

GeneMark

Version 2.5p (09.08.06)

Copyright 1993 - M. Borodovsky, J. McIninch

PROGRAM INFORMATION

Sequence : Mycobacterium phage ZygoTaiga complete sequence, 157204 bp, circularly permuted, Cluster C1
Analysis Date : 2/23/18 at 17:13:02
Pages : 76
Sequence Length : 157204 bp
GC Content : 64.69%

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,

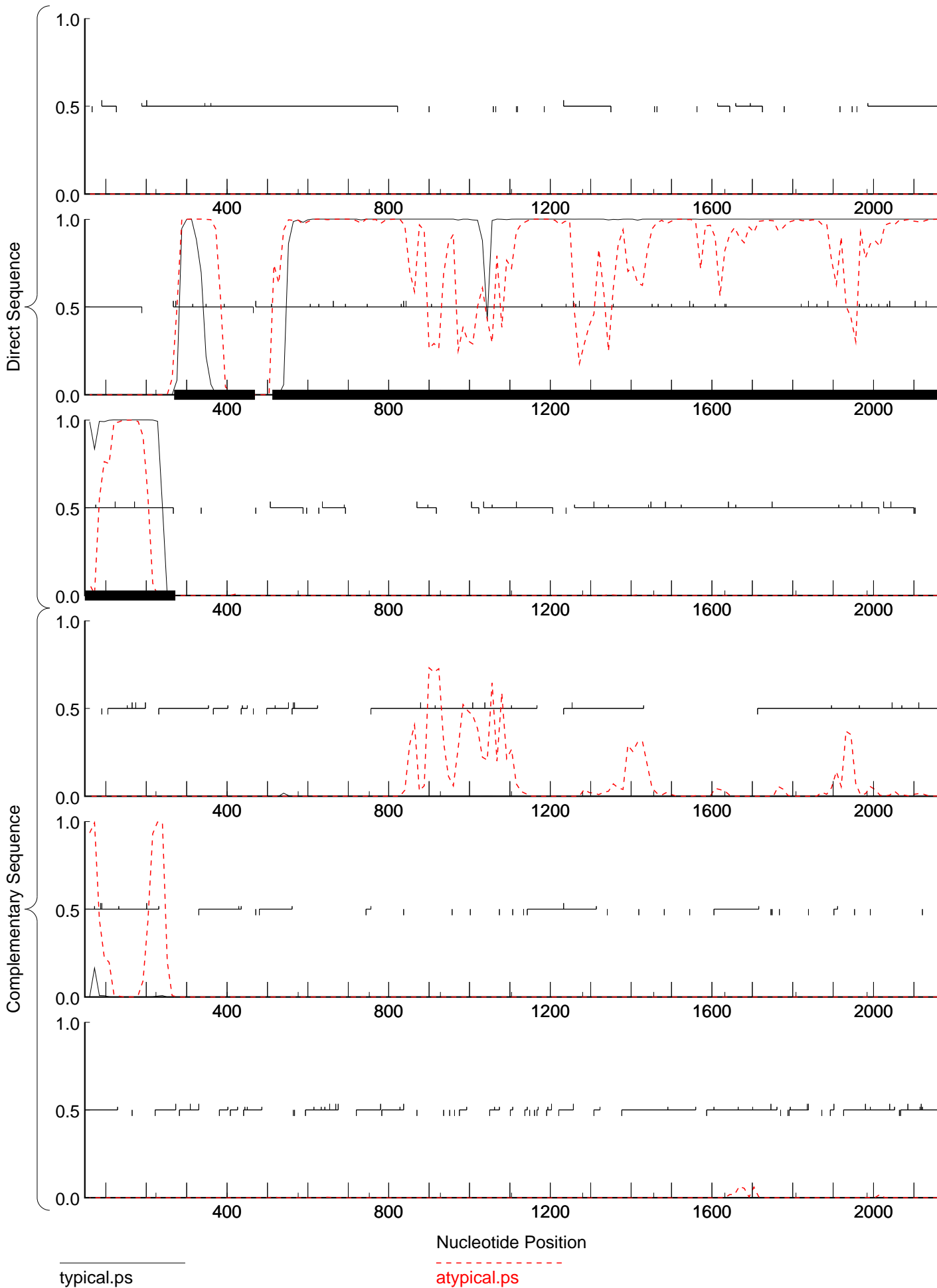
MATRIX INFORMATION

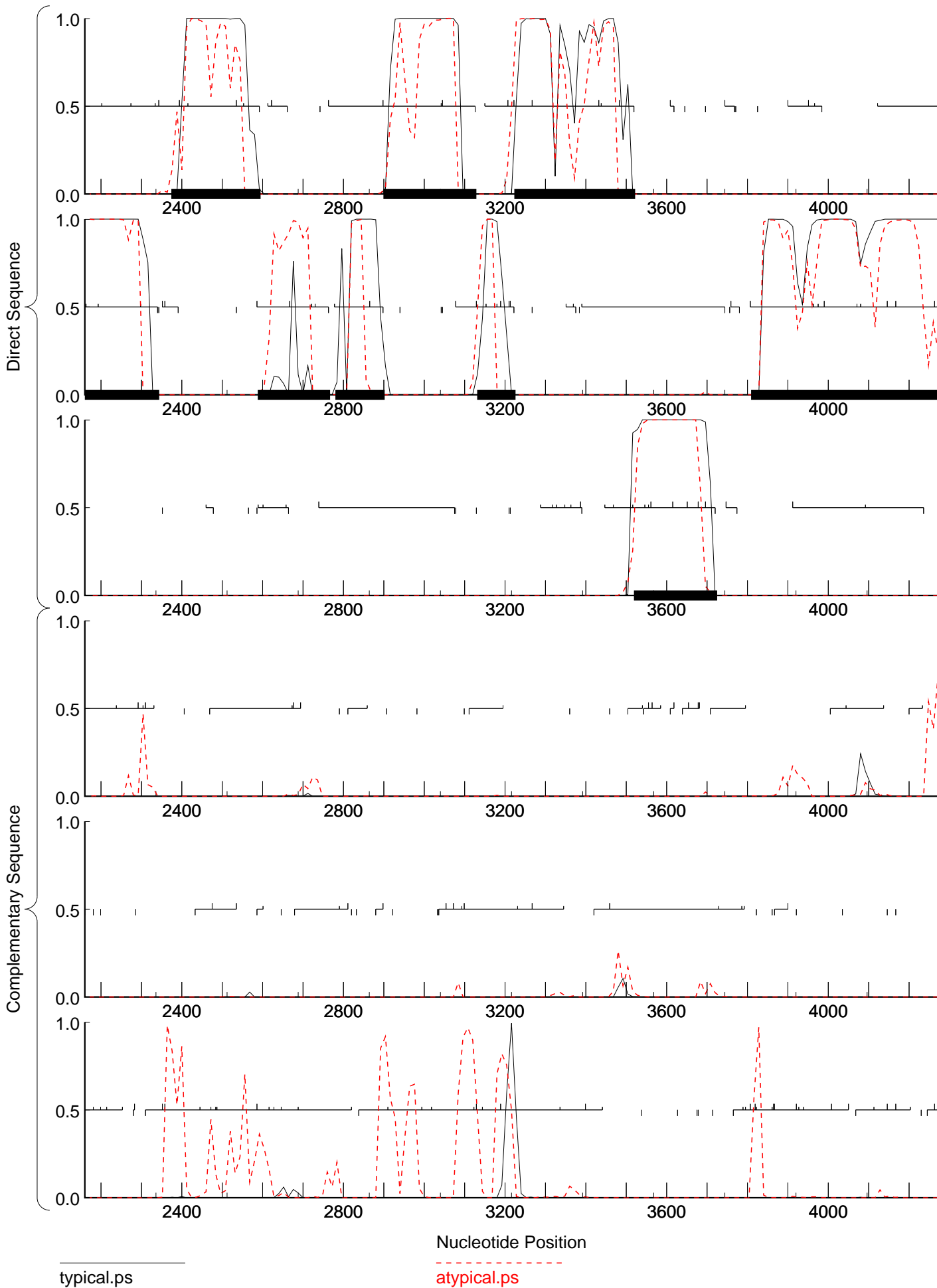
Matrix : -
Author : -
Order : 4

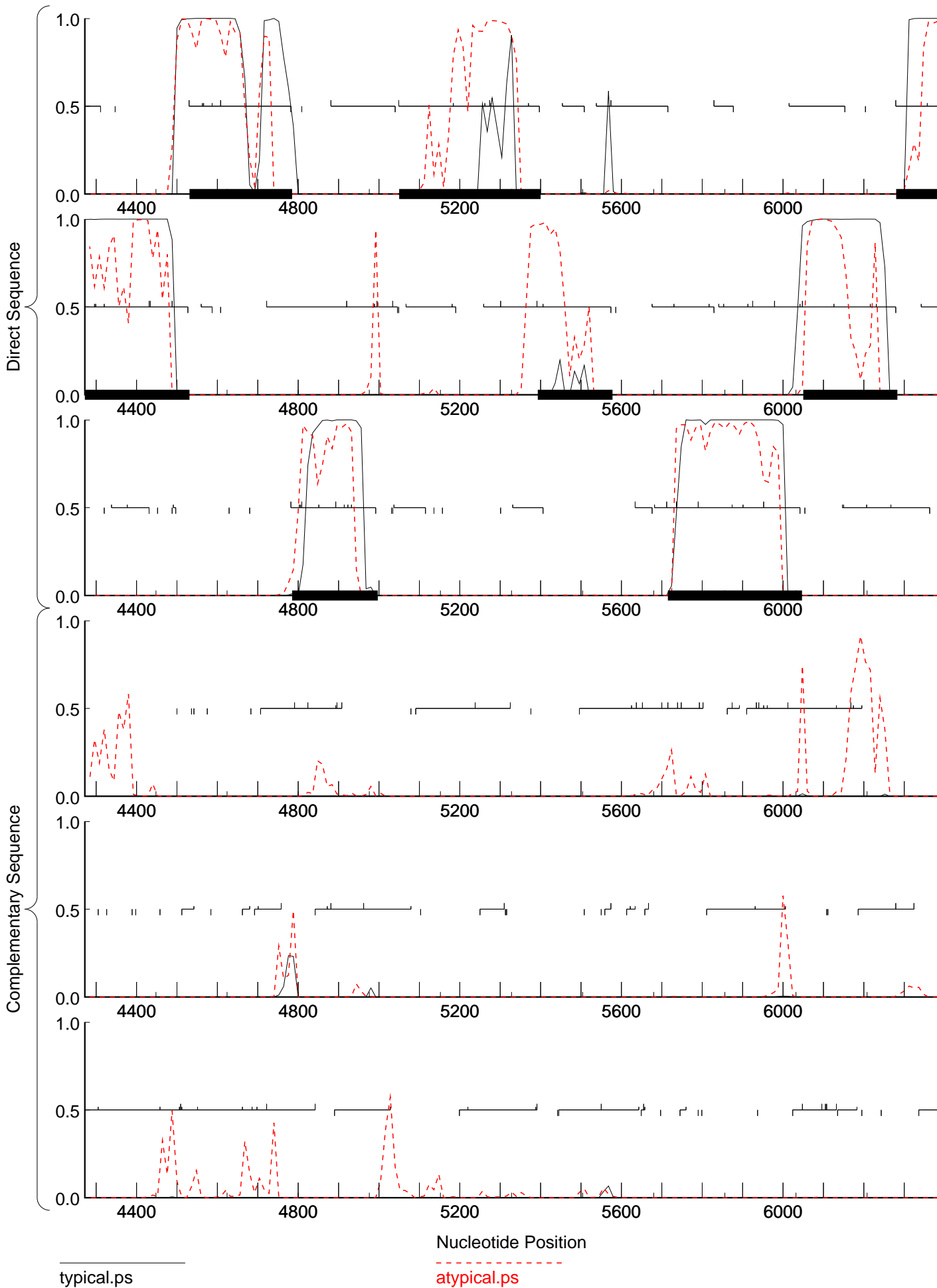
Send questions / comments to:
Dr. M. Borodovsky
Georgia Institute of Technology
School of Biology
Atlanta, GA 30332-0230

