

GeneMark

Version 2.5p (09.08.06)

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PROGRAM INFORMATION

Sequence : Mycobacterium phage Centaur complete sequence, 53001 bp including 10 bp 3' overhang (CGGTCCG)
Analysis Date : 5/23/17 at 12:53:39
Pages : 27
Sequence Length : 53001 bp
GC Content : 63.50%

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,

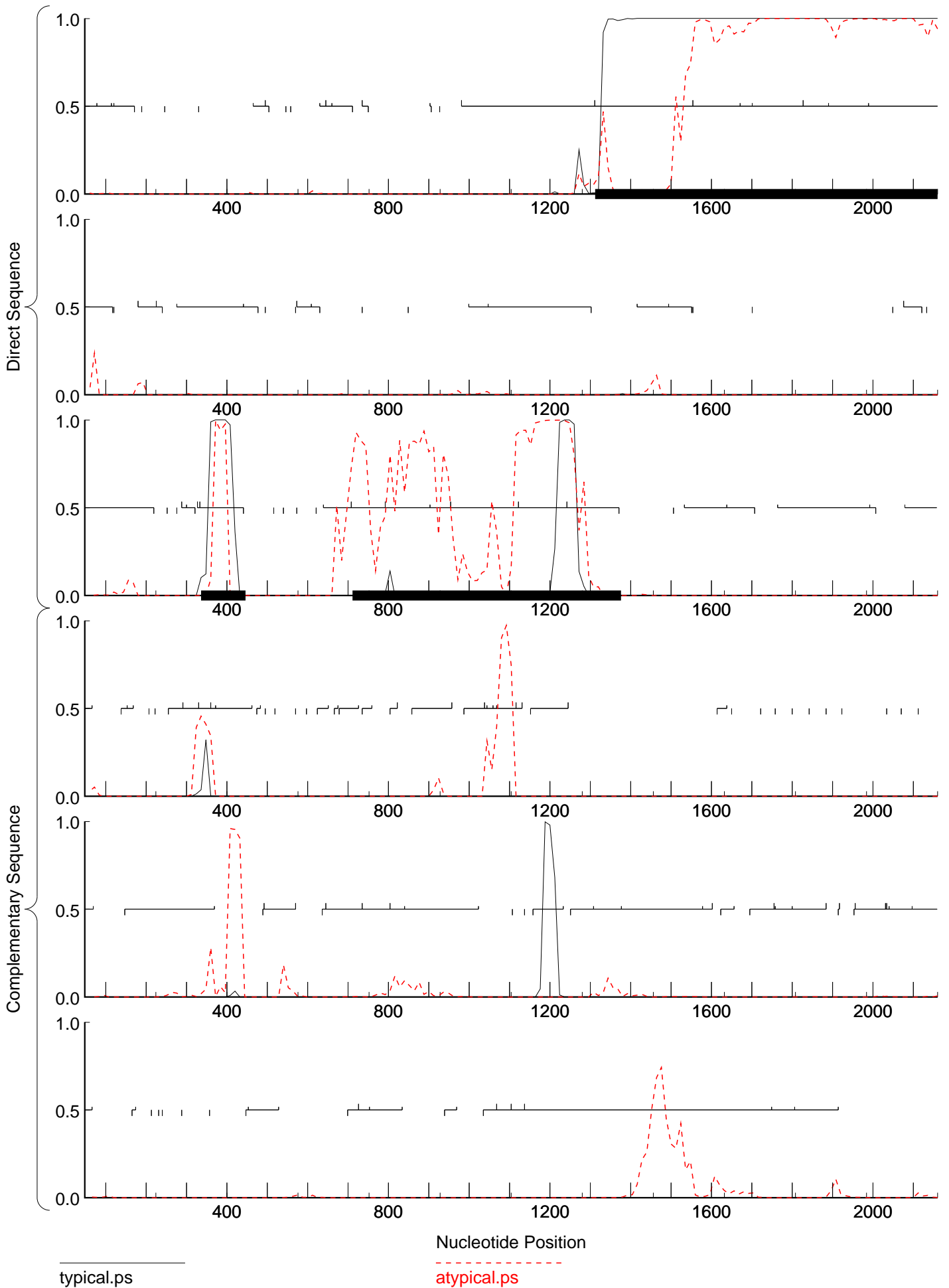
MATRIX INFORMATION

Matrix : -
Author : -
Order : 4

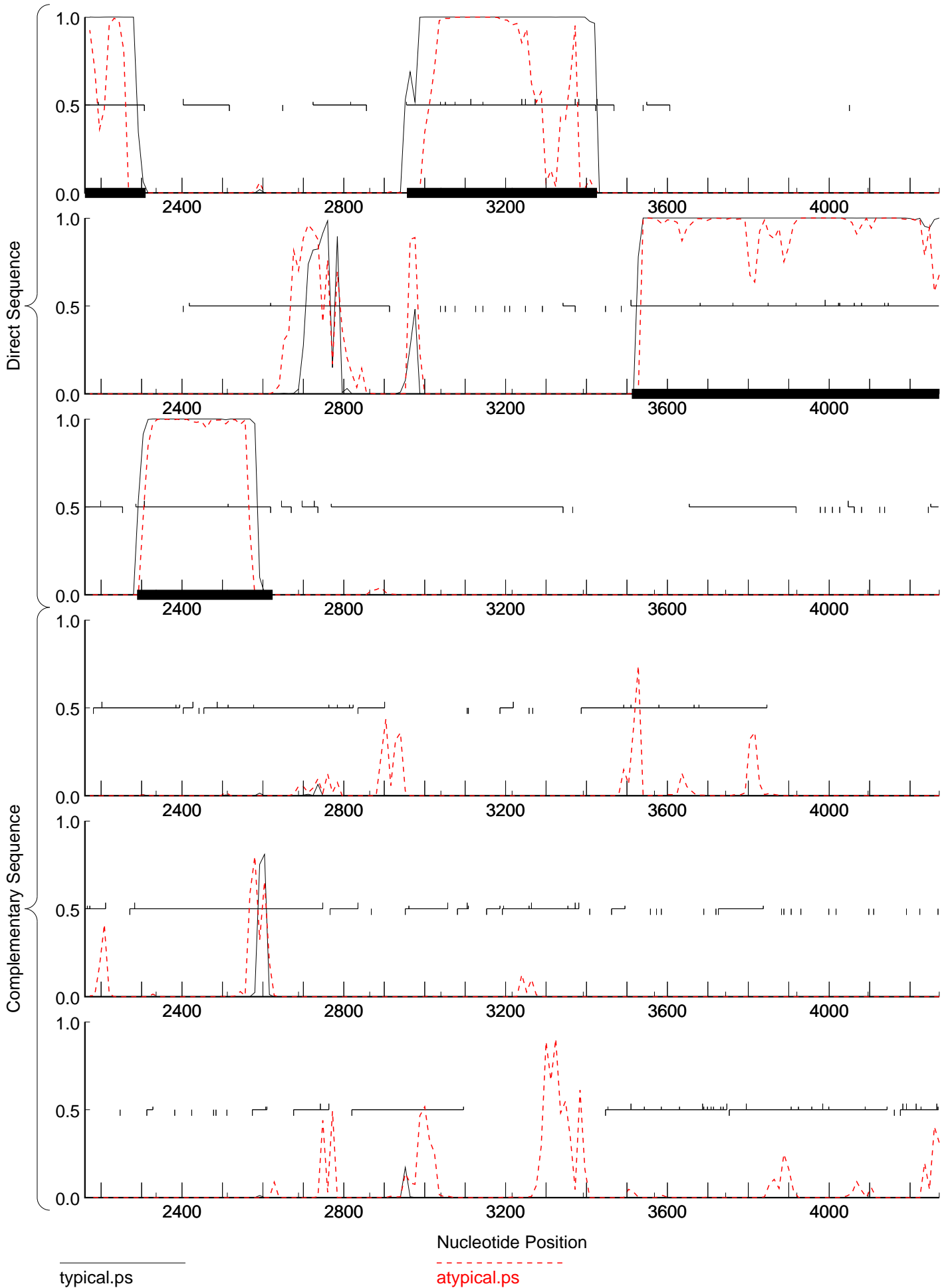
Send questions / comments to:
Dr. M. Borodovsky
Georgia Institute of Technology
School of Biology
Atlanta, GA 30332-0230

