

GeneMark

Version 2.5p (09.08.06)

Copyright 1993 - M. Borodovsky, J. McIninch

PROGRAM INFORMATION

Sequence : Mycobacteriophage StepMih complete sequence, 50841 bp including 10-base 3' overhang (CGGGTG
Analysis Date : 1/22/18 at 9:33:40
Pages : 26
Sequence Length : 50841 bp
GC Content : 64.01%

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,

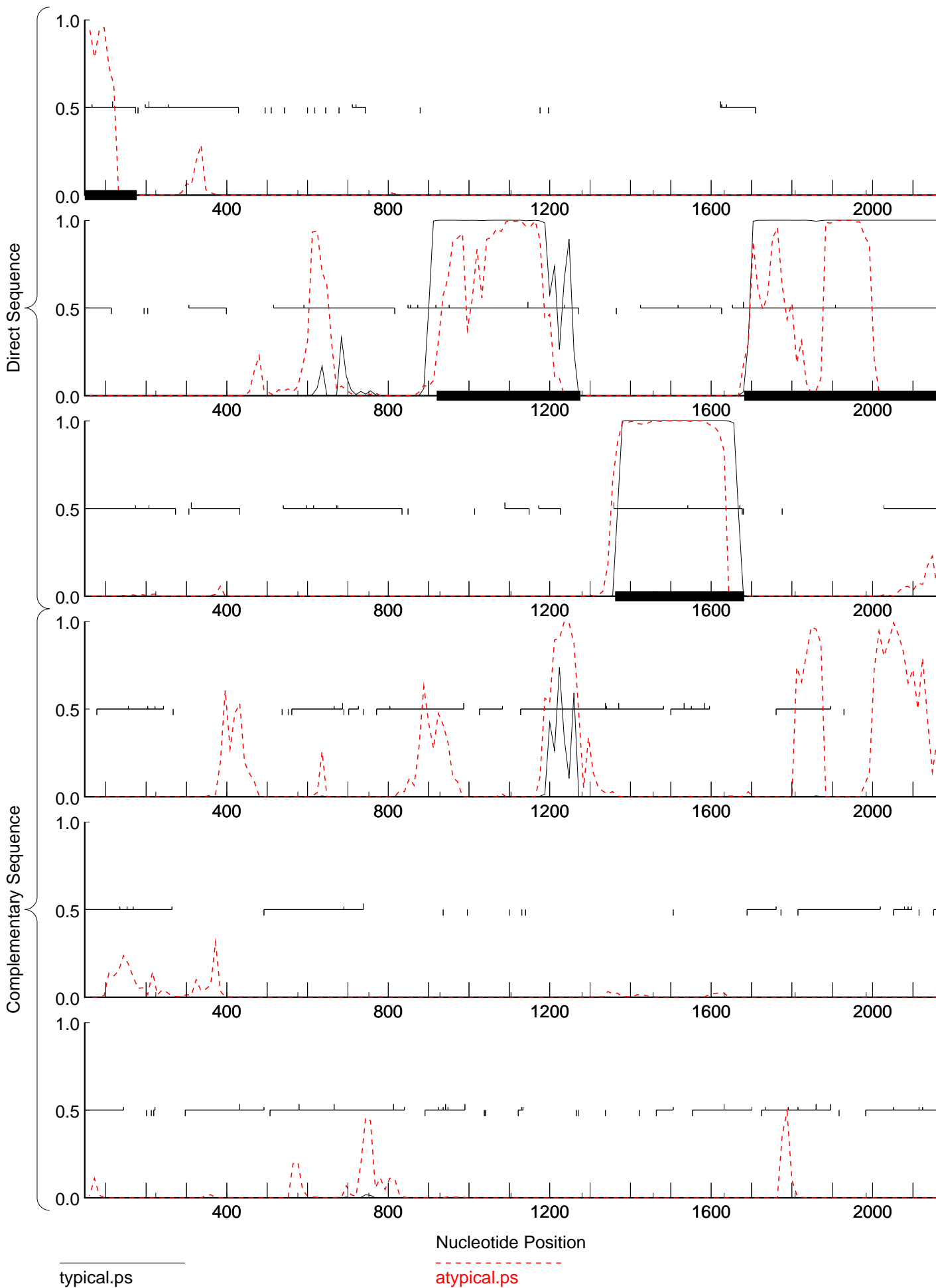
MATRIX INFORMATION

Matrix : -
Author : -
Order : 4

