

GeneMark

Version 2.5p (09.08.06)

Copyright 1993 - M. Borodovsky, J. McIninch

PROGRAM INFORMATION

Sequence : Mycobacterium phage AltPhacts complete sequence, 68056 bp, circularly permuted, Cluster B1
Analysis Date : 8/18/17 at 18:16:12
Pages : 34
Sequence Length : 68056 bp
GC Content : 66.47%

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,

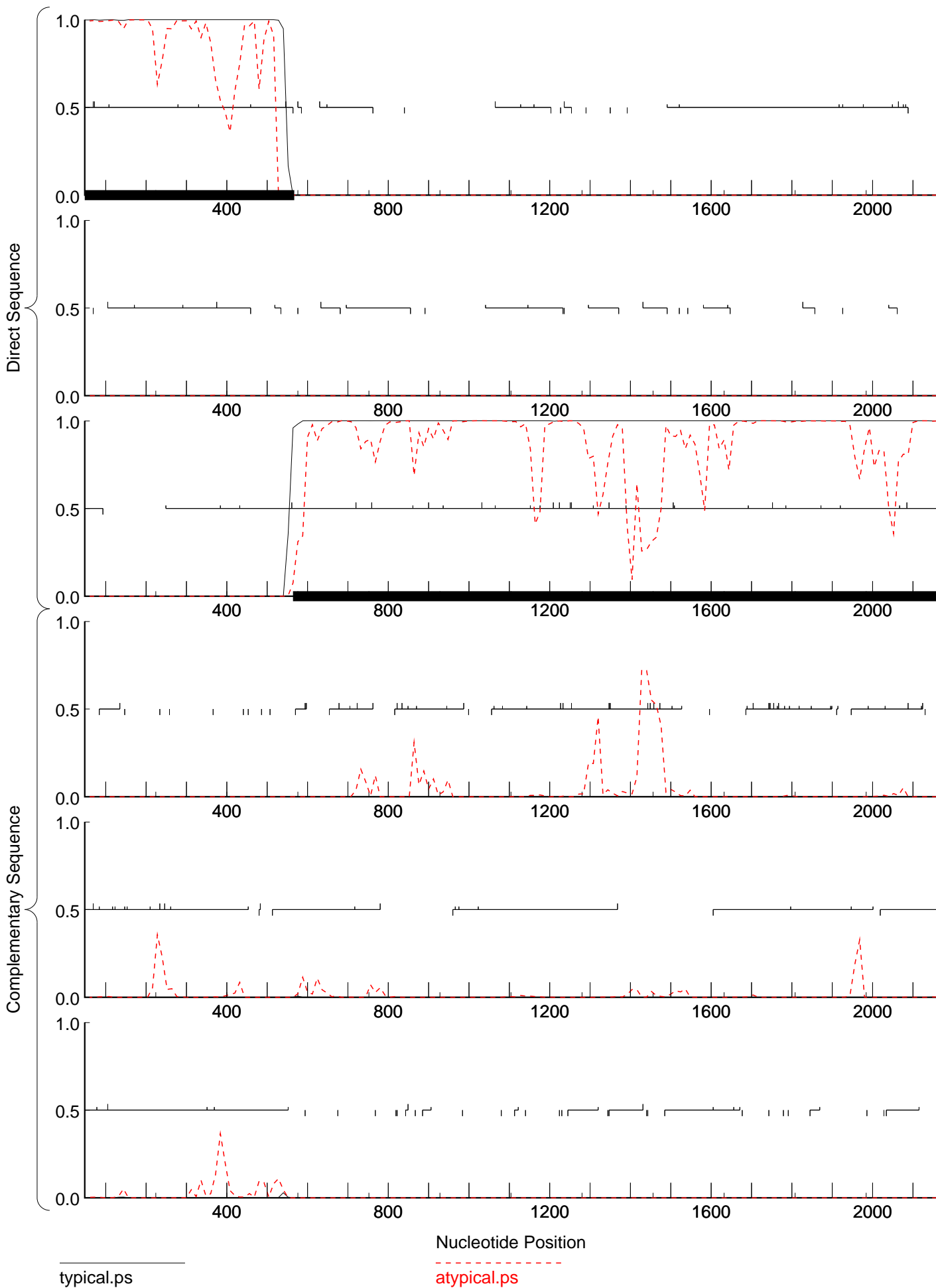
Matrix notes & comments

Training set derived by GeneMarkS, 4.30 October 2014
Fri Aug 18 18:16:12 2017

MATRIX INFORMATION

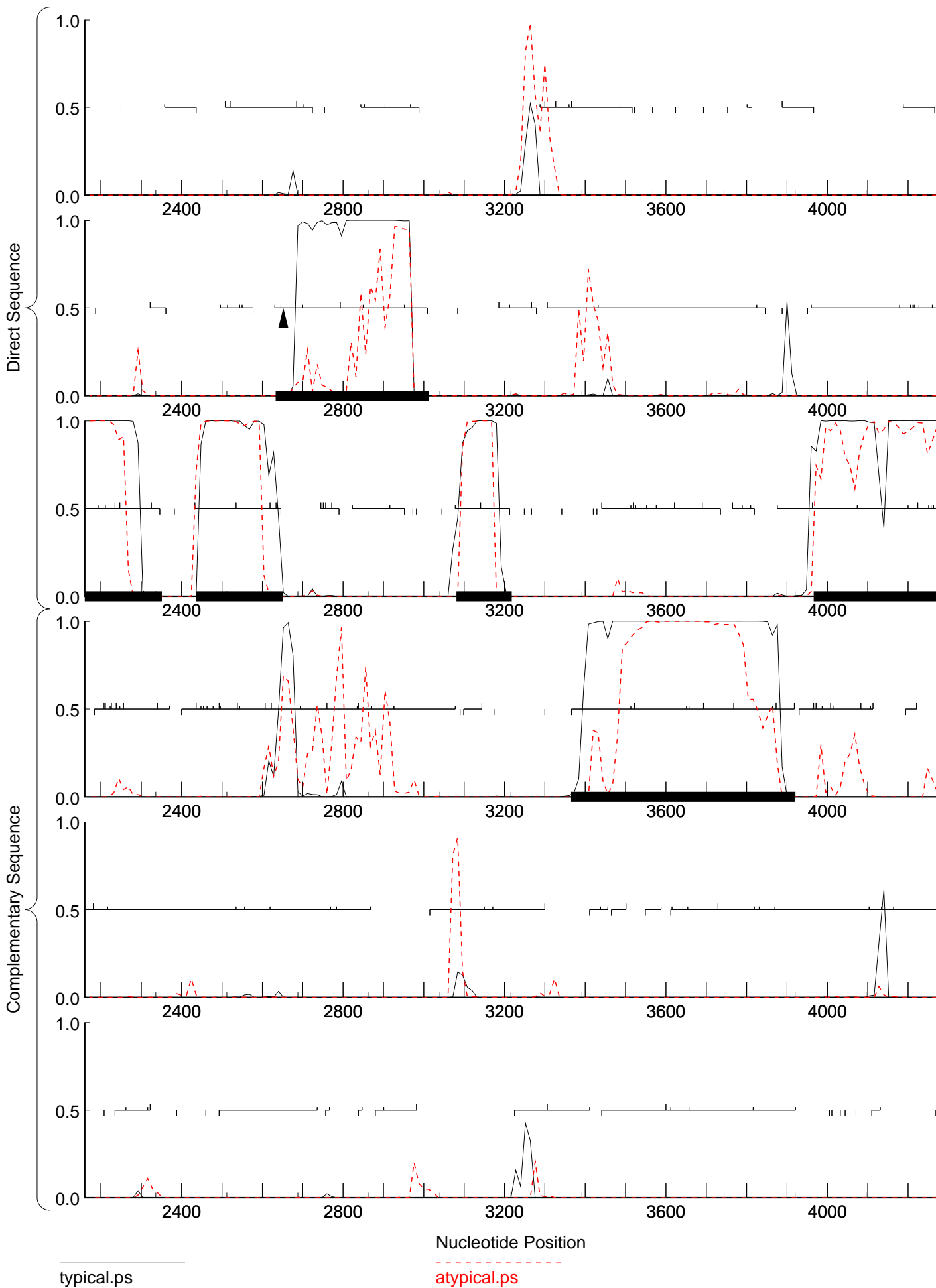
Matrix : -
Author : -
Order : 4

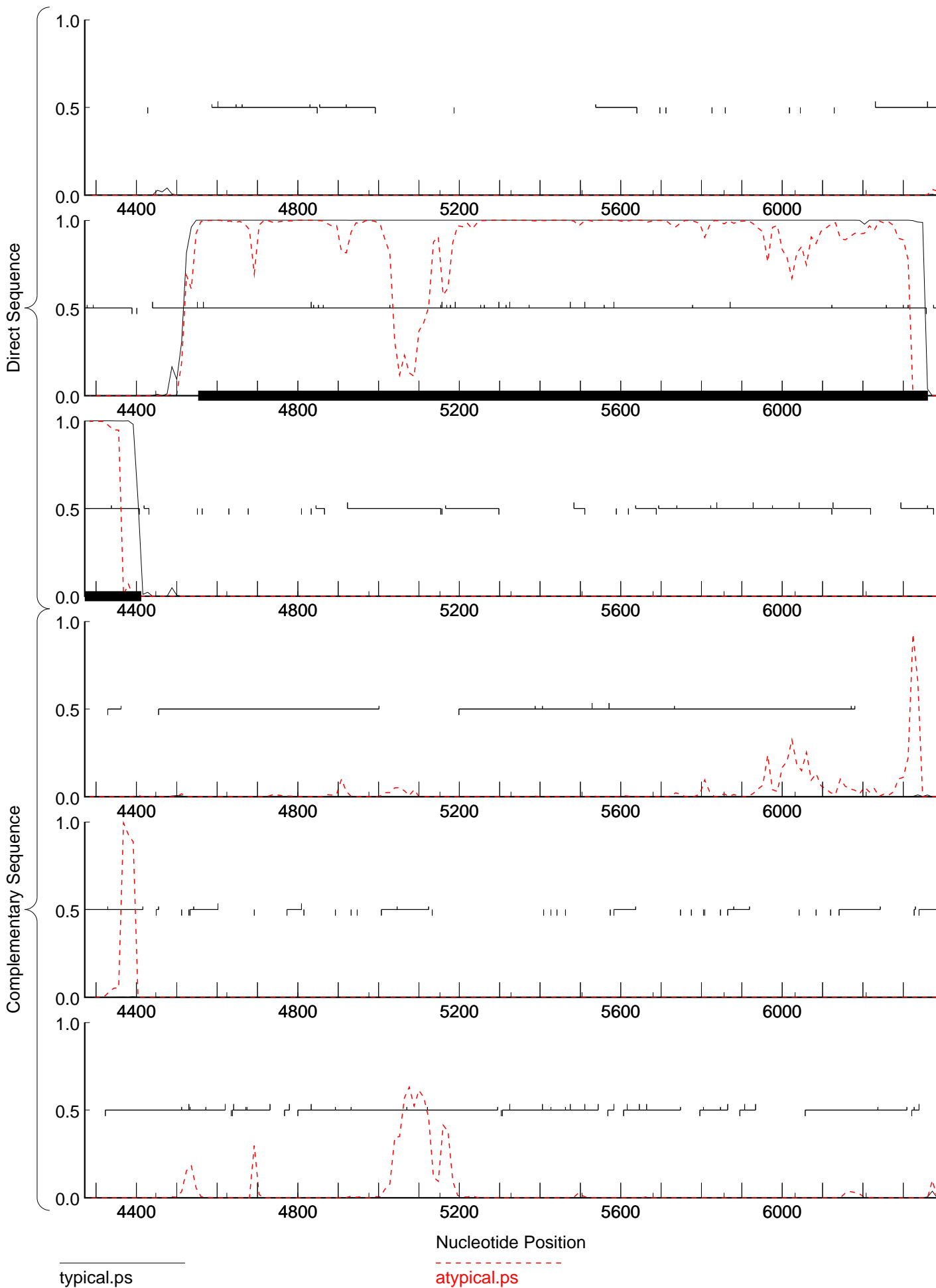
Send questions / comments to:
Dr. M. Borodovsky
Georgia Institute of Technology
School of Biology
Atlanta, GA 30332-0230

