

GeneMark

Version 2.5p (09.08.06)

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PROGRAM INFORMATION

Sequence : Mycobacterium phage Ein37 complete sequence, 53748 bp including 14-base 3' overhang (TAAGCC
Analysis Date : 8/19/17 at 5:19:46
Pages : 27
Sequence Length : 53748 bp
GC Content : 67.45%

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,

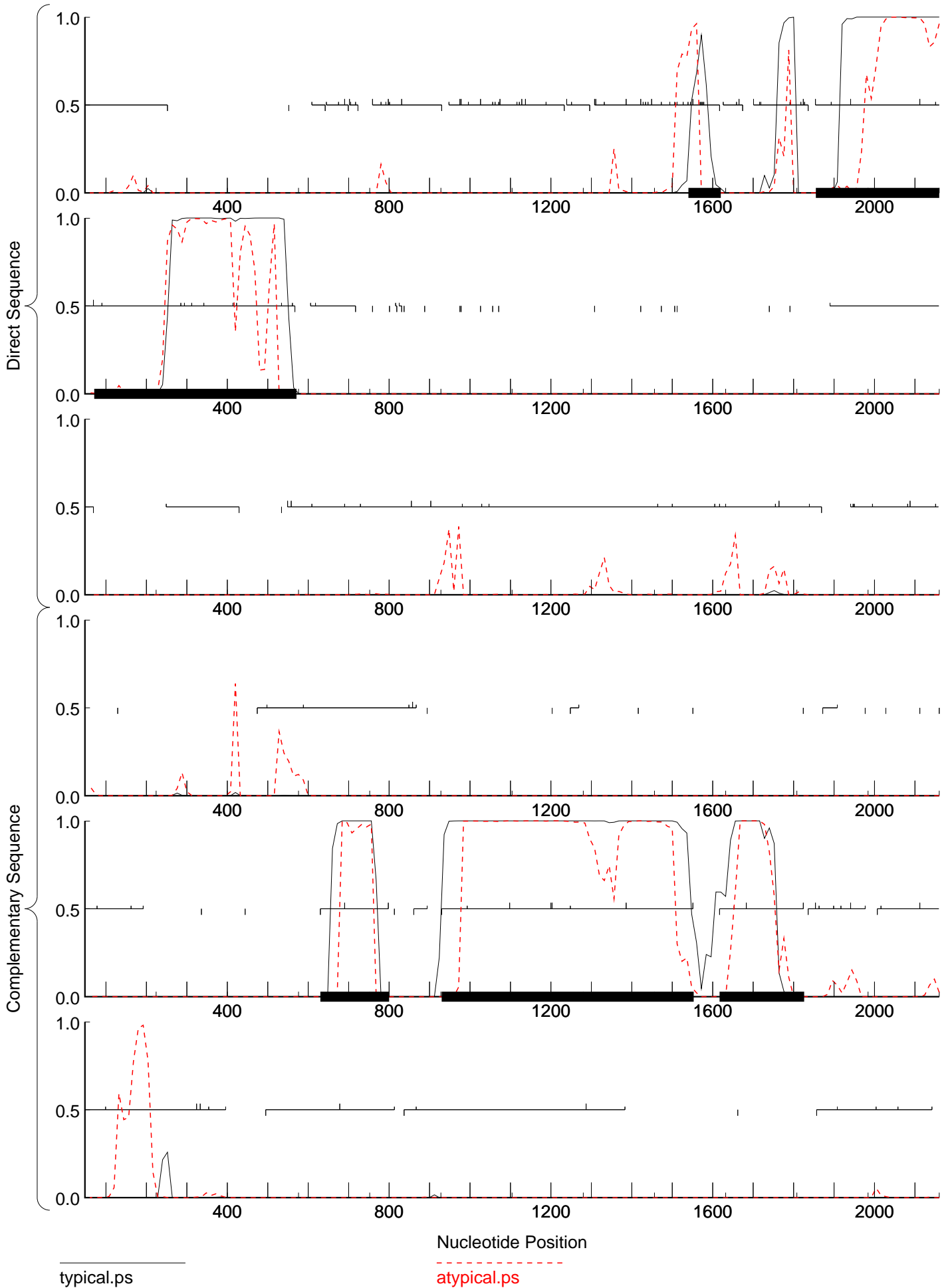
Matrix notes & comments

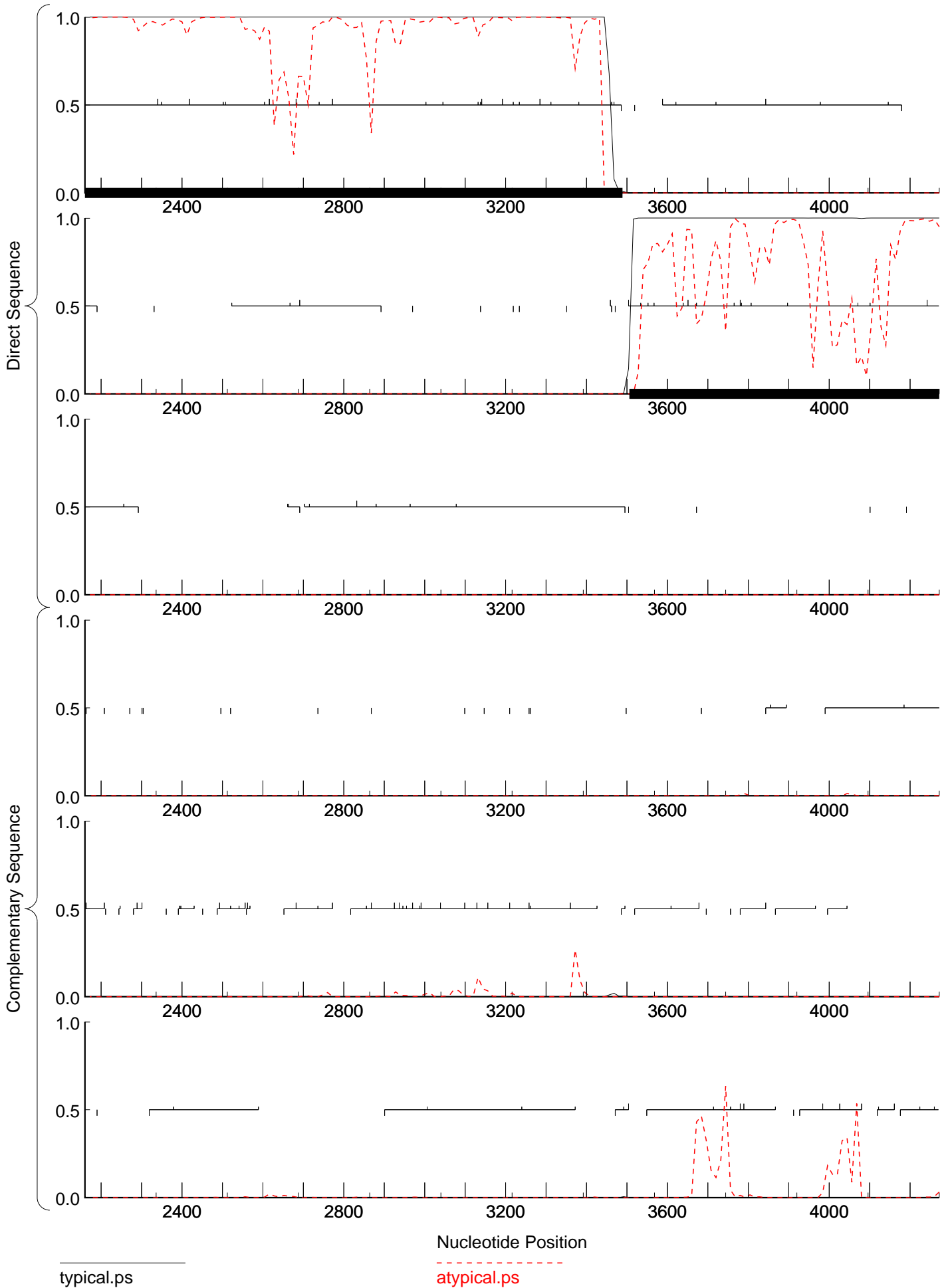
Training set derived by GeneMarkS, 4.30 October 2014
Sat Aug 19 05:19:45 2017

