

GeneMark

Version 2.5p (09.08.06)

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PROGRAM INFORMATION

Sequence : Mycobacterium phage Pistachio complete sequence, 50006 bp including 10-base 3' overhang (CG
Analysis Date : 3/21/20 at 15:54:37
Pages : 25
Sequence Length : 50006 bp
GC Content : 63.78%

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,

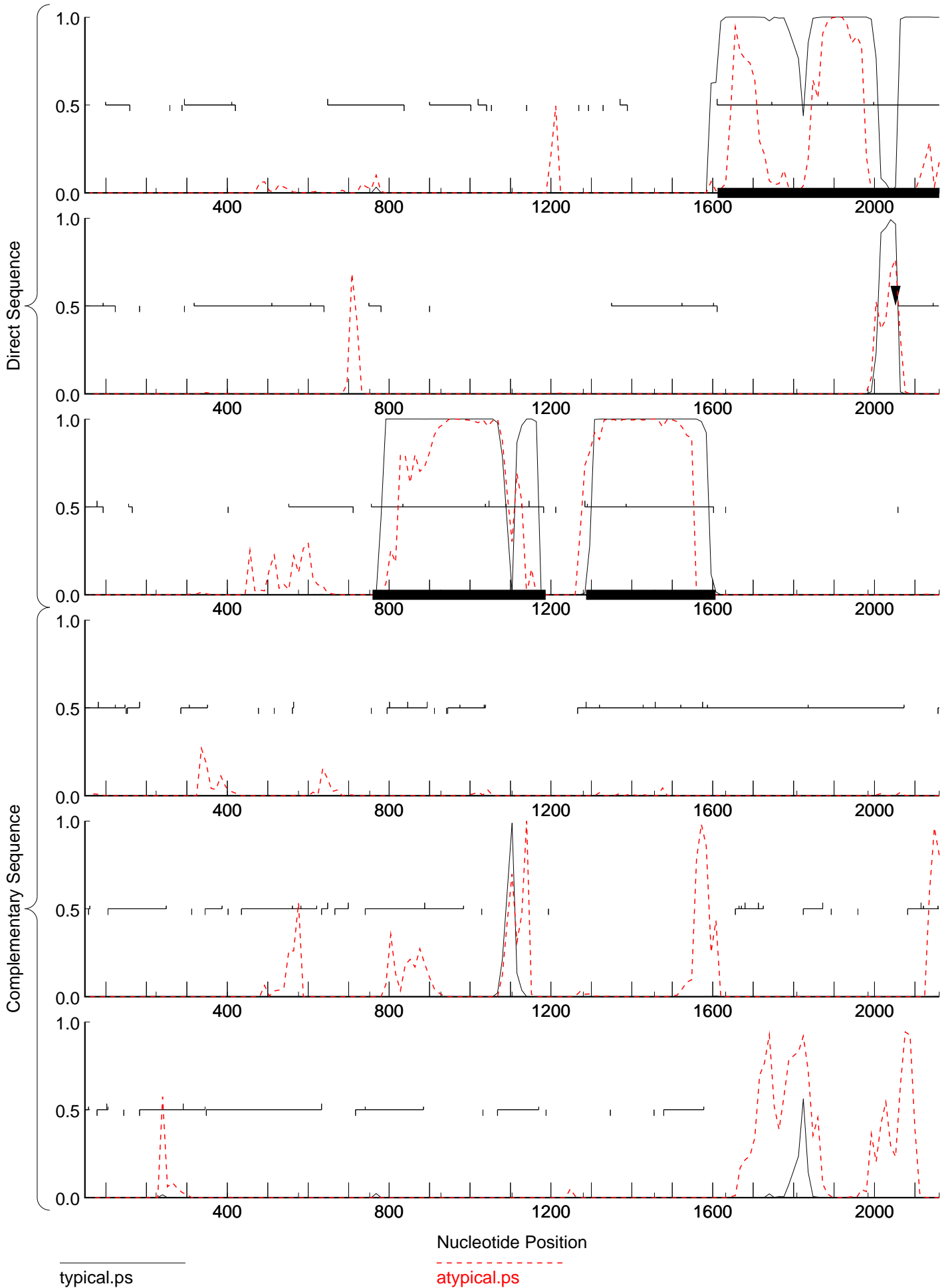
MATRIX INFORMATION

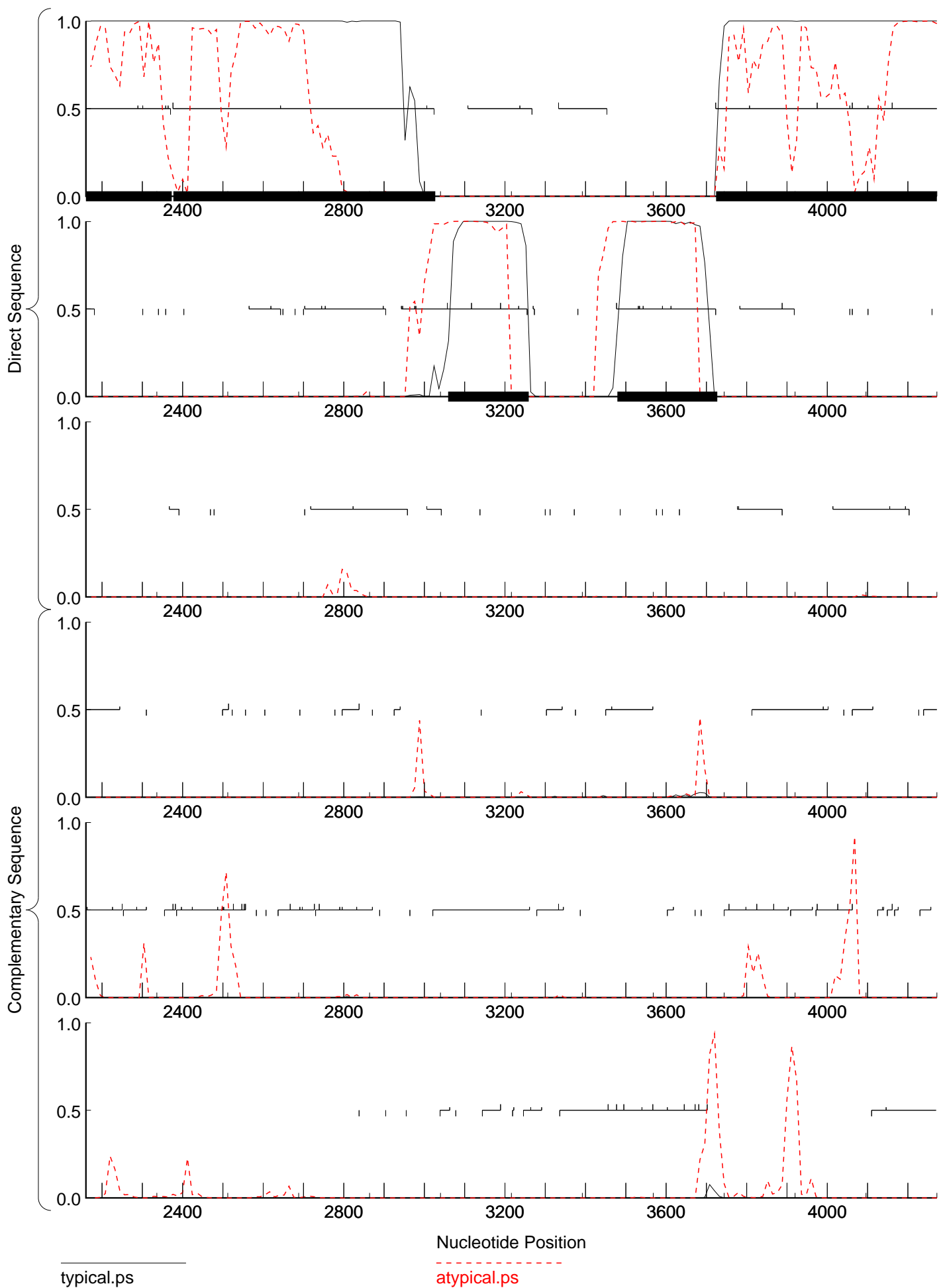
Matrix : -
Author : -
Order : 4

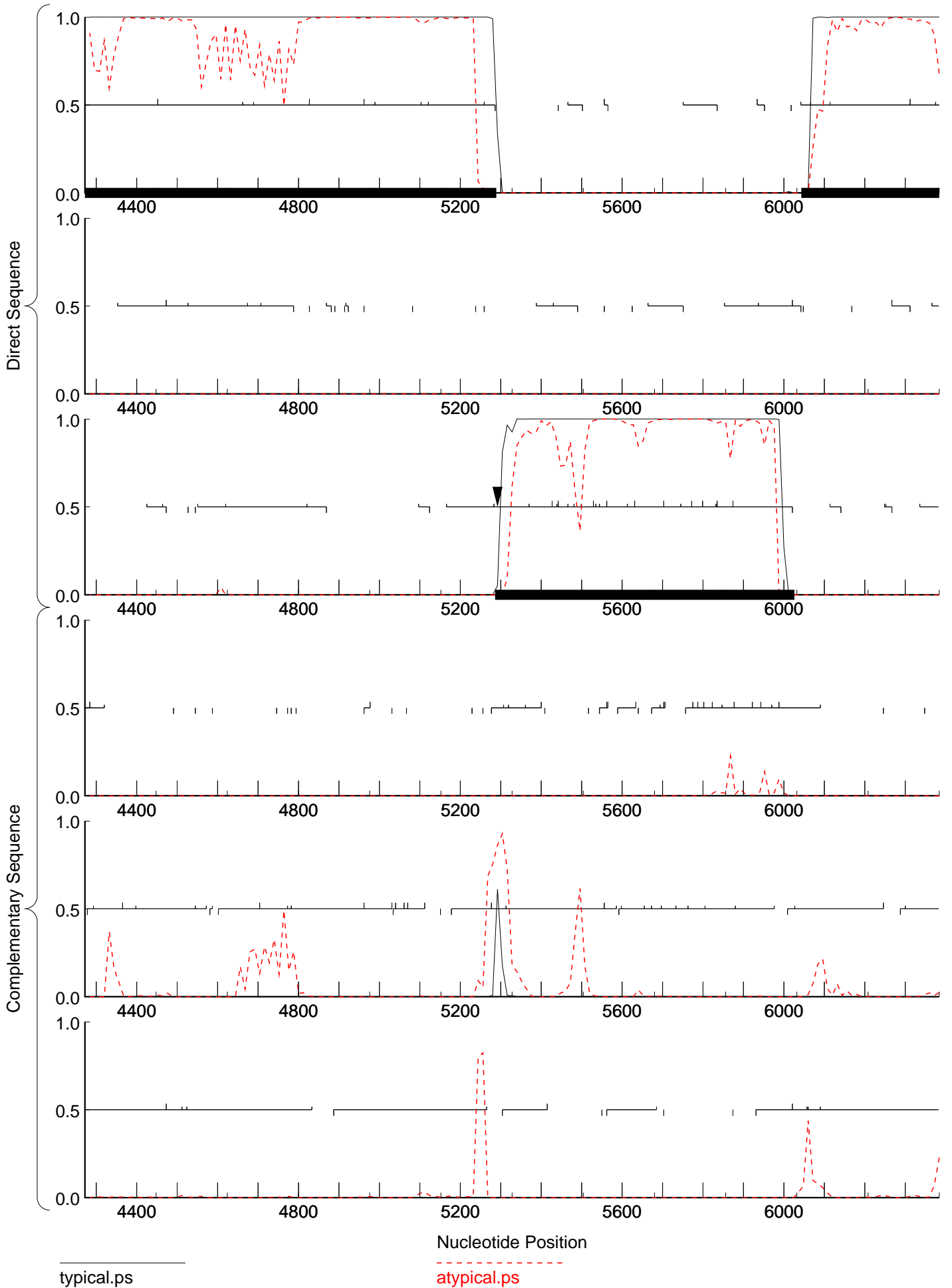
Send questions / comments to:
Dr. M. Borodovsky
Georgia Institute of Technology
School of Biology
Atlanta, GA 30332-0230

