

GeneMark

Version 2.5p (09.08.06)

Copyright 1993 - M. Borodovsky, J. McIninch

PROGRAM INFORMATION

Sequence : Mycobacterium phage JangDynasty complete sequence, 70883 bp including 4-base 3' overhang (G
Analysis Date : 7/26/17 at 16:43:14
Pages : 35
Sequence Length : 70883 bp
GC Content : 65.43%

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,

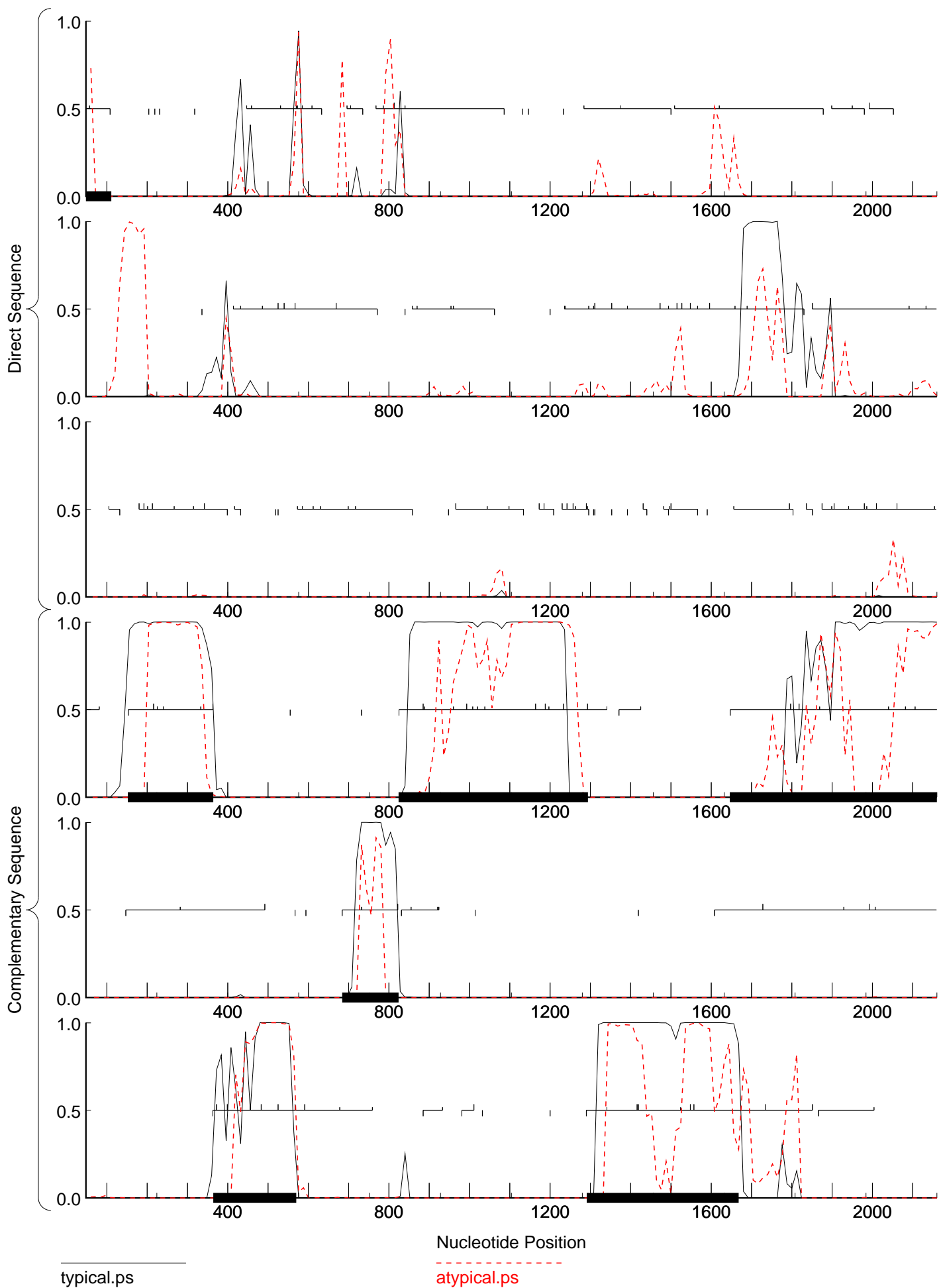
Matrix notes & comments

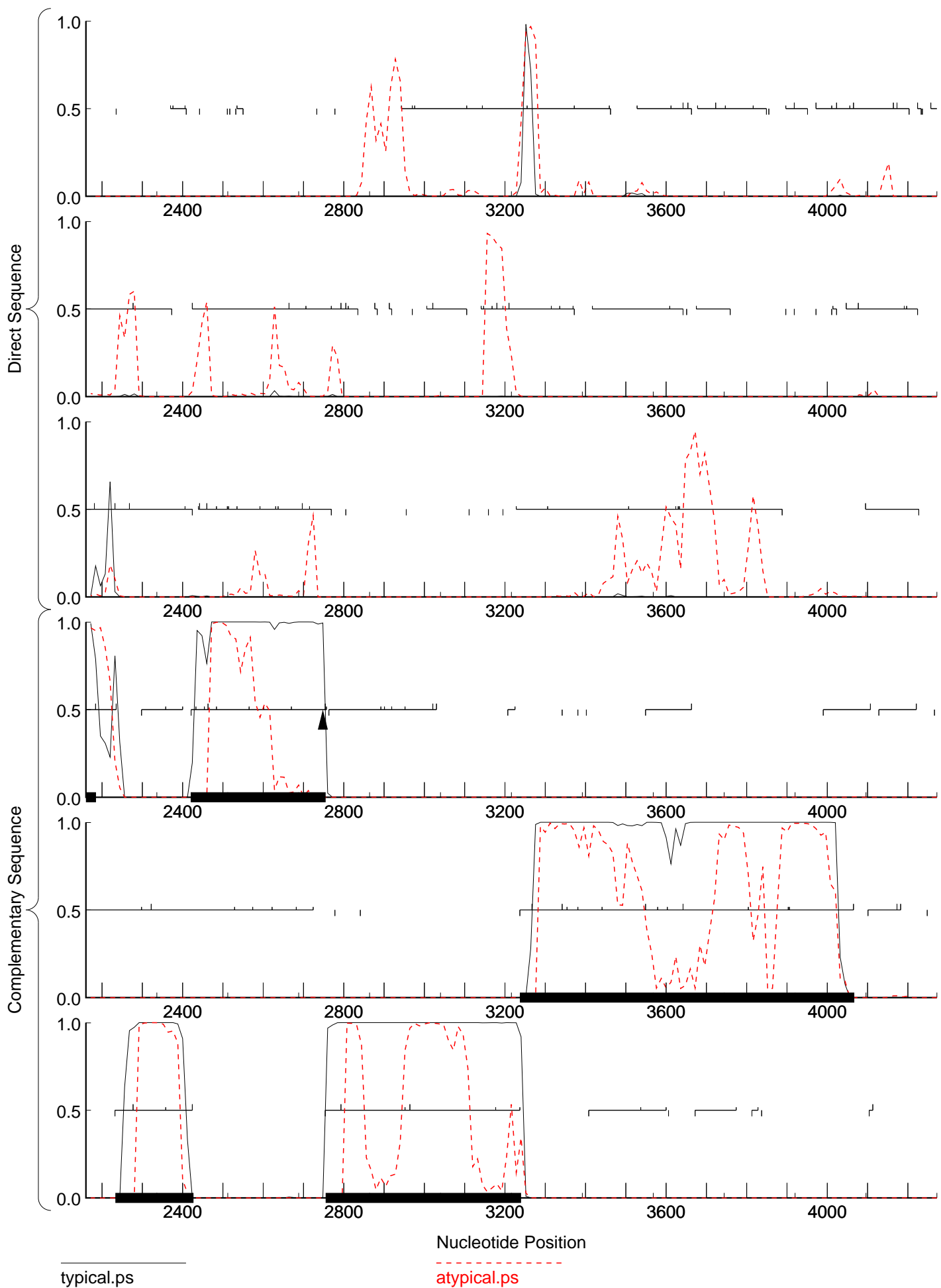
Training set derived by GeneMarkS, 4.30 October 2014
Wed Jul 26 16:43:14 2017

