

GeneMark

Version 2.5p (09.08.06)

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PROGRAM INFORMATION

Sequence : Streptomyces phage PumpkinSpice complete sequence, 132480 bp including 12588 bp terminal re
Analysis Date : 4/20/18 at 1:03:08
Pages : 64
Sequence Length : 132480 bp
GC Content : 49.44%

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,

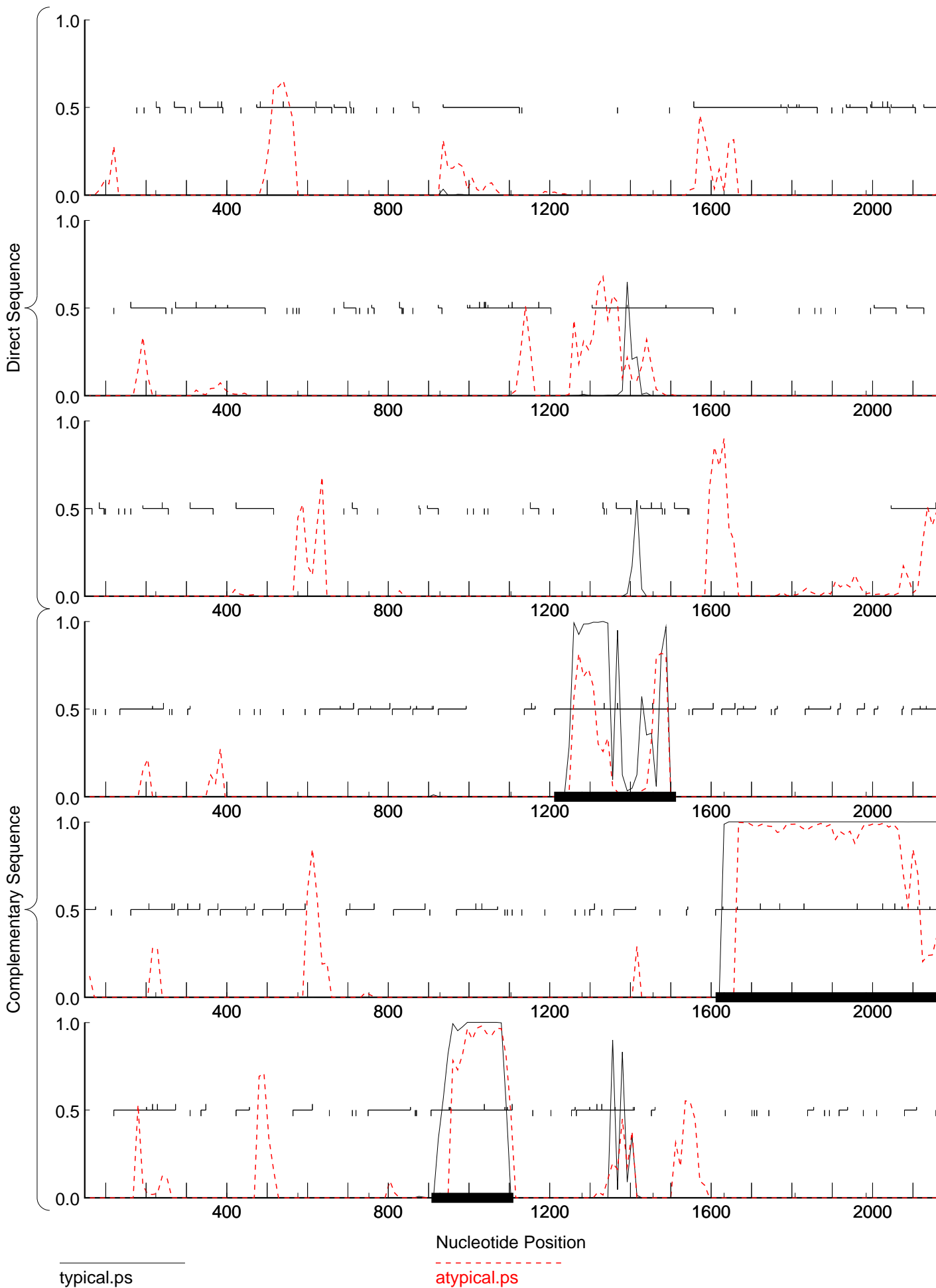
MATRIX INFORMATION

Matrix : -
Author : -
Order : 4

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

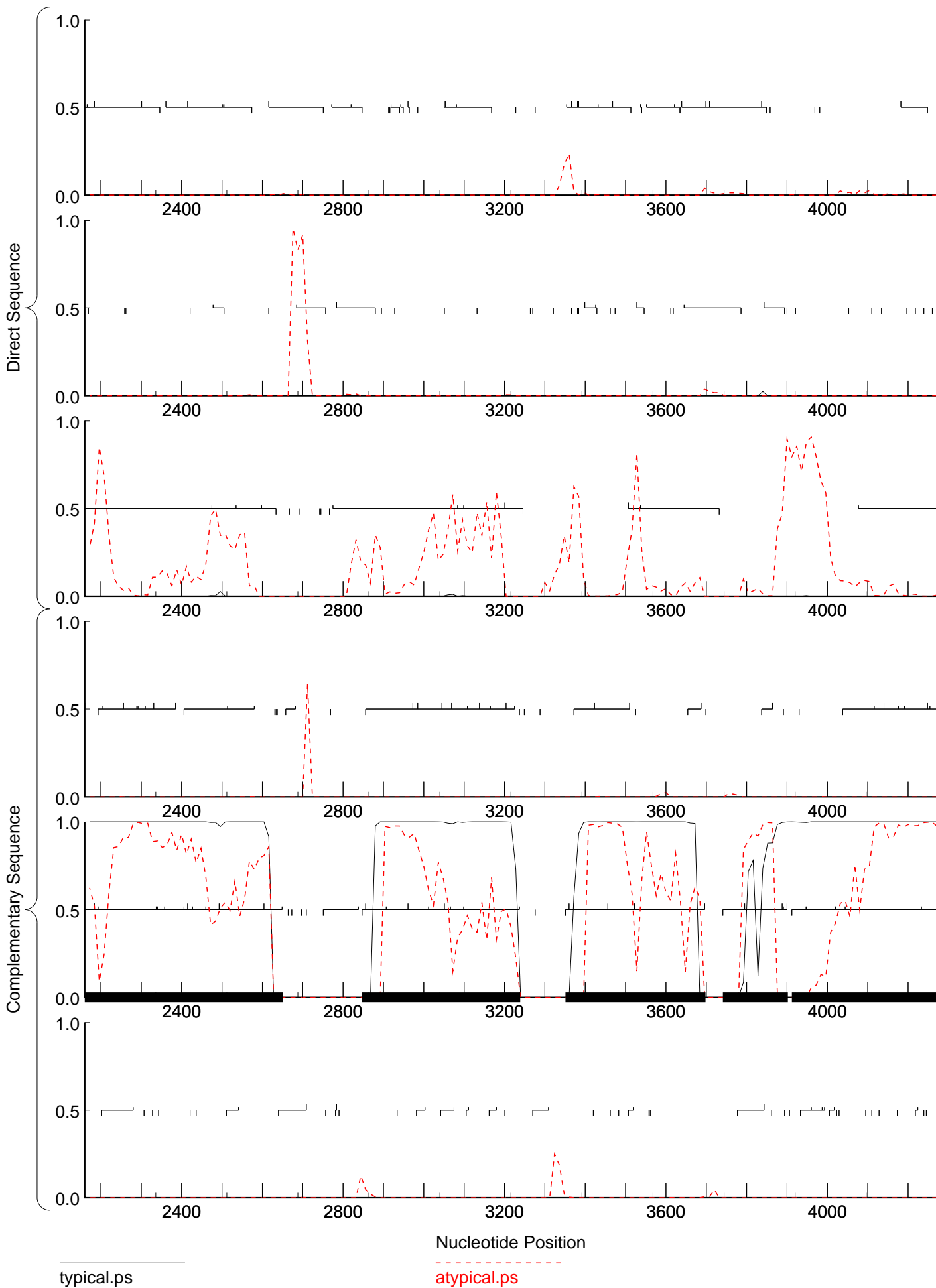
Matrix notes & comments

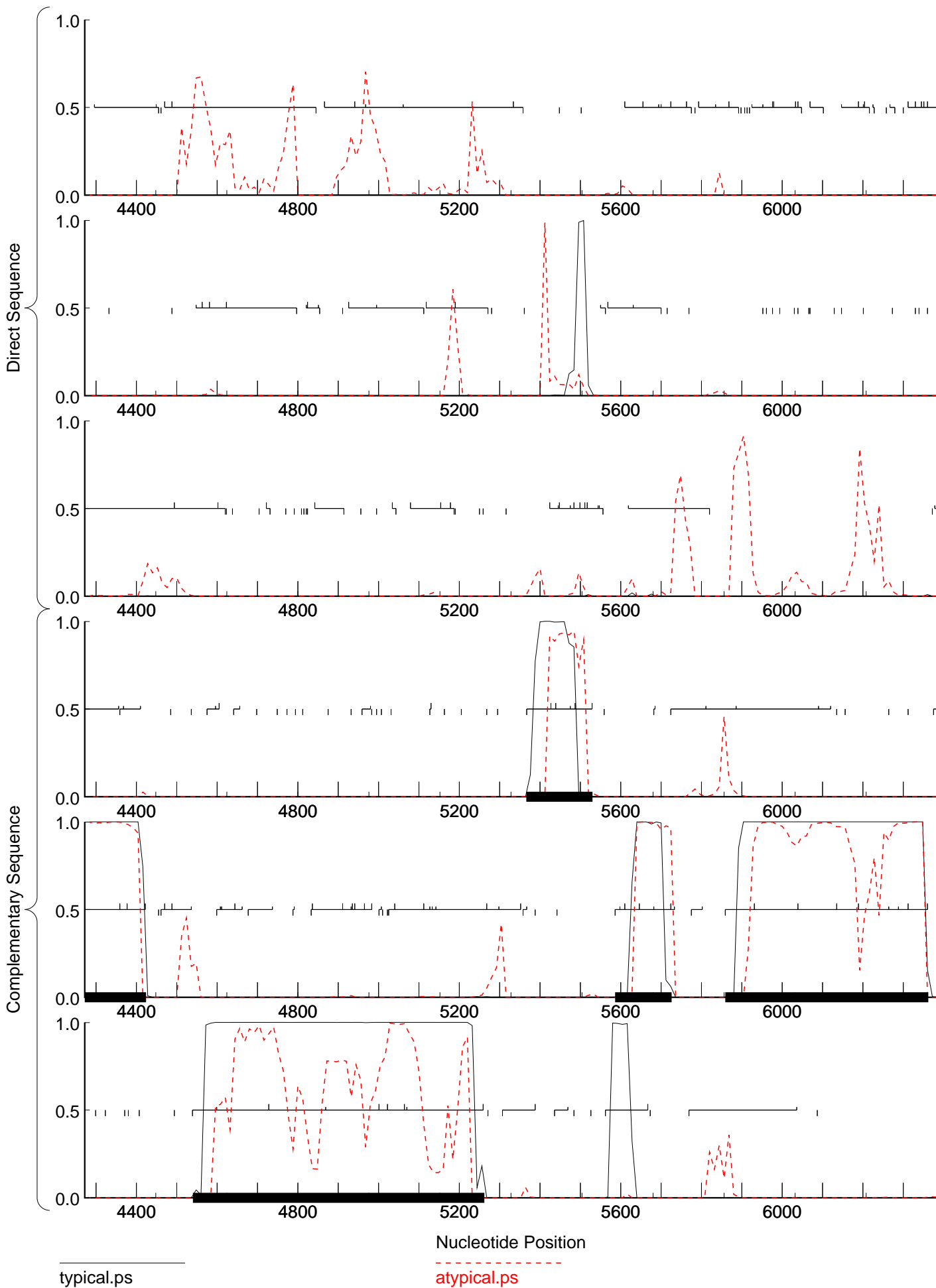
Training set derived by GeneMarkS, 4.30 October 2014
Fri Apr 20 01:03:08 2018



typical.ps

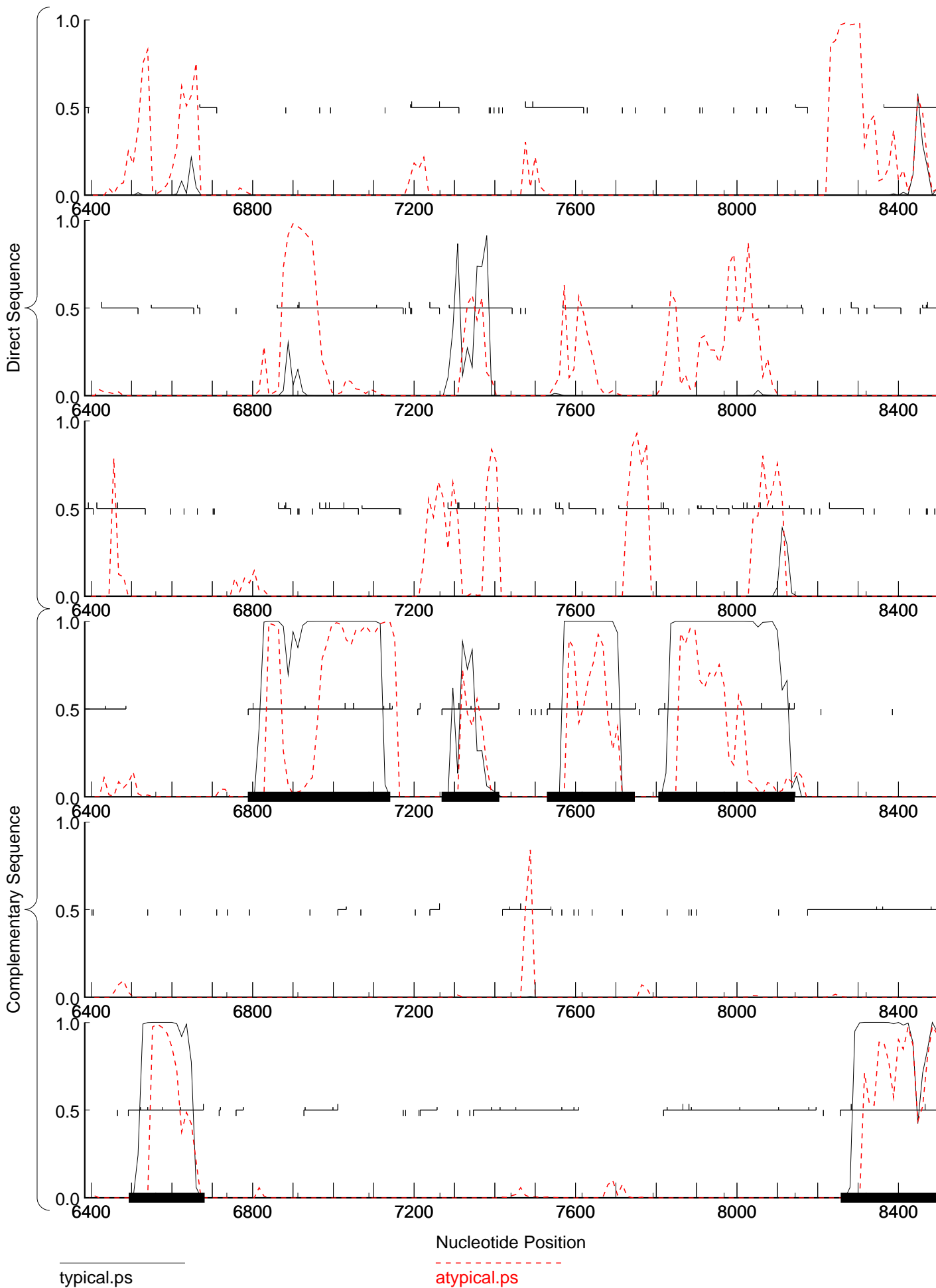
atypical.ps





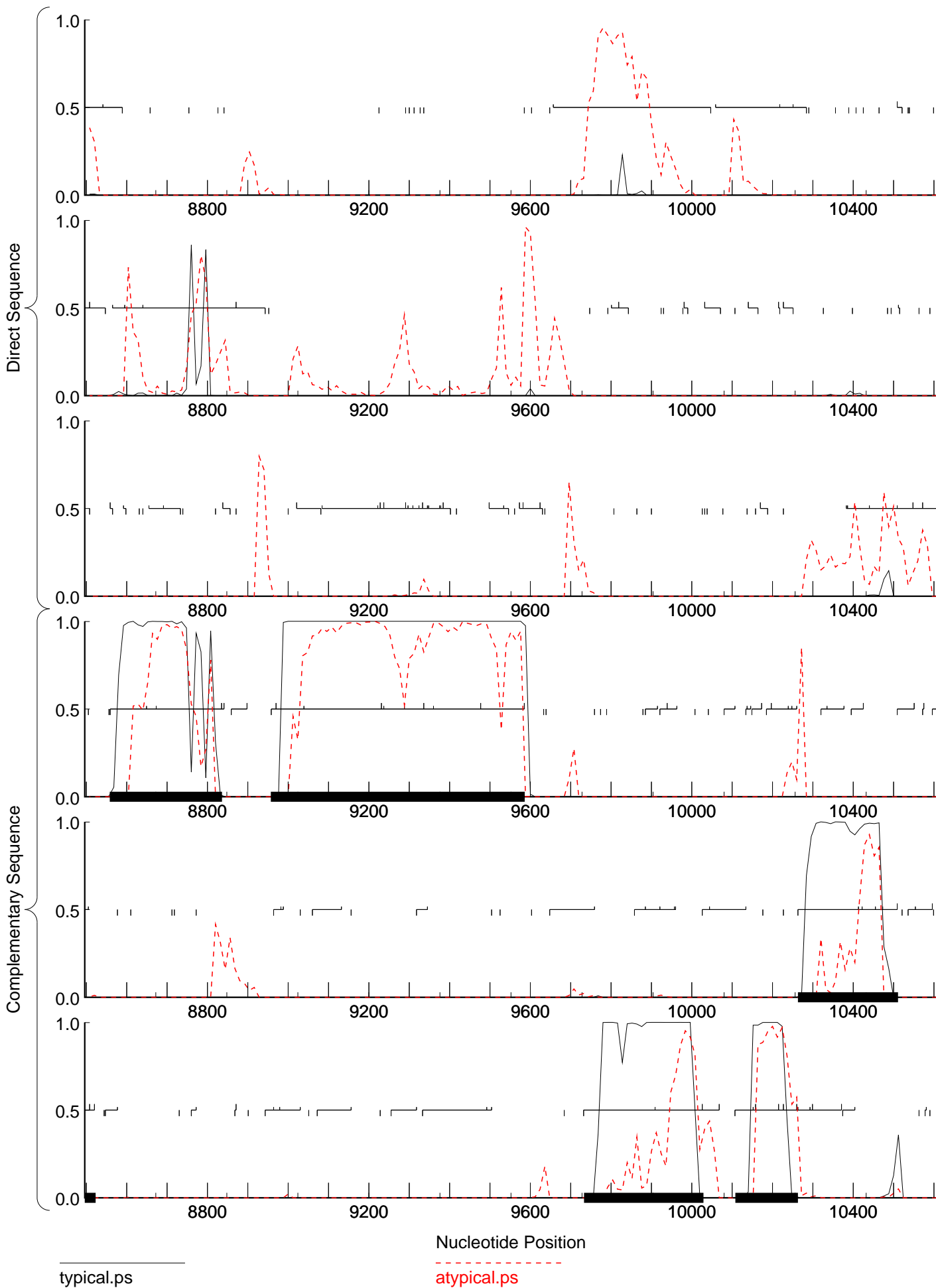
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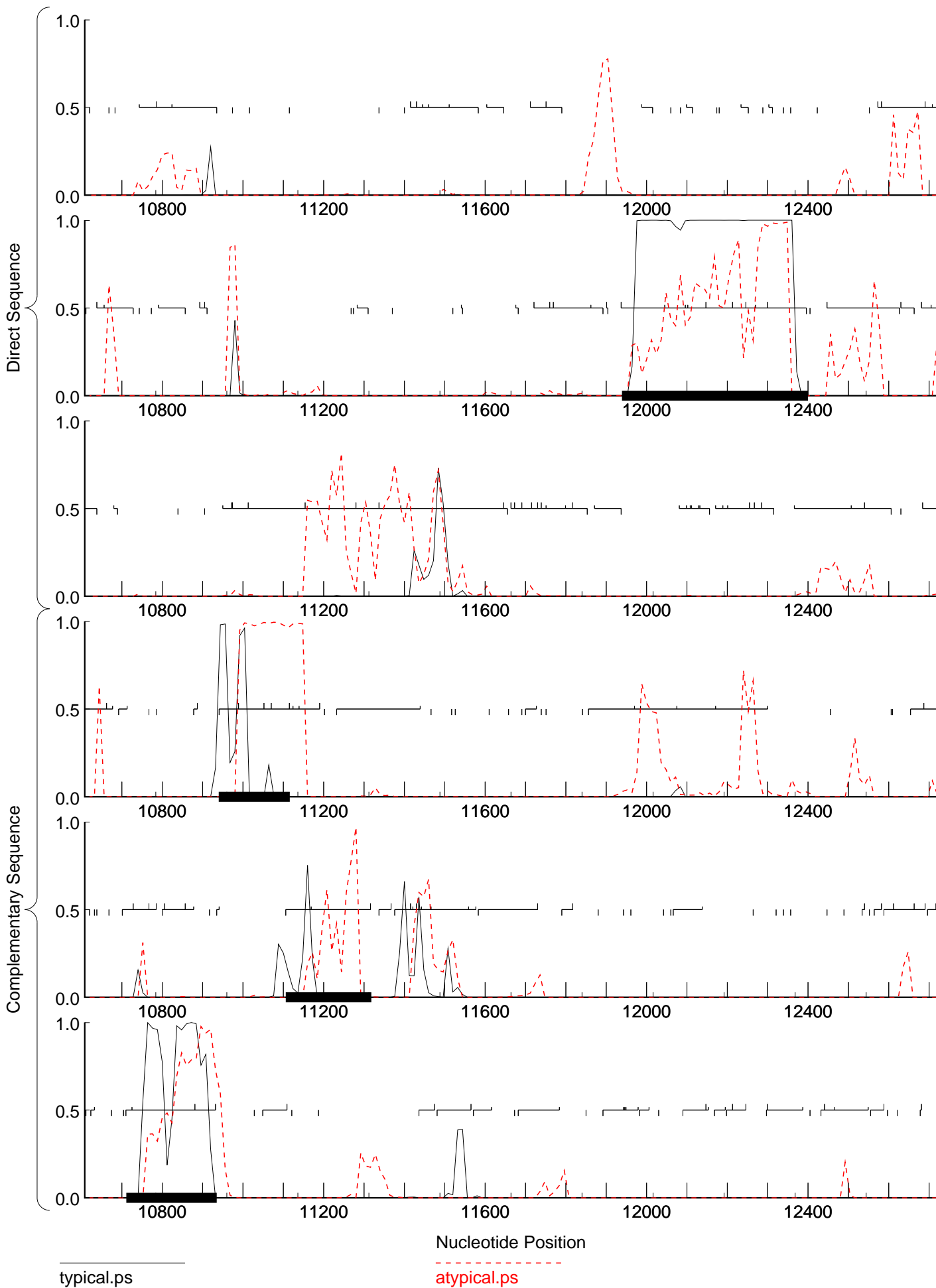
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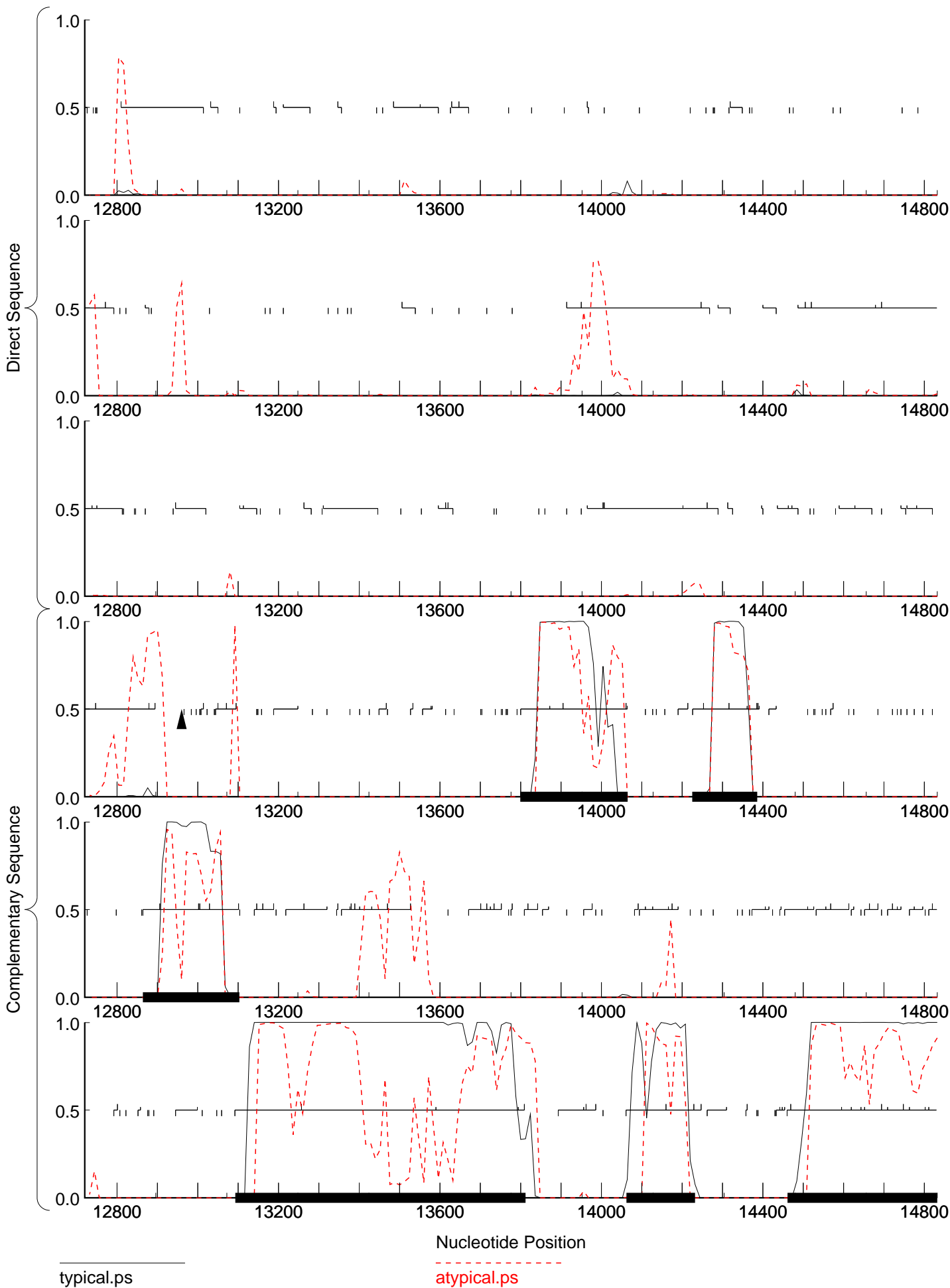


