

GeneMark

Version 2.5p (09.08.06)

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PROGRAM INFORMATION

Sequence : Mycobacterium phage Geraldini complete sequence, 59818 bp including 11-base 3' overhang (CT
Analysis Date : 5/27/18 at 21:40:01
Pages : 30
Sequence Length : 59818 bp
GC Content : 66.61%

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,

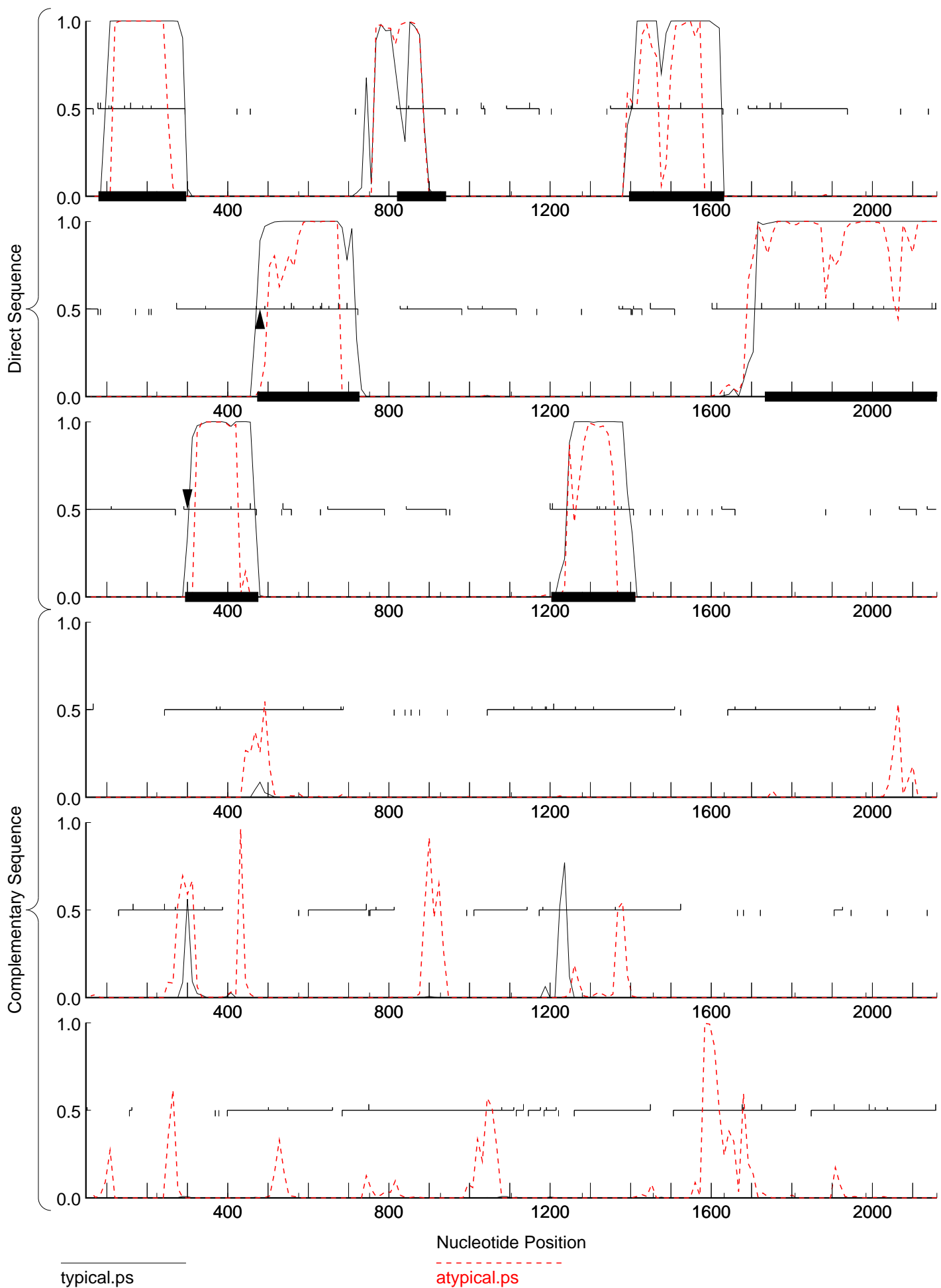
MATRIX INFORMATION

Matrix : -
Author : -
Order : 4

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

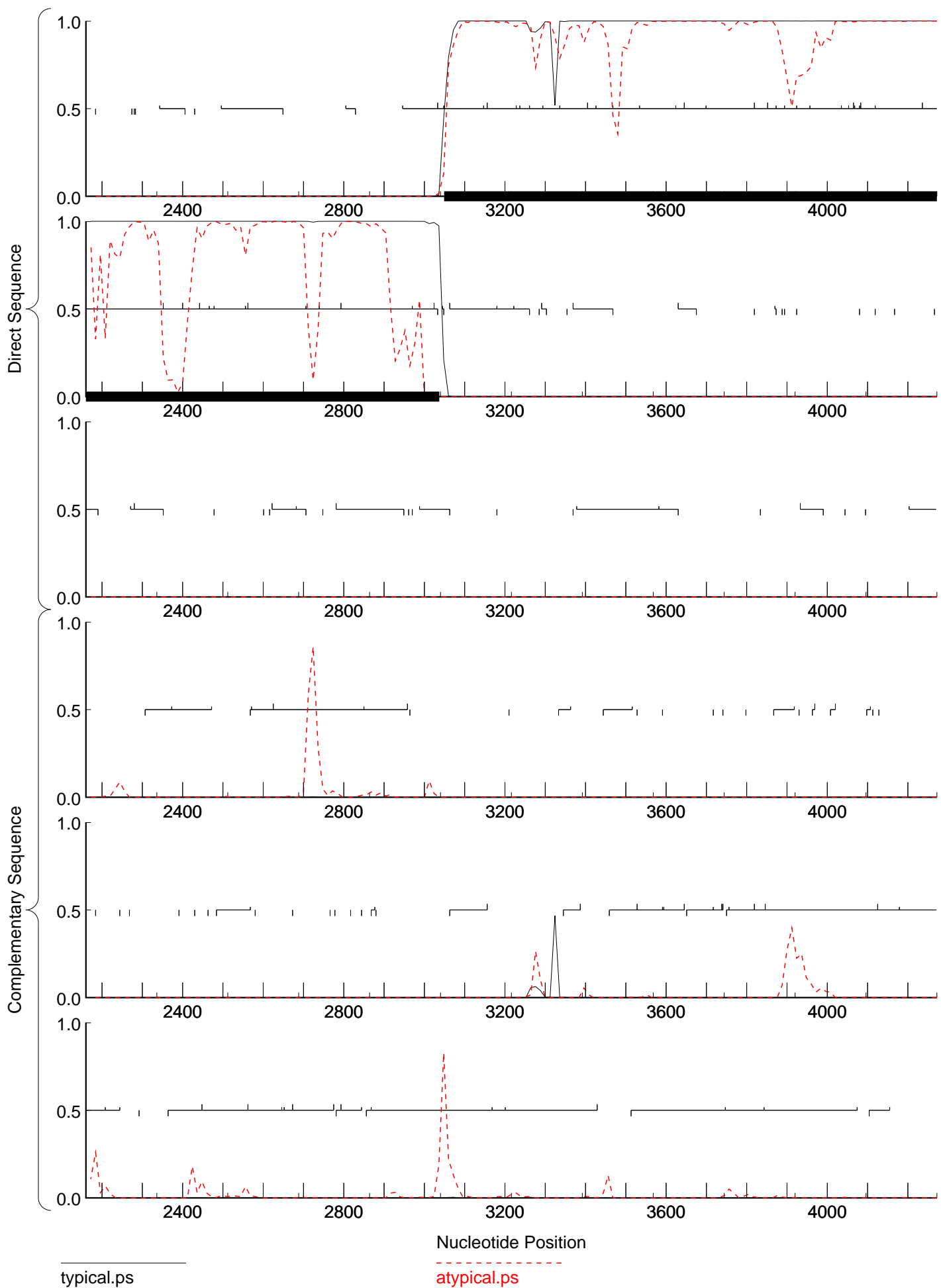
Matrix notes & comments

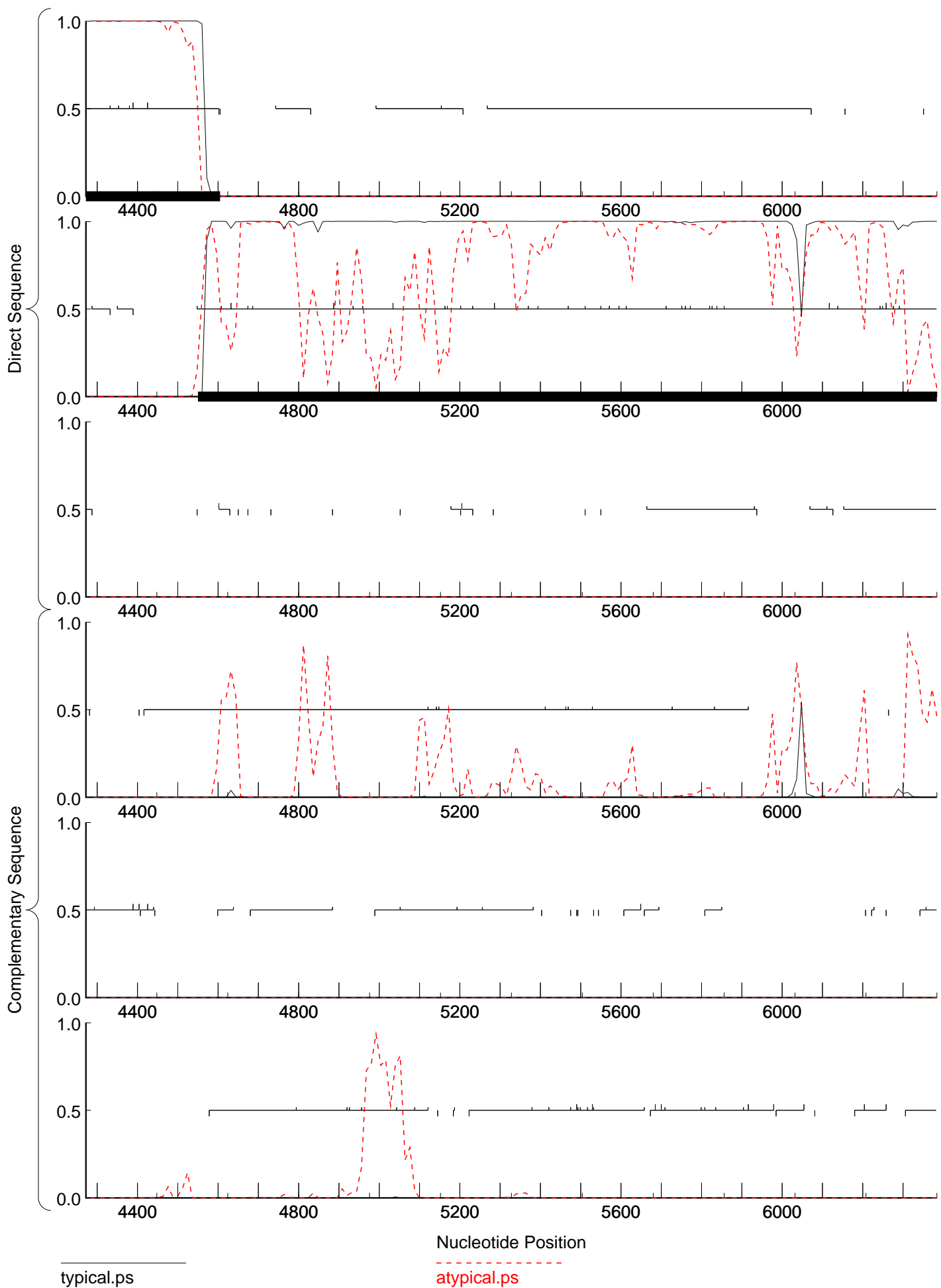
Training set derived by GeneMarkS, 4.30 October 2014
Sun May 27 21:40:00 2018



