

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 : 1/16/18 at 6:25:13
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,

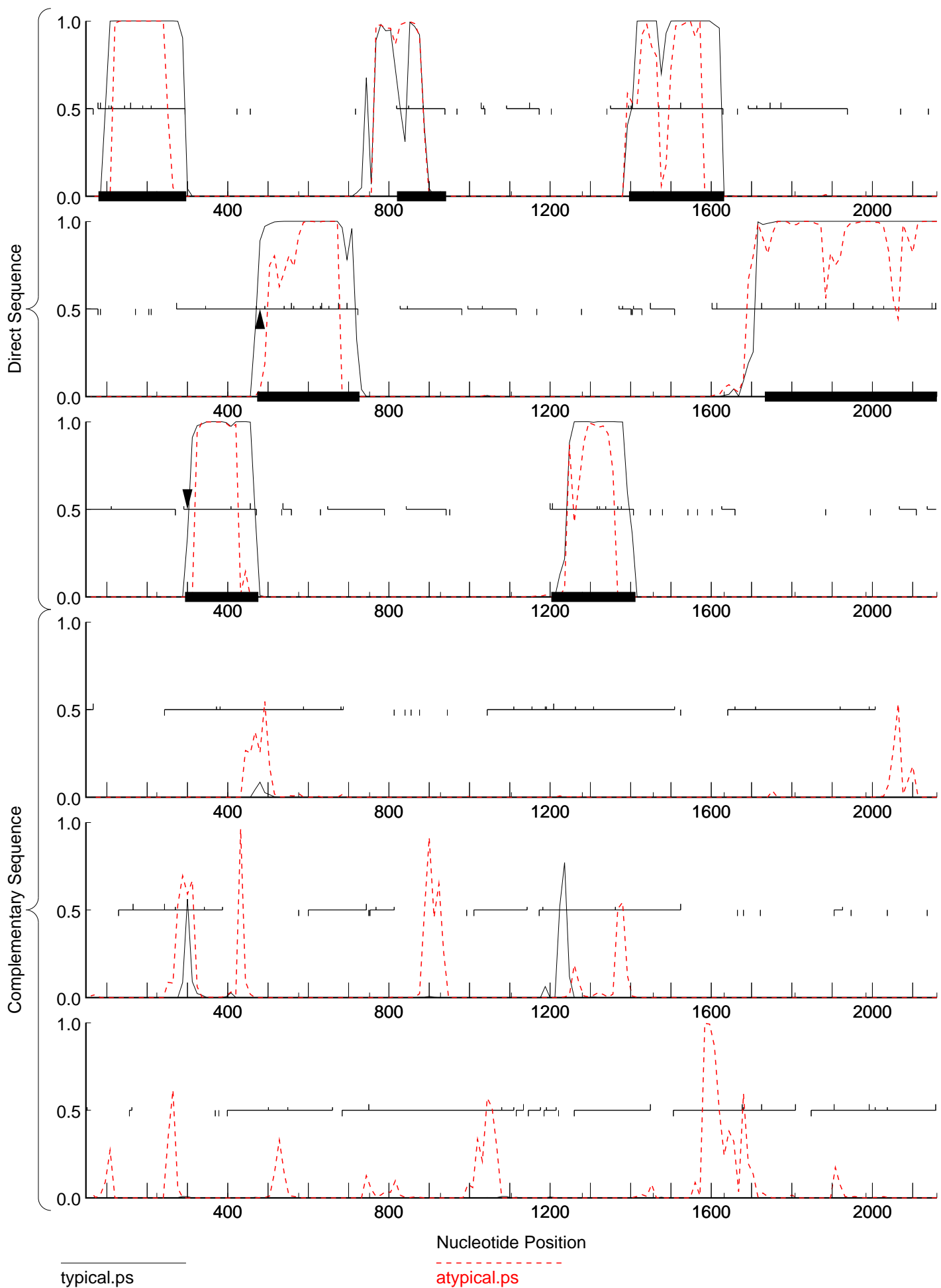
Matrix notes & comments

Training set derived by GeneMarkS, 4.30 October 2014
Tue Jan 16 06:25:13 2018

MATRIX INFORMATION

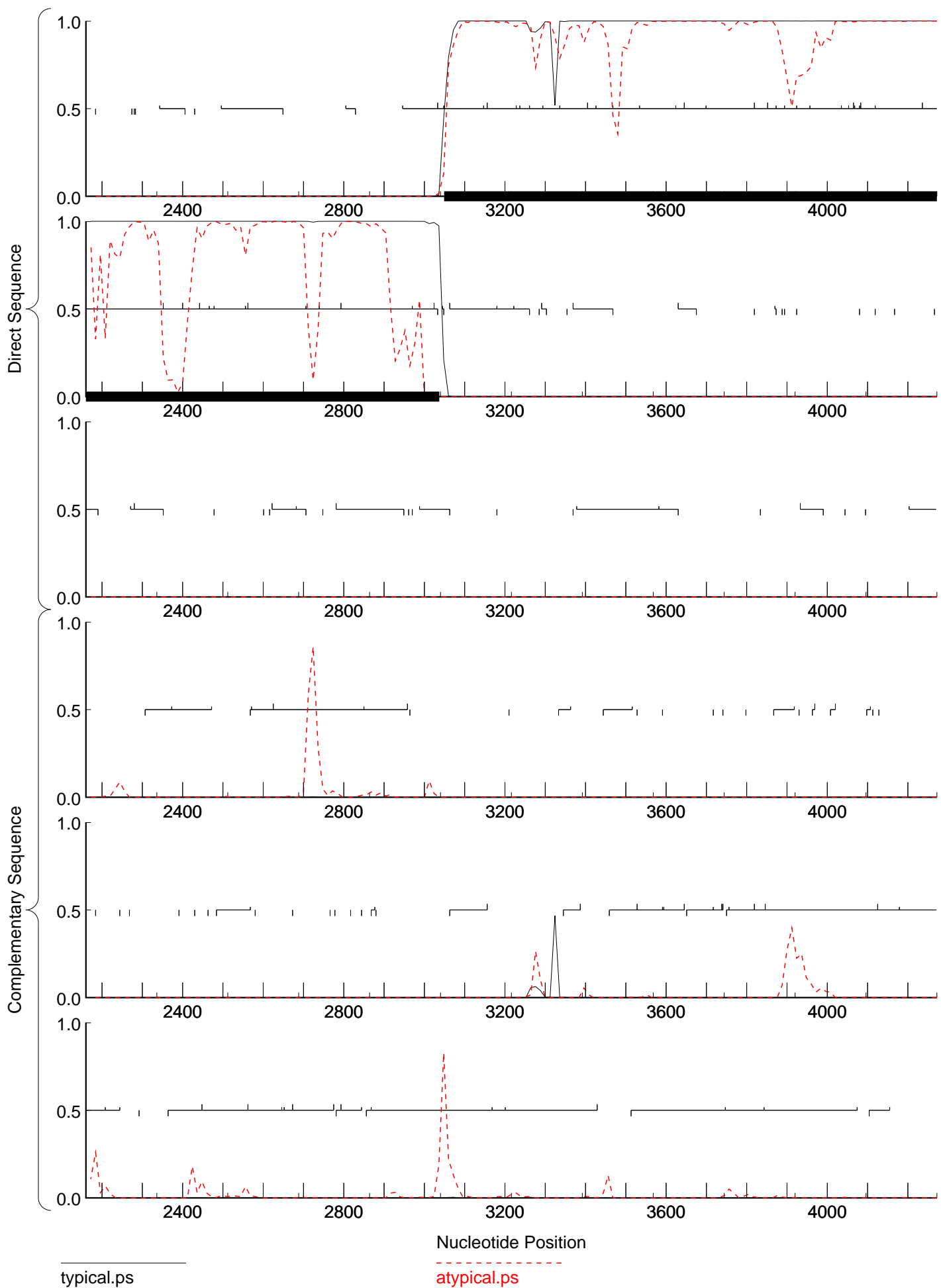
Matrix : -
Author : -
Order : 4

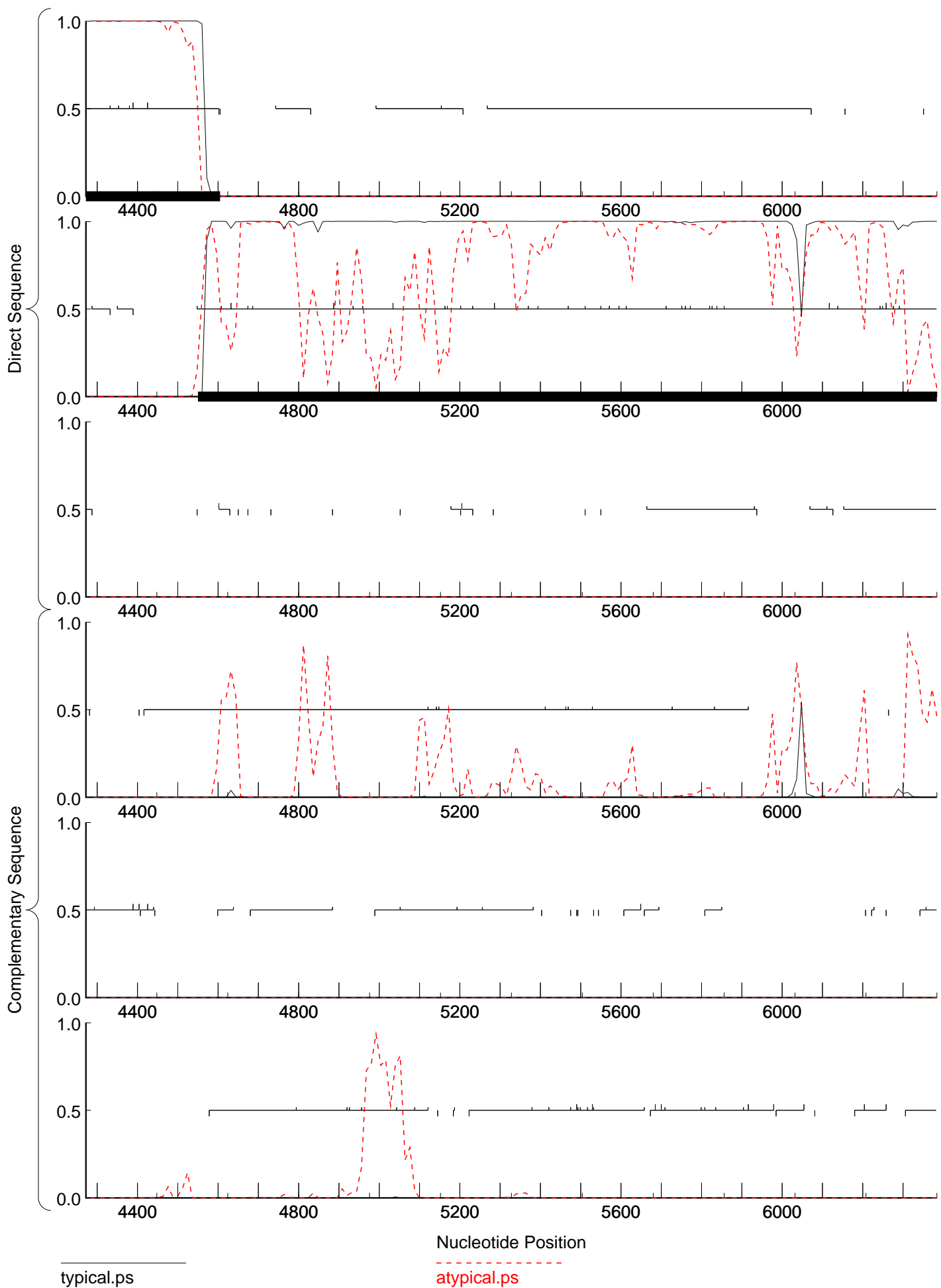
Send questions / comments to:
Dr. M. Borodovsky
Georgia Institute of Technology
School of Biology
Atlanta, GA 30332-0230



