

# GeneMark

Version 2.5p (09.08.06)

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## PROGRAM INFORMATION

Sequence : Mycobacterium phage Pippy complete sequence, 56663 bp including 10-base 3' overhang (CGGAAG  
Analysis Date : 8/19/17 at 5:17:55  
Pages : 28  
Sequence Length : 56663 bp  
GC Content : 61.50%

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,

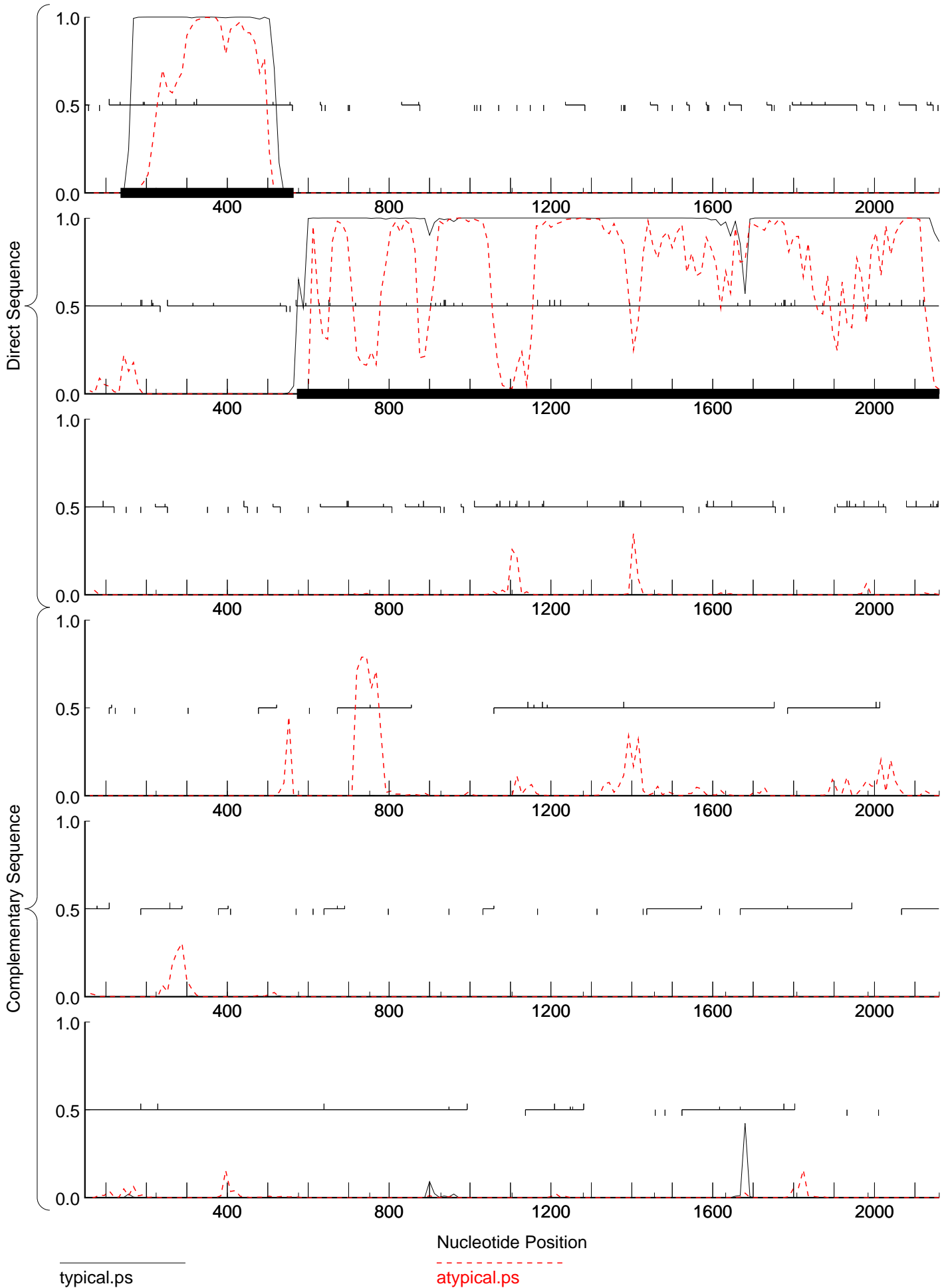
## Matrix notes & comments

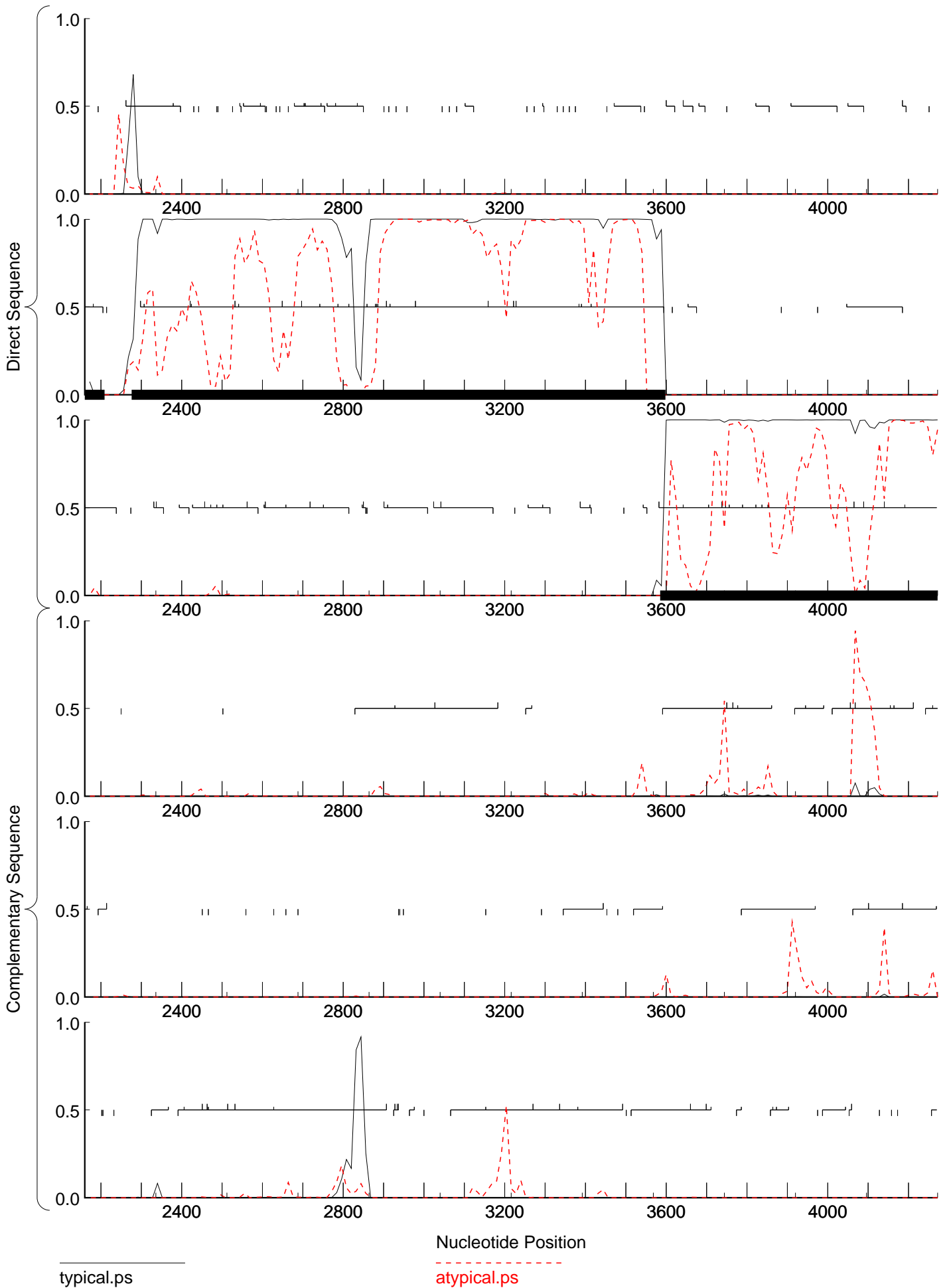
Training set derived by GeneMarkS, 4.30 October 2014  
Sat Aug 19 05:17:55 2017

