

GeneMark

Version 2.5p (09.08.06)

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PROGRAM INFORMATION

Sequence : Mycobacterium phage Sibs6 complete sequence, 50210 bp including 10-base 3' overhang (CGGATG
Analysis Date : 8/24/17 at 0:57:30
Pages : 25
Sequence Length : 50210 bp
GC Content : 63.77%

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,

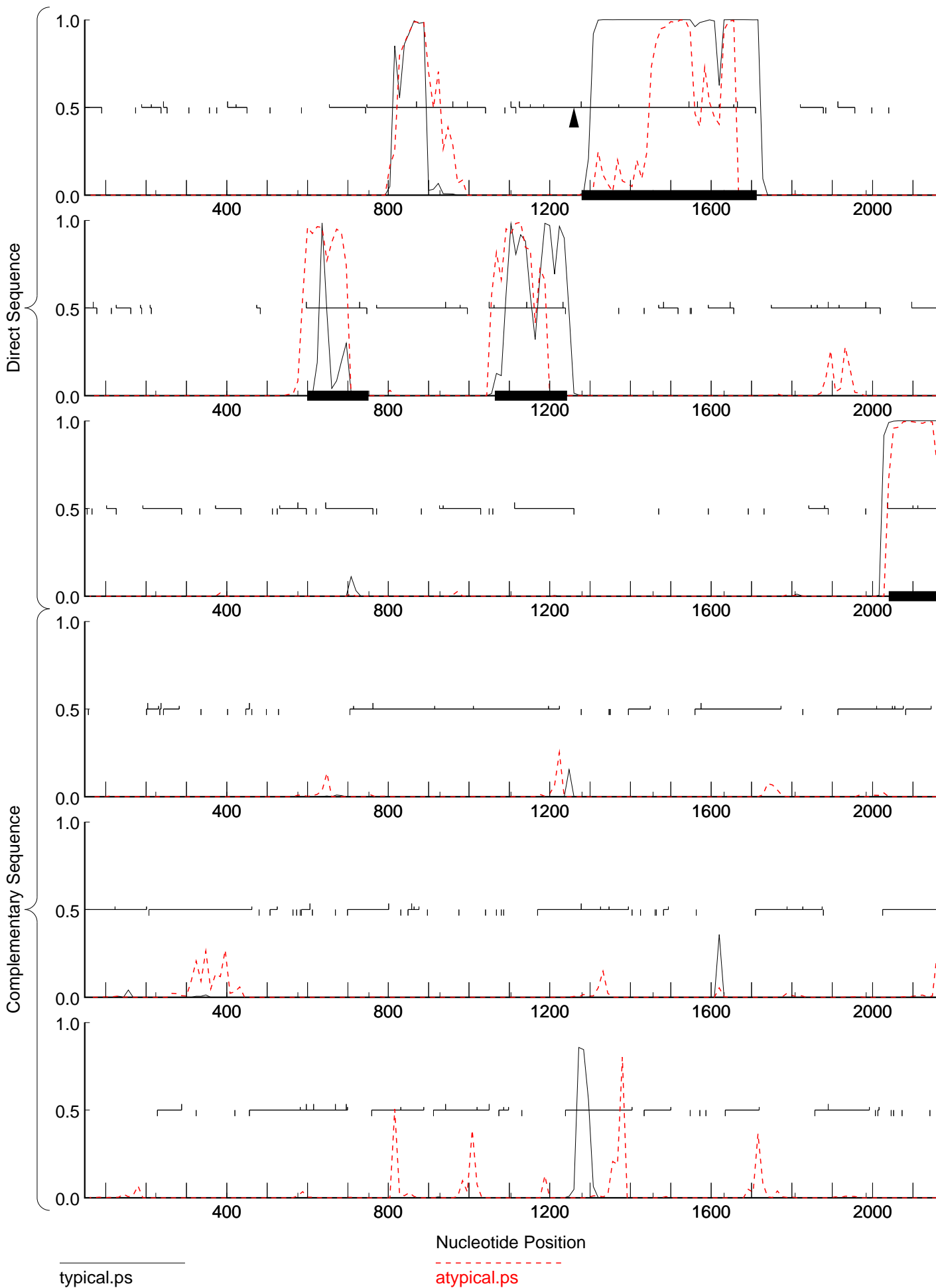
Matrix notes & comments

Training set derived by GeneMarkS, 4.30 October 2014
Thu Aug 24 00:57:30 2017

MATRIX INFORMATION

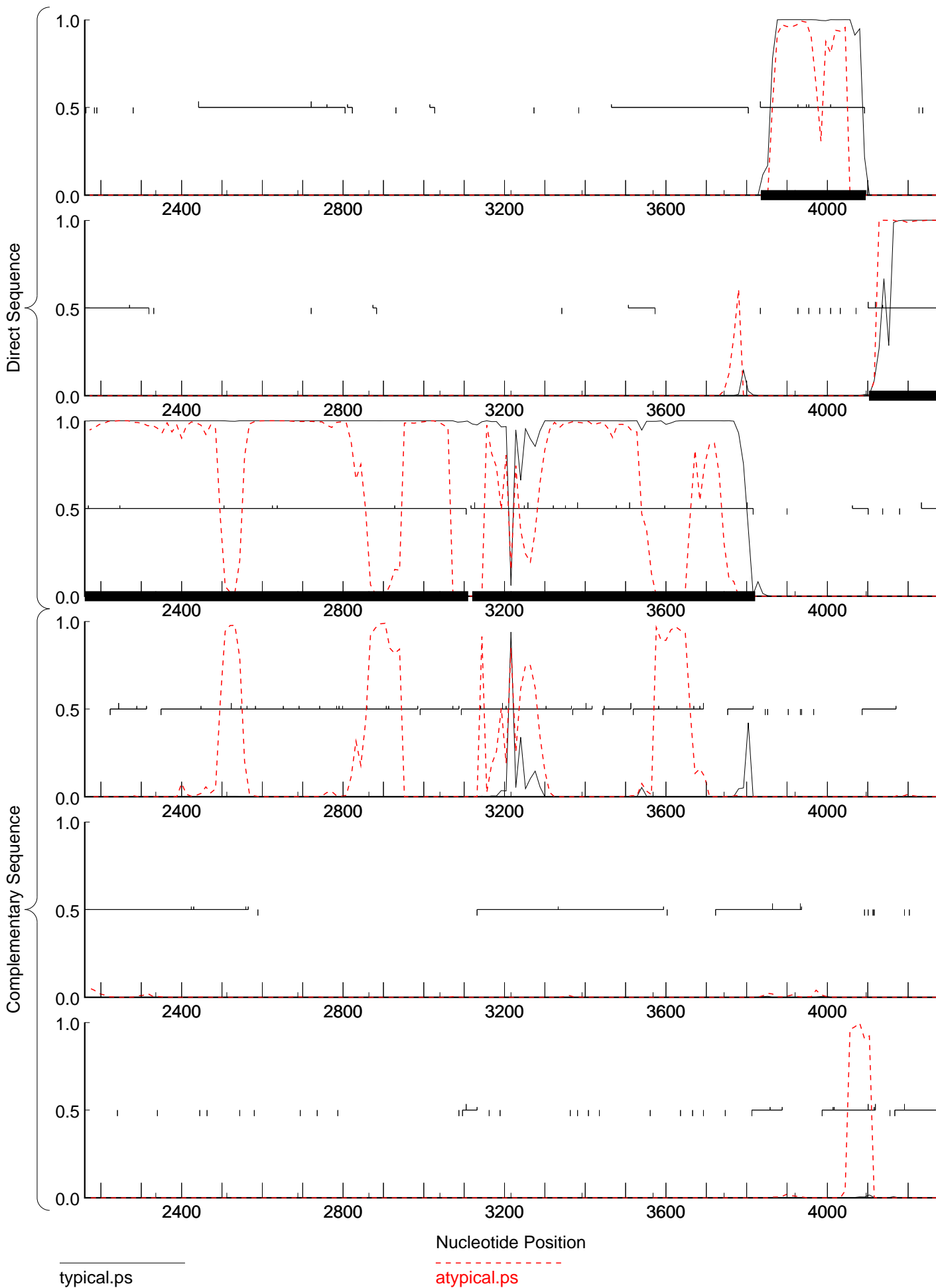
Matrix : -
Author : -
Order : 4

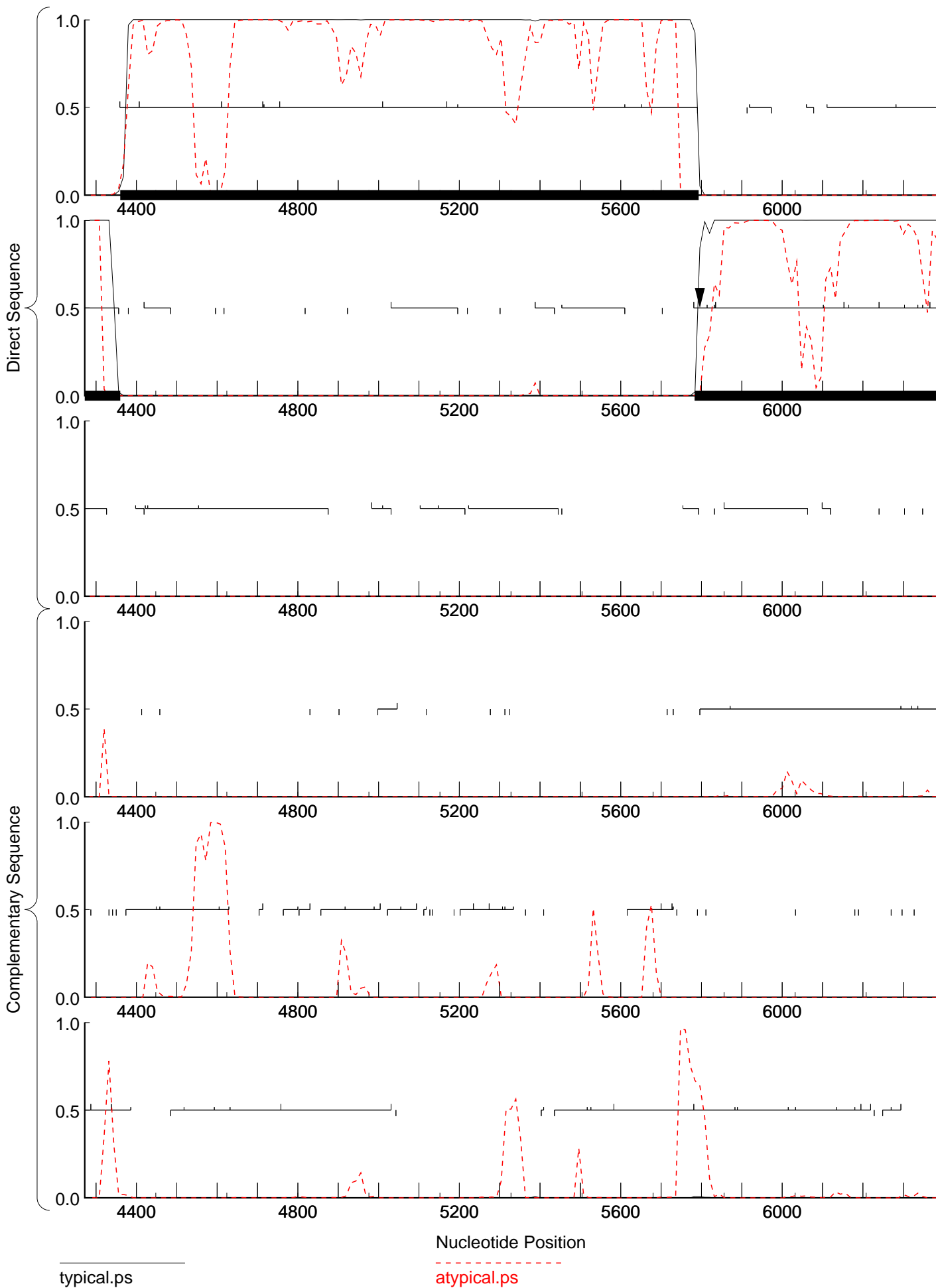
Send questions / comments to:
Dr. M. Borodovsky
Georgia Institute of Technology
School of Biology
Atlanta, GA 30332-0230

