

GeneMark

Version 2.5p (09.08.06)

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PROGRAM INFORMATION

Sequence : Mycobacterium phage Wizard007 complete sequence, 51029 bp including 10-base 3' overhang (CG
Analysis Date : 7/24/17 at 22:31:17
Pages : 26
Sequence Length : 51029 bp
GC Content : 63.85%

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

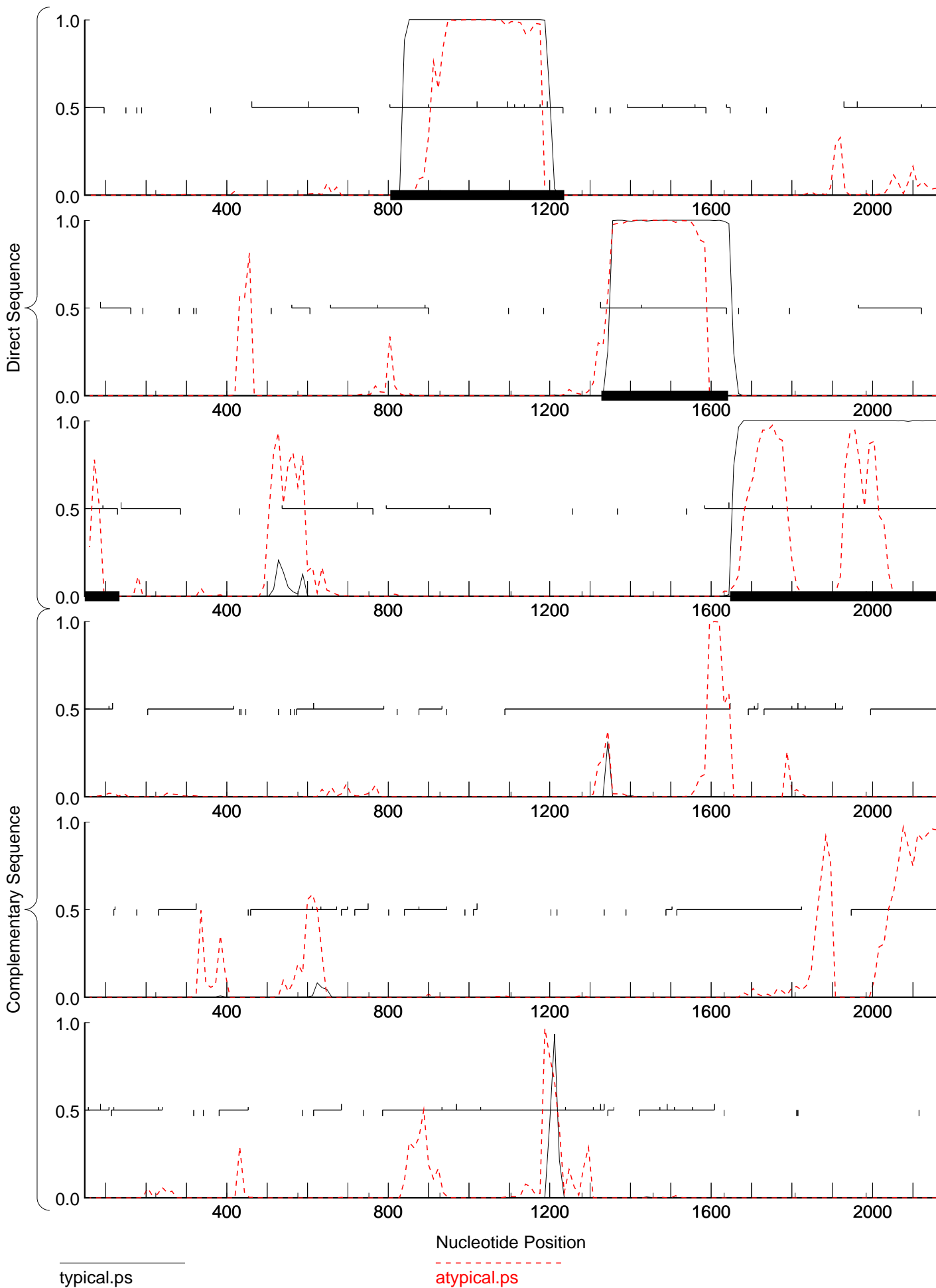
Training set derived by GeneMarkS, 4.30 October 2014
Mon Jul 24 22:31:17 2017

