

GeneMark

Version 2.5p (09.08.06)

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PROGRAM INFORMATION

Sequence : Mycobacteriophage Inca complete genome, 74070 bp including 9-base 3' overhang (CGCTTGTC),
Analysis Date : 9/23/17 at 3:36:35
Pages : 37
Sequence Length : 74070 bp
GC Content : 62.80%

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,

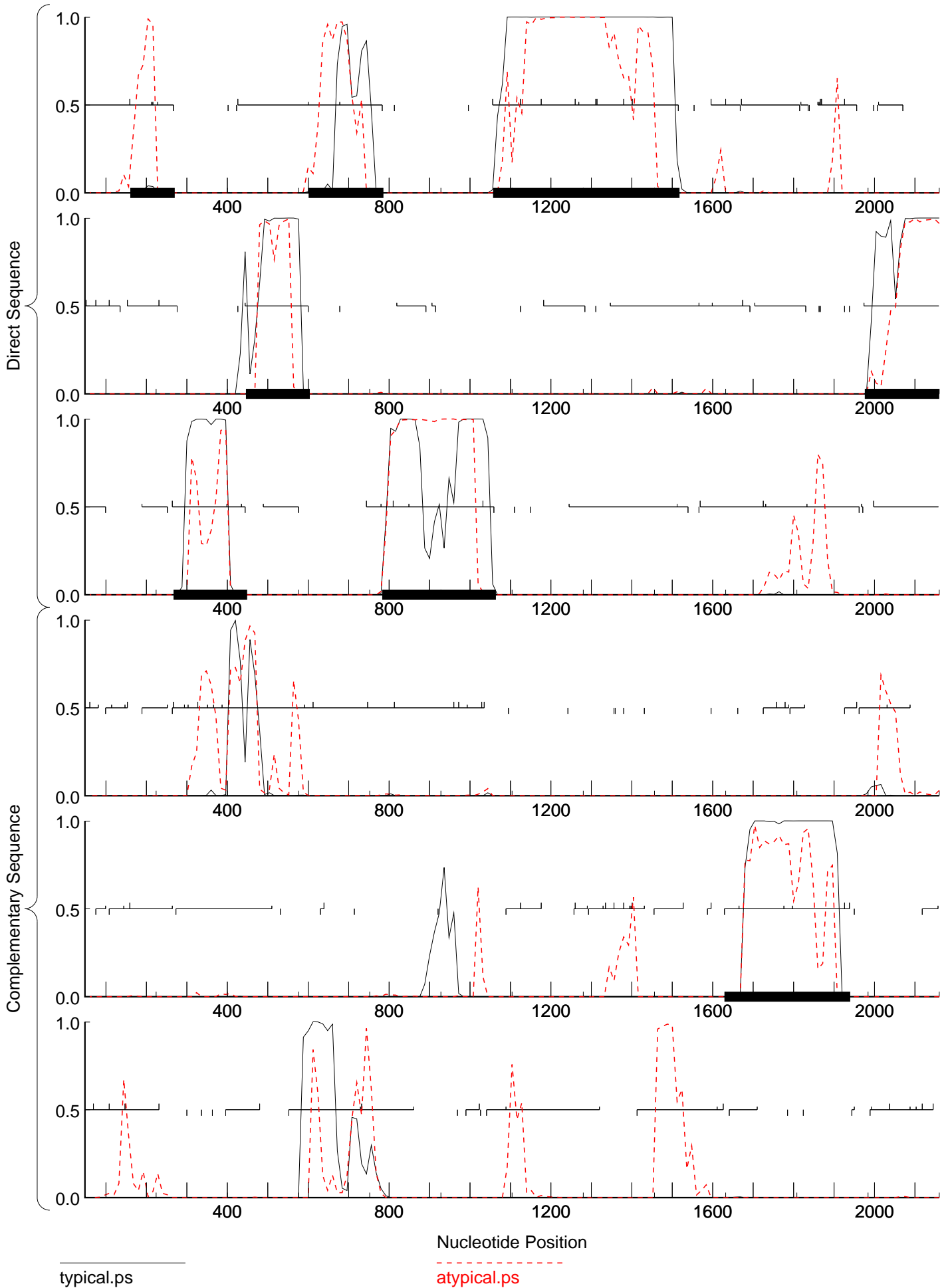
Matrix notes & comments

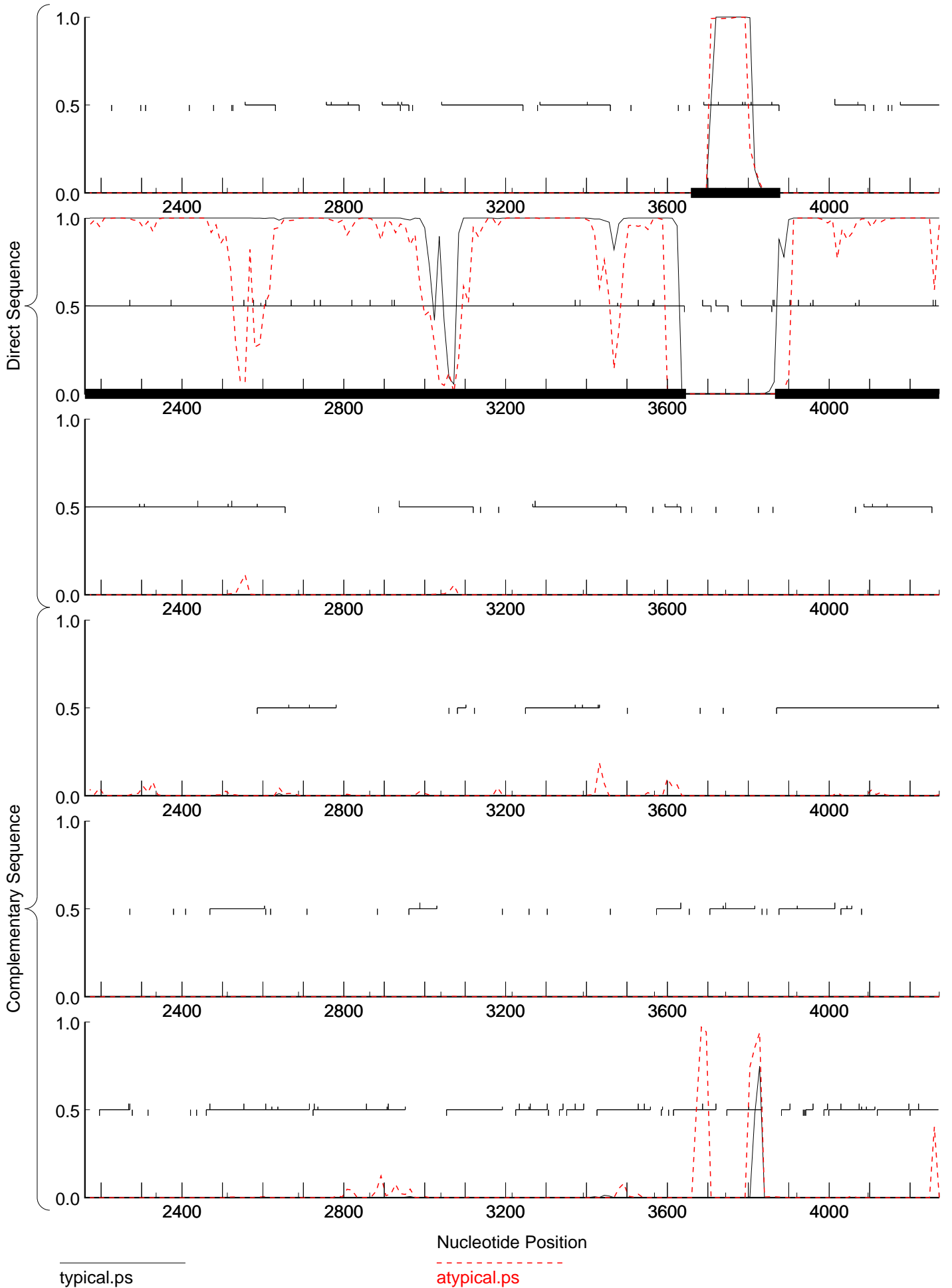
Training set derived by GeneMarkS, 4.30 October 2014
Sat Sep 23 03:36:35 2017

