

GeneMark

Version 2.5p (09.08.06)

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PROGRAM INFORMATION

Sequence : Streptomyces phage Yaboi complete sequence, 131250 bp including 12433 bp terminal repeat, C
Analysis Date : 3/20/18 at 21:29:18
Pages : 64
Sequence Length : 131250 bp
GC Content : 49.30%

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,

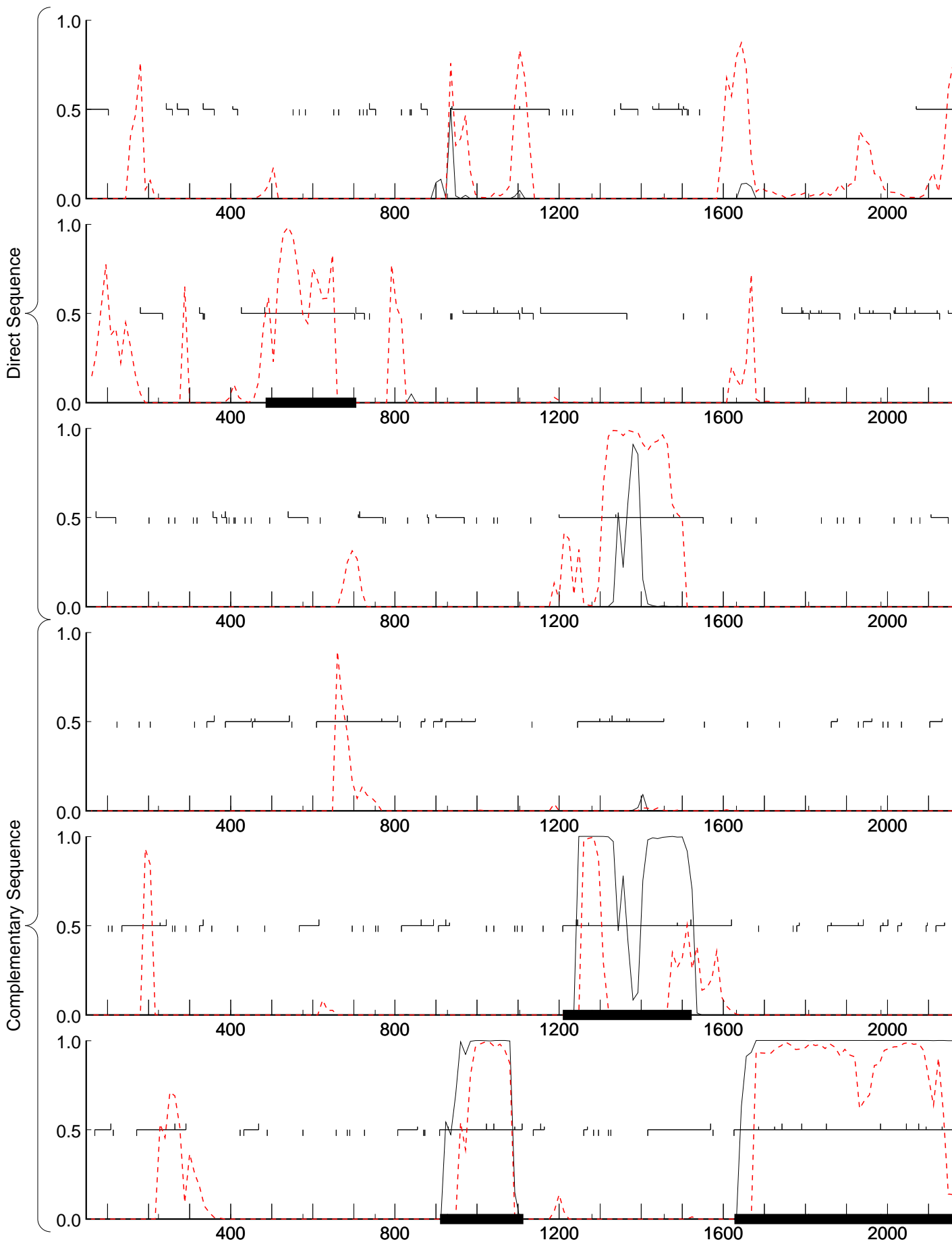
Matrix notes & comments

Training set derived by GeneMarkS, 4.30 October 2014
Tue Mar 20 21:29:17 2018

MATRIX INFORMATION

Matrix : -
Author : -
Order : 4

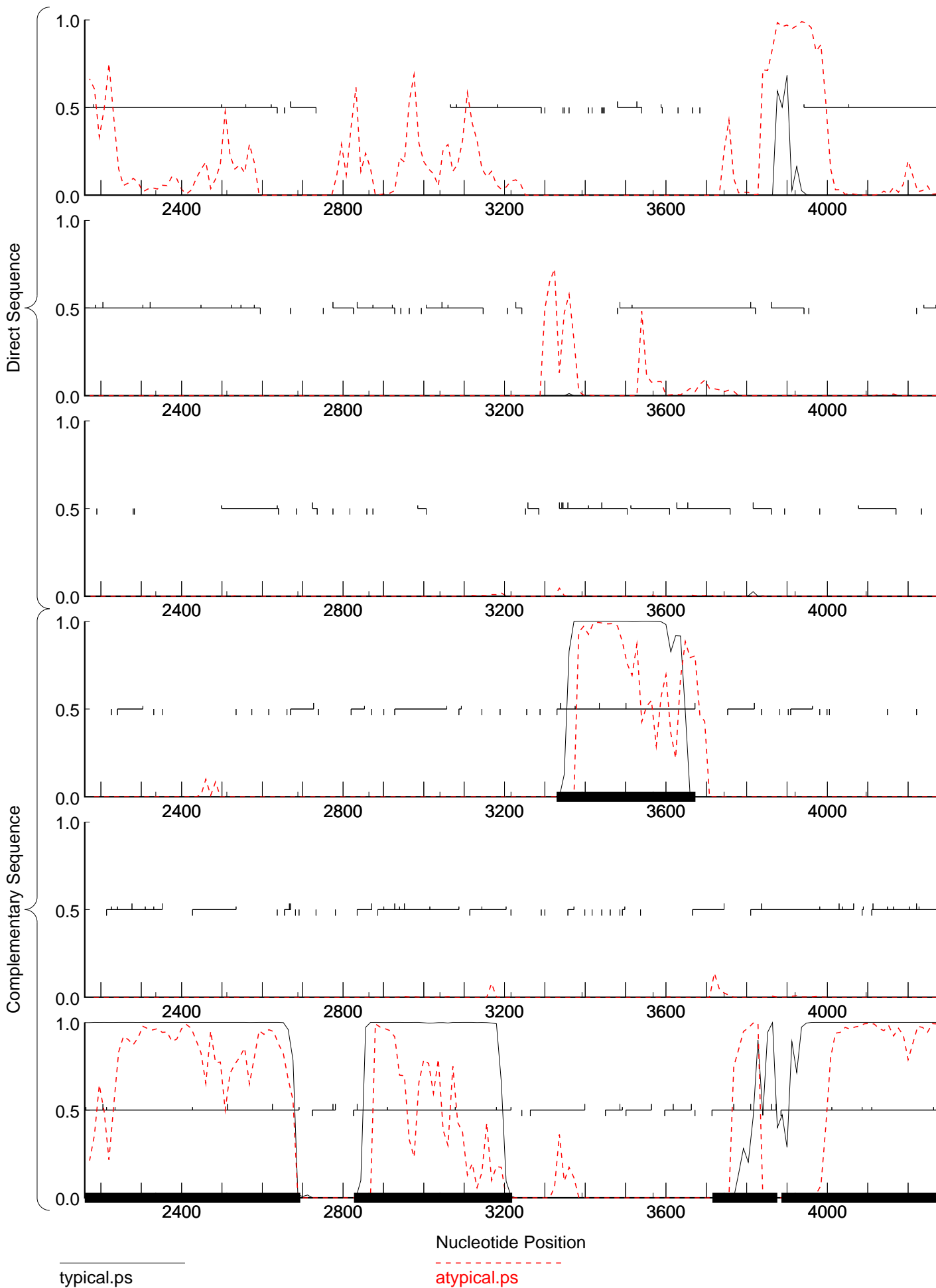
Send questions / comments to:
Dr. M. Borodovsky
Georgia Institute of Technology
School of Biology
Atlanta, GA 30332-0230



typical.ps

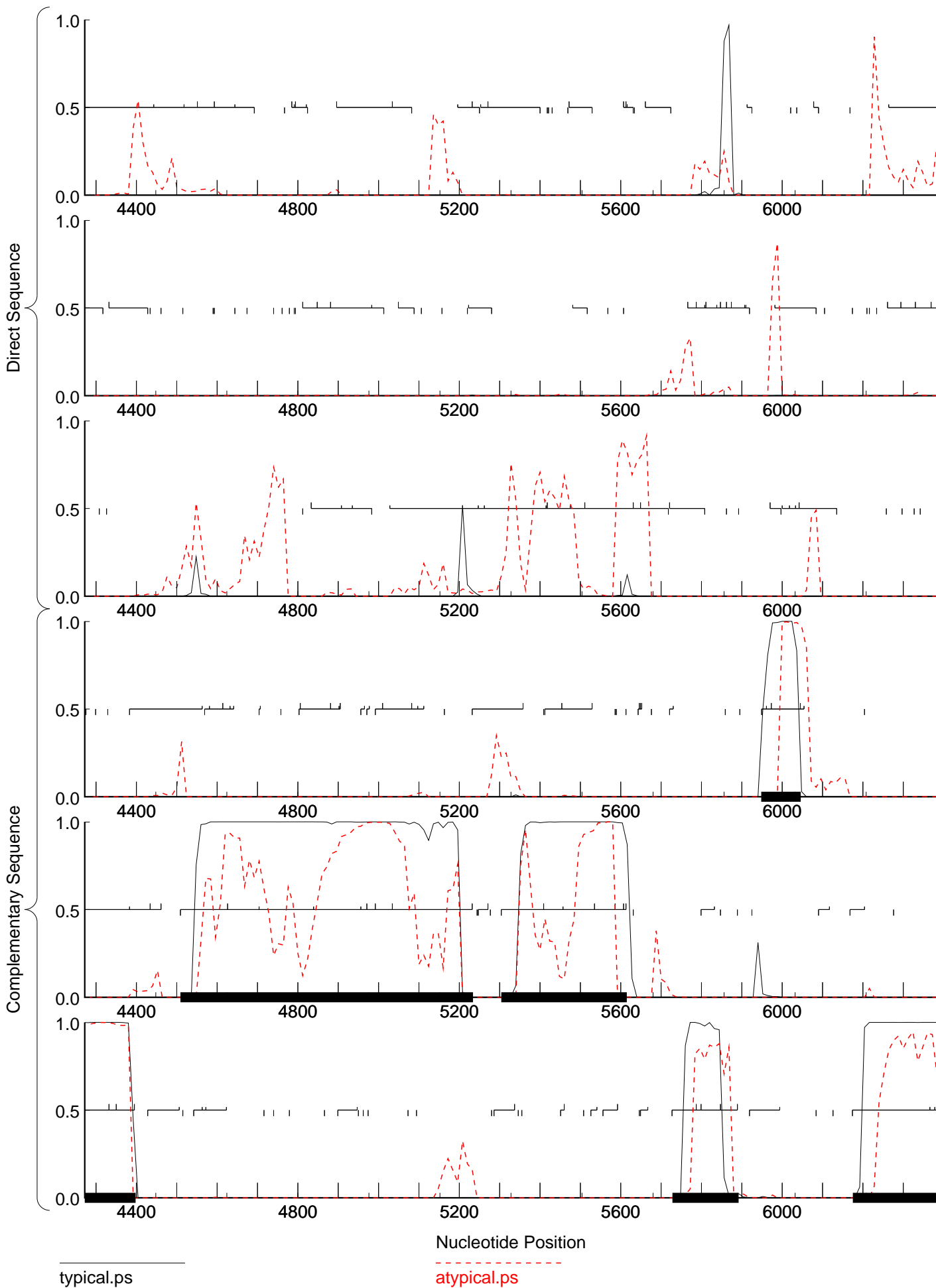
atypical.ps

phage Yaboi complete sequence, 131250 bp including 12433 bp terminal repeat, Cluster BE2, Order 4, Window 96, Step 12, 3/64
phage Yaboi complete sequence, 131250 bp including 12433 bp terminal repeat, Cluster BE2, Order 2, Window 96, Step 12, 3/64



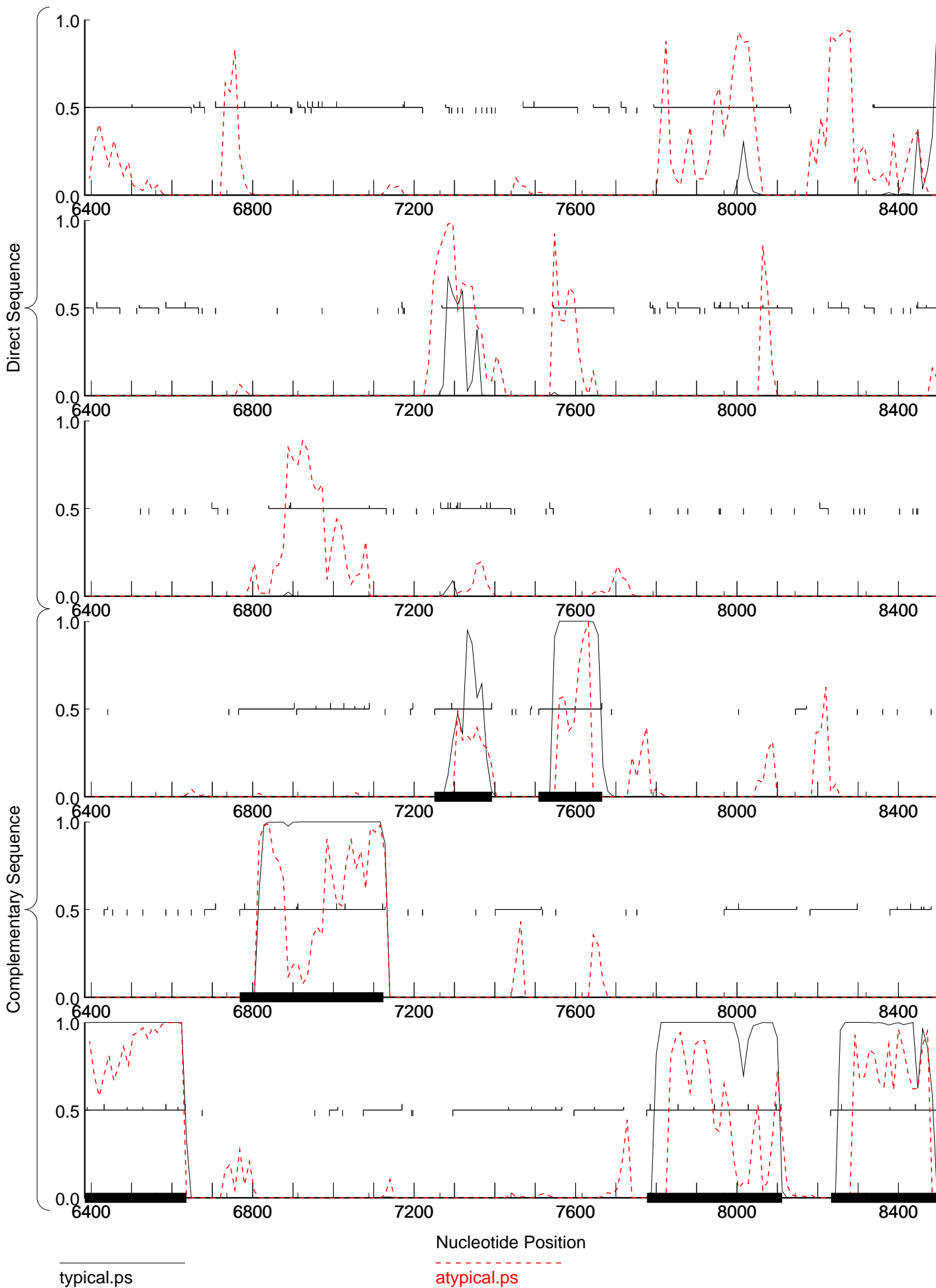
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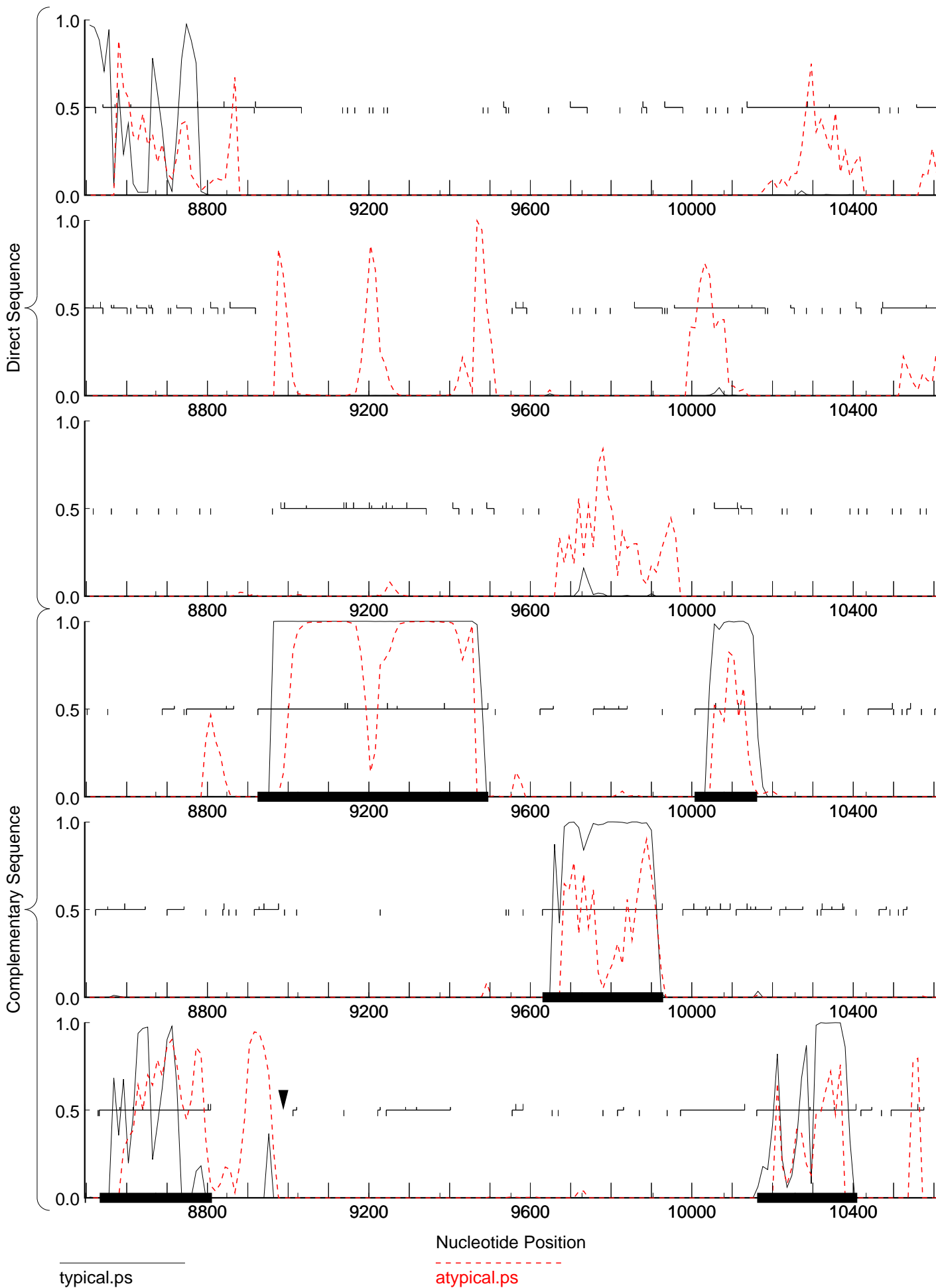
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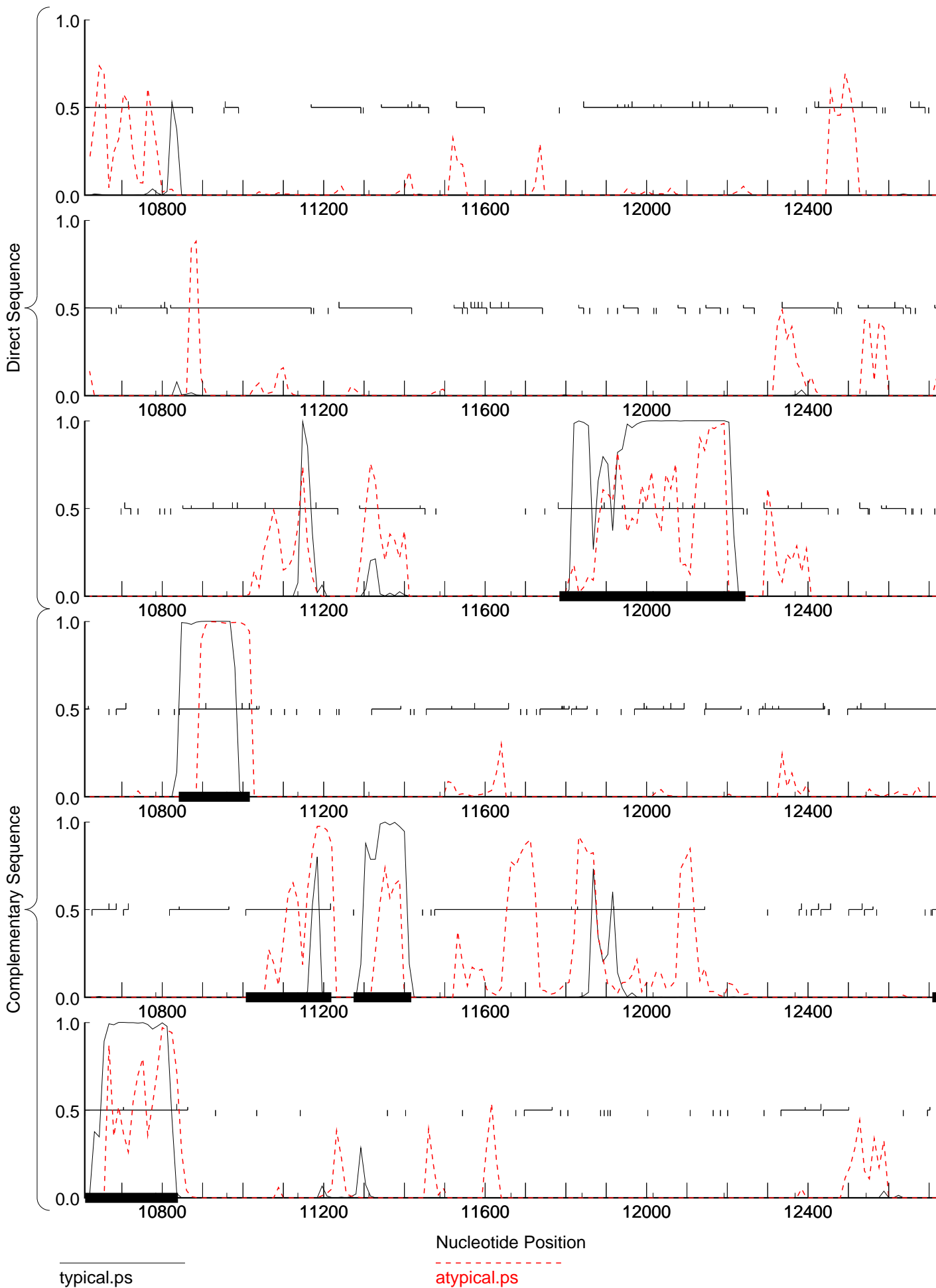
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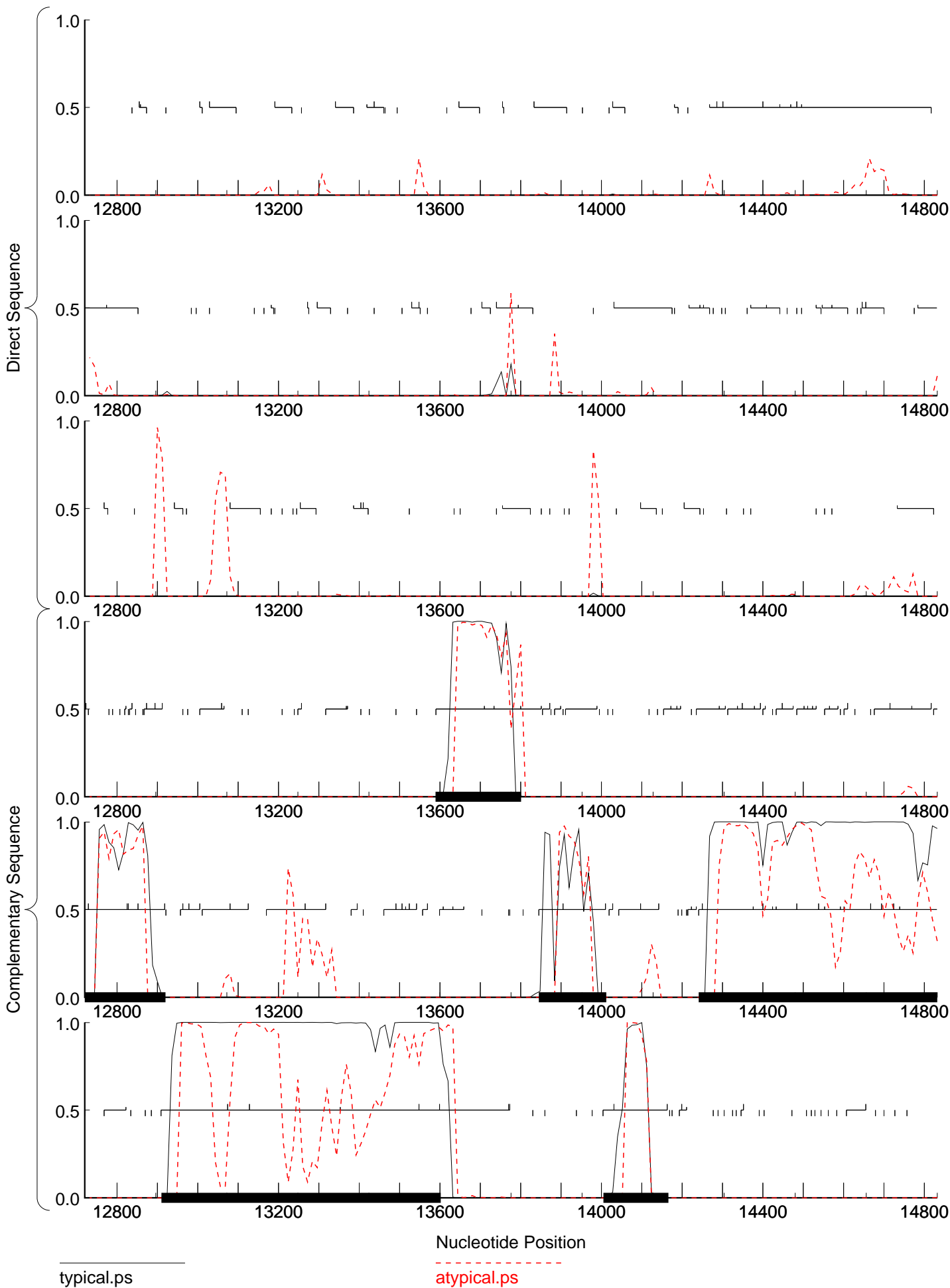