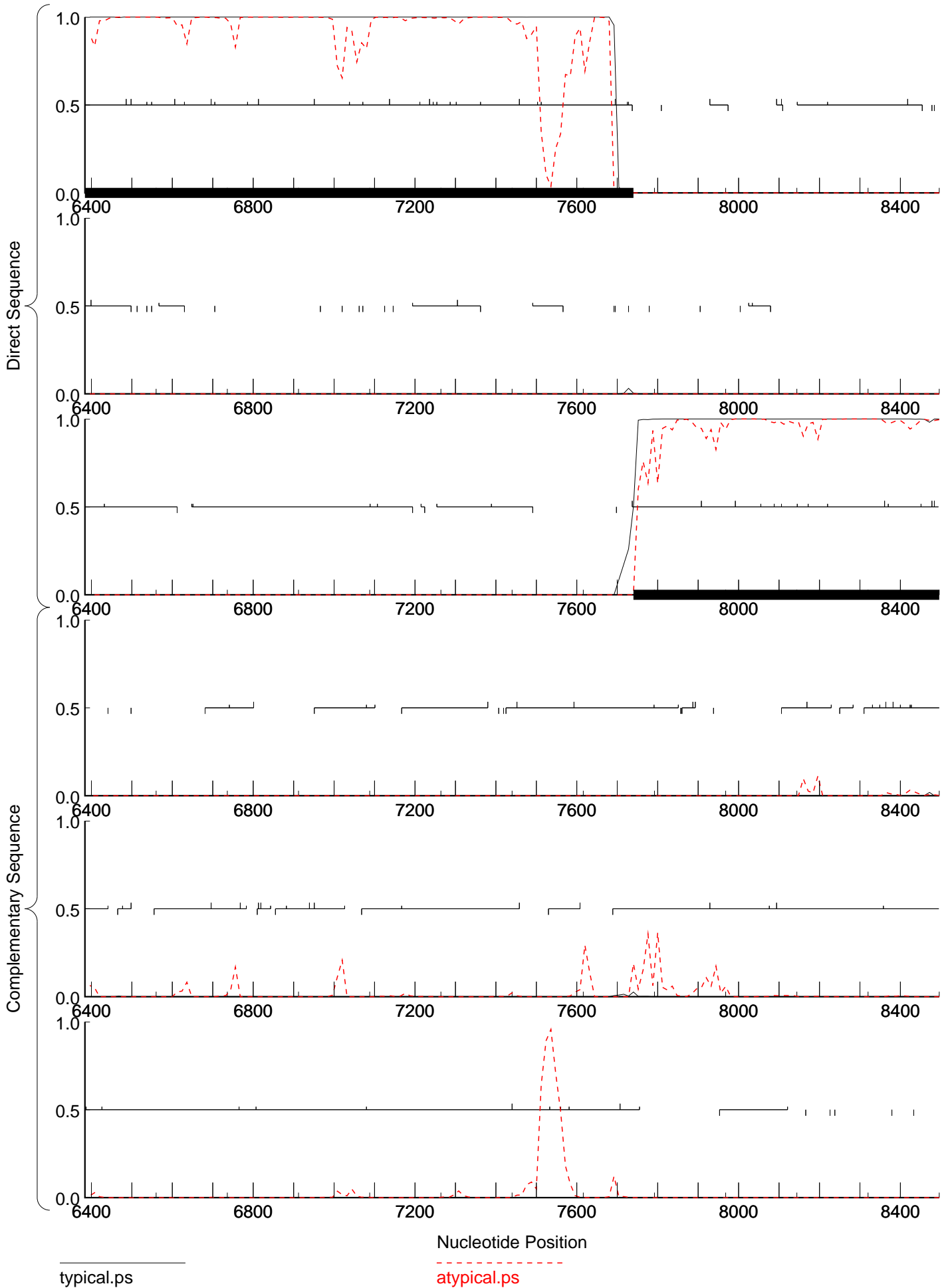
Matrix notes & comments

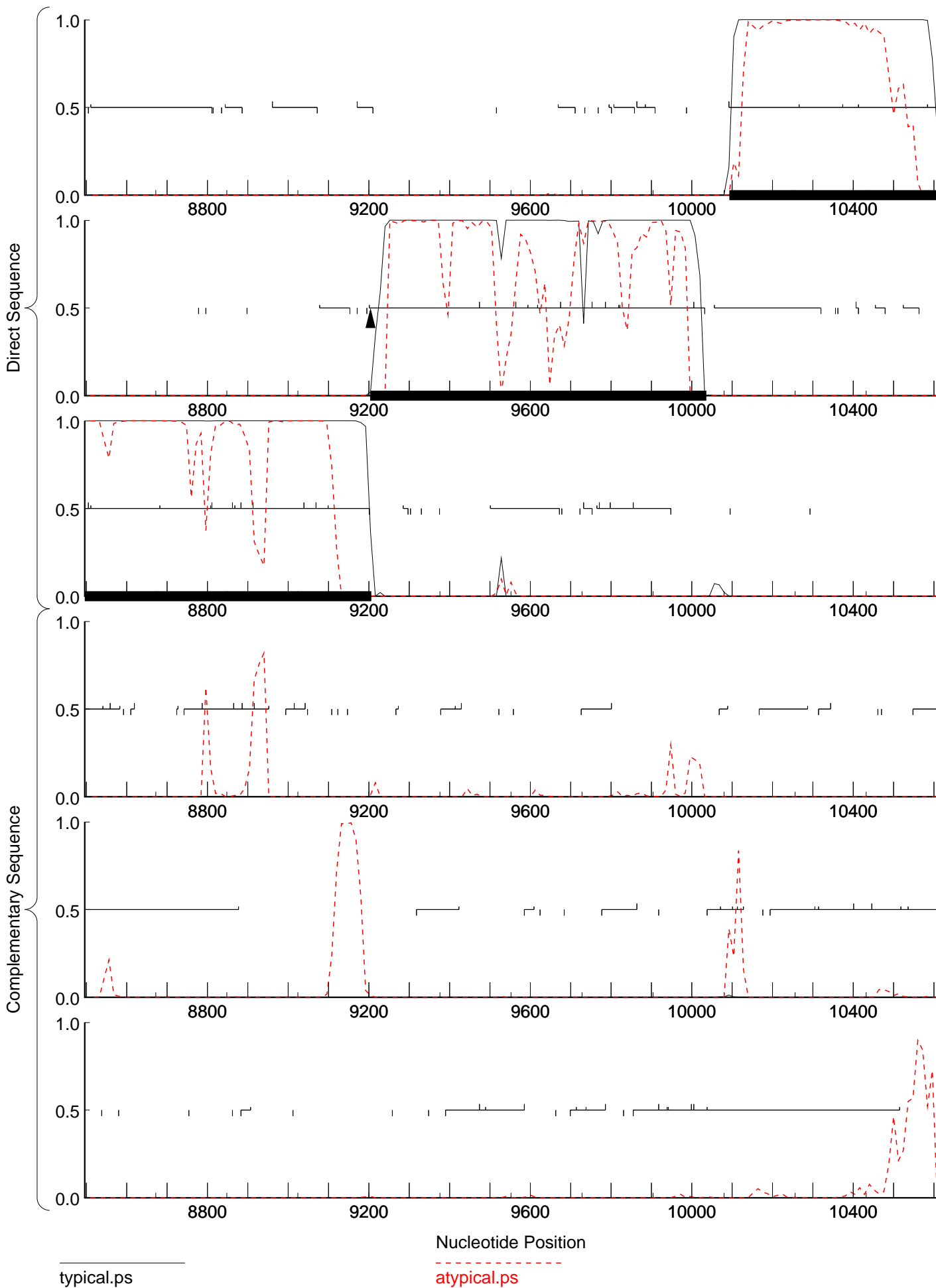
Training set derived by GeneMarkS, 4.30 October 2014
Sat Mar 21 15:54:37 2020