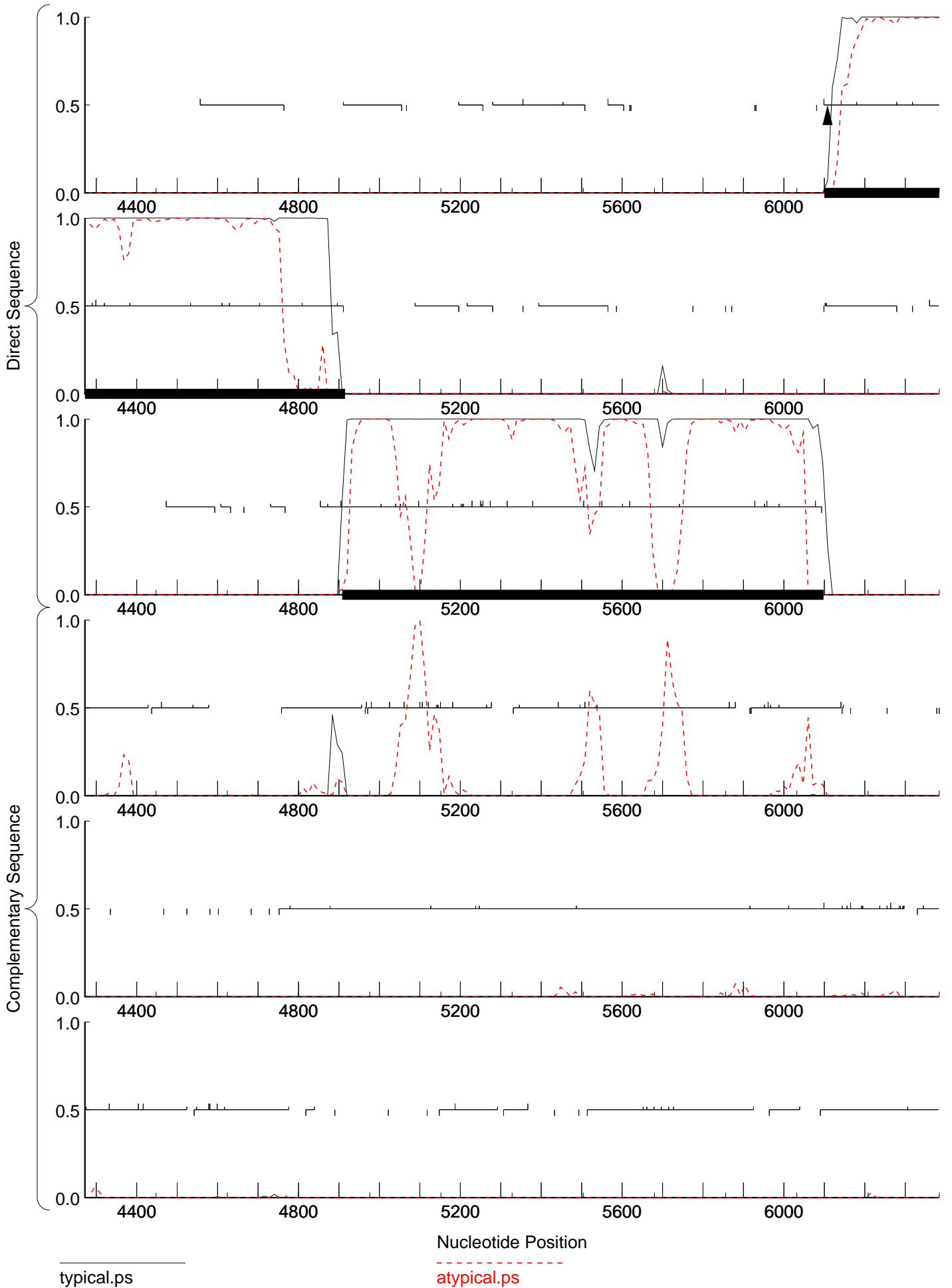
MATRIX INFORMATION

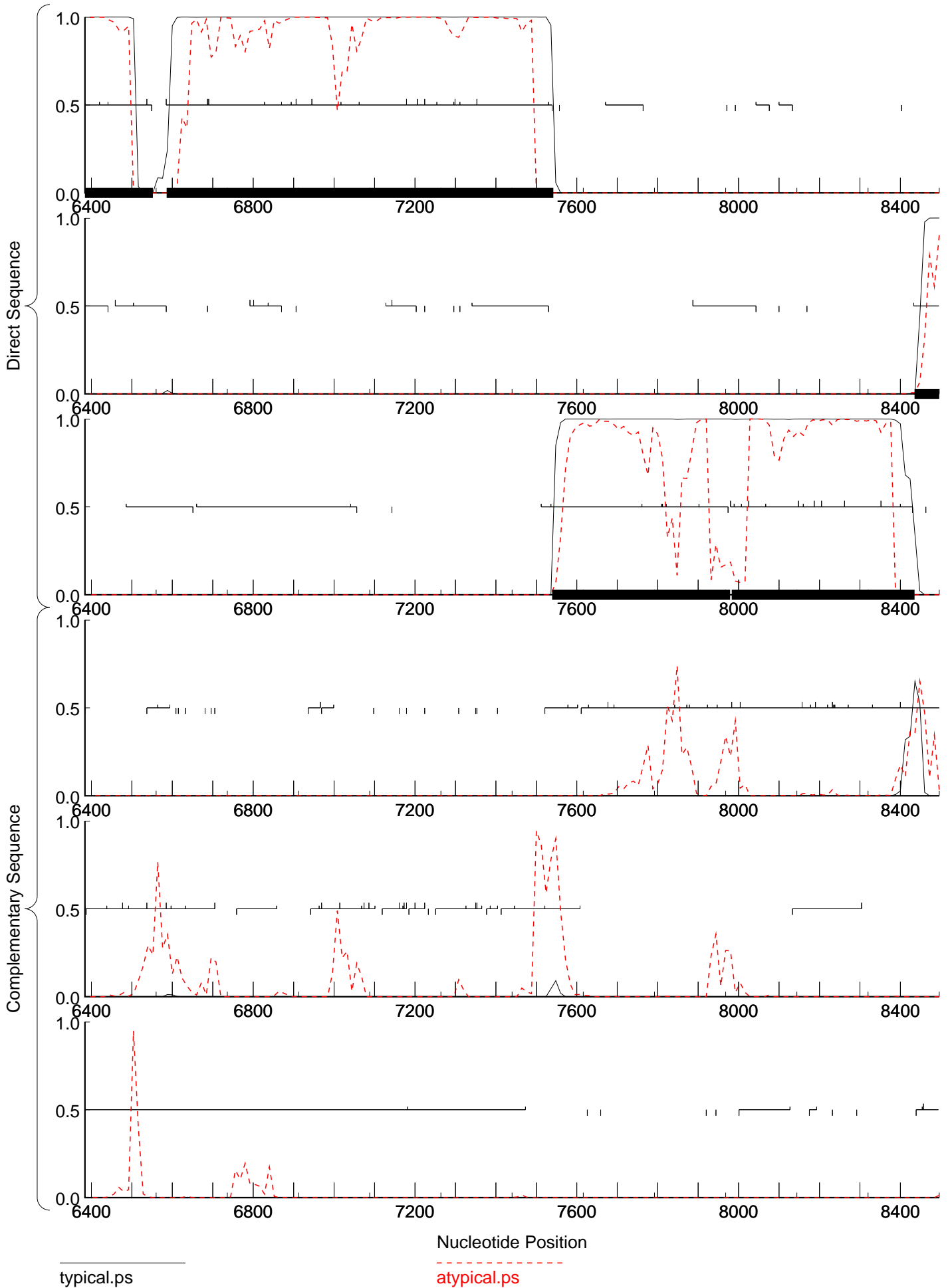
Matrix : -
Author : -
Order : 4

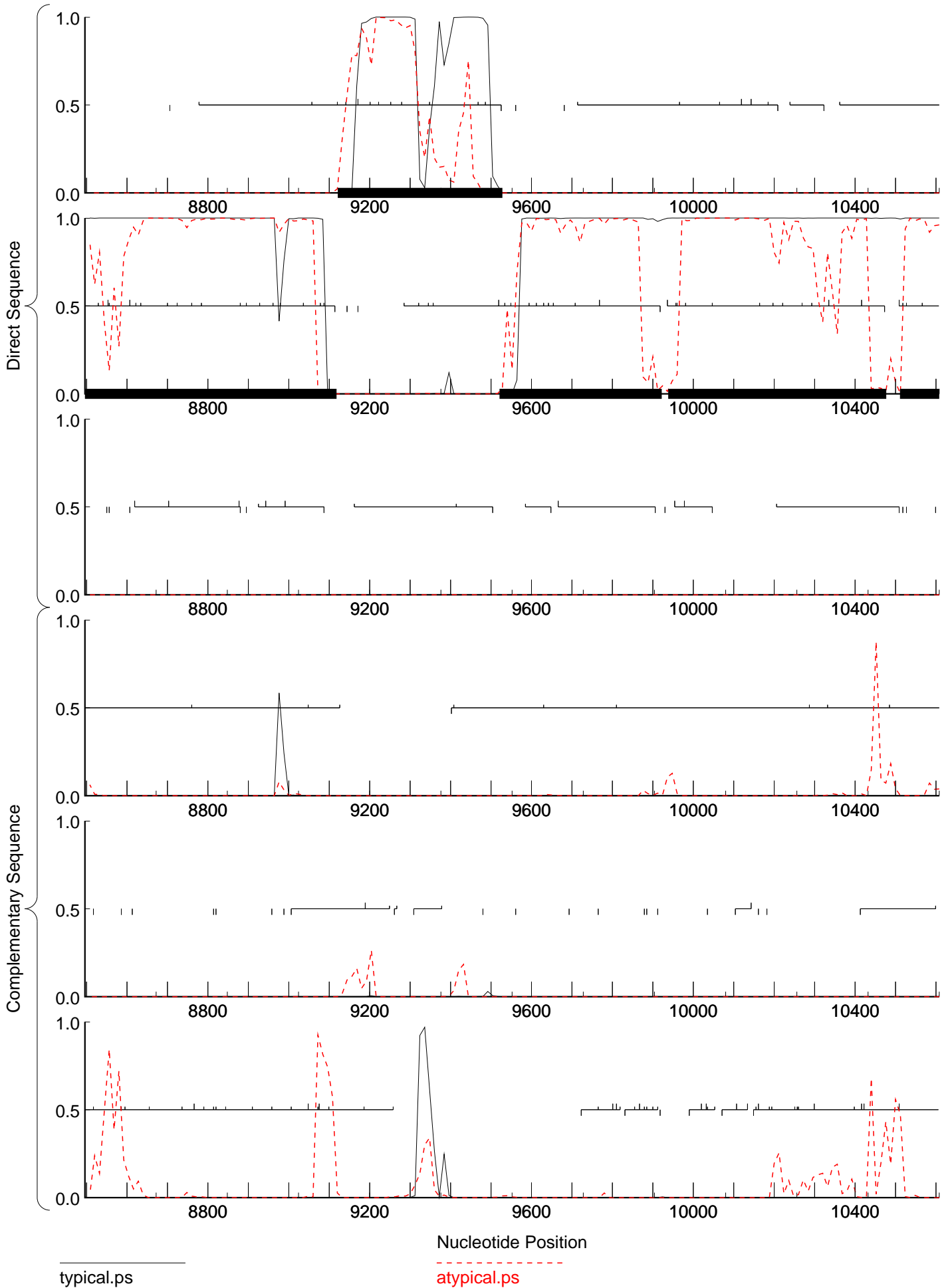
Send questions / comments to:
Dr. M. Borodovsky
Georgia Institute of Technology
School of Biology
Atlanta, GA 30332-0230





