

GeneMark

Version 2.5p (09.08.06)

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PROGRAM INFORMATION

Sequence : Mycobacterium phage BubbleTrouble complete sequence, 51397 bp including 10-base 3' overhang
Analysis Date : 2/23/18 at 17:21:55
Pages : 26
Sequence Length : 51397 bp
GC Content : 63.91%

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,

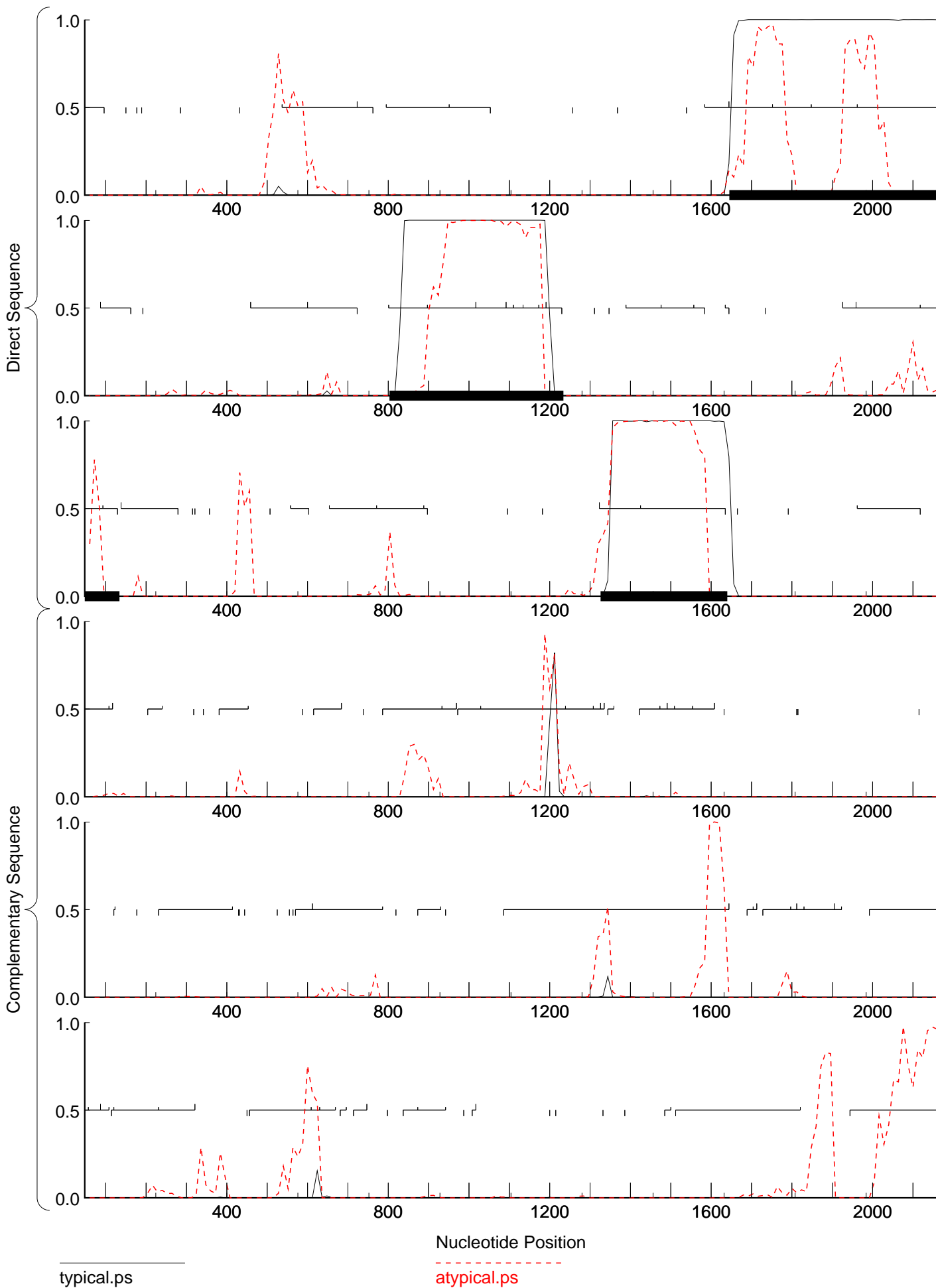
MATRIX INFORMATION

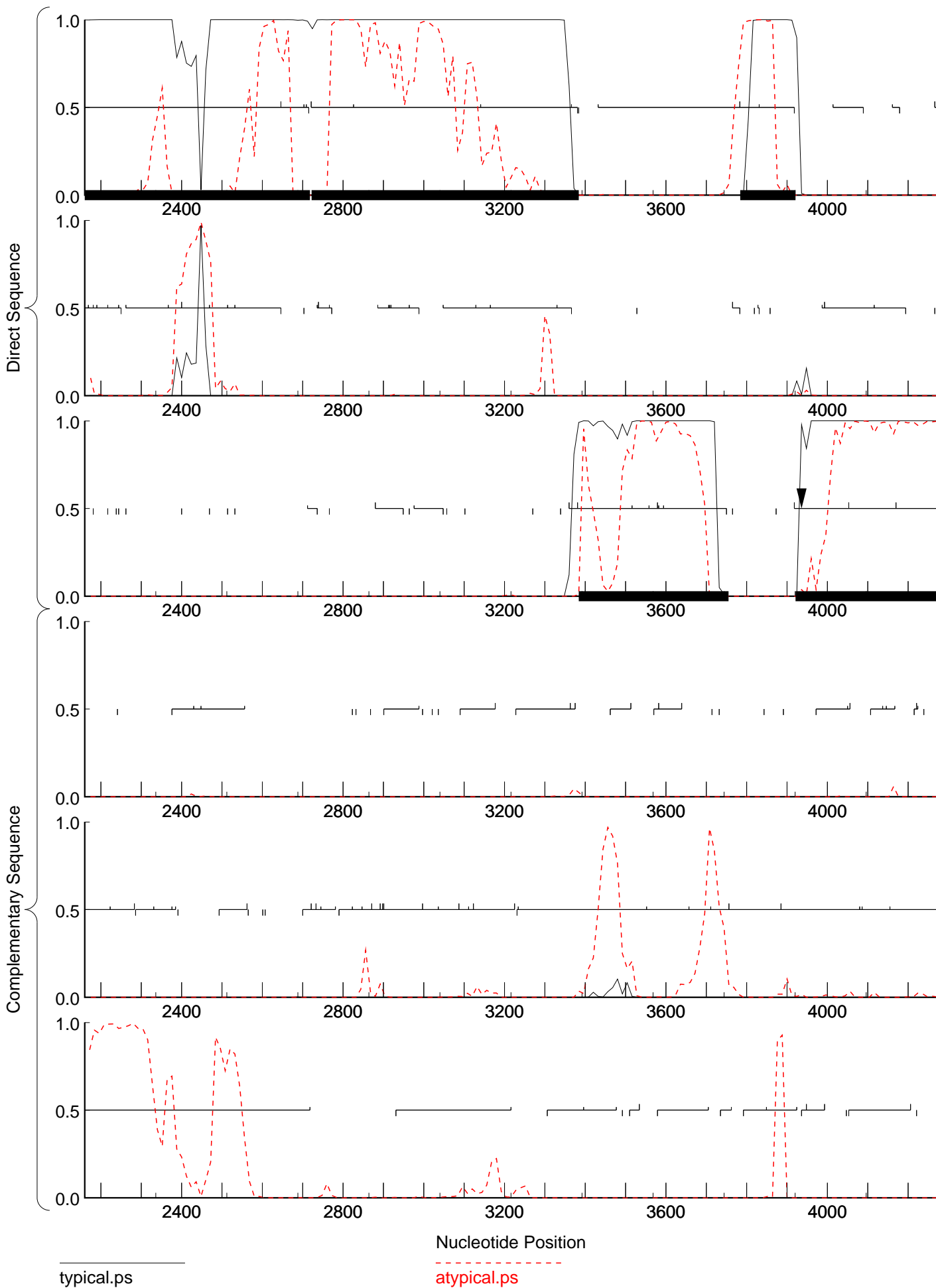
Matrix : -
Author : -
Order : 4

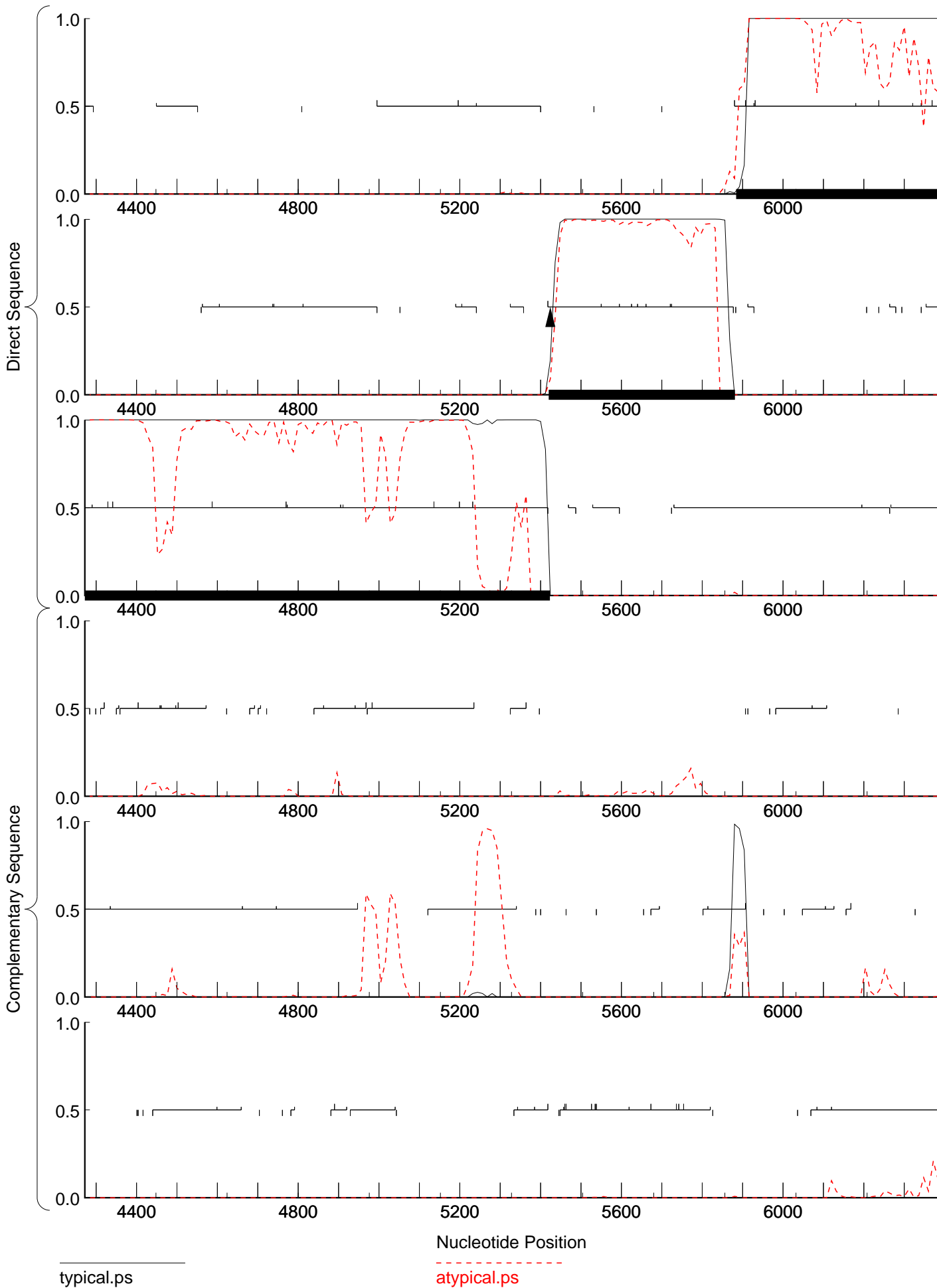
Send questions / comments to:
Dr. M. Borodovsky
Georgia Institute of Technology
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Atlanta, GA 30332-0230

