

# GeneMark

Version 2.5p (09.08.06)

Copyright 1993 - M. Borodovsky, J. McIninch

## PROGRAM INFORMATION

Sequence : Mycobacterium phage Prithvi complete sequence, 60311 bp including 11-base 3' overhang (CTCG)  
Analysis Date : 1/18/18 at 20:30:23  
Pages : 30  
Sequence Length : 60311 bp  
GC Content : 66.77%

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,

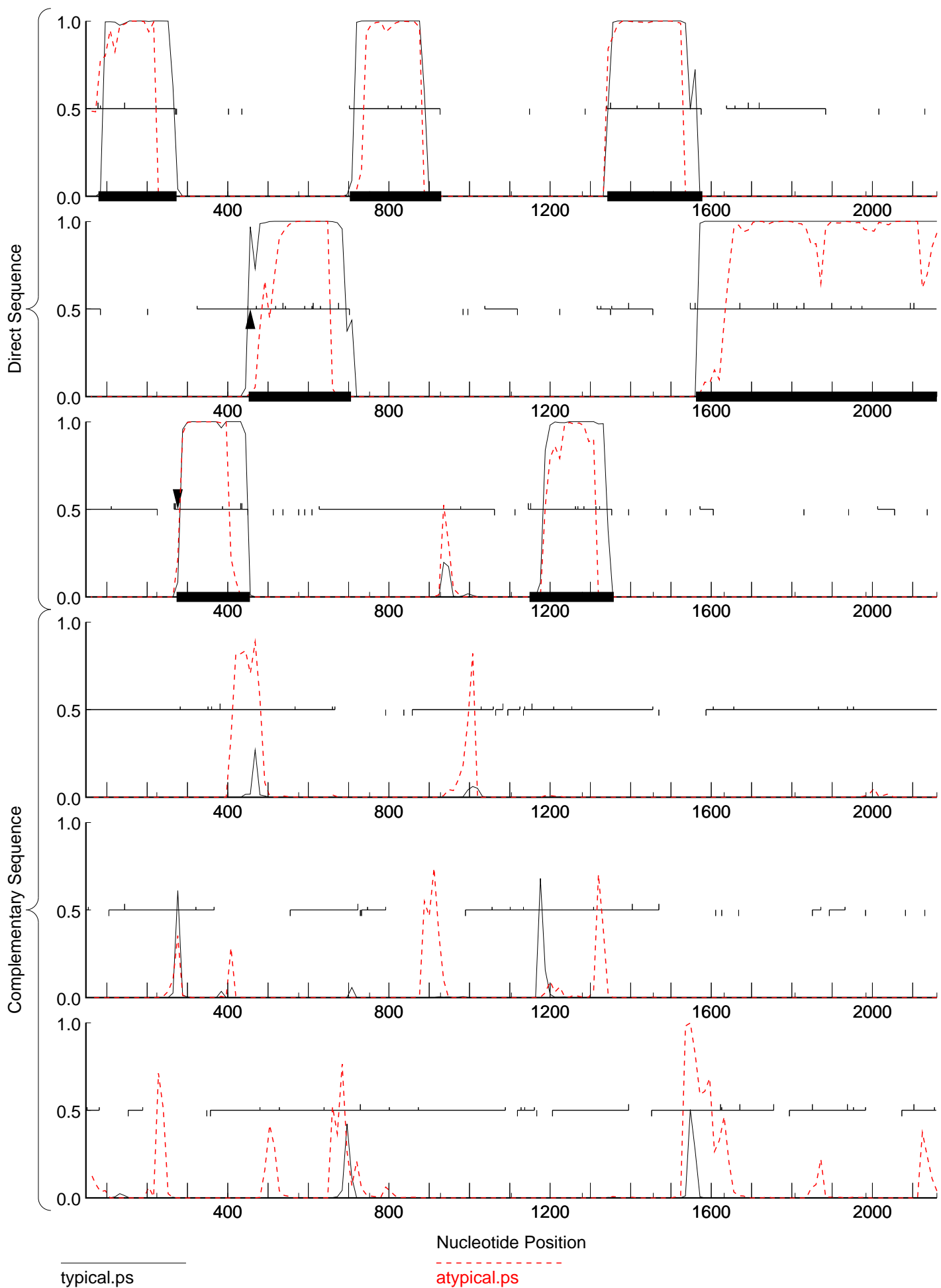
## MATRIX INFORMATION

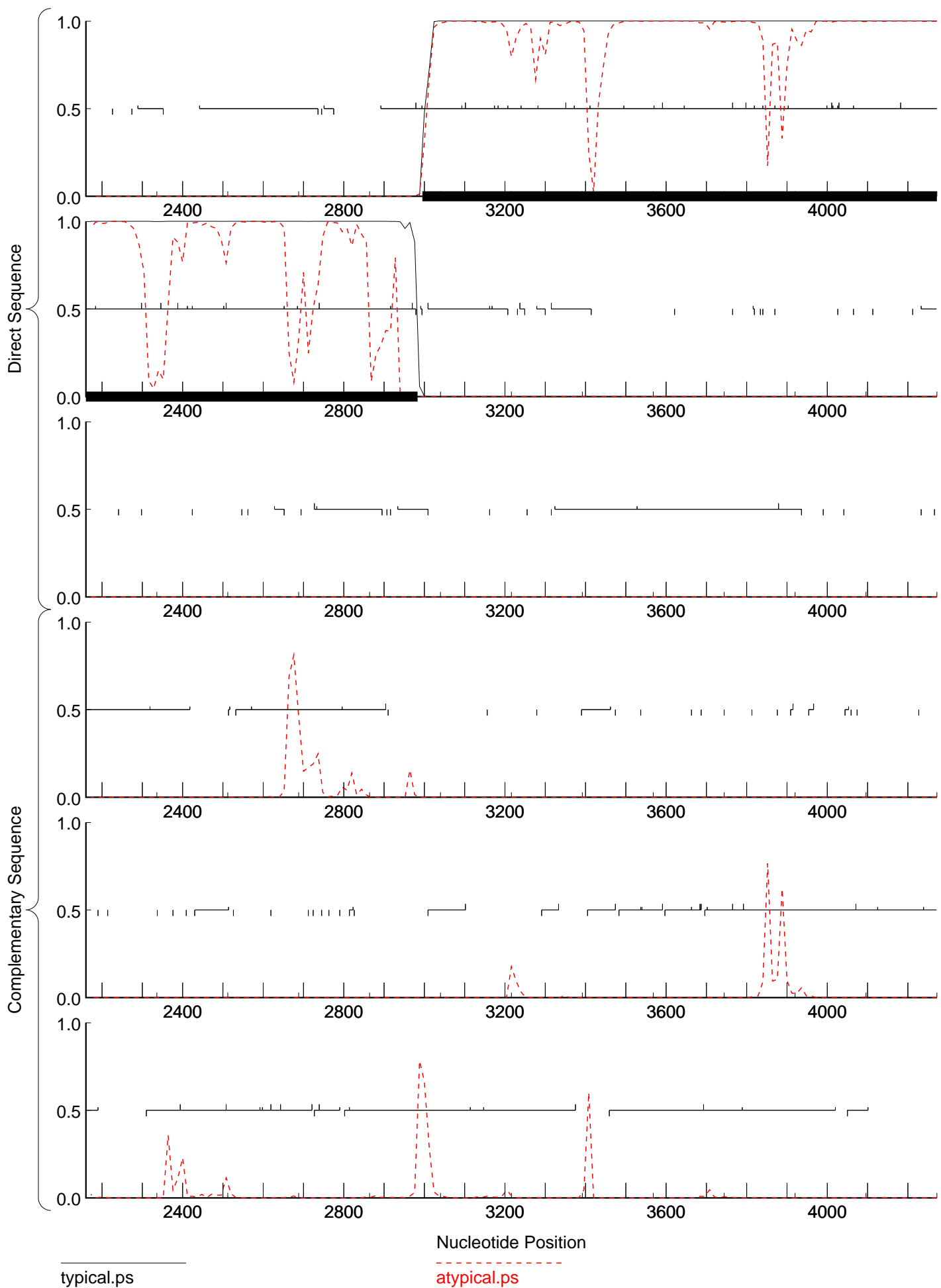
Matrix : -  
Author : -  
Order : 4

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

### Matrix notes & comments

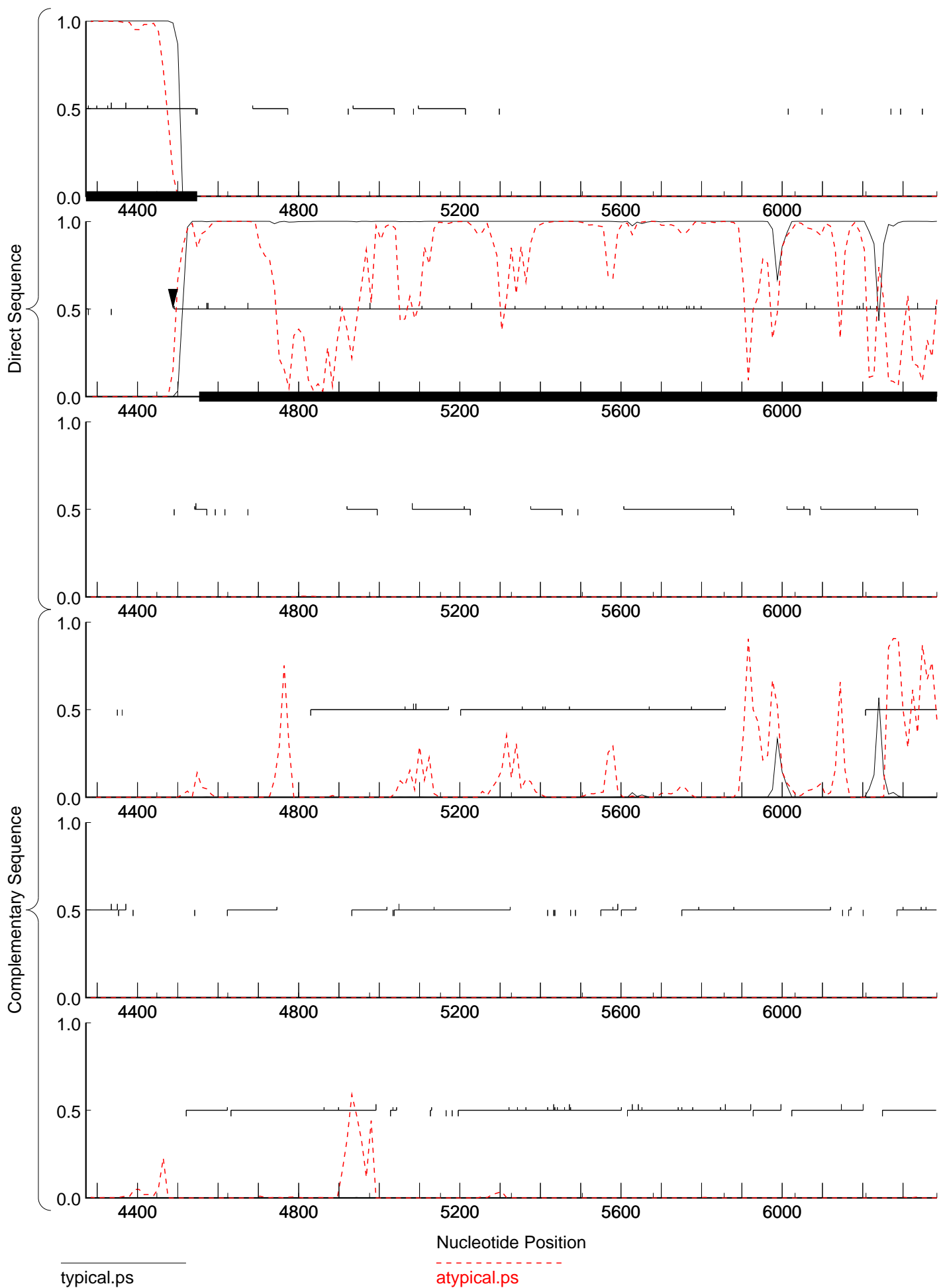
Training set derived by GeneMarkS, 4.30 October 2014  
Thu Jan 18 20:30:23 2018