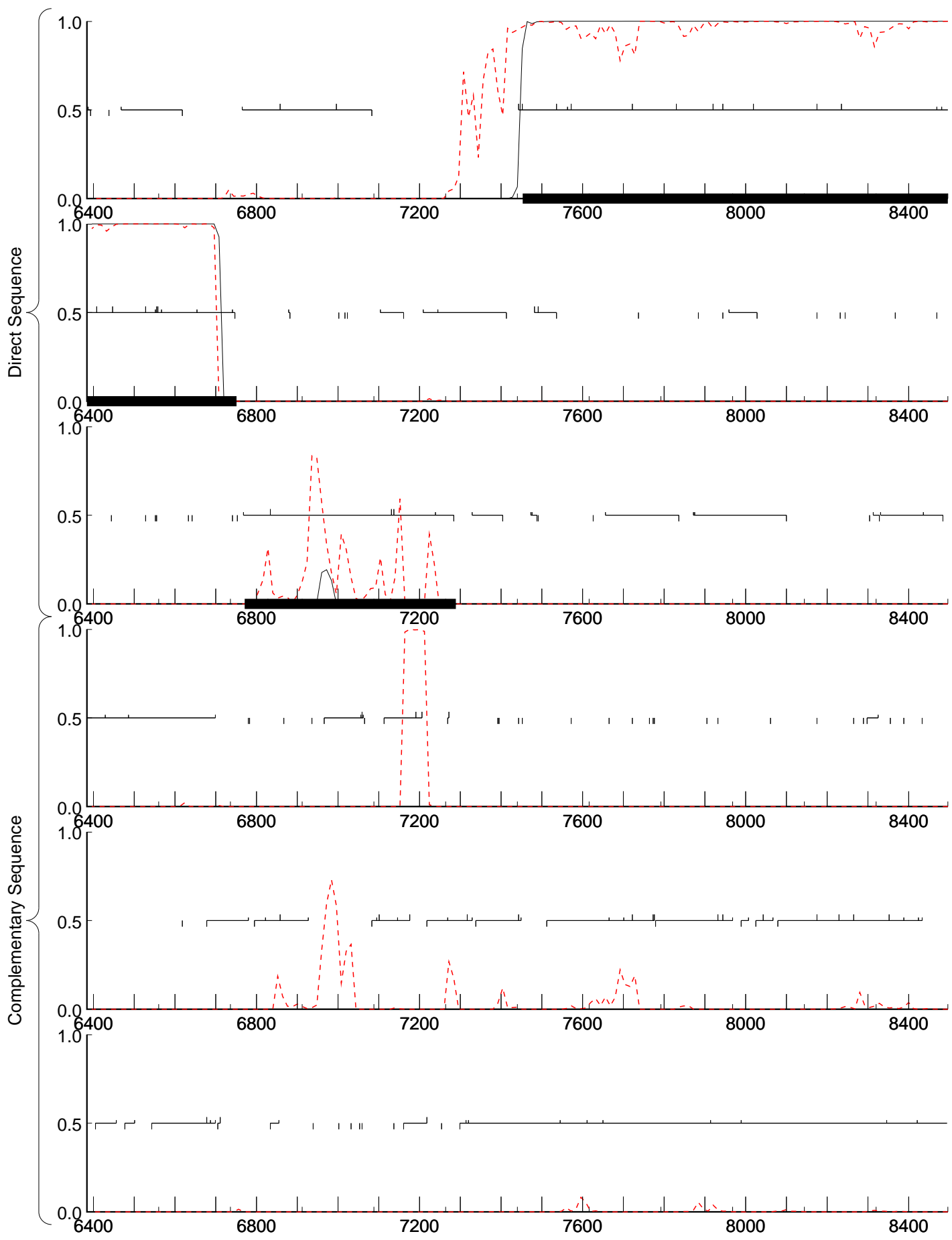


typical.ps

atypical.ps

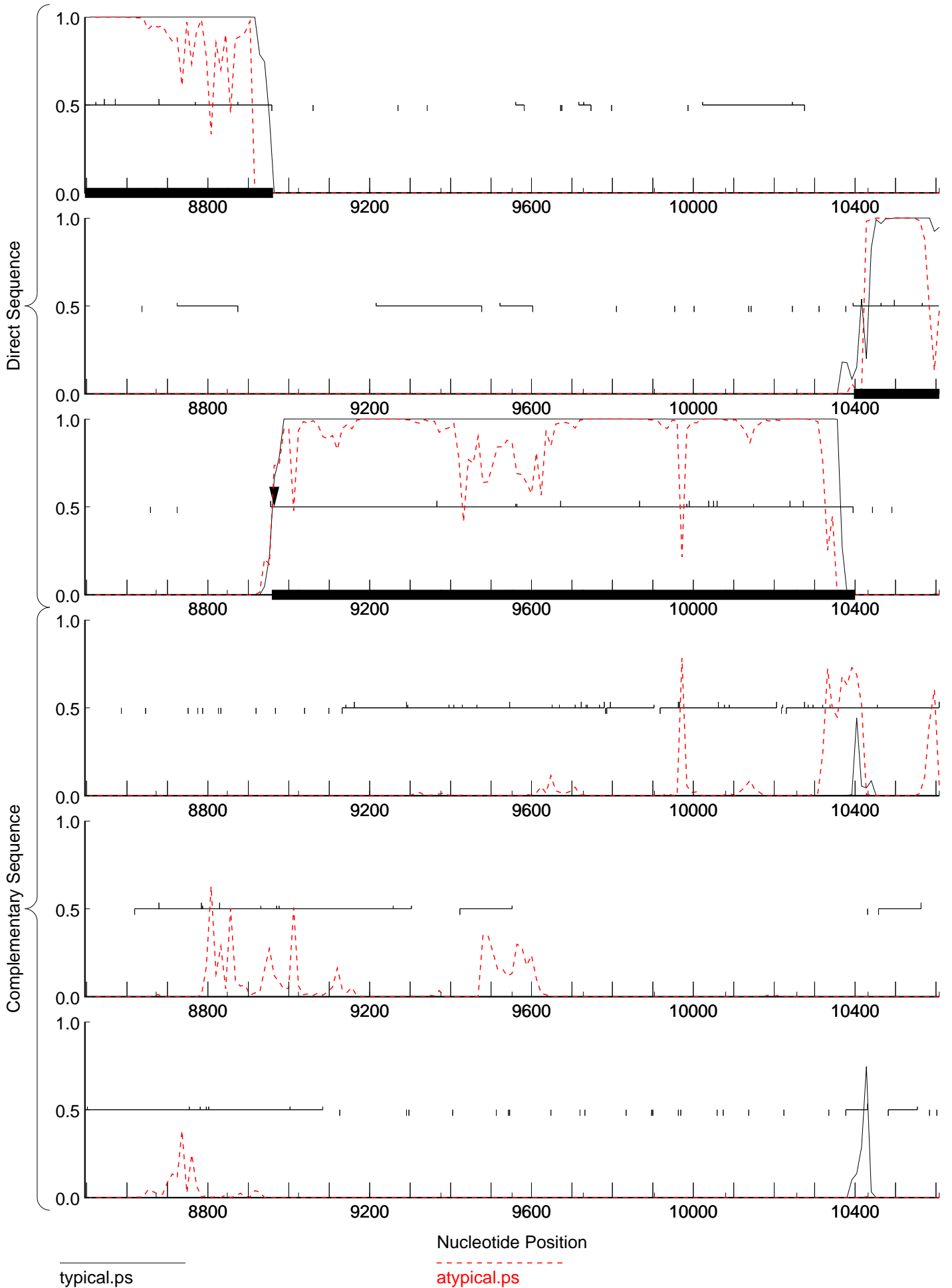


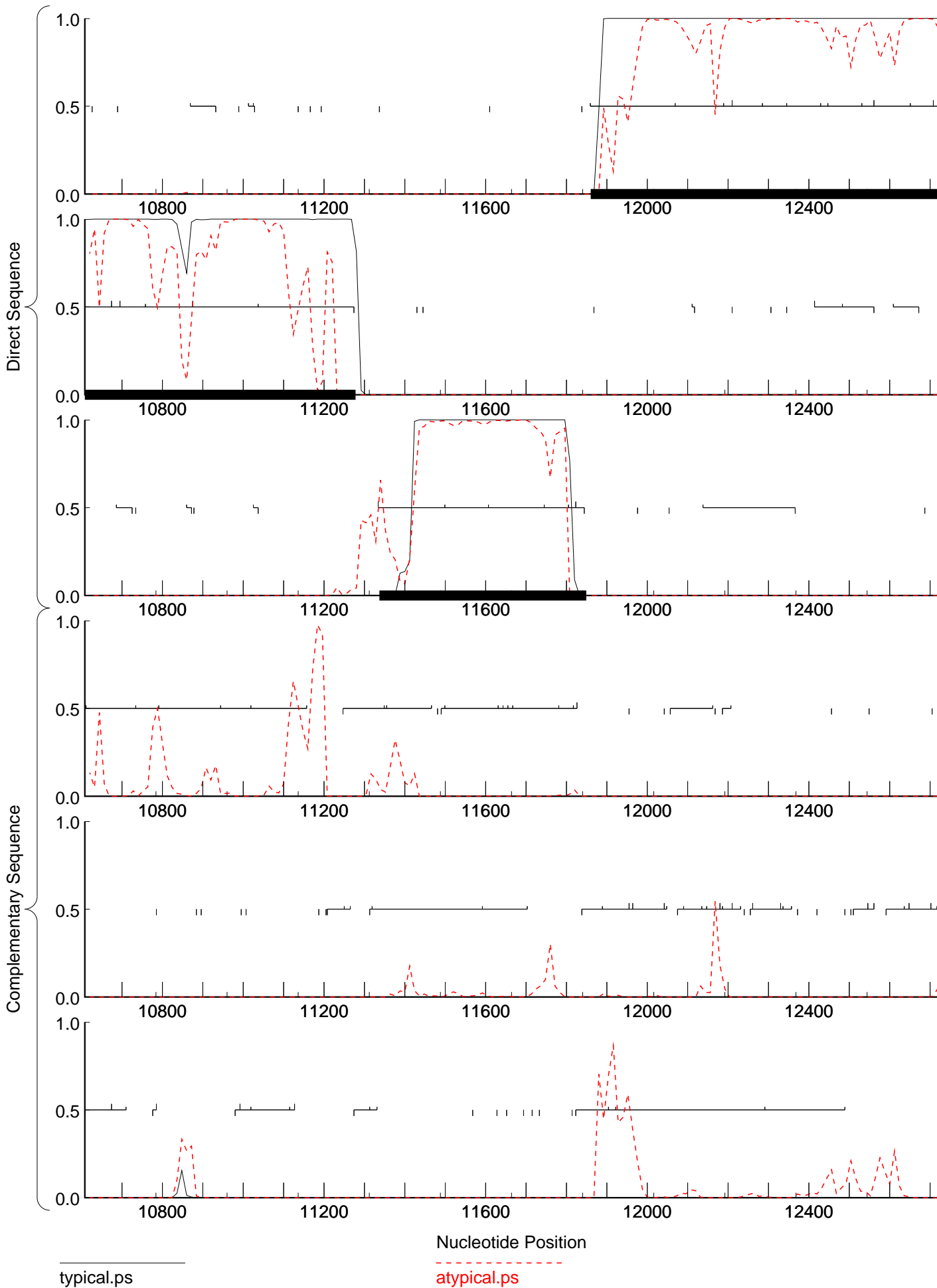


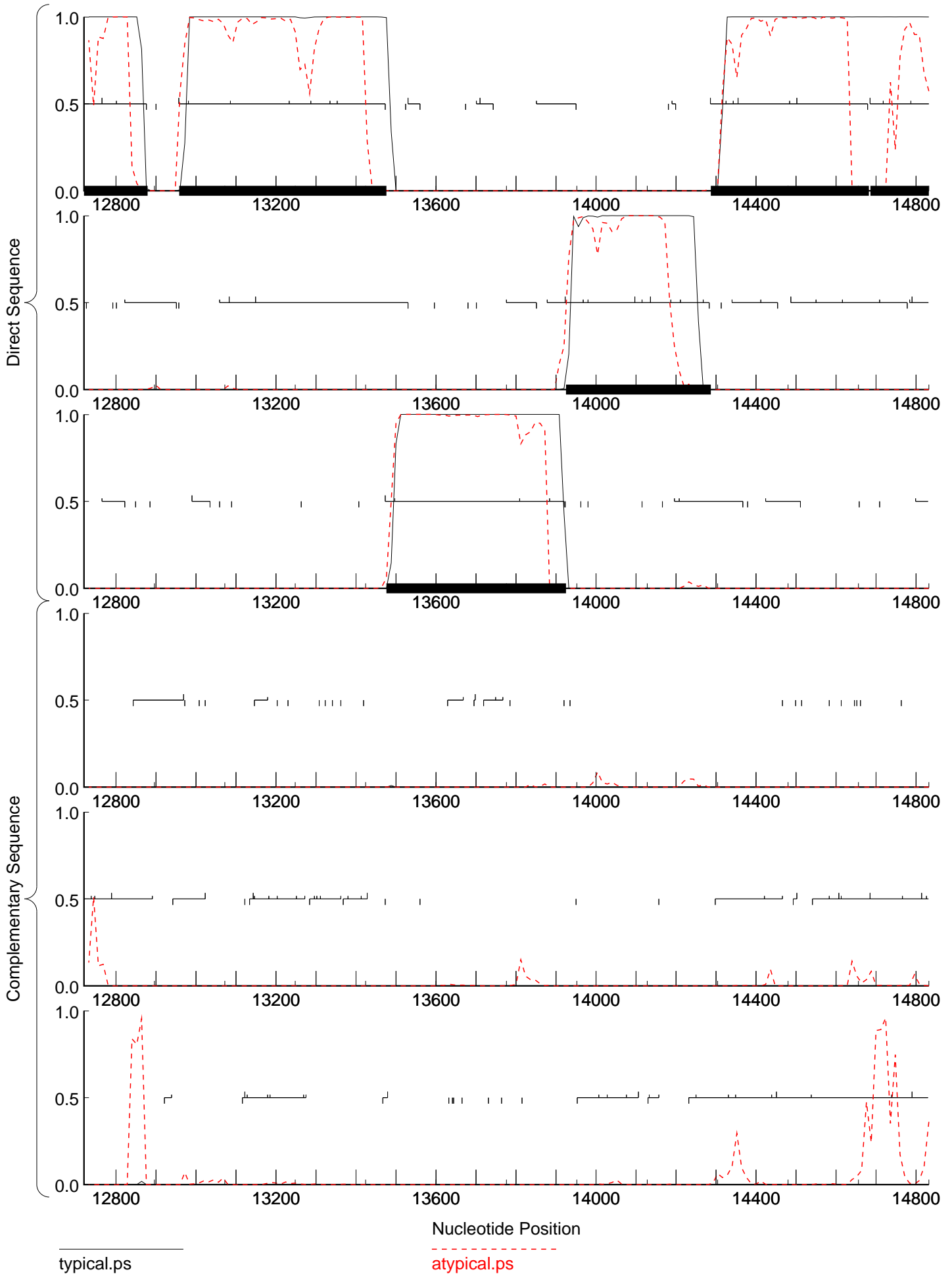