Matrix notes & comments

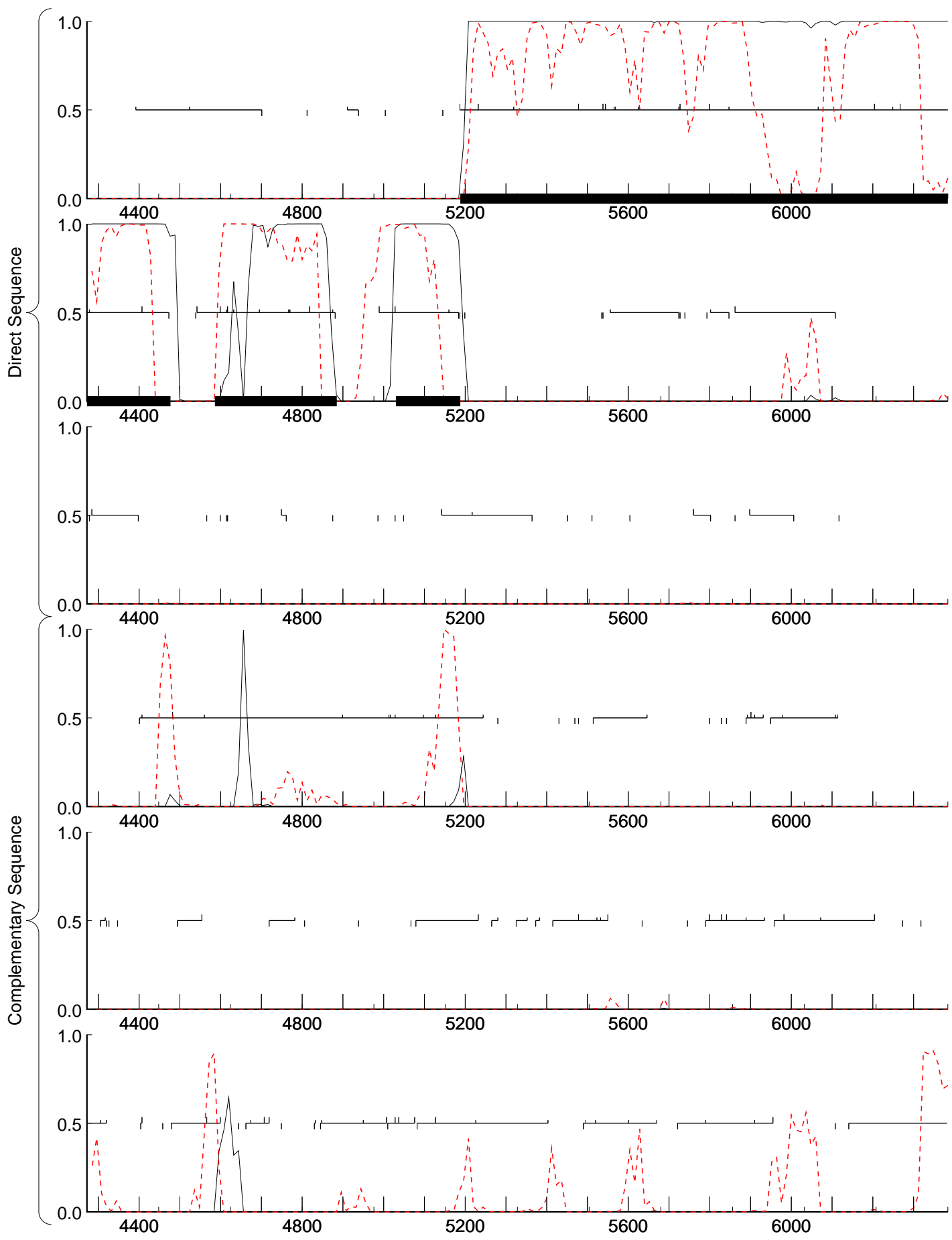
Training set derived by GeneMarkS, 4.30 October 2014
Tue May 23 12:53:39 2017



taur complete sequence, 53001 bp including 10 bp 3' overhang (CGGTCGGTTA), Cluster A2, Order 4, Window 96, Step 12, 3/27
Genemark.hmm prediction
taur complete sequence, 53001 bp including 10 bp 3' overhang (CGGTCGGTTA), Cluster A2, Order 2, Window 96, Step 12, 3/27

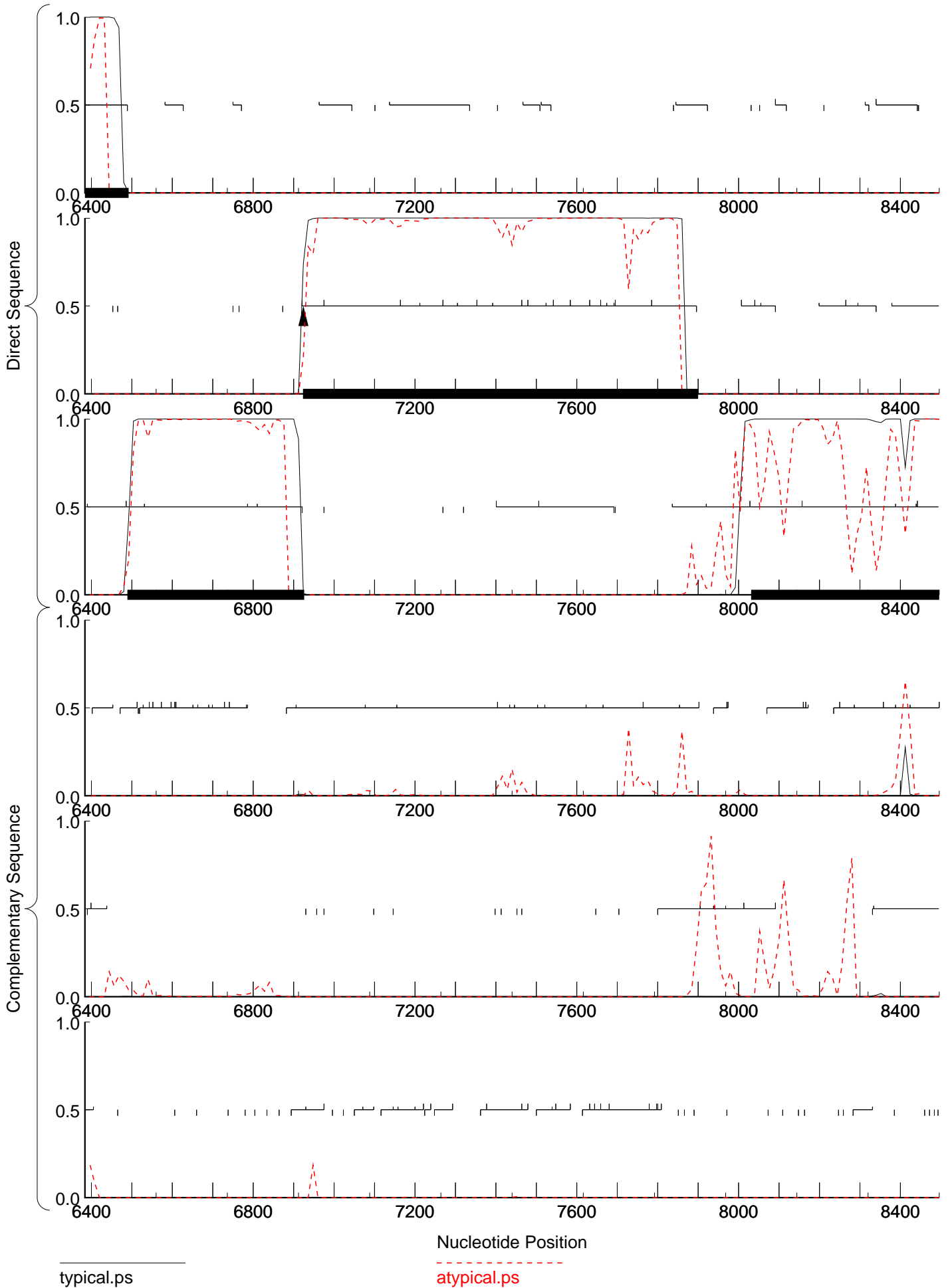


taur complete sequence, 53001 bp including 10 bp 3' overhang (CGTTCGGTTA), Cluster A2, Order 4, Window 96, Step 12, 4/27
Genemark.hmm prediction
taur complete sequence, 53001 bp including 10 bp 3' overhang (CGTTCGGTTA), Cluster A2, Order 2, Window 96, Step 12, 4/27

