

# GeneMark

Version 2.5p (09.08.06)

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## PROGRAM INFORMATION

Sequence : Mycobacterium phage Wizard007 complete sequence, 51029 bp including 10 bp 3' overhang (CGGC  
Analysis Date : 5/23/17 at 12:52:41  
Pages : 26  
Sequence Length : 51029 bp  
GC Content : 63.85%

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,

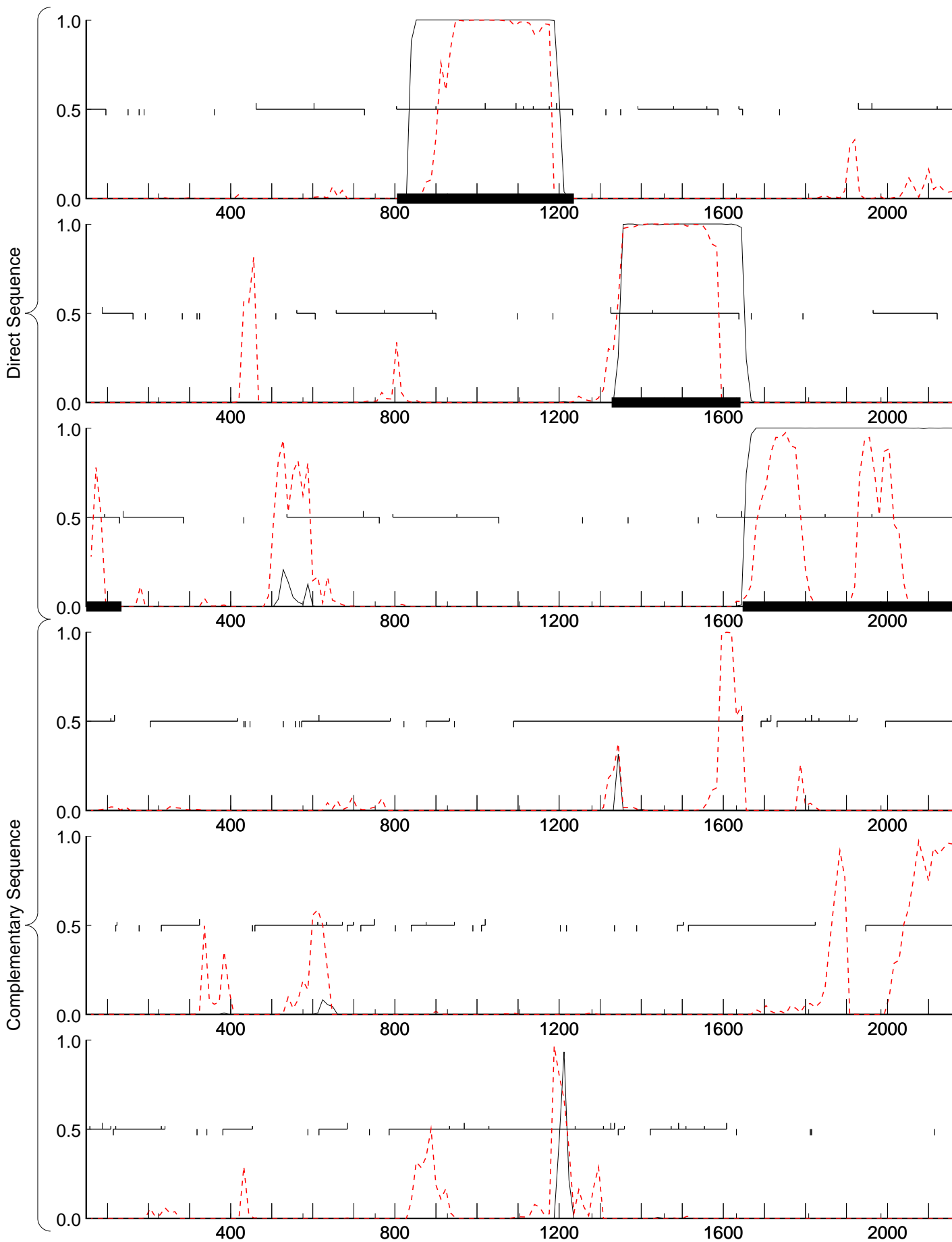
## Matrix notes & comments

Training set derived by GeneMarkS, 4.30 October 2014  
Tue May 23 12:52:41 2017

## MATRIX INFORMATION

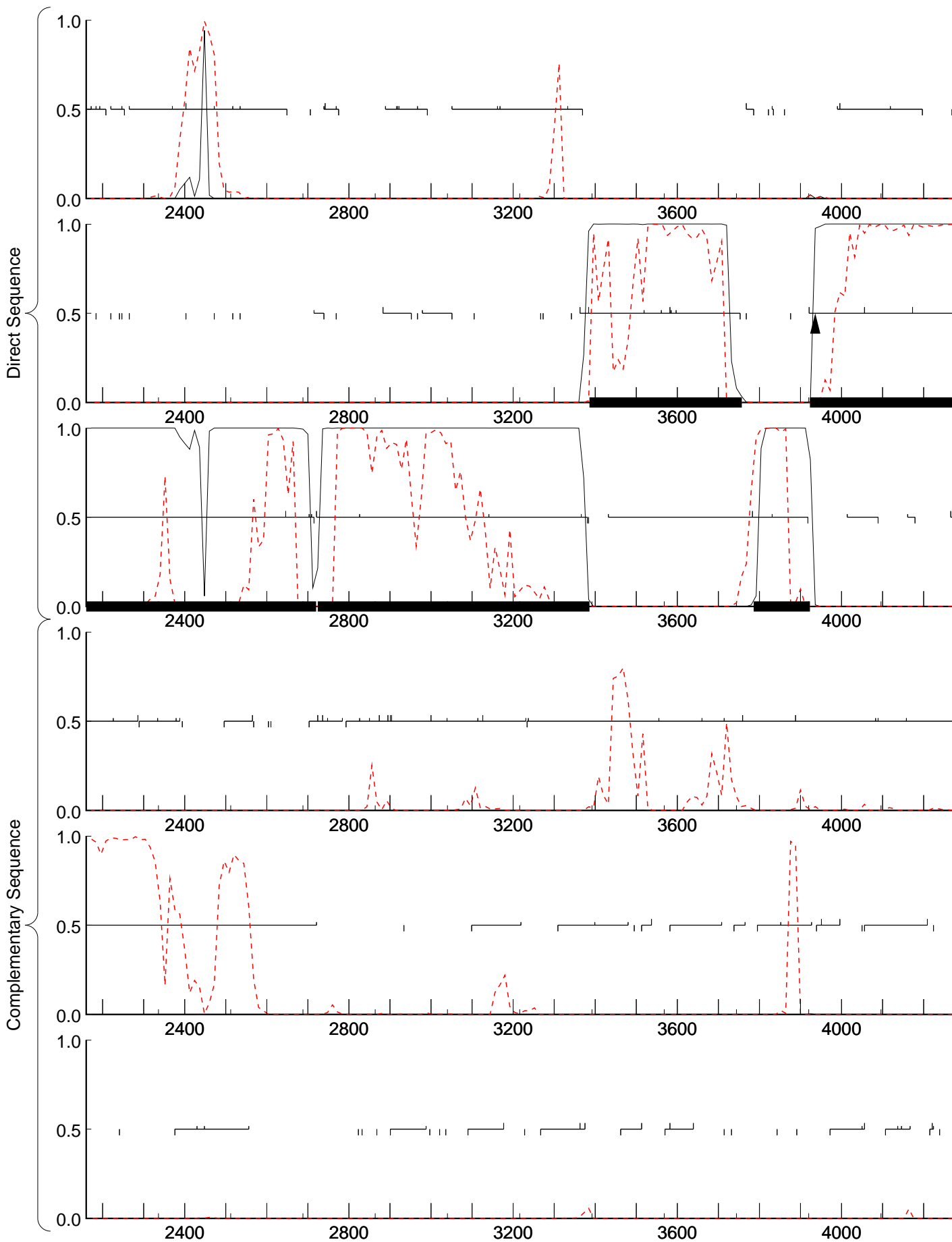
Matrix : -  
Author : -  
Order : 4

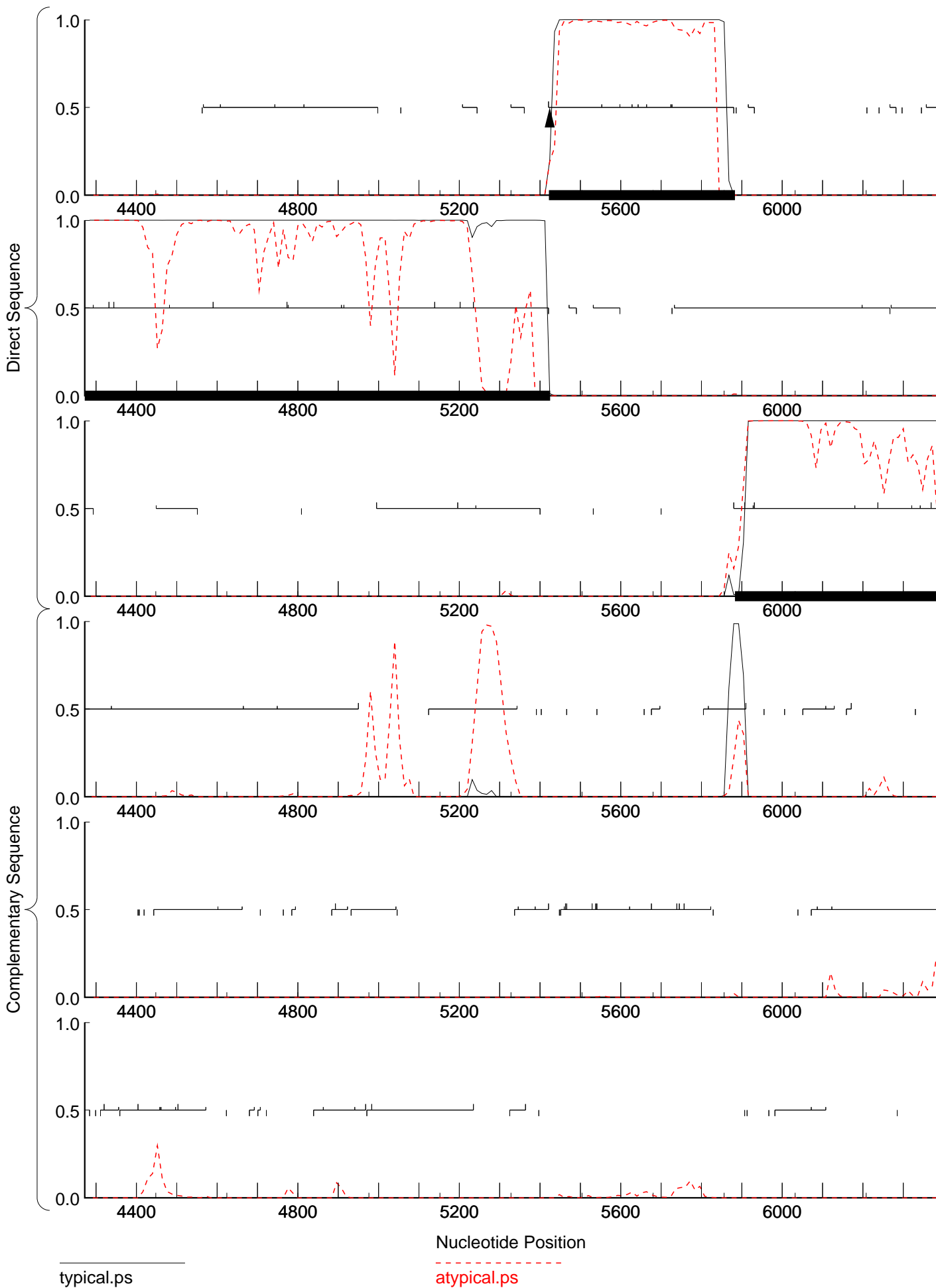
Send questions / comments to:  
Dr. M. Borodovsky  
Georgia Institute of Technology  
School of Biology  
Atlanta, GA 30332-0230