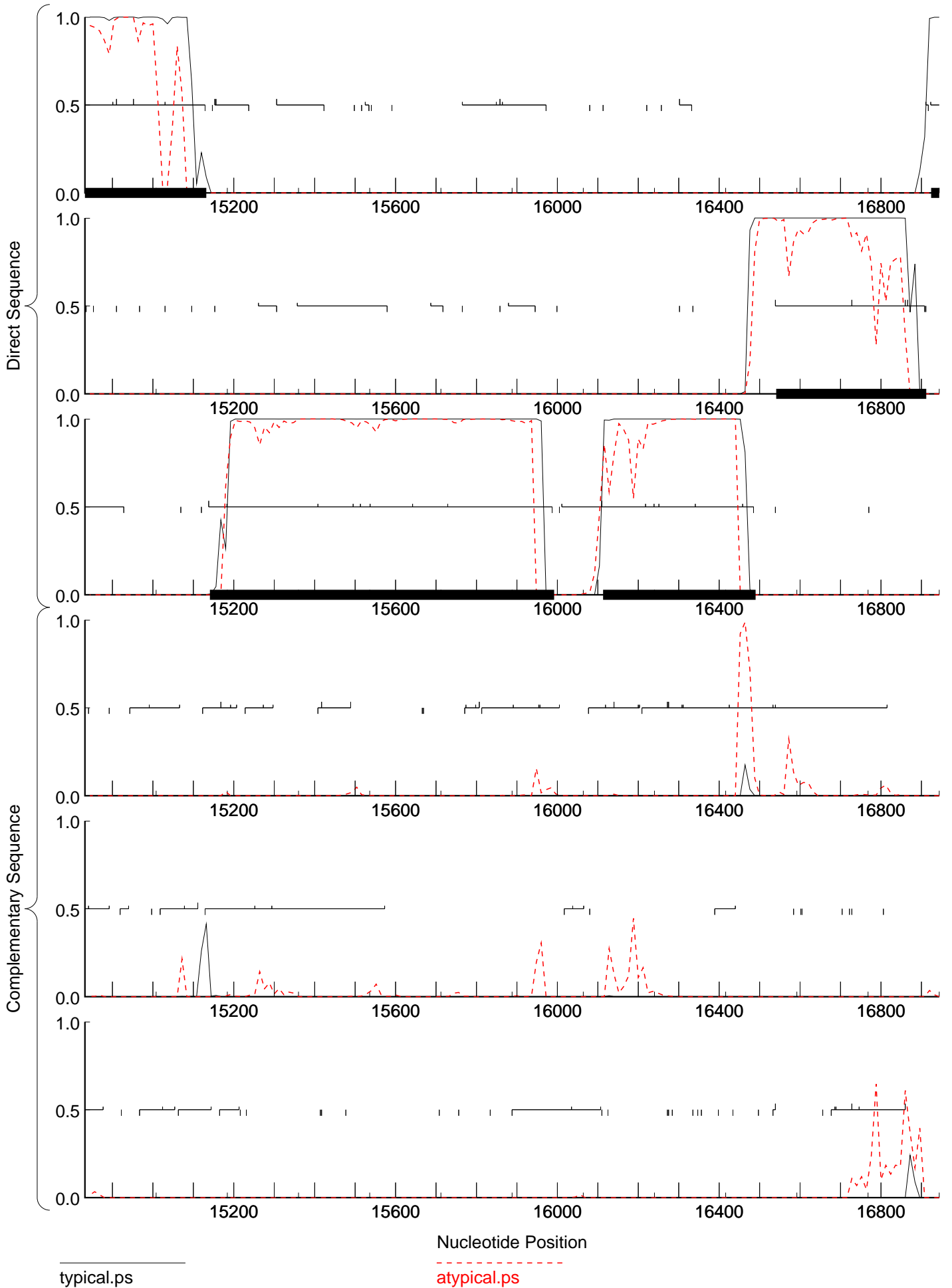
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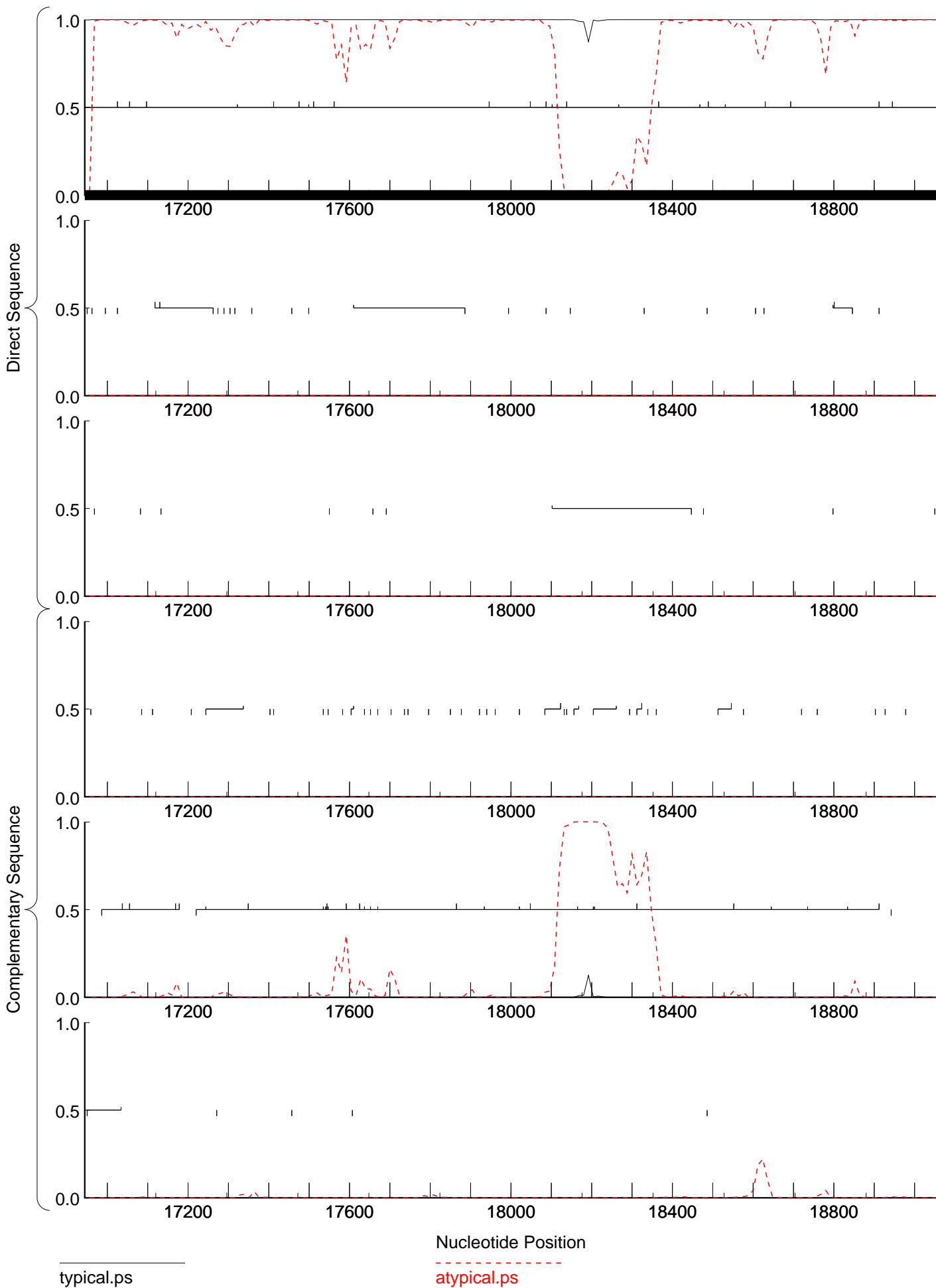
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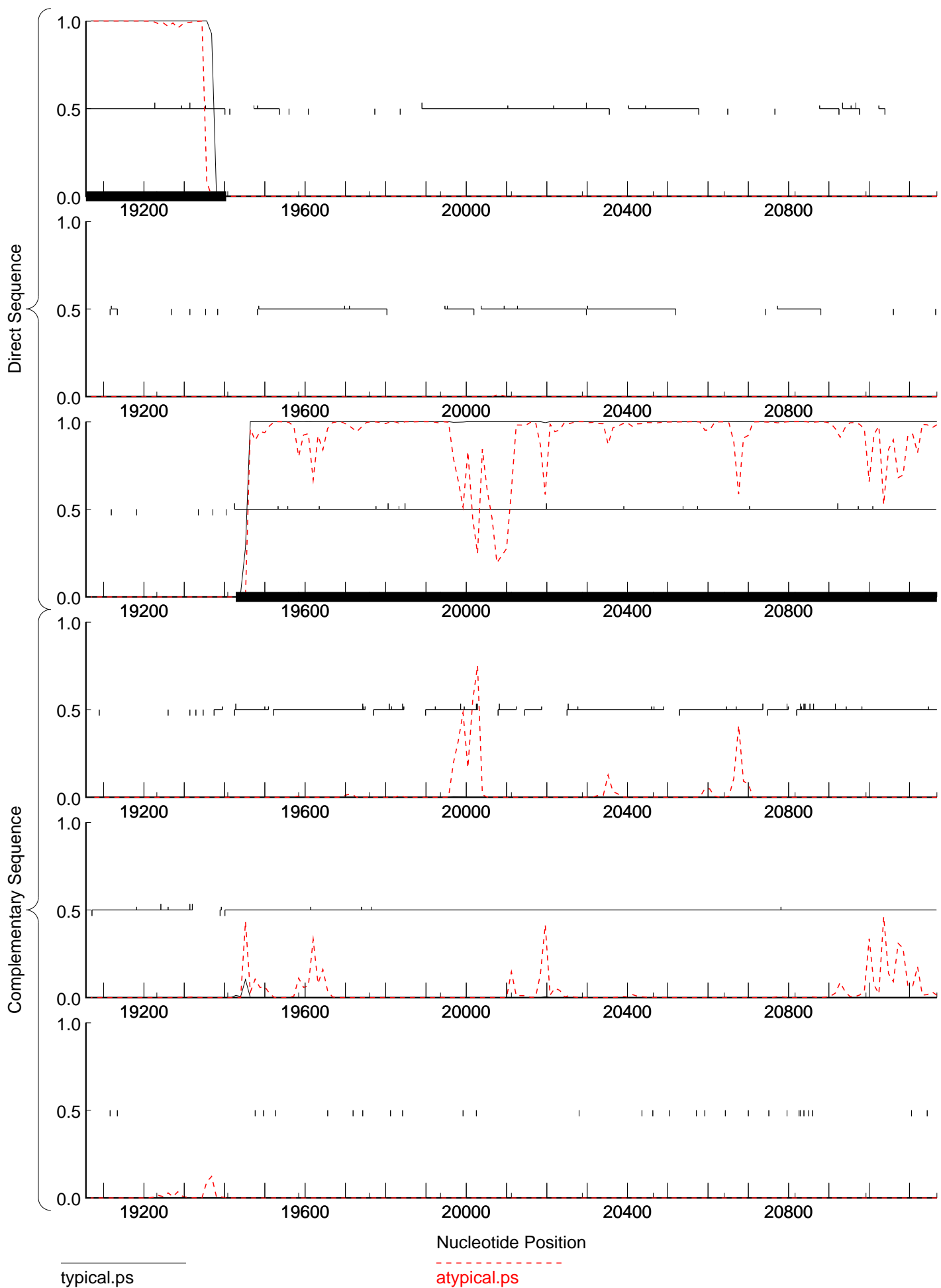