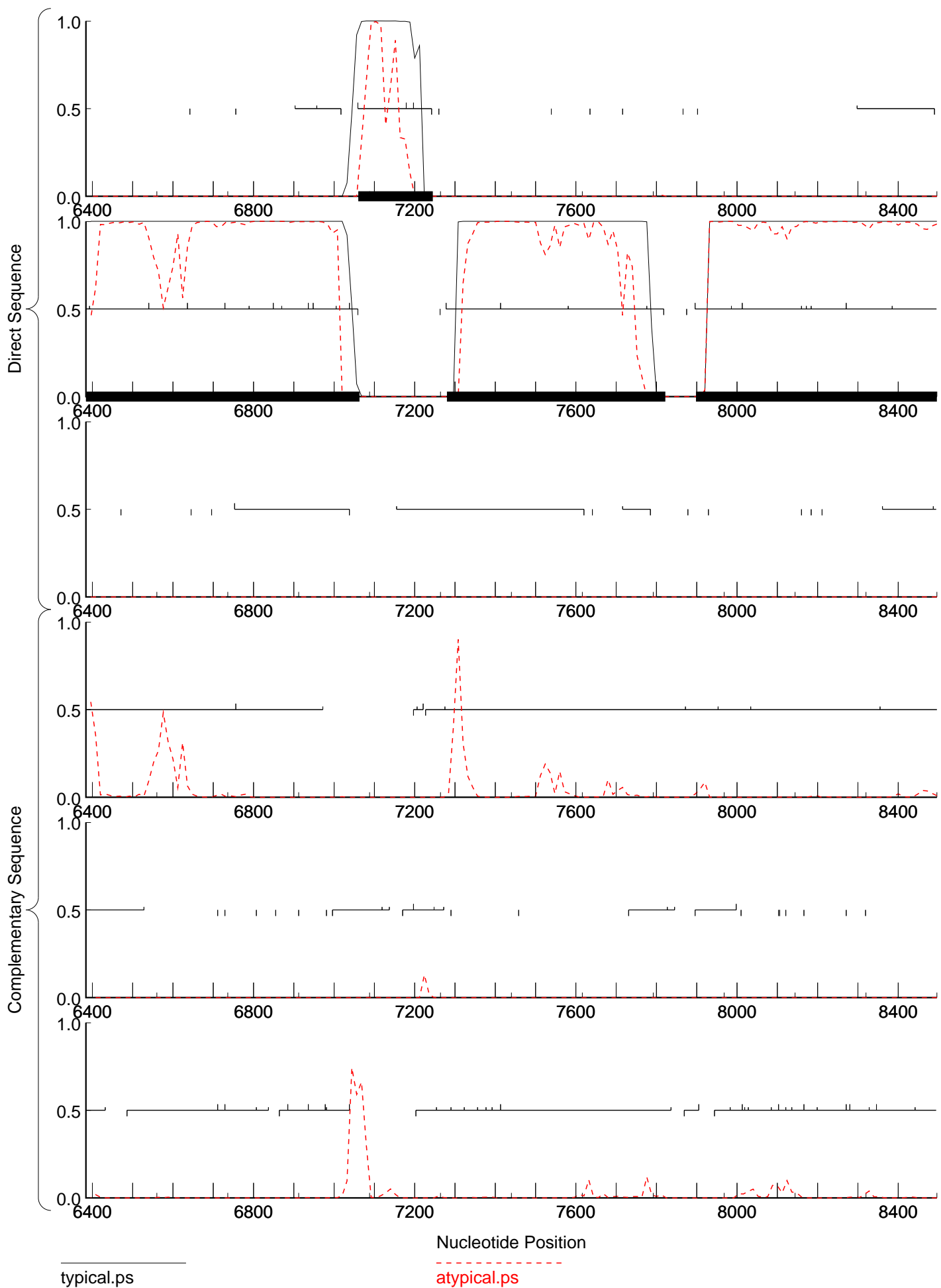




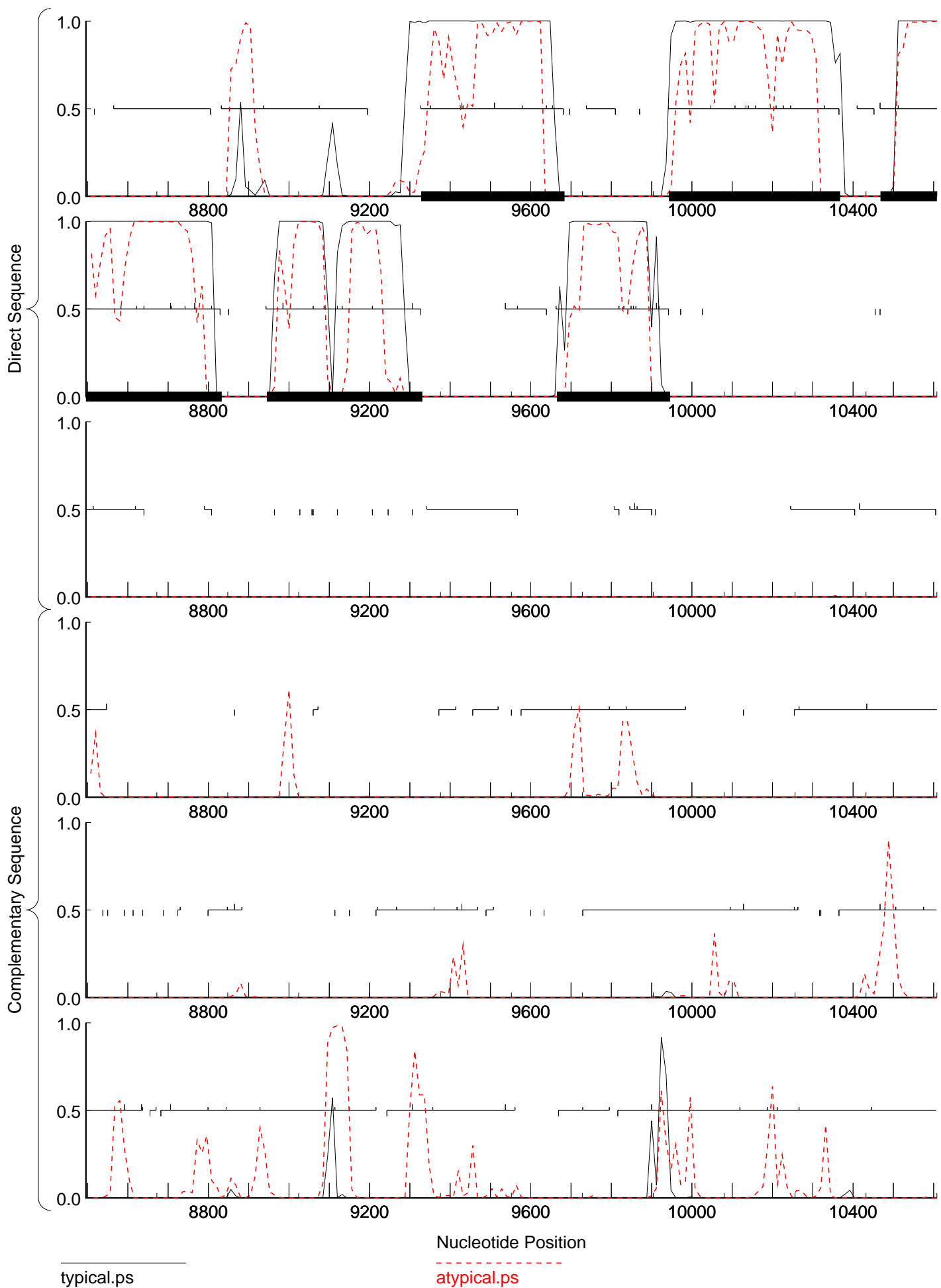
typical.ps

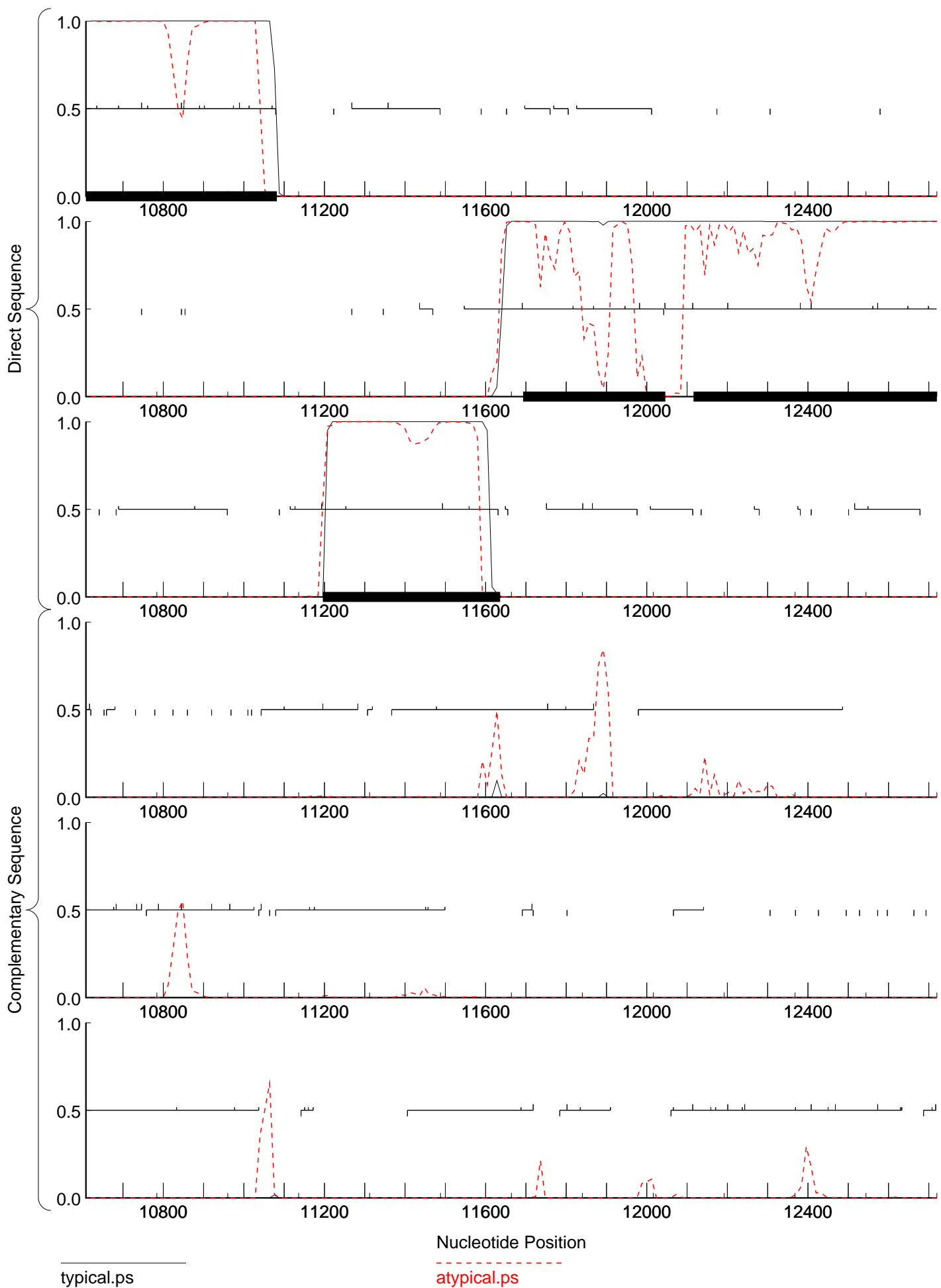
atypical.ps

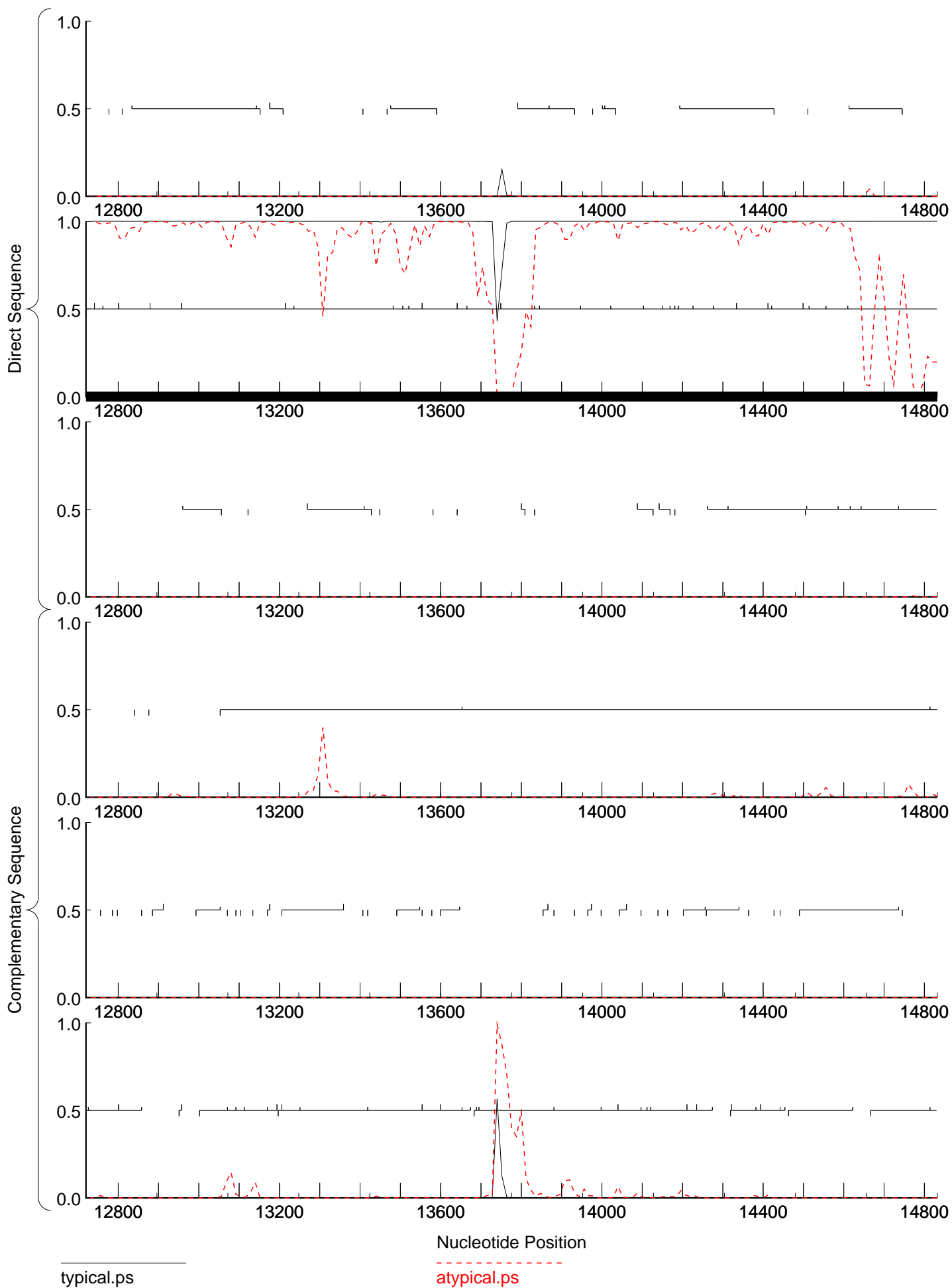