typical.ps

atypical.ps

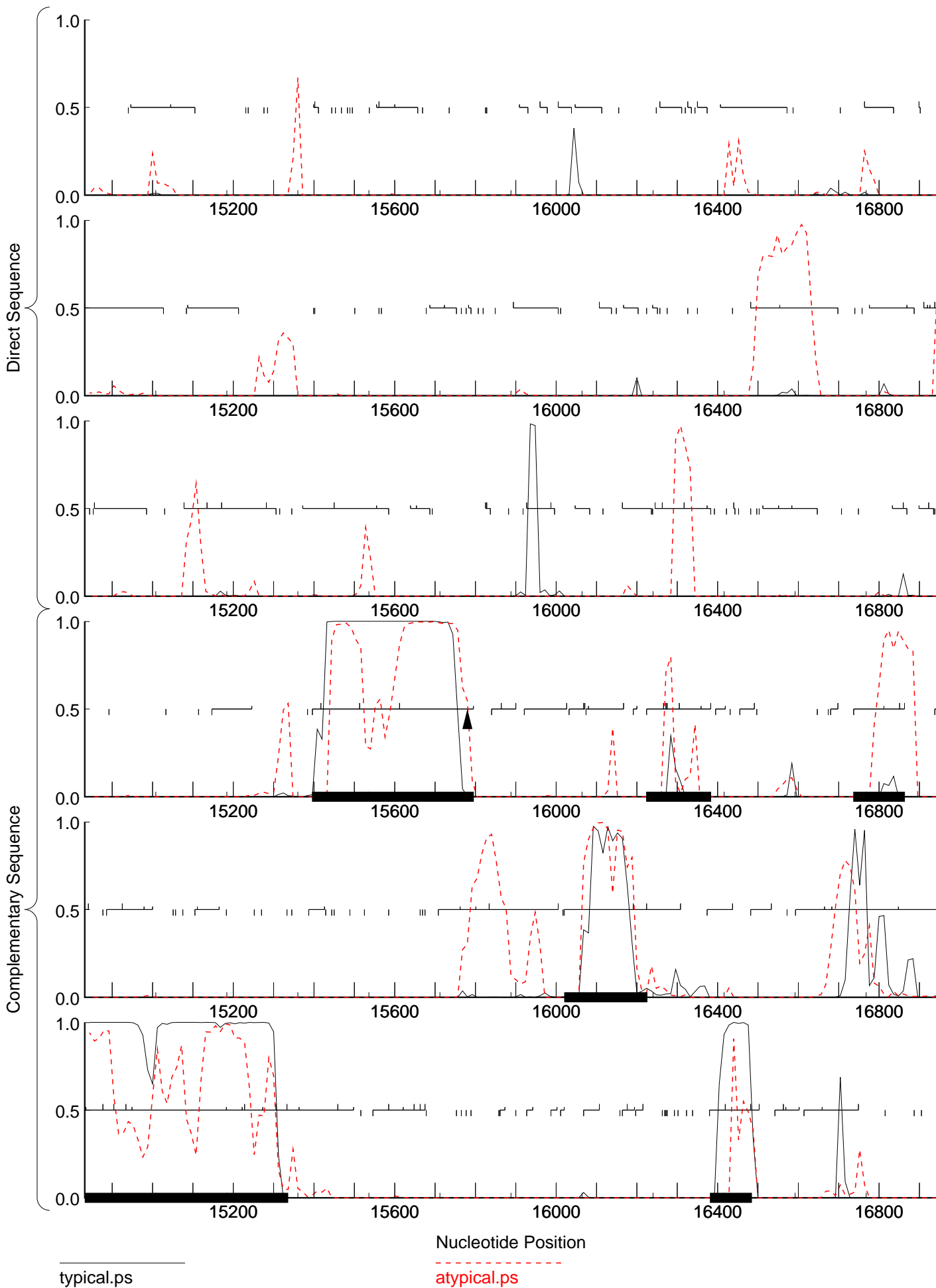


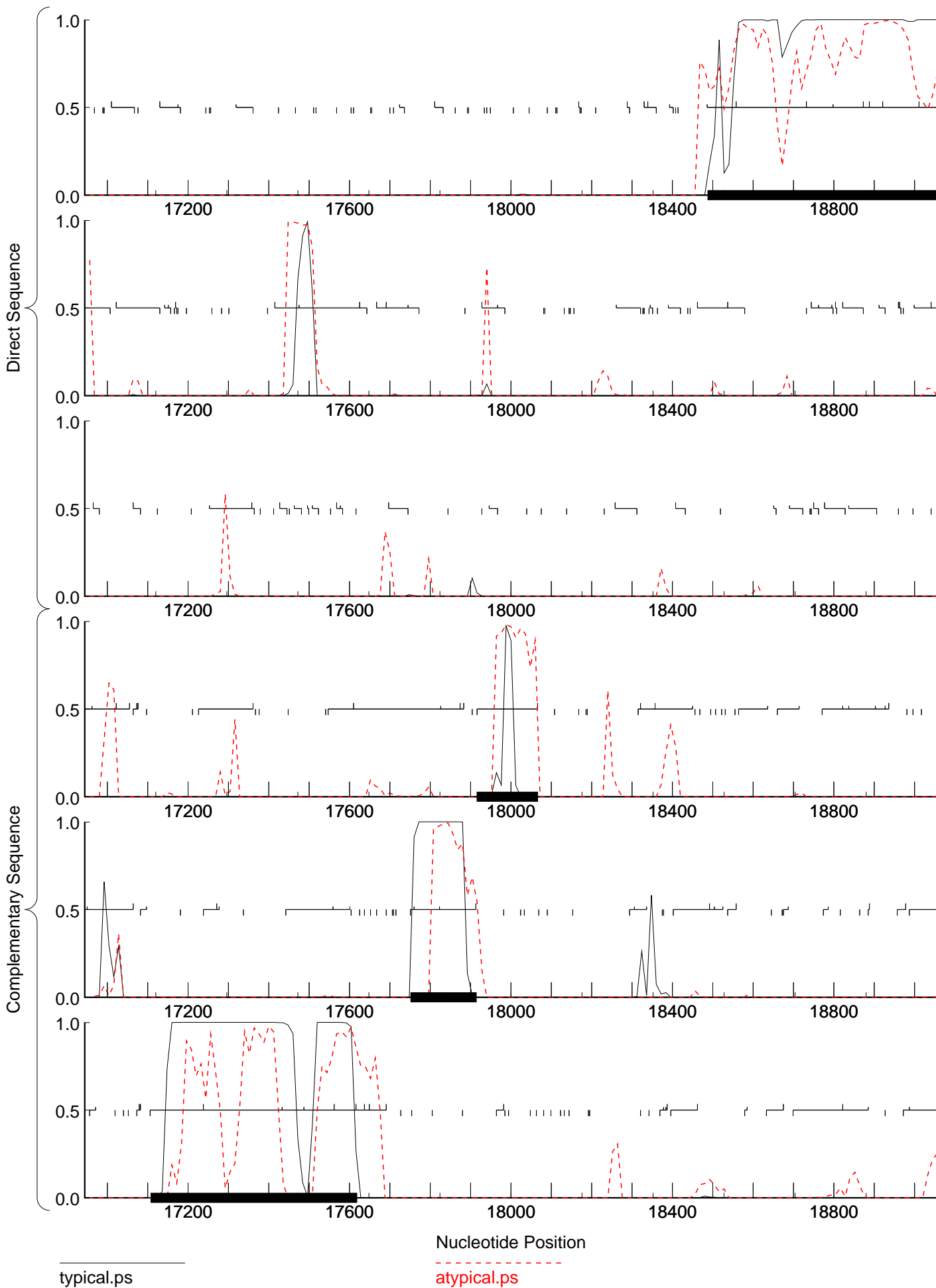


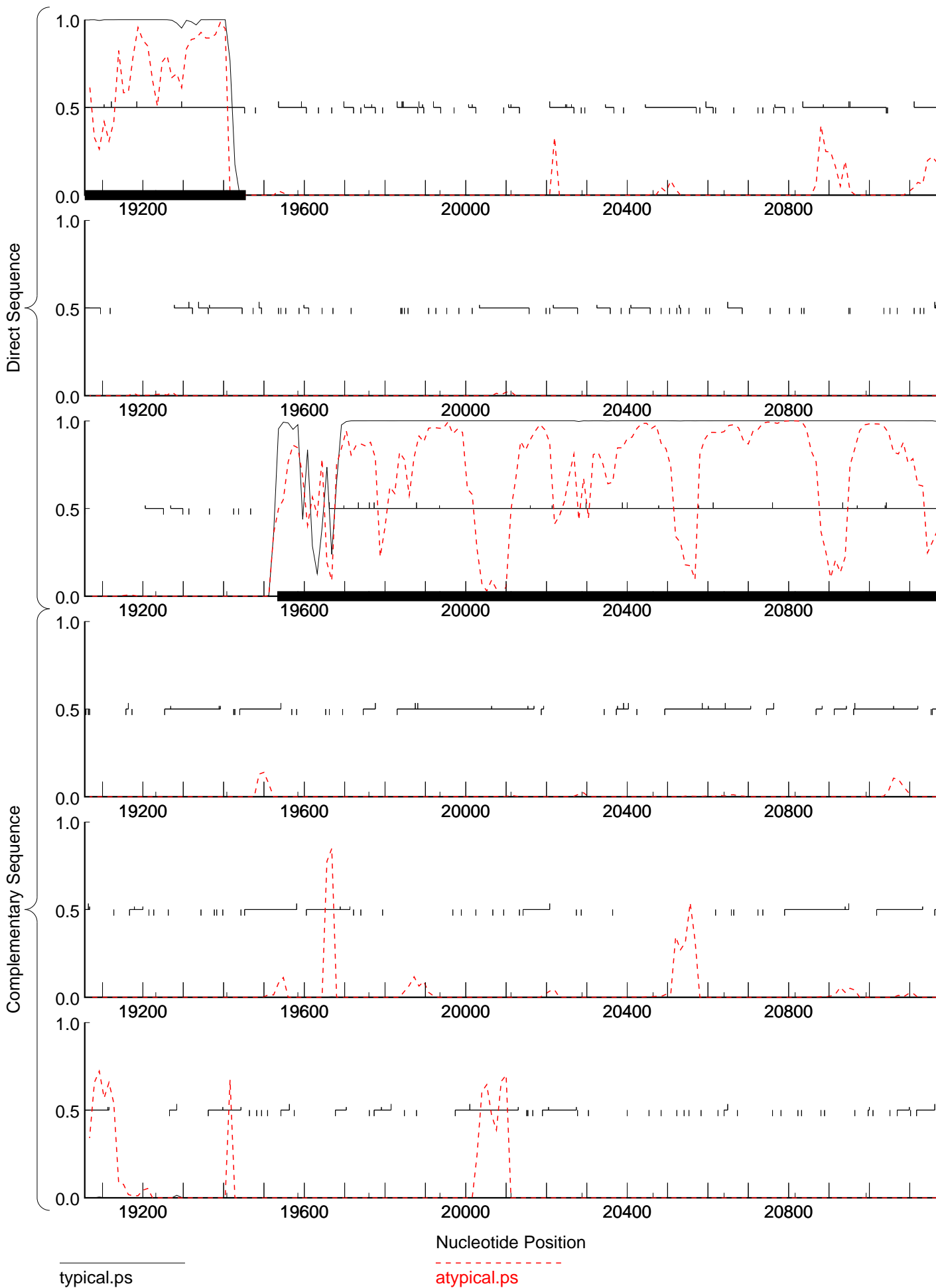


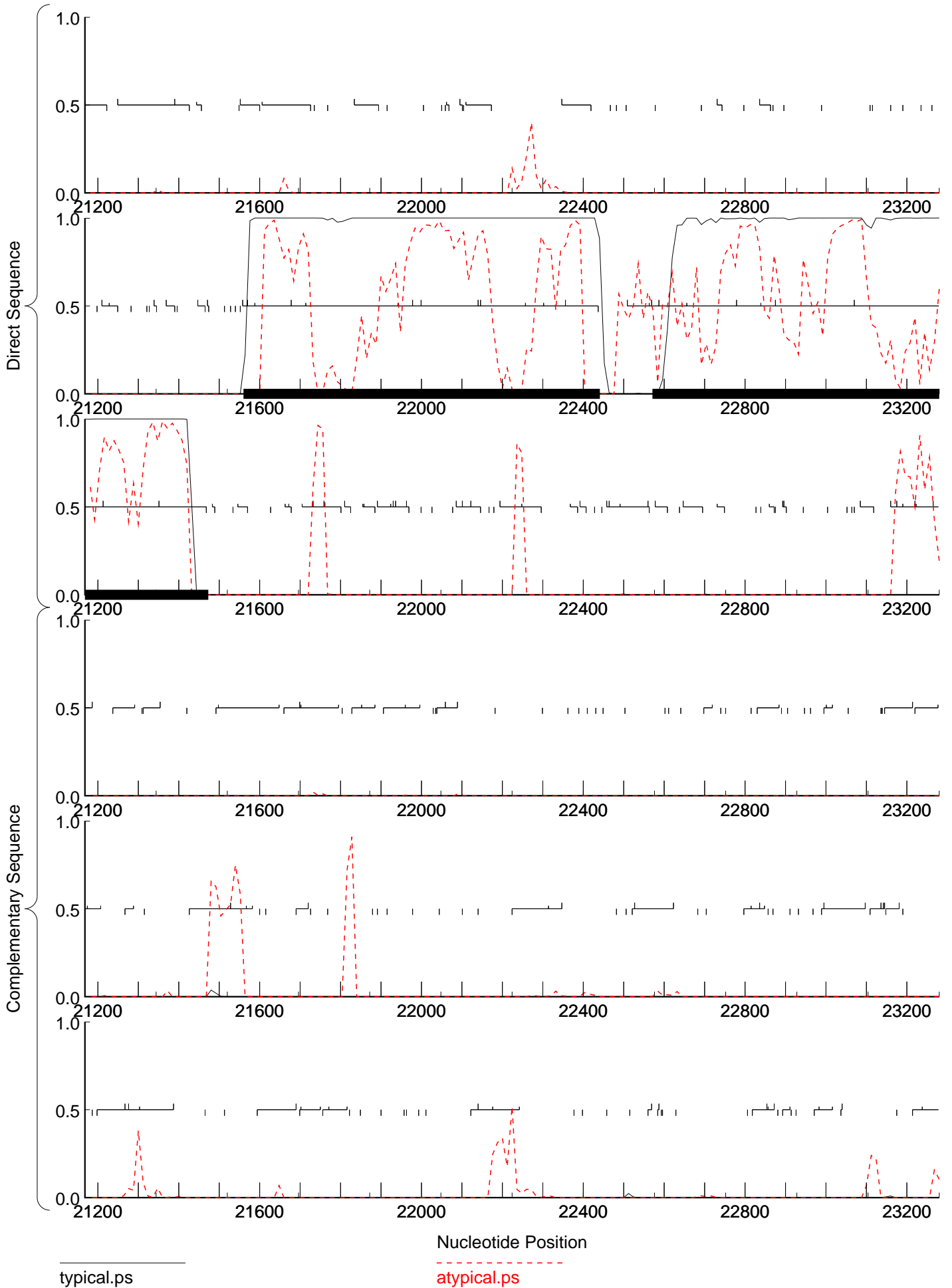
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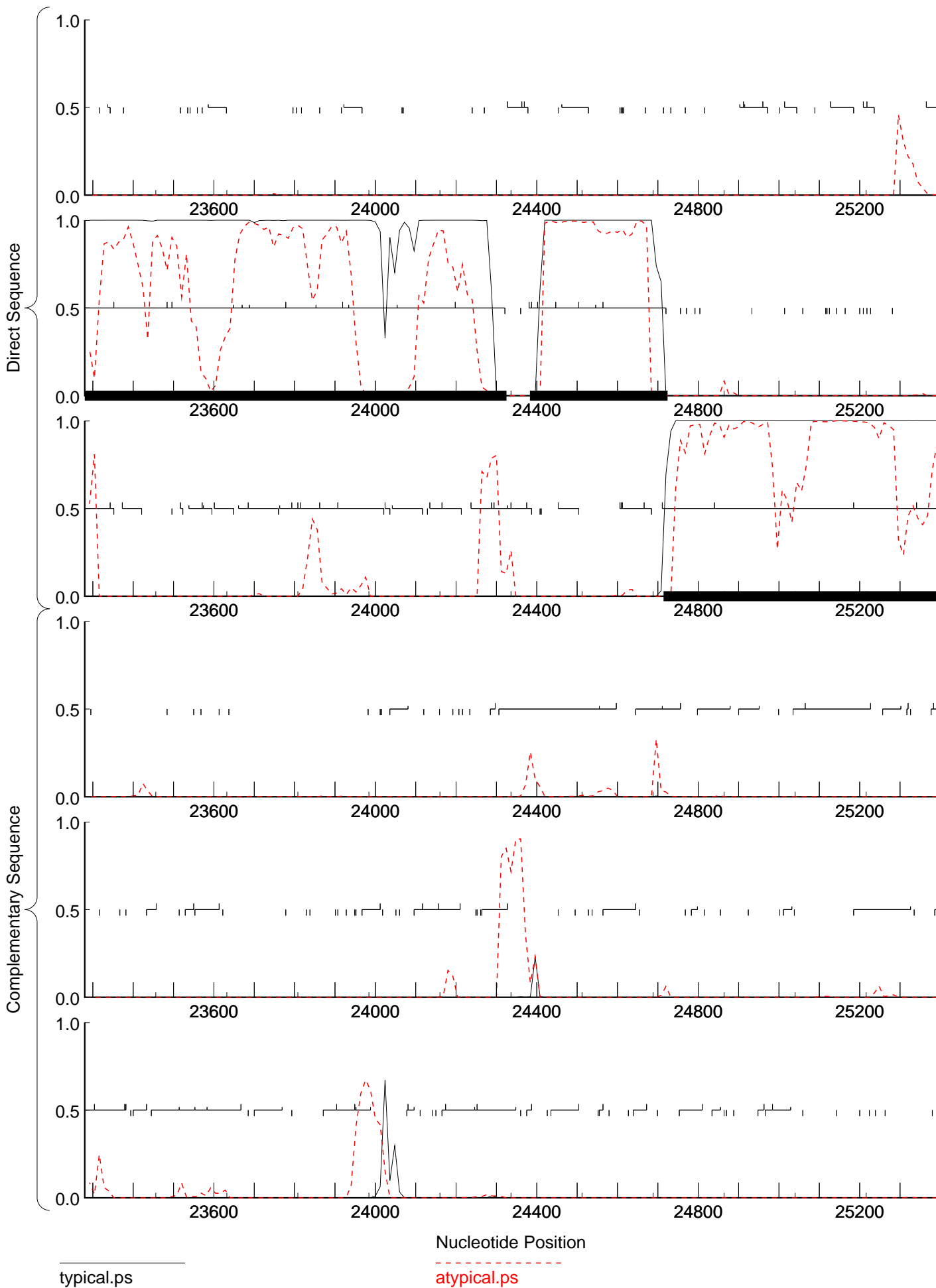
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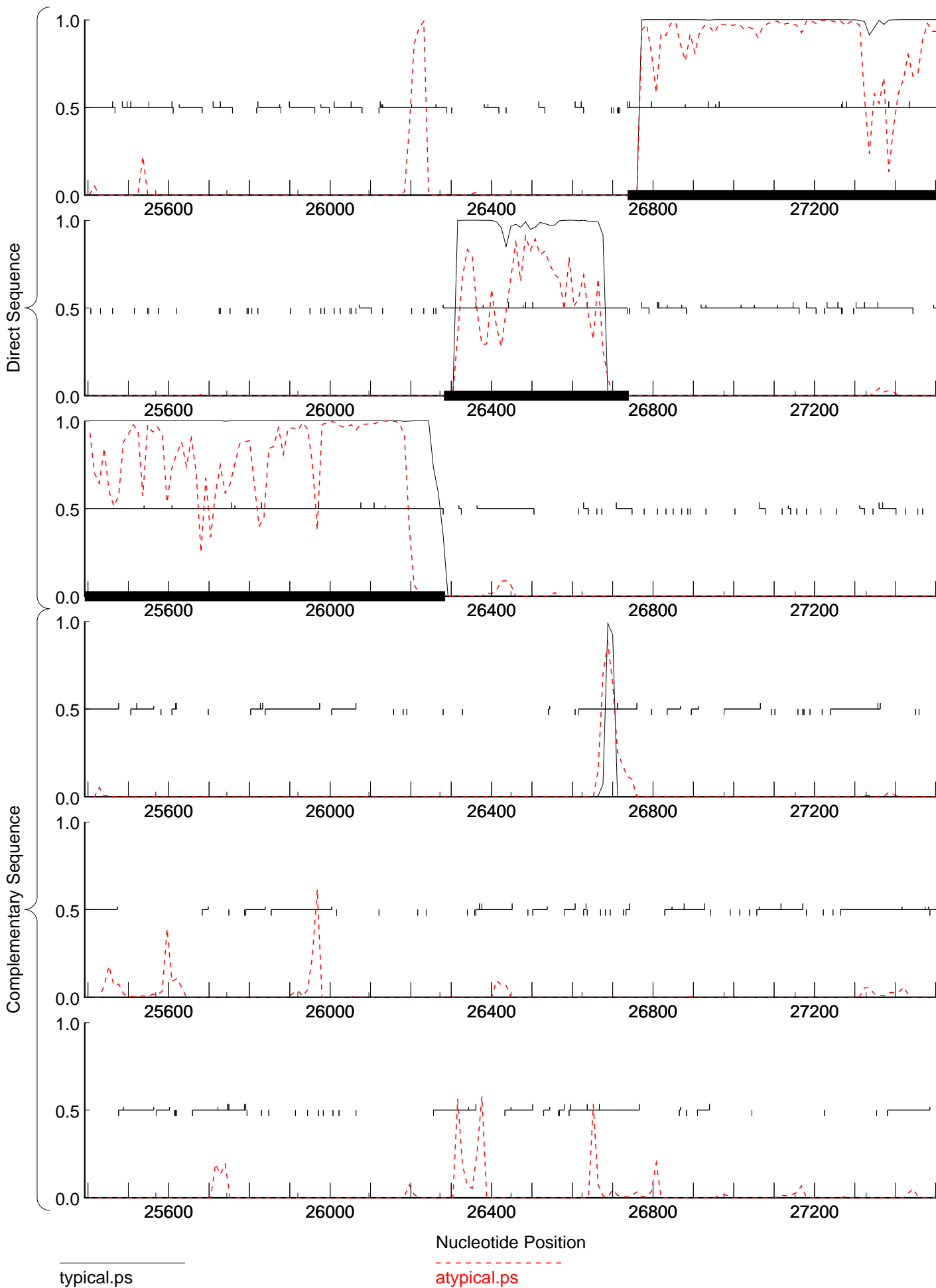