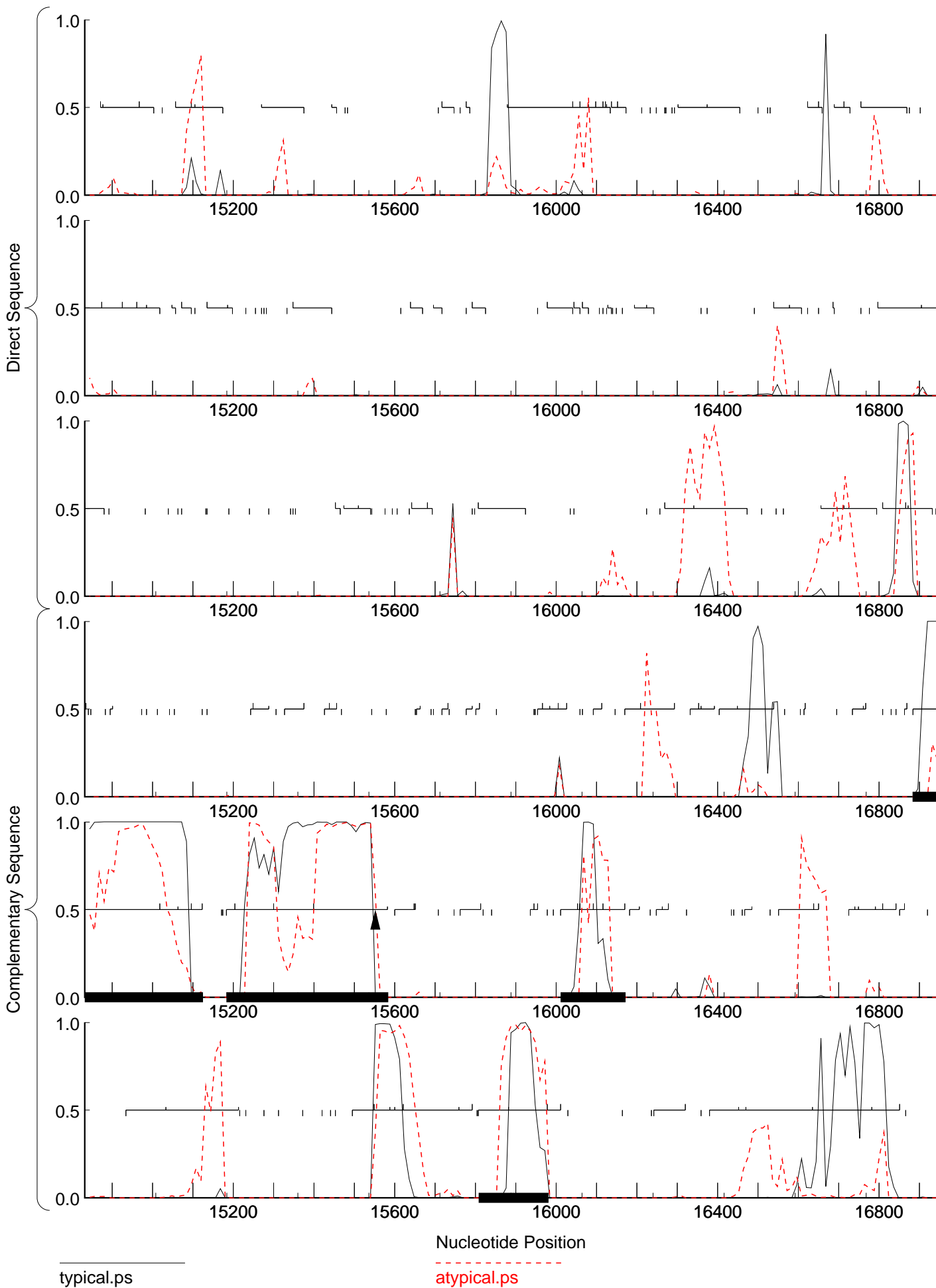
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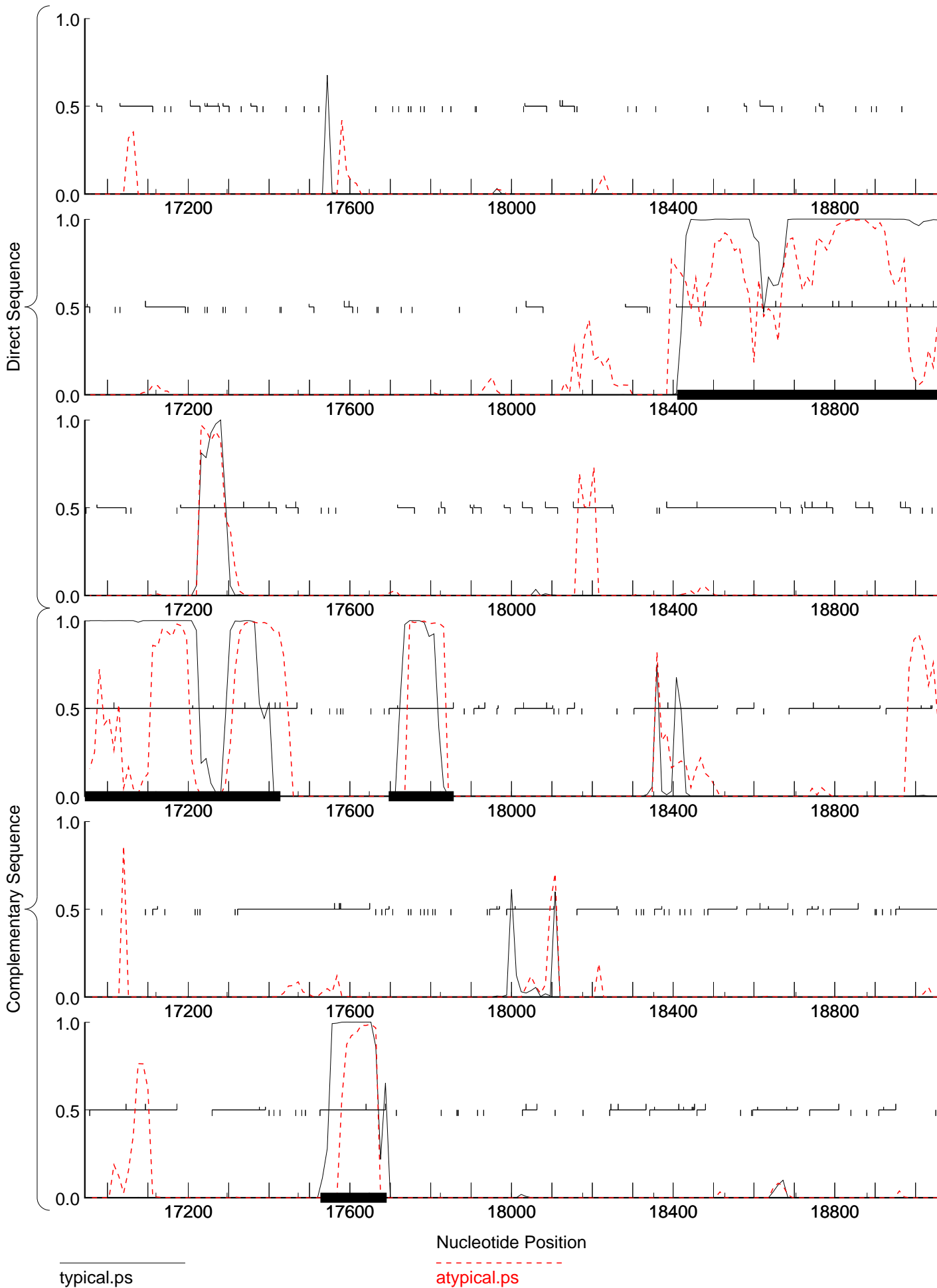
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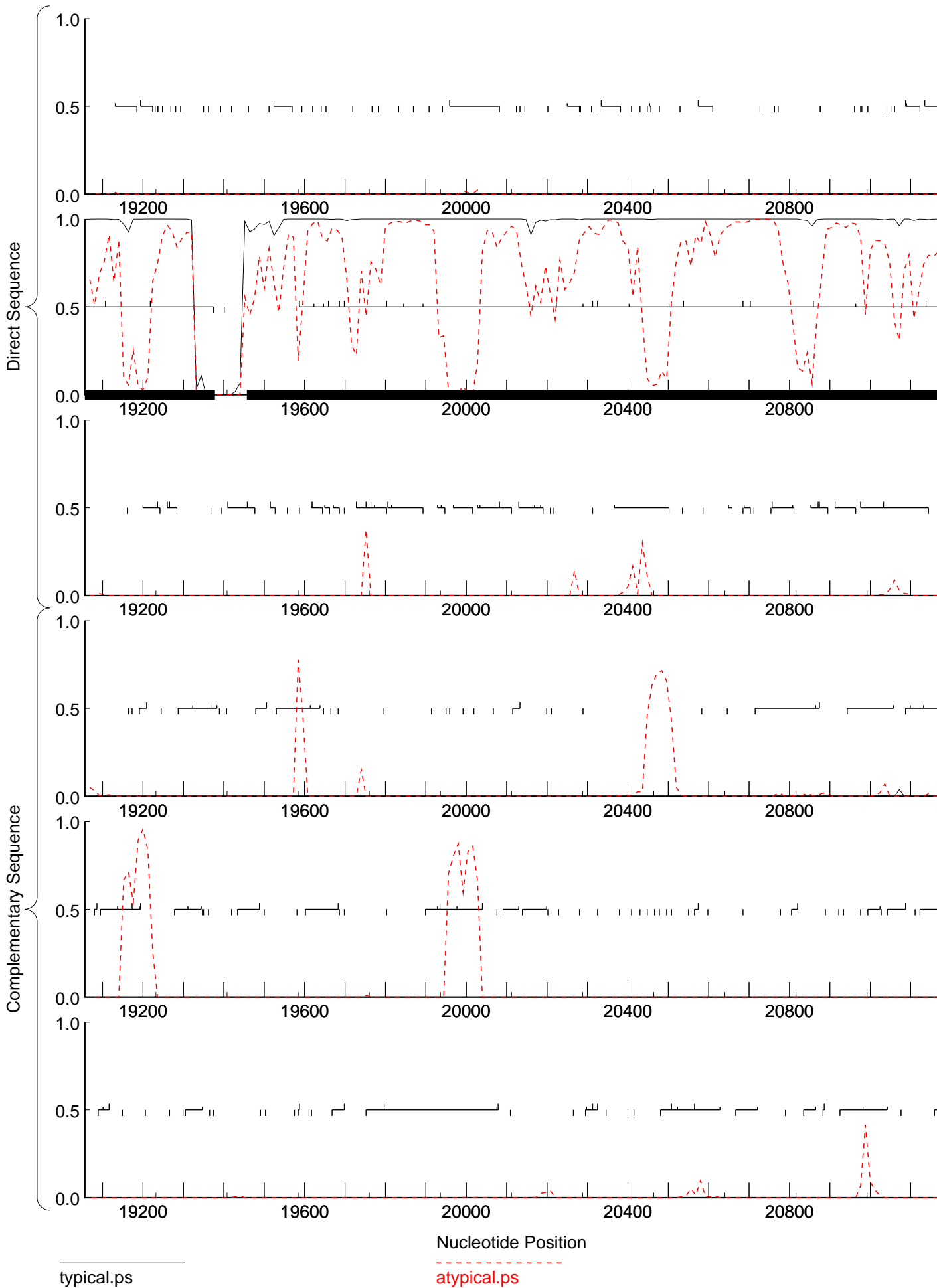


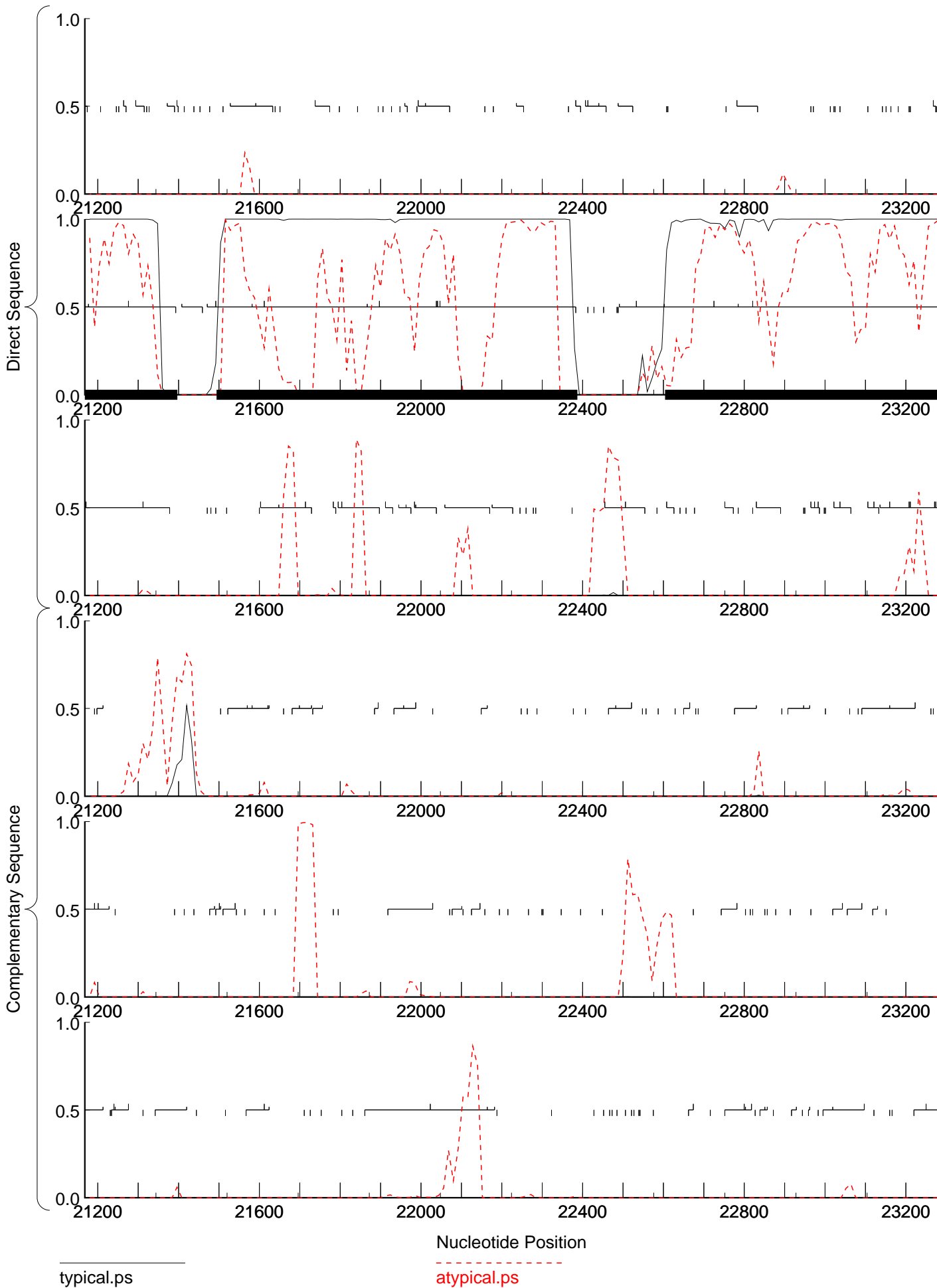


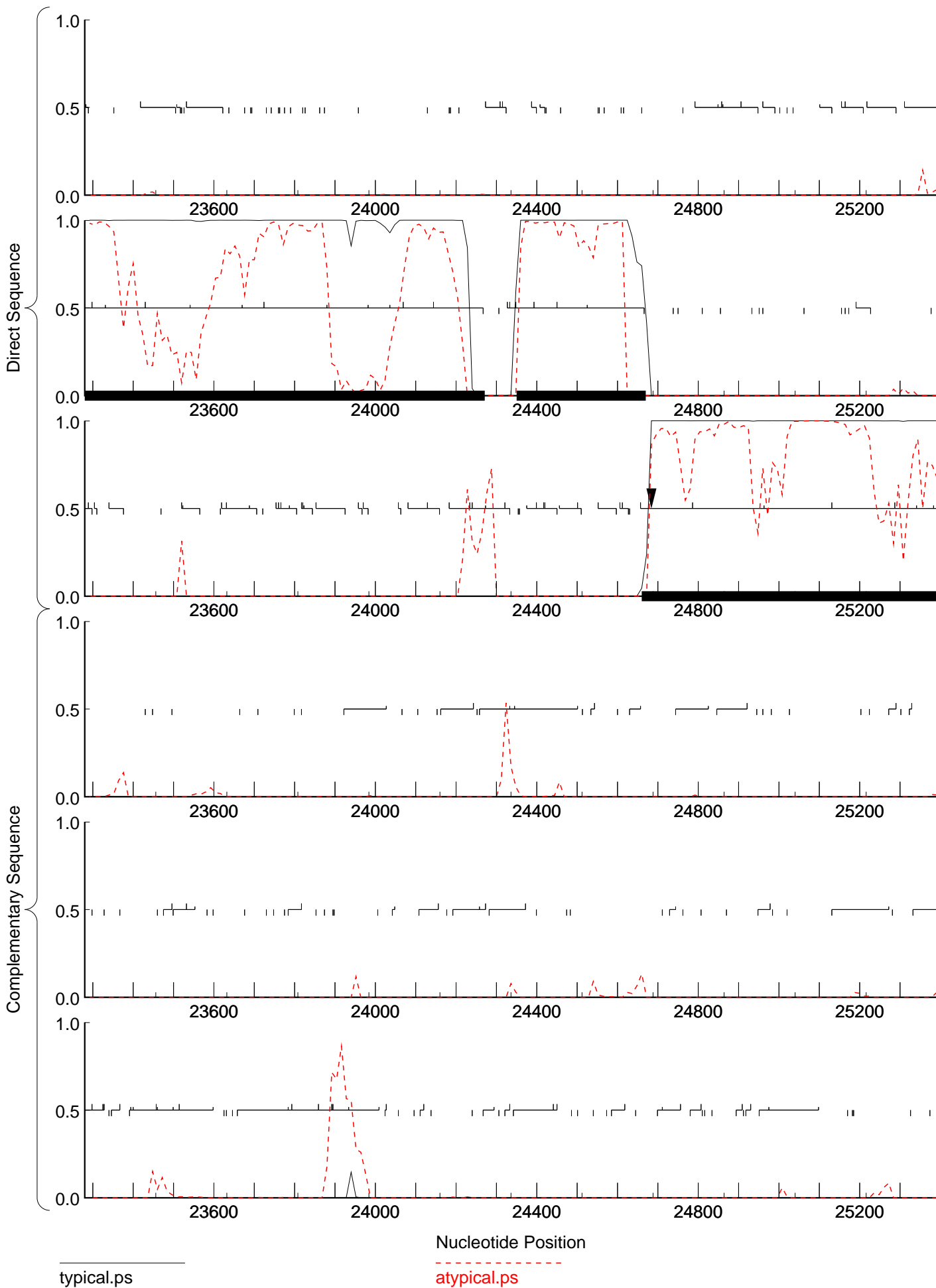


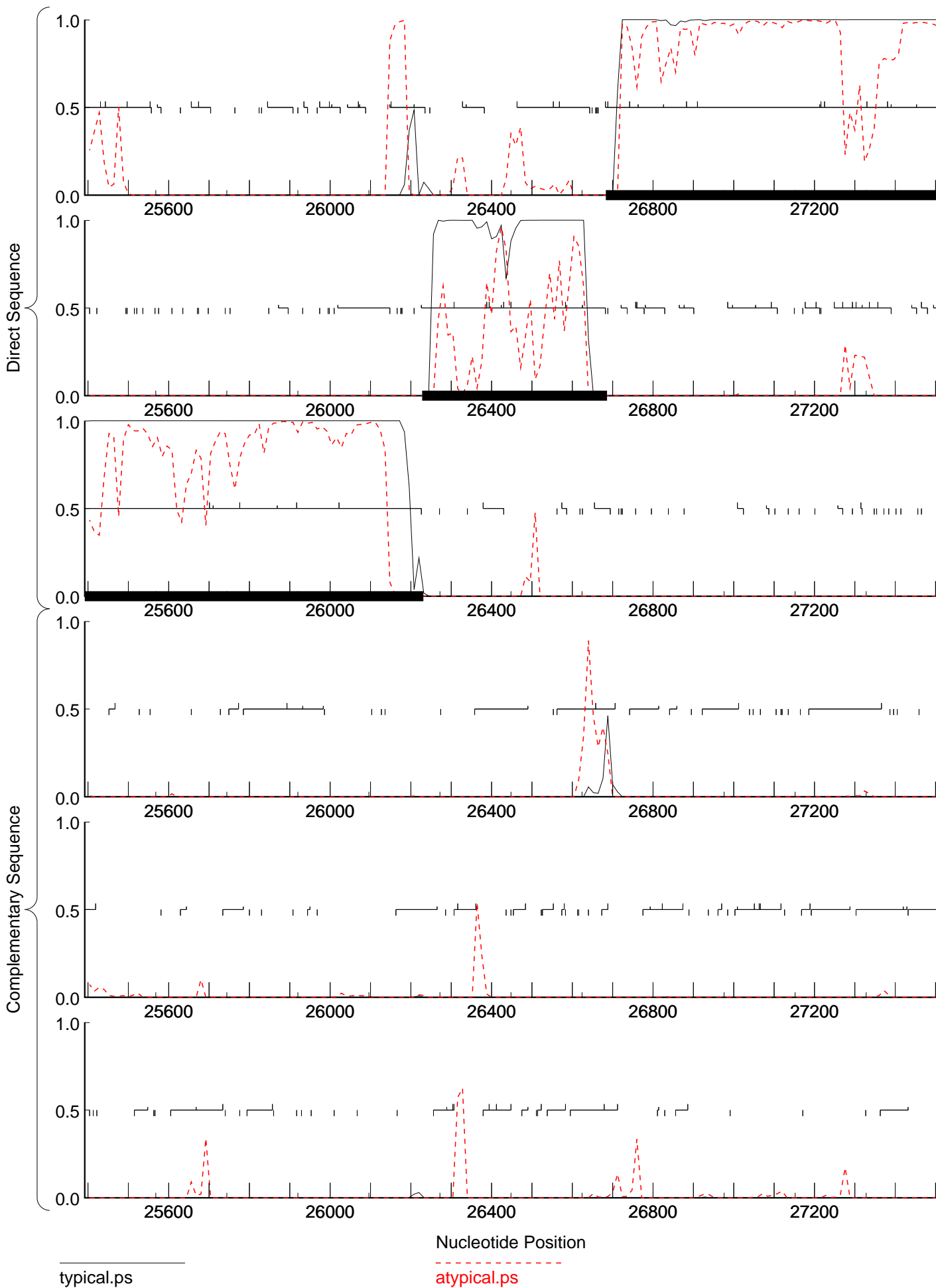
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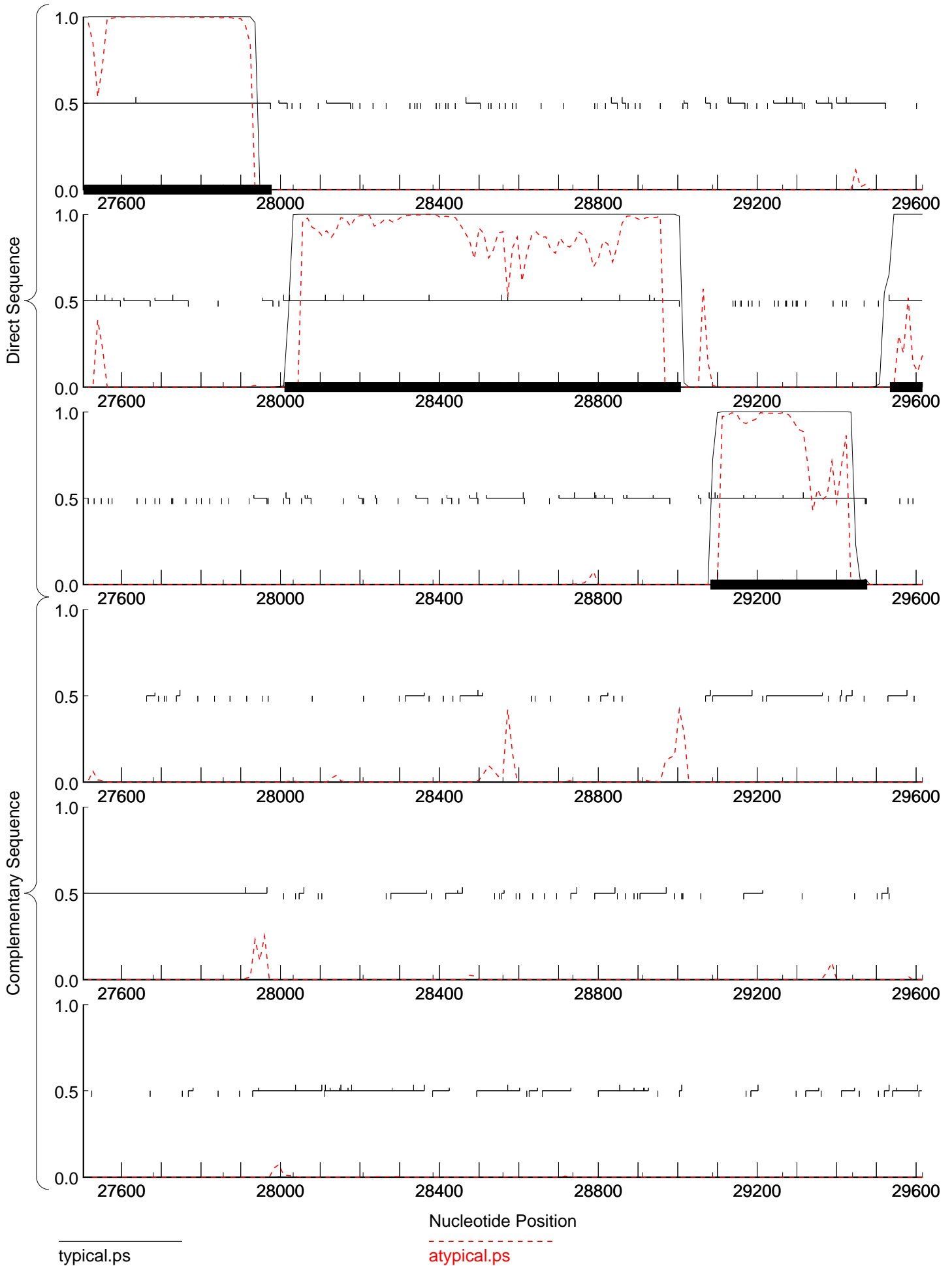
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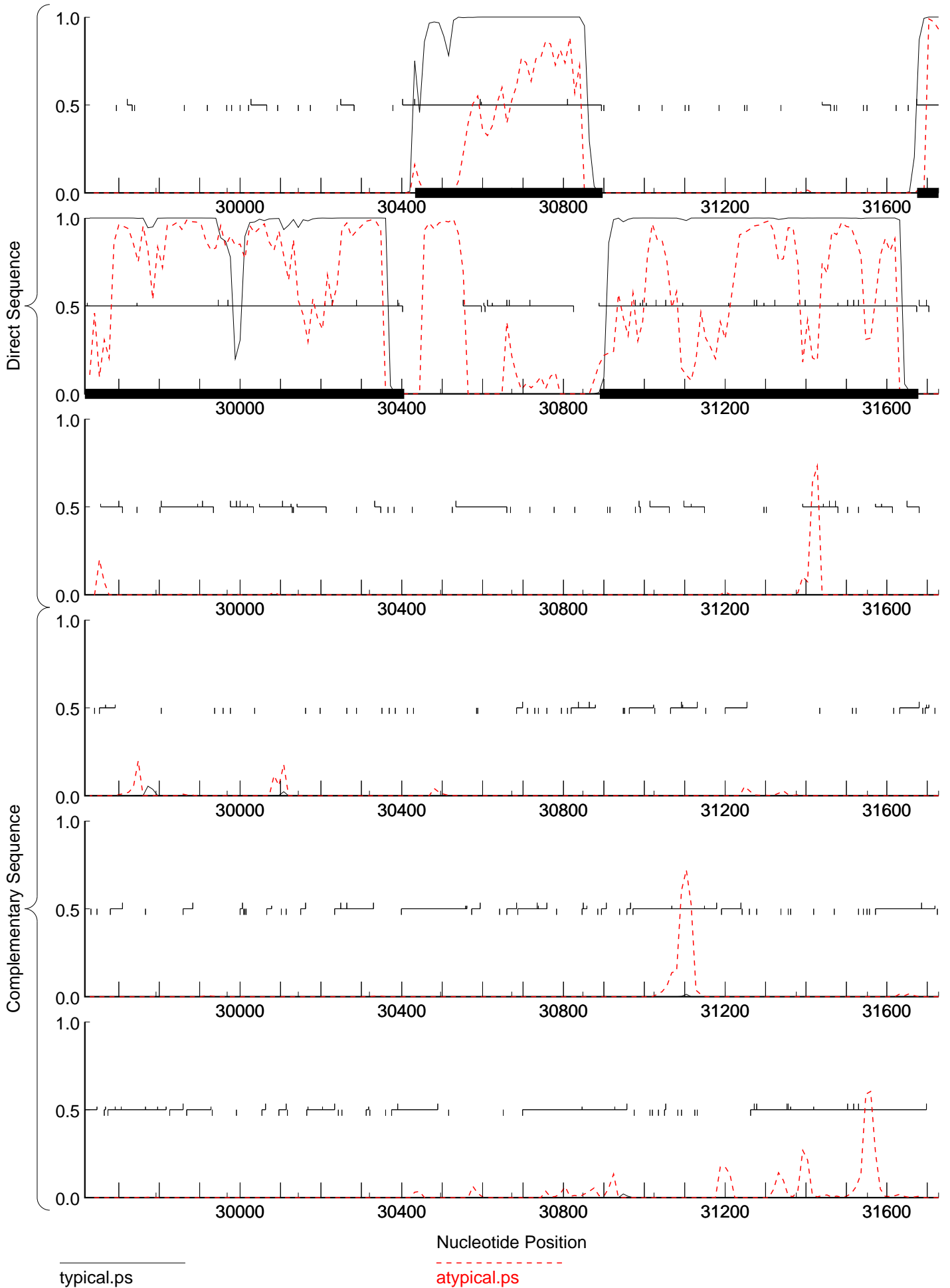