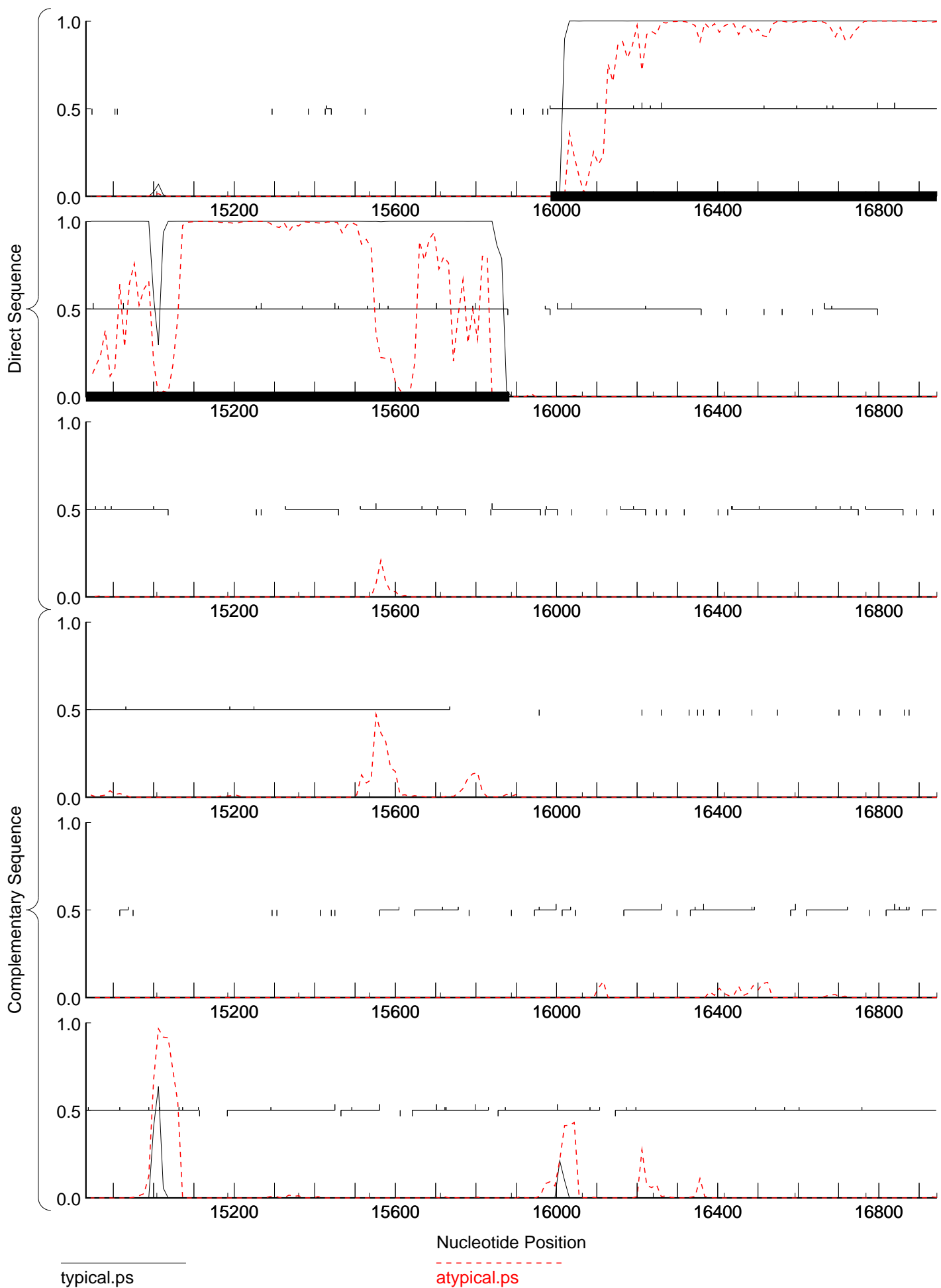
GeneMark.hmm prediction

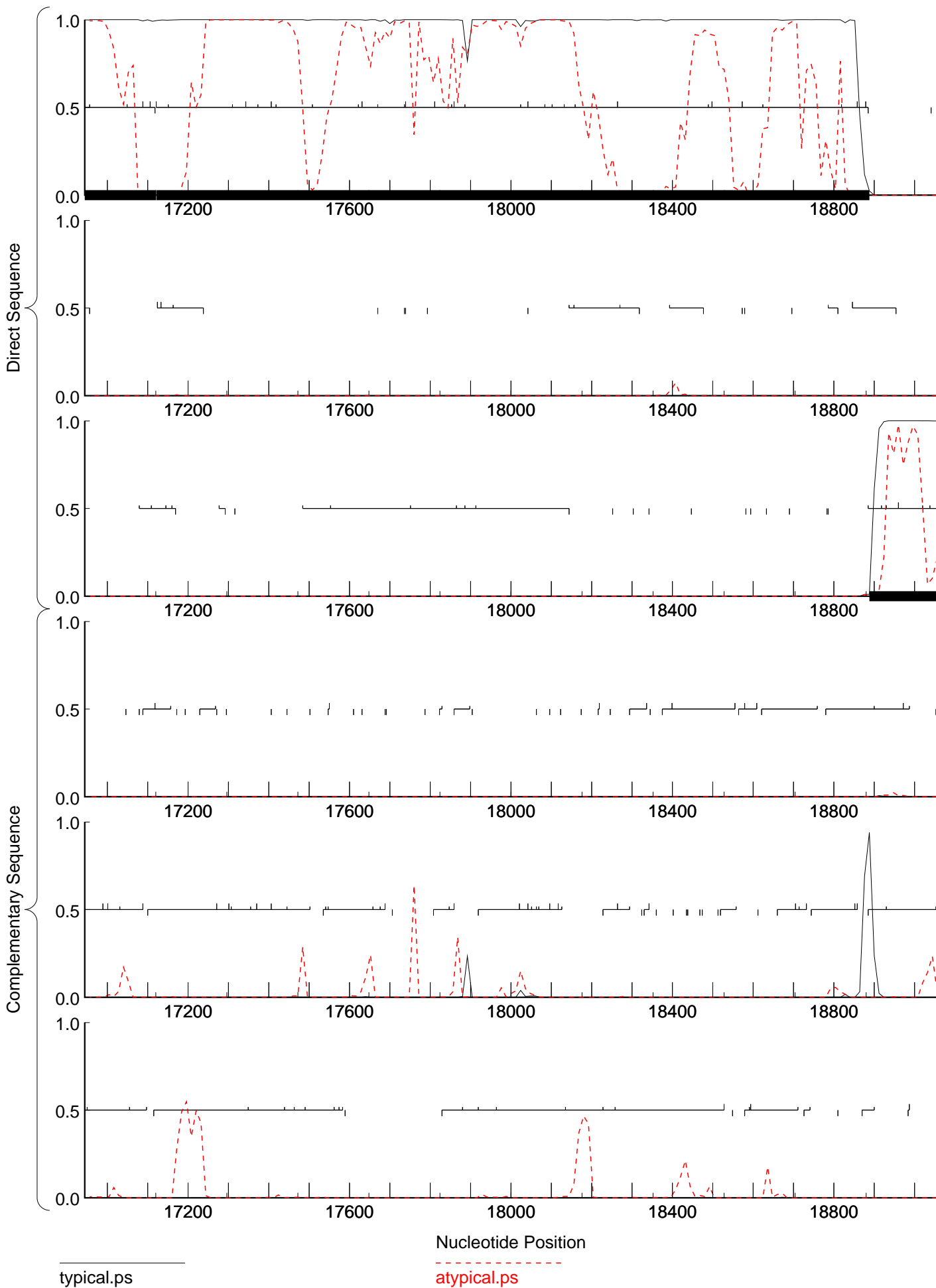






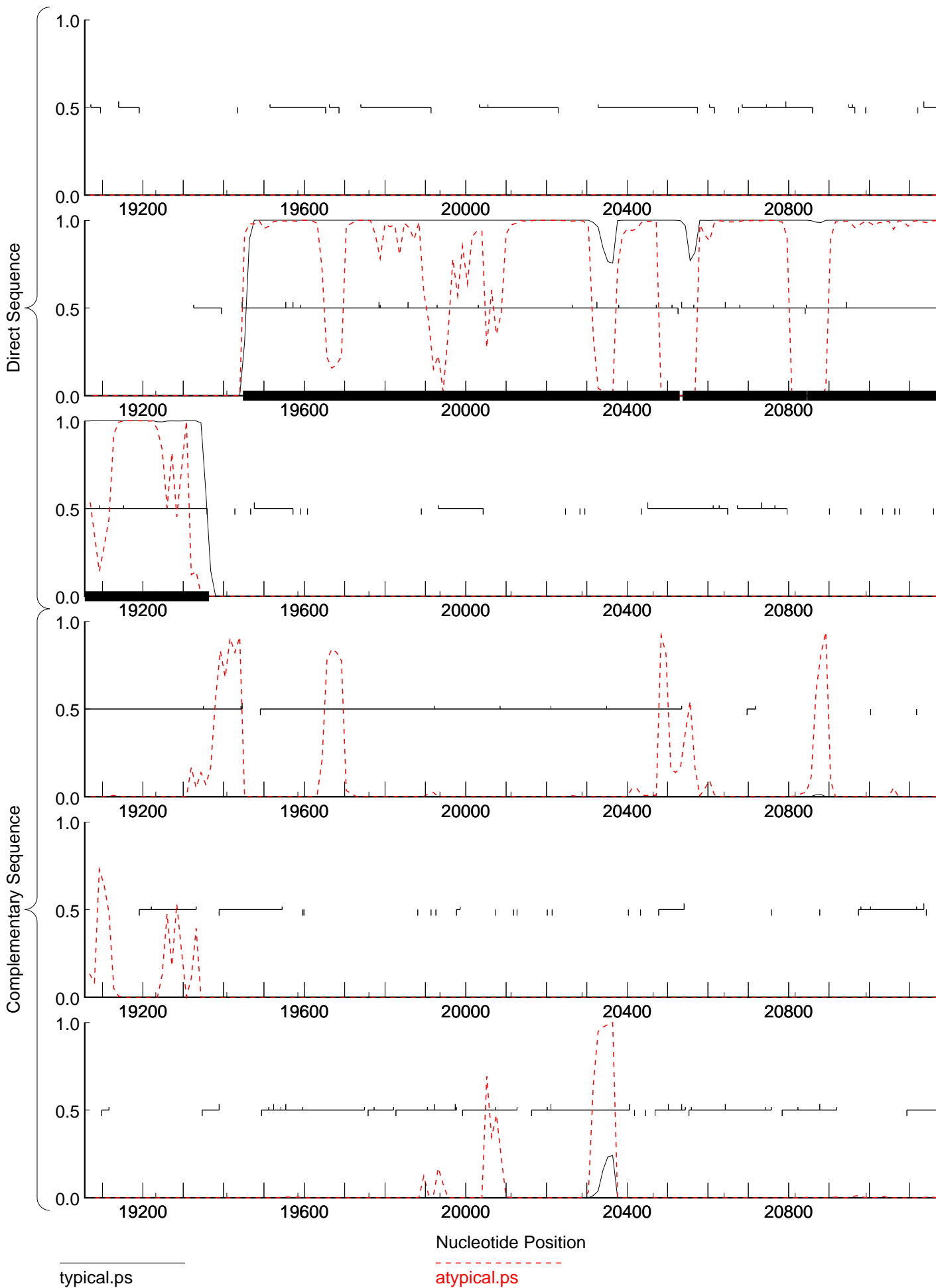


