

GeneMark

Version 2.5p (09.08.06)

Copyright 1993 - M. Borodovsky, J. McIninch

PROGRAM INFORMATION

Sequence : Mycobacterium phage Centaur complete sequence, 53001 bp including 10-base 3' overhang (CGGT
Analysis Date : 7/24/17 at 22:29:18
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,

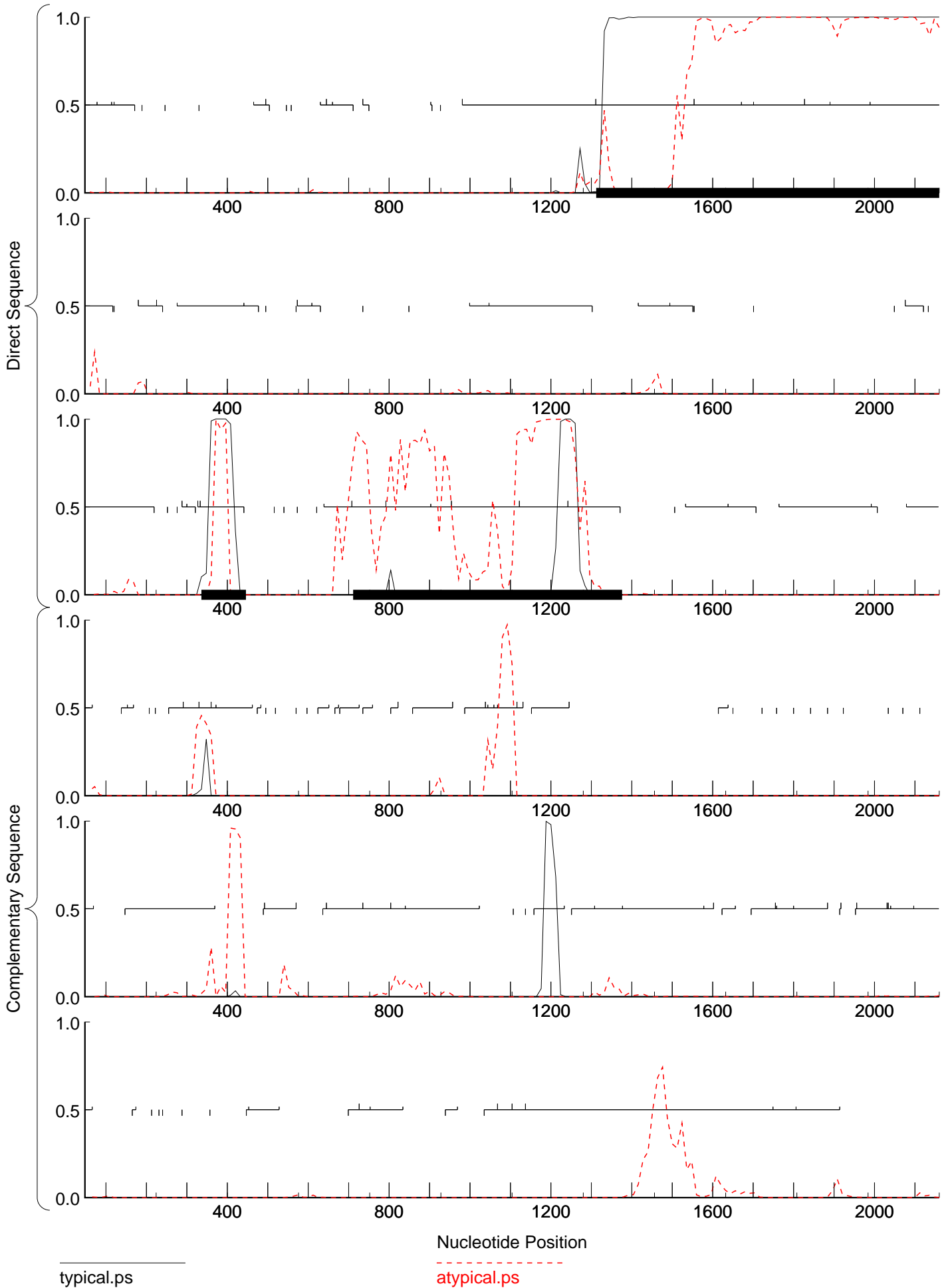
Matrix notes & comments

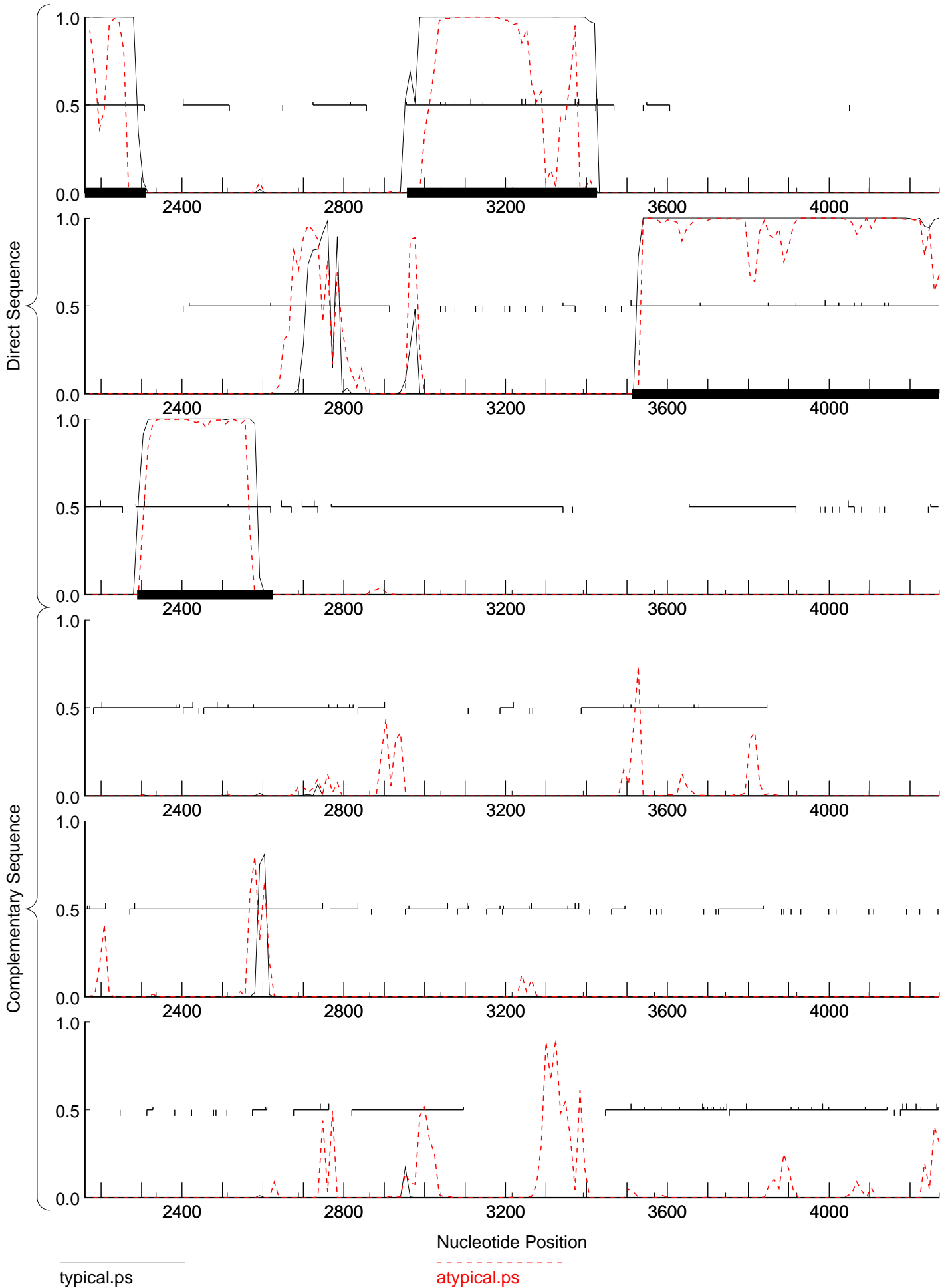
Training set derived by GeneMarkS, 4.30 October 2014
Mon Jul 24 22:29:18 2017