typical.ps

atypical.ps

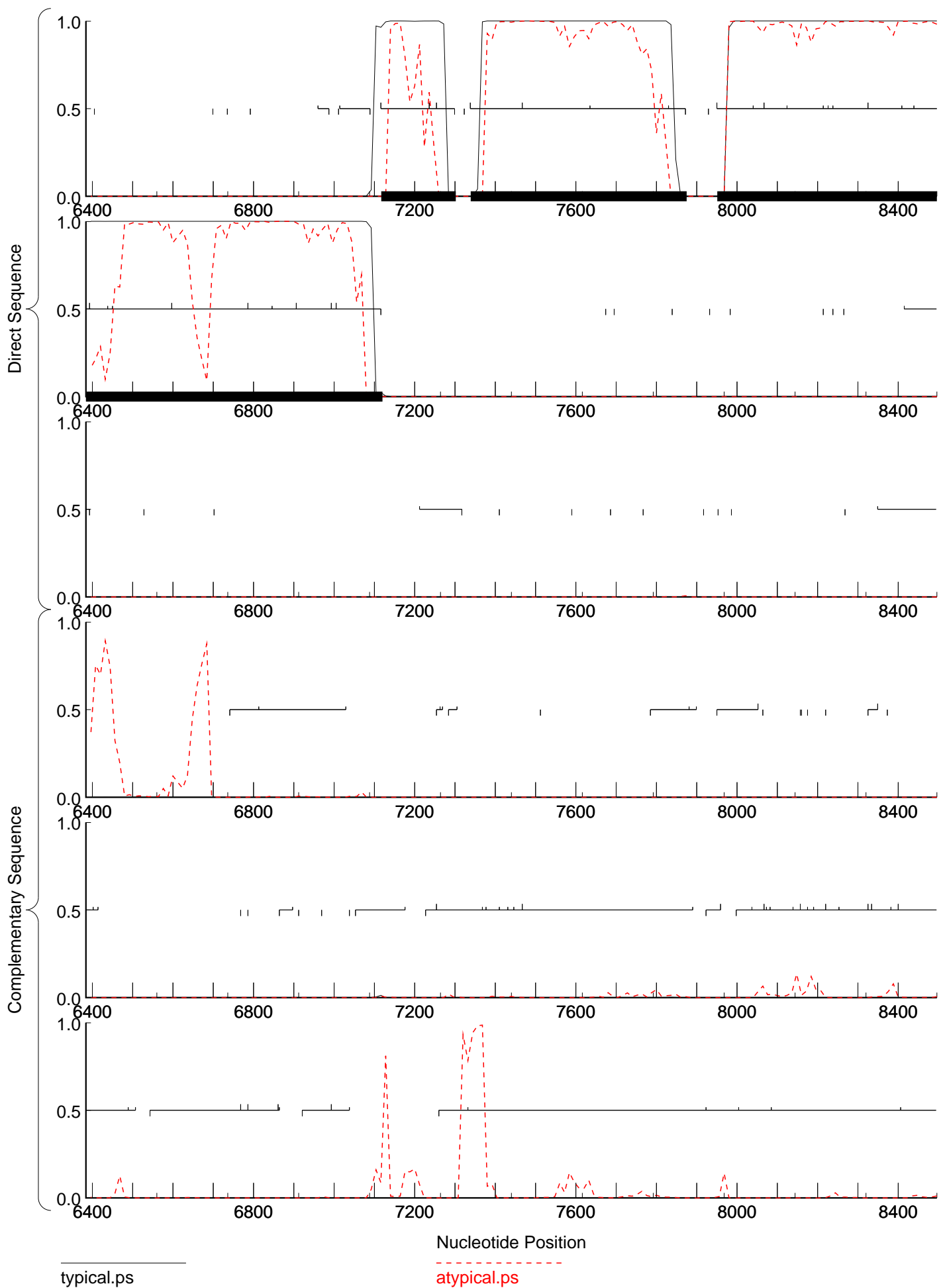


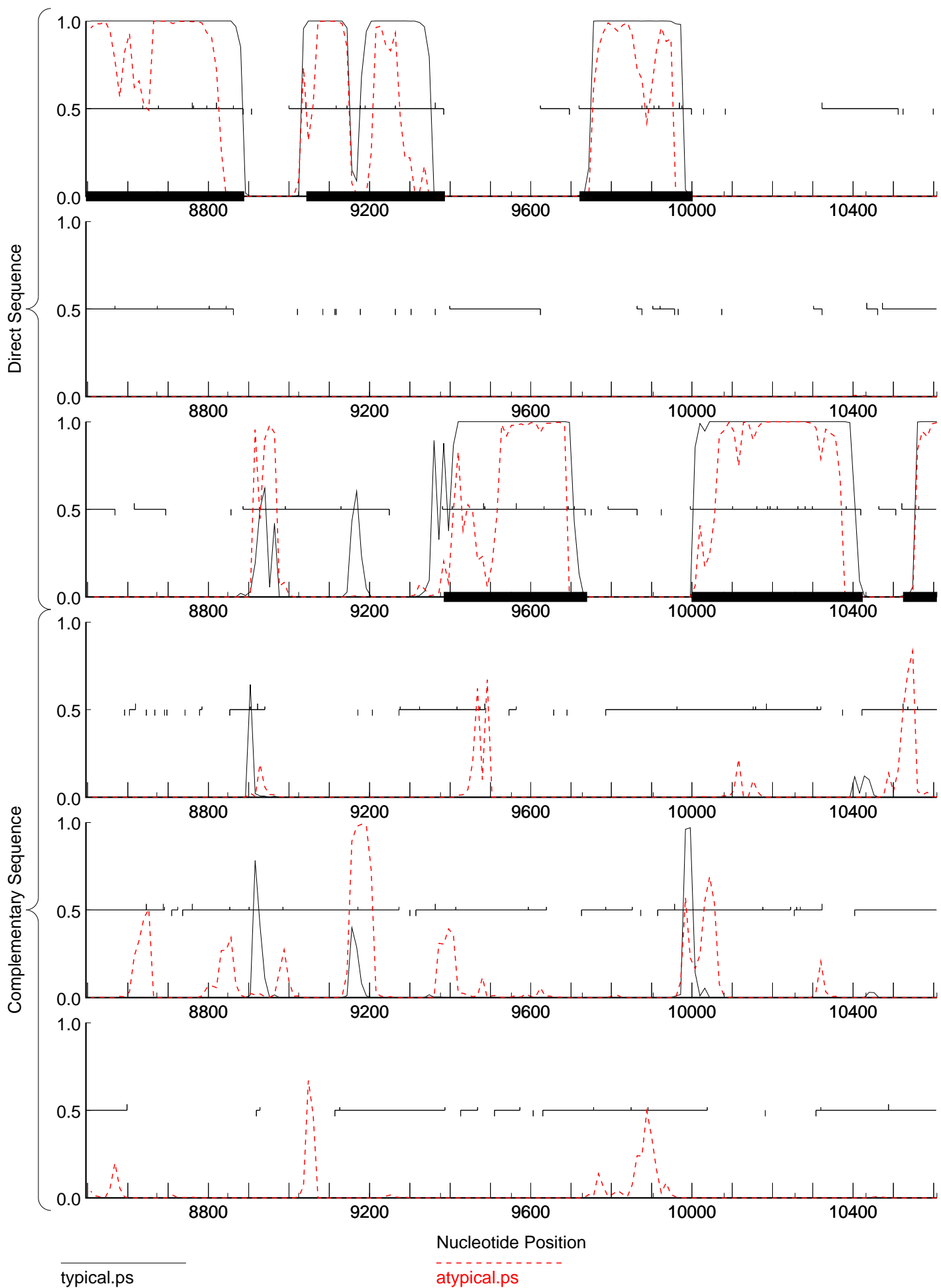


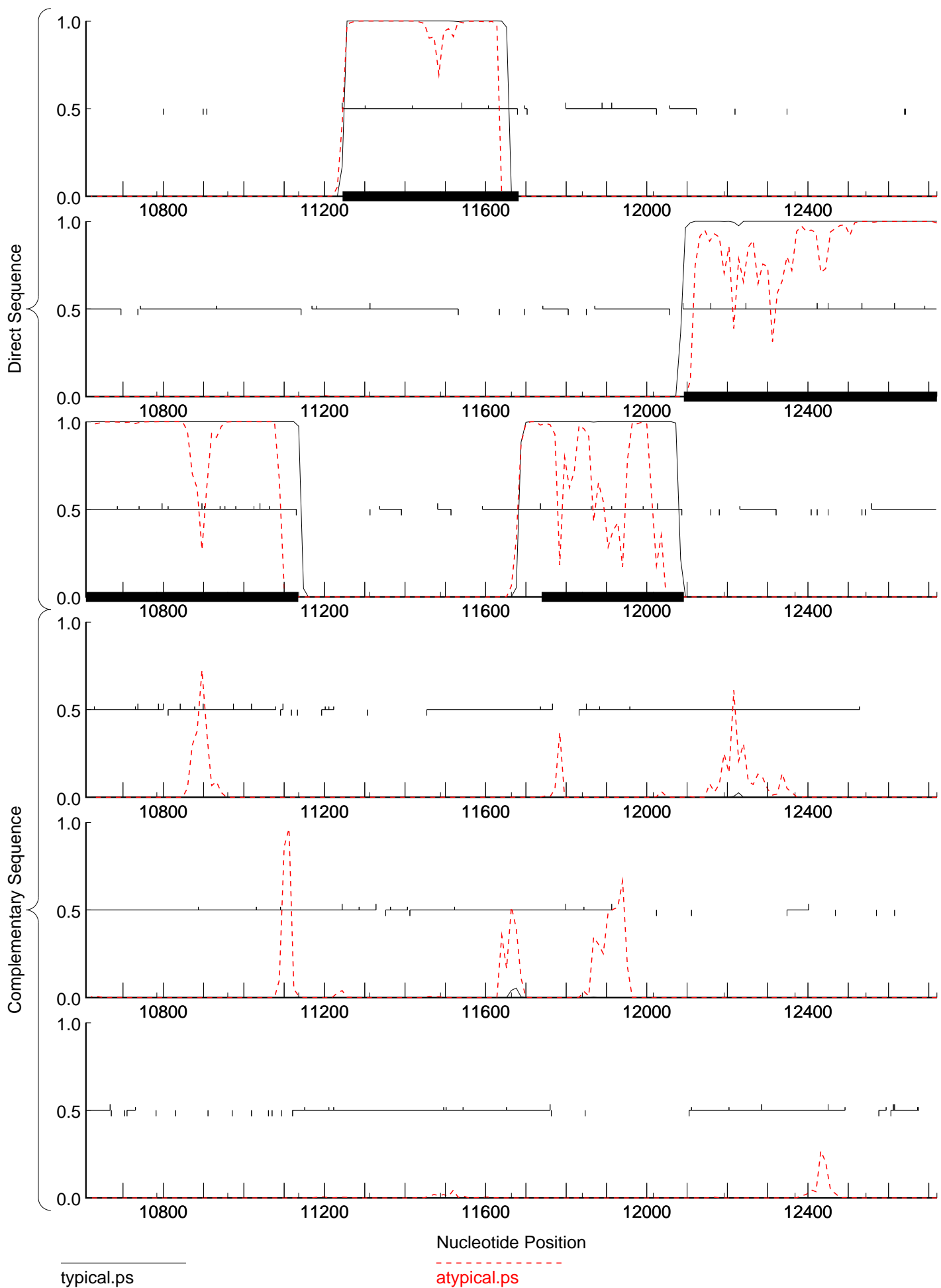
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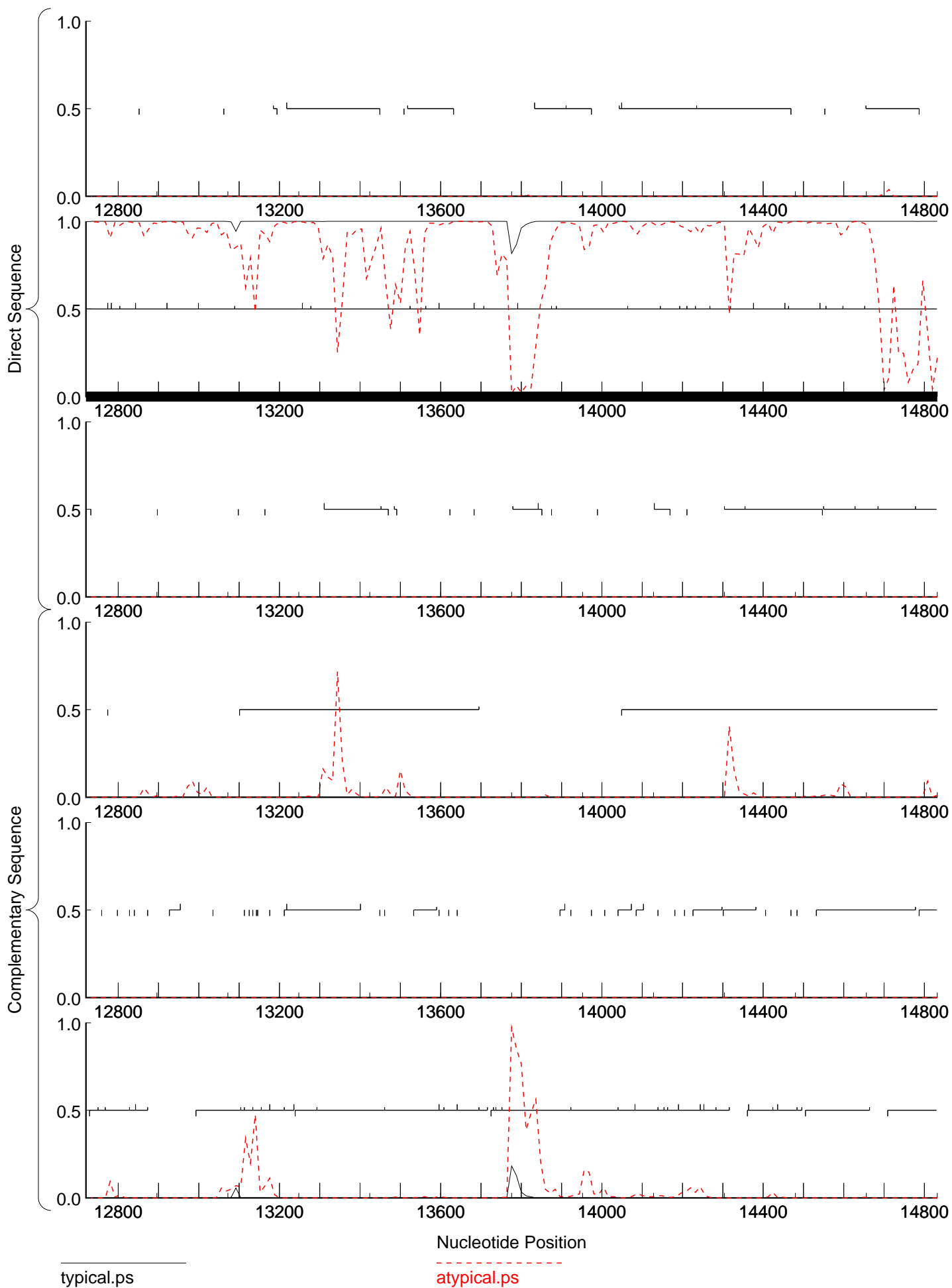
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