

# GeneMark

Version 2.5p (09.08.06)

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## PROGRAM INFORMATION

Sequence : Gordonia phage Zirinka complete sequence, 52077 bp including 10-base 3' overhang (CGGCTGGGG  
Analysis Date : 6/23/18 at 8:14:48  
Pages : 26  
Sequence Length : 52077 bp  
GC Content : 66.67%

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,

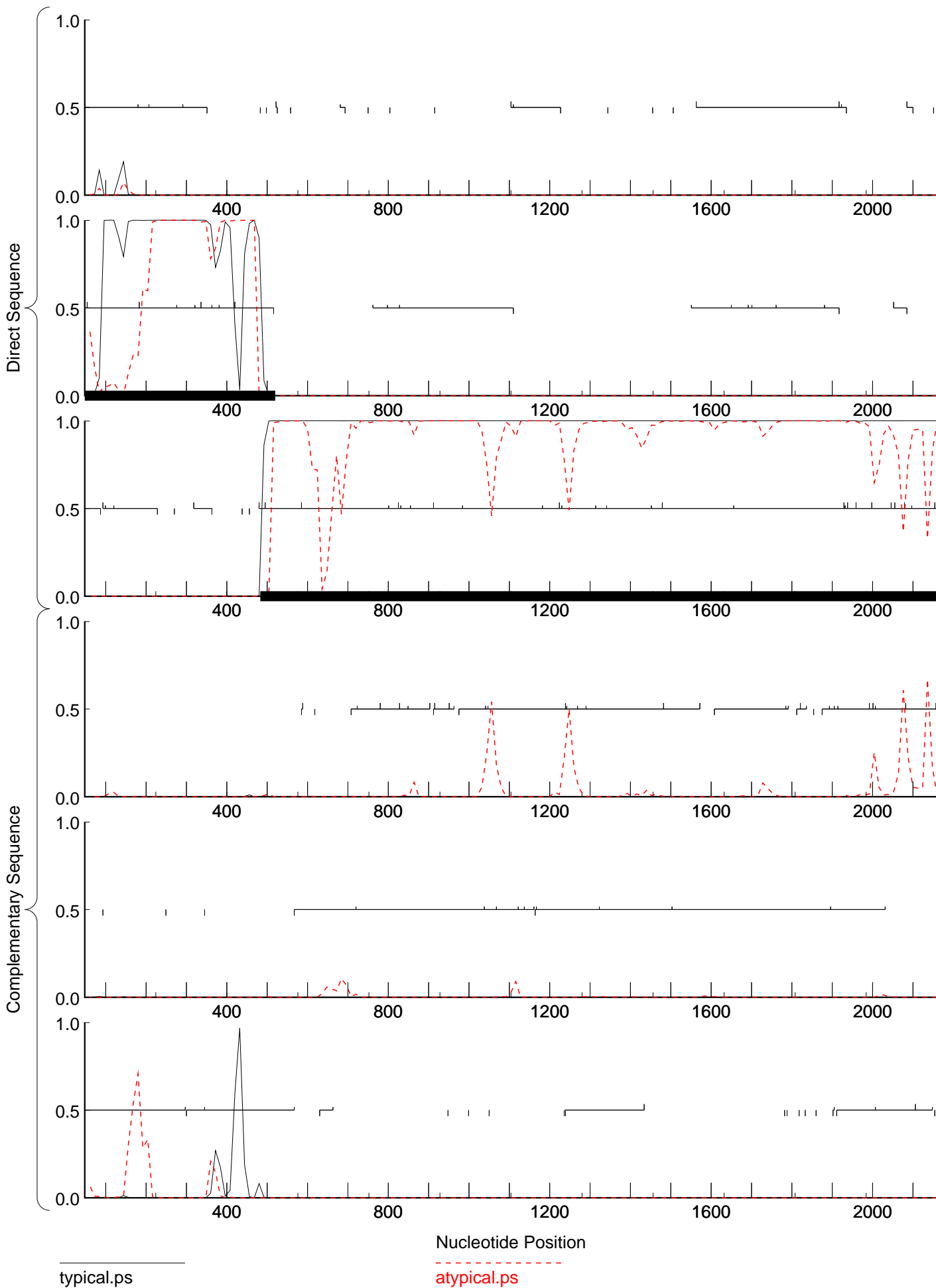
## MATRIX INFORMATION

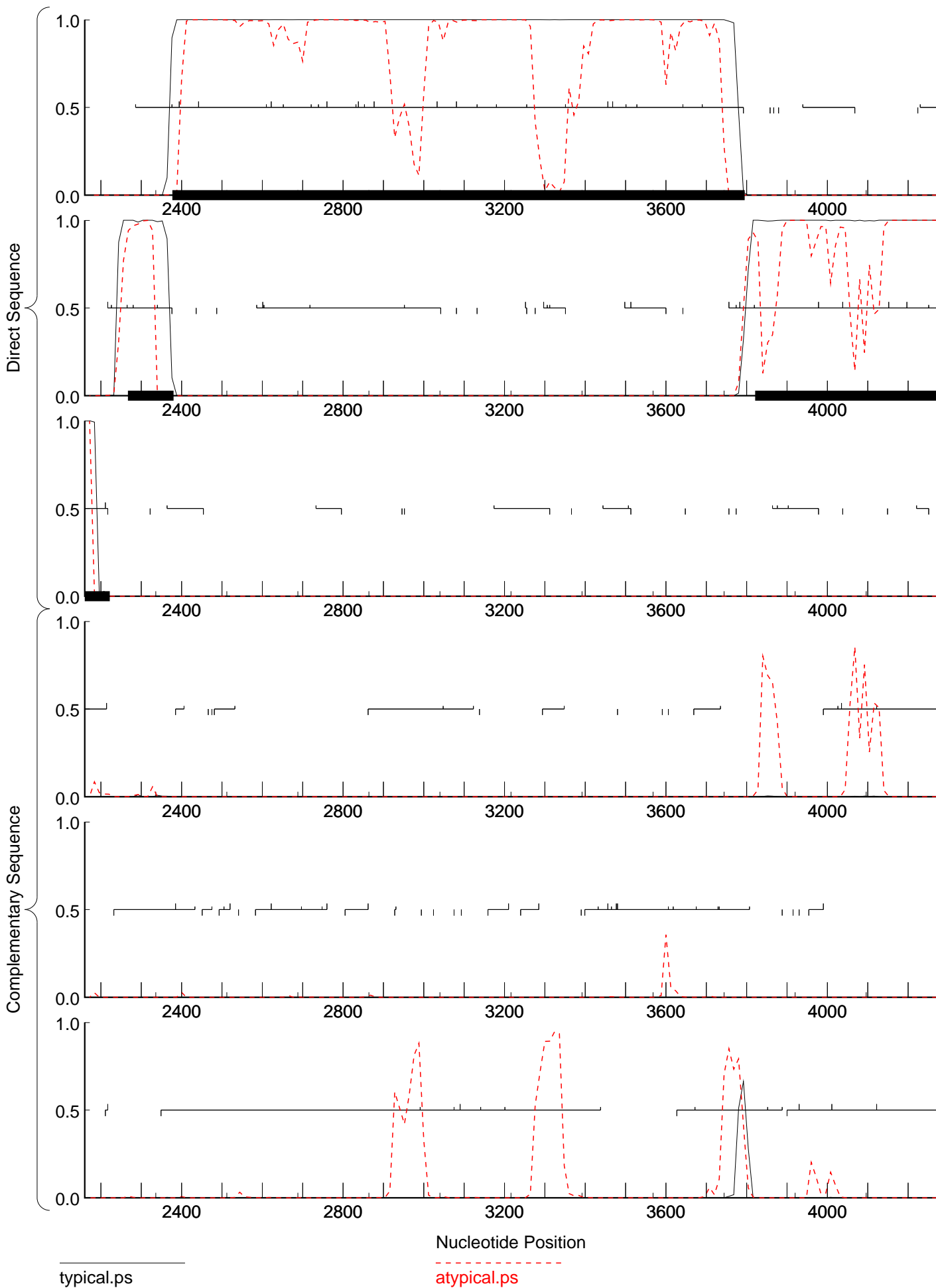
Matrix : -  
Author : -  
Order : 4

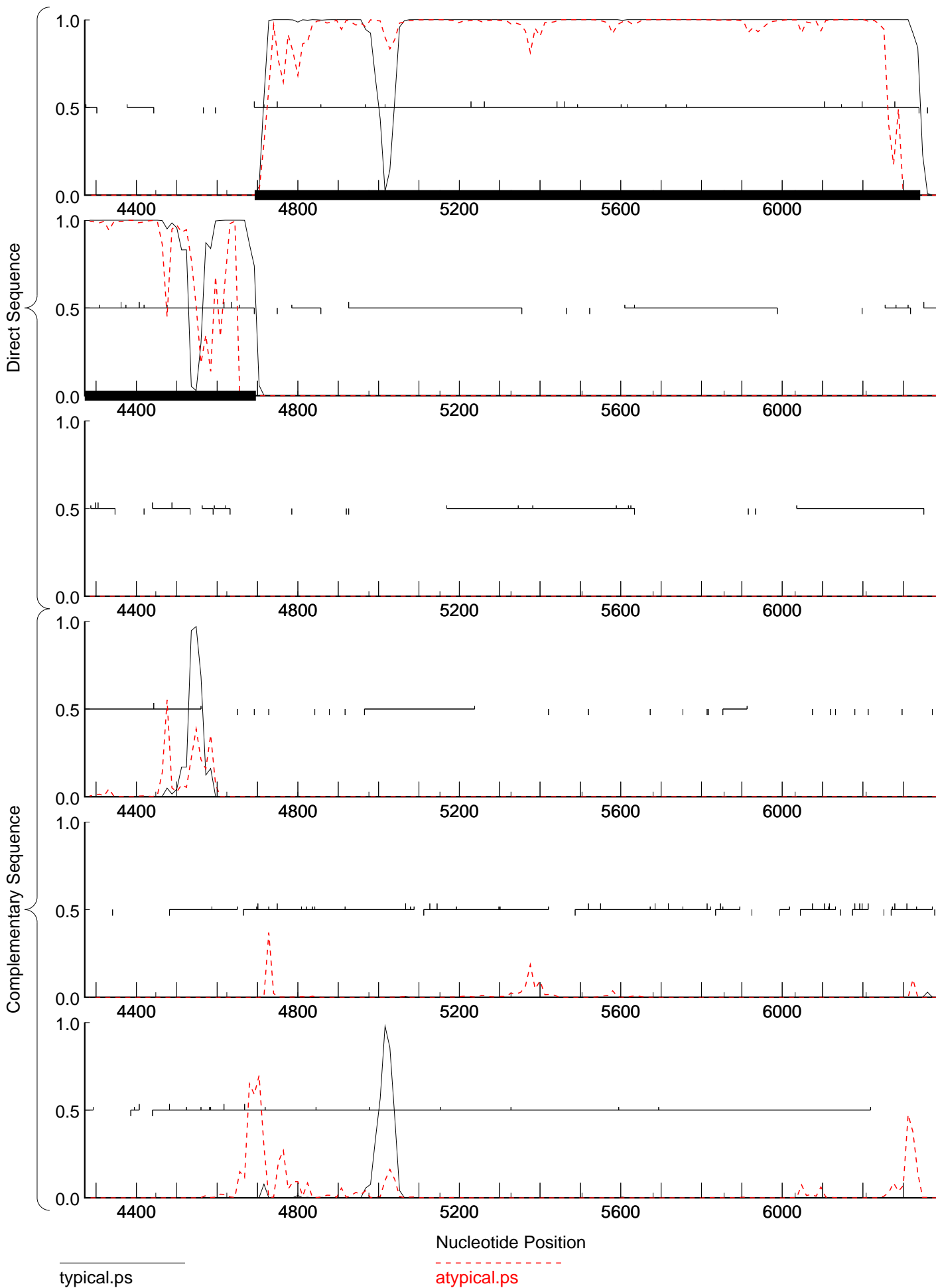
Send questions / comments to:  
Dr. M. Borodovsky  
Georgia Institute of Technology  
School of Biology  
Atlanta, GA 30332-0230

