

# GeneMark

Version 2.5p (09.08.06)

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## PROGRAM INFORMATION

Sequence : Streptomyces phage BayC complete sequence, 57243 bp, circularly permuted, Cluster BG  
Analysis Date : 11/23/17 at 6:10:47  
Pages : 29  
Sequence Length : 57243 bp  
GC Content : 69.23%

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,

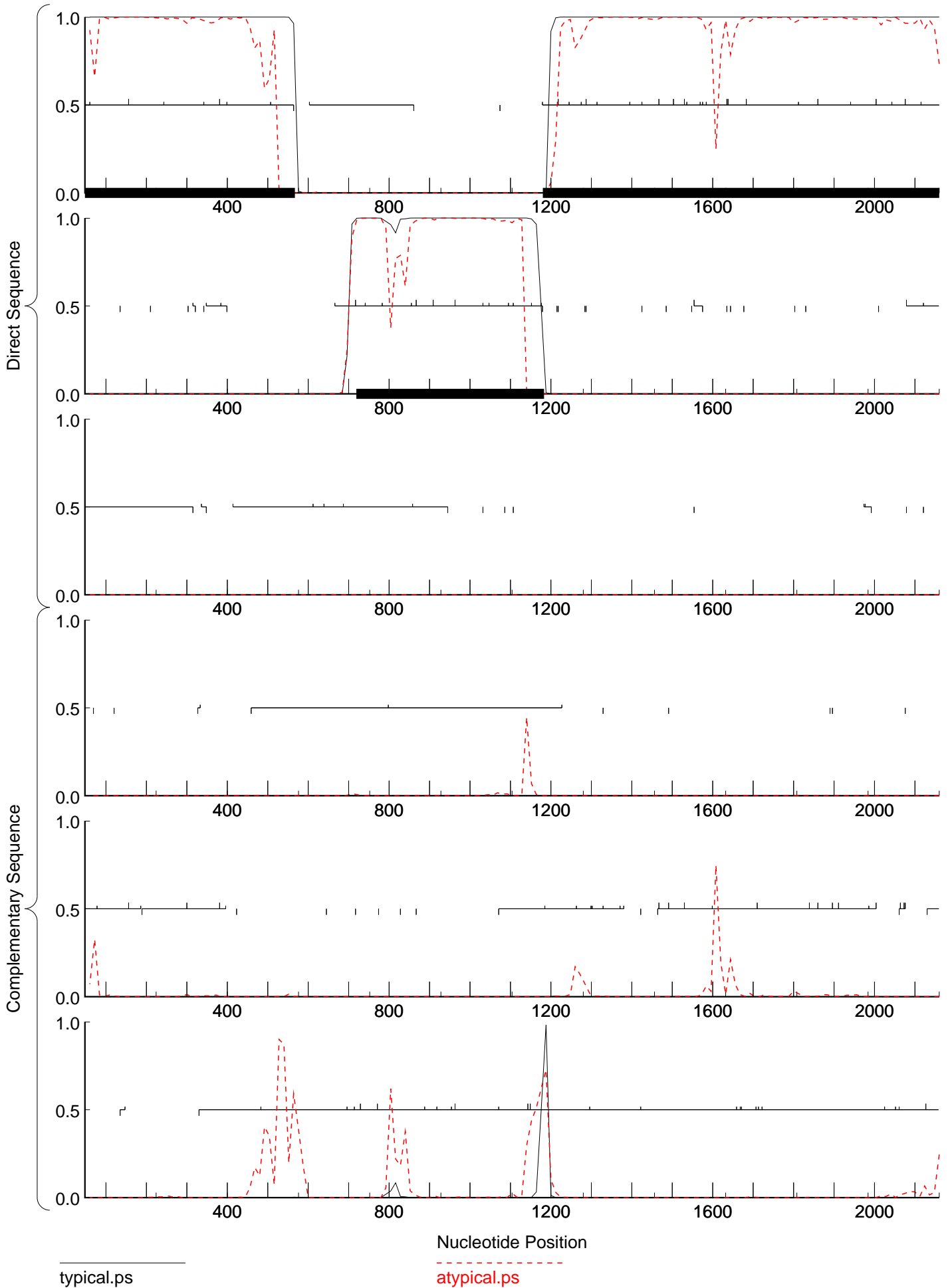
## Matrix notes & comments

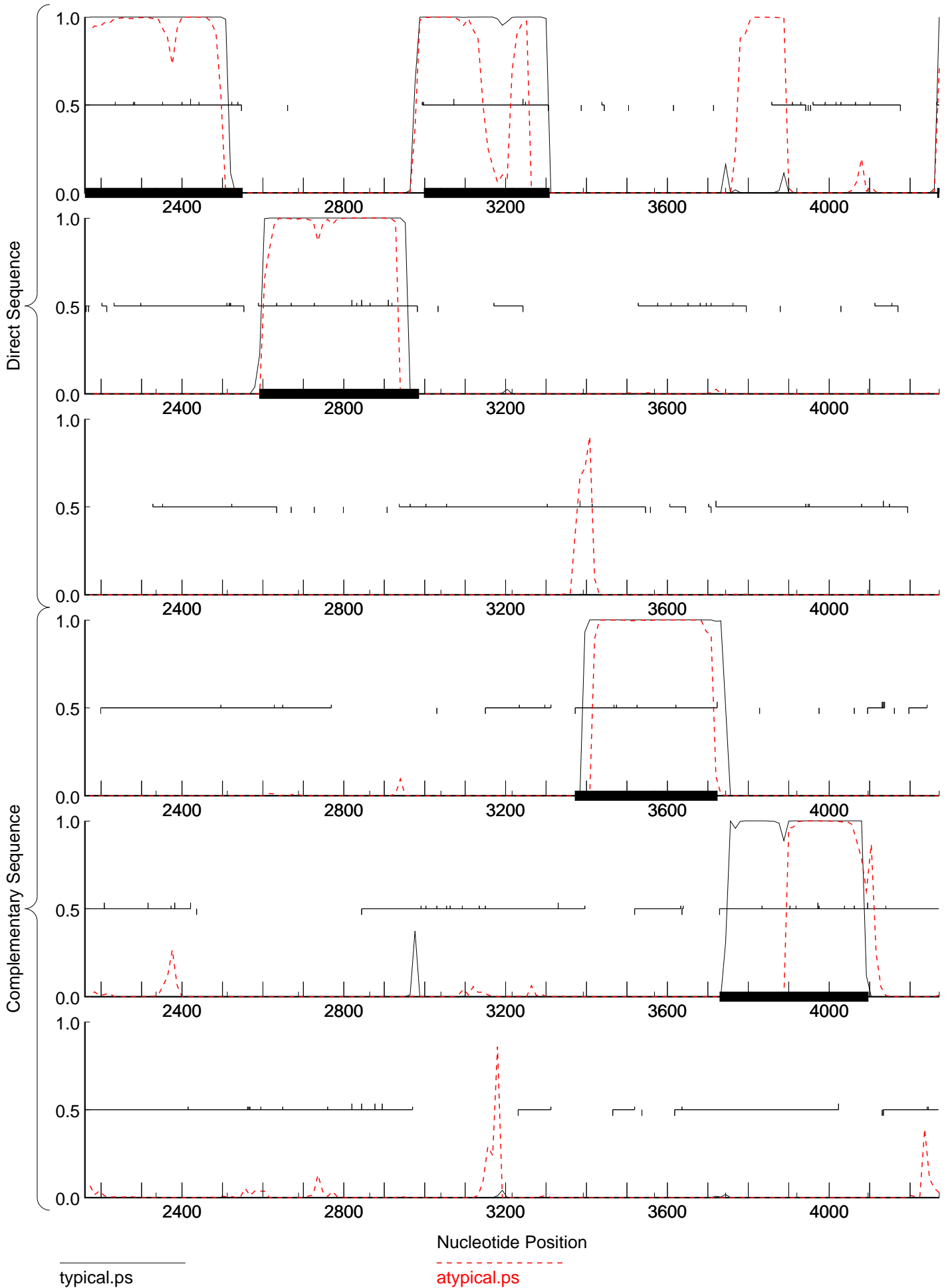
Training set derived by GeneMarkS, 4.30 October 2014  
Thu Nov 23 06:10:47 2017