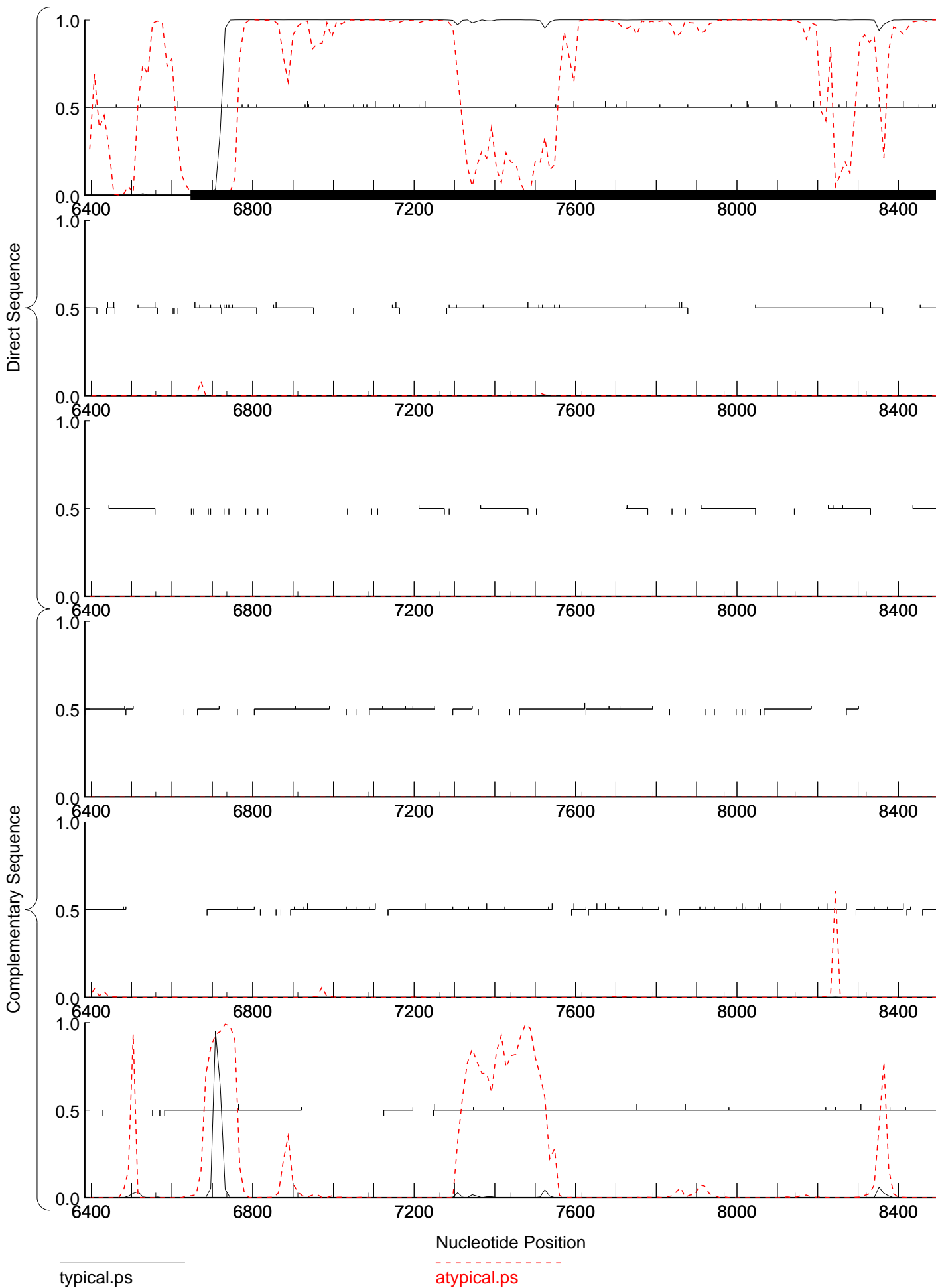


typical.ps

atypical.ps

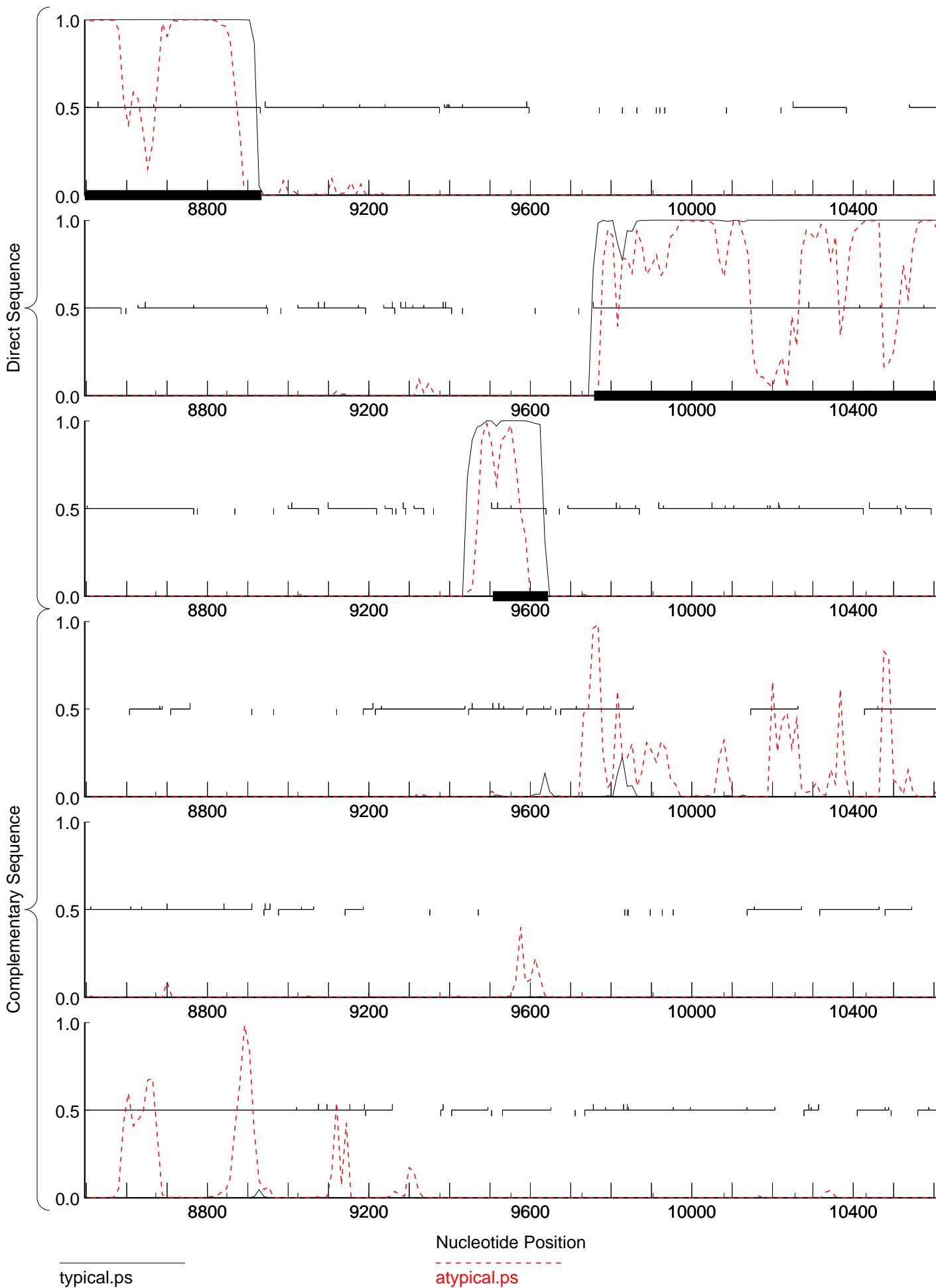


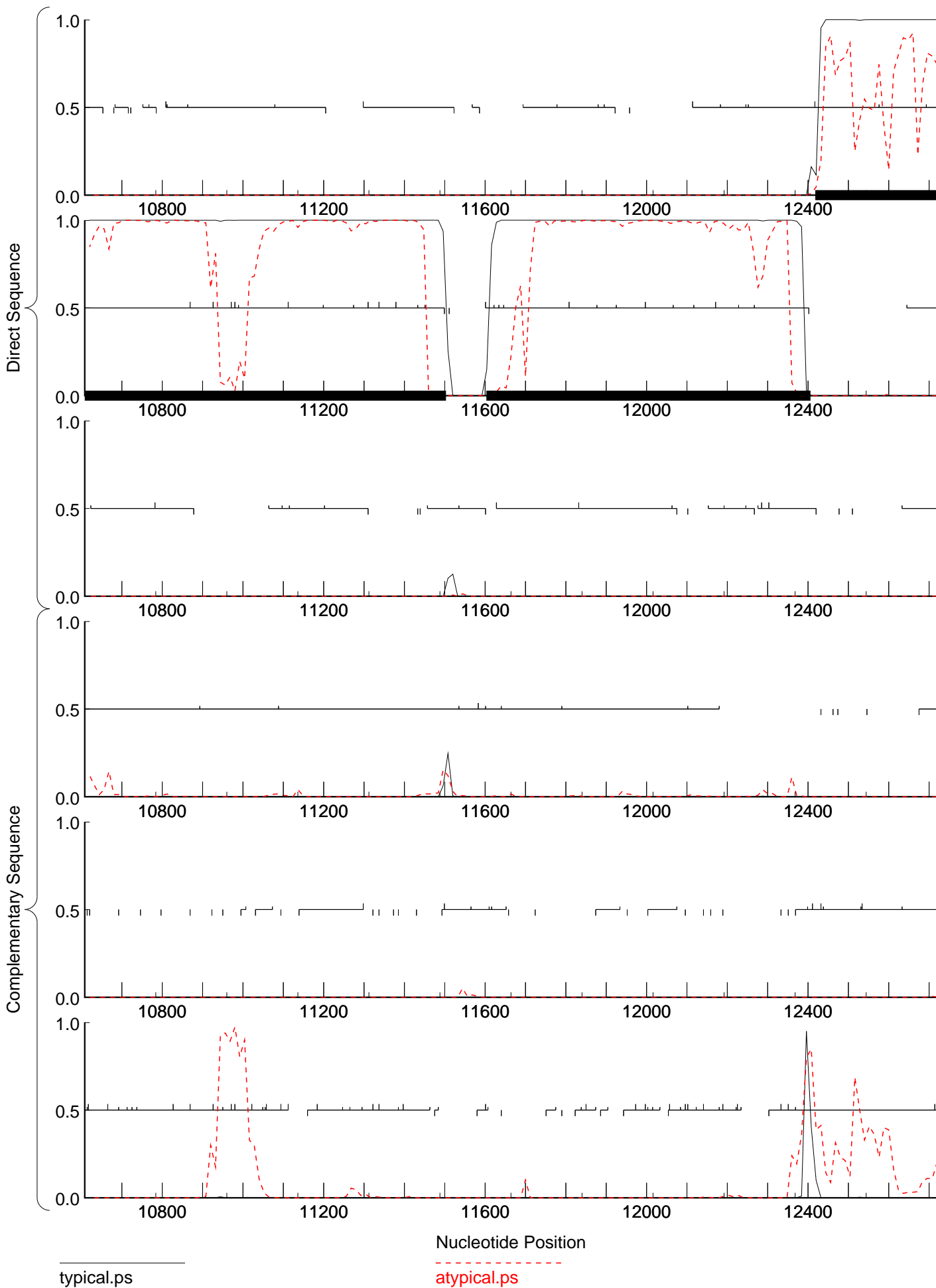


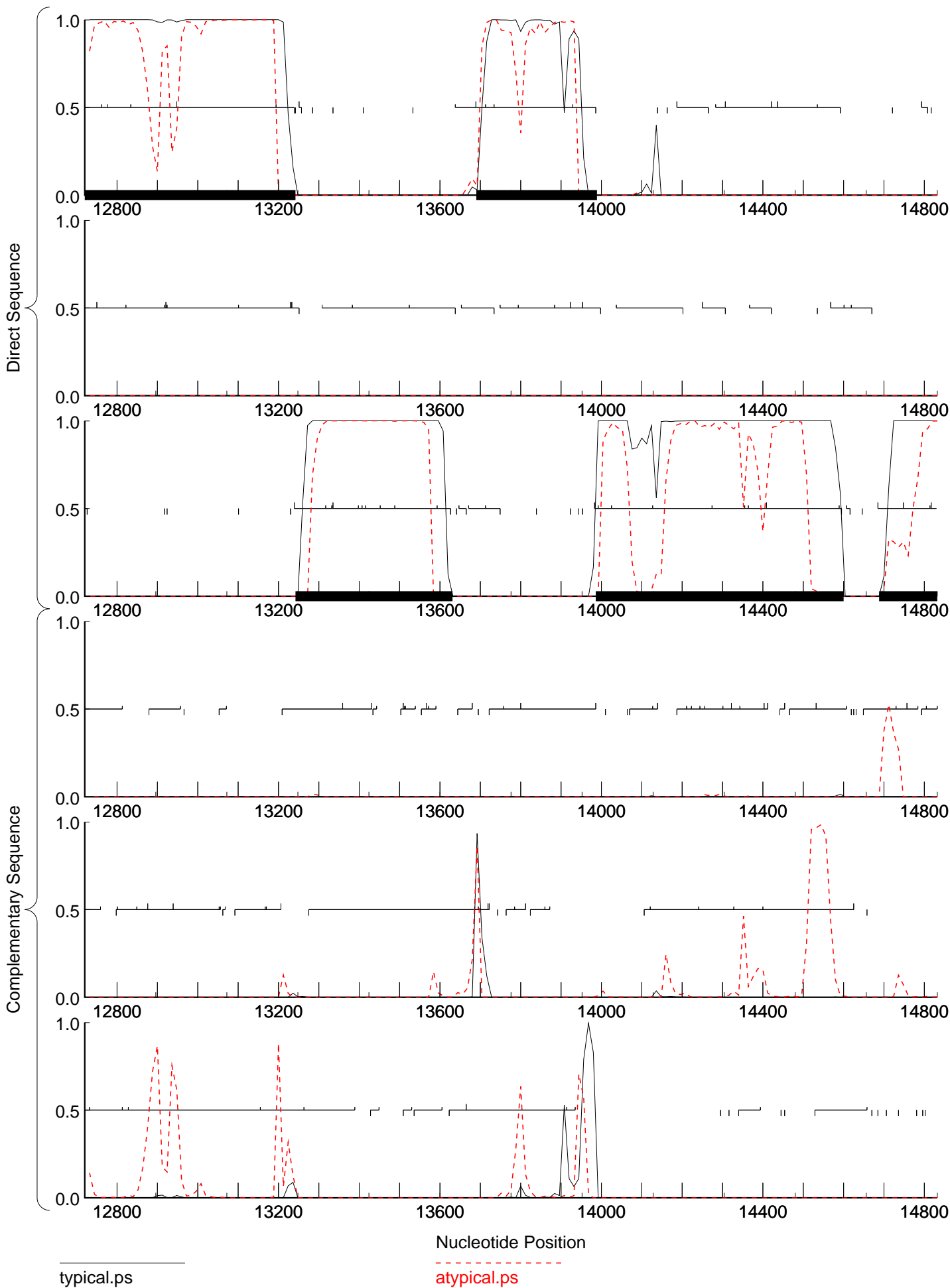


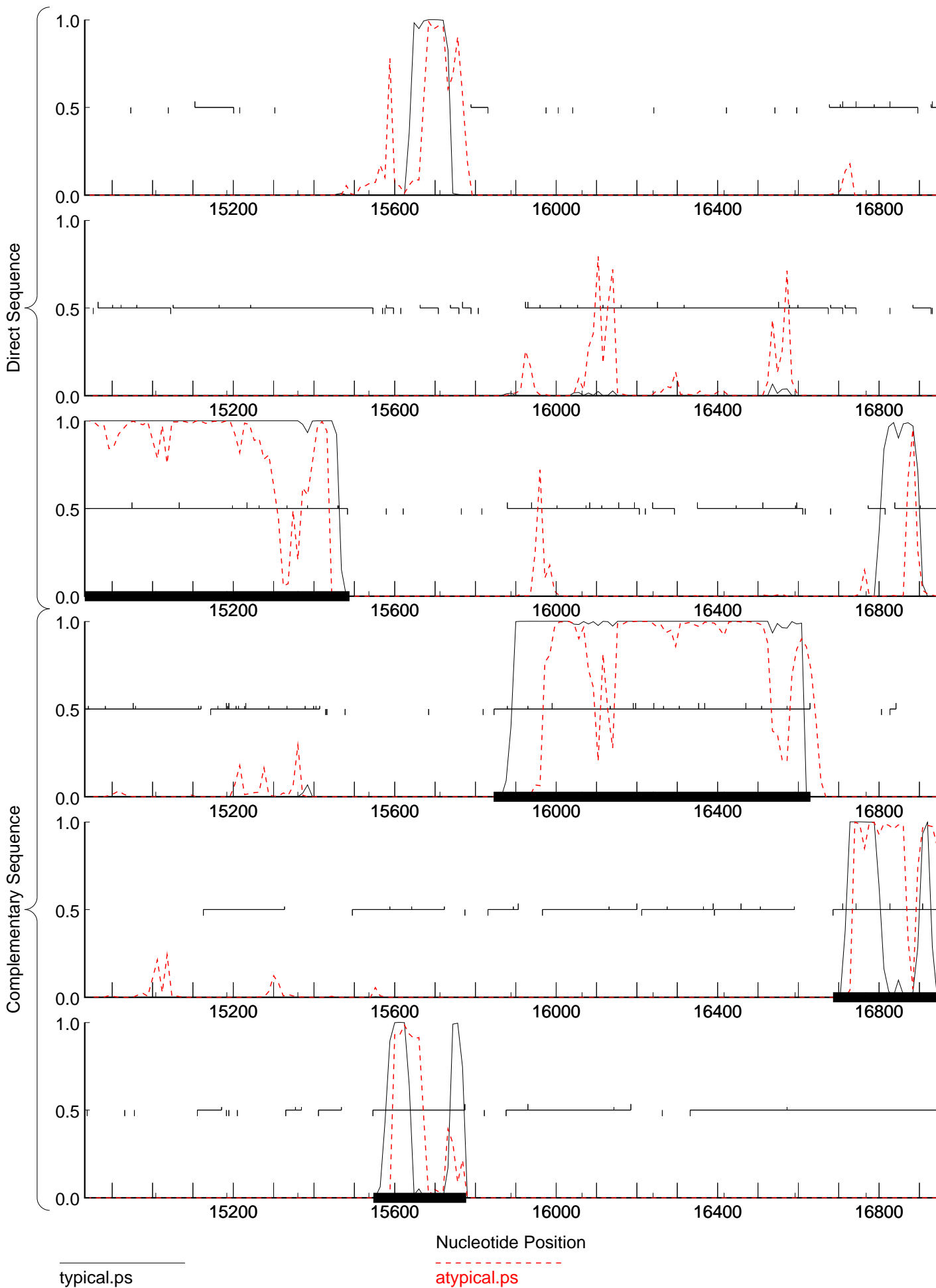