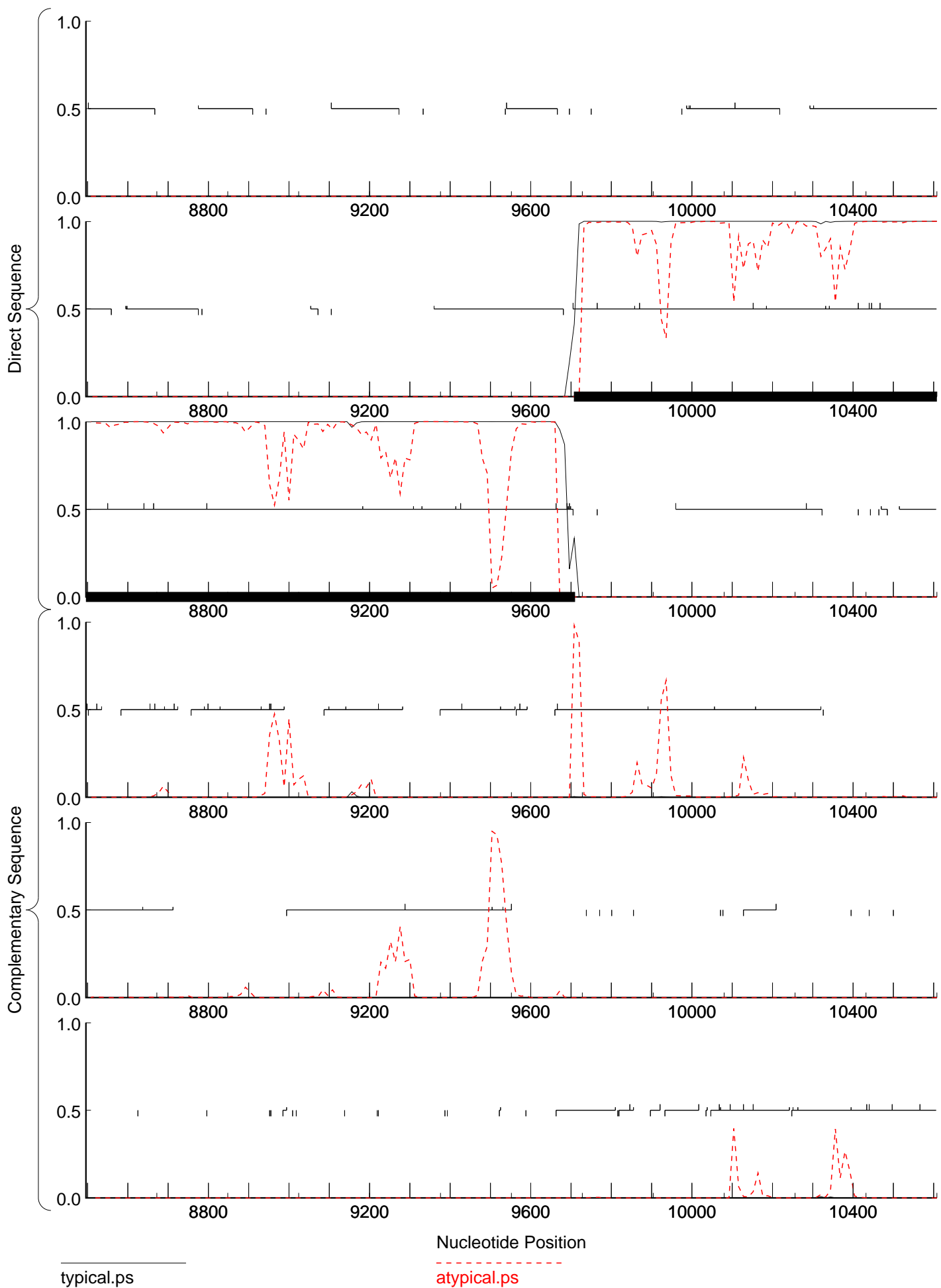


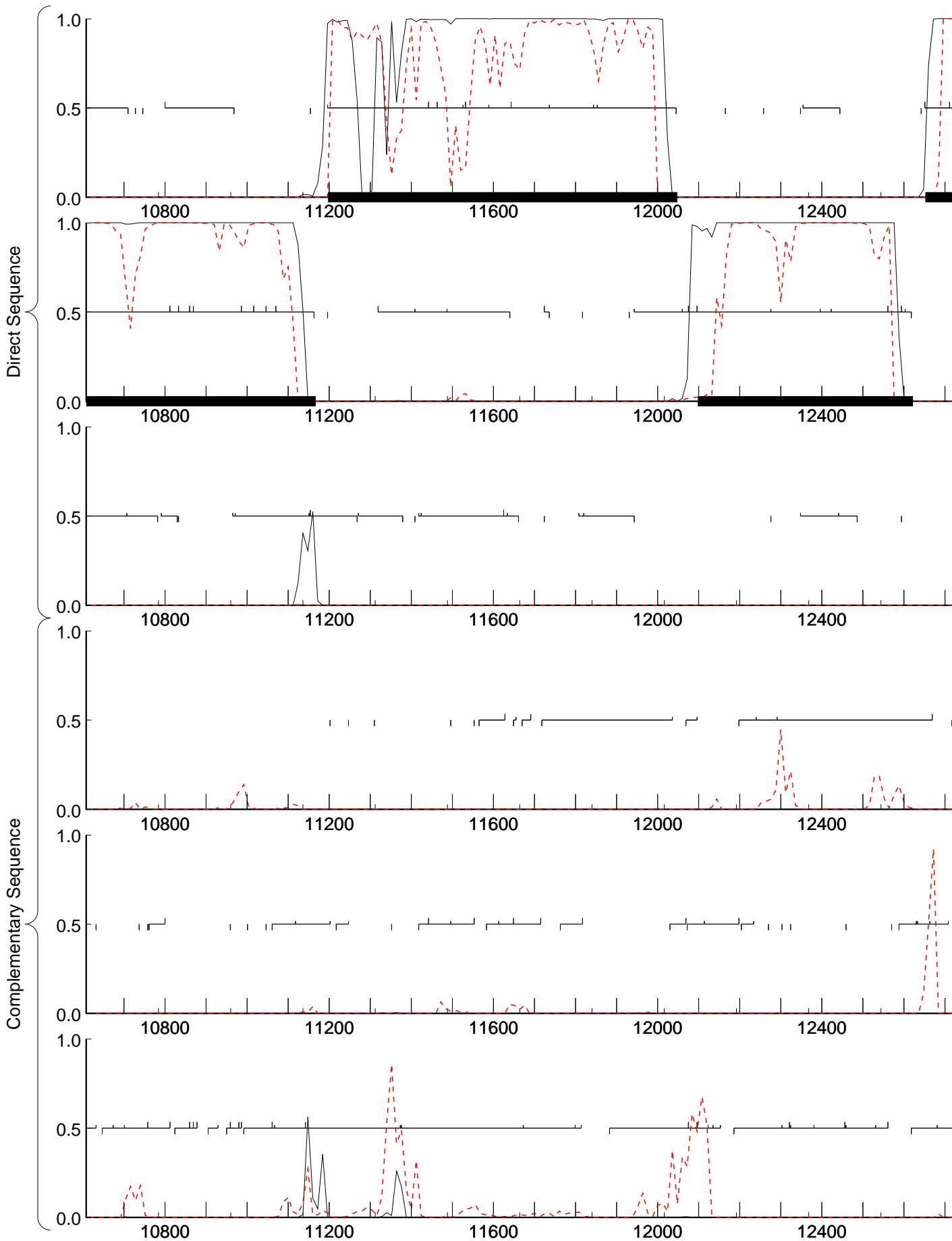
typical.ps

atypical.ps



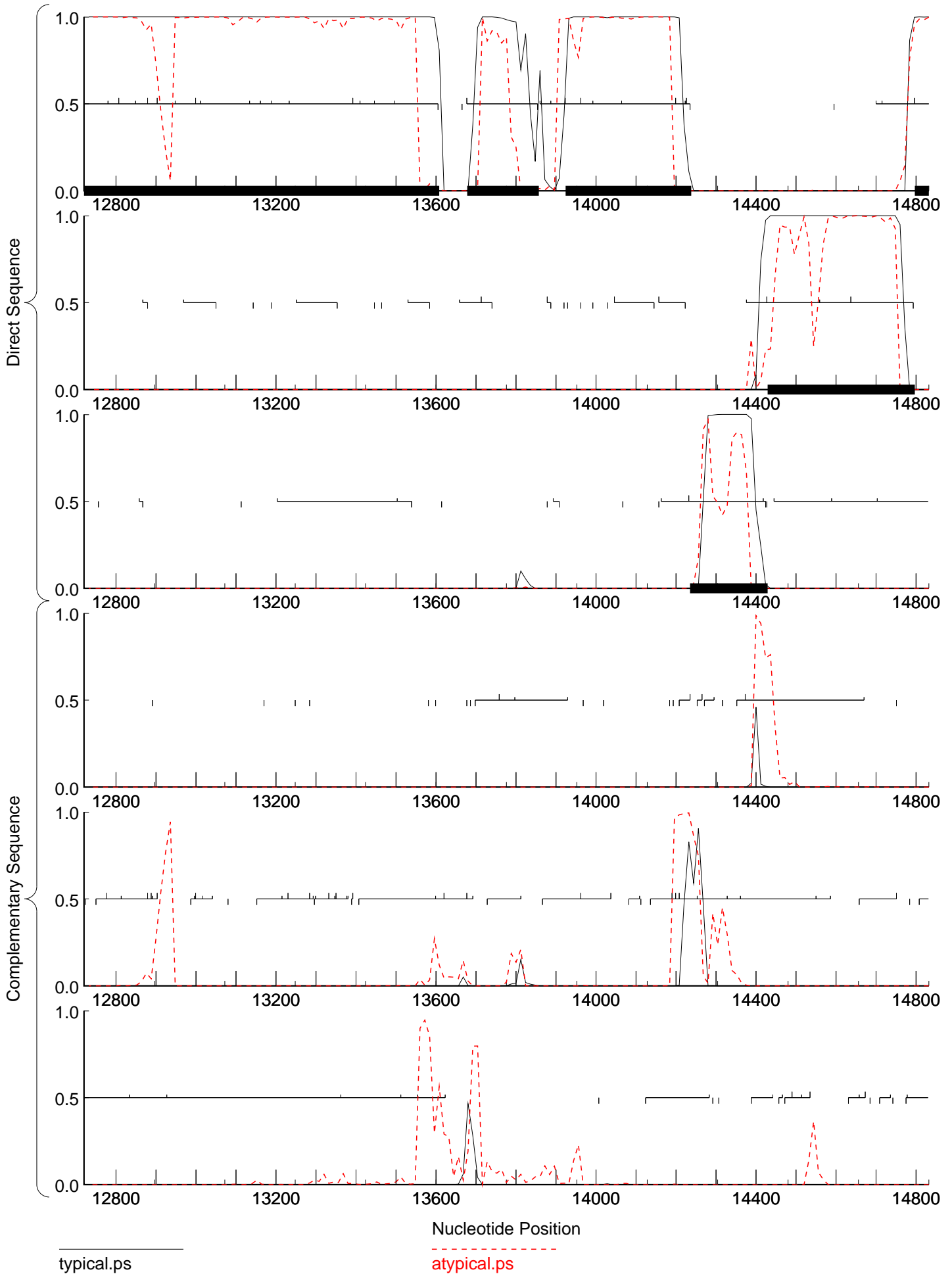
taur complete sequence, 53001 bp including 10 bp 3' overhang (CGGTCGGTTA), Cluster A2, Order 4, Window 96, Step 12, 6/27
Genemark.hmm prediction
taur complete sequence, 53001 bp including 10 bp 3' overhang (CGGTCGGTTA), Cluster A2, Order 2, Window 96, Step 12, 6/27



