

# GeneMark

Version 2.5p (09.08.06)

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## PROGRAM INFORMATION

Sequence : Mycobacterium phage Moose complete sequence, 52695 bp including 10-base 3' overhang (CGGATG  
Analysis Date : 6/24/18 at 22:36:51  
Pages : 26  
Sequence Length : 52695 bp  
GC Content : 63.71%

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,

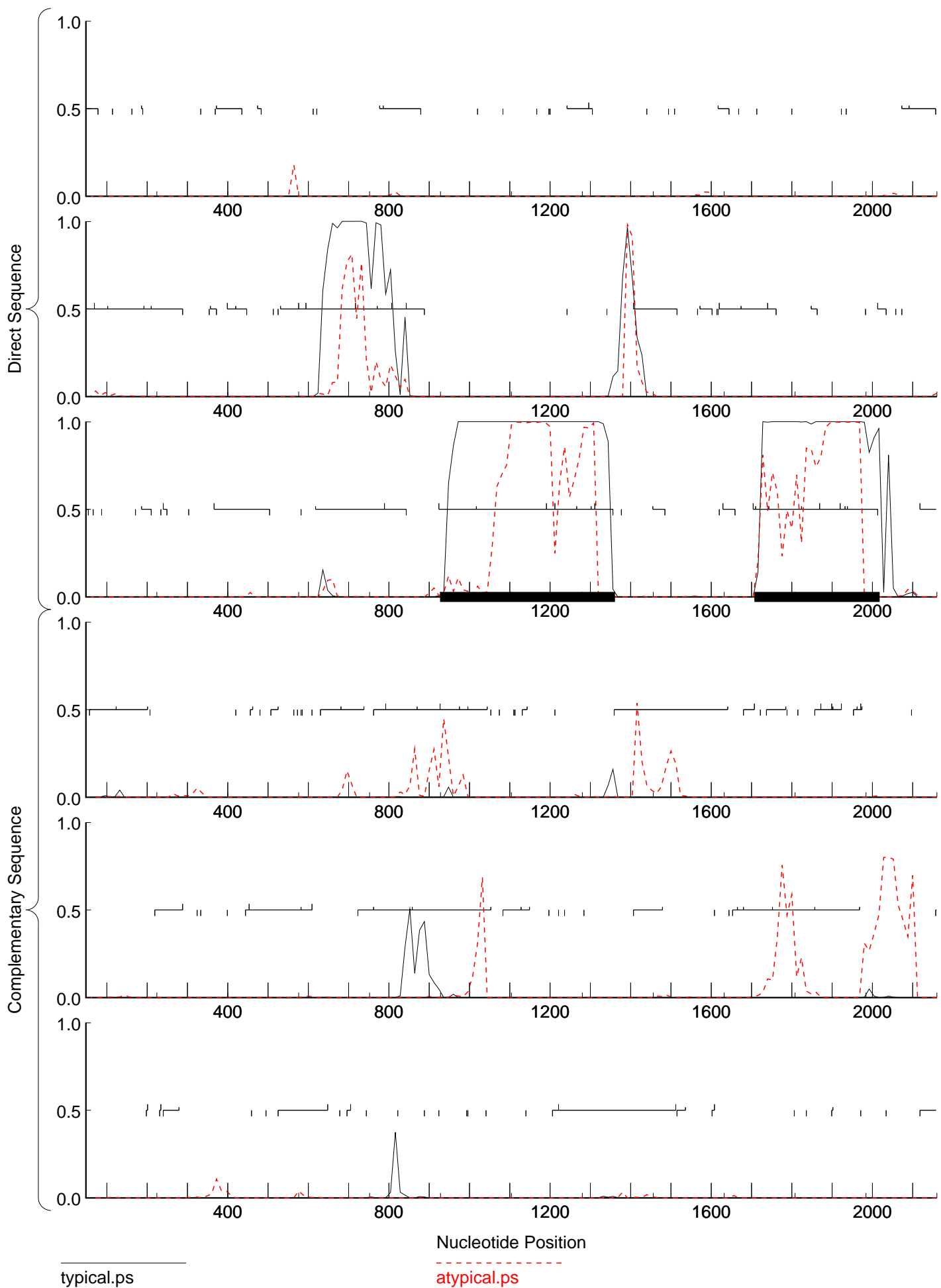
## Matrix notes & comments

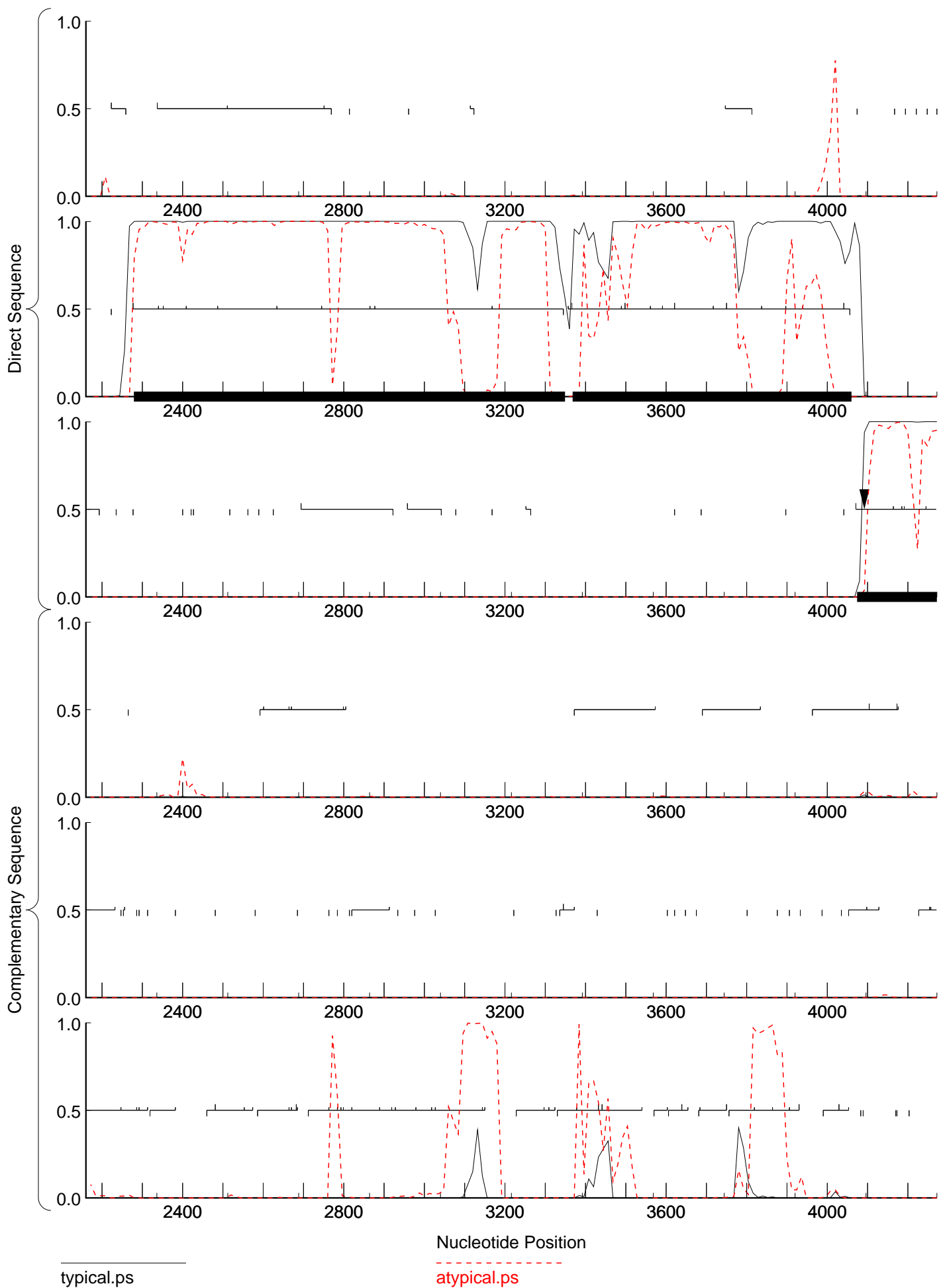
Training set derived by GeneMarkS, 4.30 October 2014  
Sun Jun 24 22:36:51 2018

