

GeneMark

Version 2.5p (09.08.06)

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PROGRAM INFORMATION

Sequence : Streptomyces phage Alvy complete sequence, 53011 bp including 11-base 3' overhang (CGCCACGT
Analysis Date : 4/20/18 at 0:53:54
Pages : 27
Sequence Length : 53011 bp
GC Content : 67.55%

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,

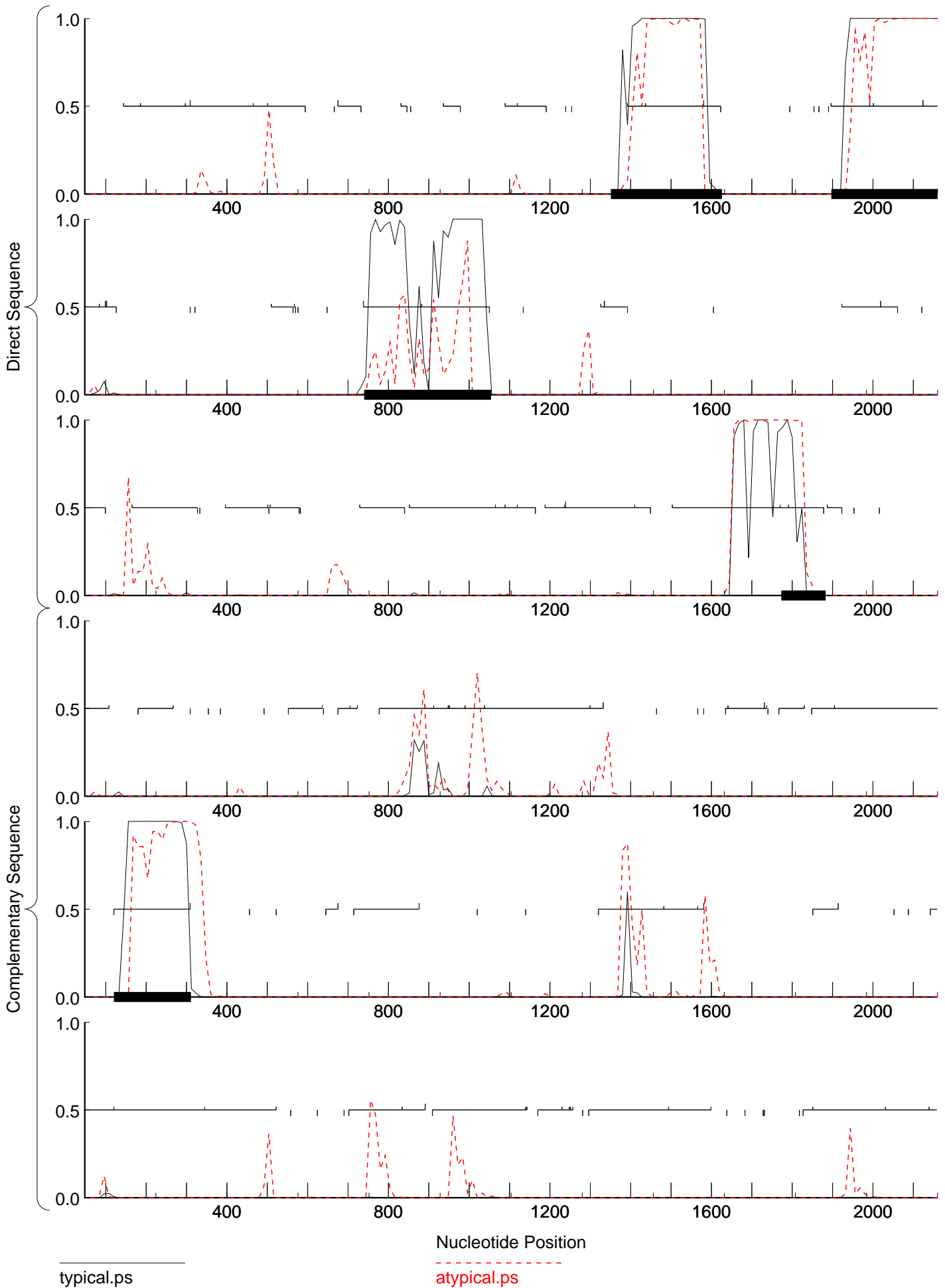
Matrix notes & comments

Training set derived by GeneMarkS, 4.30 October 2014
Fri Apr 20 00:53:54 2018

MATRIX INFORMATION

Matrix : -
Author : -
Order : 4

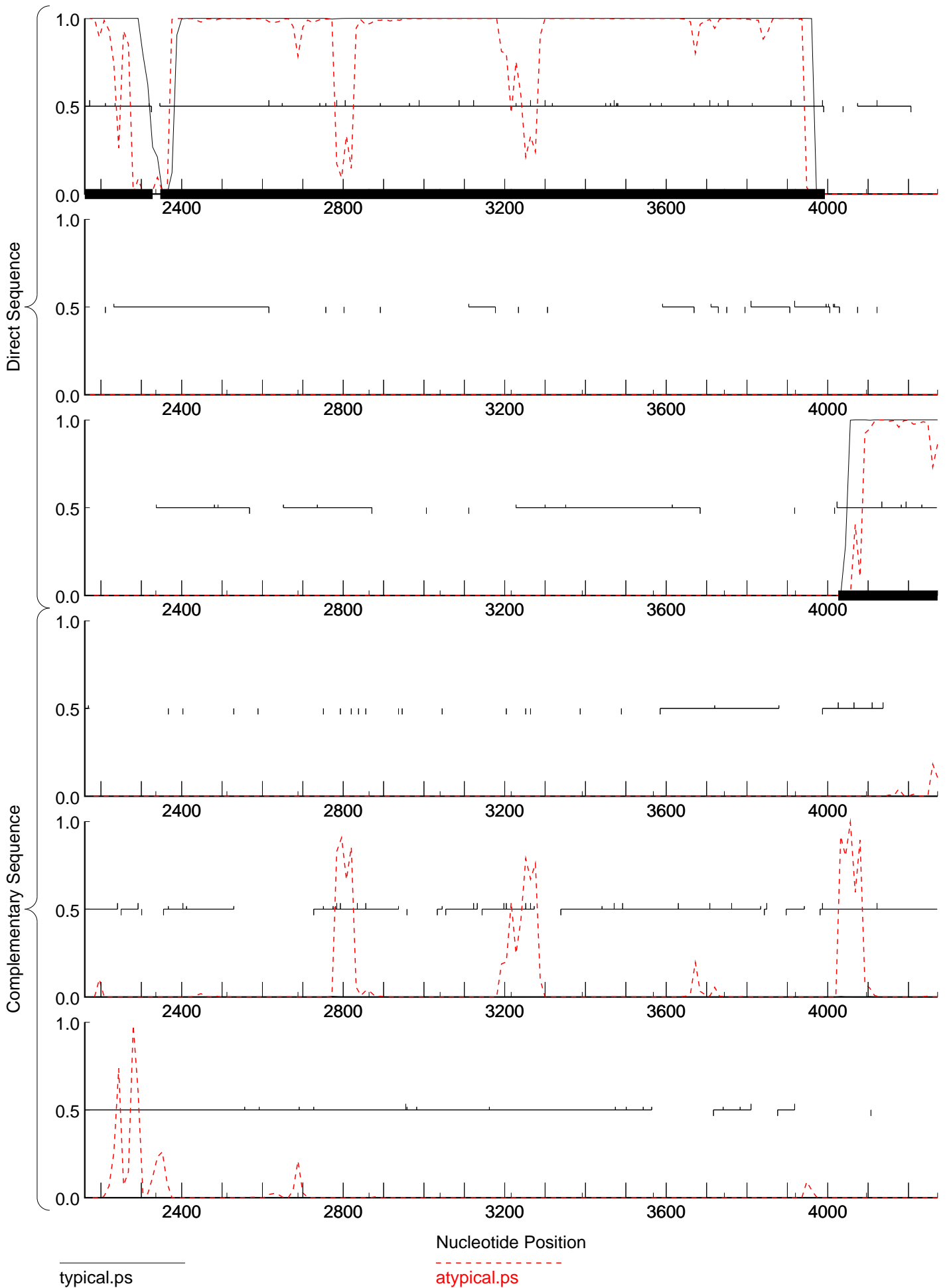
Send questions / comments to:
Dr. M. Borodovsky
Georgia Institute of Technology
School of Biology
Atlanta, GA 30332-0230