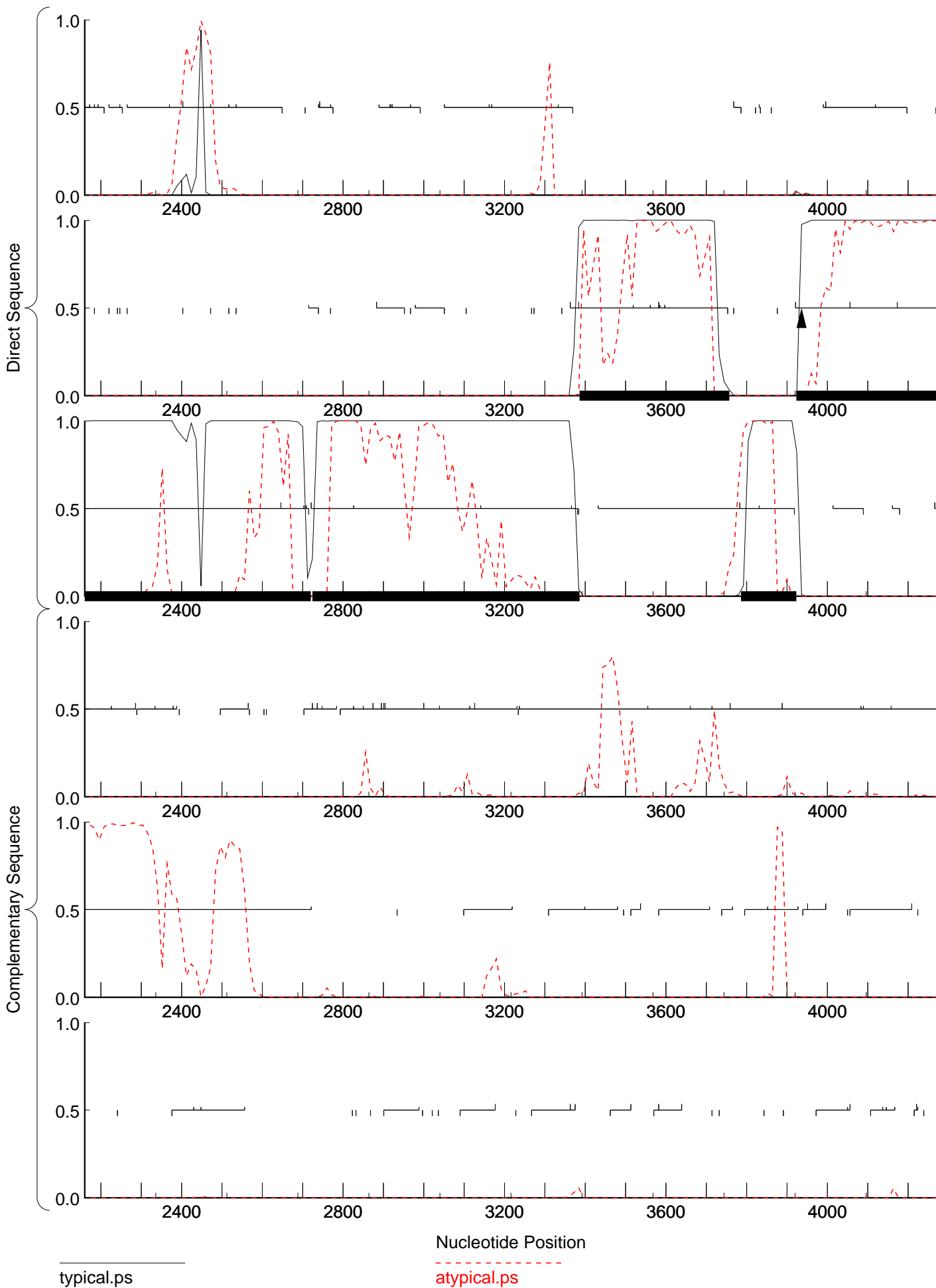
MATRIX INFORMATION

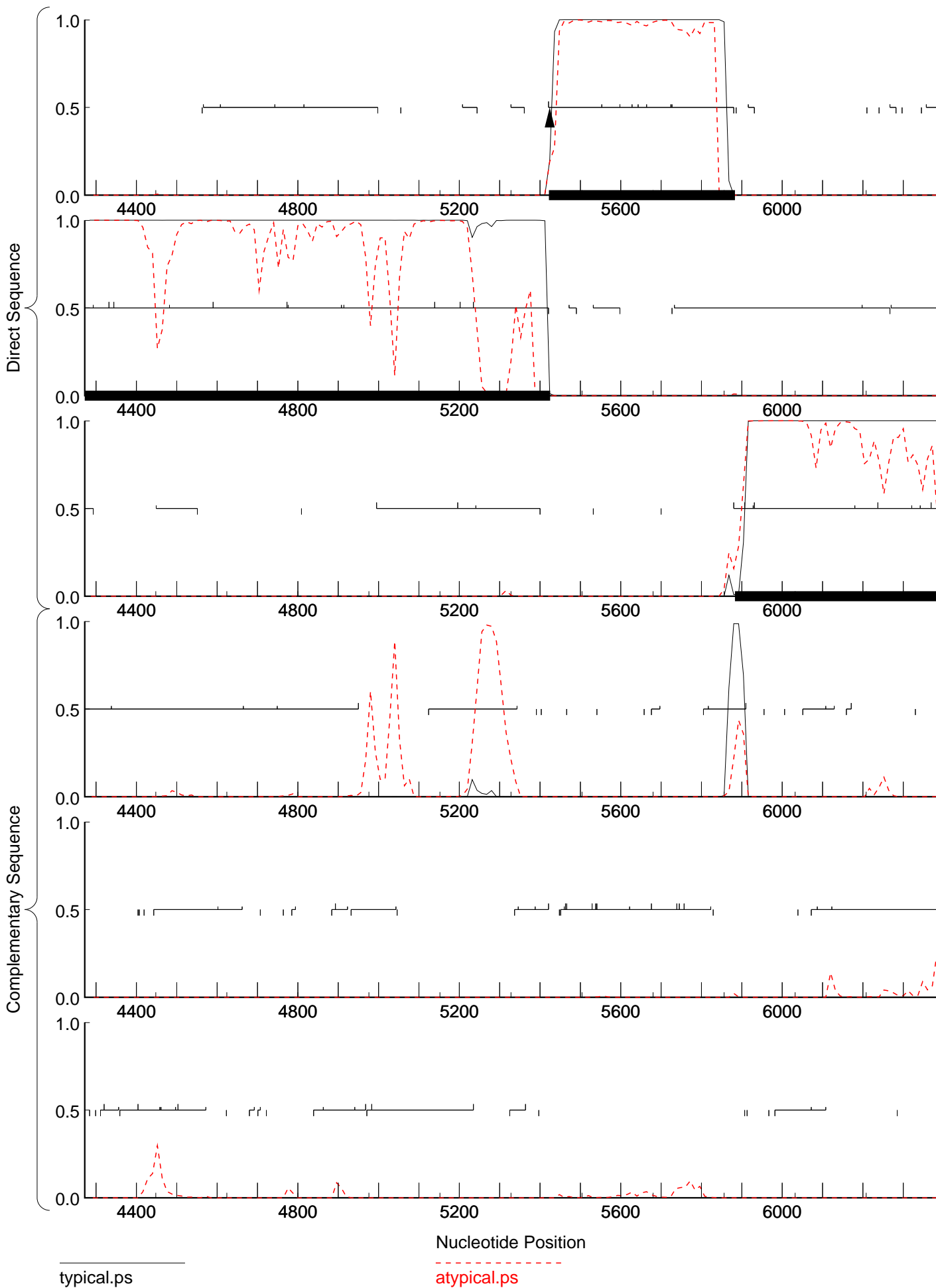
Matrix : -
Author : -
Order : 4

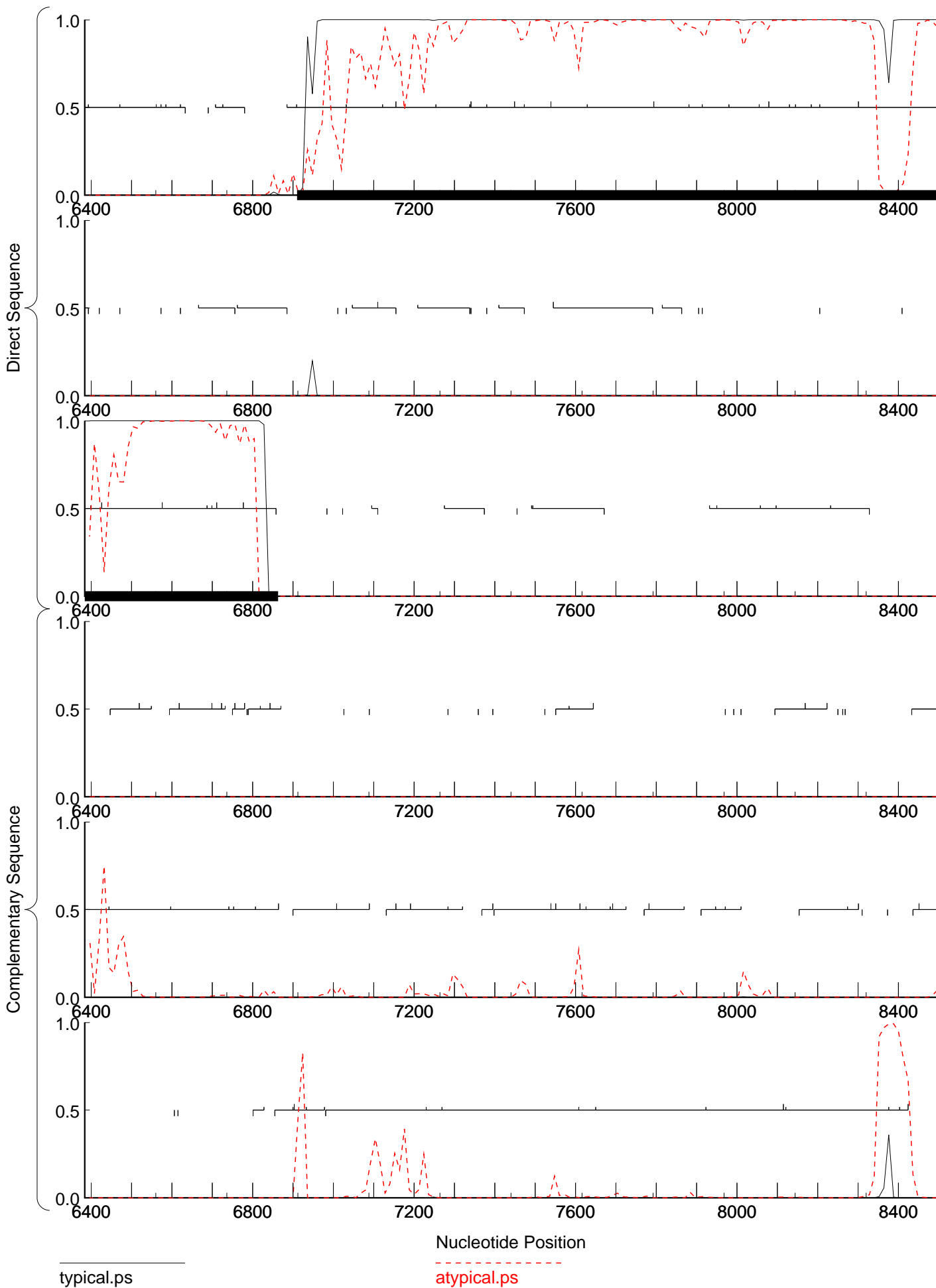
Send questions / comments to:
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Georgia Institute of Technology
School of Biology
Atlanta, GA 30332-0230

For help or information, send electronic mail to: custserv@genepro.com, or mark@amber.biology.gatech.edu