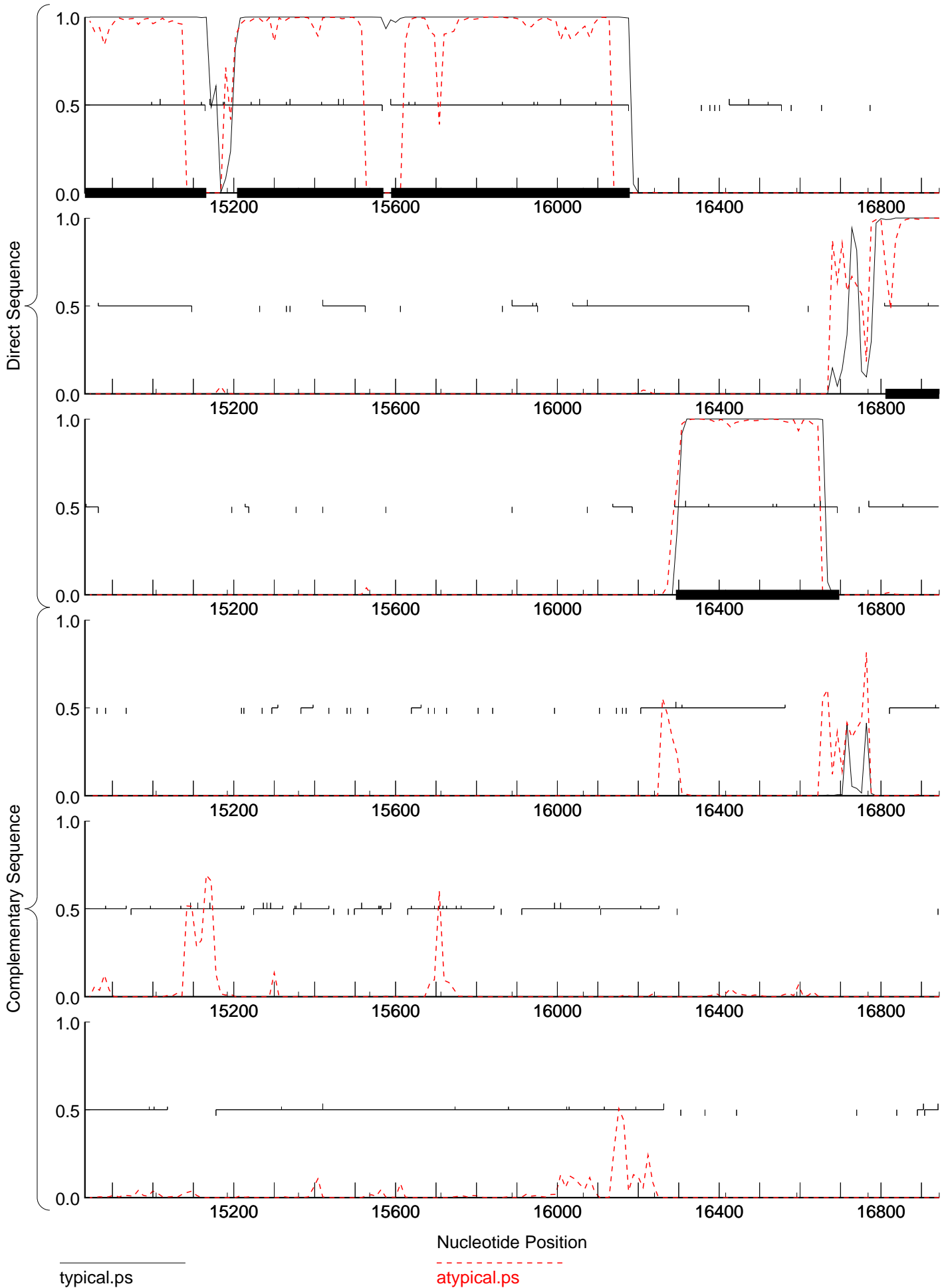


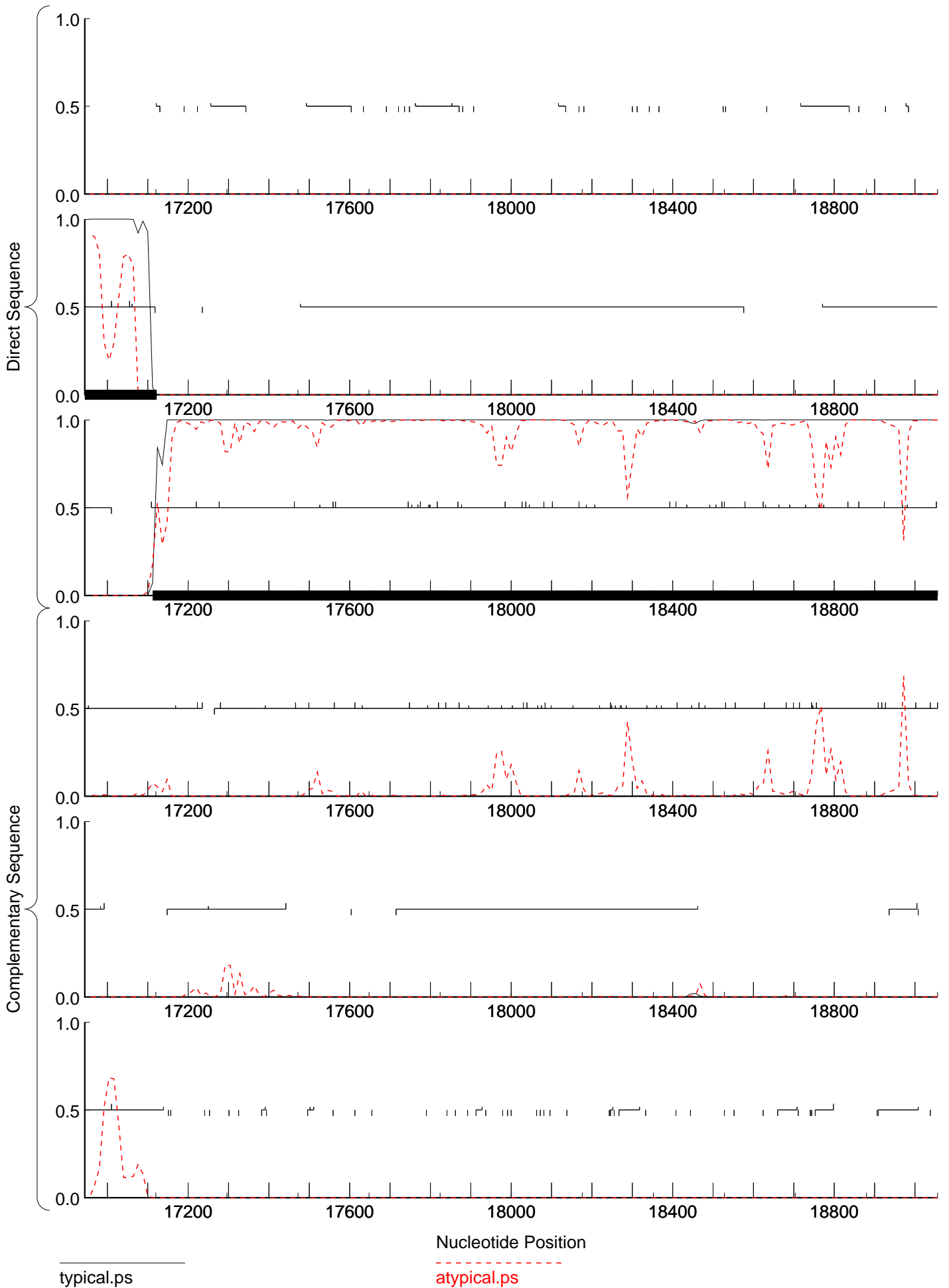
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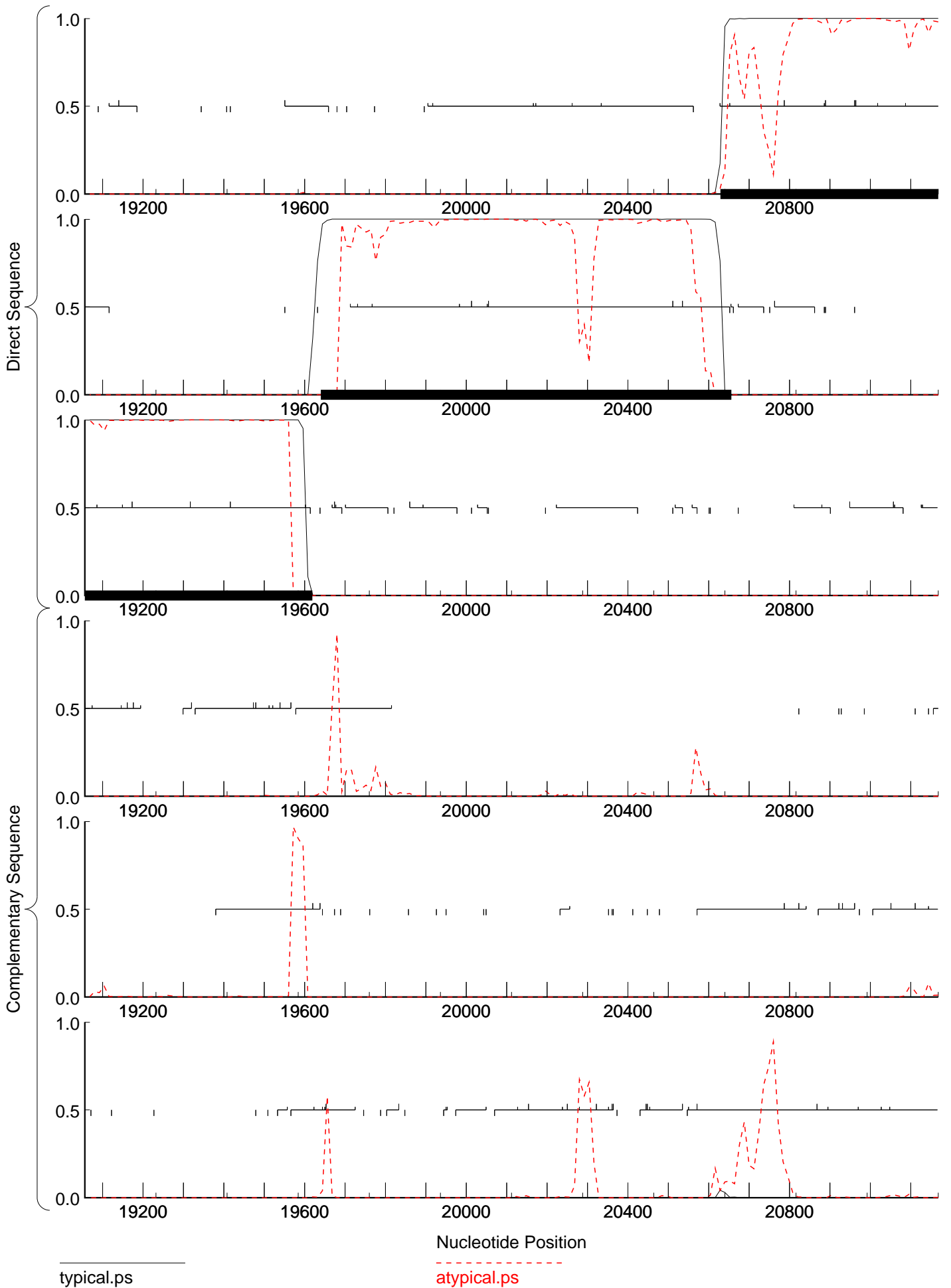
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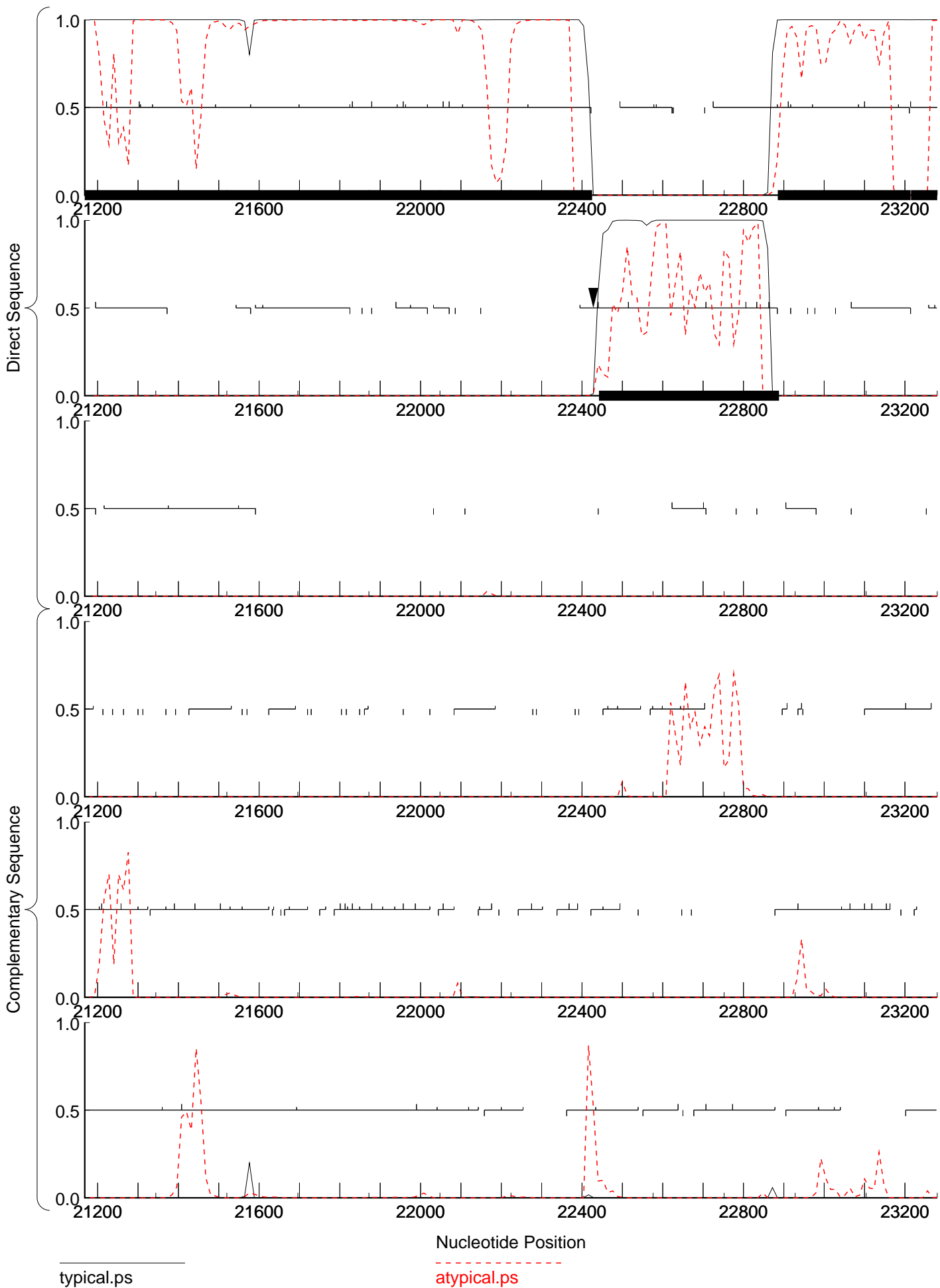


taur complete sequence, 53001 bp including 10 bp 3' overhang (CGGTCGGTTA), Cluster A2, Order 4, Window 96, Step 12, 9/27
Genemark.hmm prediction
taur complete sequence, 53001 bp including 10 bp 3' overhang (CGGTCGGTTA), Cluster A2, Order 2, Window 96, Step 12, 9/27



