

GeneMark

Version 2.5p (09.08.06)

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PROGRAM INFORMATION

Sequence : Microbacterium phage Lyell complete sequence, 62716 bp including 3549 bp terminal repeat, C
Analysis Date : 6/24/18 at 10:40:44
Pages : 31
Sequence Length : 62716 bp
GC Content : 61.60%

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

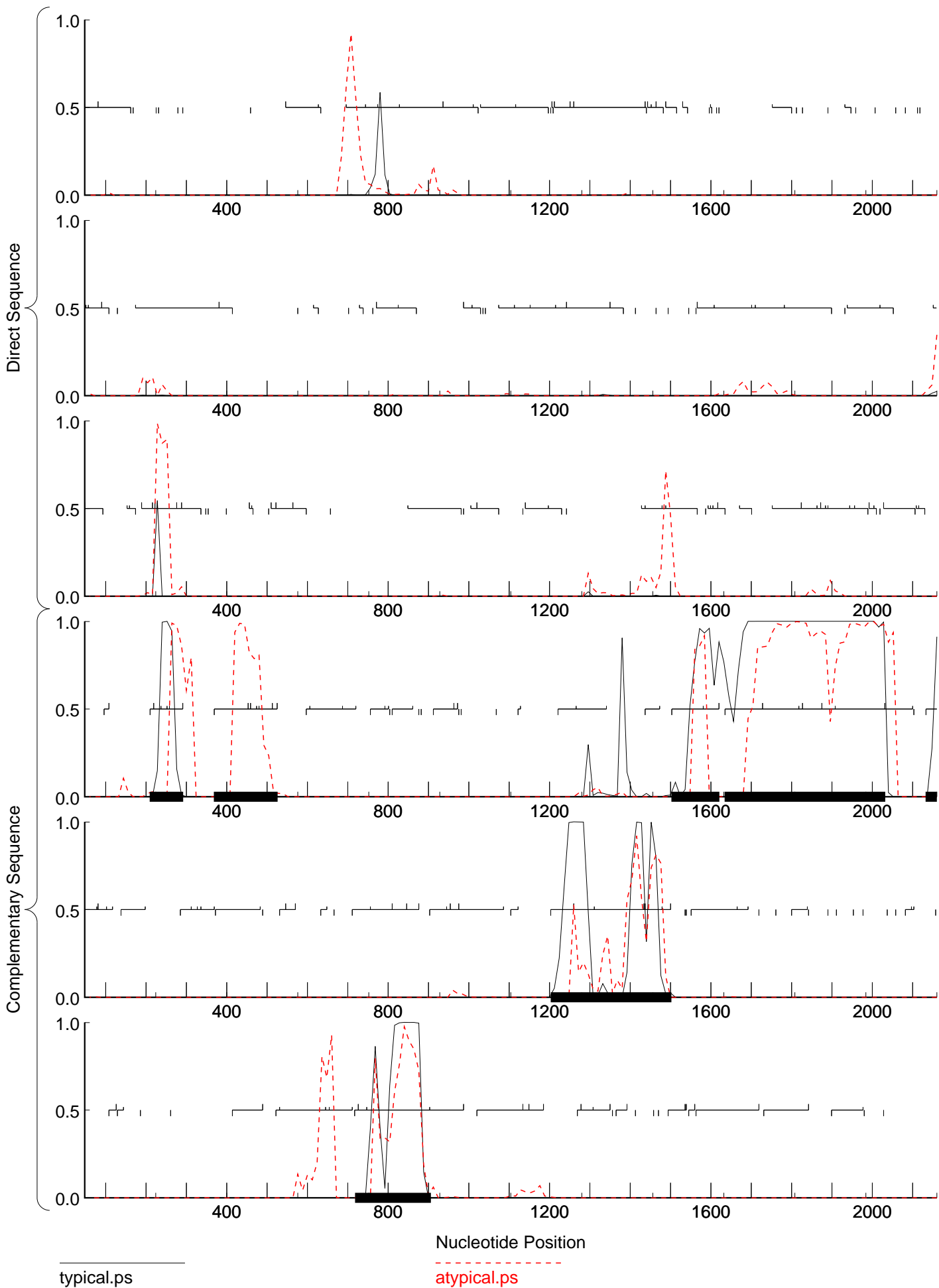
Training set derived by GeneMarkS, 4.30 October 2014
Sun Jun 24 10:40:44 2018

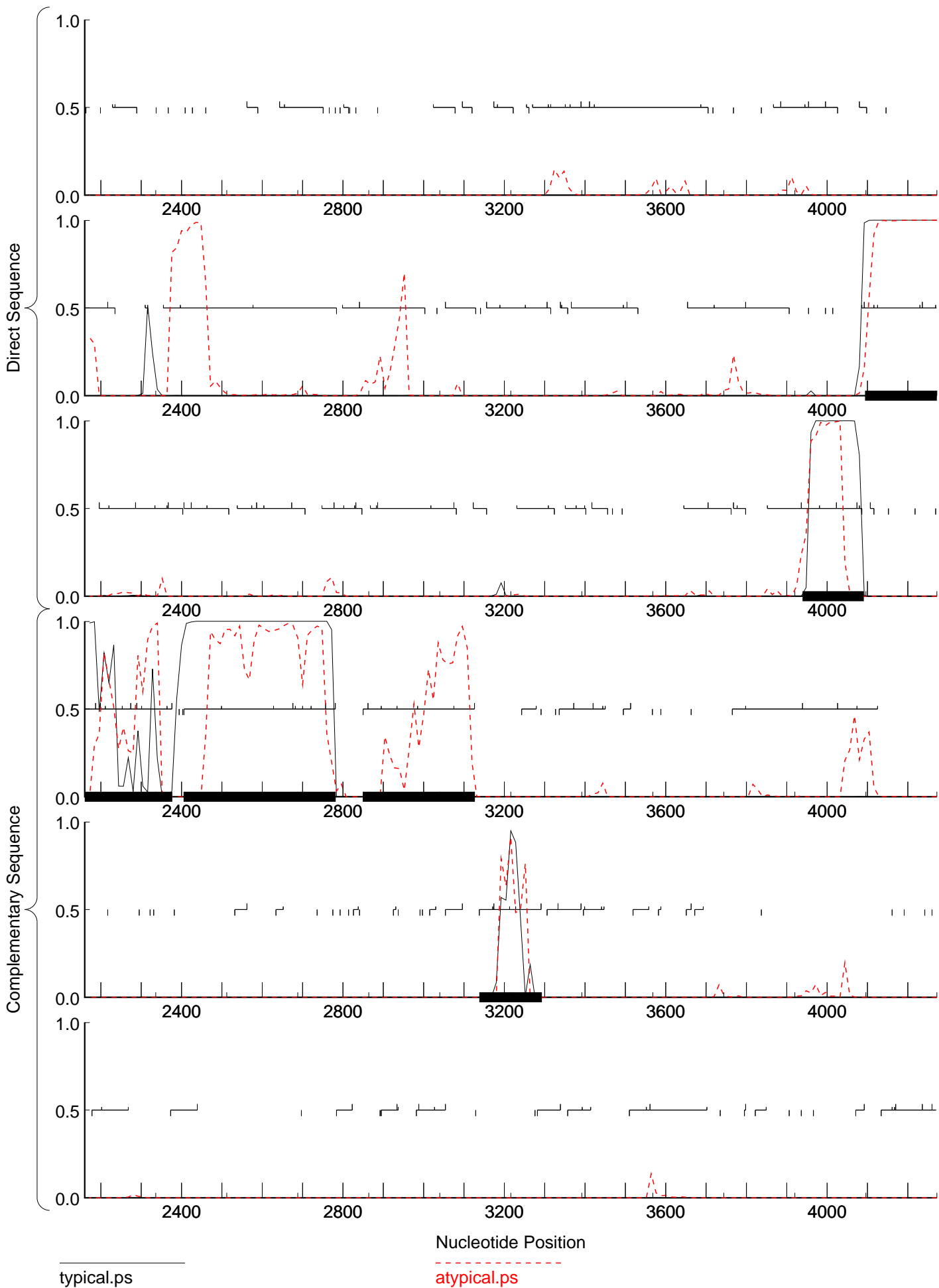
MATRIX INFORMATION

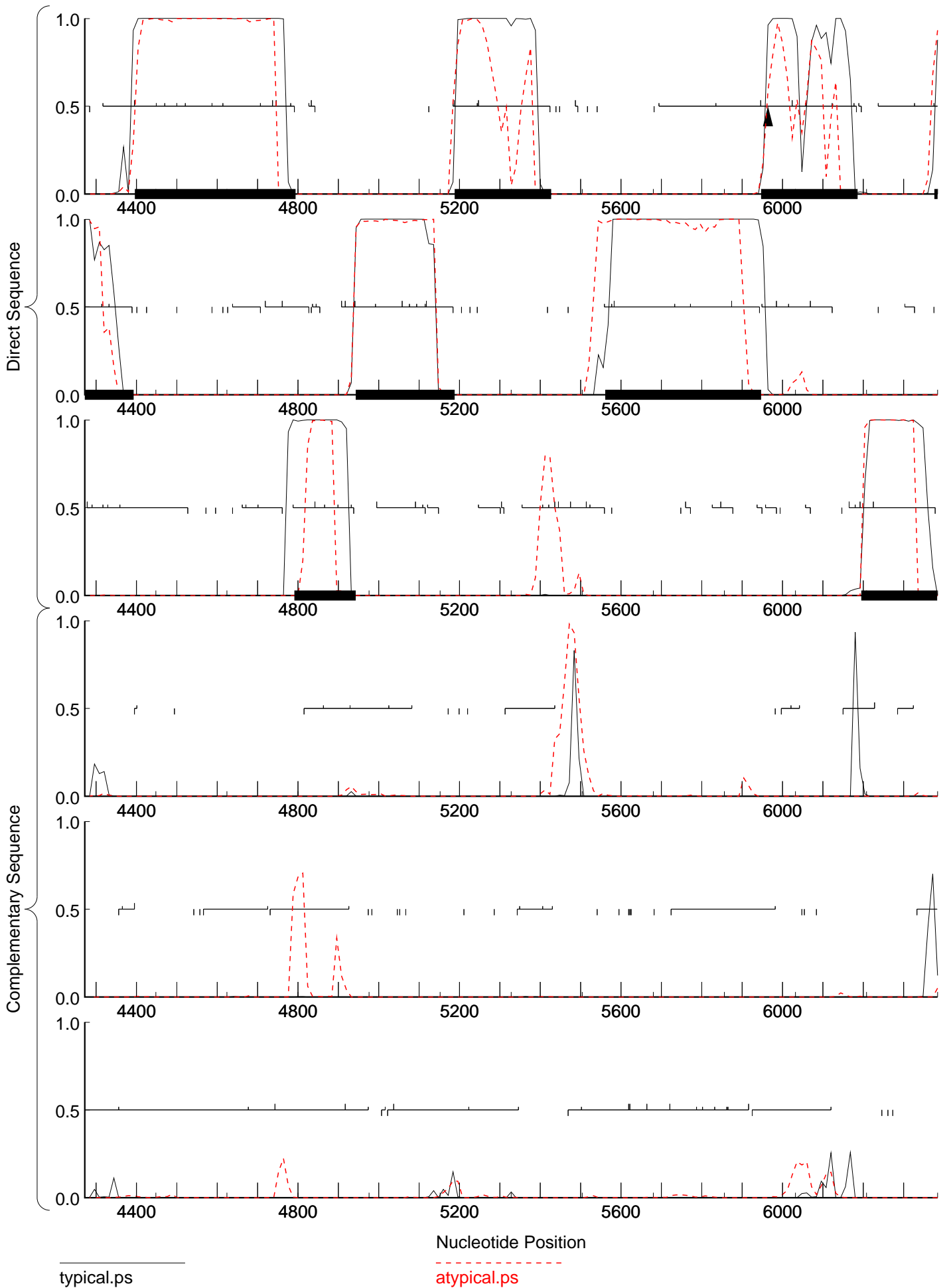
Matrix : -
Author : -
Order : 4

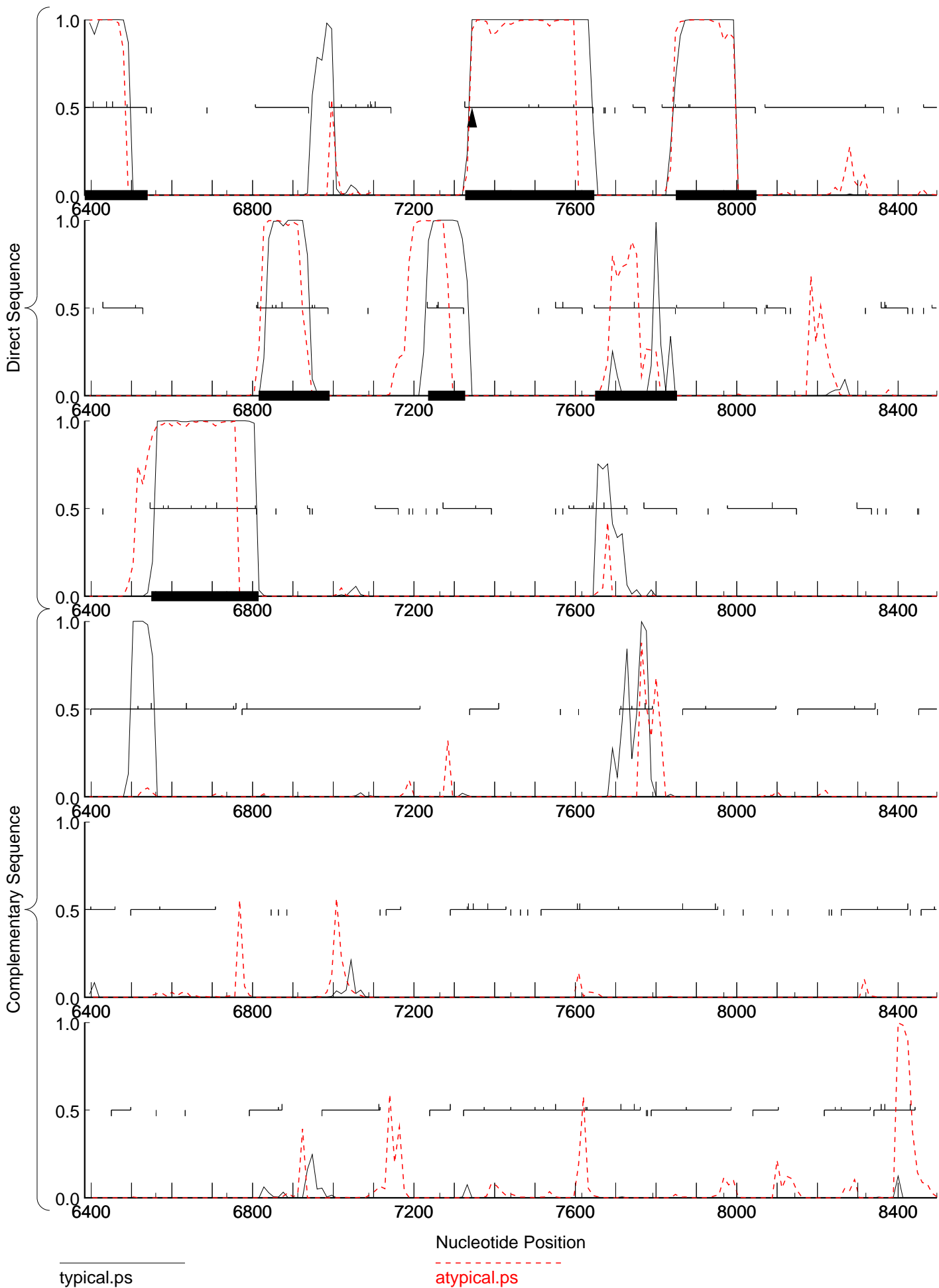
Send questions / comments to:
Dr. M. Borodovsky
Georgia Institute of Technology
School of Biology
Atlanta, GA 30332-0230