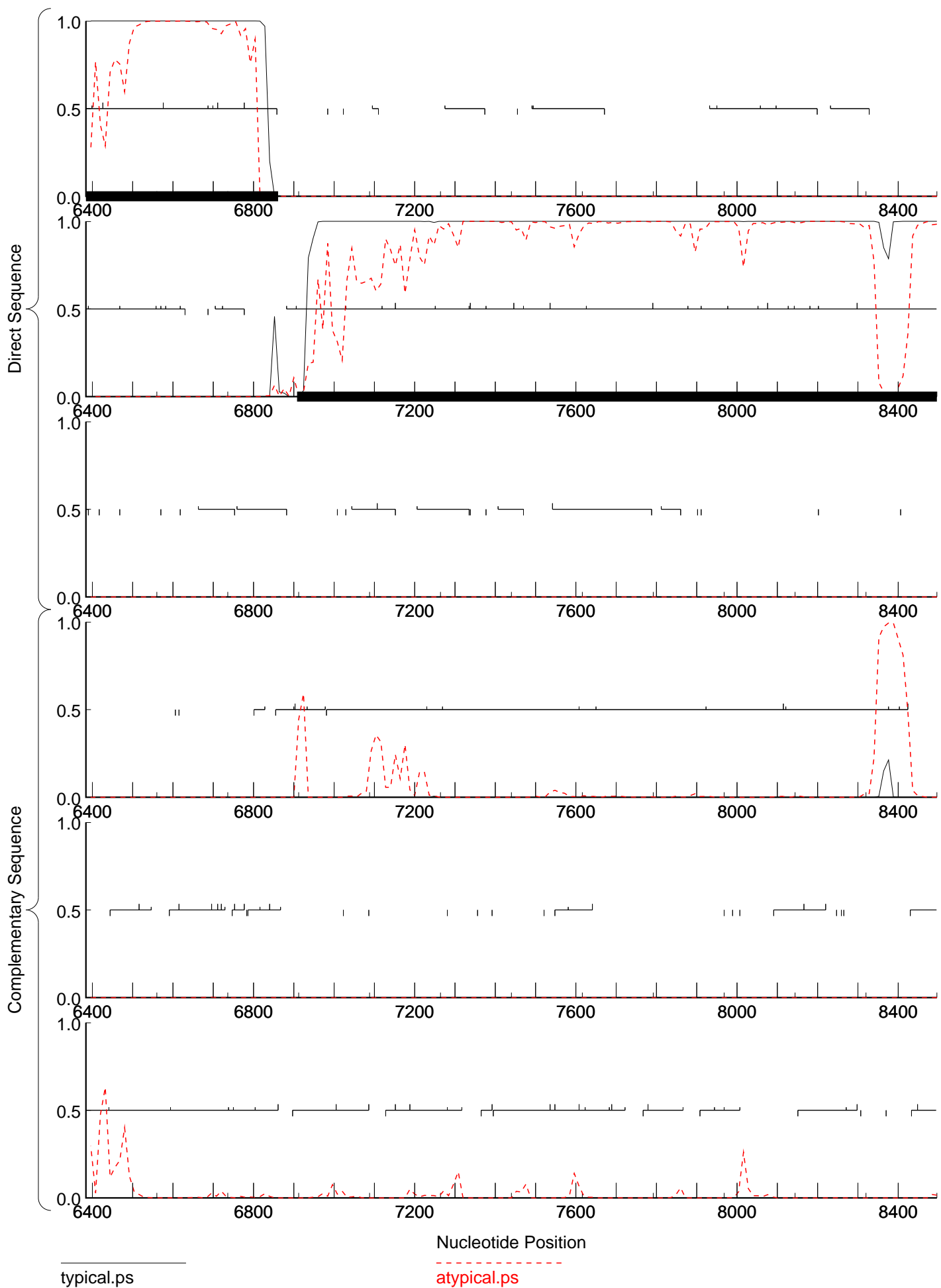
Matrix notes & comments

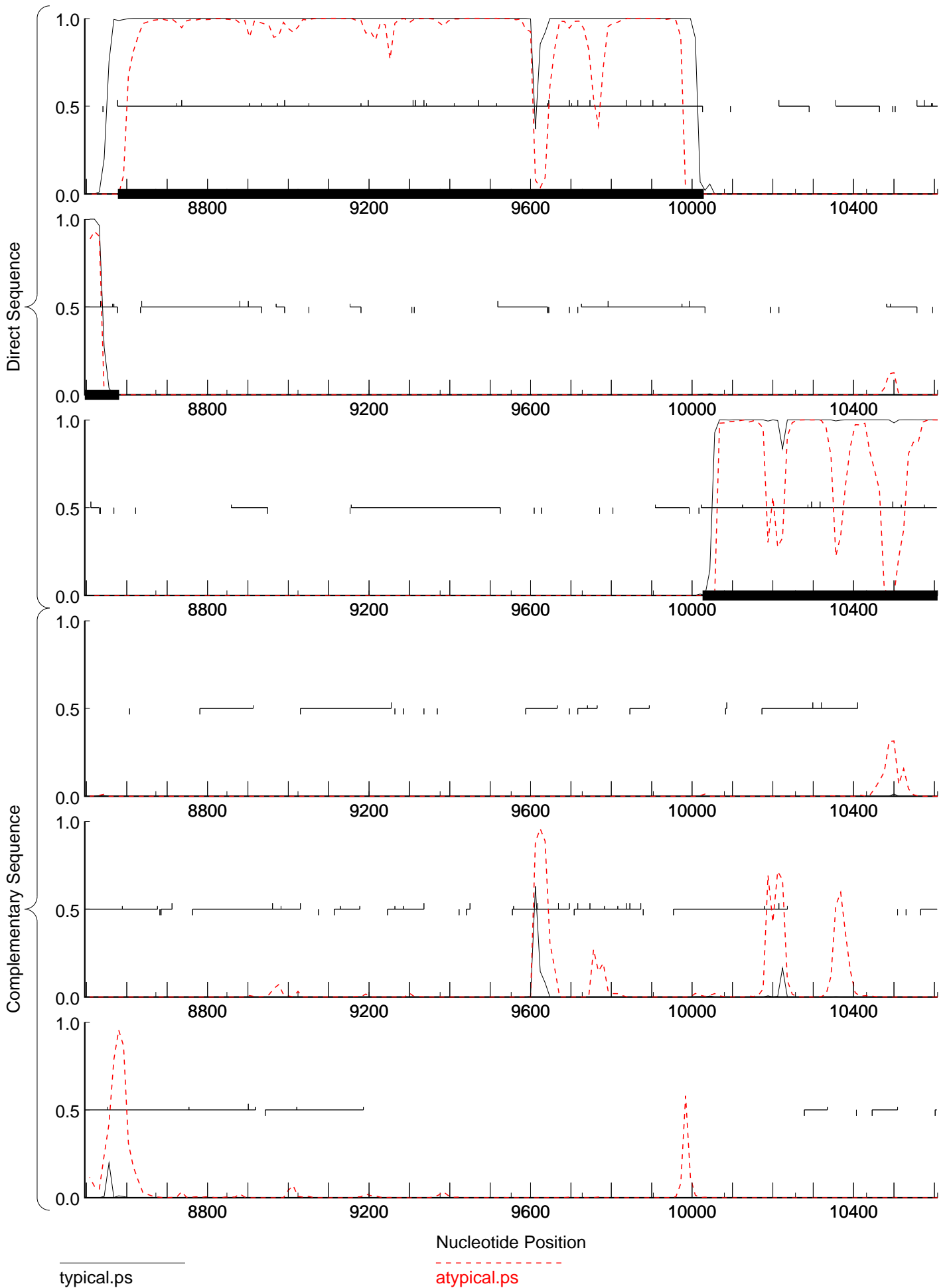
Training set derived by GeneMarkS, 4.30 October 2014
Fri Feb 23 17:21:55 2018