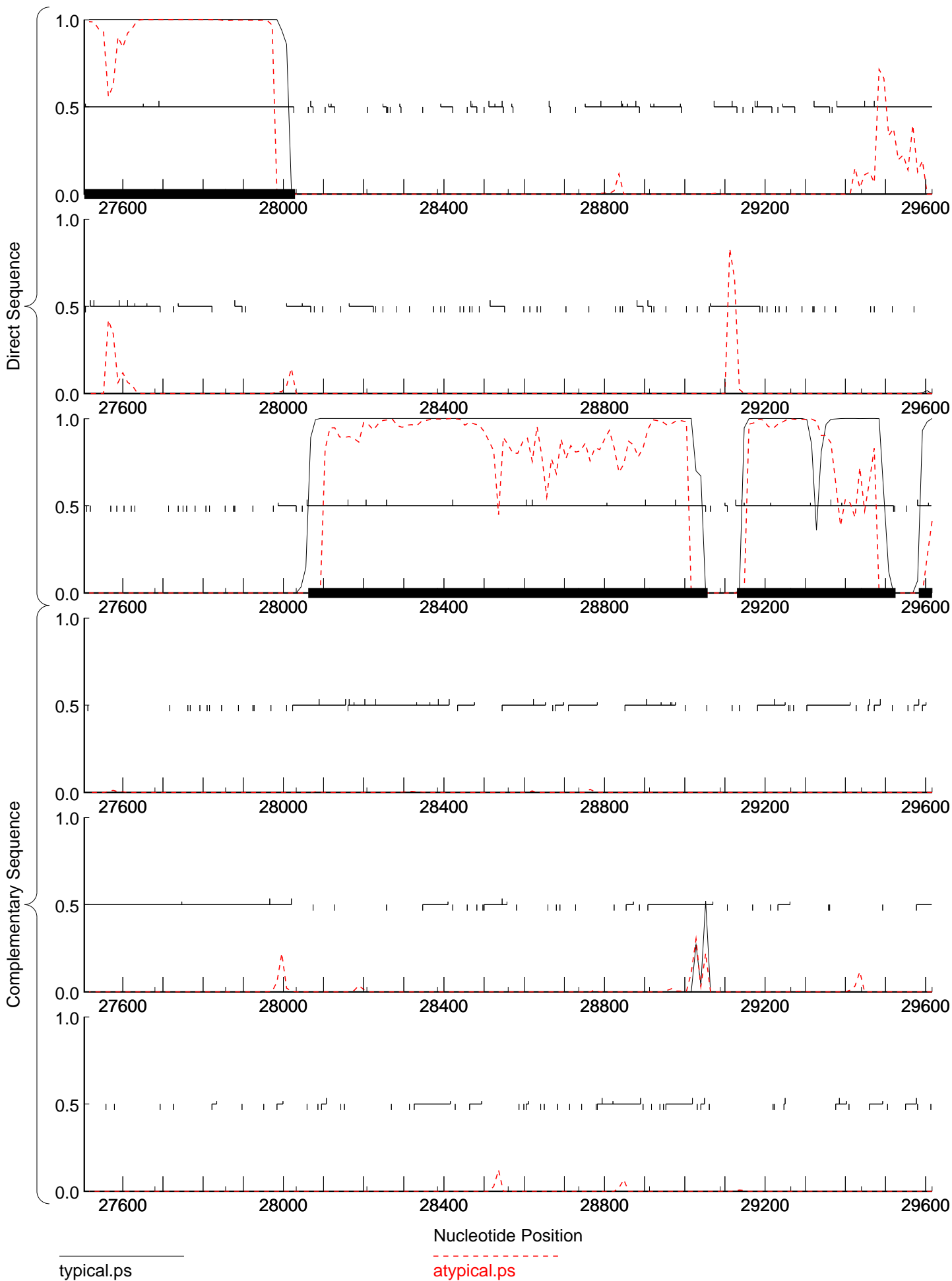






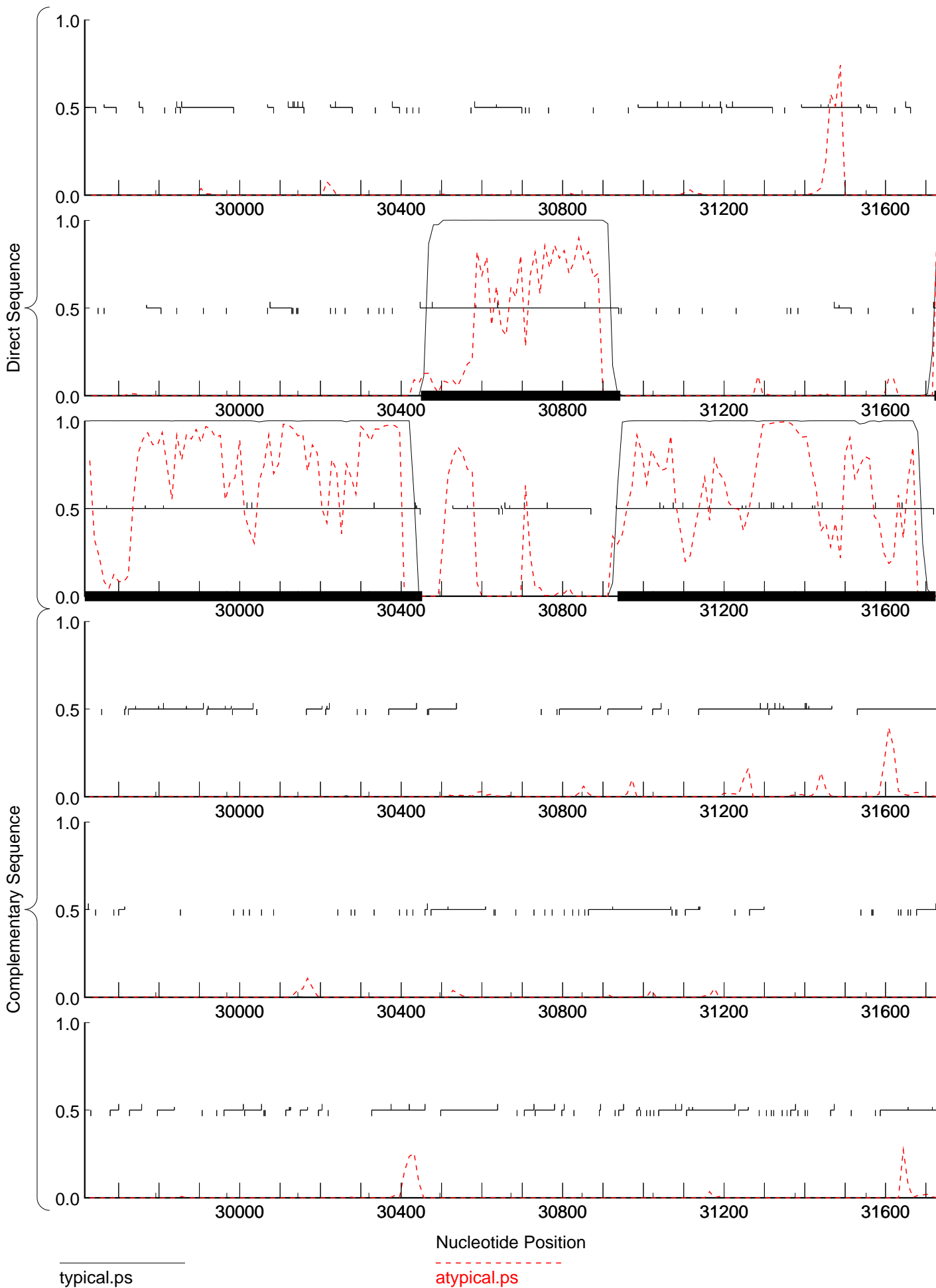






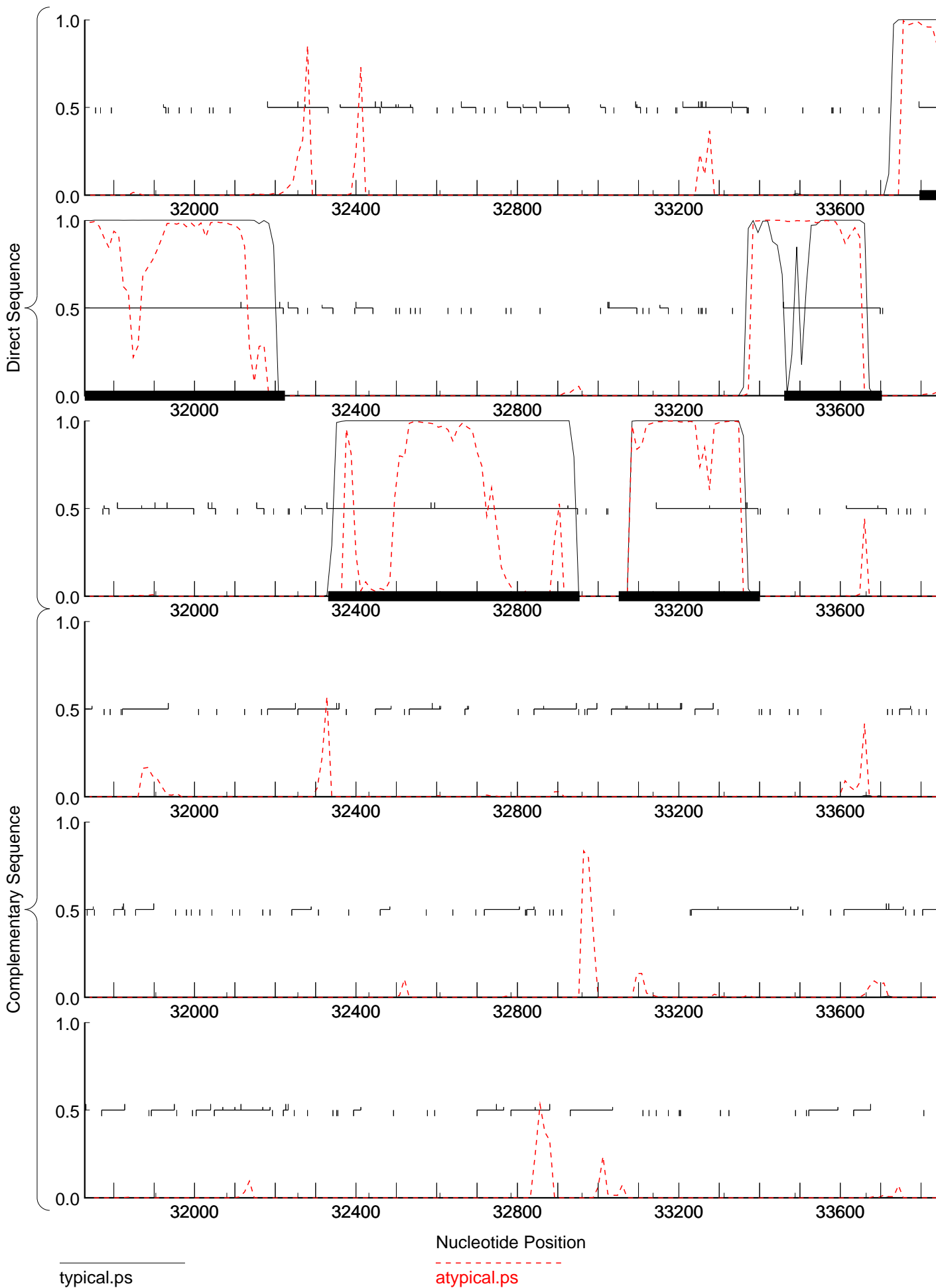
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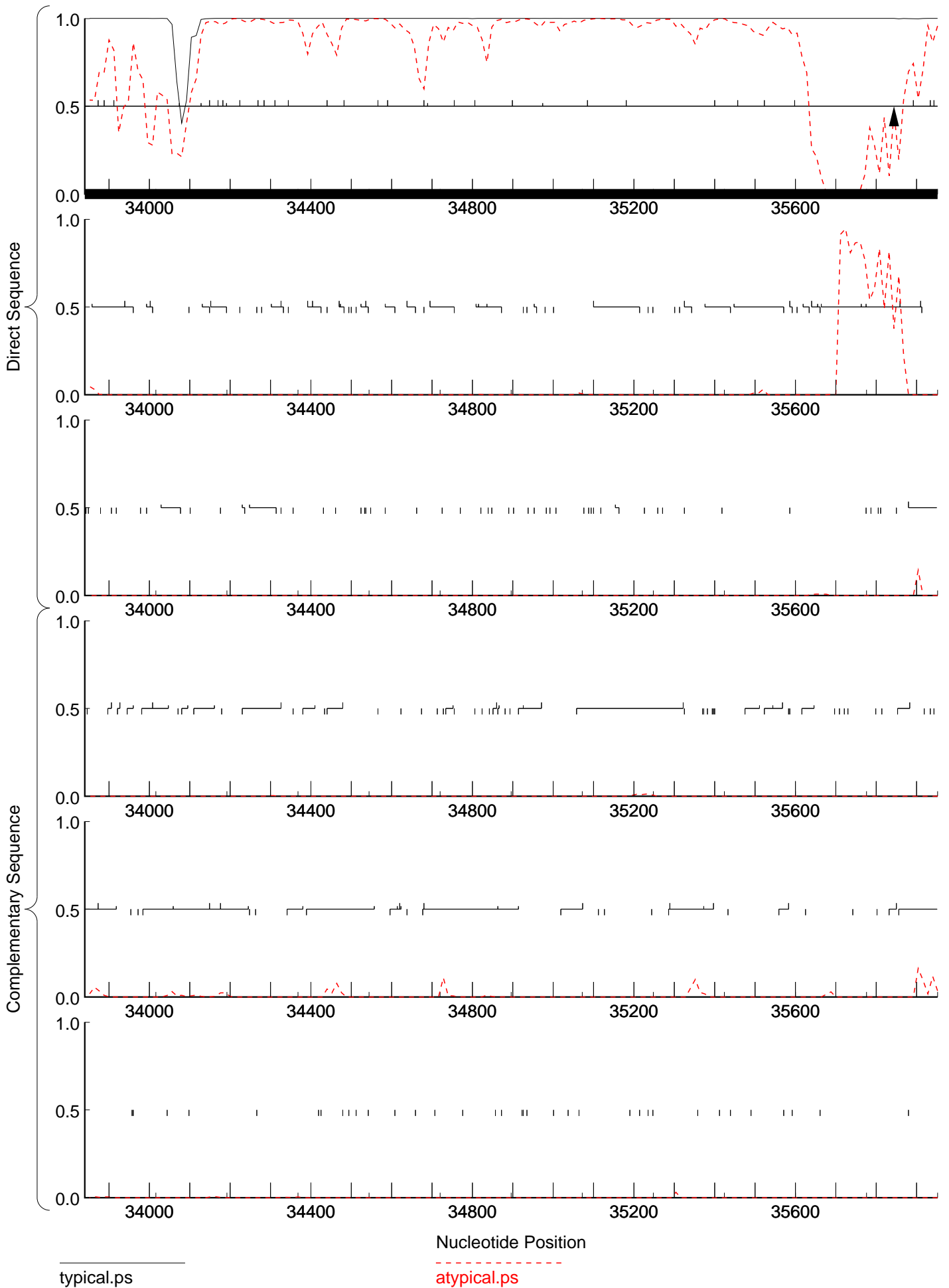
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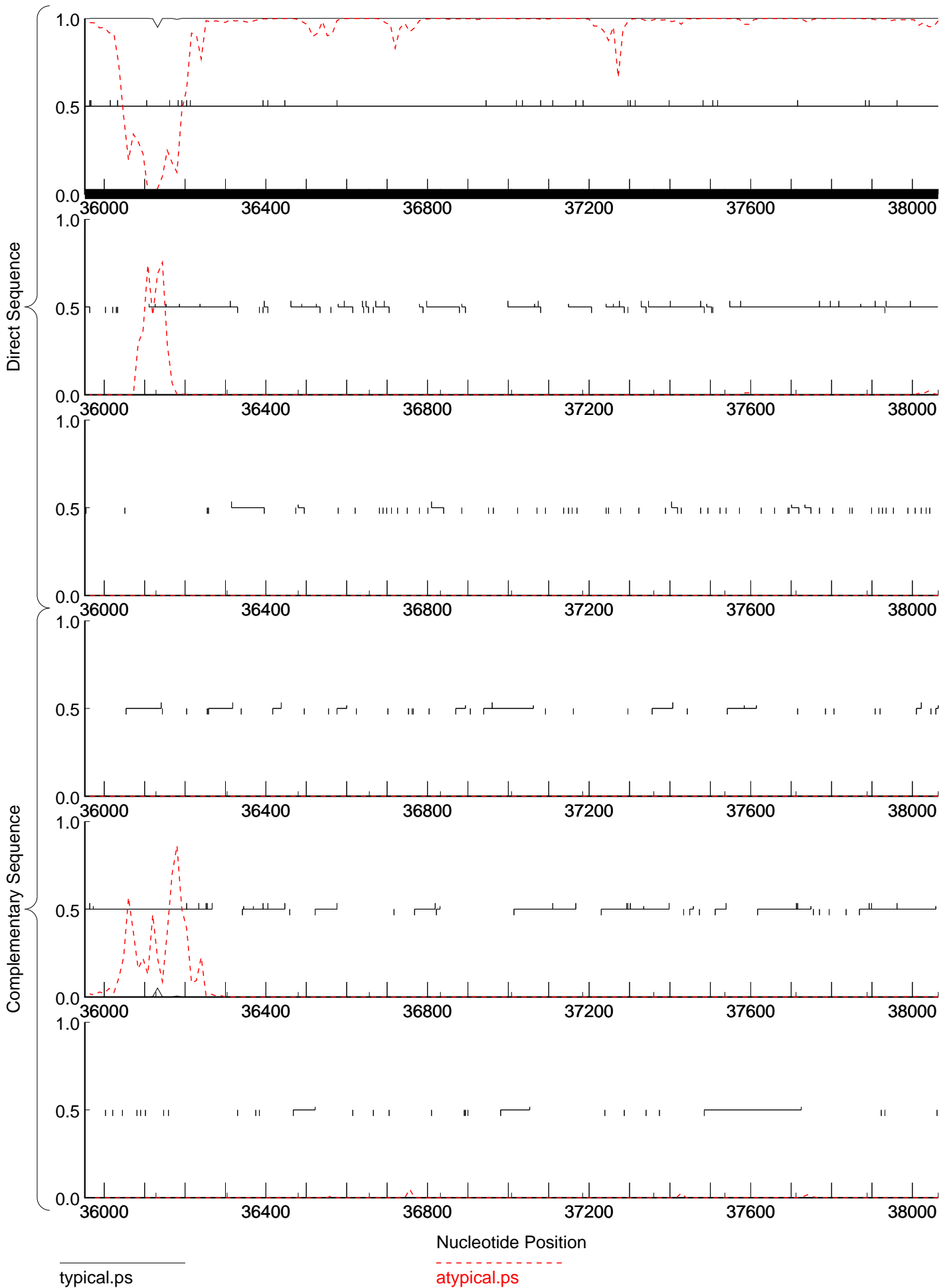


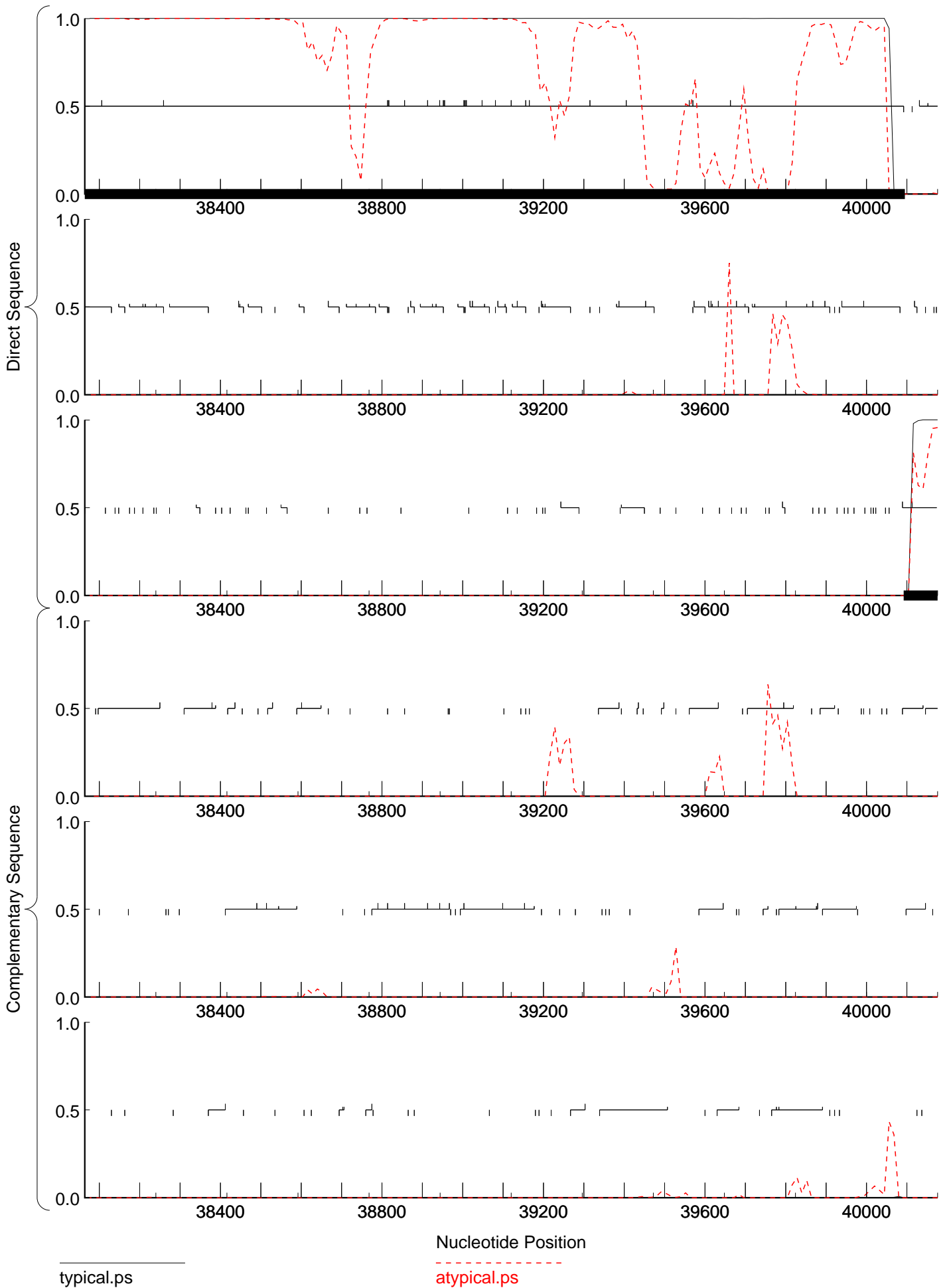
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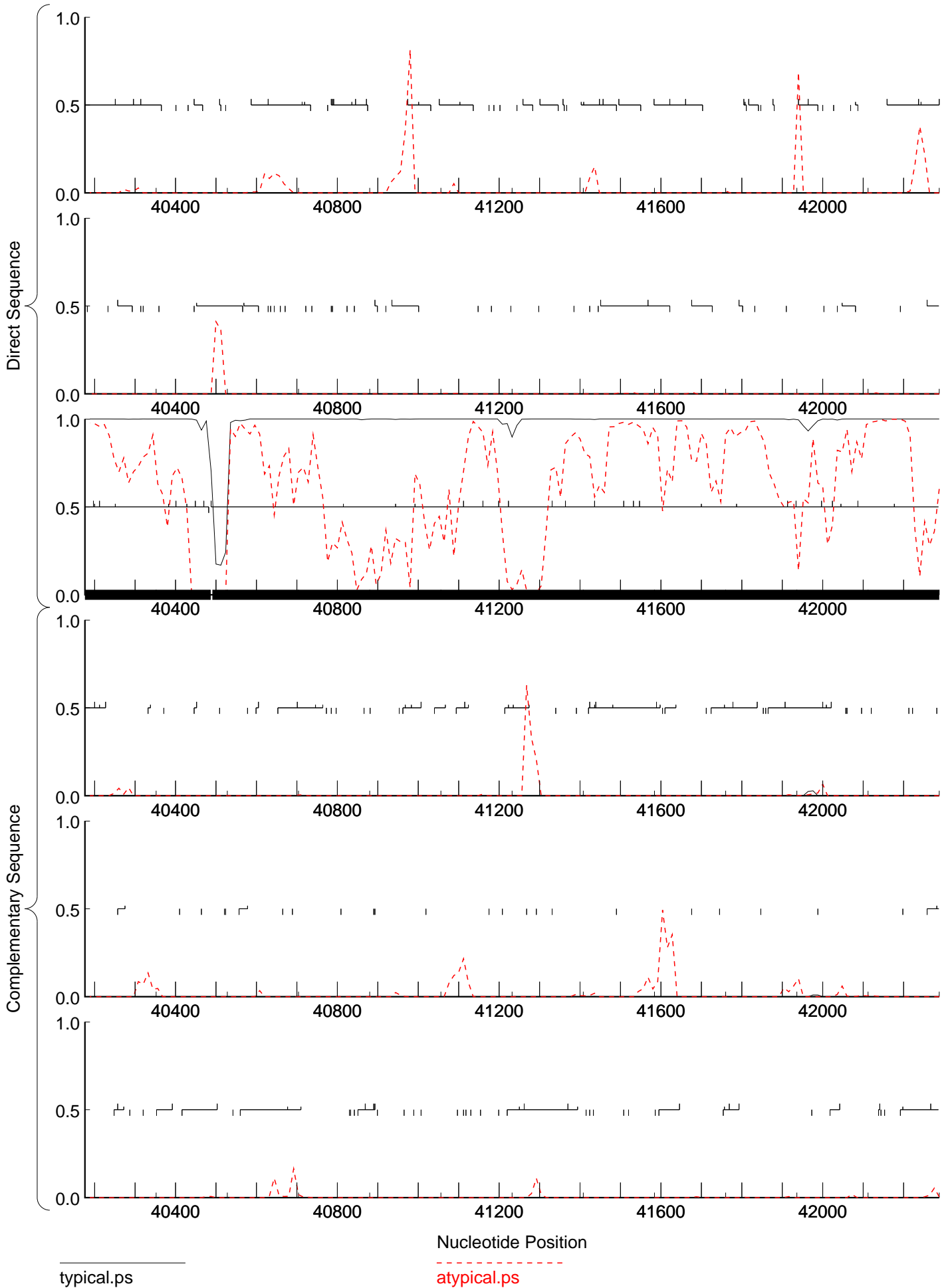
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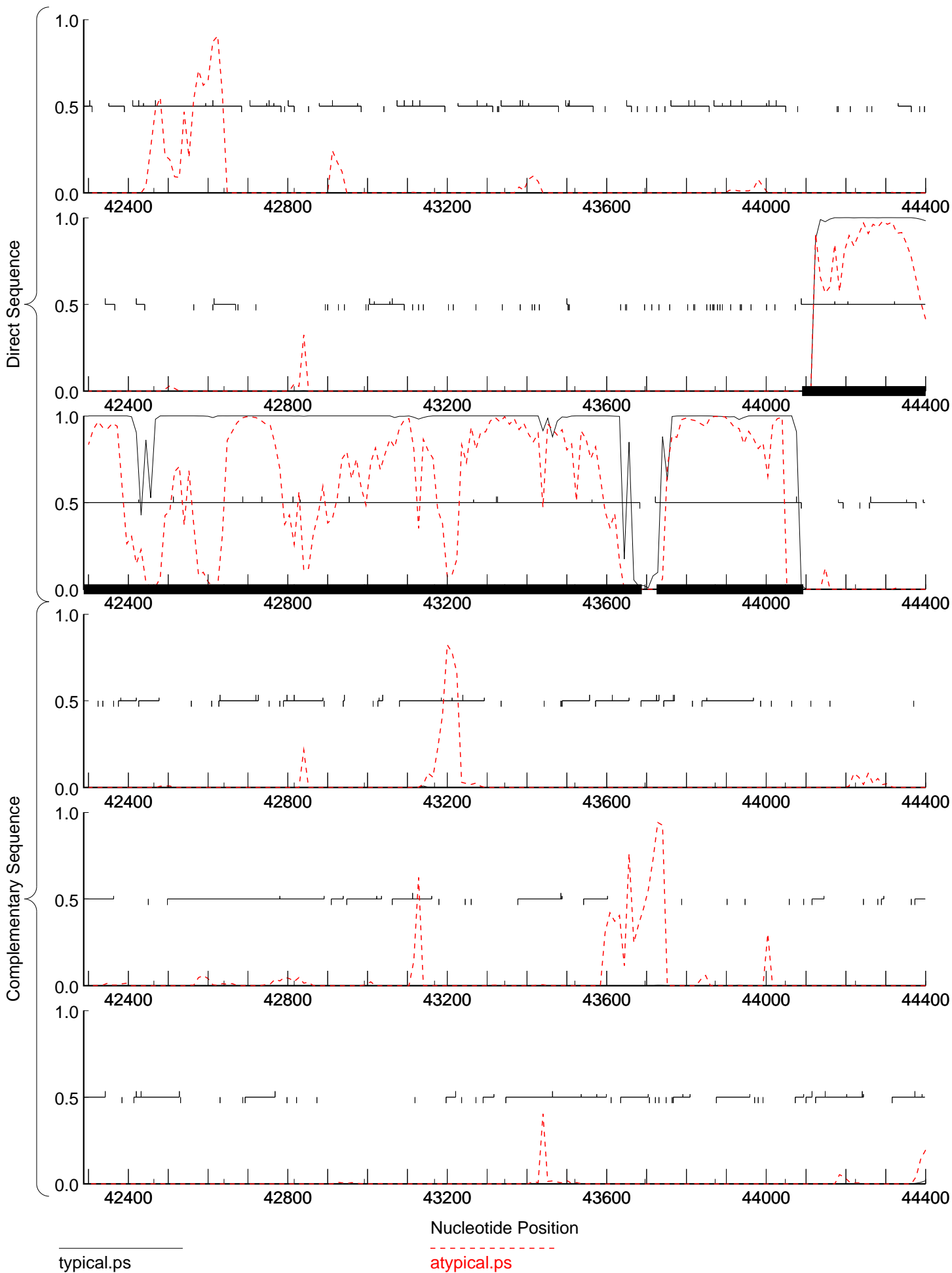






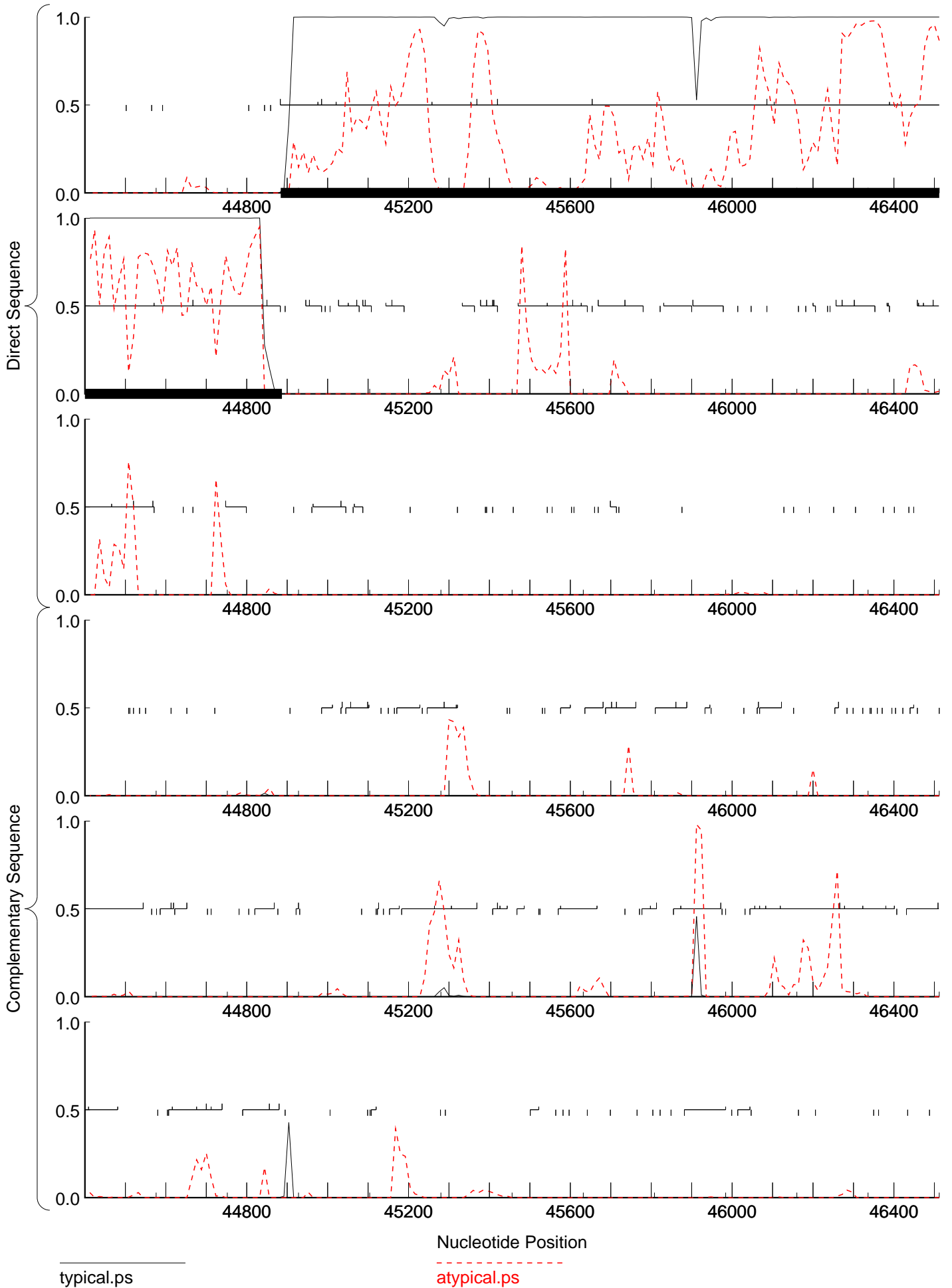
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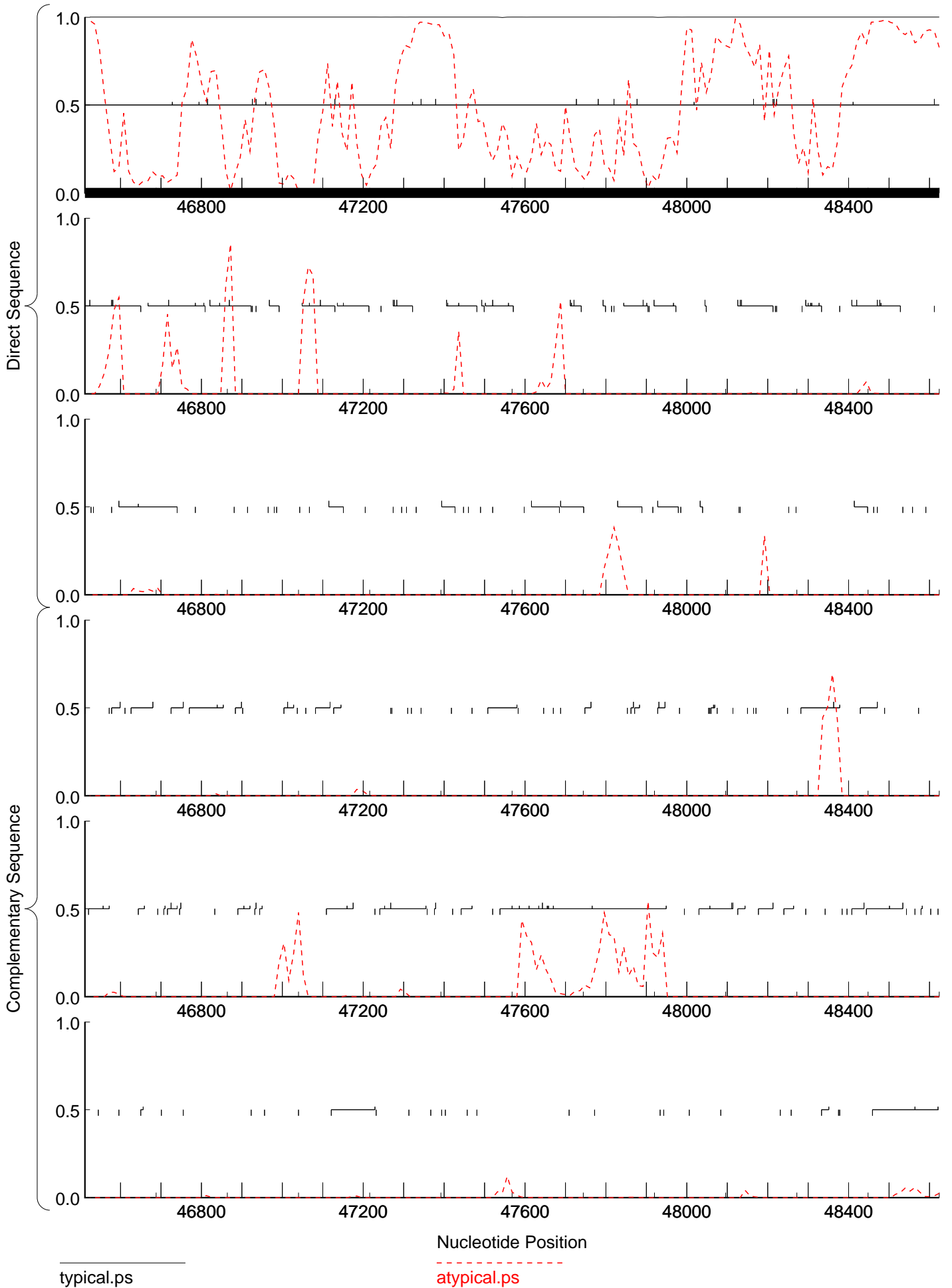
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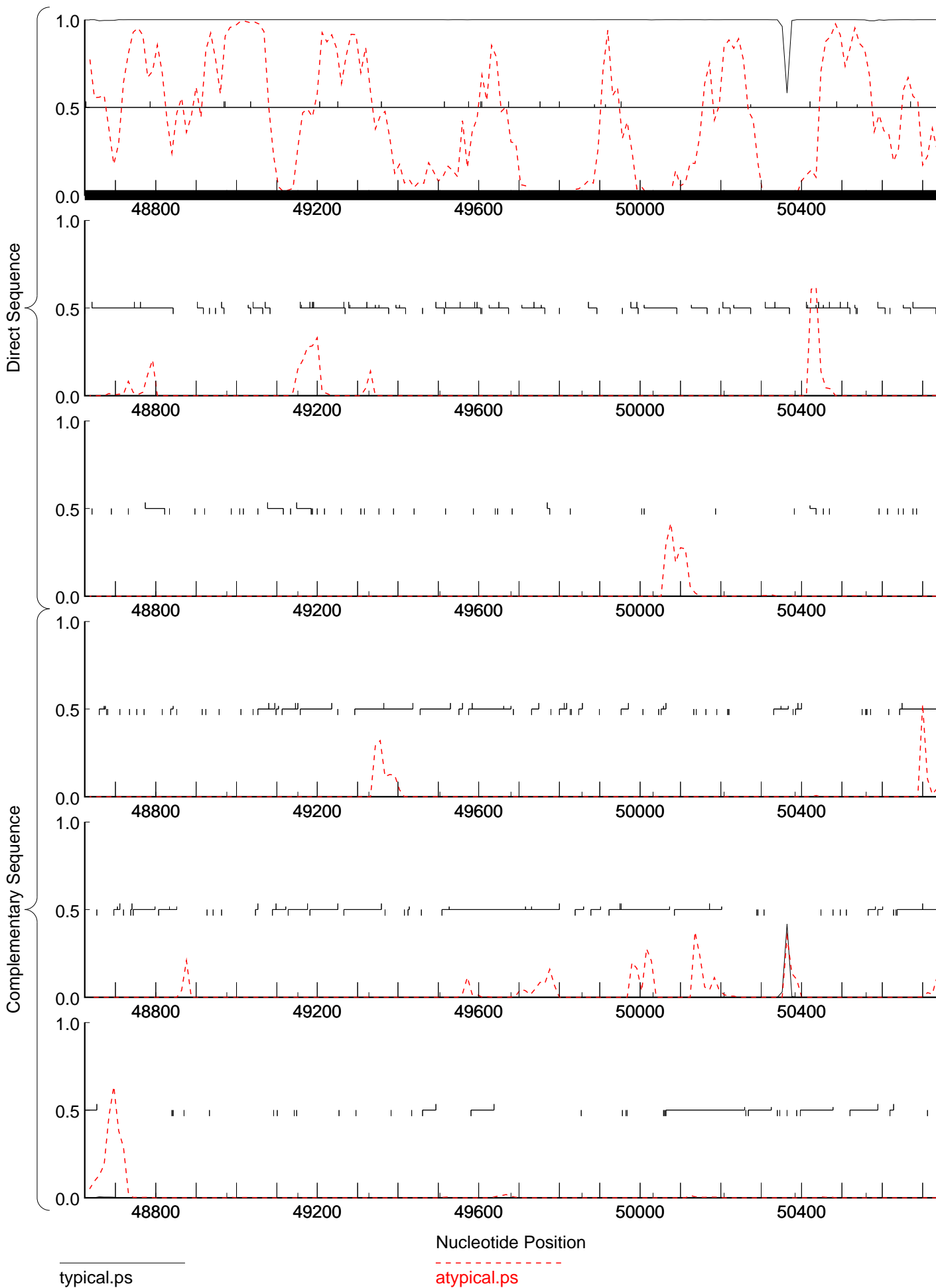