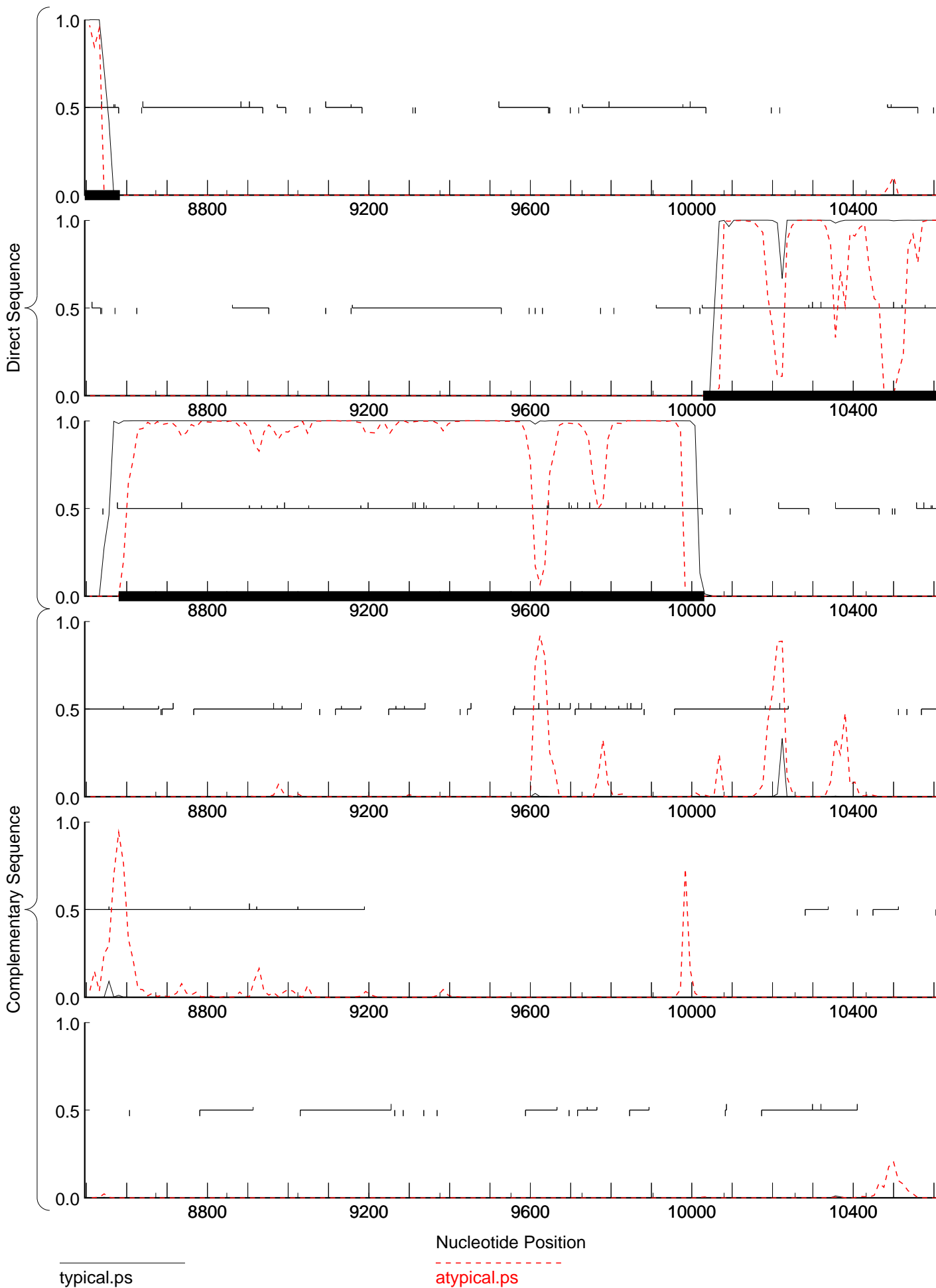


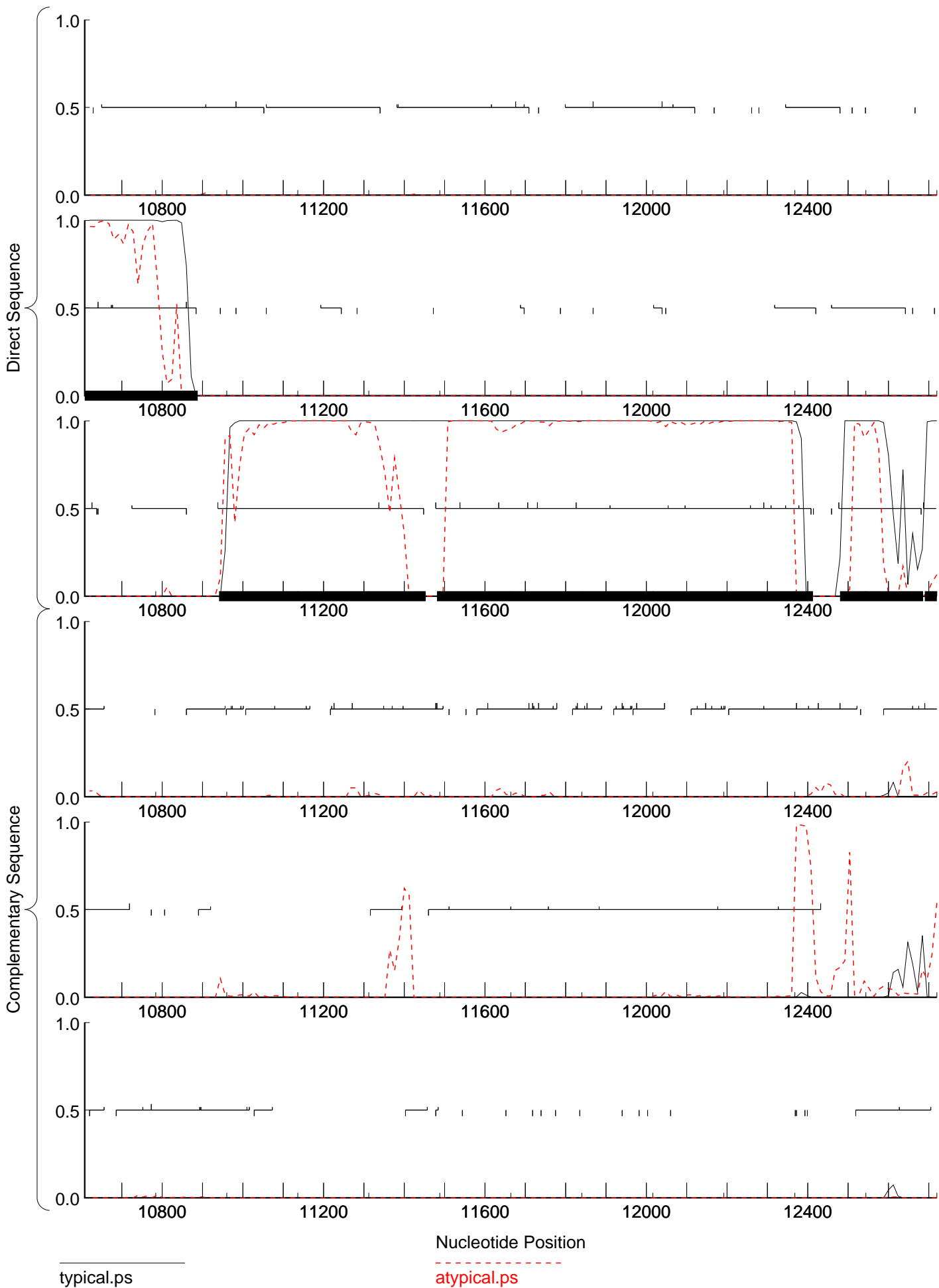


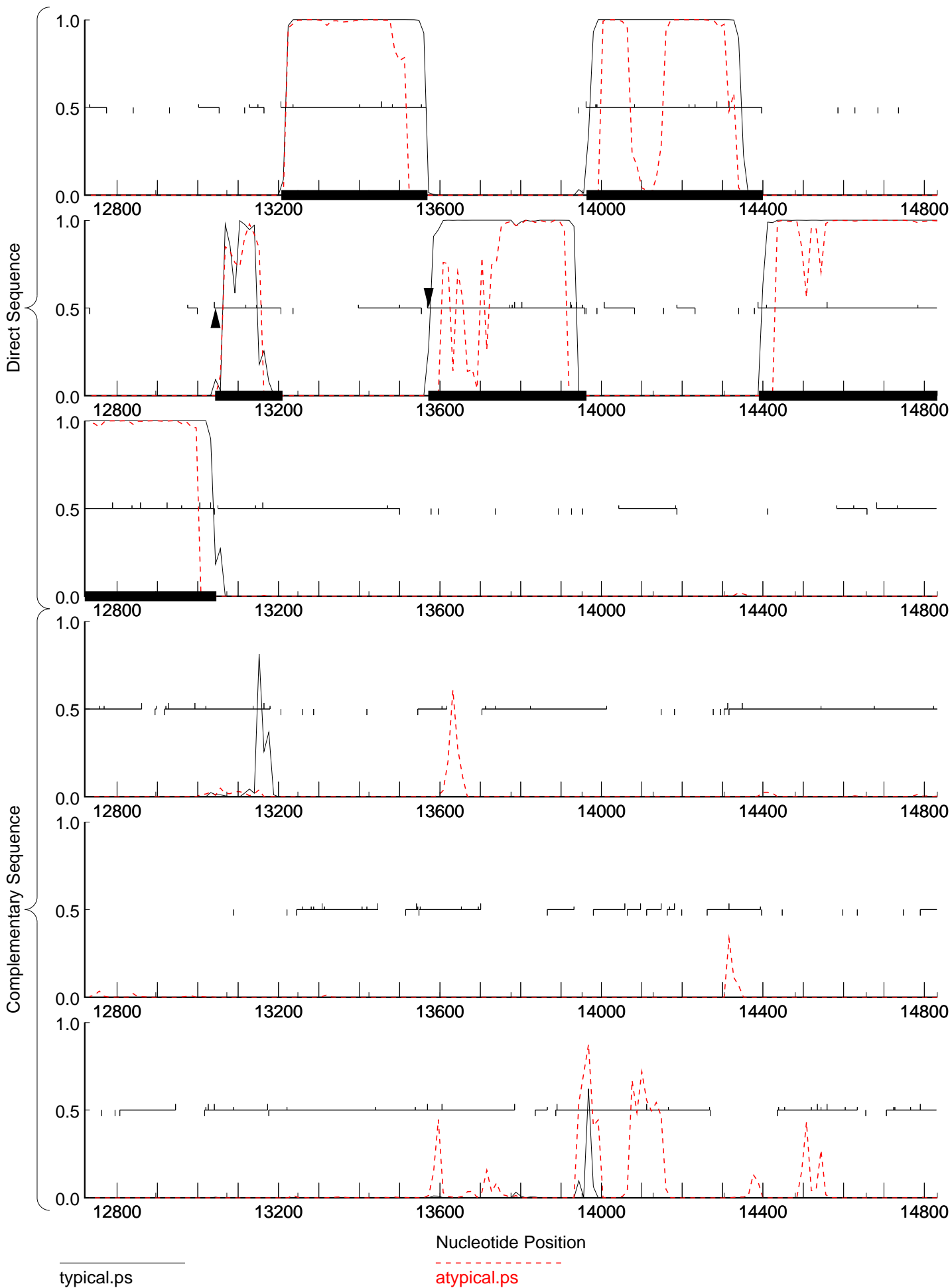


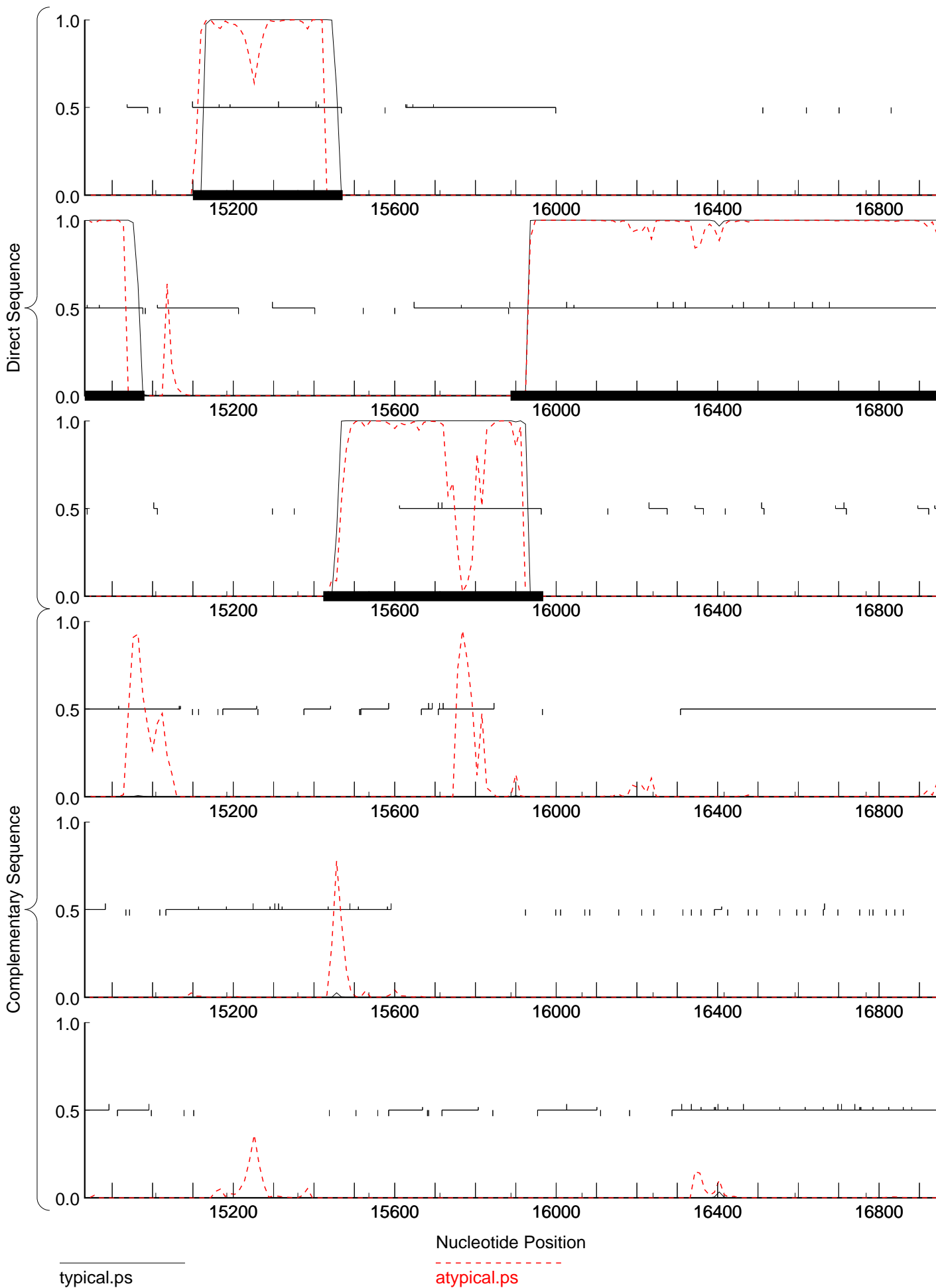
