

GeneMark

Version 2.5p (09.08.06)

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PROGRAM INFORMATION

Sequence : Mycobacterium phage Lorde complete sequence, 53852 bp including 10-base 3' overhang (CCGATG
Analysis Date : 5/26/18 at 7:53:23
Pages : 27
Sequence Length : 53852 bp
GC Content : 61.20%

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,

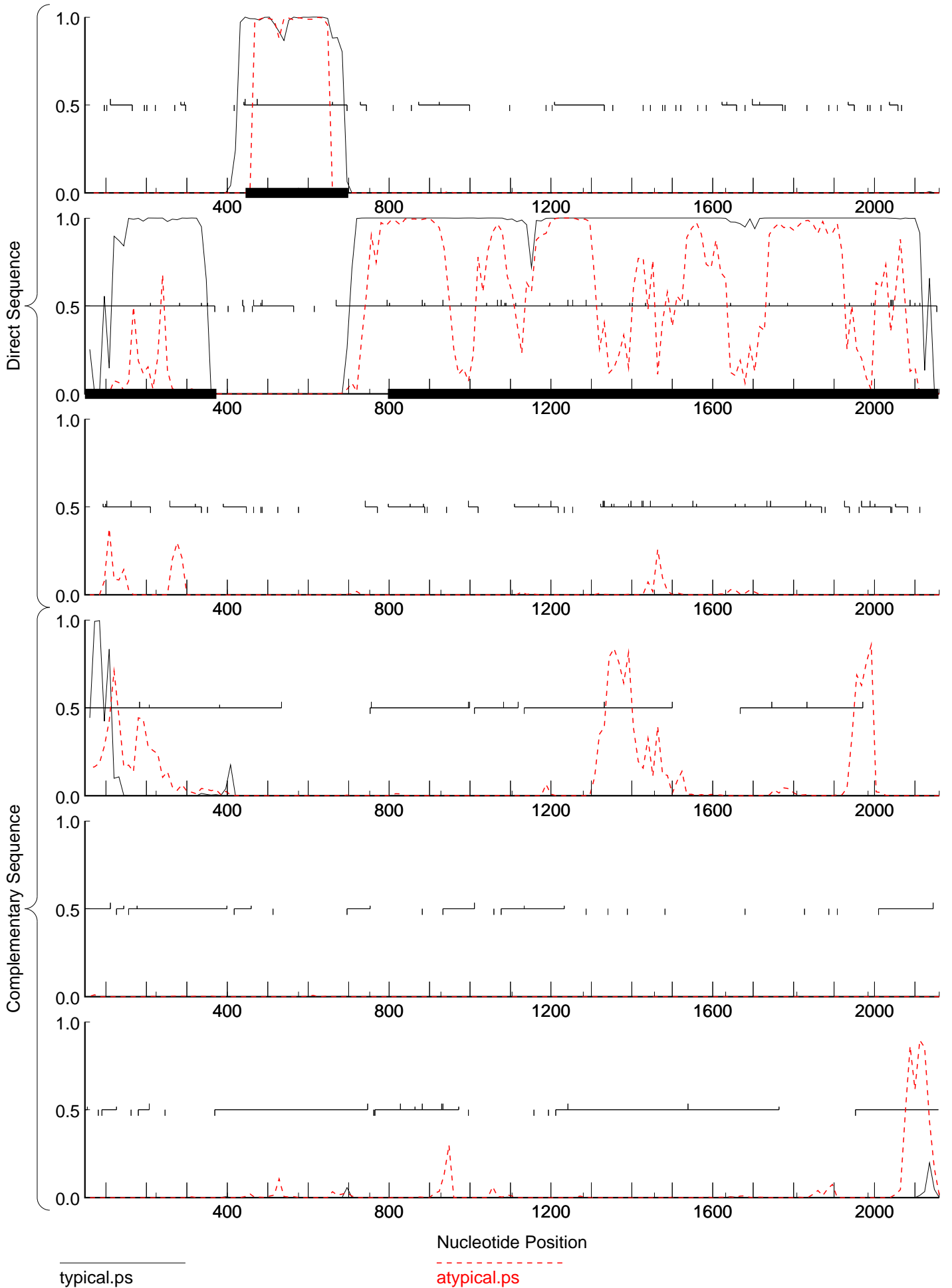
MATRIX INFORMATION

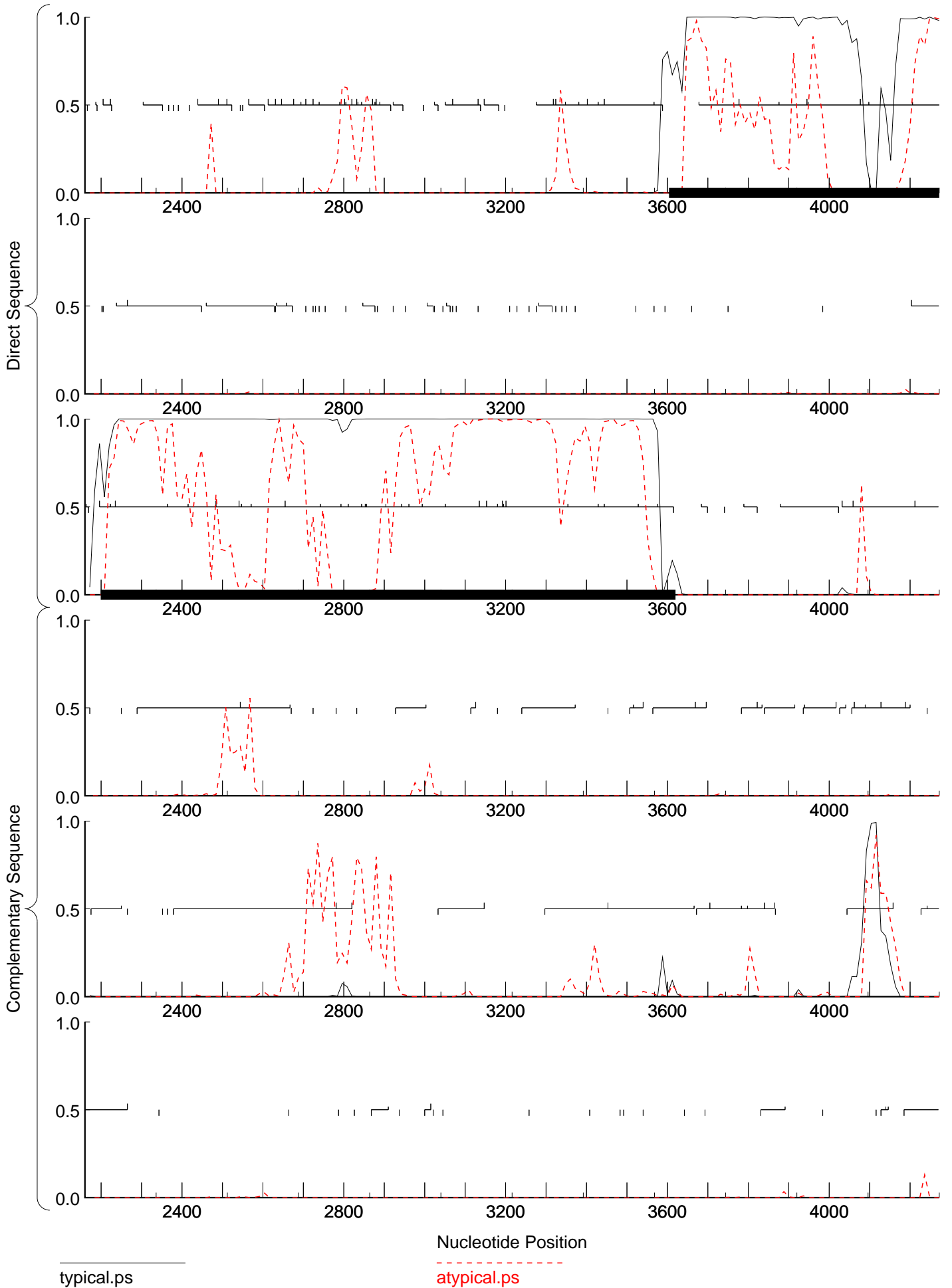
Matrix : -
Author : -
Order : 4

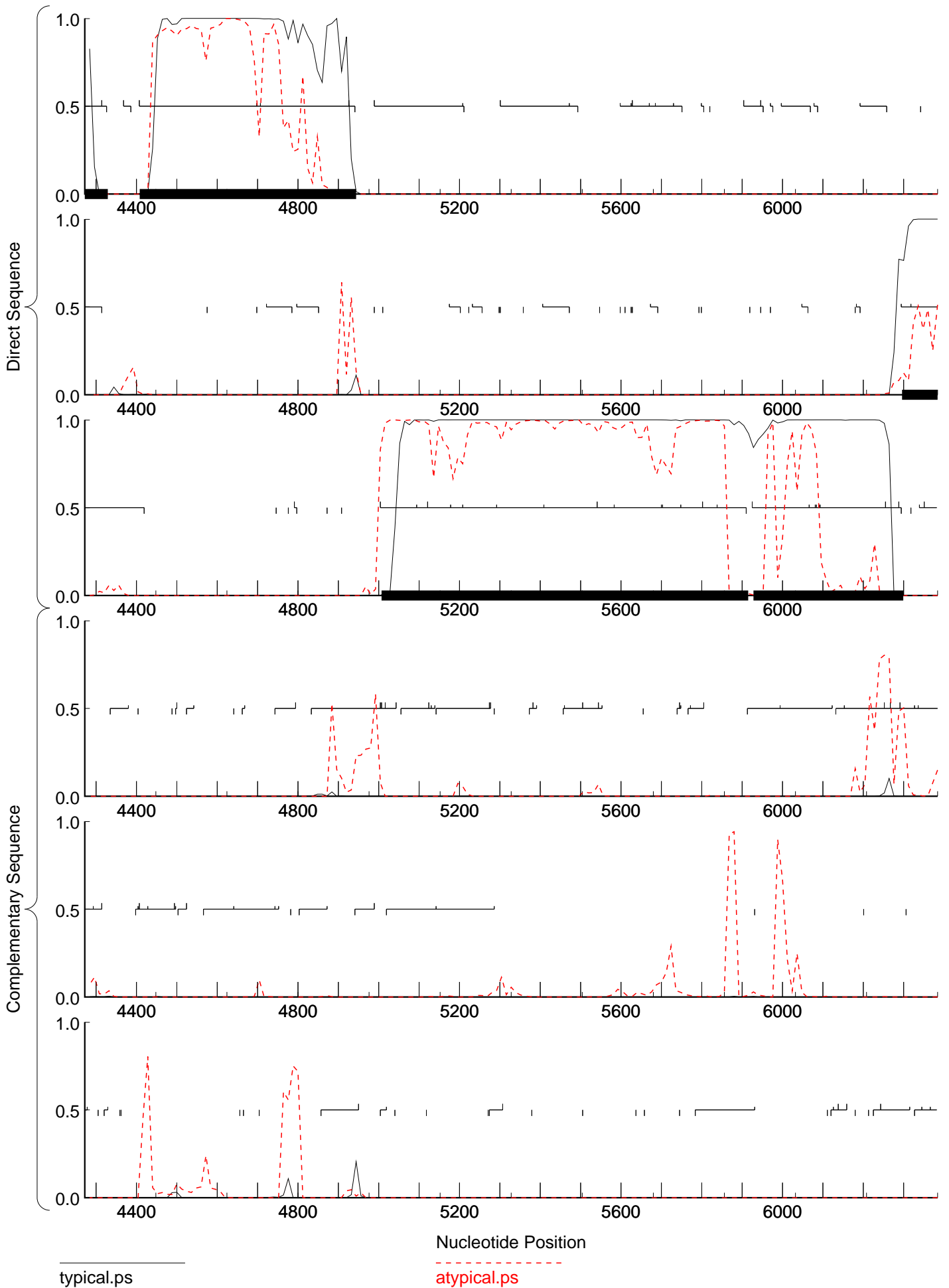
Send questions / comments to:
Dr. M. Borodovsky
Georgia Institute of Technology
School of Biology
Atlanta, GA 30332-0230

