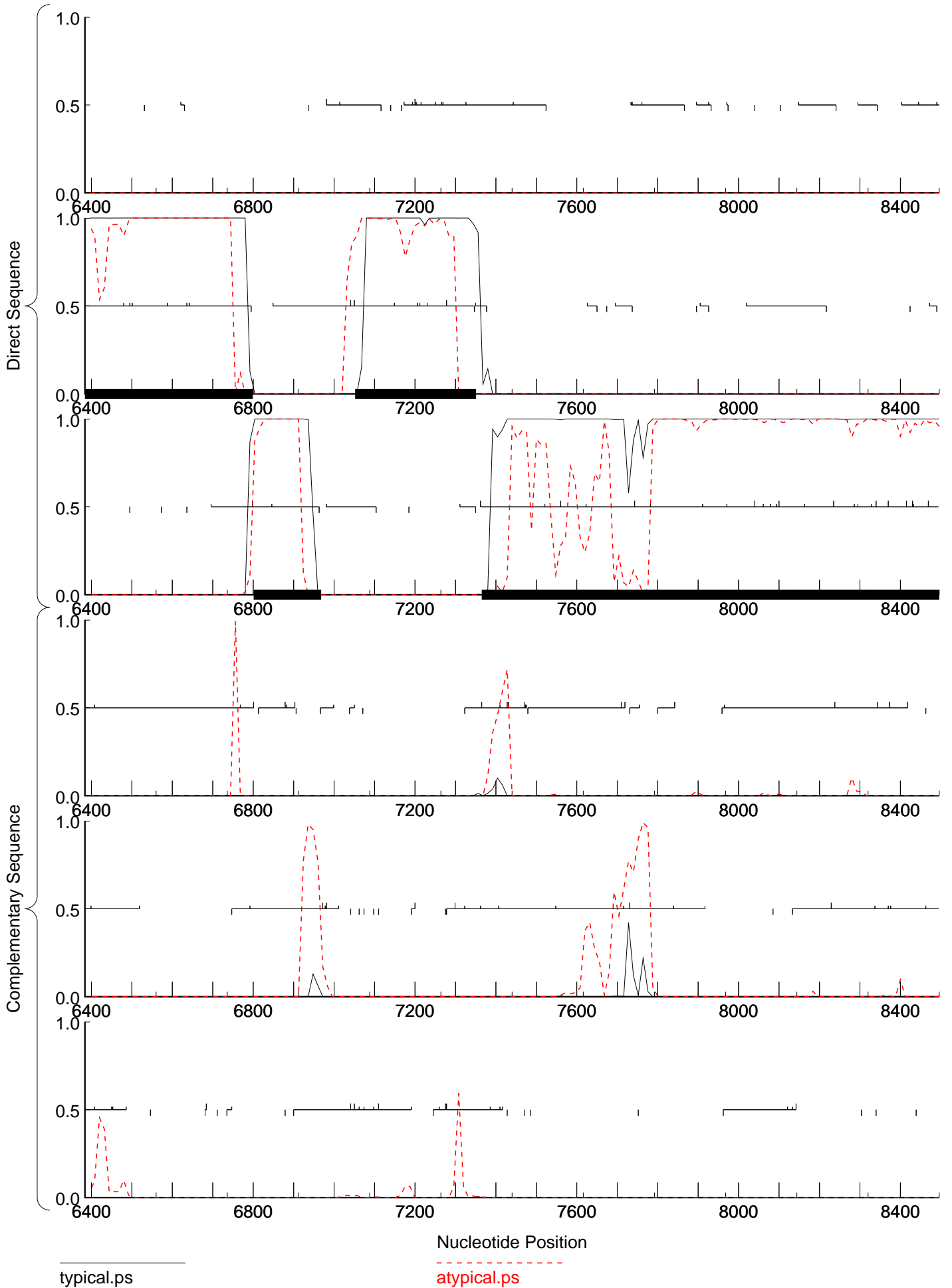
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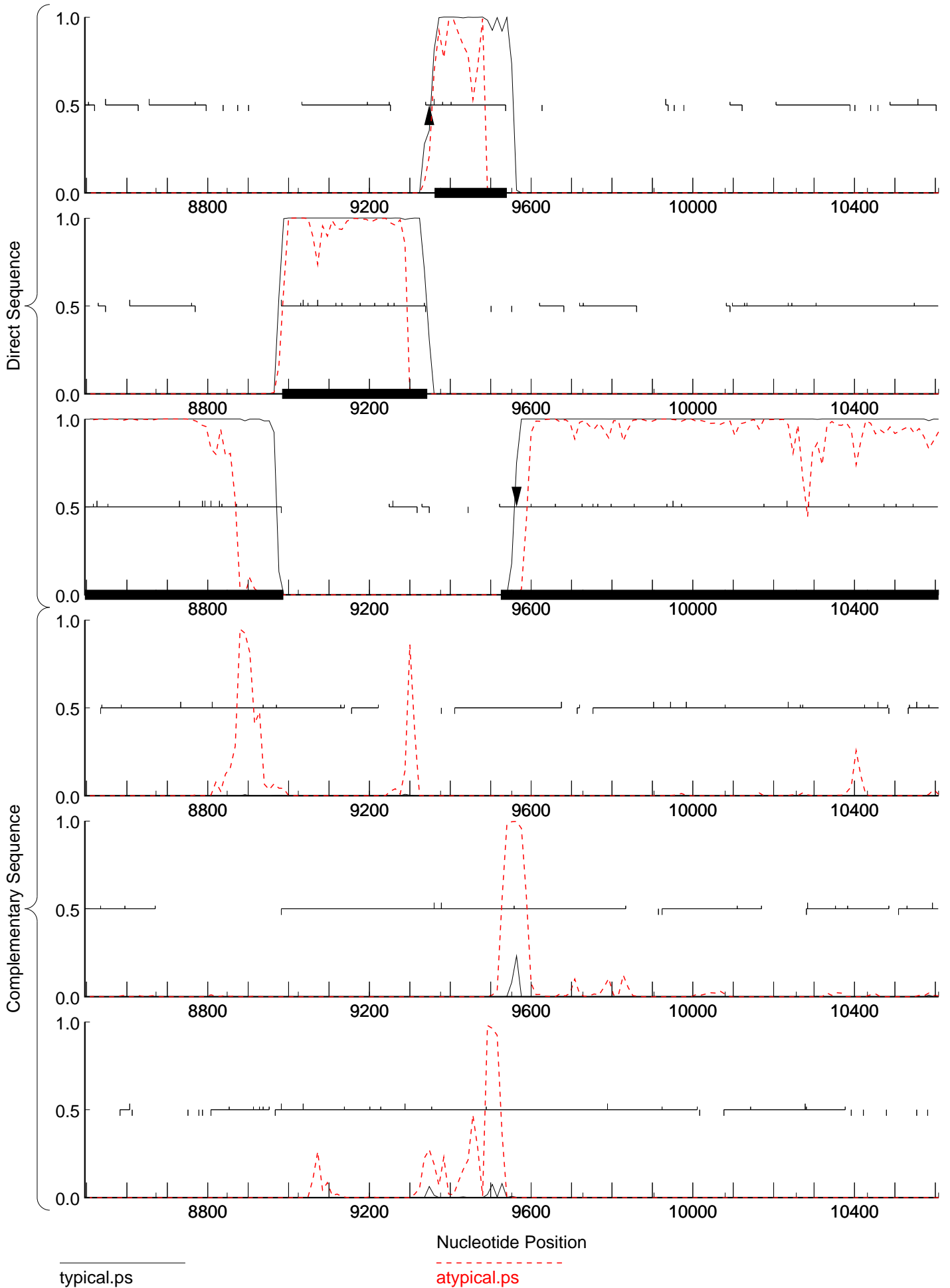


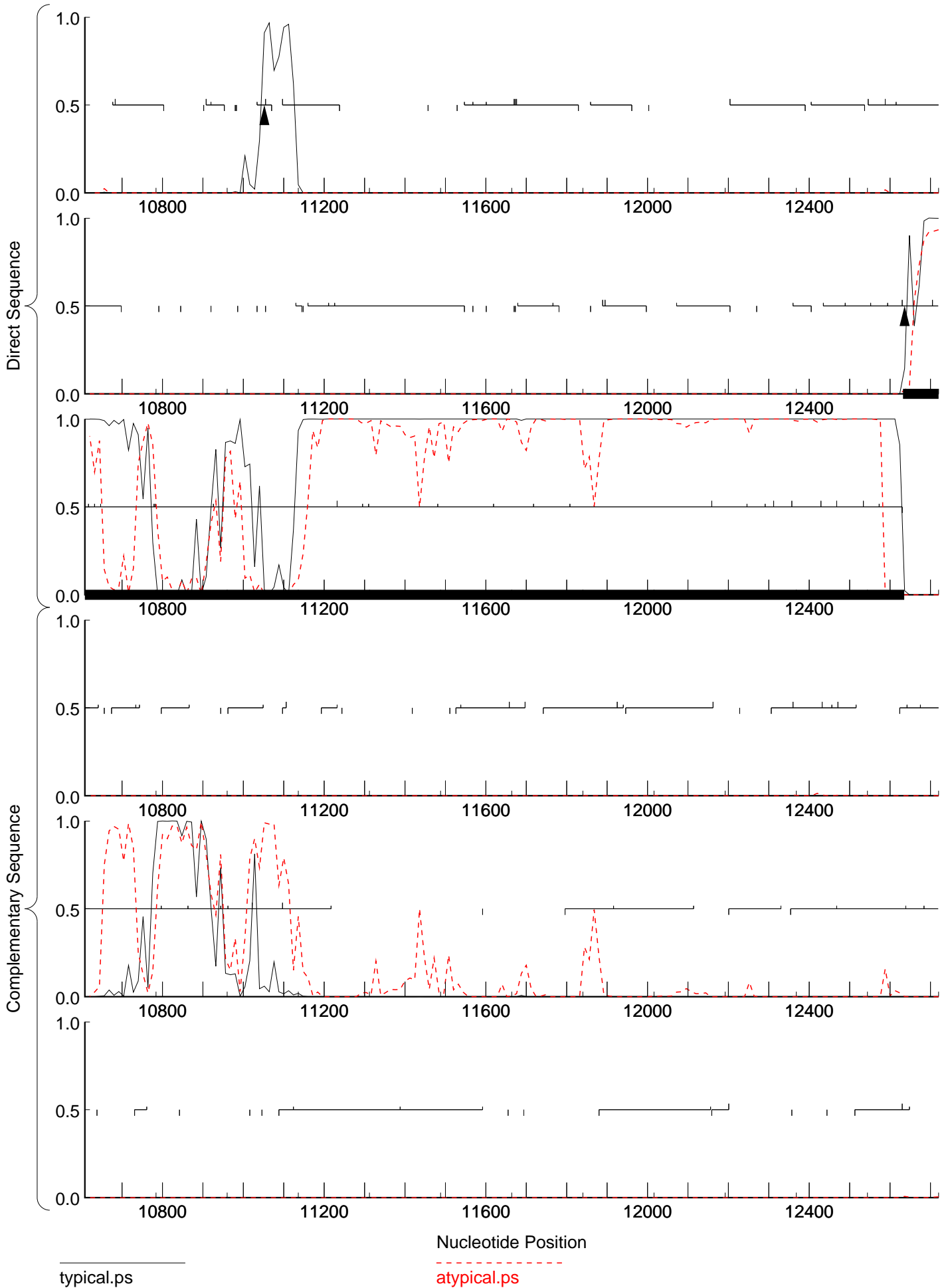
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atypical.ps