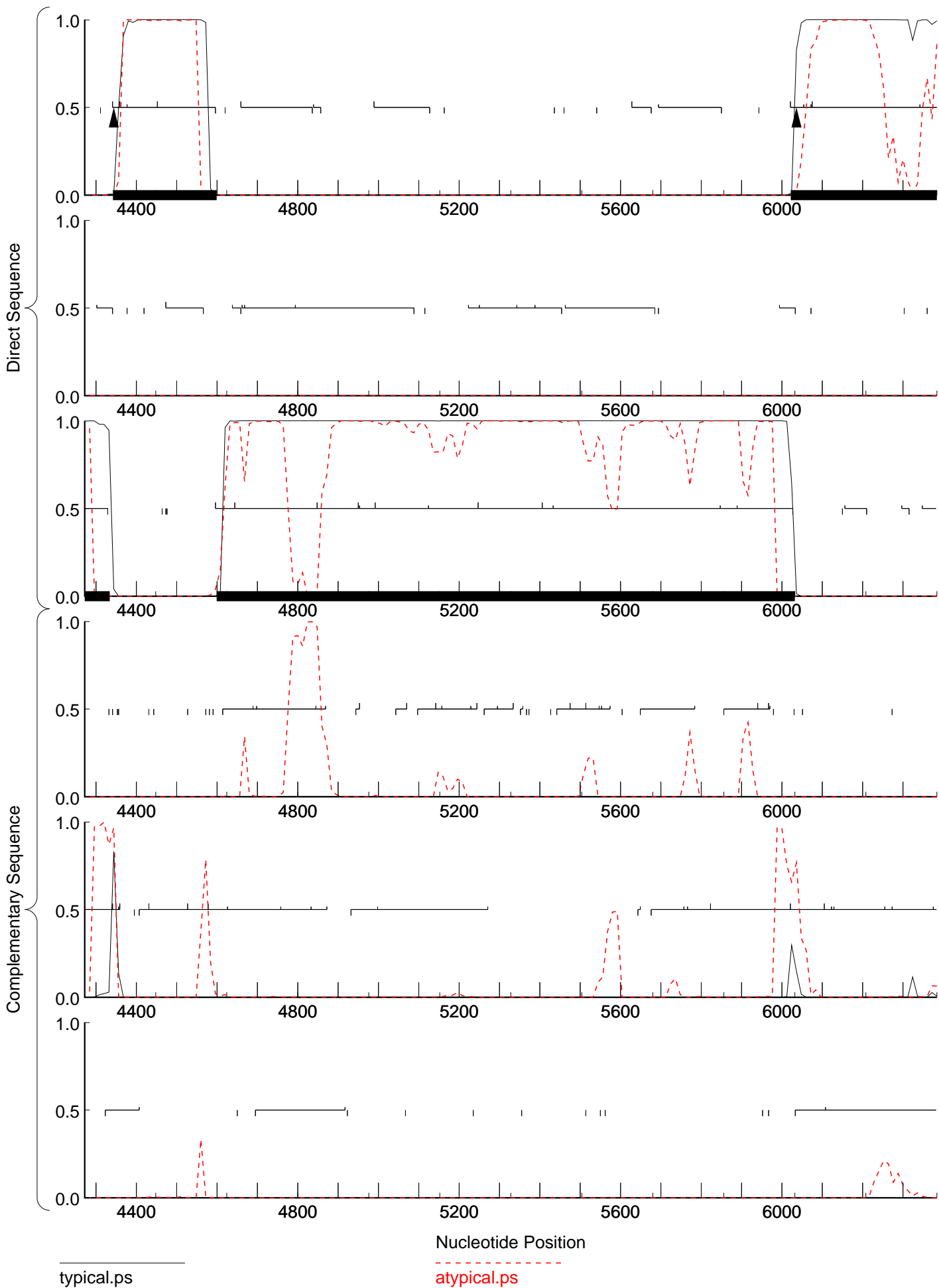
## MATRIX INFORMATION

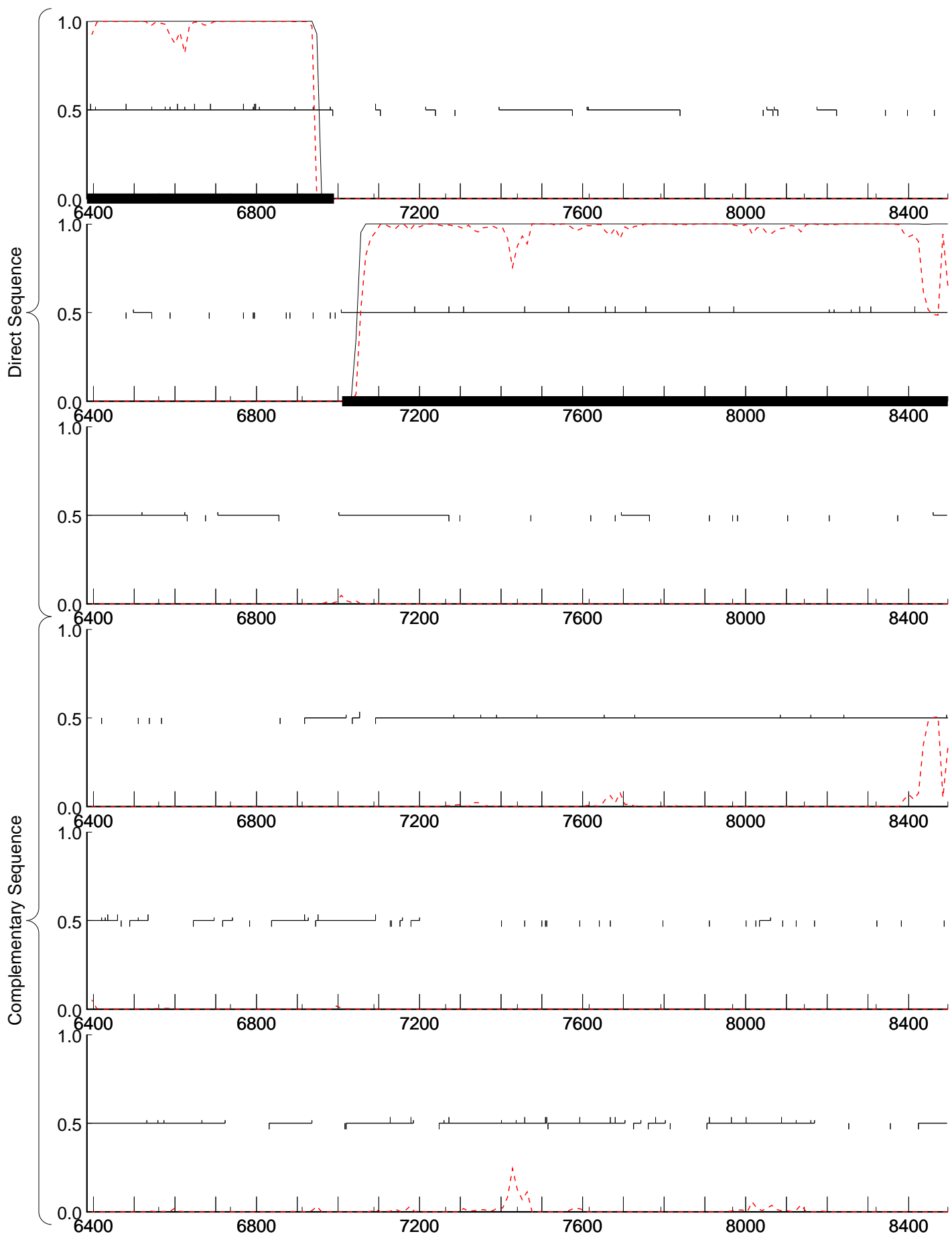
Matrix : -  
Author : -  
Order : 4

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



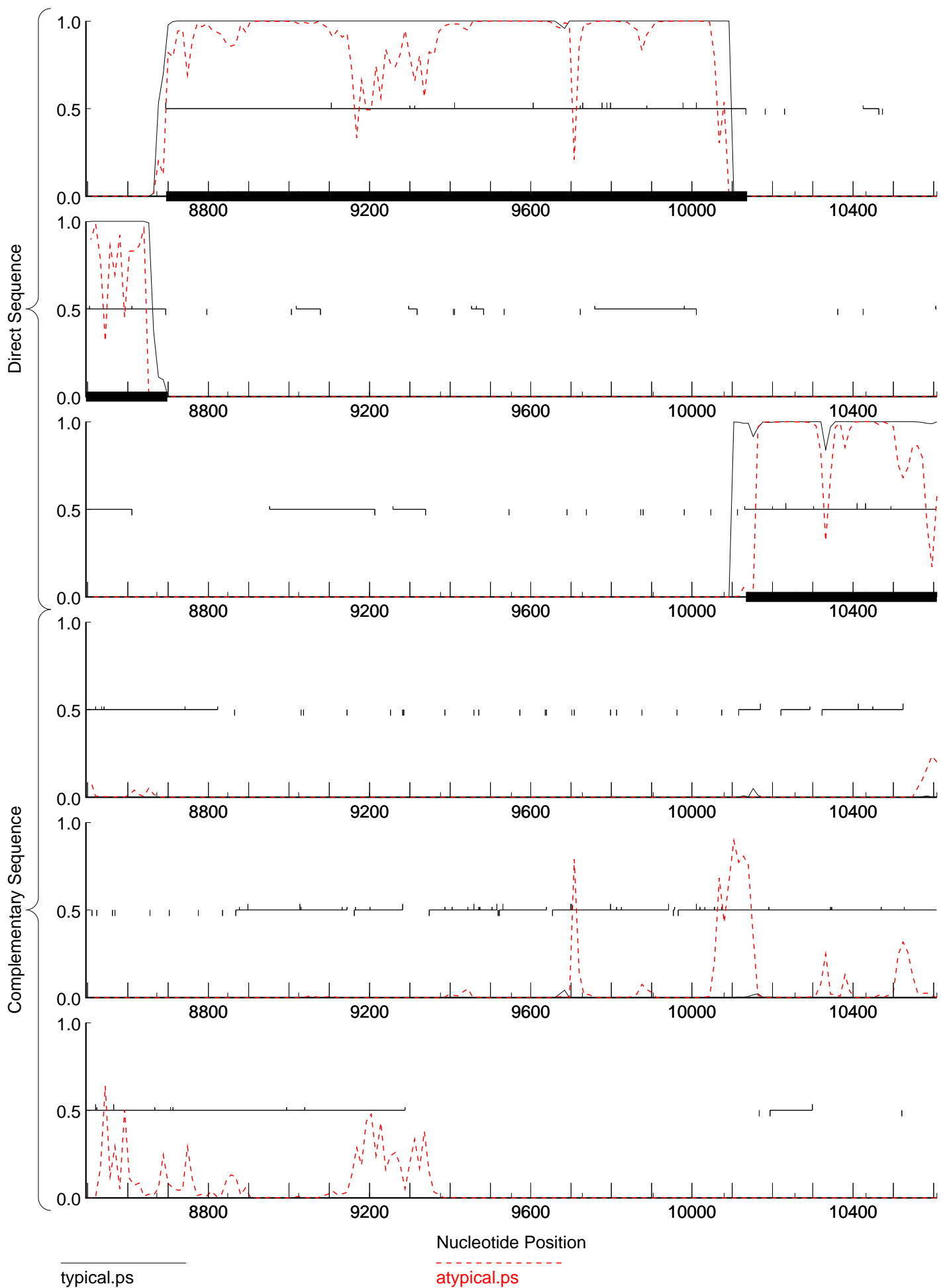






