

# GeneMark

Version 2.5p (09.08.06)

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## PROGRAM INFORMATION

Sequence : Mycobacterium phage Centaur complete sequence, 53001 bp including 10-base 3' overhang (CGGT  
Analysis Date : 10/23/17 at 15:02:04  
Pages : 27  
Sequence Length : 53001 bp  
GC Content : 63.50%

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,

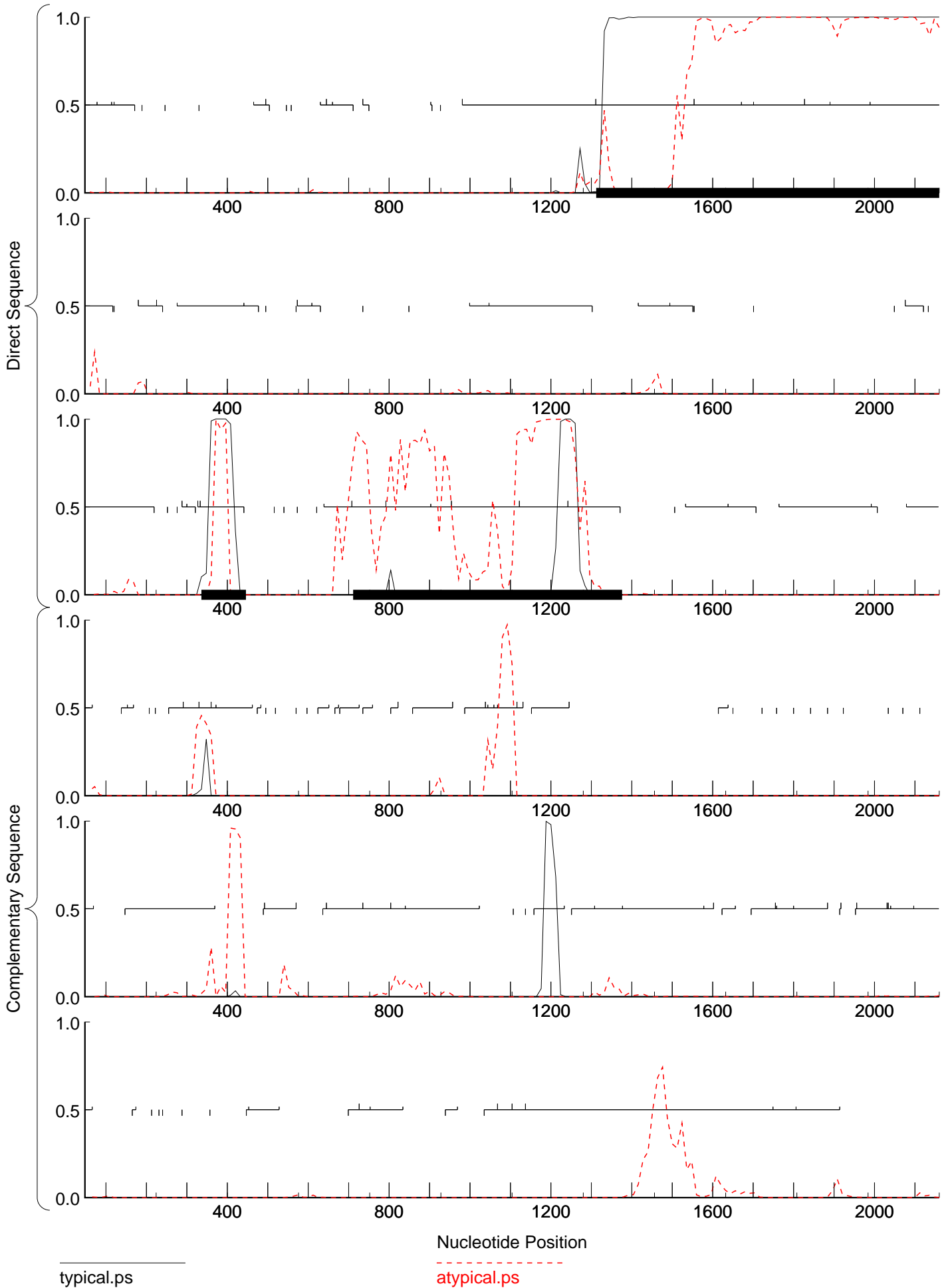
## Matrix notes & comments

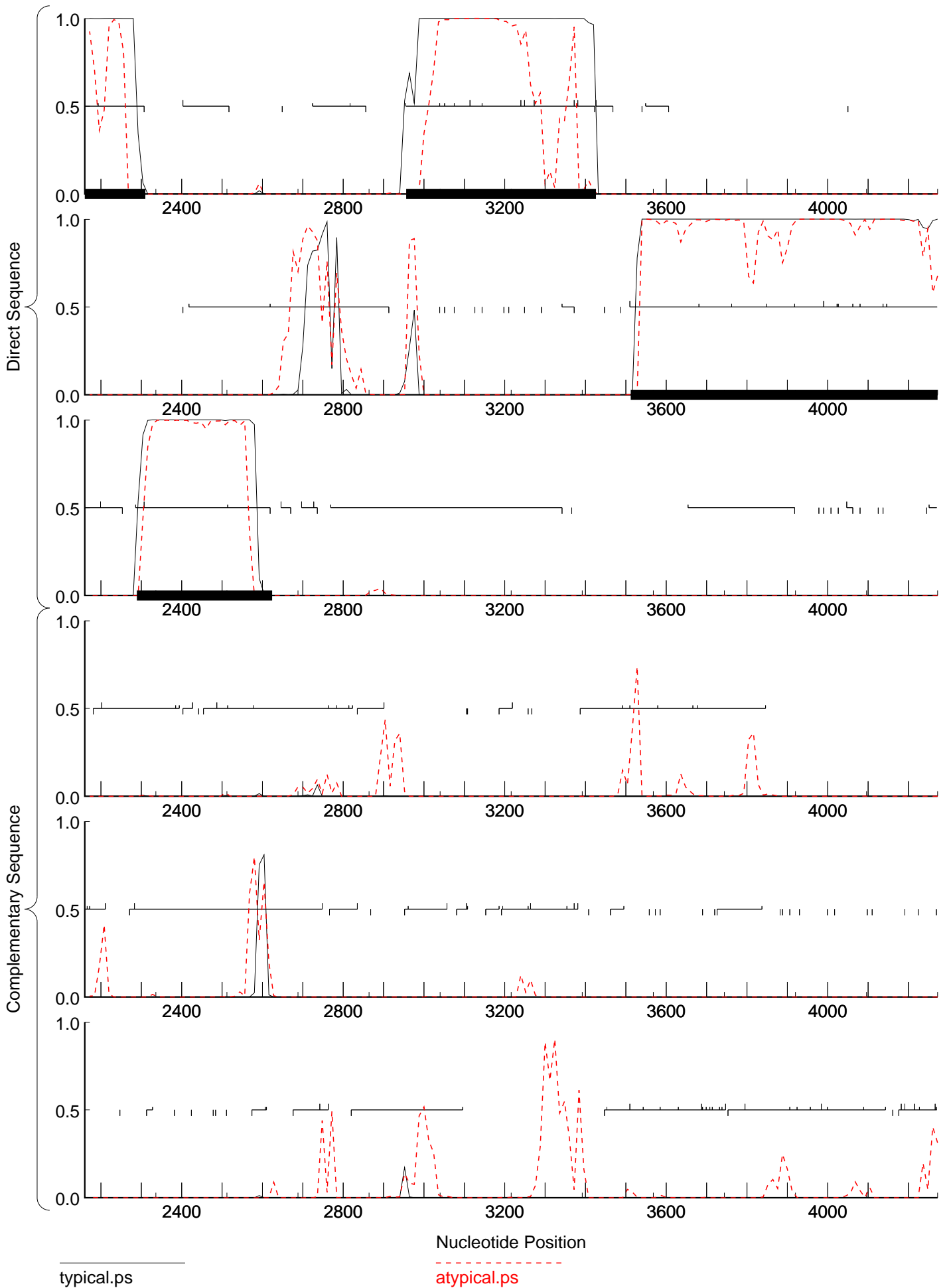
Training set derived by GeneMarkS, 4.30 October 2014  
Mon Oct 23 15:02:04 2017

