

GeneMark

Version 2.5p (09.08.06)

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PROGRAM INFORMATION

Sequence : Mycobacterium phage Godines complete sequence, 67277 bp, circularly permuted, Cluster B2
Analysis Date : 10/23/17 at 7:25:19
Pages : 33
Sequence Length : 67277 bp
GC Content : 69.03%

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,

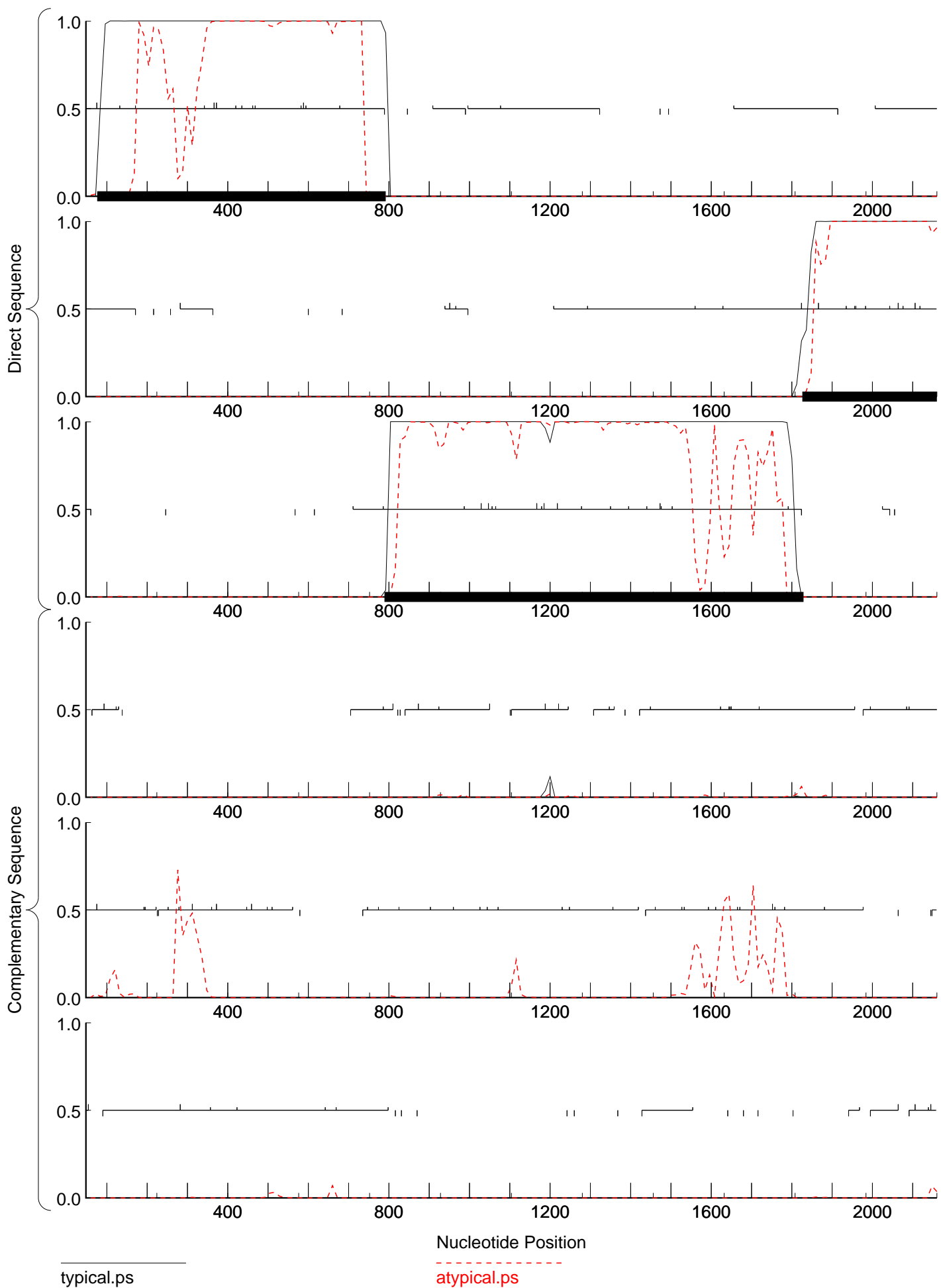
MATRIX INFORMATION

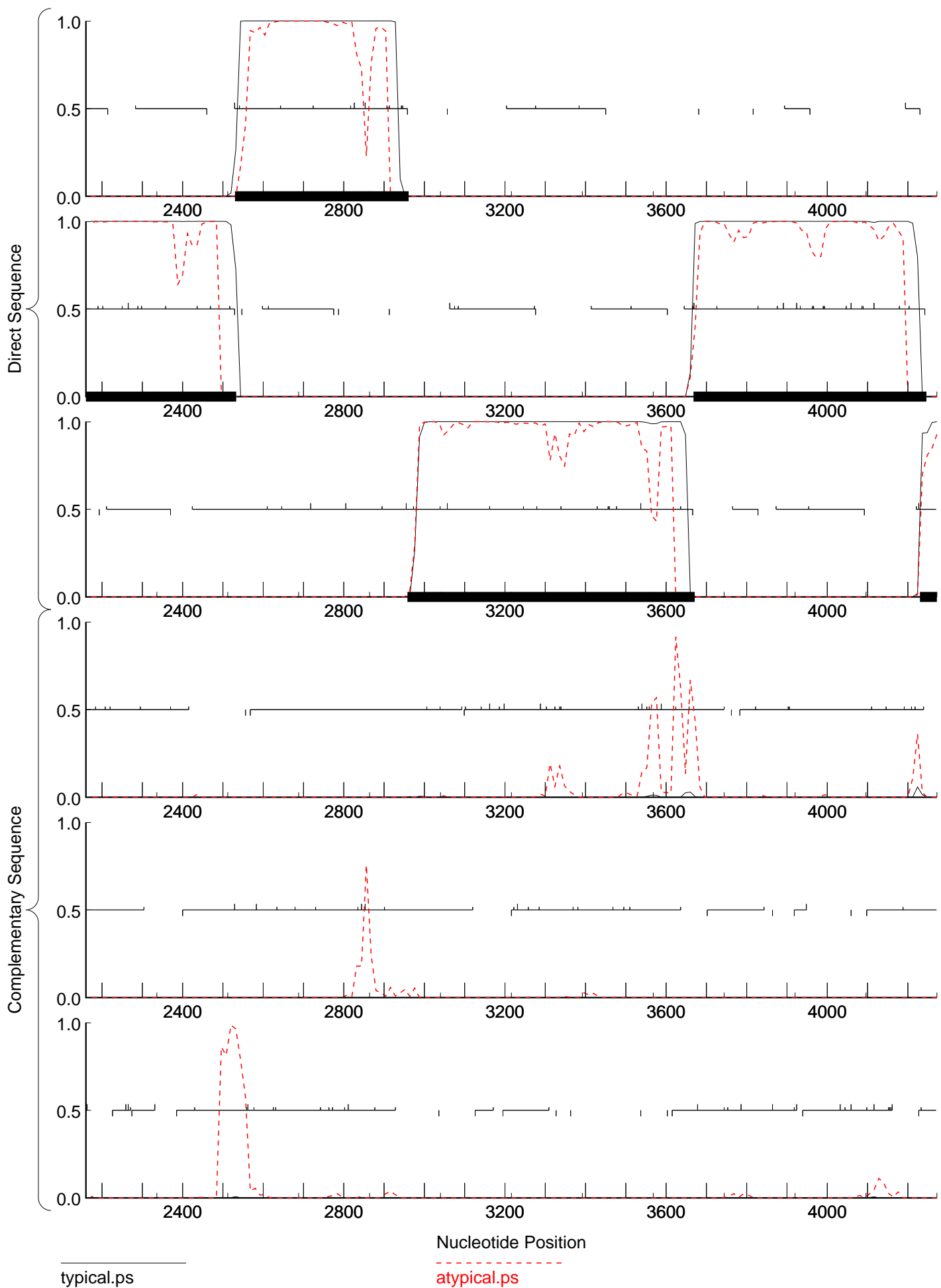
Matrix : -
Author : -
Order : 4

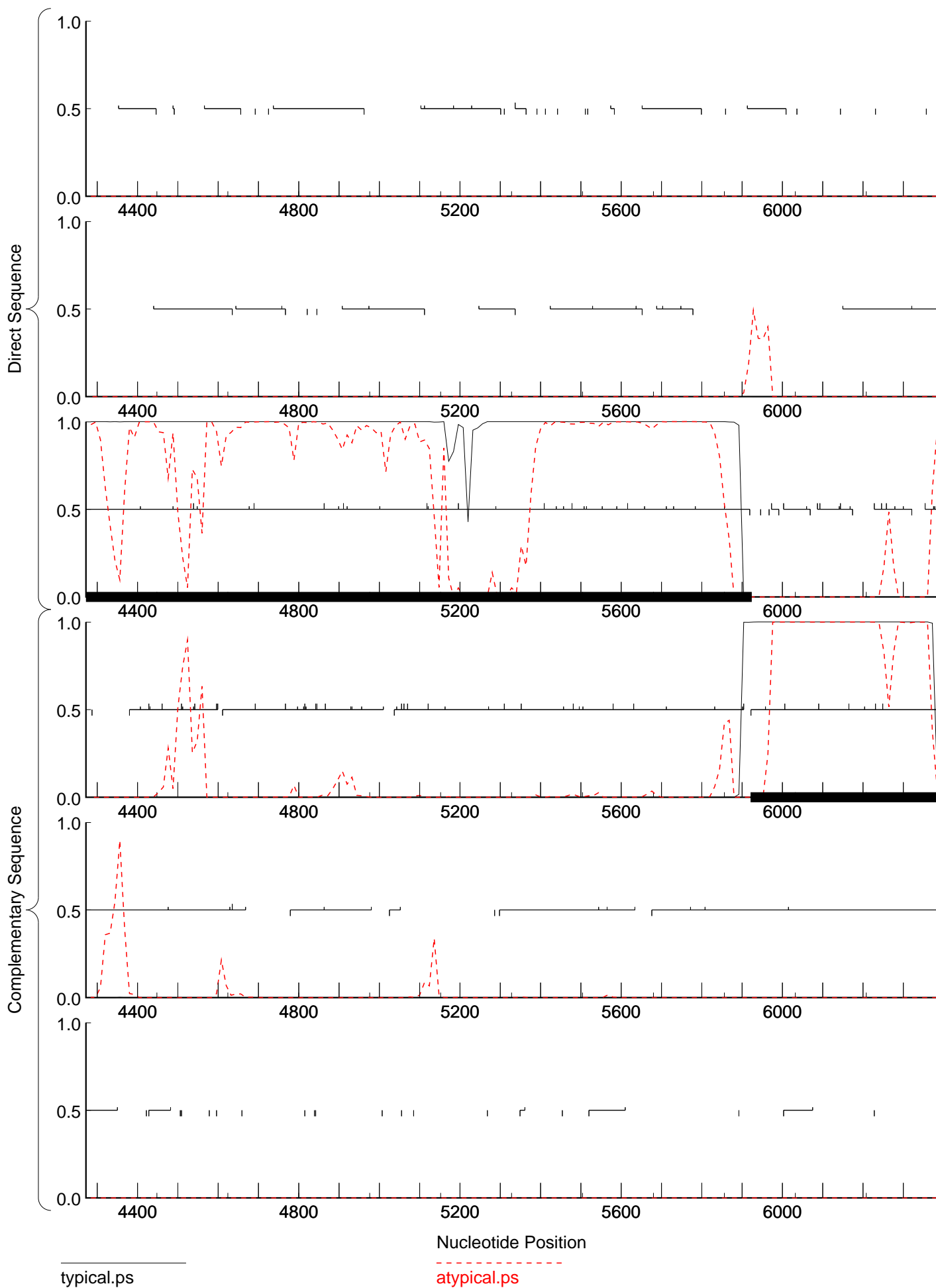
Send questions / comments to:
Dr. M. Borodovsky
Georgia Institute of Technology
School of Biology
Atlanta, GA 30332-0230