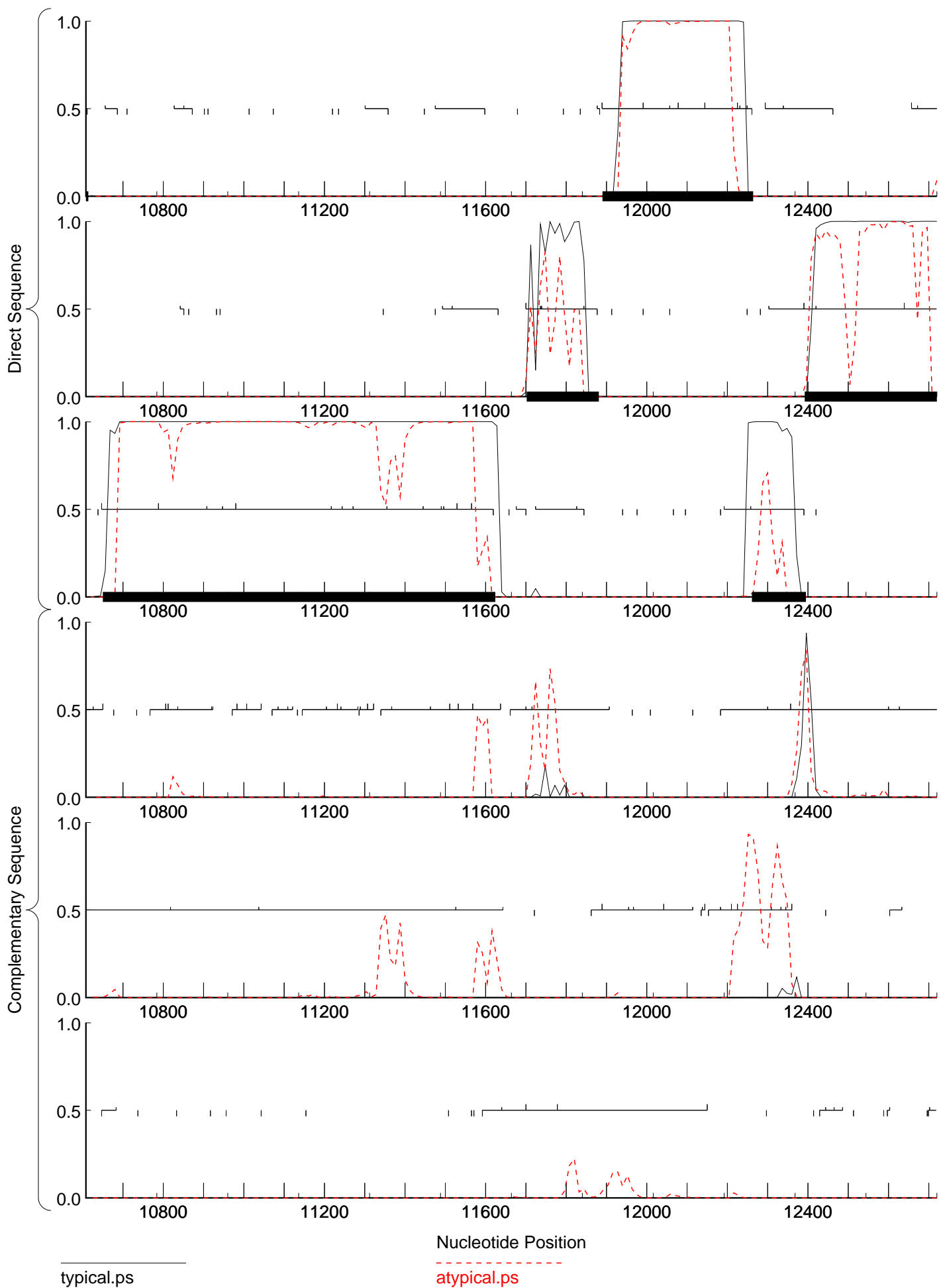






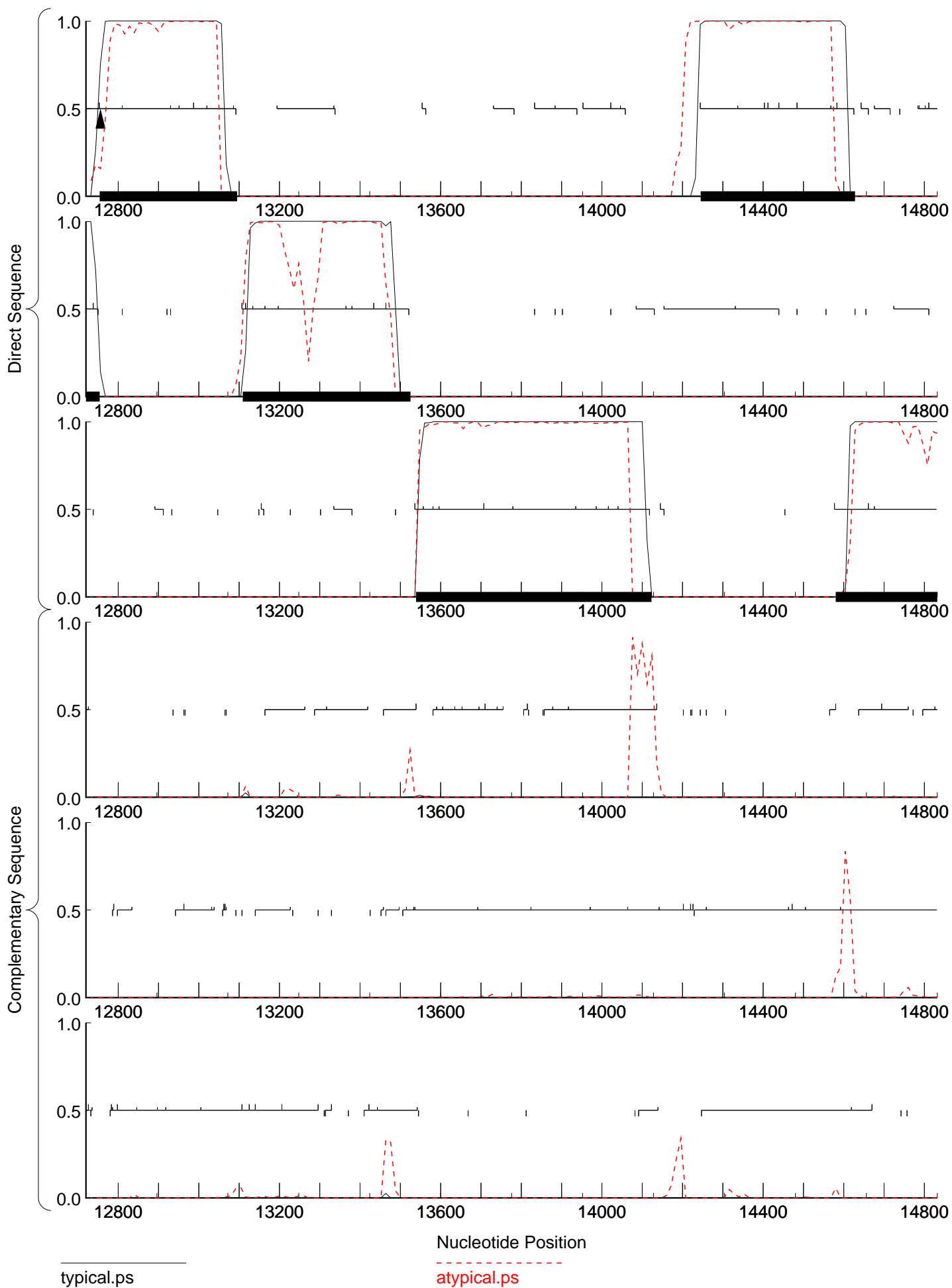






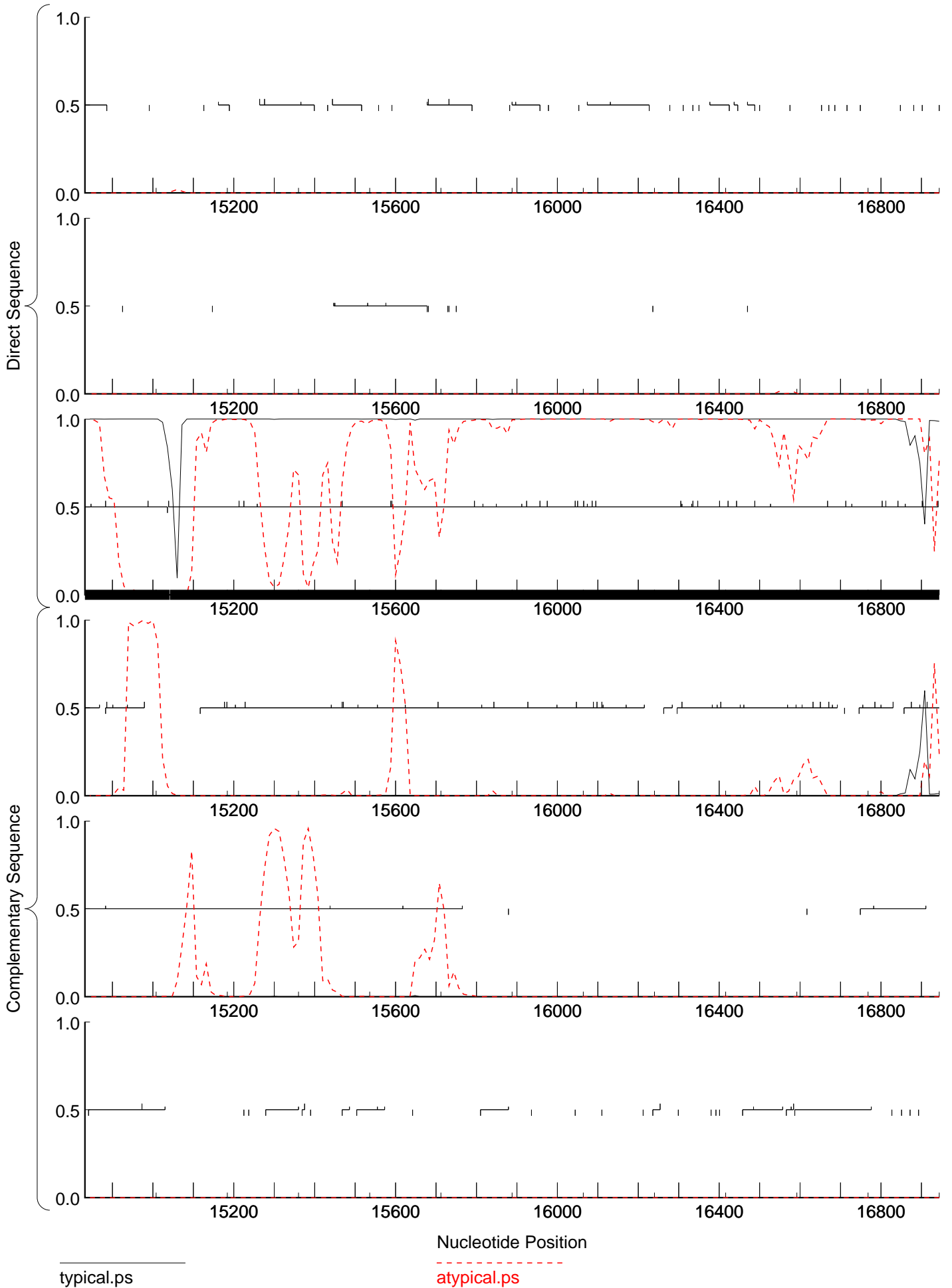
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