## MATRIX INFORMATION

Matrix : -  
Author : -  
Order : 4

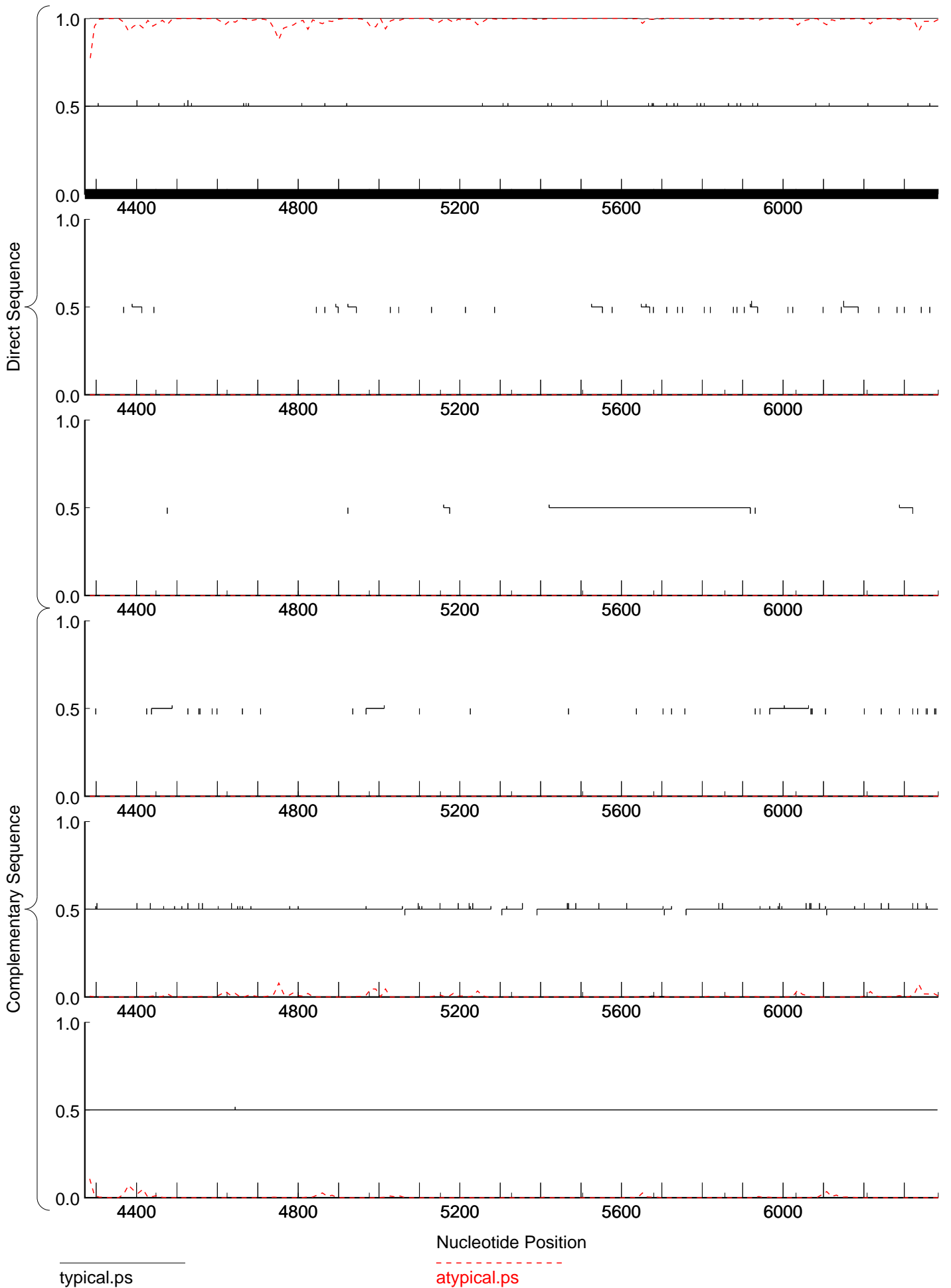
Send questions / comments to:  
Dr. M. Borodovsky  
Georgia Institute of Technology  
School of Biology  
Atlanta, GA 30332-0230