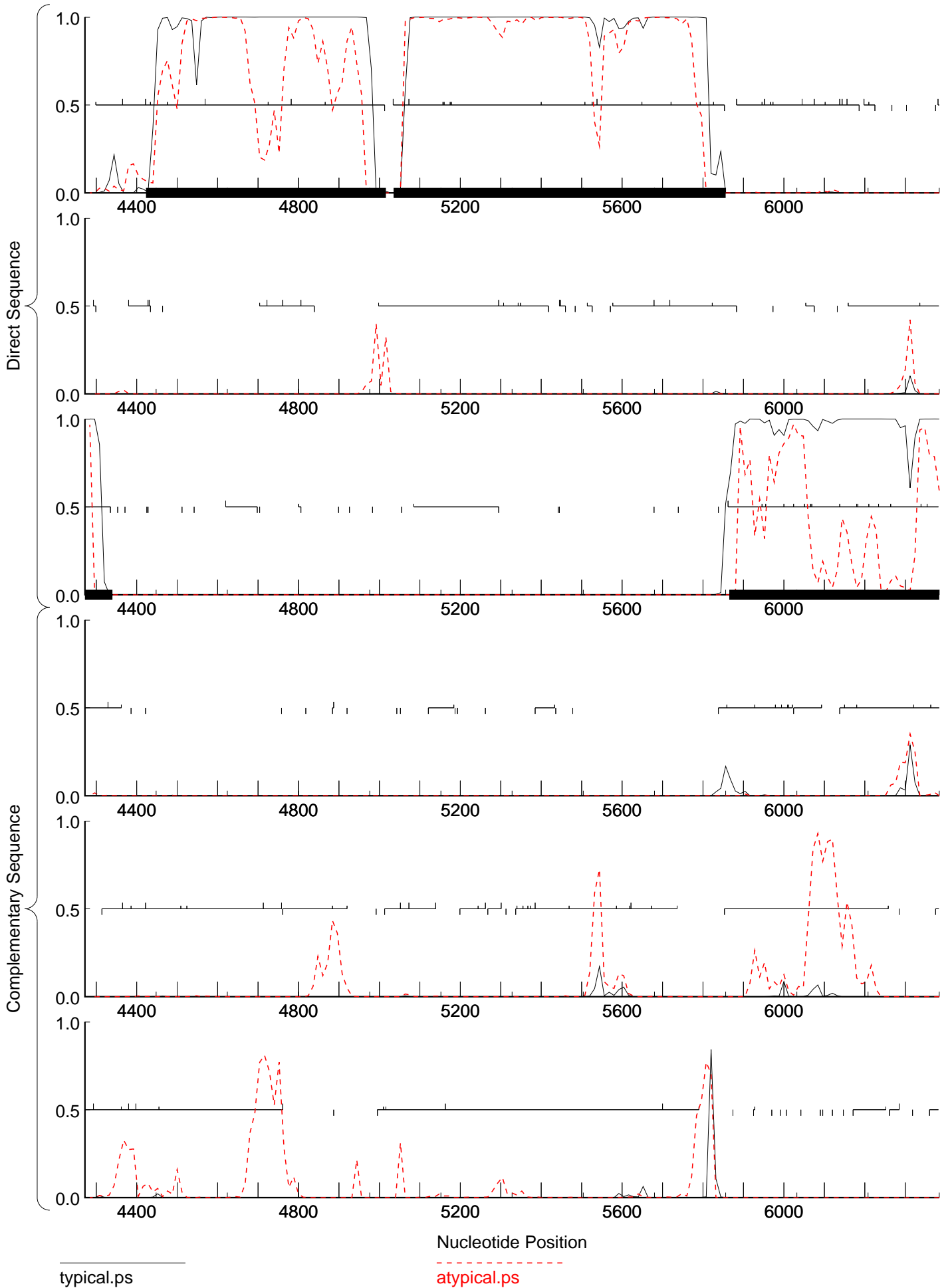
## MATRIX INFORMATION

Matrix : -  
Author : -  
Order : 4

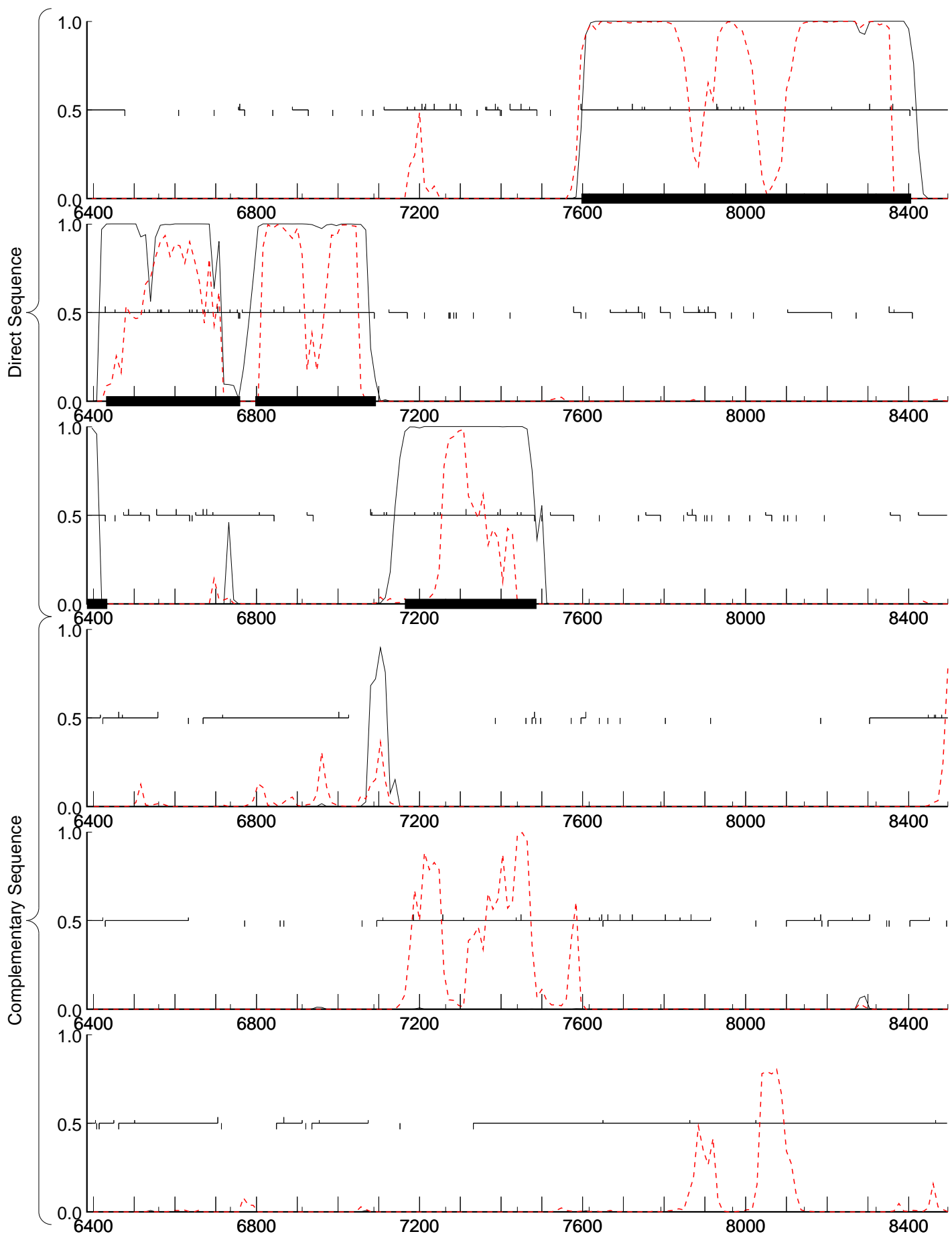
Send questions / comments to:  
Dr. M. Borodovsky  
Georgia Institute of Technology  
School of Biology  
Atlanta, GA 30332-0230