## MATRIX INFORMATION

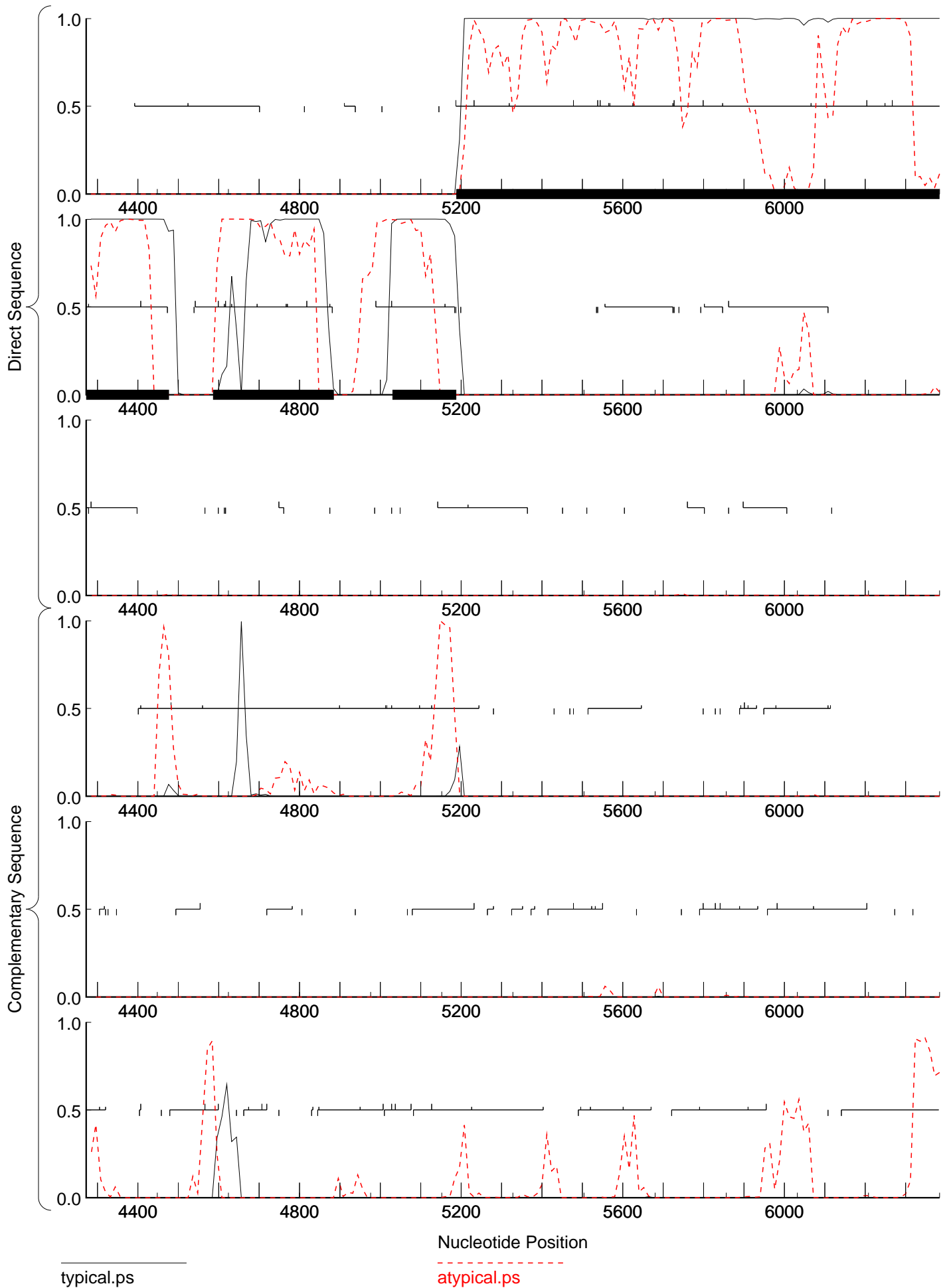
Matrix : -  
Author : -  
Order : 4

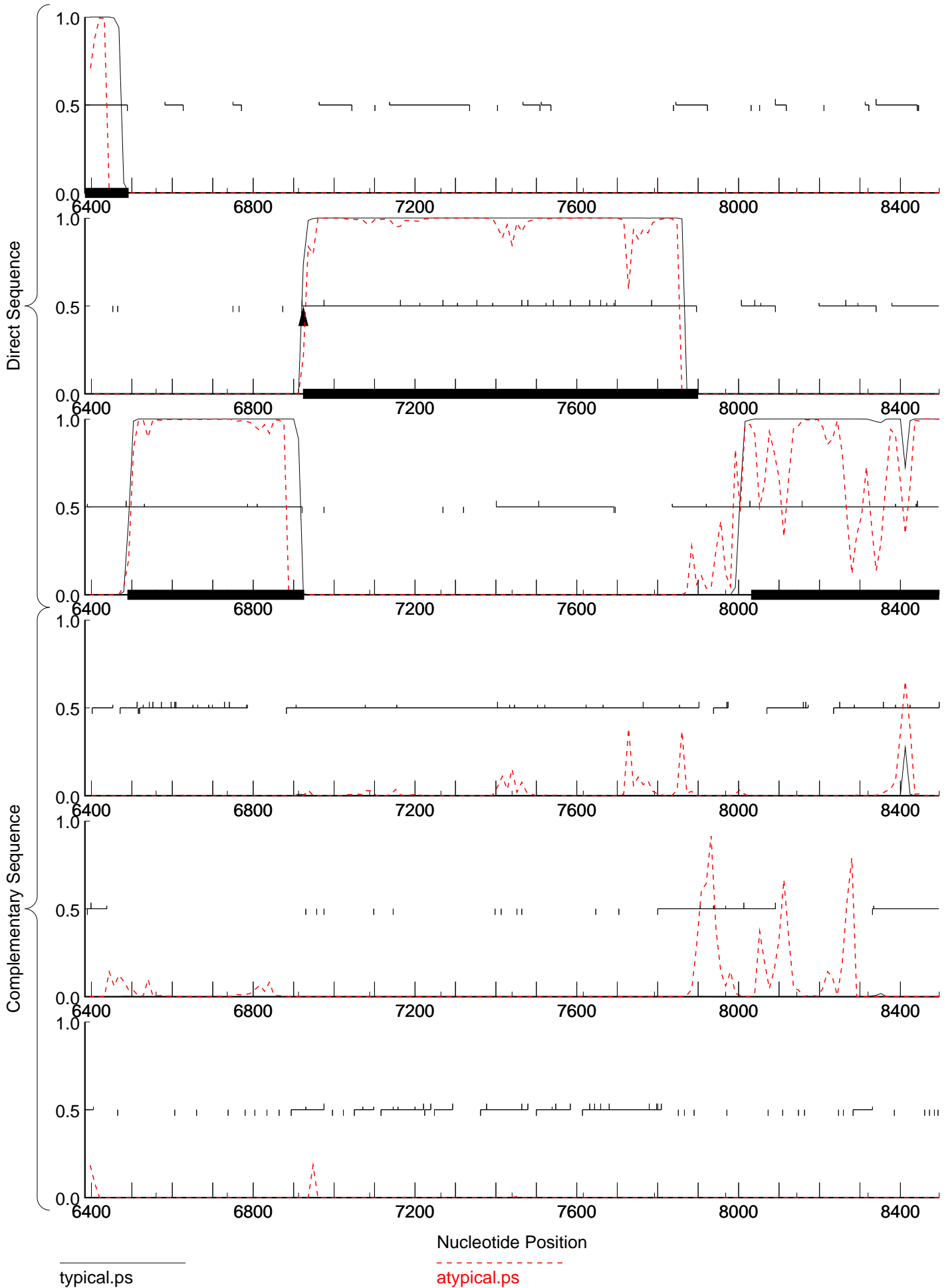
Send questions / comments to:  
Dr. M. Borodovsky  
Georgia Institute of Technology  
School of Biology  
Atlanta, GA 30332-0230