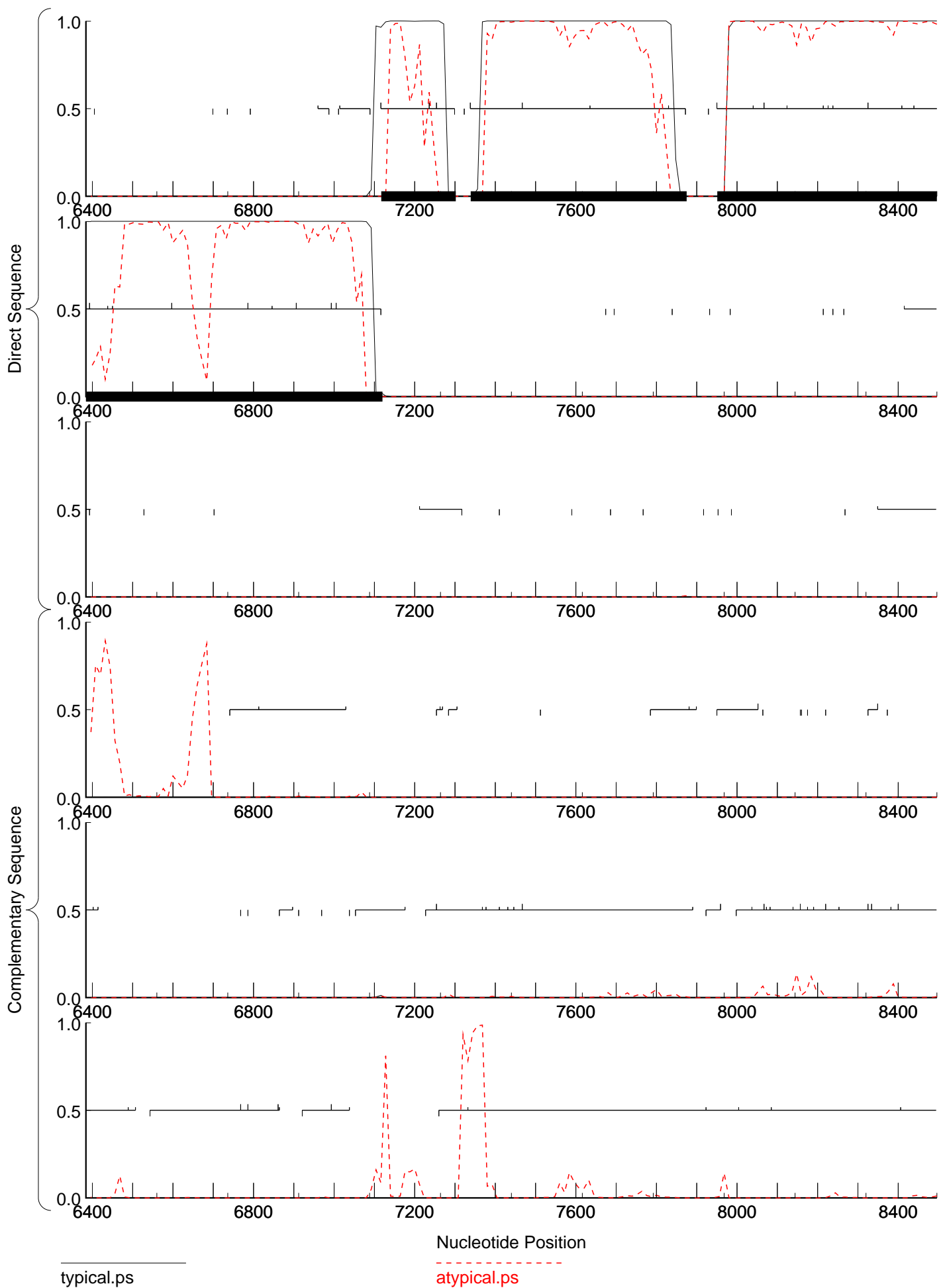
typical.ps

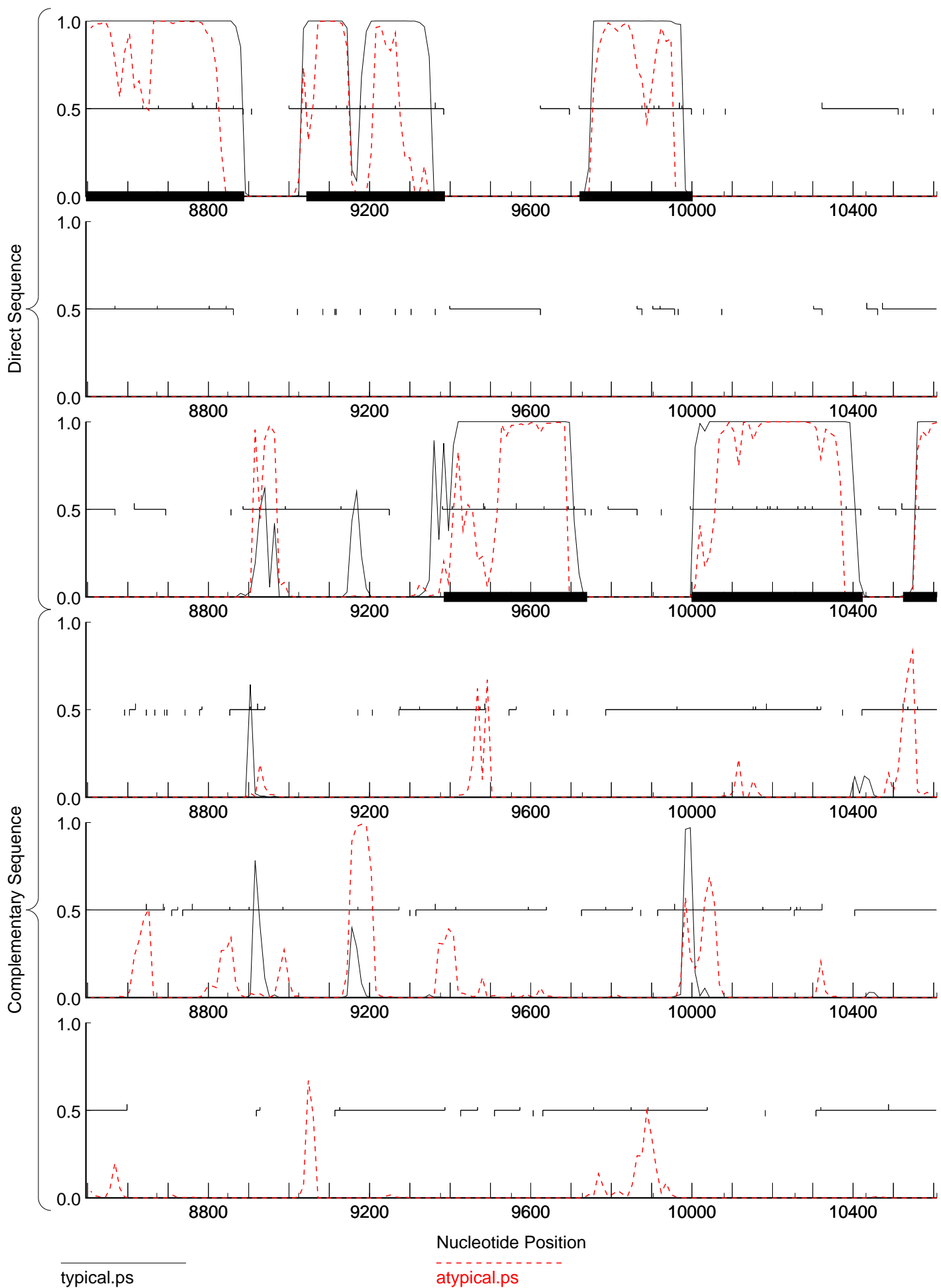
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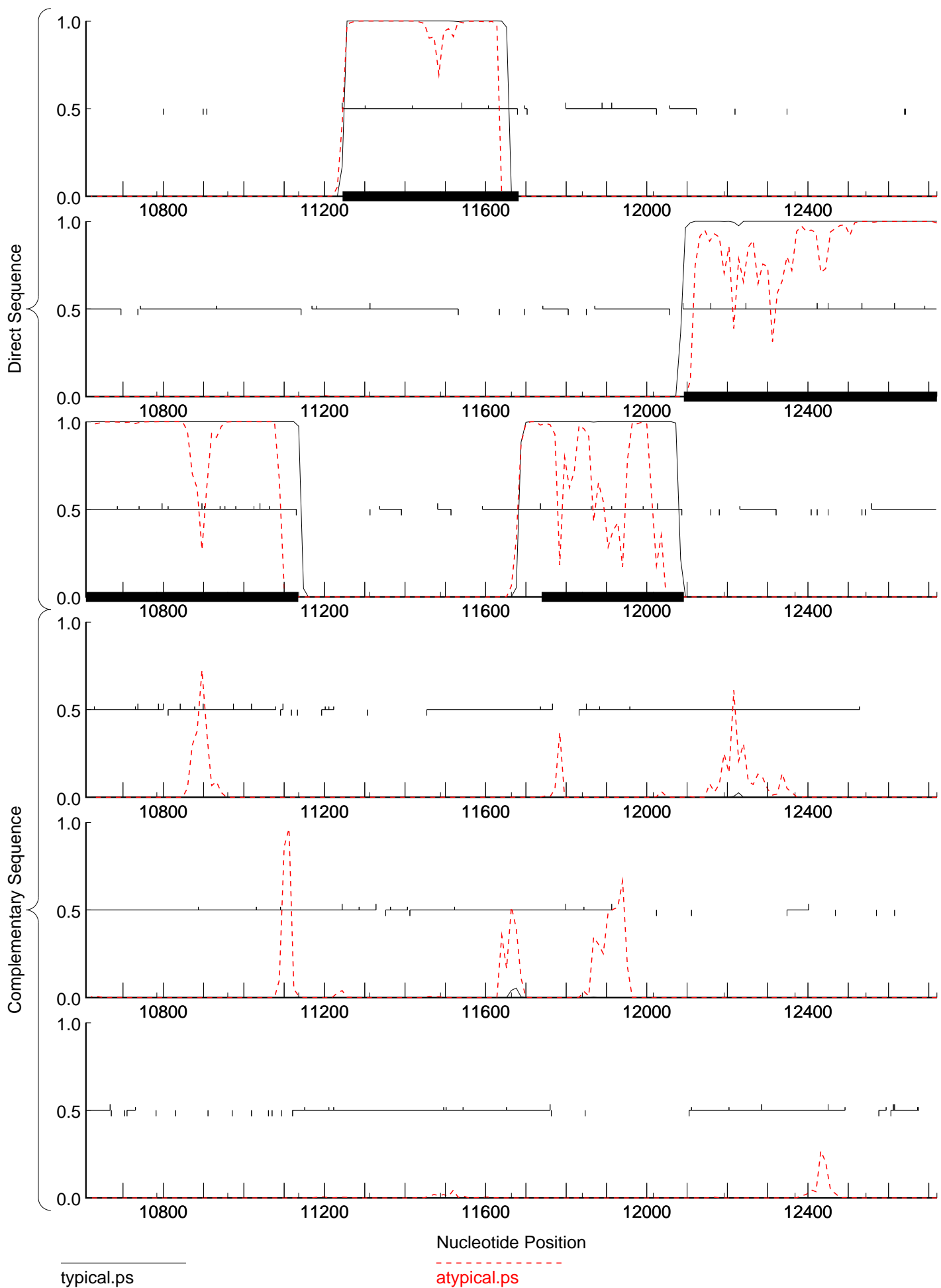