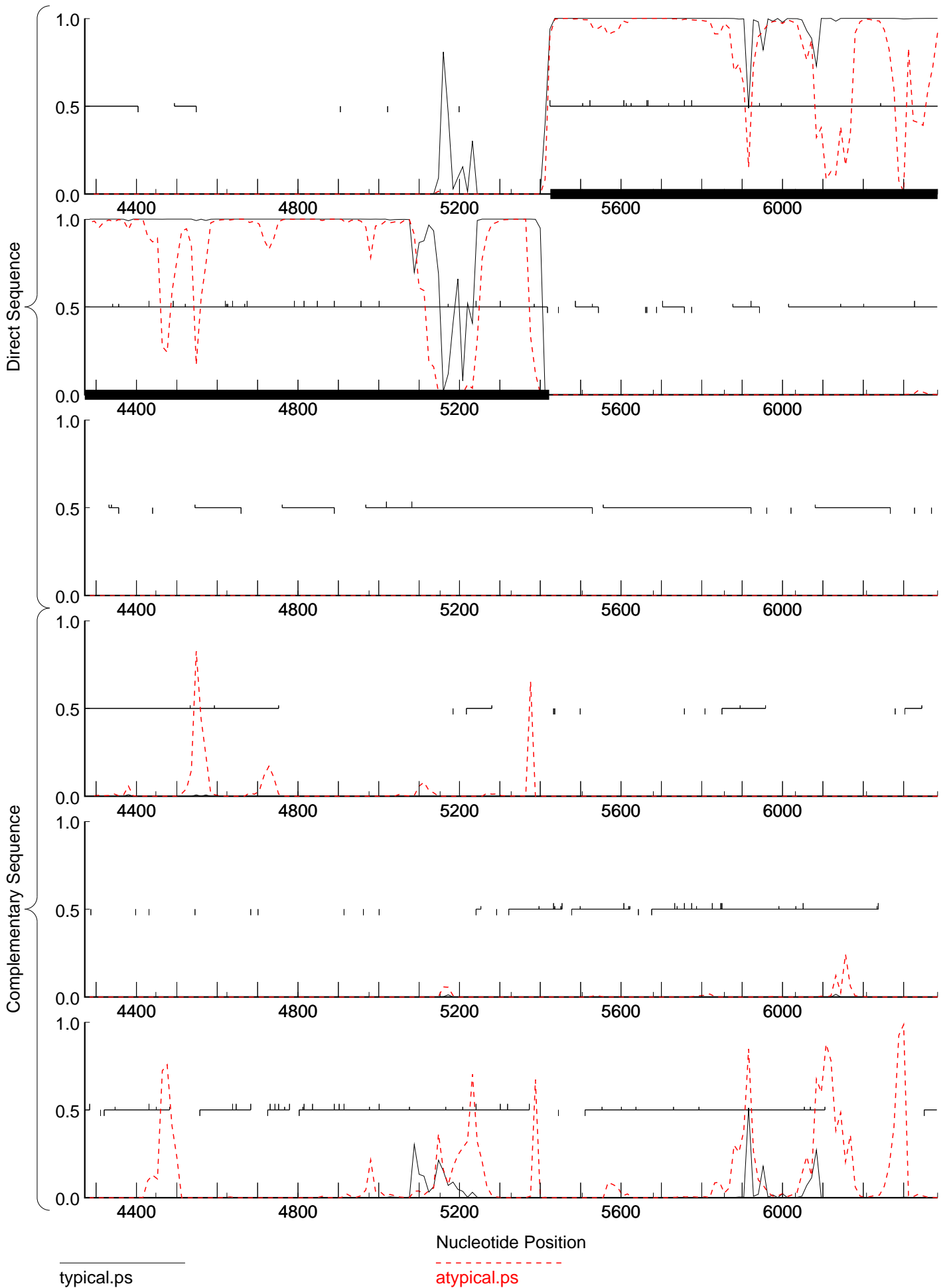
MATRIX INFORMATION

Matrix : -
Author : -
Order : 4

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

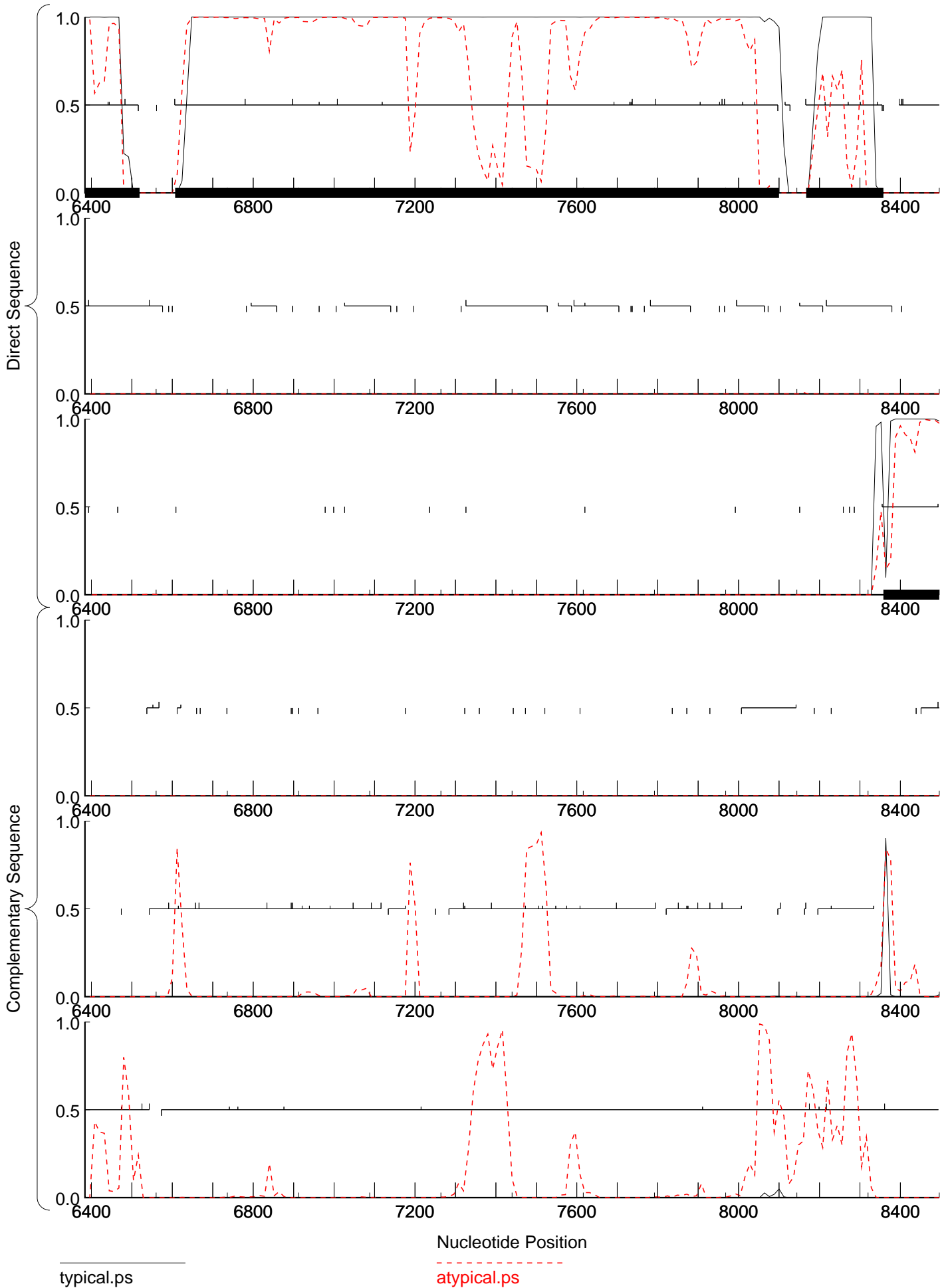


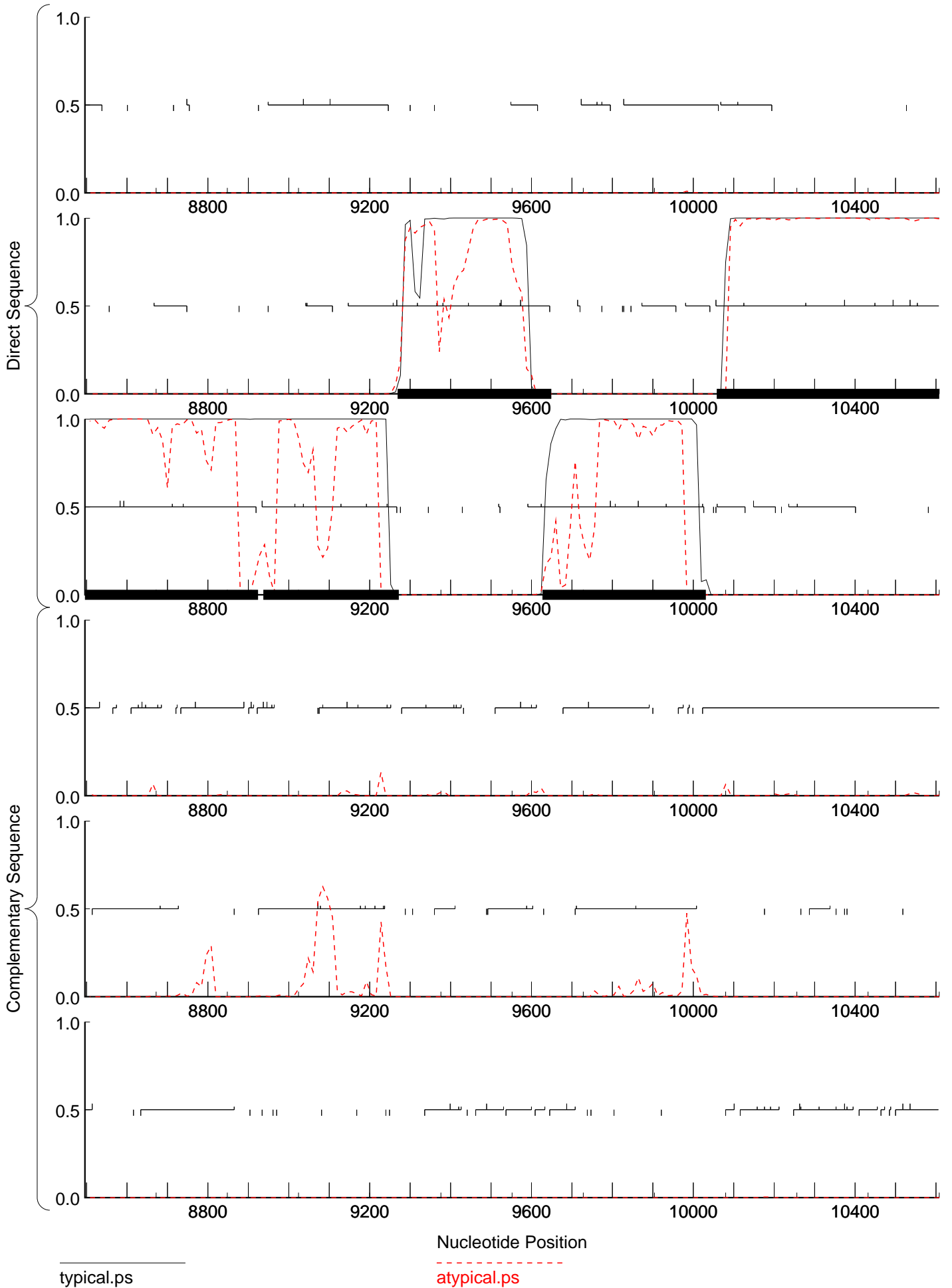