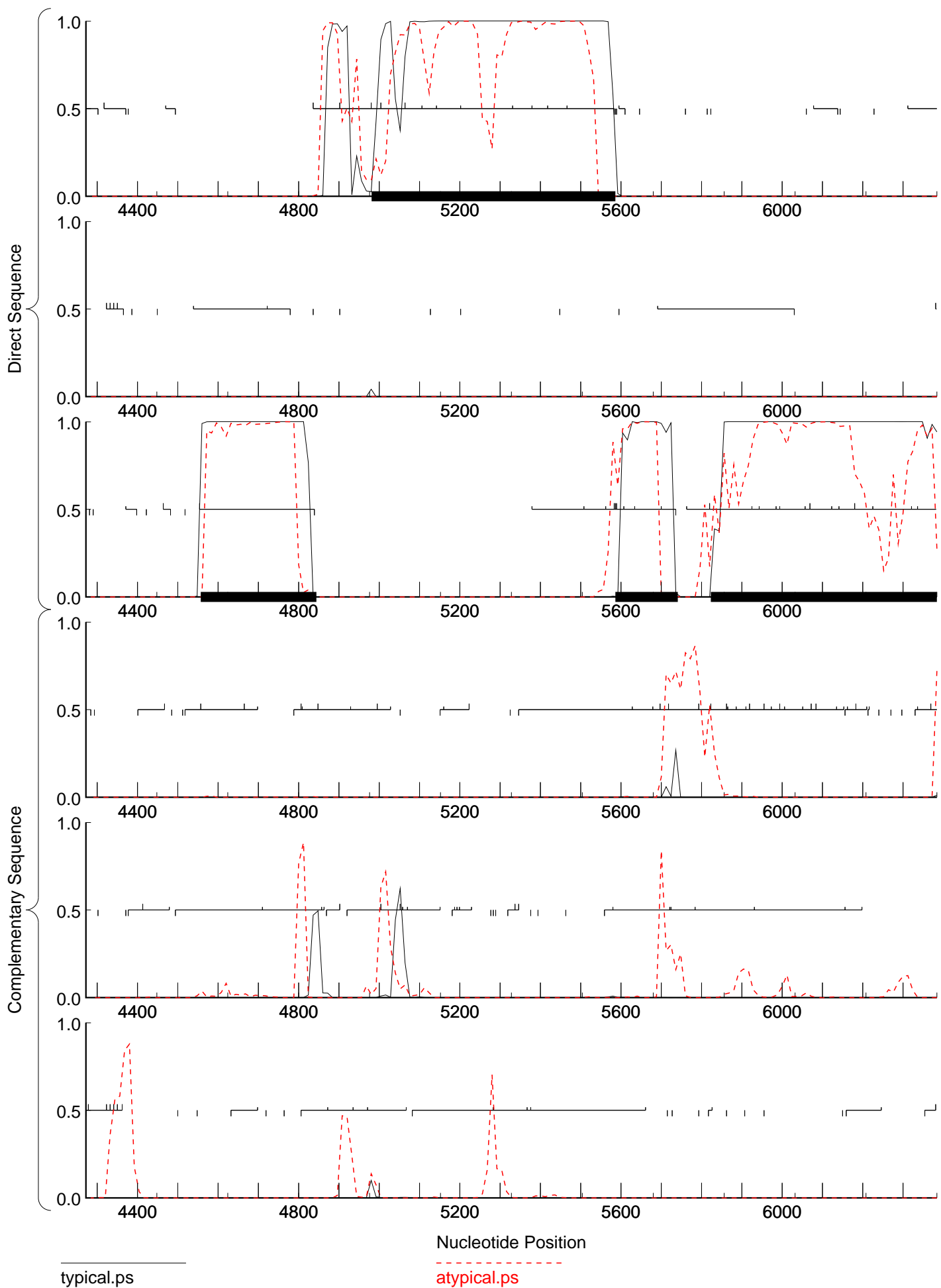
MATRIX INFORMATION

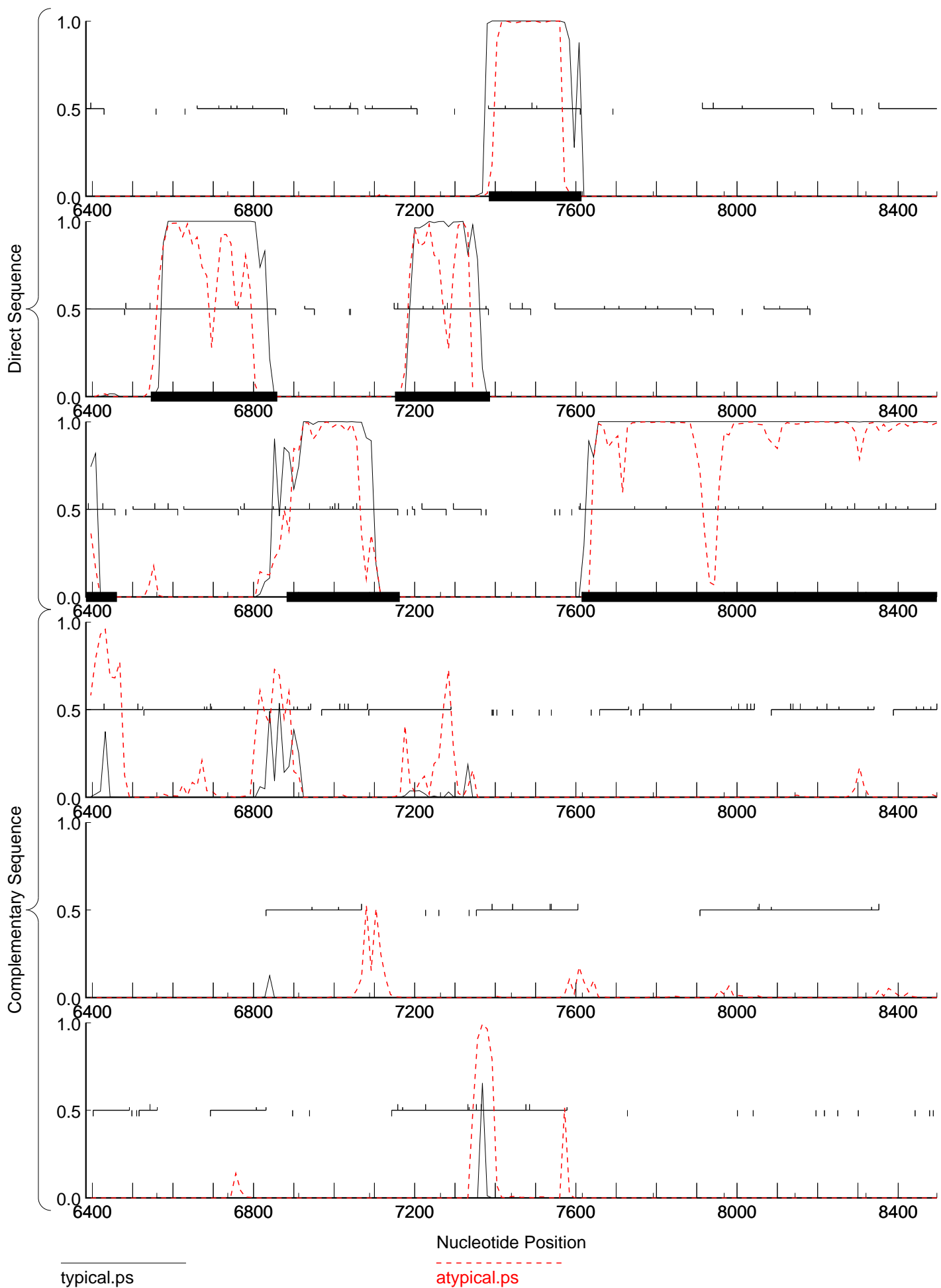
Matrix : -
Author : -
Order : 4

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



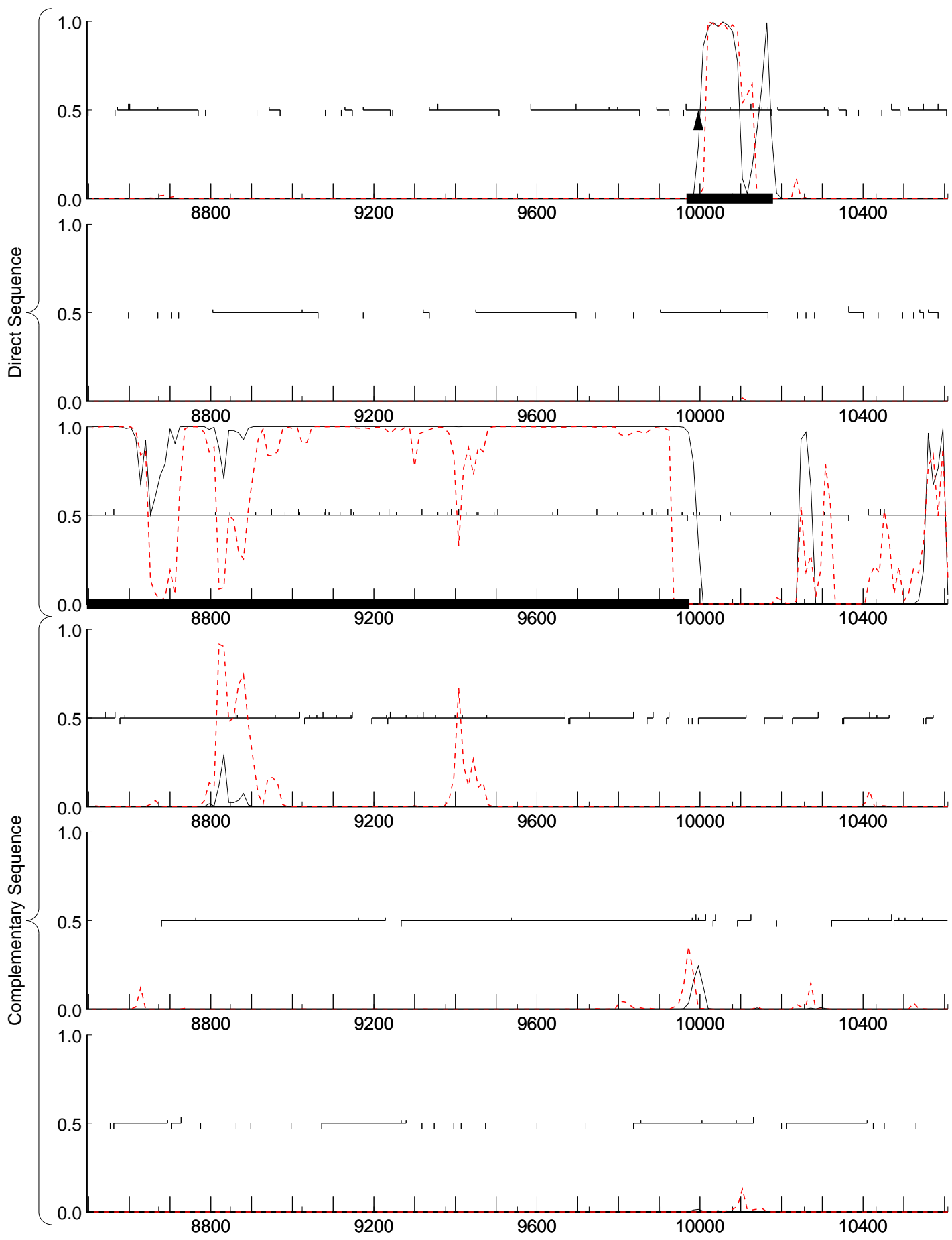