MATRIX INFORMATION

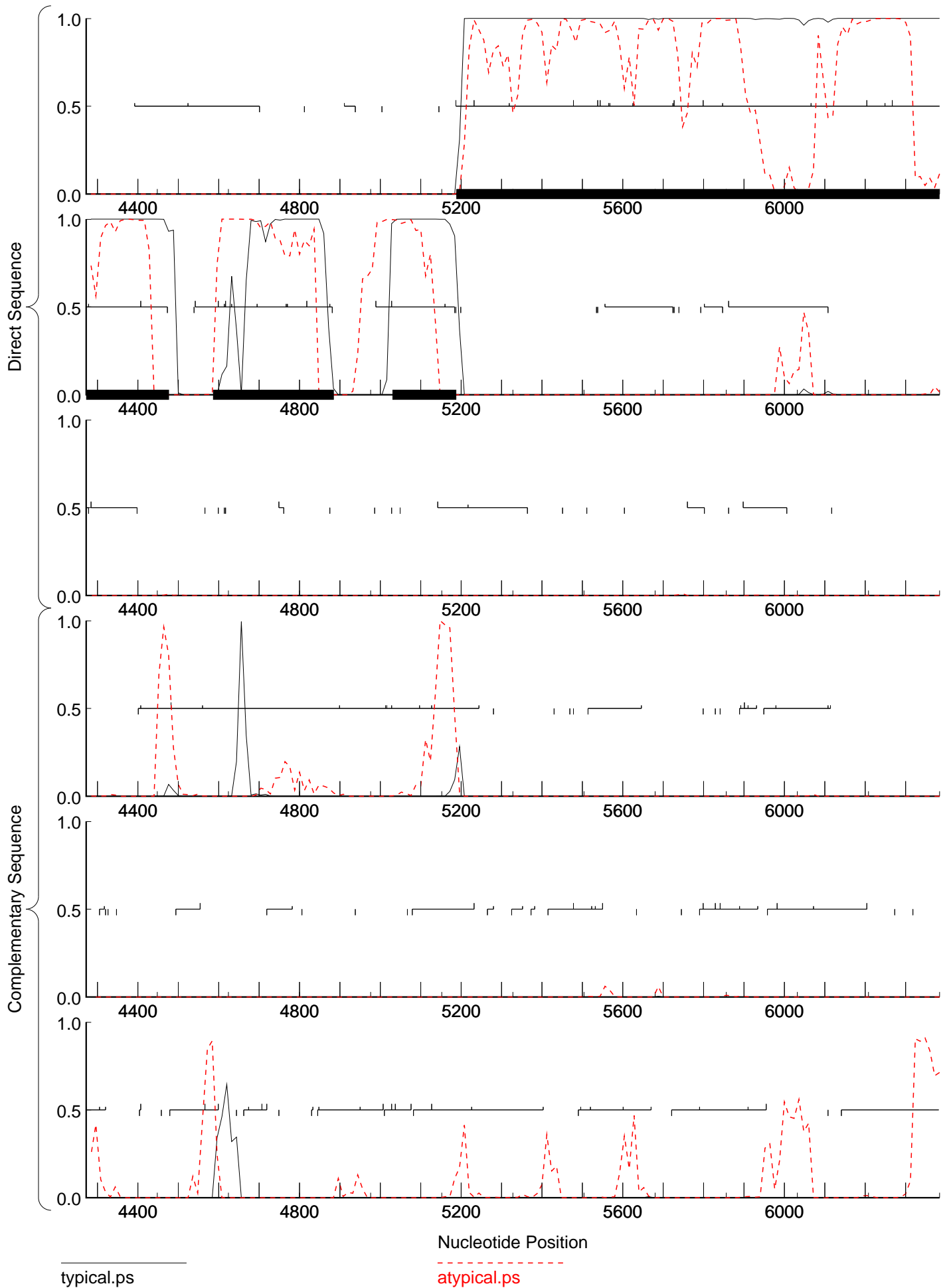
Matrix : -
Author : -
Order : 4

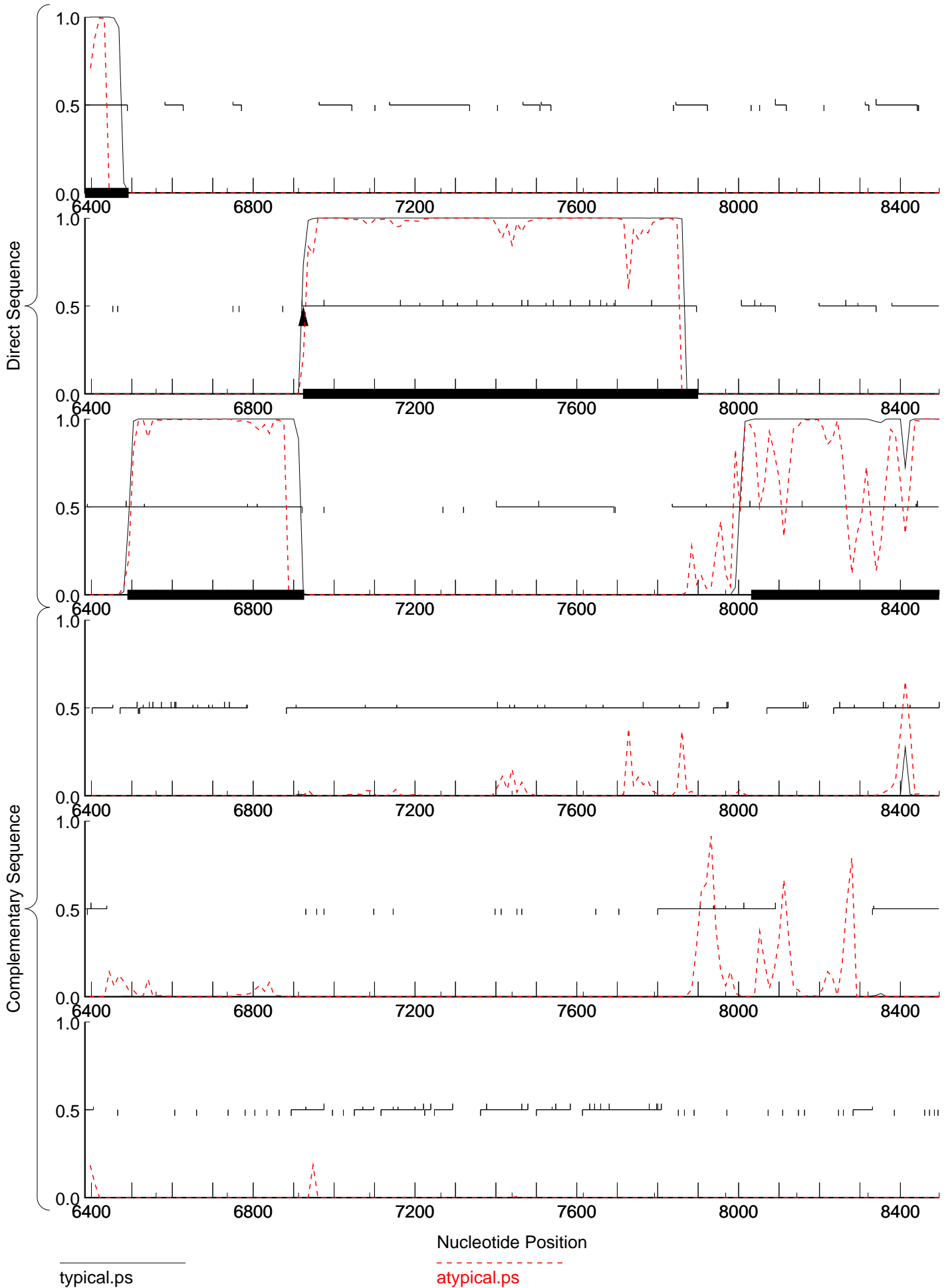
Send questions / comments to:
Dr. M. Borodovsky
Georgia Institute of Technology
School of Biology
Atlanta, GA 30332-0230