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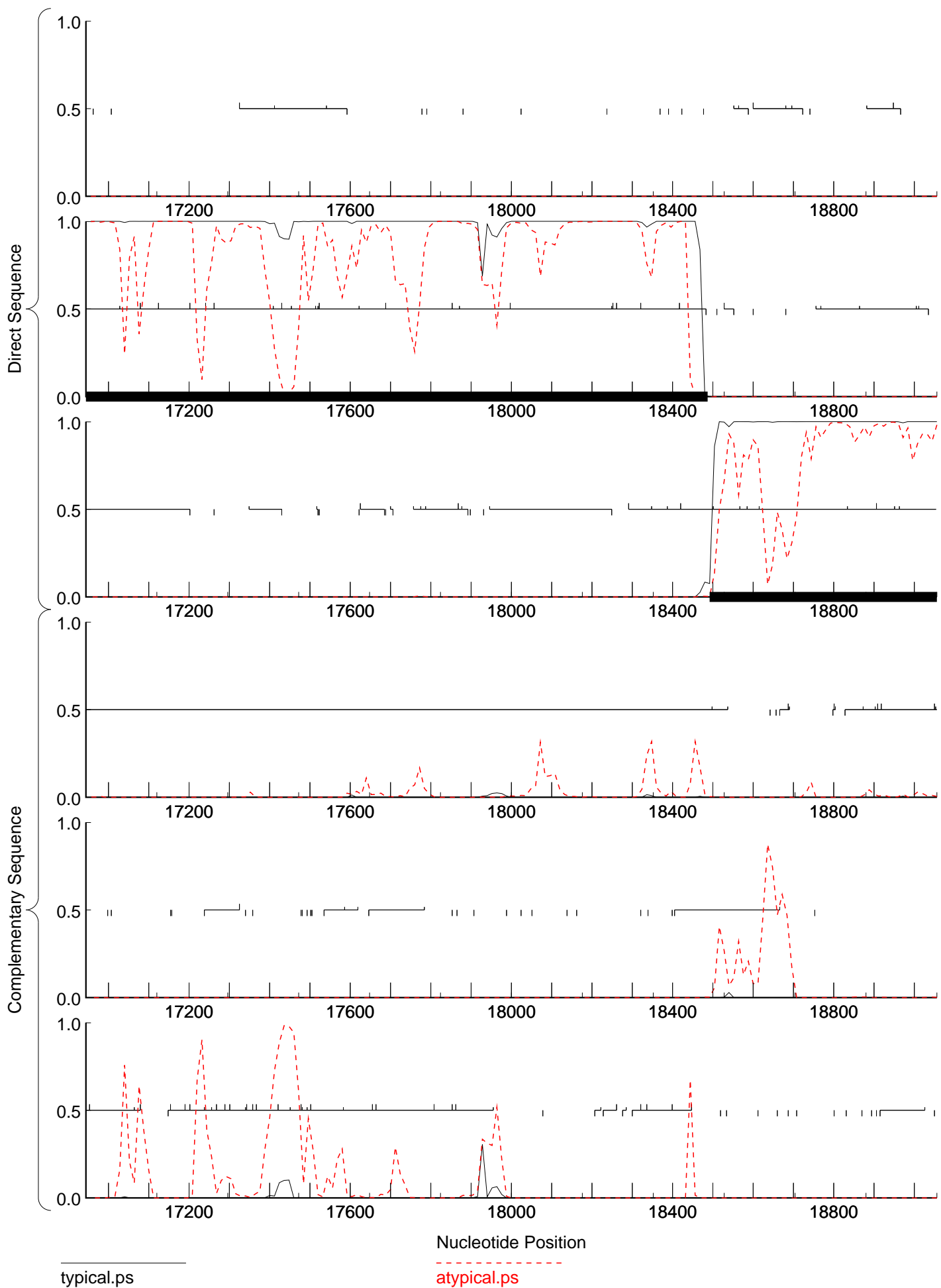
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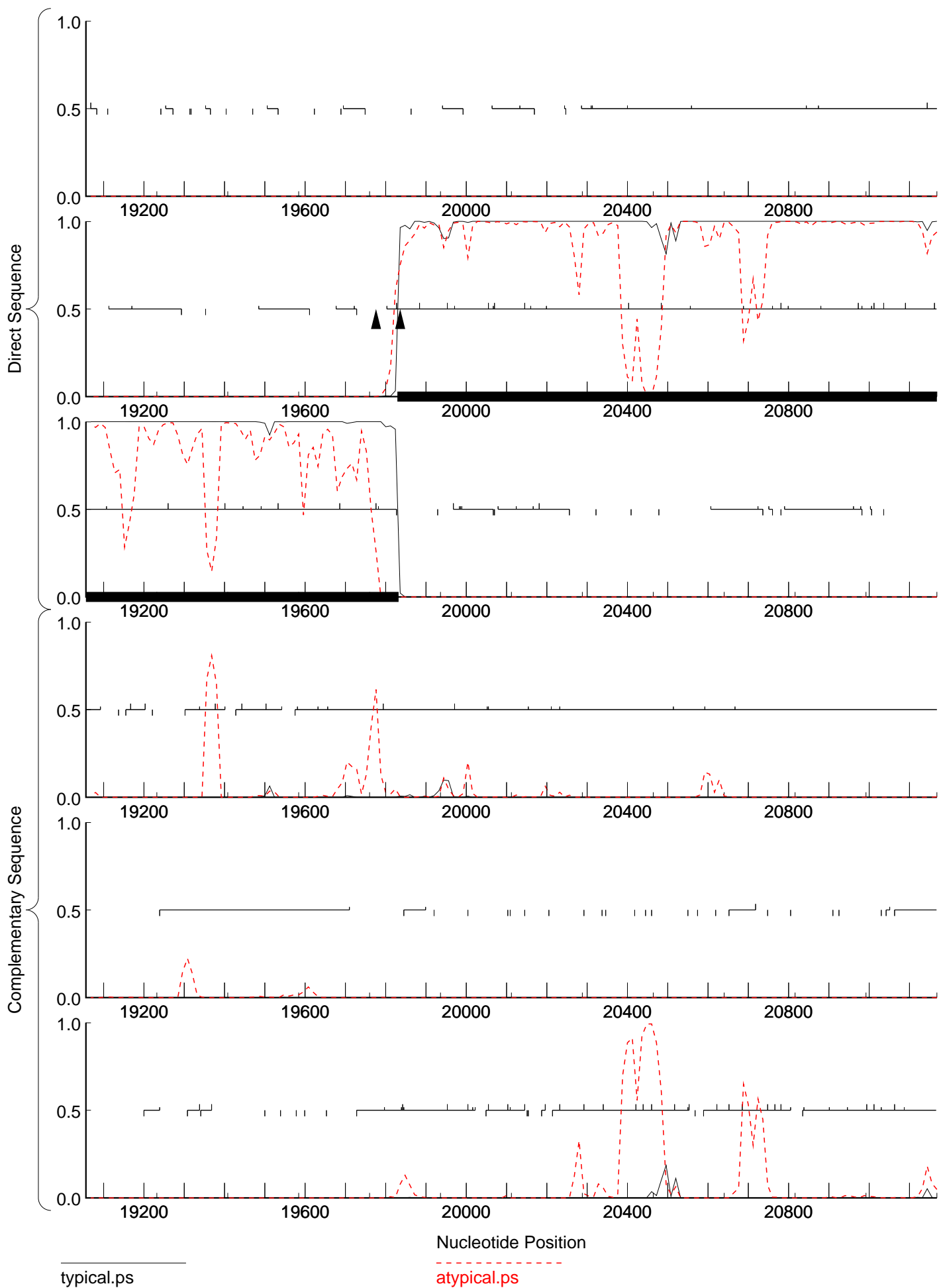