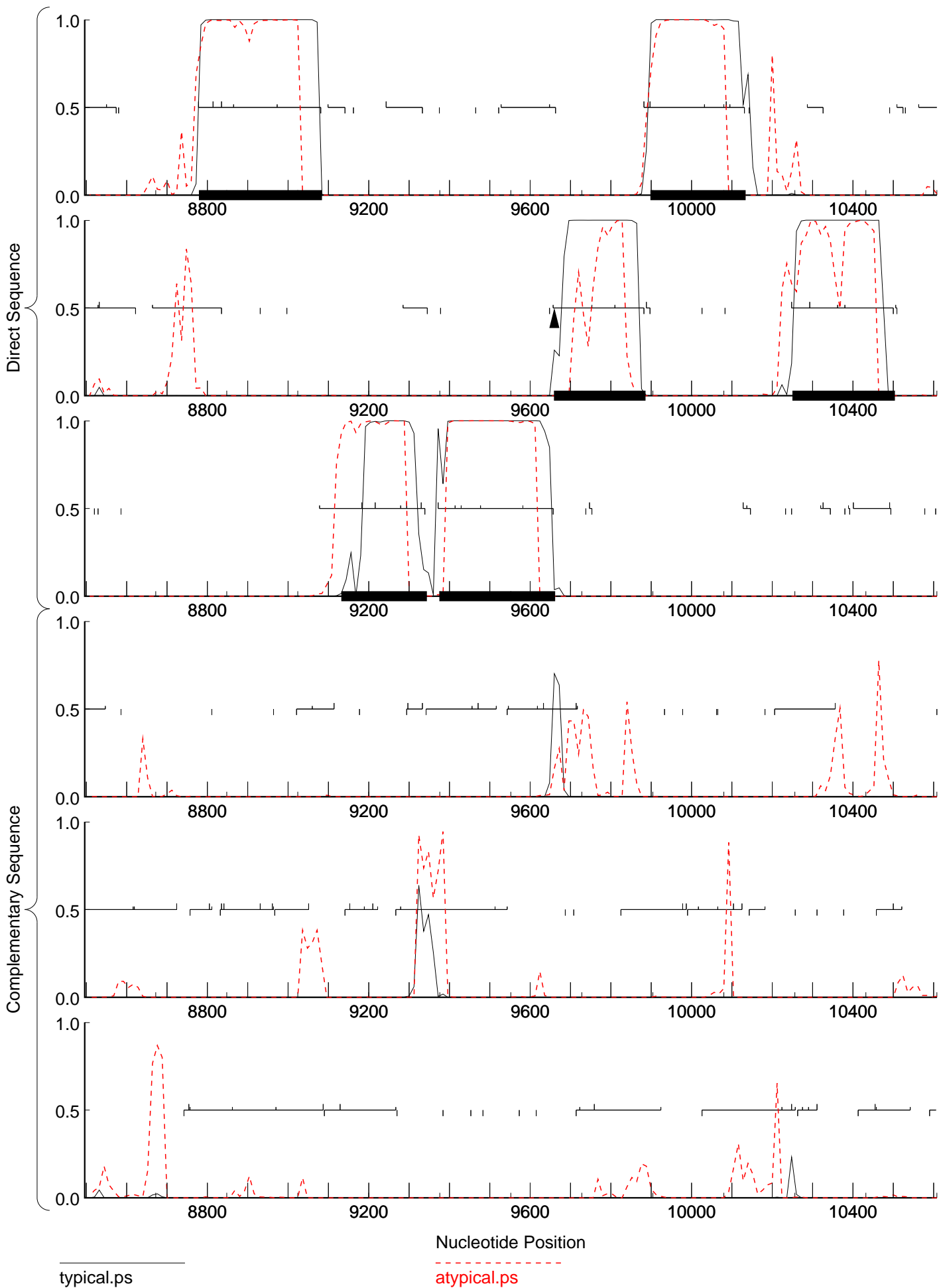
For help or information, send electronic mail to: custserv@genepro.com, or mark@amber.biology.gatech.edu