typical.ps

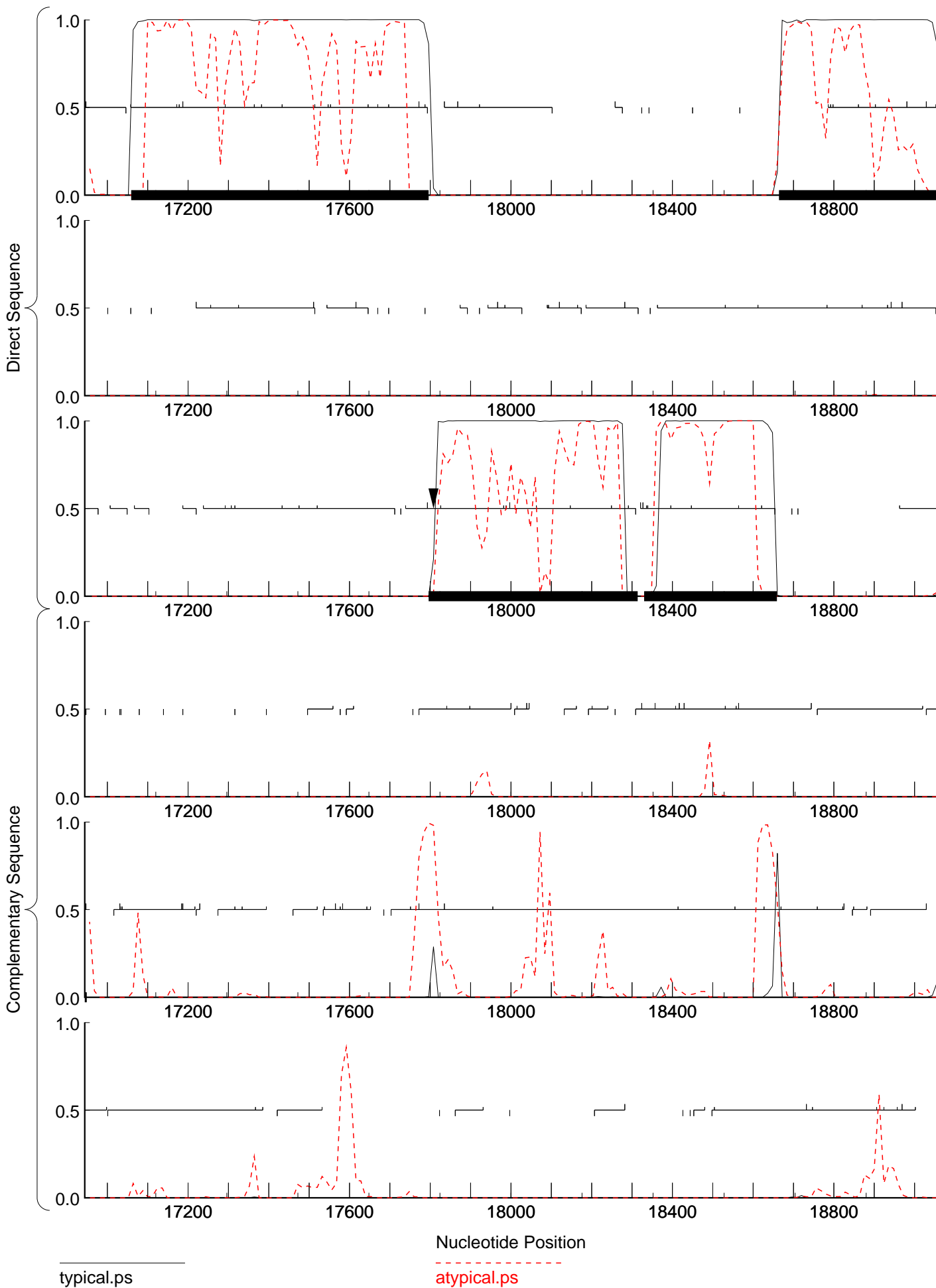
atypical.ps

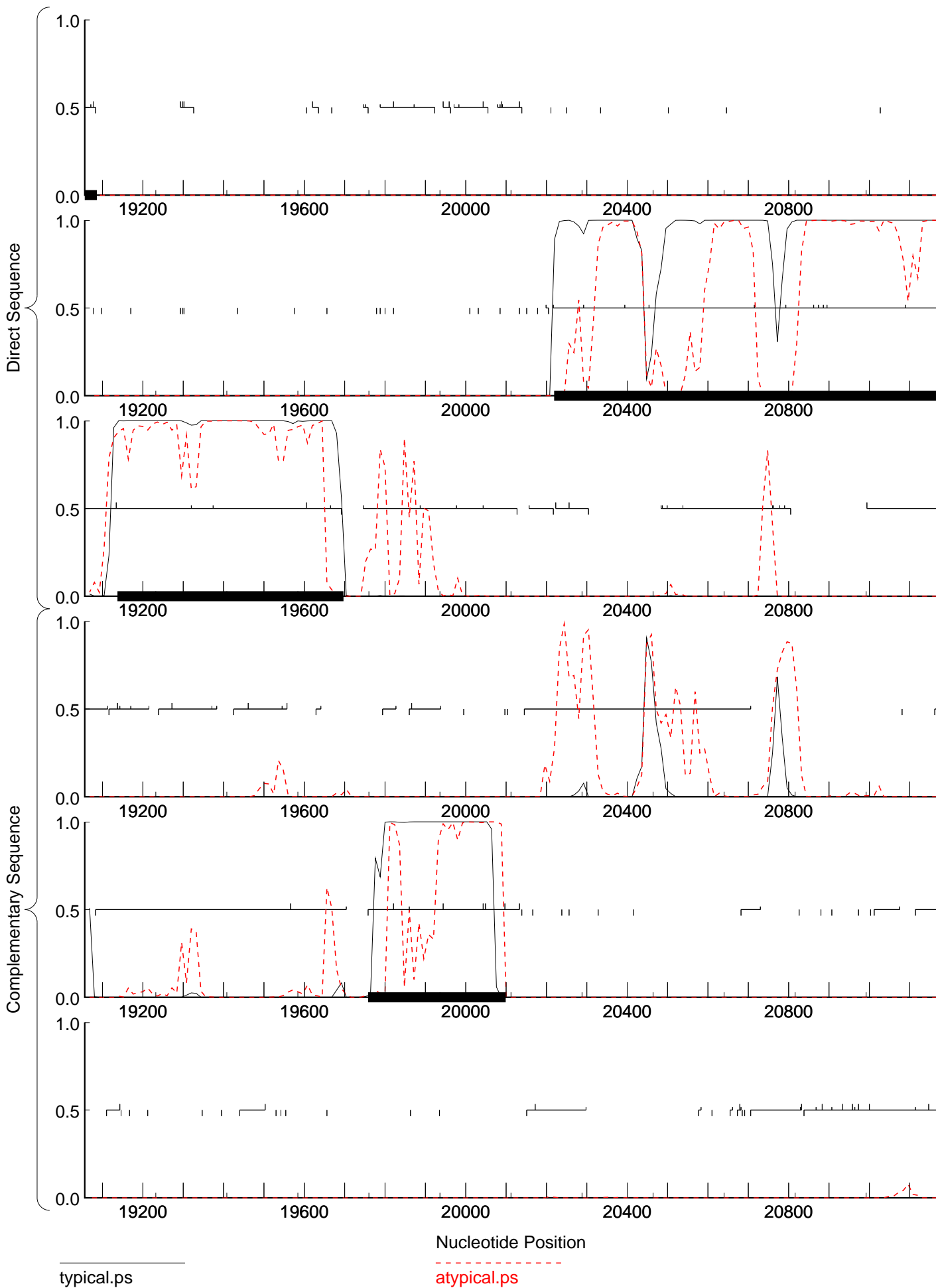






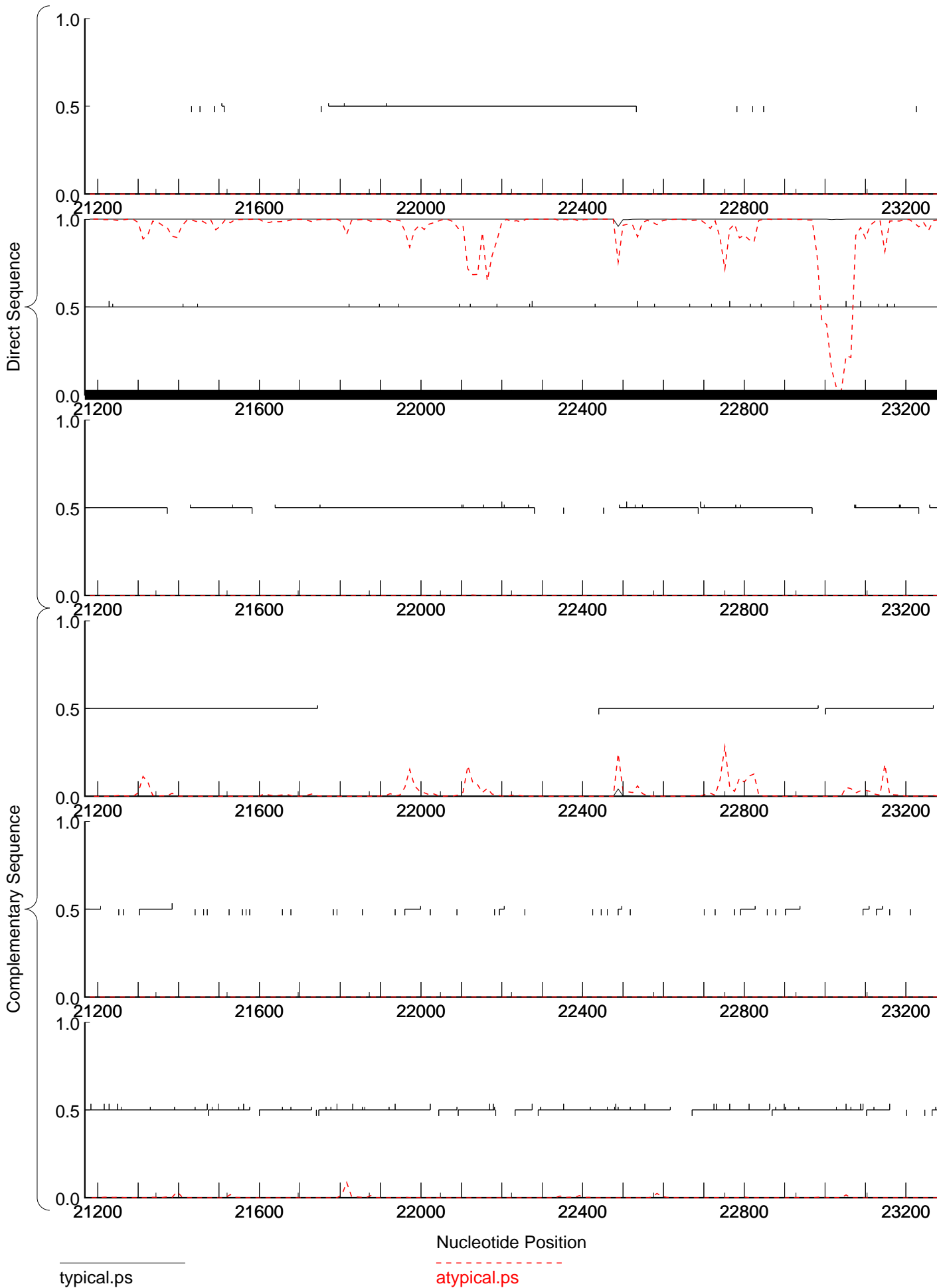






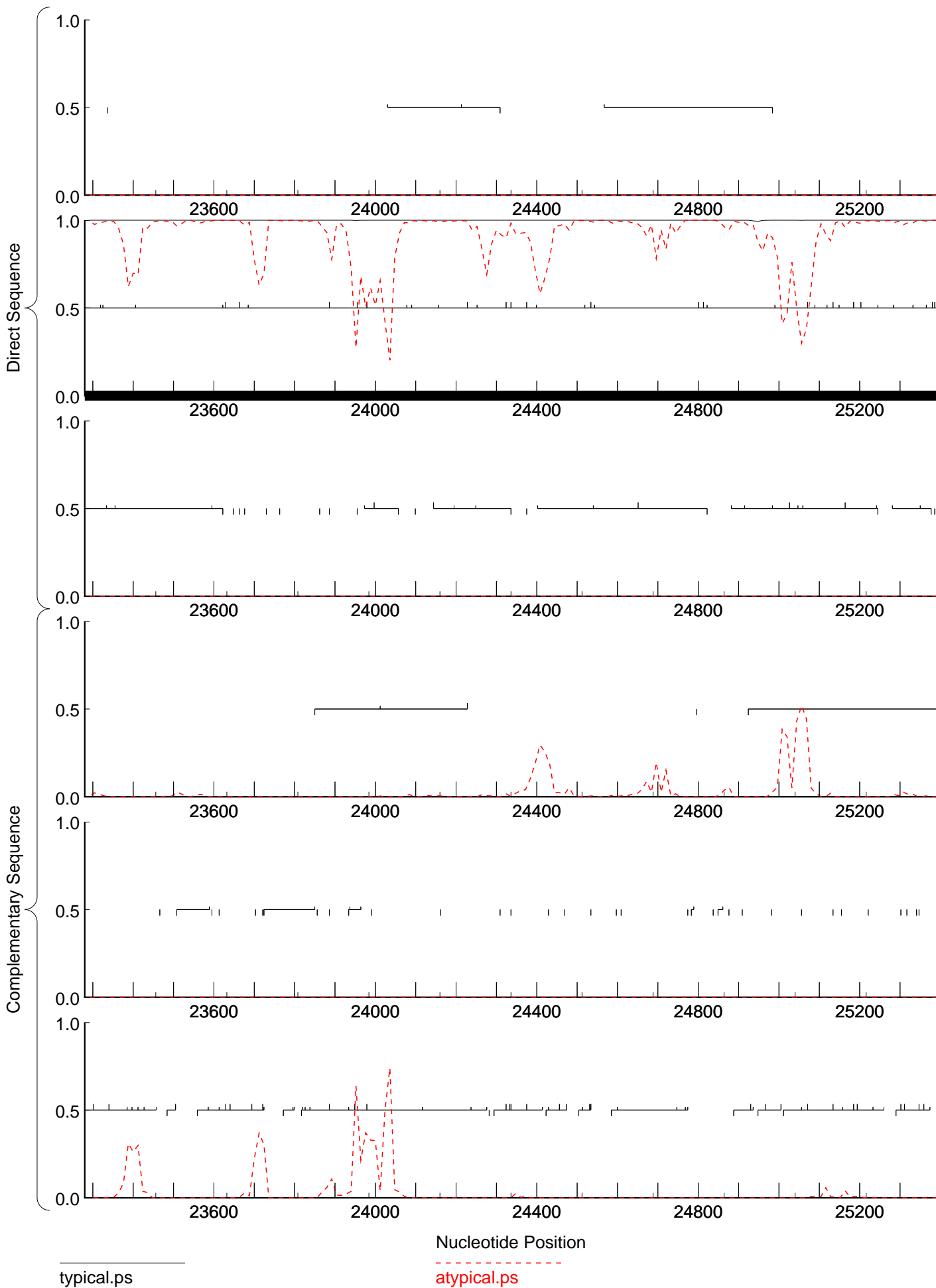