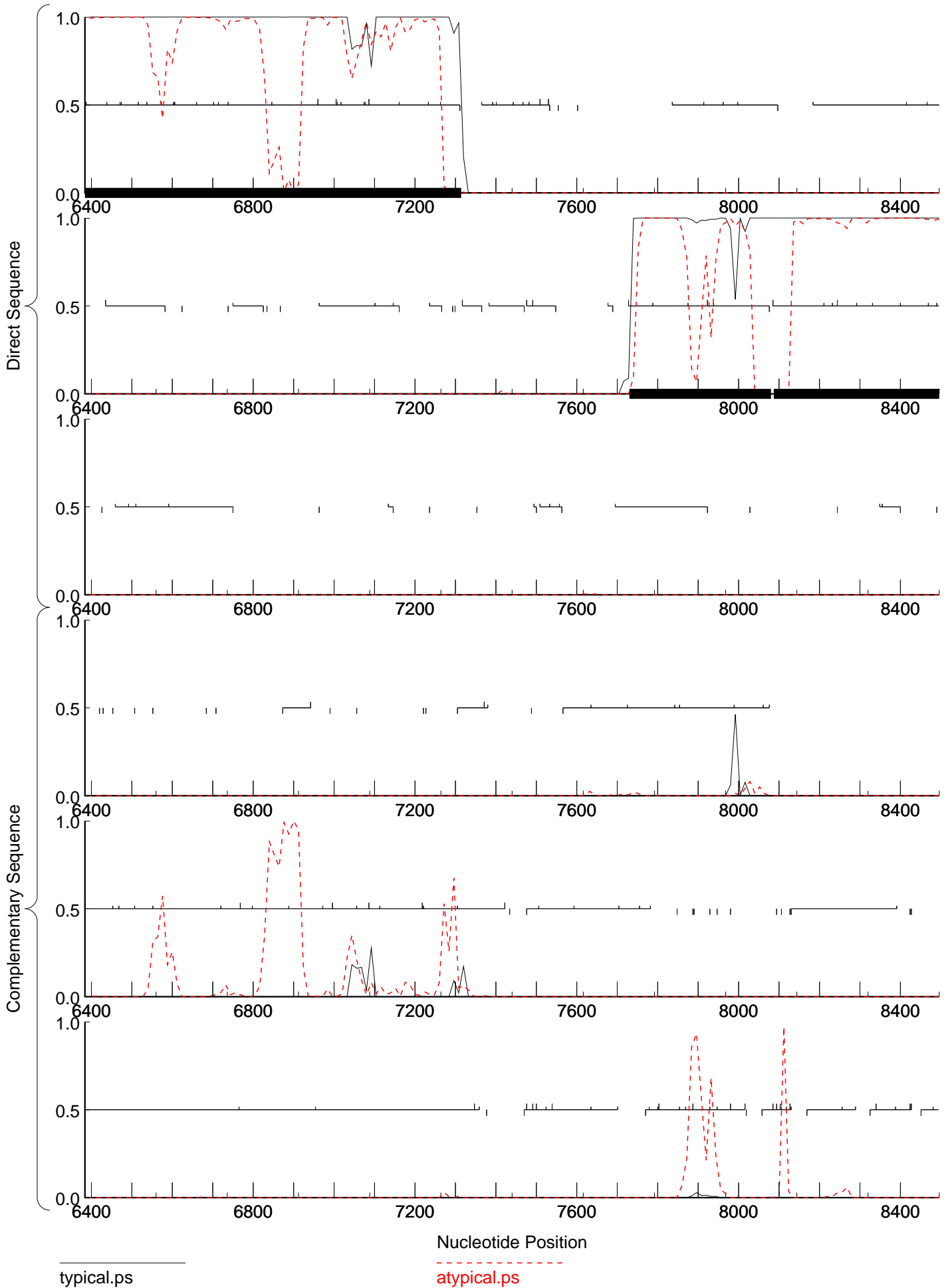




typical.ps

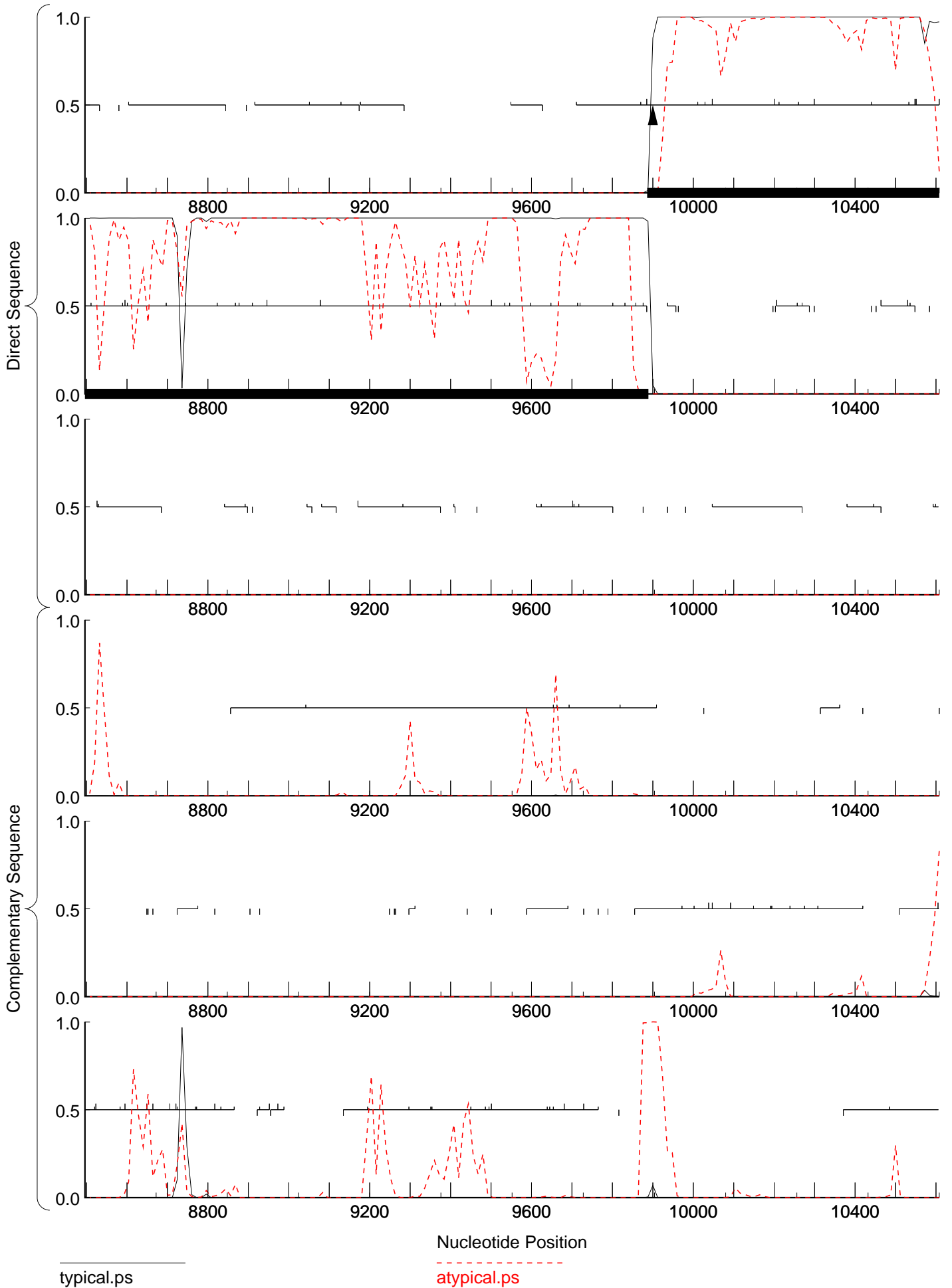
atypical.ps

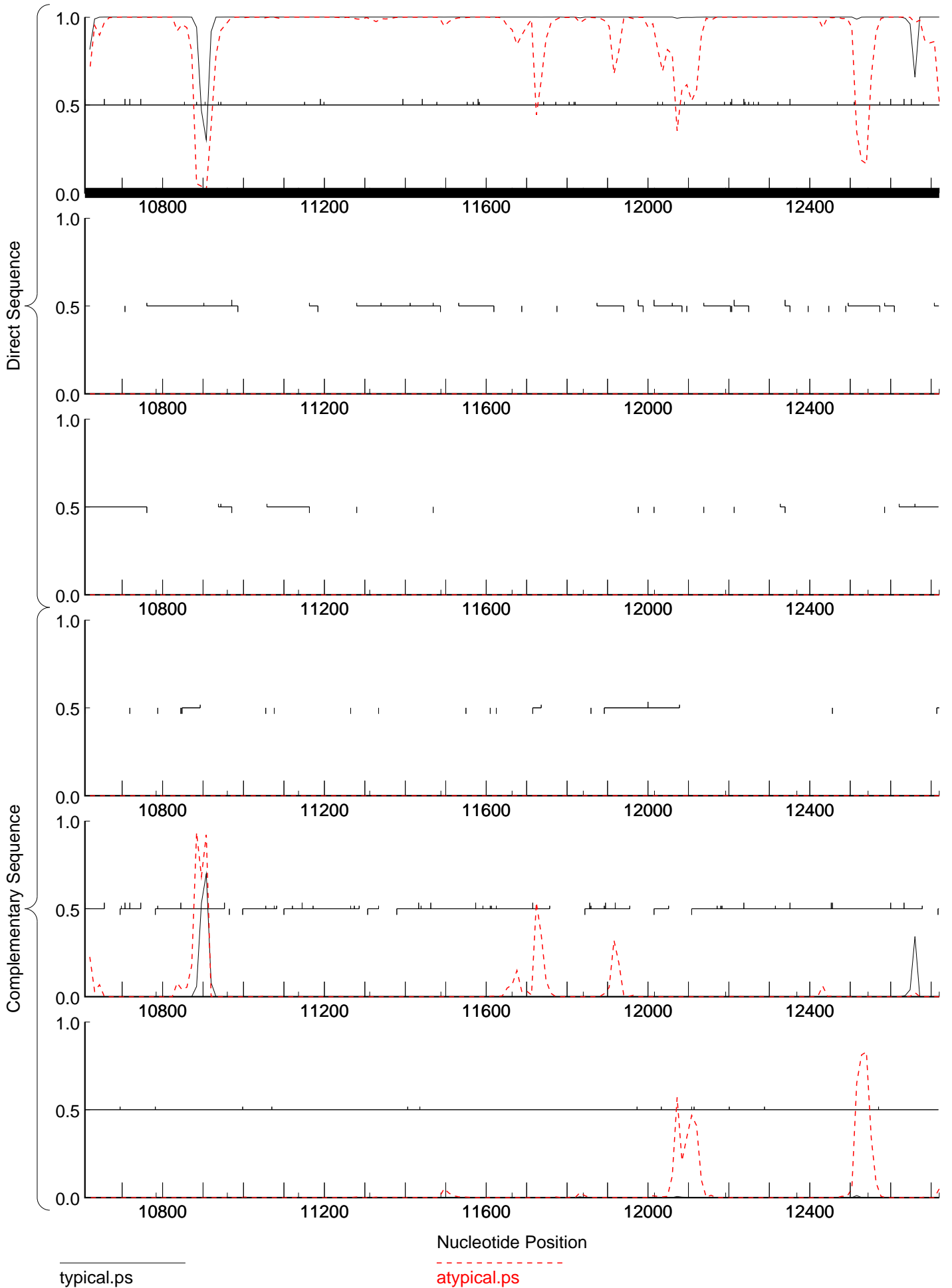


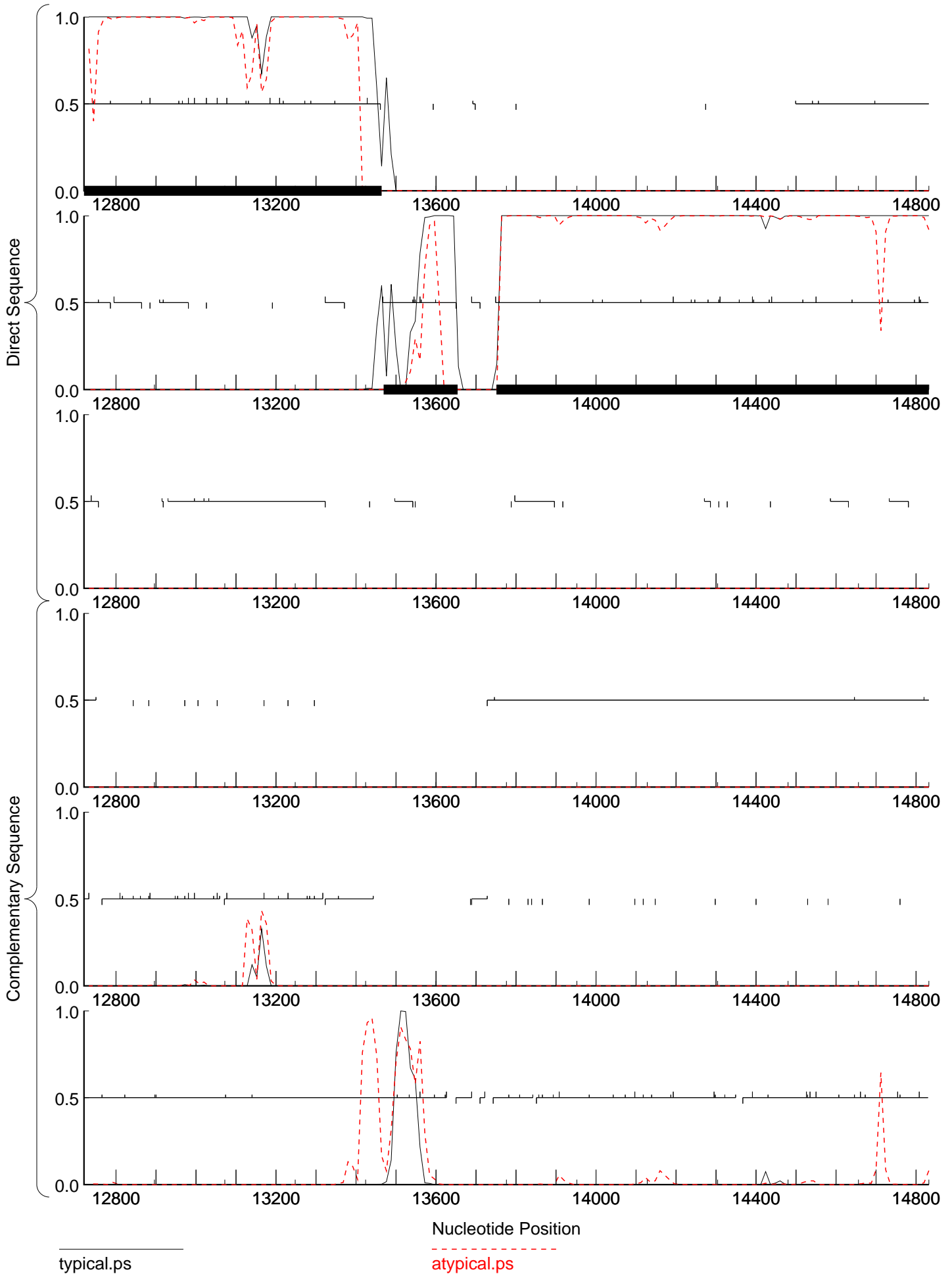