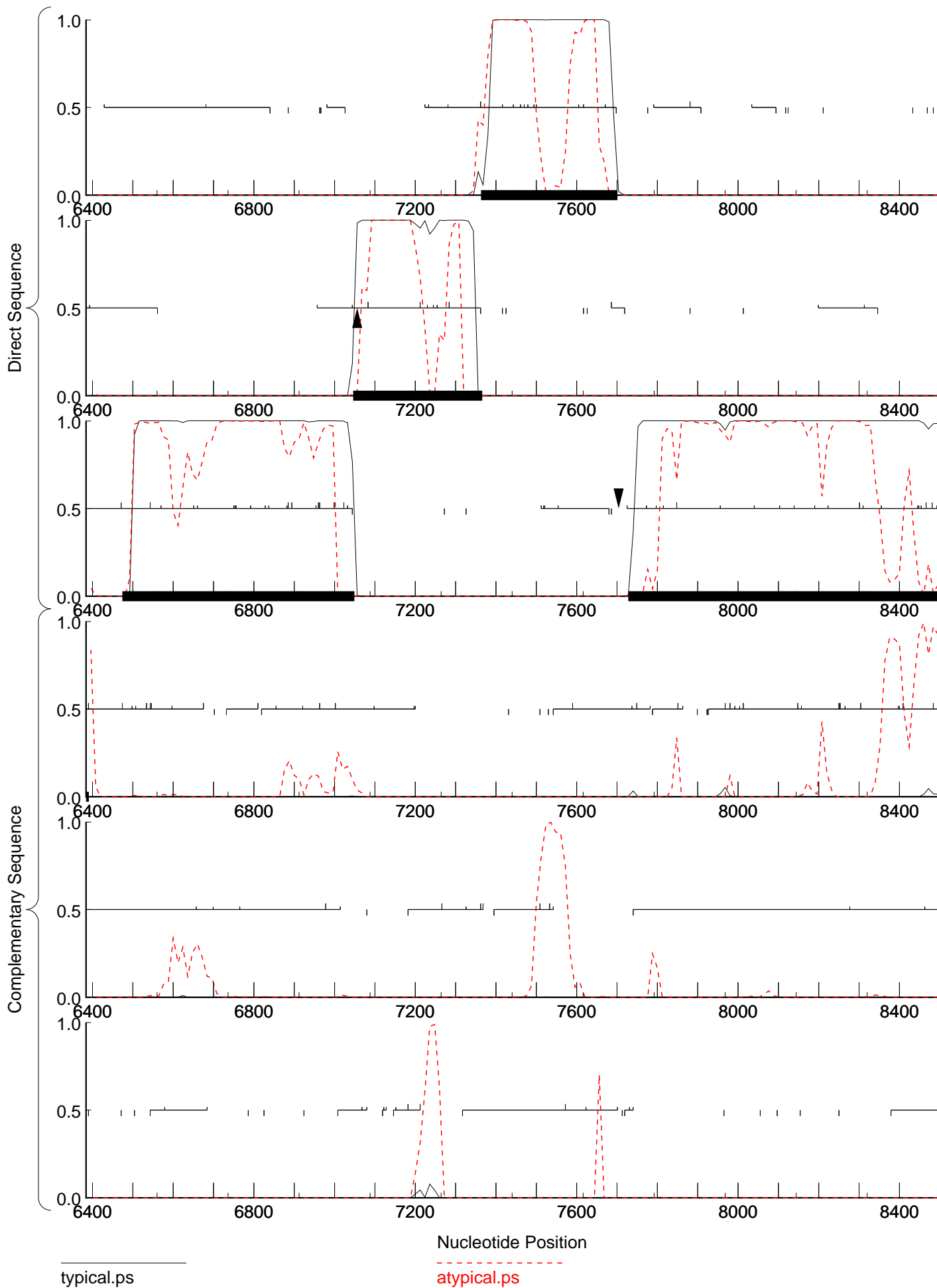
Matrix notes & comments

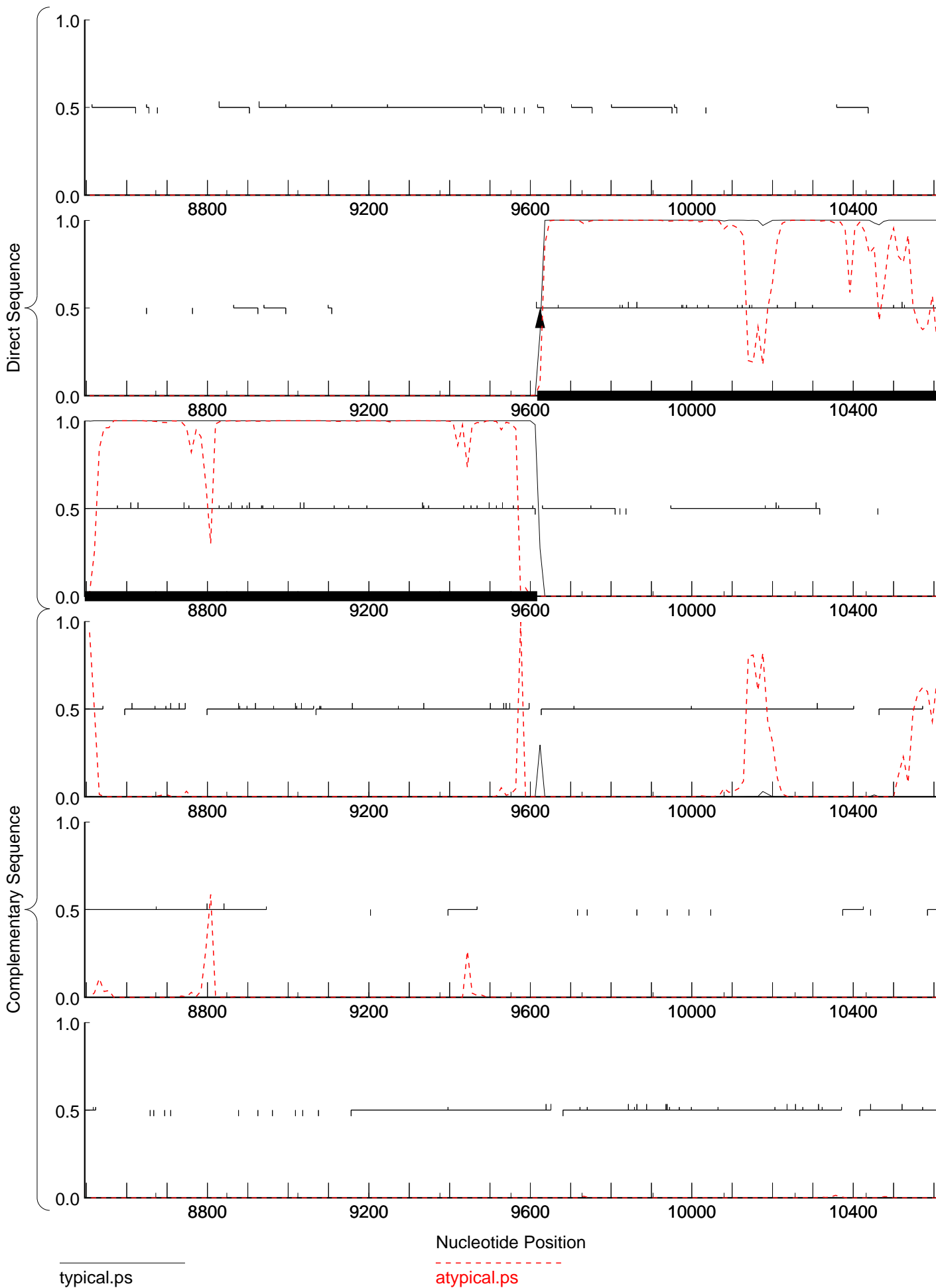
Training set derived by GeneMarkS, 4.30 October 2014
Mon Oct 23 07:25:19 2017

