

GeneMark

Version 2.5p (09.08.06)

Copyright 1993 - M. Borodovsky, J. McIninch

PROGRAM INFORMATION

Sequence : Mycobacterium phage Haleema complete sequence, 68415 bp, circularly permuted, Cluster B1
Analysis Date : 12/15/17 at 7:14:18
Pages : 34
Sequence Length : 68415 bp
GC Content : 66.50%

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,

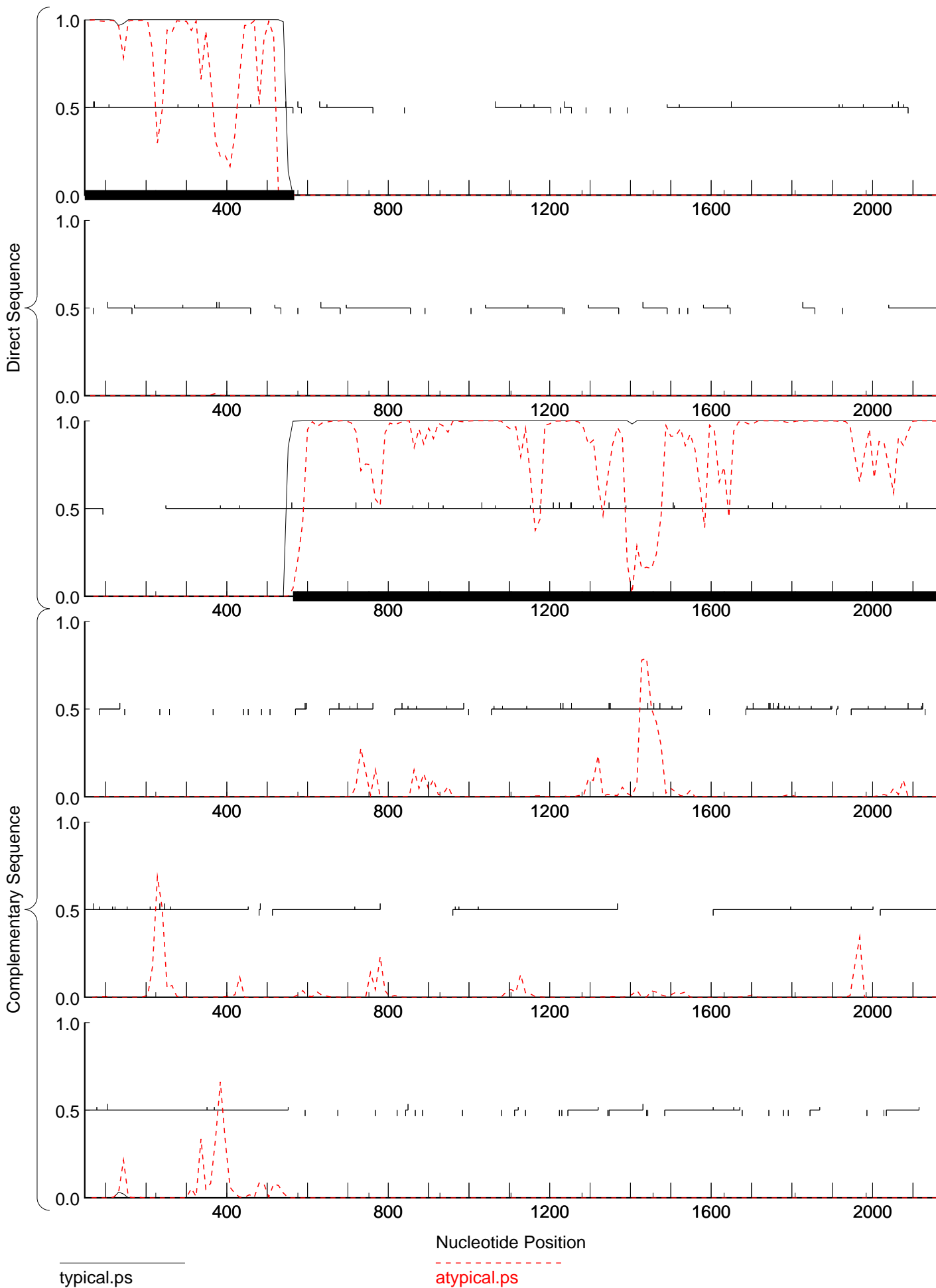
MATRIX INFORMATION

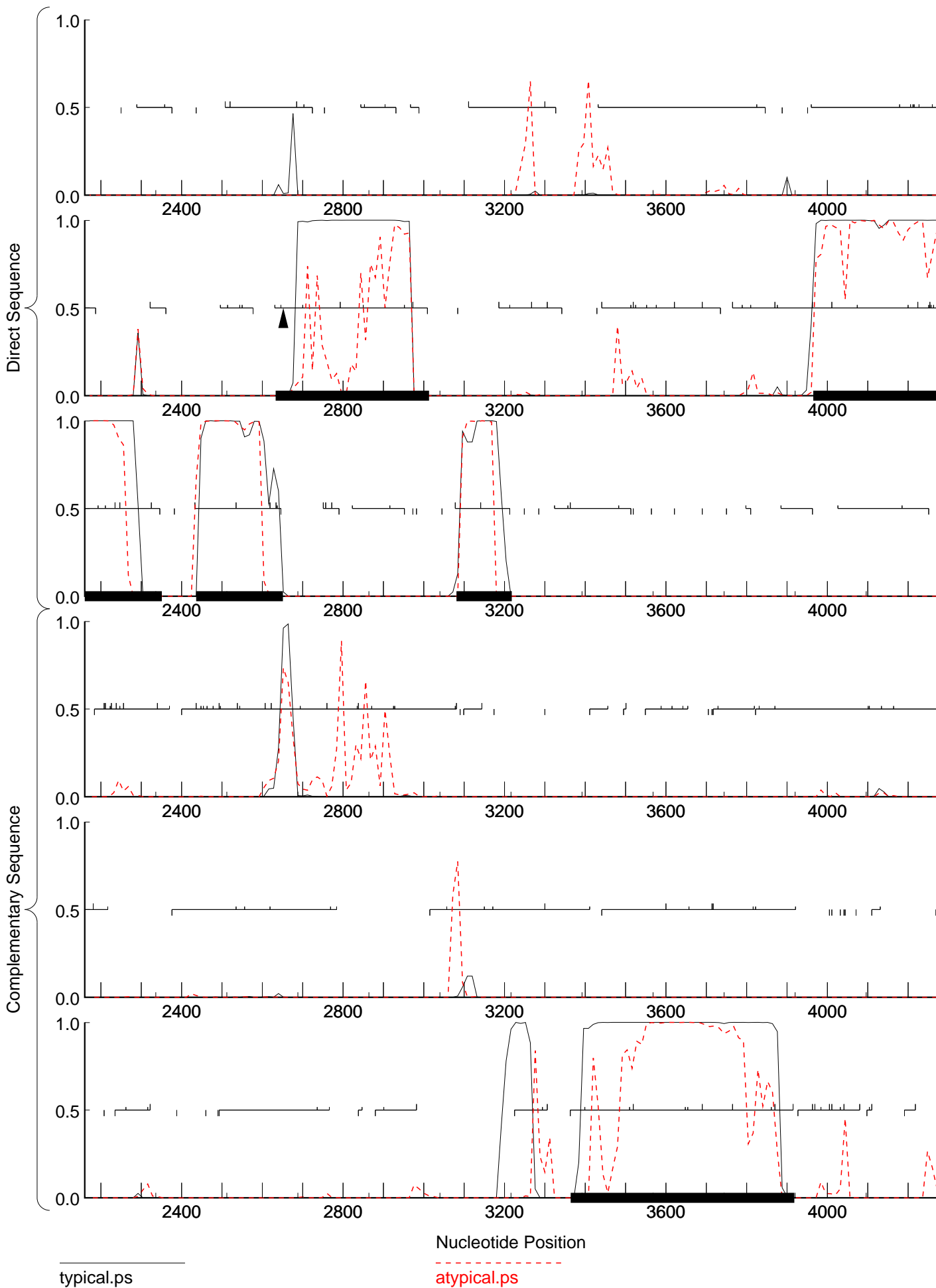
Matrix : -
Author : -
Order : 4

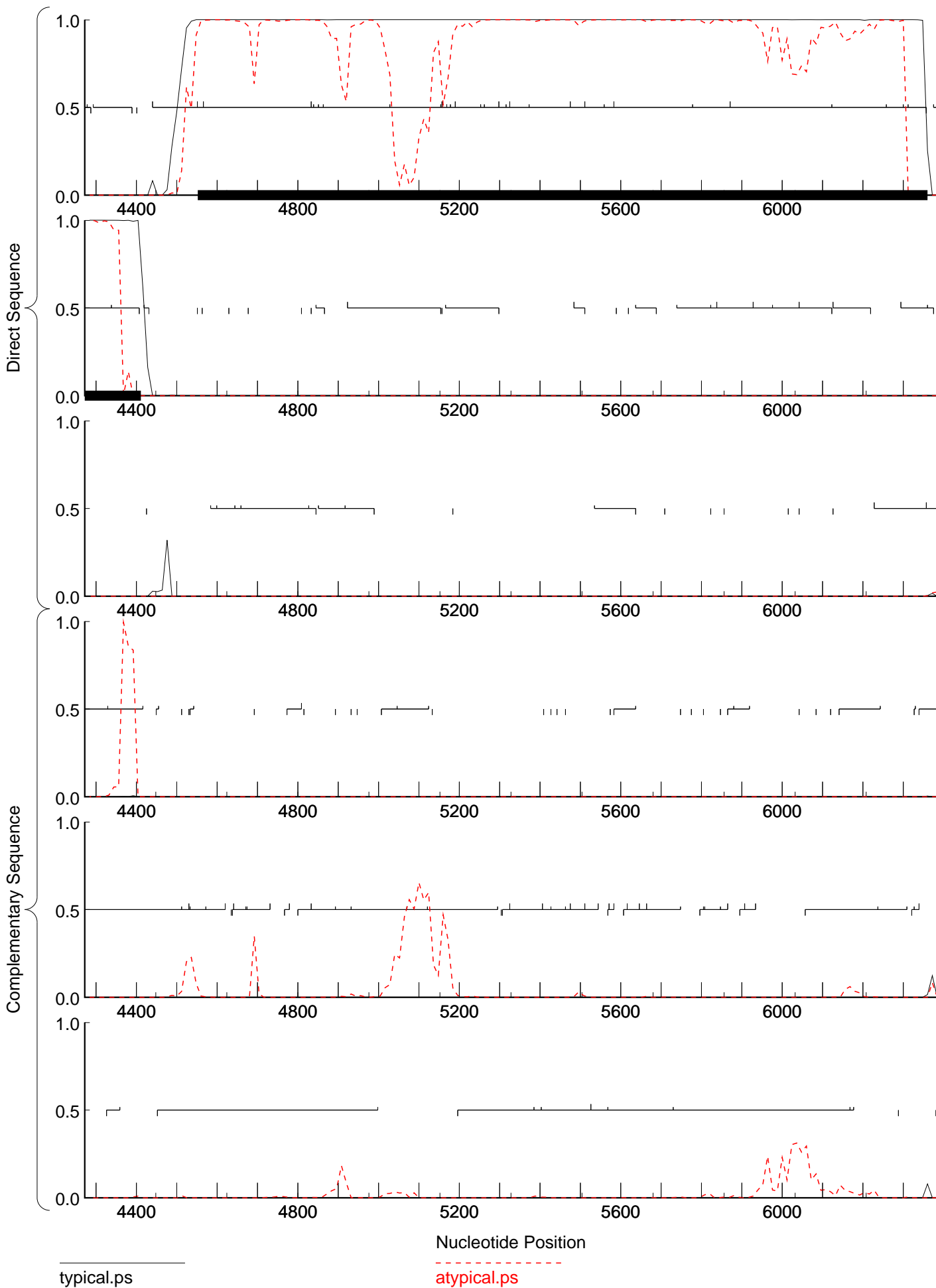
Send questions / comments to:
Dr. M. Borodovsky
Georgia Institute of Technology
School of Biology
Atlanta, GA 30332-0230