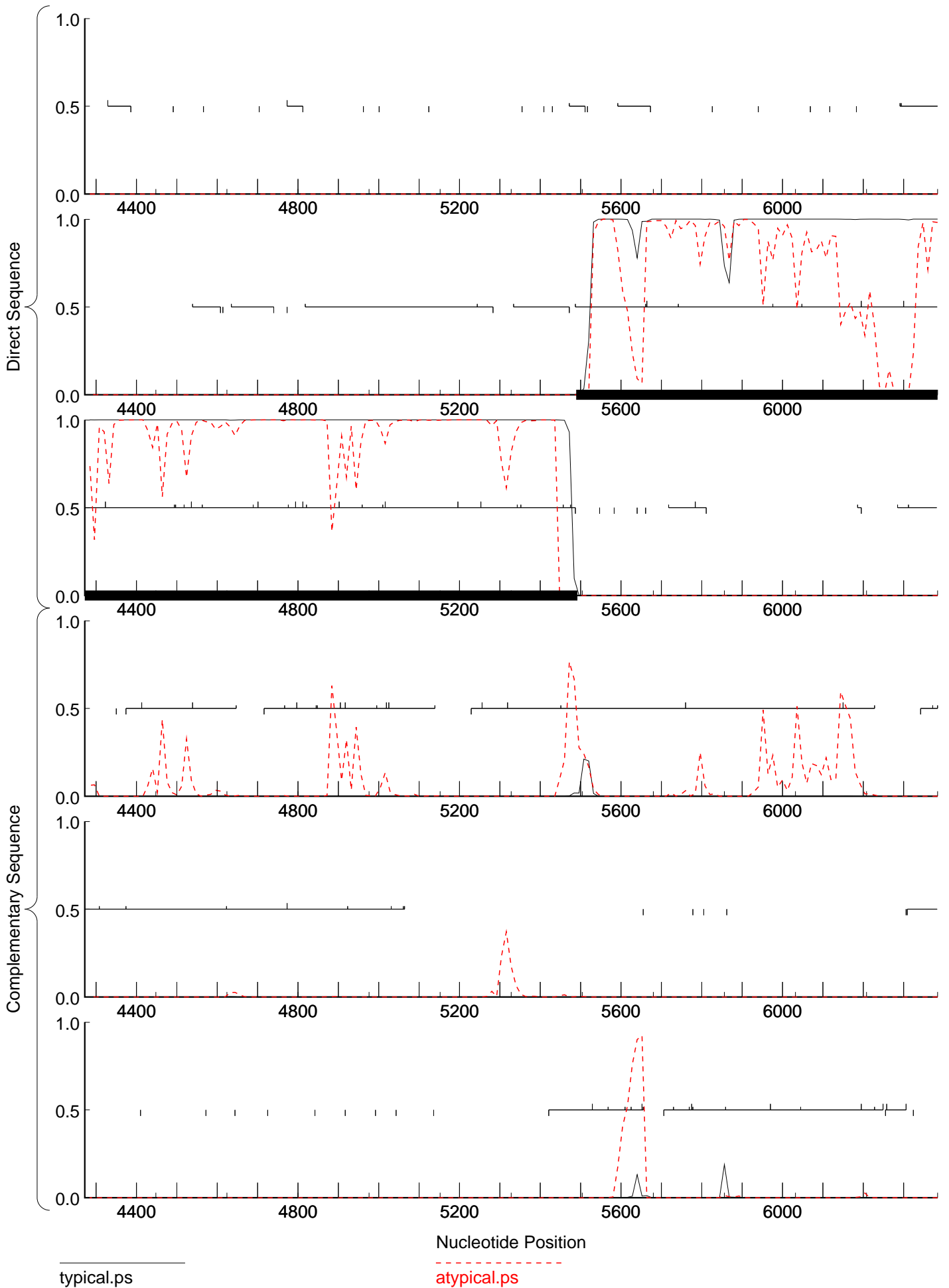
typical.ps

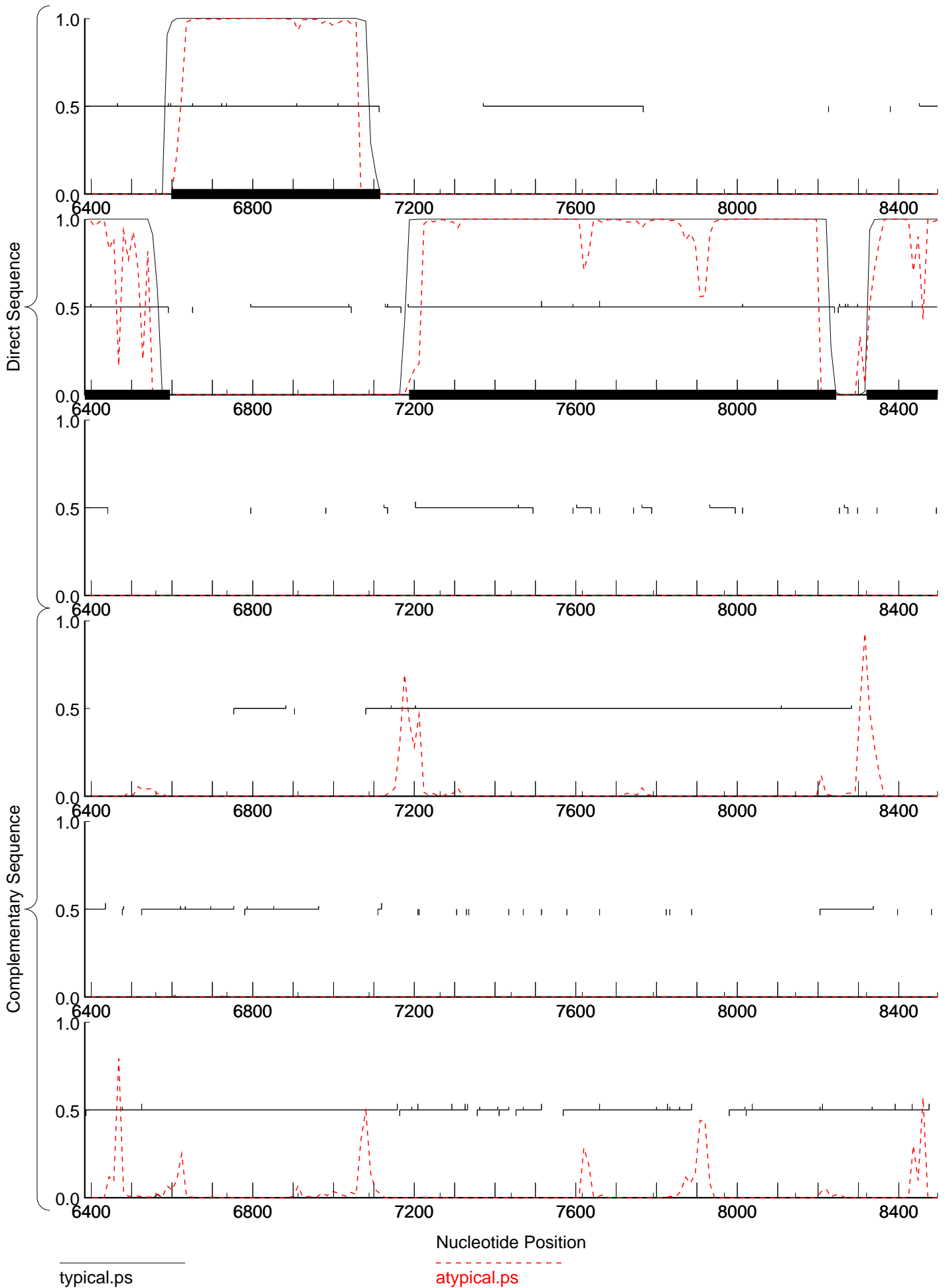
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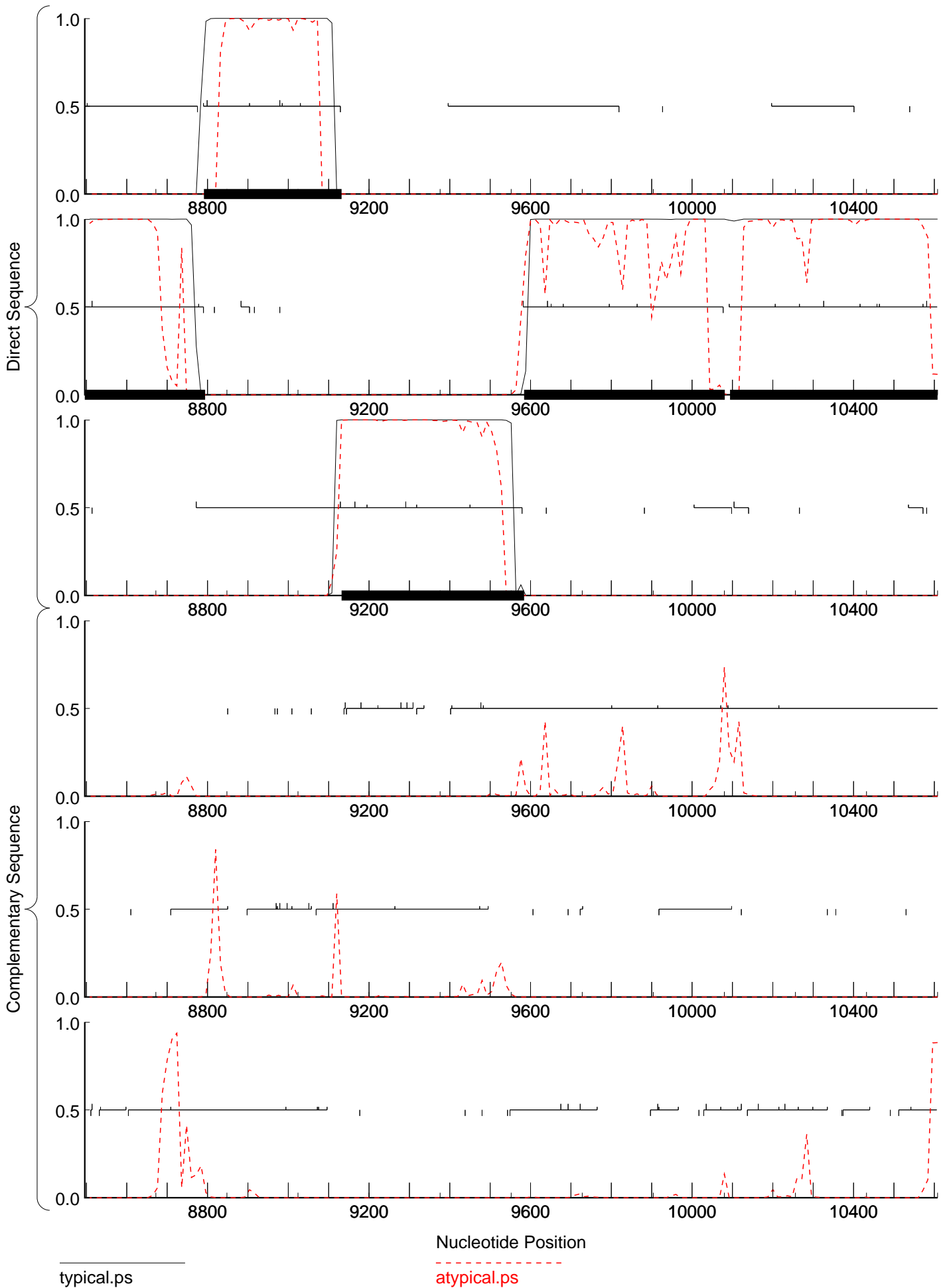
complete sequence, 53011 bp including 11-base 3' overhang (CGCCACGTCTT), Cluster BD2, Order 4, Window 96, Step 12, 3/27
complete sequence, 53011 bp including 11-base 3' overhang (CGCCACGTCTT), Cluster BD2, Order 2, Window 96, Step 12, 3/27