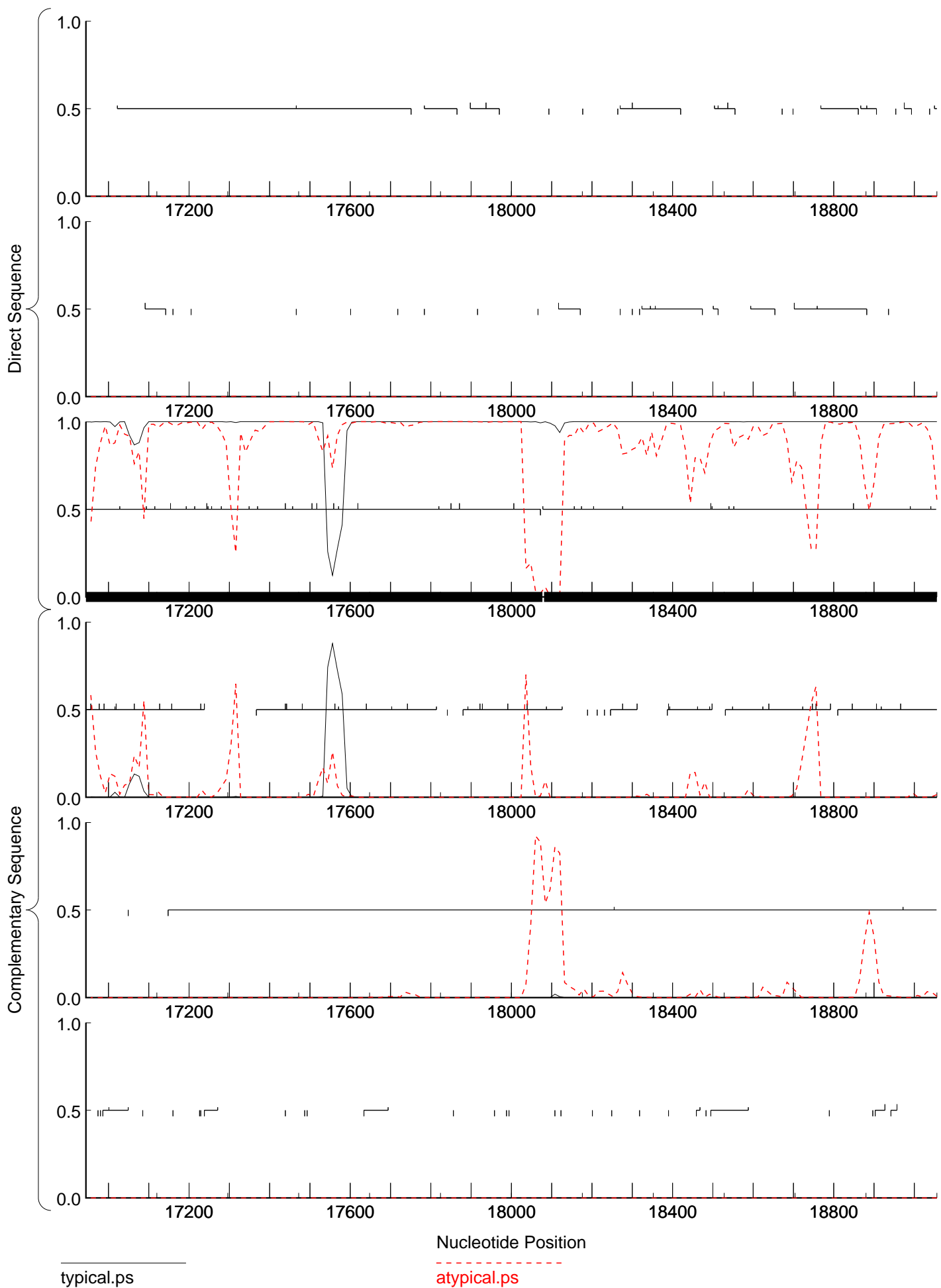
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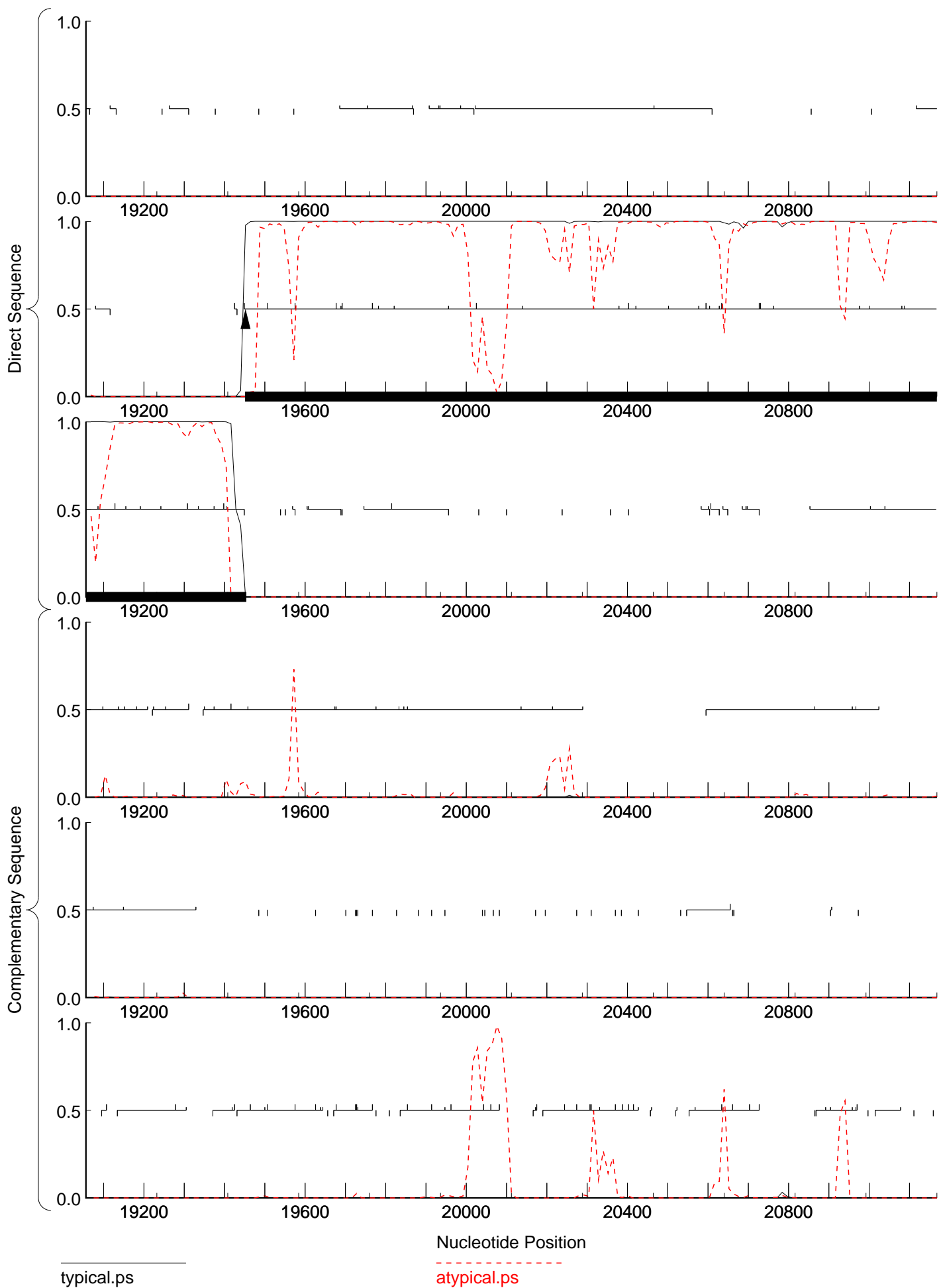