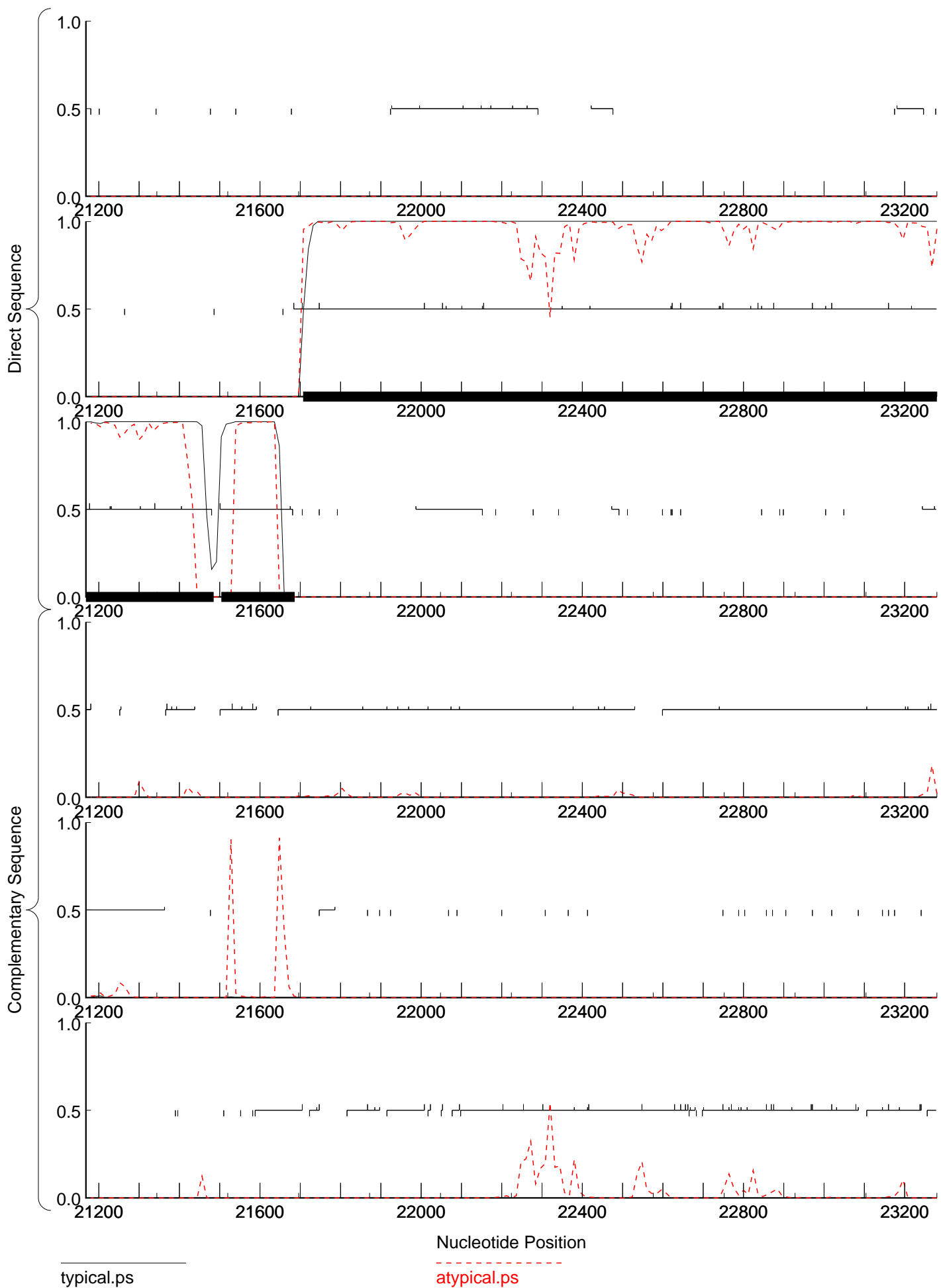


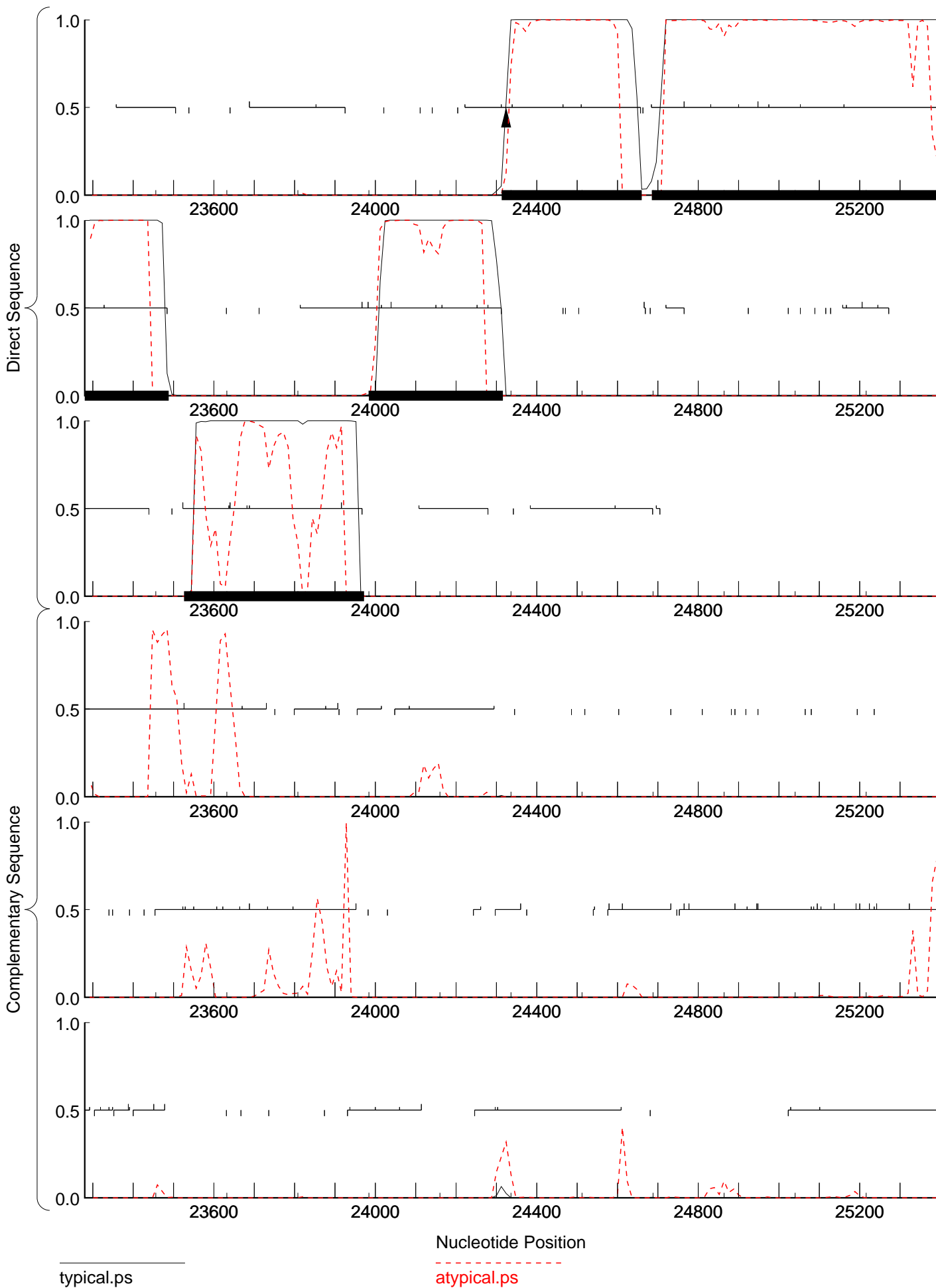


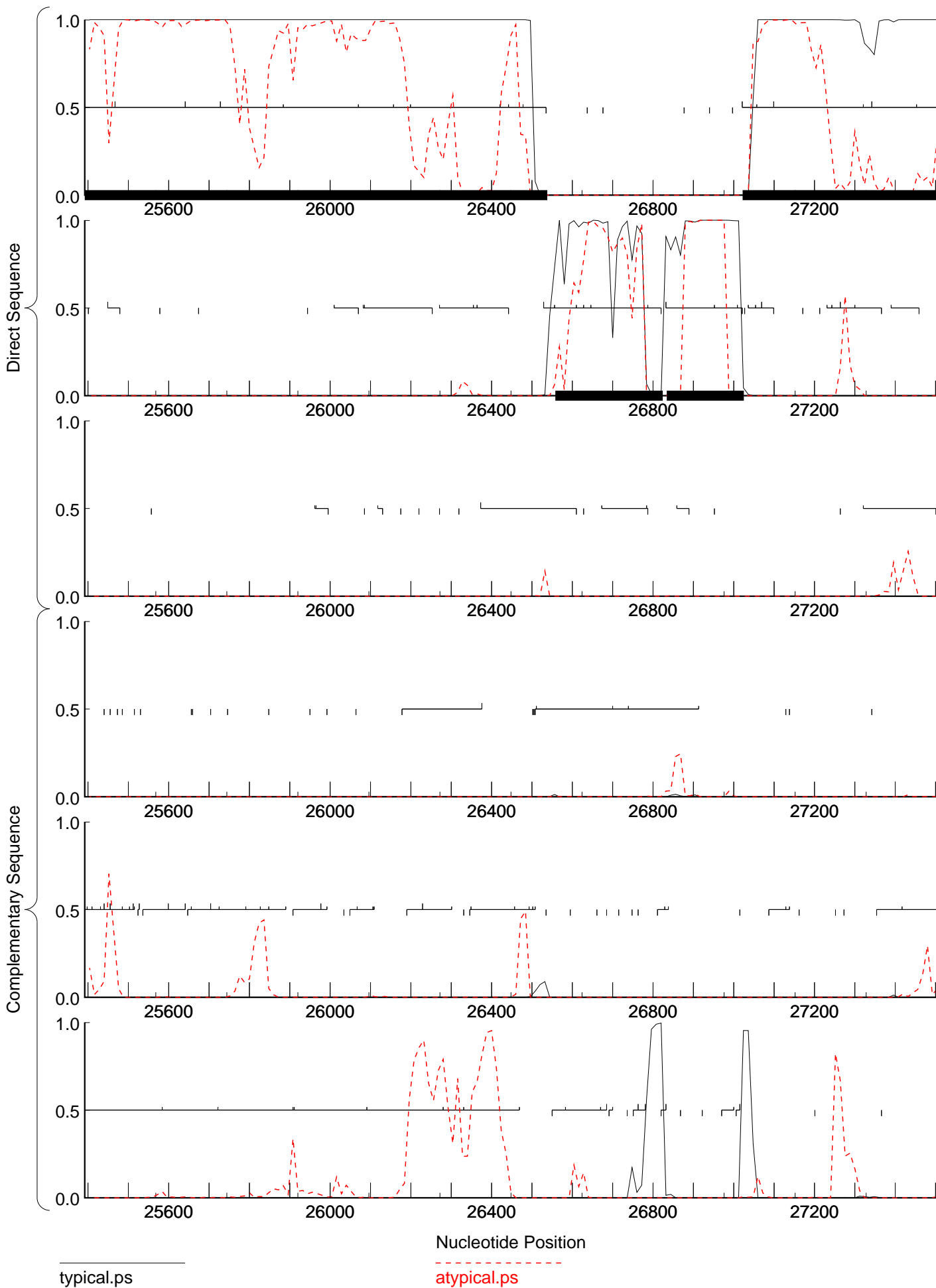