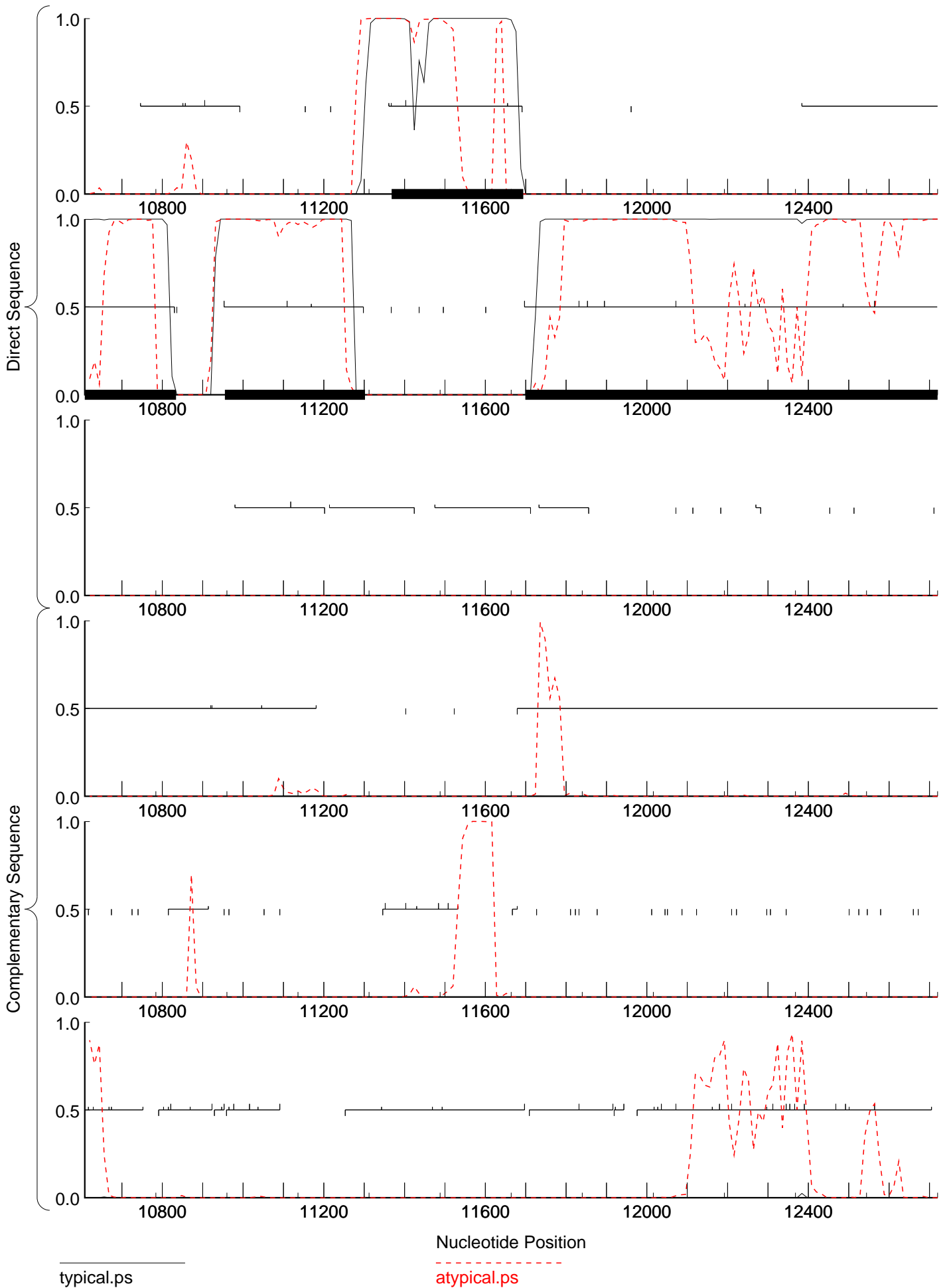


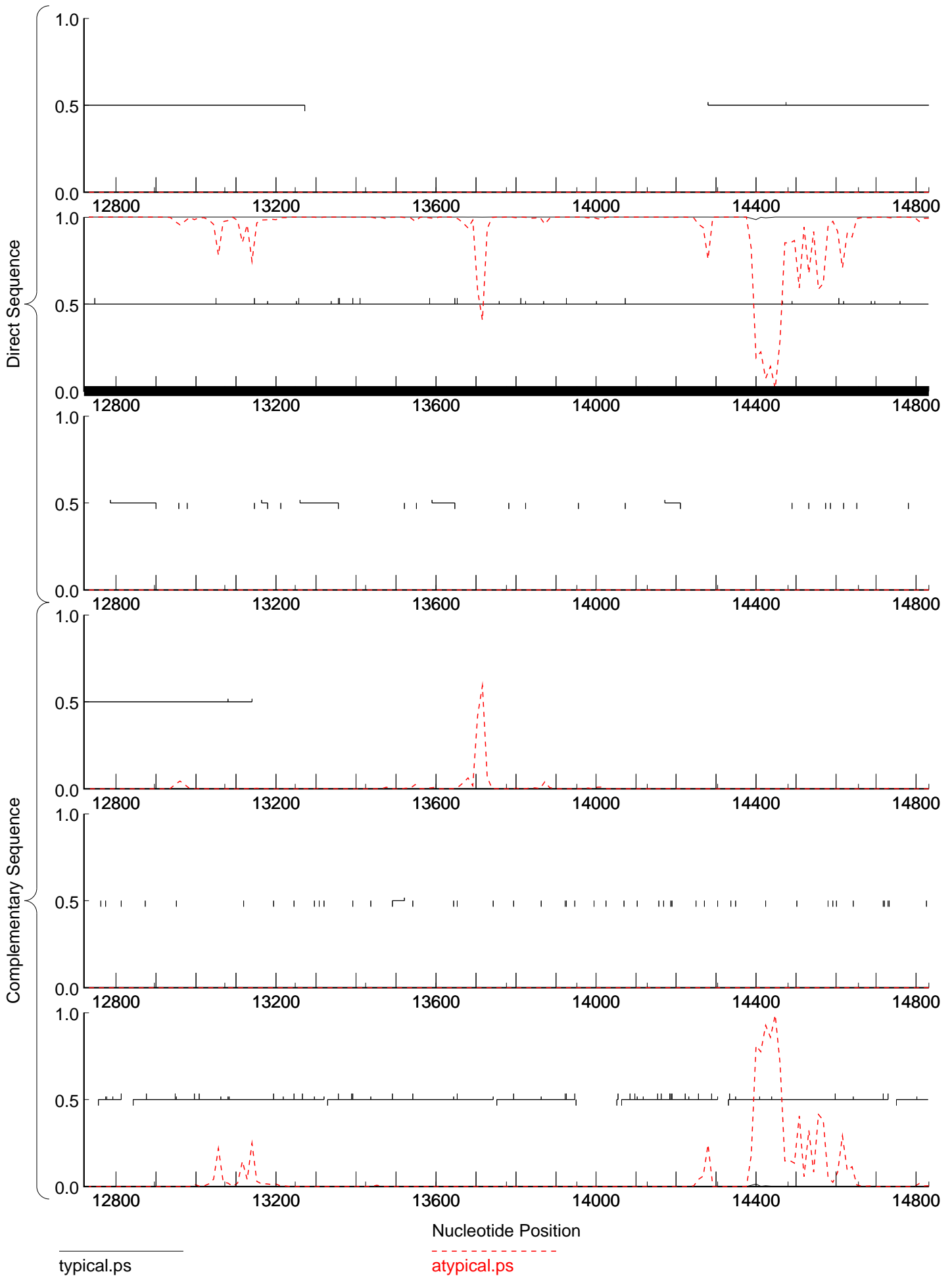
complete sequence, 53011 bp including 11-base 3' overhang (CGCCACGTCTT), Cluster BD2, Order 4, Window 96, Step 12, 4/27
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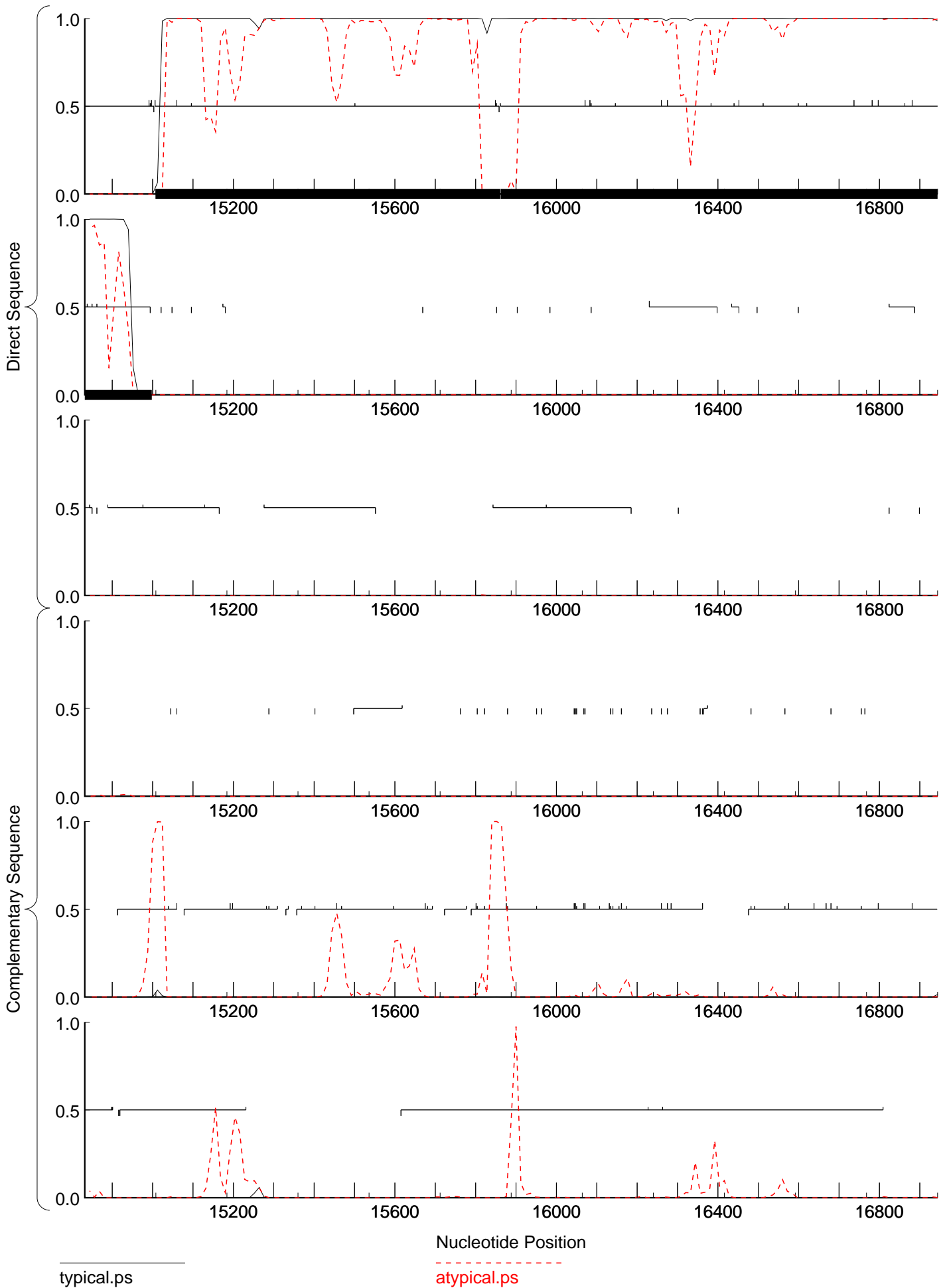