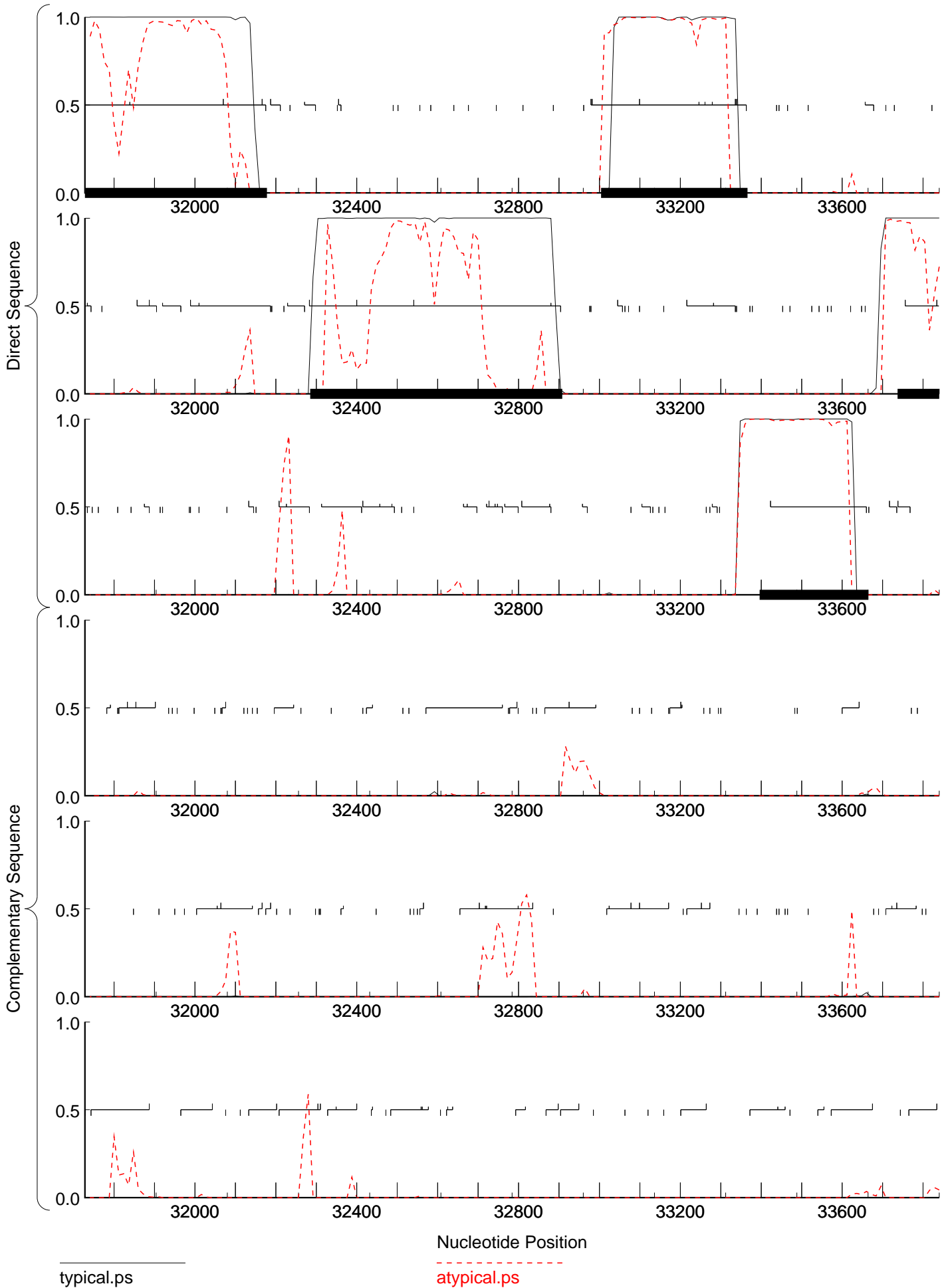


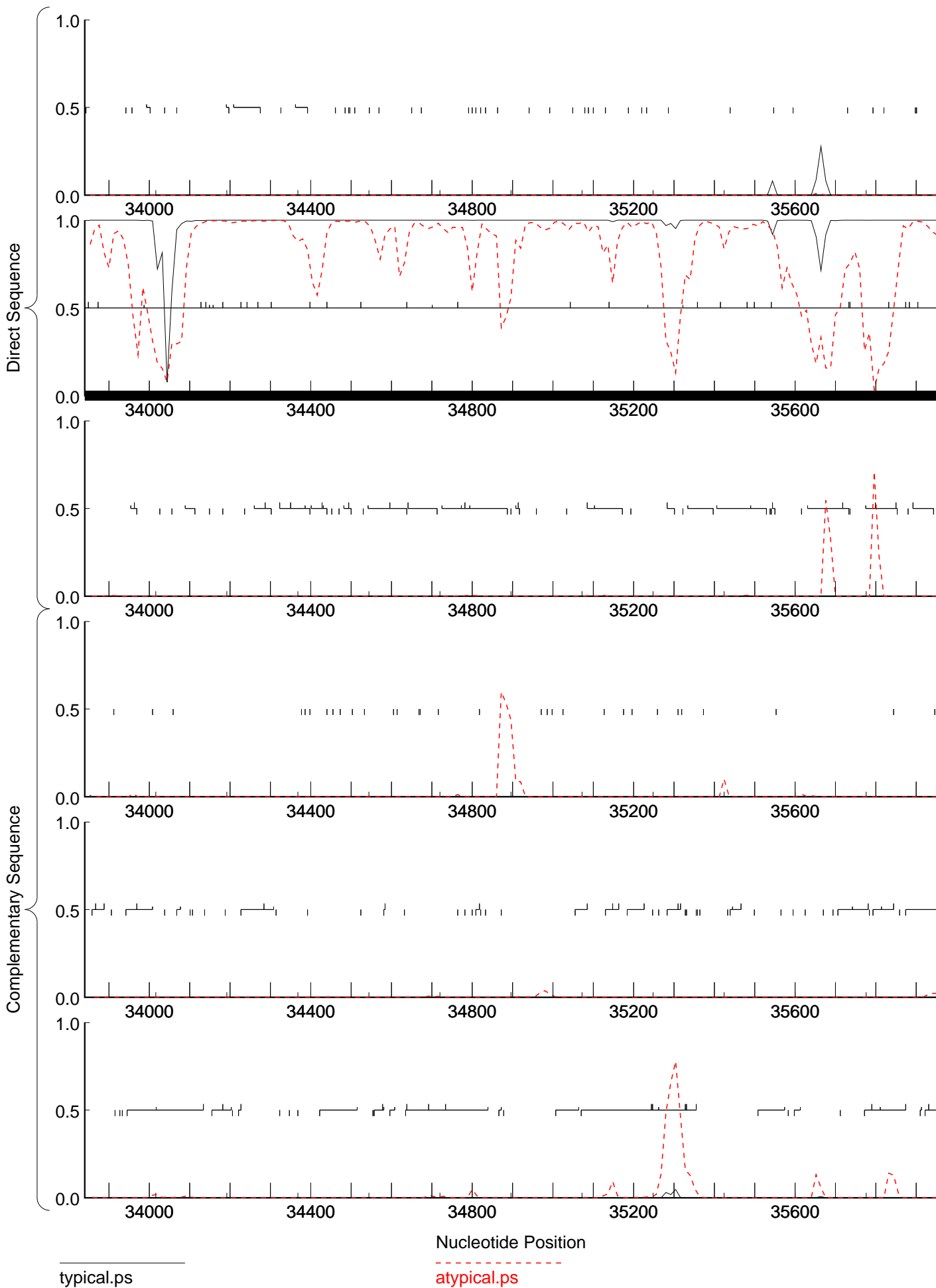


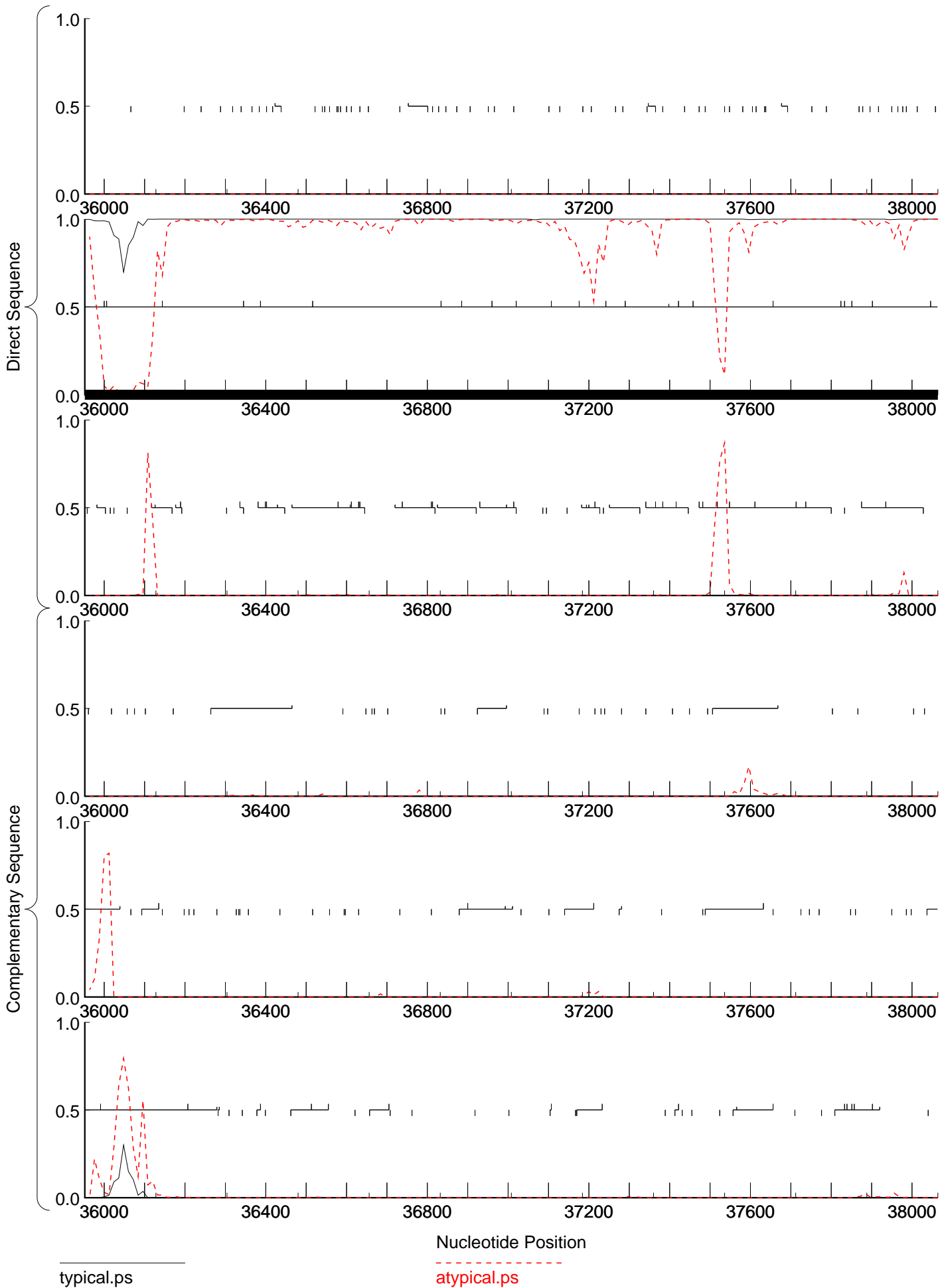
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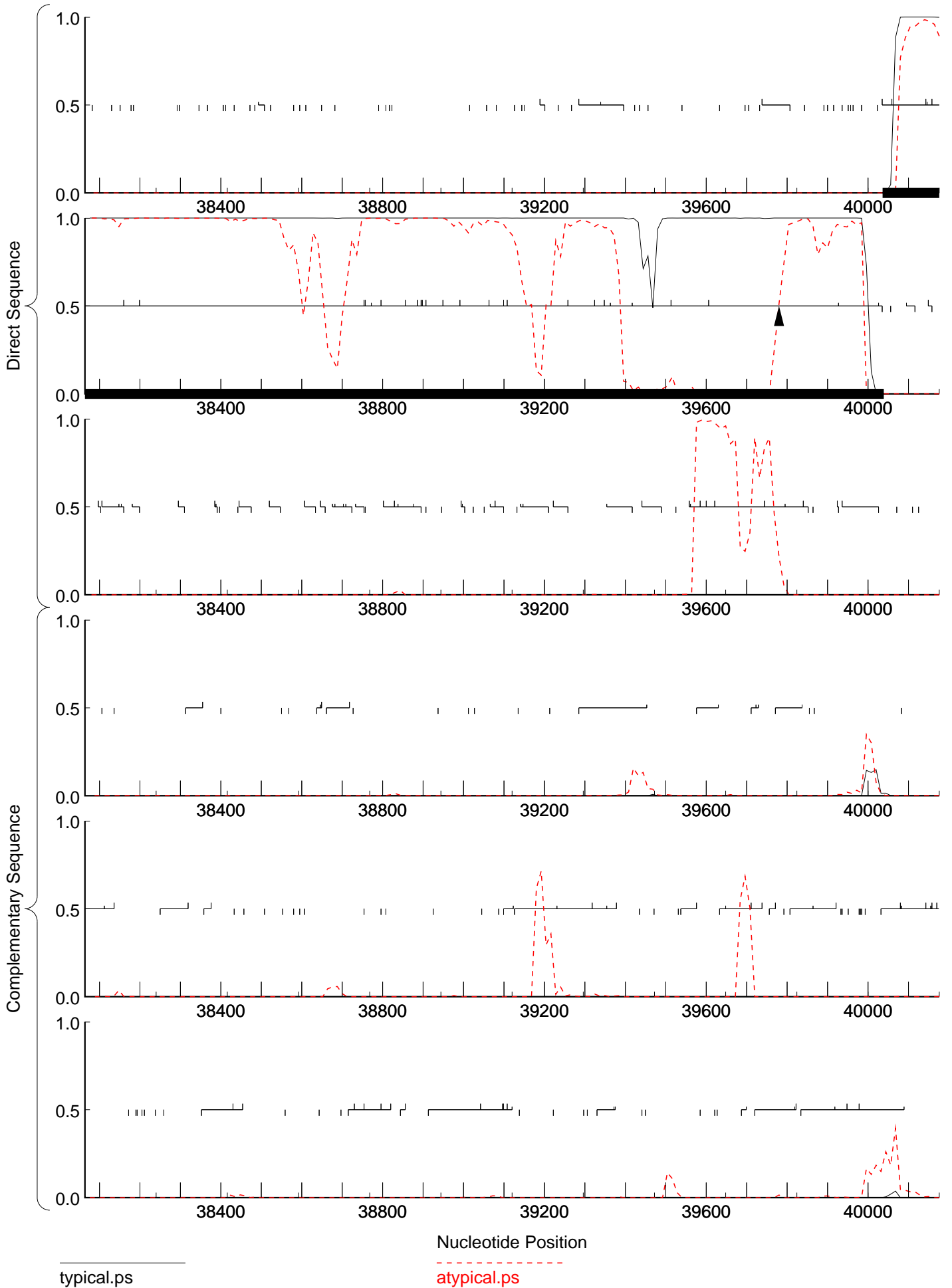
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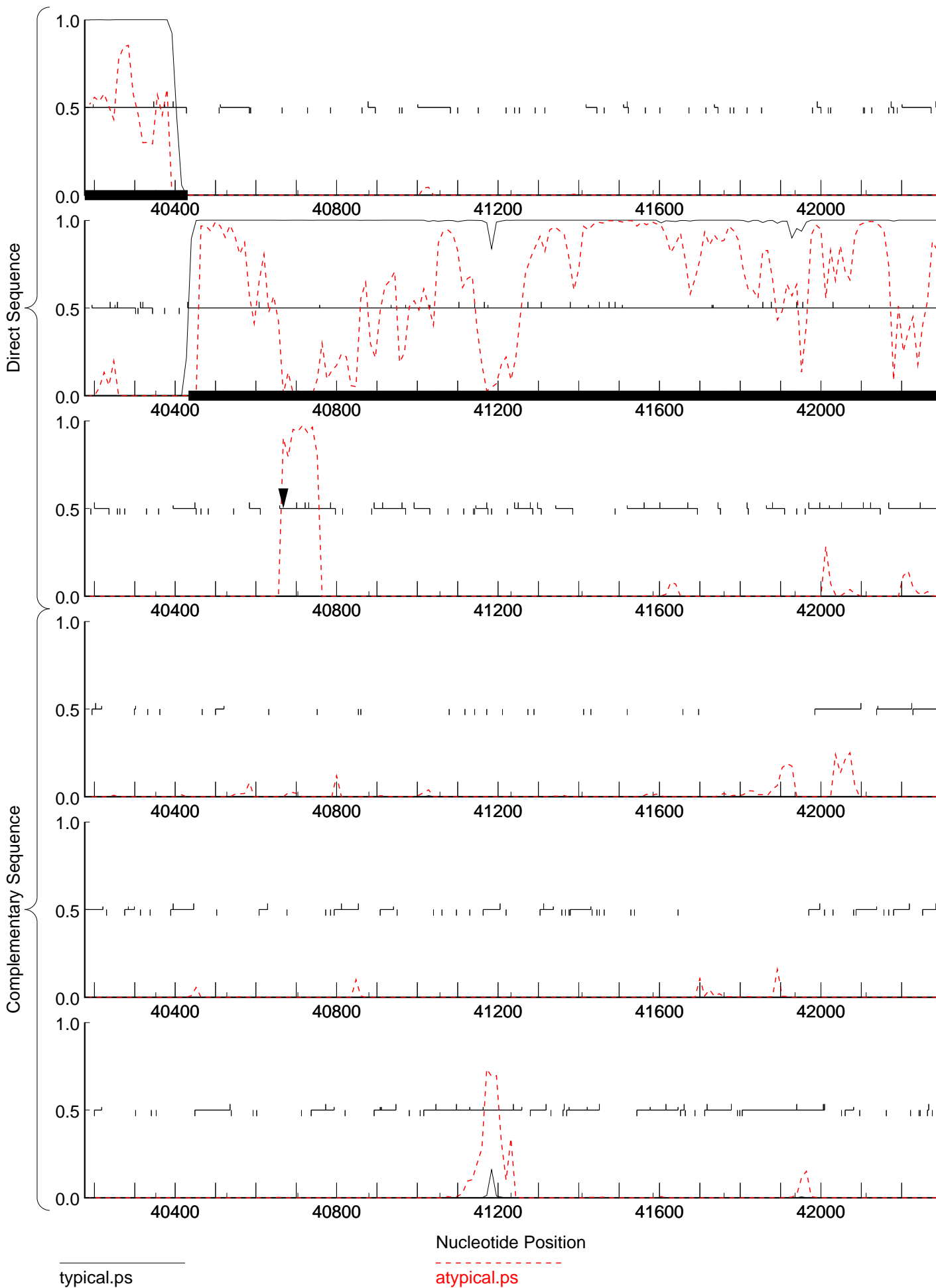


