

GeneMark

Version 2.5p (09.08.06)

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PROGRAM INFORMATION

Sequence : Mycobacterium phage Yecey3 complete sequence, 53277 bp including 10-base 3' overhang (CGGAC
Analysis Date : 8/20/17 at 11:31:07
Pages : 27
Sequence Length : 53277 bp
GC Content : 62.86%

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,

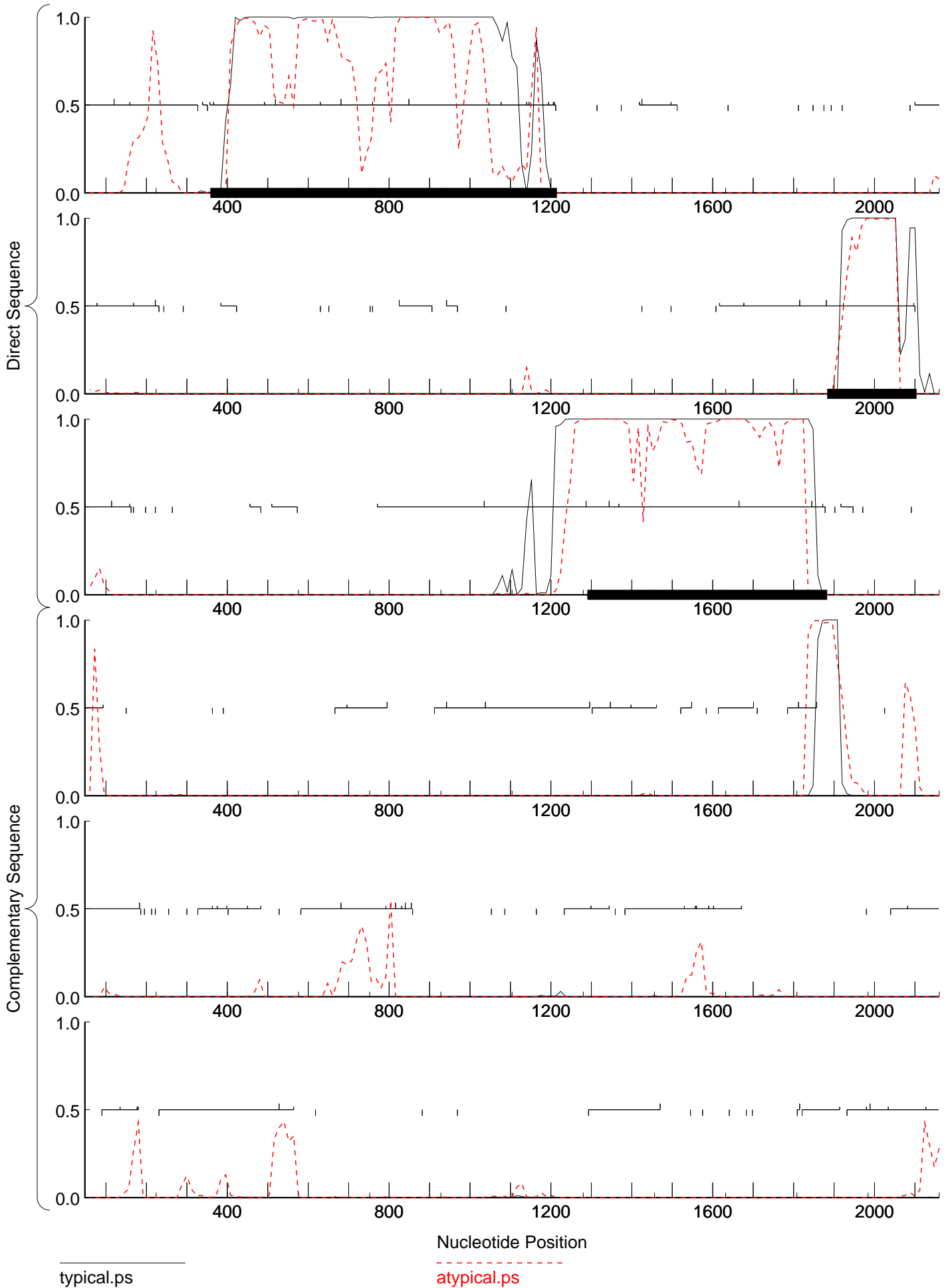
Matrix notes & comments

Training set derived by GeneMarkS, 4.30 October 2014
Sun Aug 20 11:31:07 2017

MATRIX INFORMATION

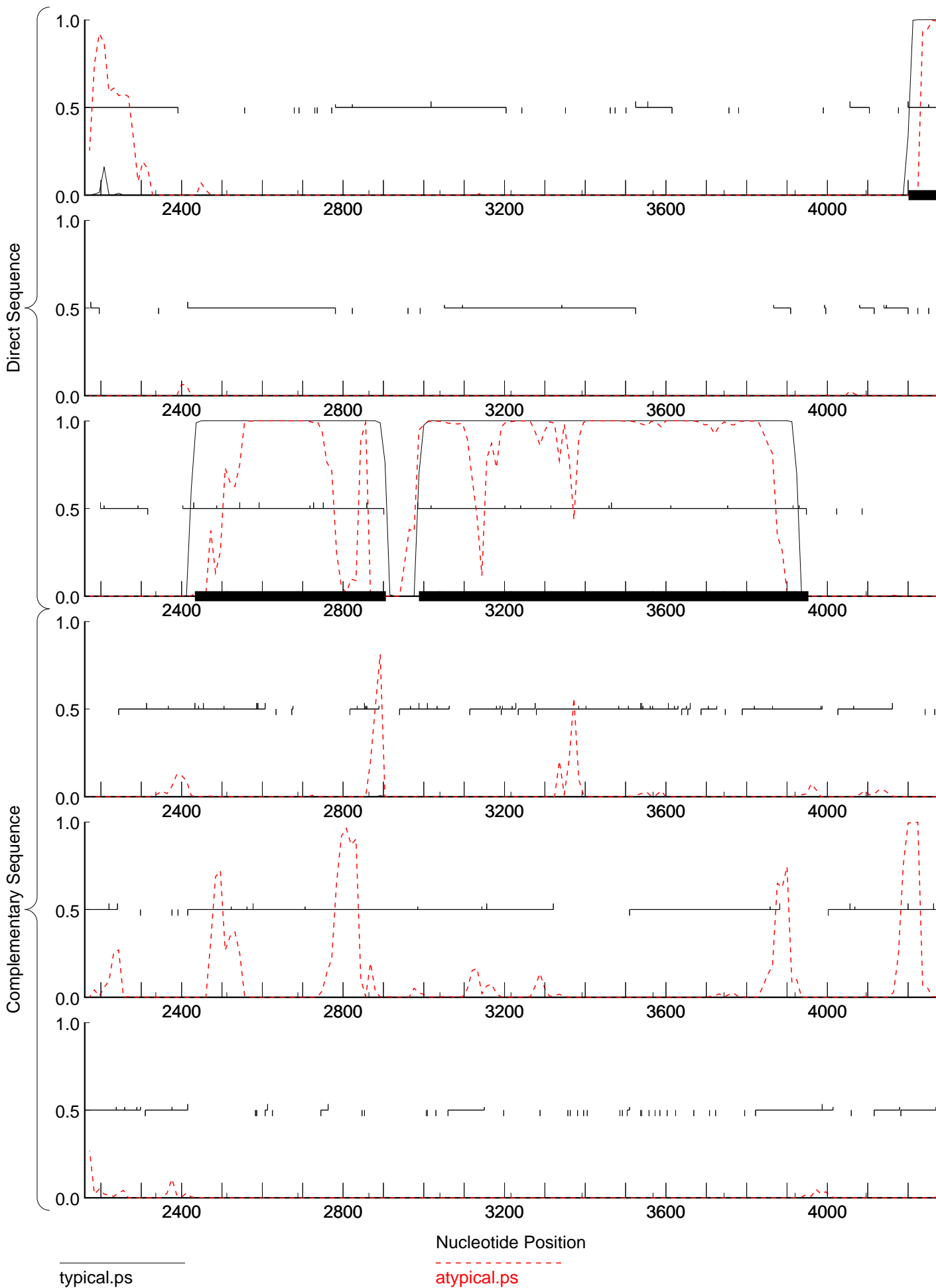
Matrix : -
Author : -
Order : 4

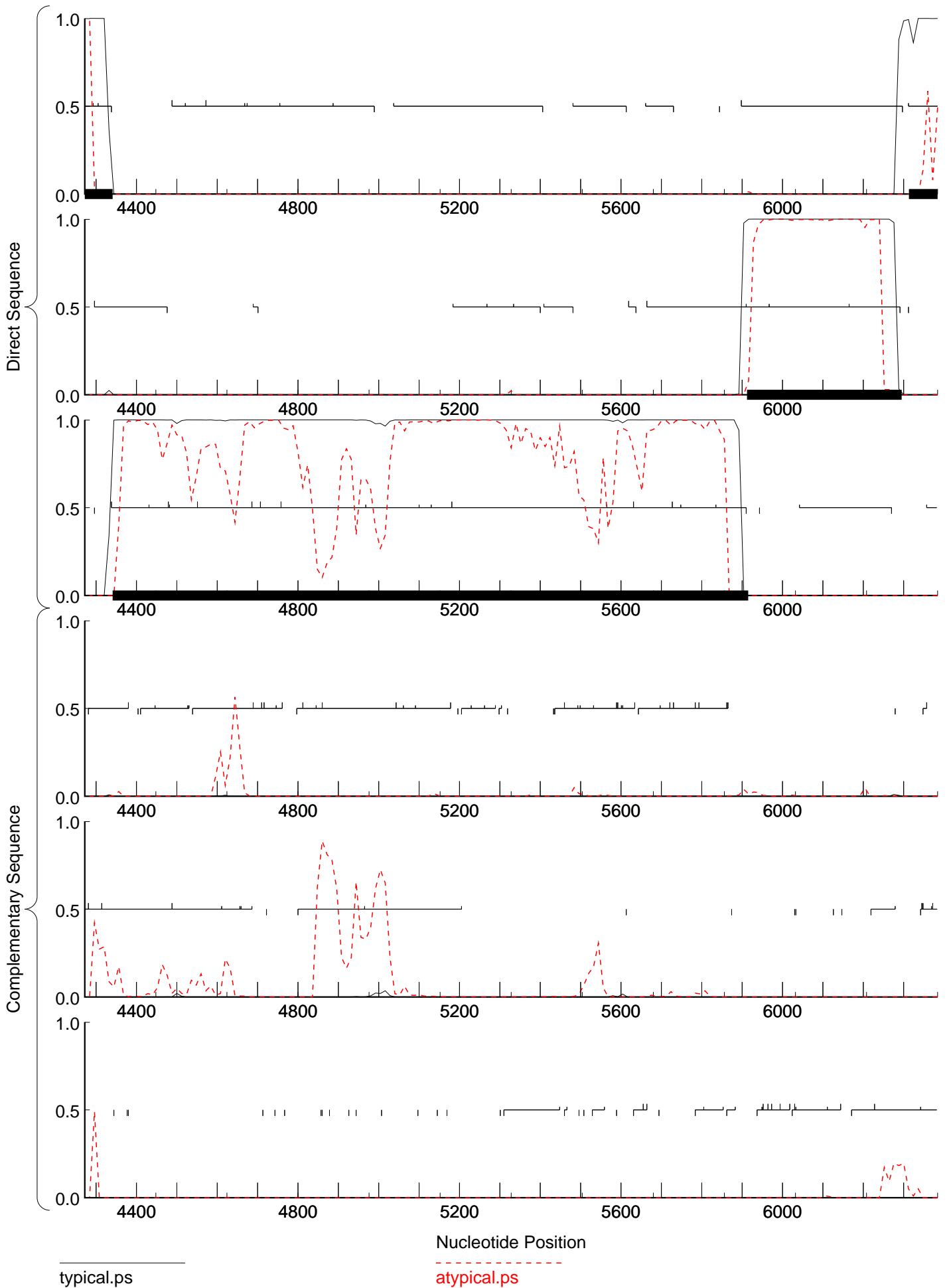
Send questions / comments to:
Dr. M. Borodovsky
Georgia Institute of Technology
School of Biology
Atlanta, GA 30332-0230