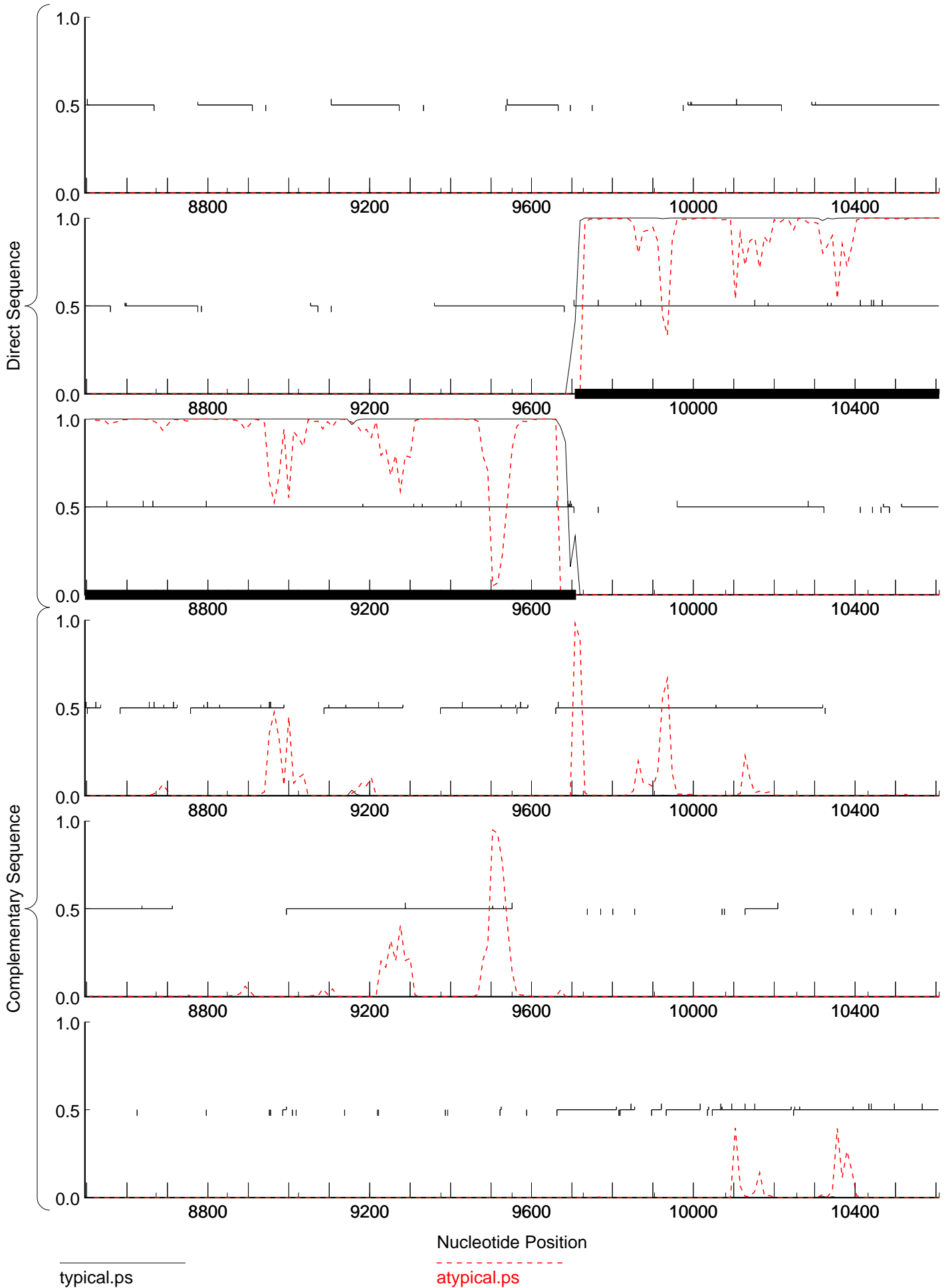


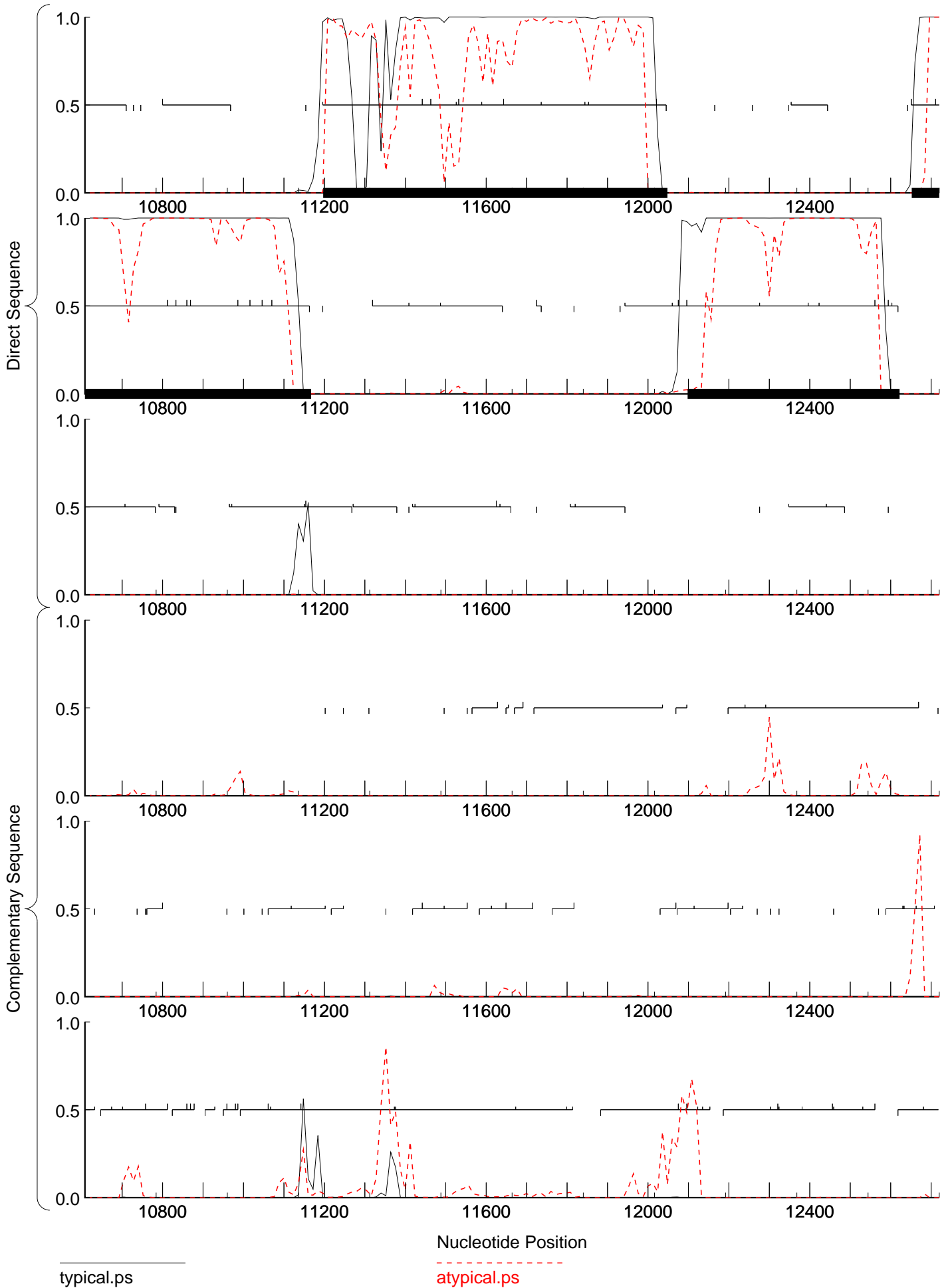


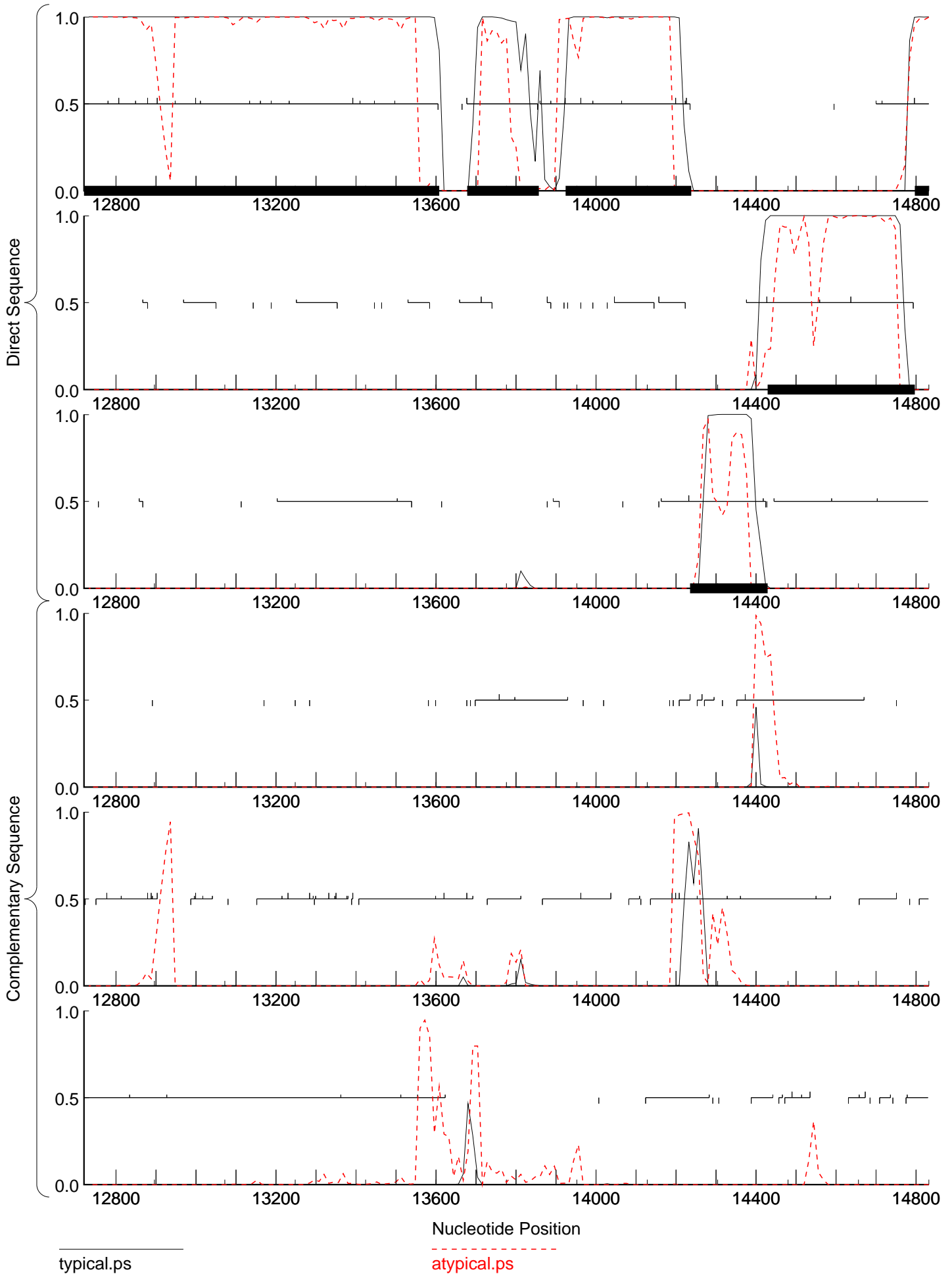
GeneMark.hmm prediction