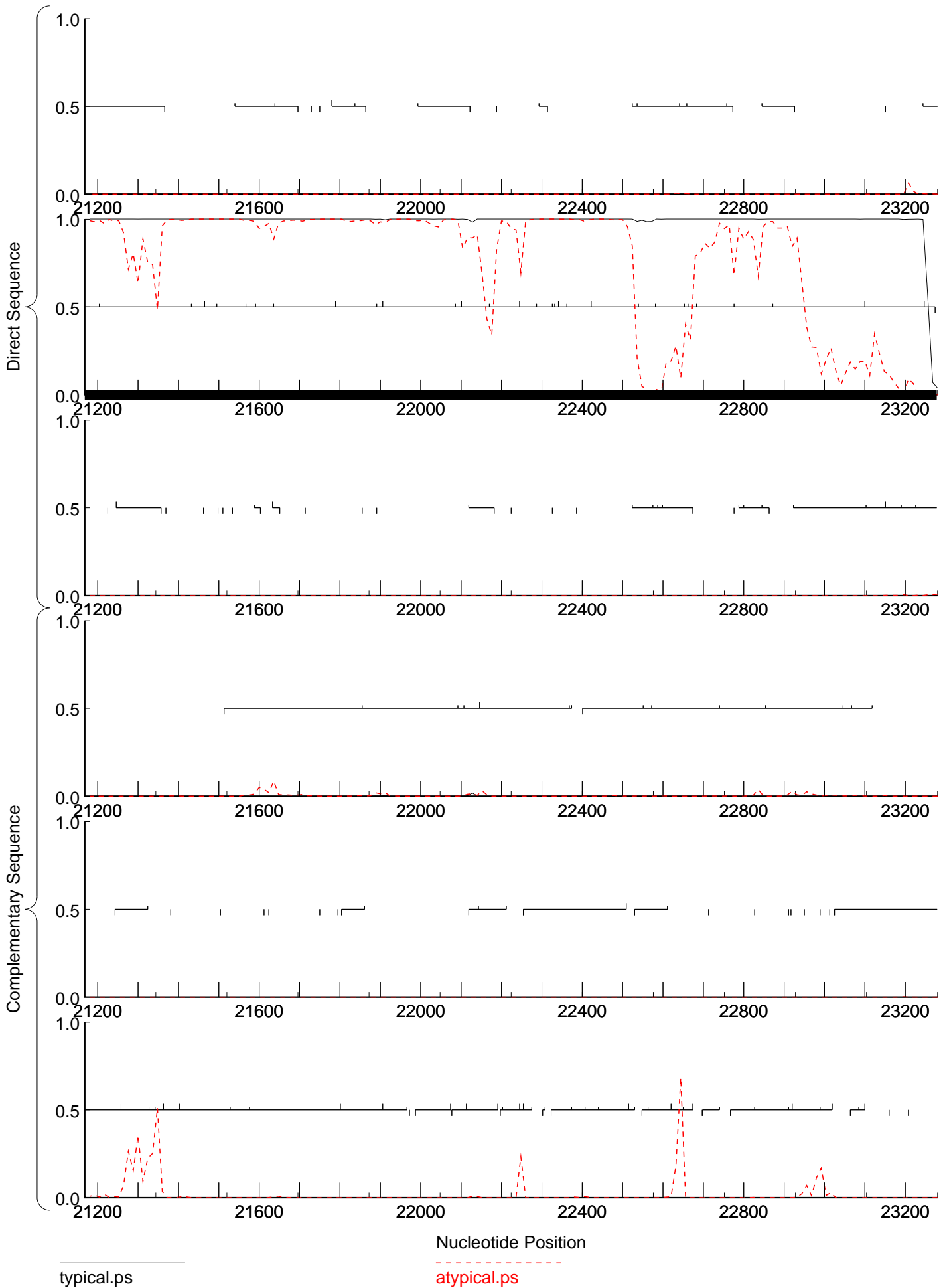


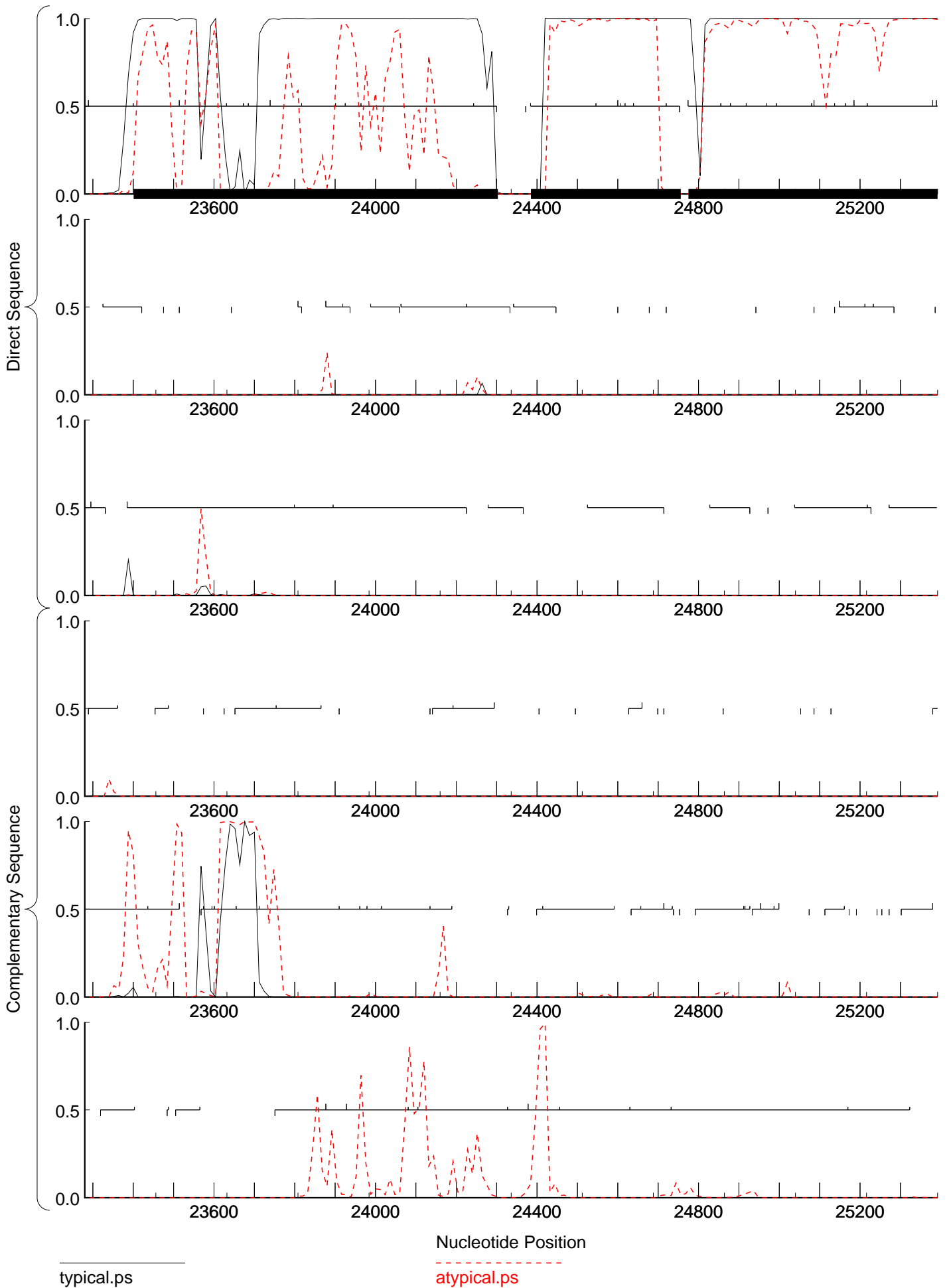


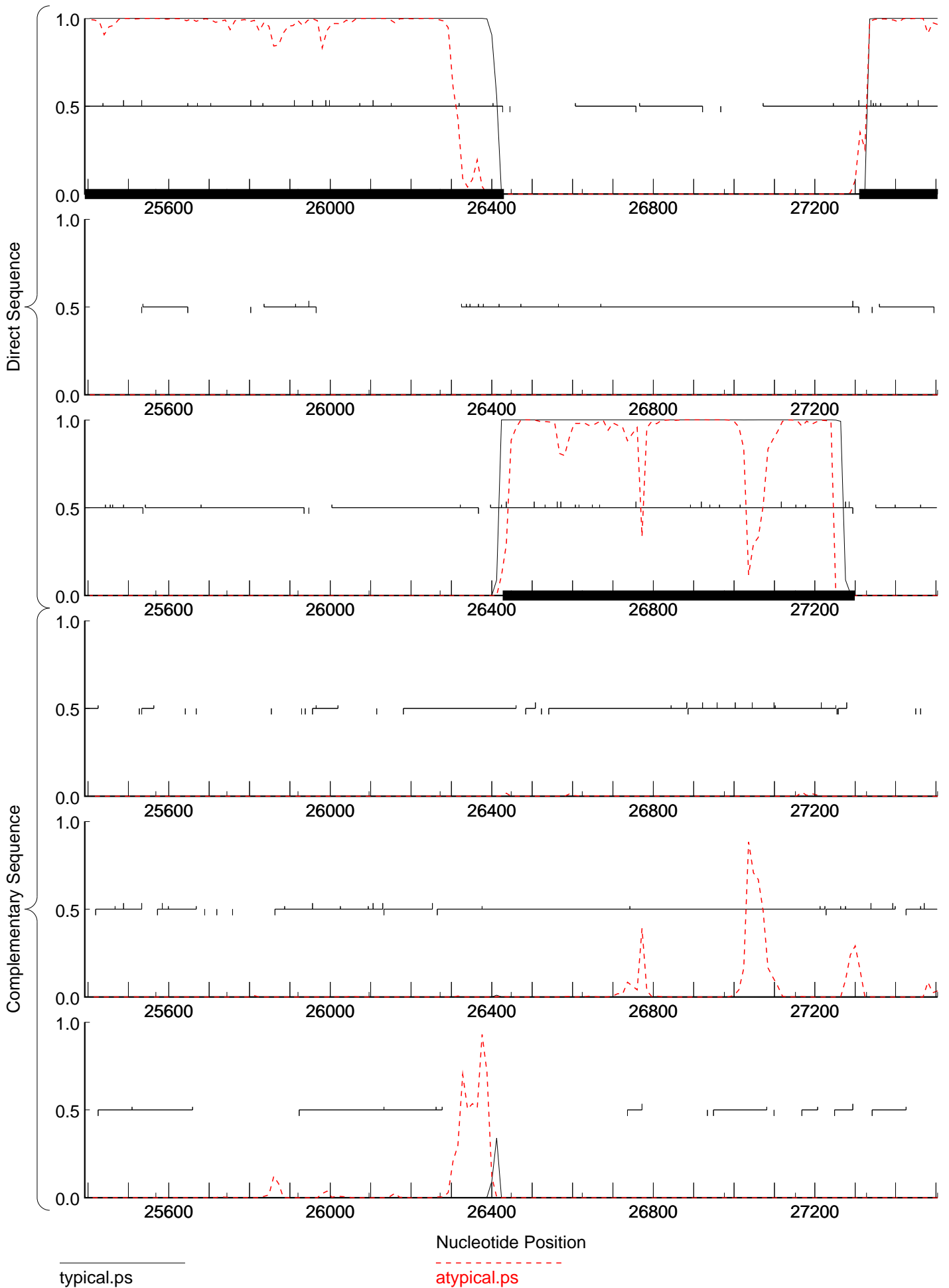
typical.ps

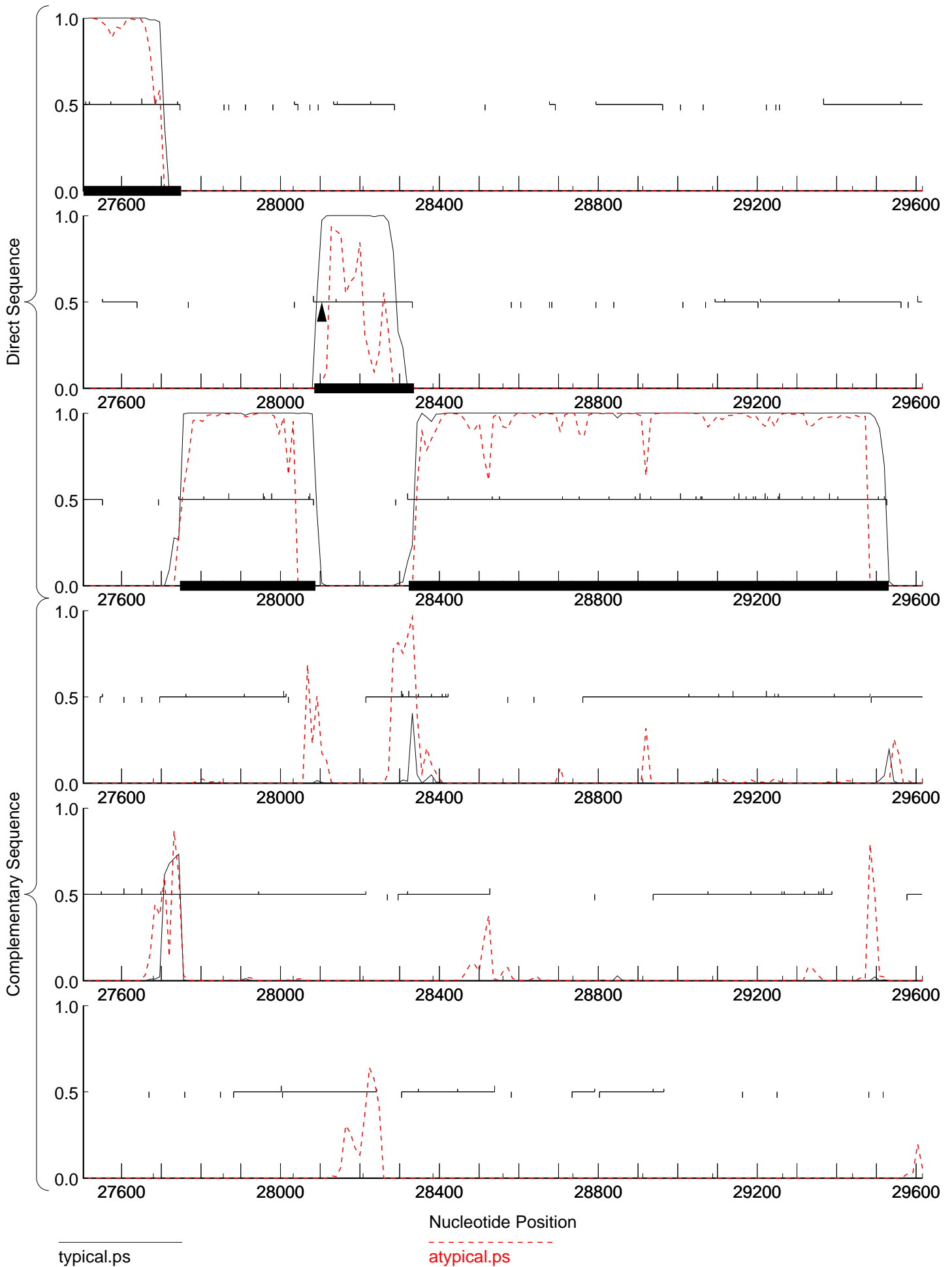
atypical.ps