typical.ps

atypical.ps

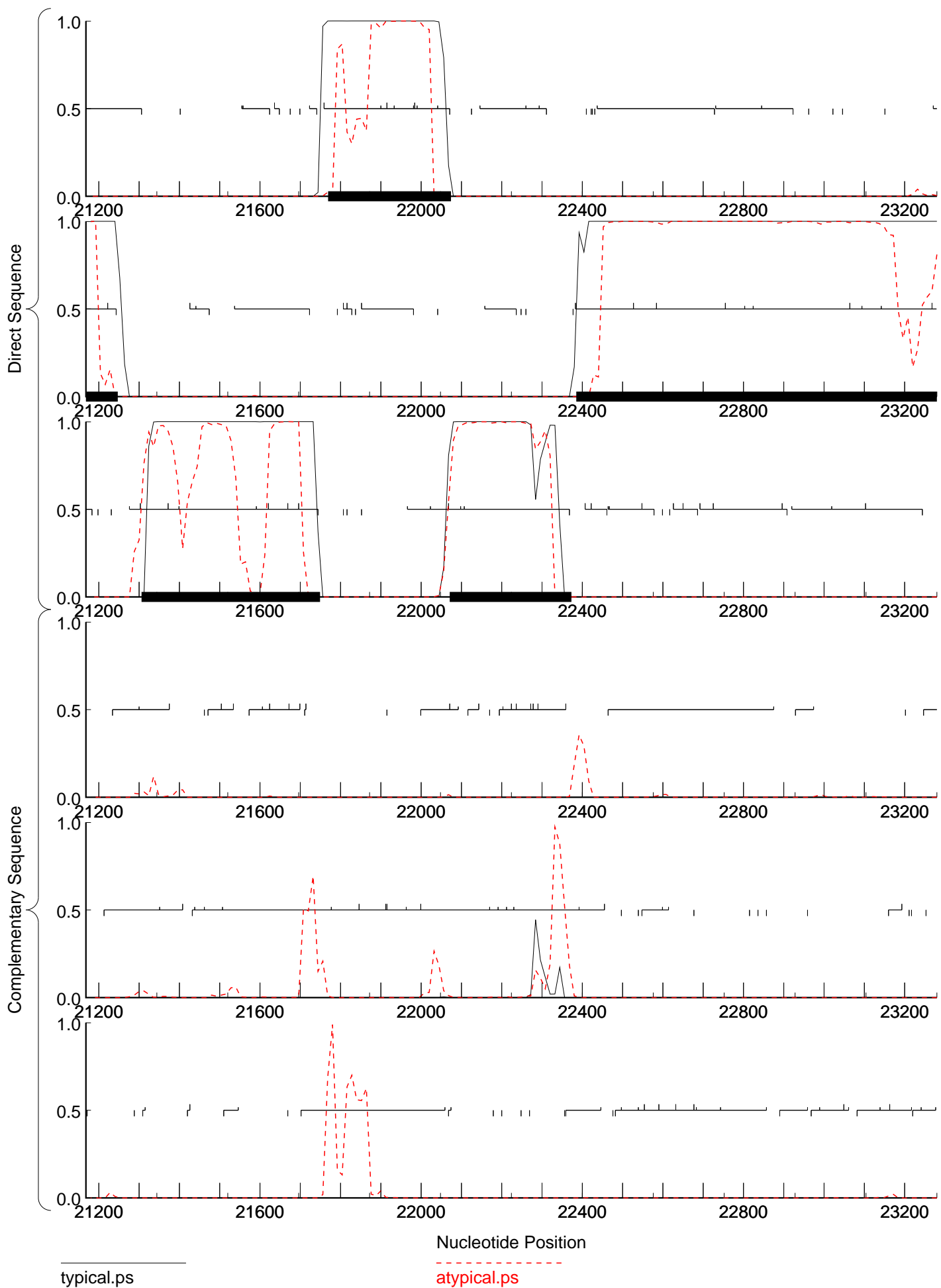






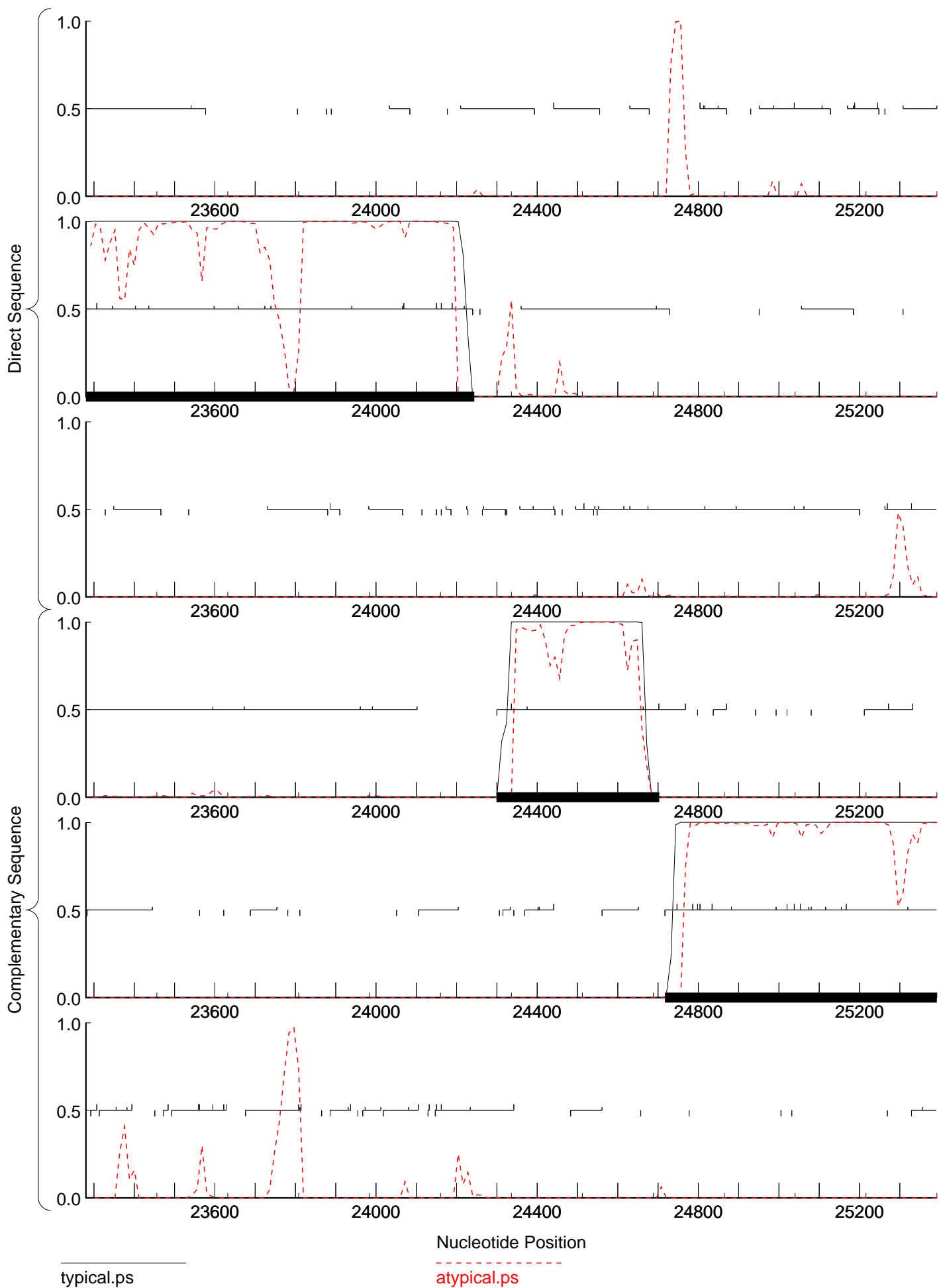
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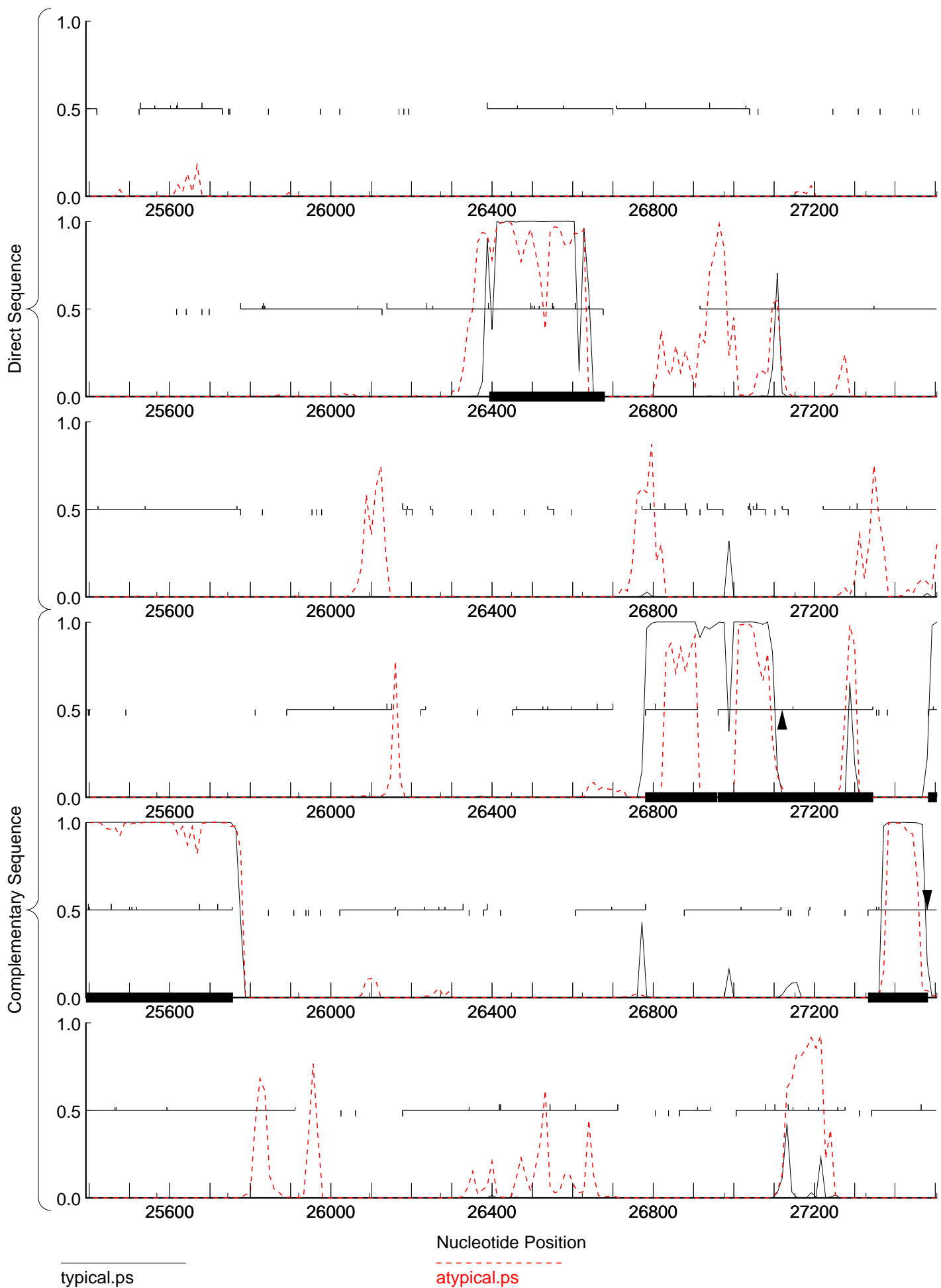
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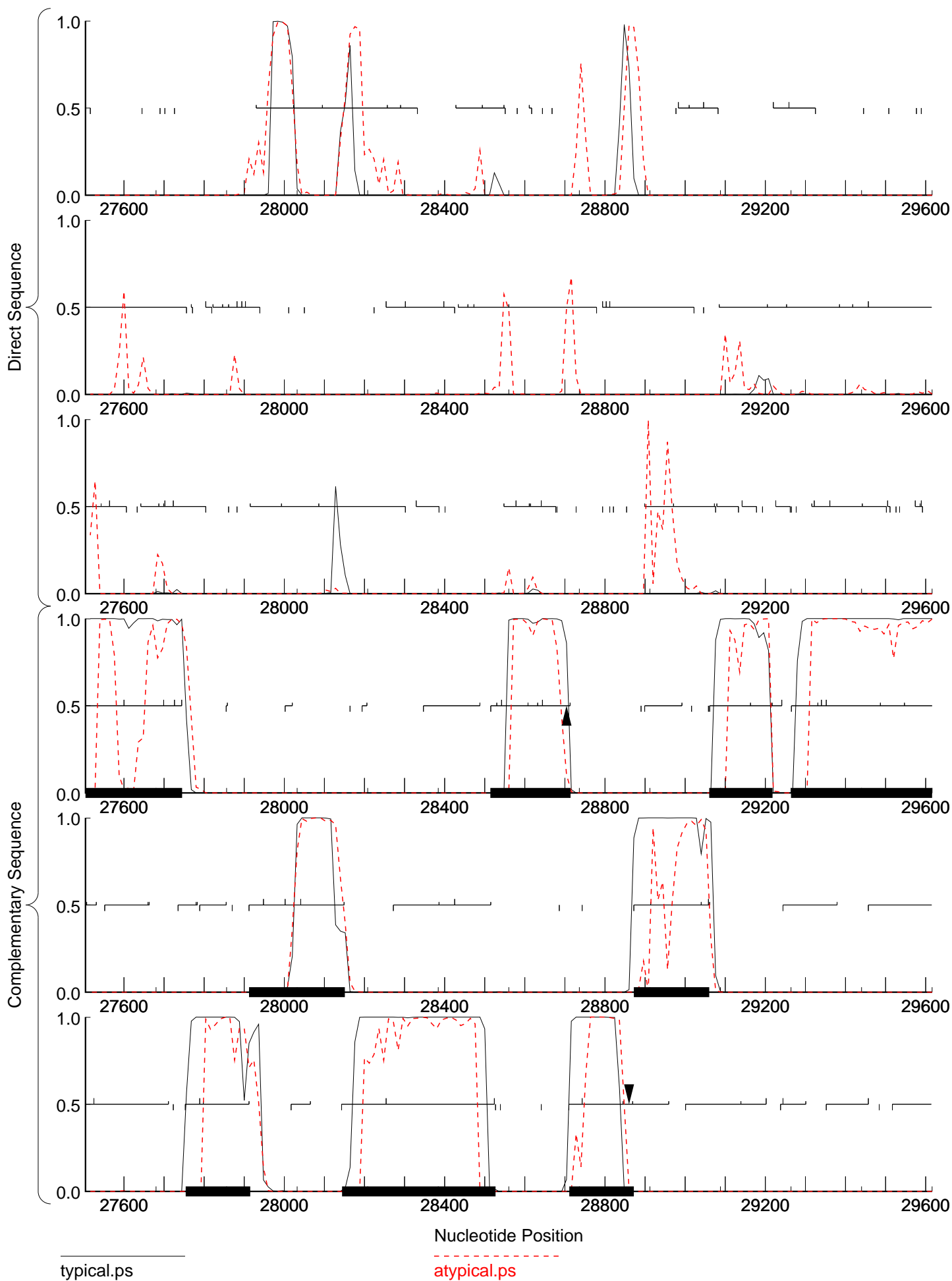