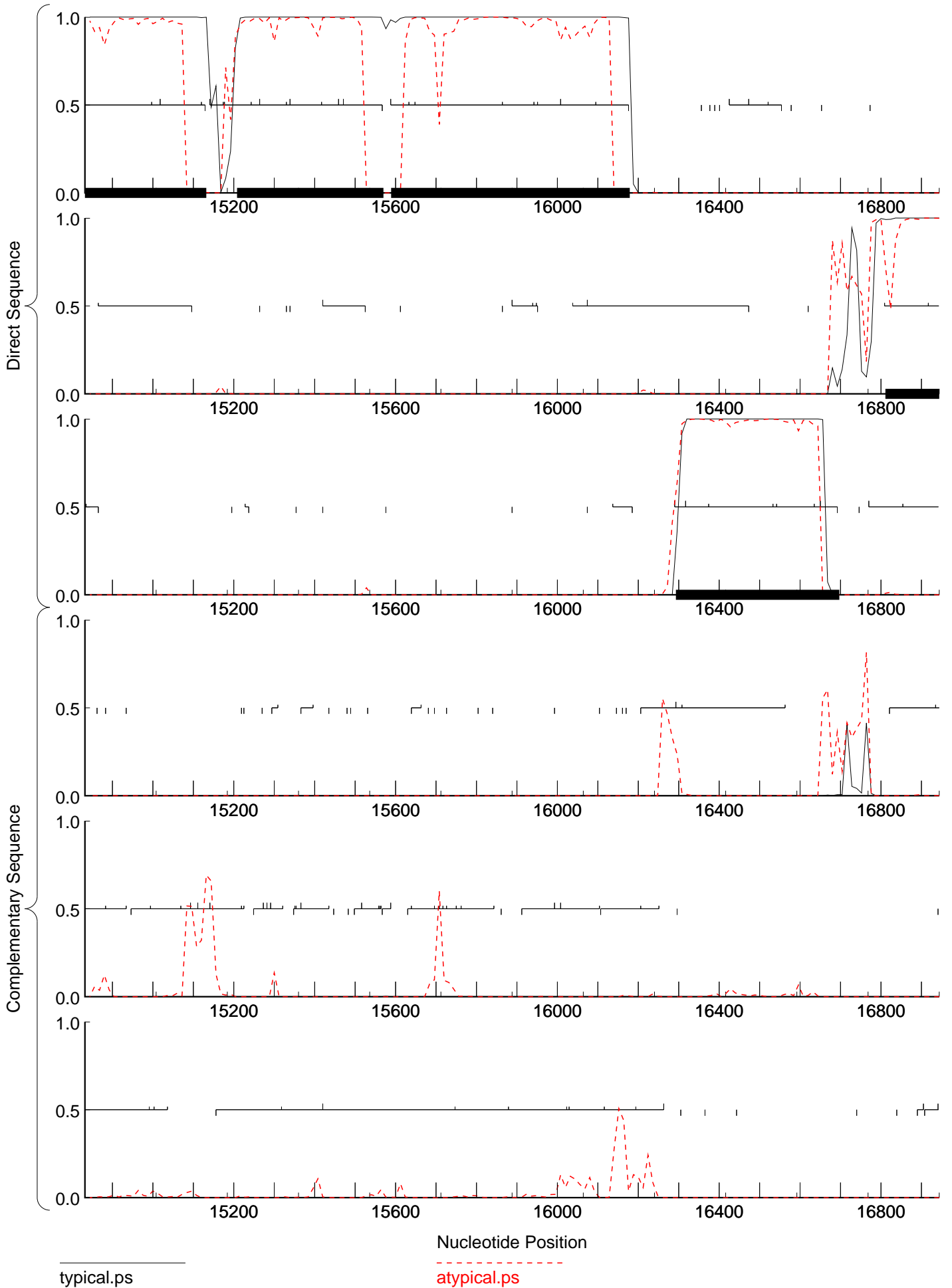






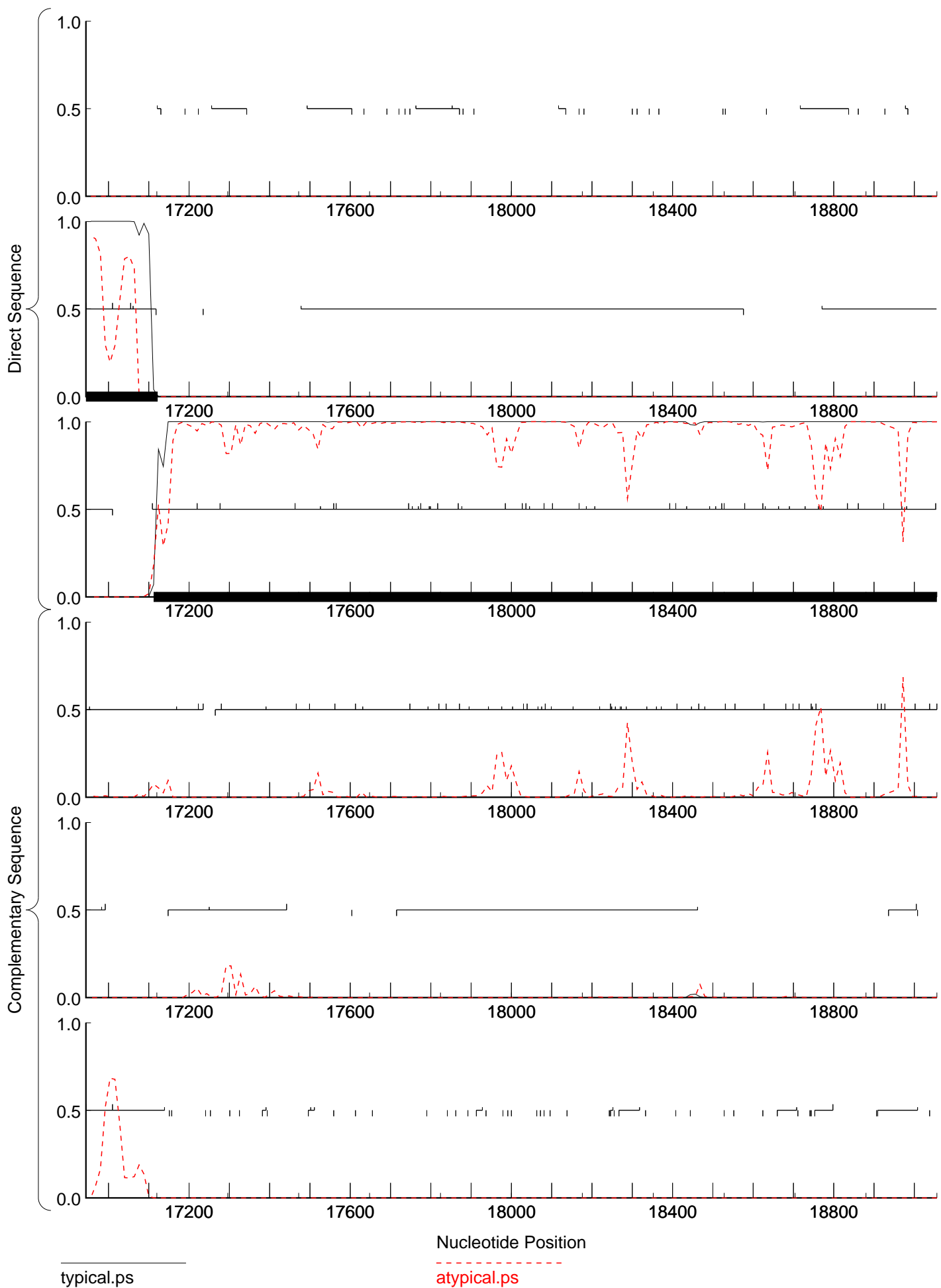


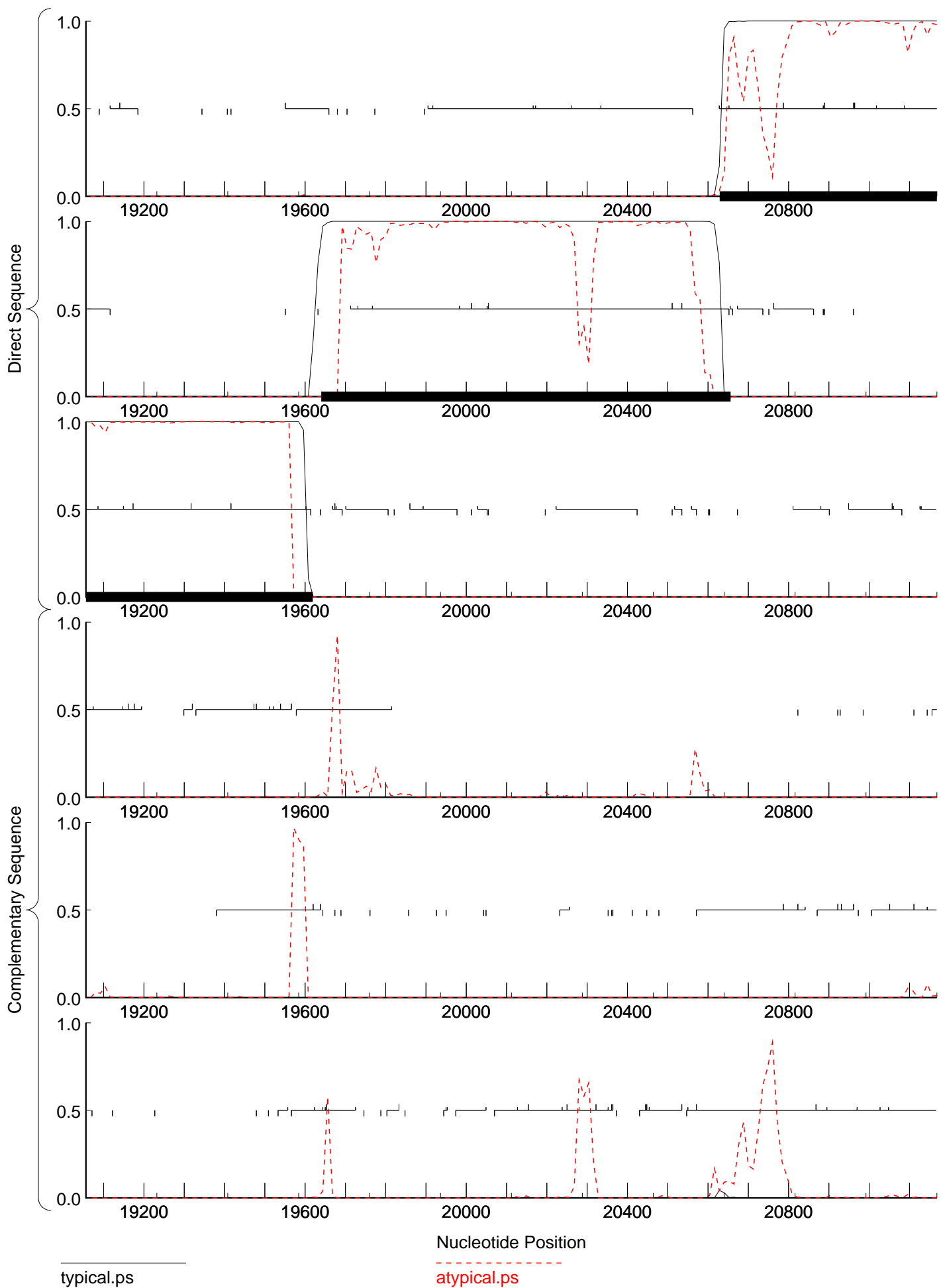


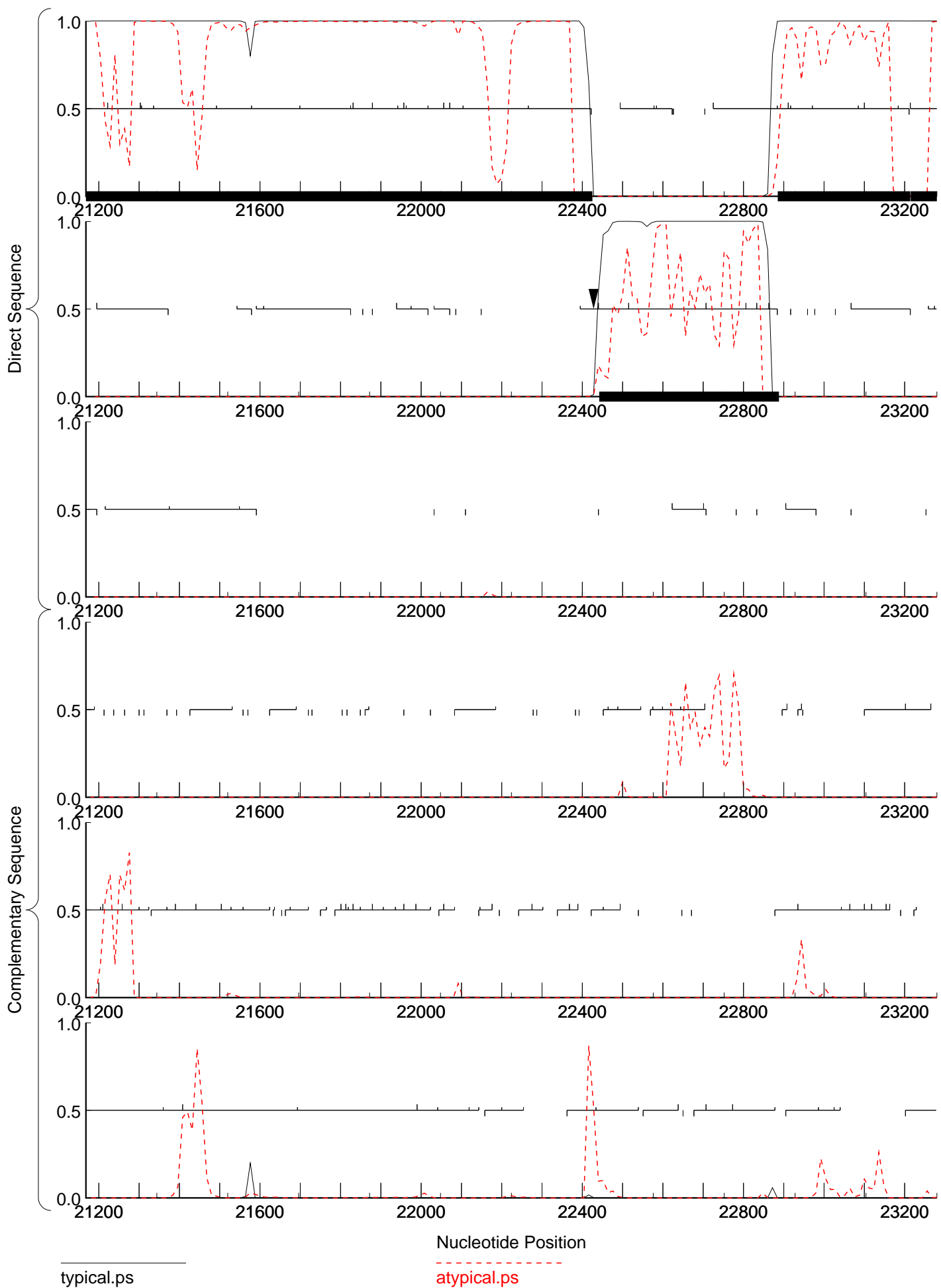