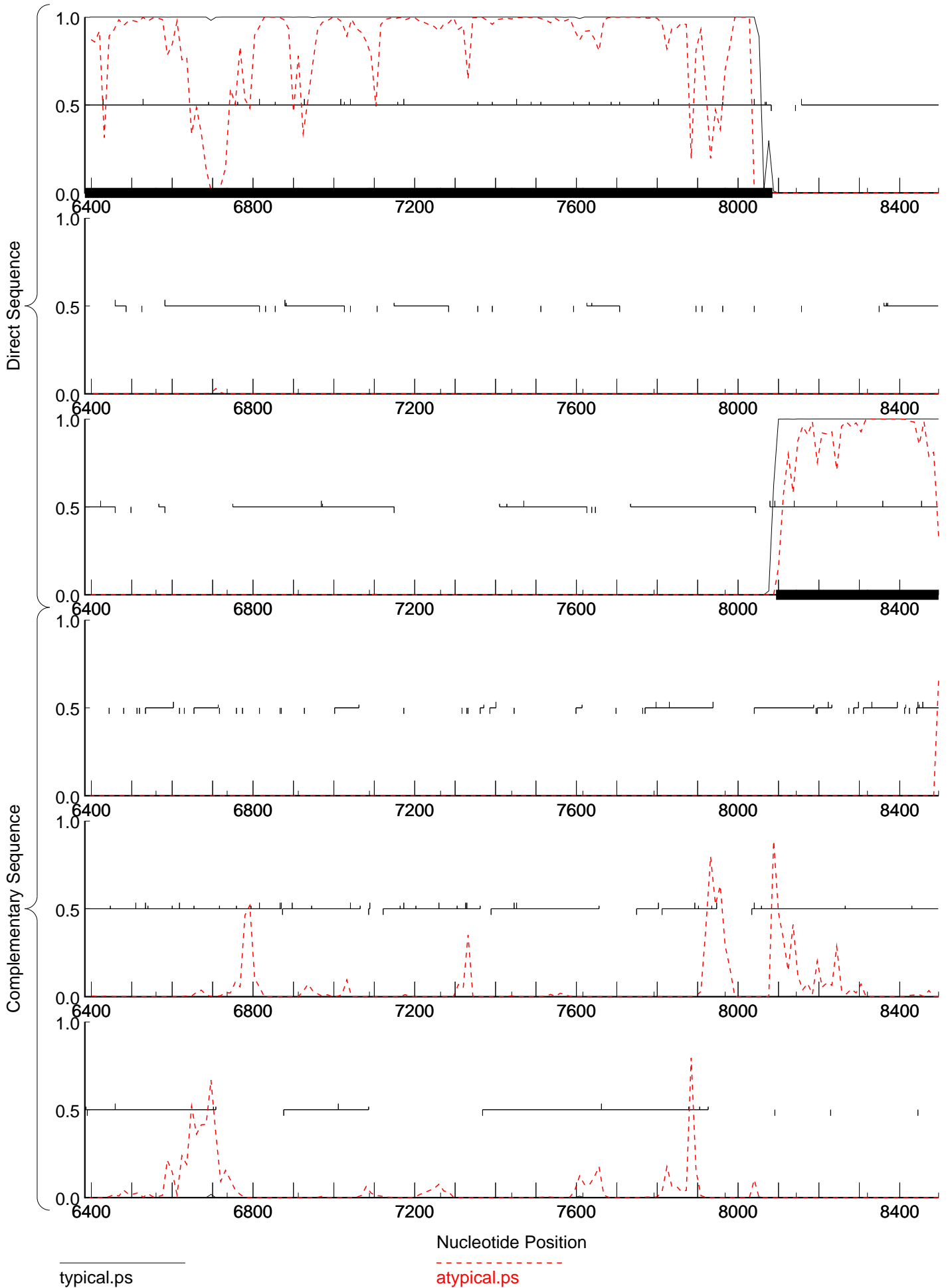
typical.ps

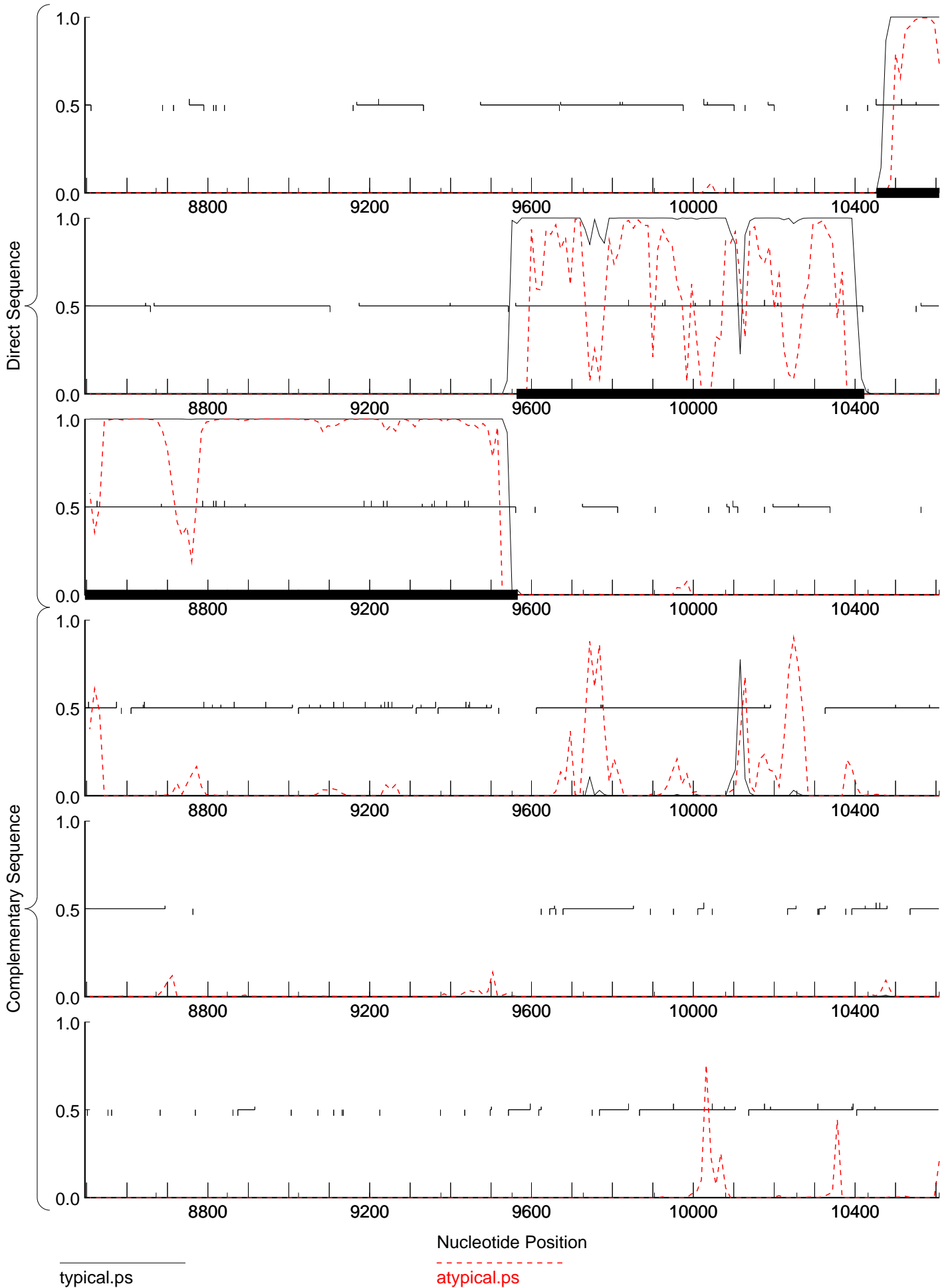
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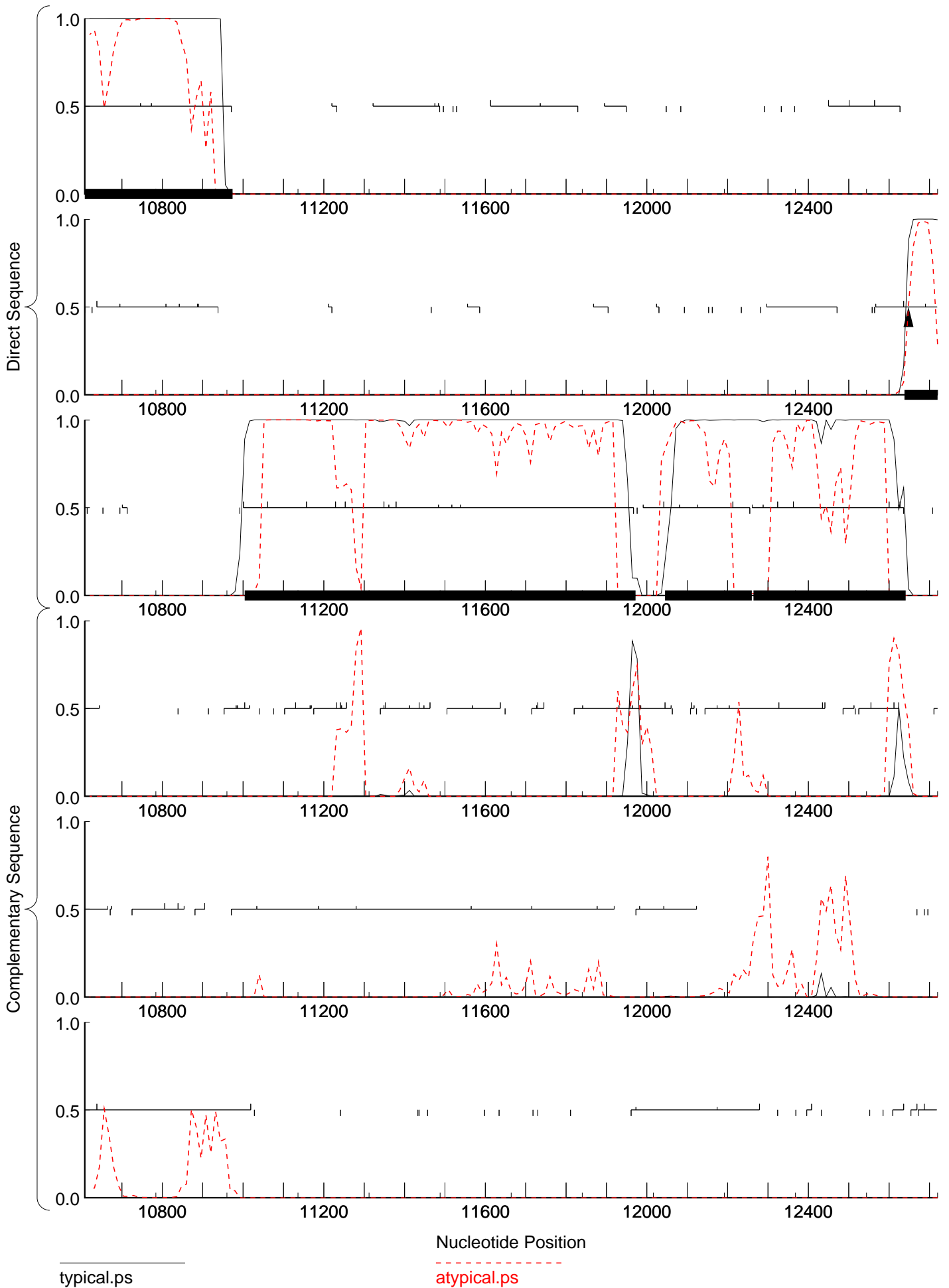




3 complete sequence, 53277 bp including 10-base 3' overhang (CGGACGGTTA), Cluster A9, Order 4, Window 96, Step 12, 5/27
3 complete sequence, 53277 bp including 10-base 3' overhang (CGGACGGTTA), Cluster A9, Order 2, Window 96, Step 12, 5/27