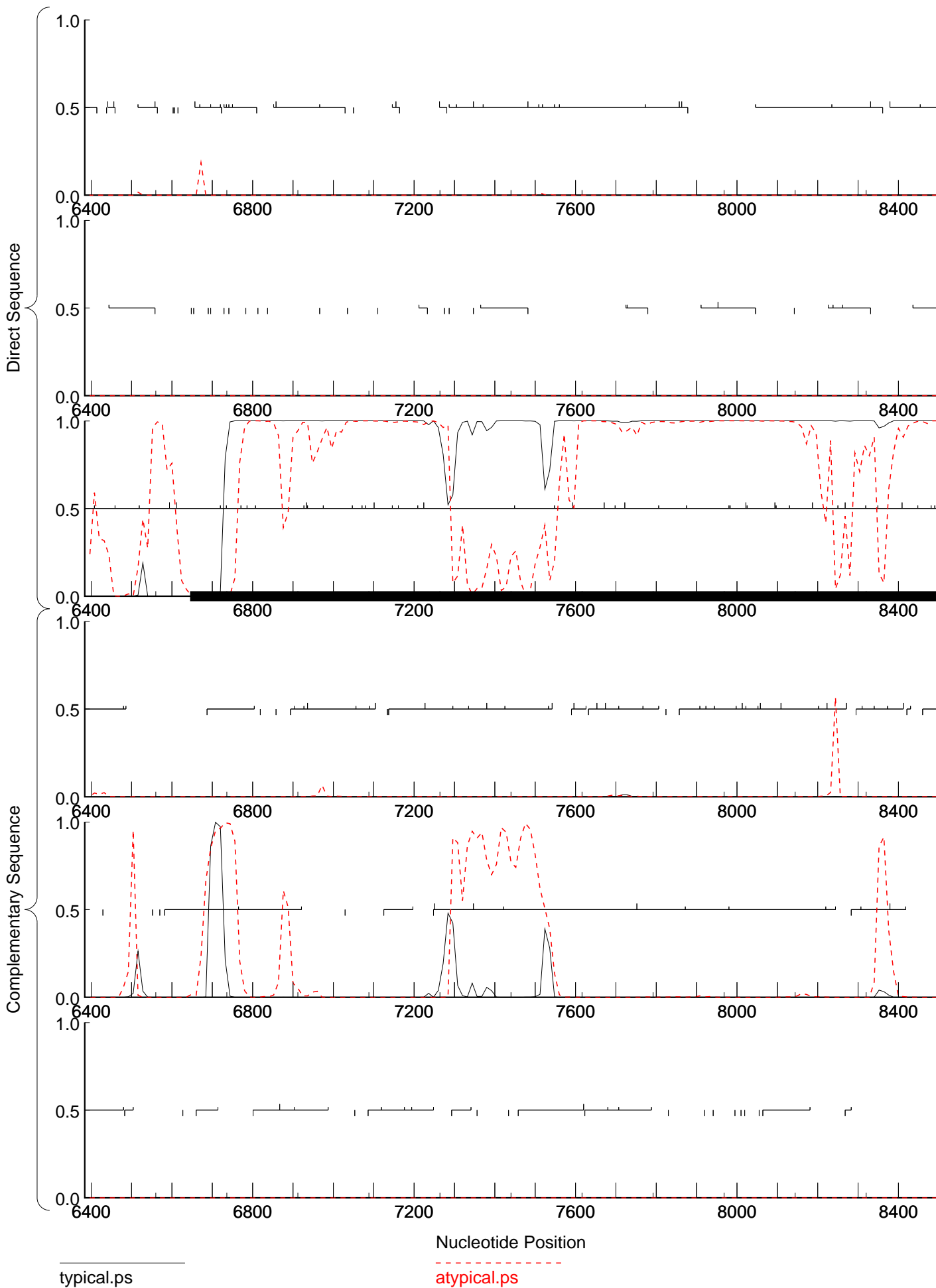
Matrix notes & comments

Training set derived by GeneMarkS, 4.30 October 2014
Fri Dec 15 07:14:18 2017



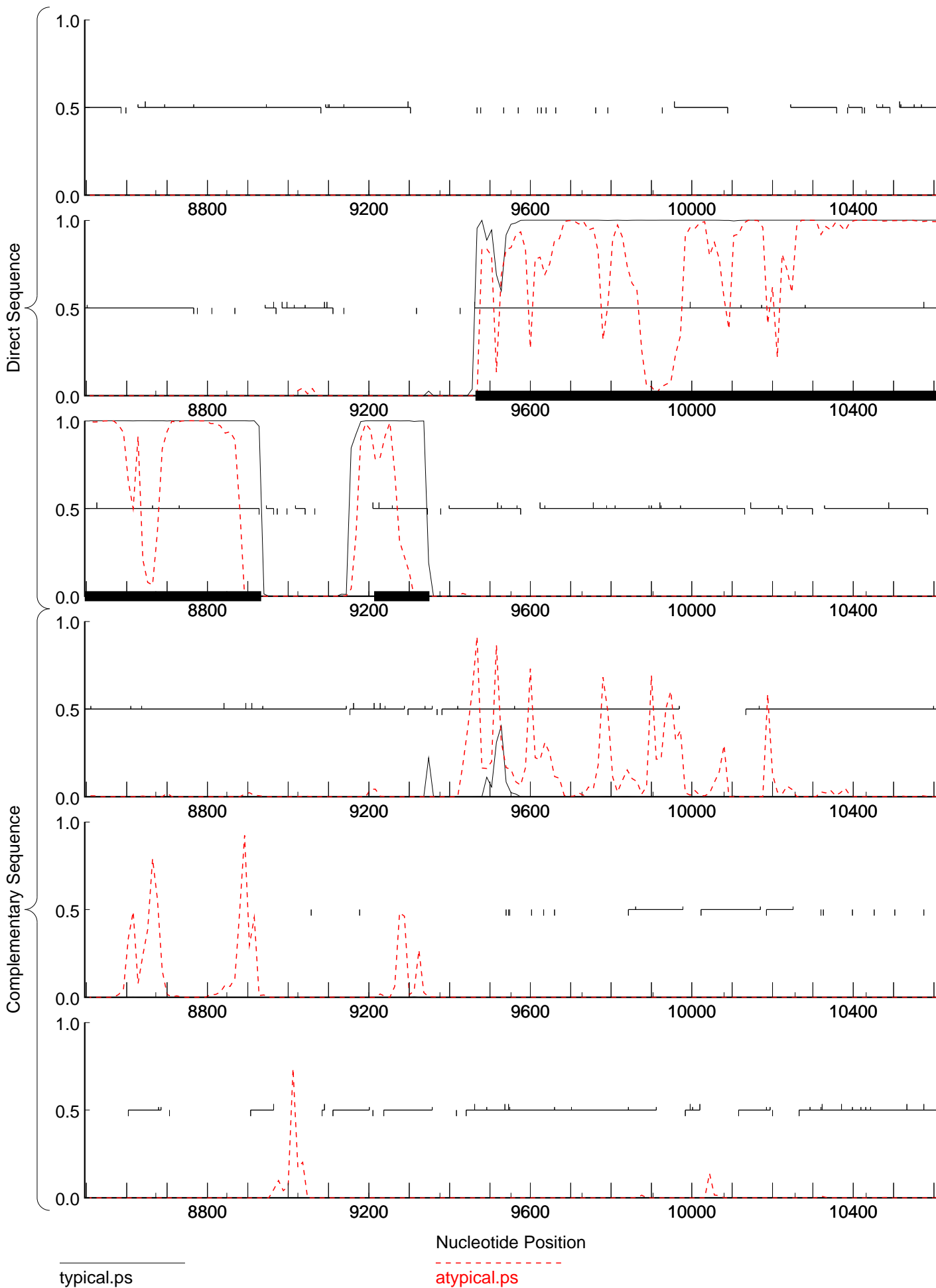


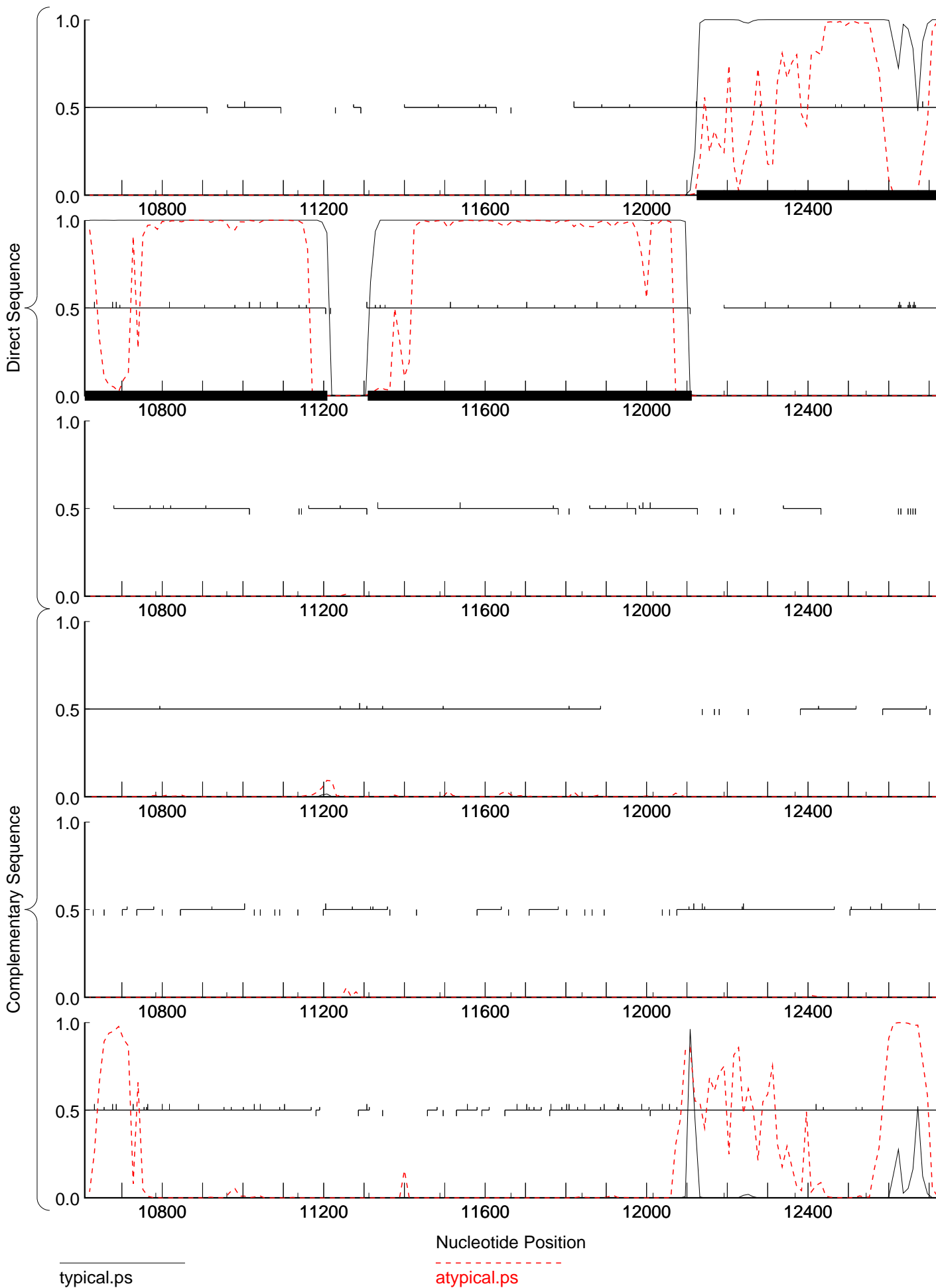