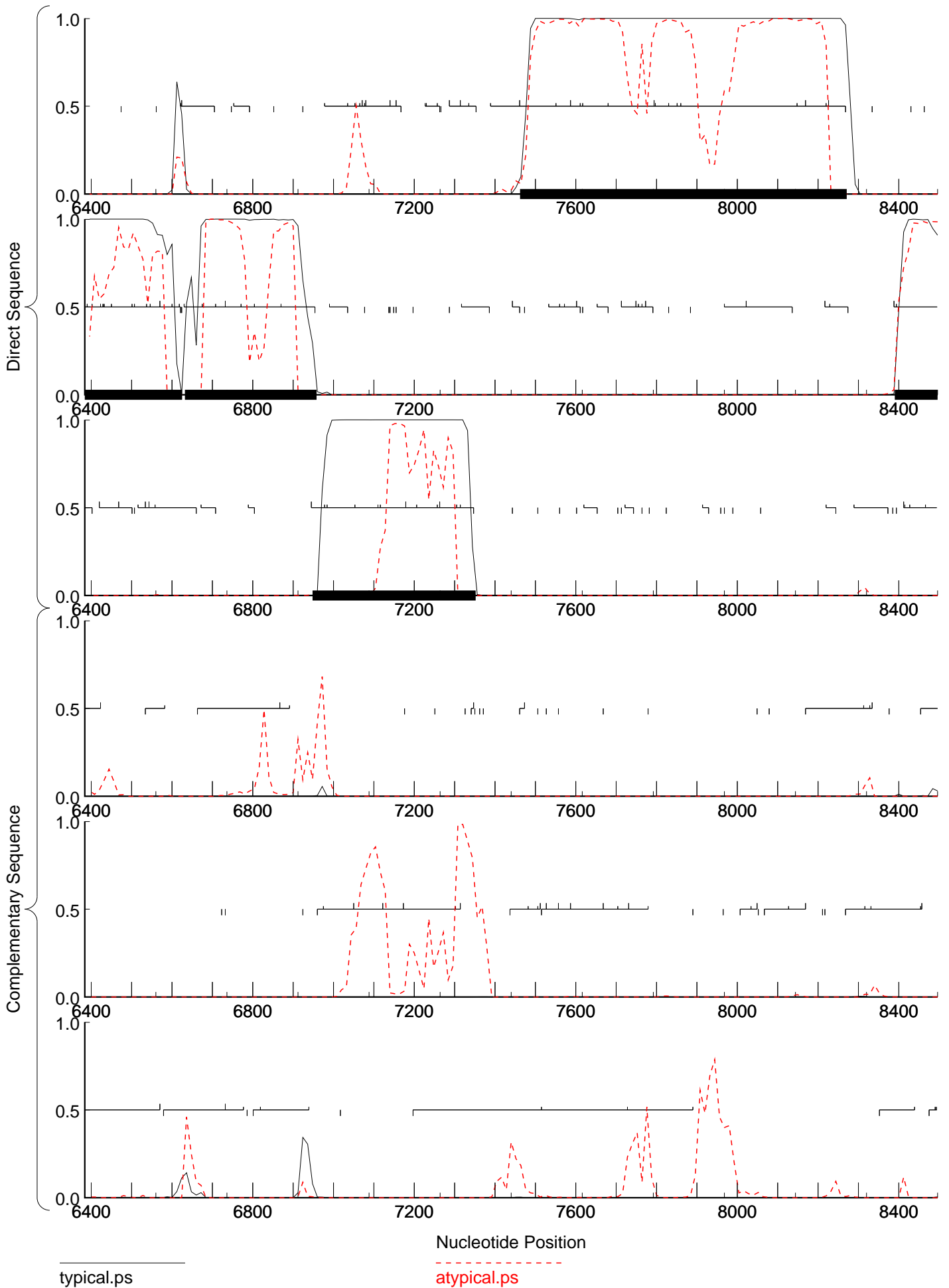
Matrix notes & comments

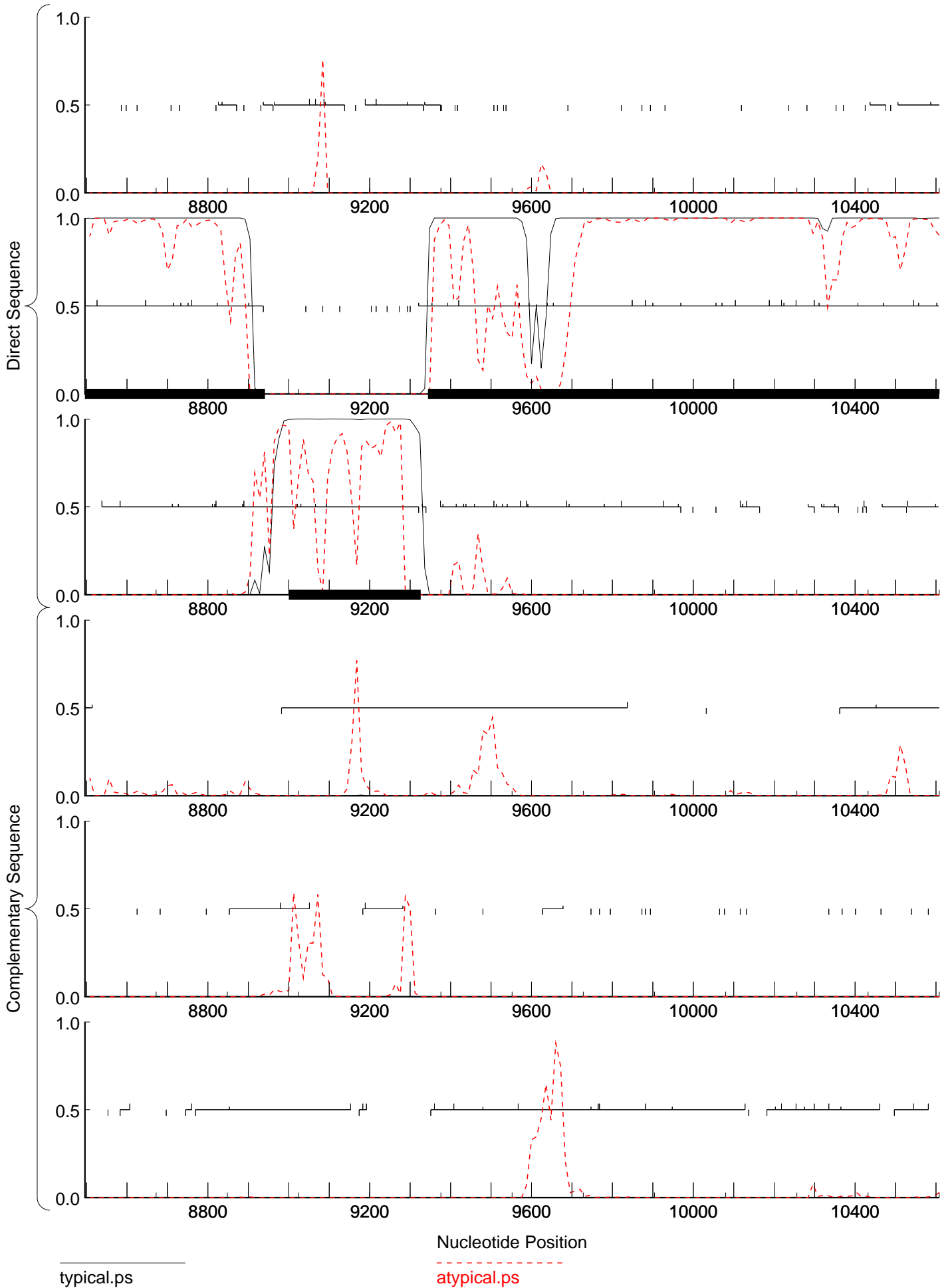
Training set derived by GeneMarkS, 4.30 October 2014
Sat May 26 07:53:23 2018