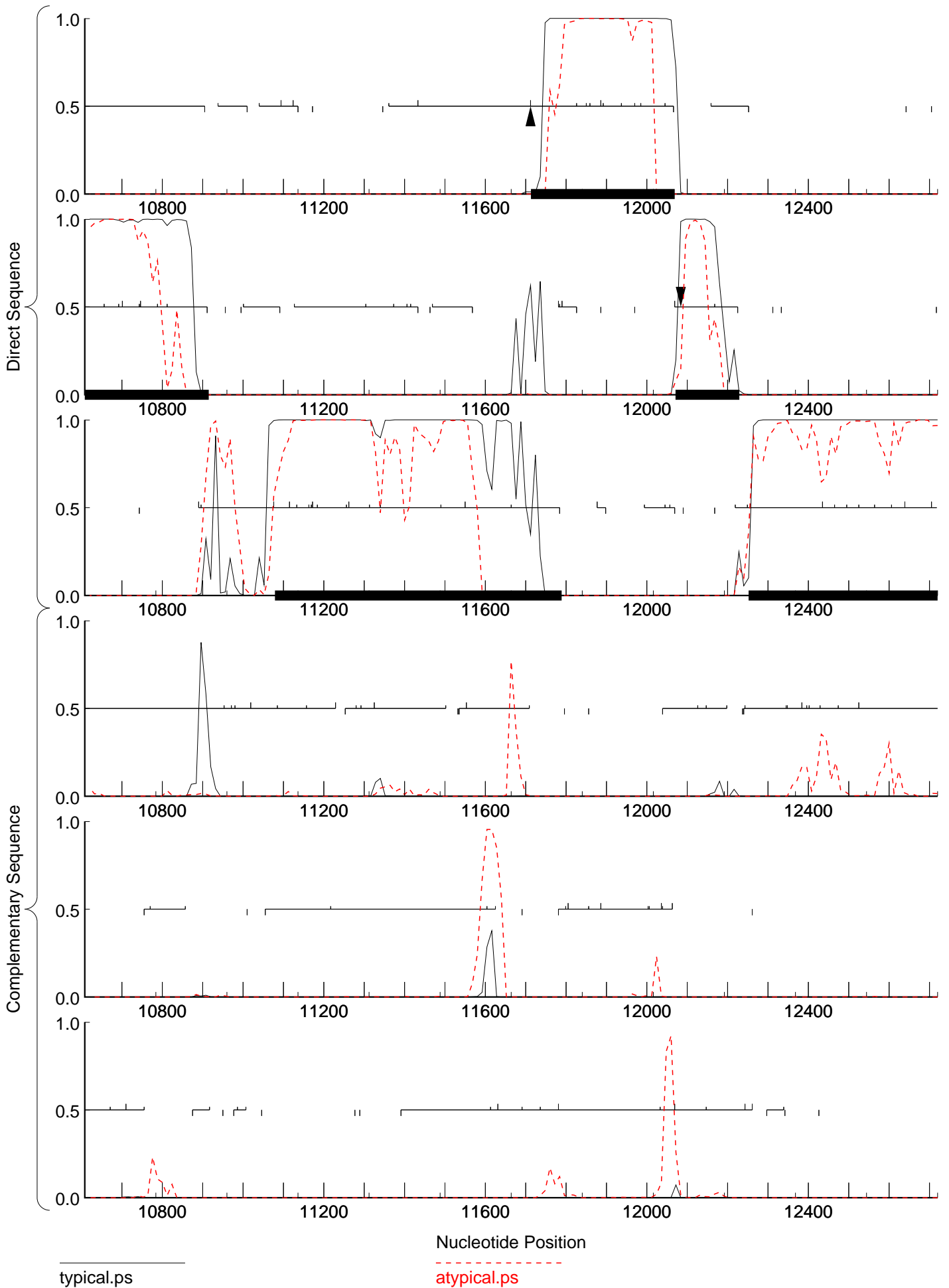


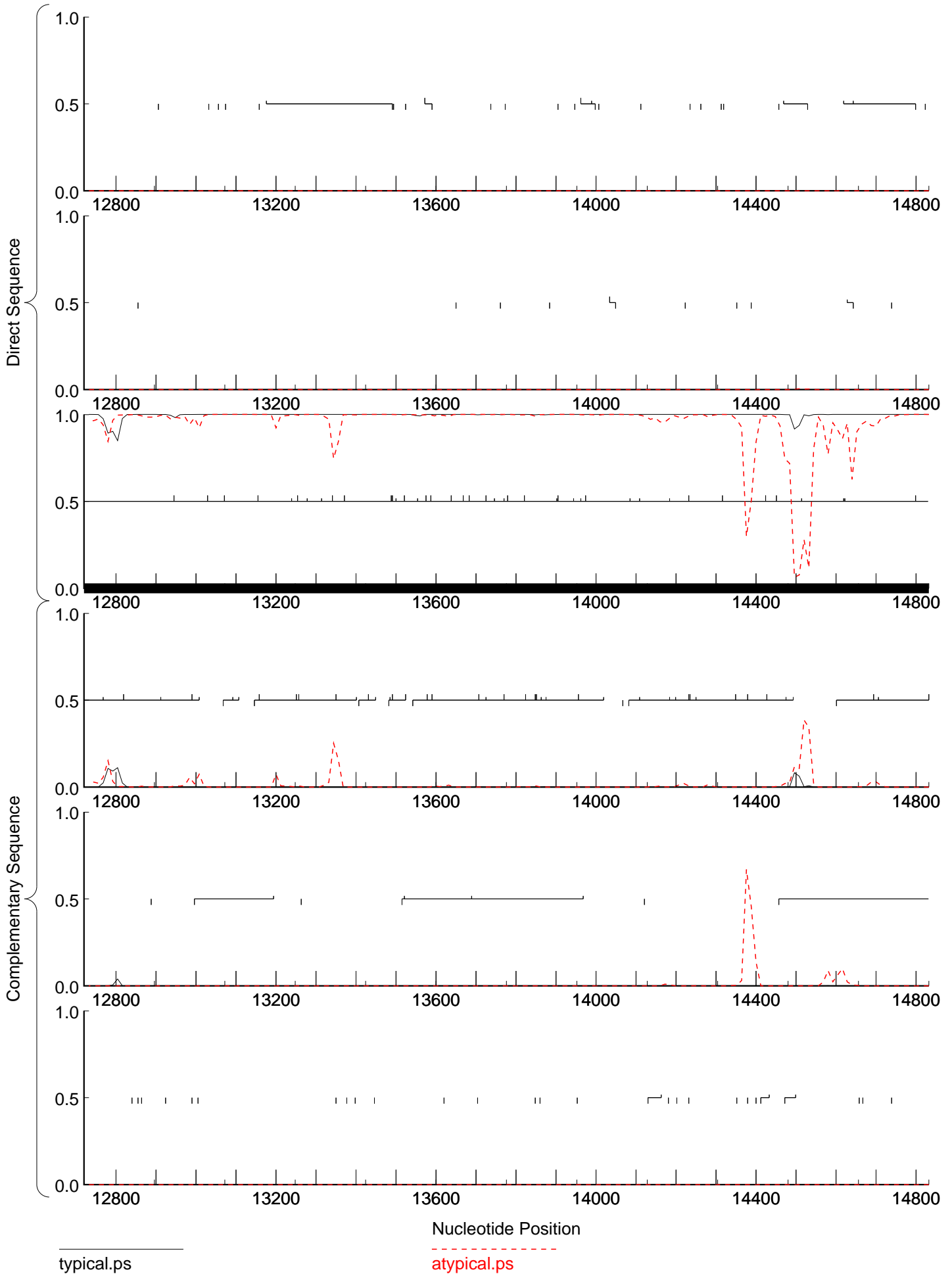


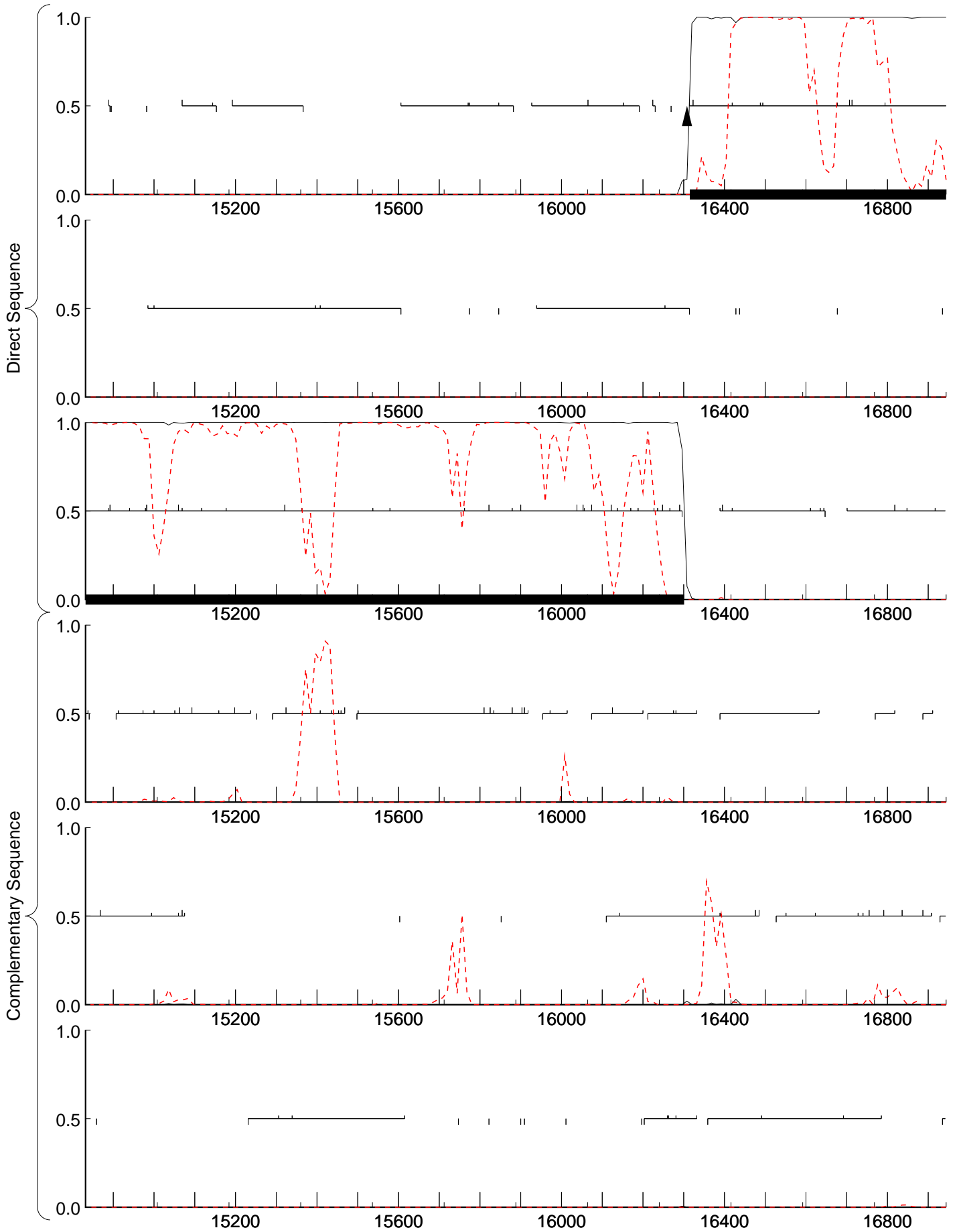


typical.ps