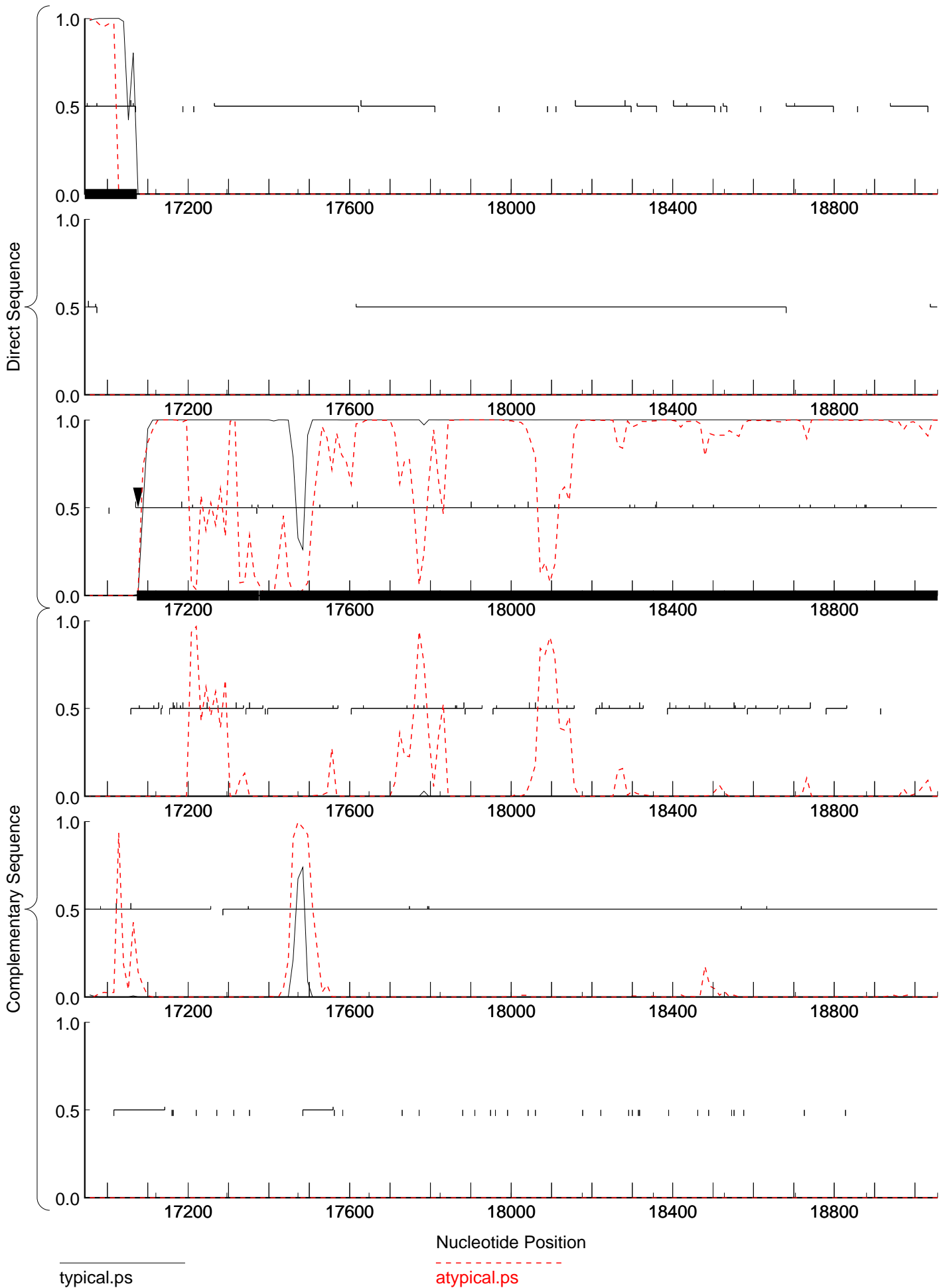


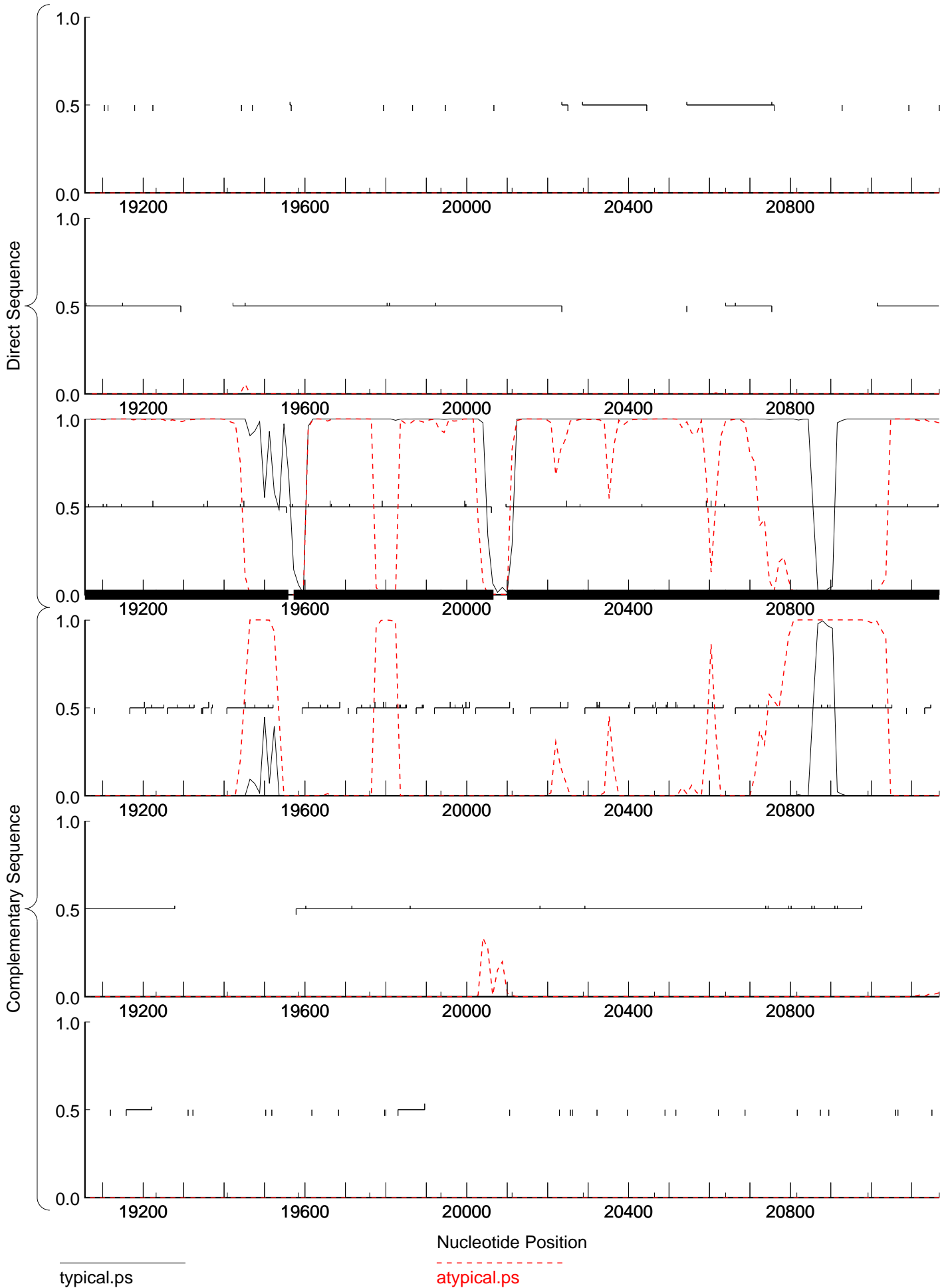




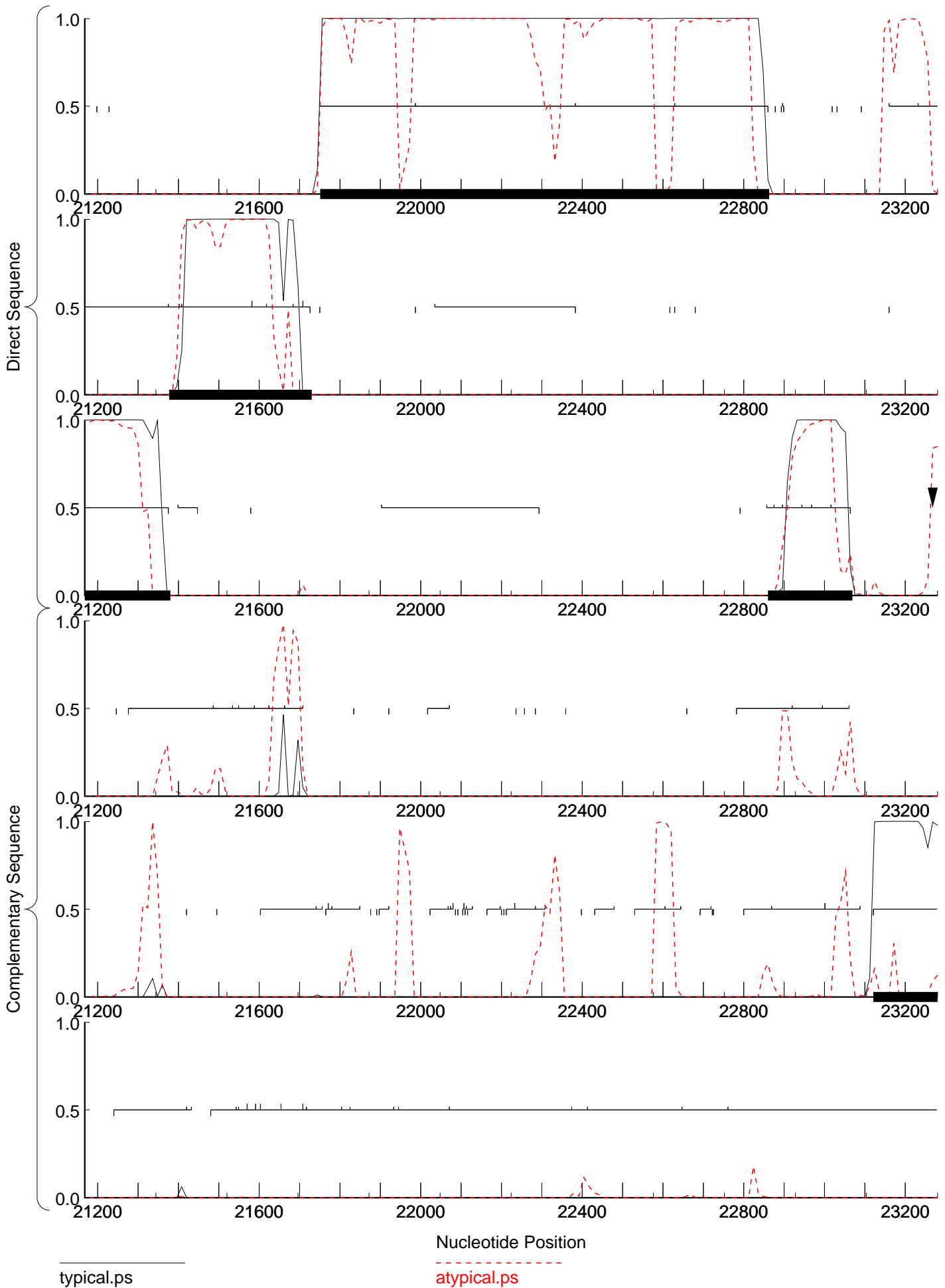
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complete sequence, 53011 bp including 11-base 3' overhang (CGCCACGTCTT), Cluster BD2, Order 2, Window 96, Step 12, 9/27