typical.ps

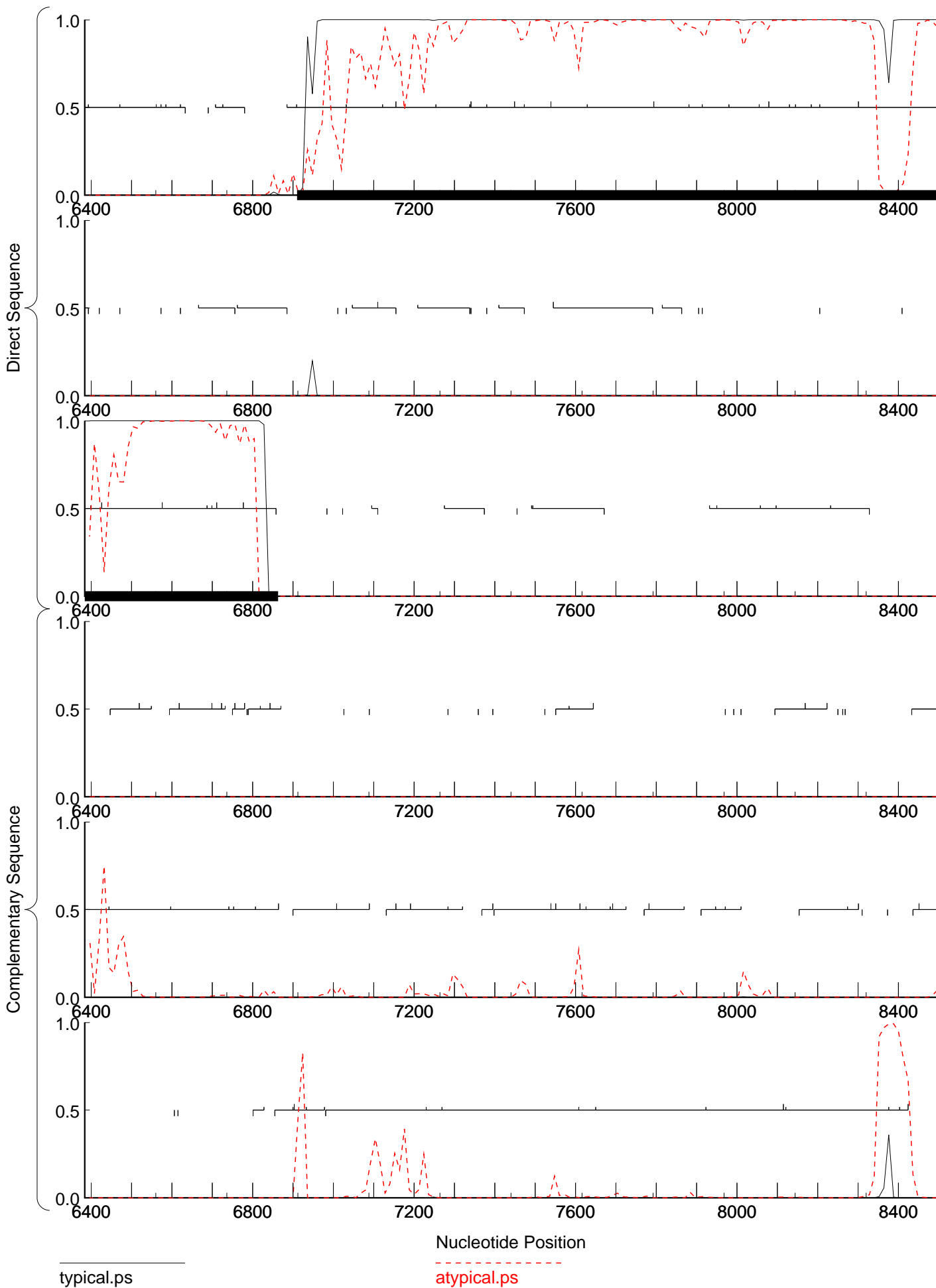
atypical.ps

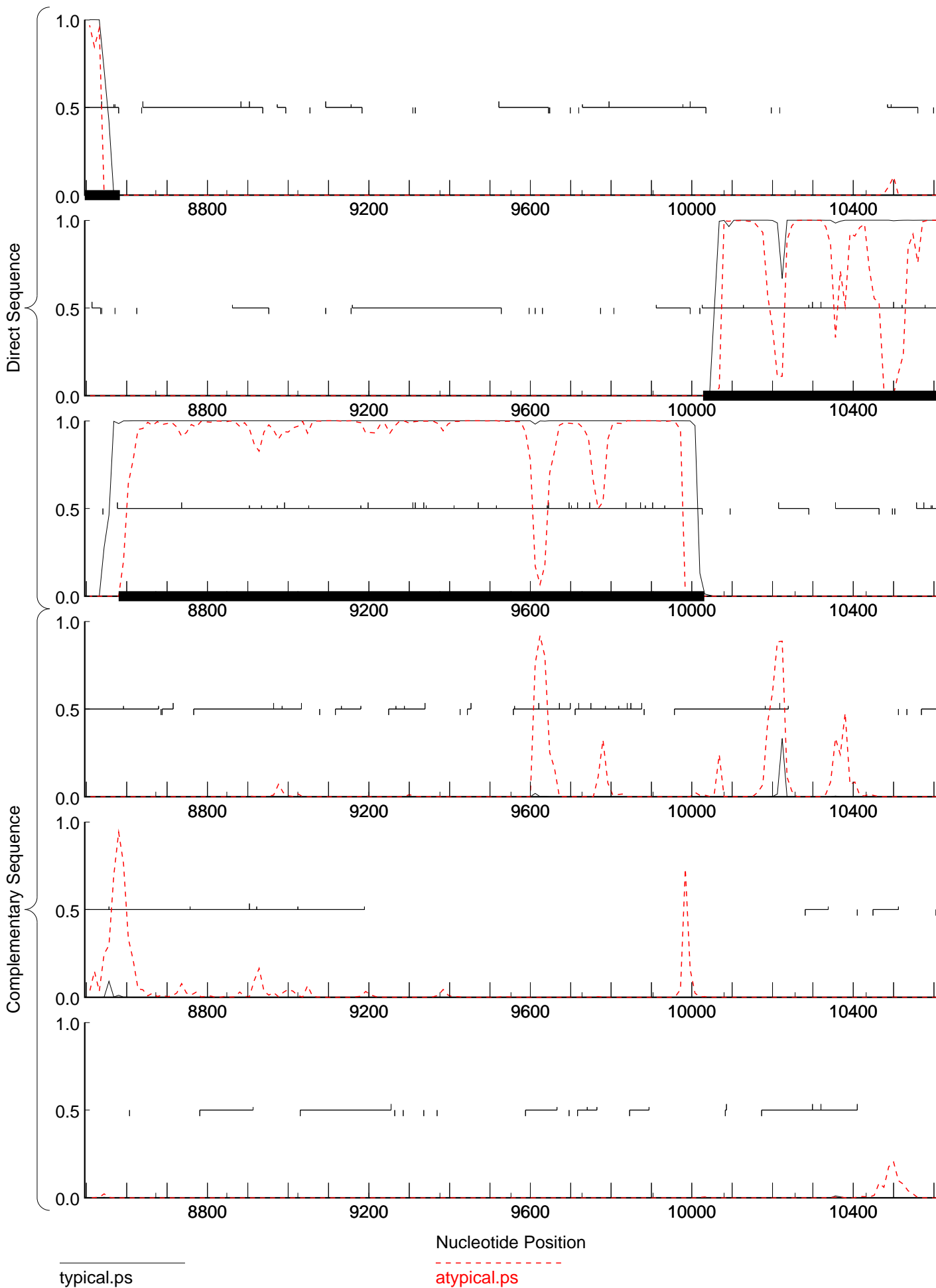


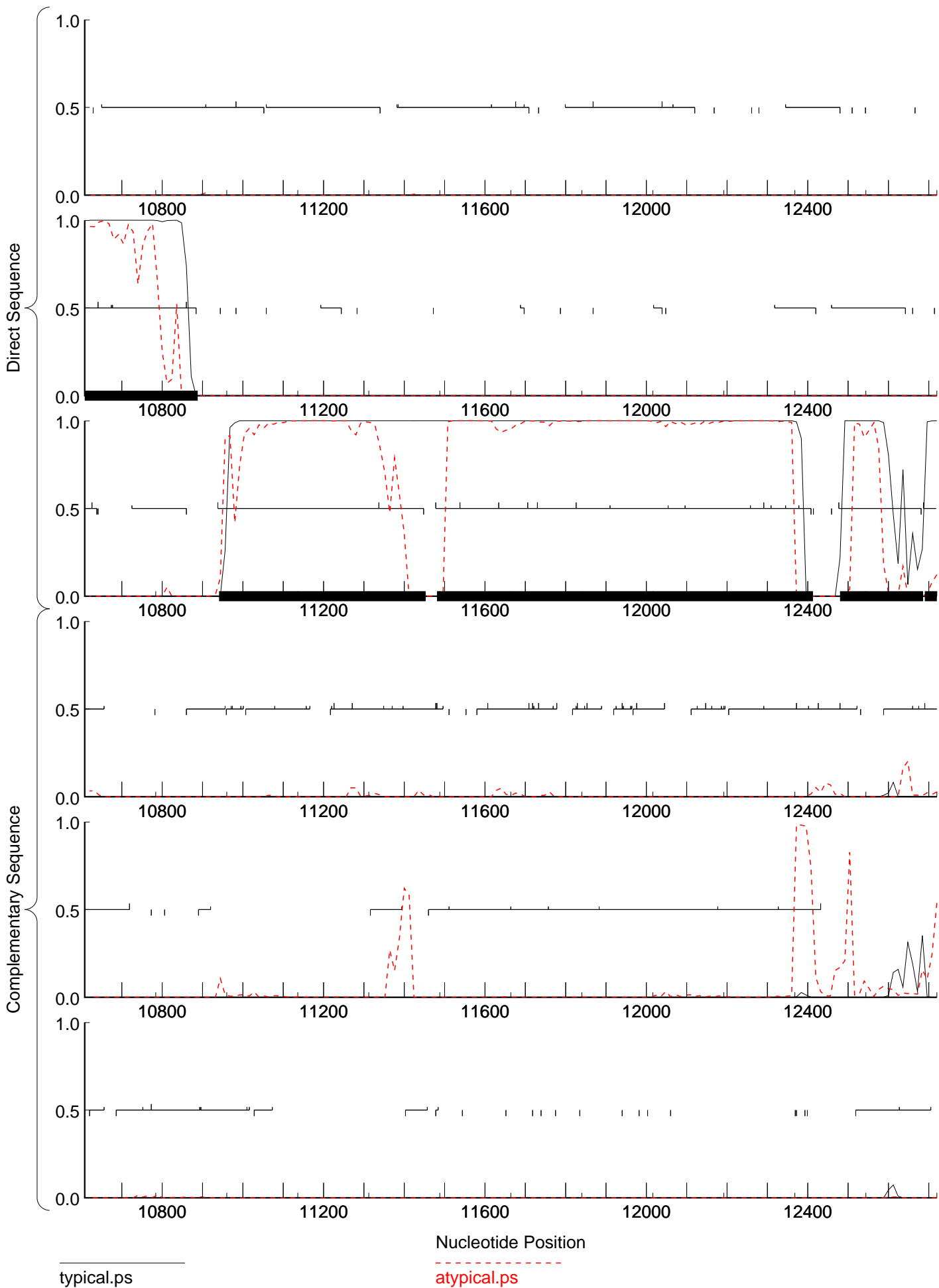


