

GeneMark

Version 2.5p (09.08.06)

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PROGRAM INFORMATION

Sequence : Microbacterium phage Hyperion complete squence, 61769 bp including 203 bp terminal repeat,
Analysis Date : 6/24/17 at 10:02:46
Pages : 31
Sequence Length : 61769 bp
GC Content : 67.01%

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,

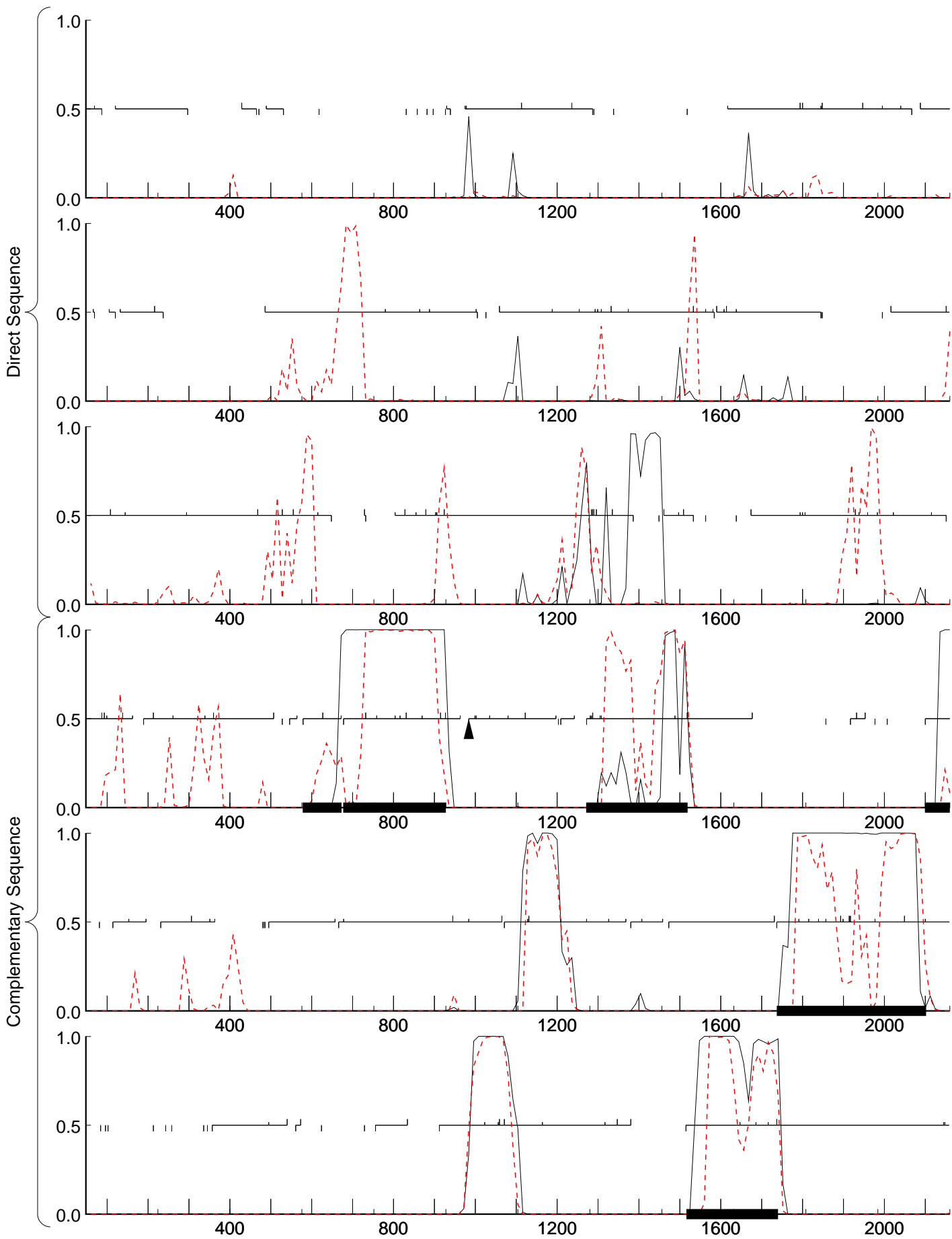
Matrix notes & comments

Training set derived by GeneMarkS, 4.30 October 2014
Sat Jun 24 10:02:46 2017

MATRIX INFORMATION

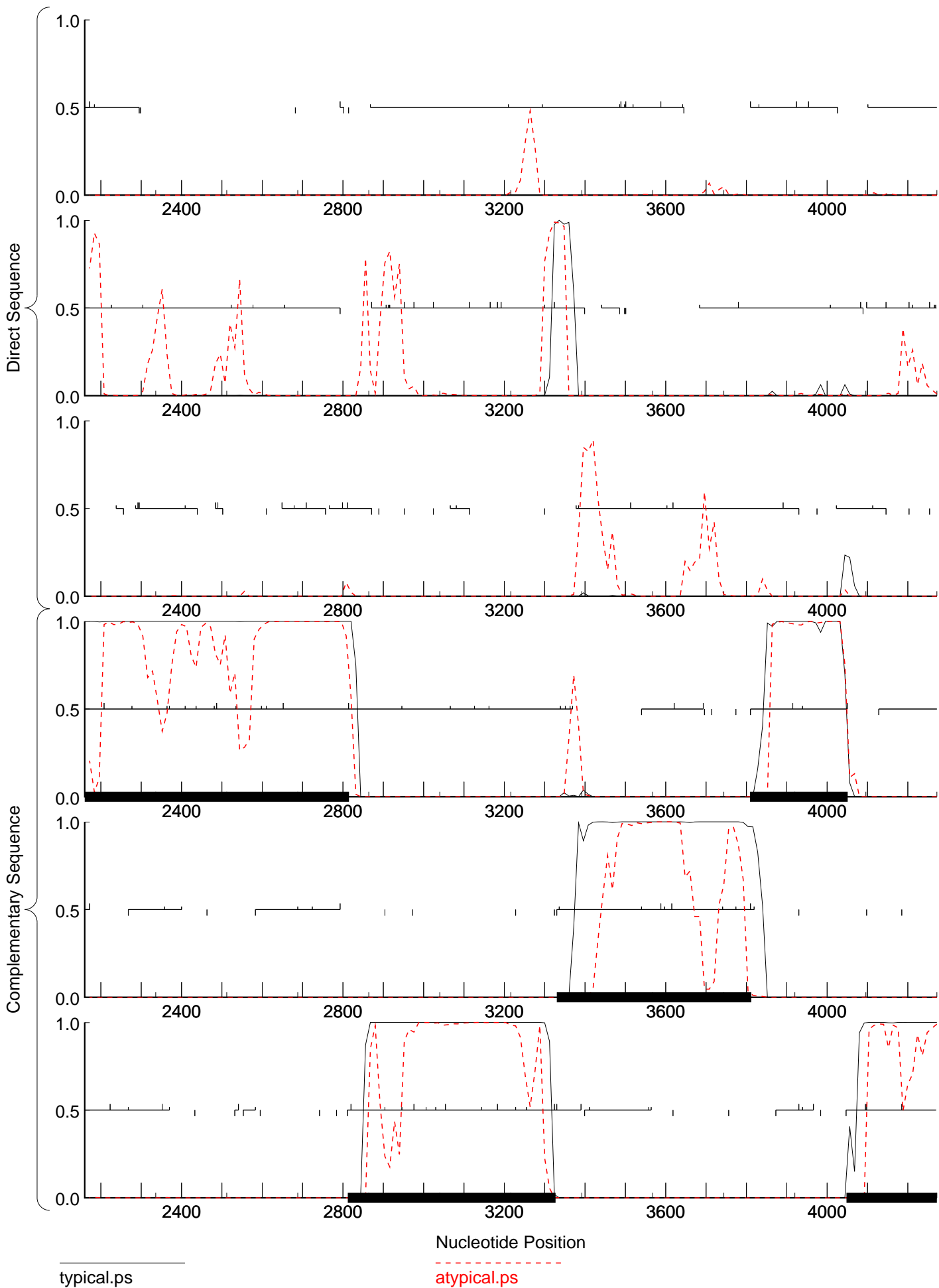
Matrix : -
Author : -
Order : 4

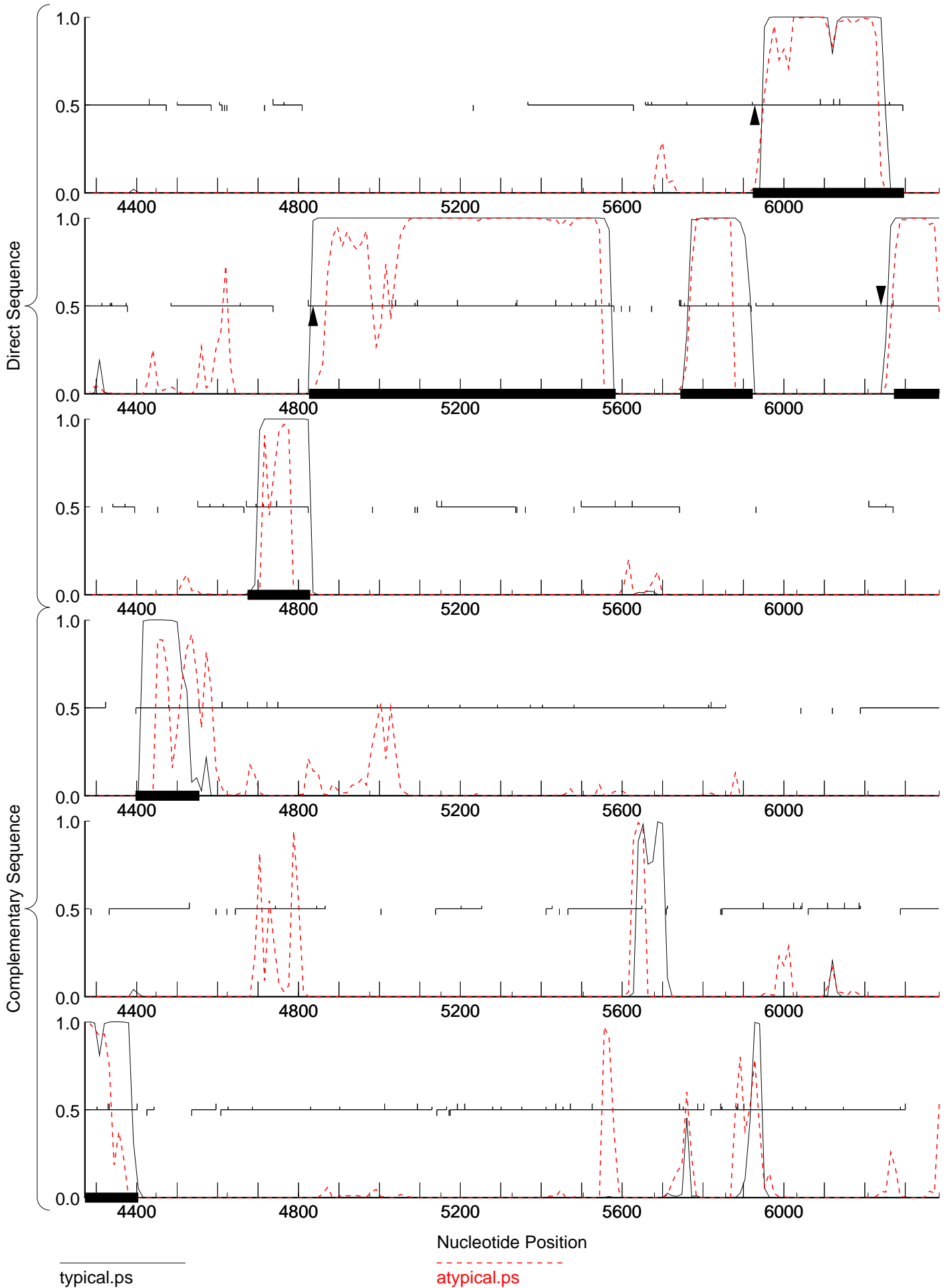
Send questions / comments to:
Dr. M. Borodovsky
Georgia Institute of Technology
School of Biology
Atlanta, GA 30332-0230