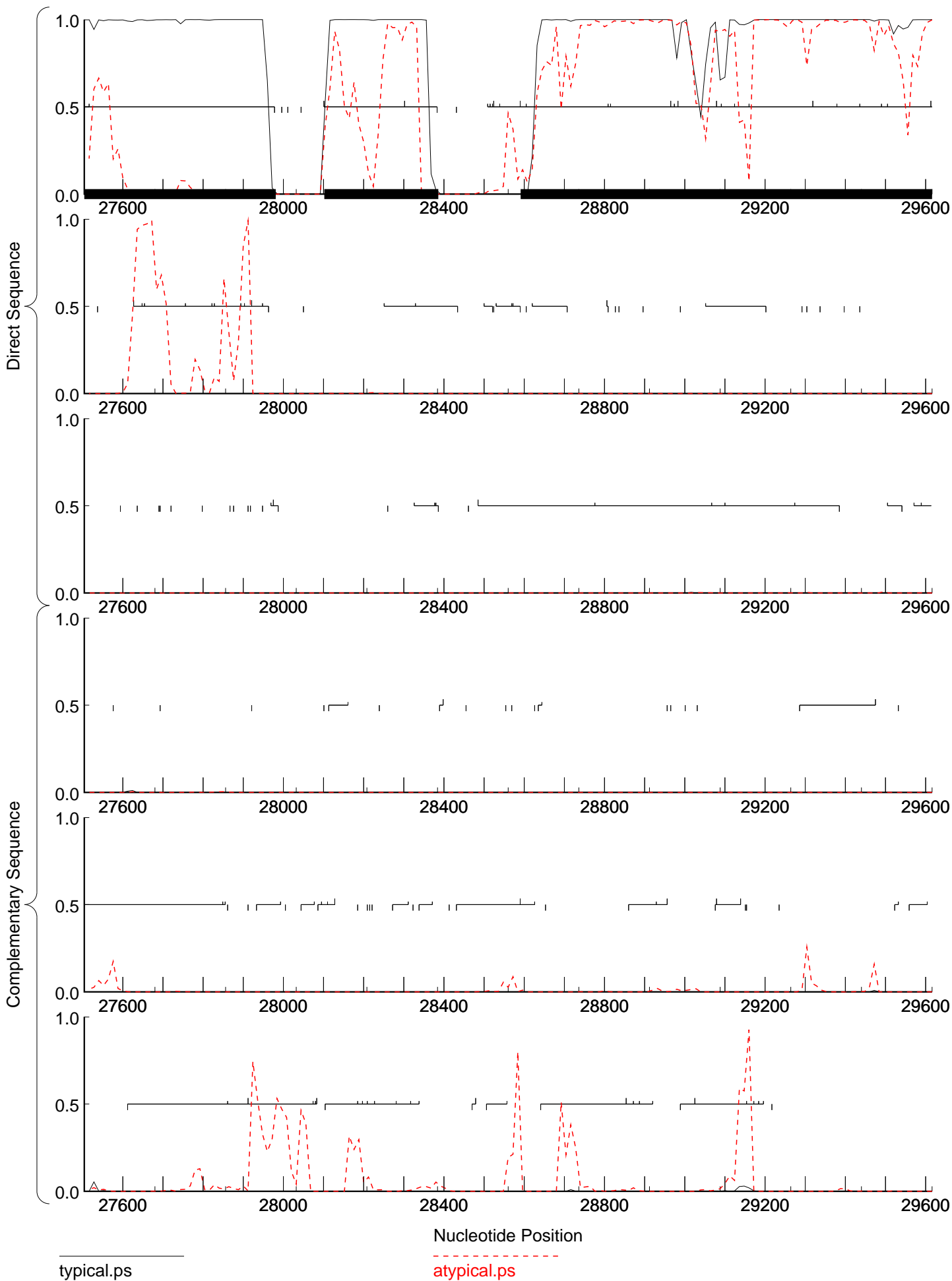


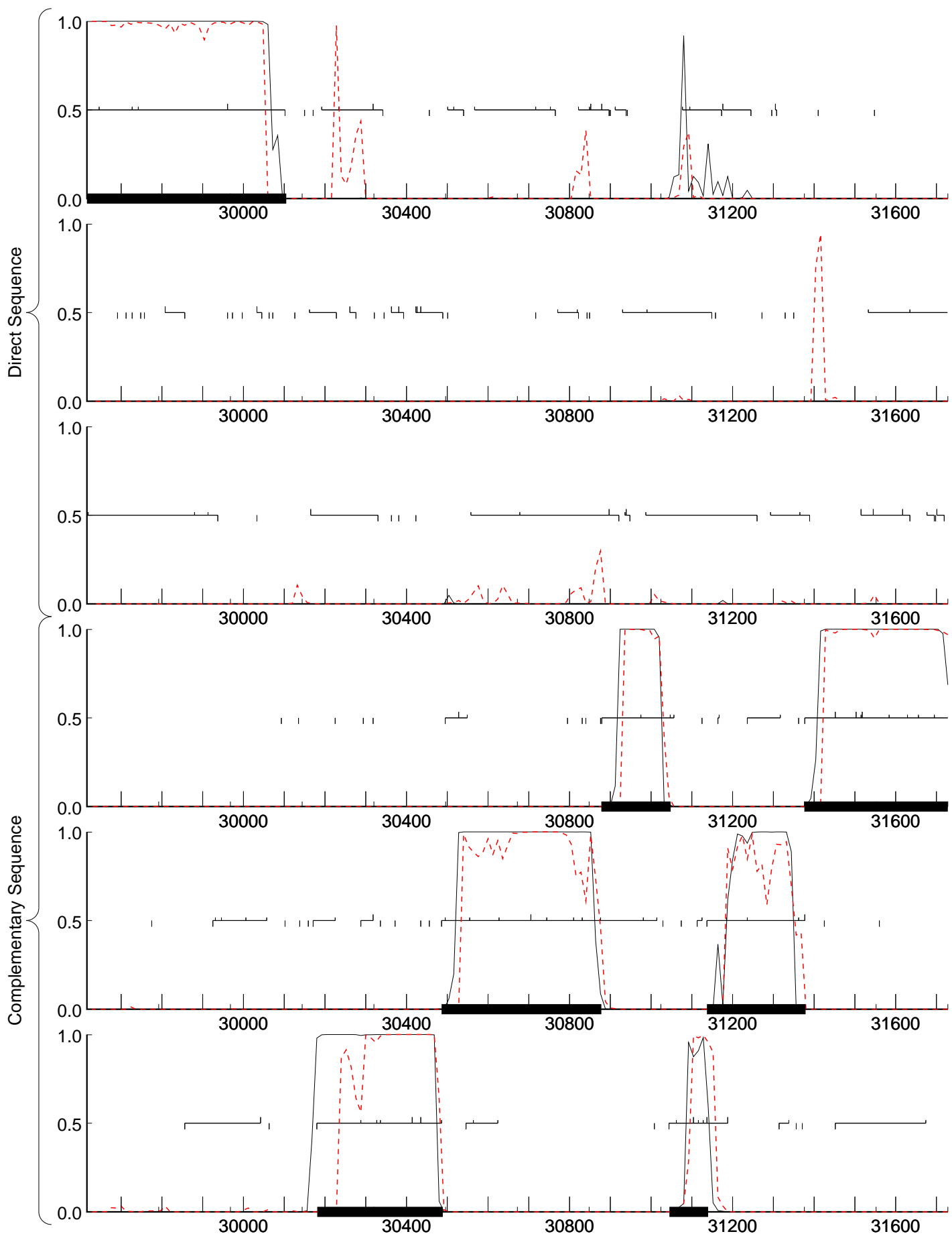






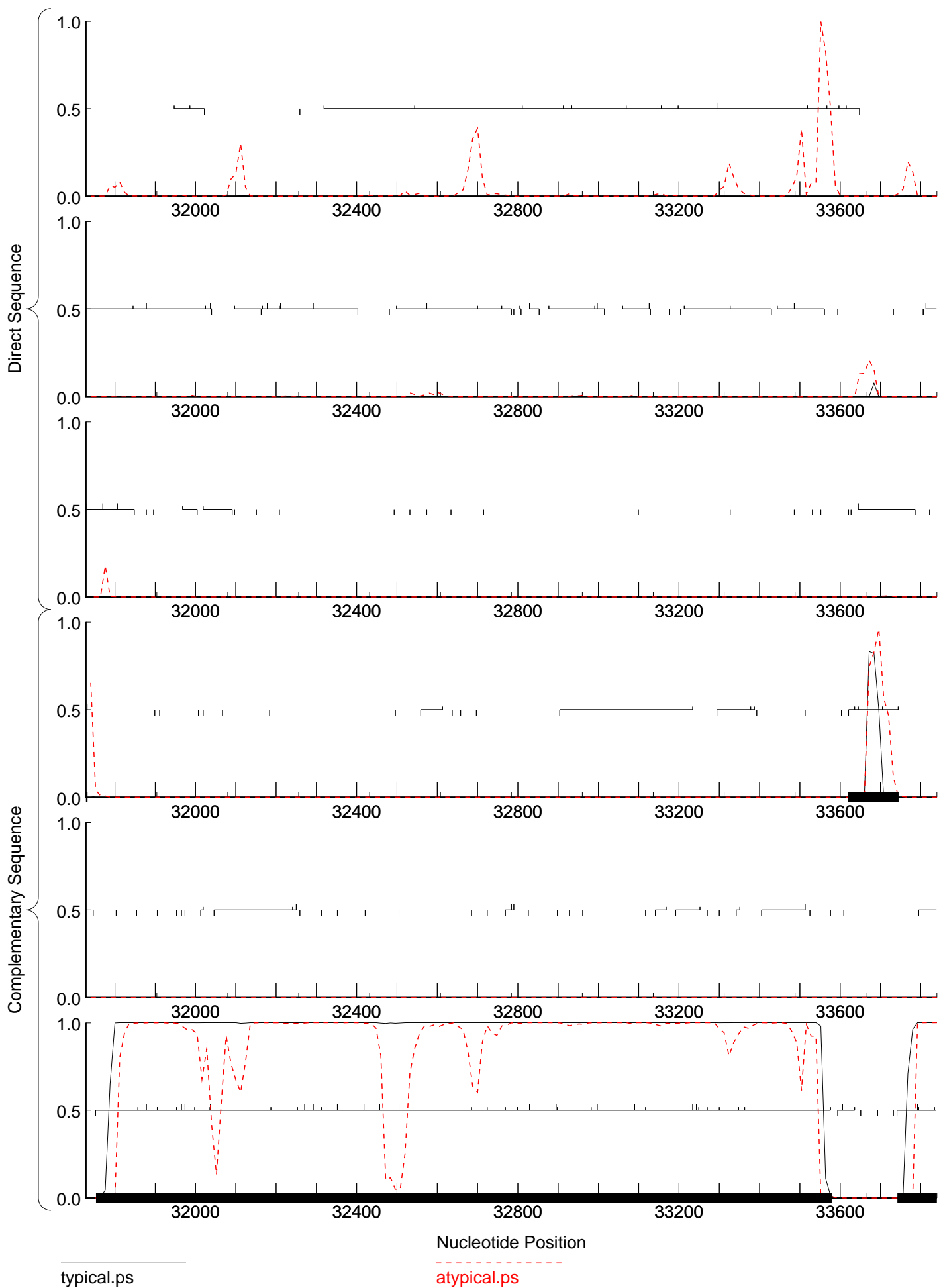