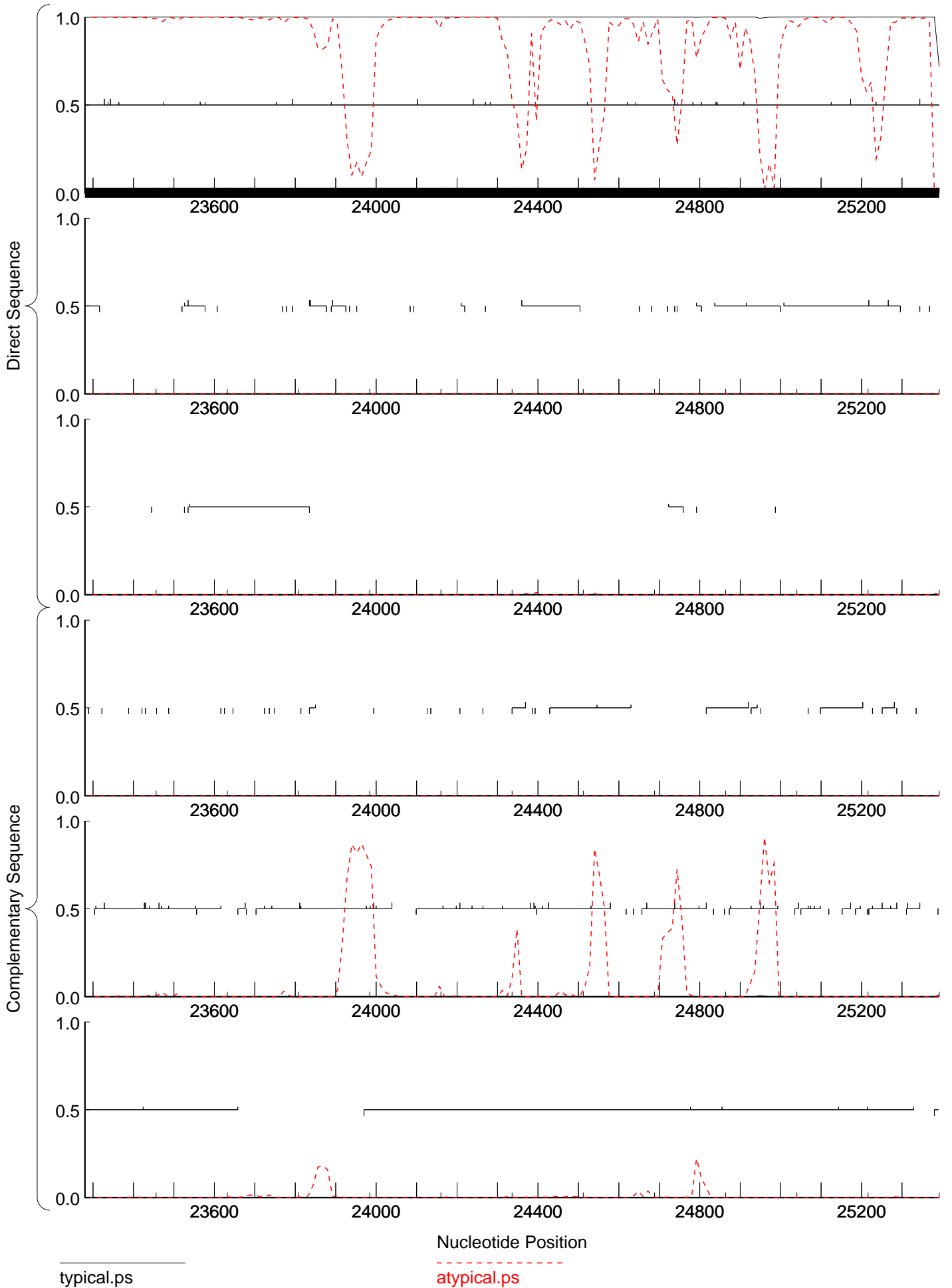


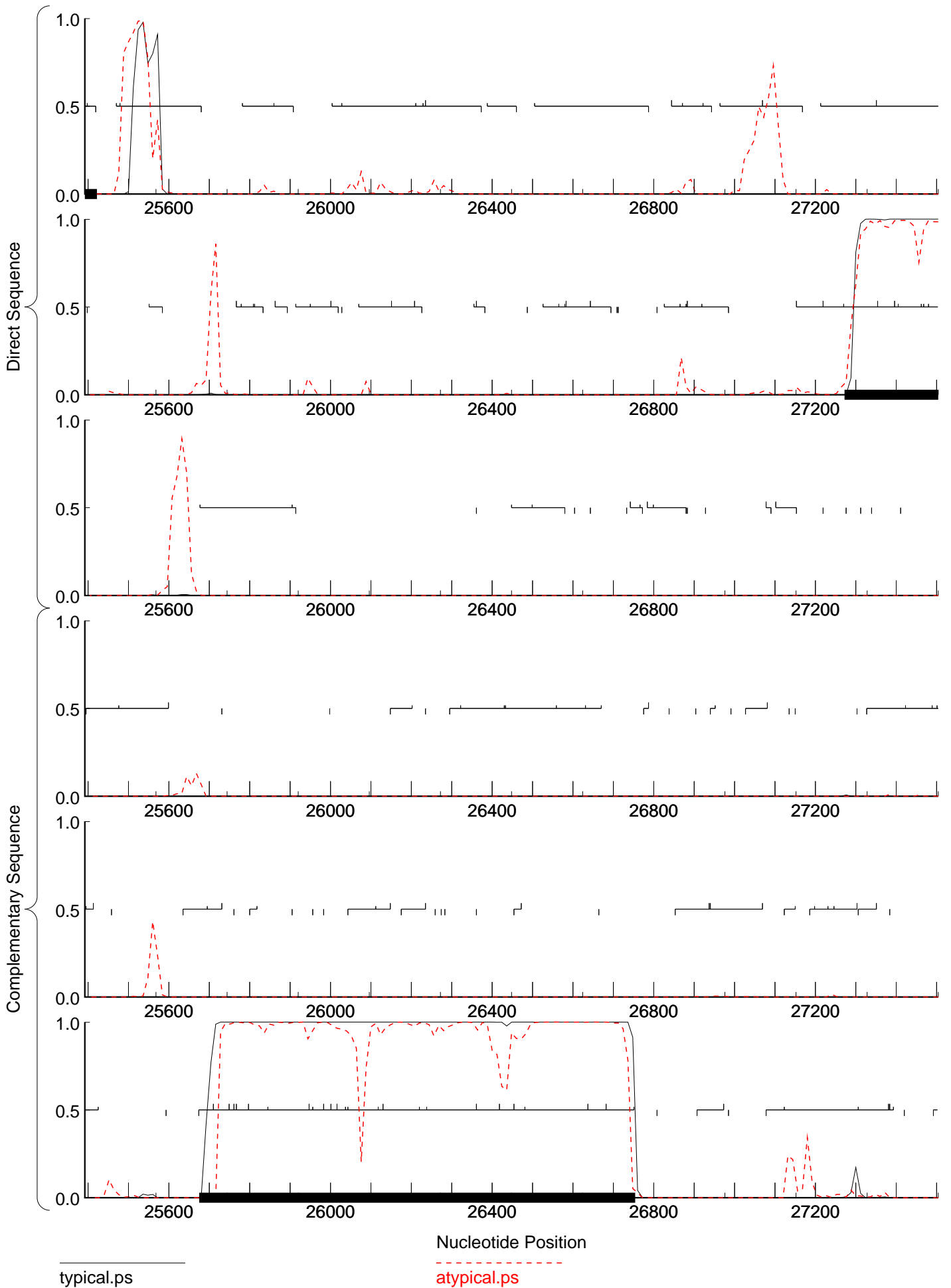


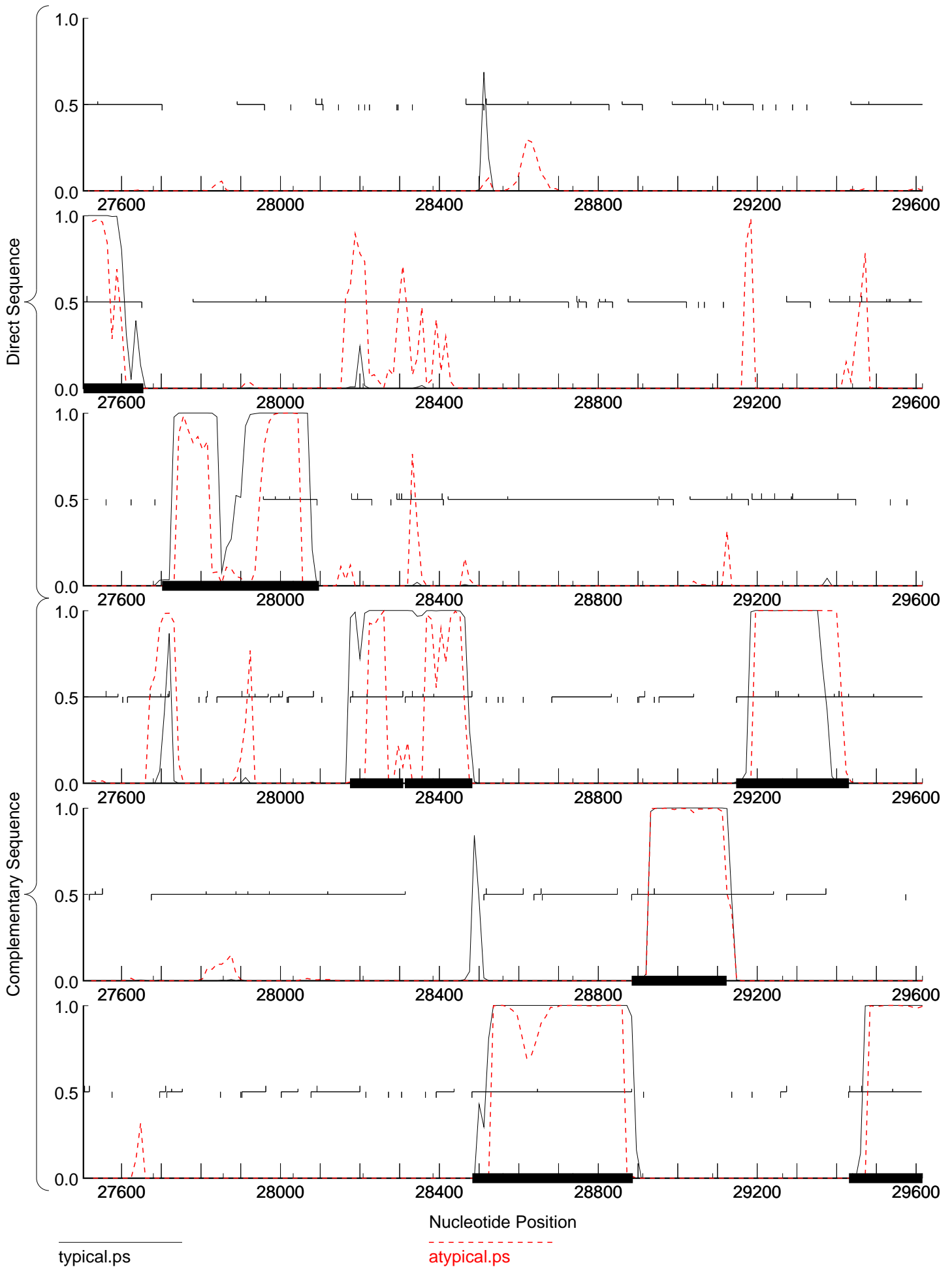


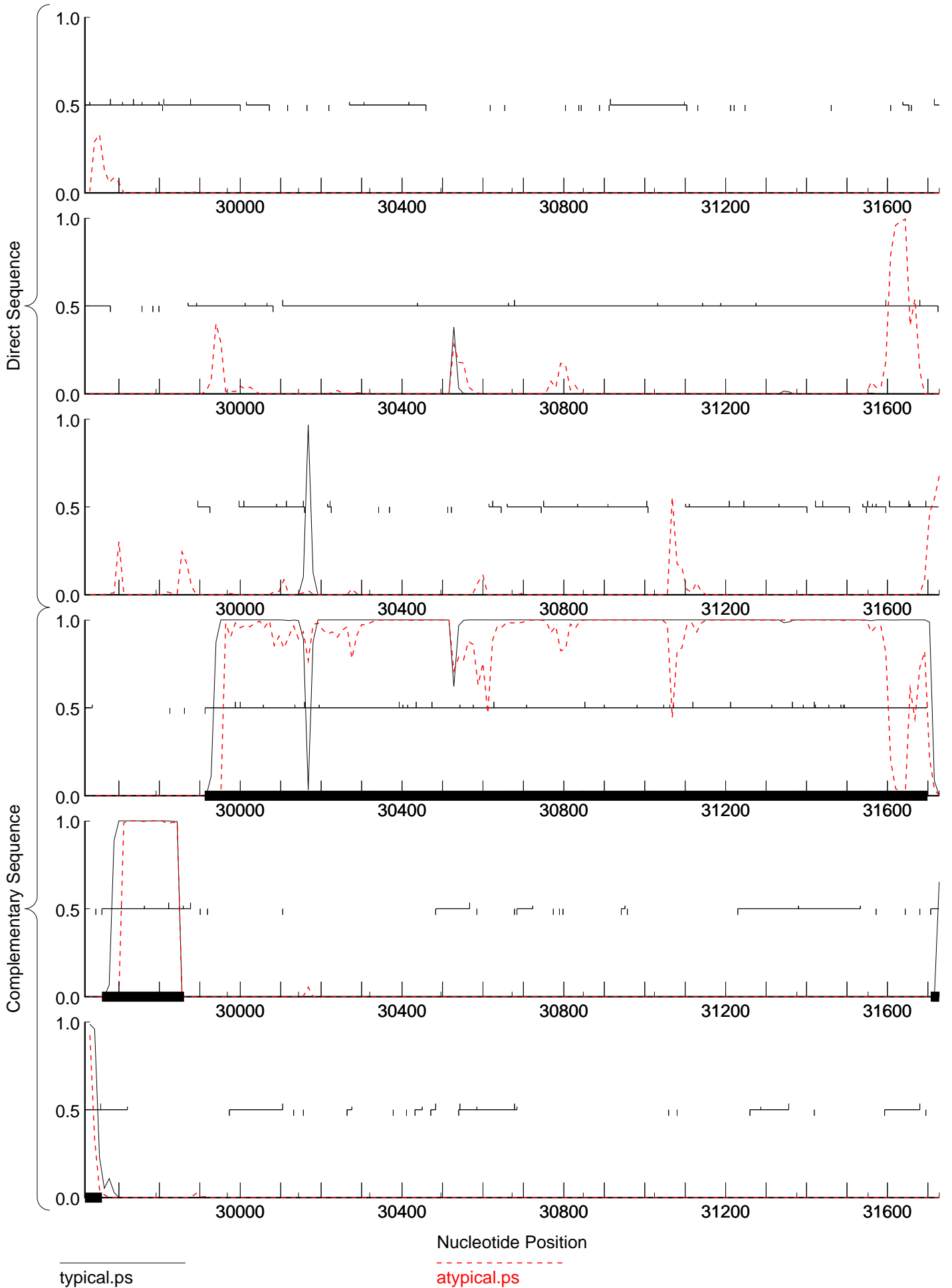