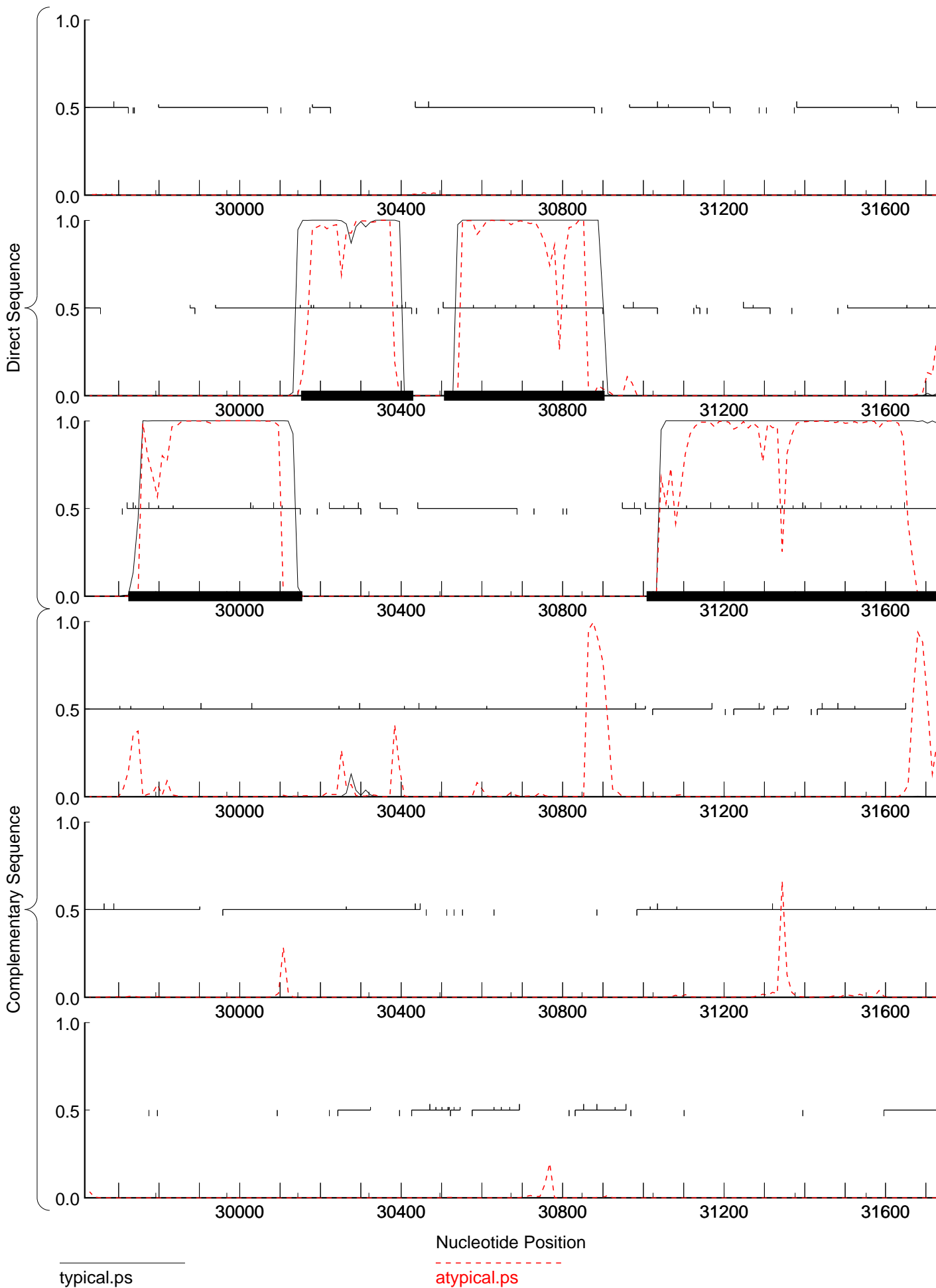




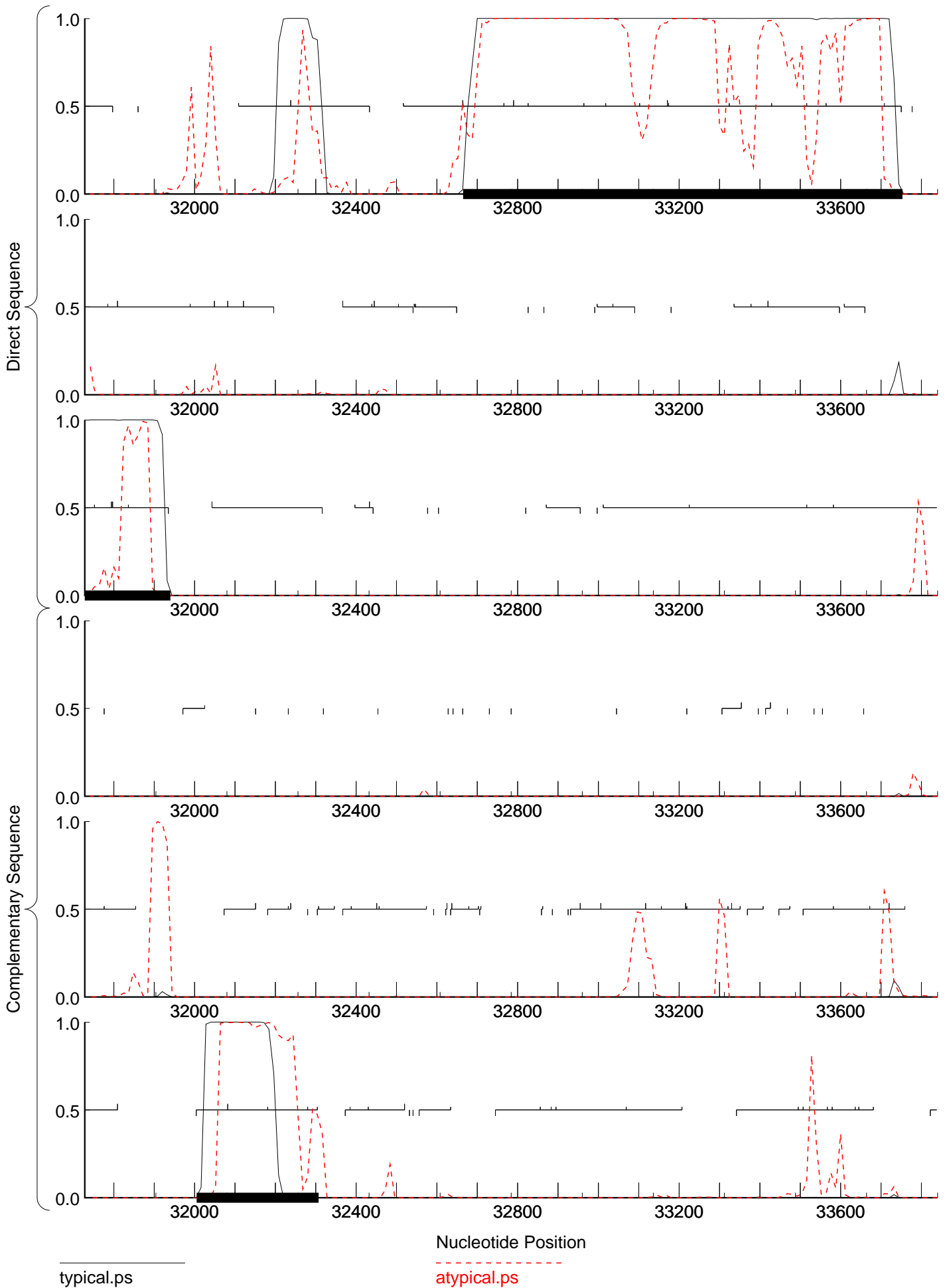


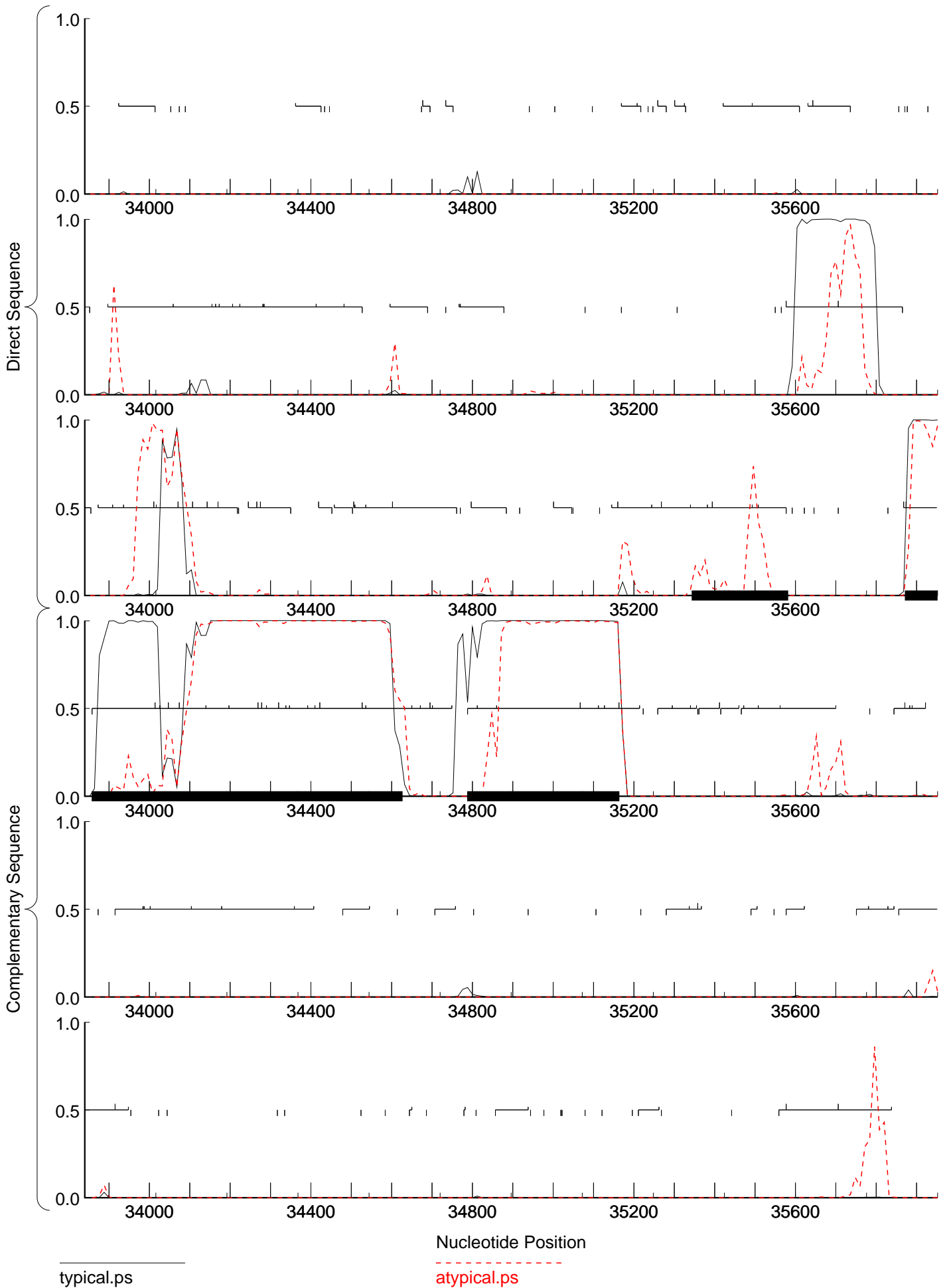


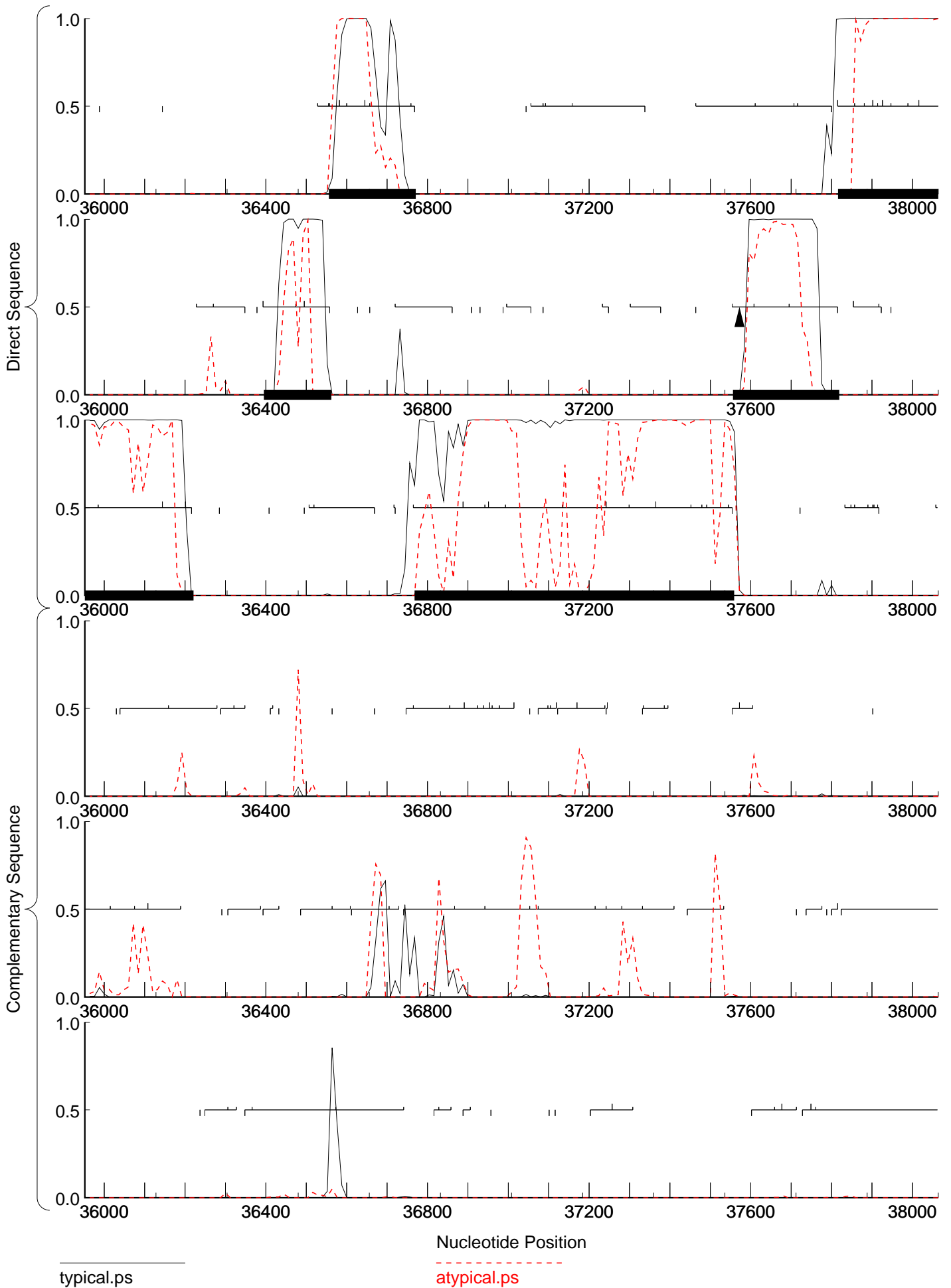