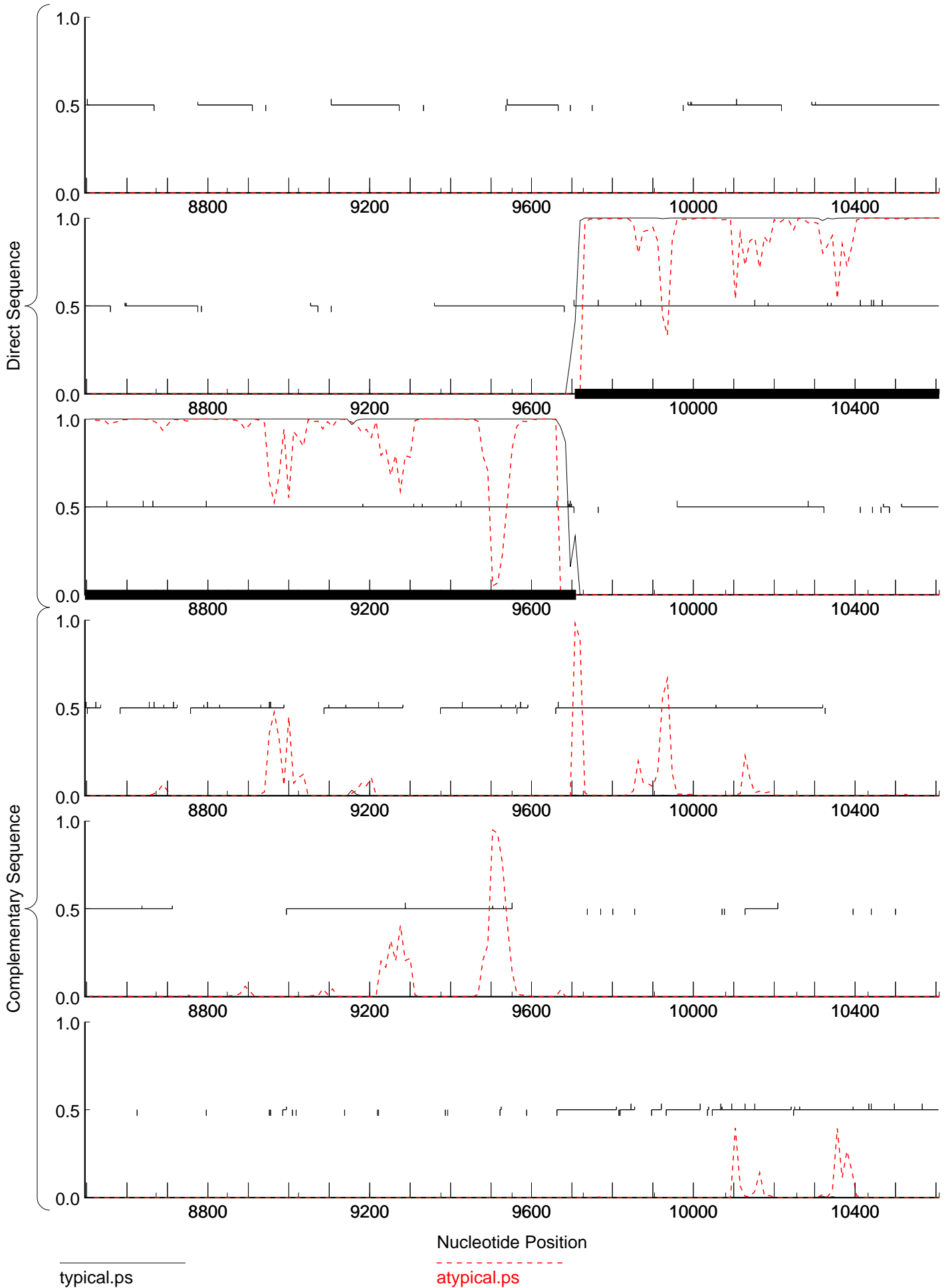


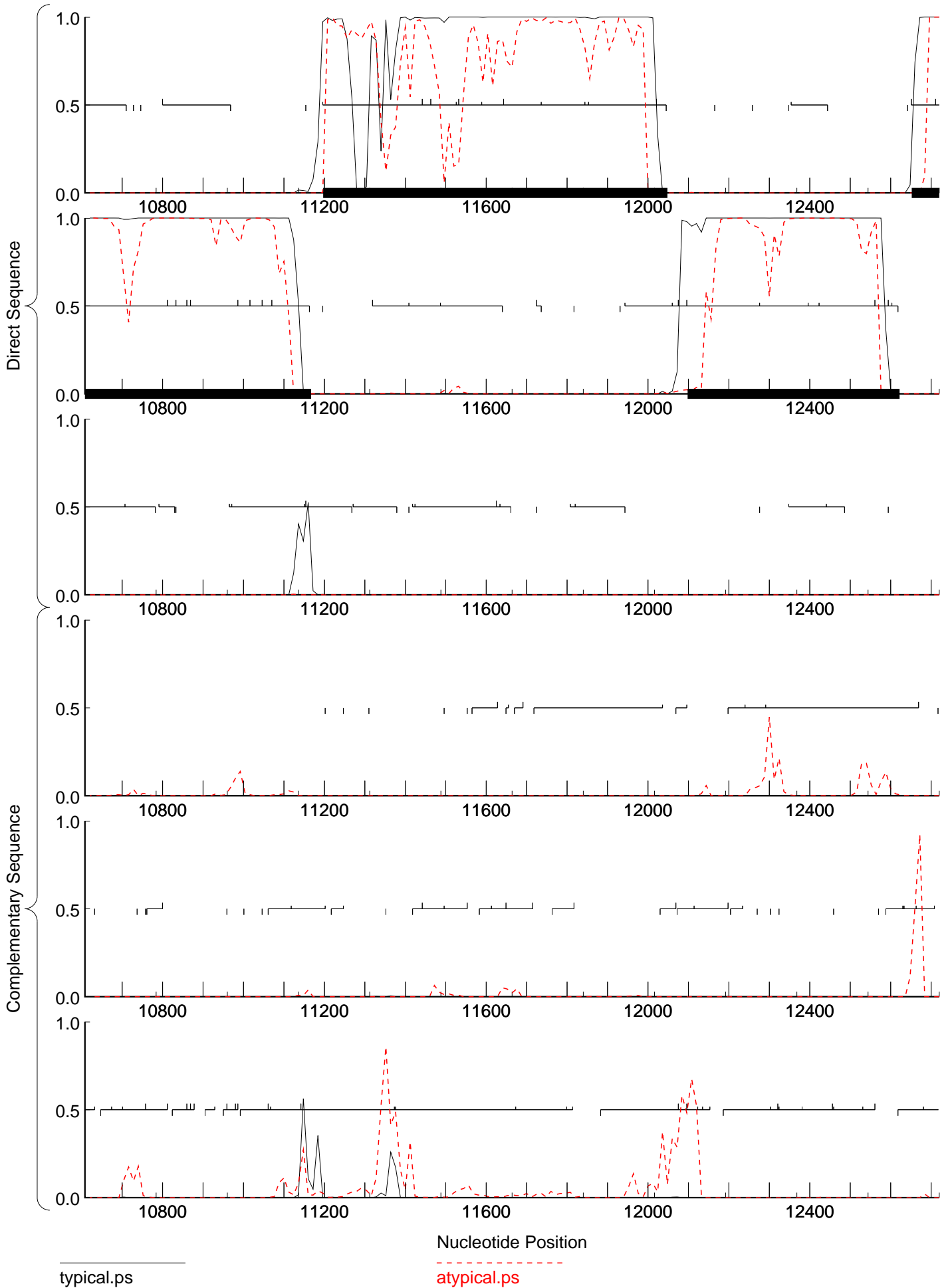


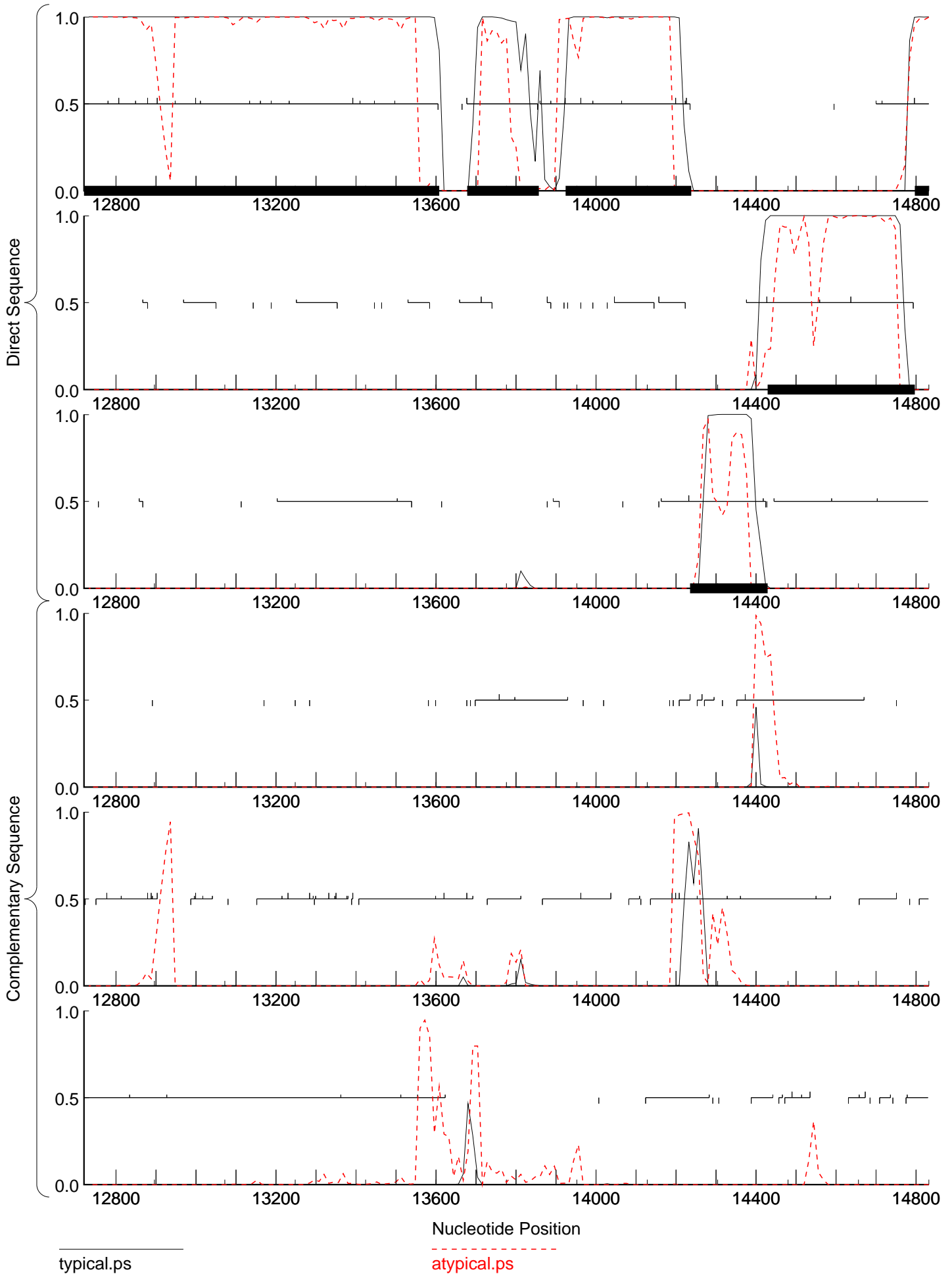