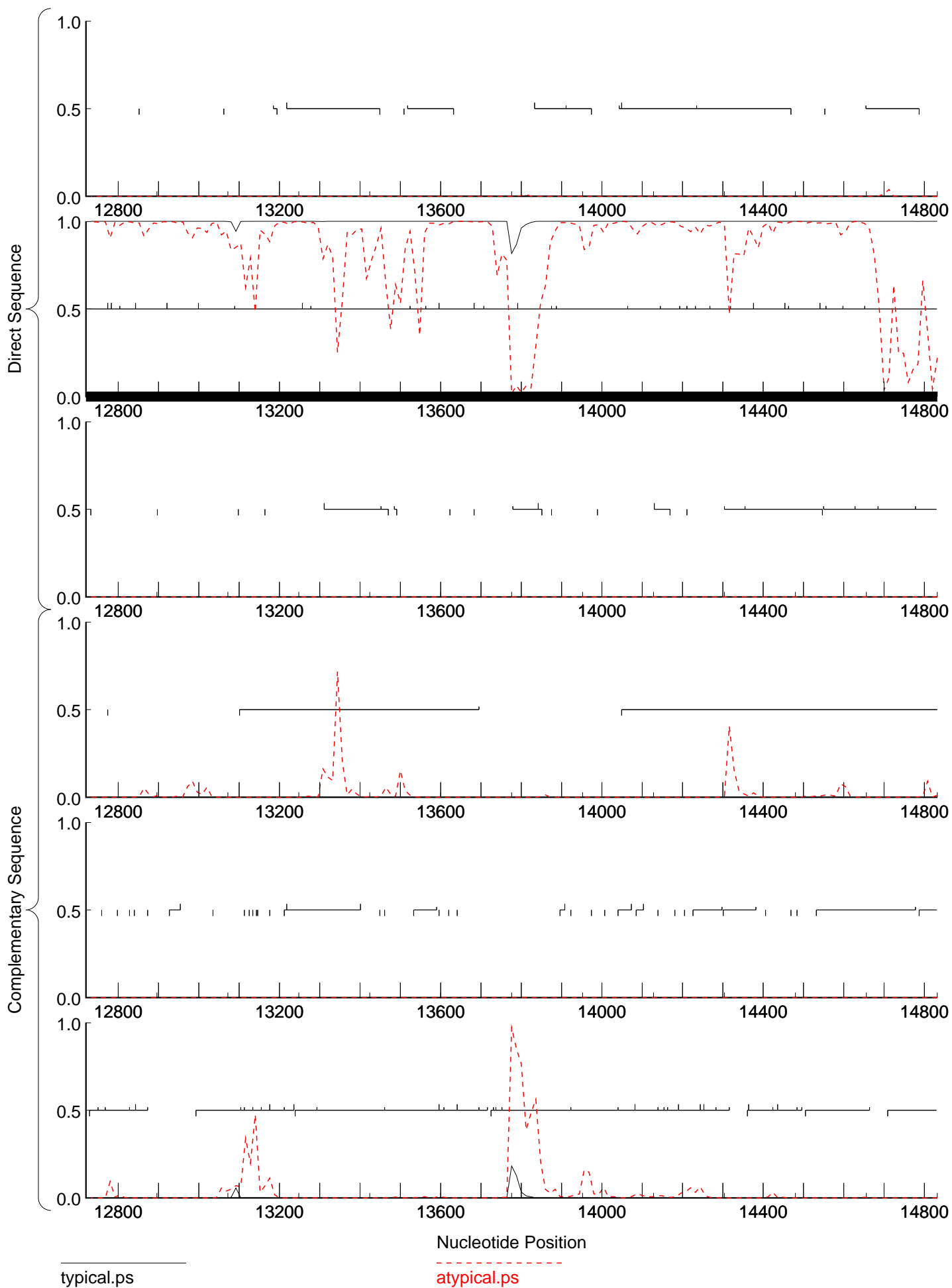


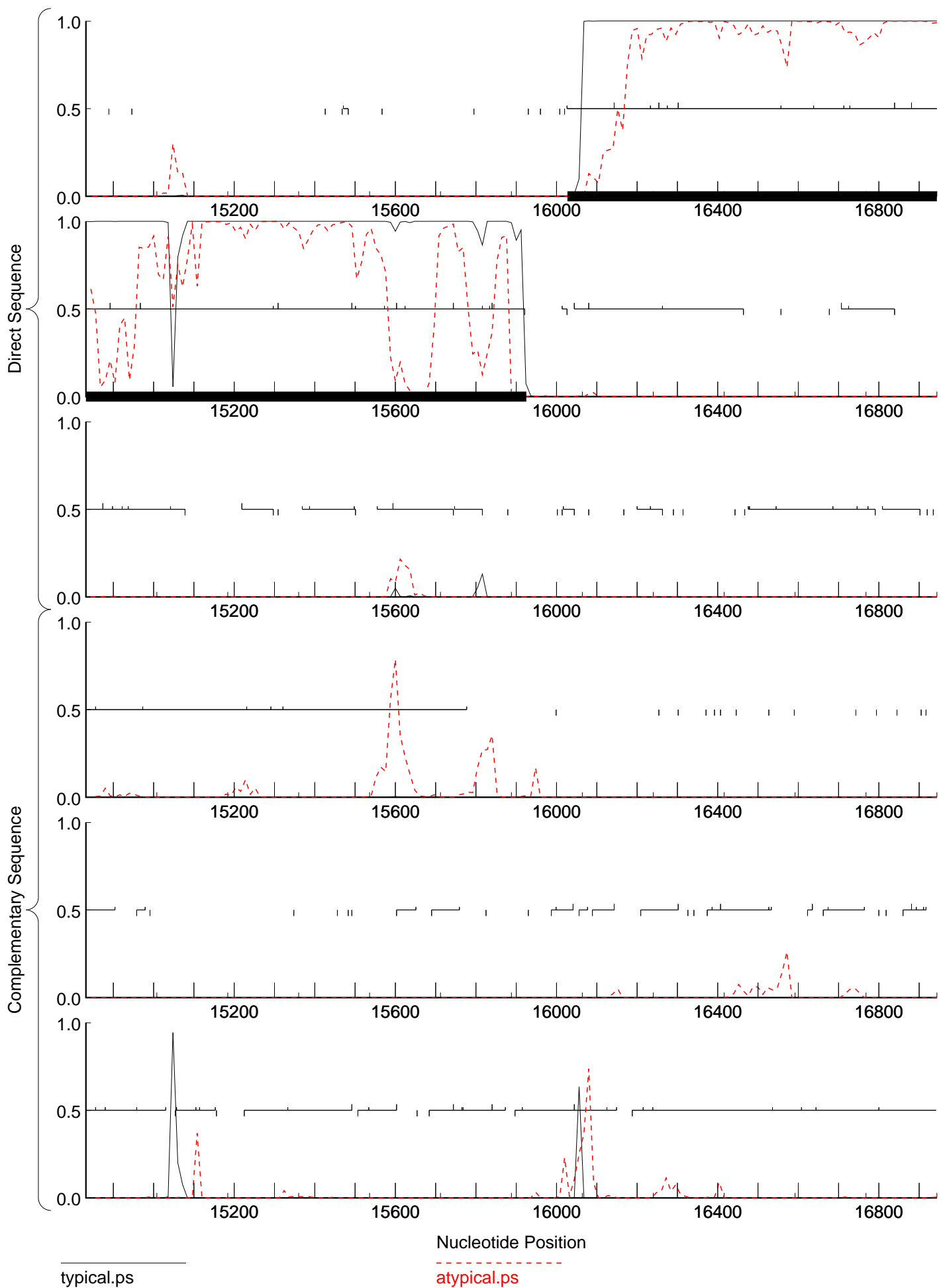
complete sequence, 59818 bp including 11-base 3' overhang (CTCGTAGGCAT), Cluster K1, Order 4, Window 96, Step 12, 5/30
complete sequence, 59818 bp including 11-base 3' overhang (CTCGTAGGCAT), Cluster K1, Order 2, Window 96, Step 12, 5/30

