

GeneMark

Version 2.5p (09.08.06)

Copyright 1993 - M. Borodovsky, J. McIninch

PROGRAM INFORMATION

Sequence : Mycobacterium phage Manatee Complete Sequence, 51040 bp including 10-base 3' overhang (CGGA
Analysis Date : 9/22/17 at 3:53:22
Pages : 26
Sequence Length : 51040 bp
GC Content : 63.79%

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,

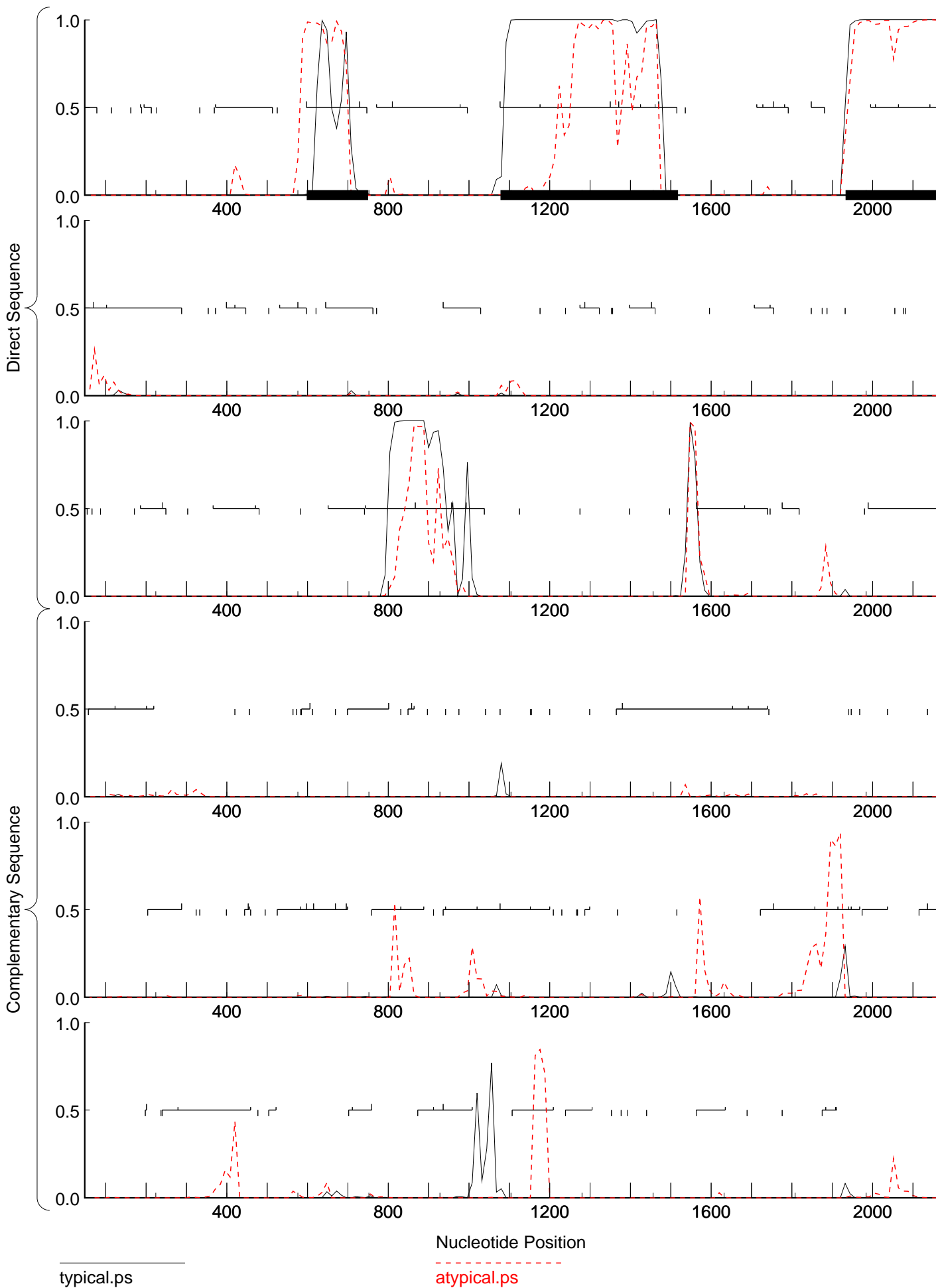
MATRIX INFORMATION