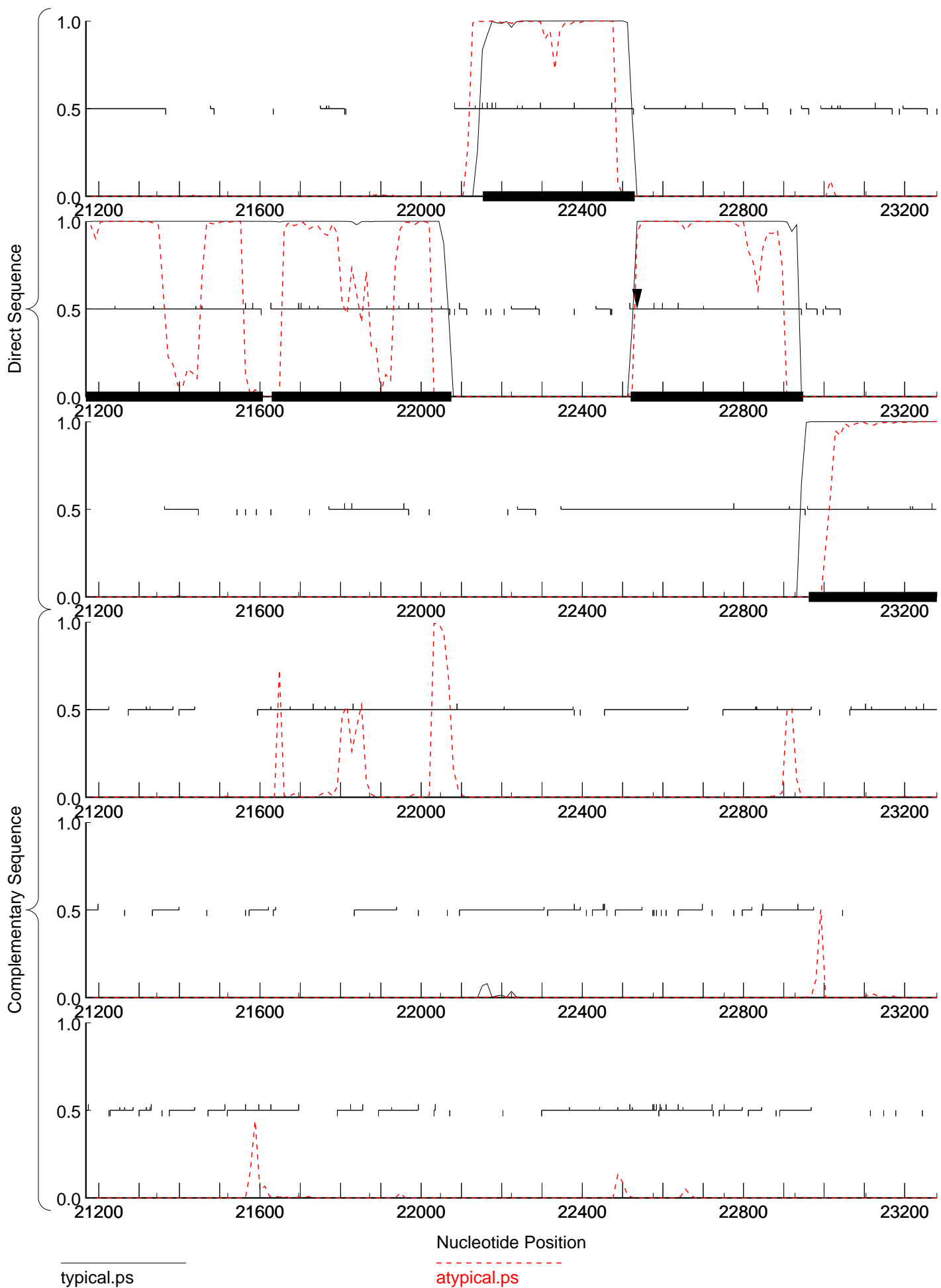
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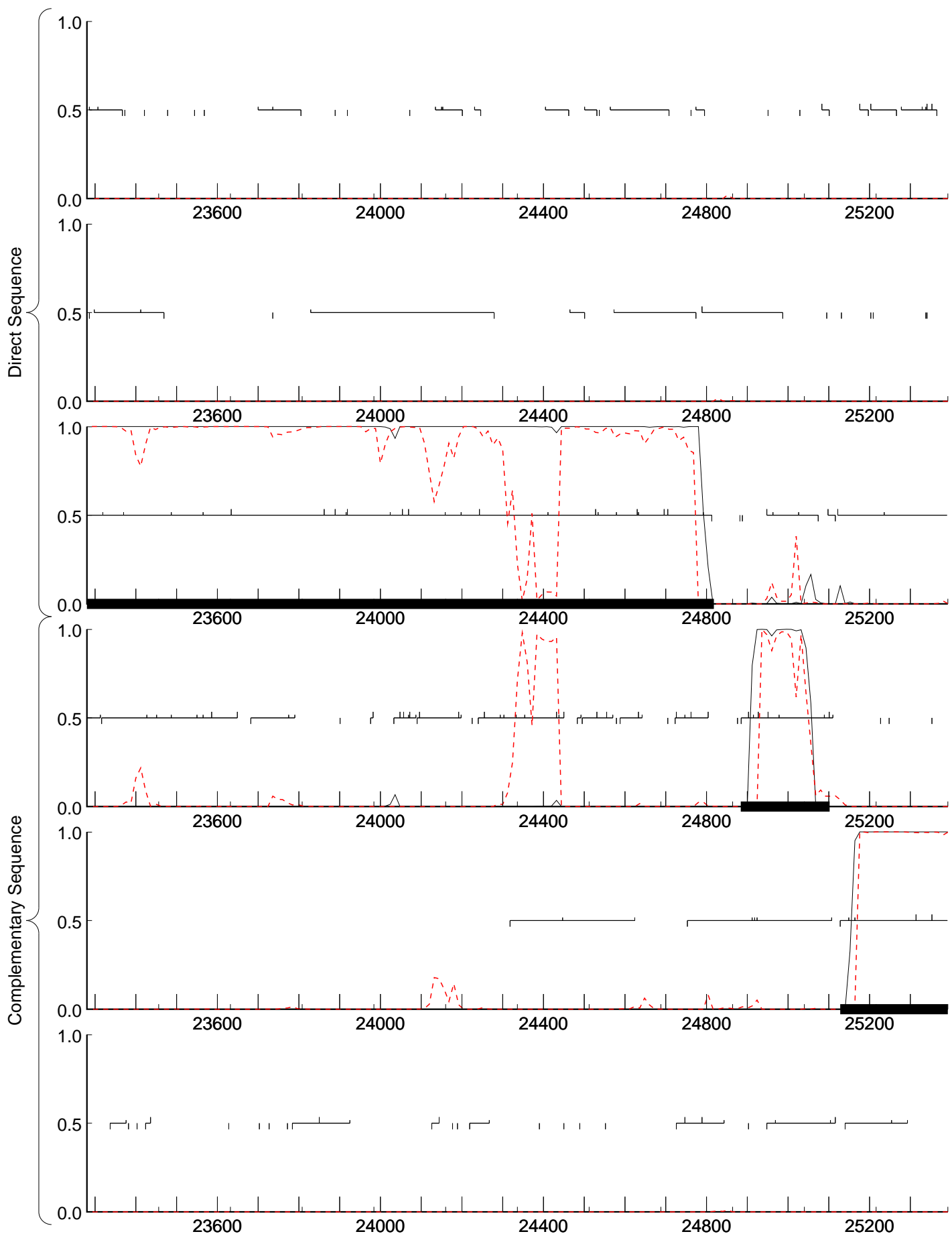
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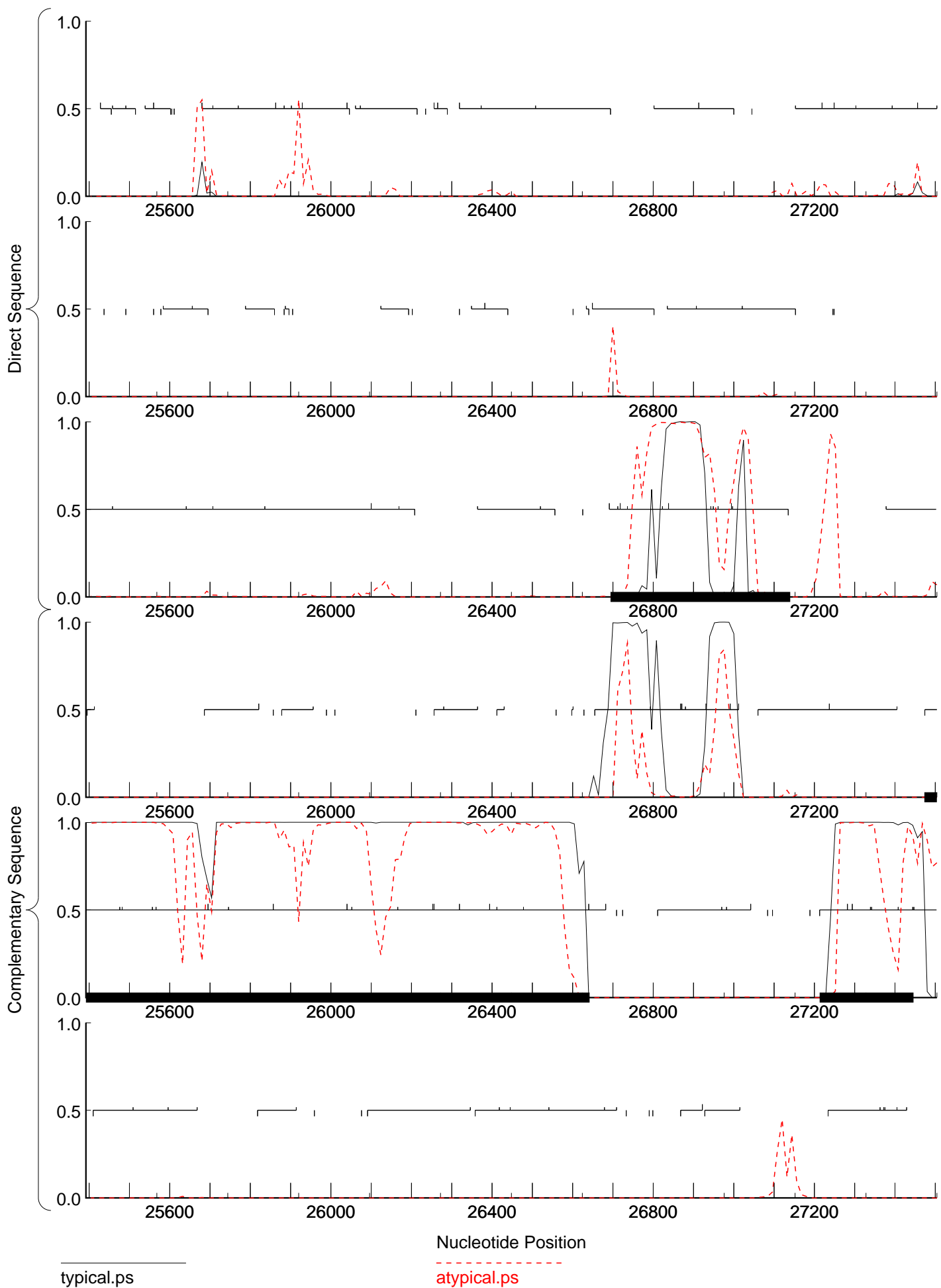