typical.ps

atypical.ps

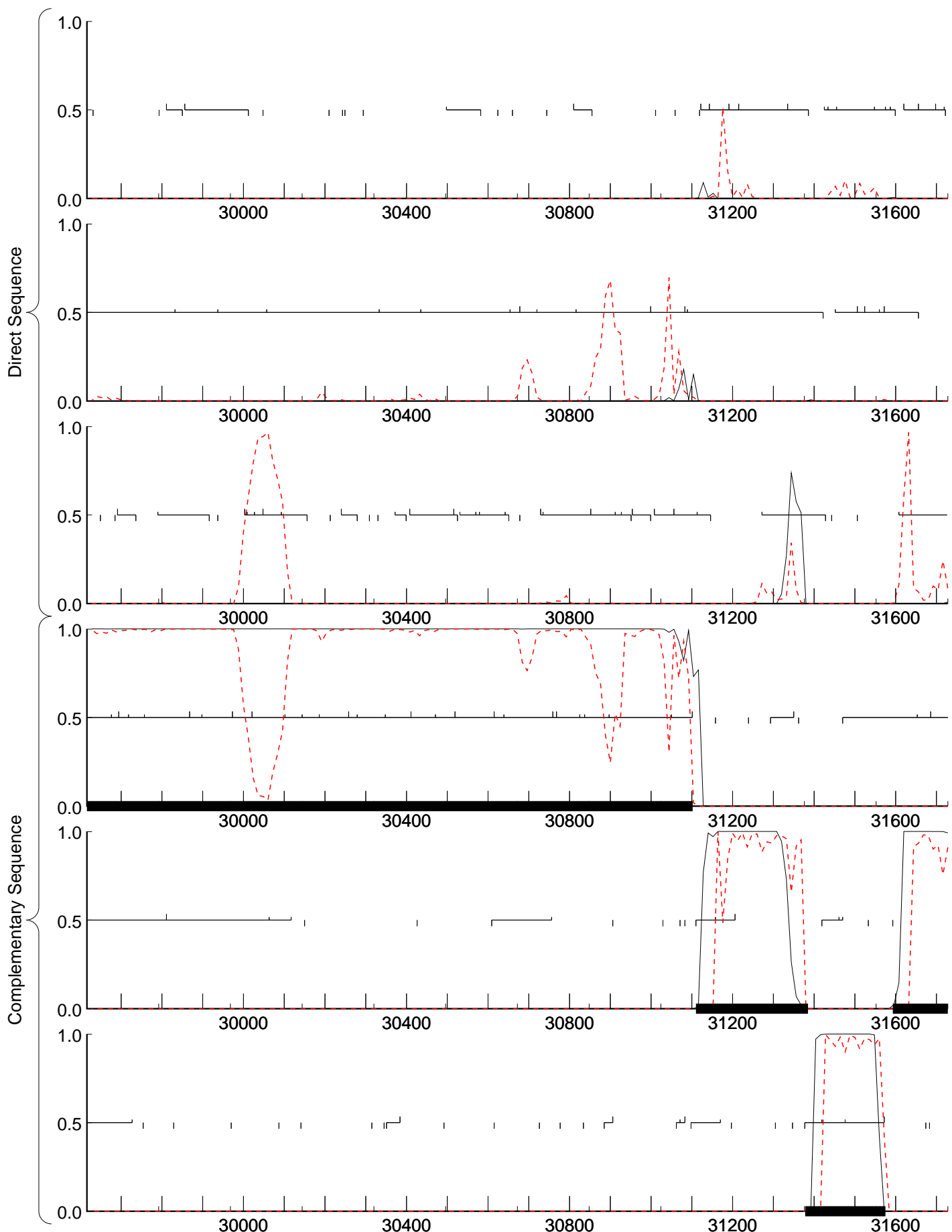






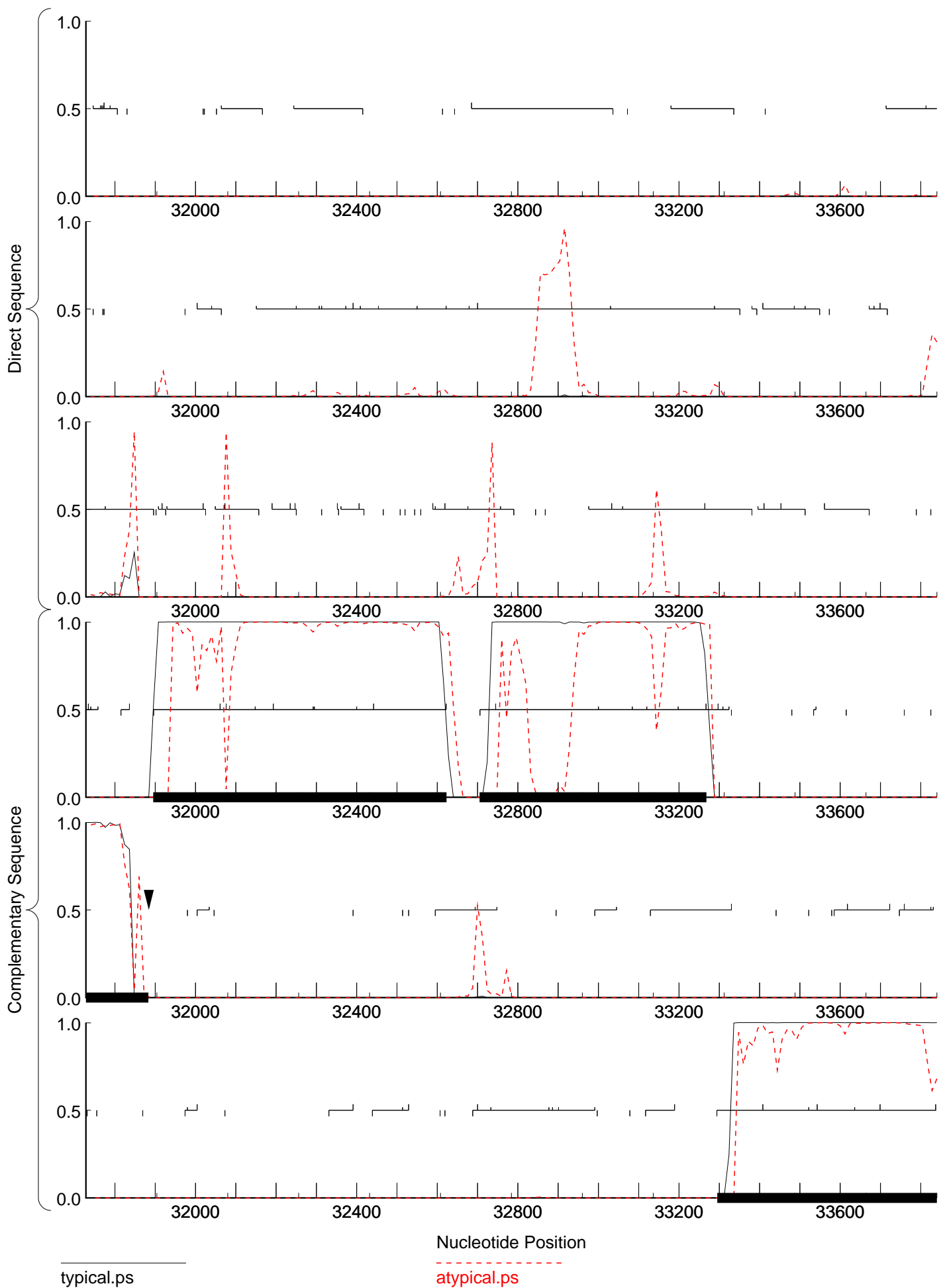
typical.ps

atypical.ps



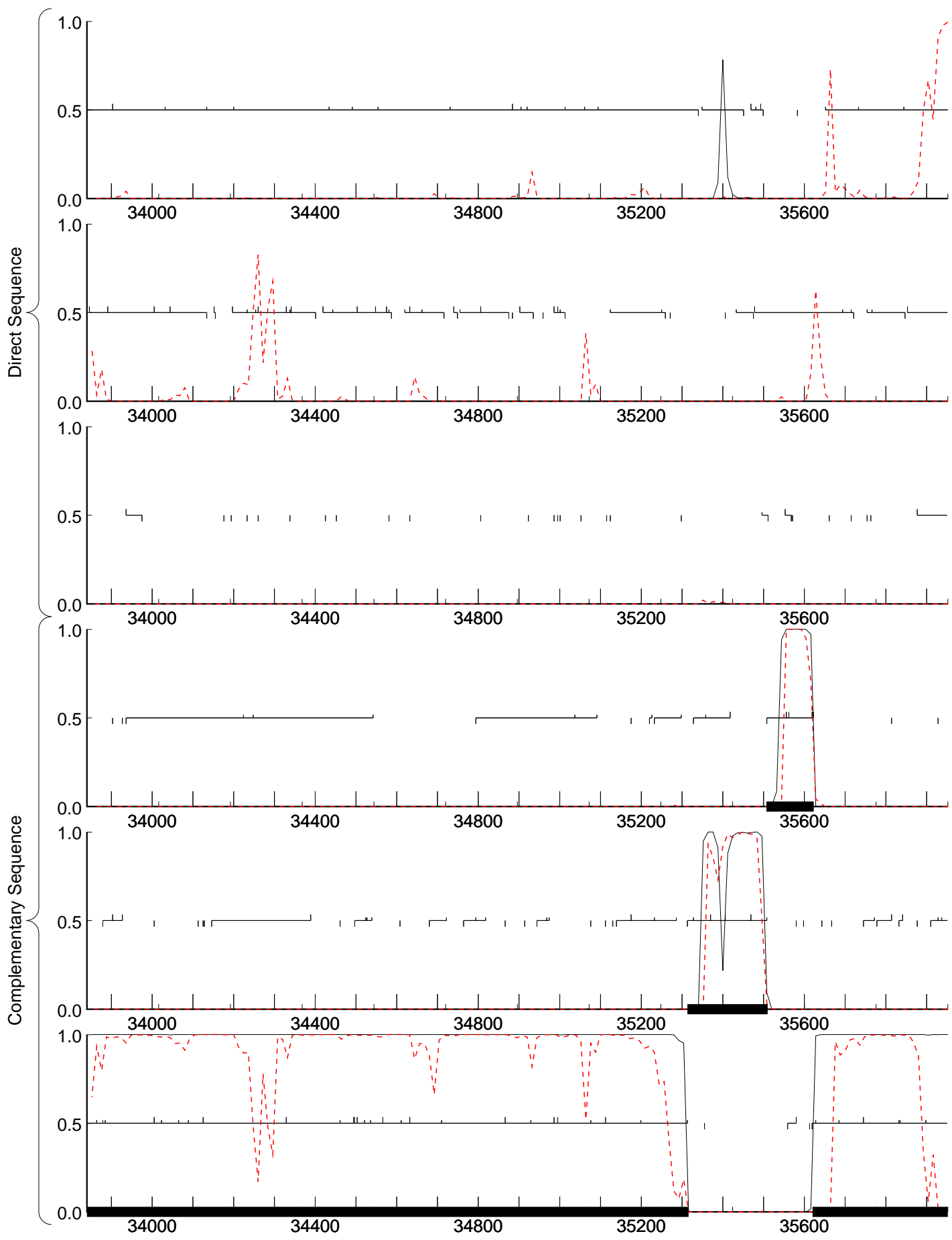
typical.ps

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