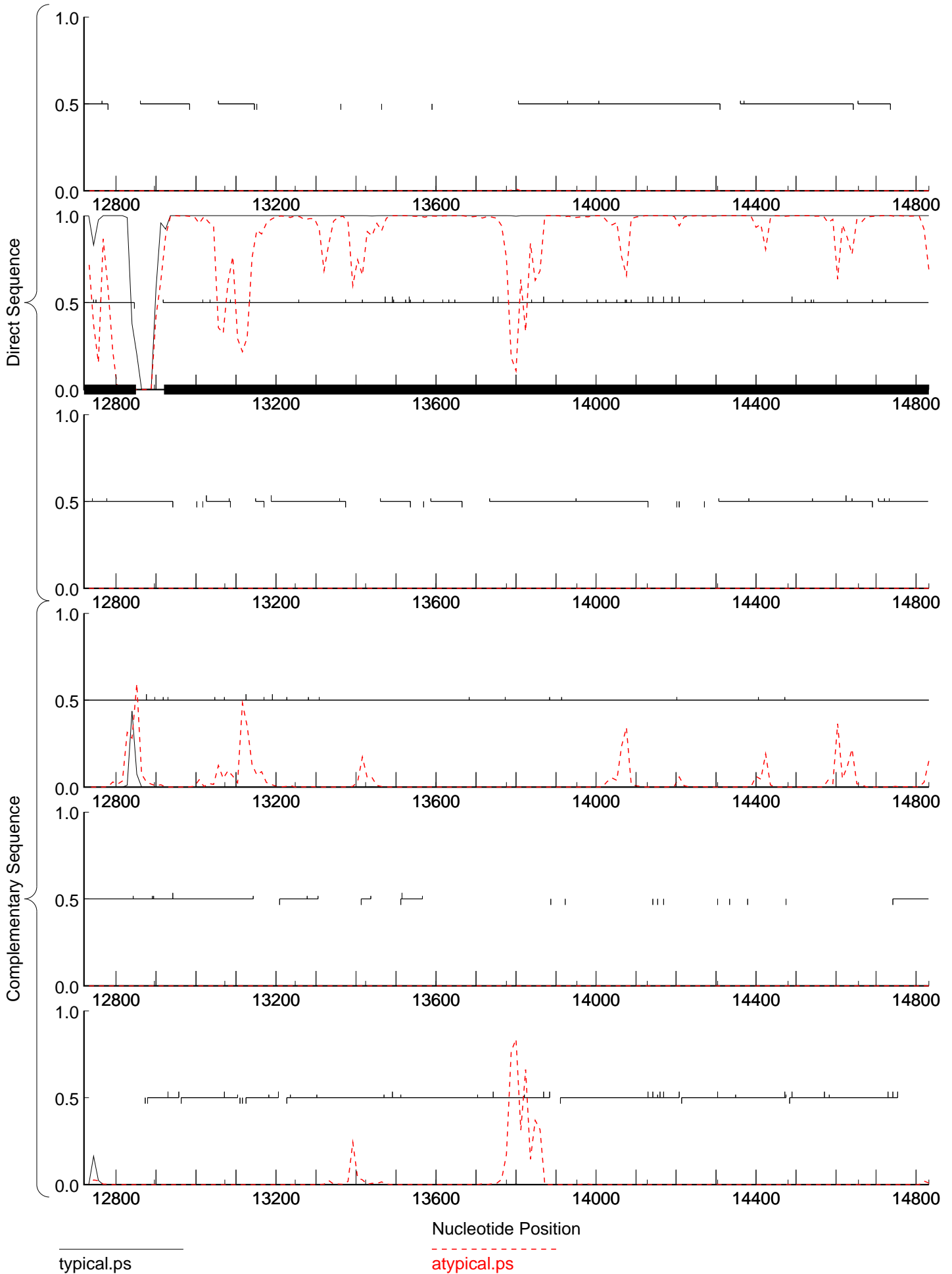




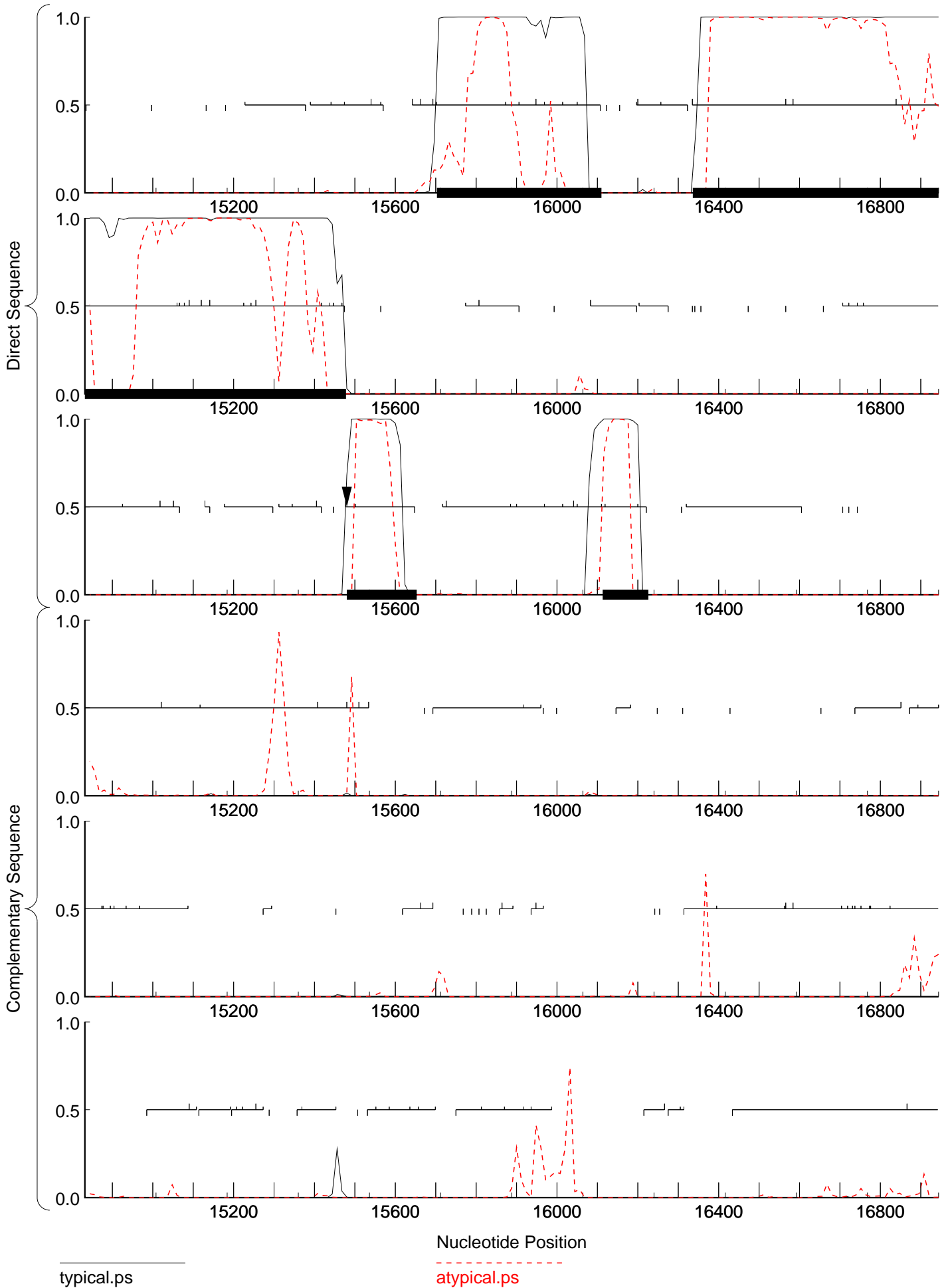


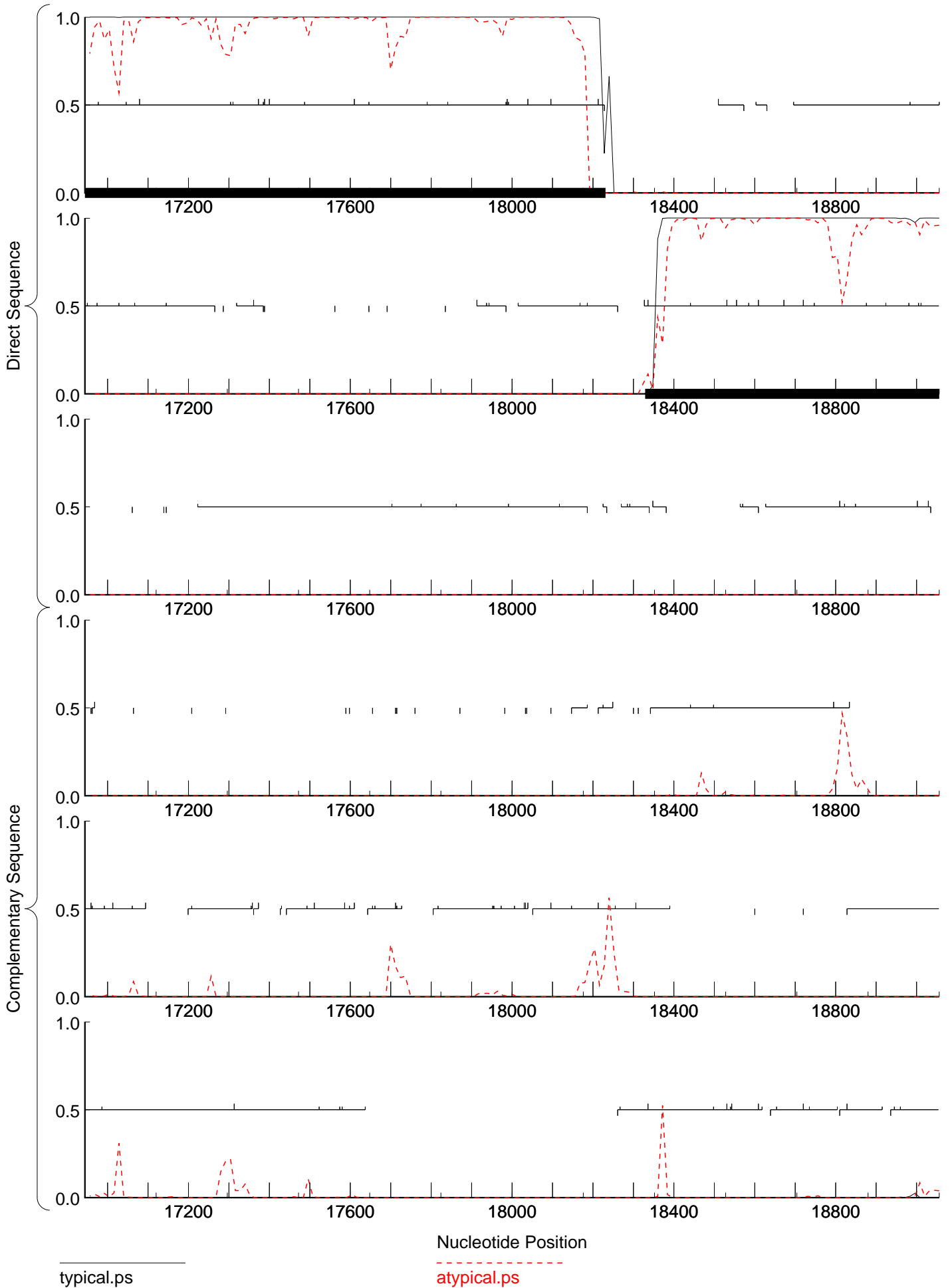
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