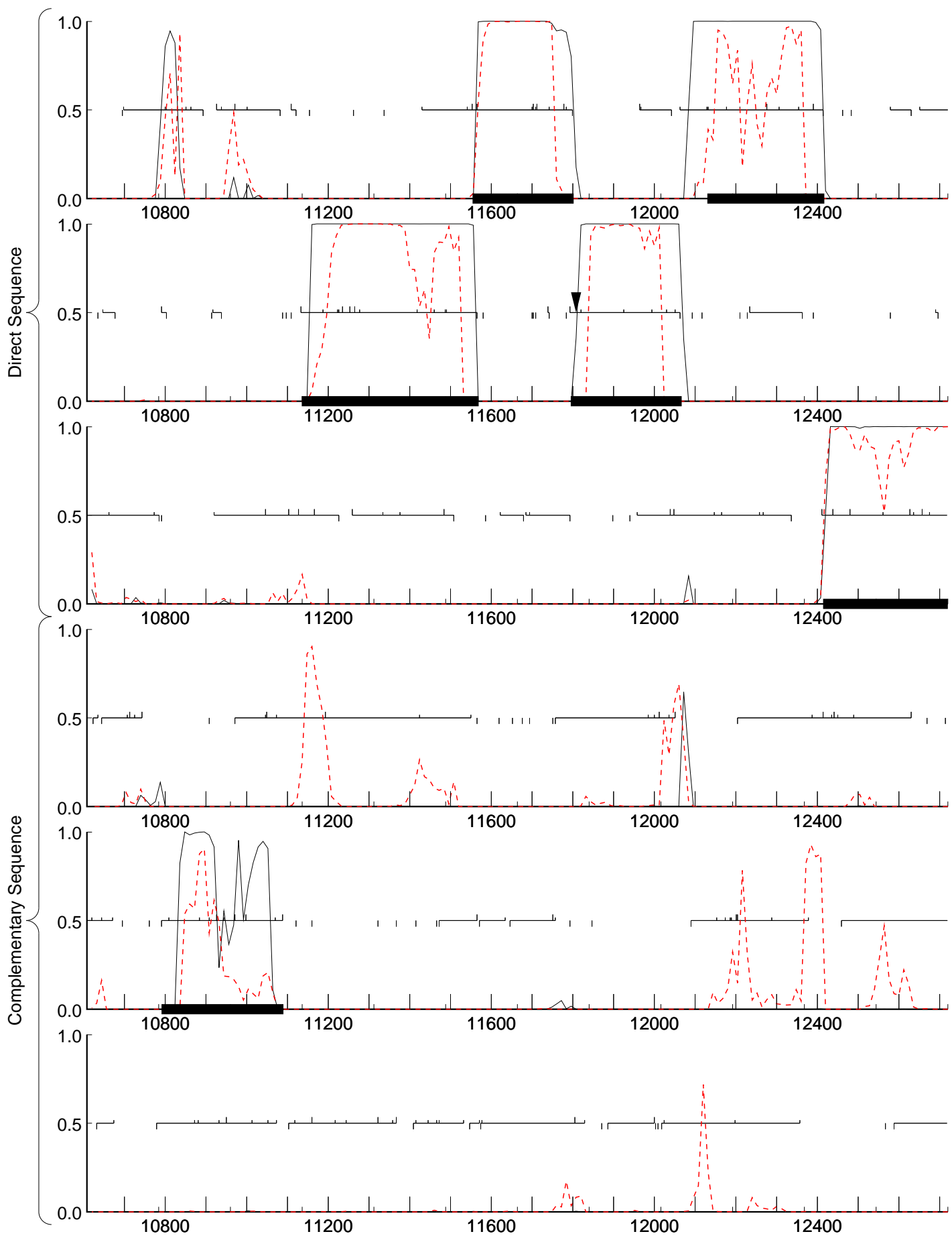
typical.ps

atypical.ps



typical.ps

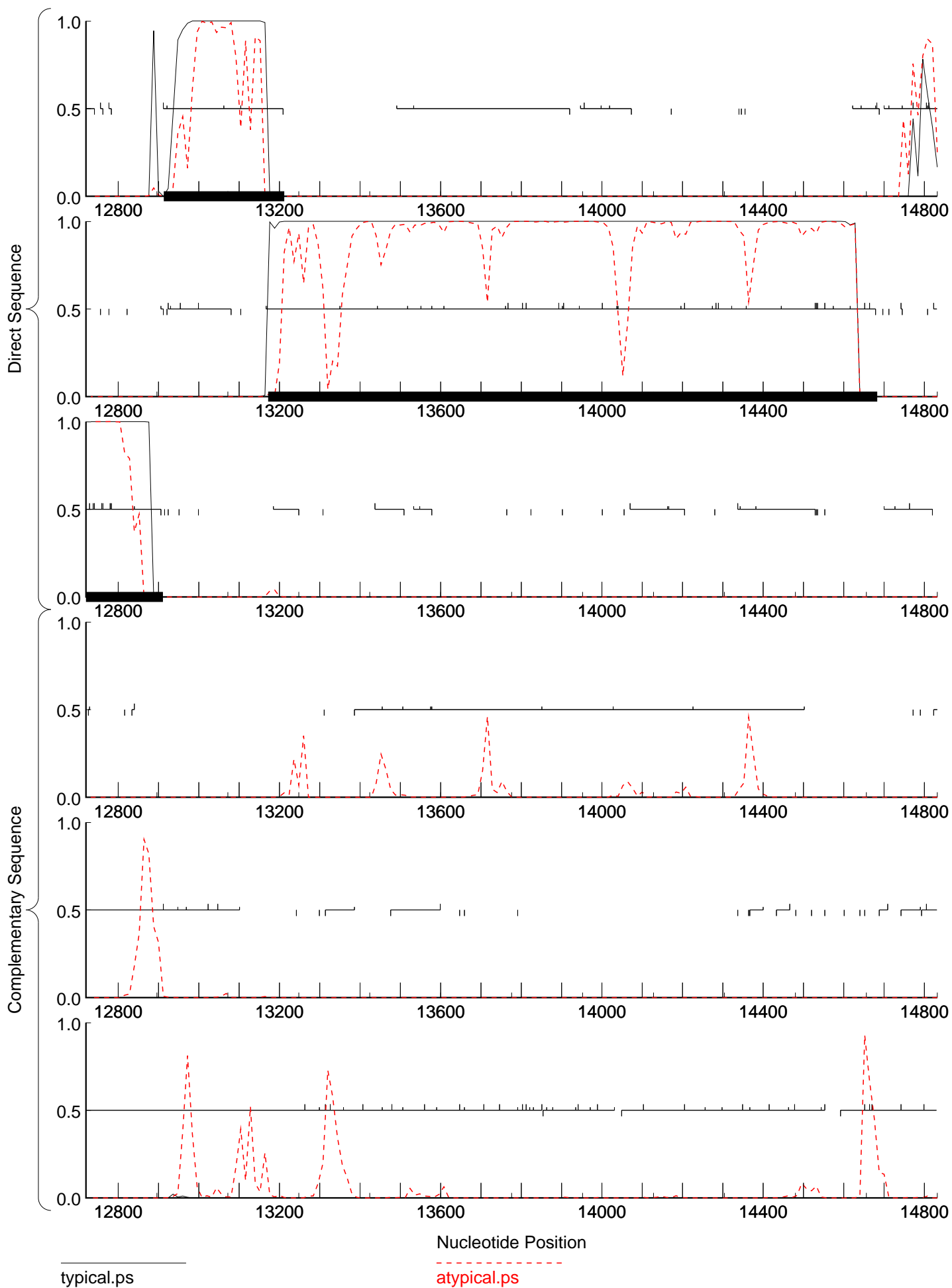
atypical.ps

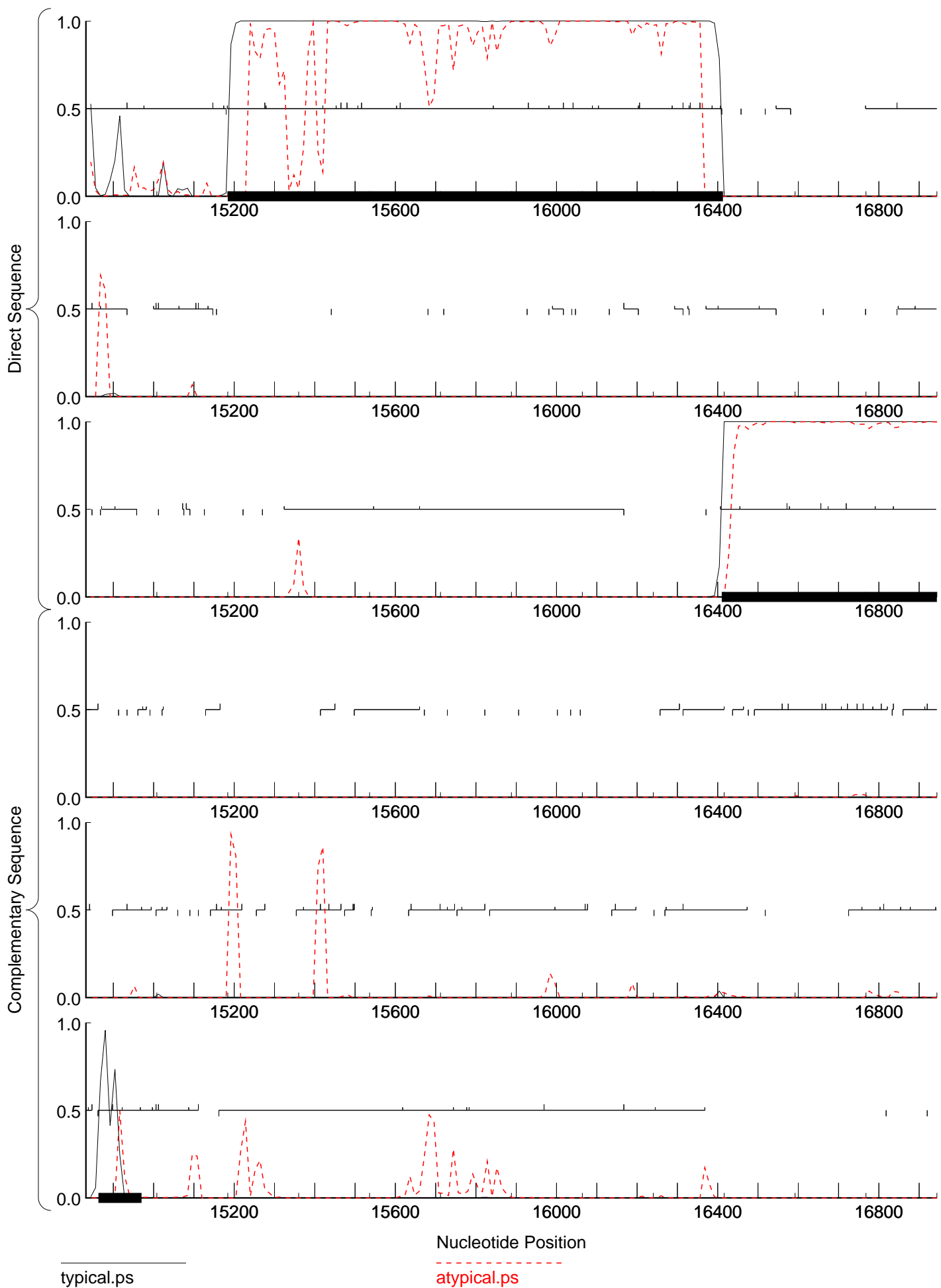


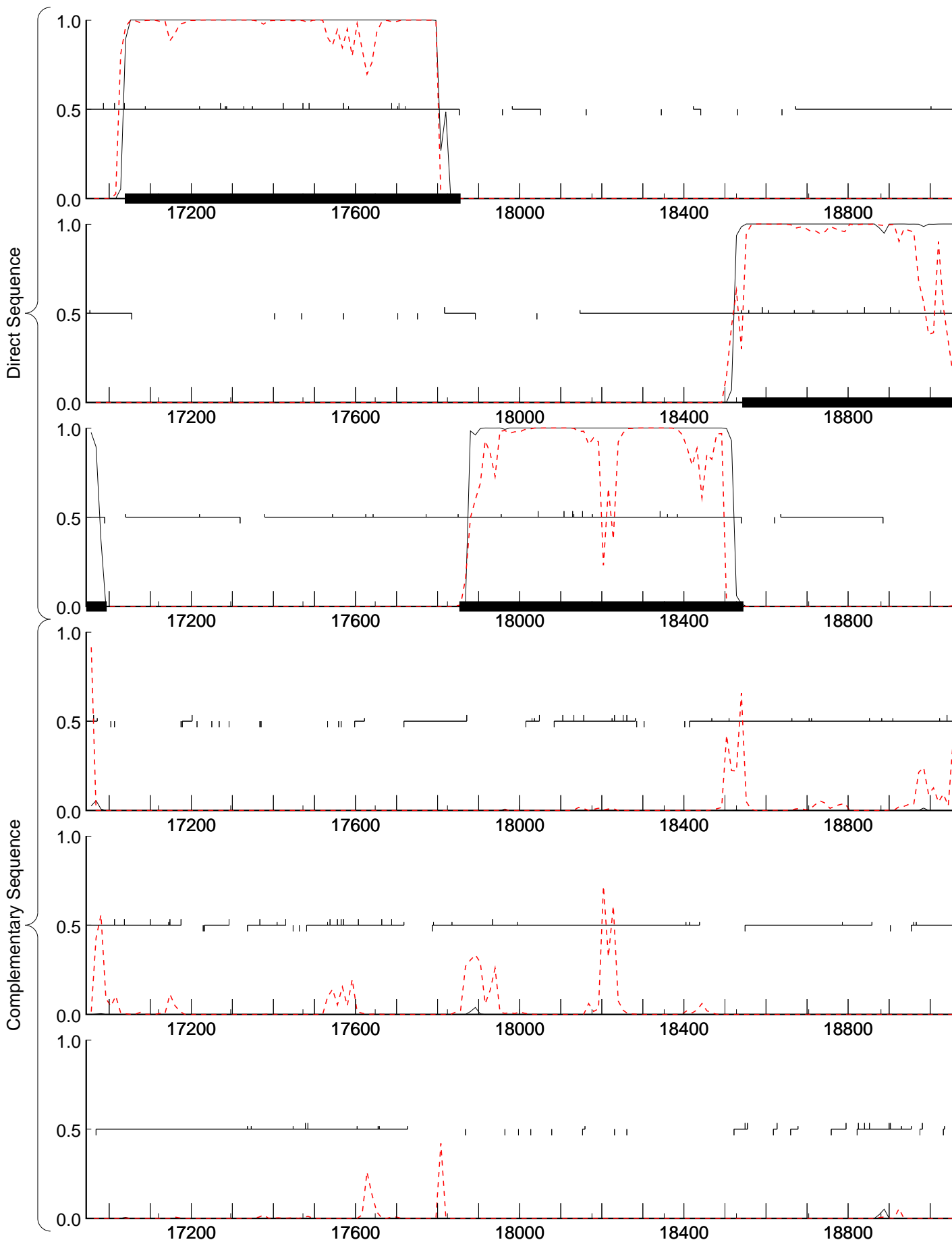
Nucleotide Position