Matrix : -
Author : -
Order : 4

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

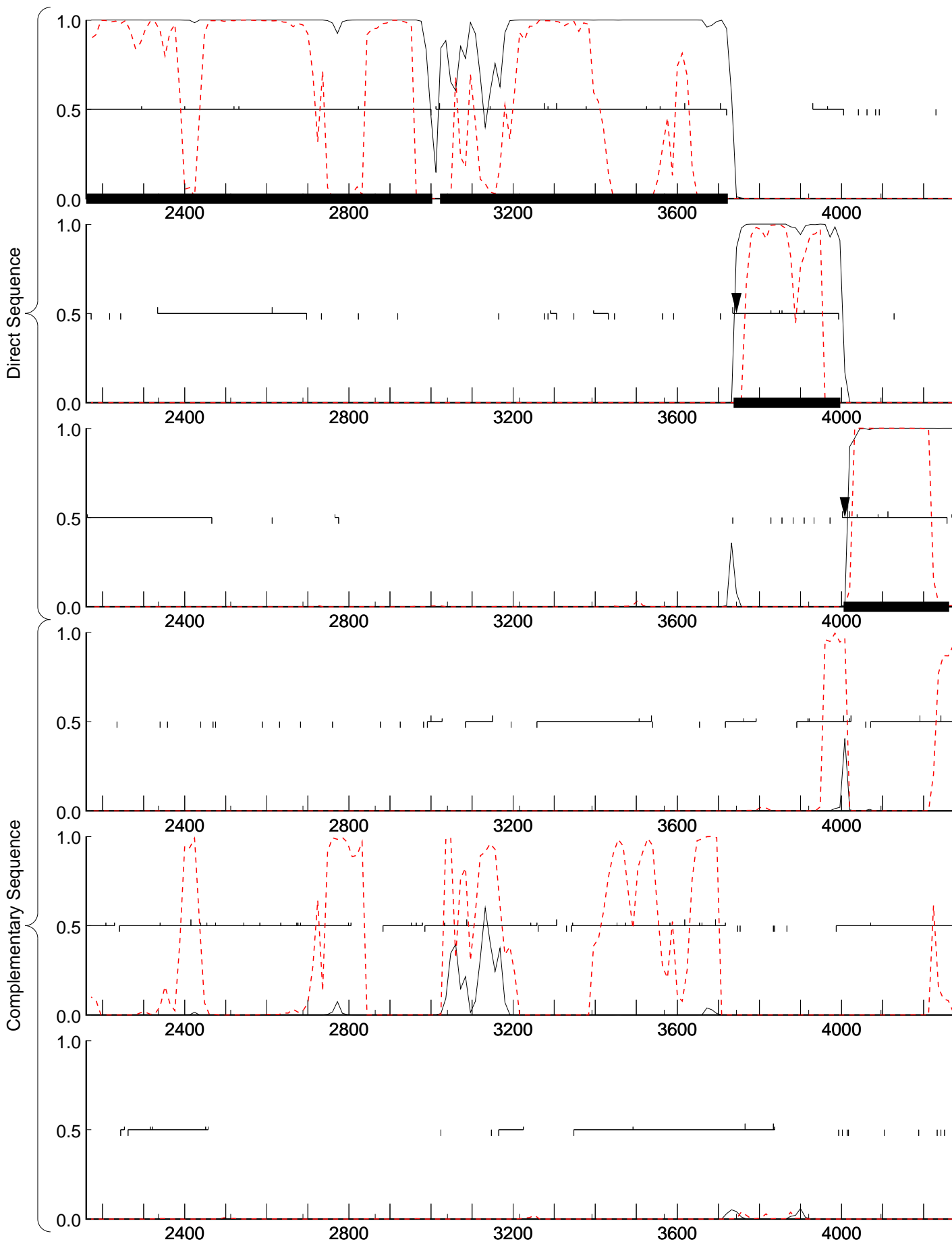
Matrix notes & comments

Training set derived by GeneMarkS, 4.30 October 2014