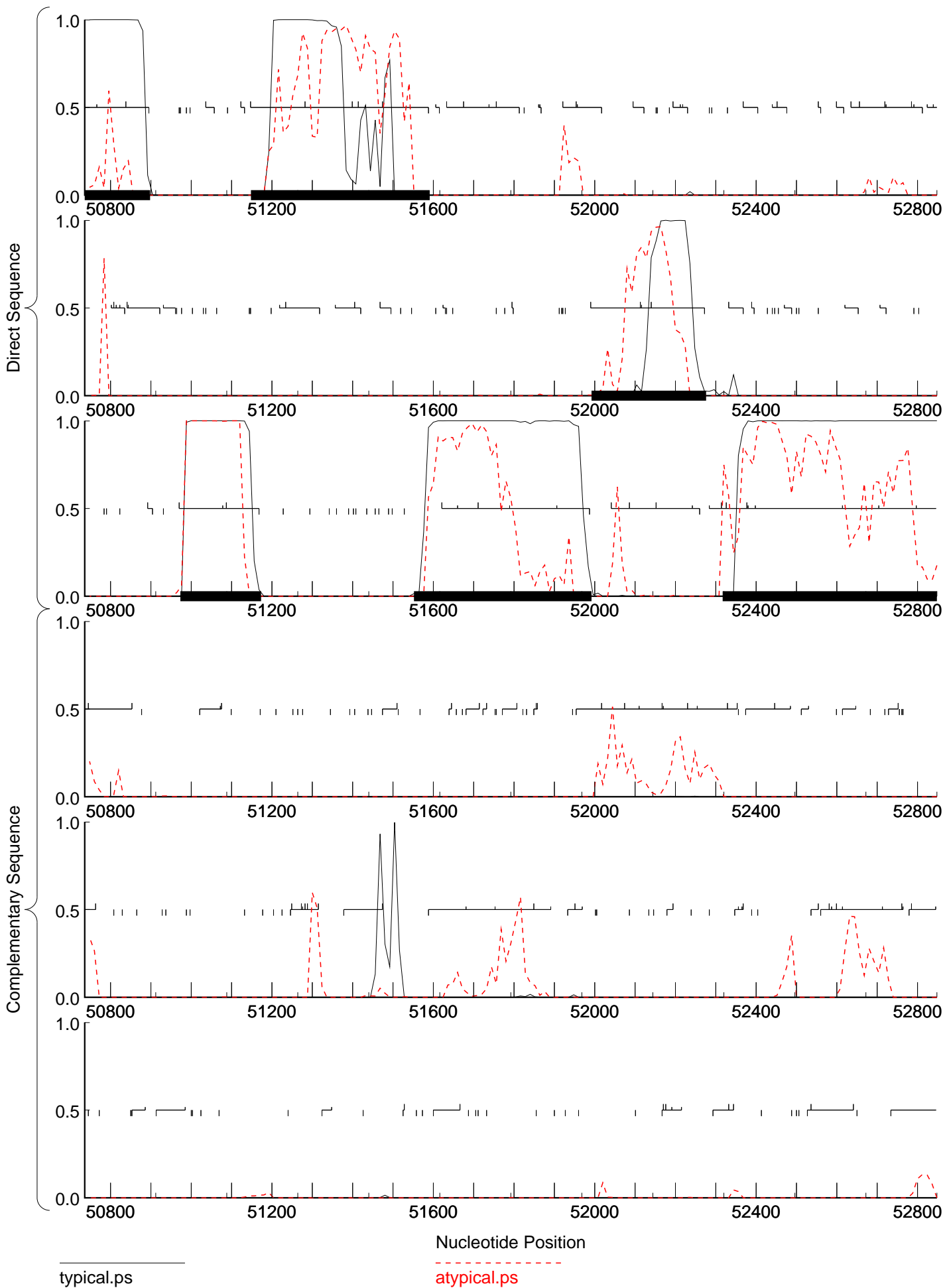
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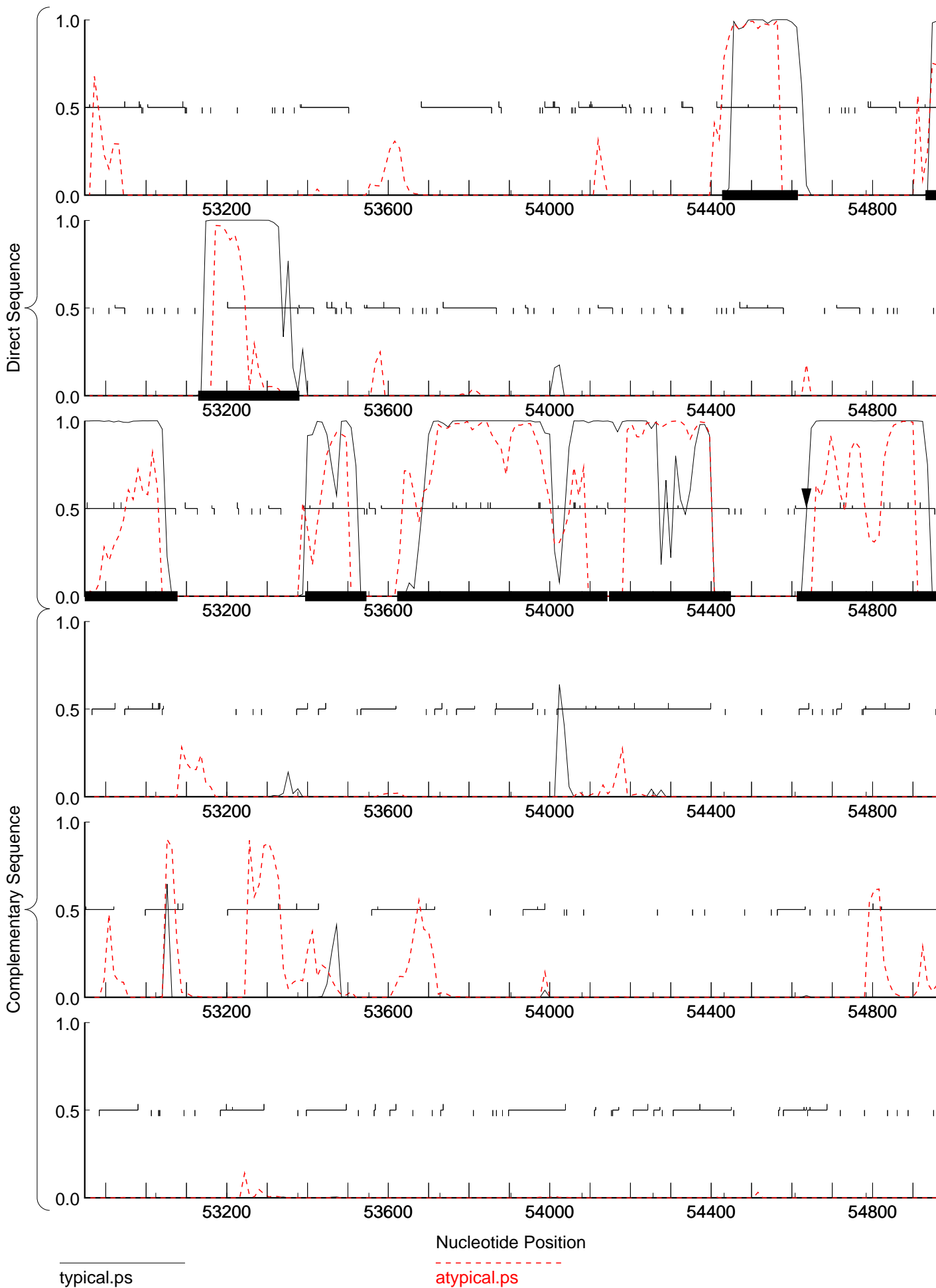
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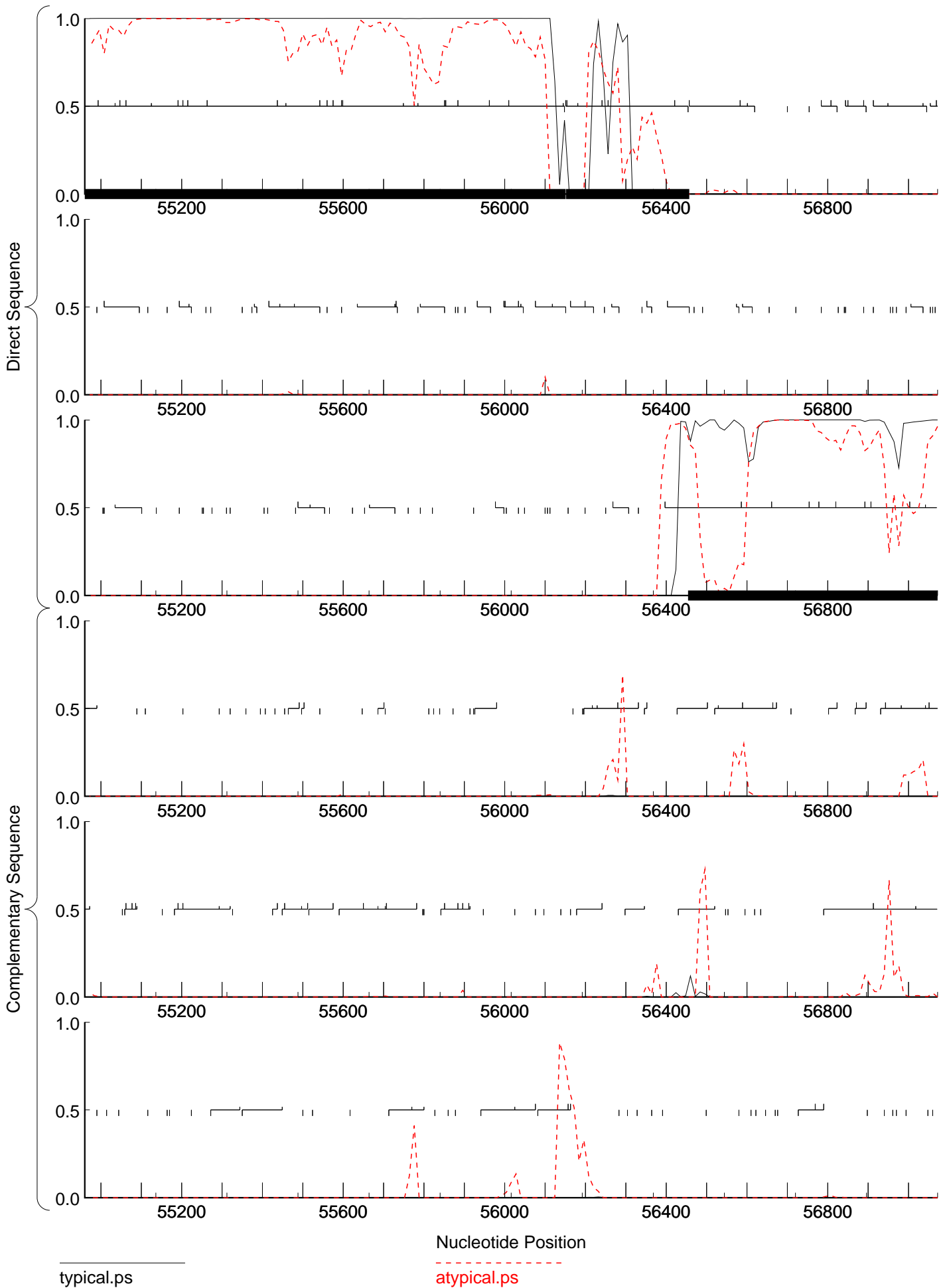