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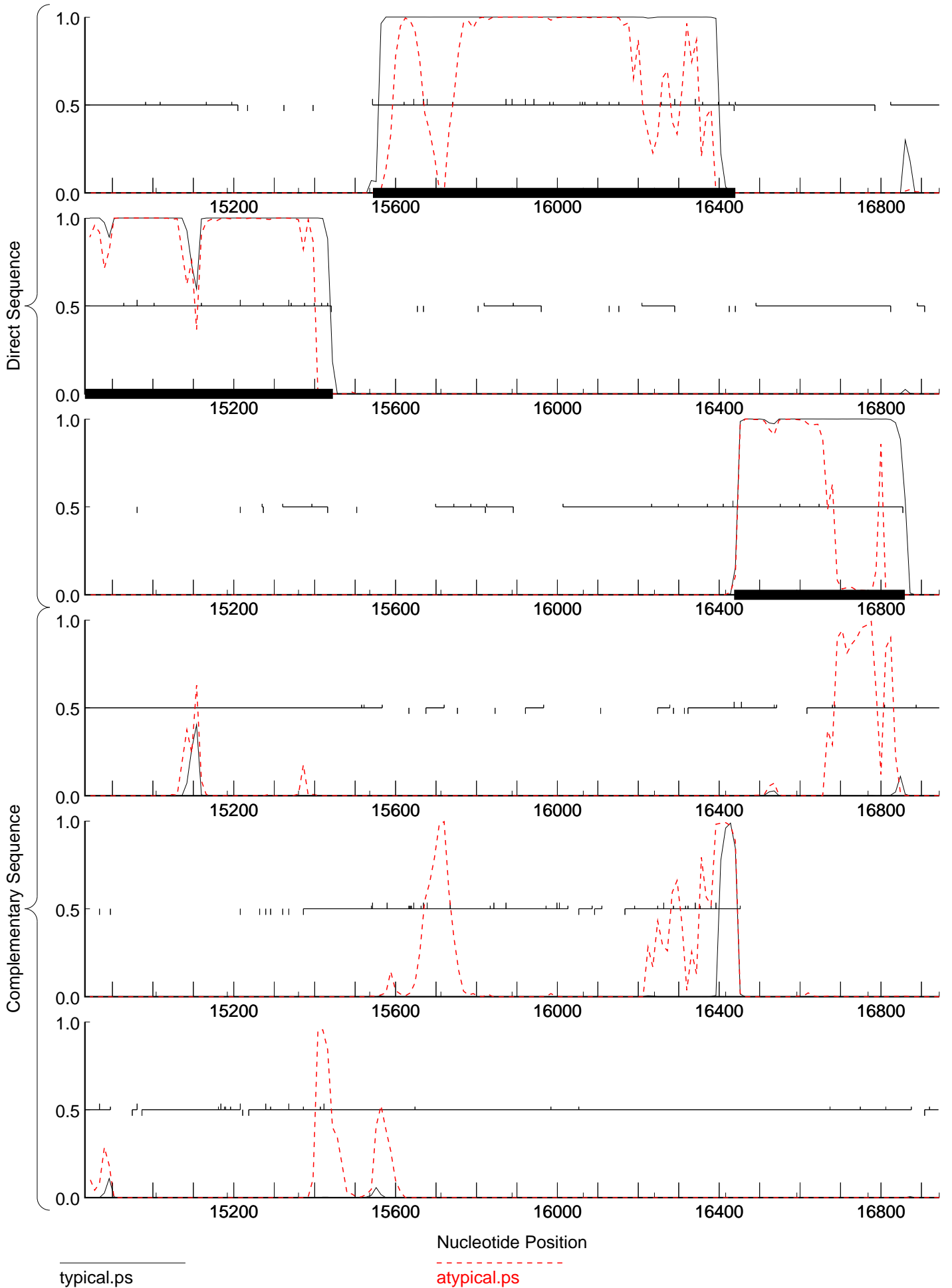
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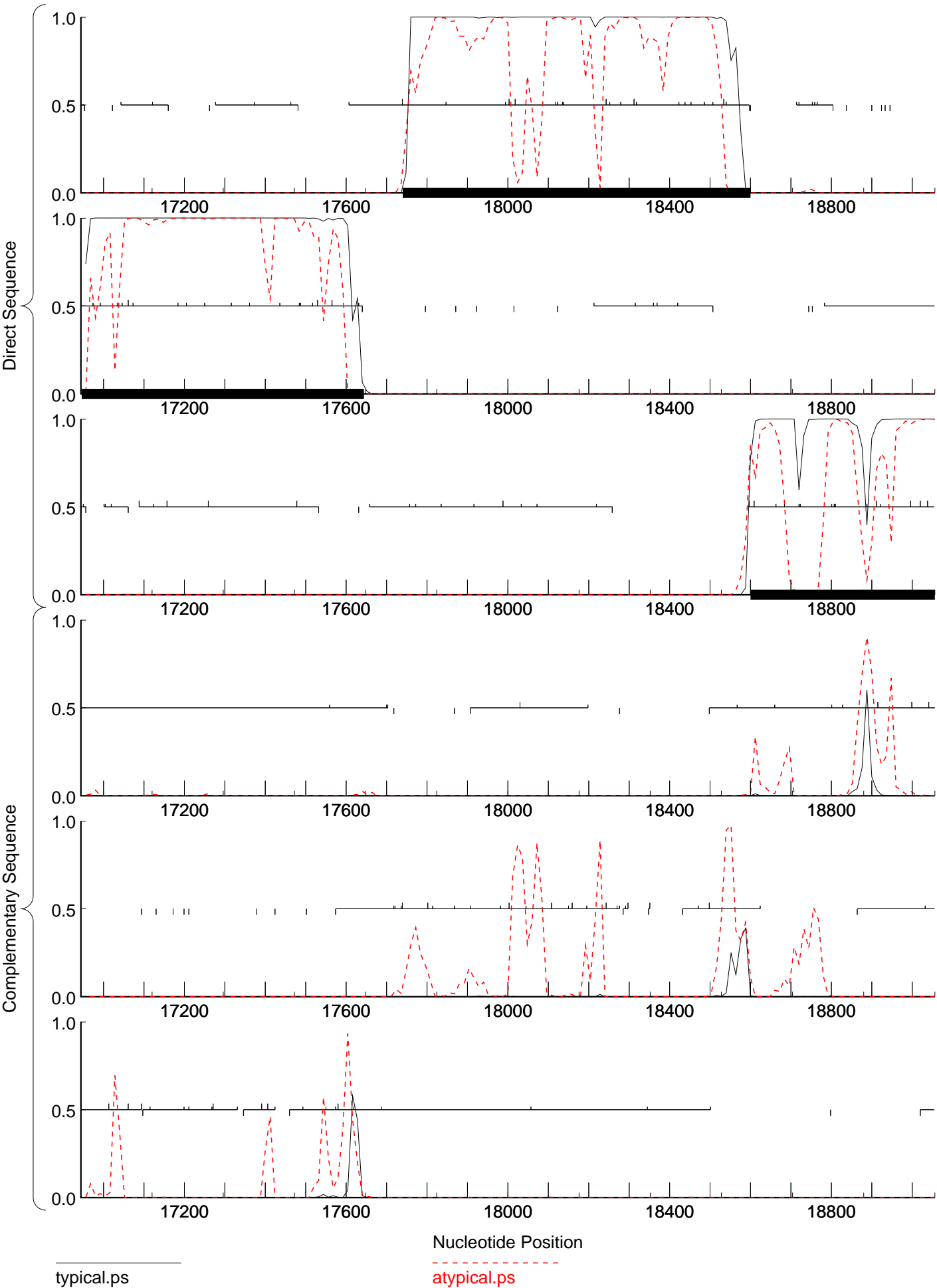