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

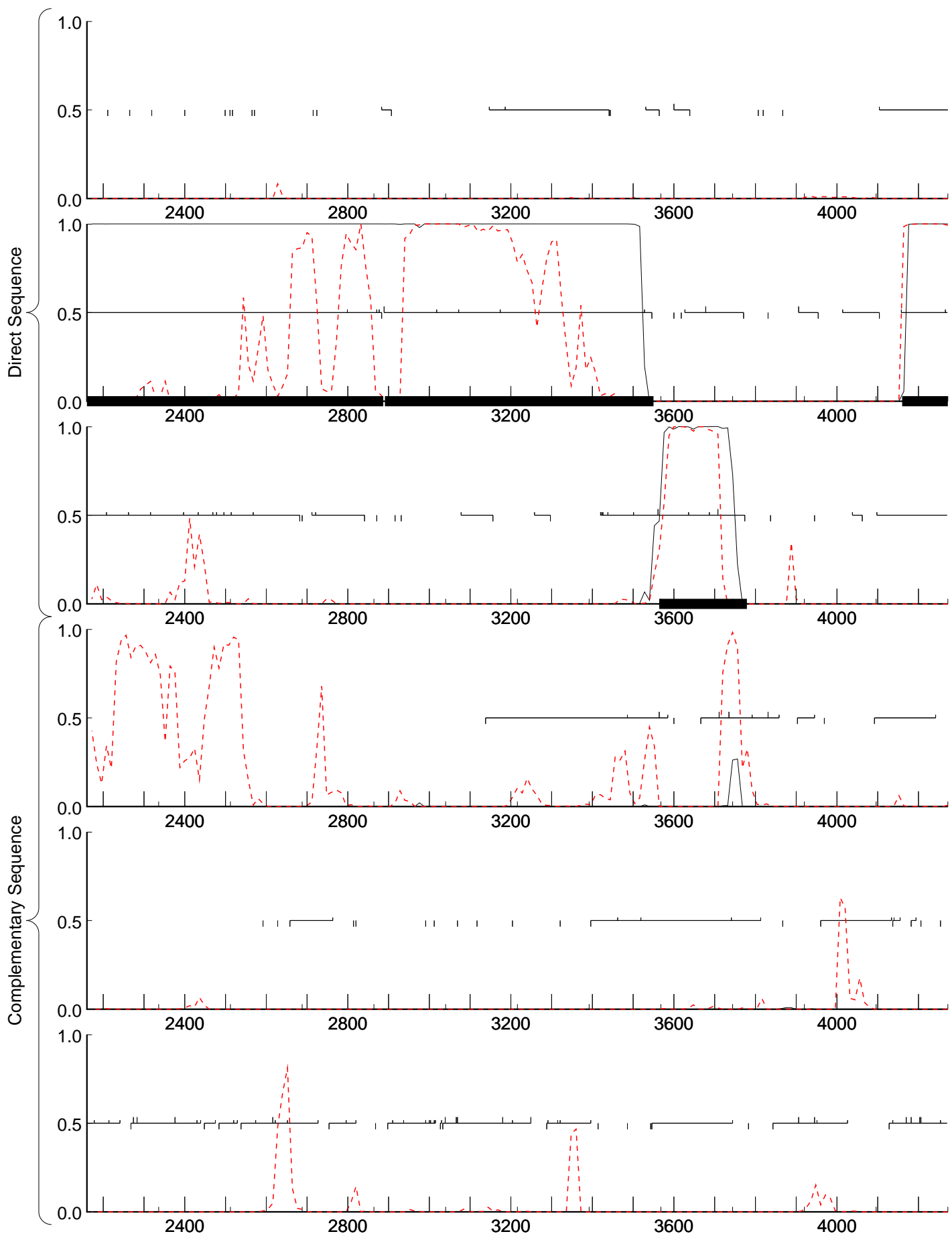
Matrix notes & comments

Training set derived by GeneMarkS, 4.30 October 2014
Mon Jan 22 09:33:40 2018



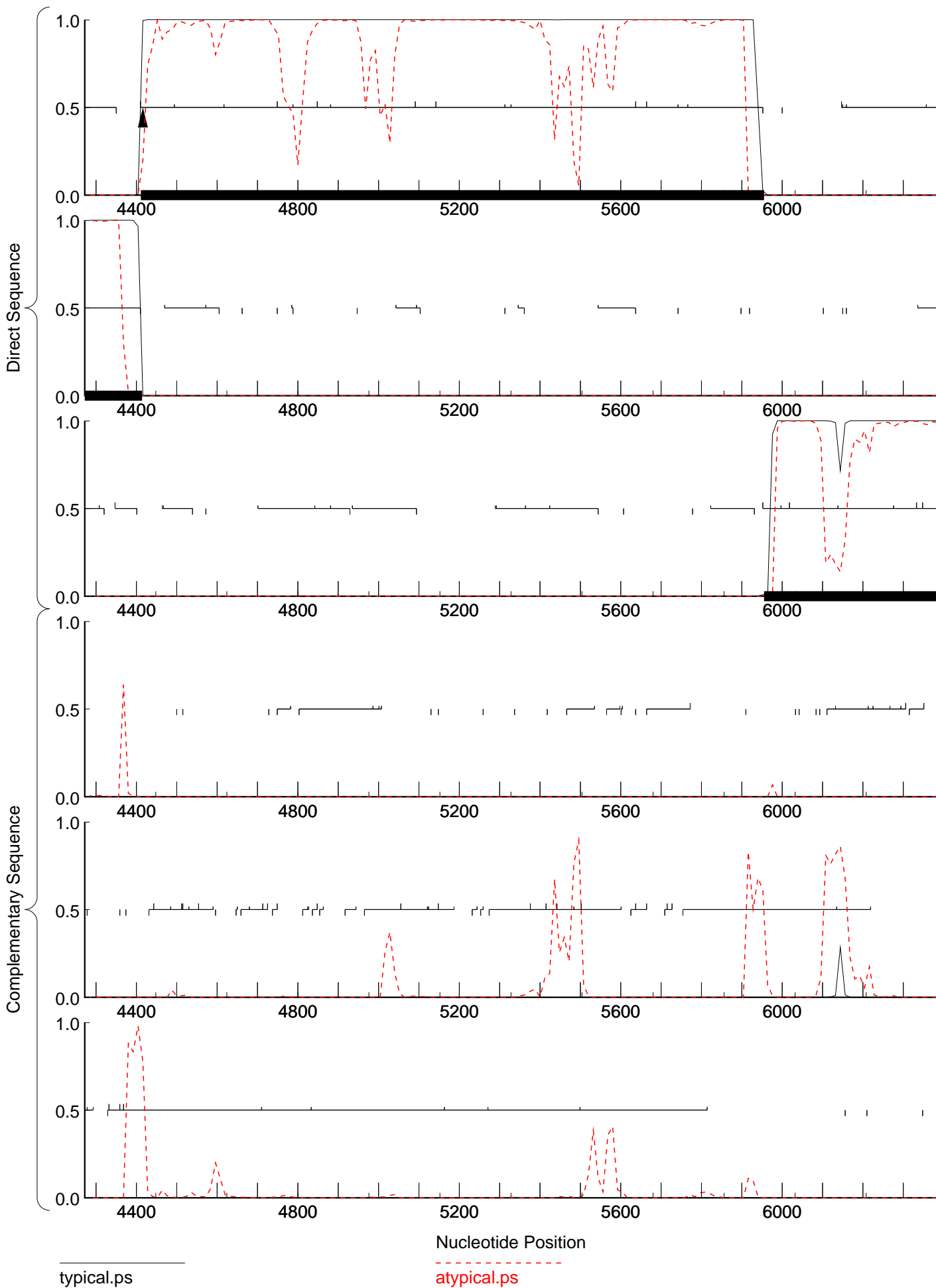
typical.ps