Fri Sep 22 03:53:22 2017



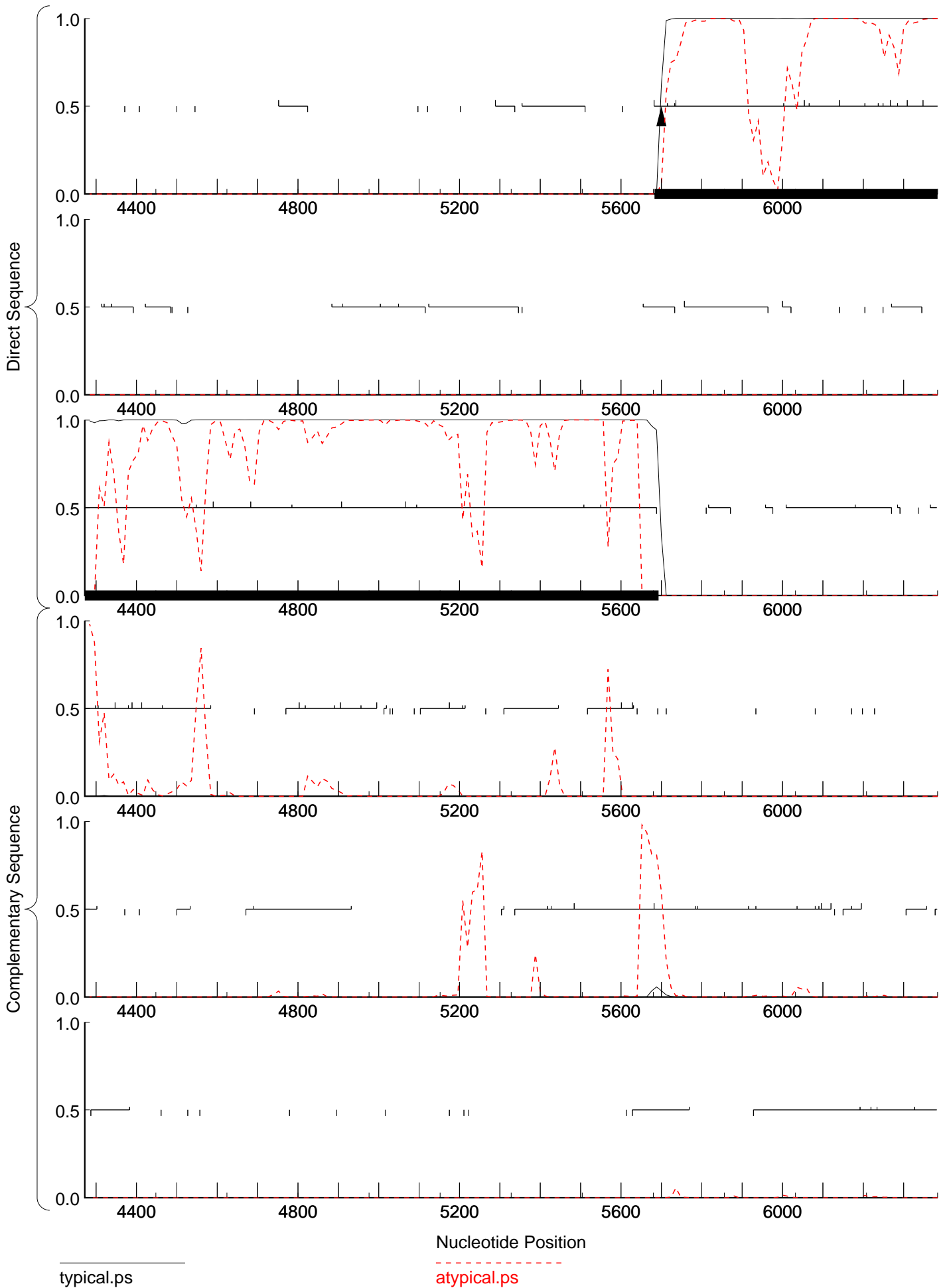
typical.ps

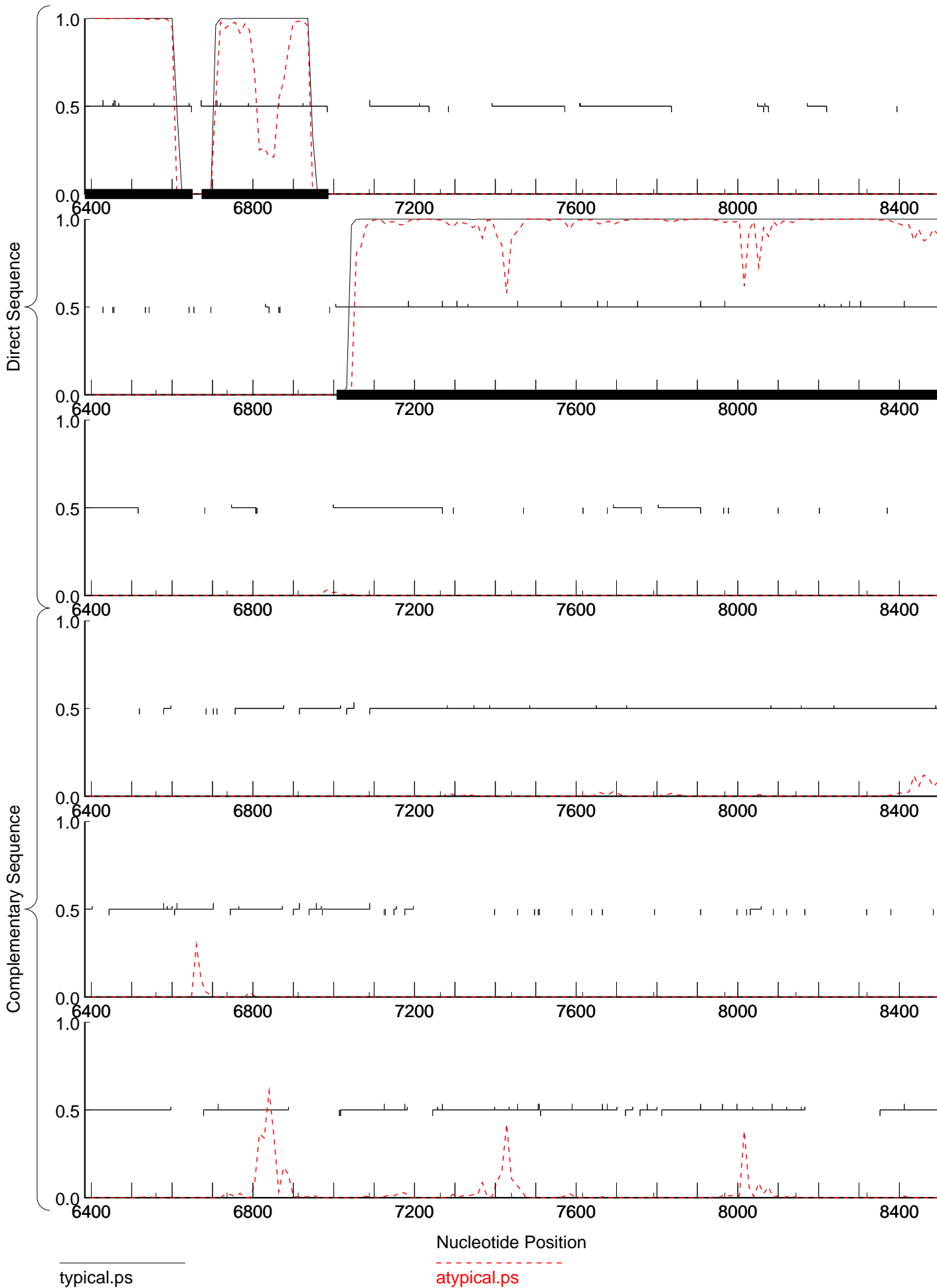
atypical.ps

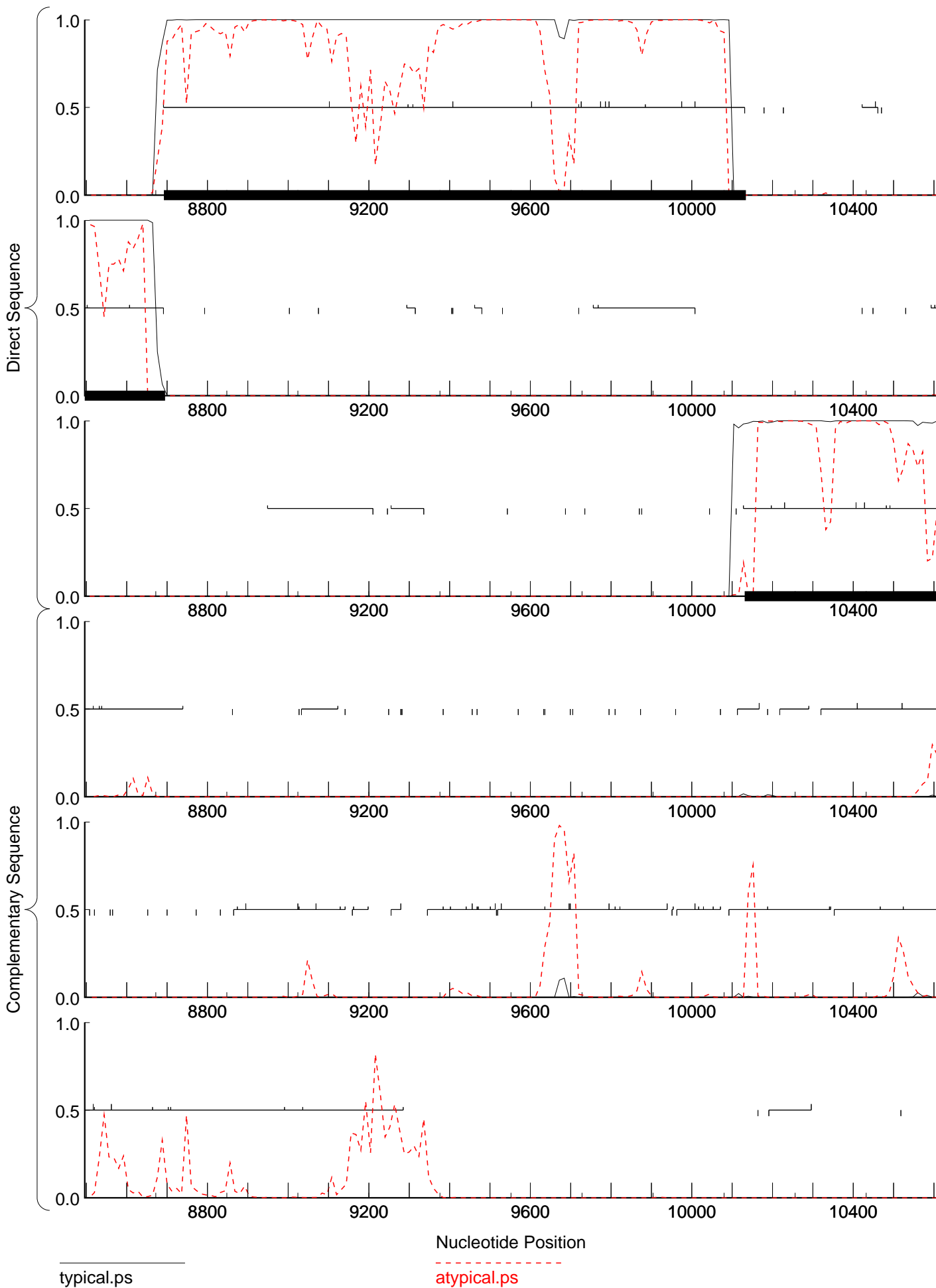


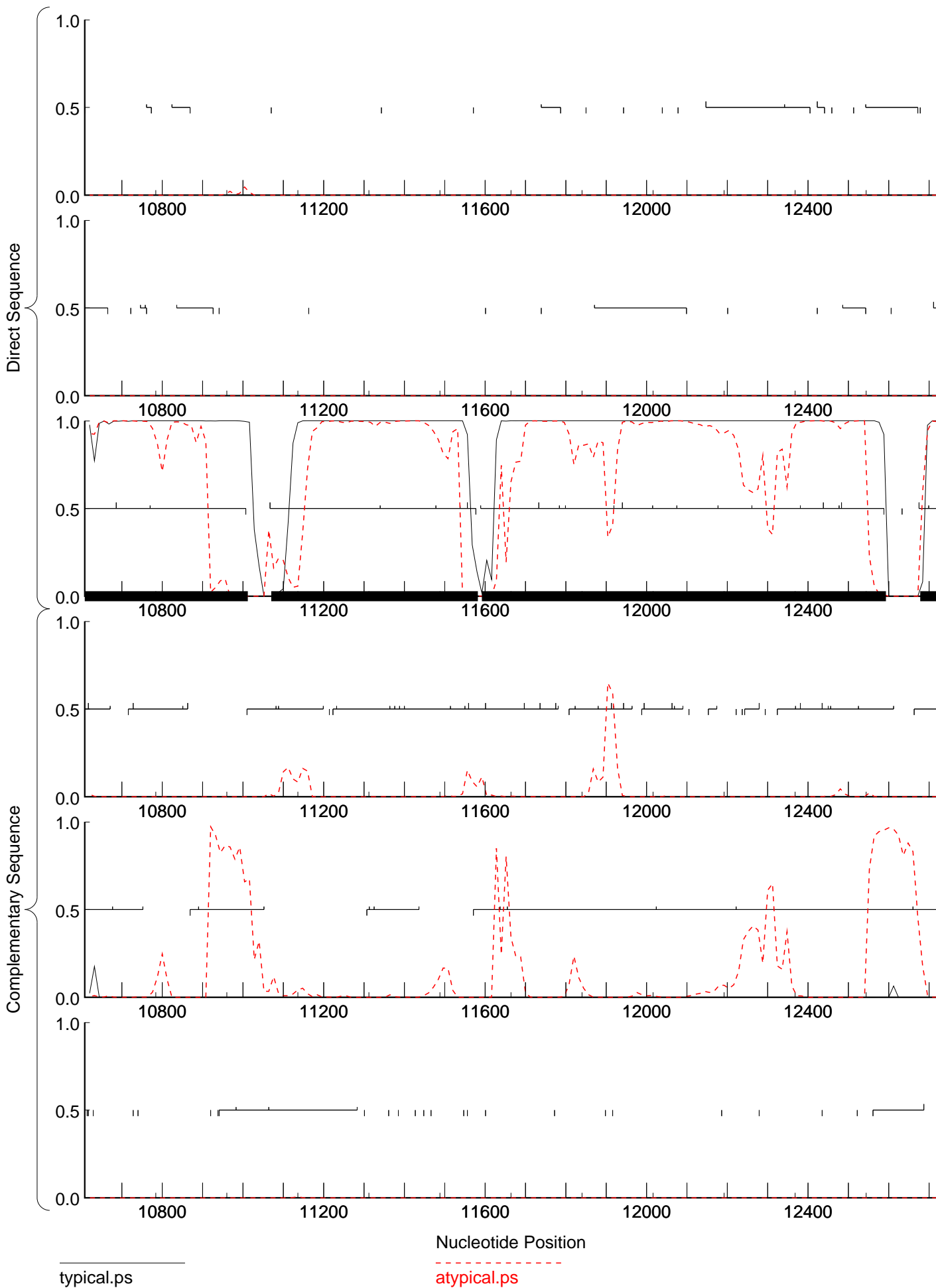