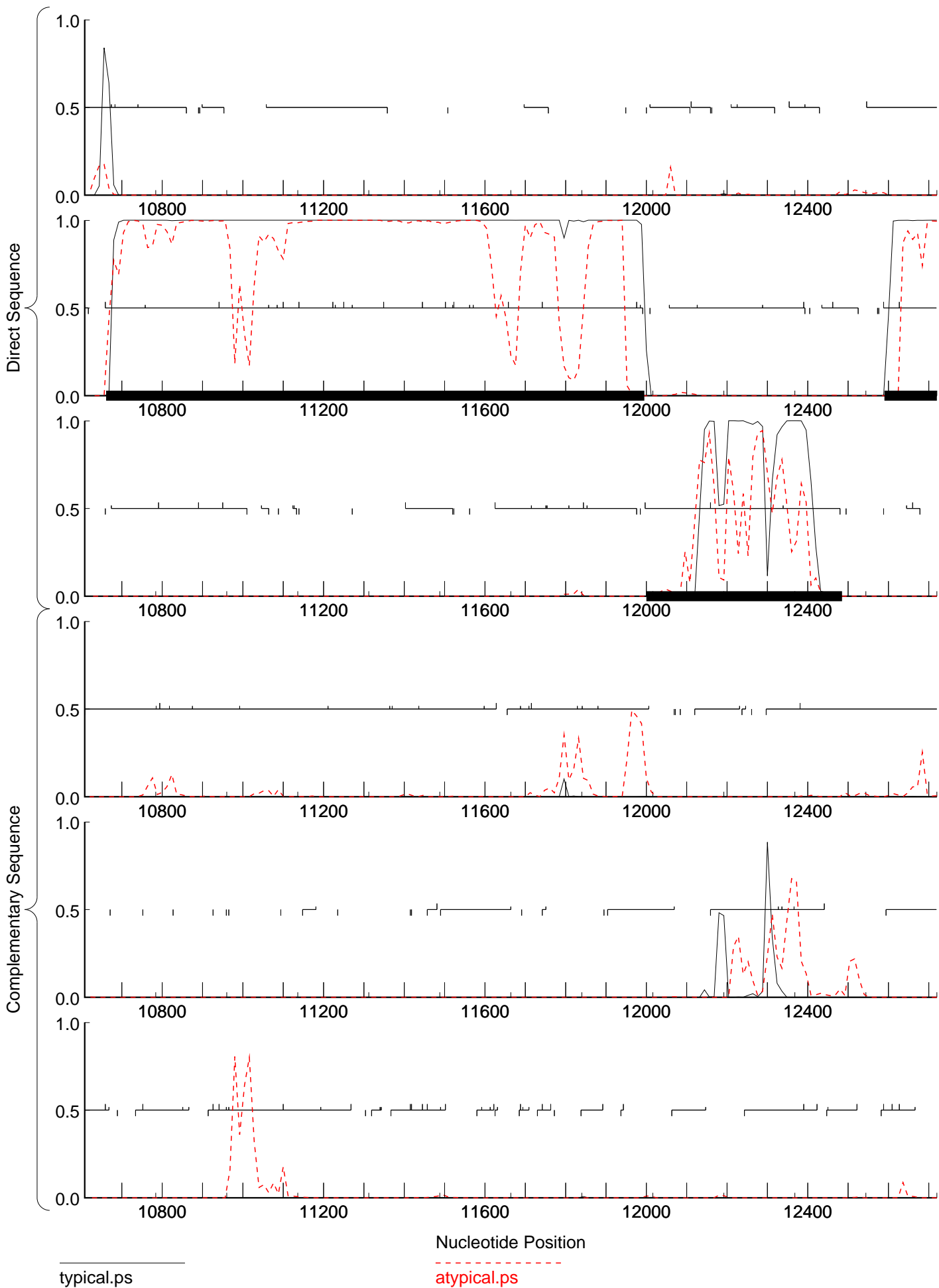






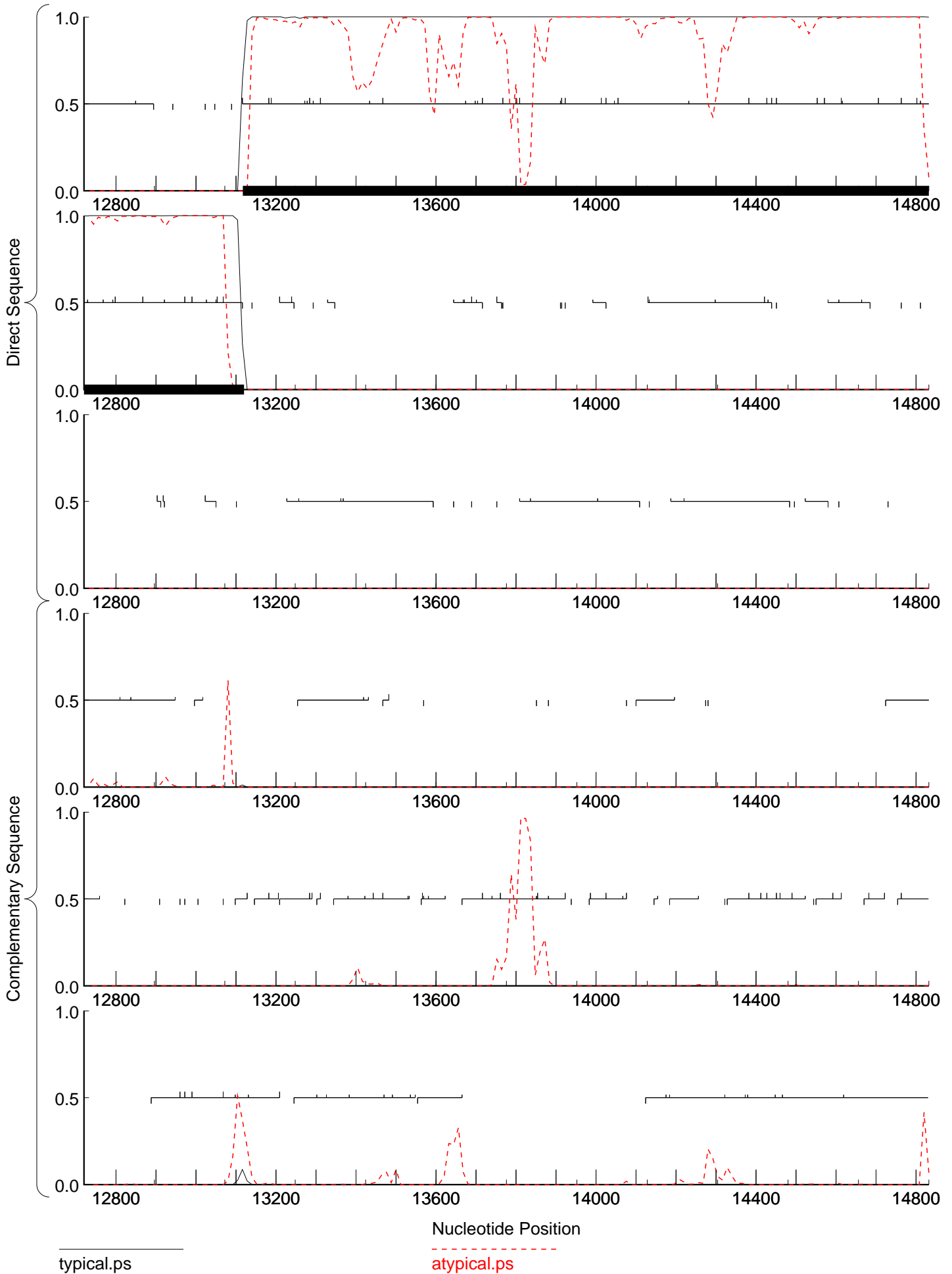


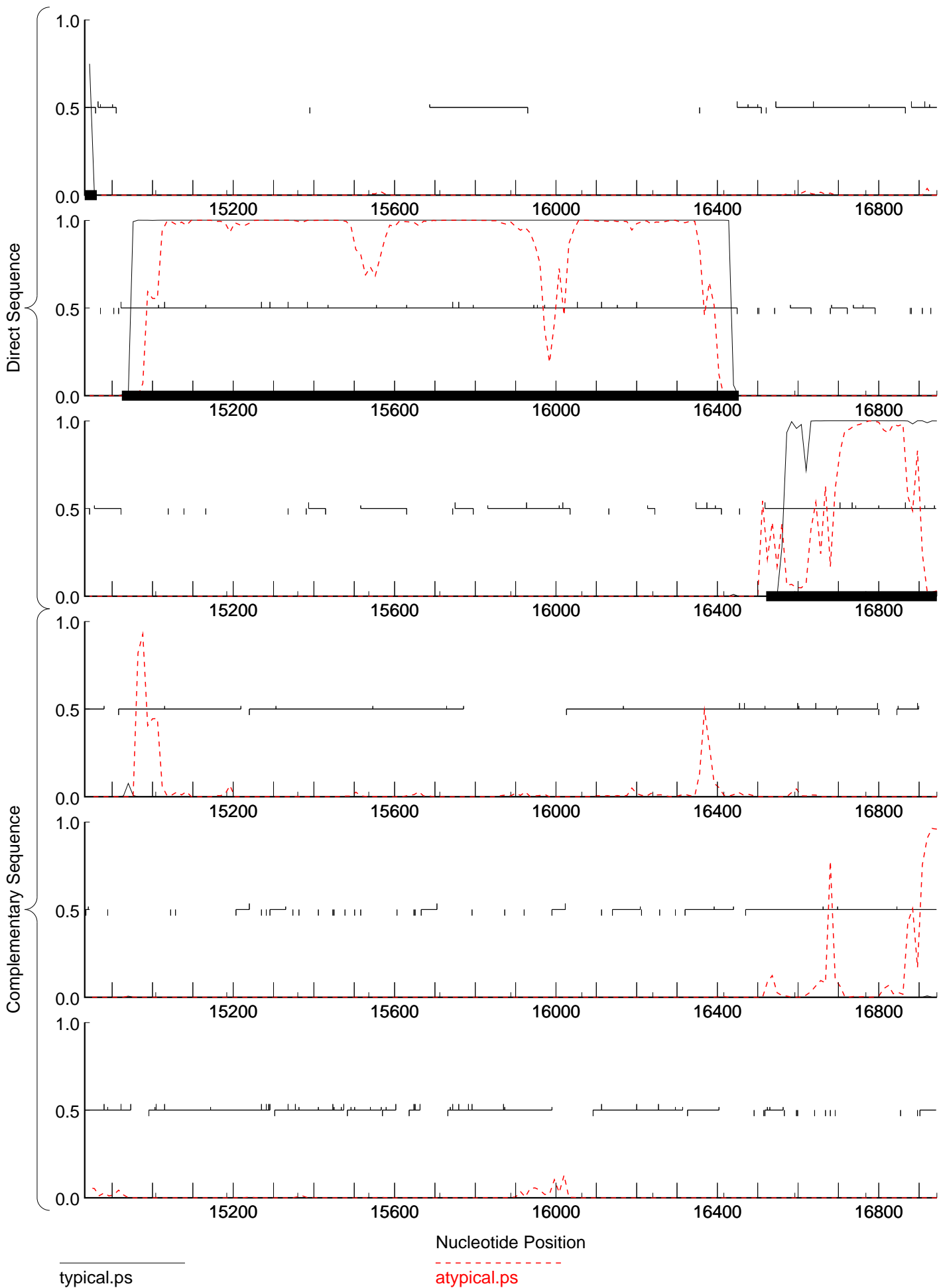


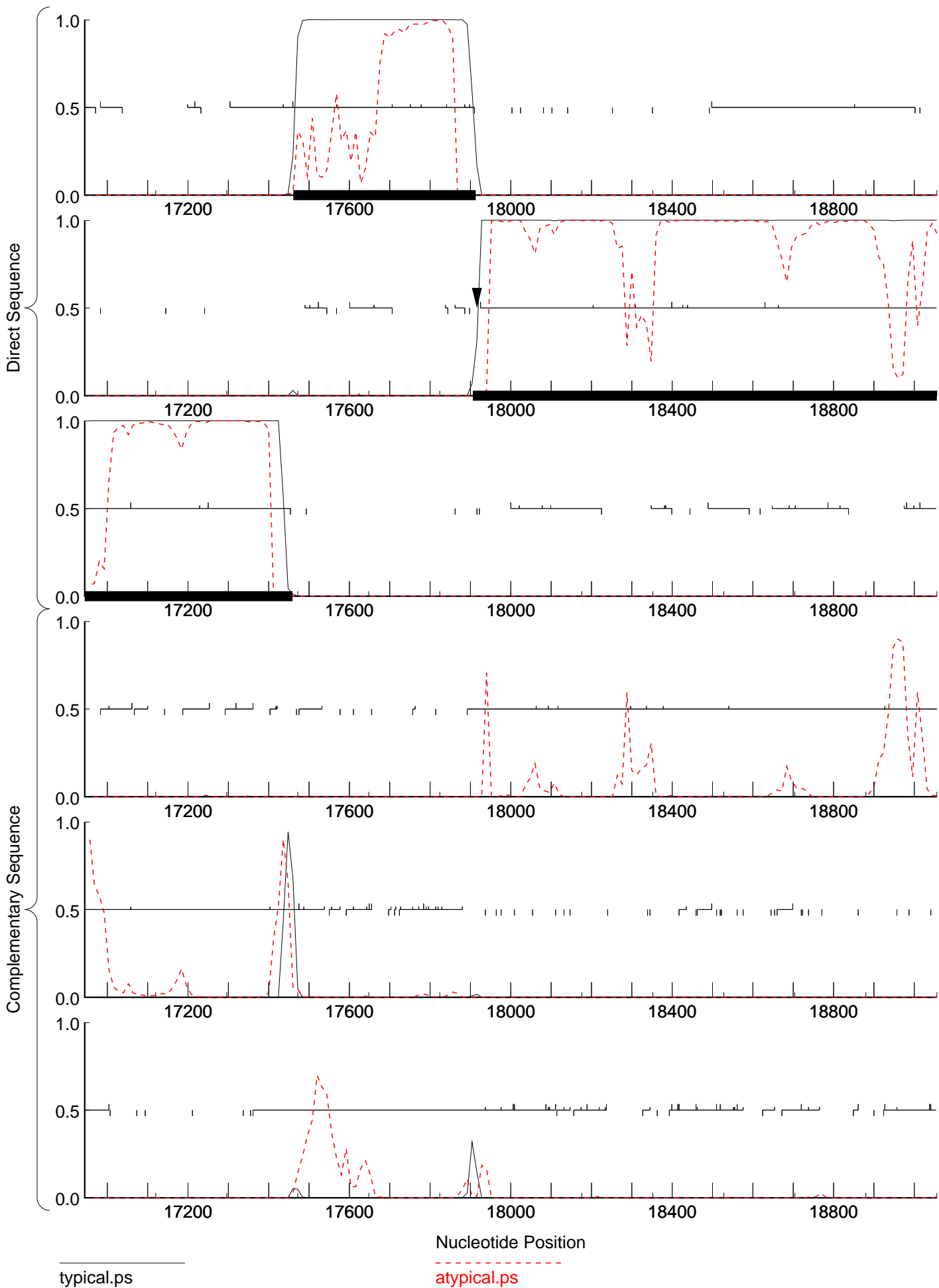