atypical.ps

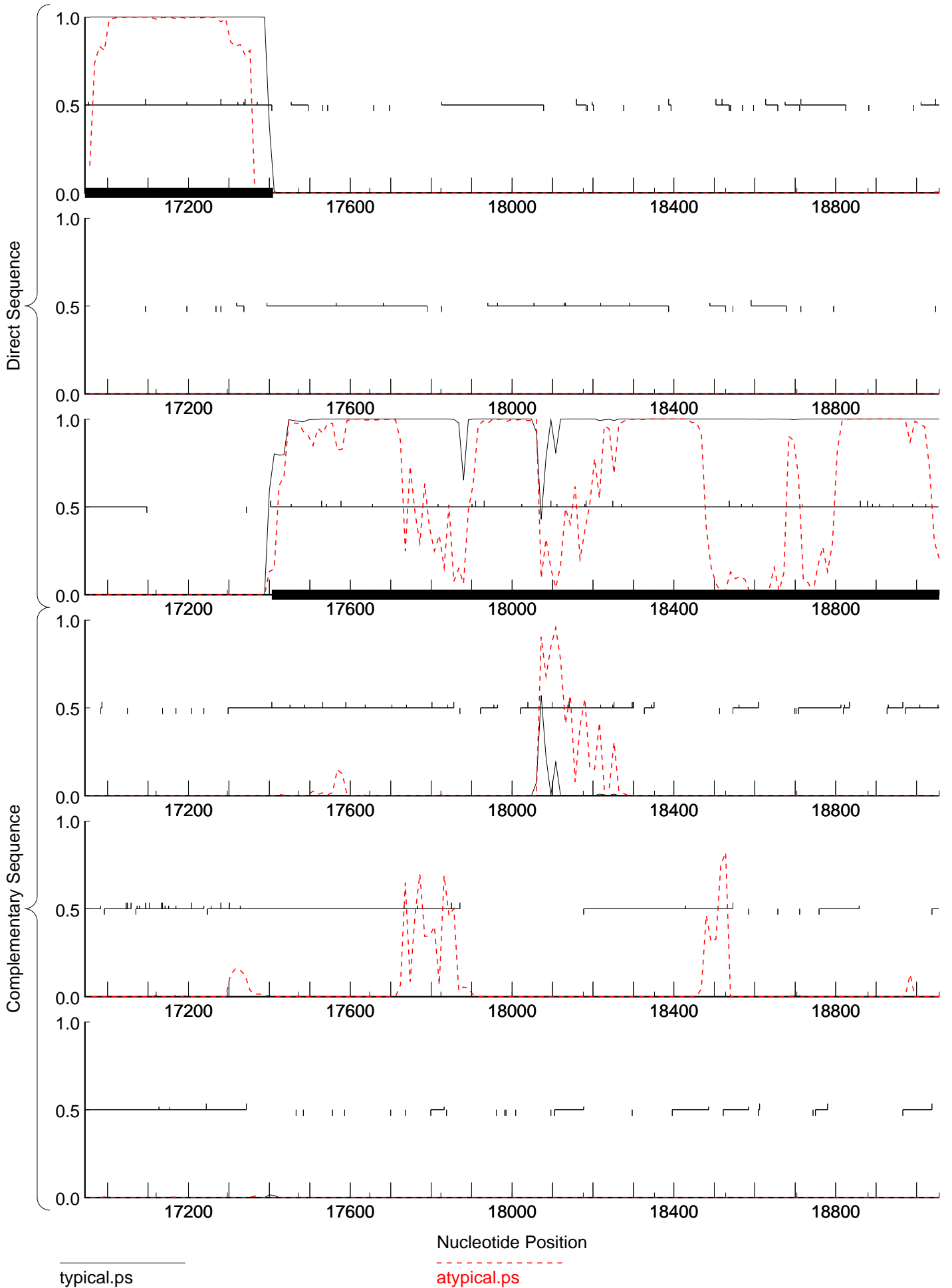


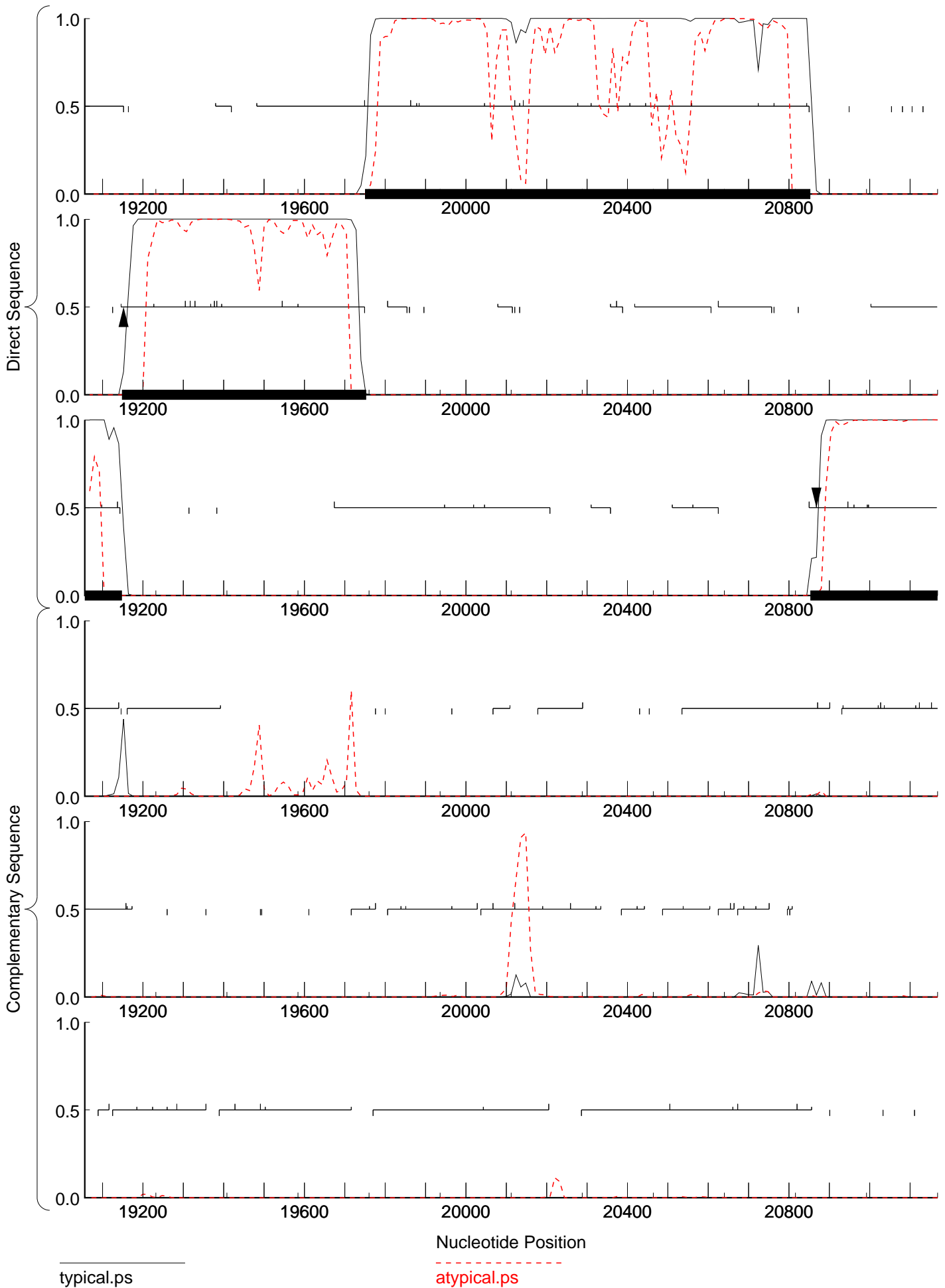


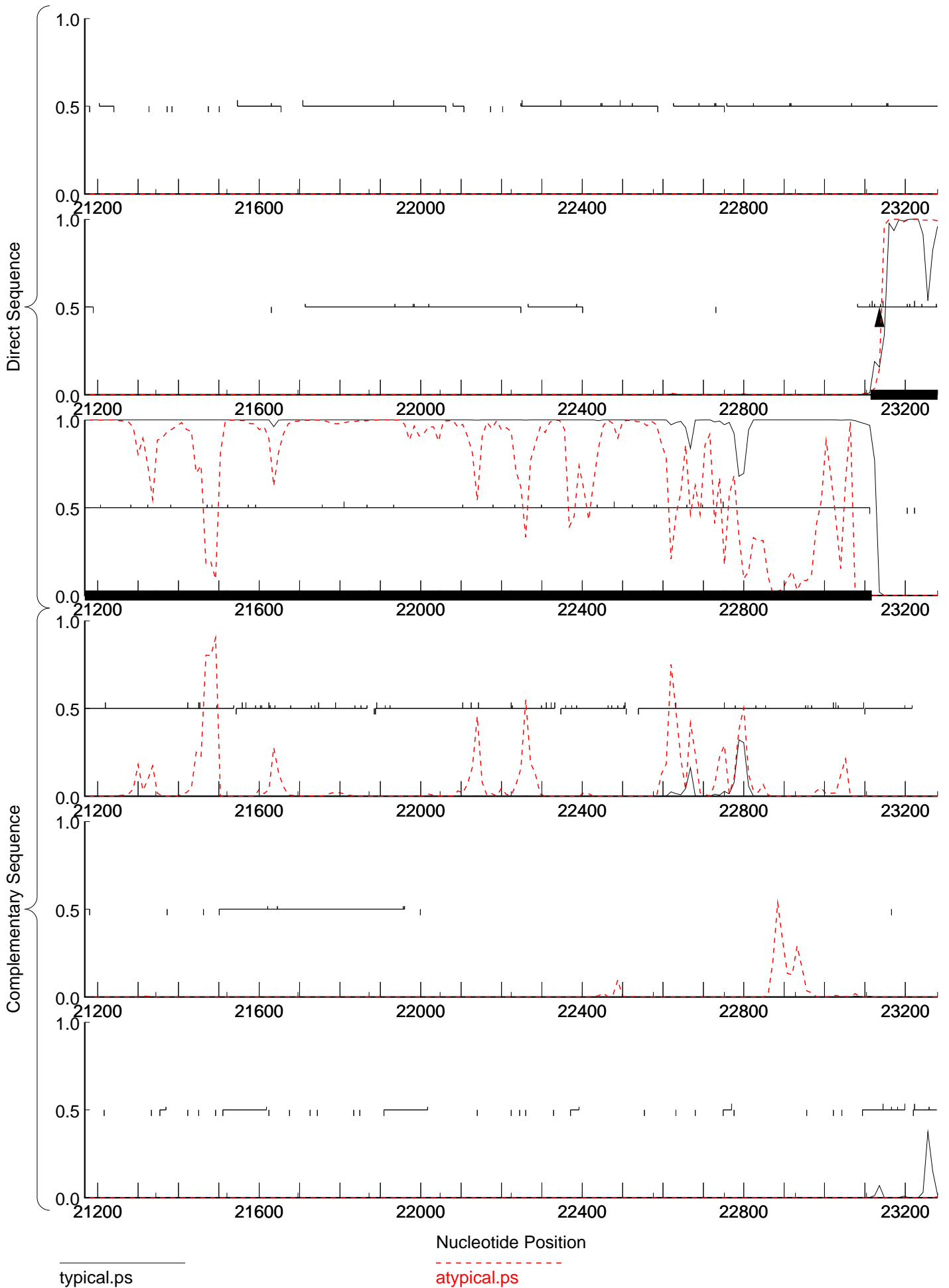