typical.ps

atypical.ps

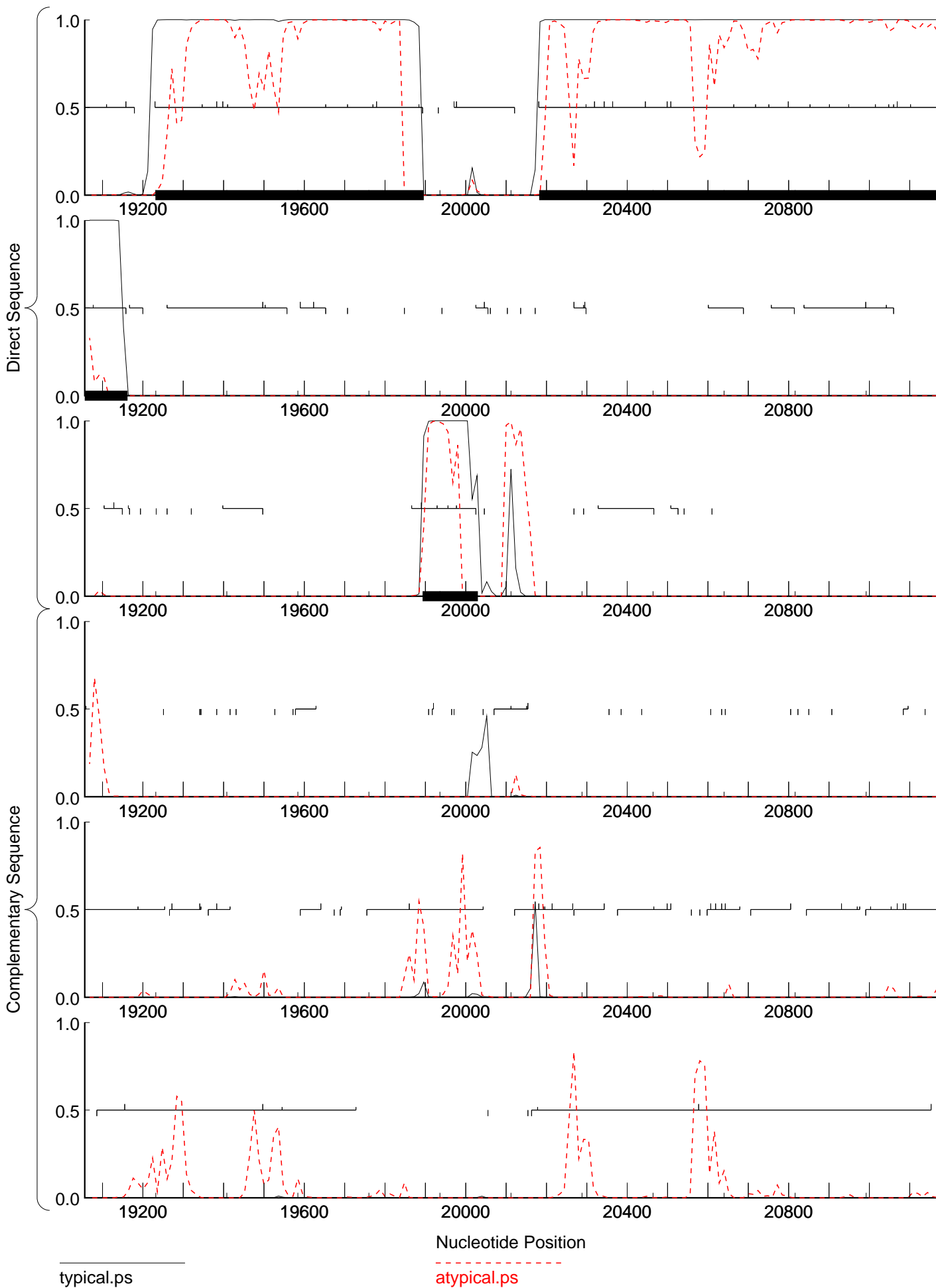


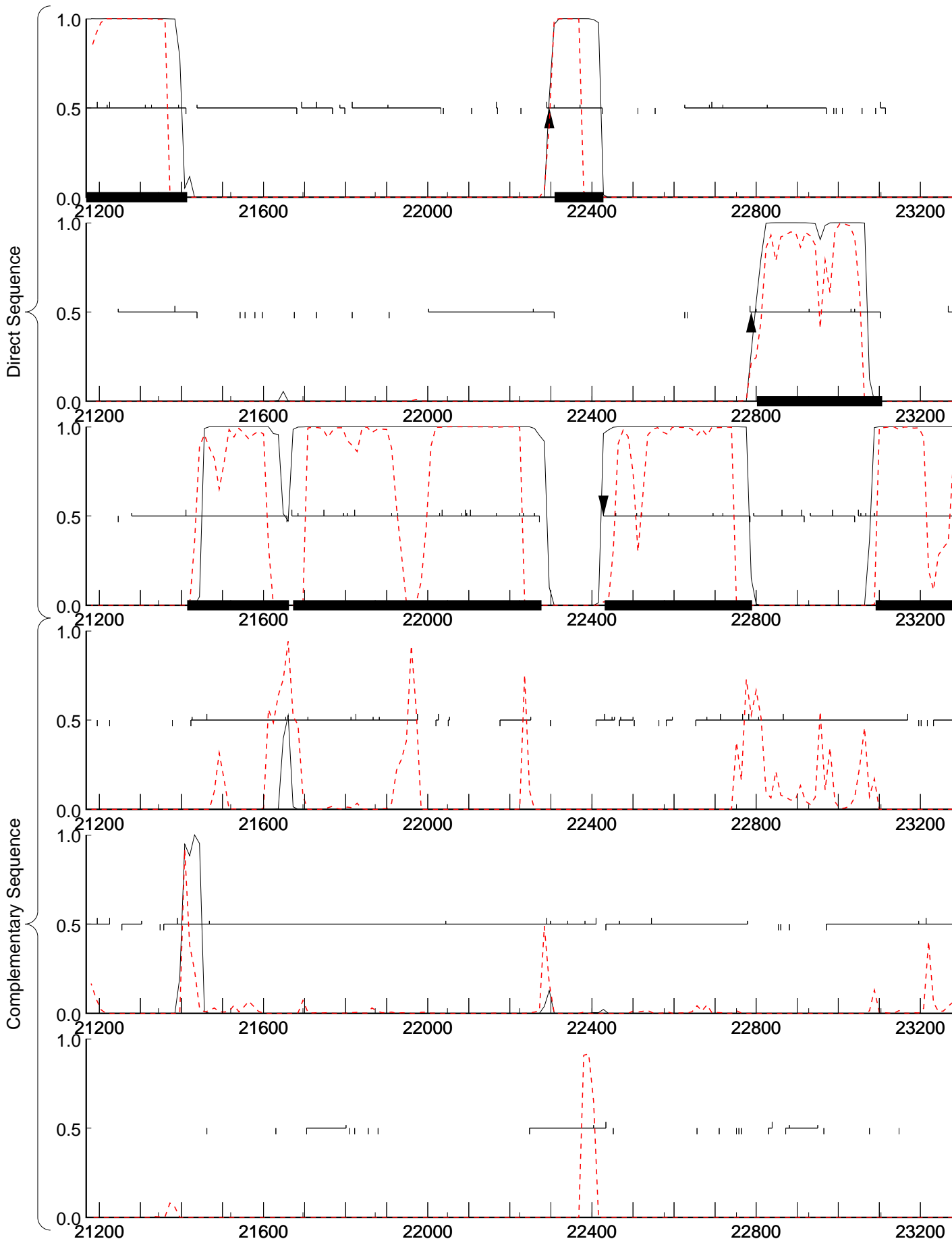




typical.ps

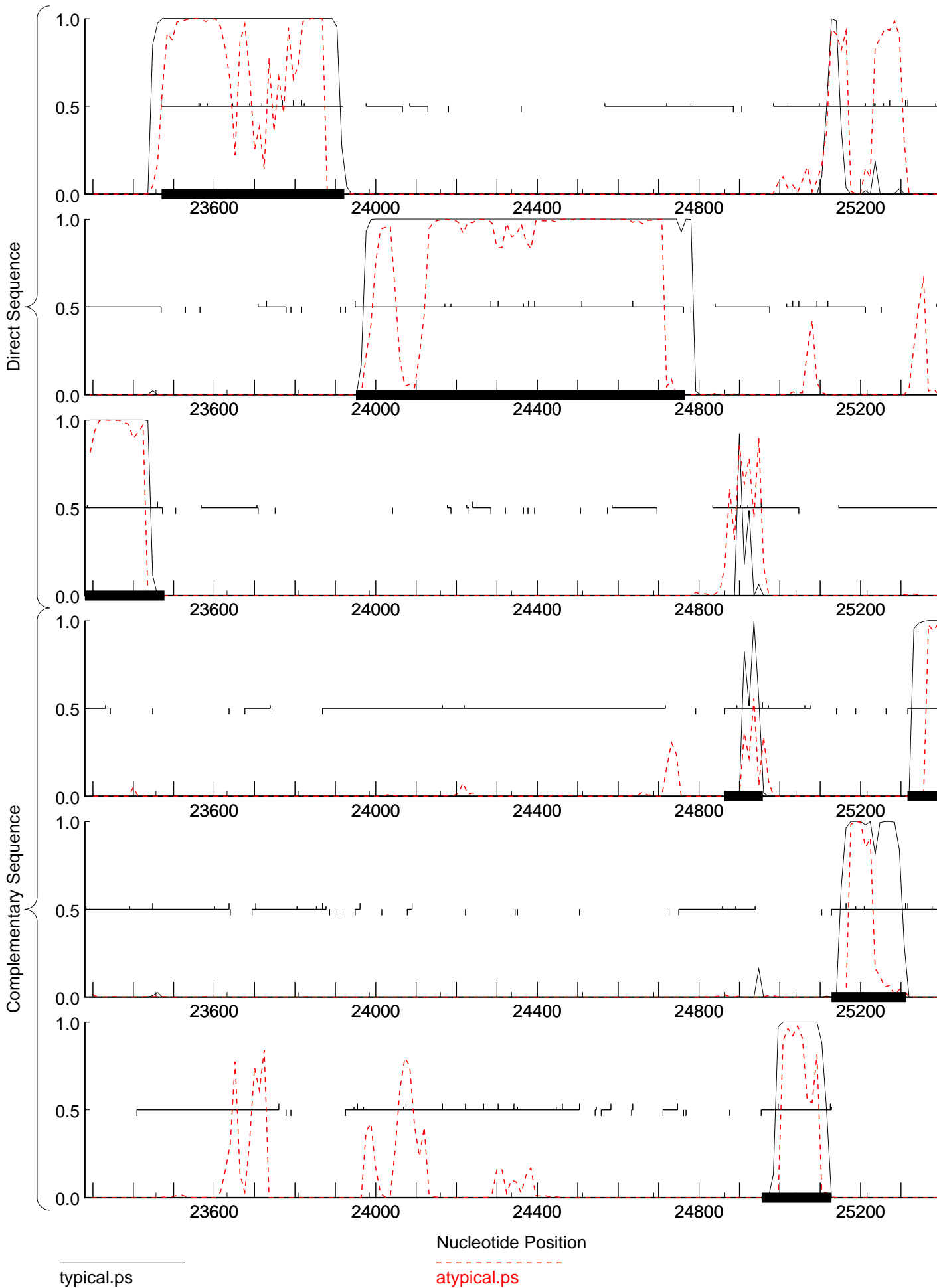
atypical.ps





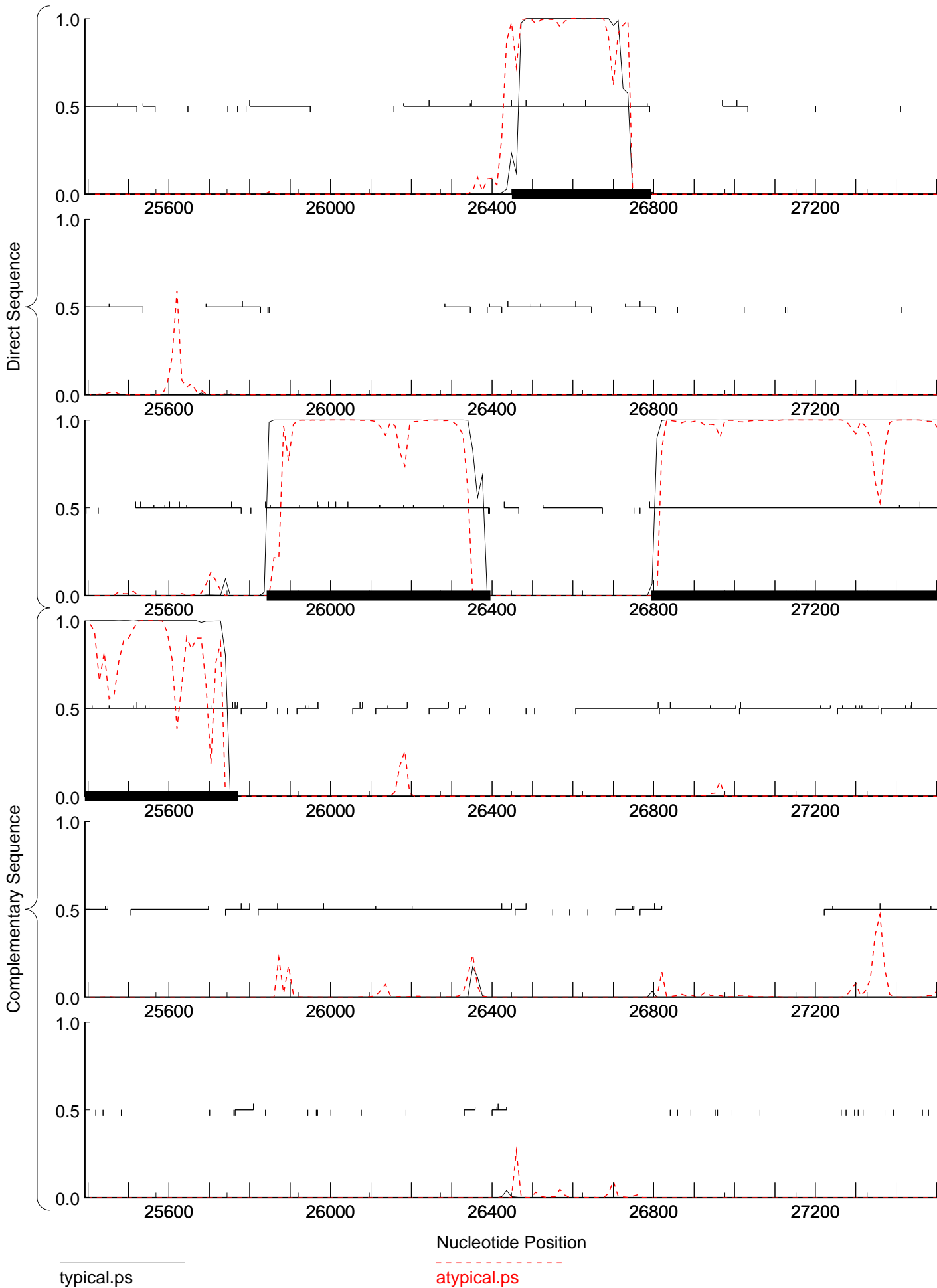
typical.ps