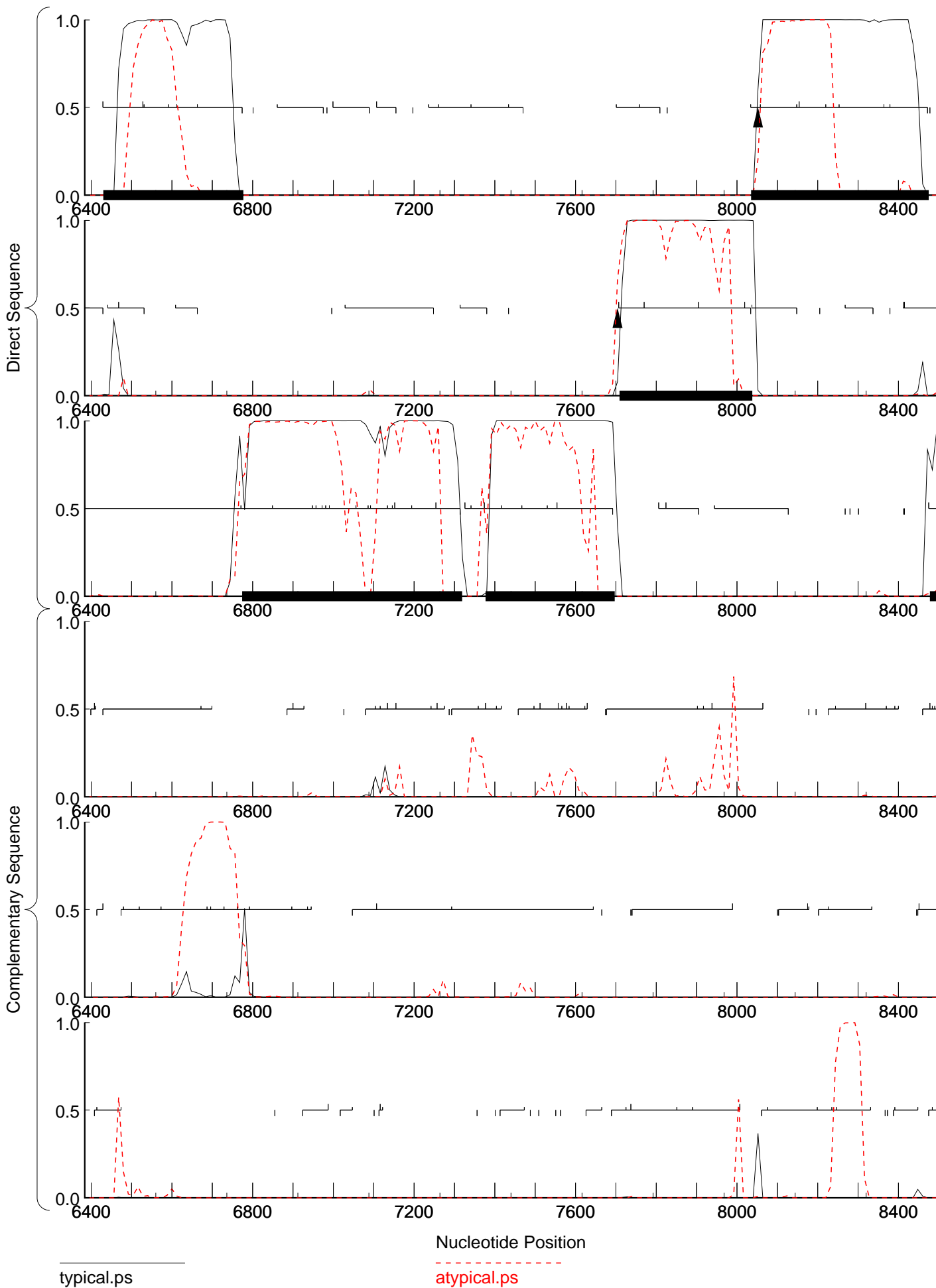
### Matrix notes & comments

Training set derived by GeneMarkS, 4.30 October 2014  
Sat Jun 23 08:14:48 2018

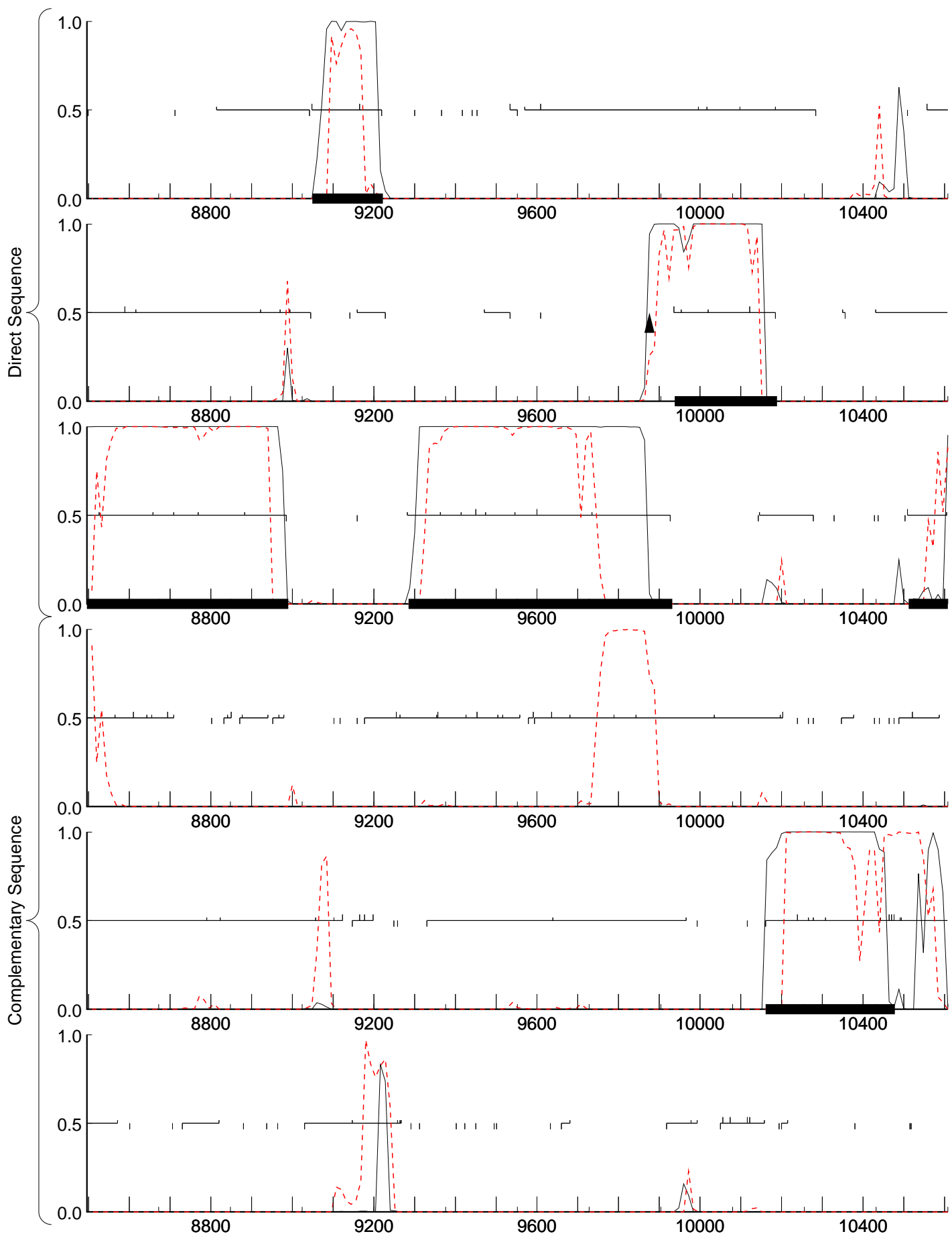






