

GeneMark

Version 2.5p (09.08.06)

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PROGRAM INFORMATION

Sequence : Mycobacterium phage Ohfah complete sequence, 51412 bp including 10-base 3' overhang (CGGCCG
Analysis Date : 5/27/18 at 21:40:14
Pages : 26
Sequence Length : 51412 bp
GC Content : 63.84%

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,

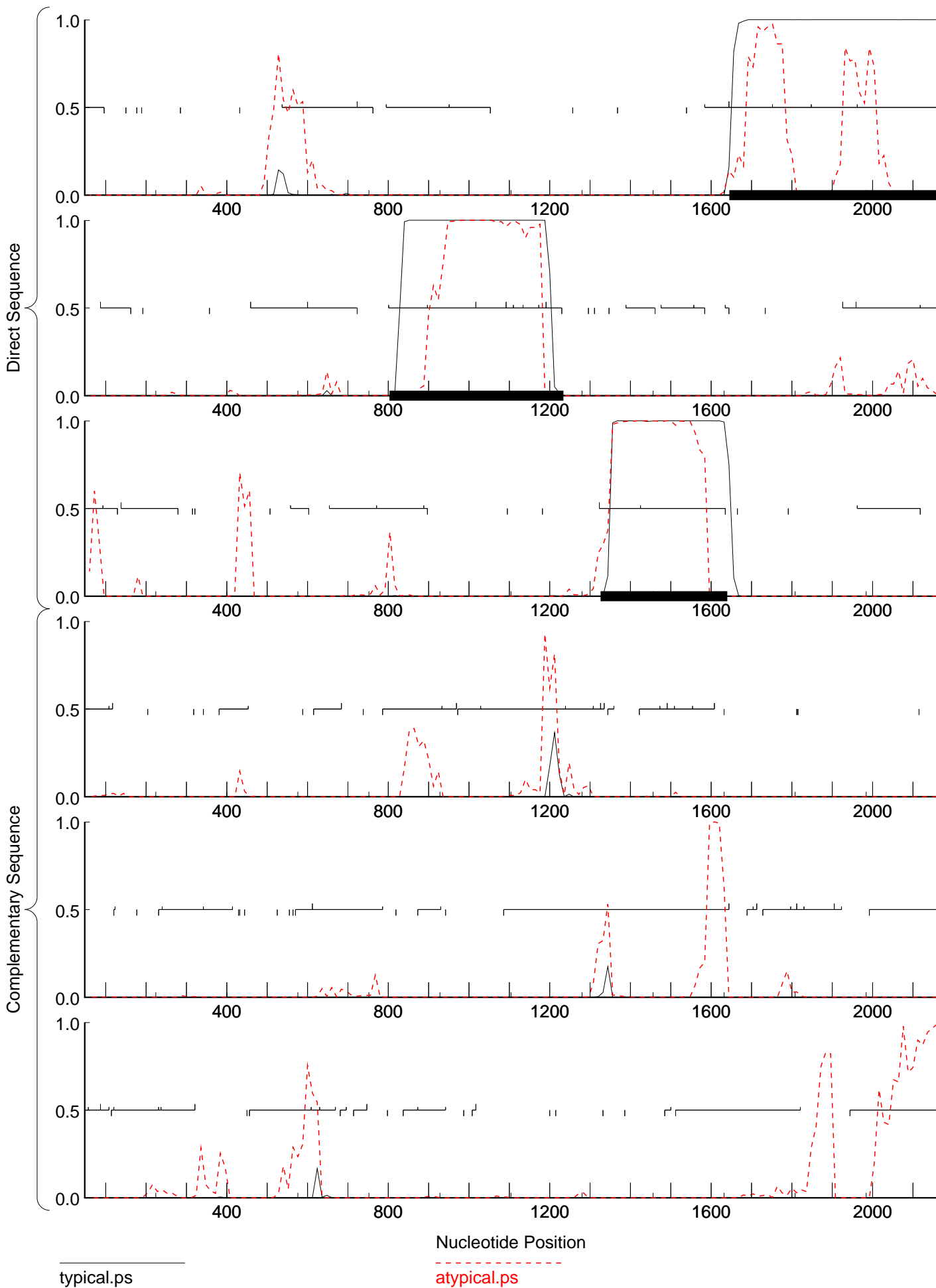
MATRIX INFORMATION

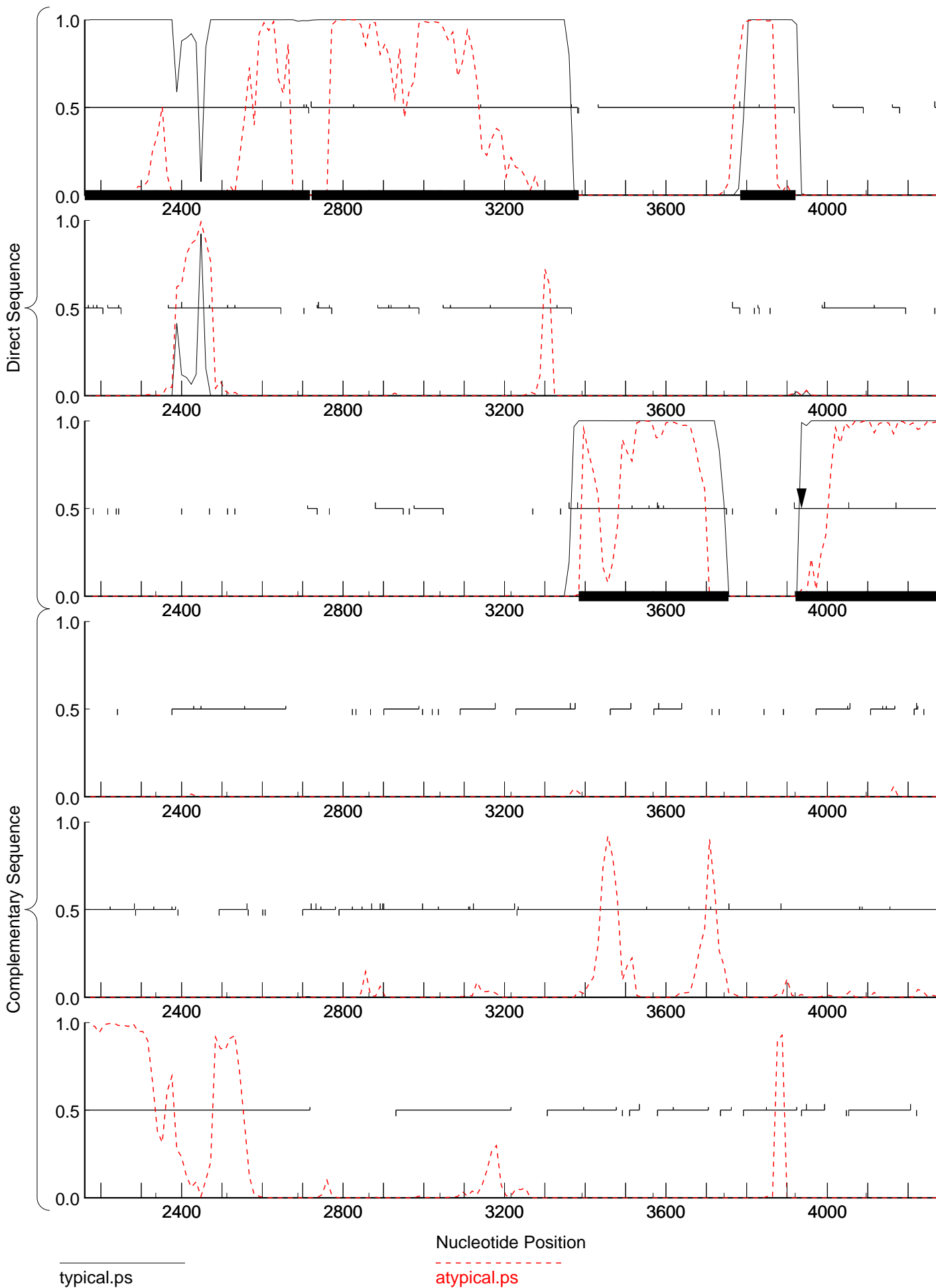
Matrix : -
Author : -
Order : 4

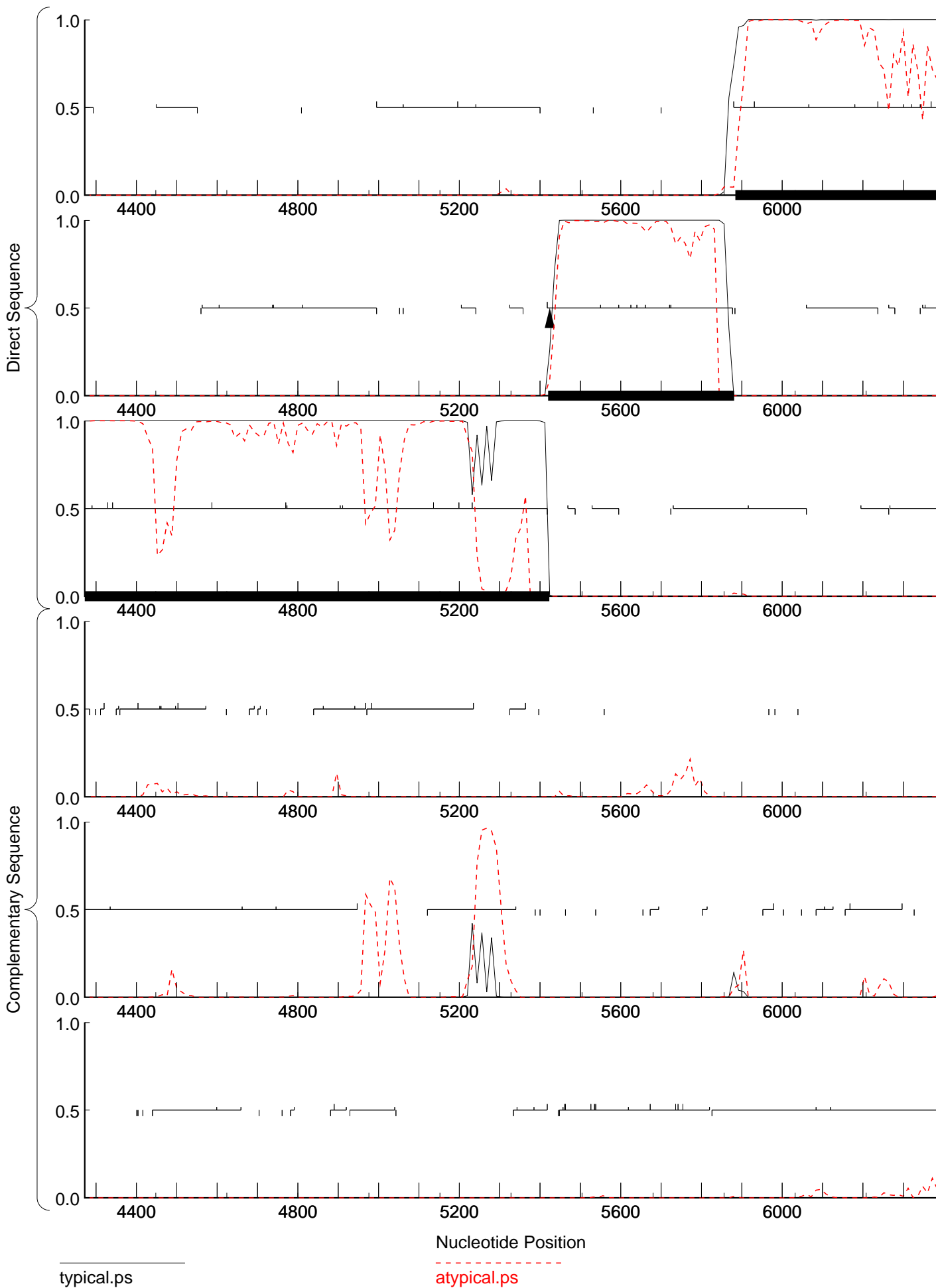
Send questions / comments to:
Dr. M. Borodovsky
Georgia Institute of Technology
School of Biology
Atlanta, GA 30332-0230