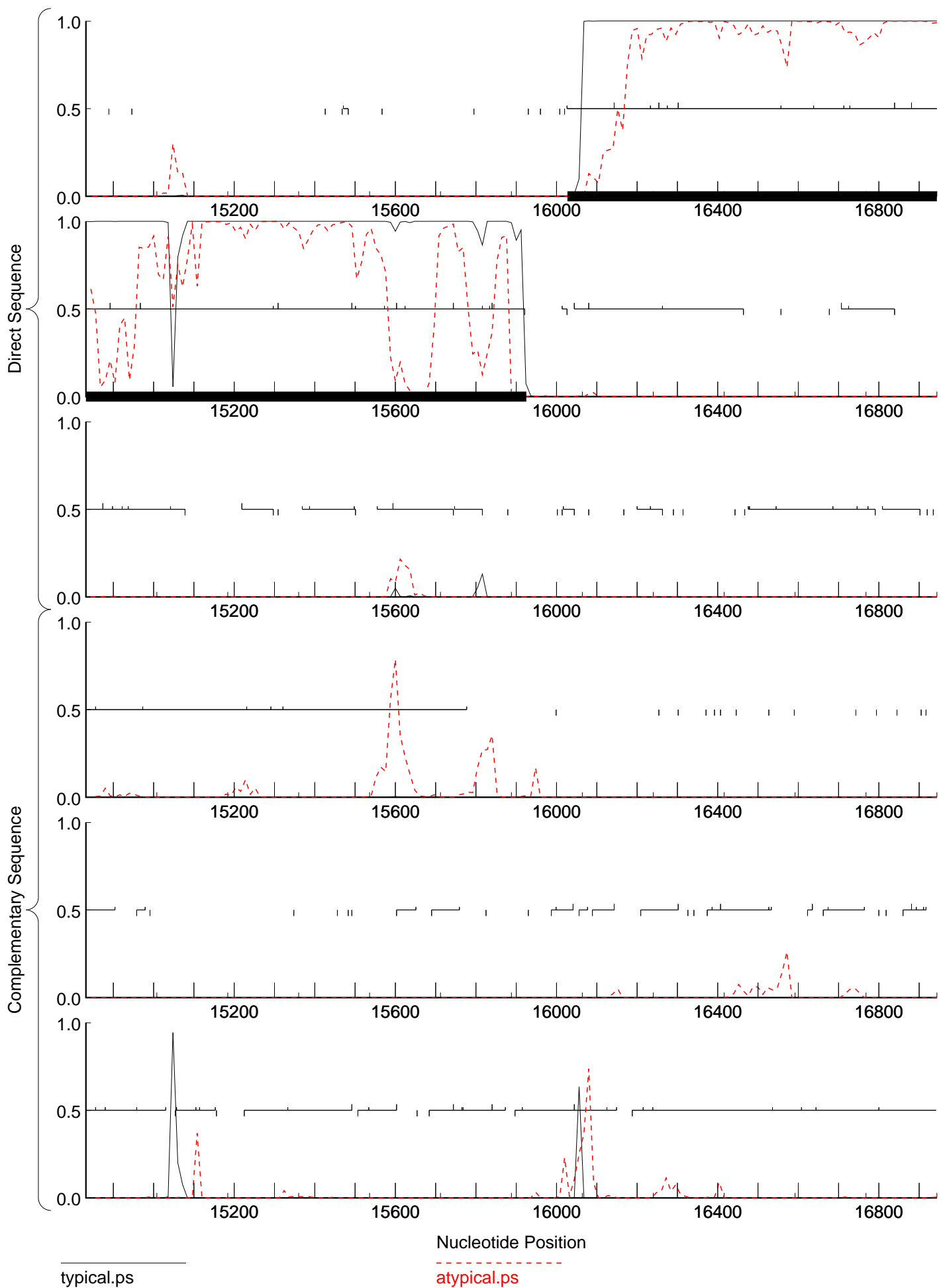
complete sequence, 59818 bp including 11-base 3' overhang (CTCGTAGGCAT), Cluster K1, Order 4, Window 96, Step 12, 5/30
complete sequence, 59818 bp including 11-base 3' overhang (CTCGTAGGCAT), Cluster K1, Order 2, Window 96, Step 12, 5/30