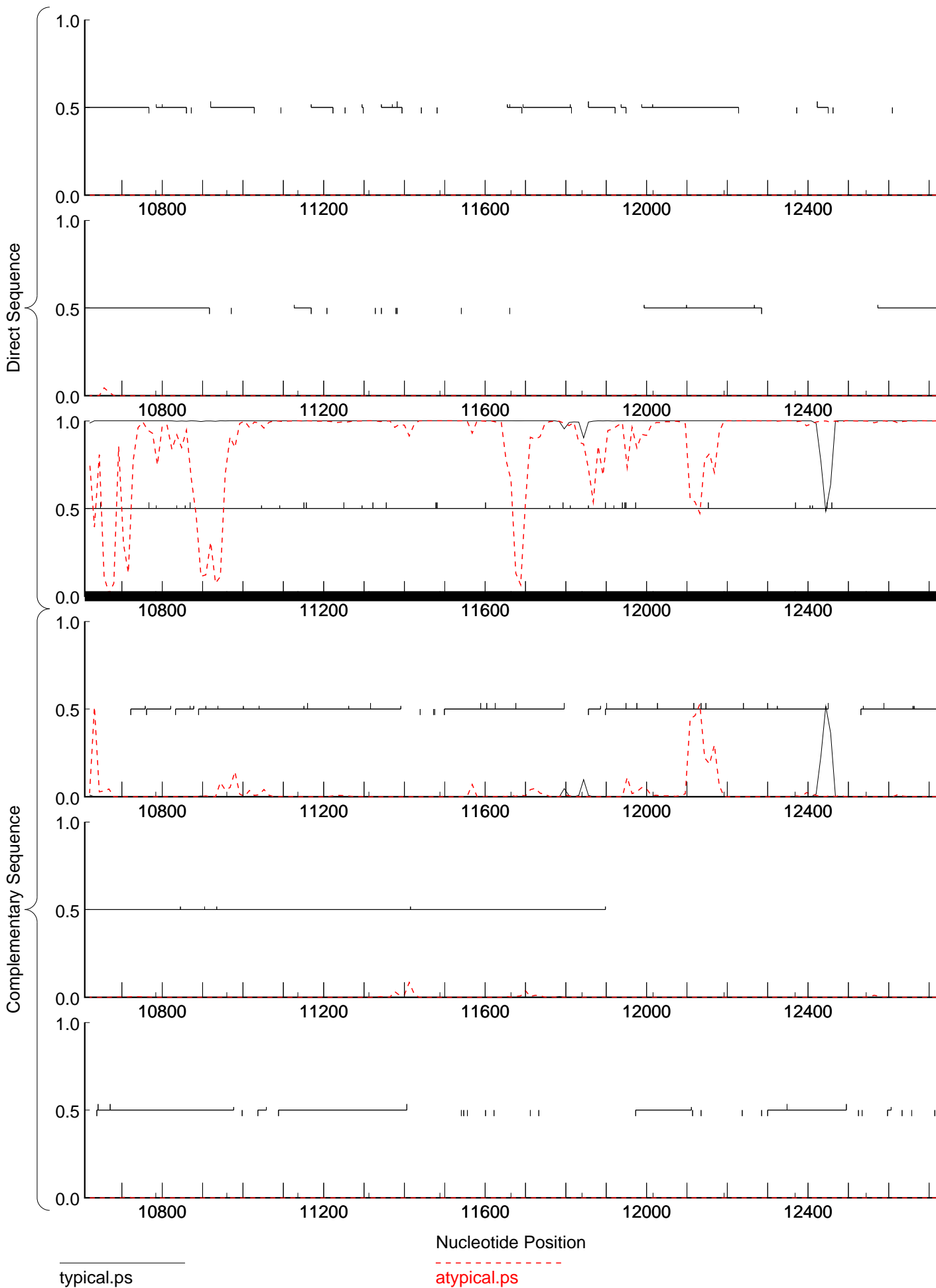


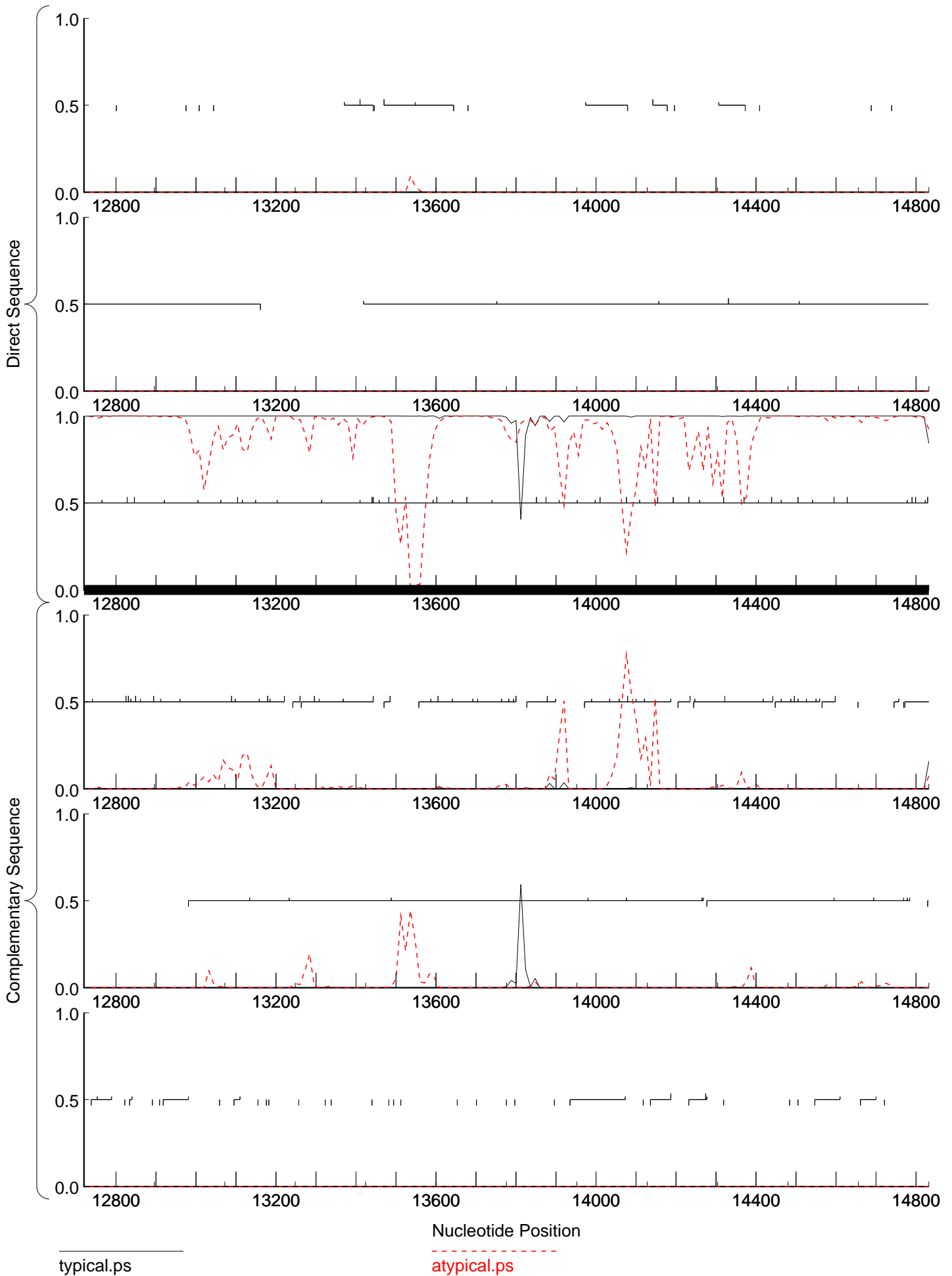
GeneMark.hmm prediction



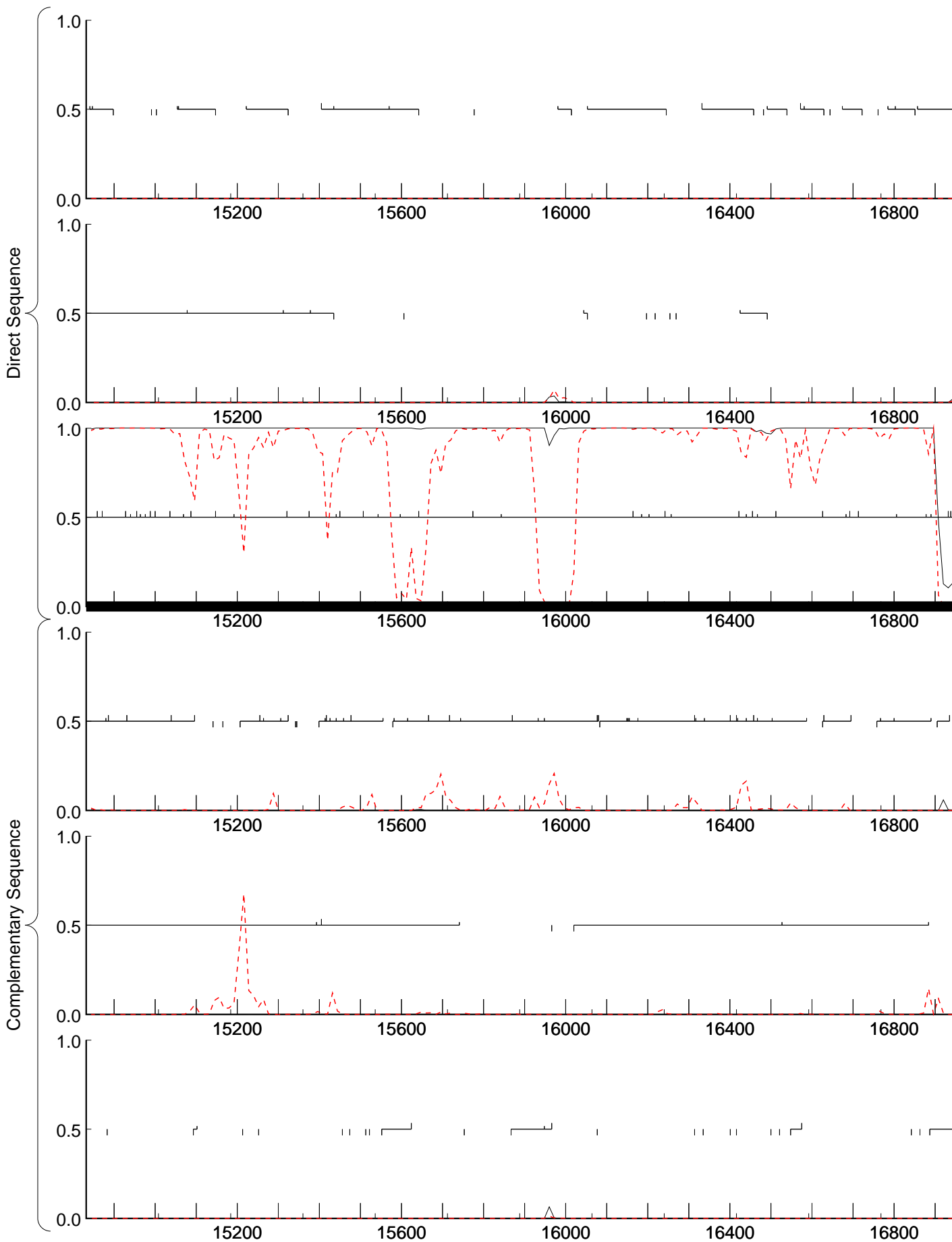
typical.ps

atypical.ps



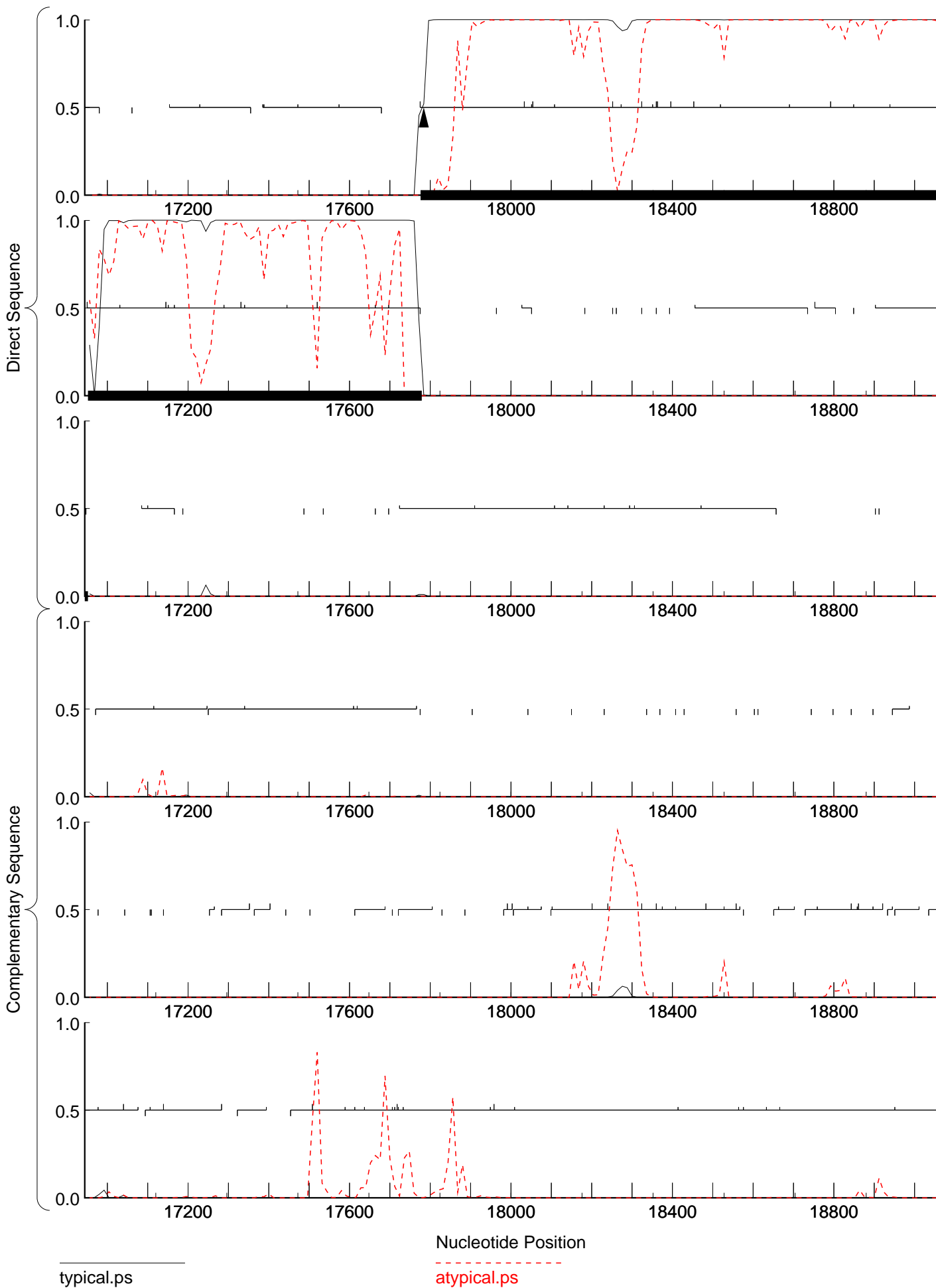