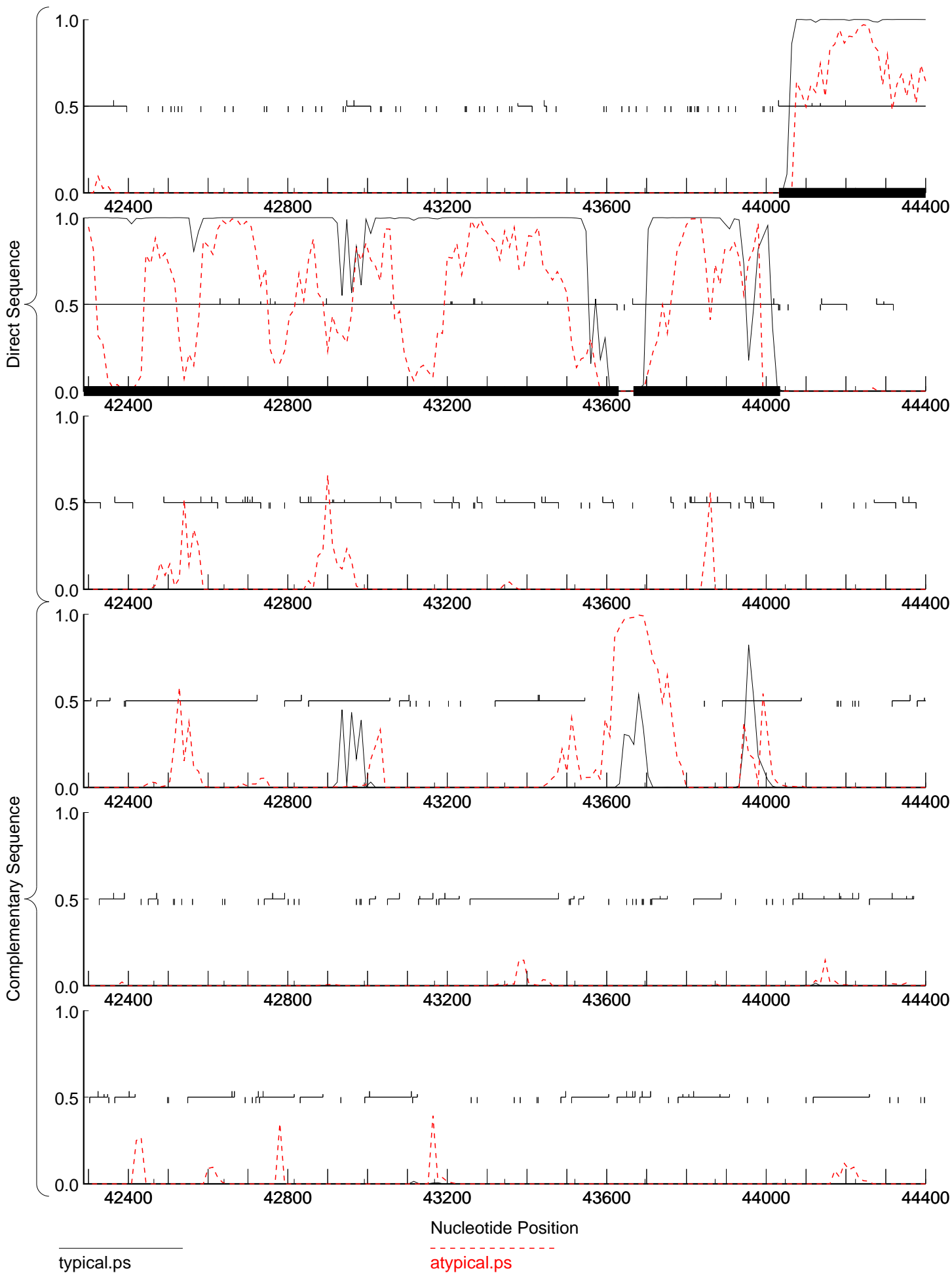


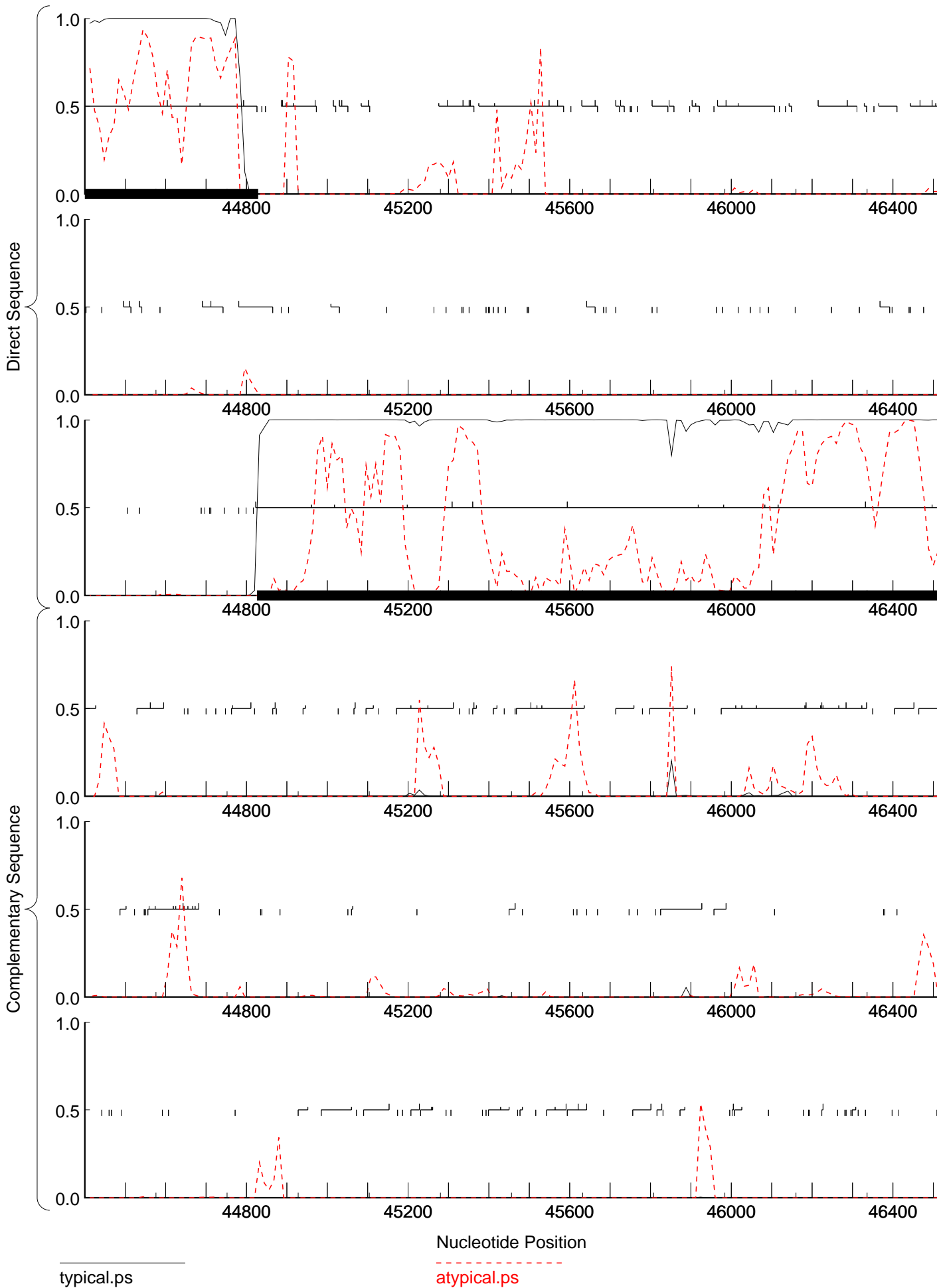






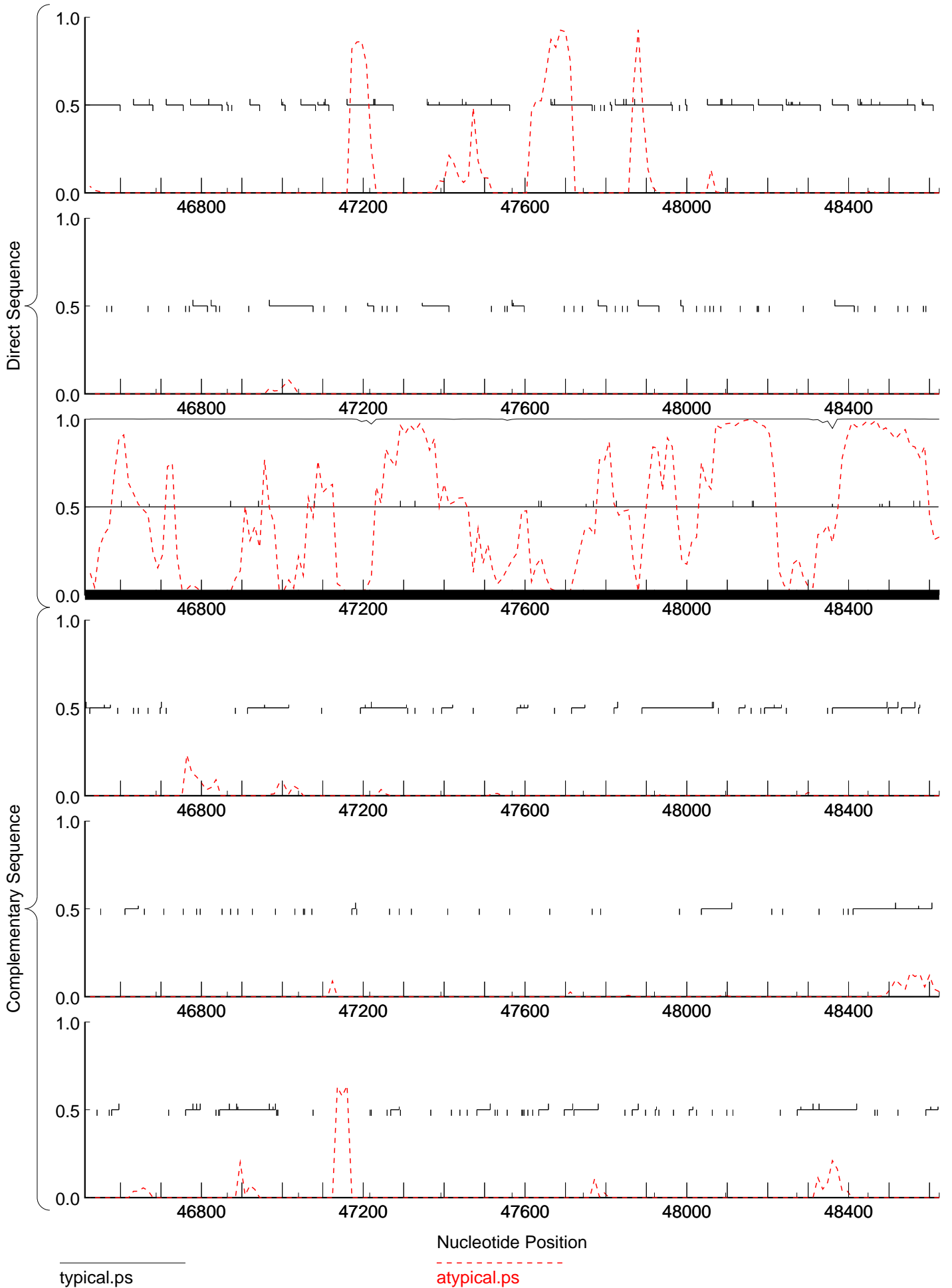


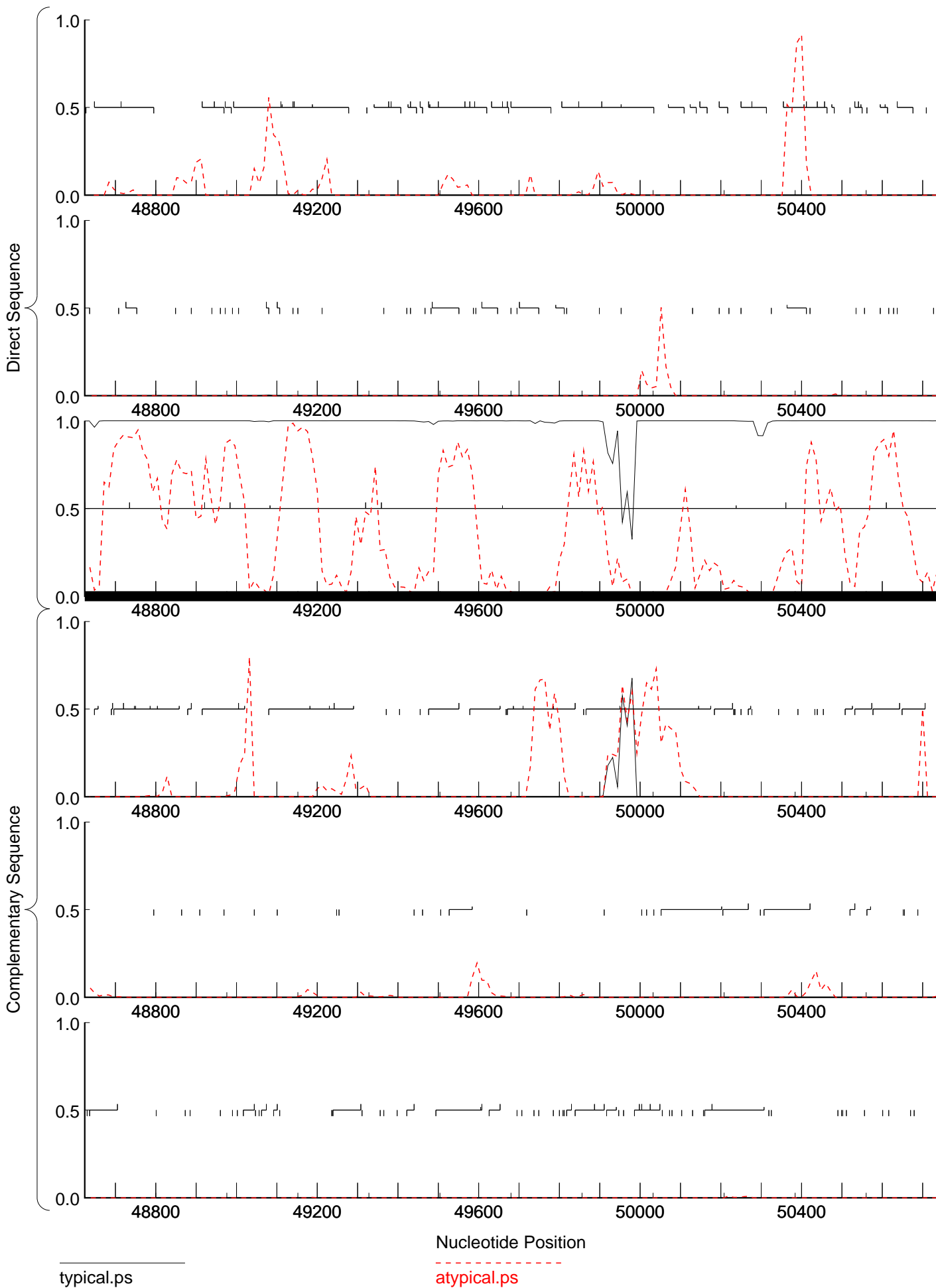




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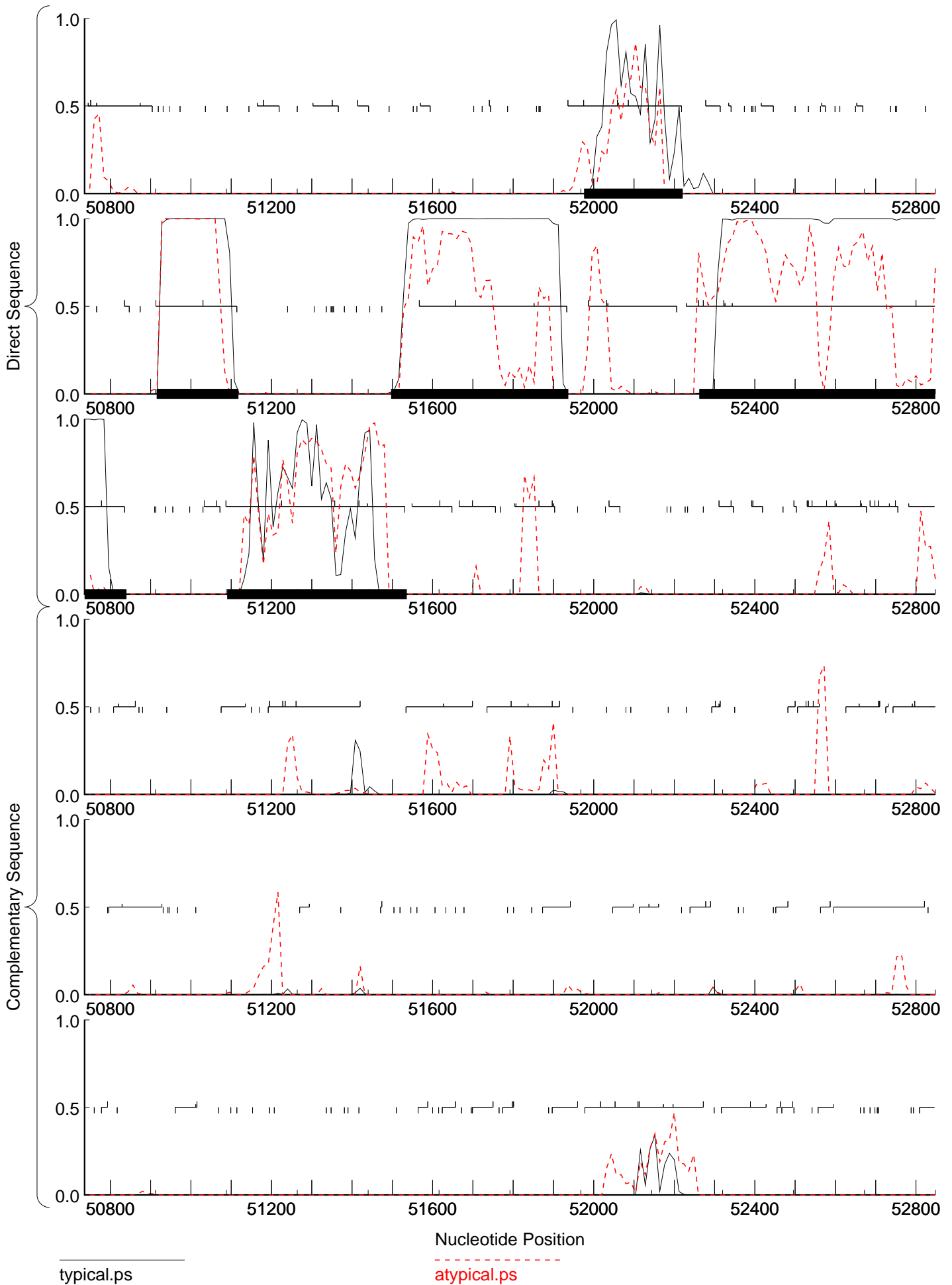
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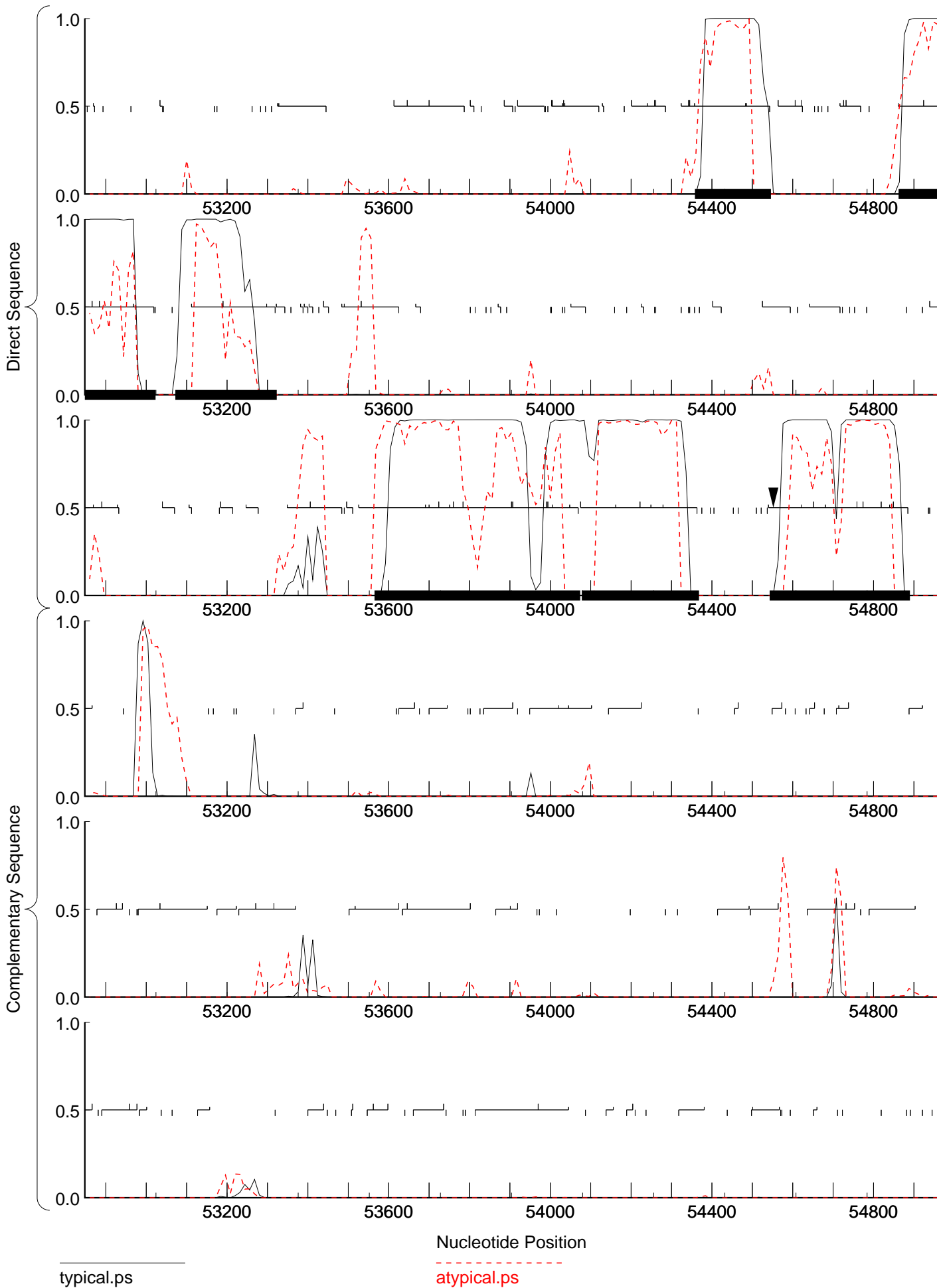


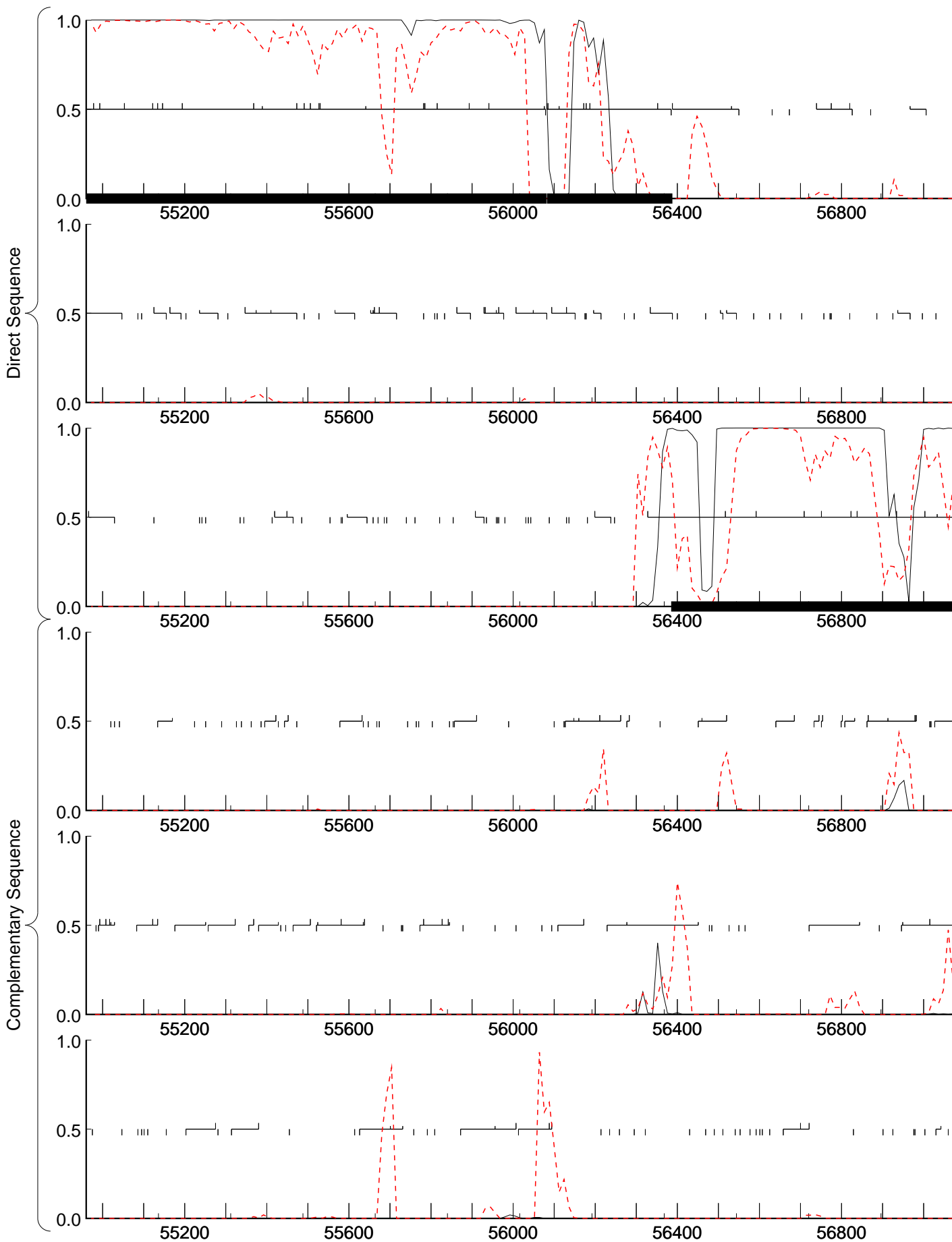


typical.ps

atypical.ps

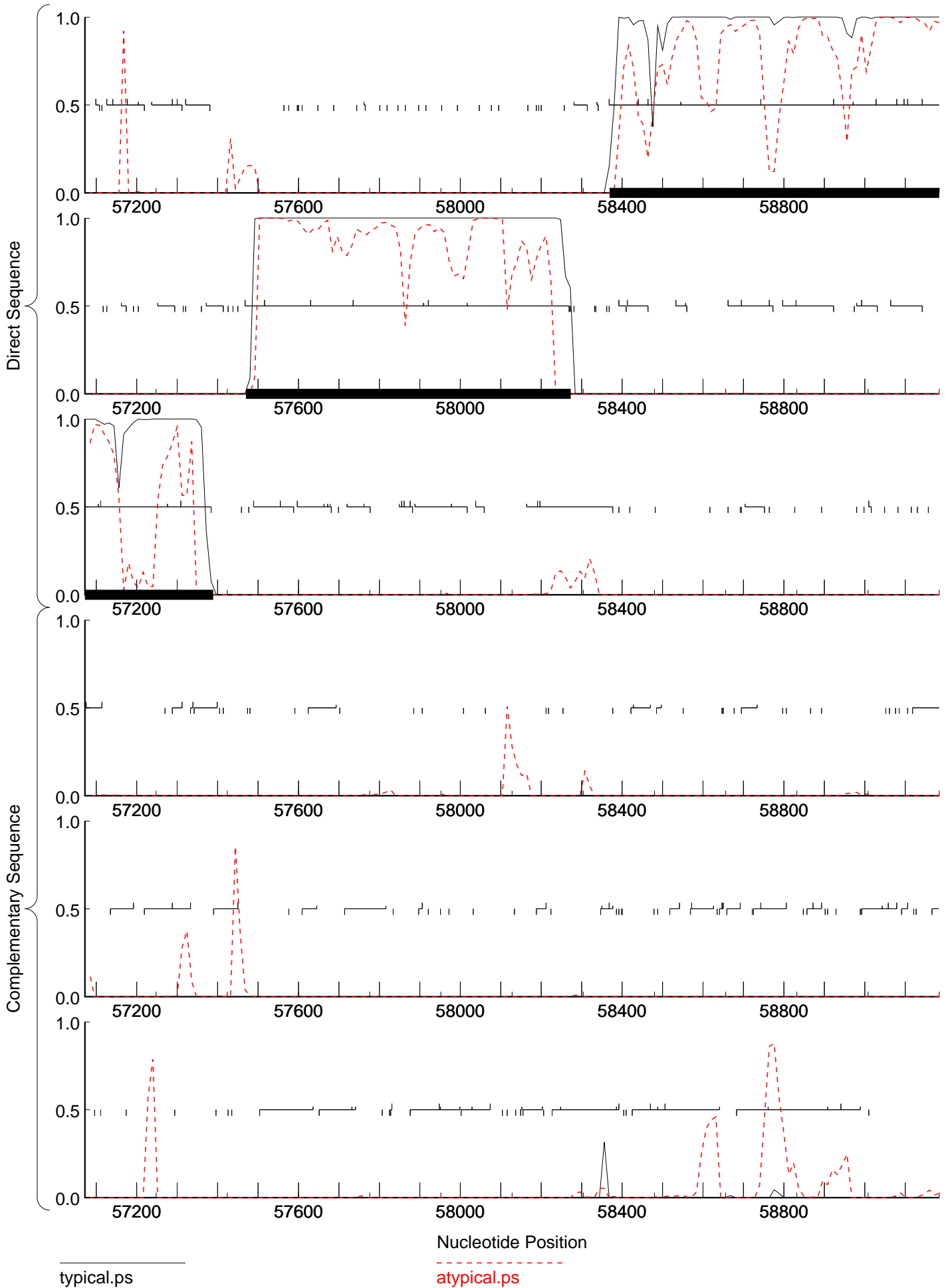


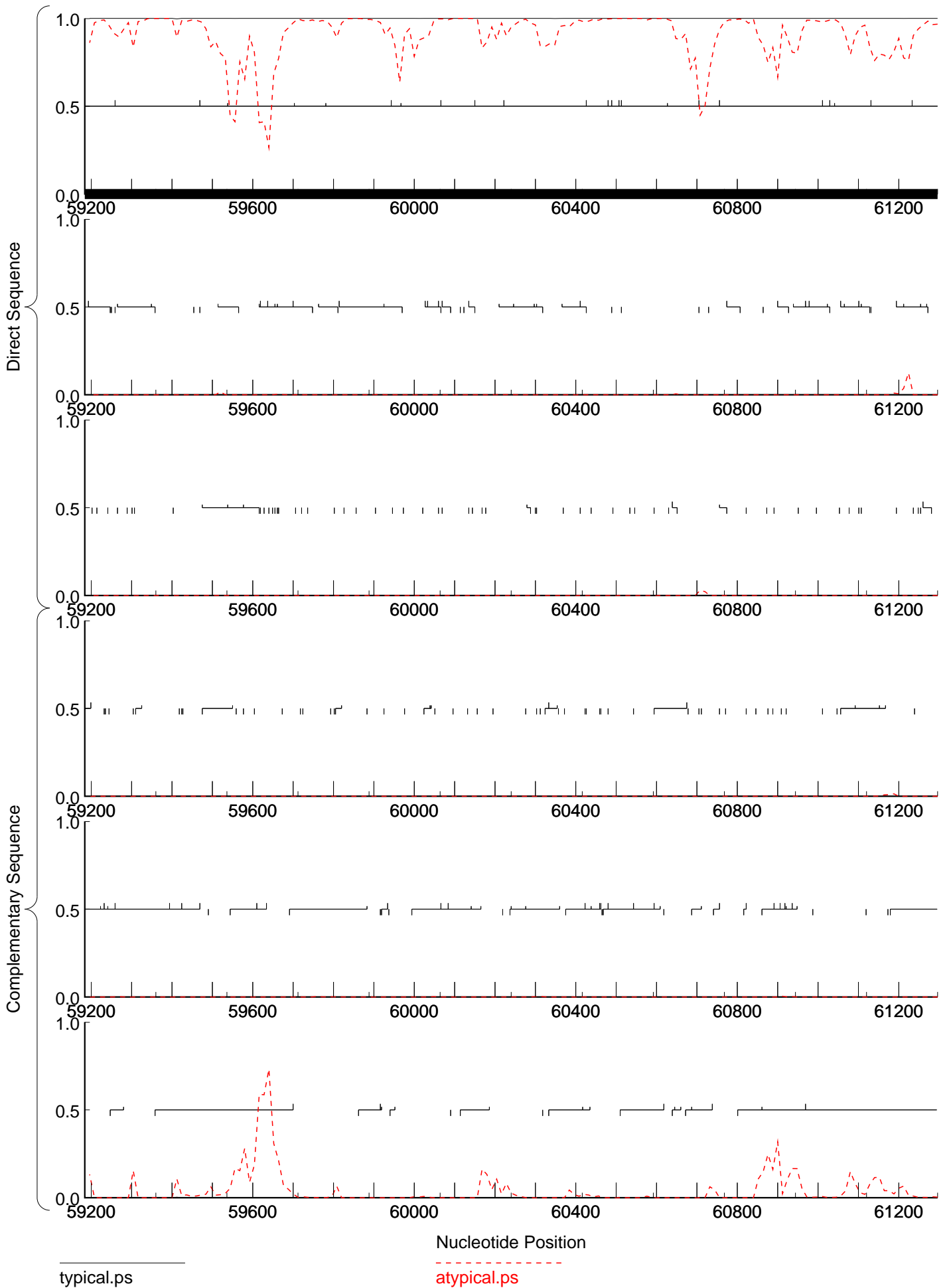


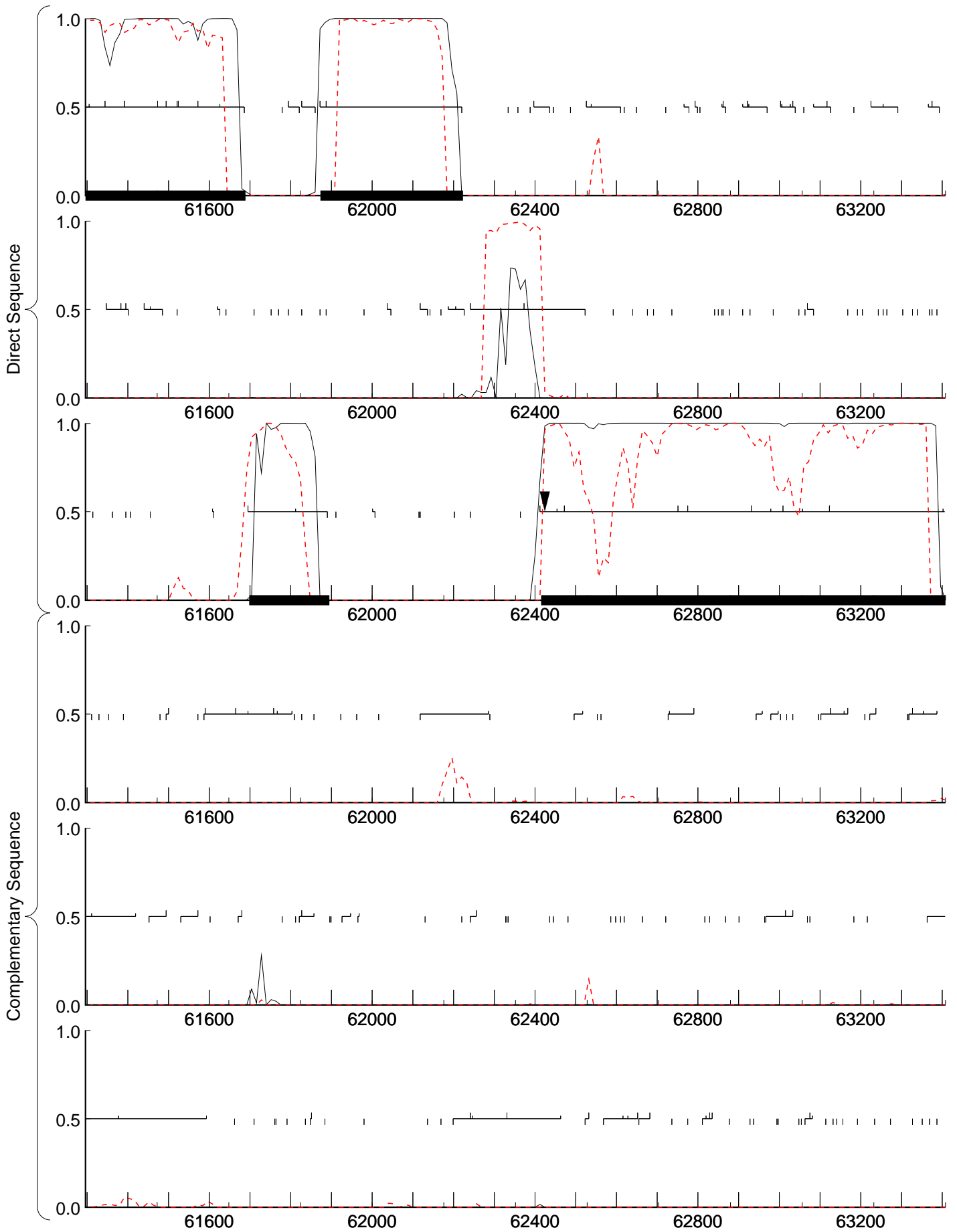


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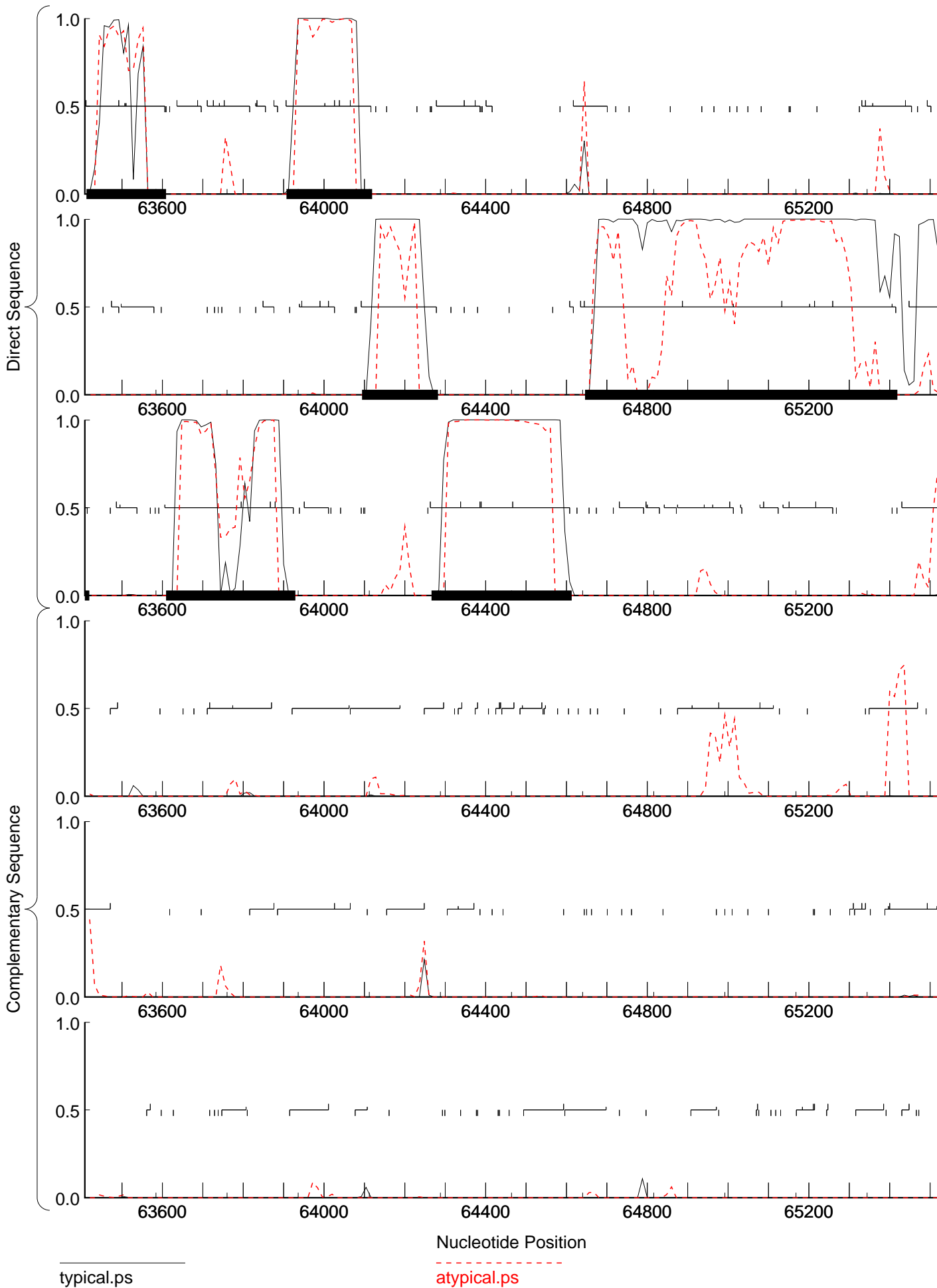