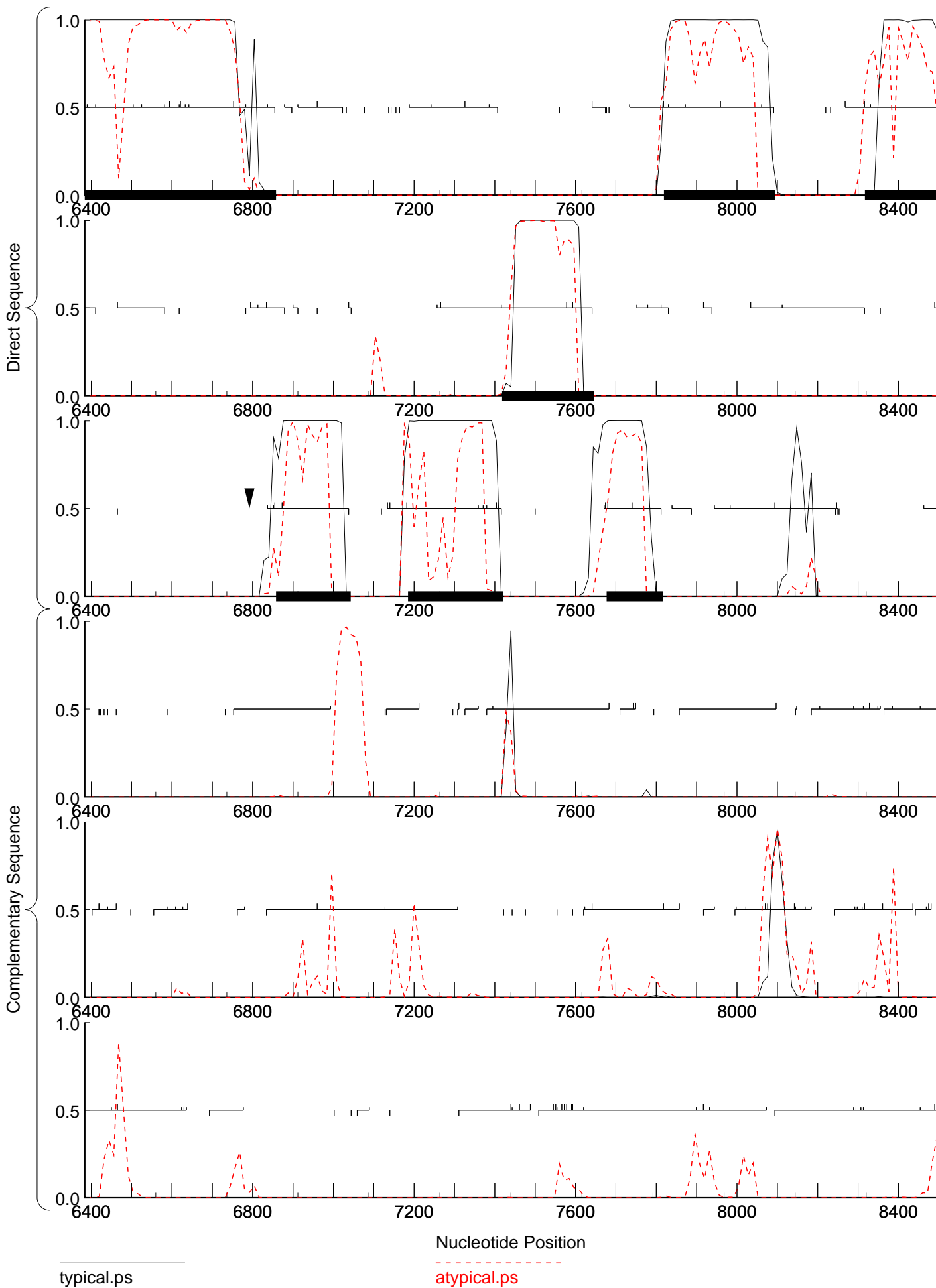
Matrix notes & comments

Training set derived by GeneMarkS, 4.30 October 2014
Fri Feb 23 17:13:02 2018



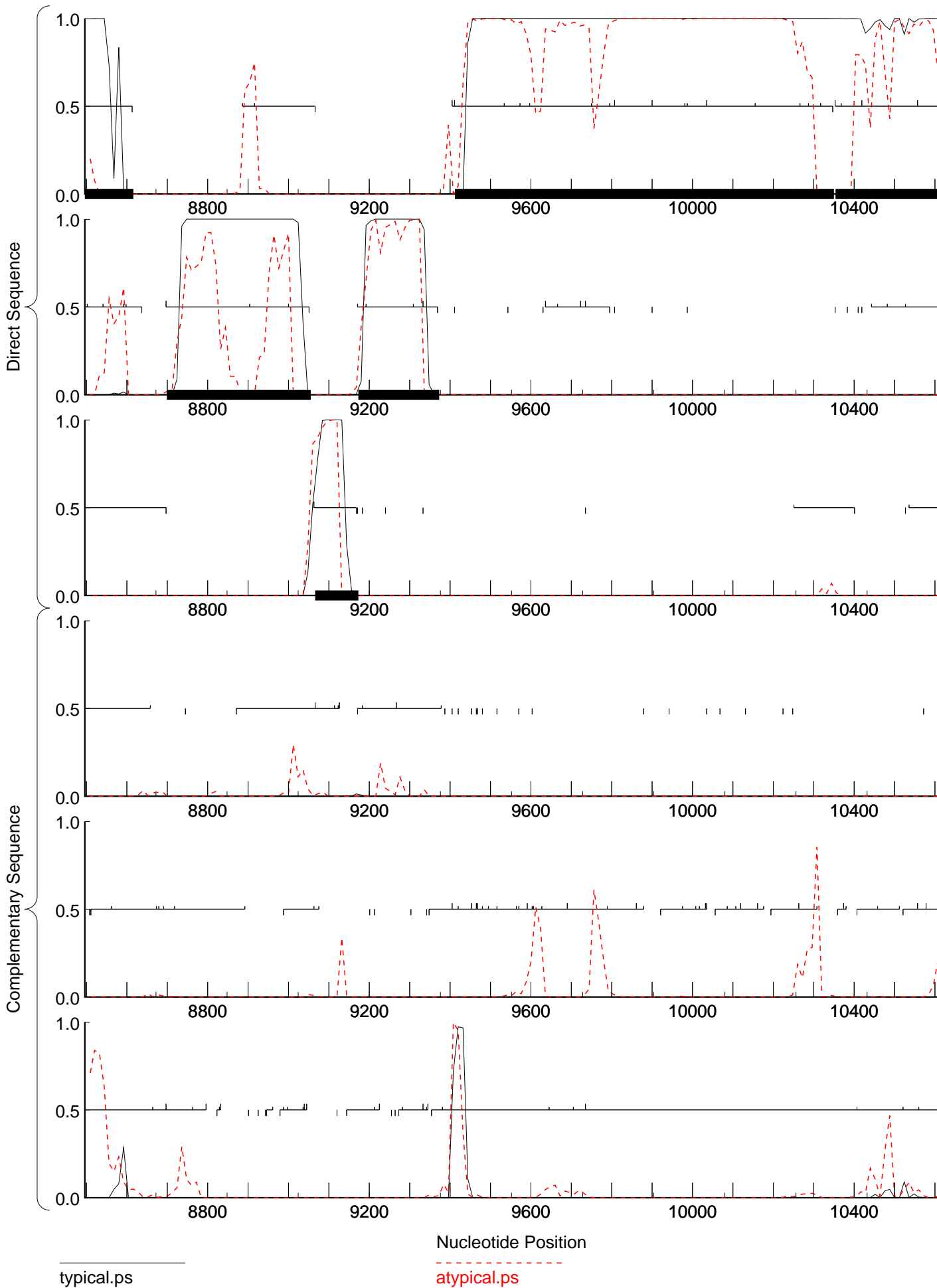


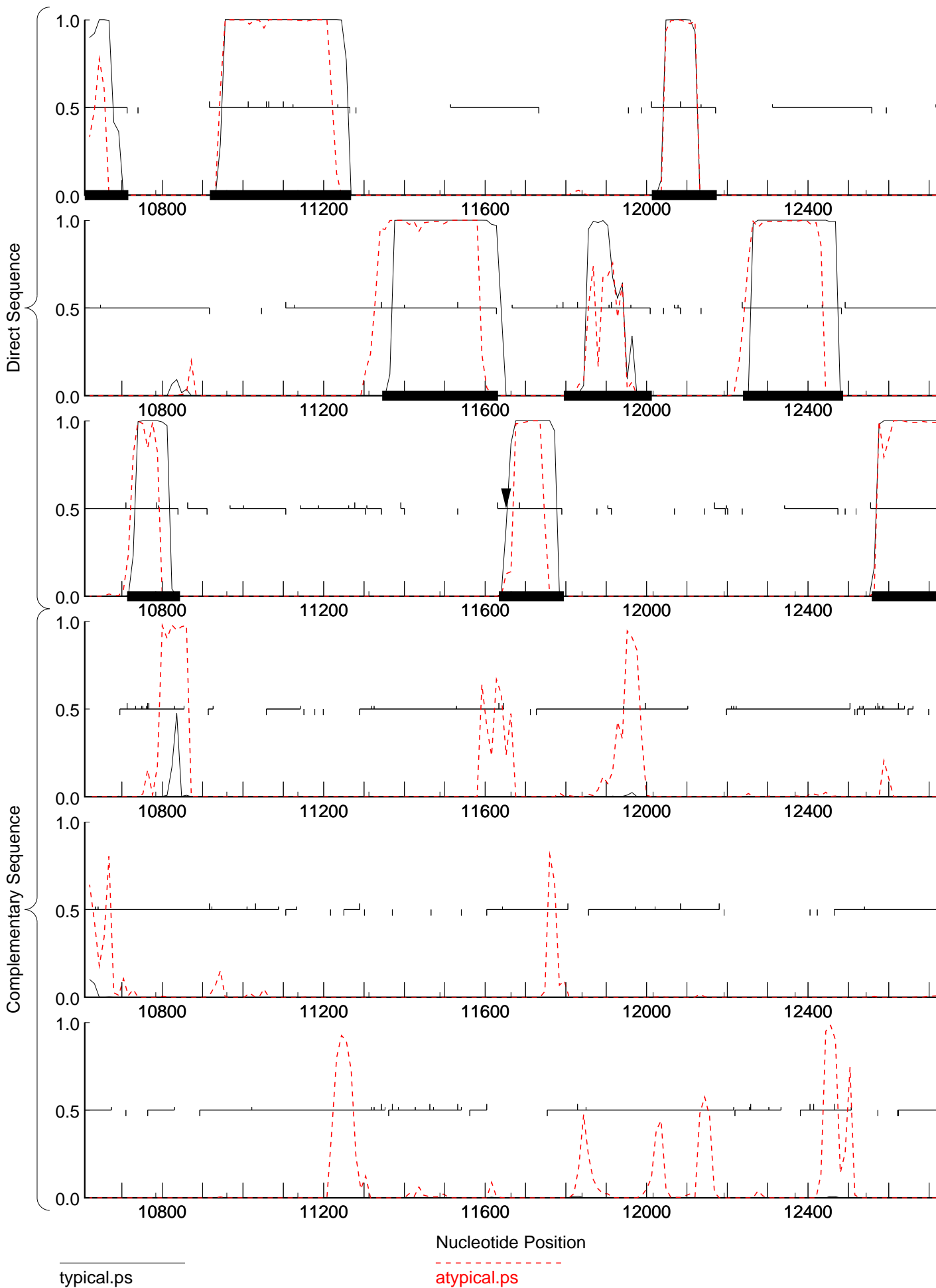


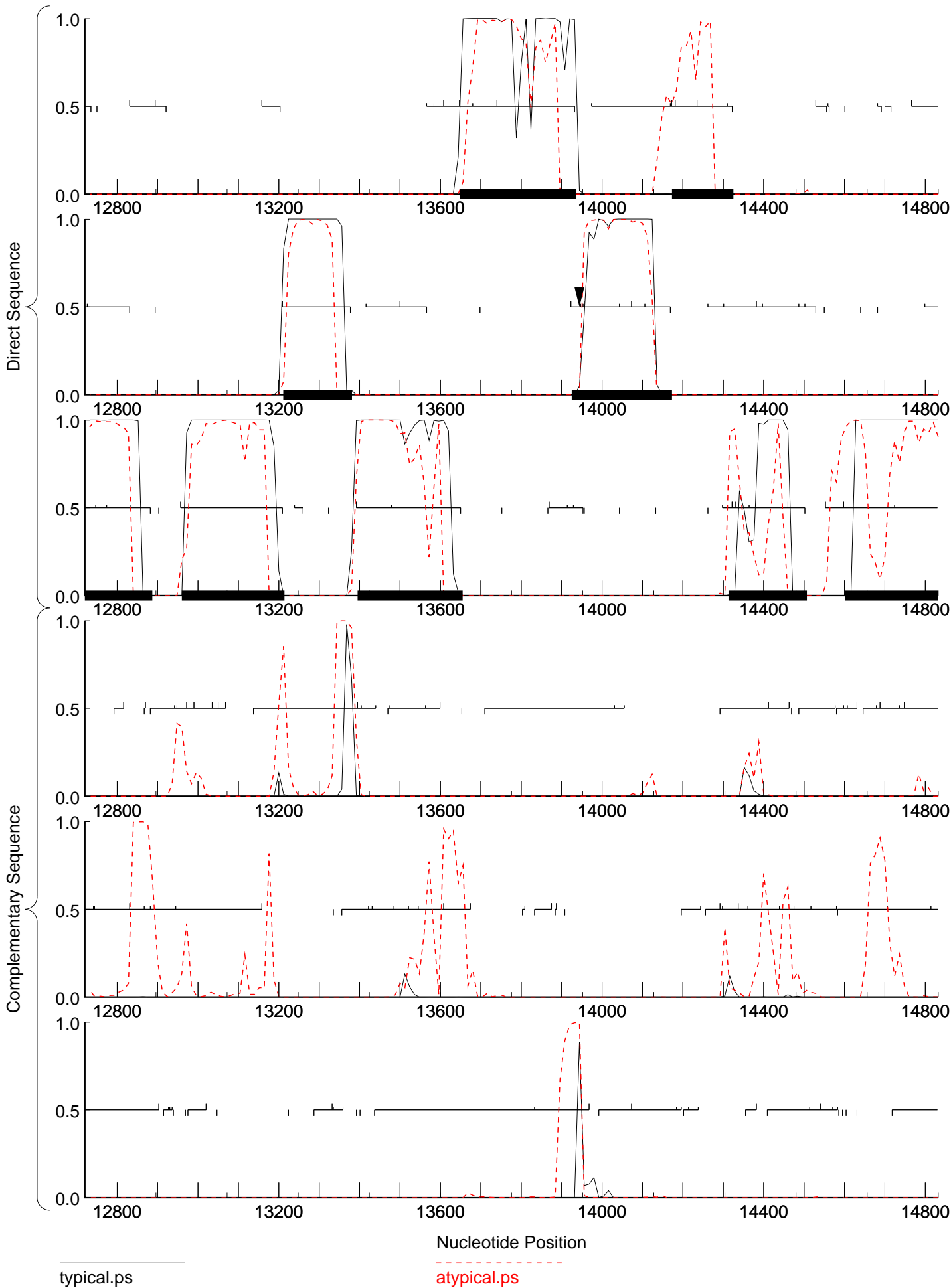


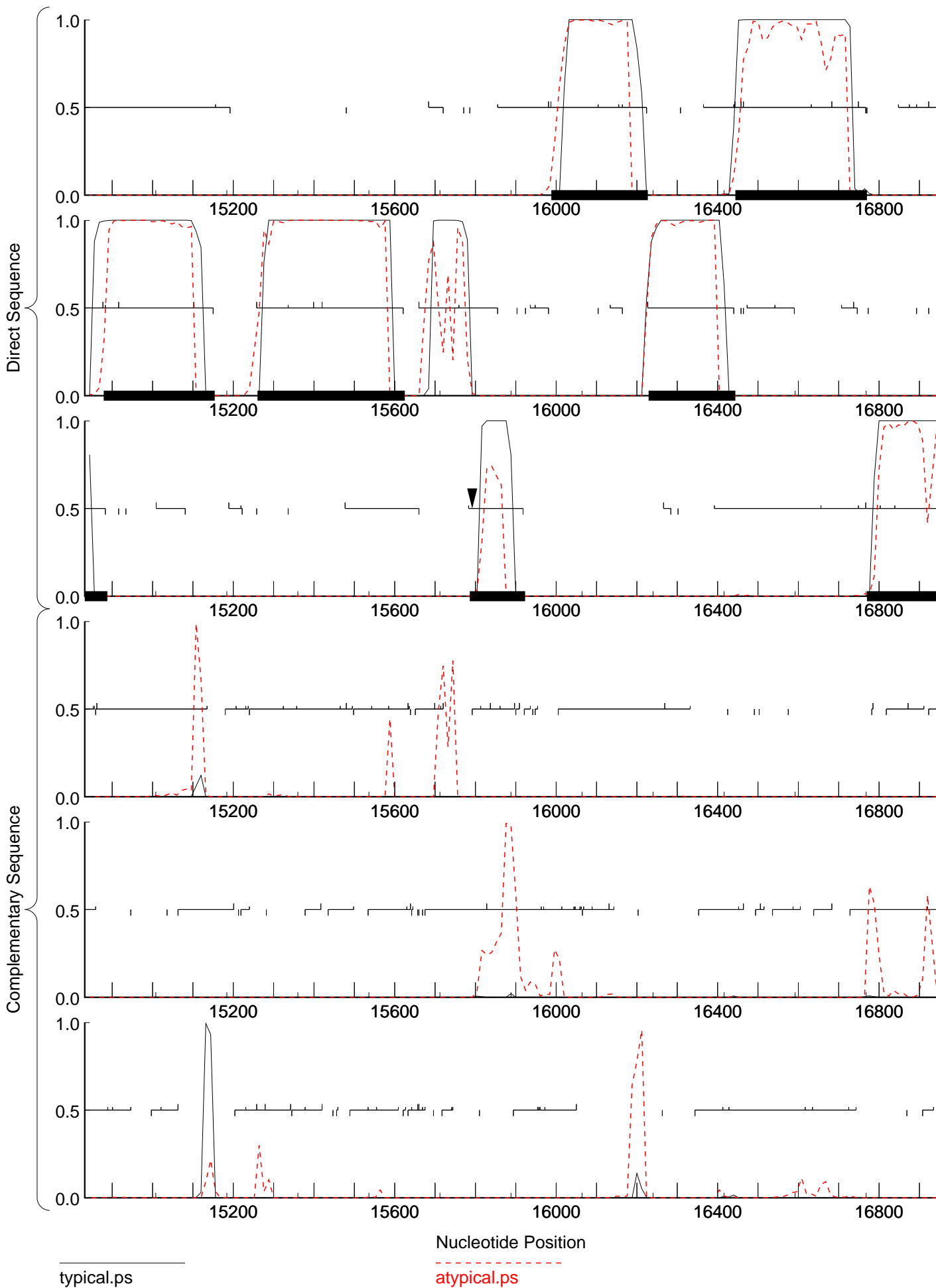
typical.ps

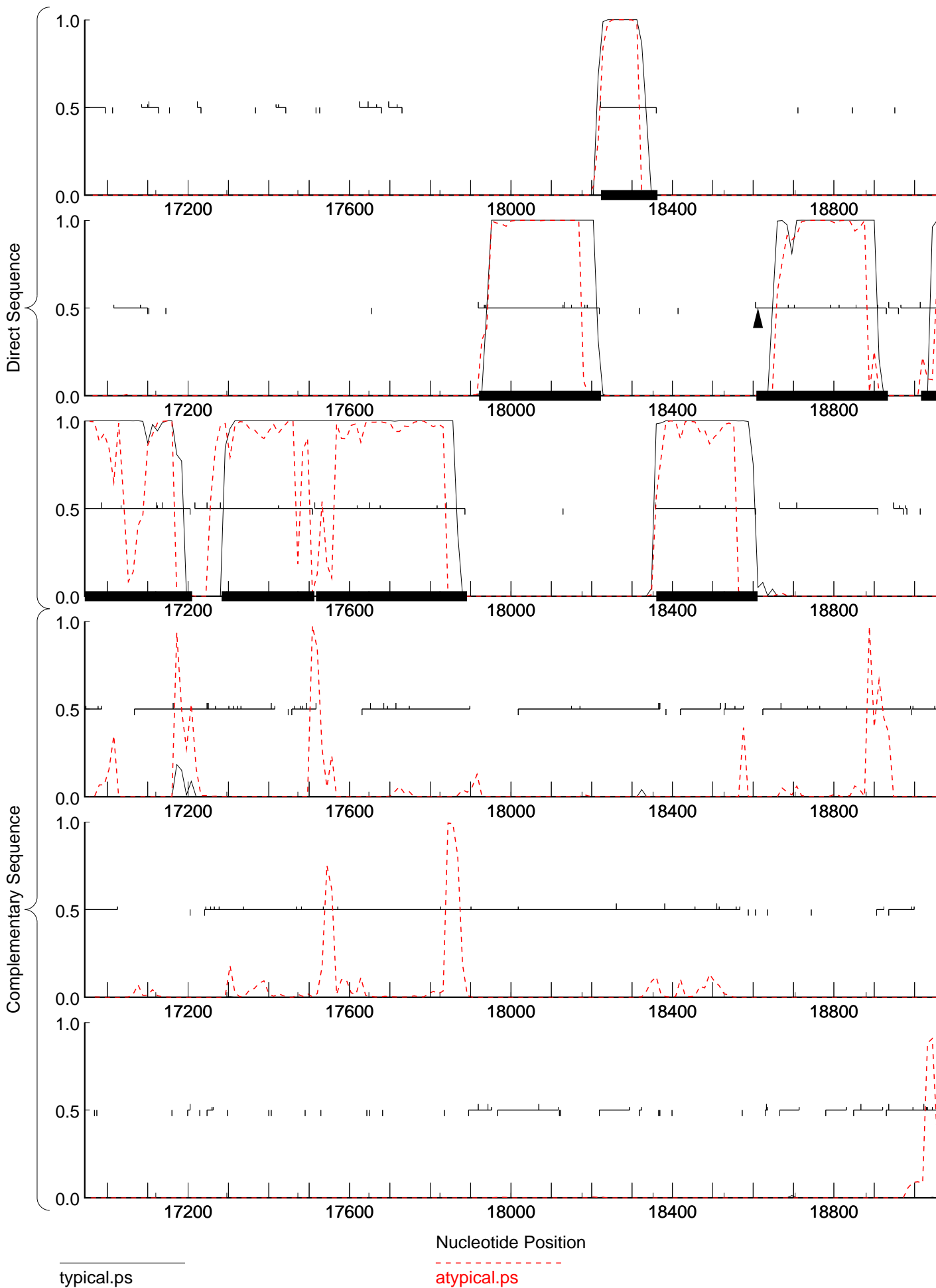
atypical.ps

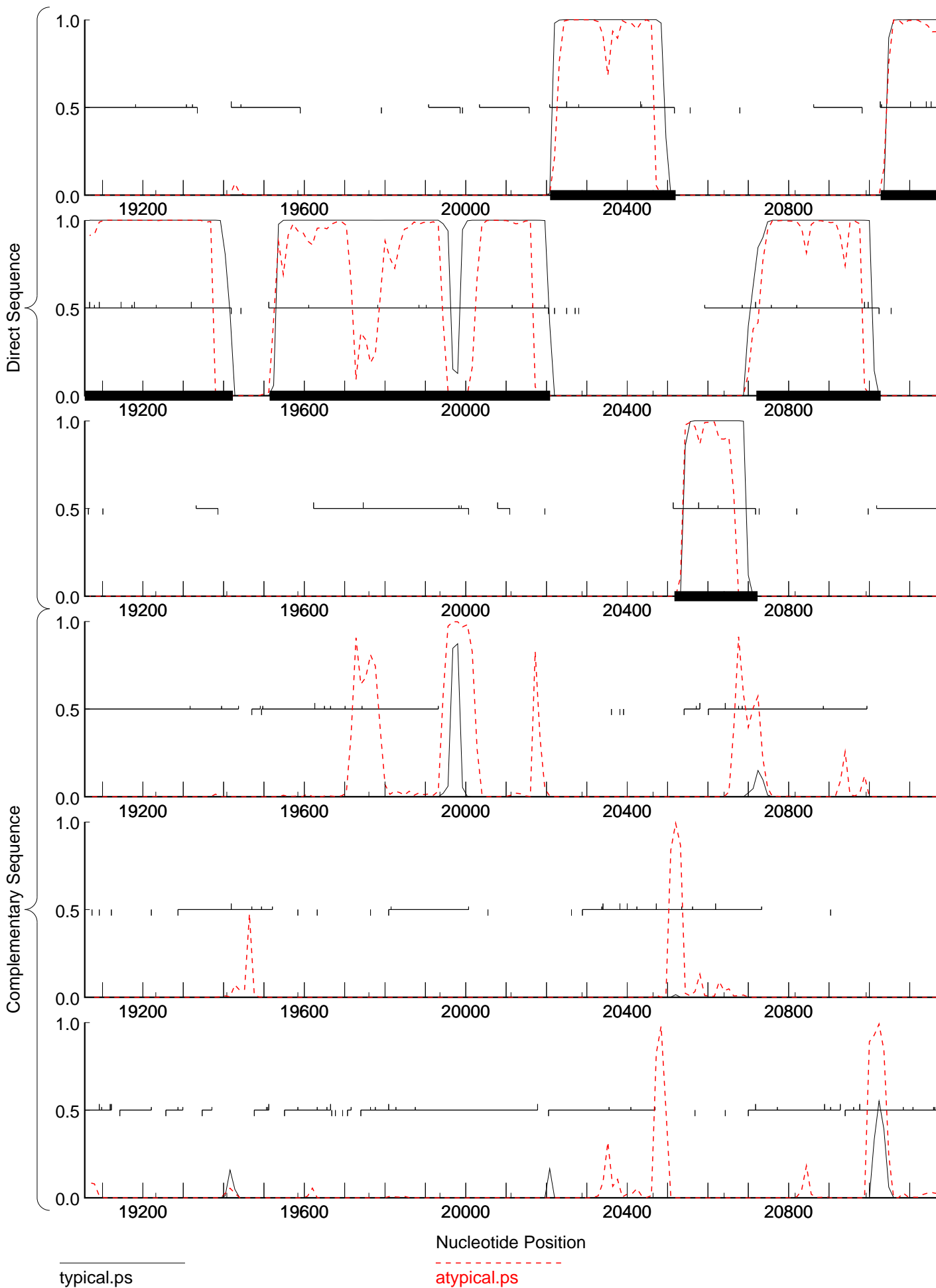


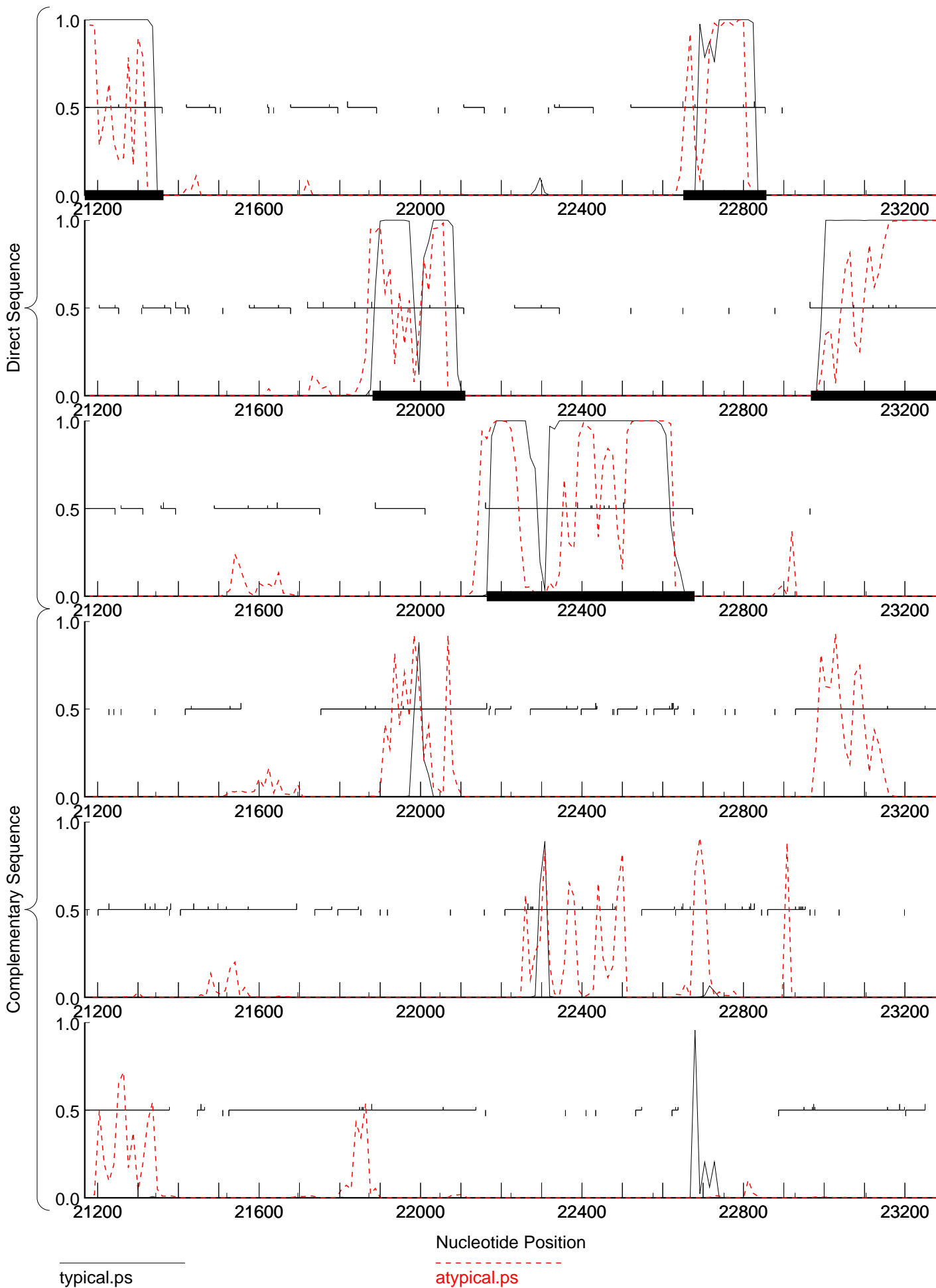


