

GeneMark

Version 2.5p (09.08.06)

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PROGRAM INFORMATION

Sequence : Mycobacterium phage PhatBacter complete sequence, 76217 bp including 9-base 3' overhang (CG
Analysis Date : 6/22/18 at 13:19:30
Pages : 38
Sequence Length : 76217 bp
GC Content : 62.98%

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,

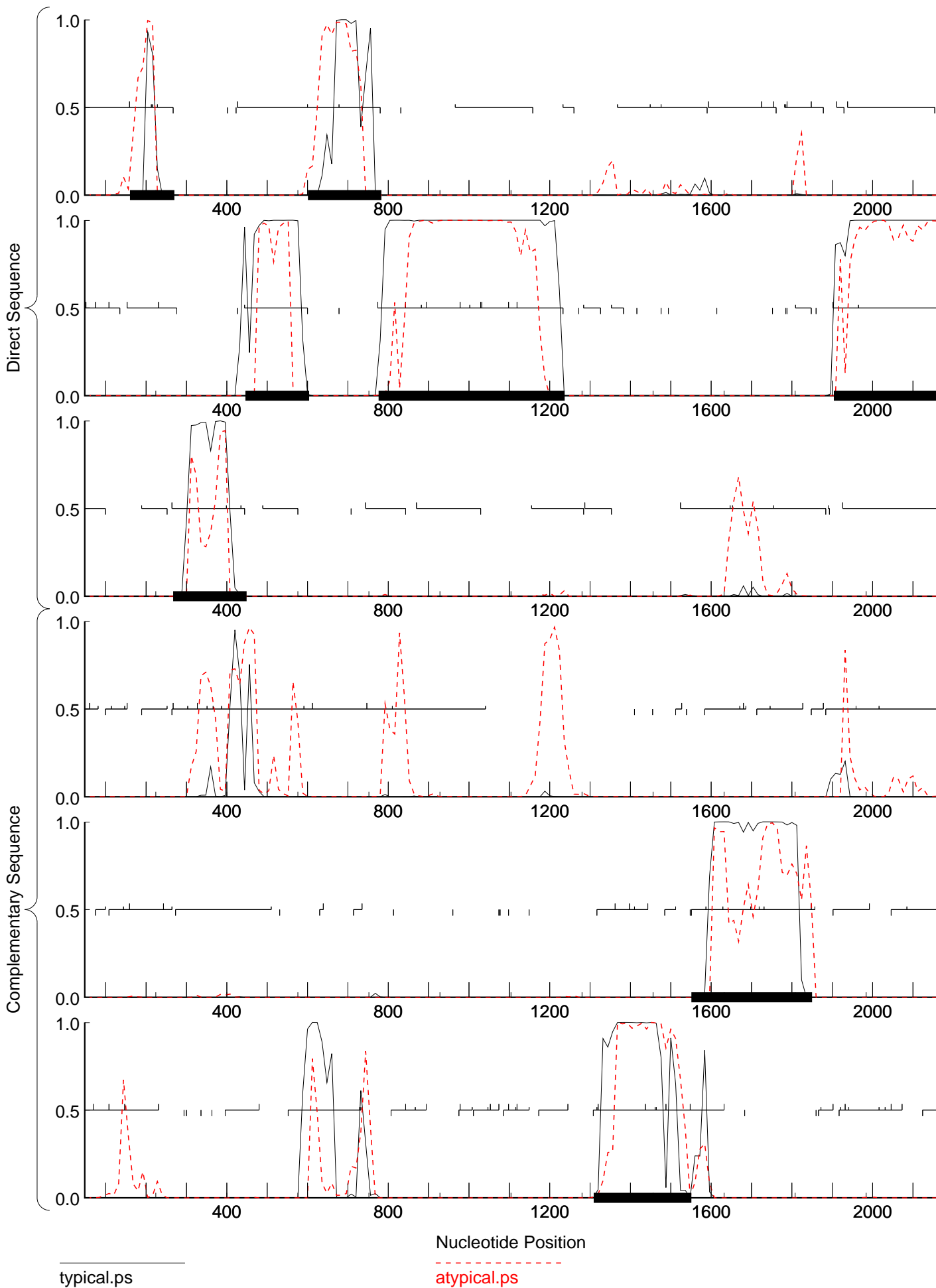
Matrix notes & comments

Training set derived by GeneMarkS, 4.30 October 2014
Fri Jun 22 13:19:30 2018