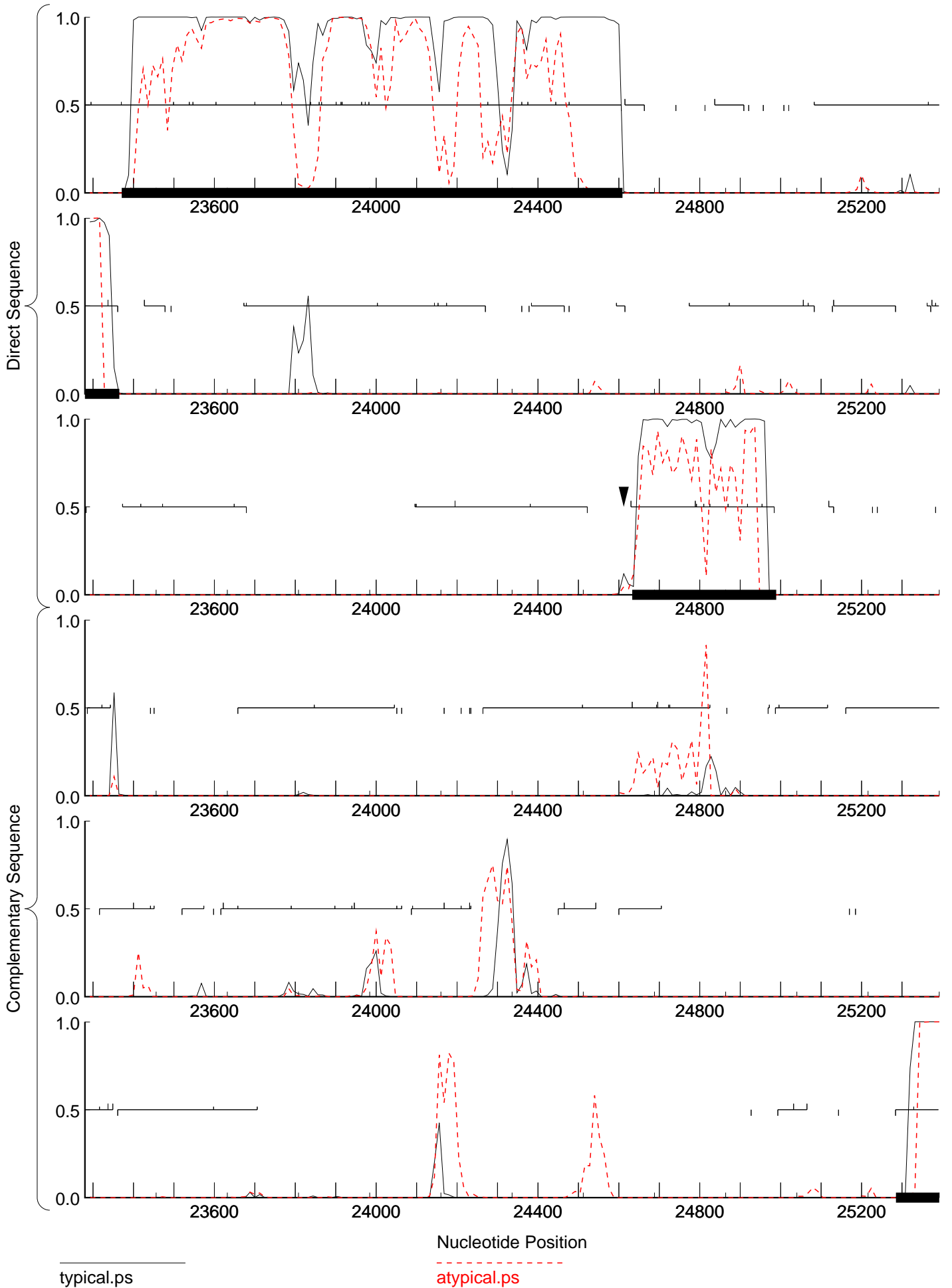
typical.ps

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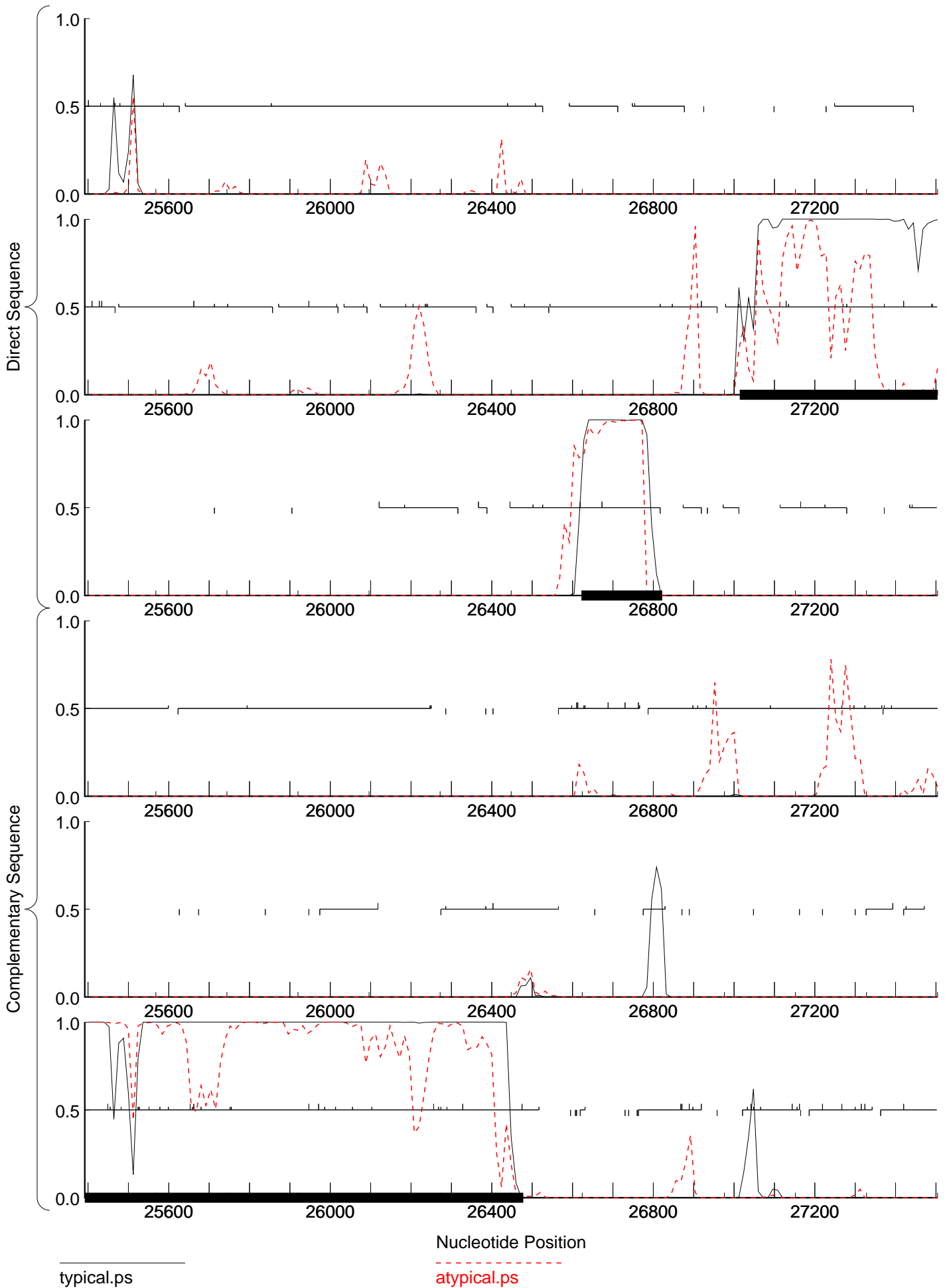