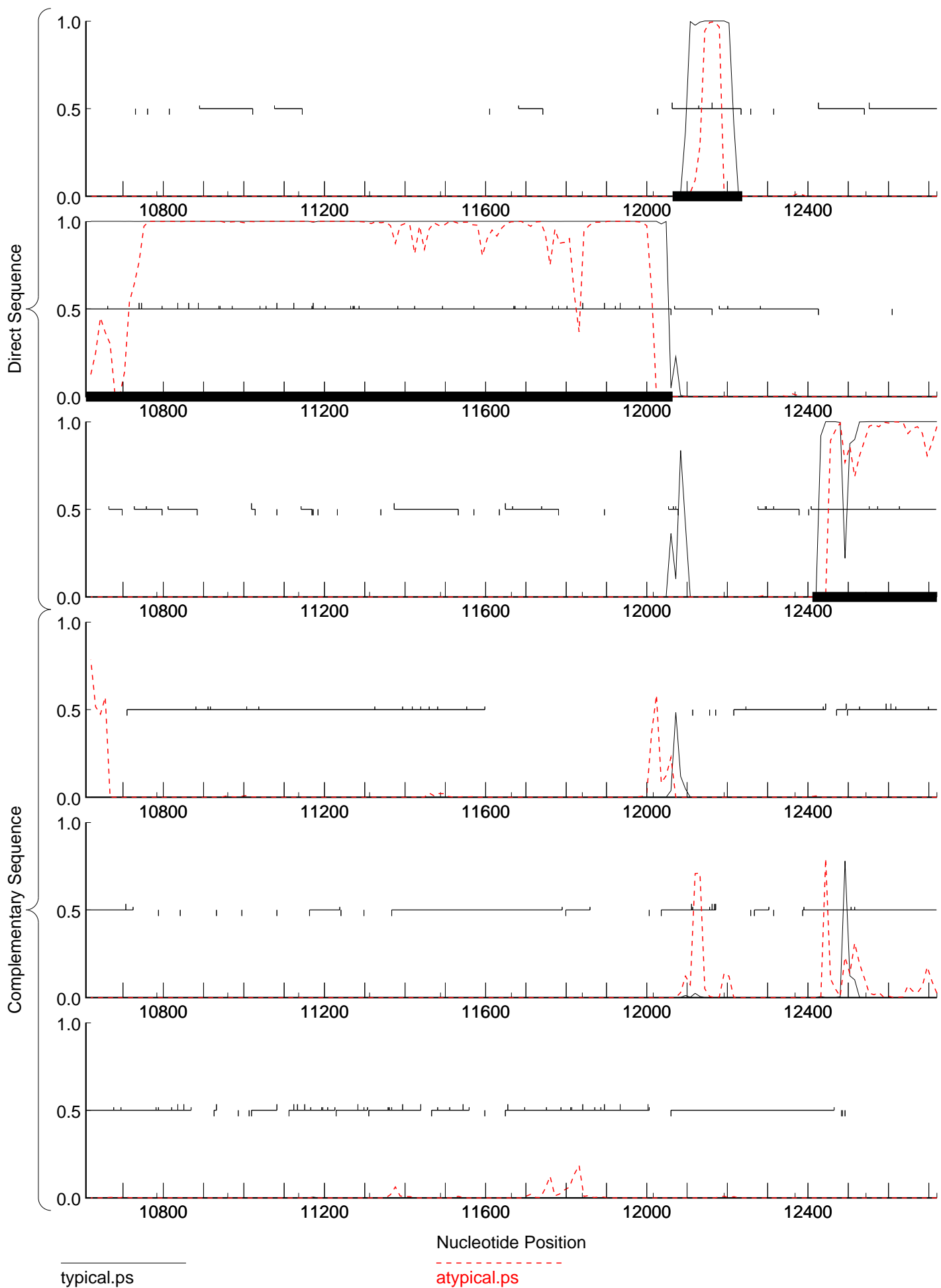


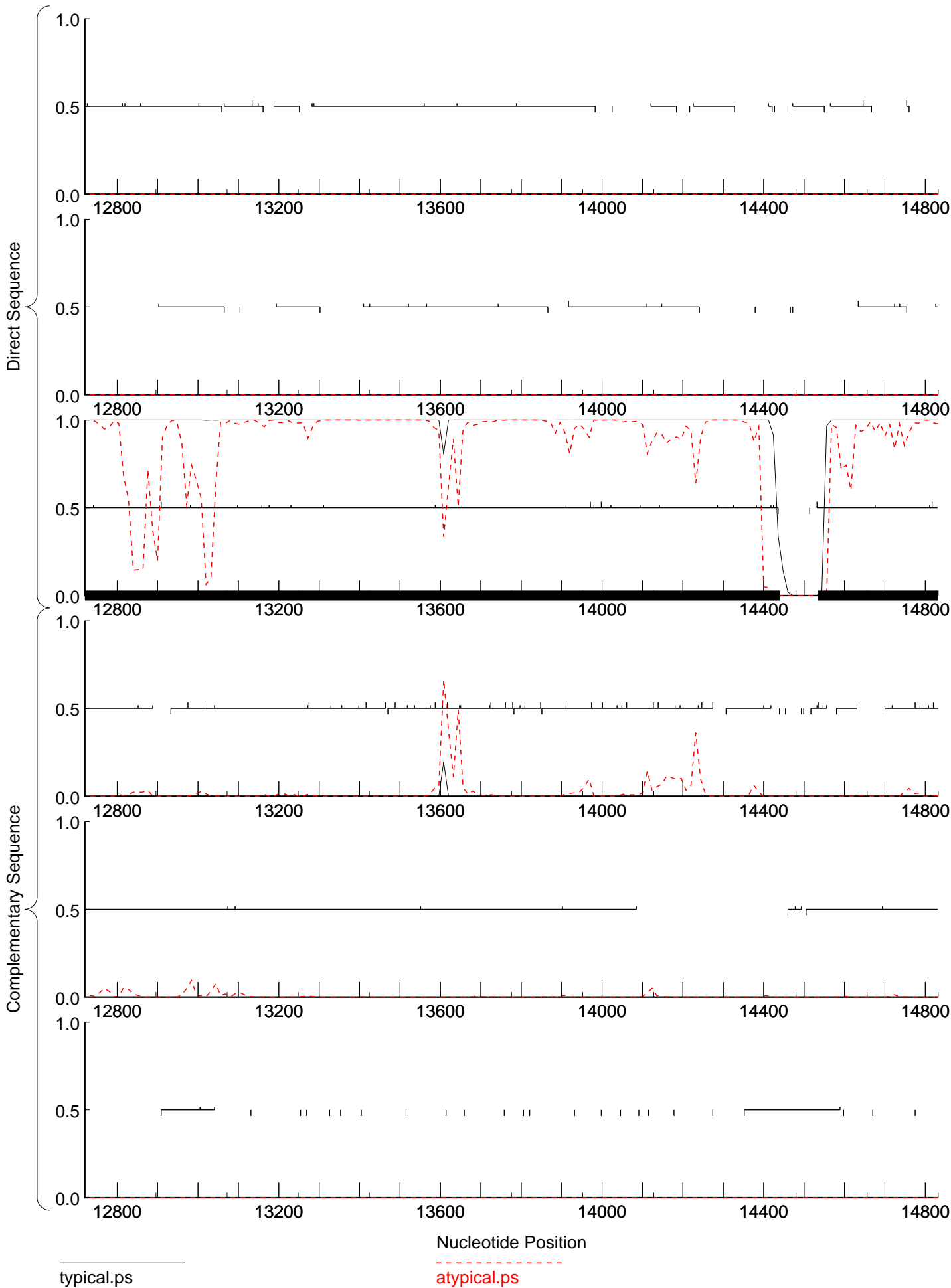


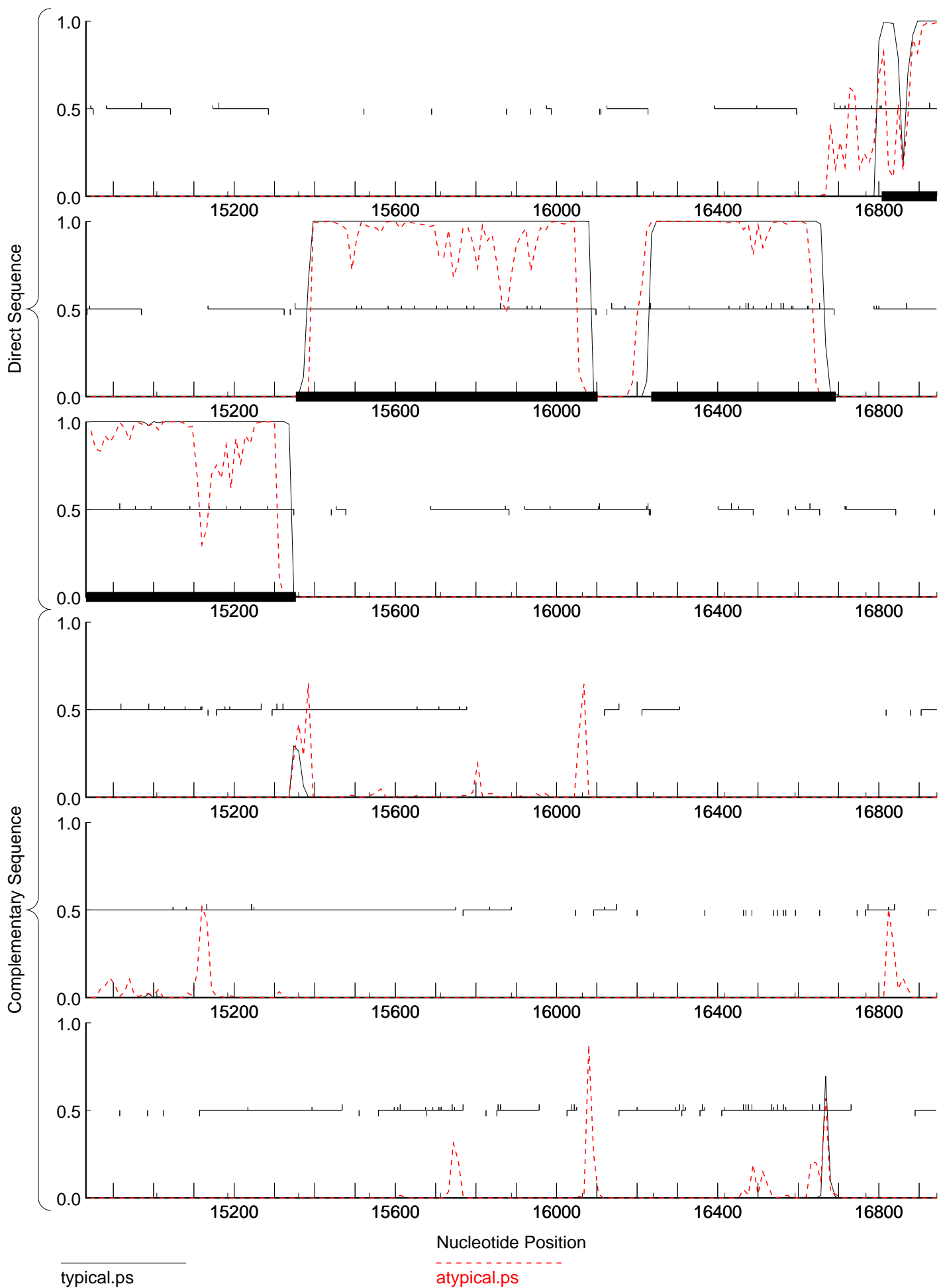


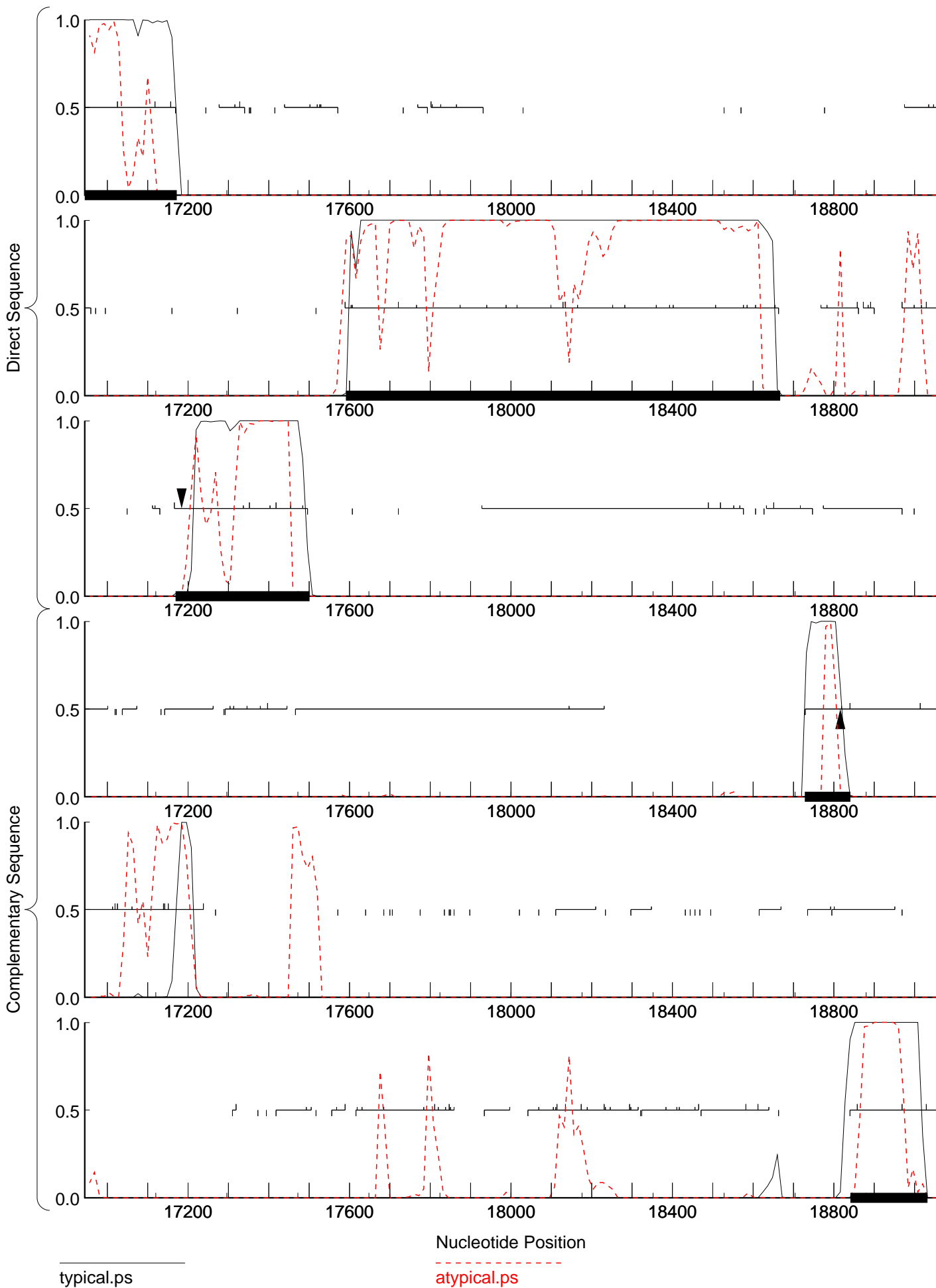






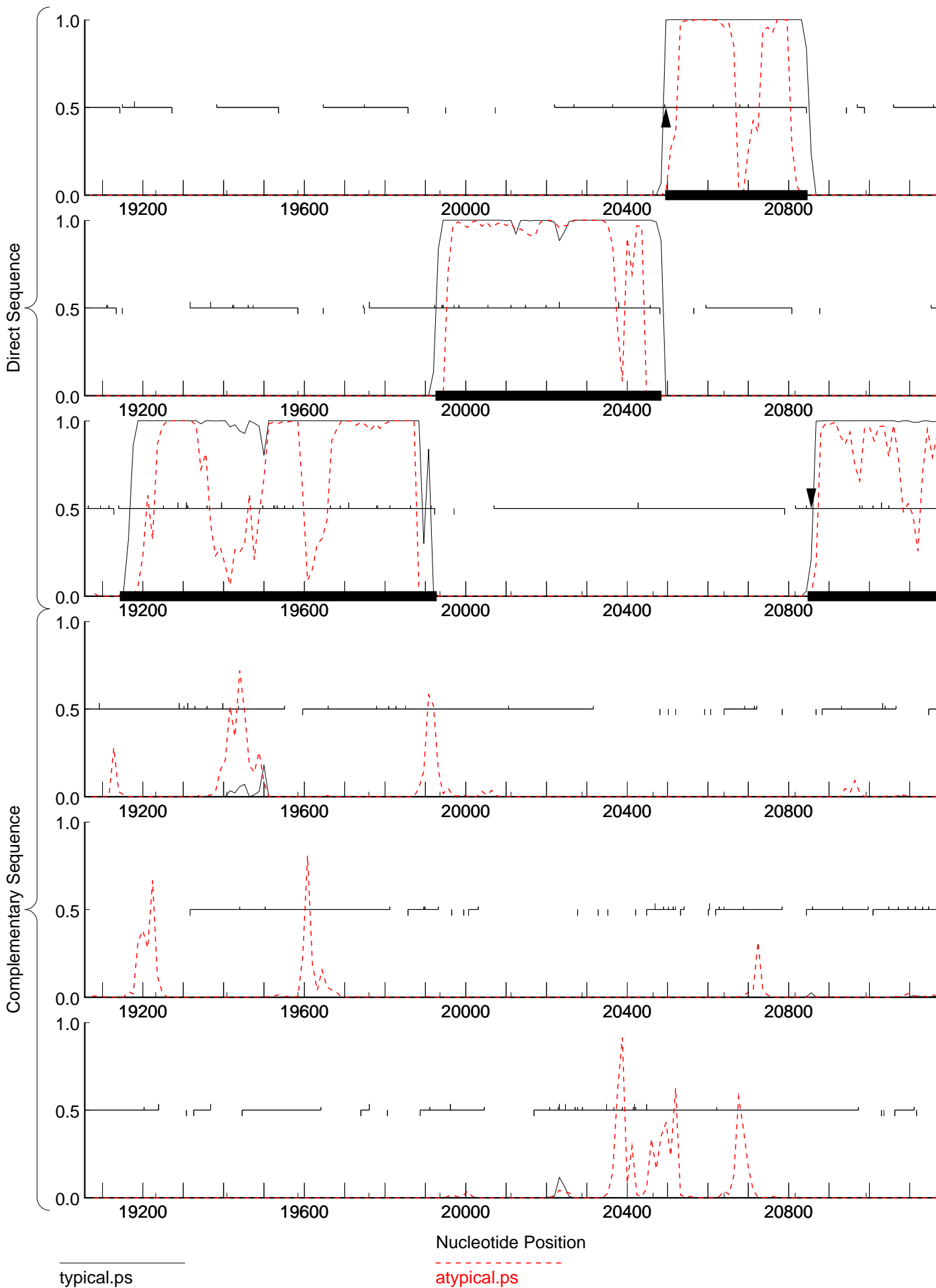


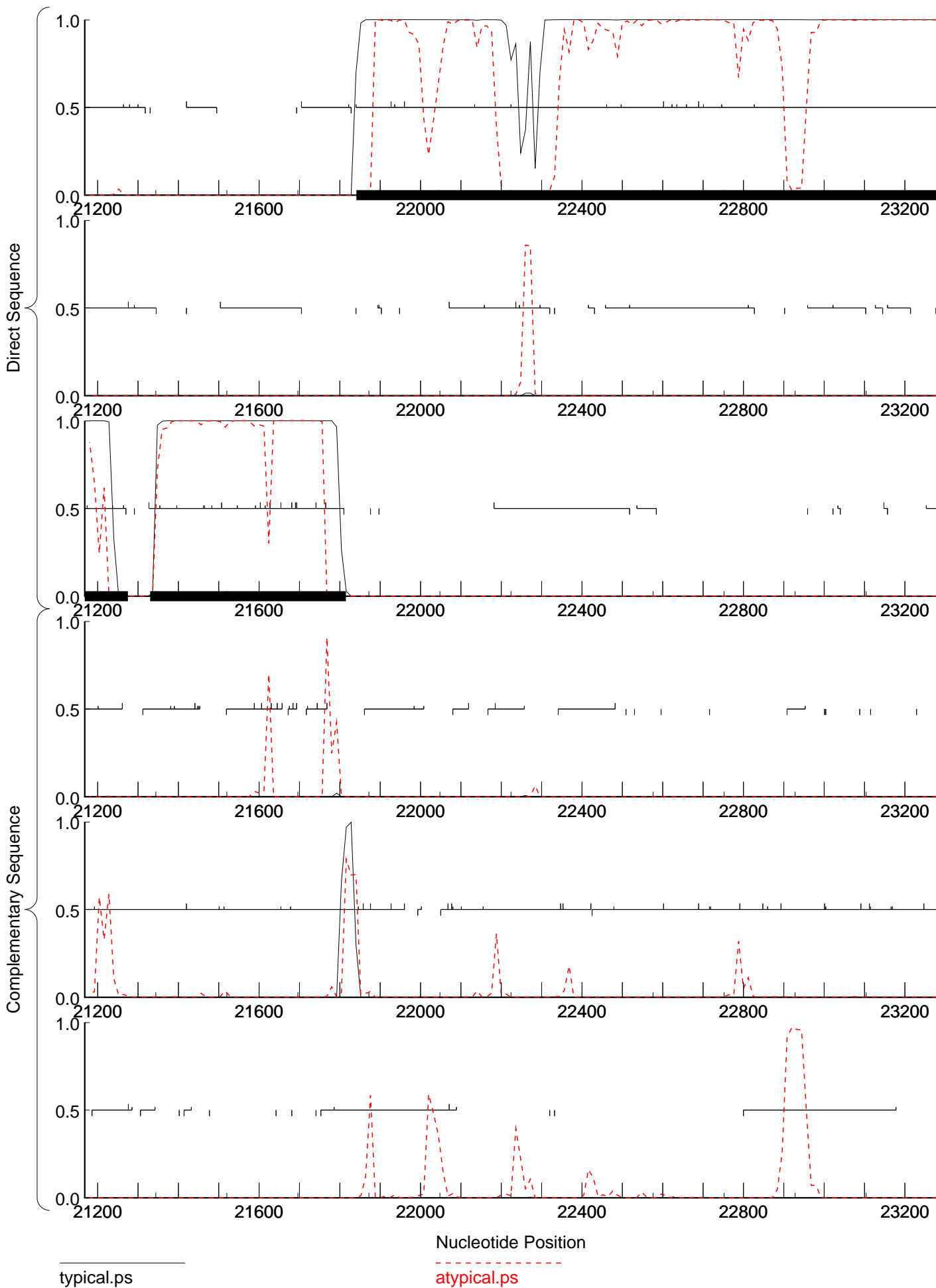