GeneMark.hmm prediction







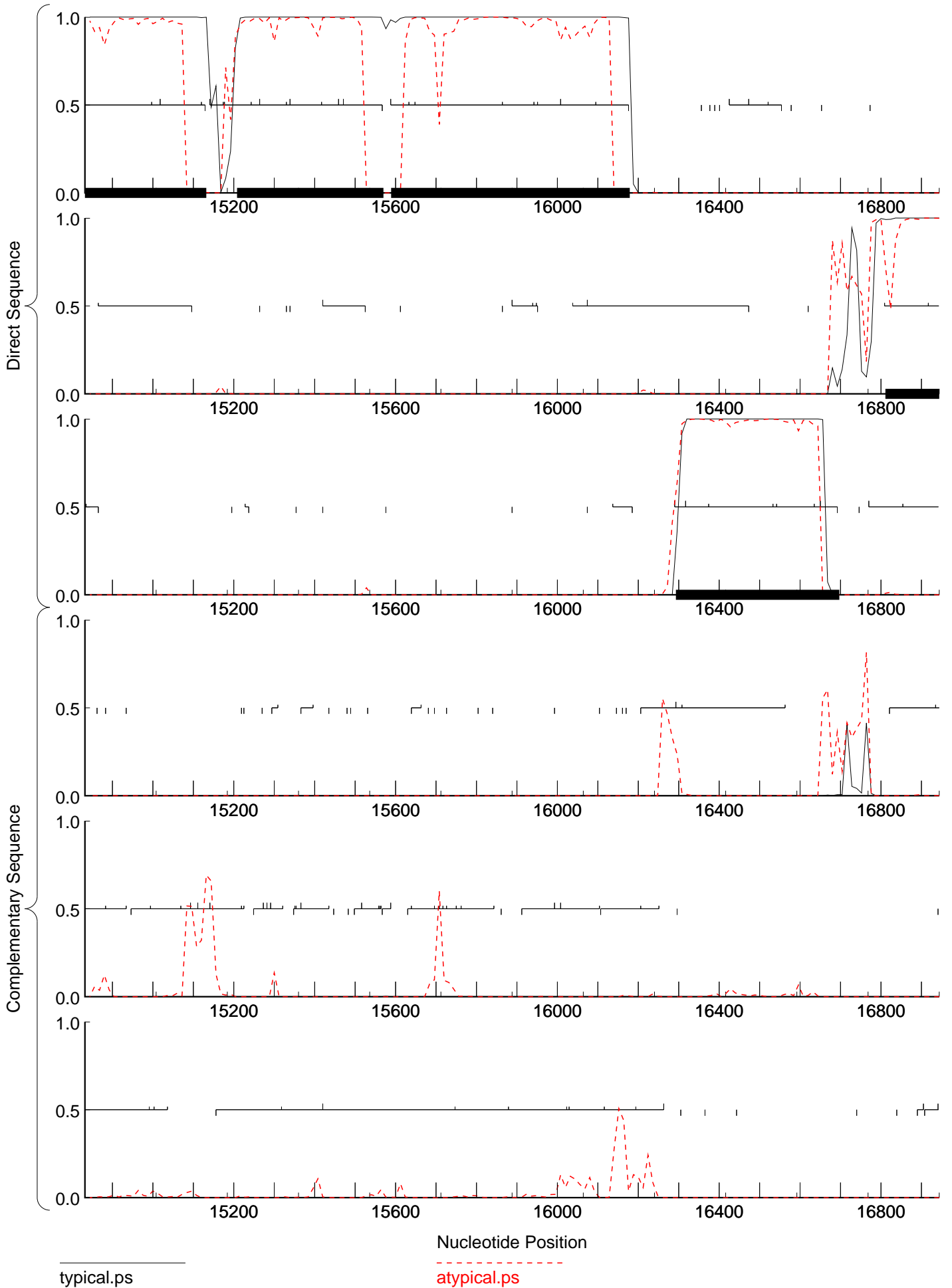


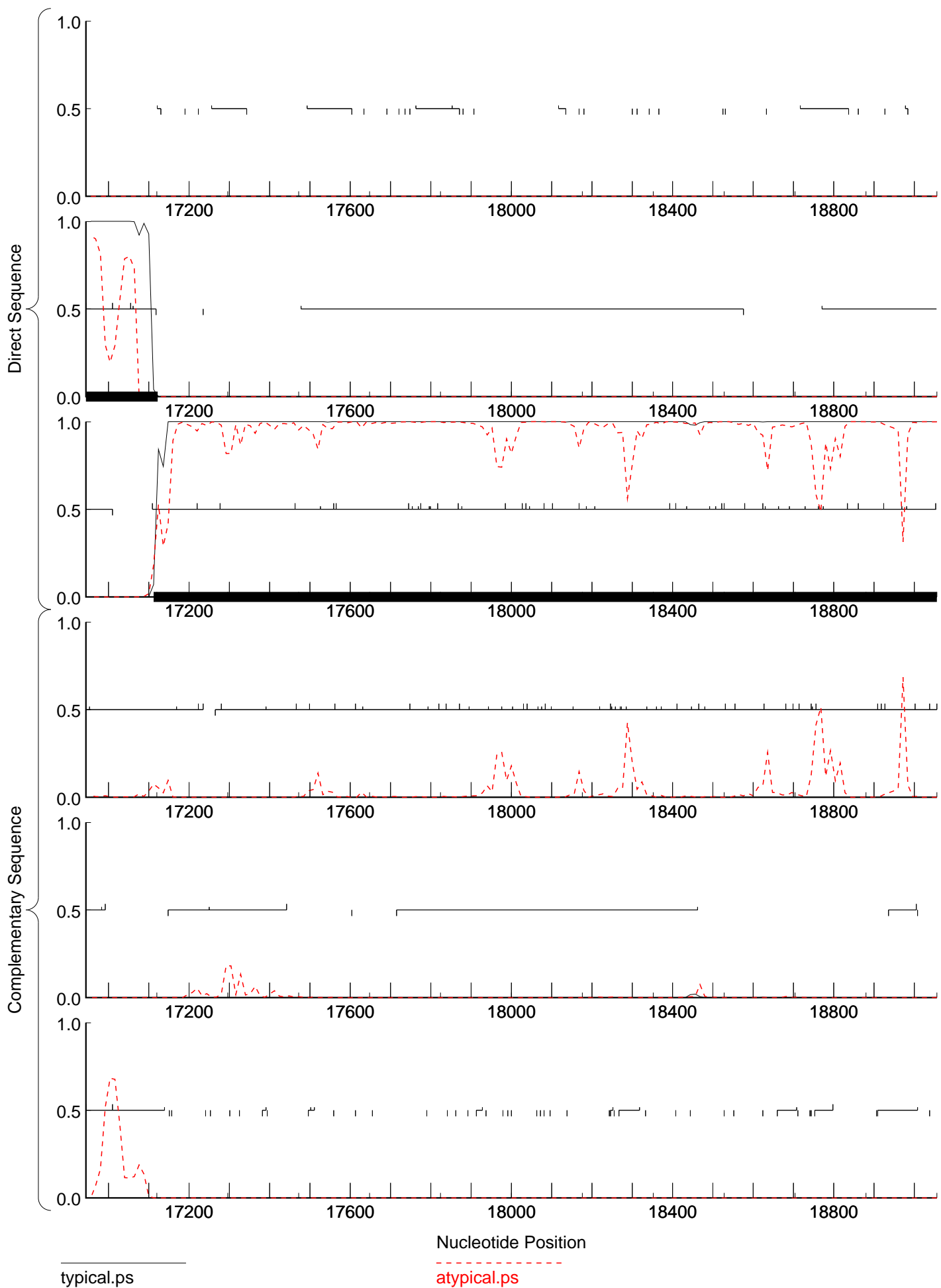