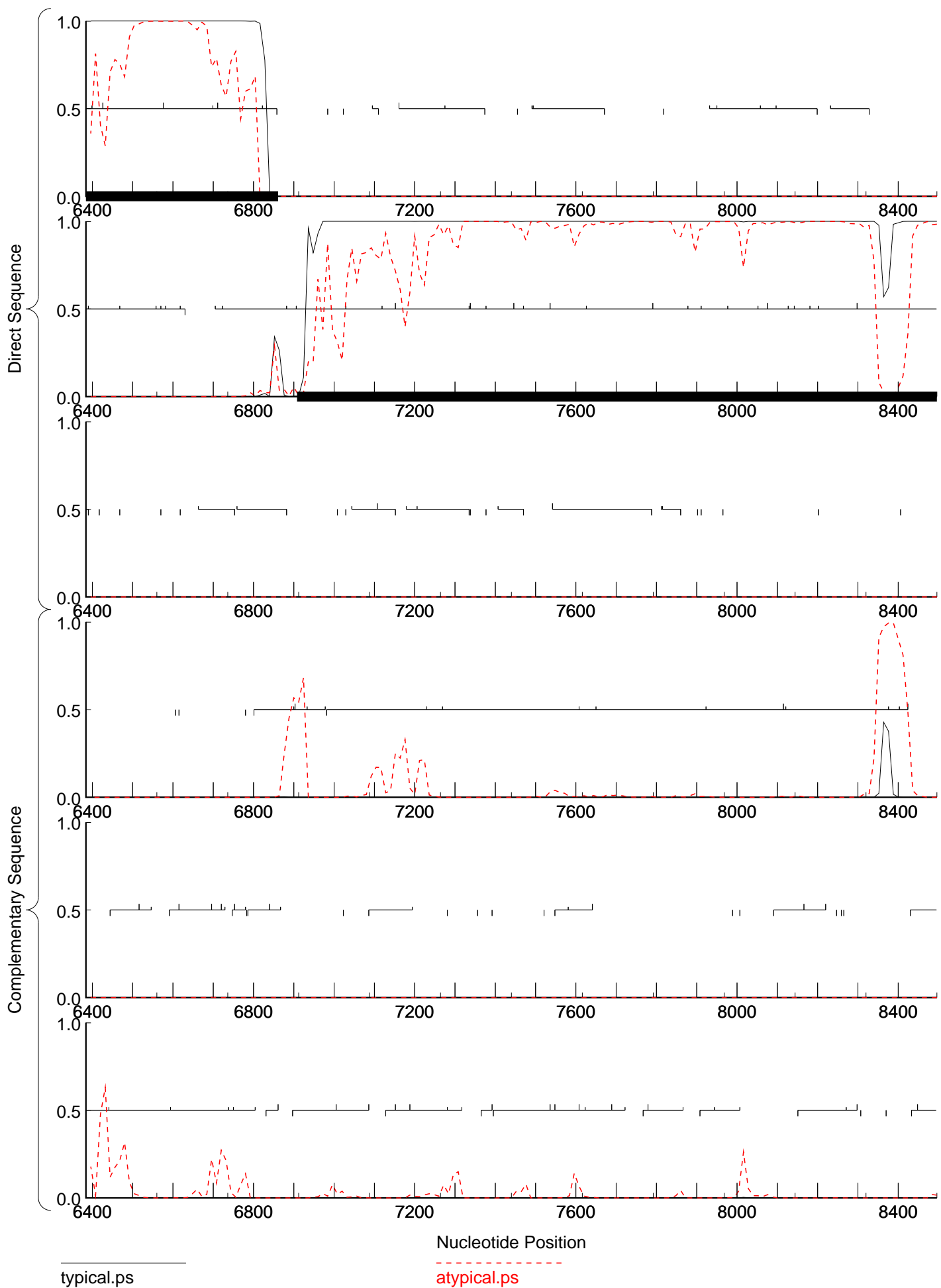
Matrix notes & comments

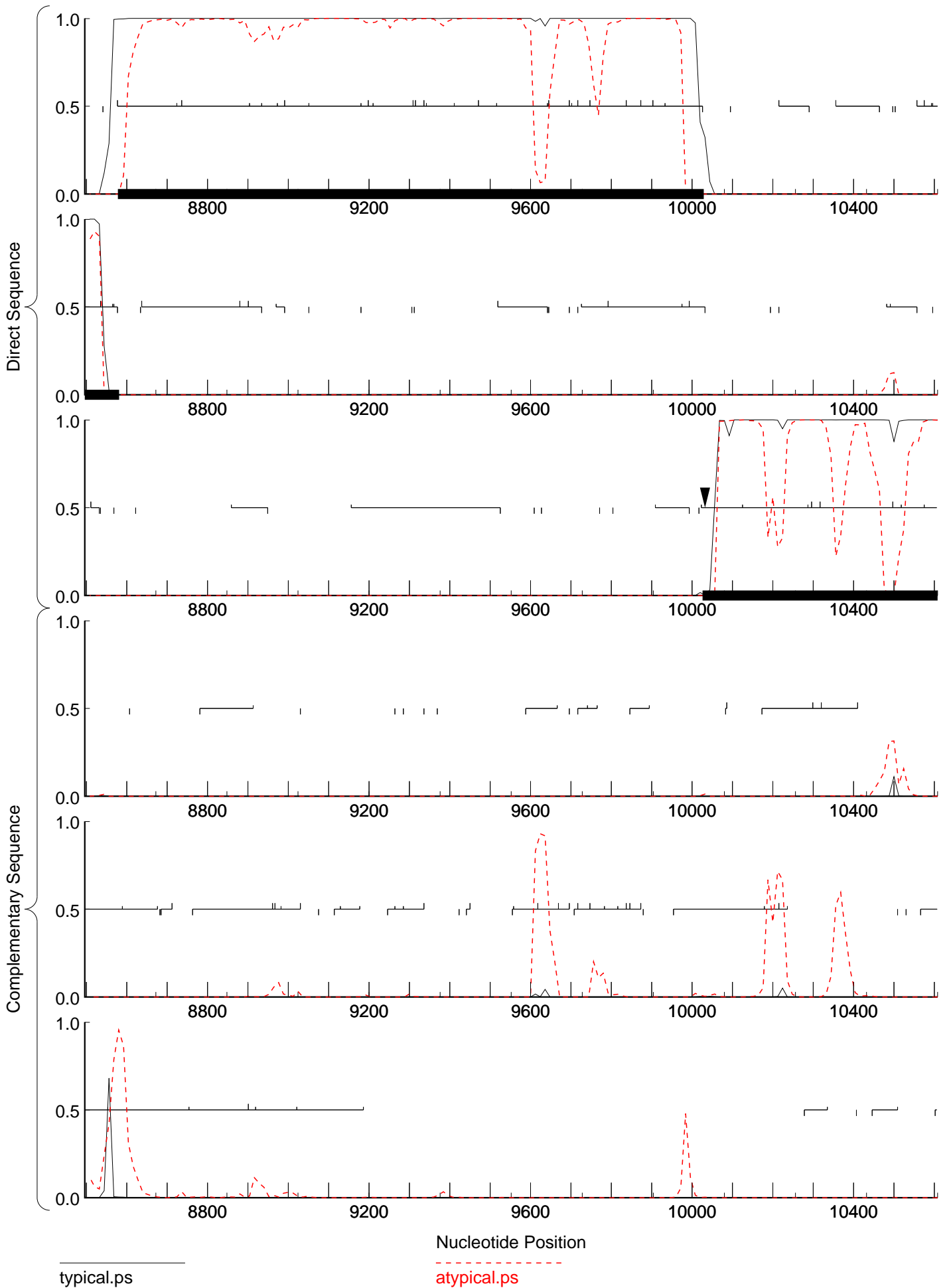
Training set derived by GeneMarkS, 4.30 October 2014
Sun May 27 21:40:13 2018