typical.ps

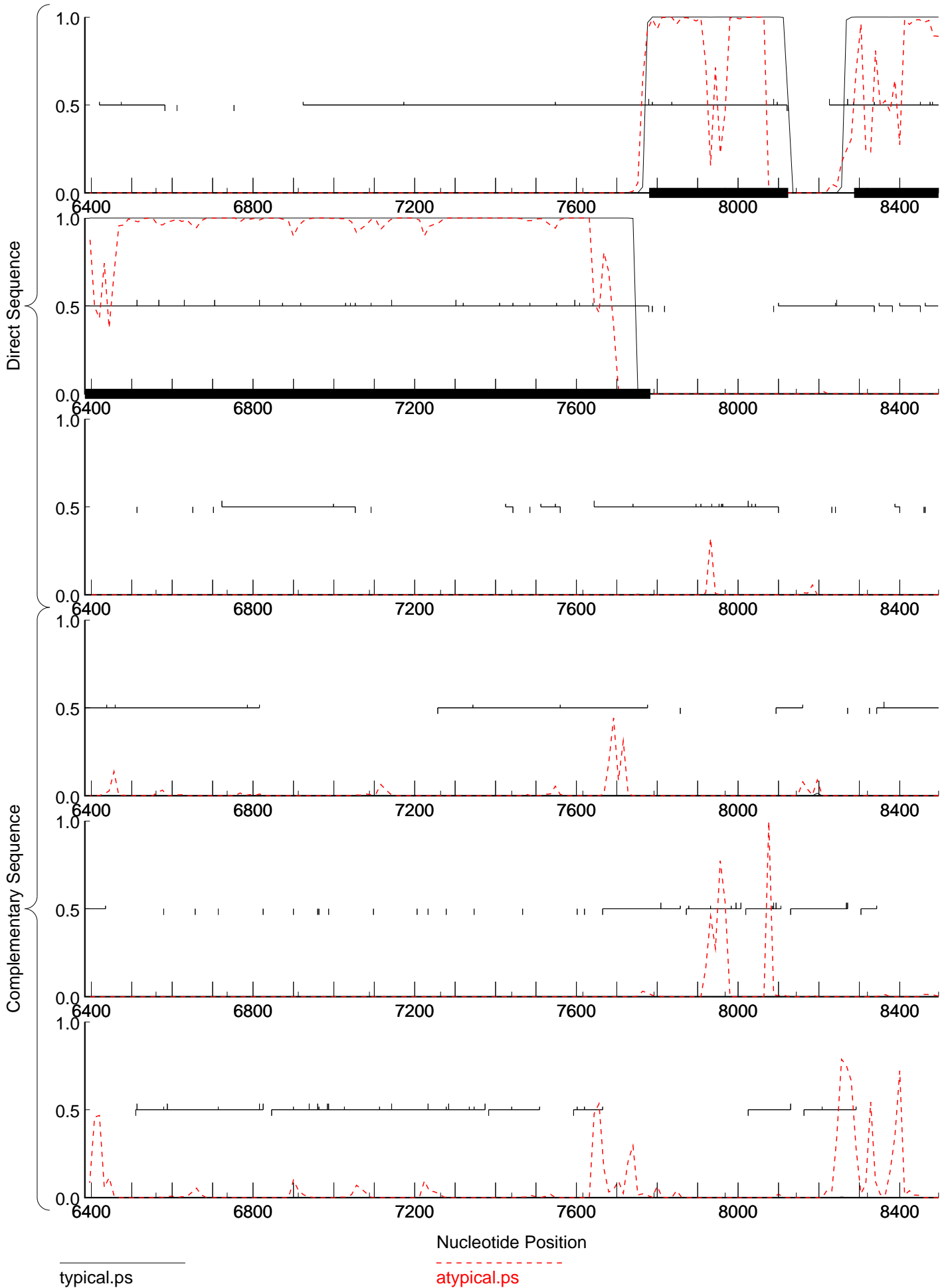
atypical.ps

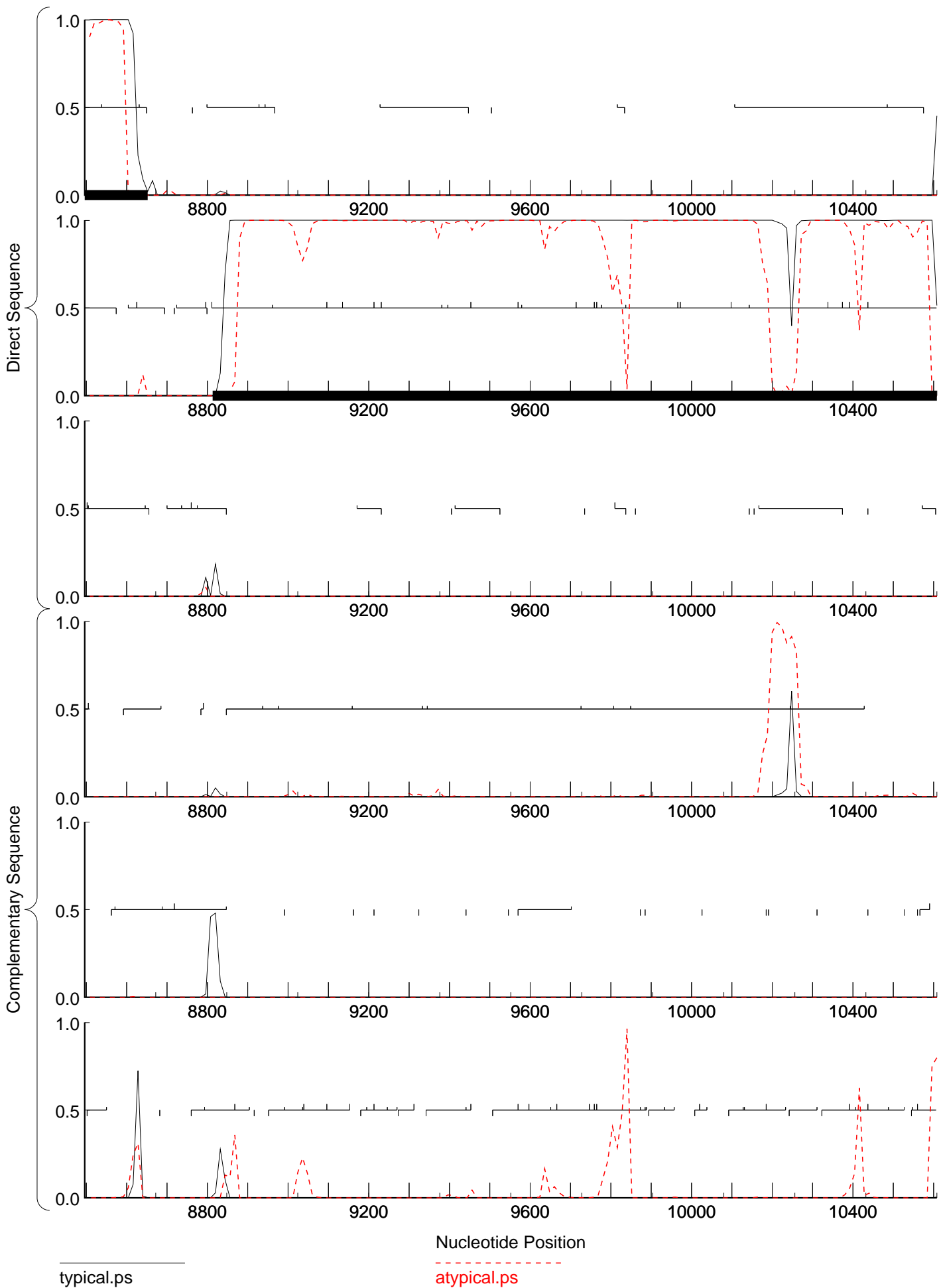


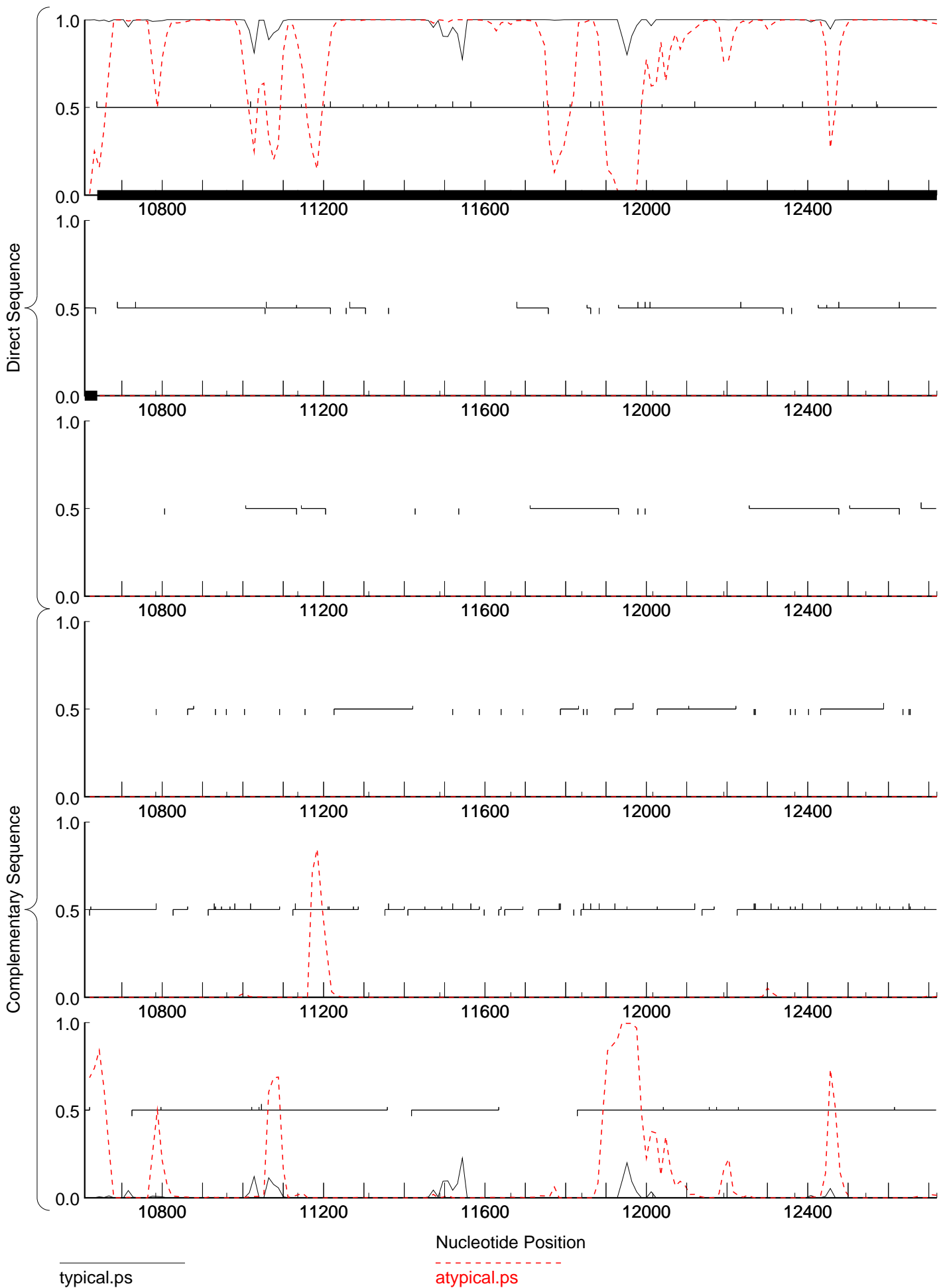