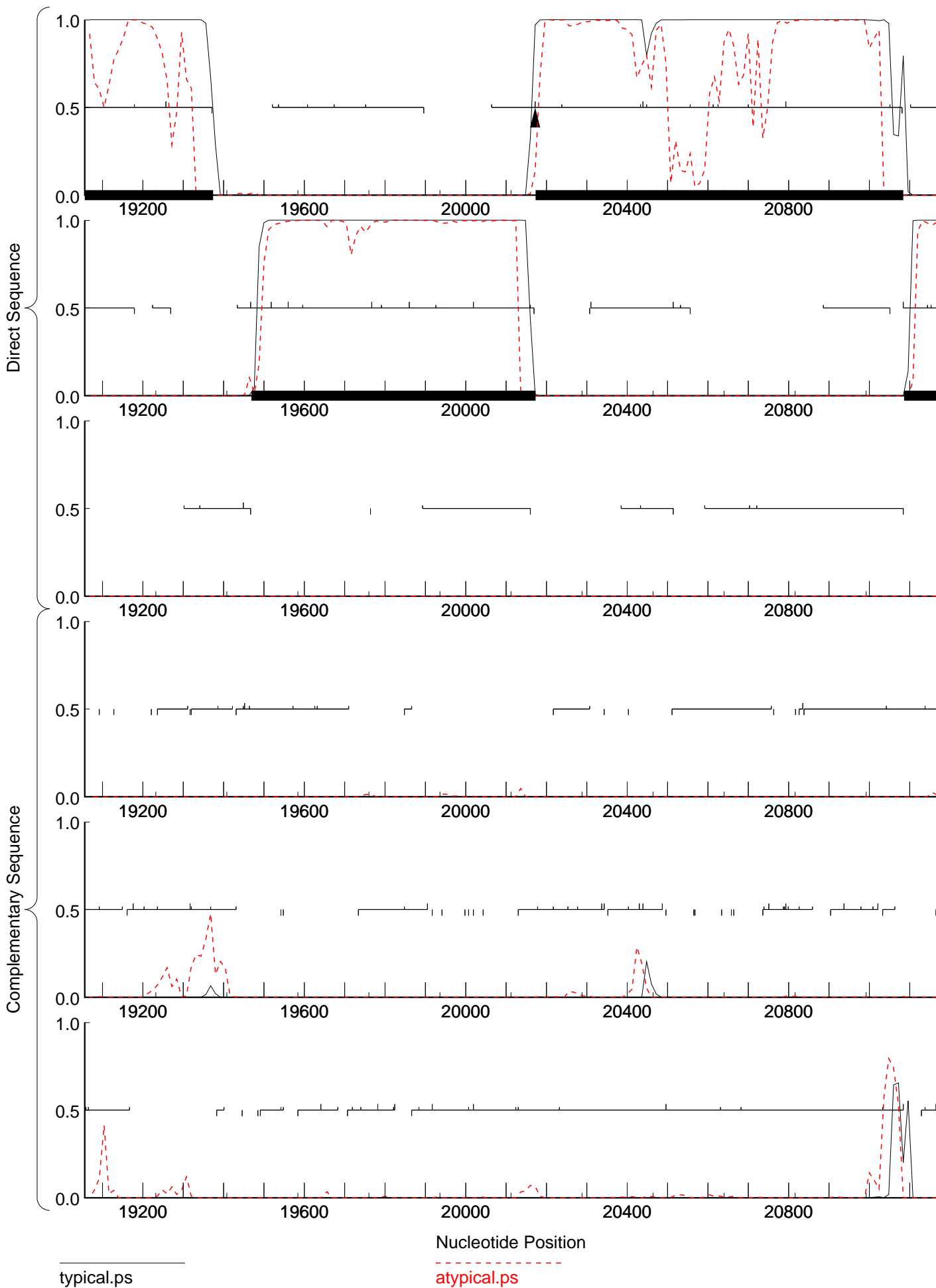




typical.ps

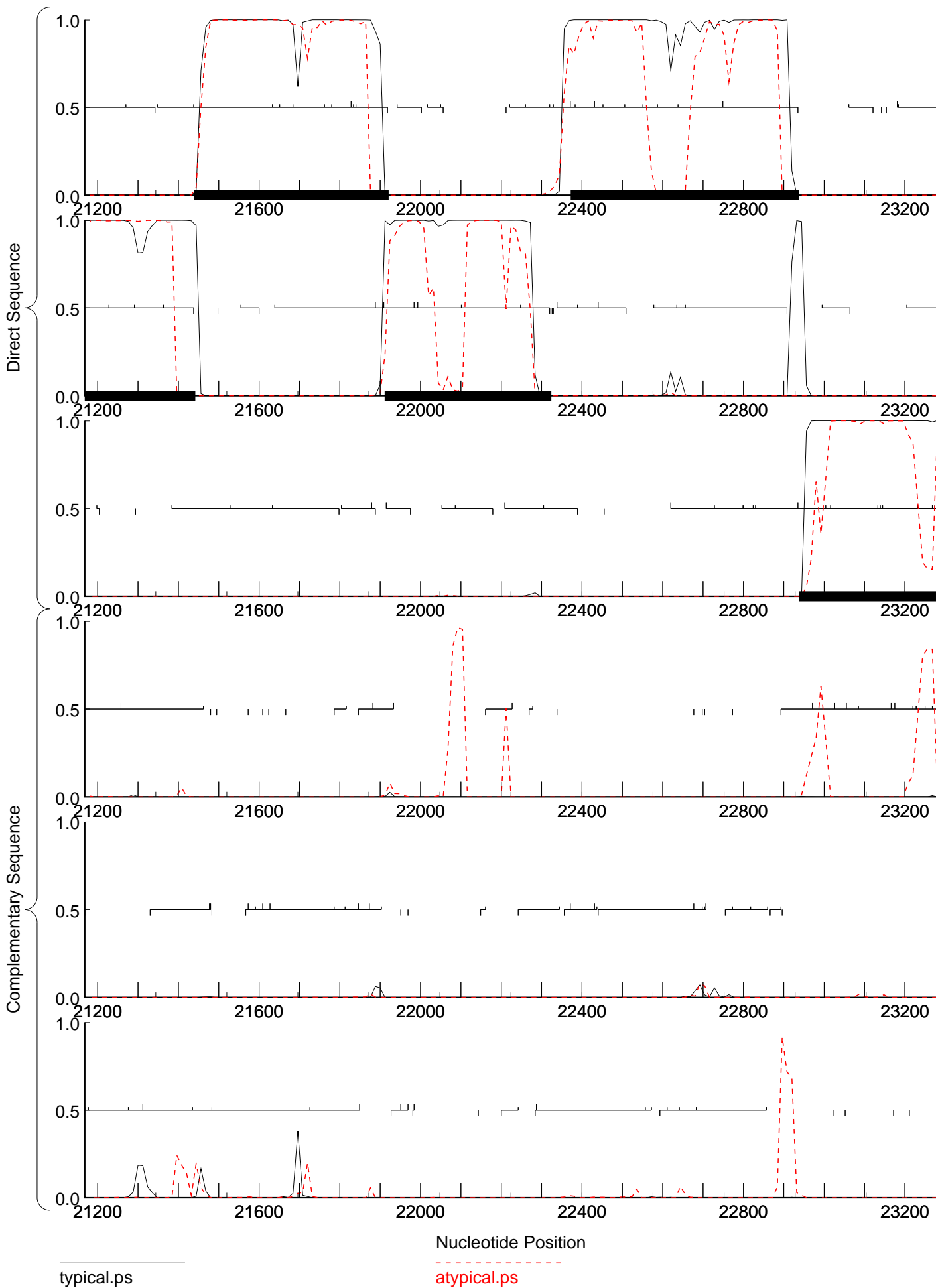
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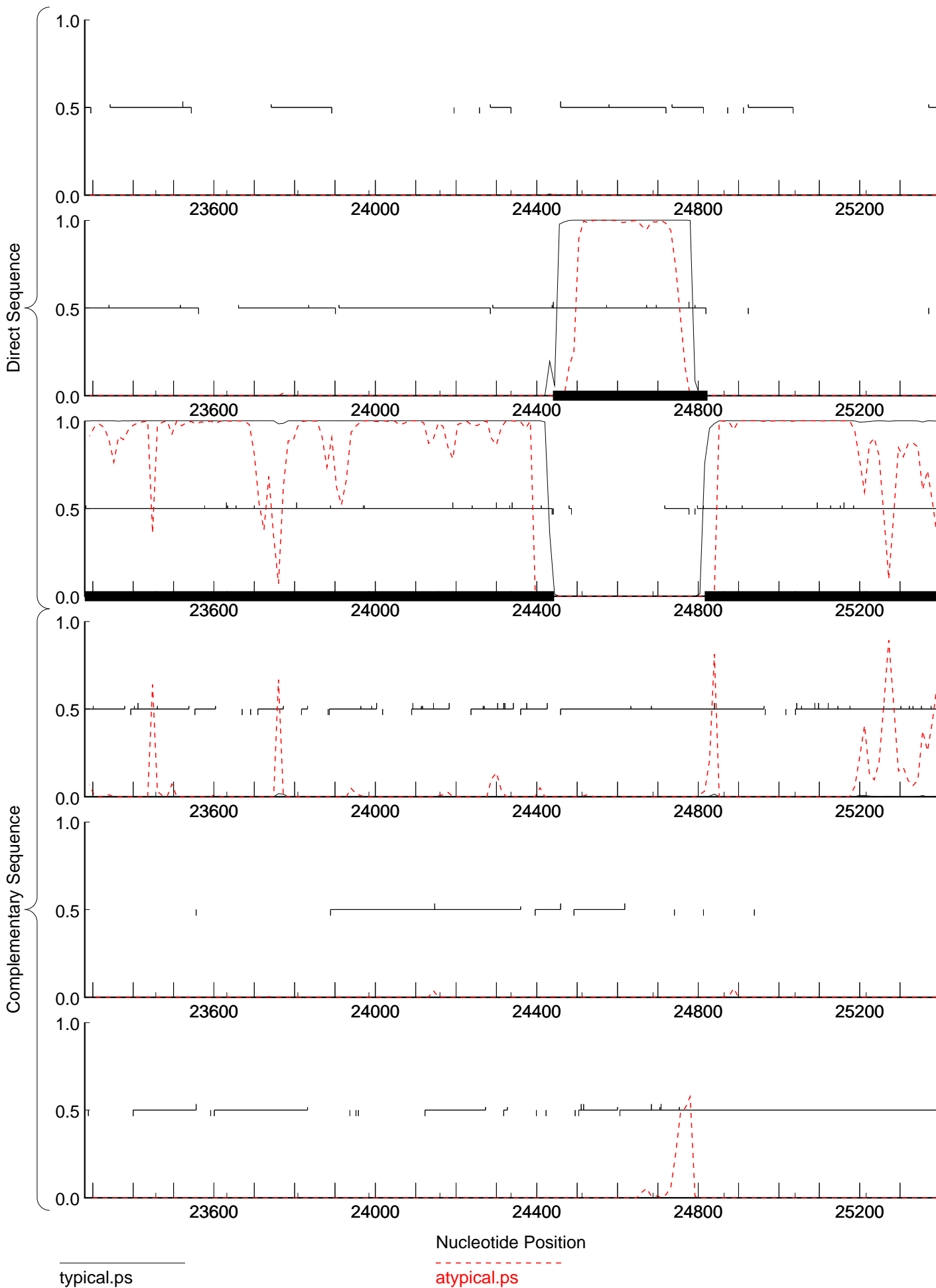