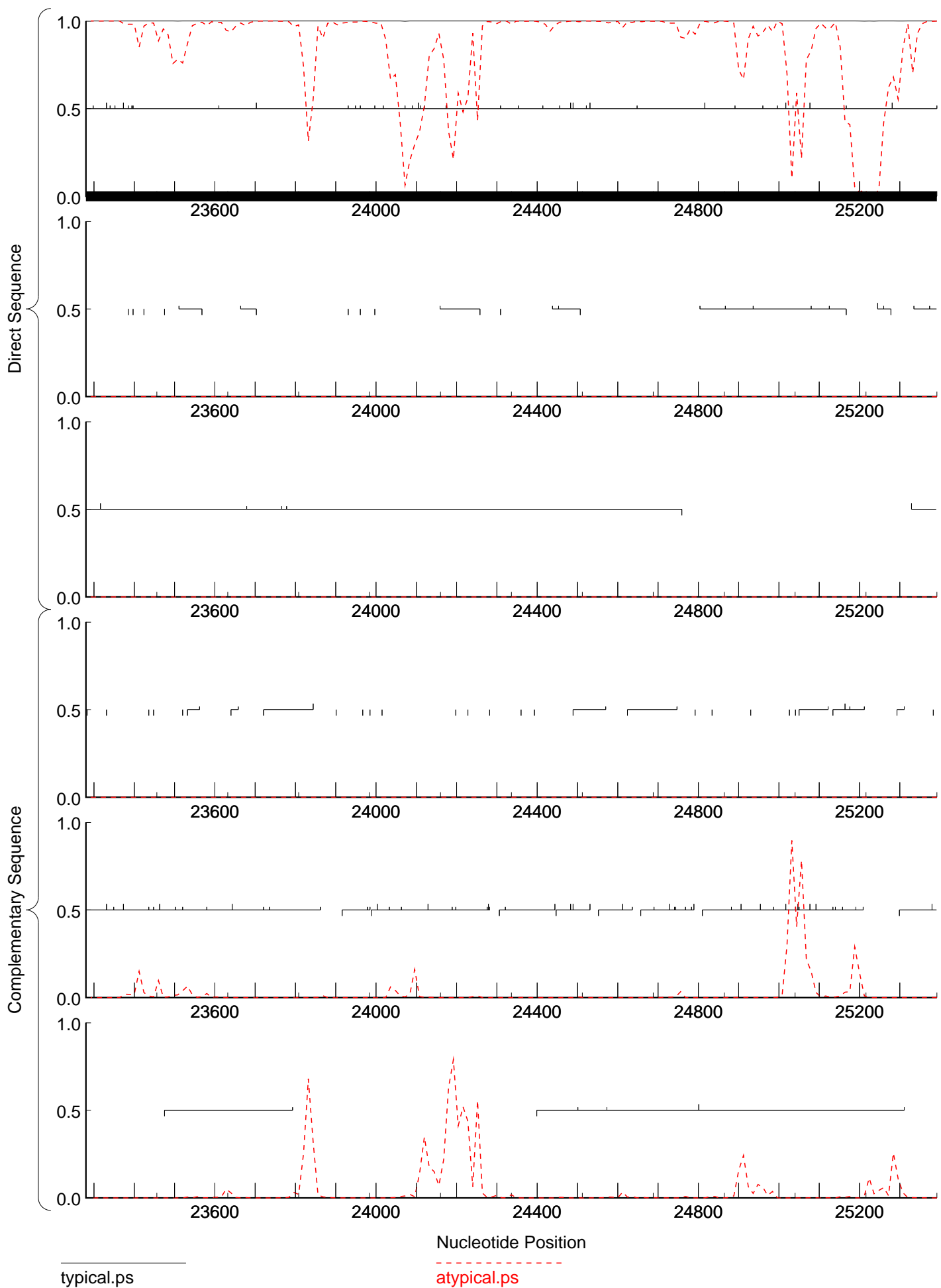


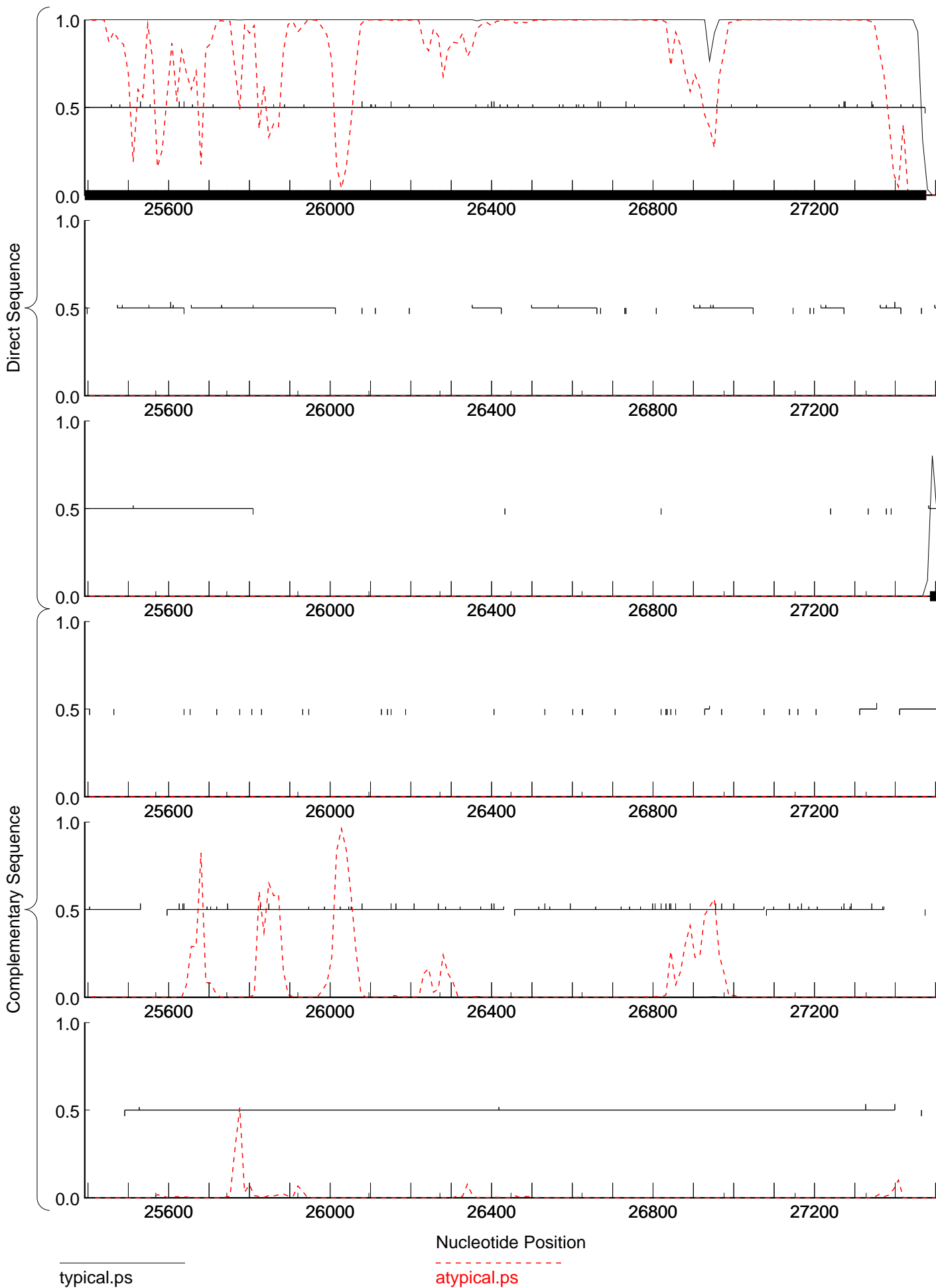
typical.ps

atypical.ps



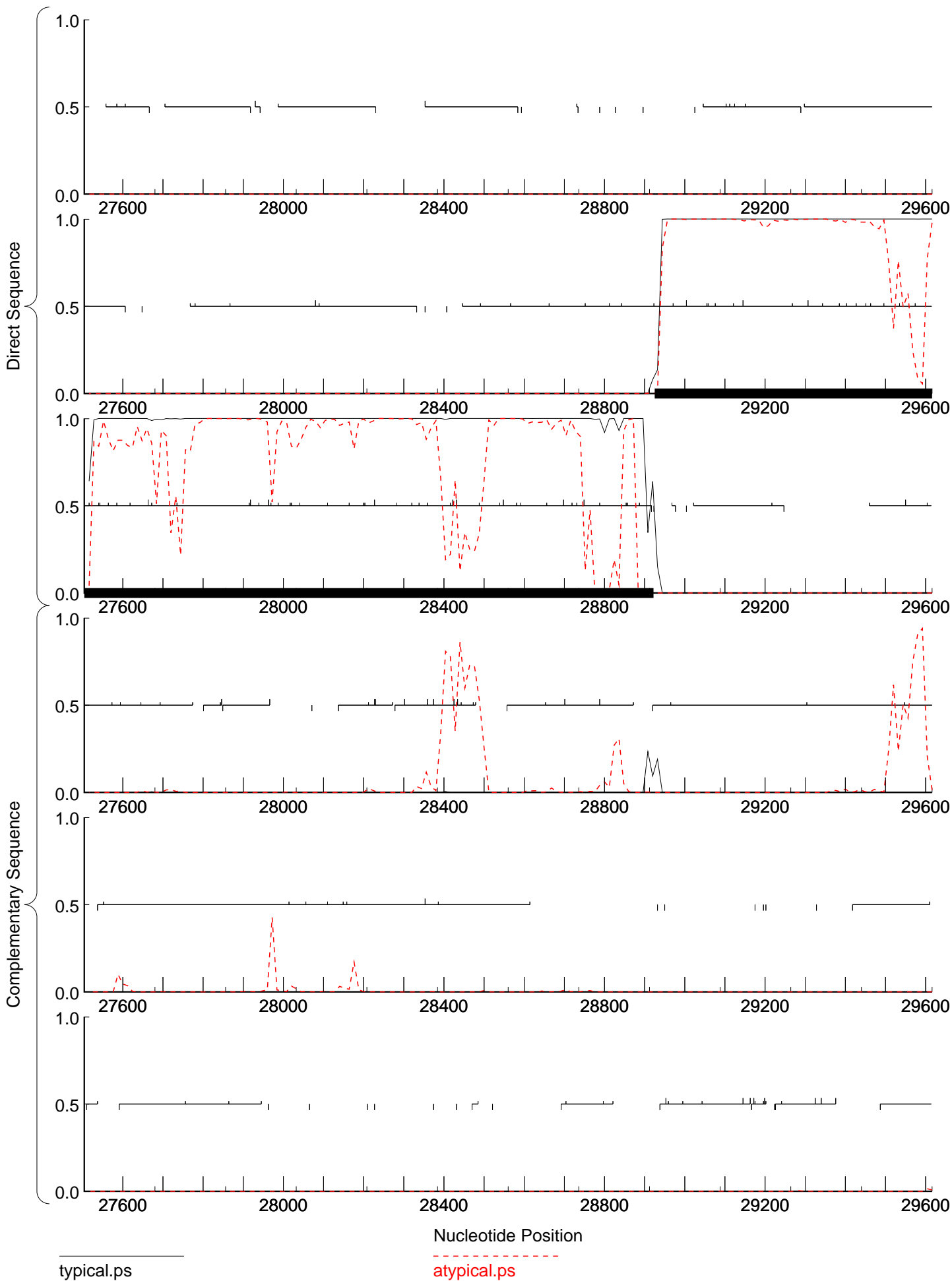


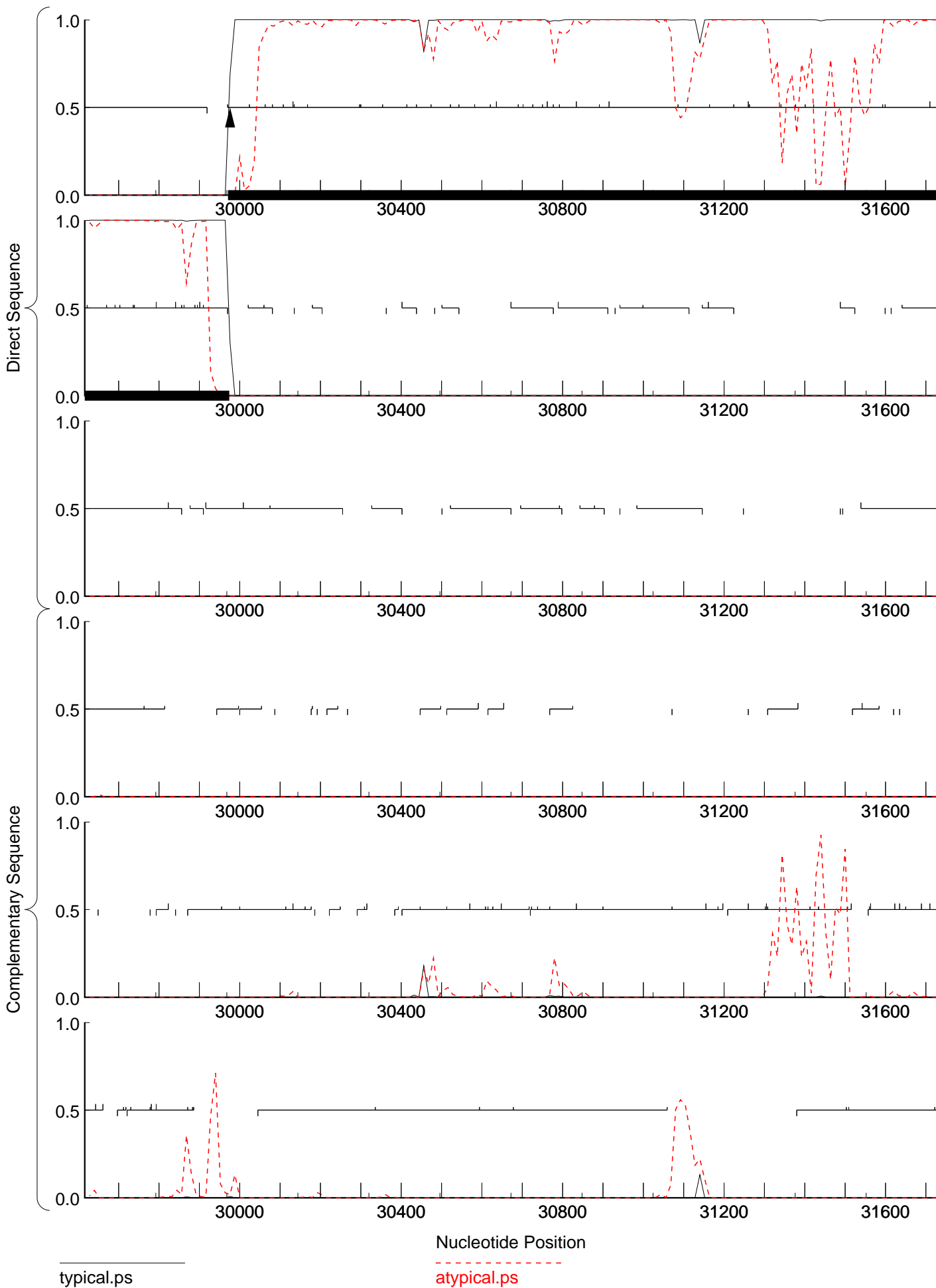


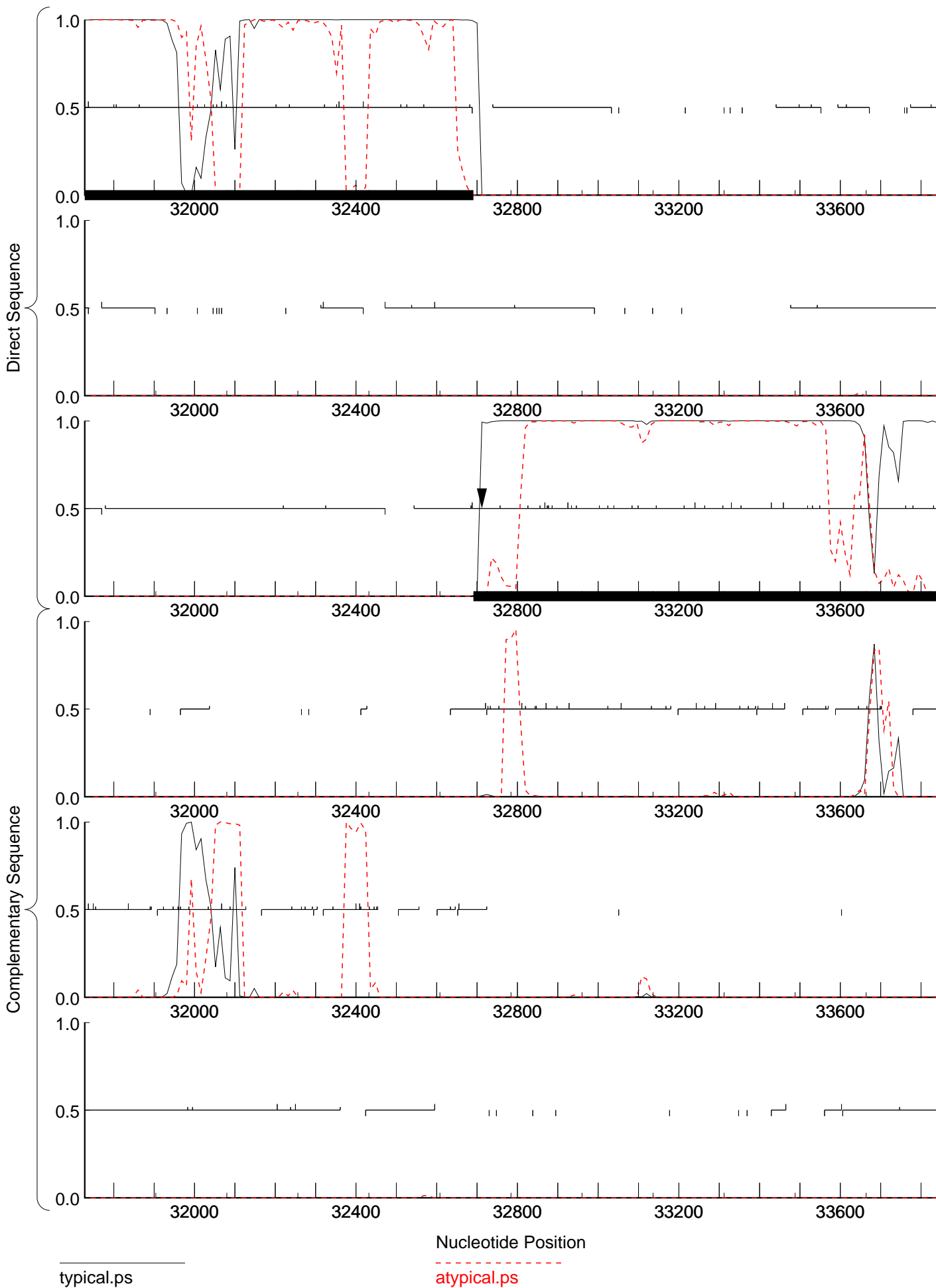