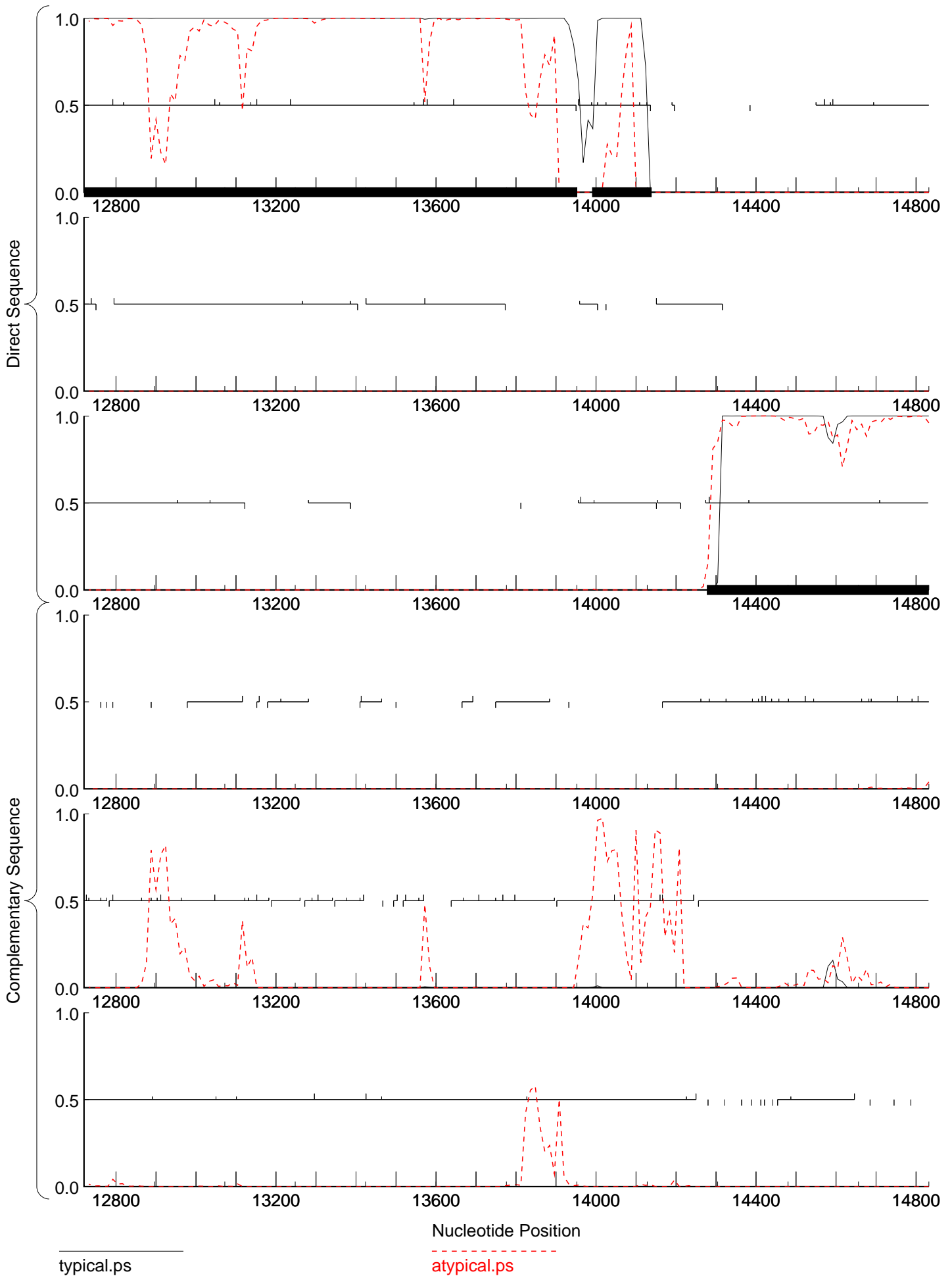
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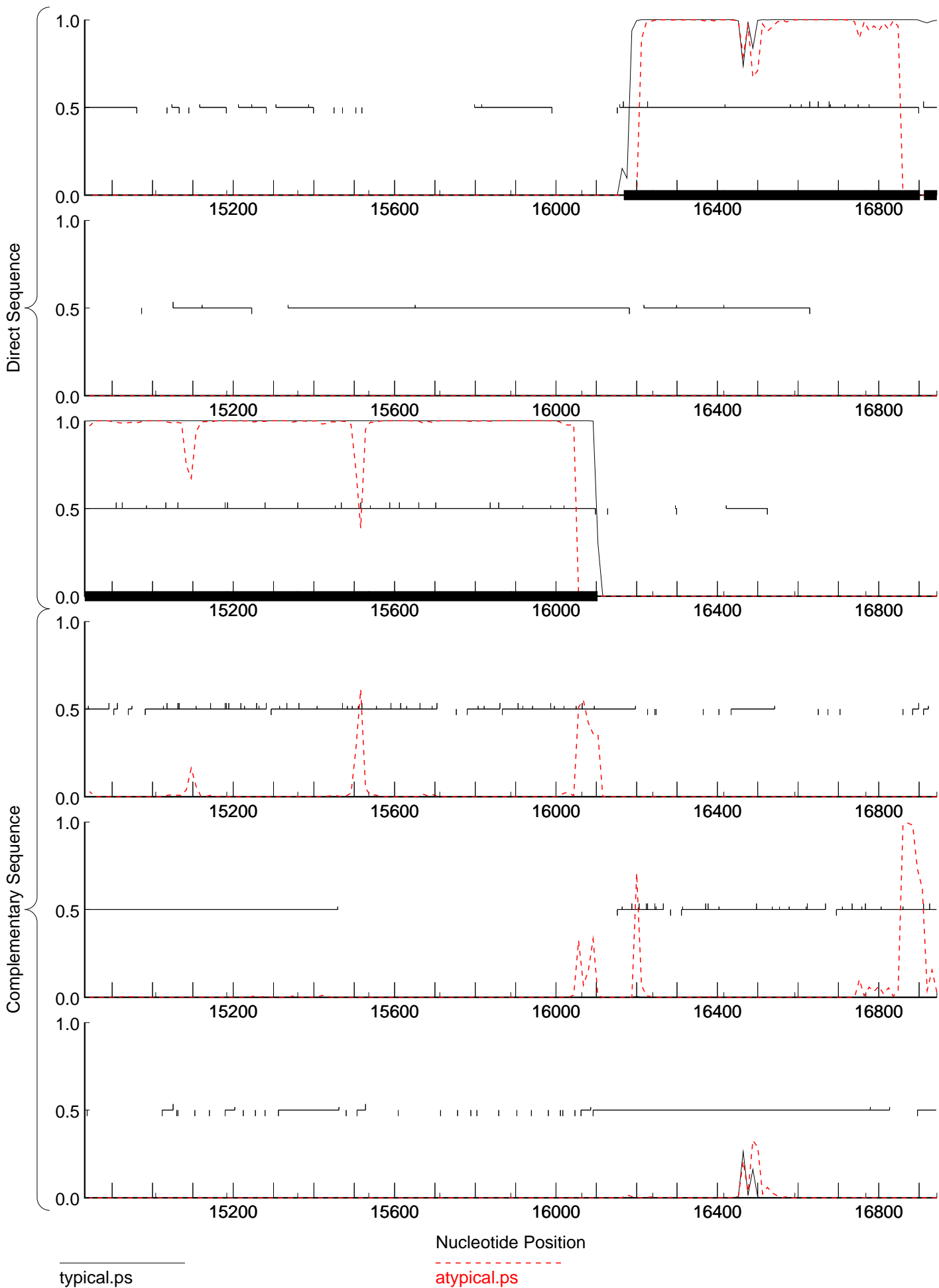