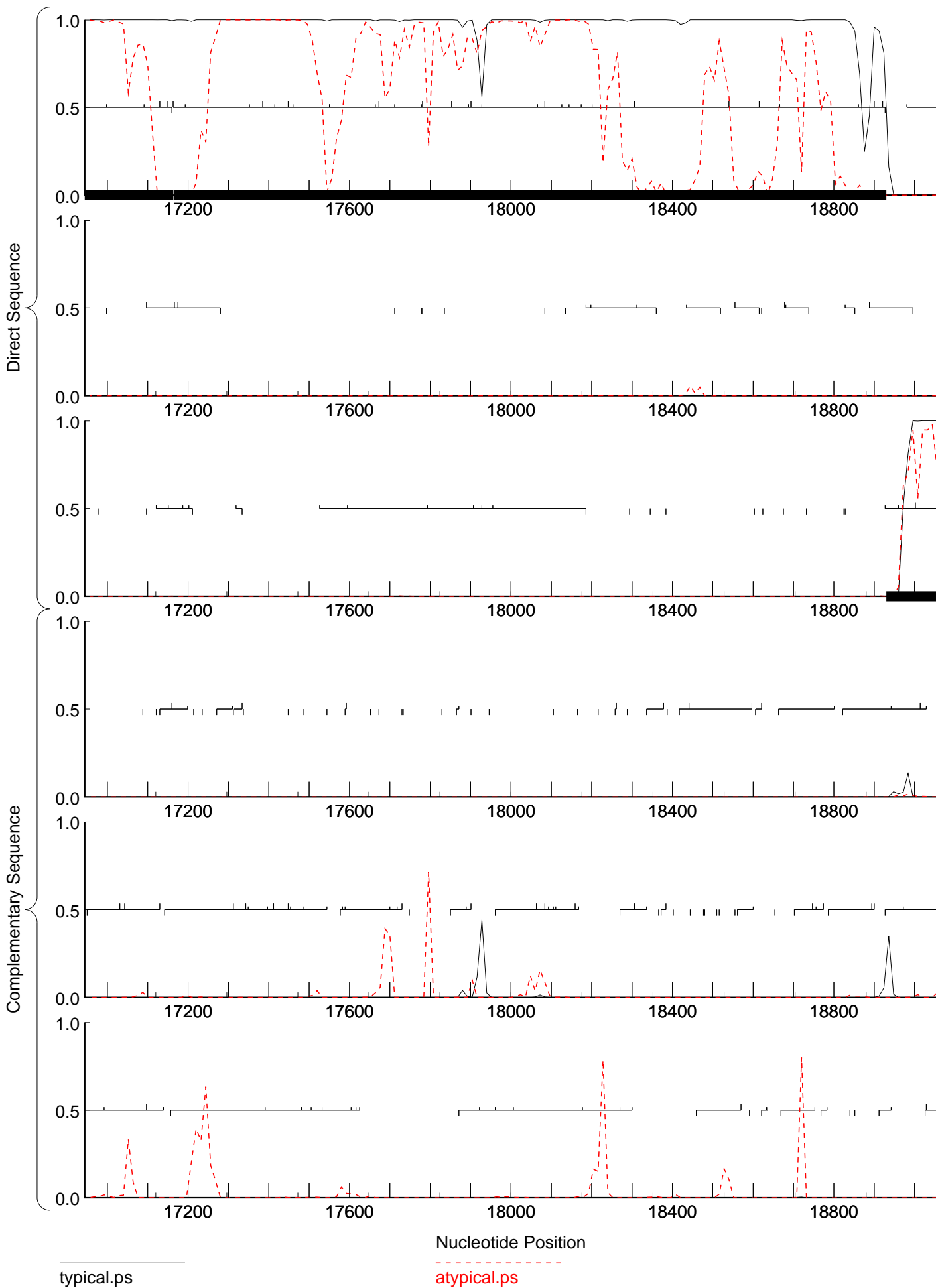






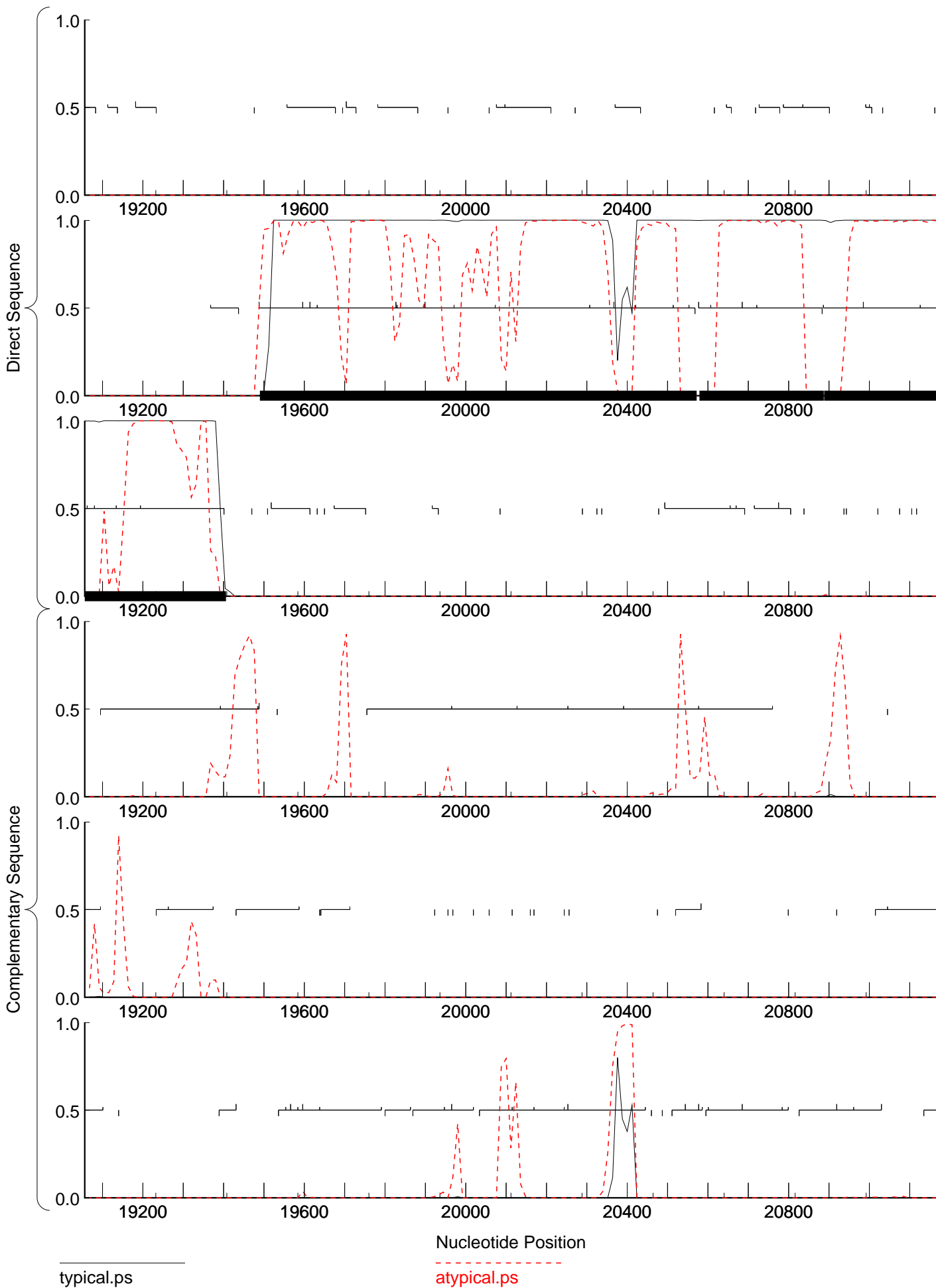


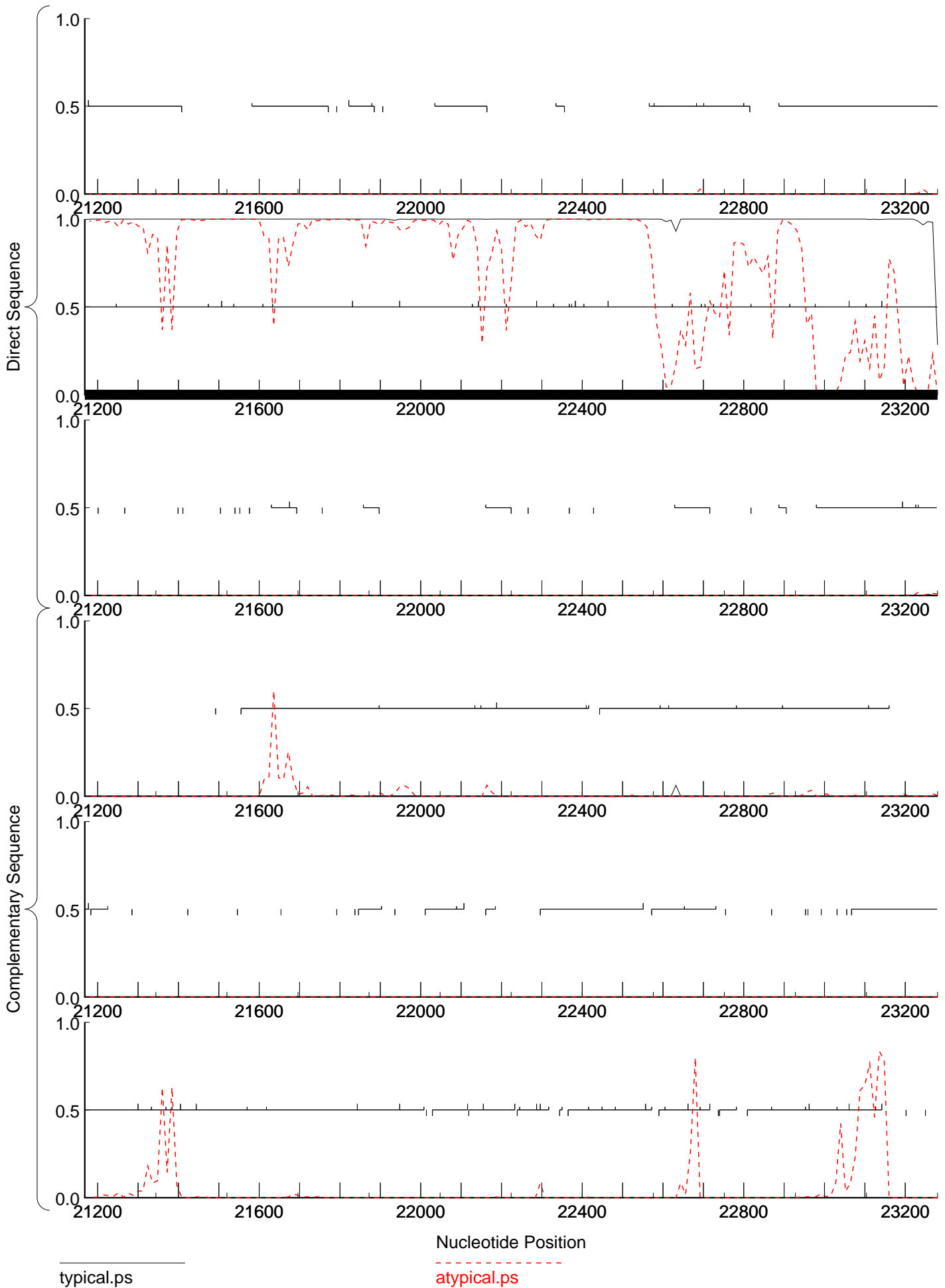