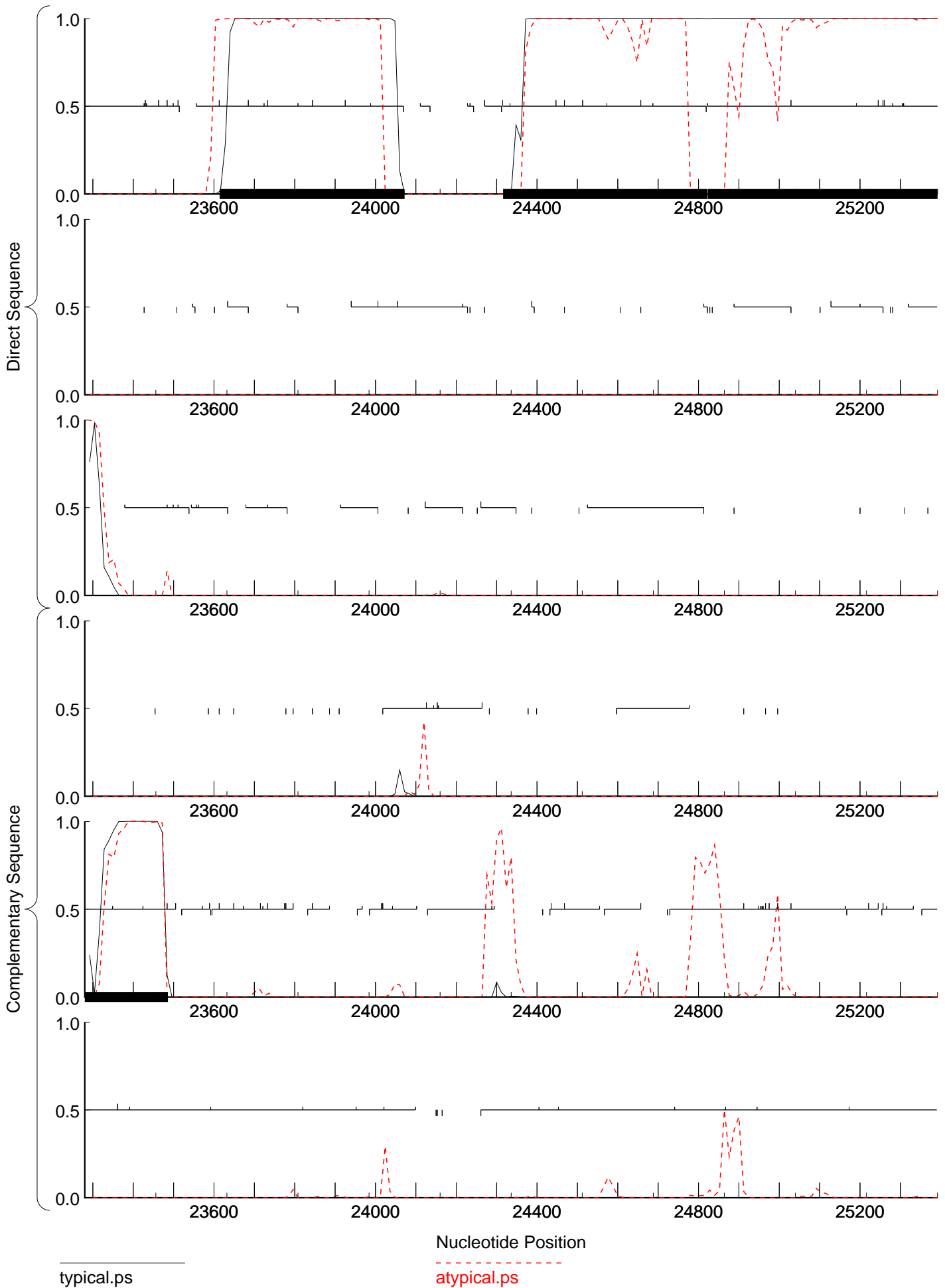






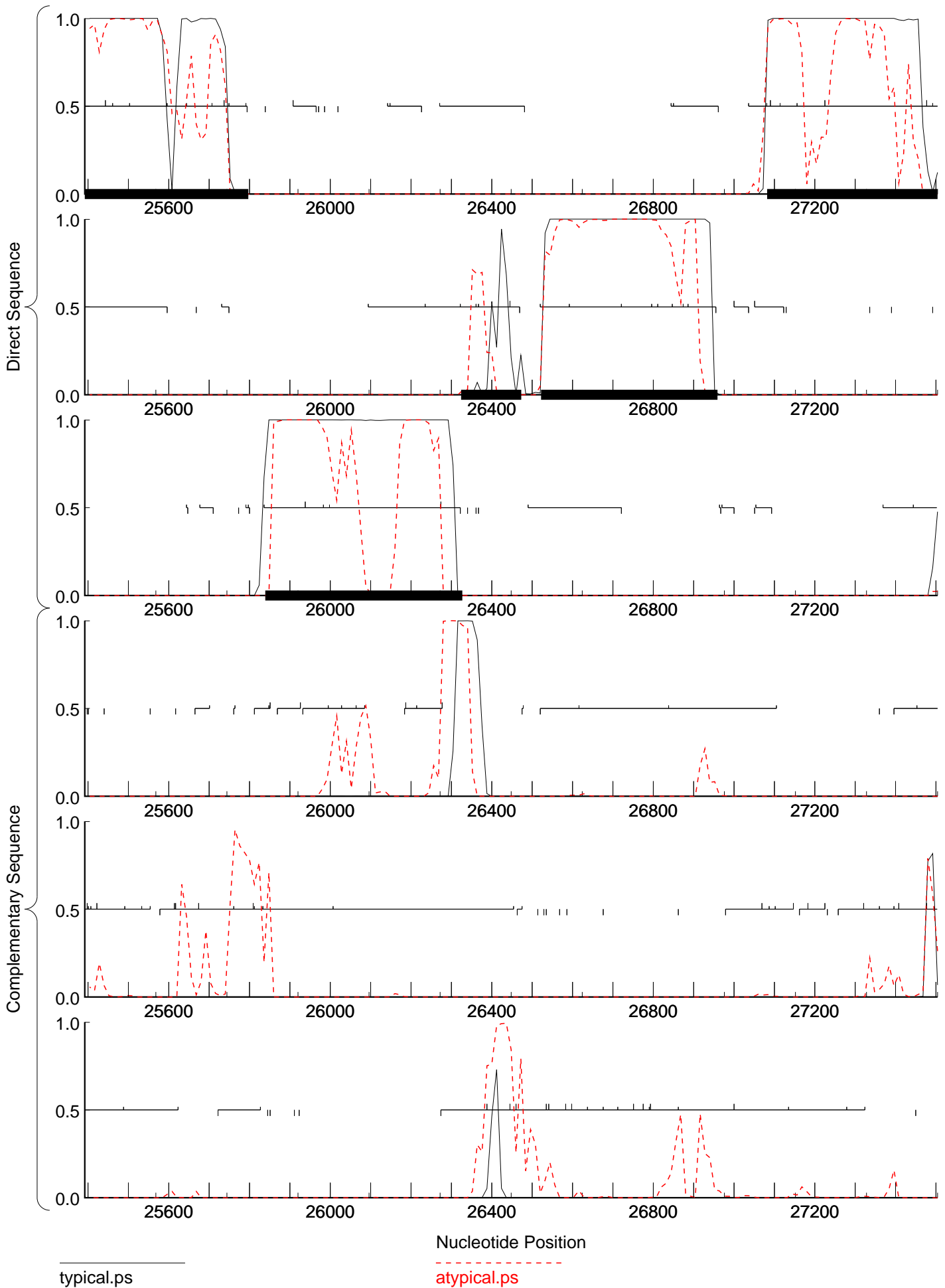
GeneMark.hmm prediction

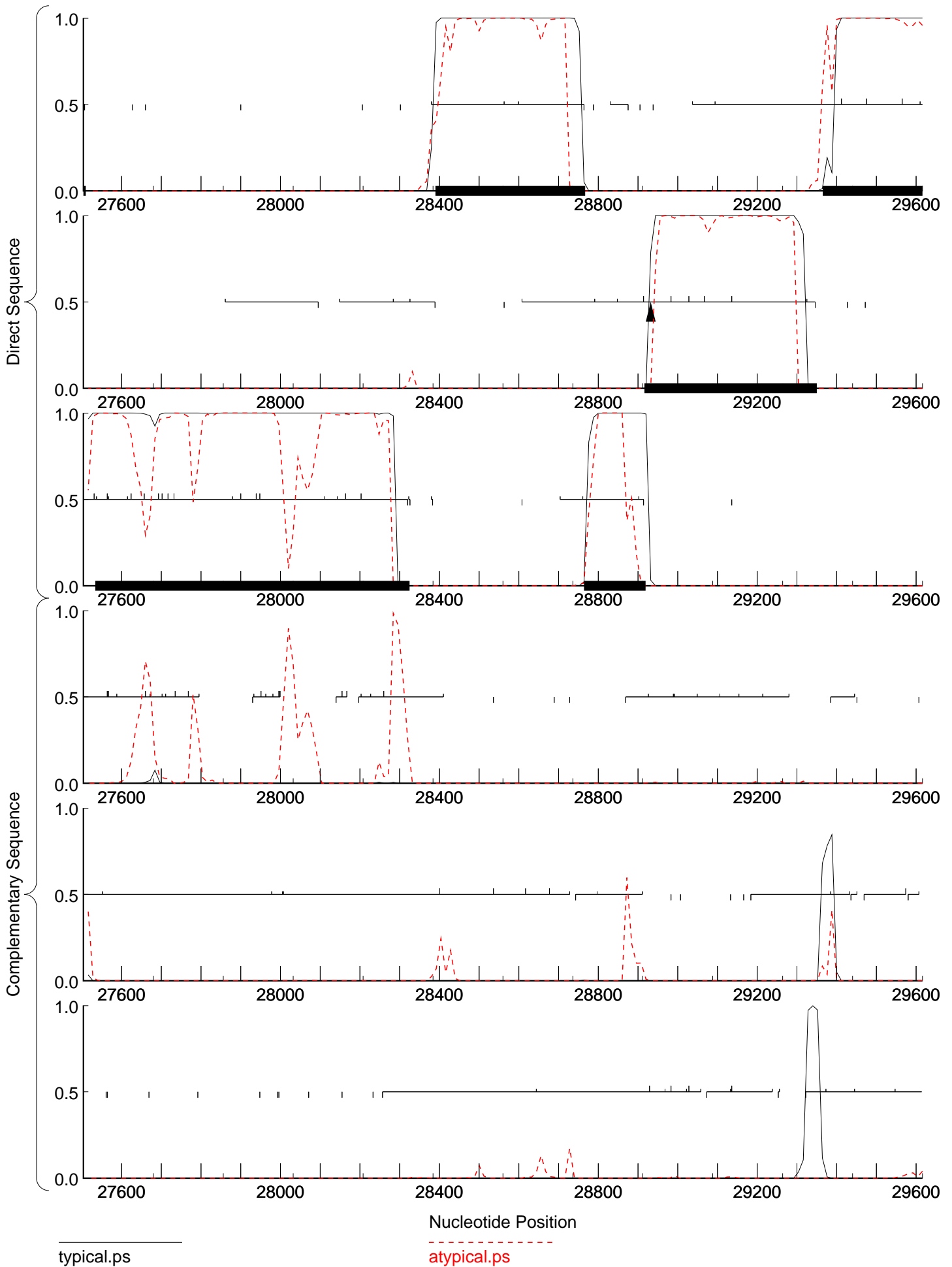




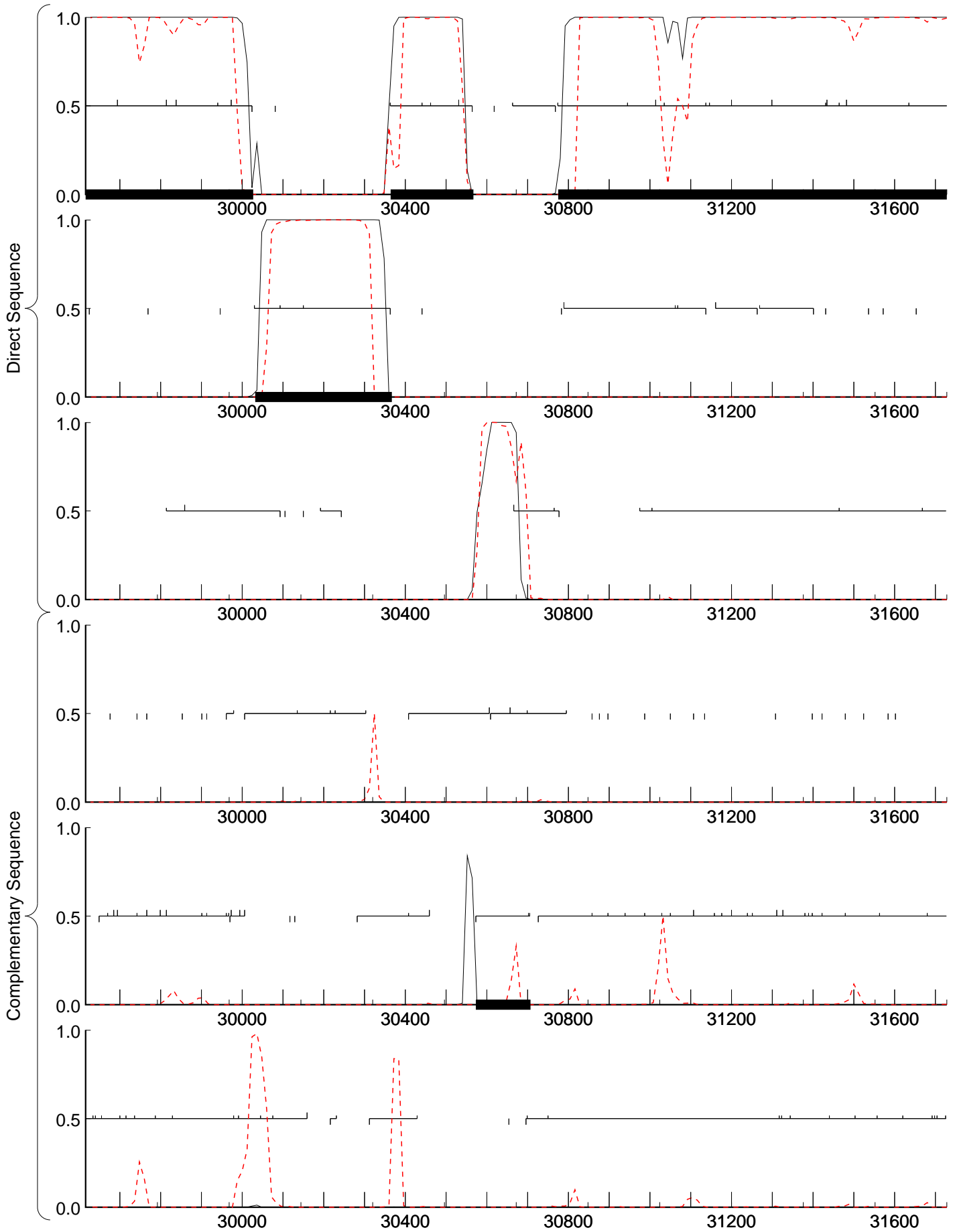
typical.ps

atypical.ps





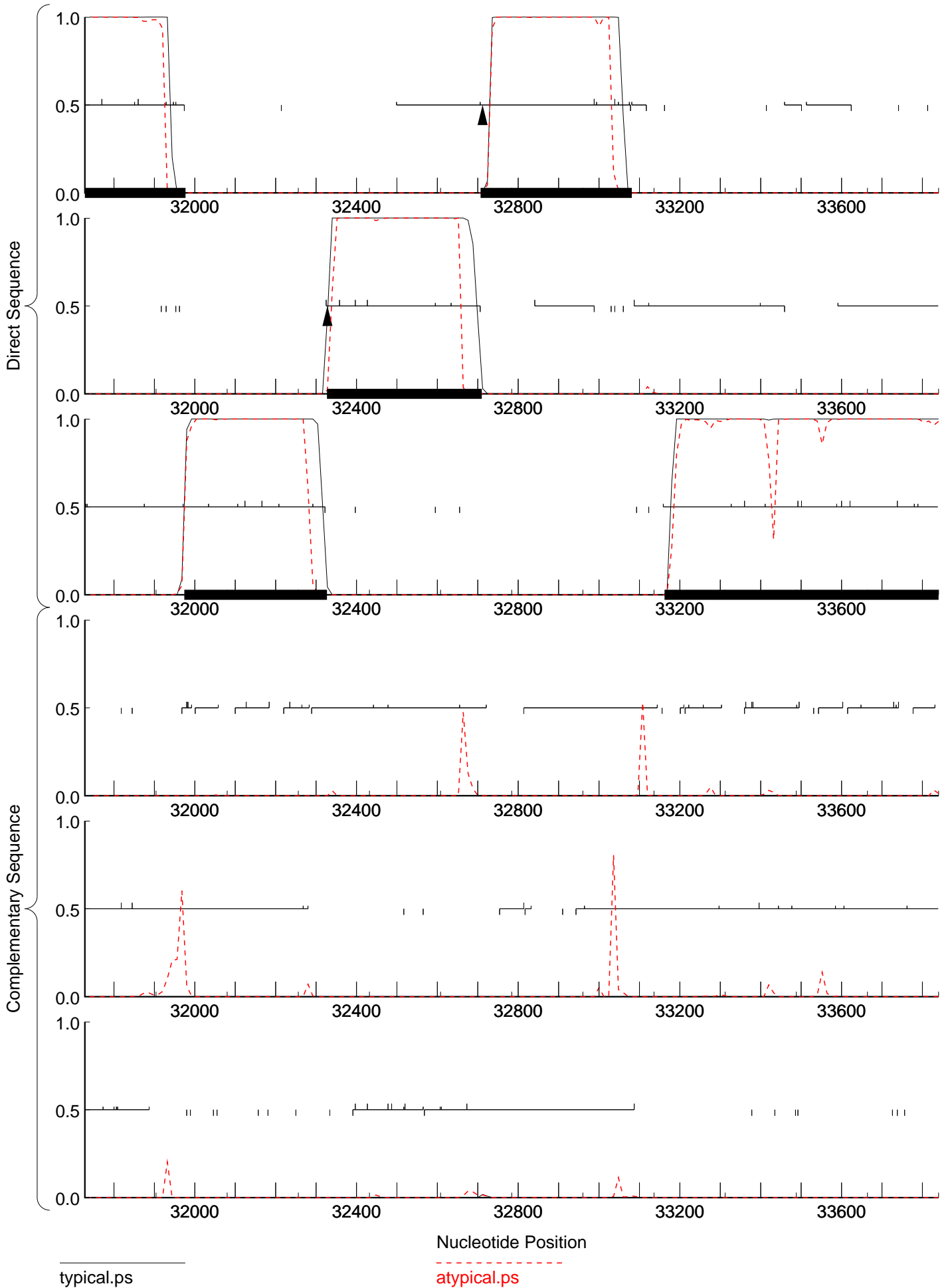