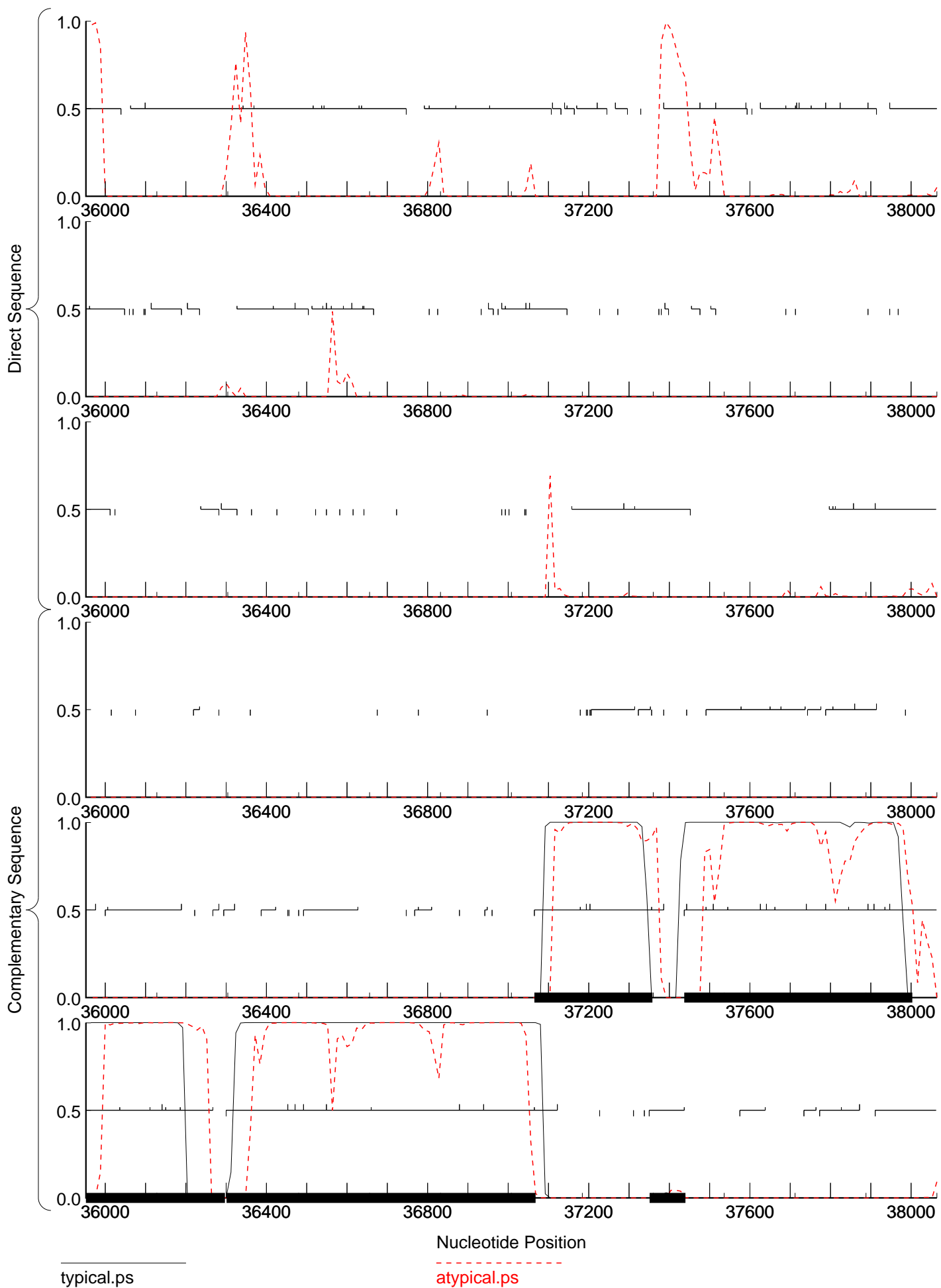
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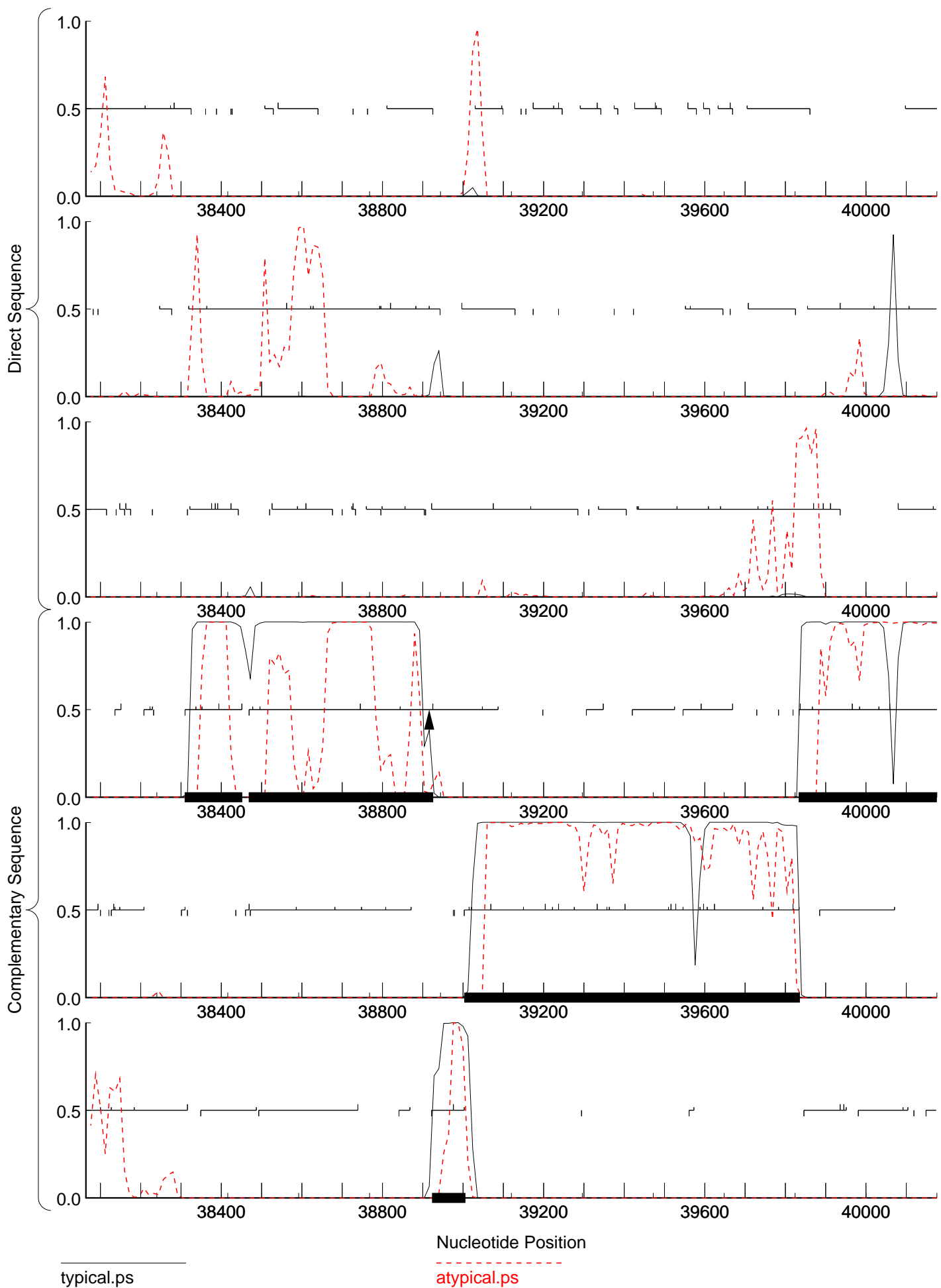
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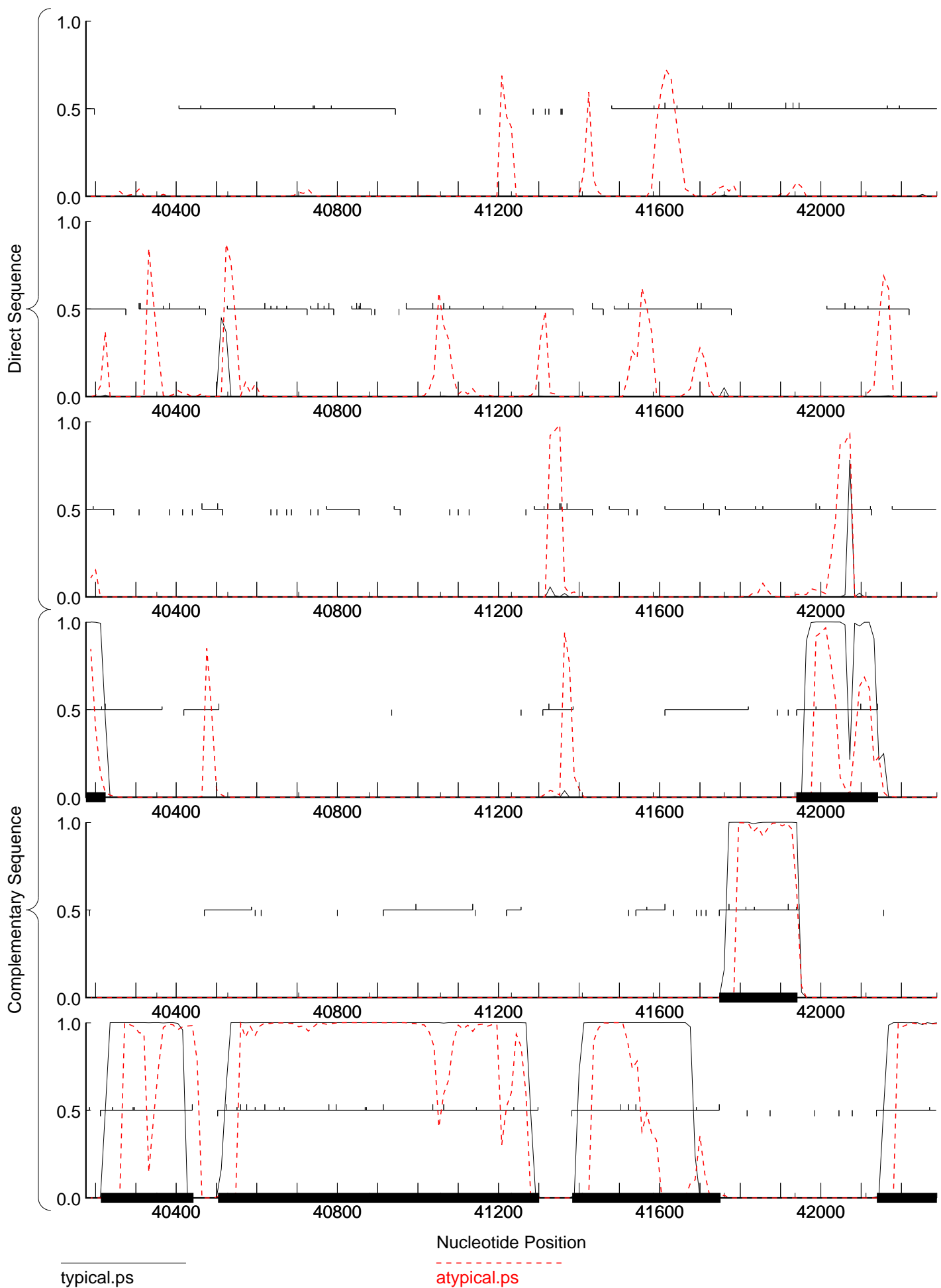