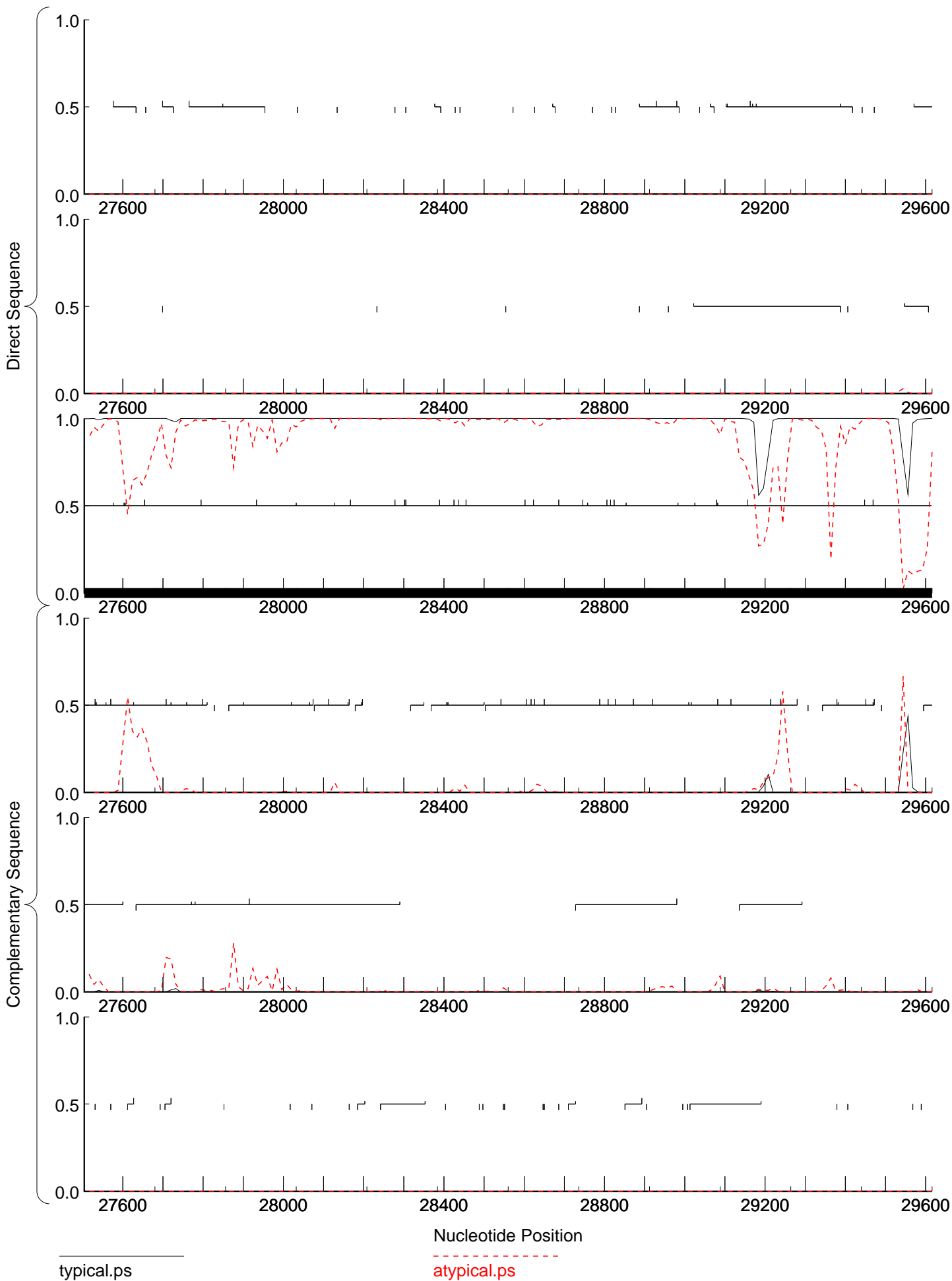
atypical.ps

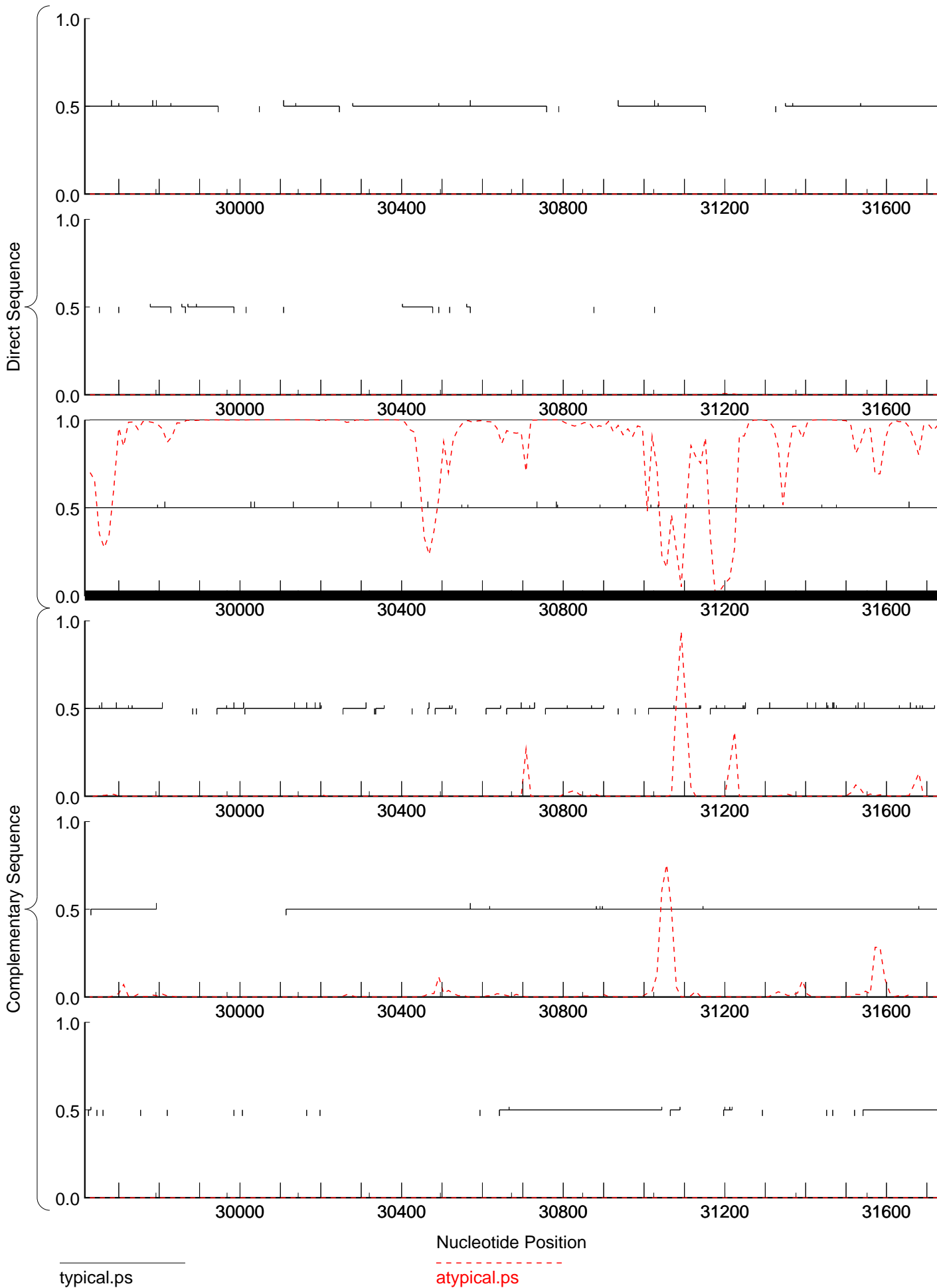


typical.ps

atypical.ps

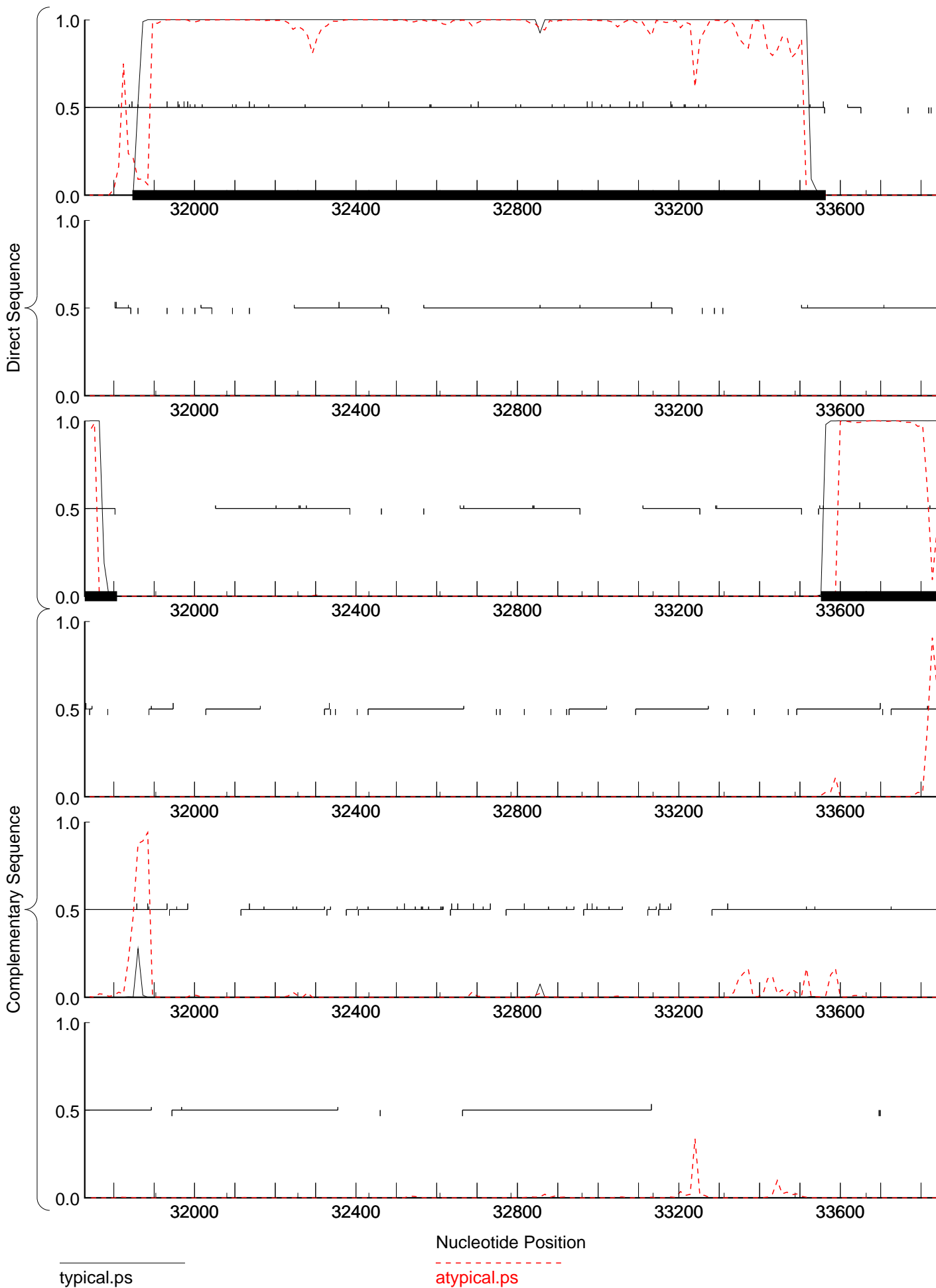






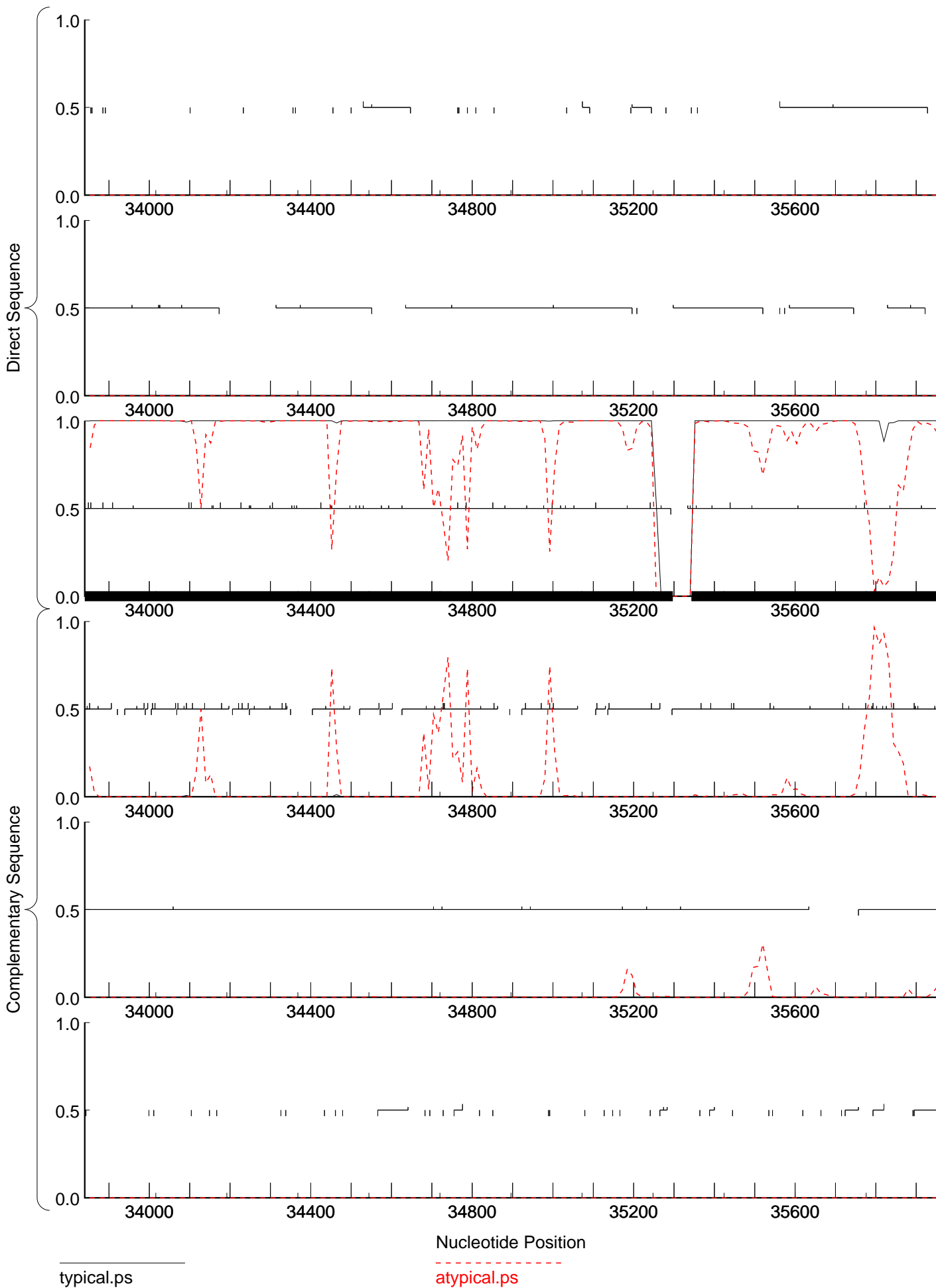
typical.ps

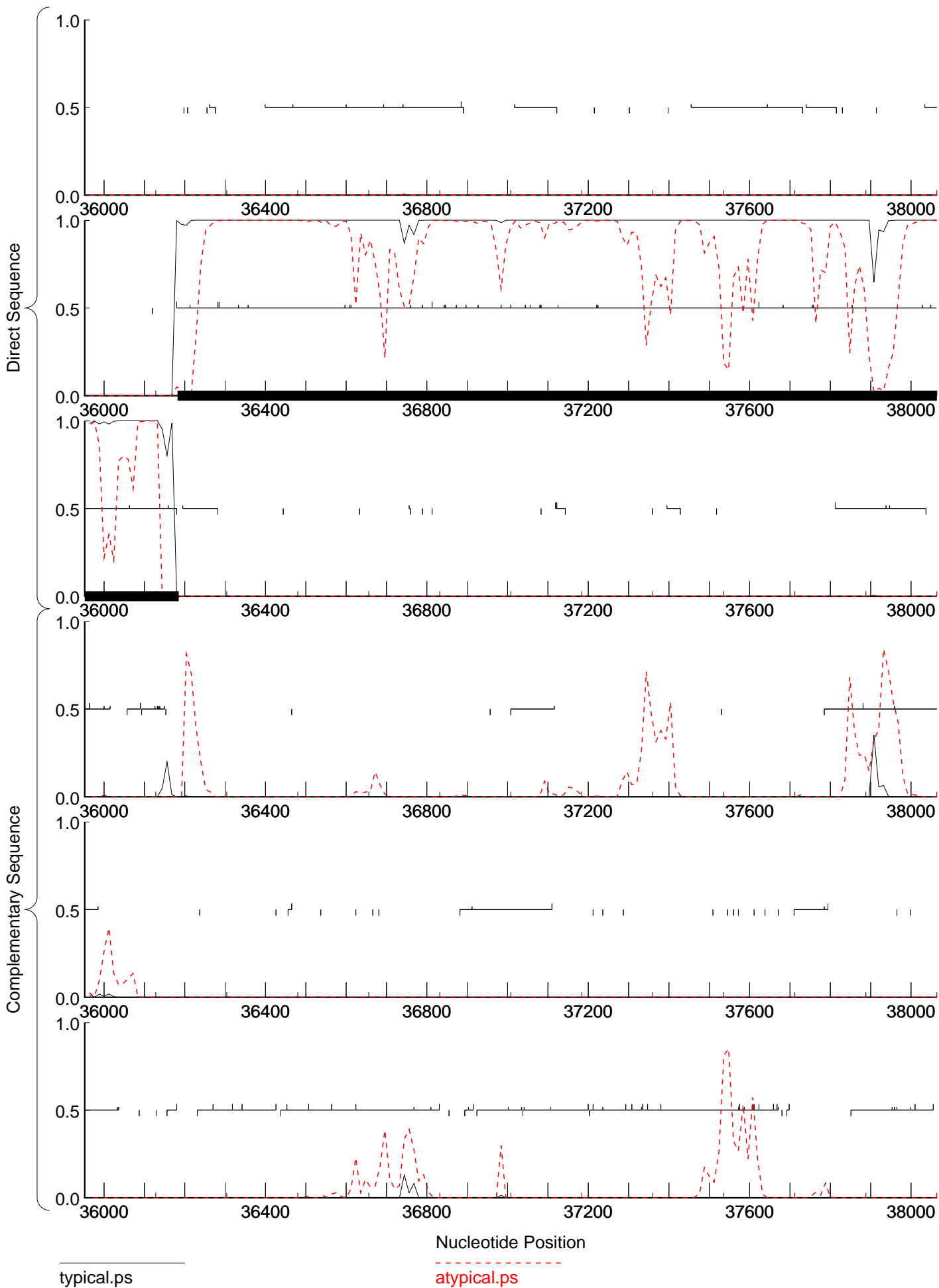