GeneMark.hmm prediction

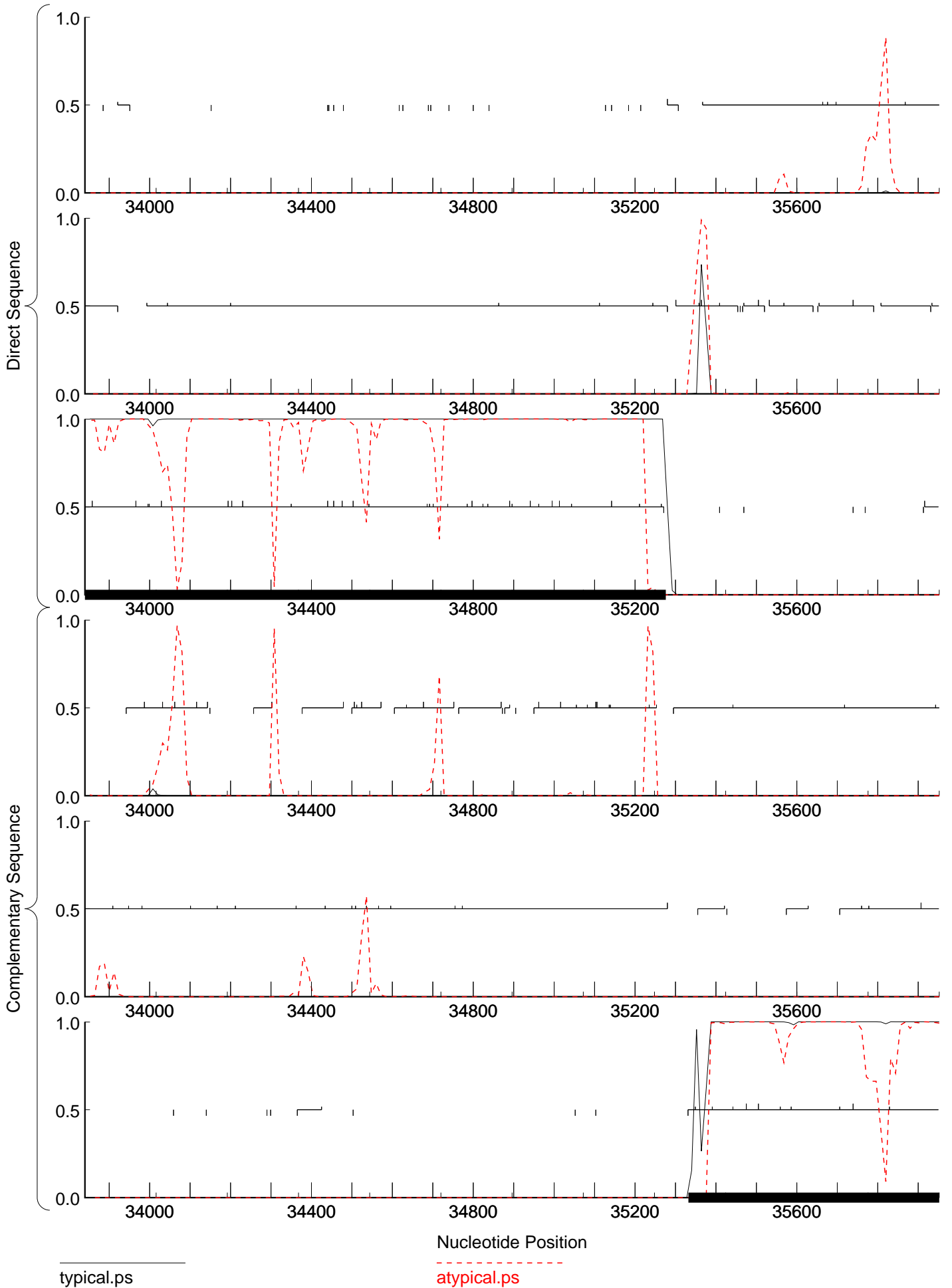


typical.ps

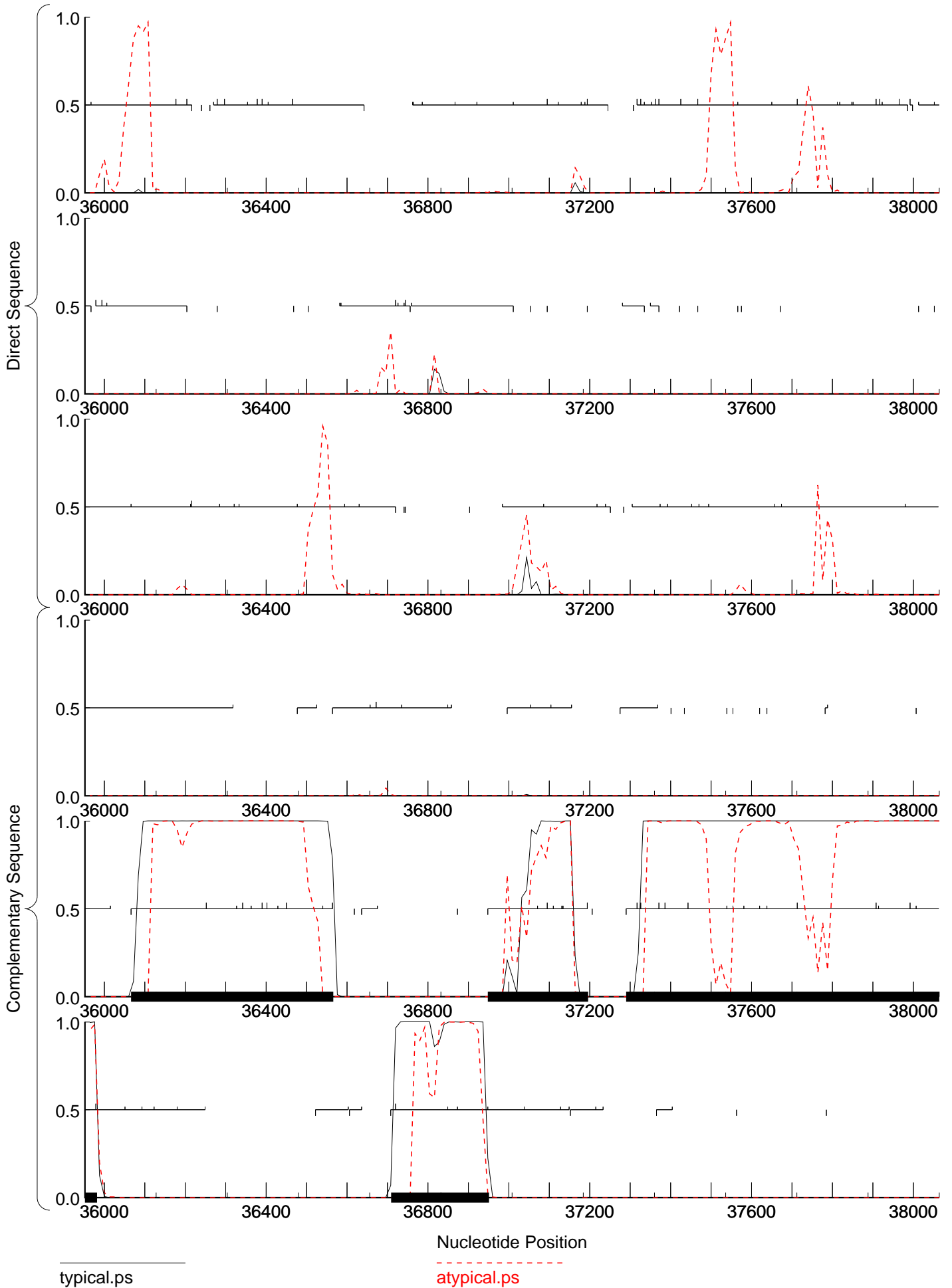
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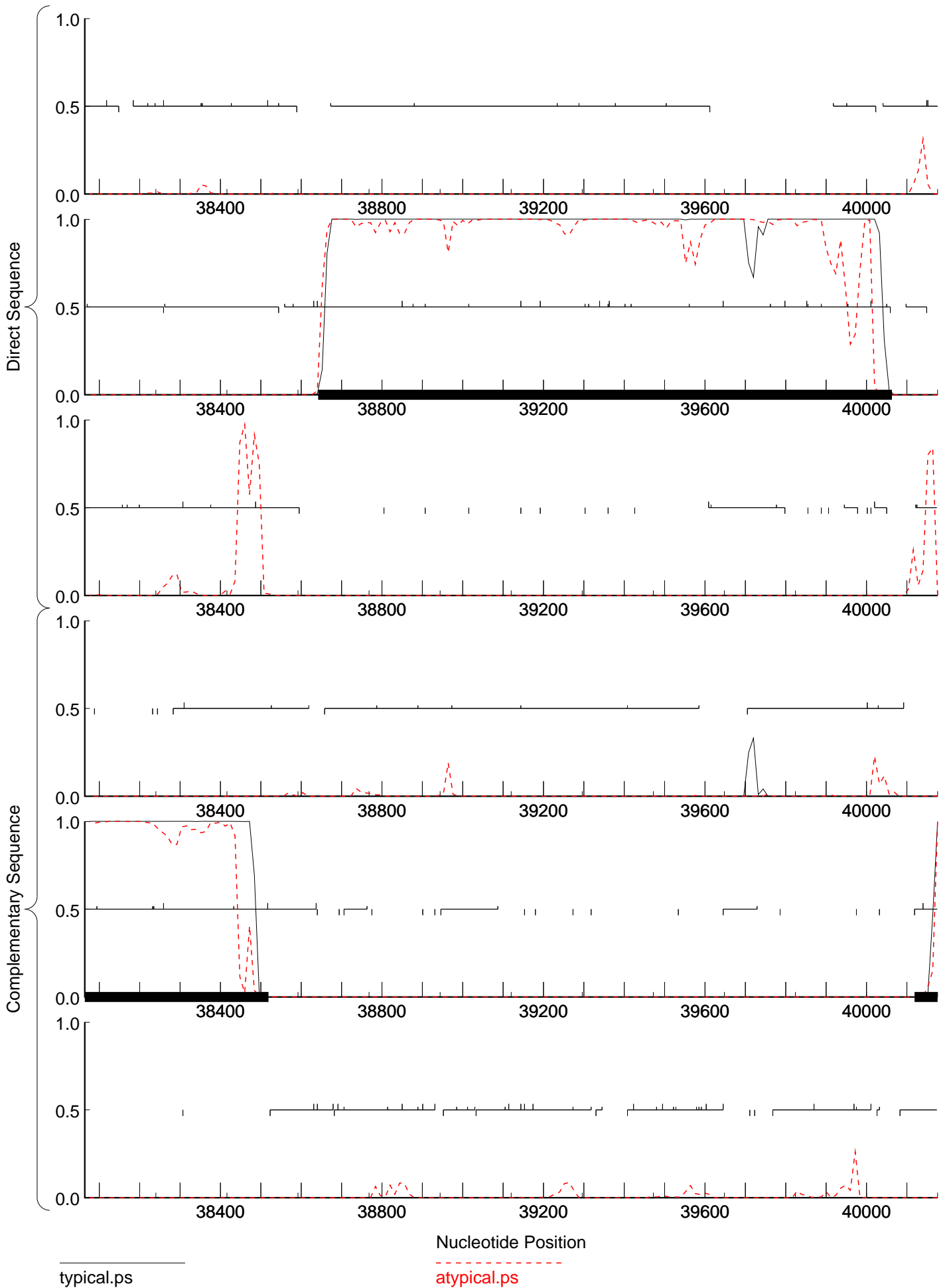
GeneMark.hmm prediction