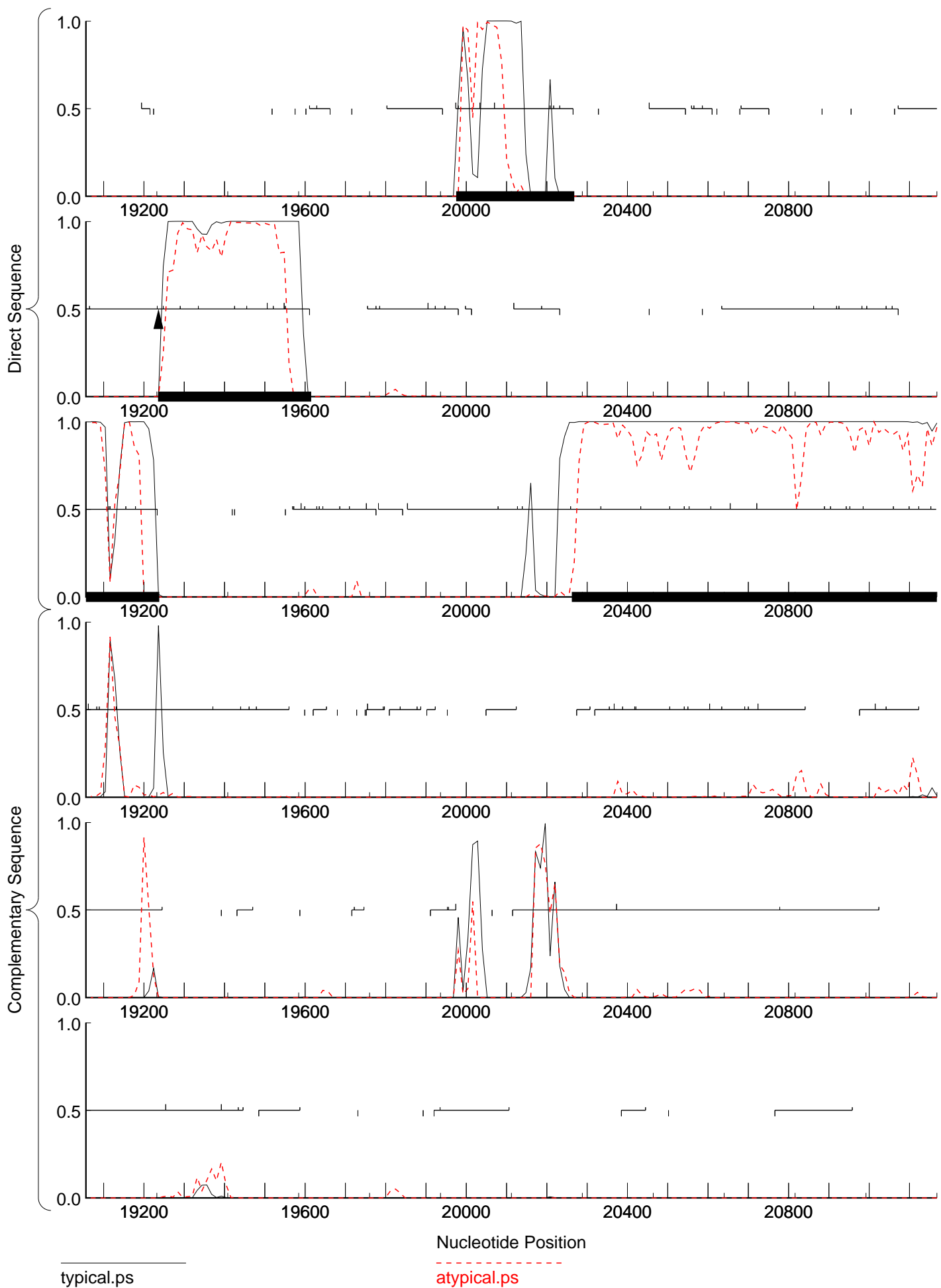


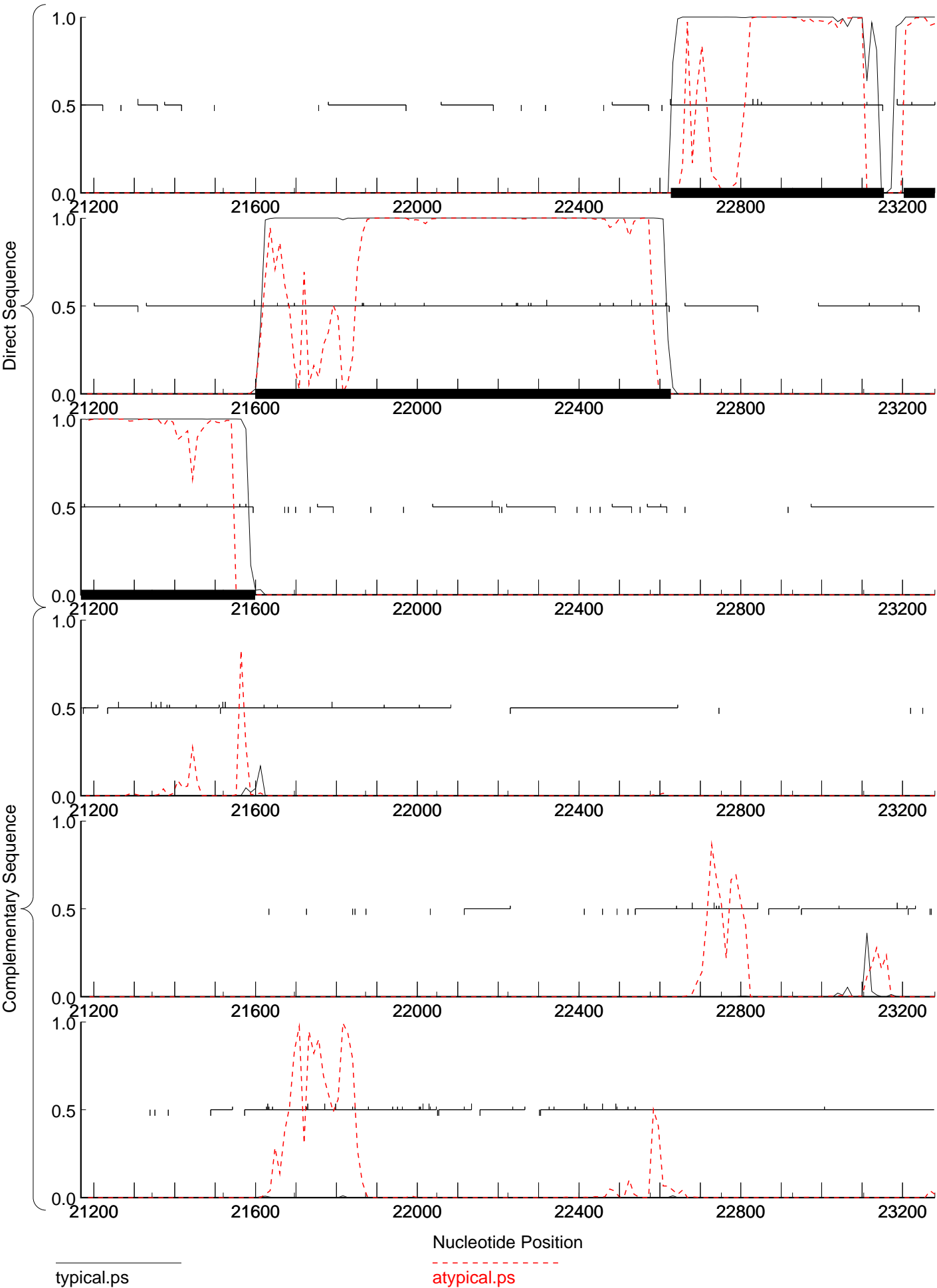






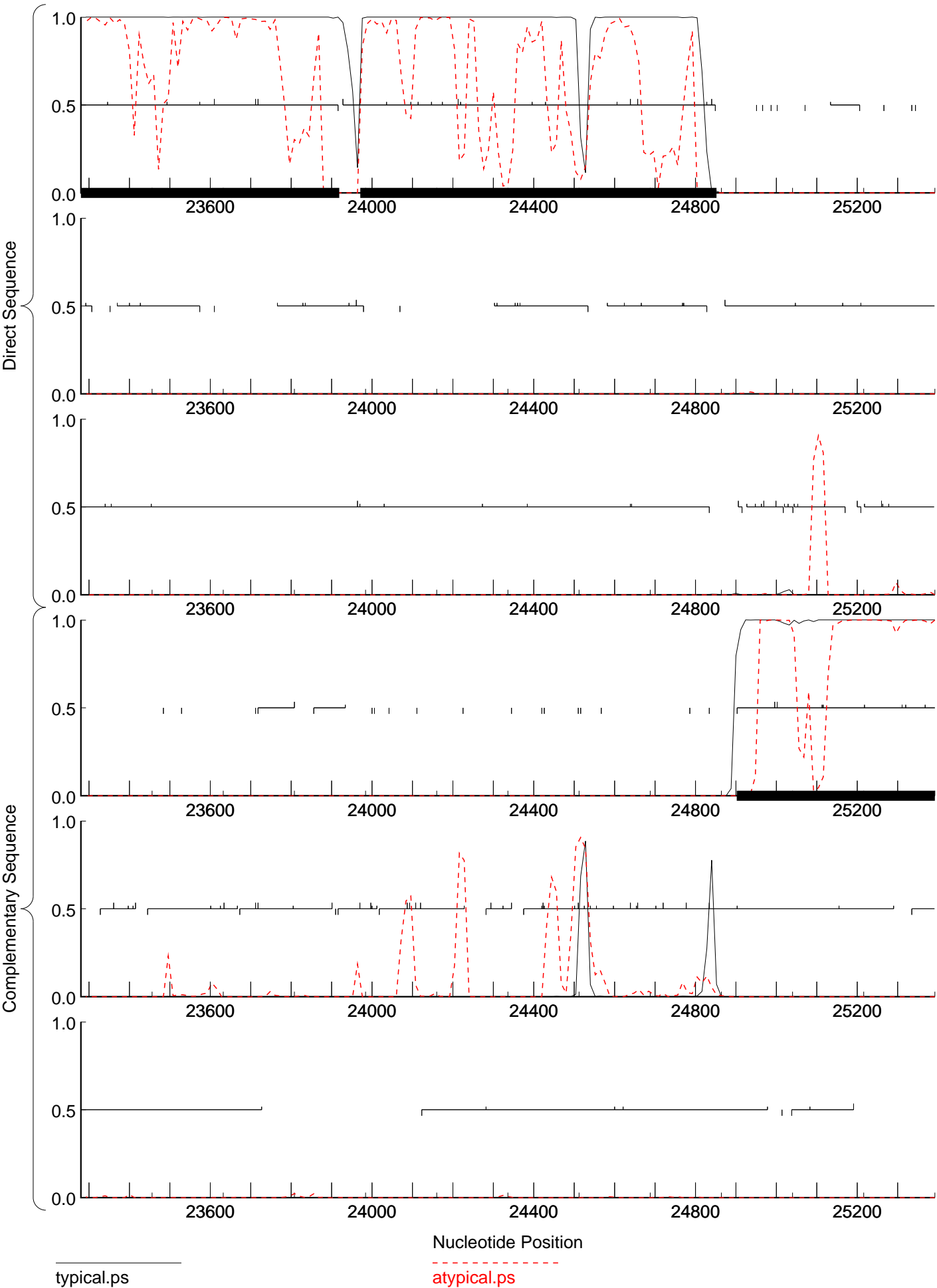


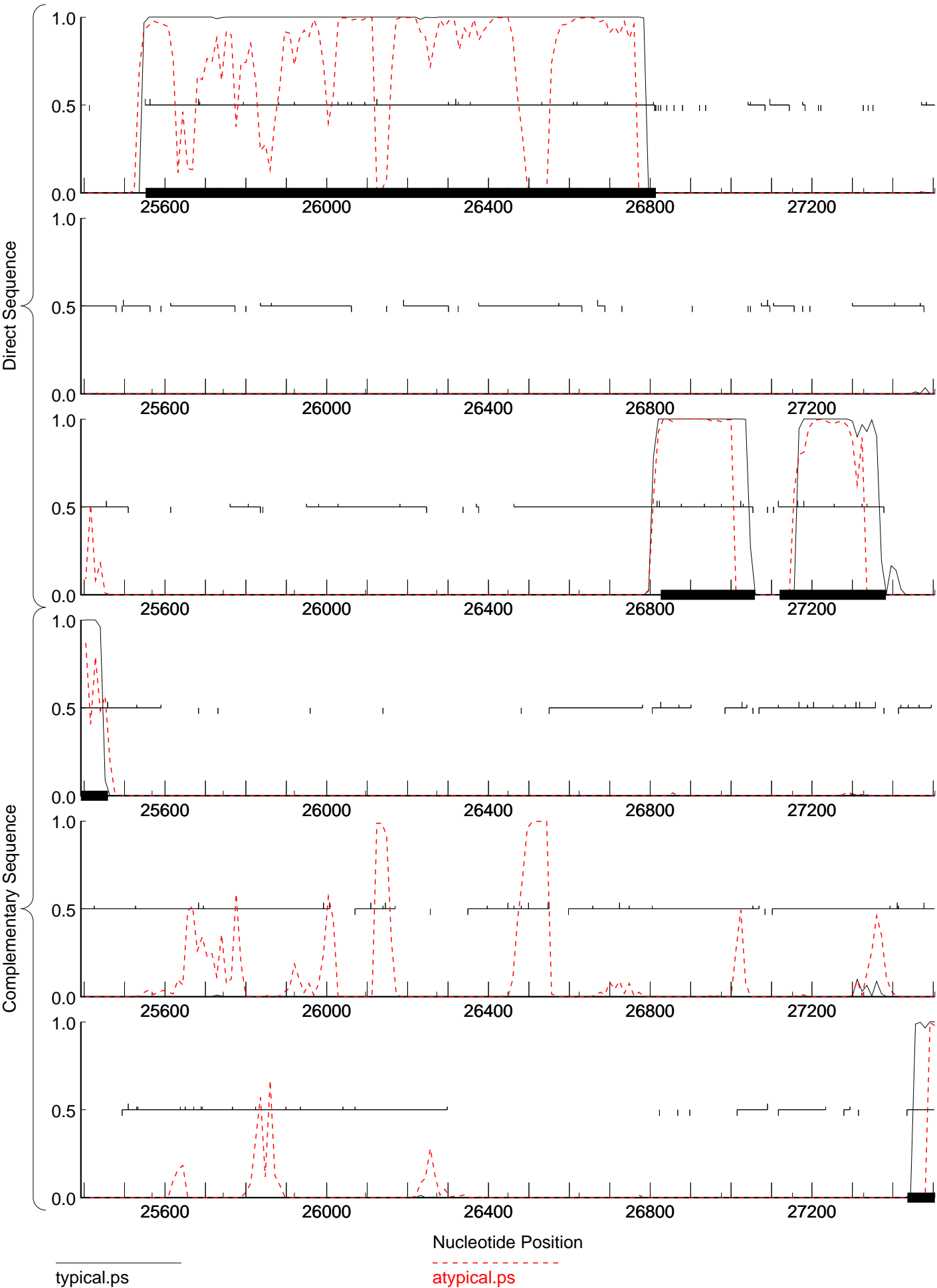