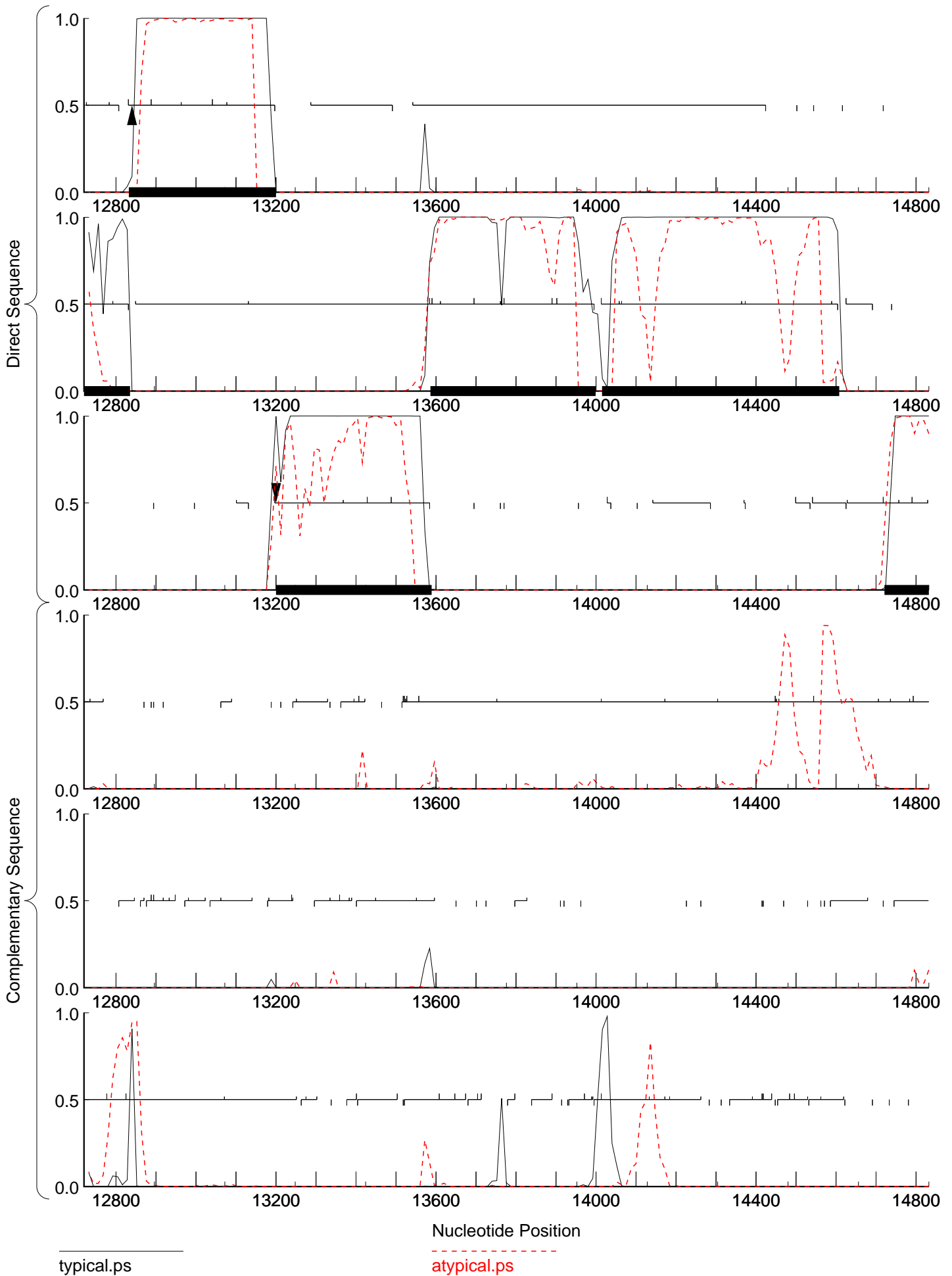




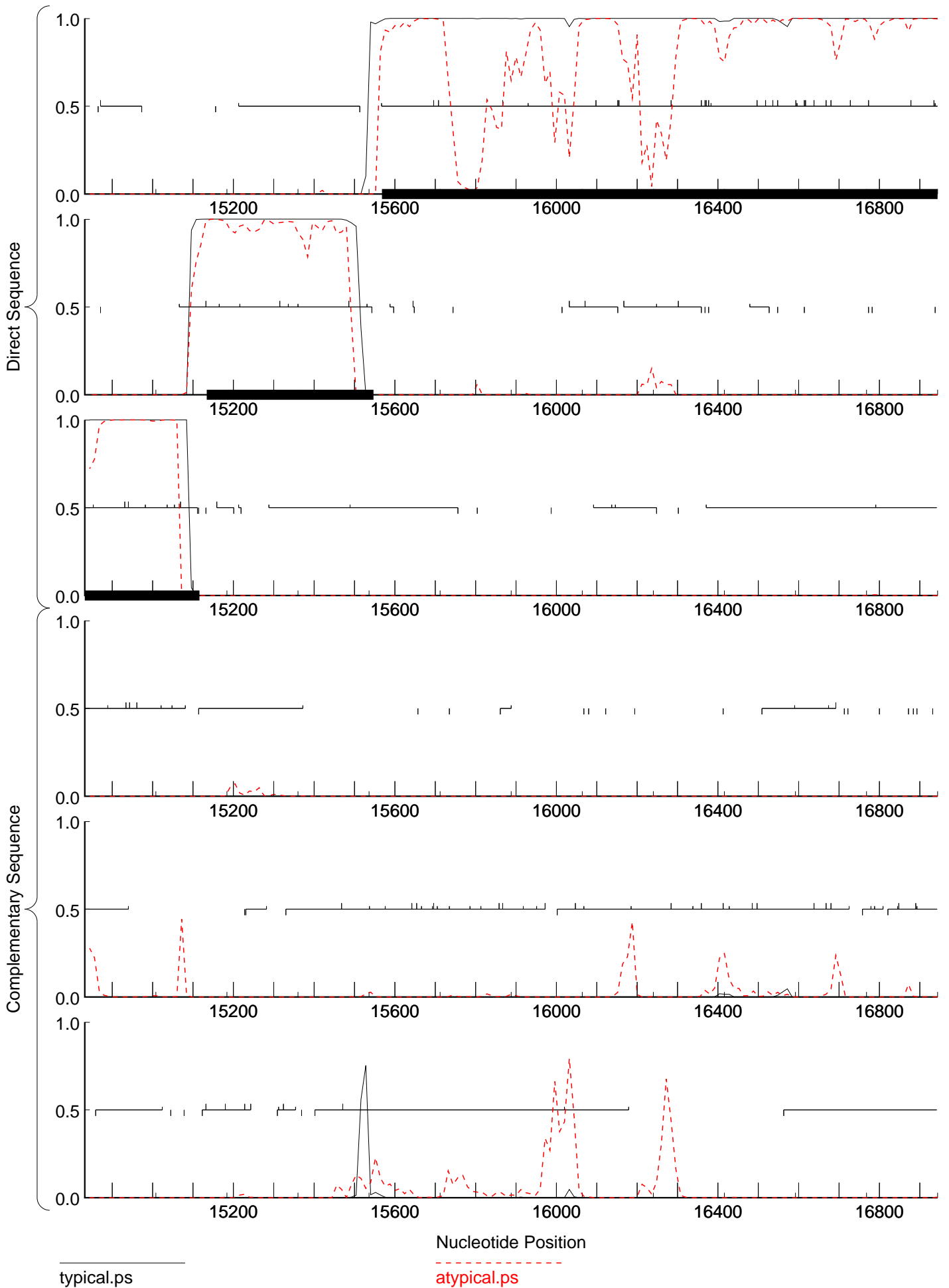


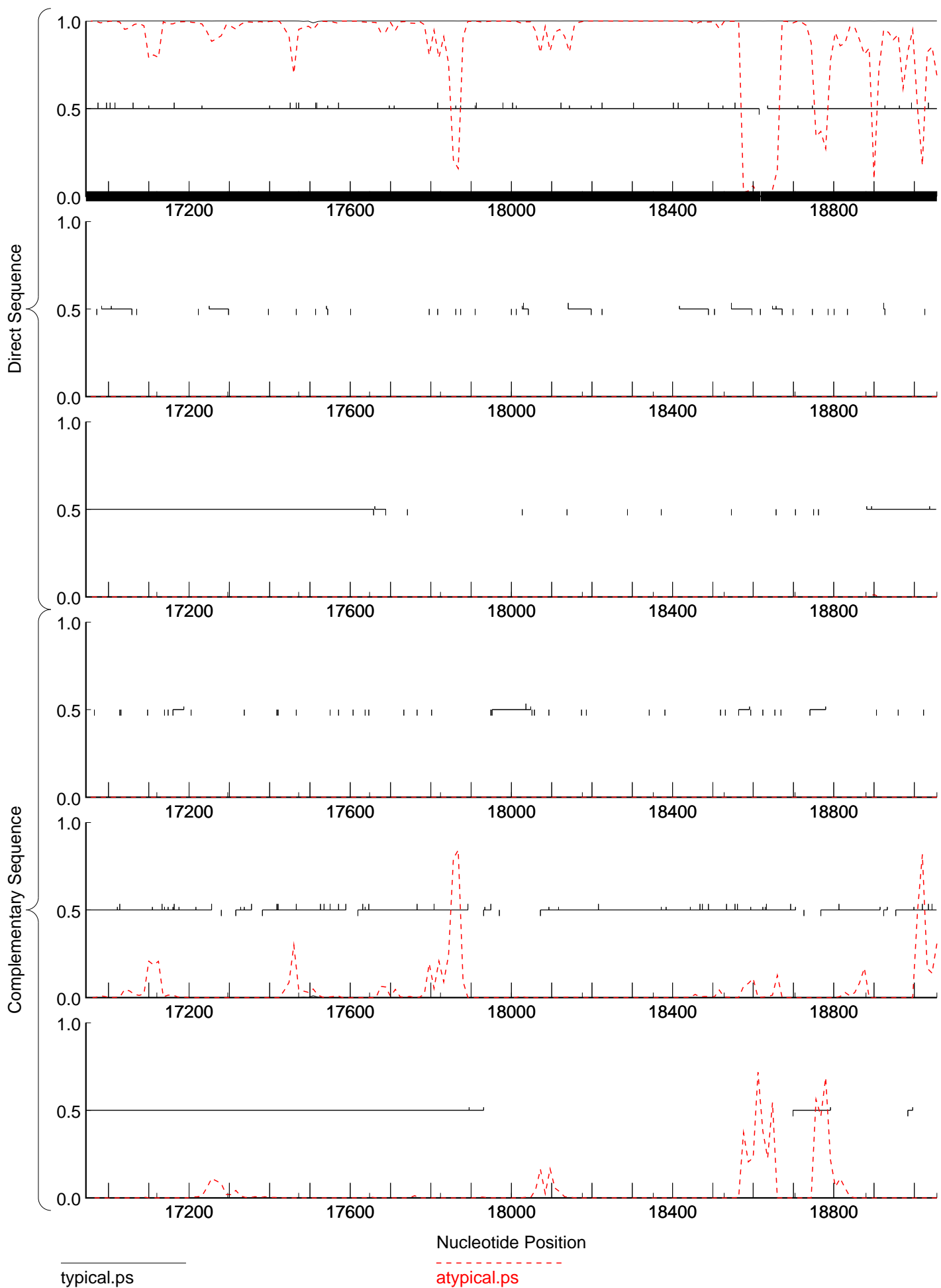
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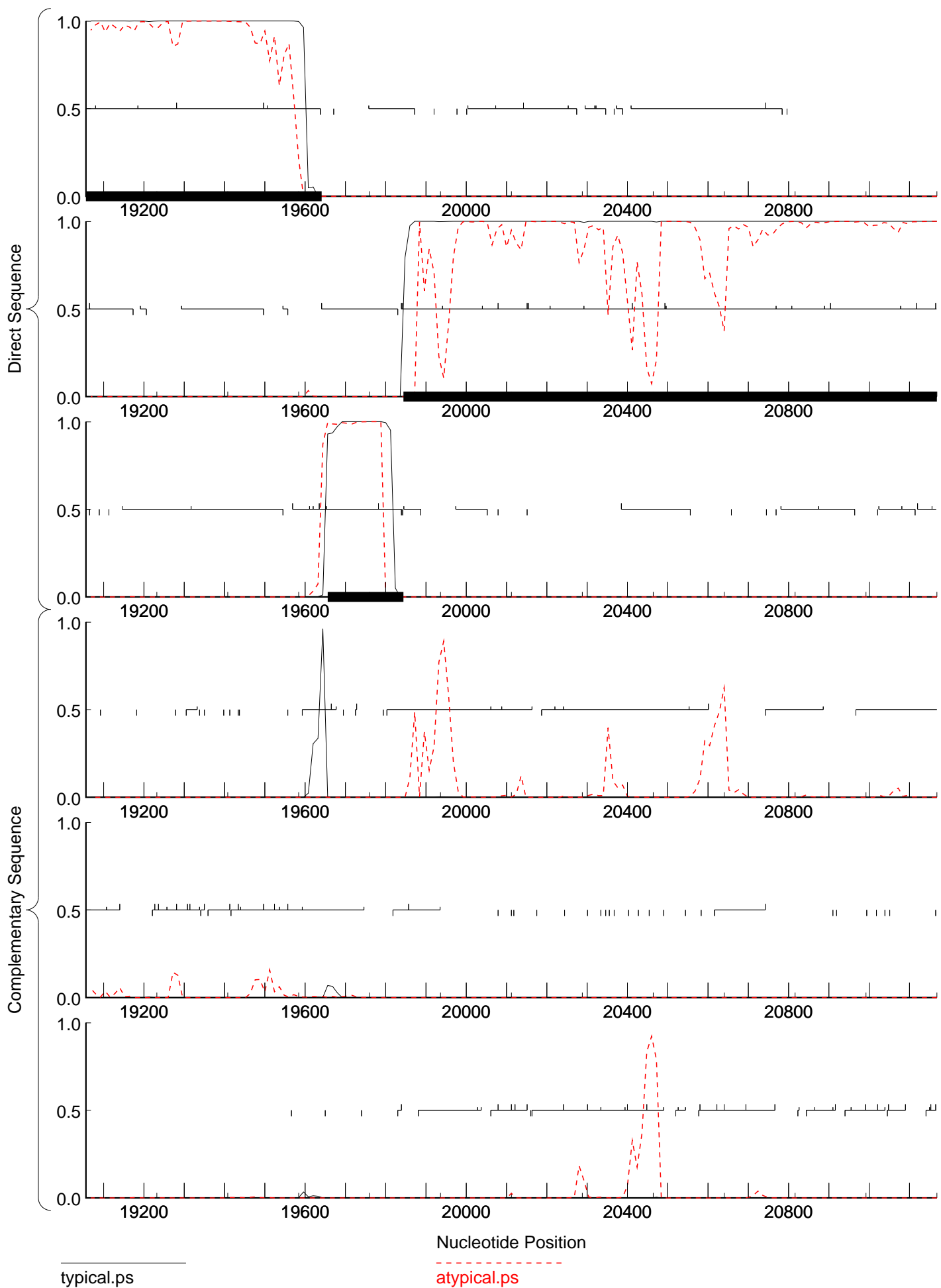