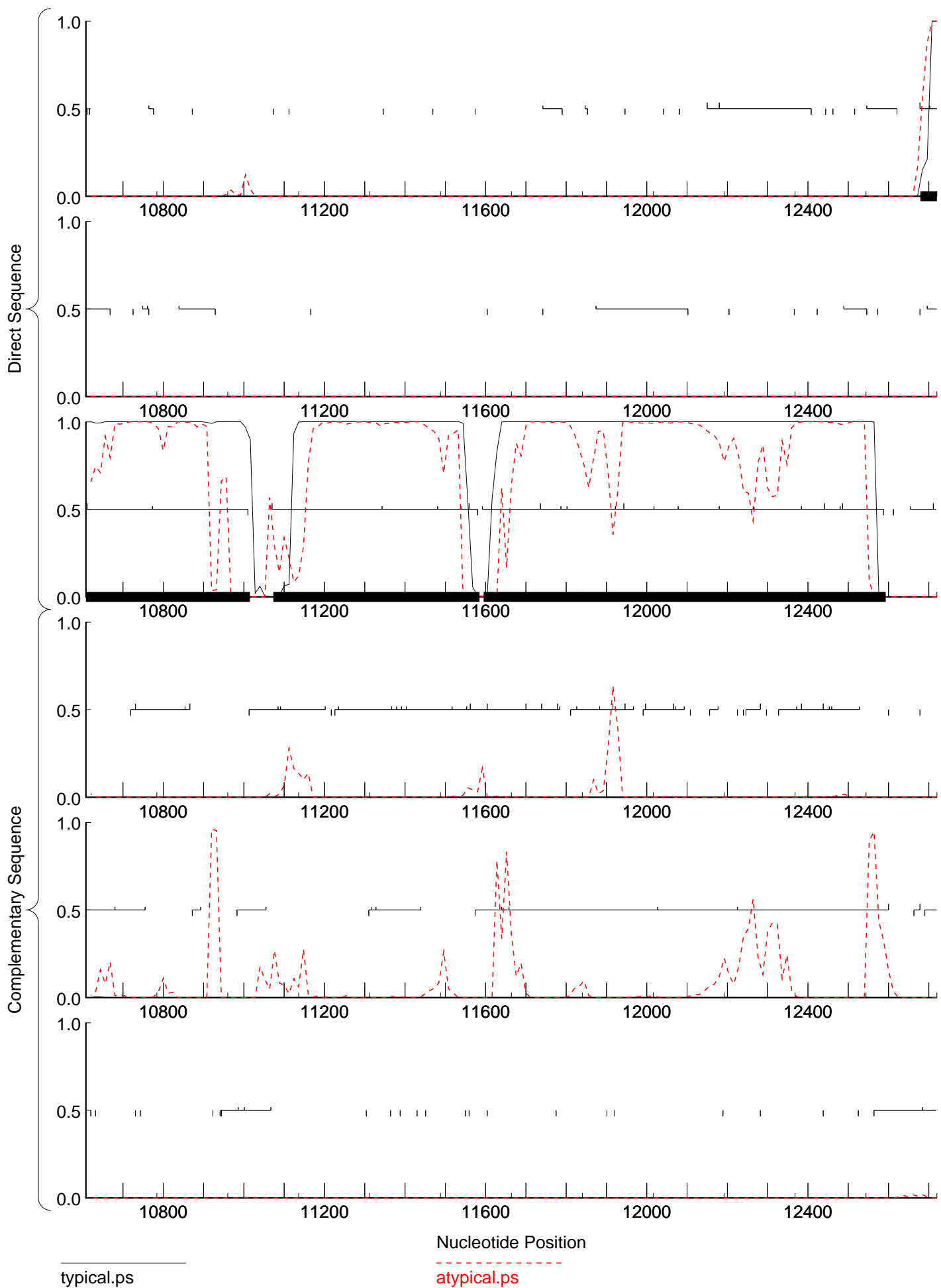
typical.ps

atypical.ps

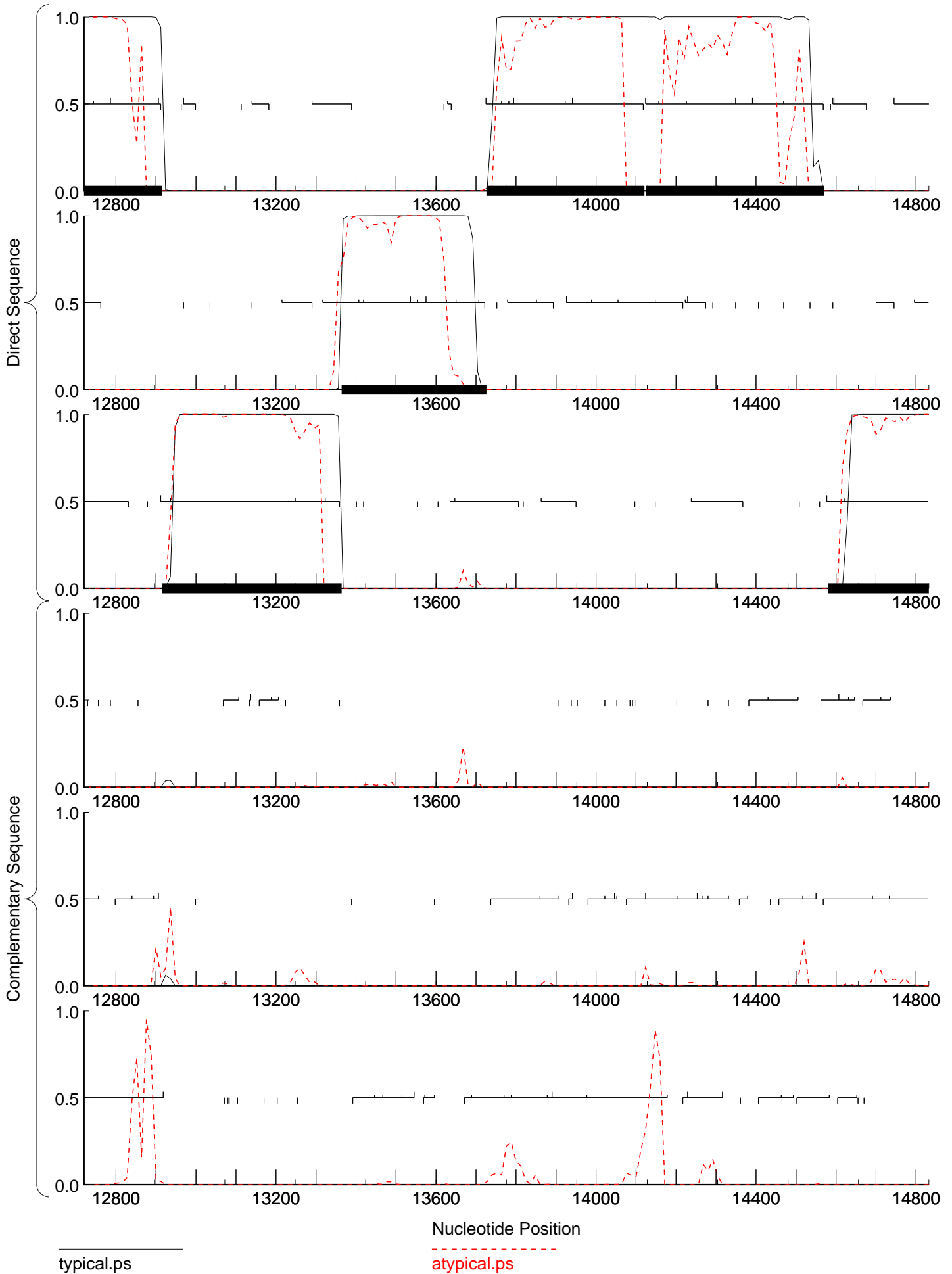


typical.ps

atypical.ps



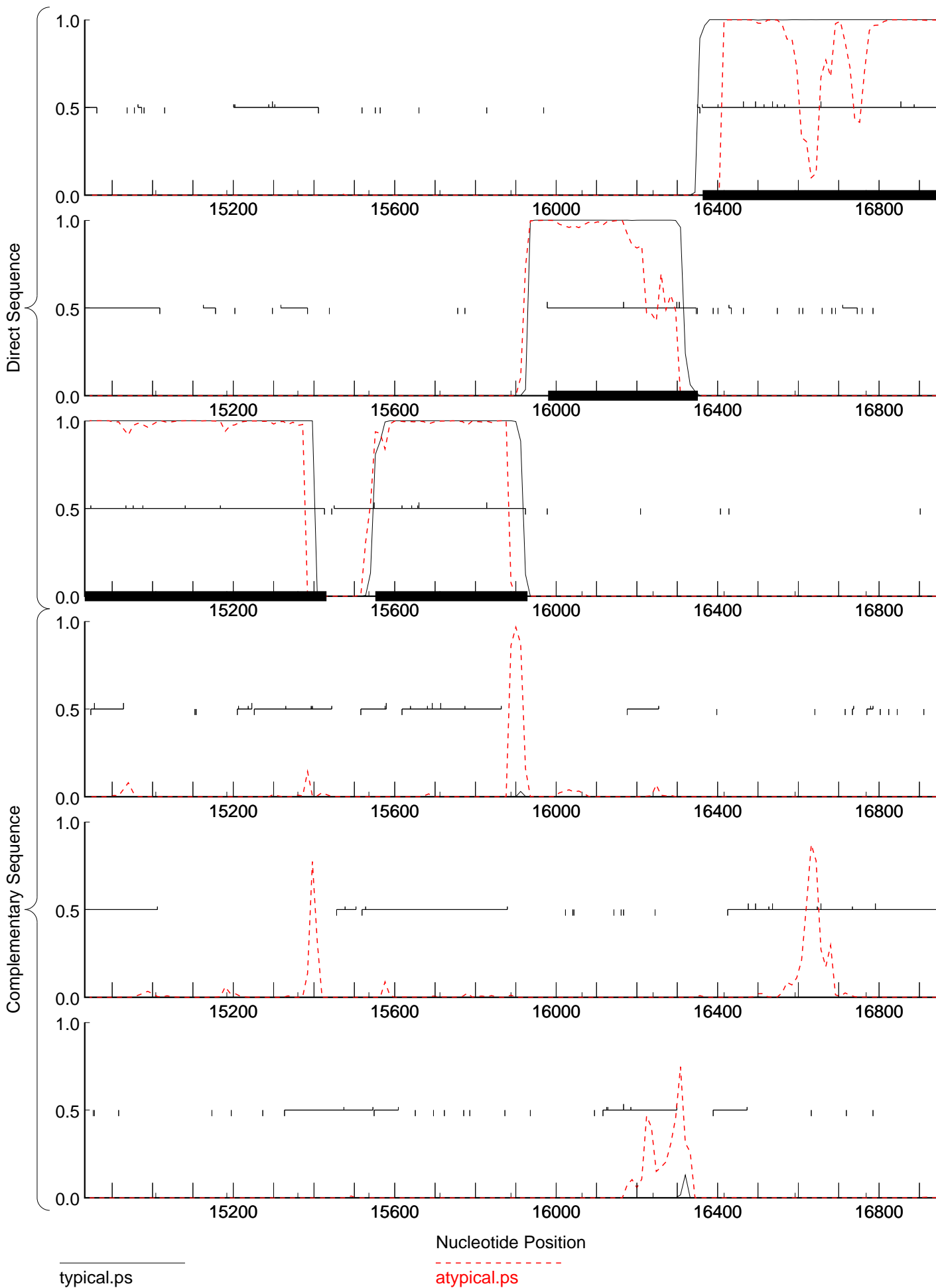
GeneMark.hmm prediction

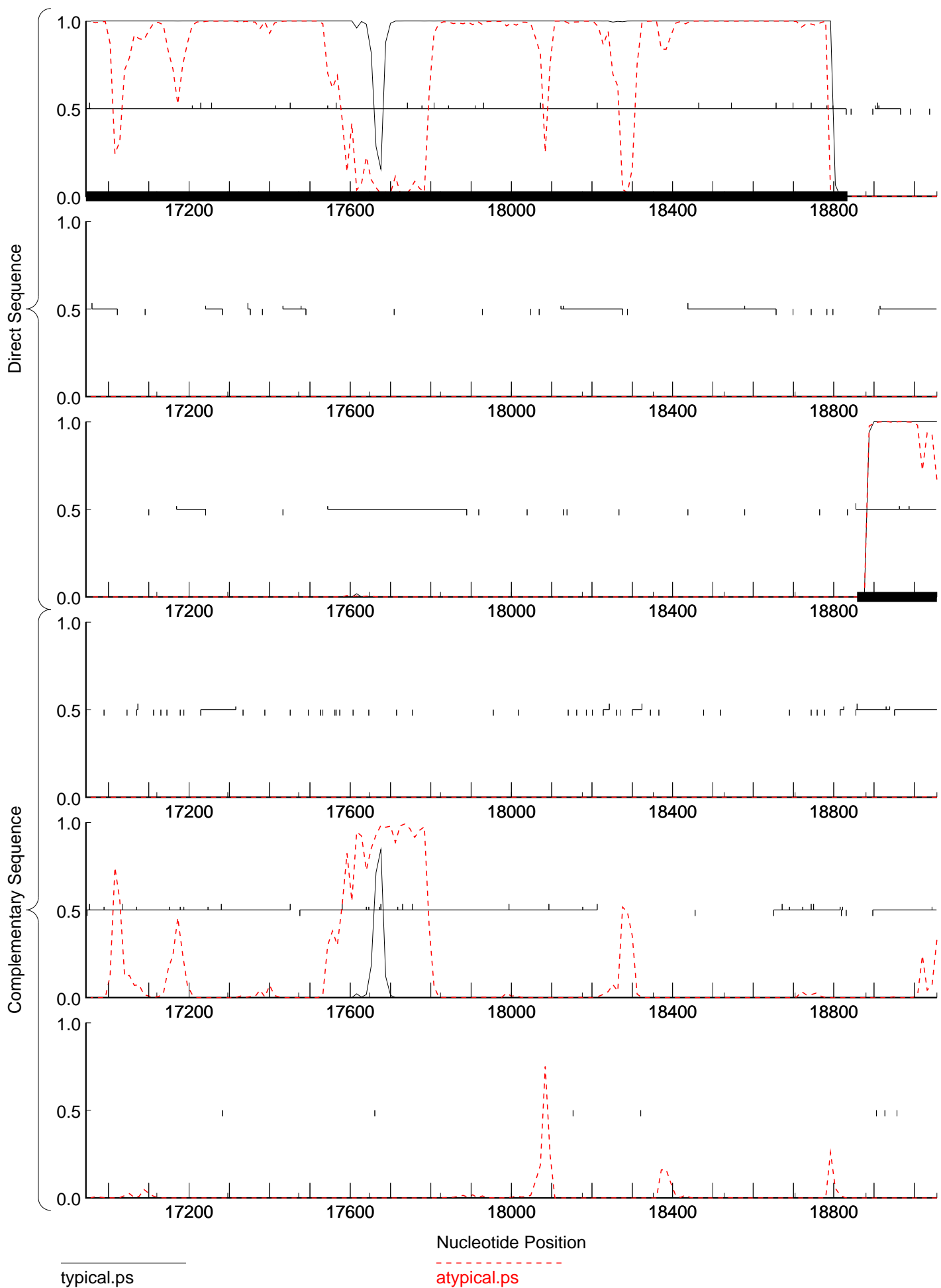