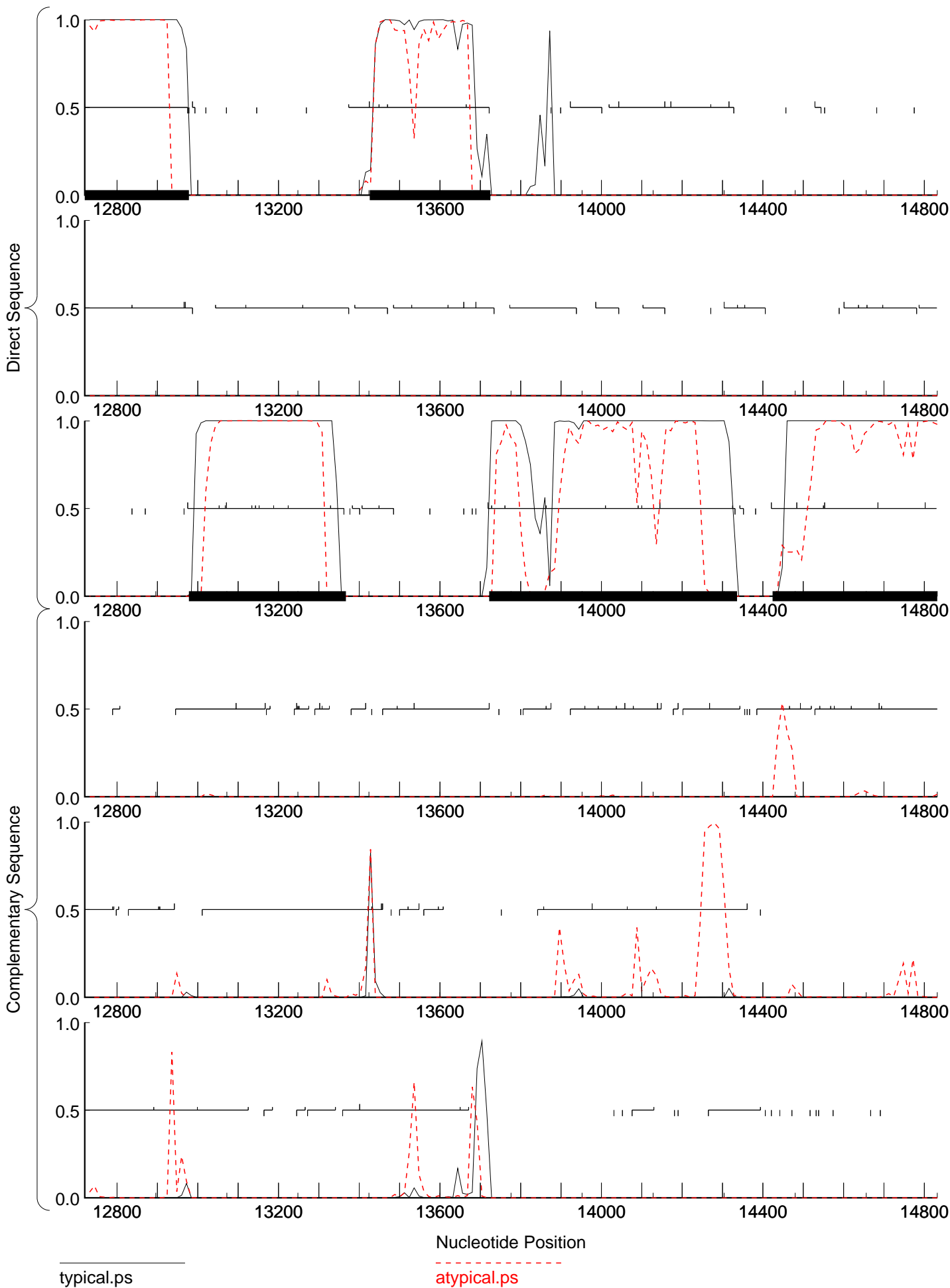


typical.ps

atypical.ps

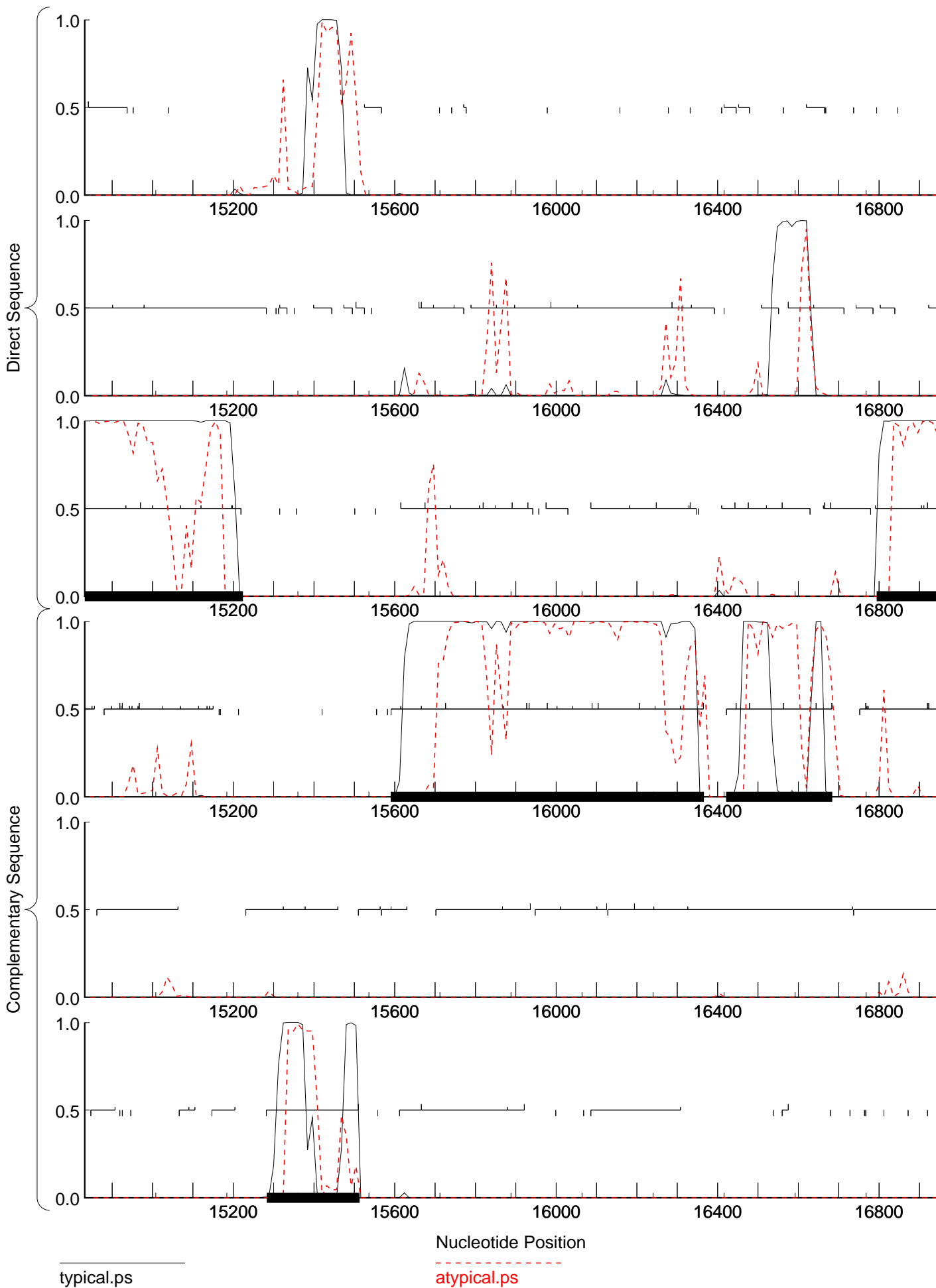


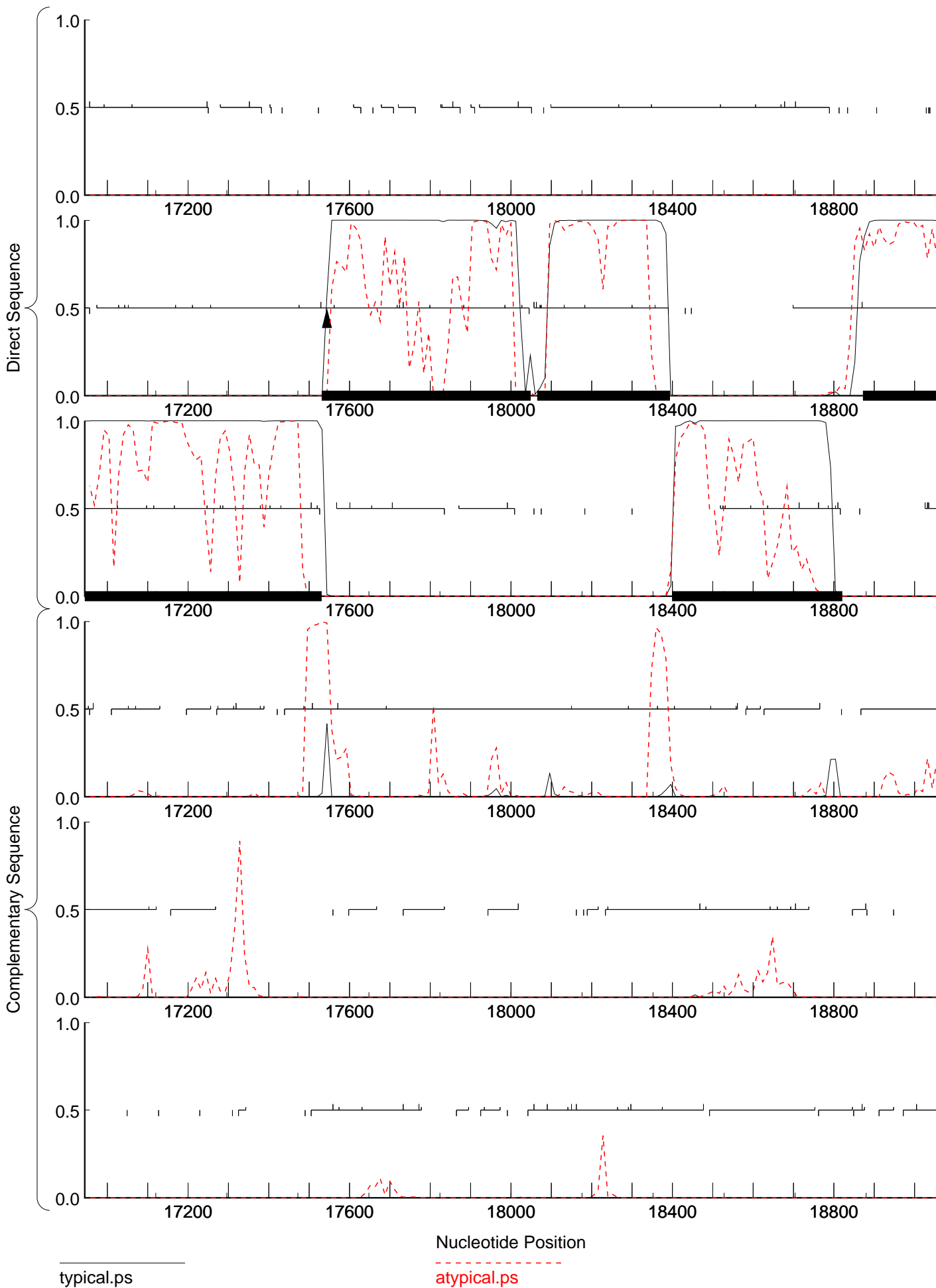


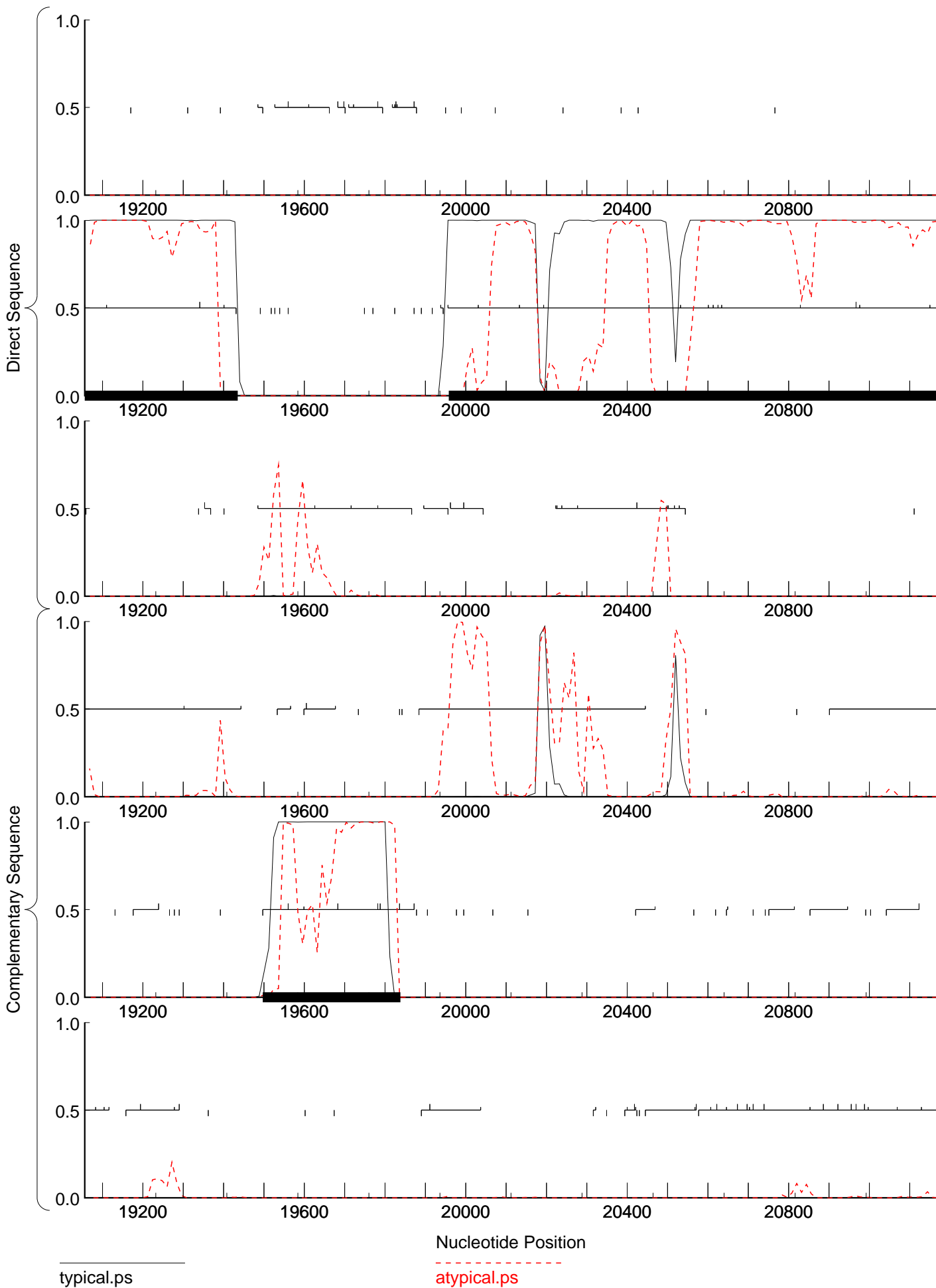