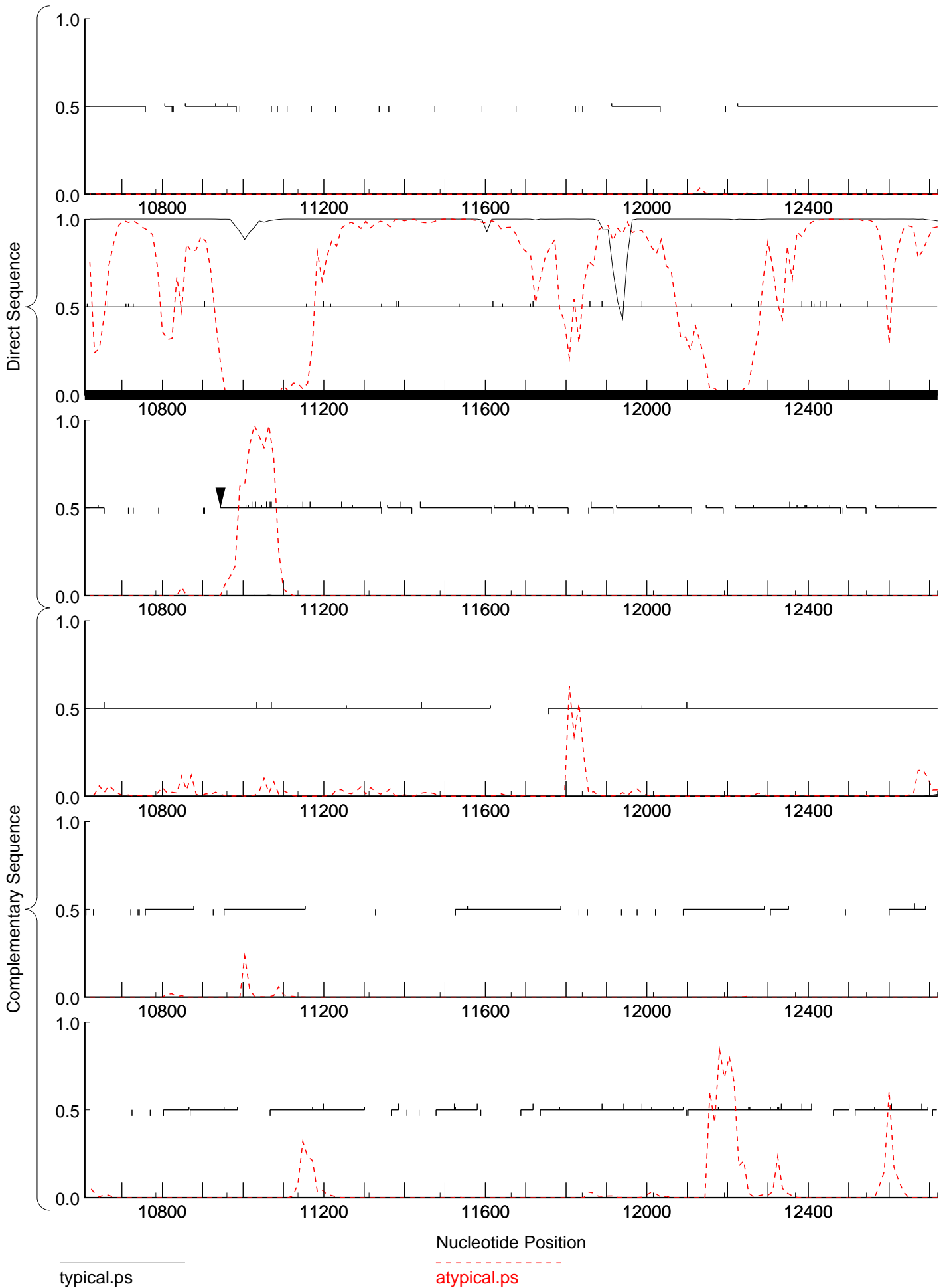


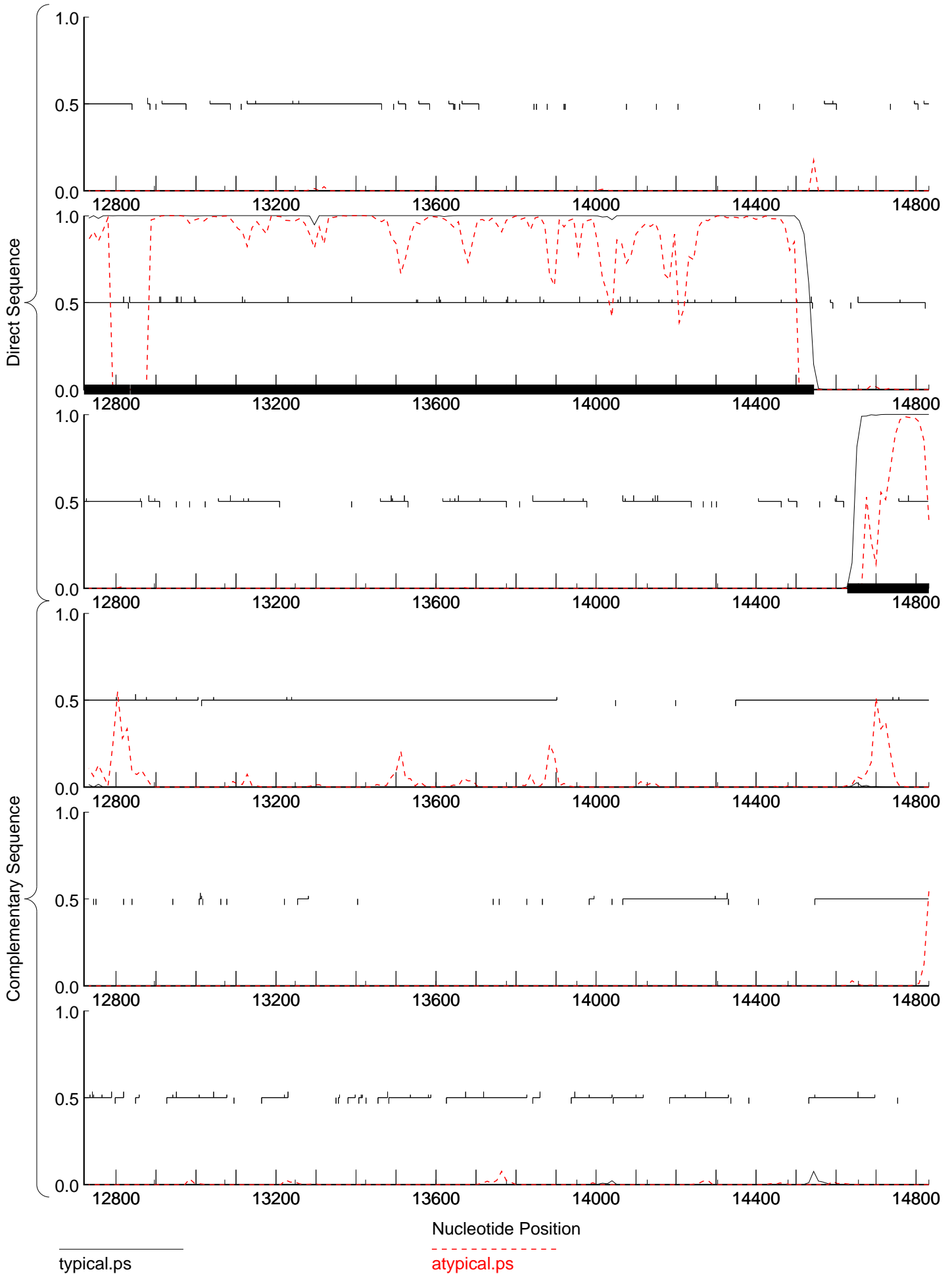


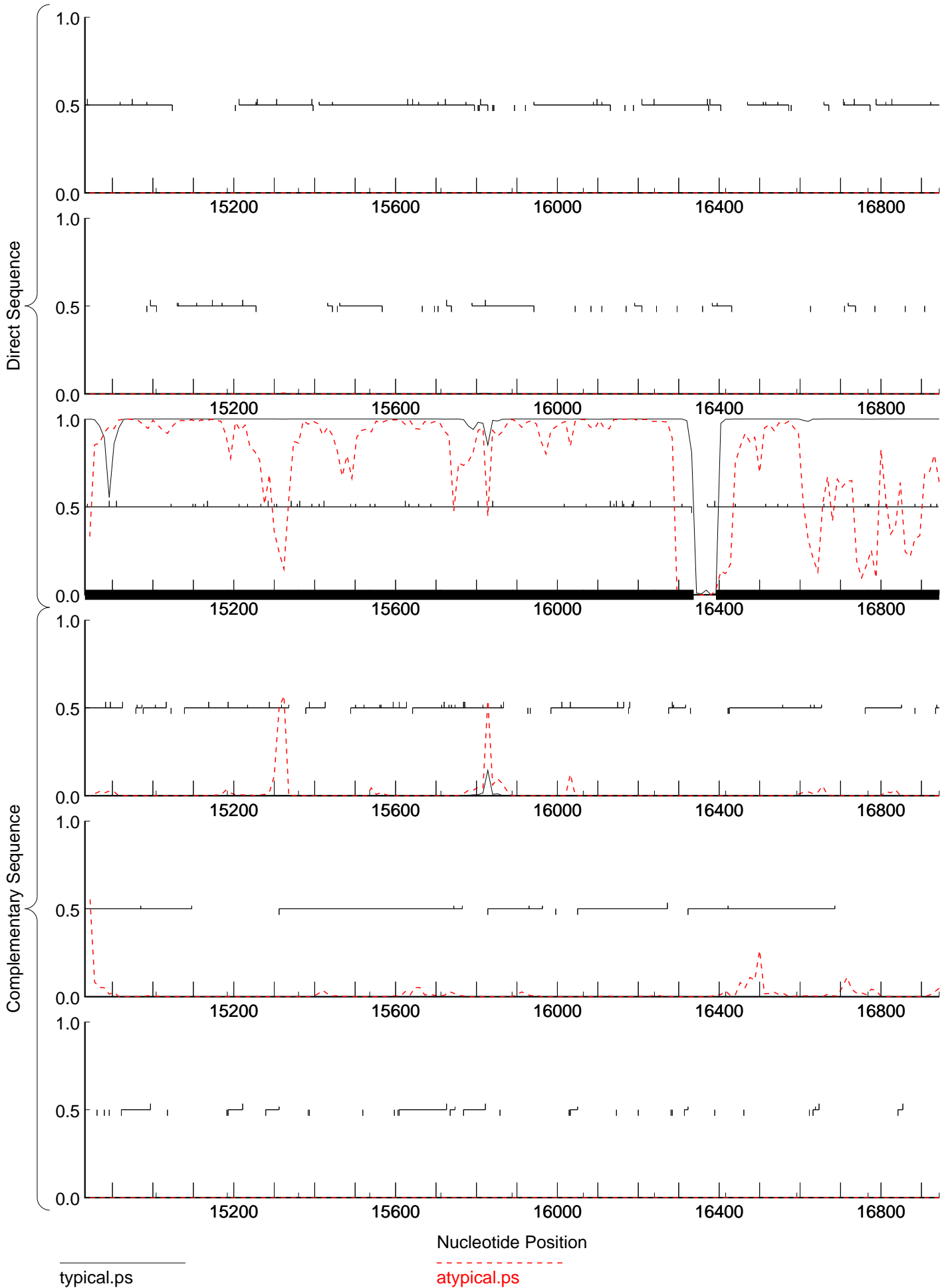