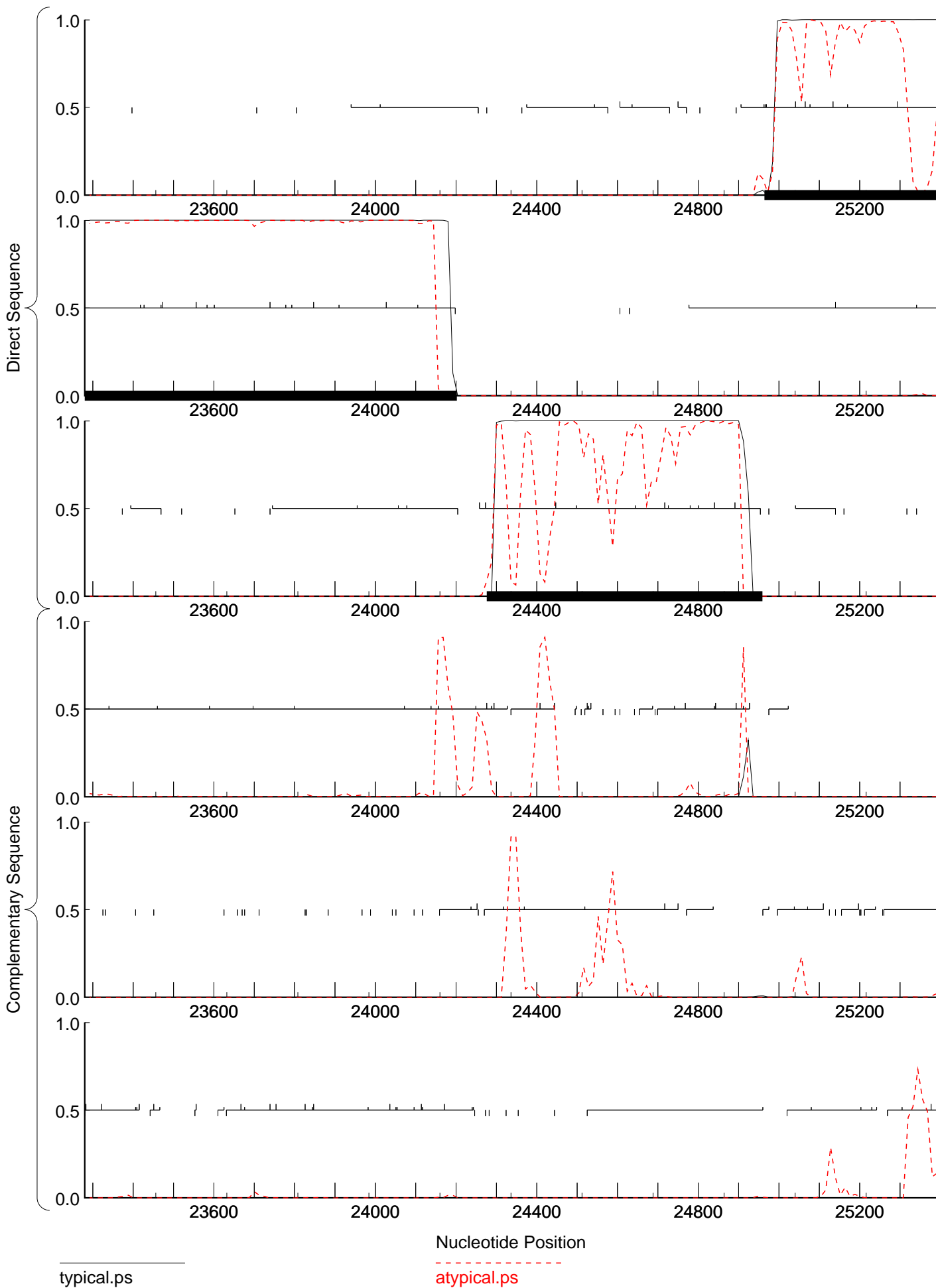






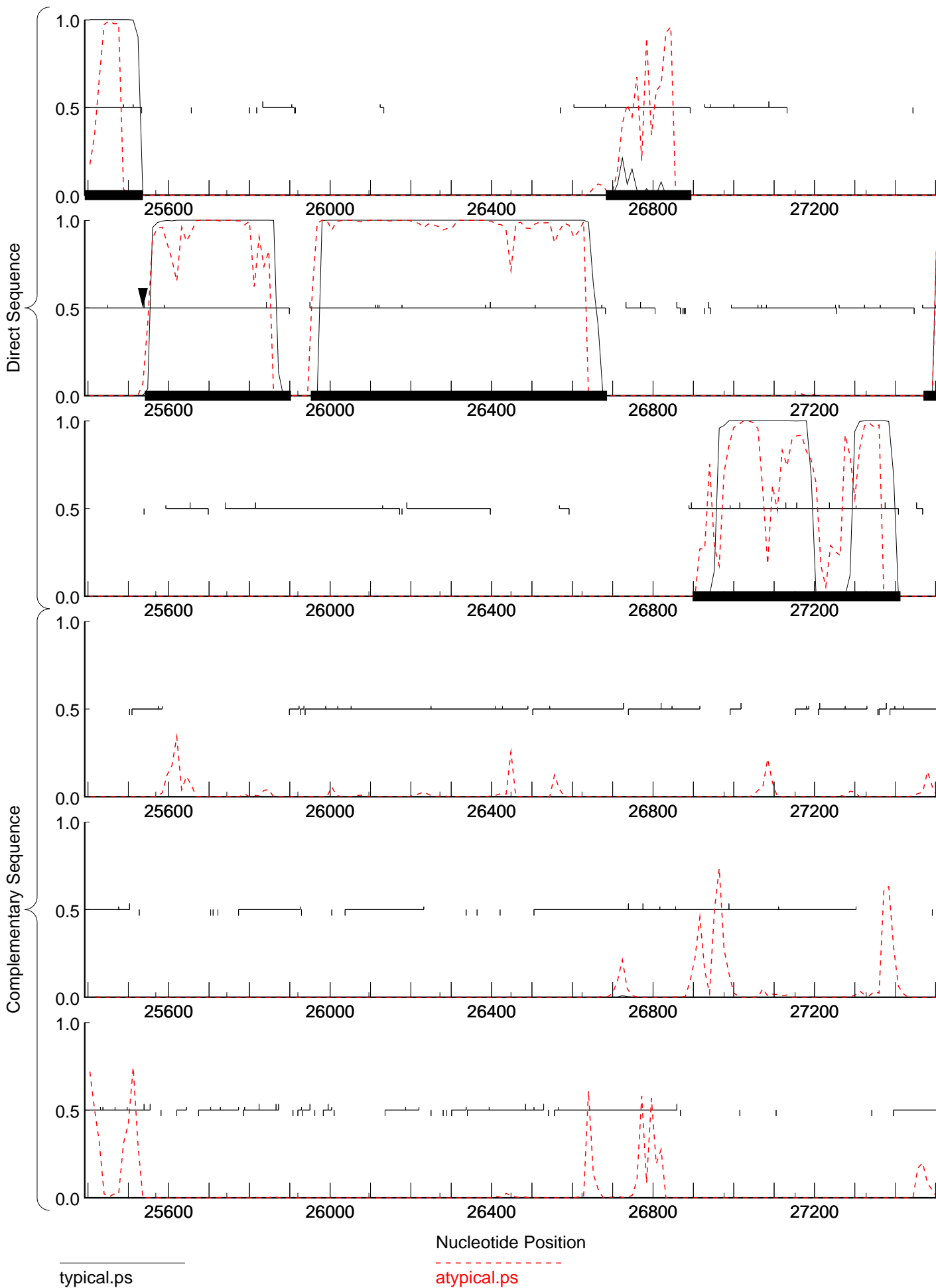


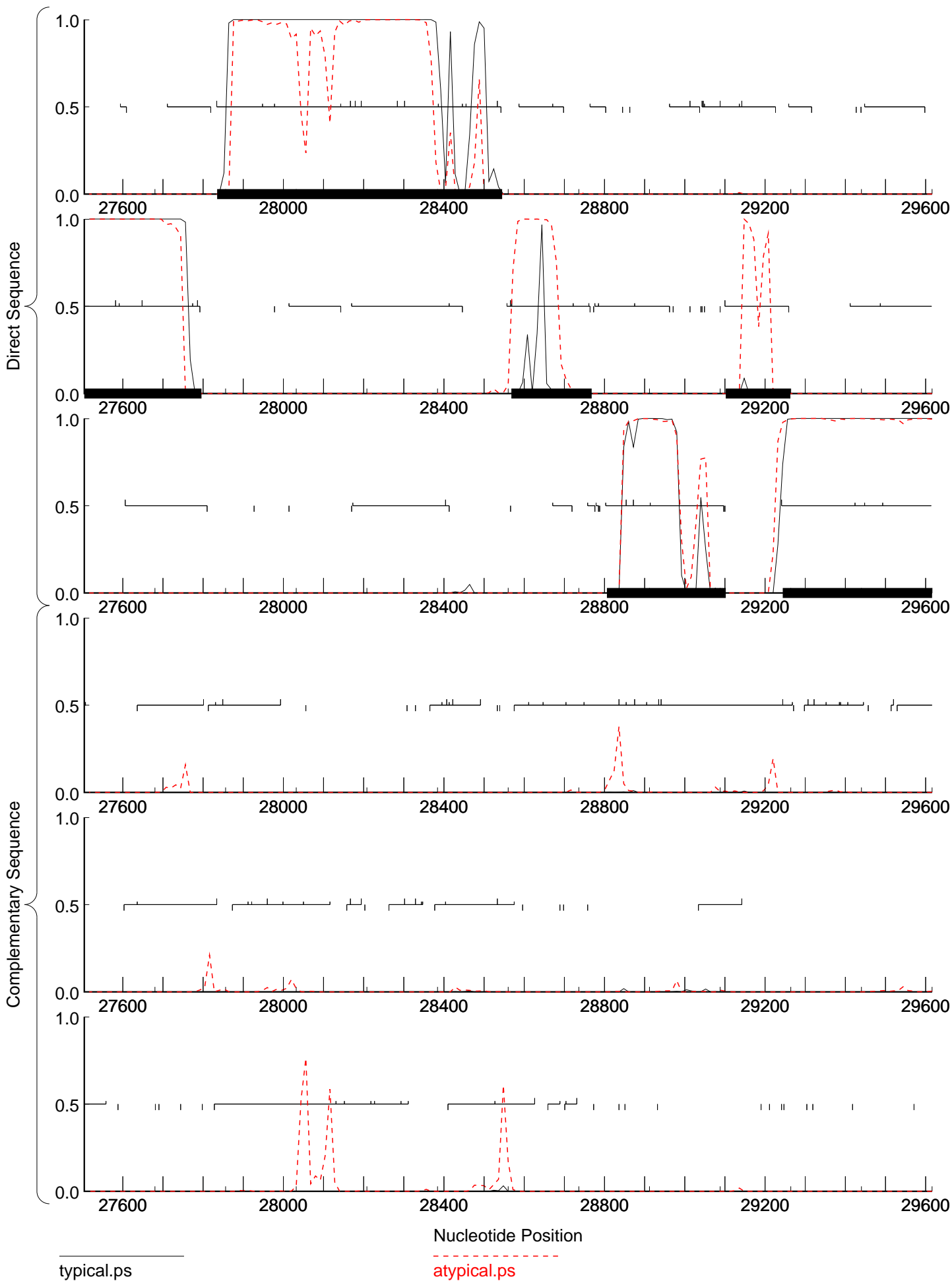


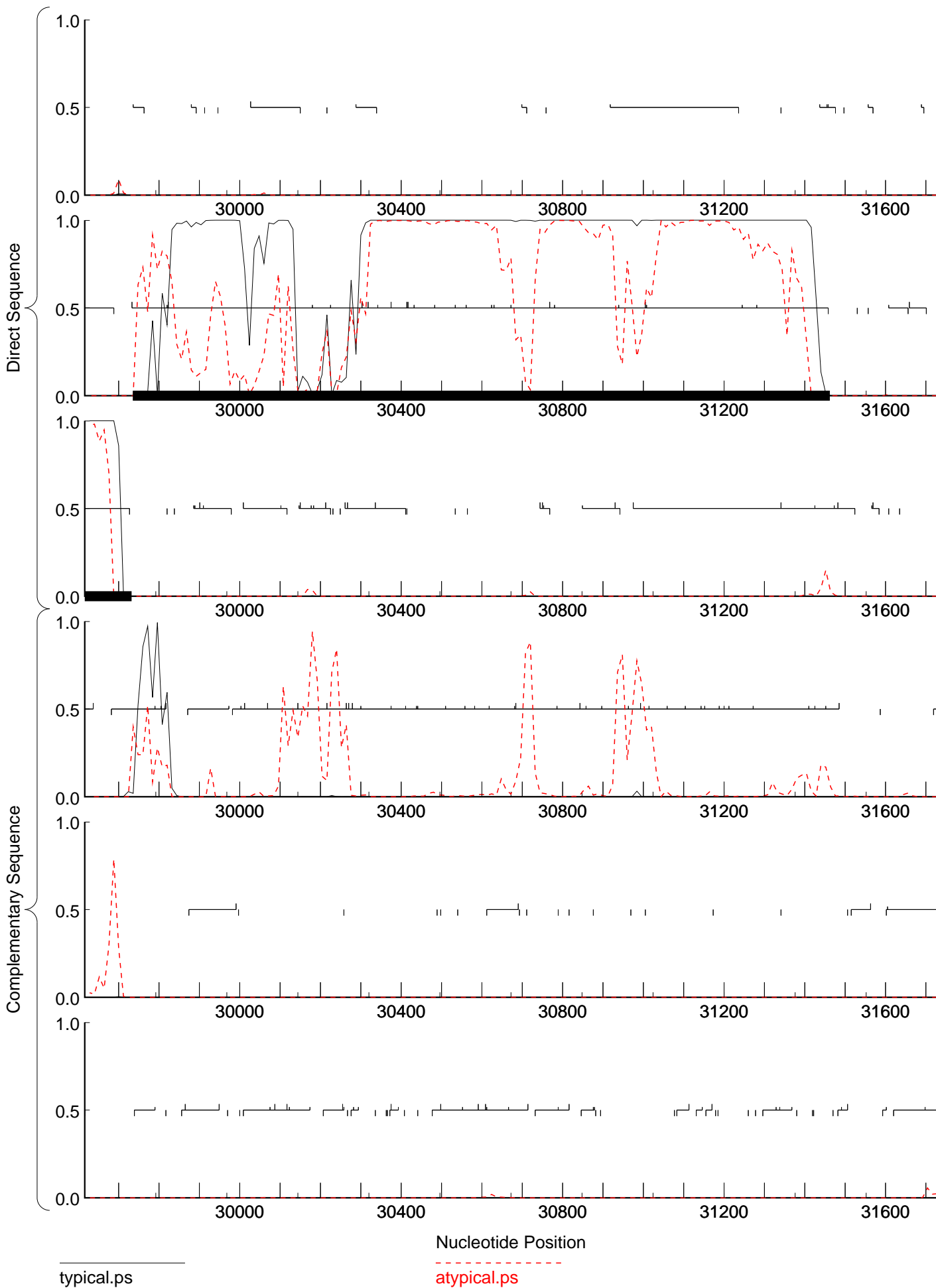


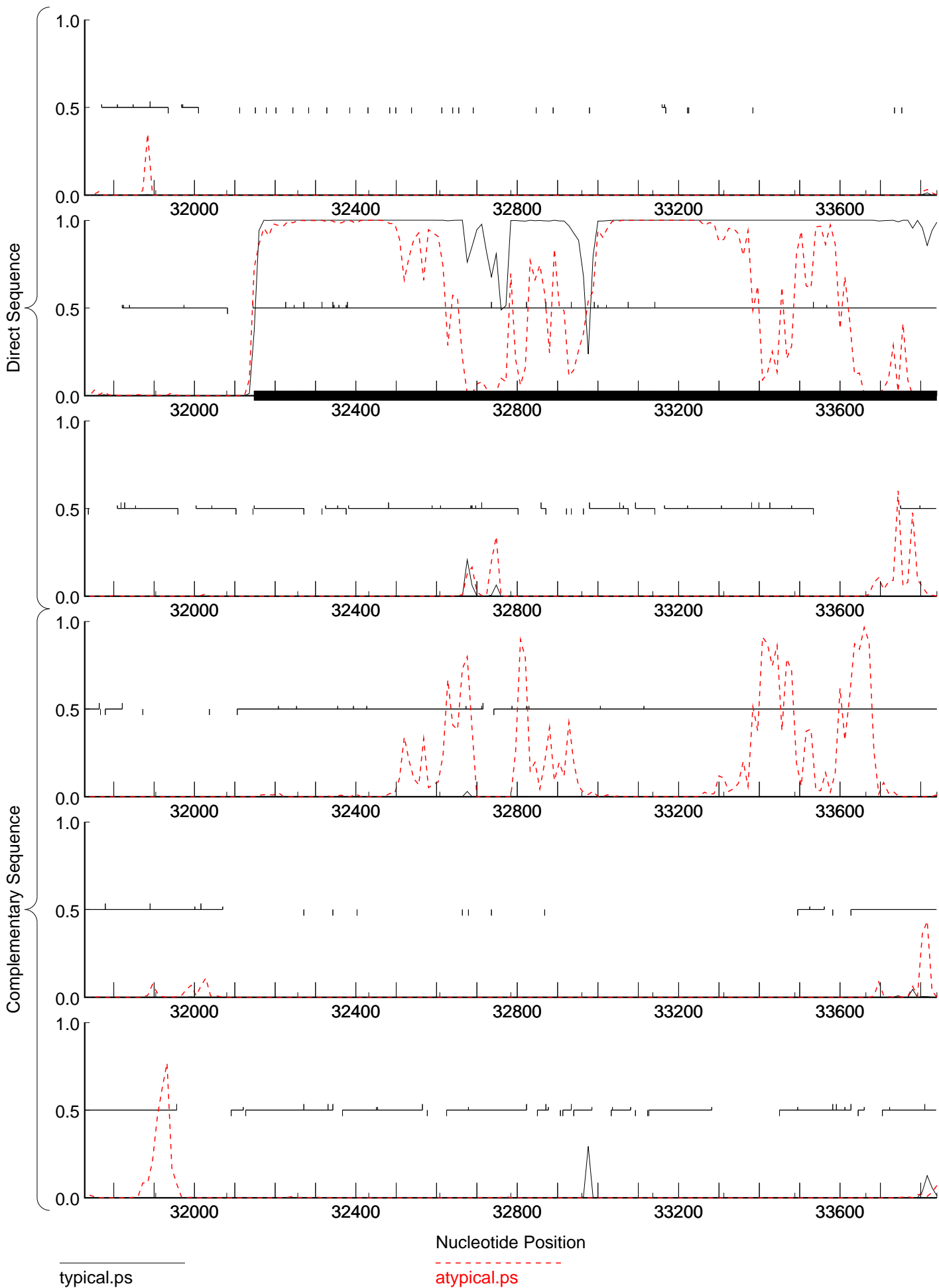
typical.ps

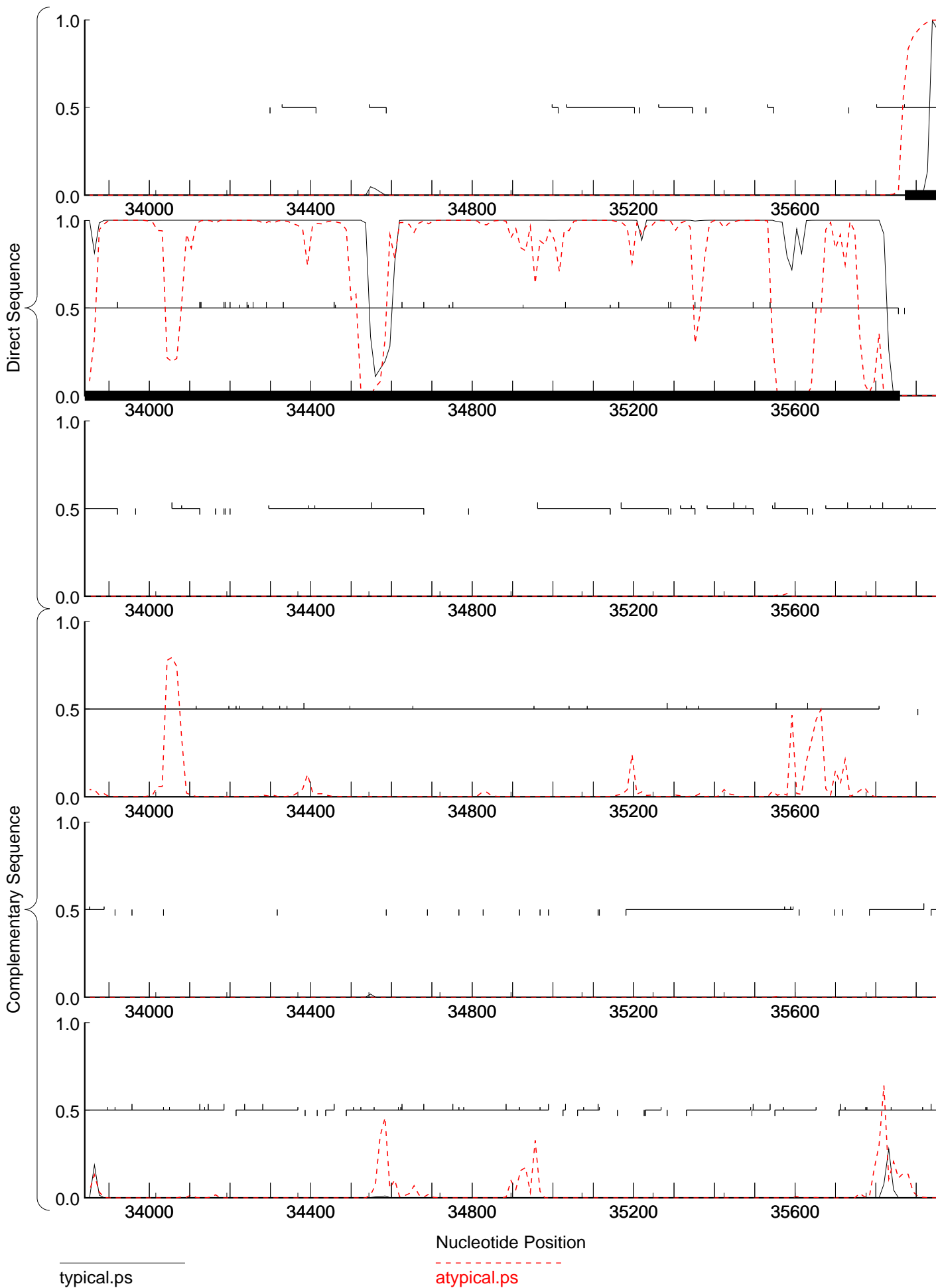
atypical.ps

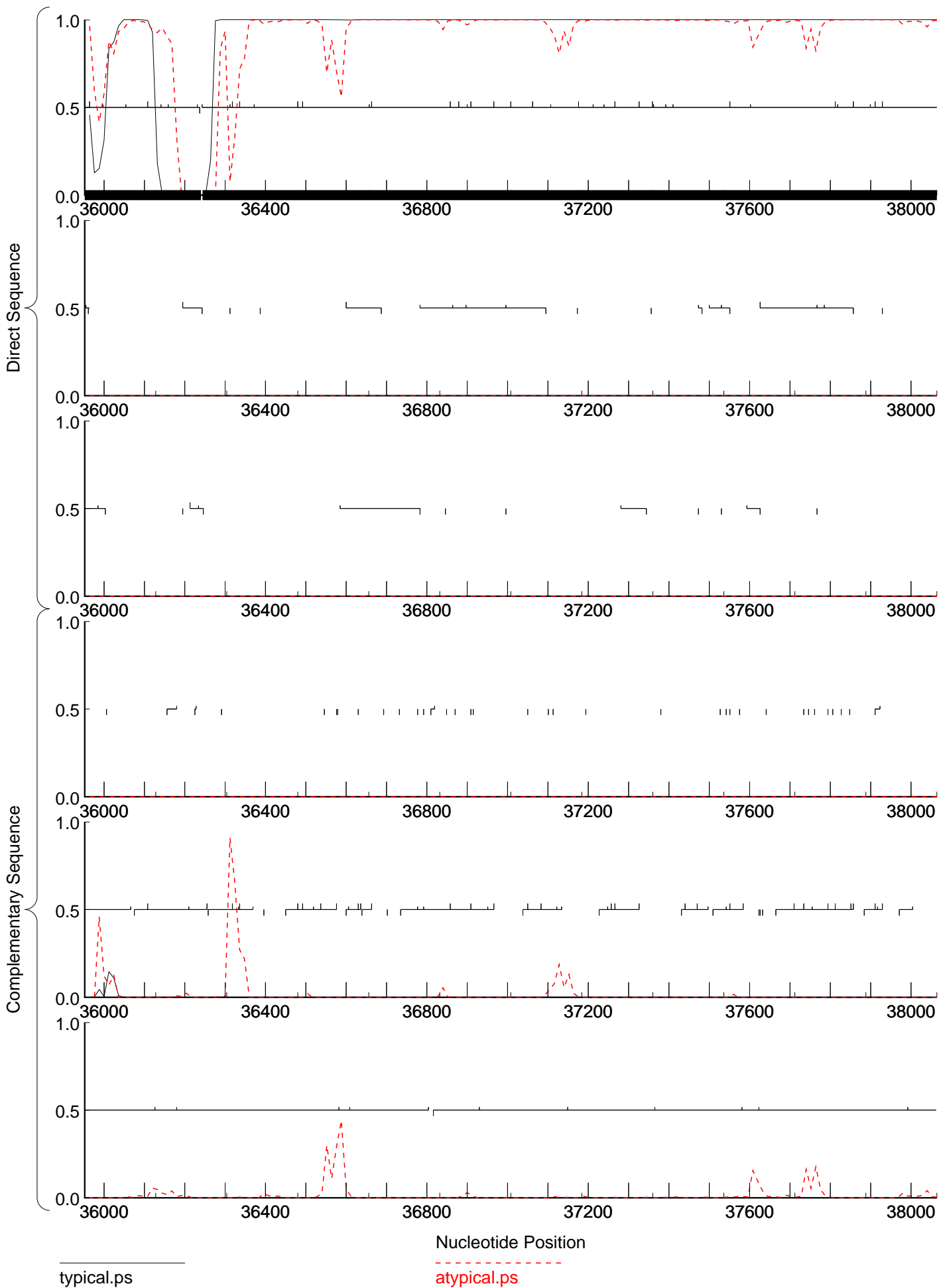


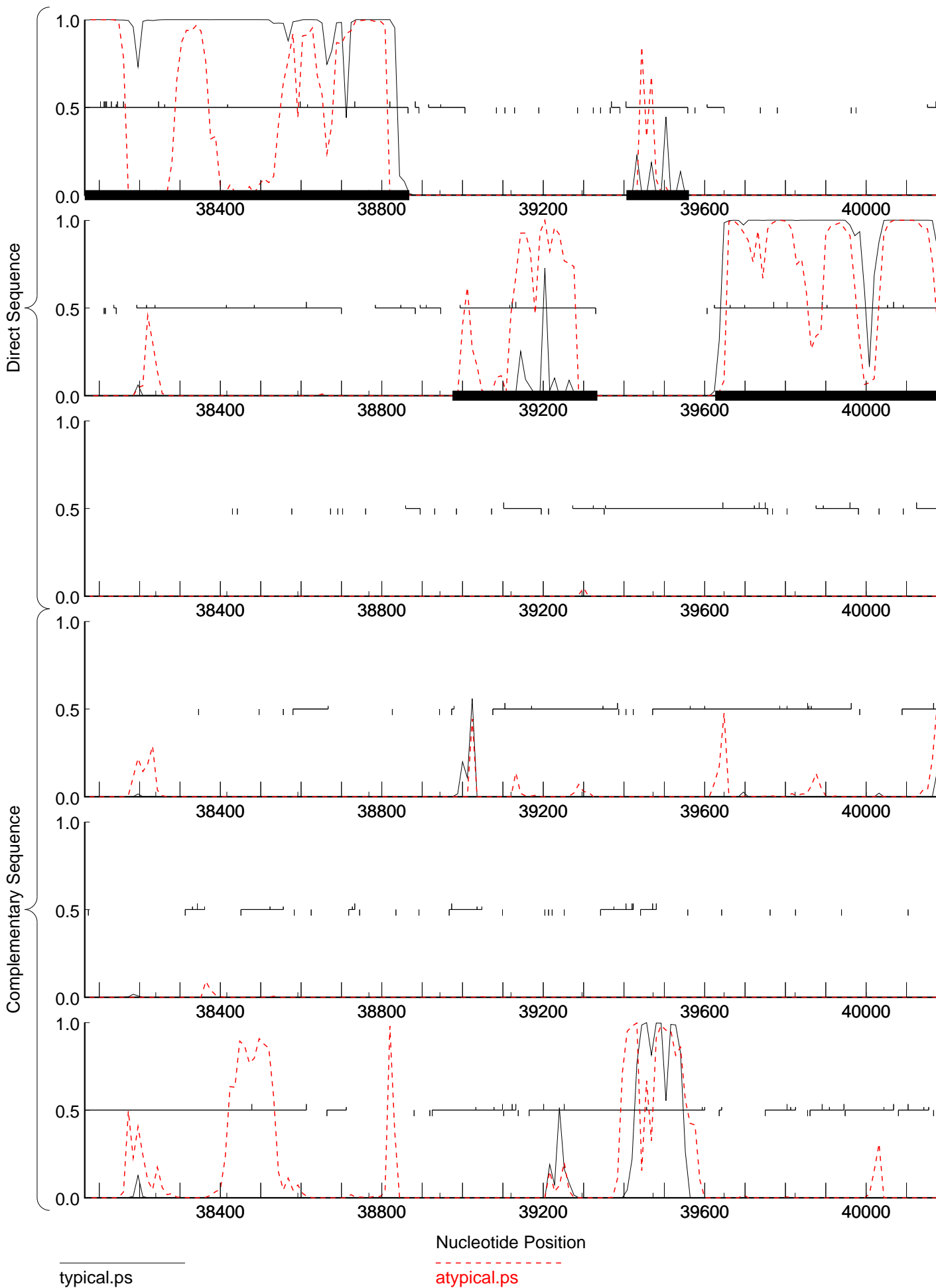


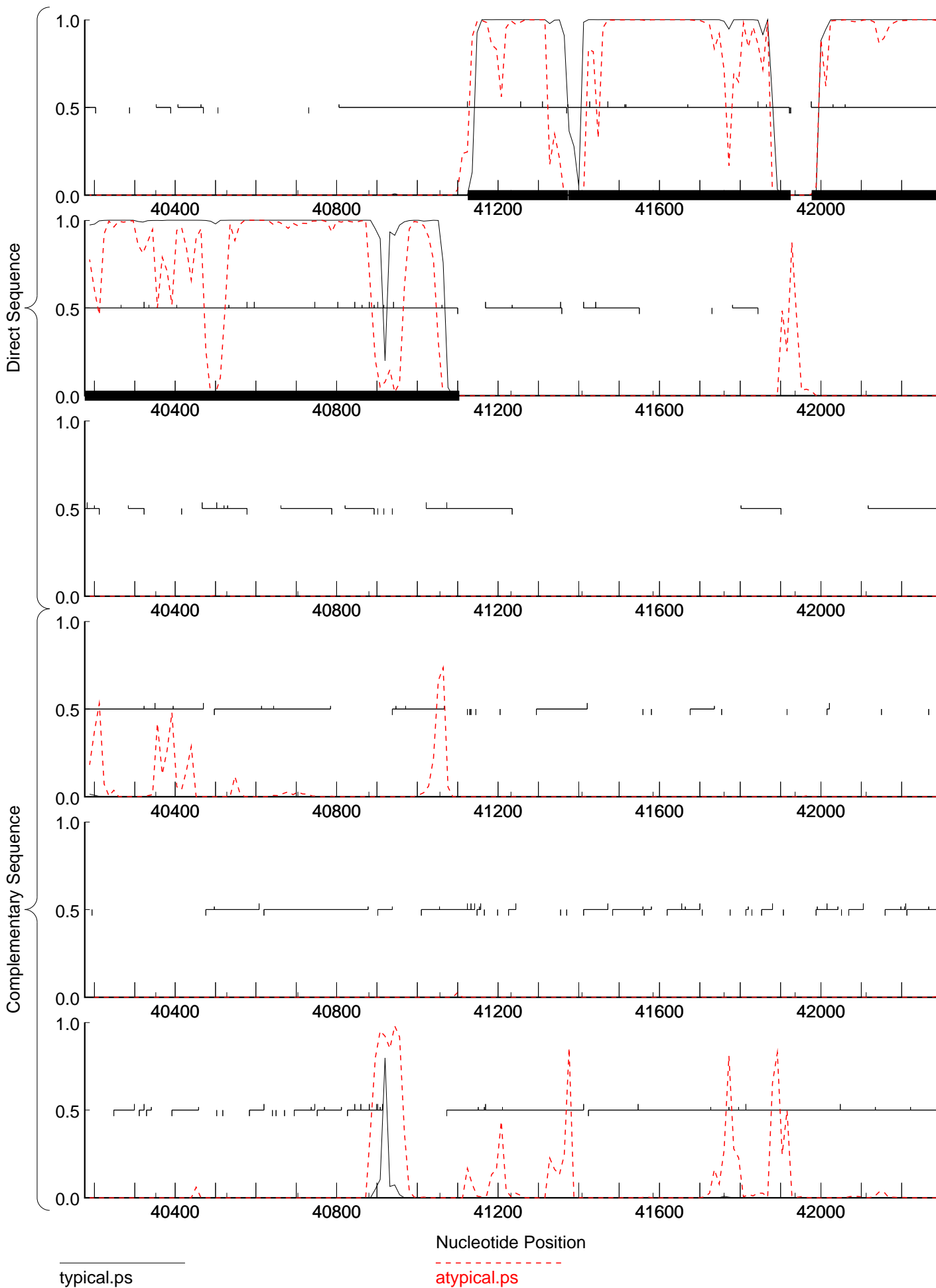


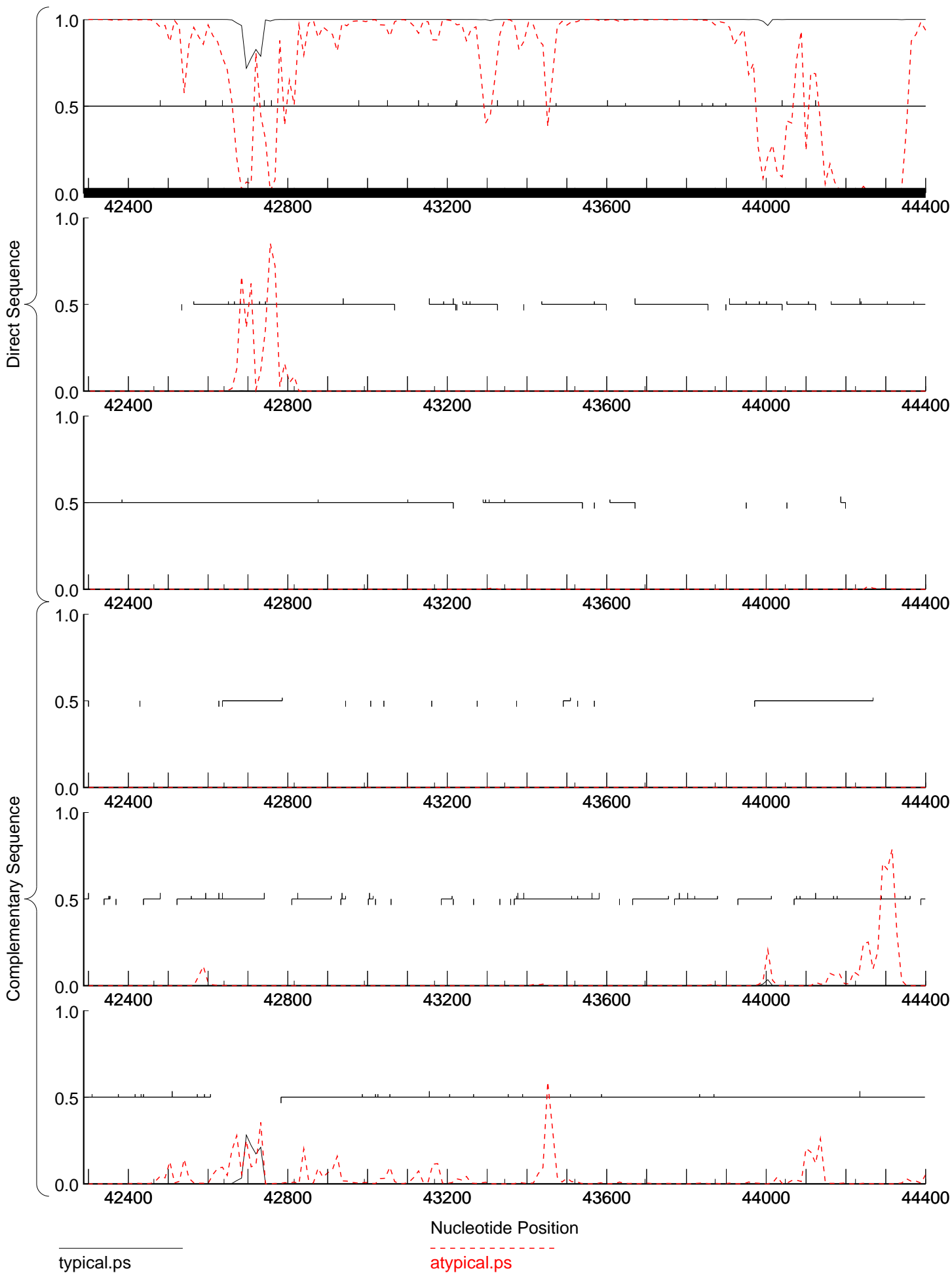






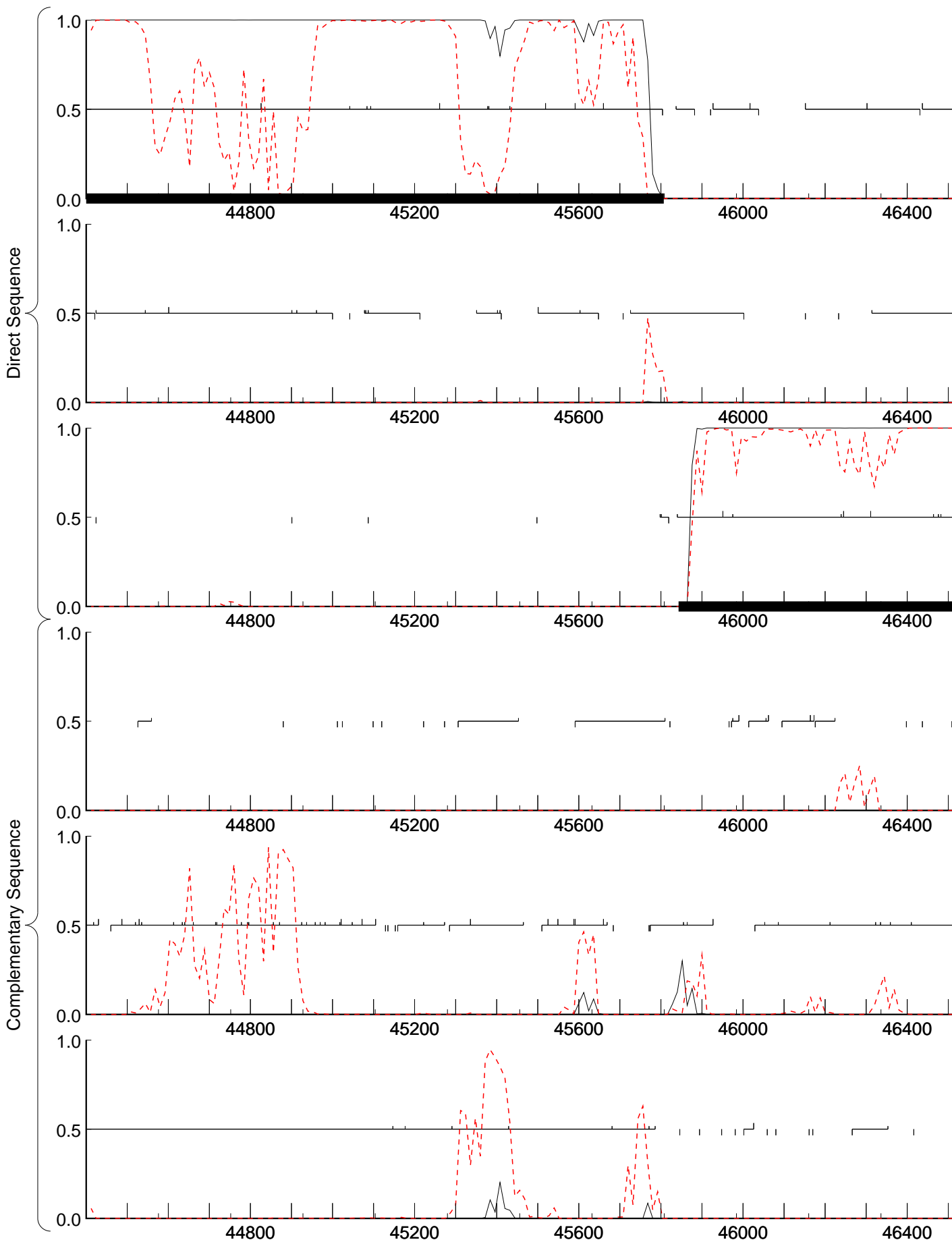