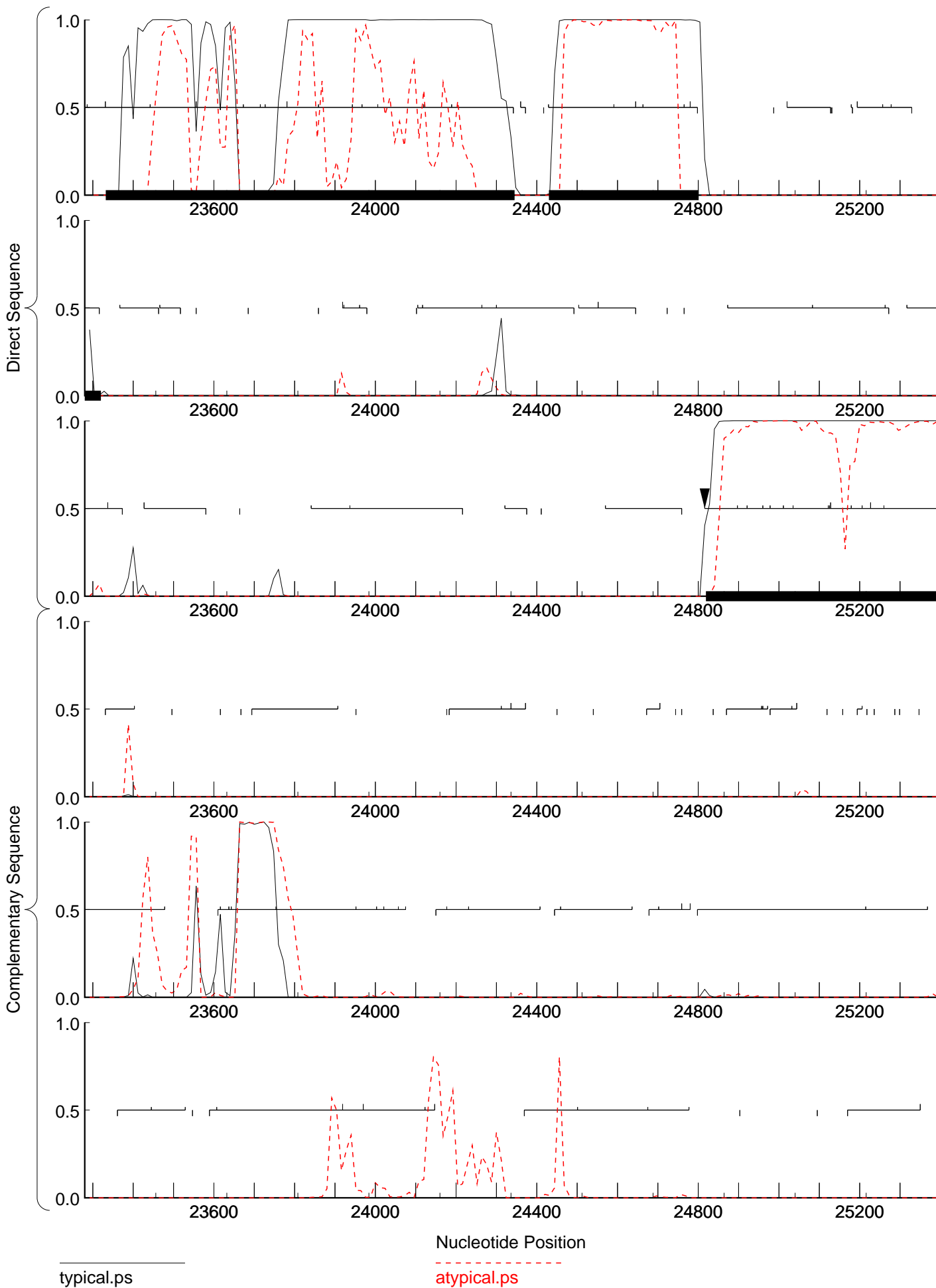


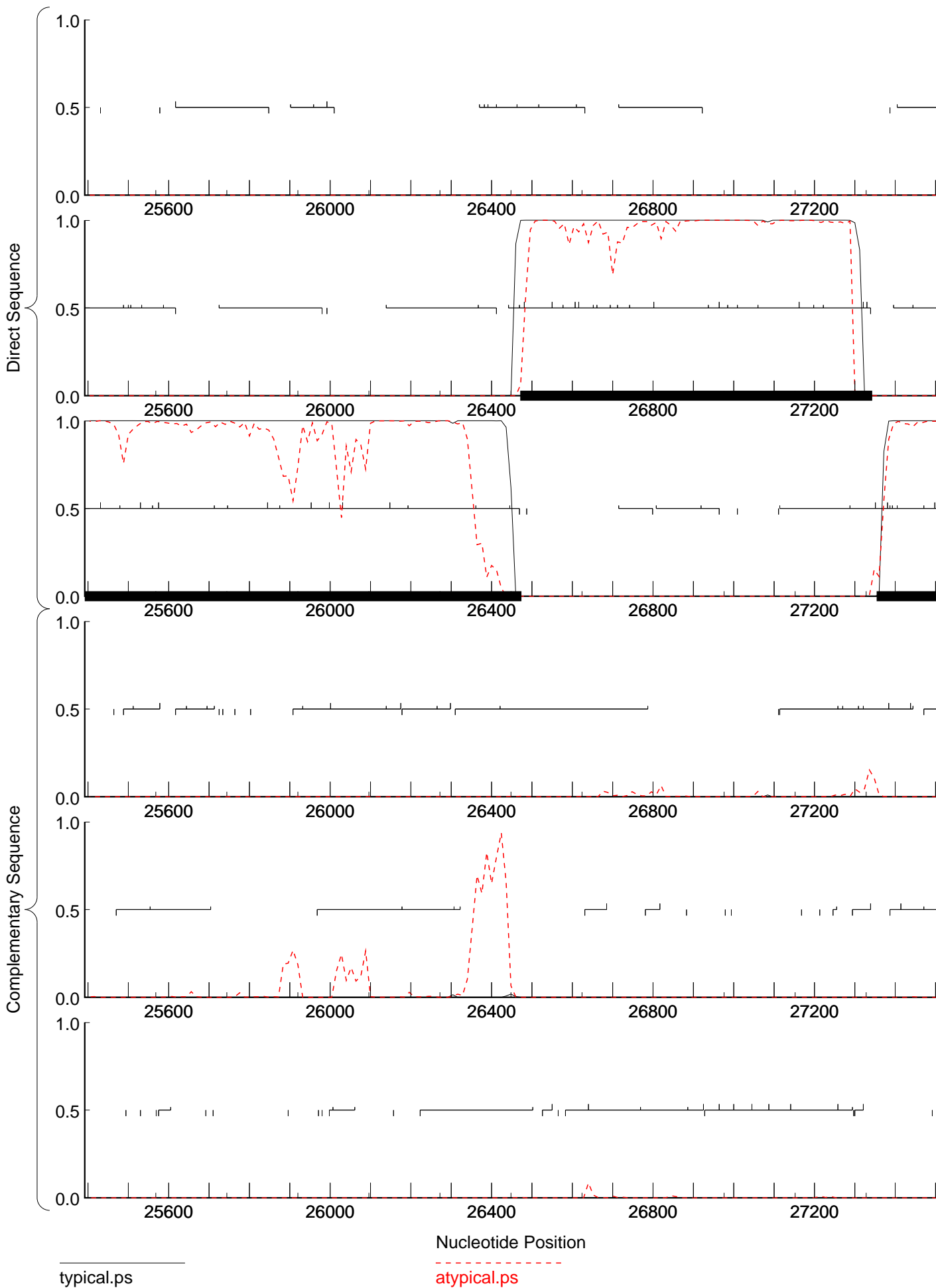
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