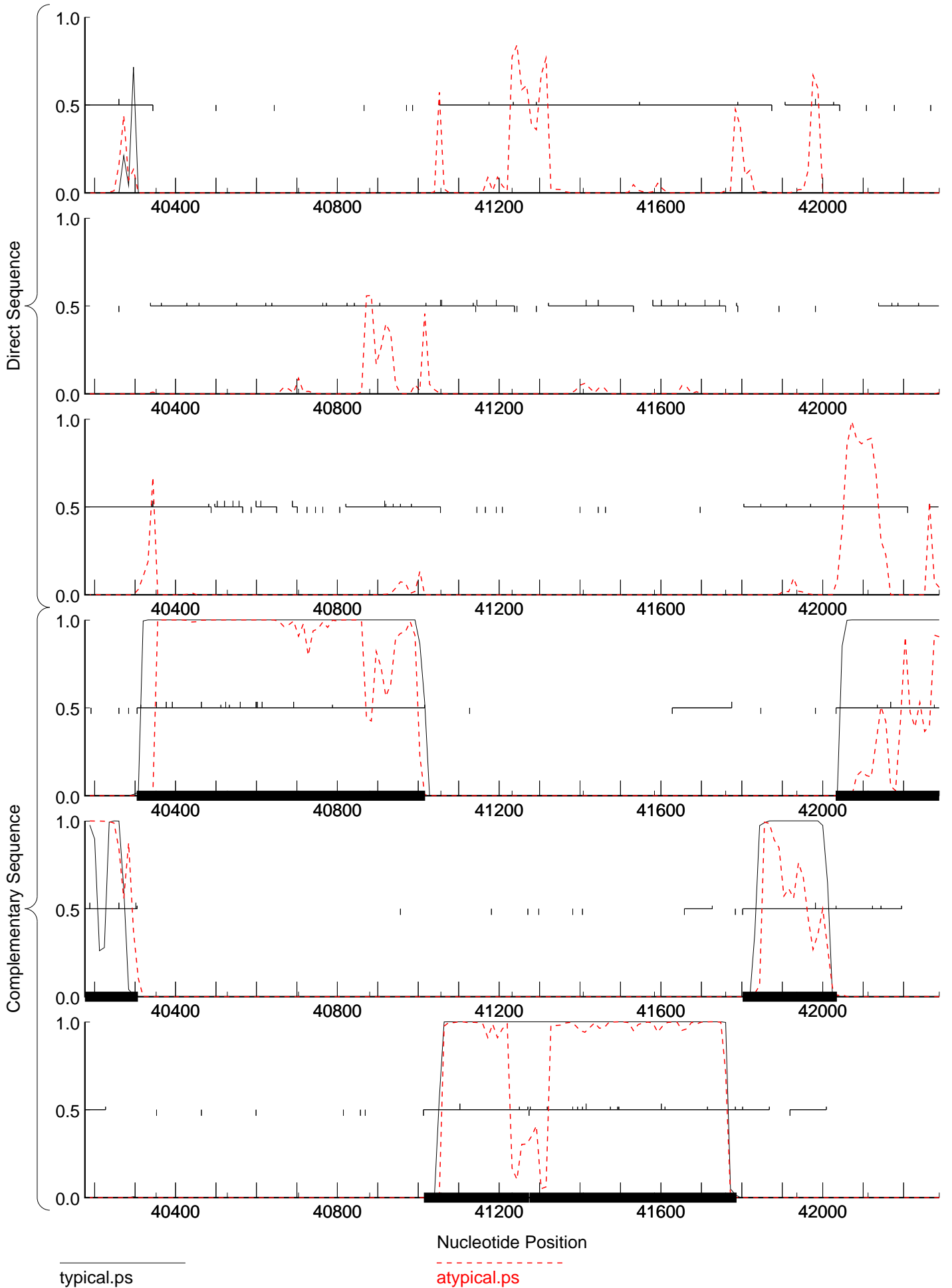


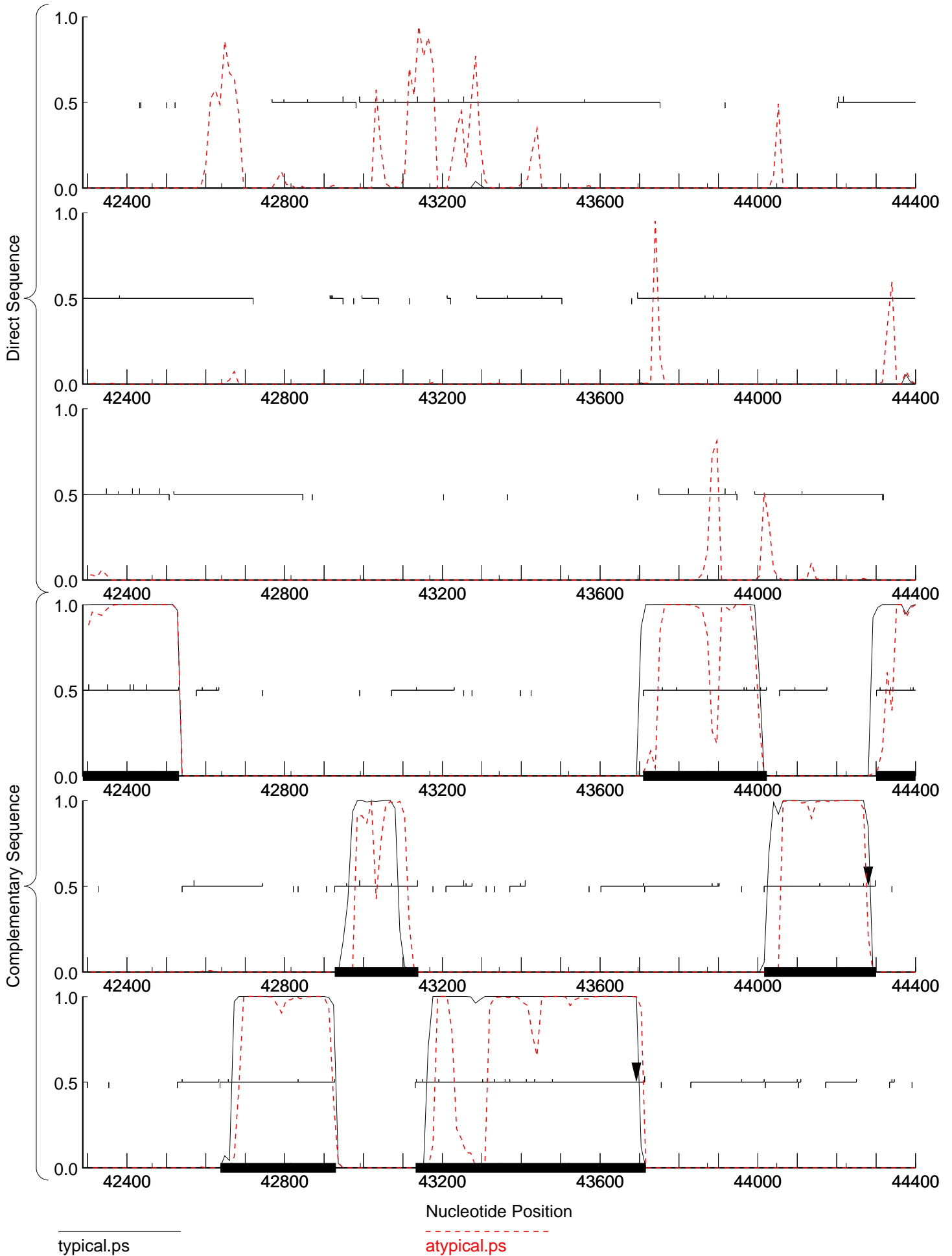


GeneMark.hmm prediction

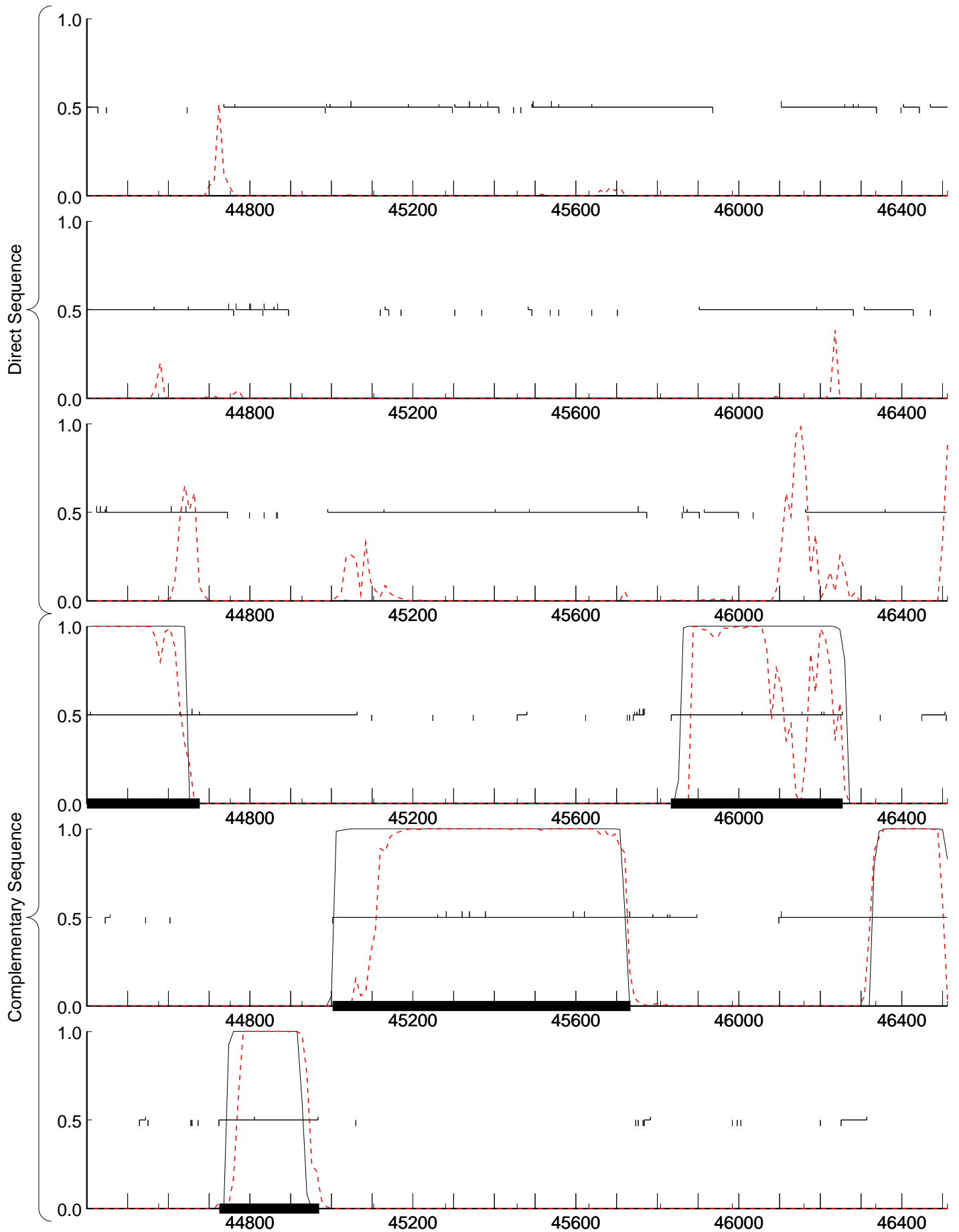