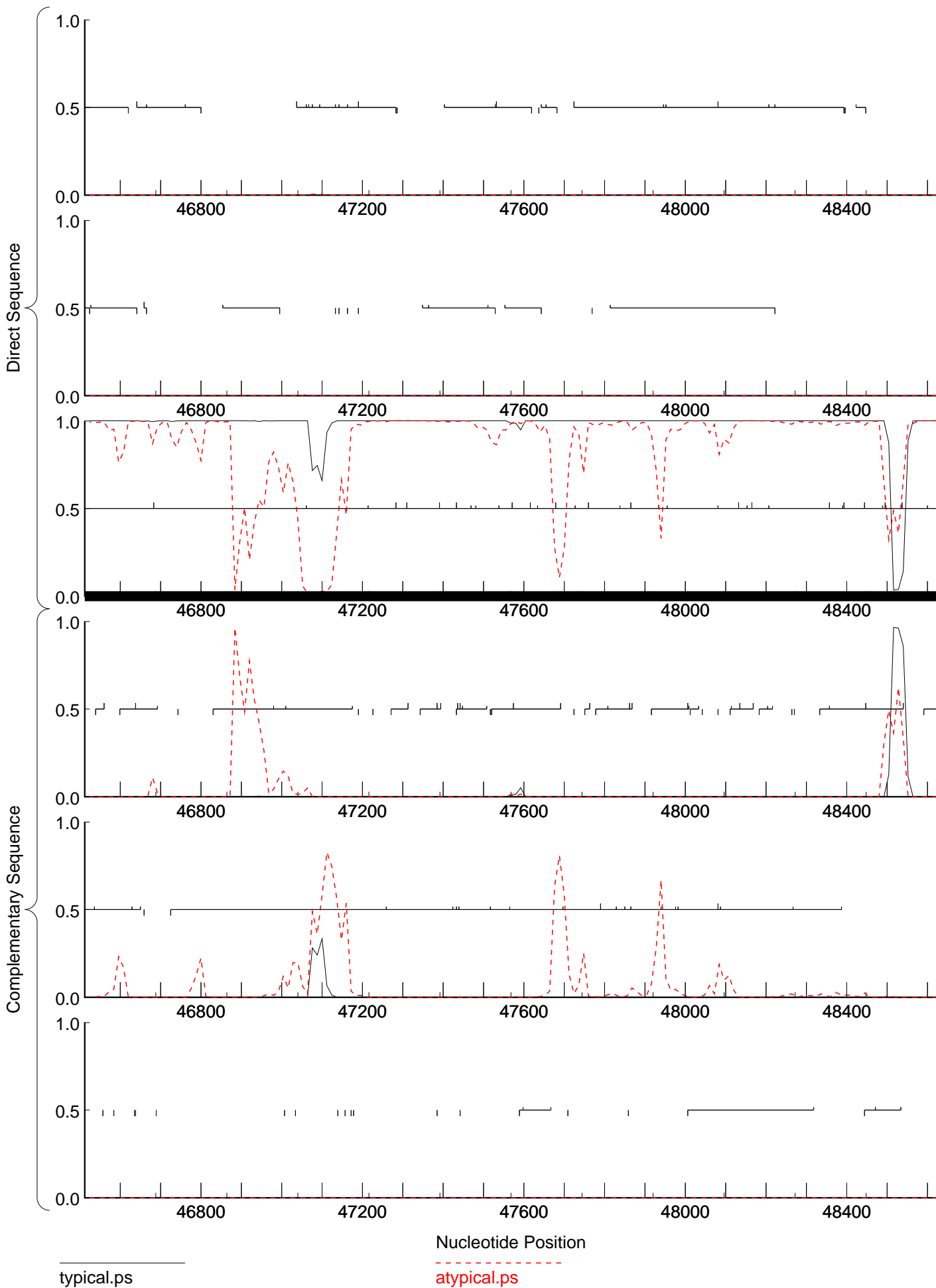
typical.ps

atypical.ps



typical.ps

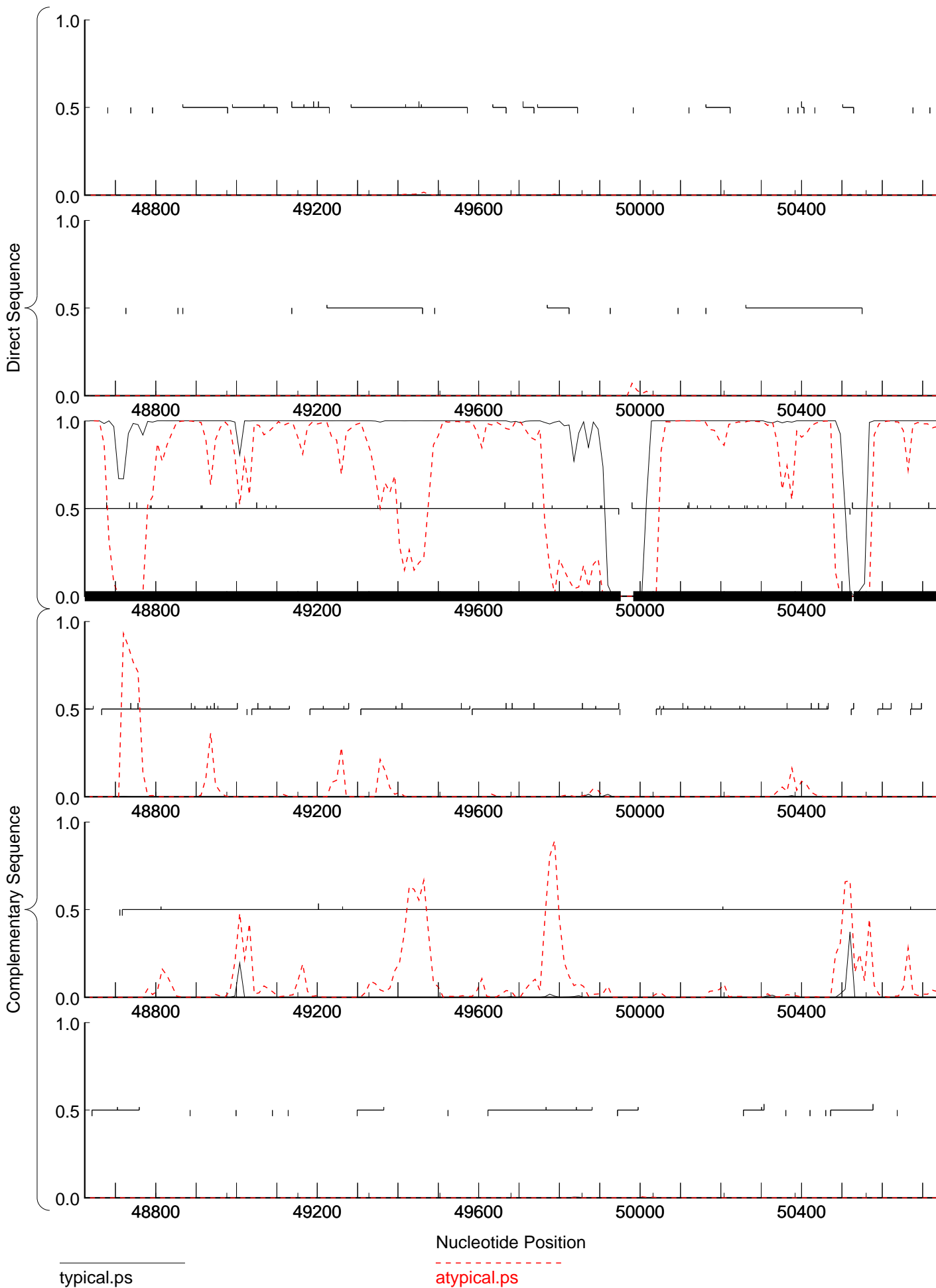
atypical.ps



typical.ps

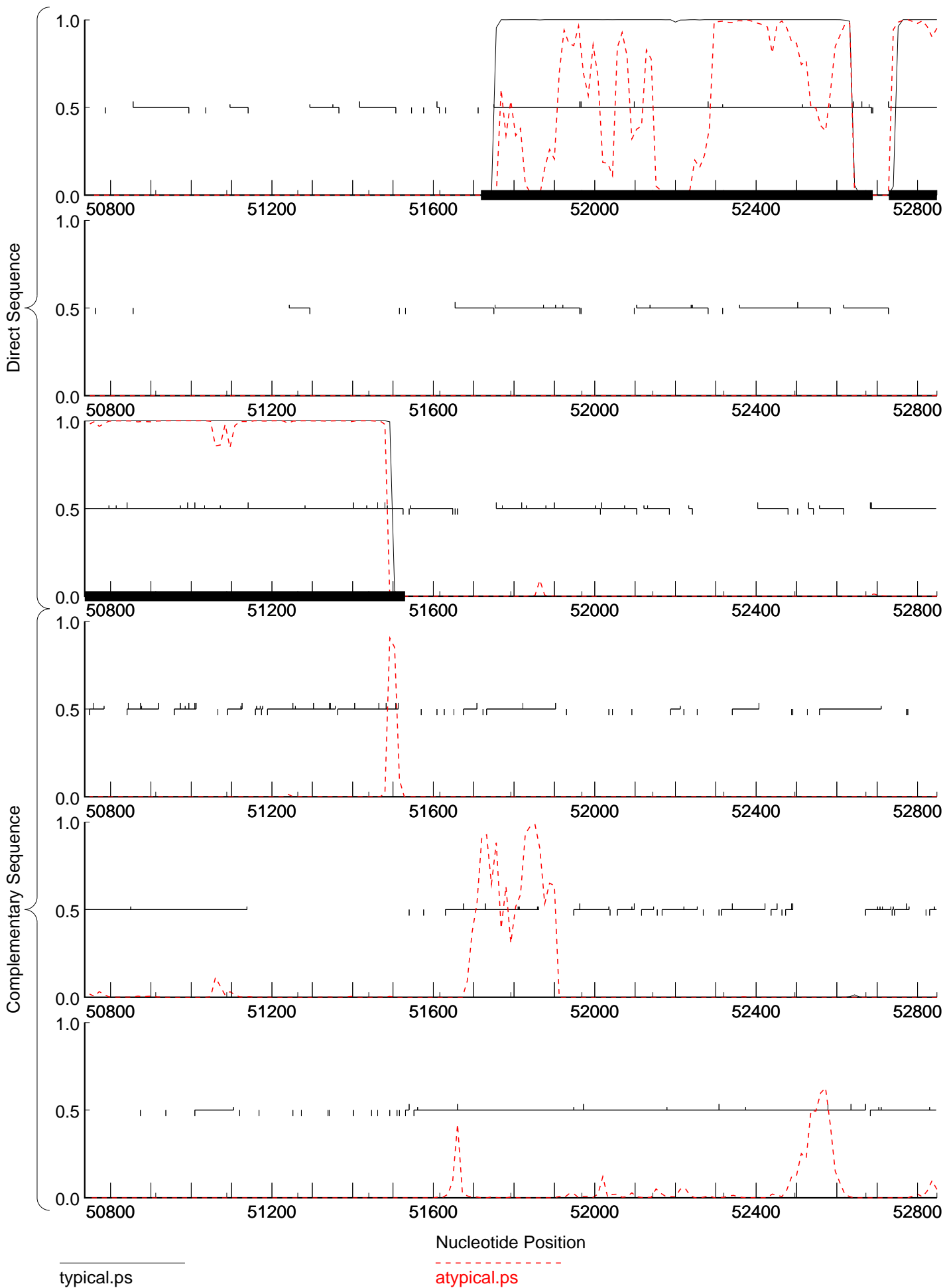
Nucleotide Position

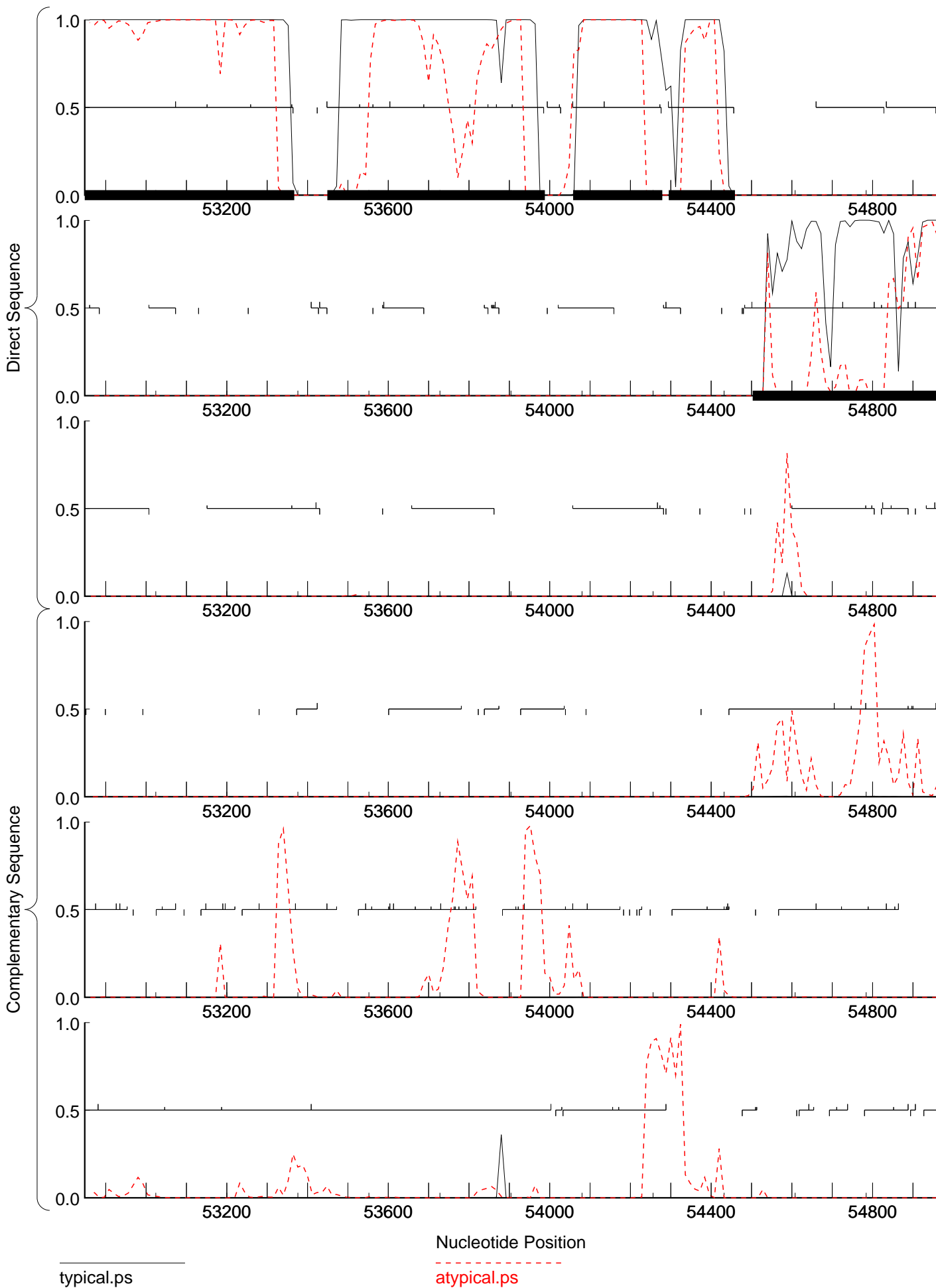
atypical.ps

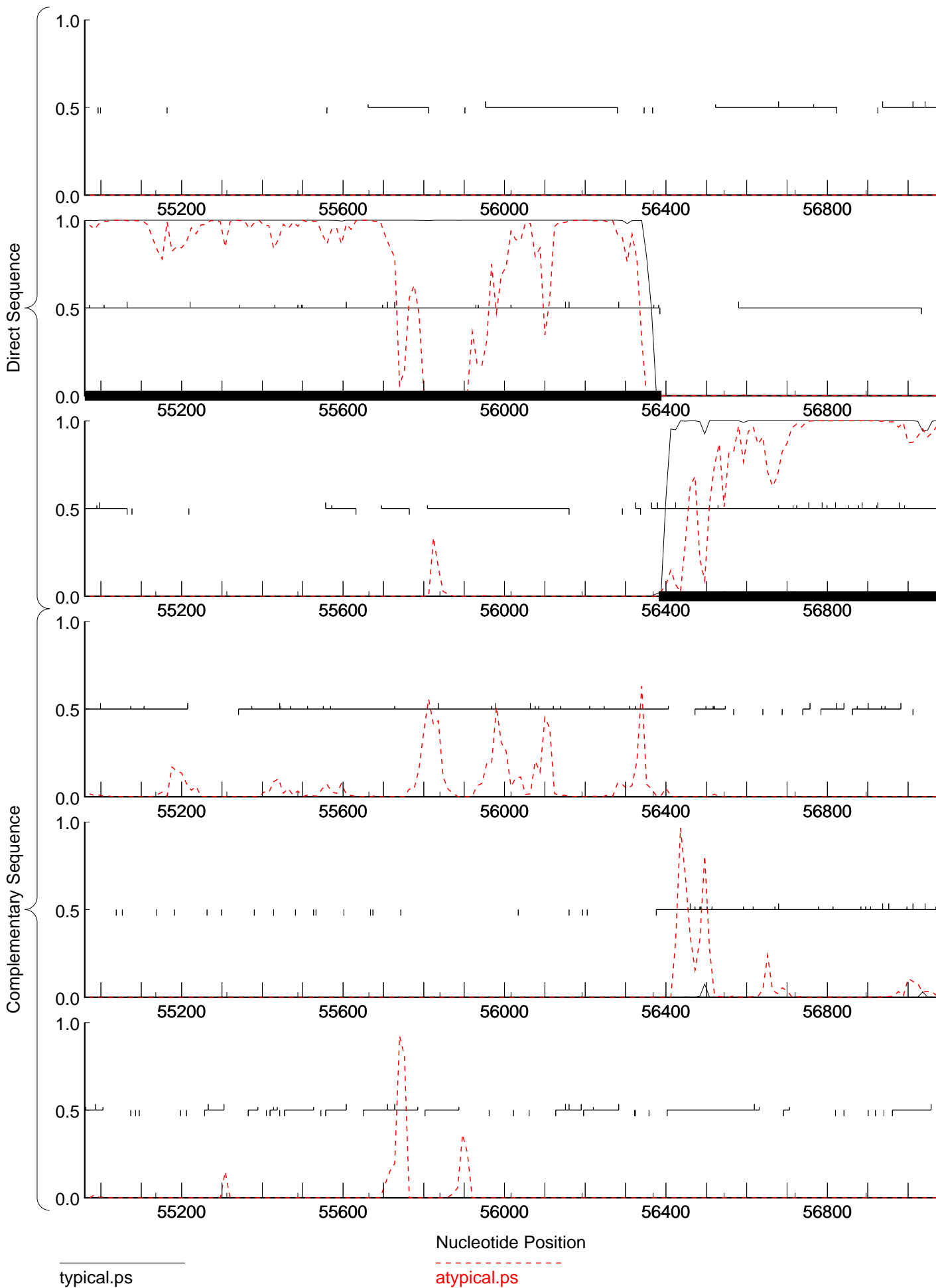


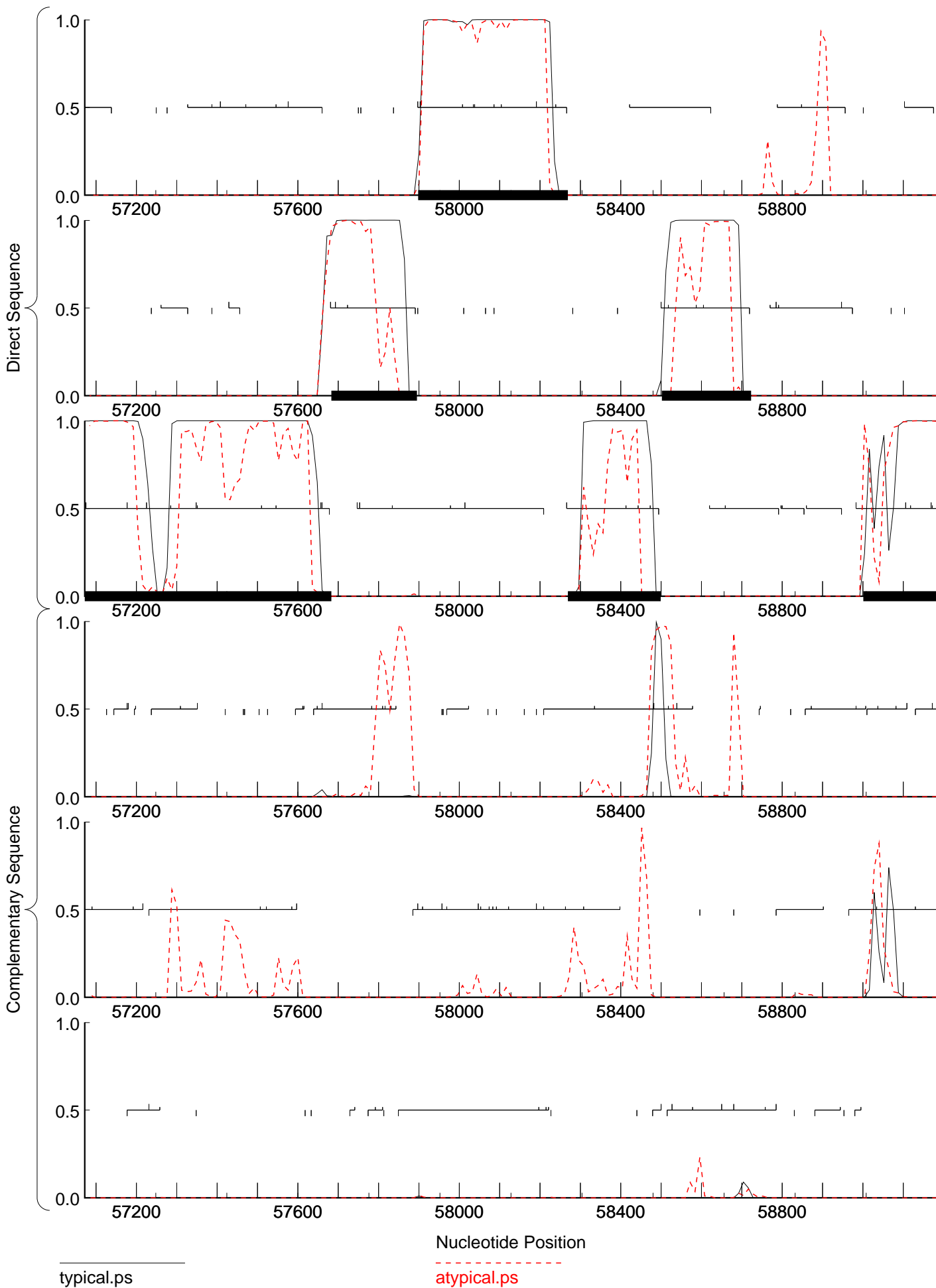
typical.ps

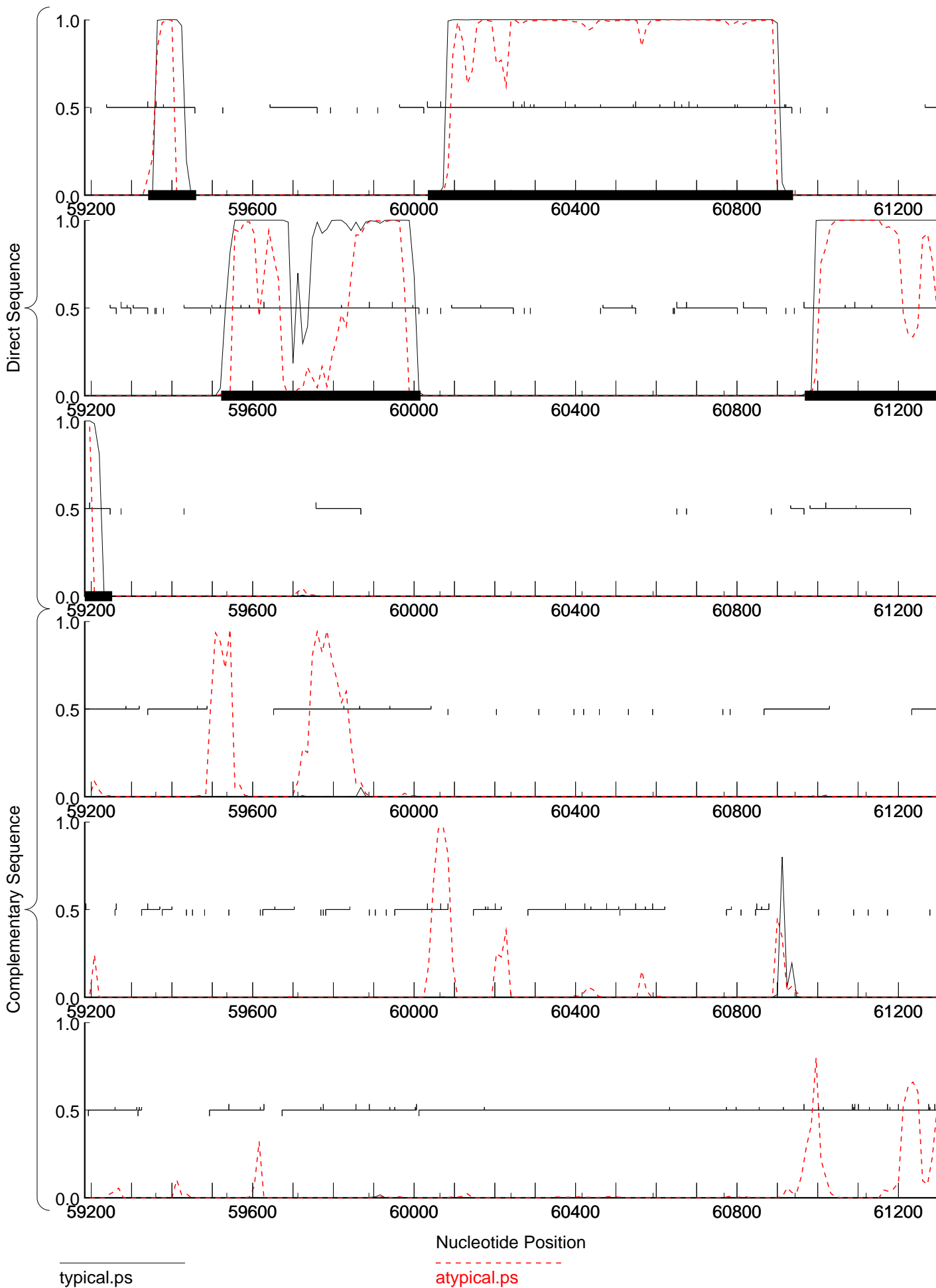
atypical.ps

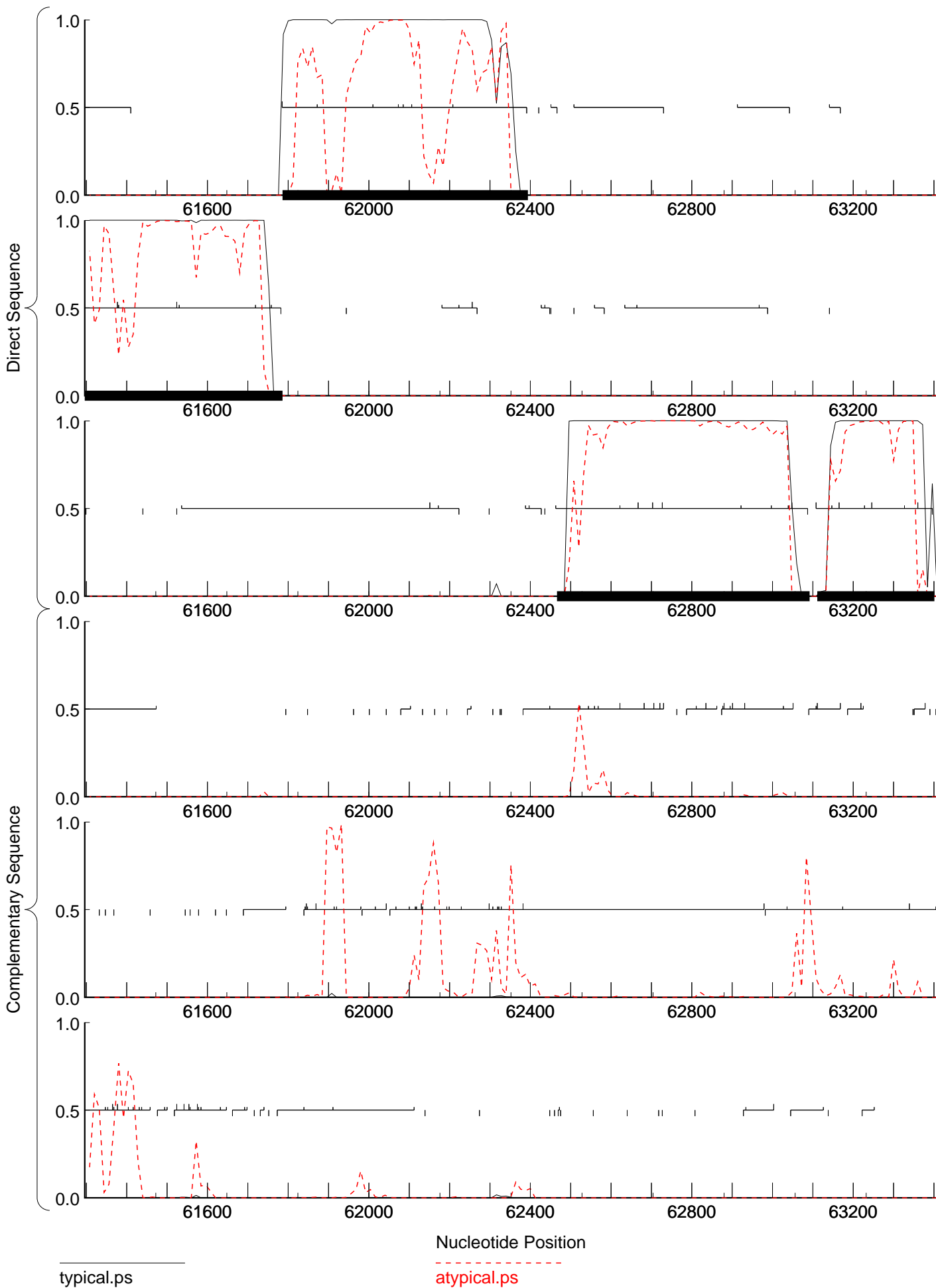