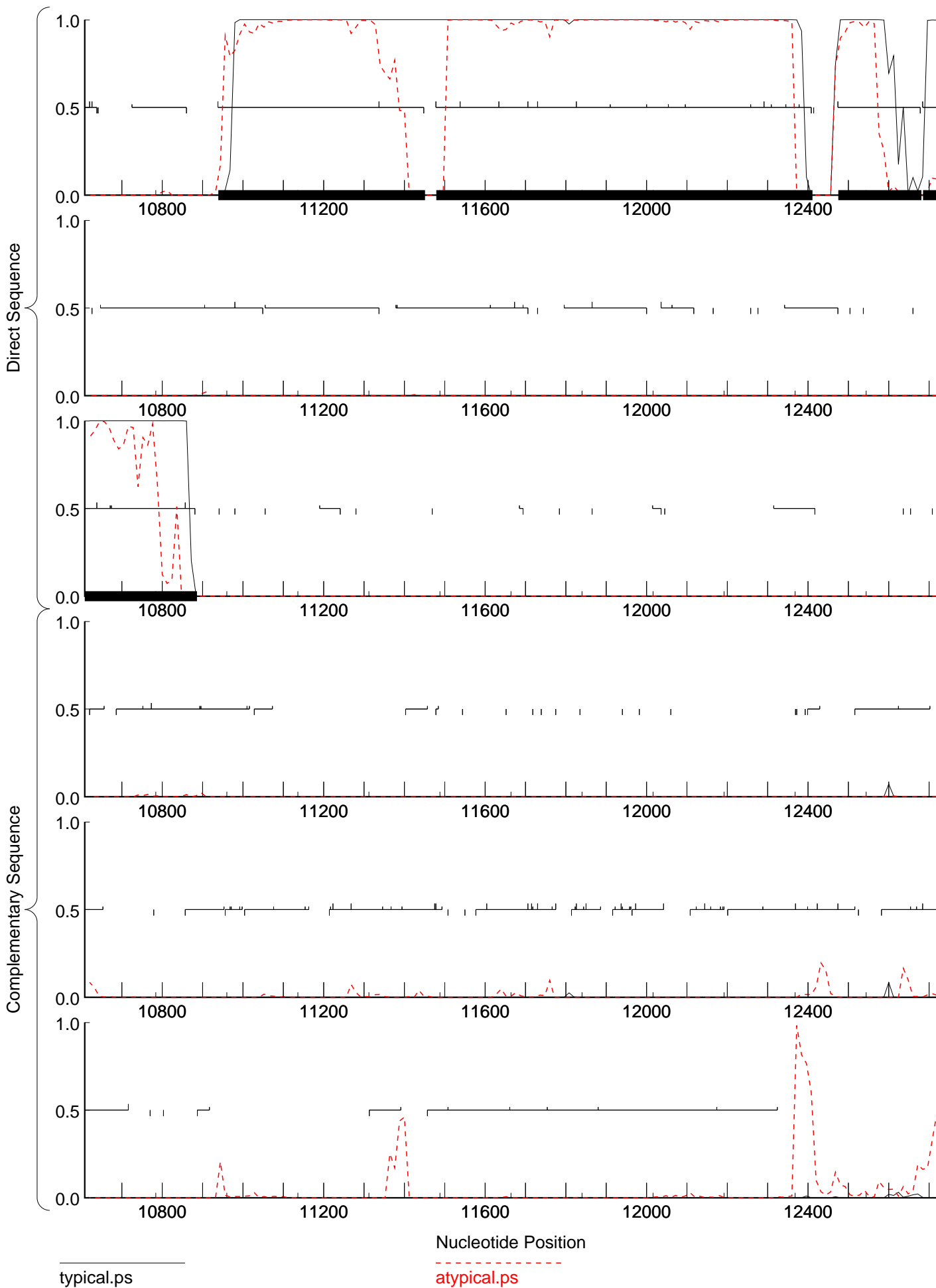


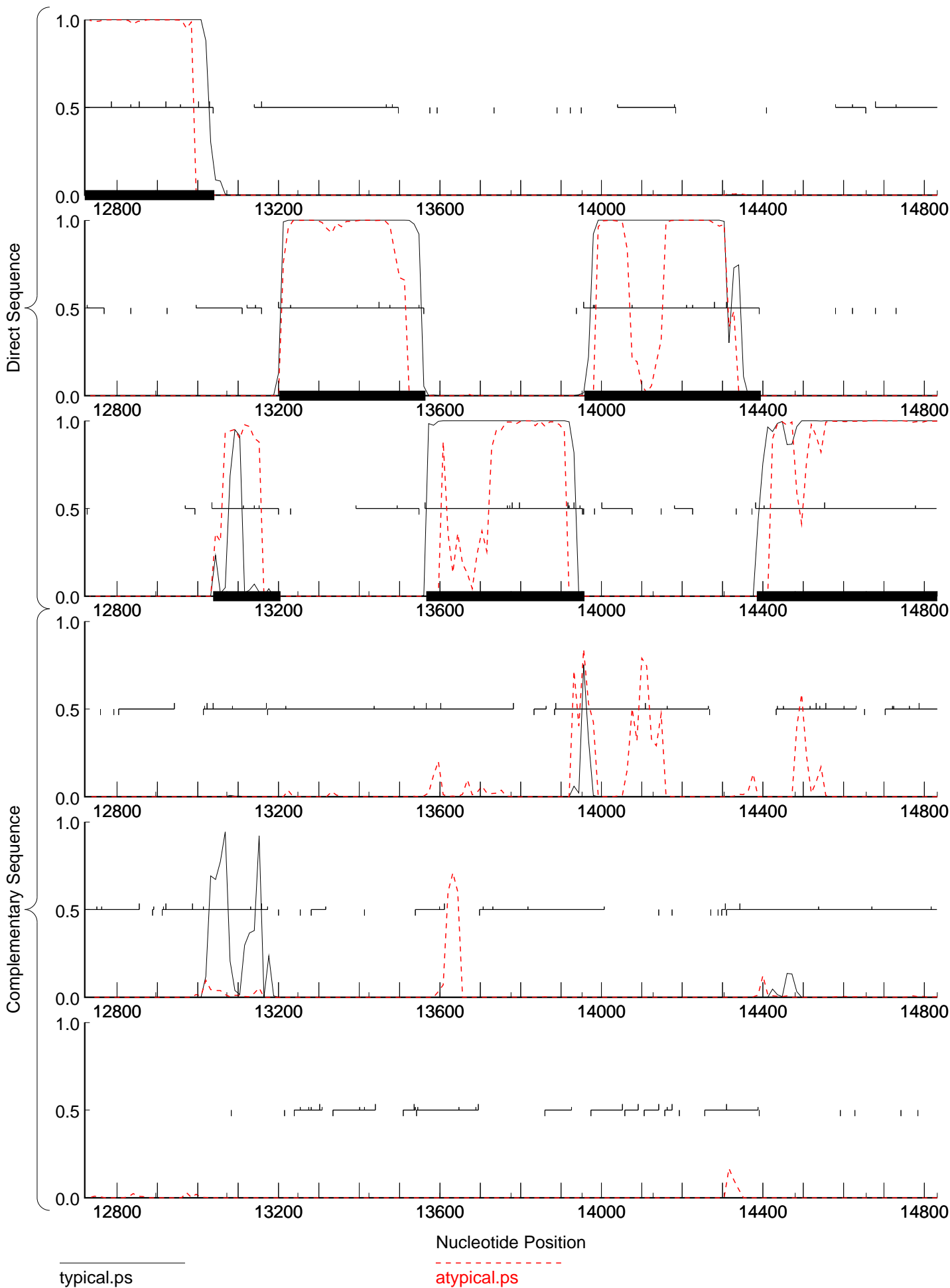


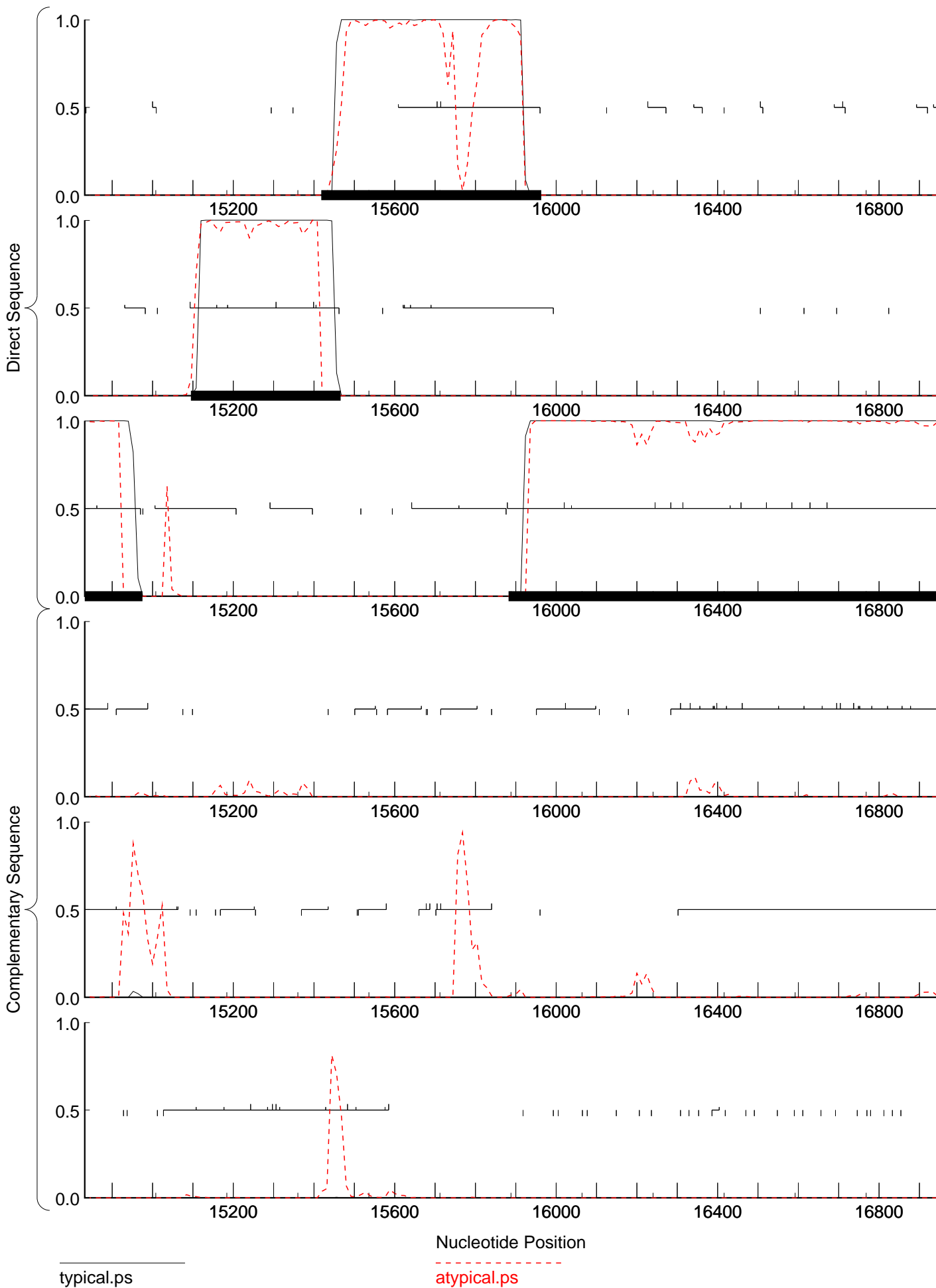






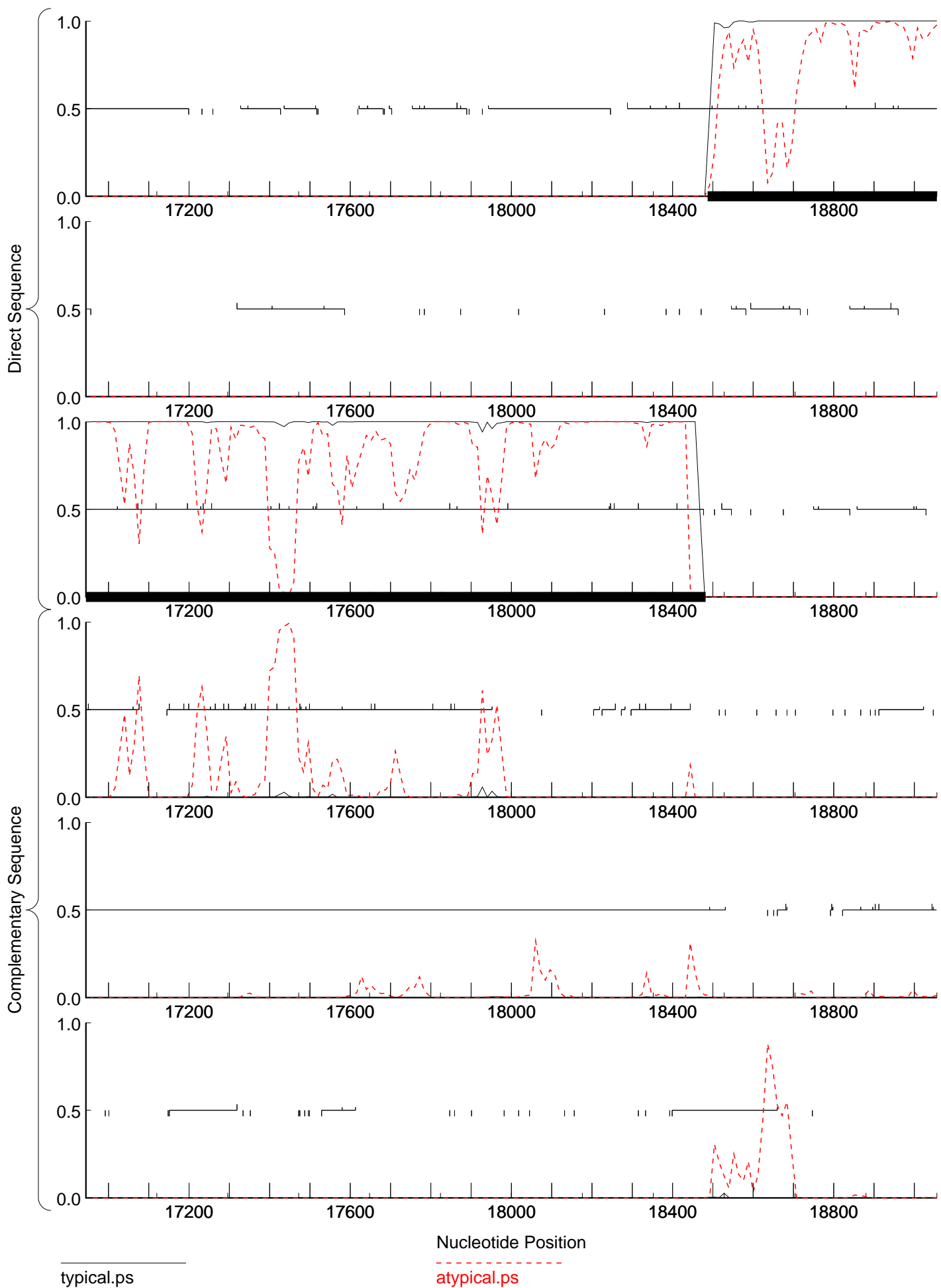