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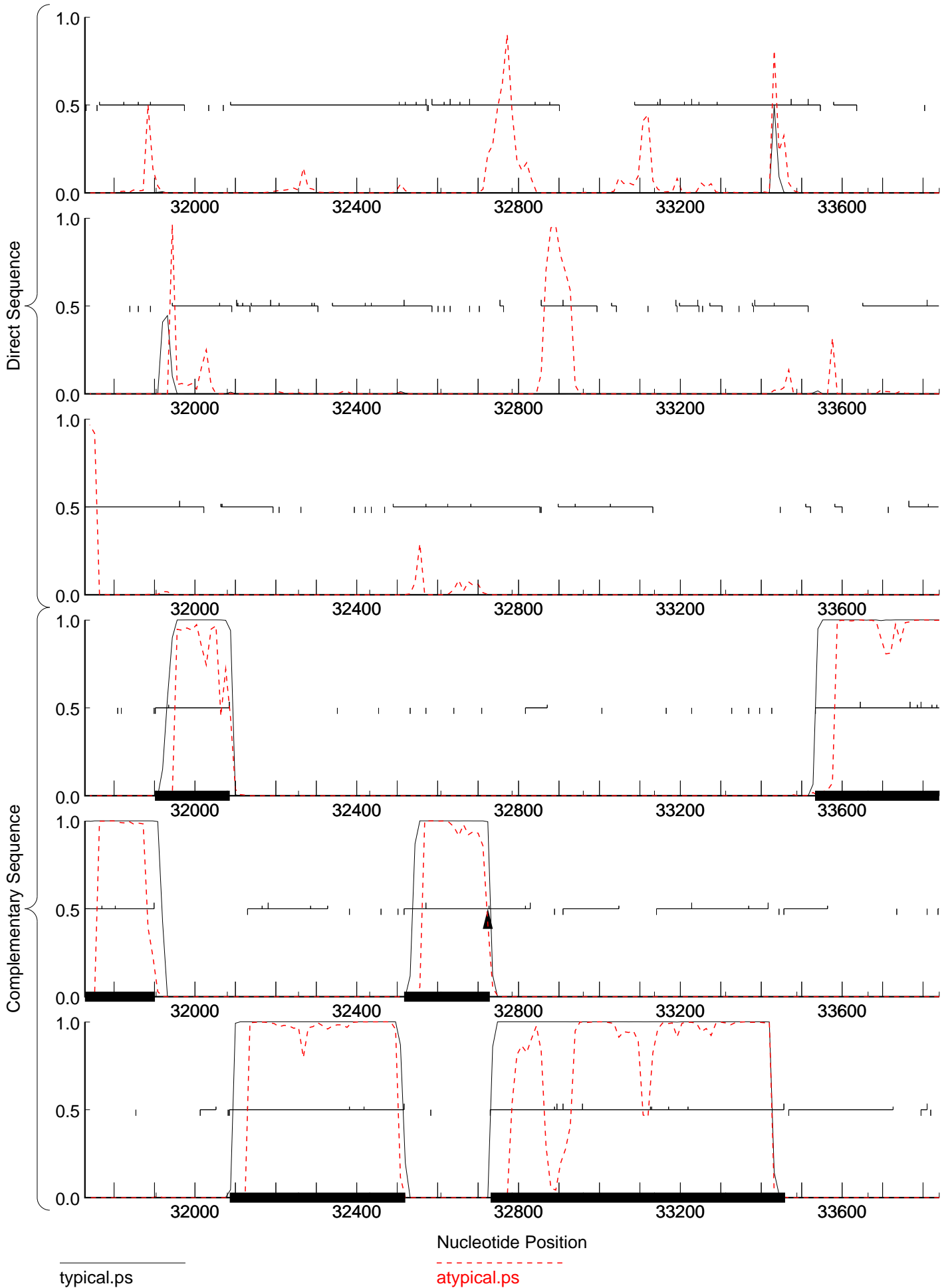
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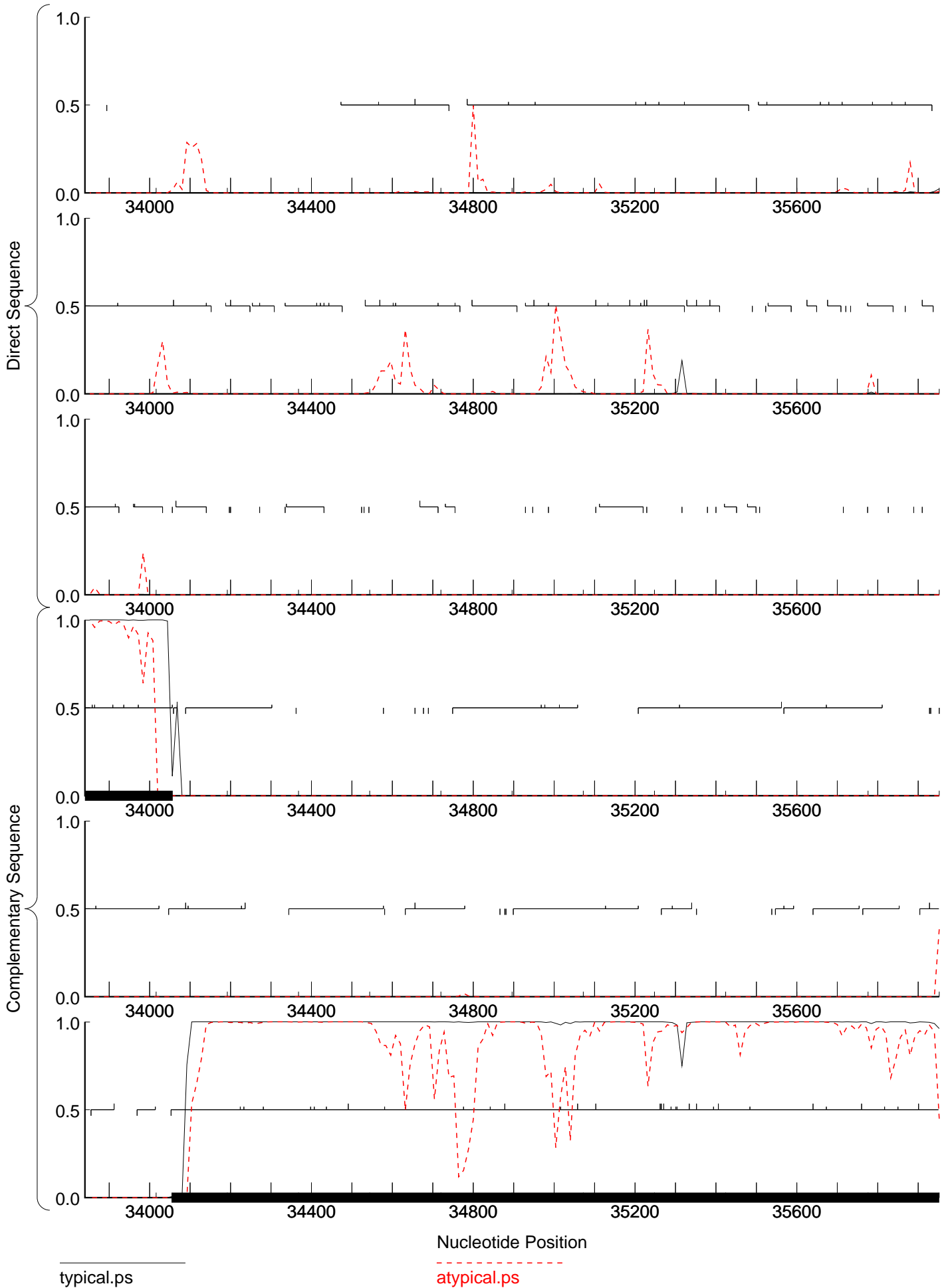