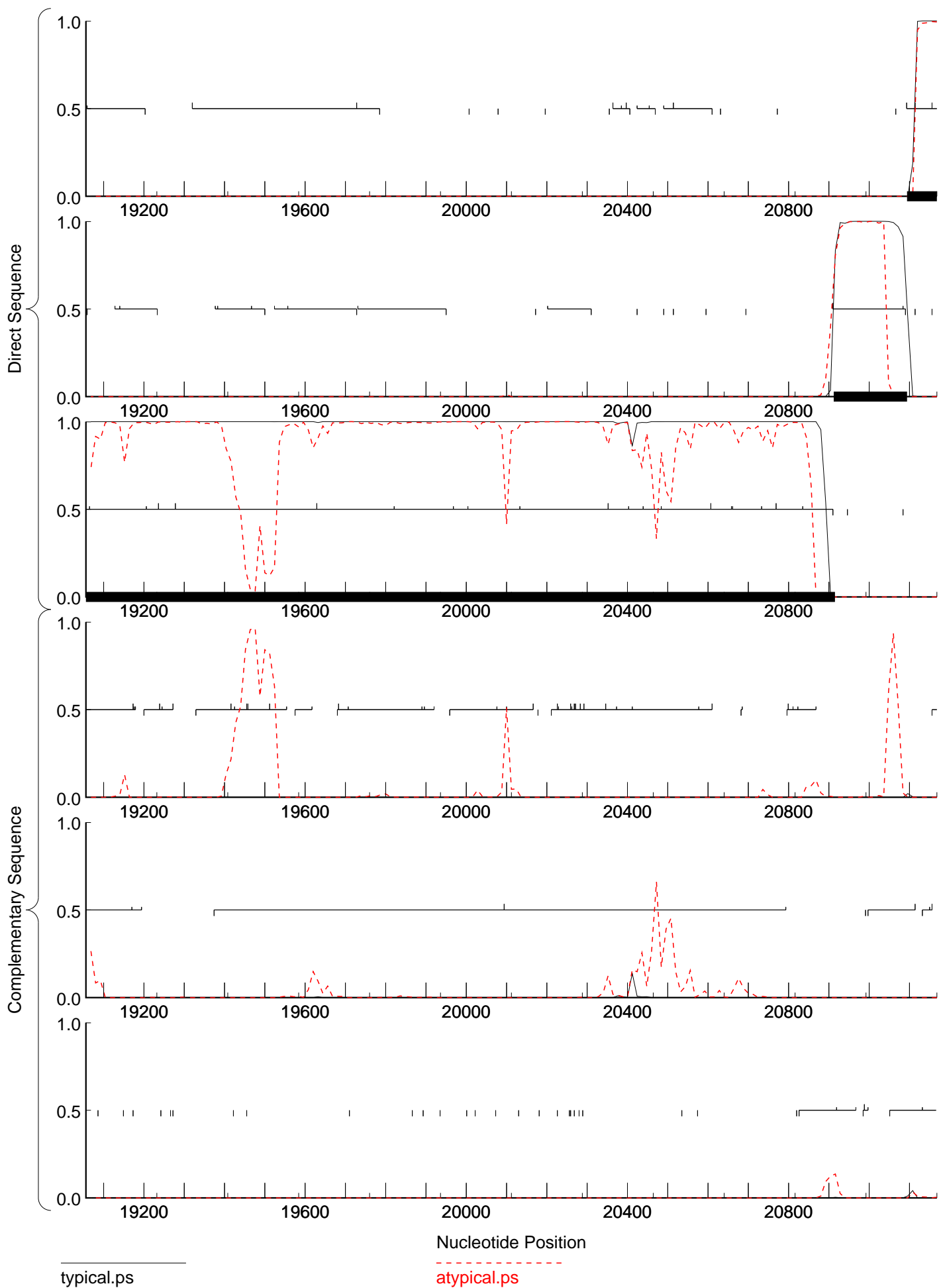
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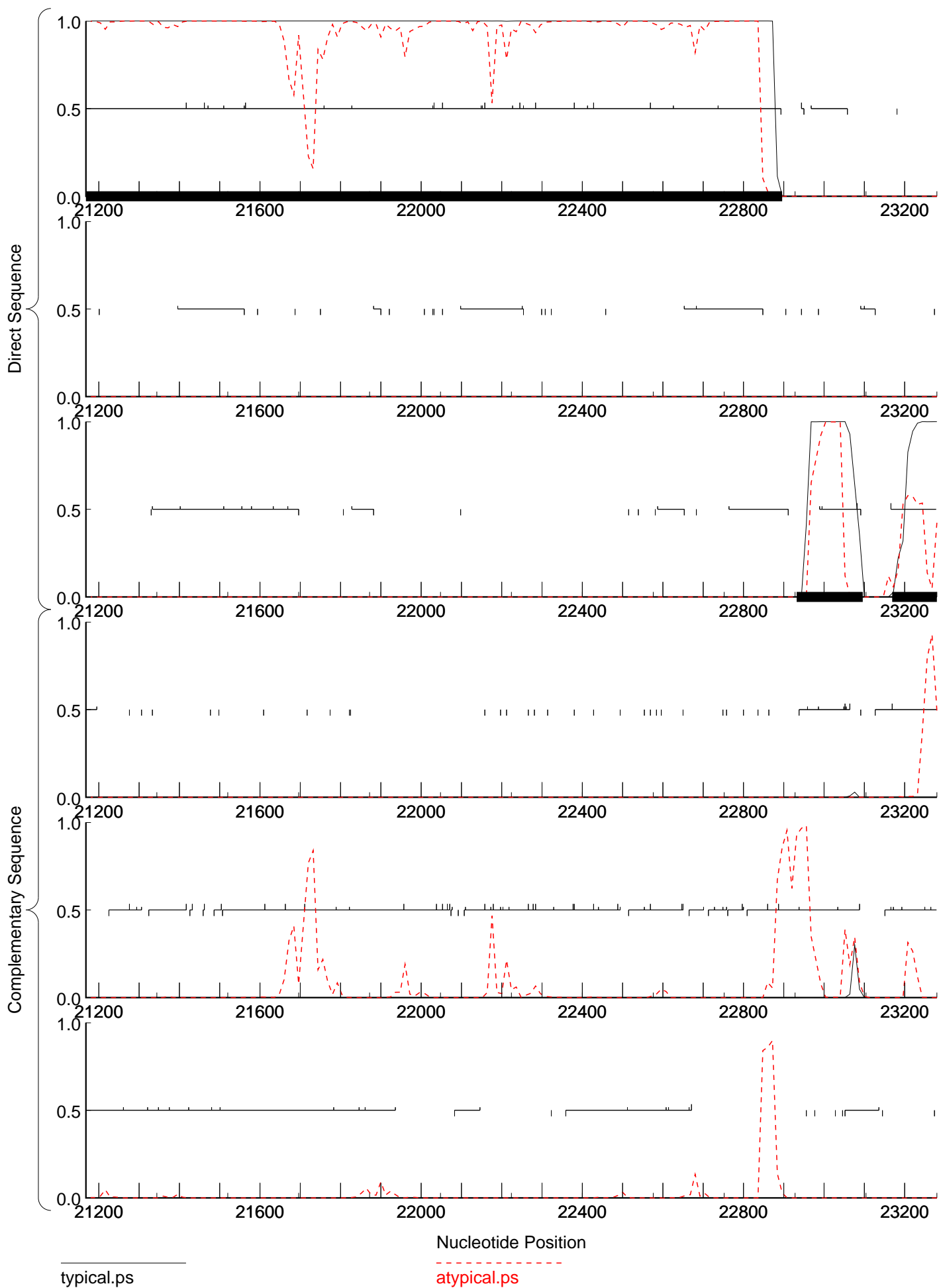
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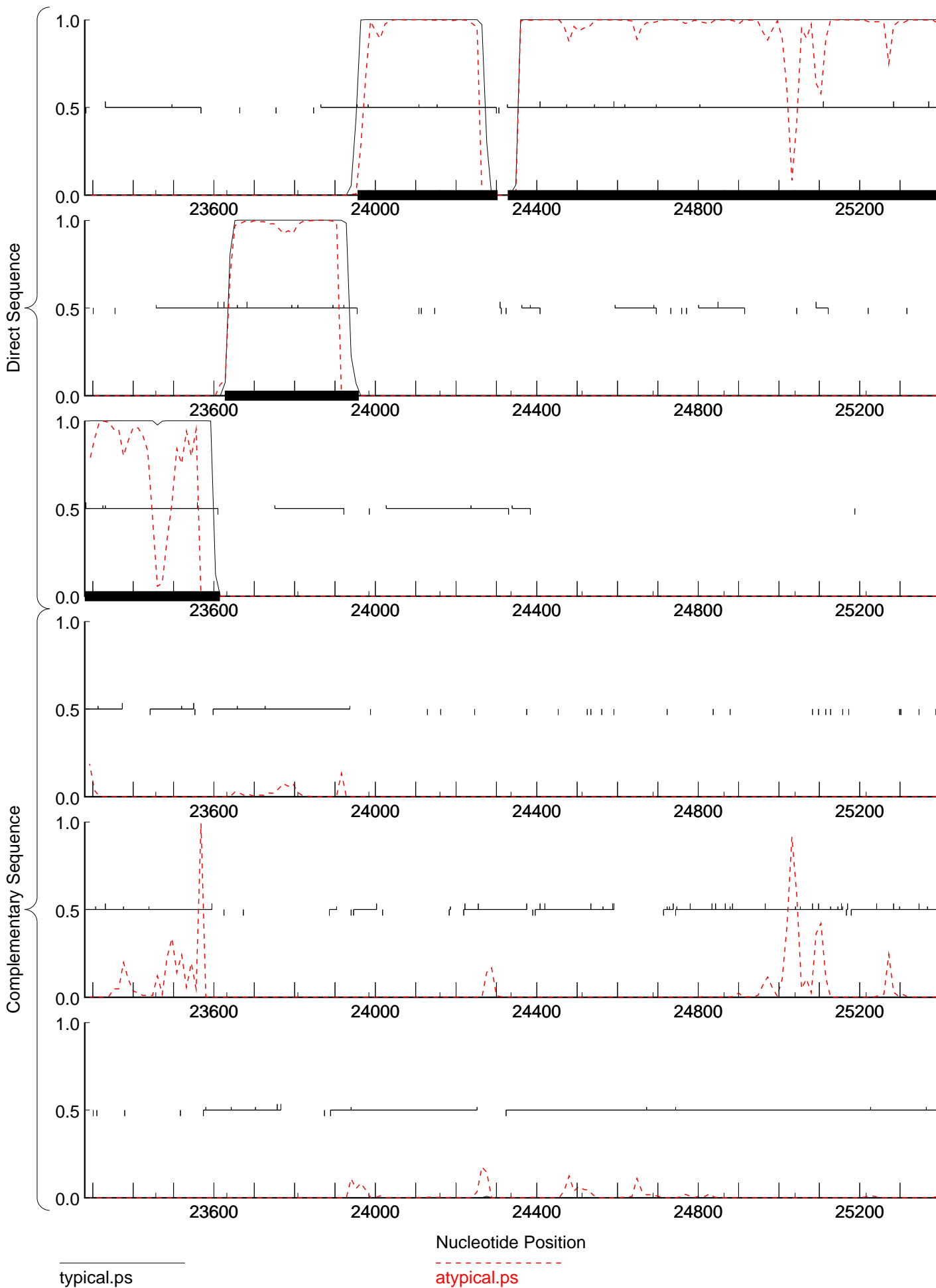