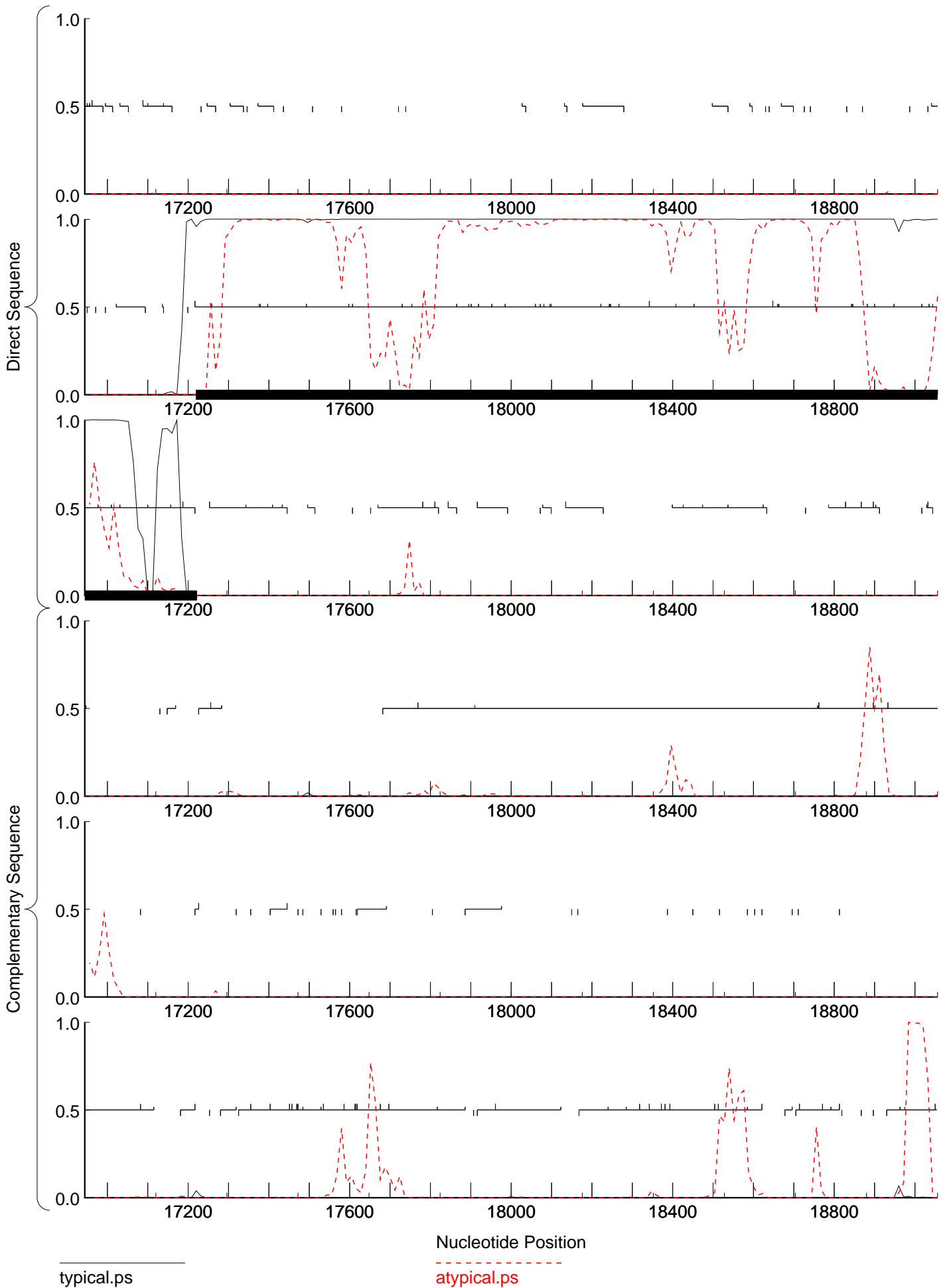


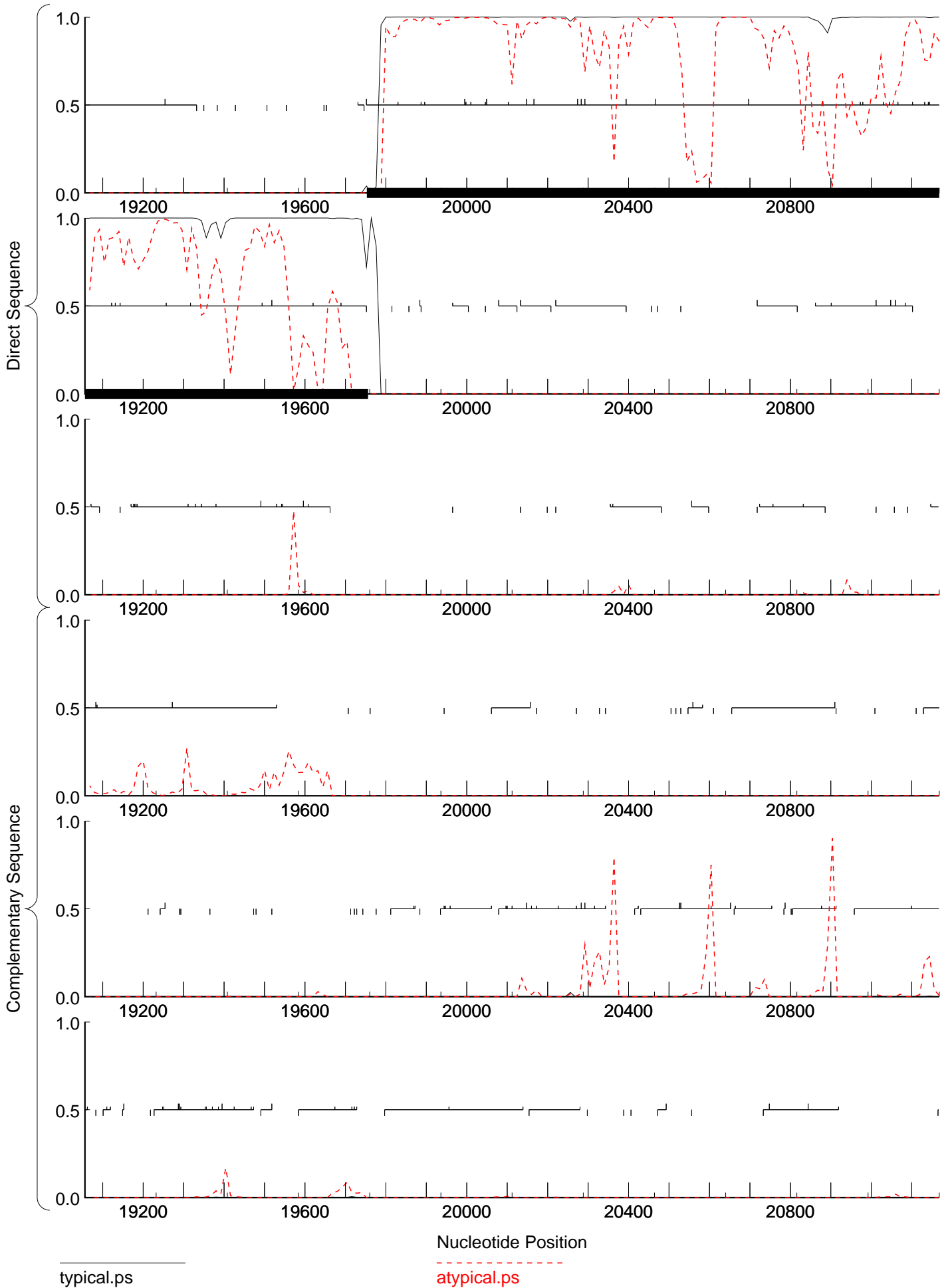


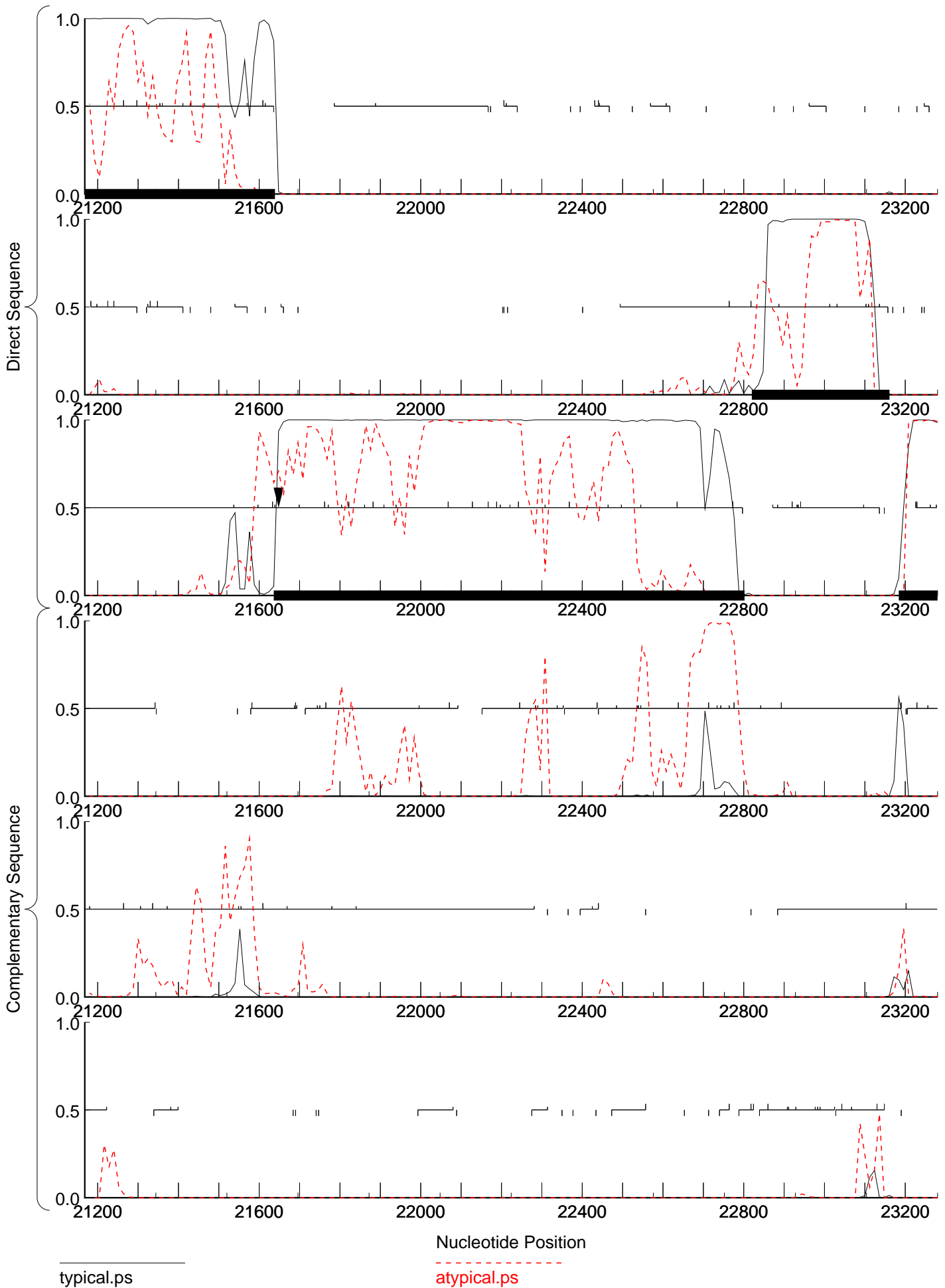


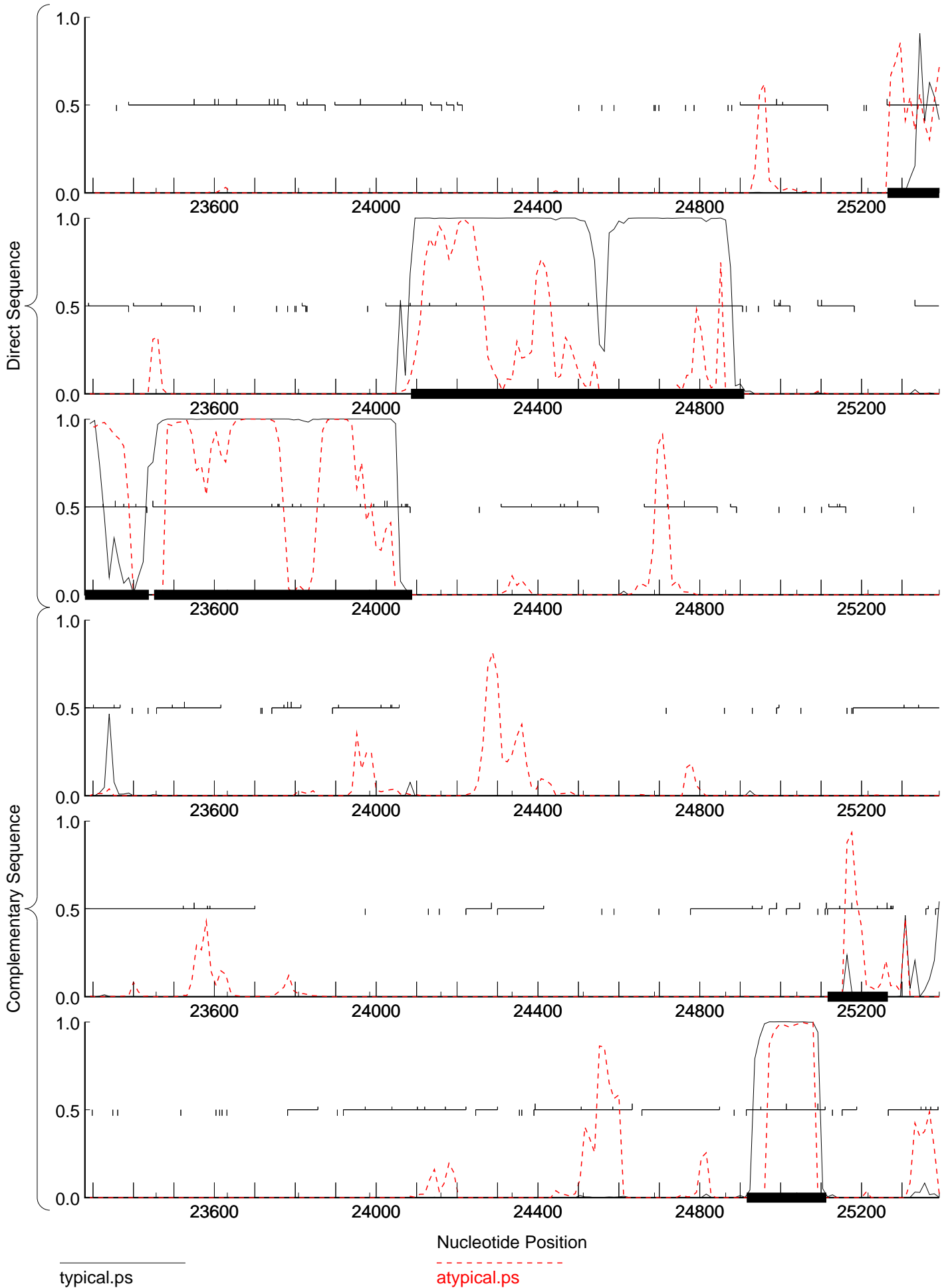