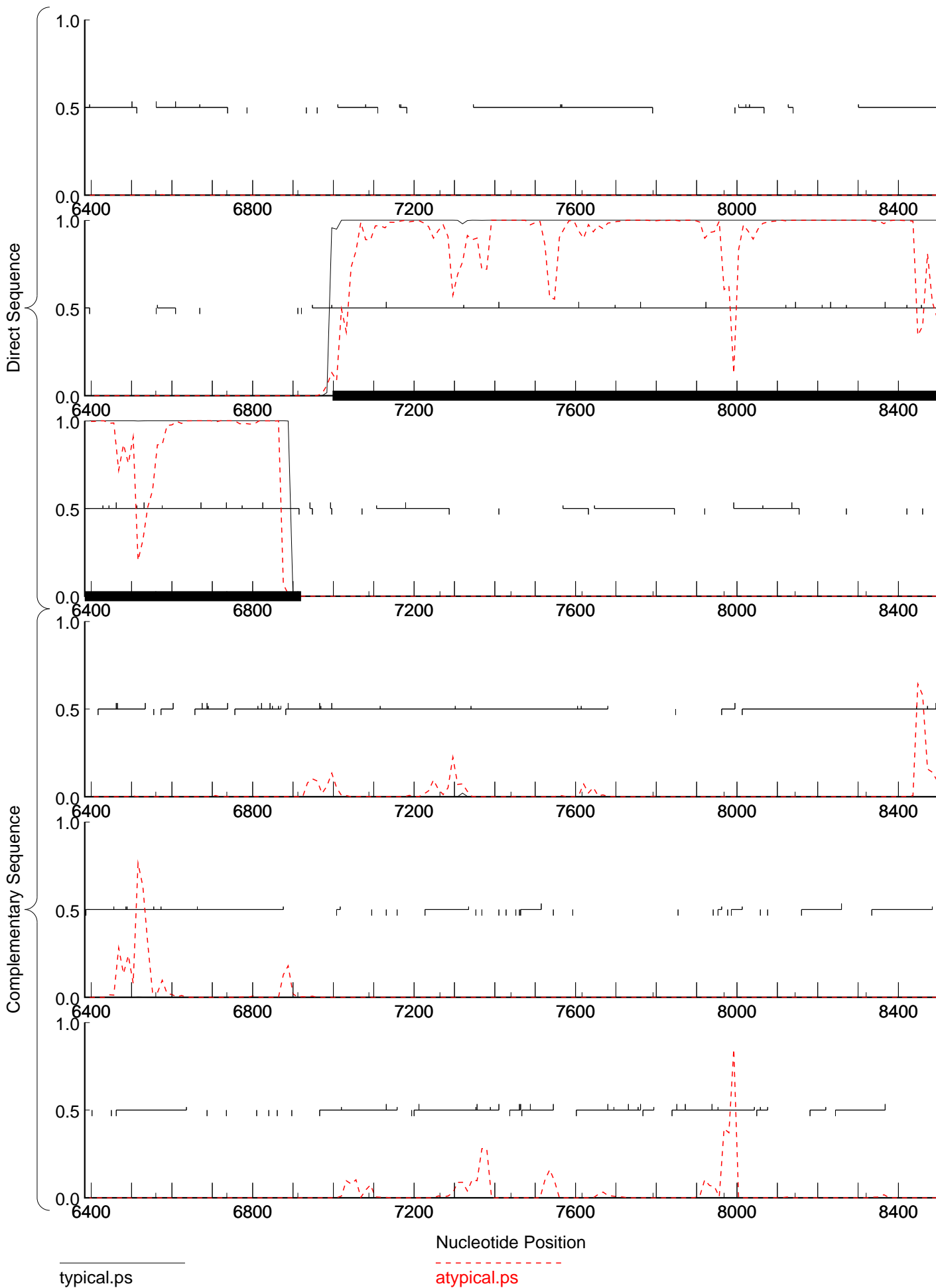
atypical.ps

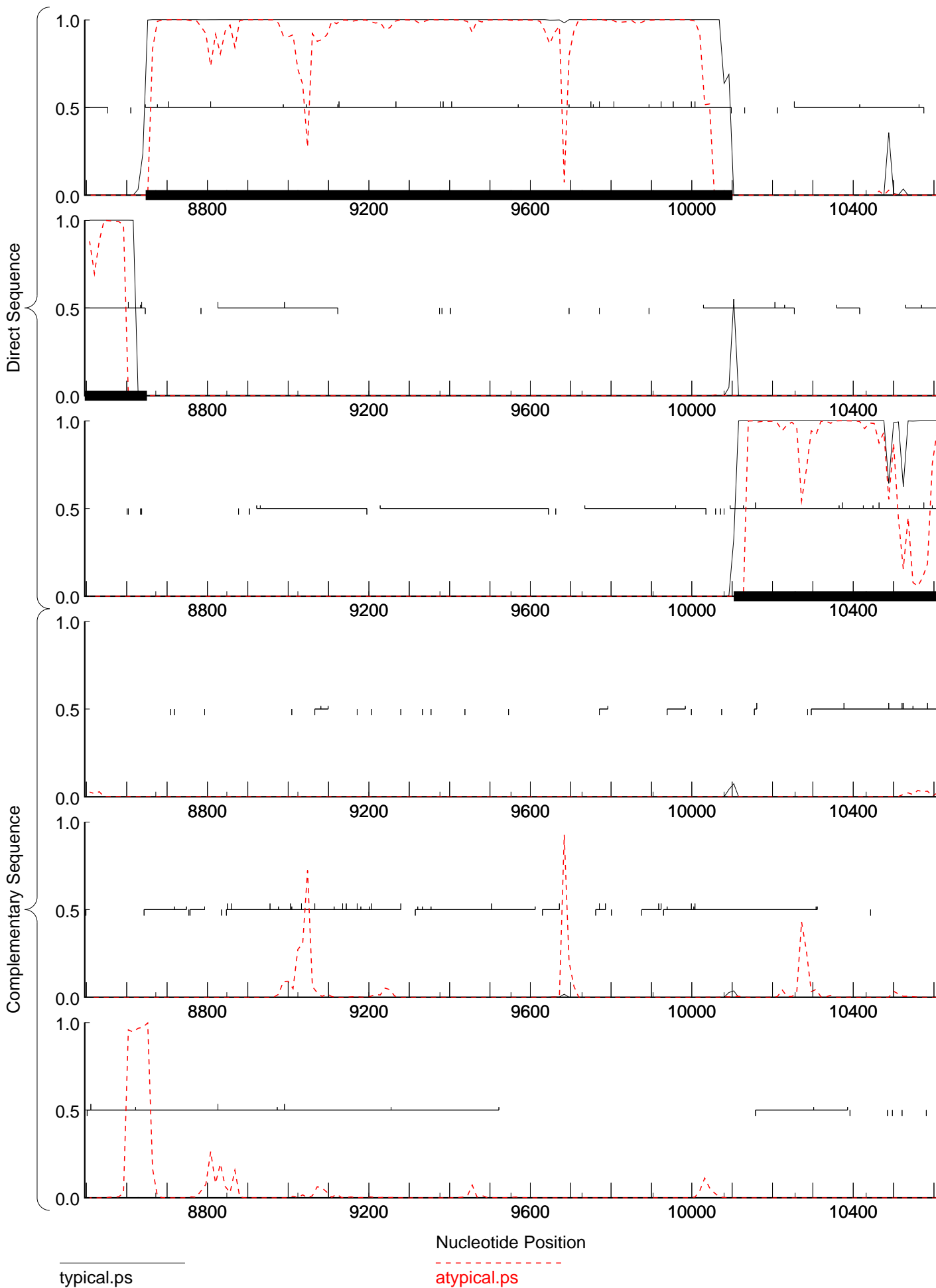


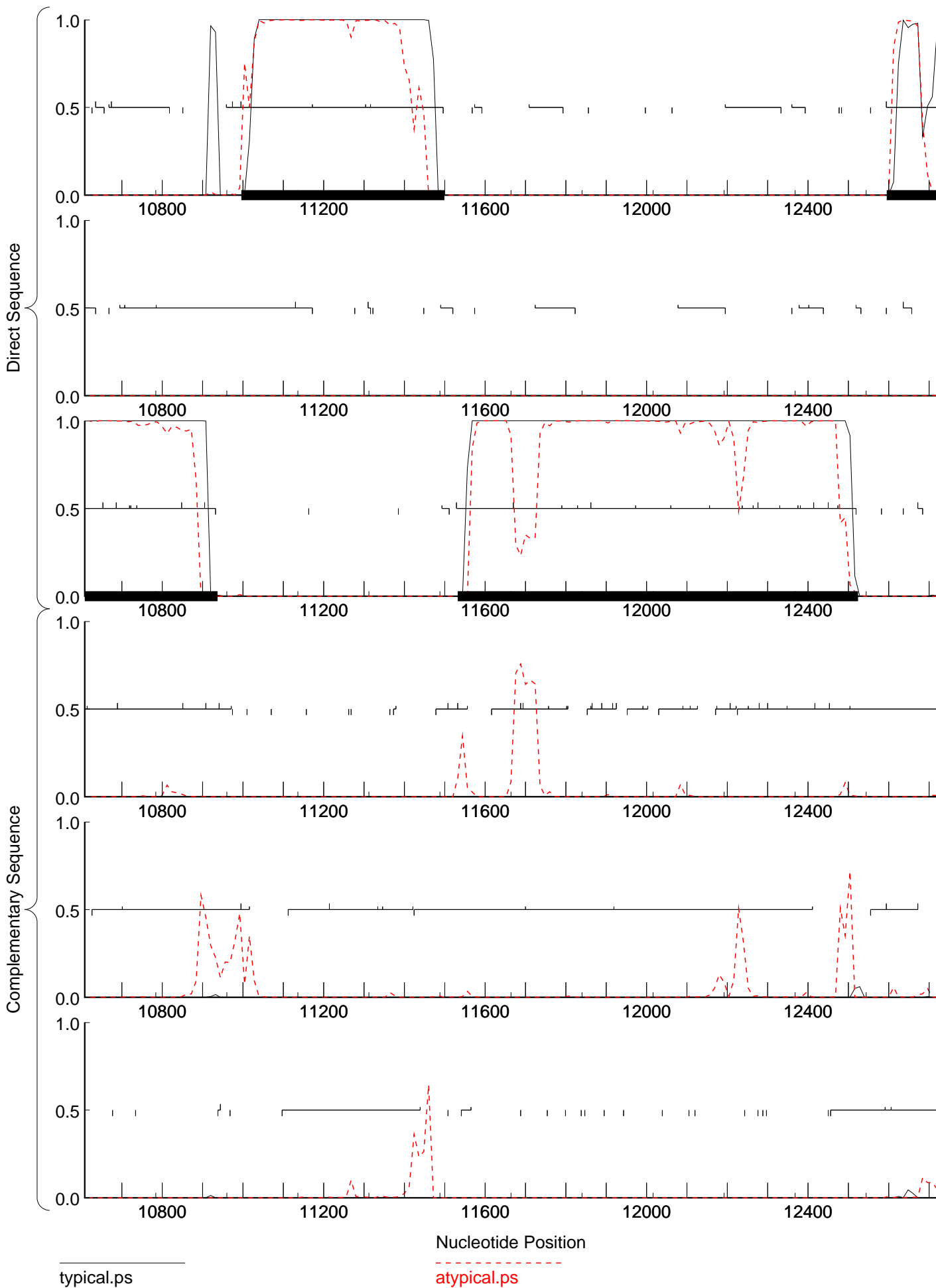
typical.ps

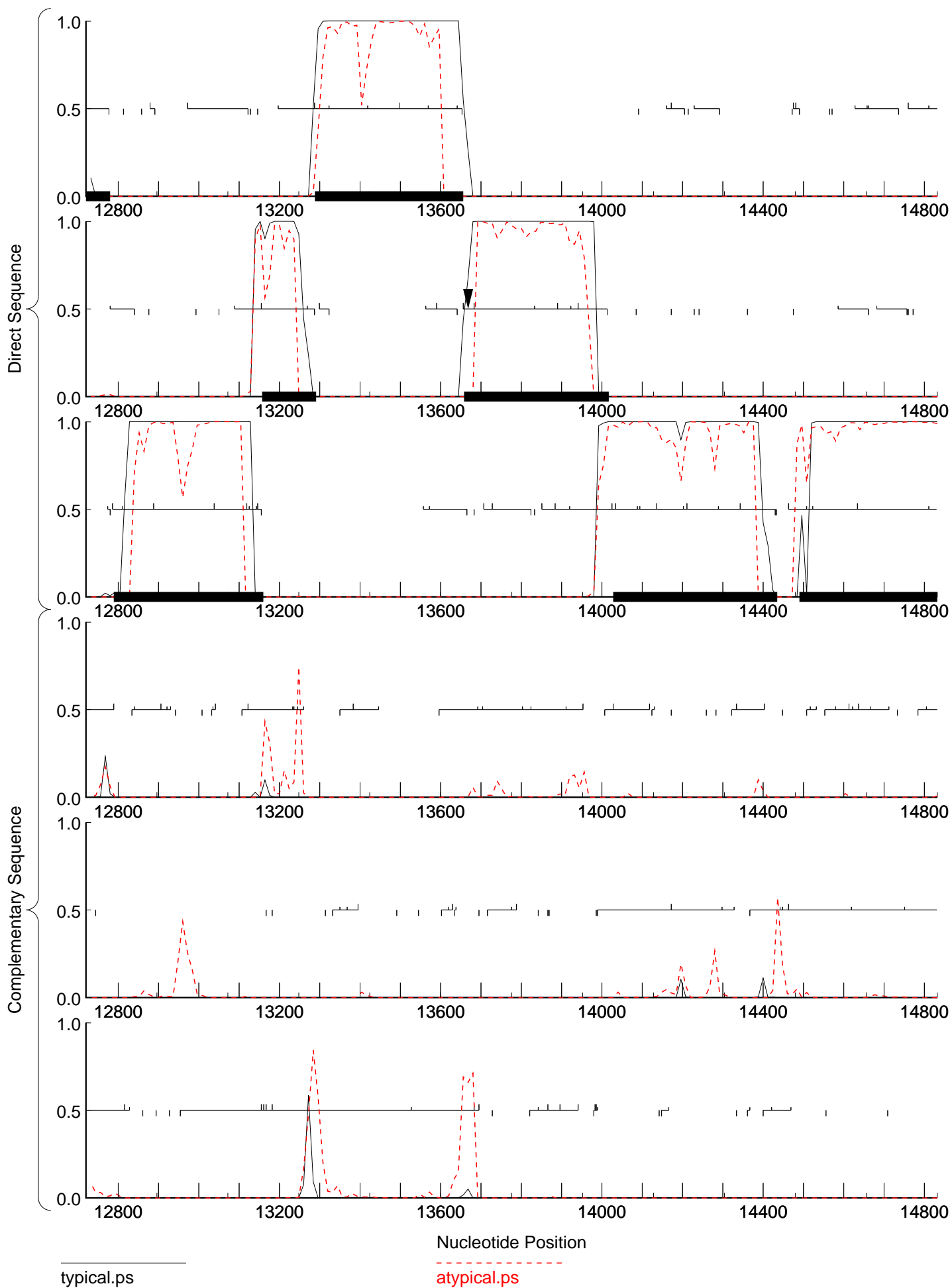
atypical.ps





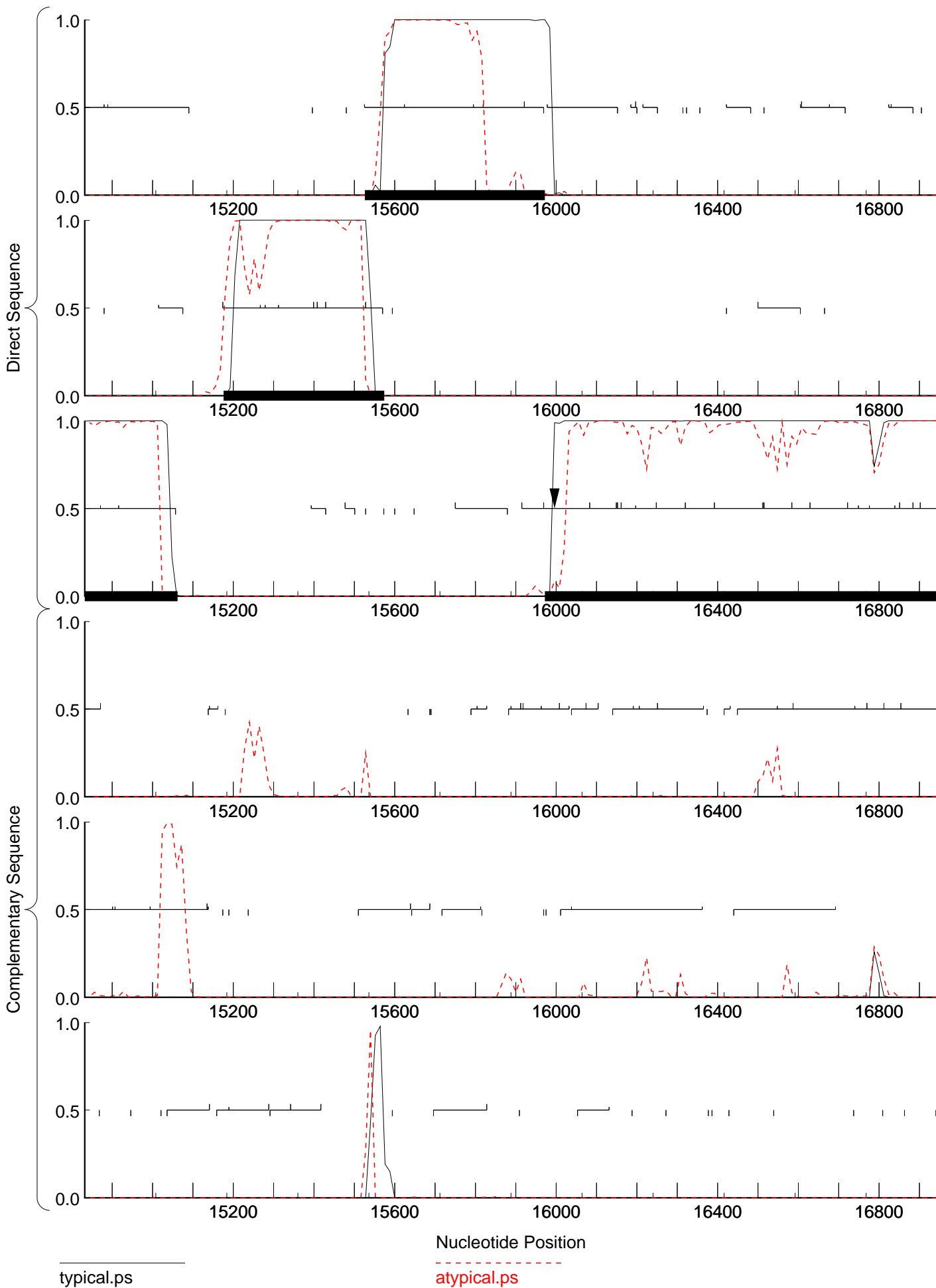