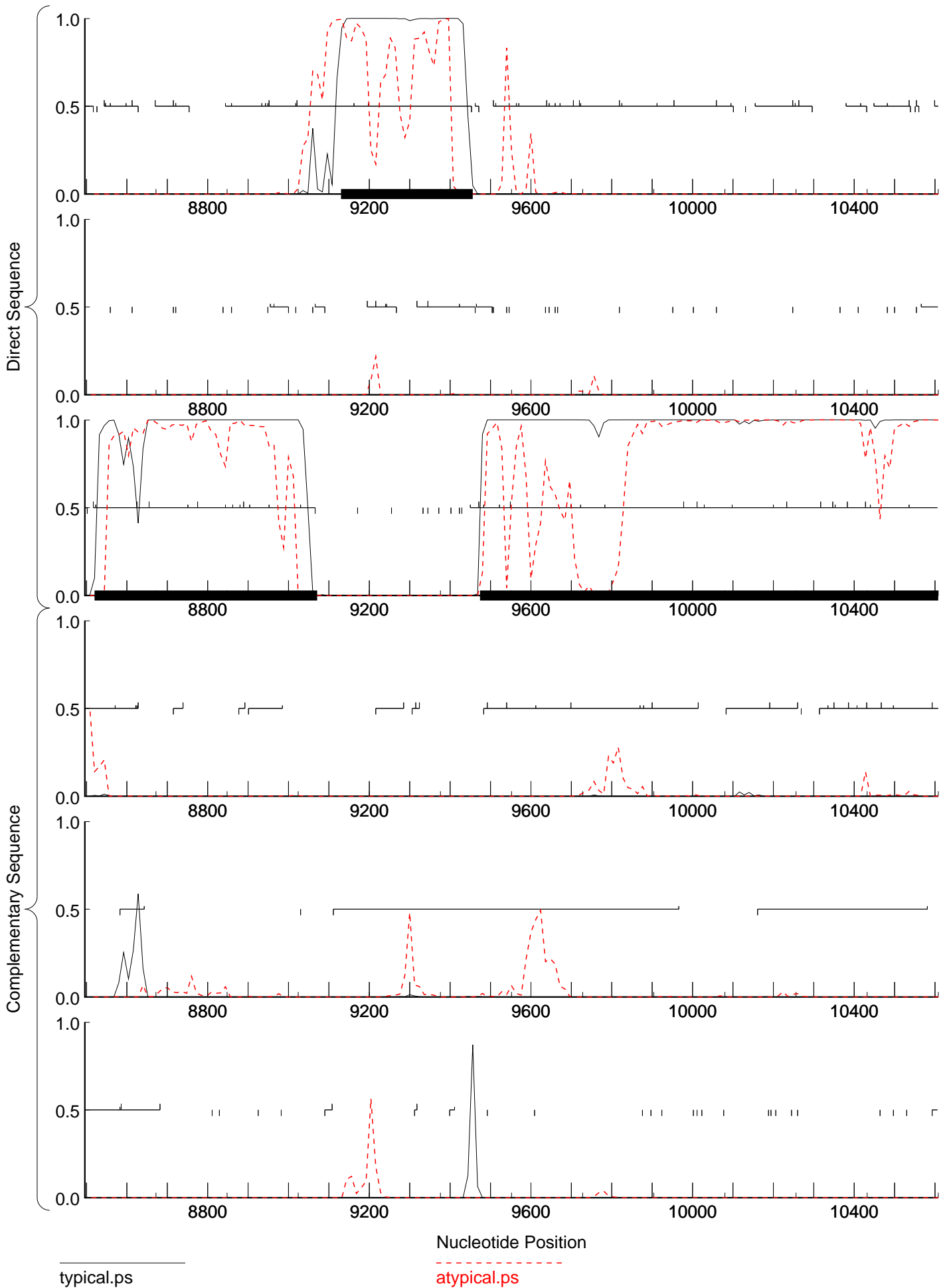


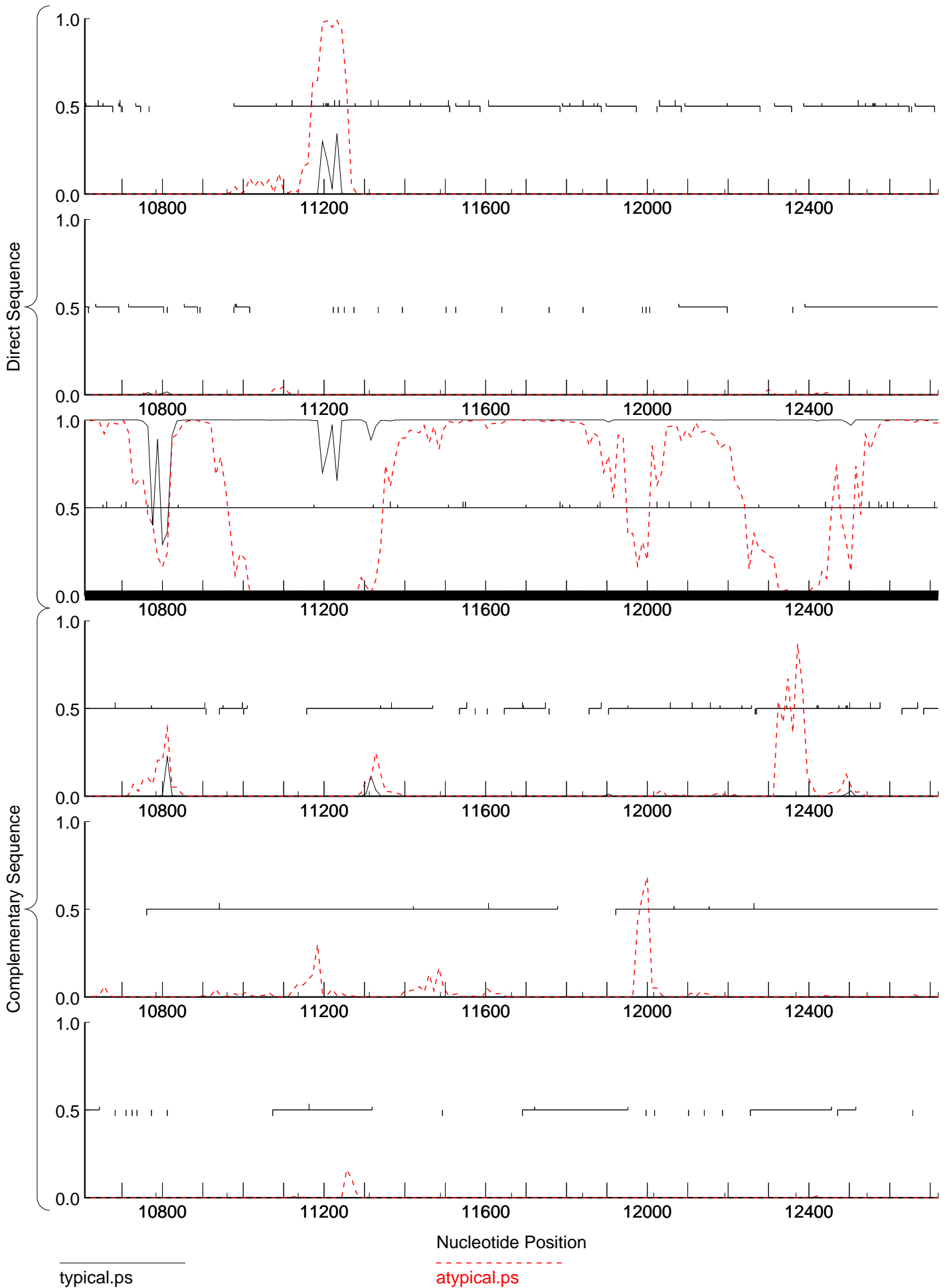
GeneMark.hmm prediction

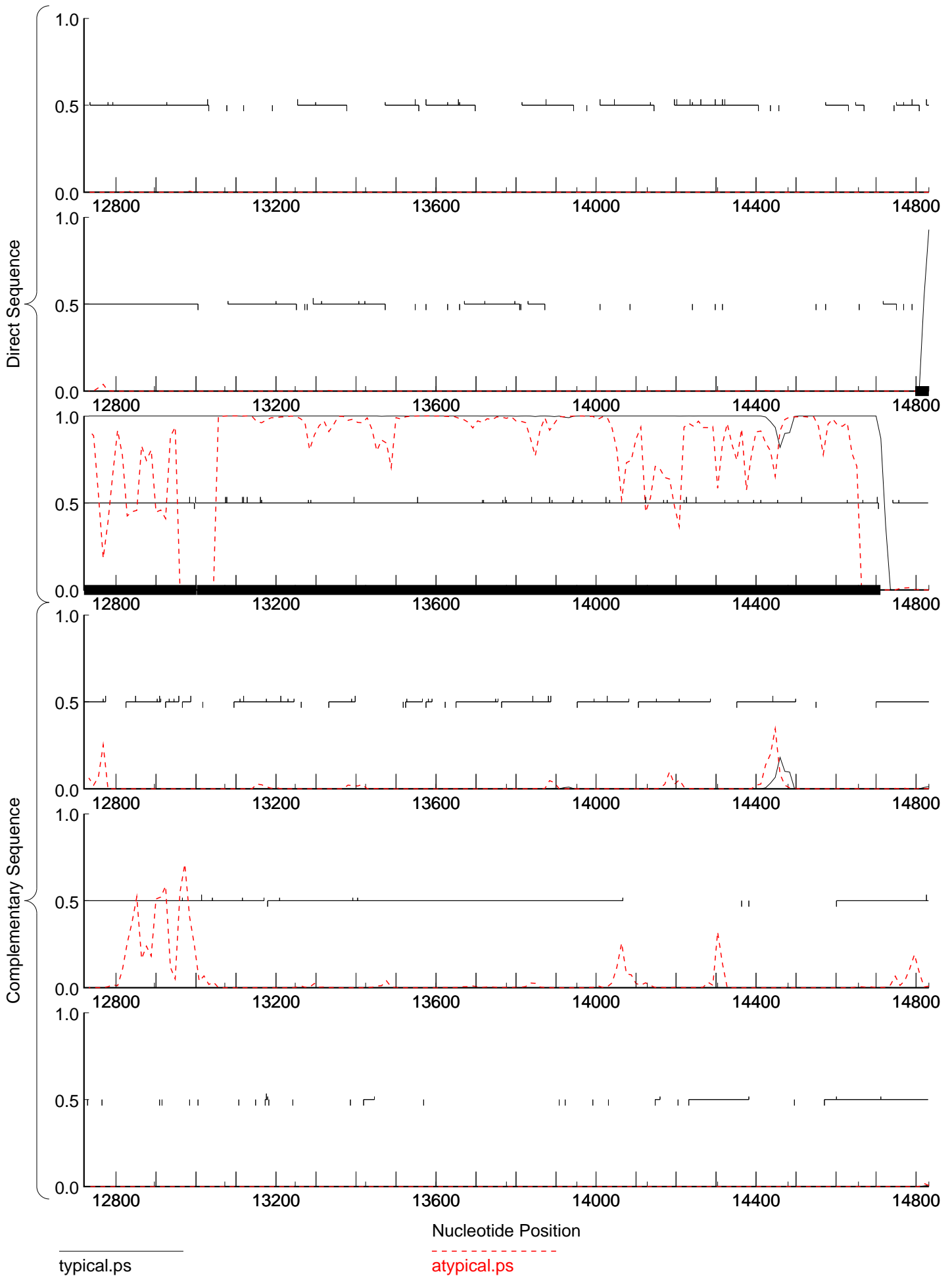


typical.ps

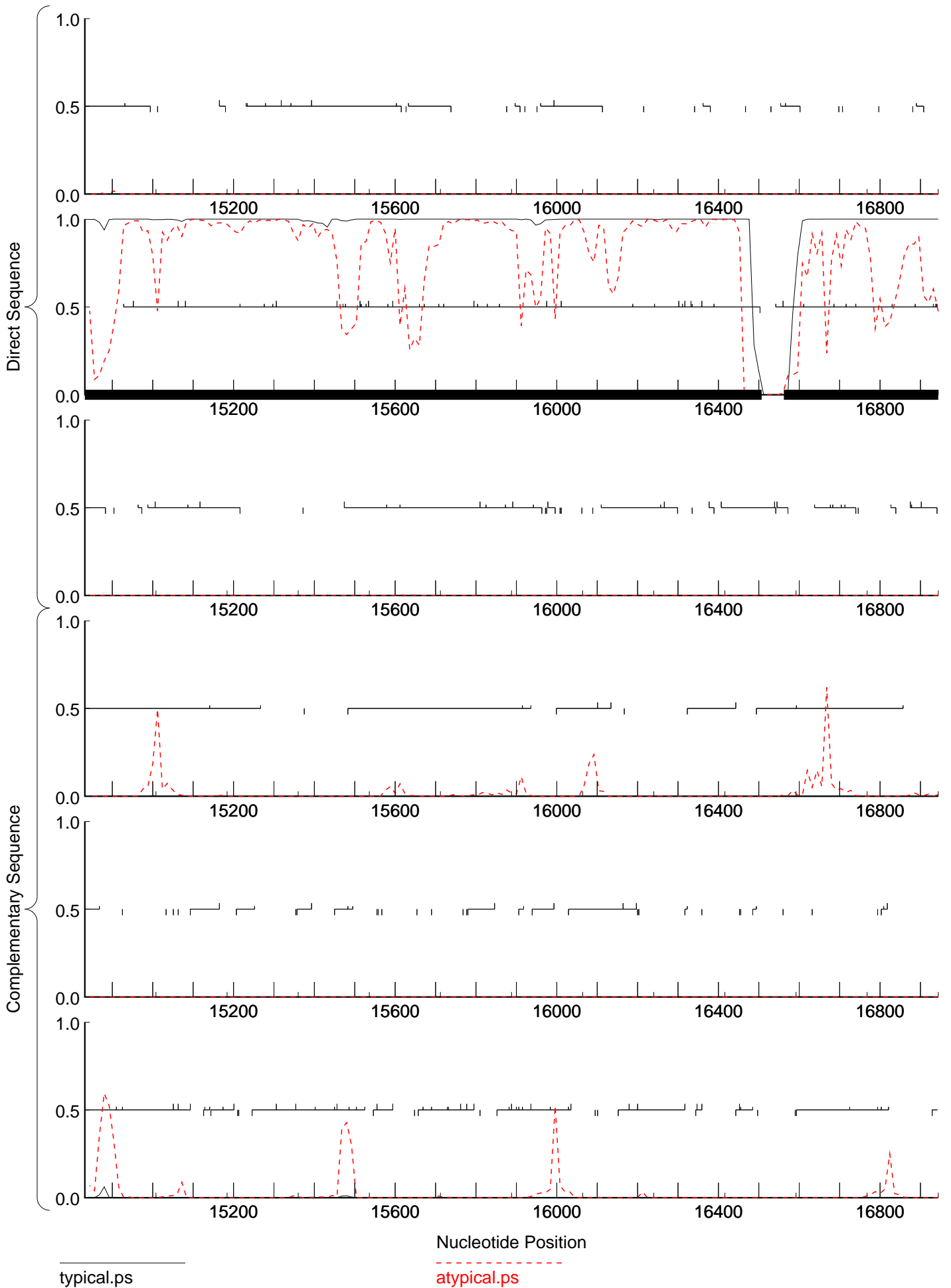
atypical.ps