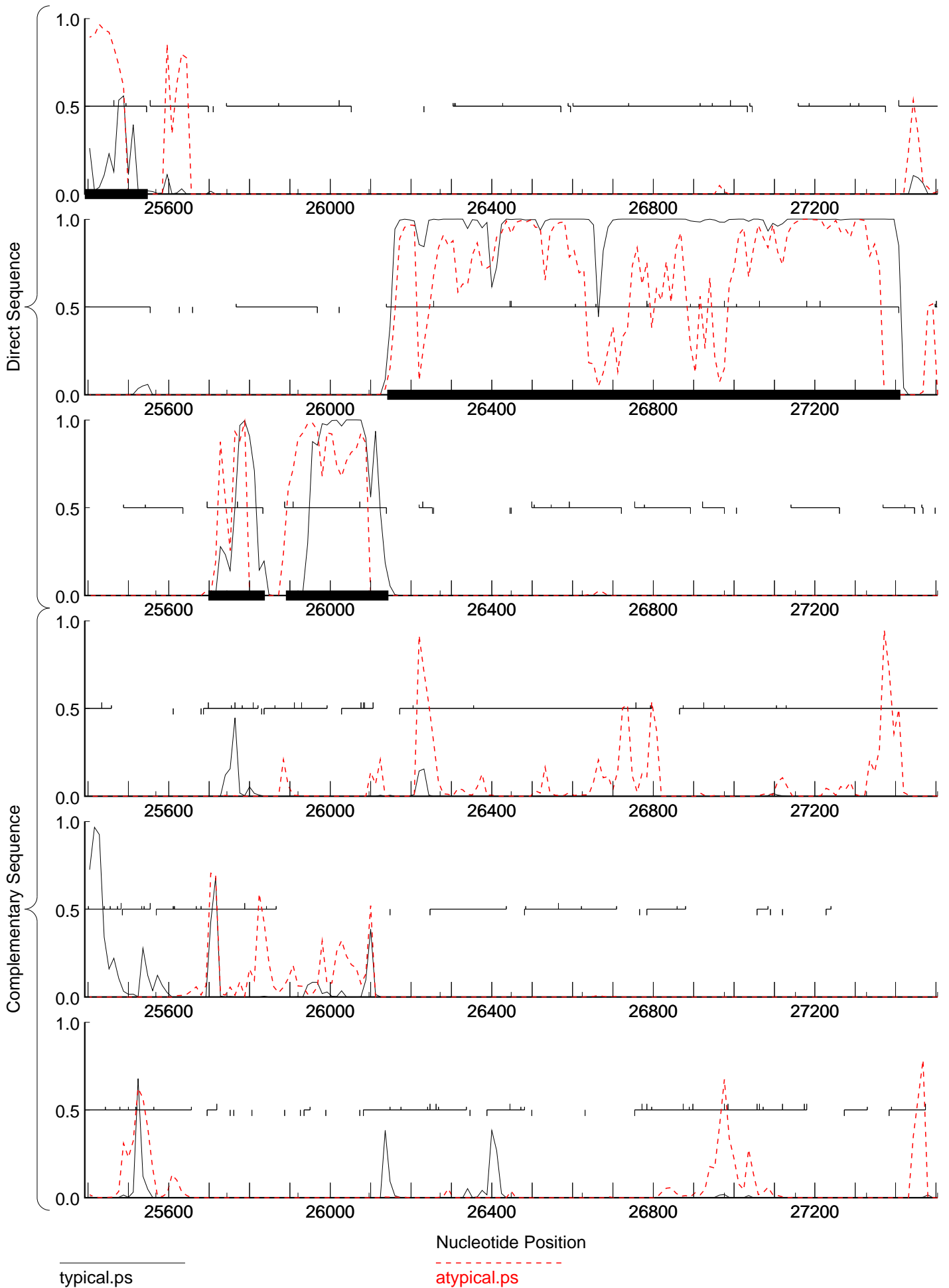


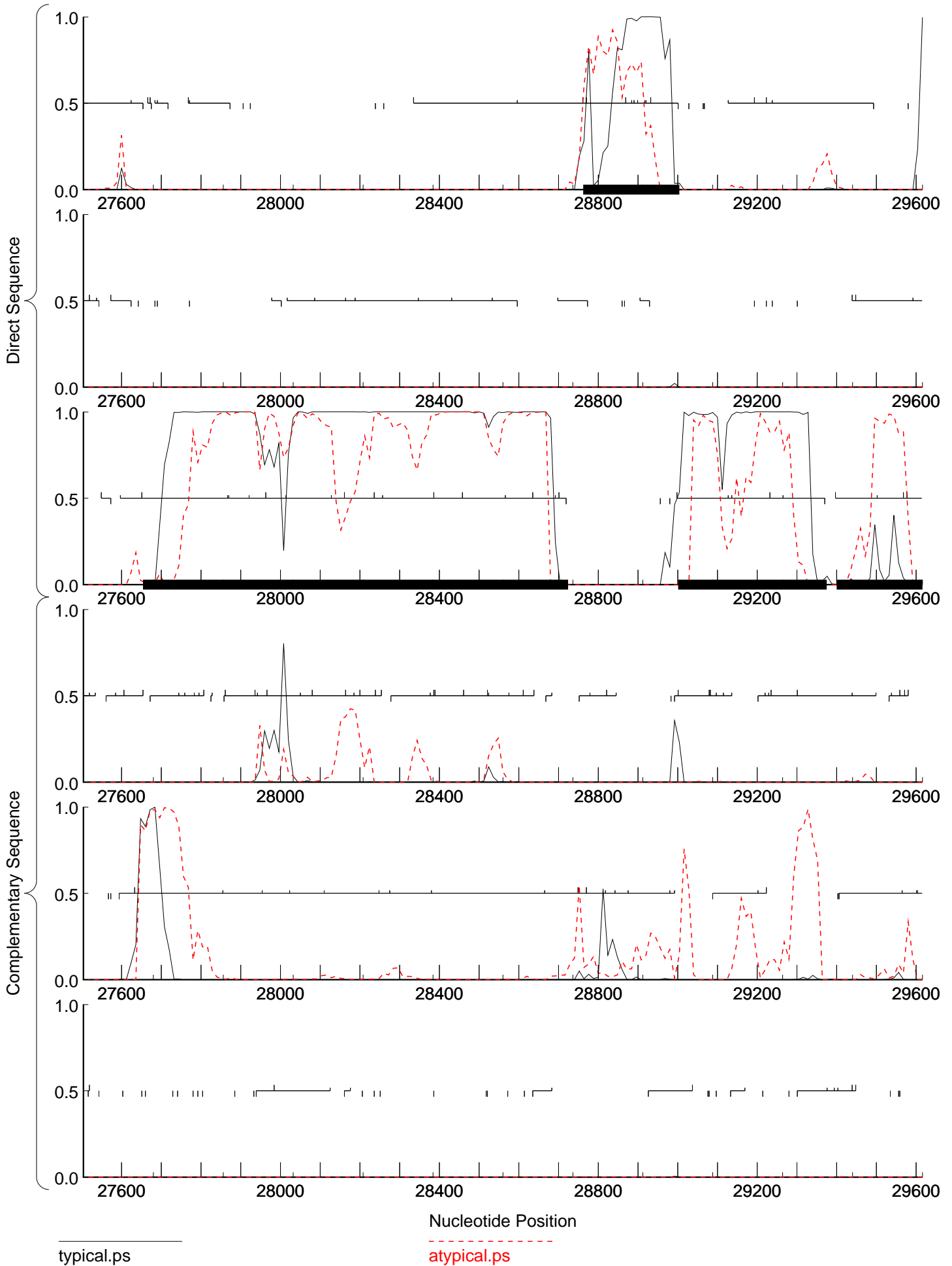






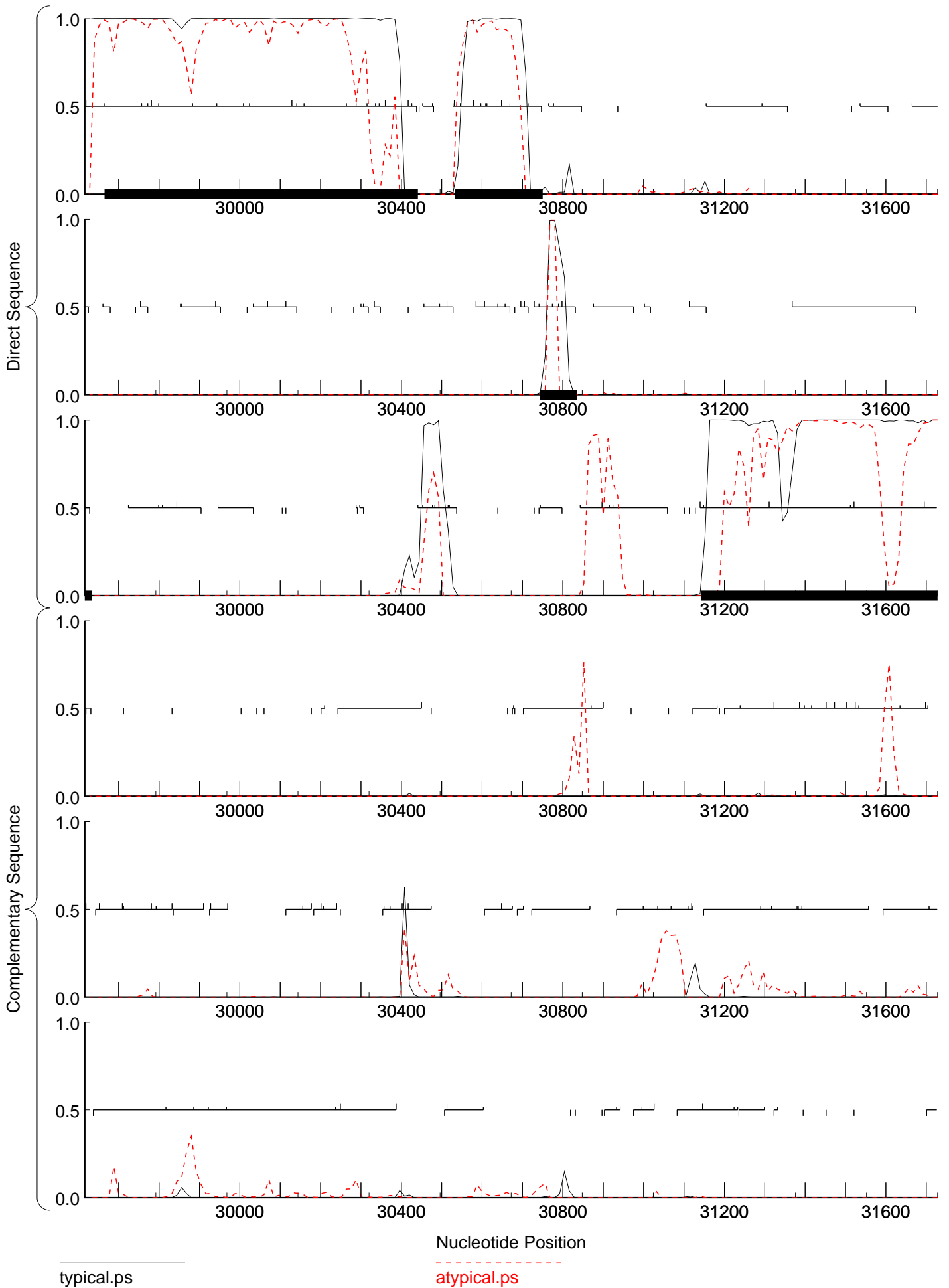