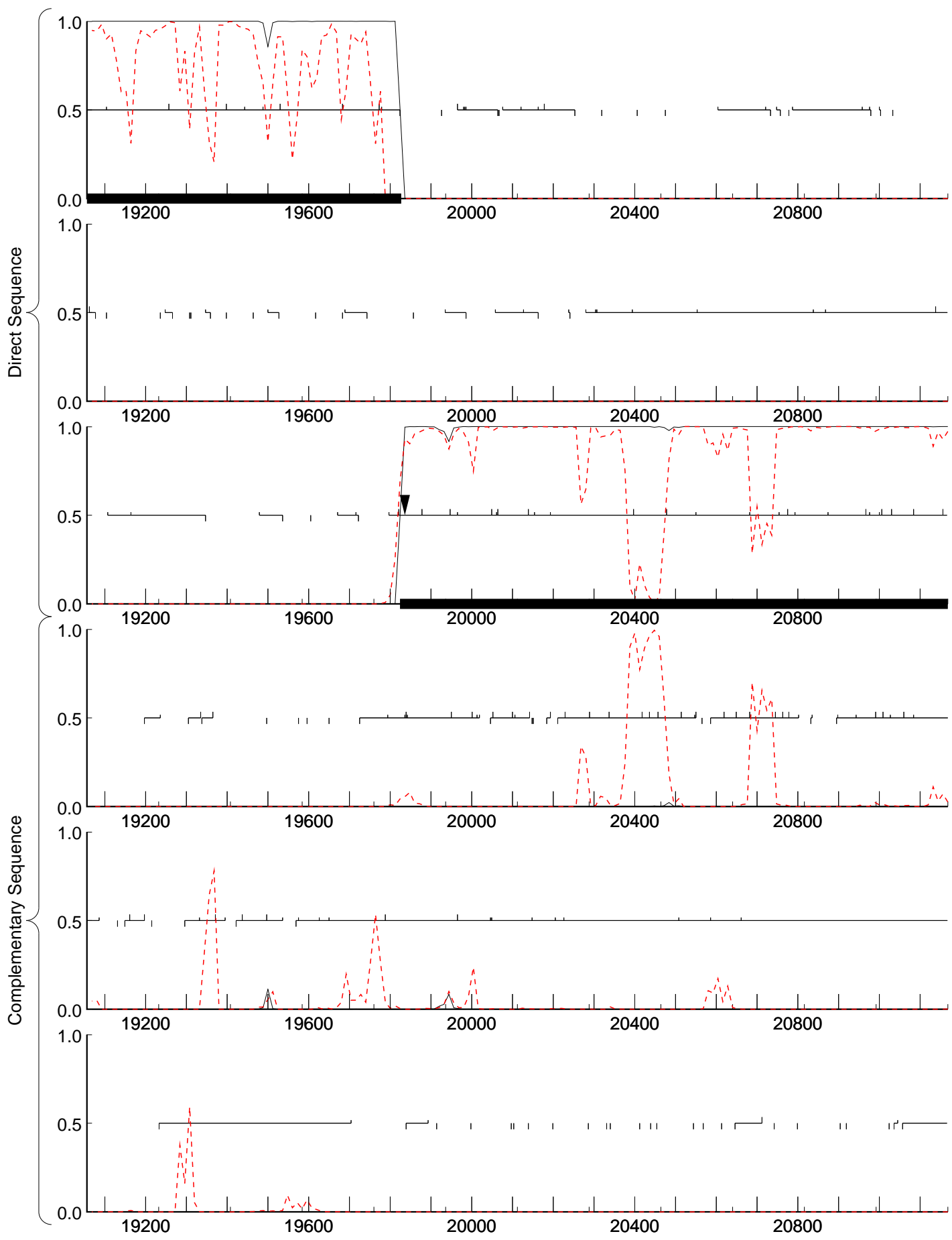
typical.ps

atypical.ps



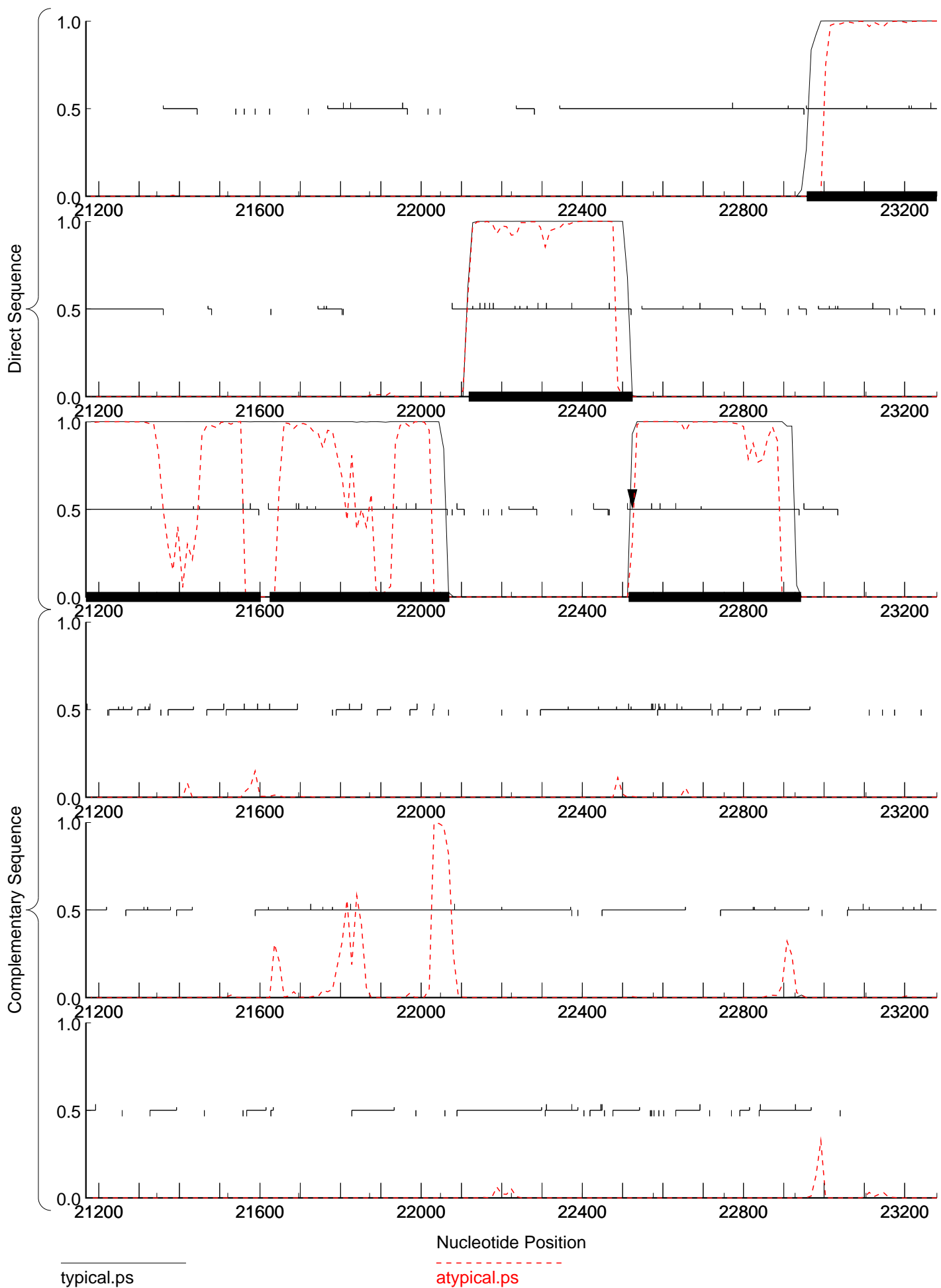
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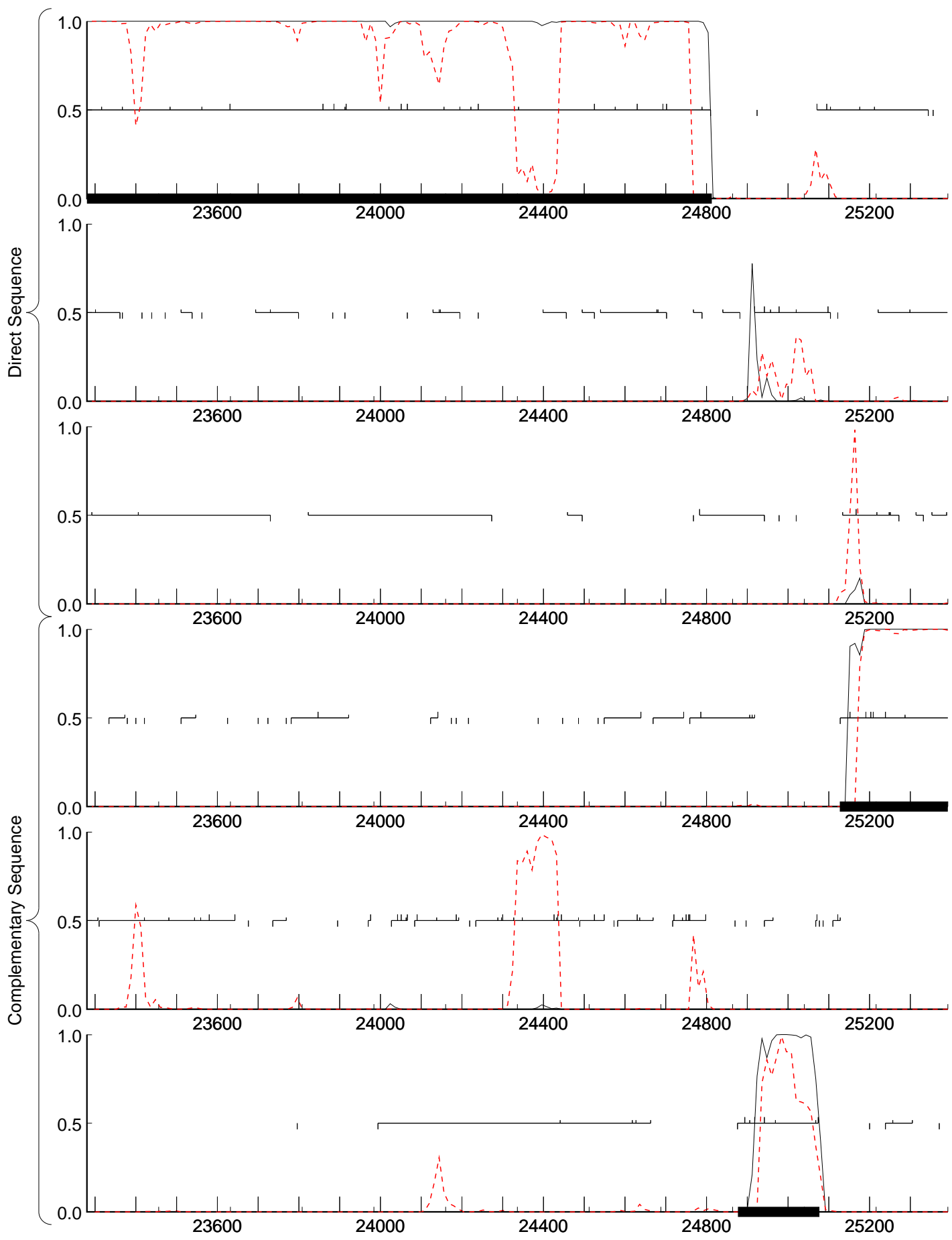
atypical.ps