typical.ps

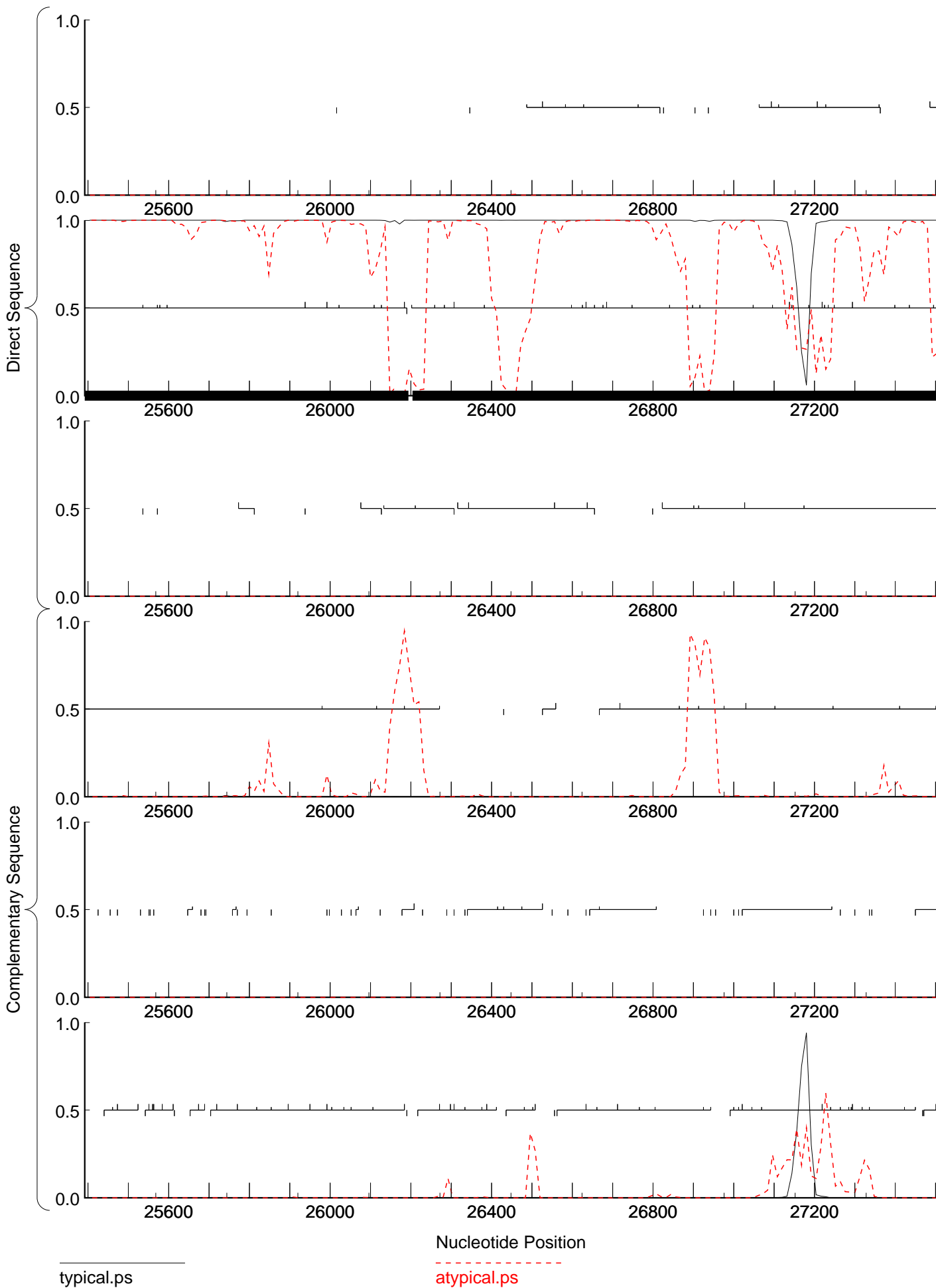
atypical.ps

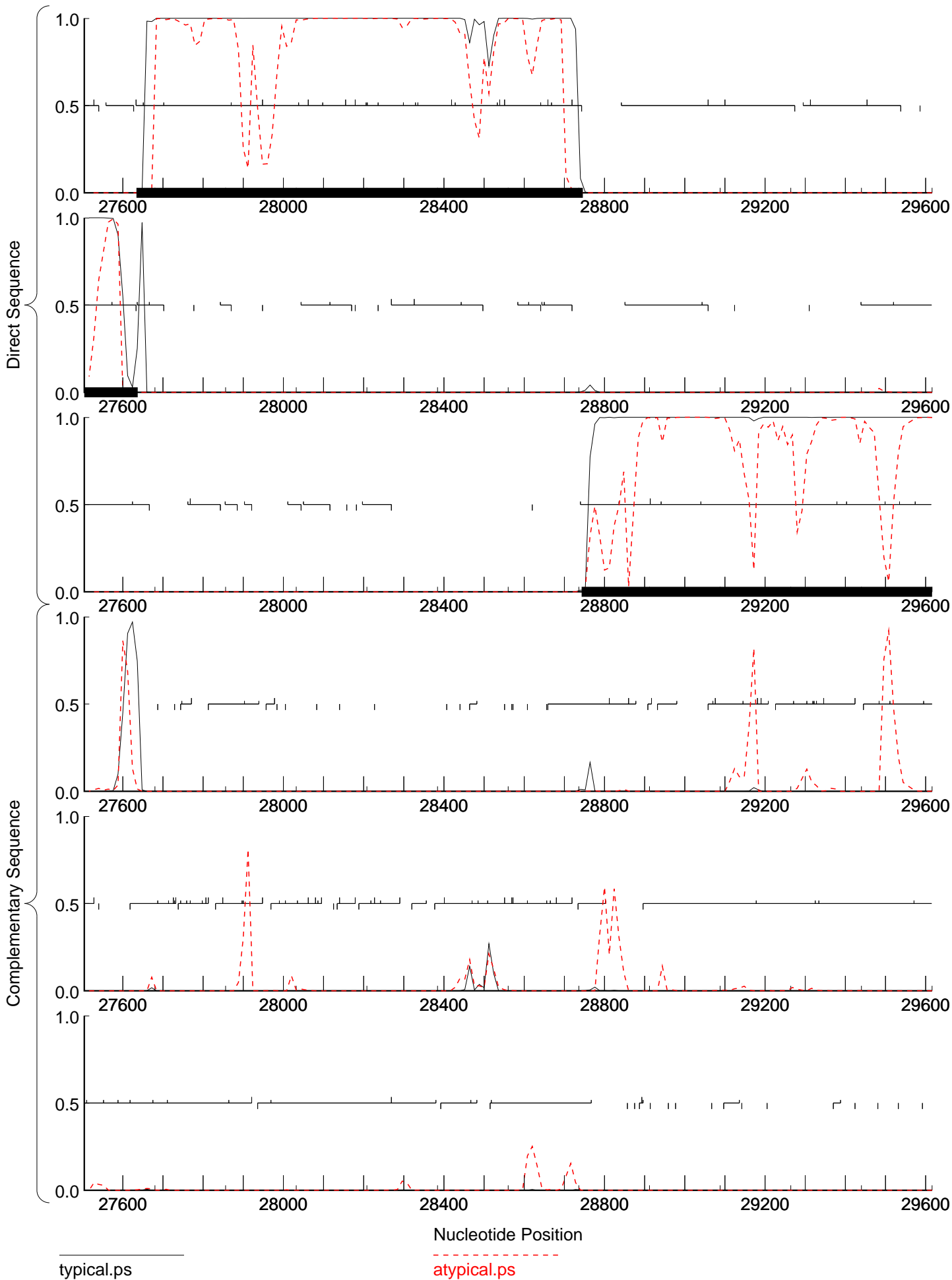


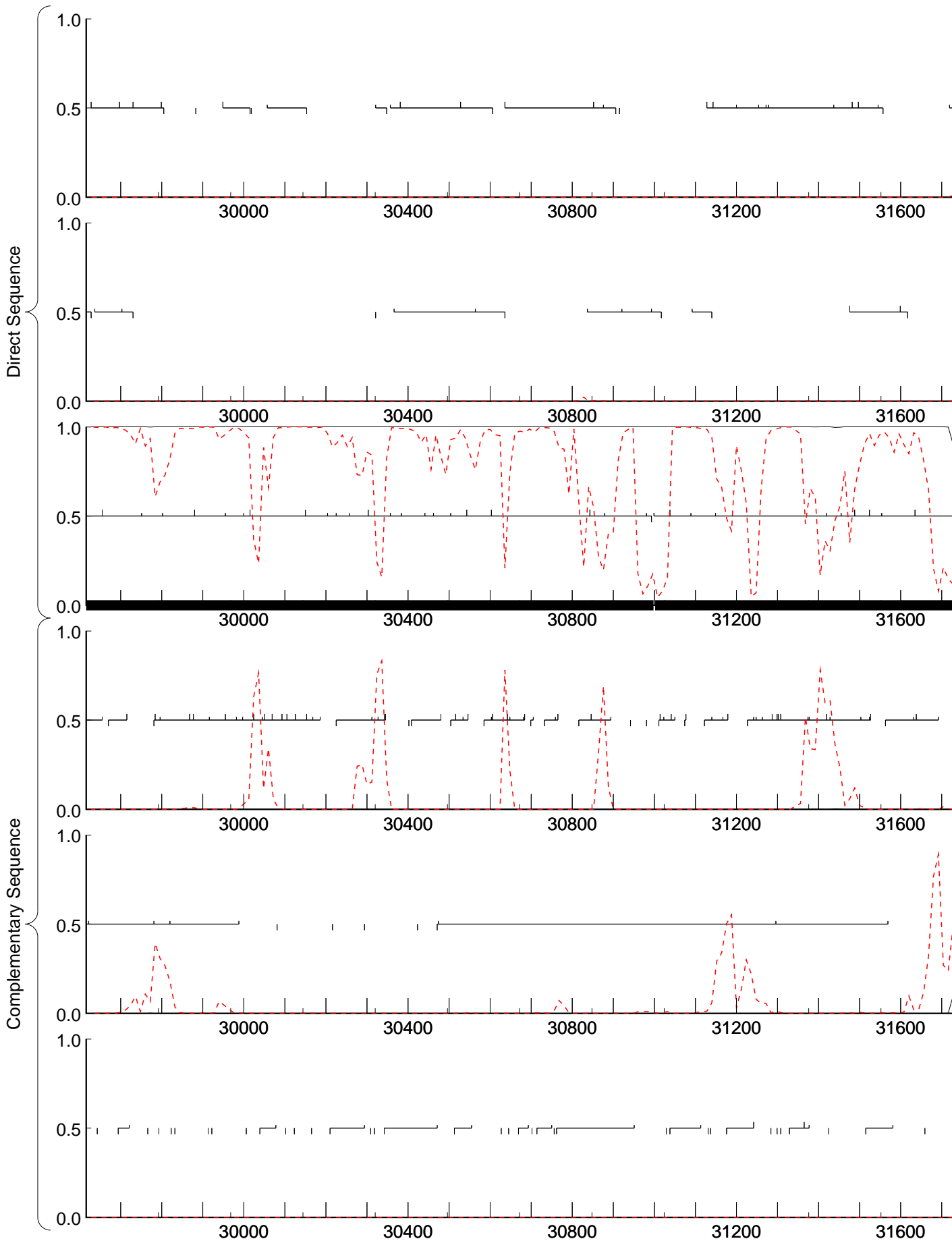
typical.ps

atypical.ps



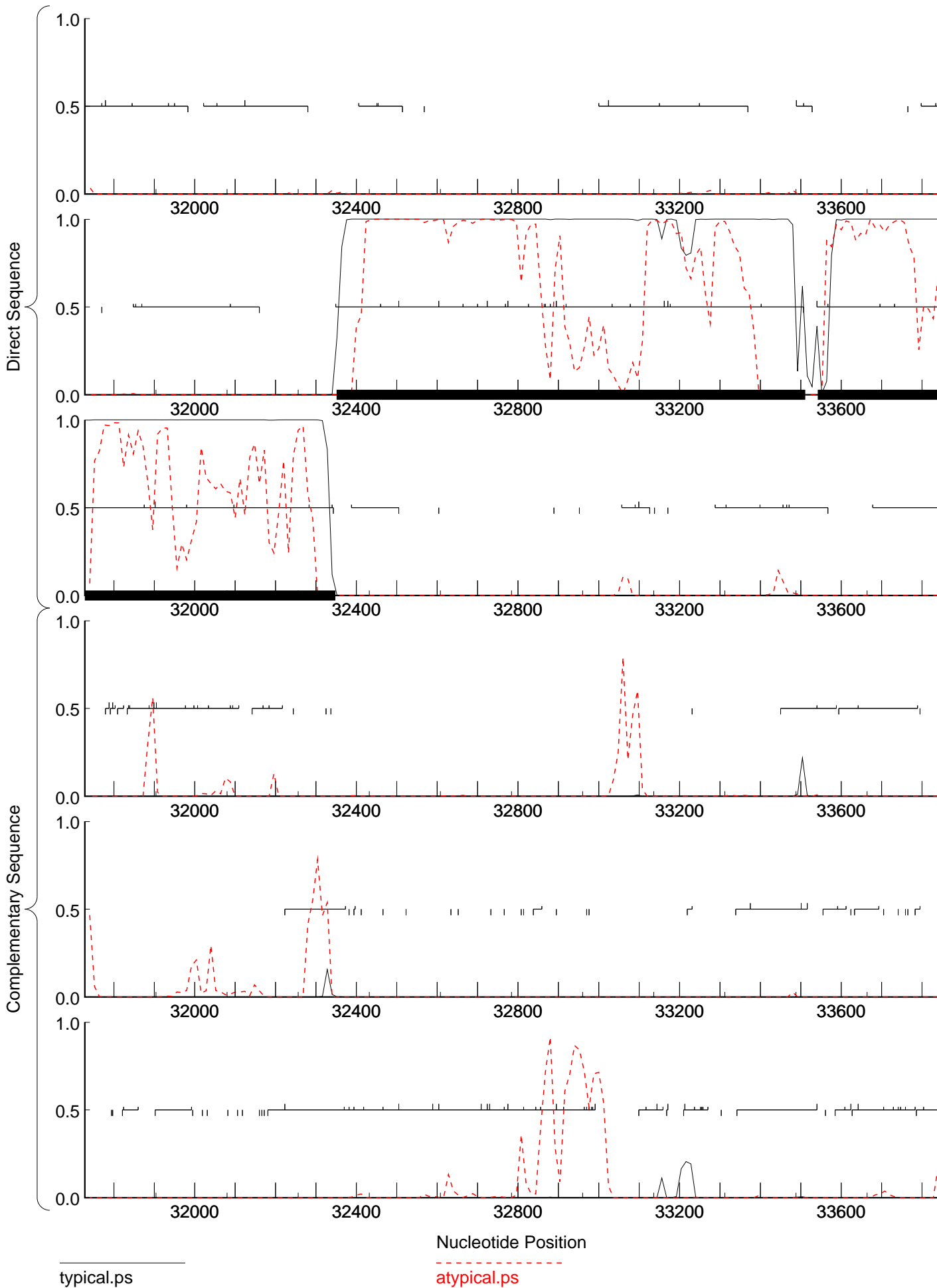