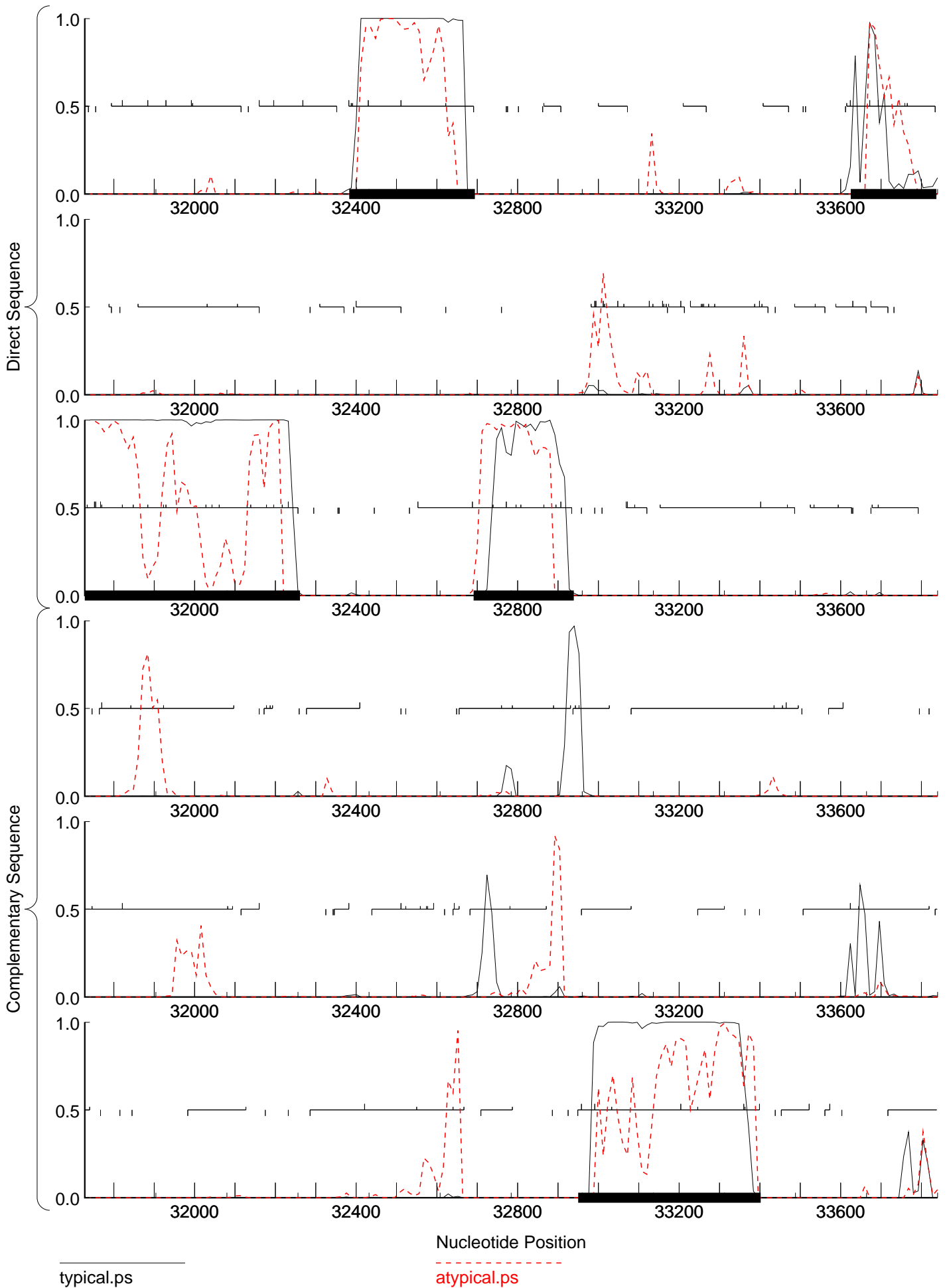


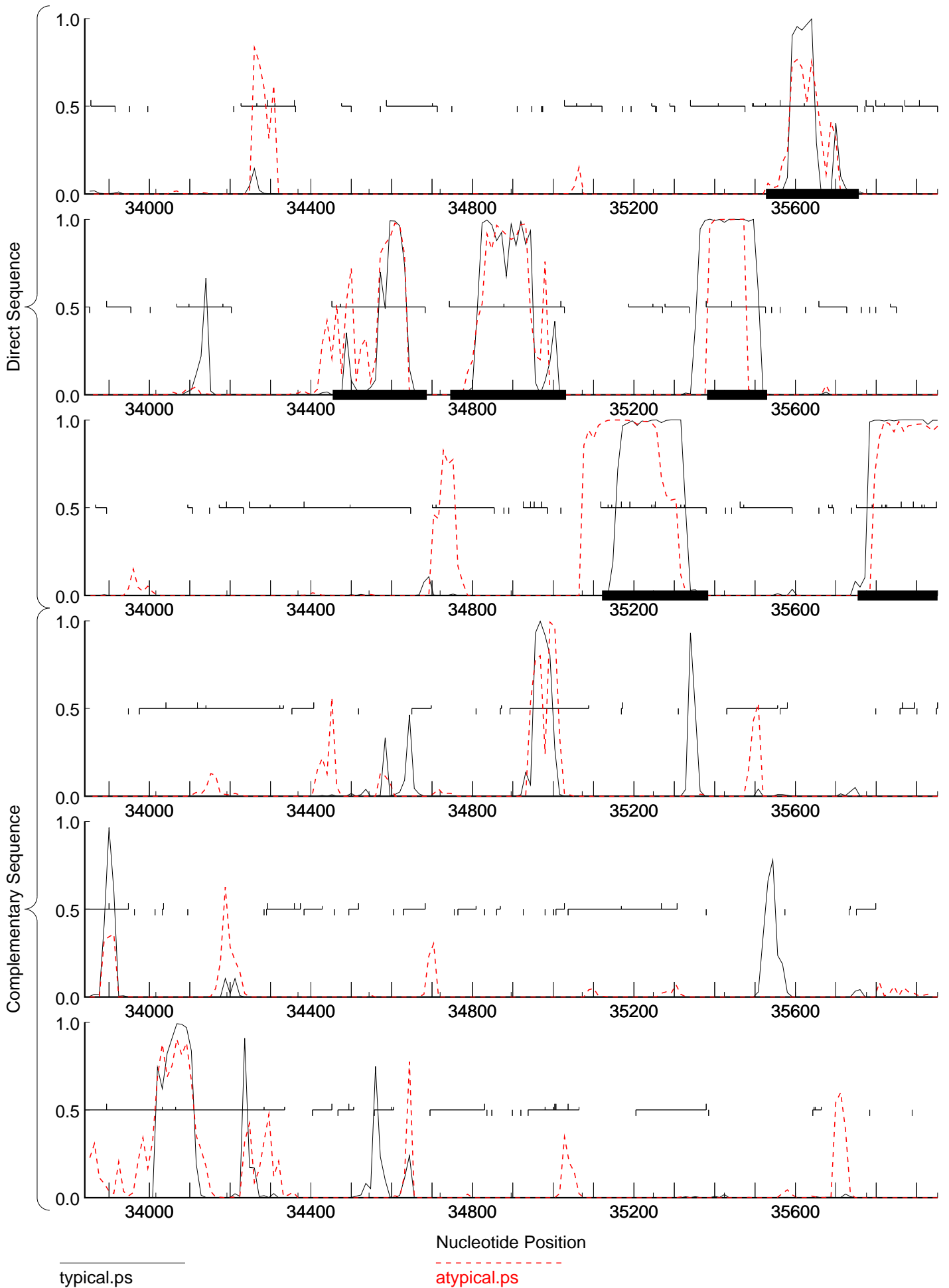


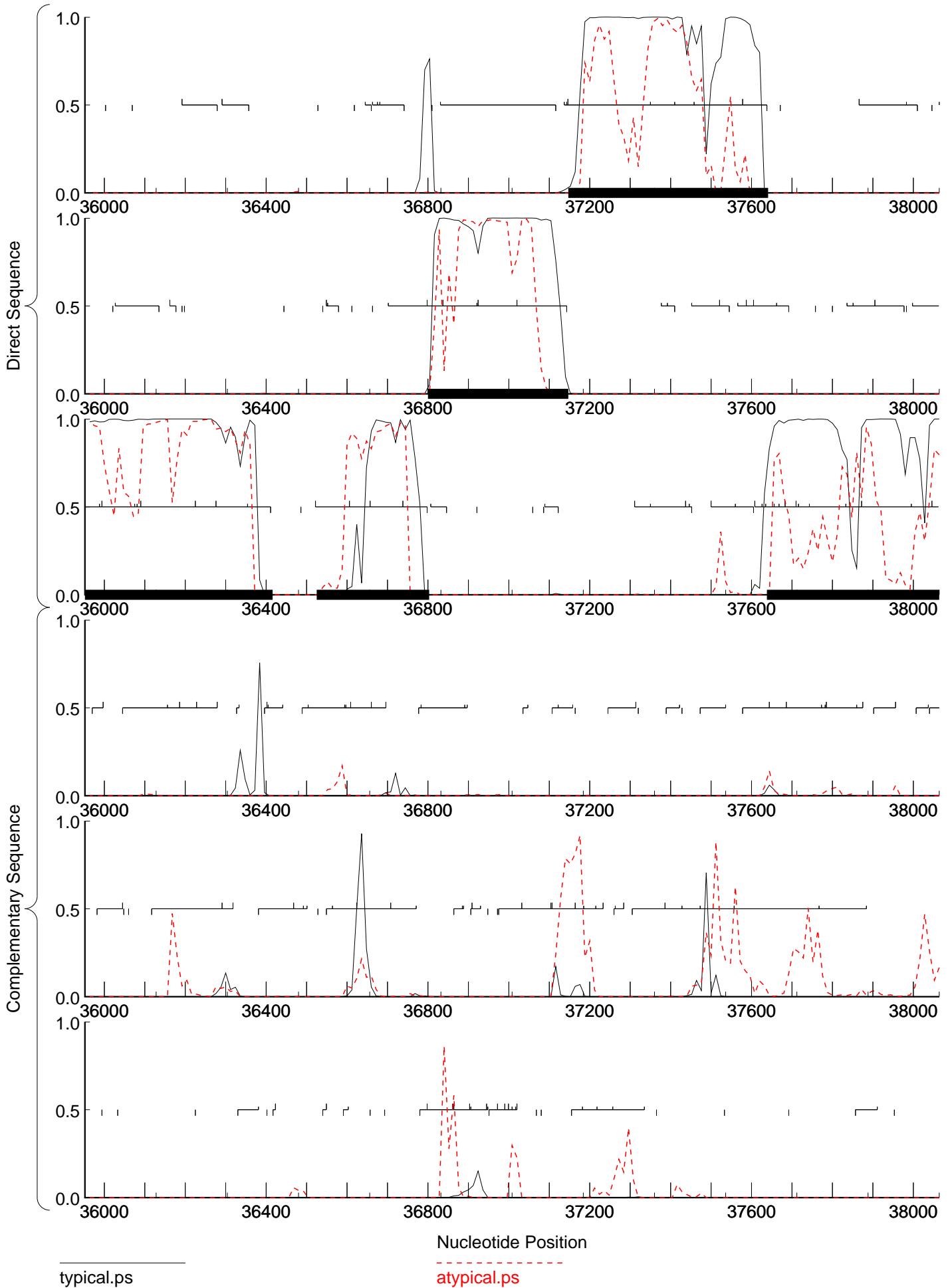
typical.ps