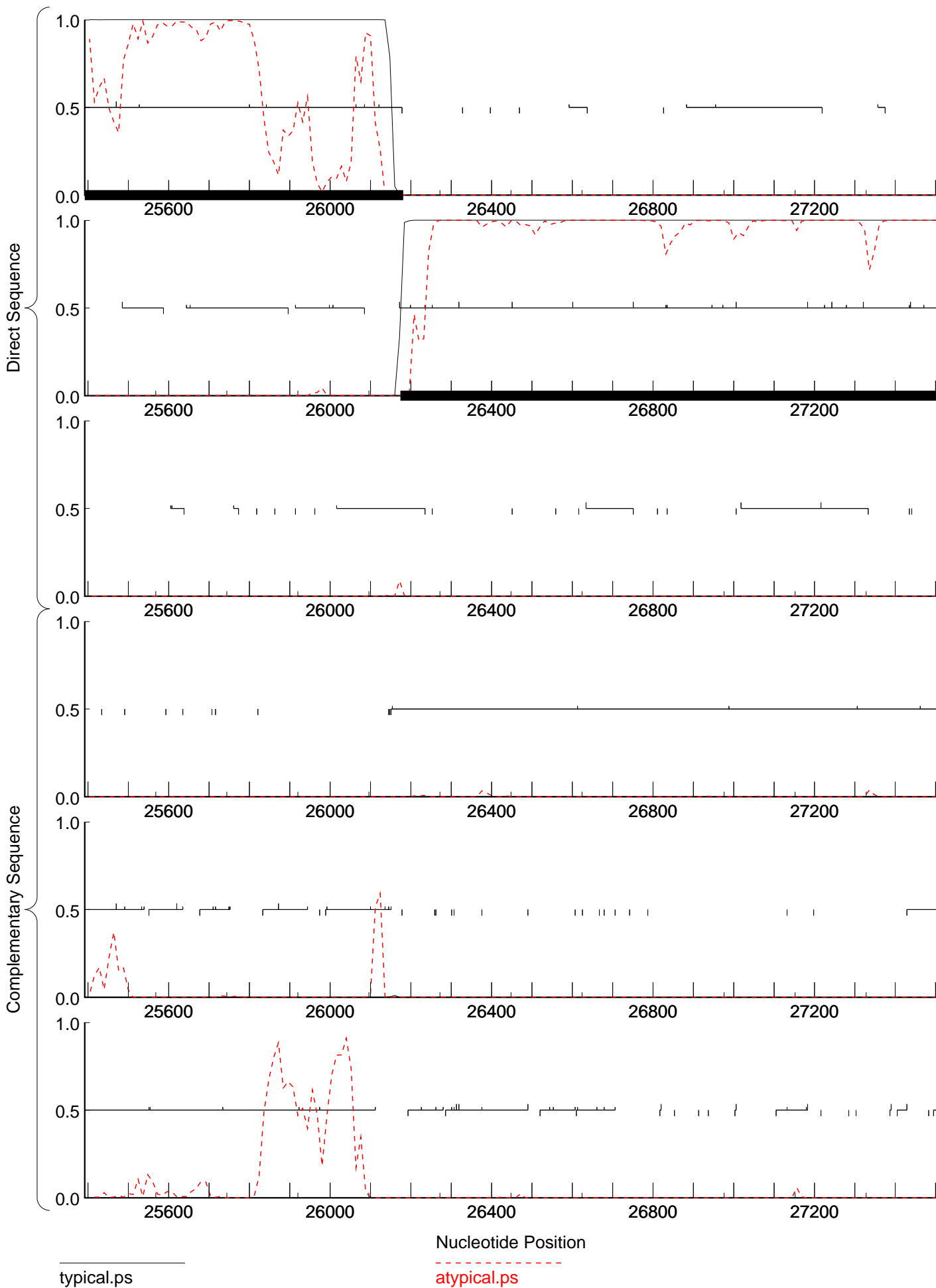


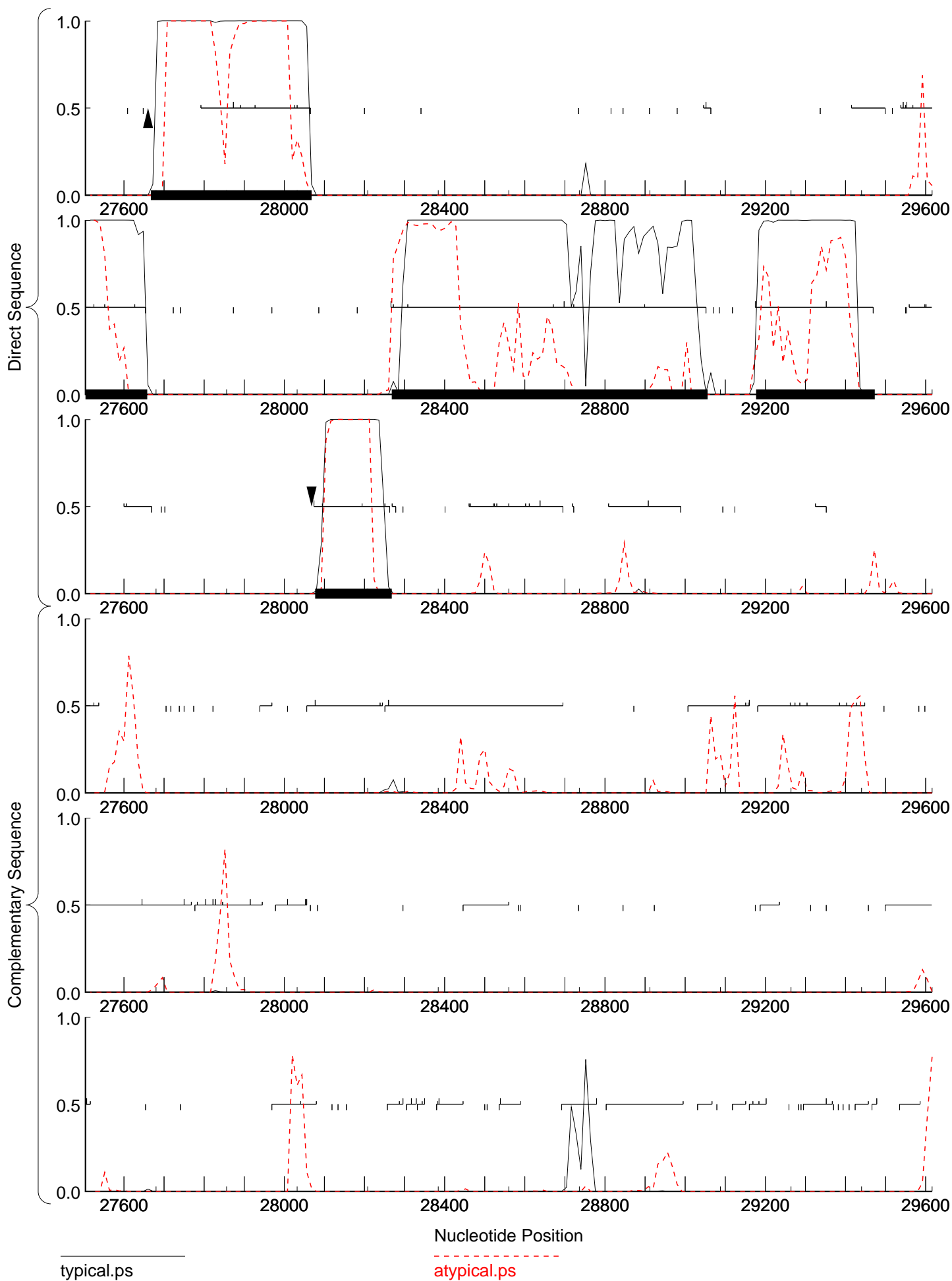






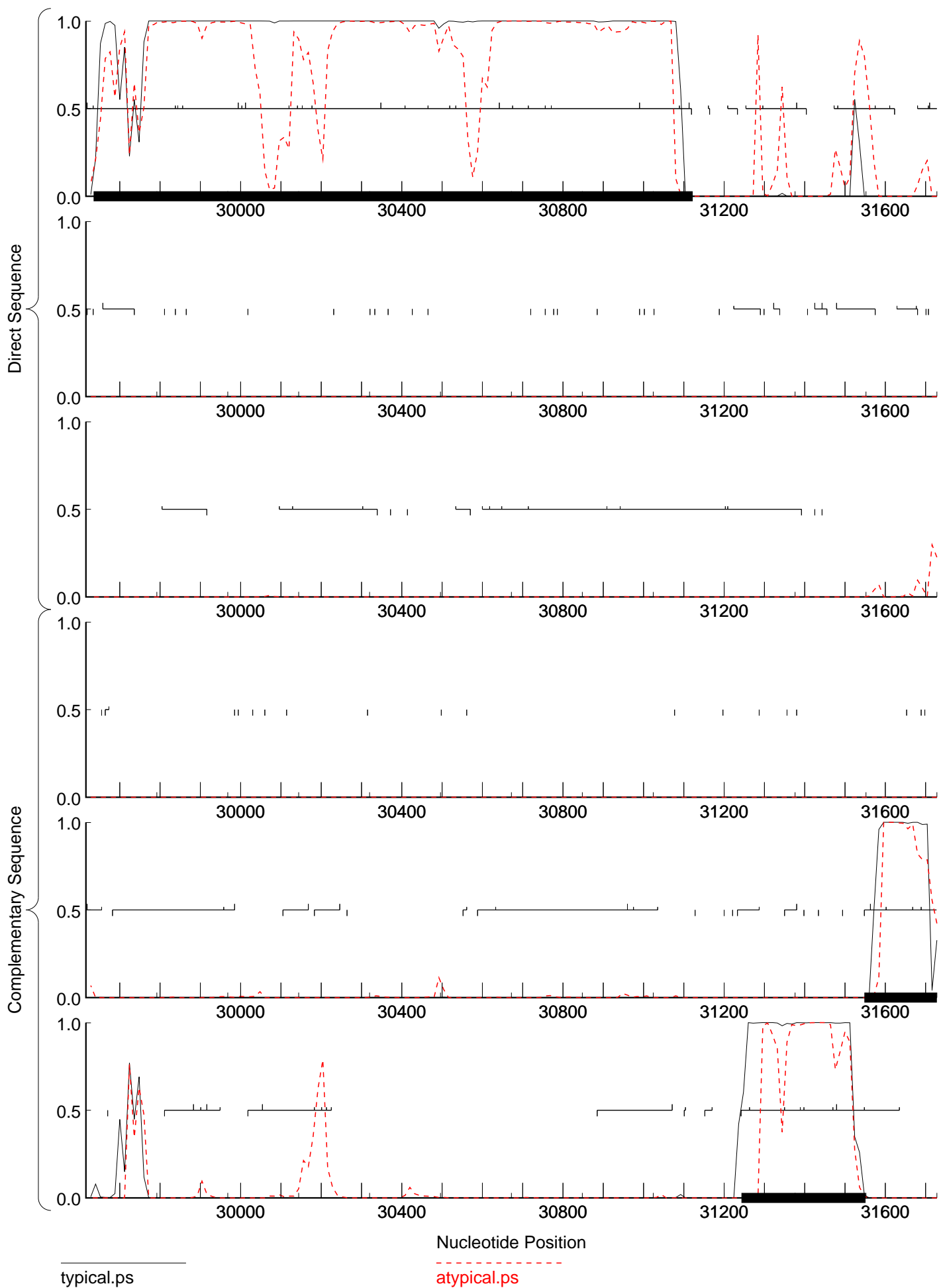




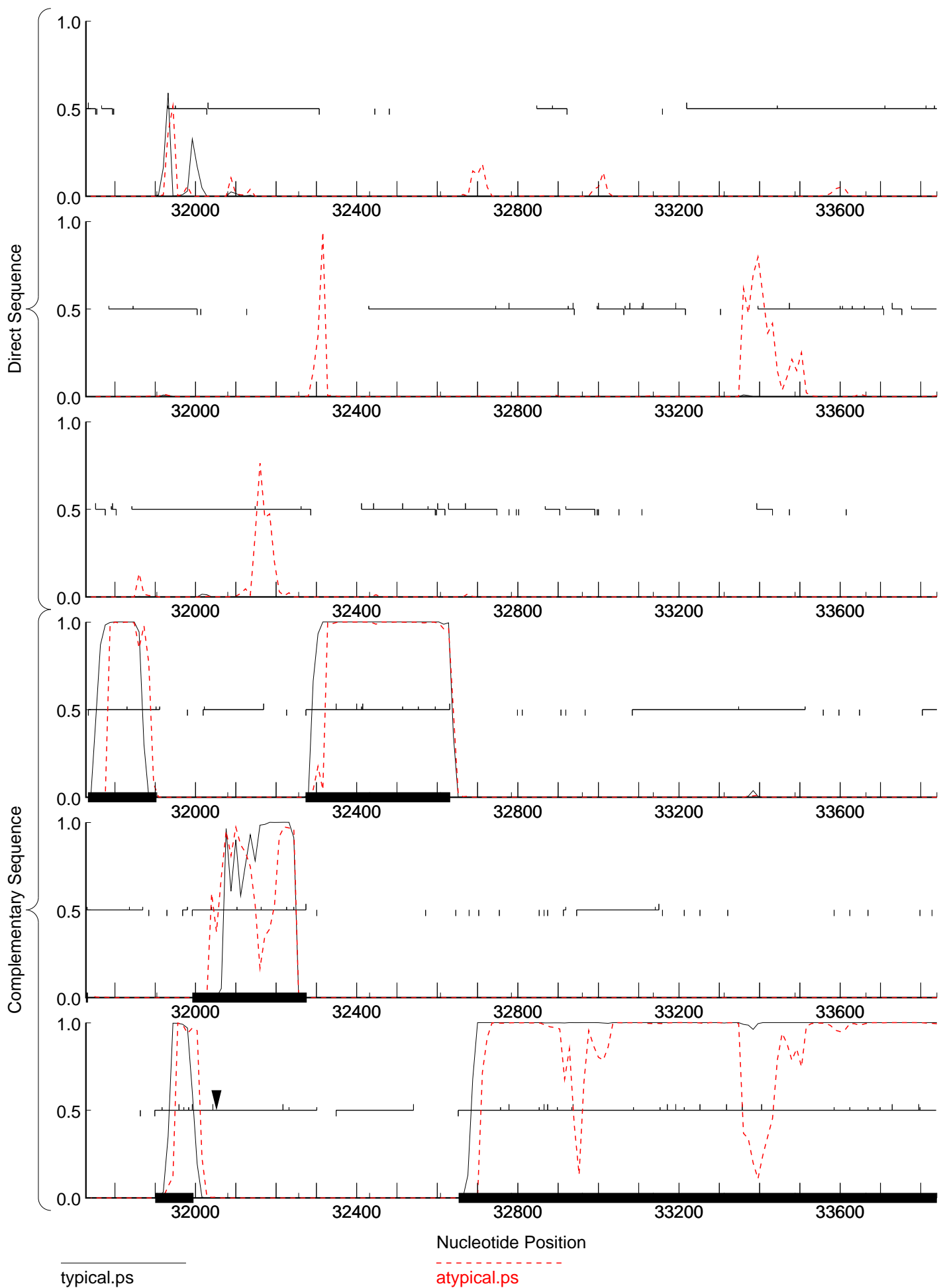