typical.ps

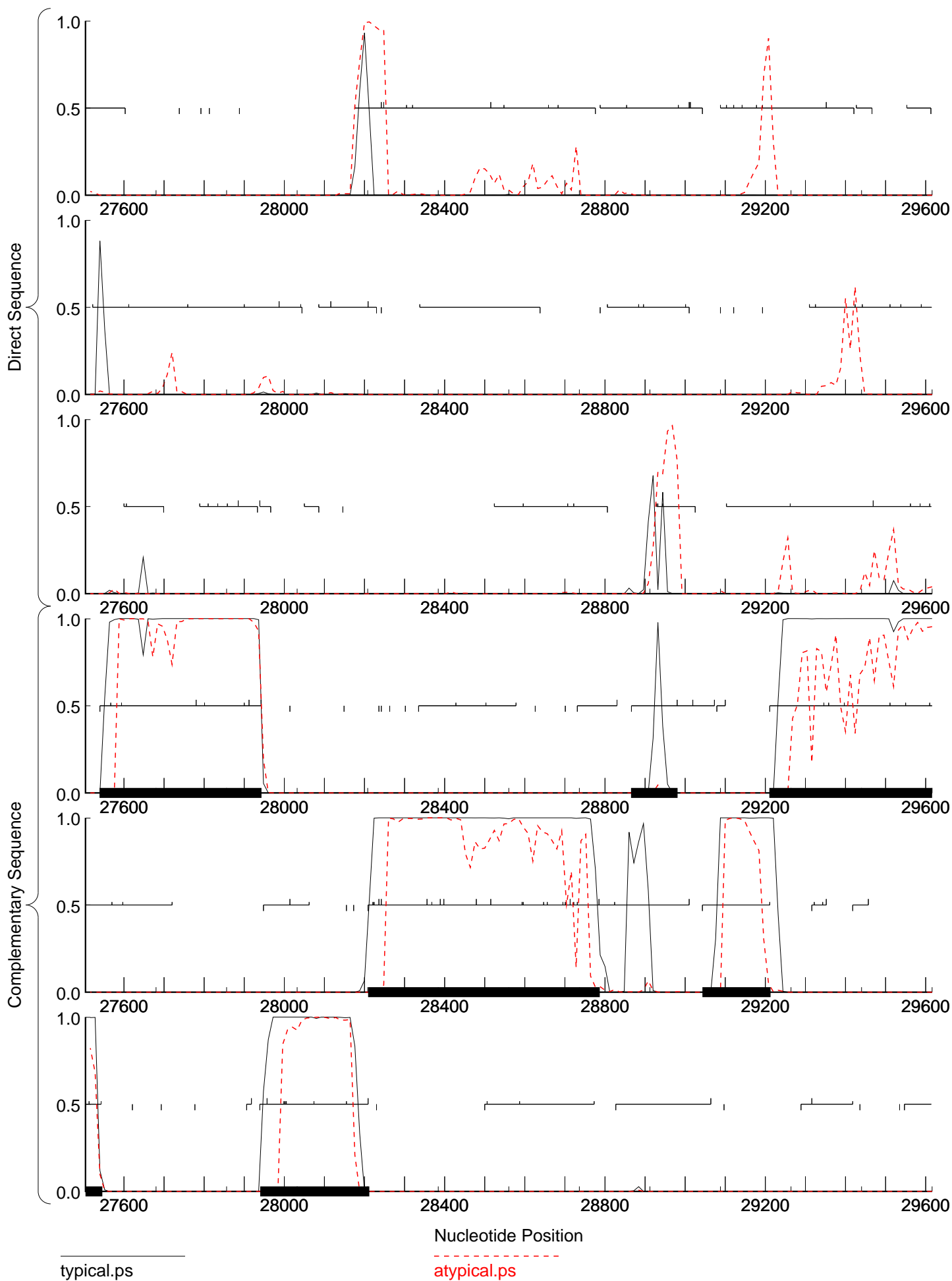
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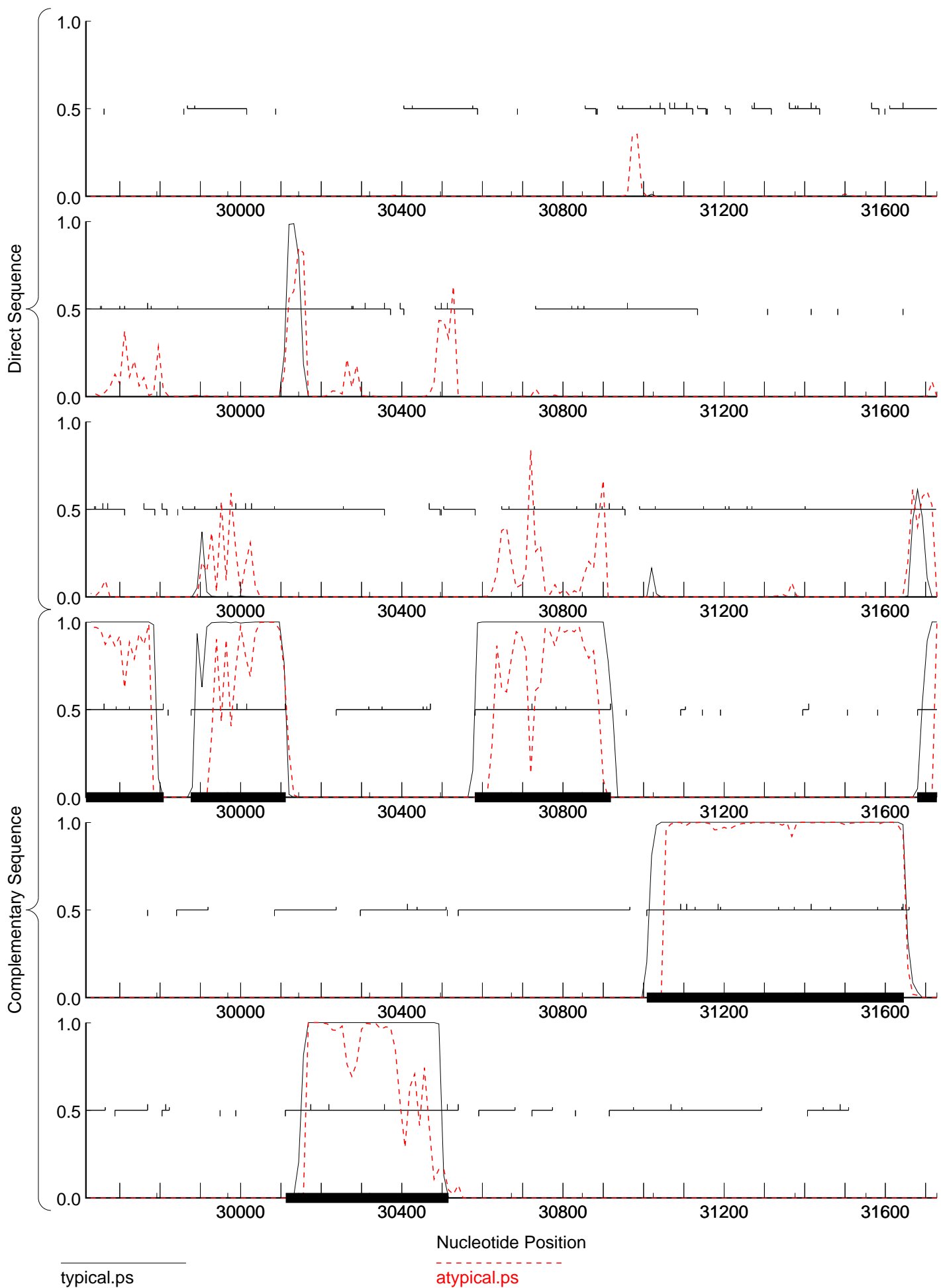