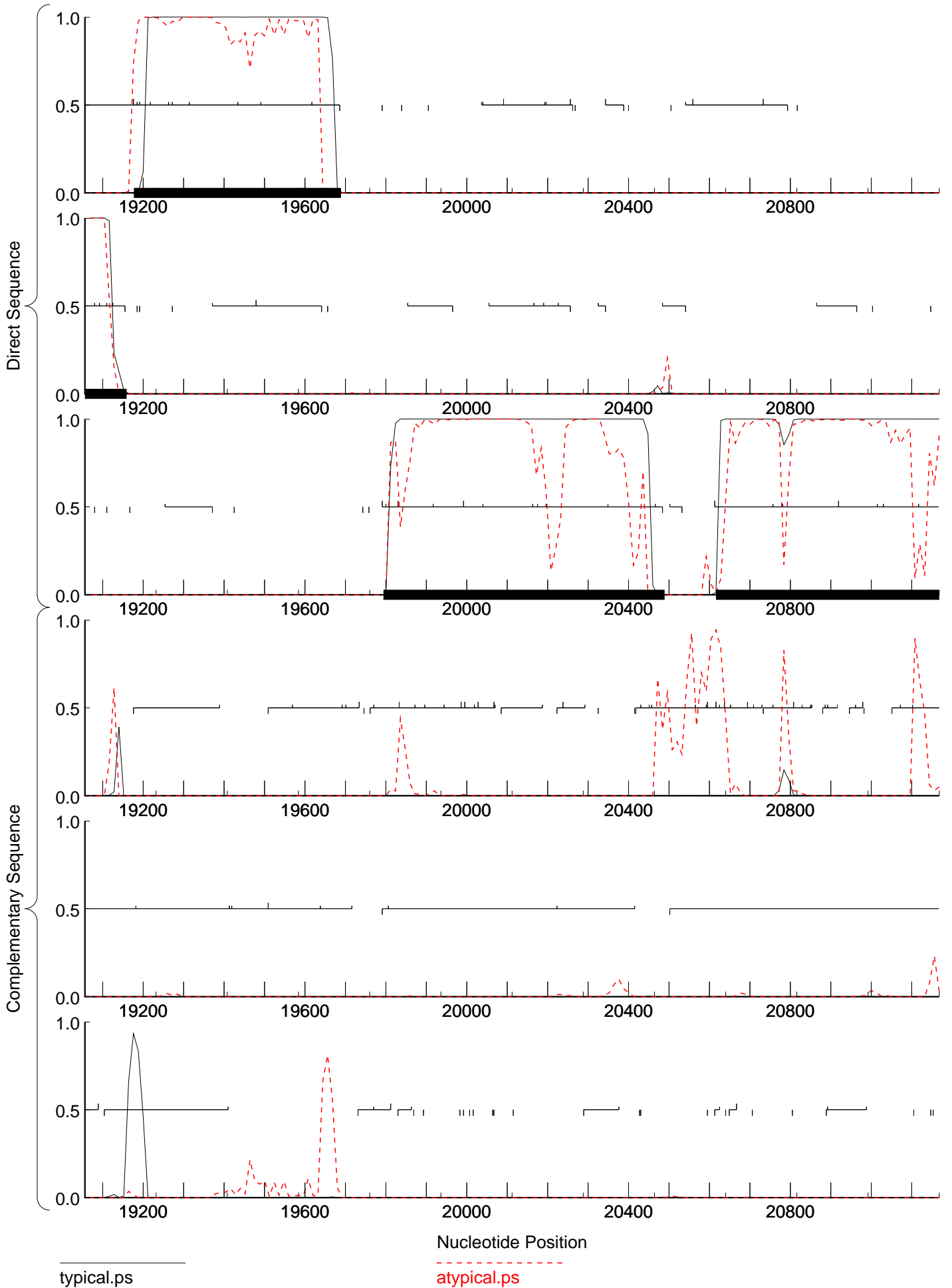
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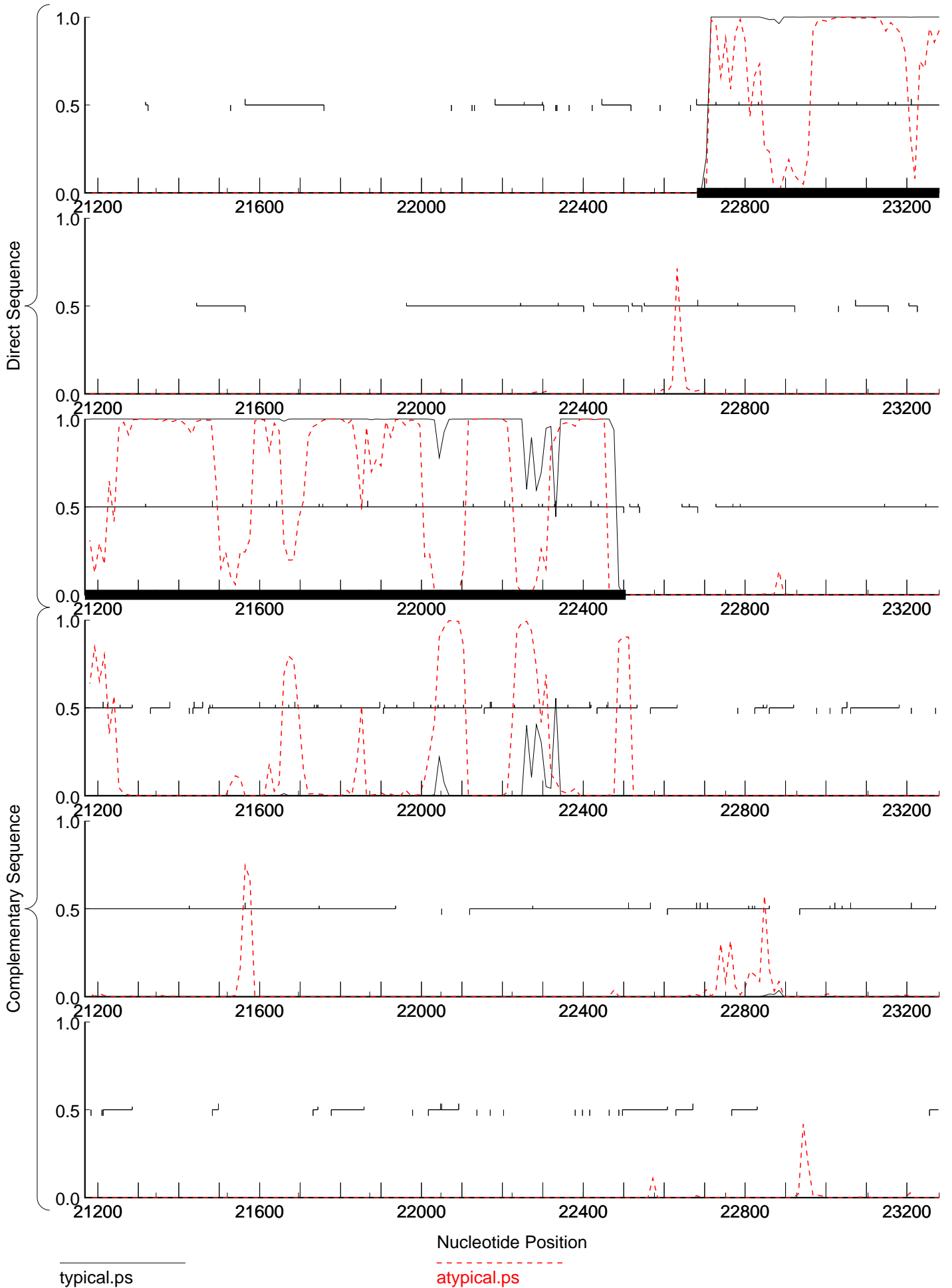