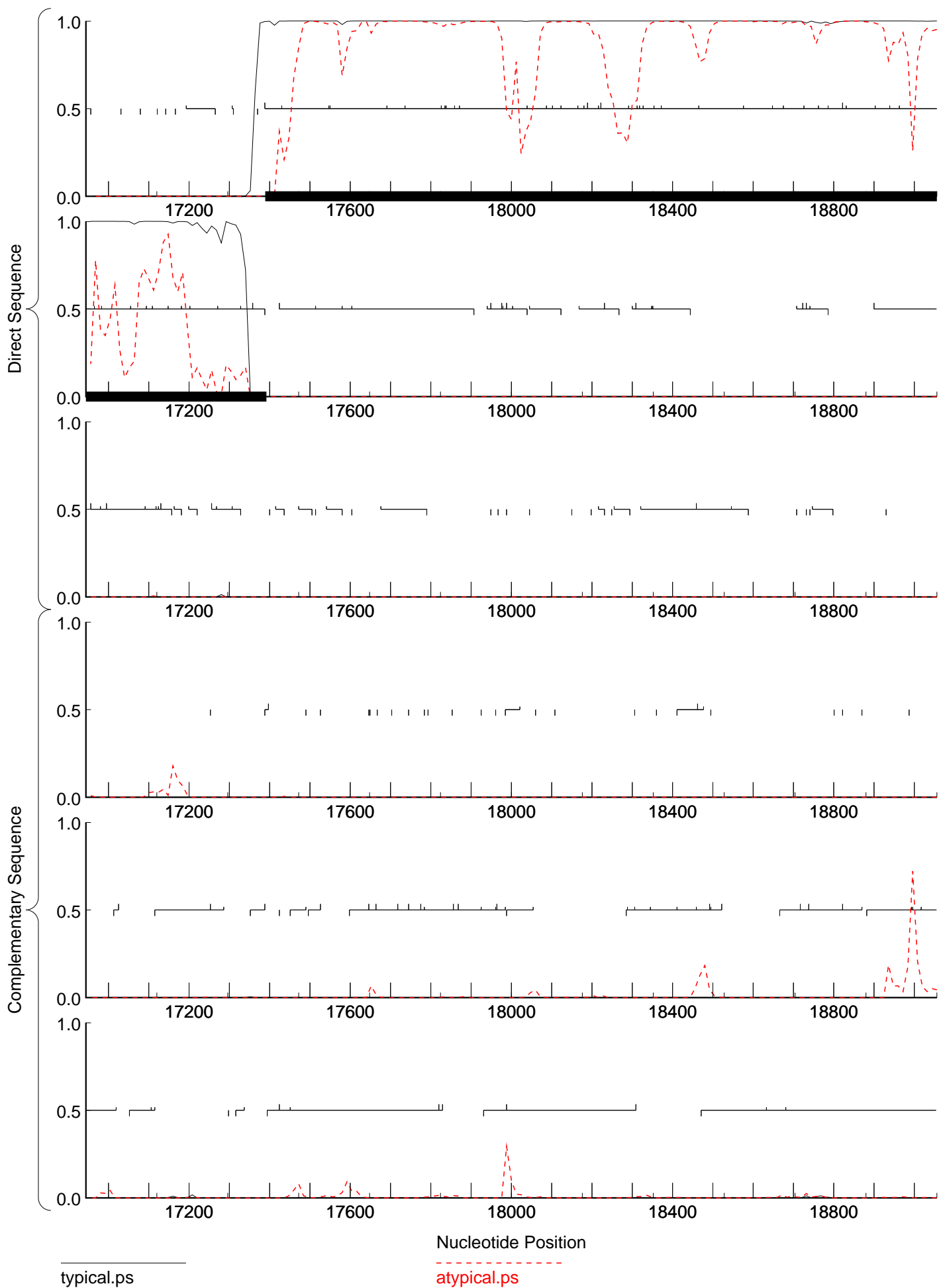






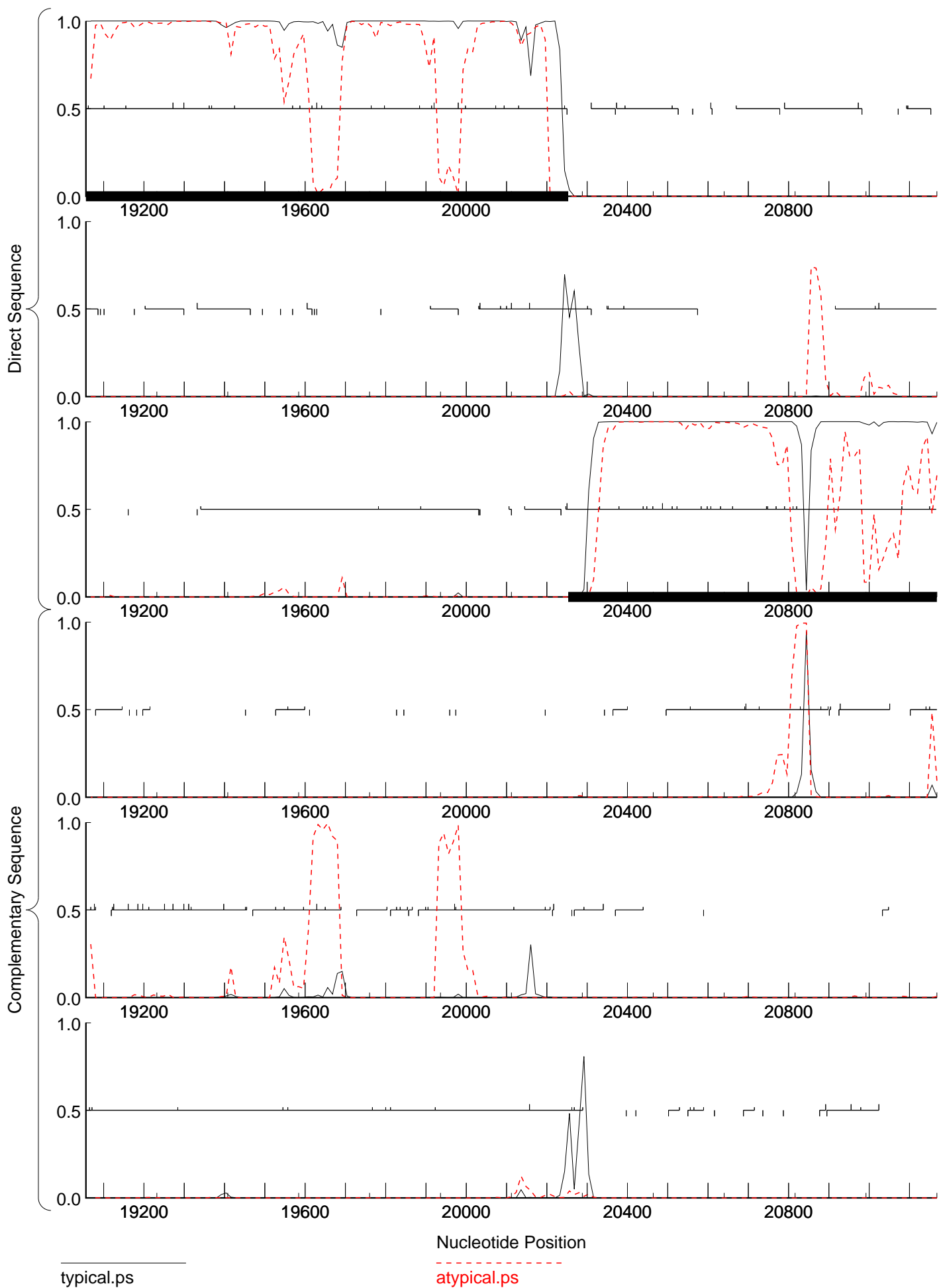






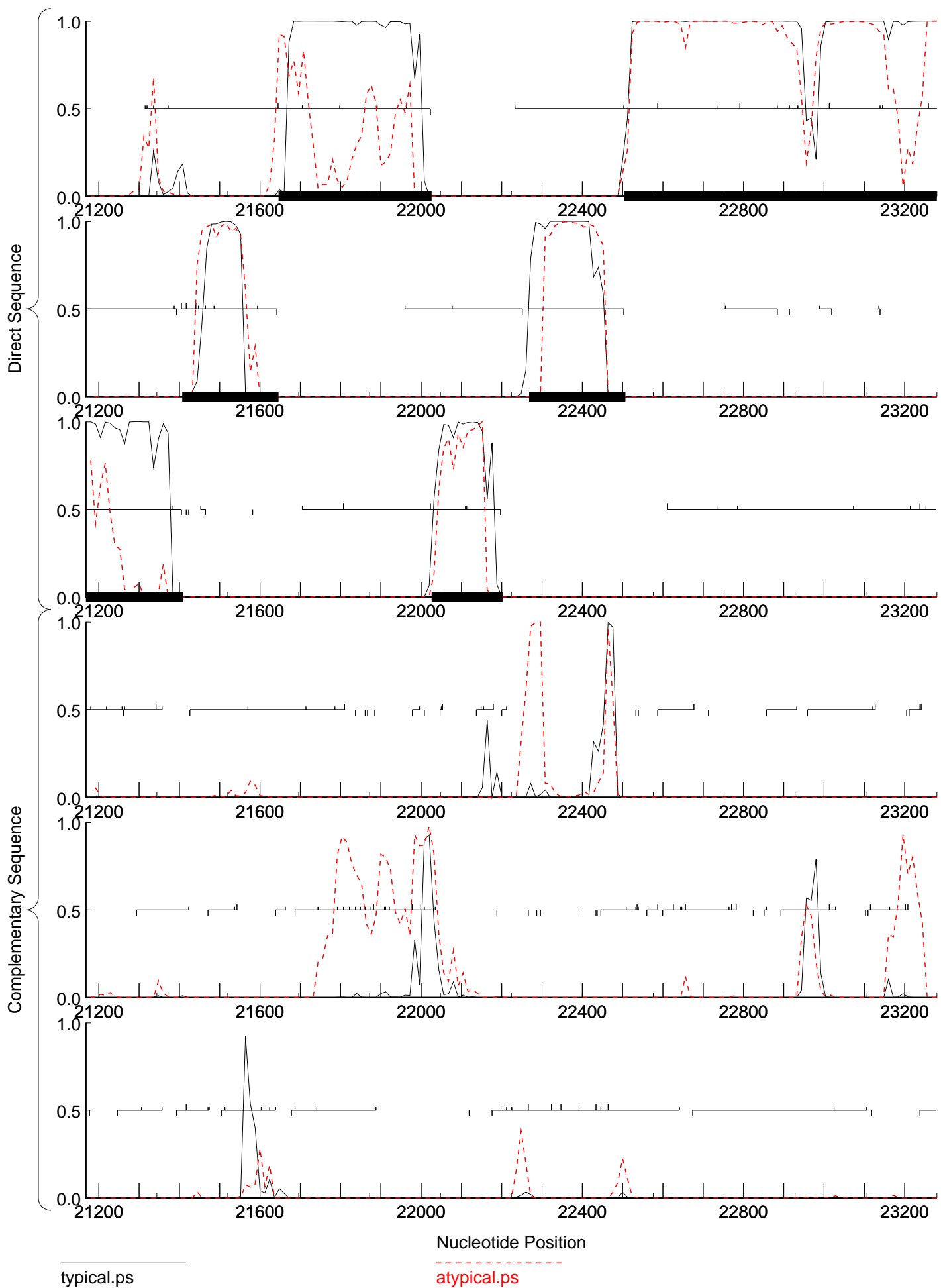
typical.ps

atypical.ps



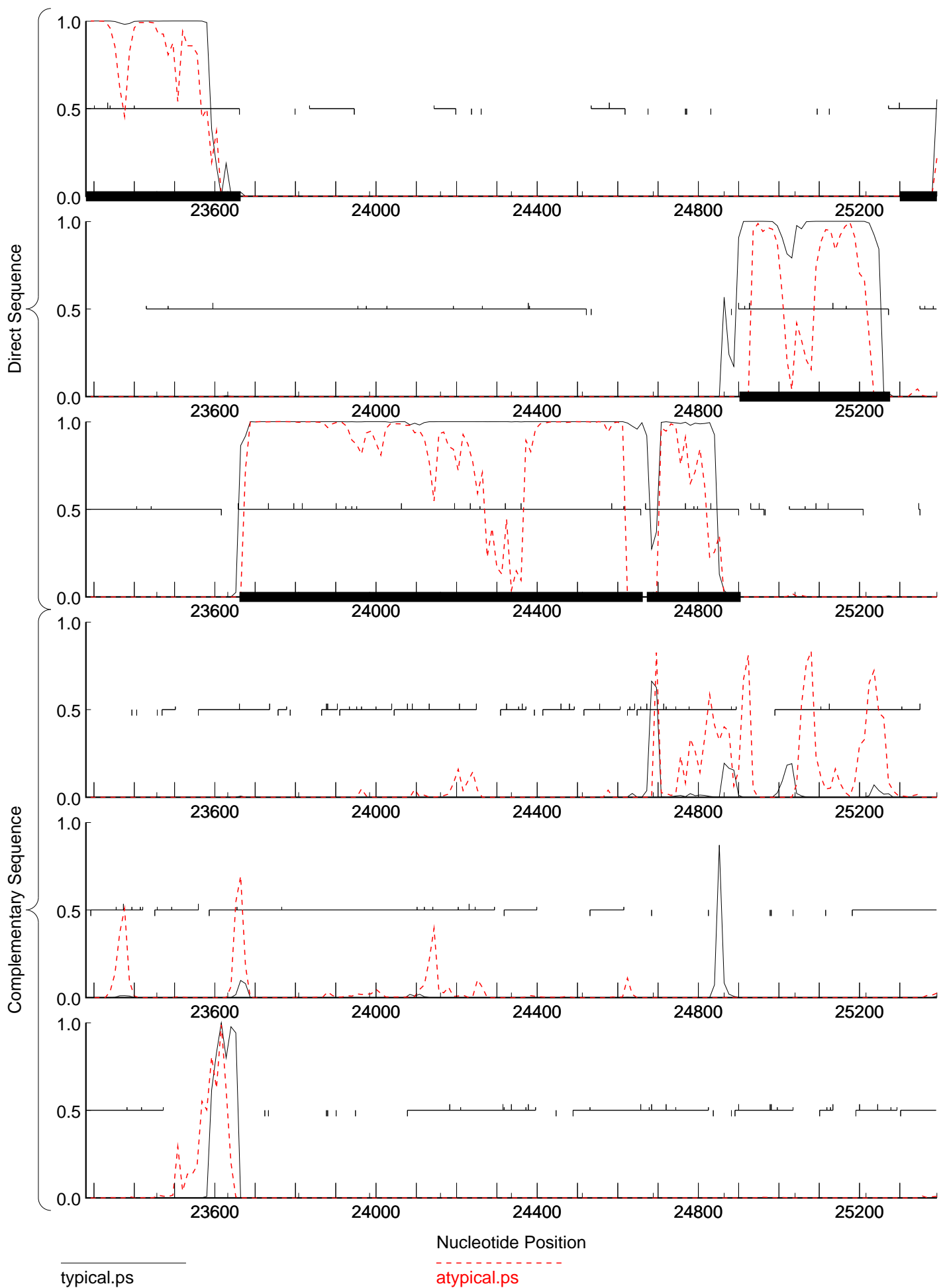
typical.ps

atypical.ps



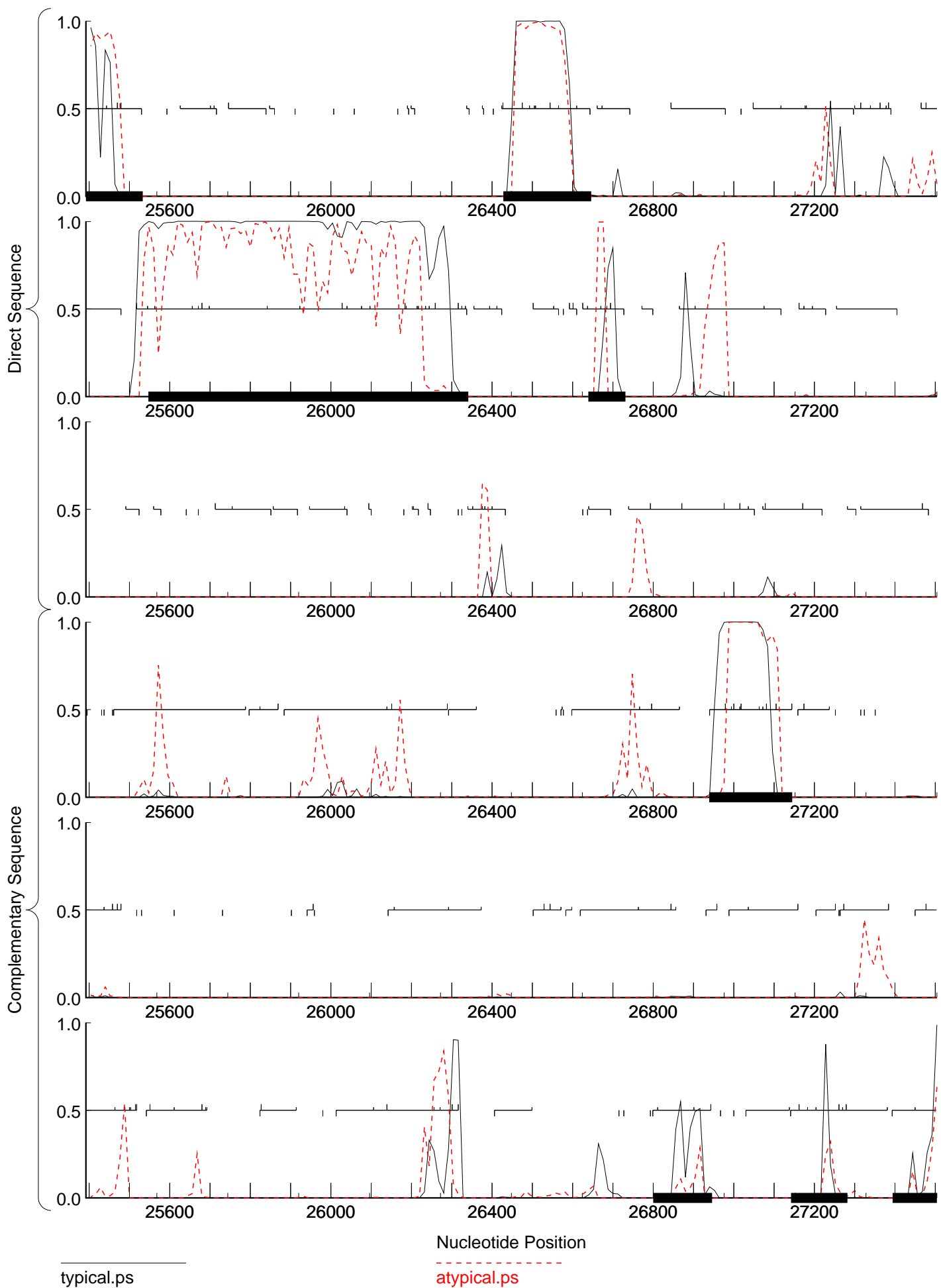