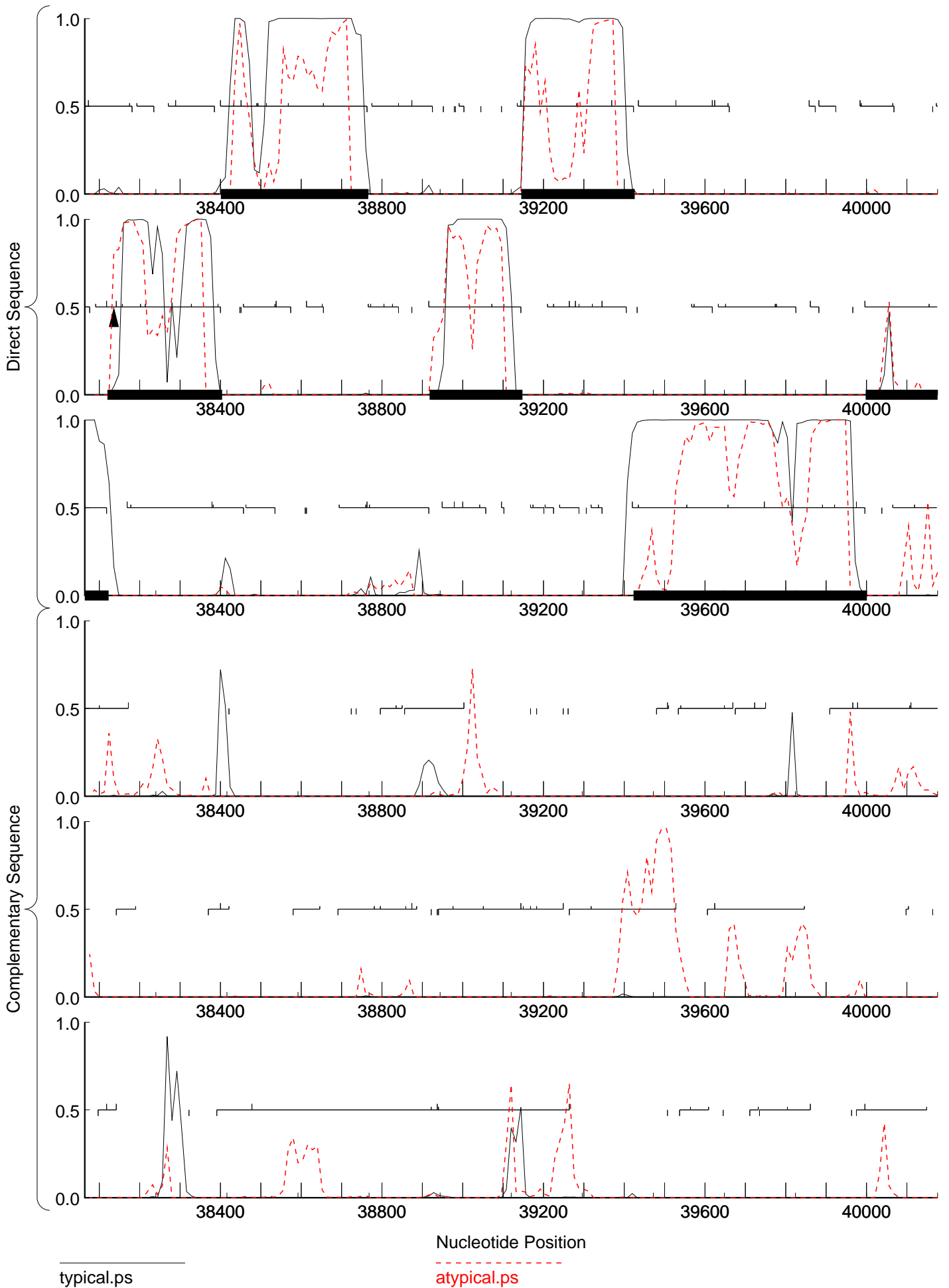
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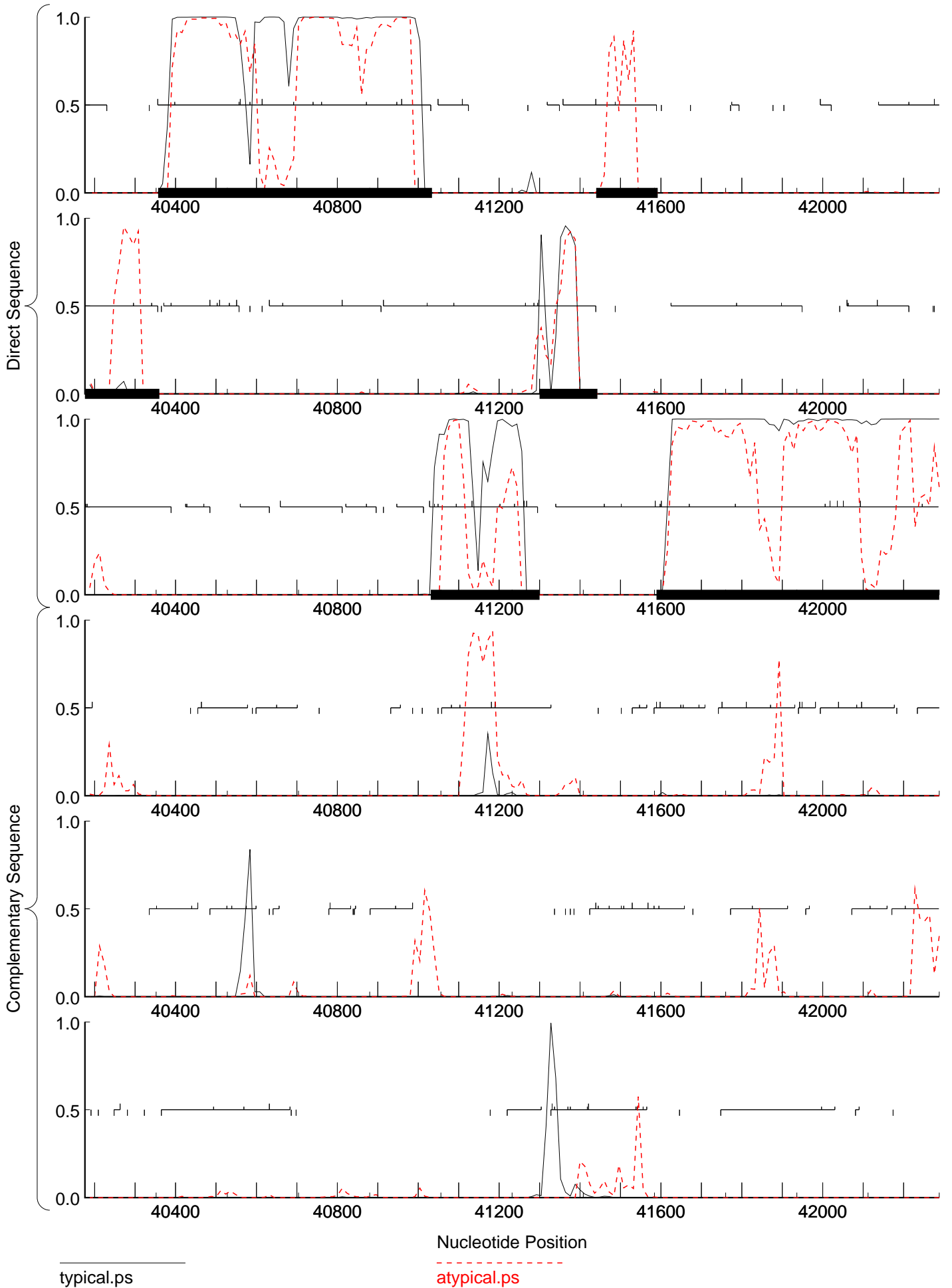