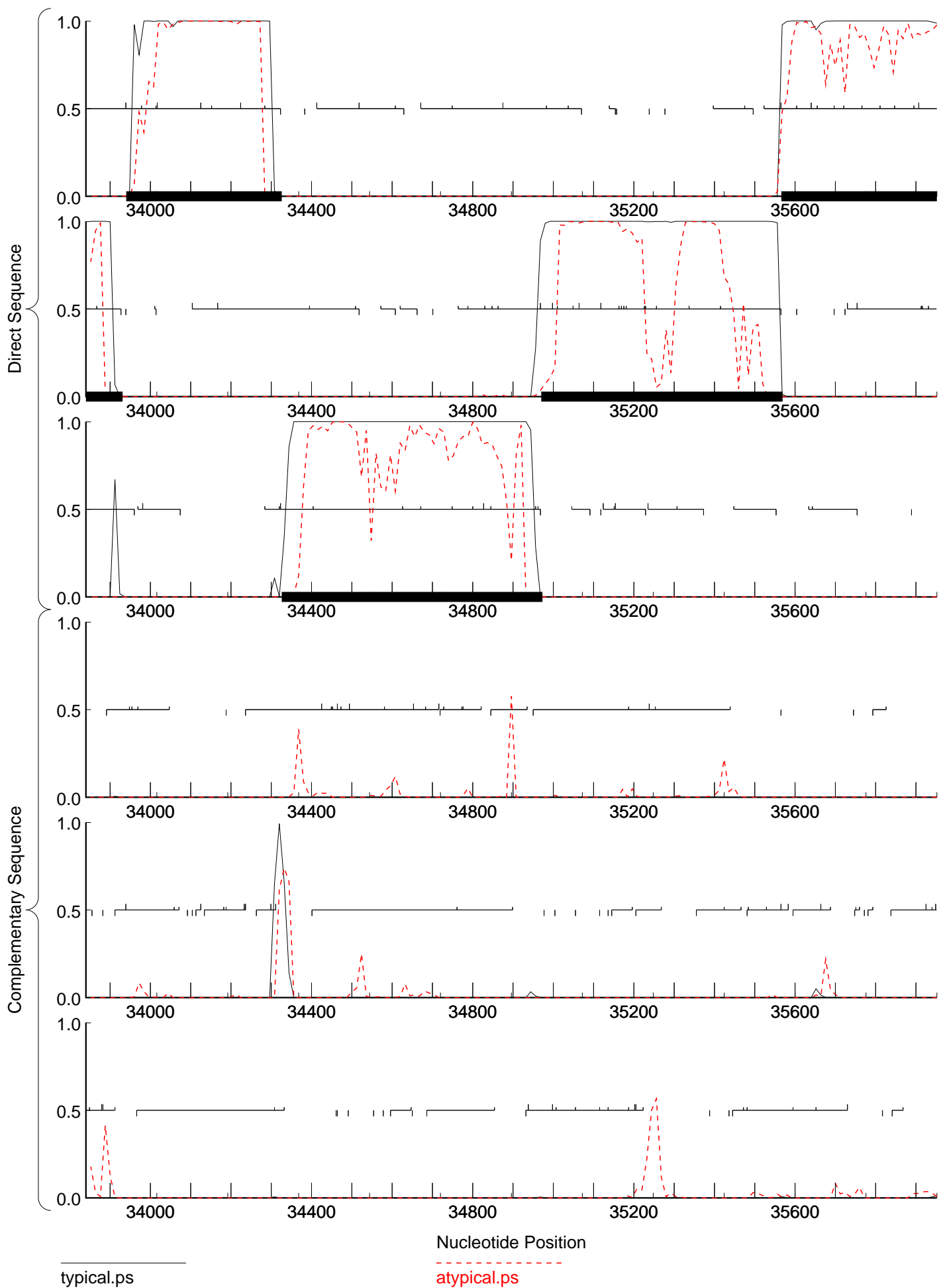
typical.ps

atypical.ps



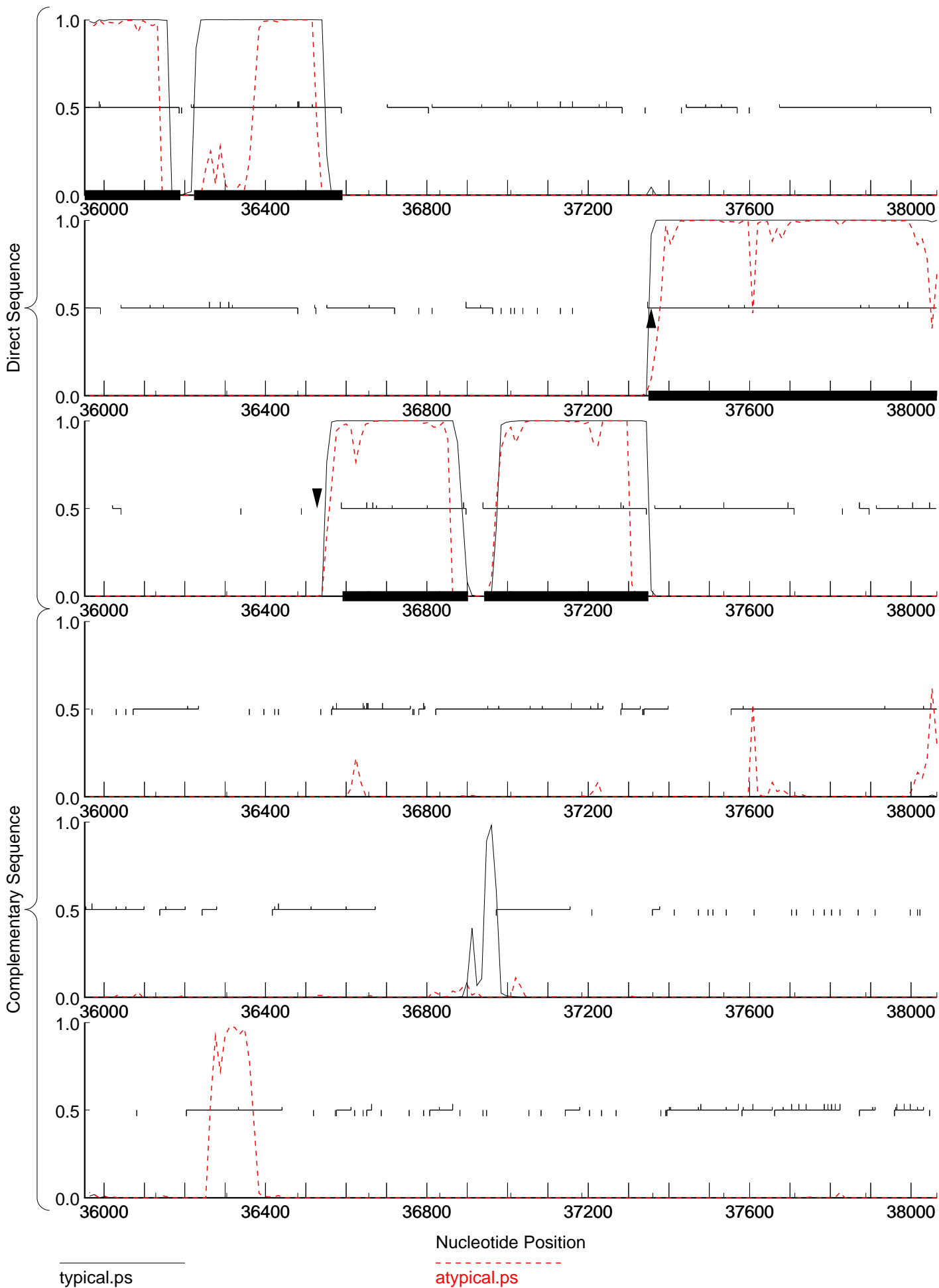
typical.ps

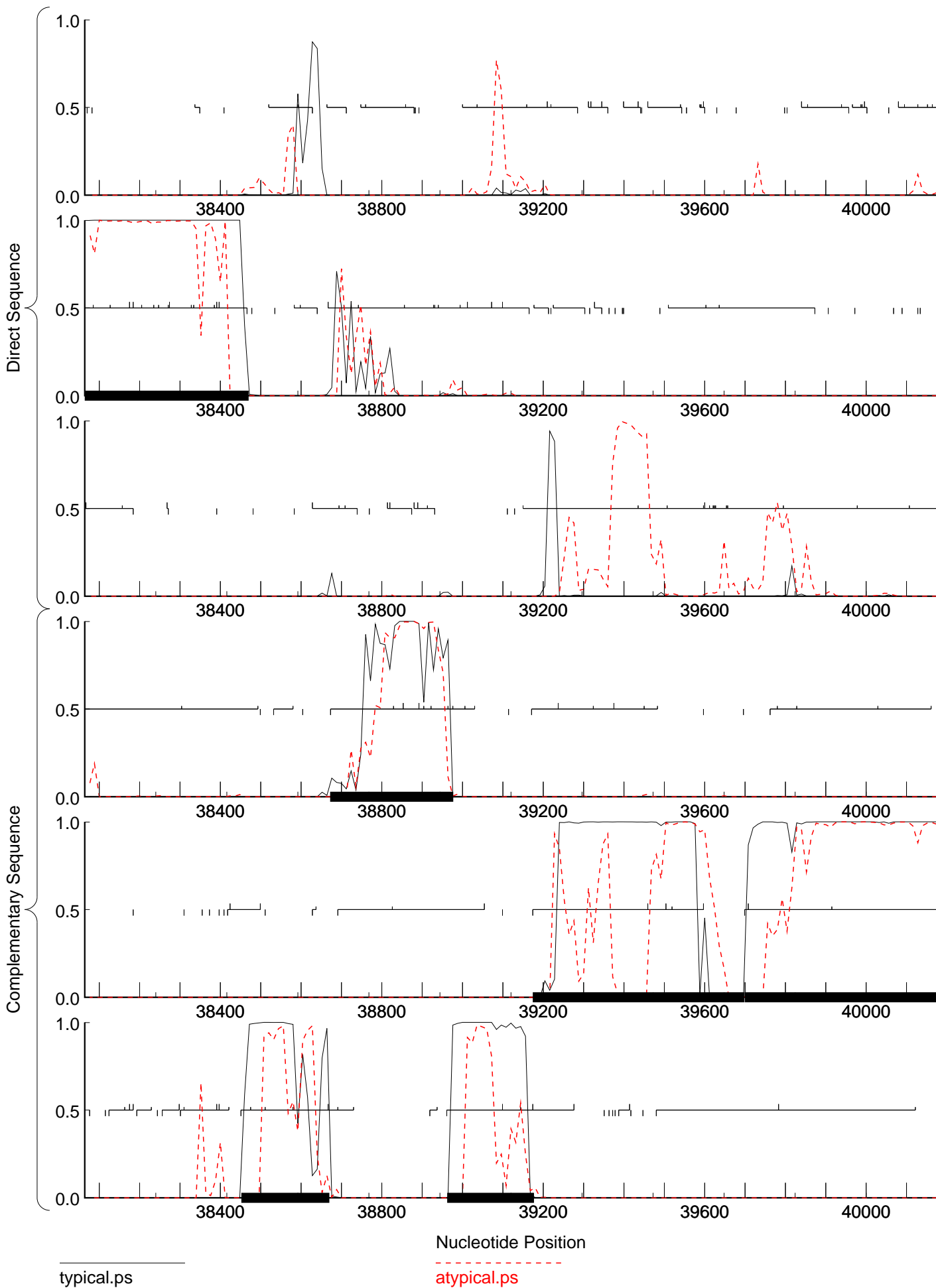
atypical.ps



typical.ps

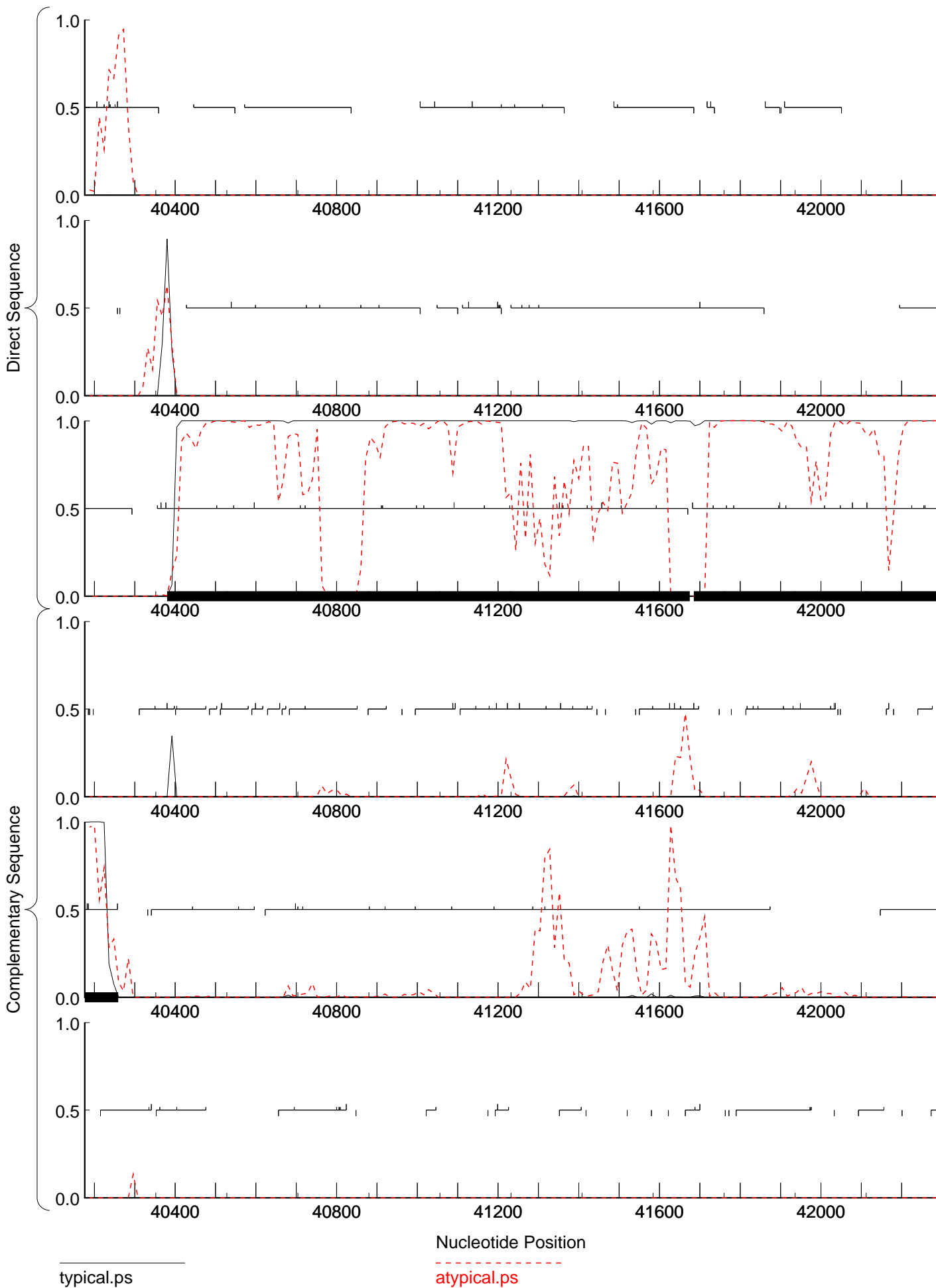