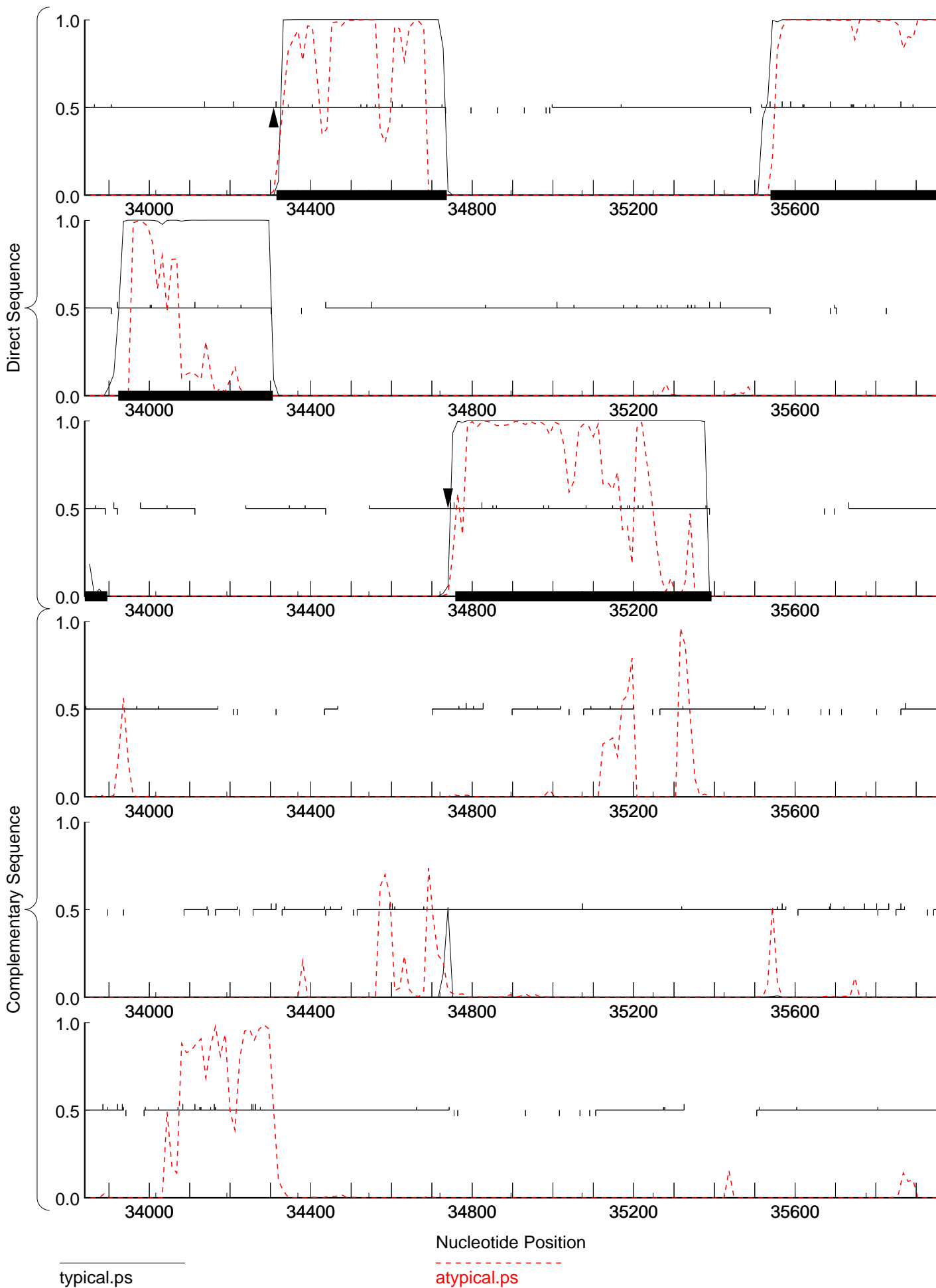
typical.ps

atypical.ps



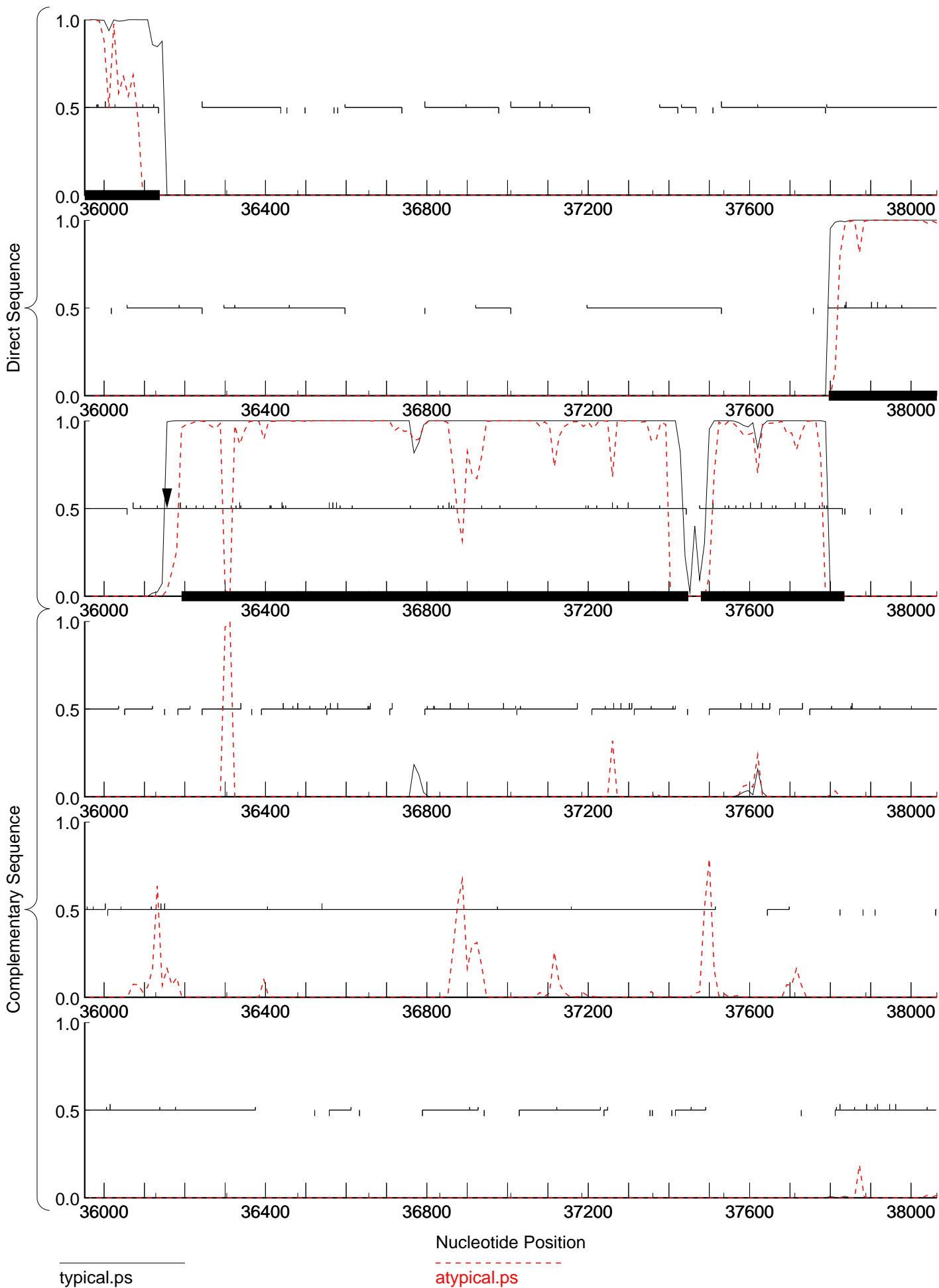


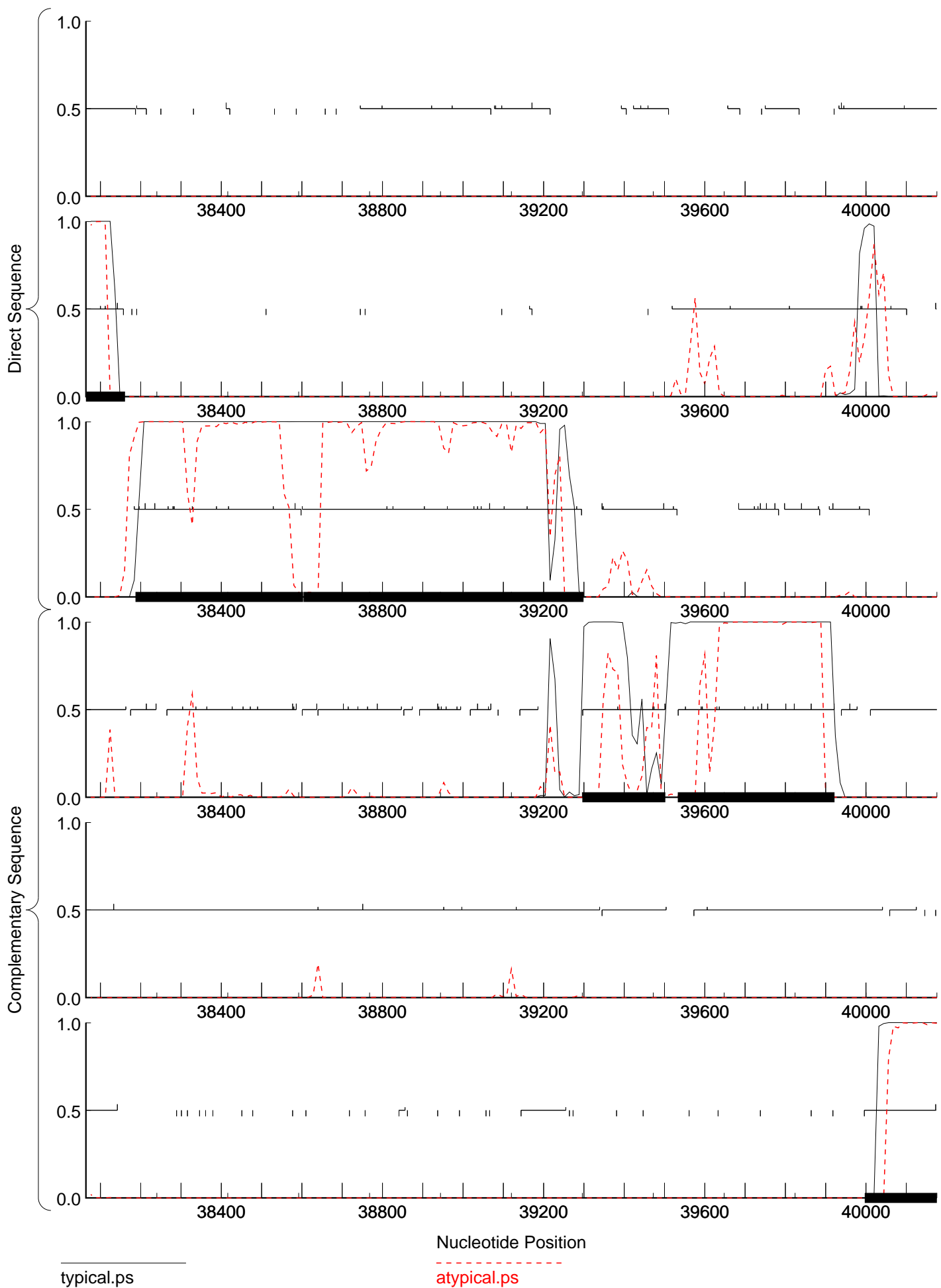




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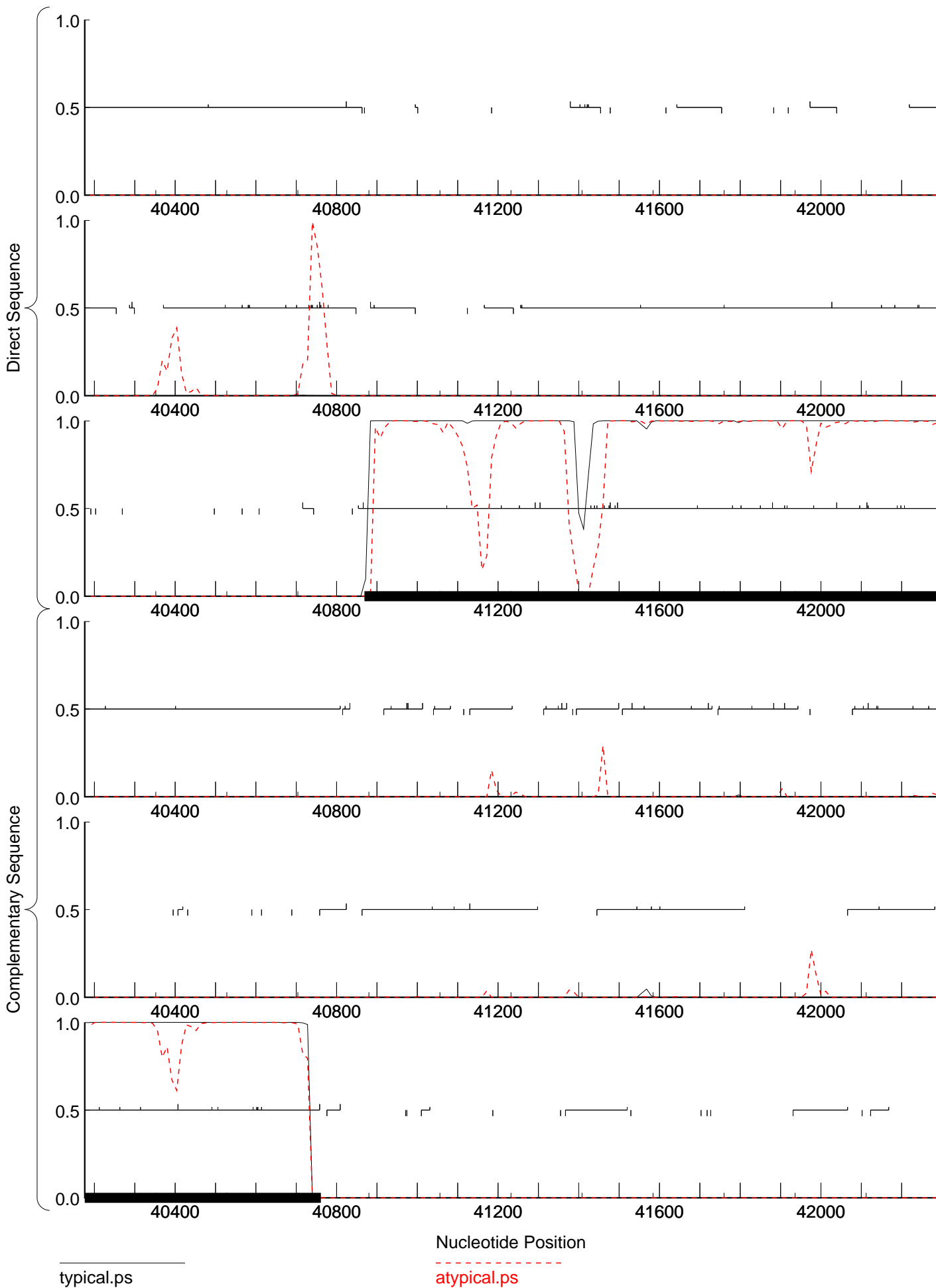