typical.ps

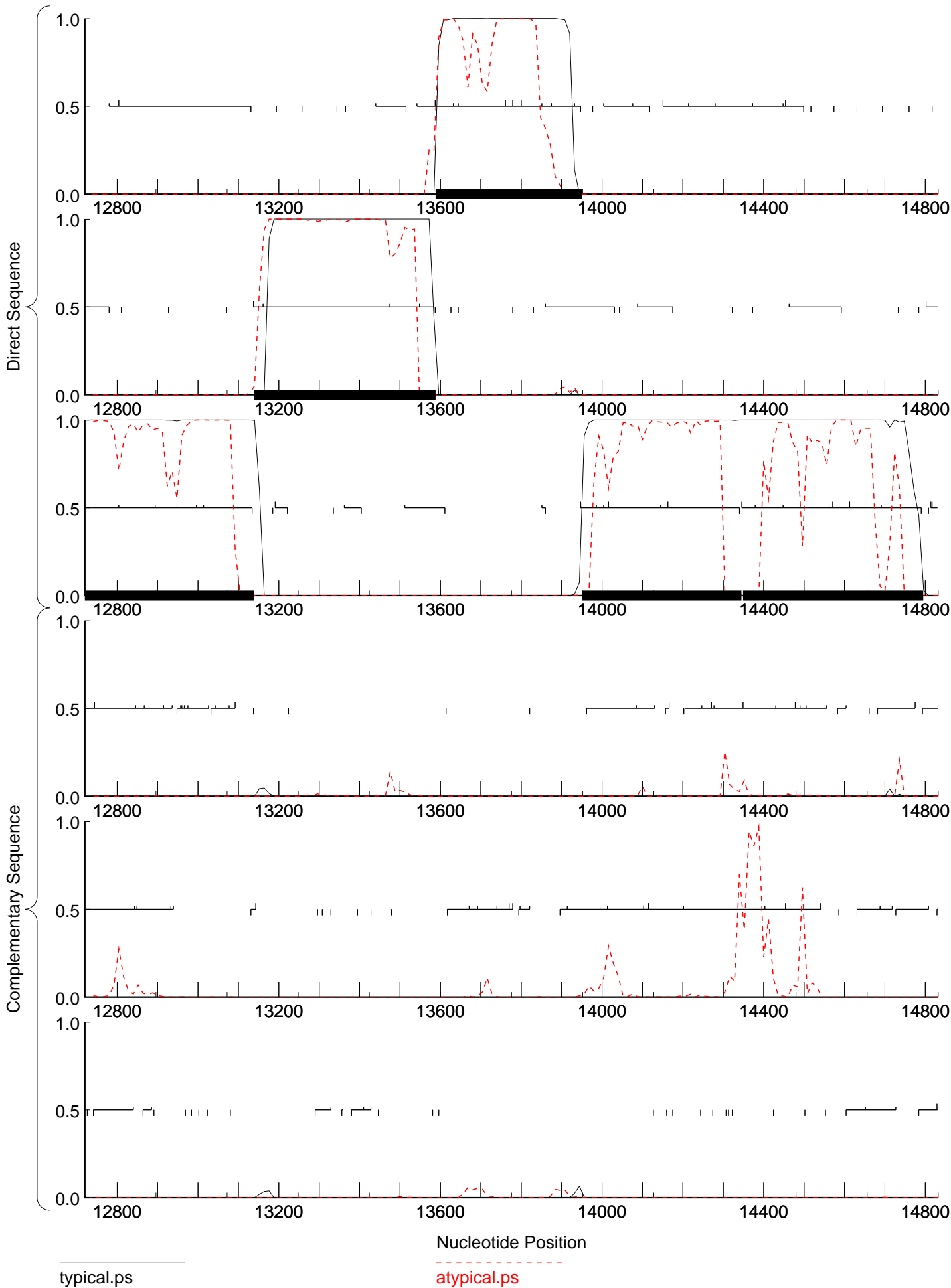
atypical.ps

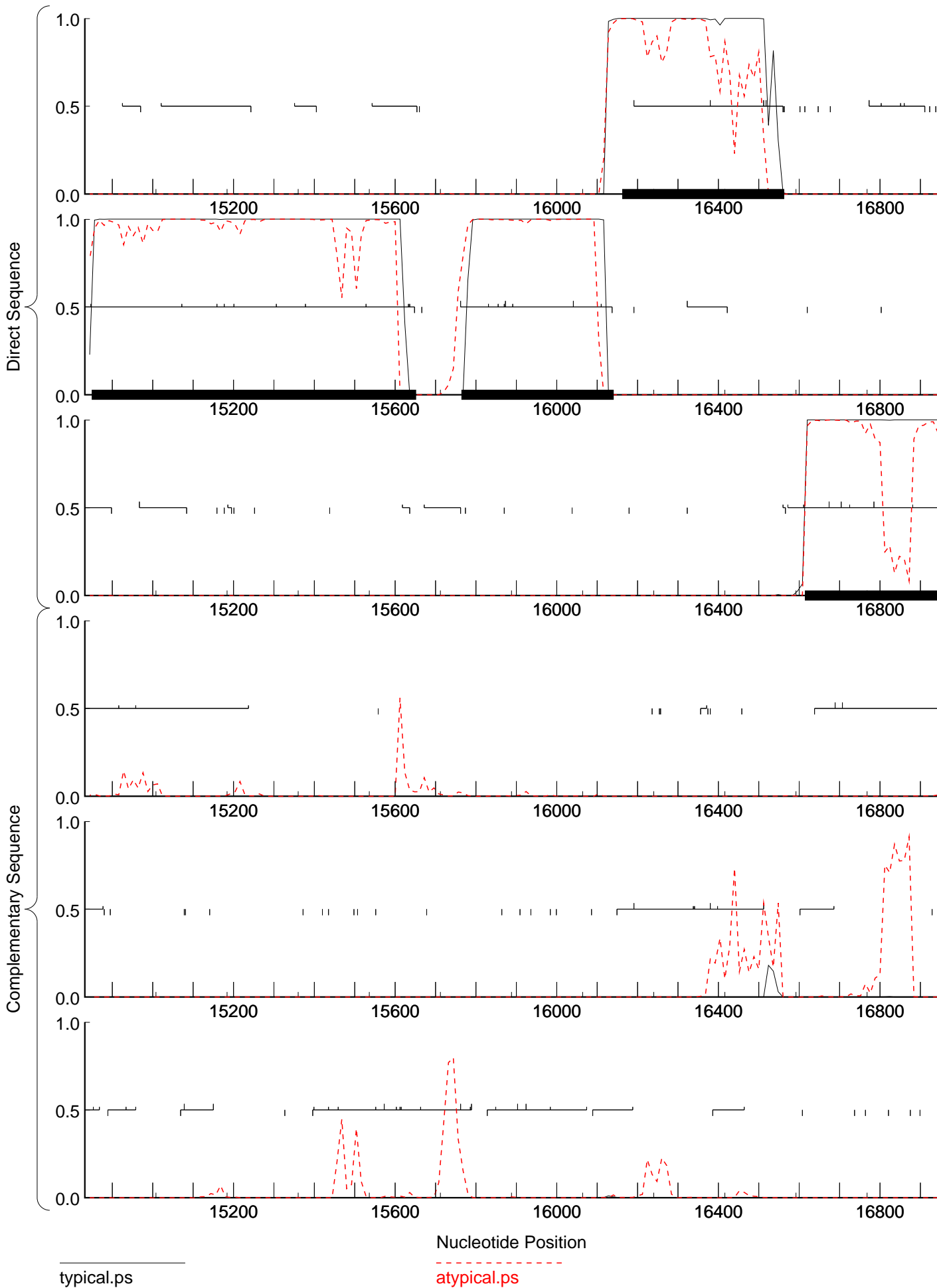






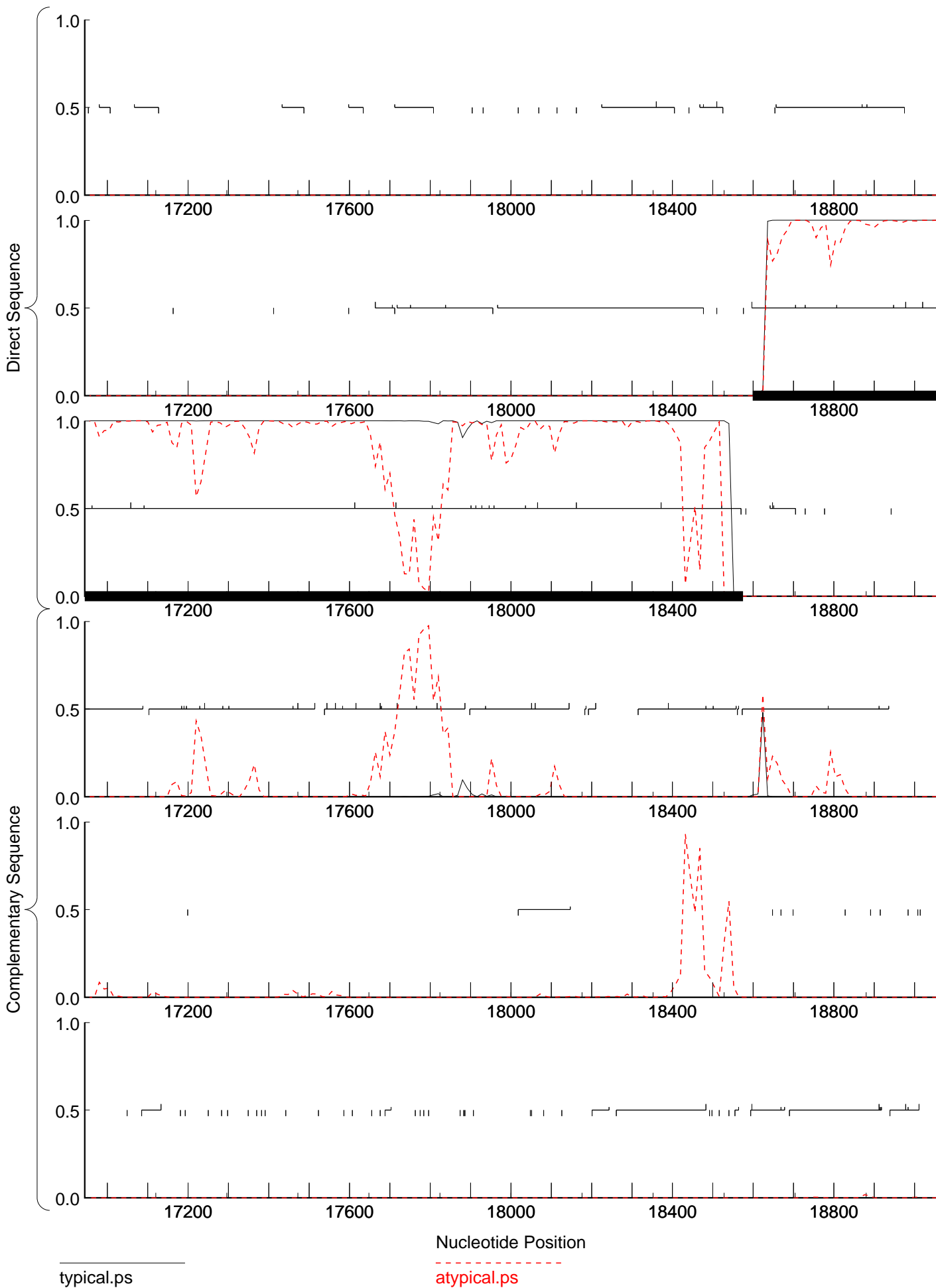


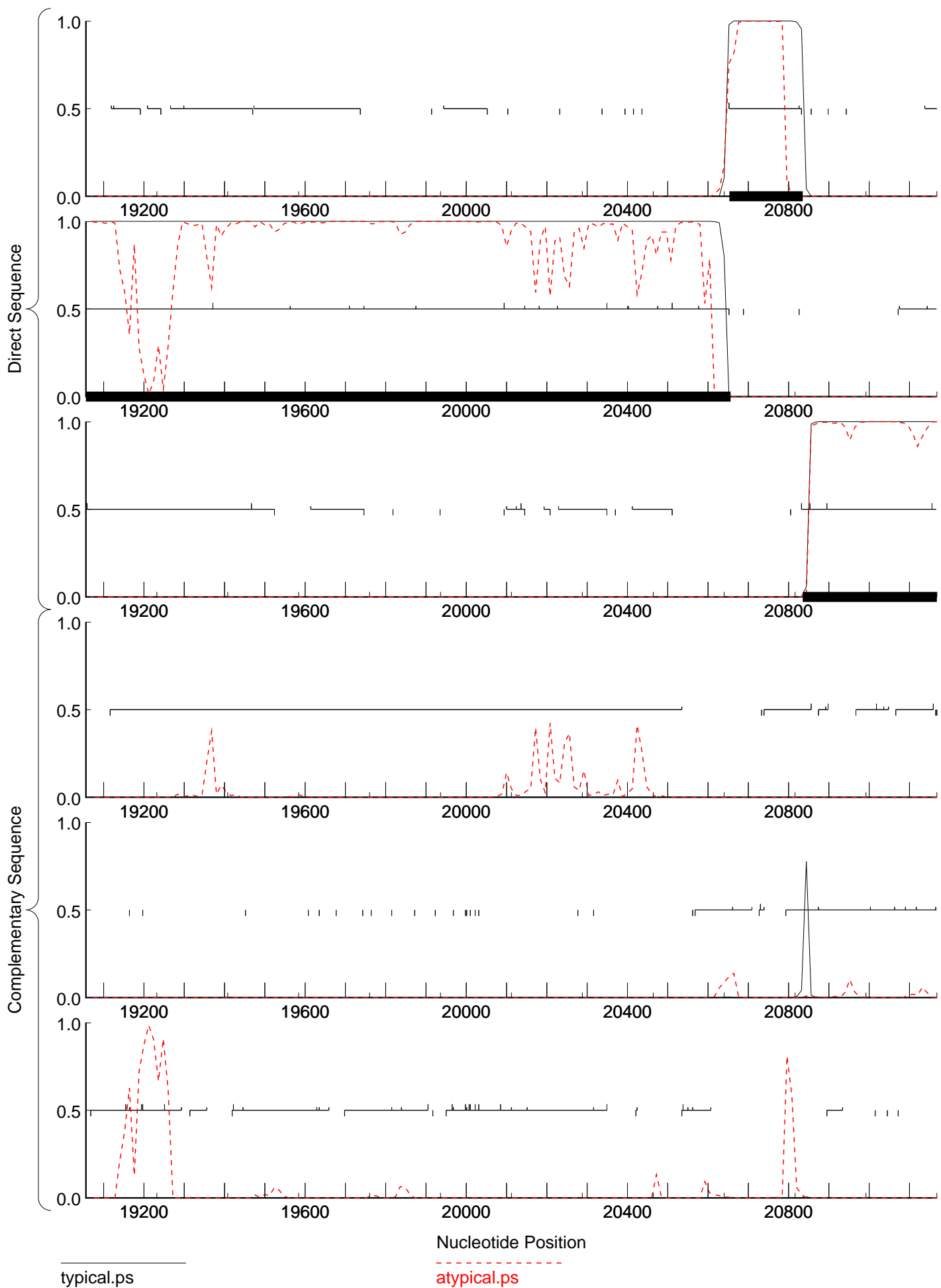