typical.ps

atypical.ps



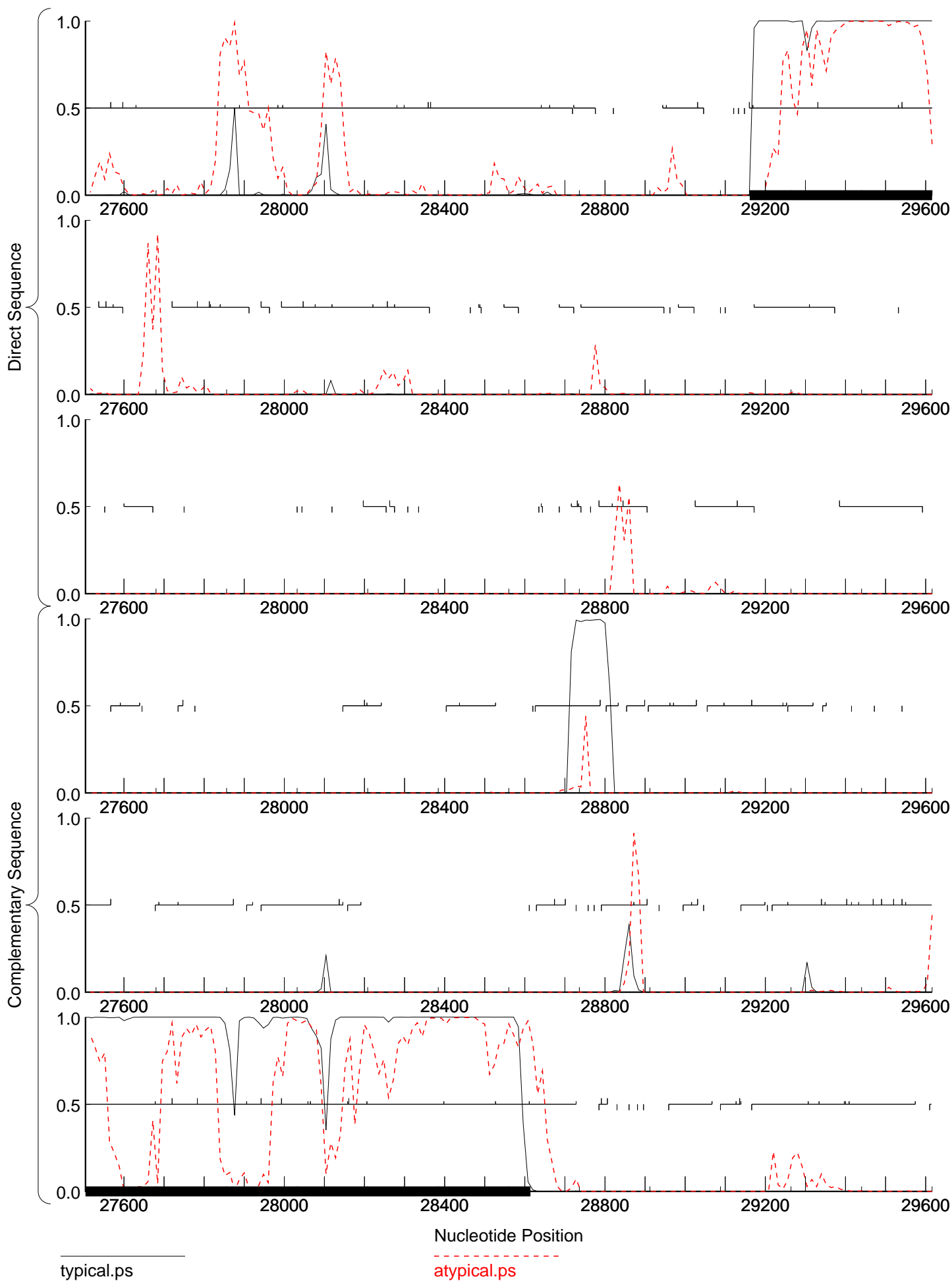
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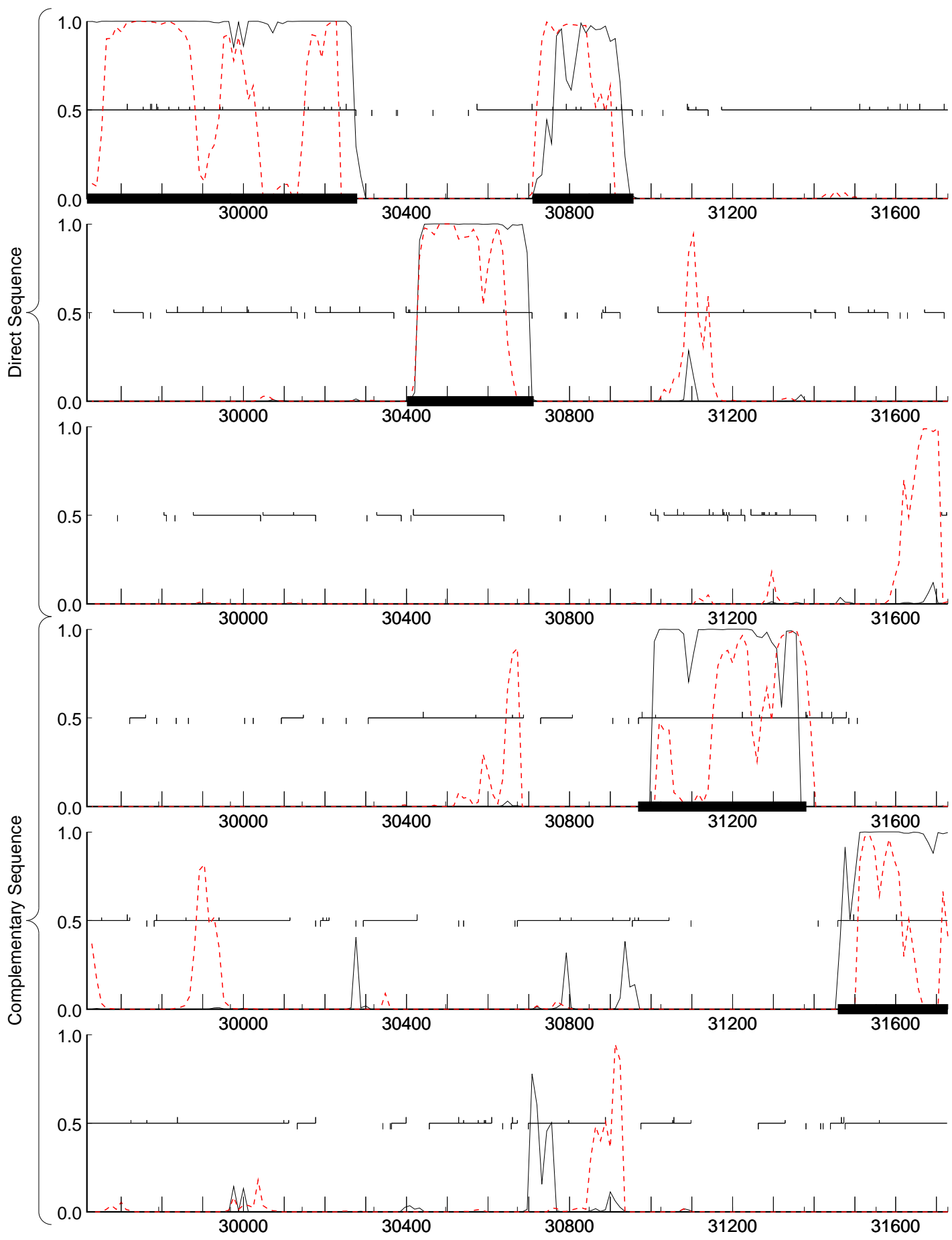
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typical.ps