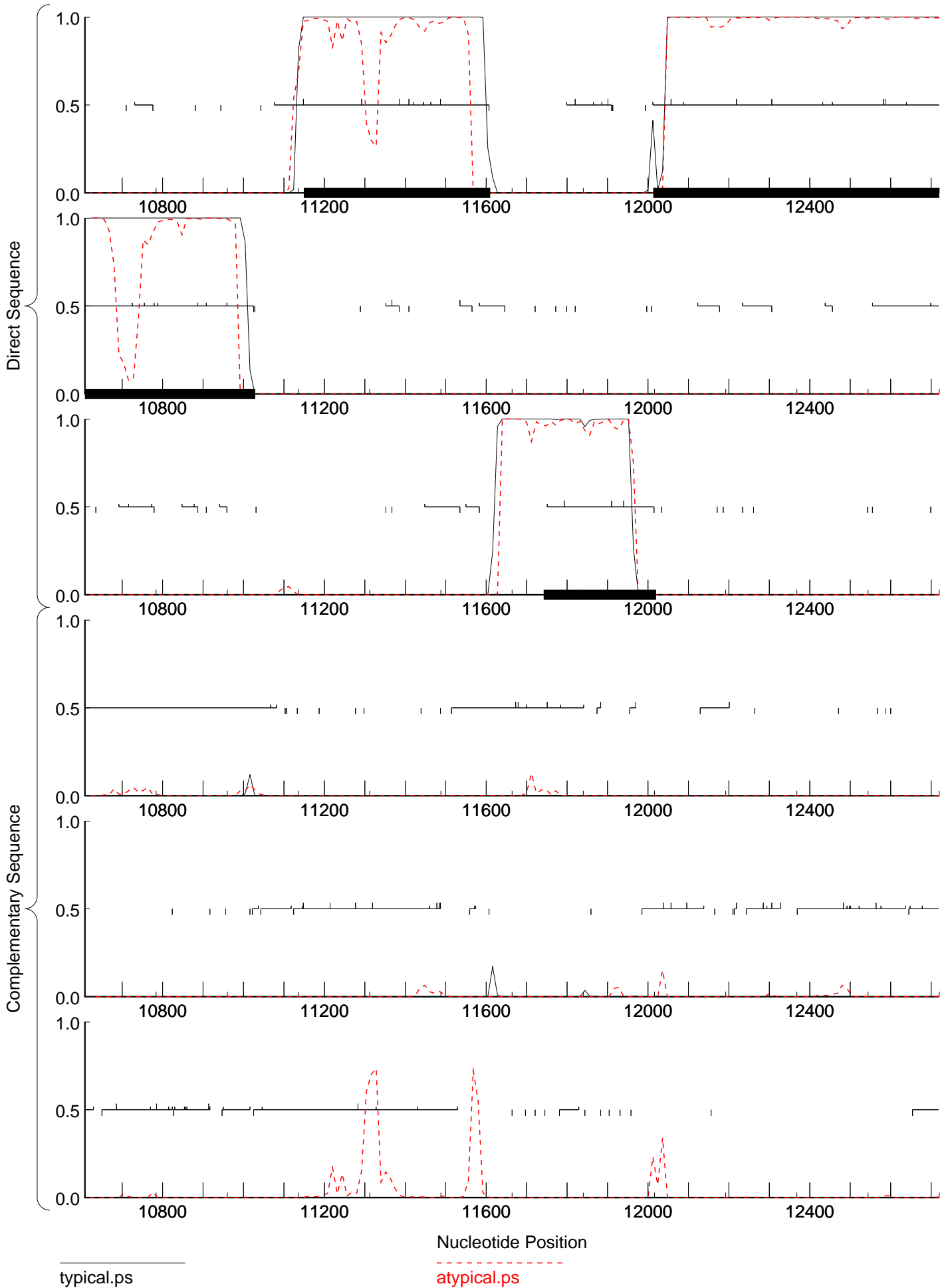


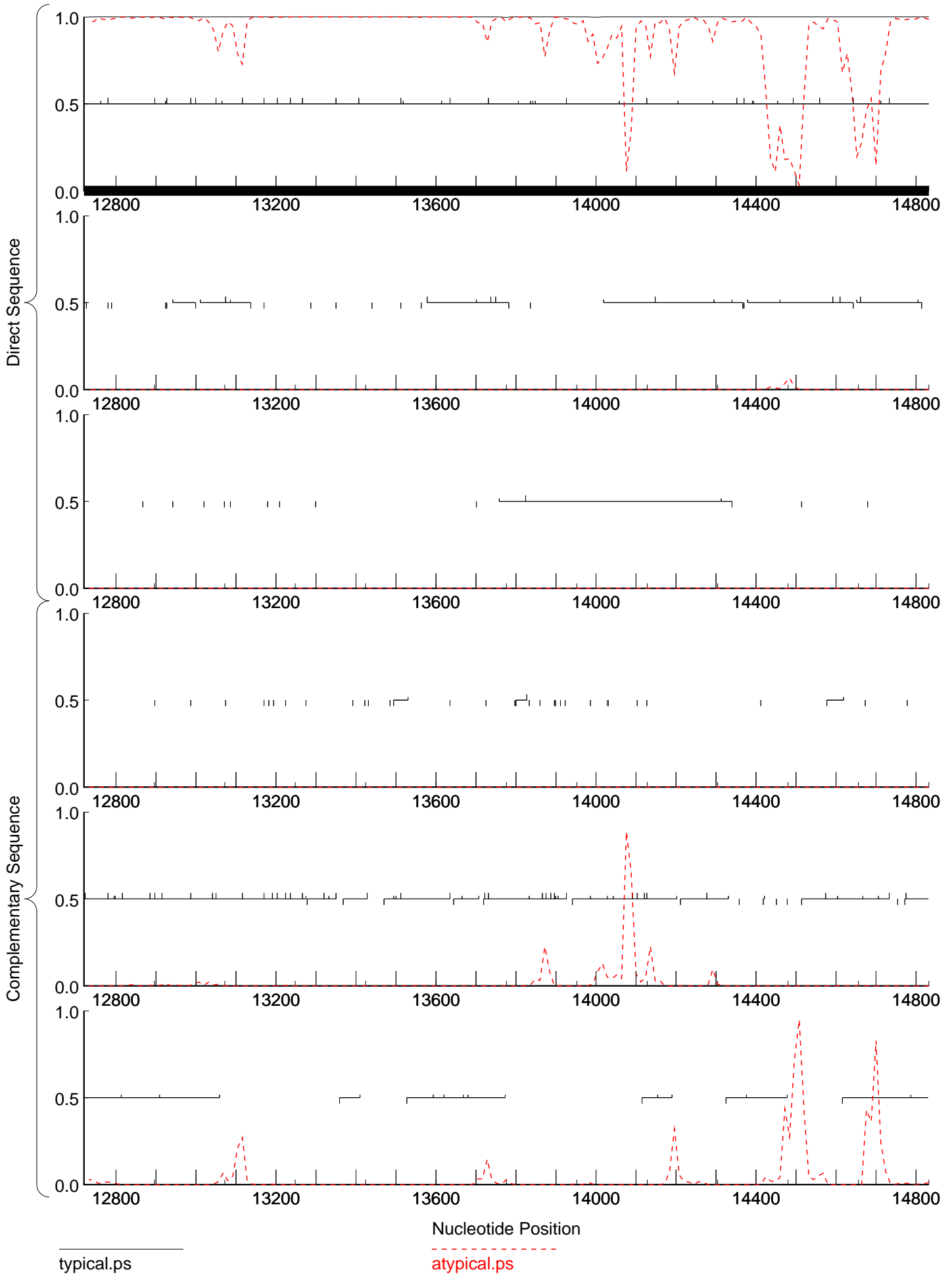
typical.ps

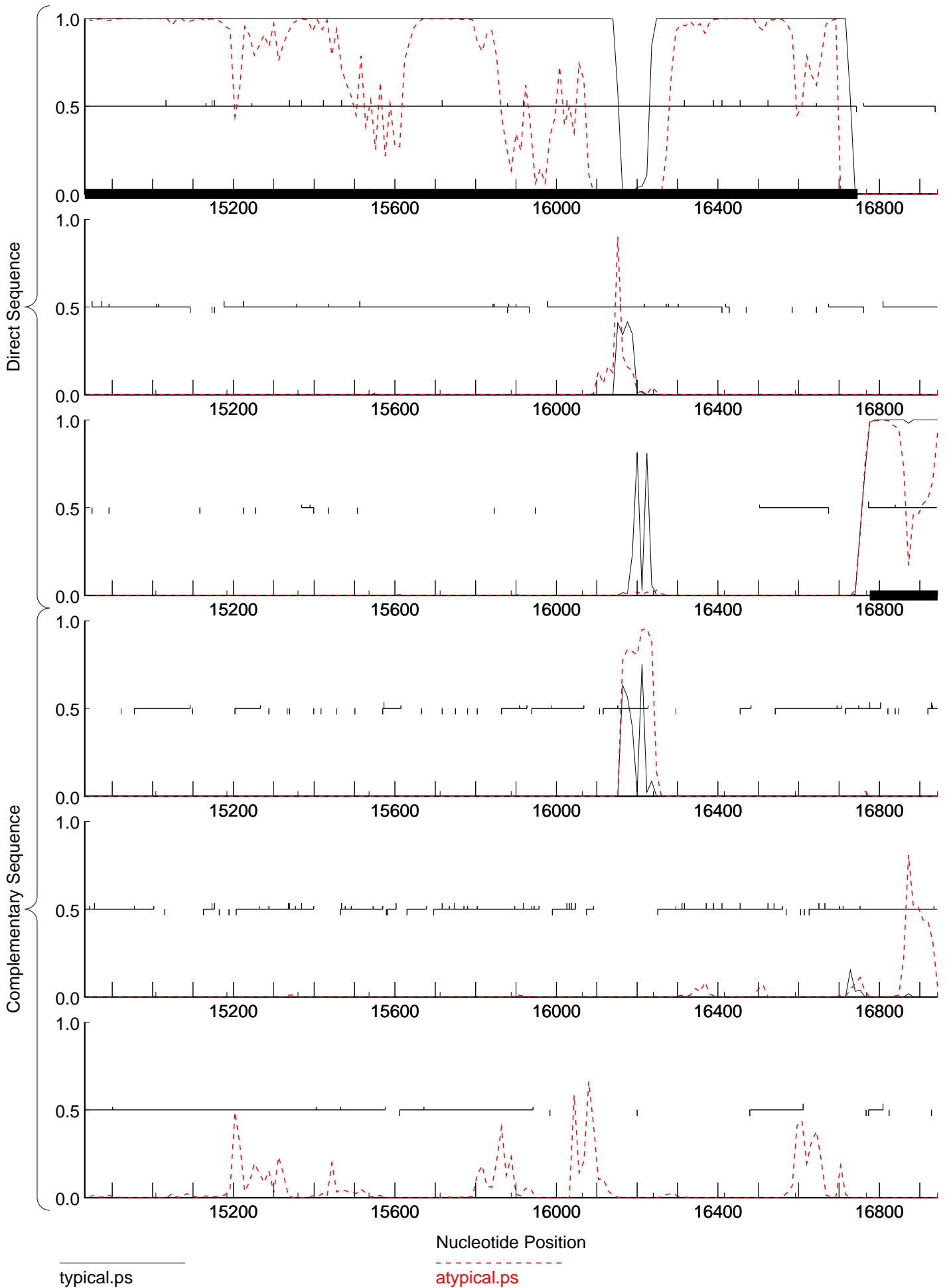
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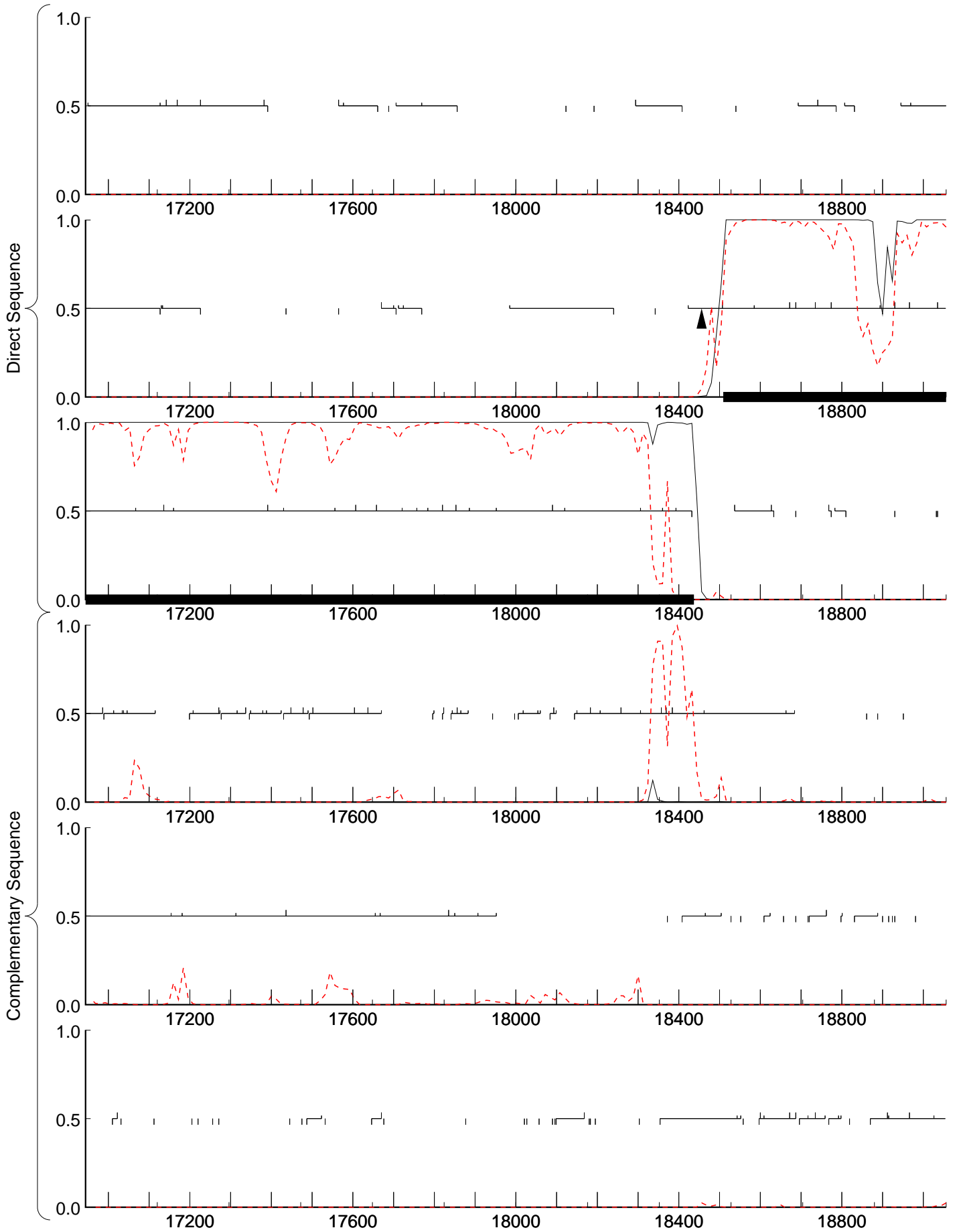