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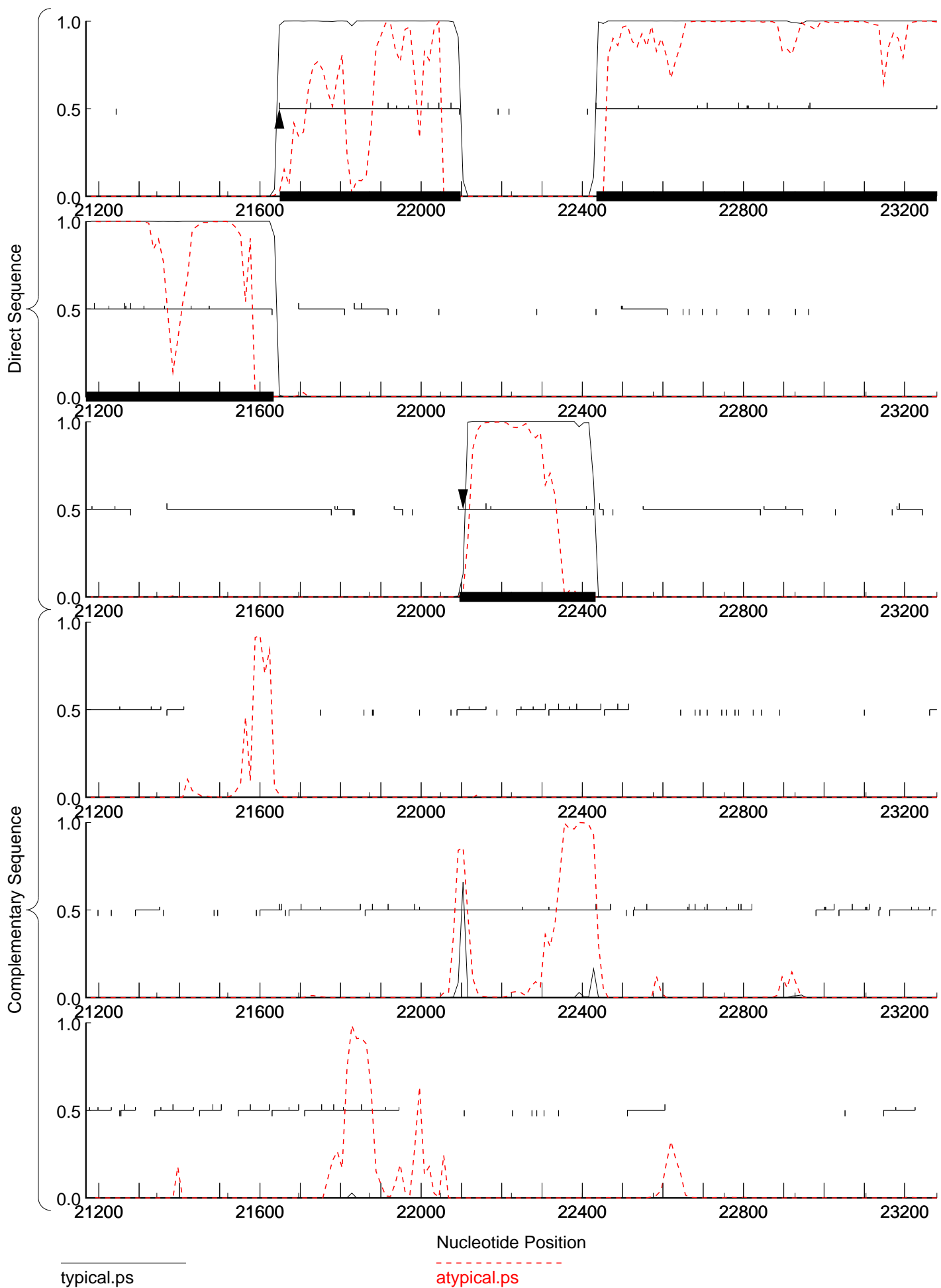


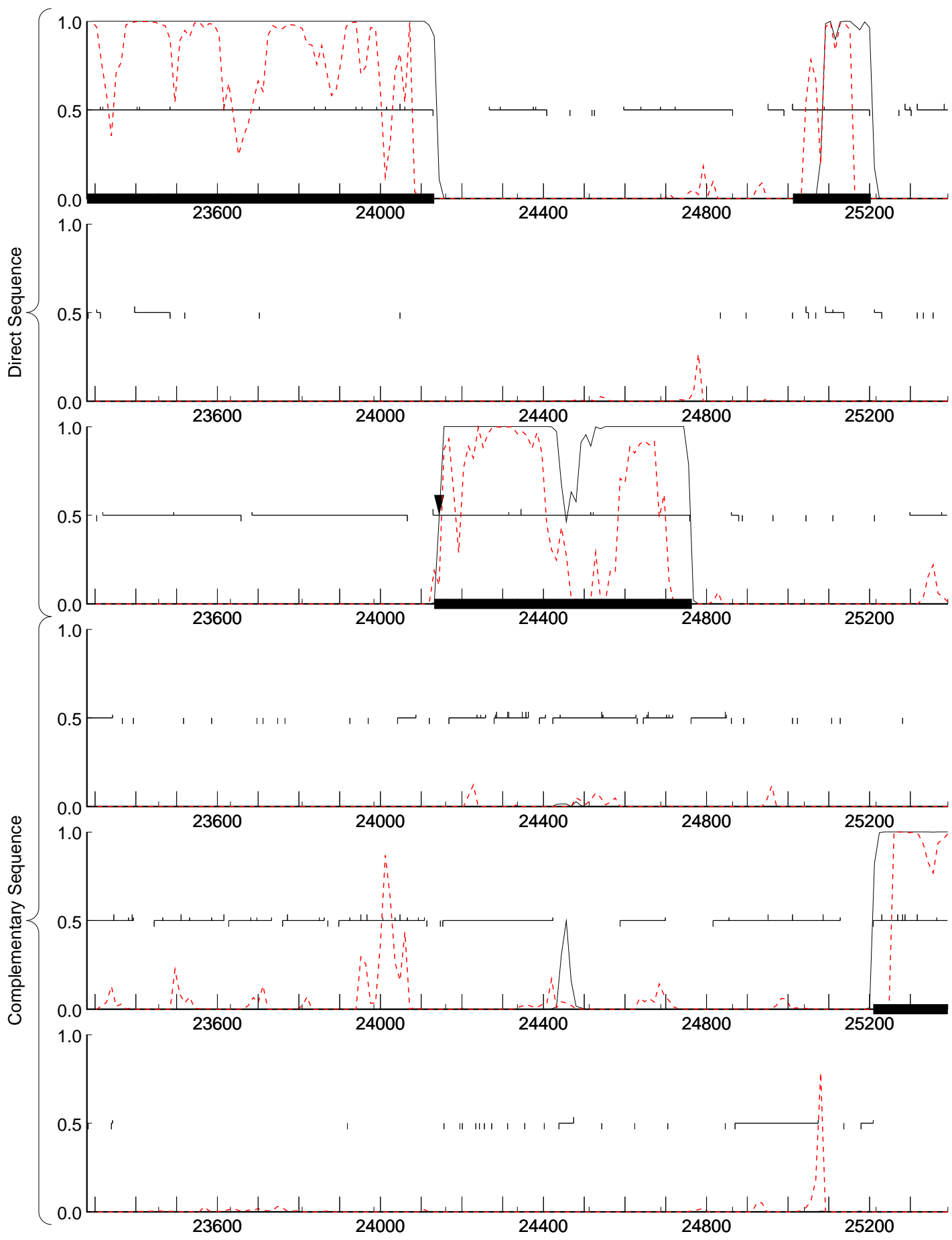
3 complete sequence, 53277 bp including 10-base 3' overhang (CGGACGGTTA), Cluster A9, Order 4, Window 96, Step 12, 9/27
3 complete sequence, 53277 bp including 10-base 3' overhang (CGGACGGTTA), Cluster A9, Order 2, Window 96, Step 12, 9/27