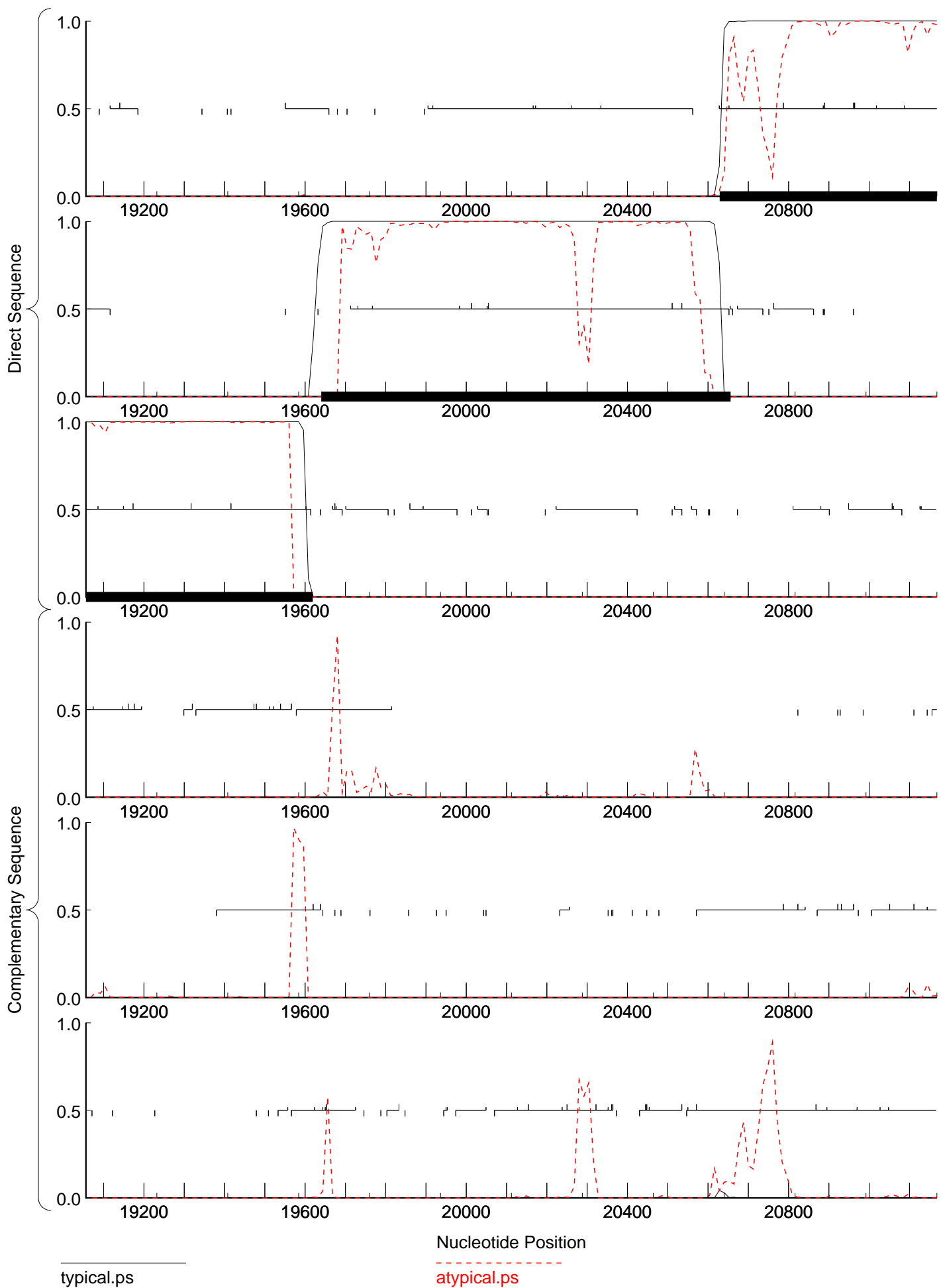
typical.ps

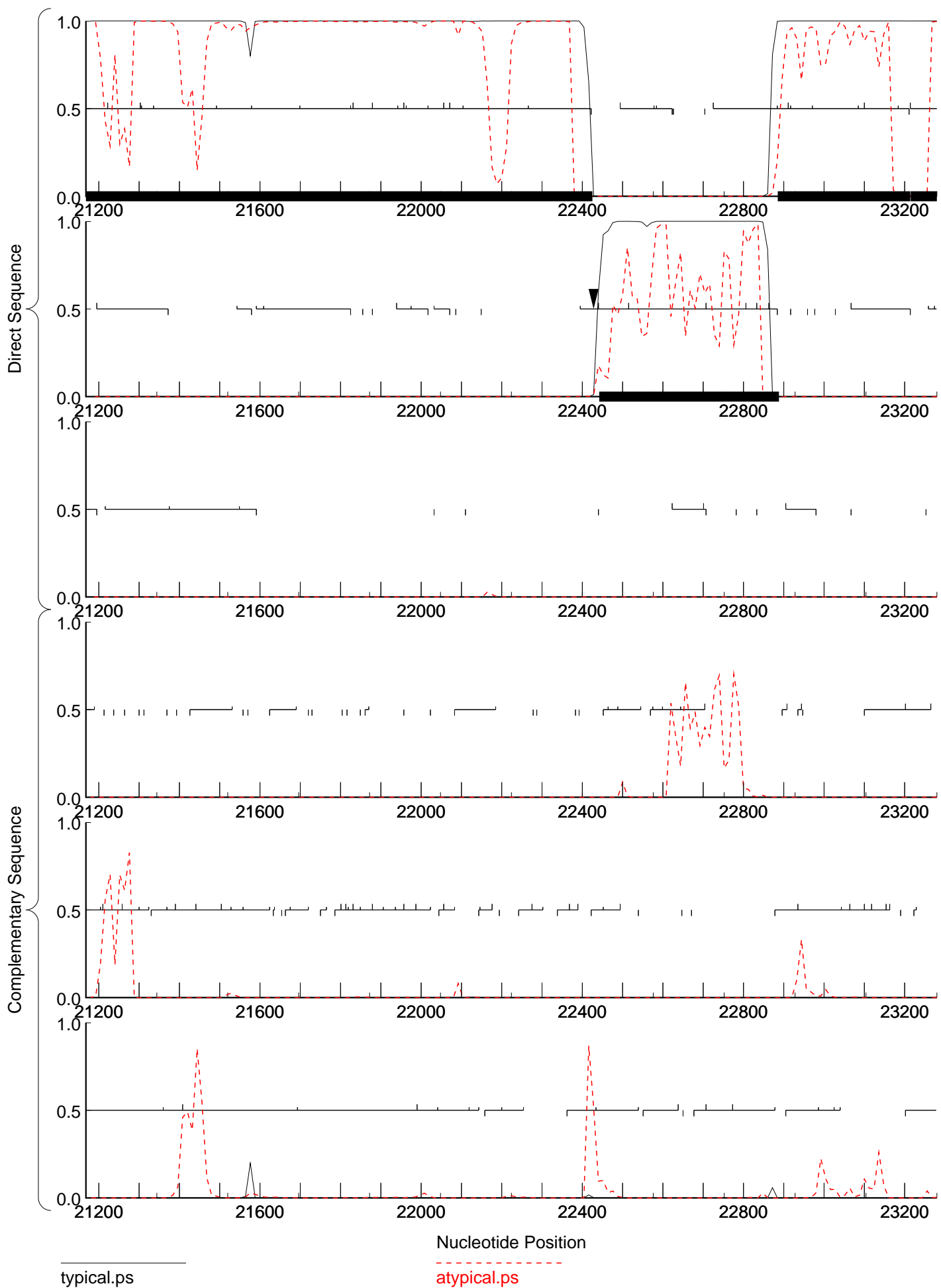
atypical.ps