typical.ps

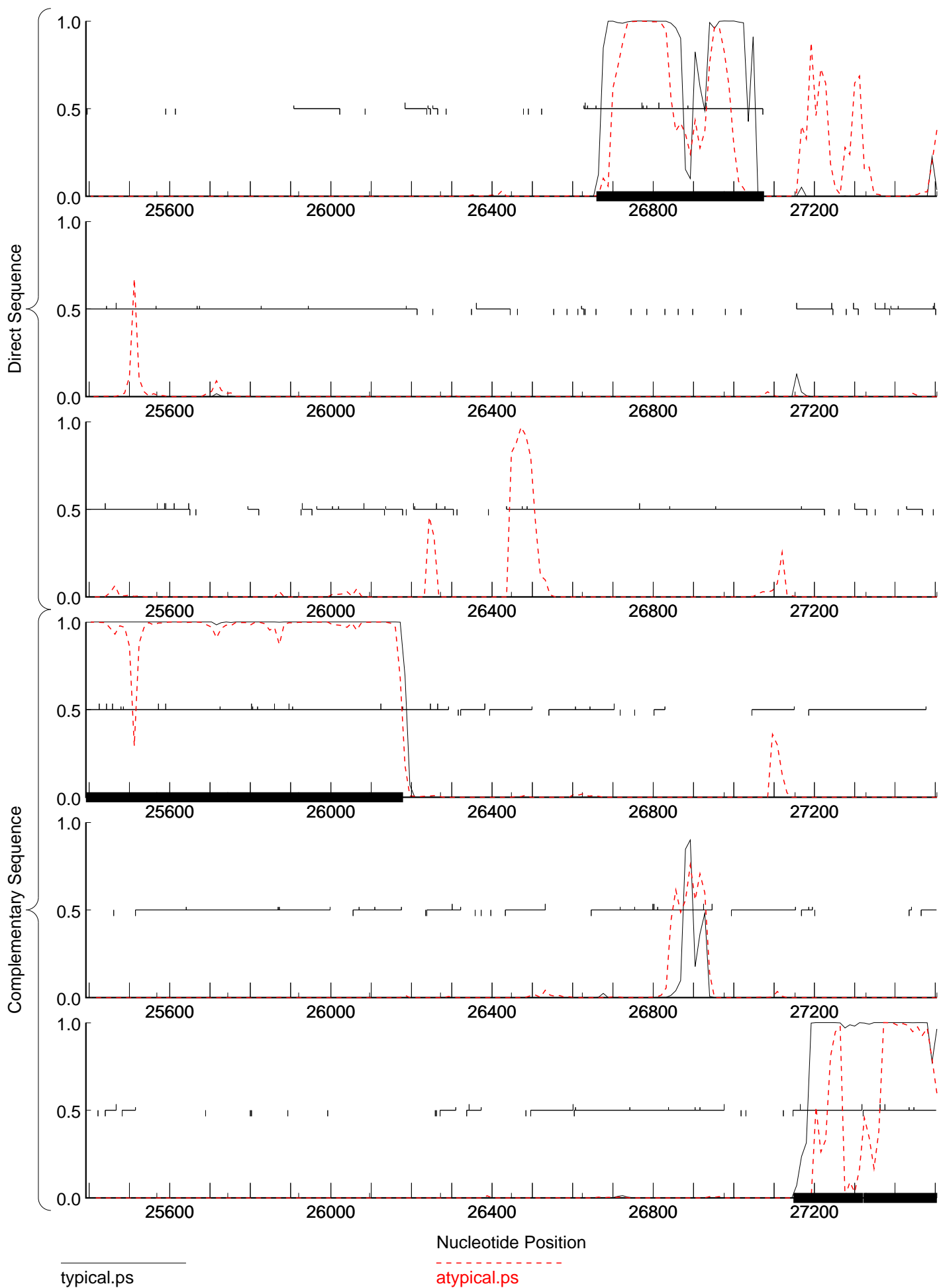
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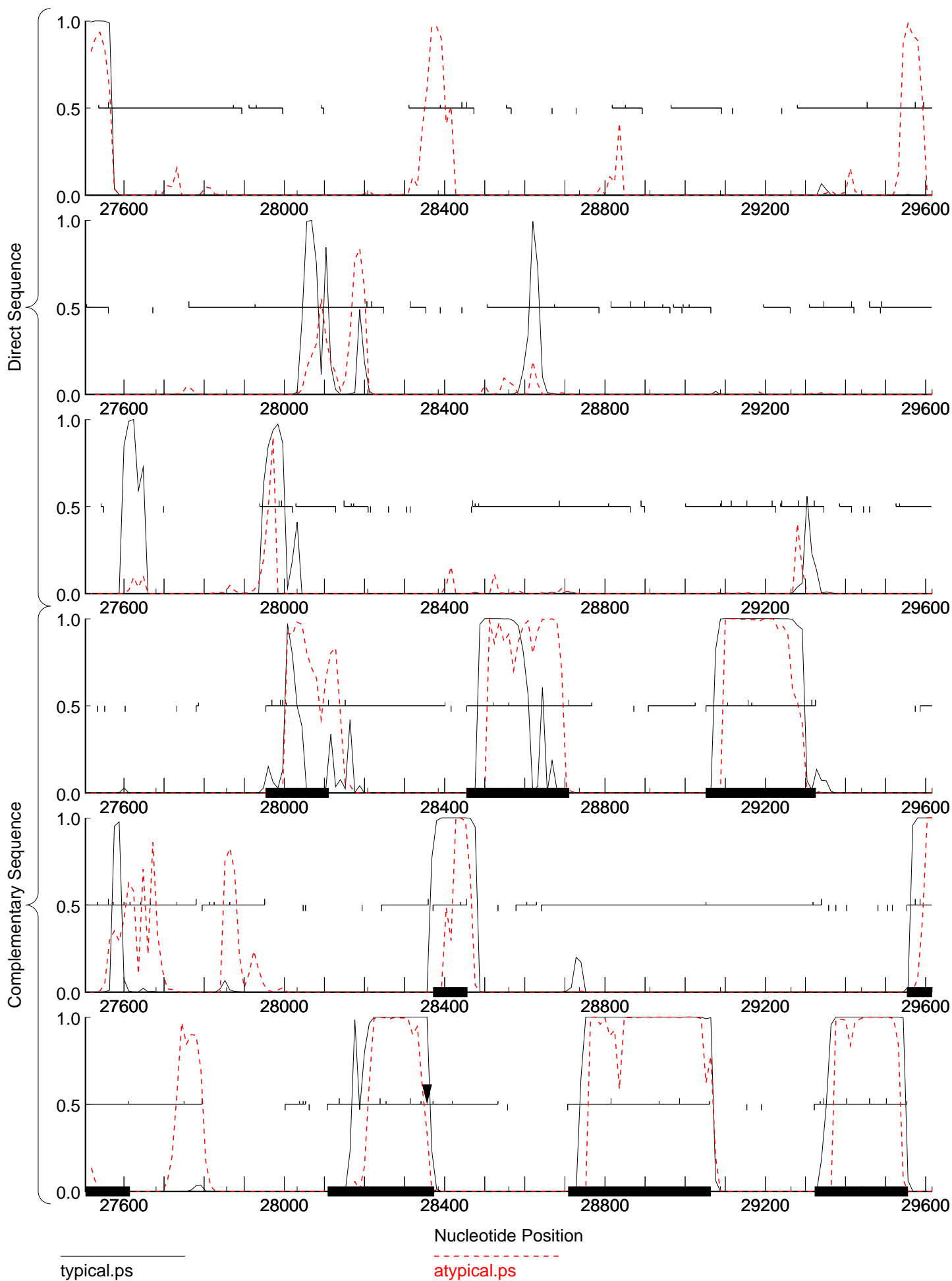