typical.ps

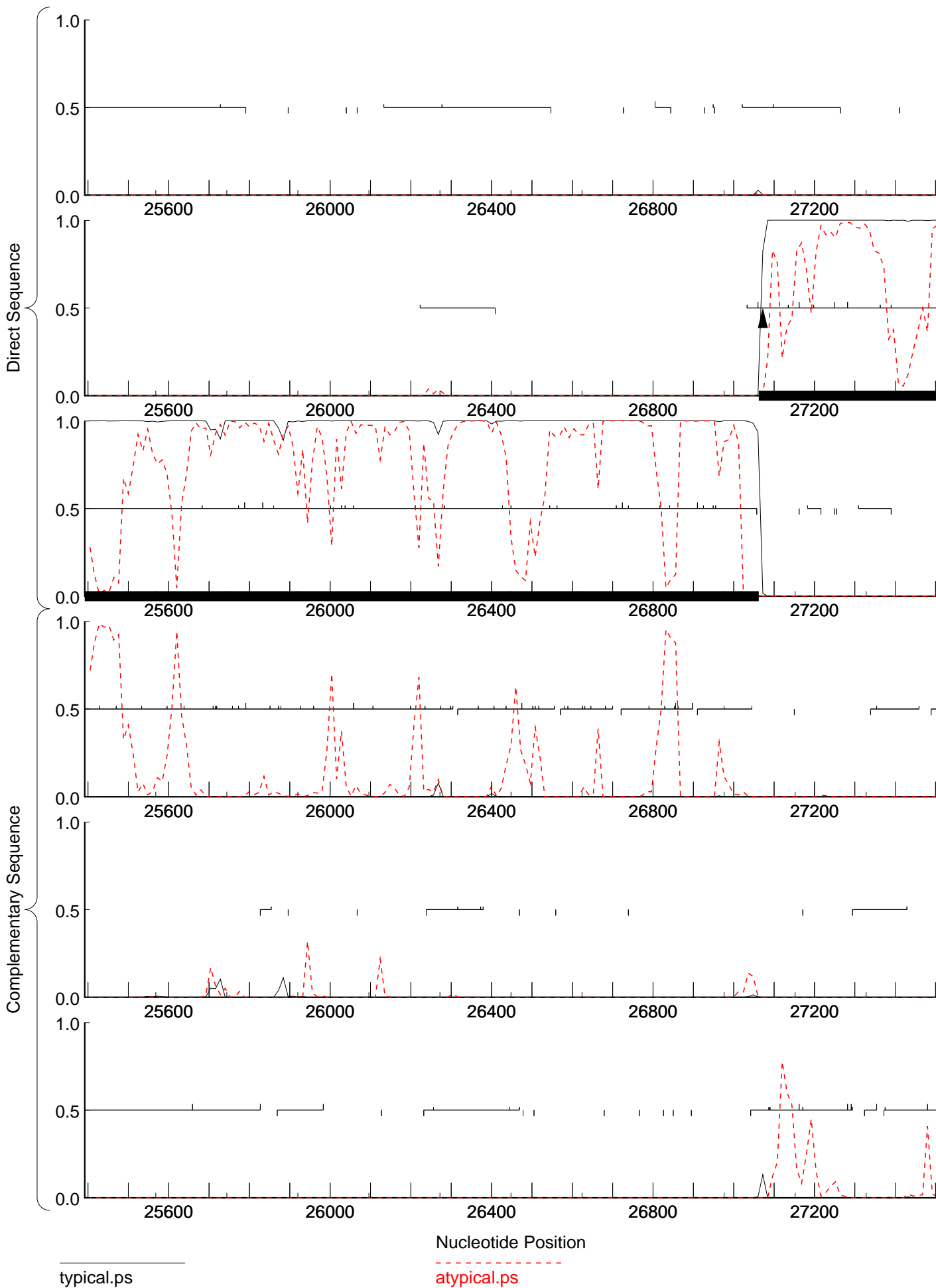
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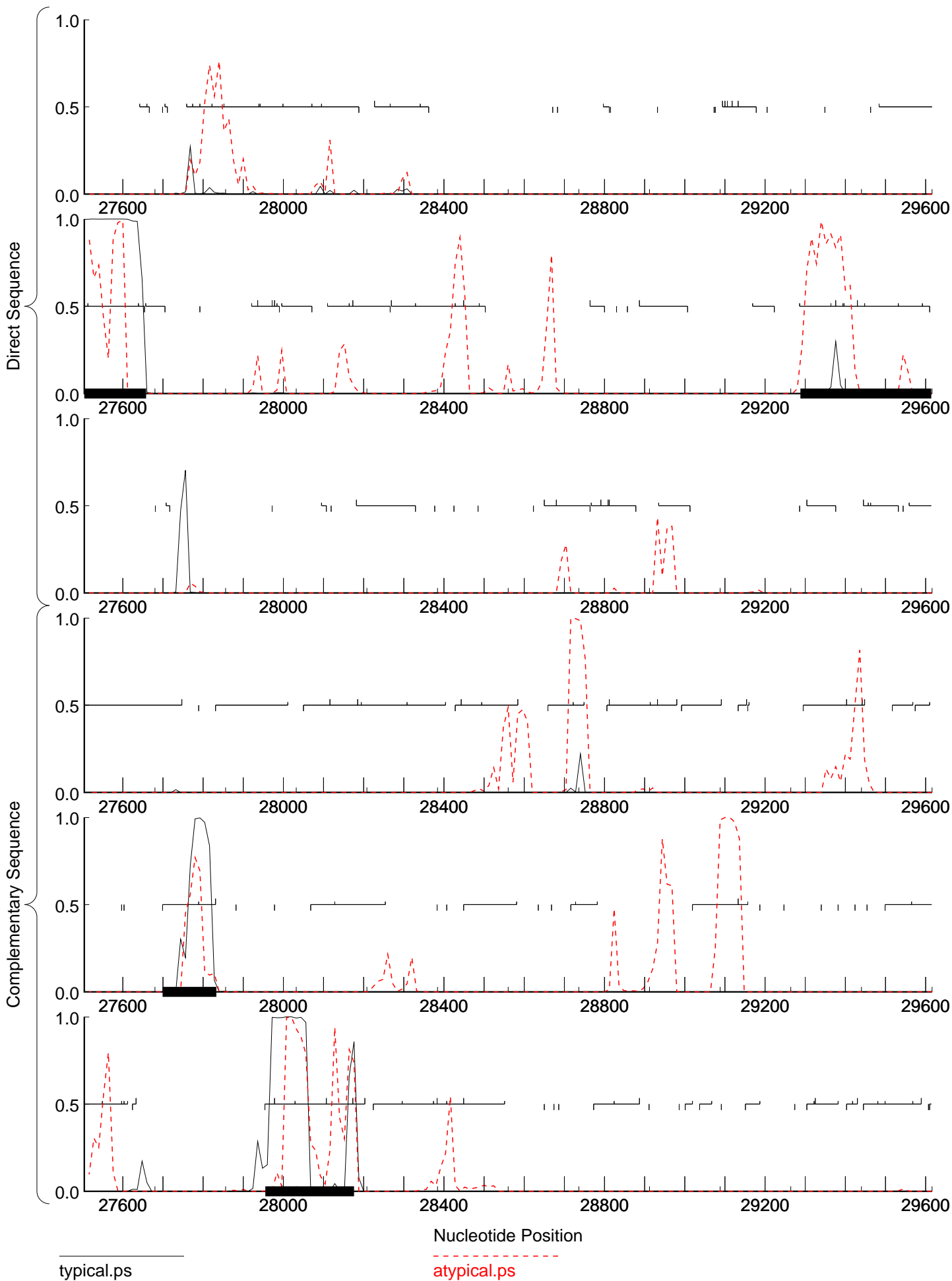