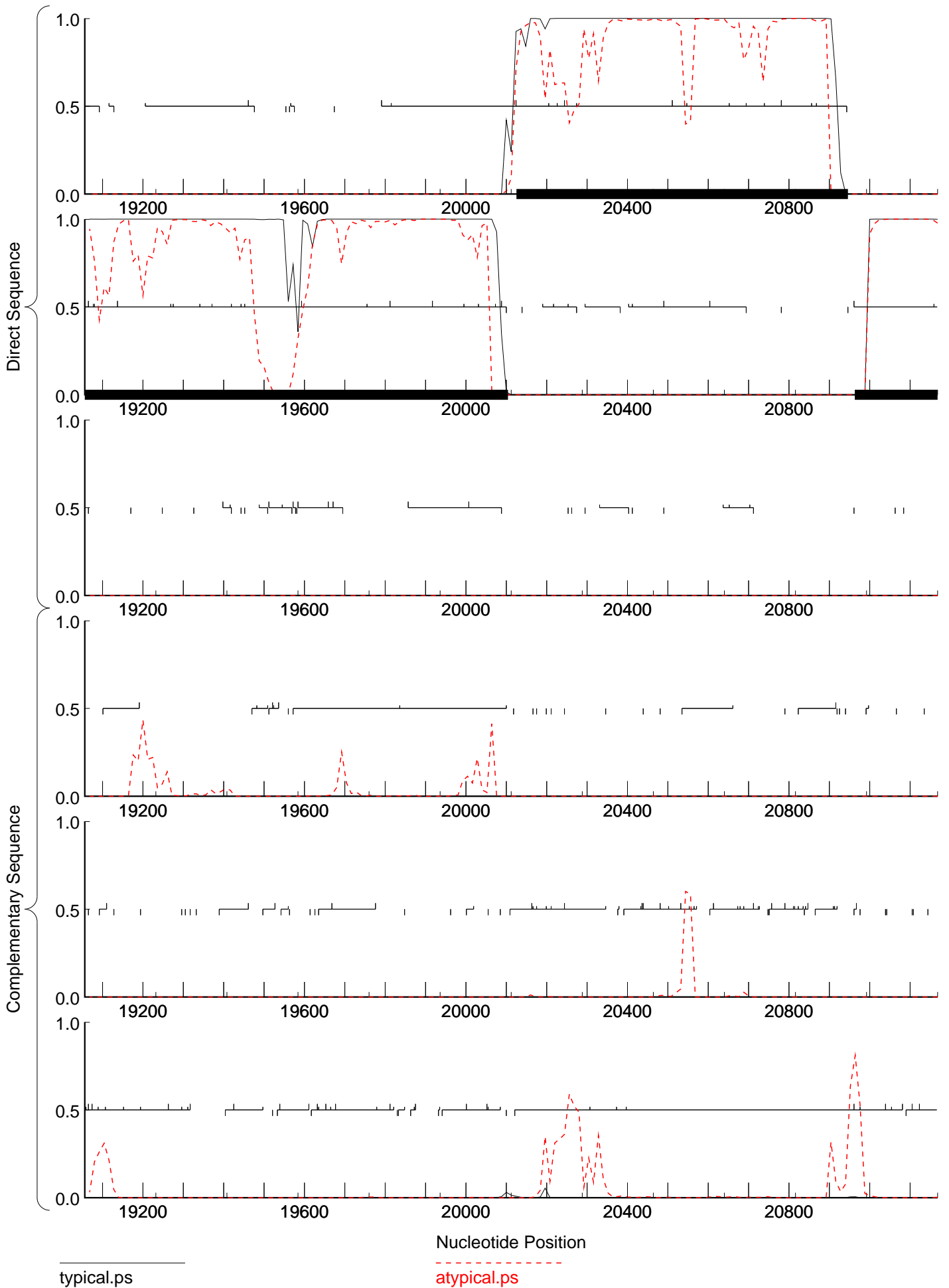


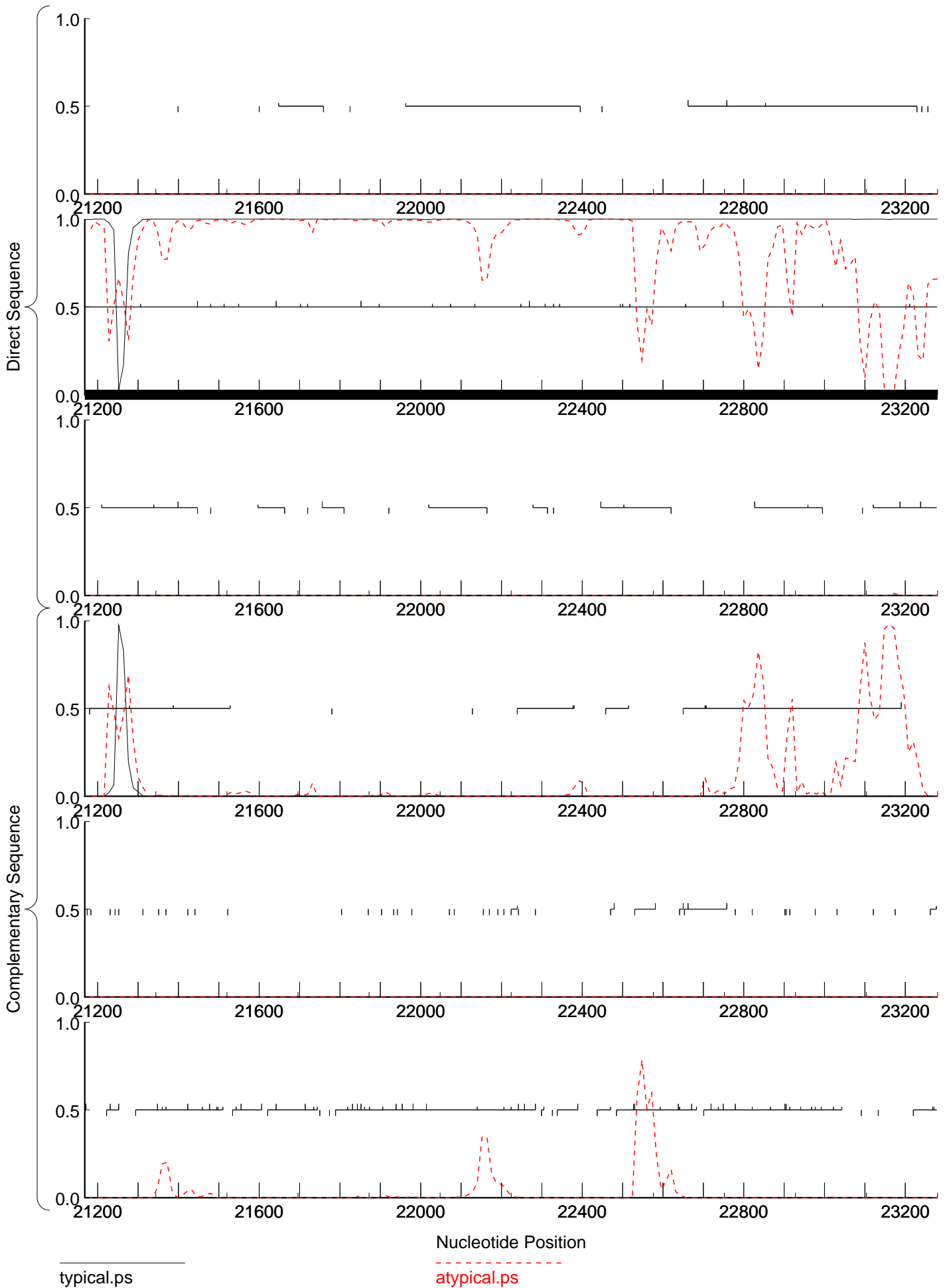


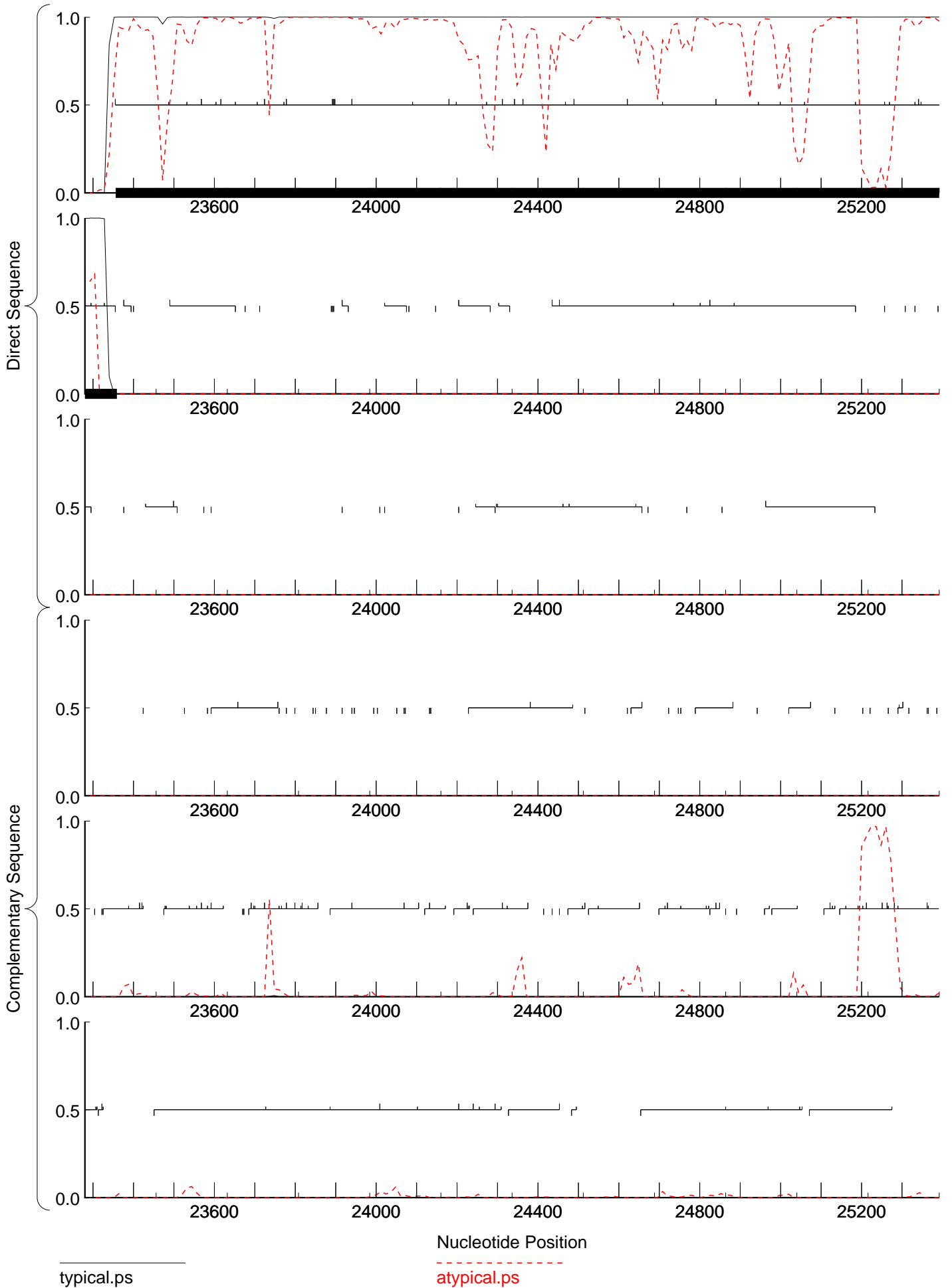
typical.ps

Nucleotide Position

atypical.ps

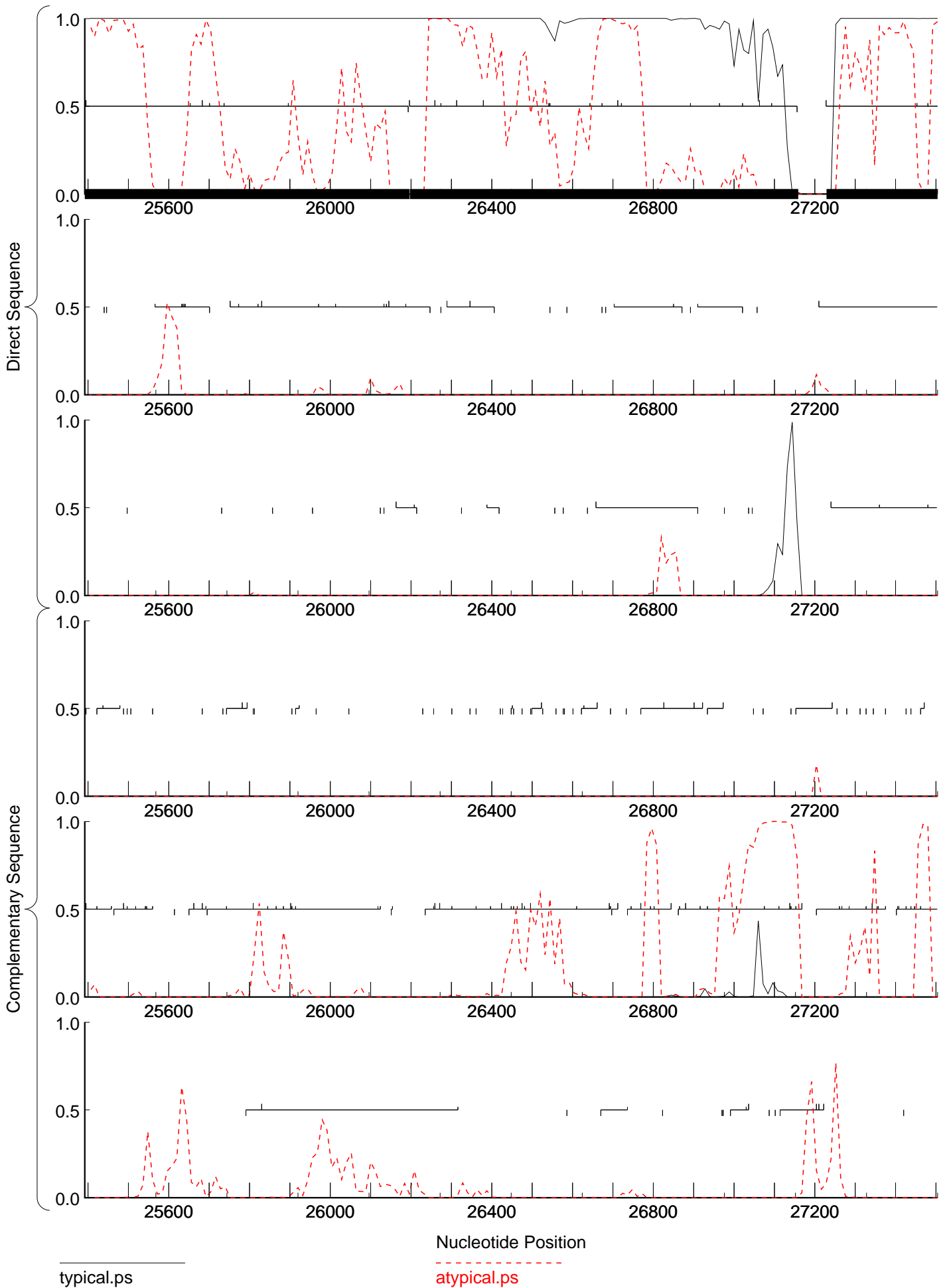


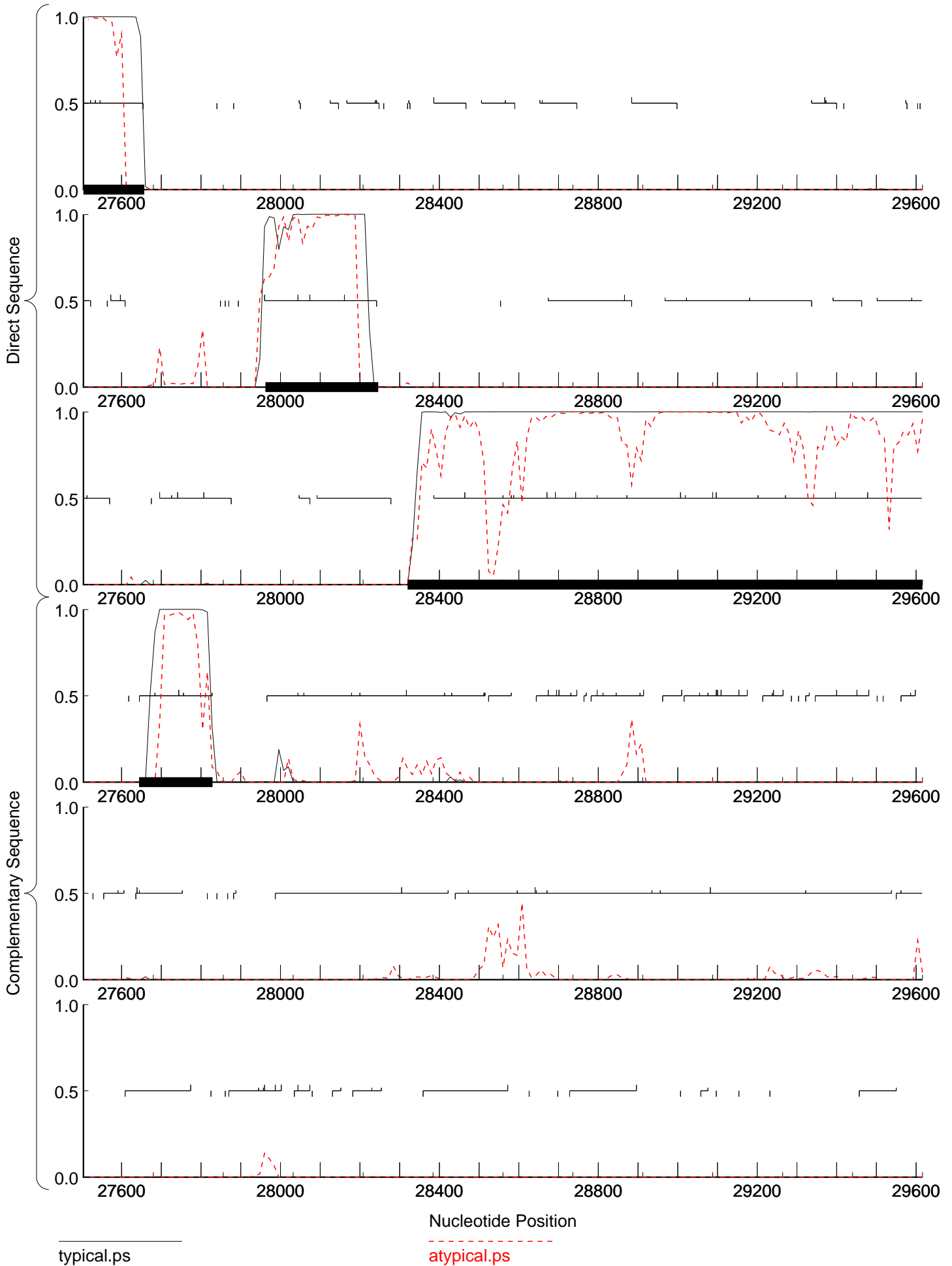


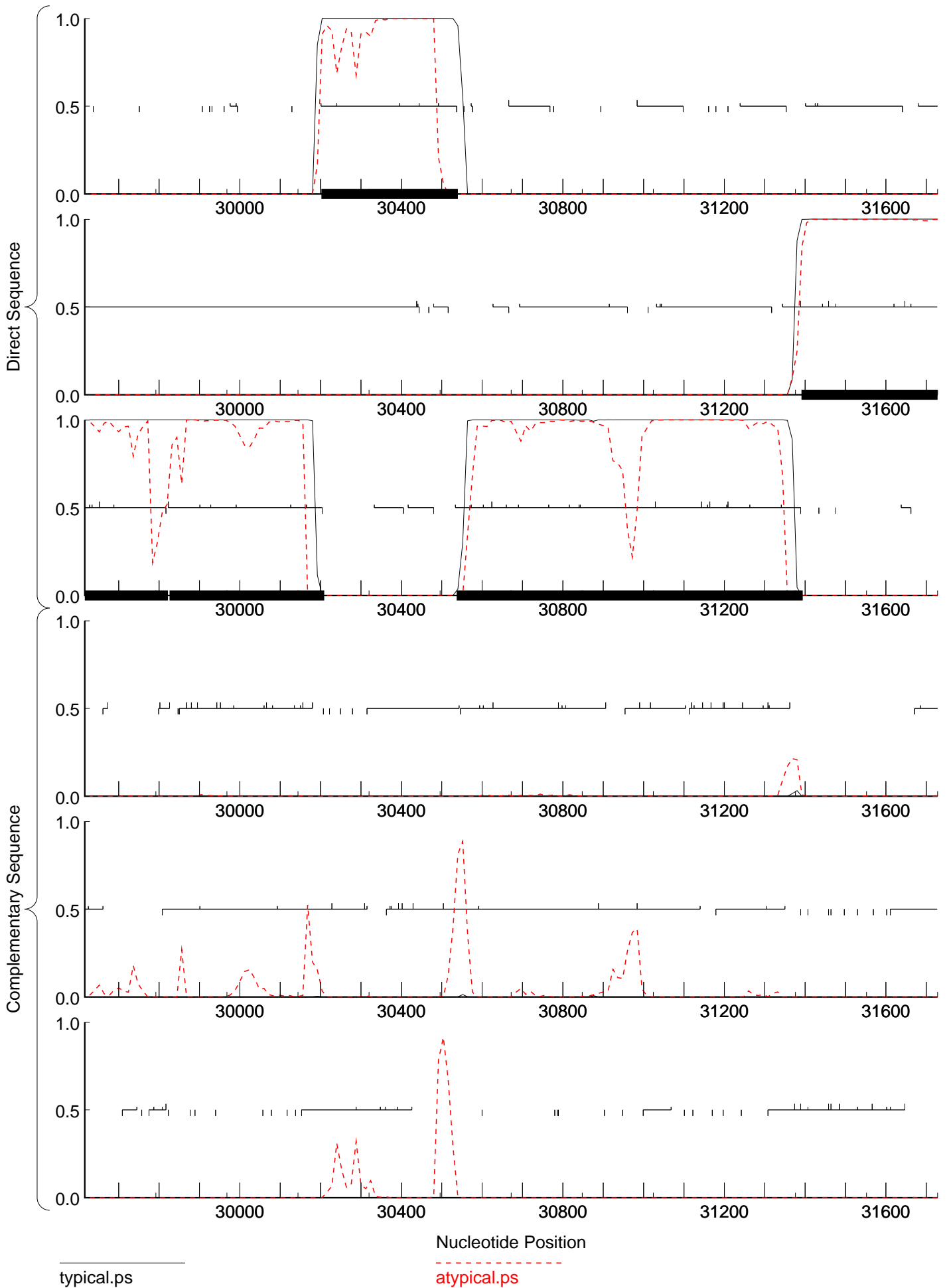