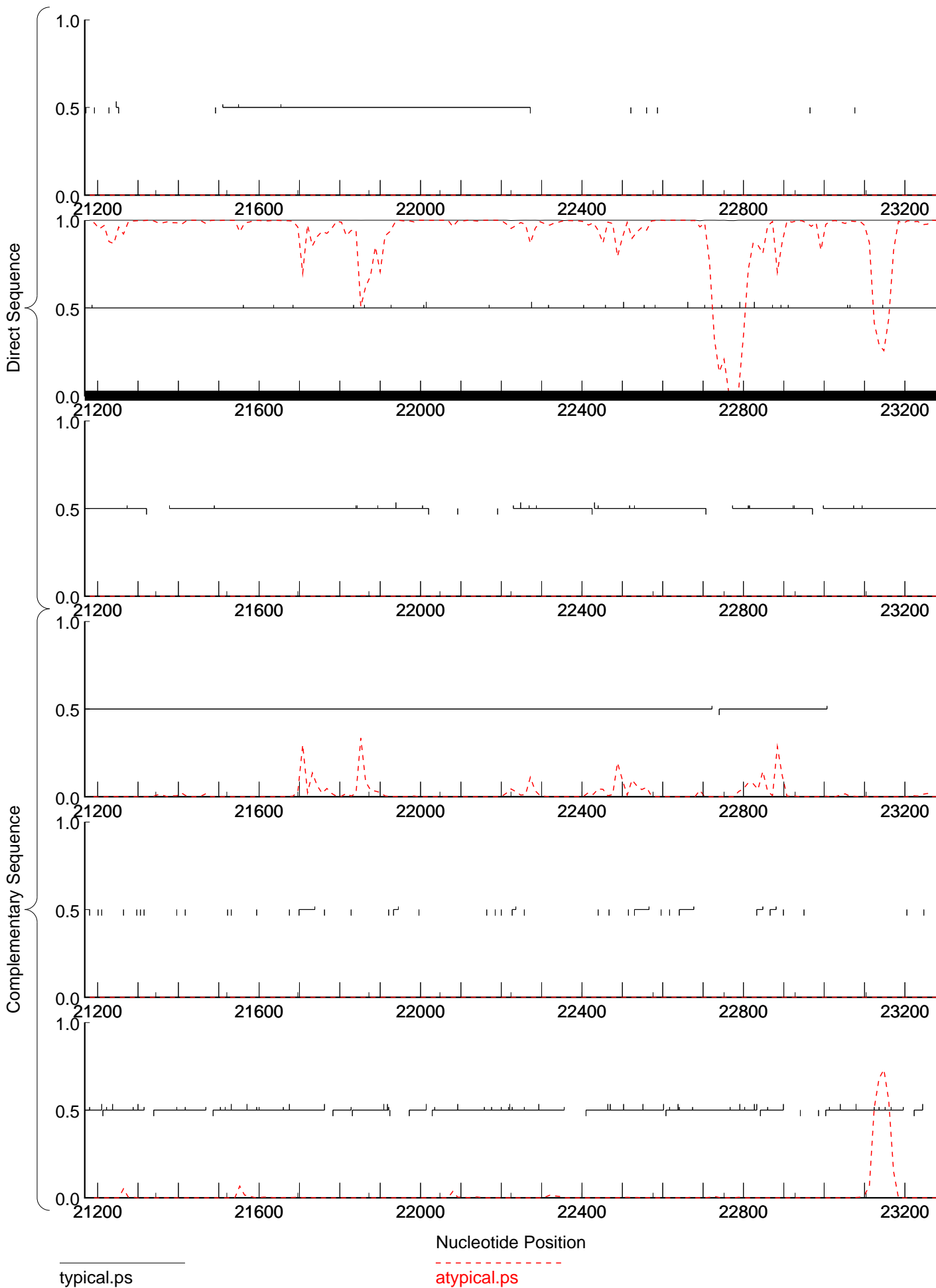
typical.ps

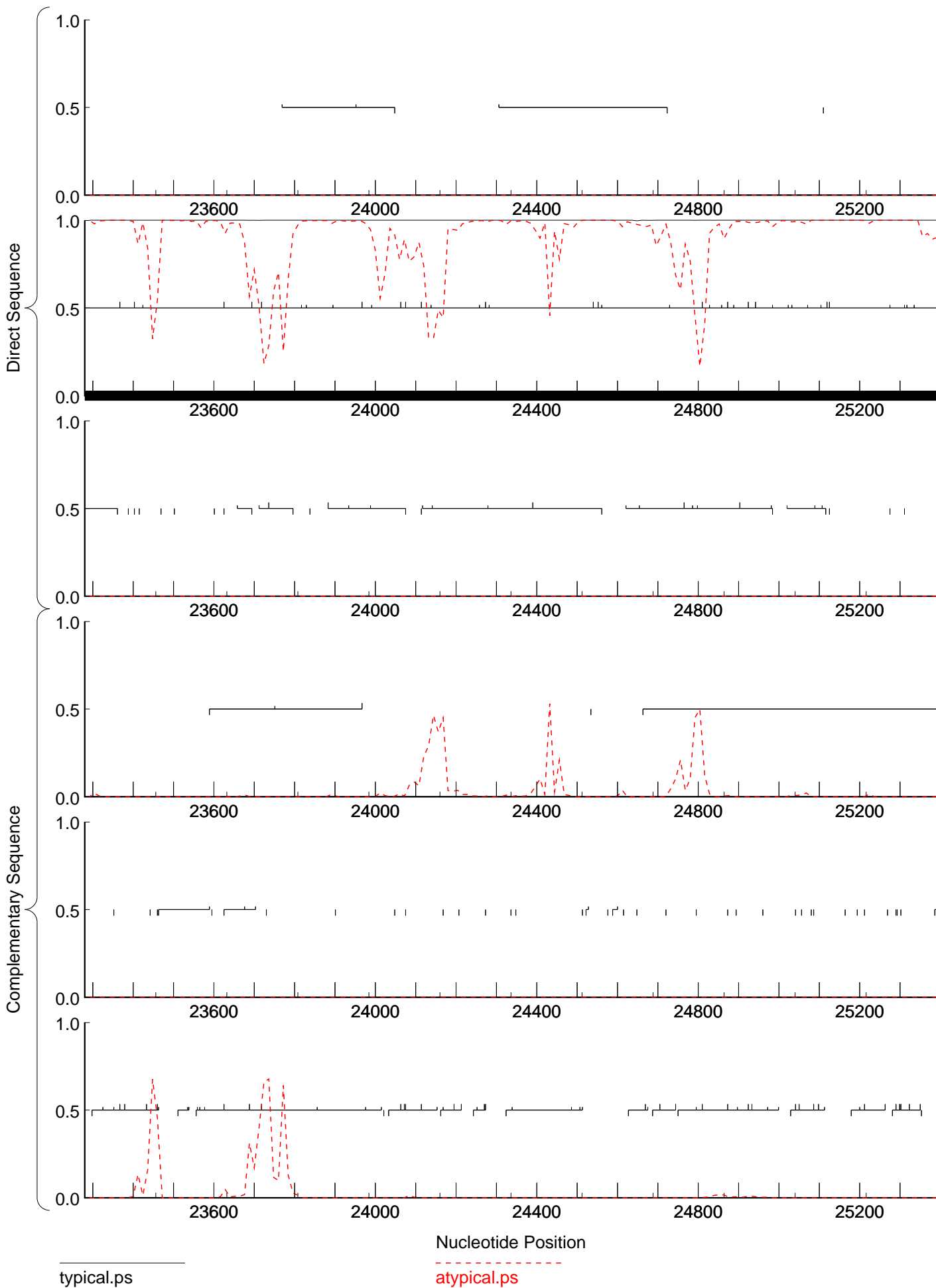
atypical.ps

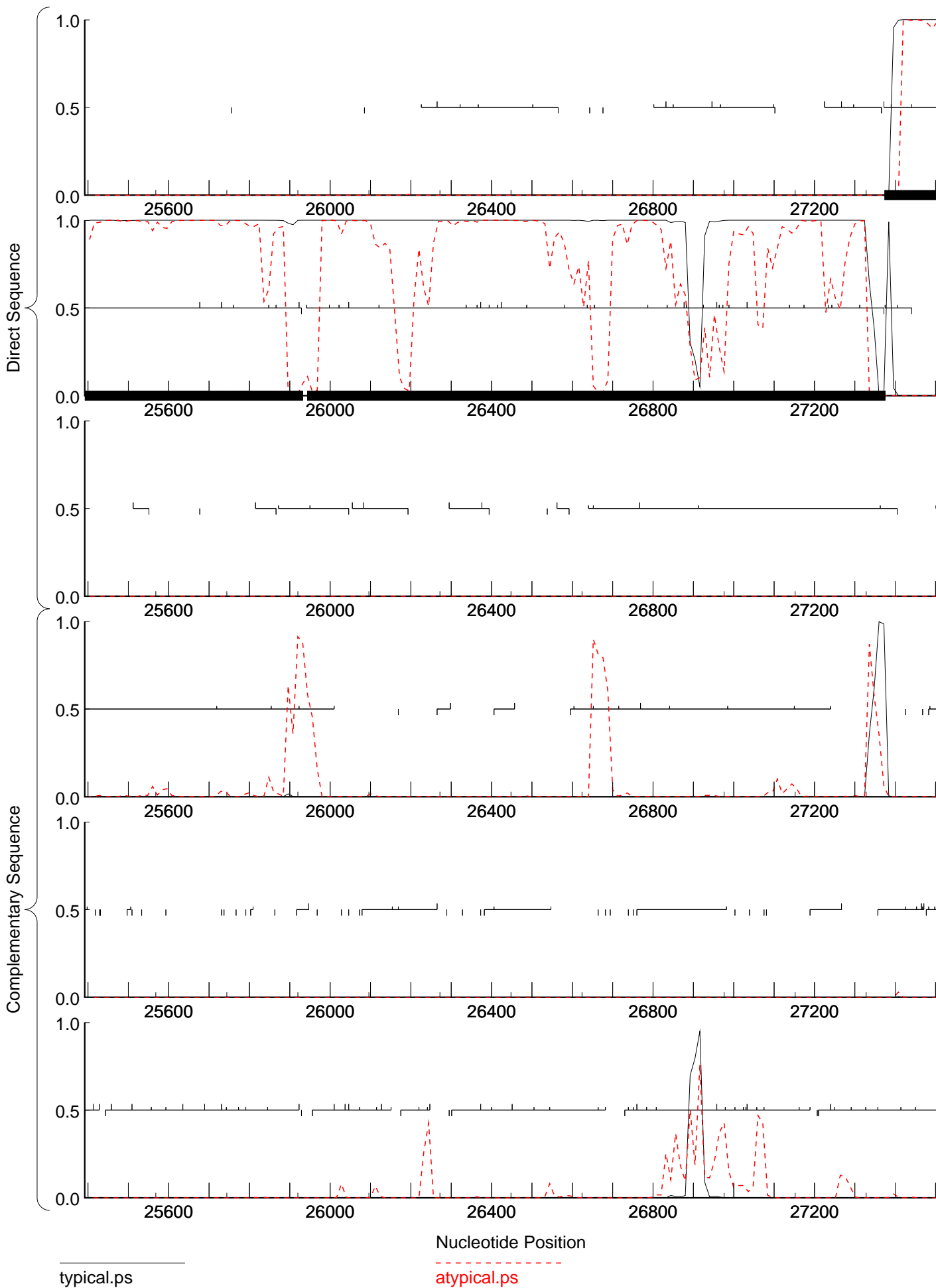


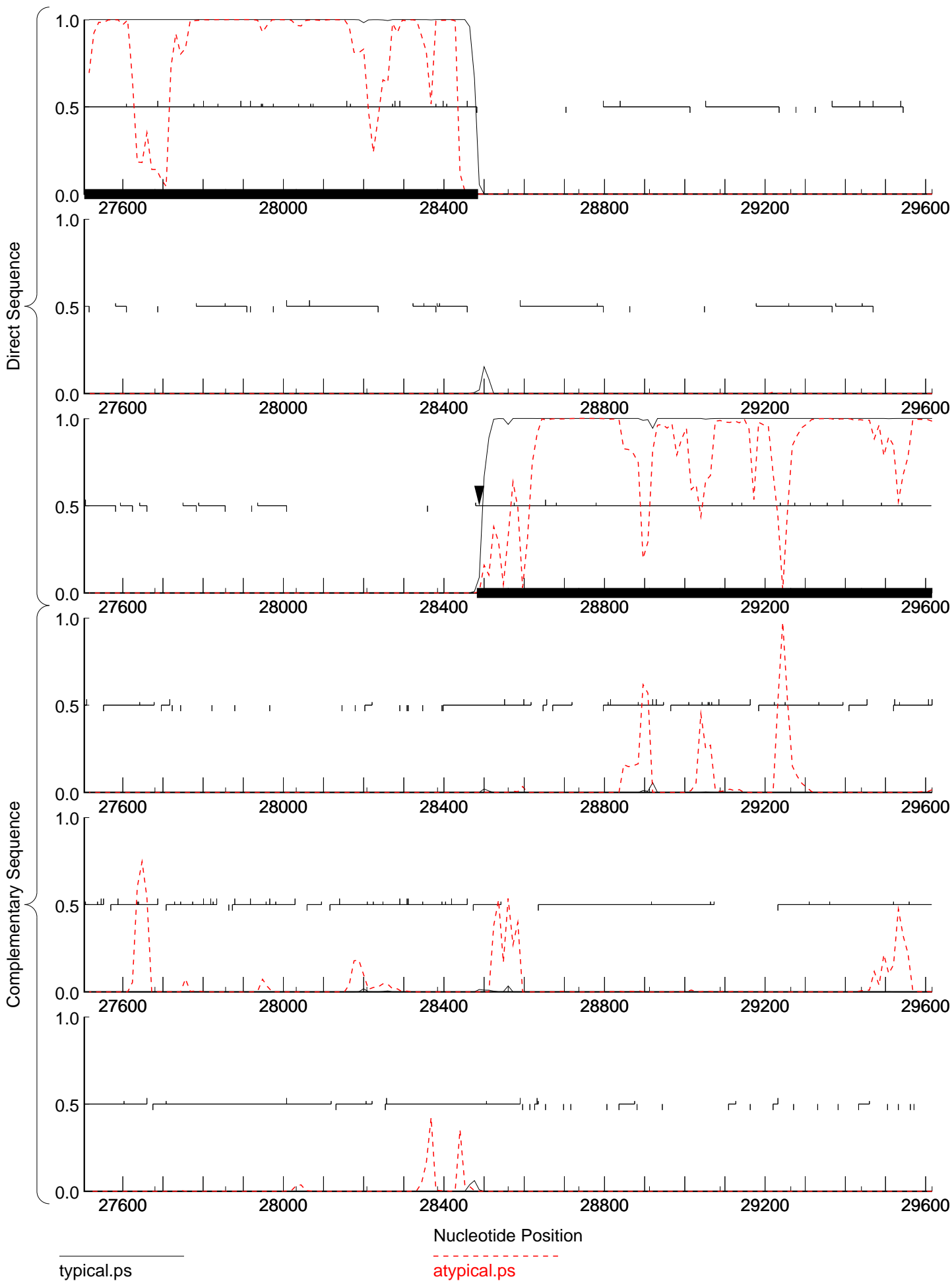


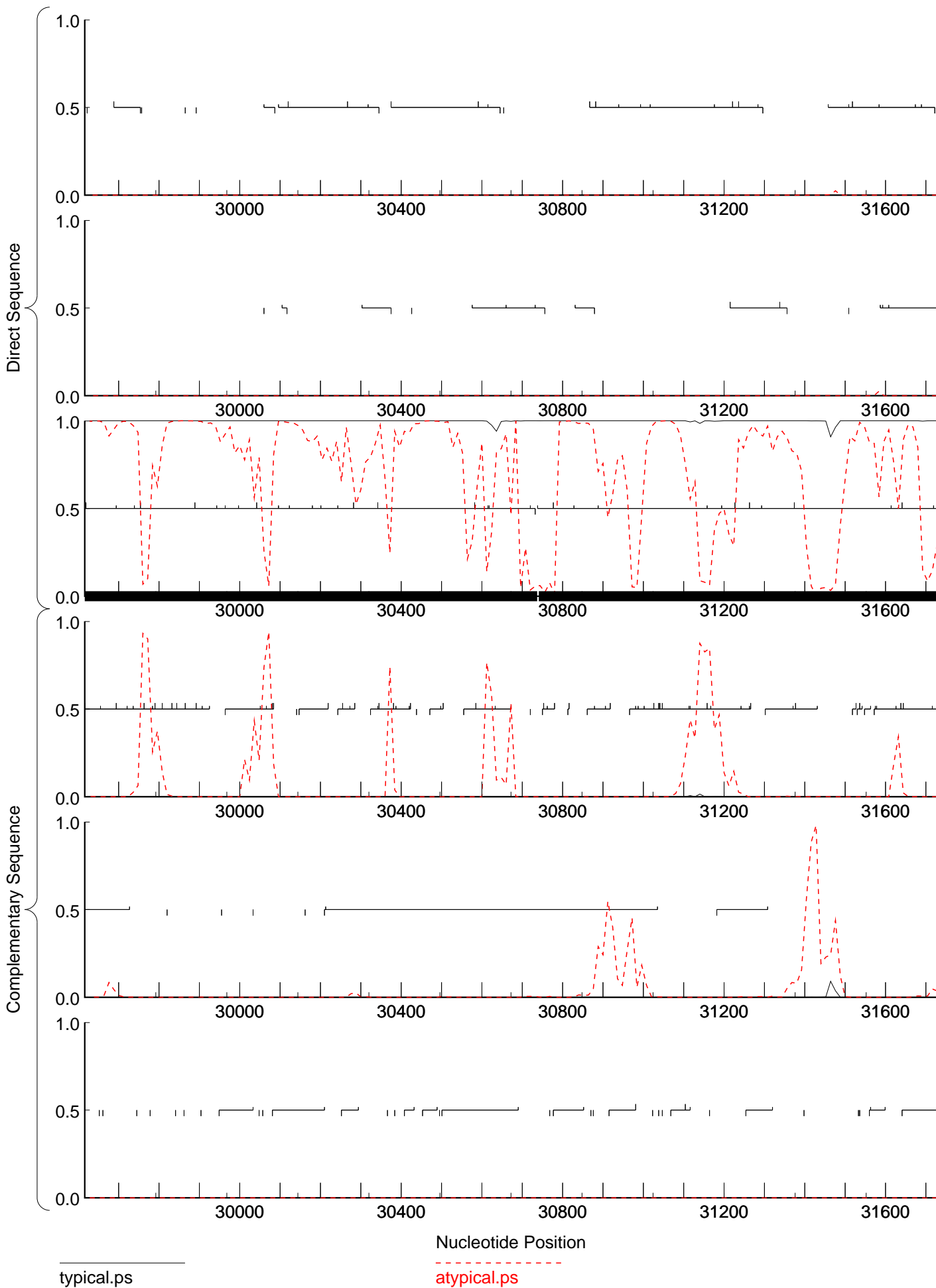