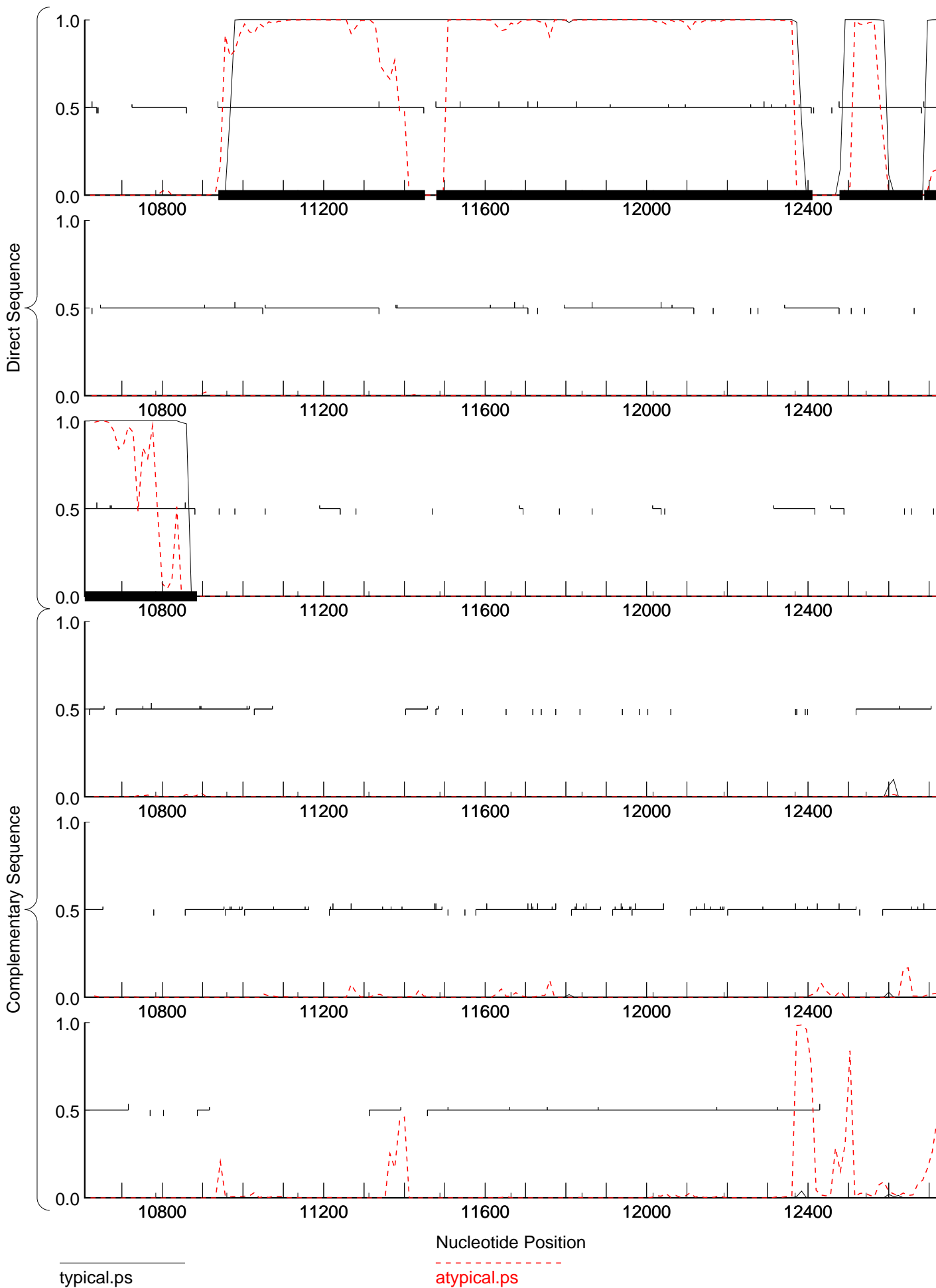


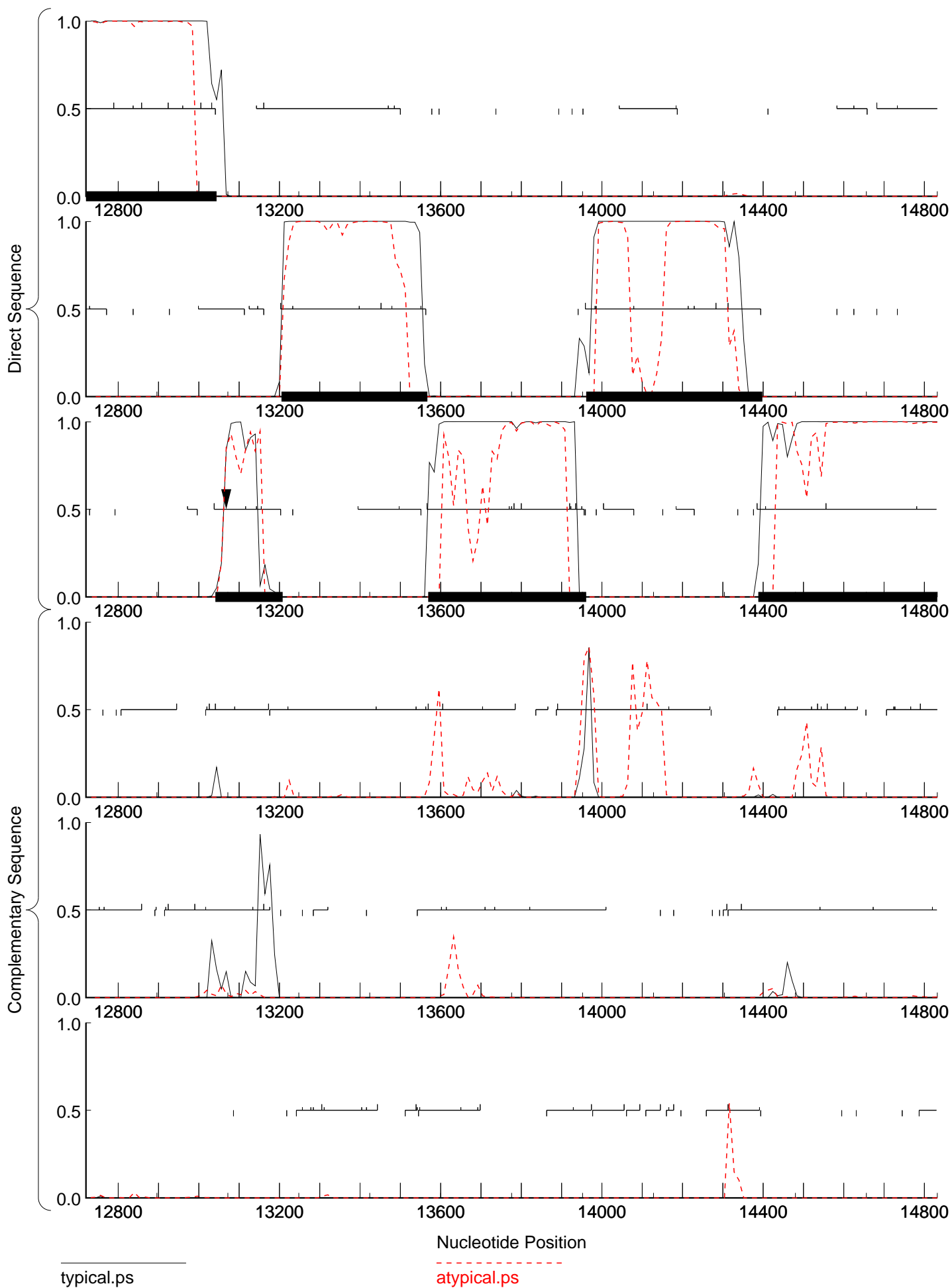


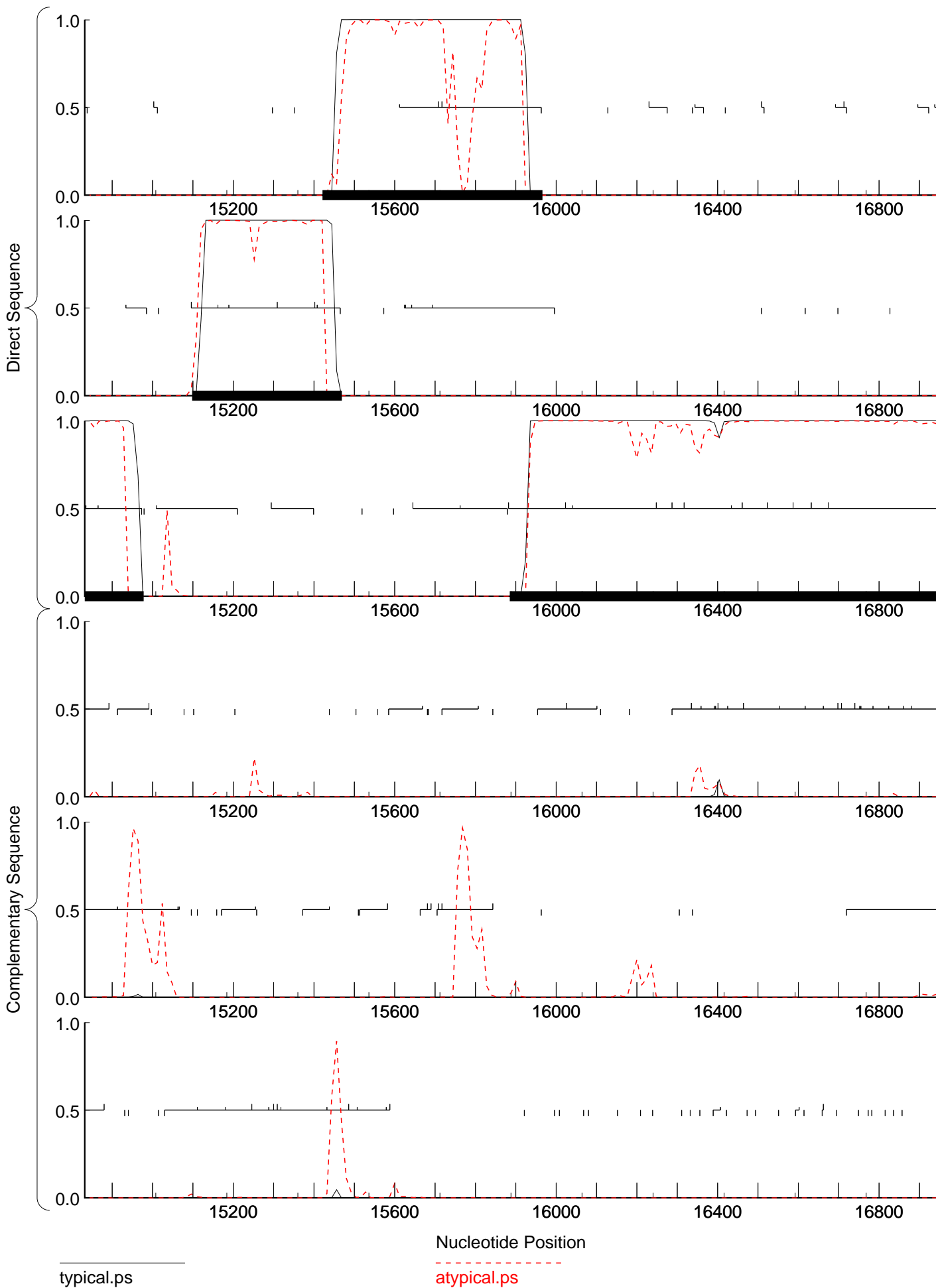


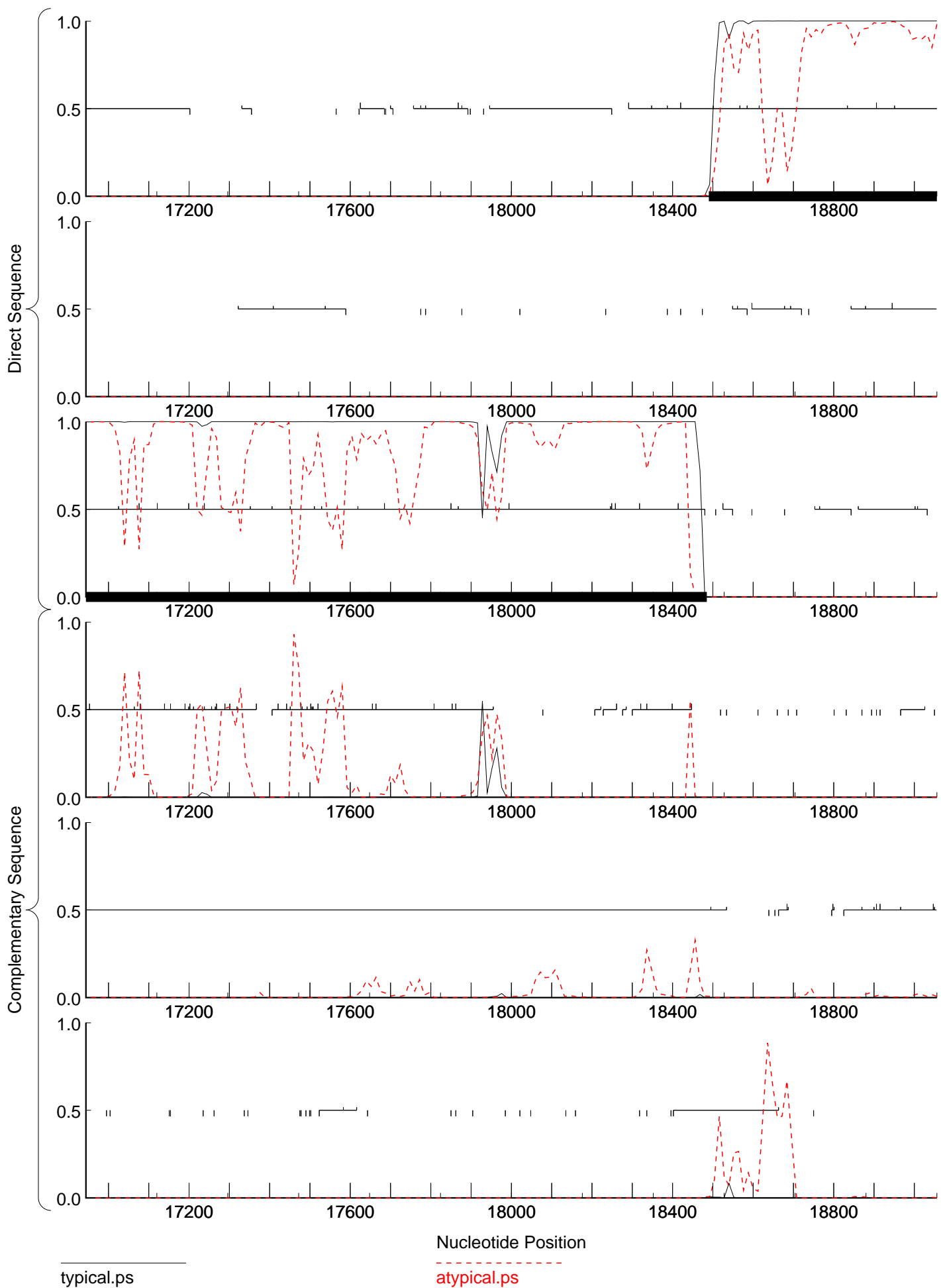


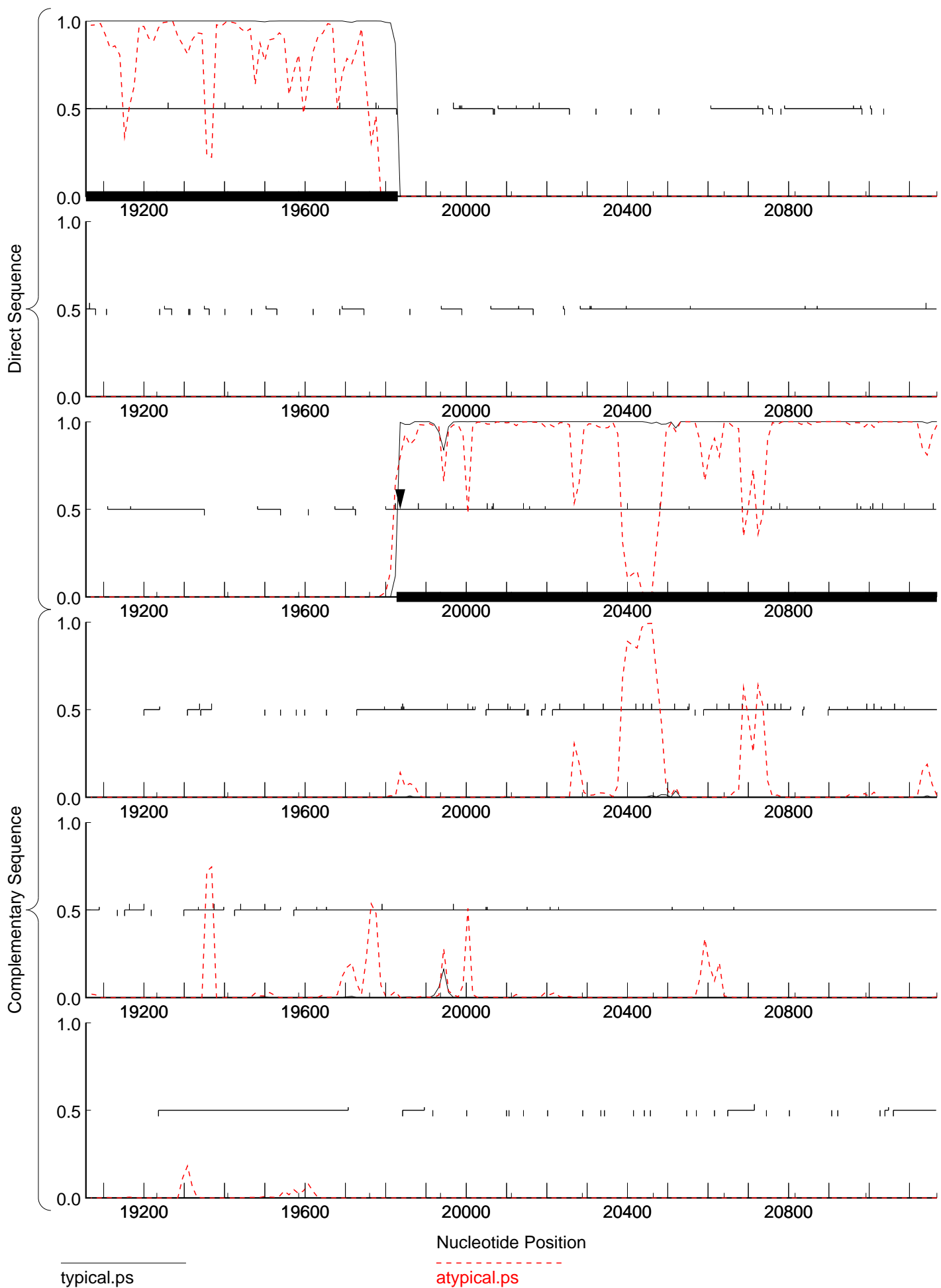