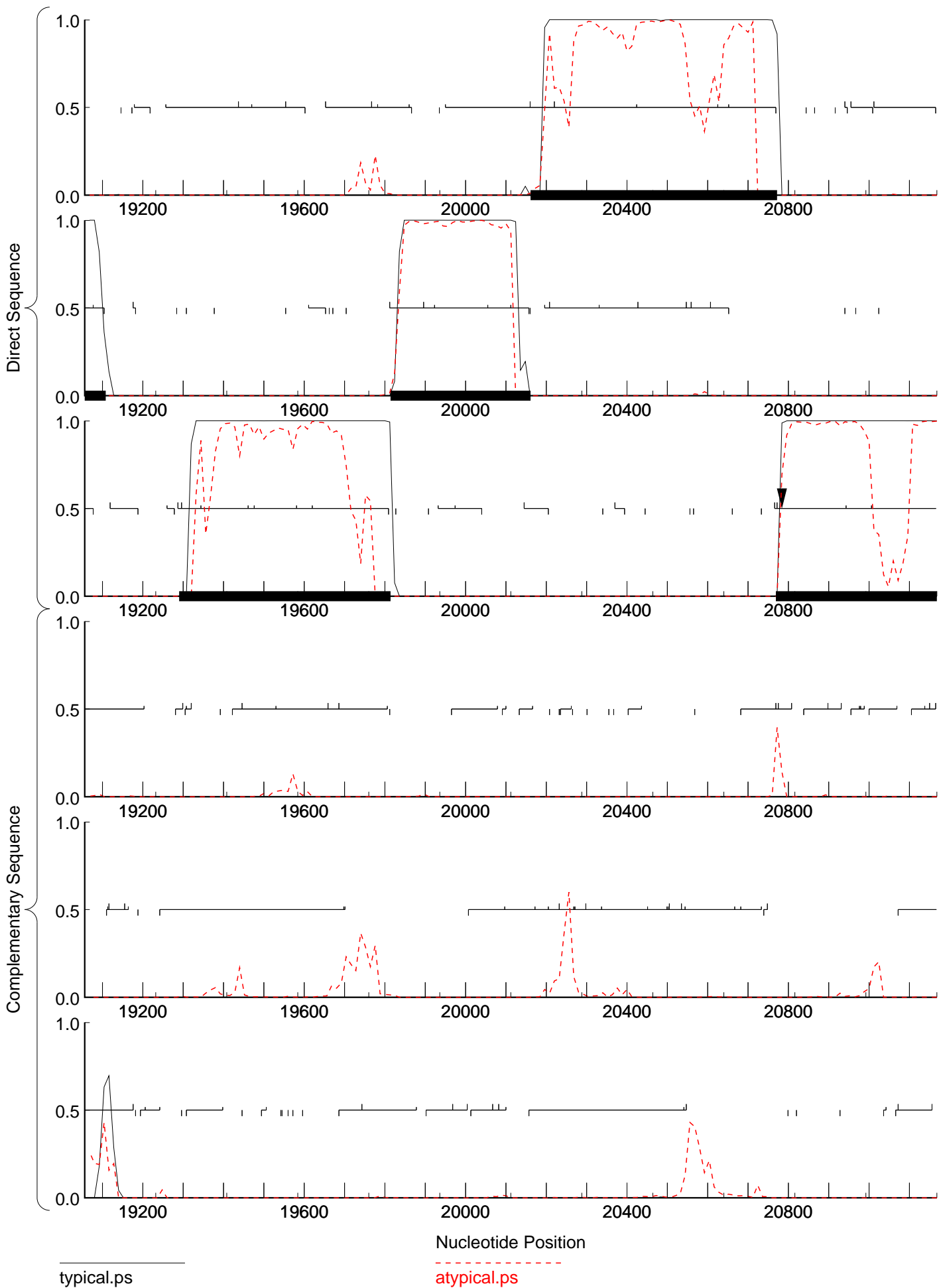
typical.ps

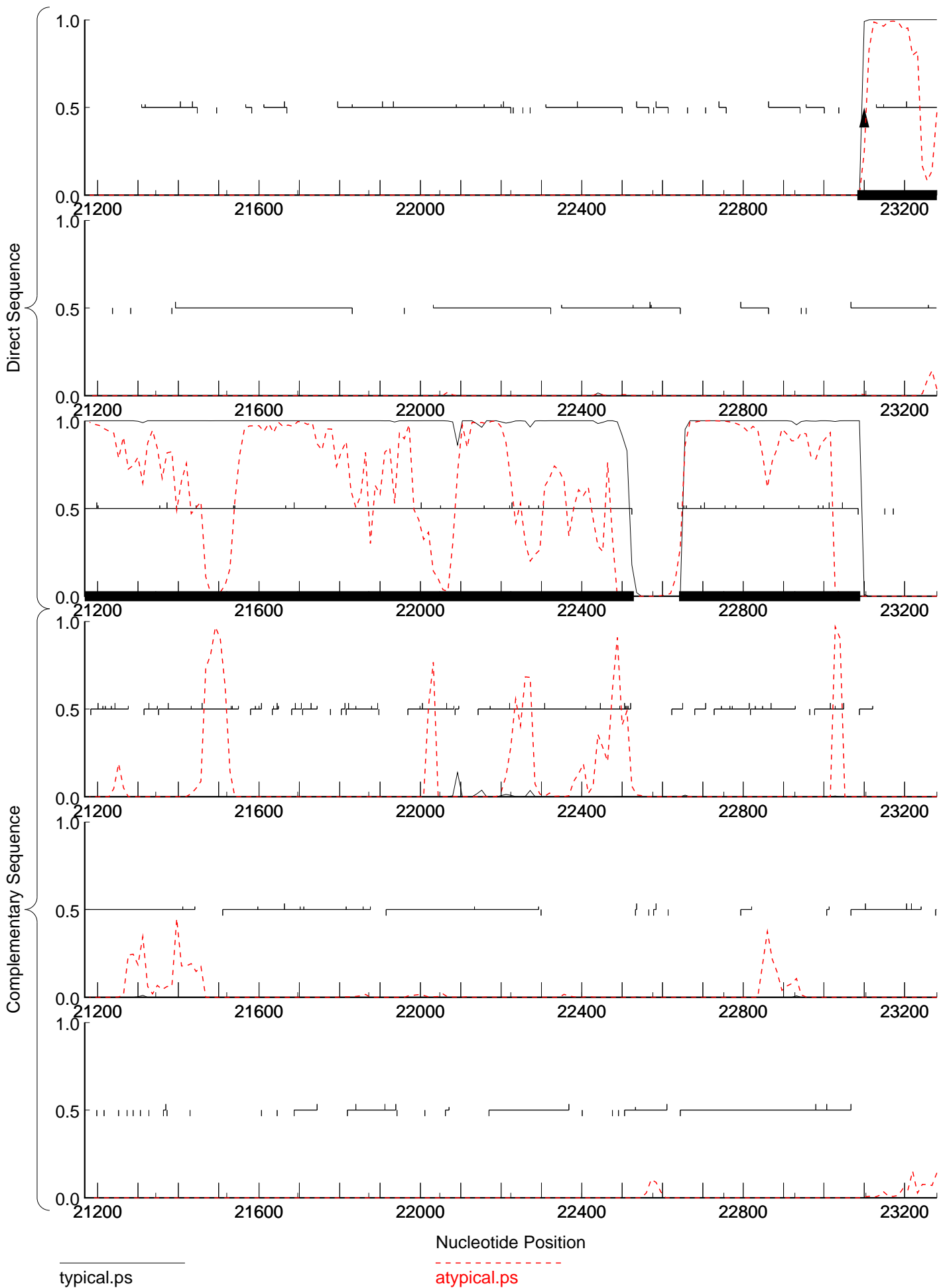
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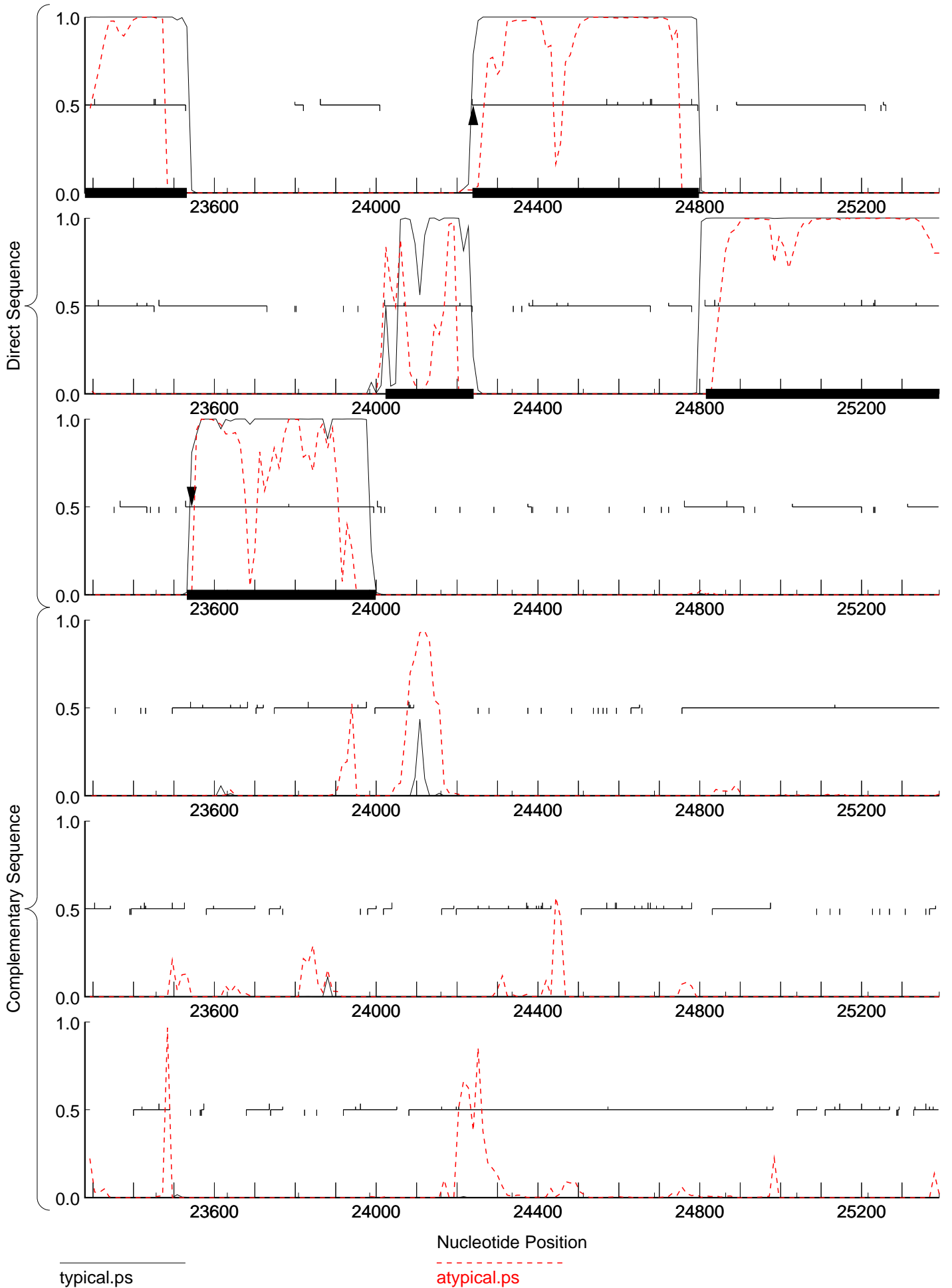