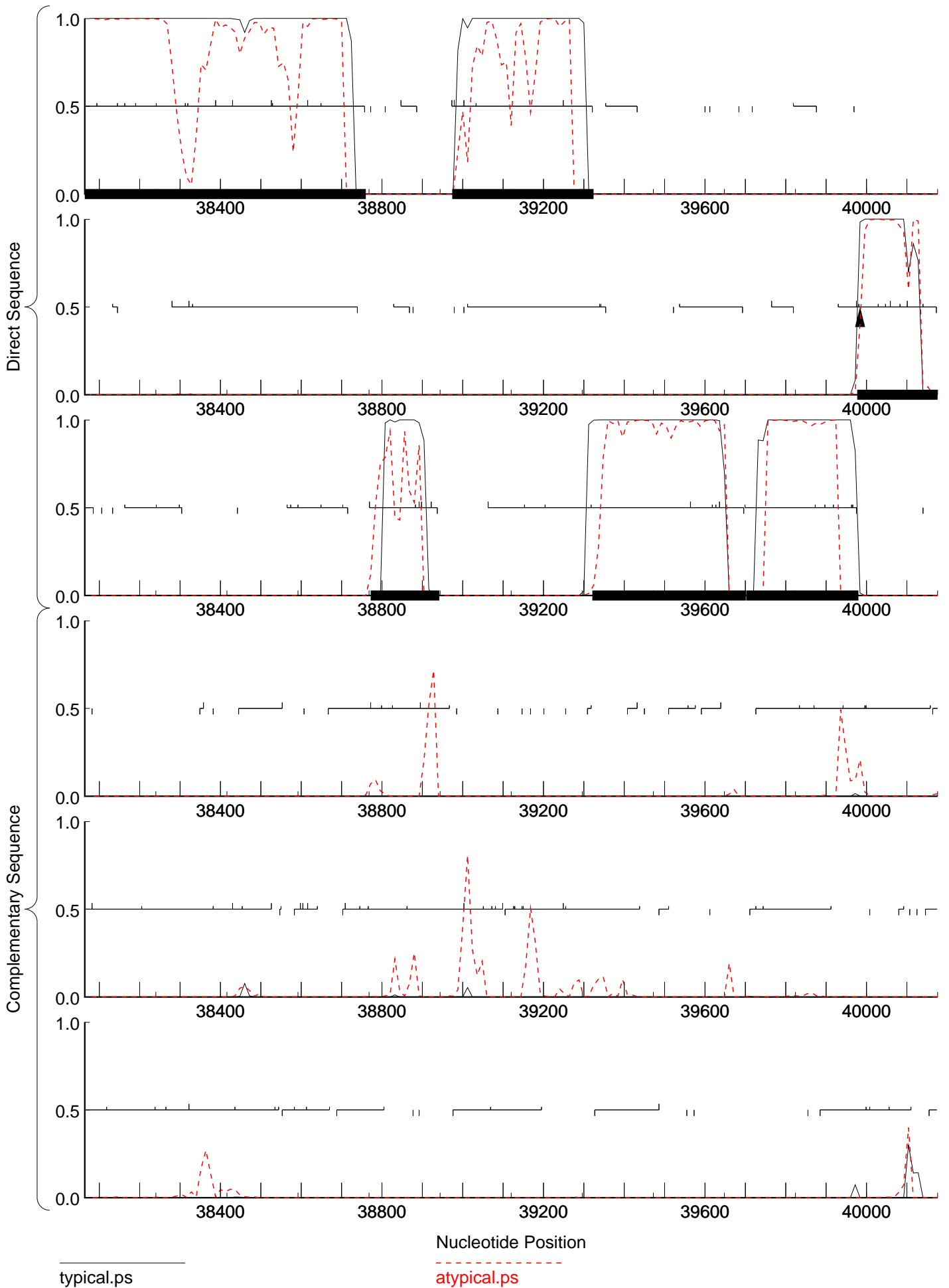






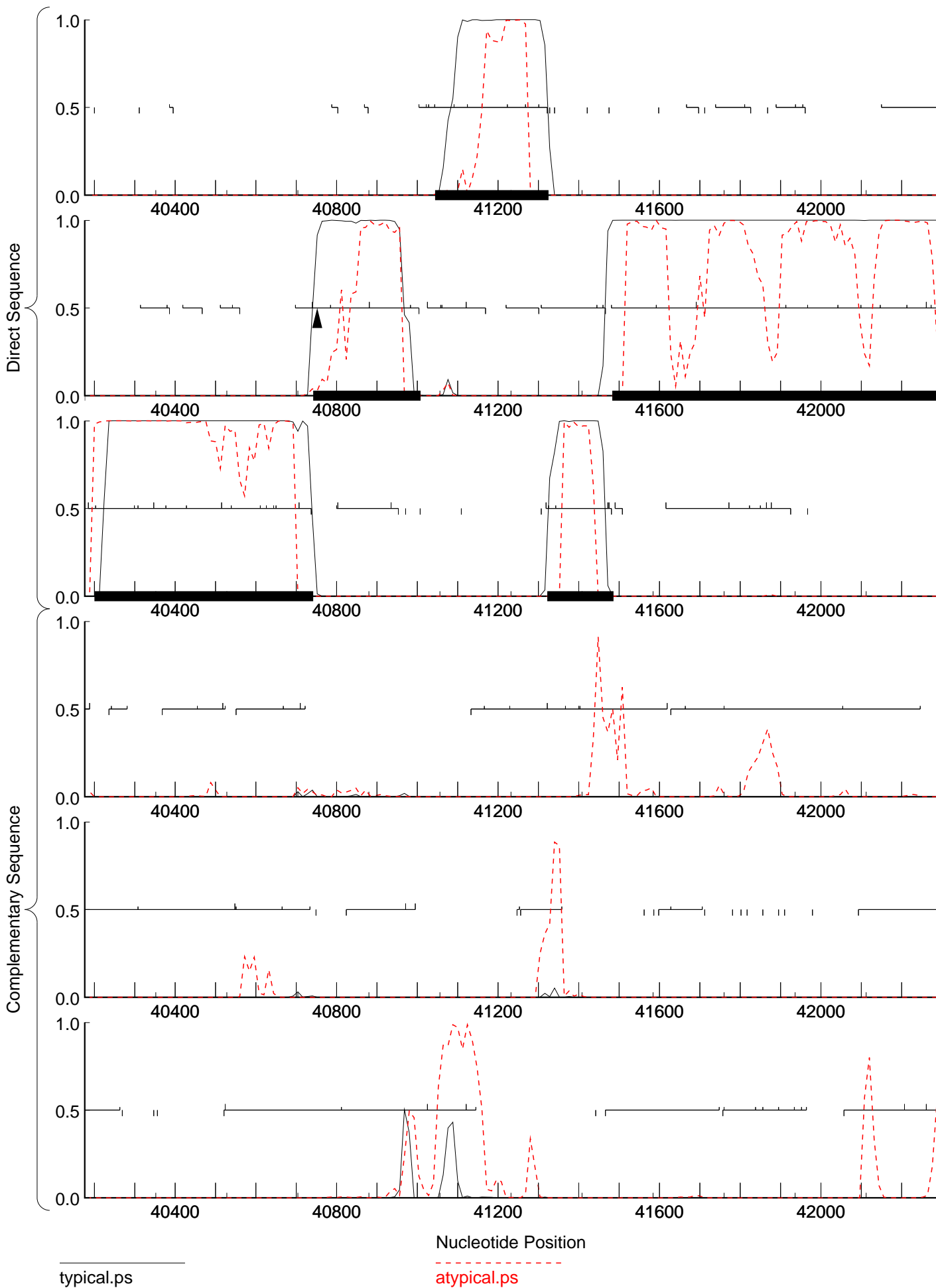


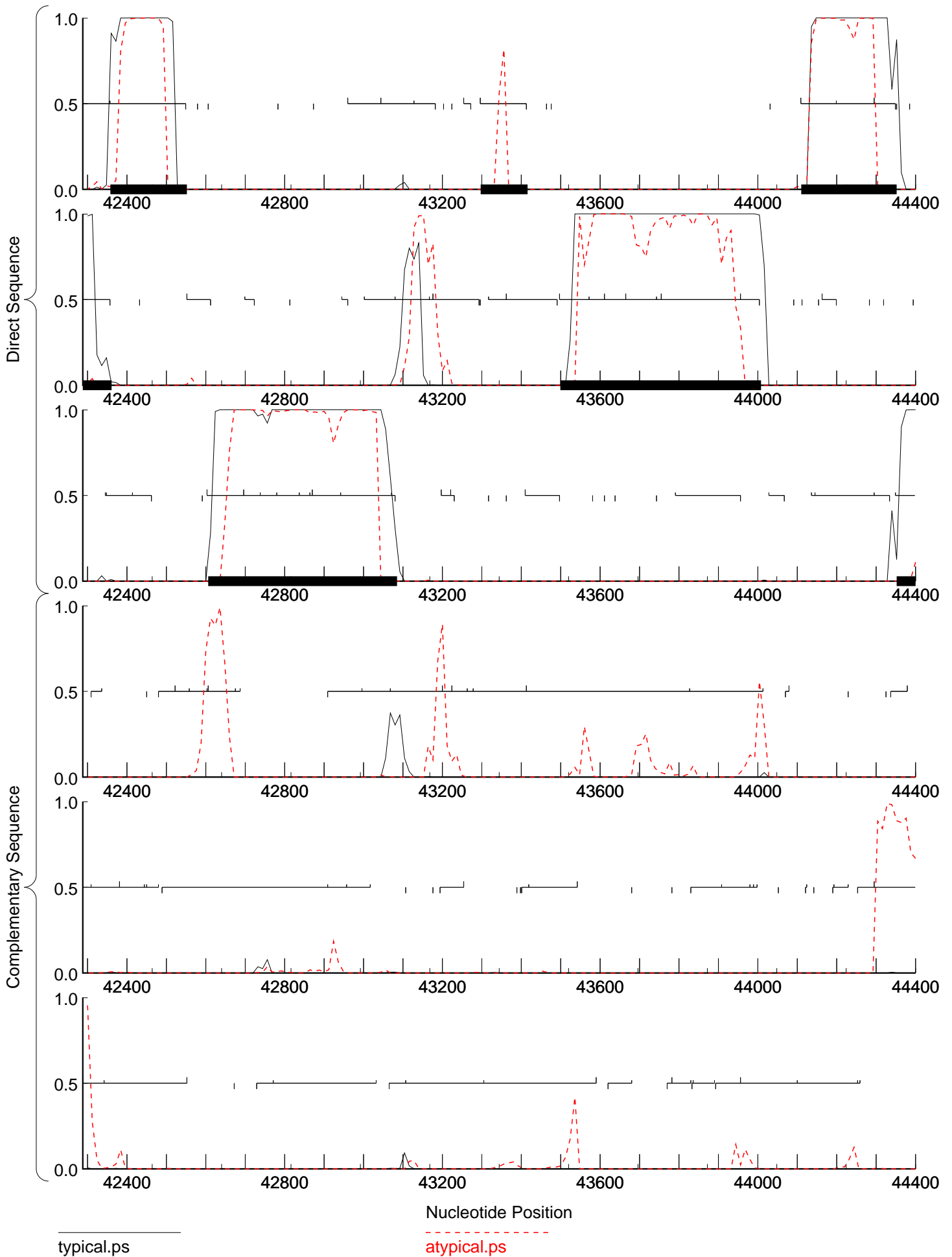




typical.ps

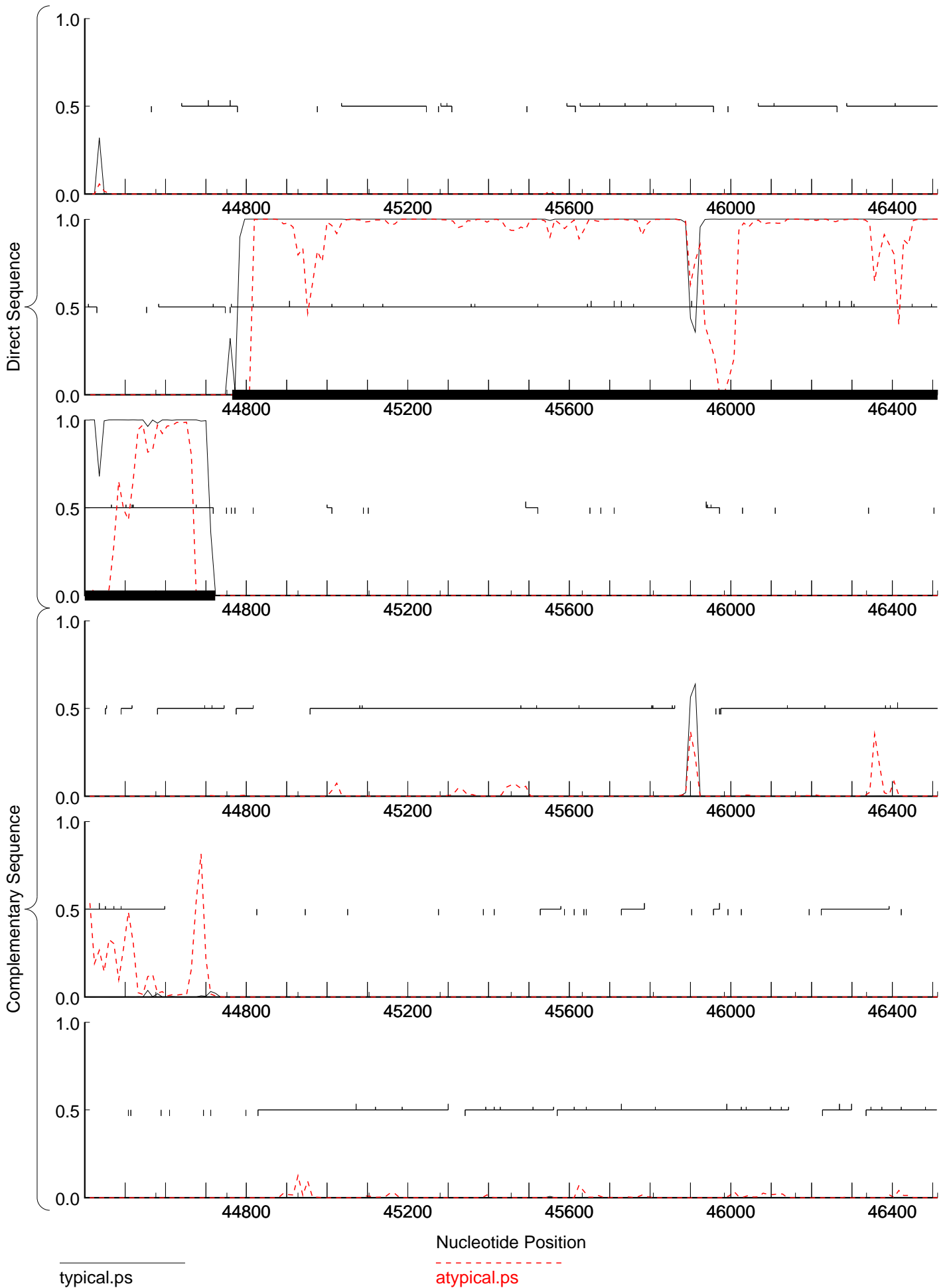