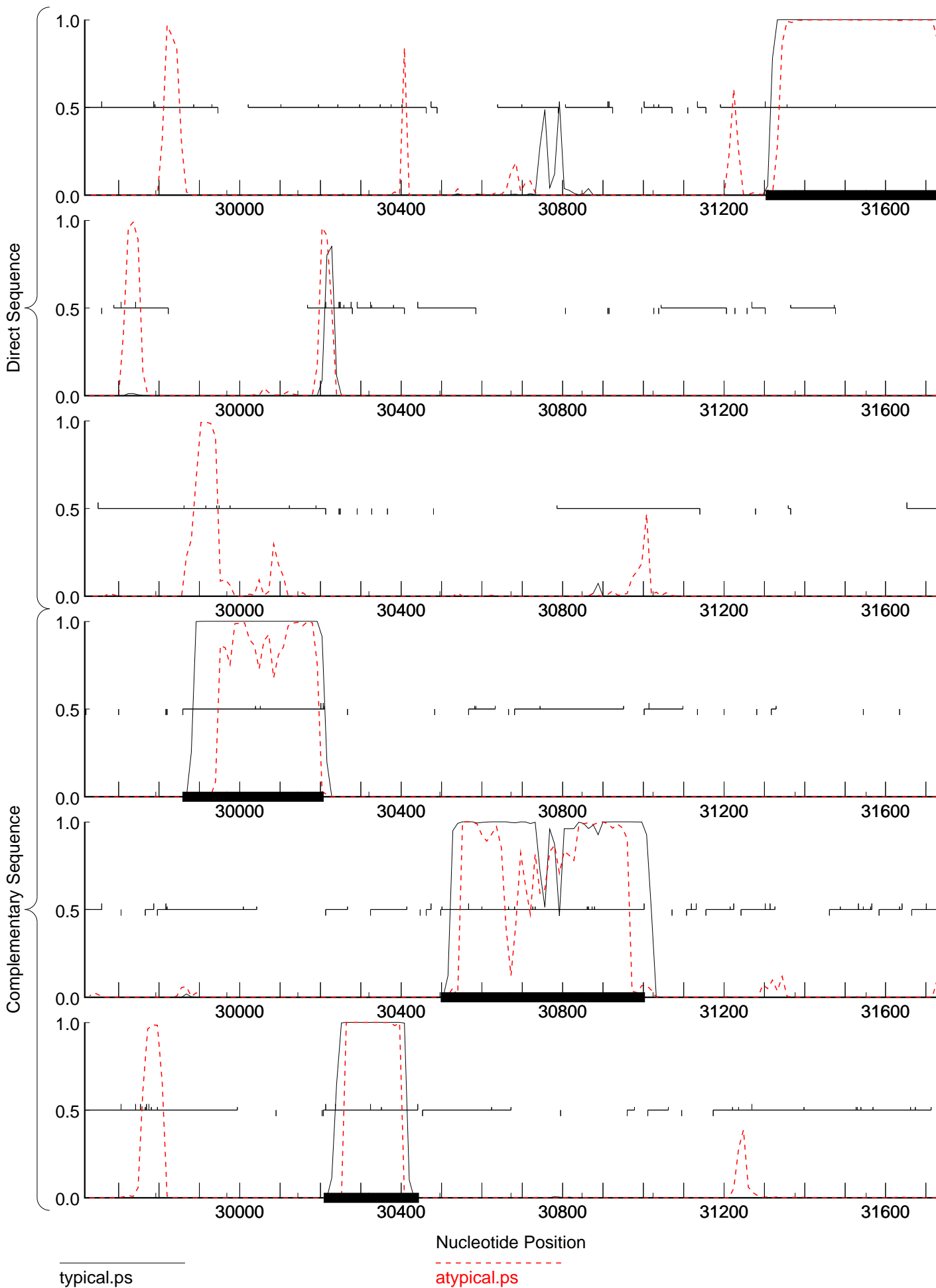


typical.ps

atypical.ps



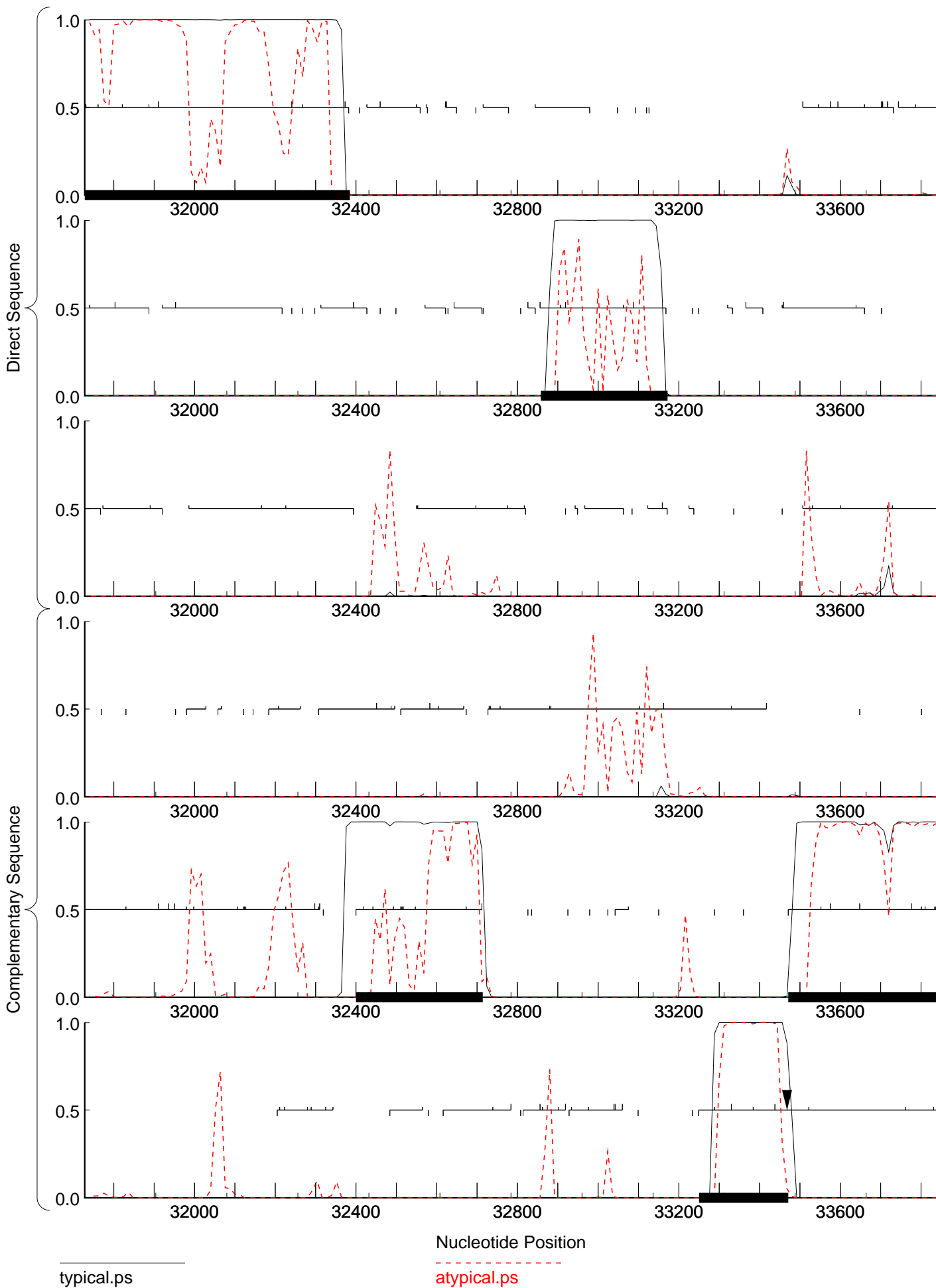


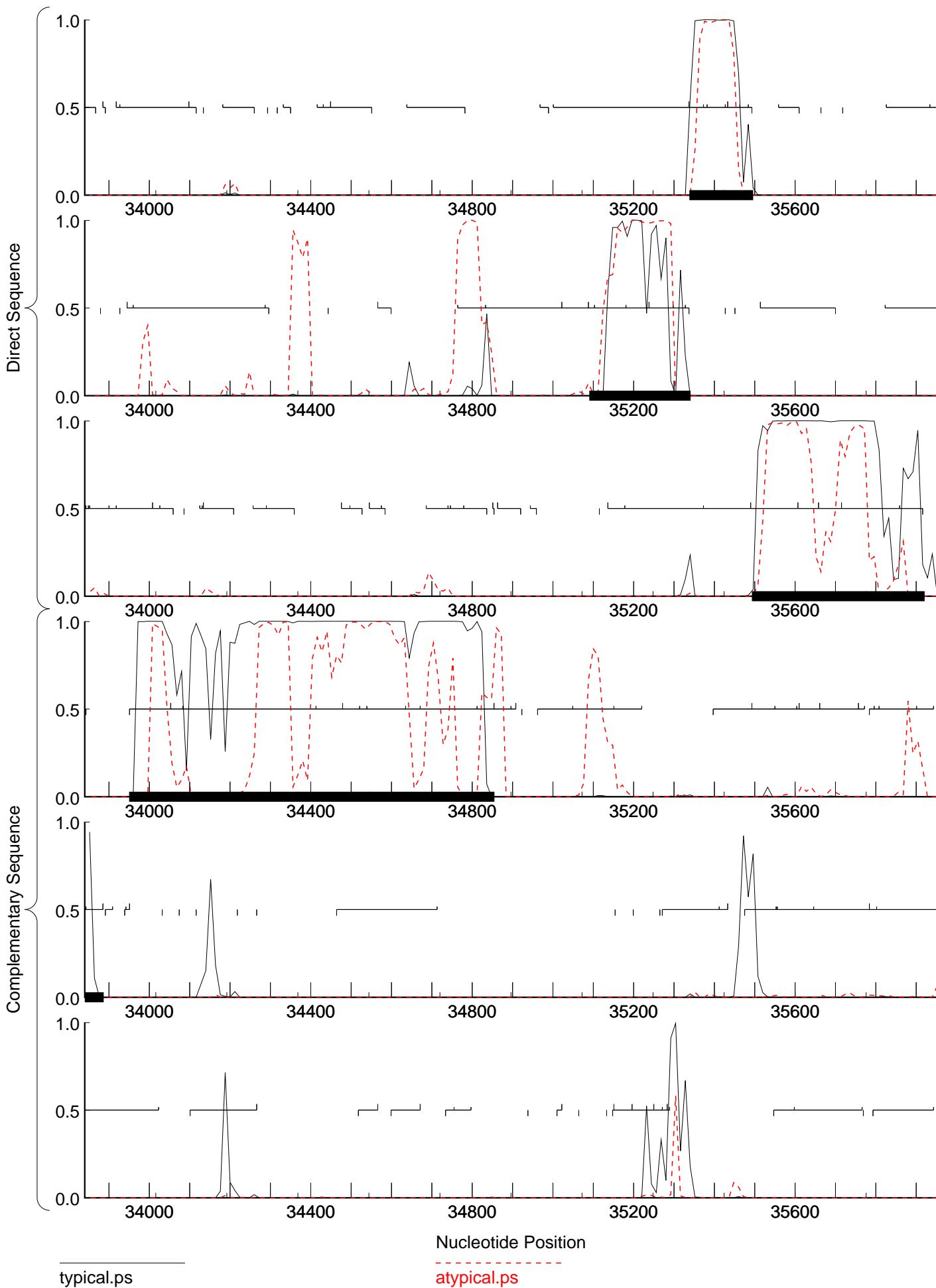


typical.ps

atypical.ps

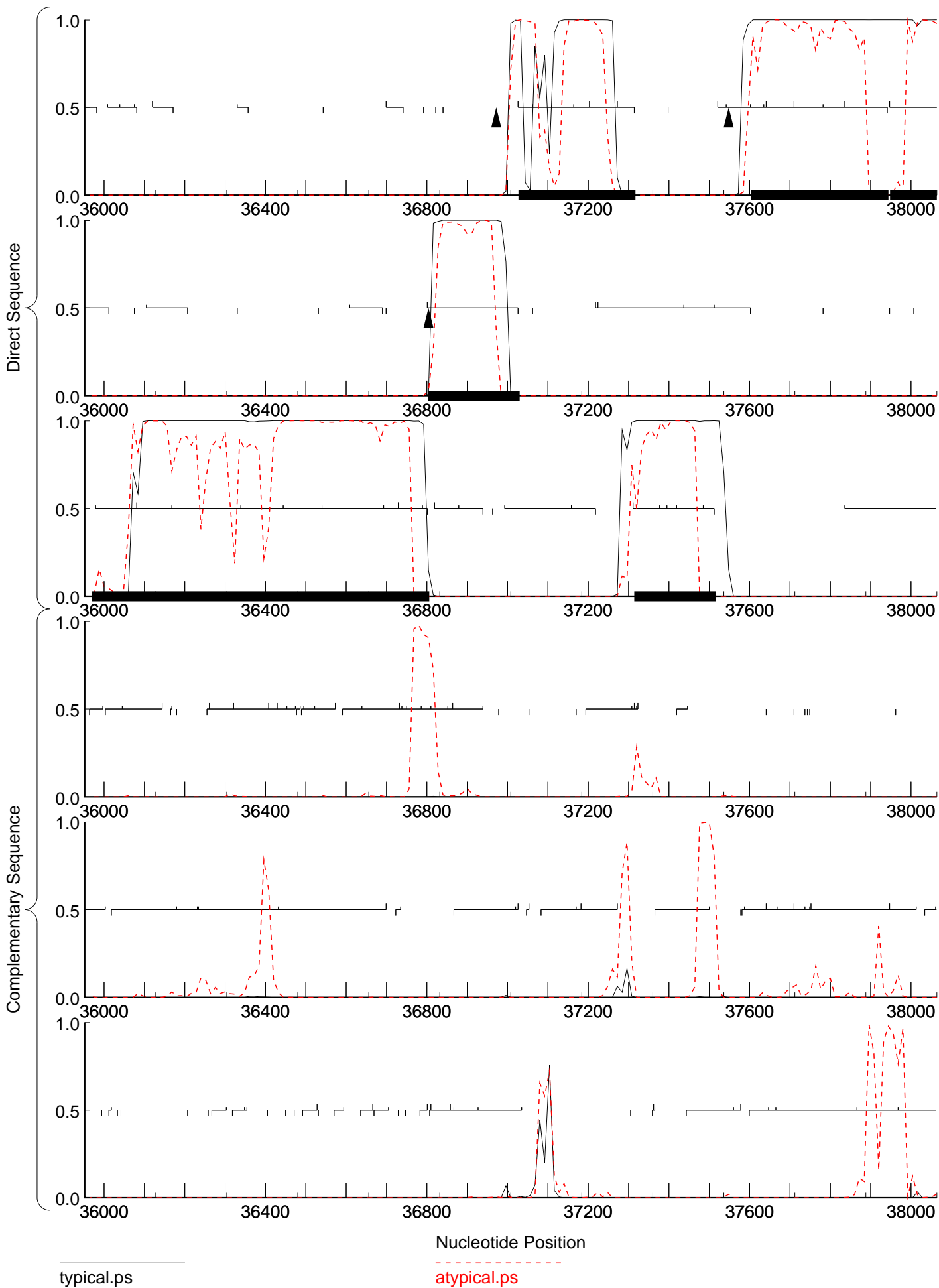