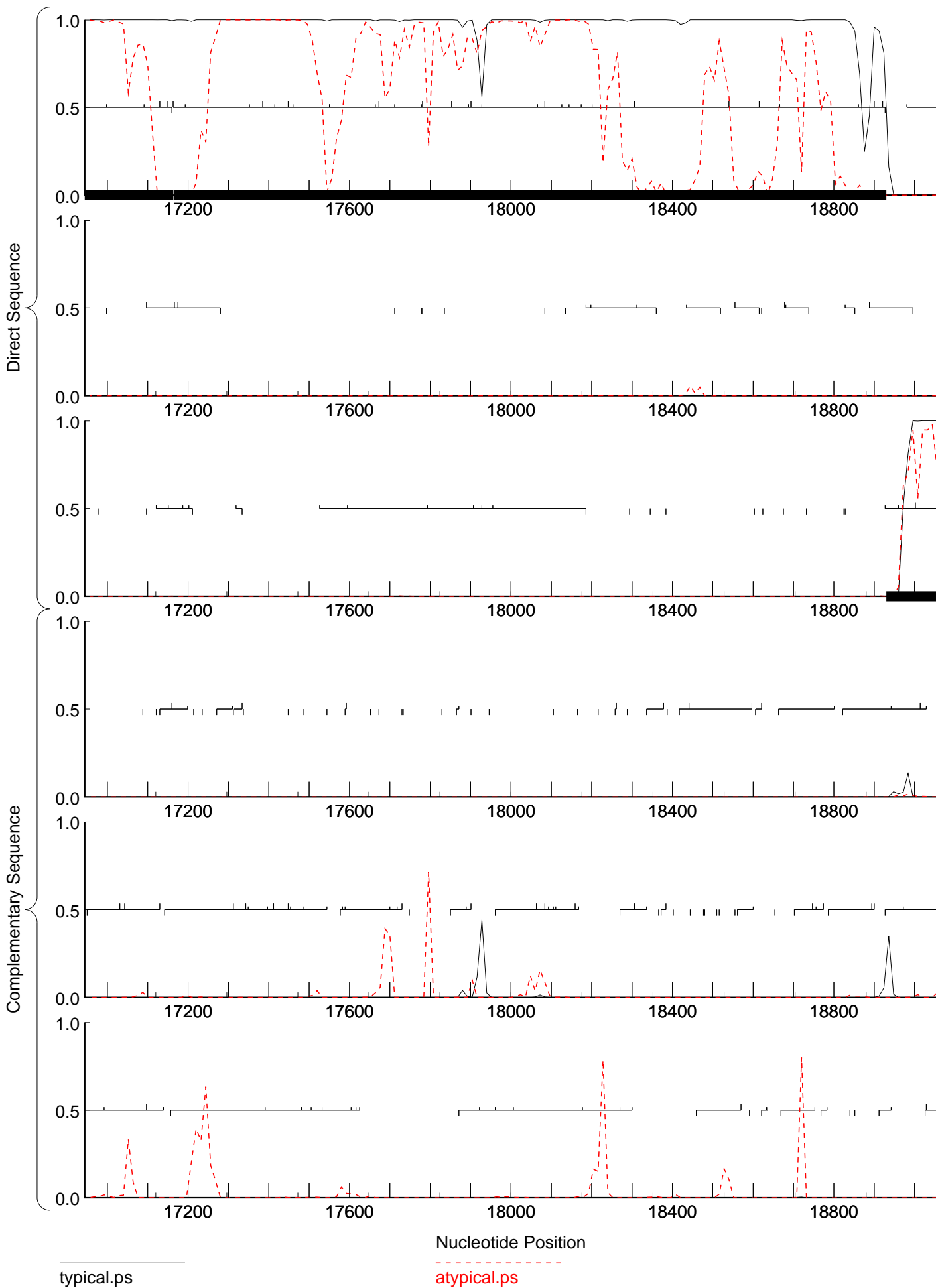






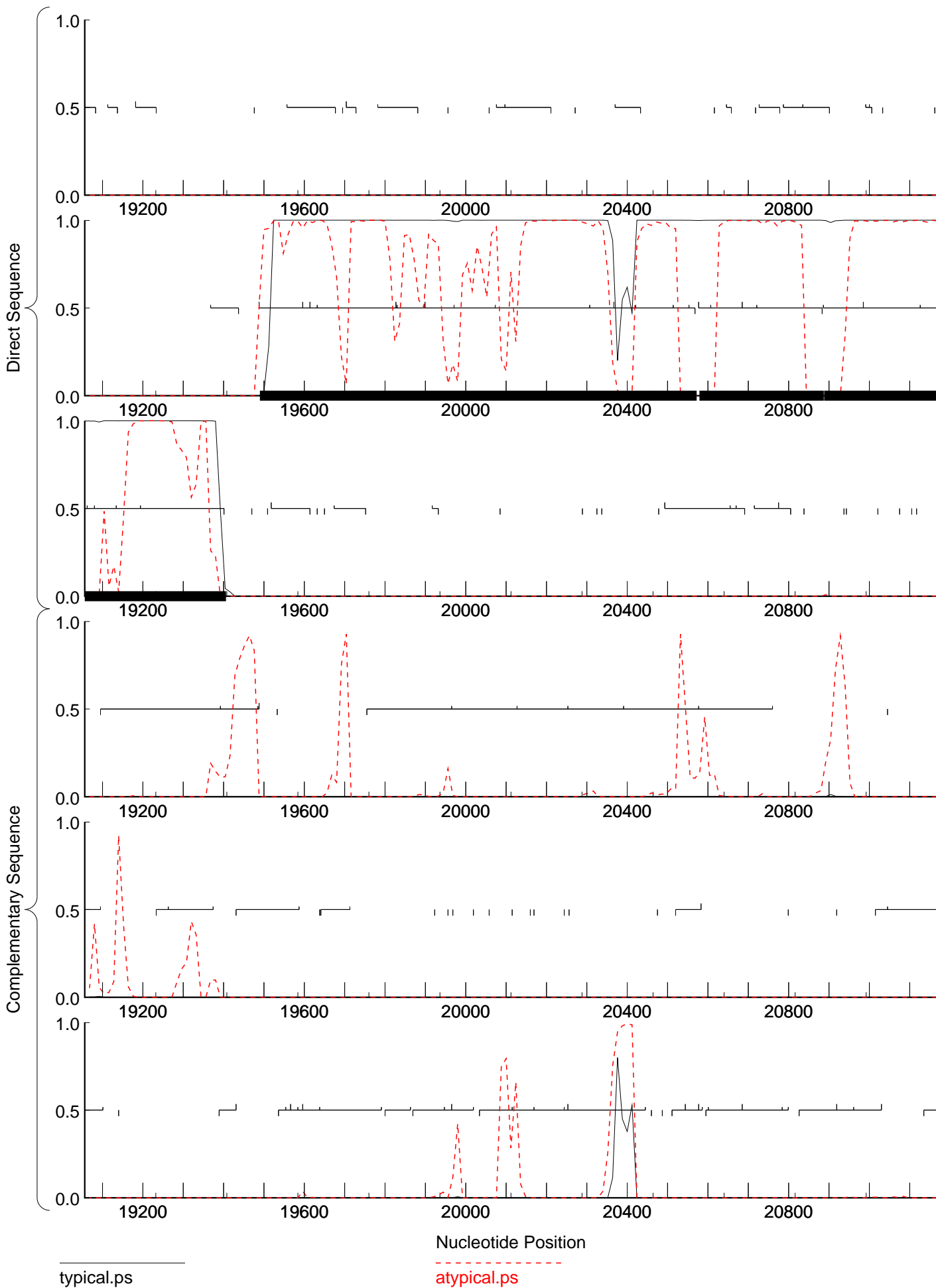


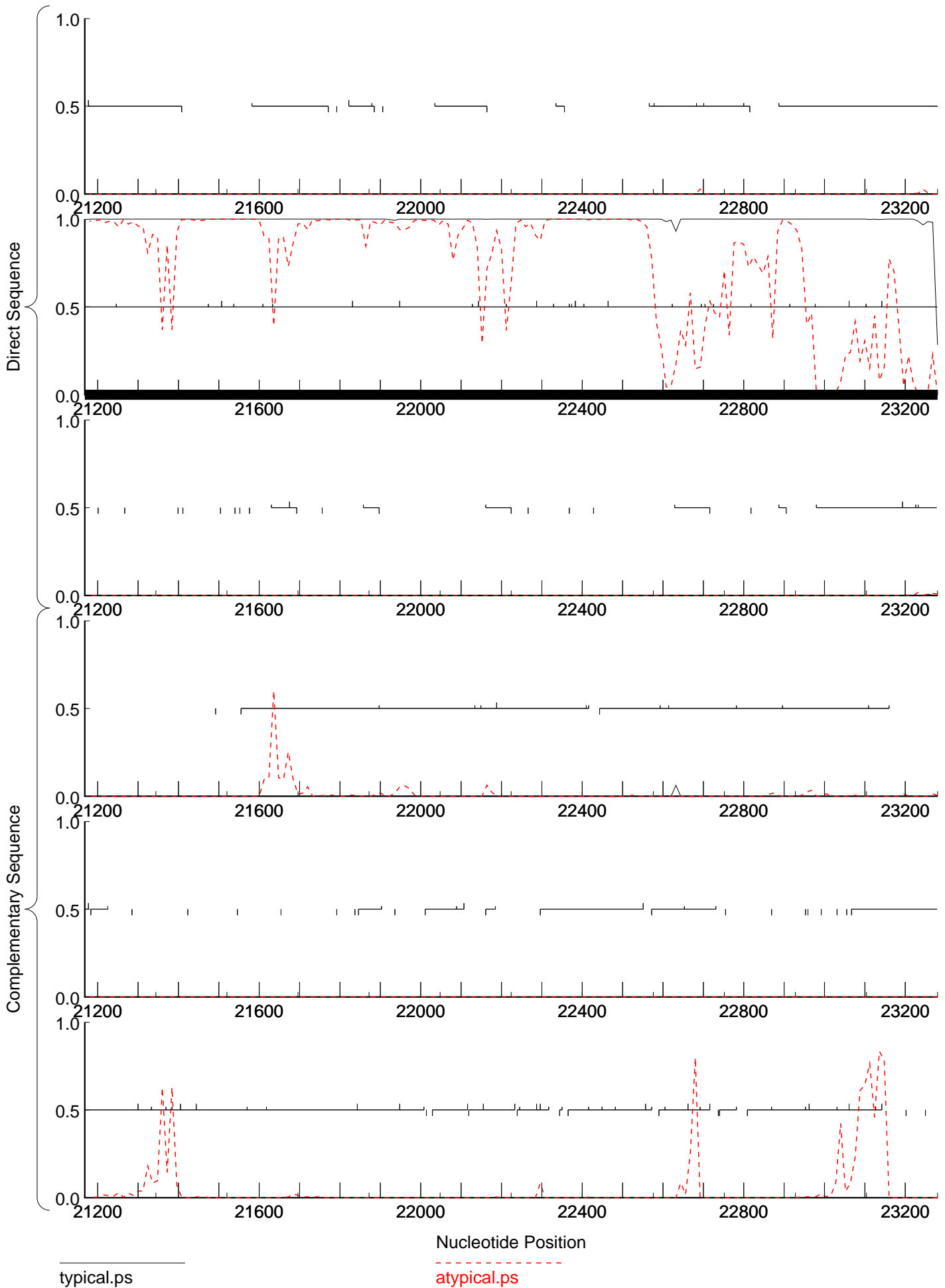


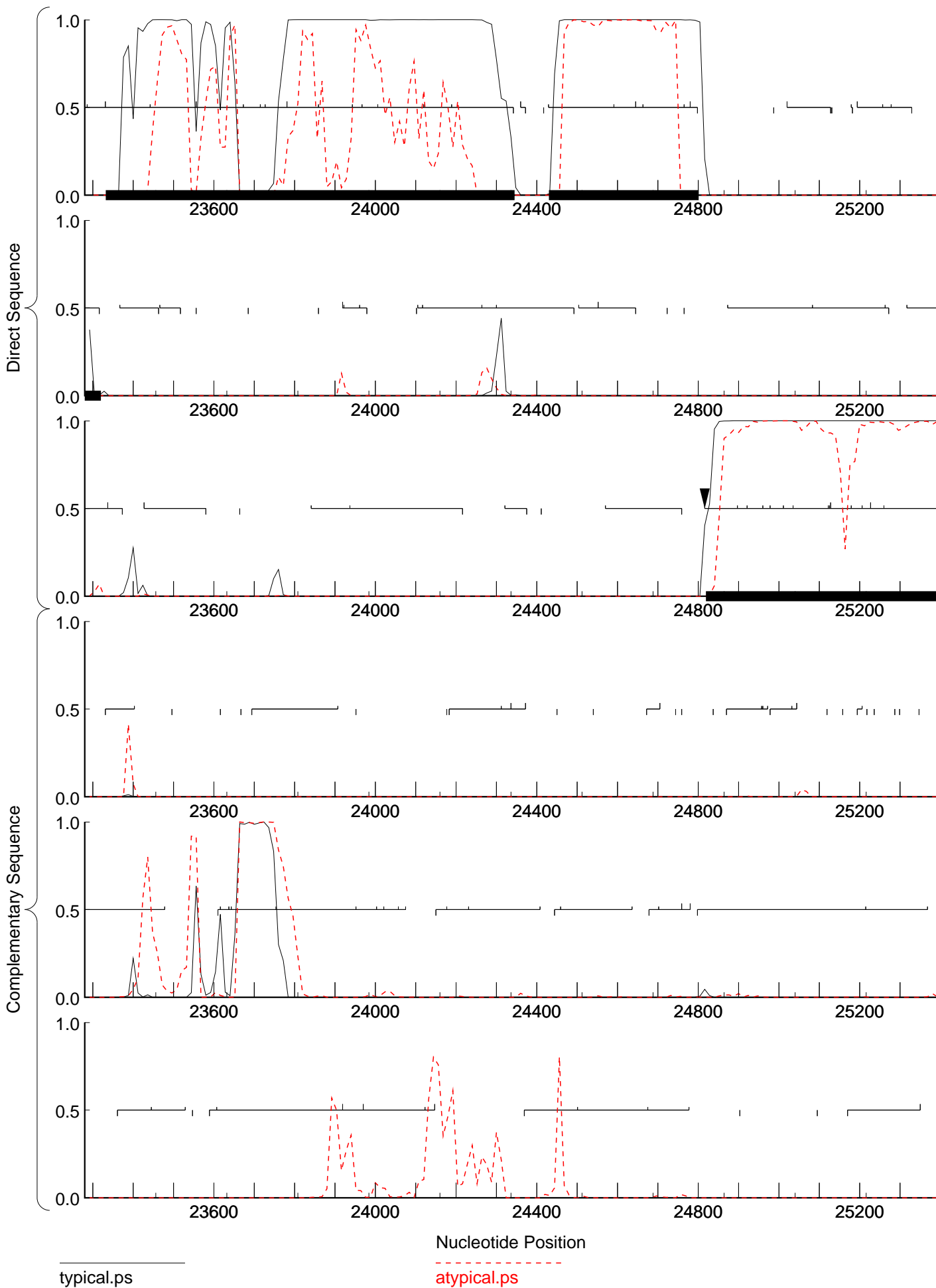


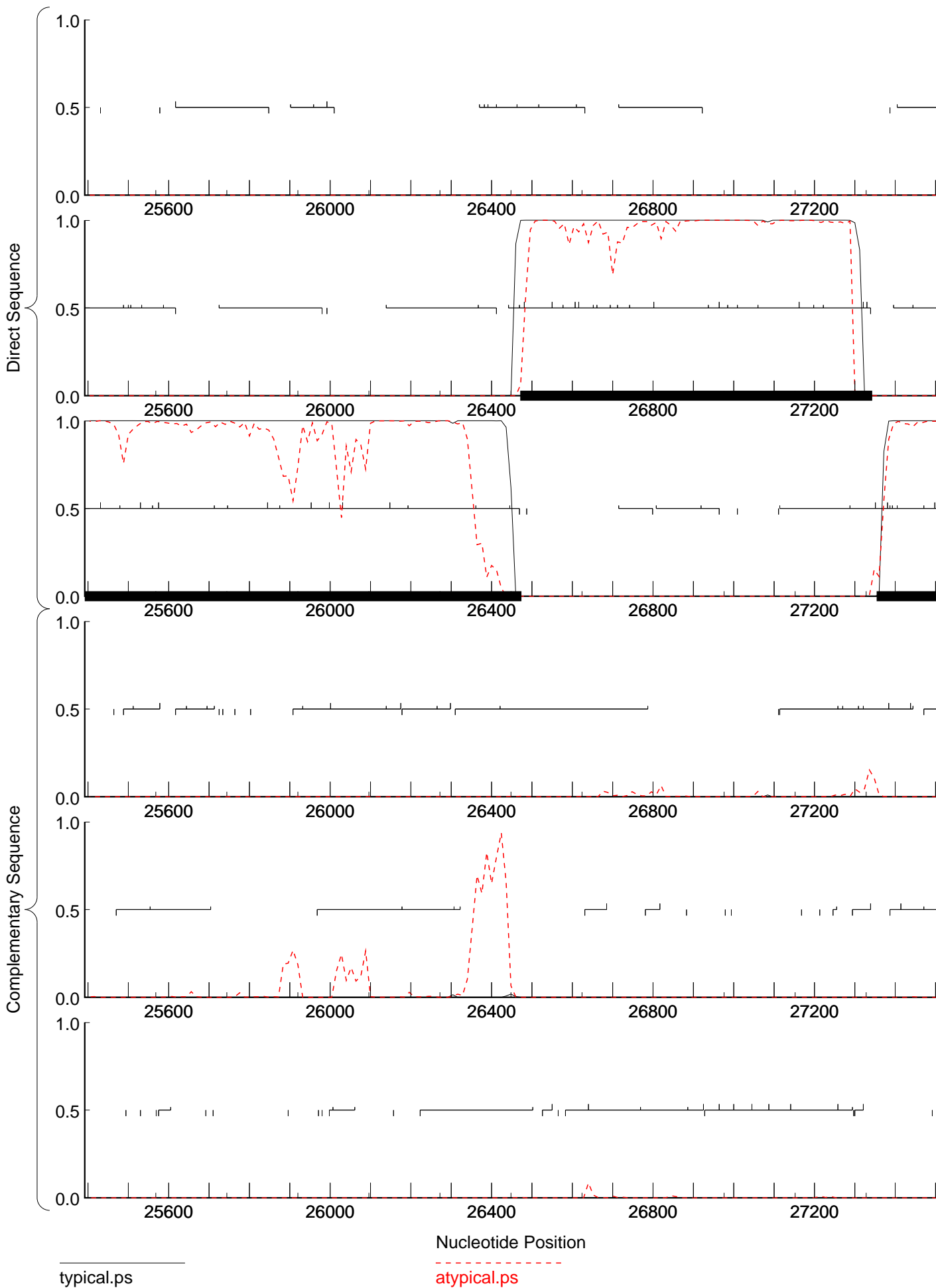