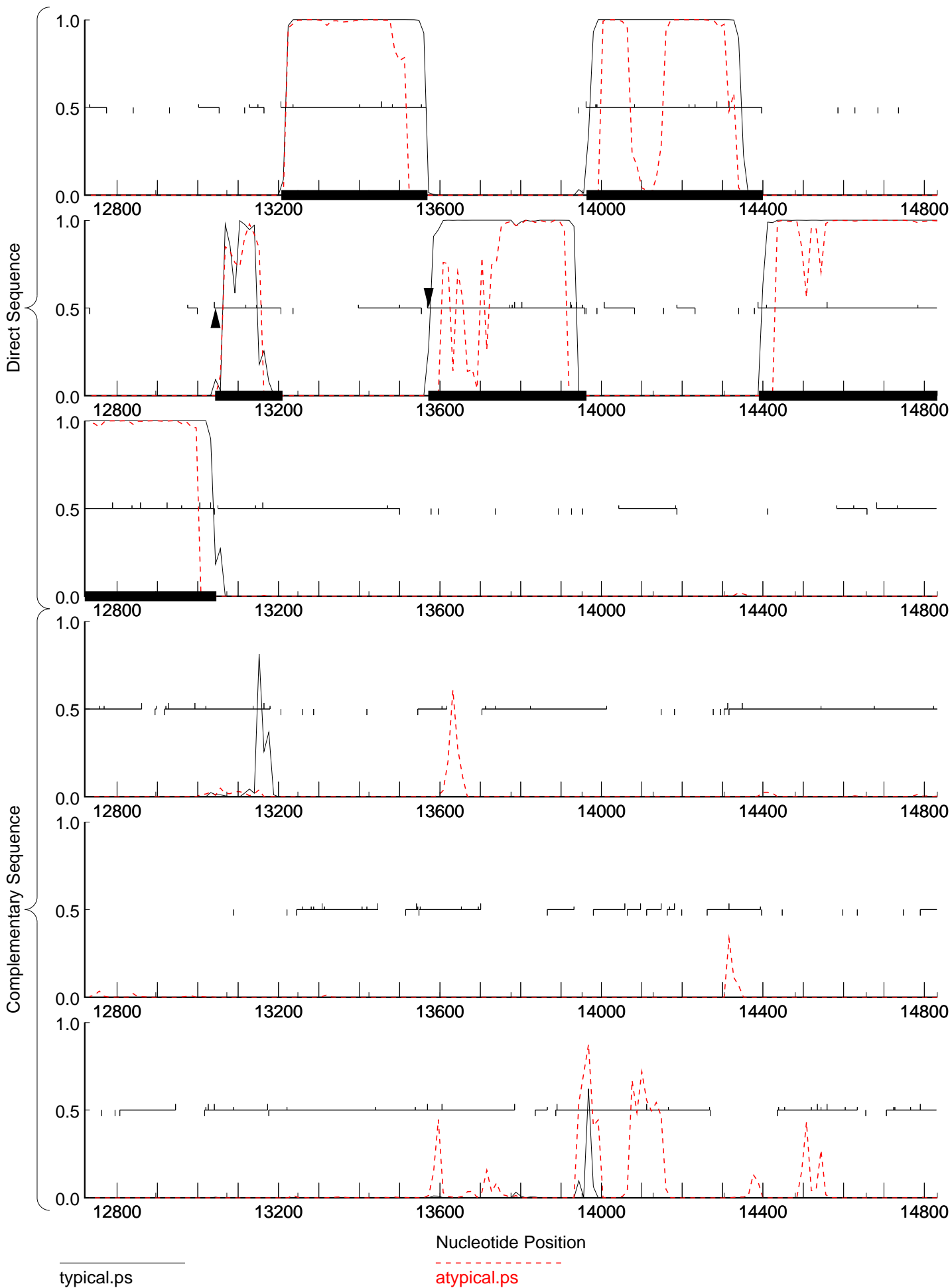
typical.ps

atypical.ps

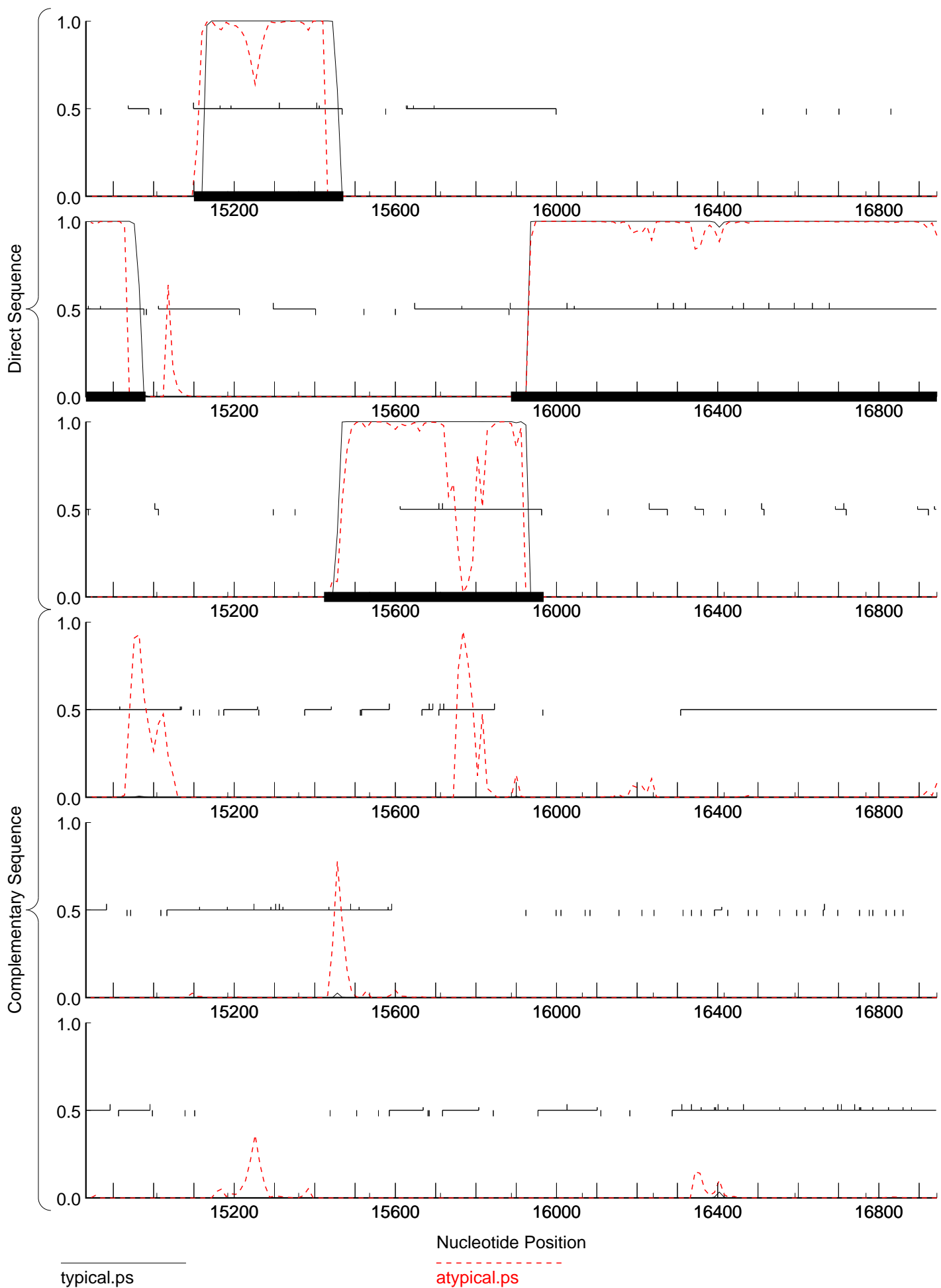