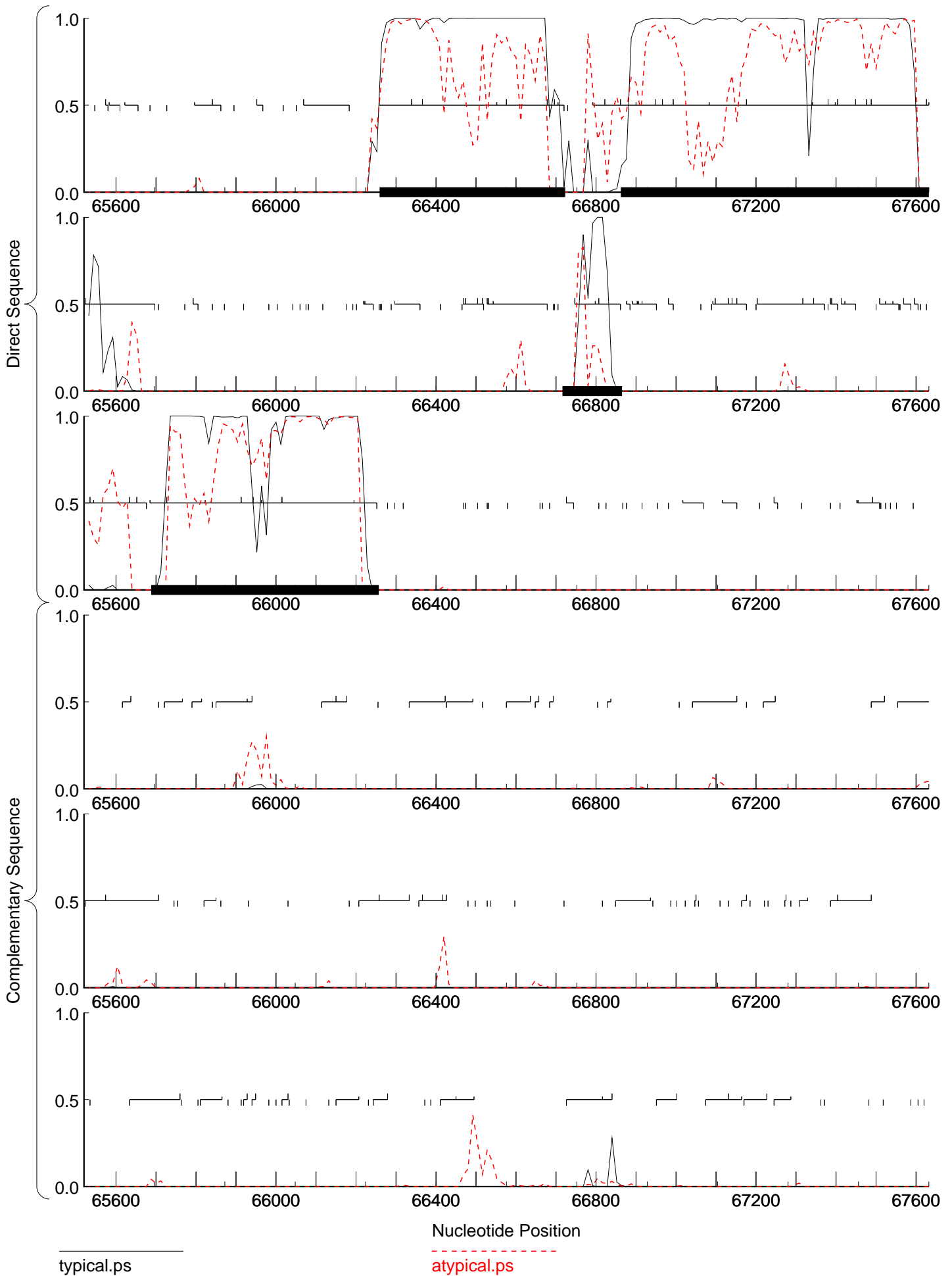
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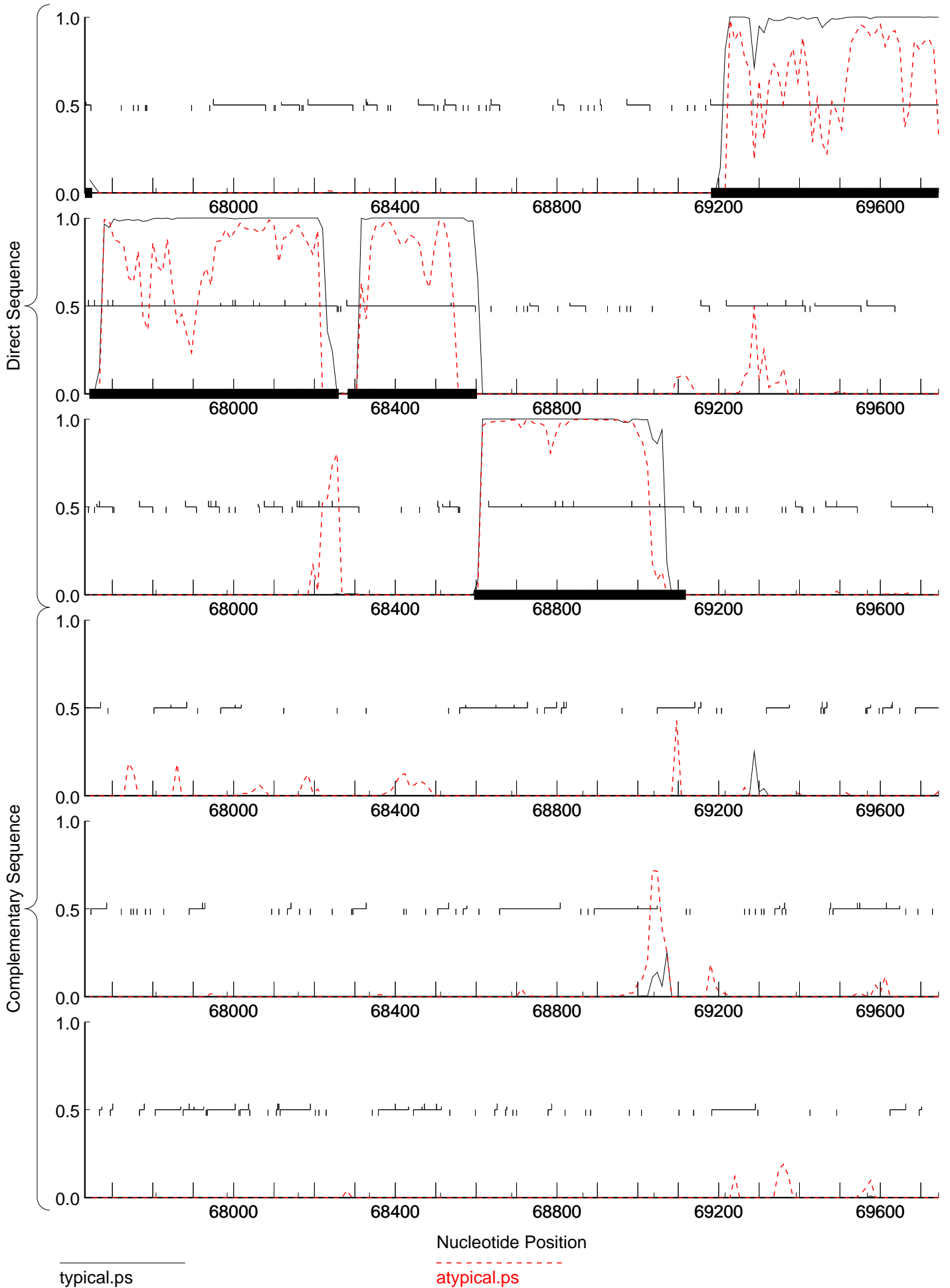
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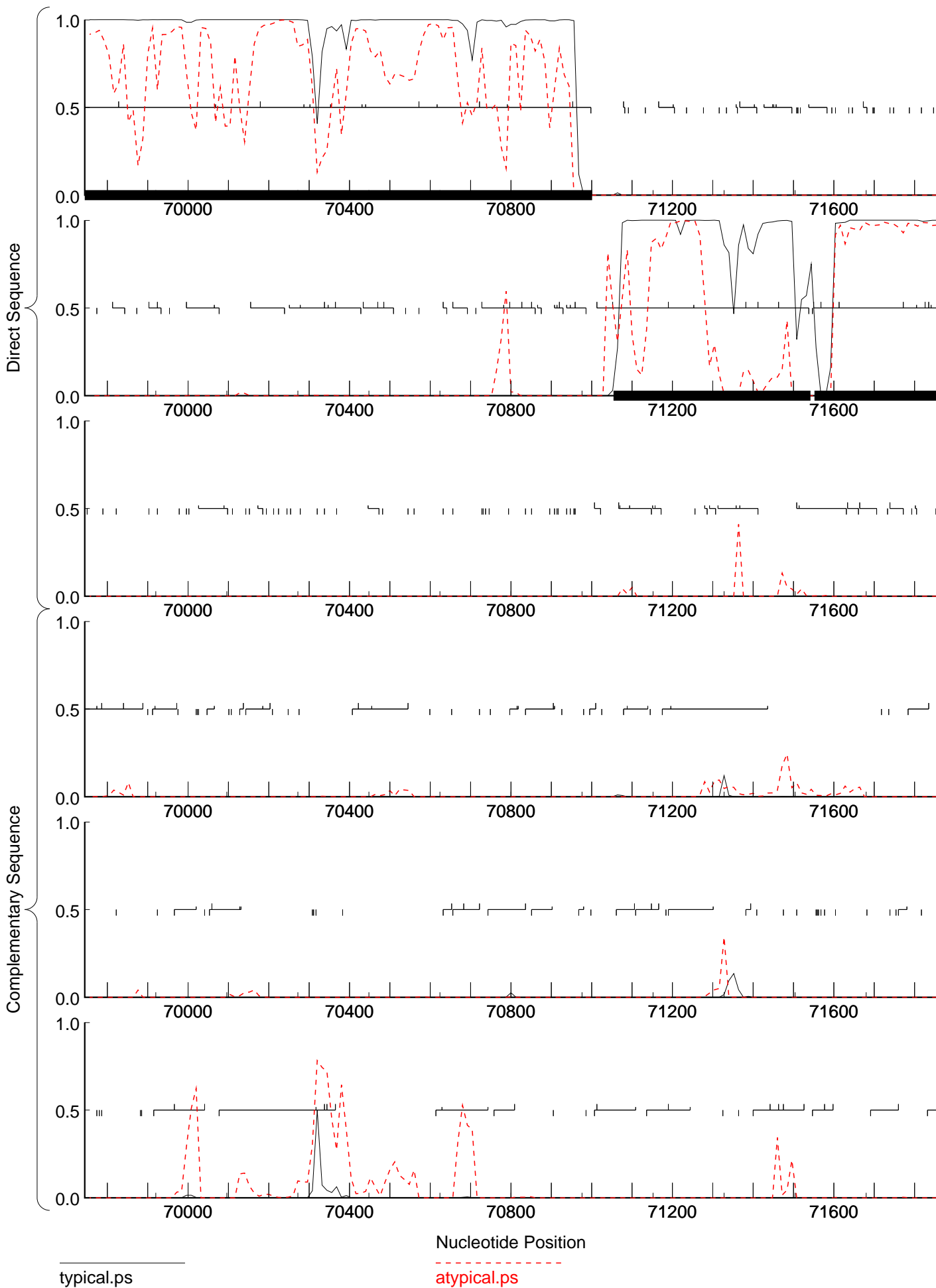
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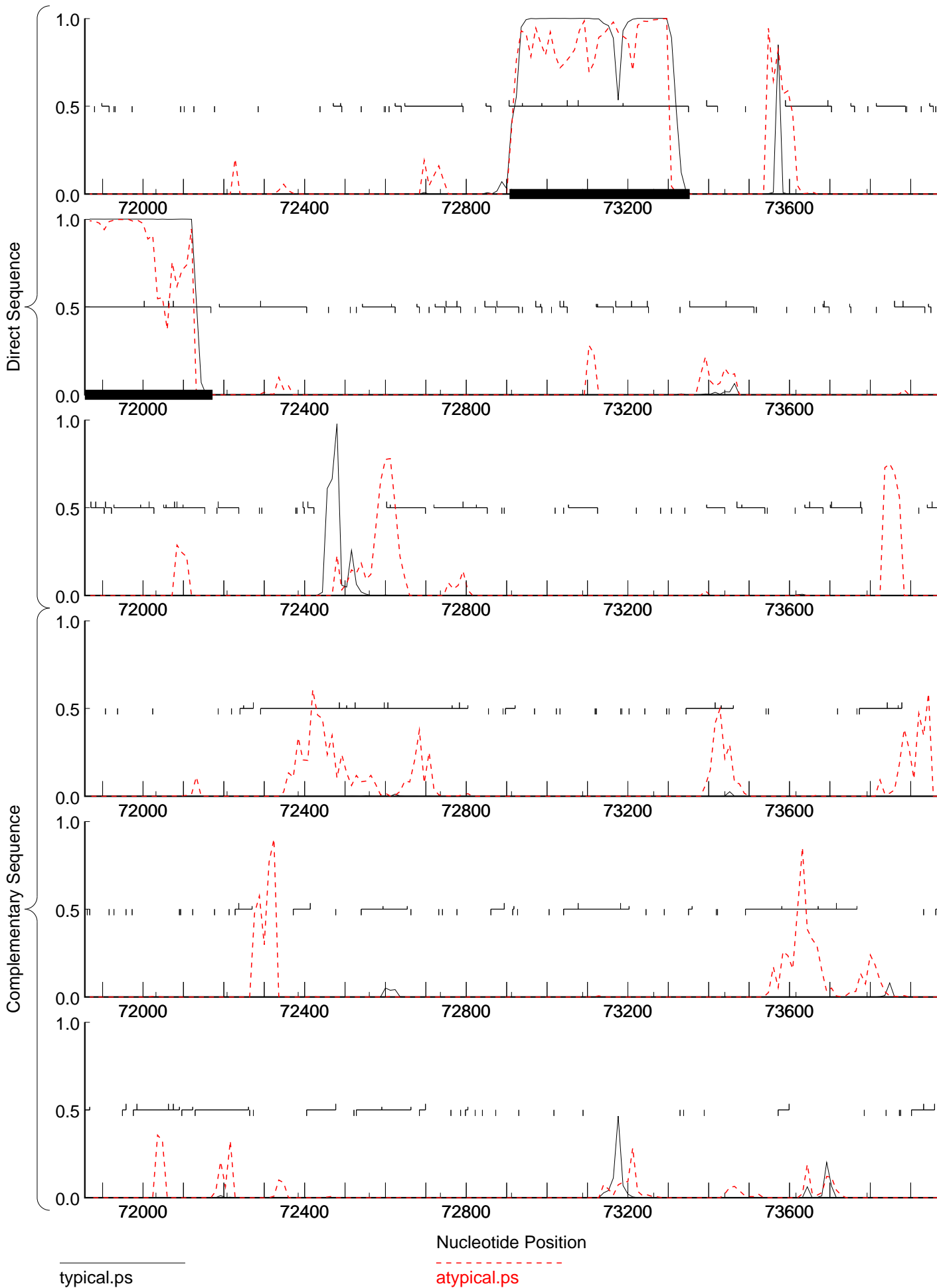
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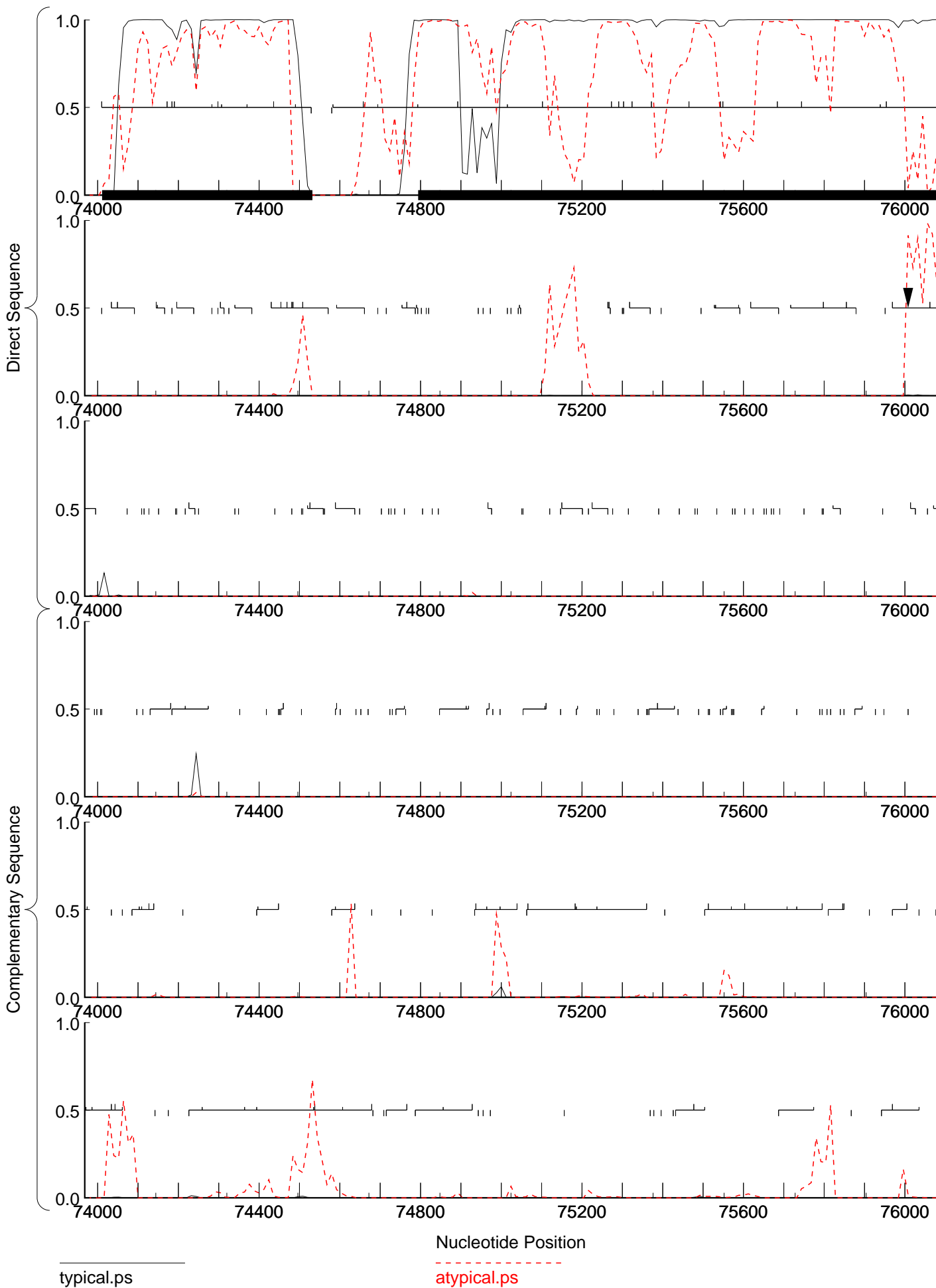
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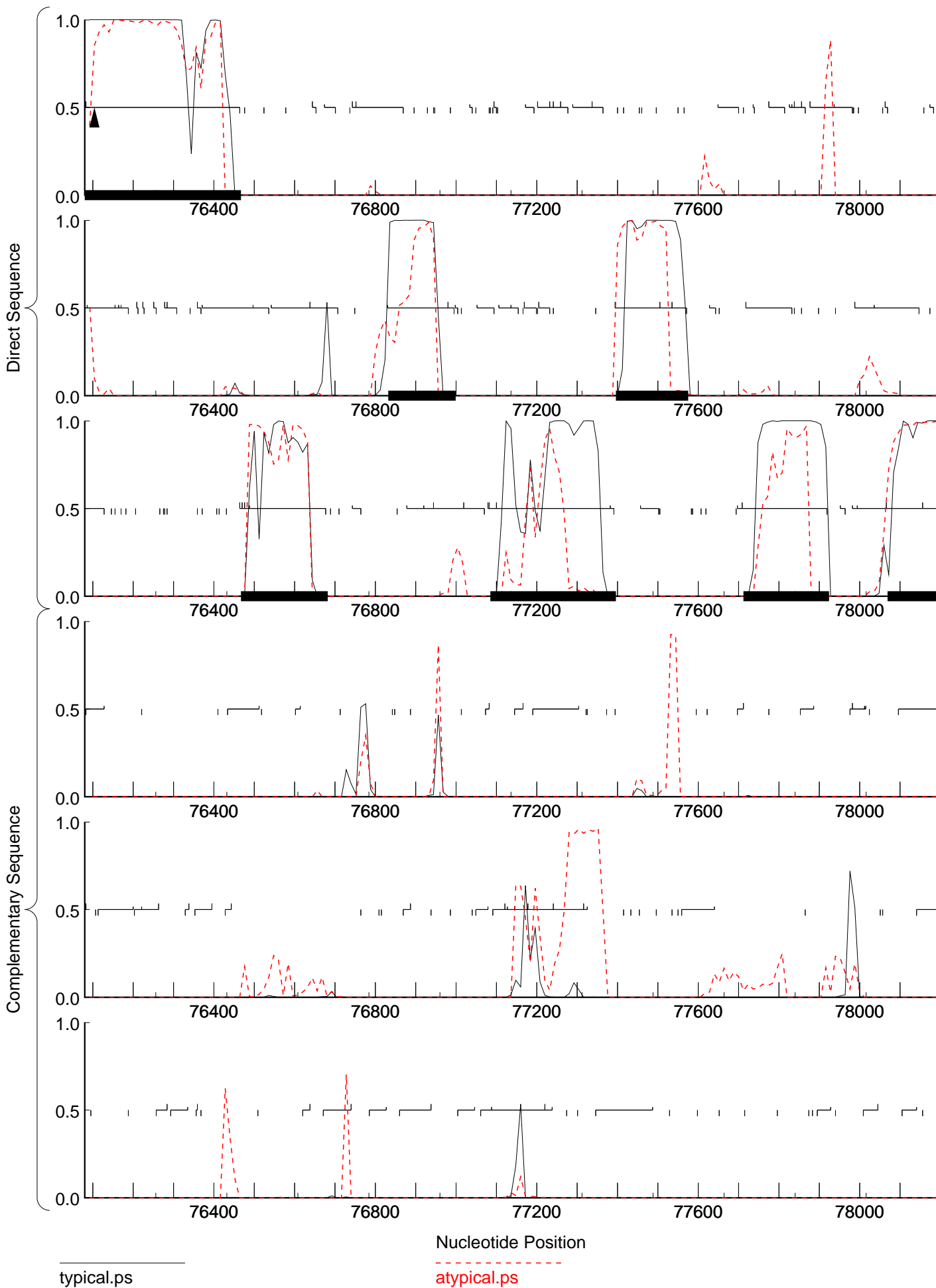
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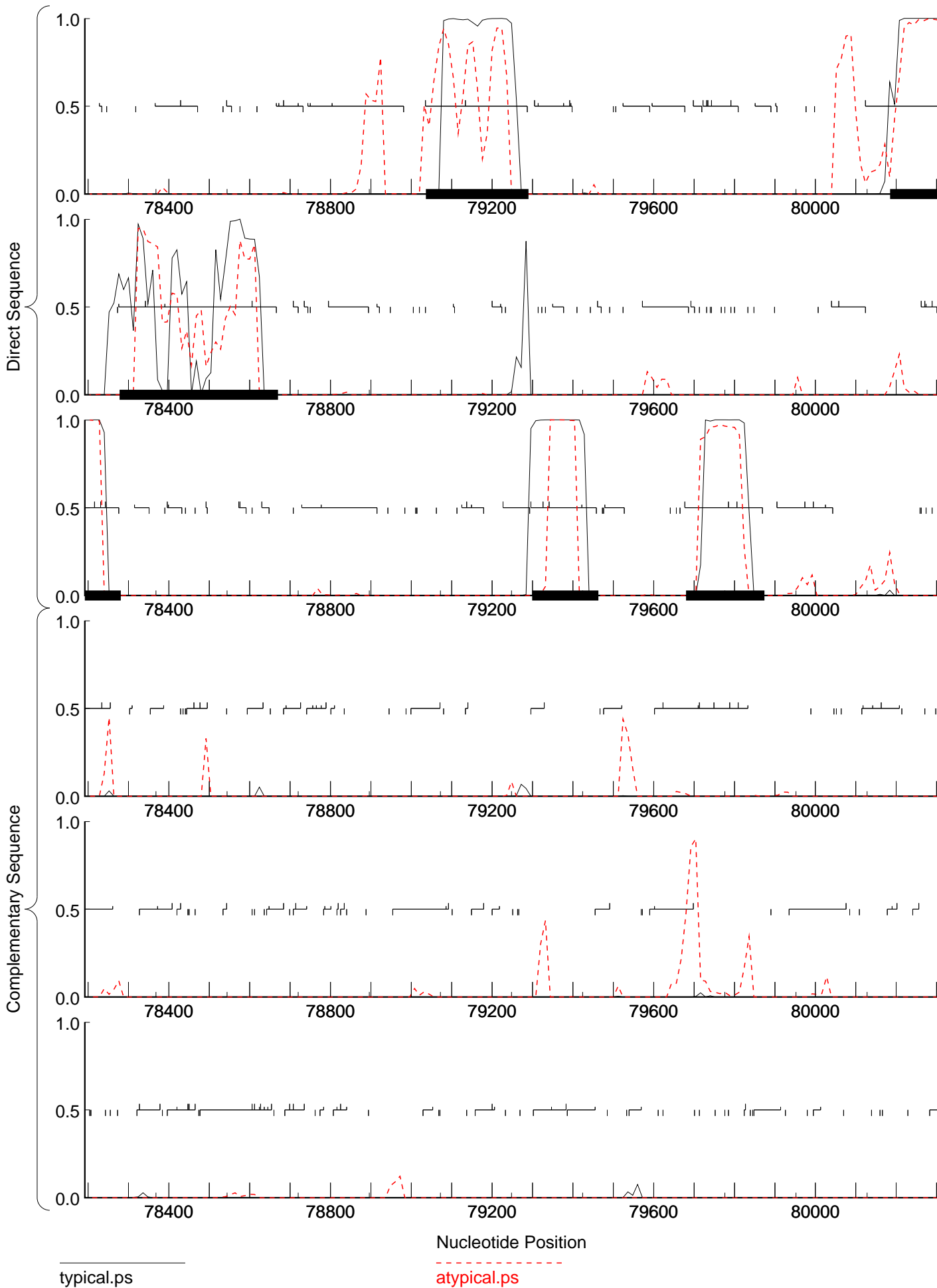
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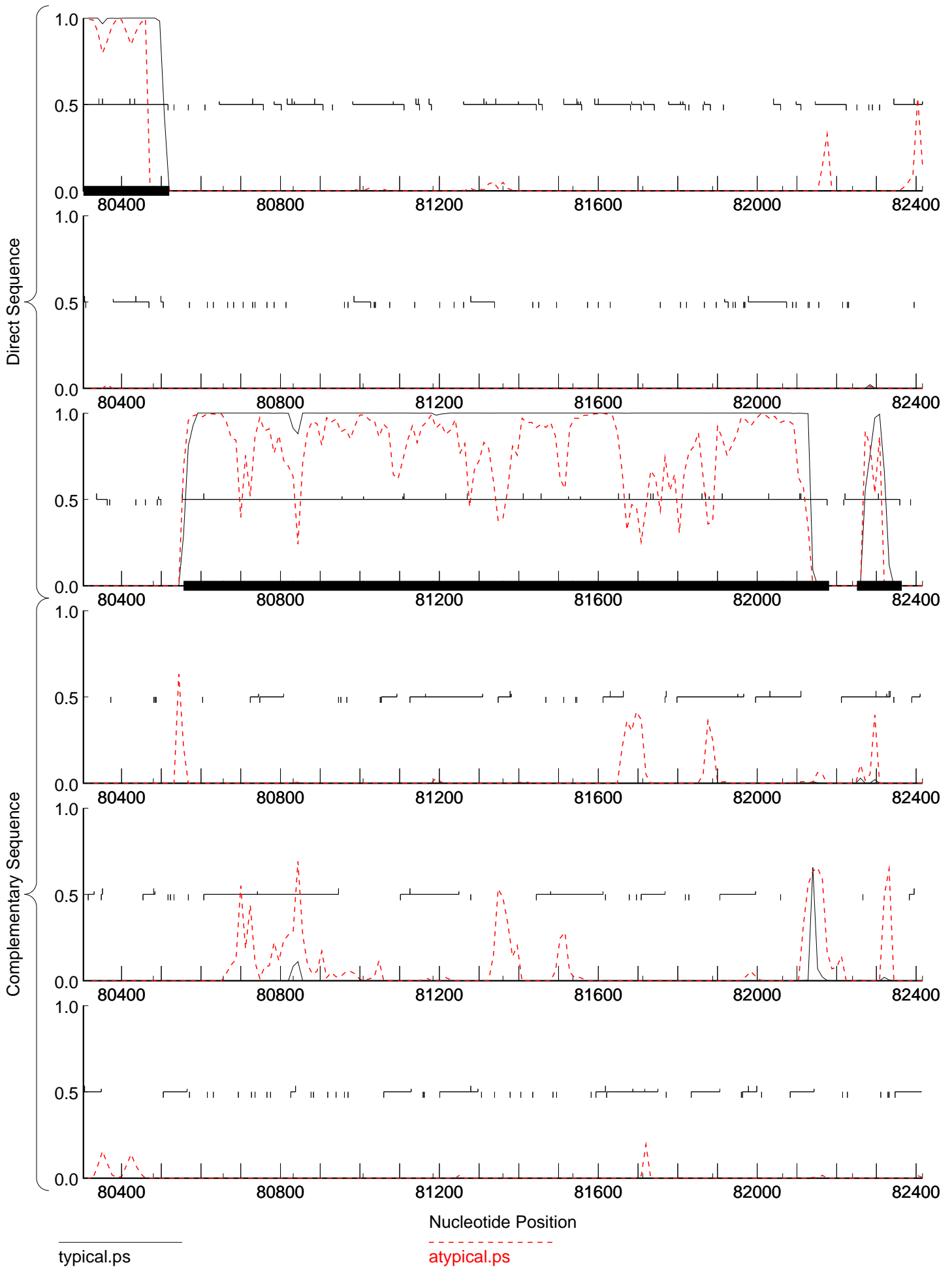