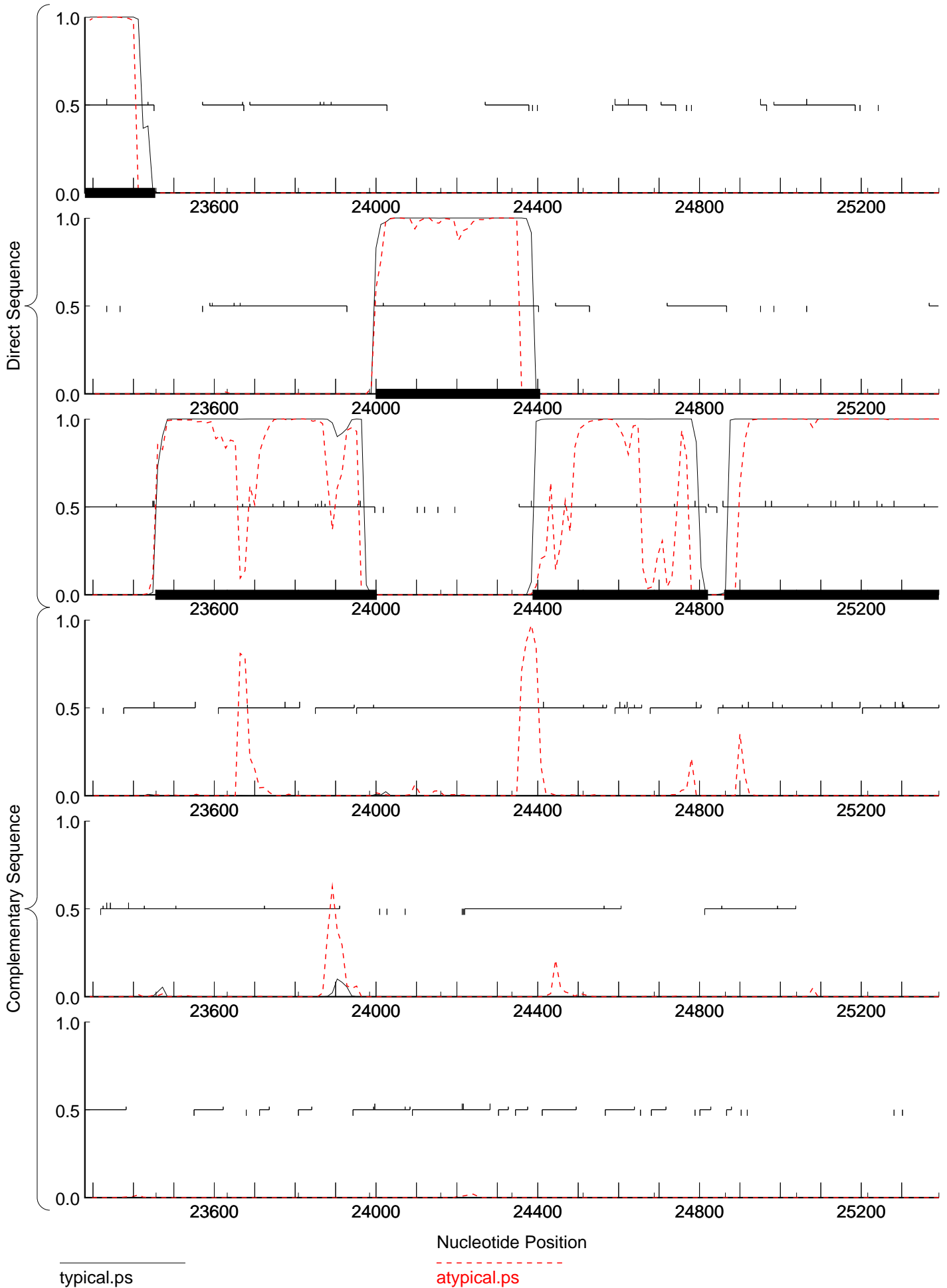


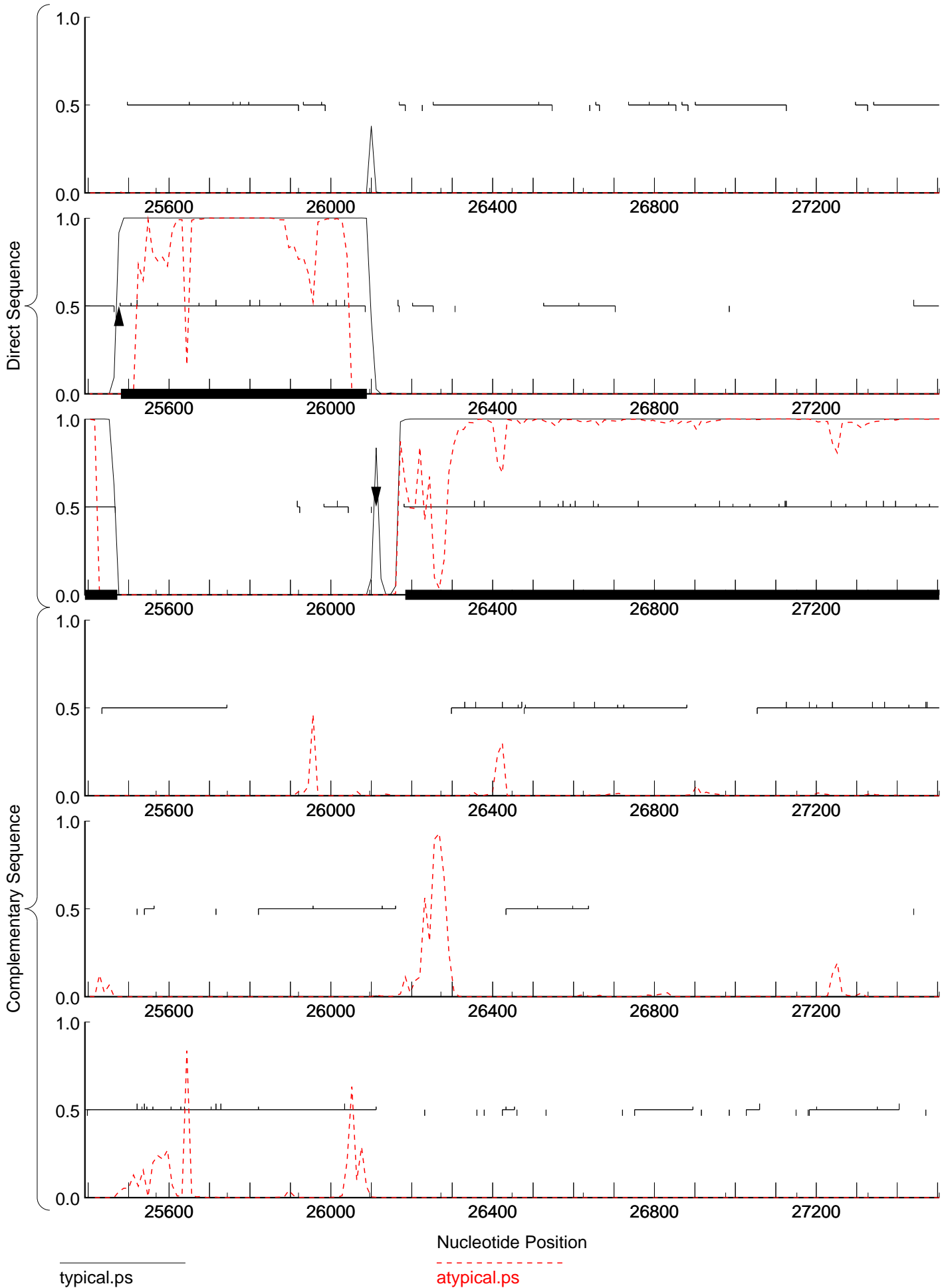


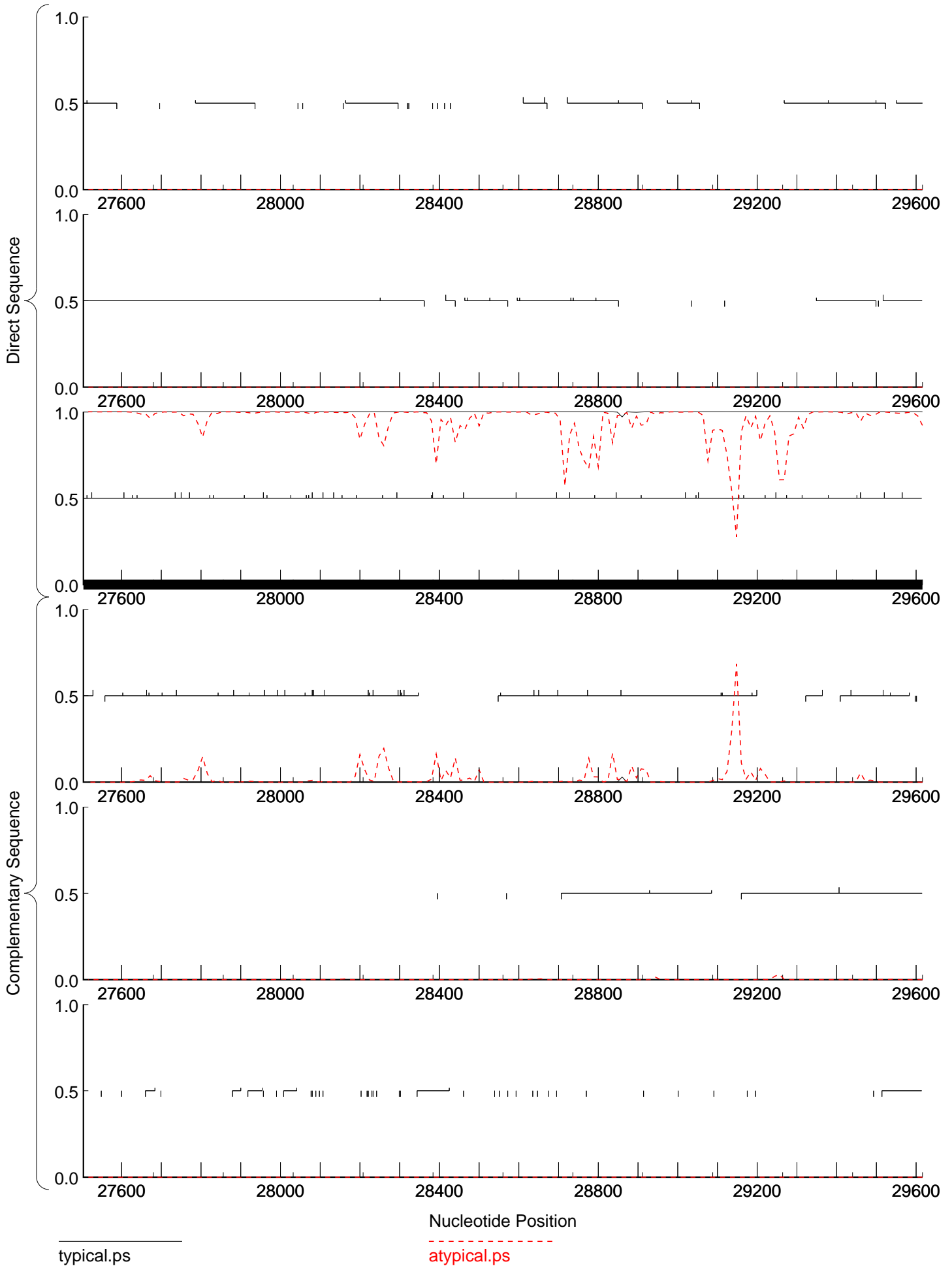


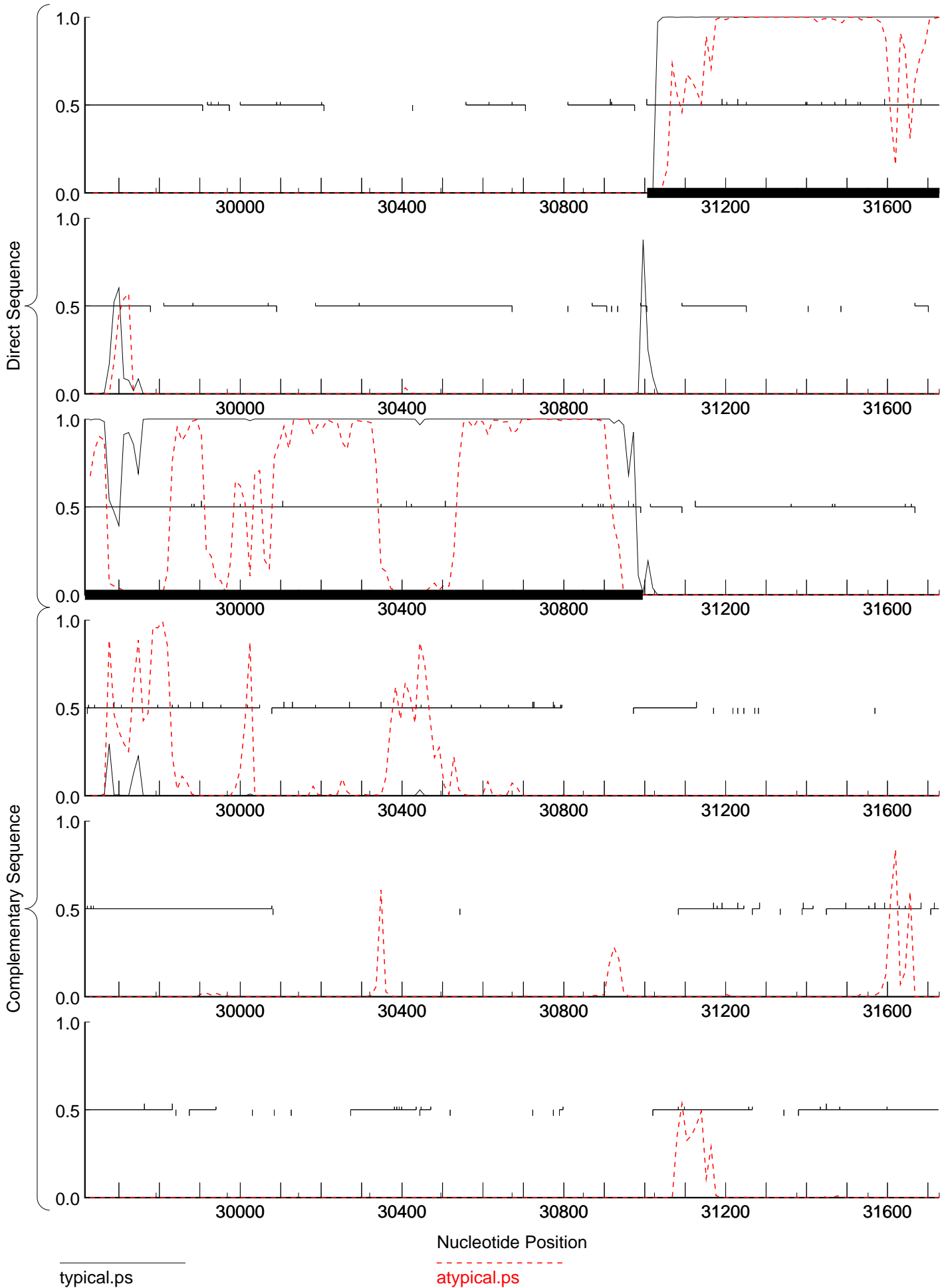