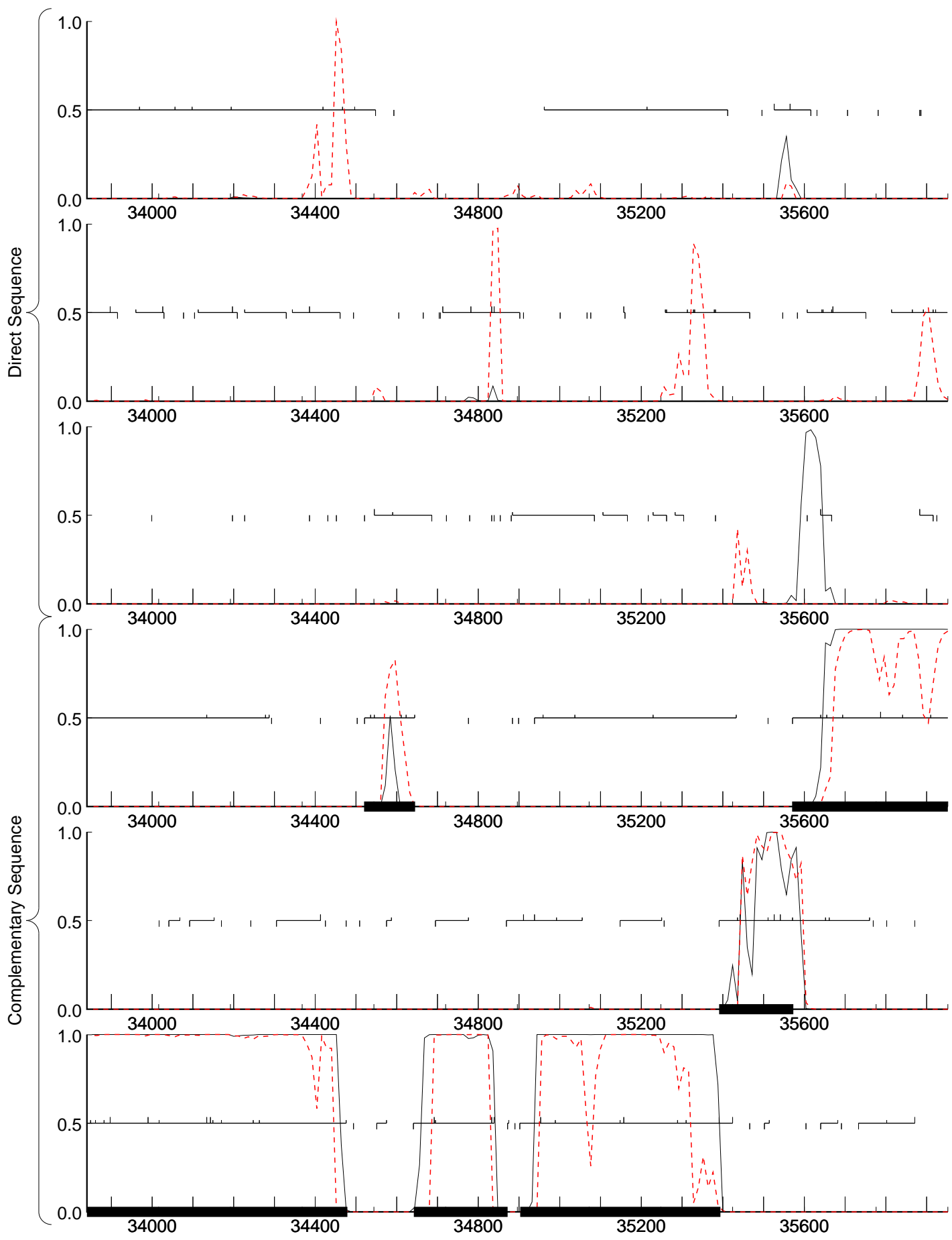
typical.ps

atypical.ps



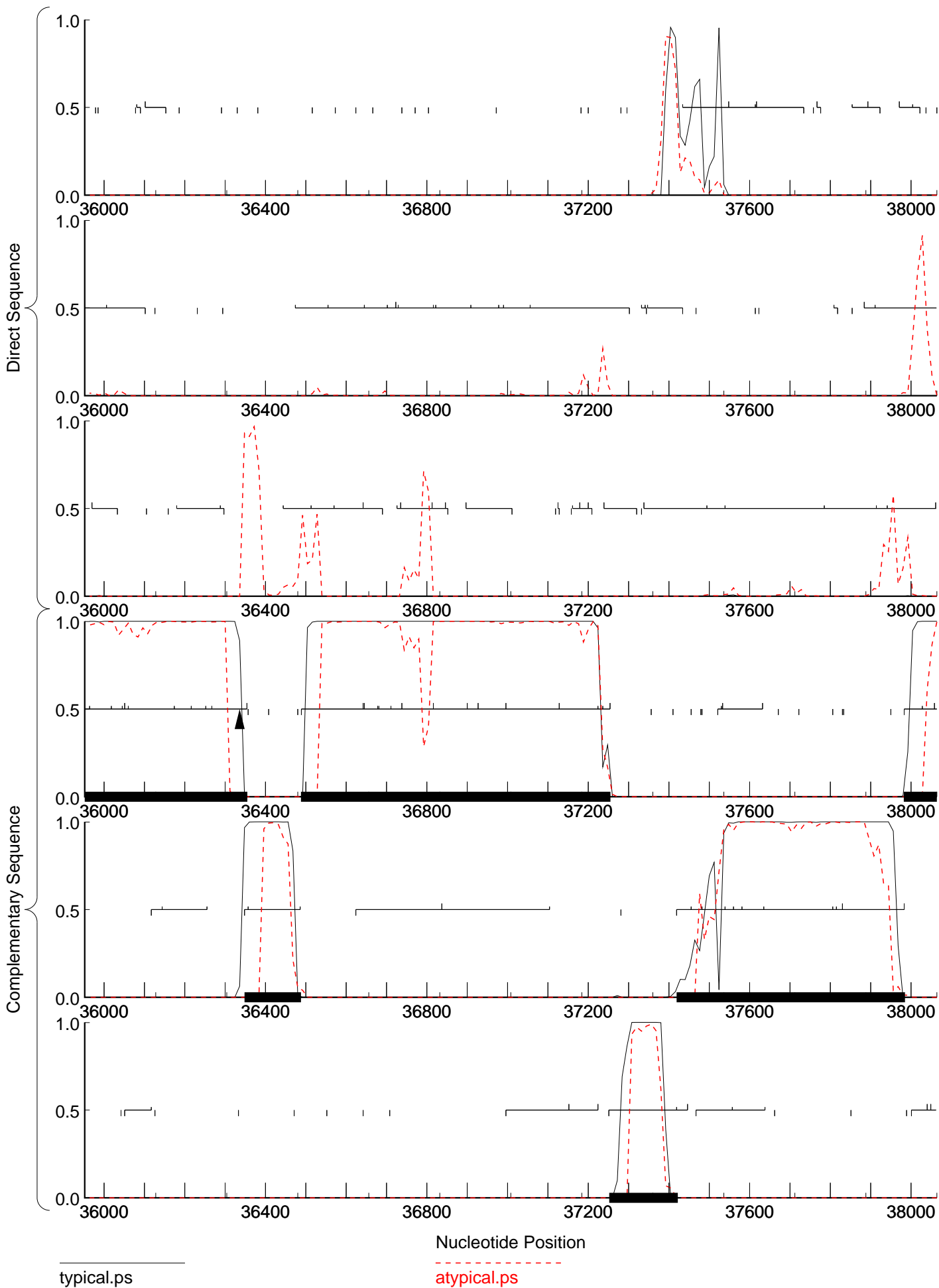


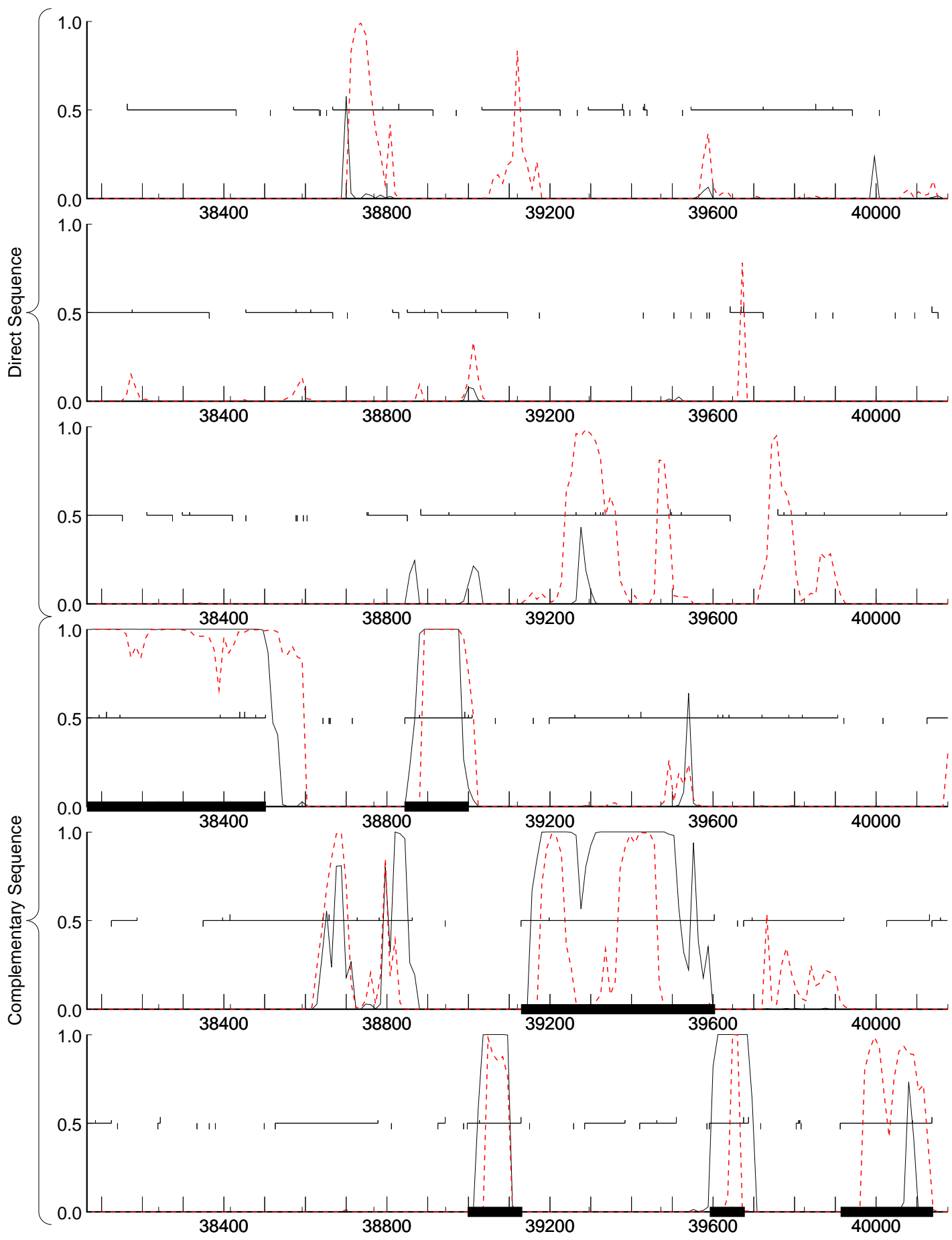