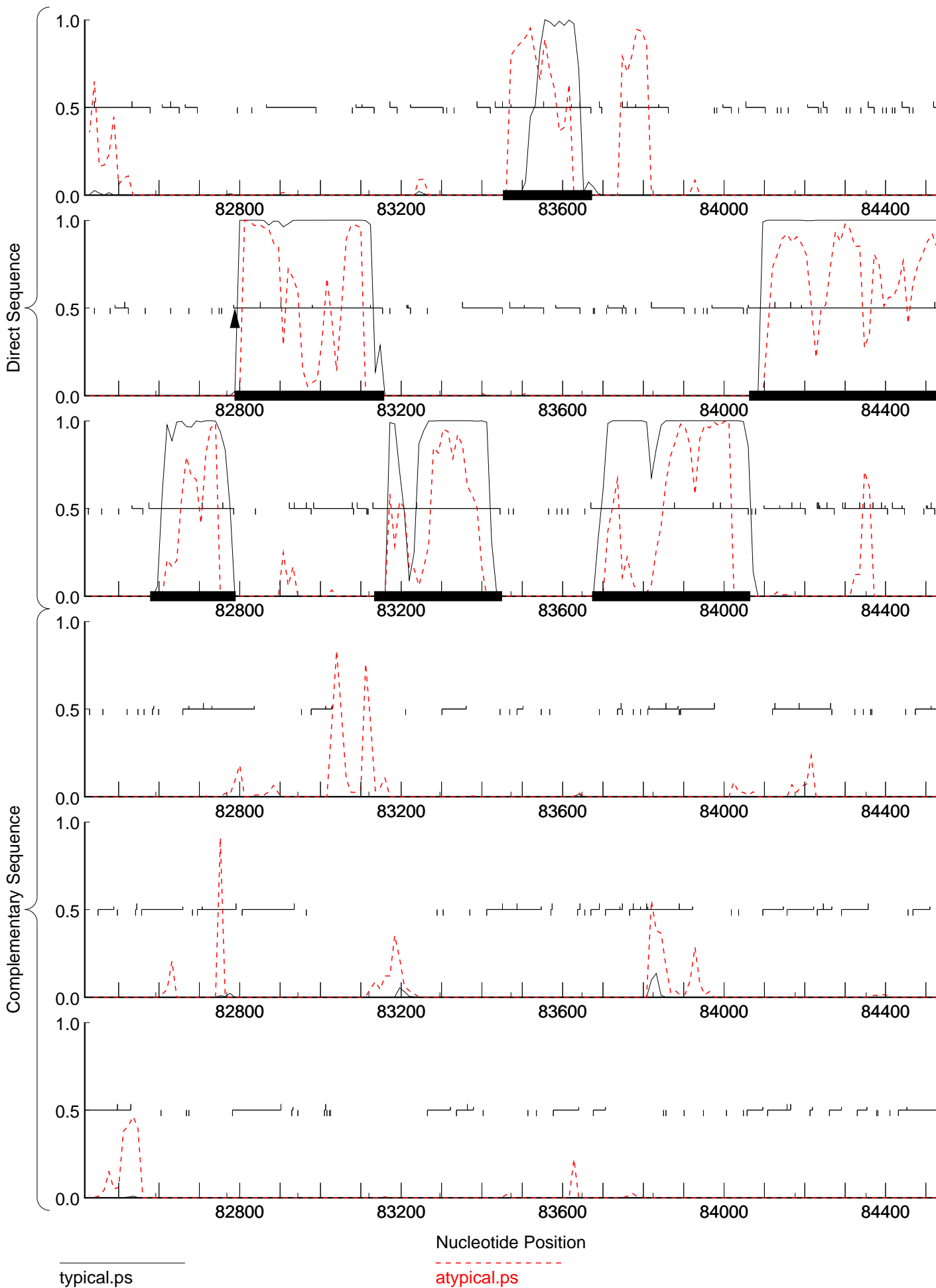
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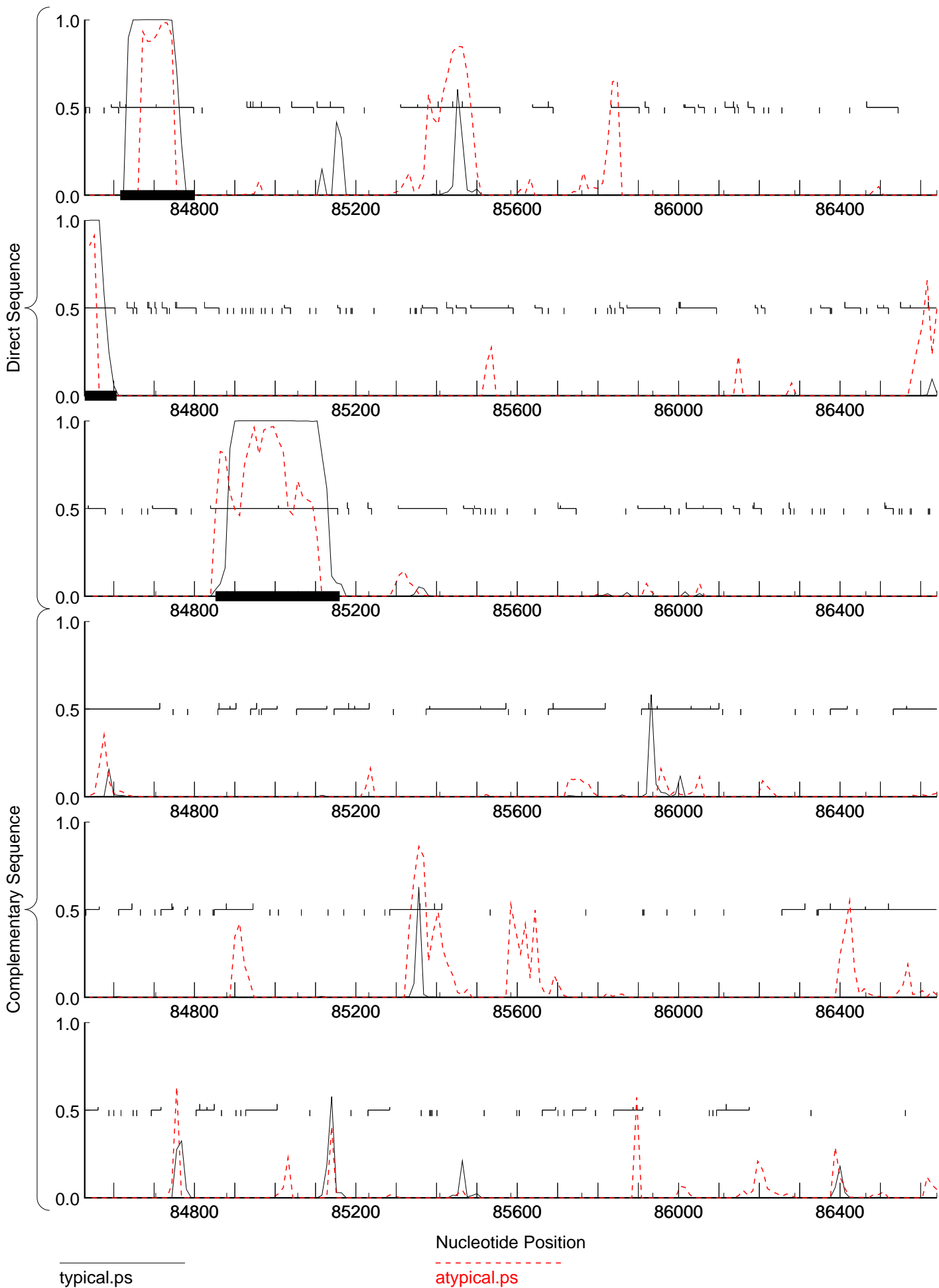
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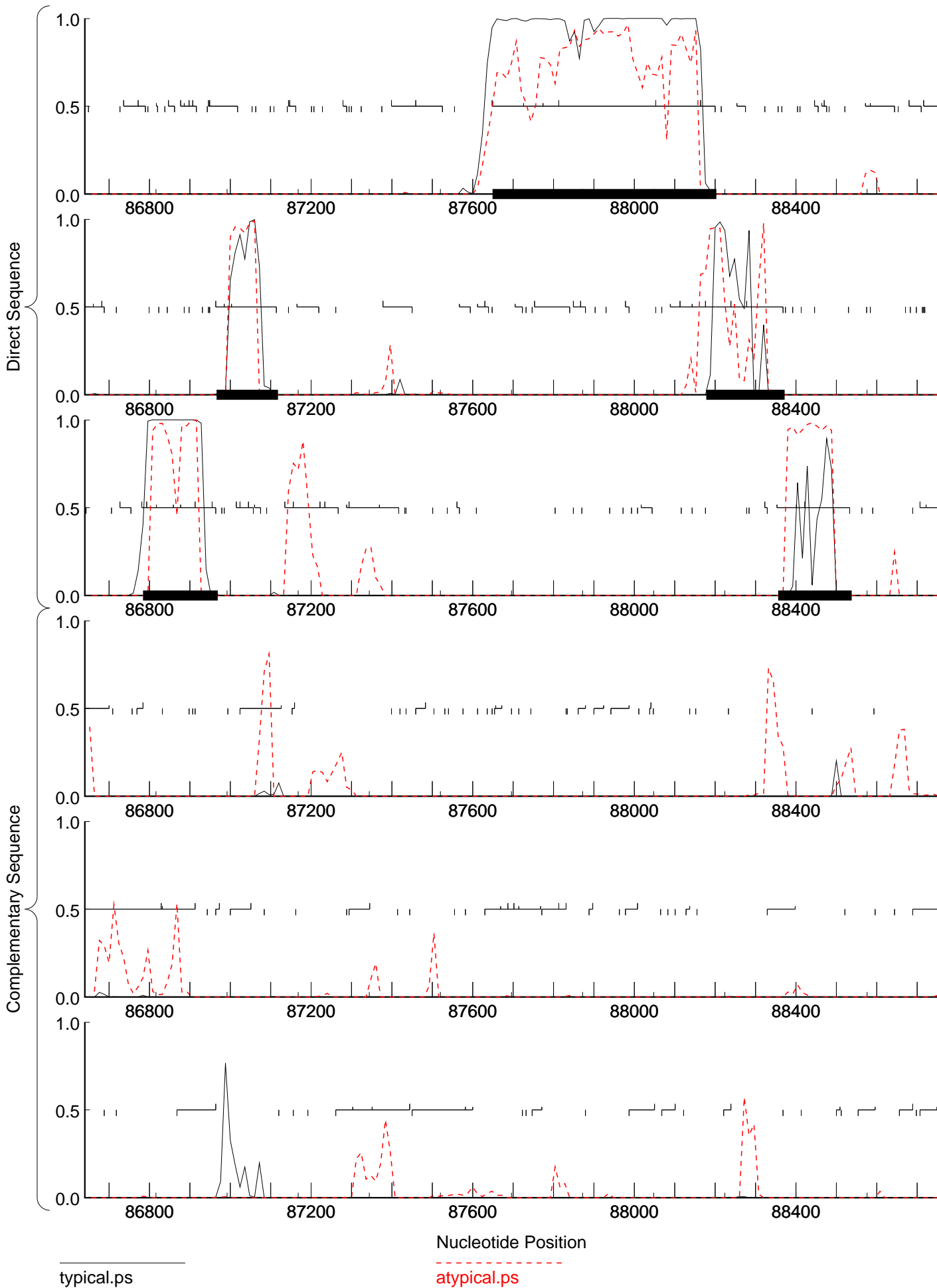