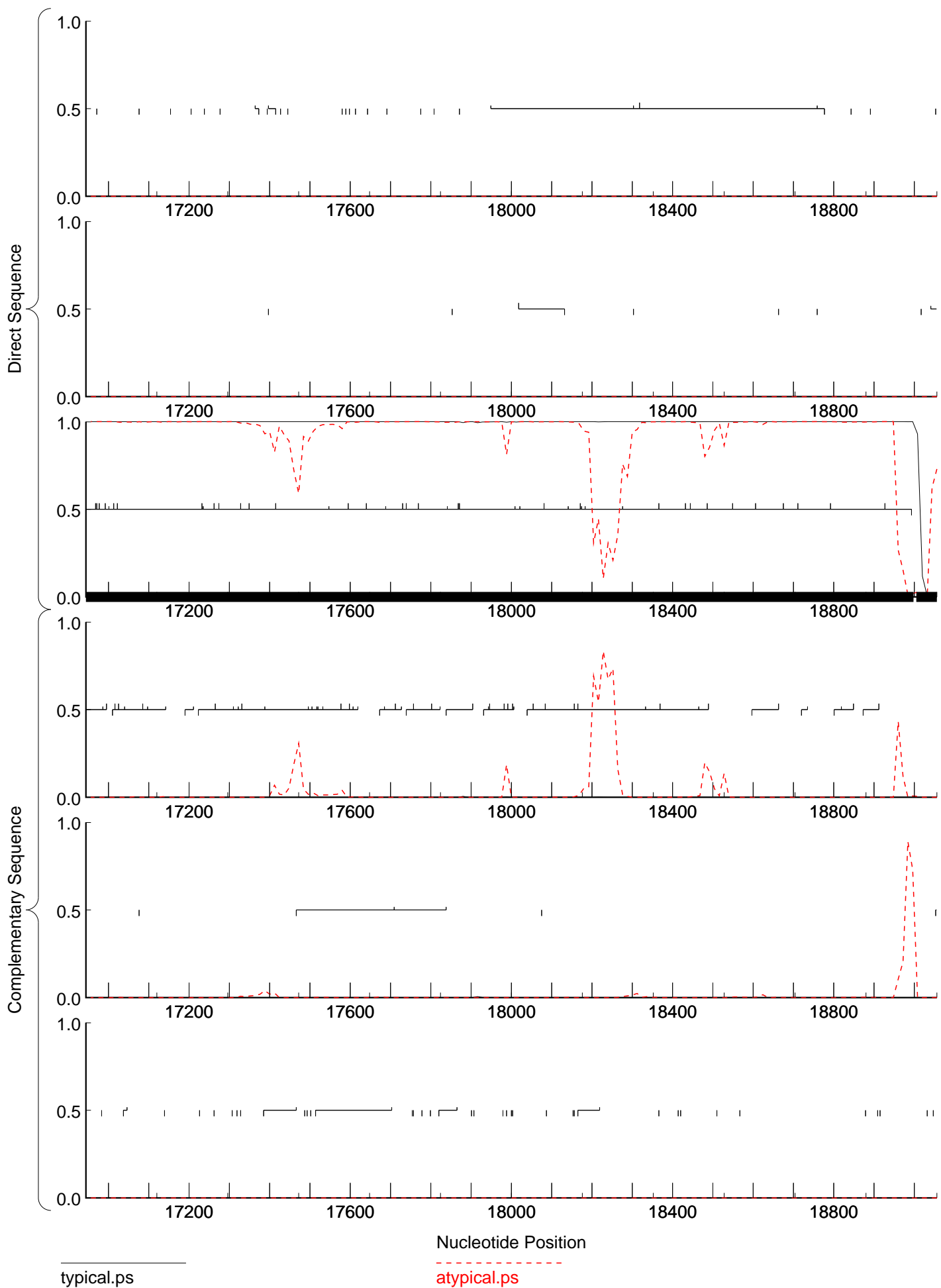


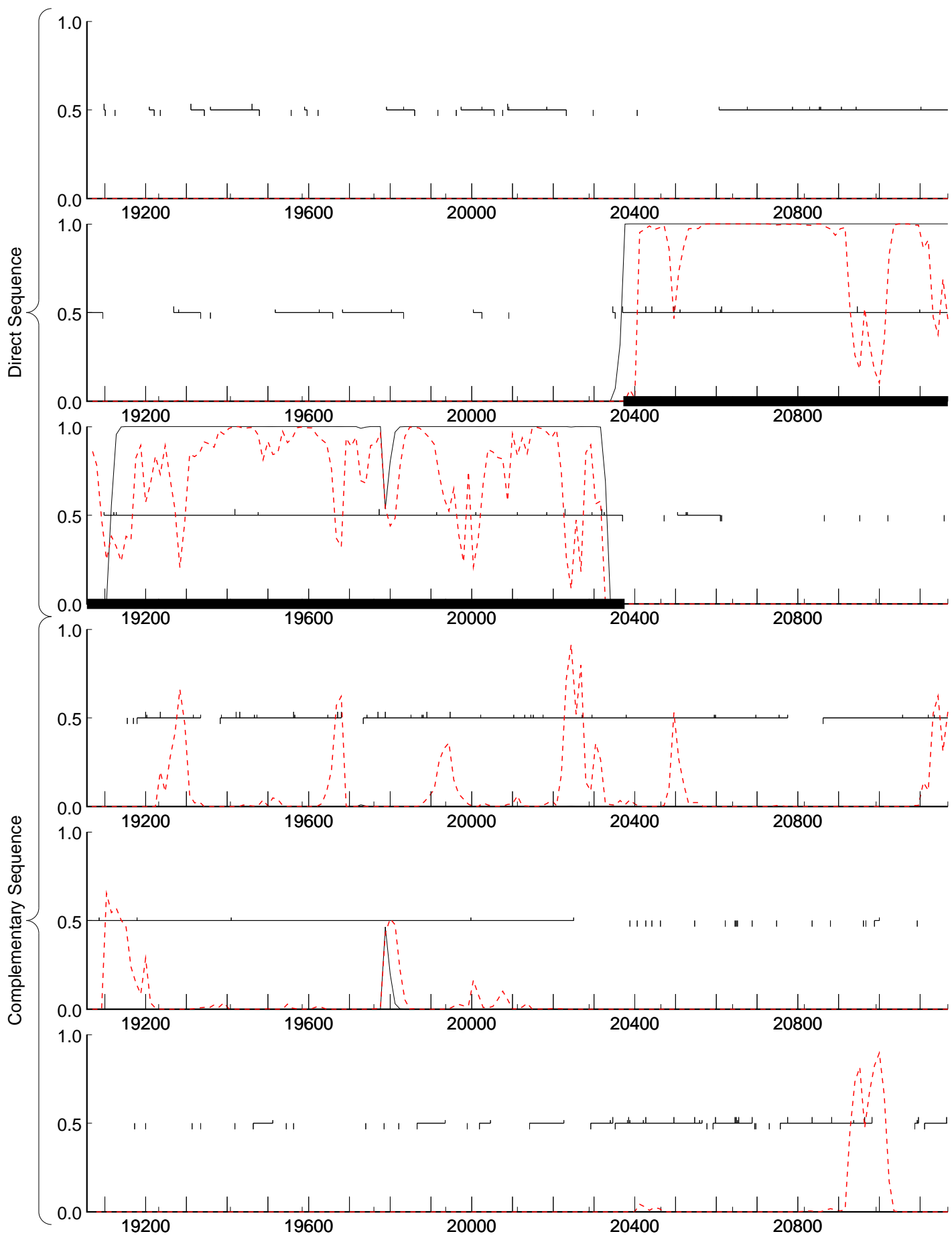


typical.ps

atypical.ps

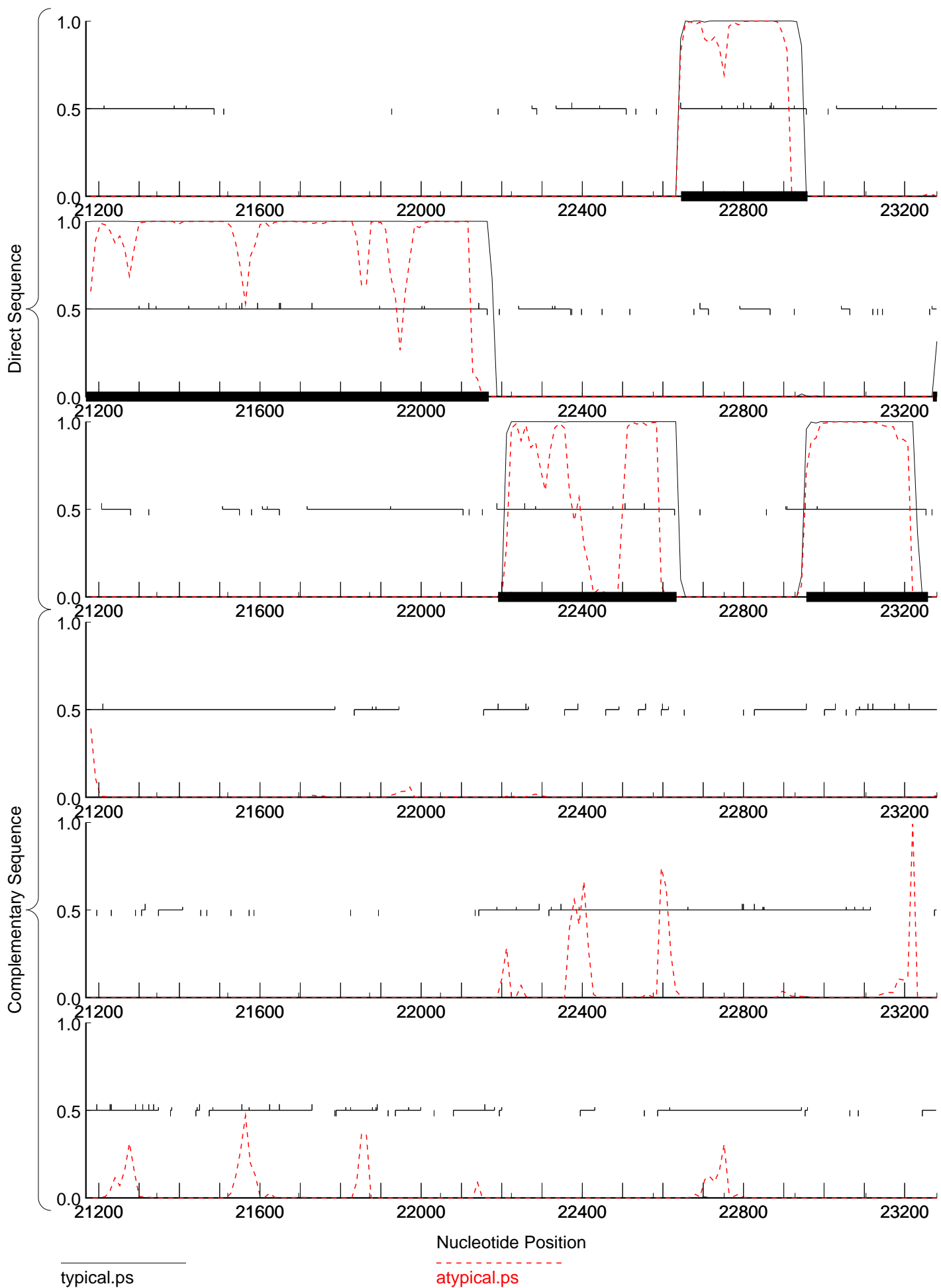


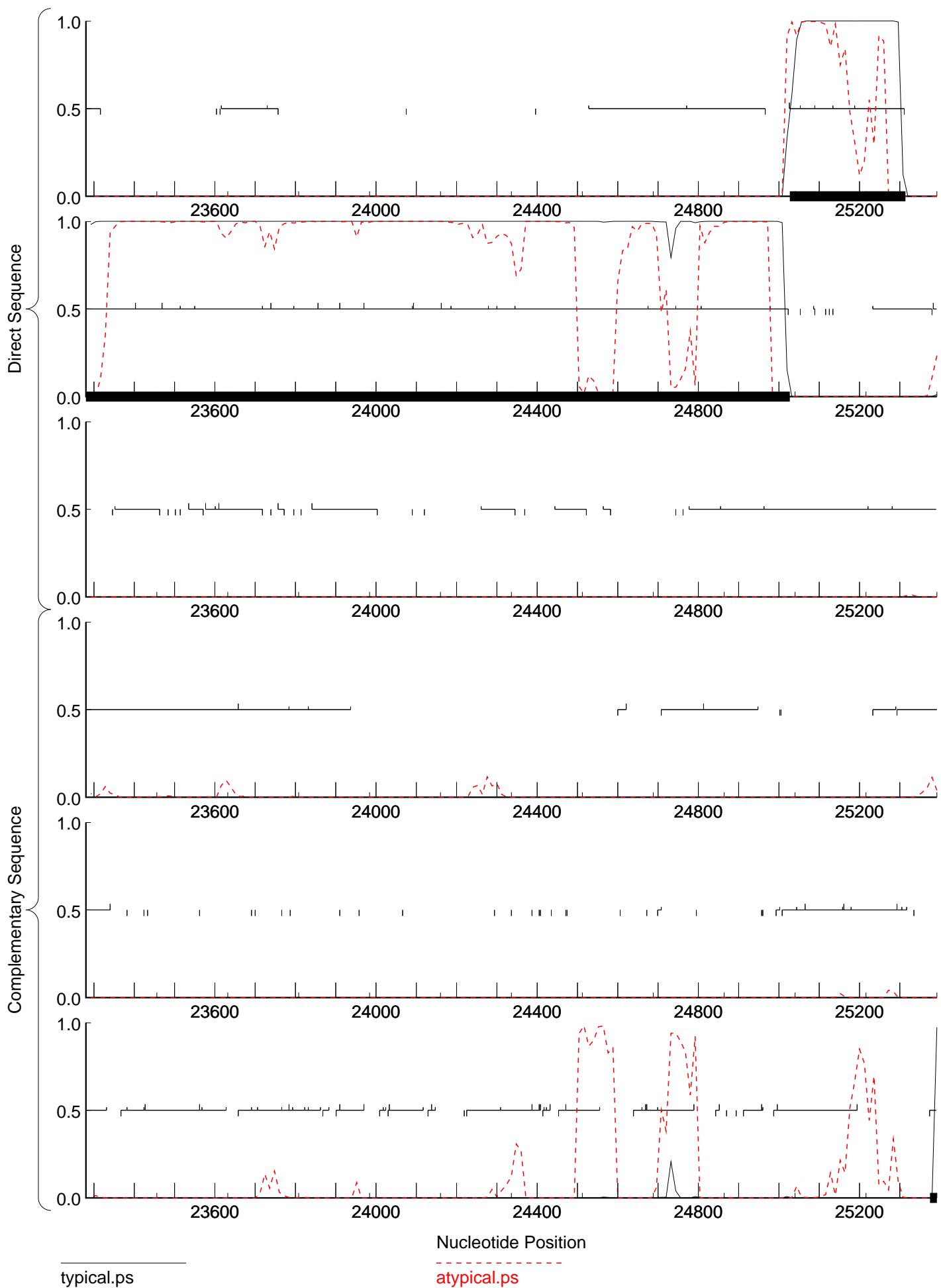


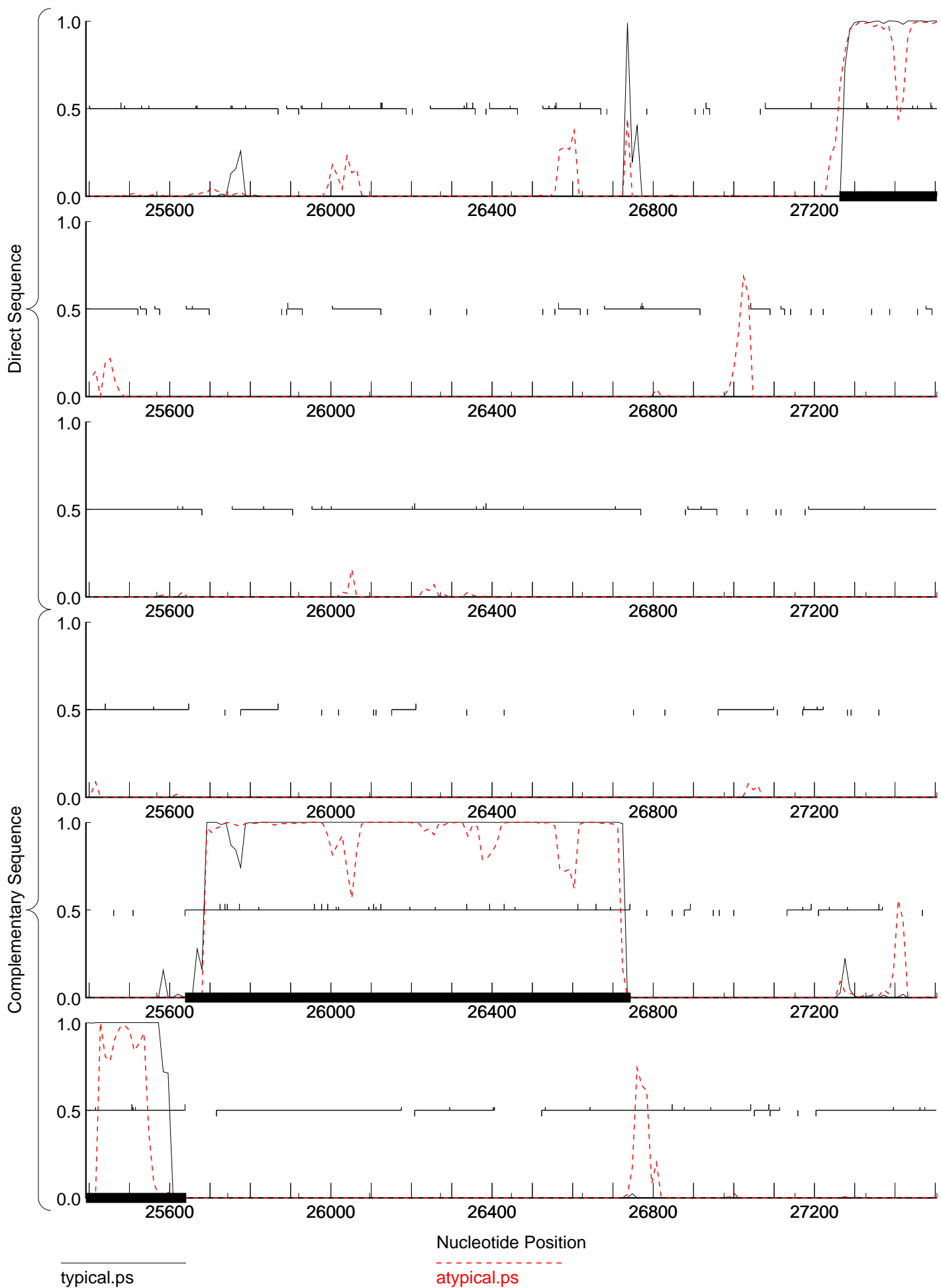