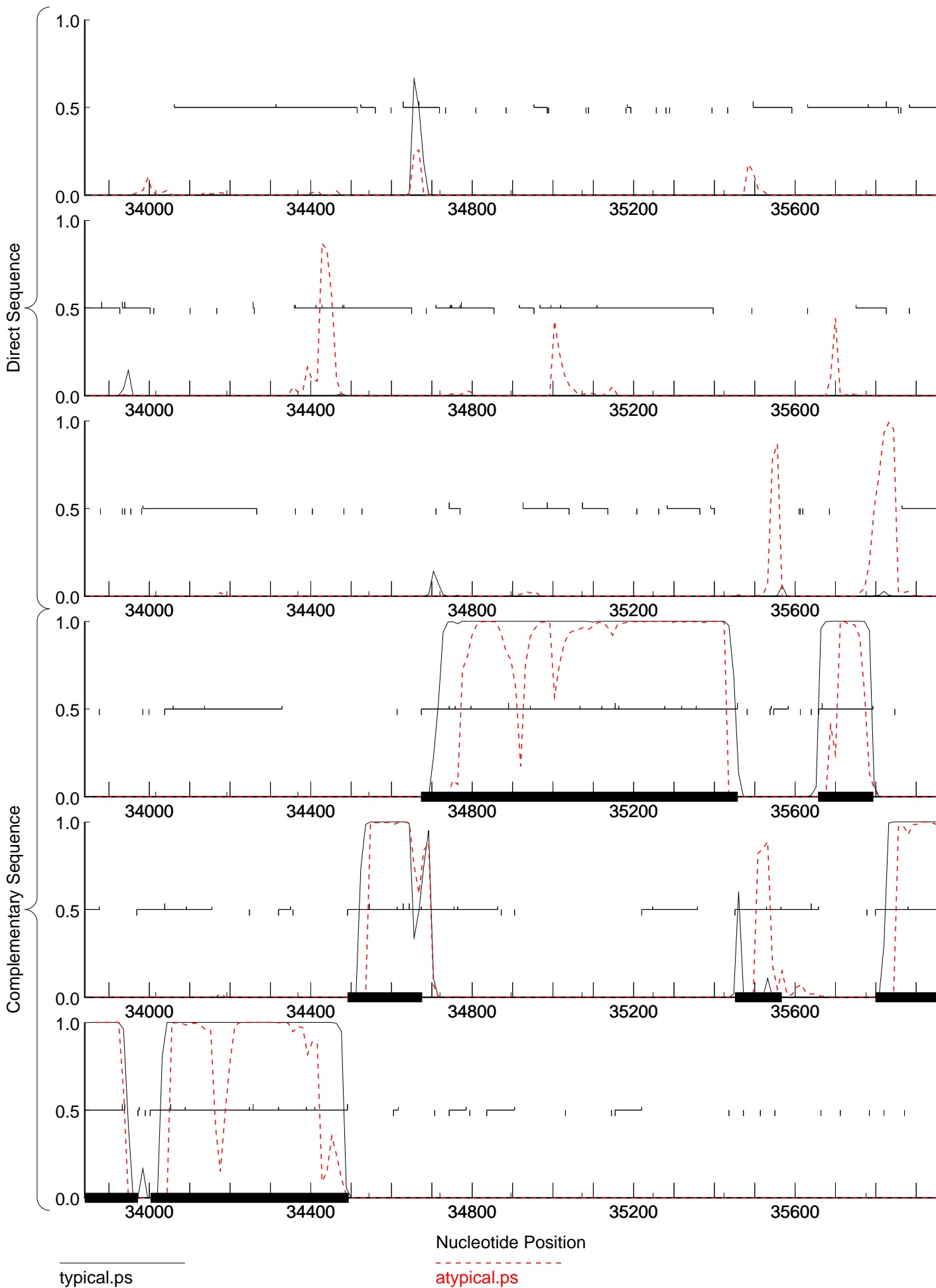


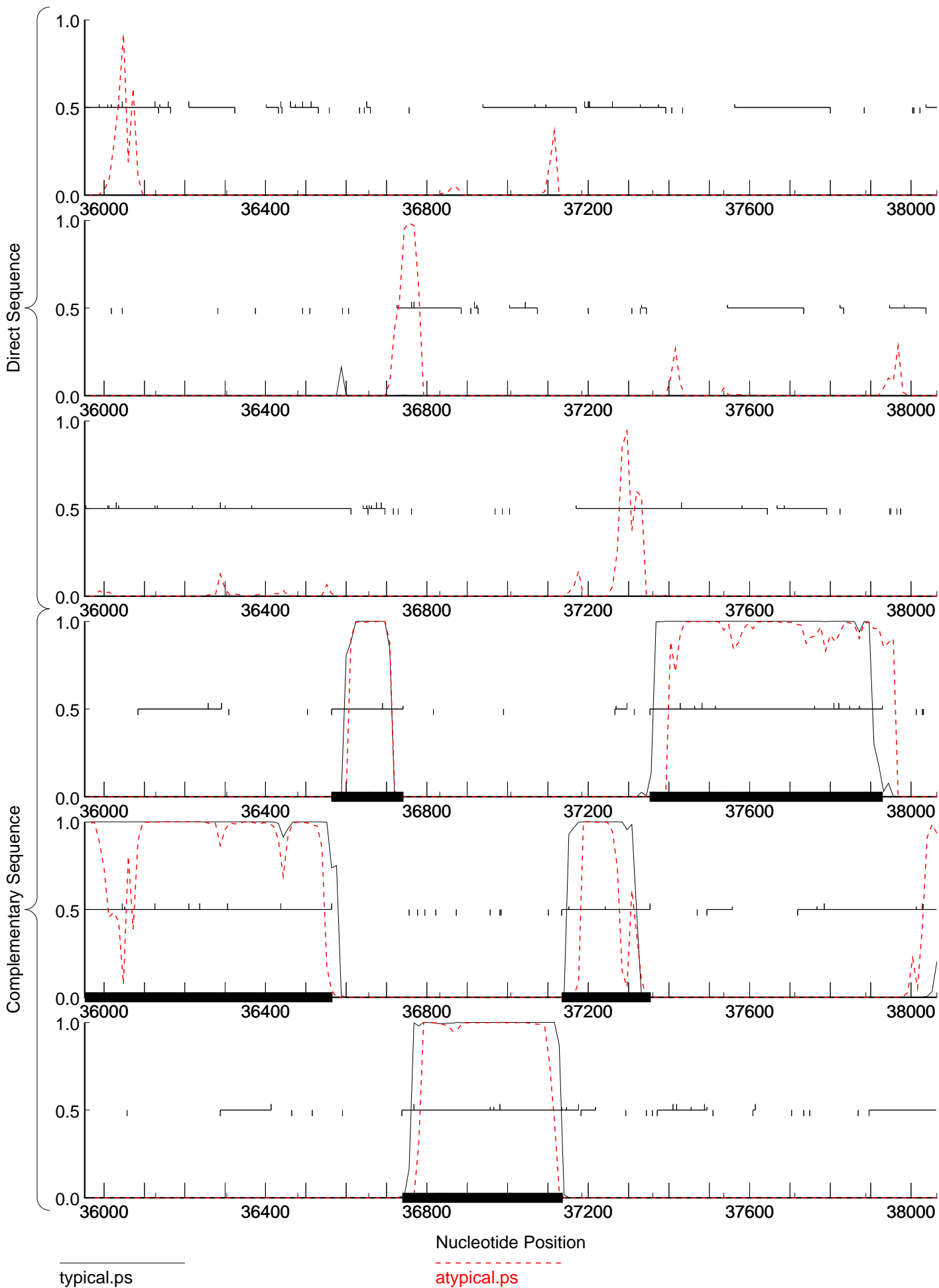


typical.ps

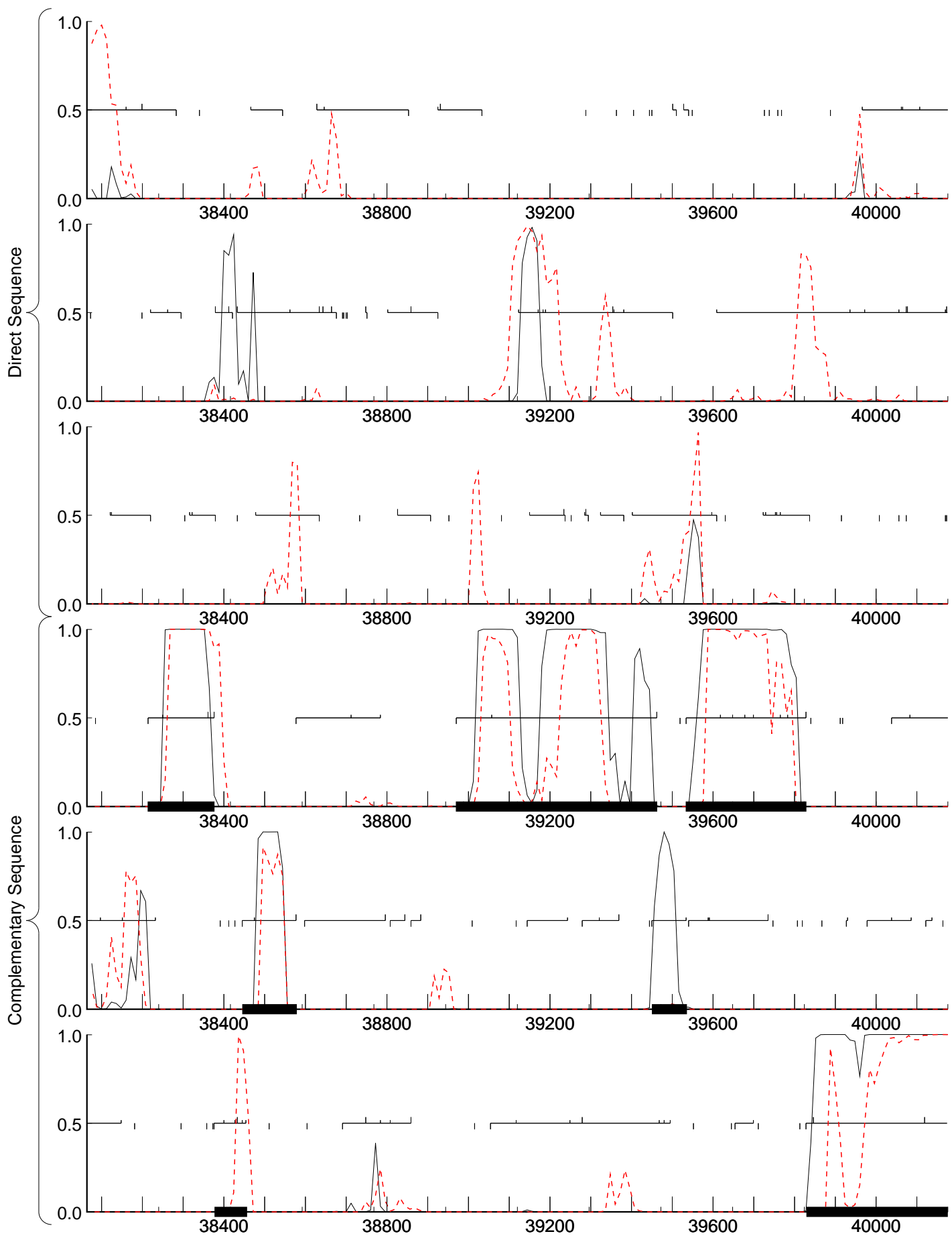