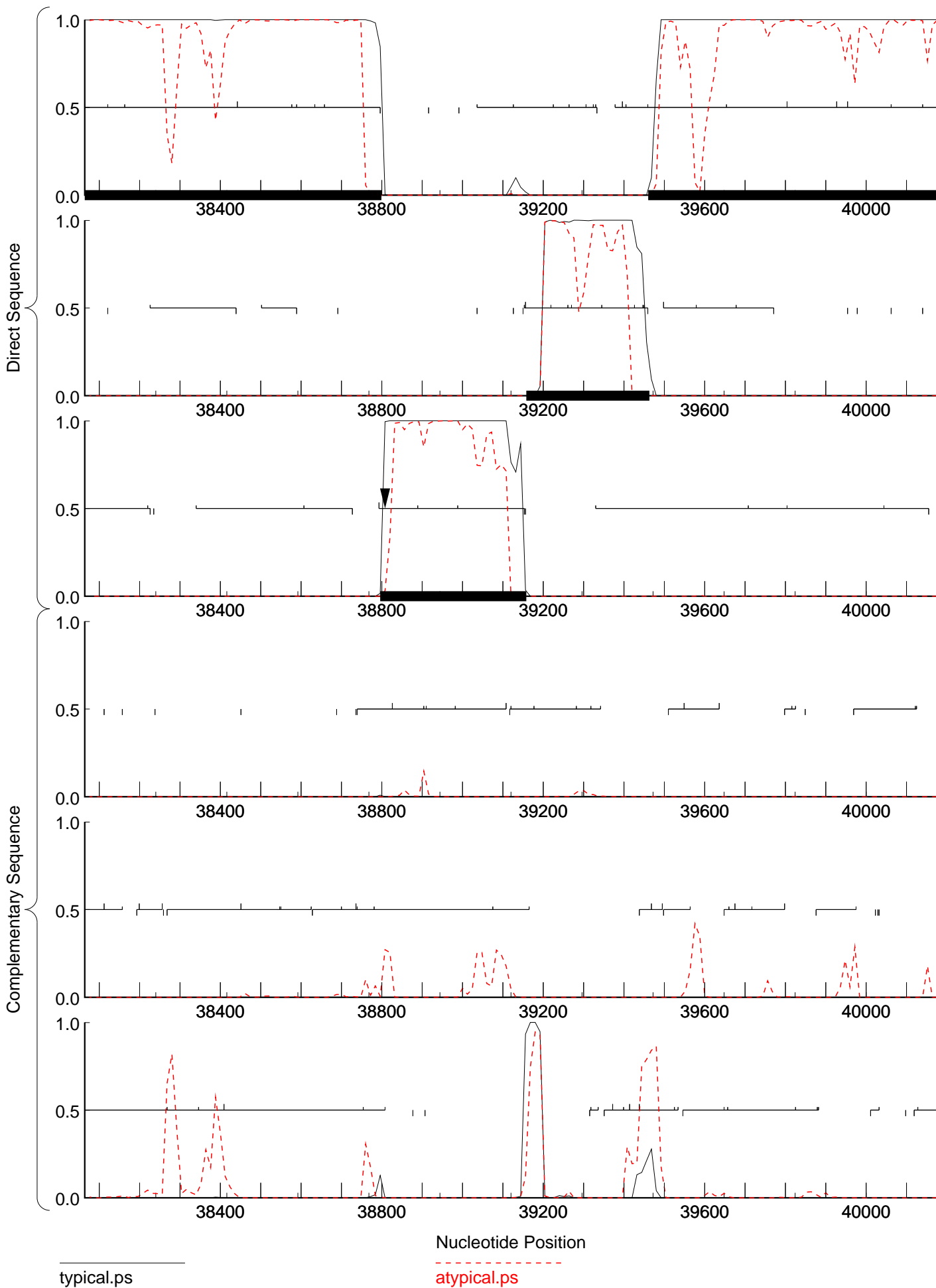


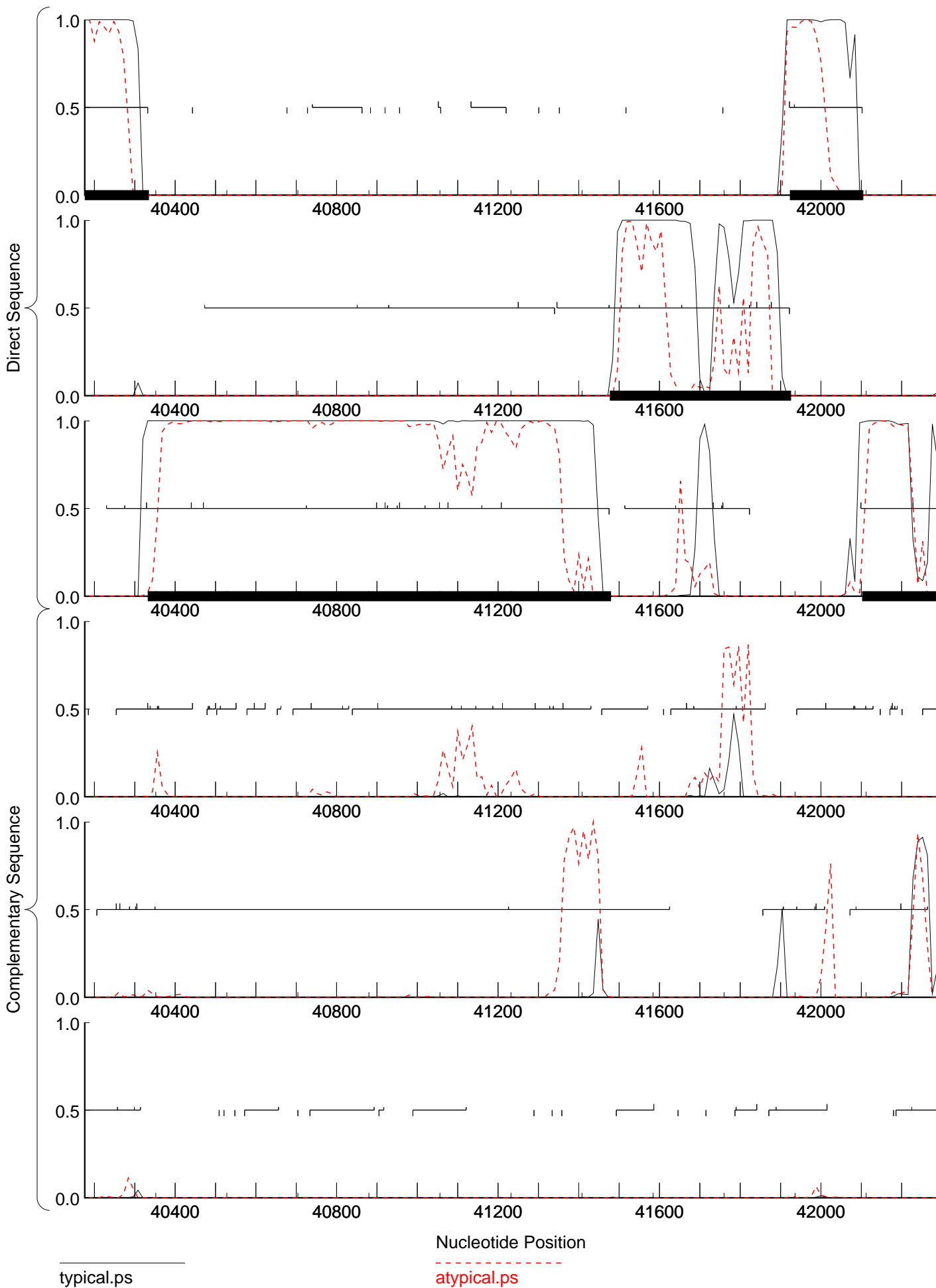


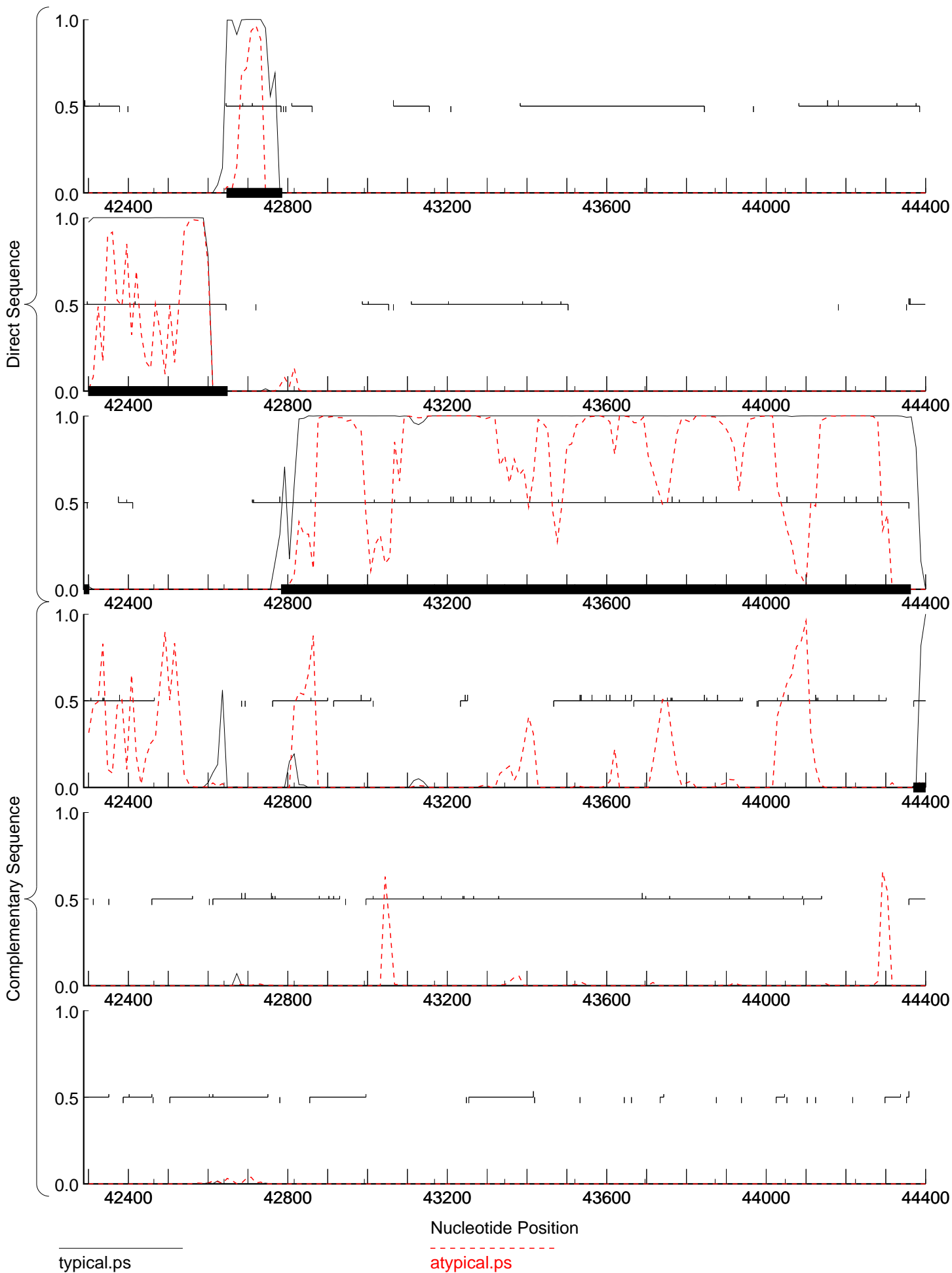
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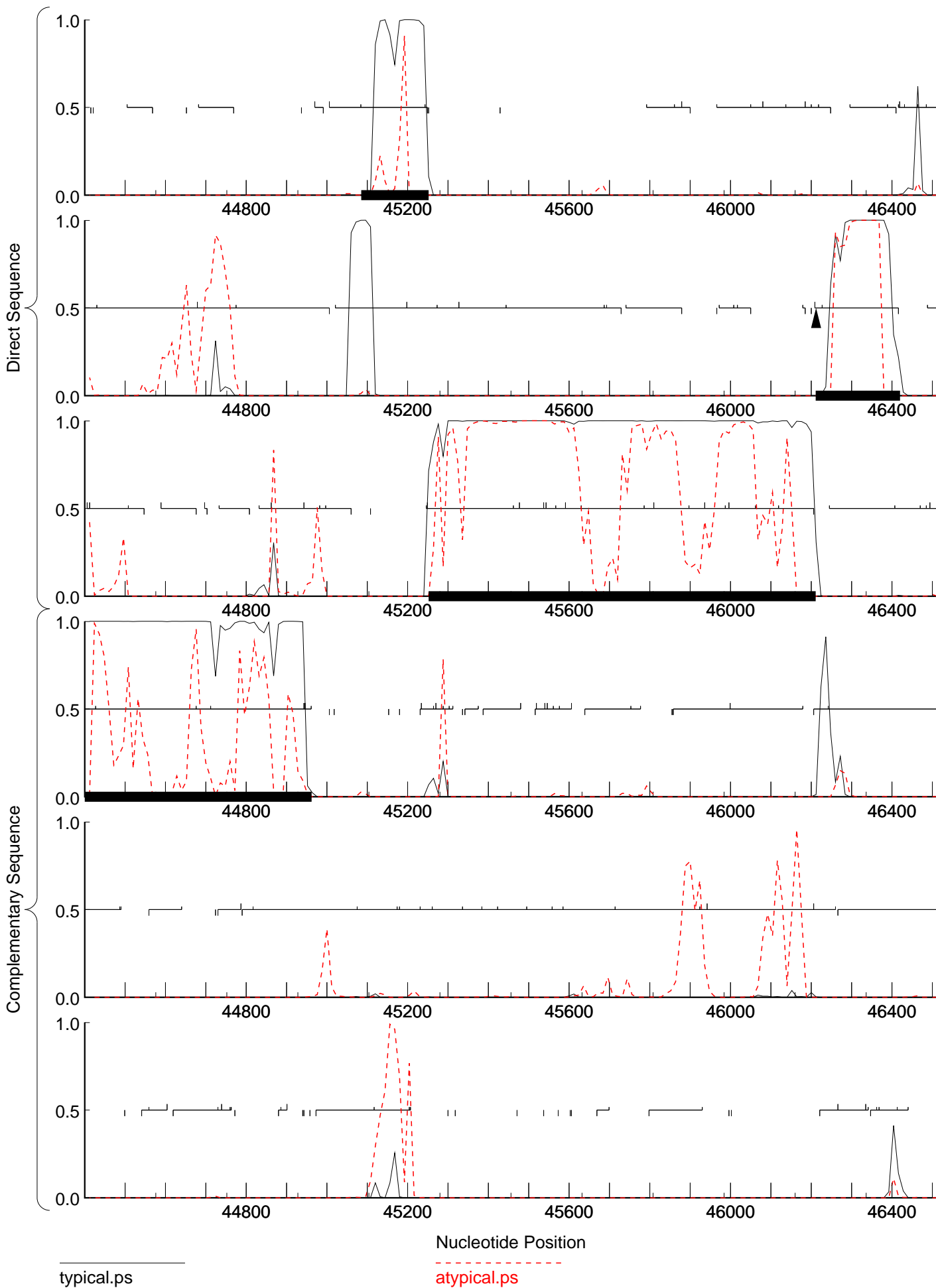