typical.ps

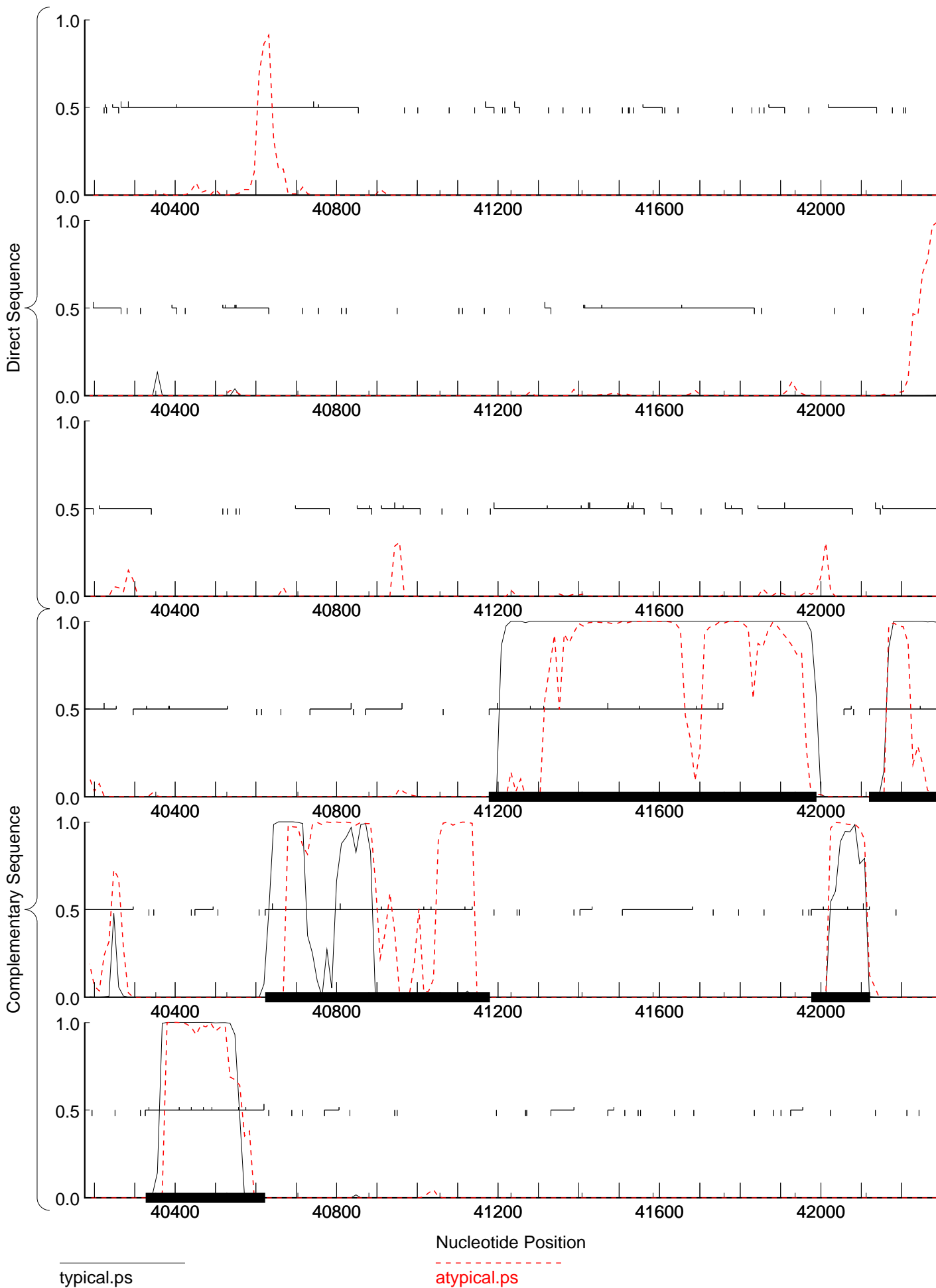
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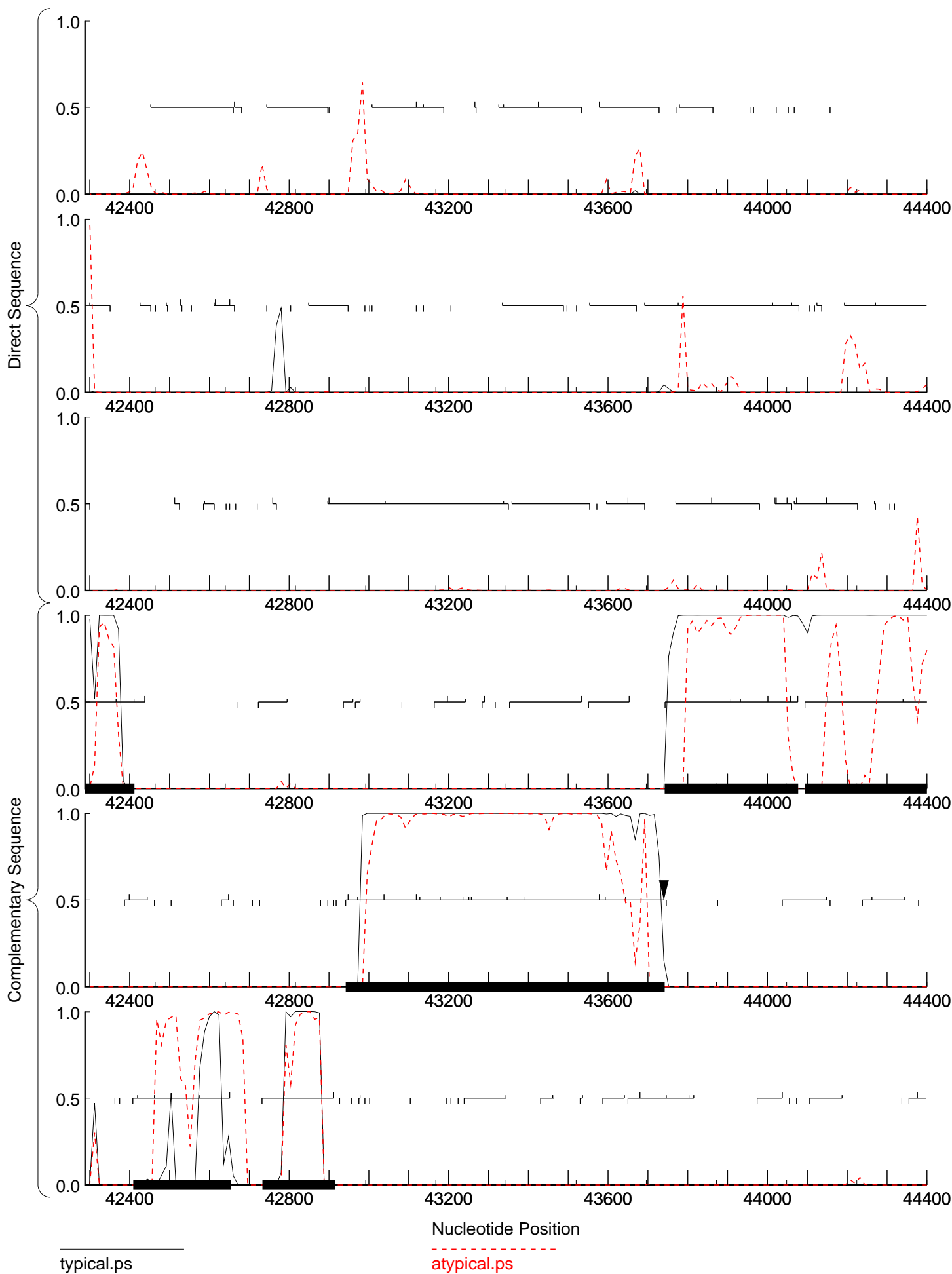