MATRIX INFORMATION

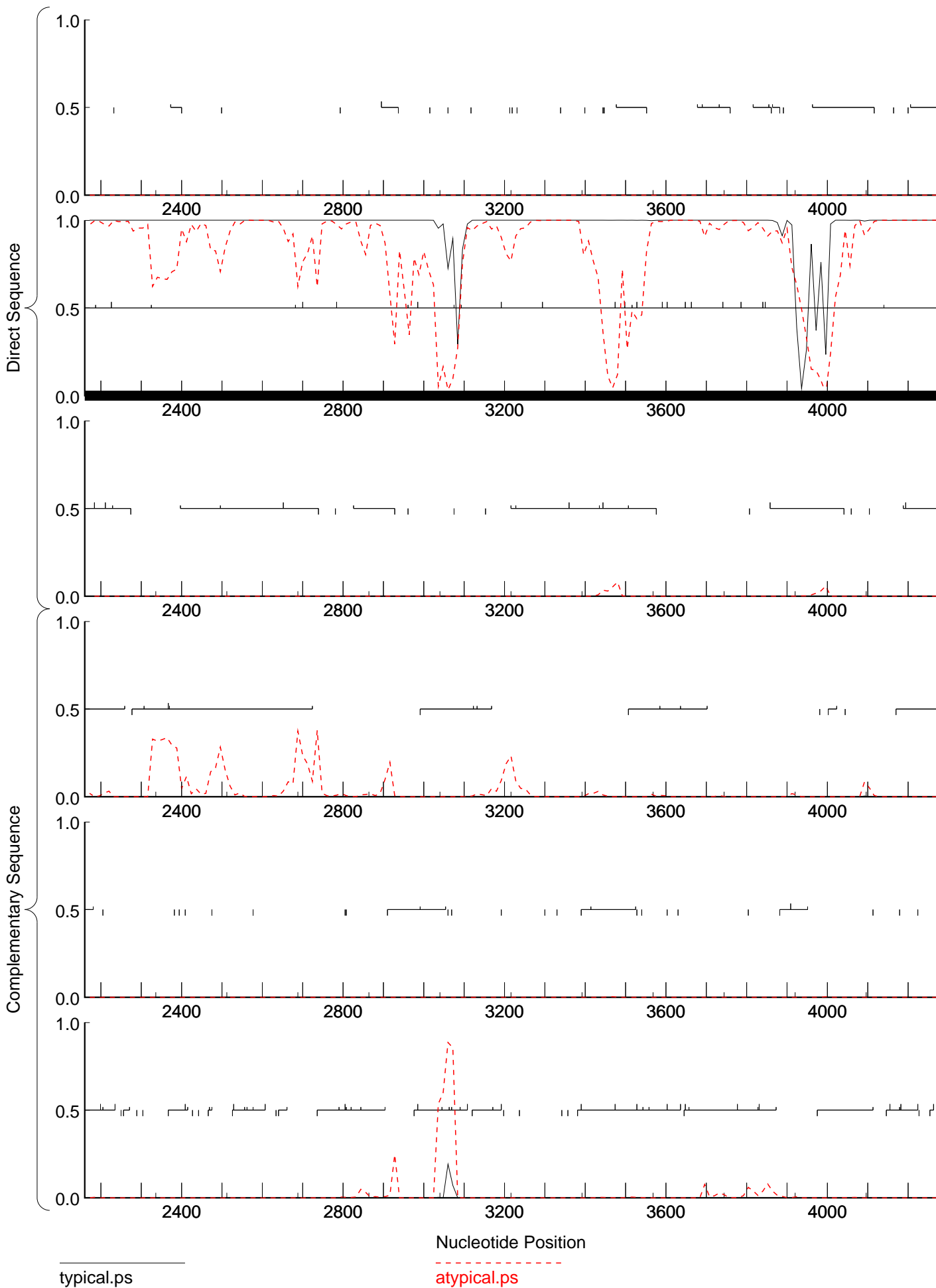
Matrix : -
Author : -
Order : 4

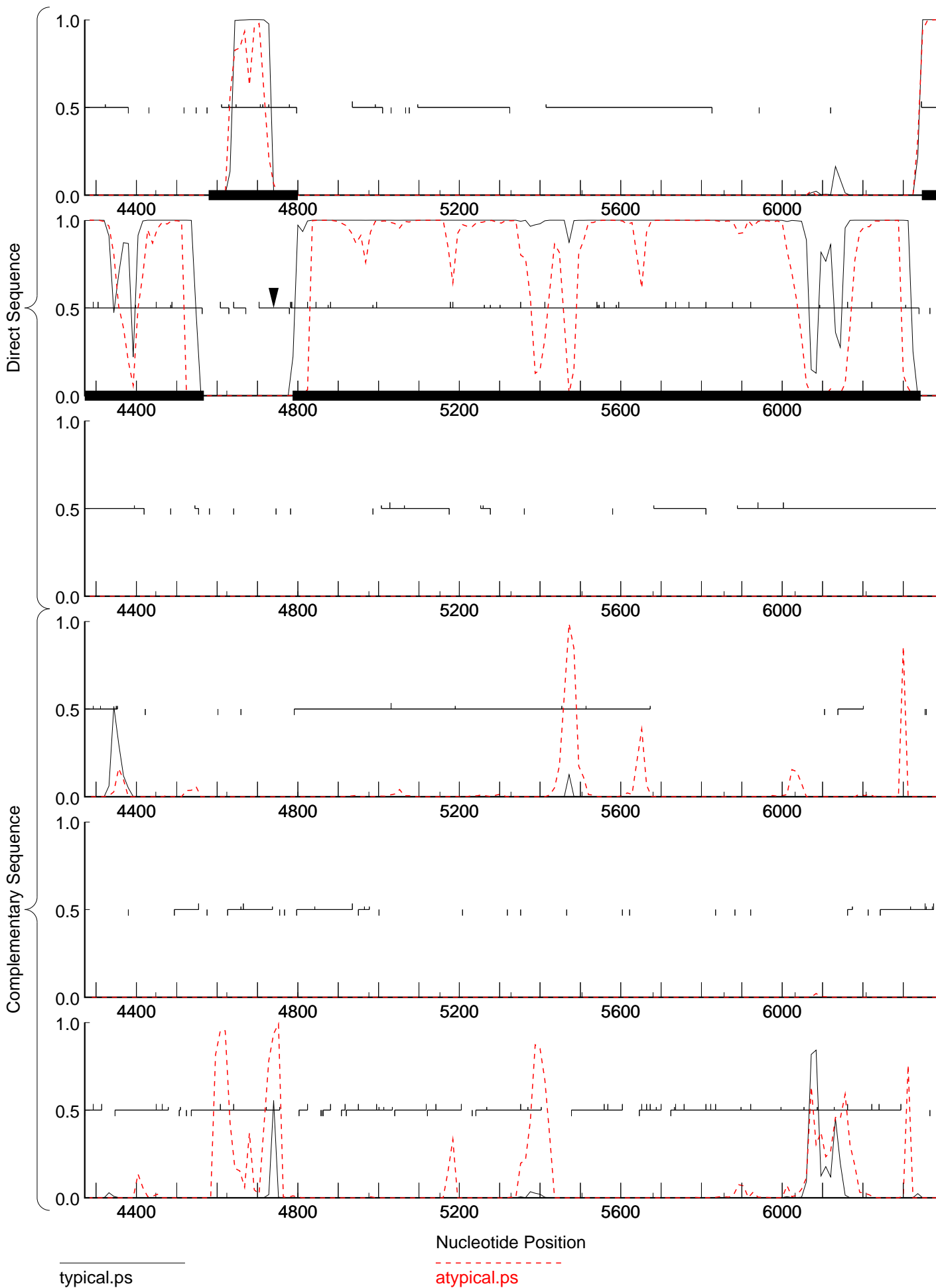
Send questions / comments to:
Dr. M. Borodovsky
Georgia Institute of Technology
School of Biology
Atlanta, GA 30332-0230