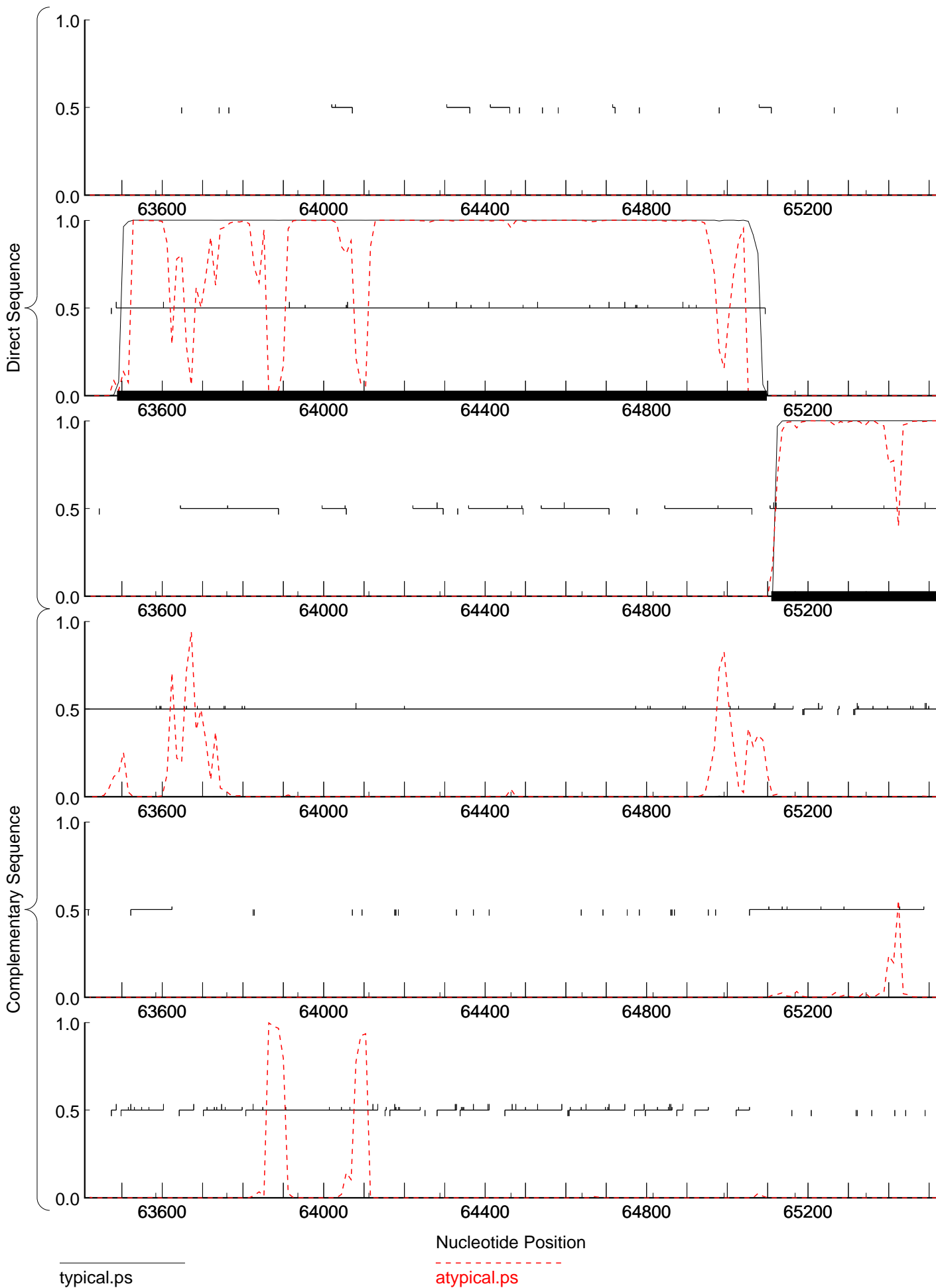


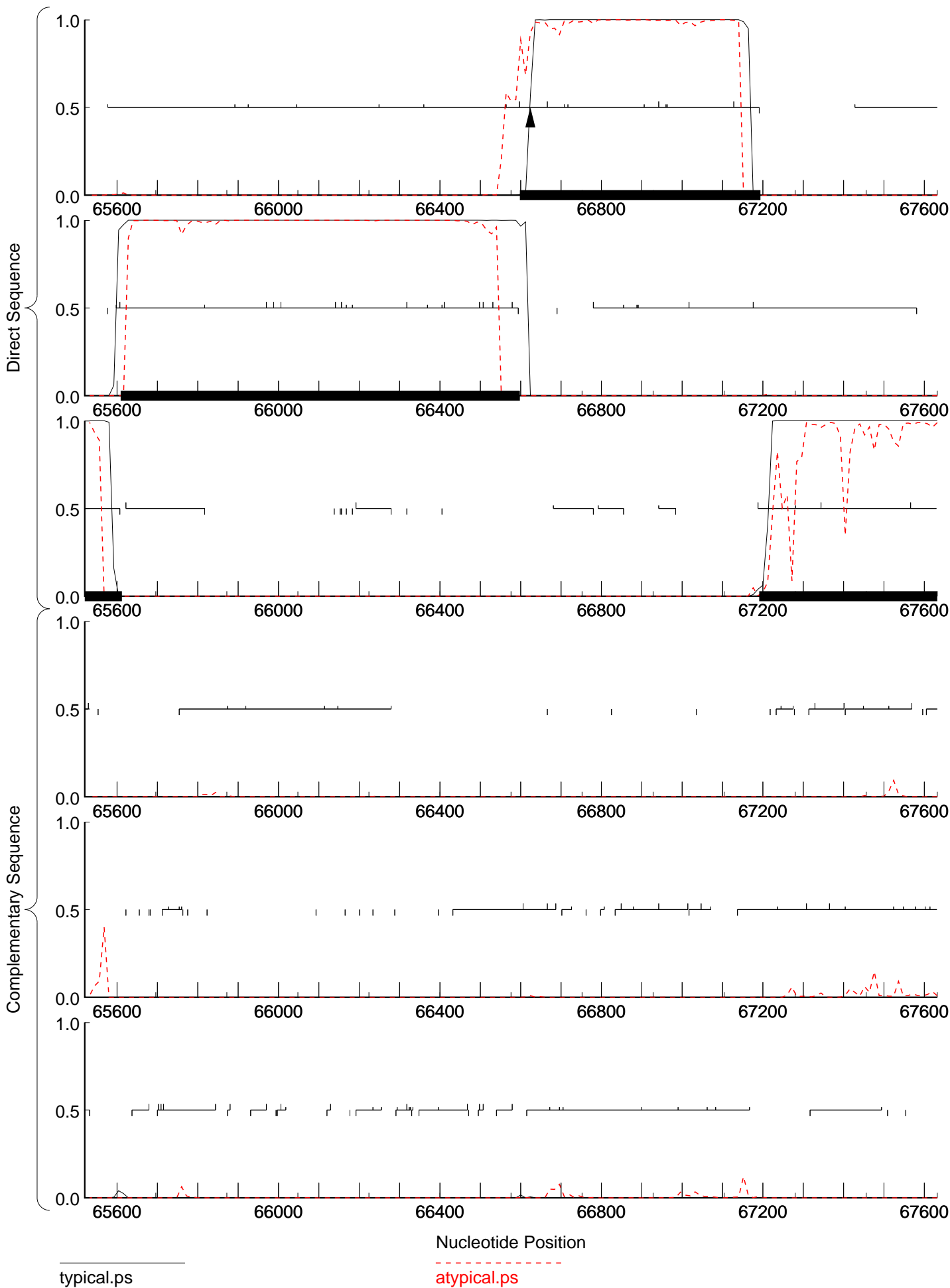


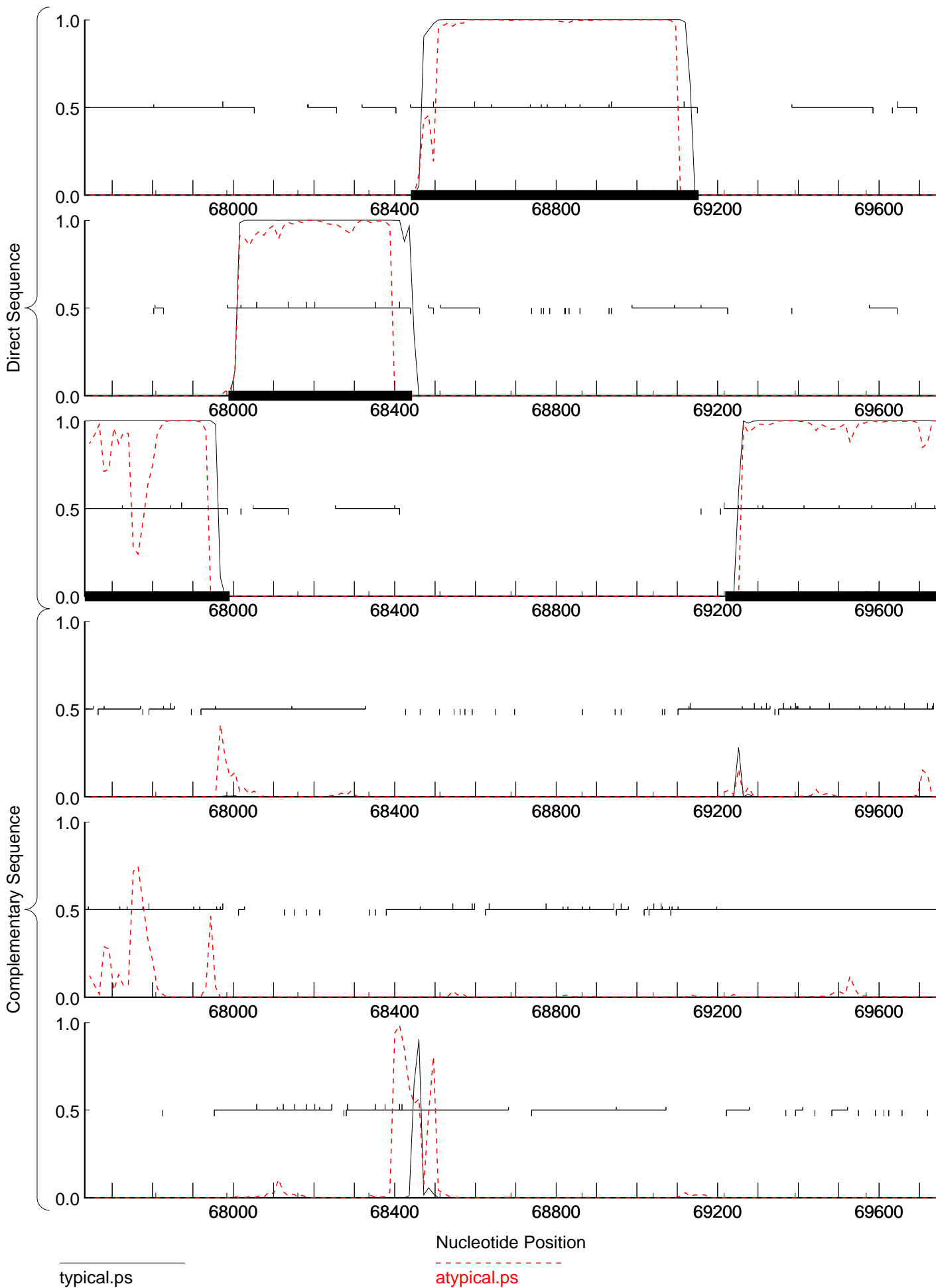


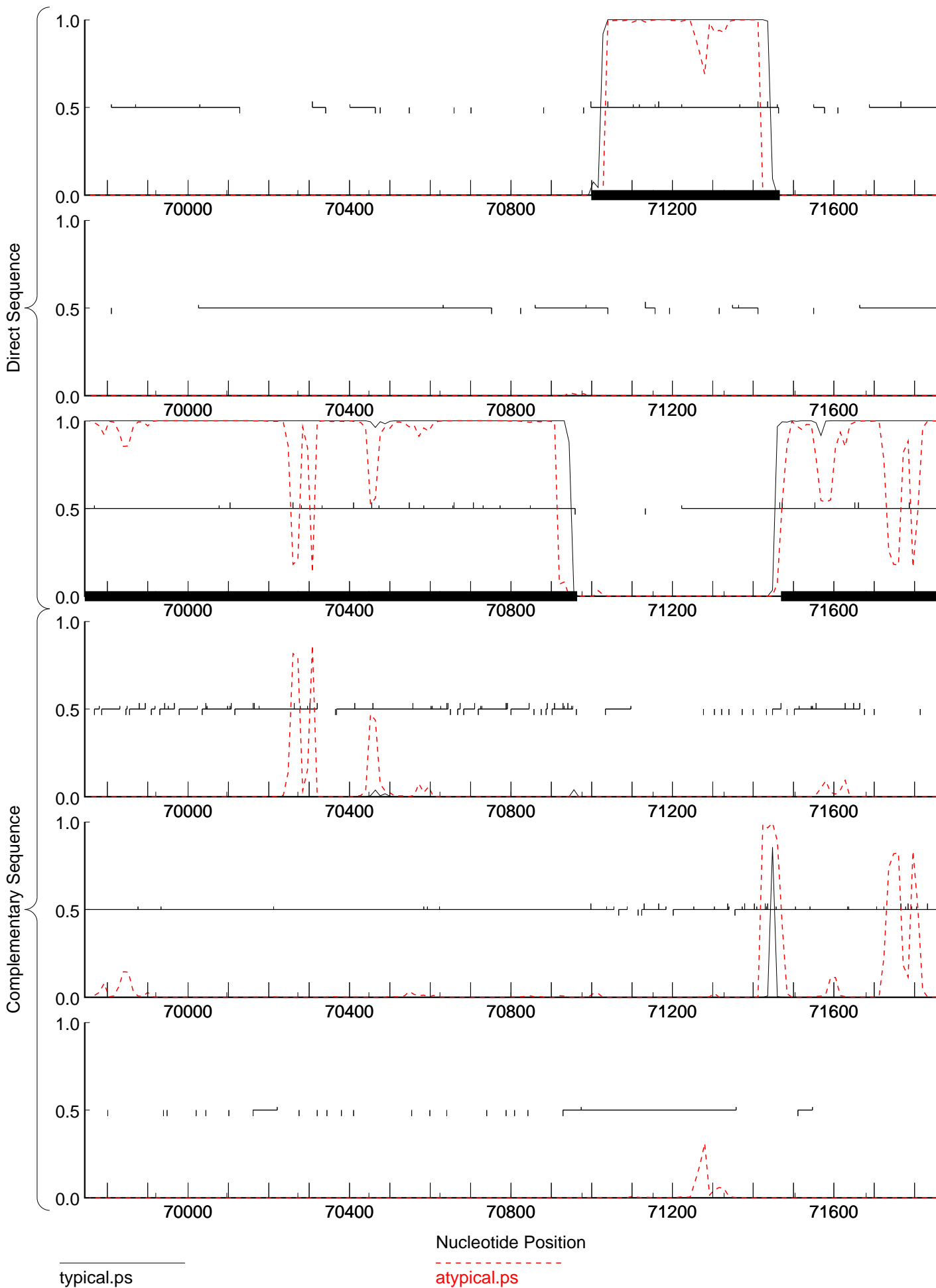


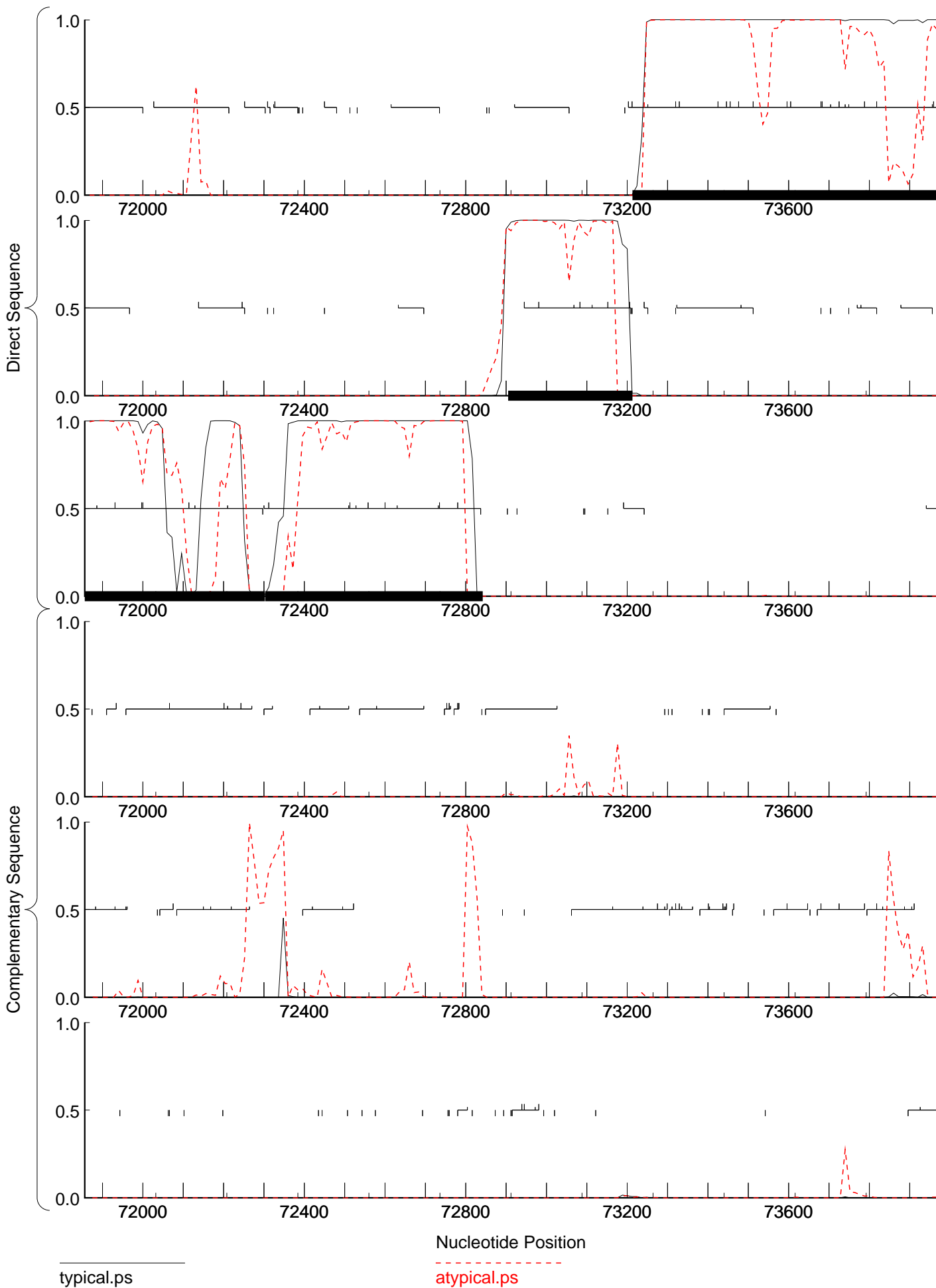


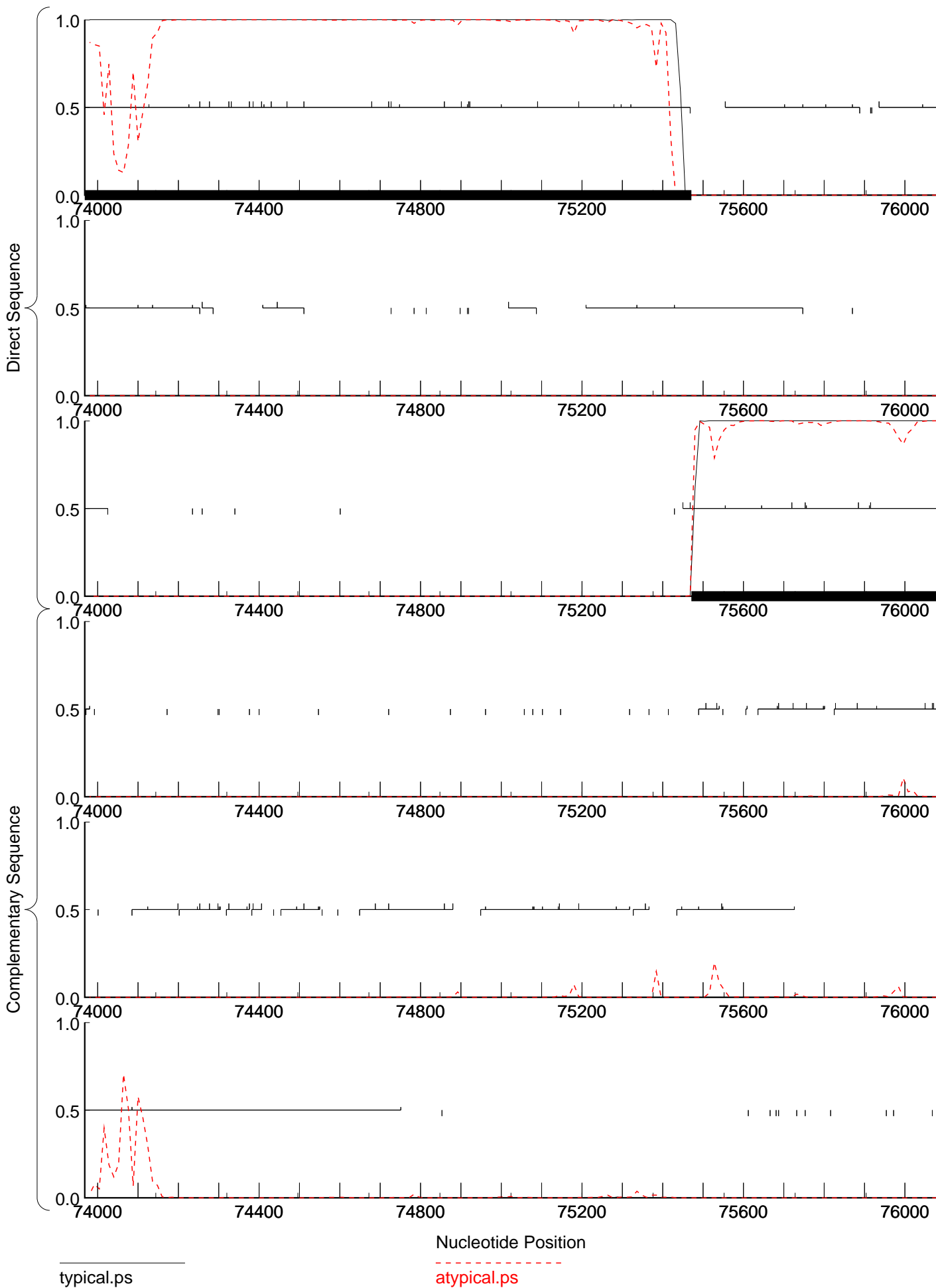






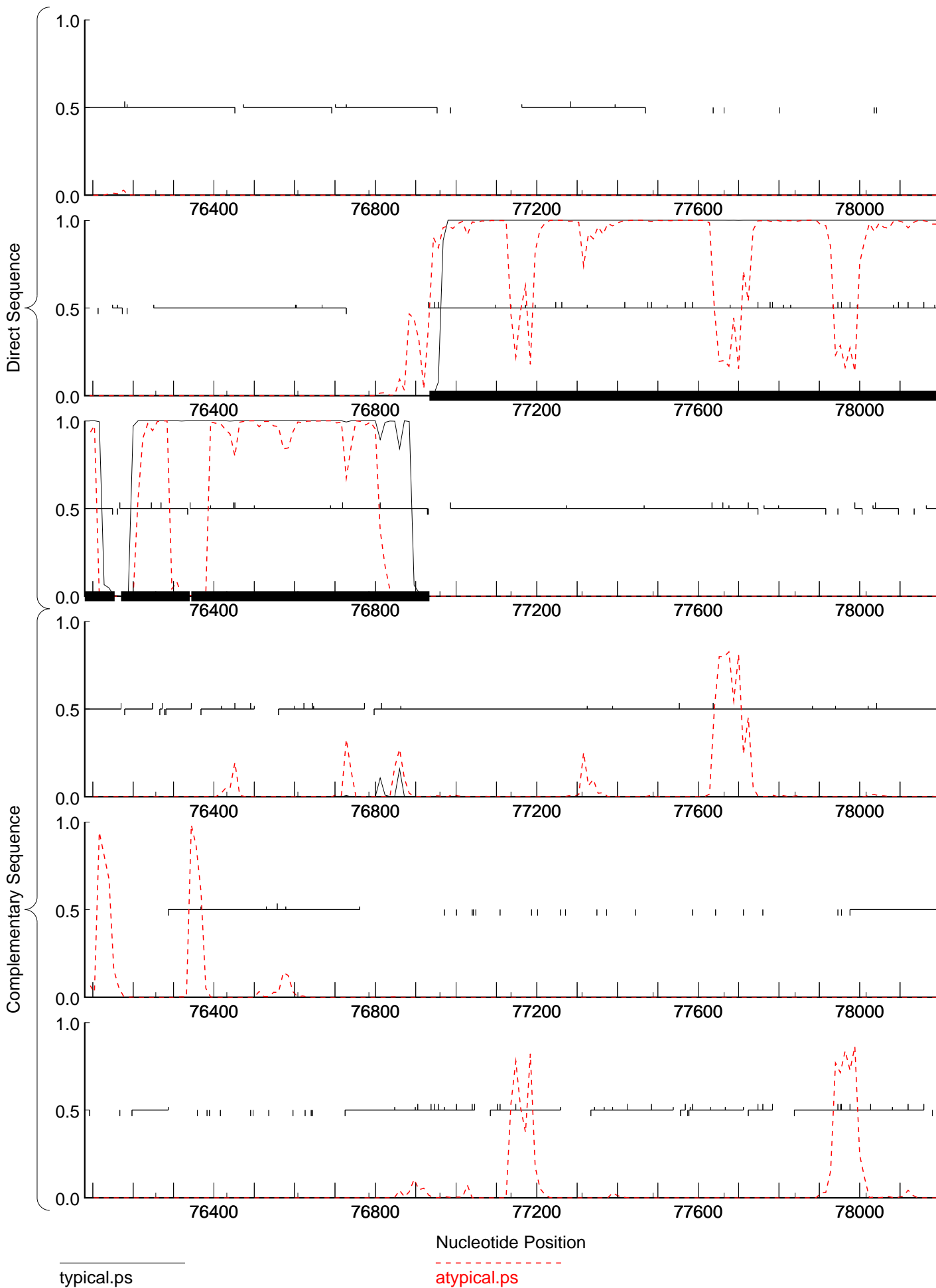


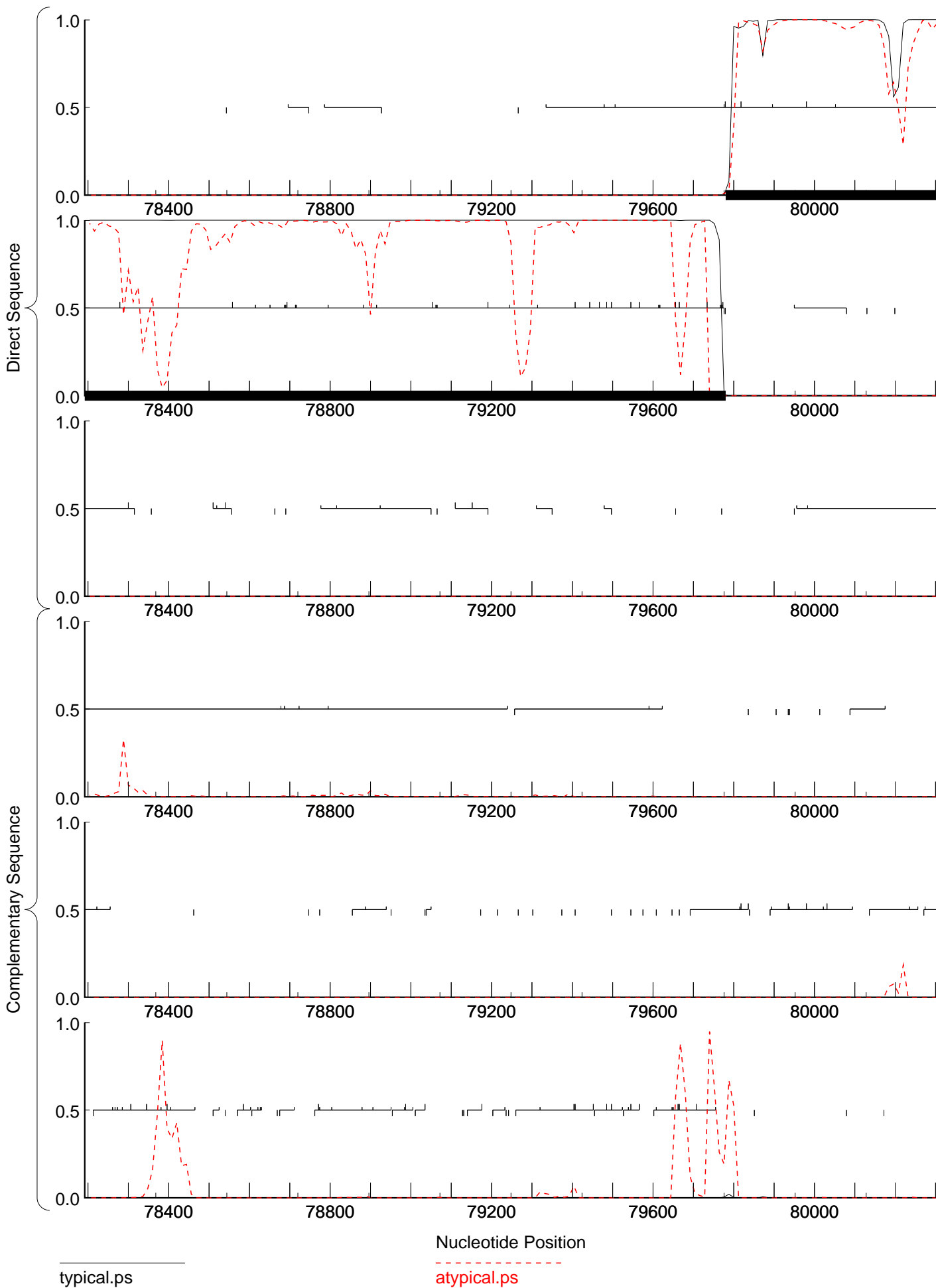


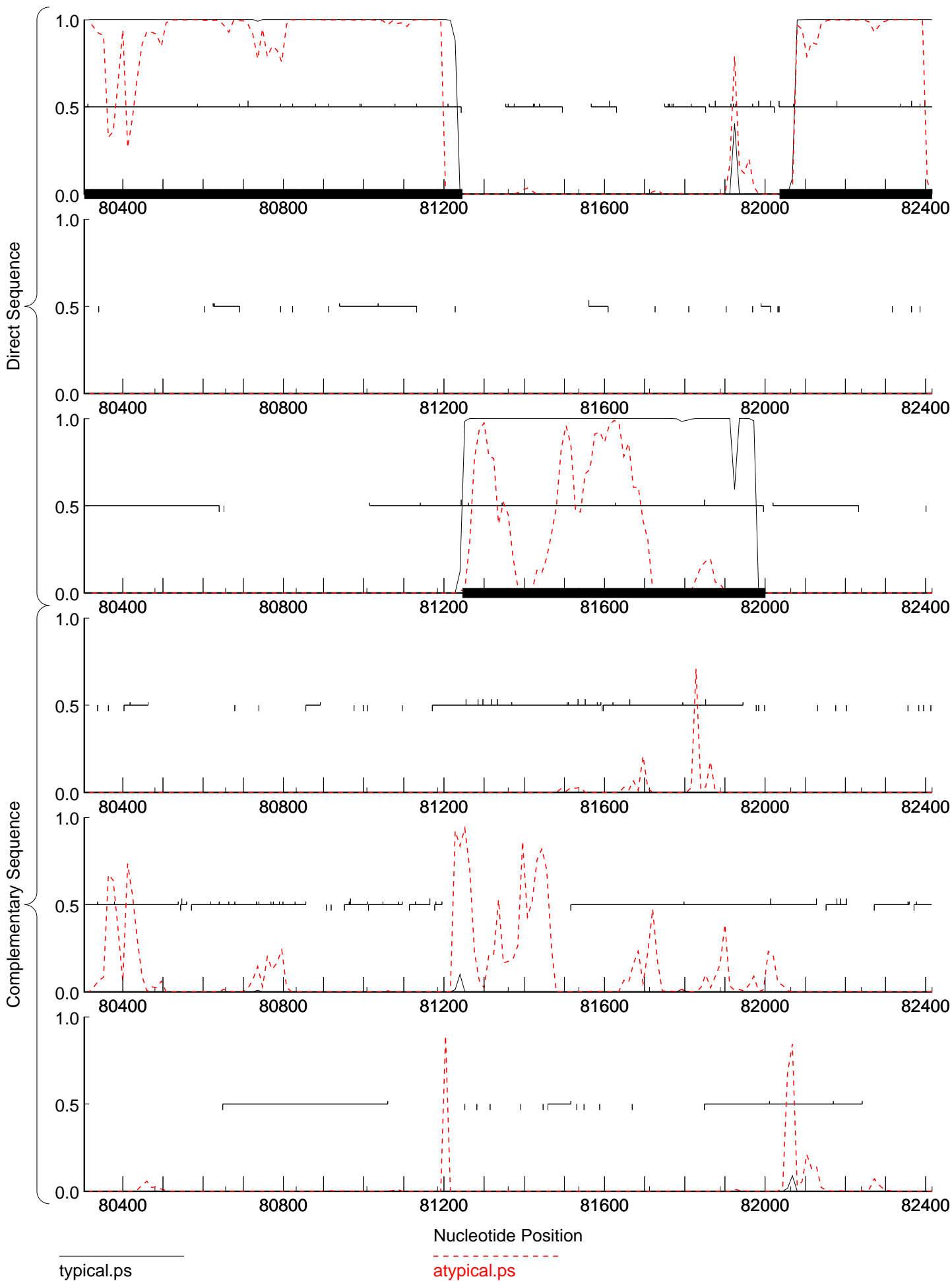


typical.ps

atypical.ps

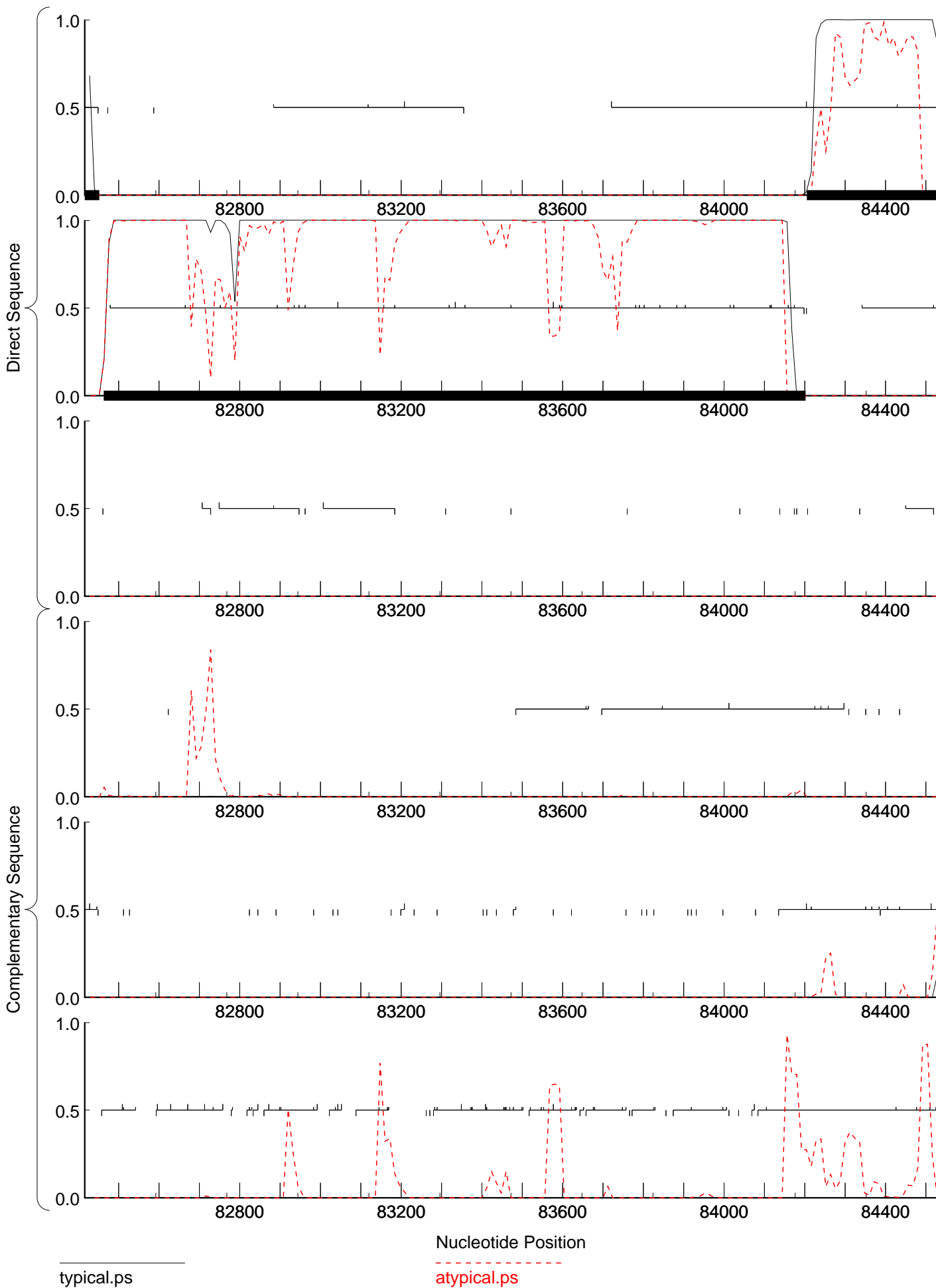


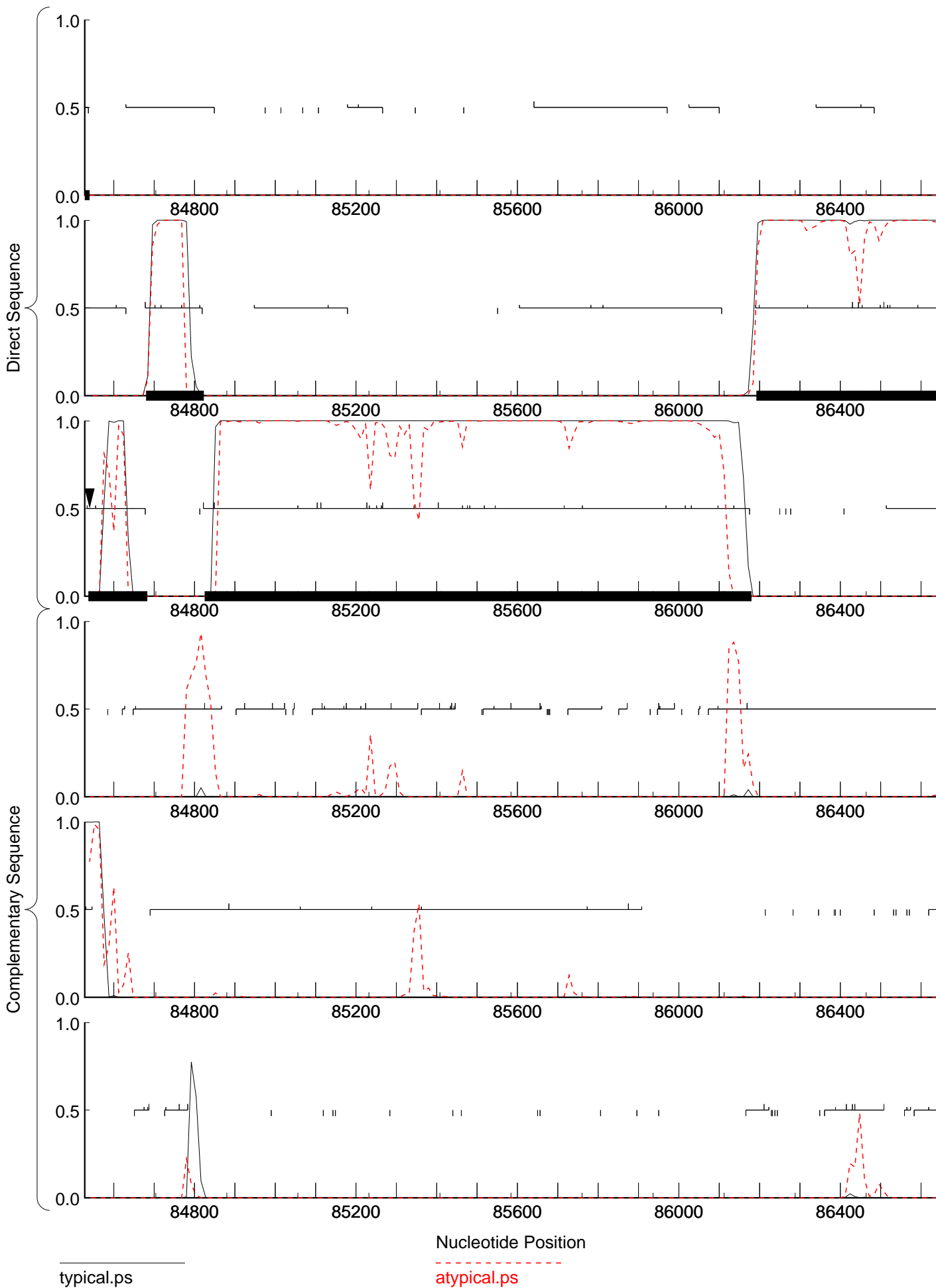