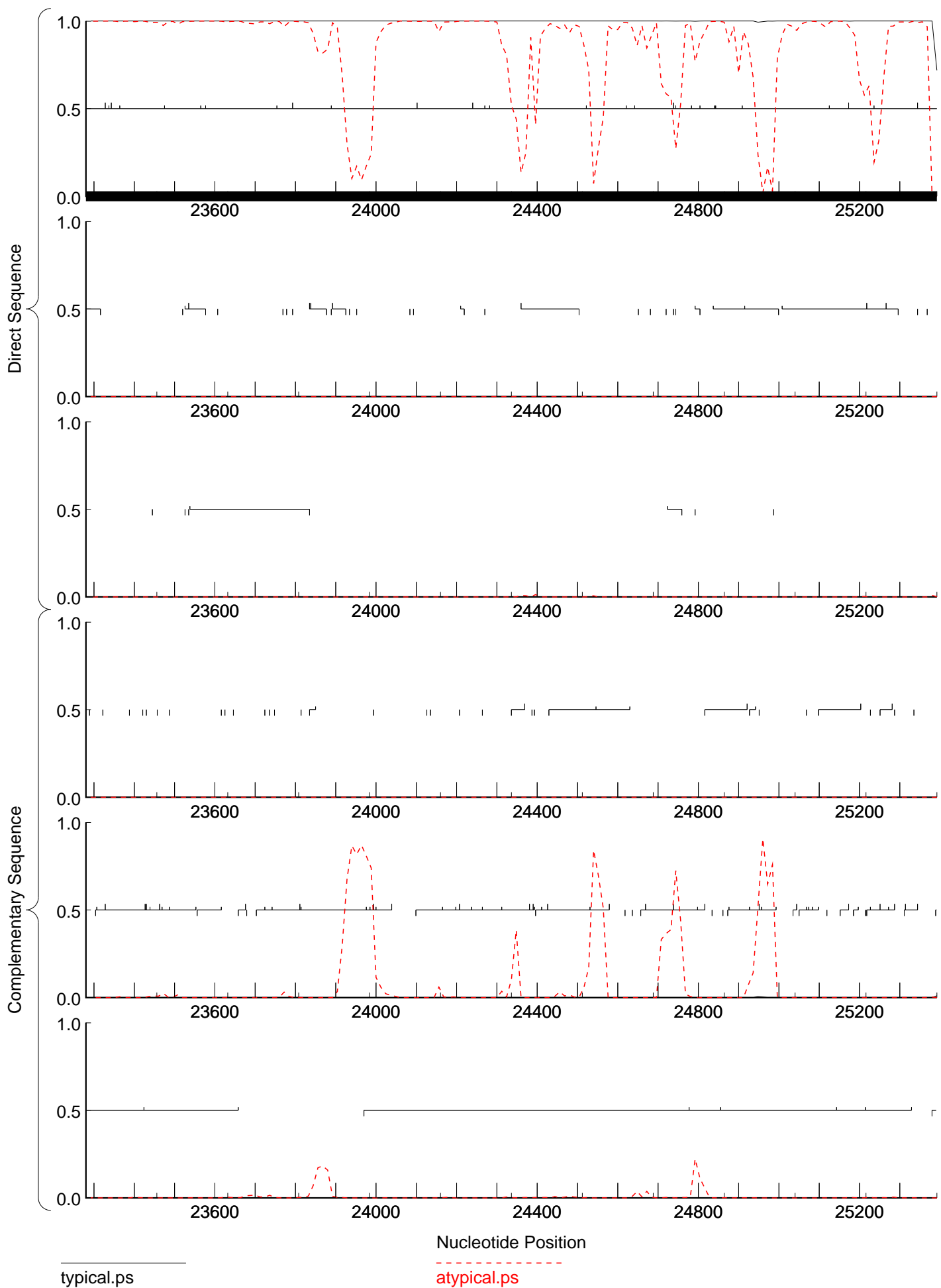






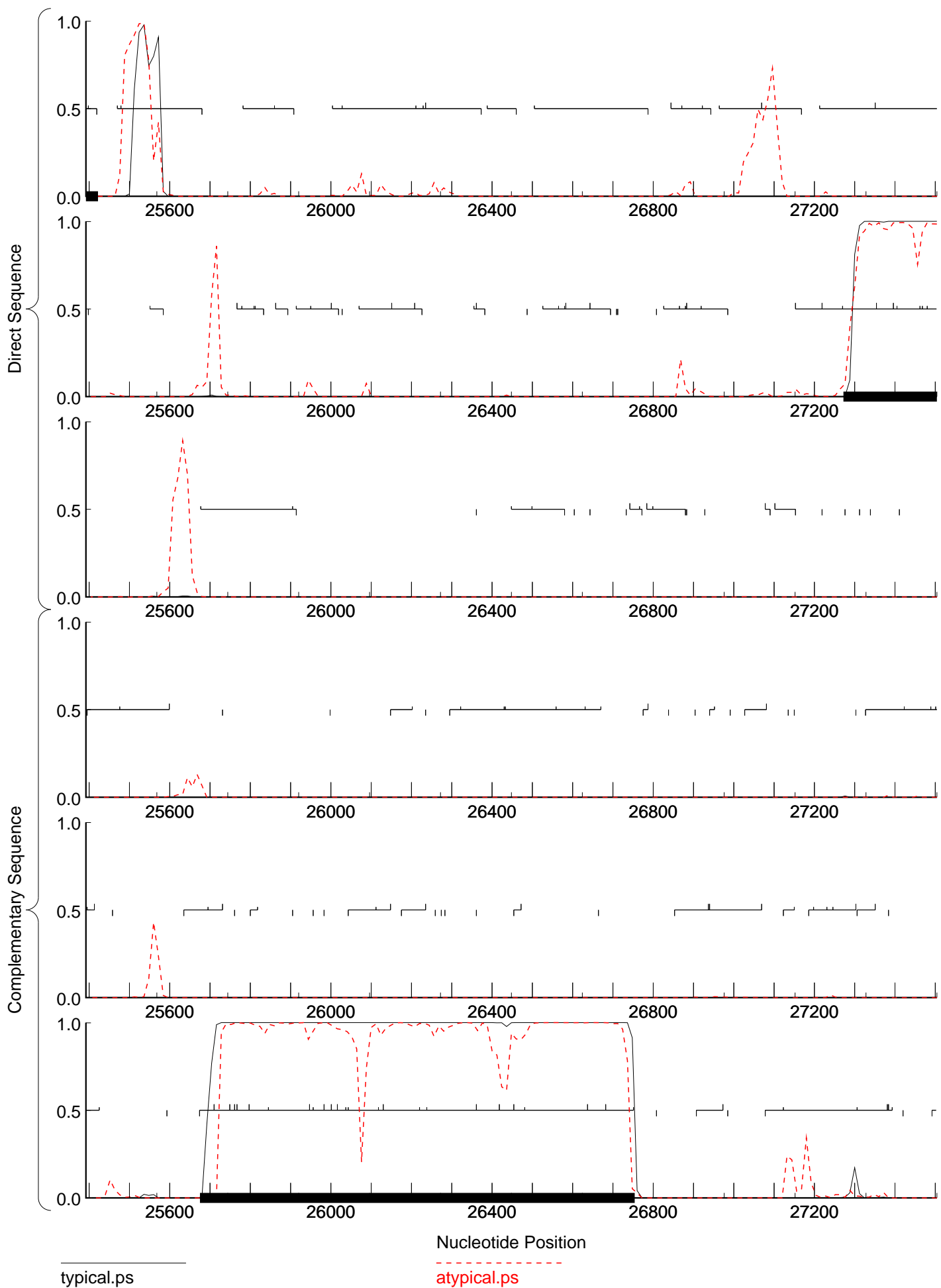


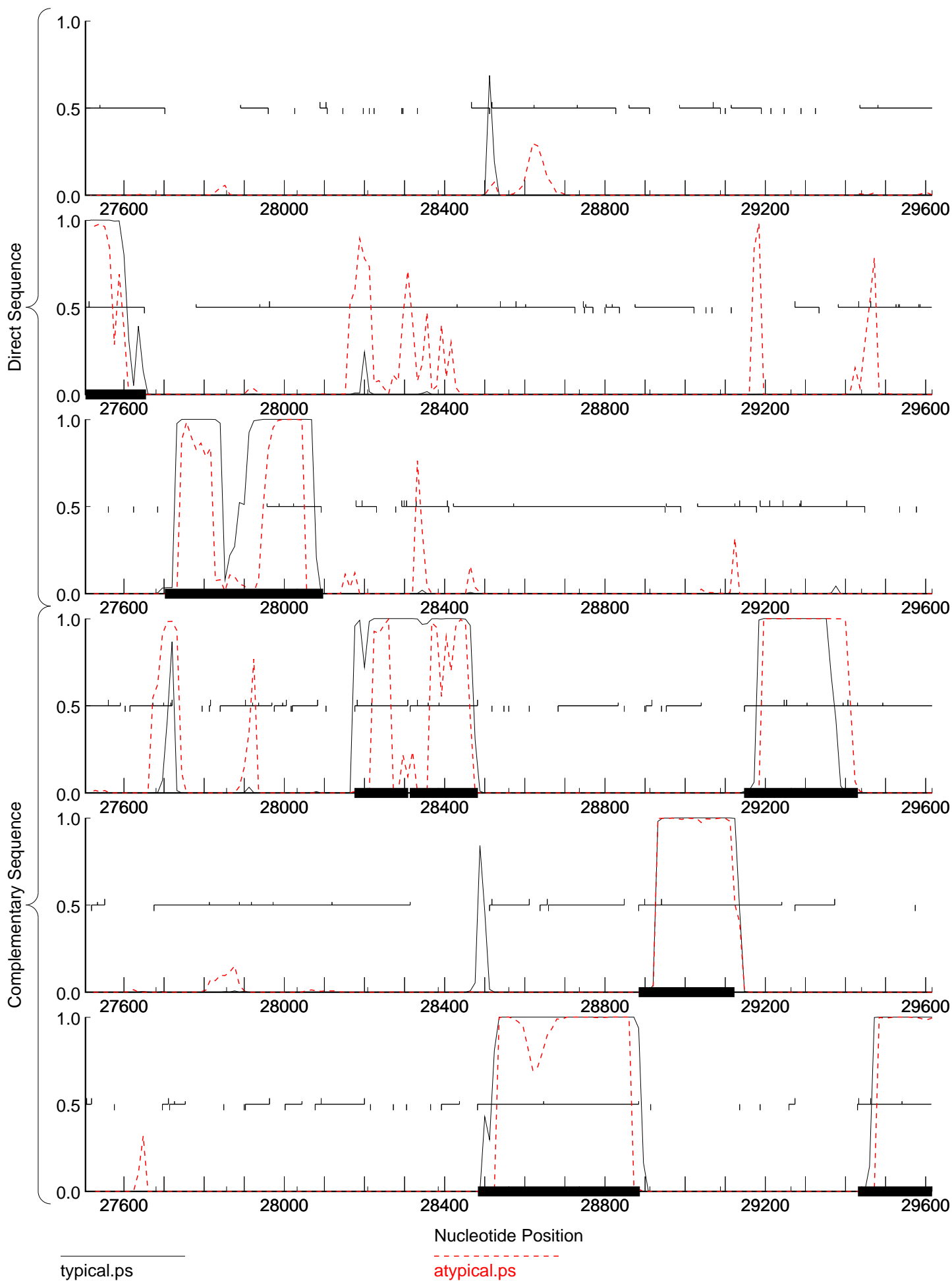