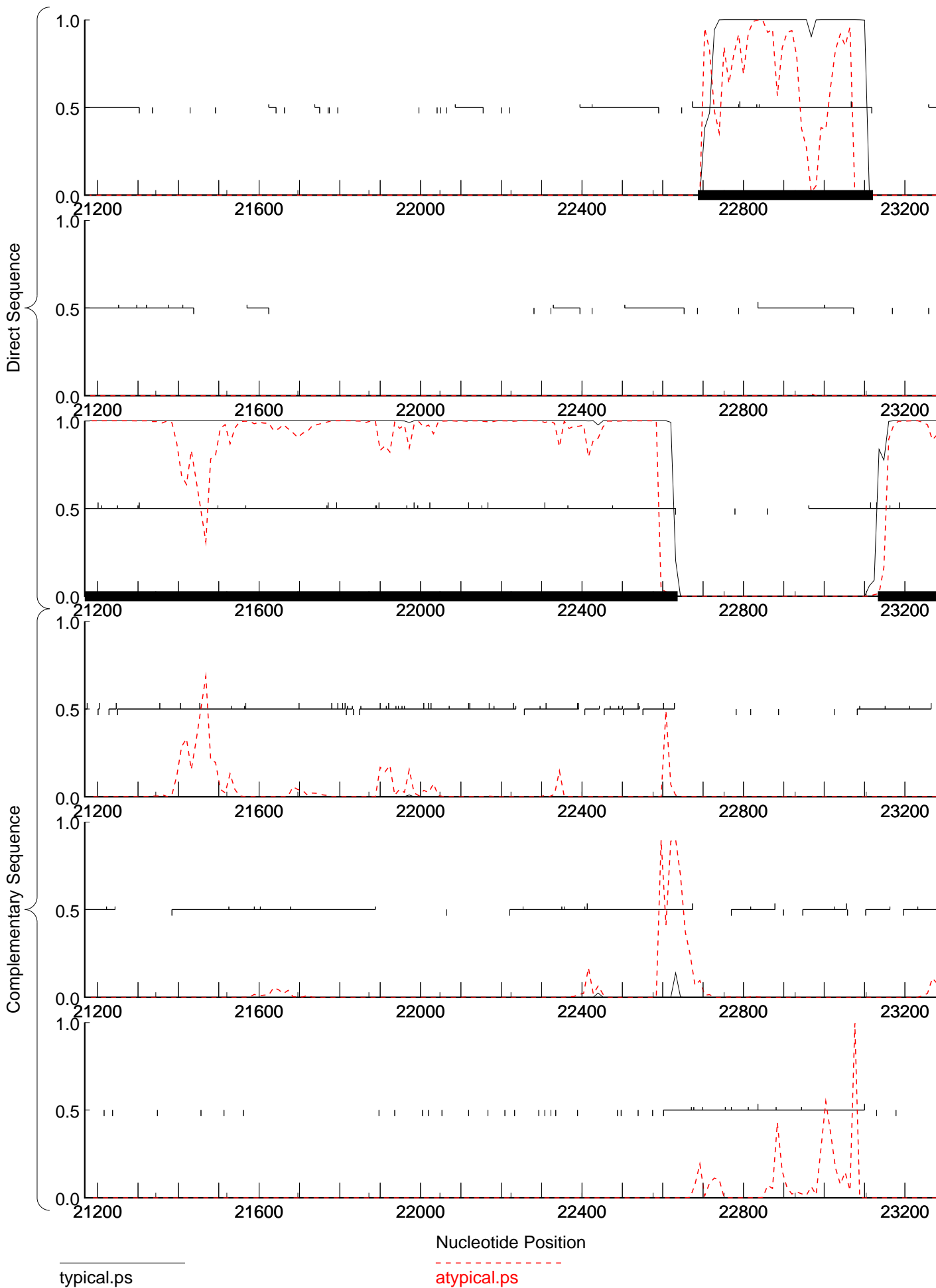


typical.ps

atypical.ps

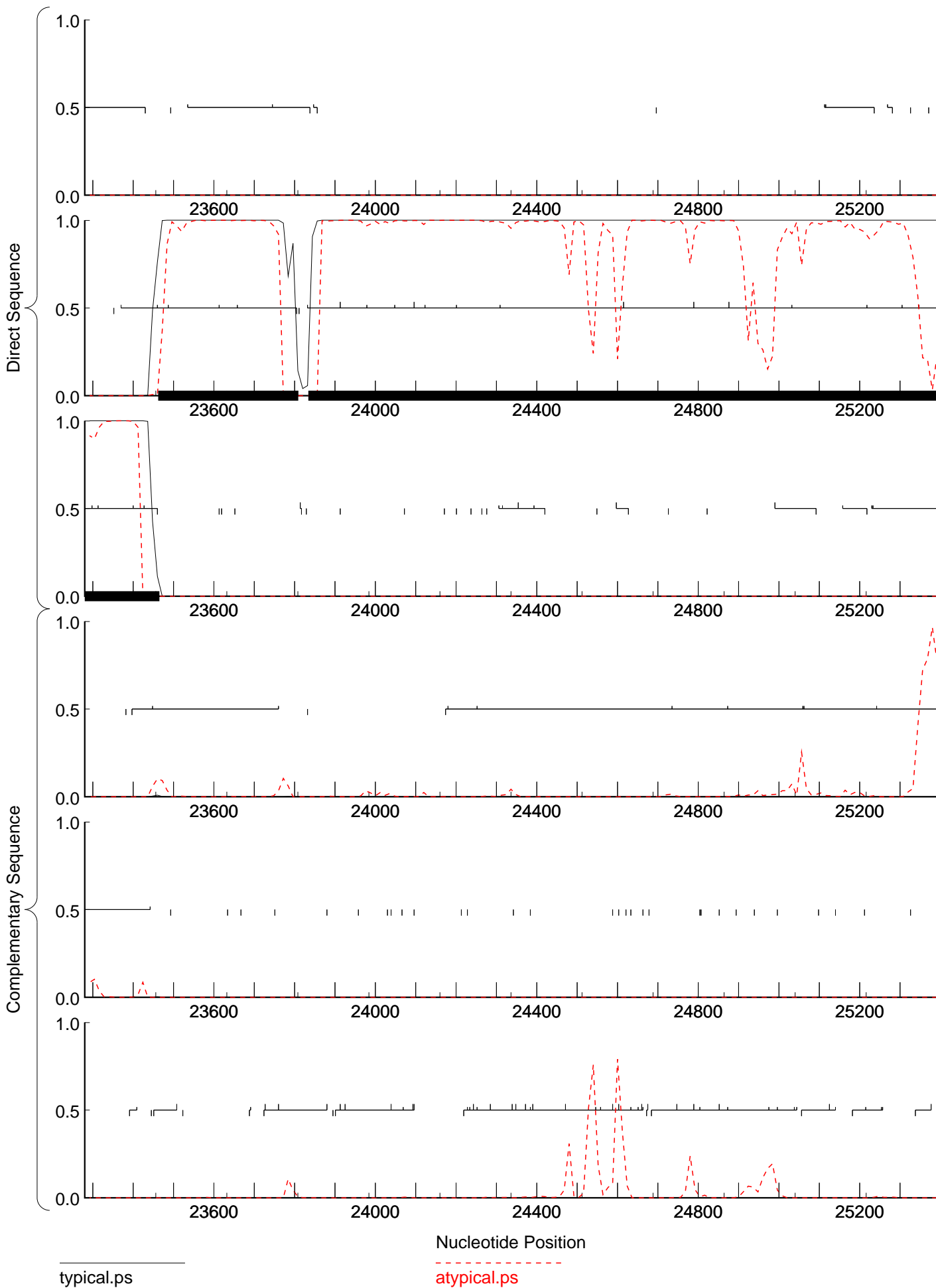


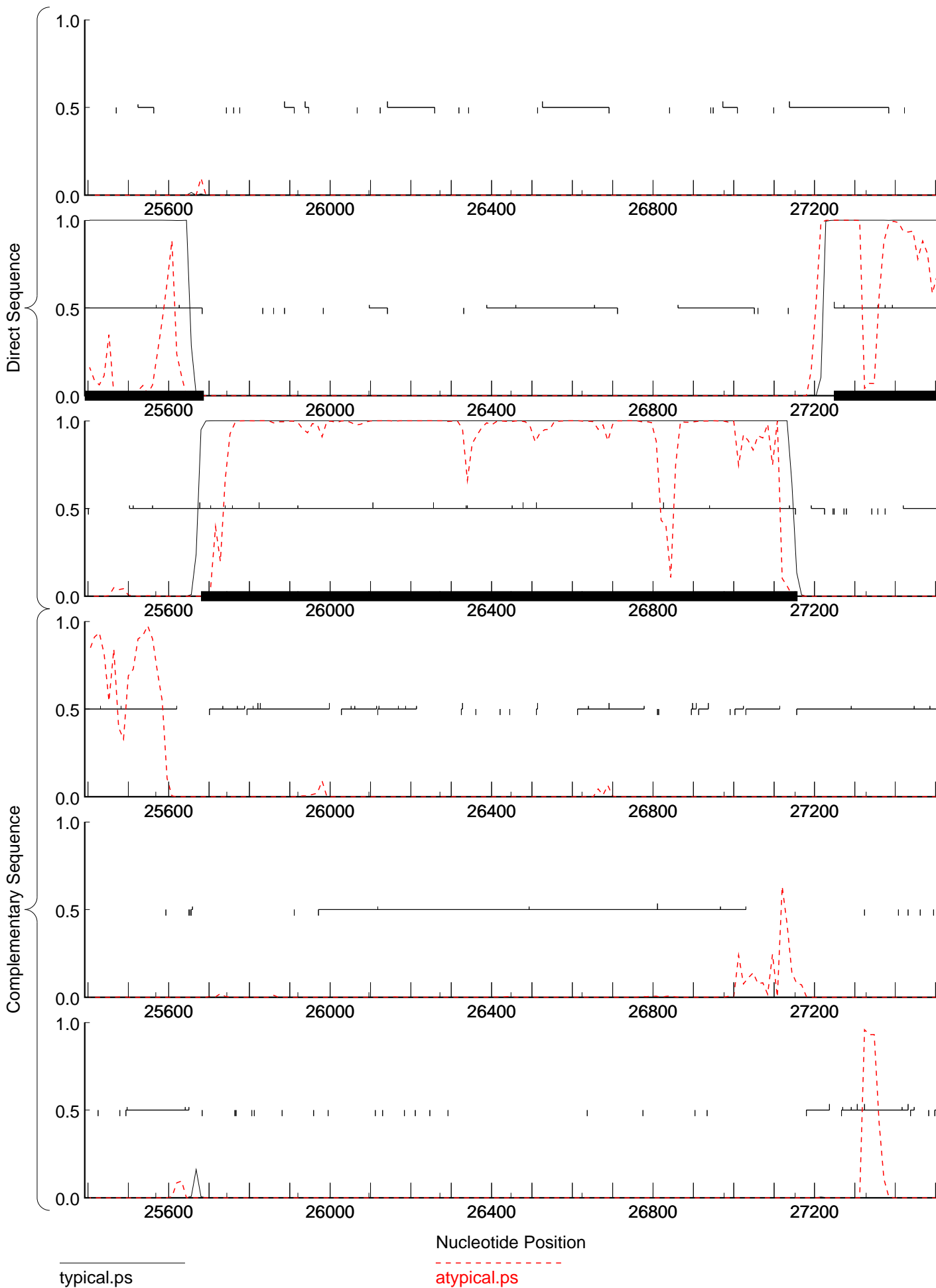


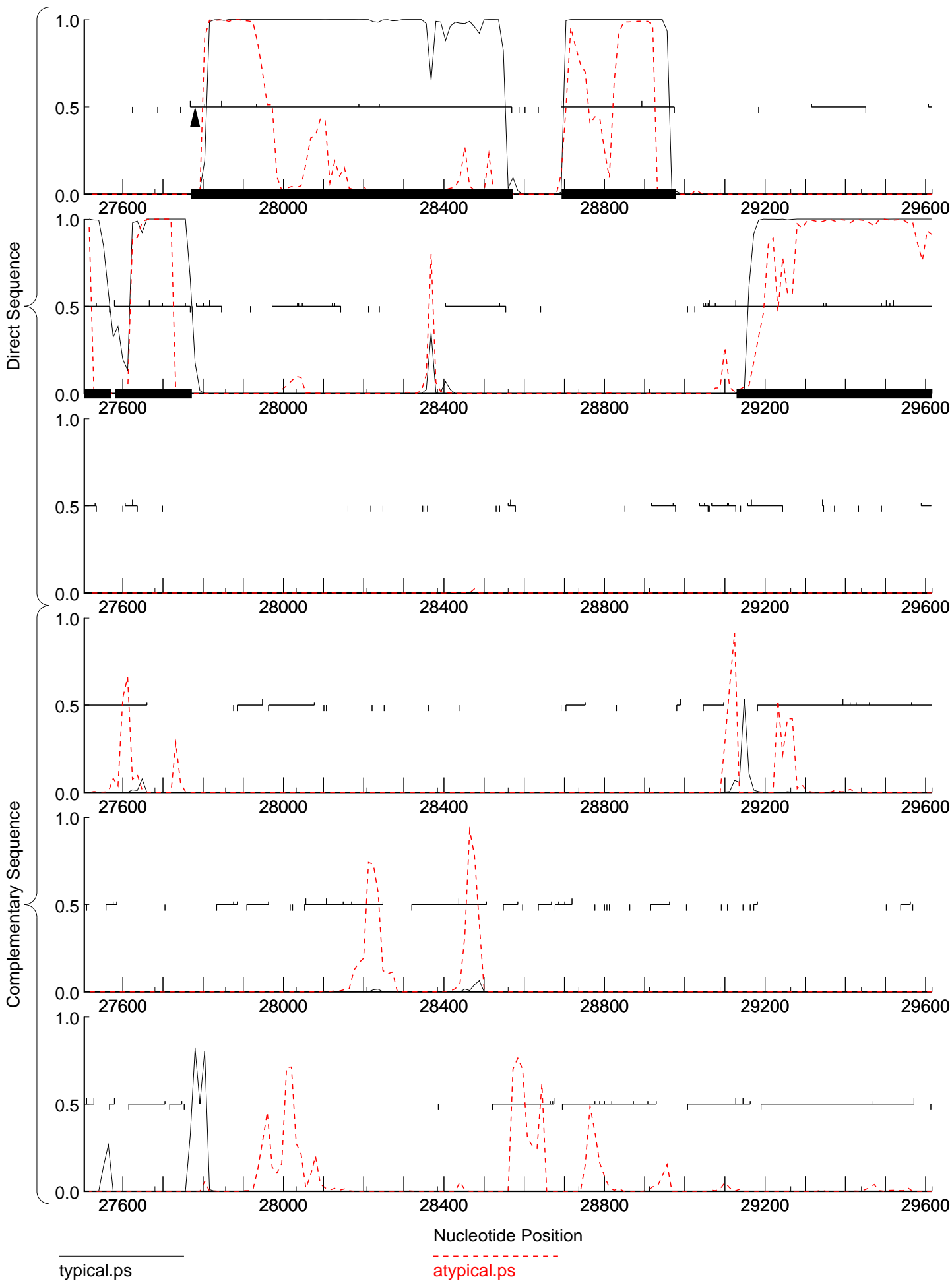


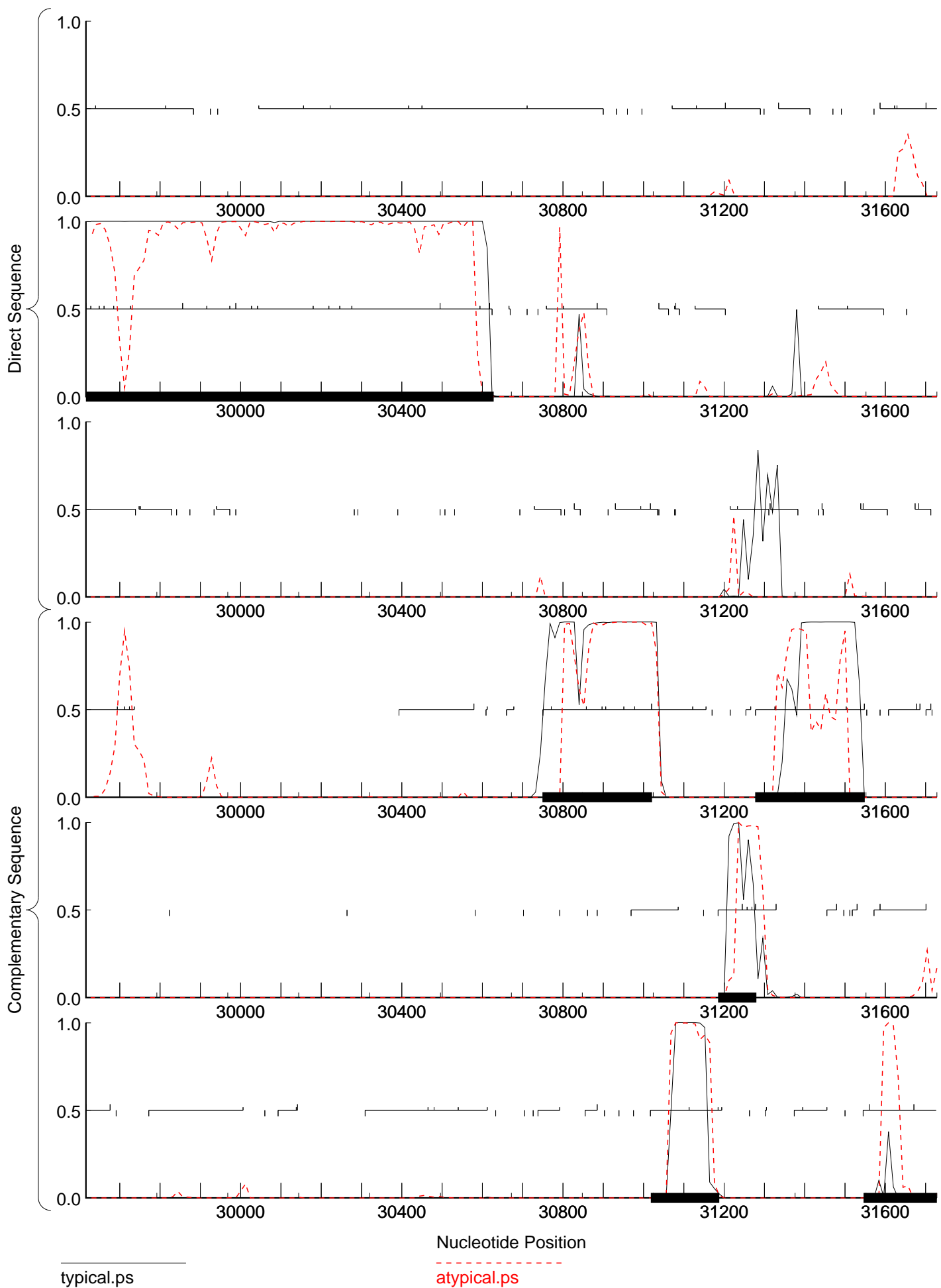