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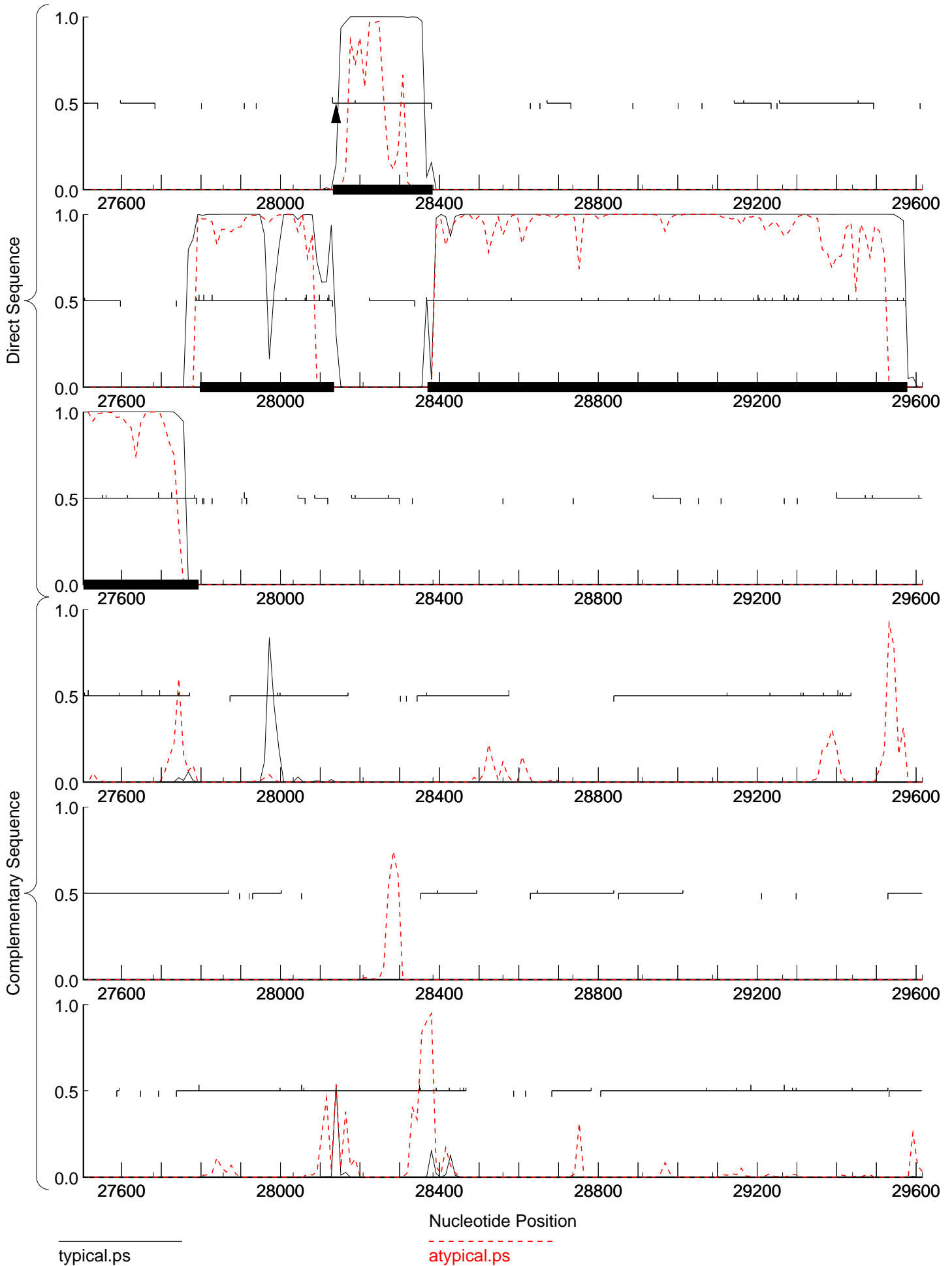
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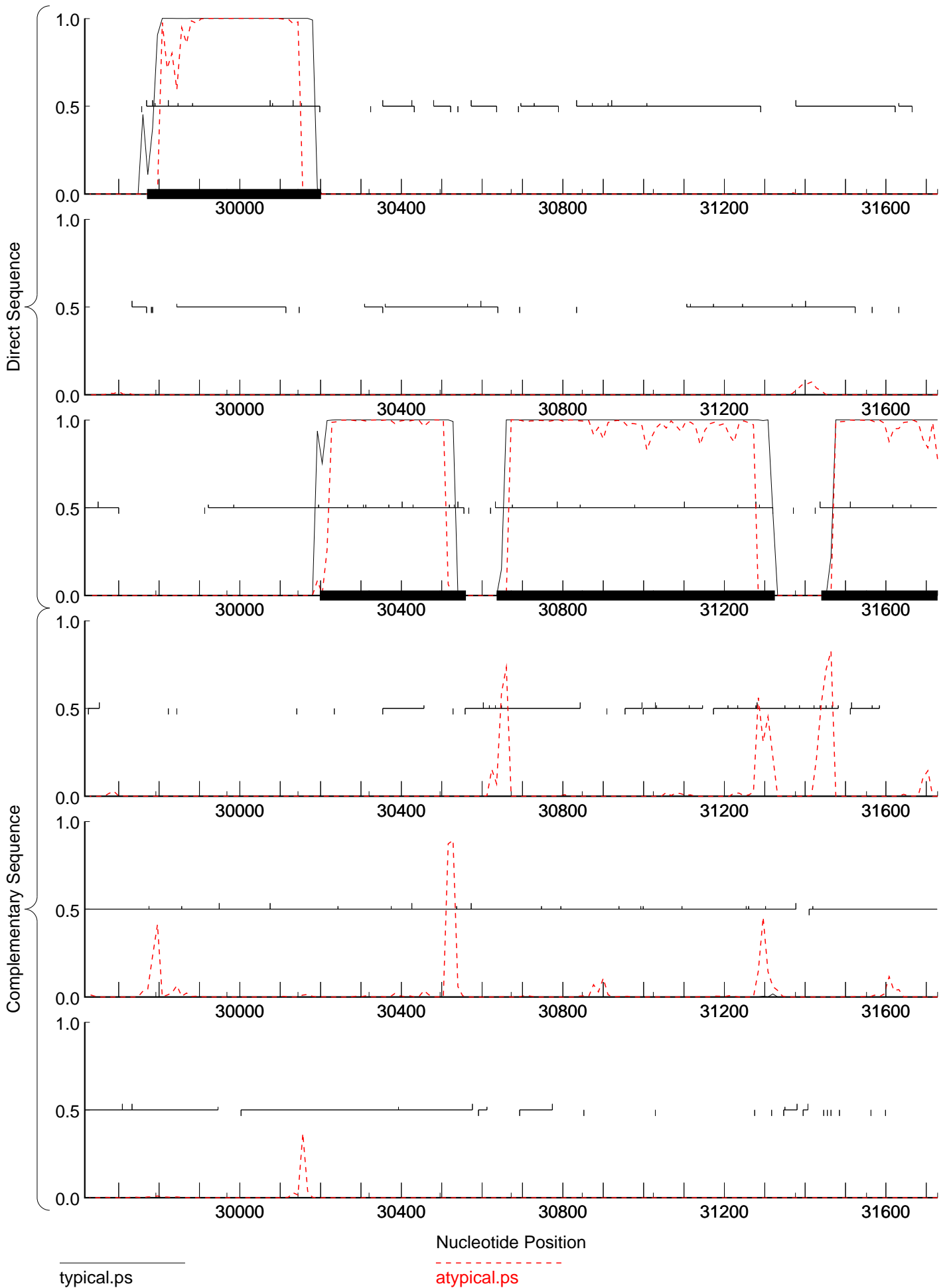