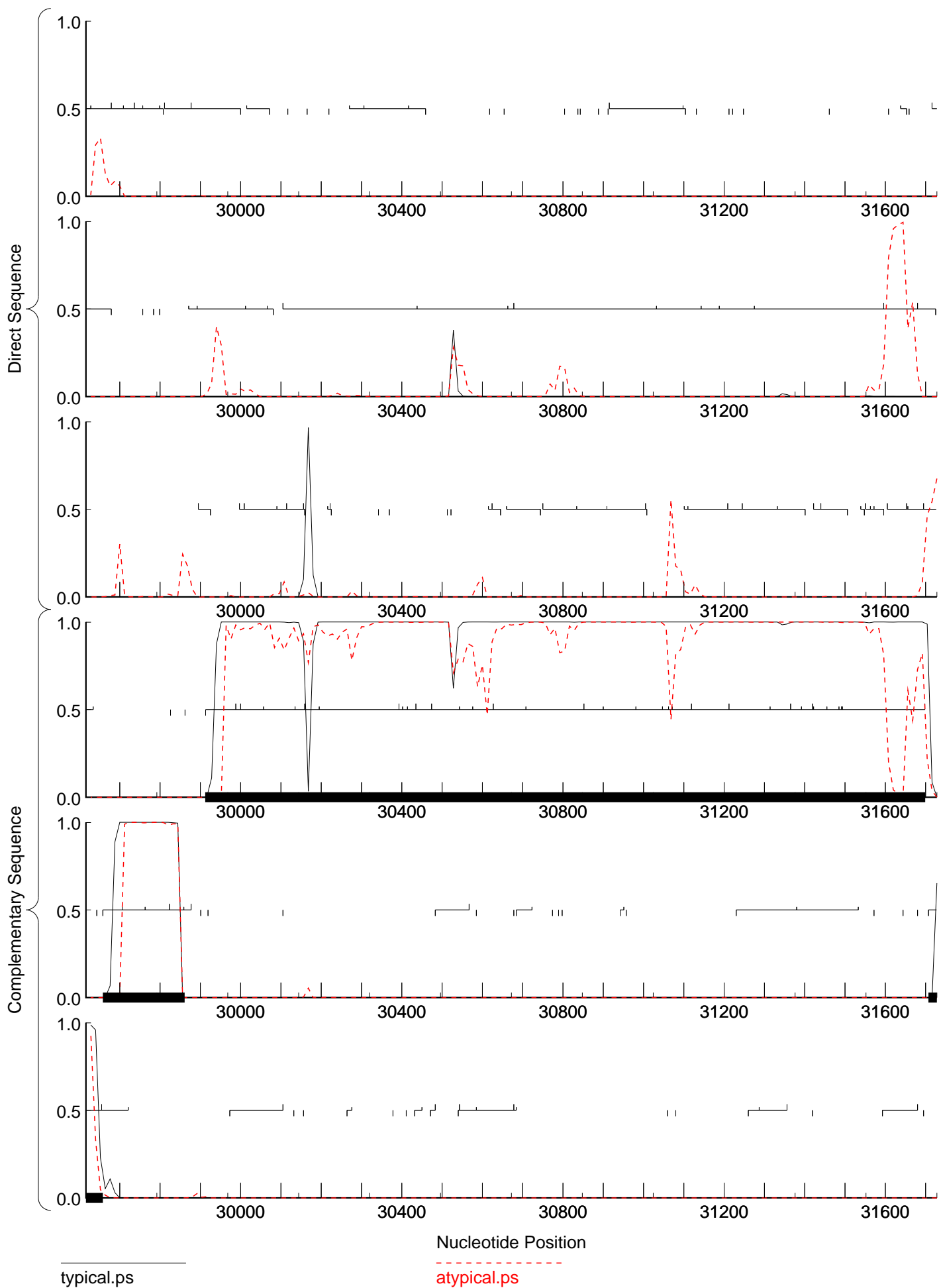


typical.ps

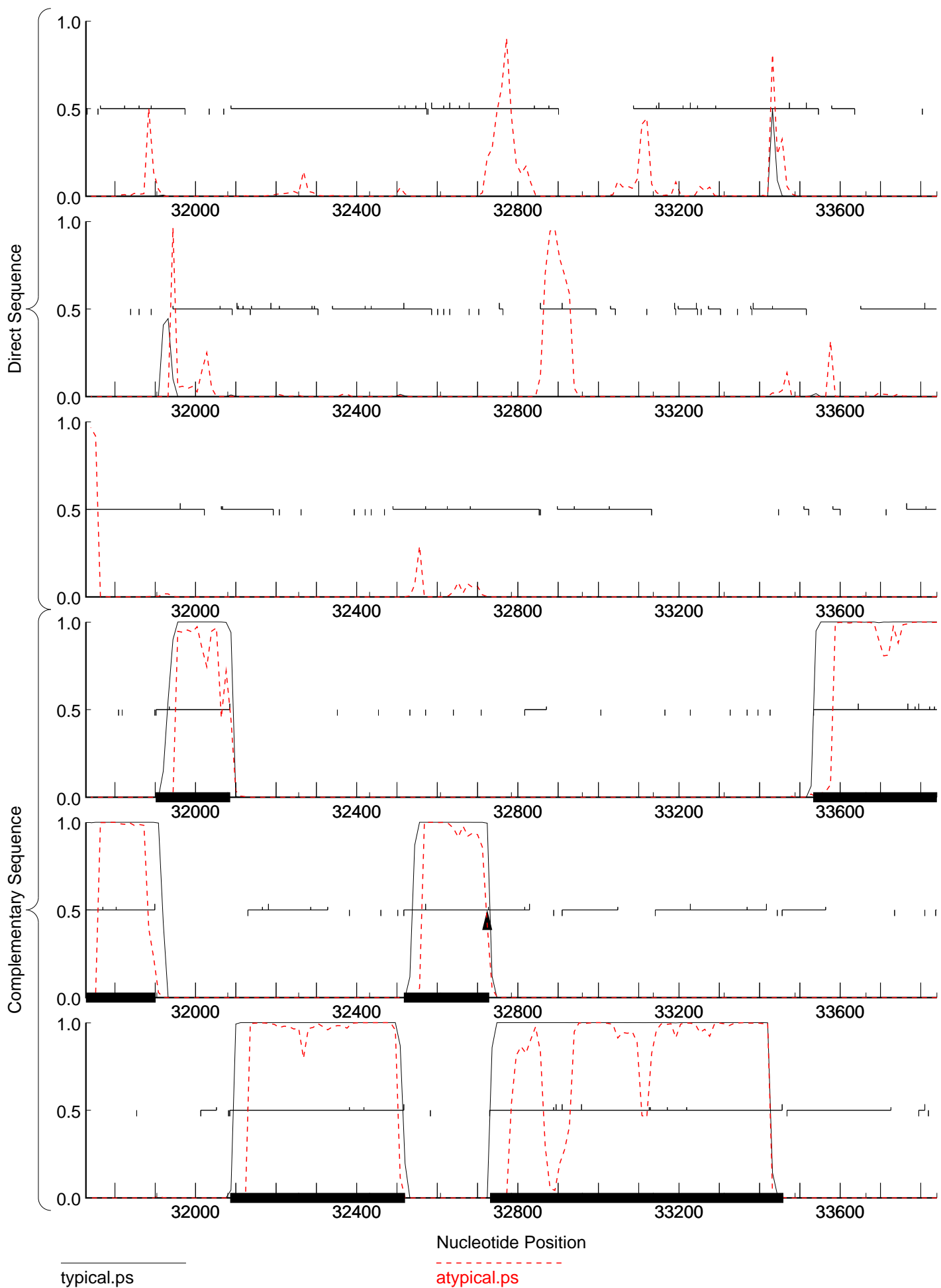