typical.ps

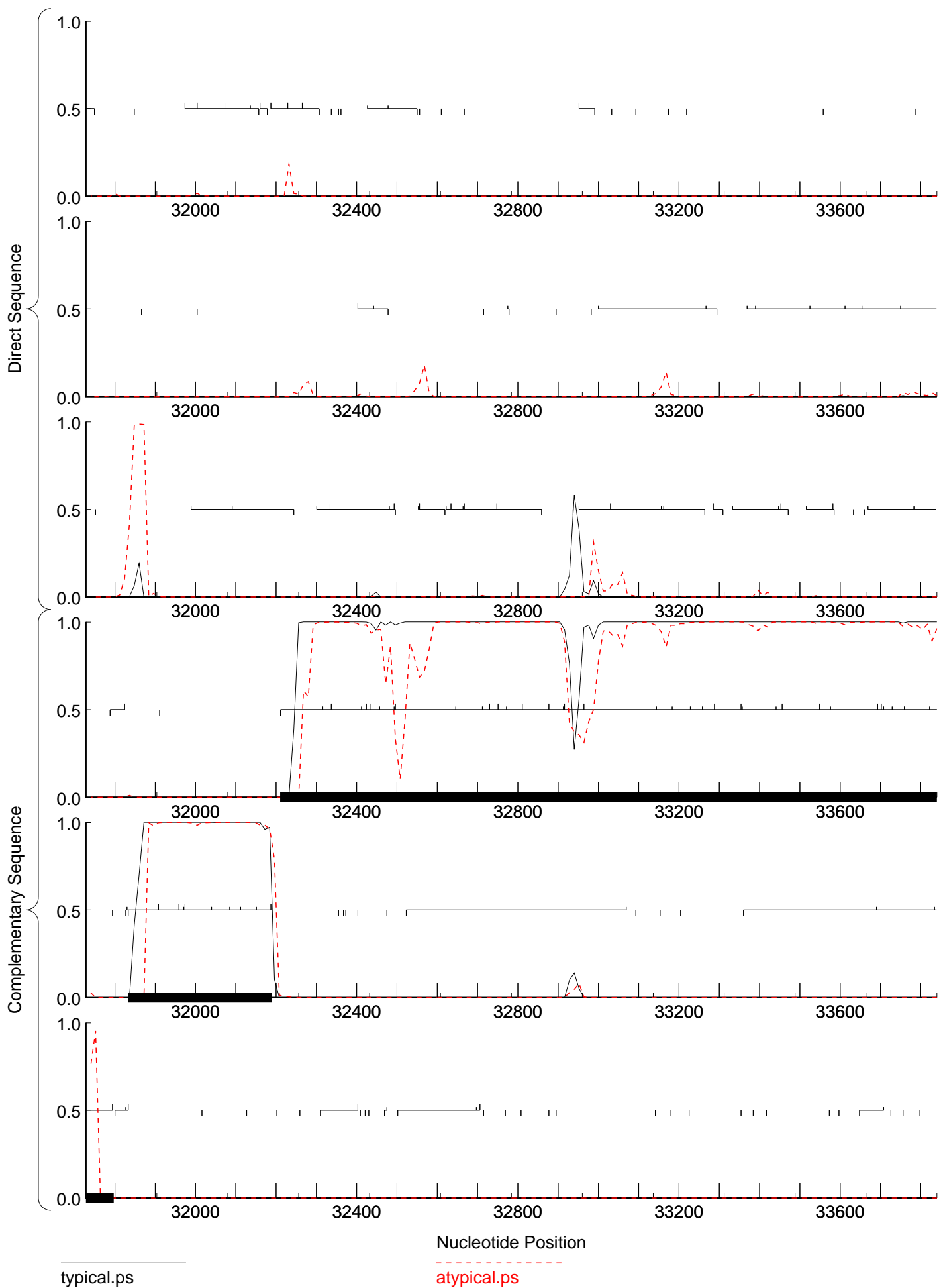
atypical.ps

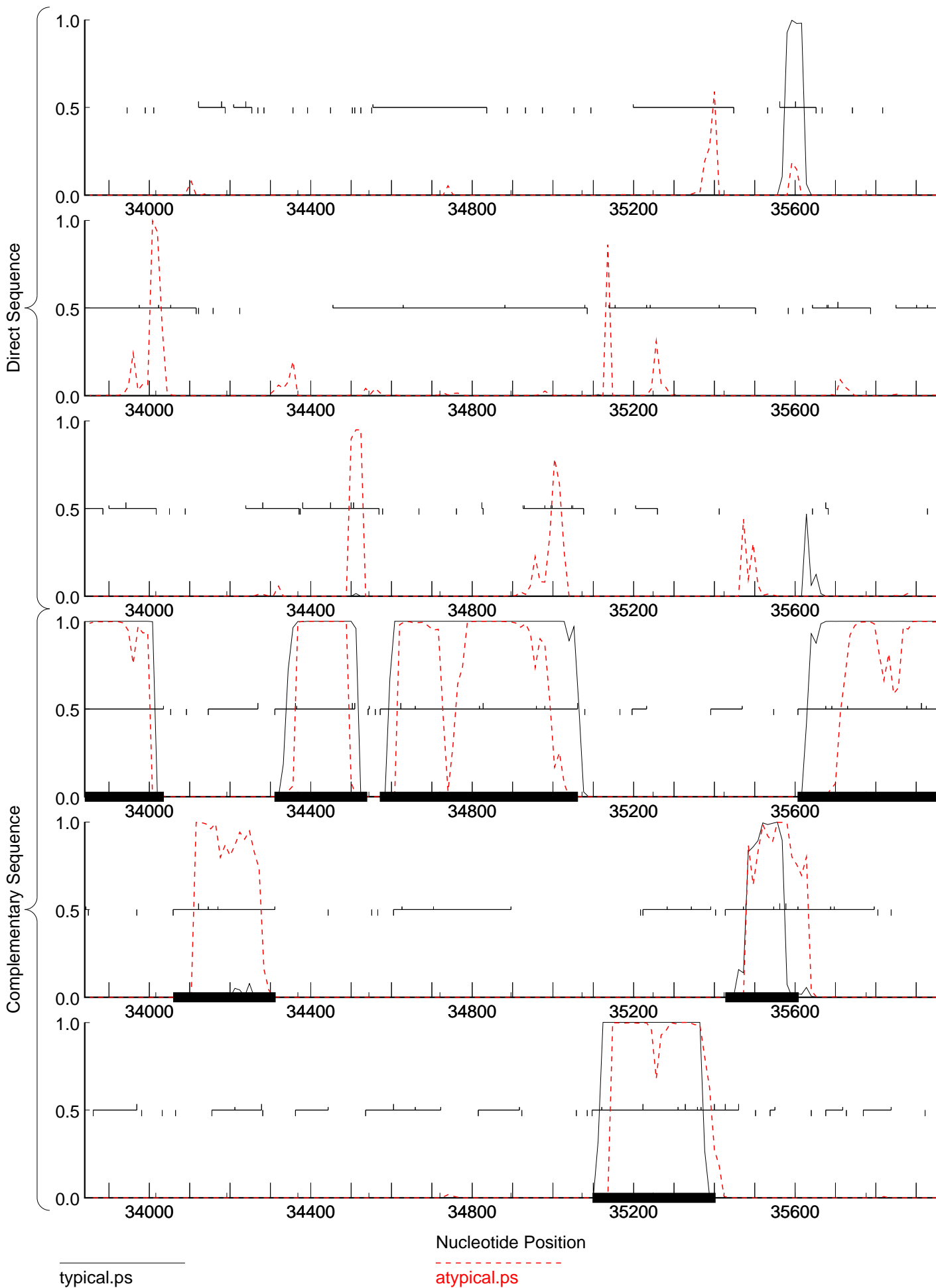


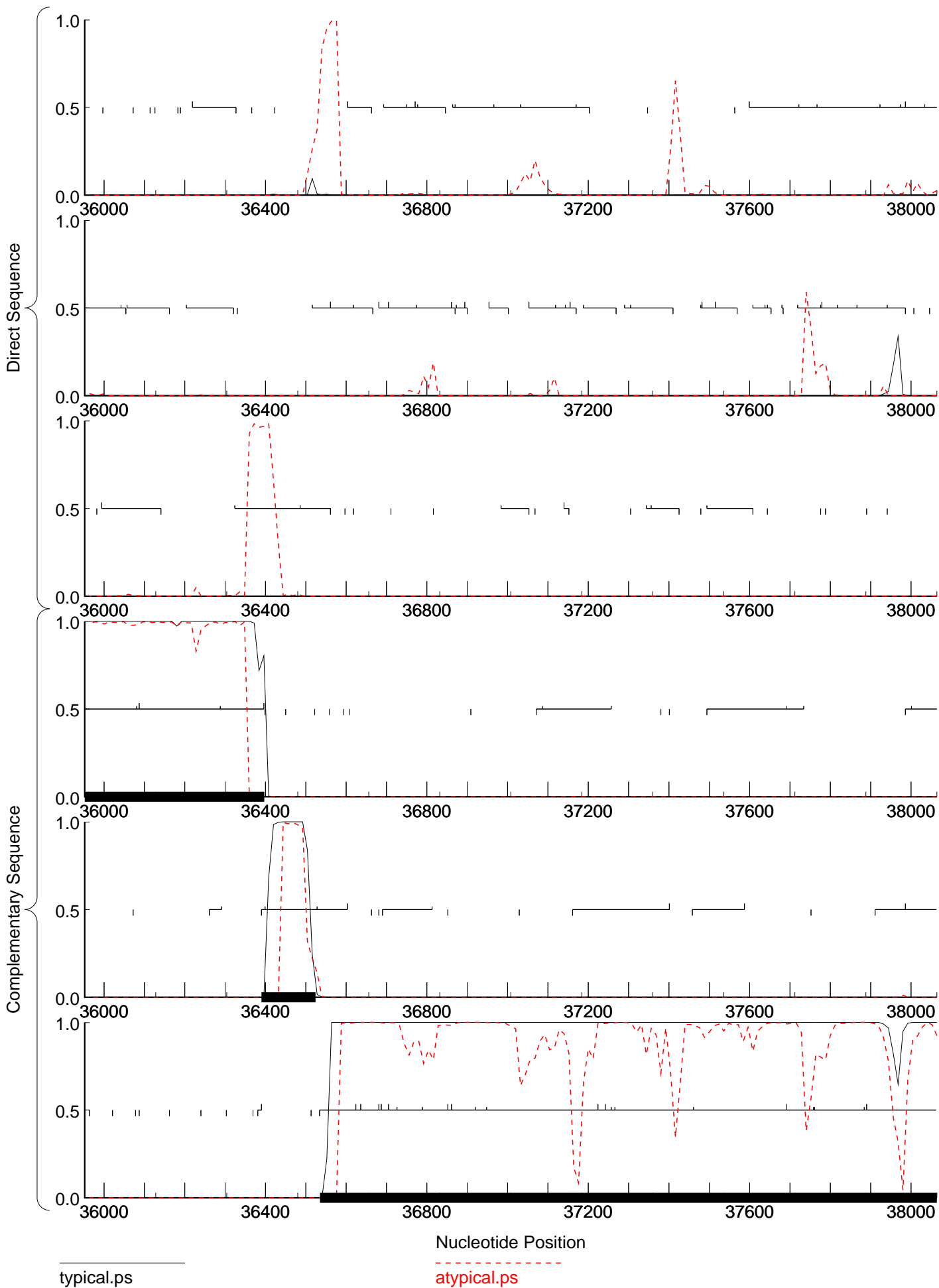


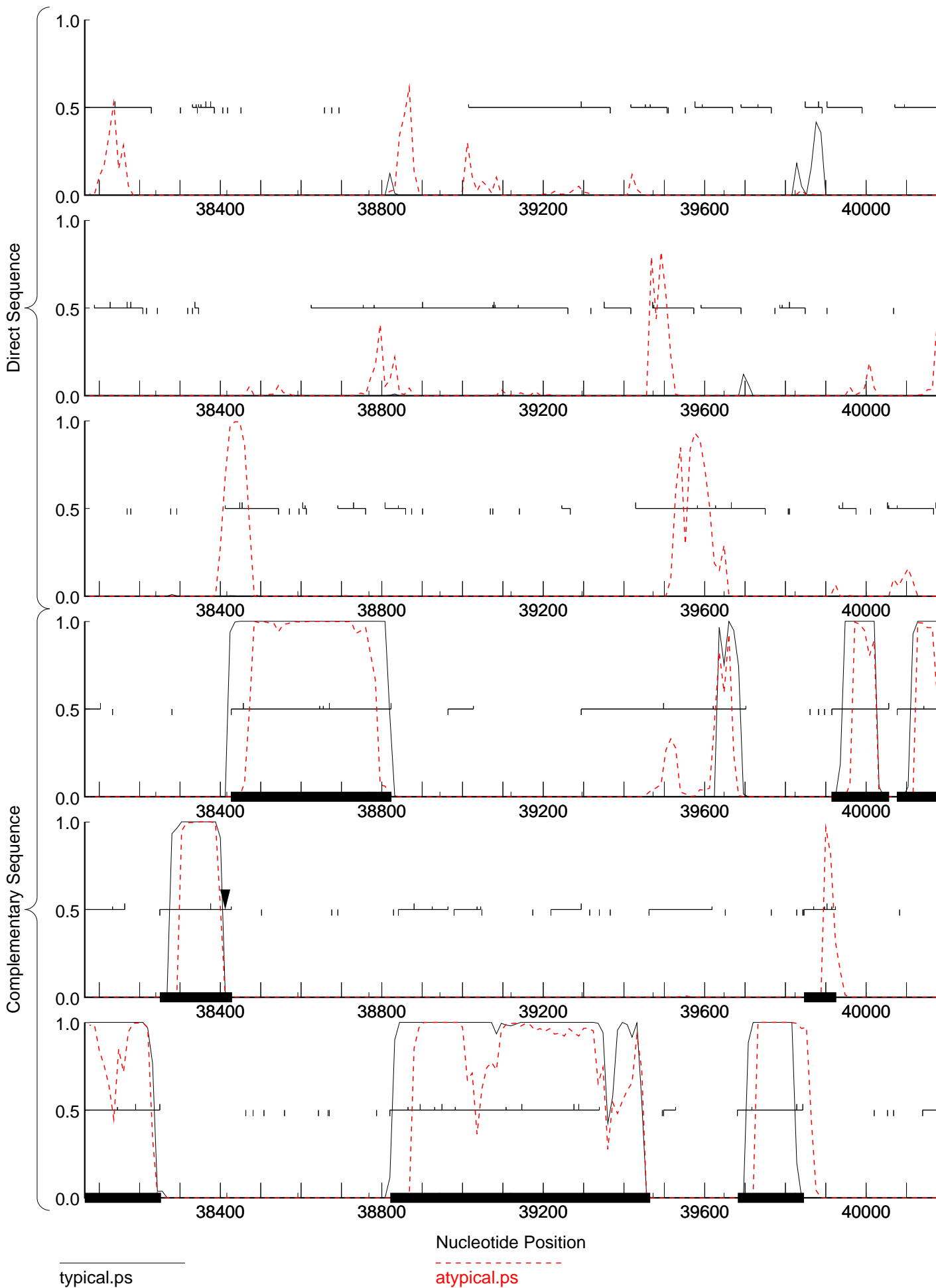