atypical.ps

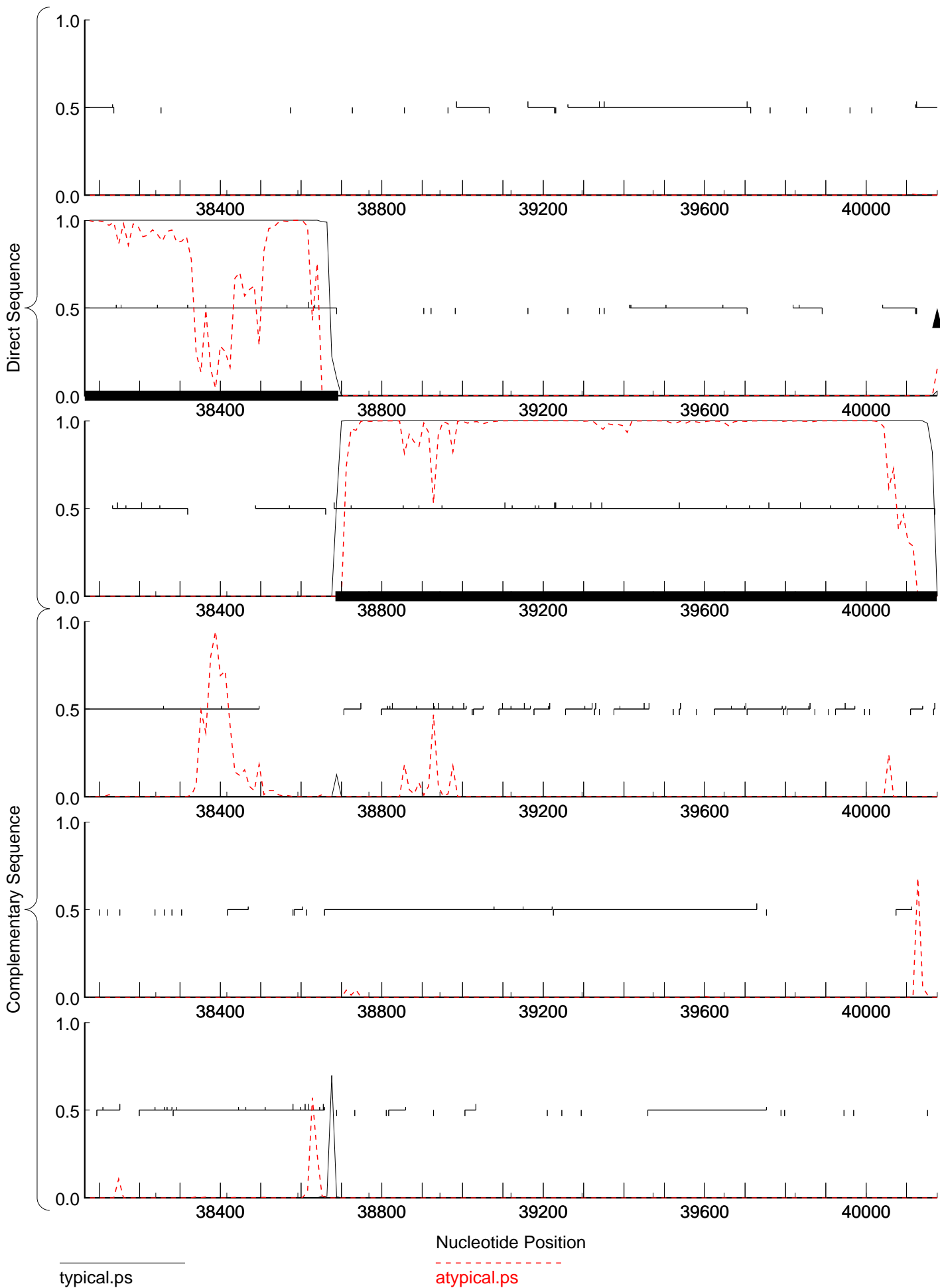


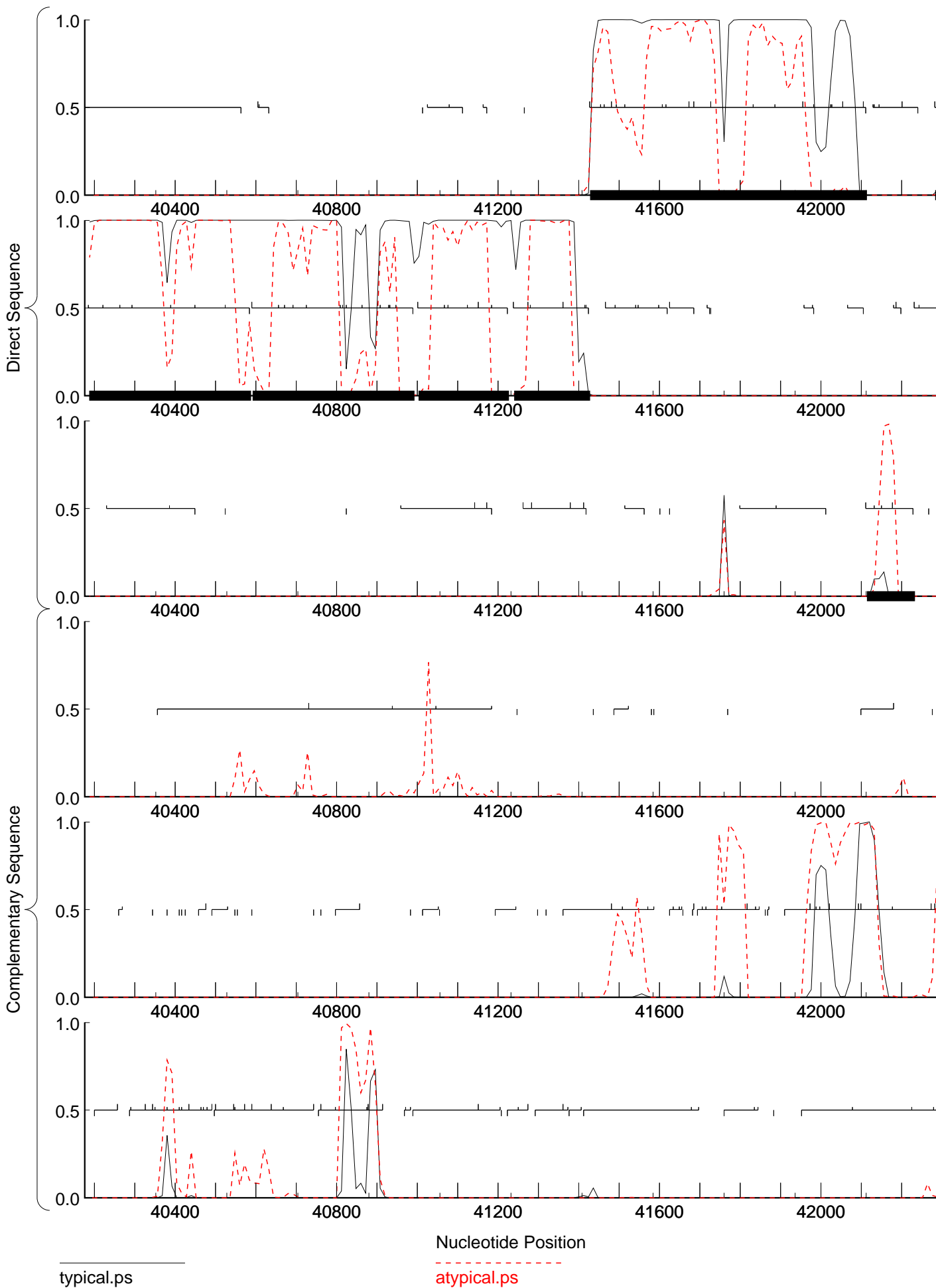
typical.ps

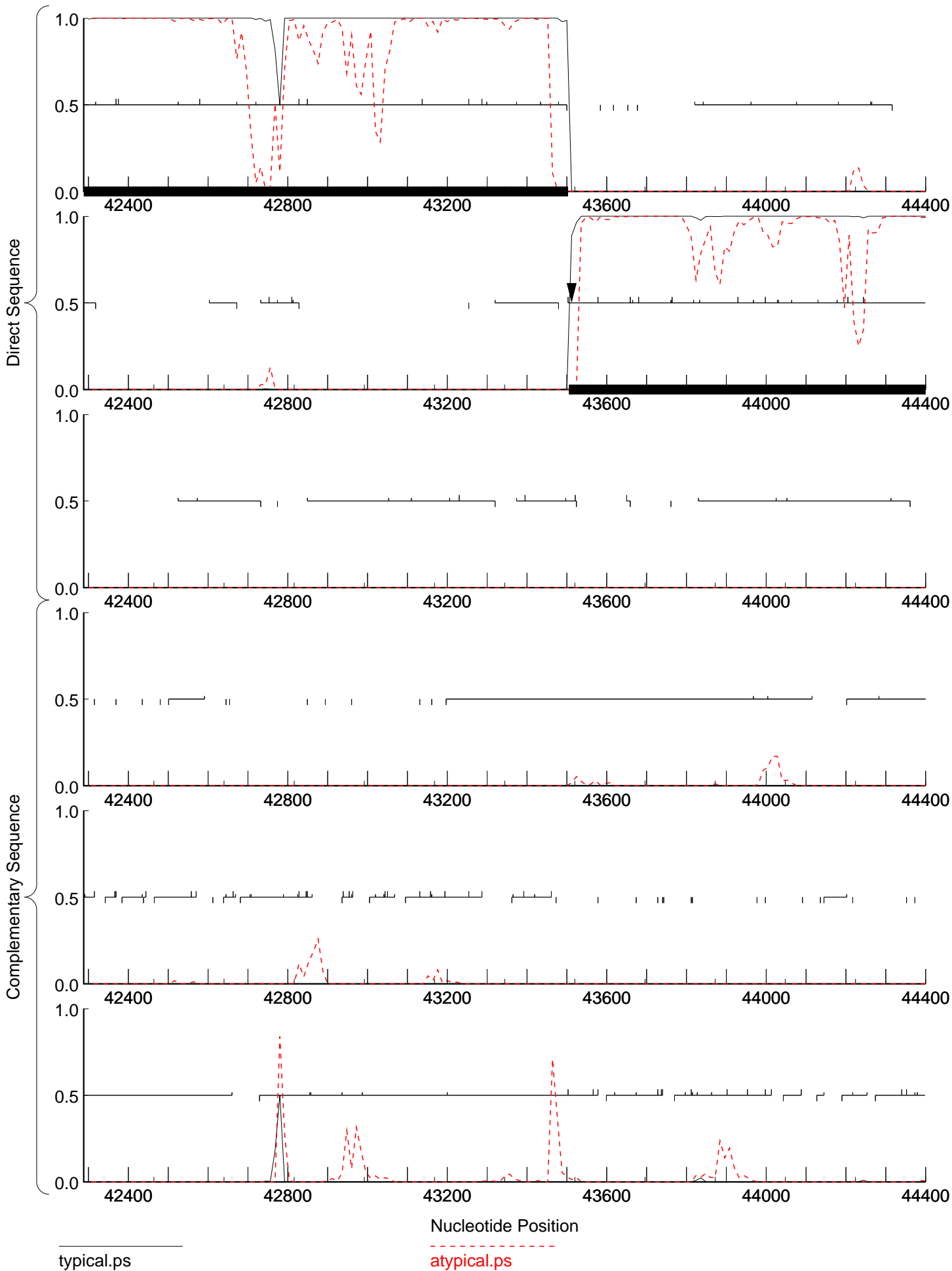
atypical.ps

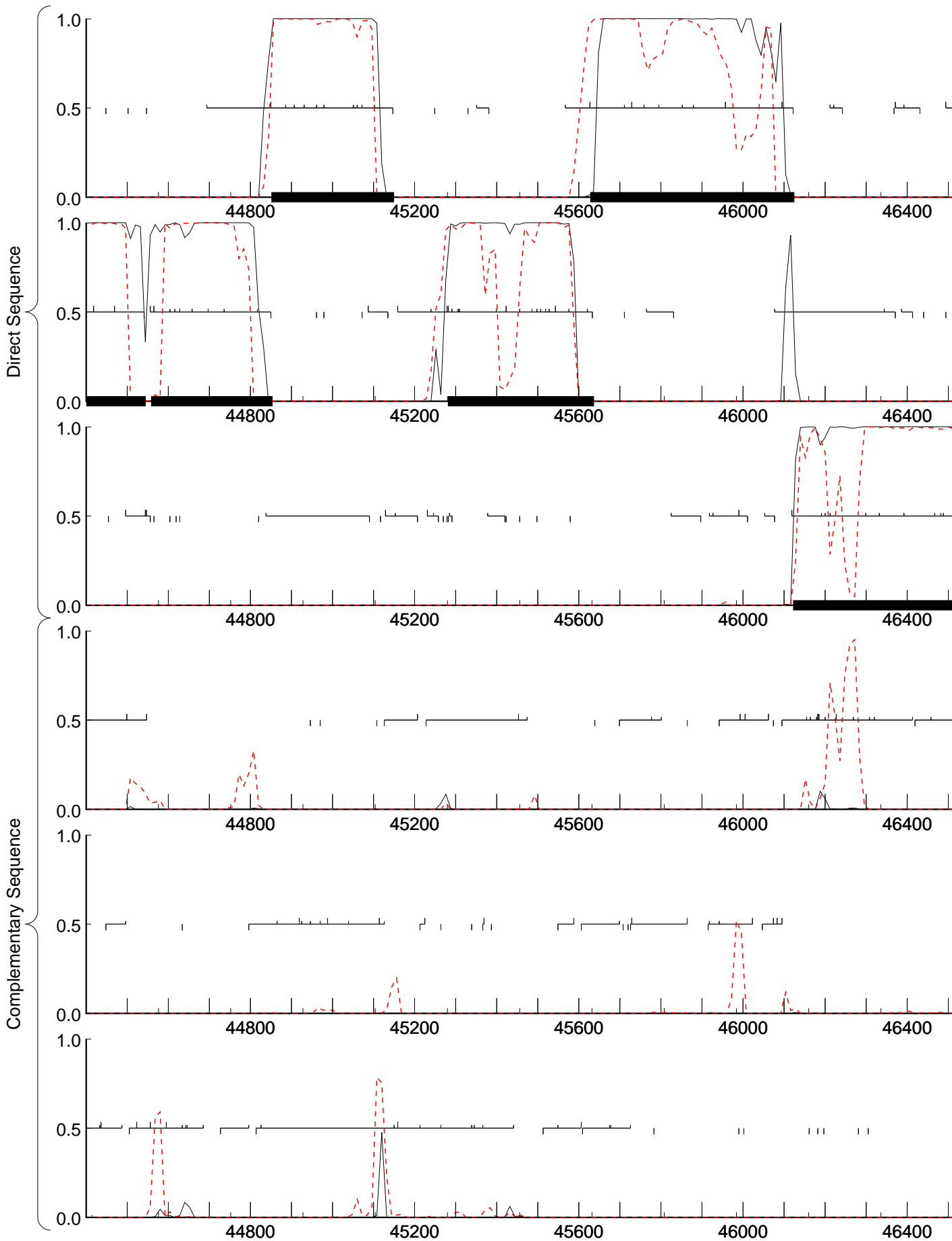