atypical.ps

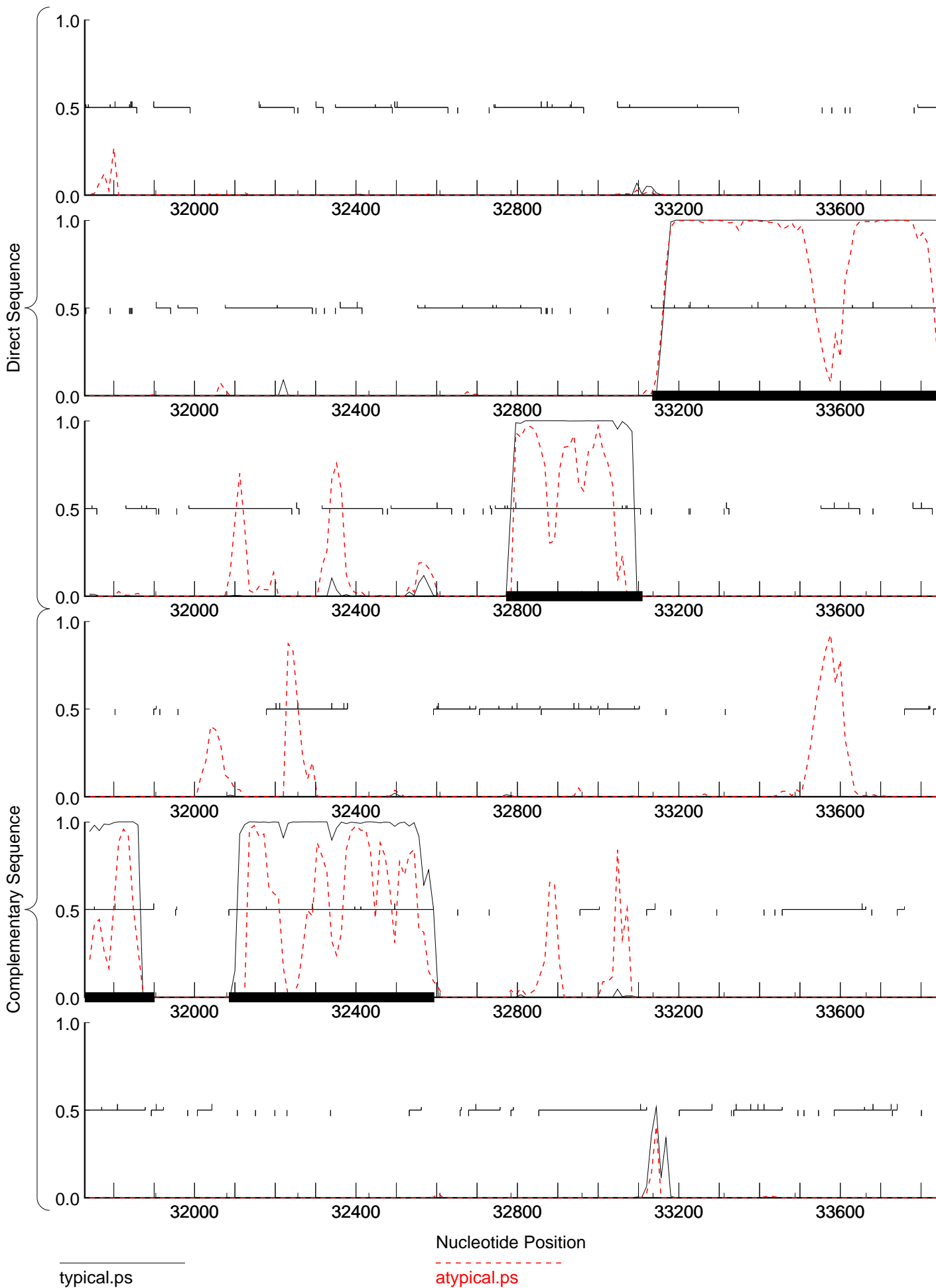


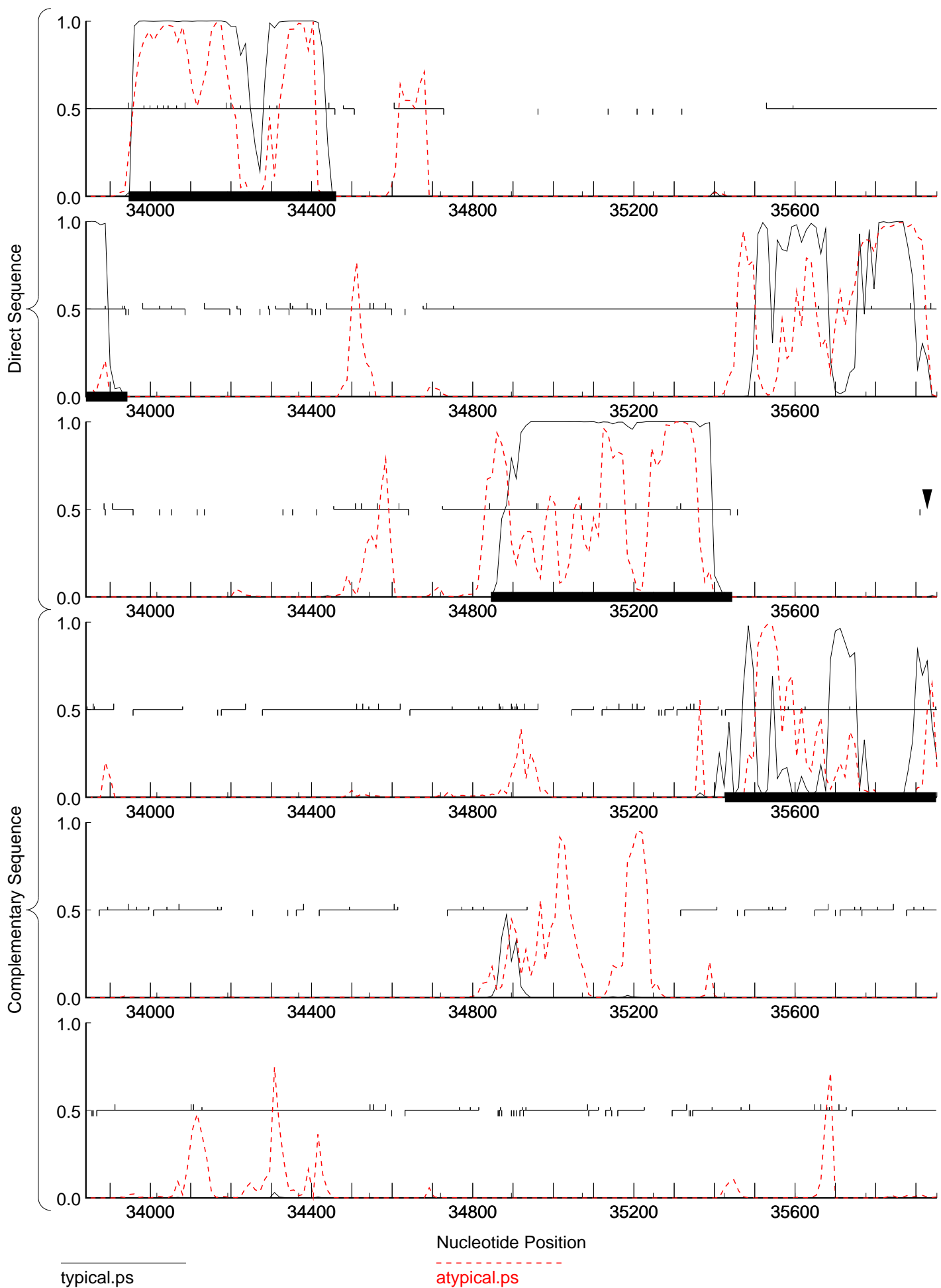


typical.ps

atypical.ps

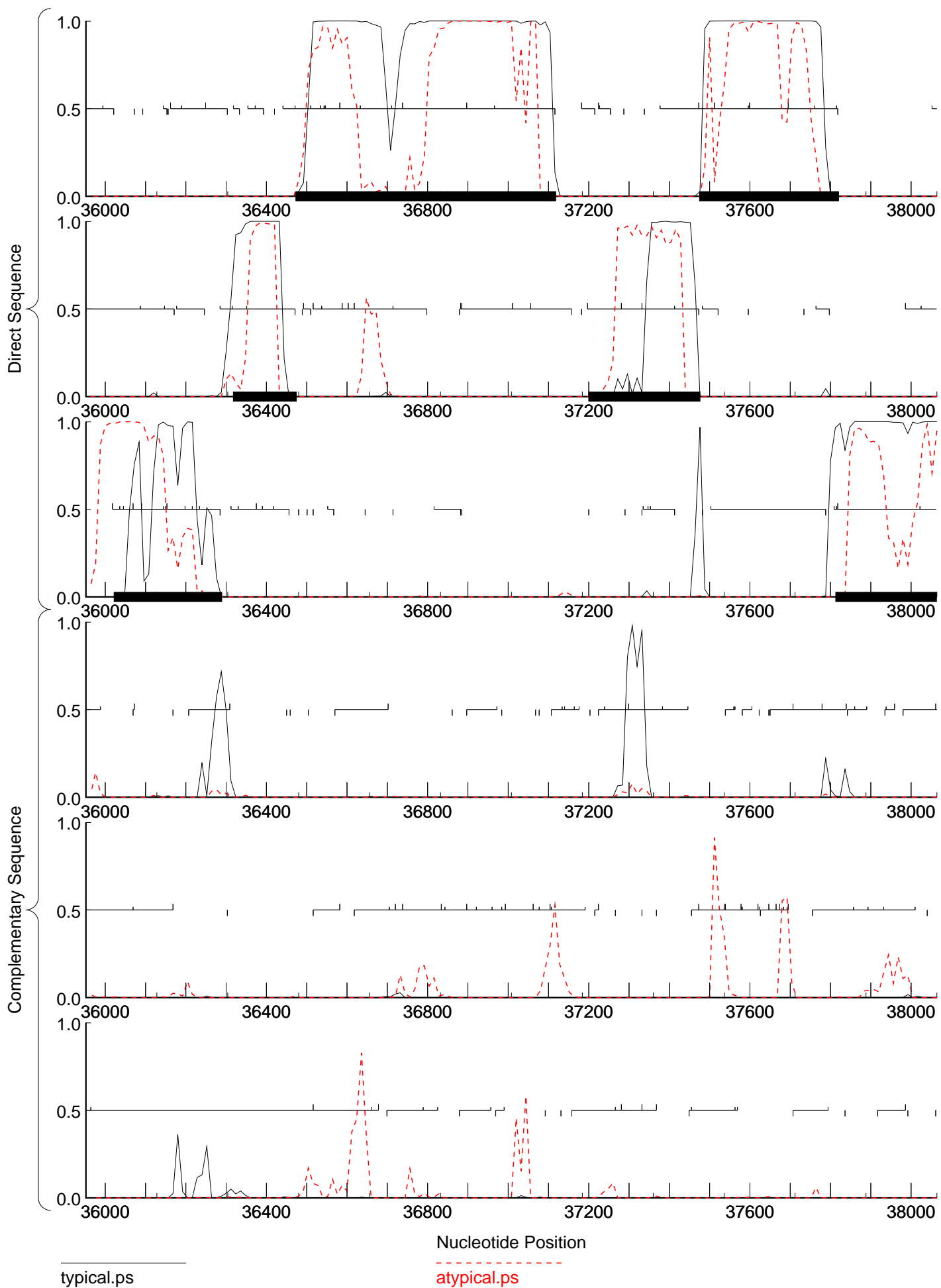






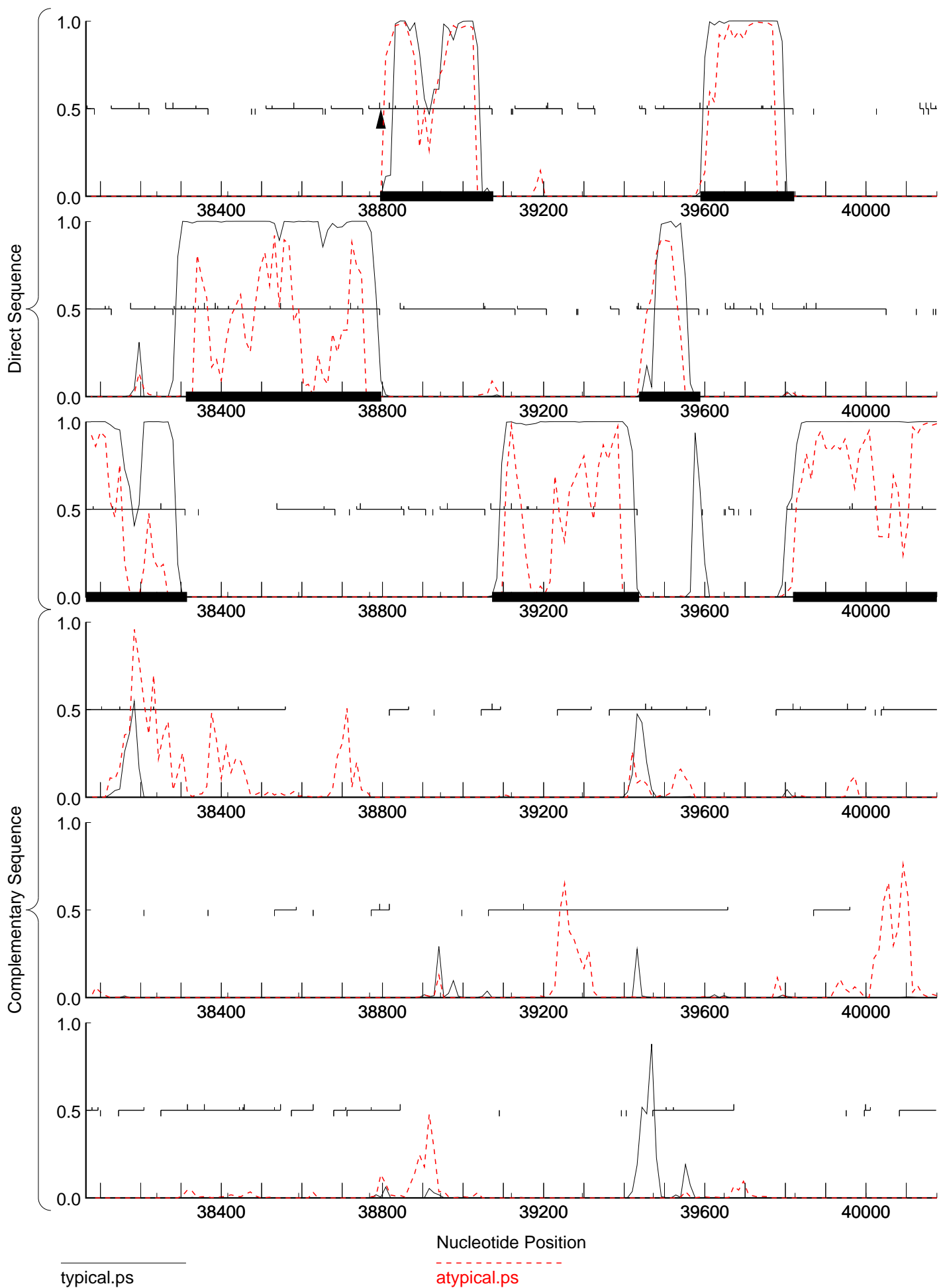
typical.ps

atypical.ps