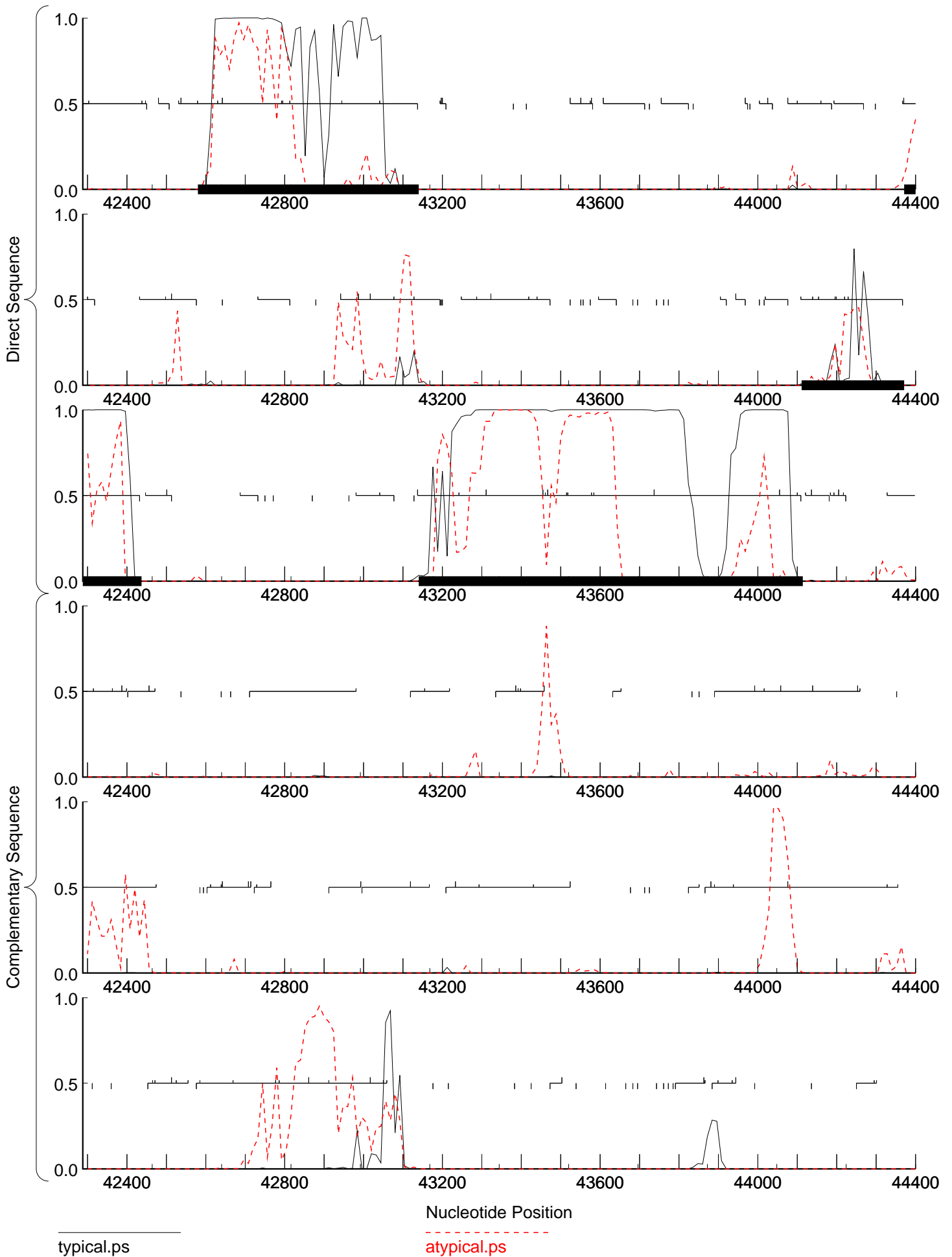


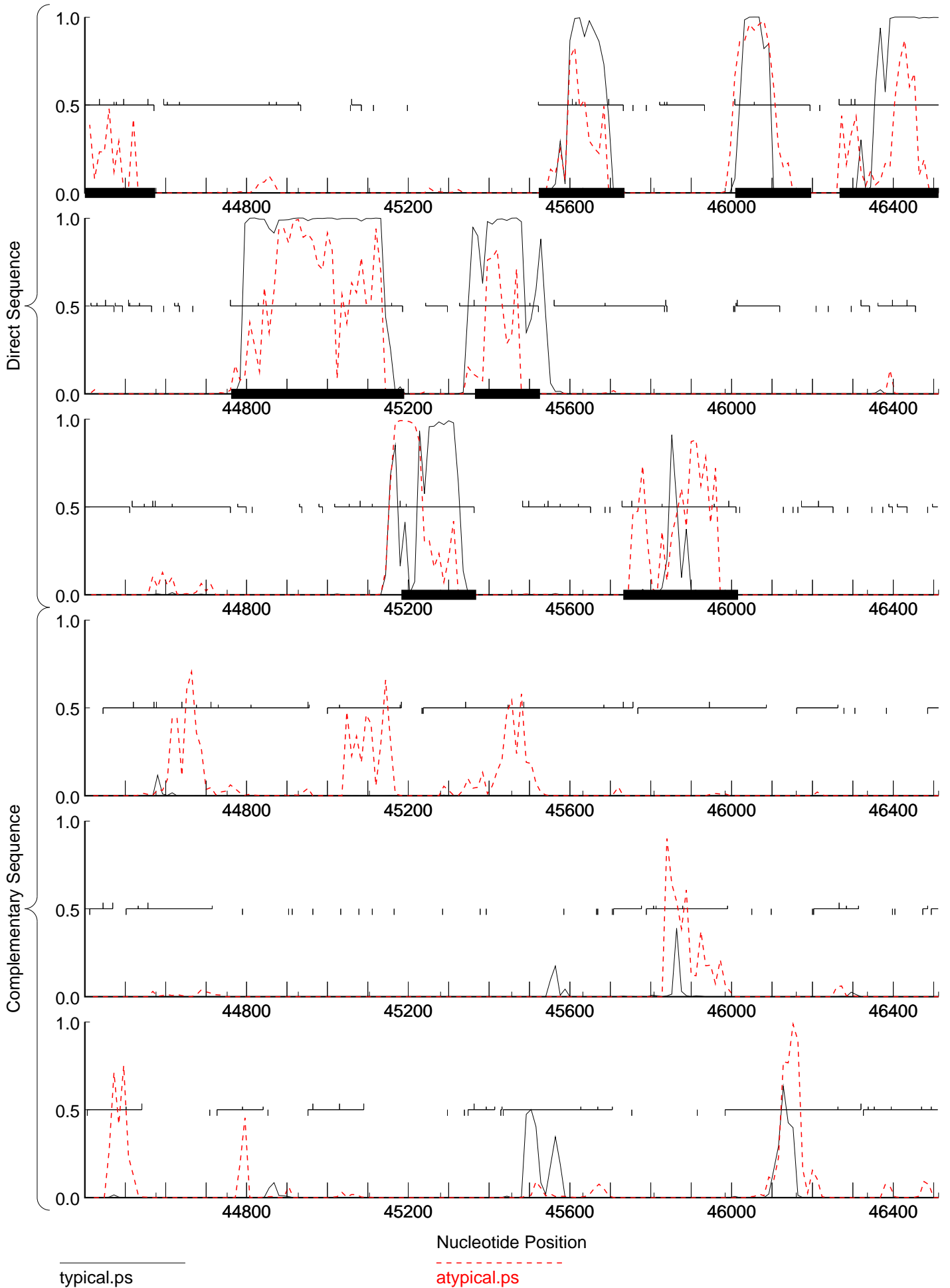


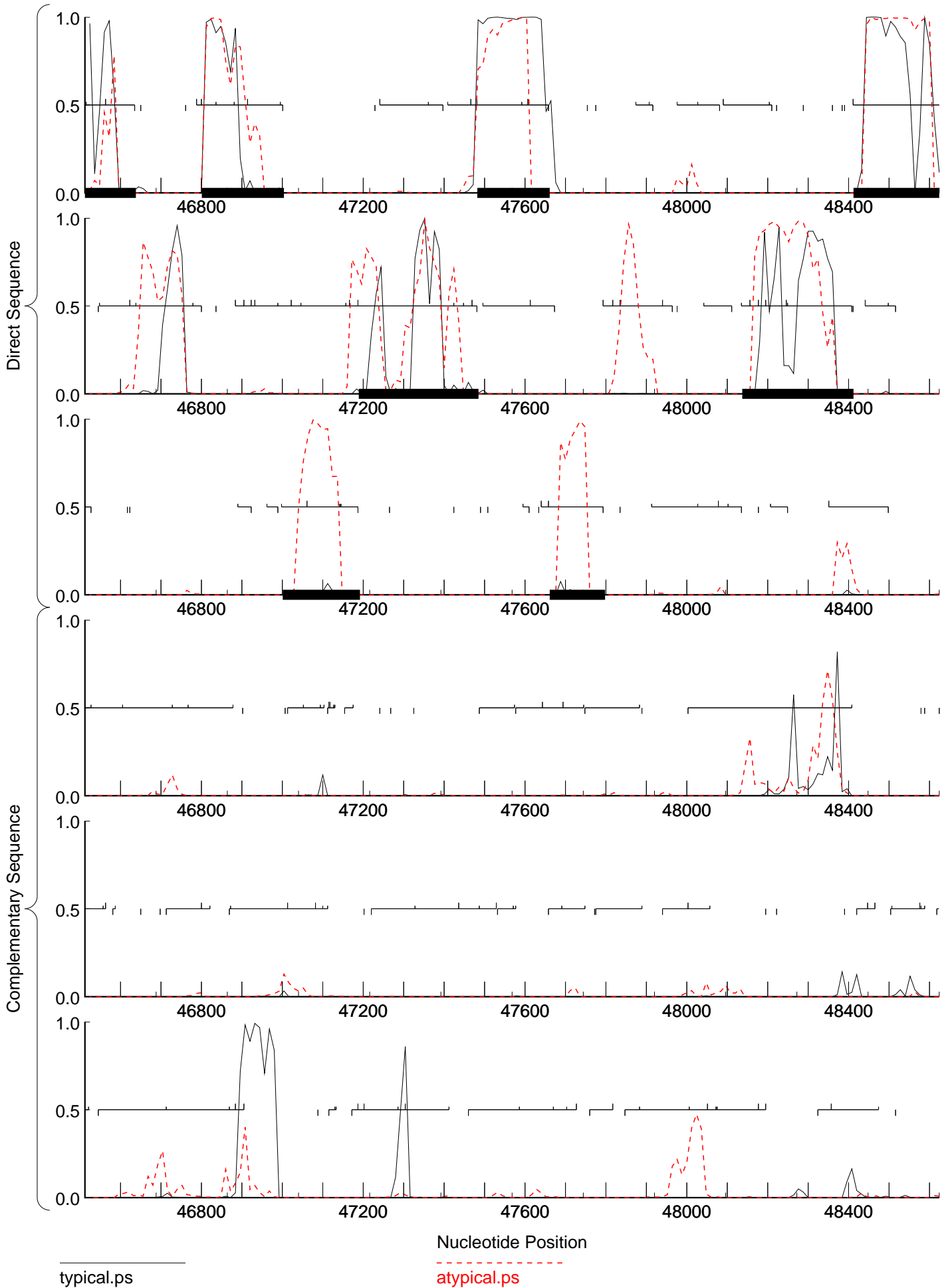


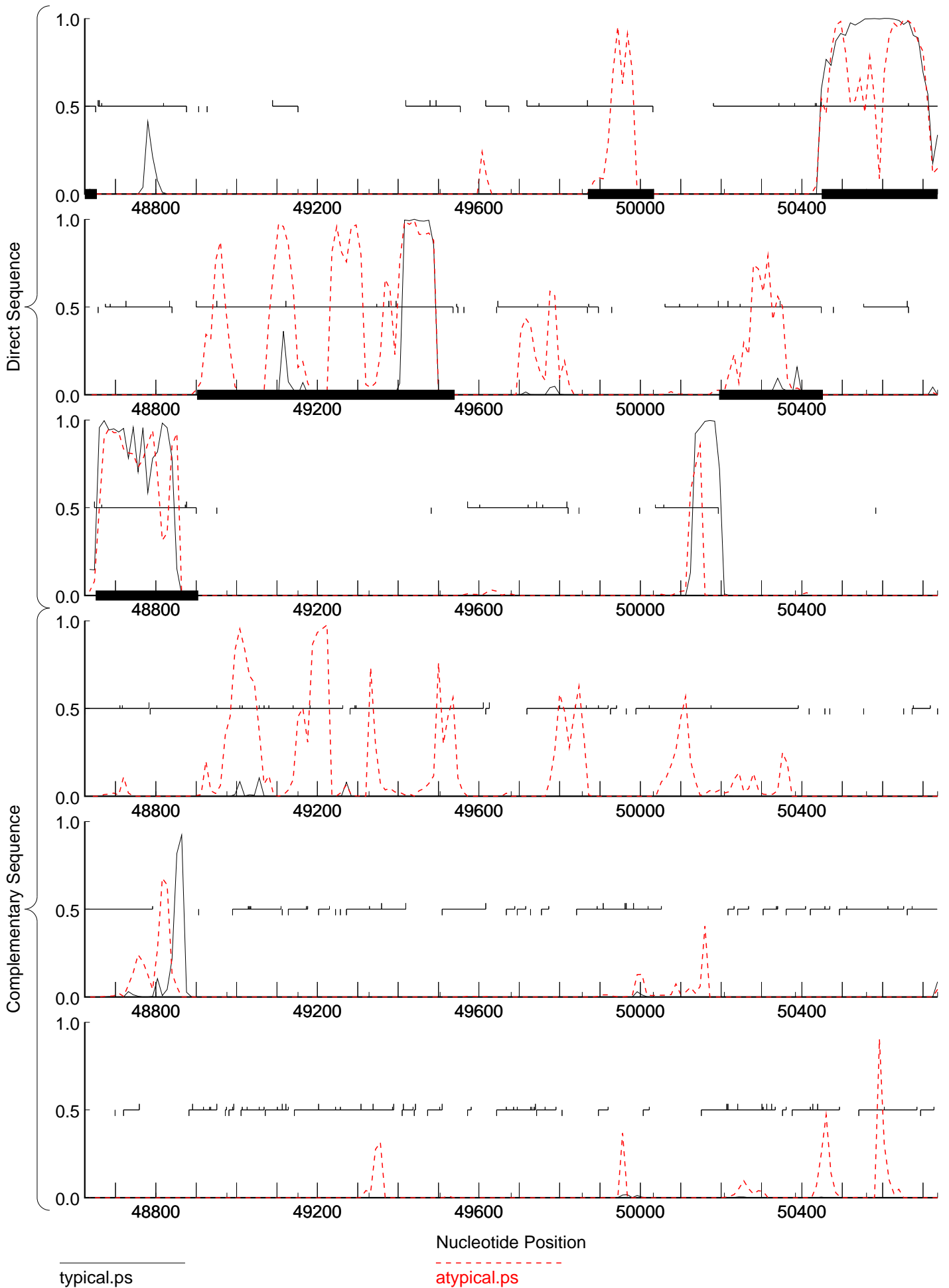


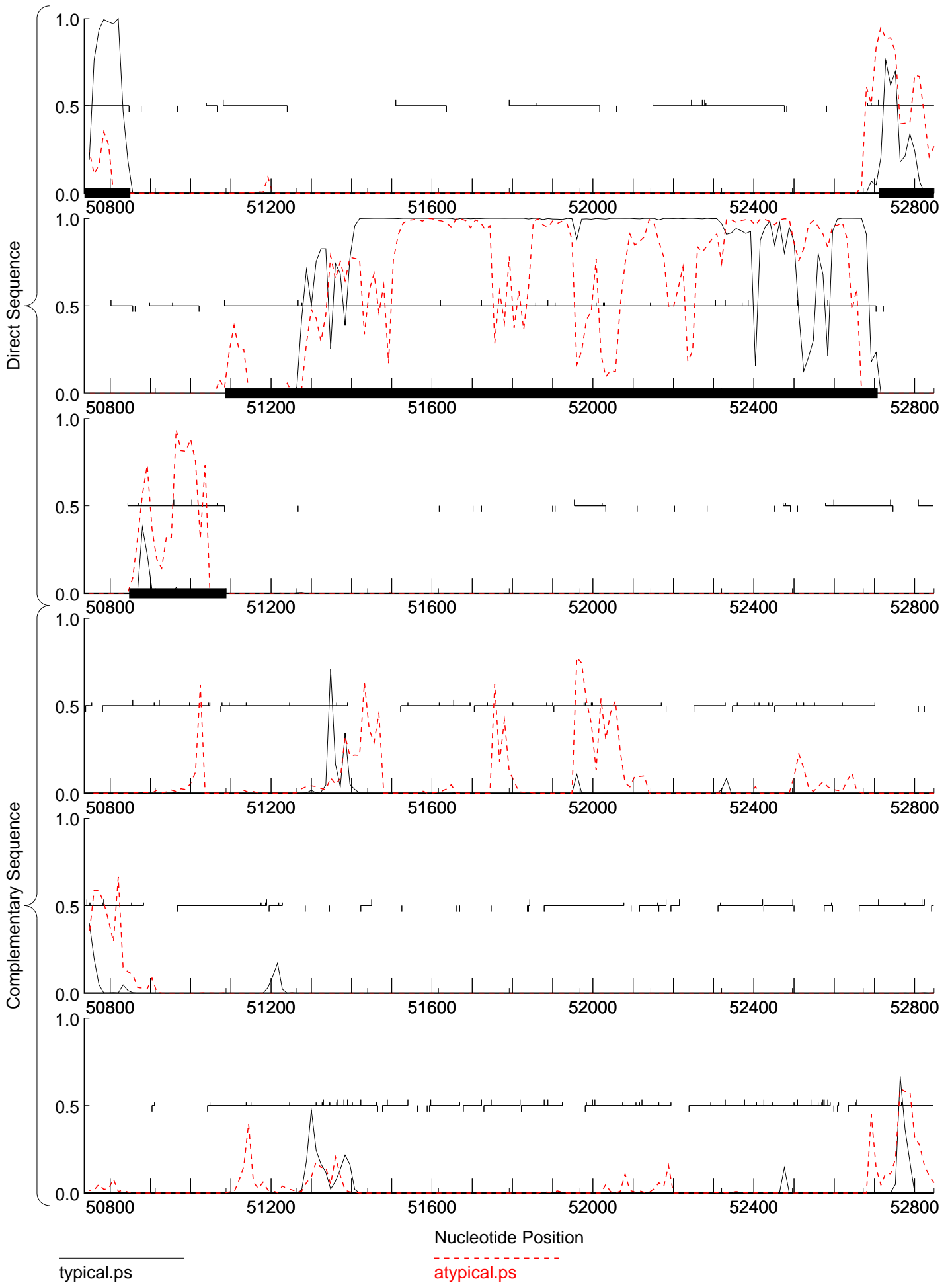












typical.ps

atypical.ps