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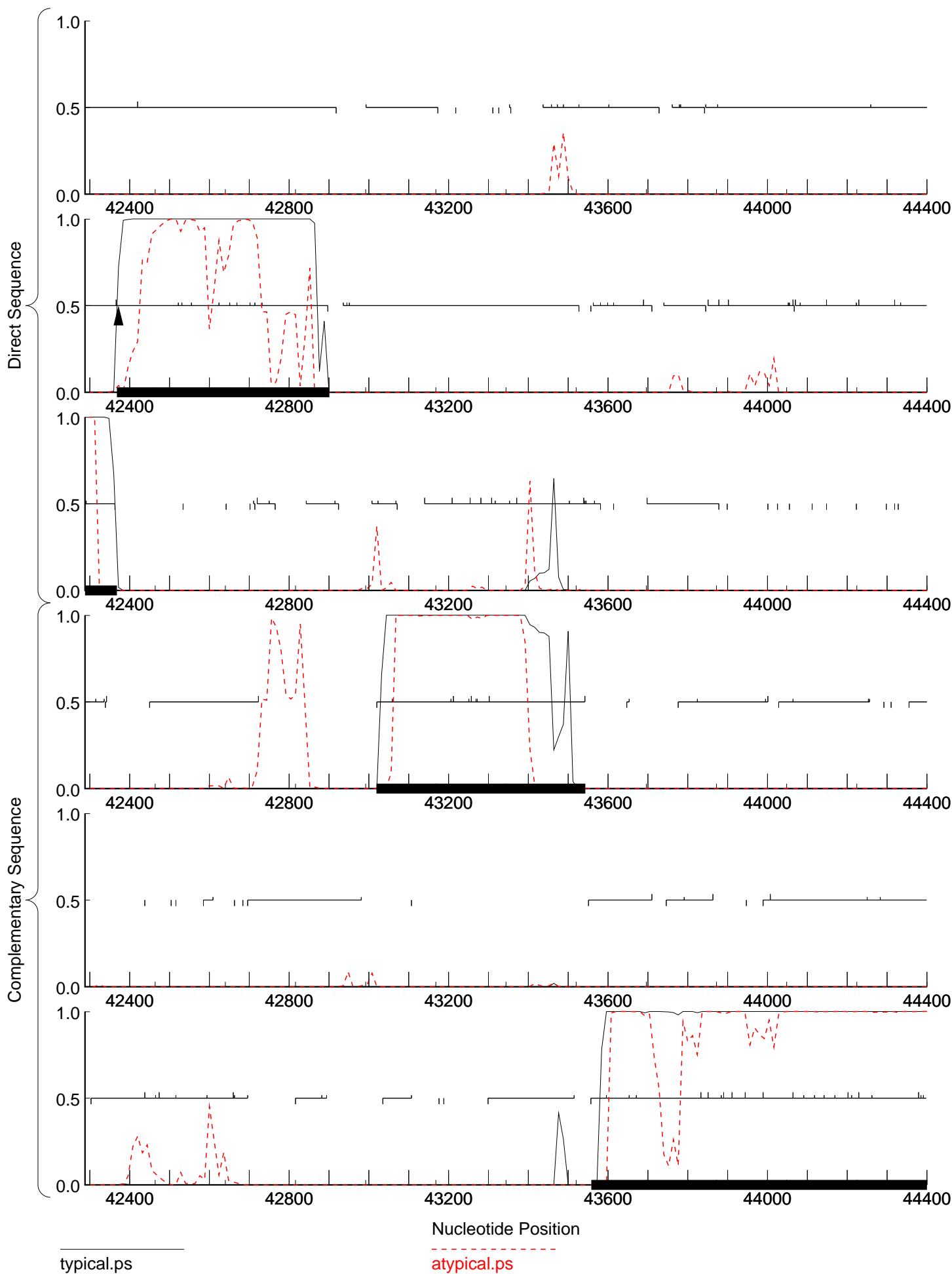
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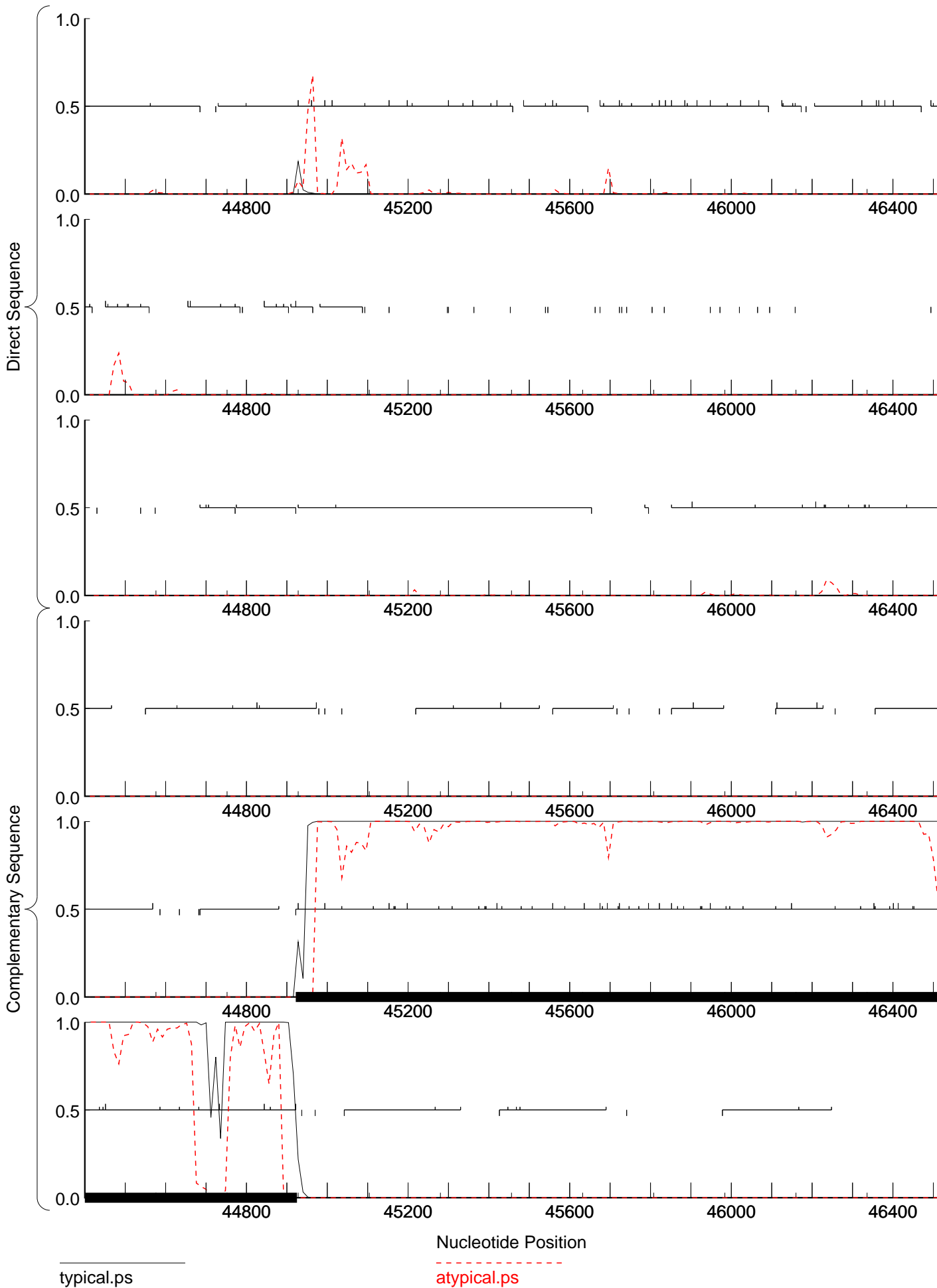
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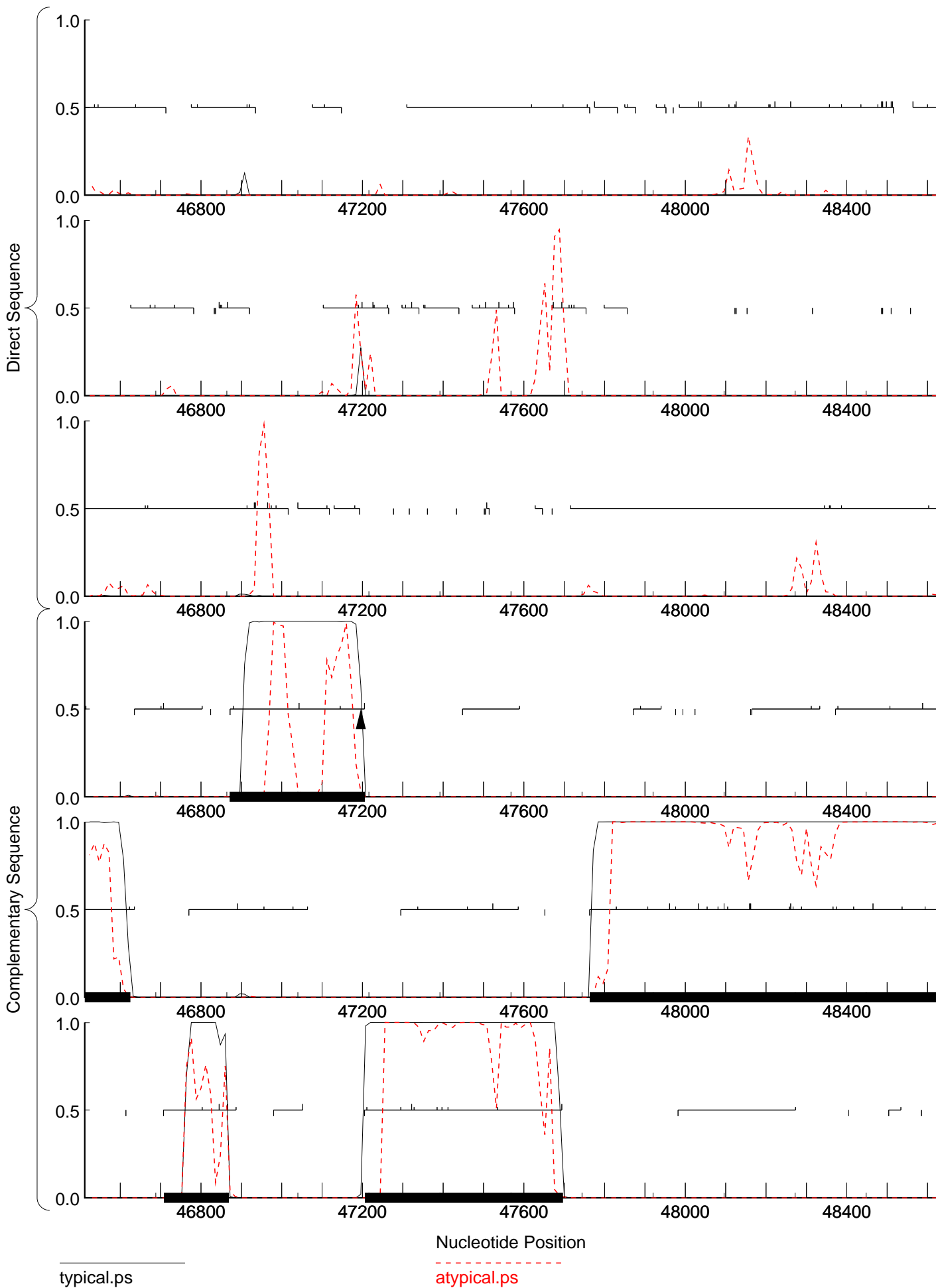