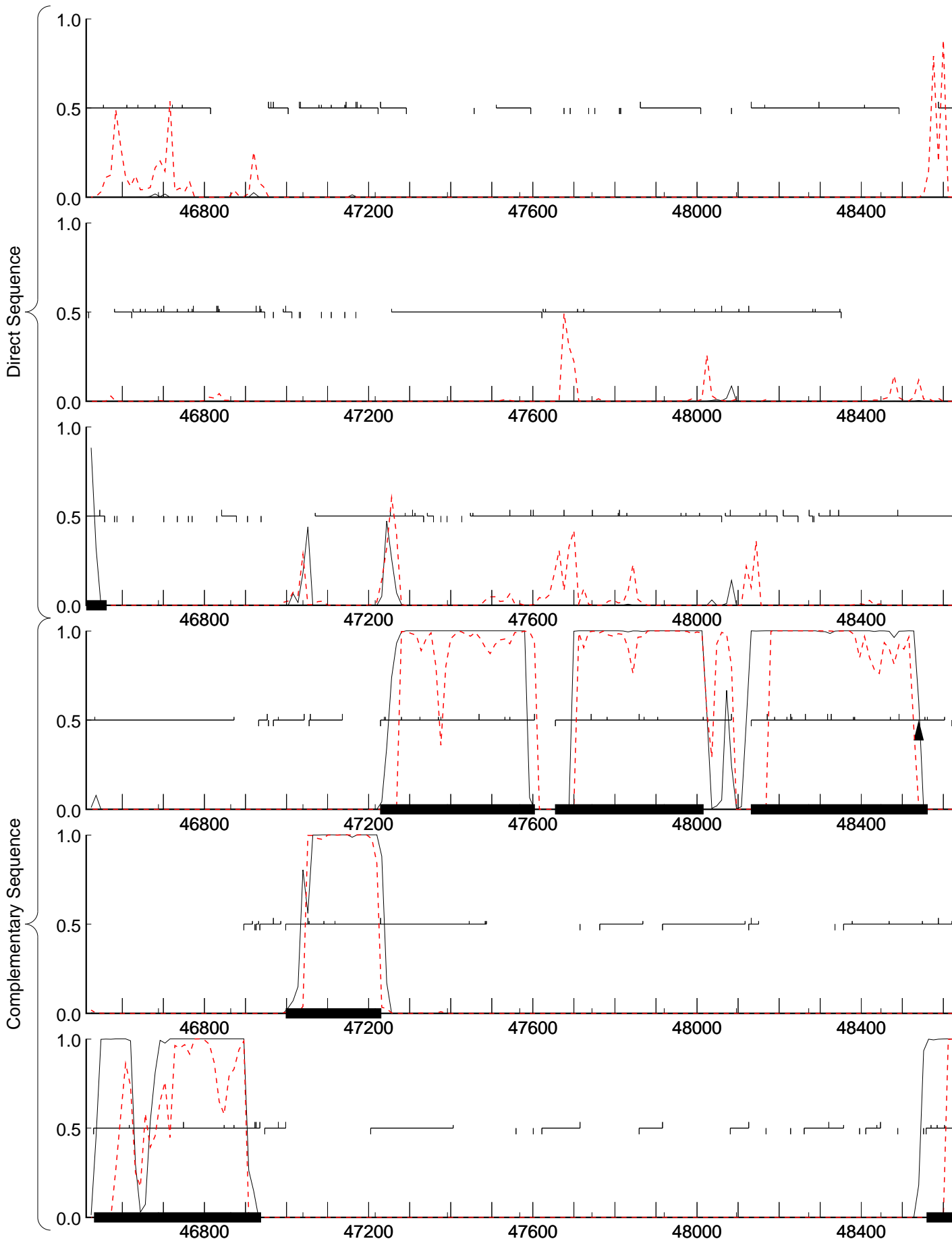






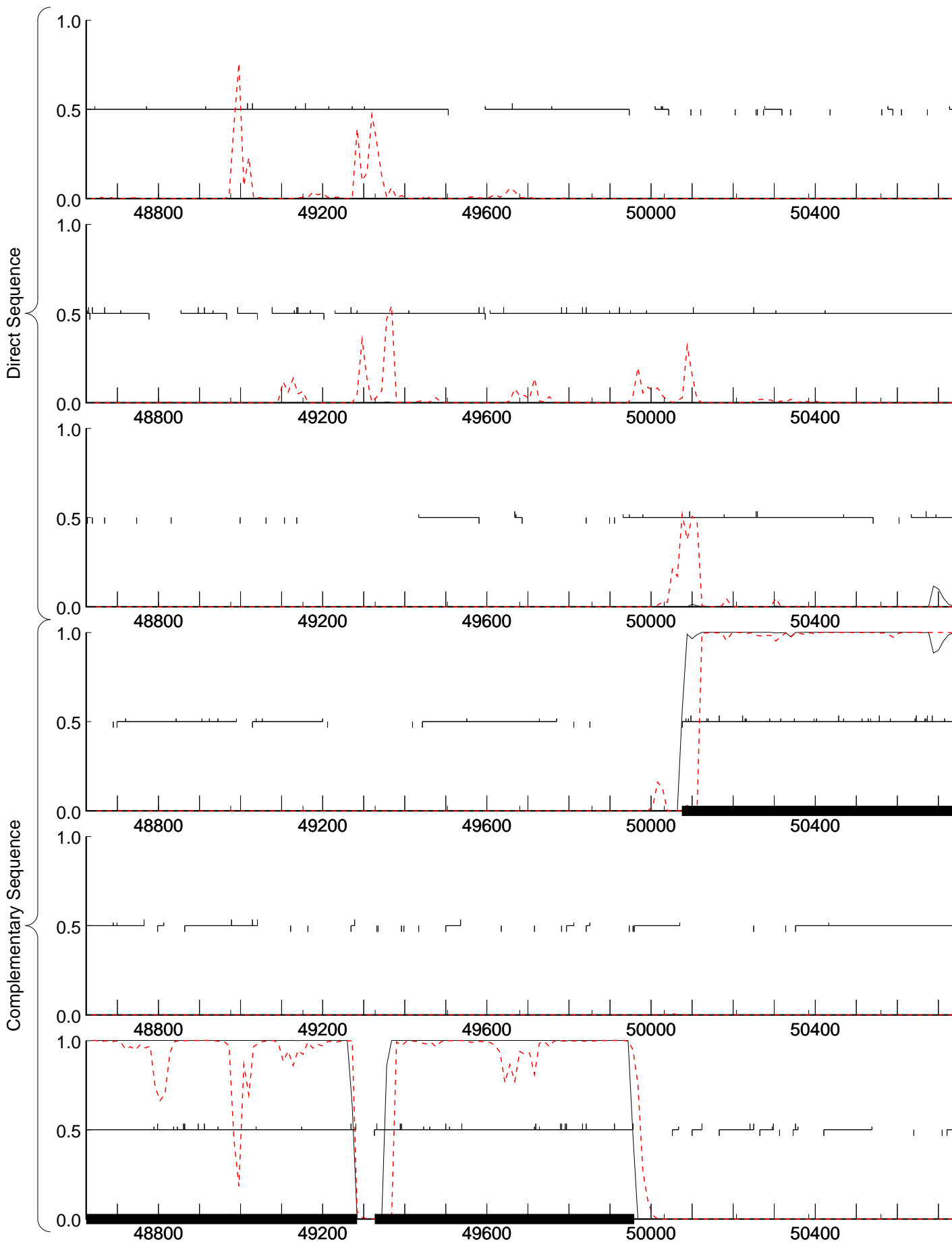






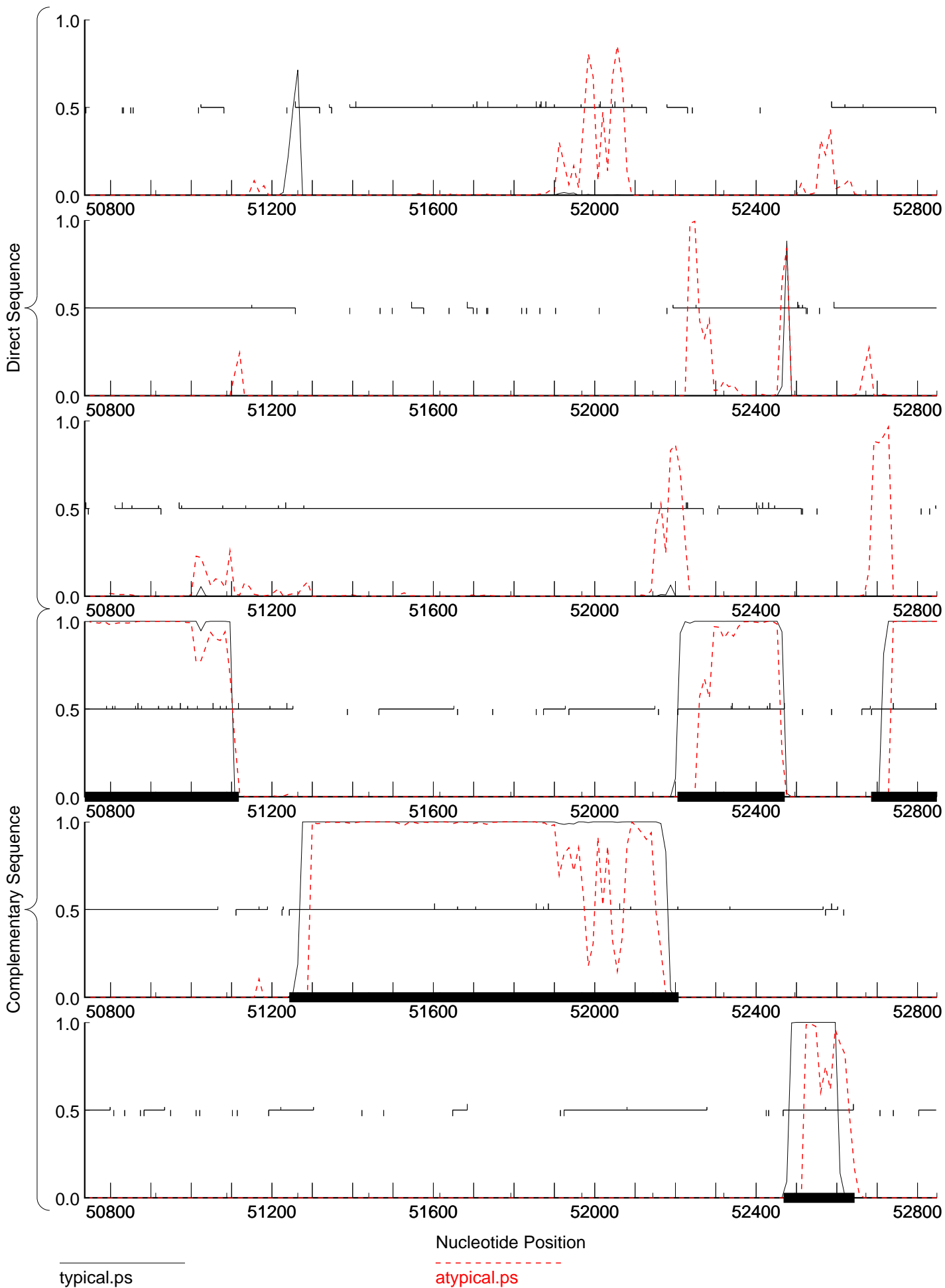
typical.ps

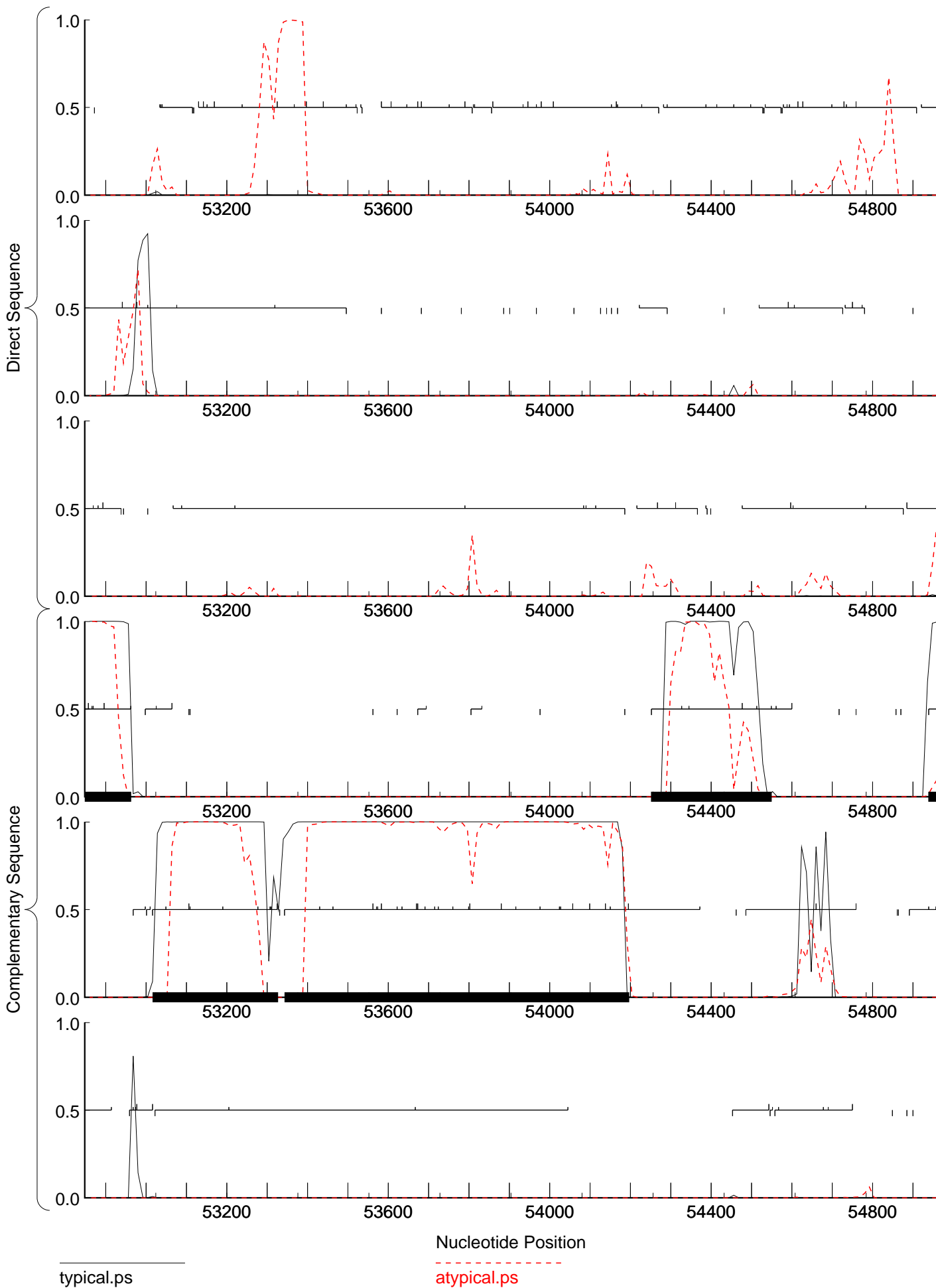
atypical.ps

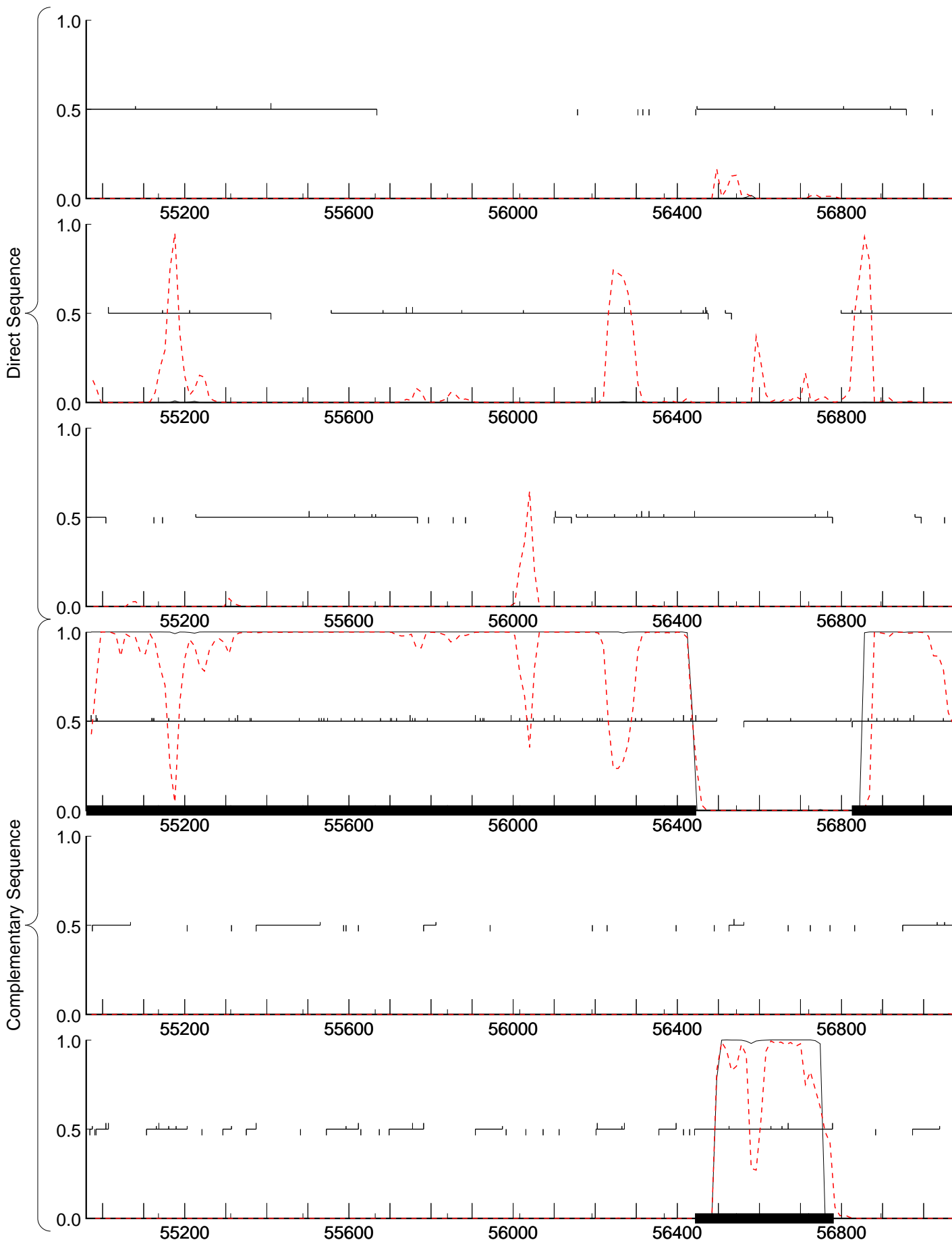


typical.ps

atypical.ps