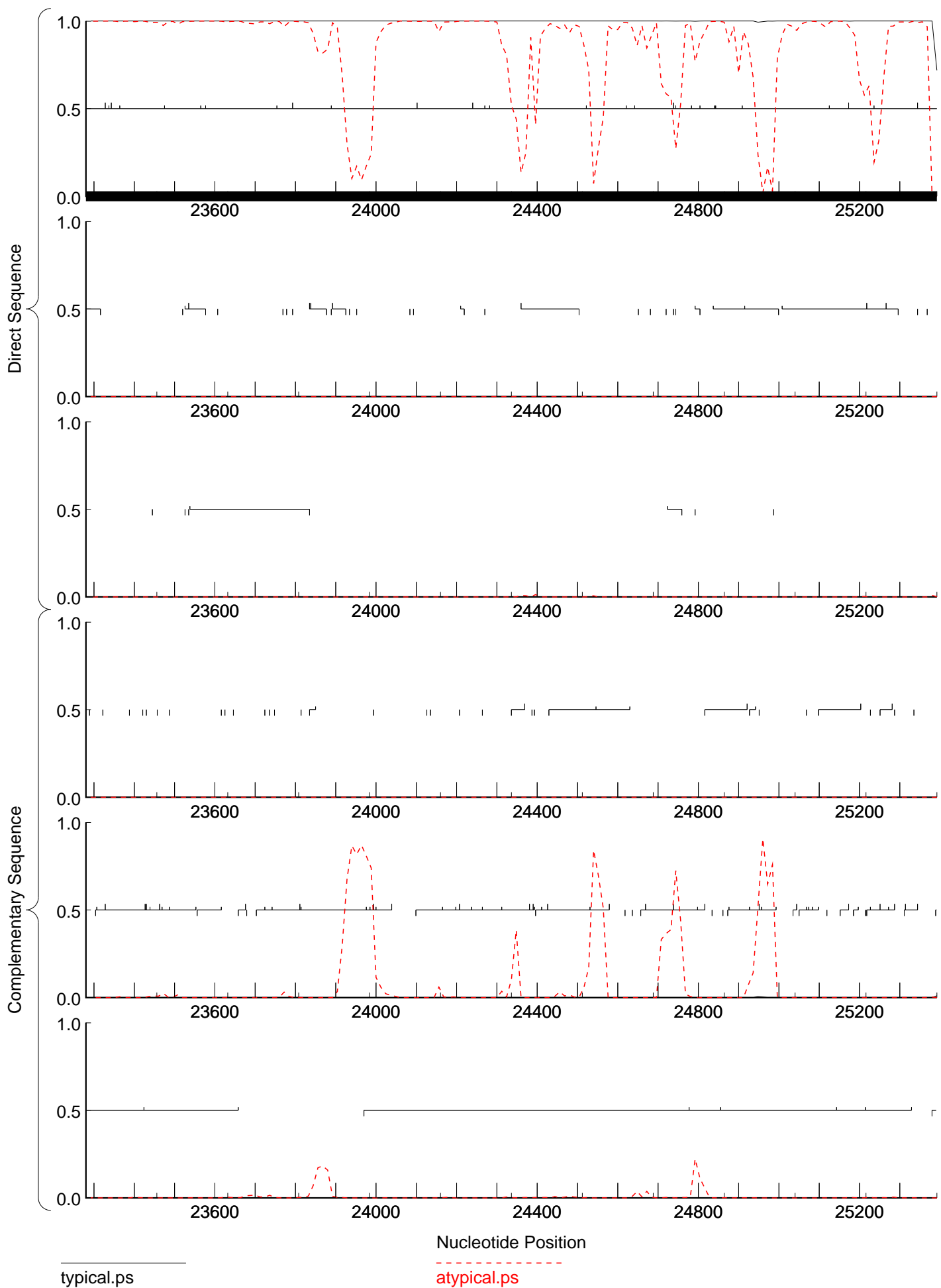
typical.ps

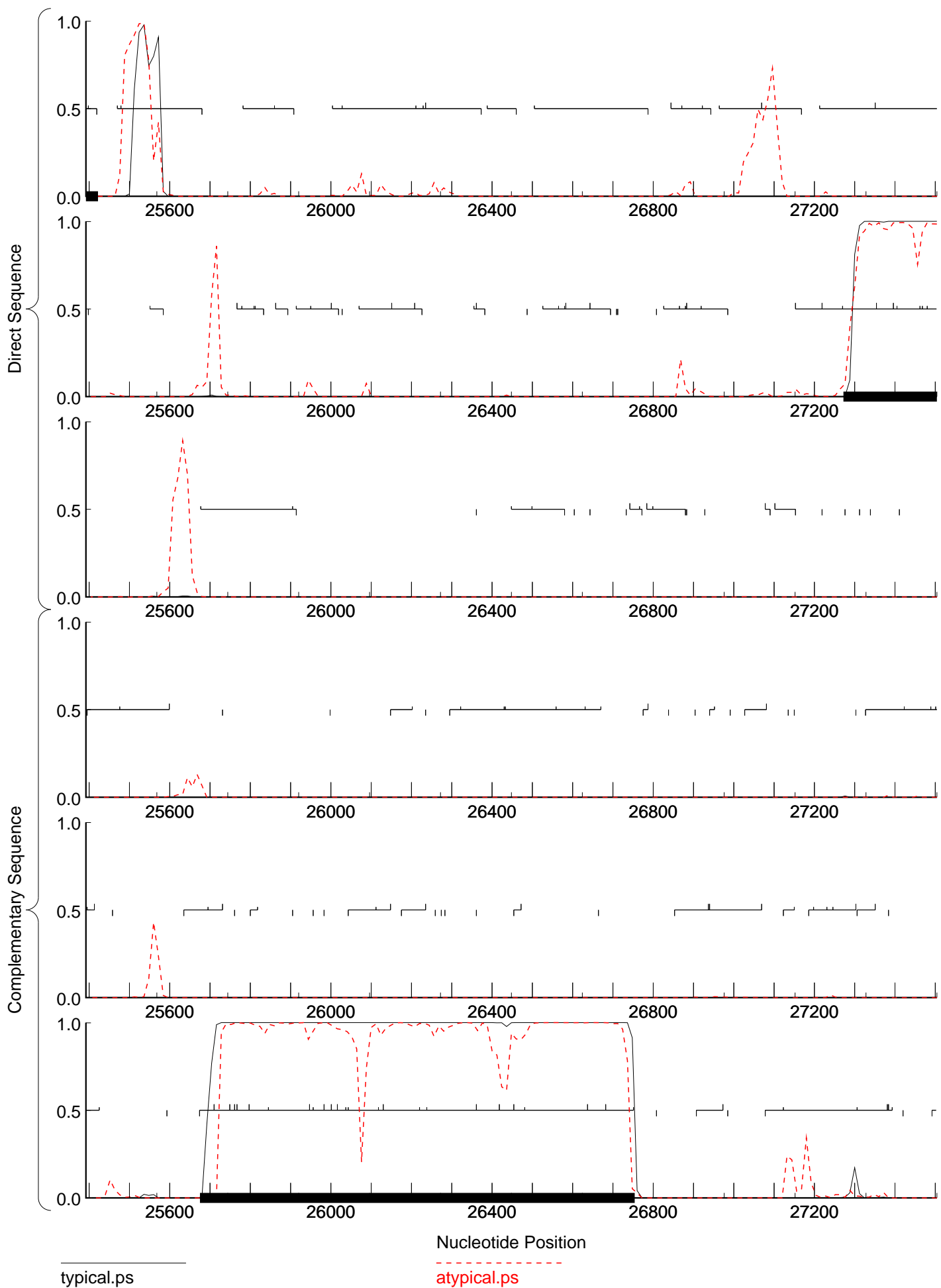
atypical.ps

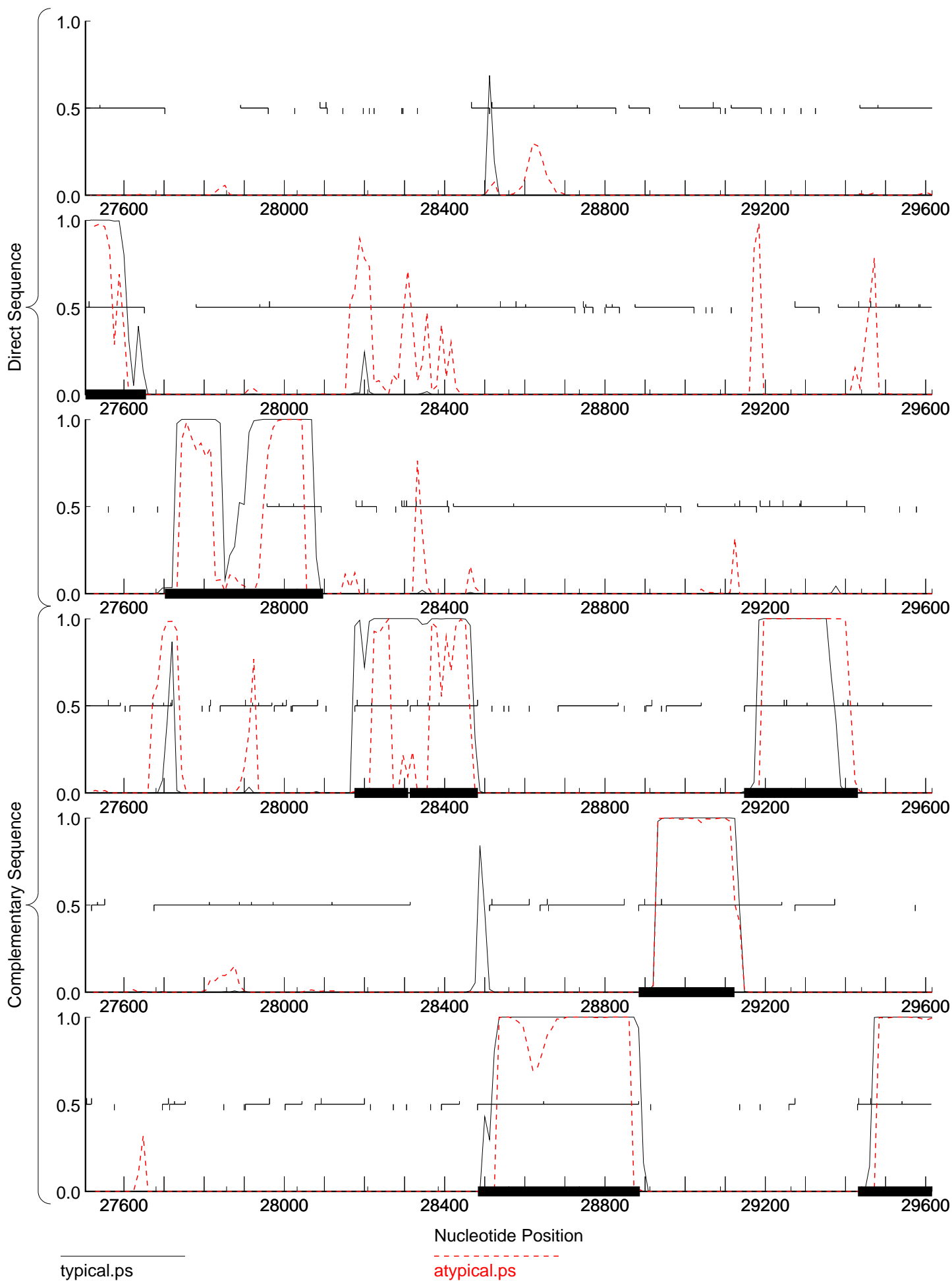