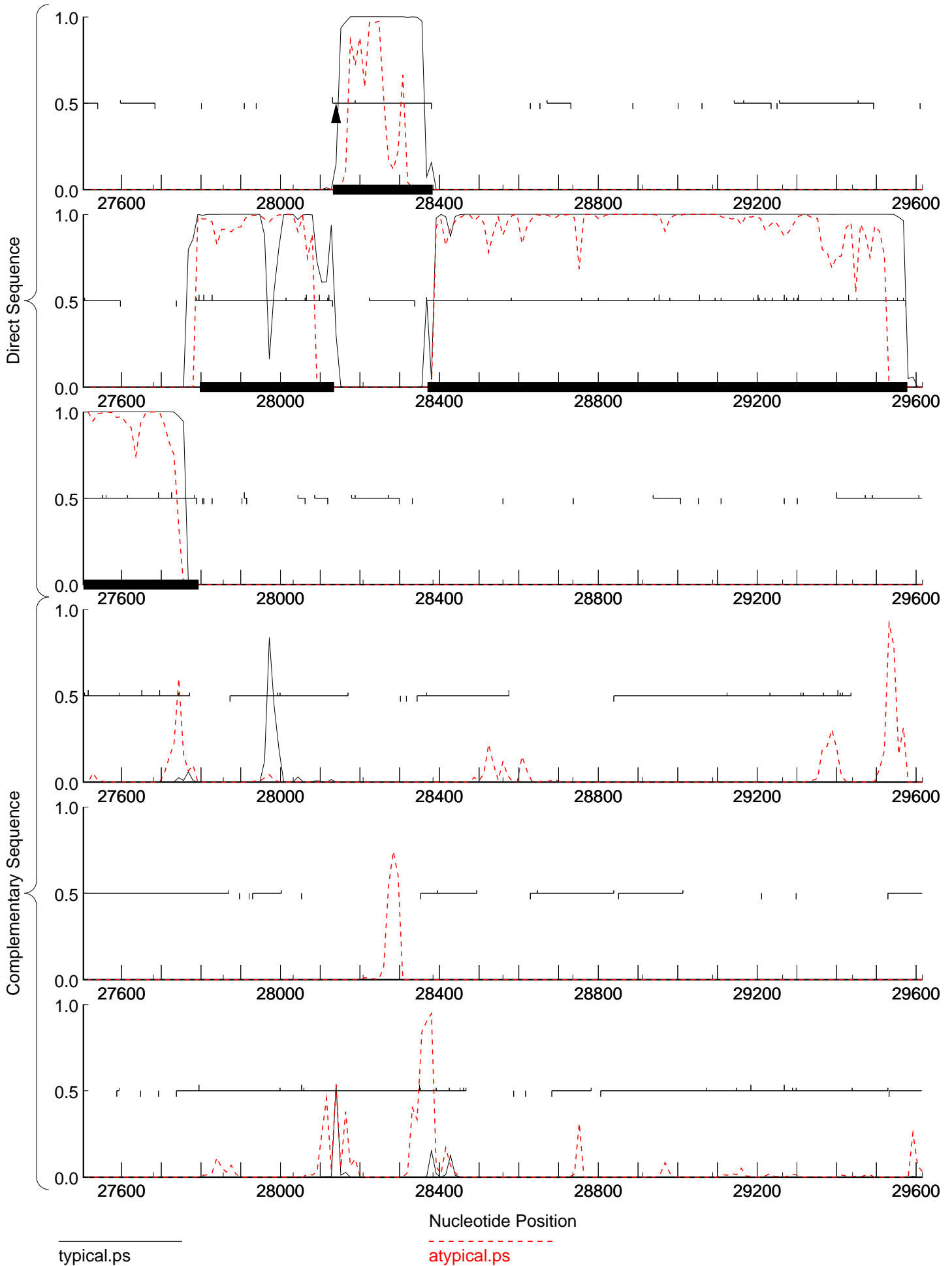
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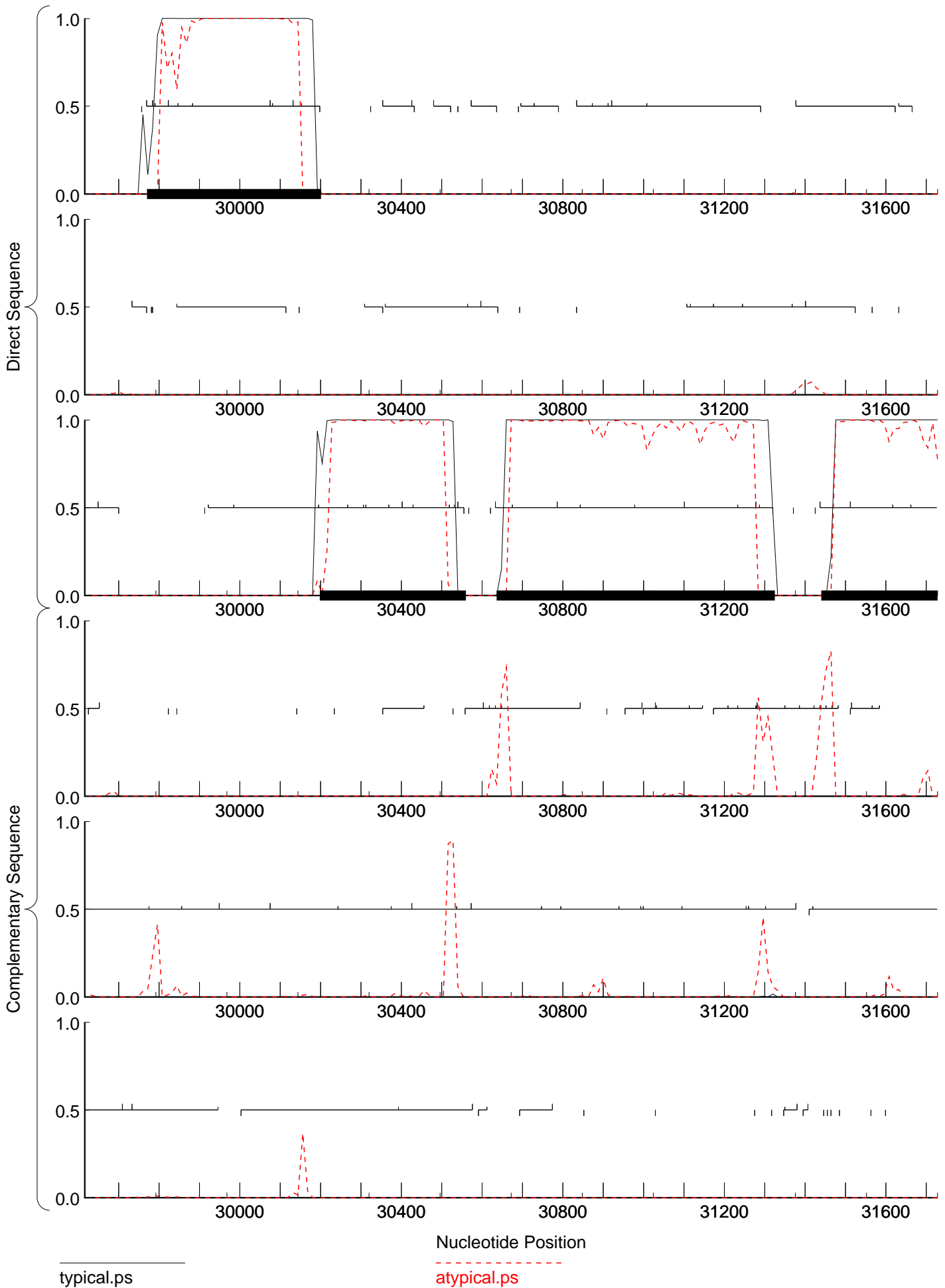