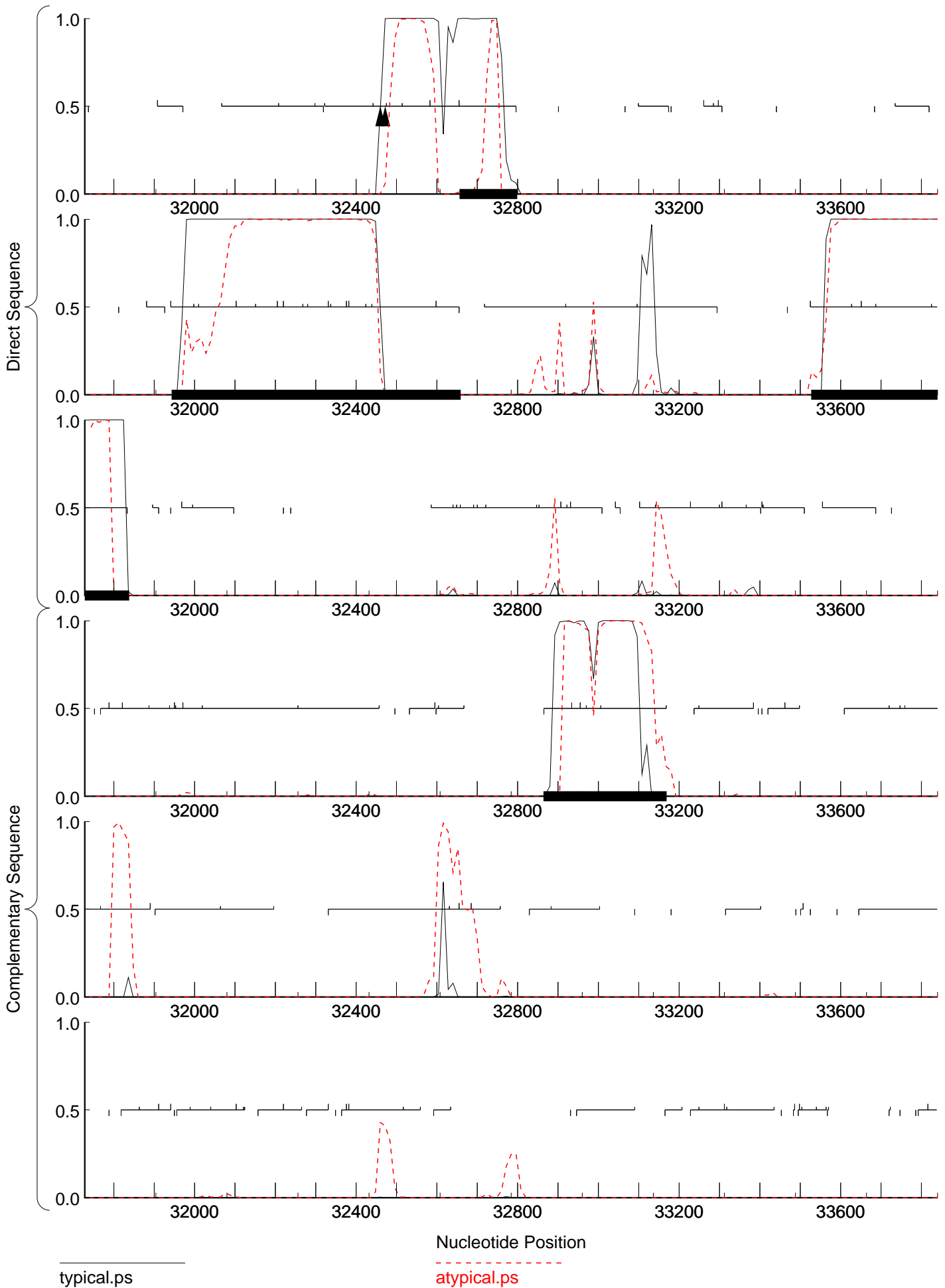


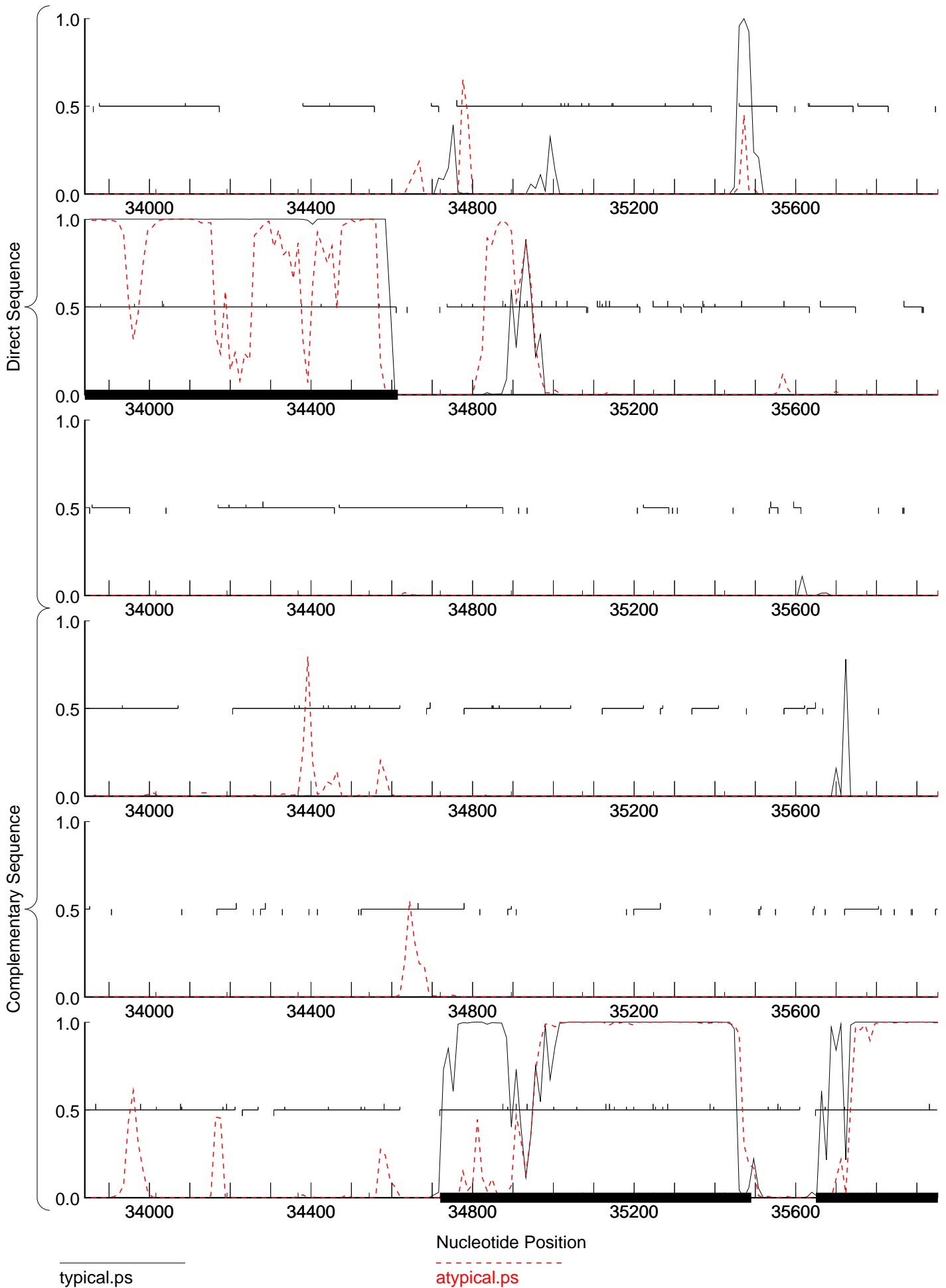


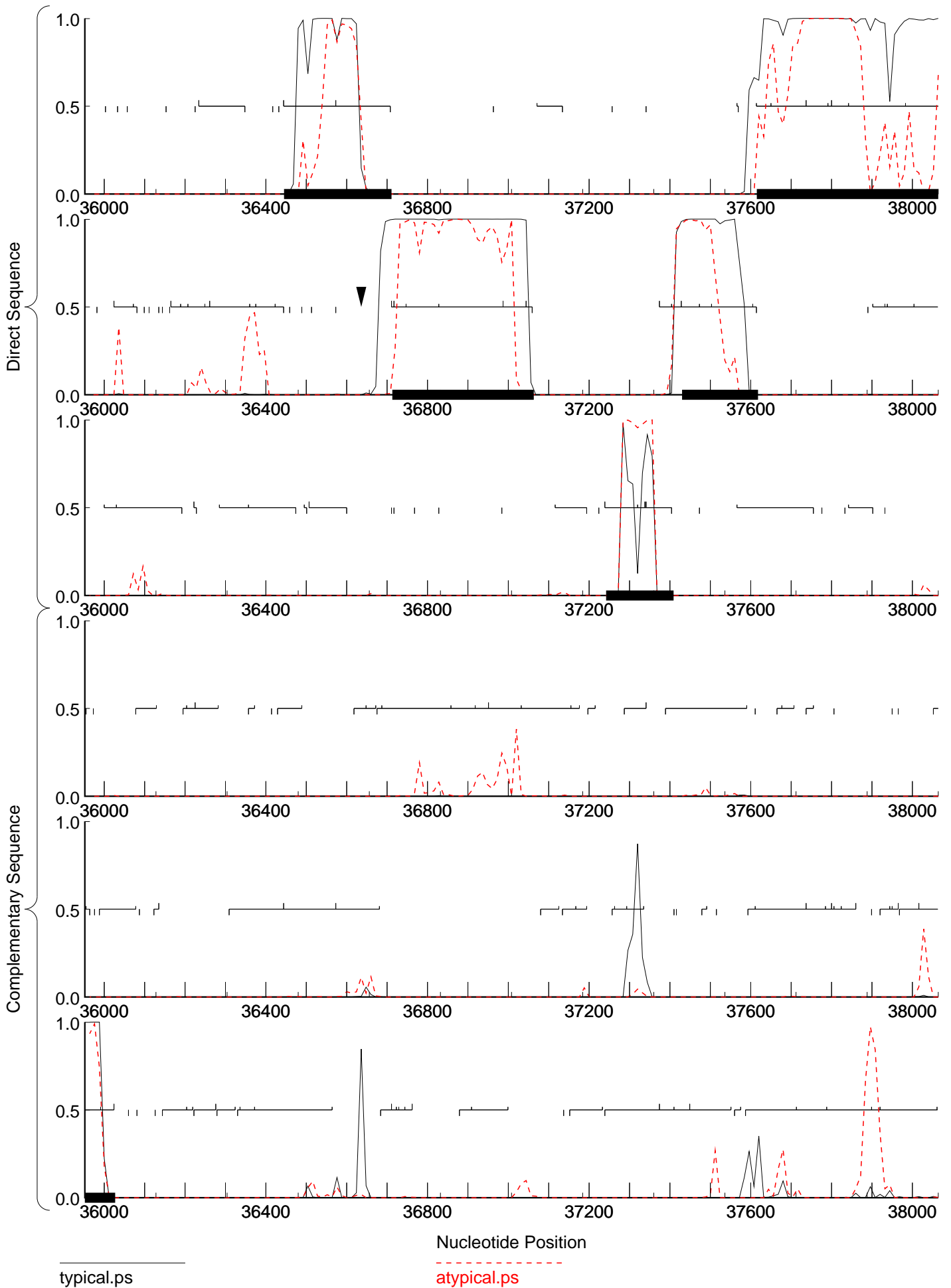
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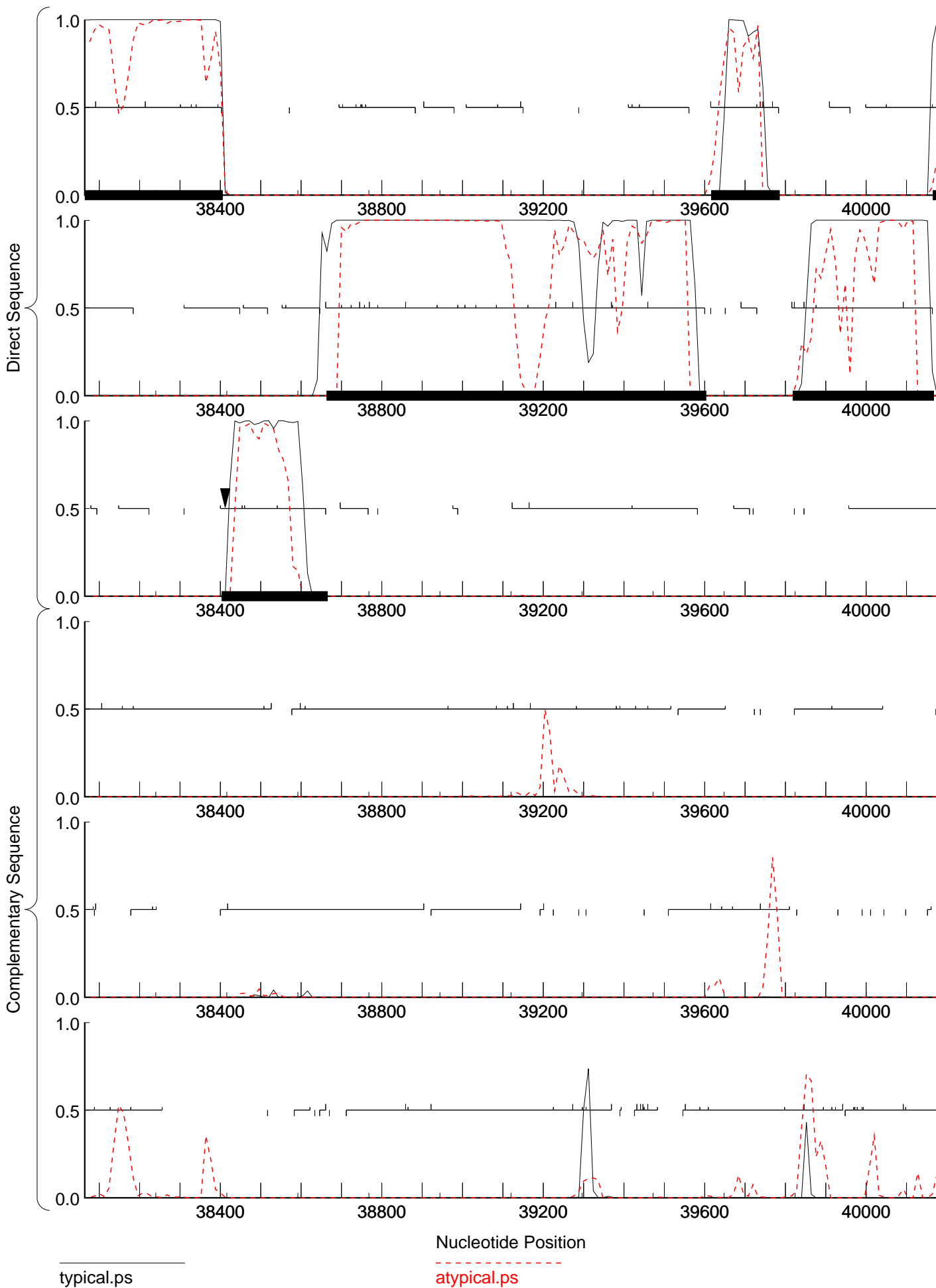