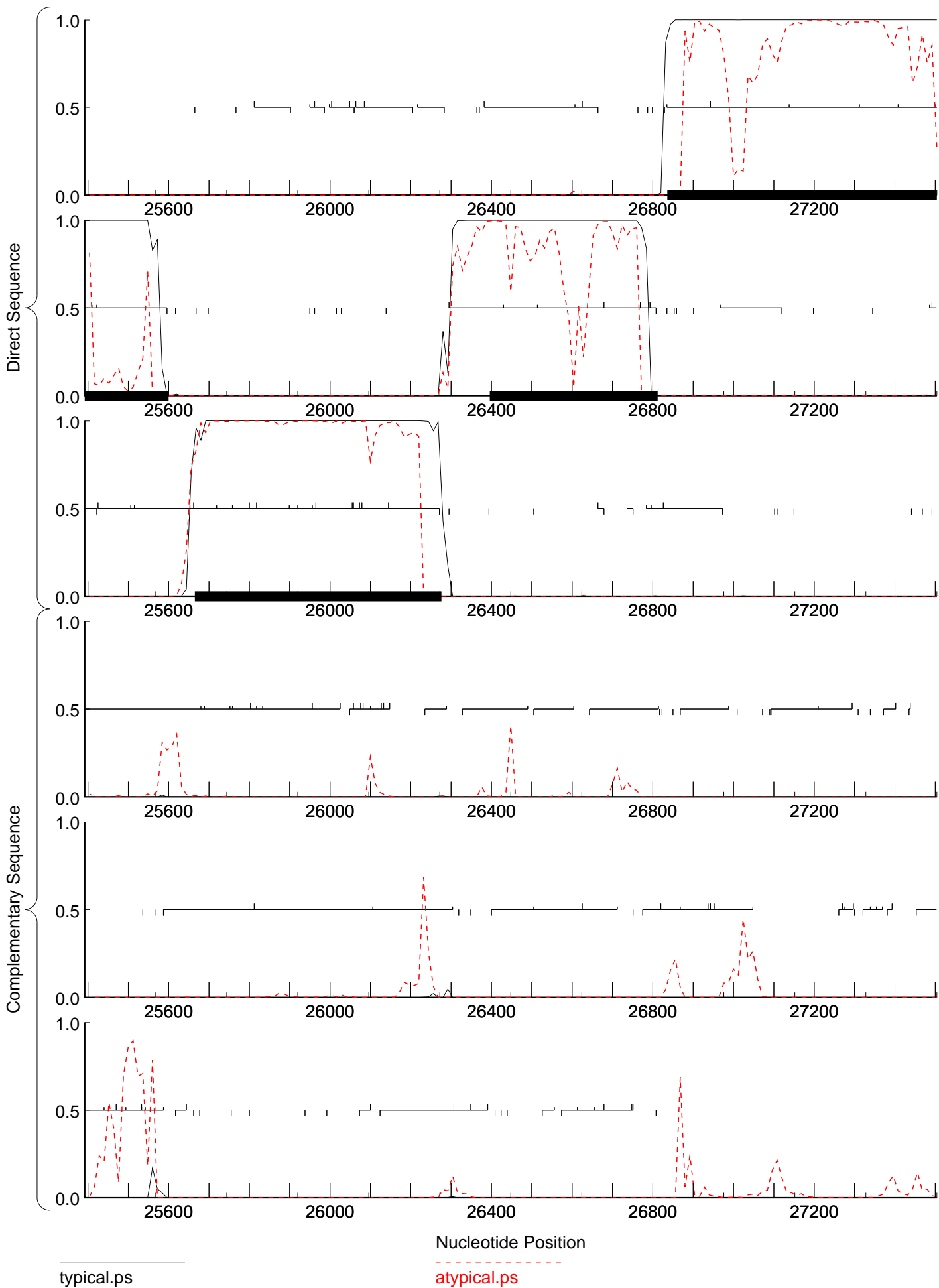




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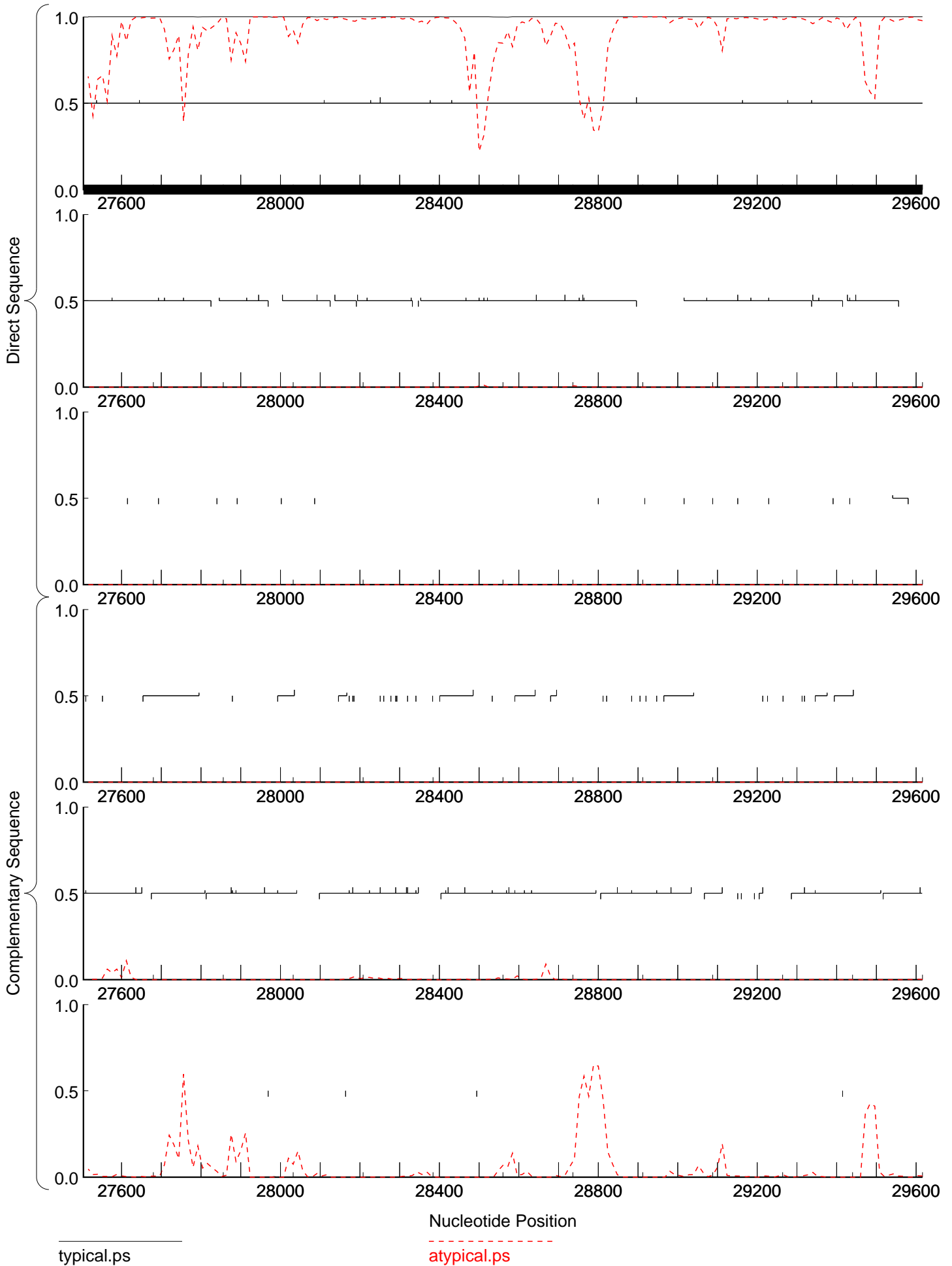
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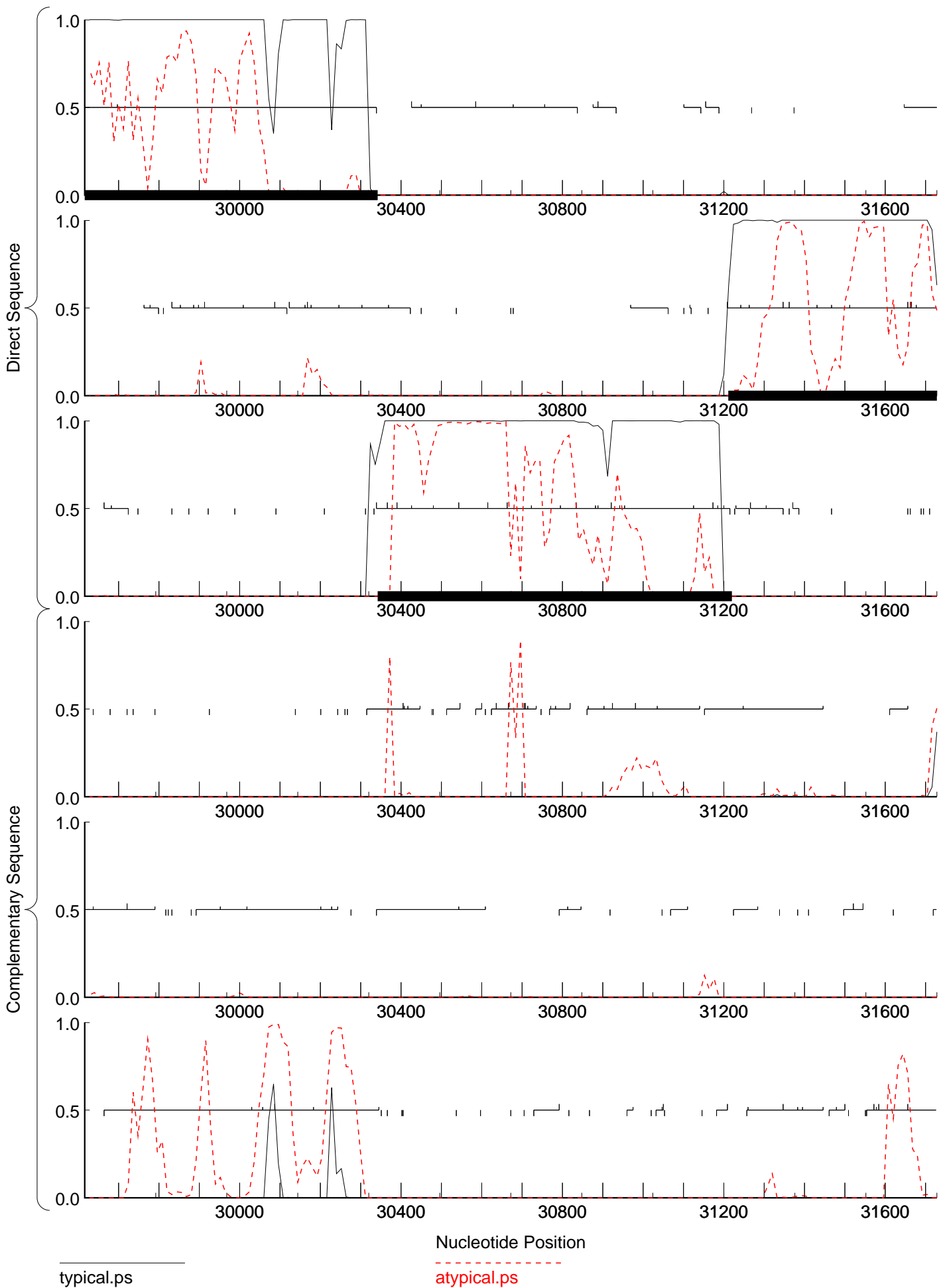
GeneMark.hmm prediction