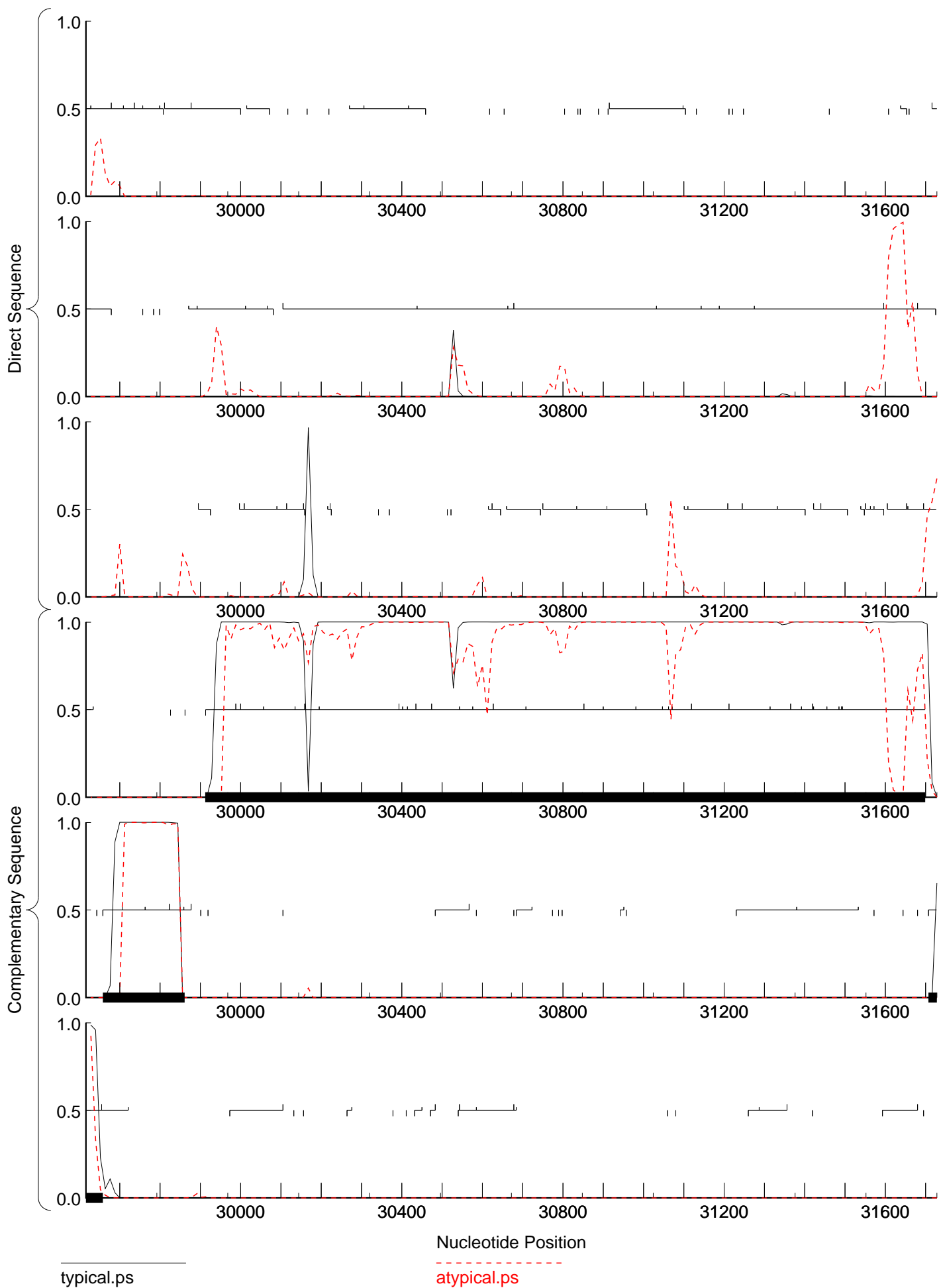


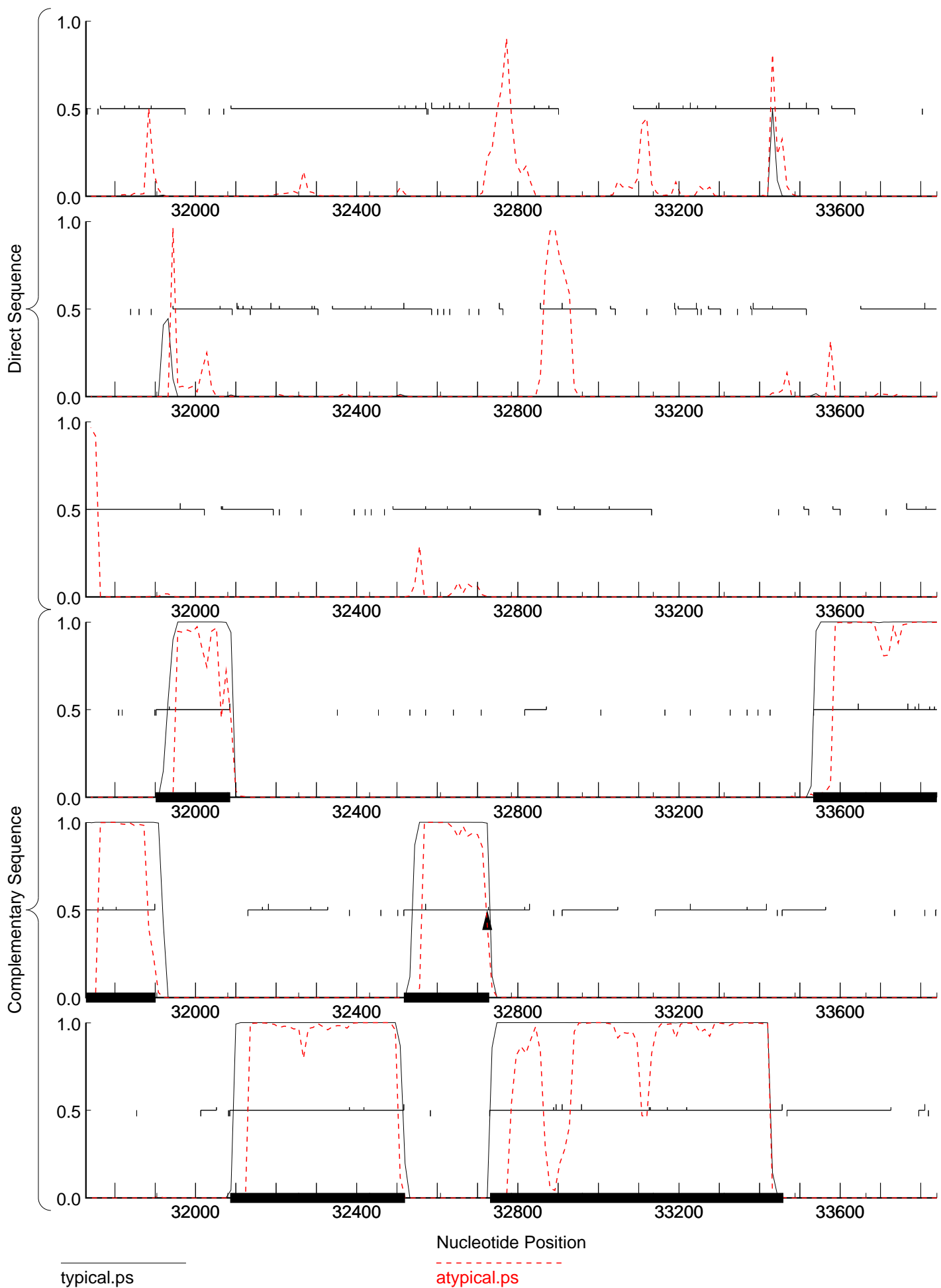






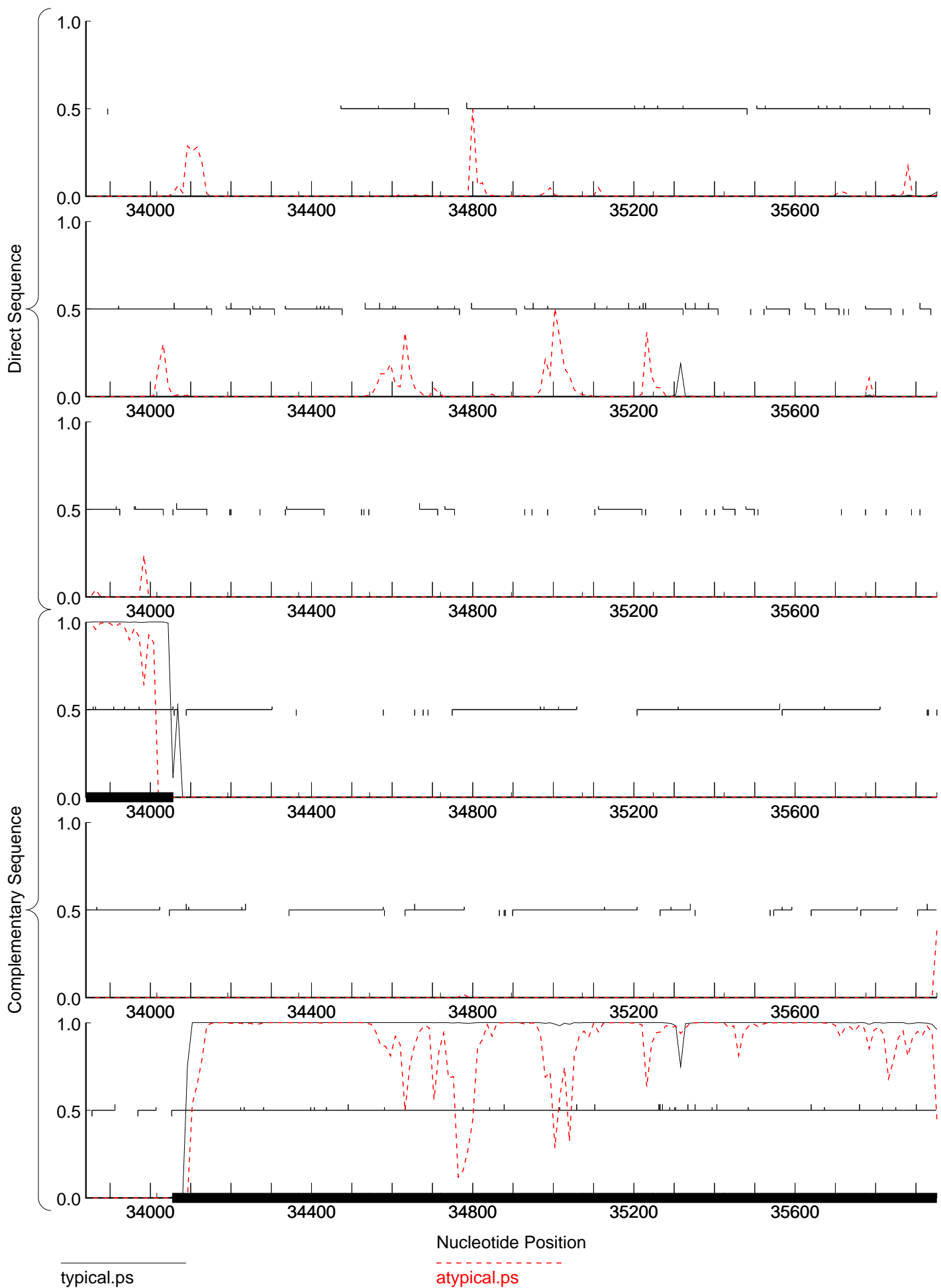






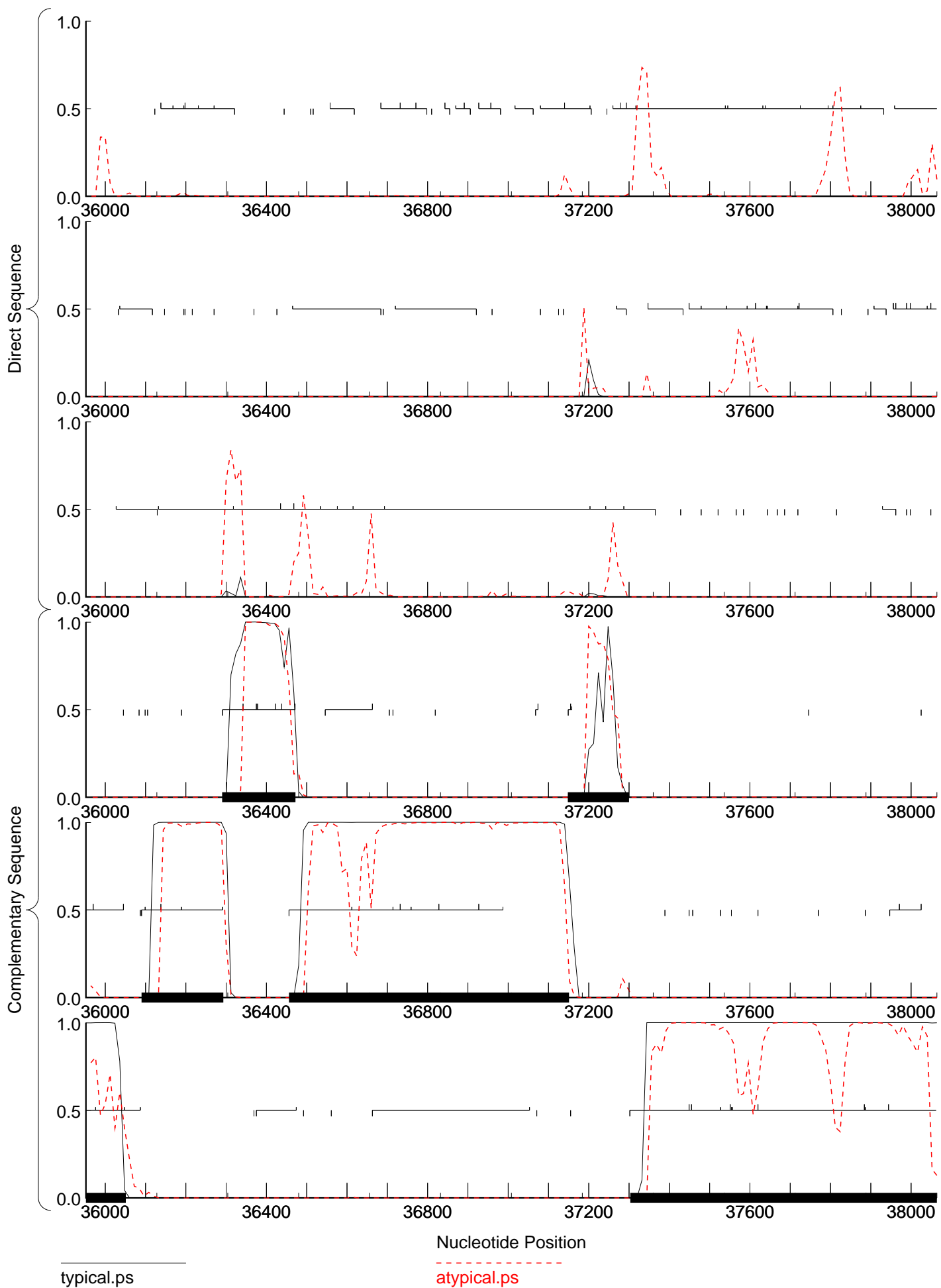