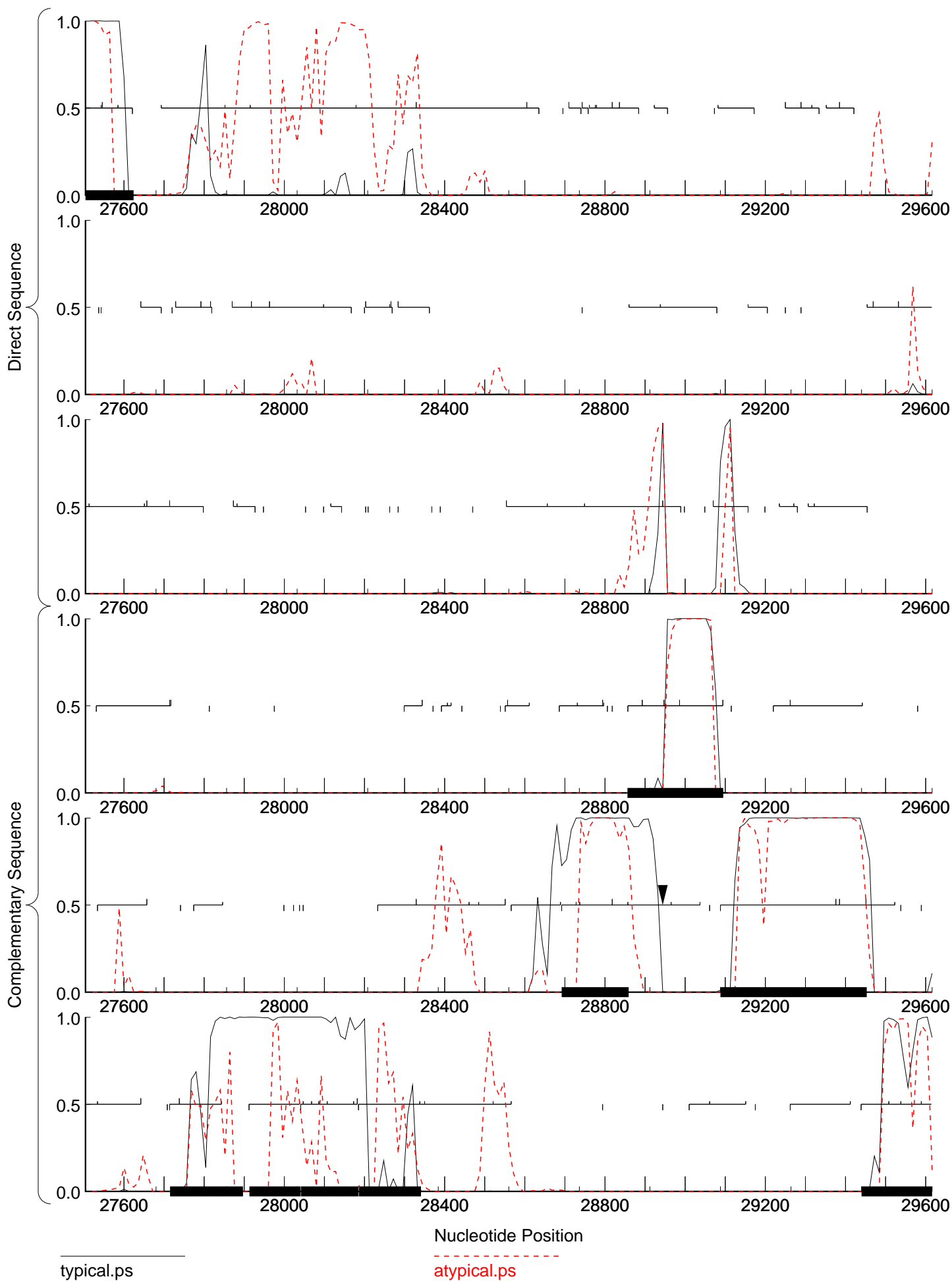
typical.ps

atypical.ps



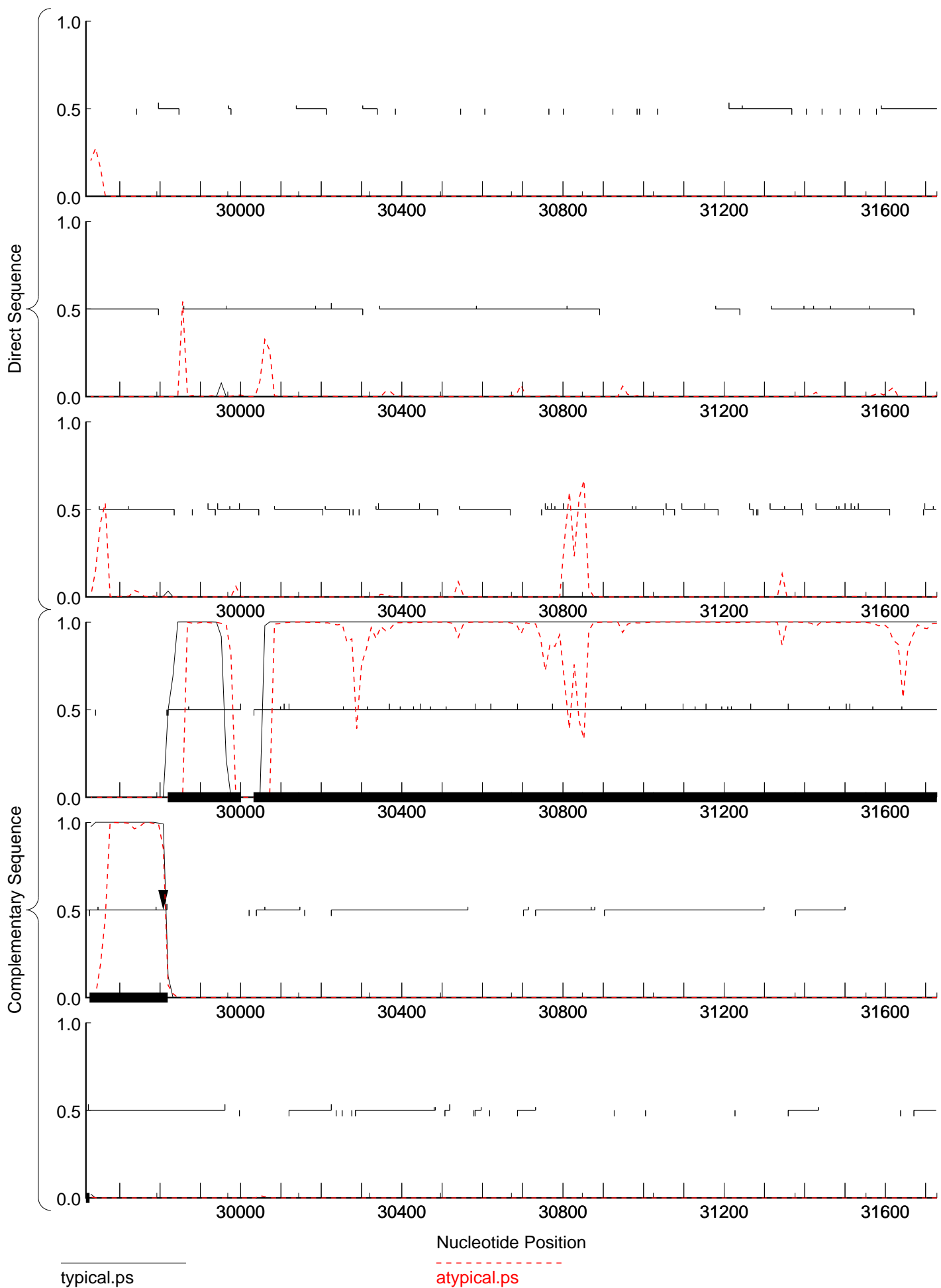


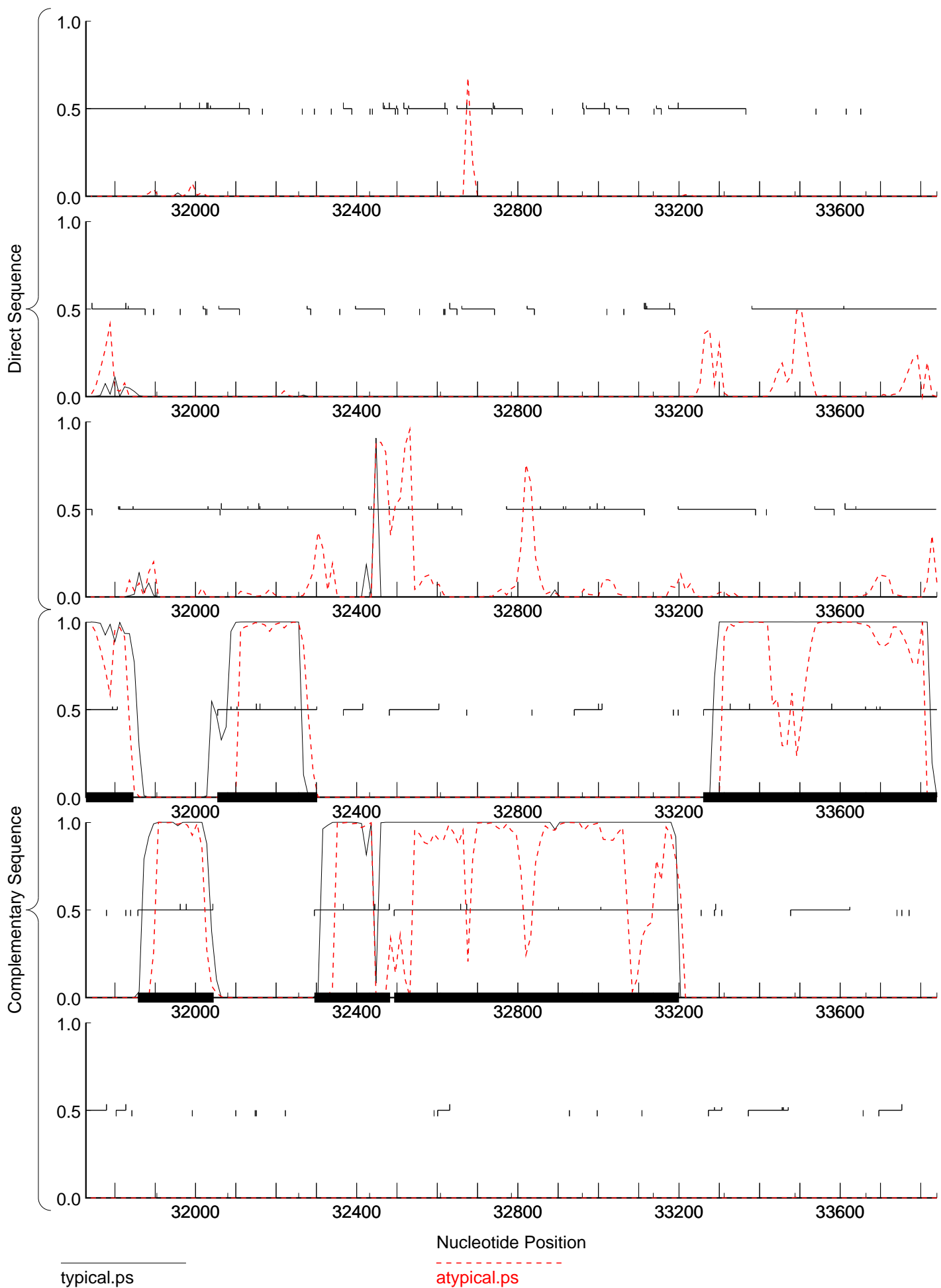


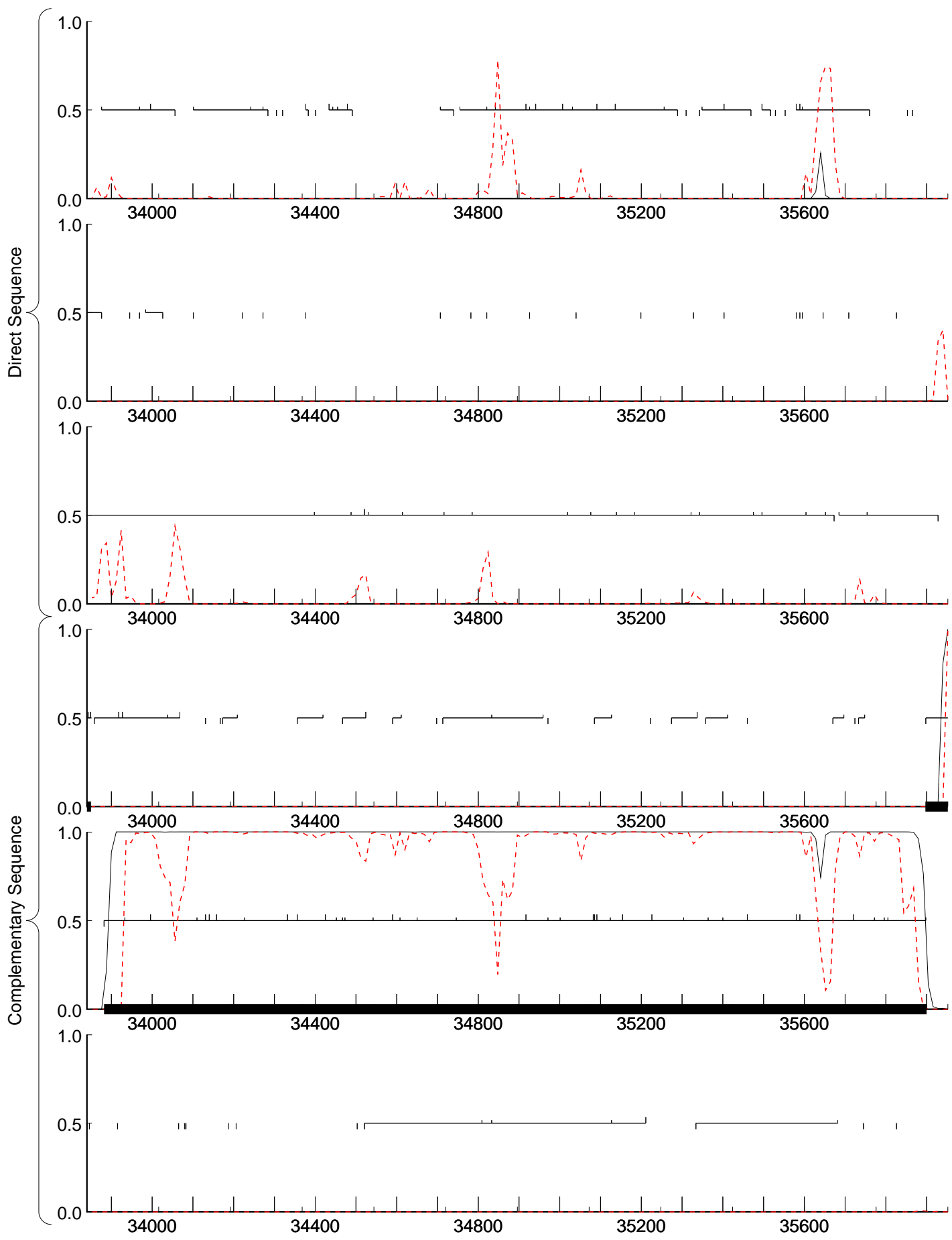


typical.ps

atypical.ps