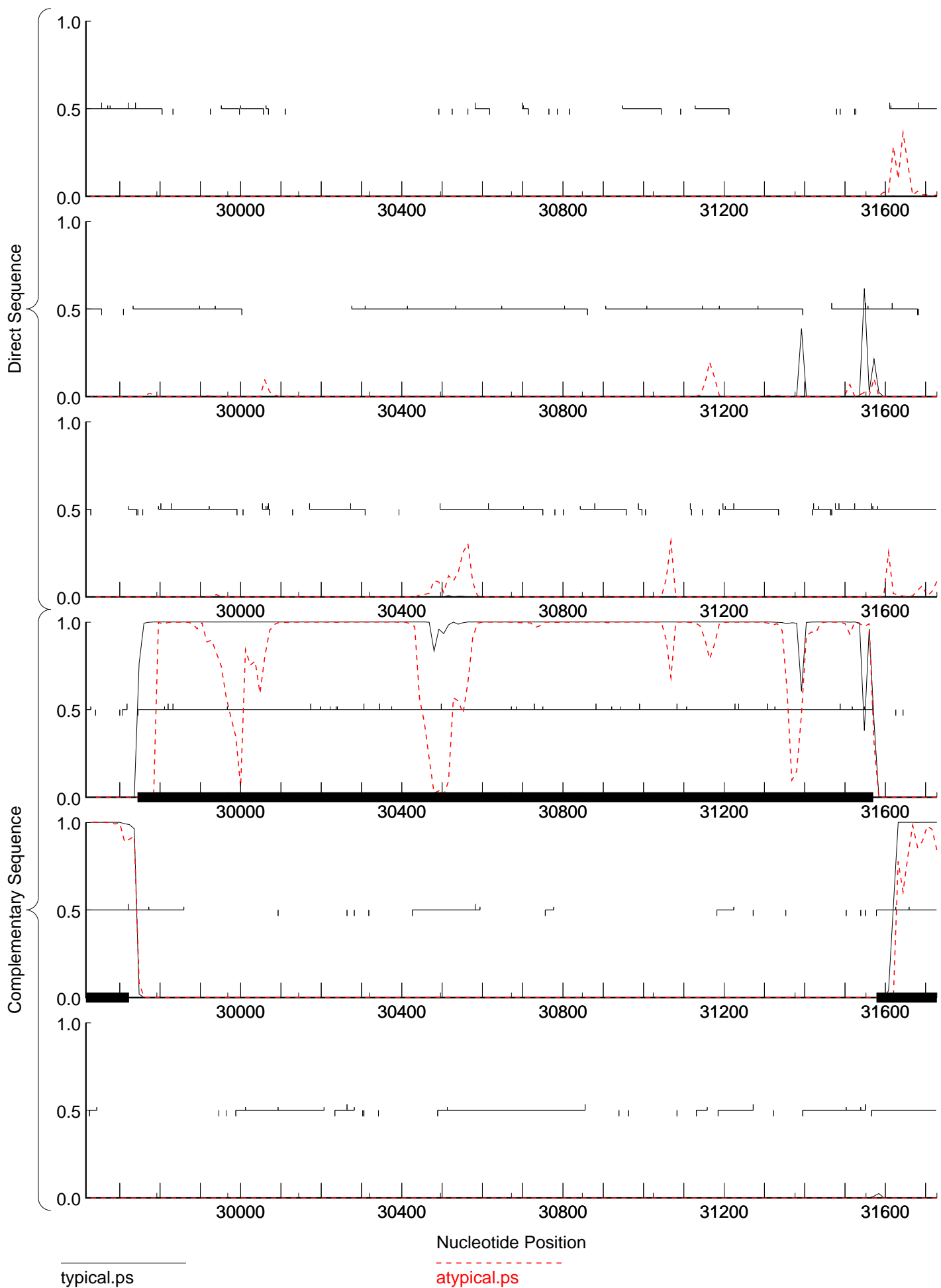


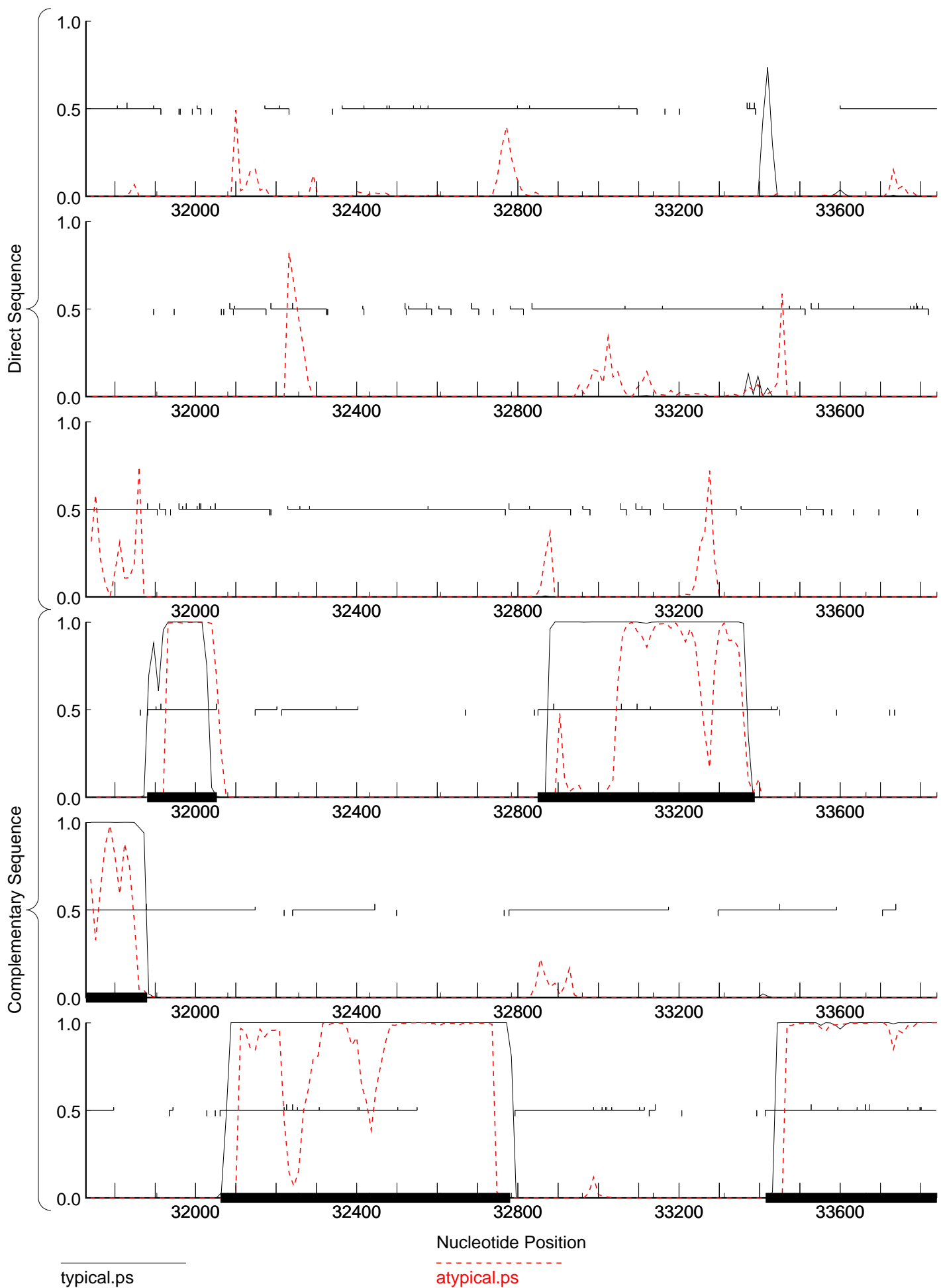
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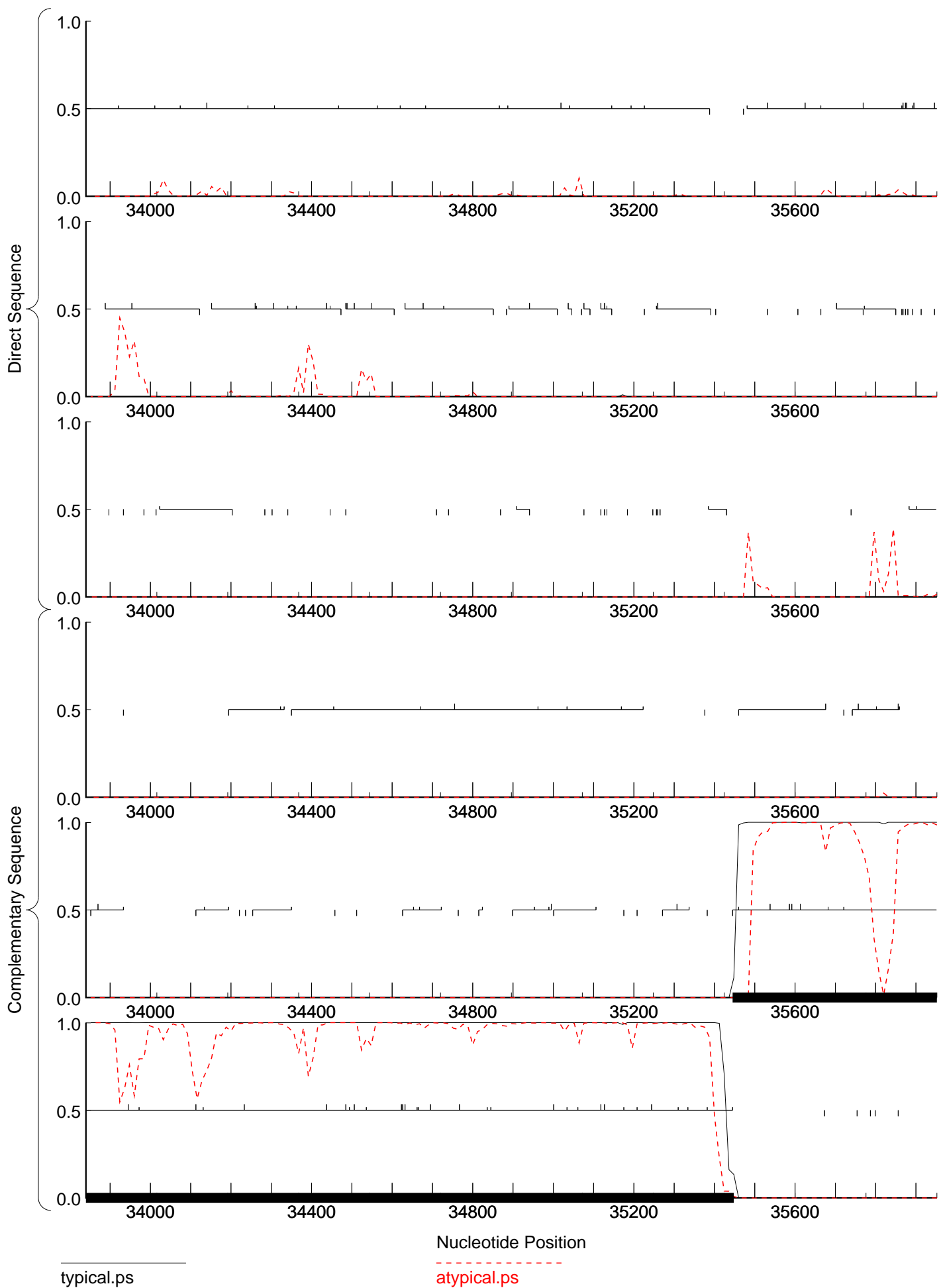
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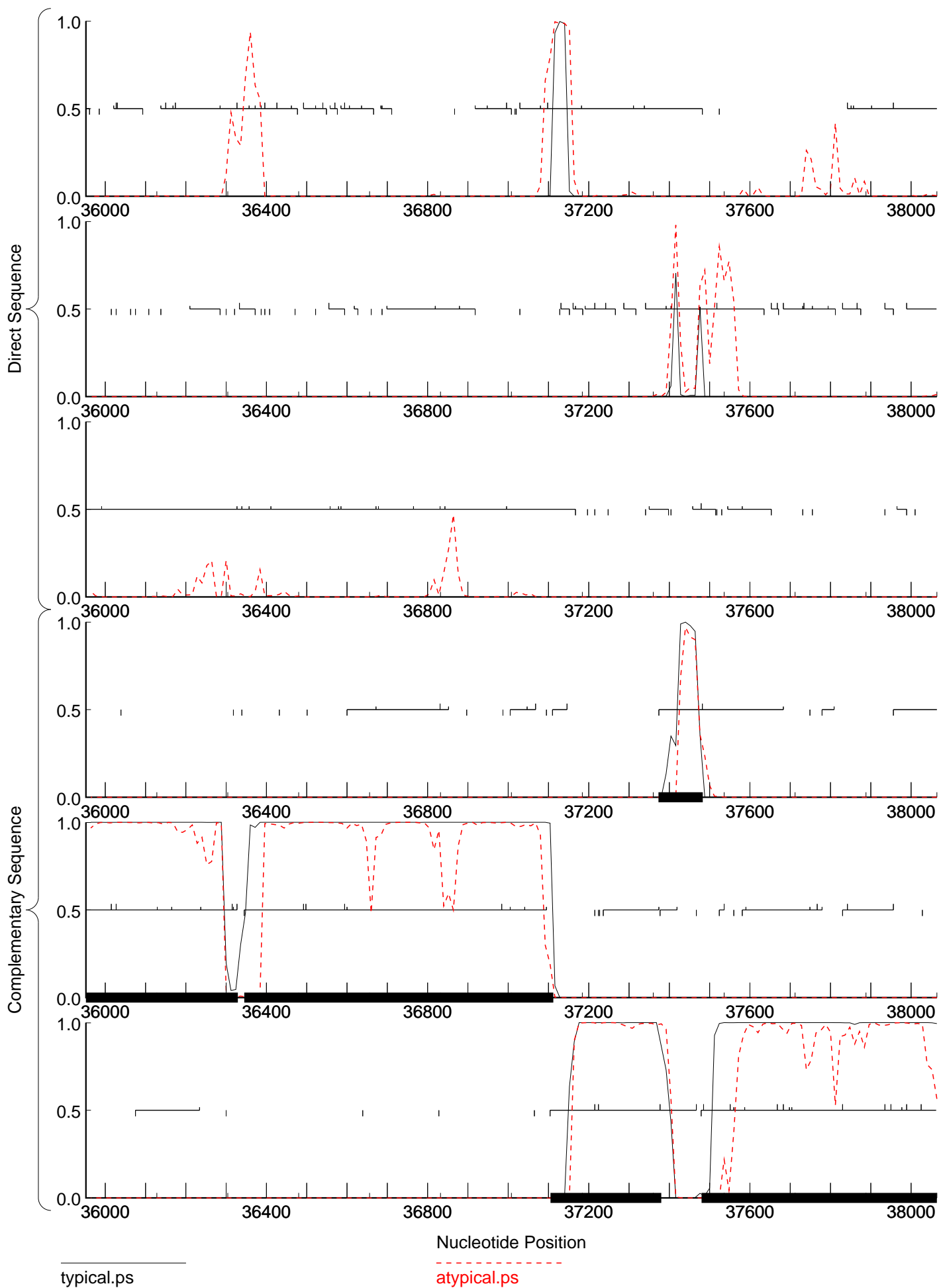