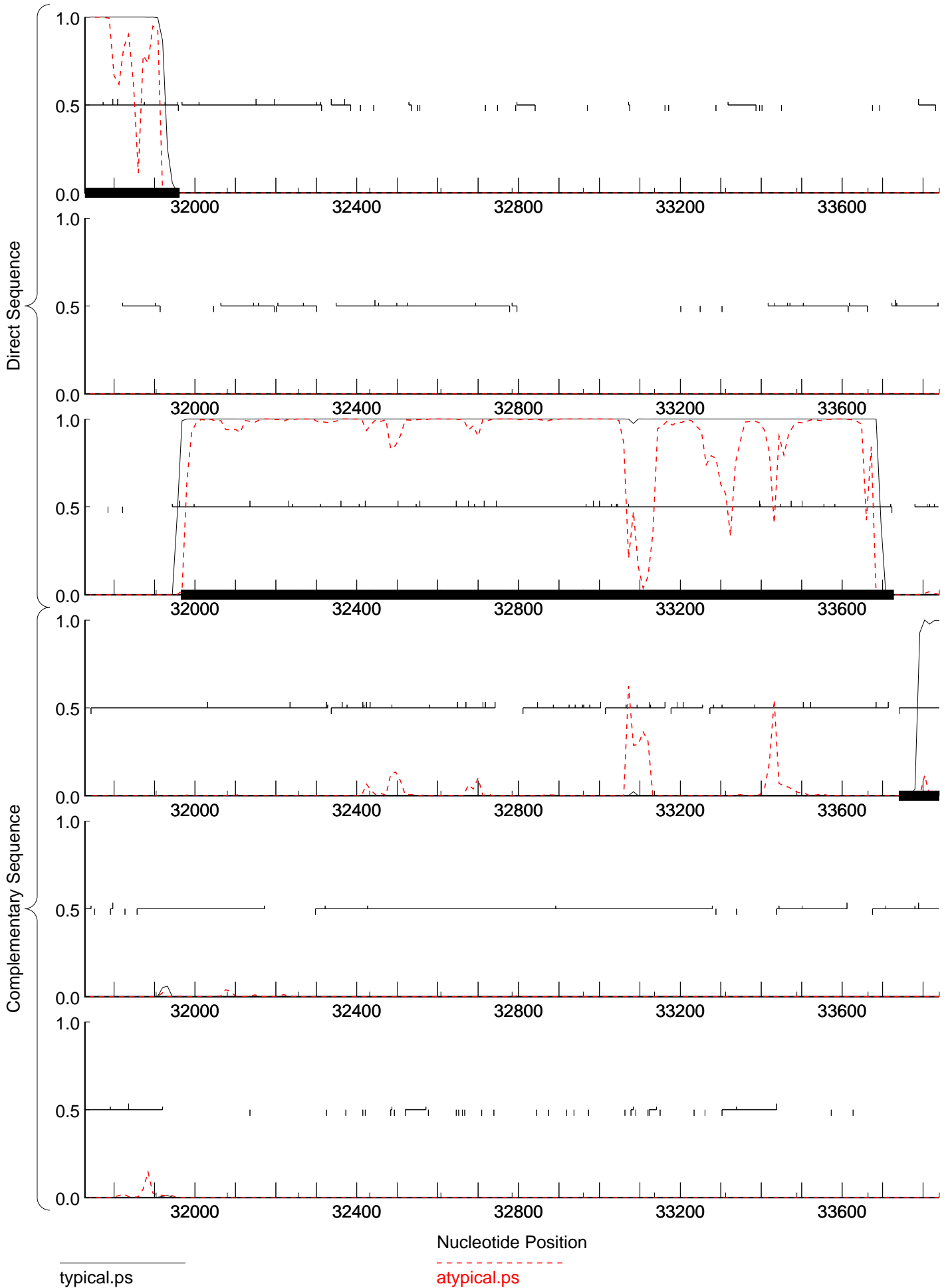






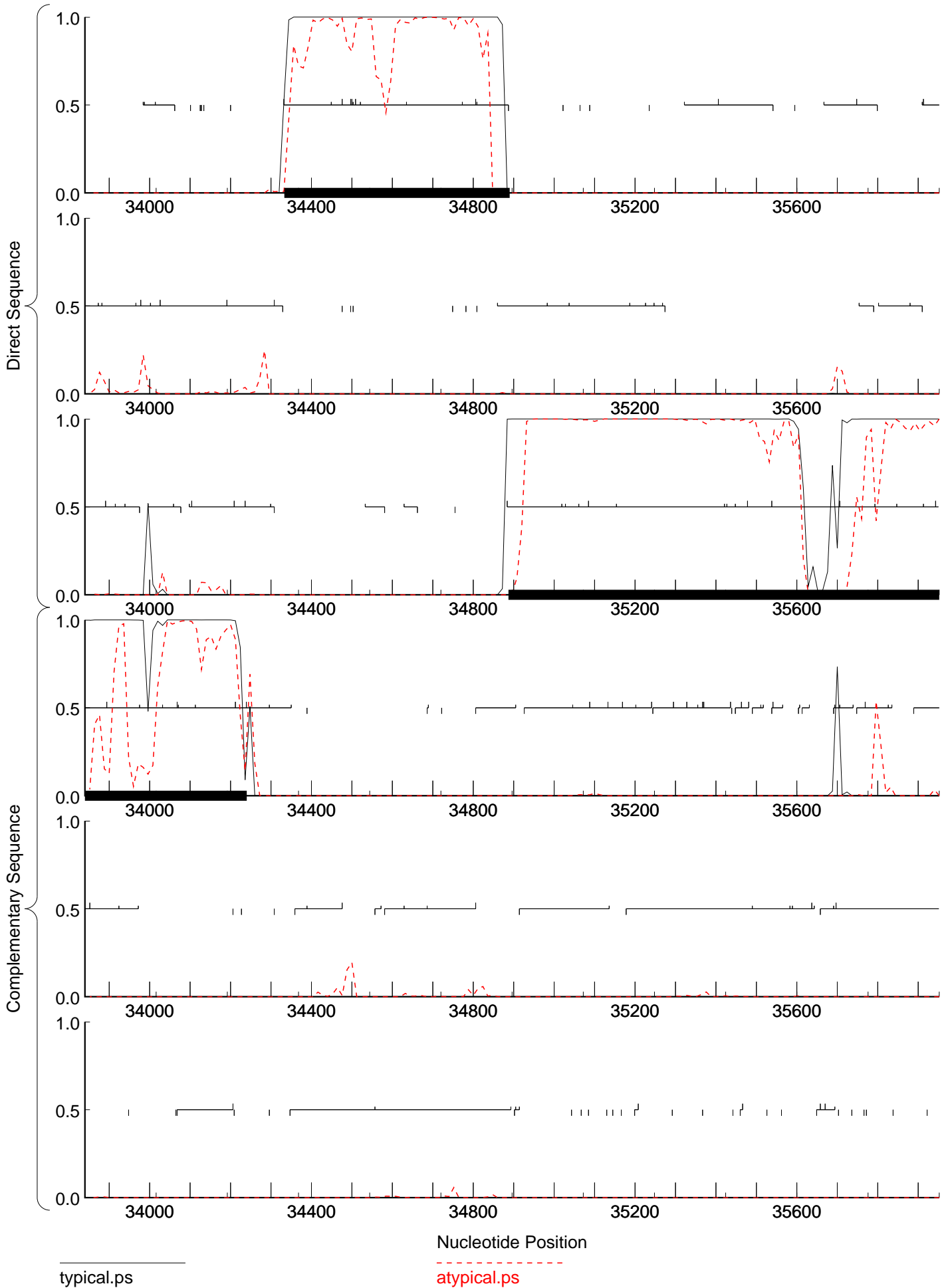


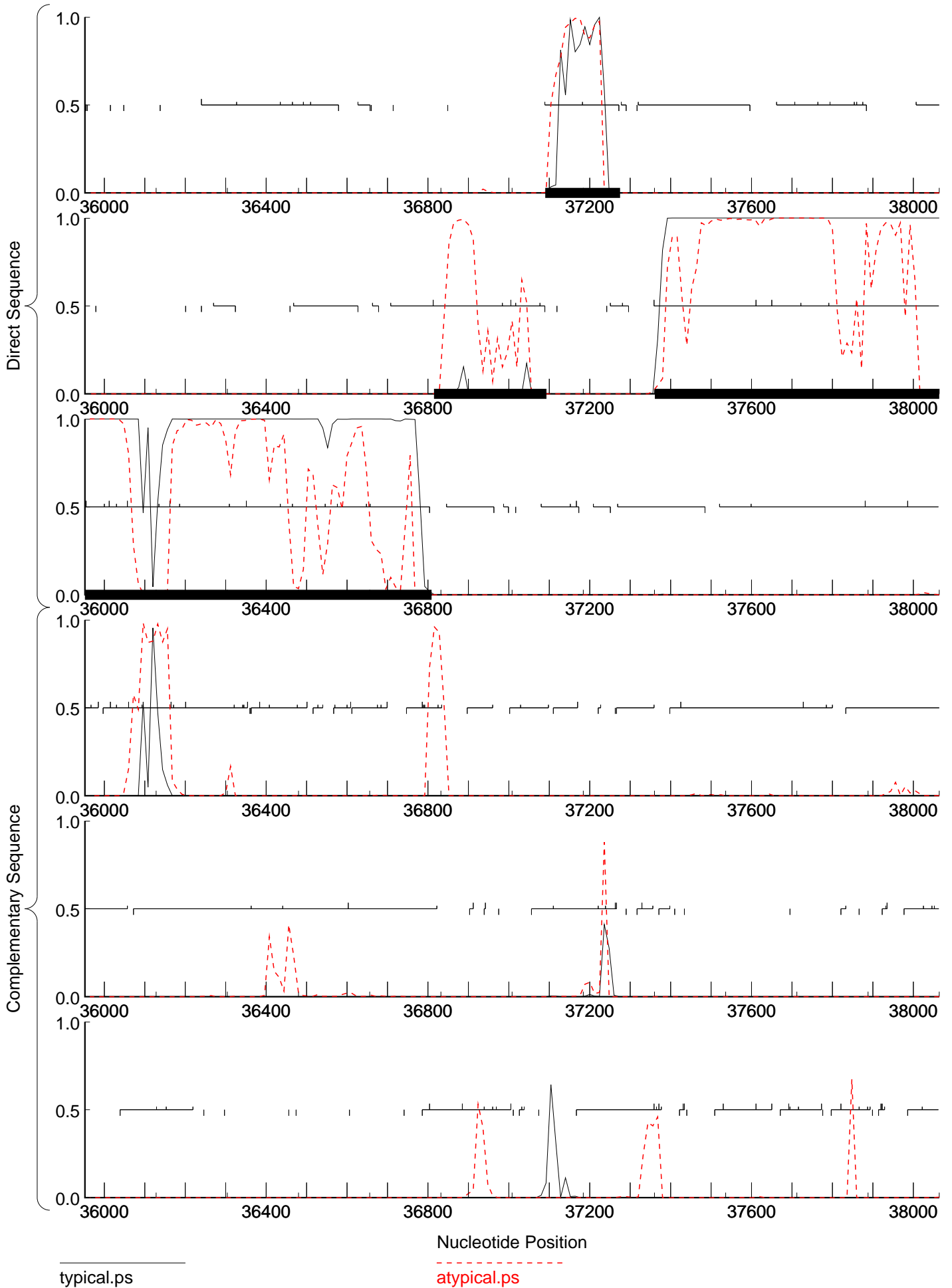


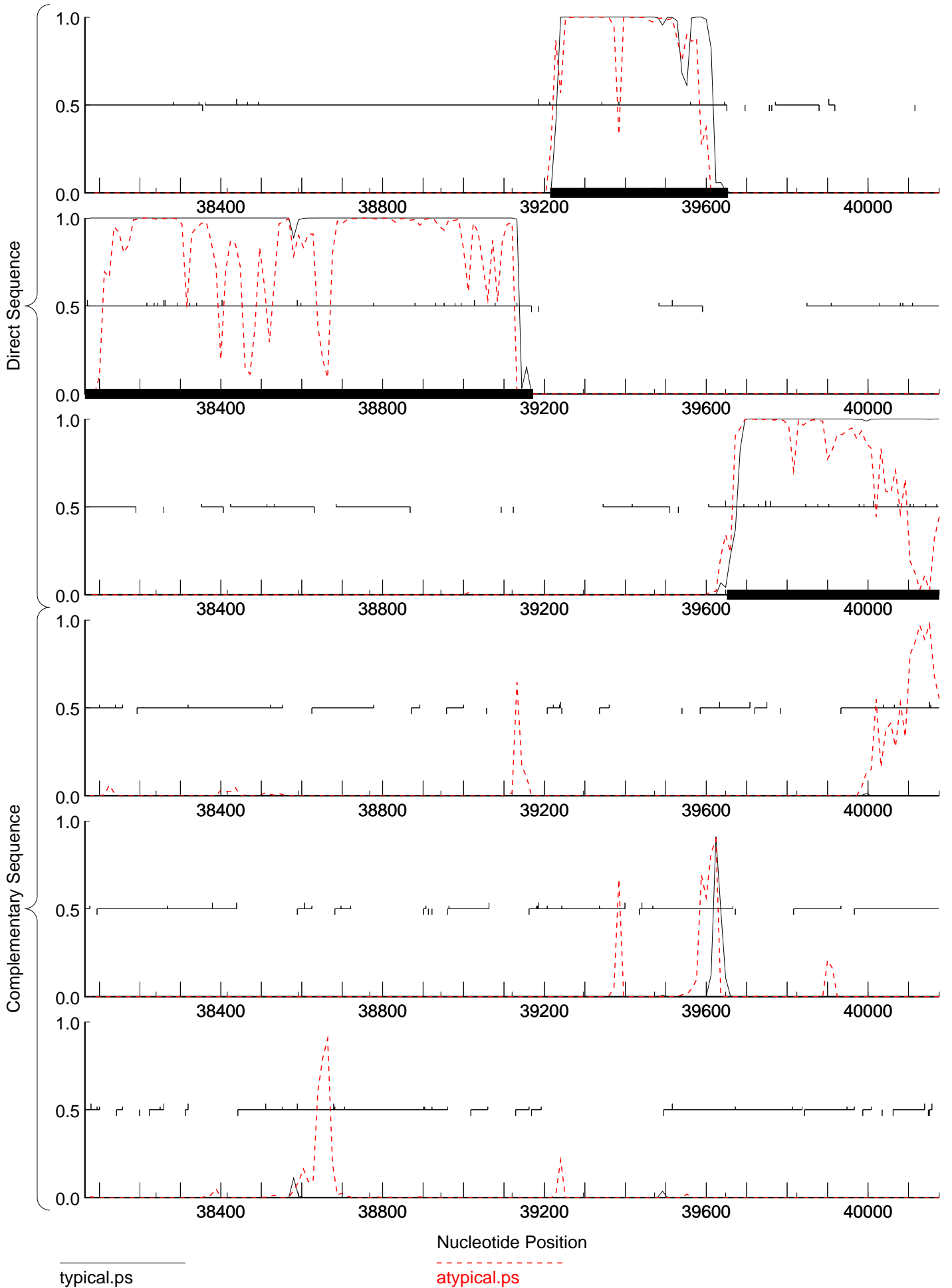