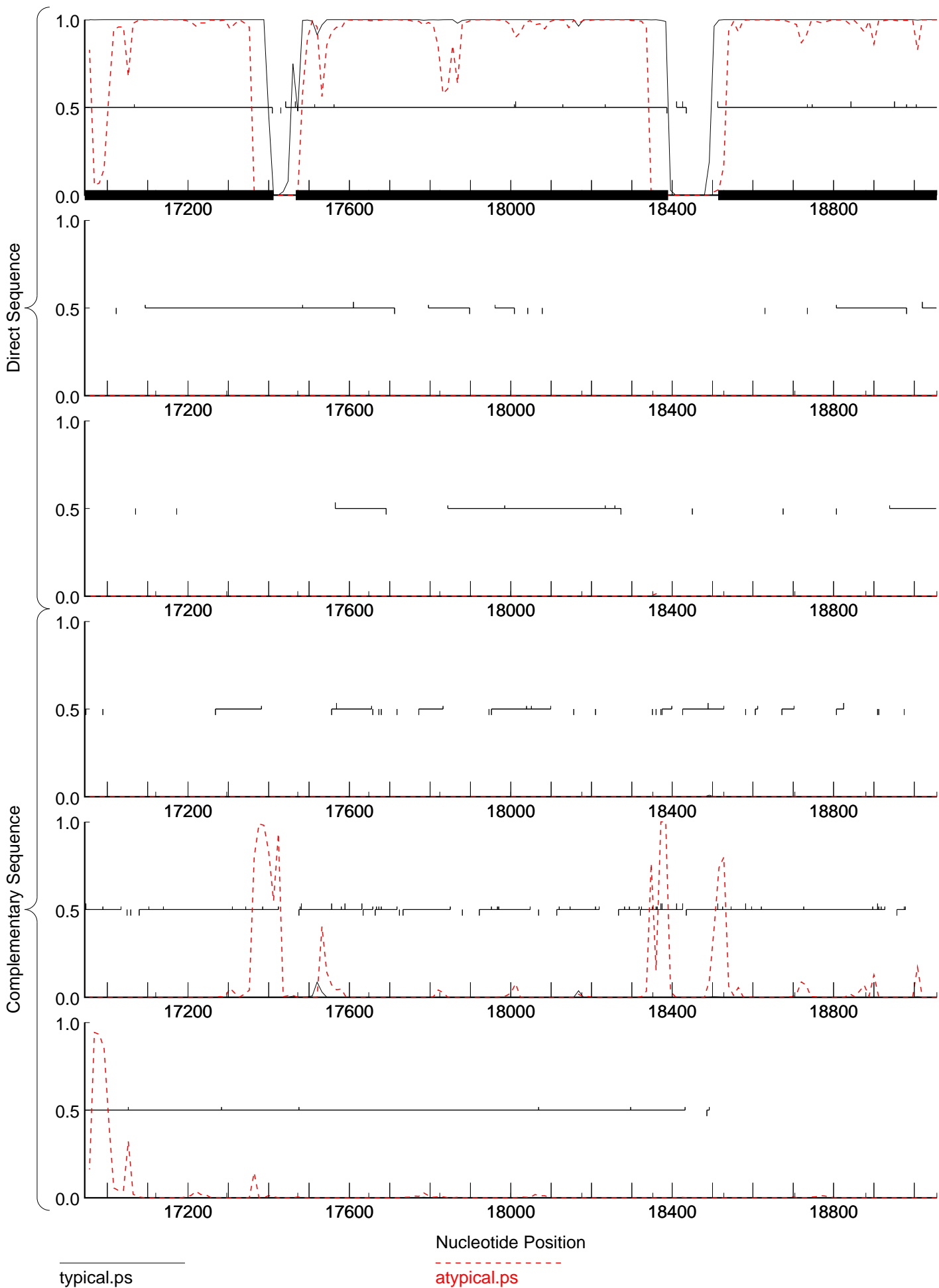


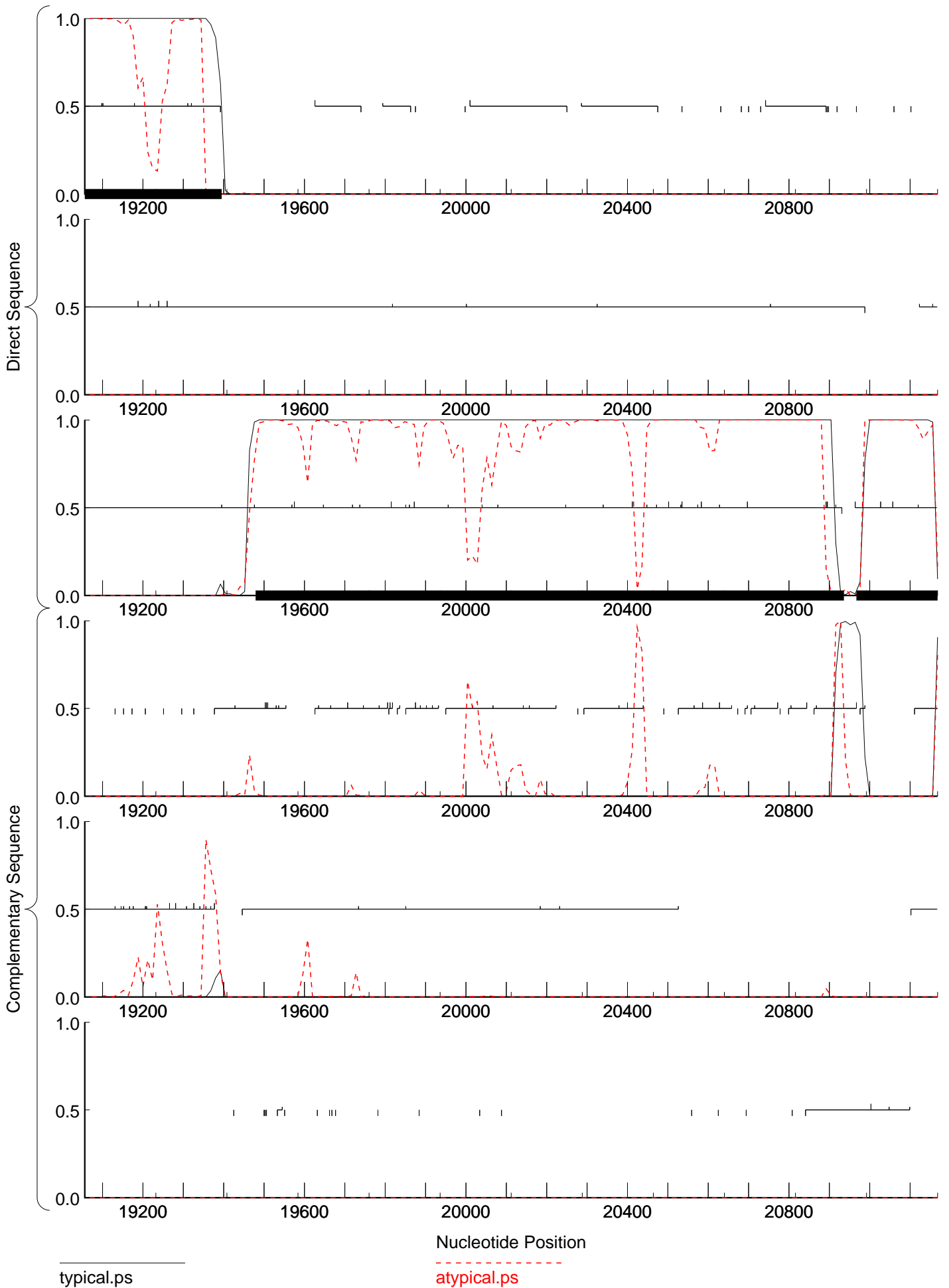


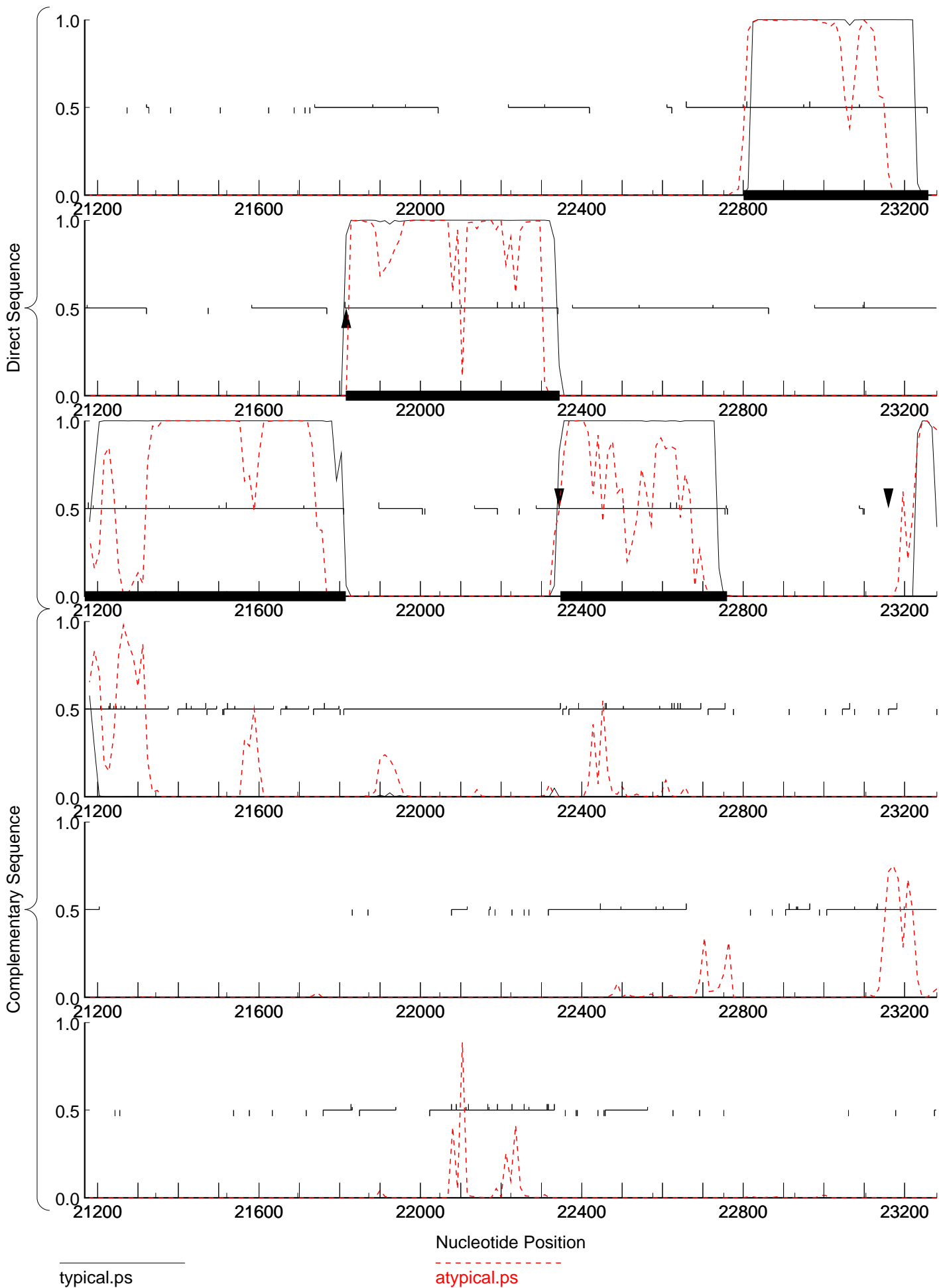
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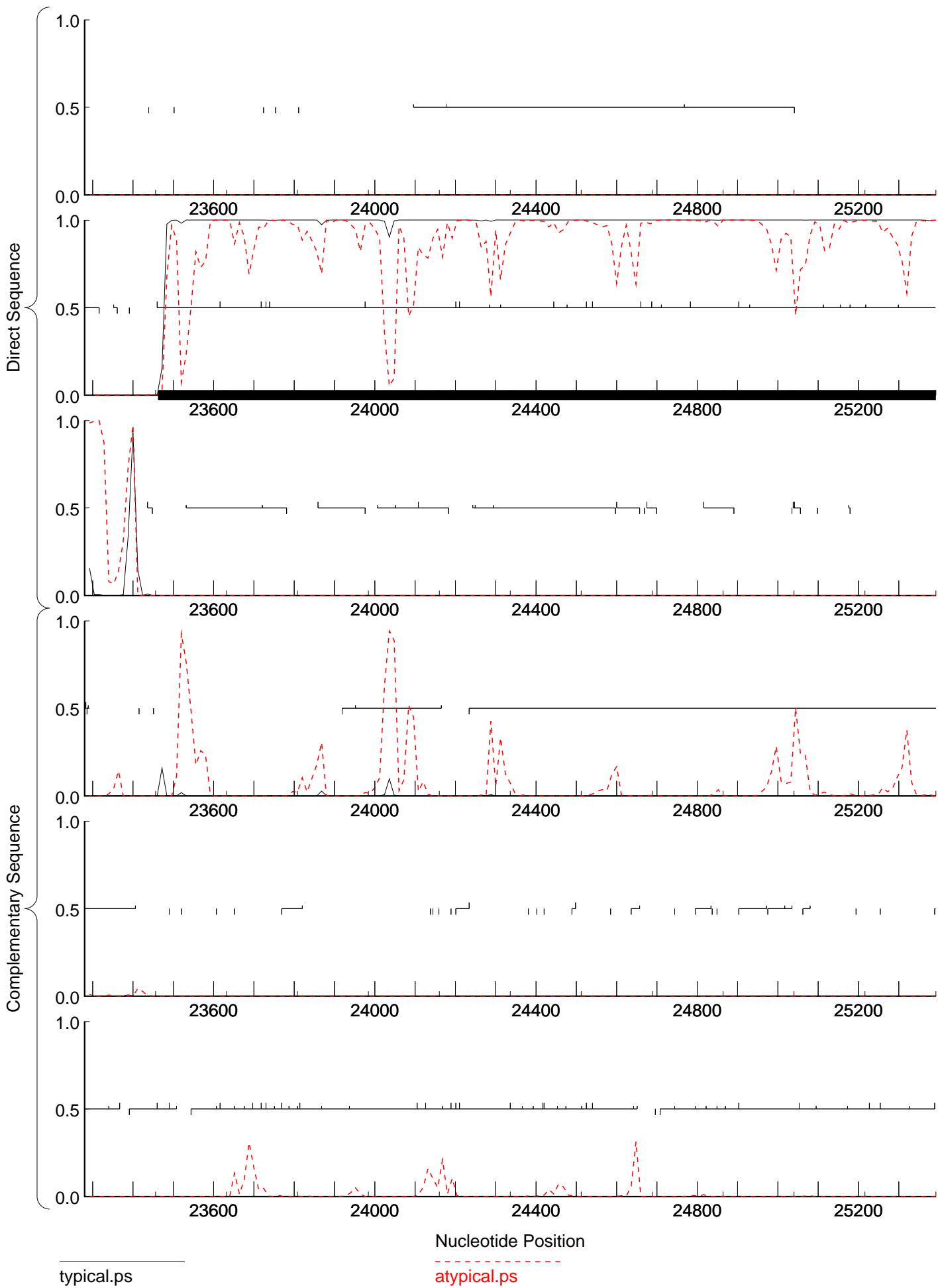
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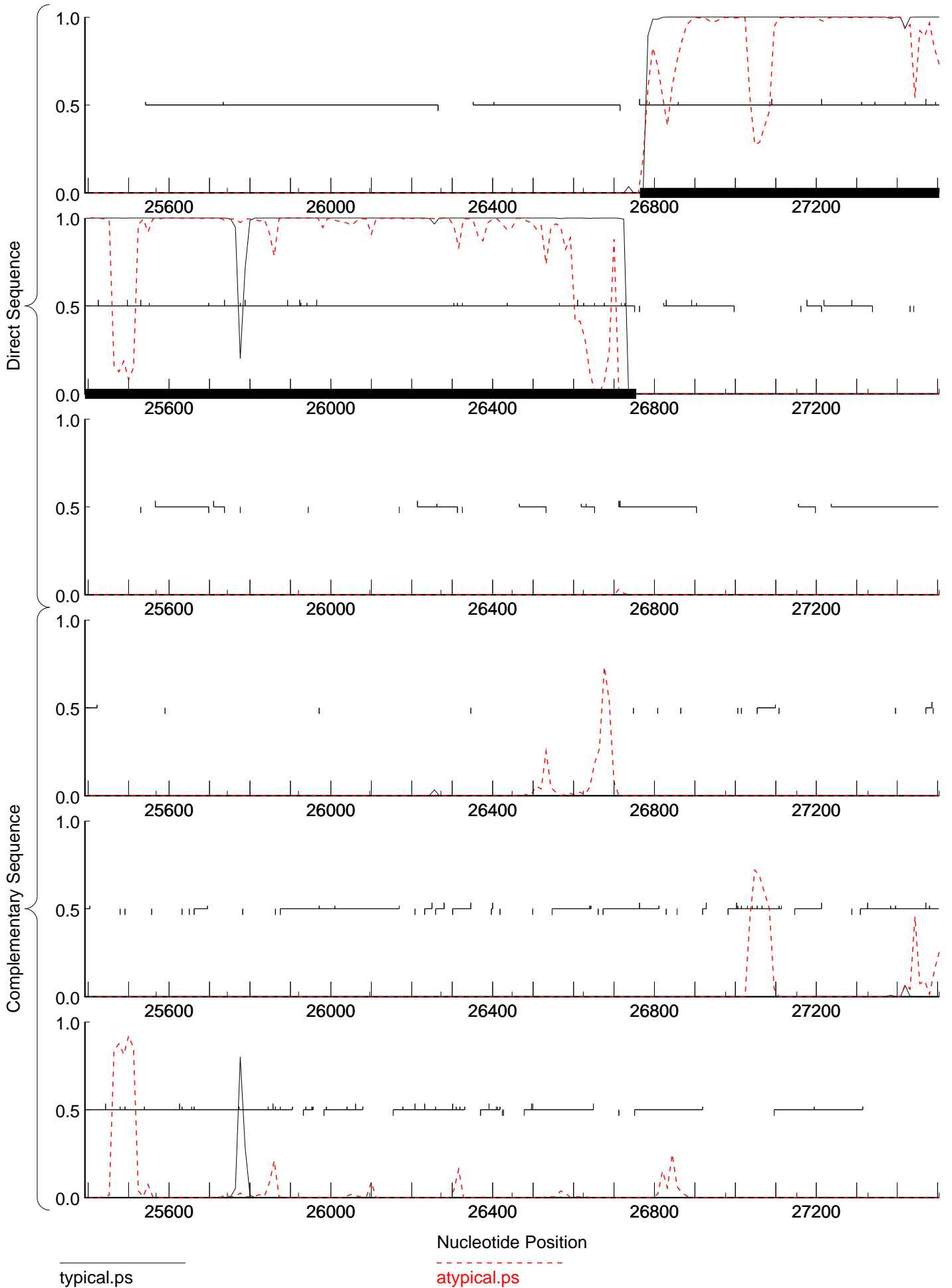