atypical.ps

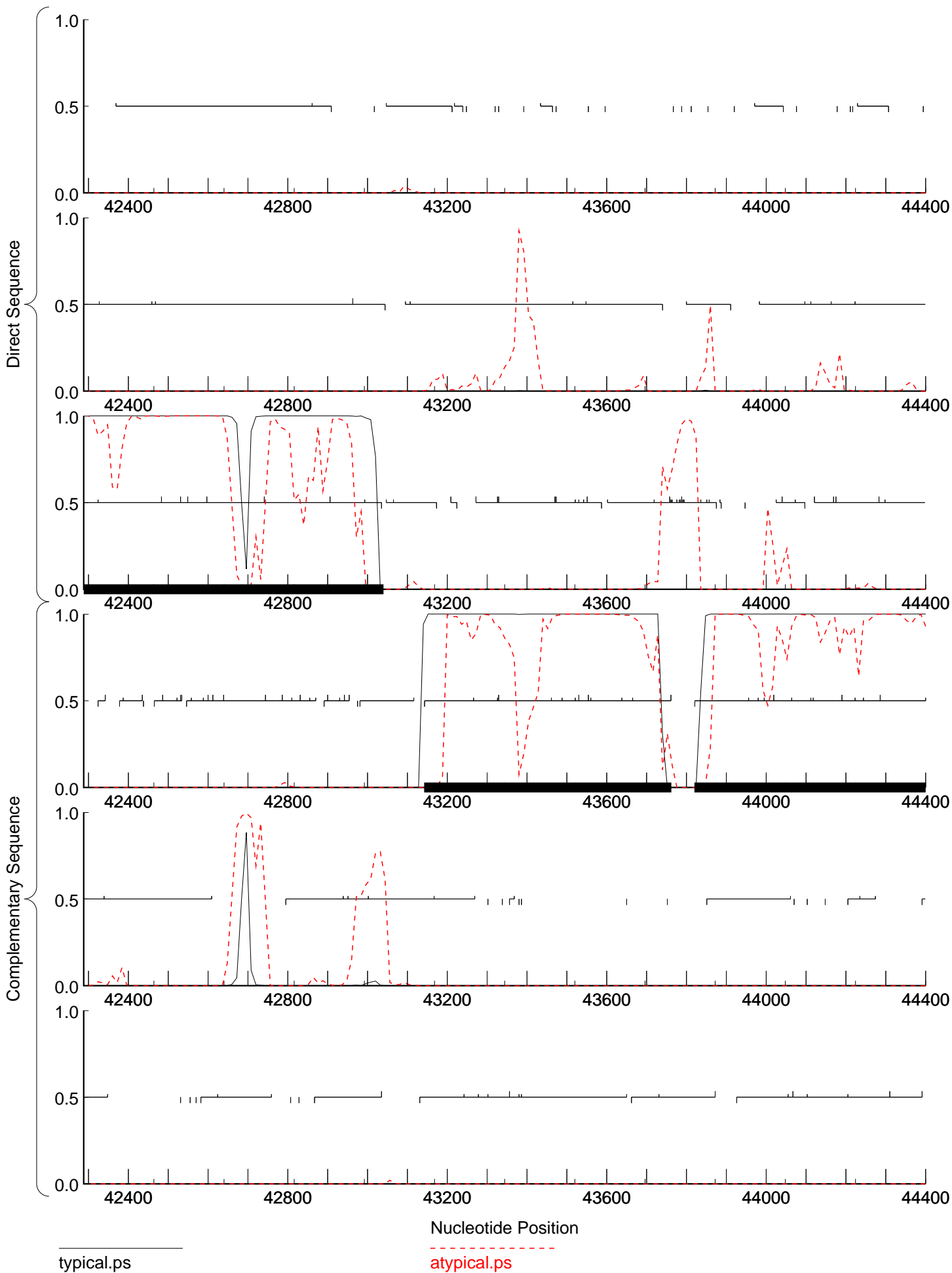




typical.ps

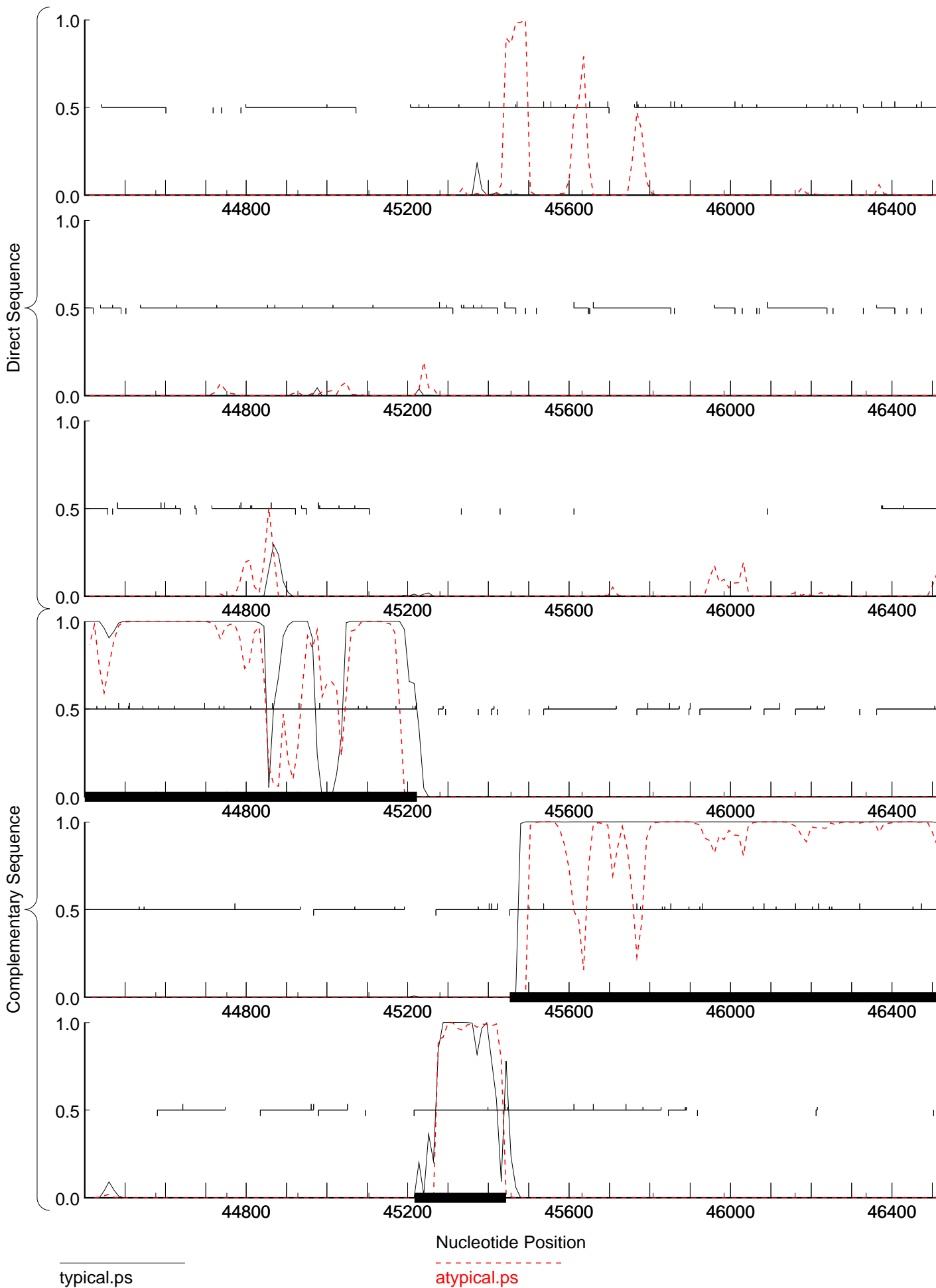
atypical.ps

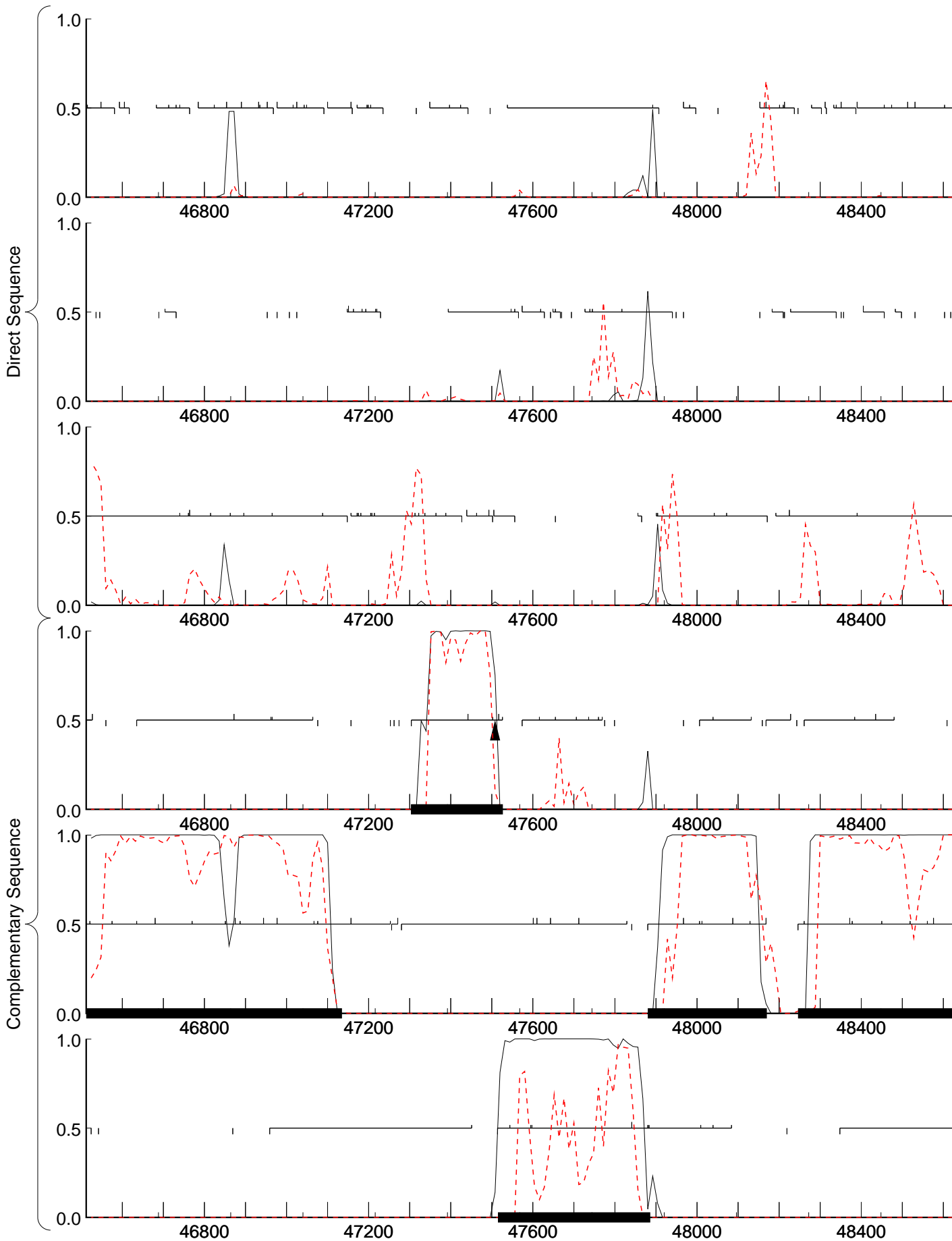




typical.ps

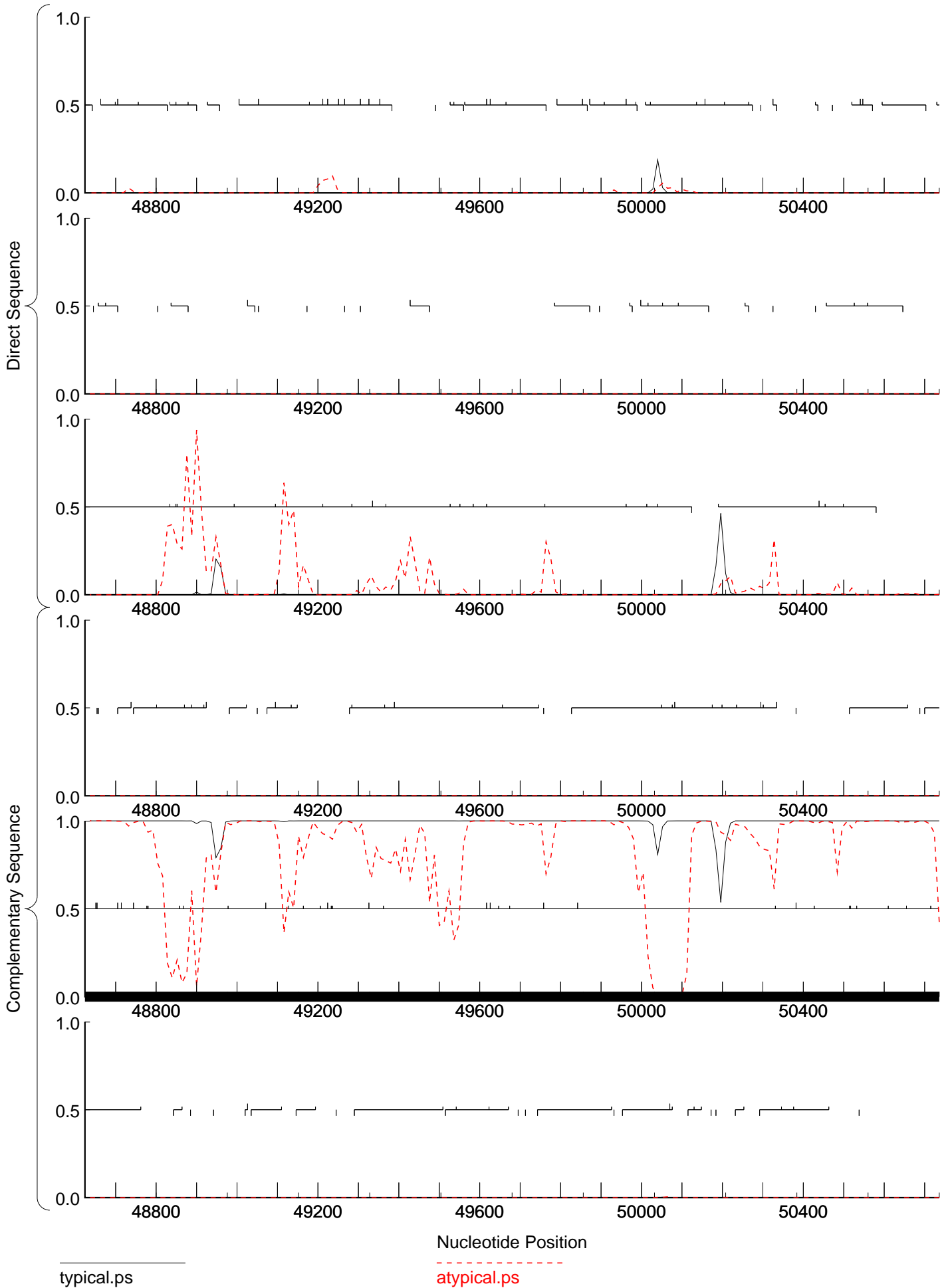