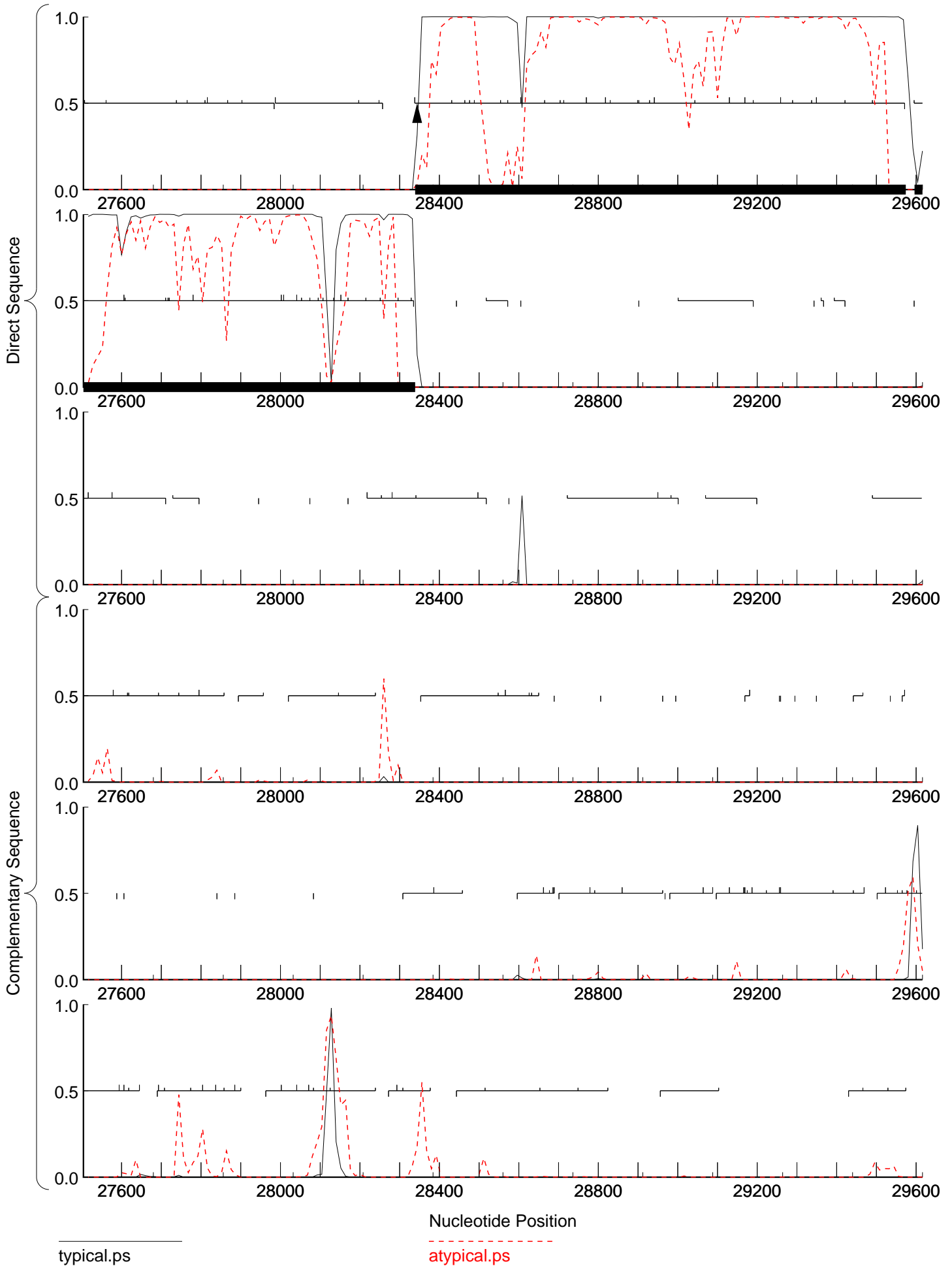


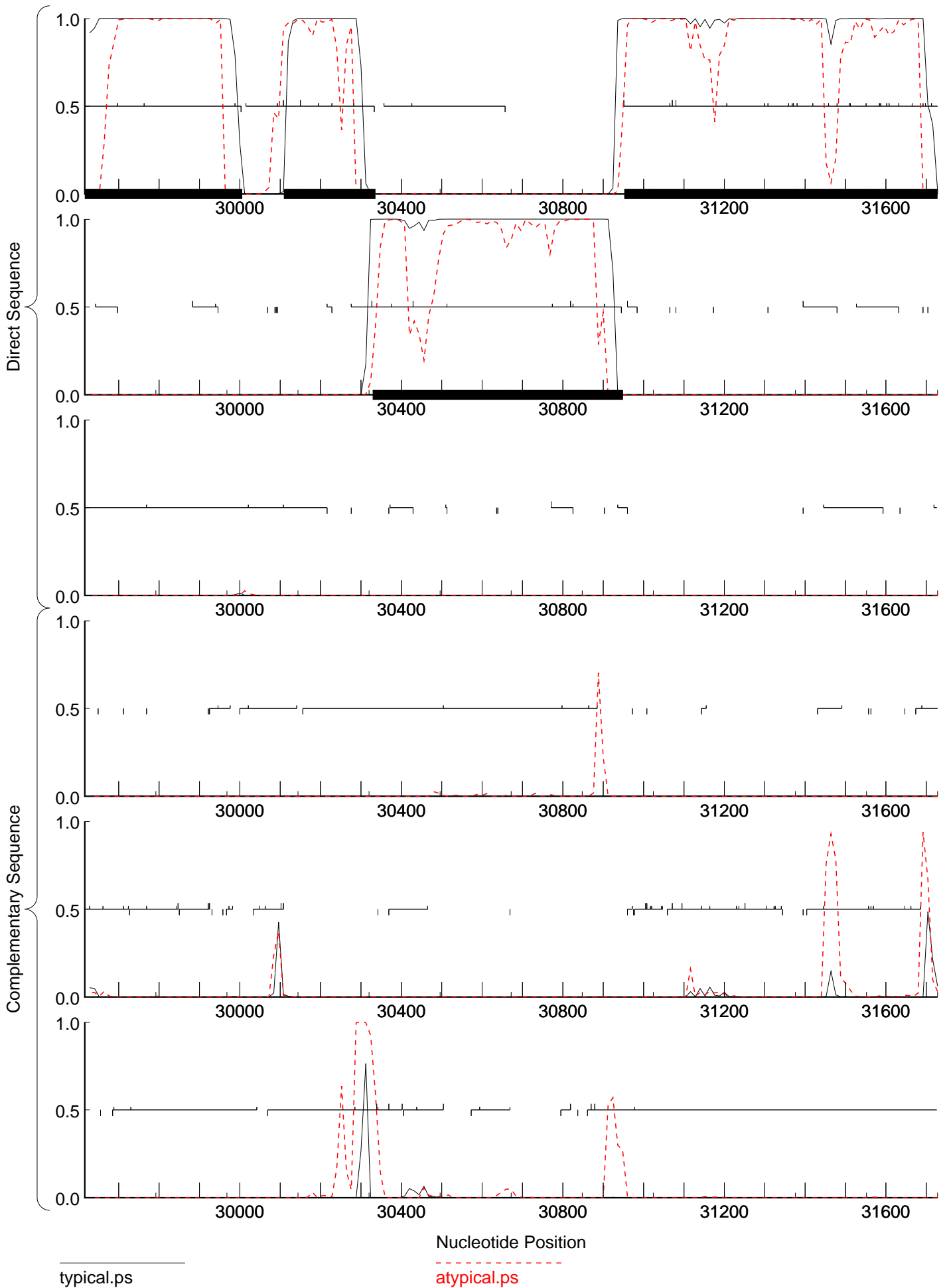


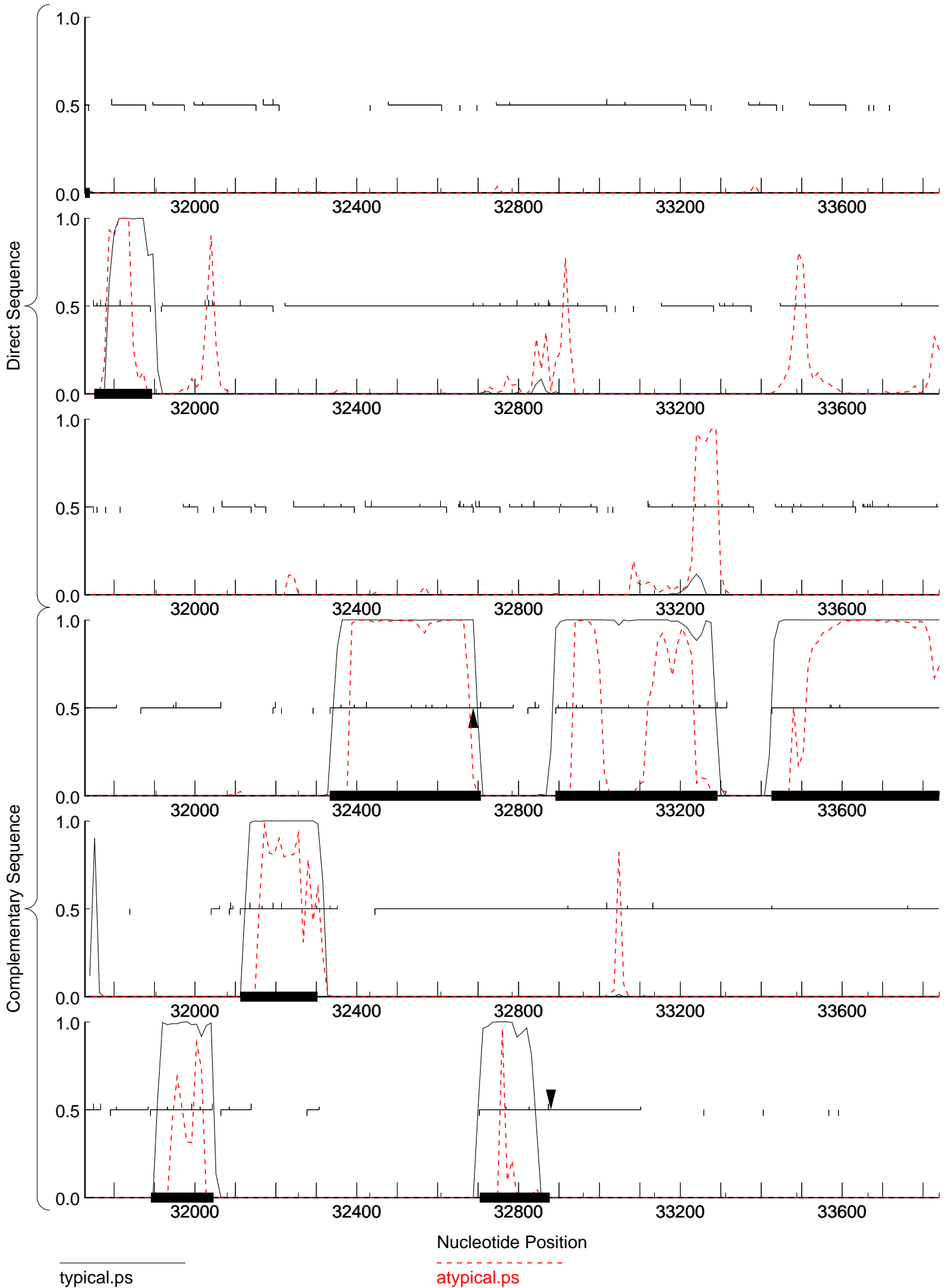
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