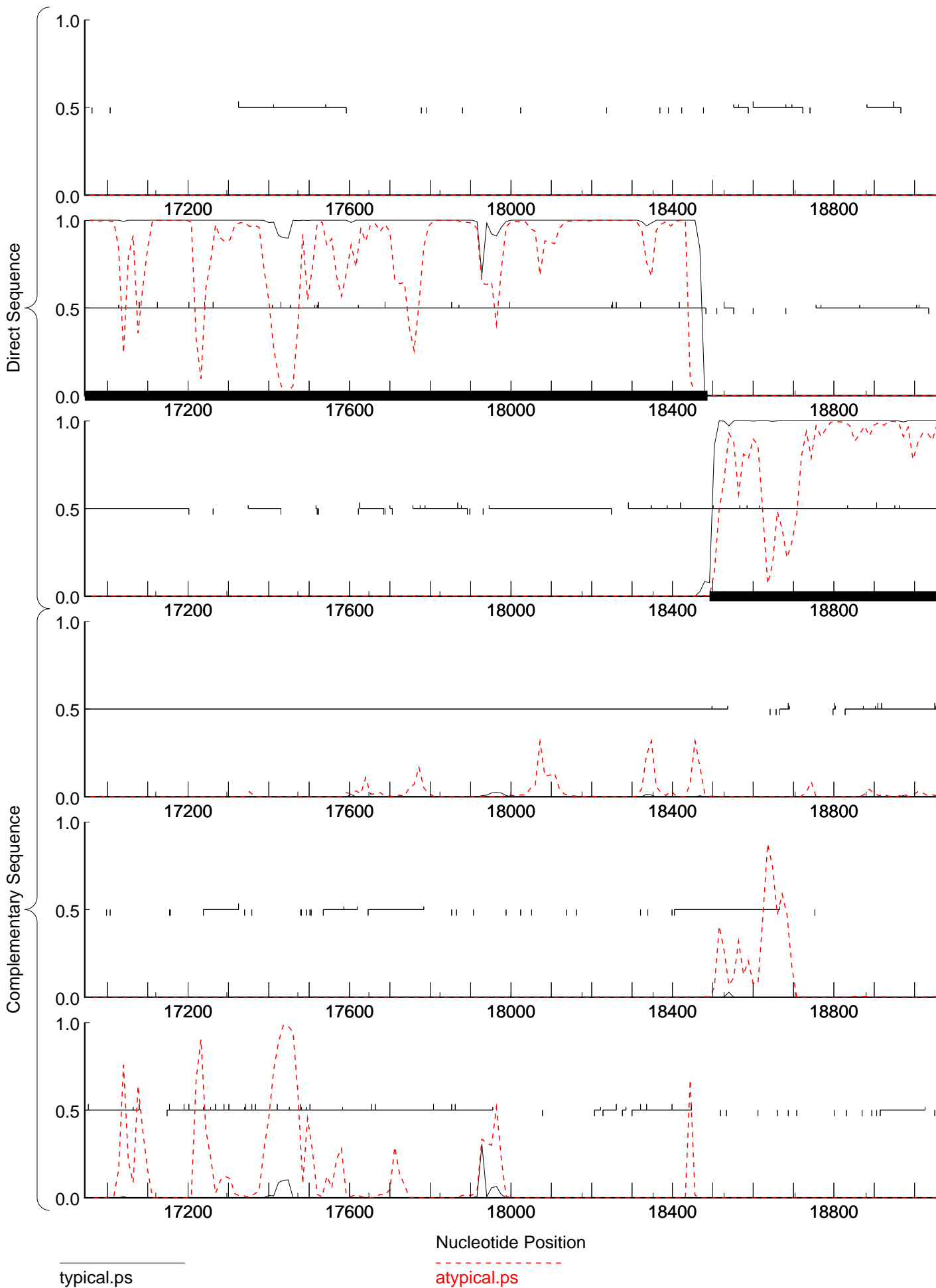


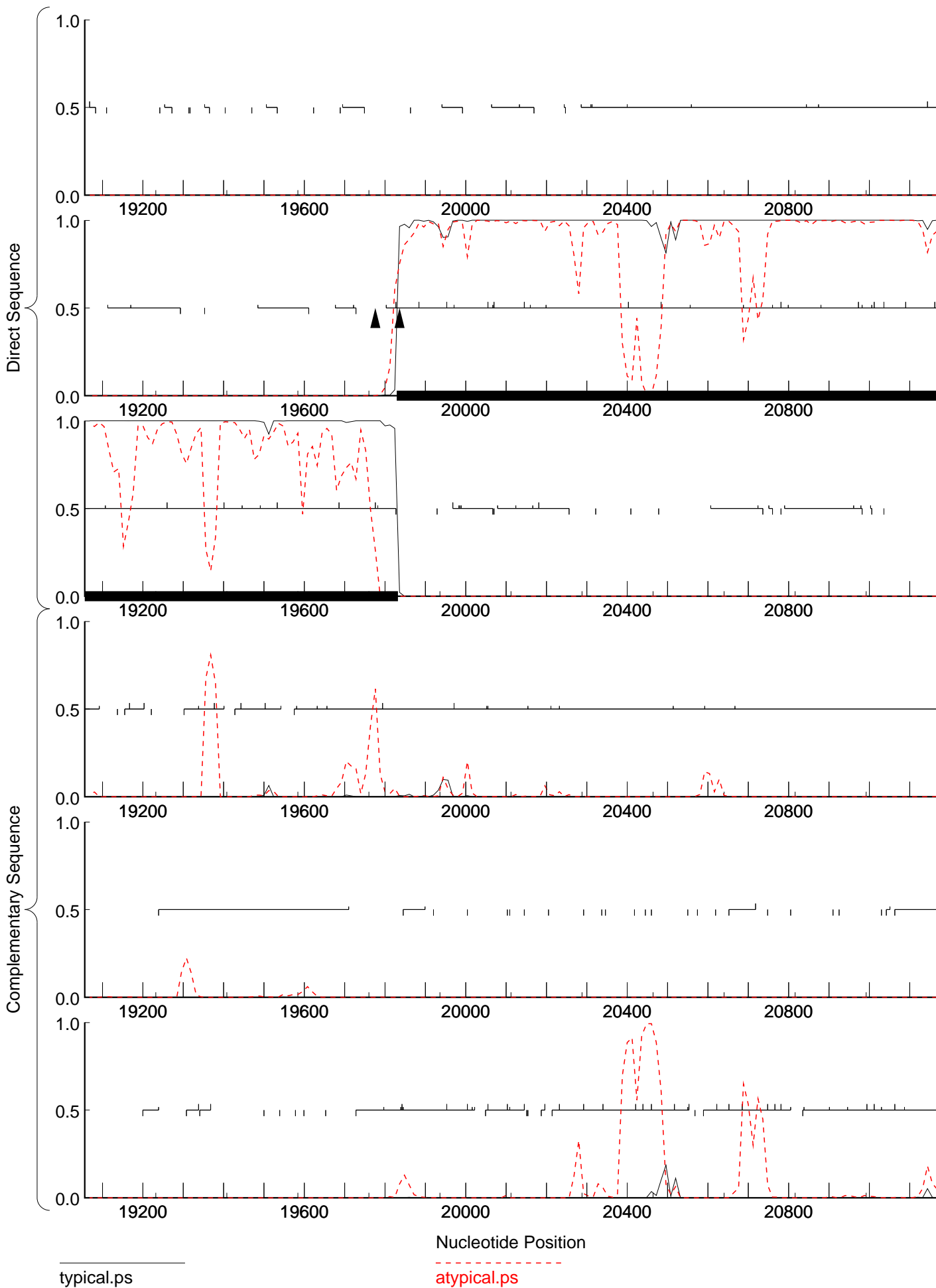


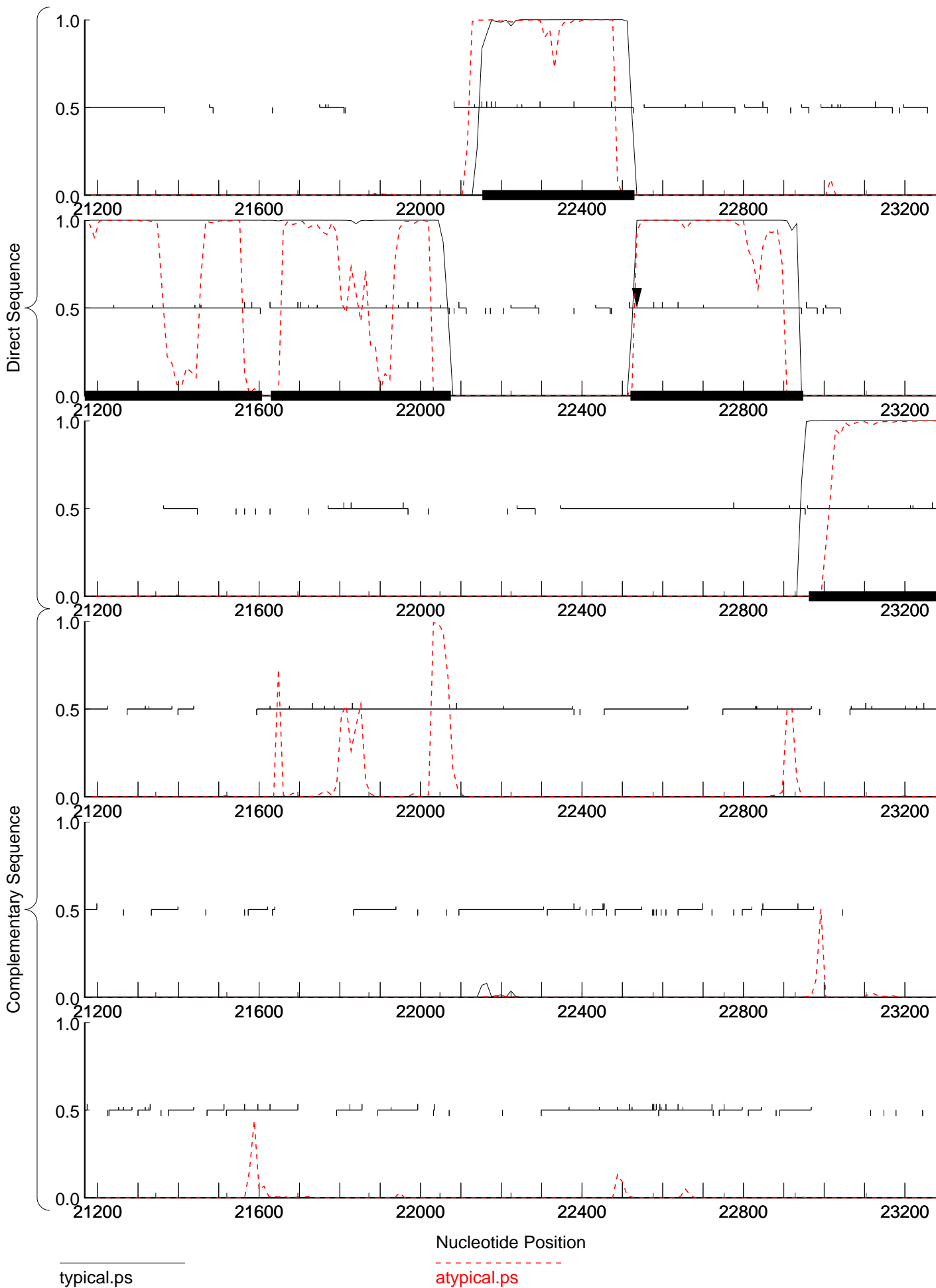


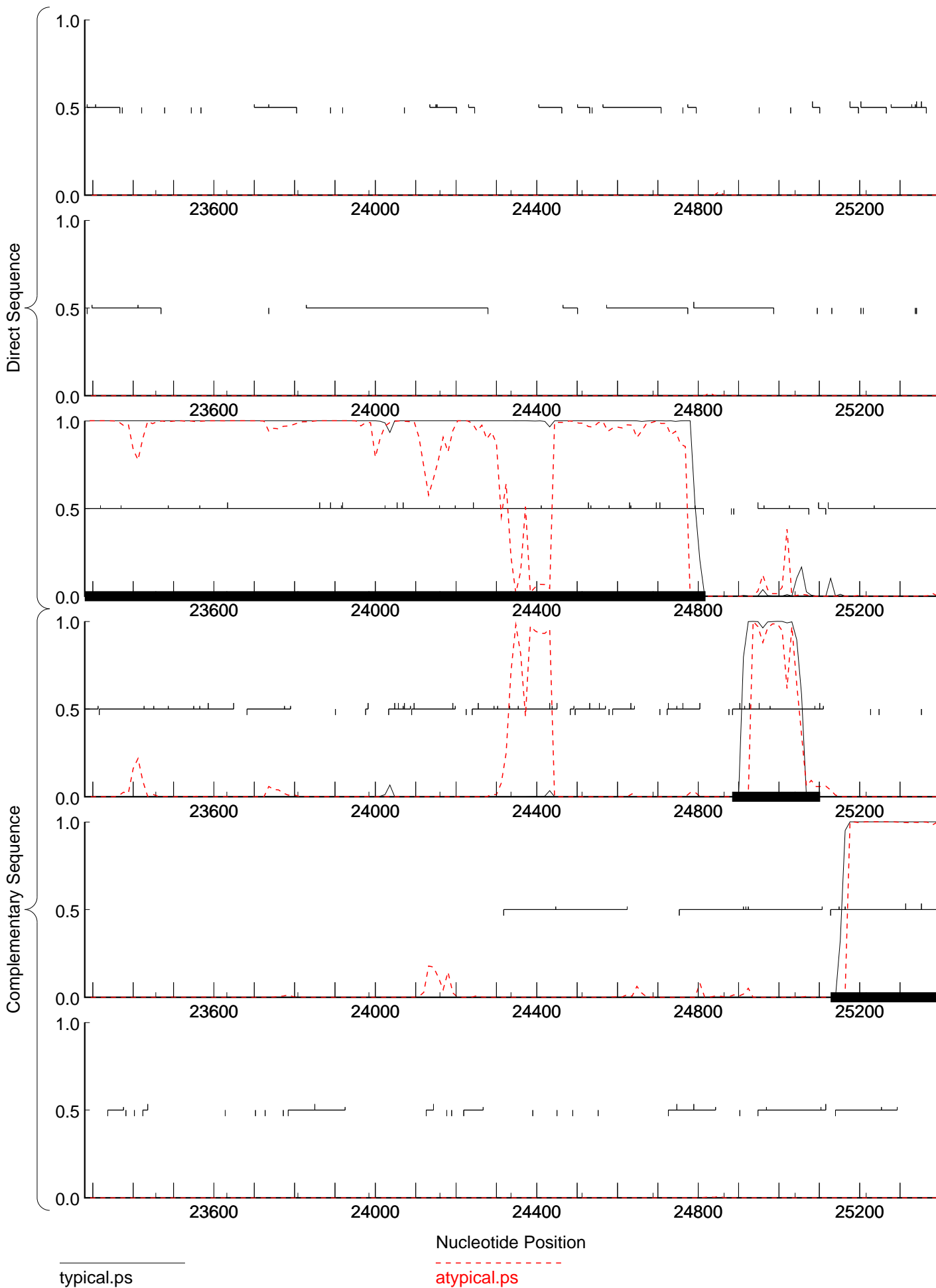