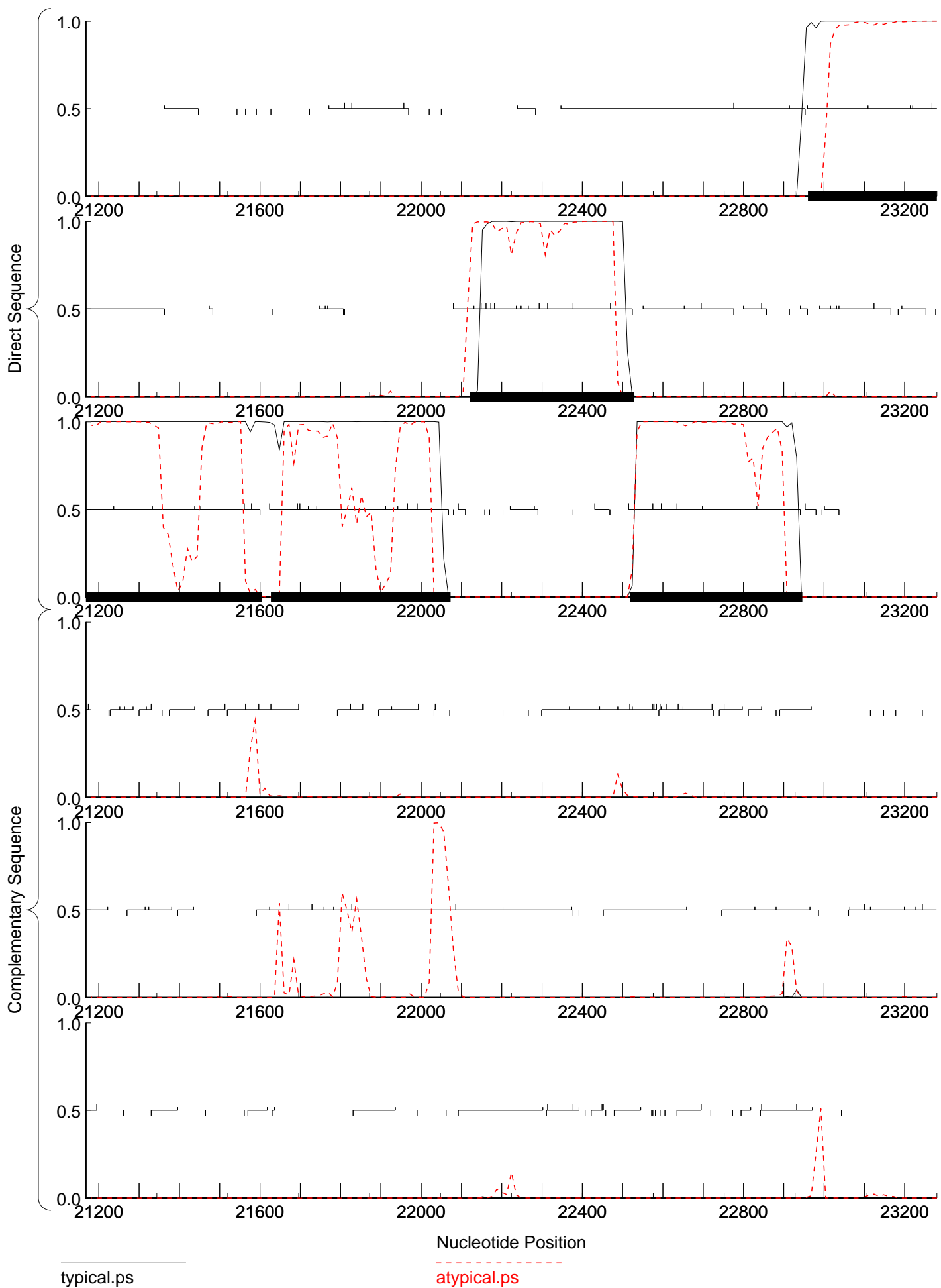




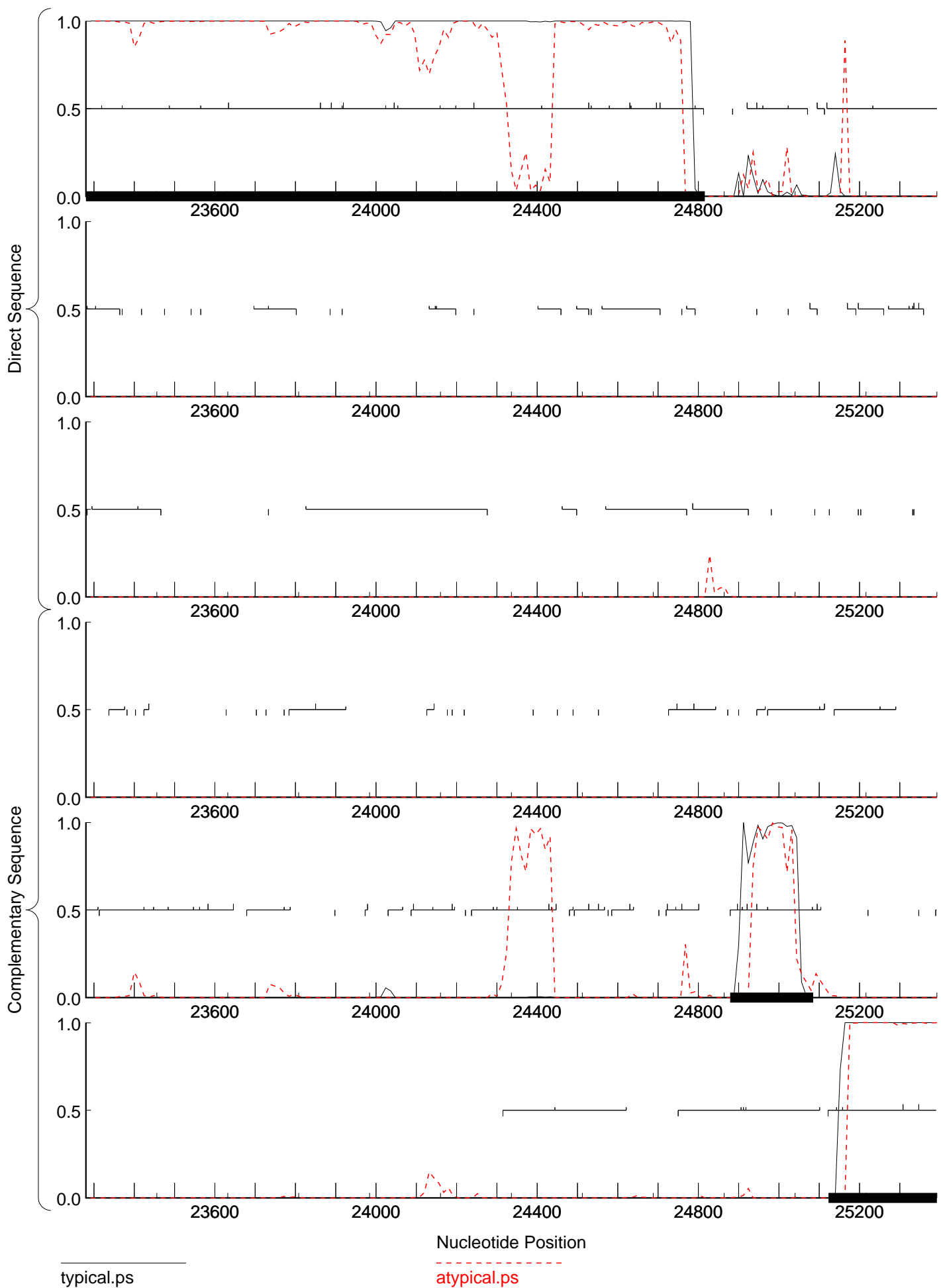


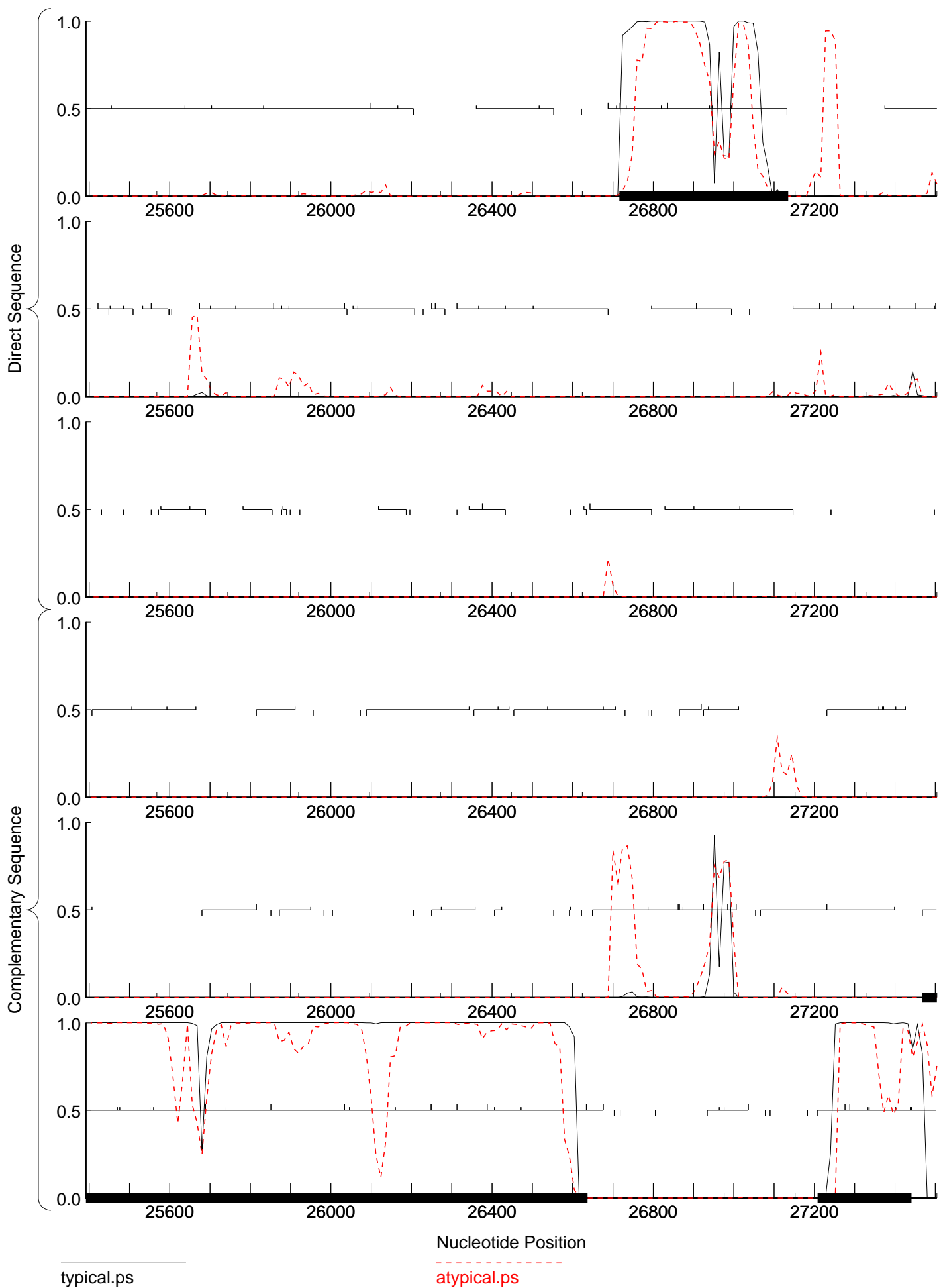






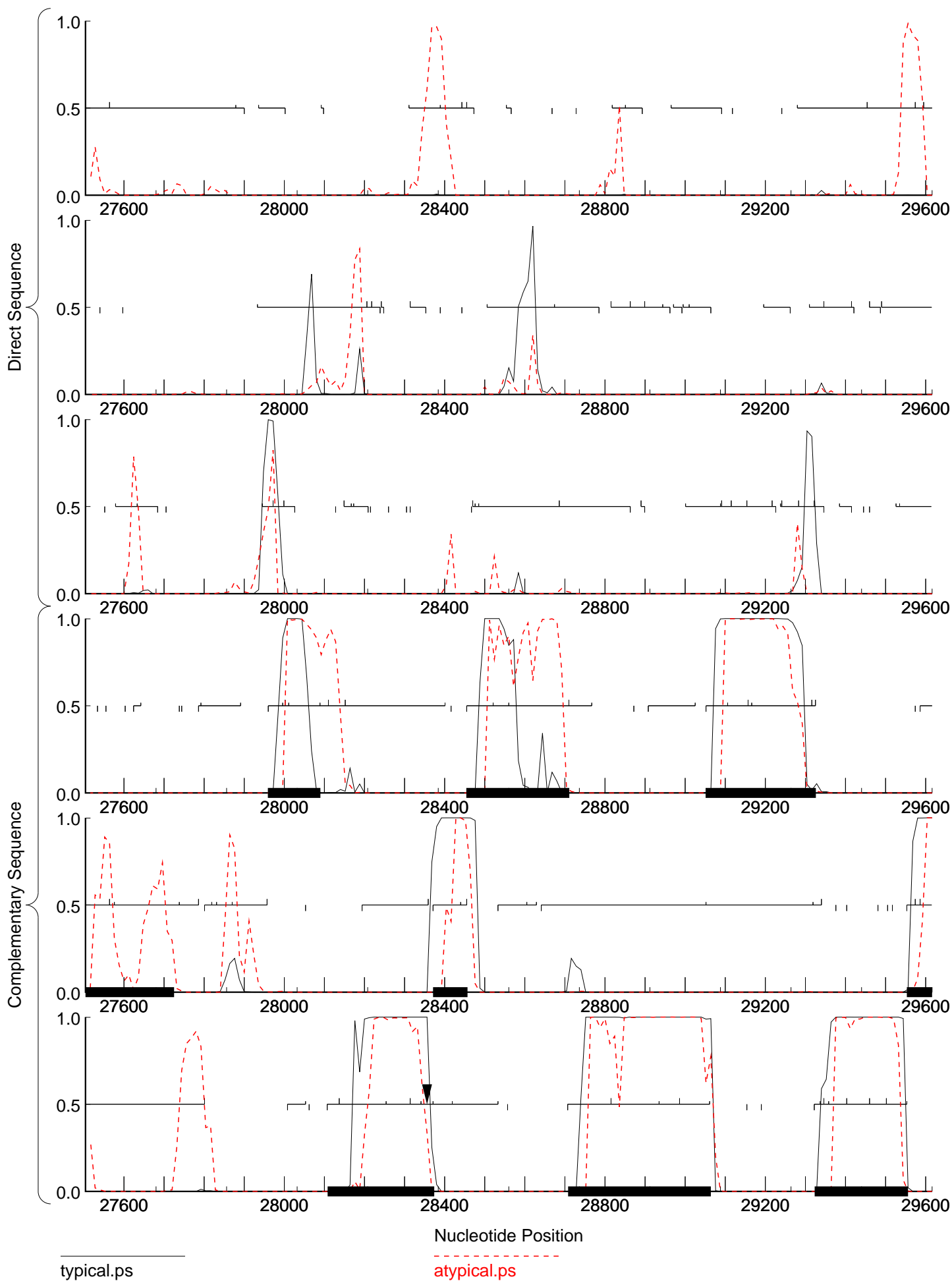
complete sequence, 51397 bp including 10-base 3' overhang (CGGCCGGTAA), Cluster A4, Order 4, Window 96, Step 12, 13/26
complete sequence, 51397 bp including 10-base 3' overhang (CGGCCGGTAA), Cluster A4, Order 2, Window 96, Step 12, 13/26