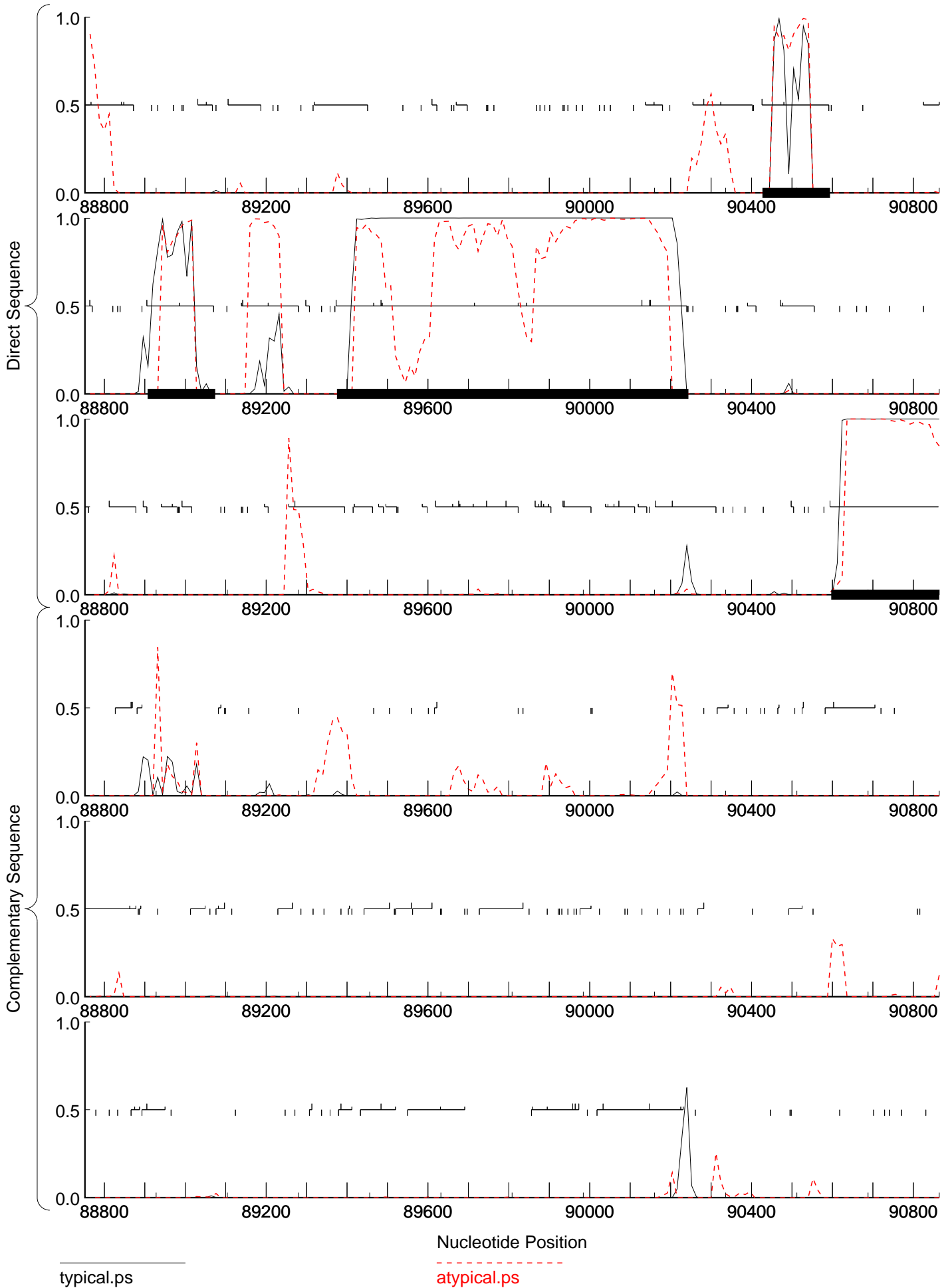


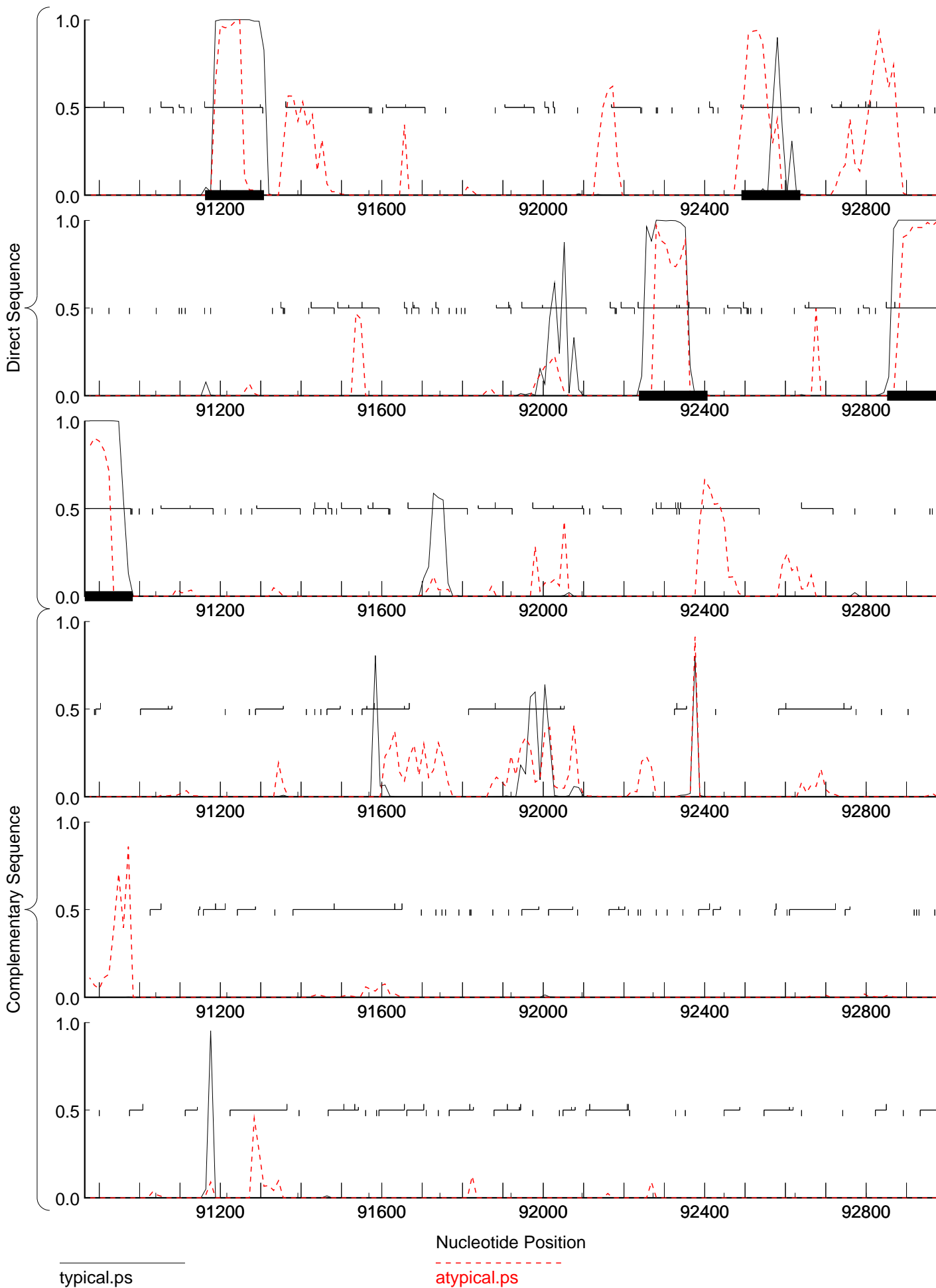


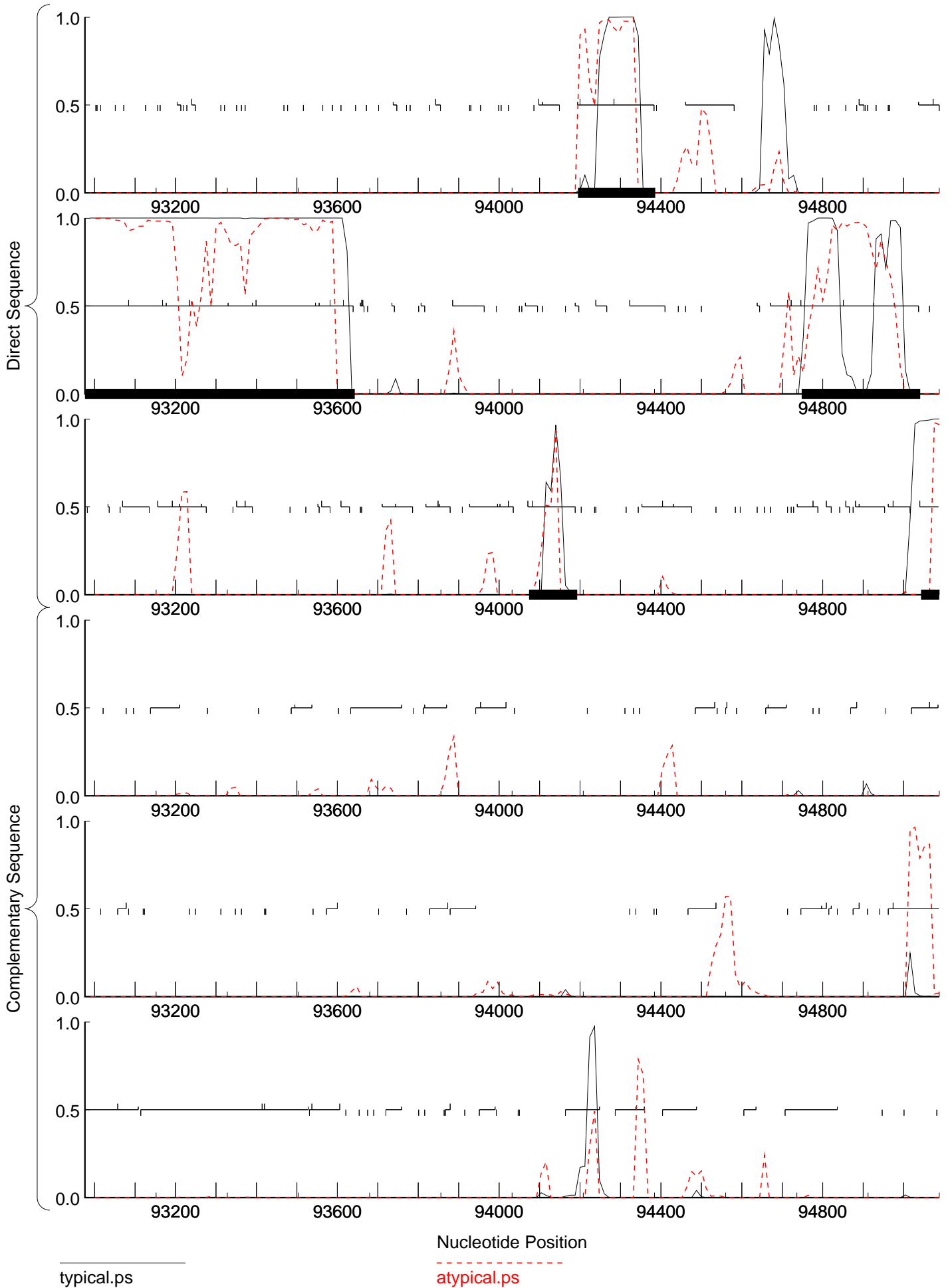






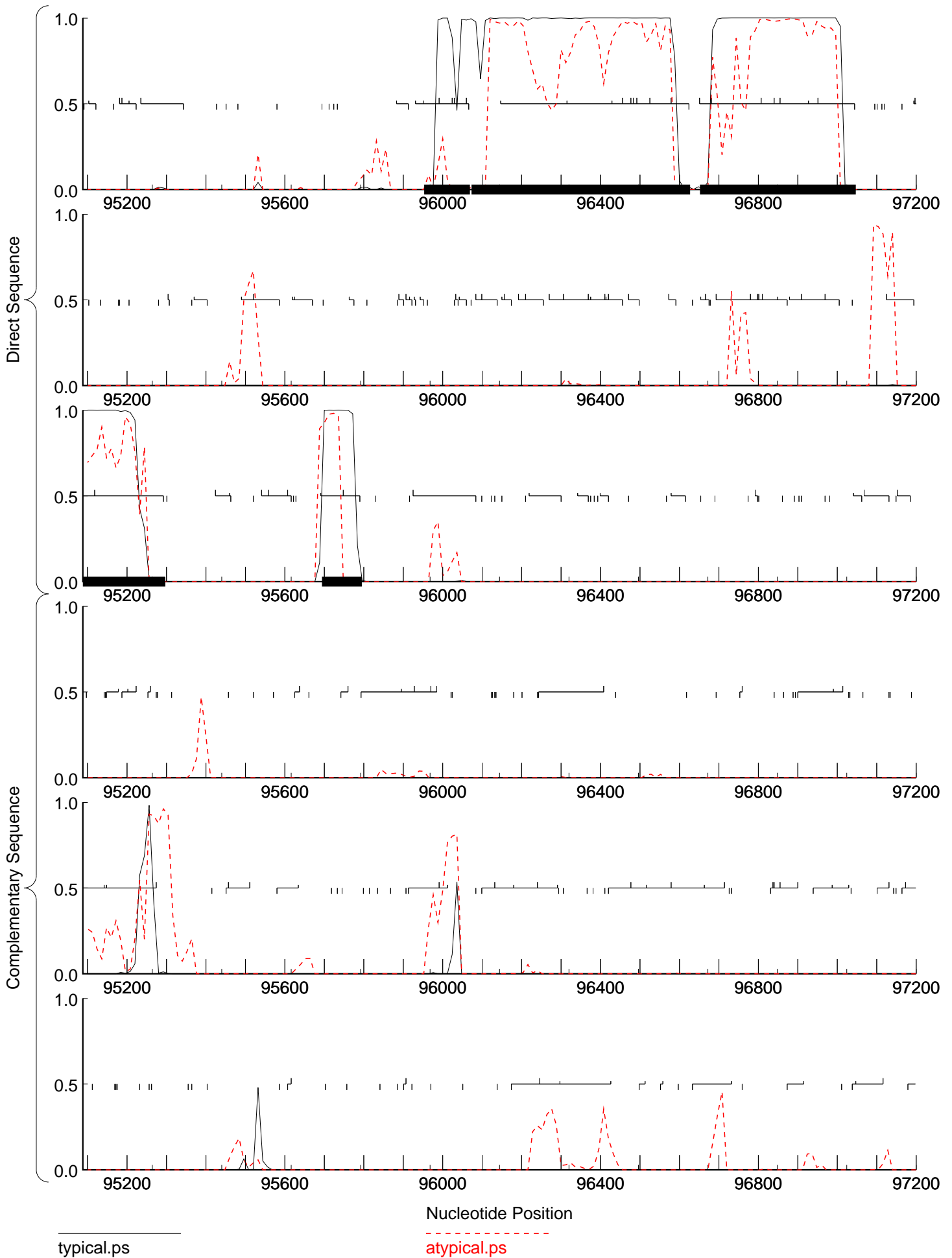


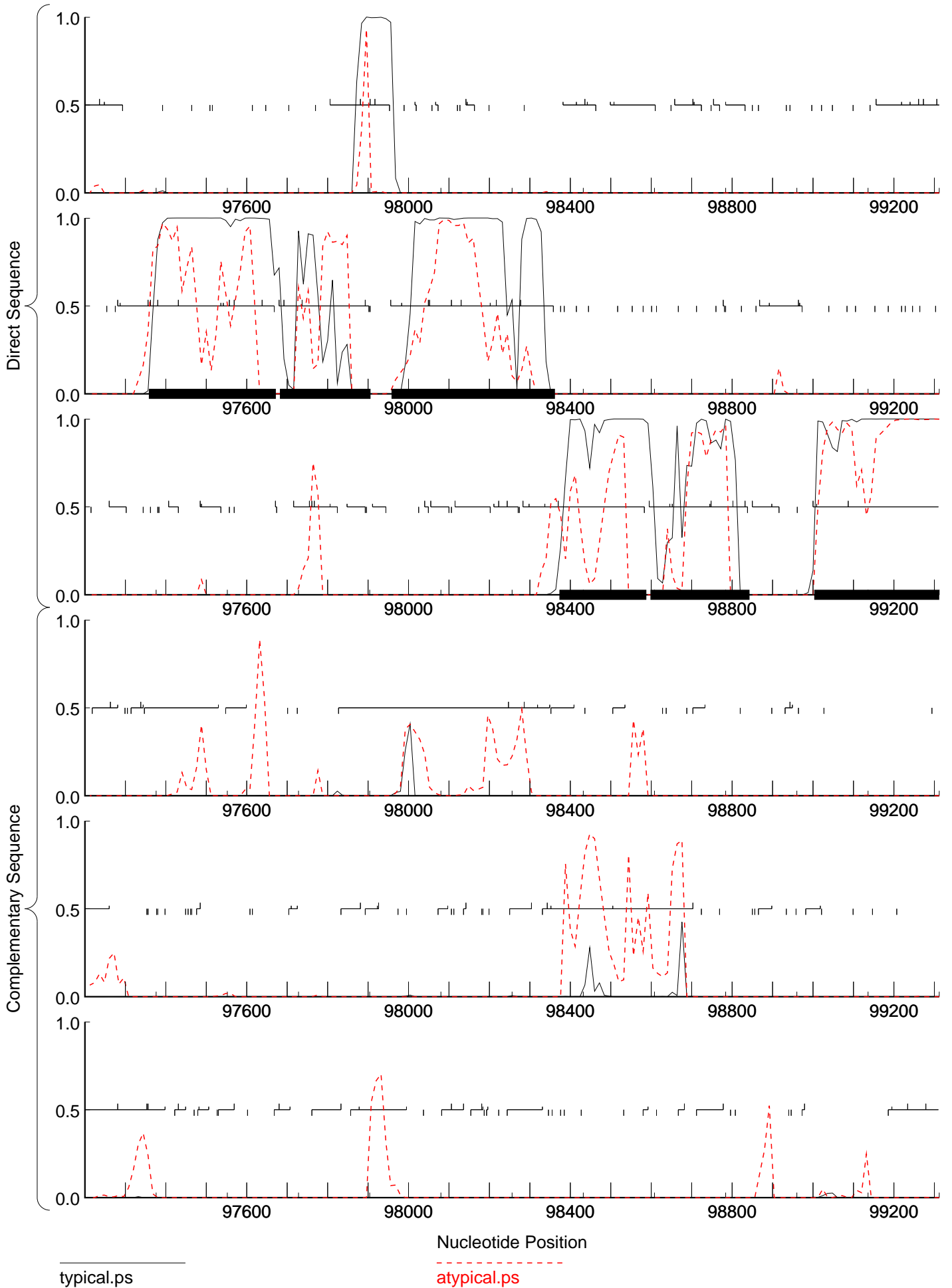




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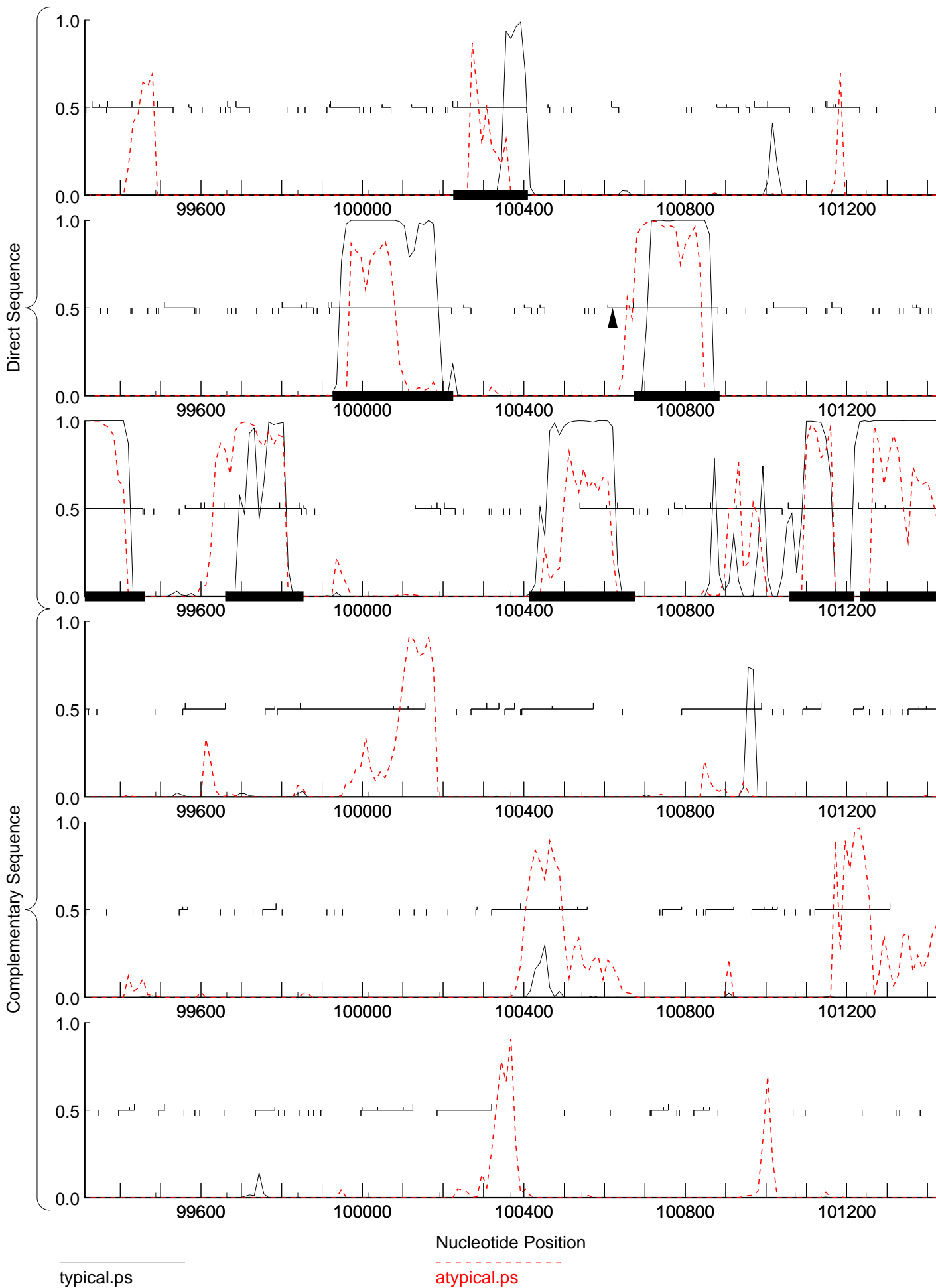
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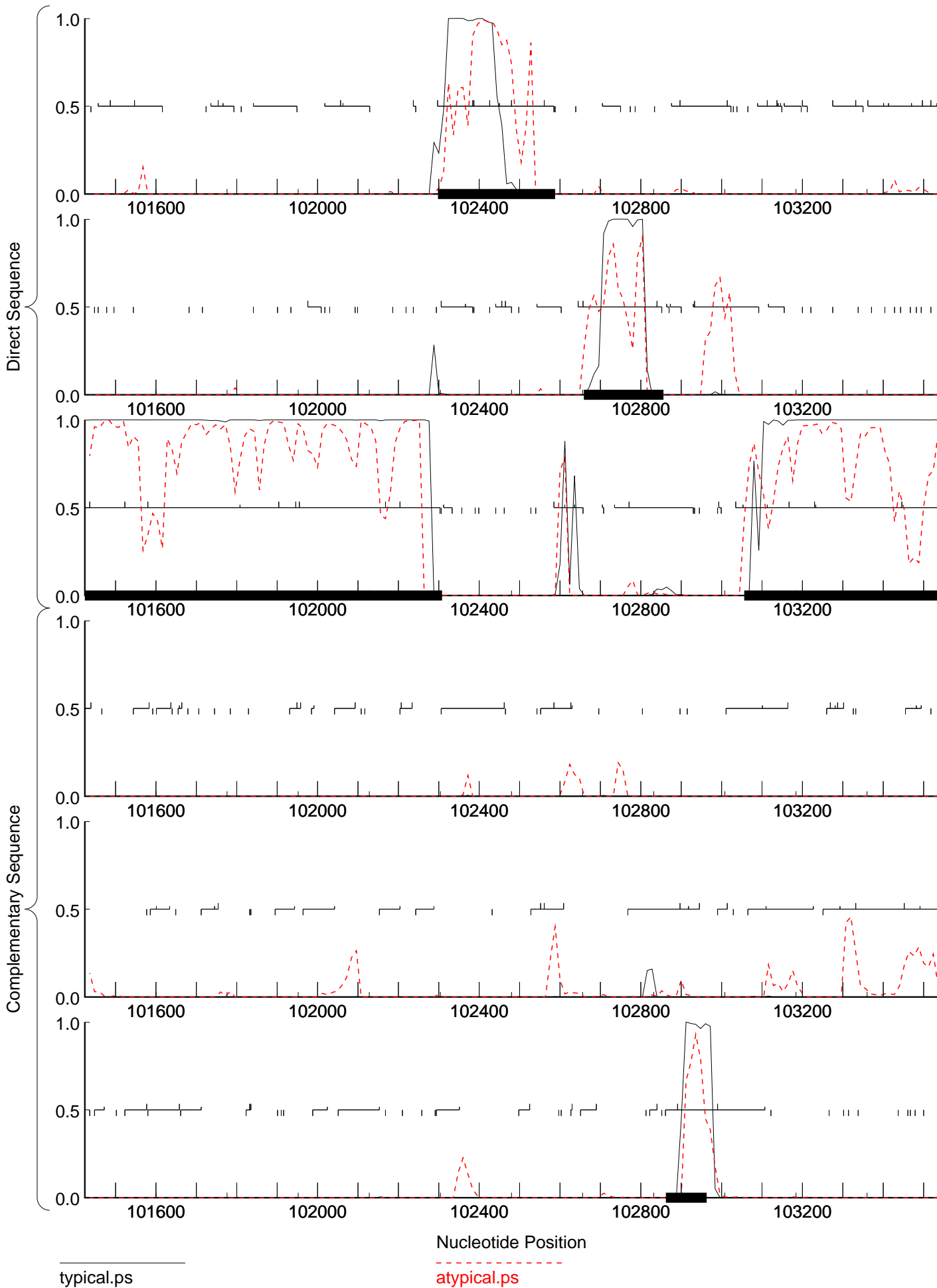
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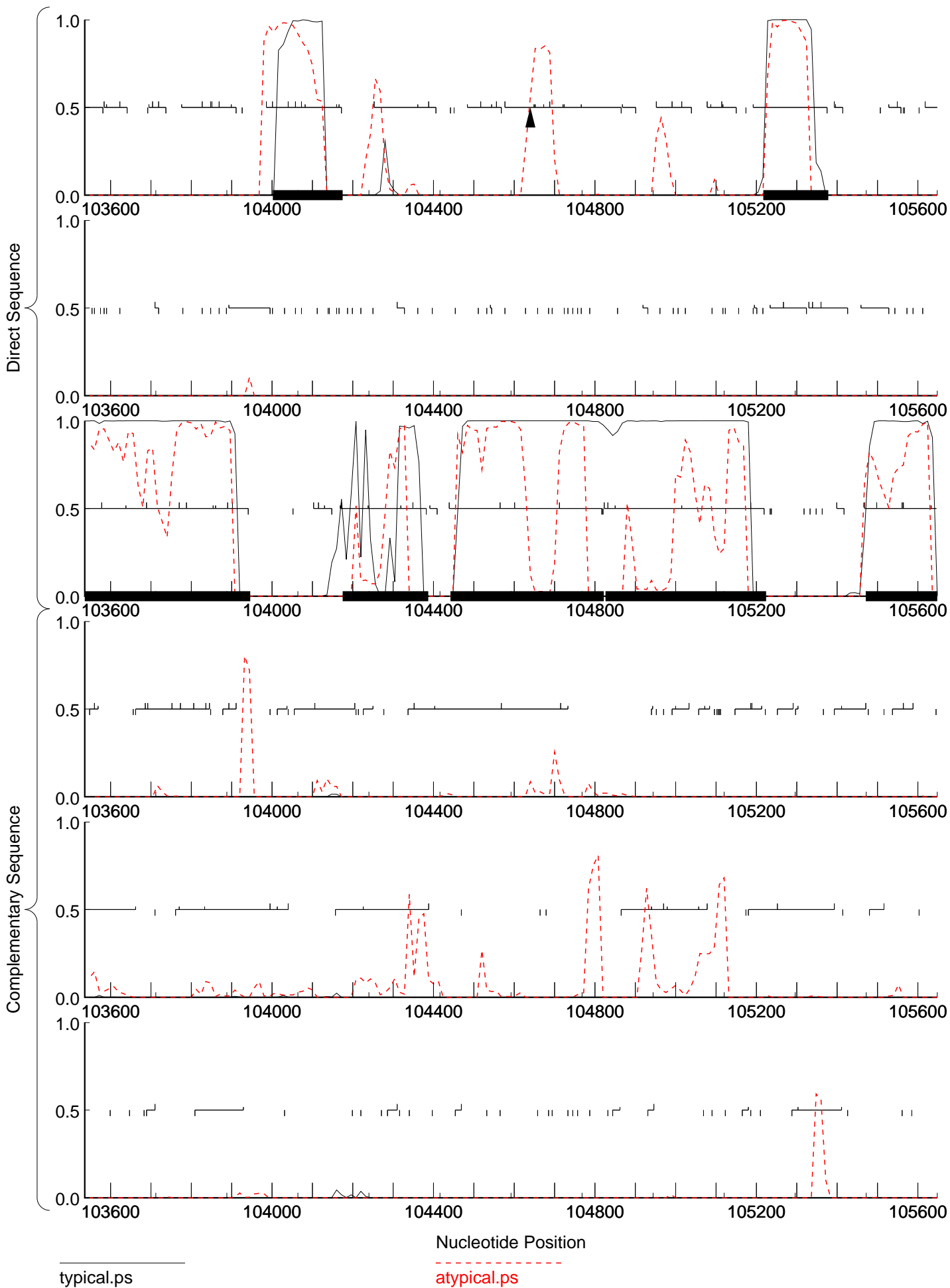
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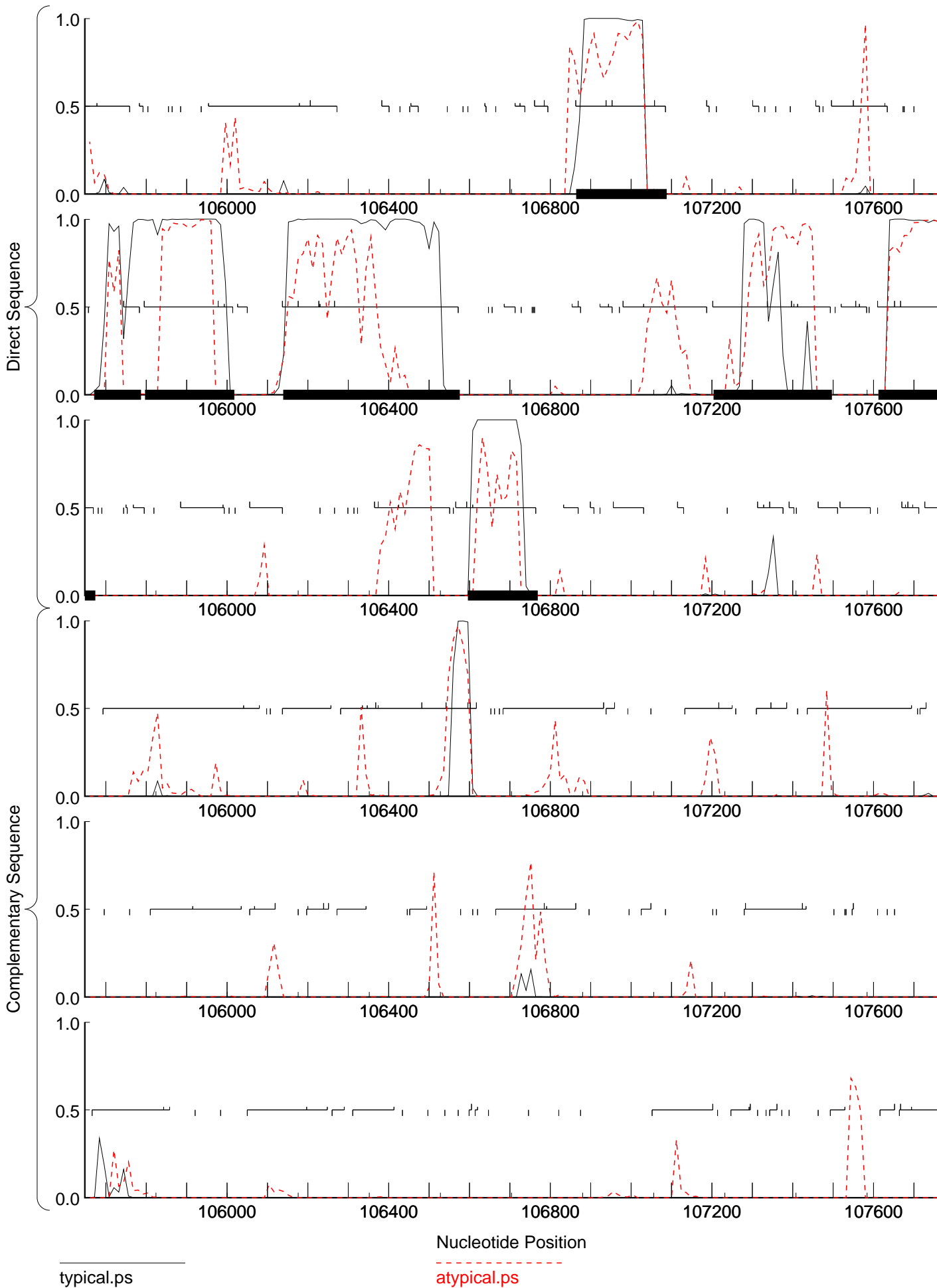
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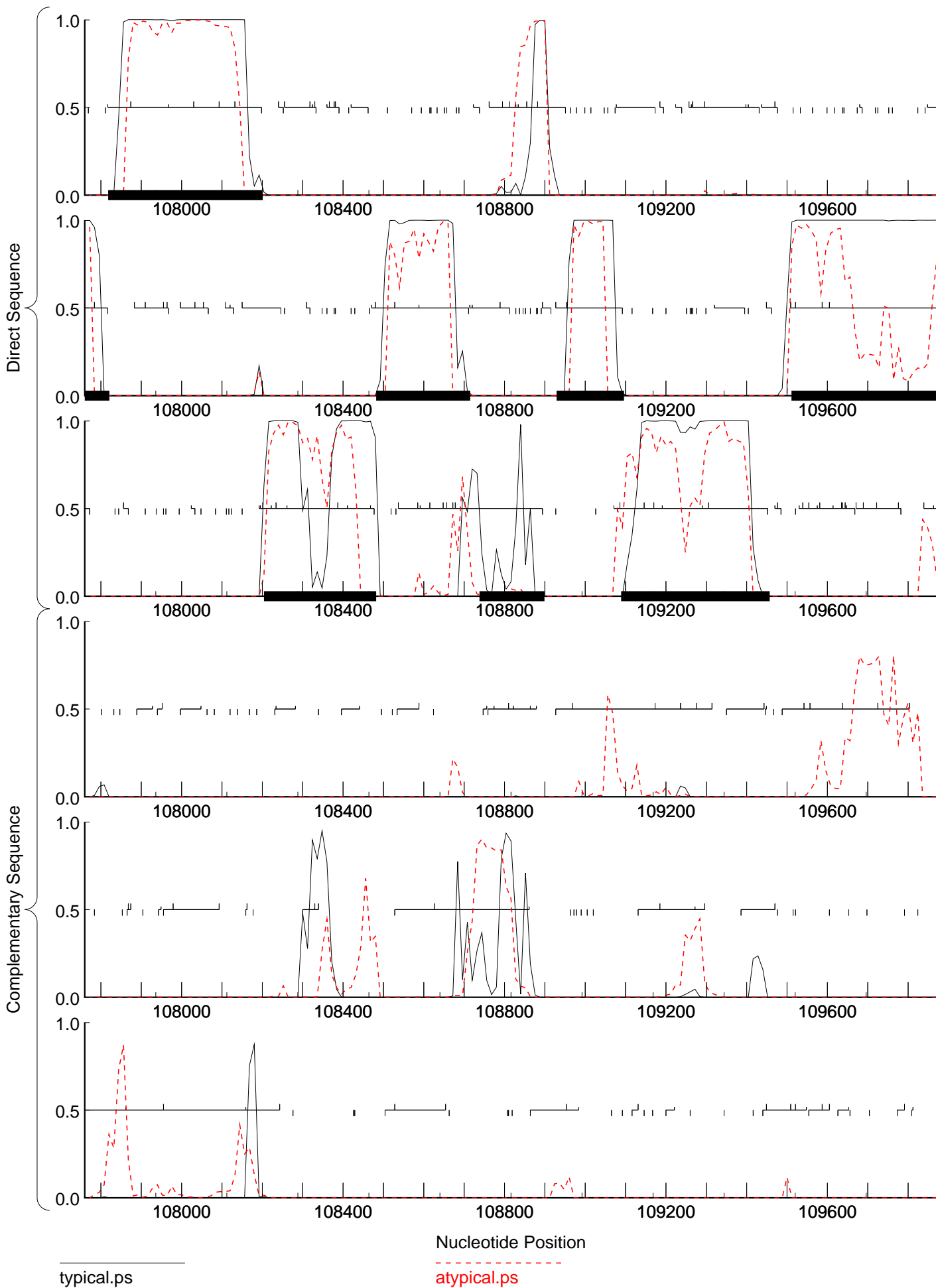
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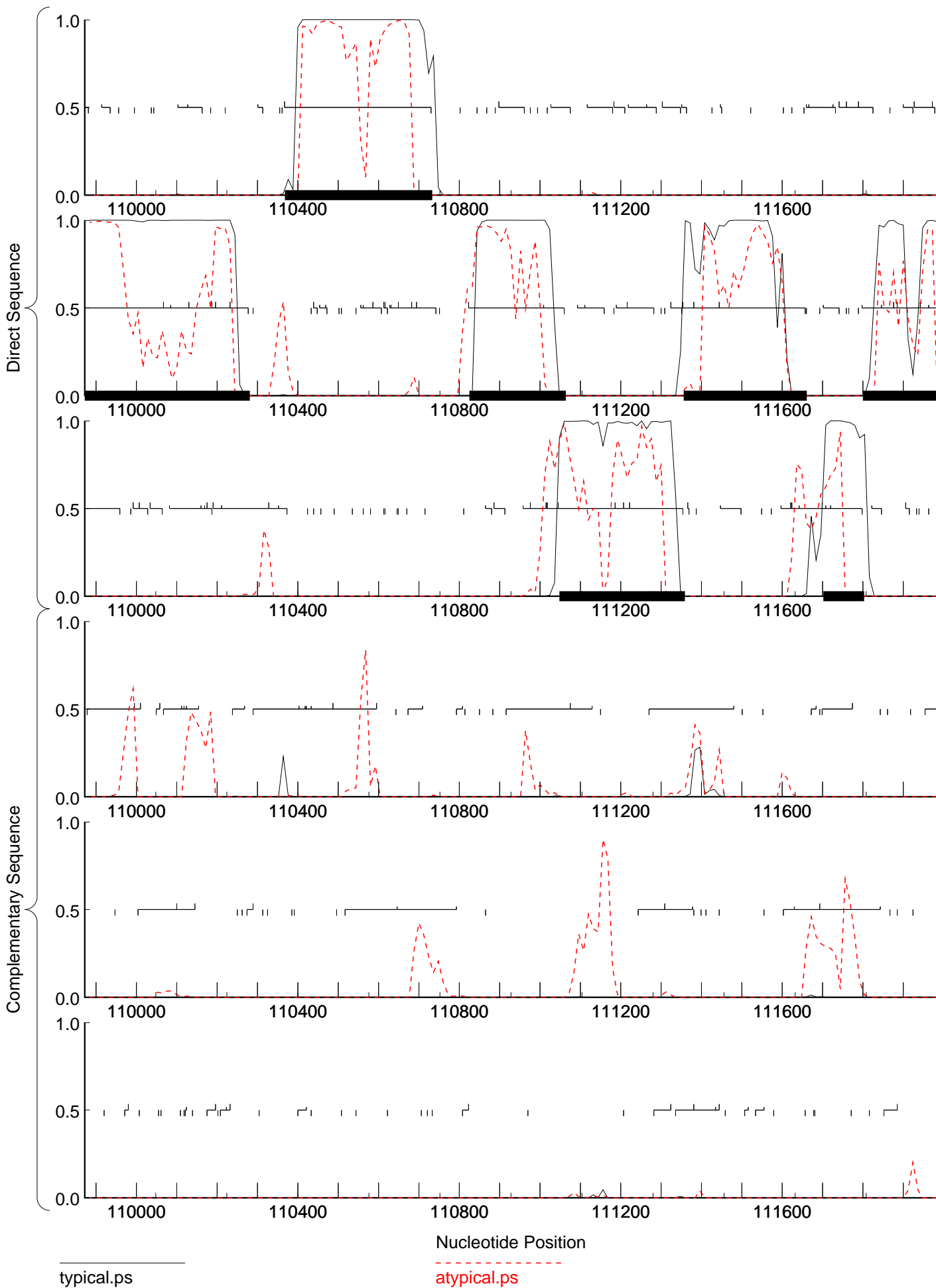
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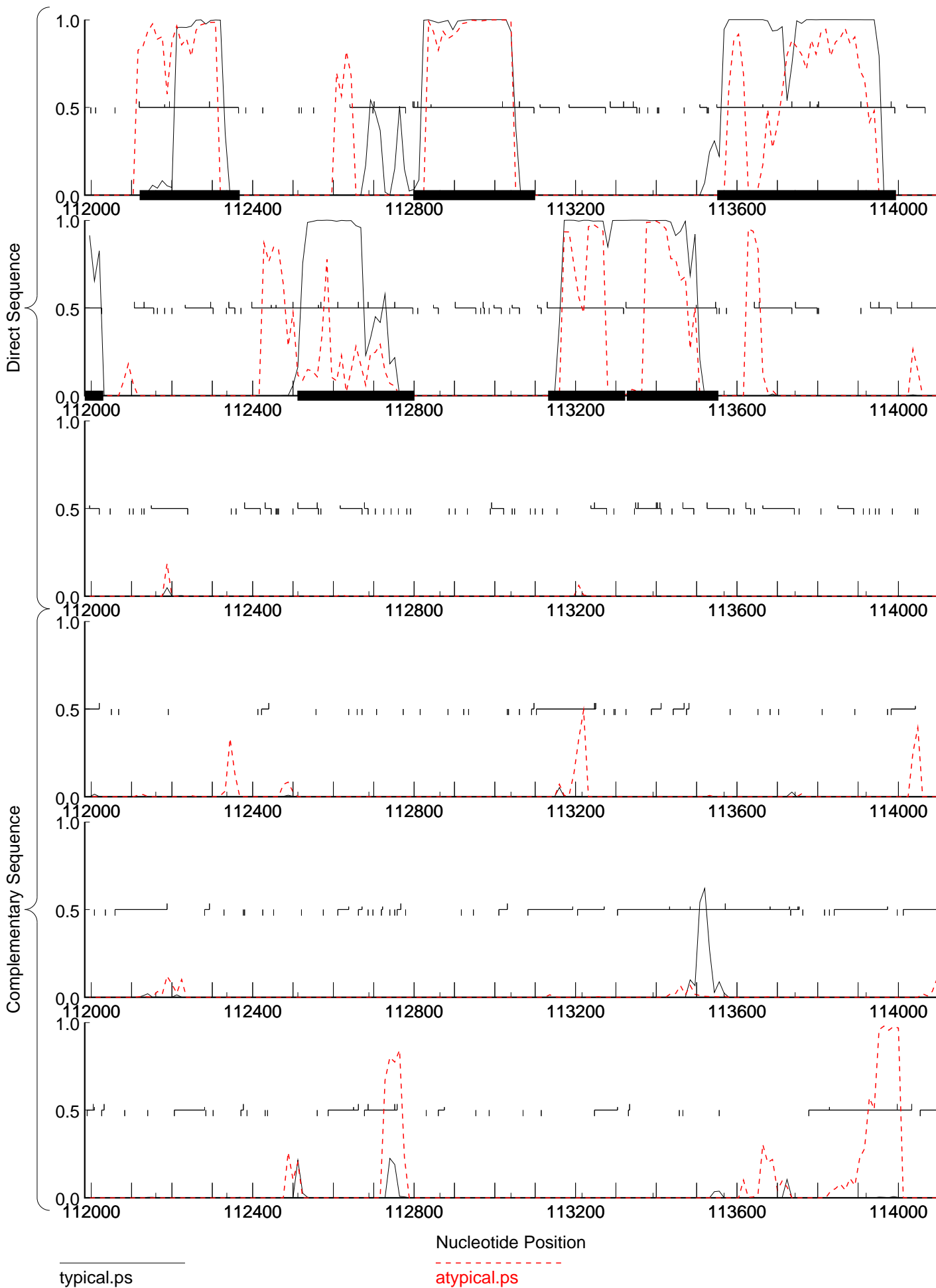
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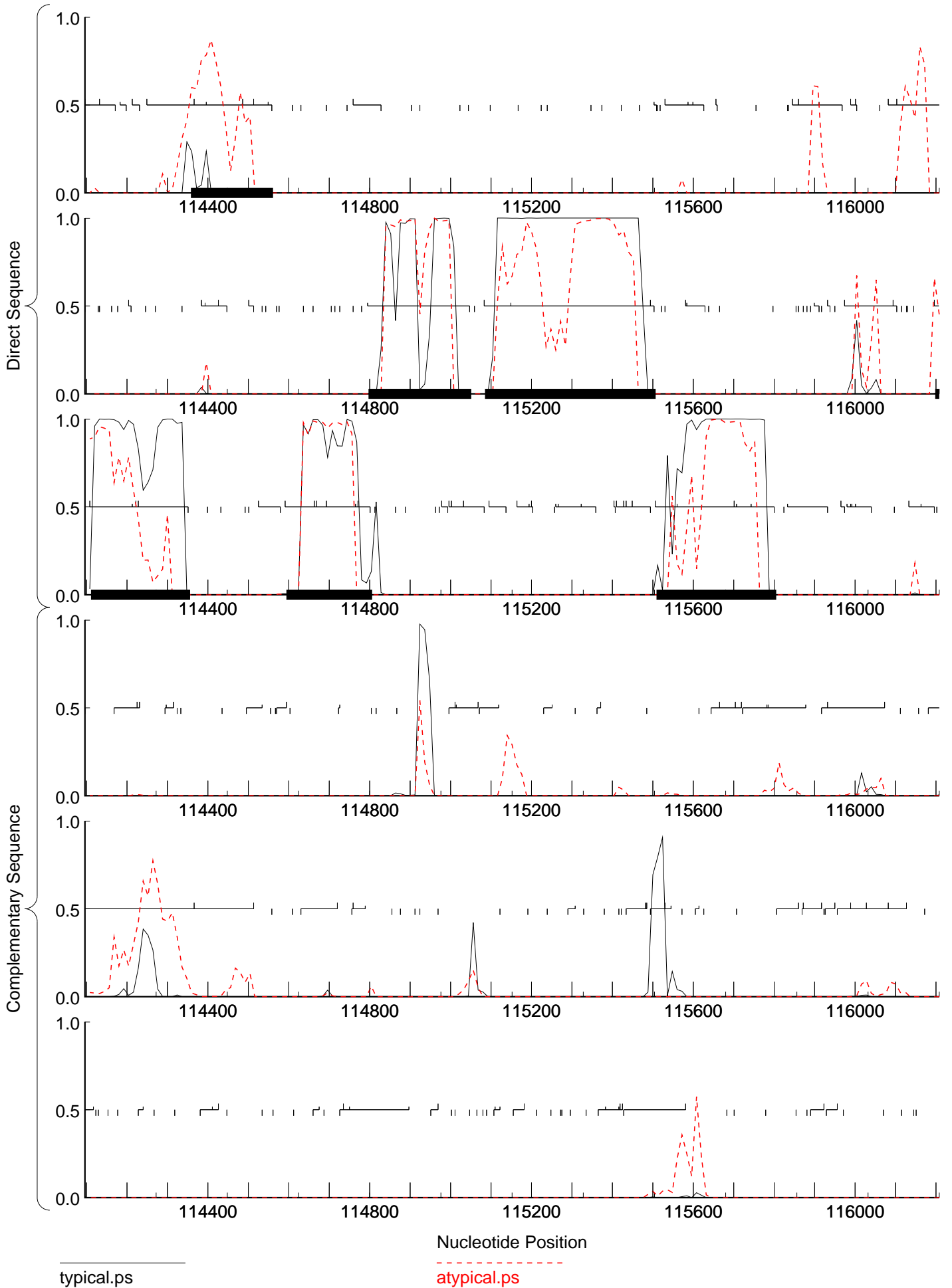