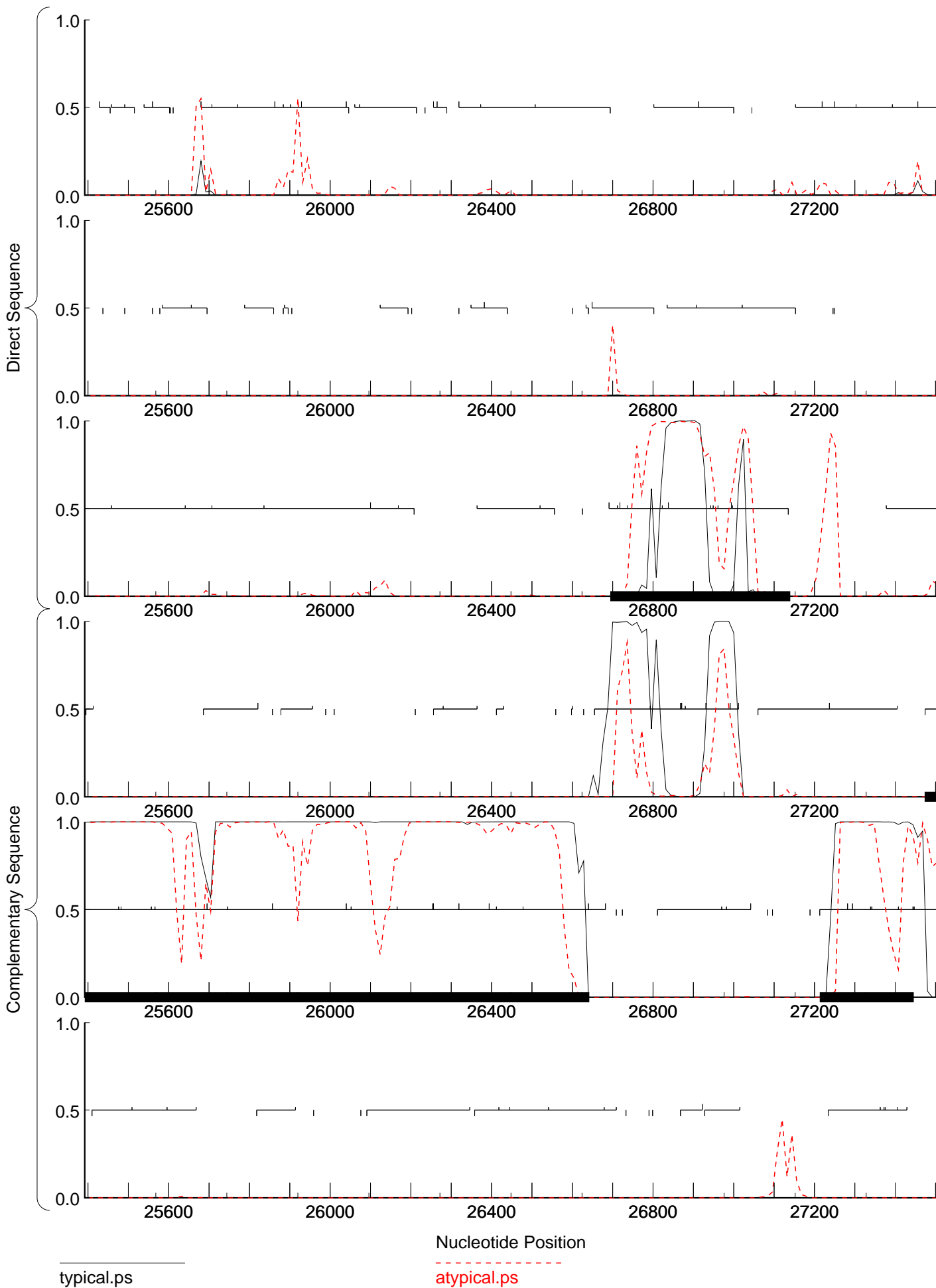


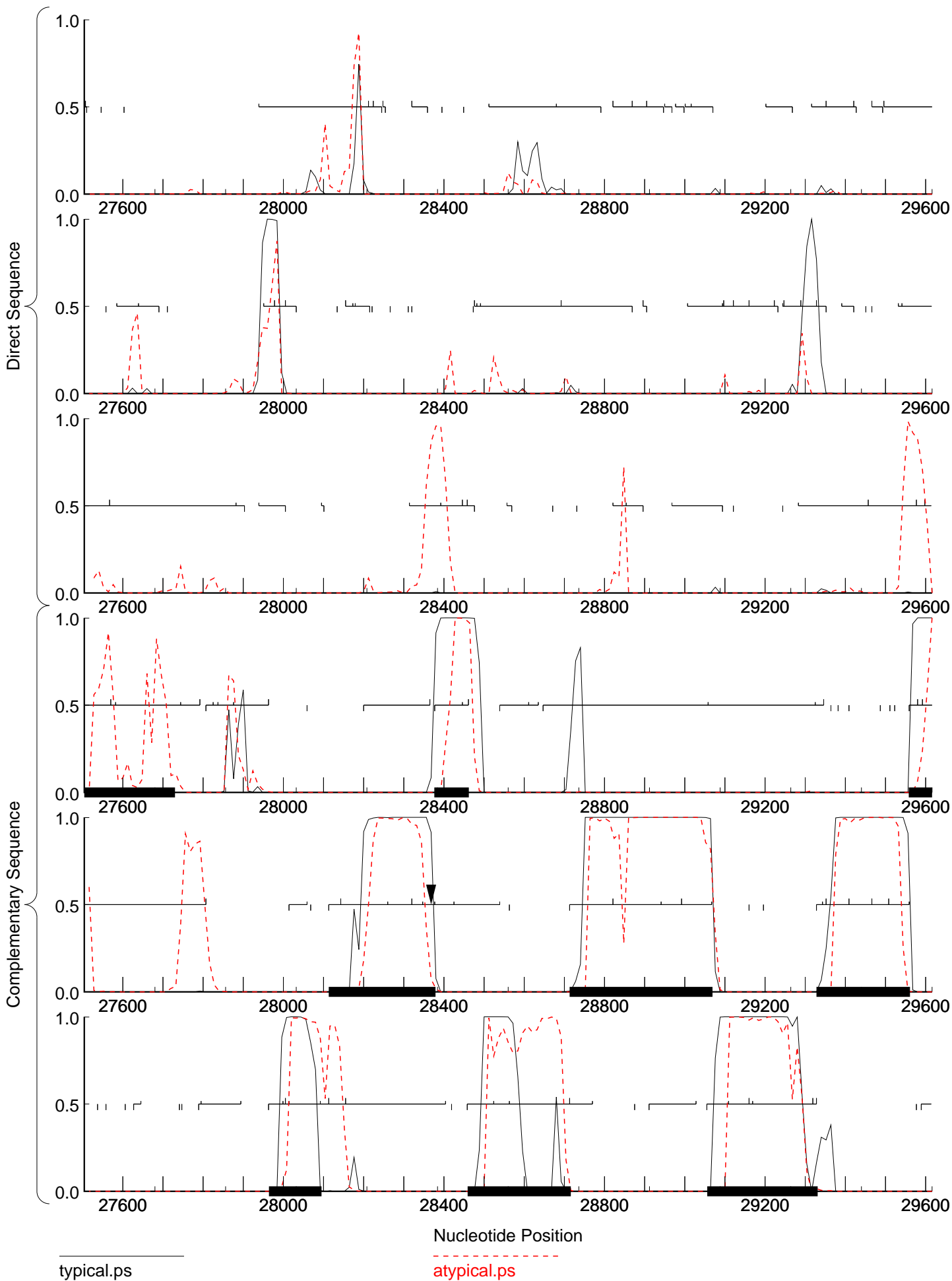


typical.ps

Nucleotide Position