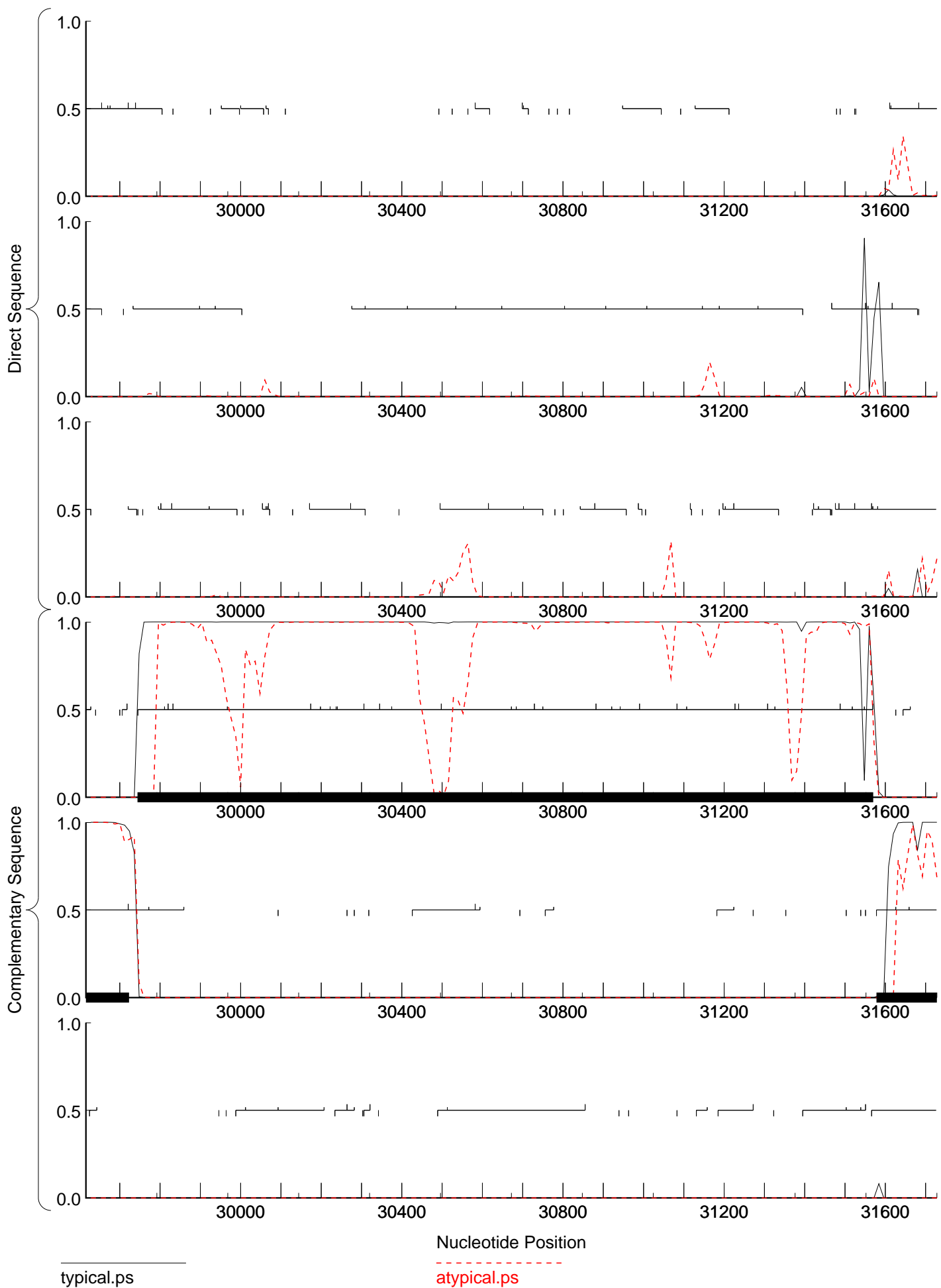


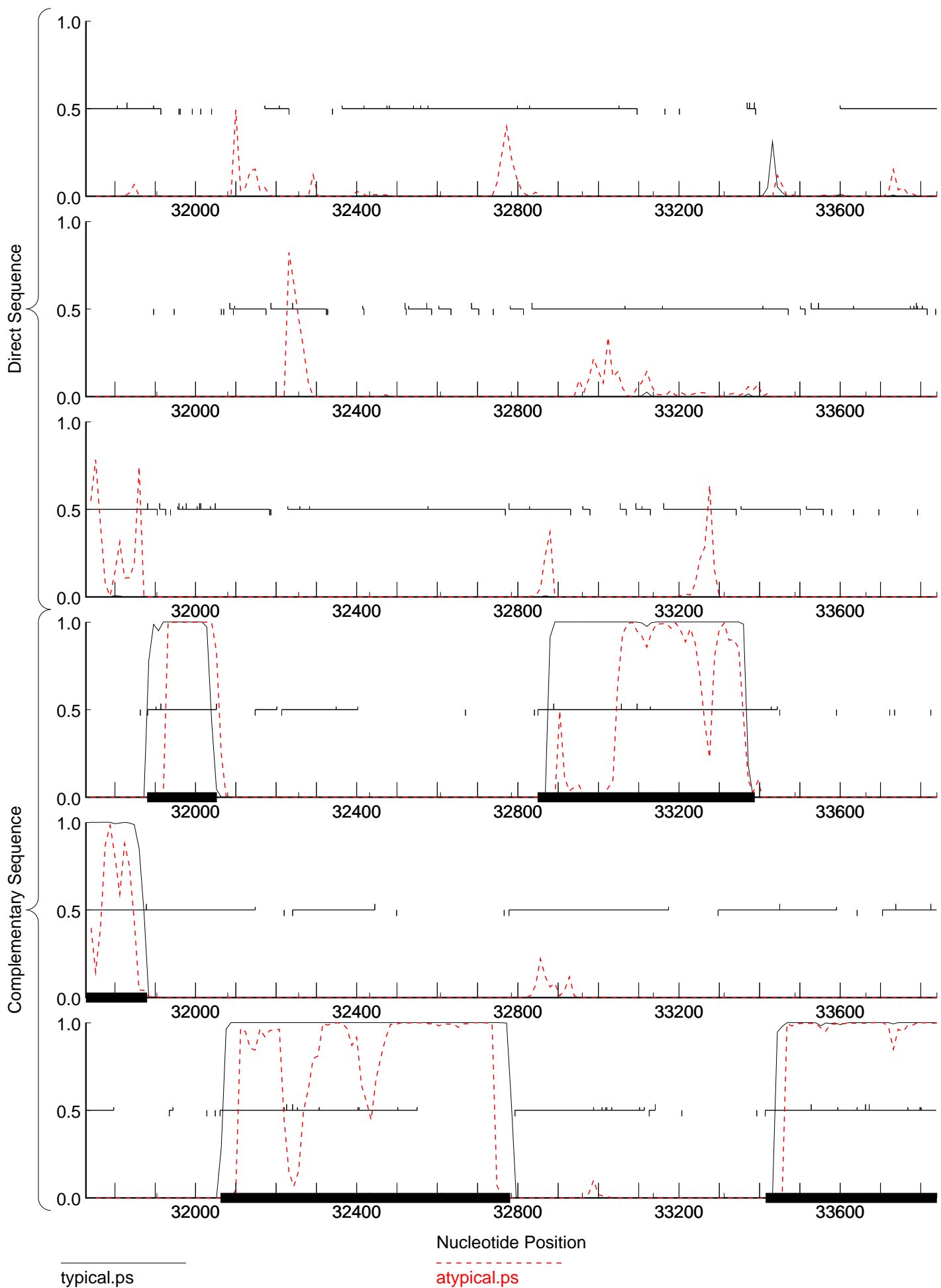


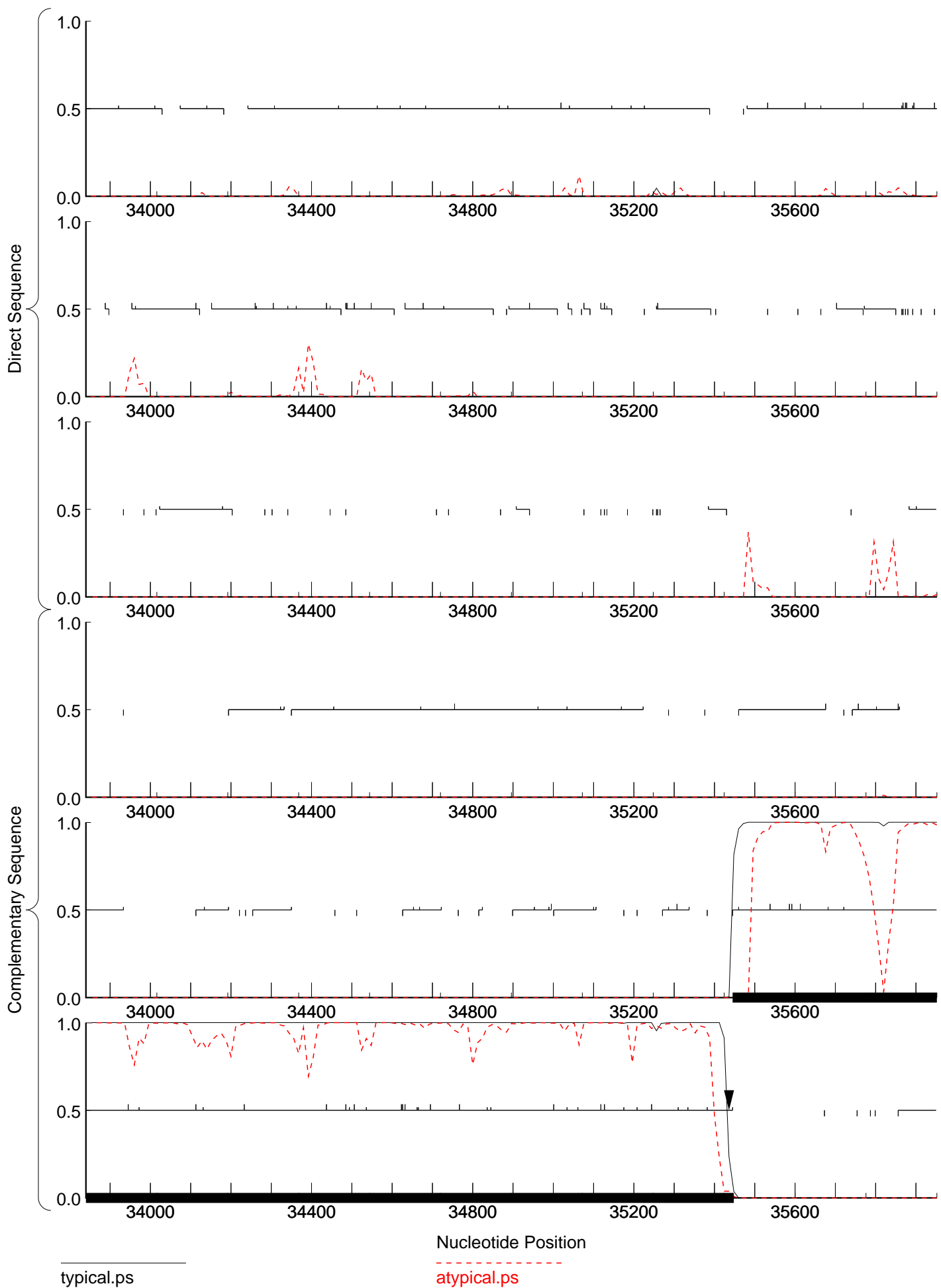
typical.ps