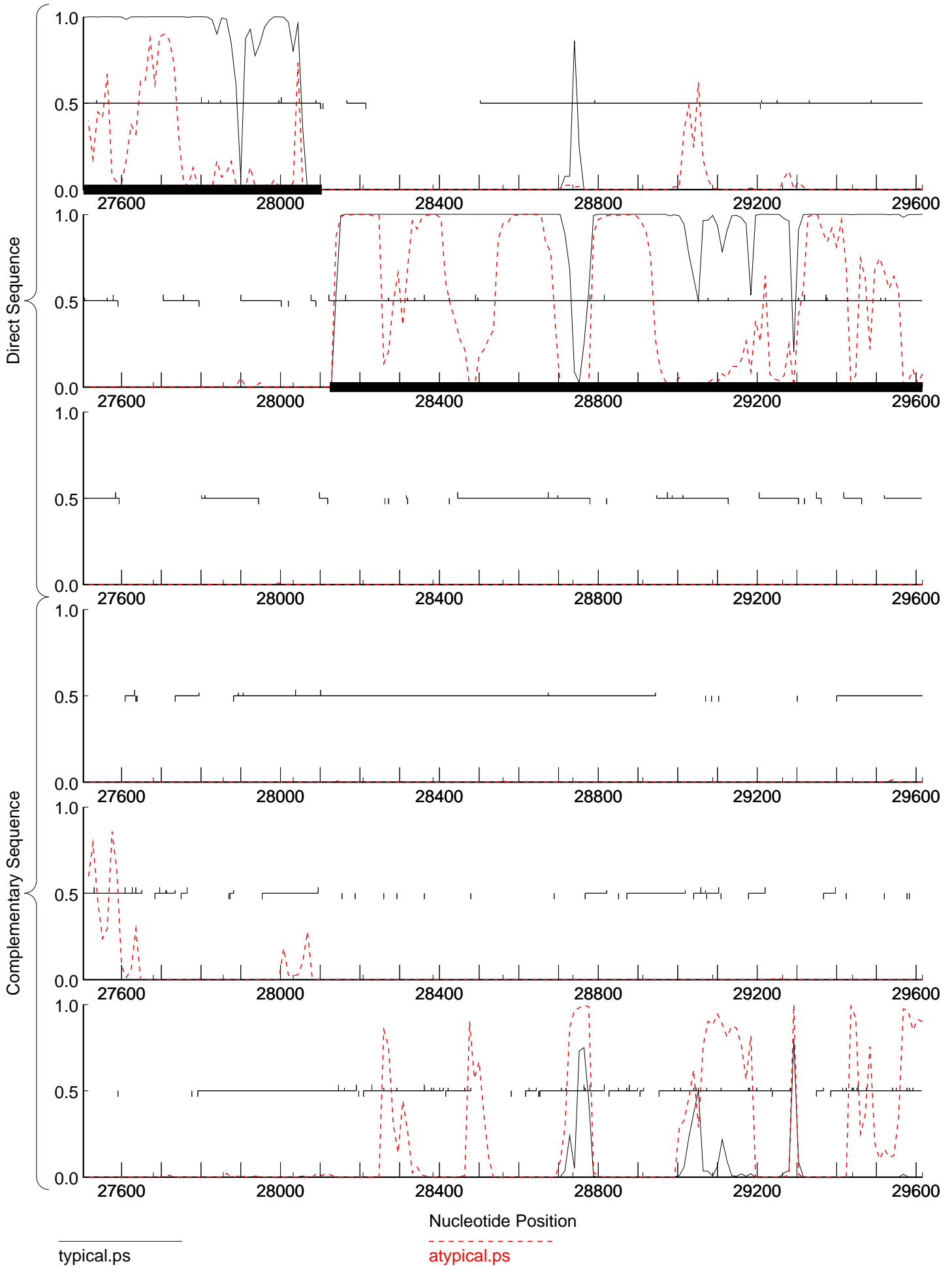


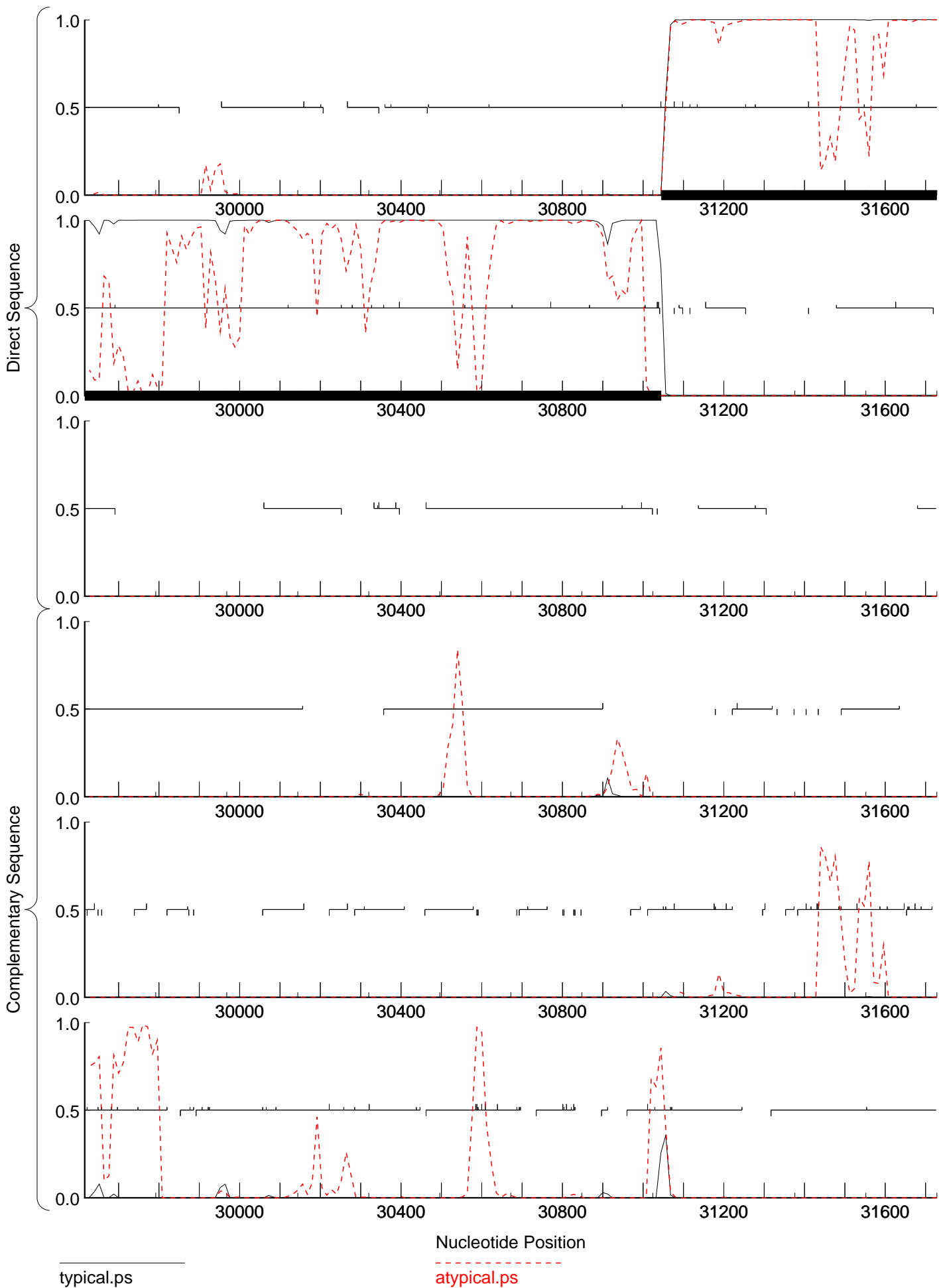


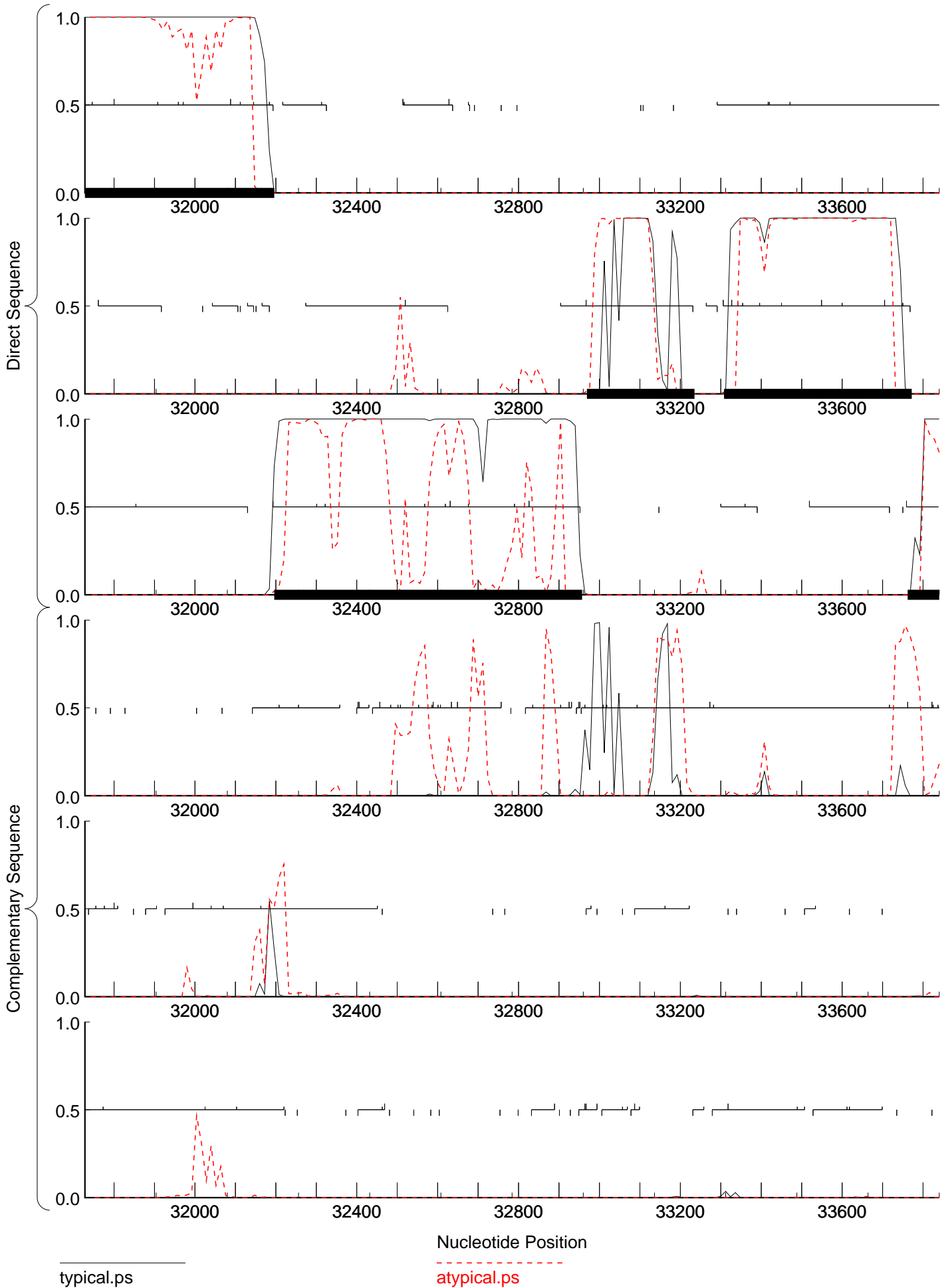








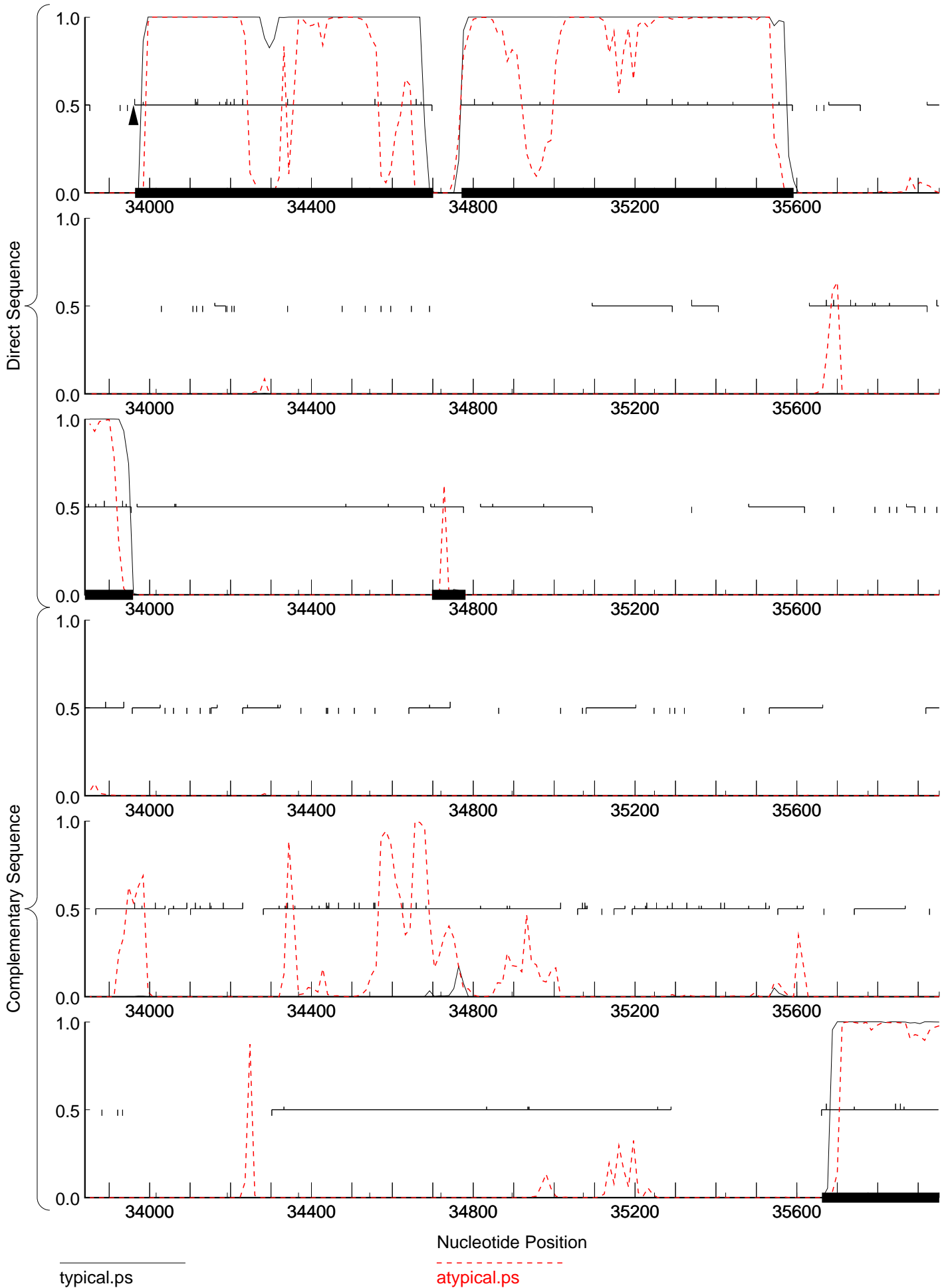