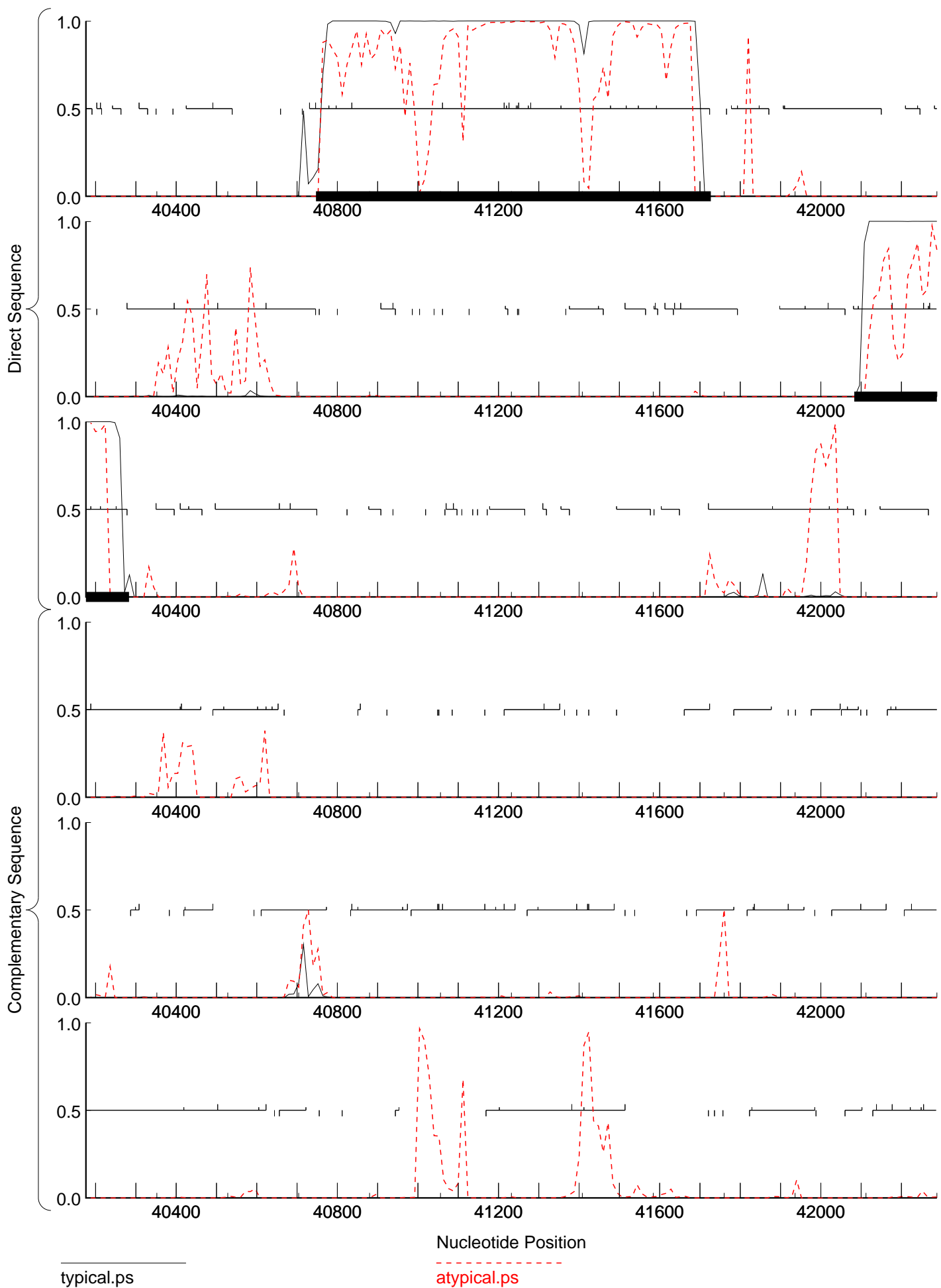
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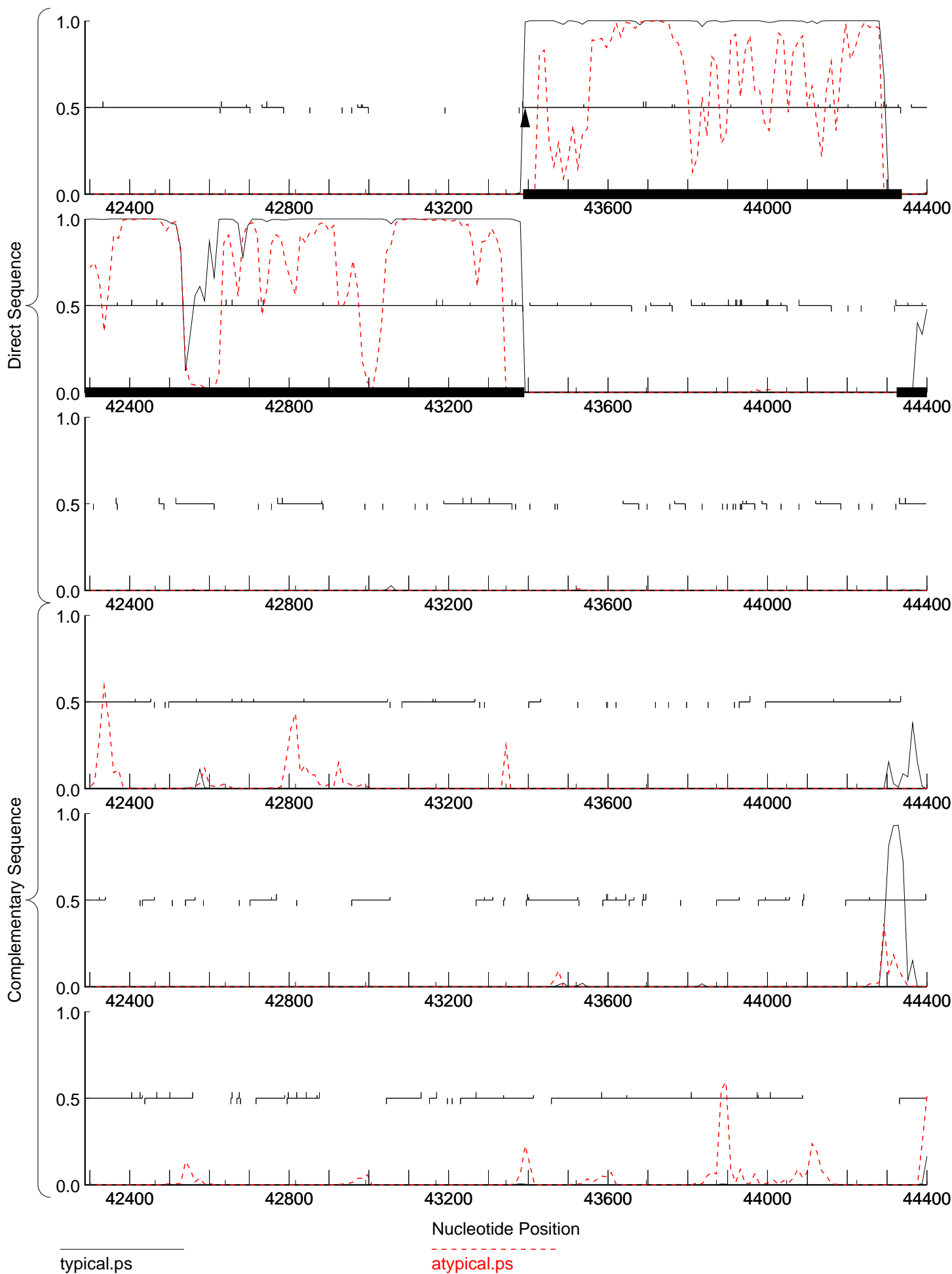
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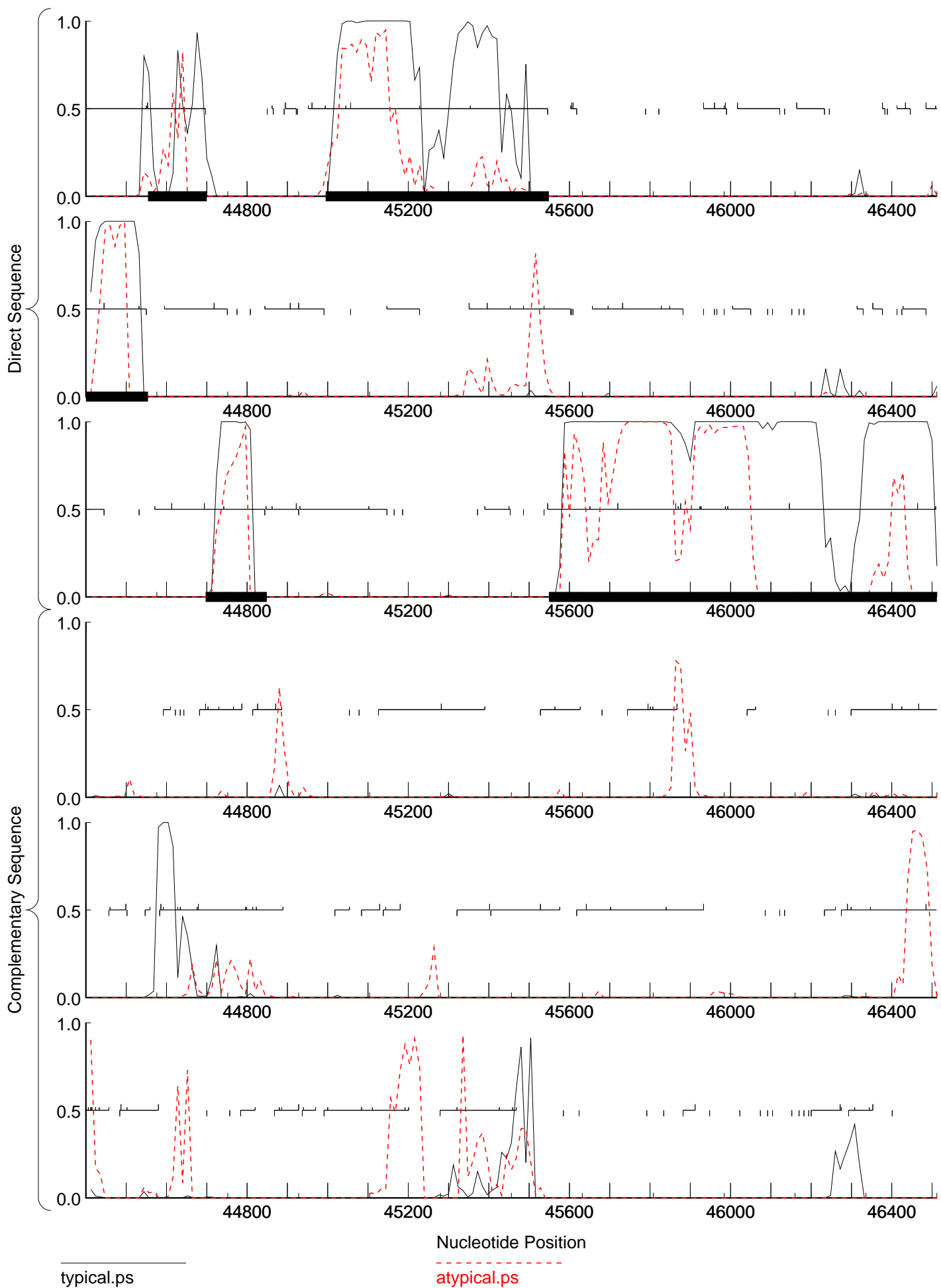