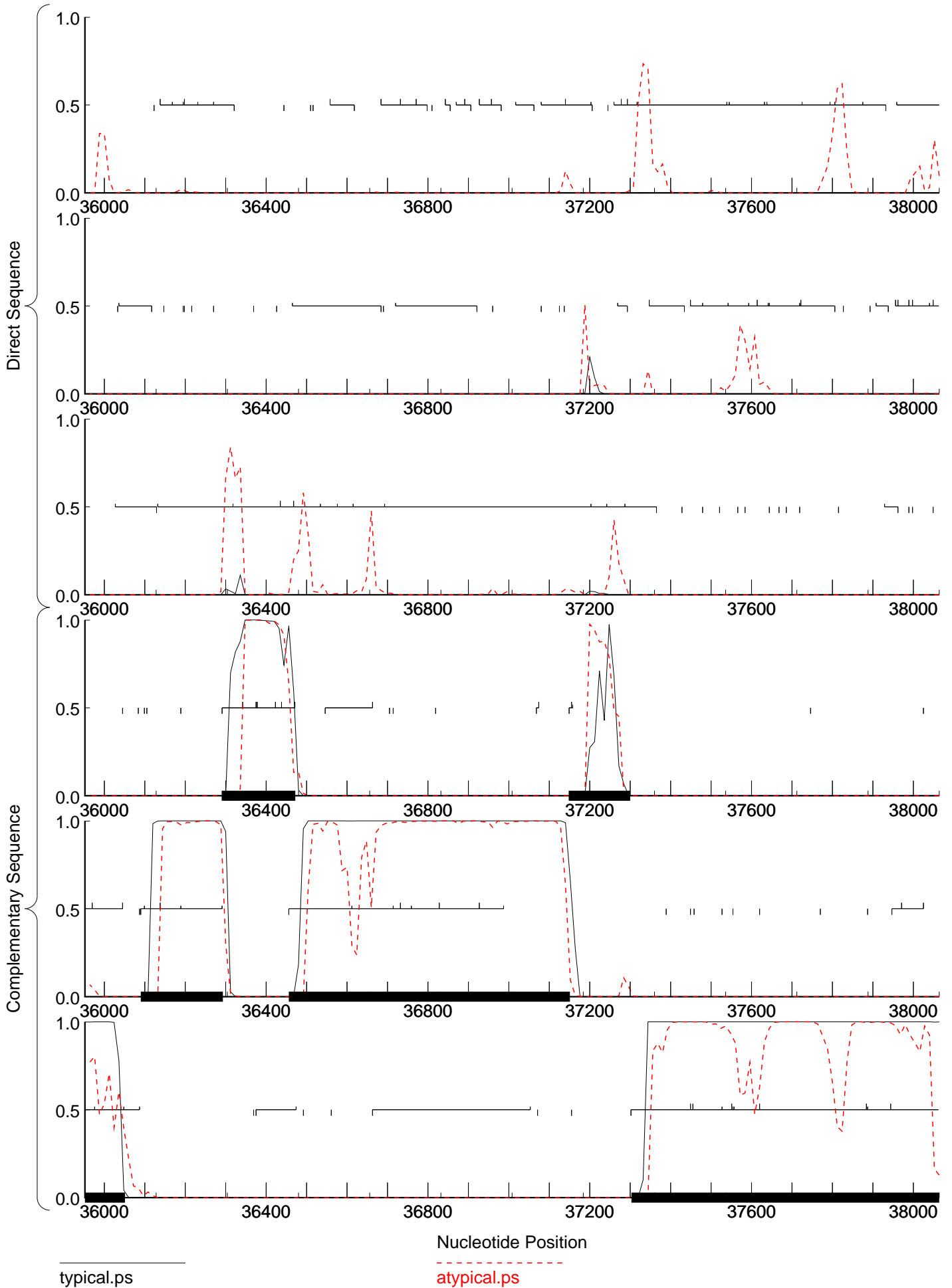


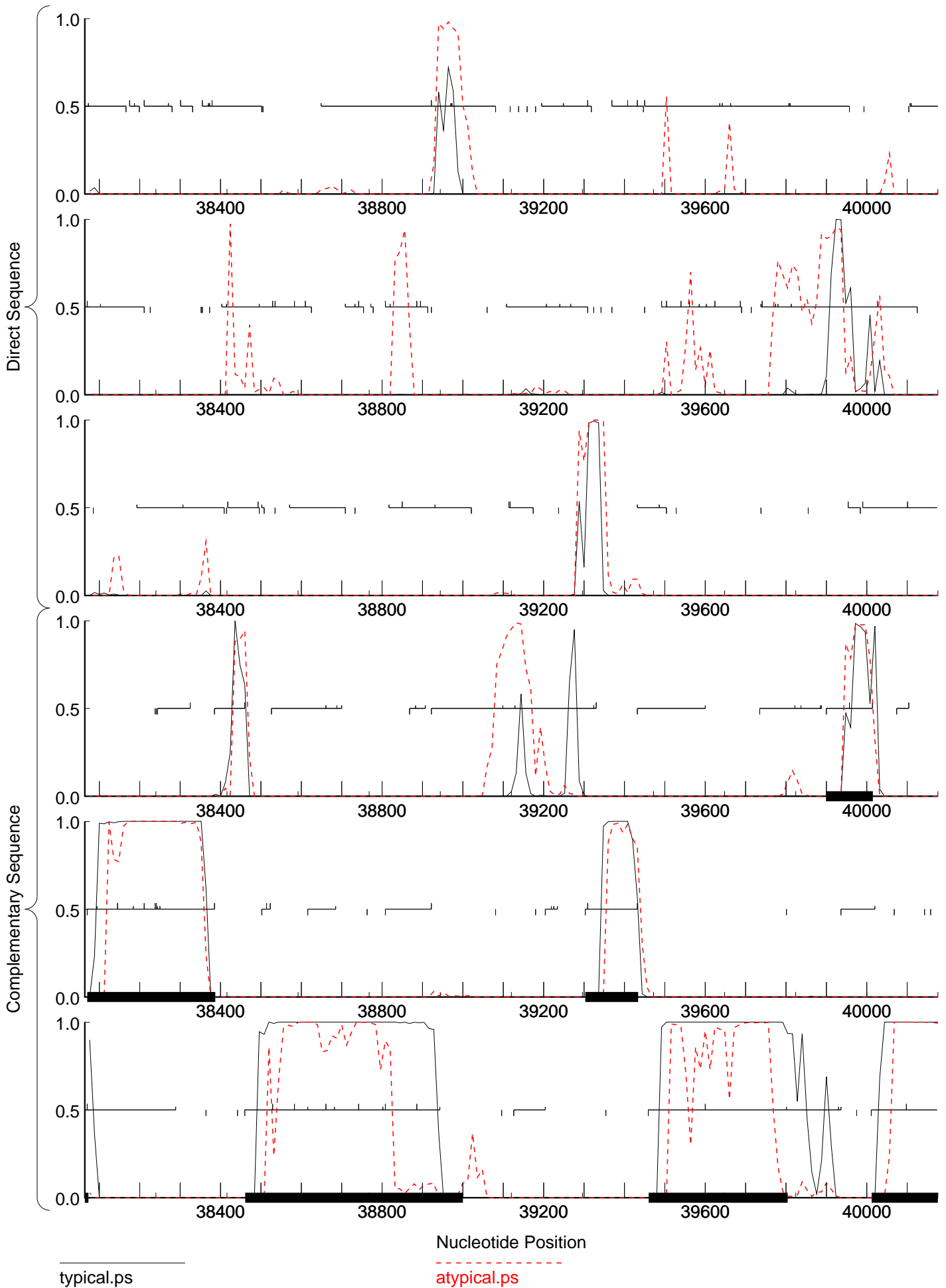












typical.ps

atypical.ps