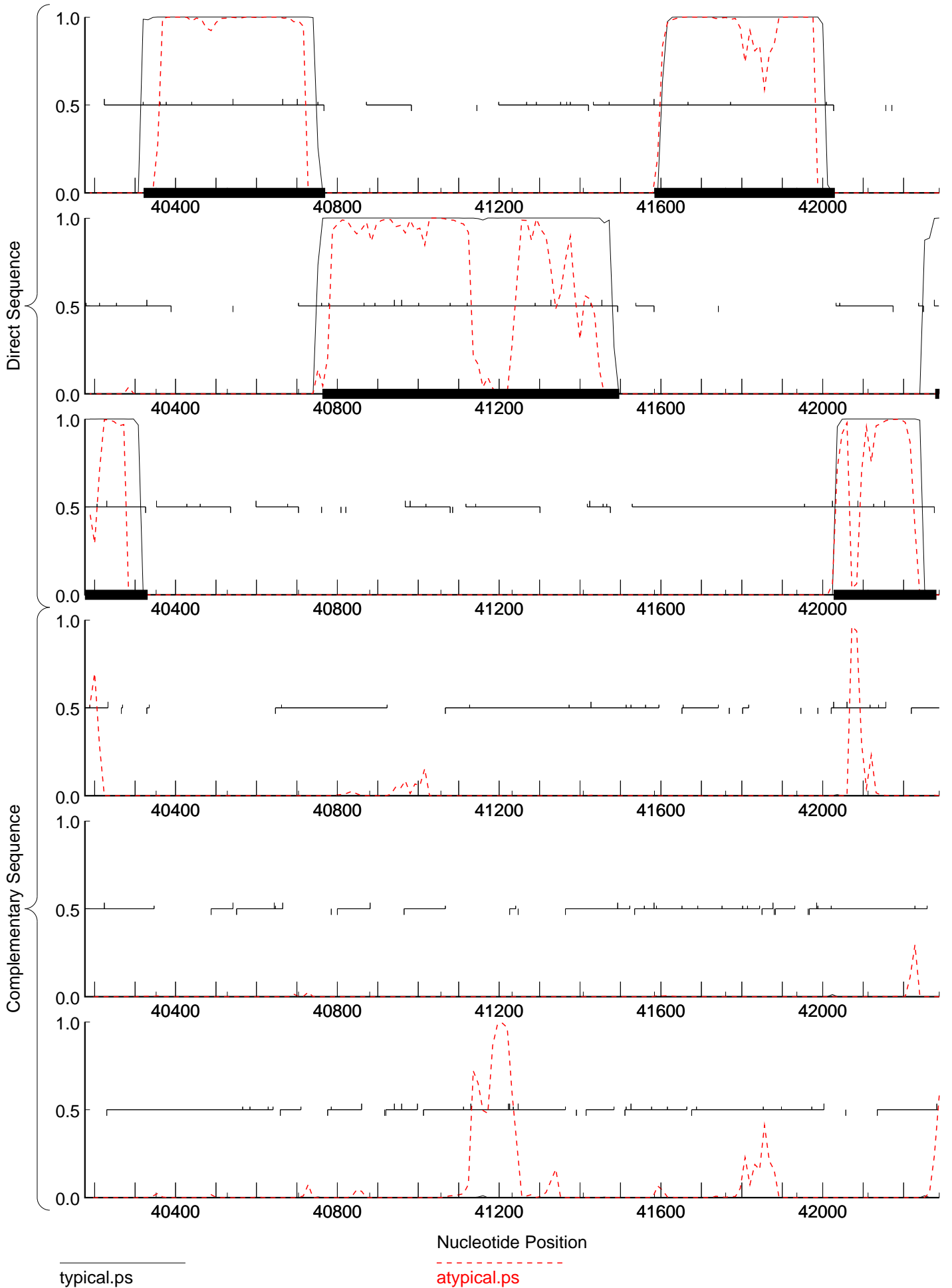
typical.ps

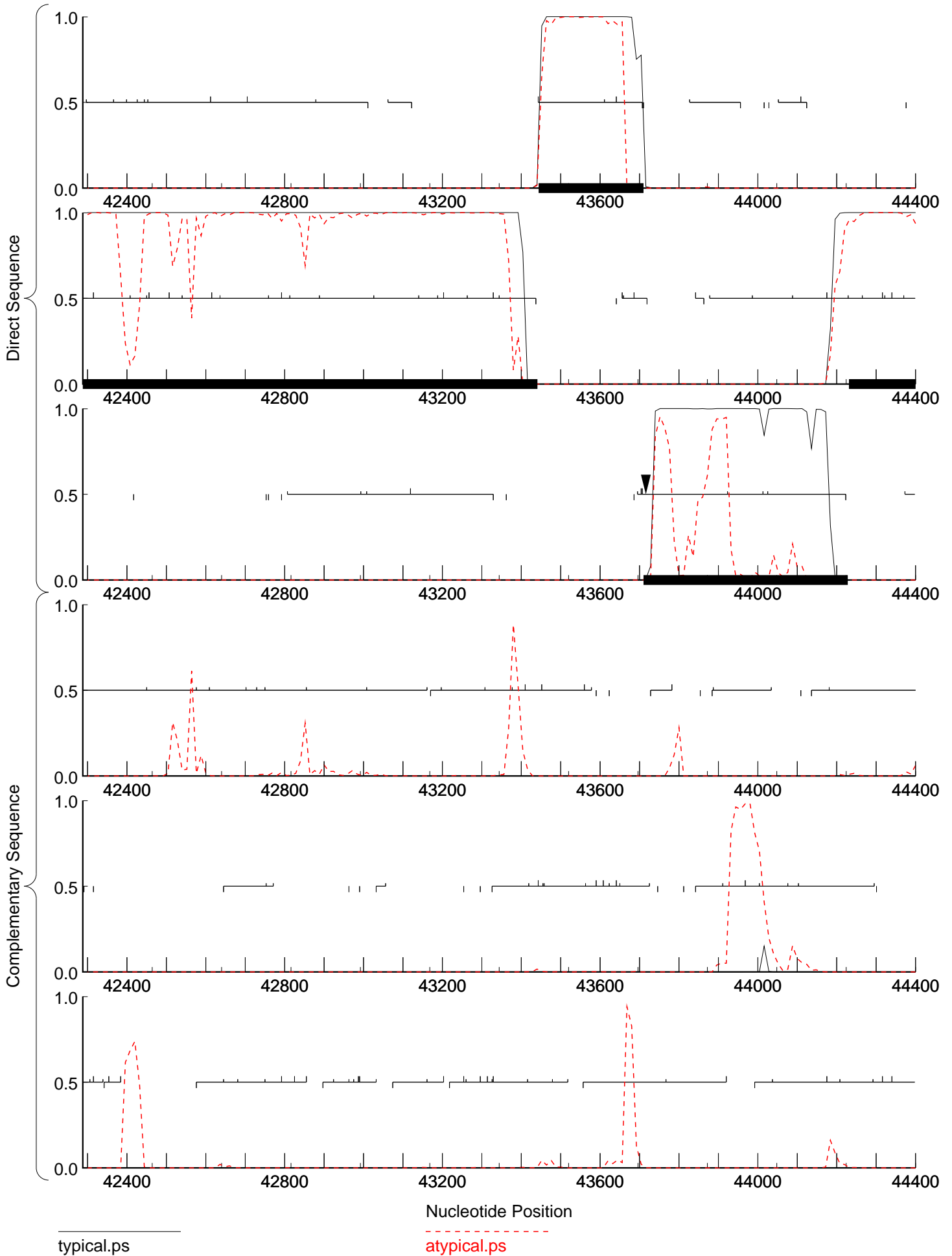
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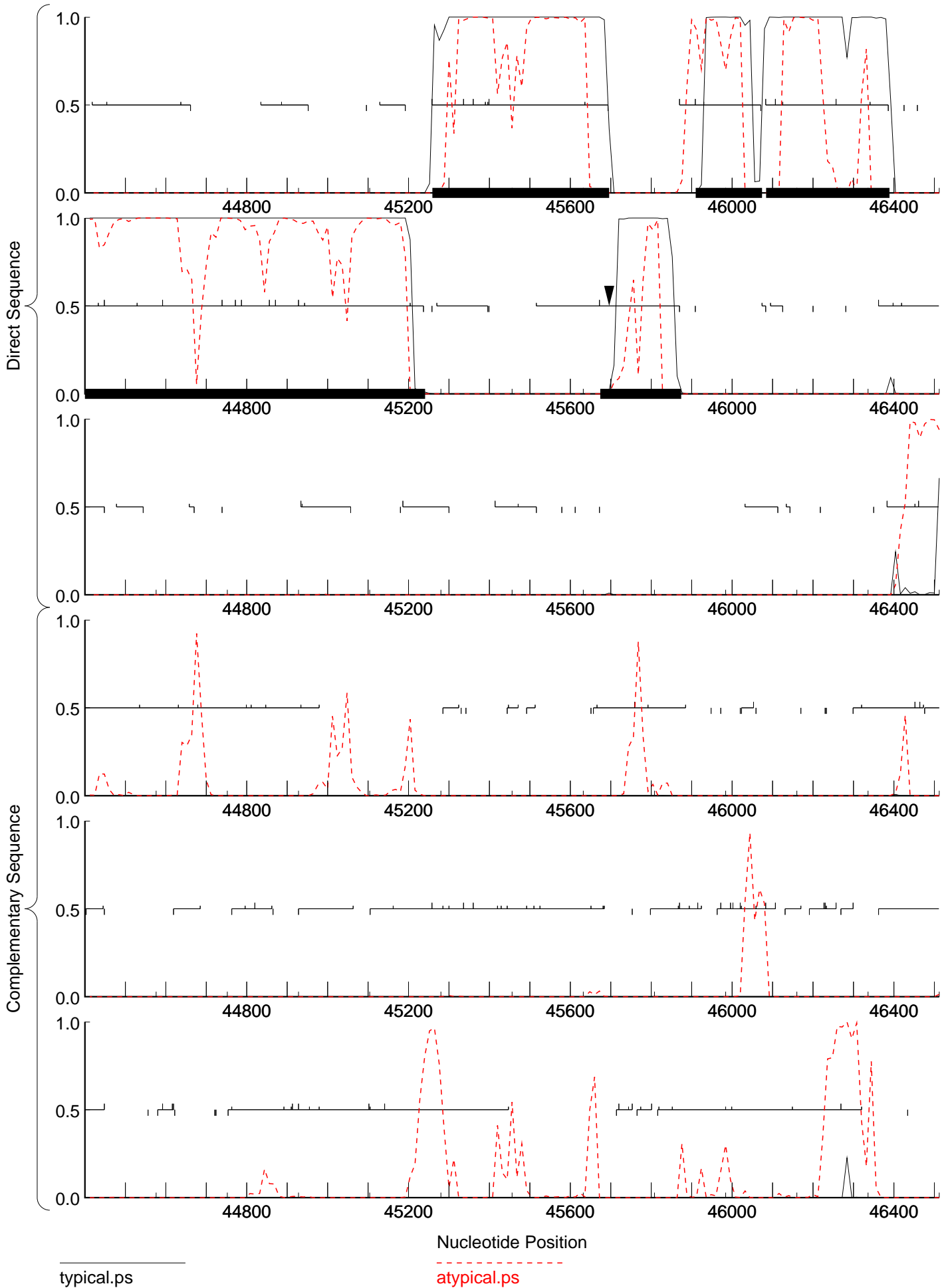