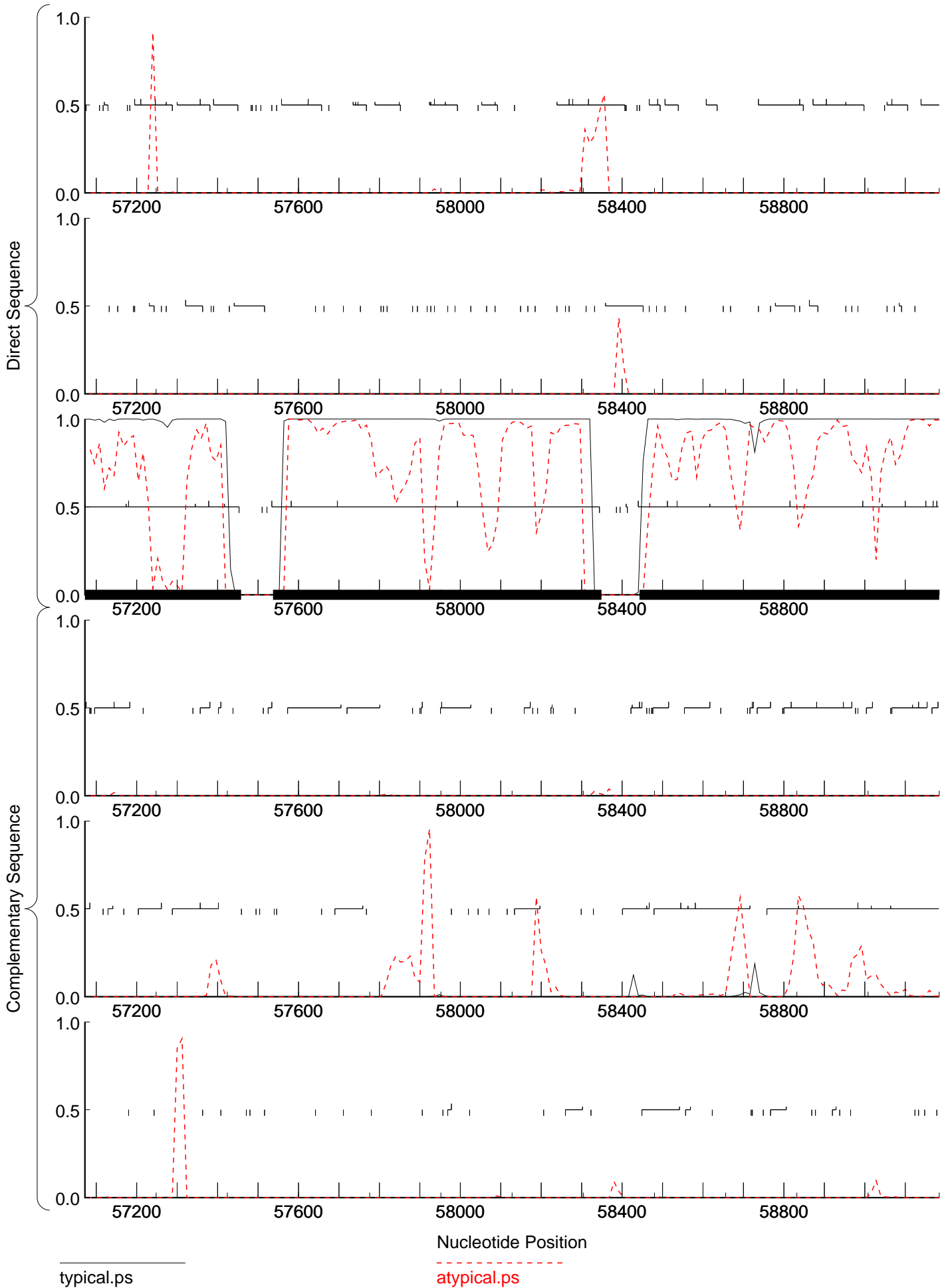


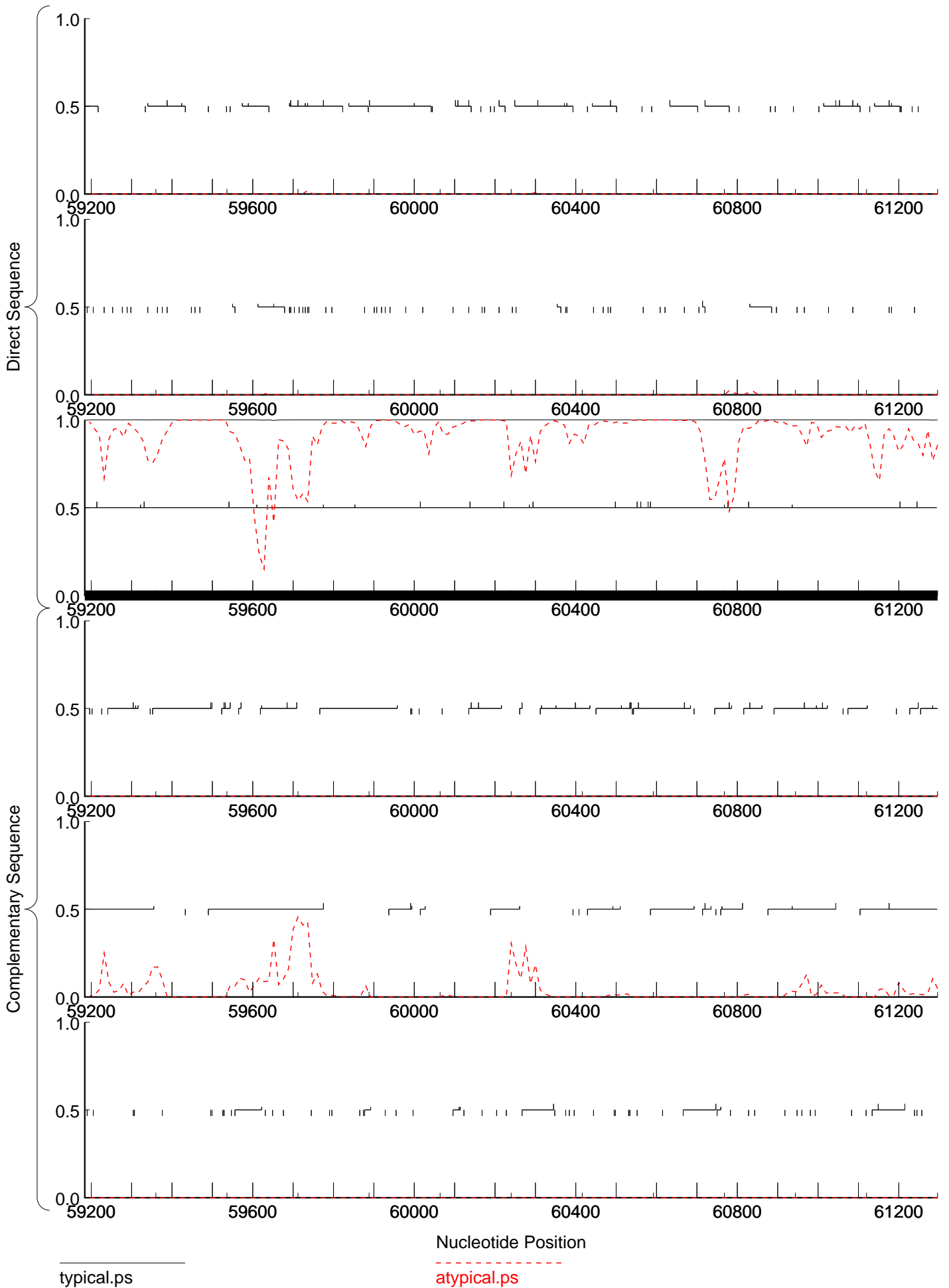




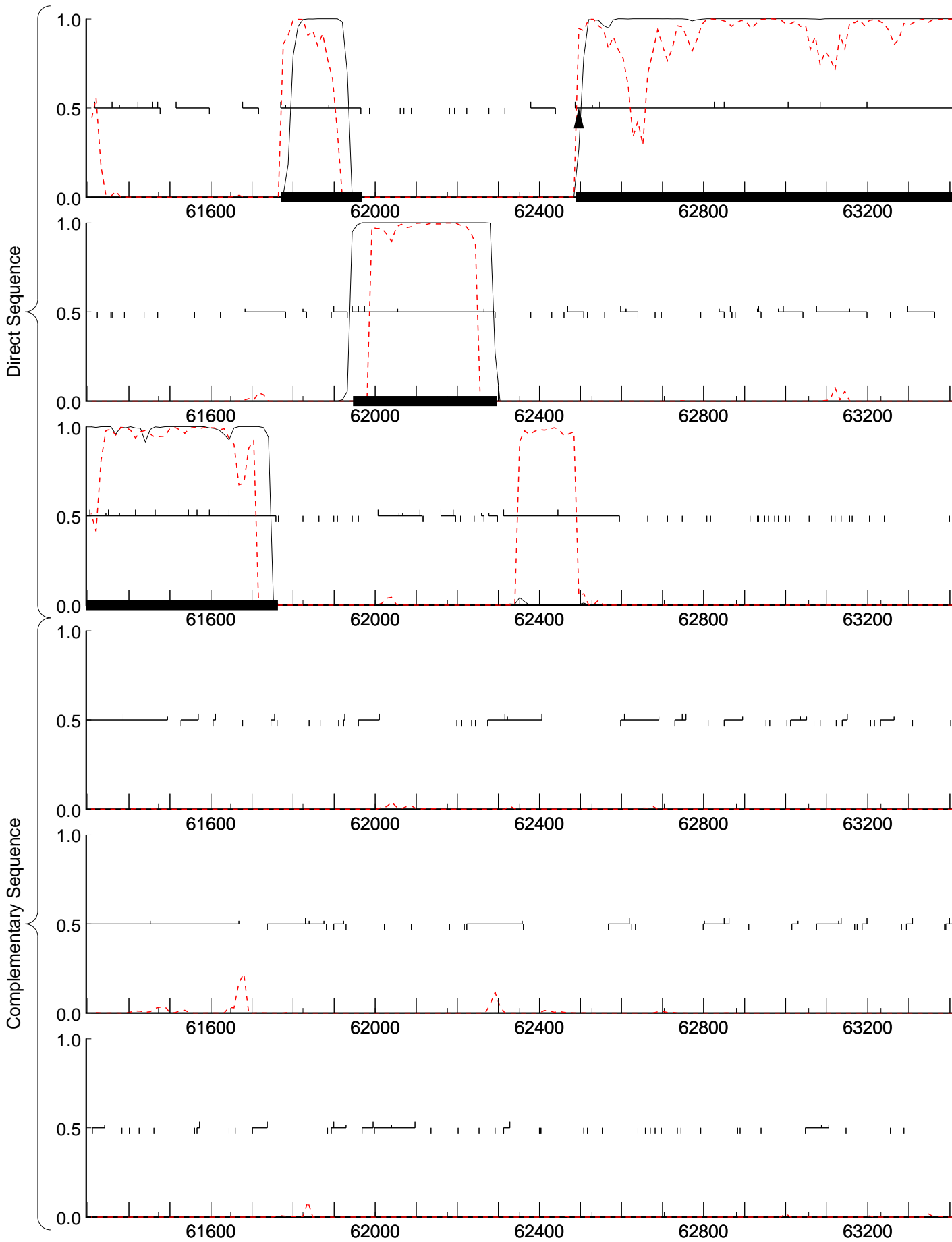








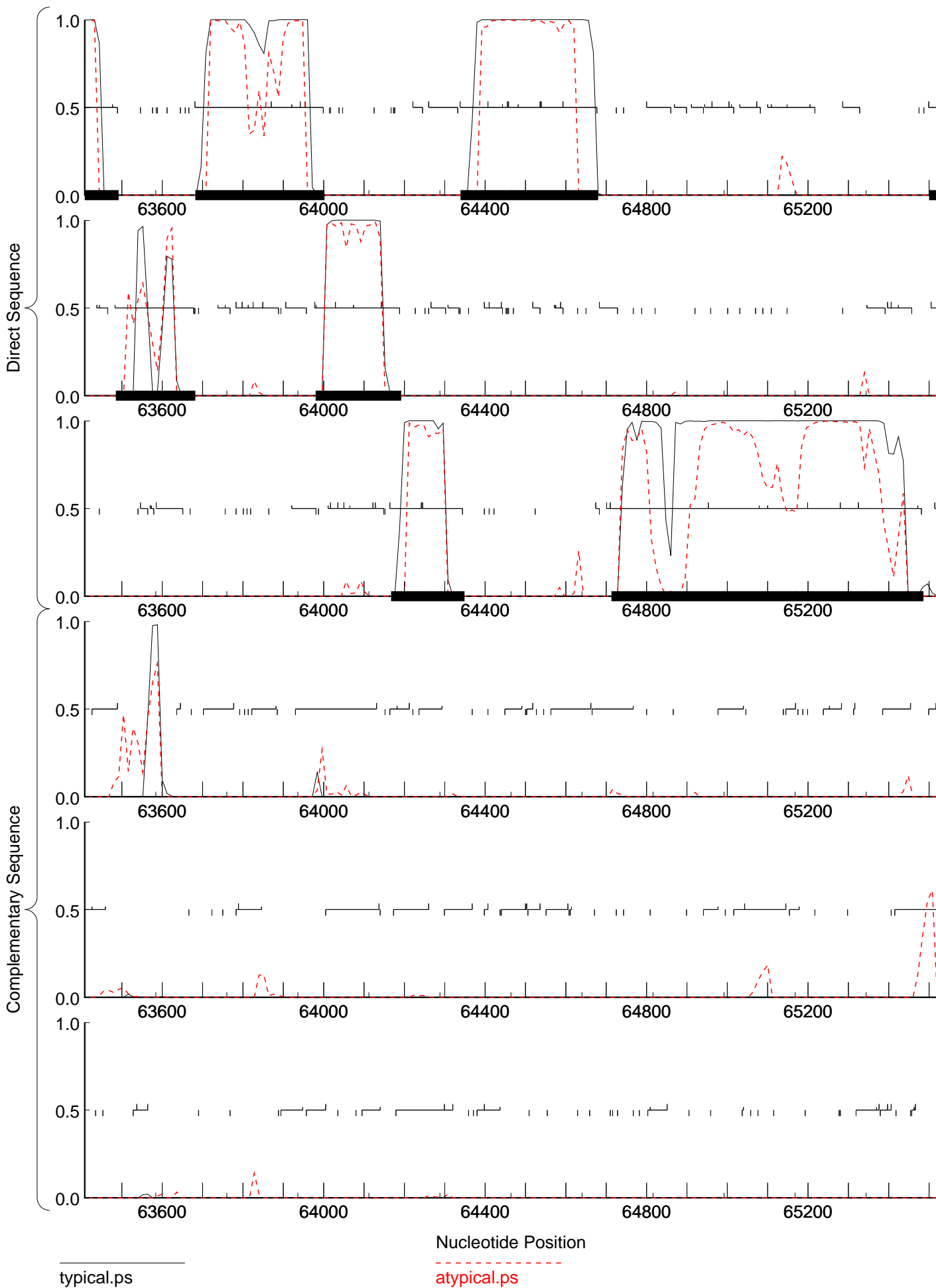
GeneMark.hmm prediction

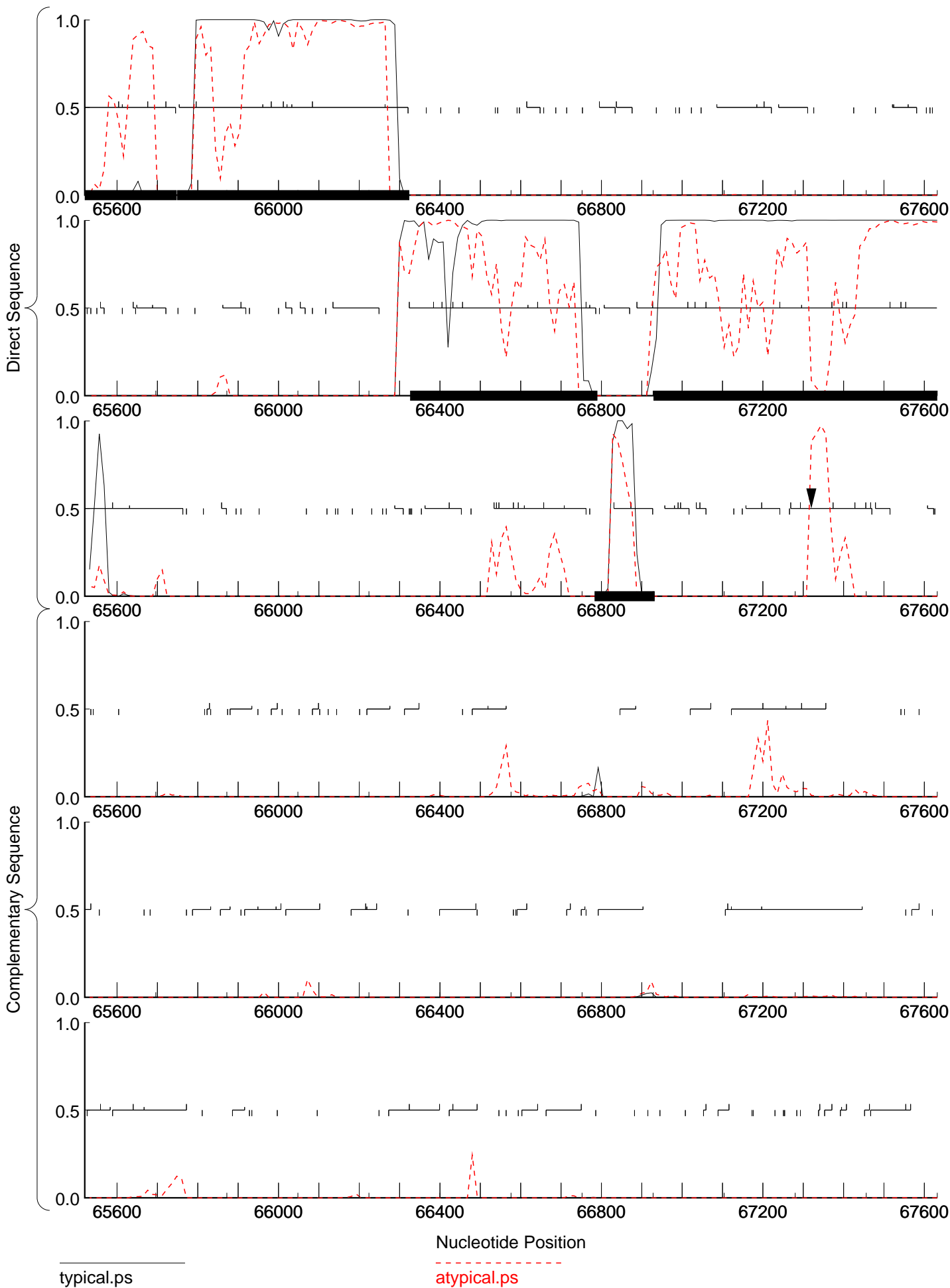


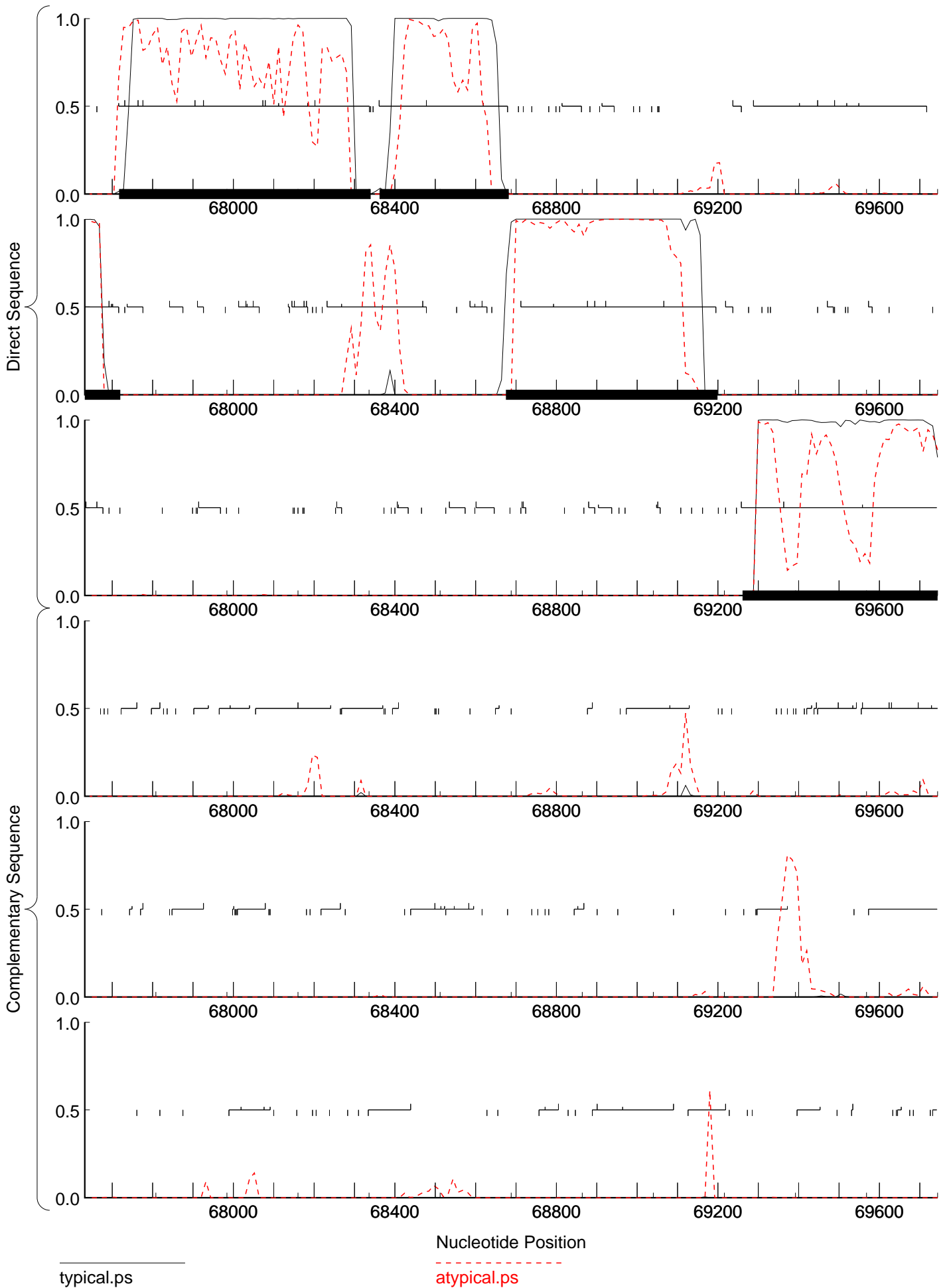
typical.ps

Nucleotide Position

atypical.ps

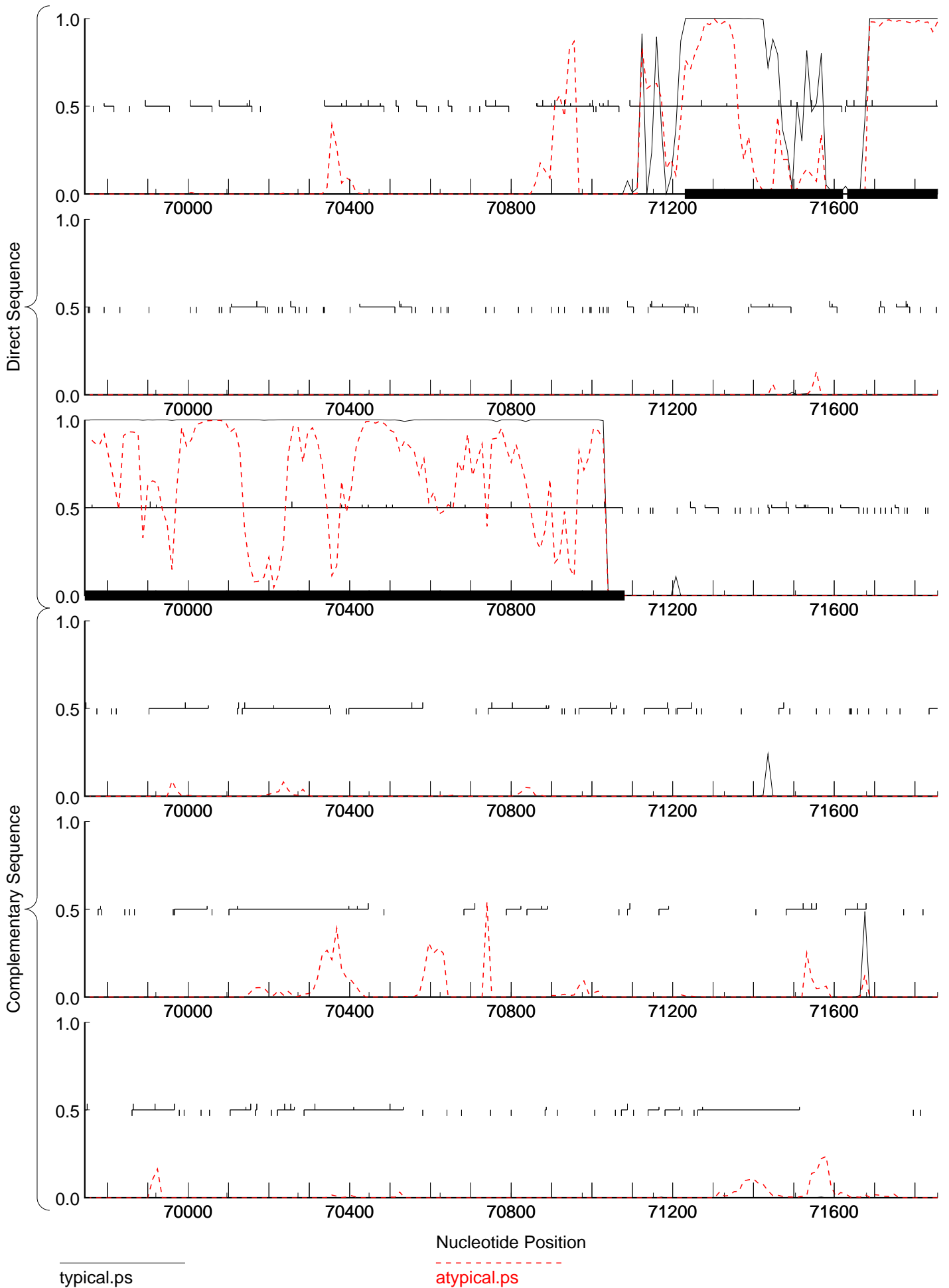


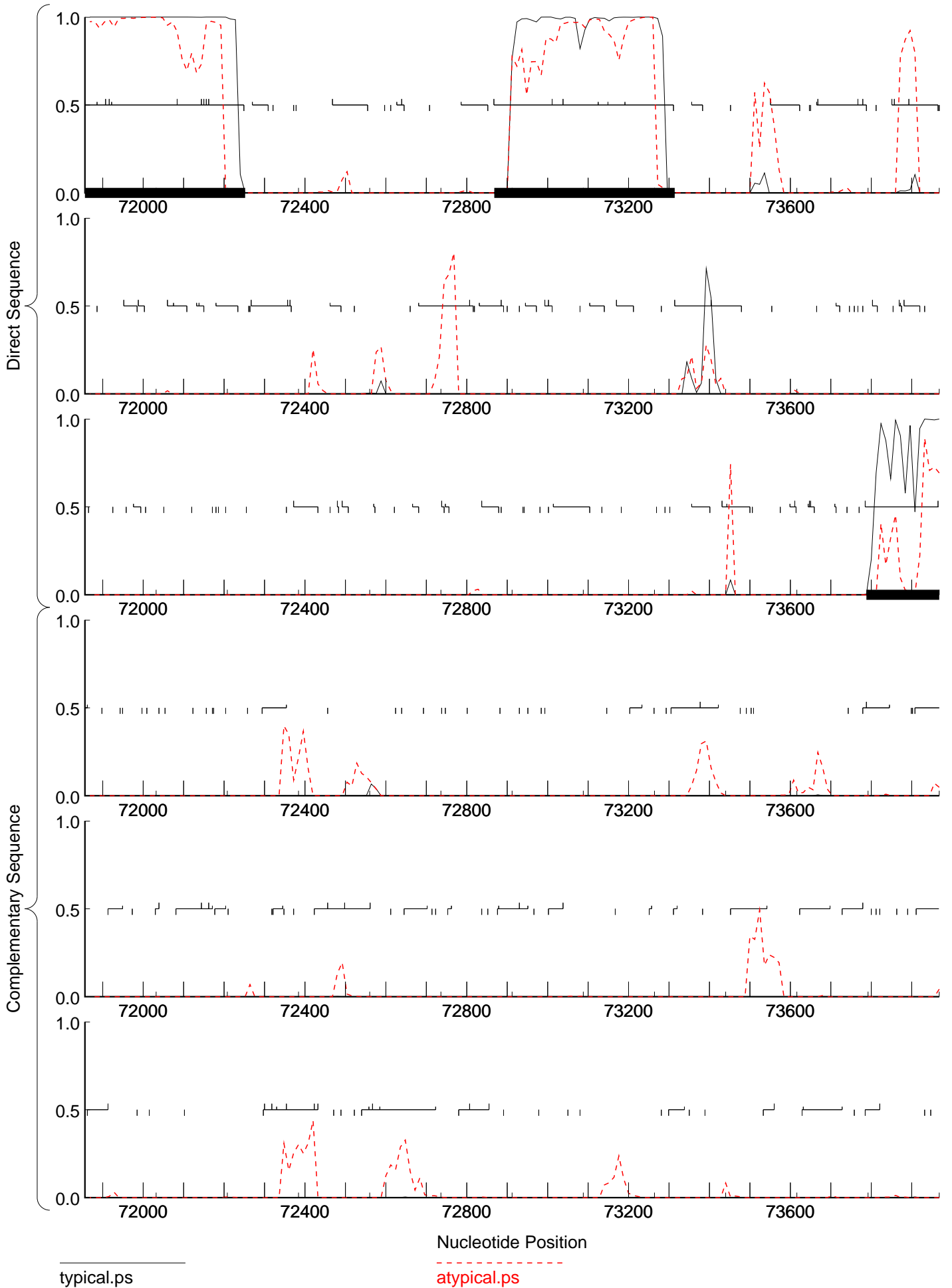




typical.ps

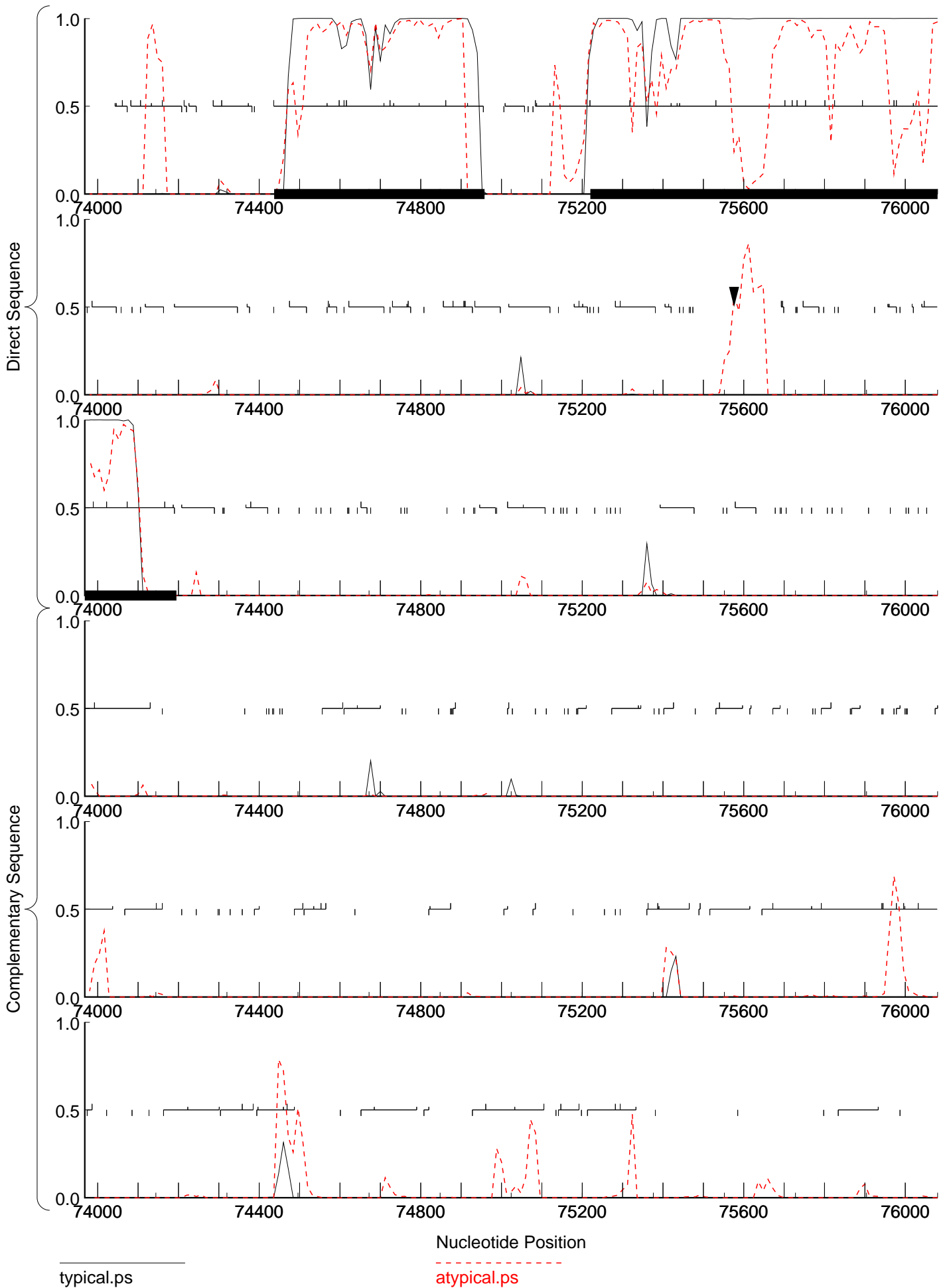
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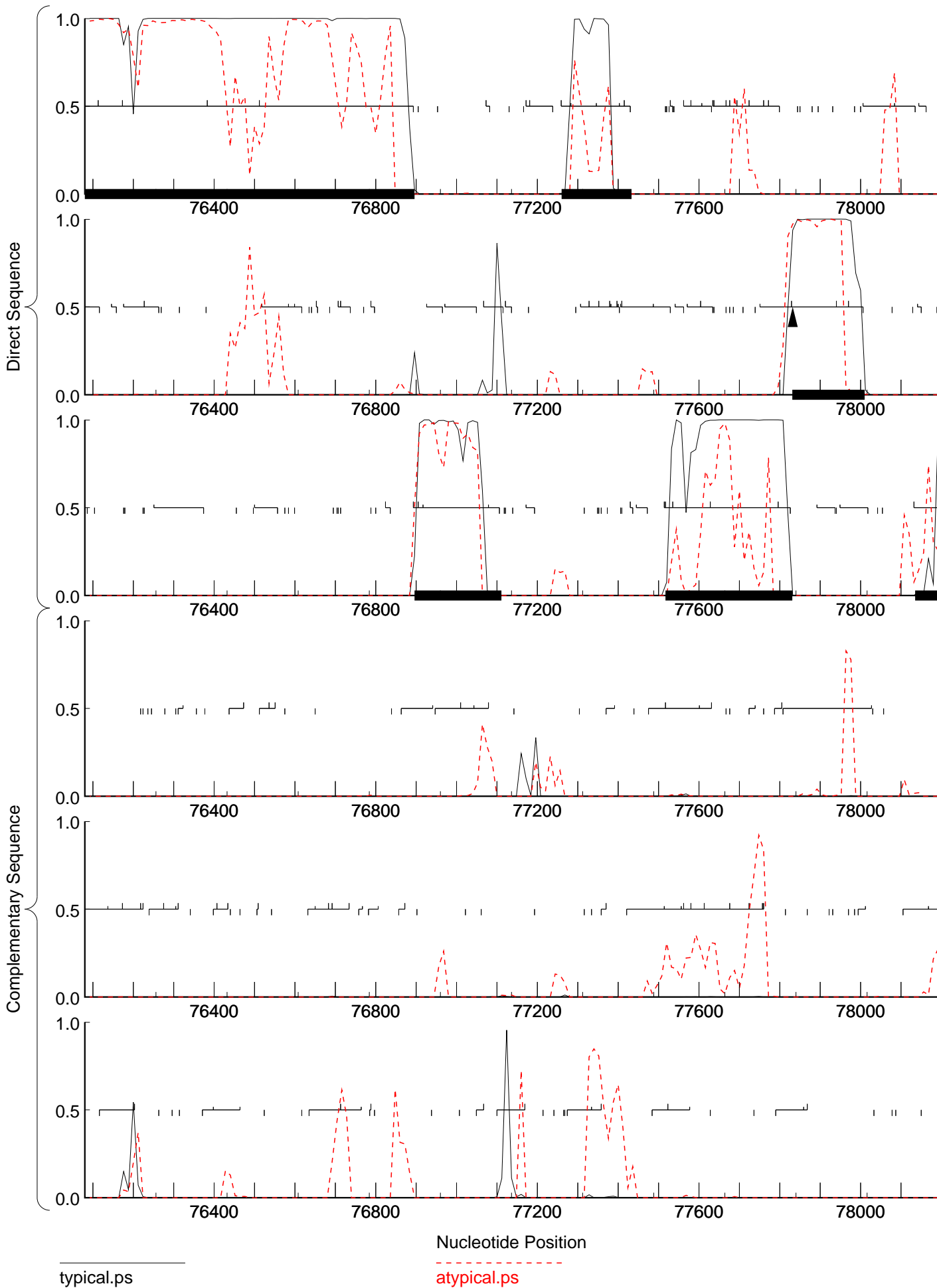
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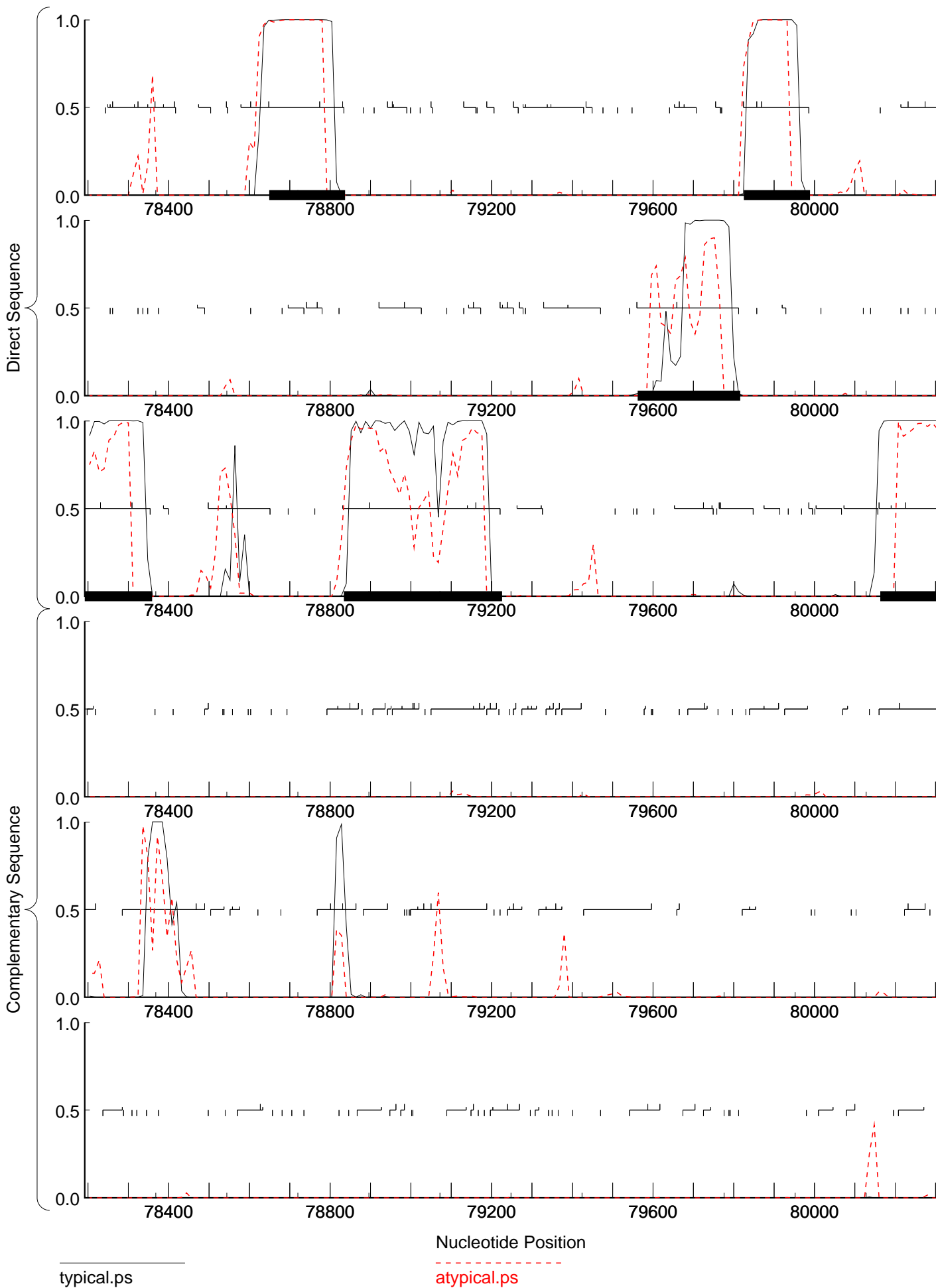
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typical.ps