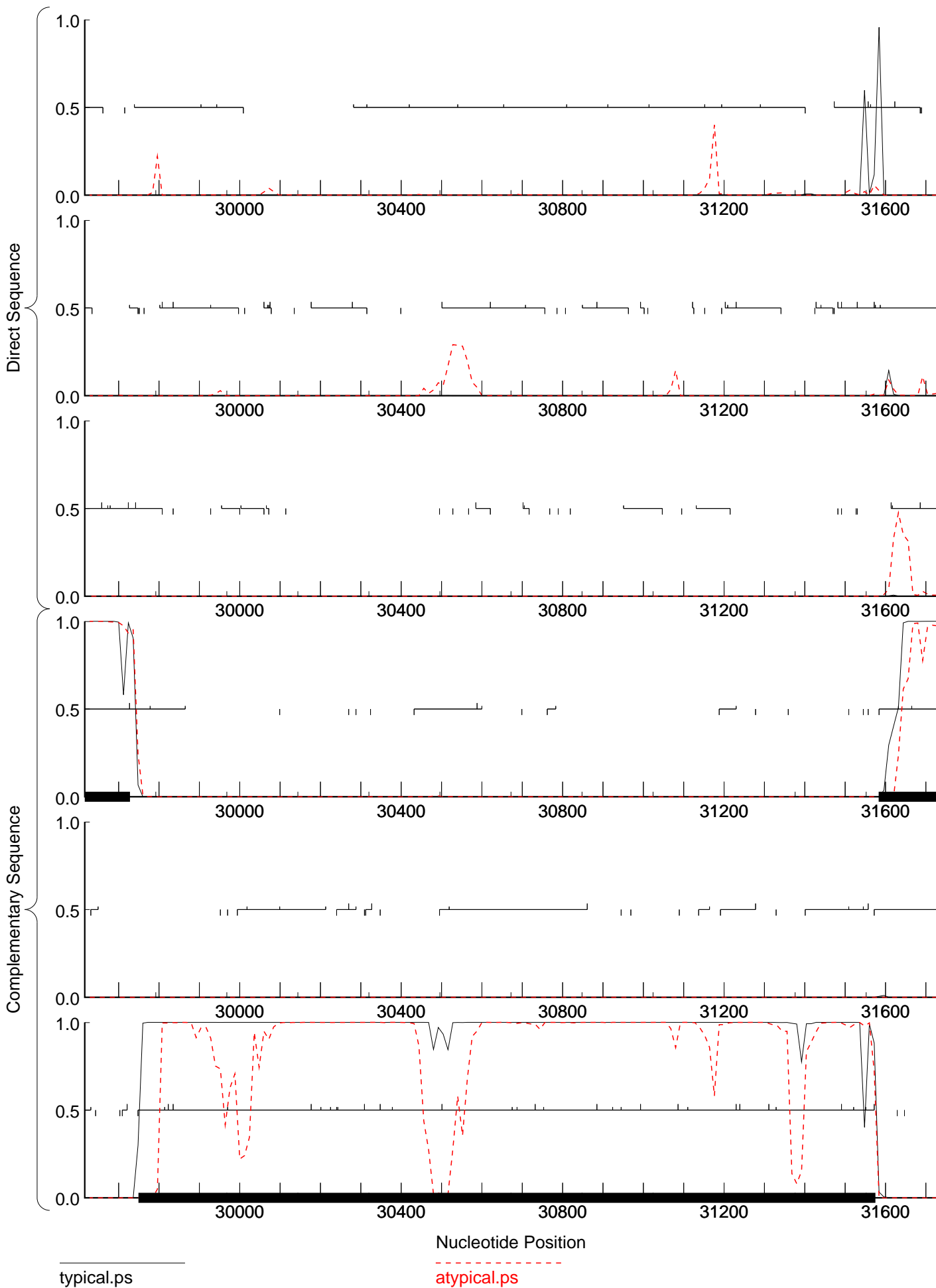
atypical.ps



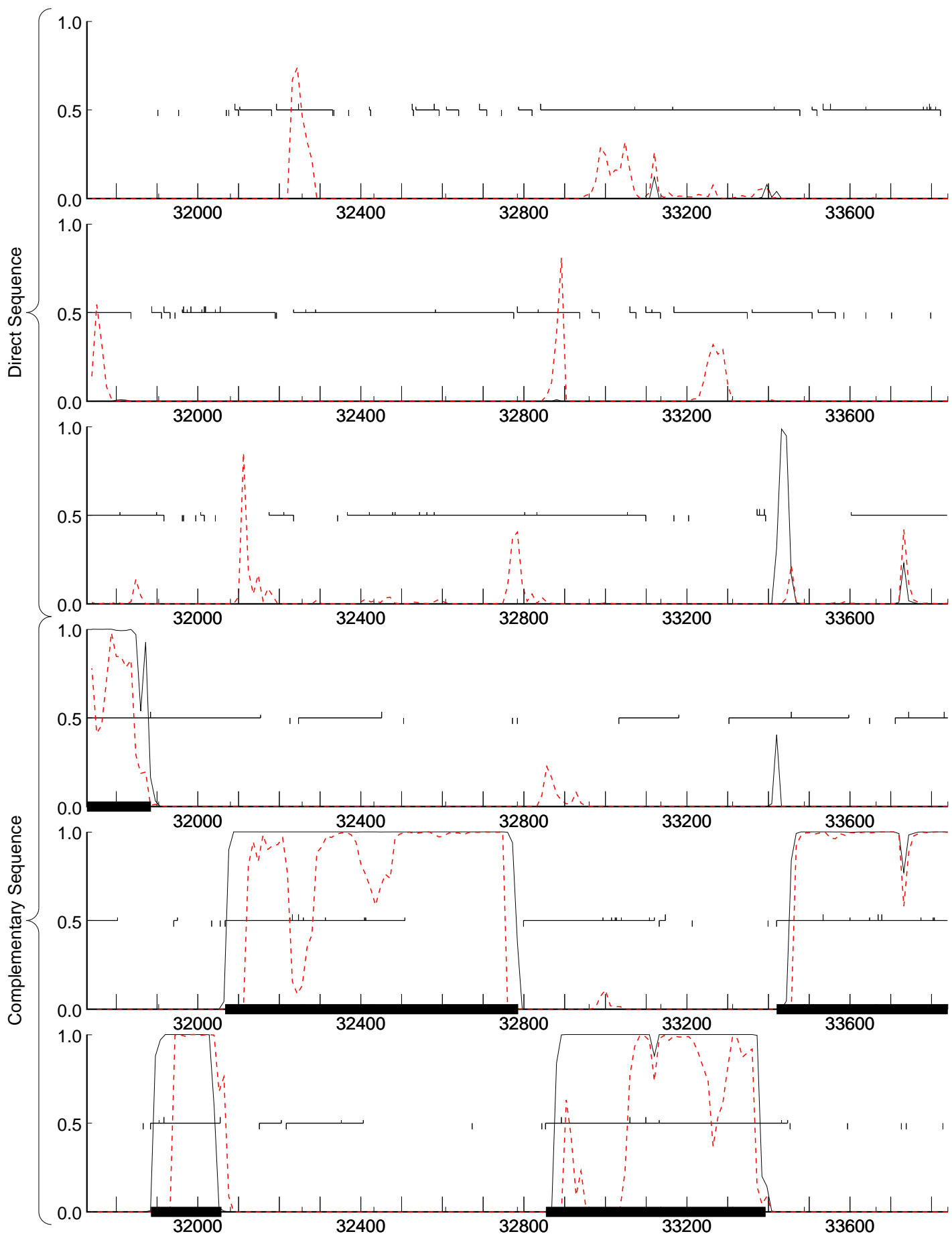


typical.ps

atypical.ps