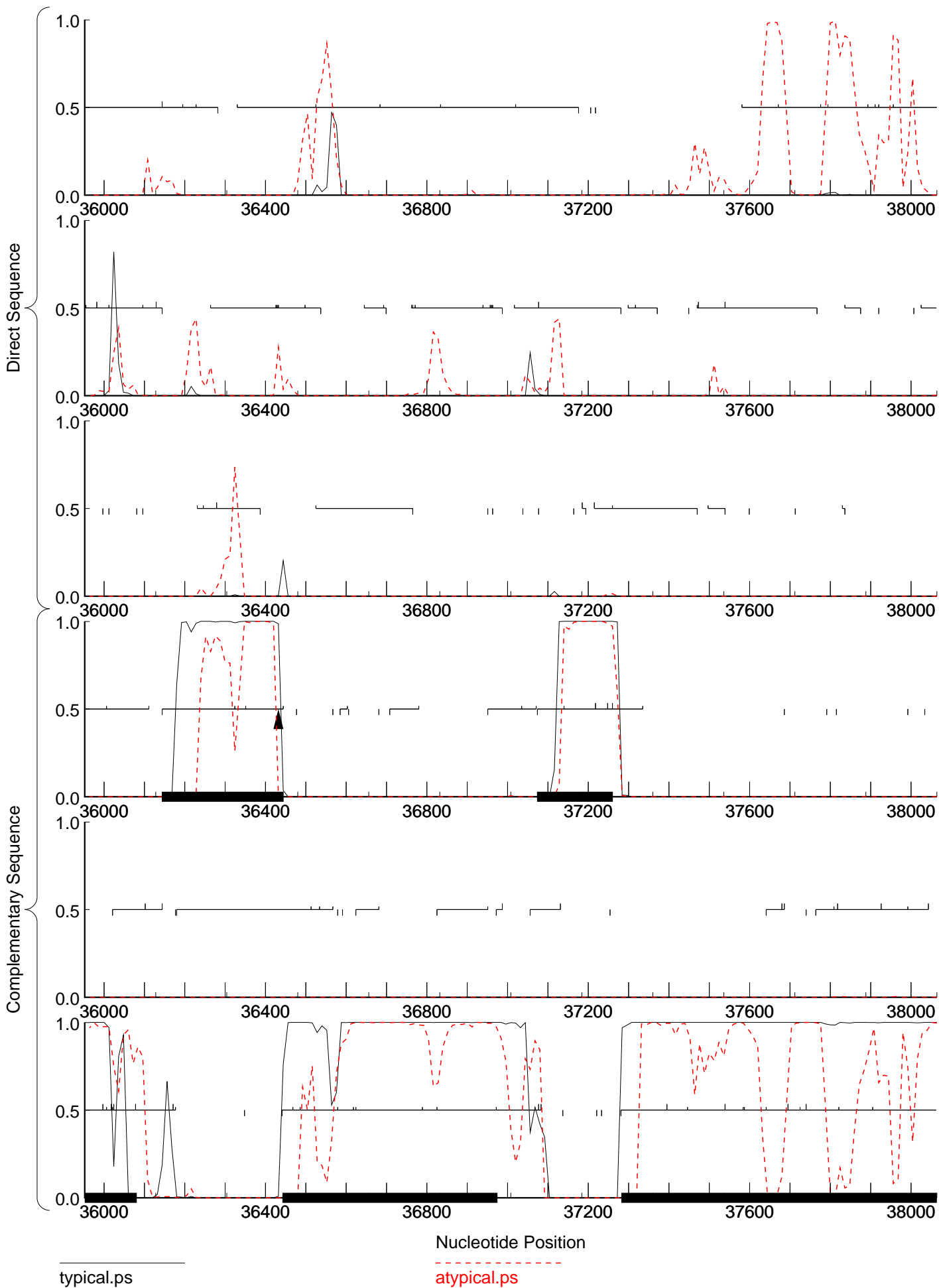


typical.ps

Nucleotide Position

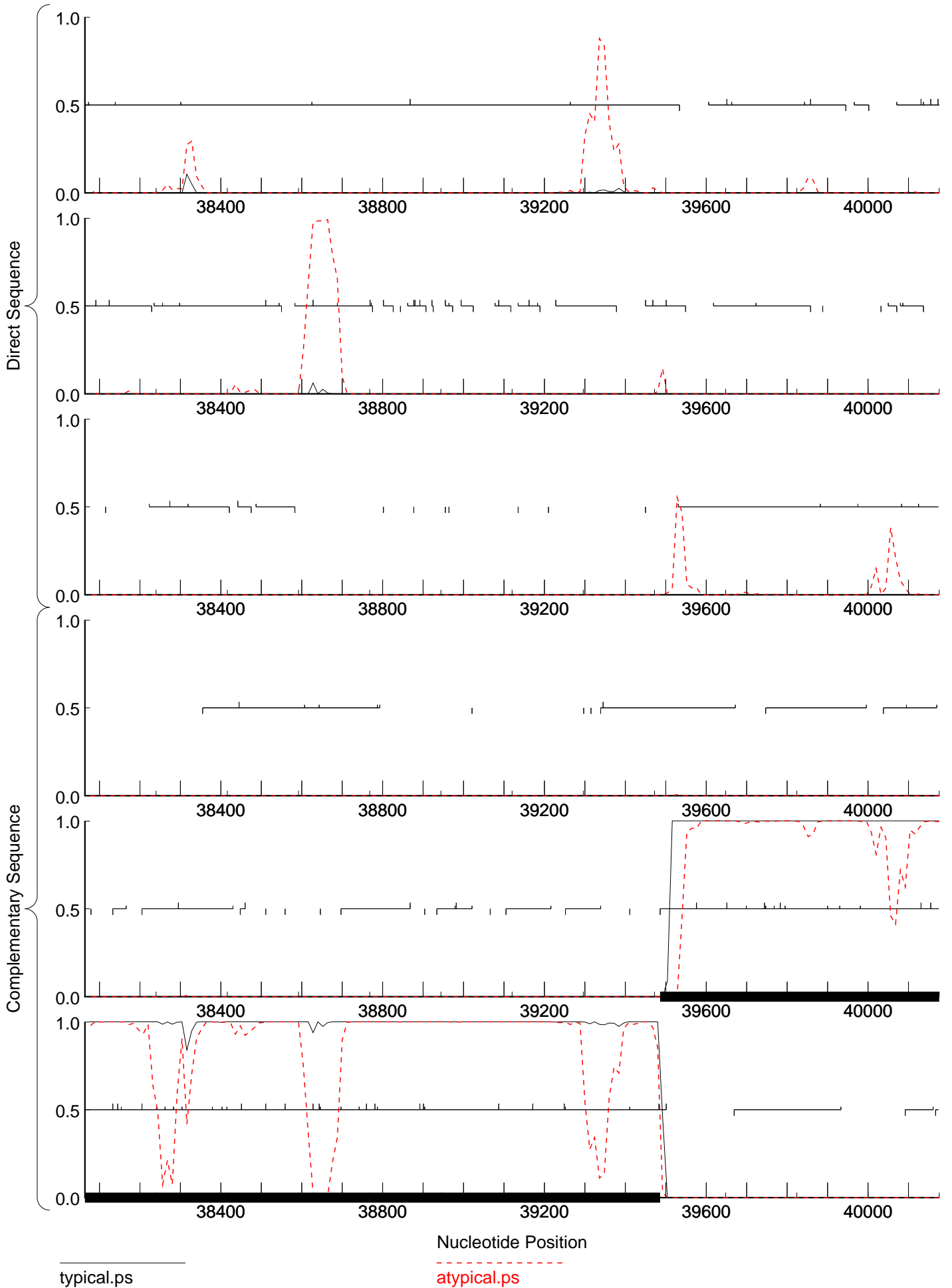
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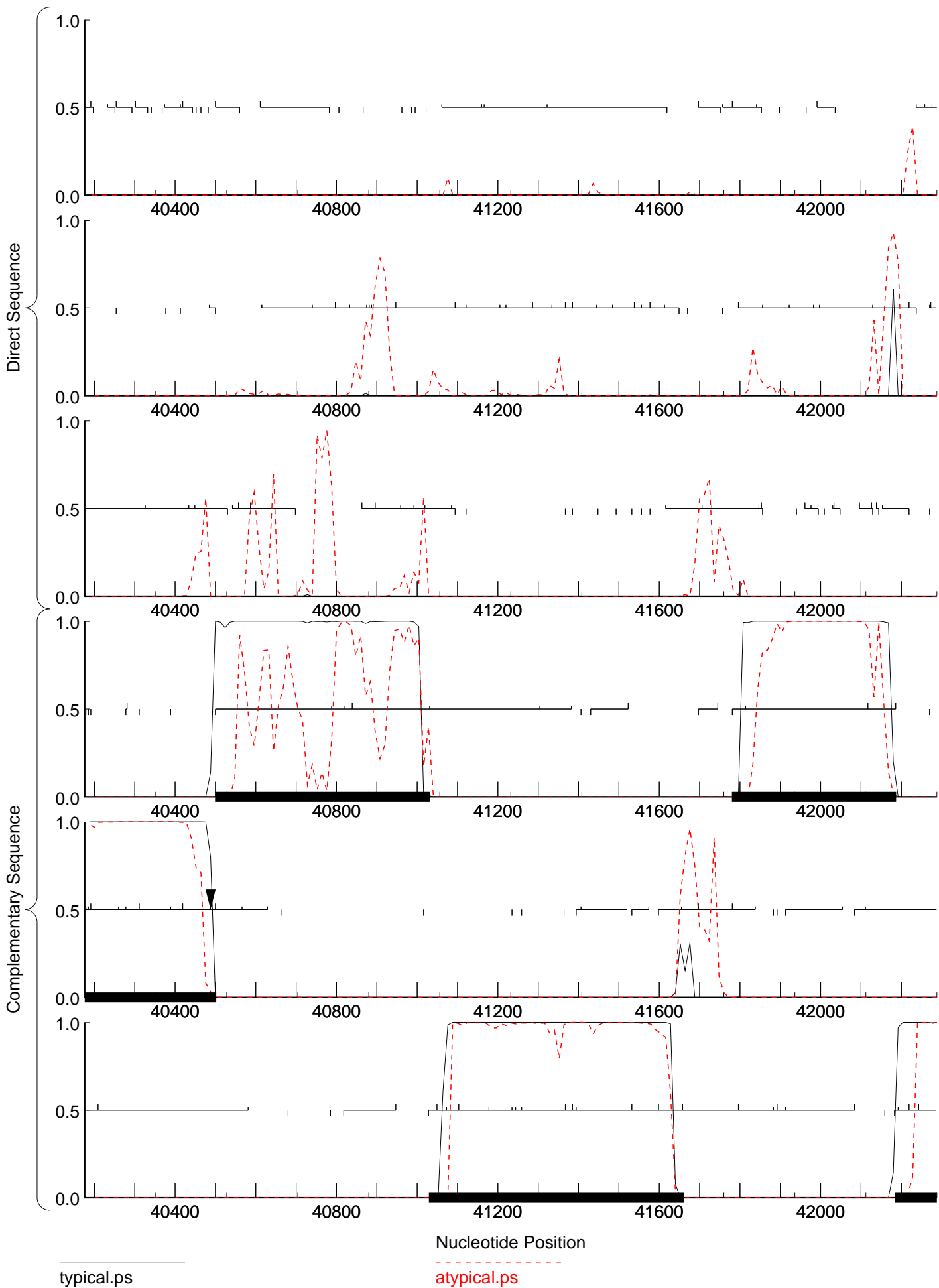