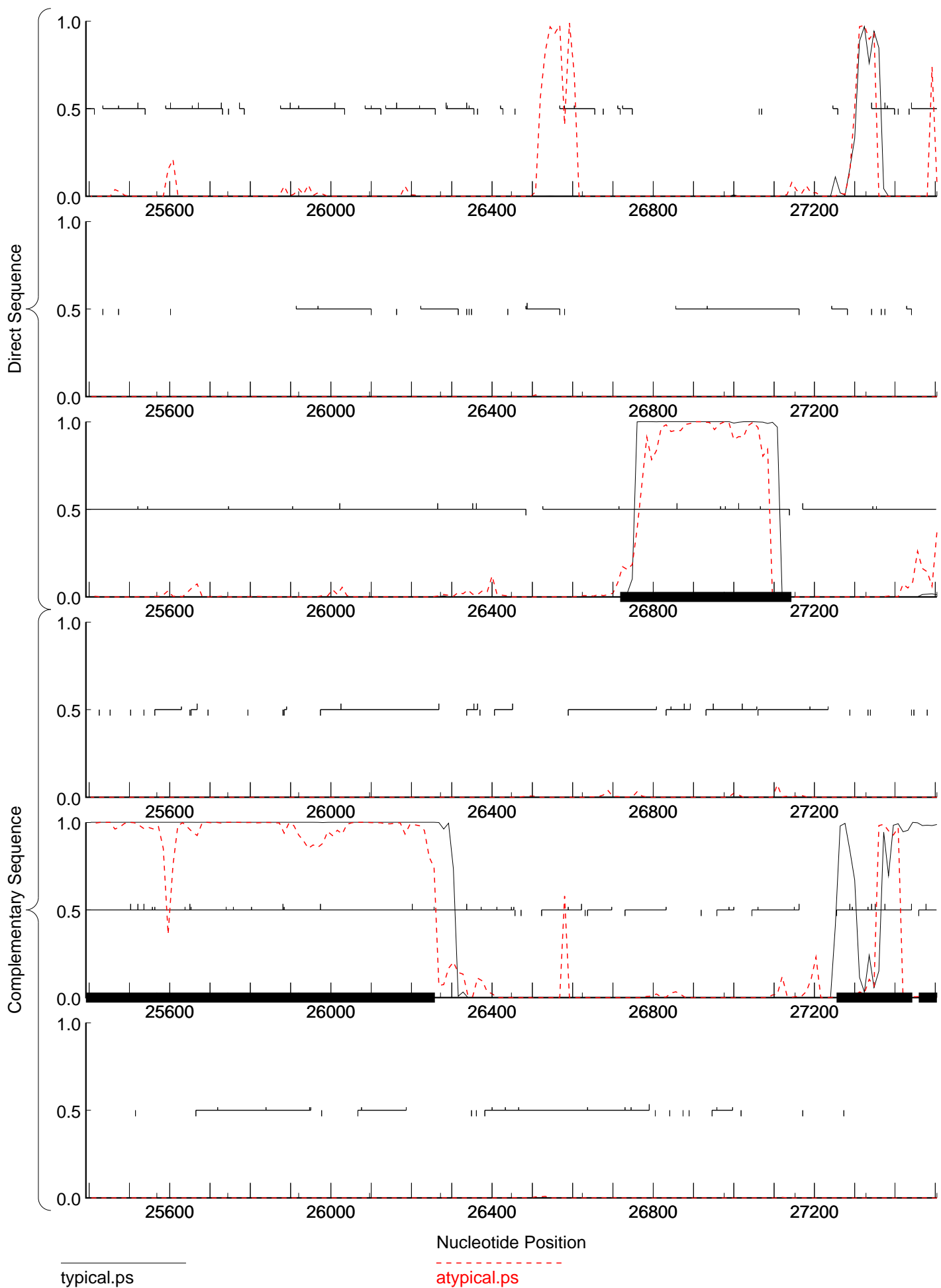






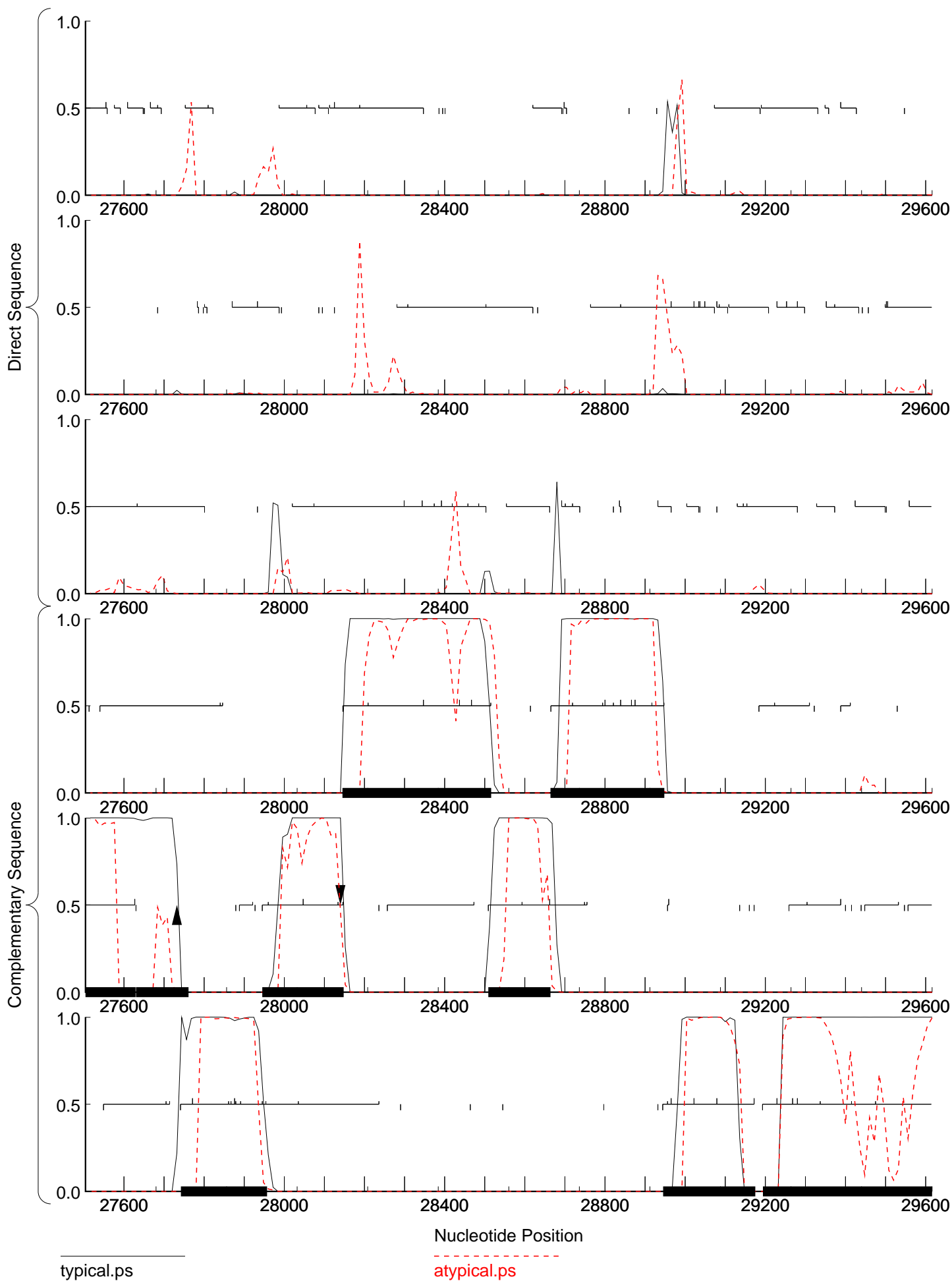
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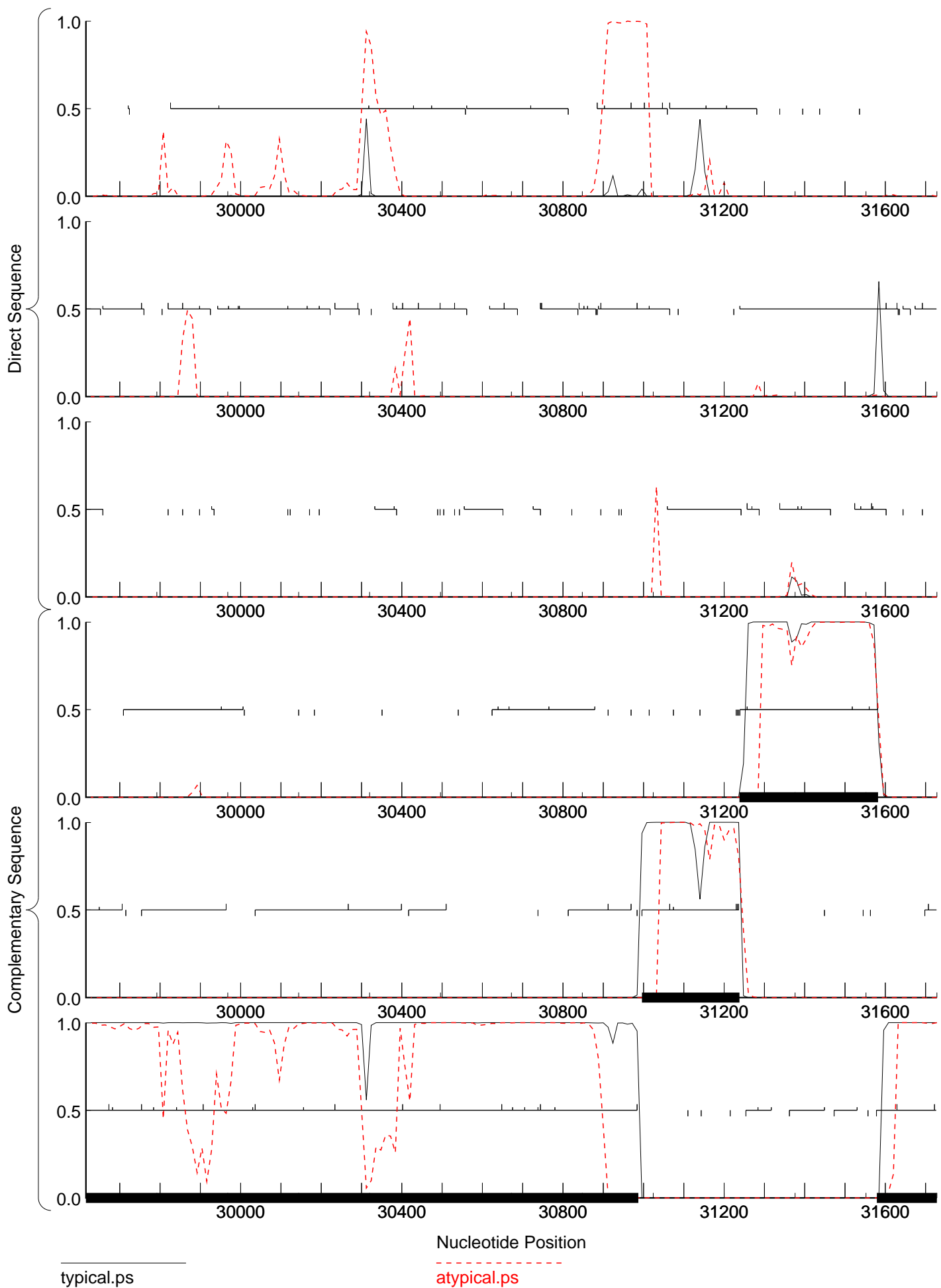