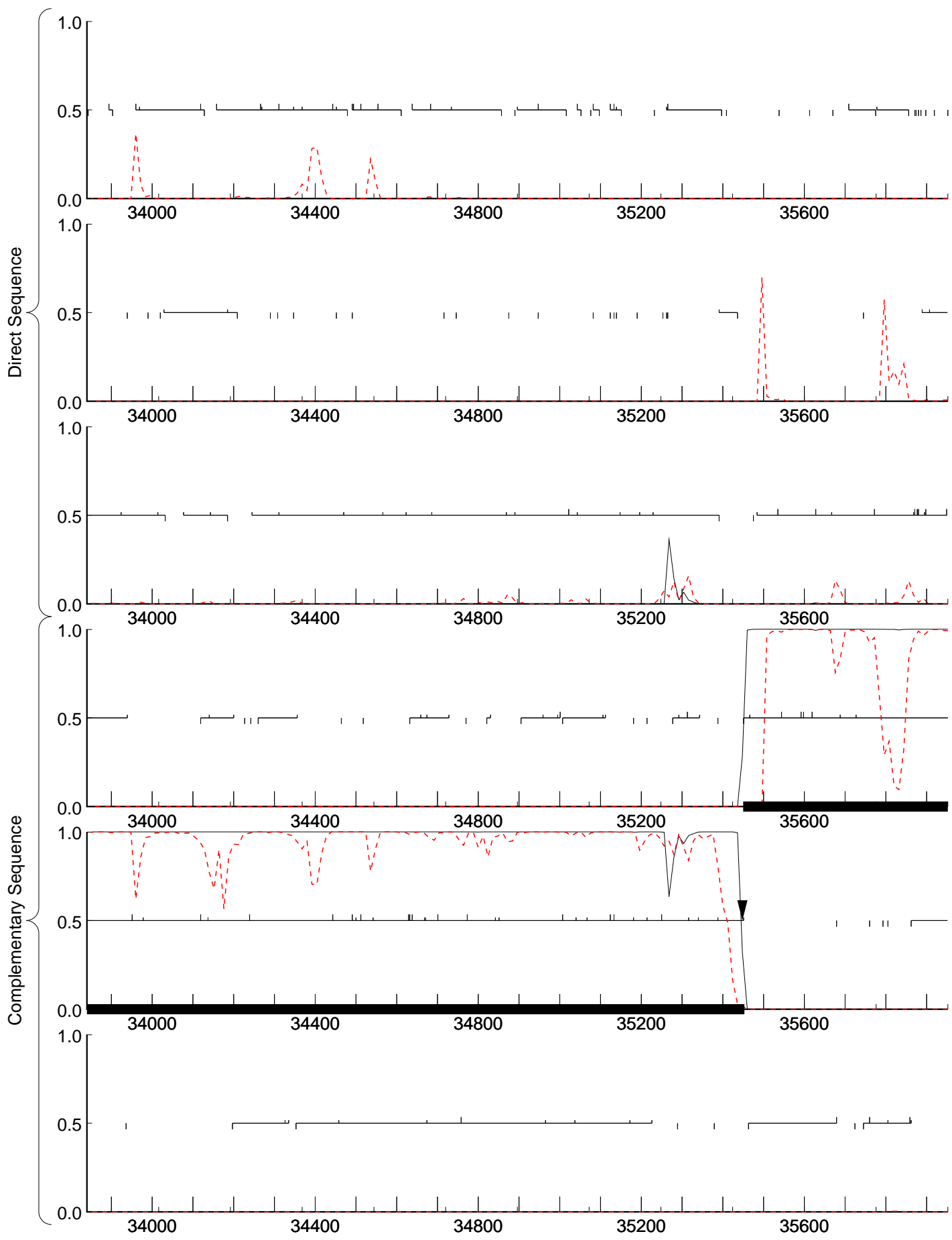






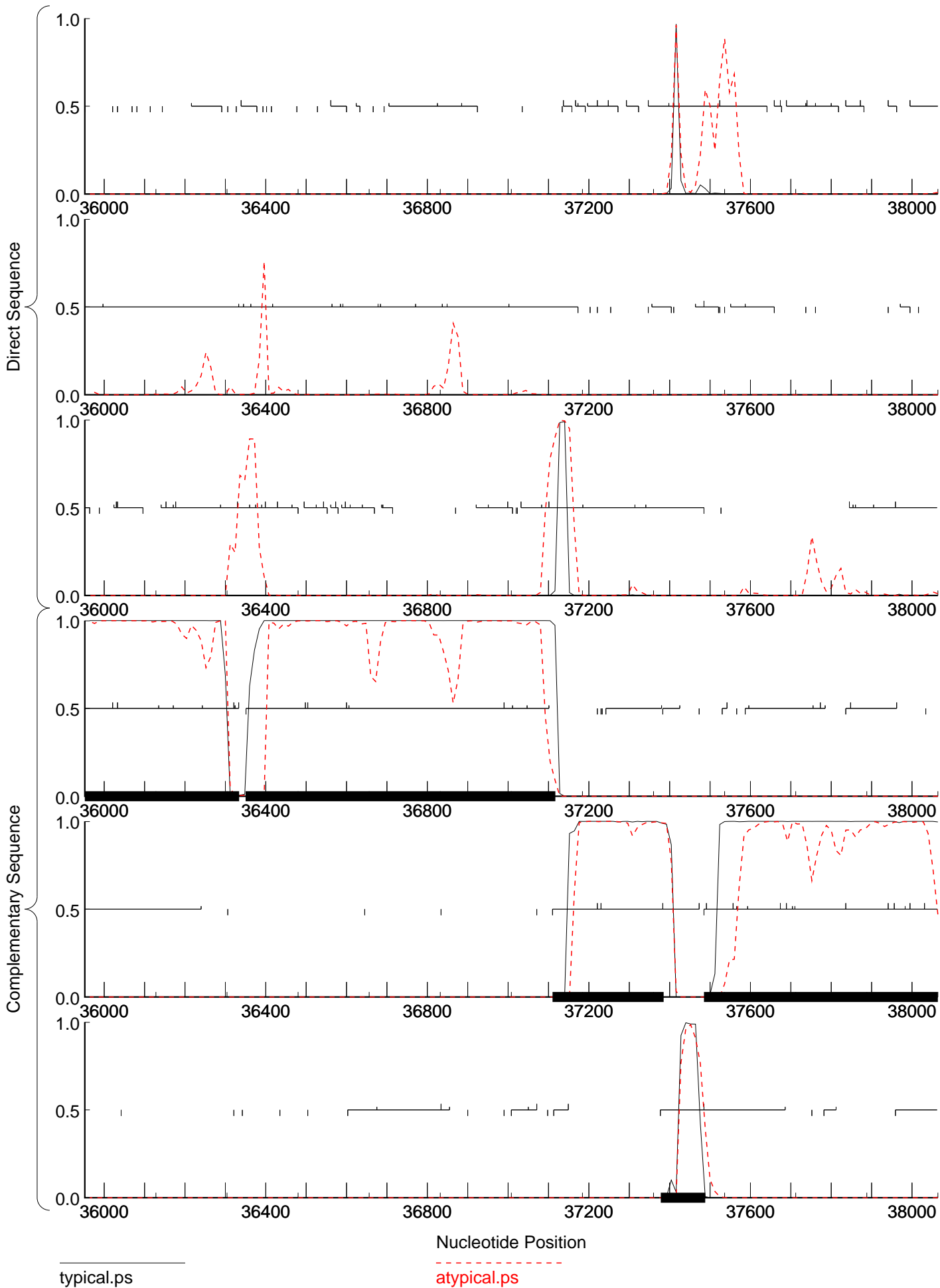
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