typical.ps

atypical.ps

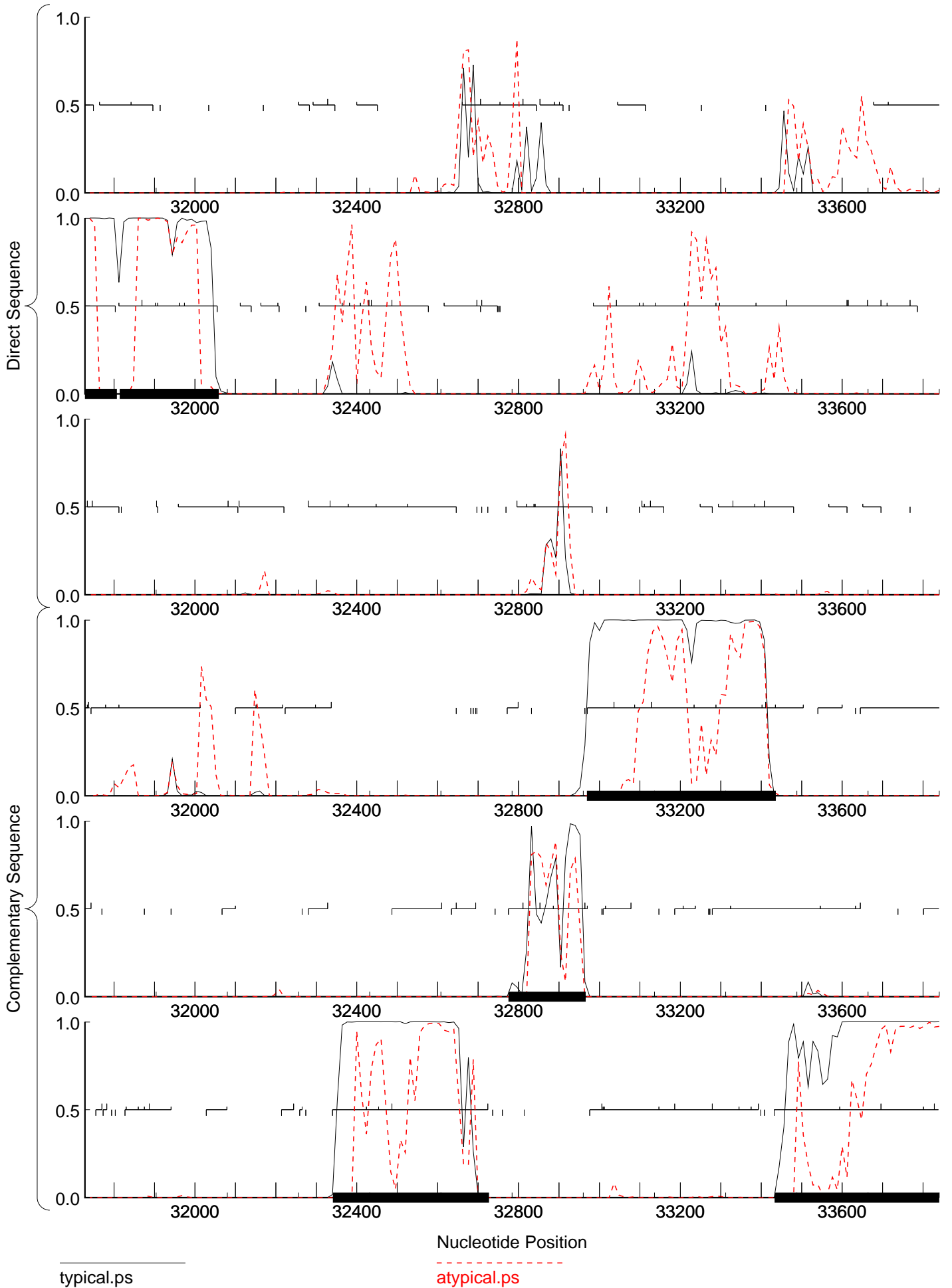






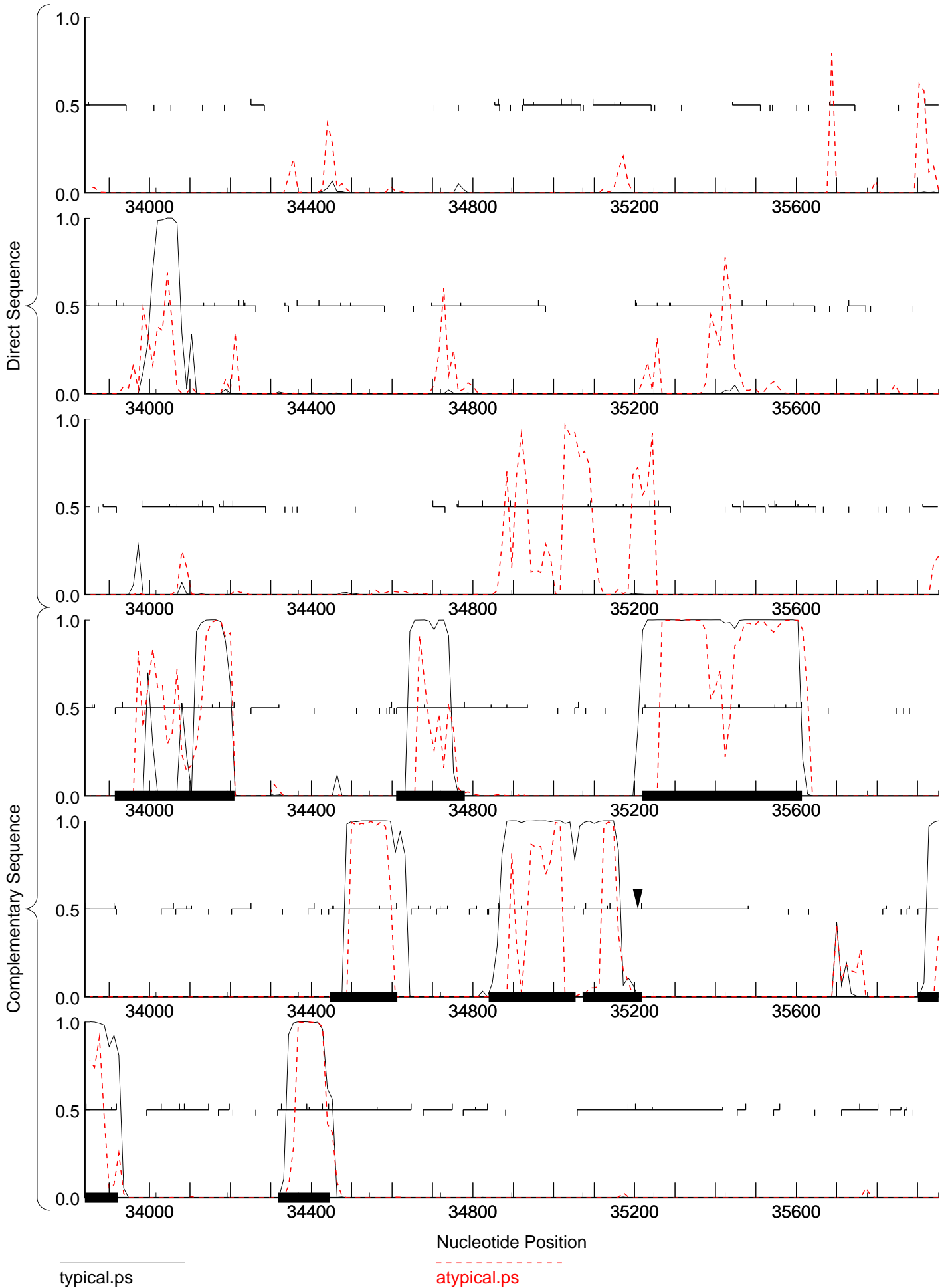
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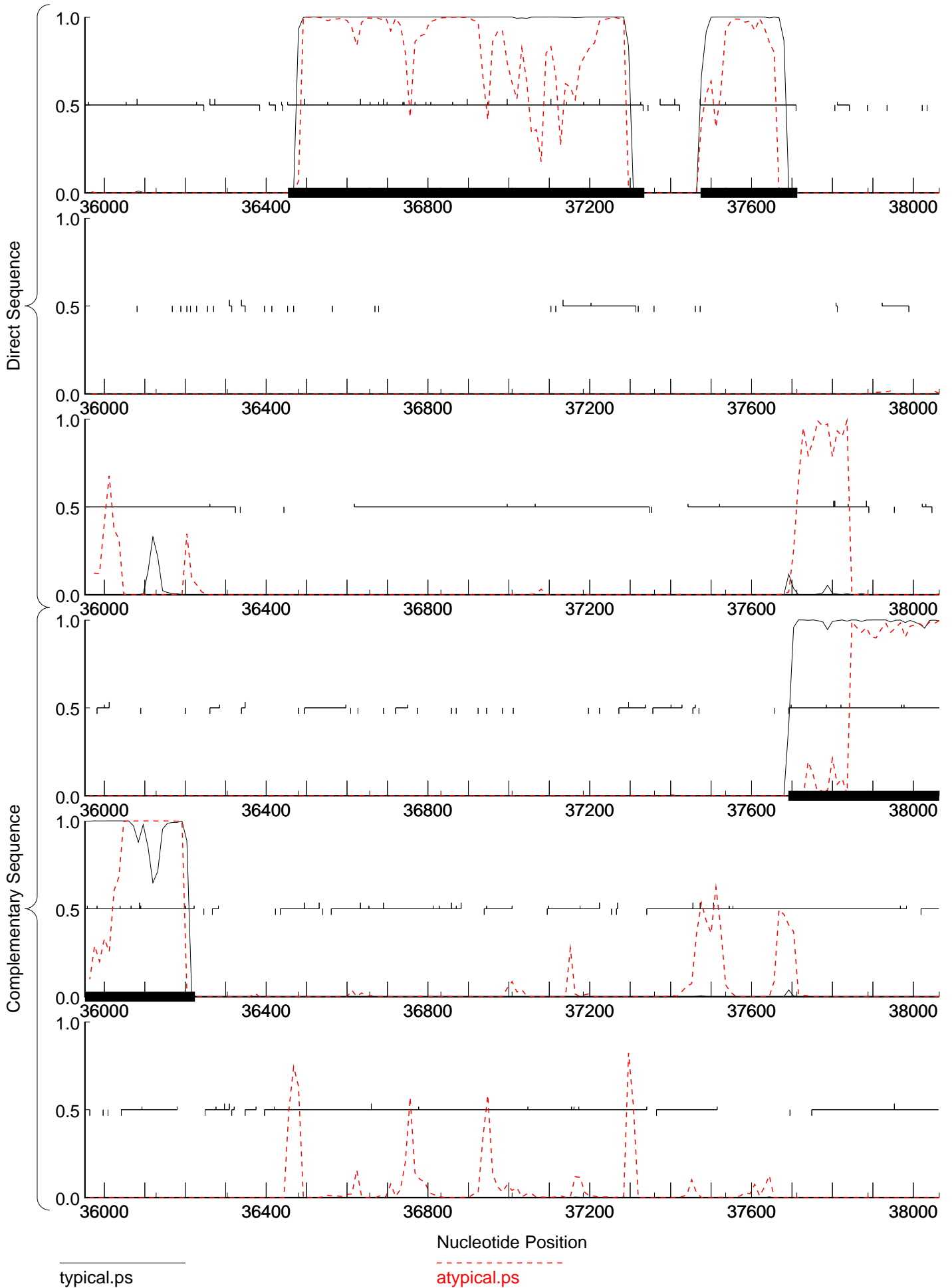
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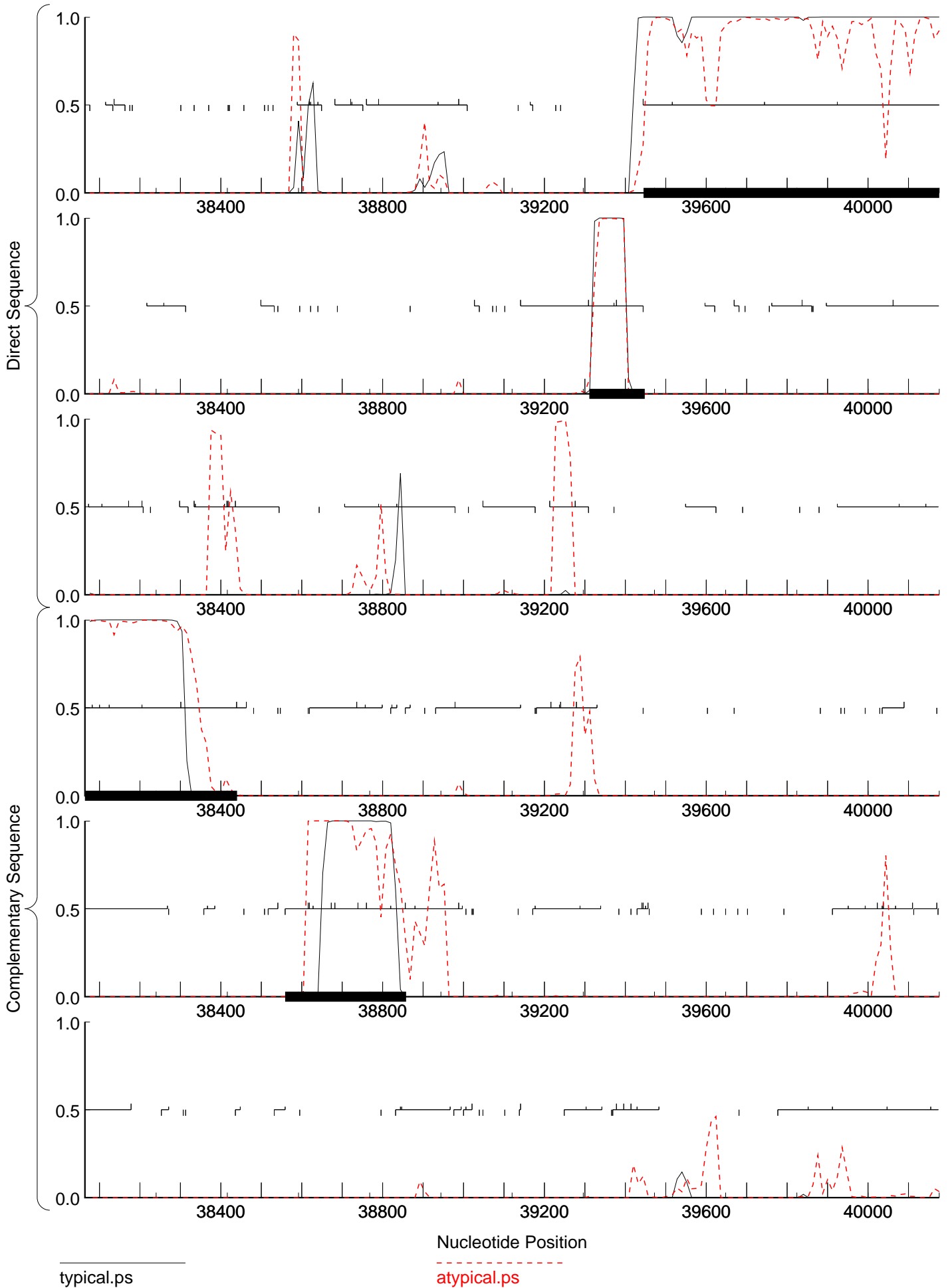