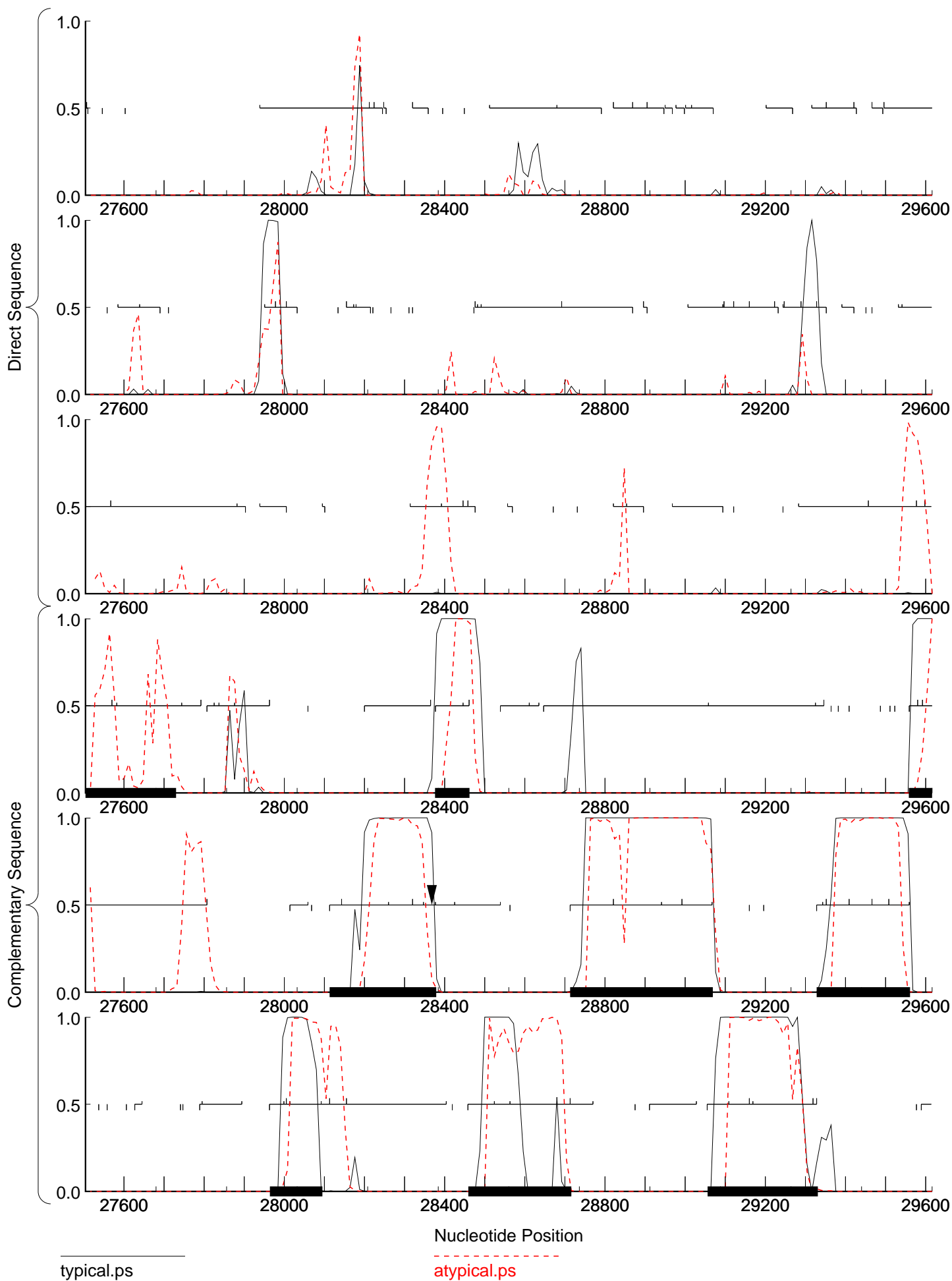
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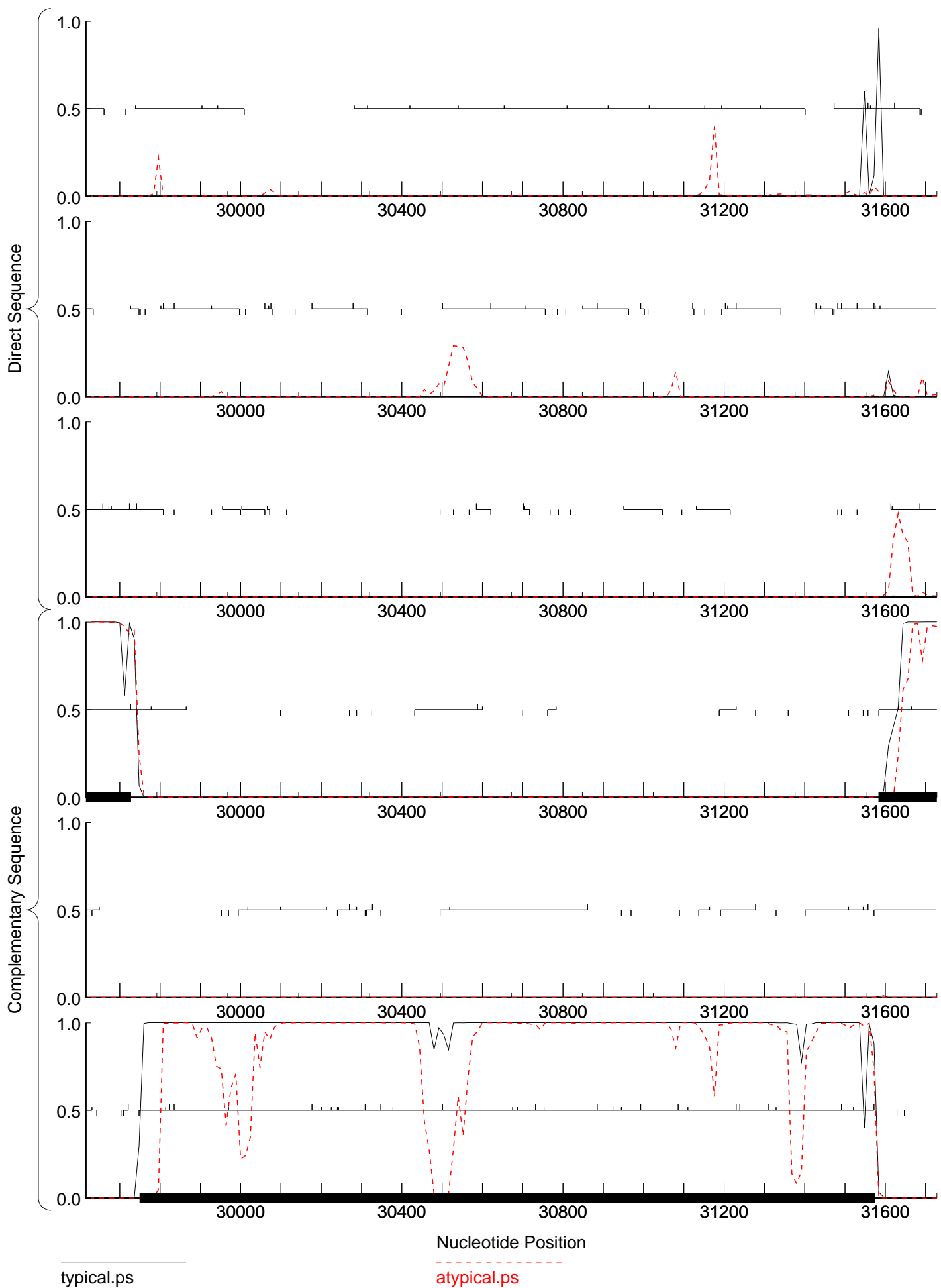