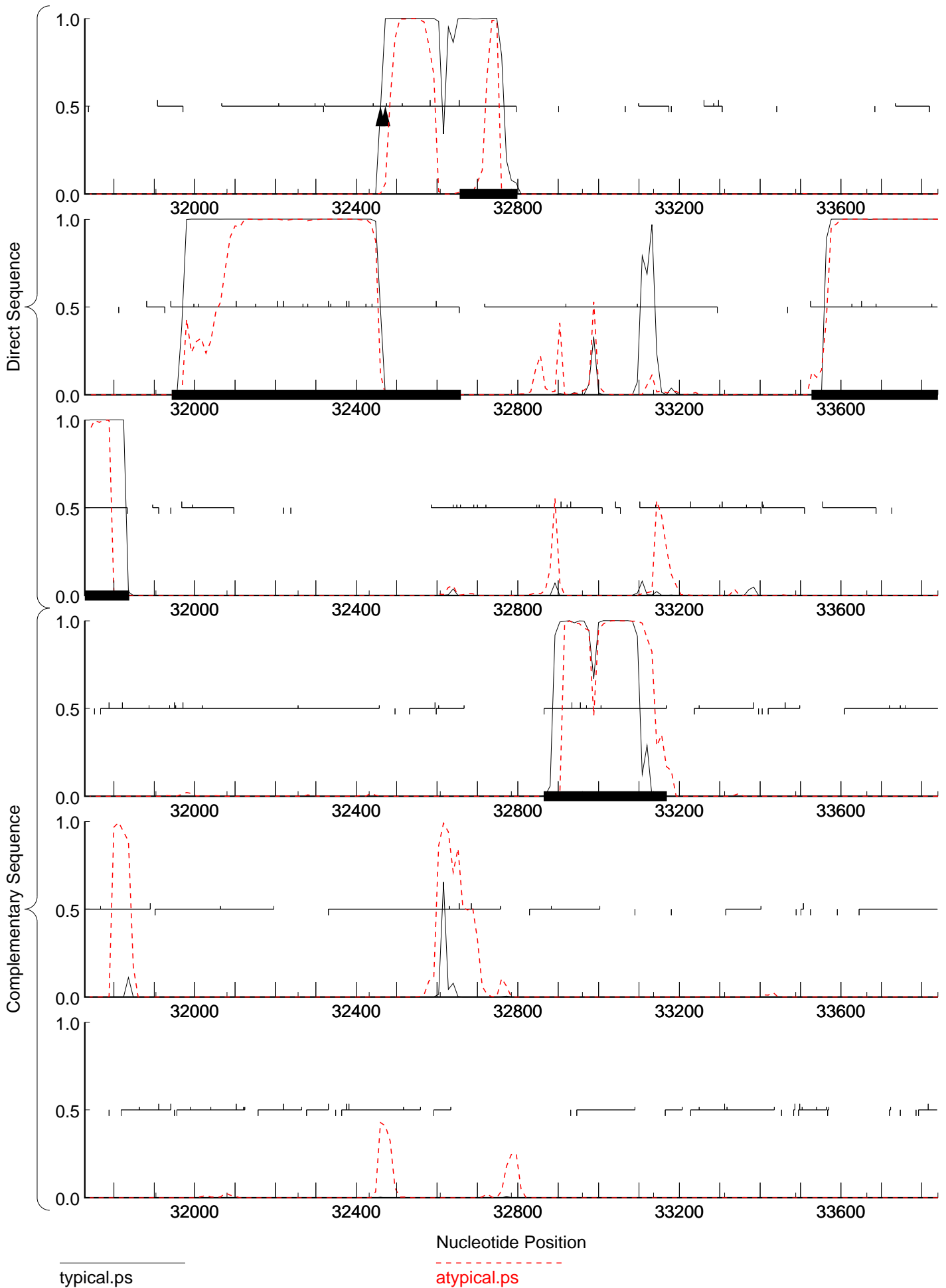


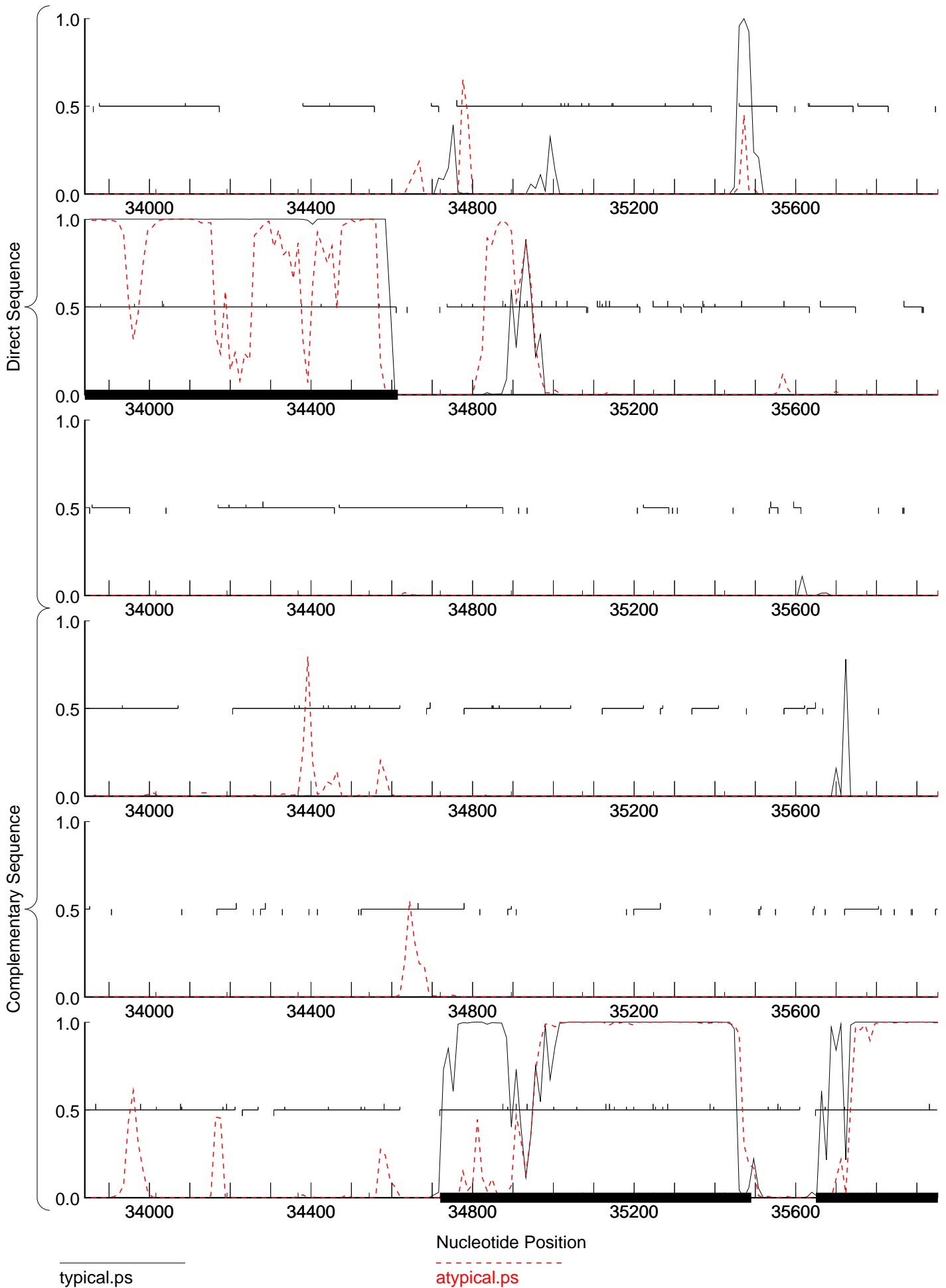


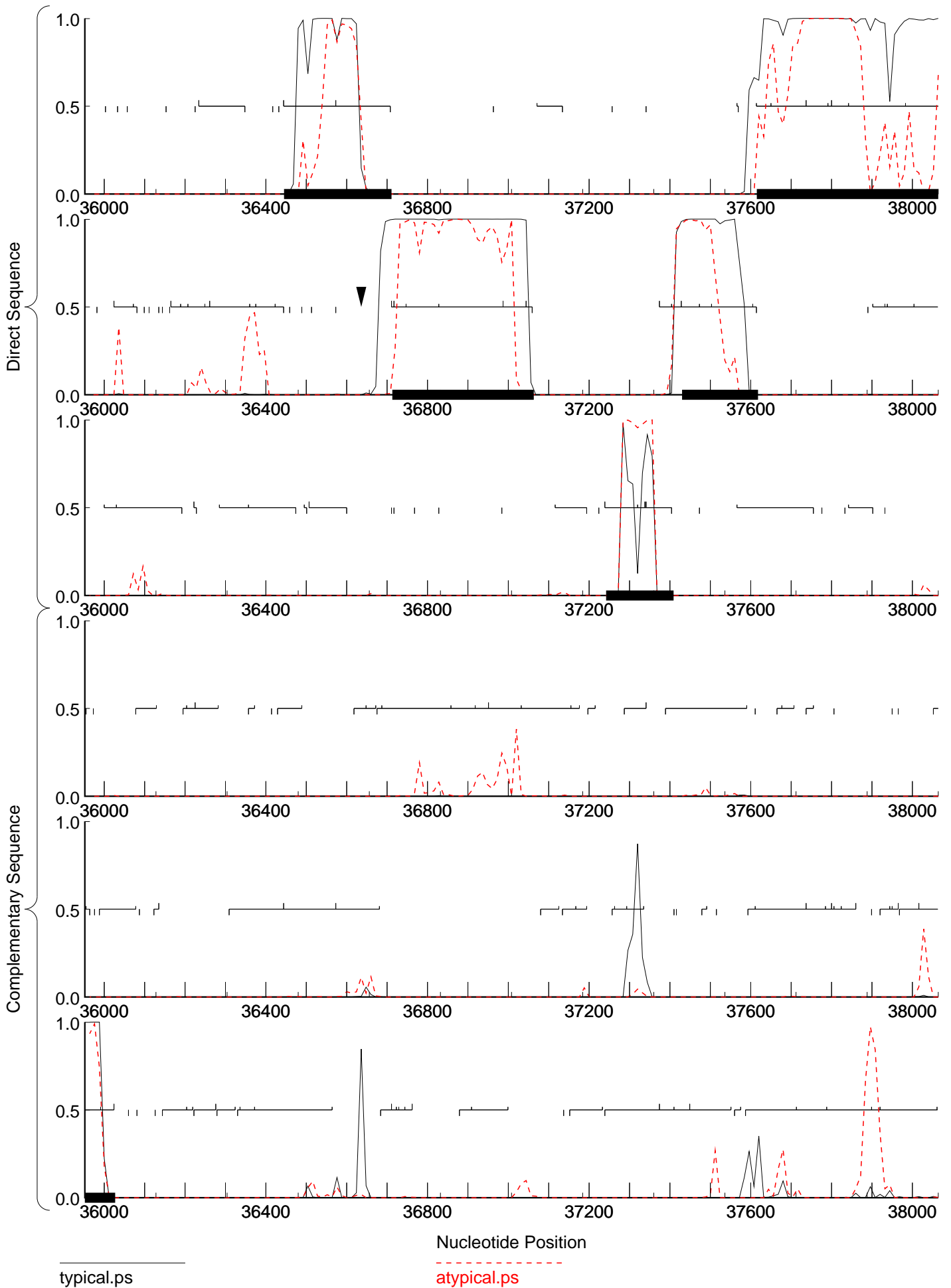