typical.ps

atypical.ps

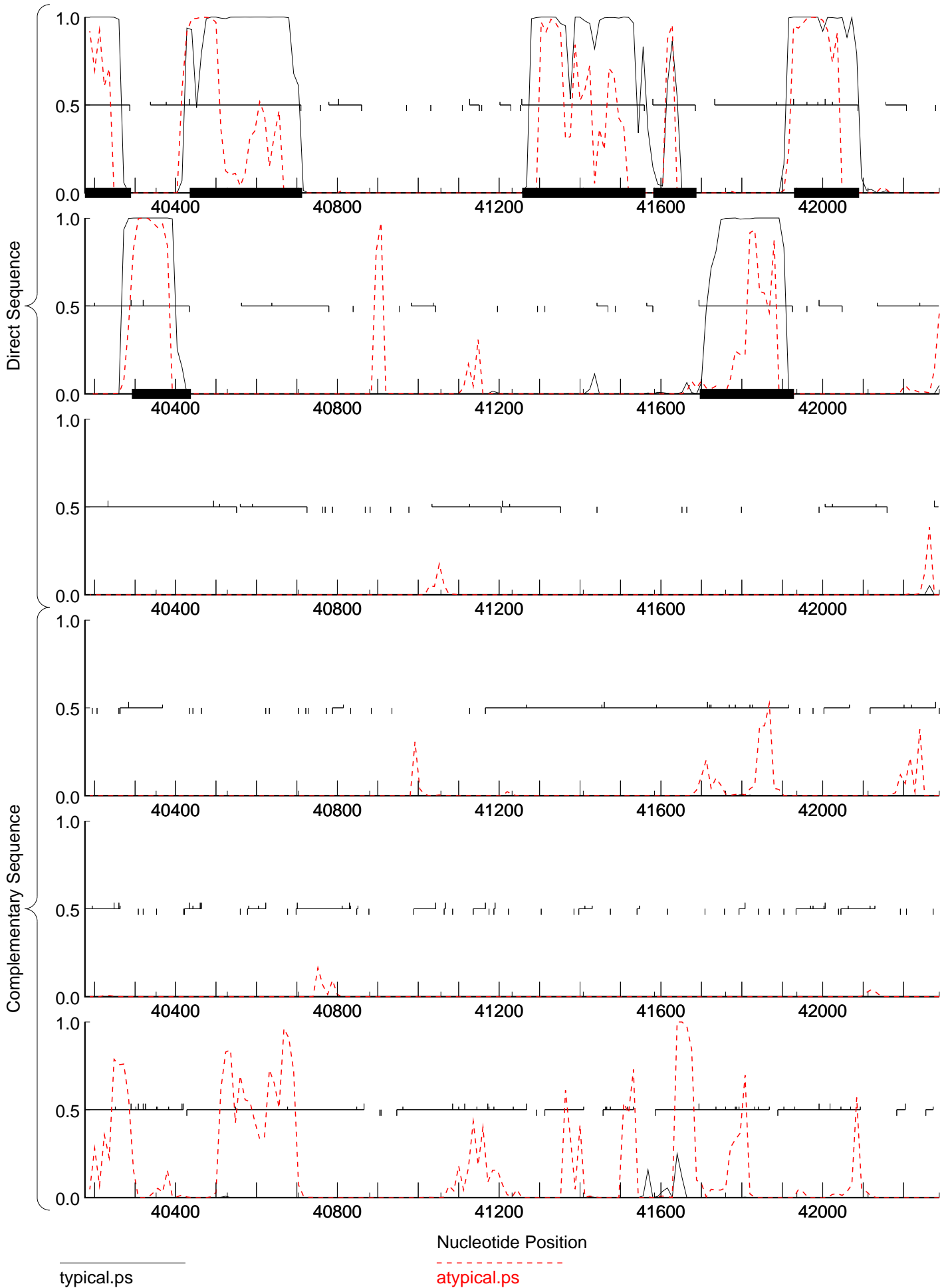


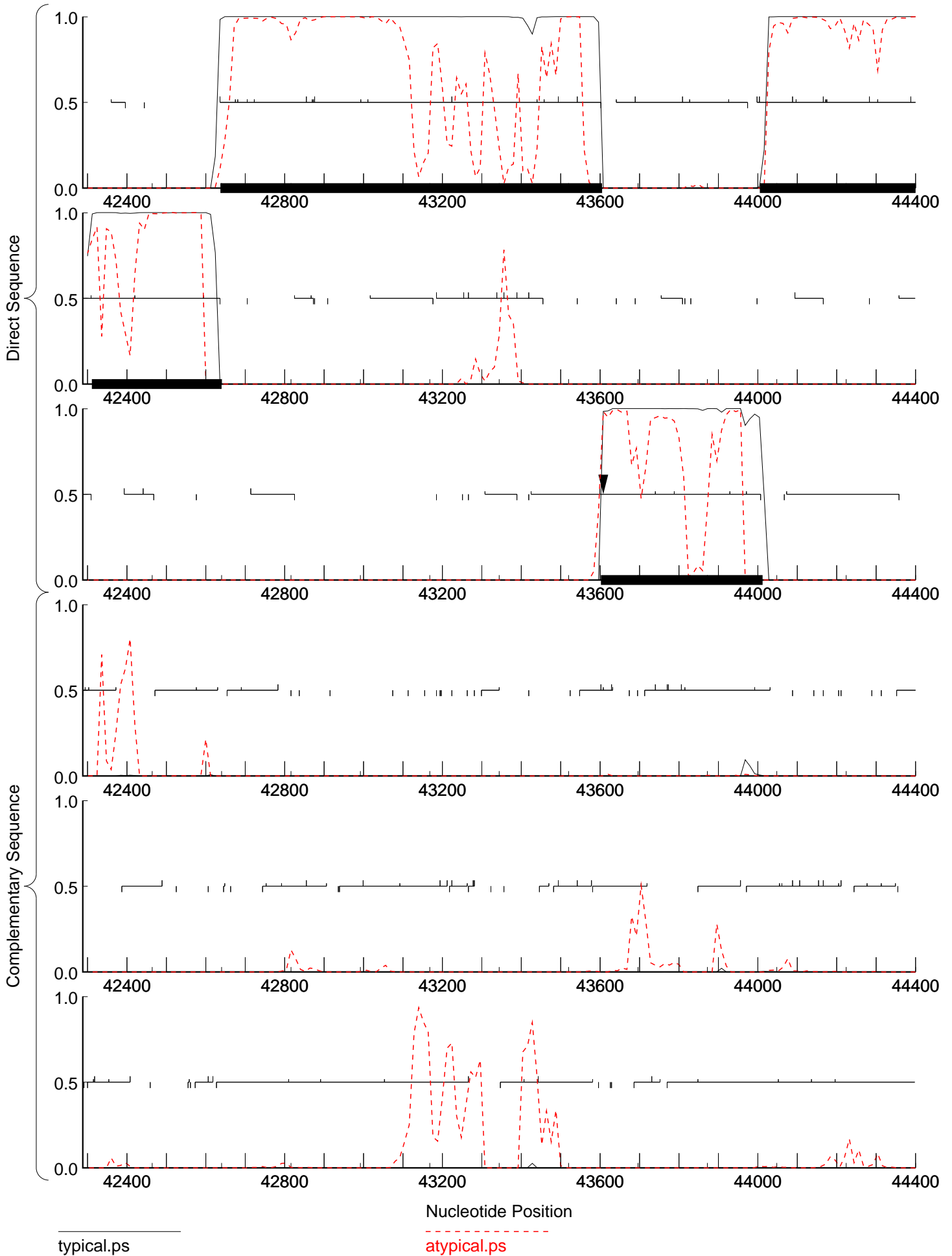


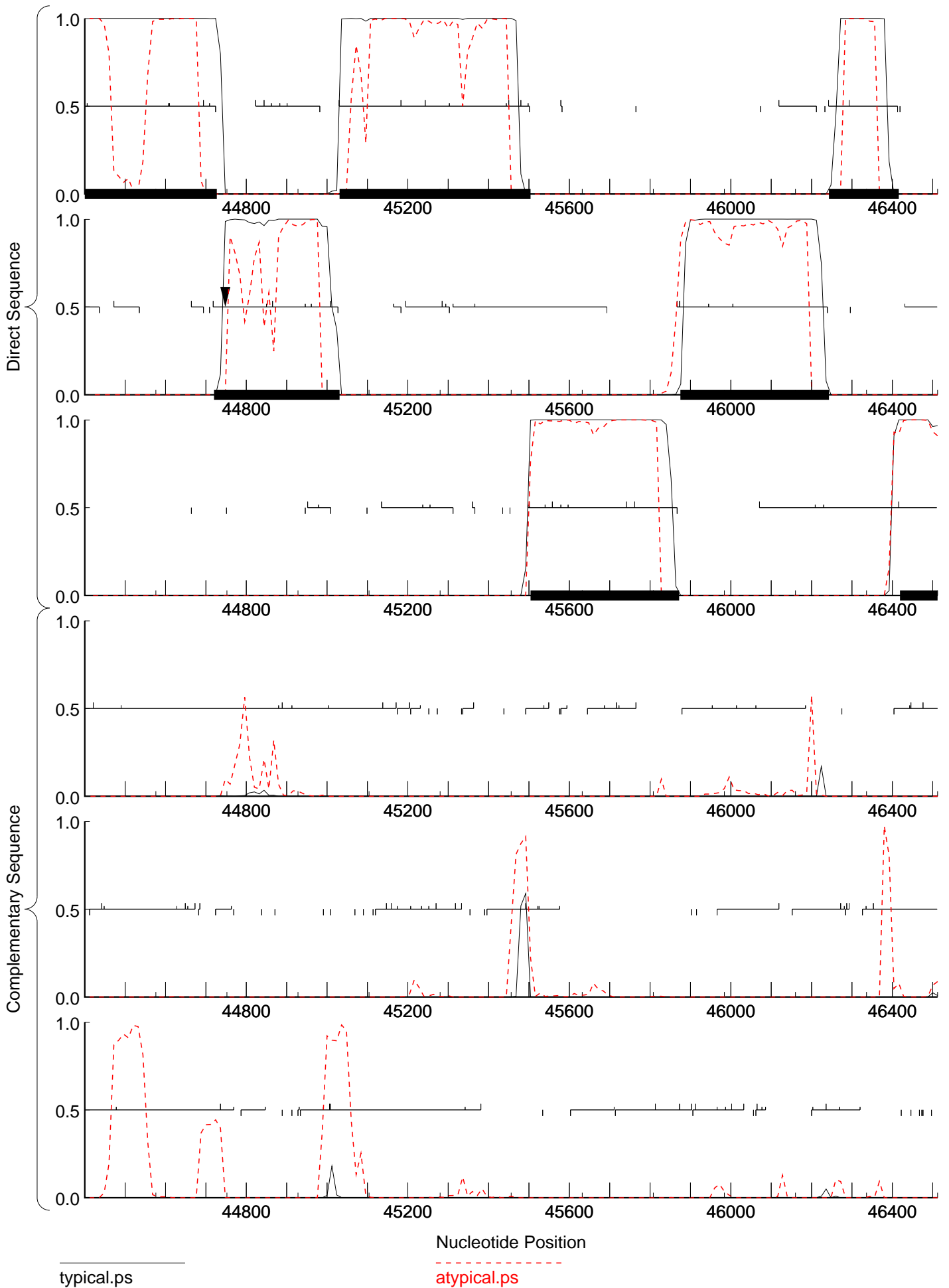


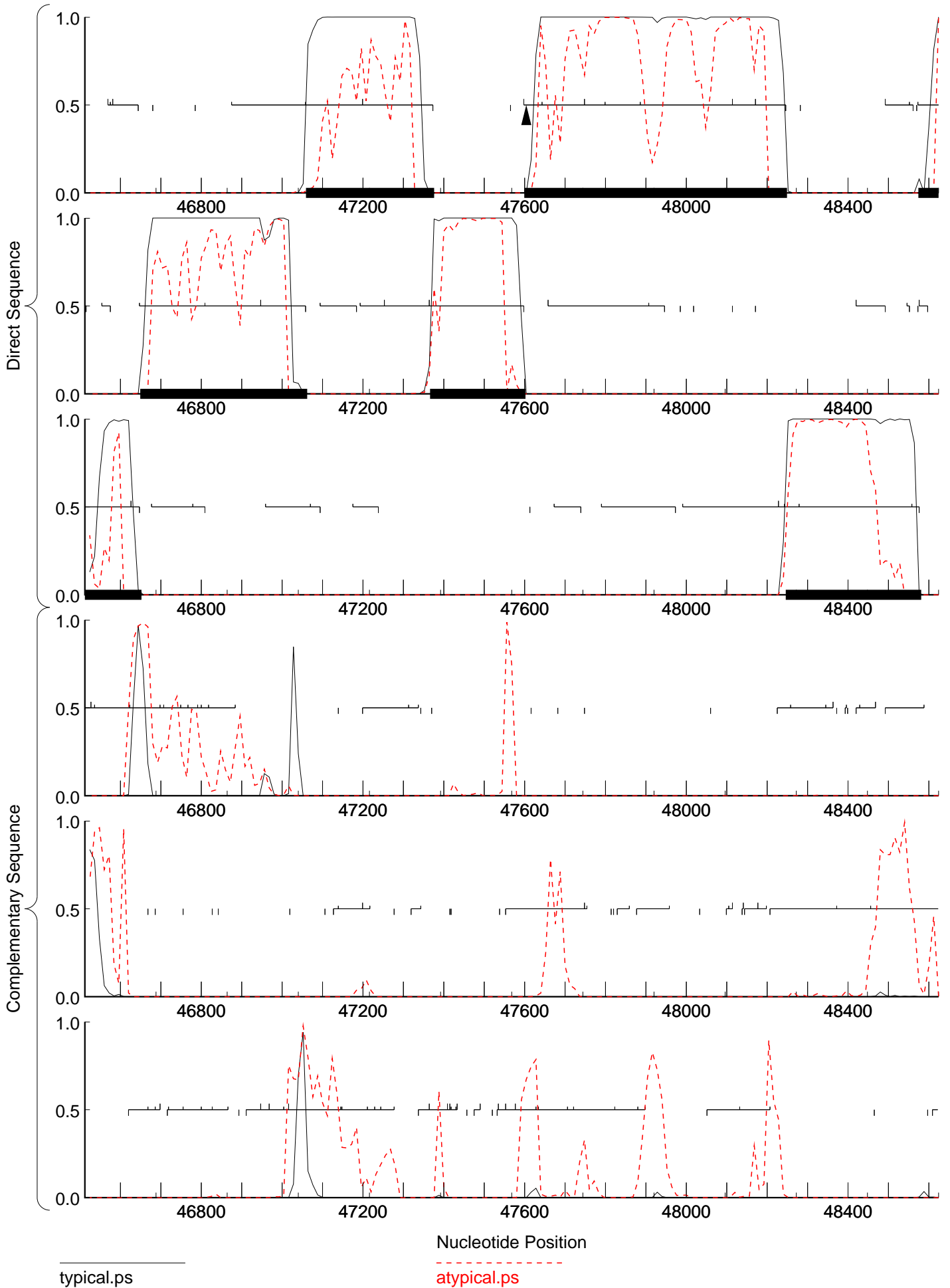
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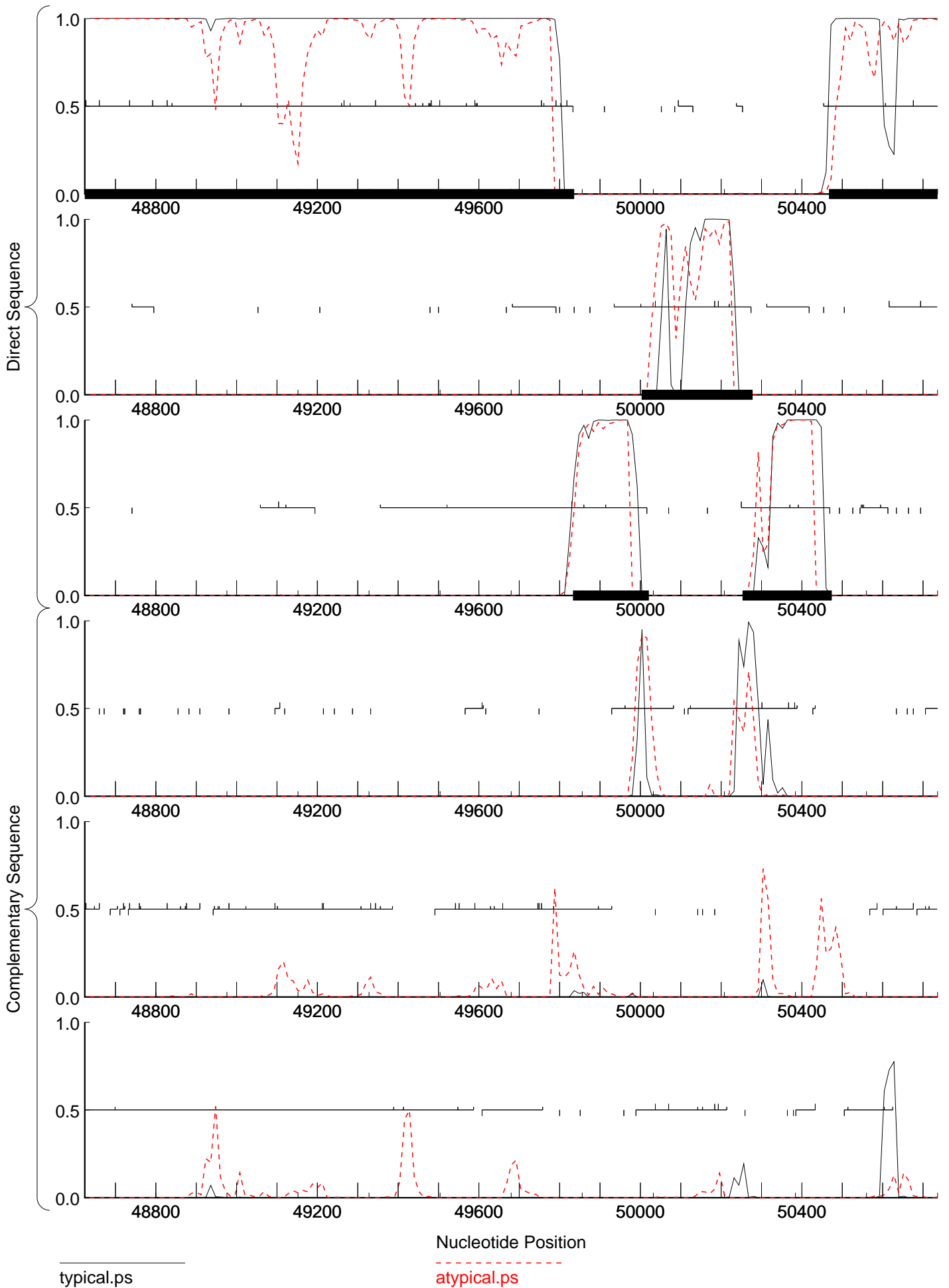
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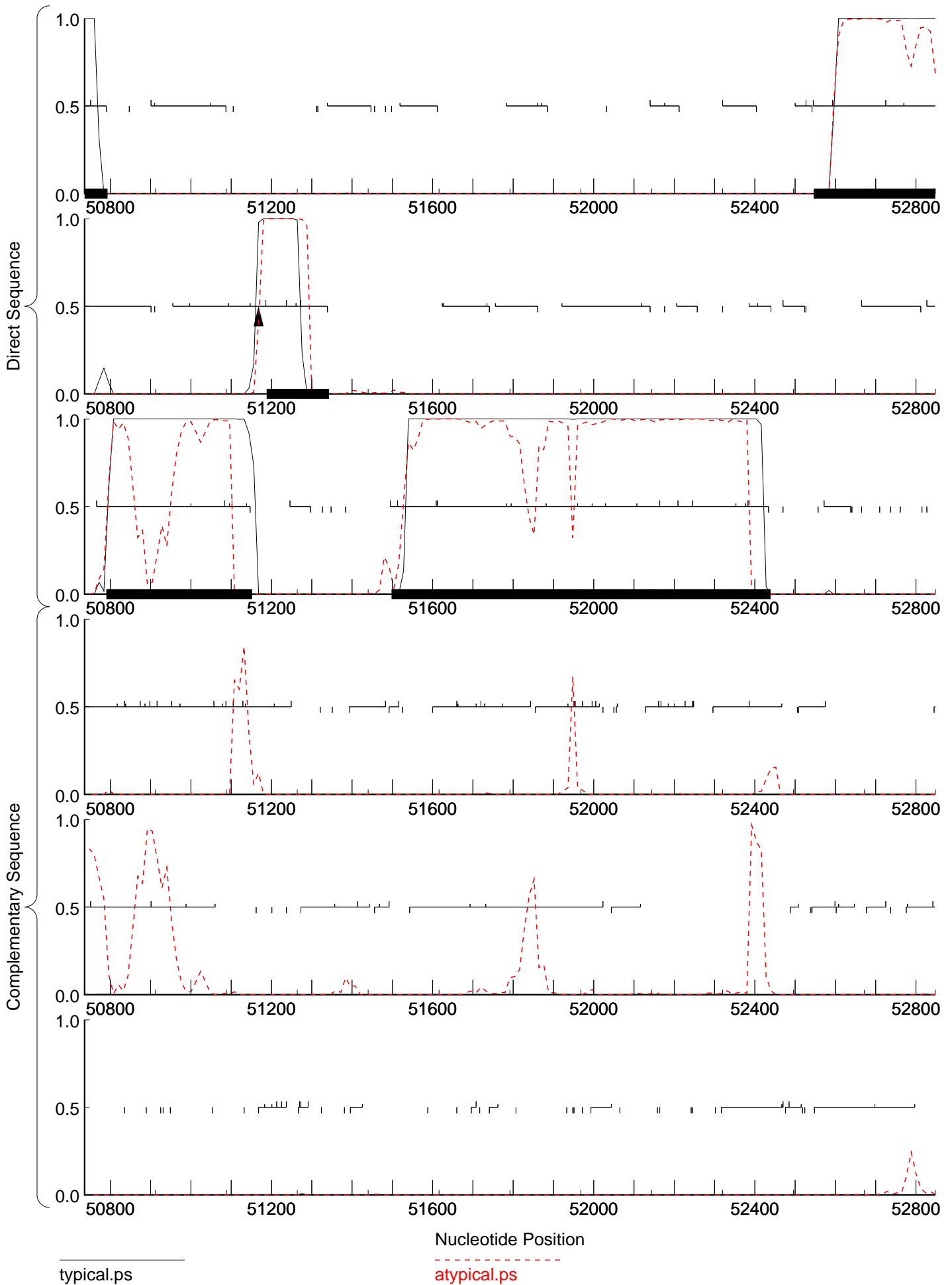