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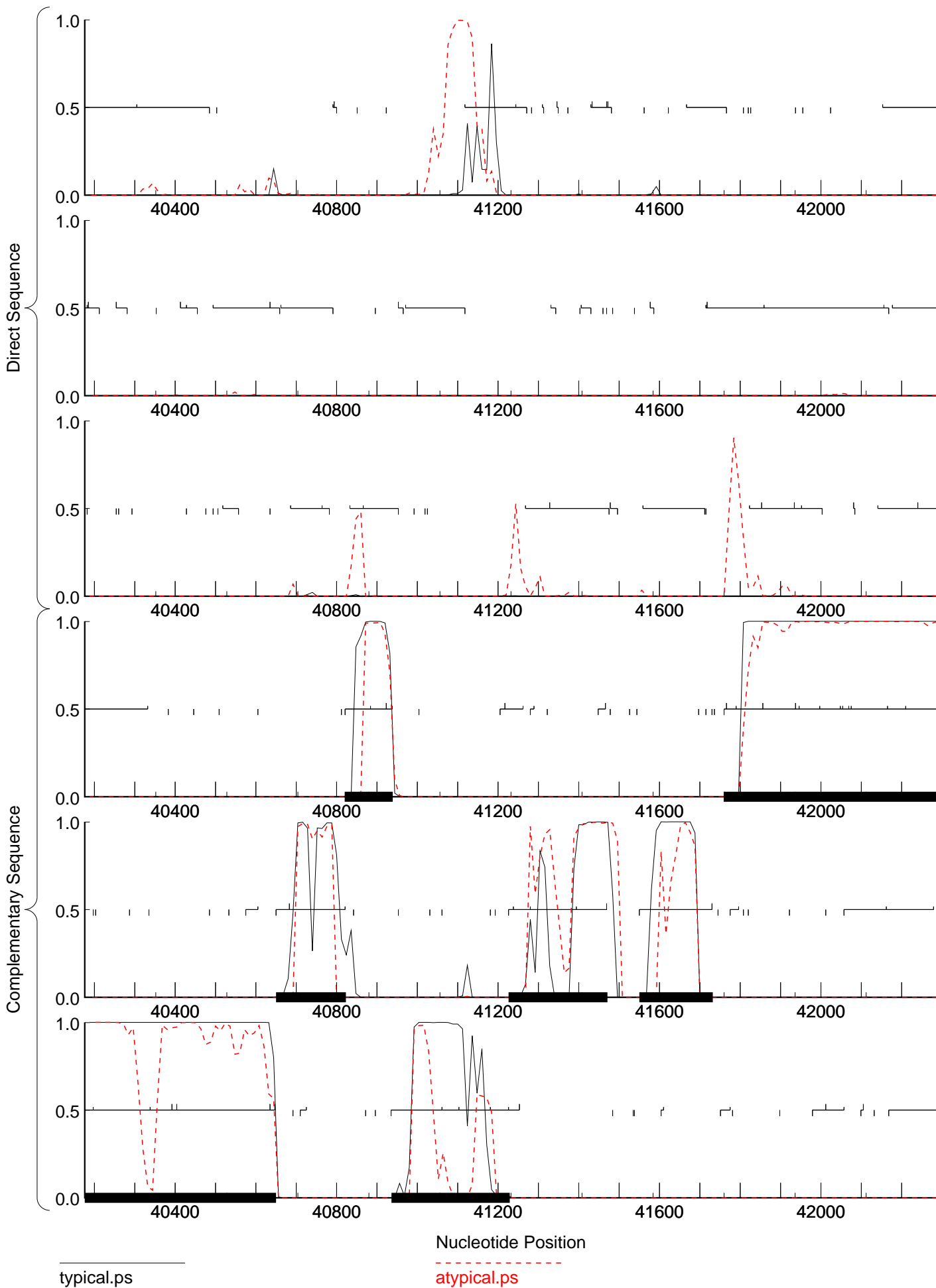
GeneMark.hmm prediction