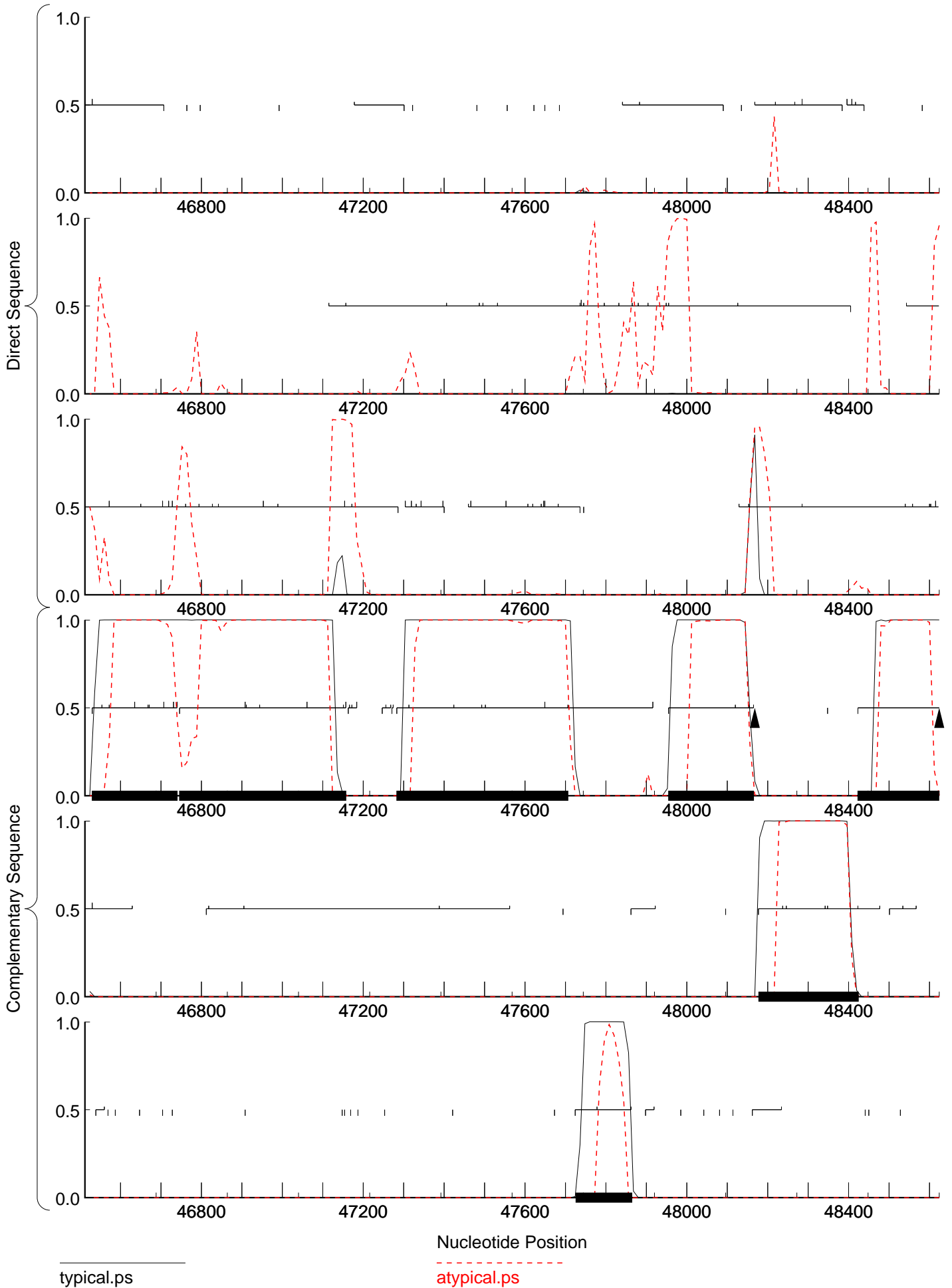


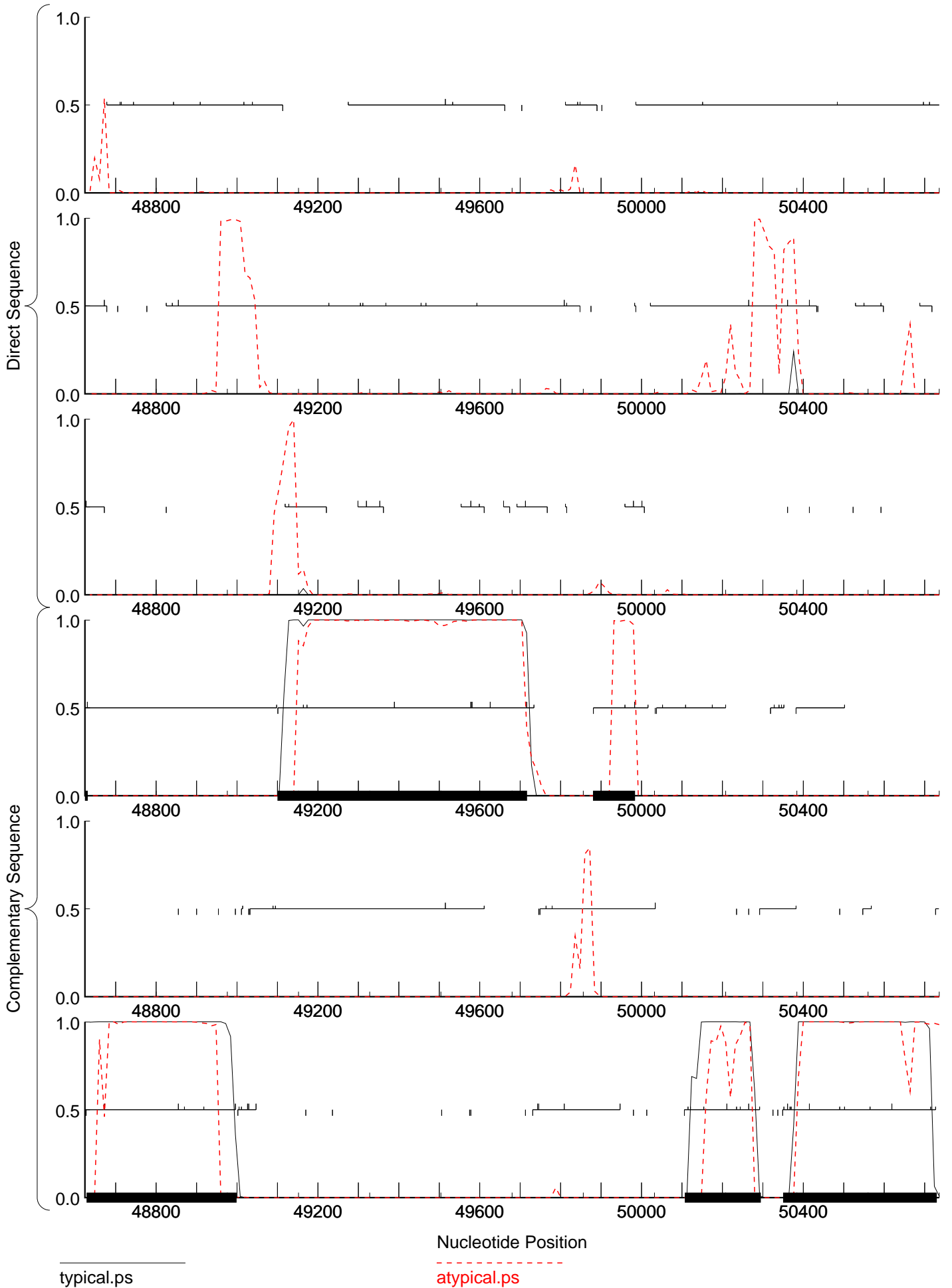
GeneMark.hmm prediction



typical.ps

atypical.ps





GeneMark.hmm prediction