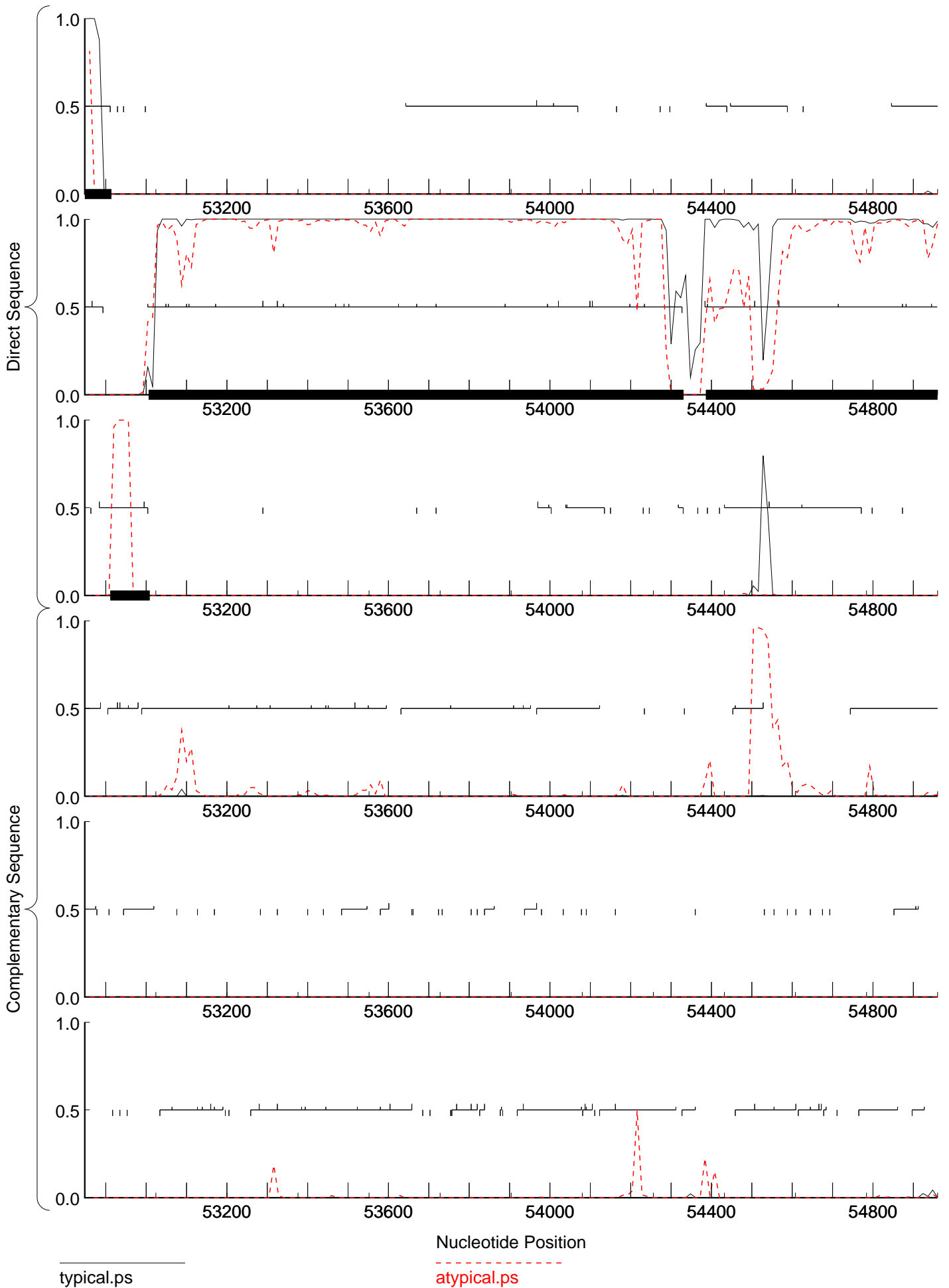


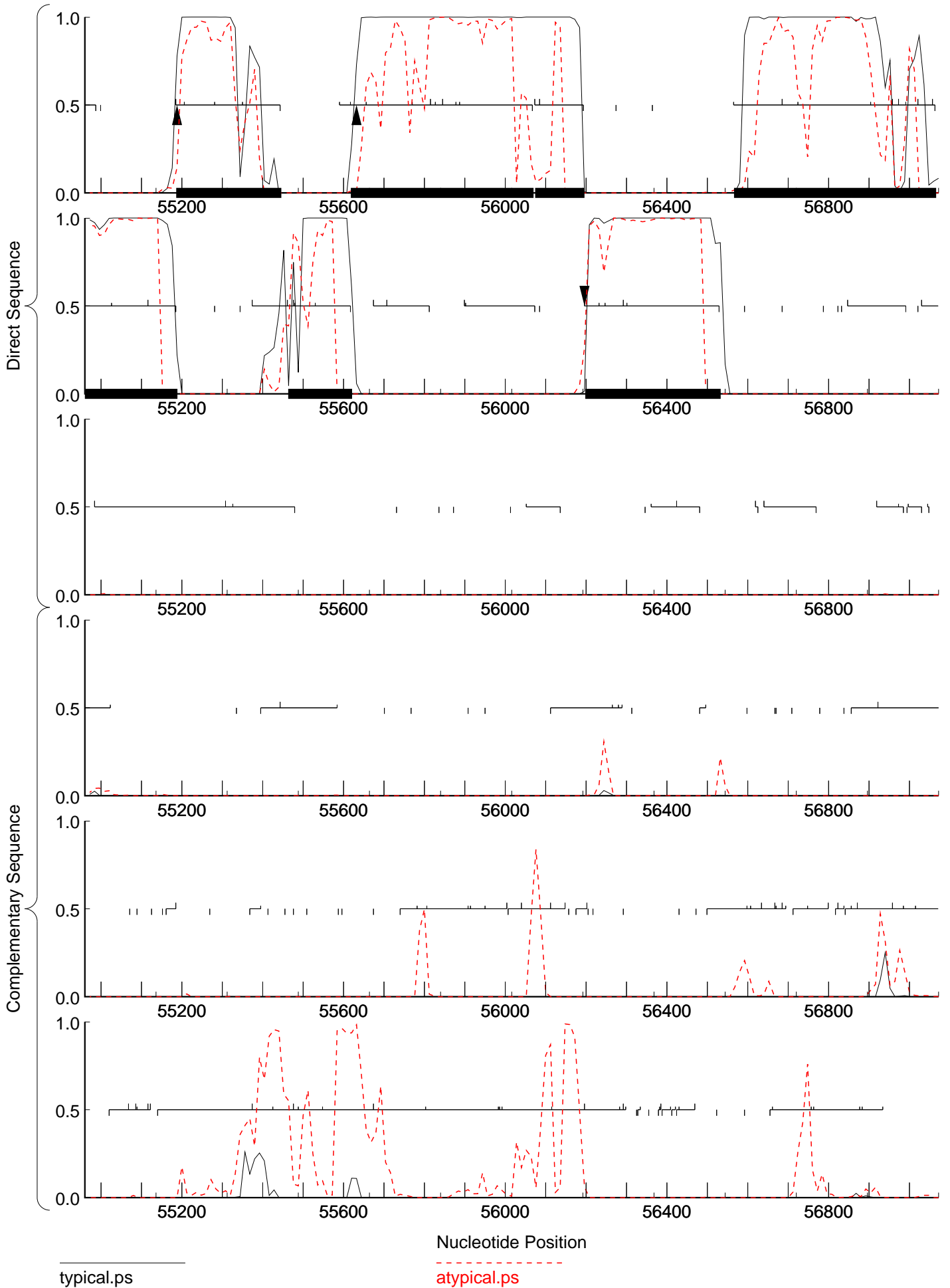


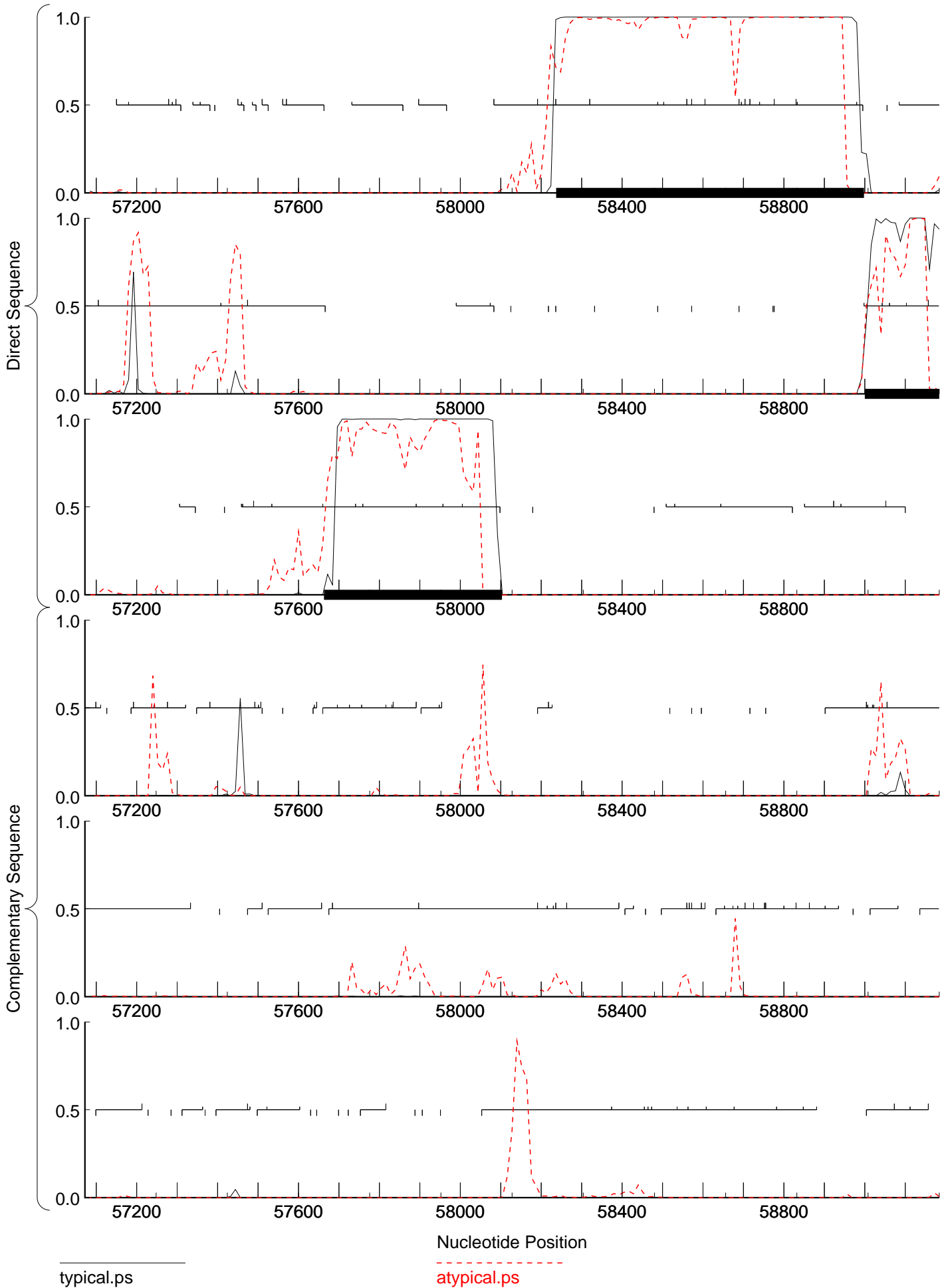


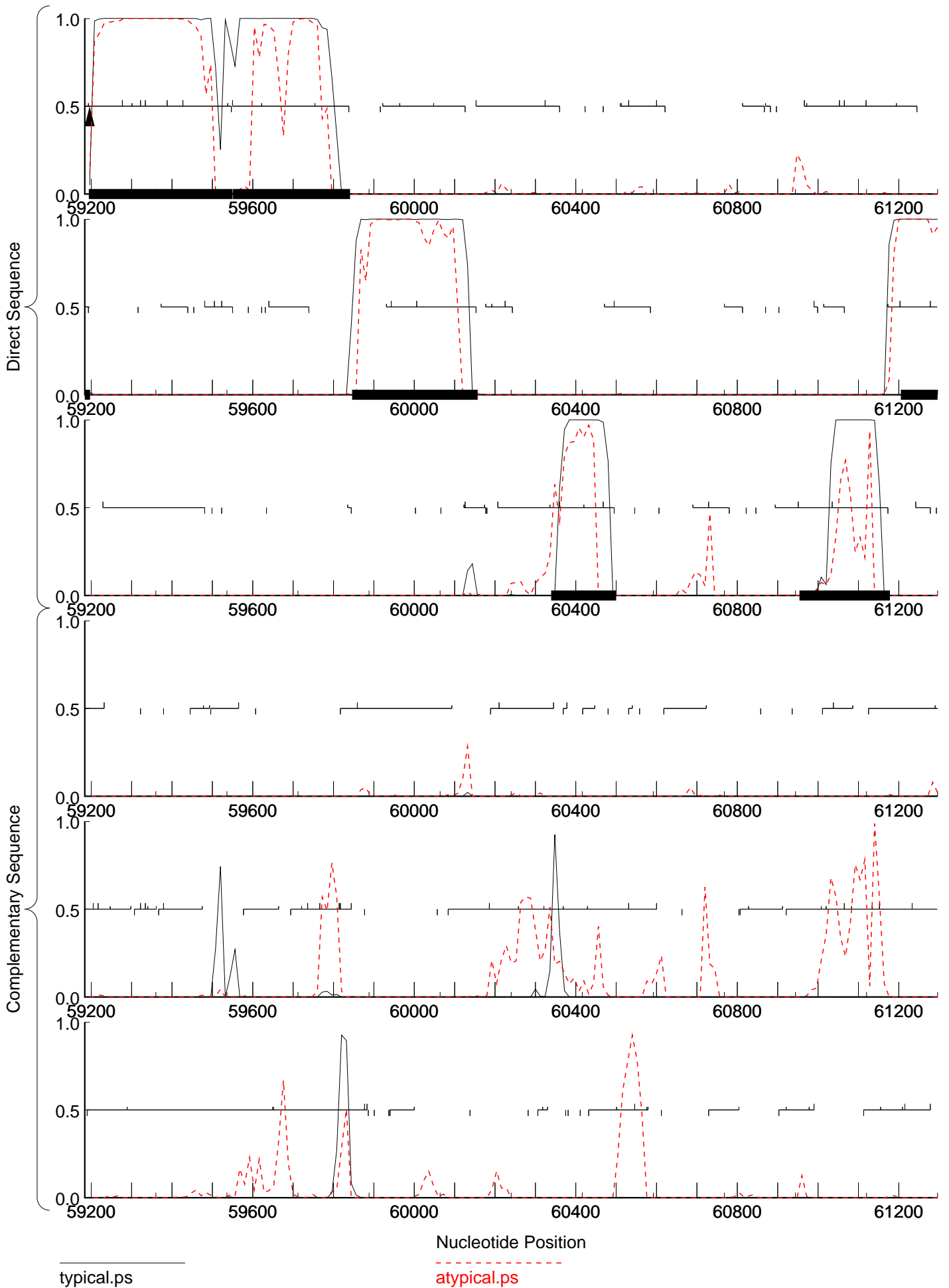


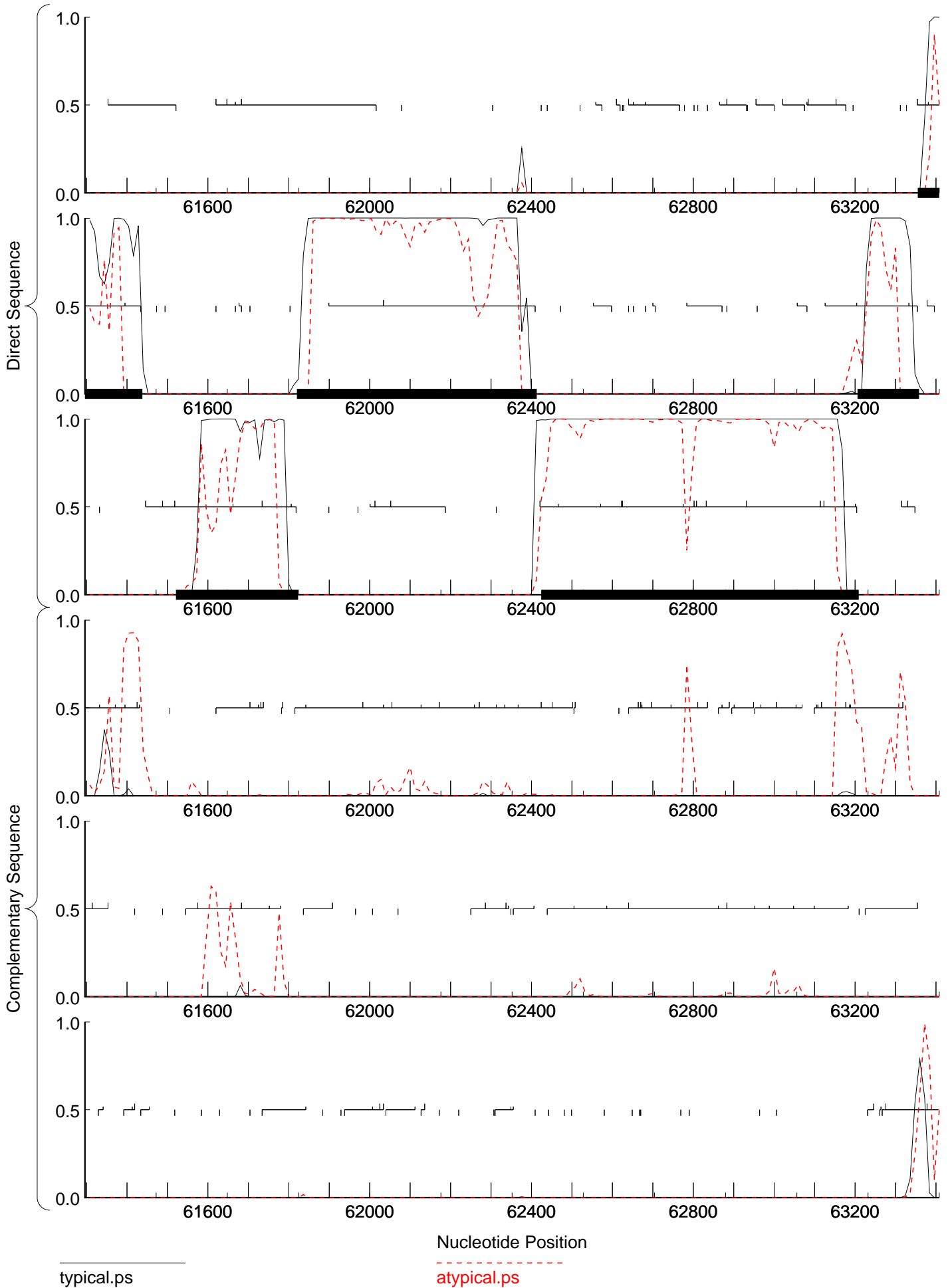