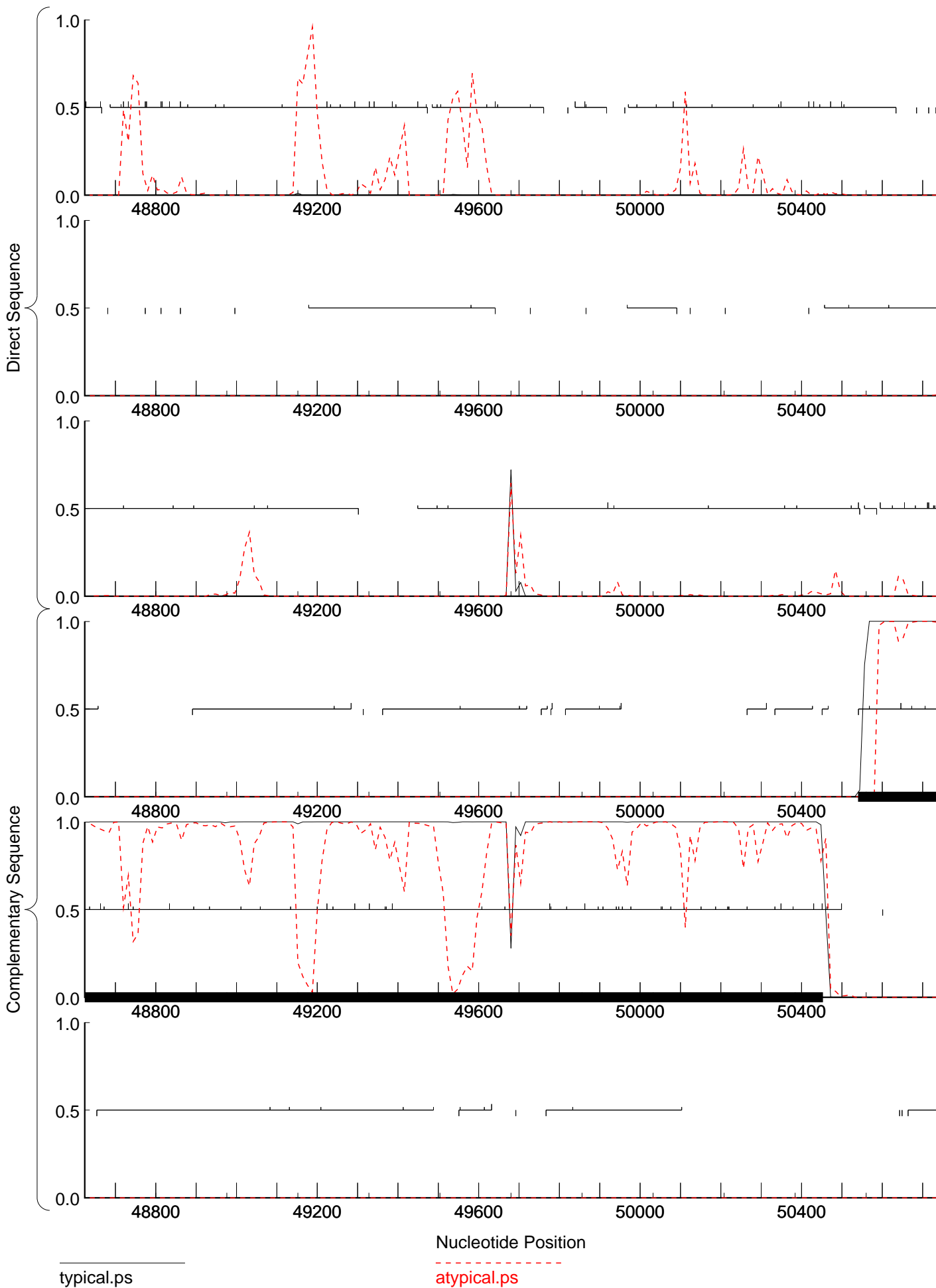
typical.ps

atypical.ps



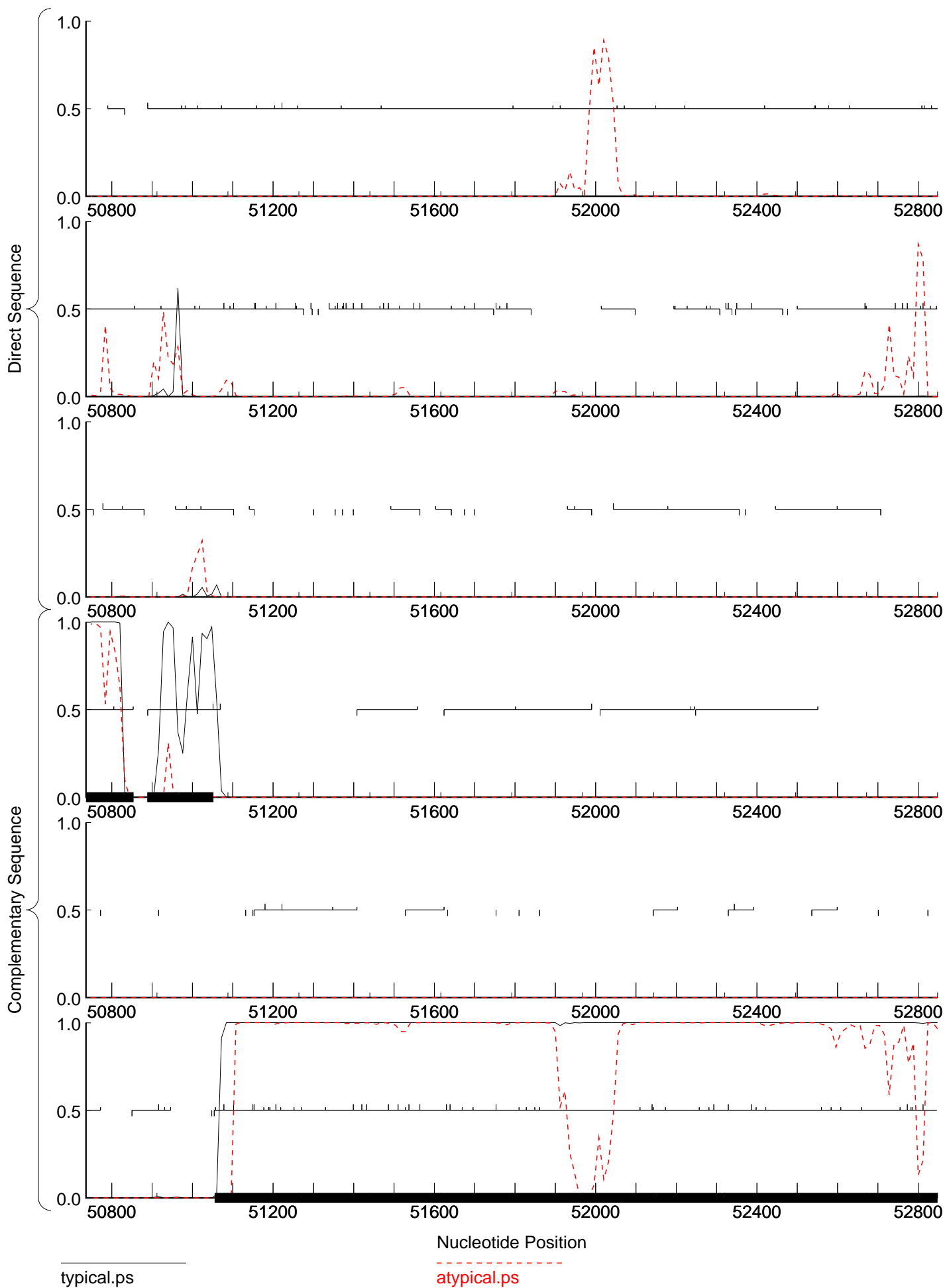


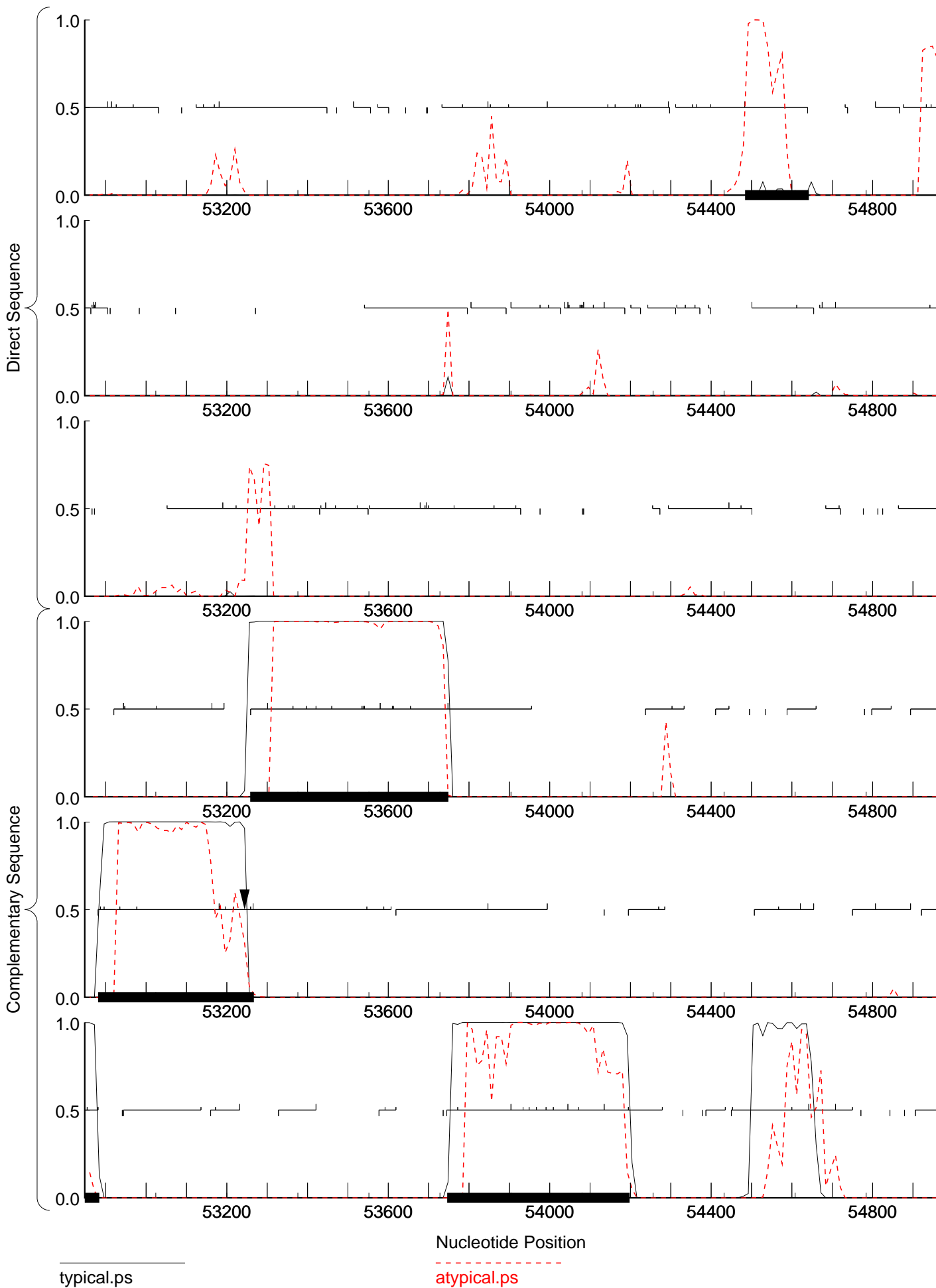


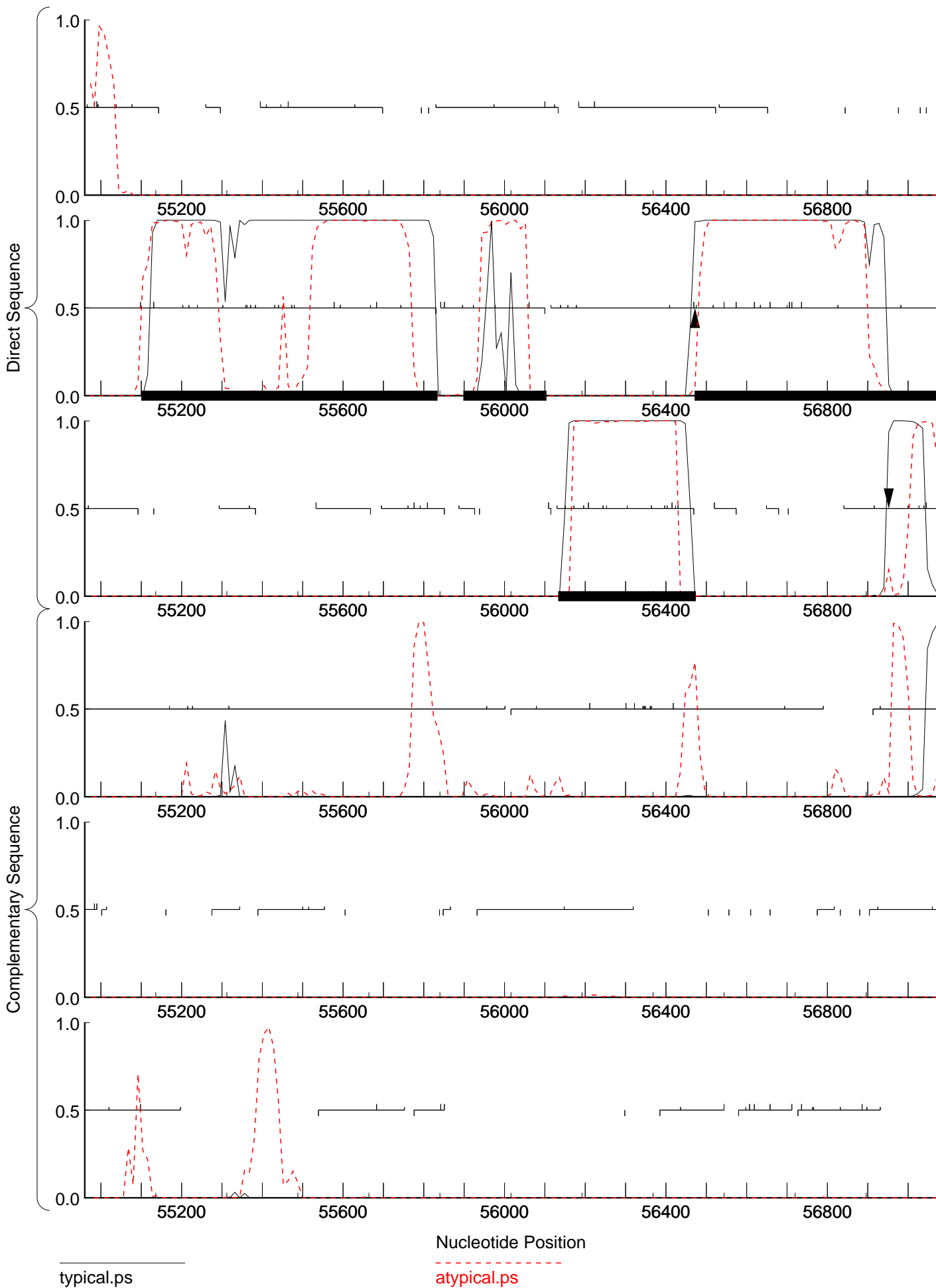


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atypical.ps