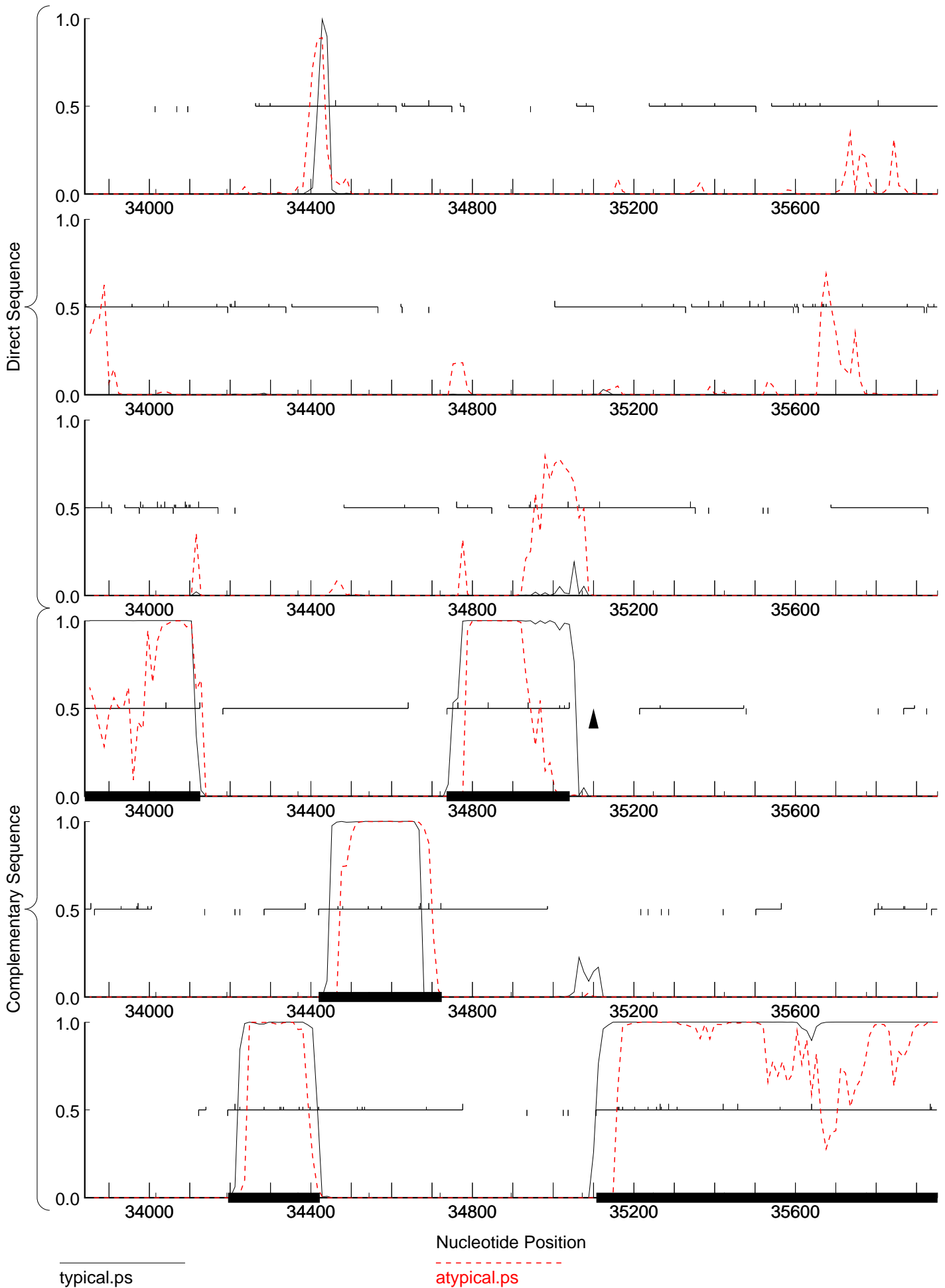
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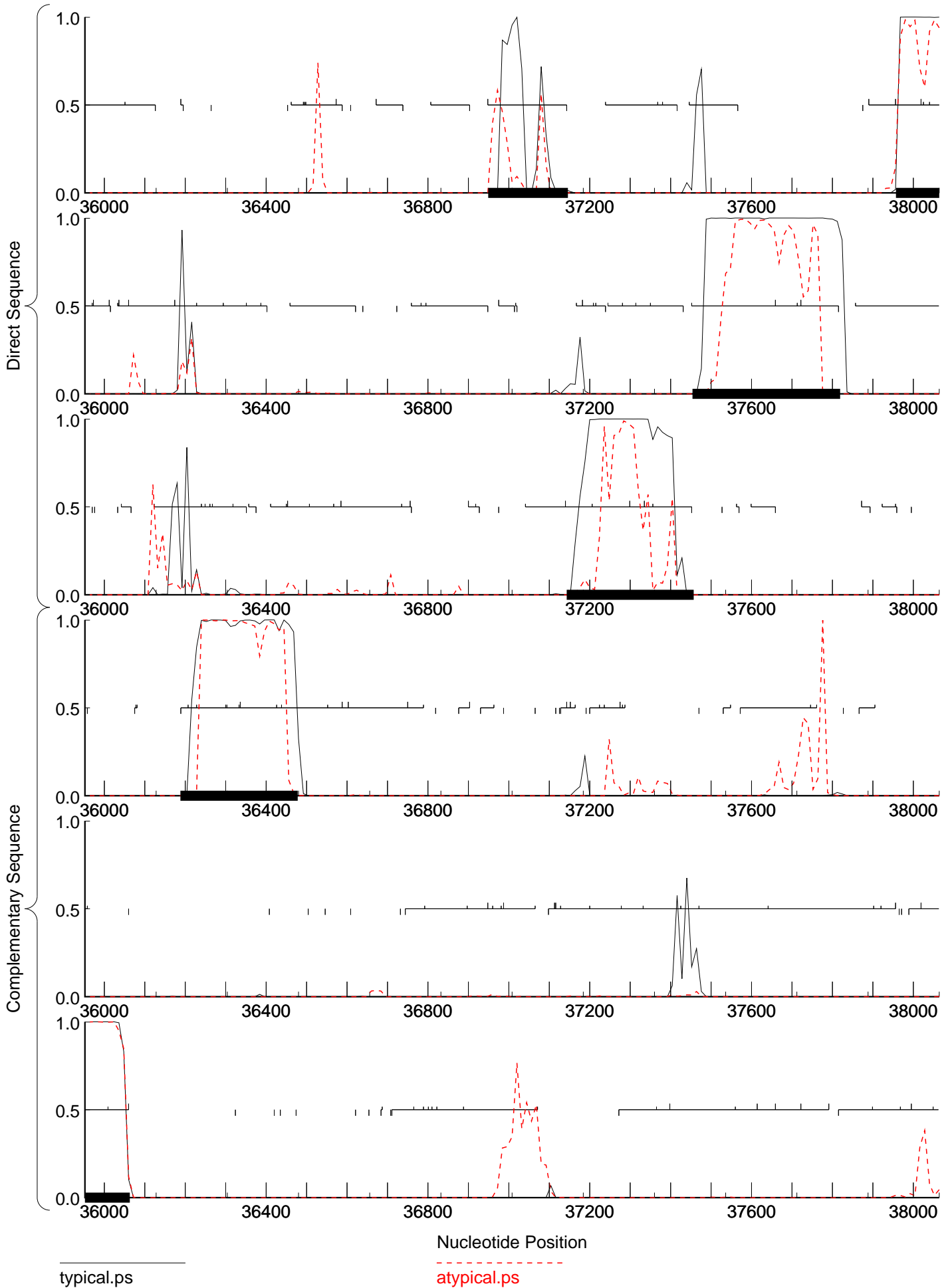