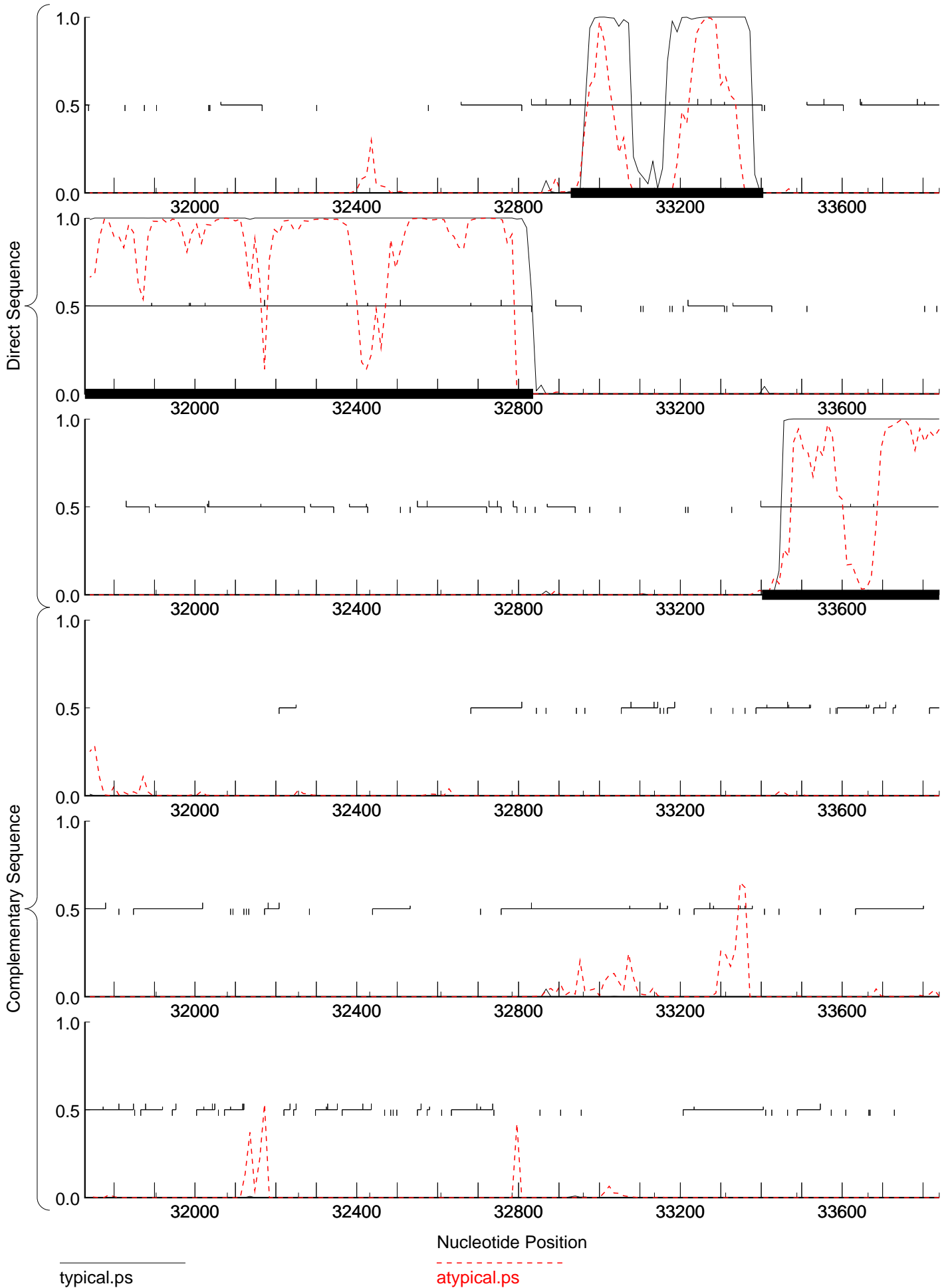


typical.ps

atypical.ps

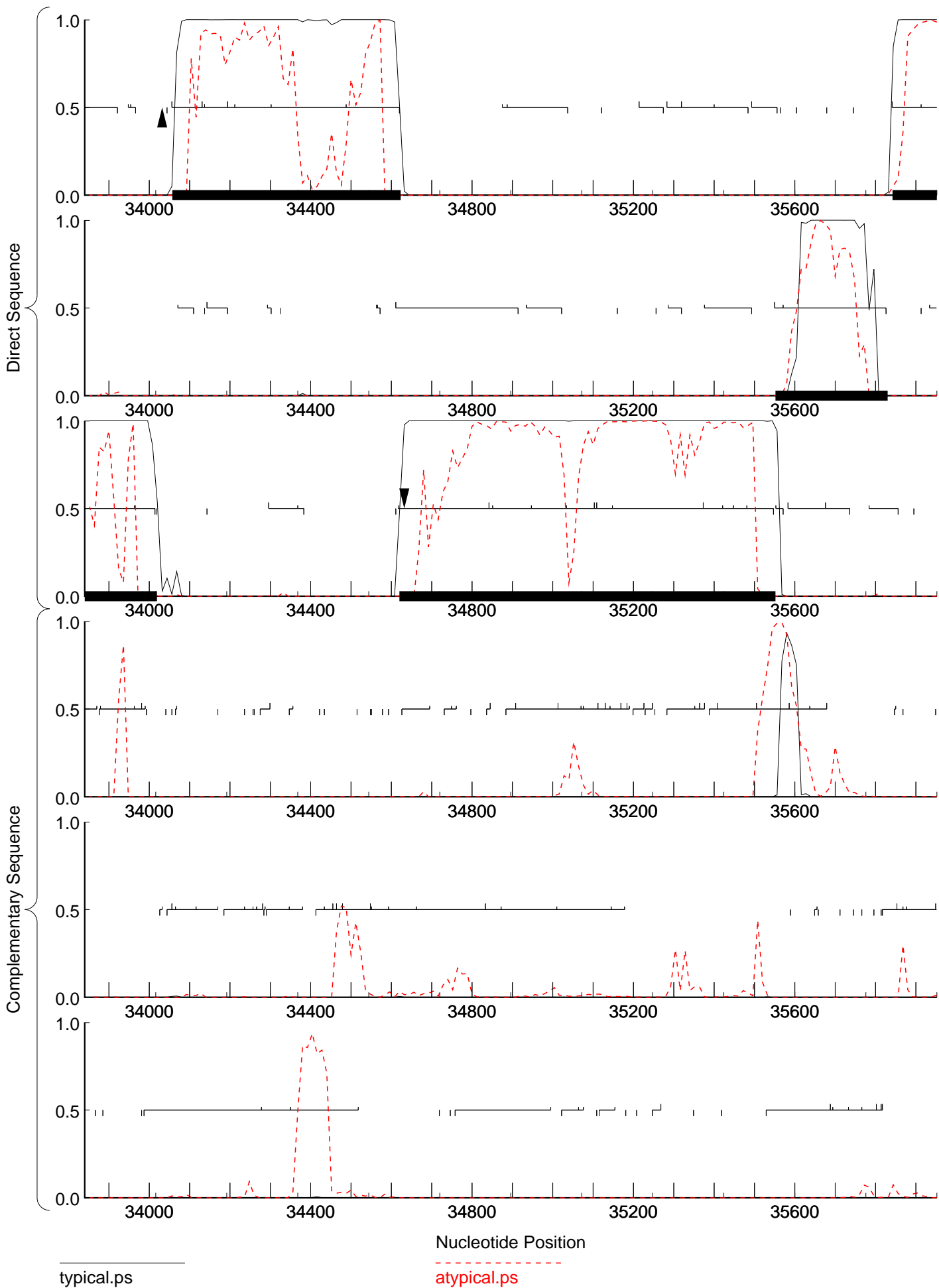


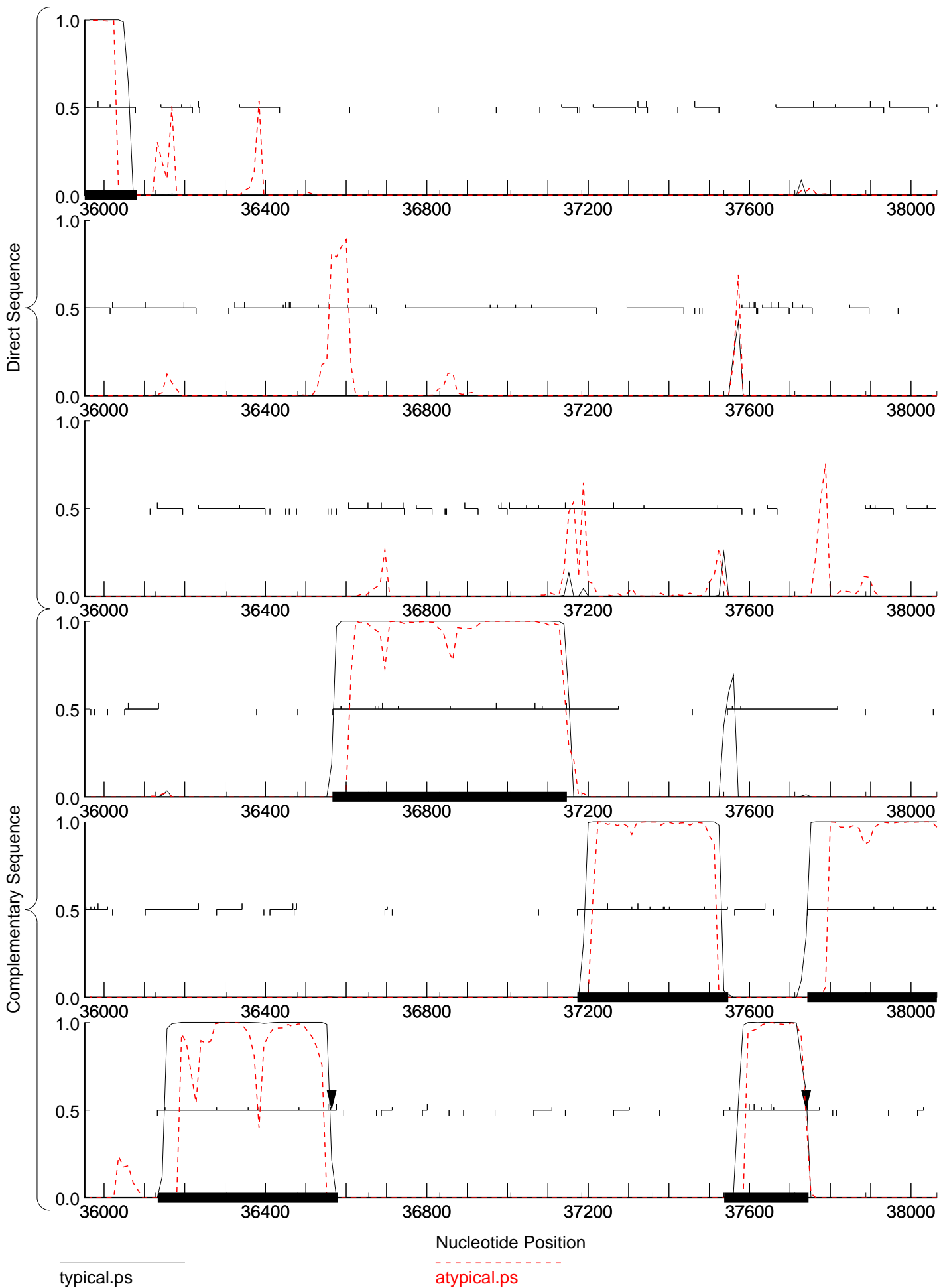




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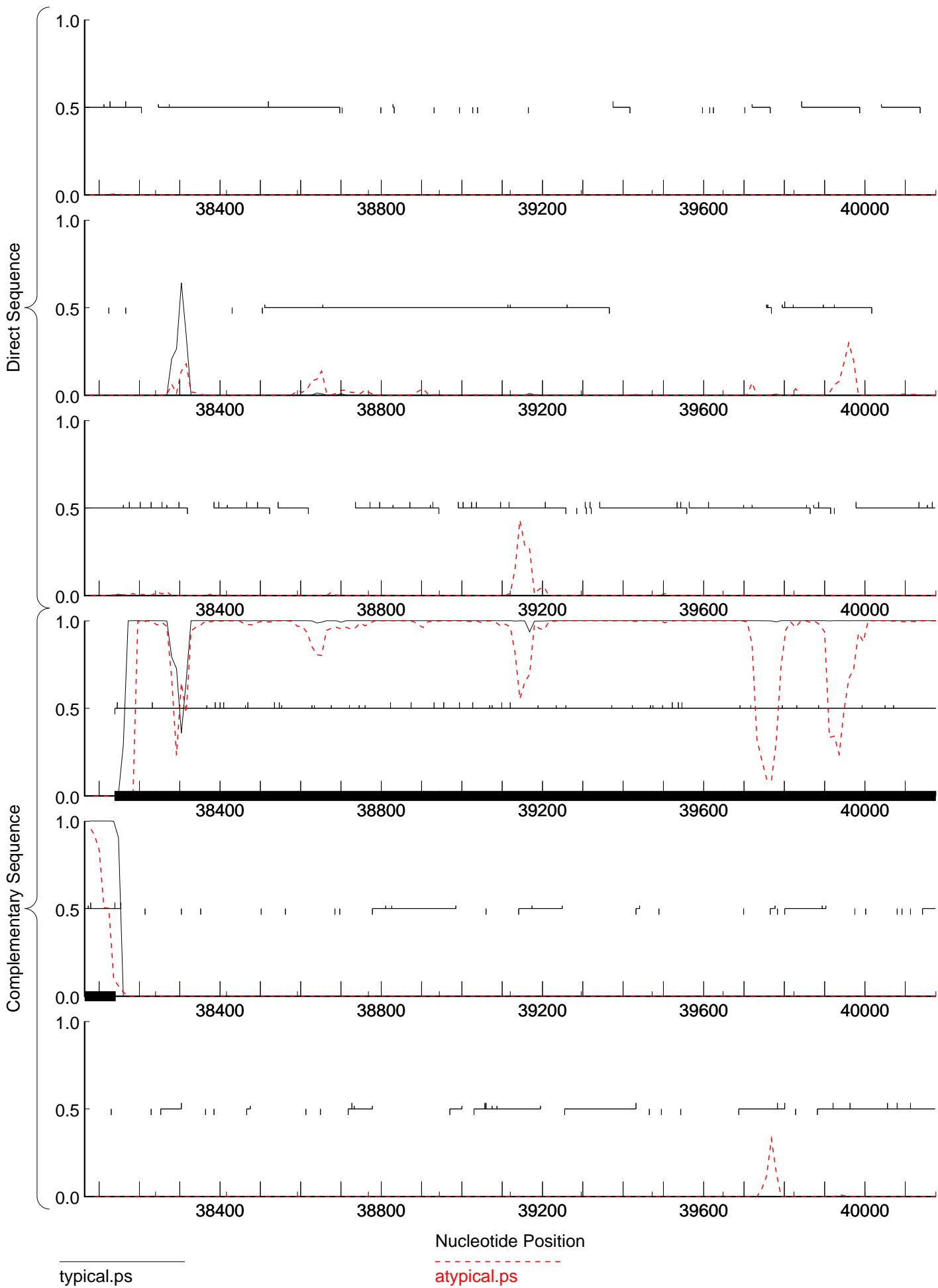
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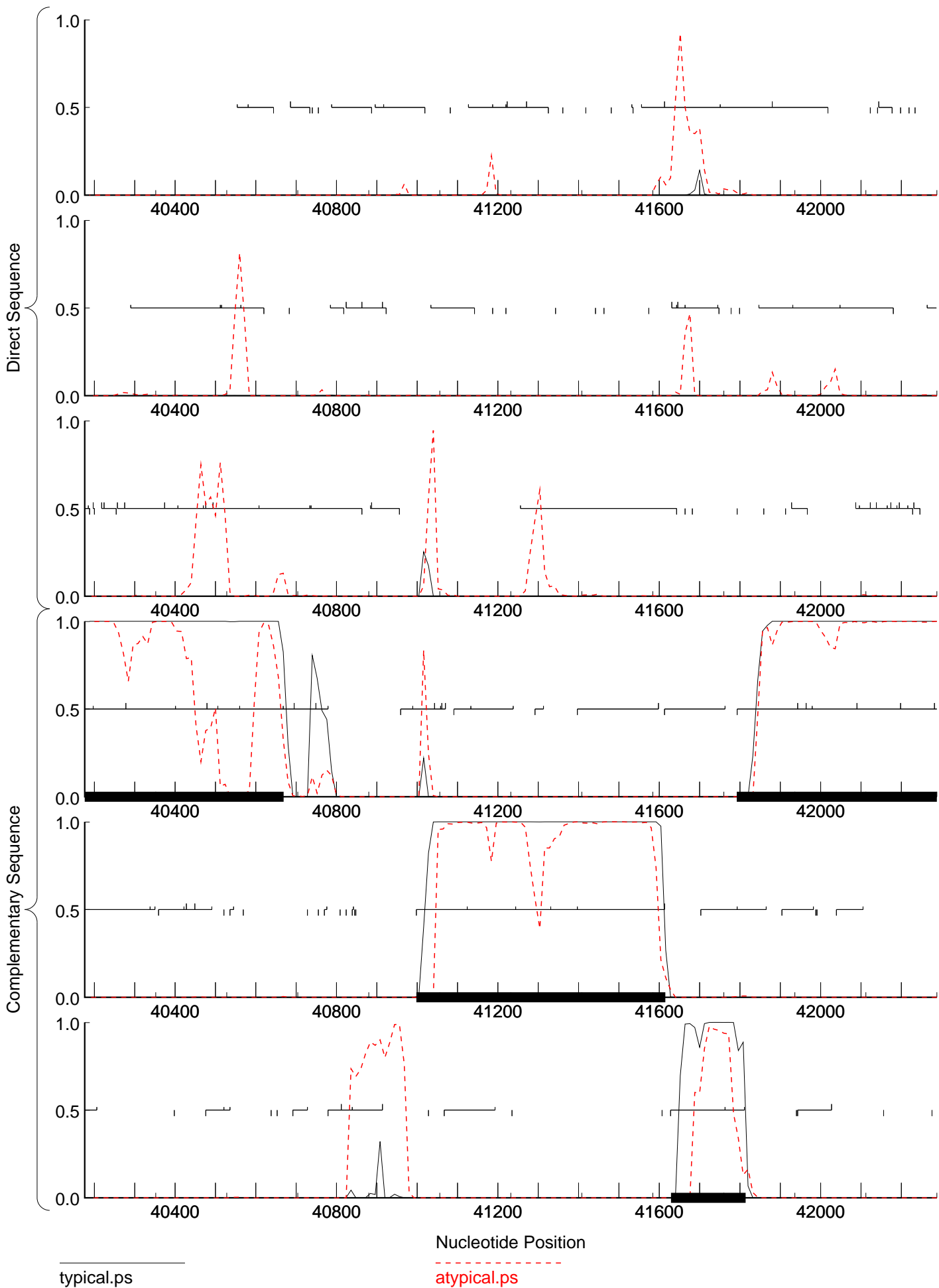