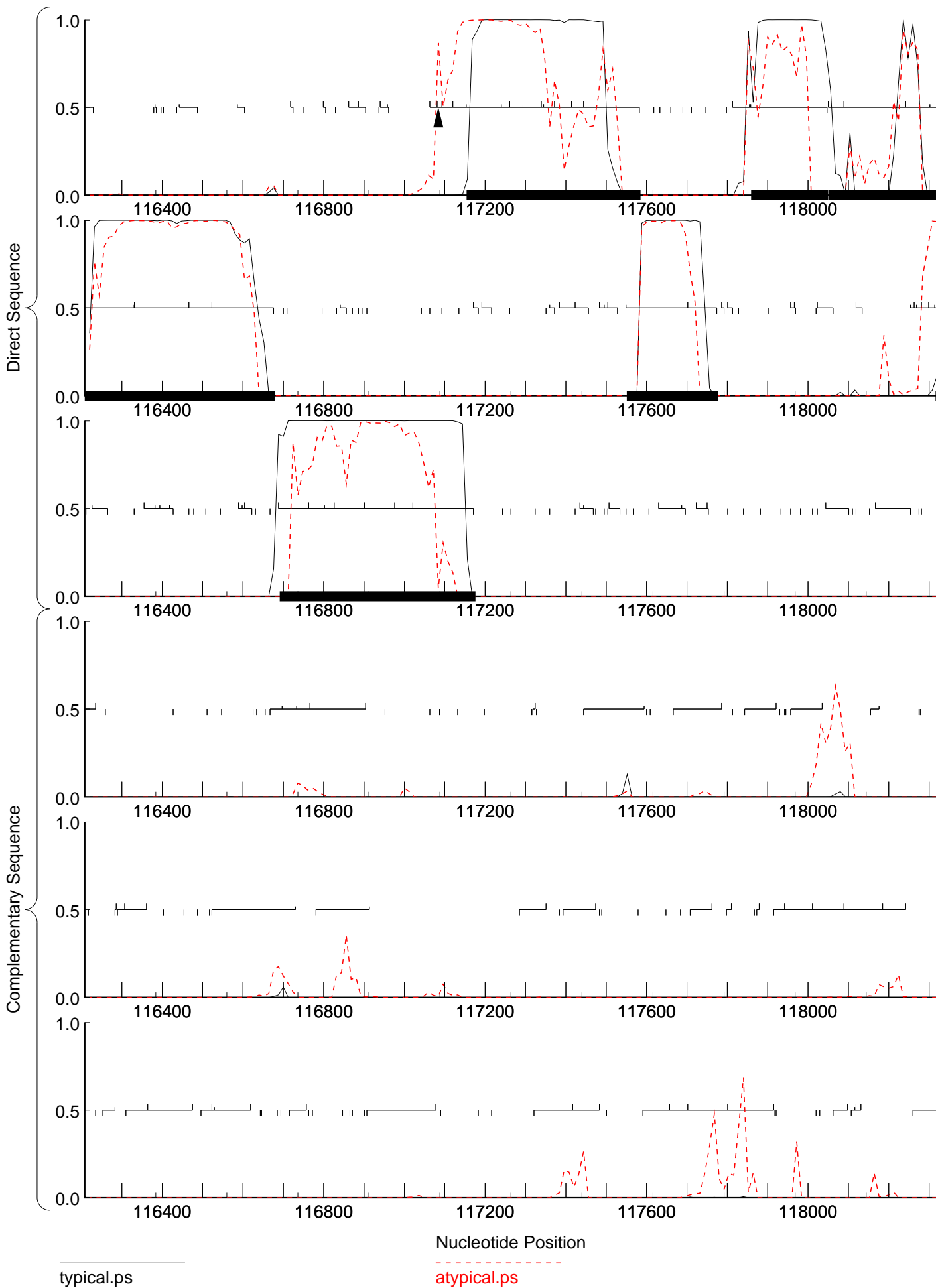
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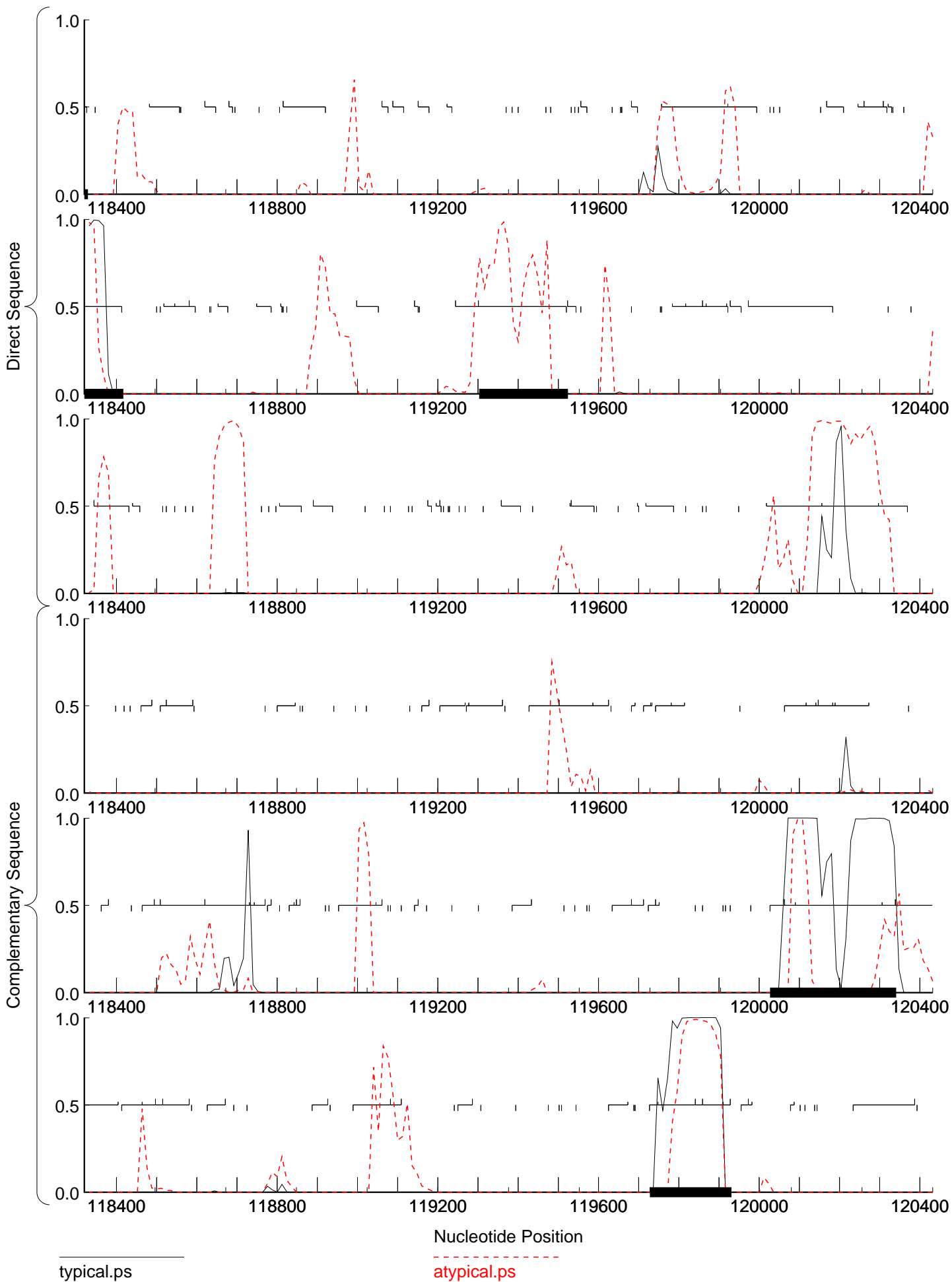
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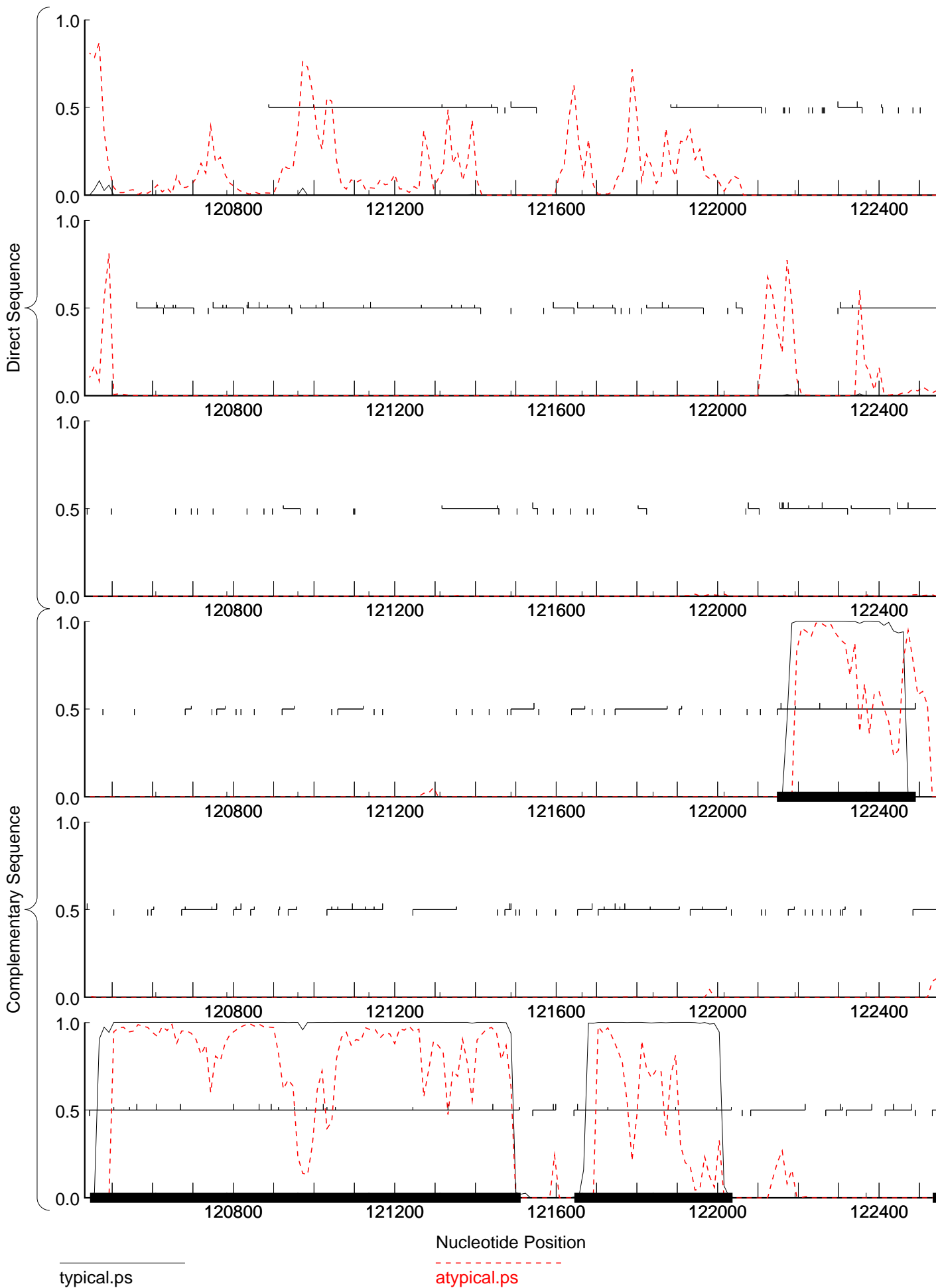


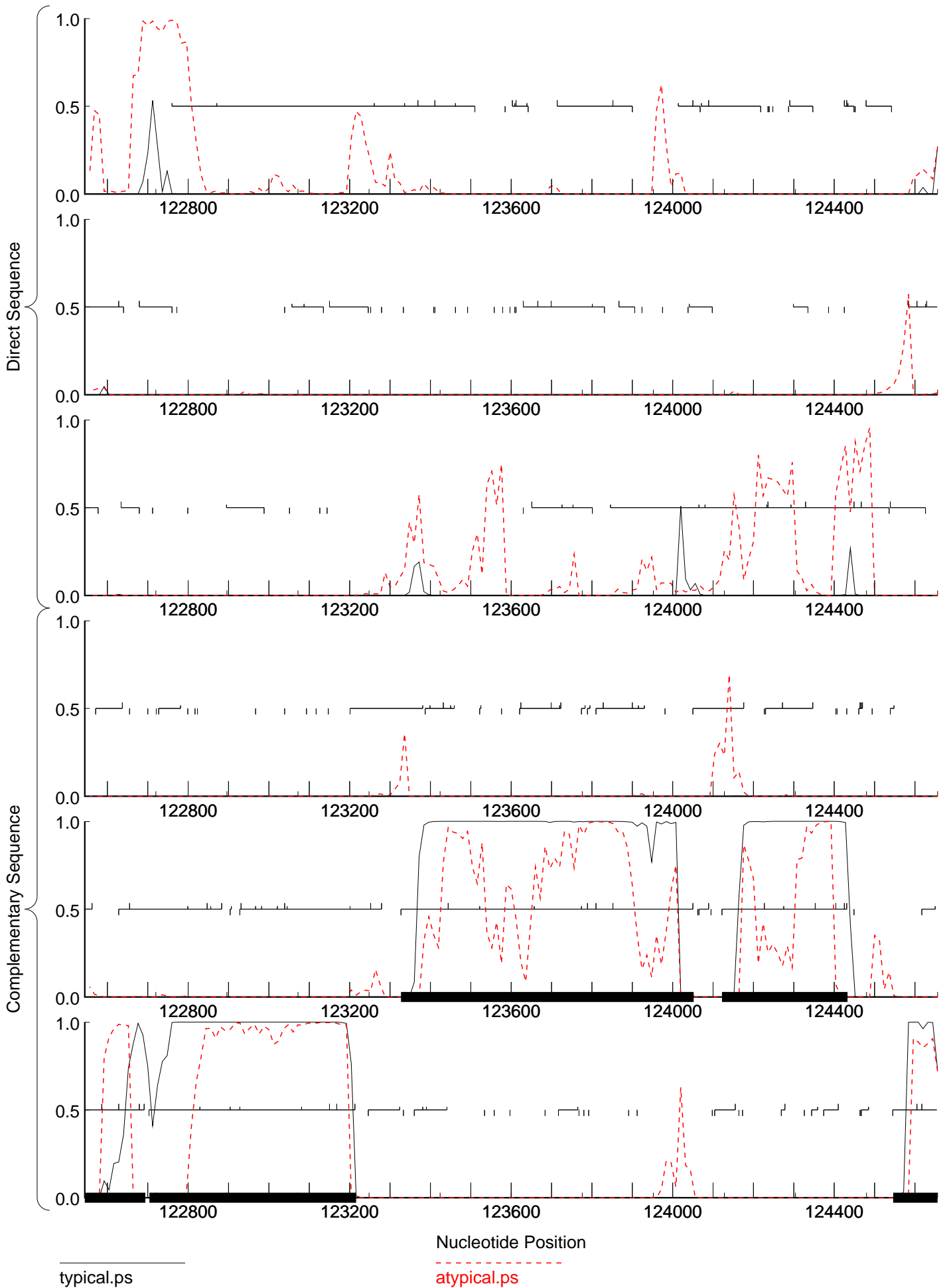


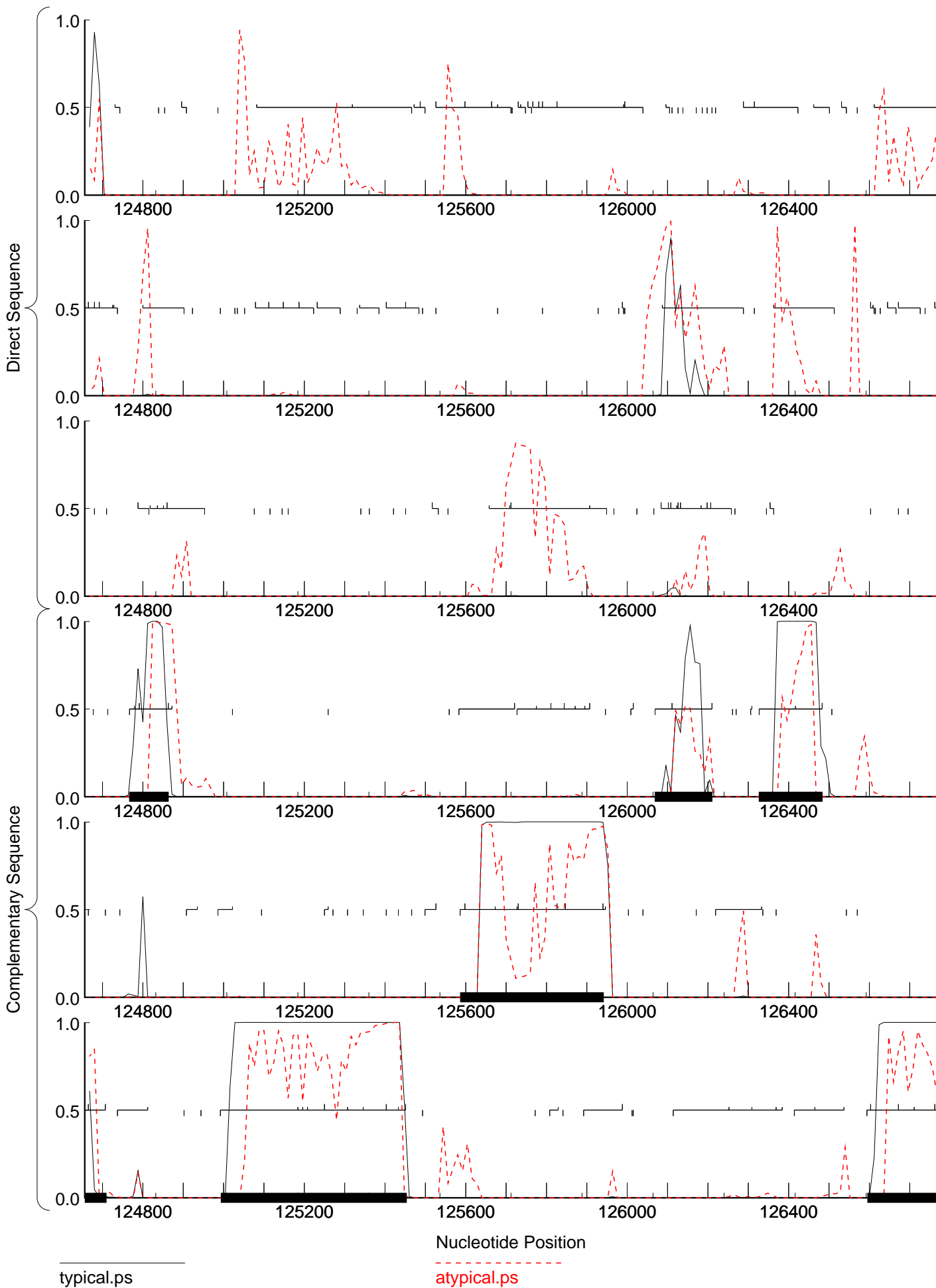


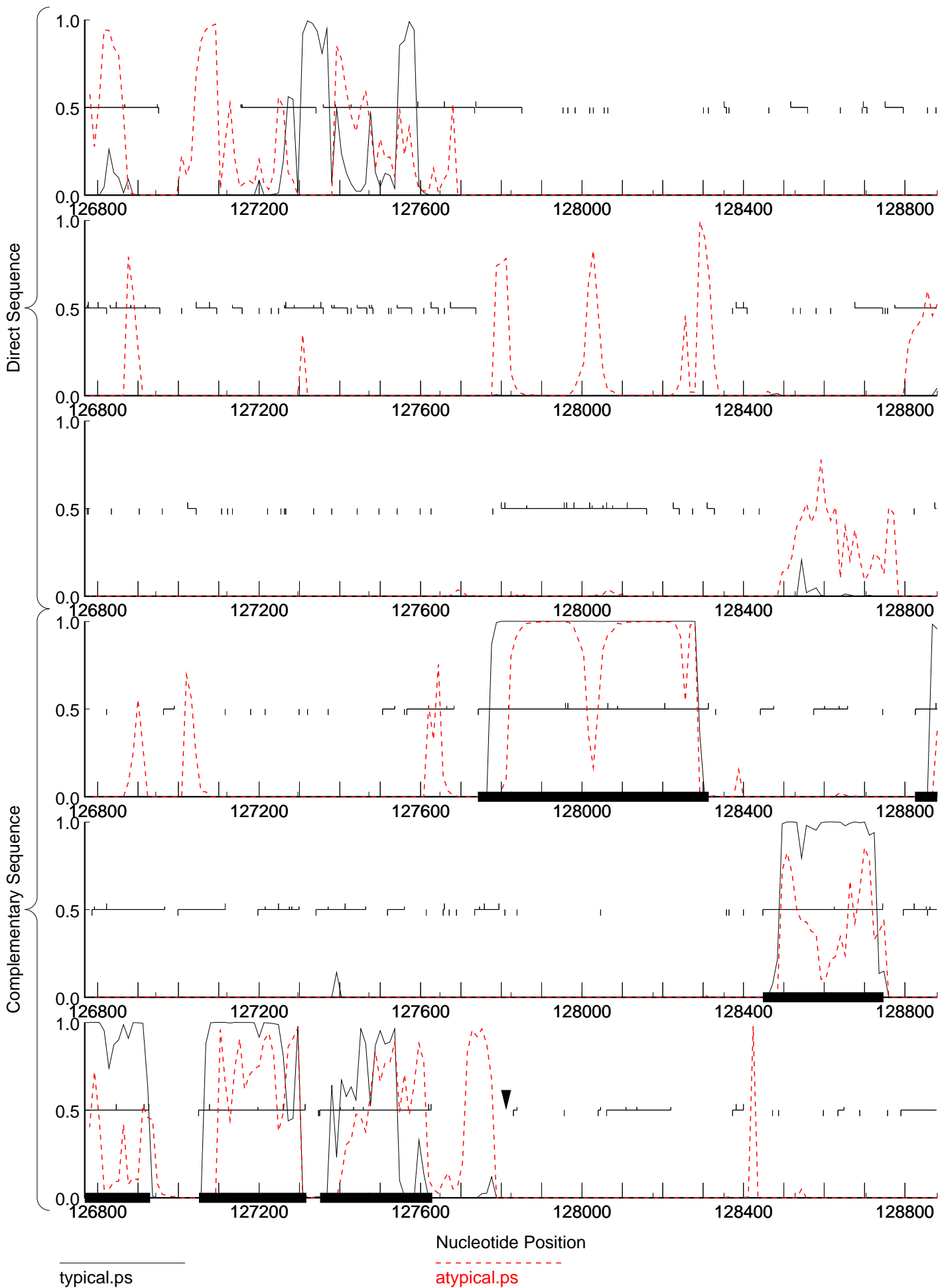
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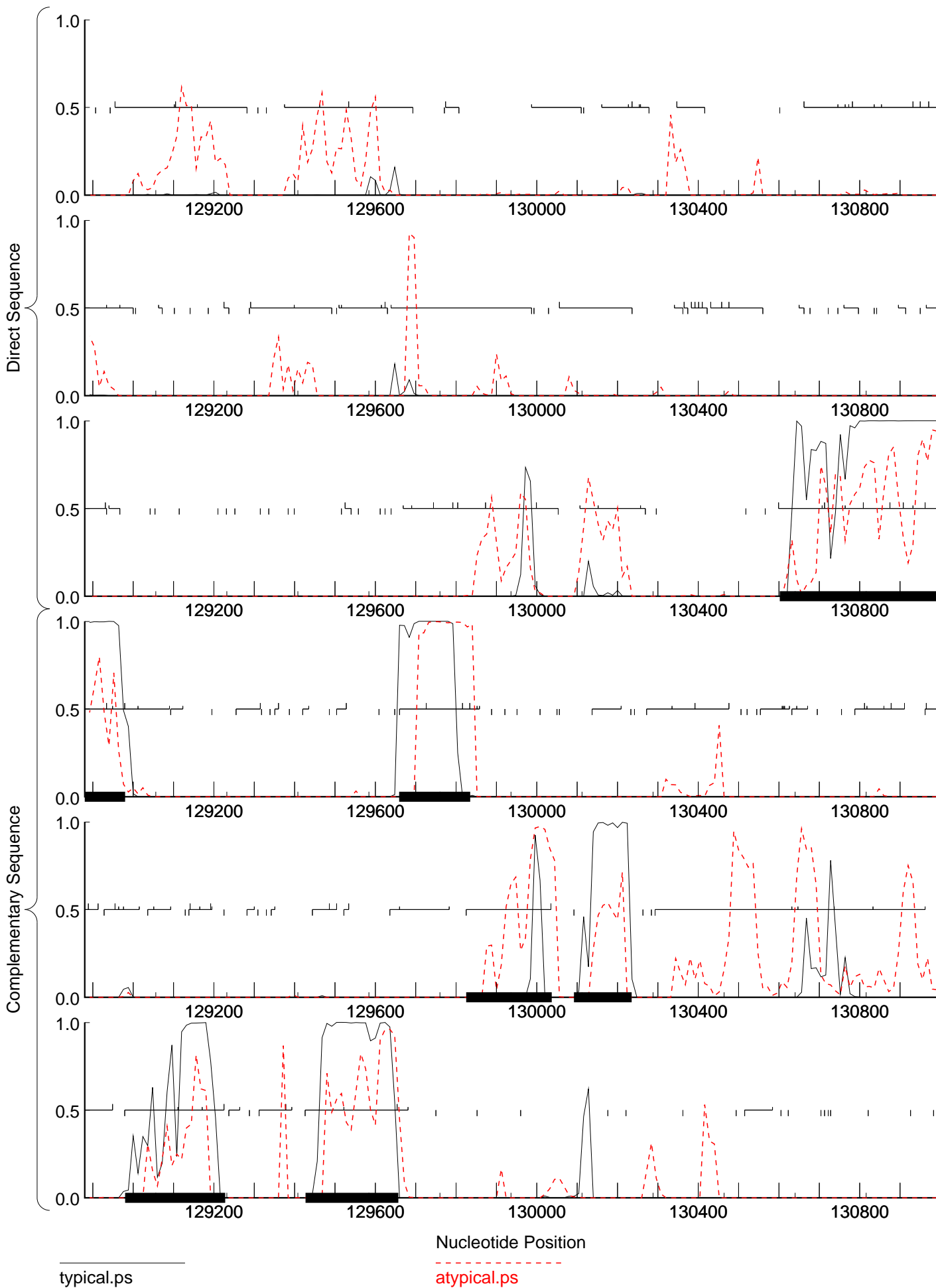
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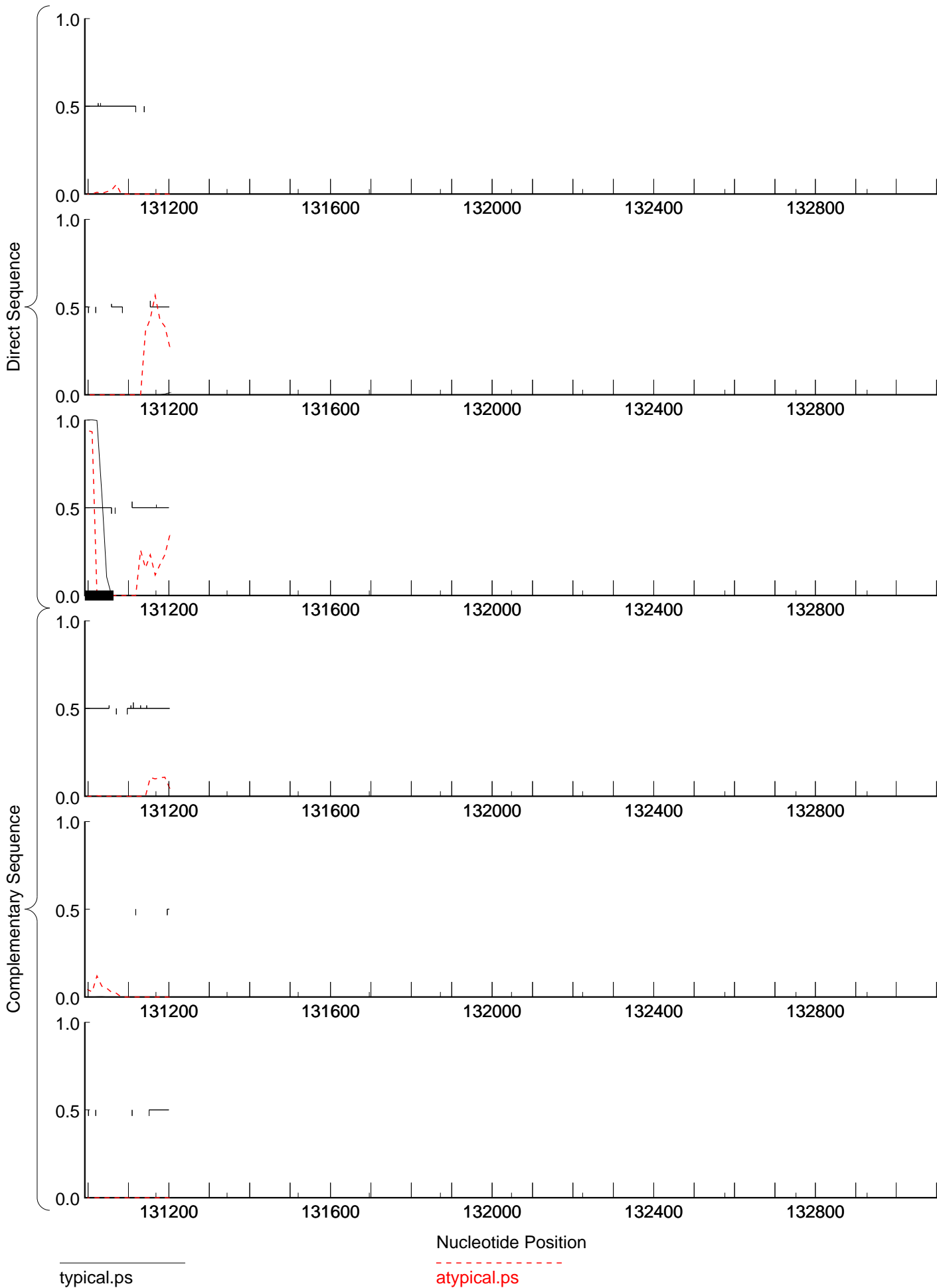












typical.ps

Nucleotide Position

atypical.ps