atypical.ps

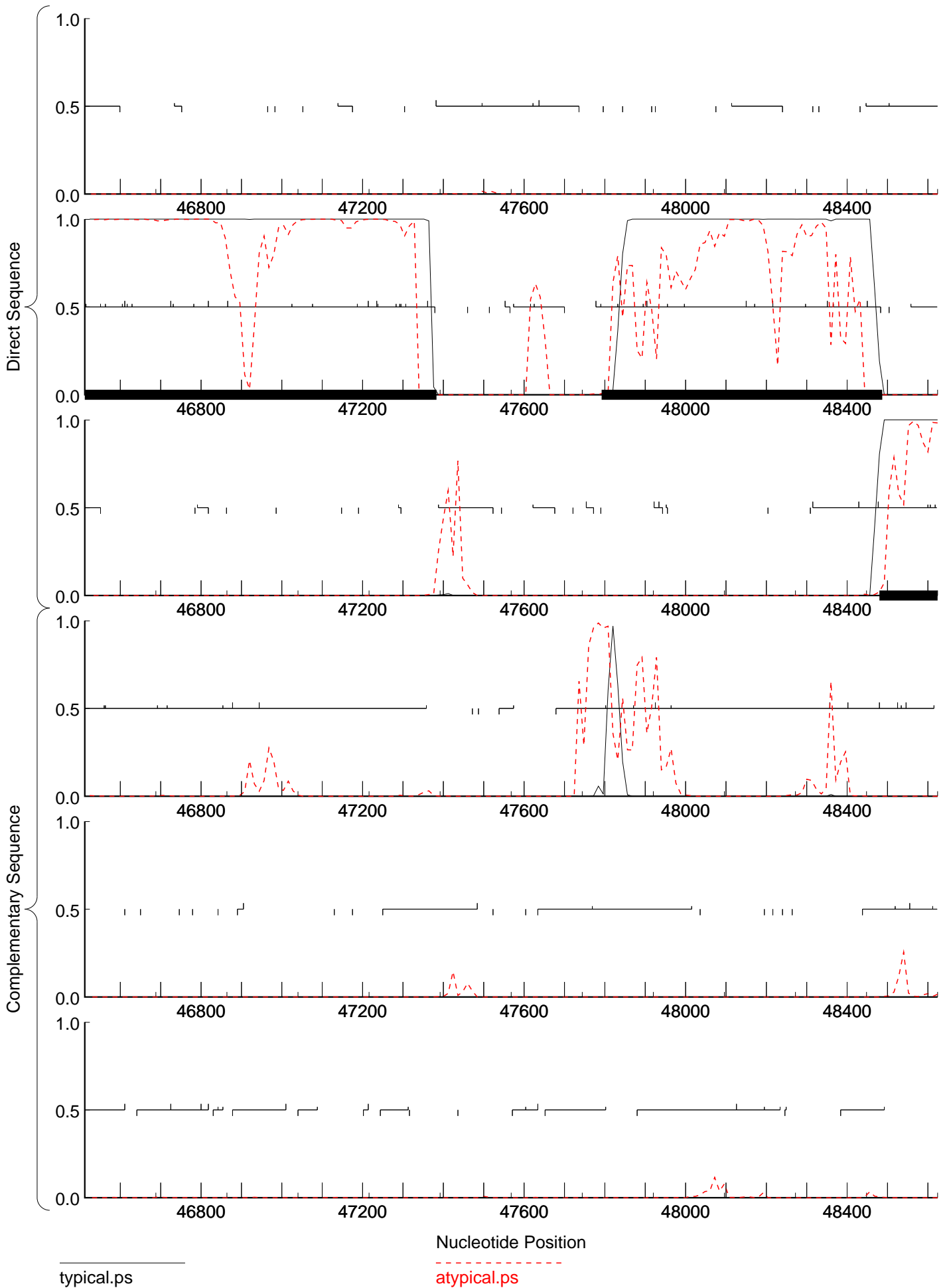




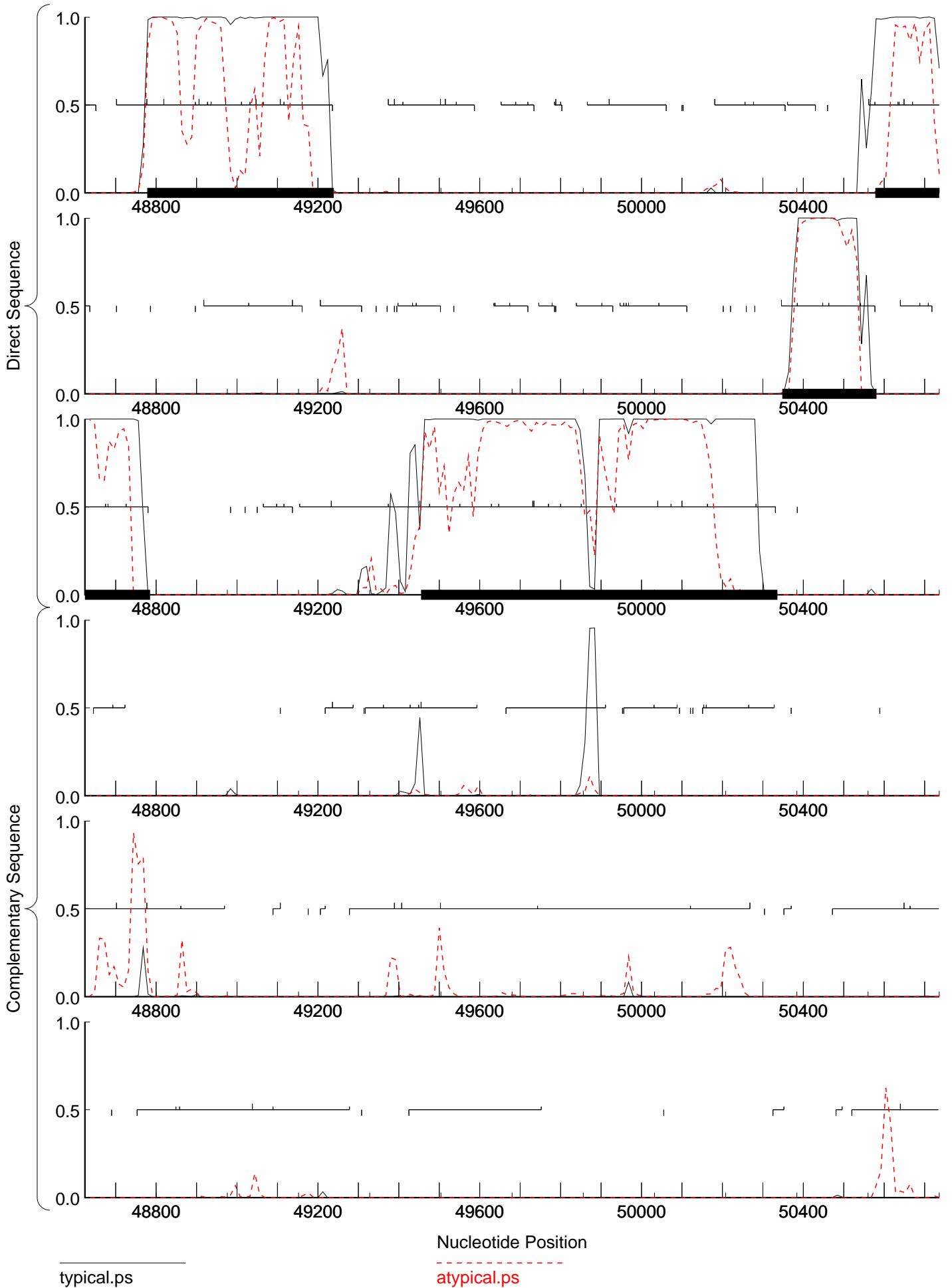
typical.ps

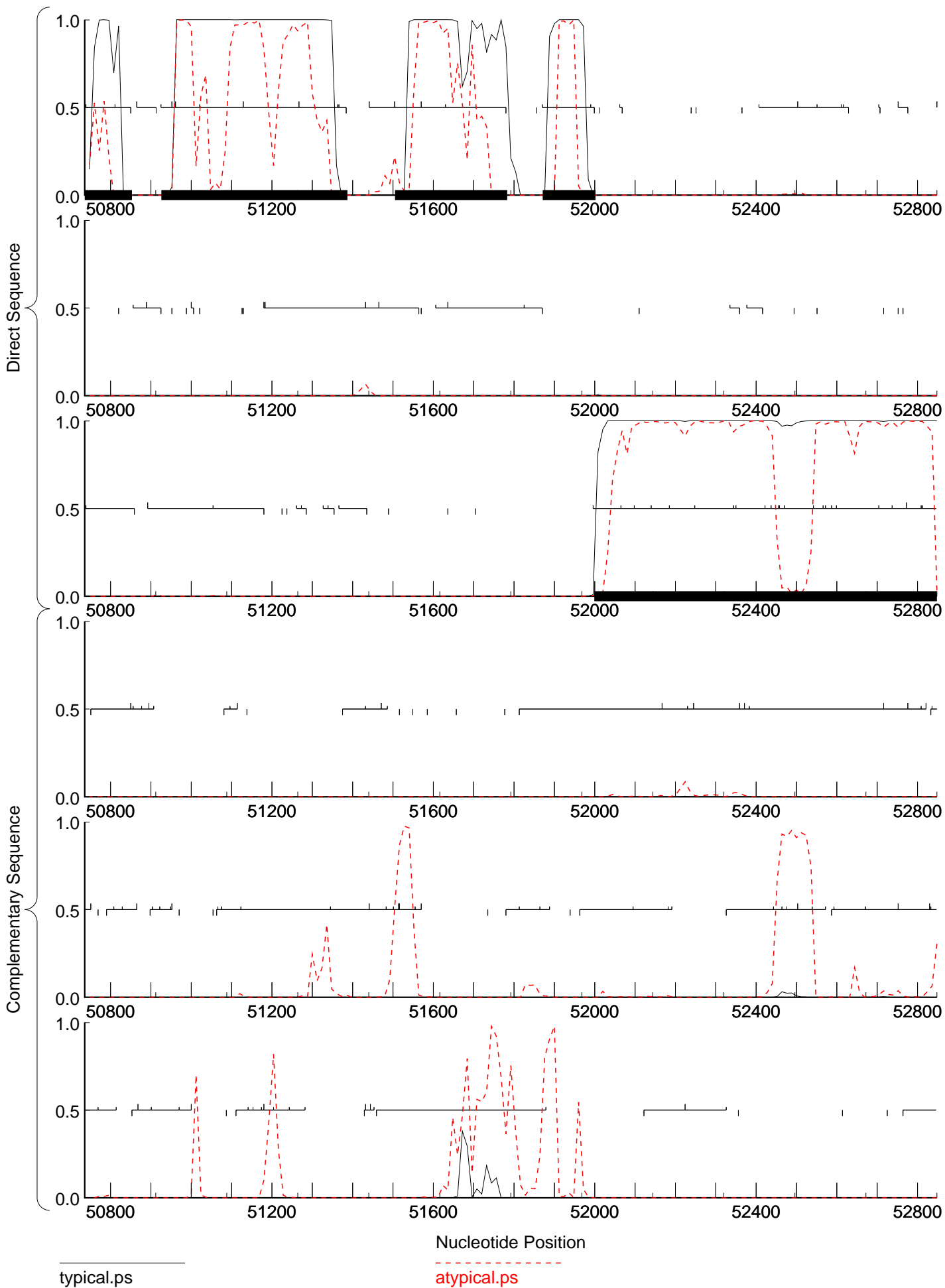
atypical.ps

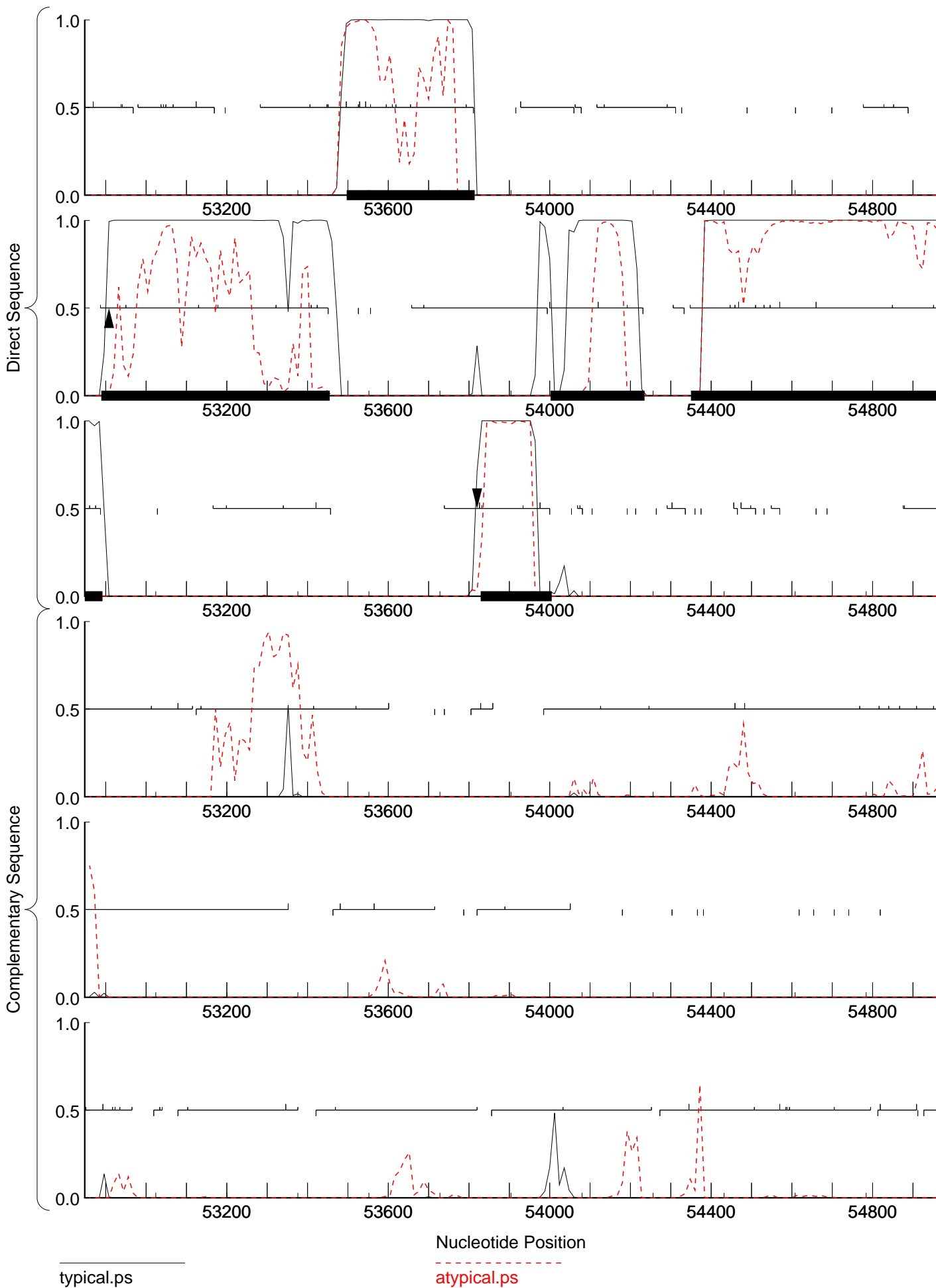