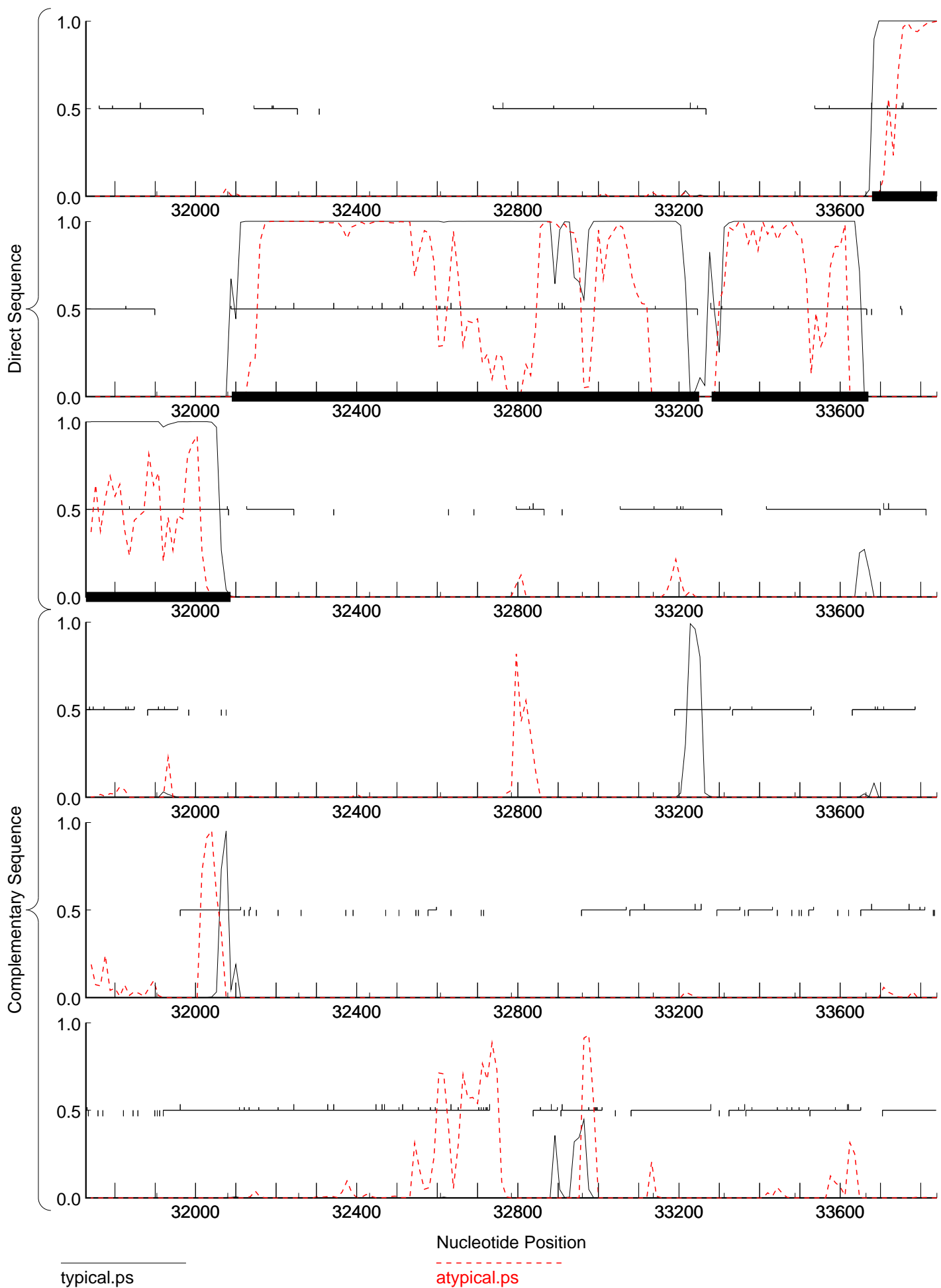


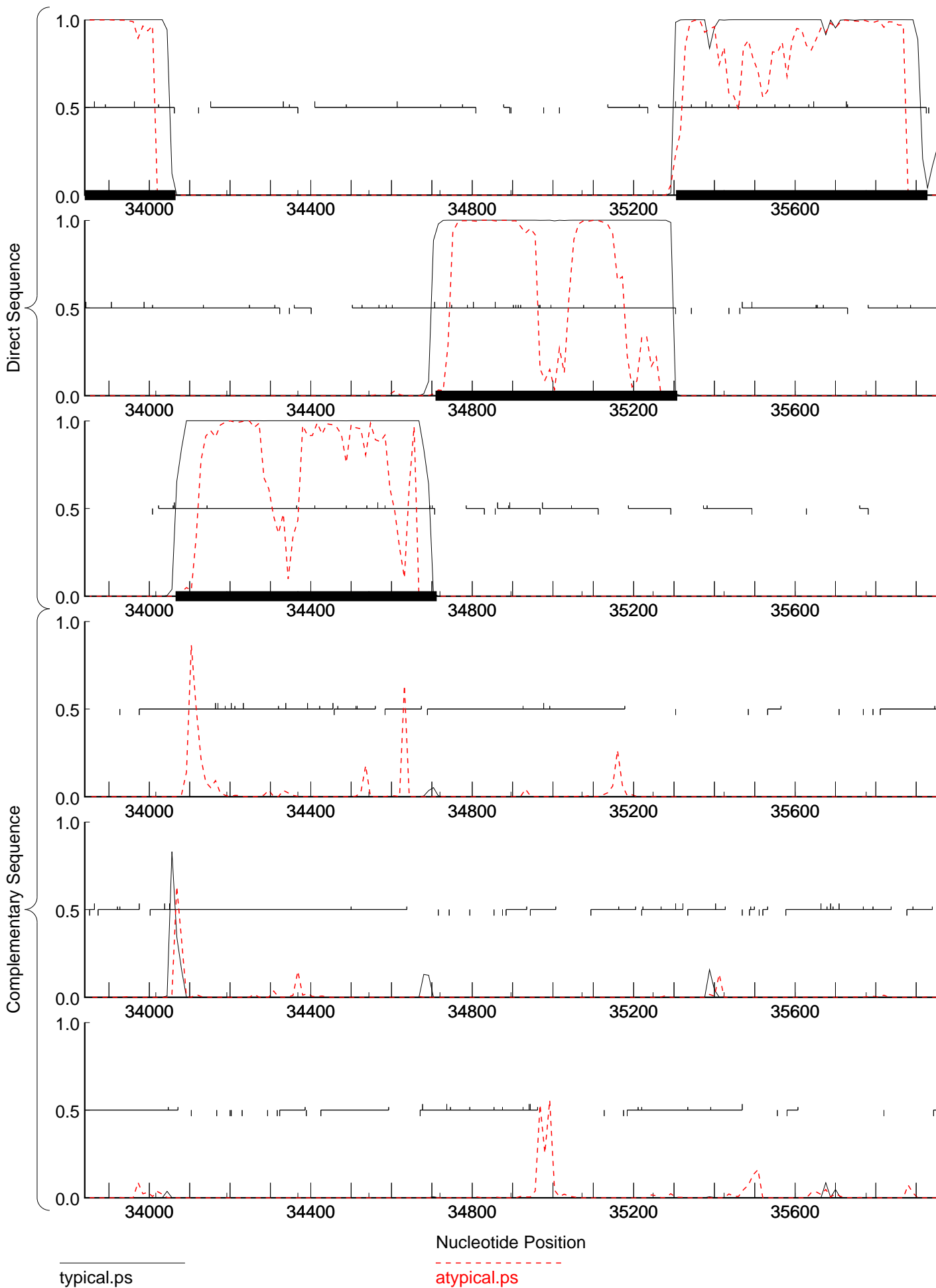


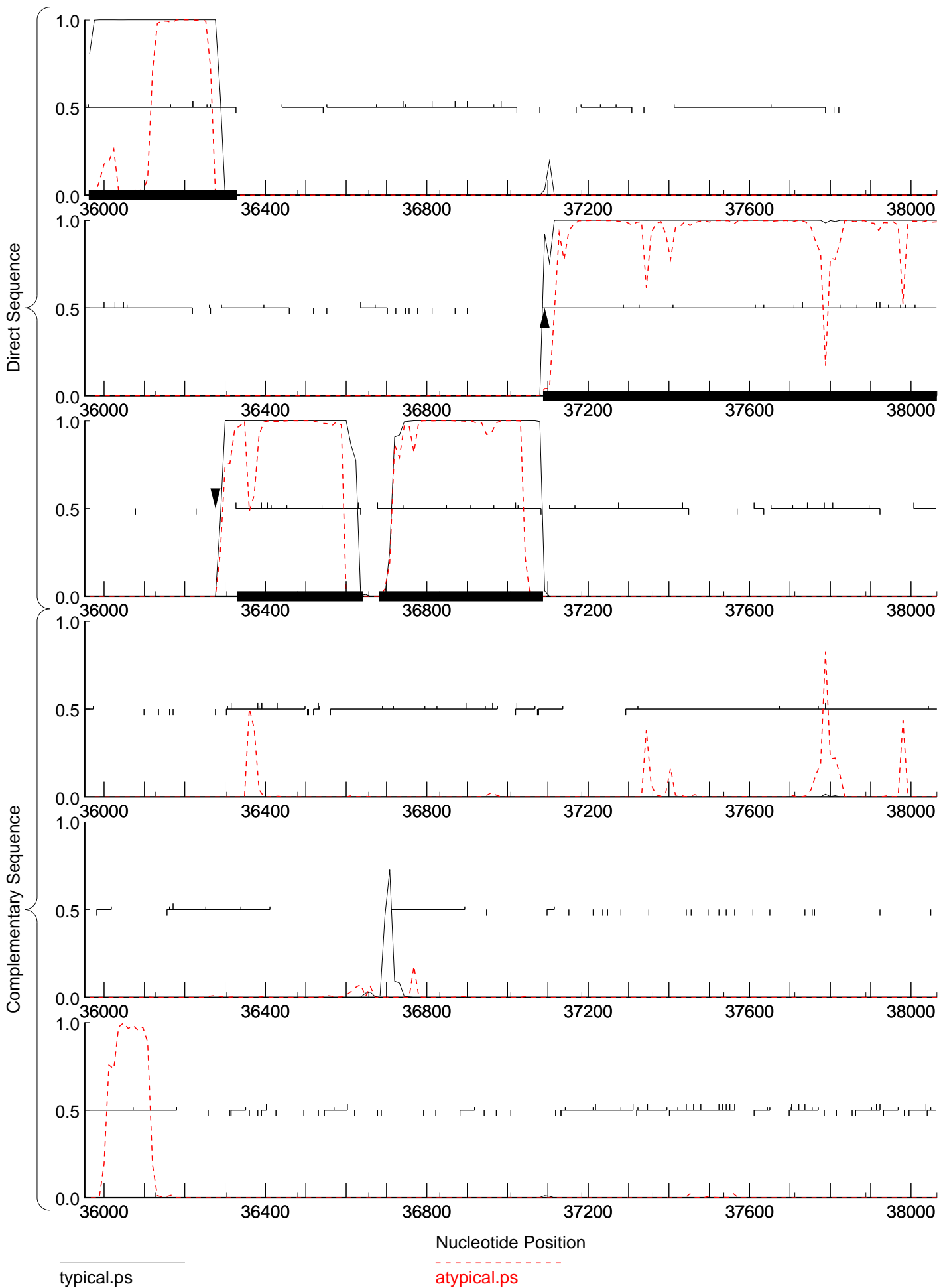


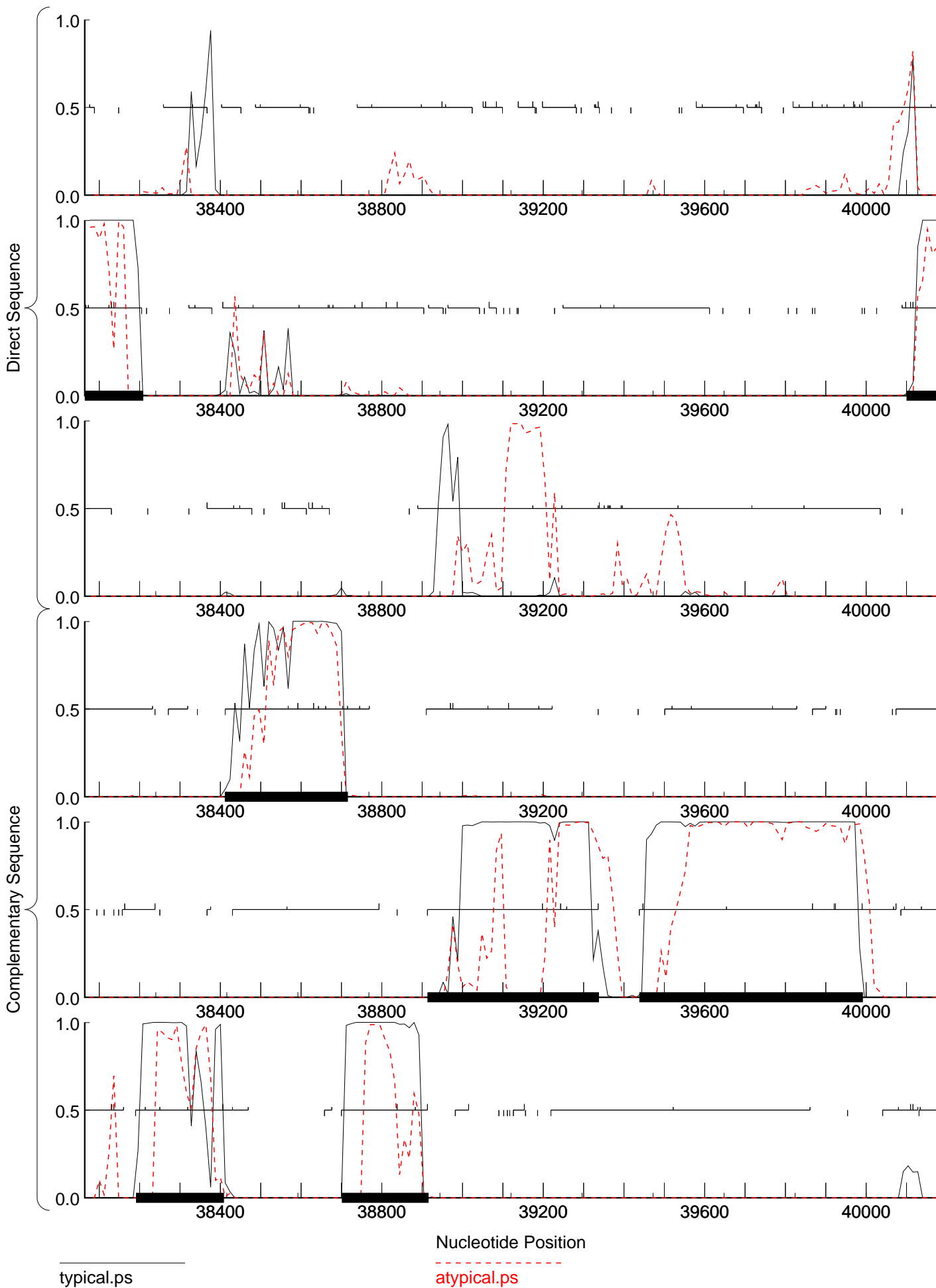






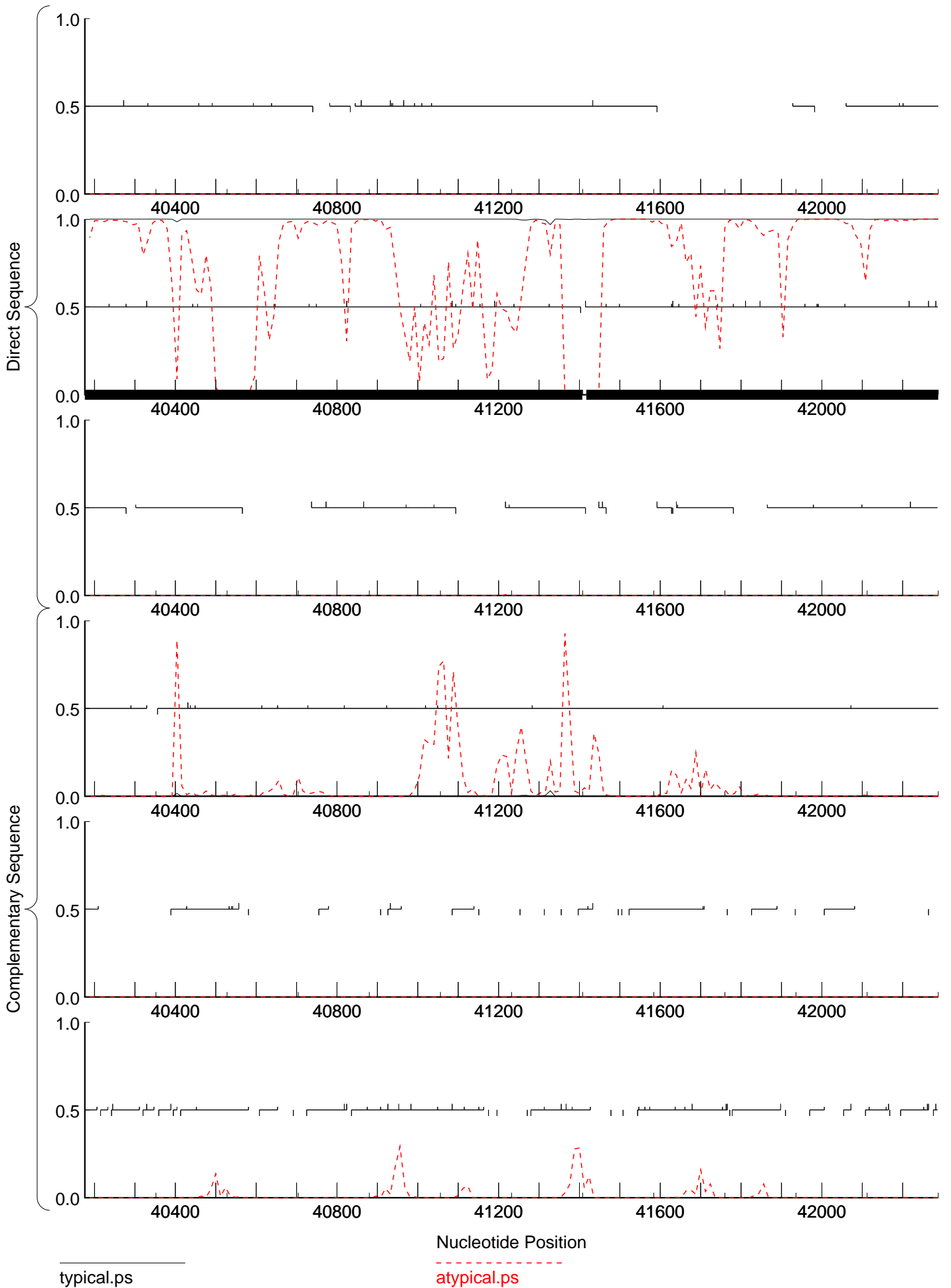