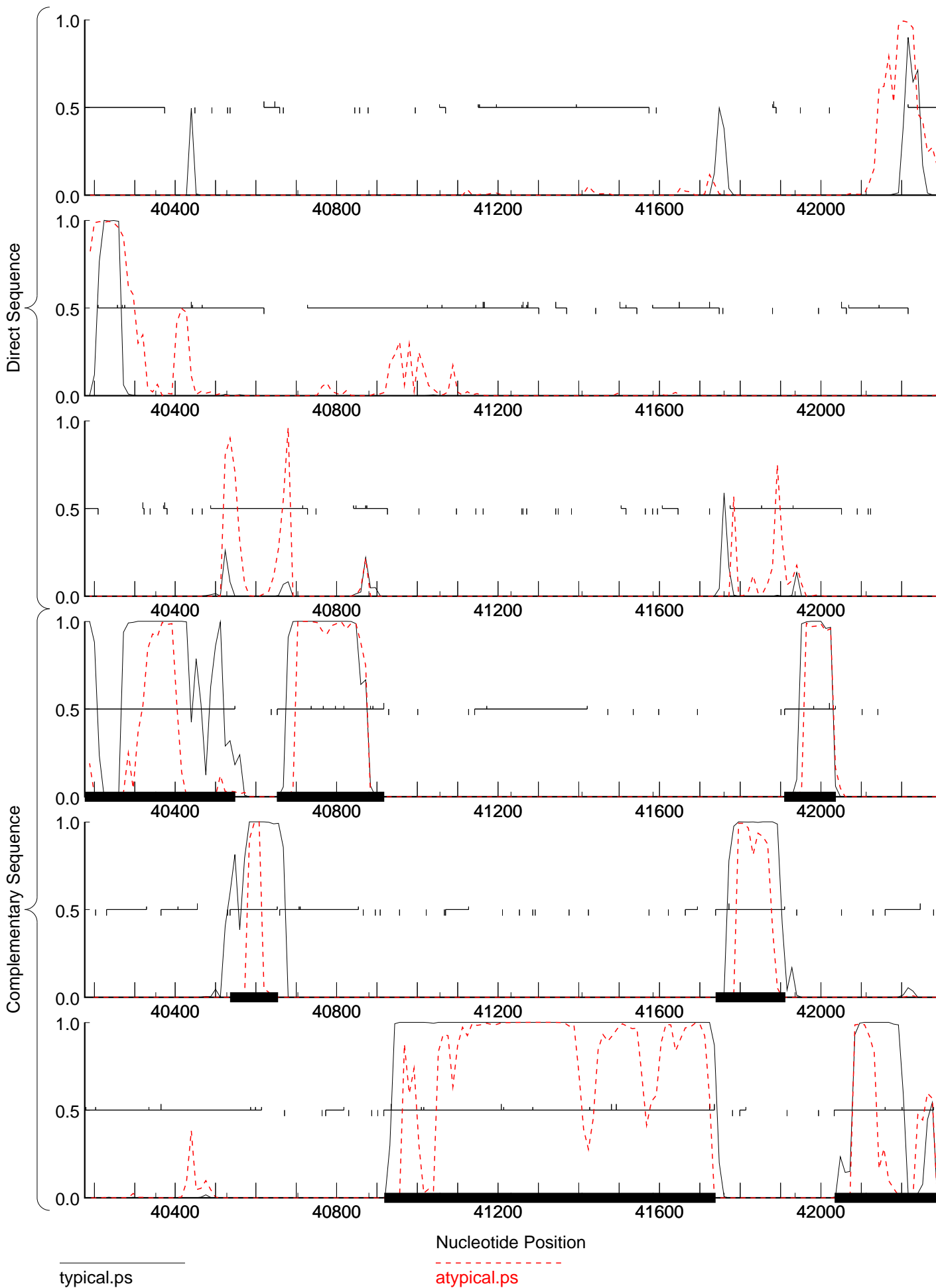


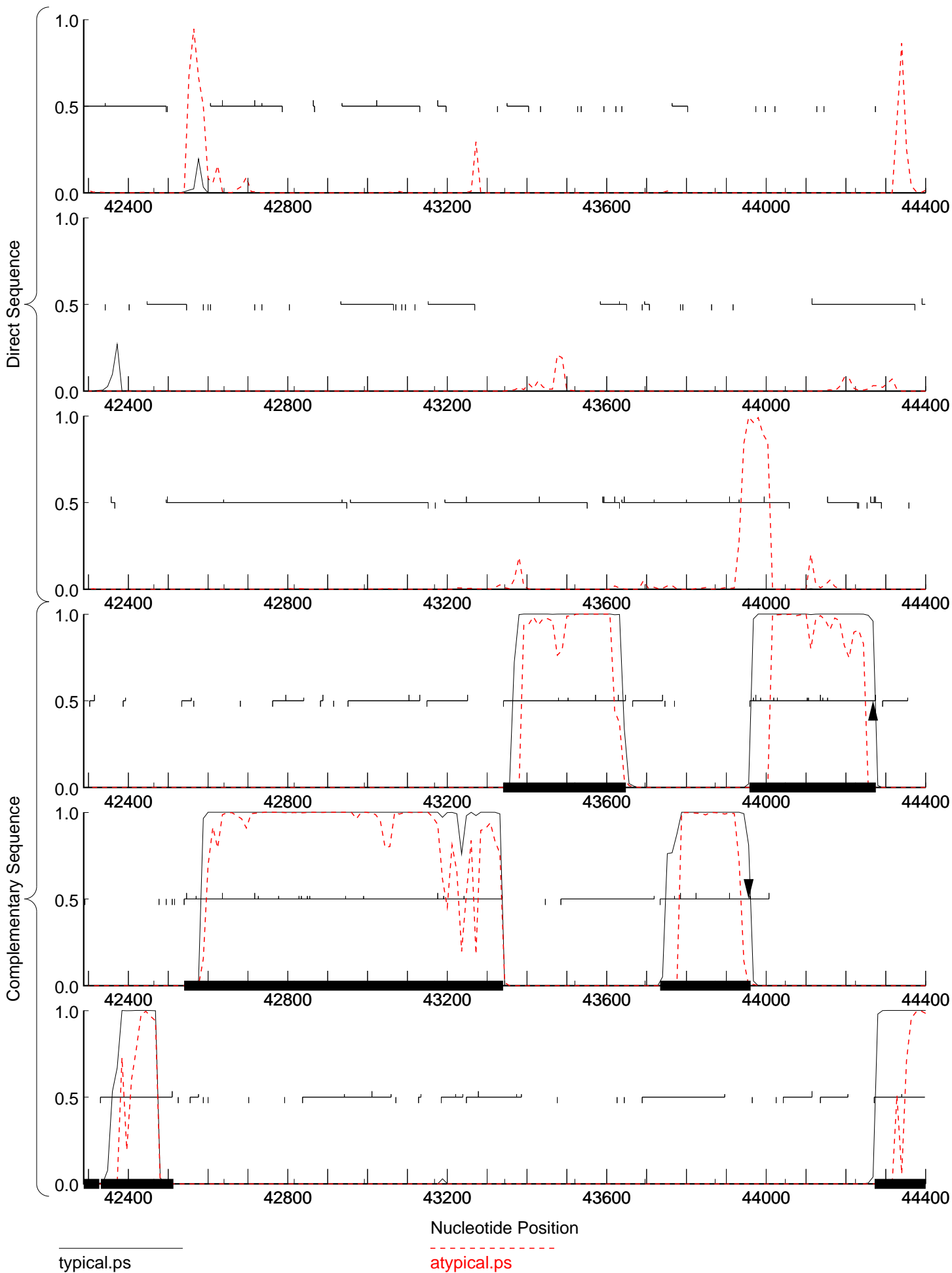




typical.ps

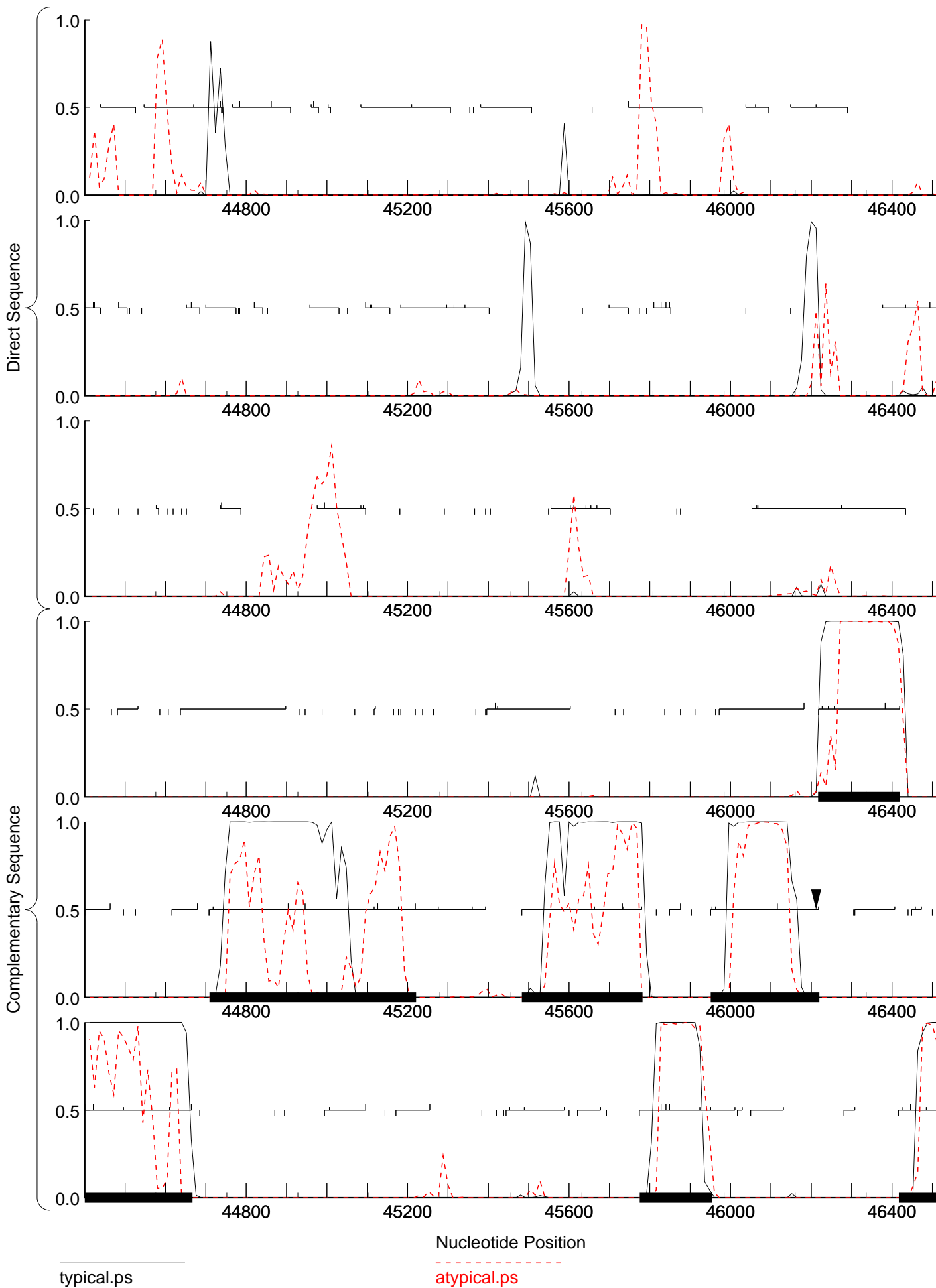
atypical.ps





typical.ps

atypical.ps



typical.ps

atypical.ps