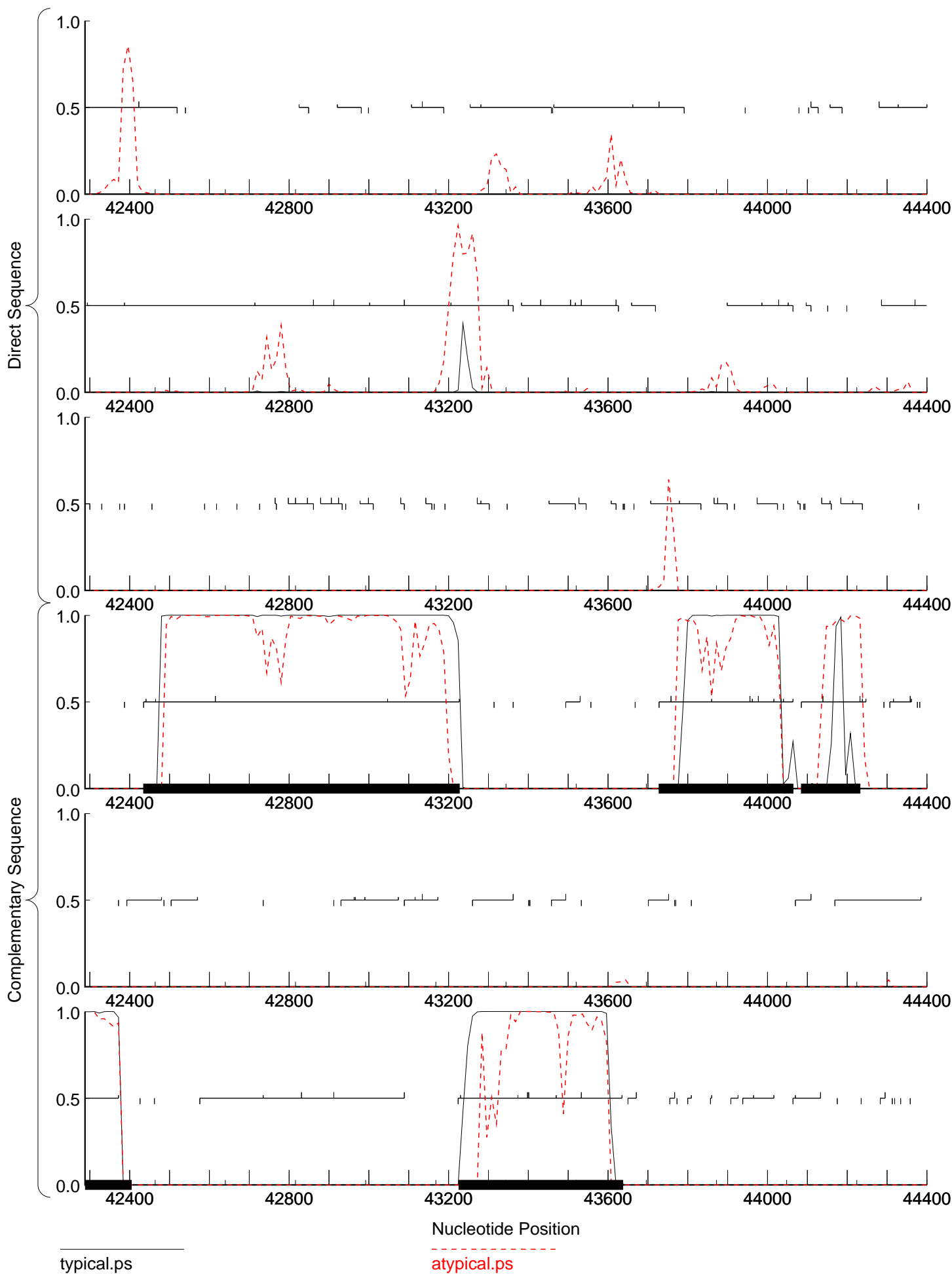
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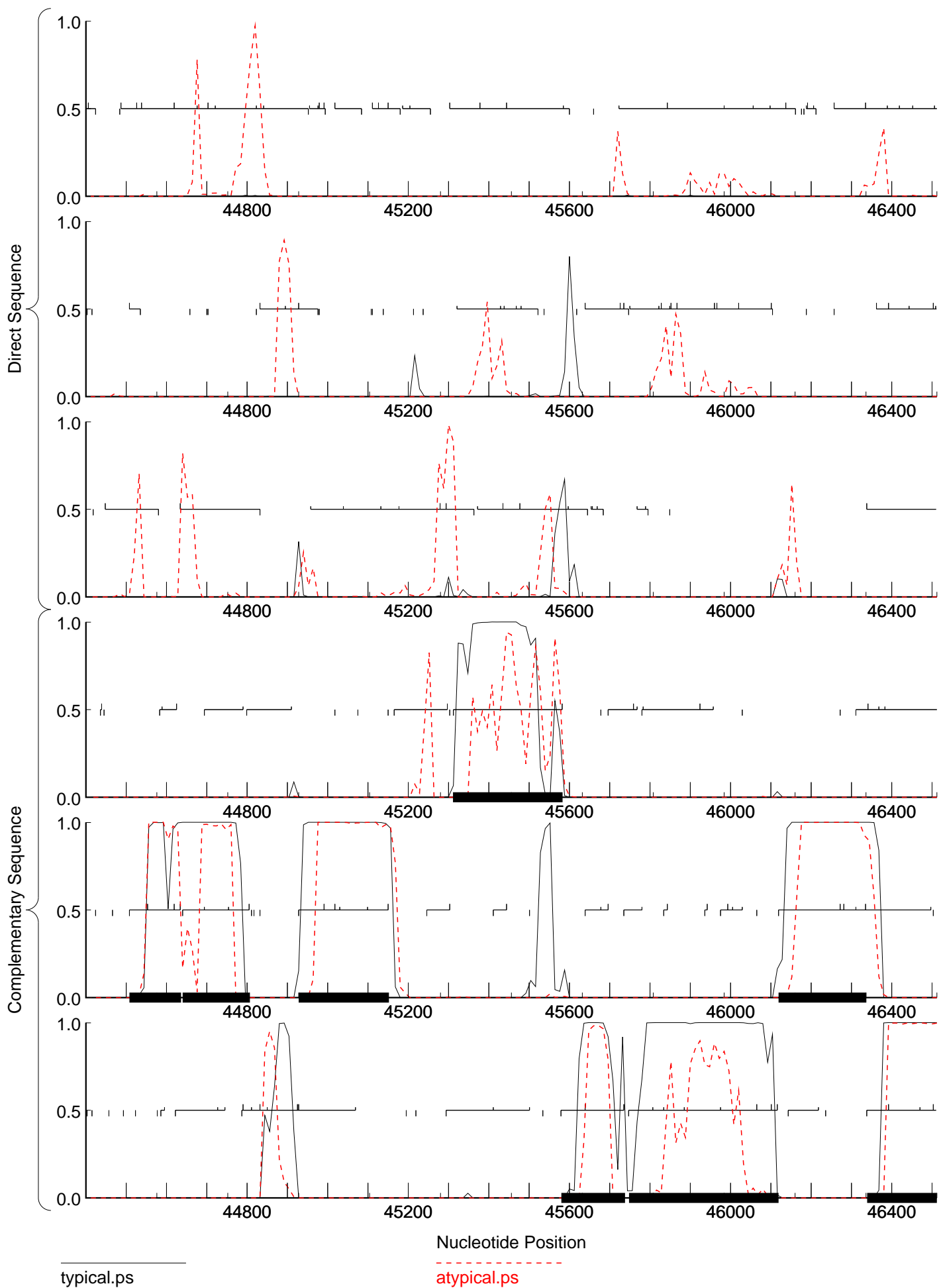
atypical.ps











typical.ps

atypical.ps