atypical.ps

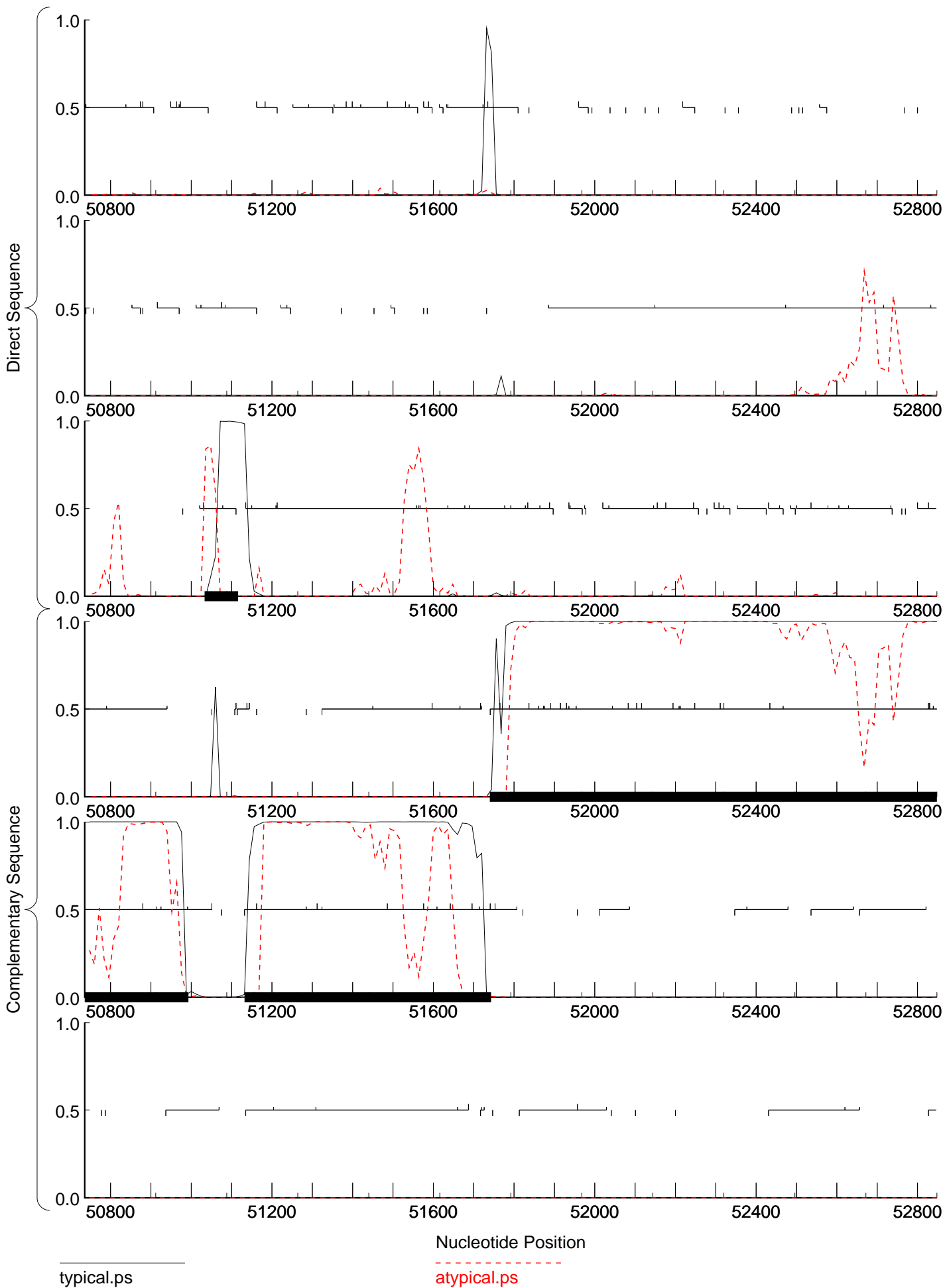




typical.ps

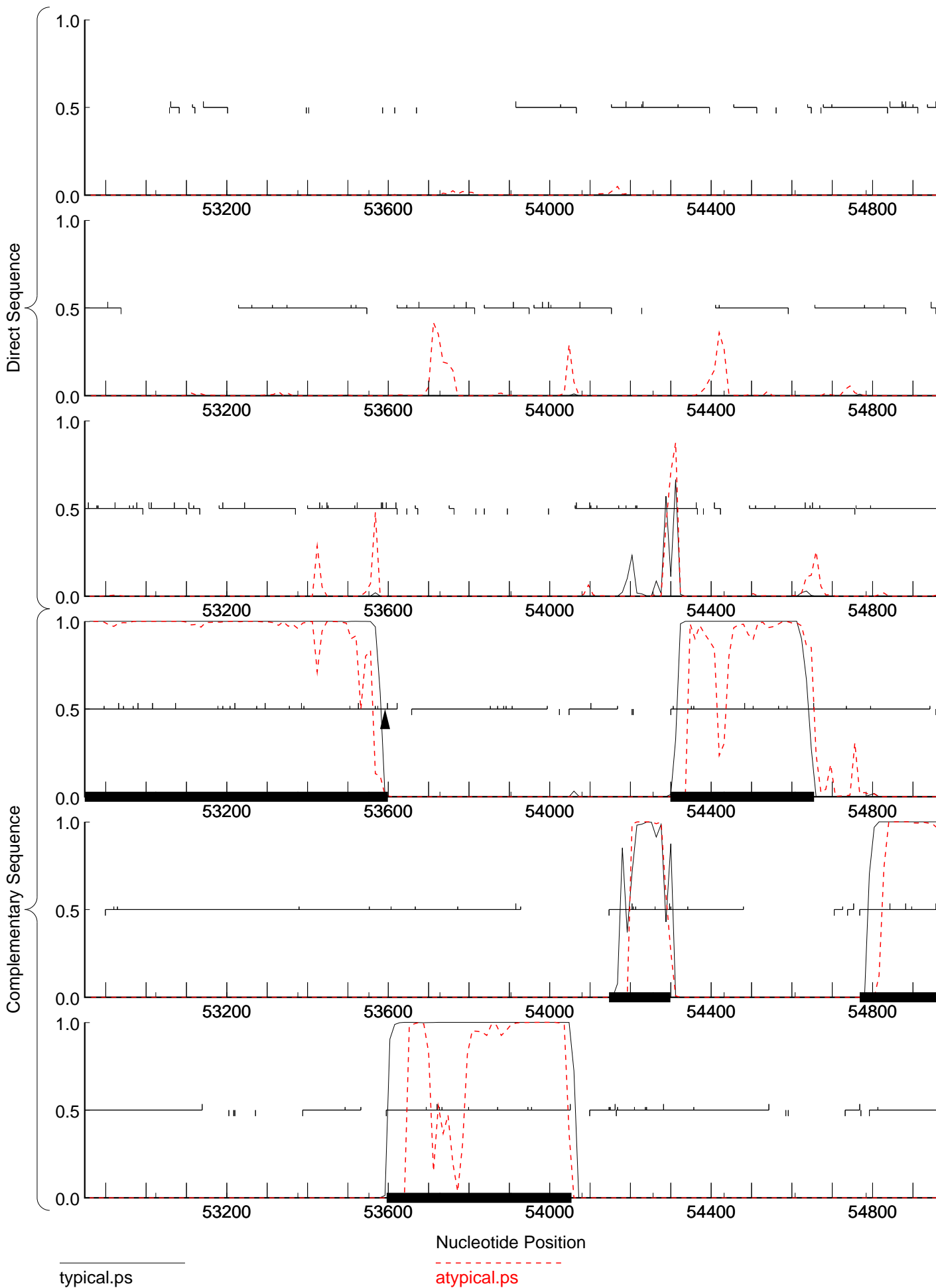
atypical.ps





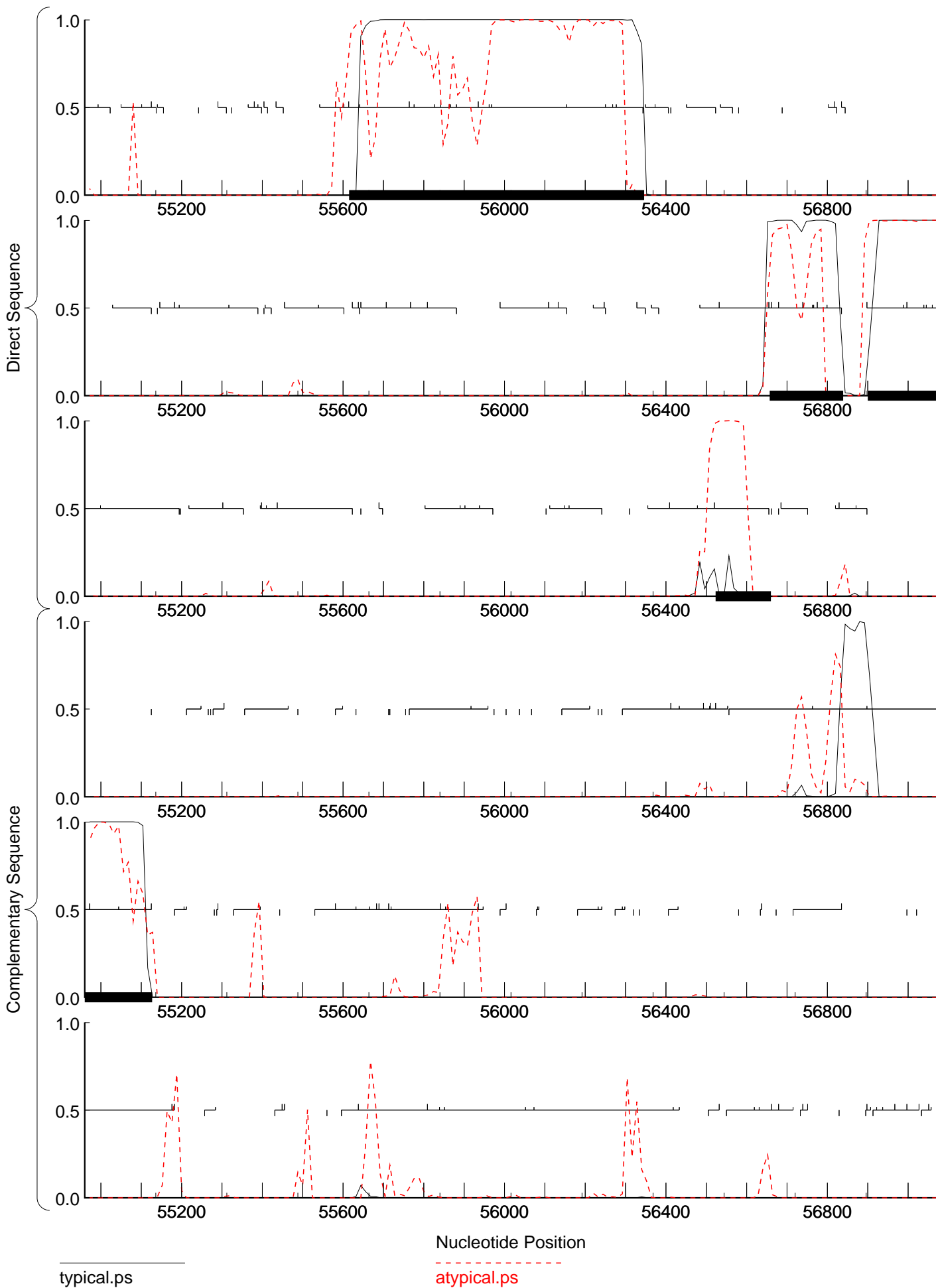
typical.ps

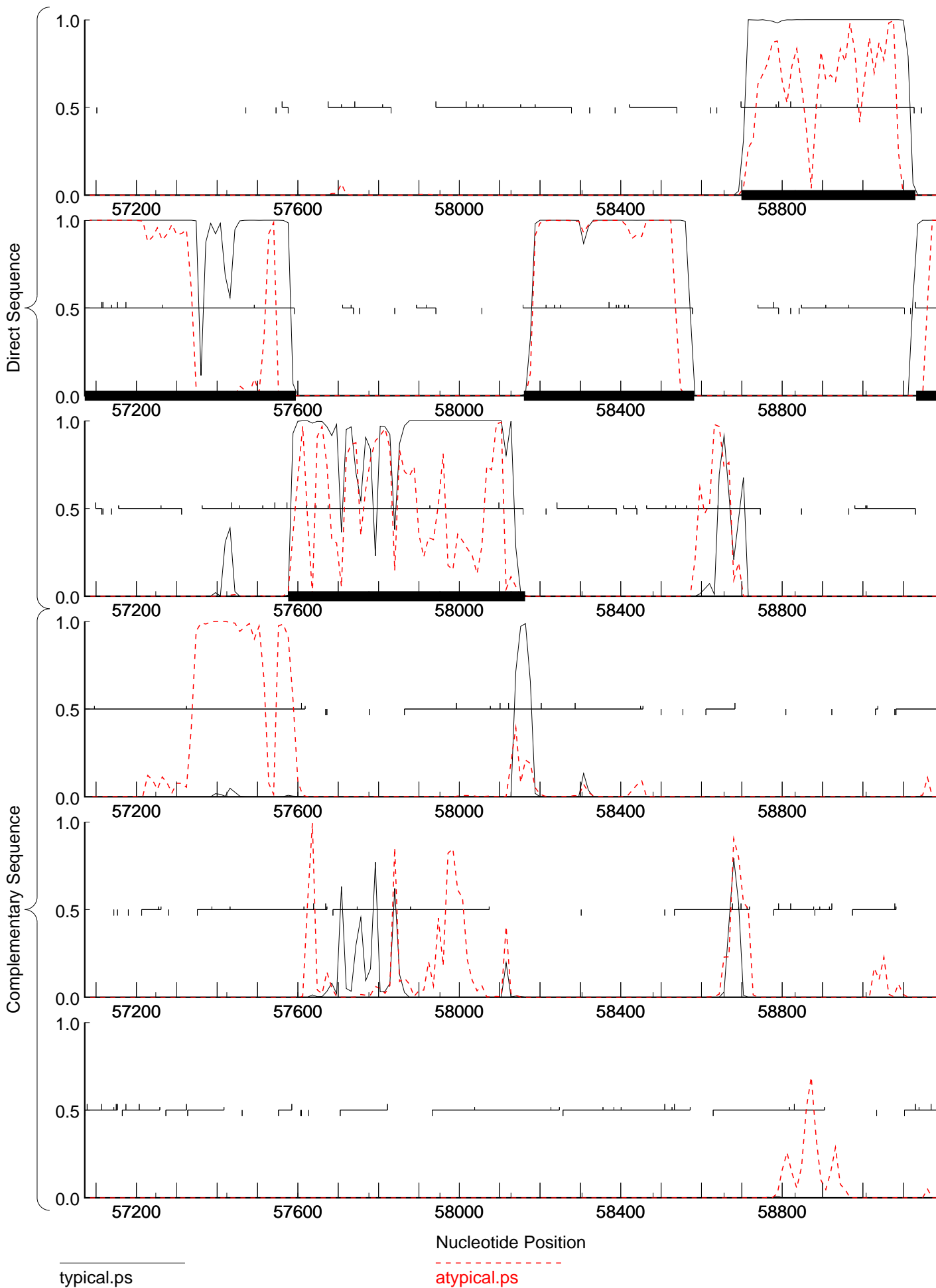