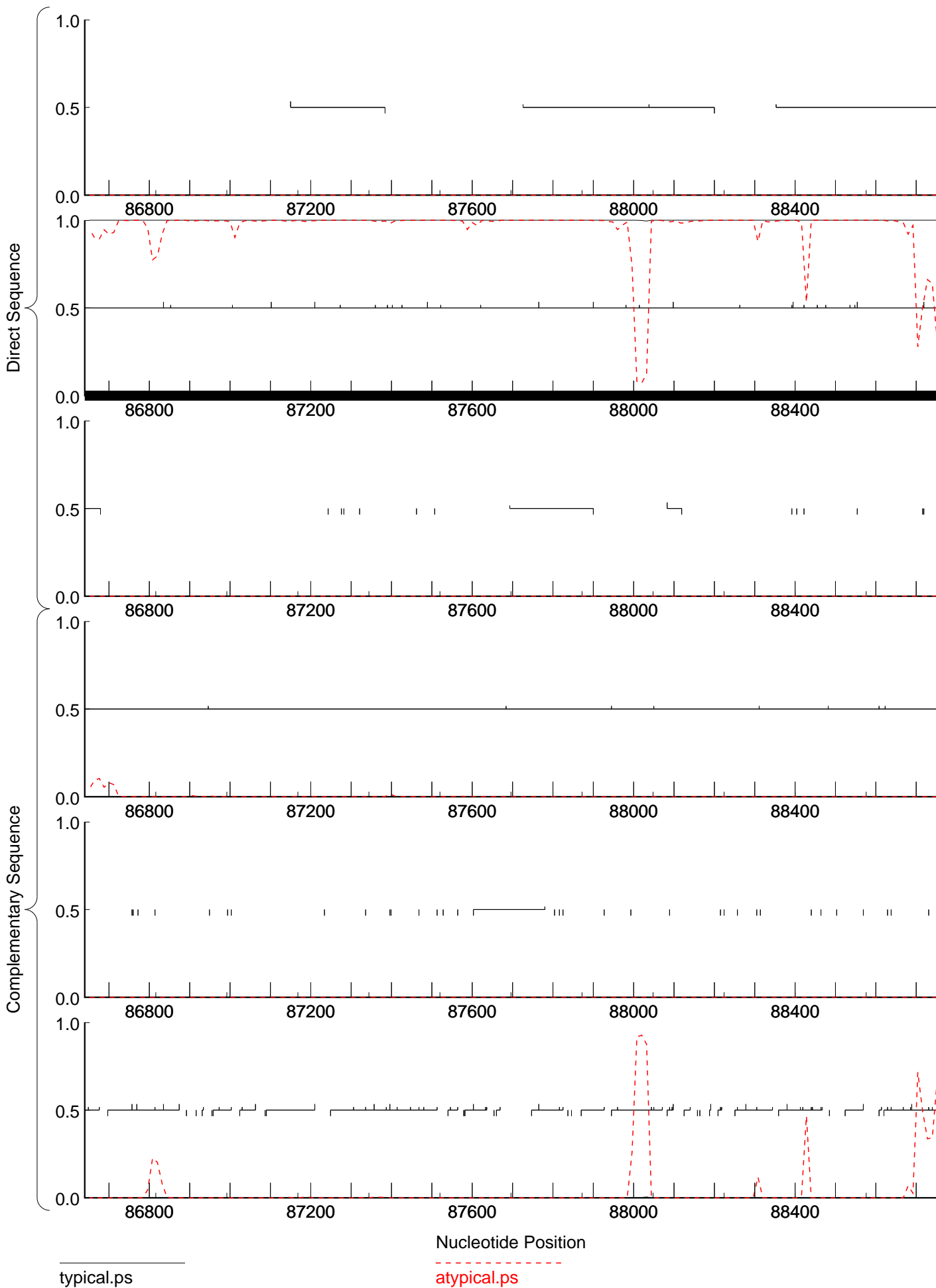


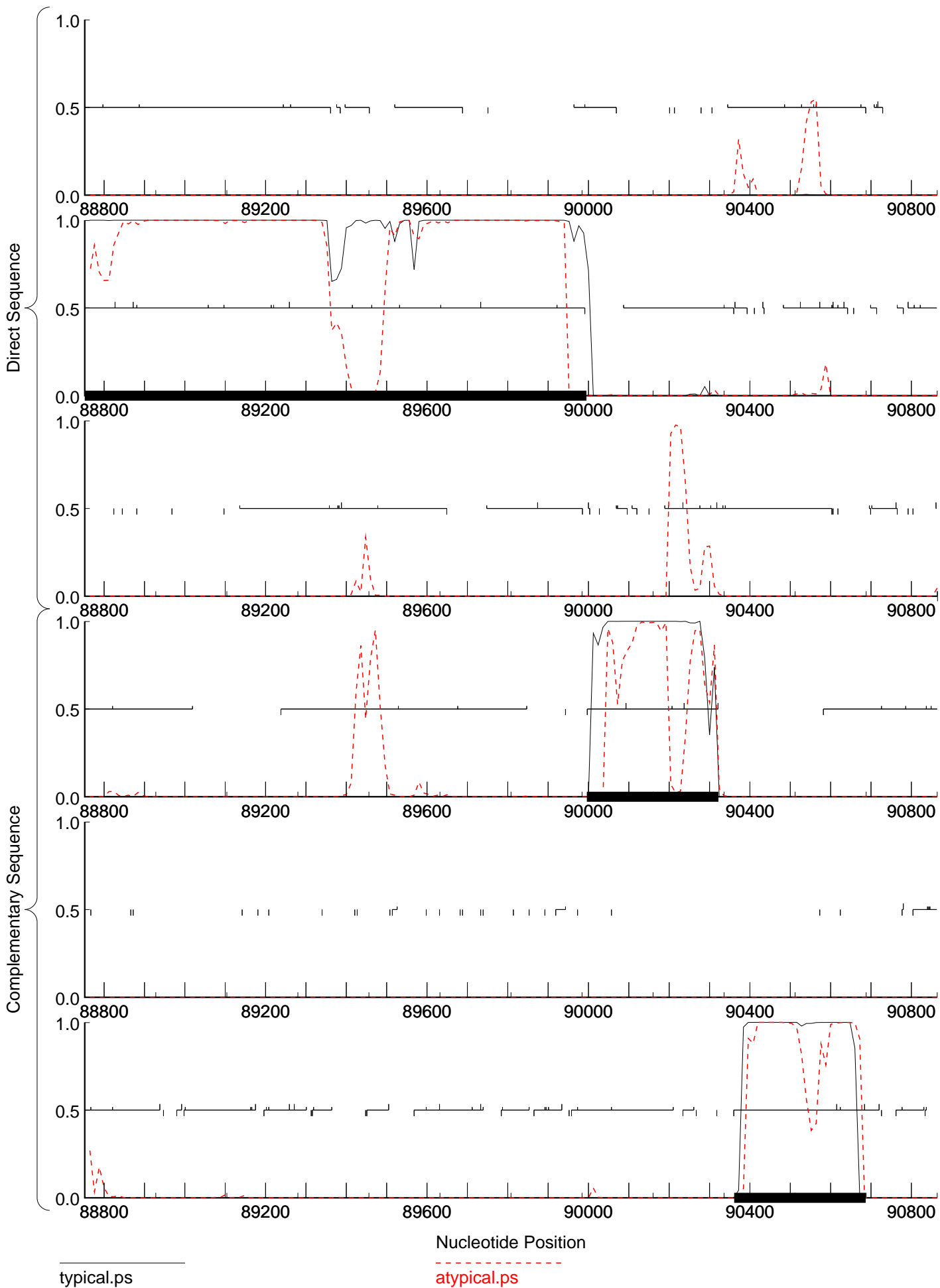
typical.ps

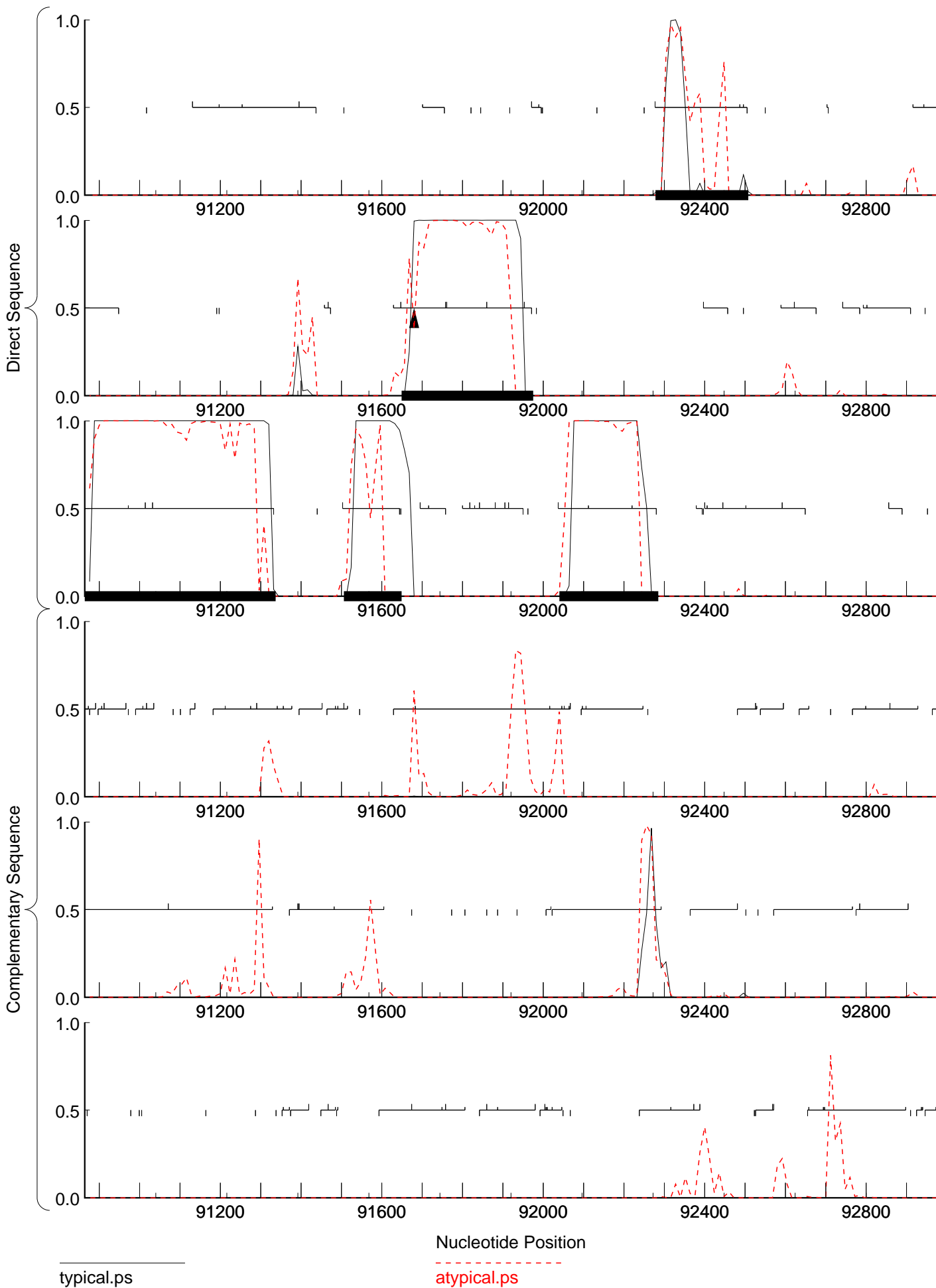
atypical.ps





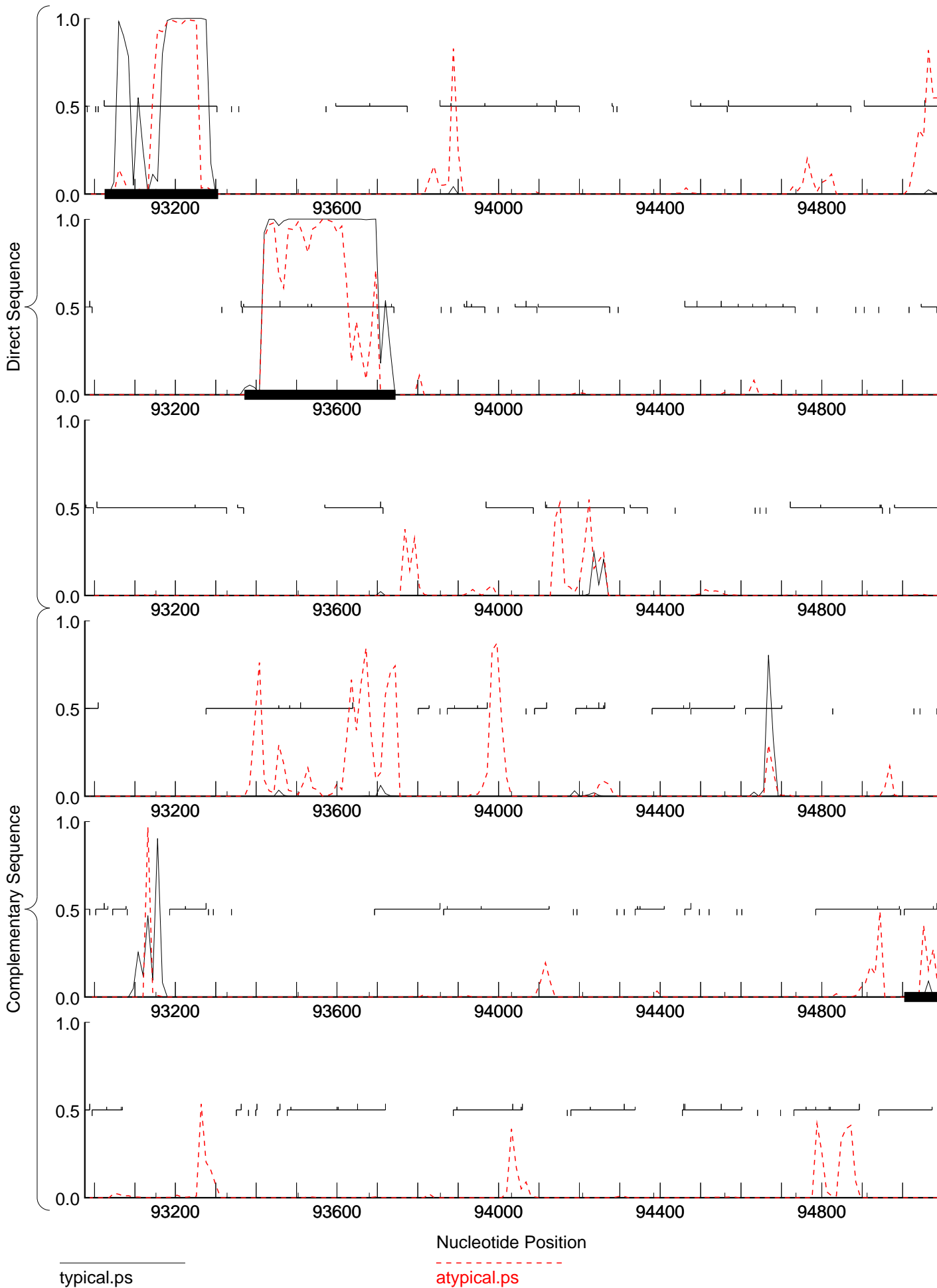


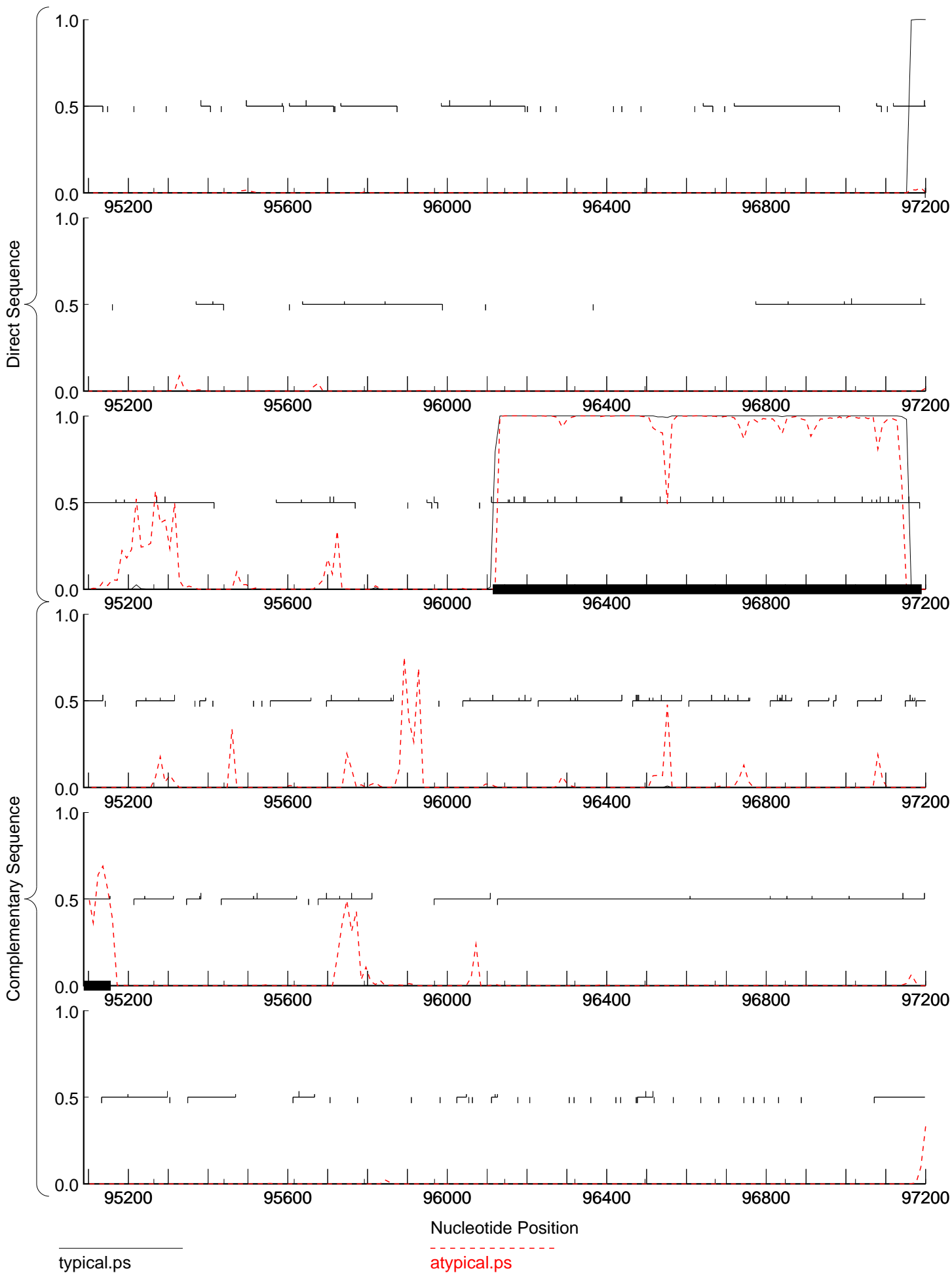


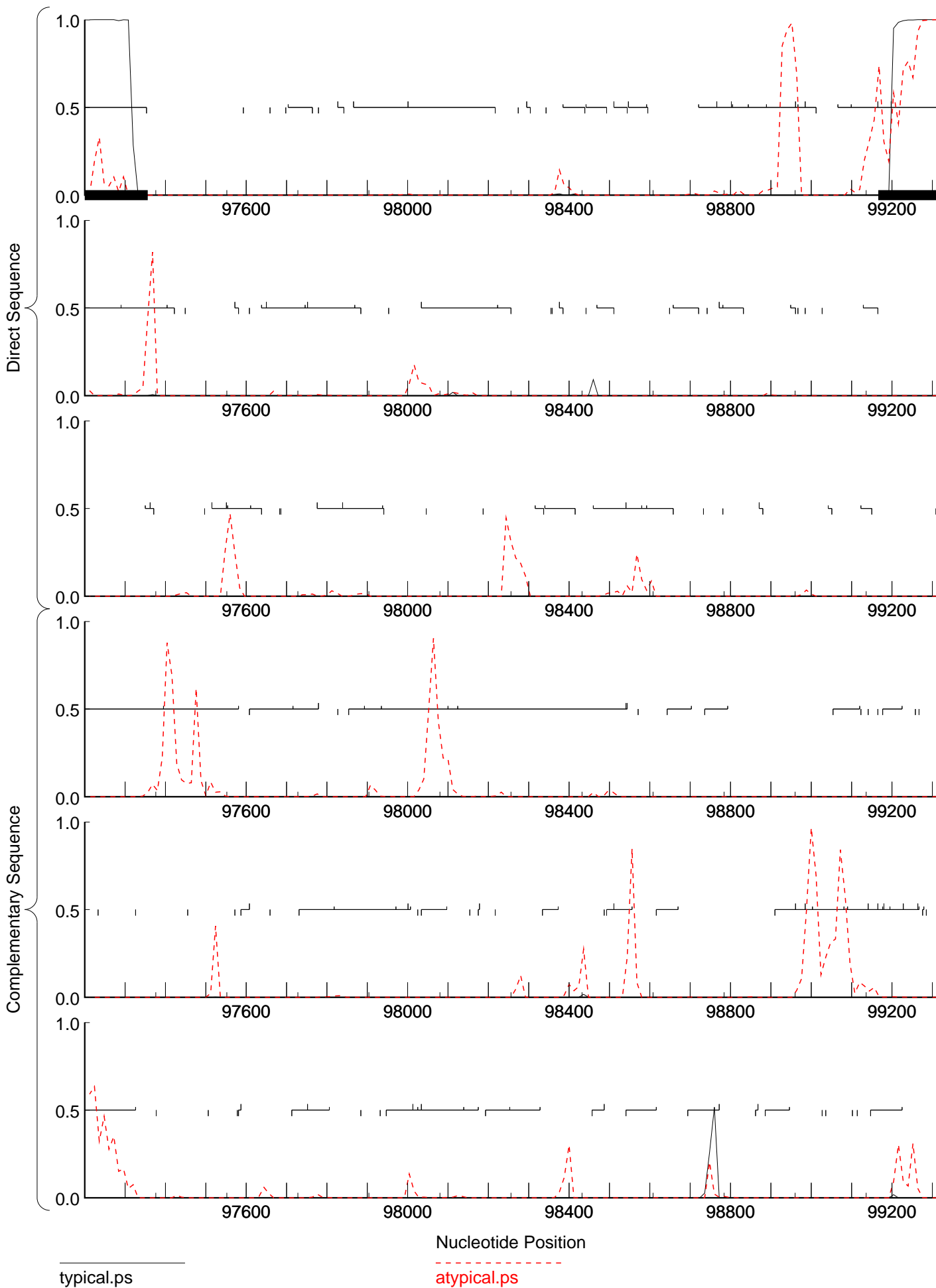


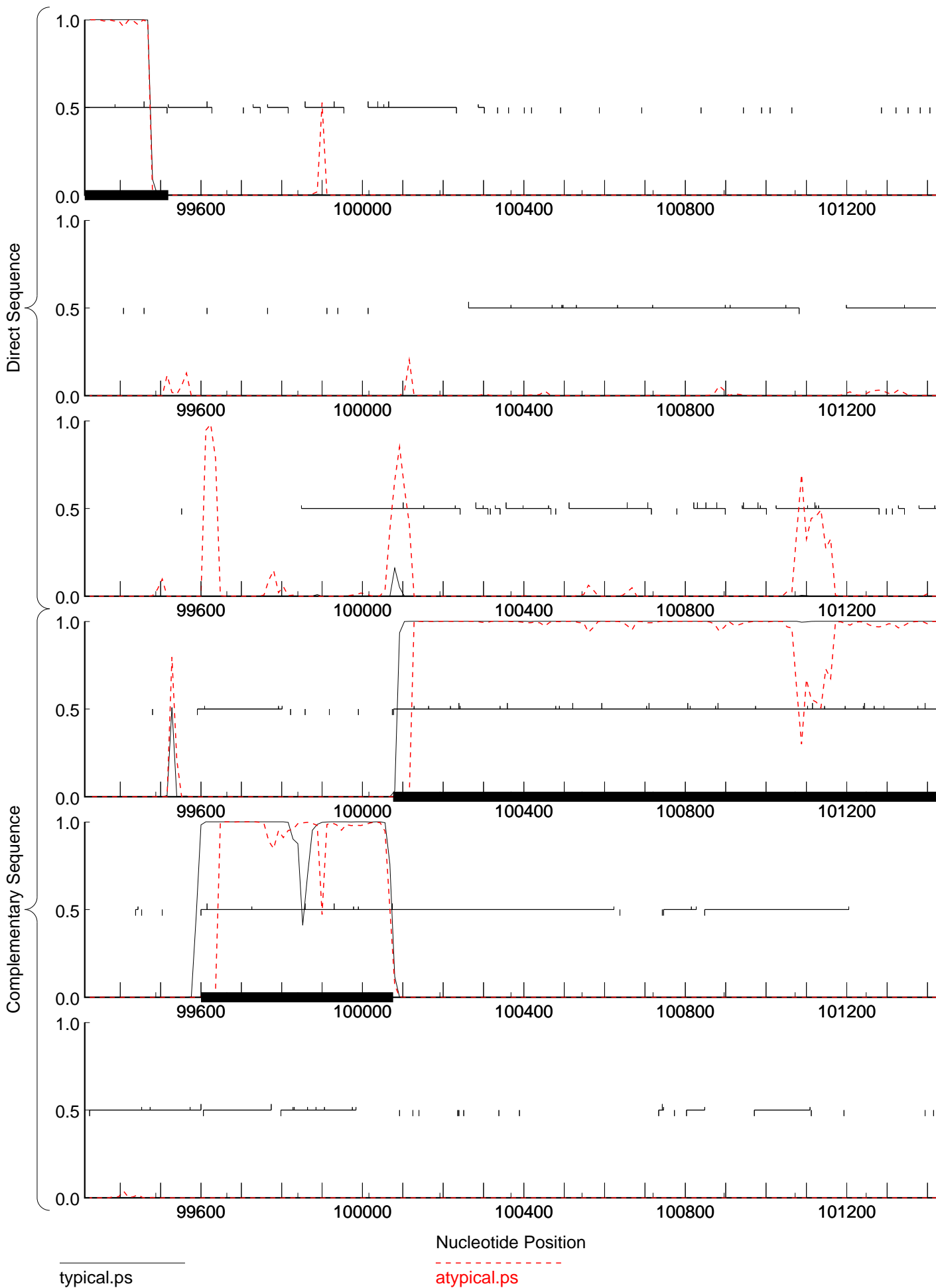
typical.ps

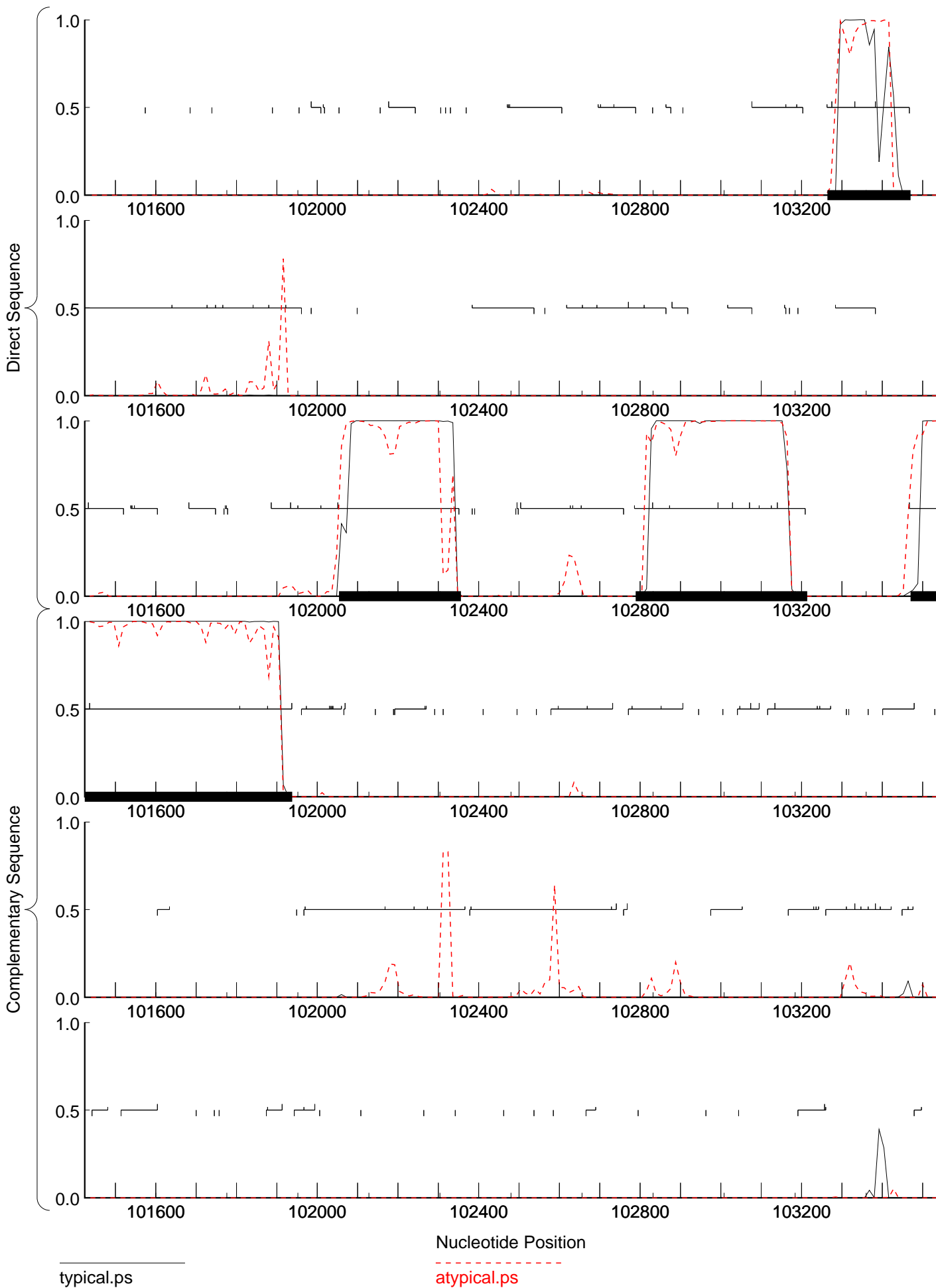
atypical.ps

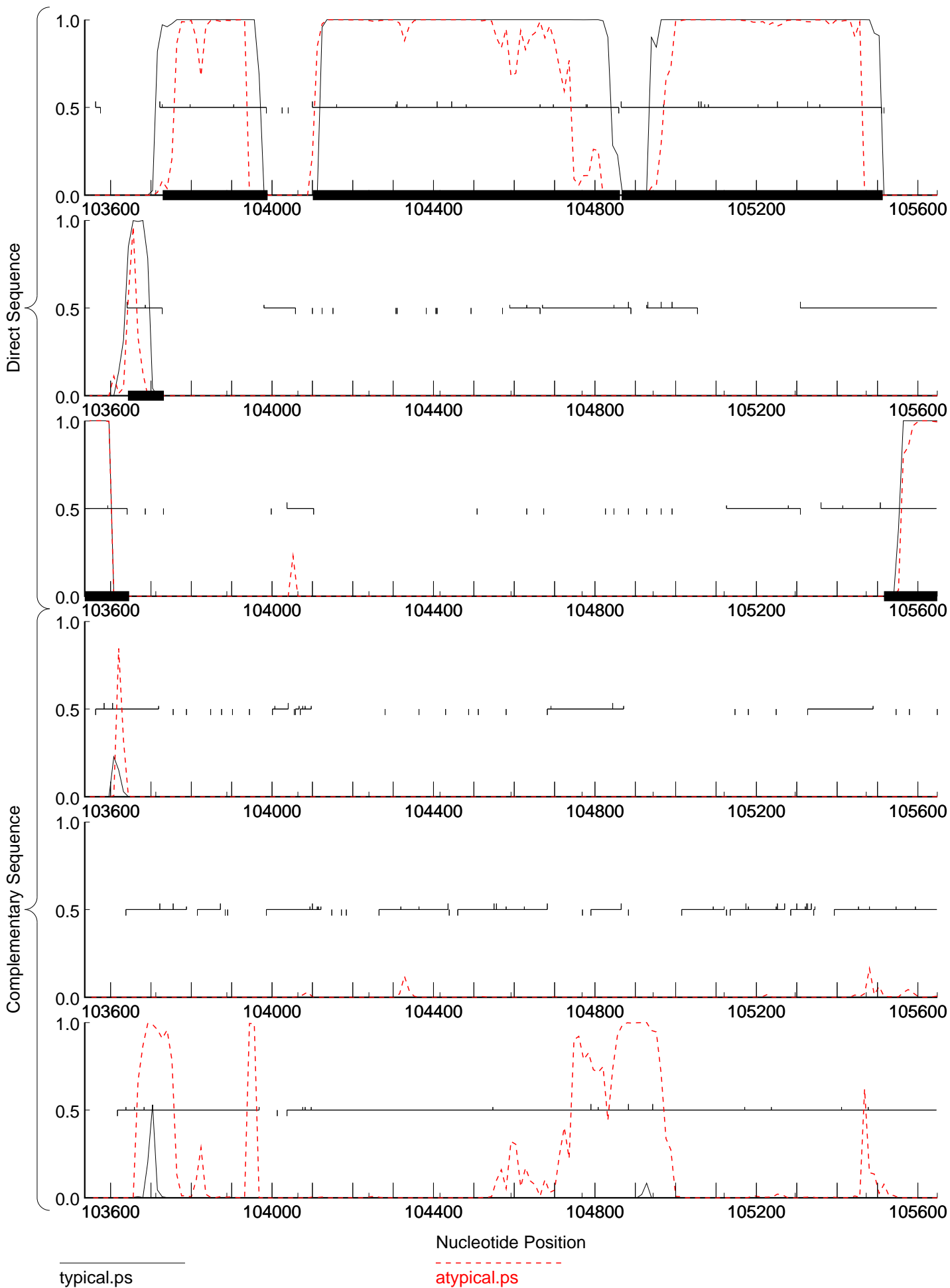


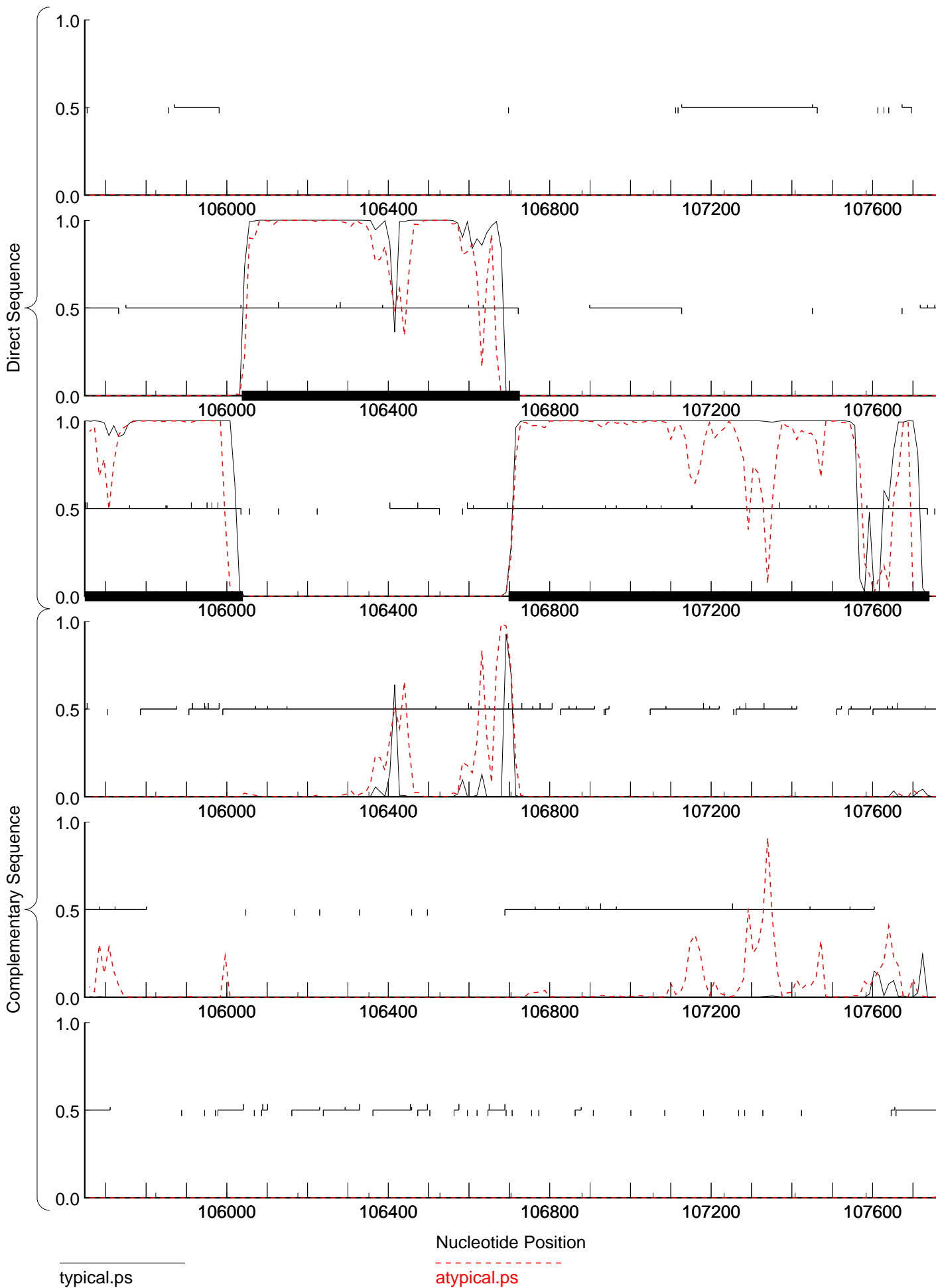