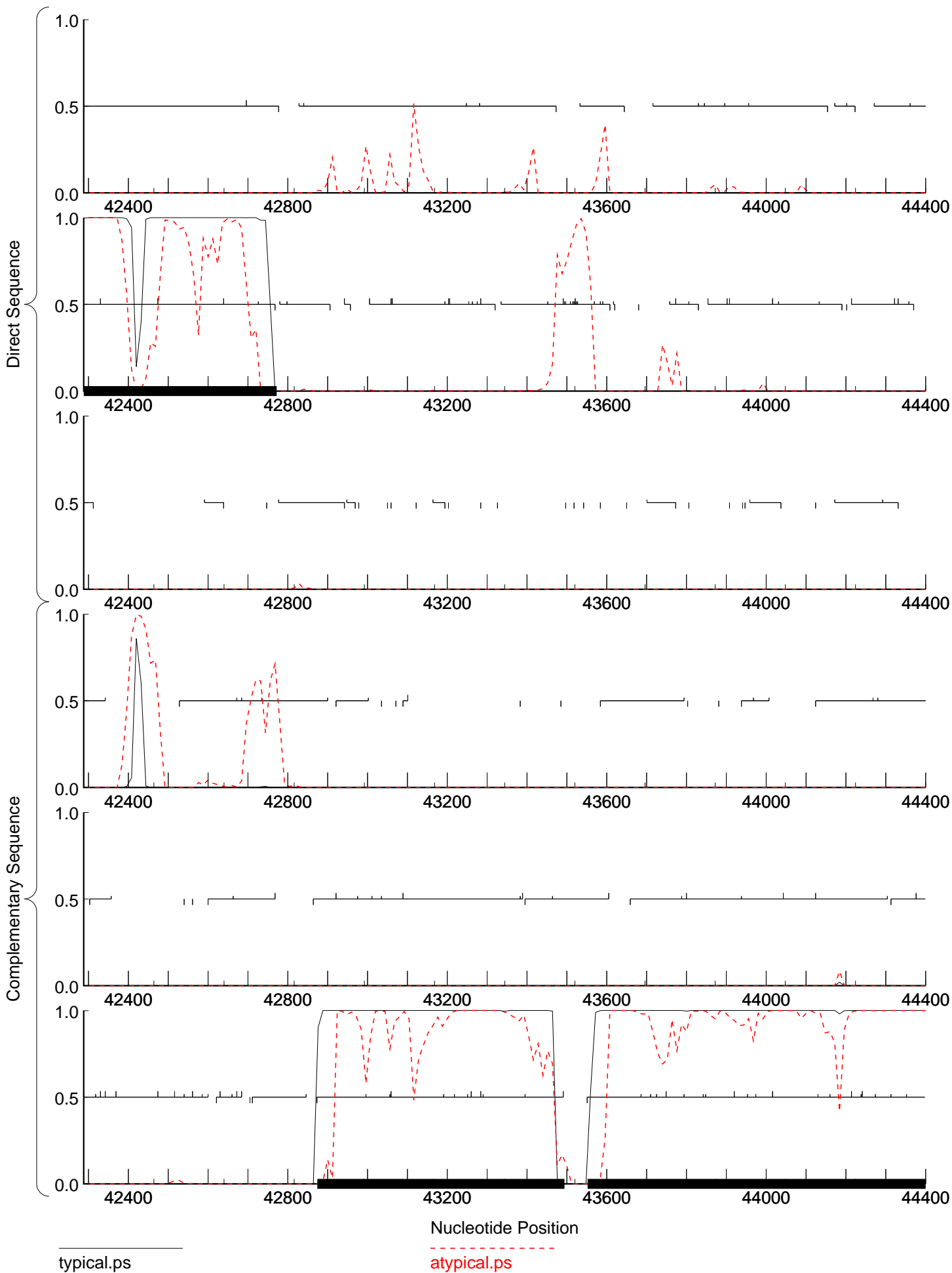


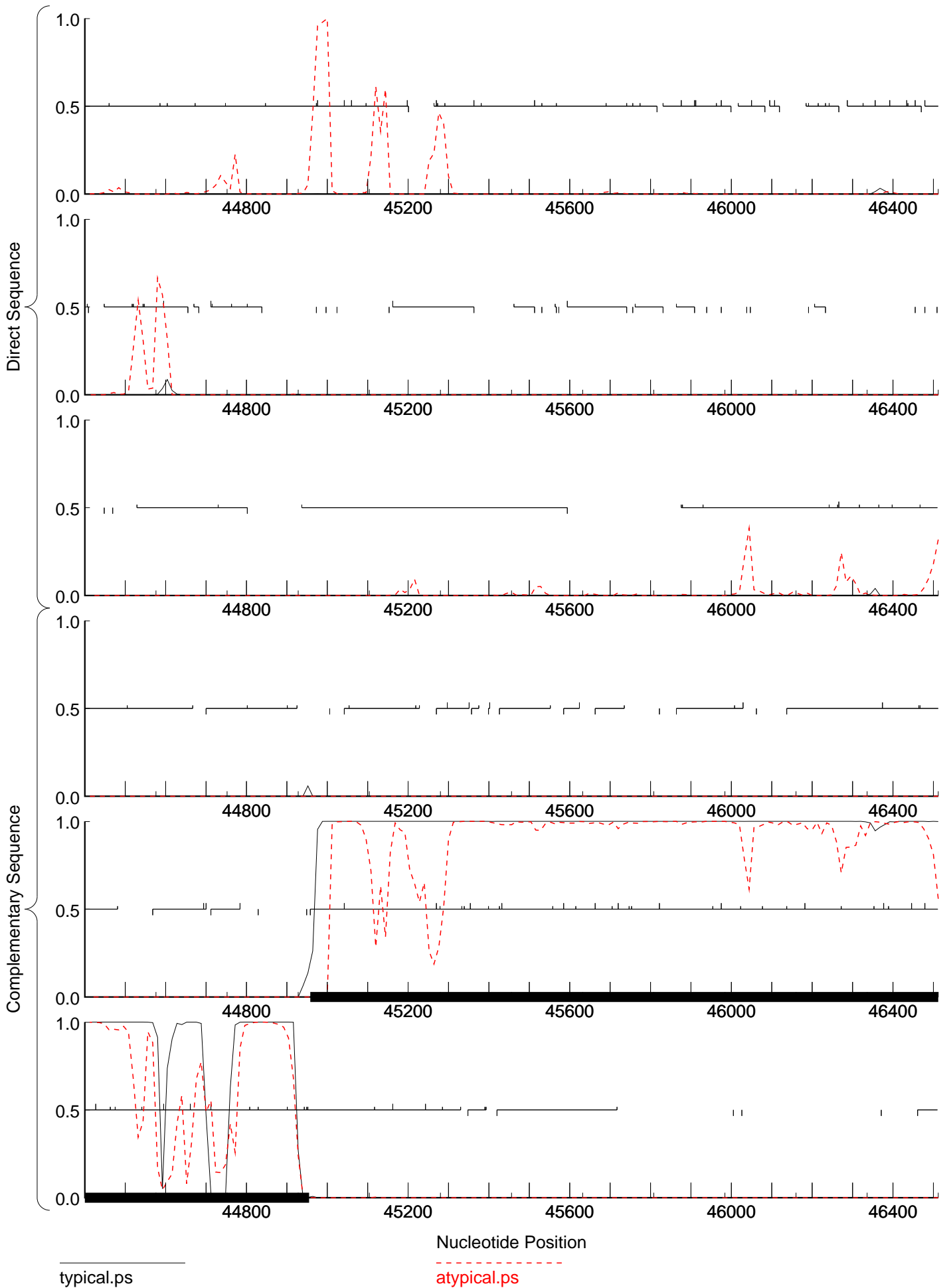


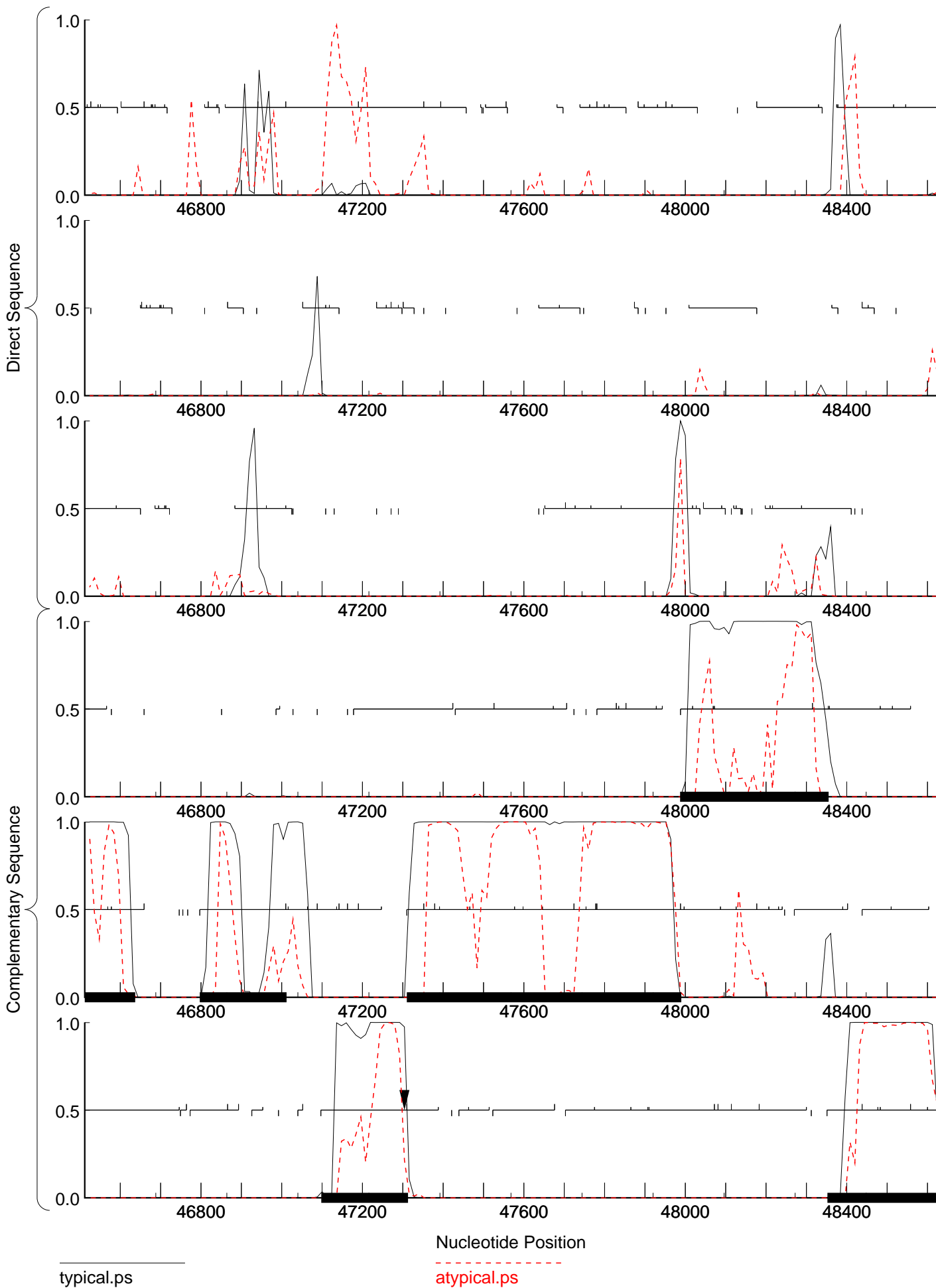
typical.ps

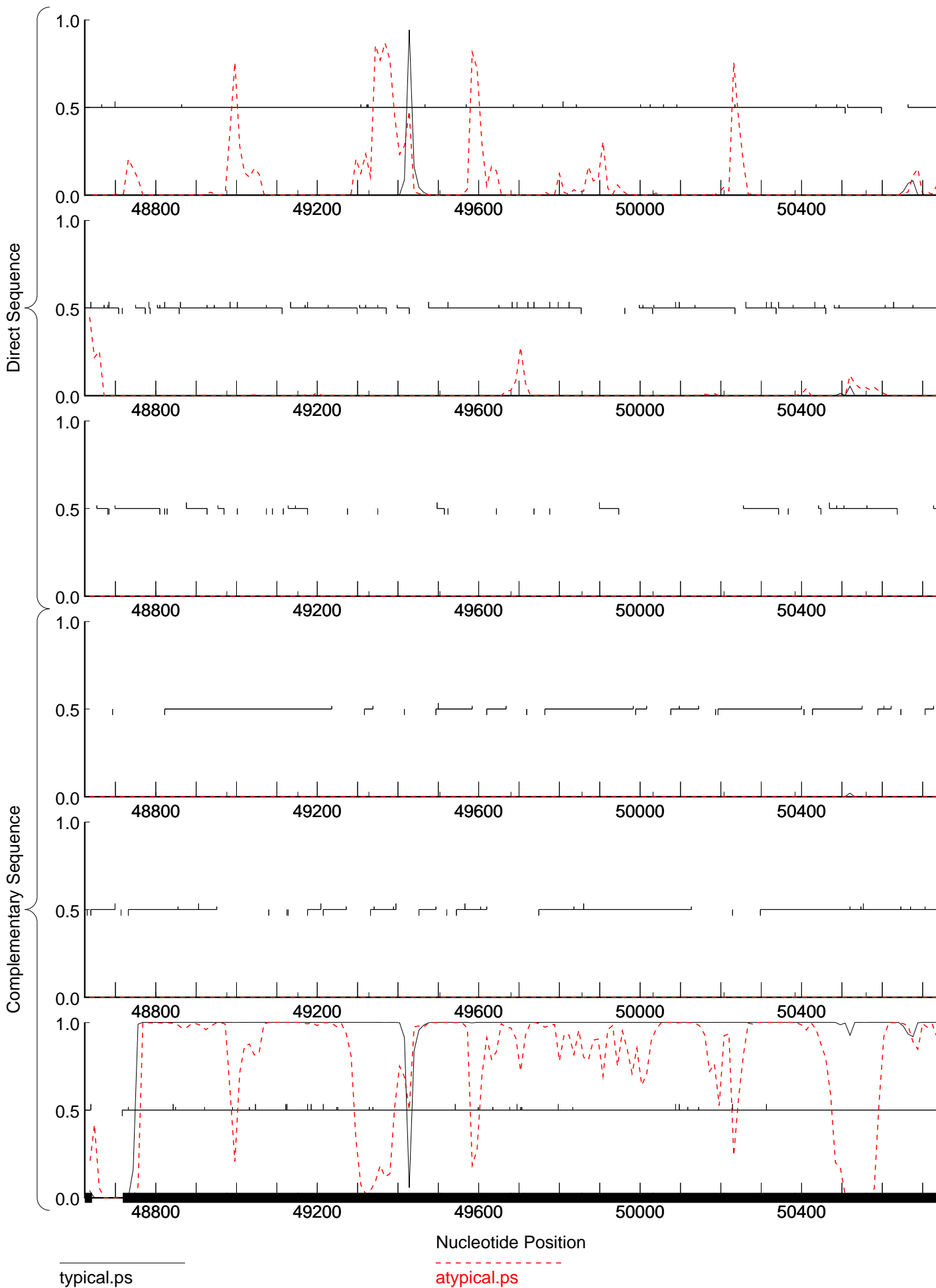
atypical.ps