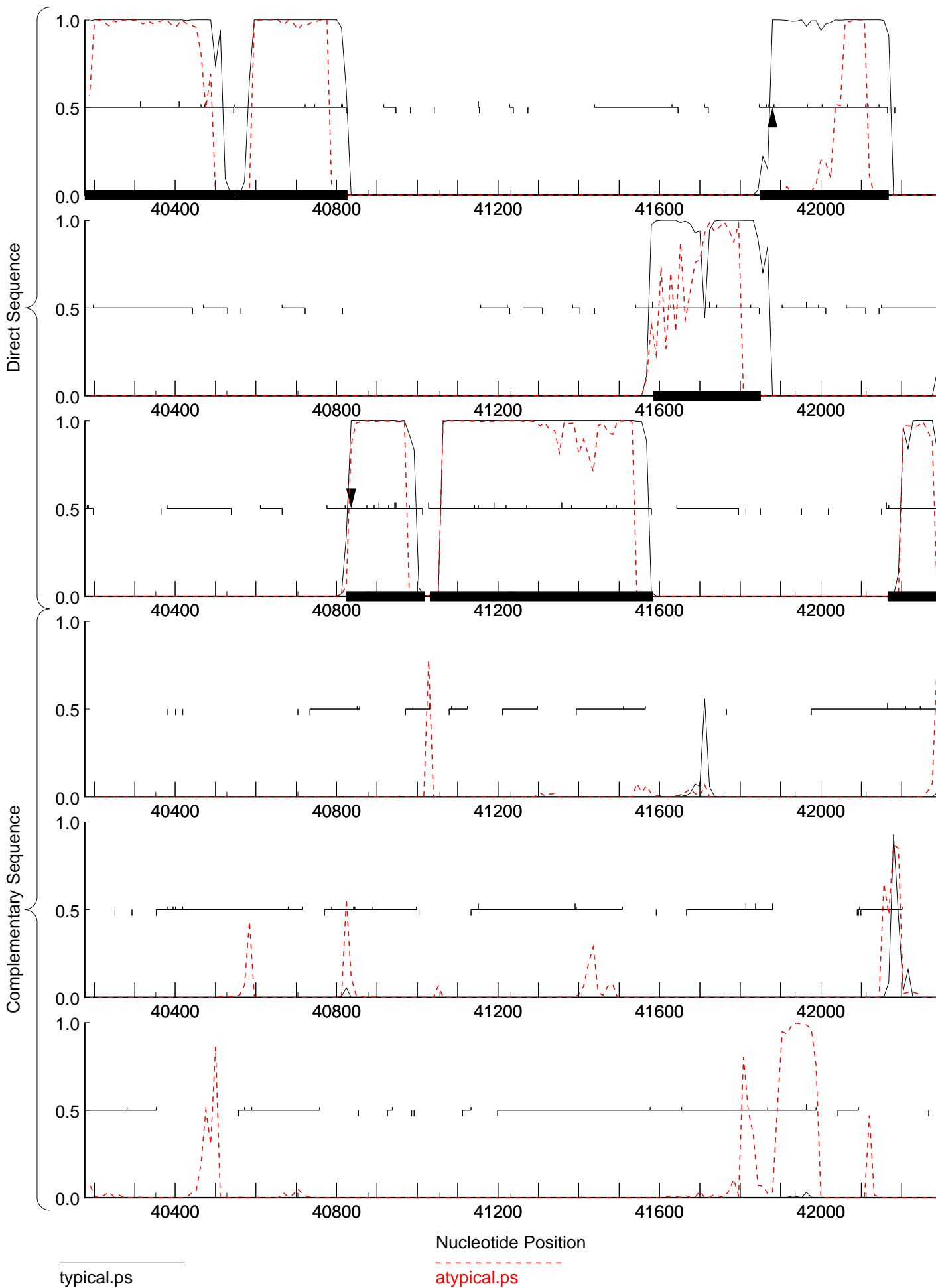


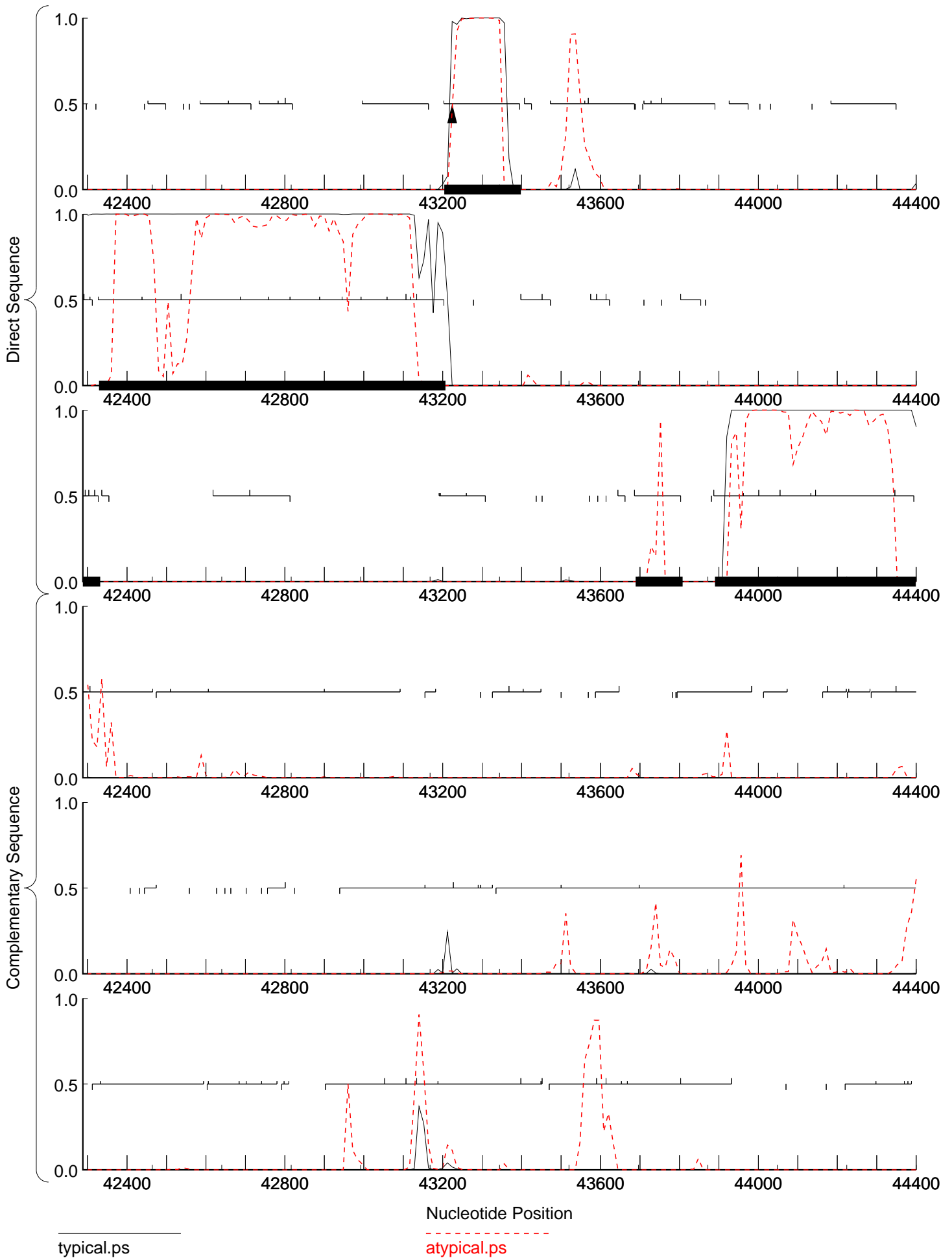


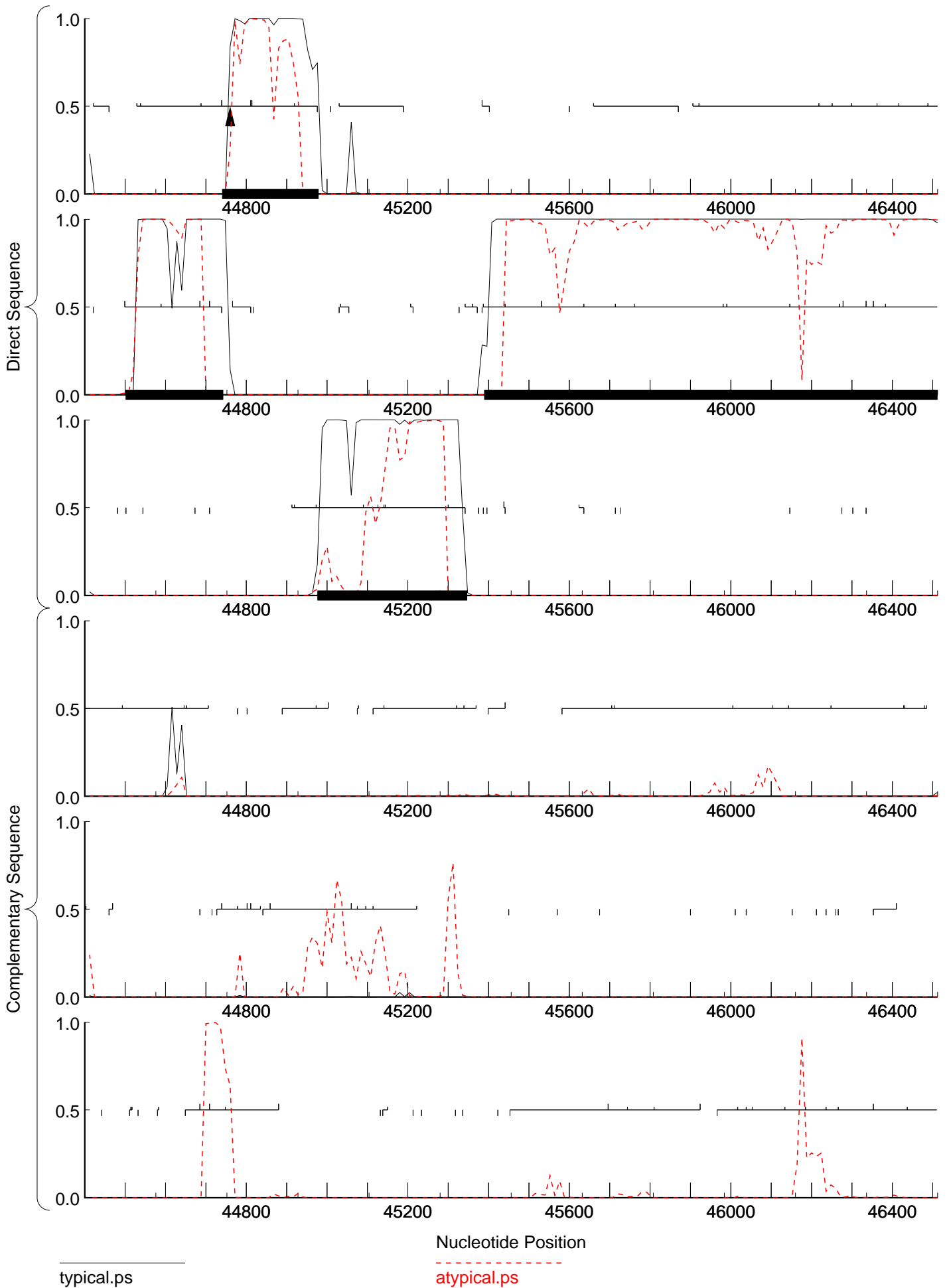
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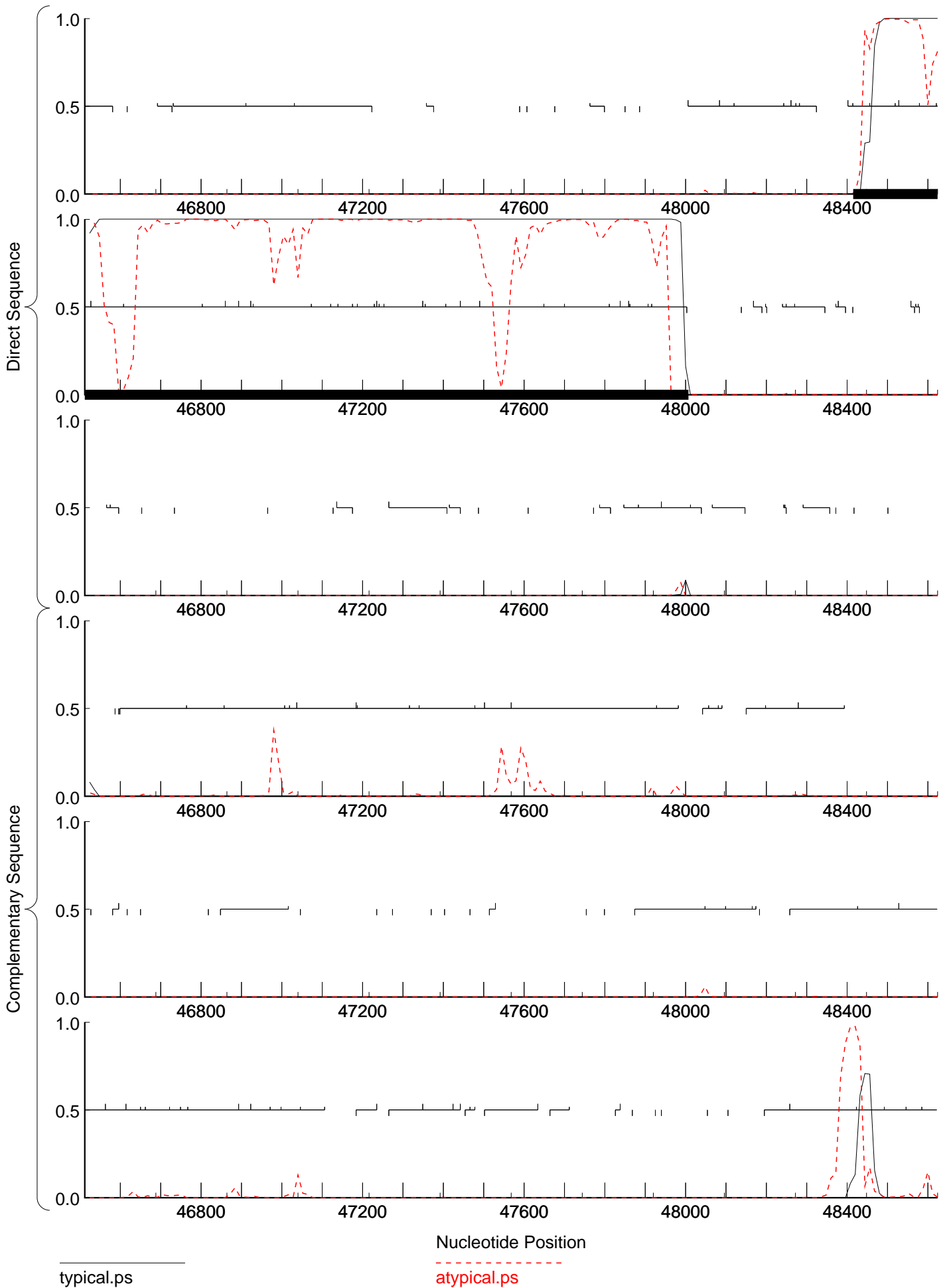