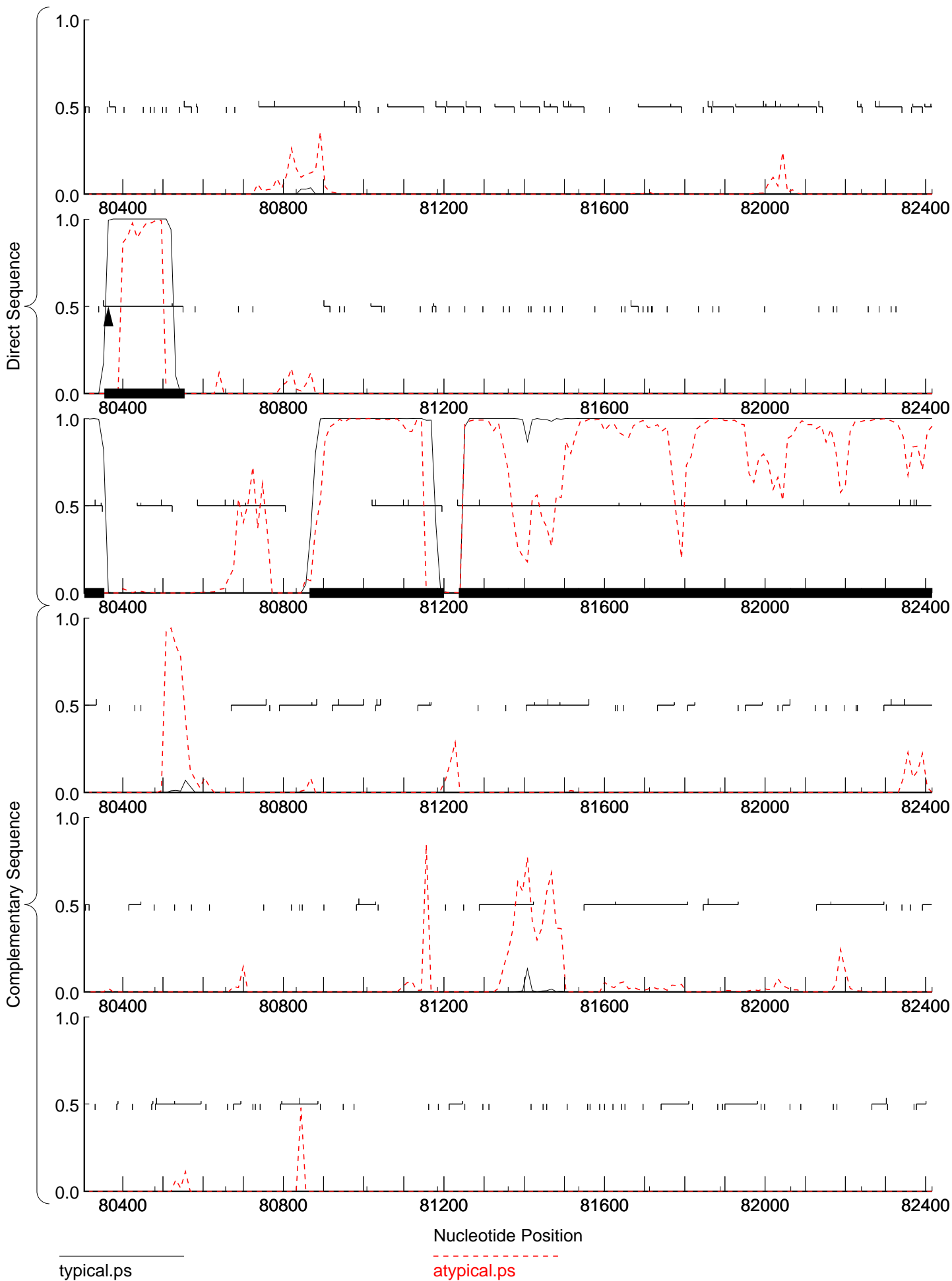
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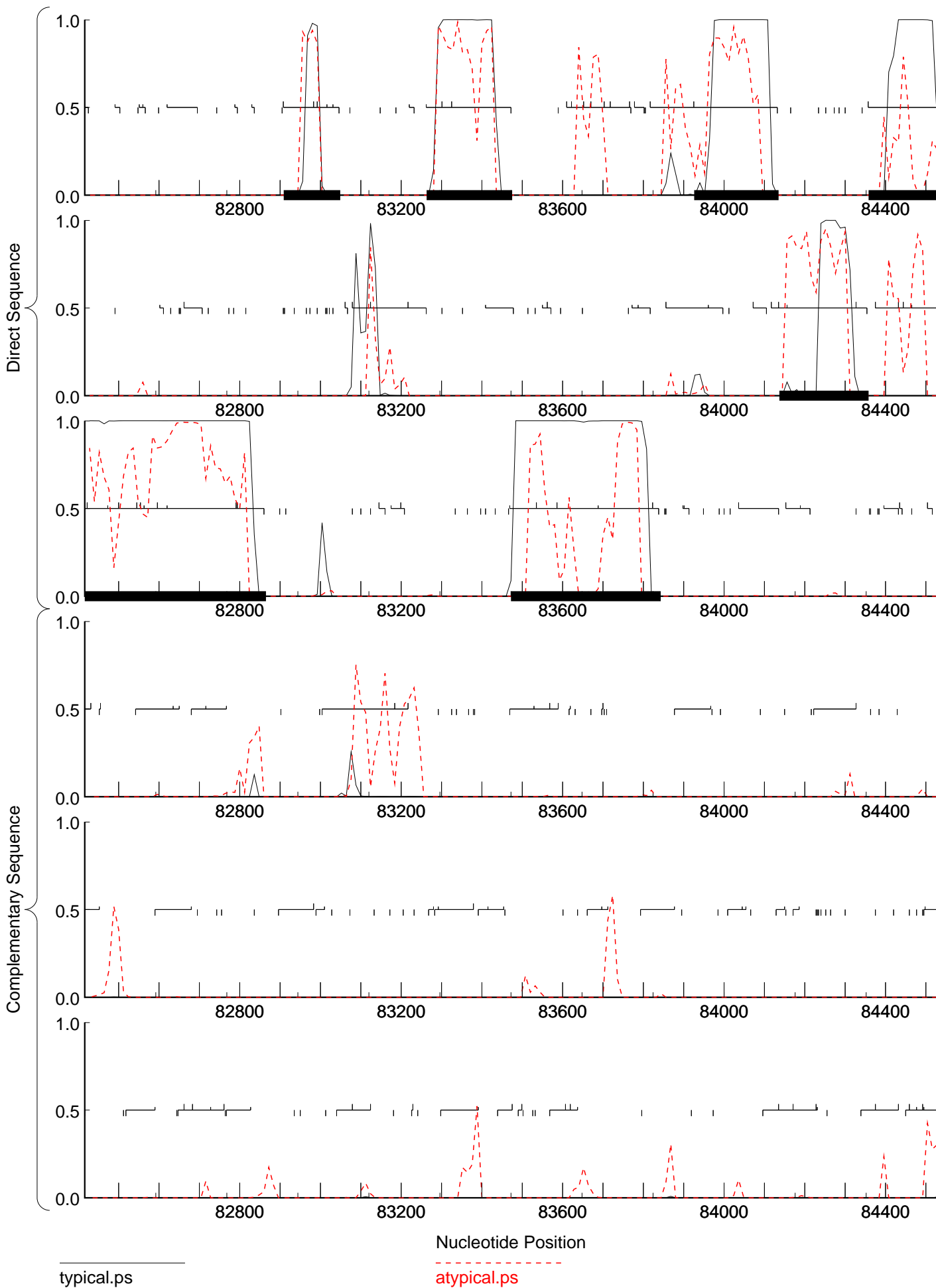




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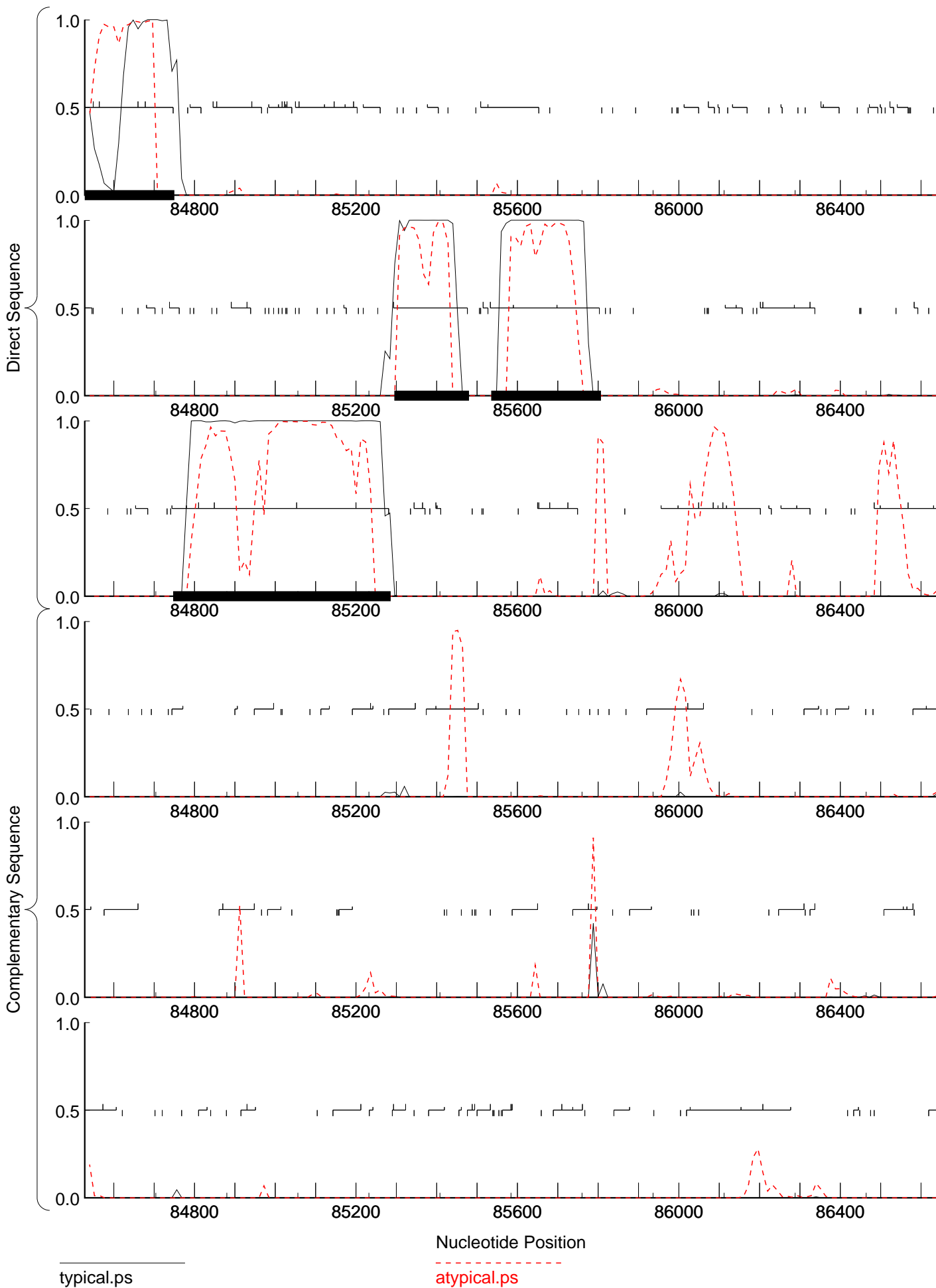
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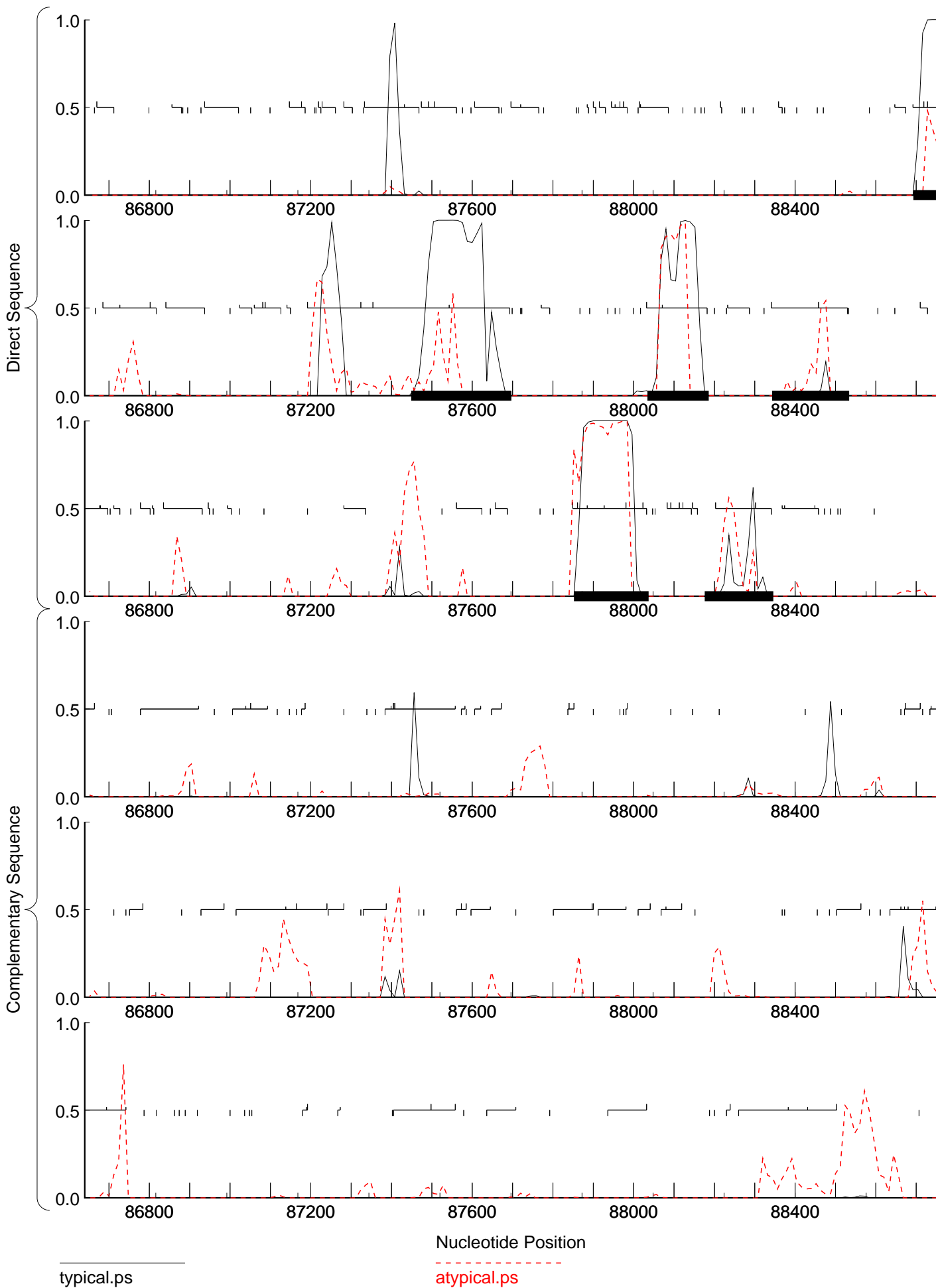


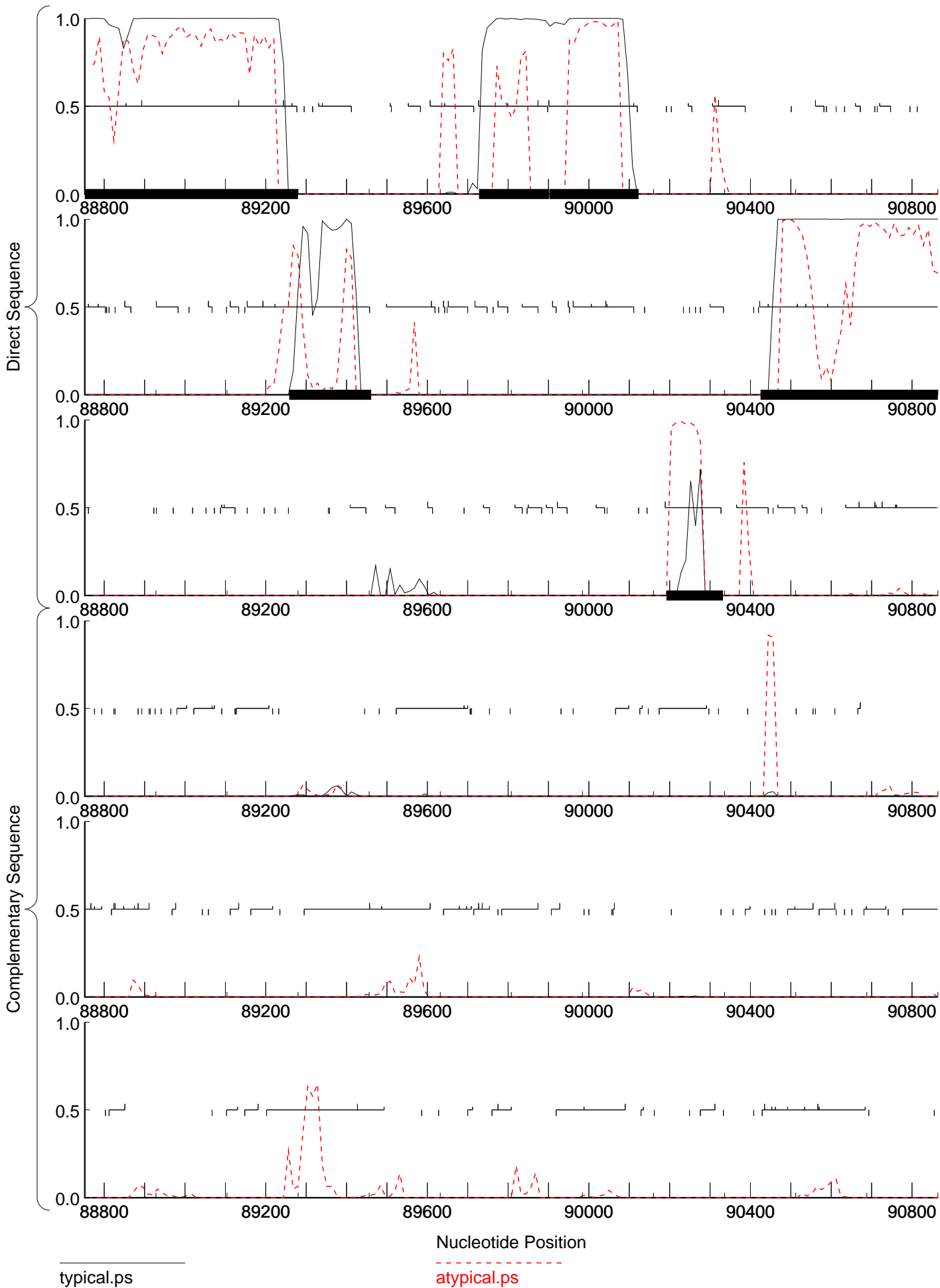


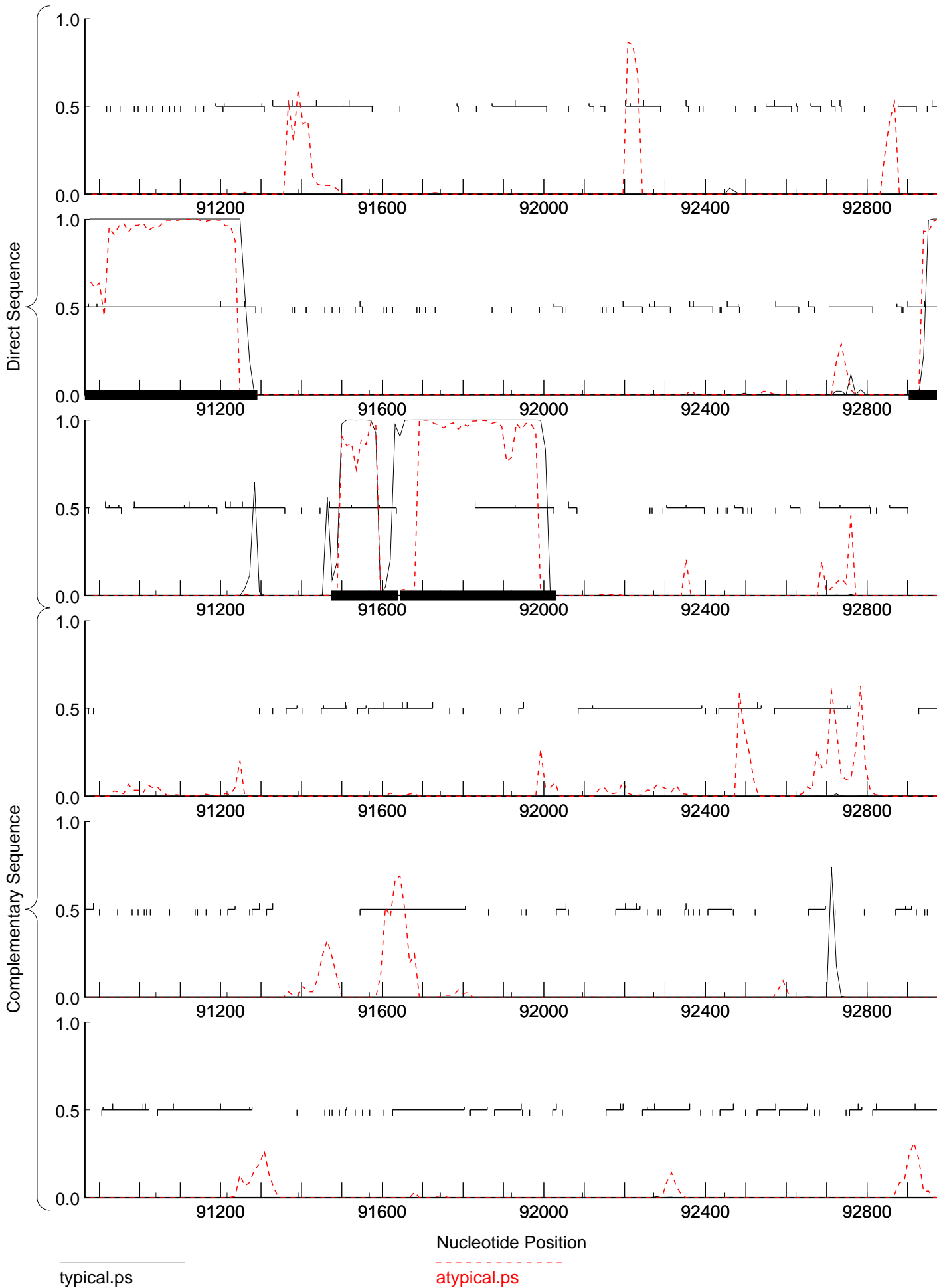
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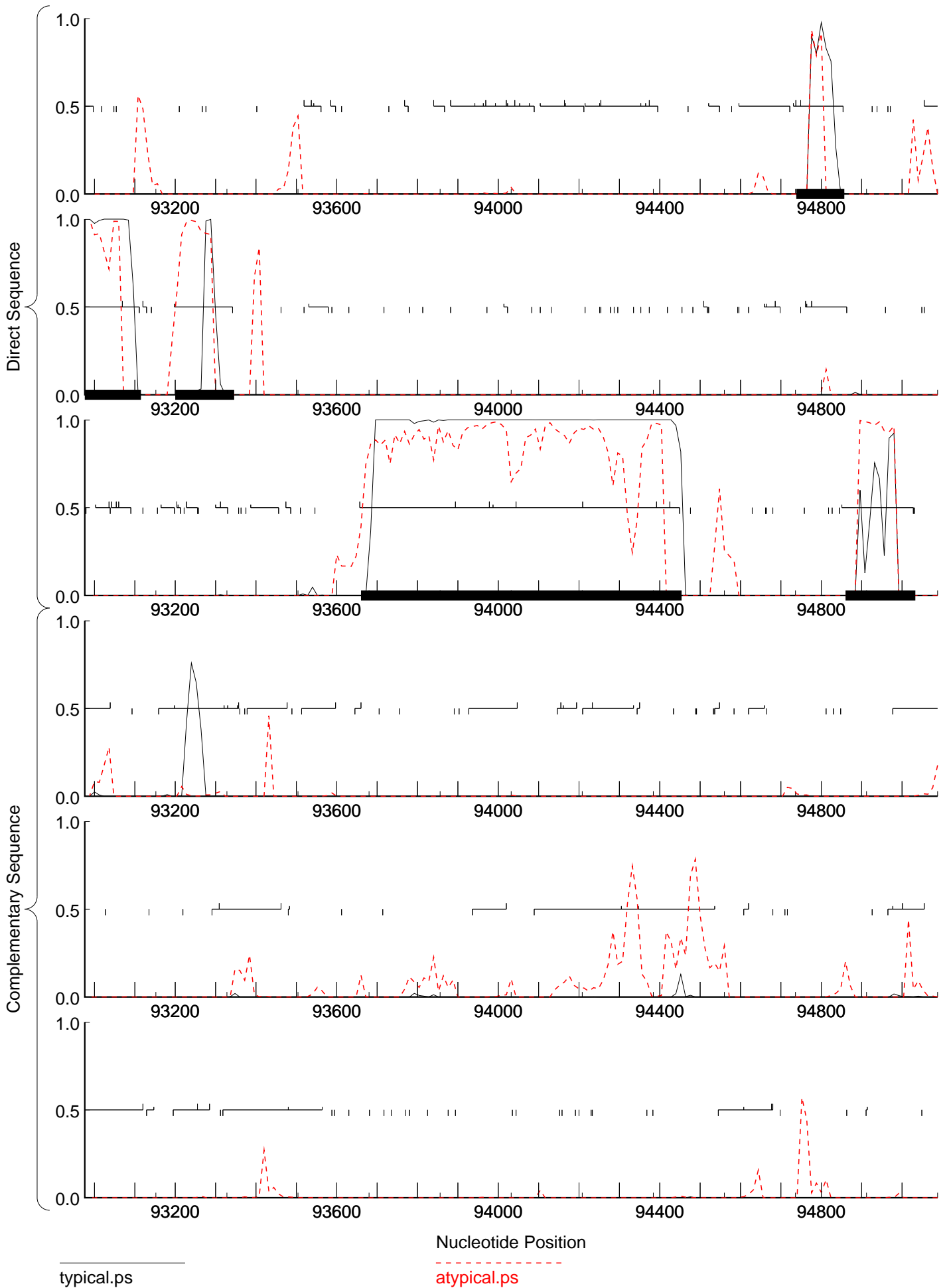
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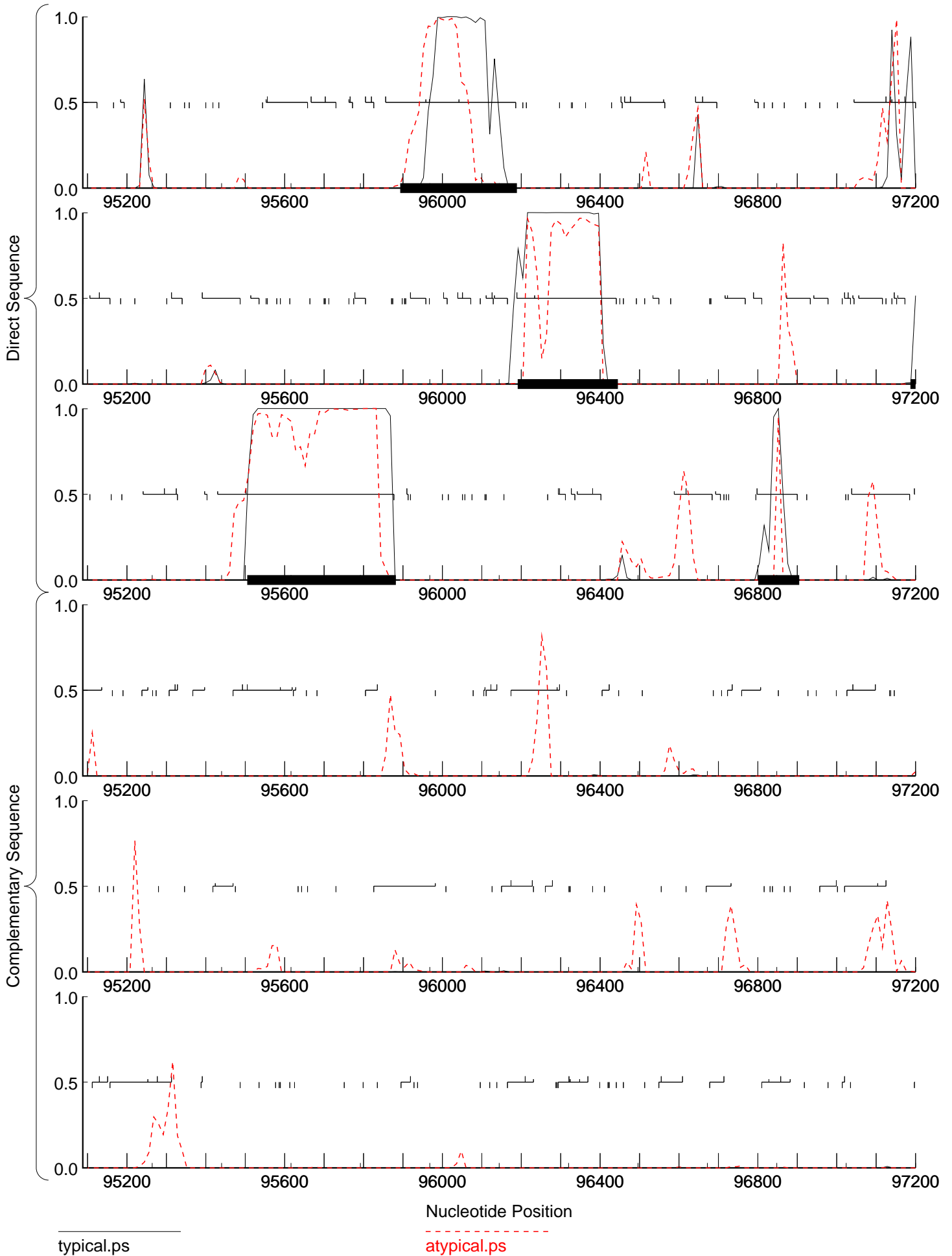