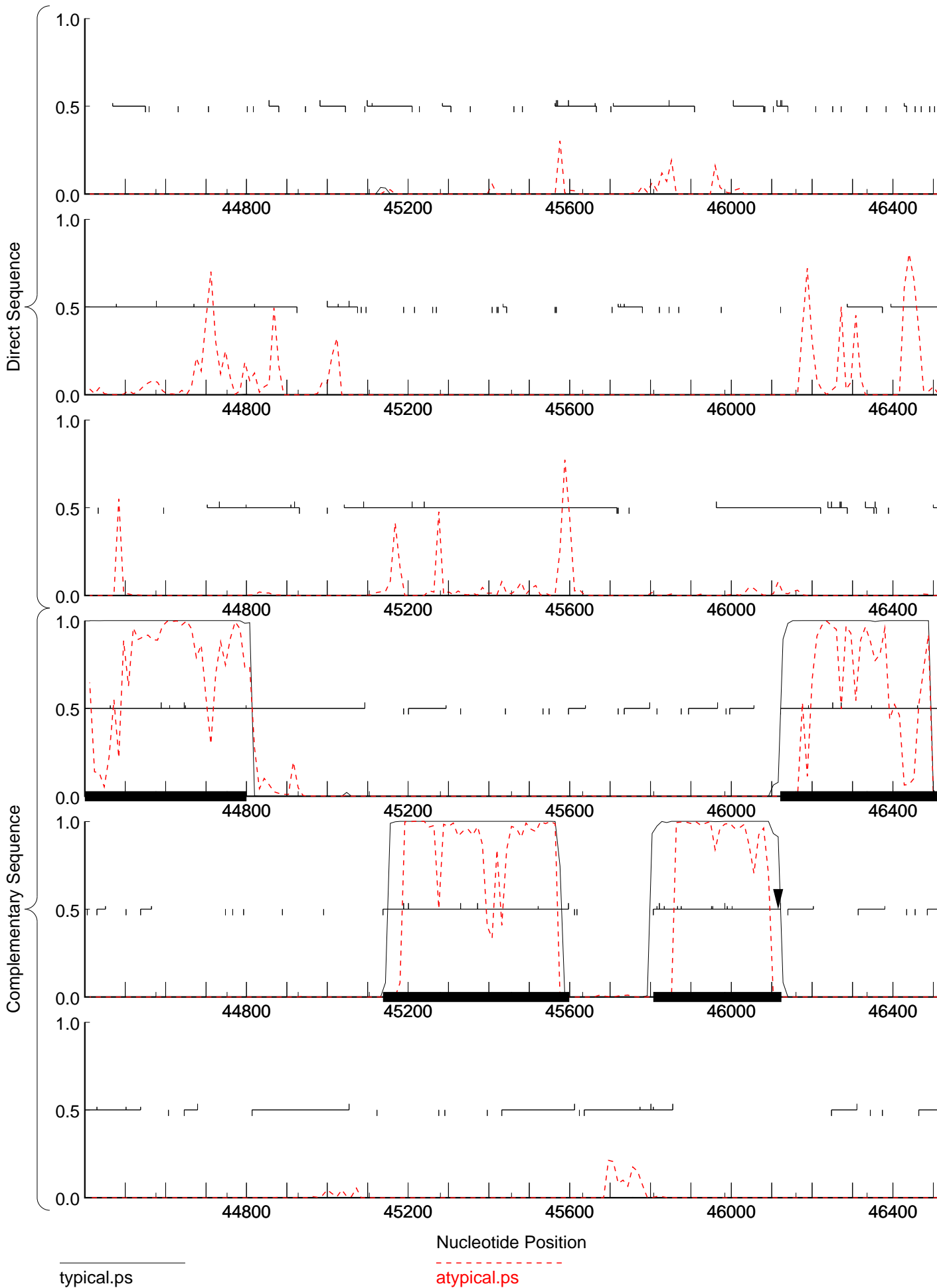


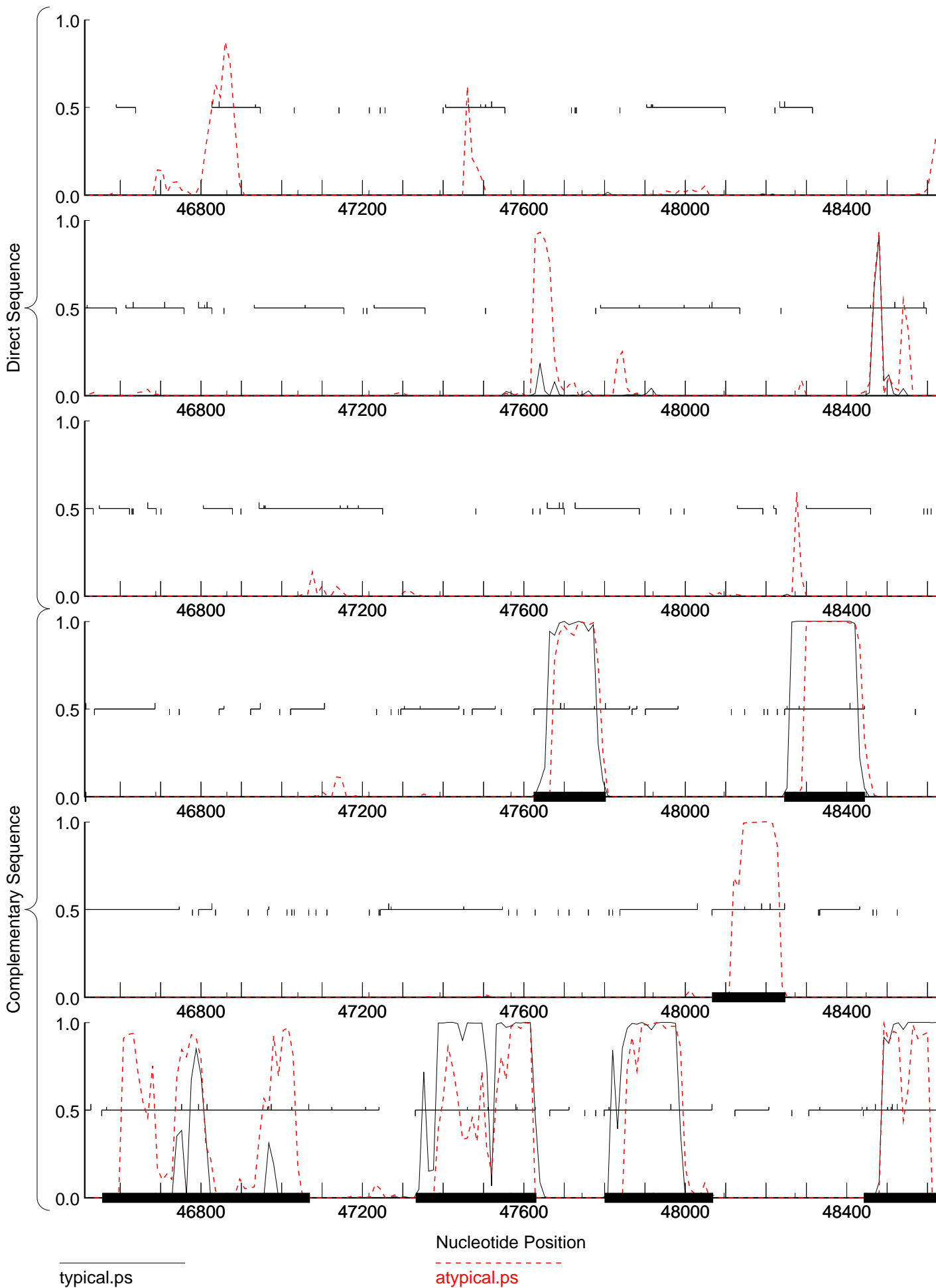
typical.ps

atypical.ps

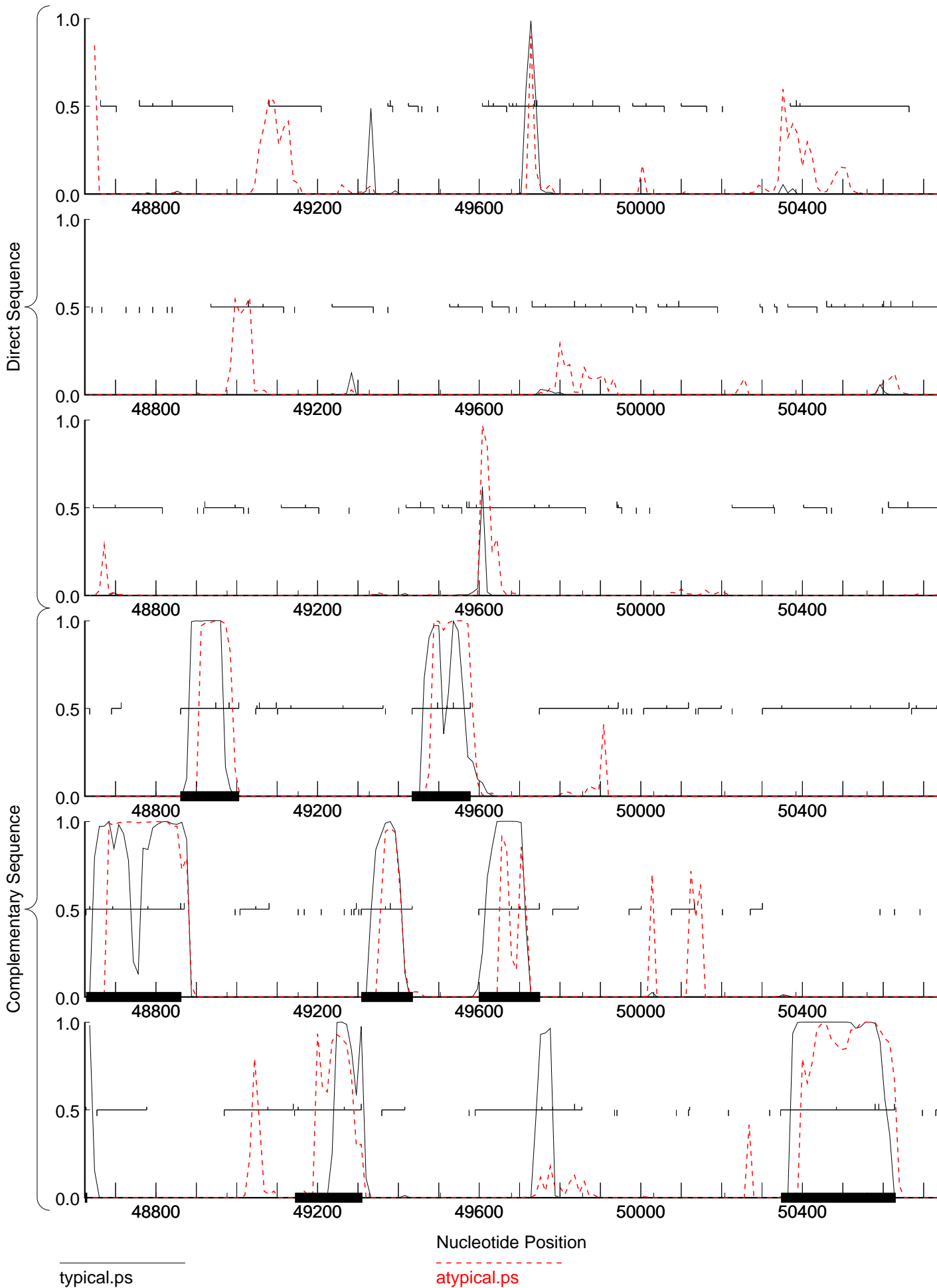












typical.ps

atypical.ps