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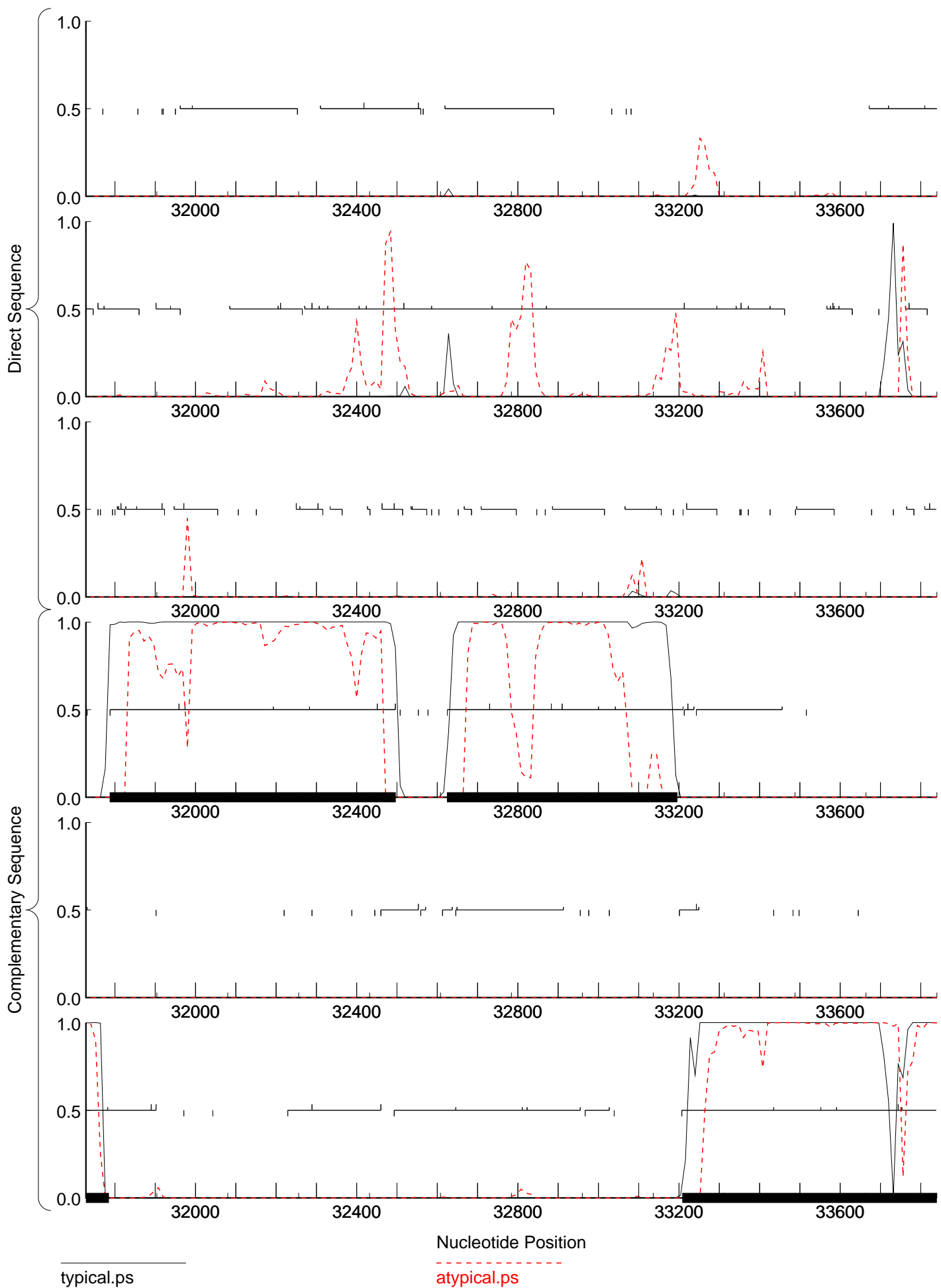
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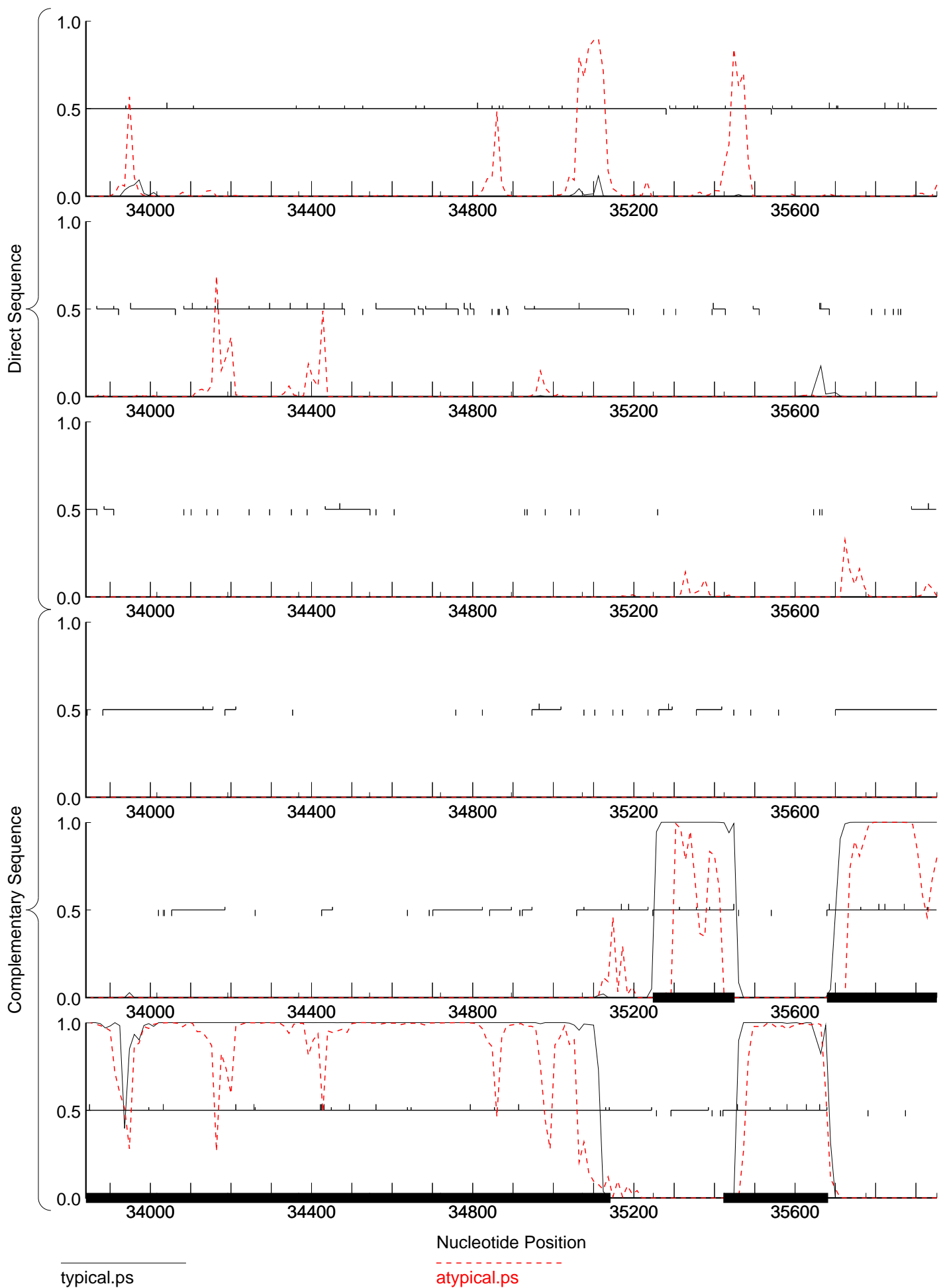