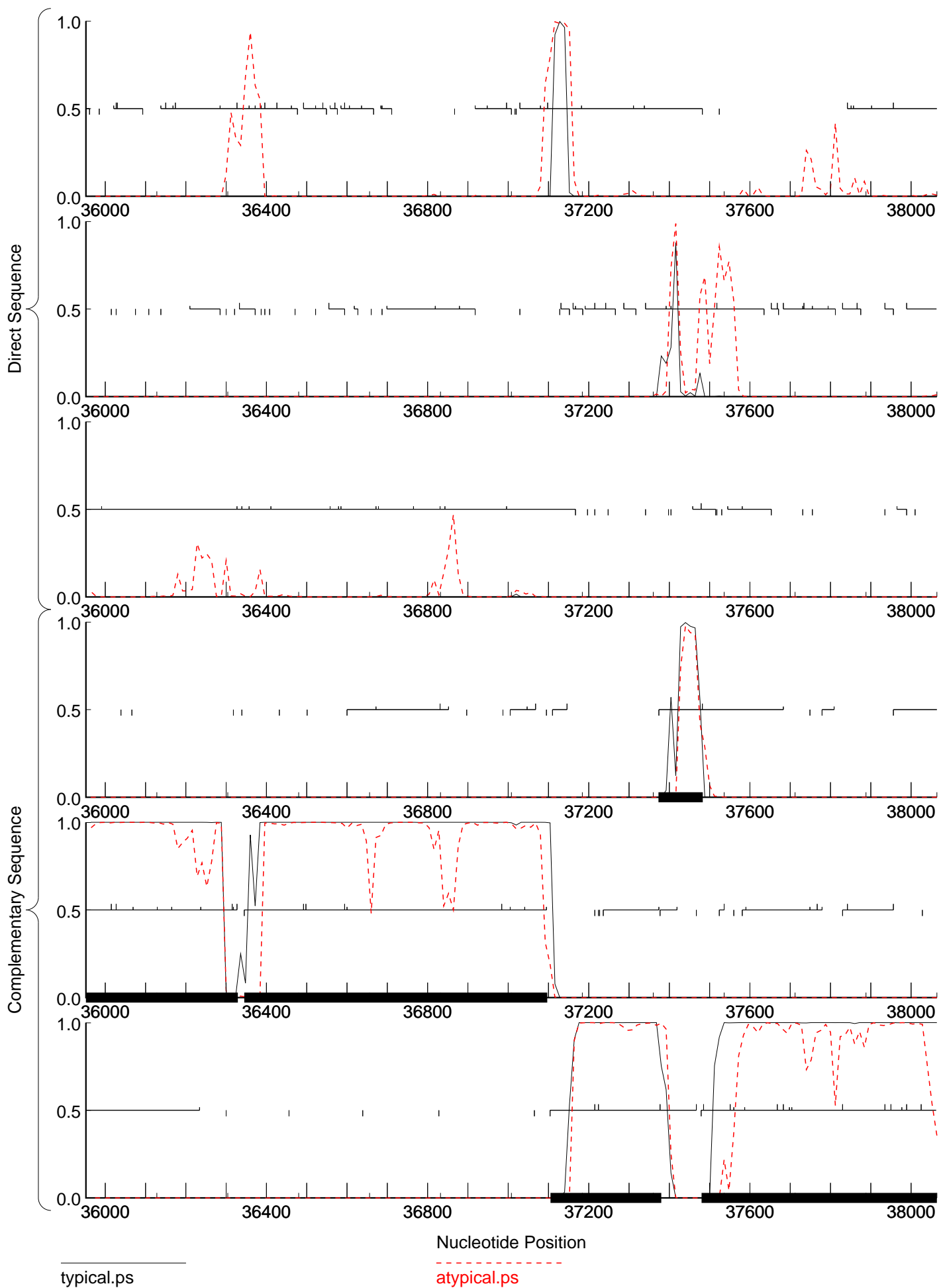
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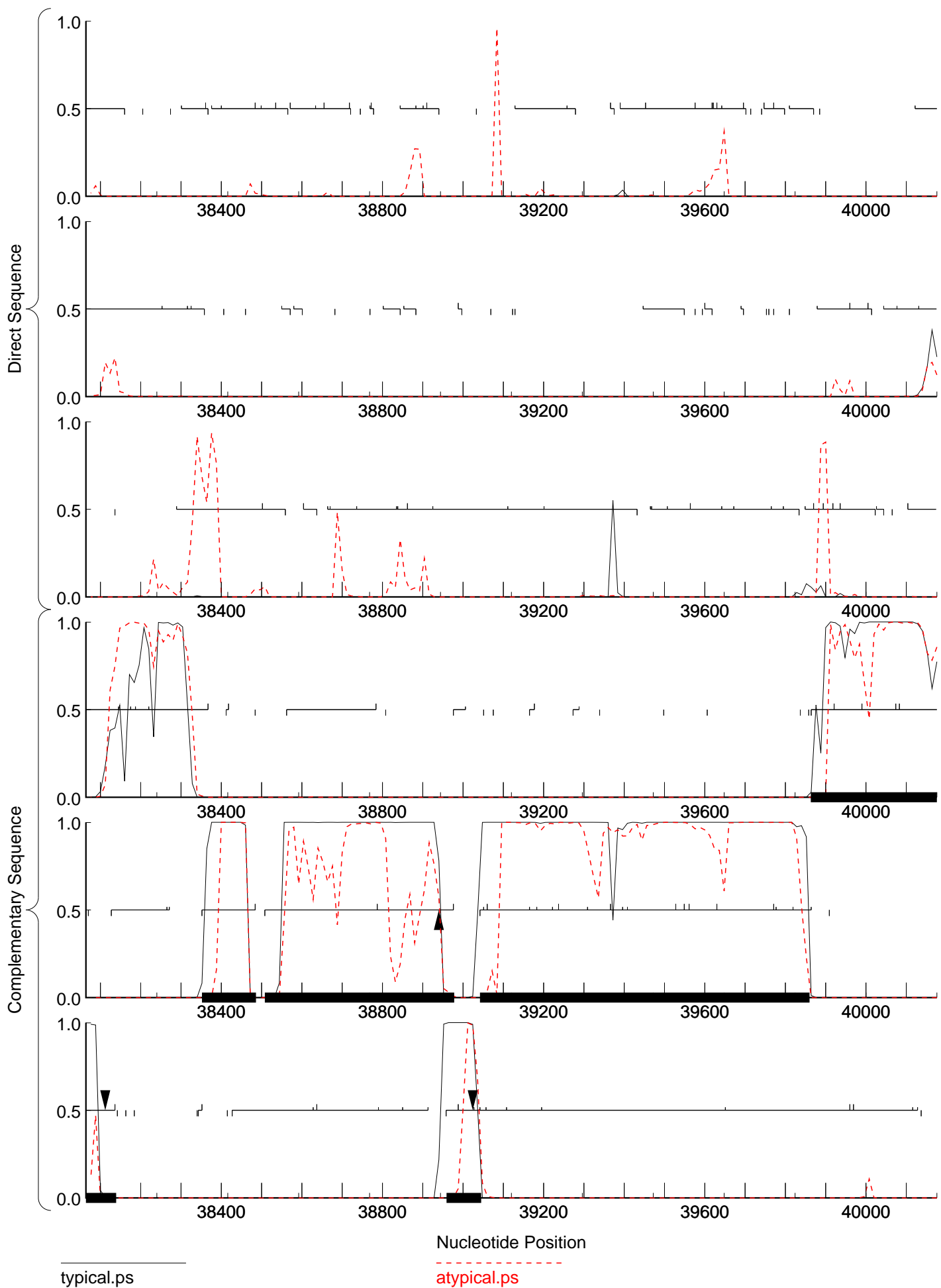