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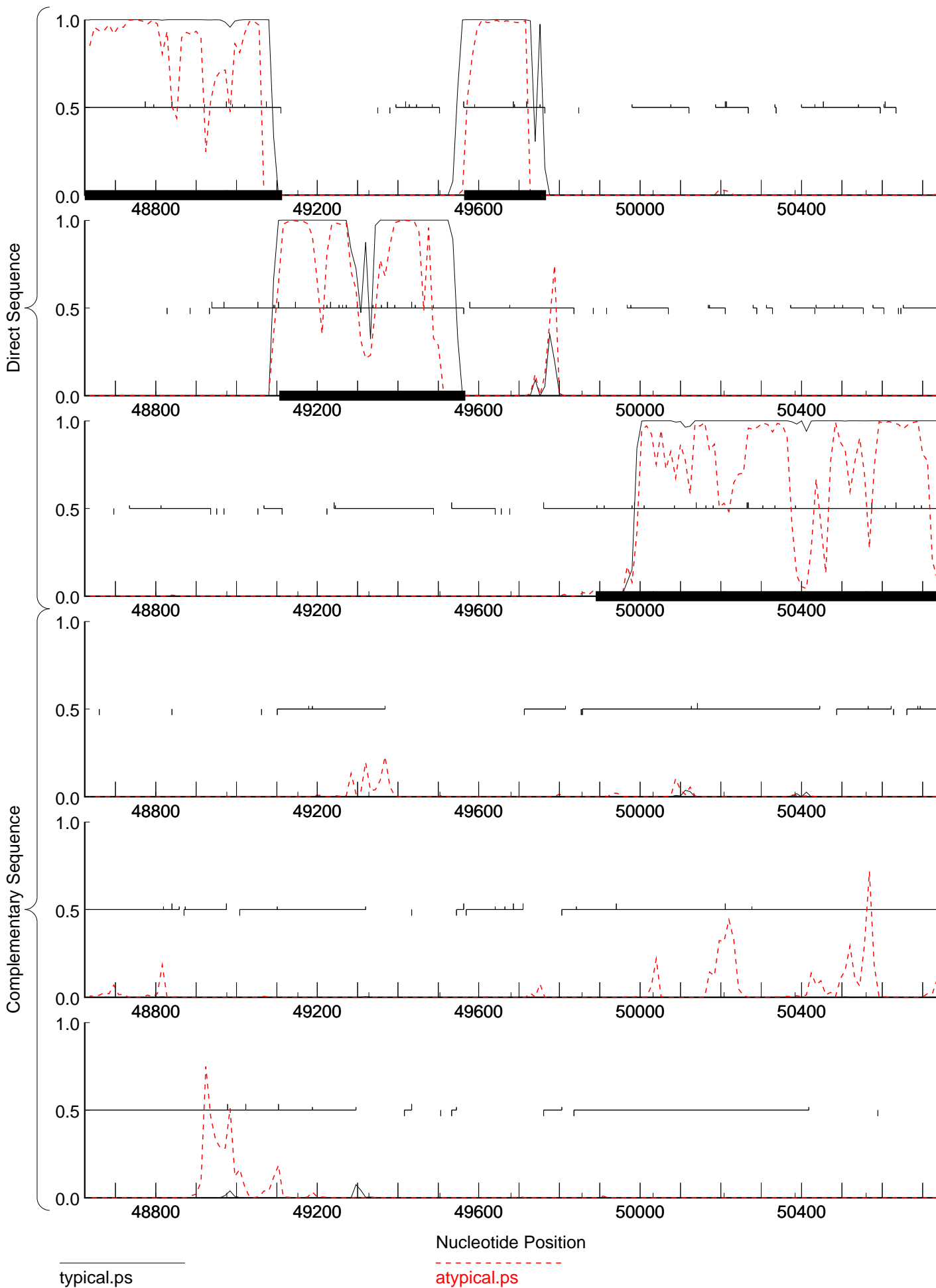






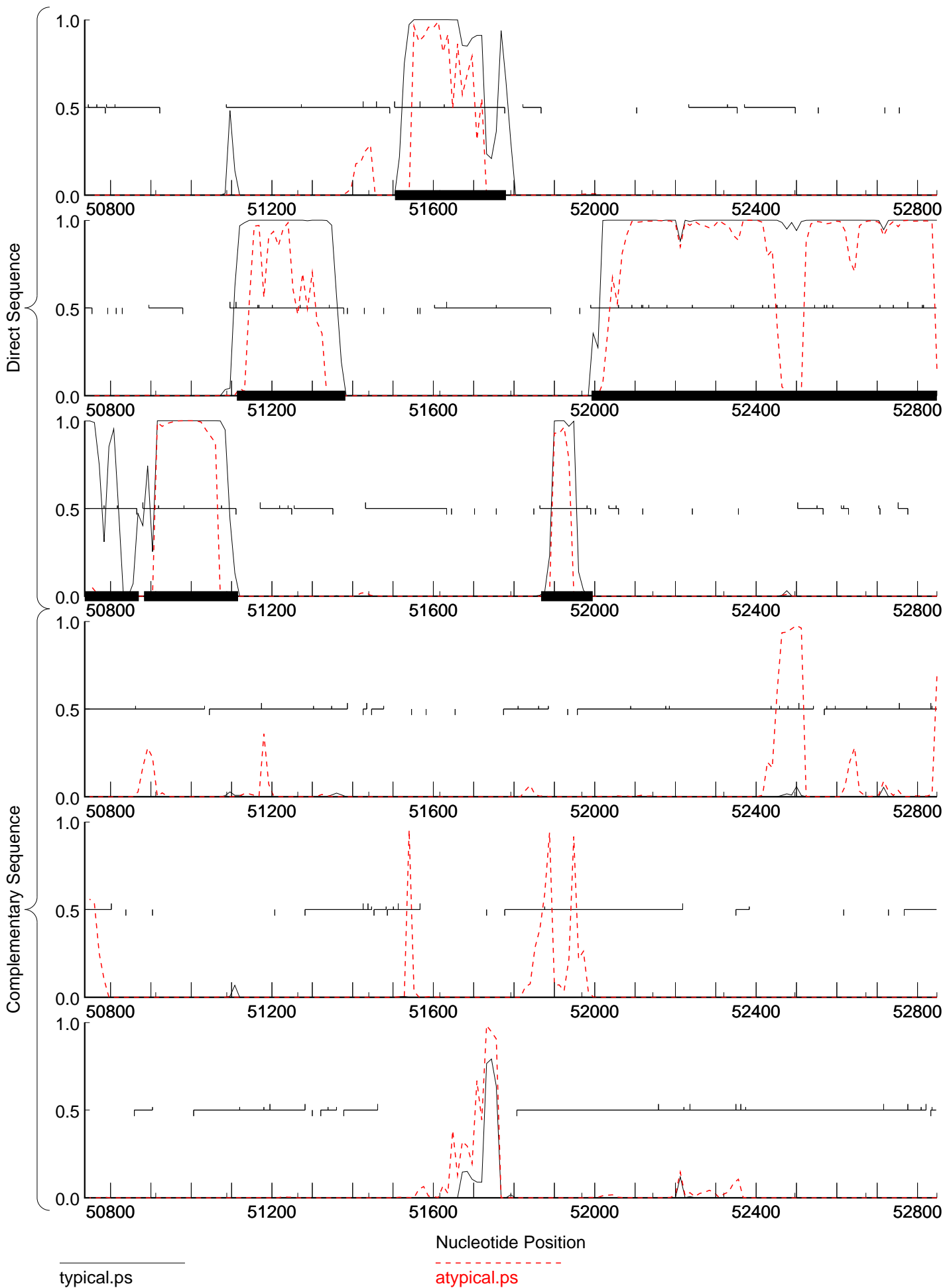






typical.ps

atypical.ps



typical.ps

atypical.ps