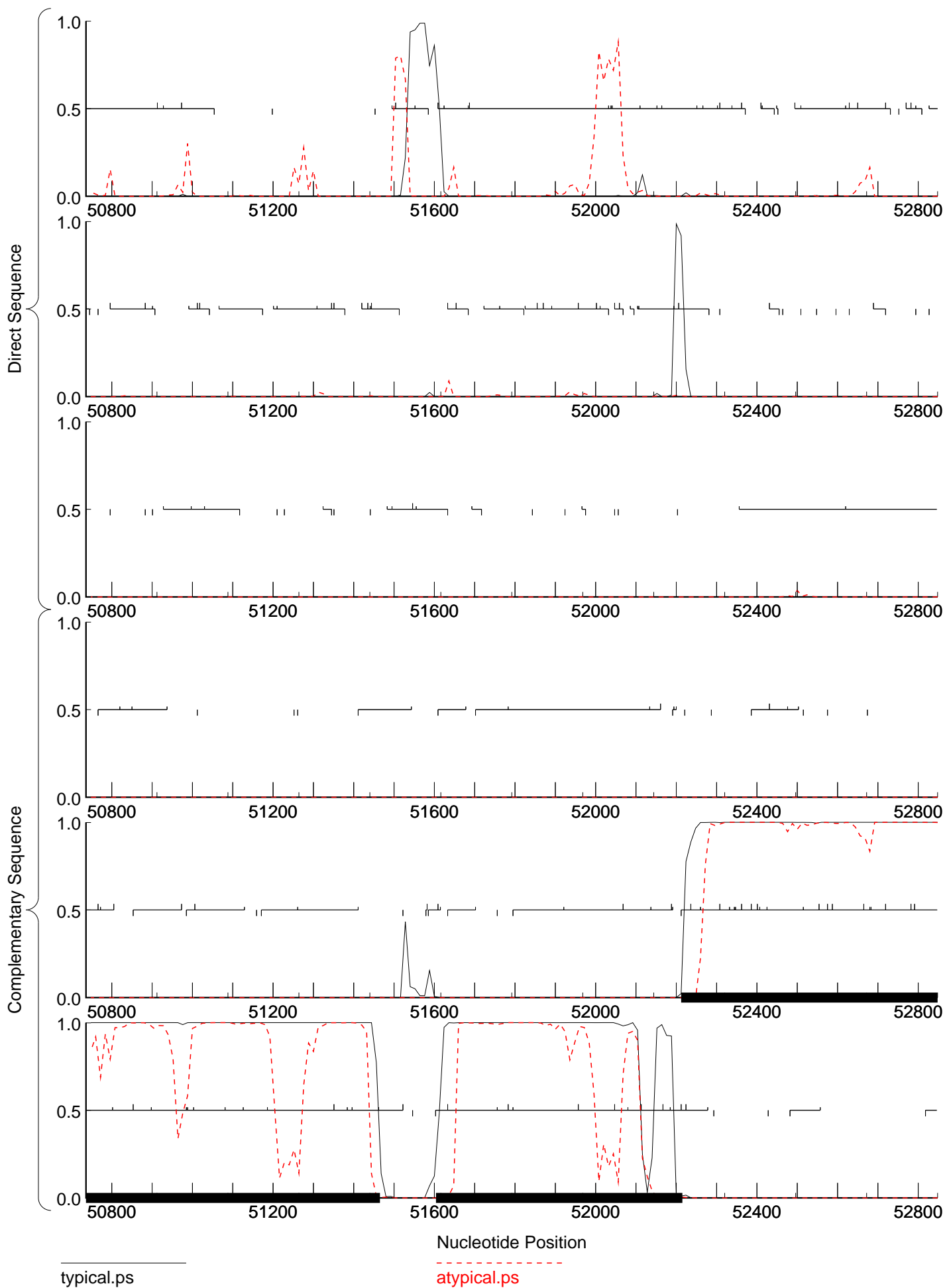






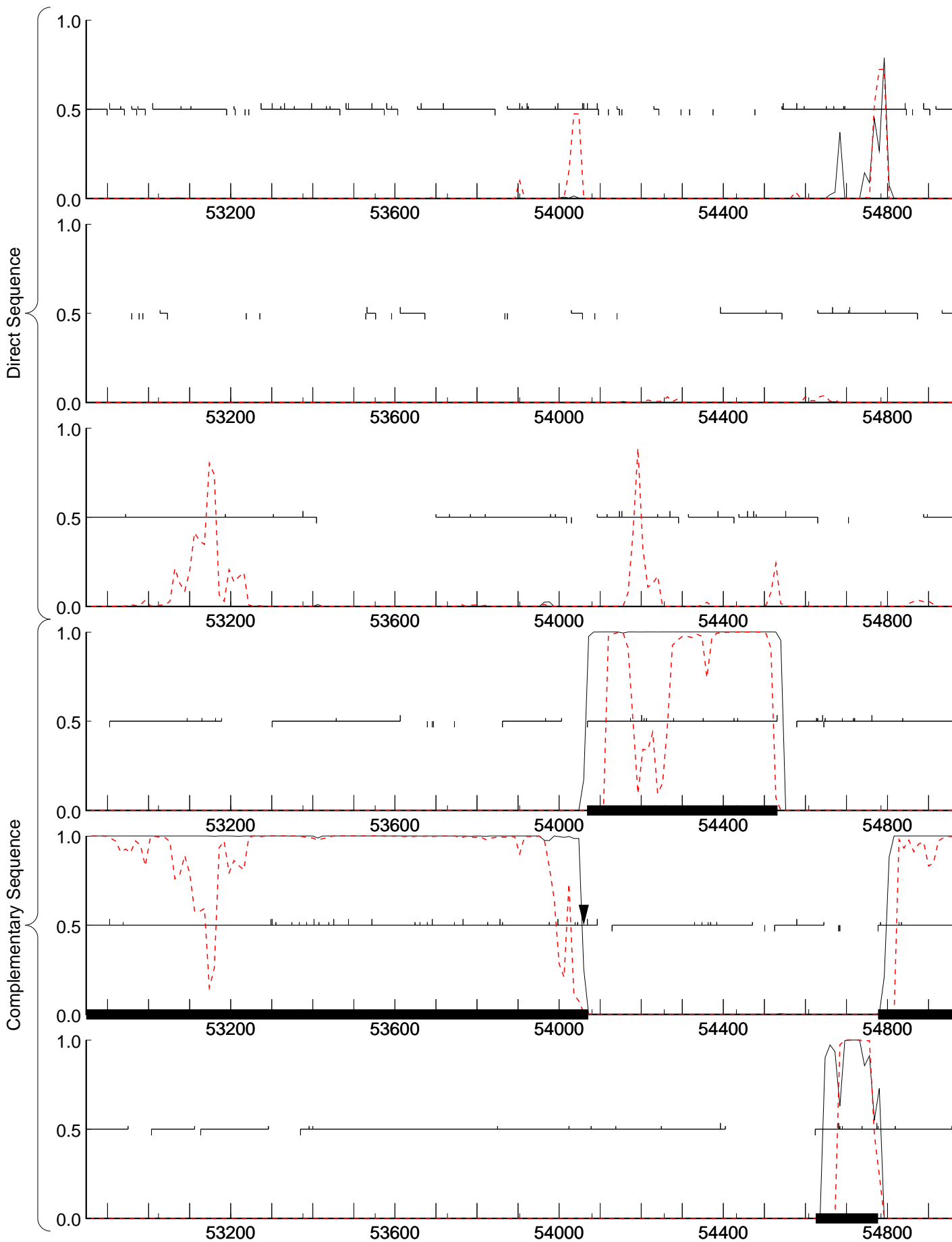






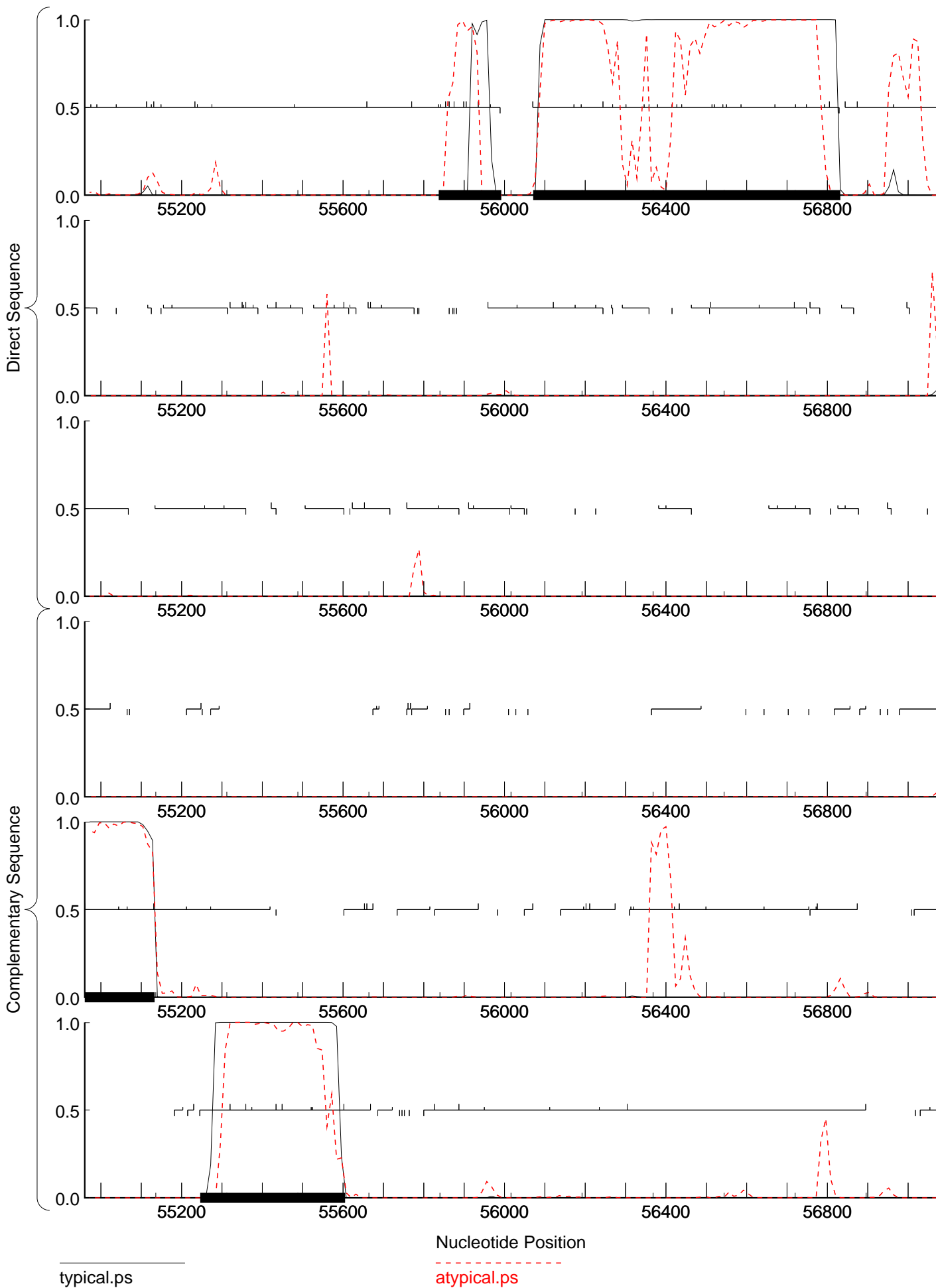
typical.ps

atypical.ps



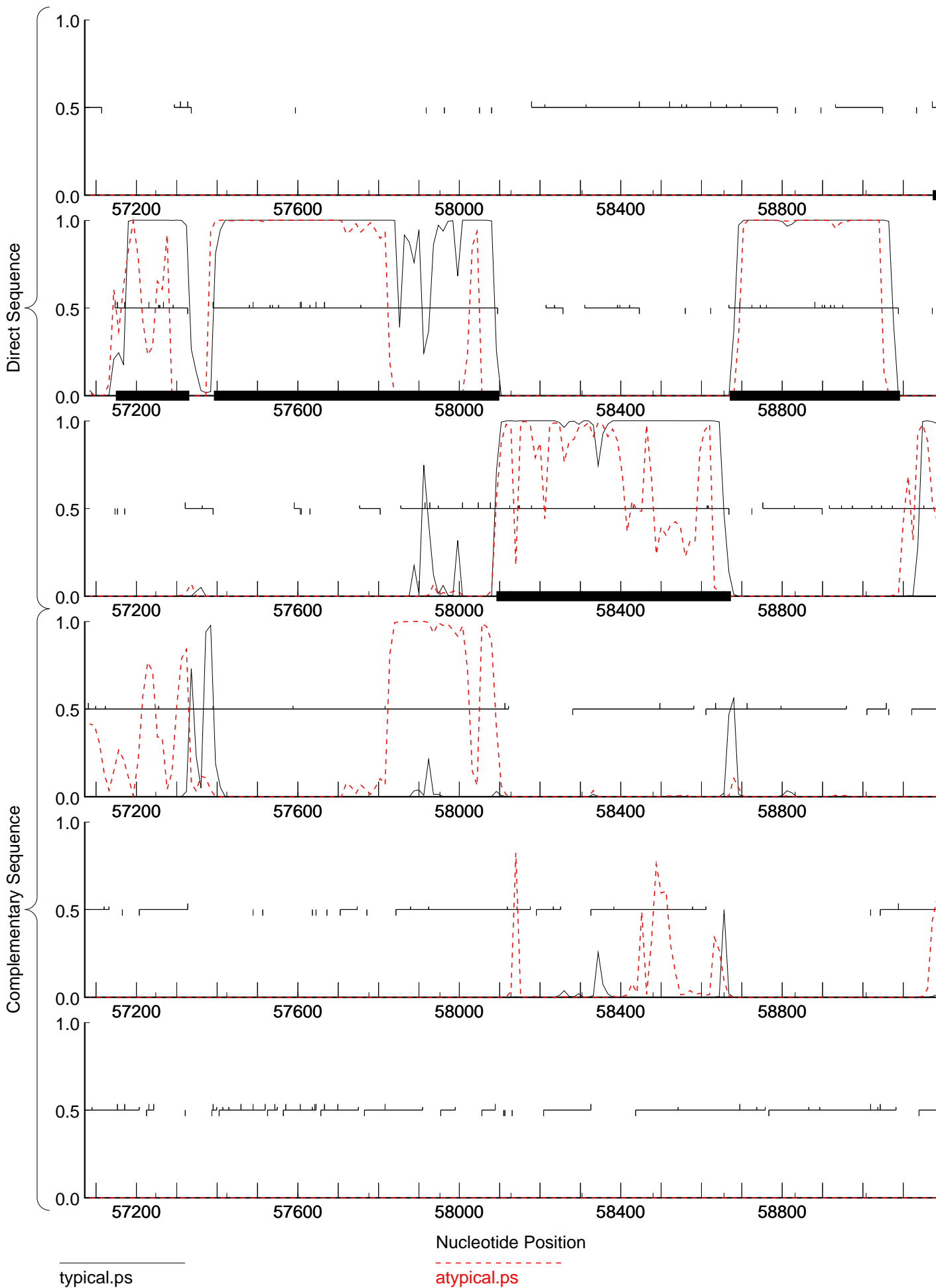
typical.ps

atypical.ps