atypical.ps



typical.ps

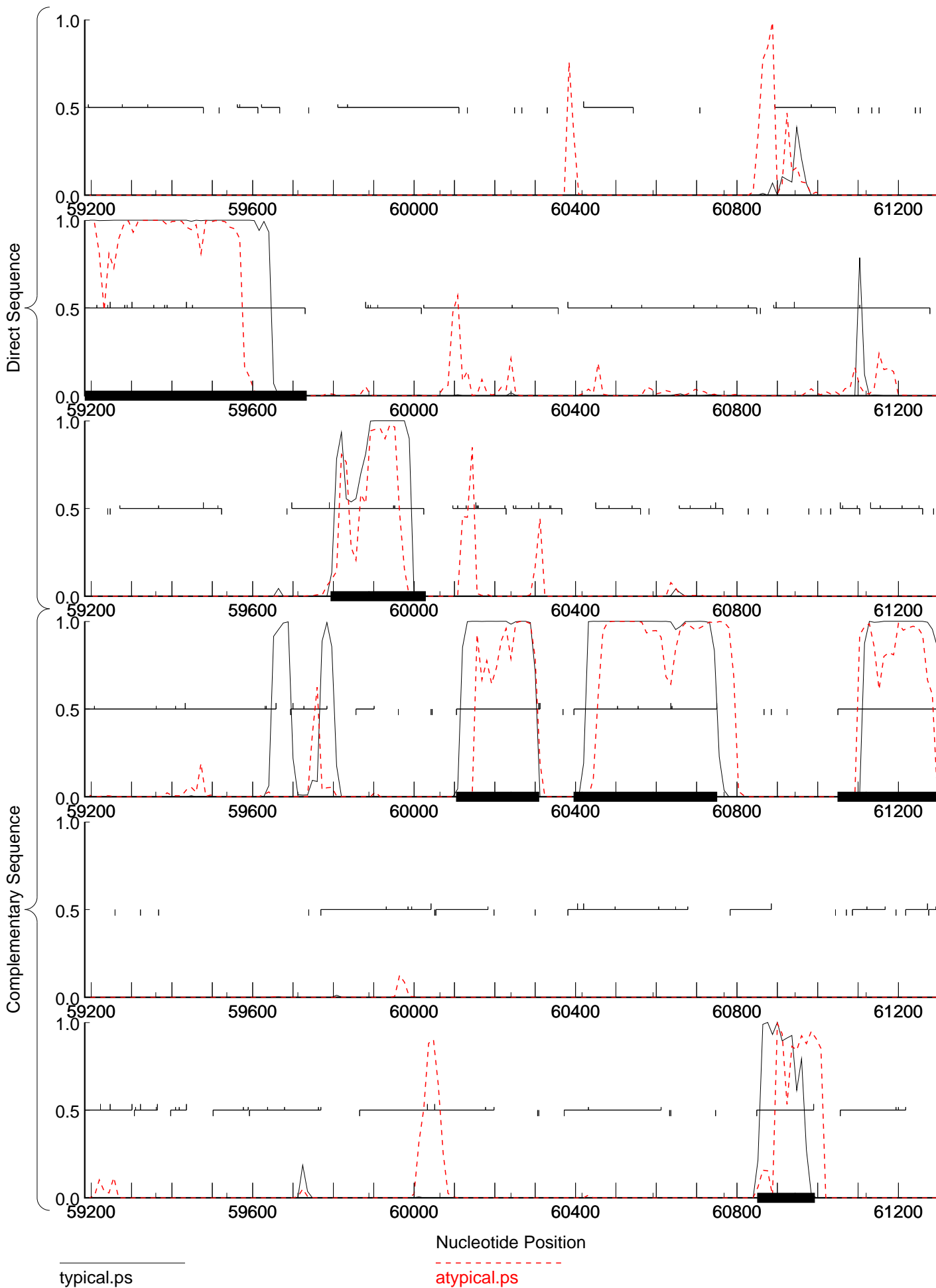
atypical.ps





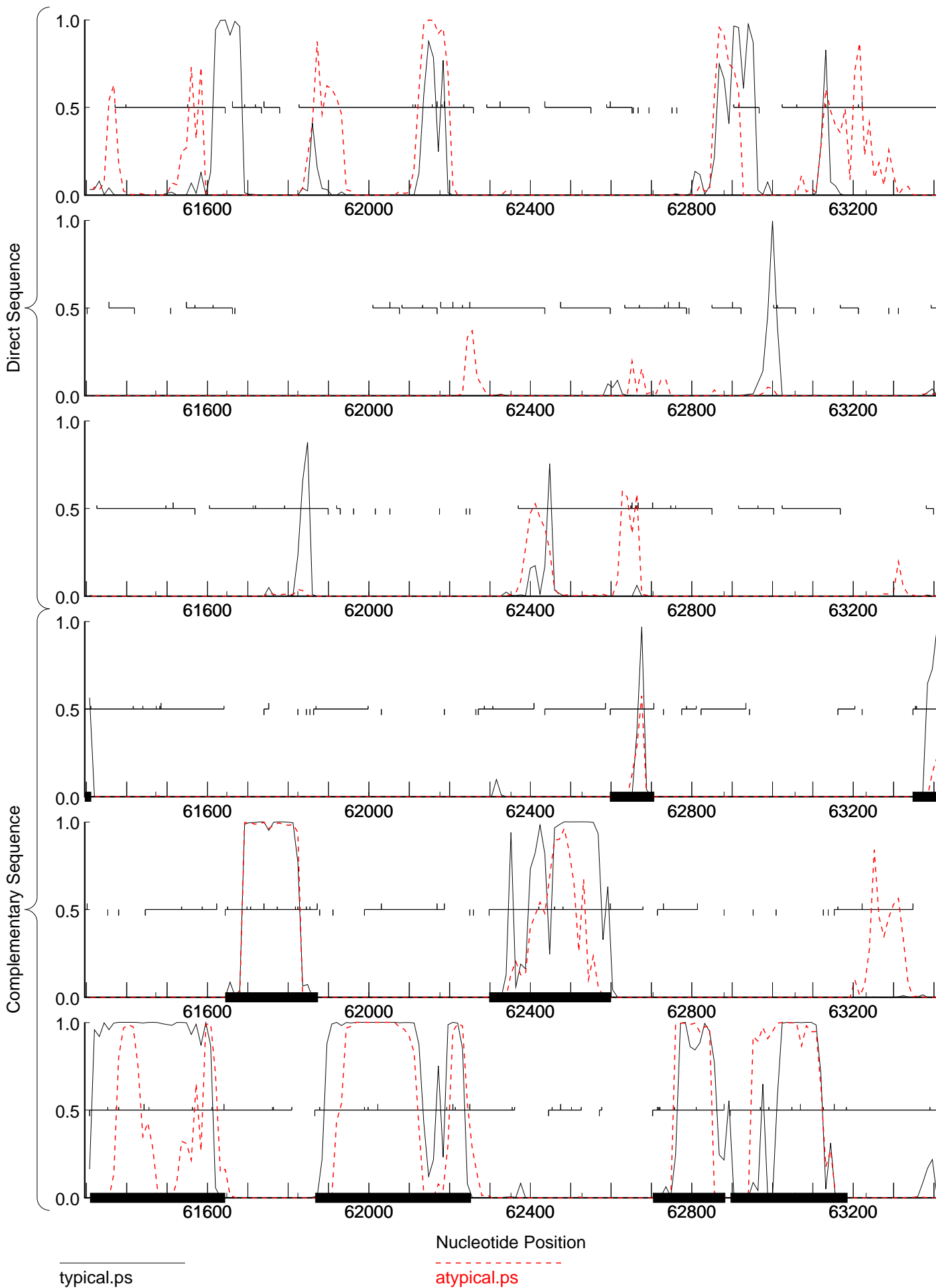
typical.ps

atypical.ps



typical.ps

atypical.ps



typical.ps

atypical.ps