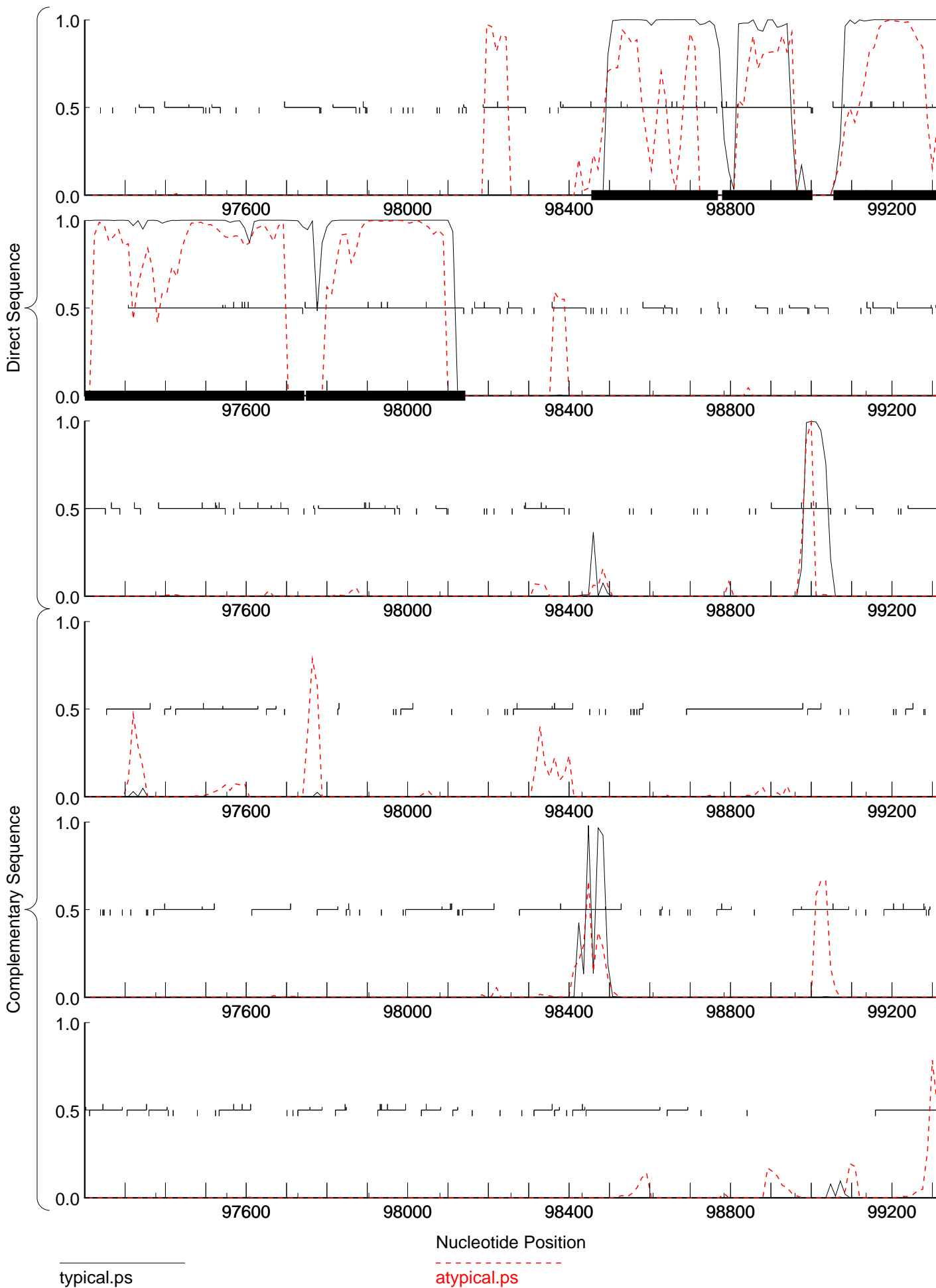


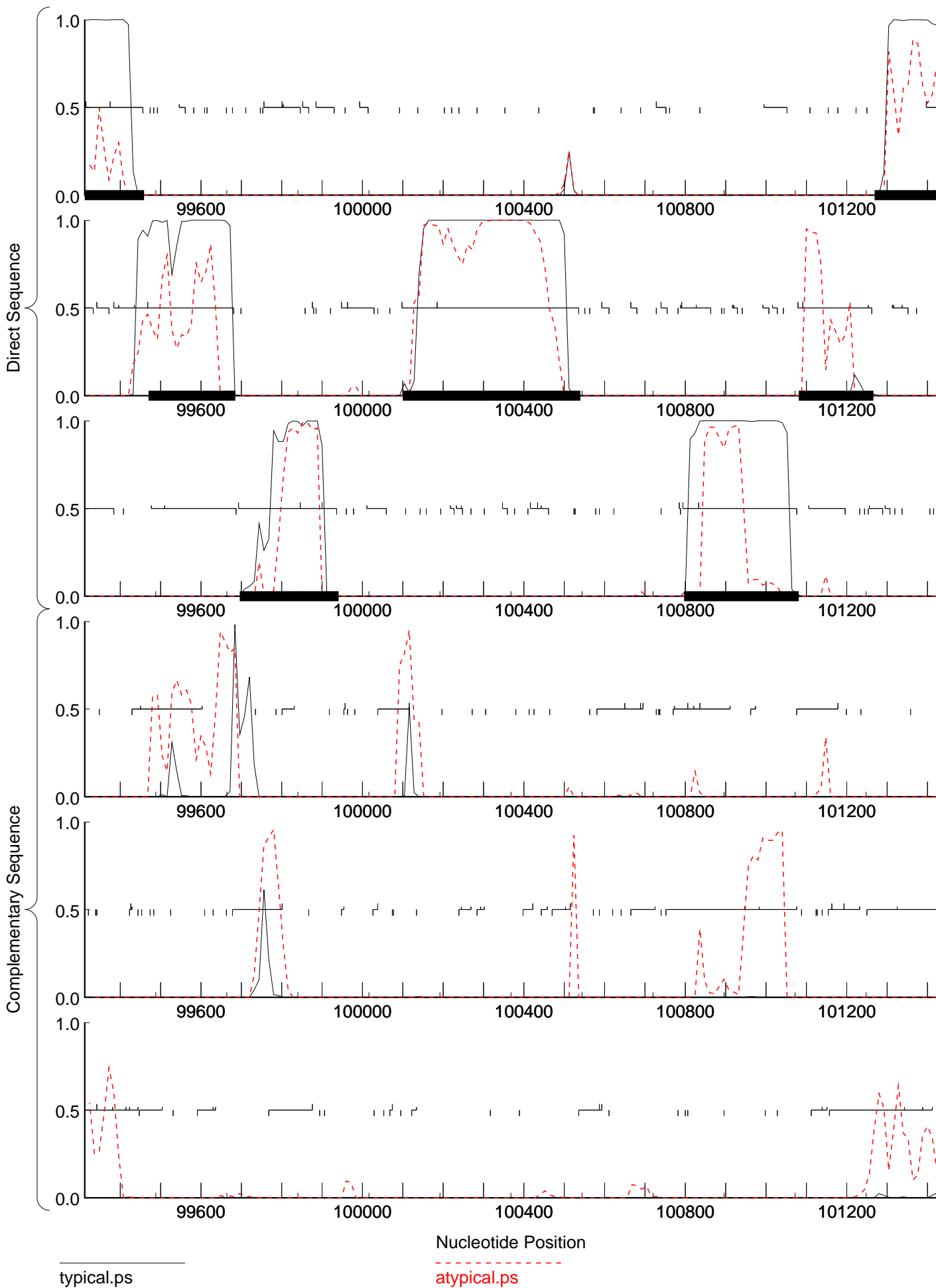


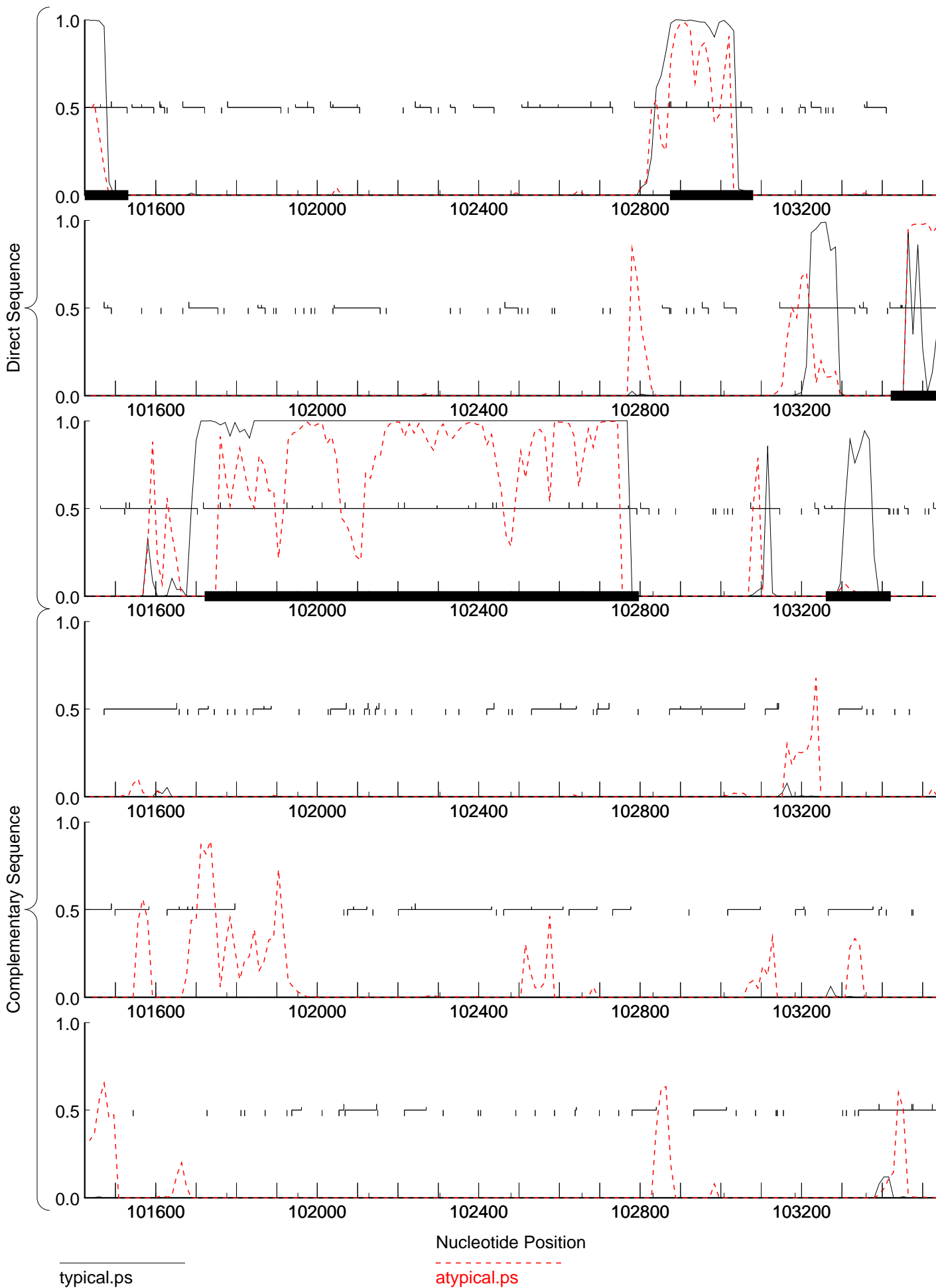






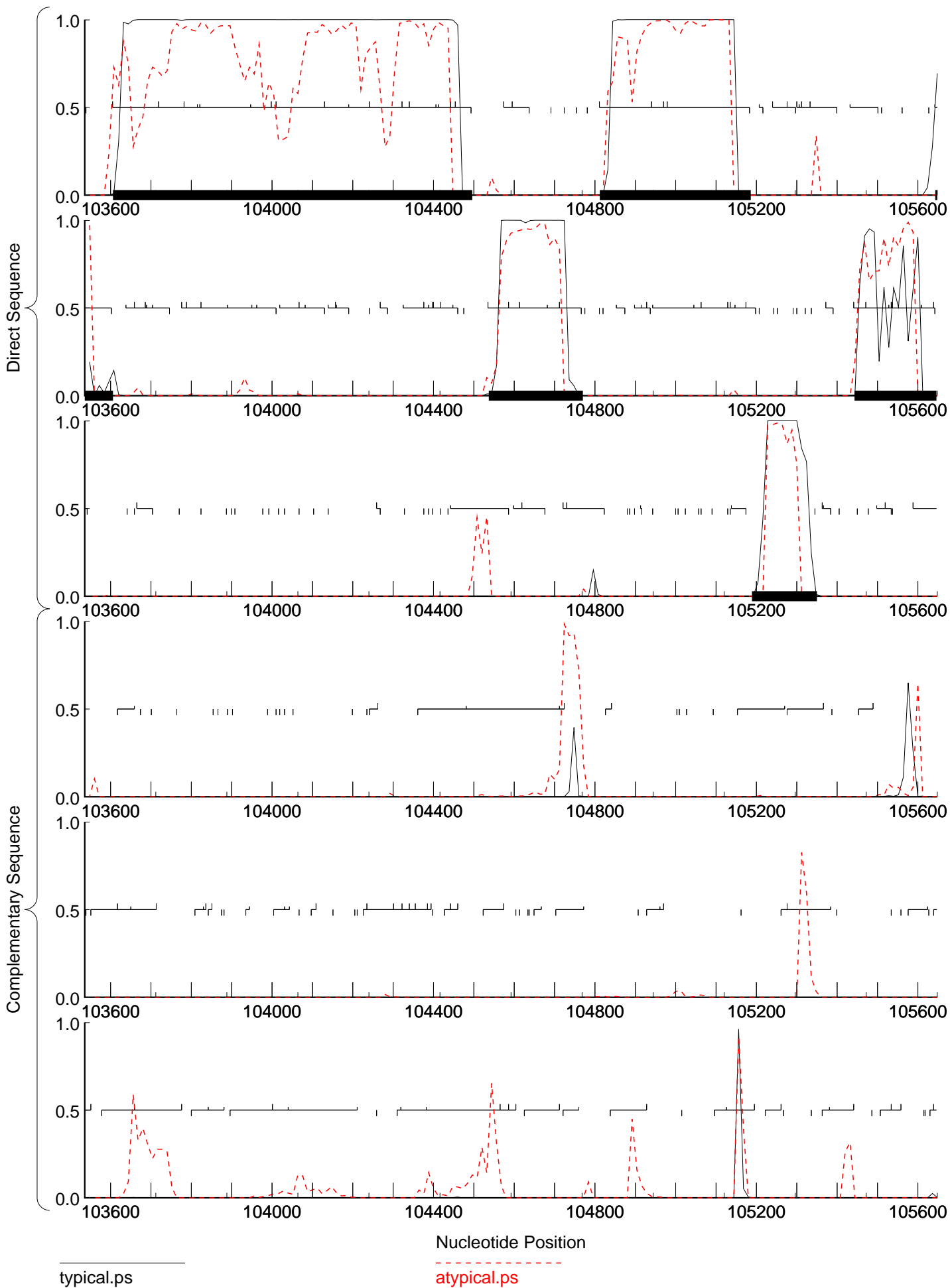


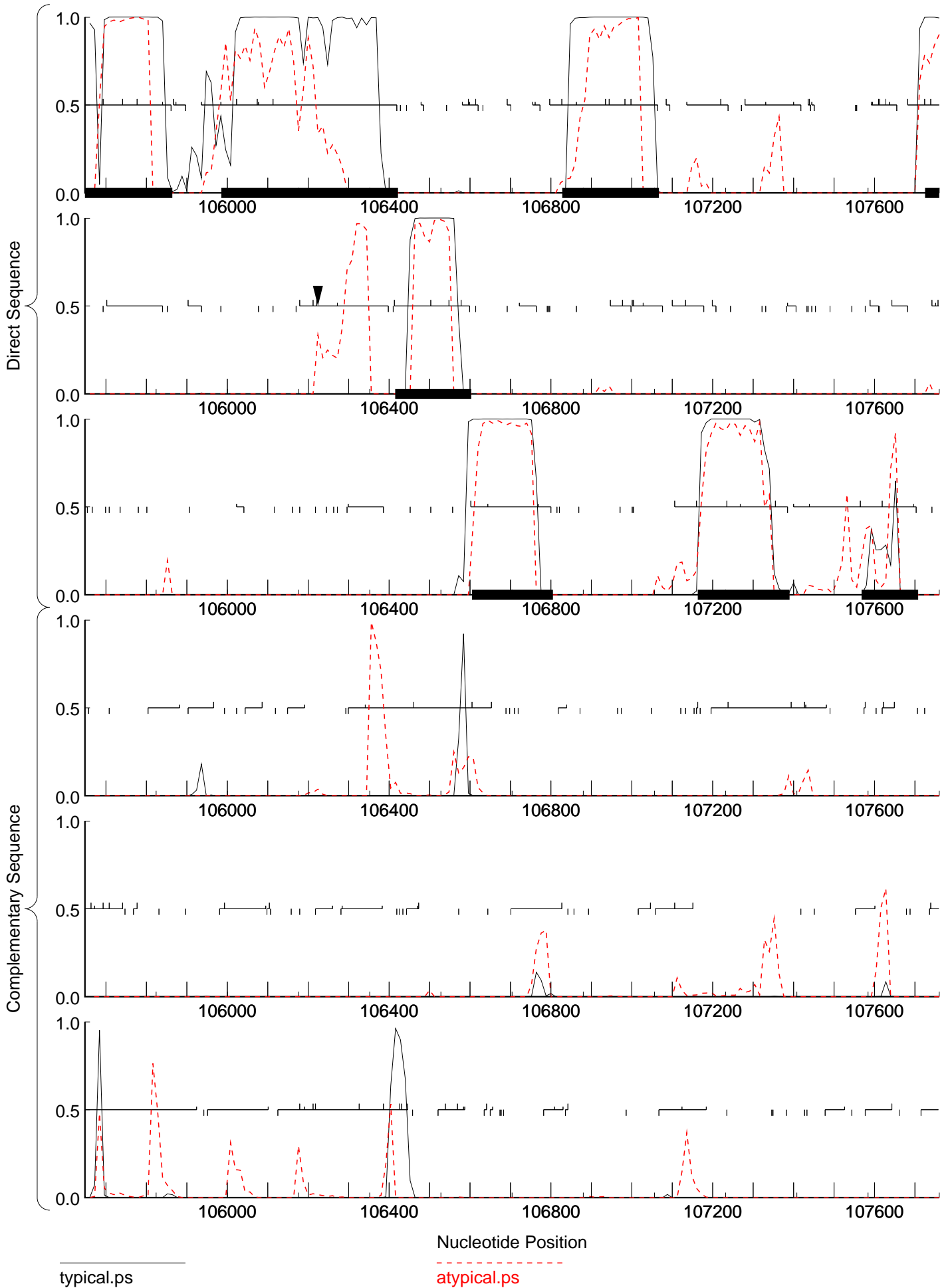




typical.ps

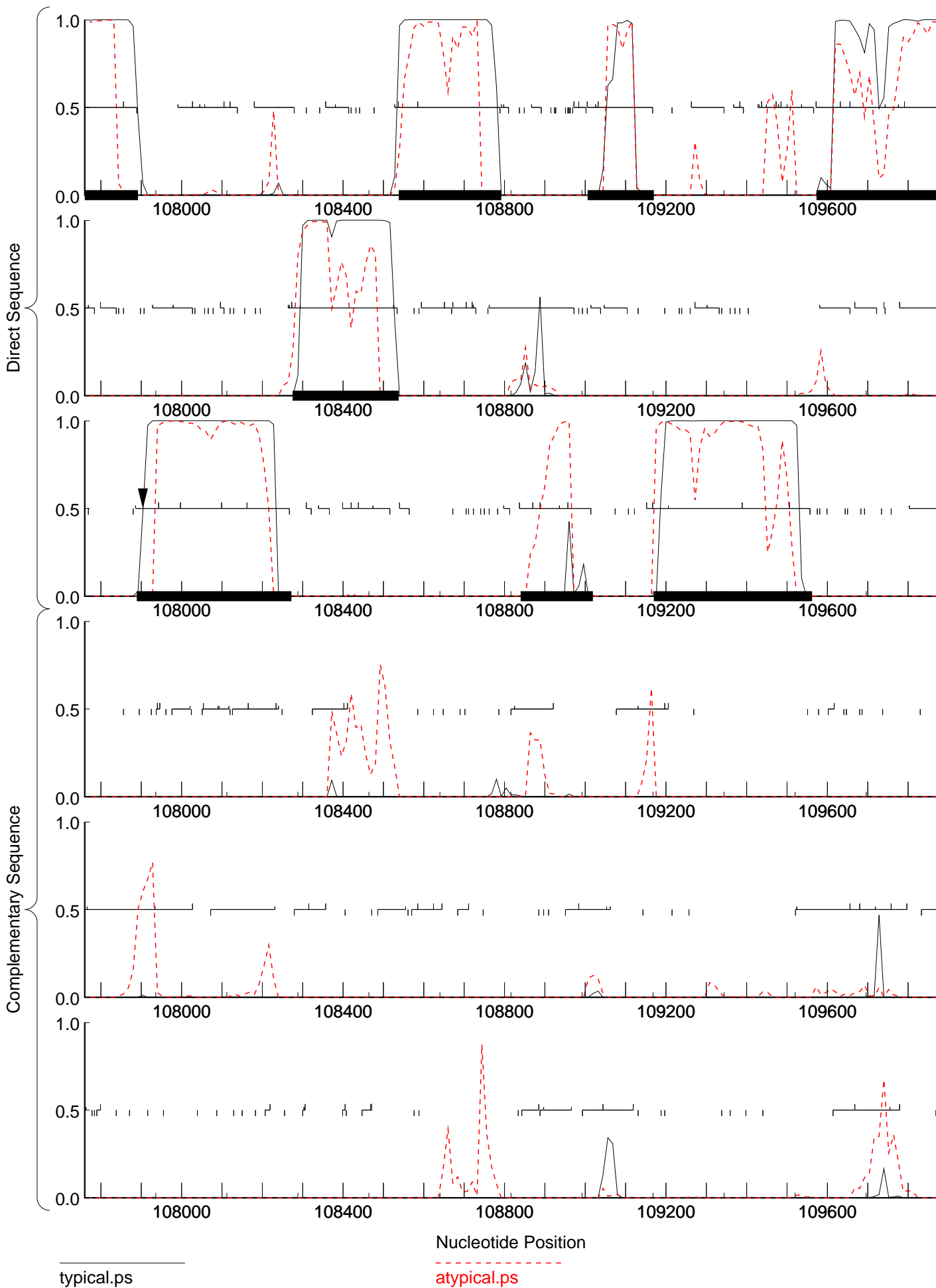
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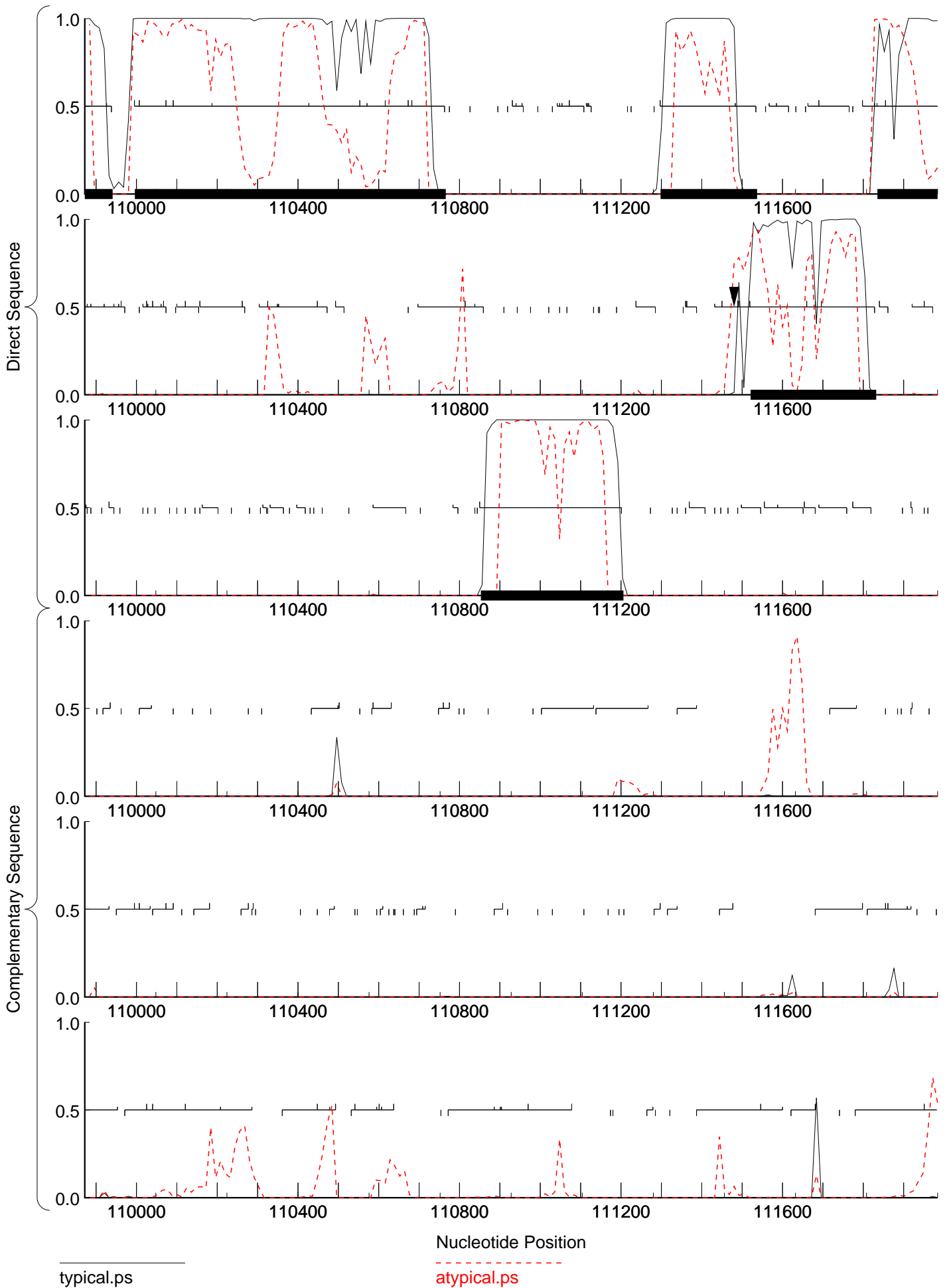