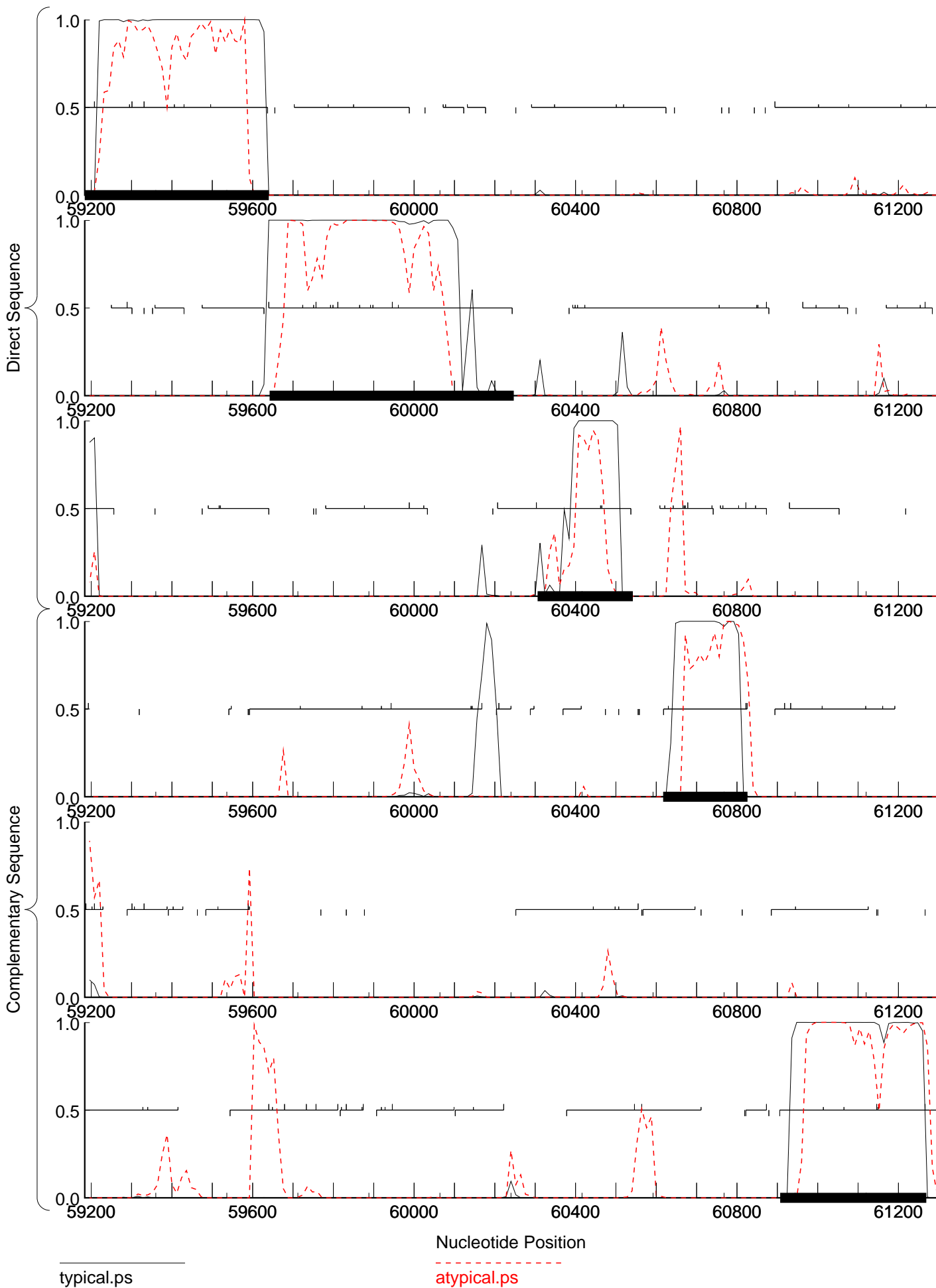
typical.ps

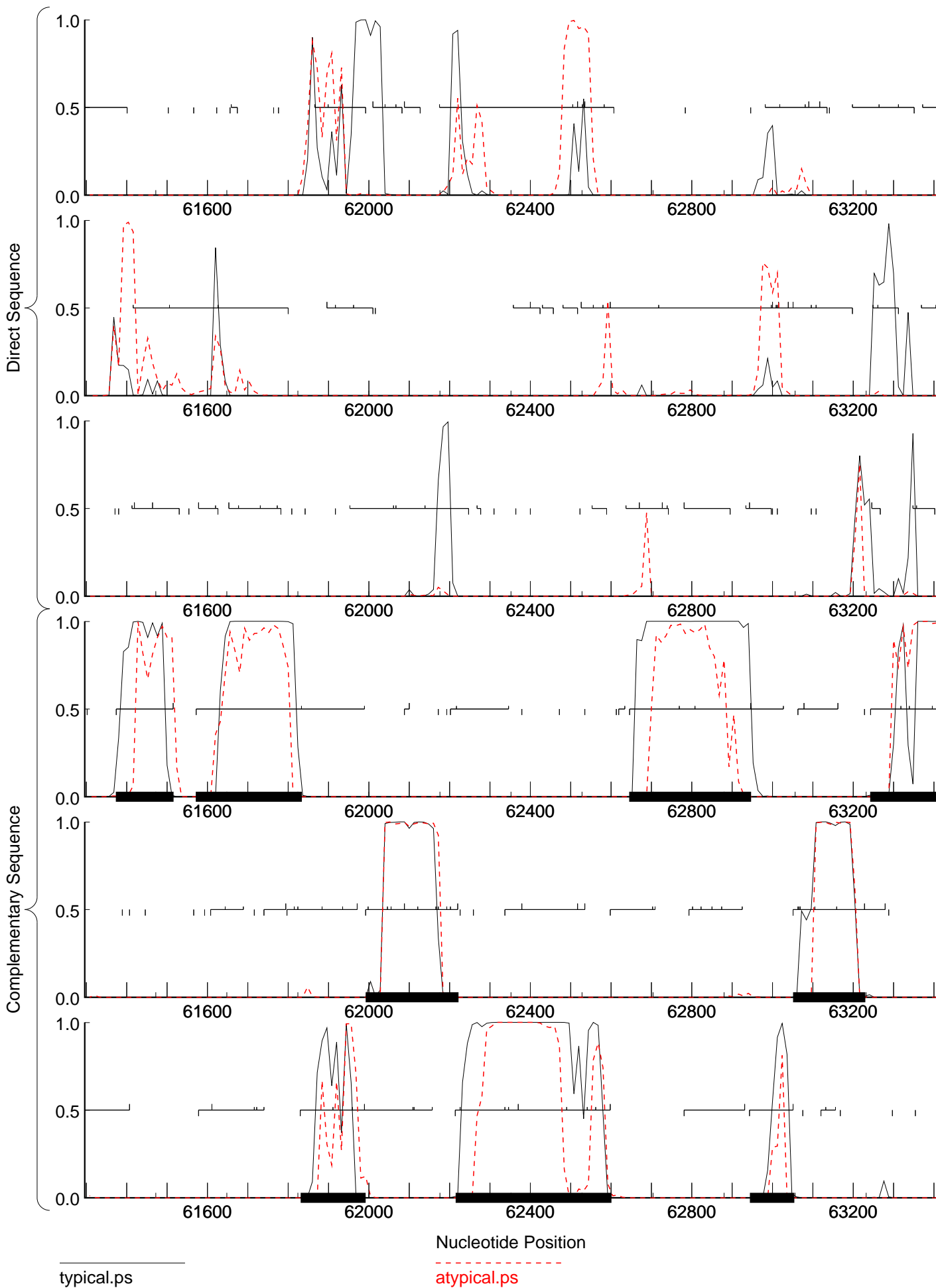
atypical.ps



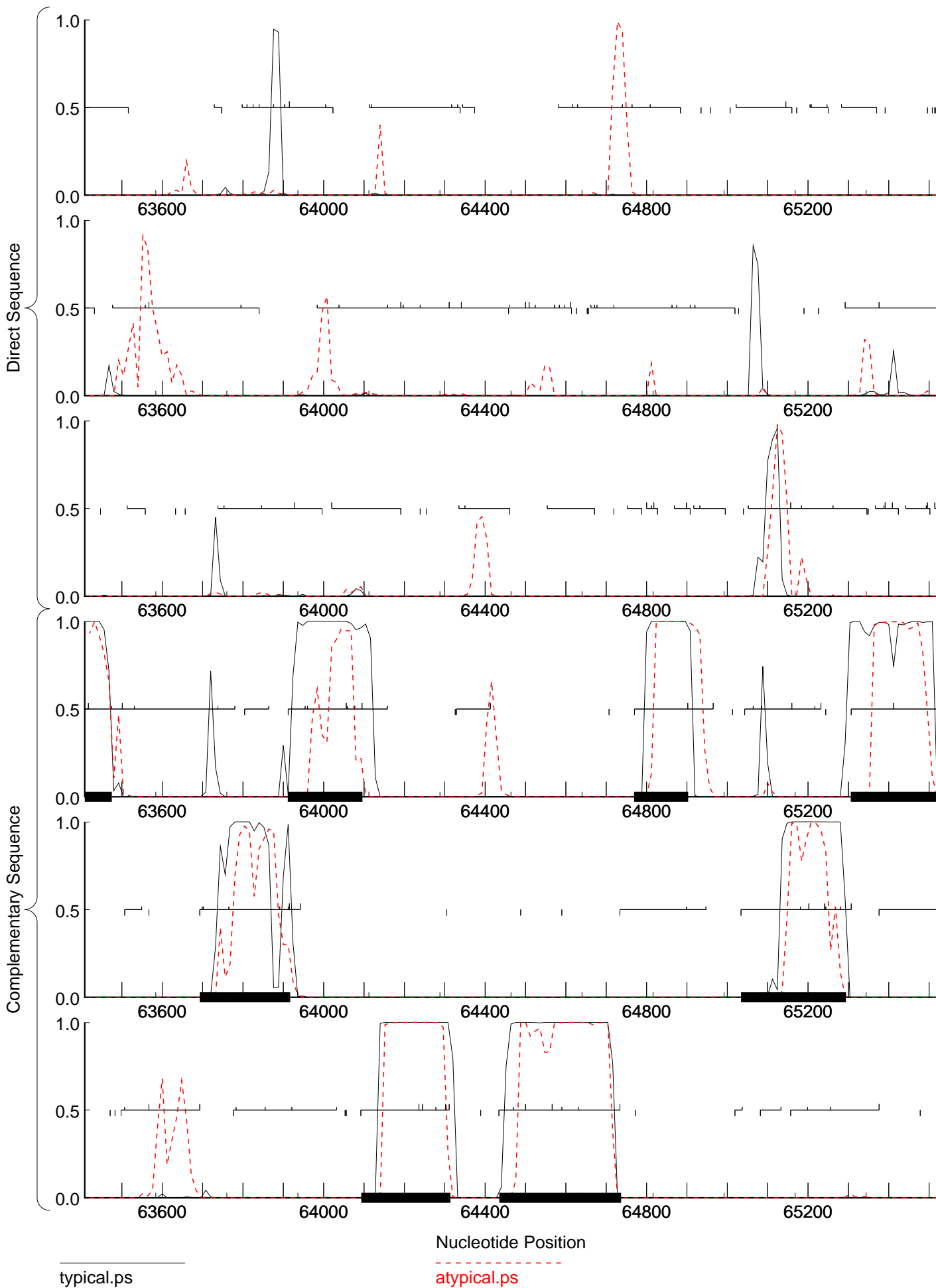
typical.ps

atypical.ps





GeneMark.hmm prediction



typical.ps

atypical.ps