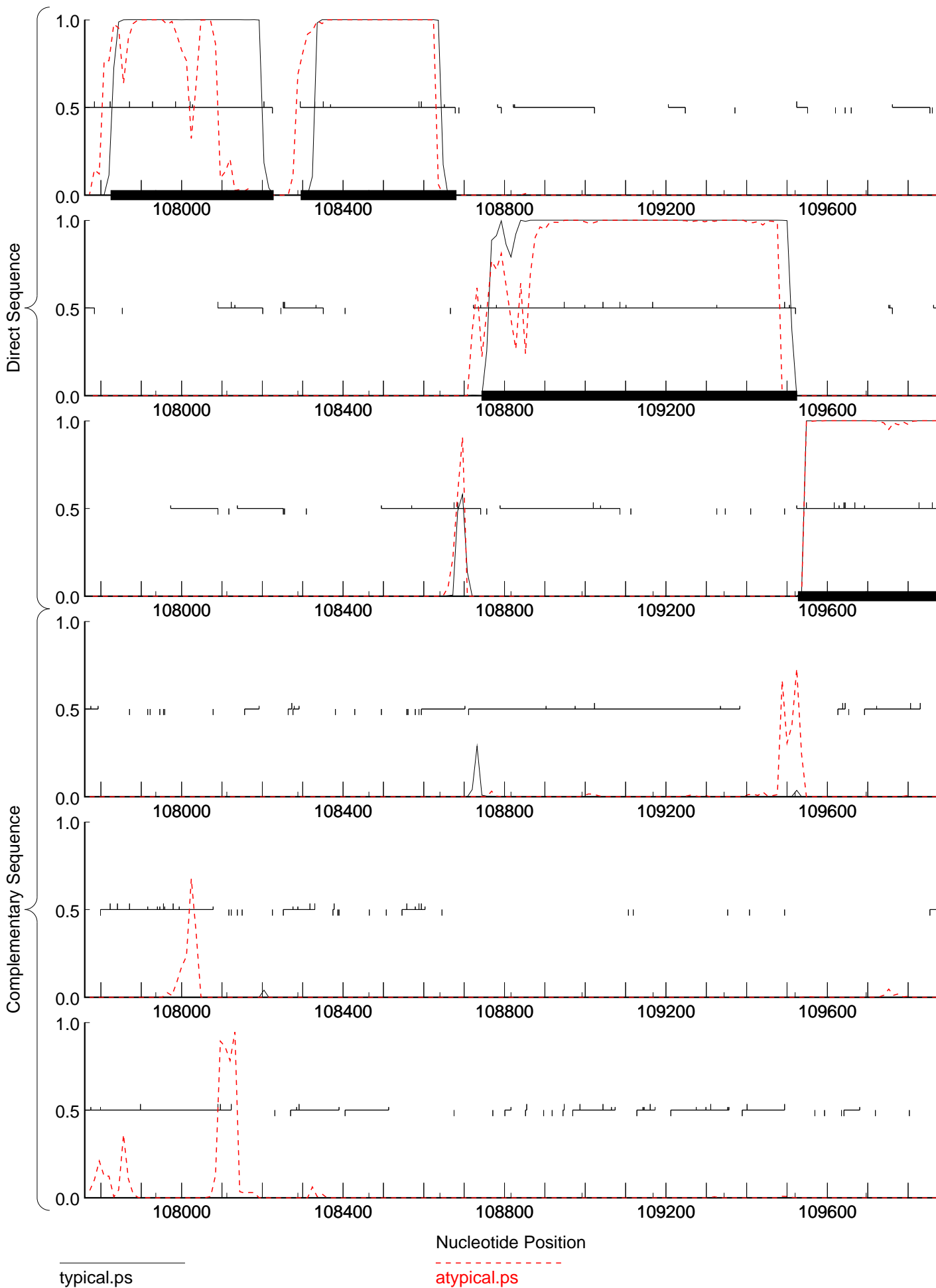


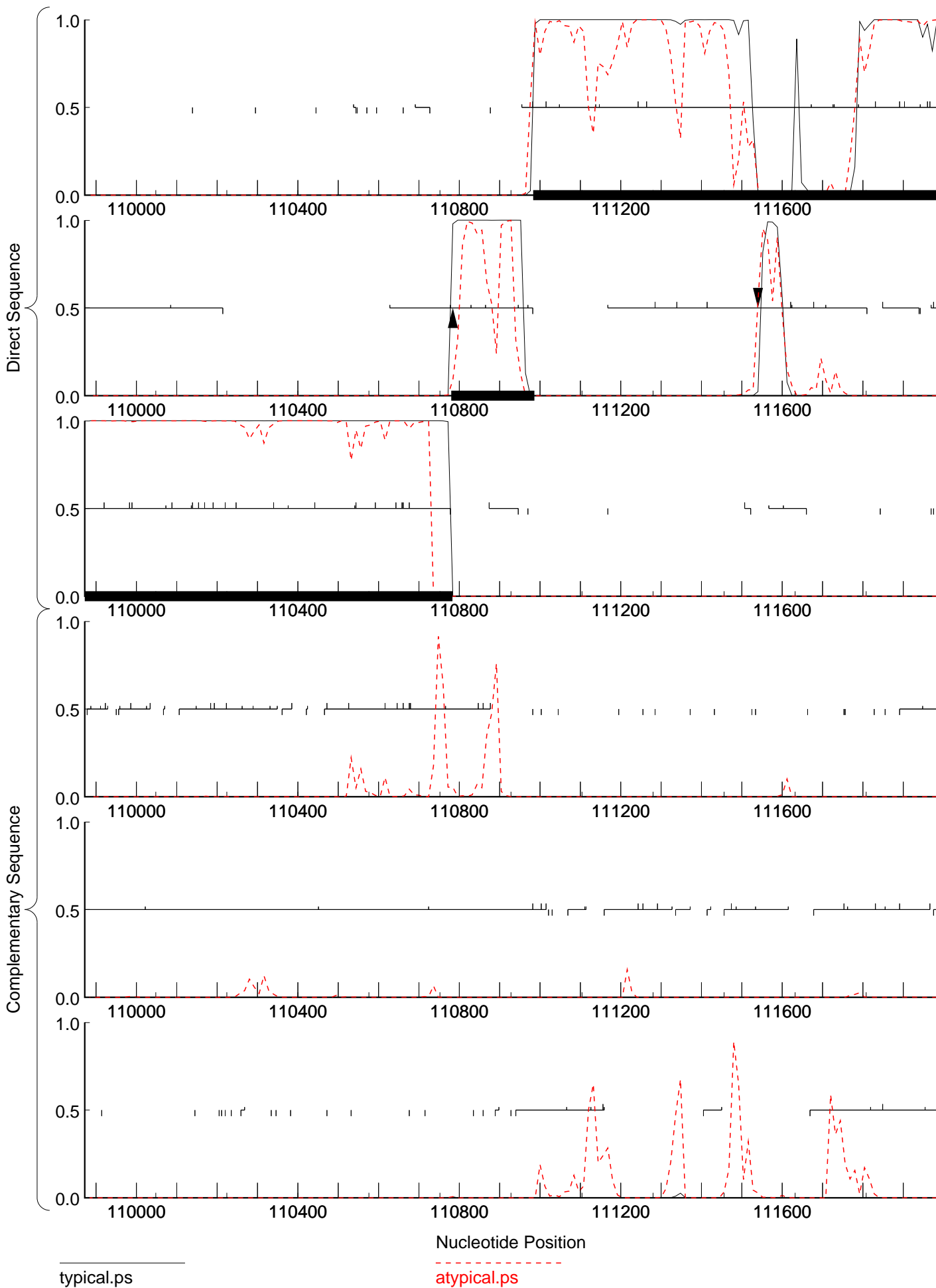


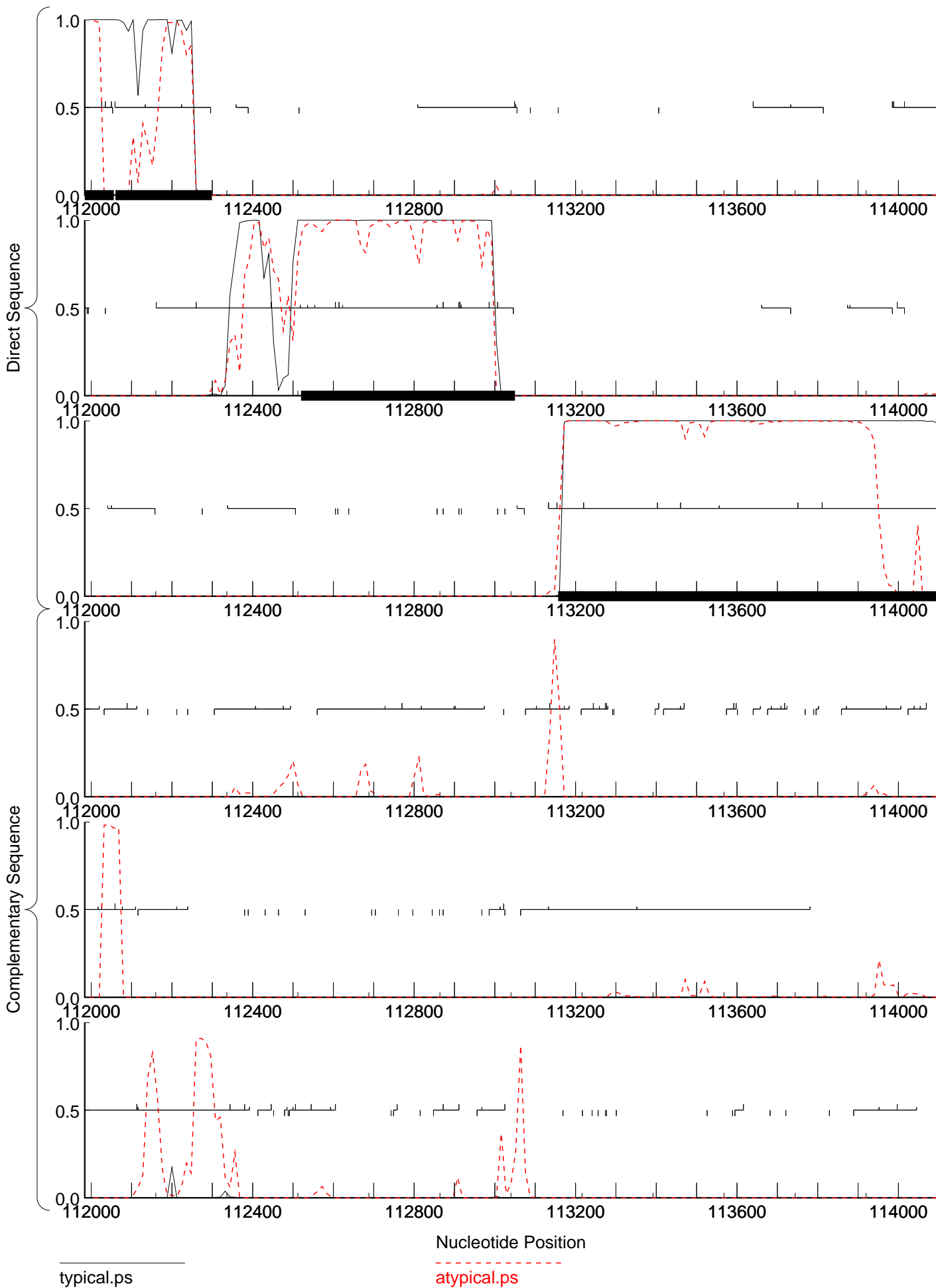


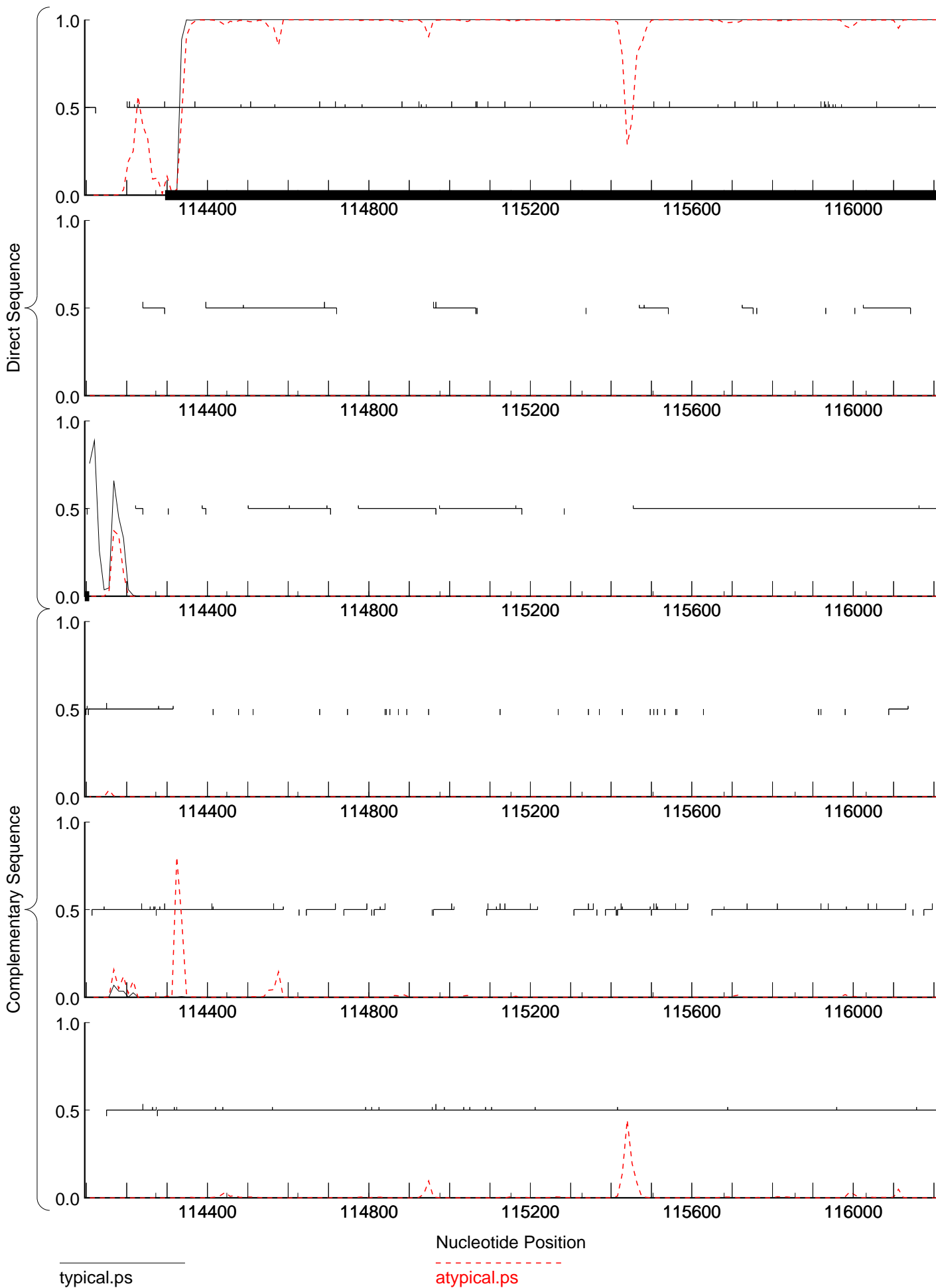


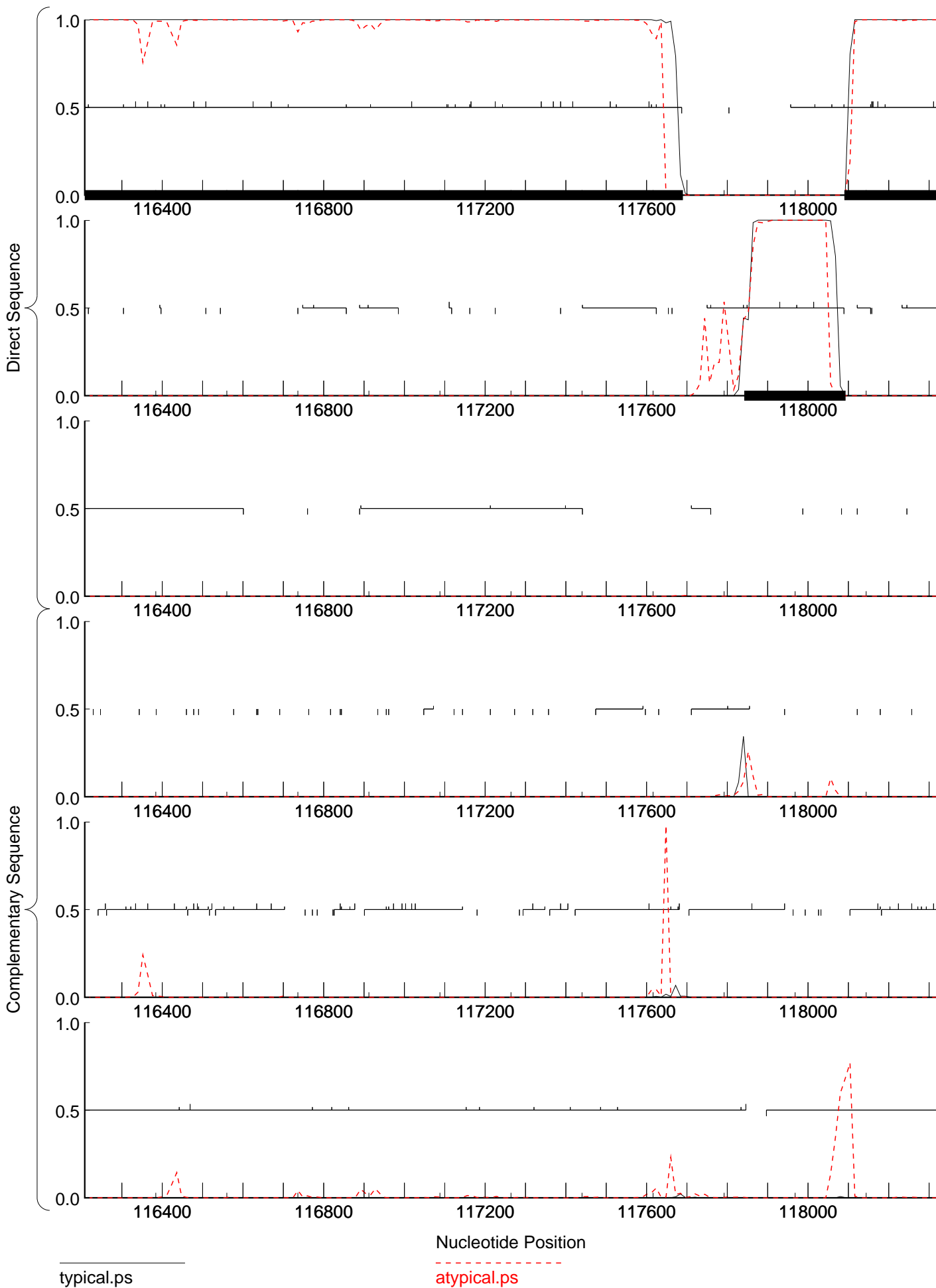


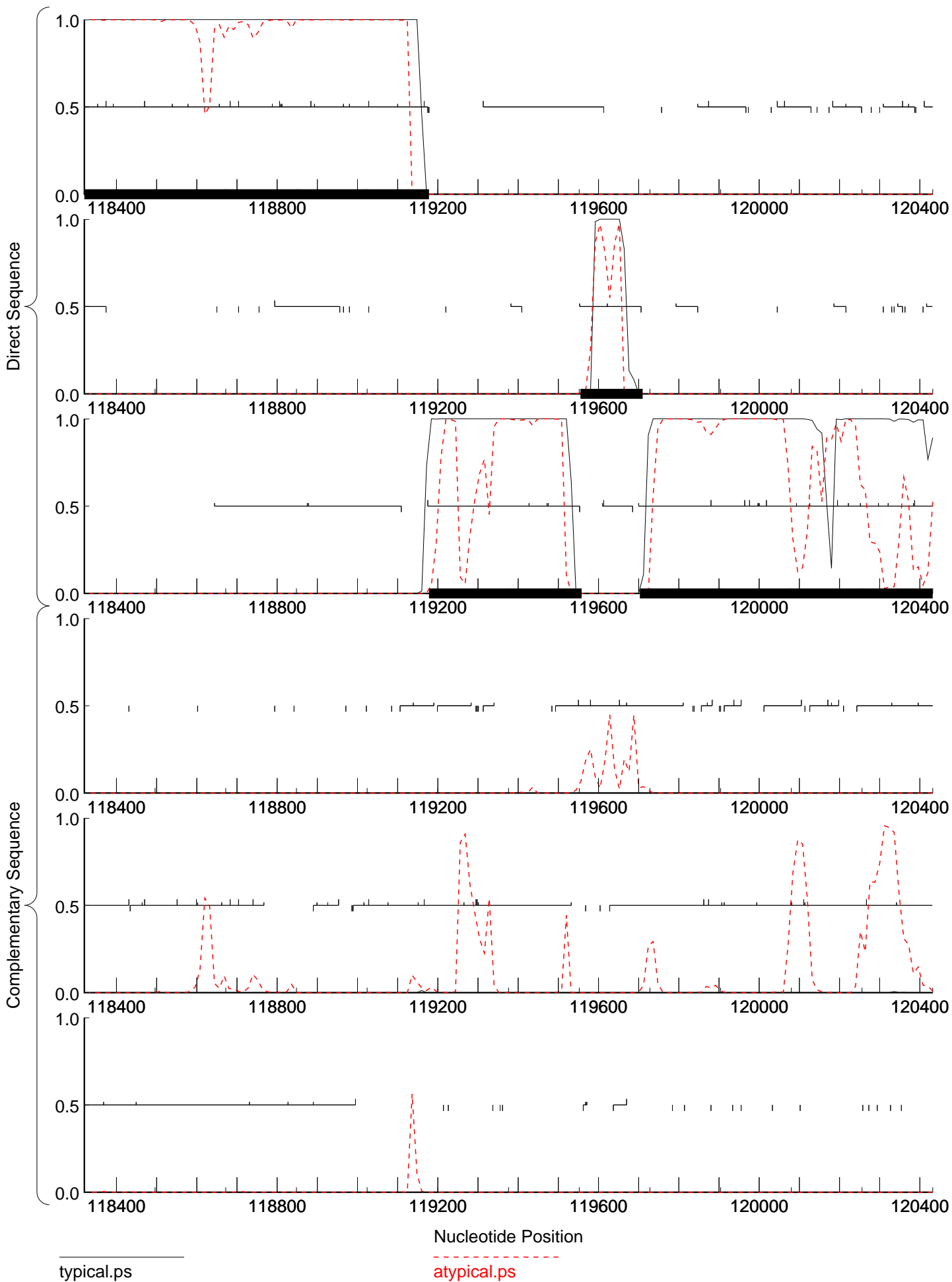


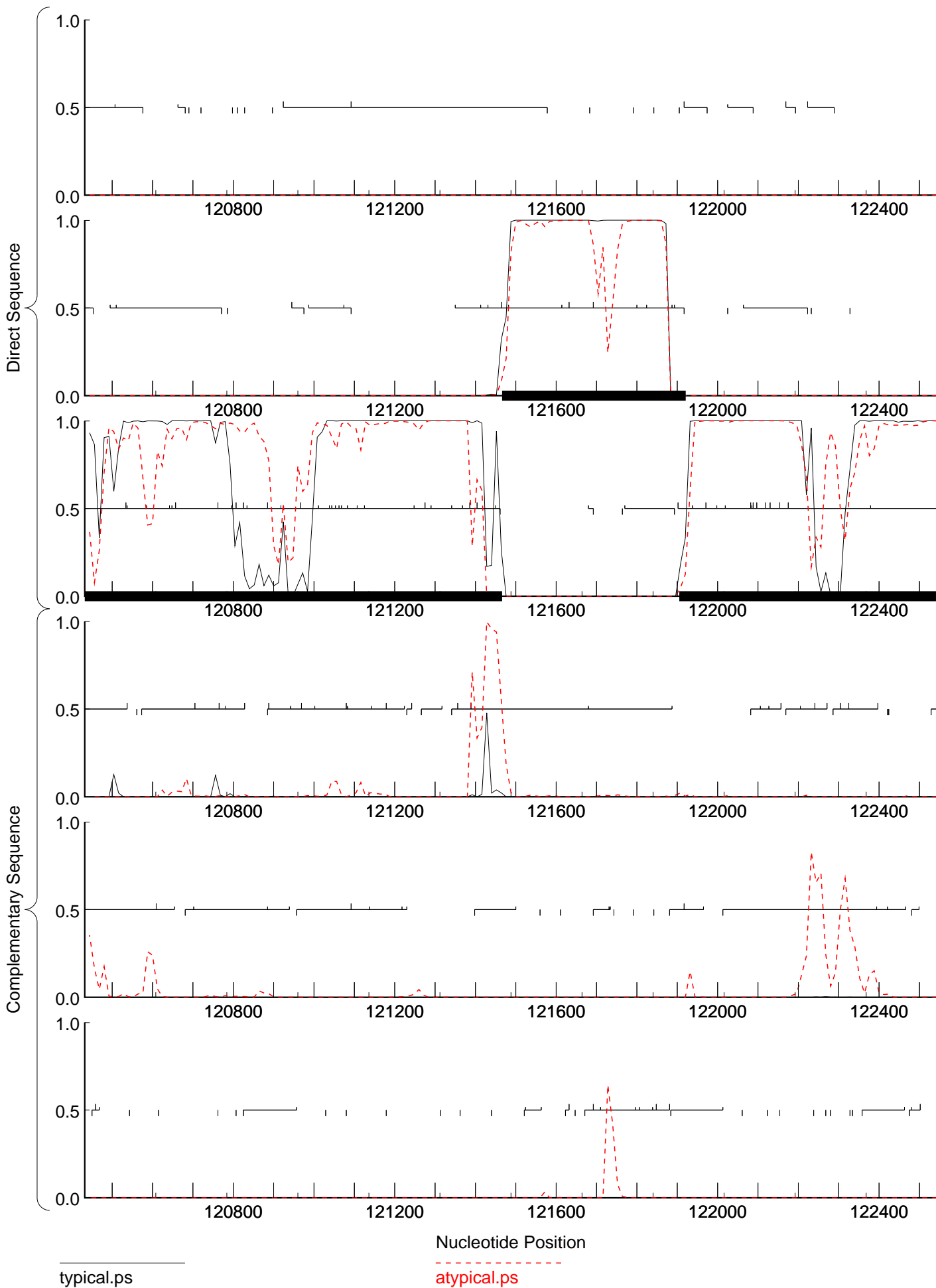


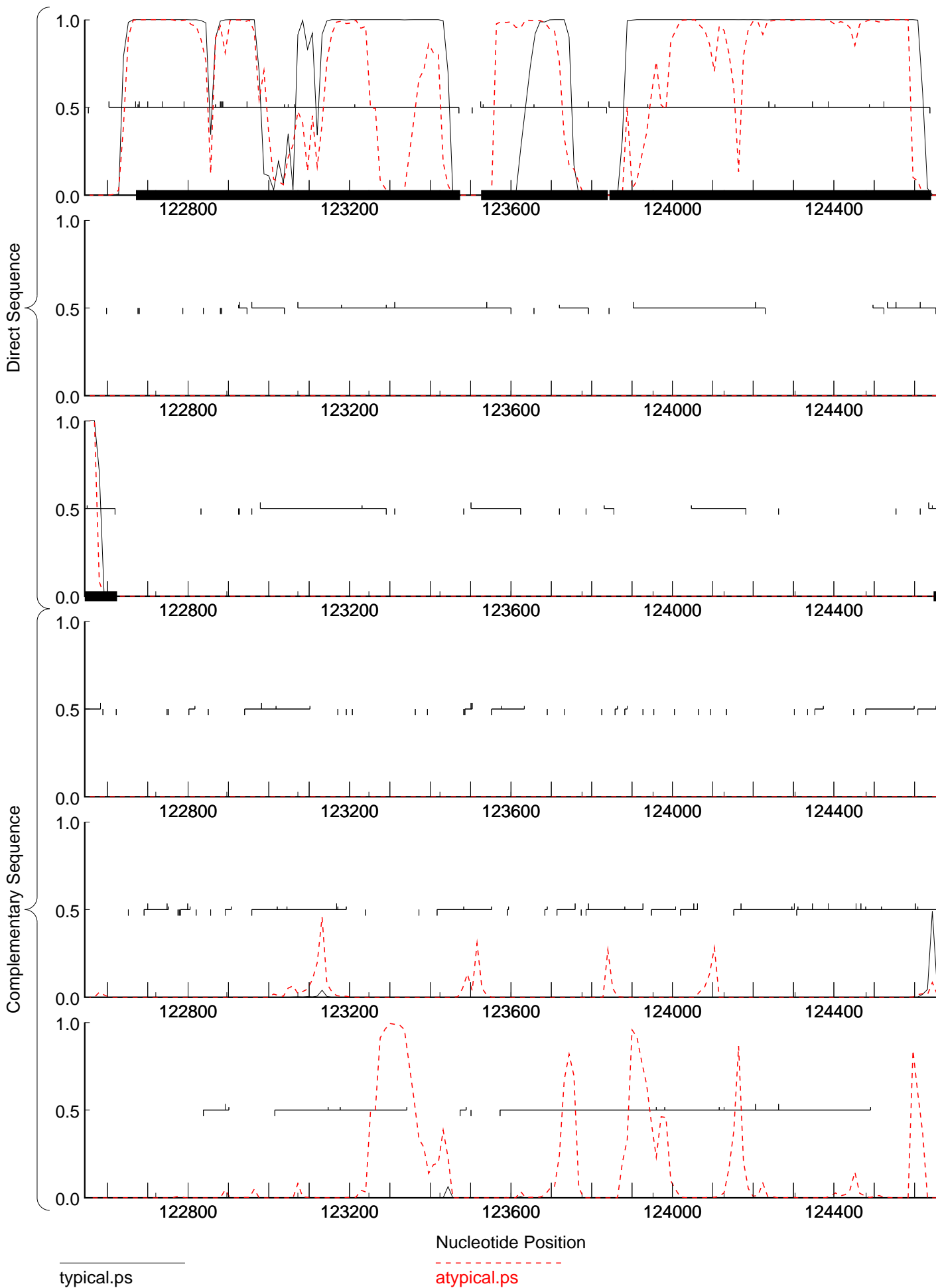






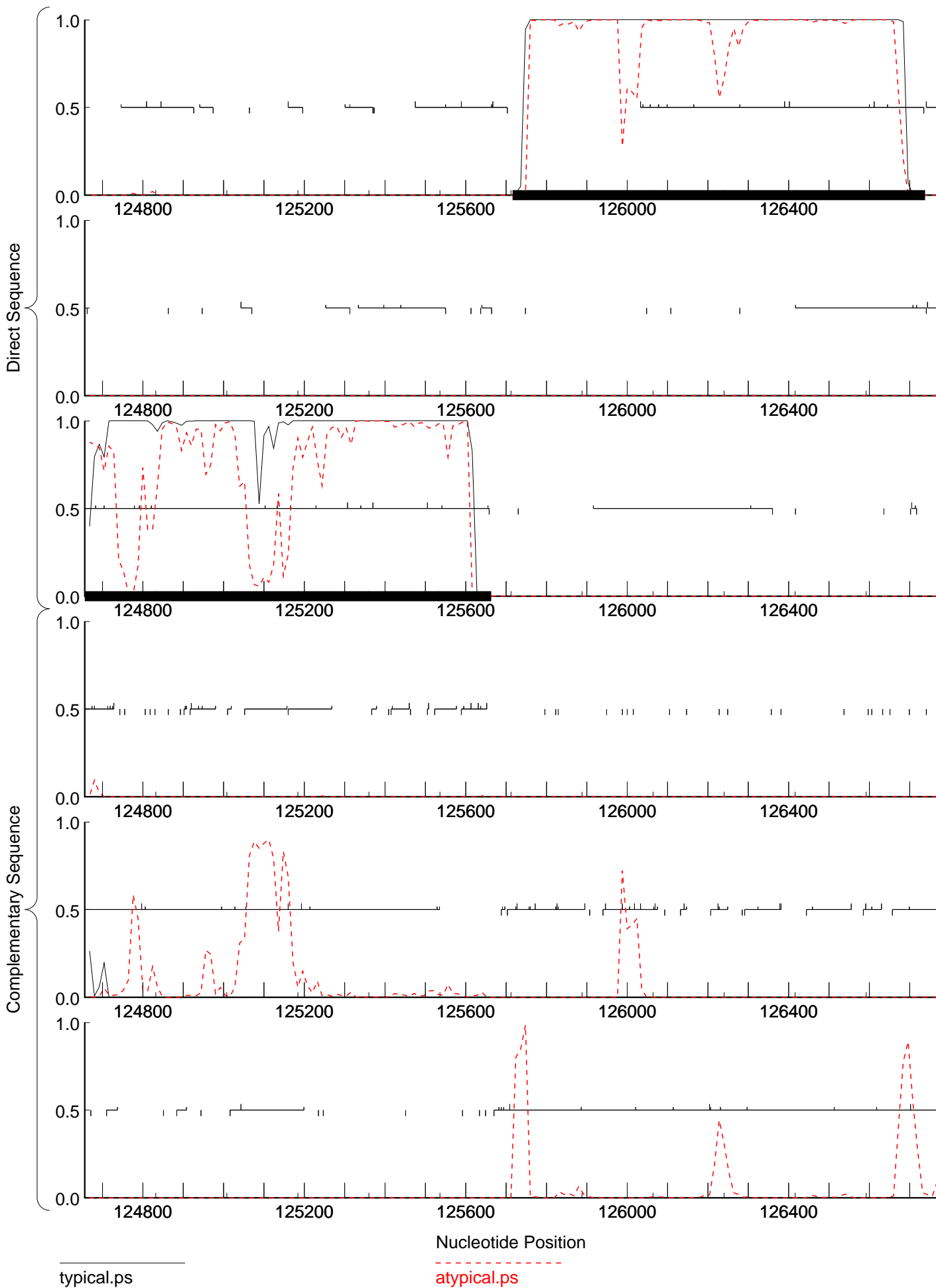


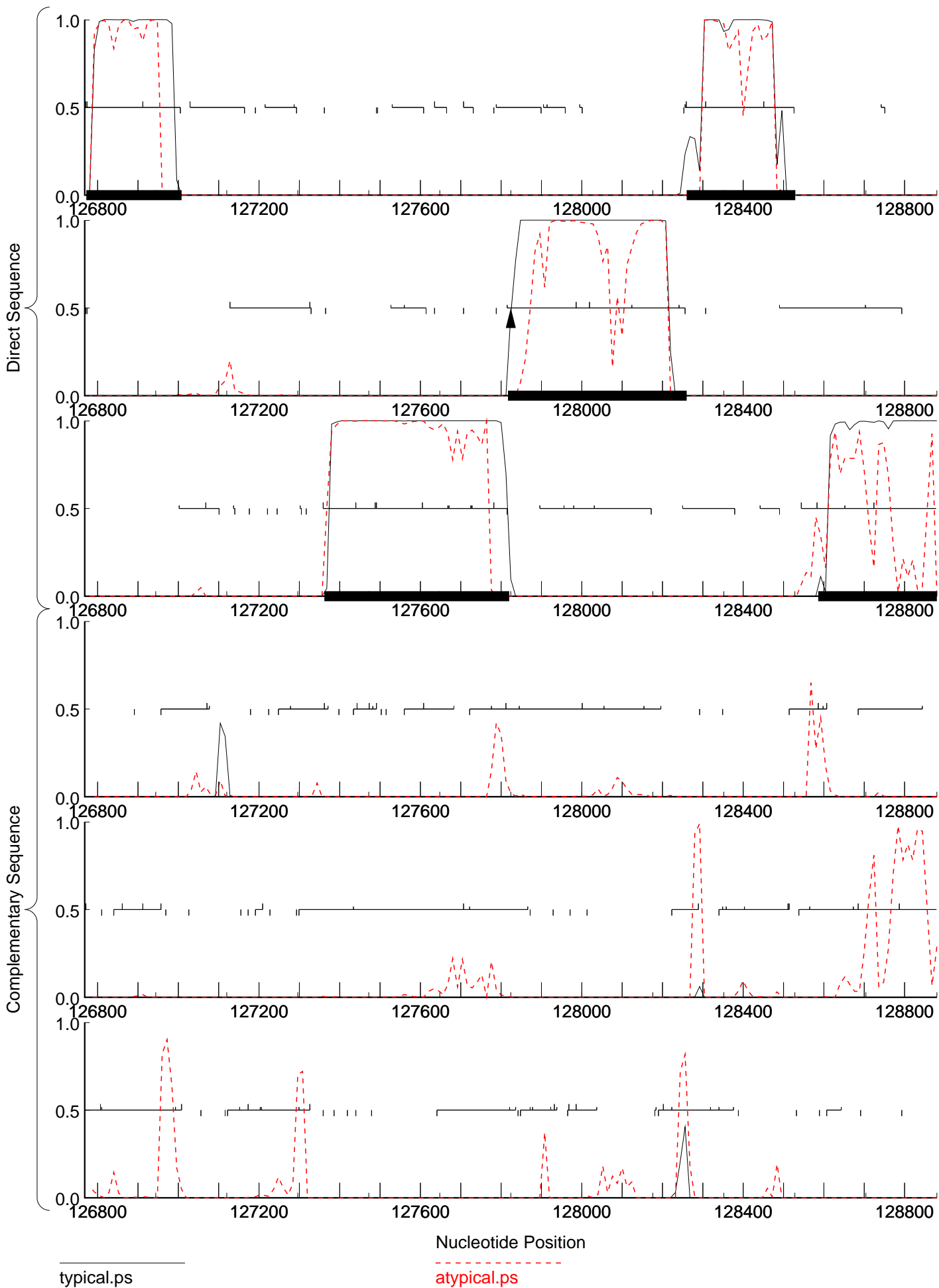


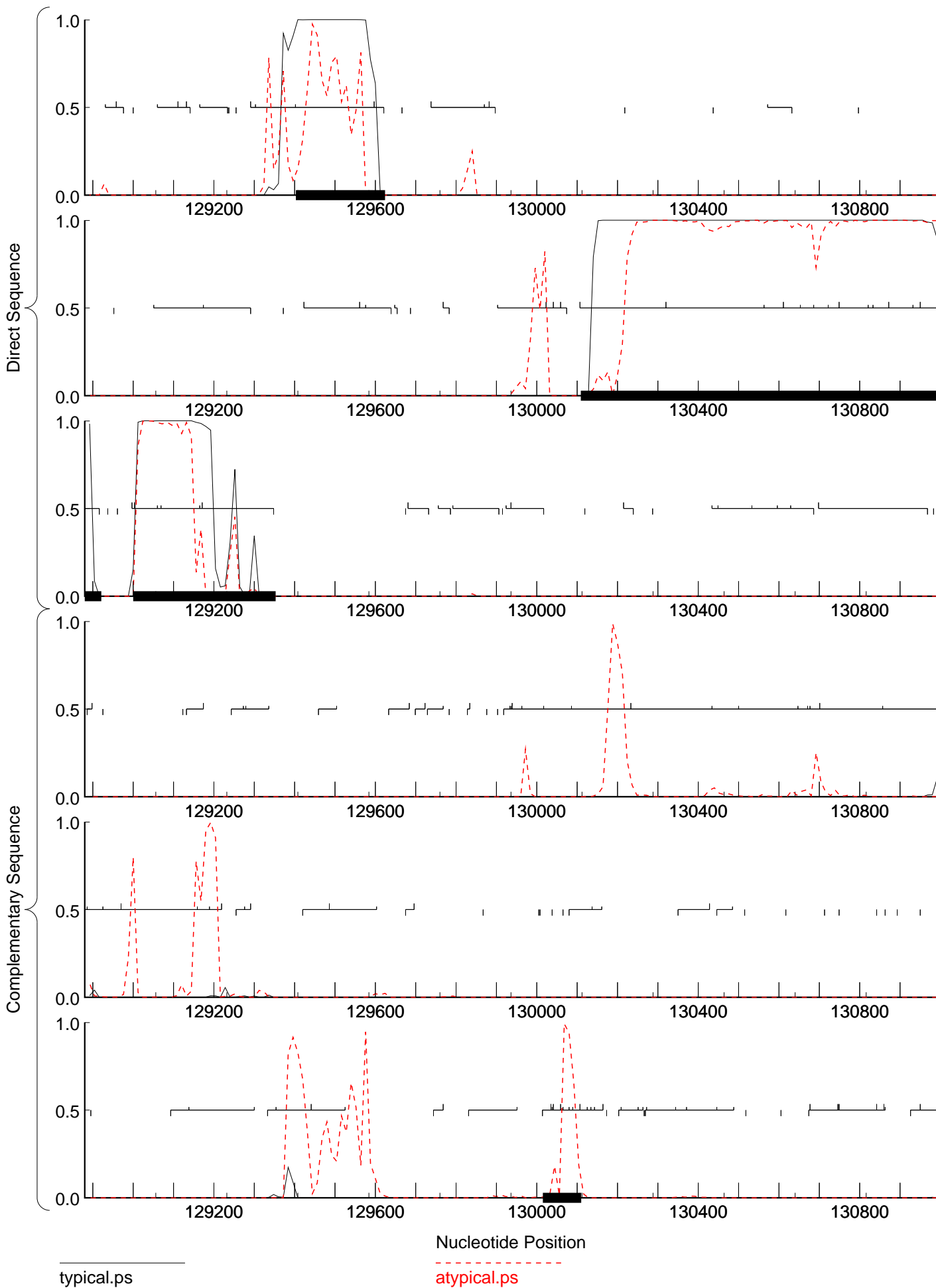