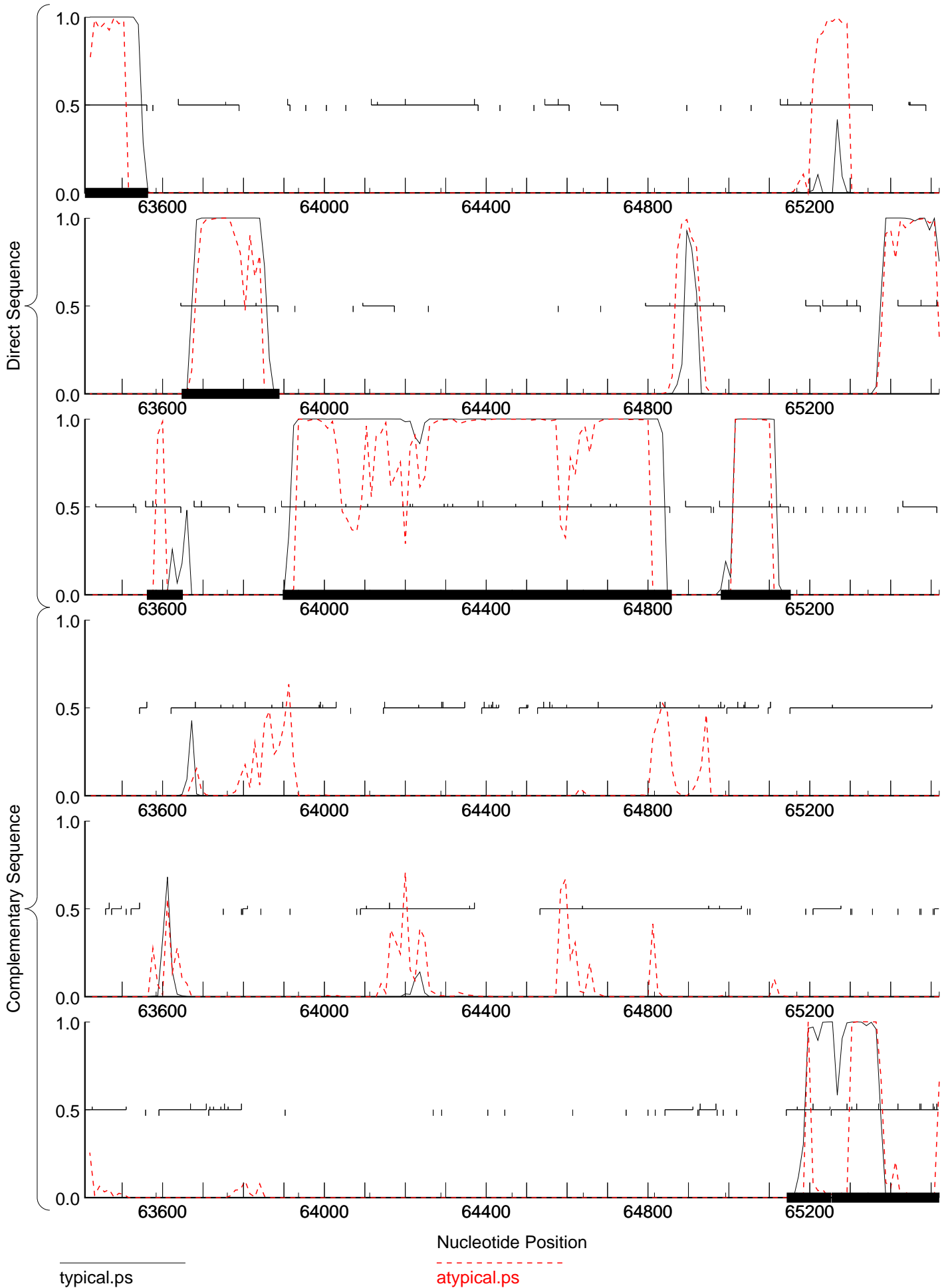


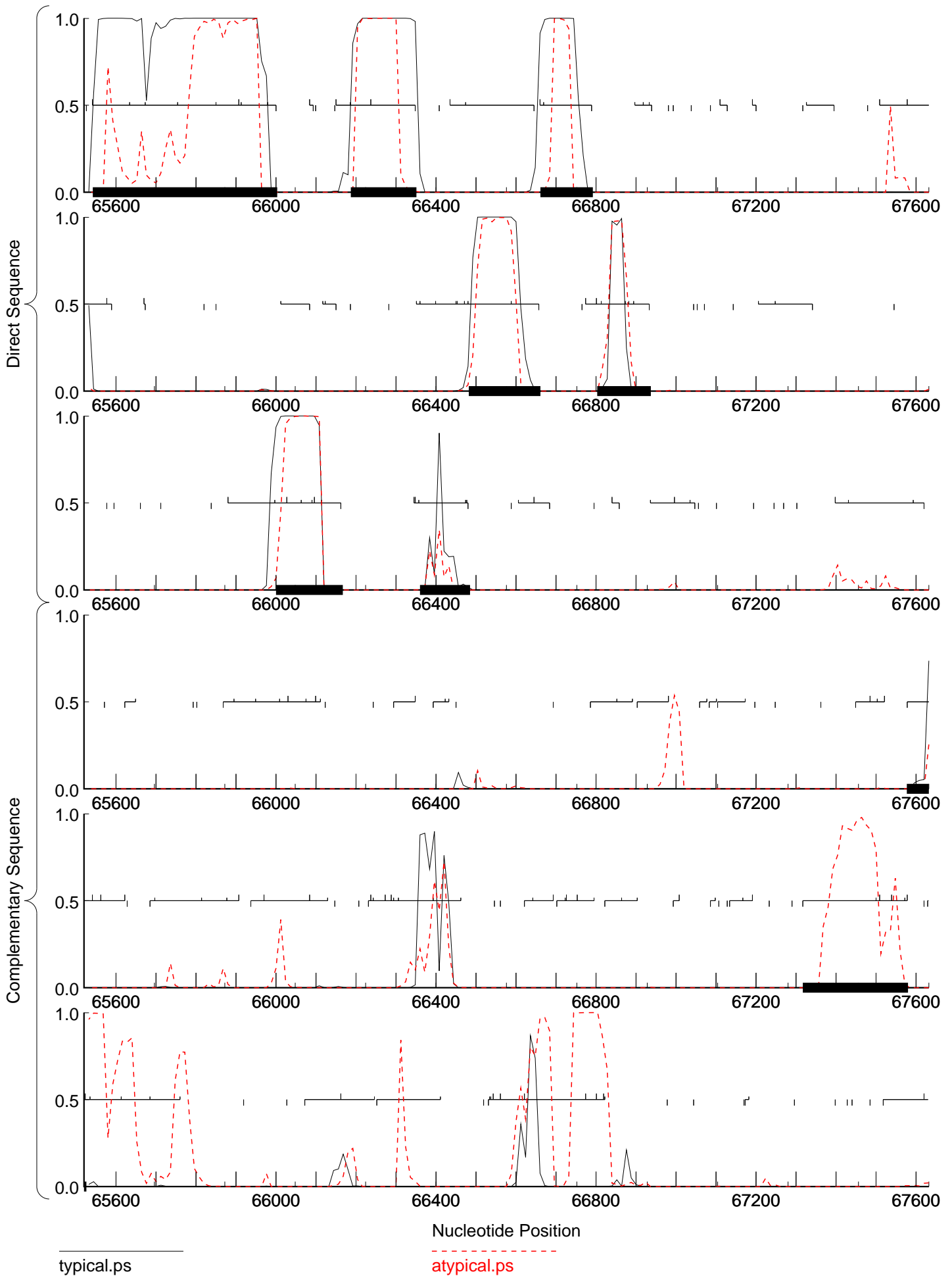


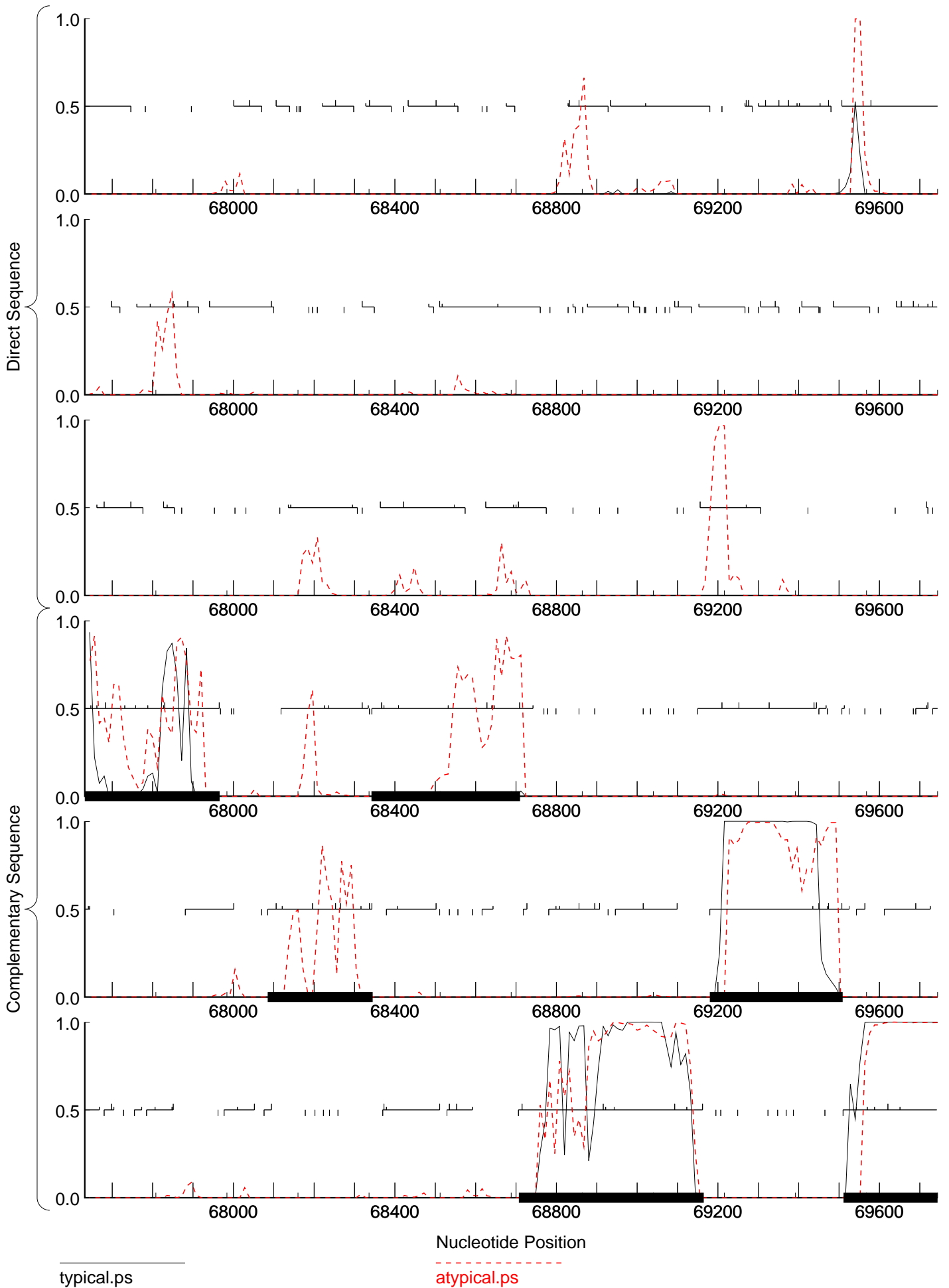


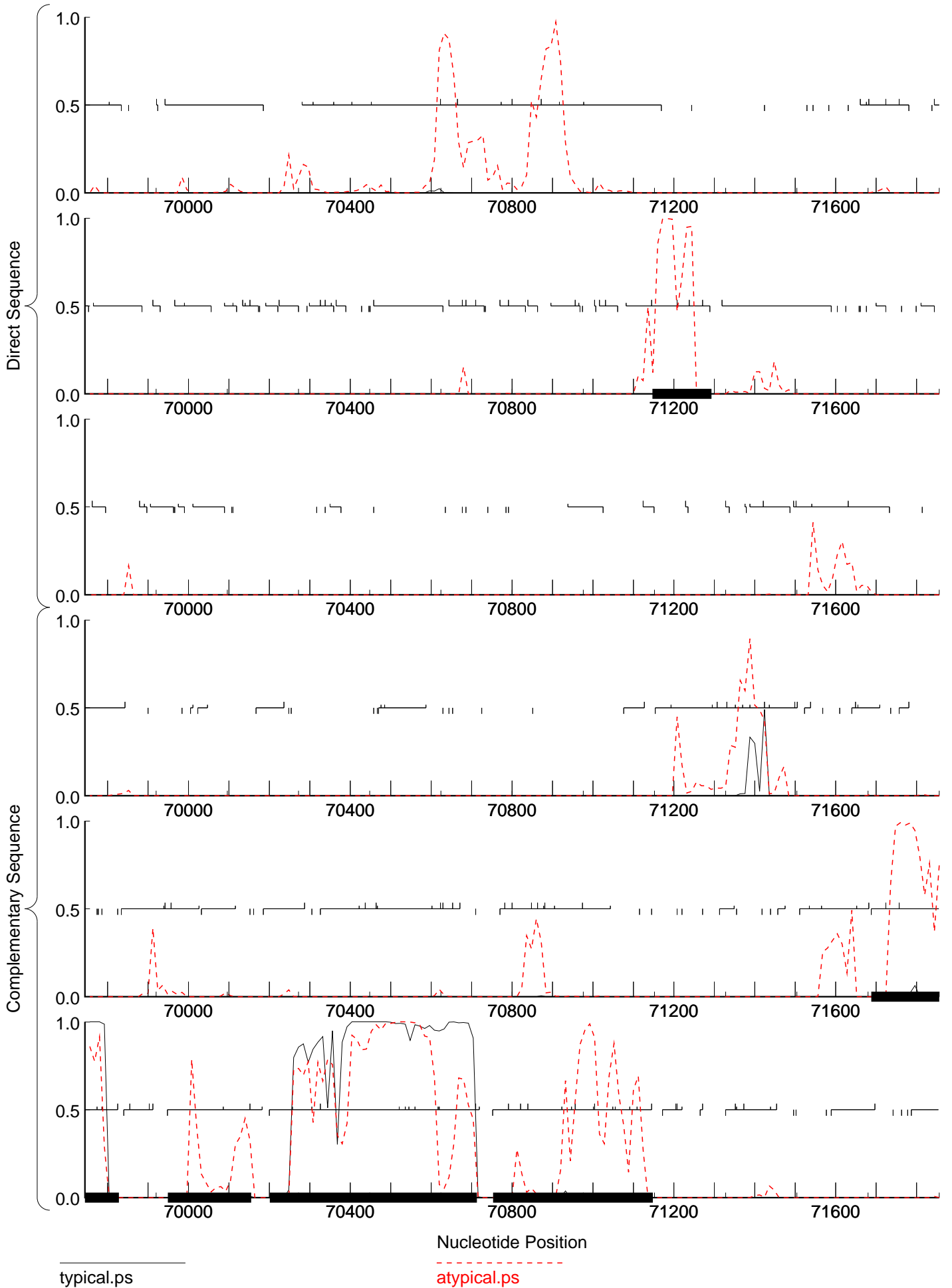


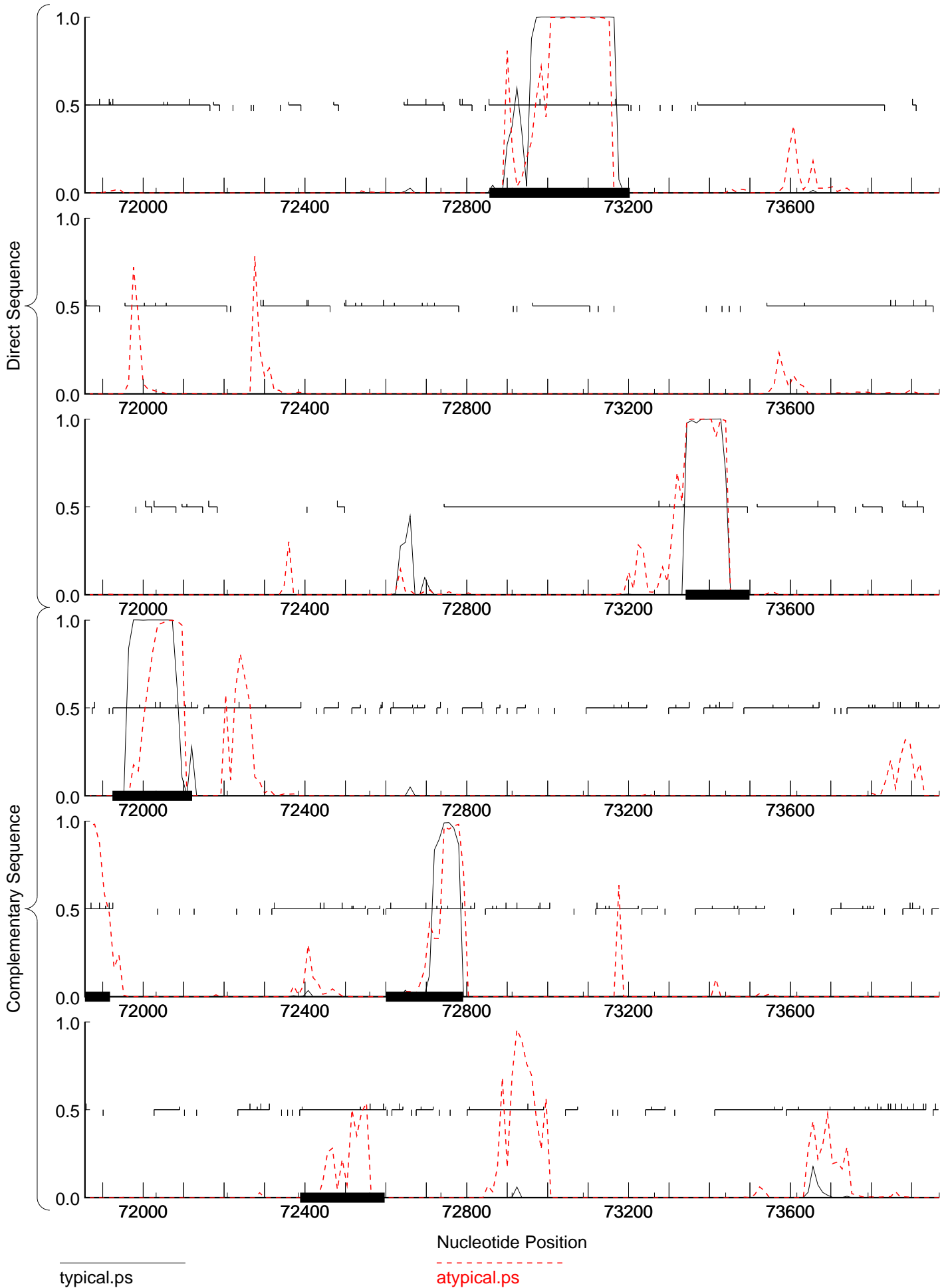












typical.ps

atypical.ps