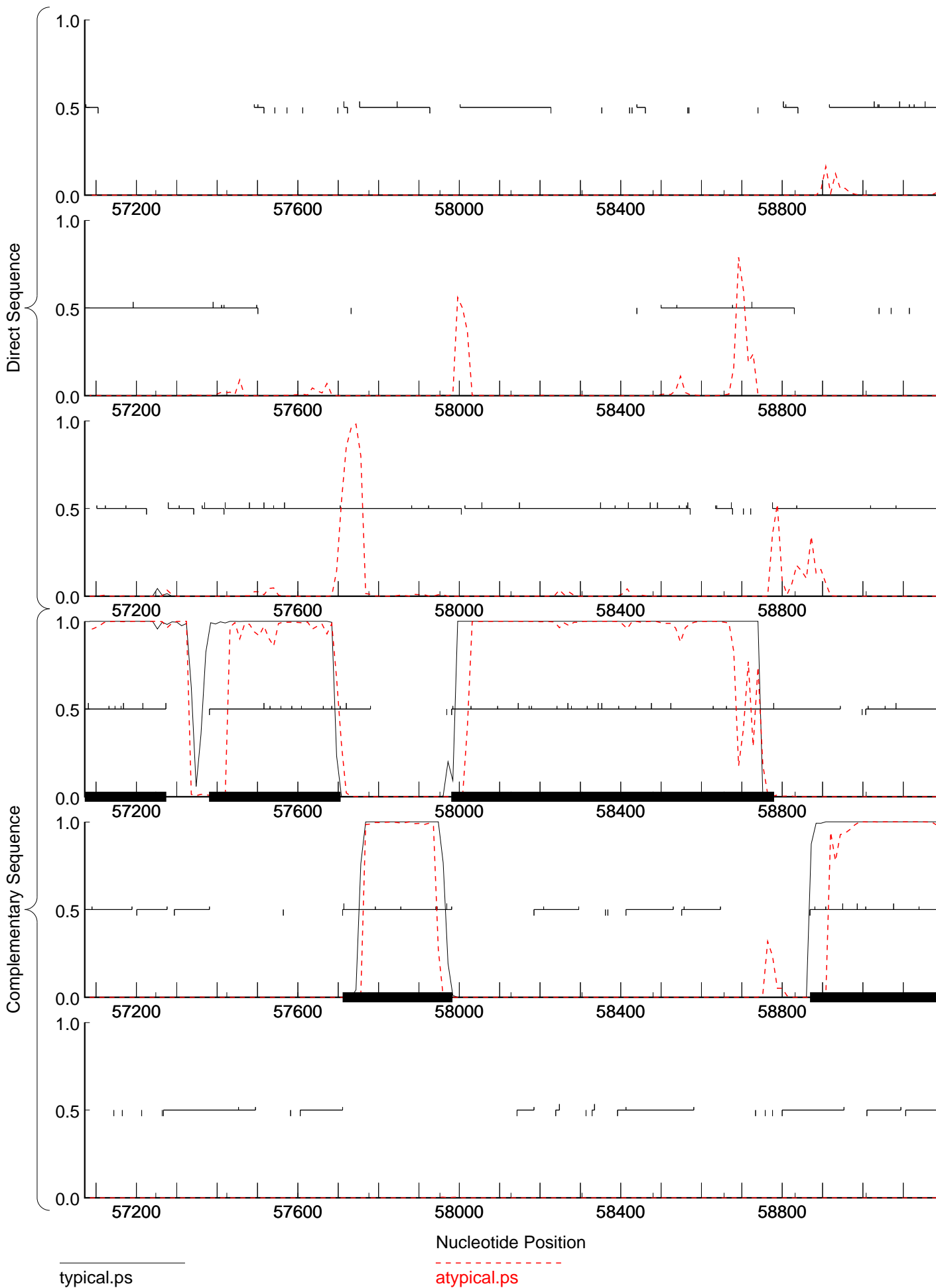






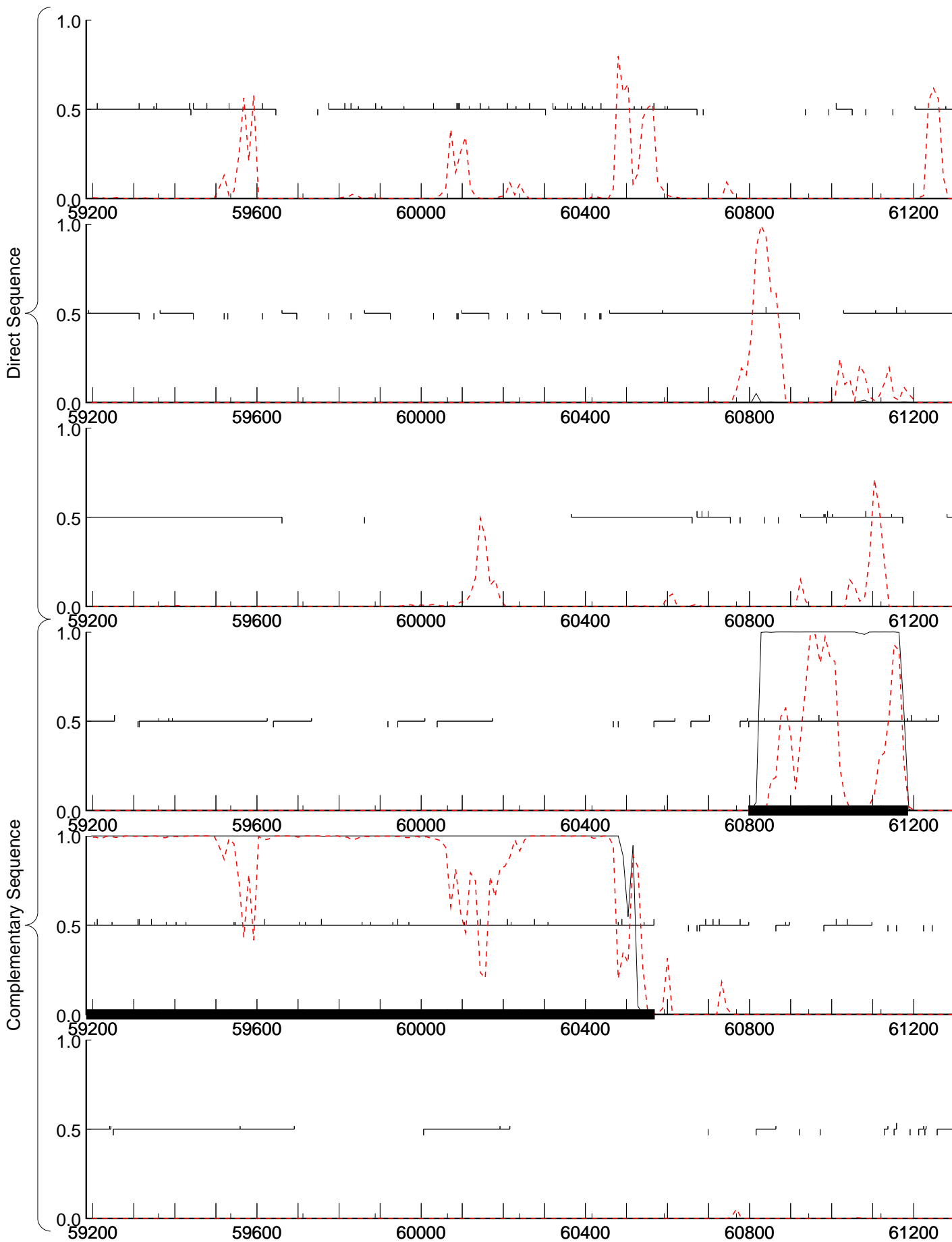
typical.ps

atypical.ps



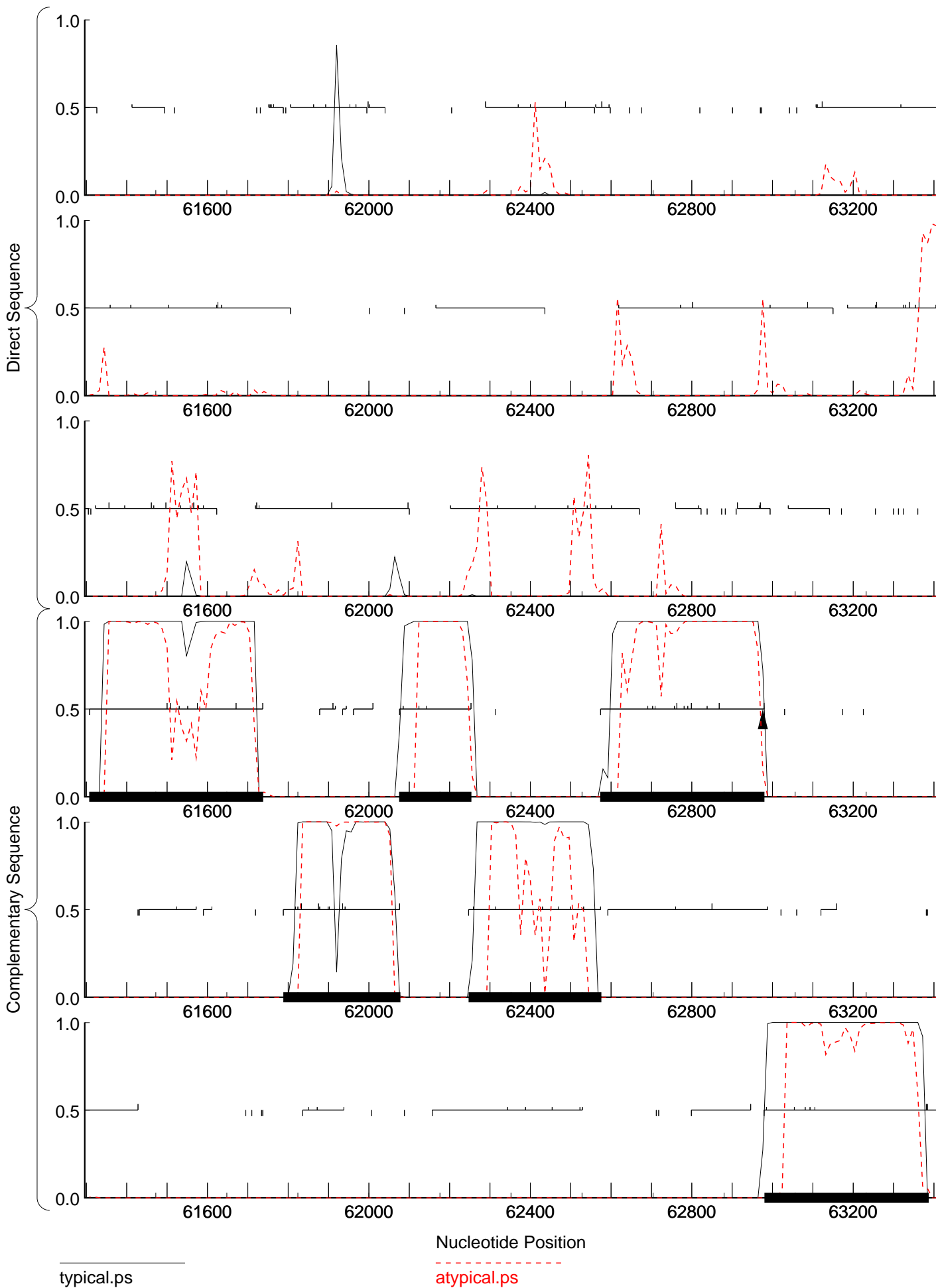
typical.ps

atypical.ps



typical.ps

atypical.ps



typical.ps

atypical.ps