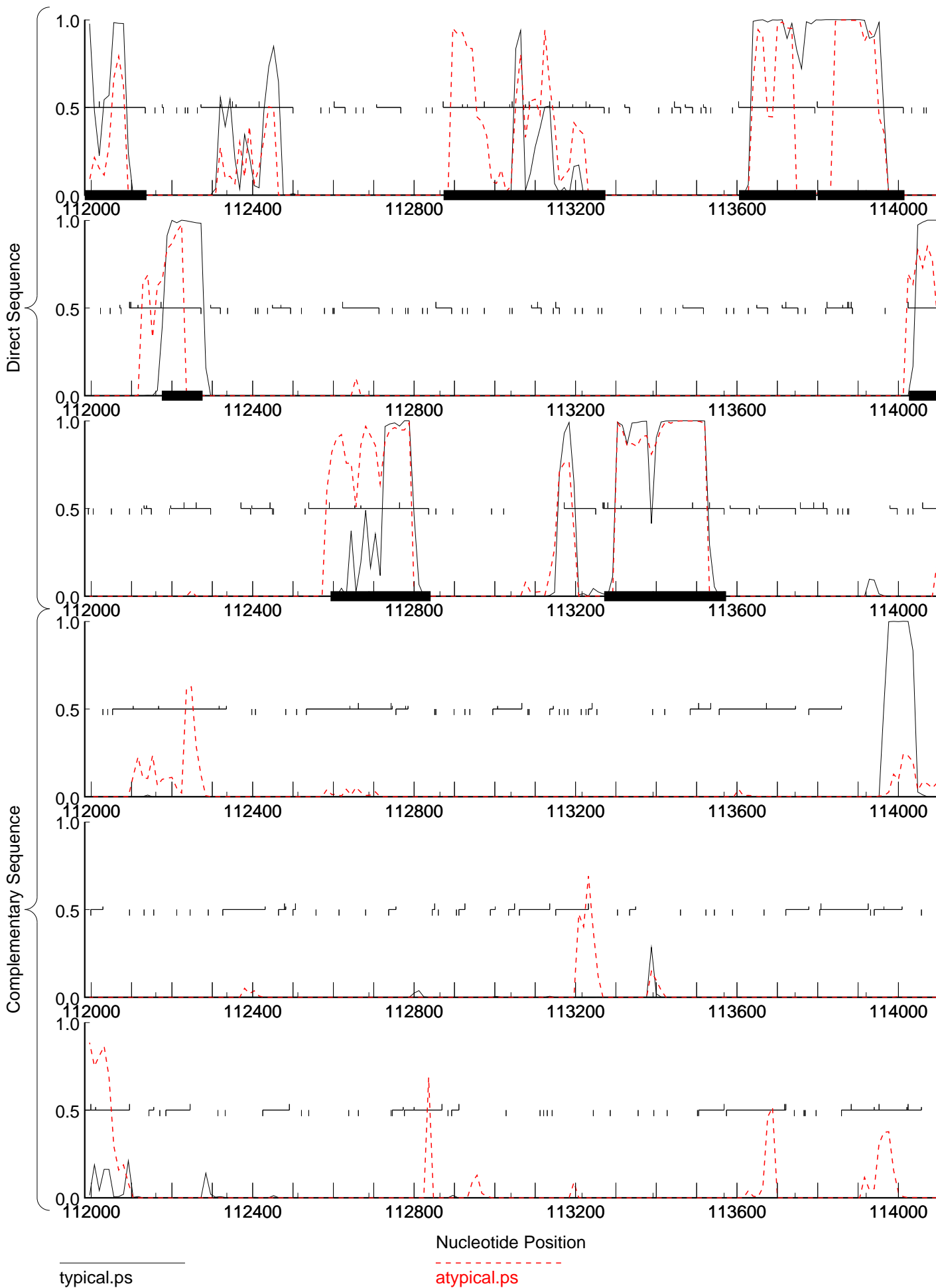


typical.ps

atypical.ps

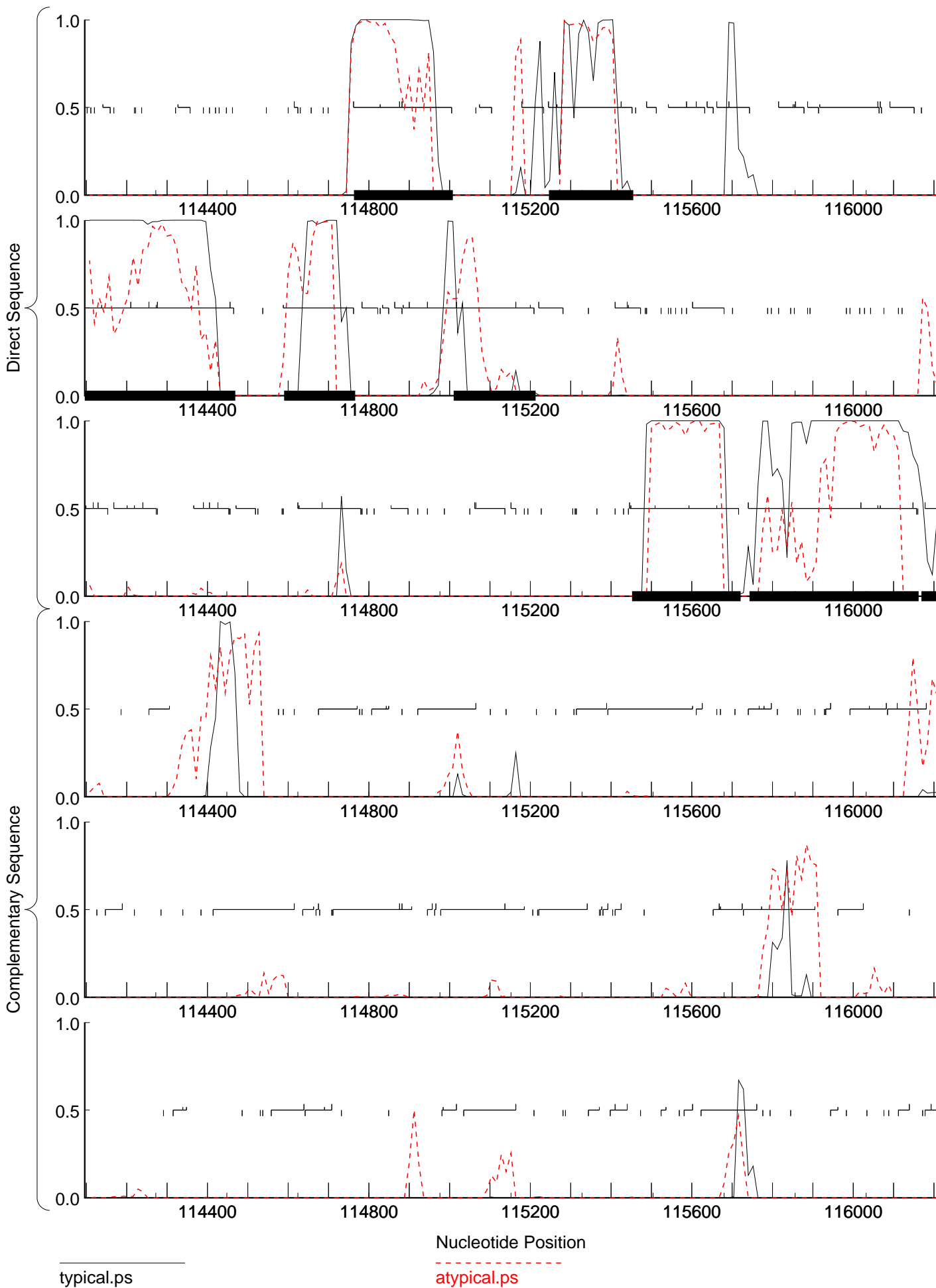






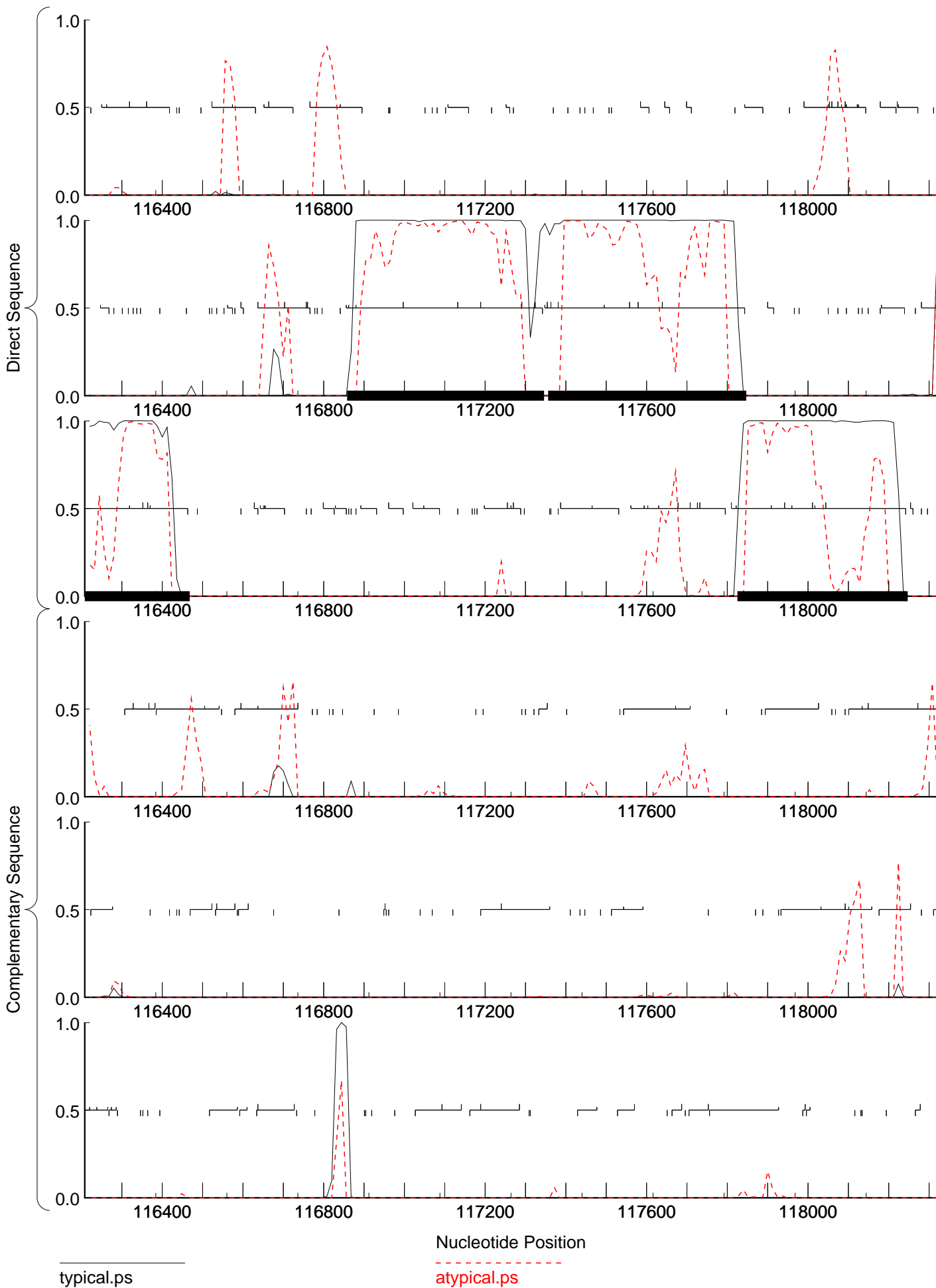
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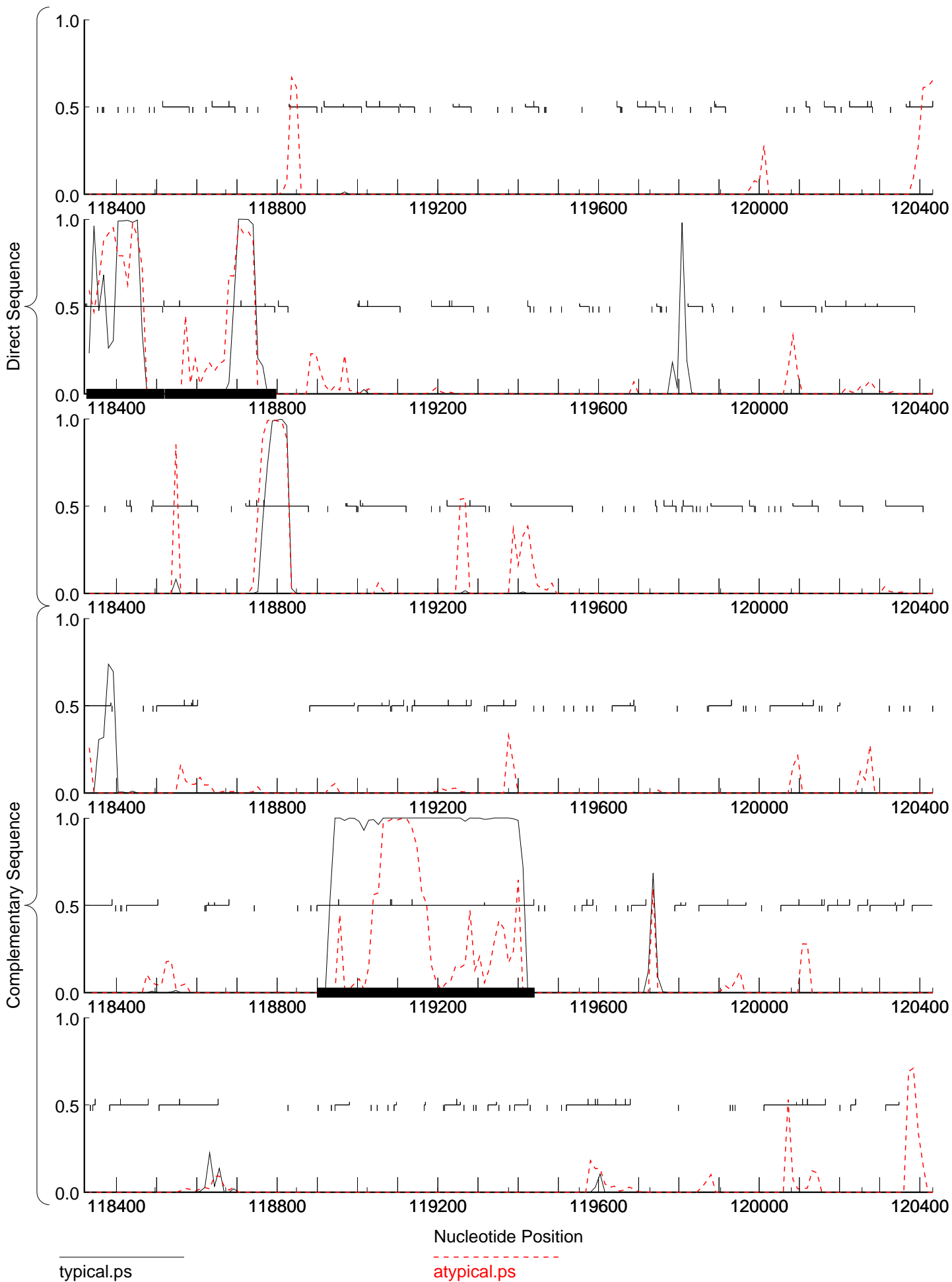
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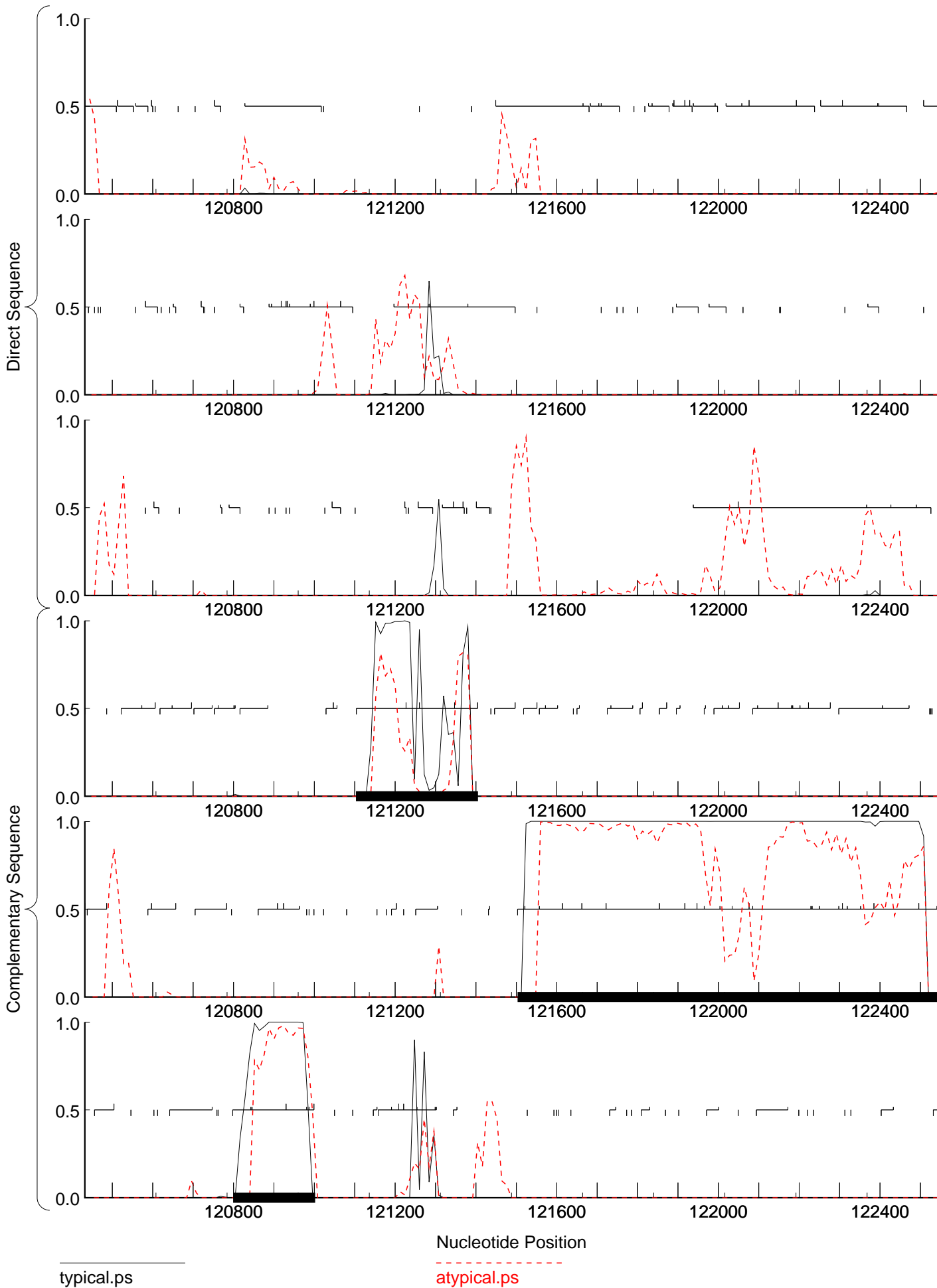
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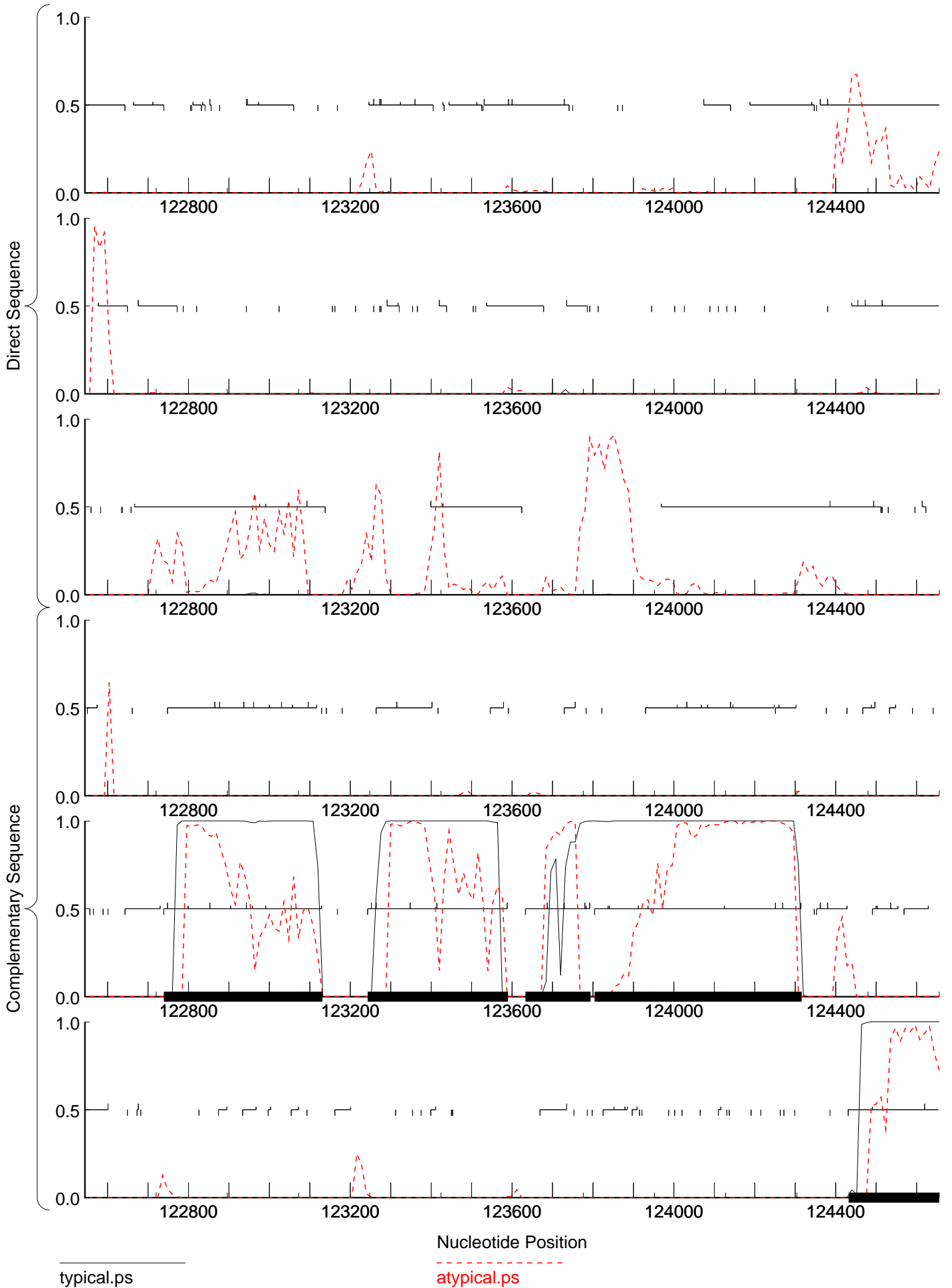


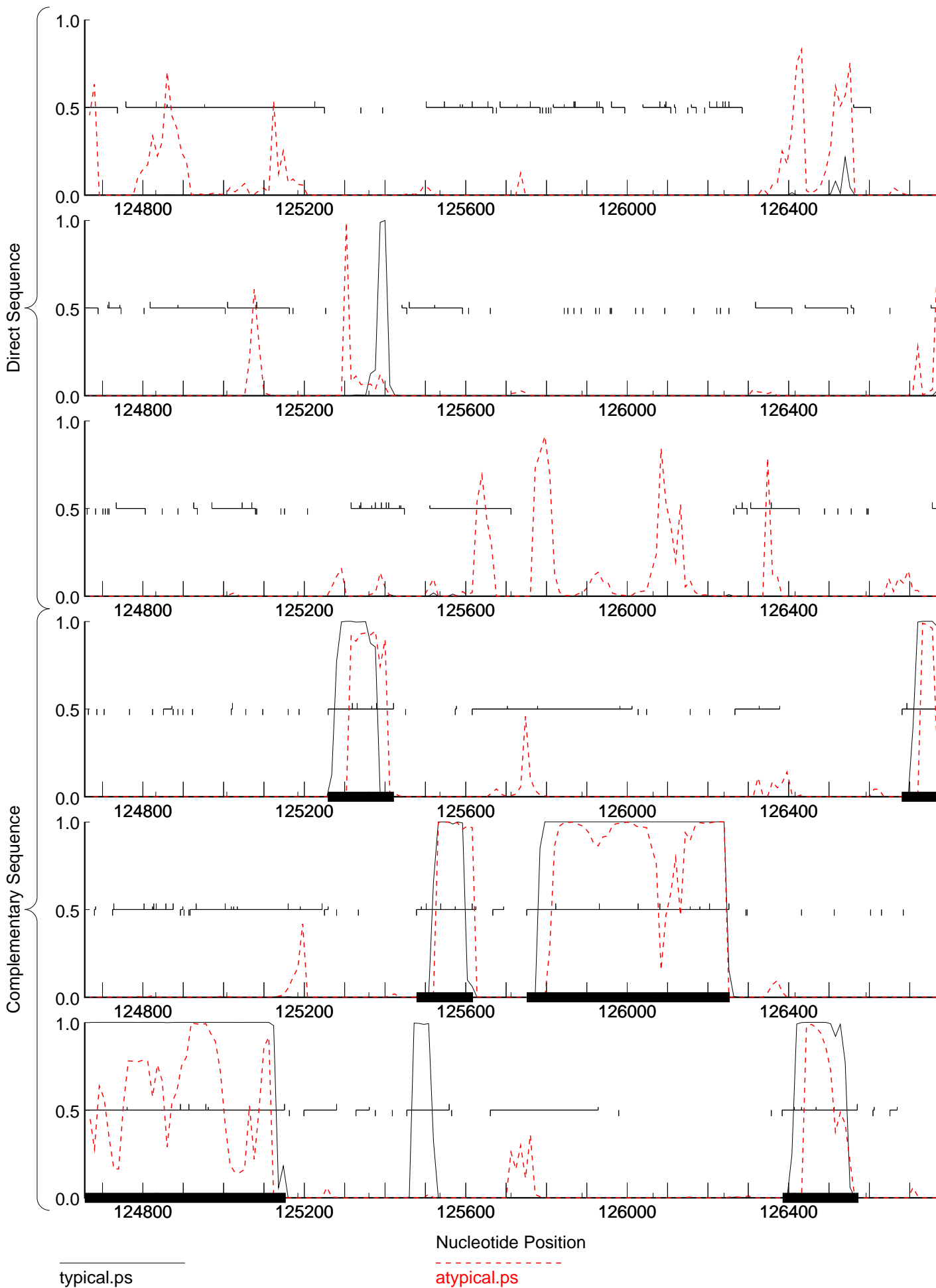


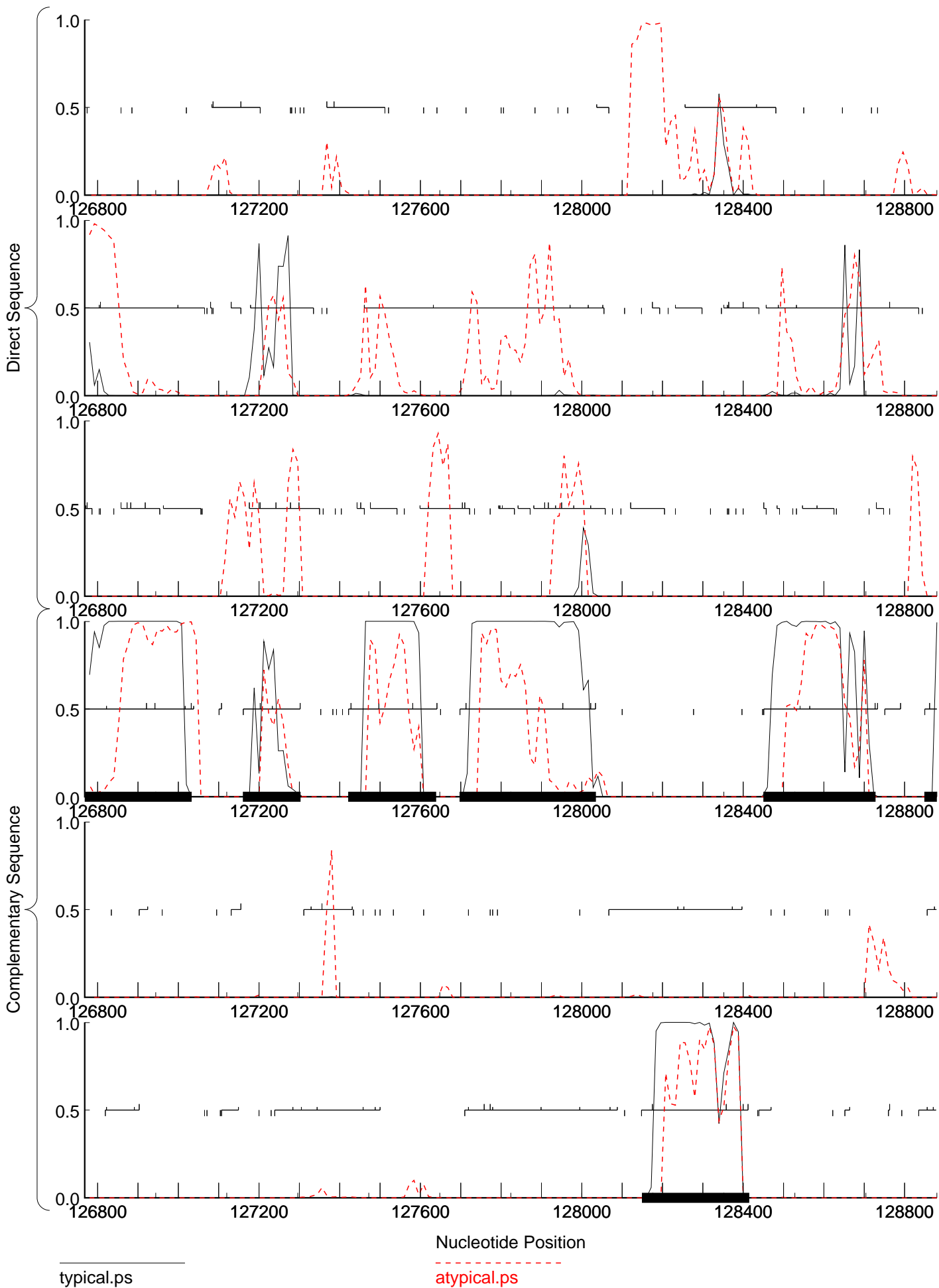
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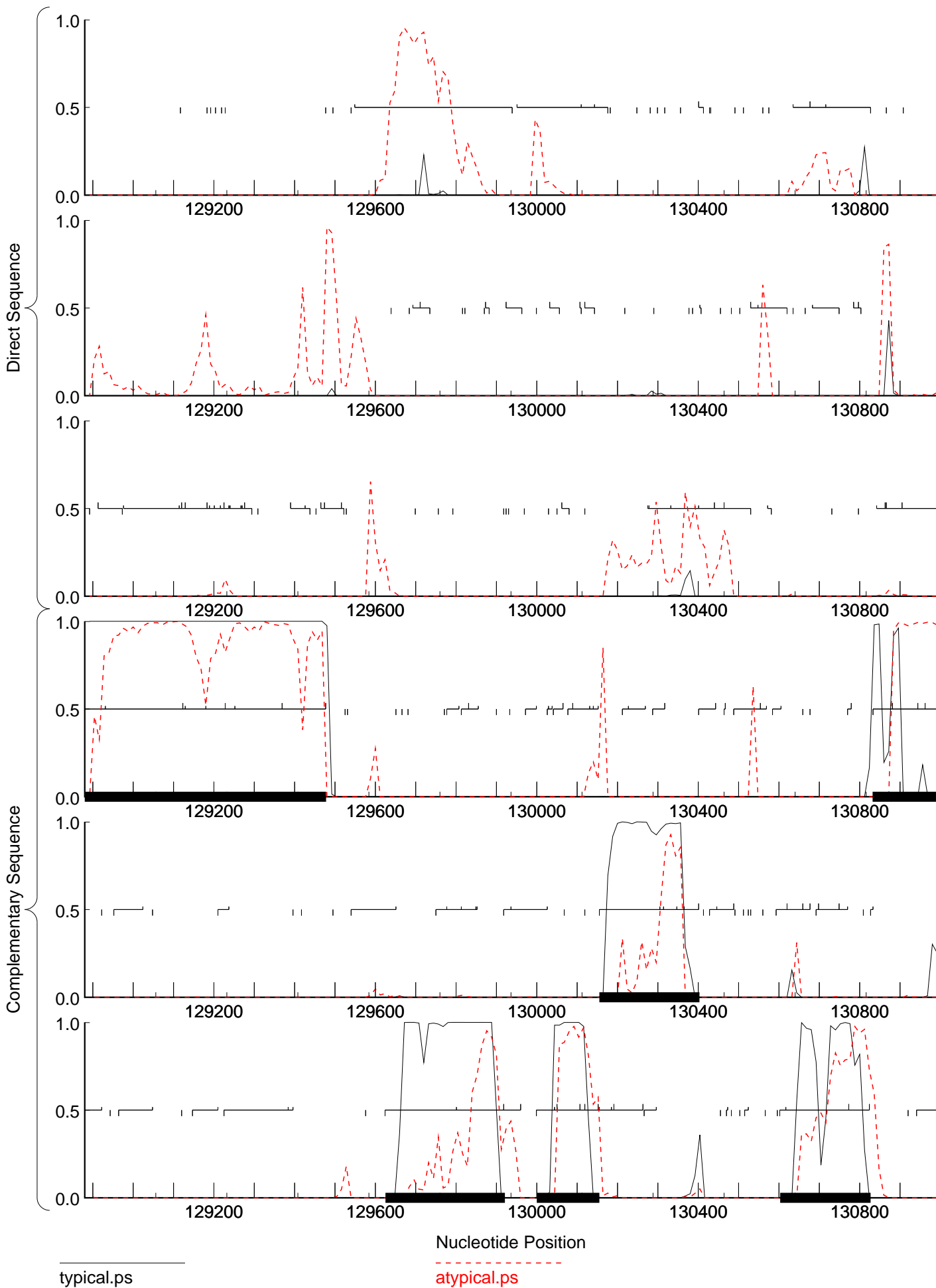






typical.ps

atypical.ps



typical.ps

atypical.ps