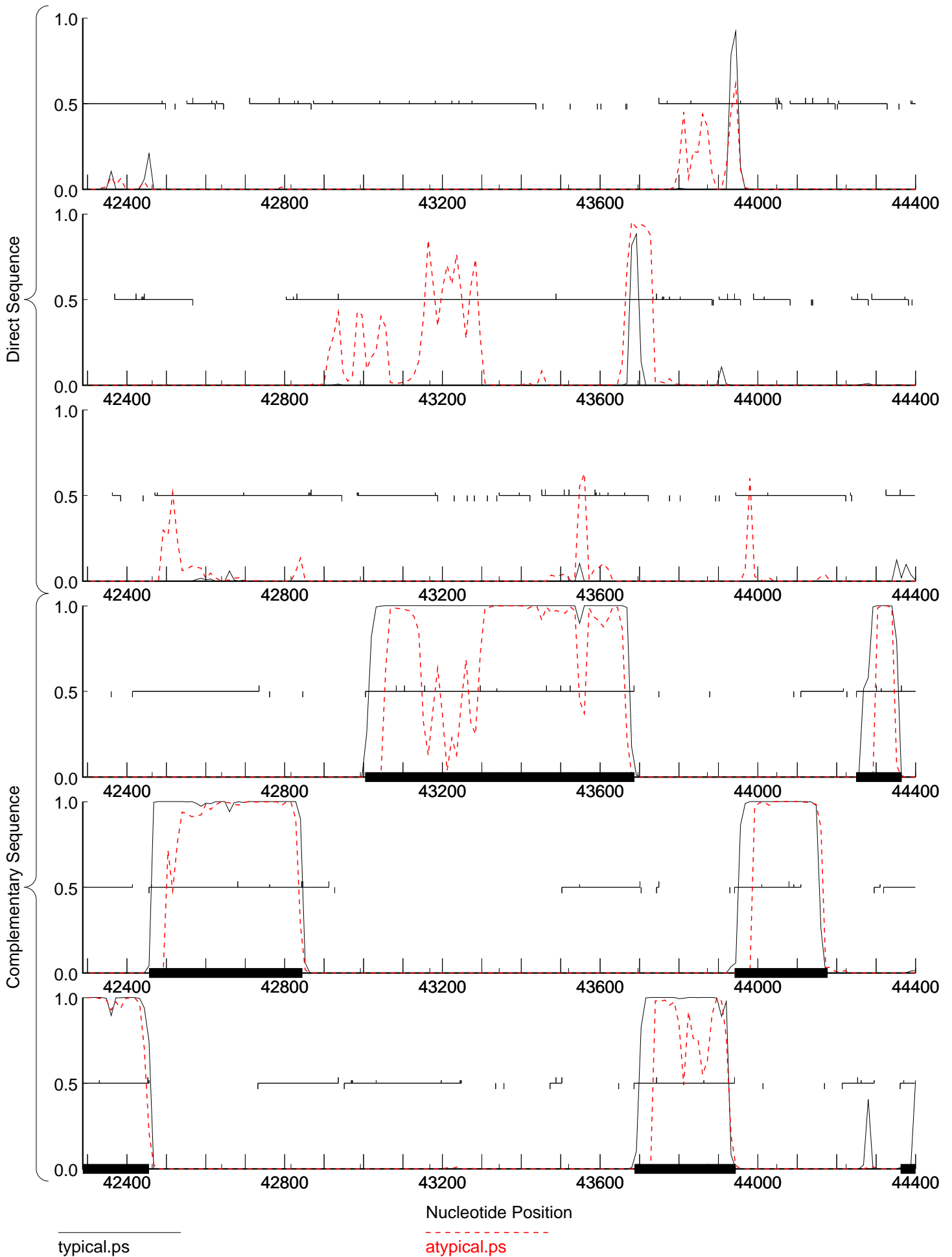


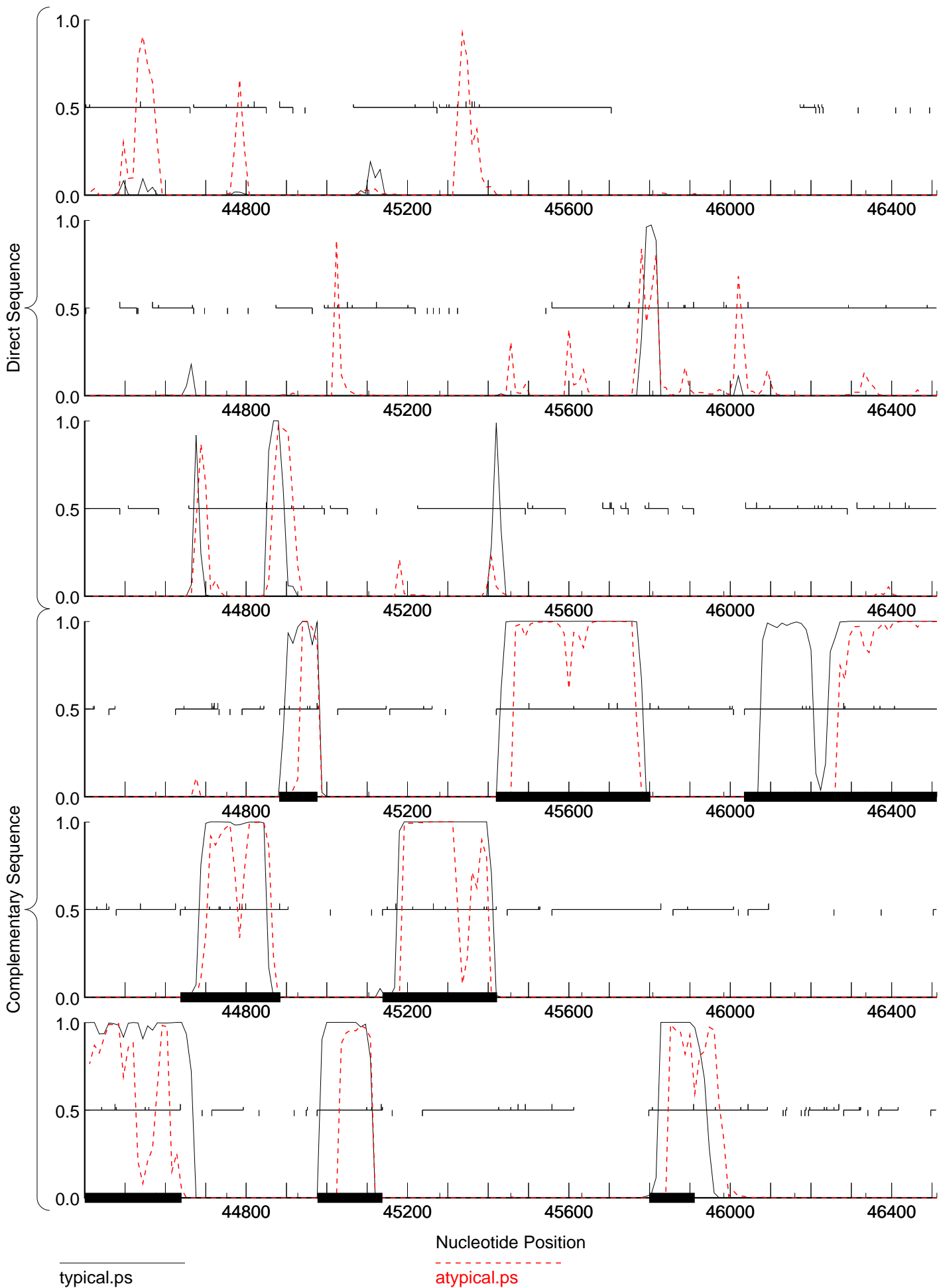
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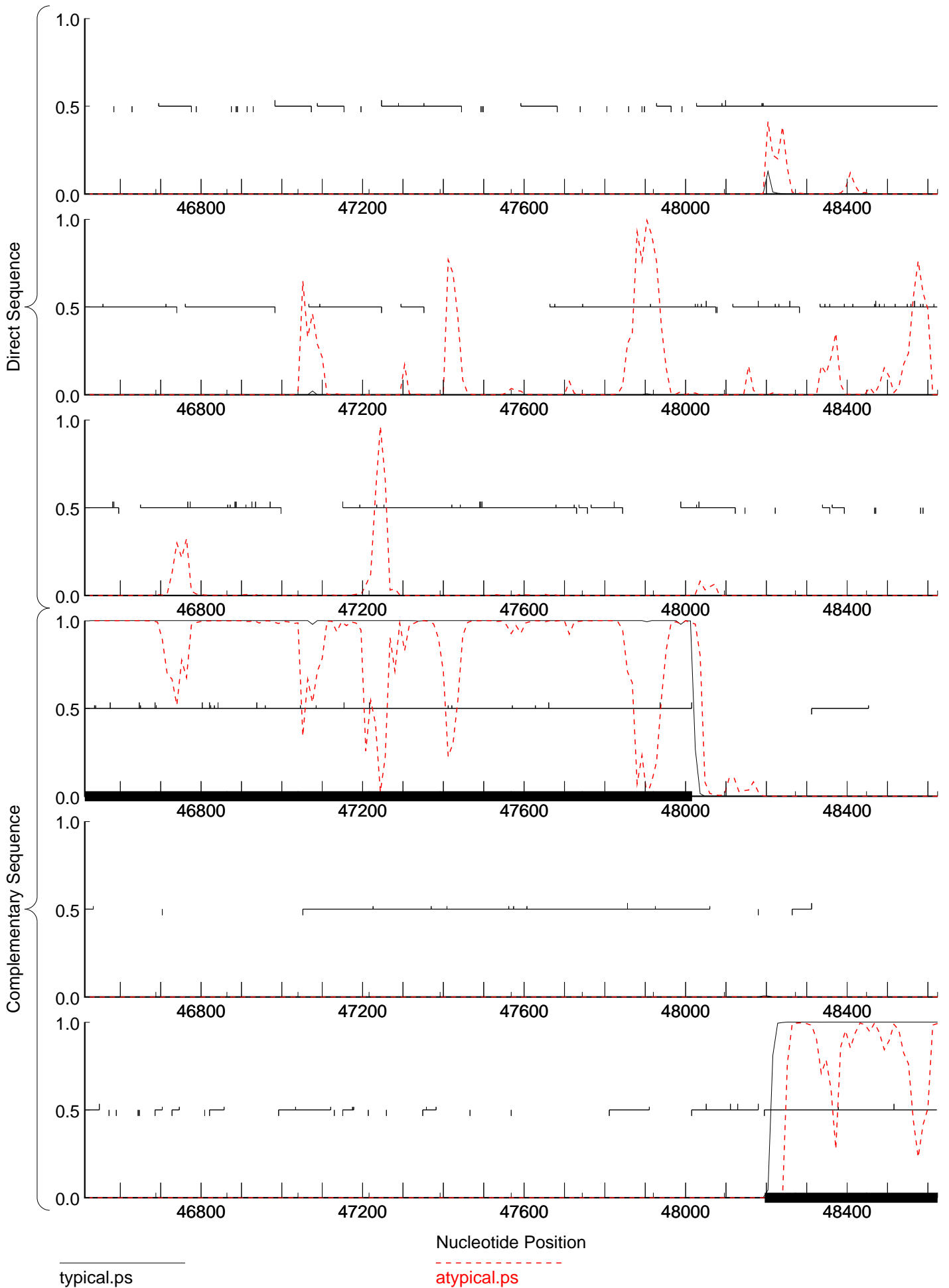
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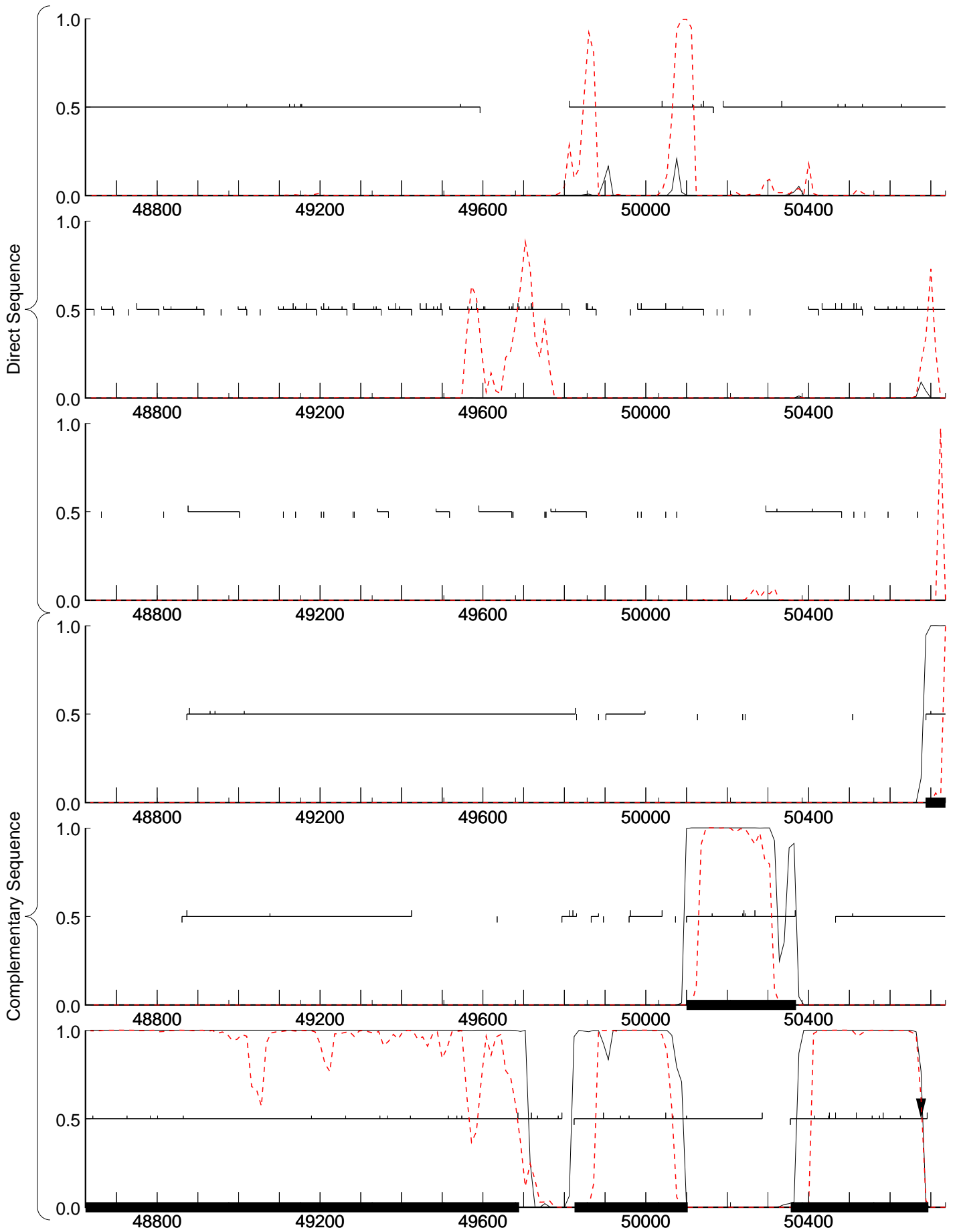