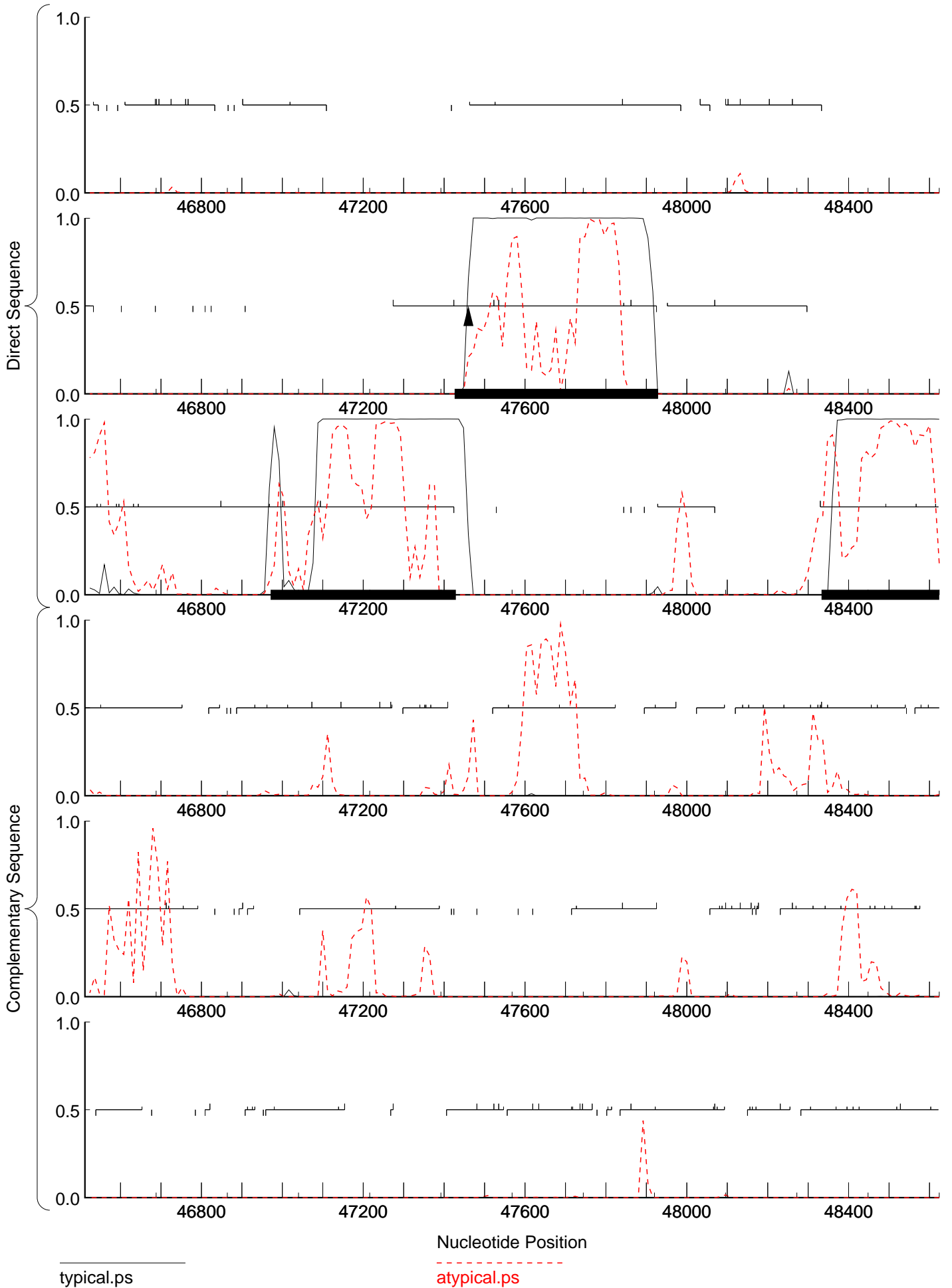
typical.ps

atypical.ps

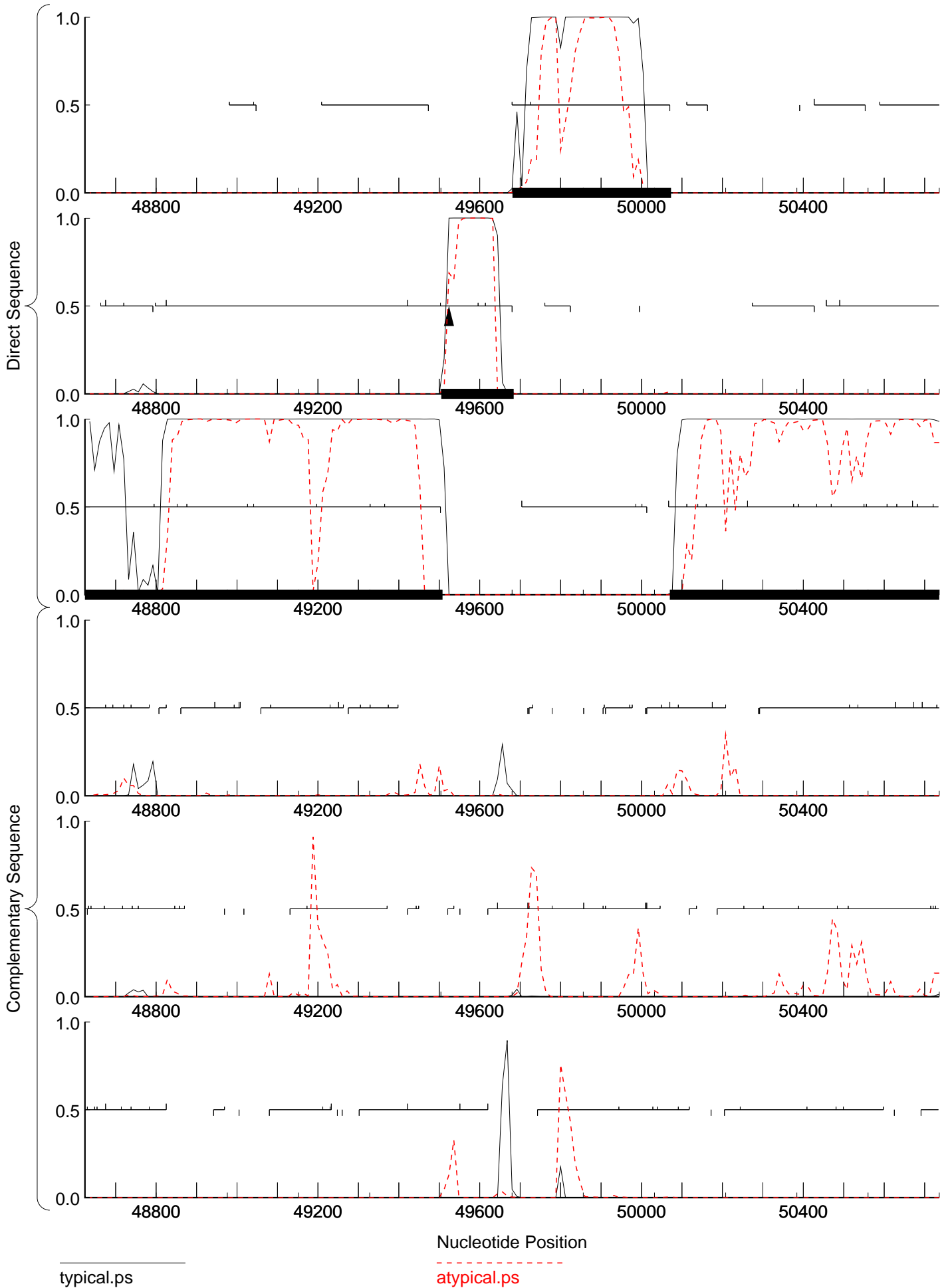


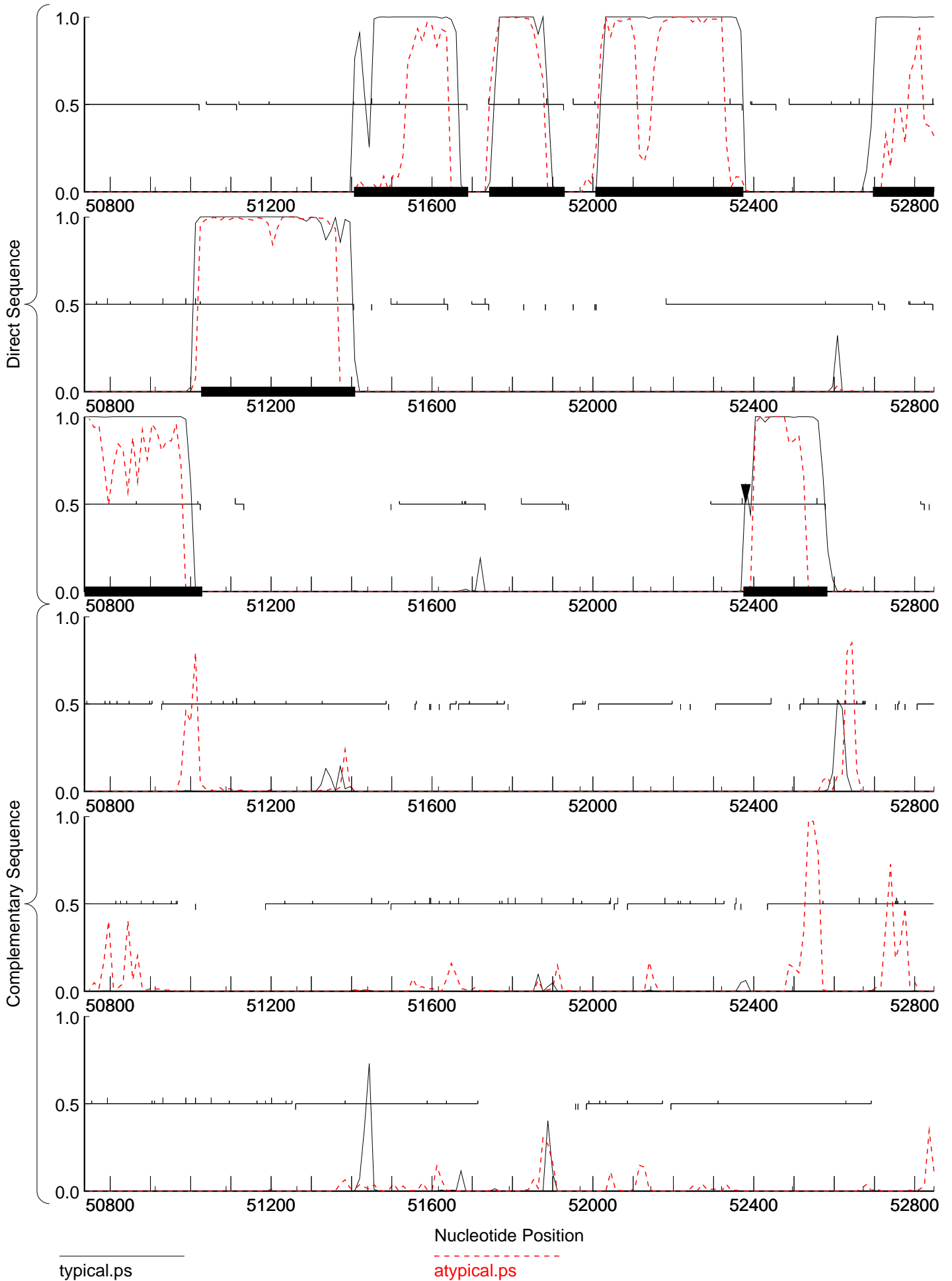
typical.ps