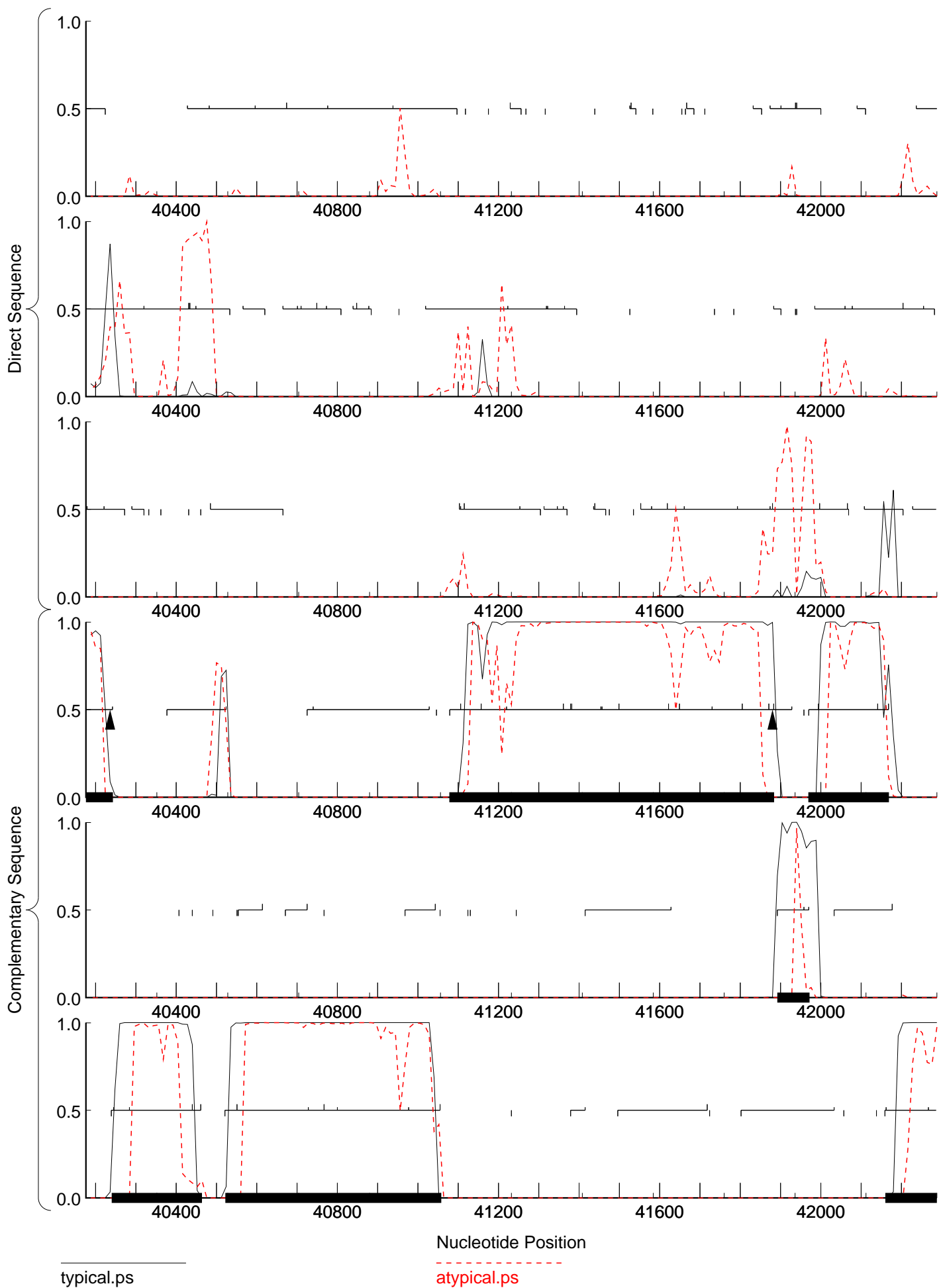


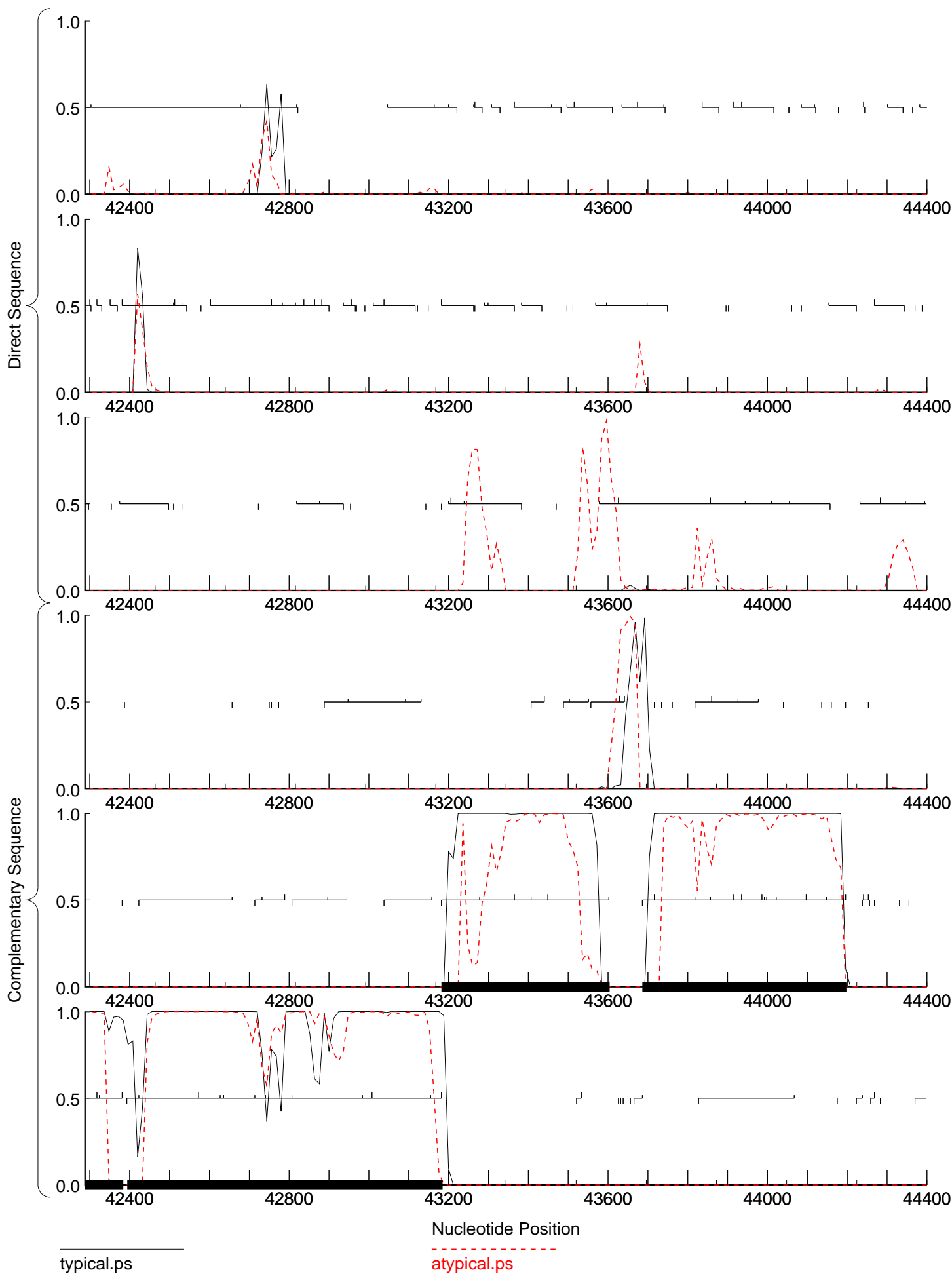












typical.ps

atypical.ps