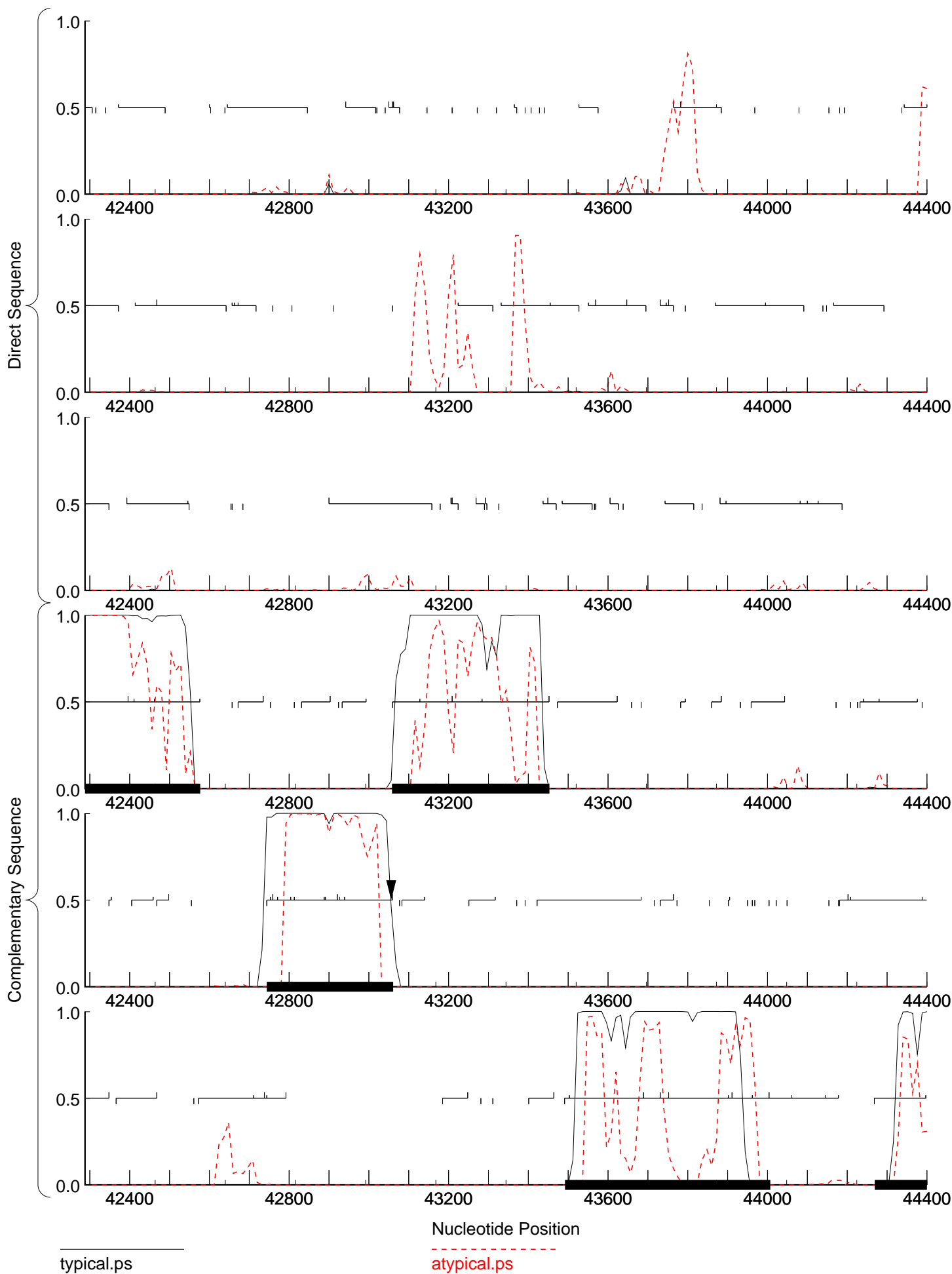


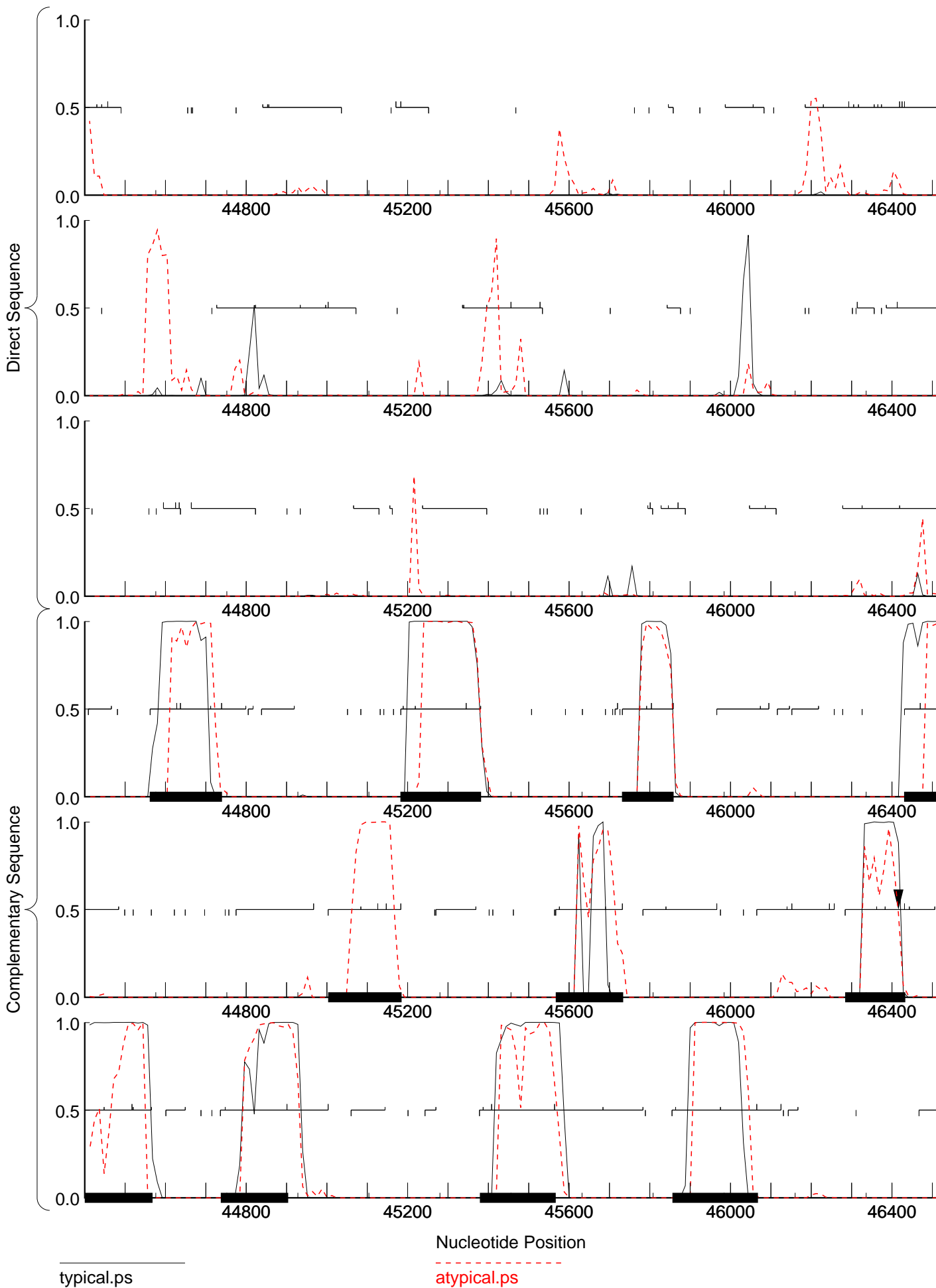
typical.ps

Nucleotide Position

atypical.ps







typical.ps

atypical.ps