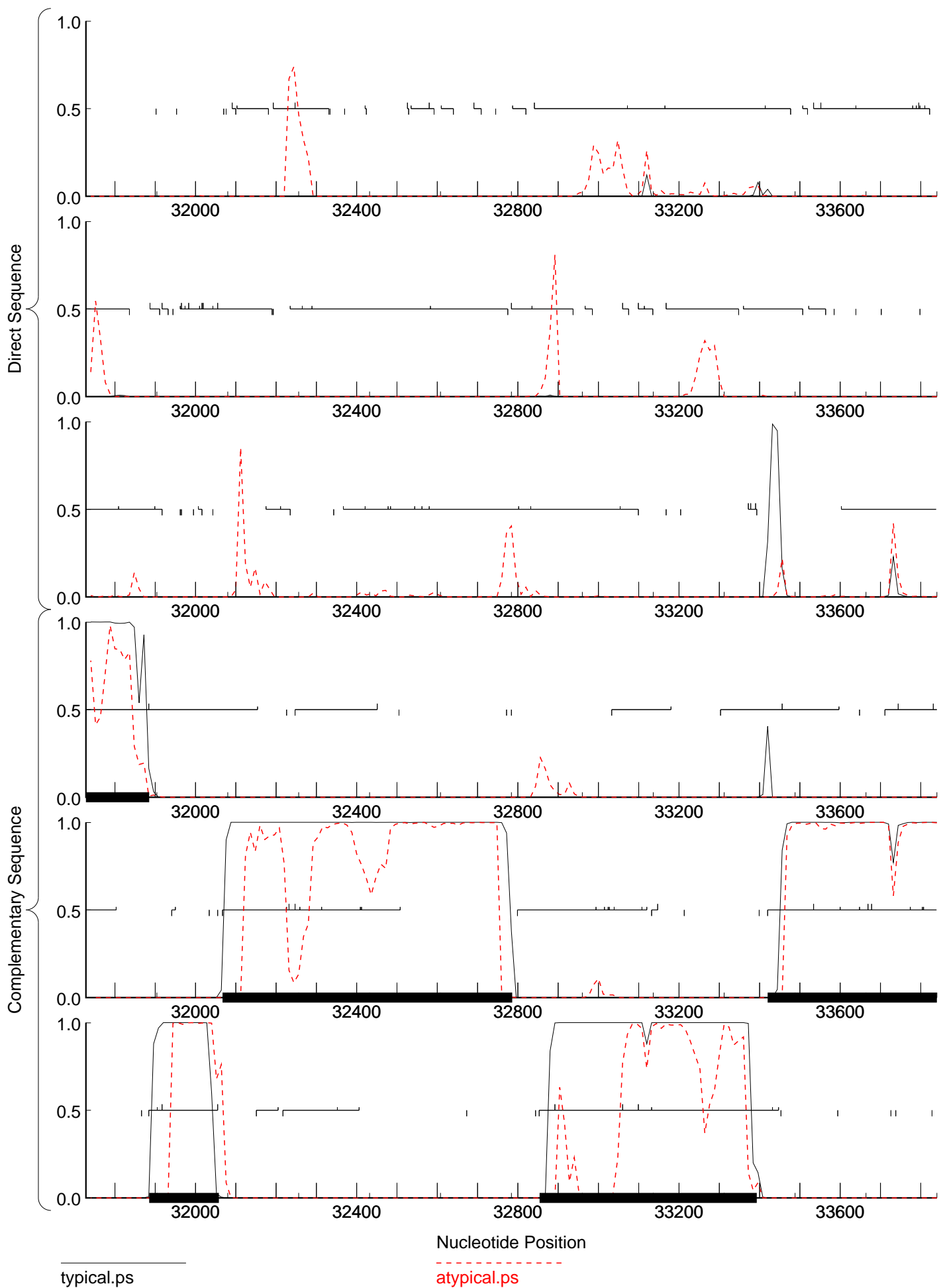




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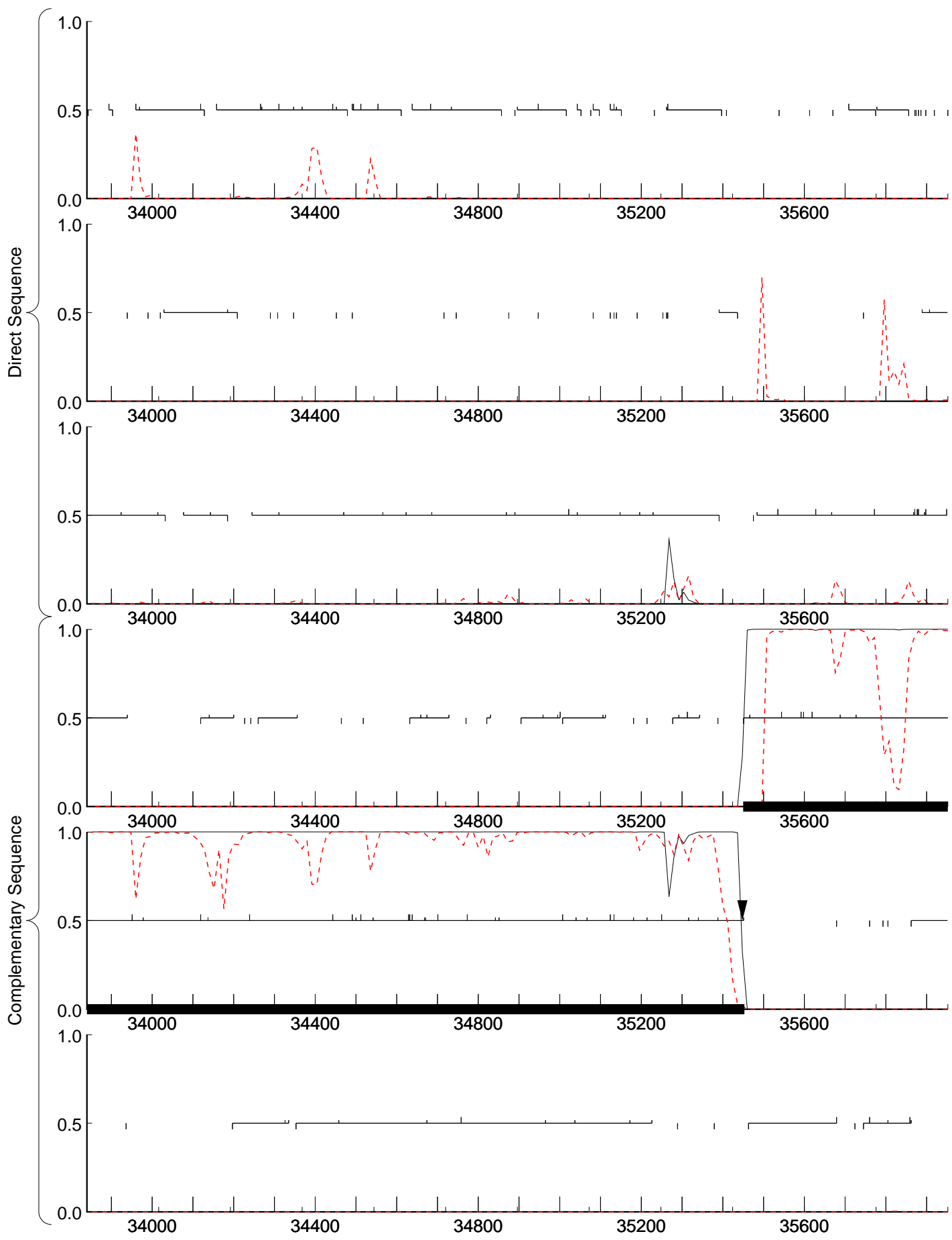
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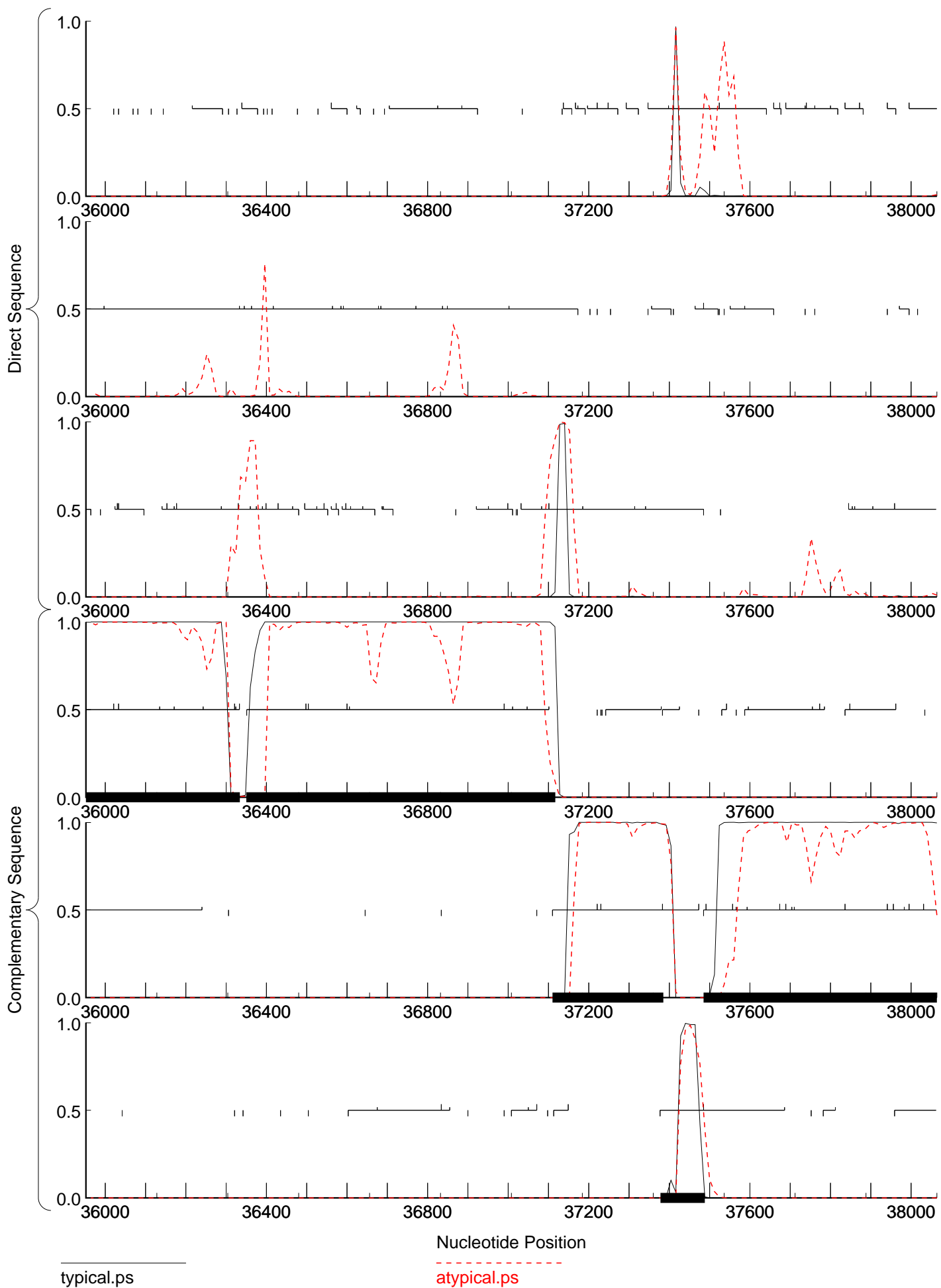
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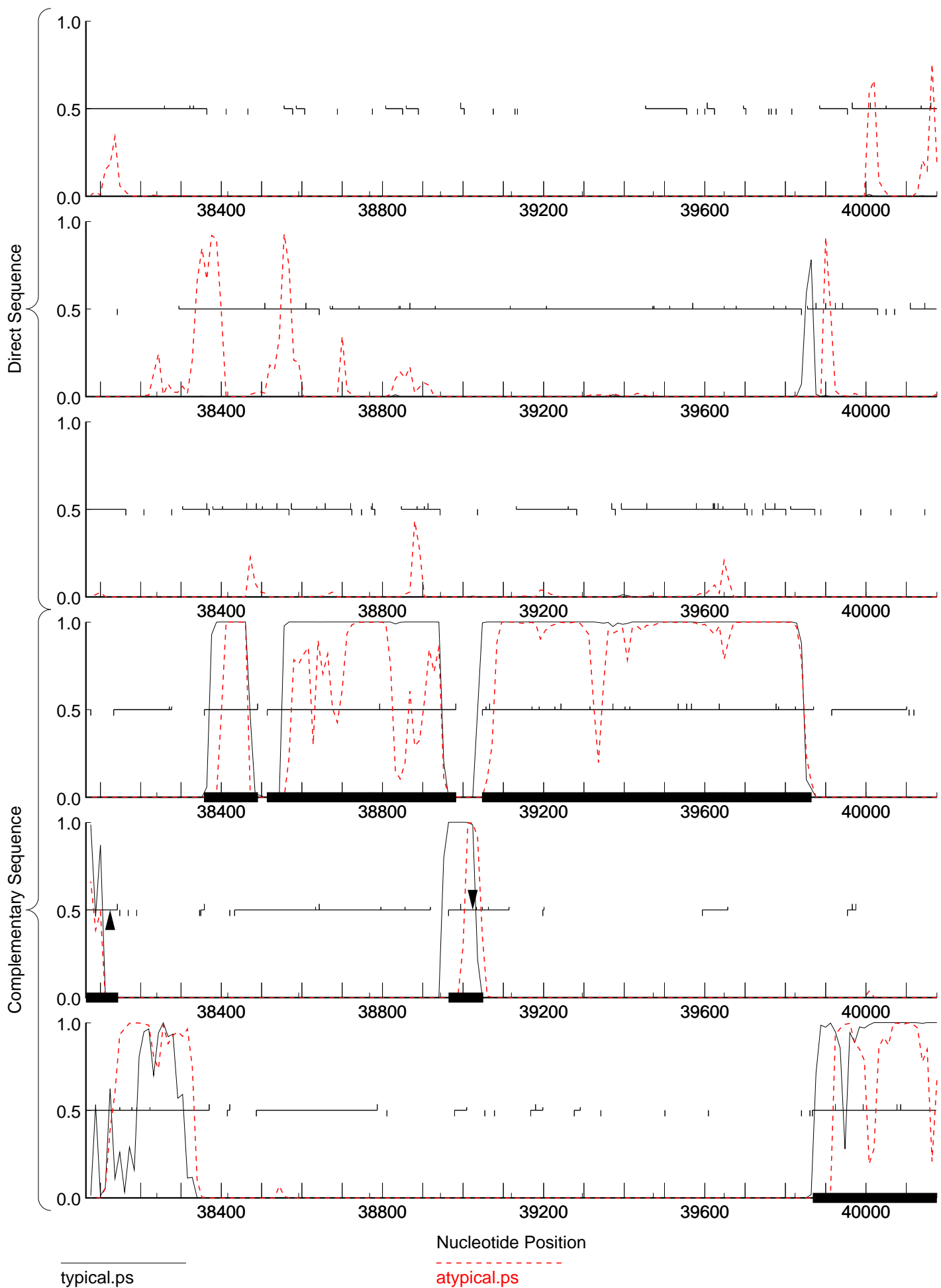
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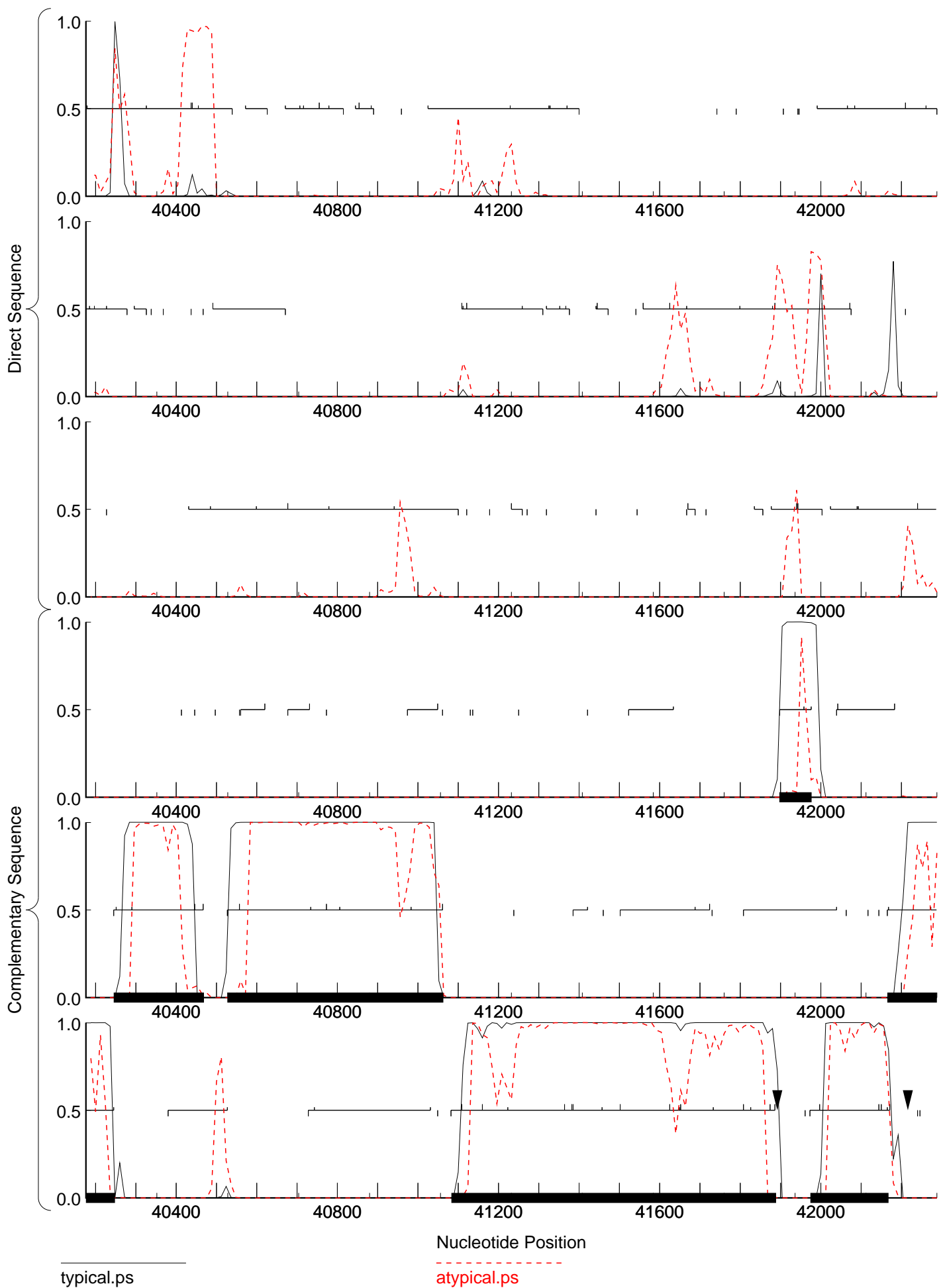