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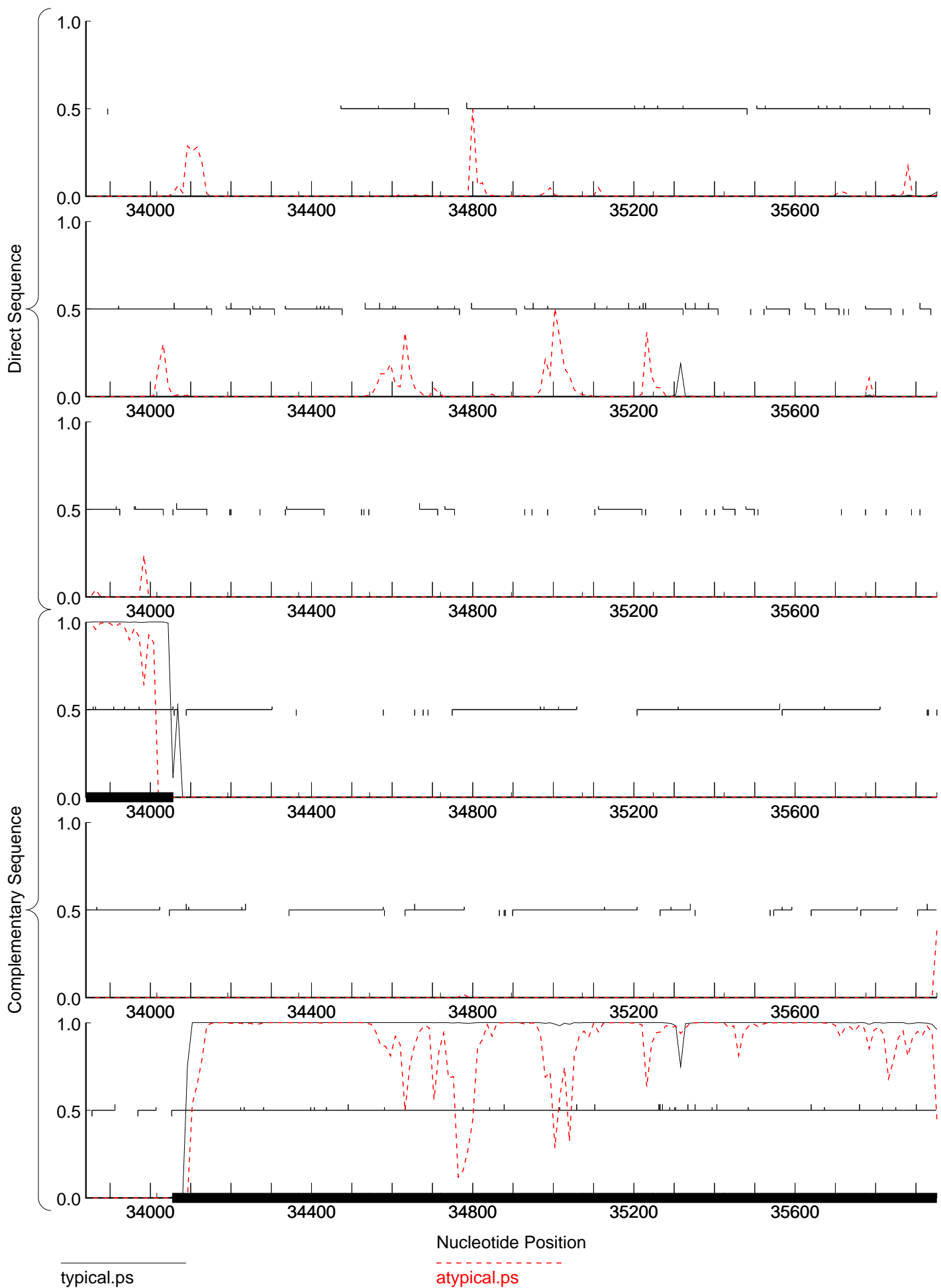












typical.ps

atypical.ps