typical.ps

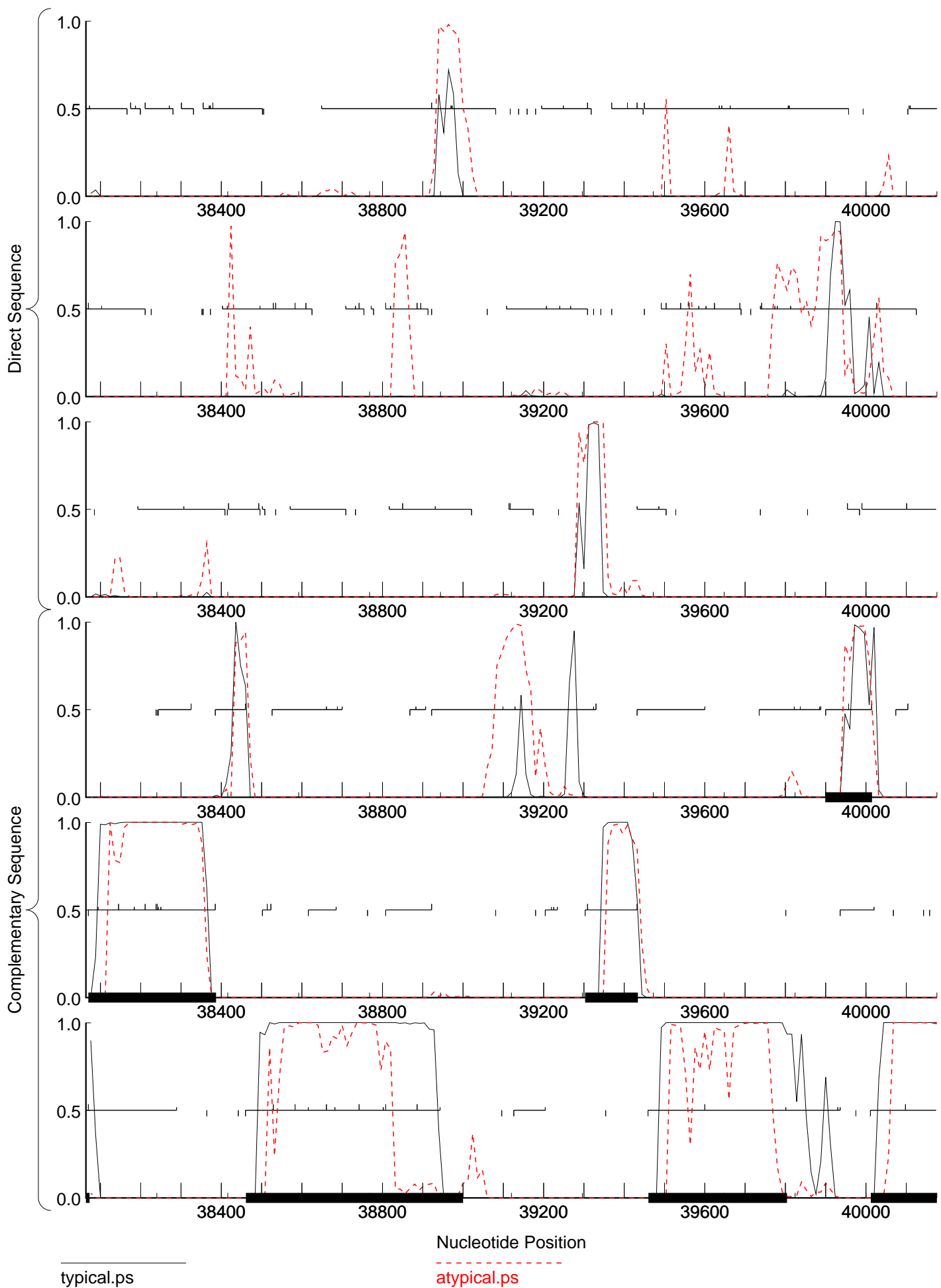
atypical.ps

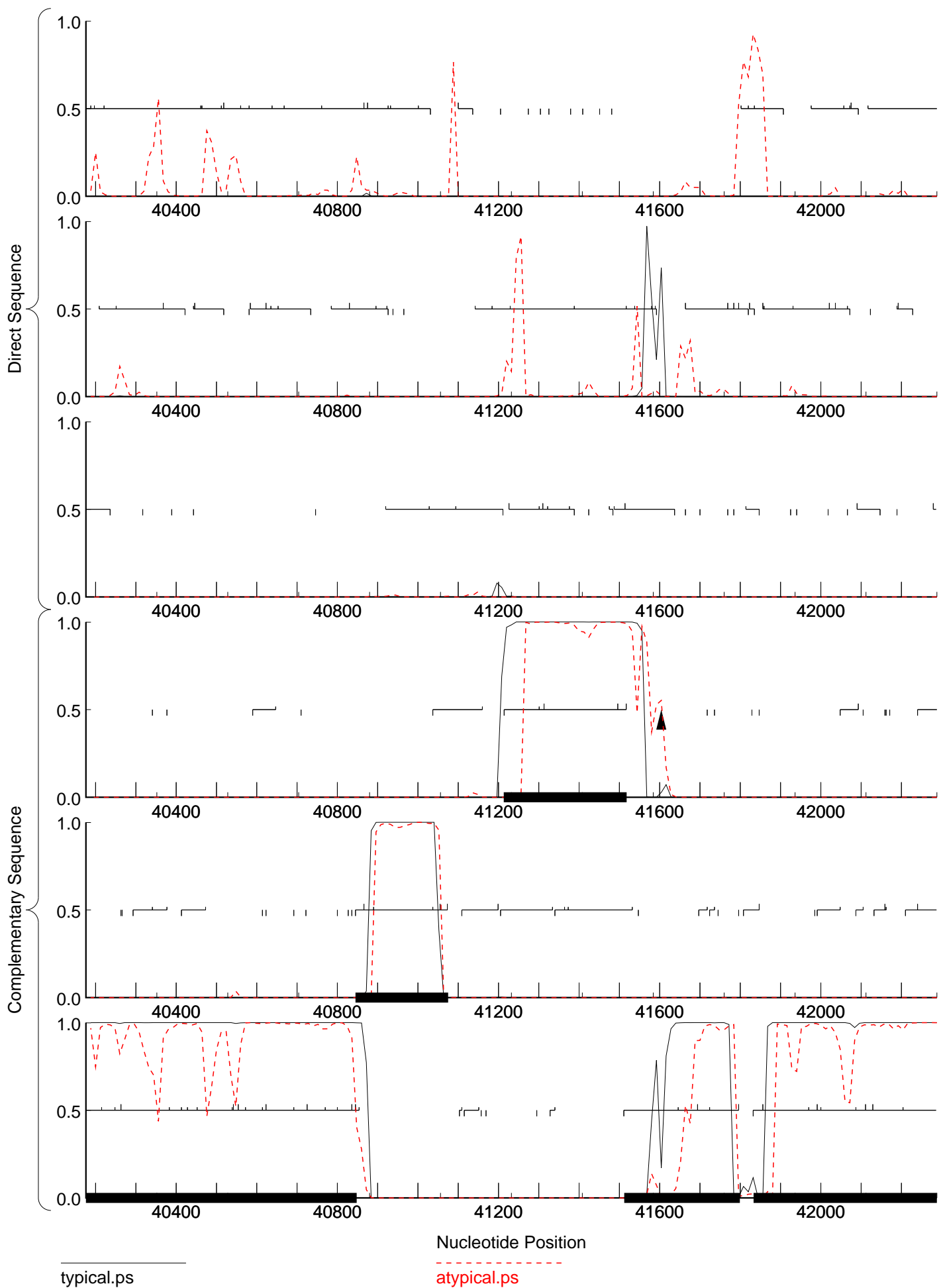


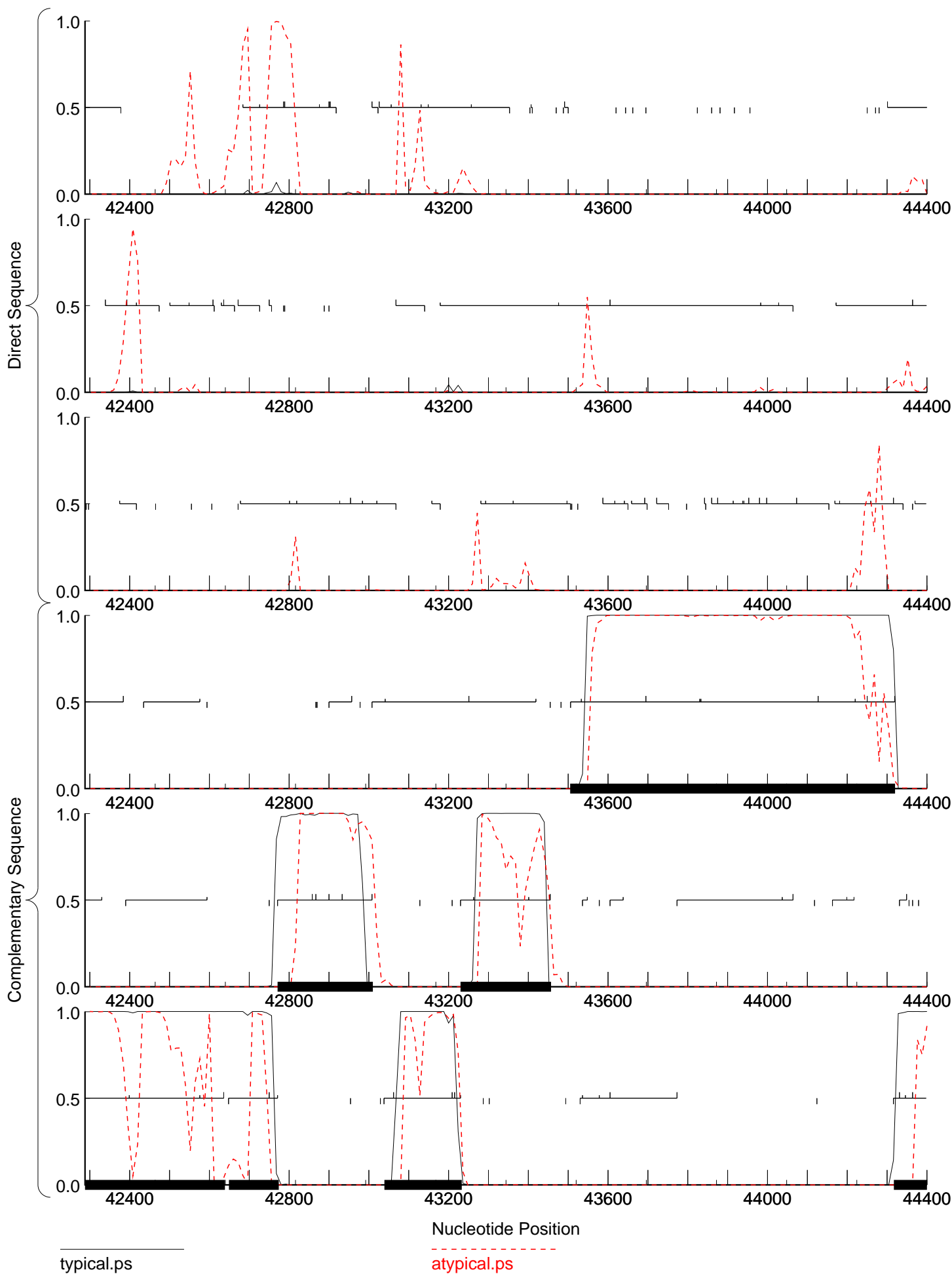
typical.ps