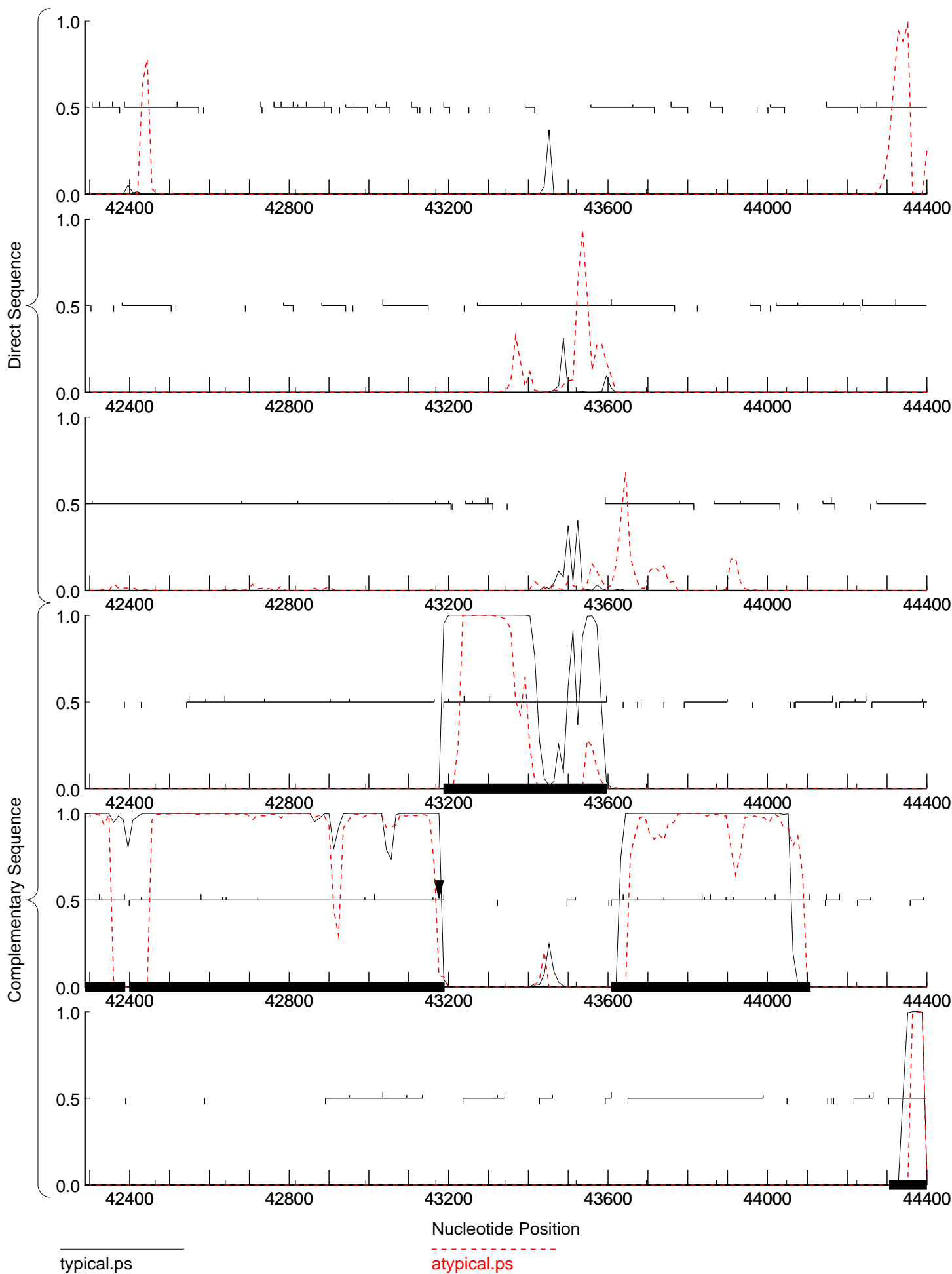
typical.ps

atypical.ps



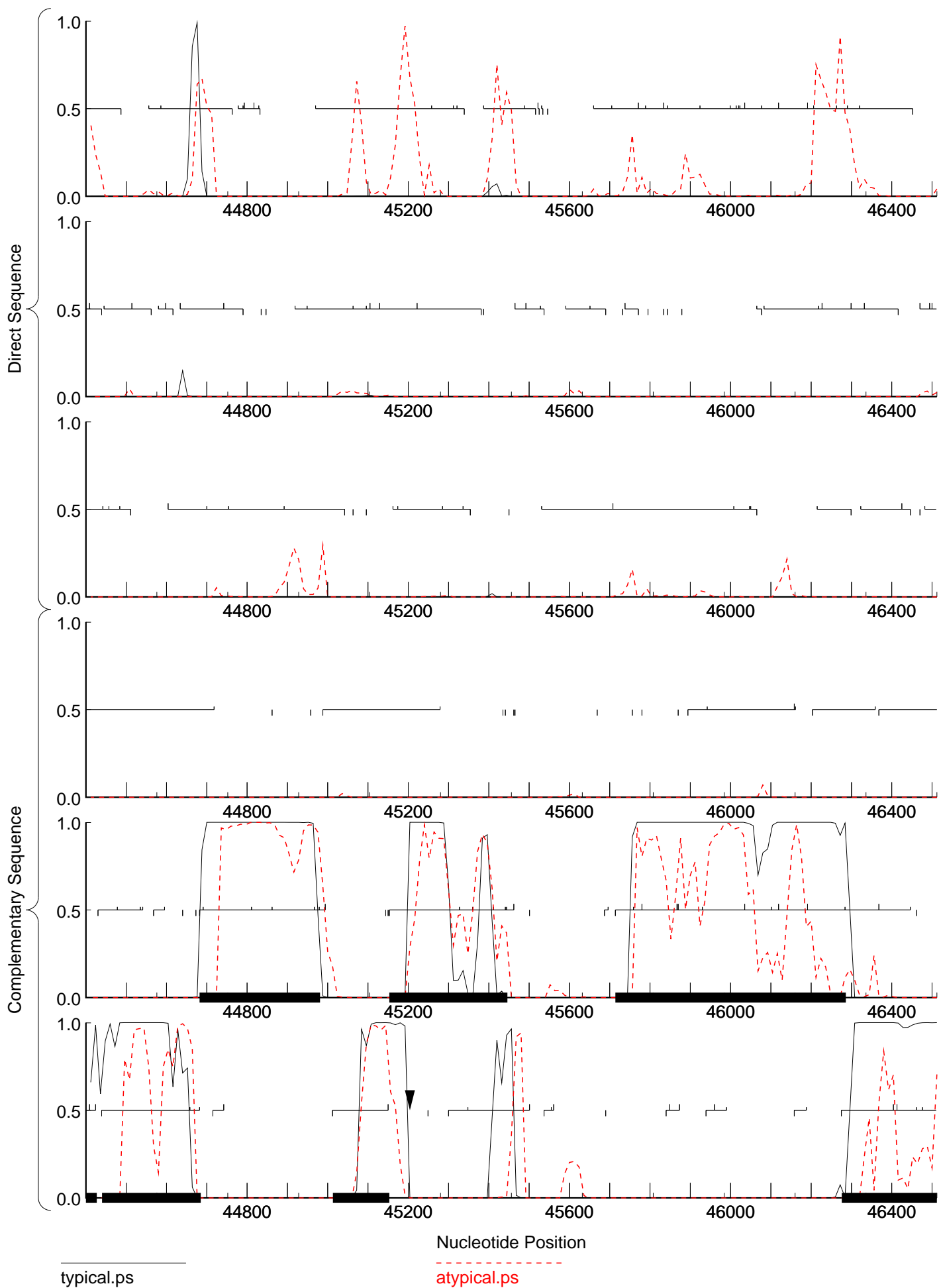


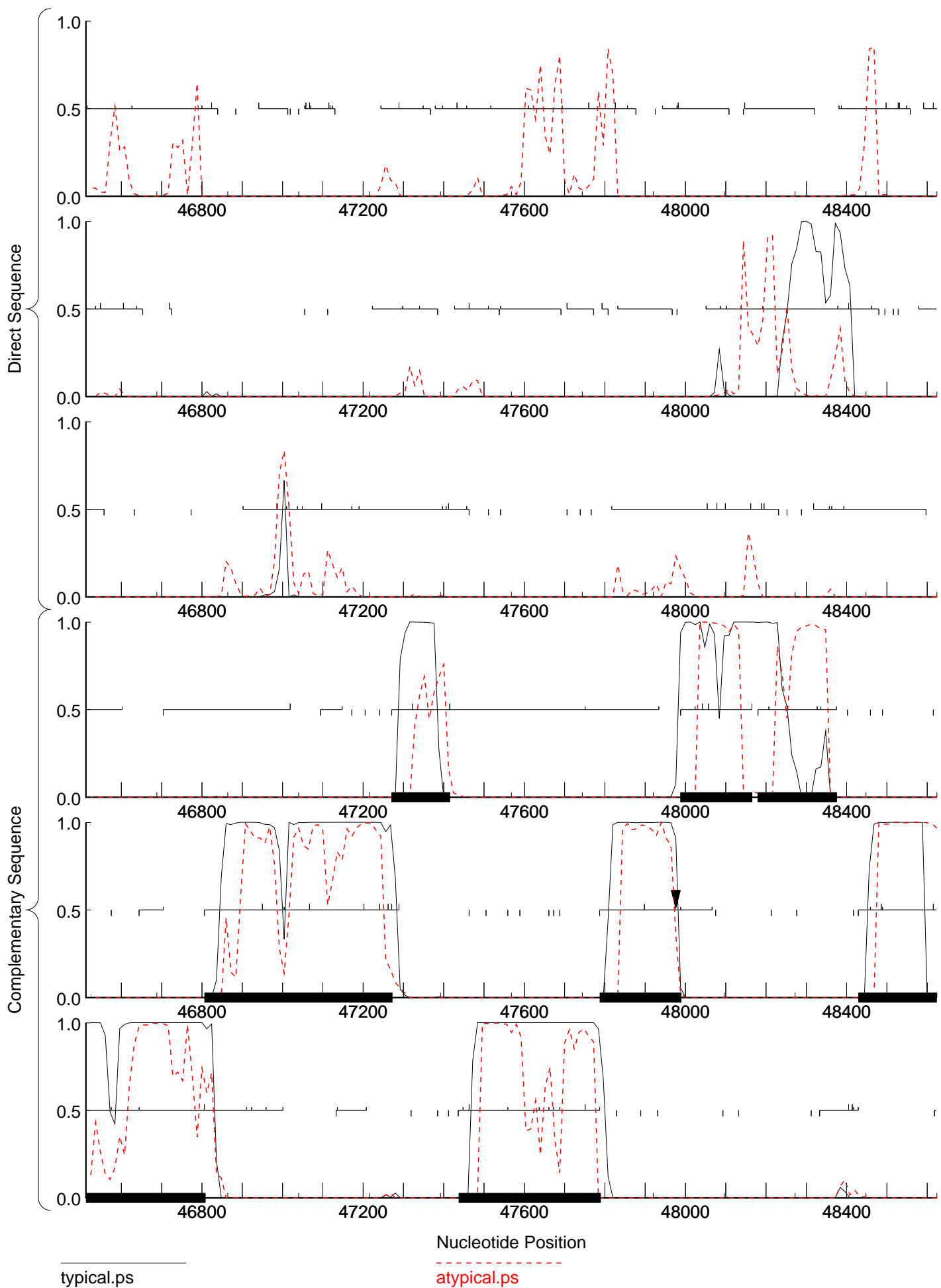




typical.ps

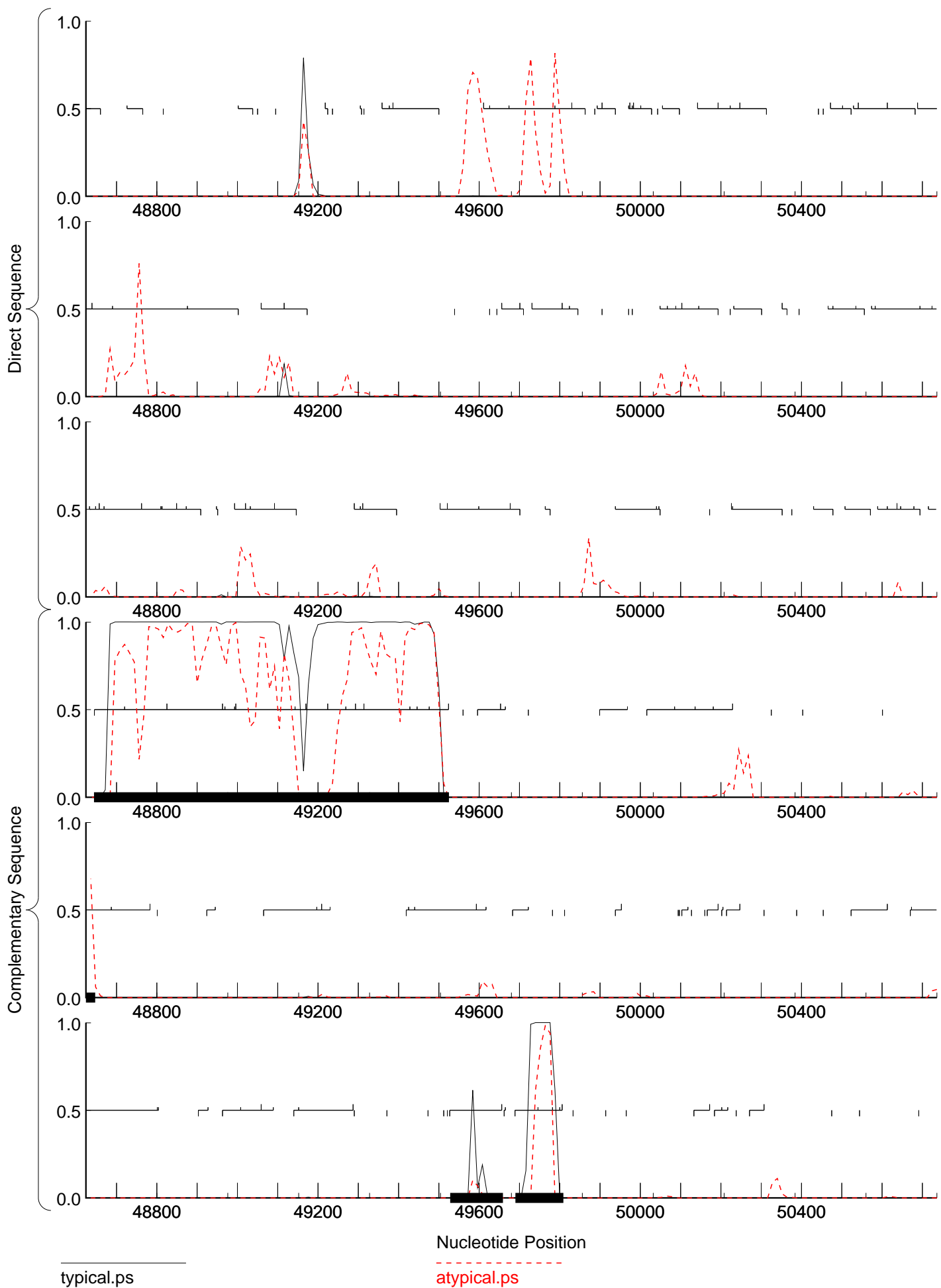
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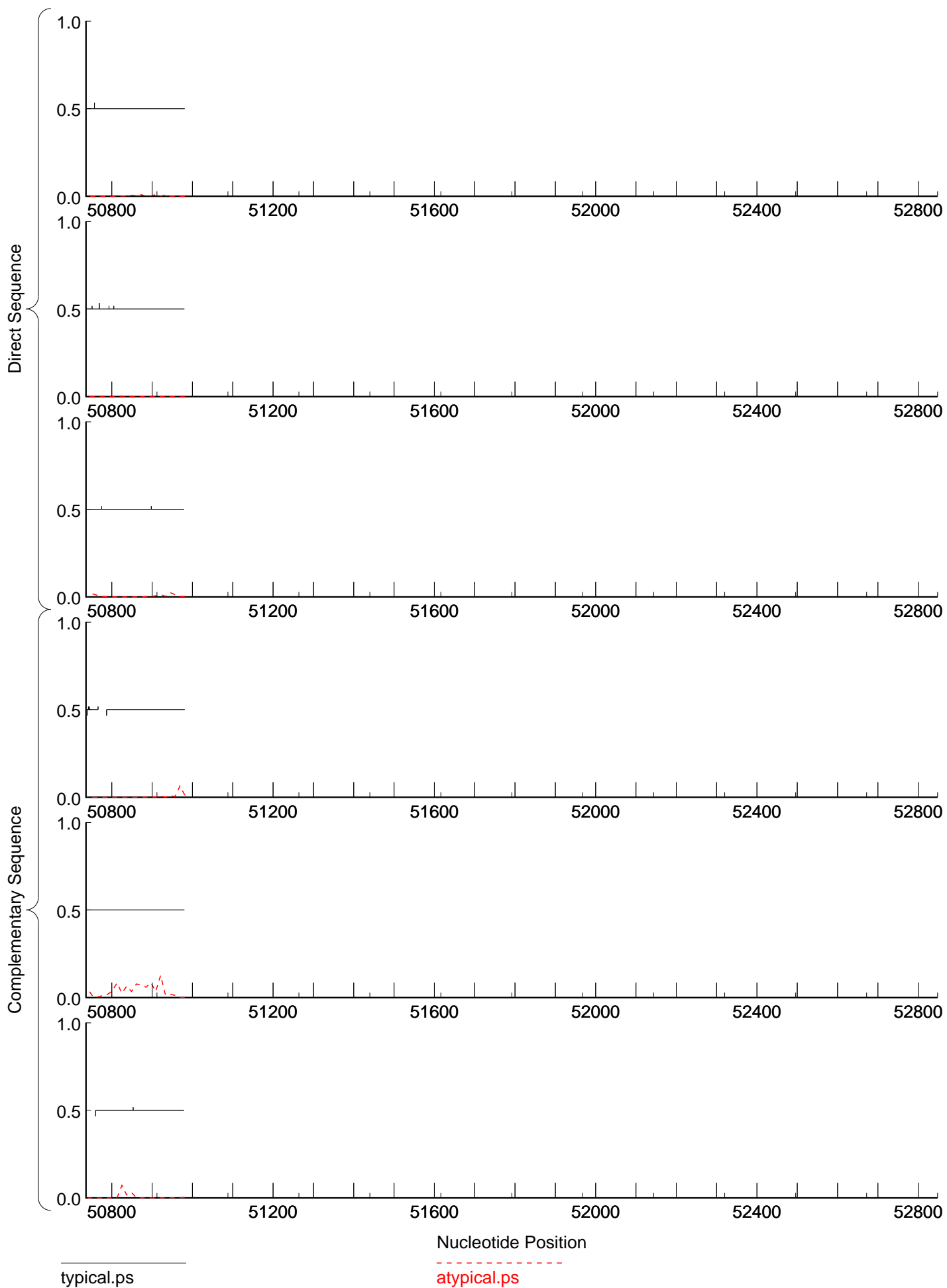
typical.ps

atypical.ps



typical.ps

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