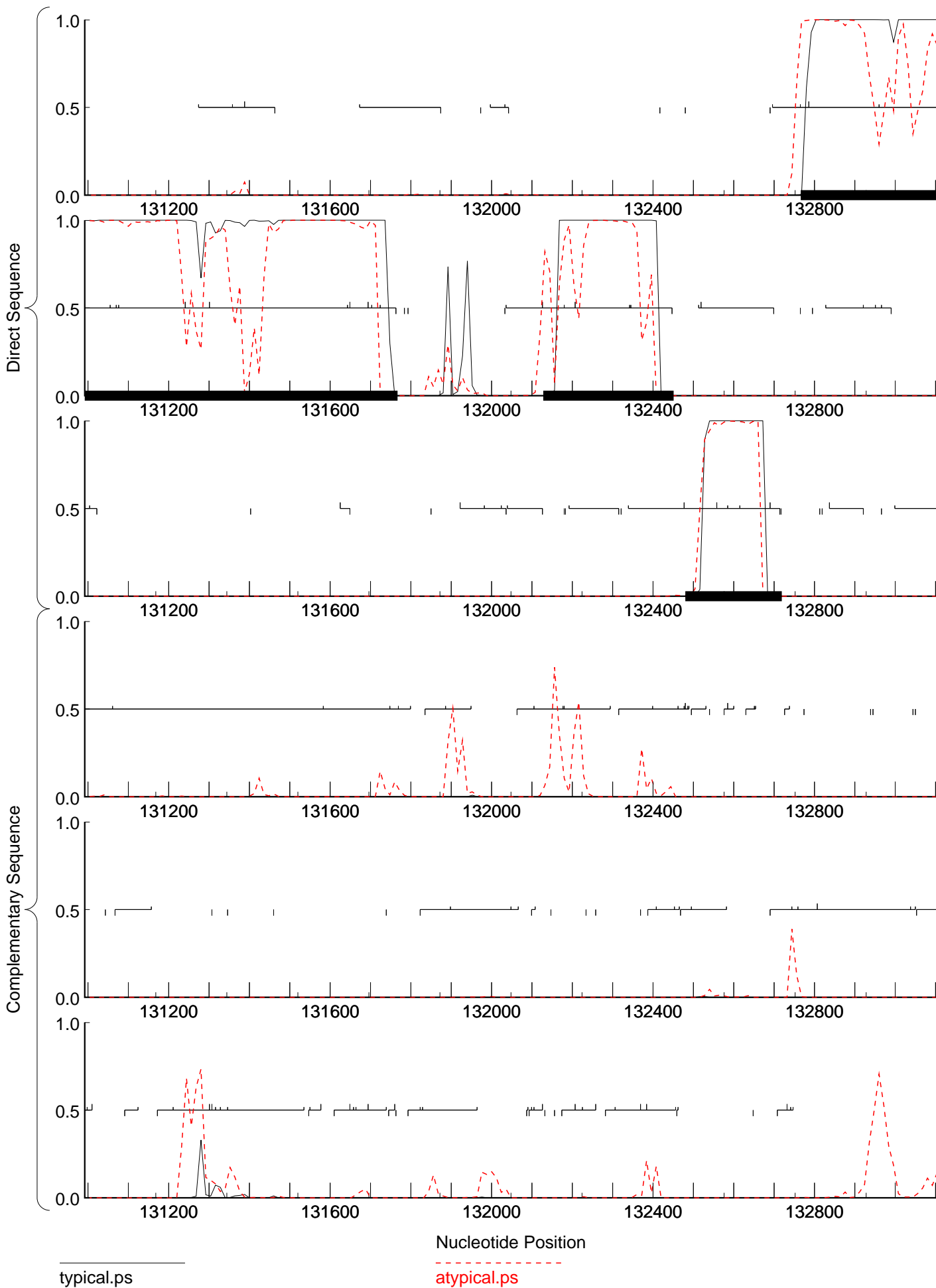
typical.ps

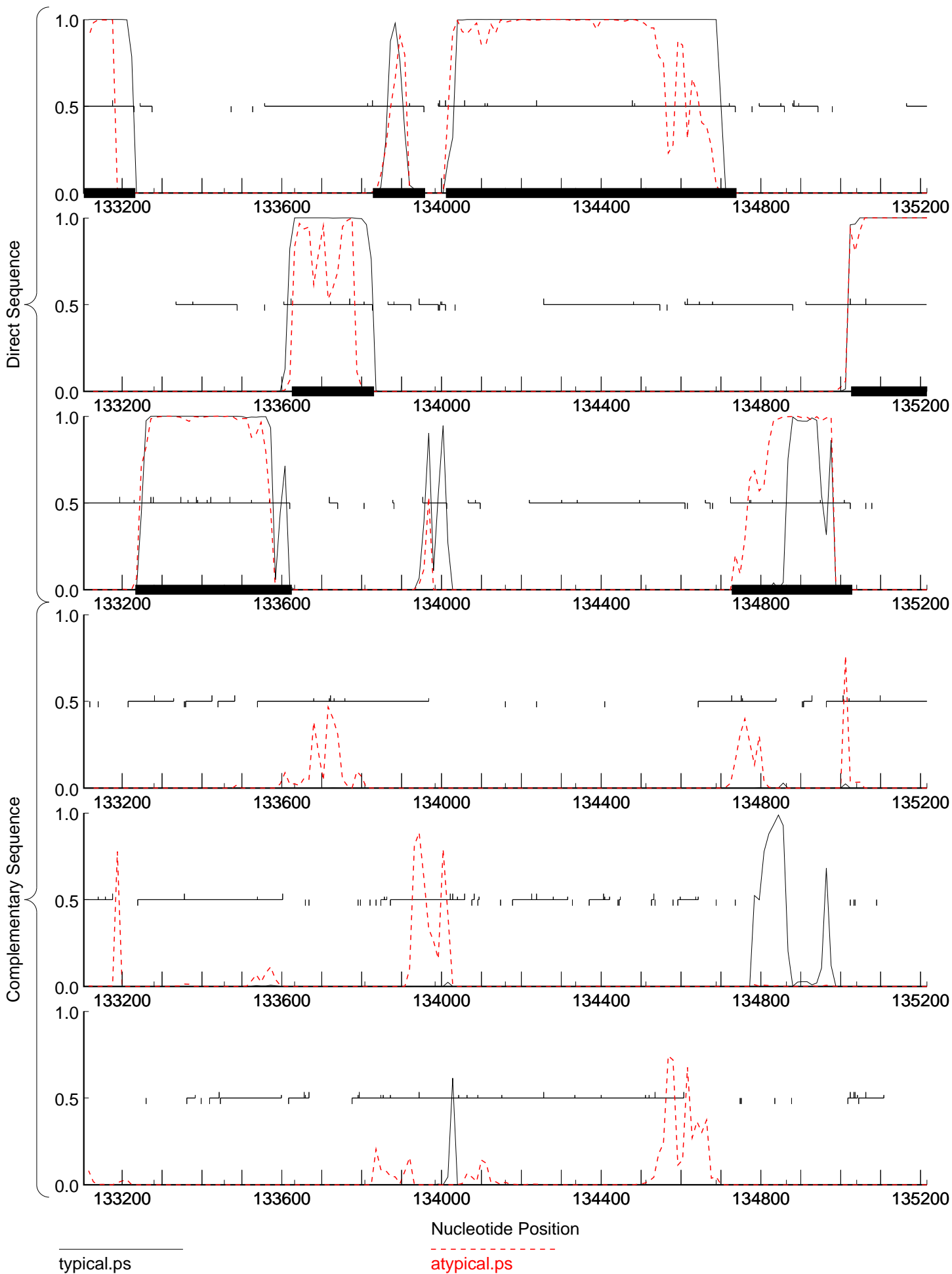
atypical.ps





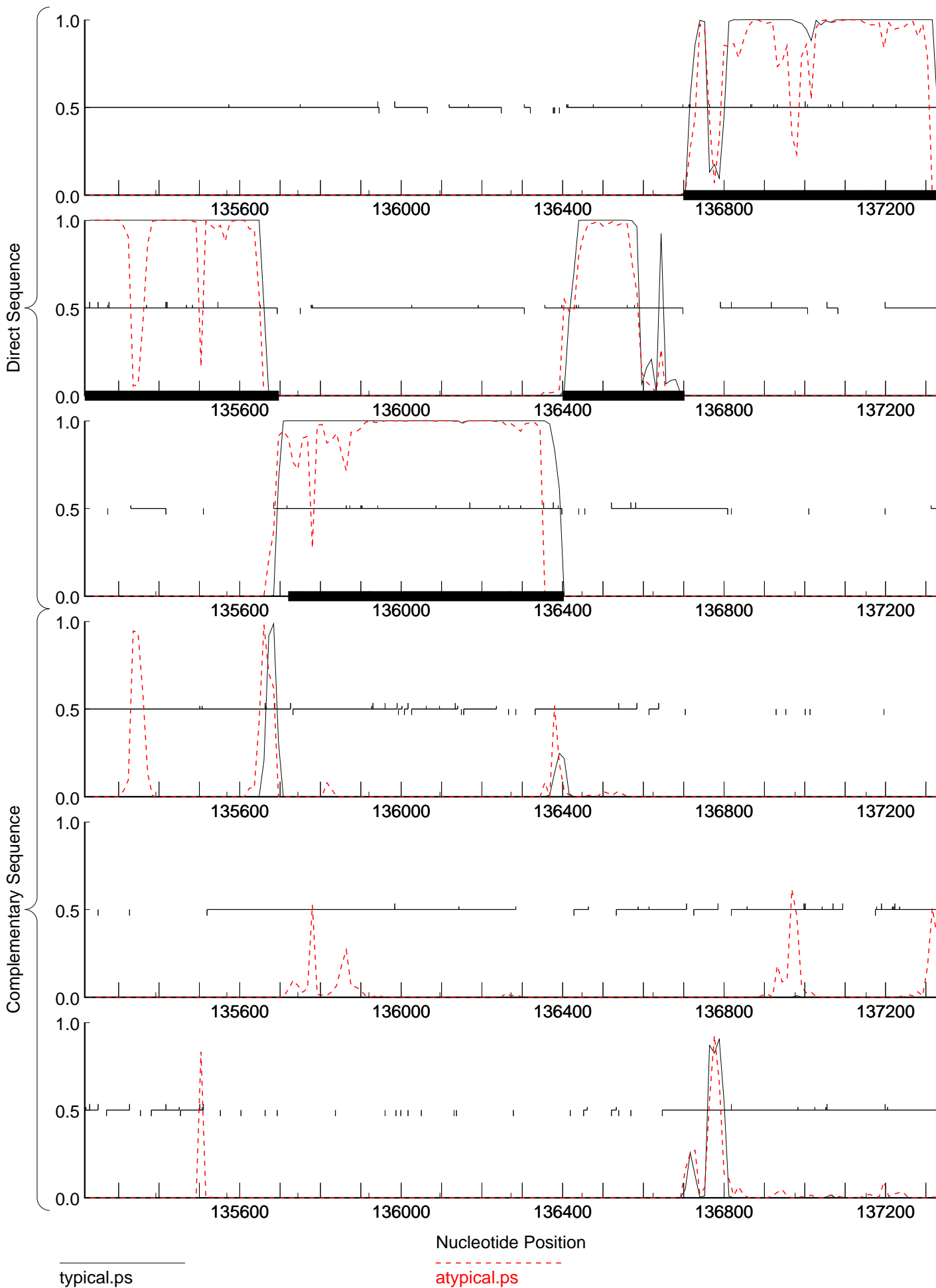


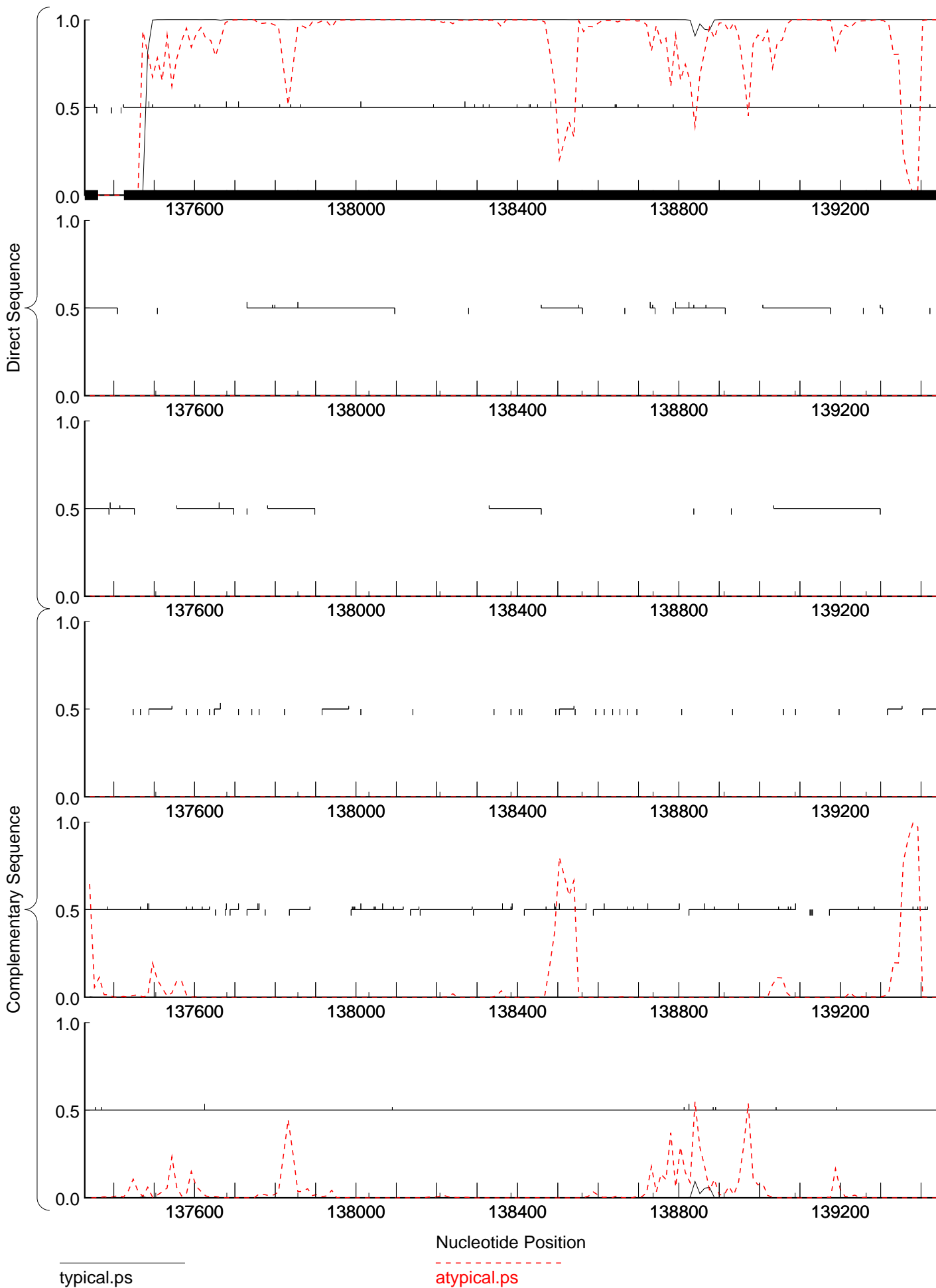


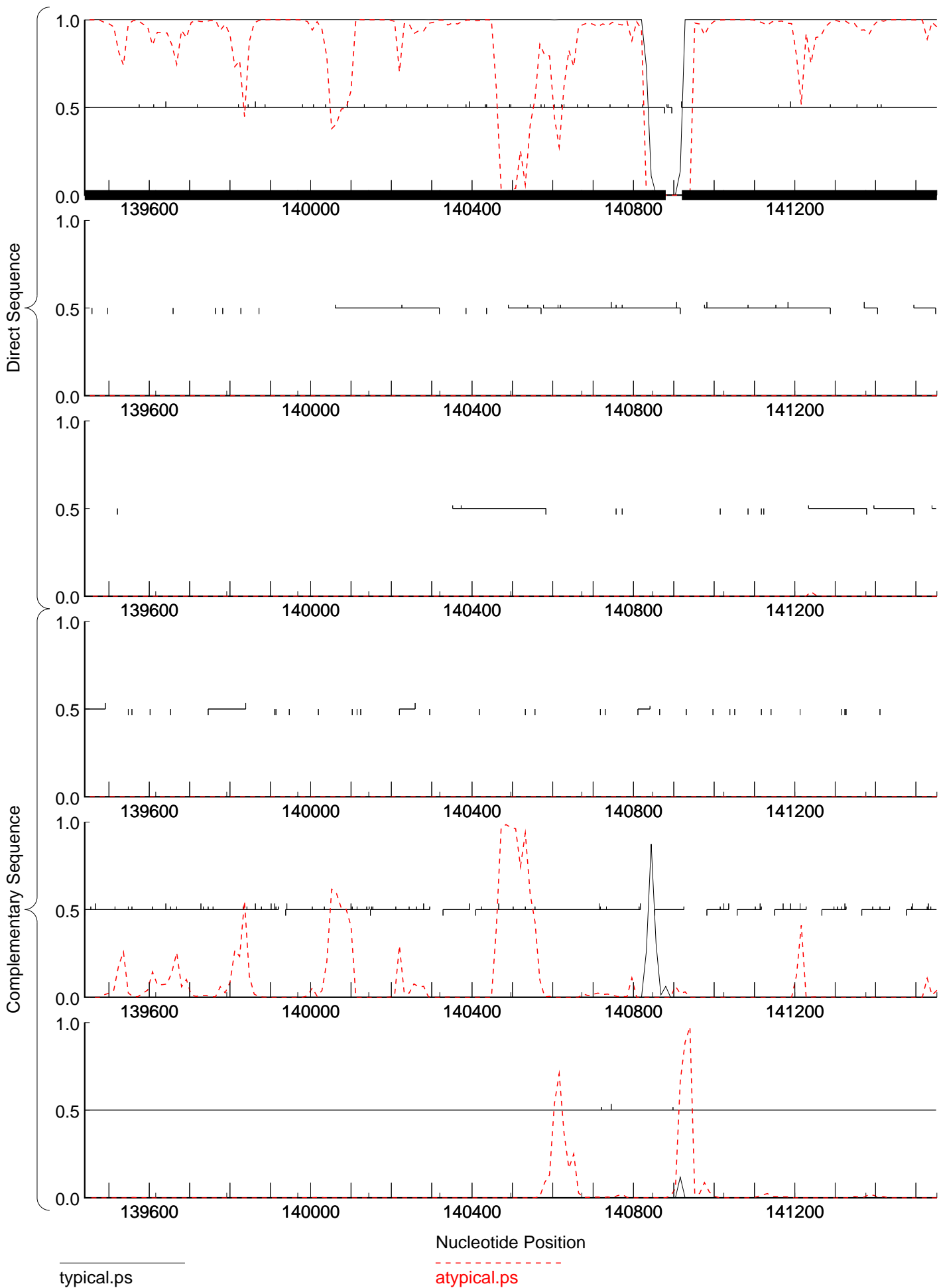


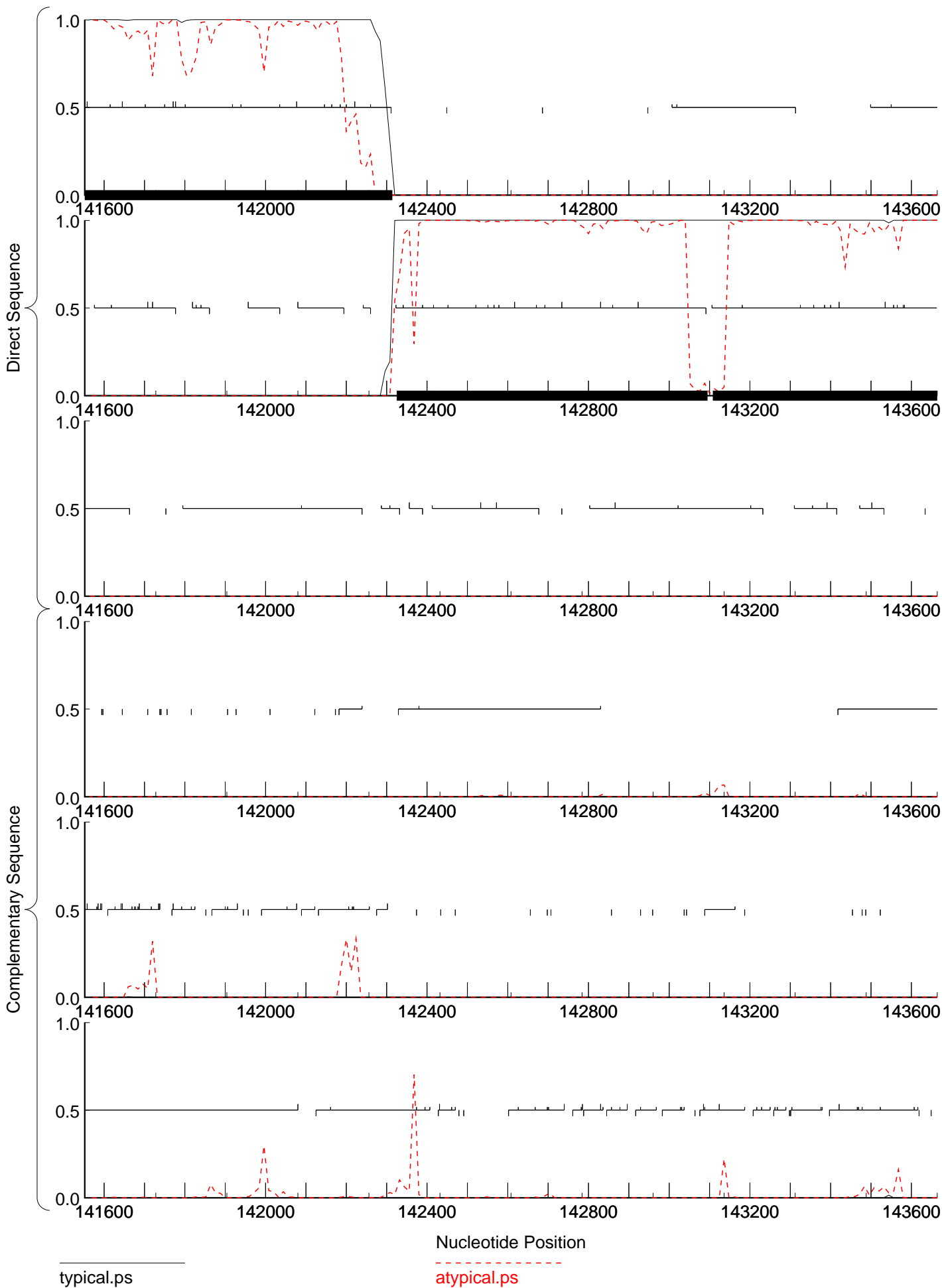
typical.ps

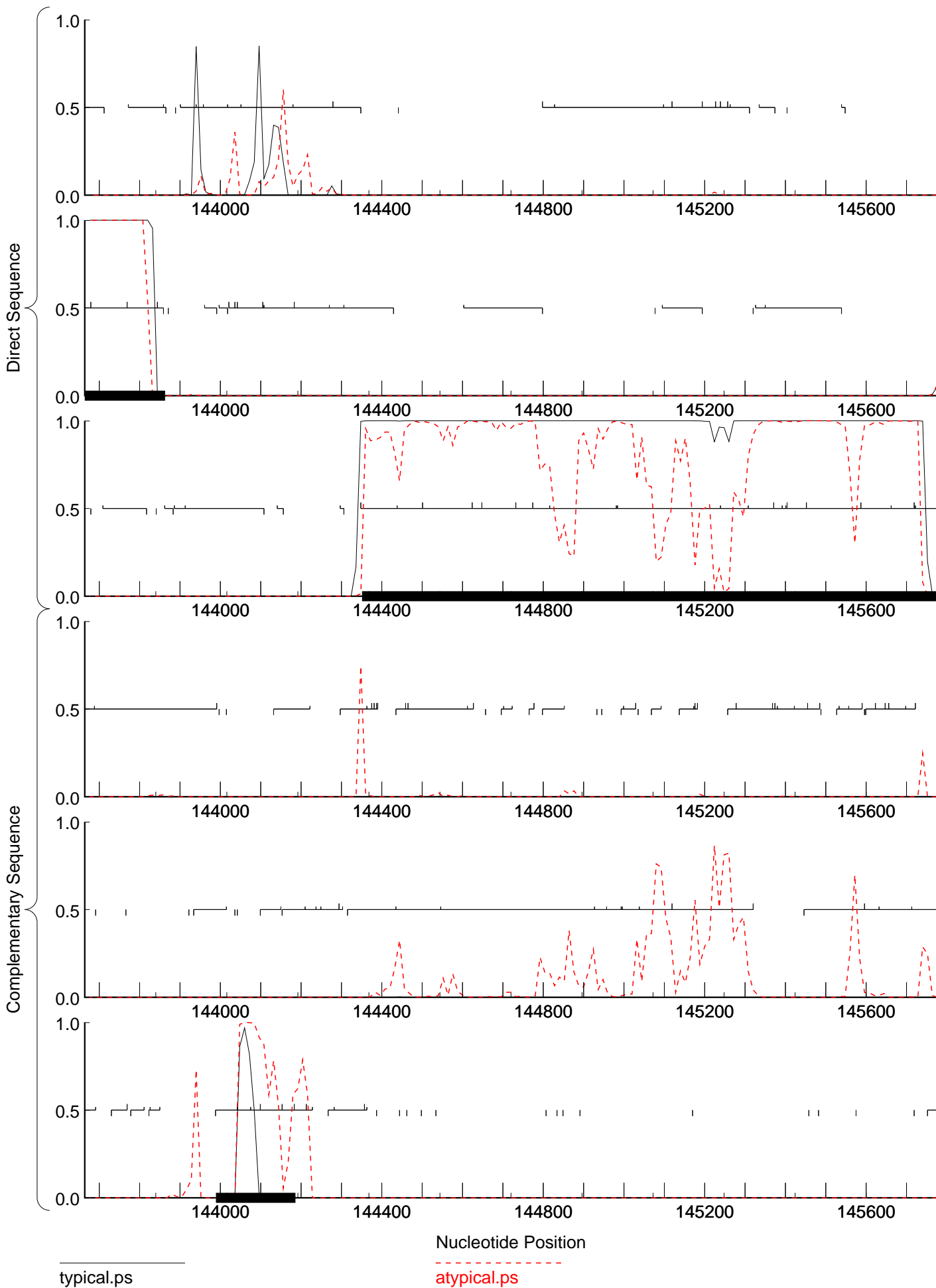
atypical.ps

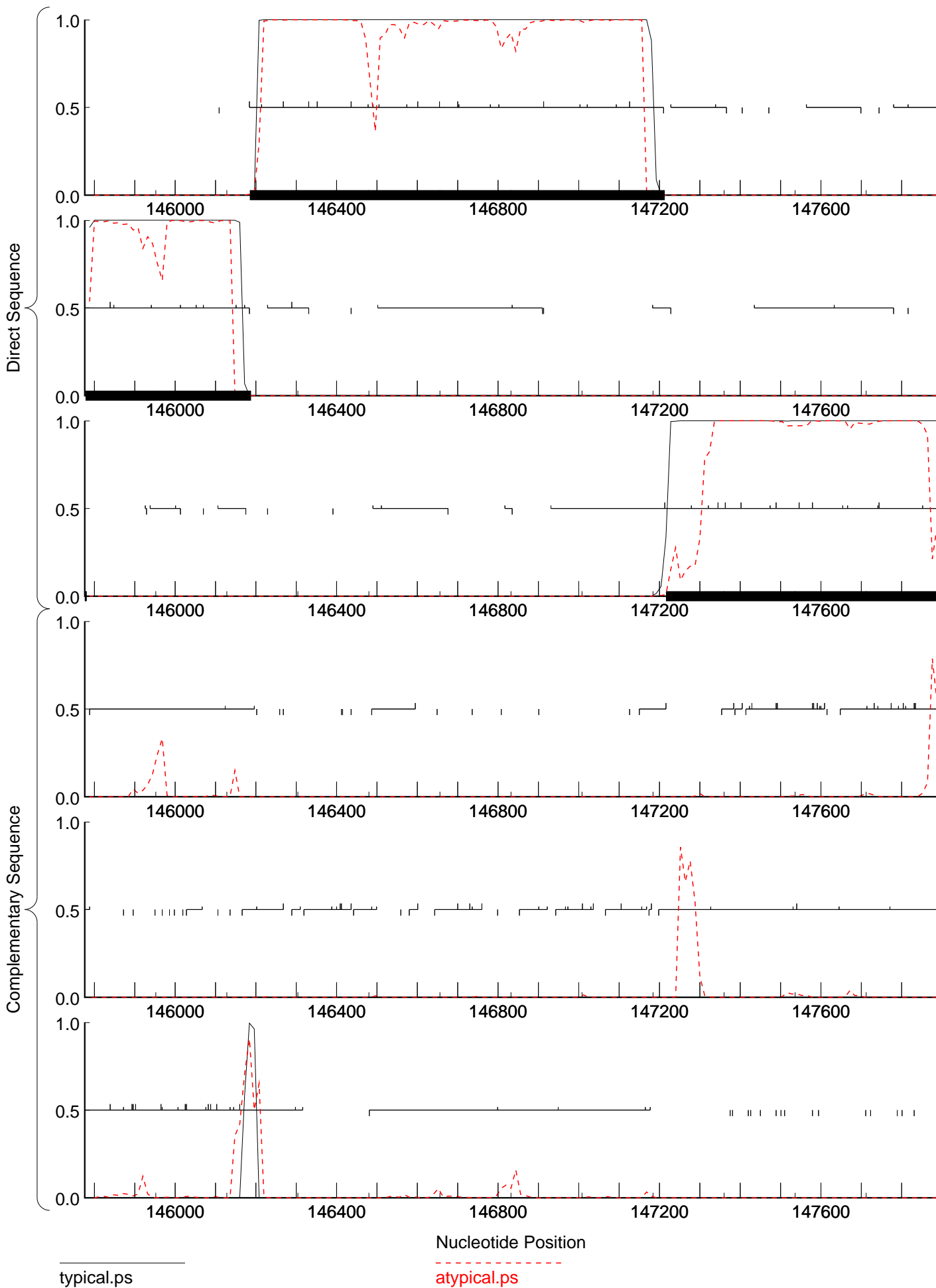


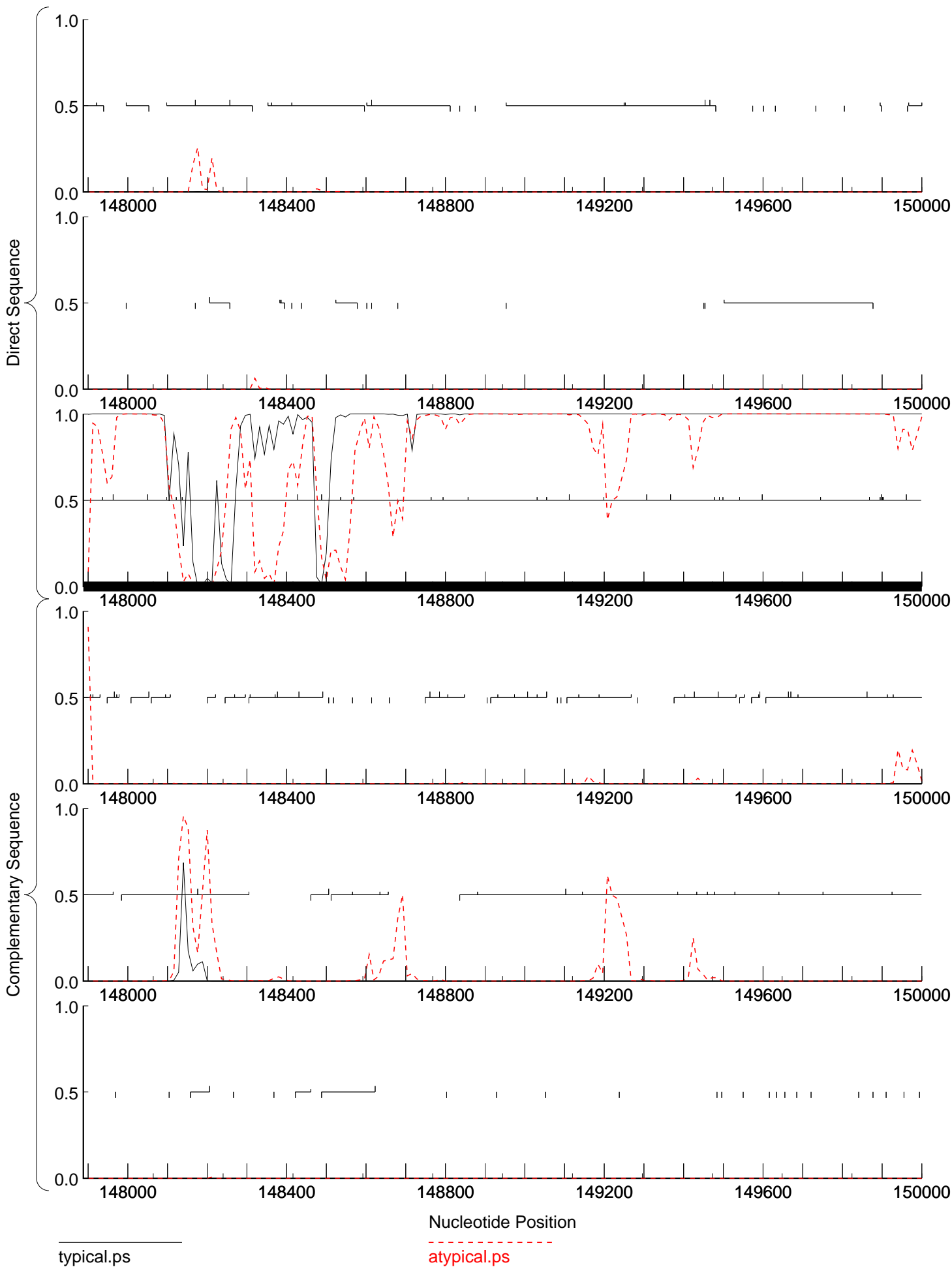












typical.ps

atypical.ps