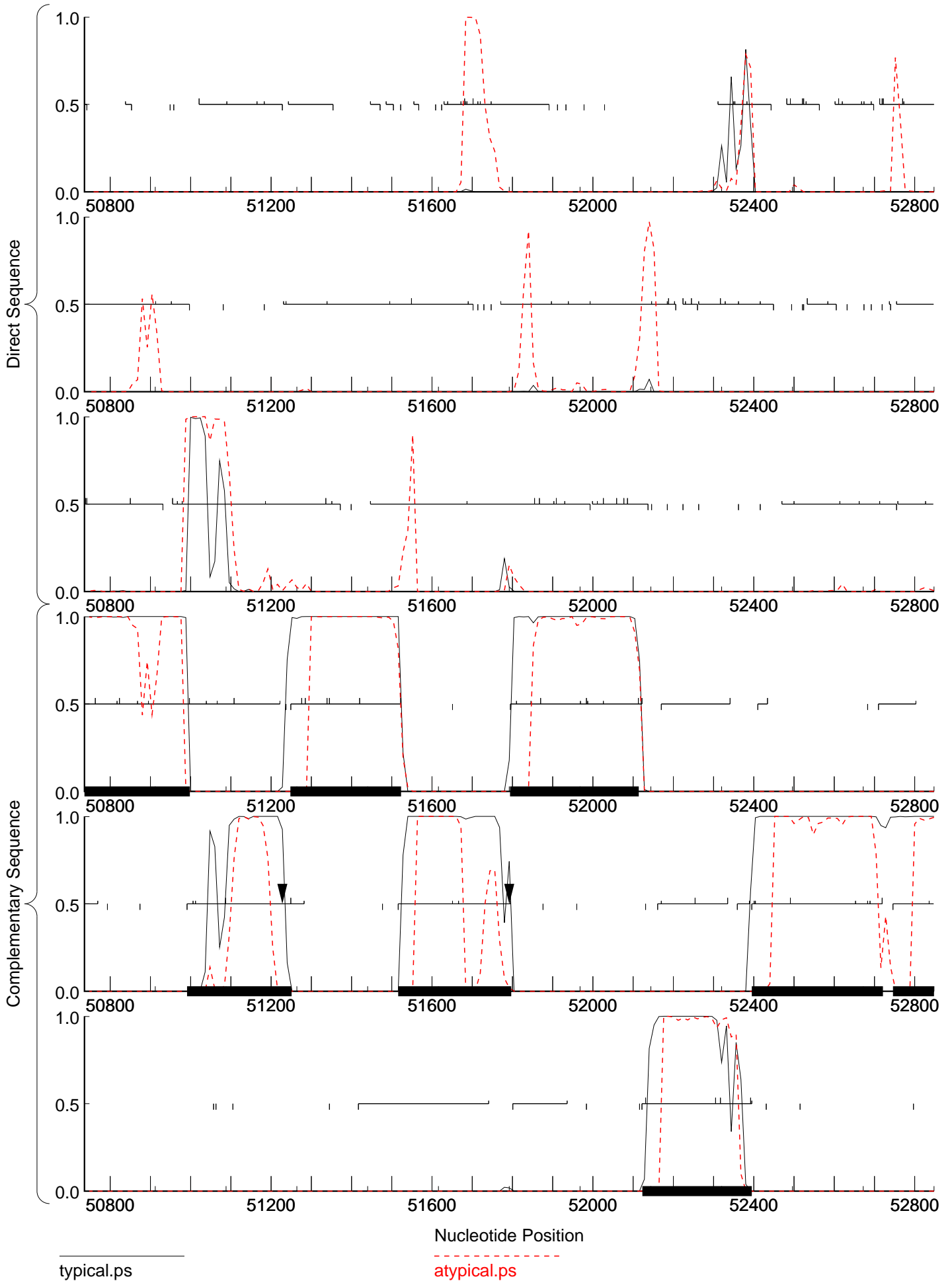


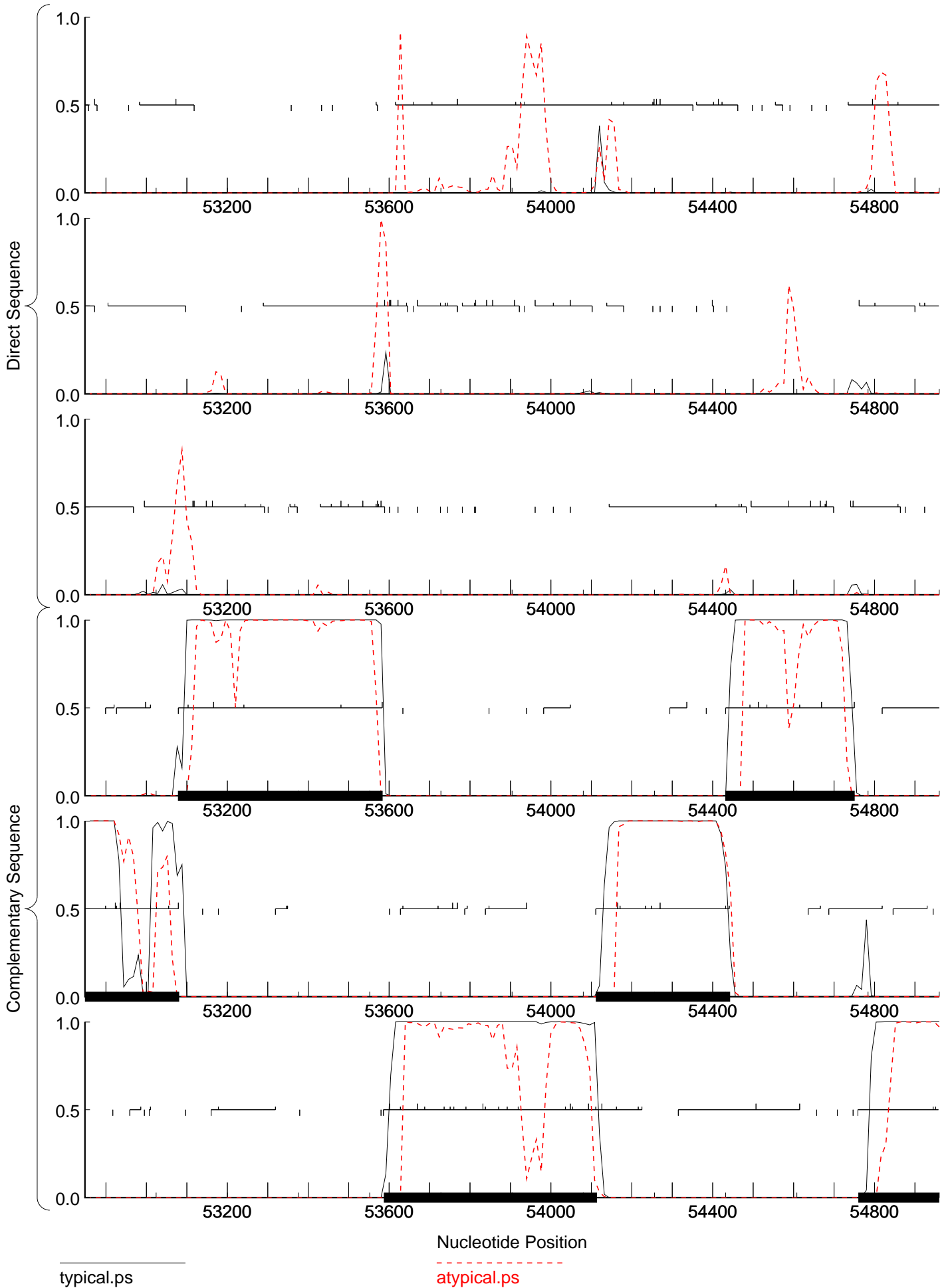


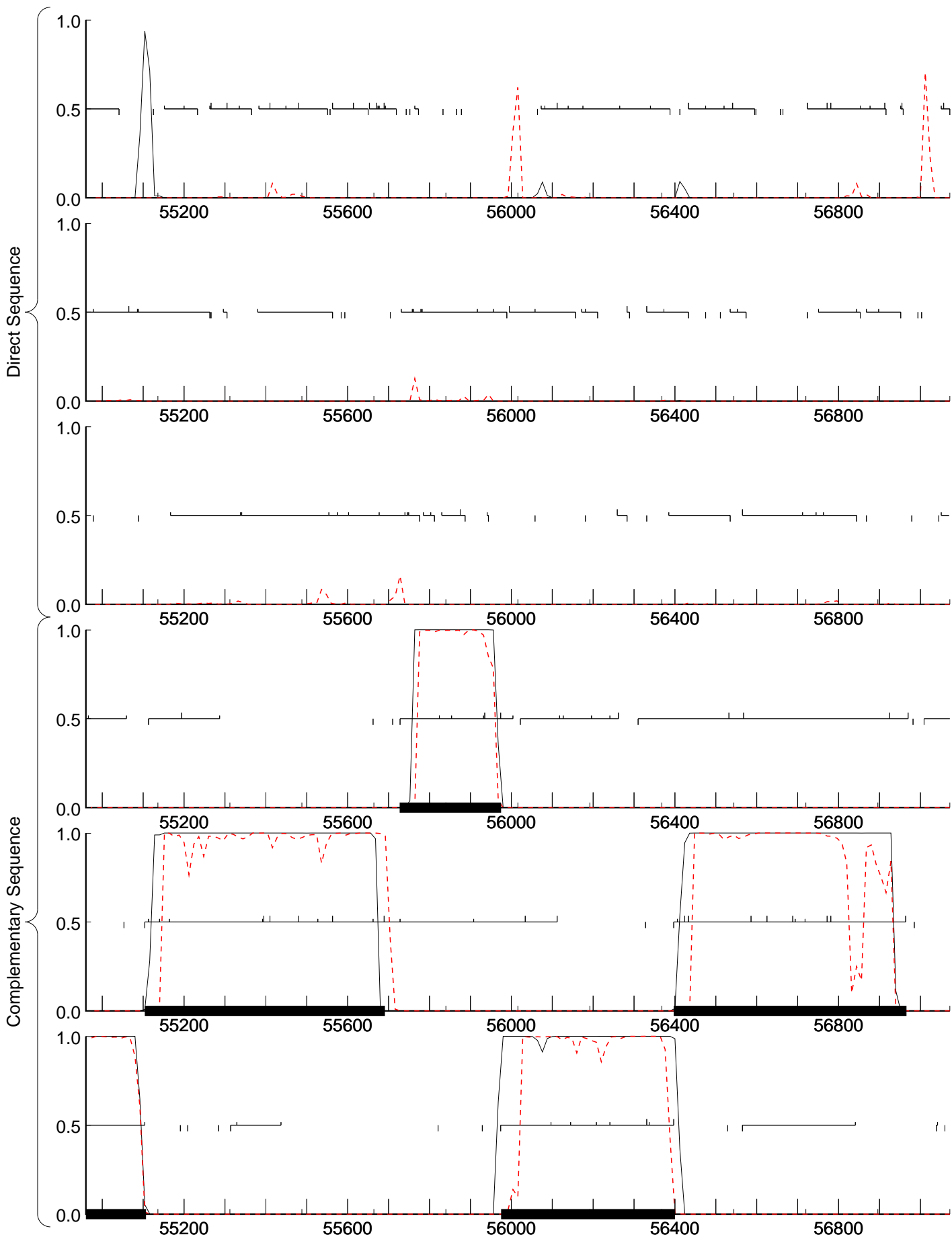


typical.ps

atypical.ps

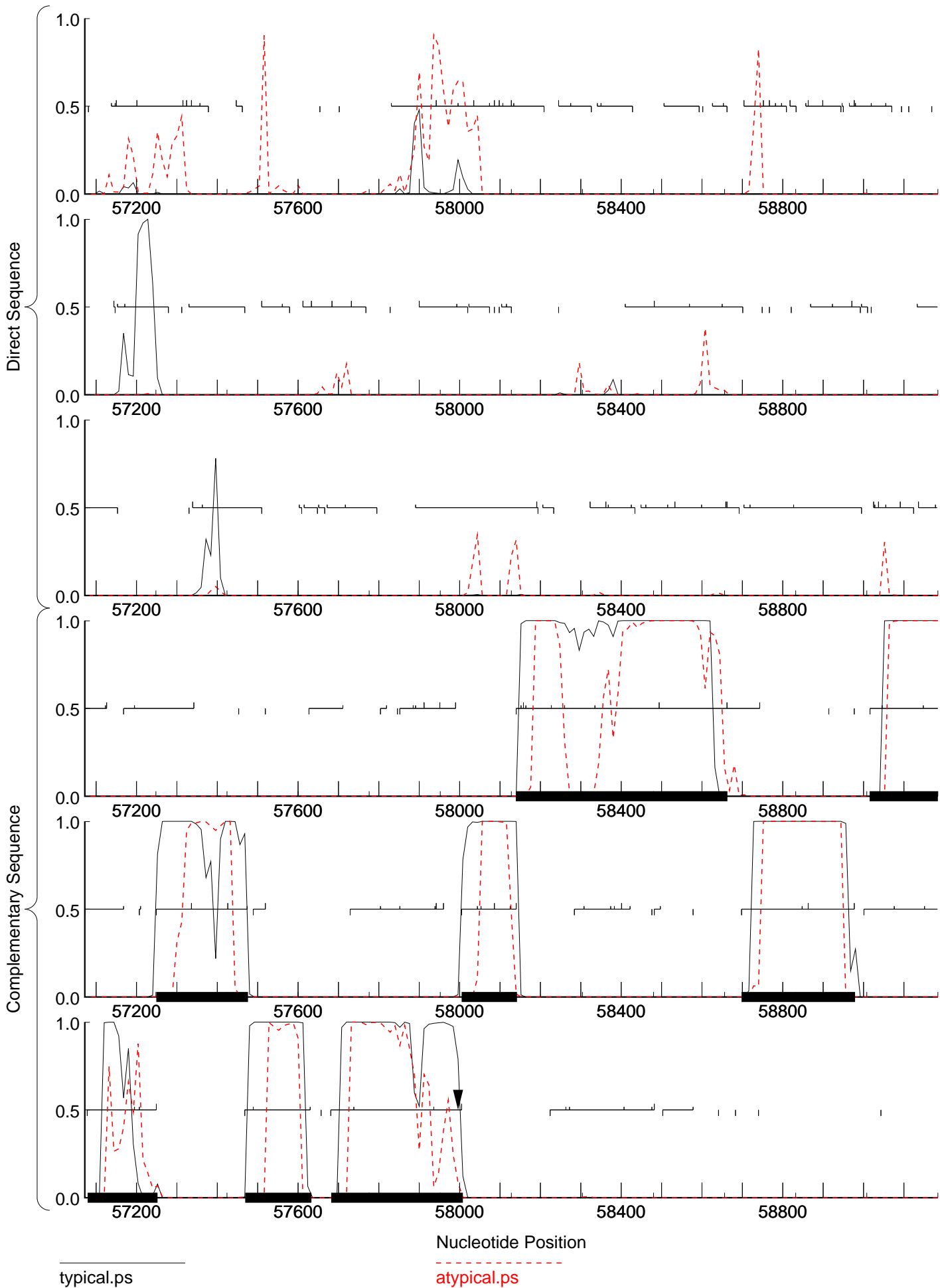






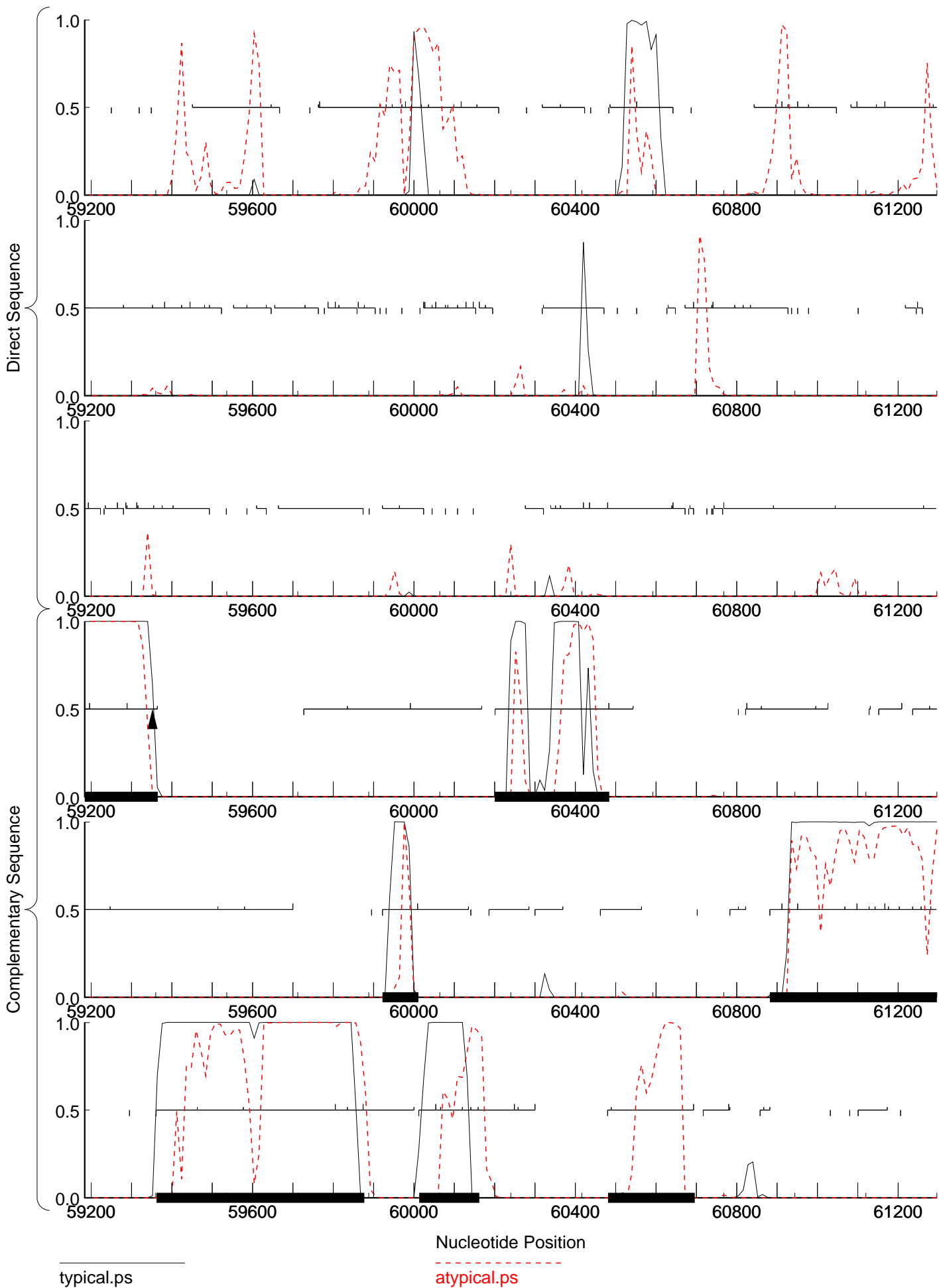
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