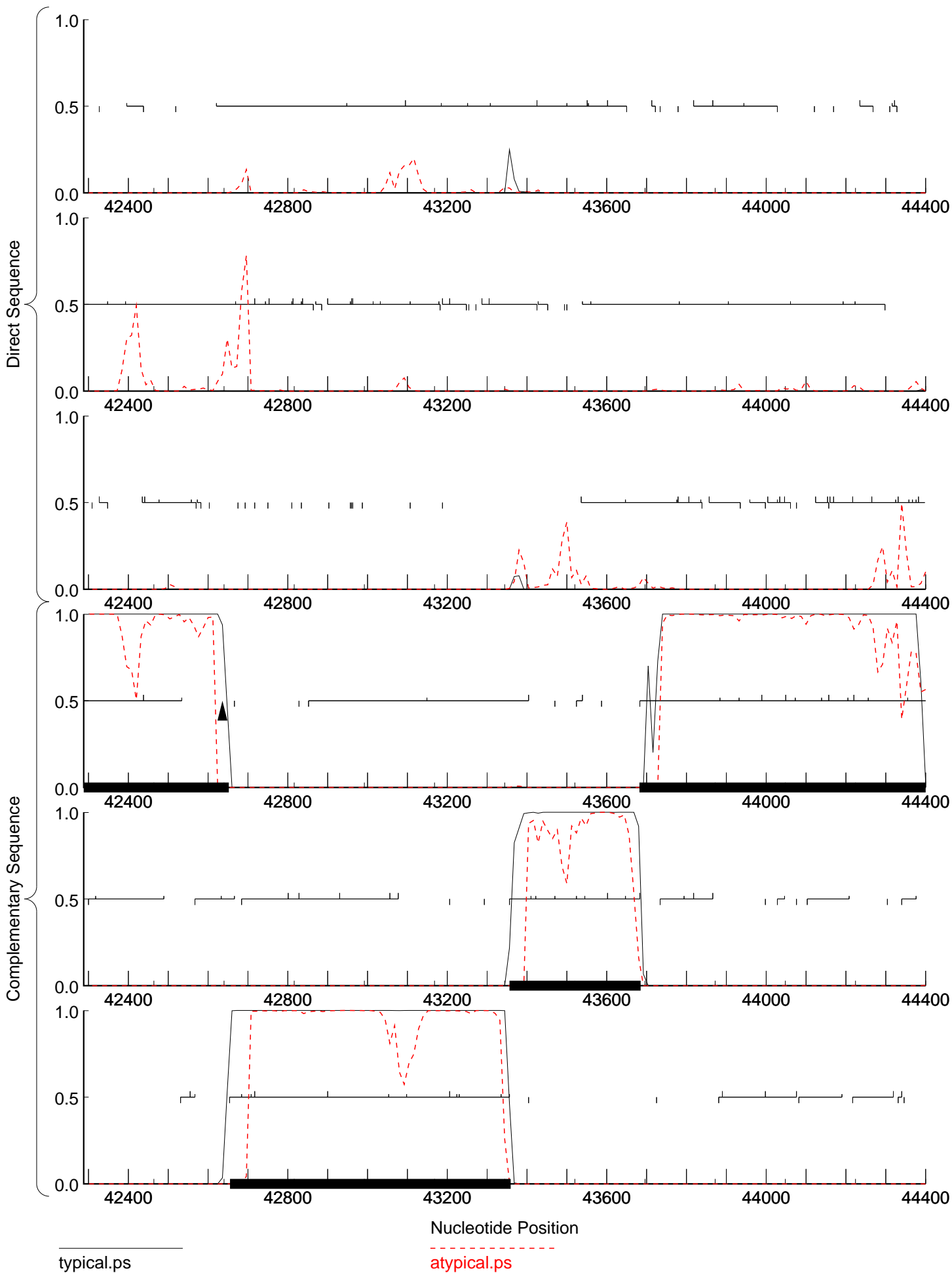


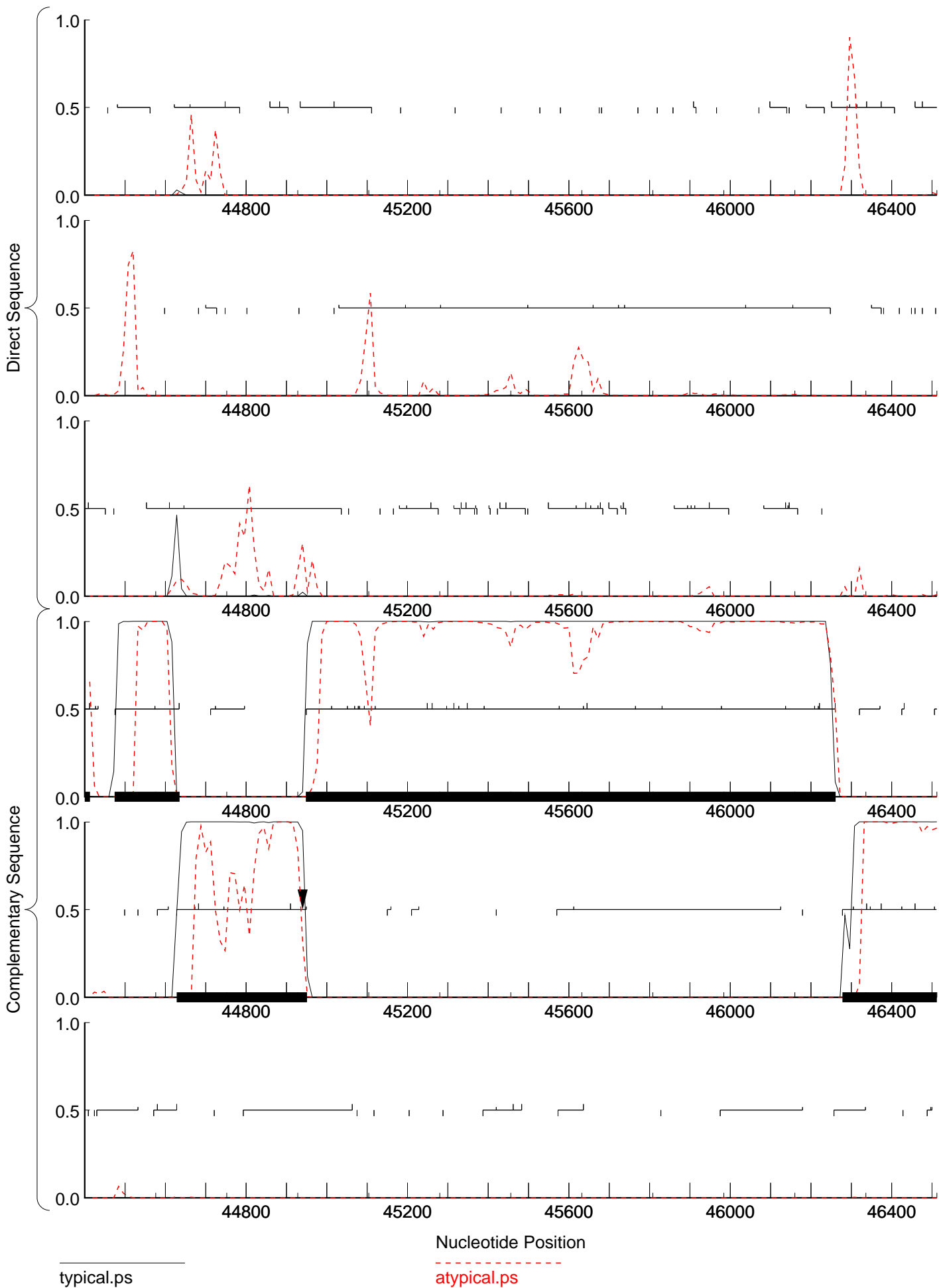
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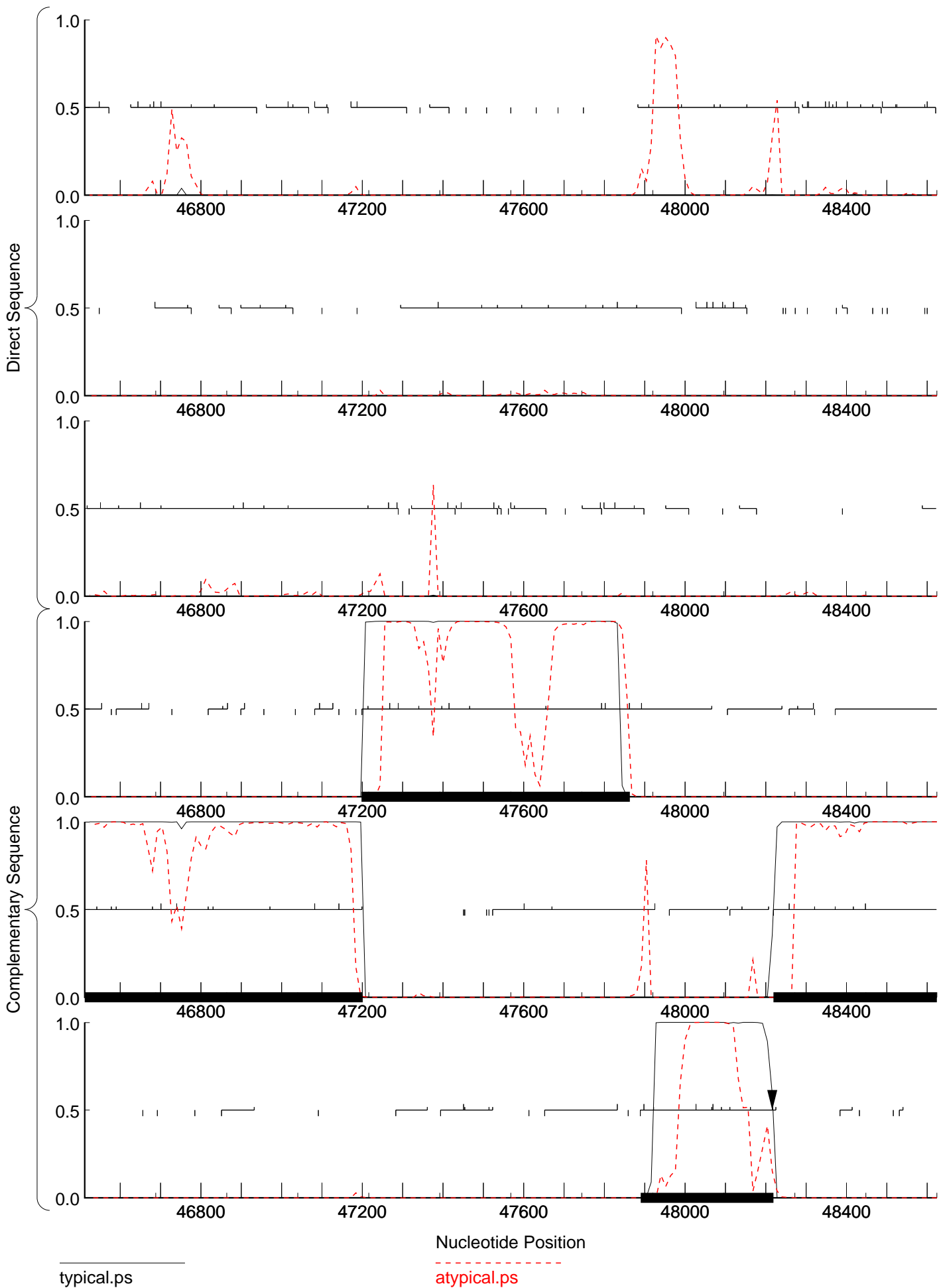
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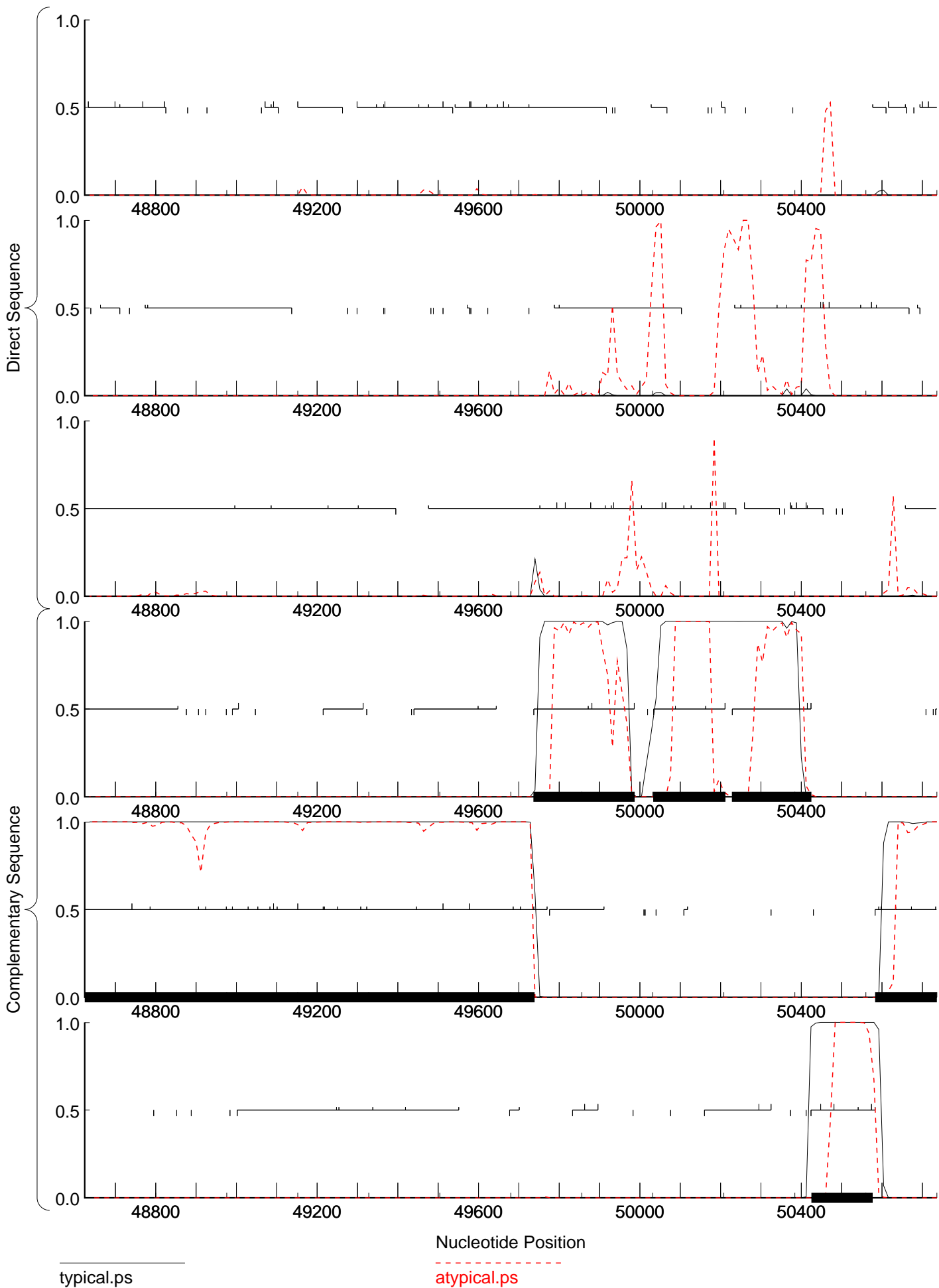