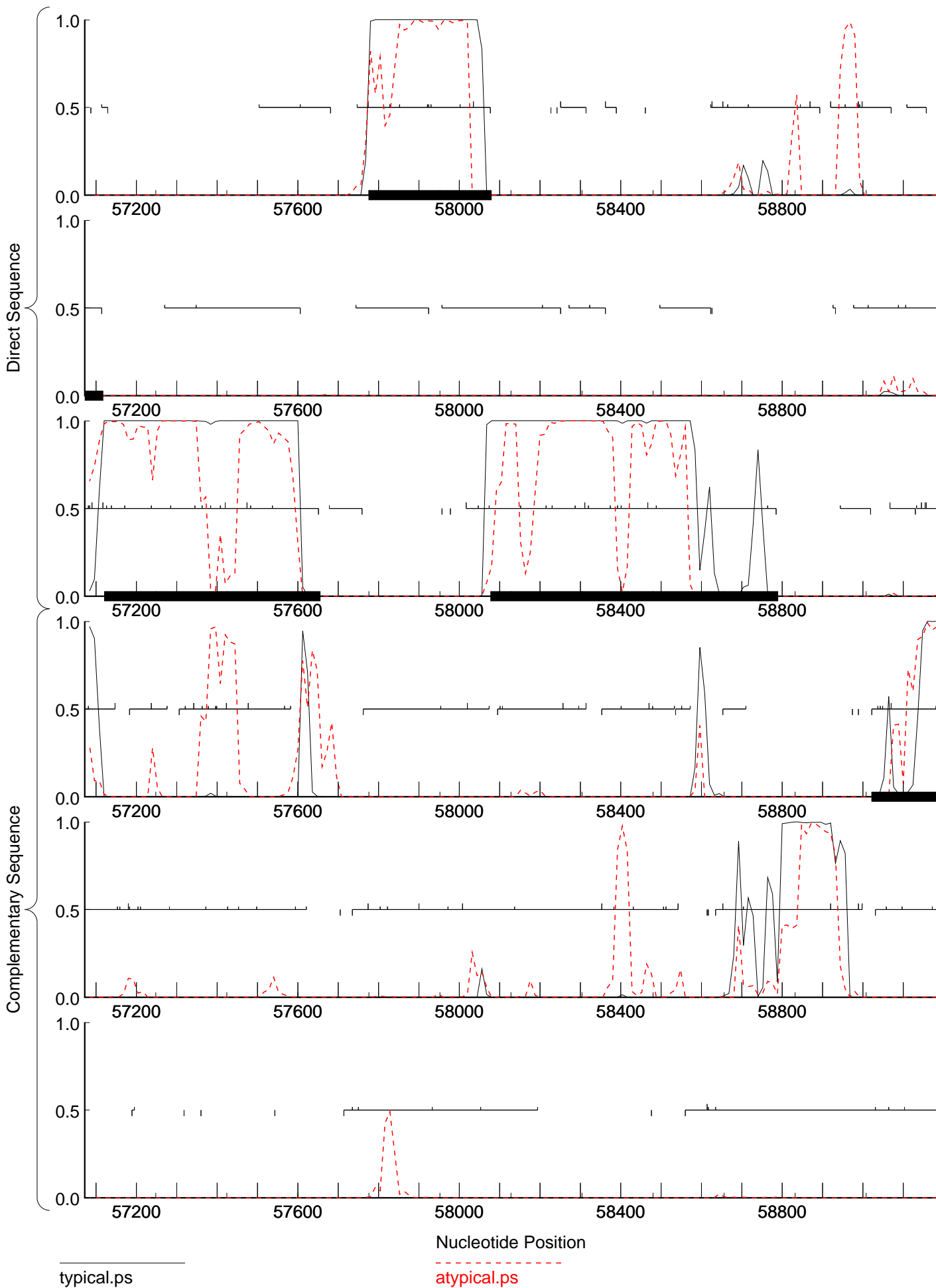


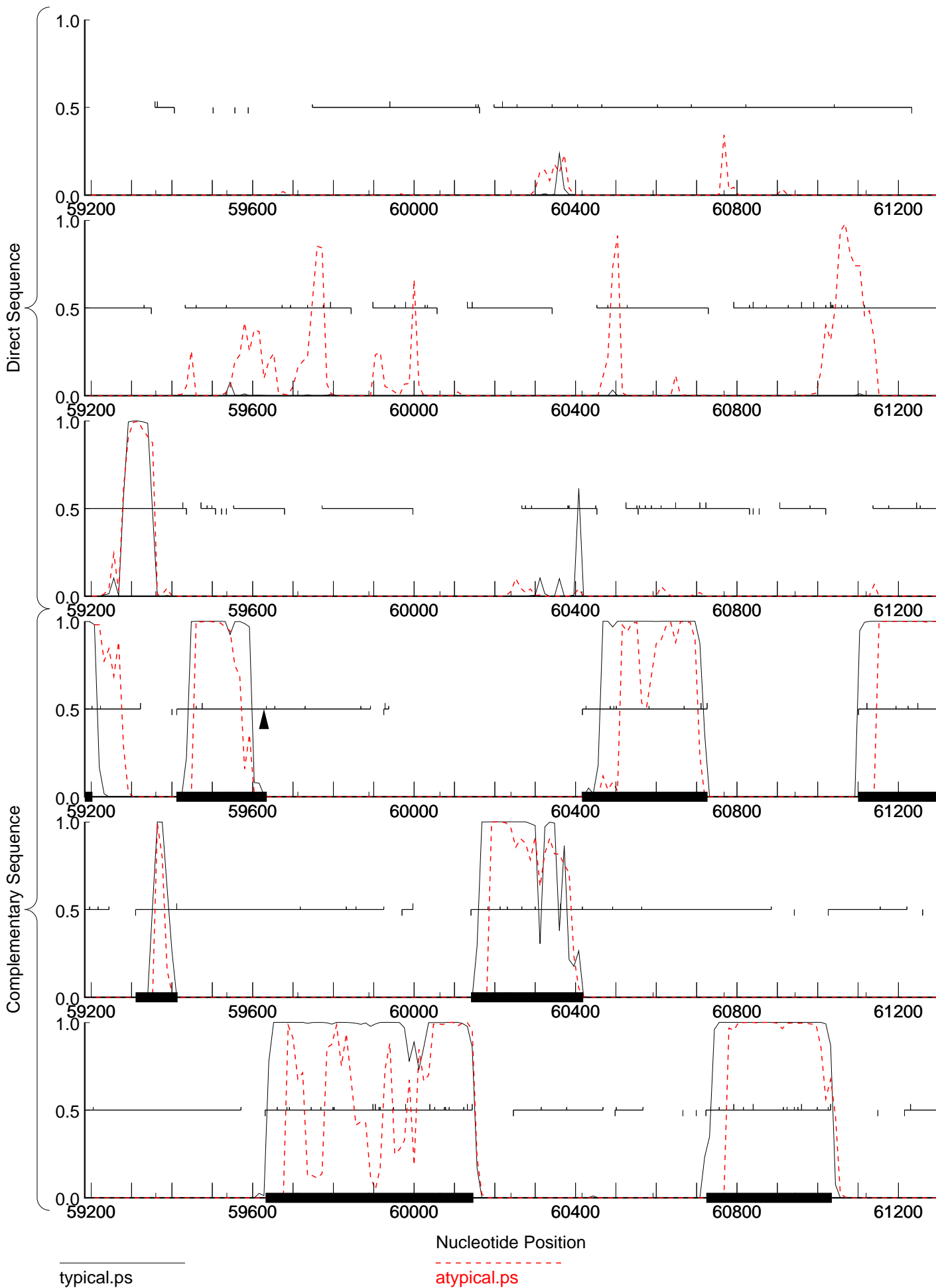


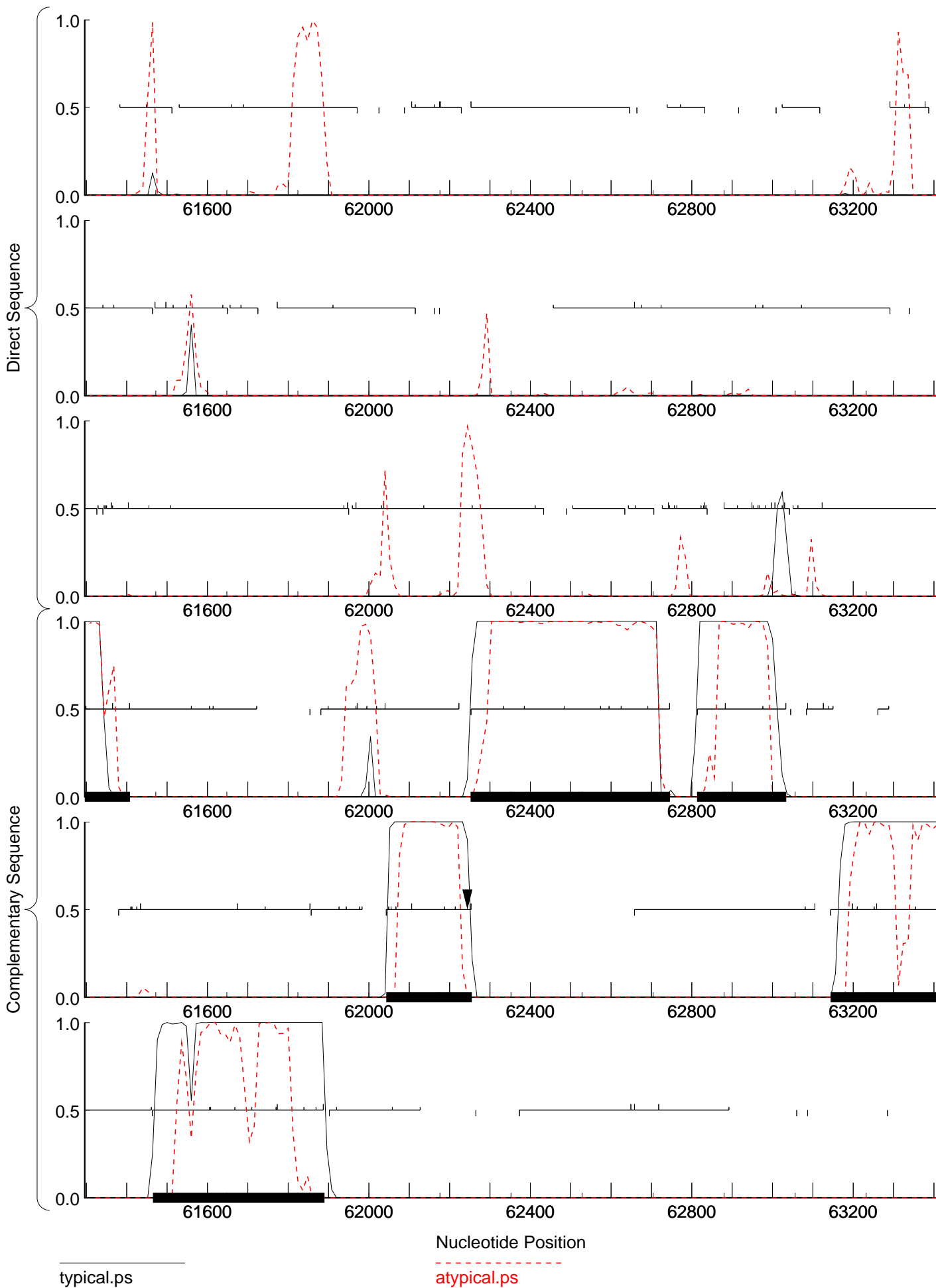


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atypical.ps







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atypical.ps