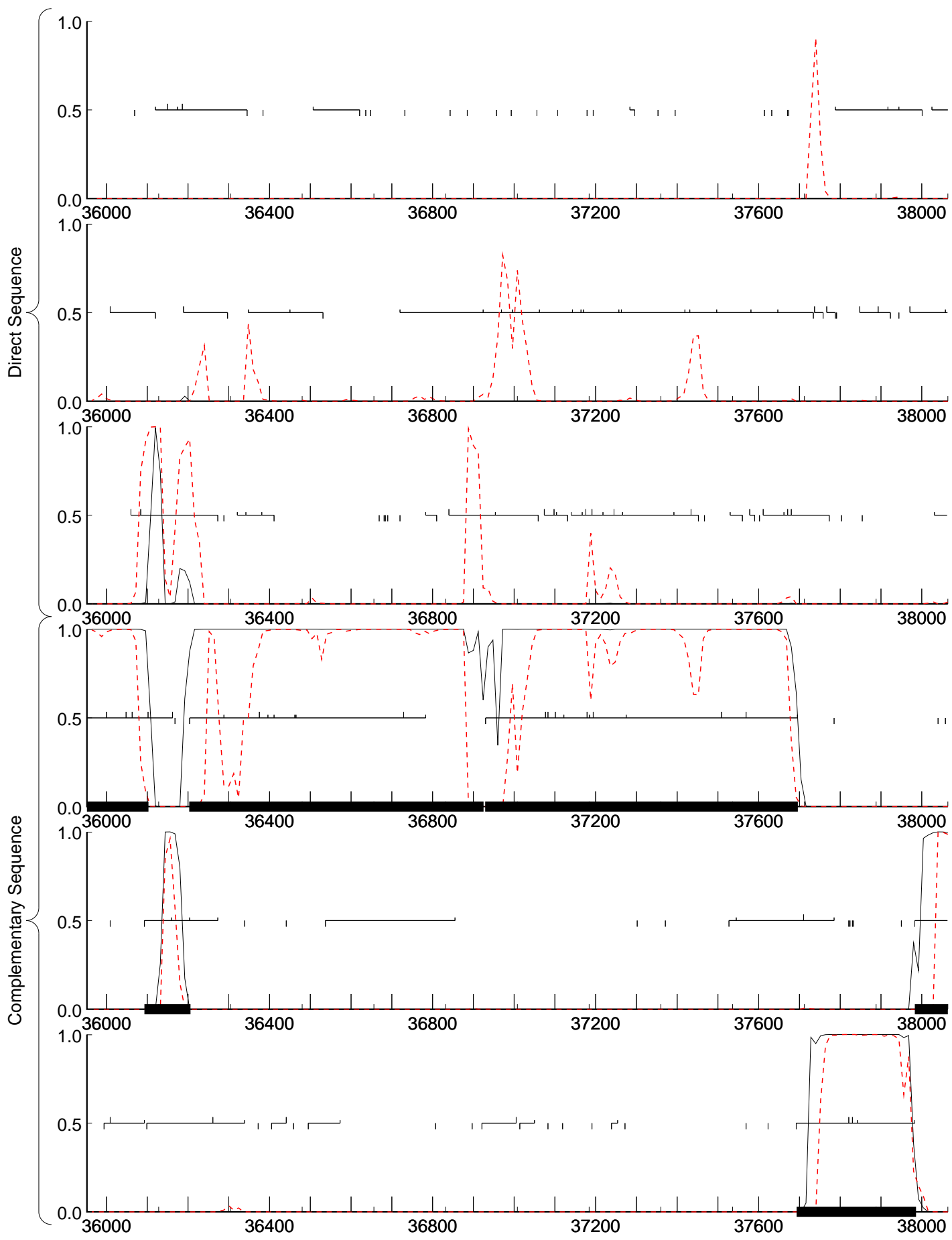


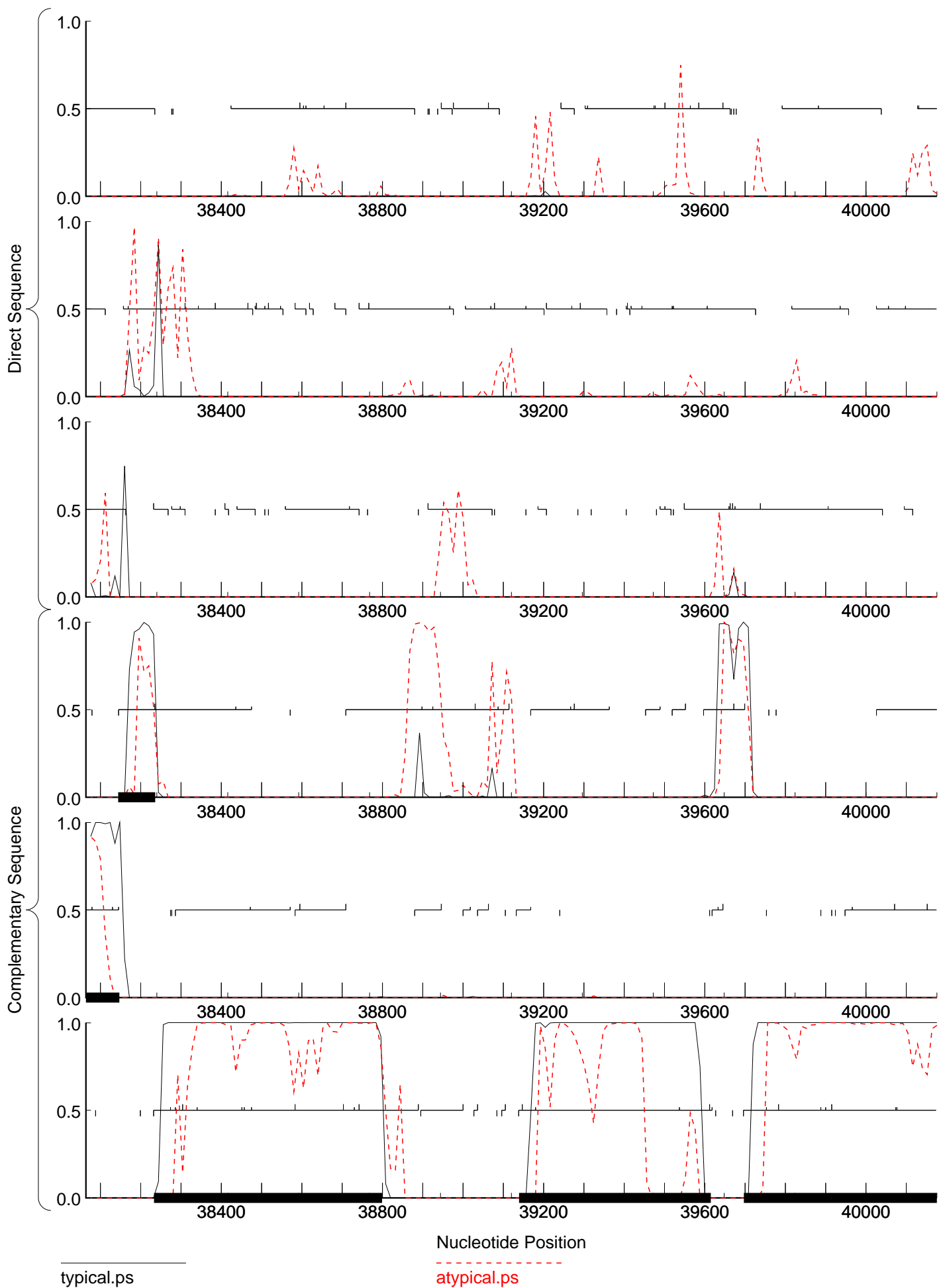


typical.ps

atypical.ps

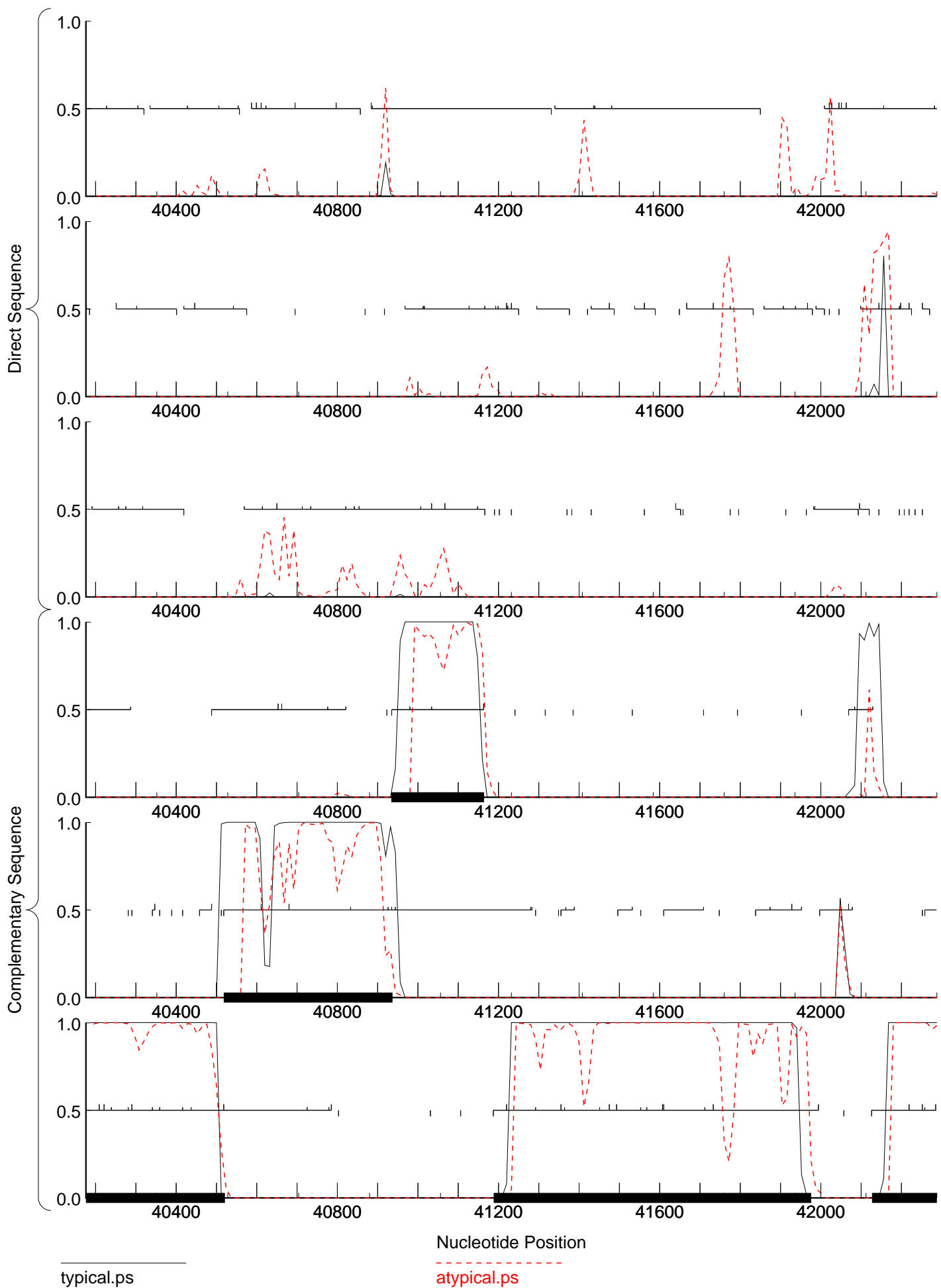
GeneMark.hmm prediction





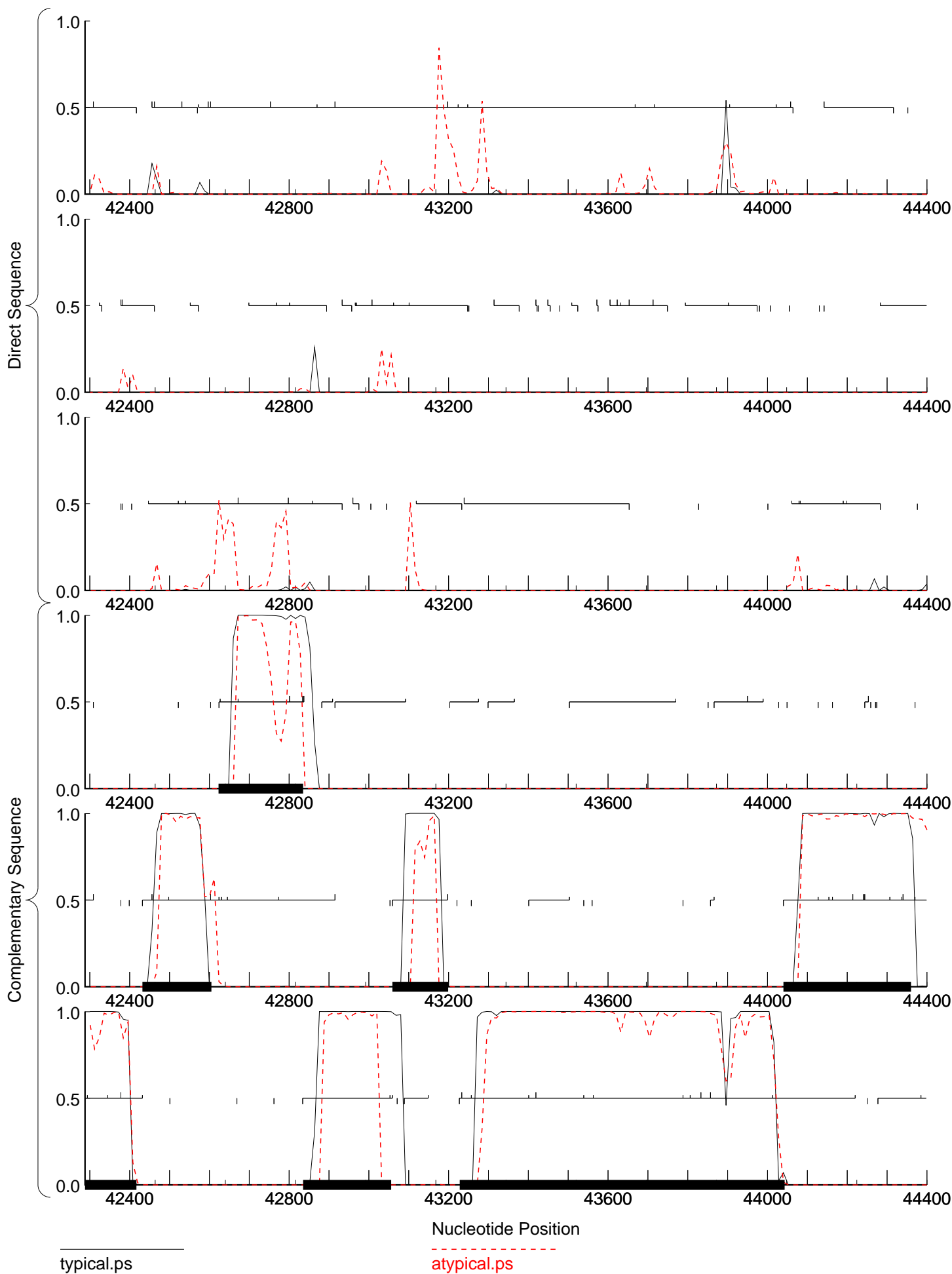
typical.ps