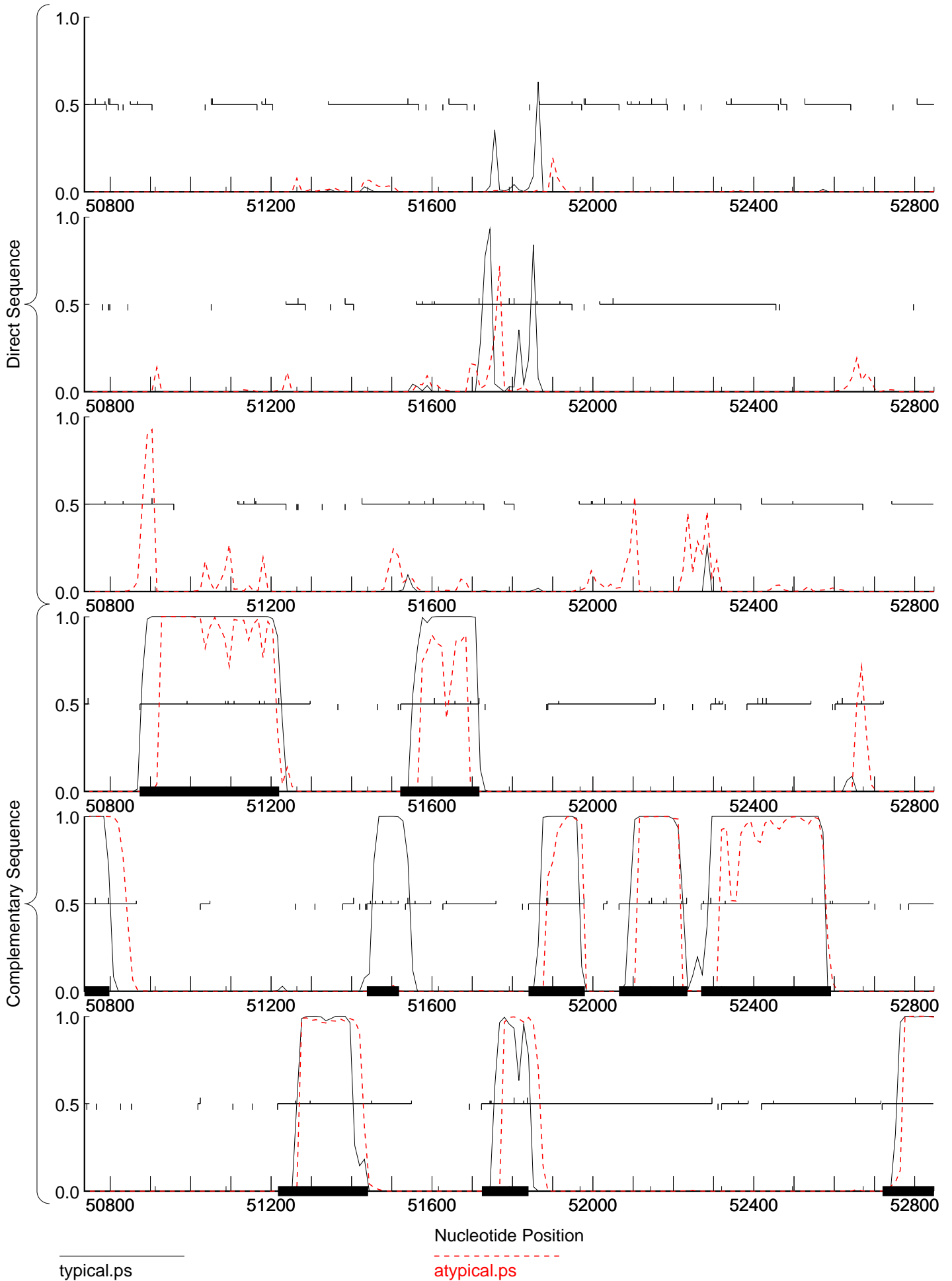


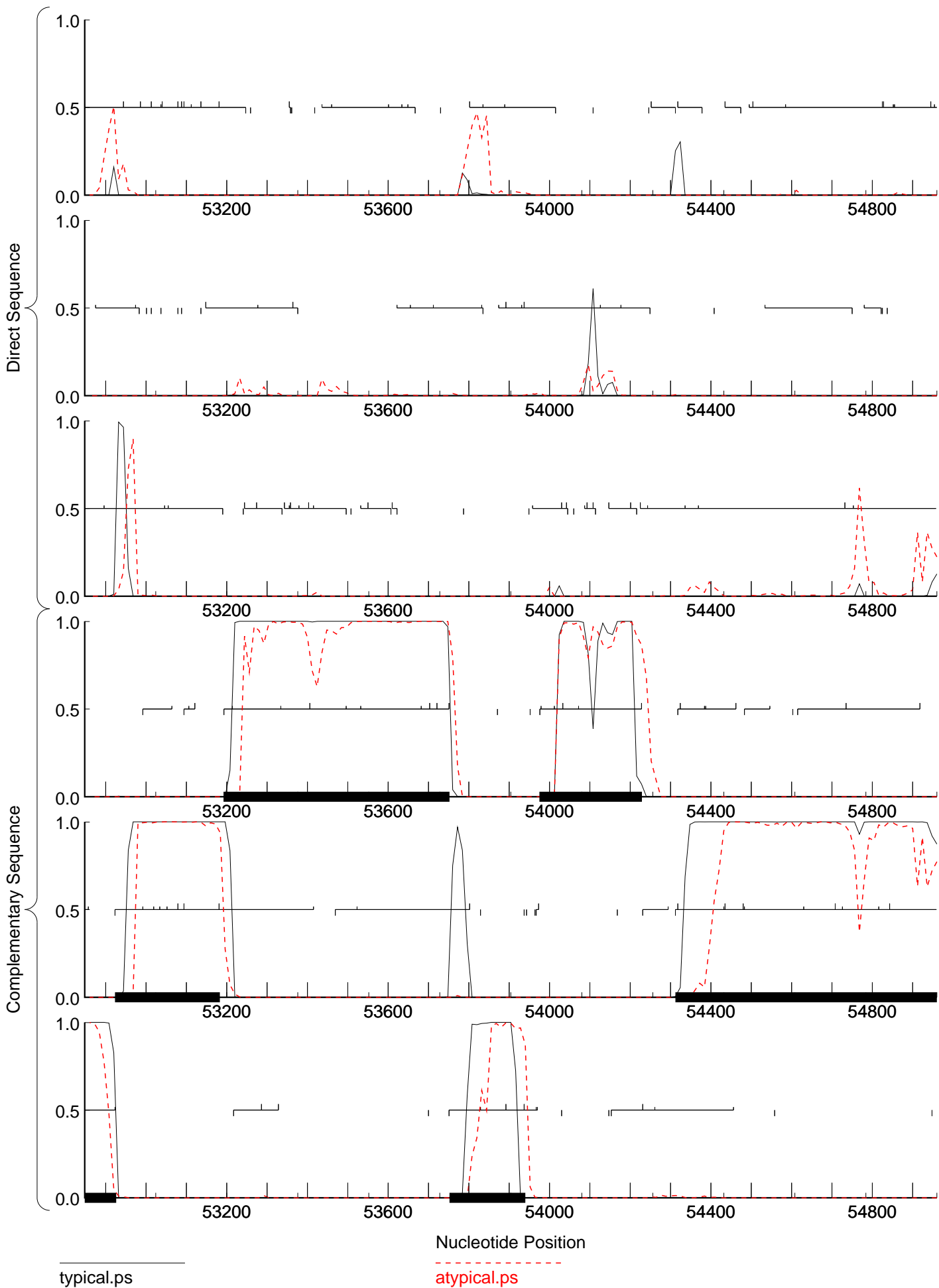


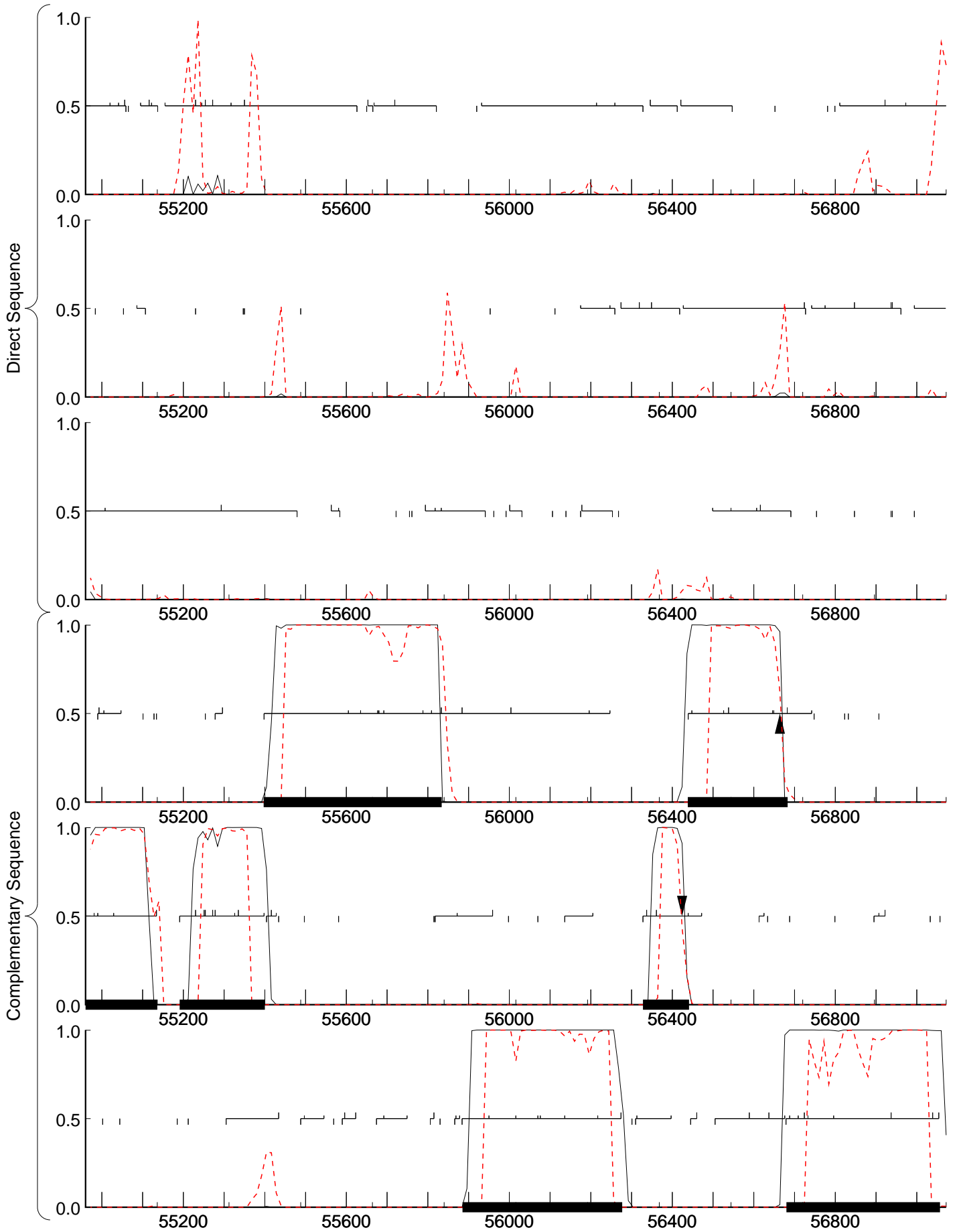
typical.ps

atypical.ps



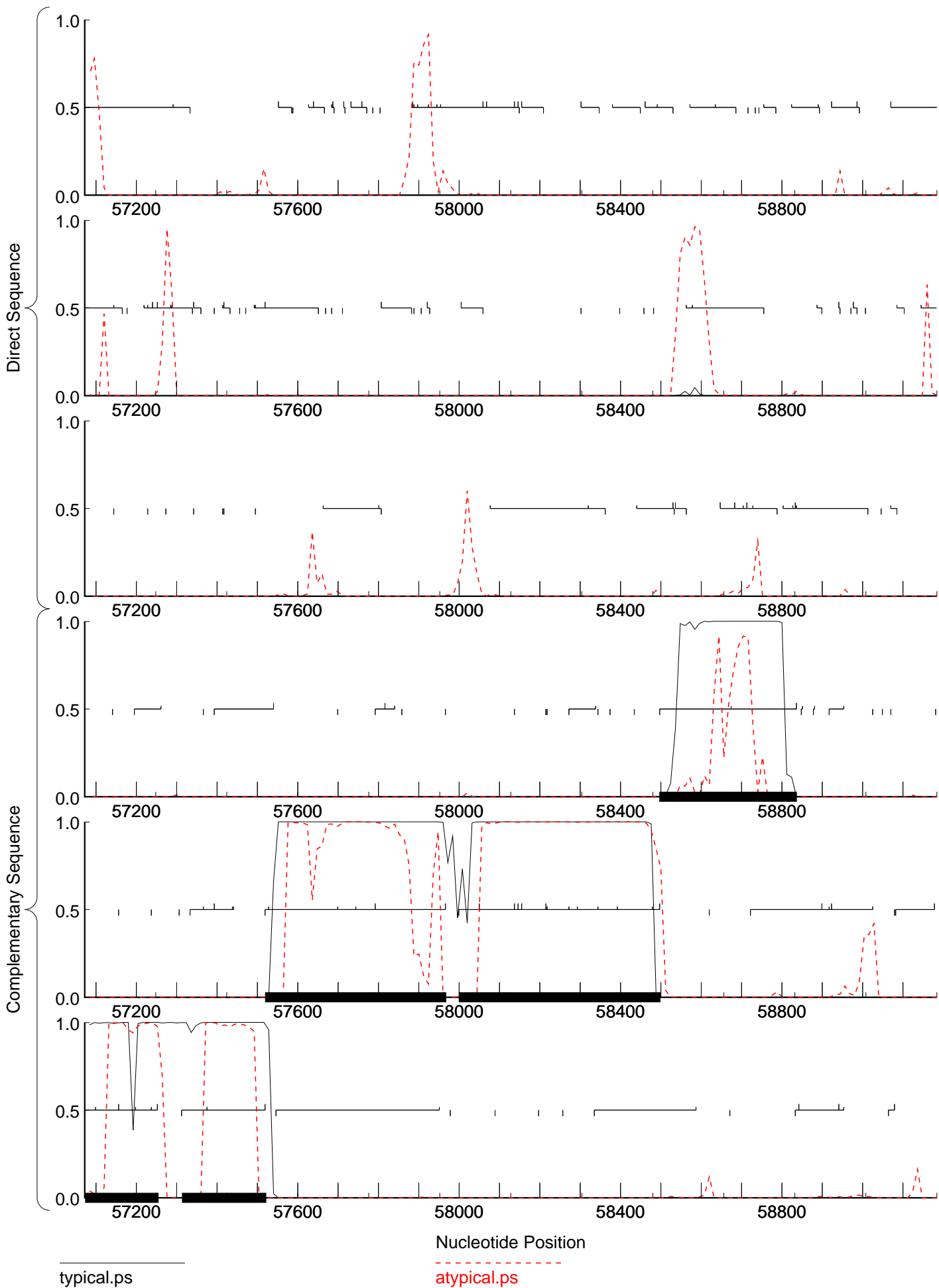






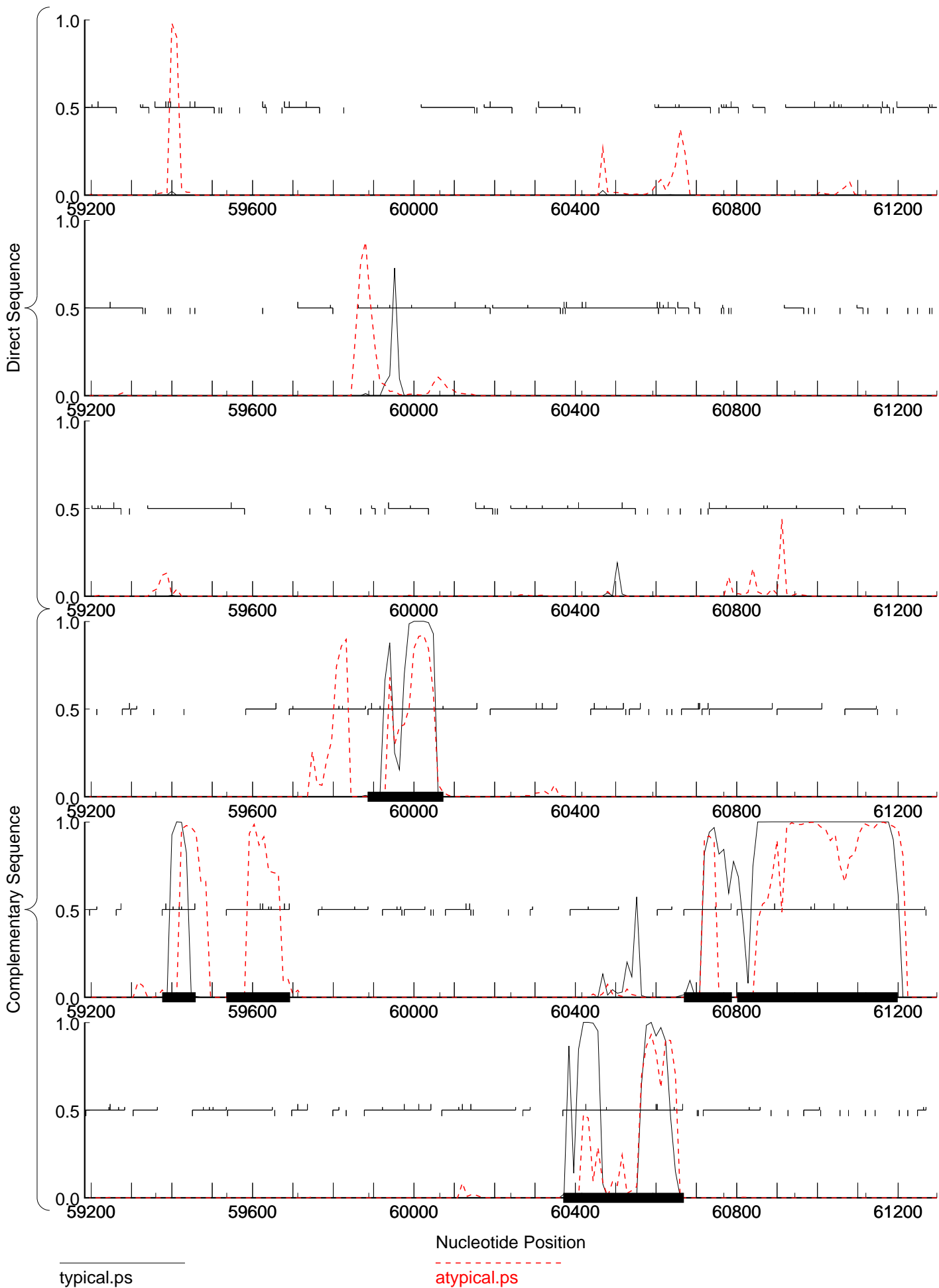
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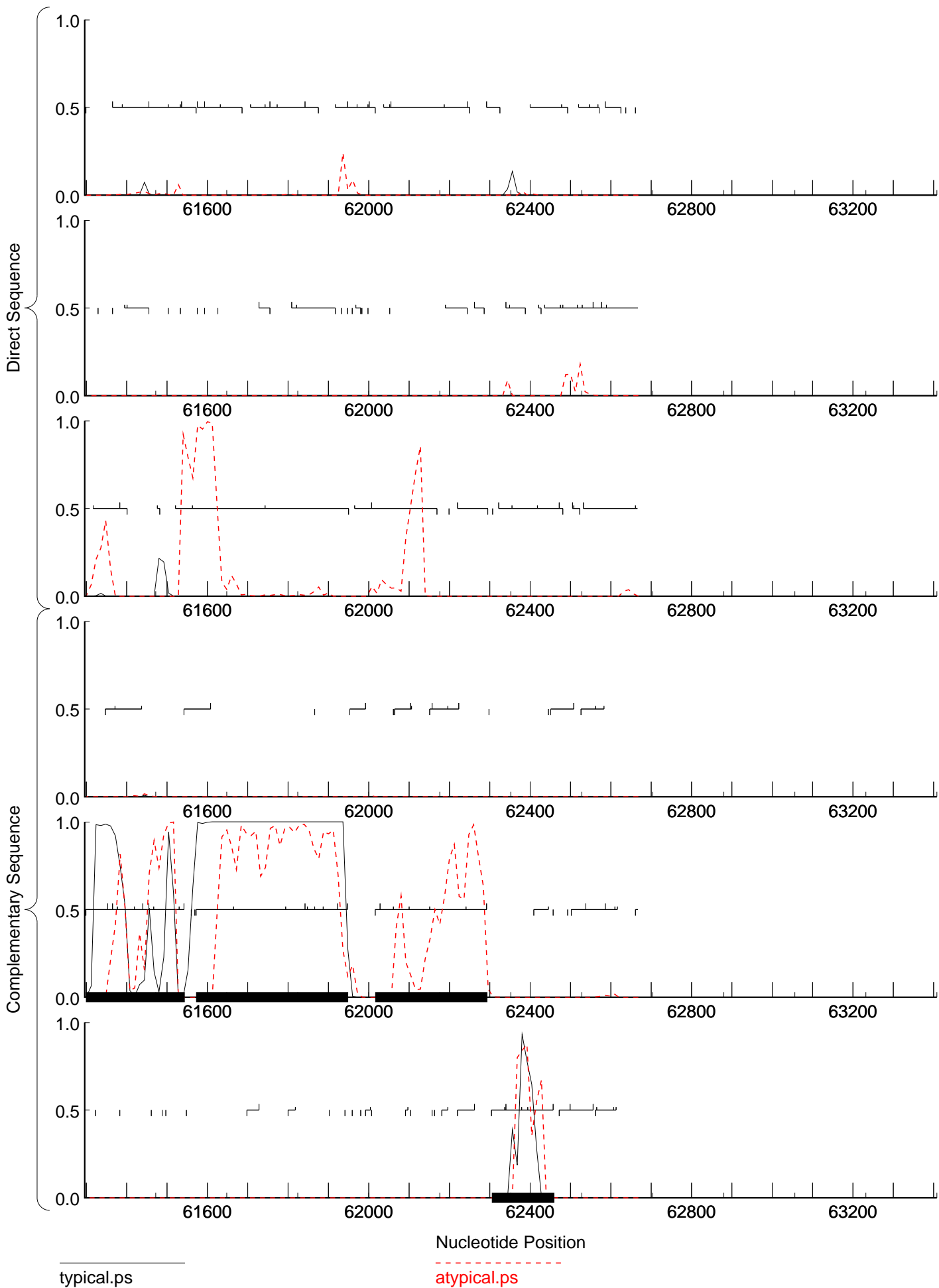
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