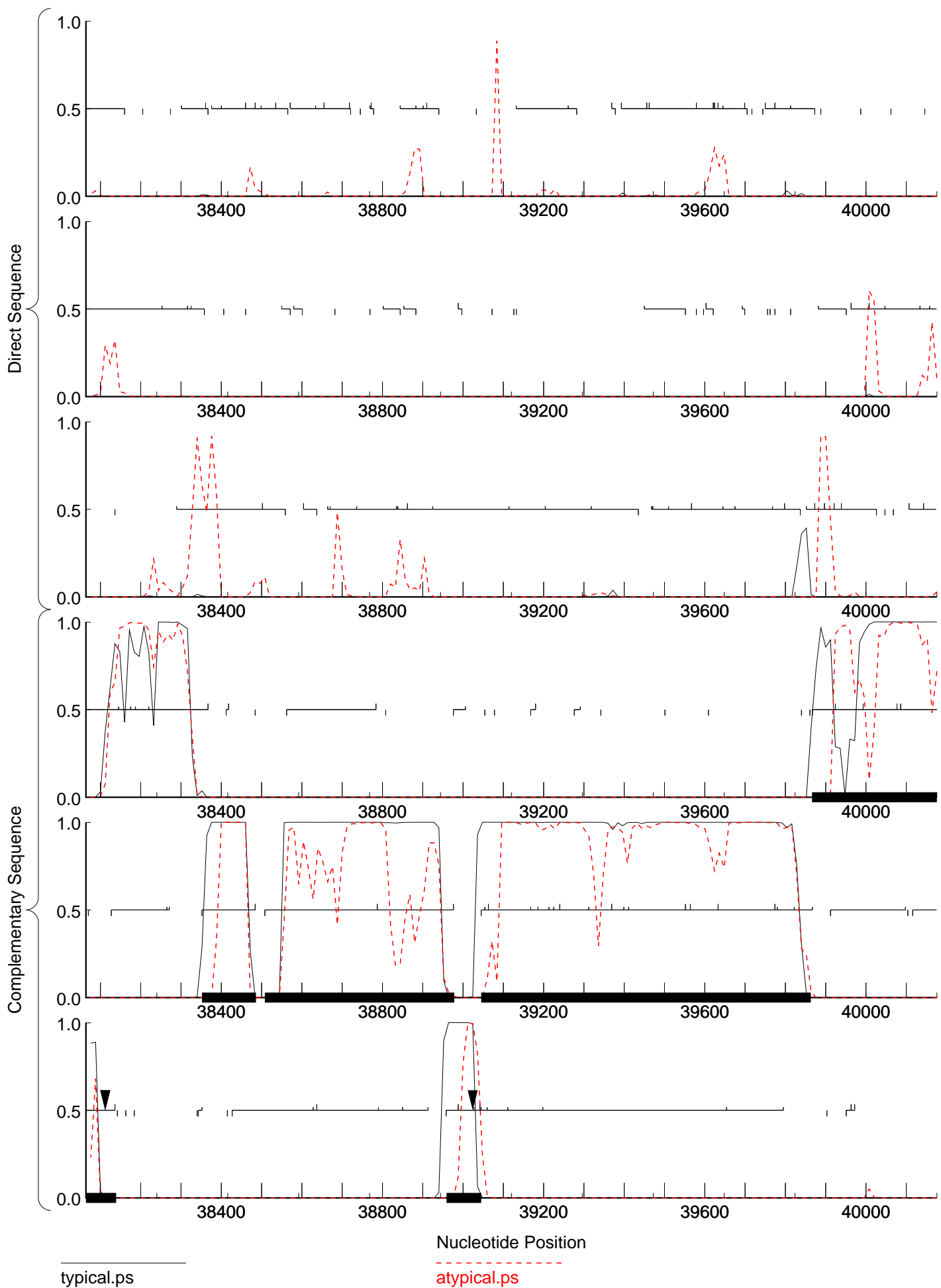






typical.ps

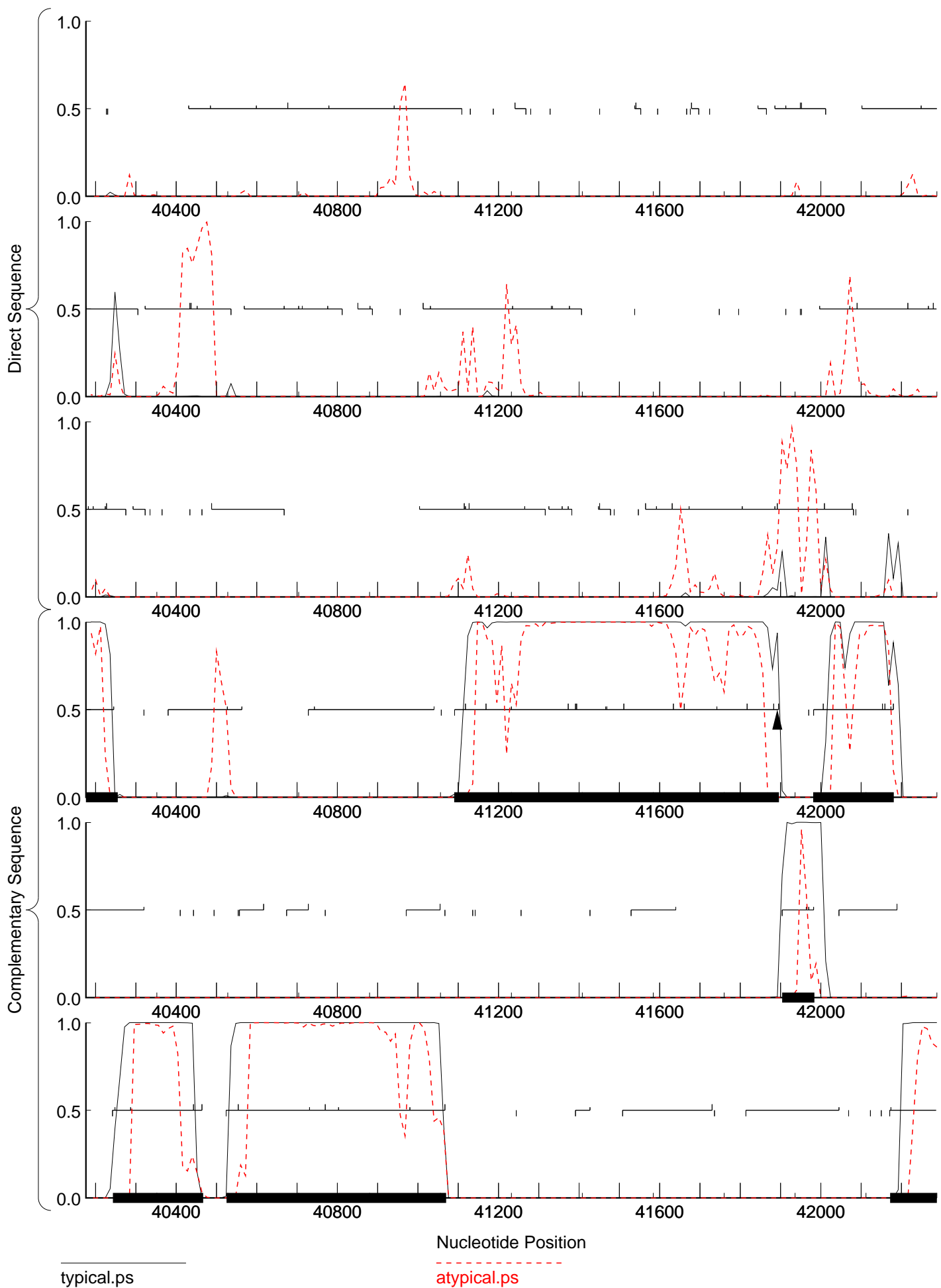
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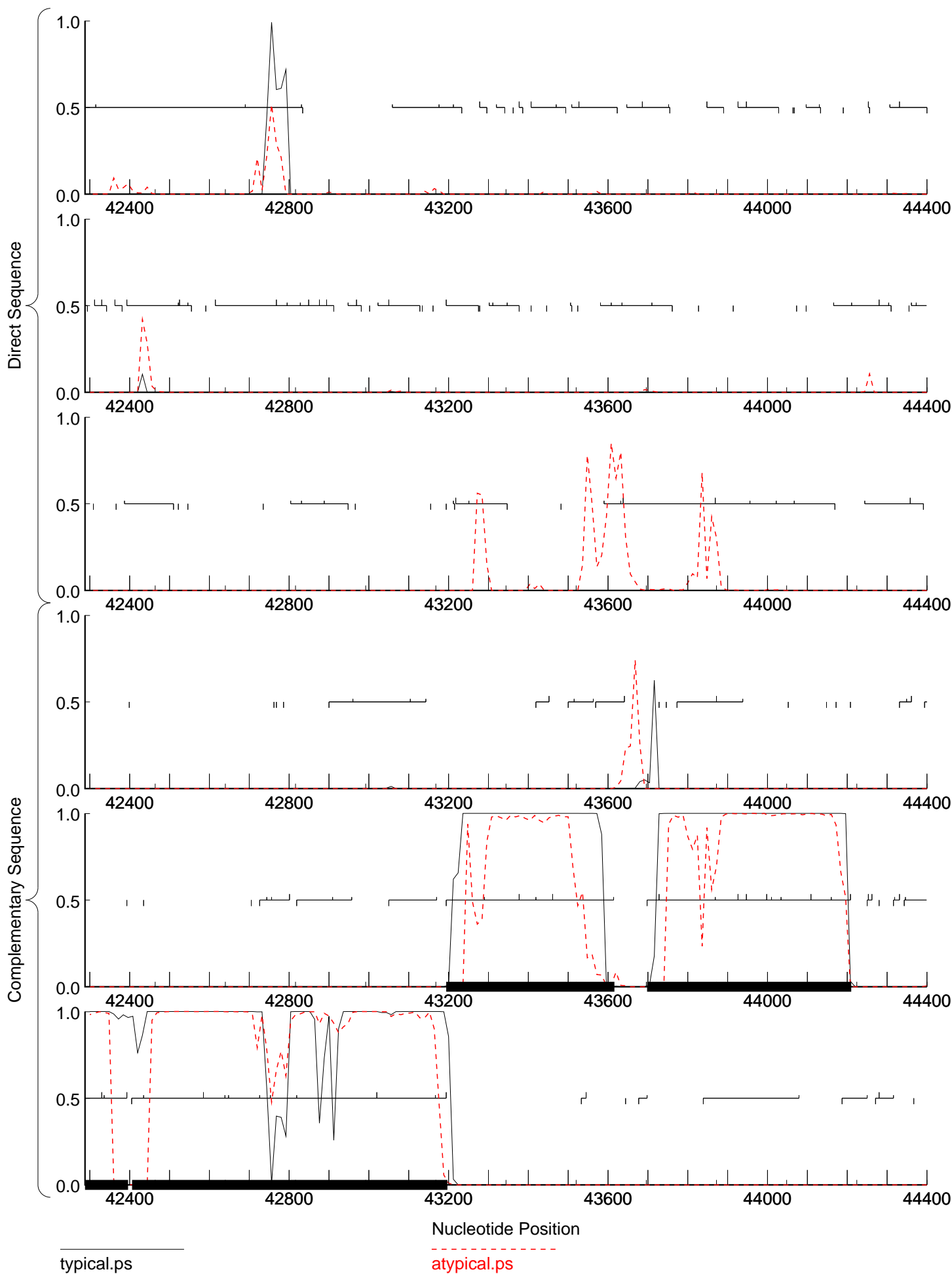