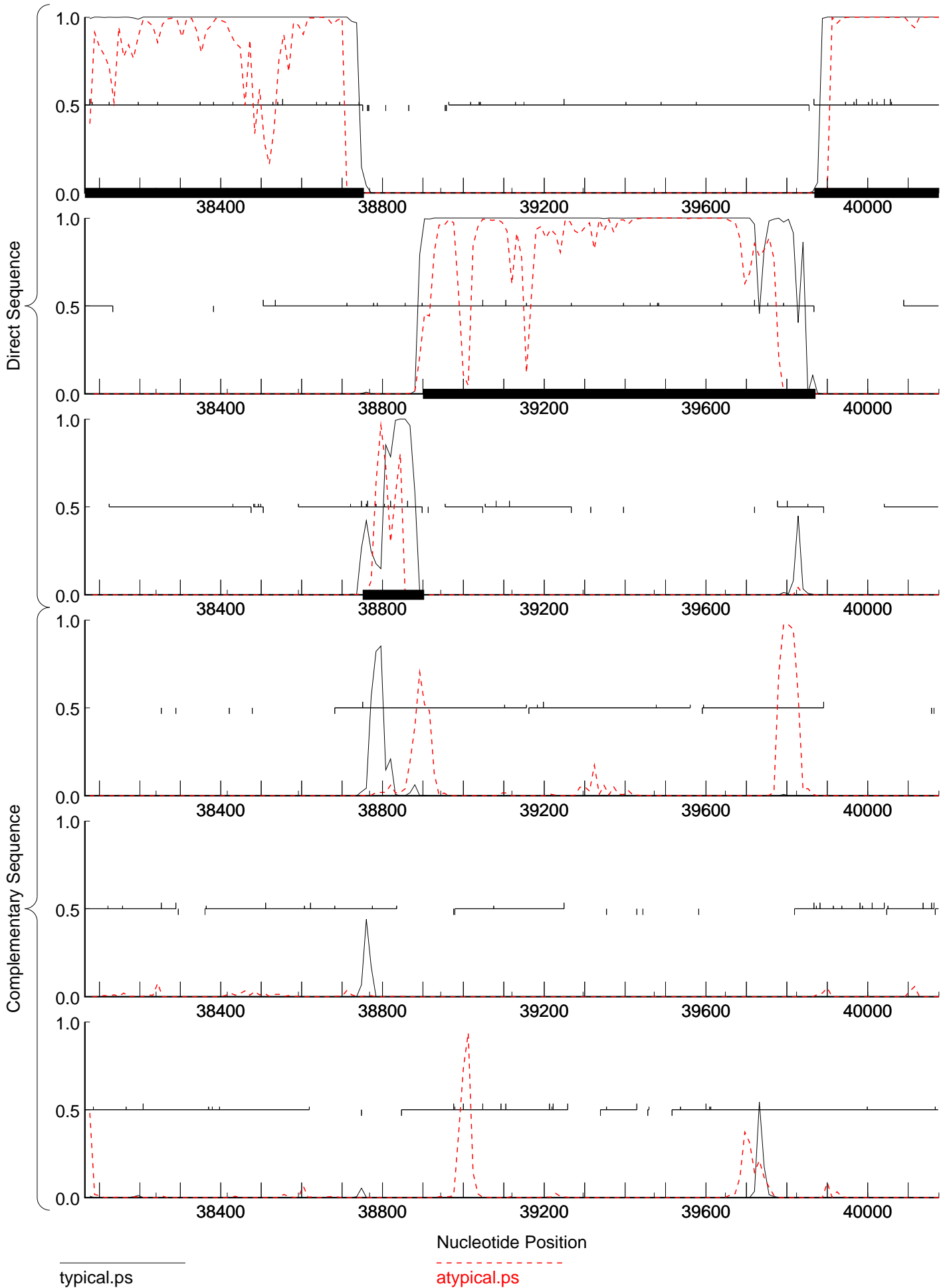


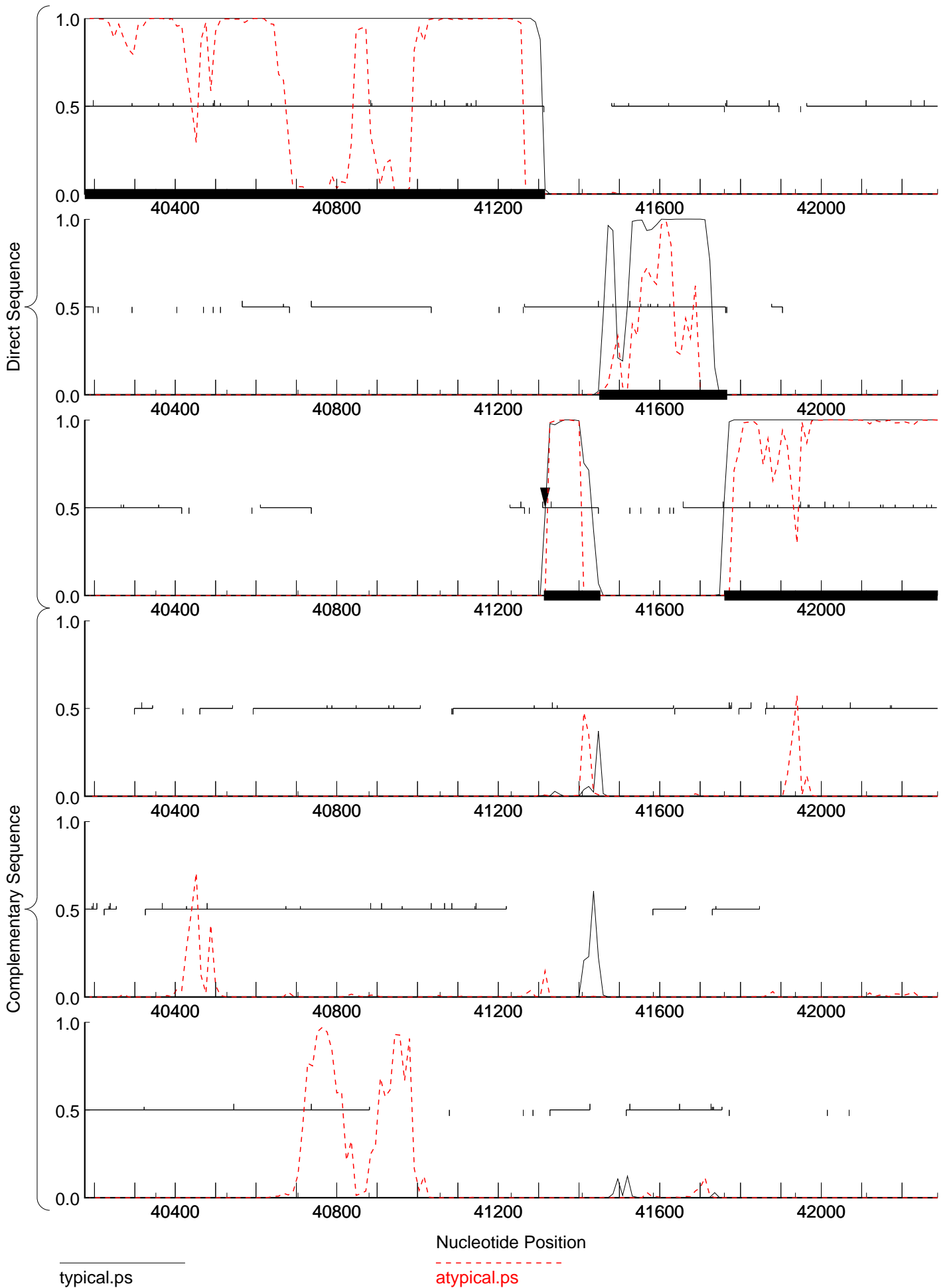


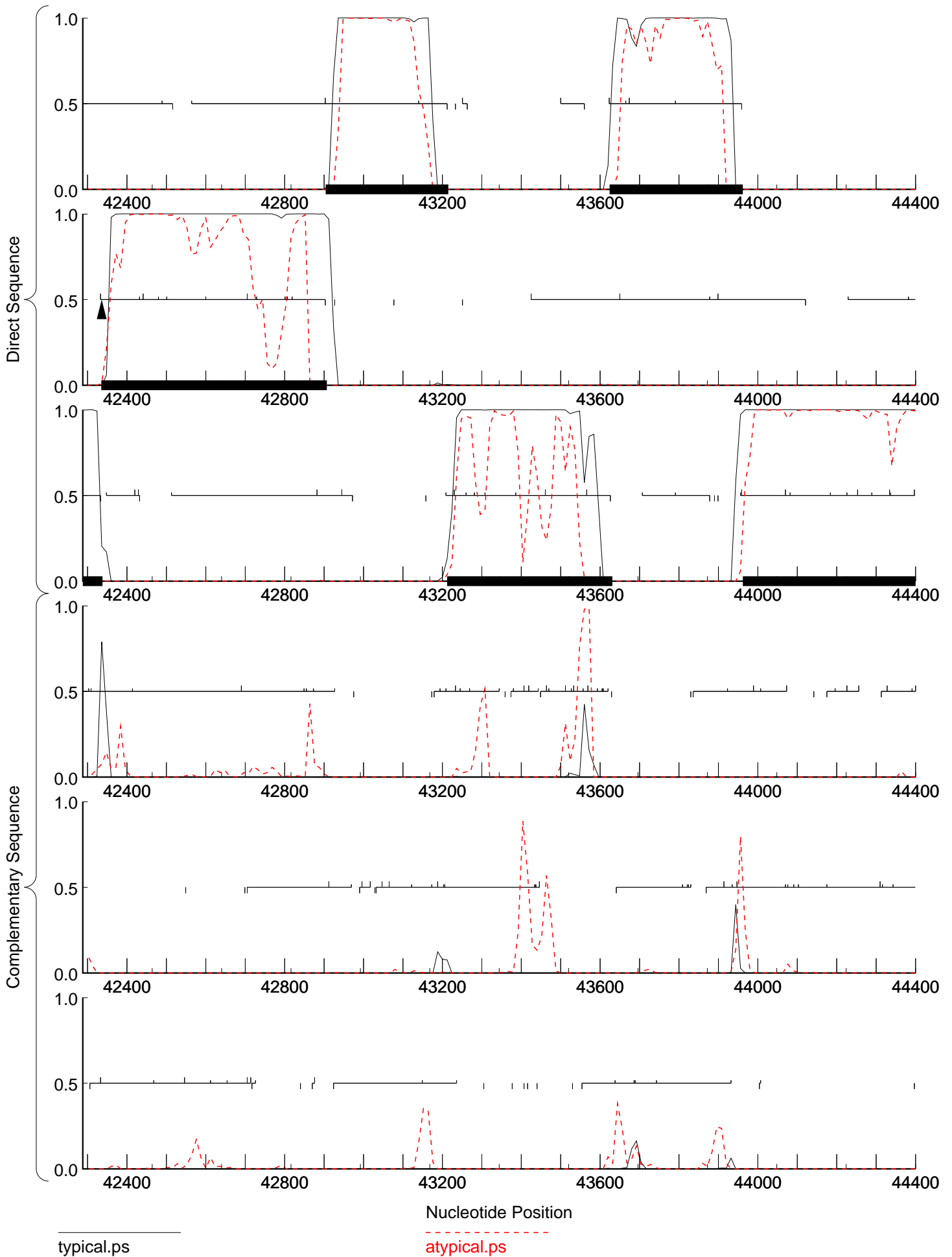


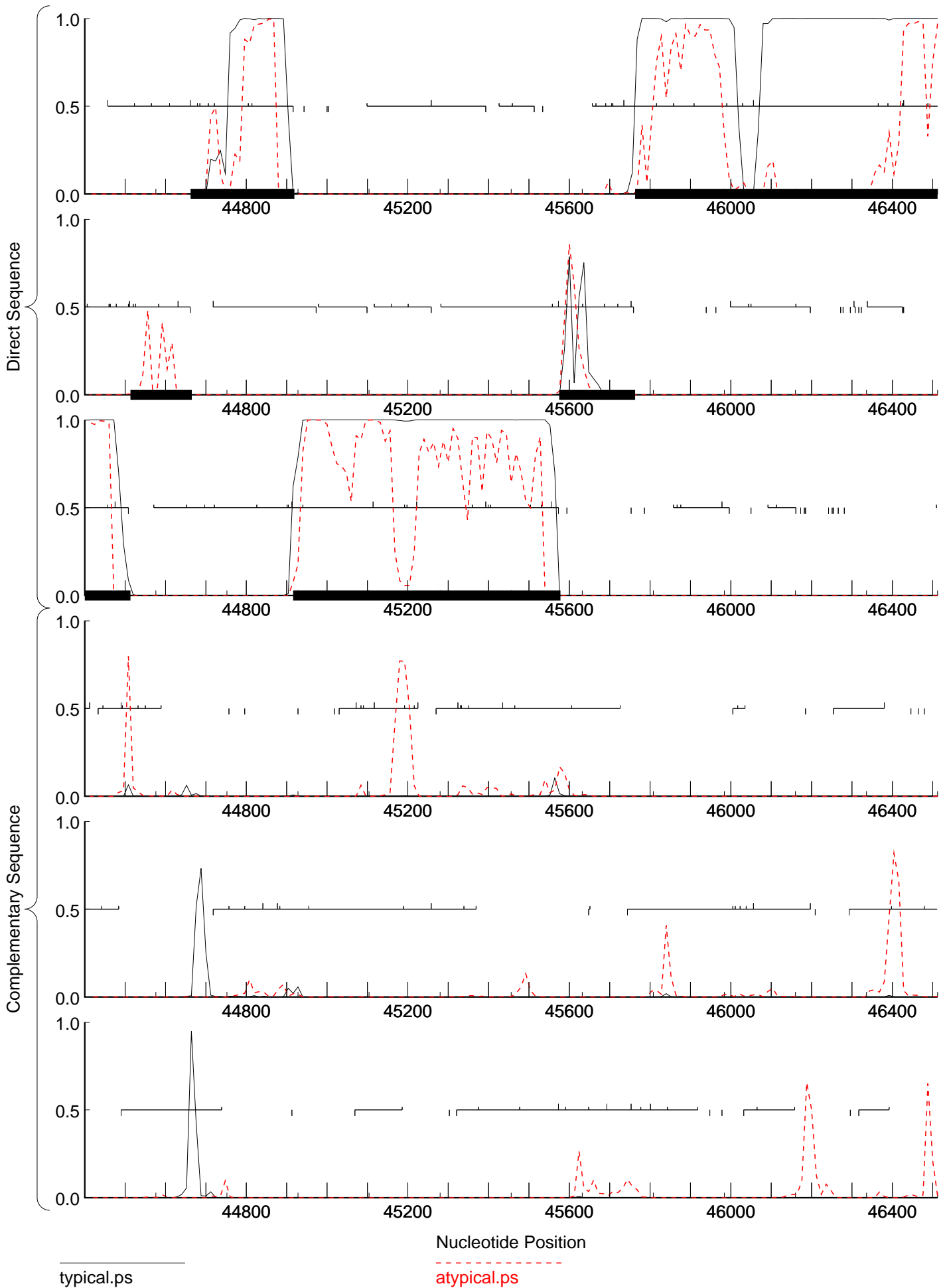






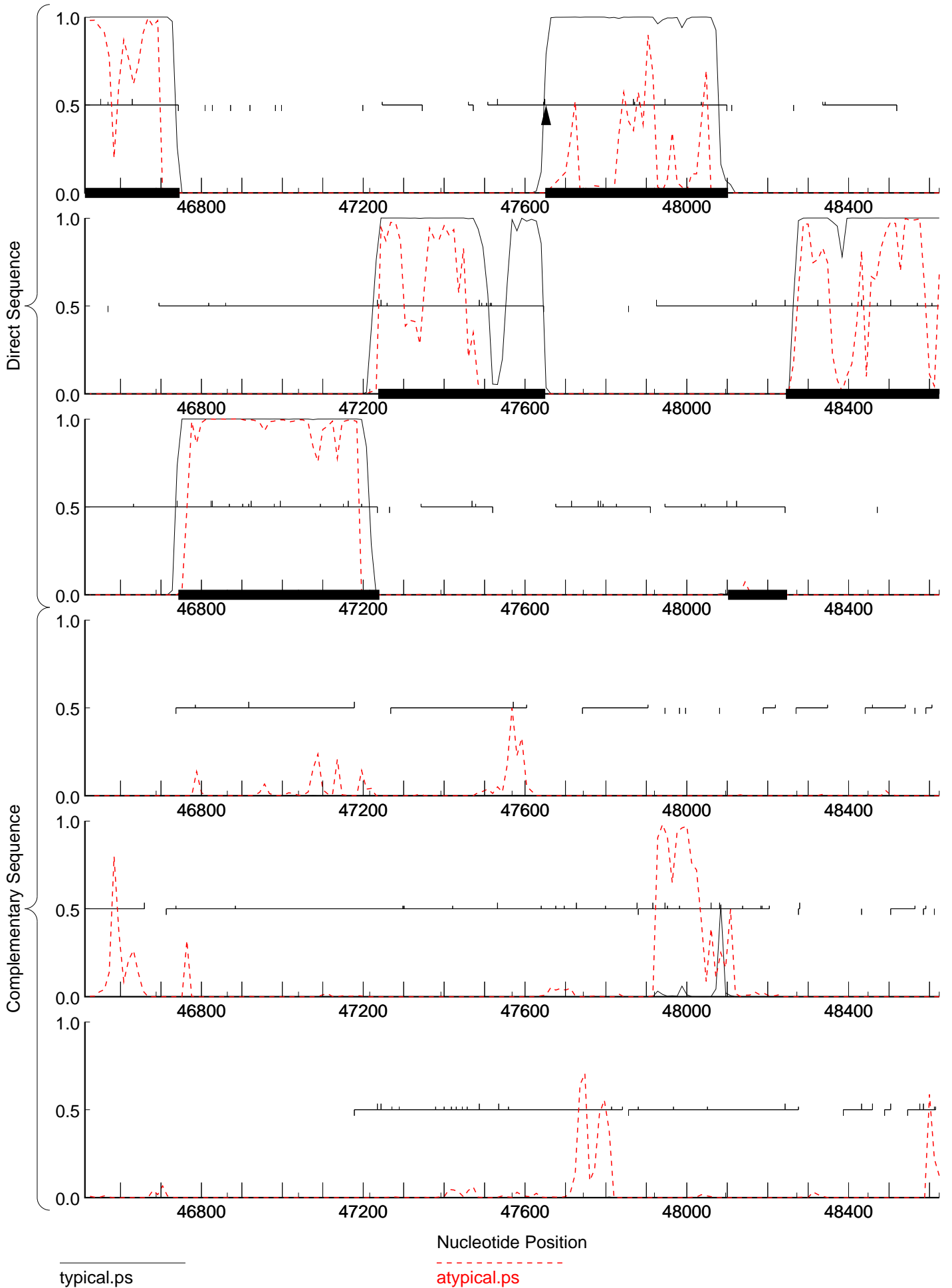






typical.ps

atypical.ps



typical.ps

atypical.ps