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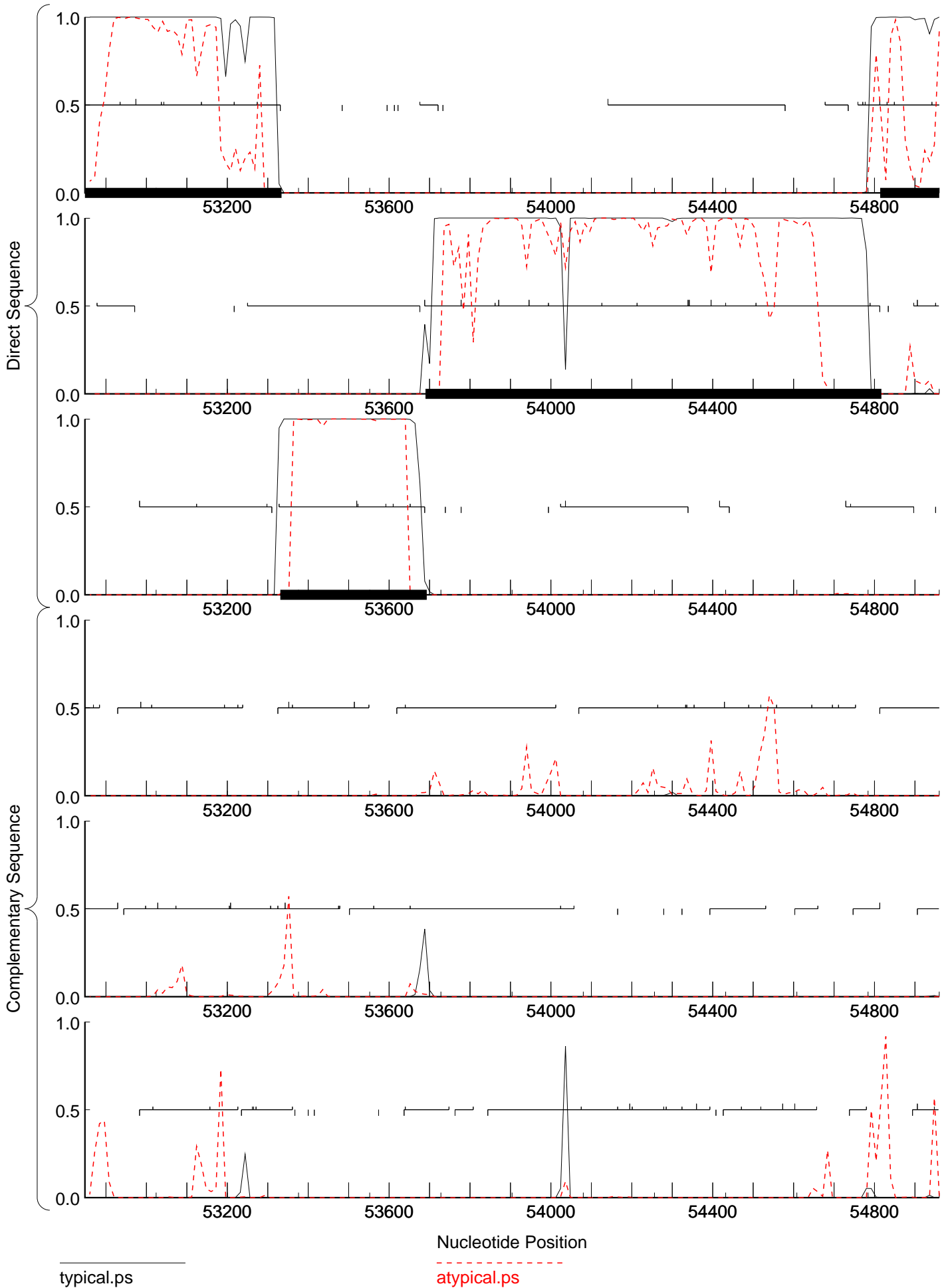


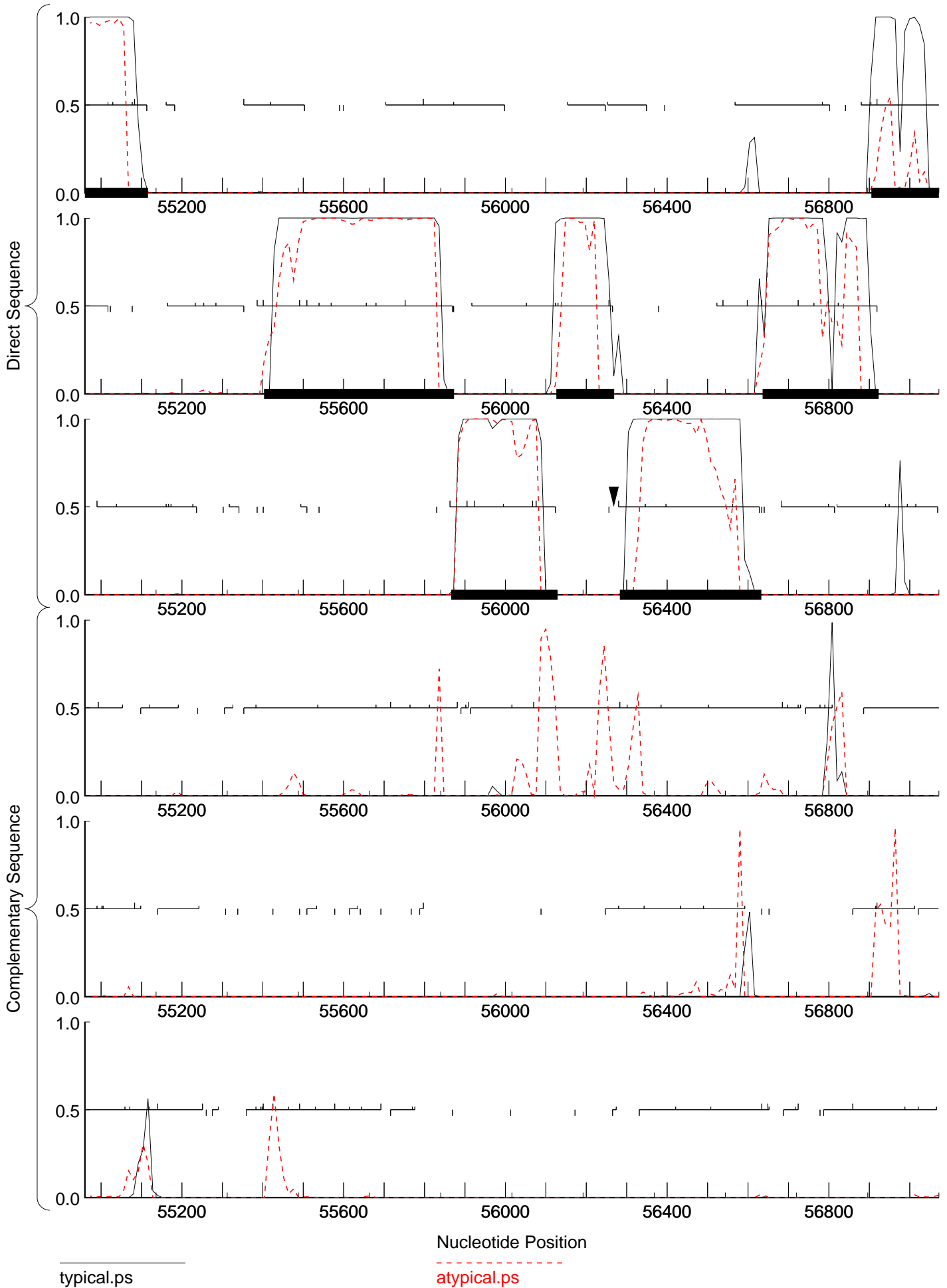




typical.ps

atypical.ps





typical.ps

atypical.ps