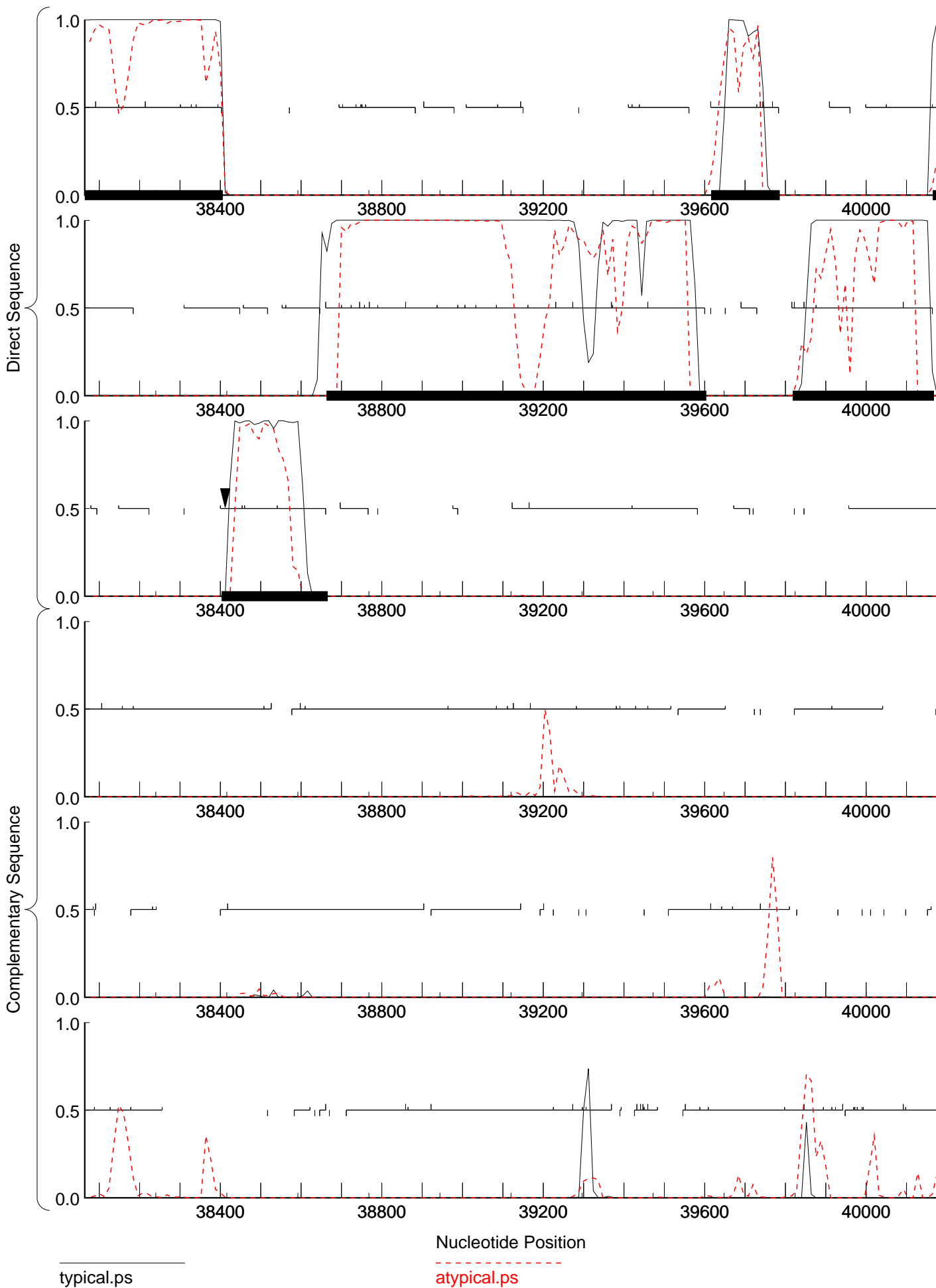


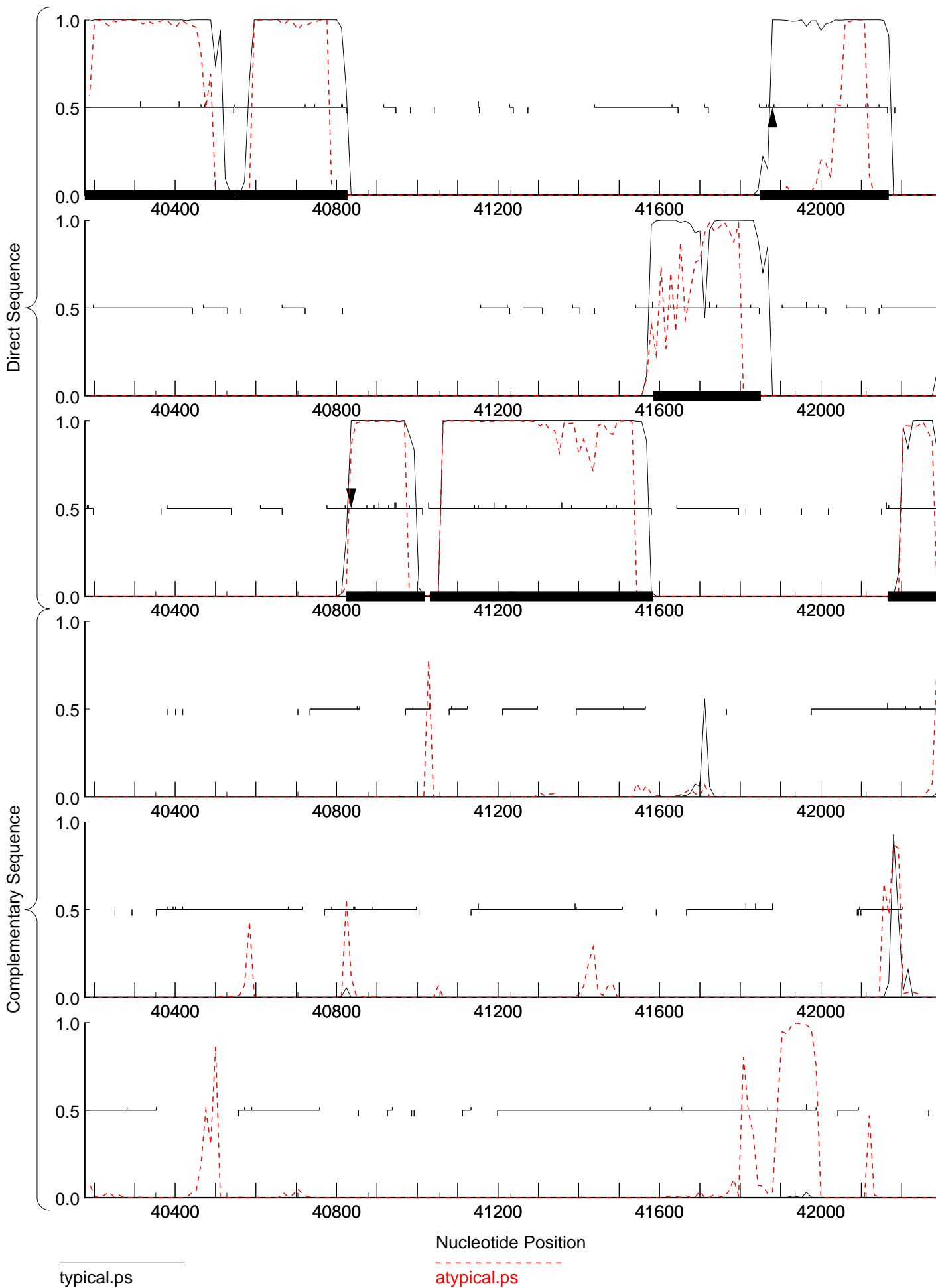












typical.ps

atypical.ps