typical.ps

atypical.ps

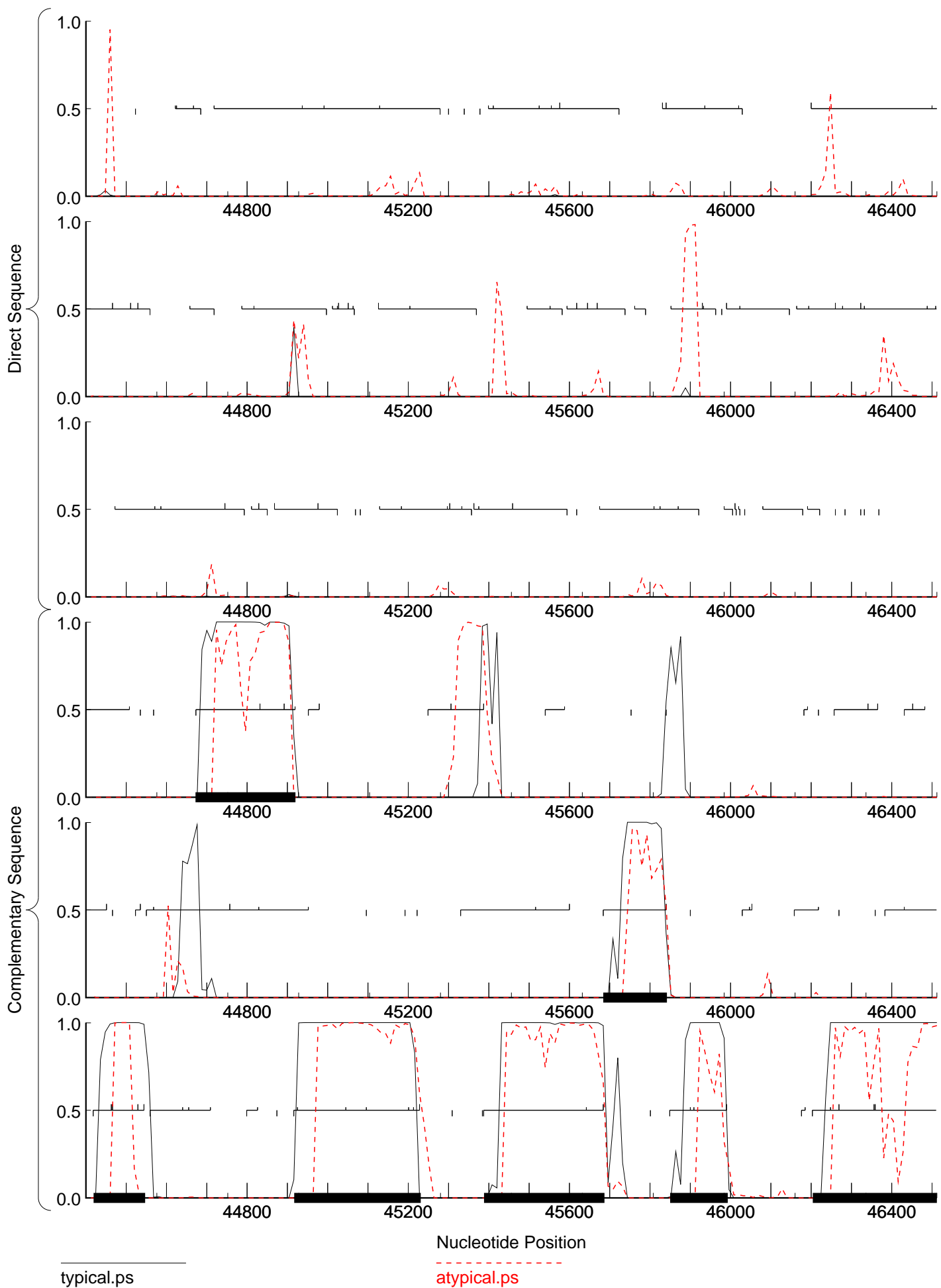
GeneMark.hmm prediction





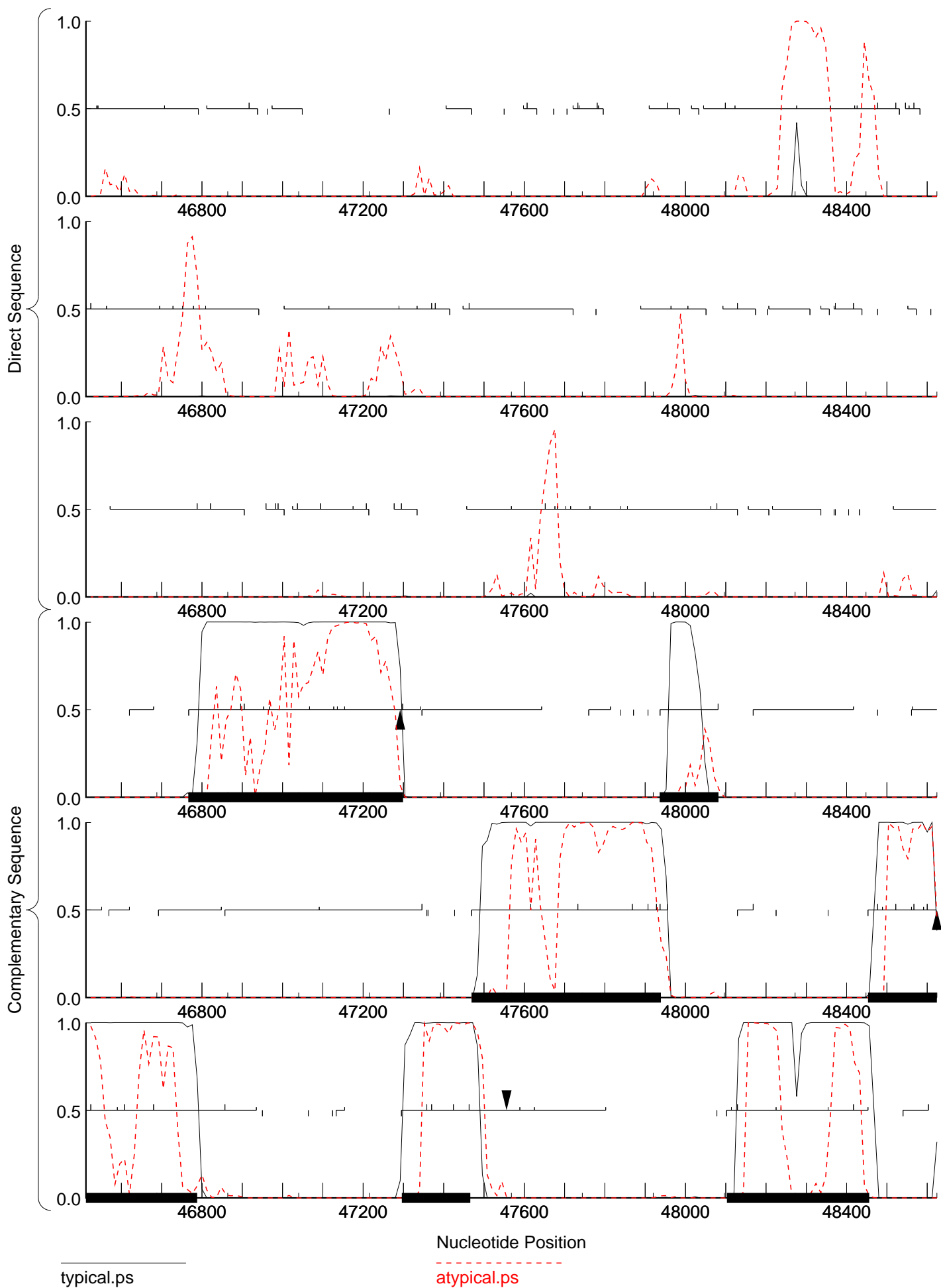
typical.ps

atypical.ps



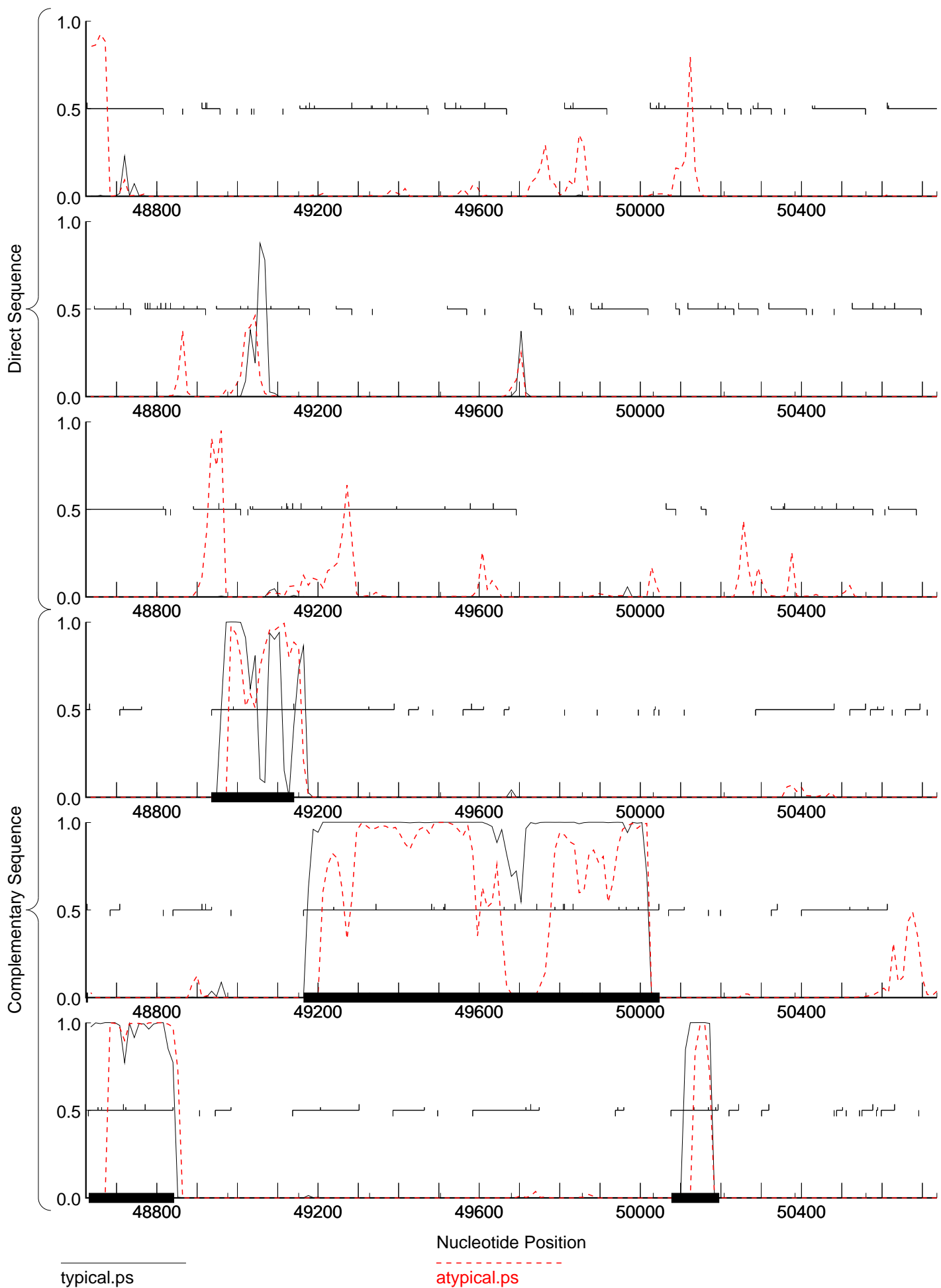
typical.ps

atypical.ps



typical.ps

atypical.ps



typical.ps

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