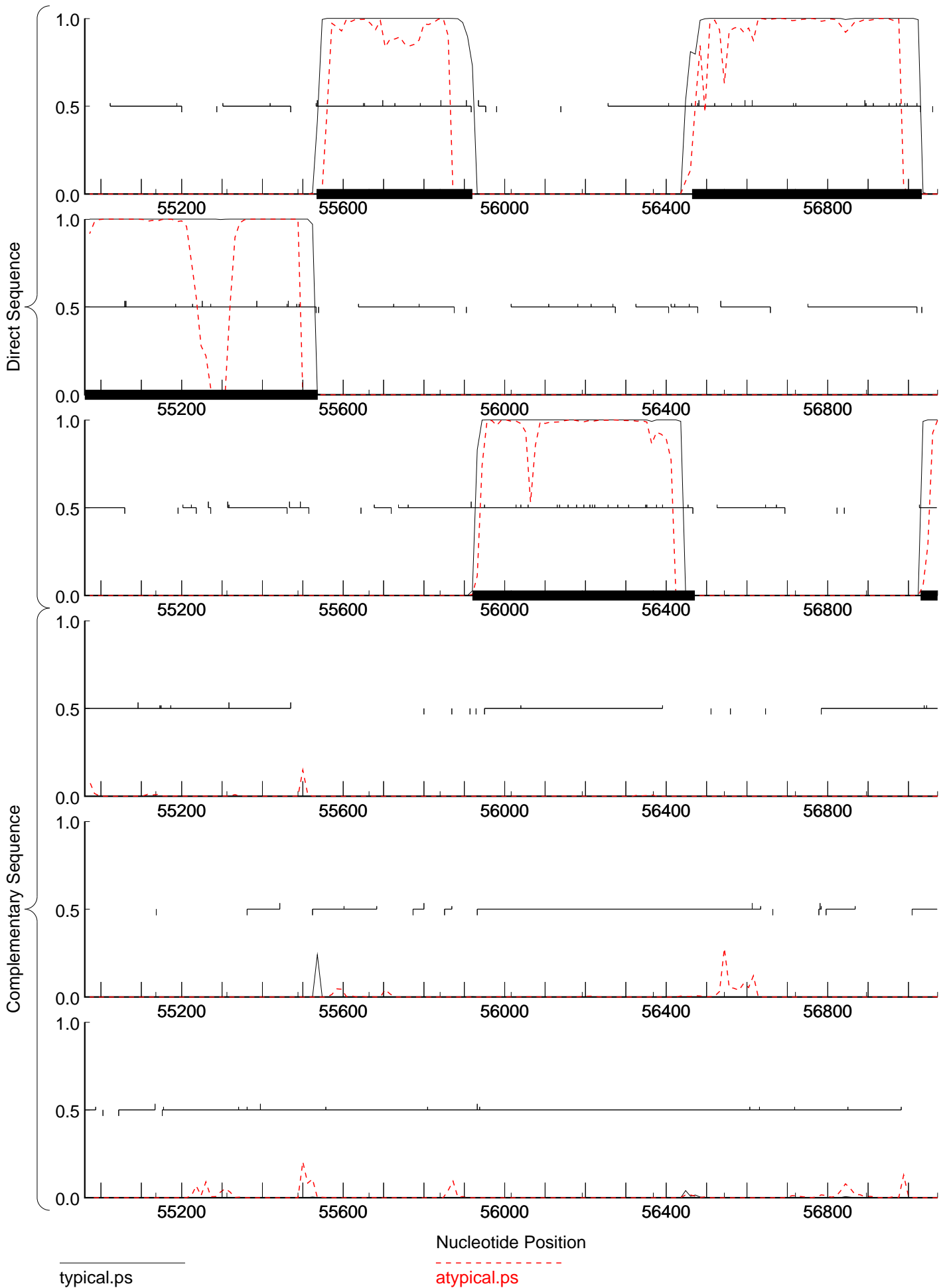


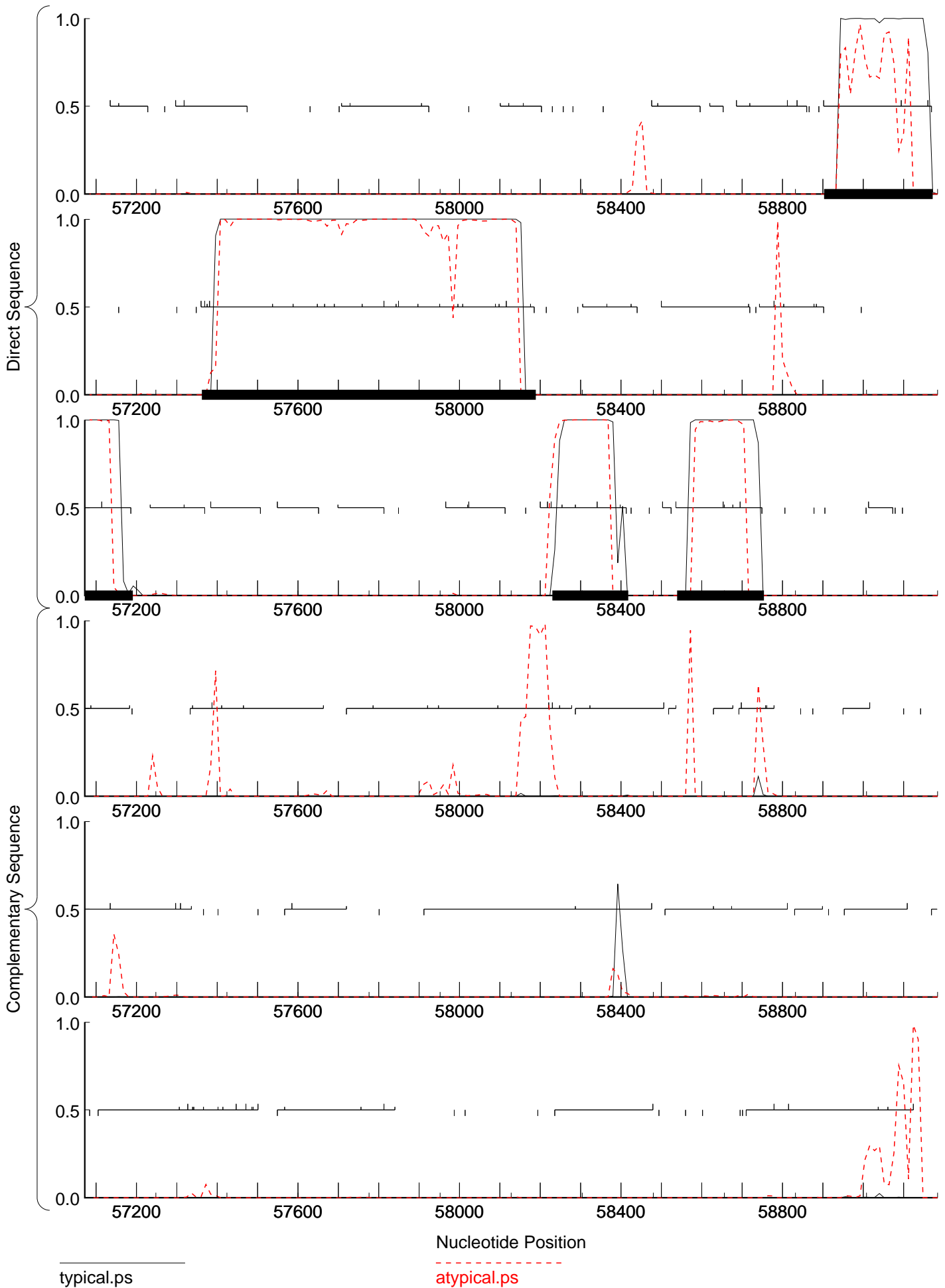


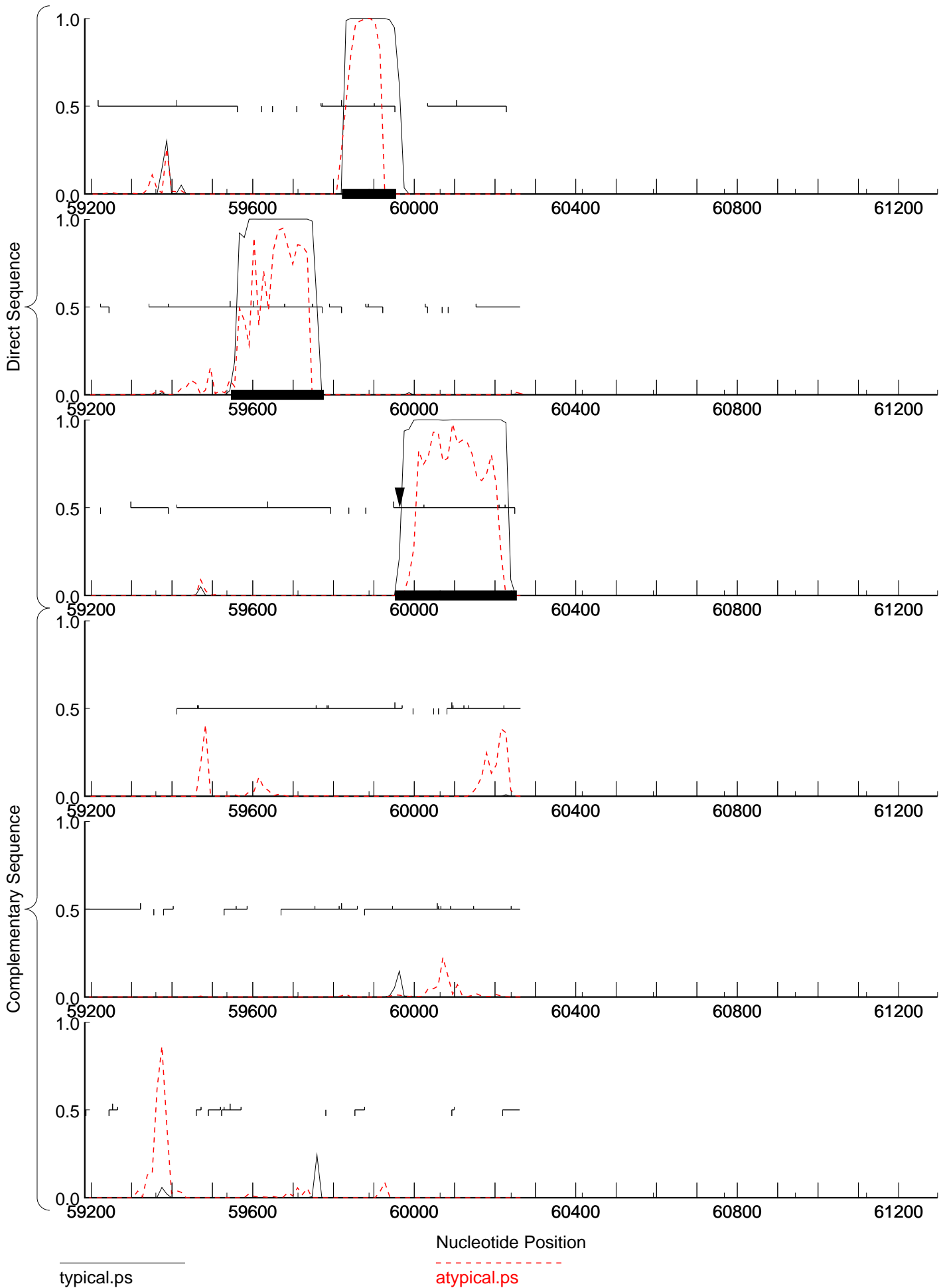












typical.ps

atypical.ps