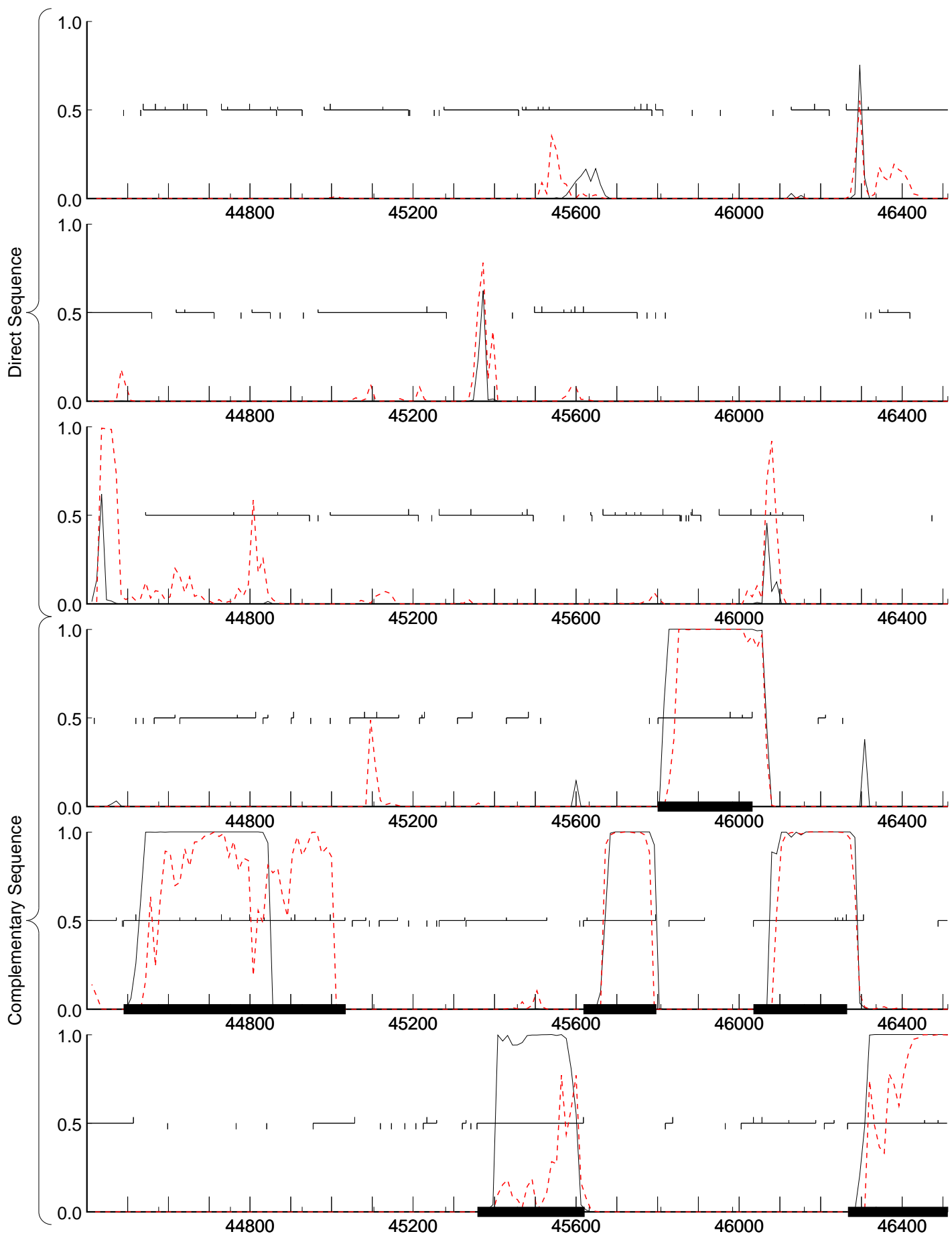
atypical.ps



typical.ps

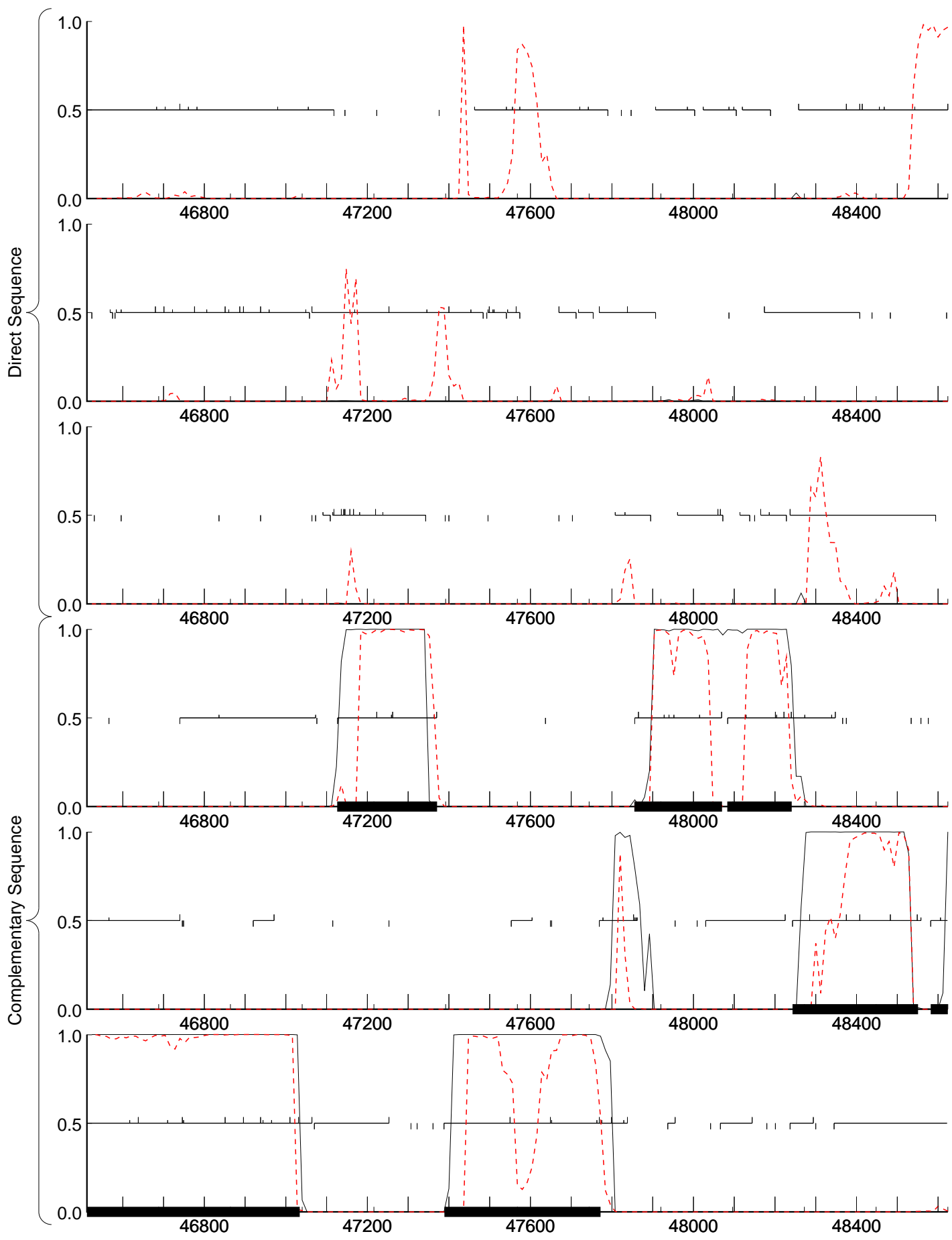
atypical.ps





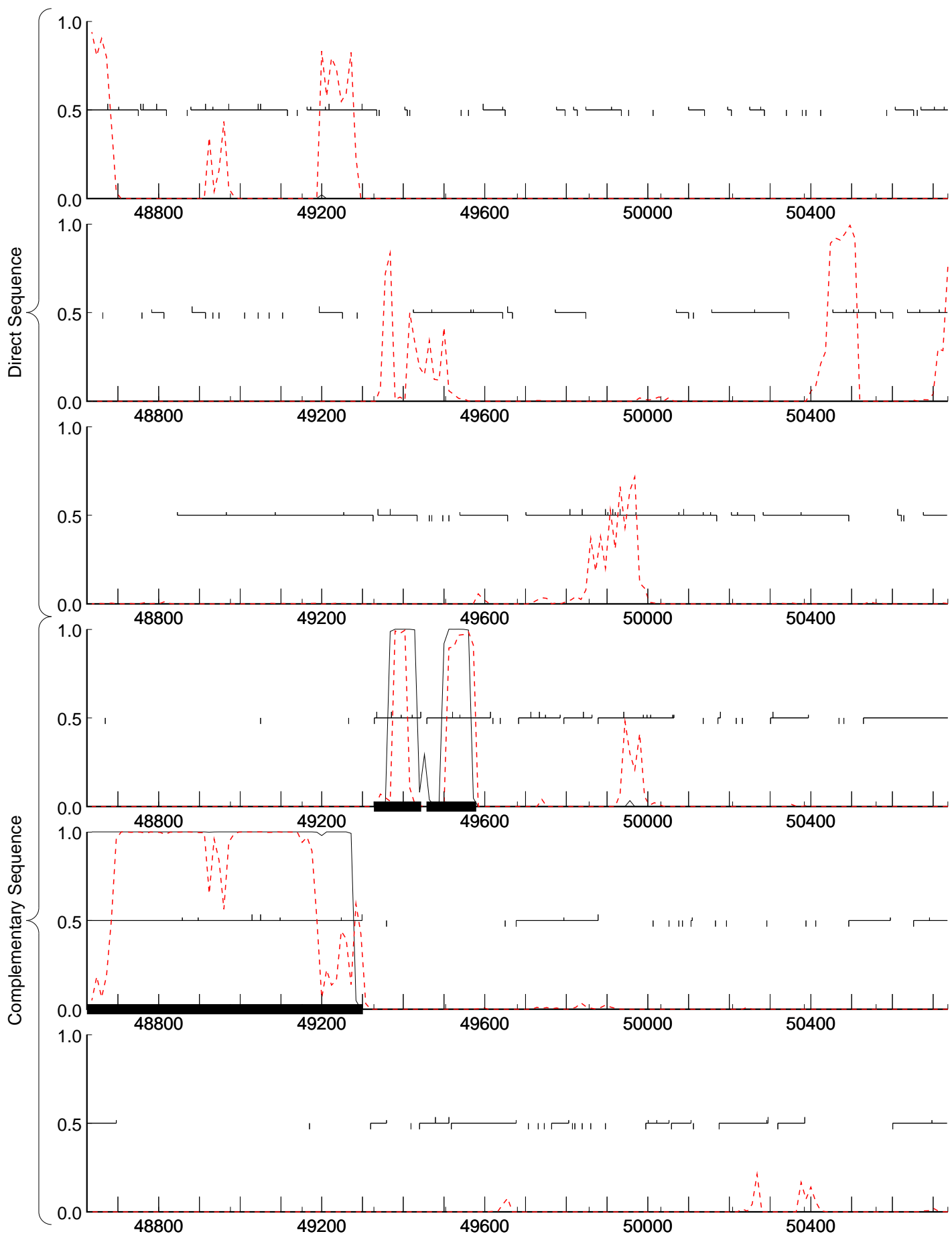
typical.ps

atypical.ps



typical.ps

atypical.ps



typical.ps

atypical.ps