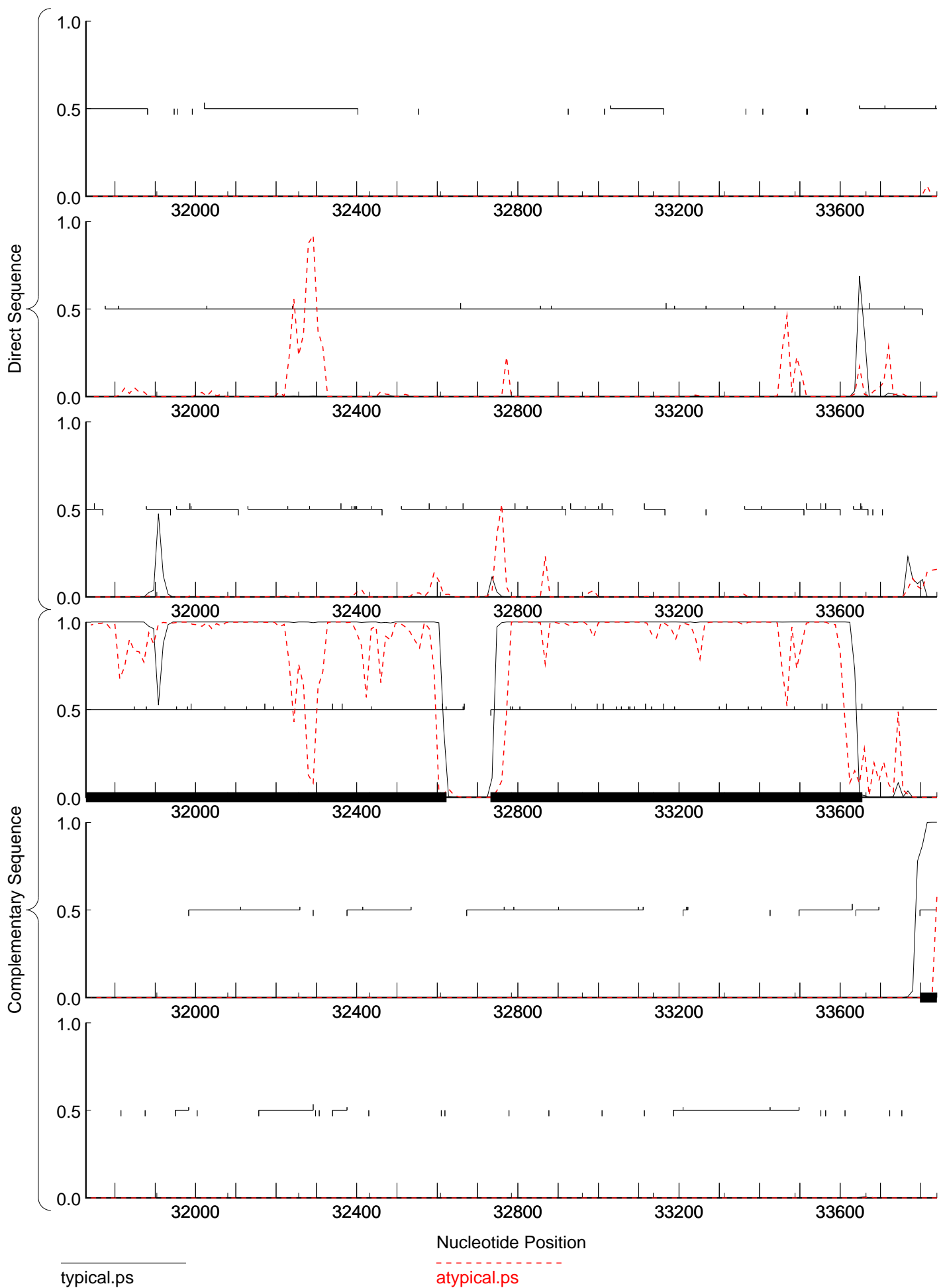
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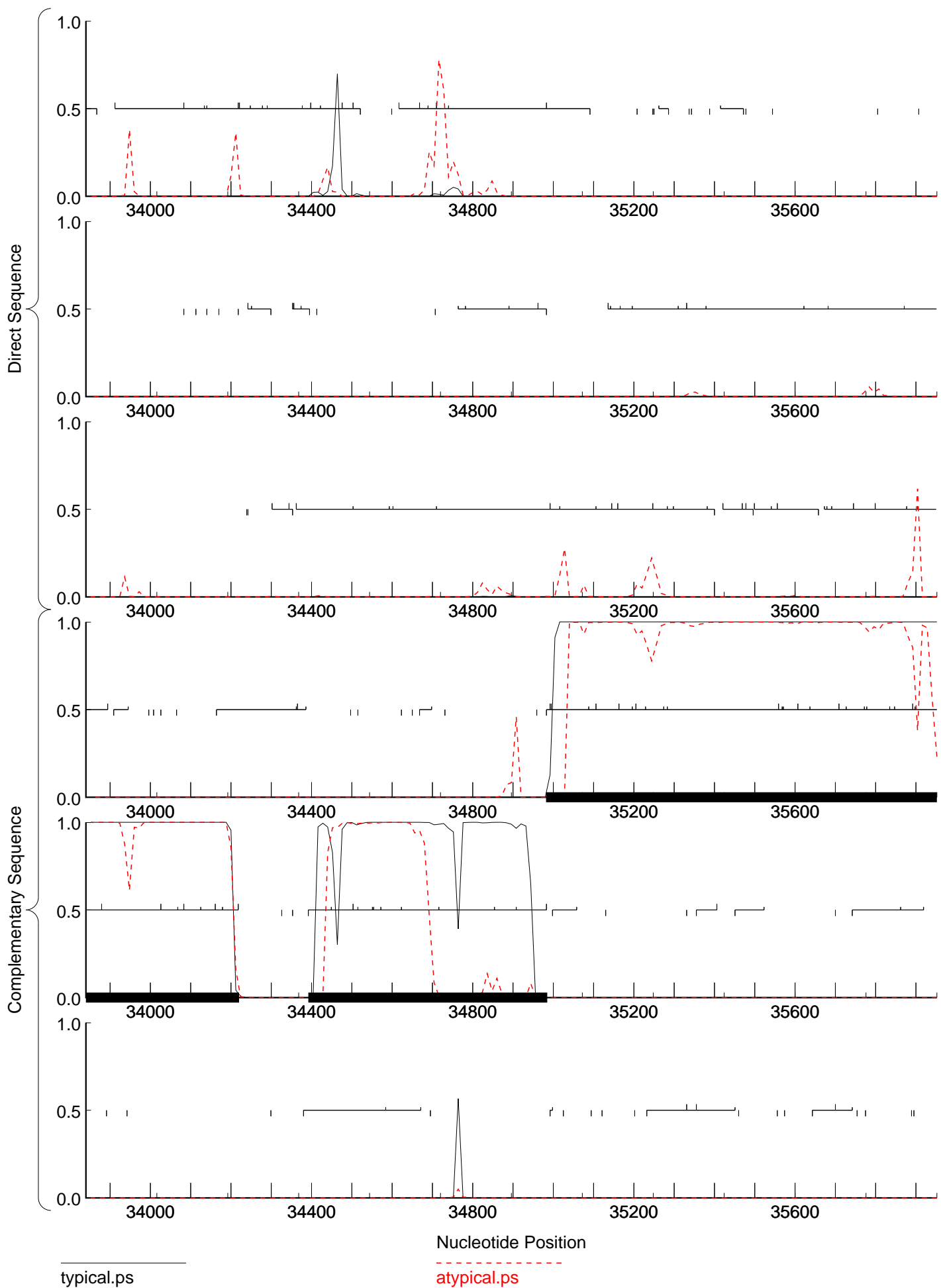
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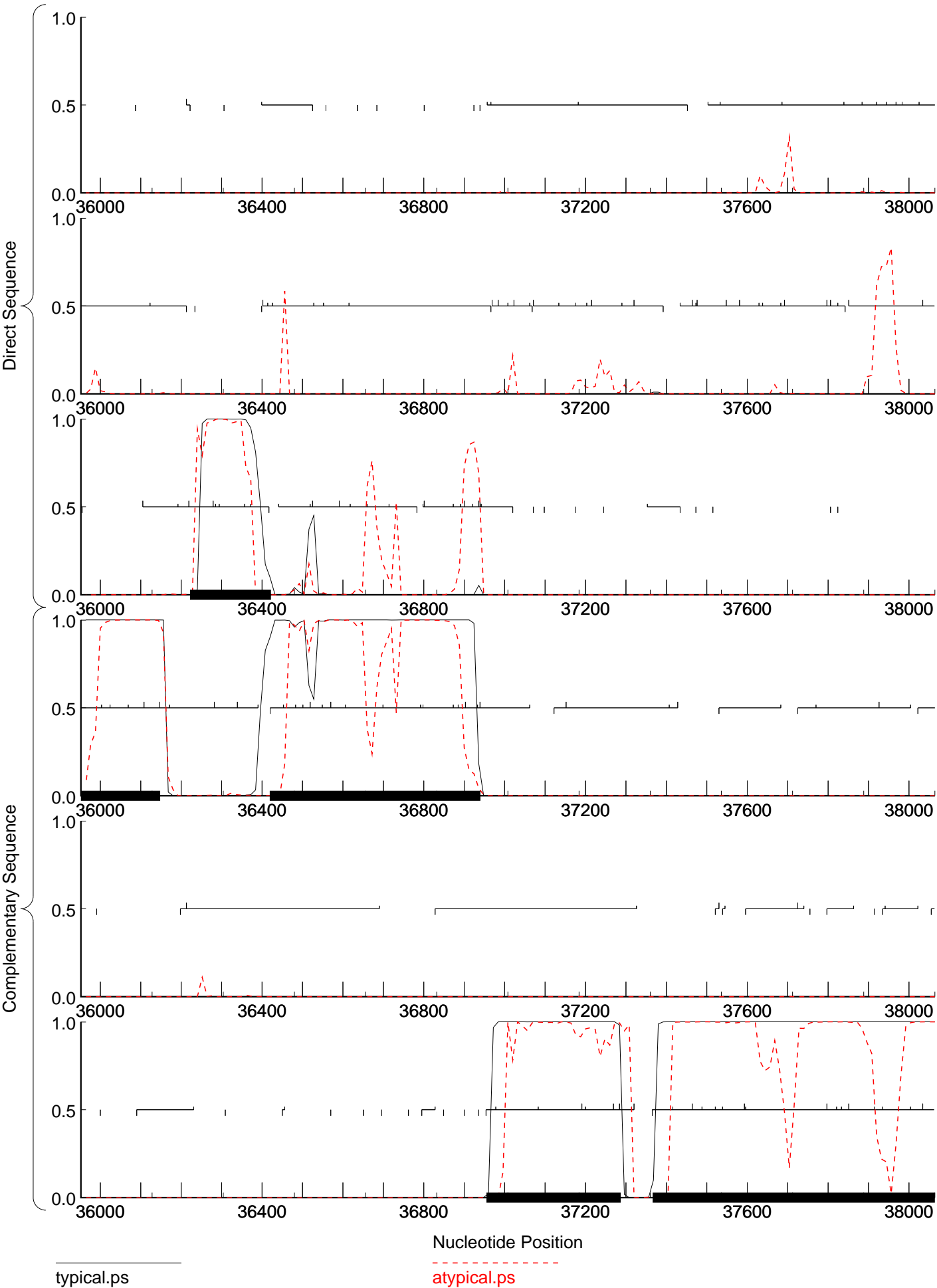