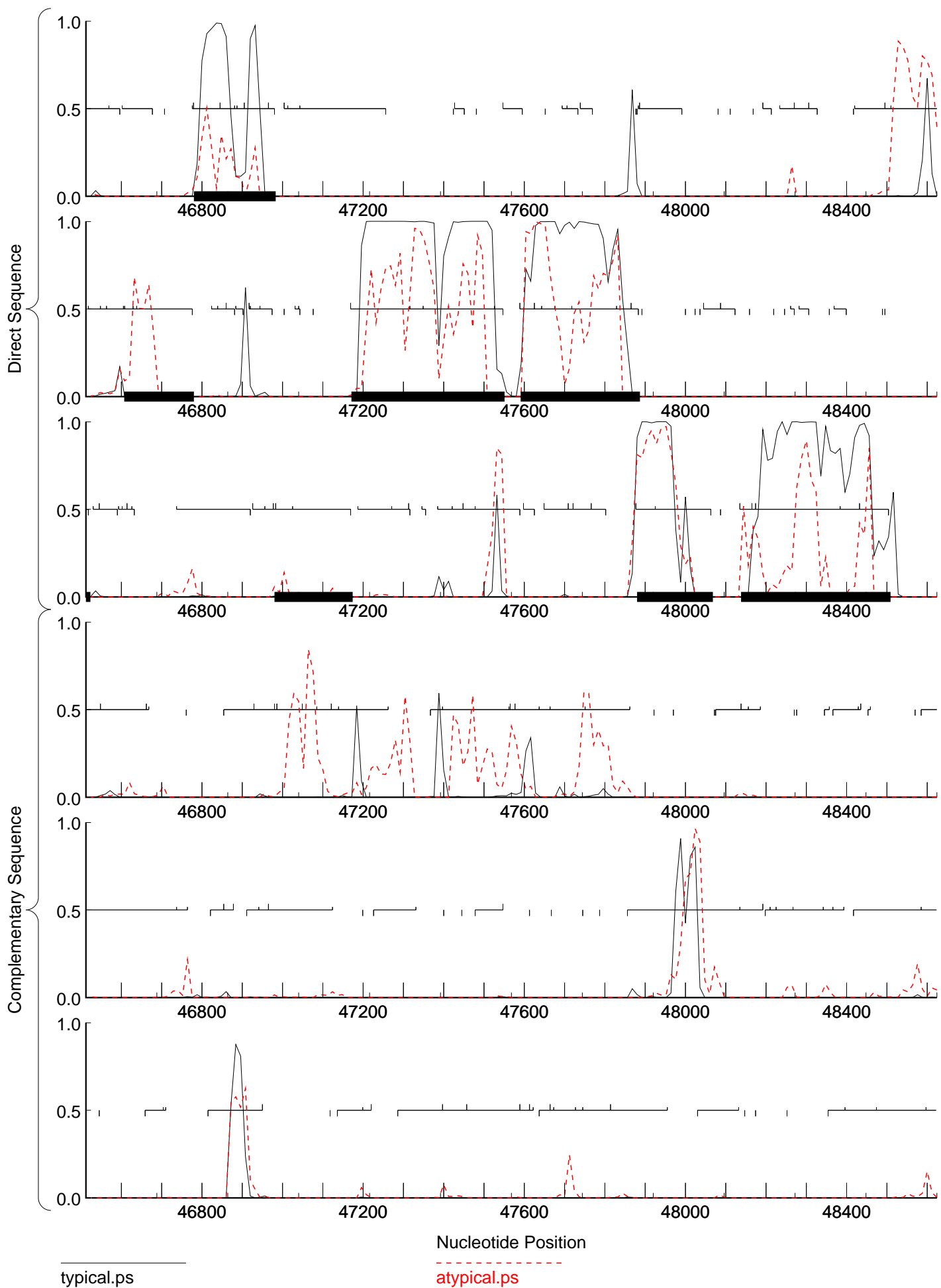
typical.ps

atypical.ps

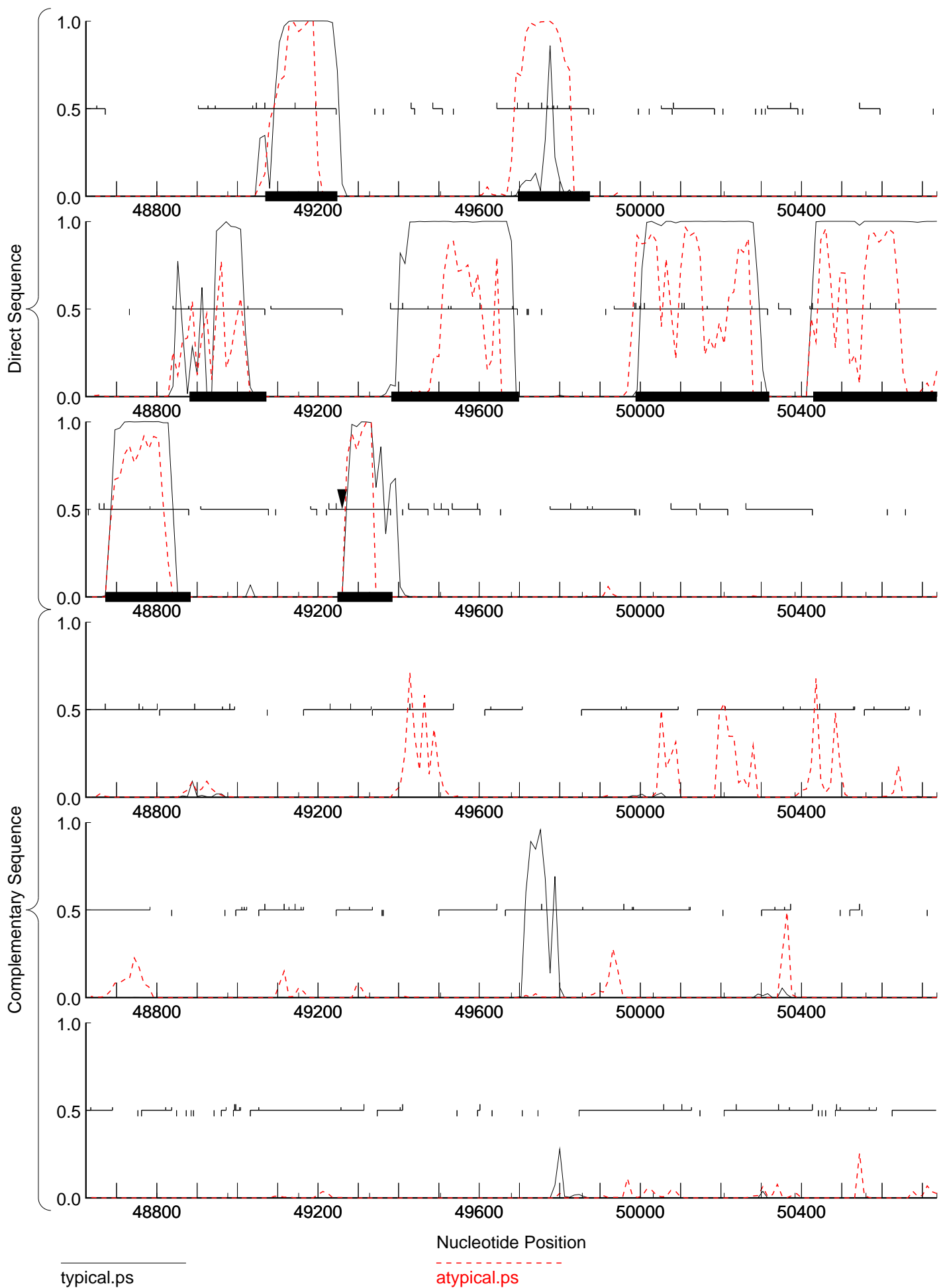


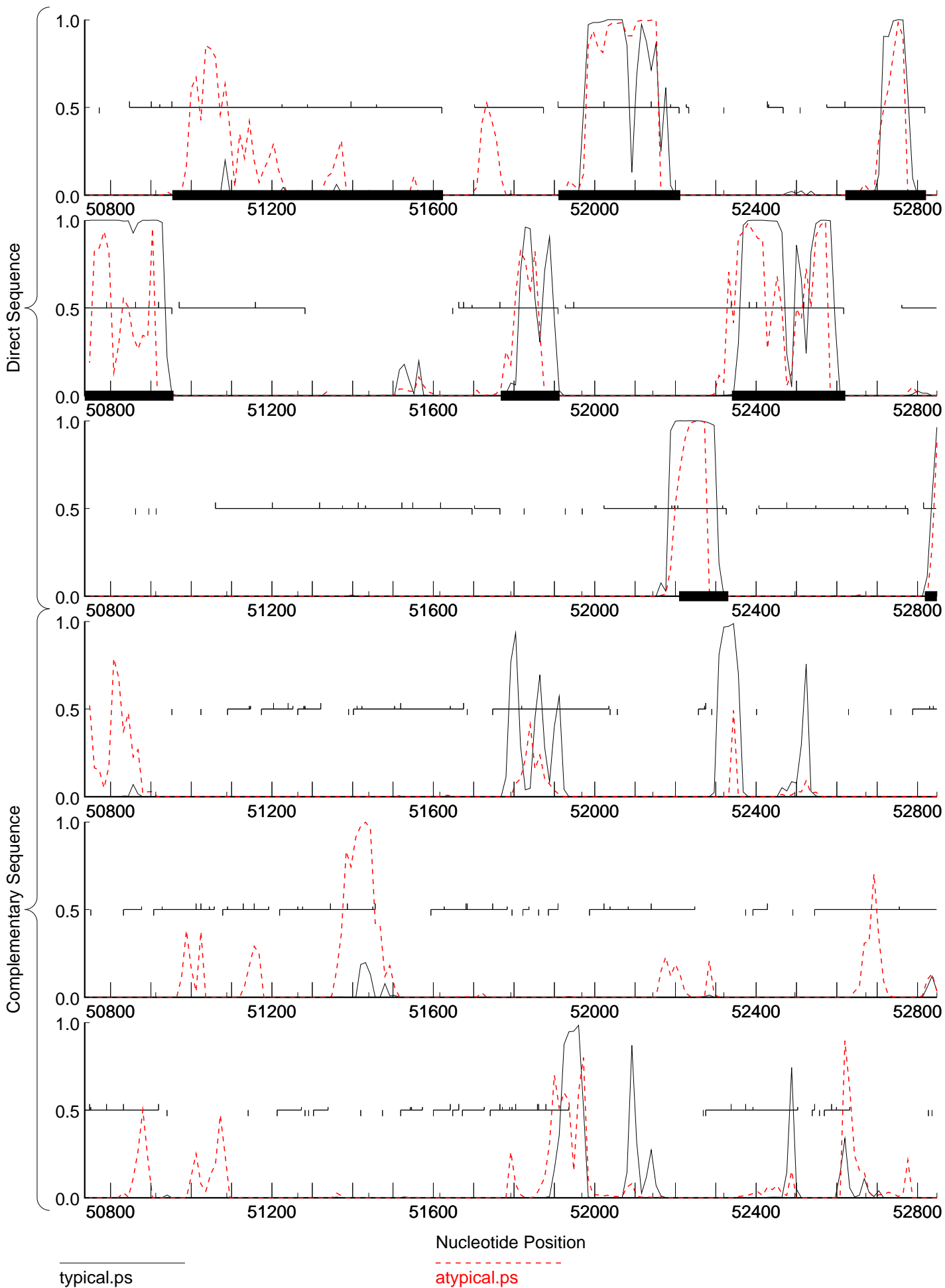


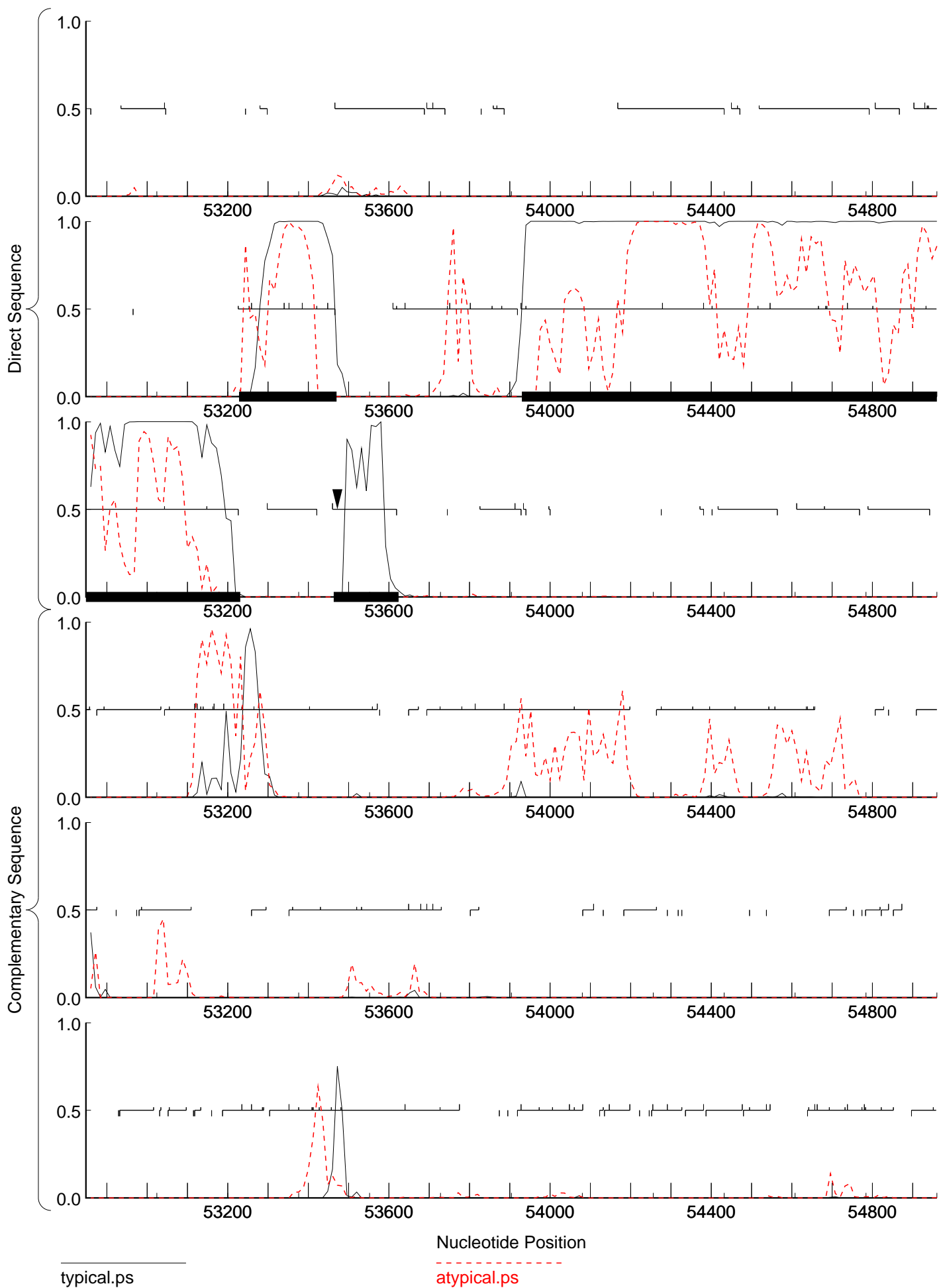


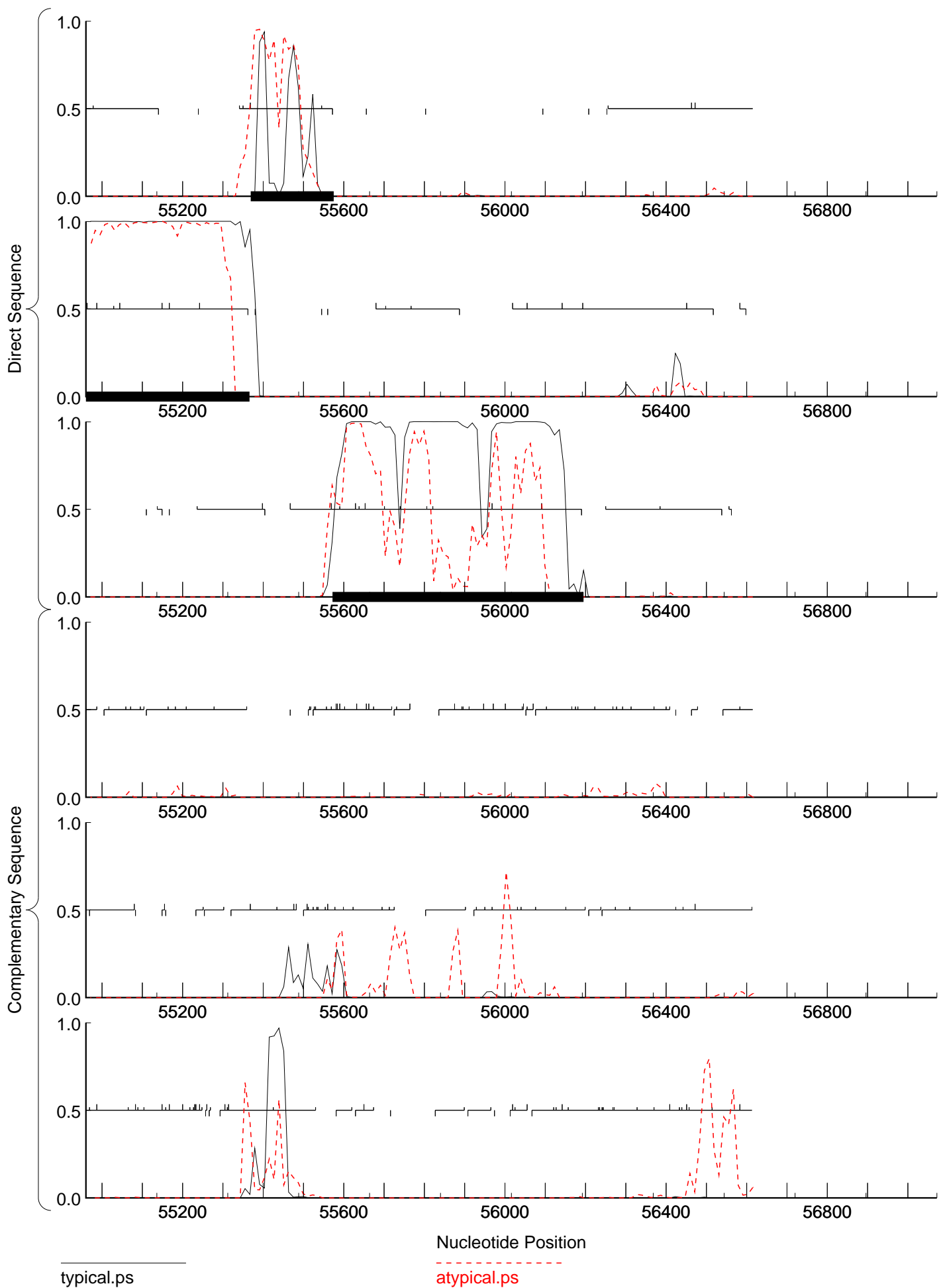












typical.ps

atypical.ps