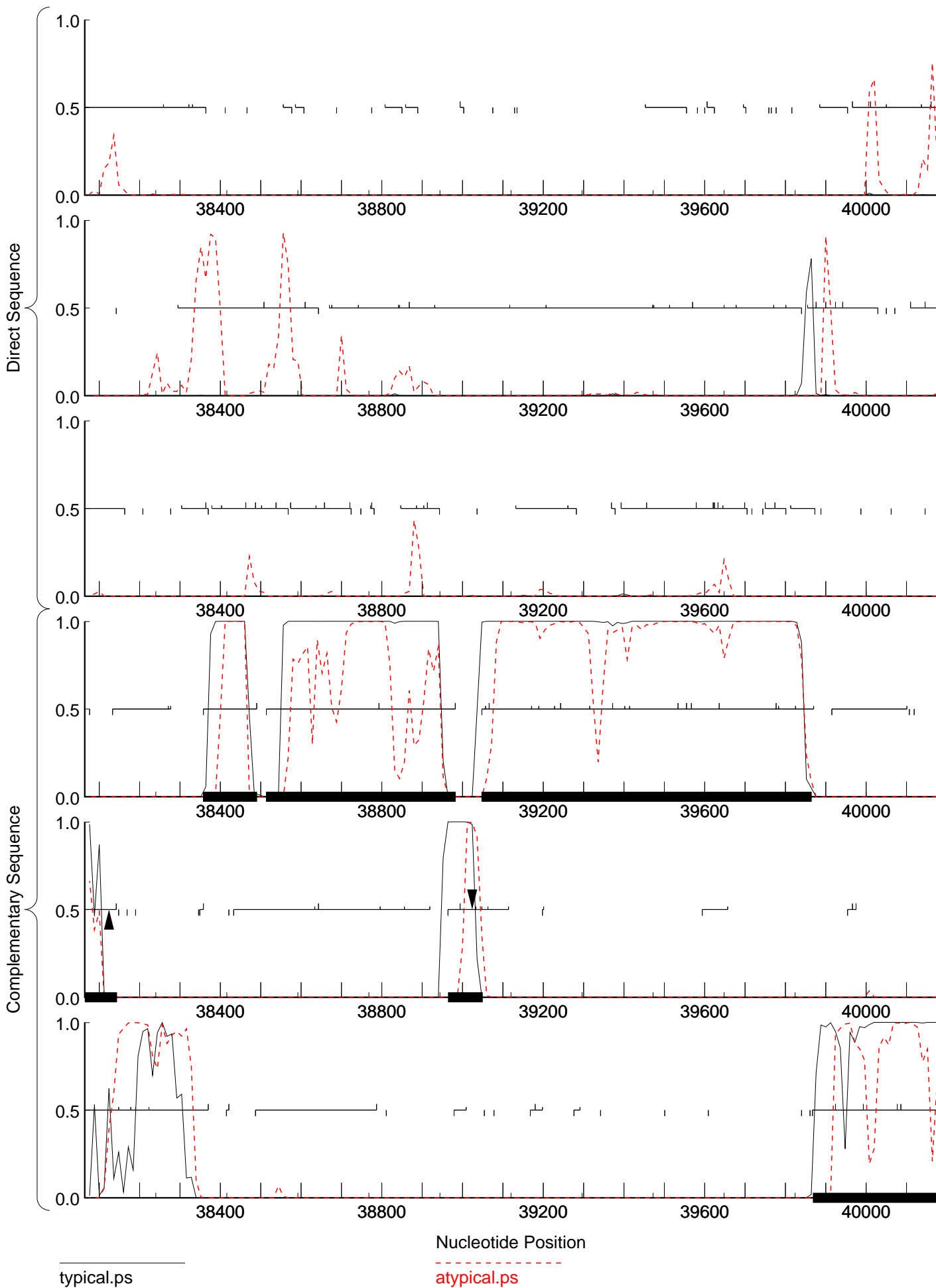
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typical.ps

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