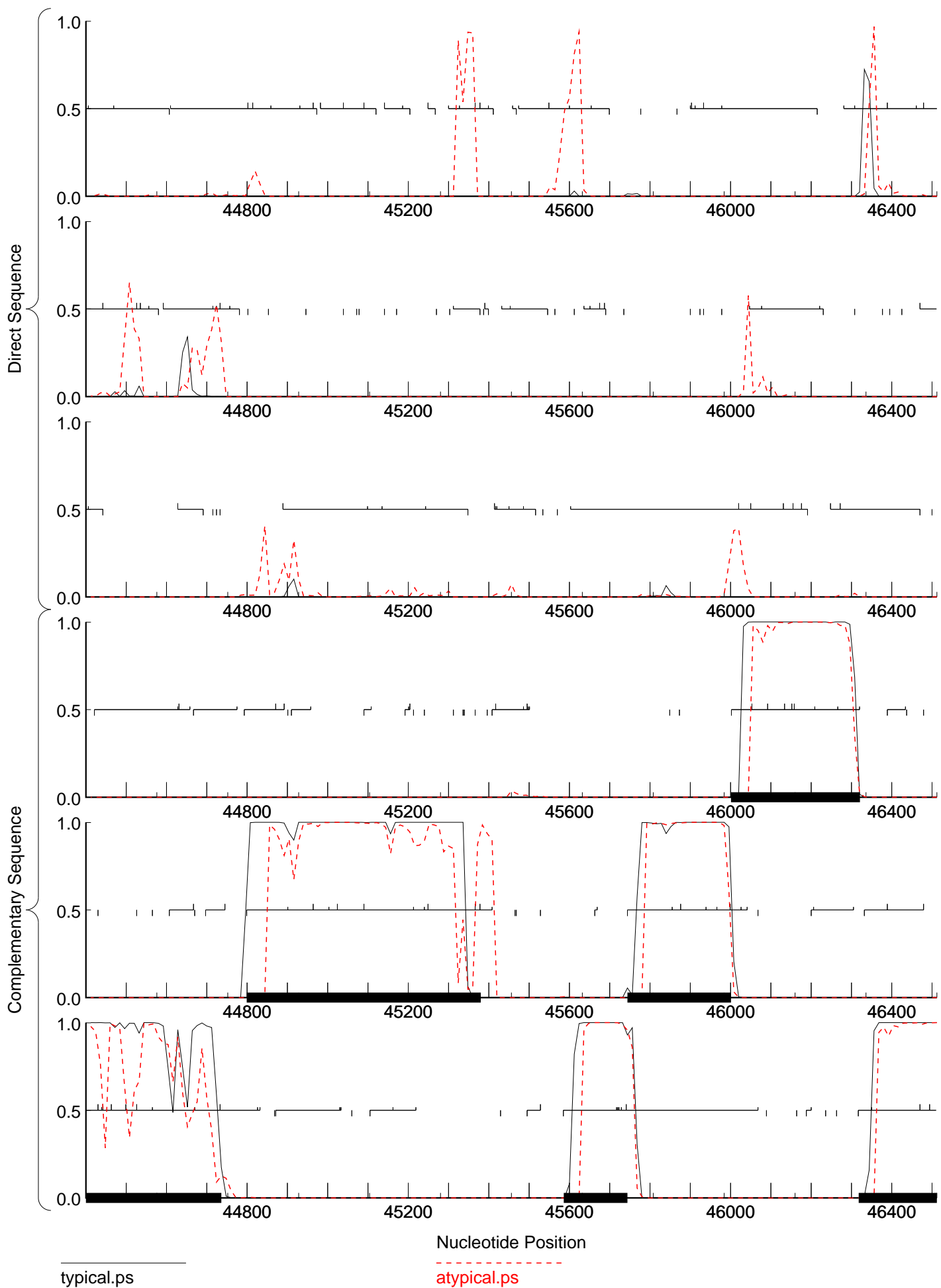
atypical.ps

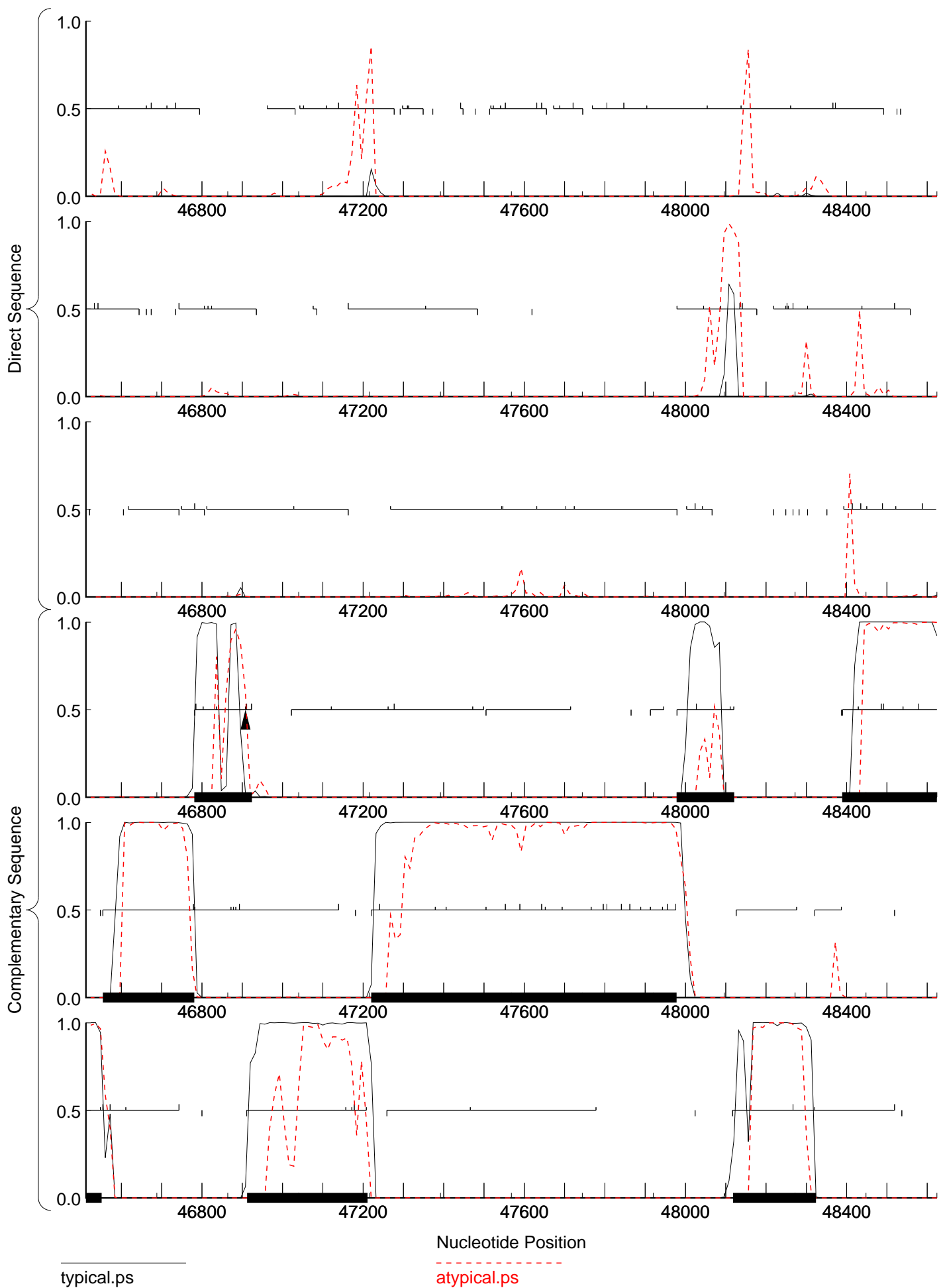


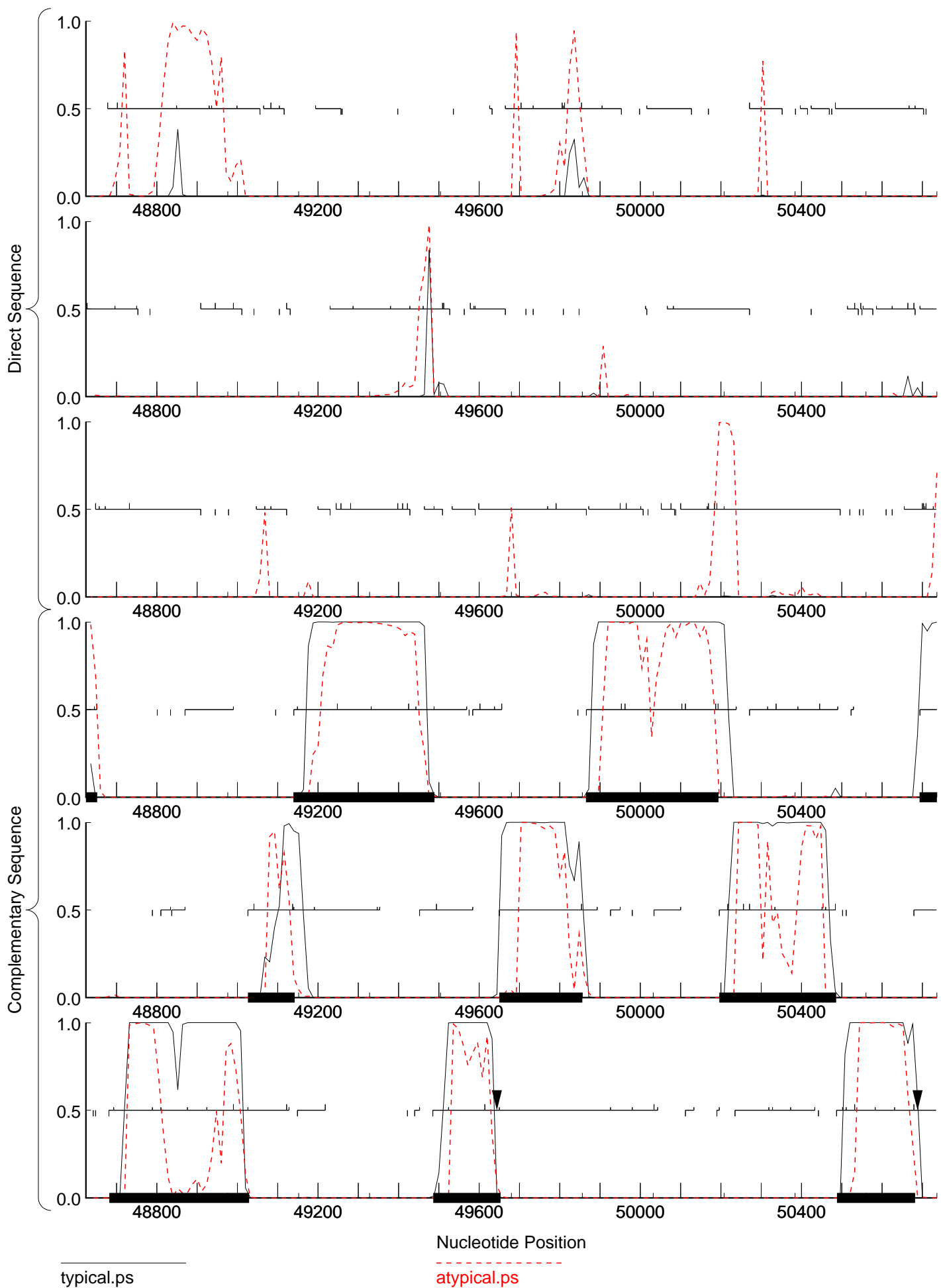












typical.ps

atypical.ps