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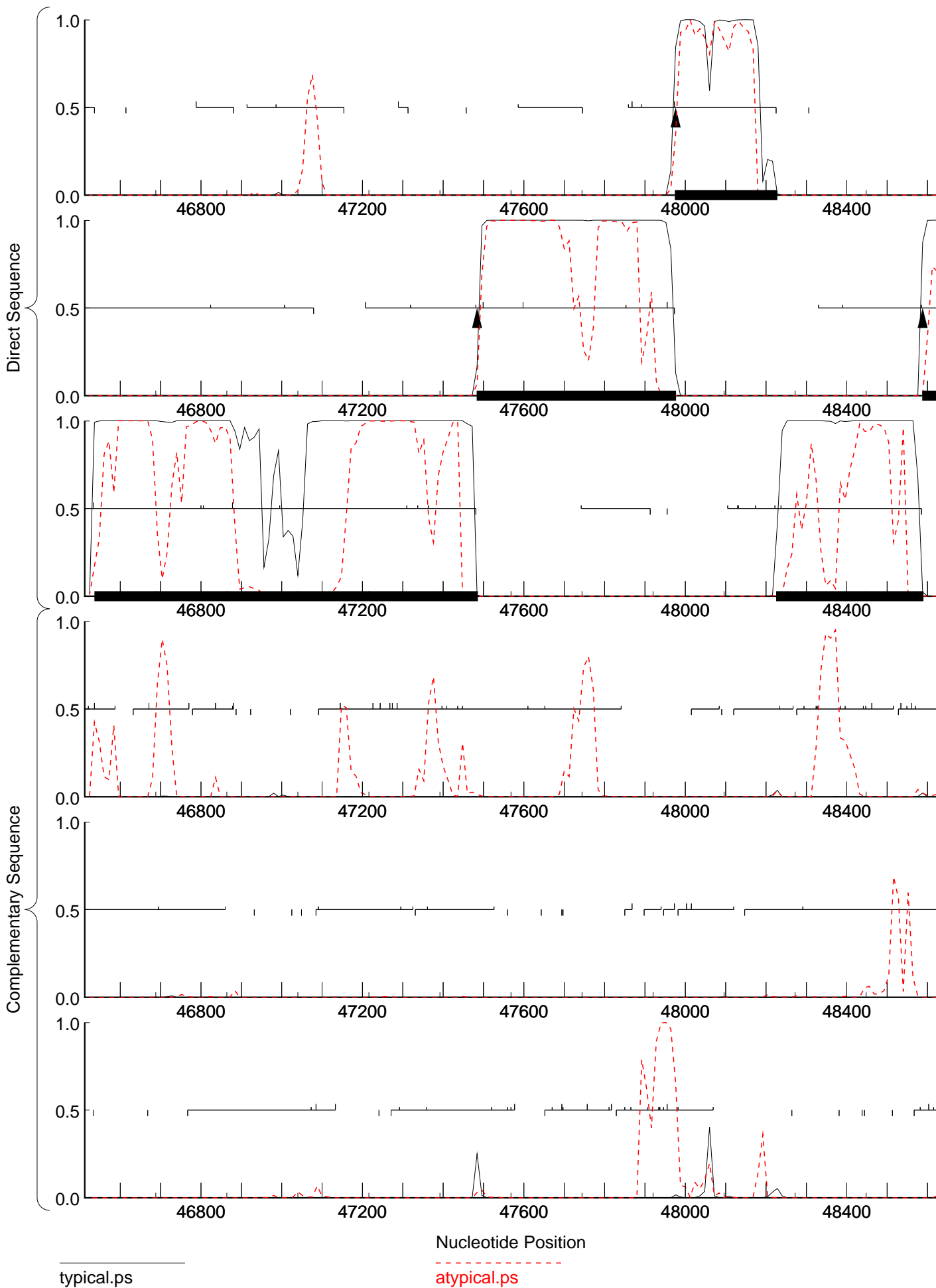








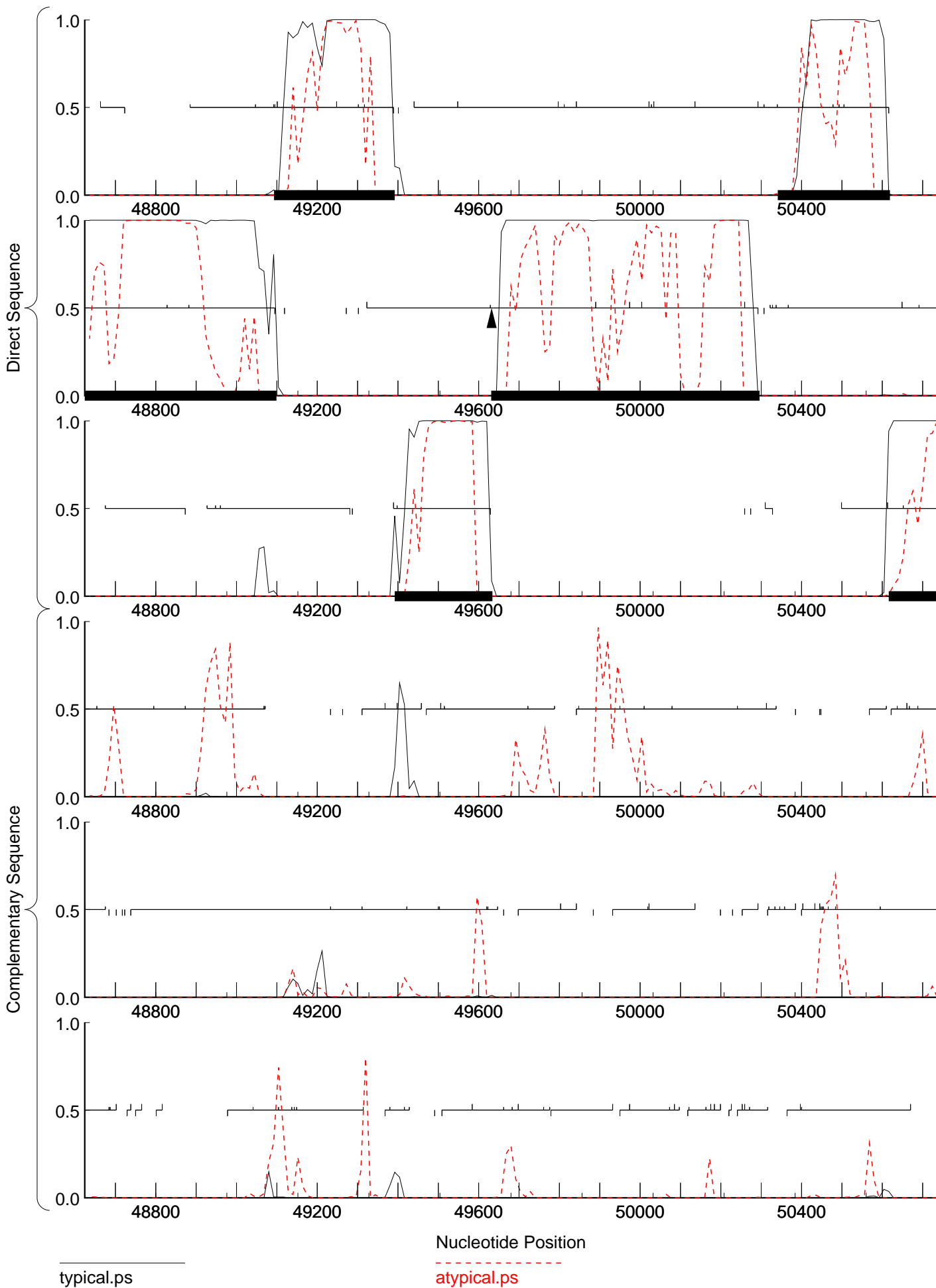




typical.ps

atypical.ps





typical.ps

atypical.ps