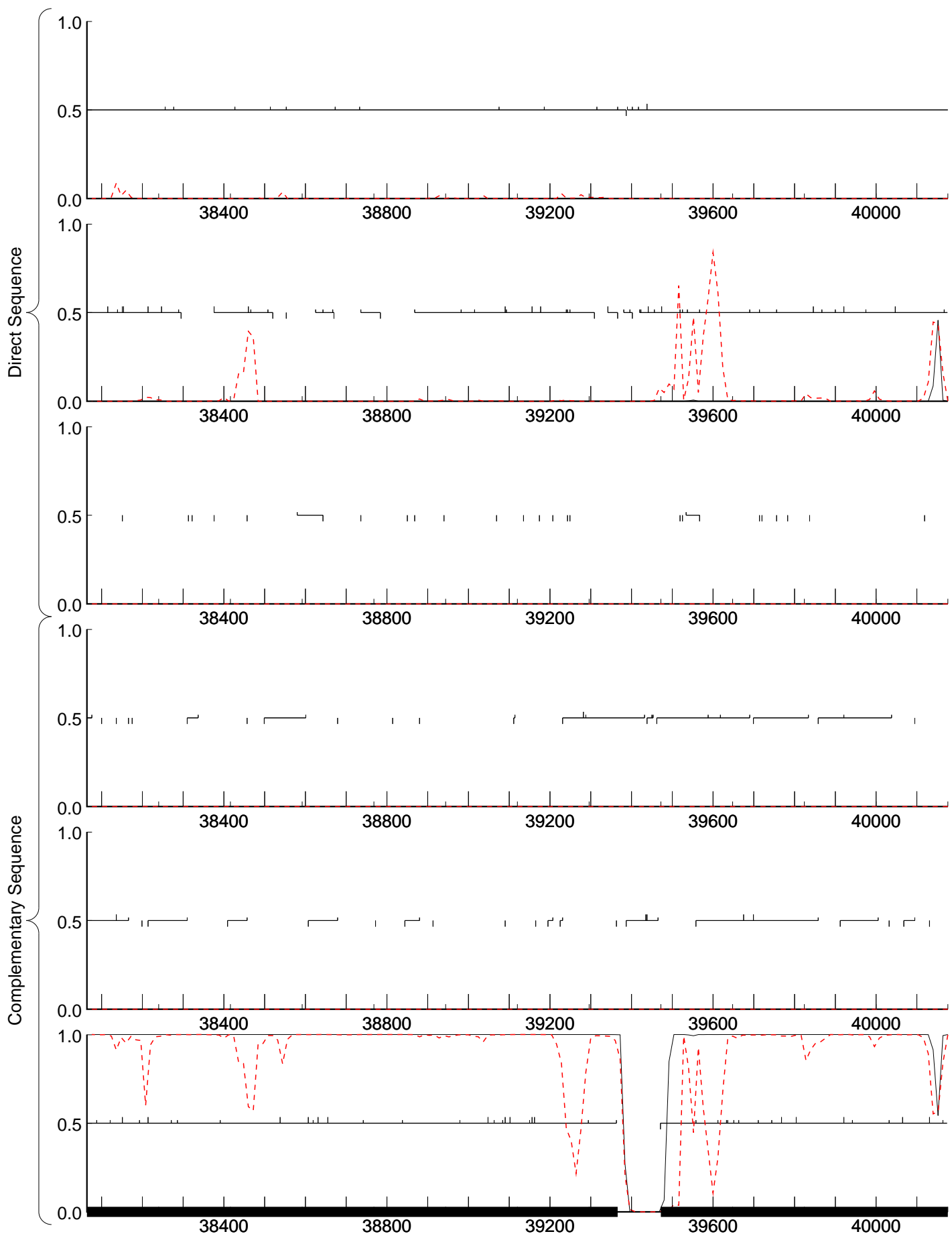








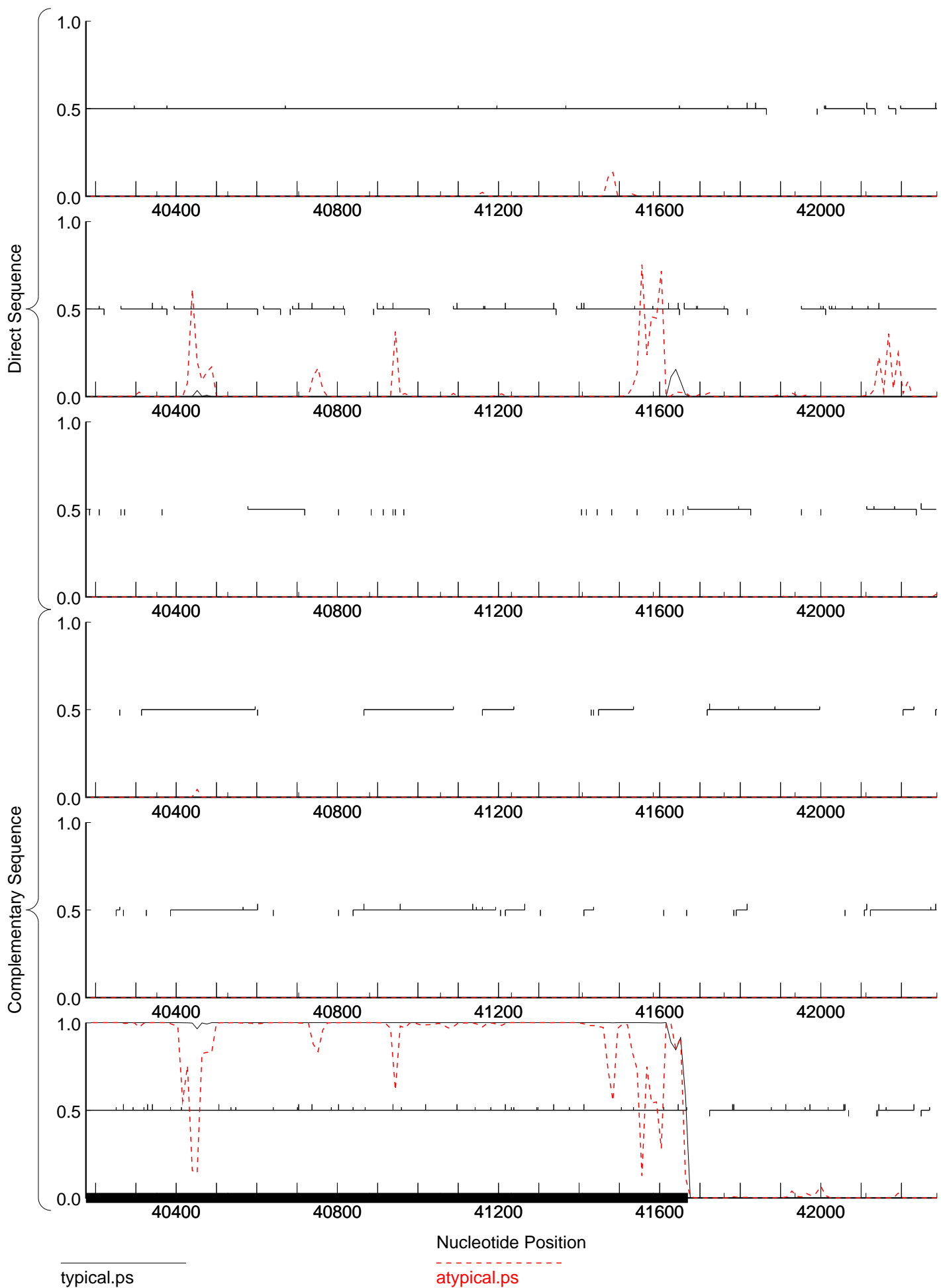




typical.ps

Nucleotide Position

atypical.ps



typical.ps

atypical.ps