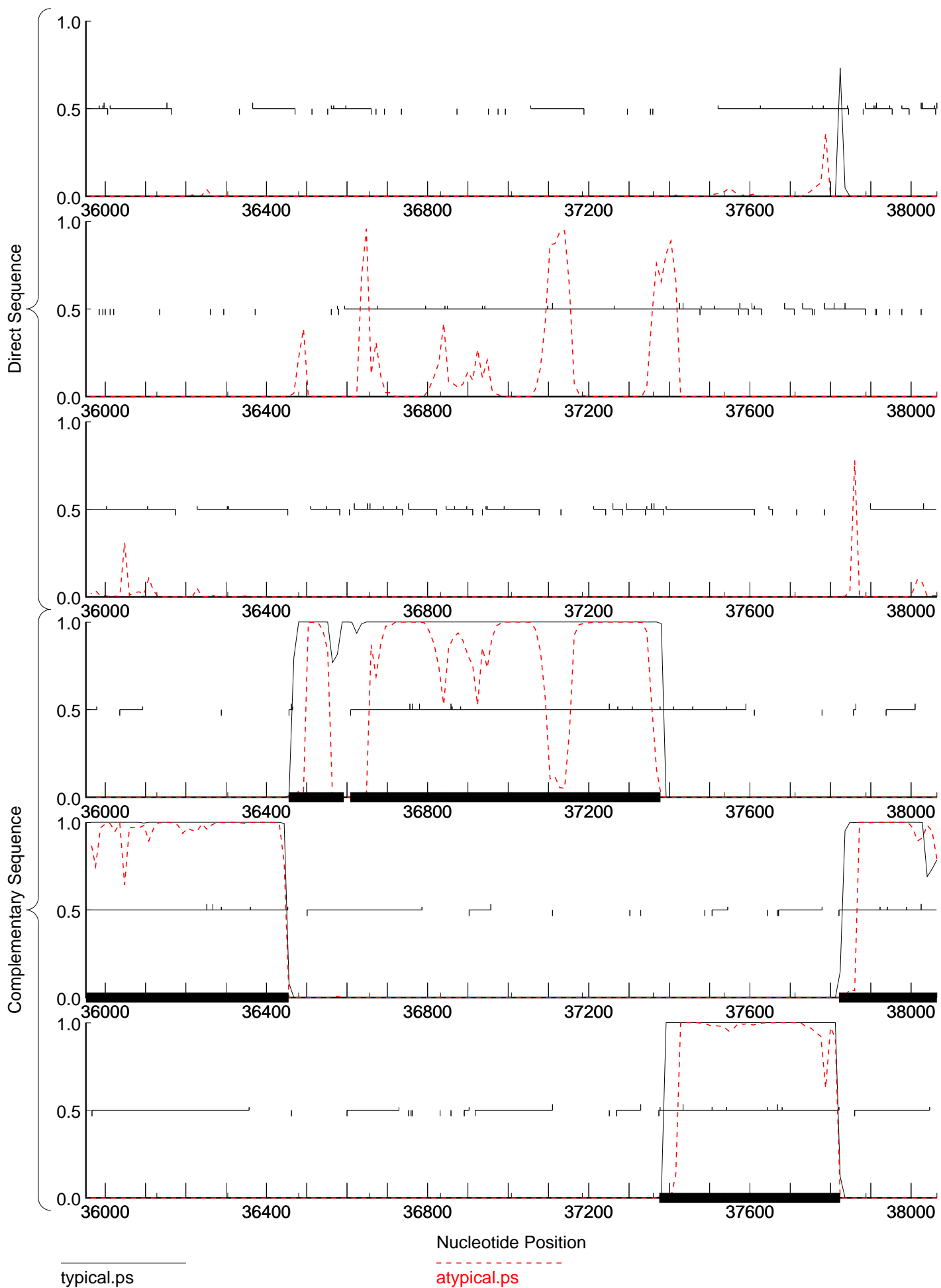
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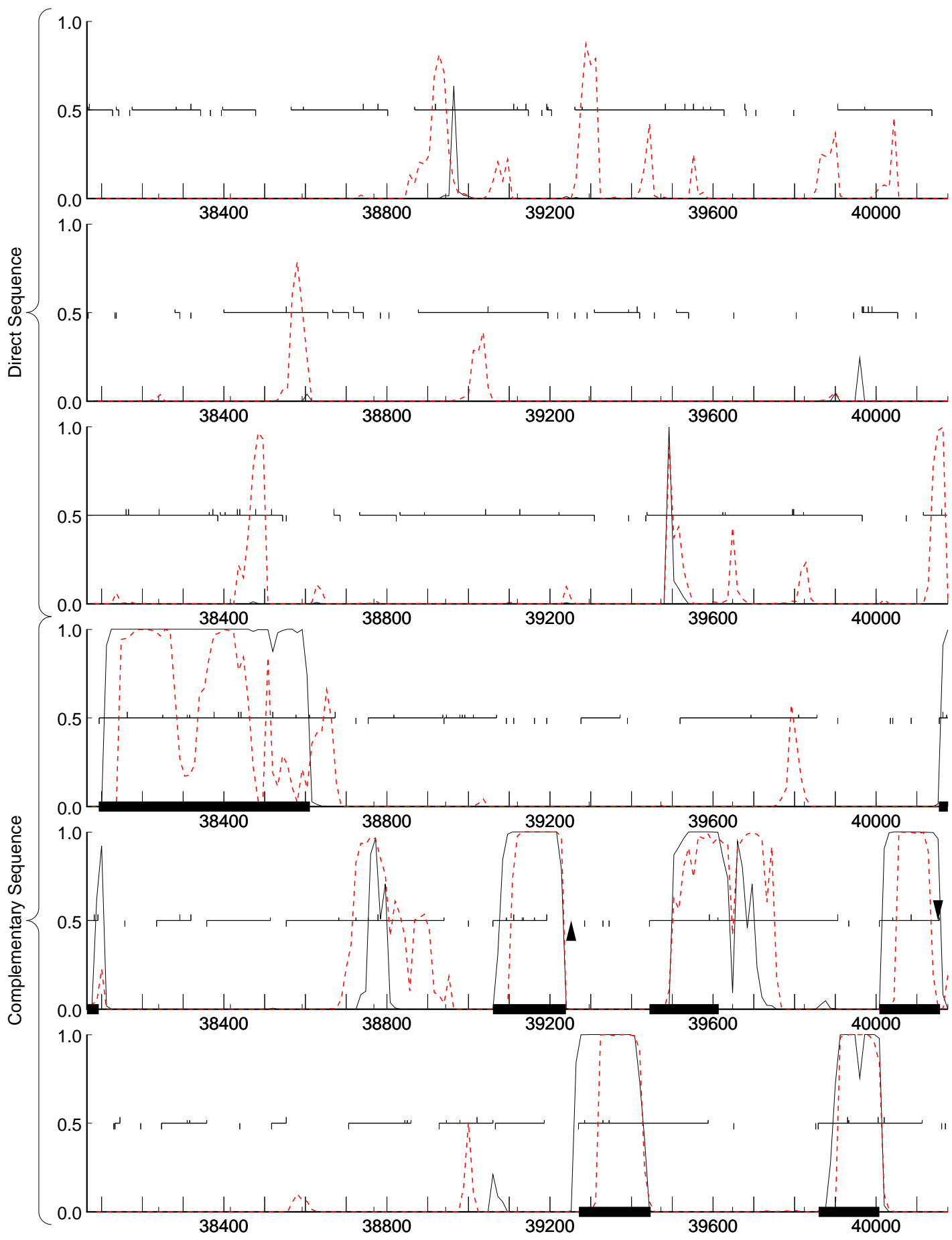
atypical.ps











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