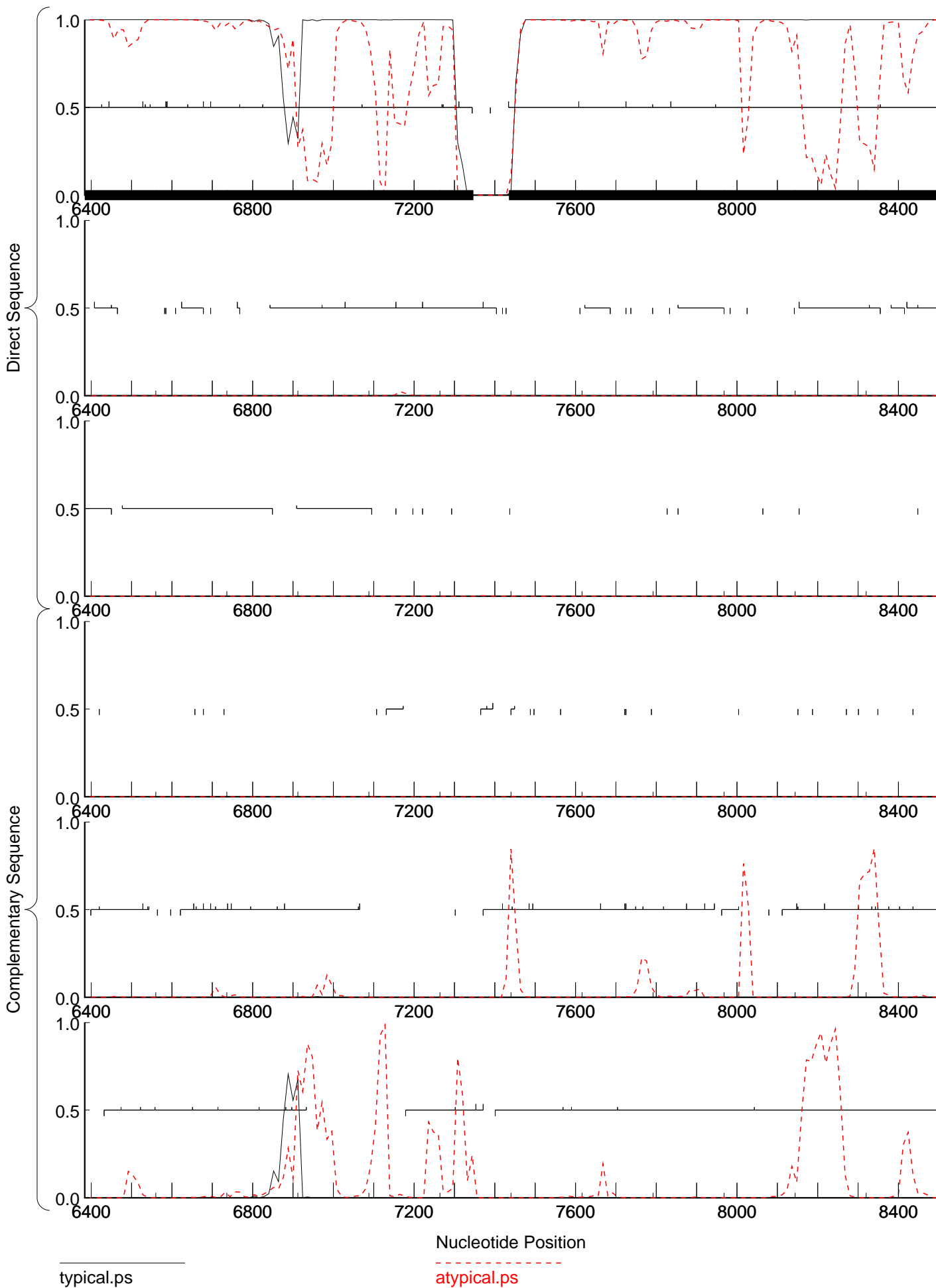


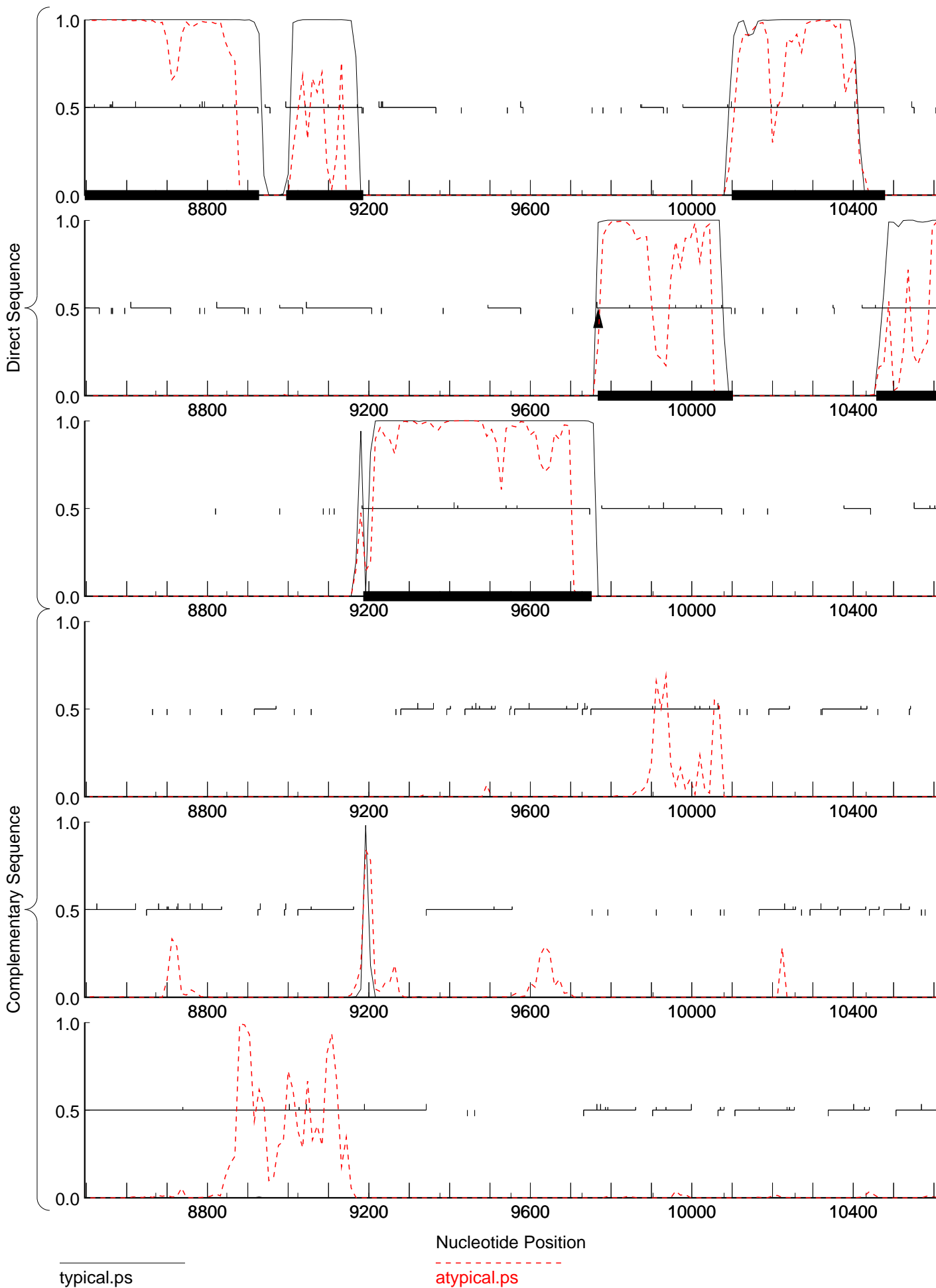
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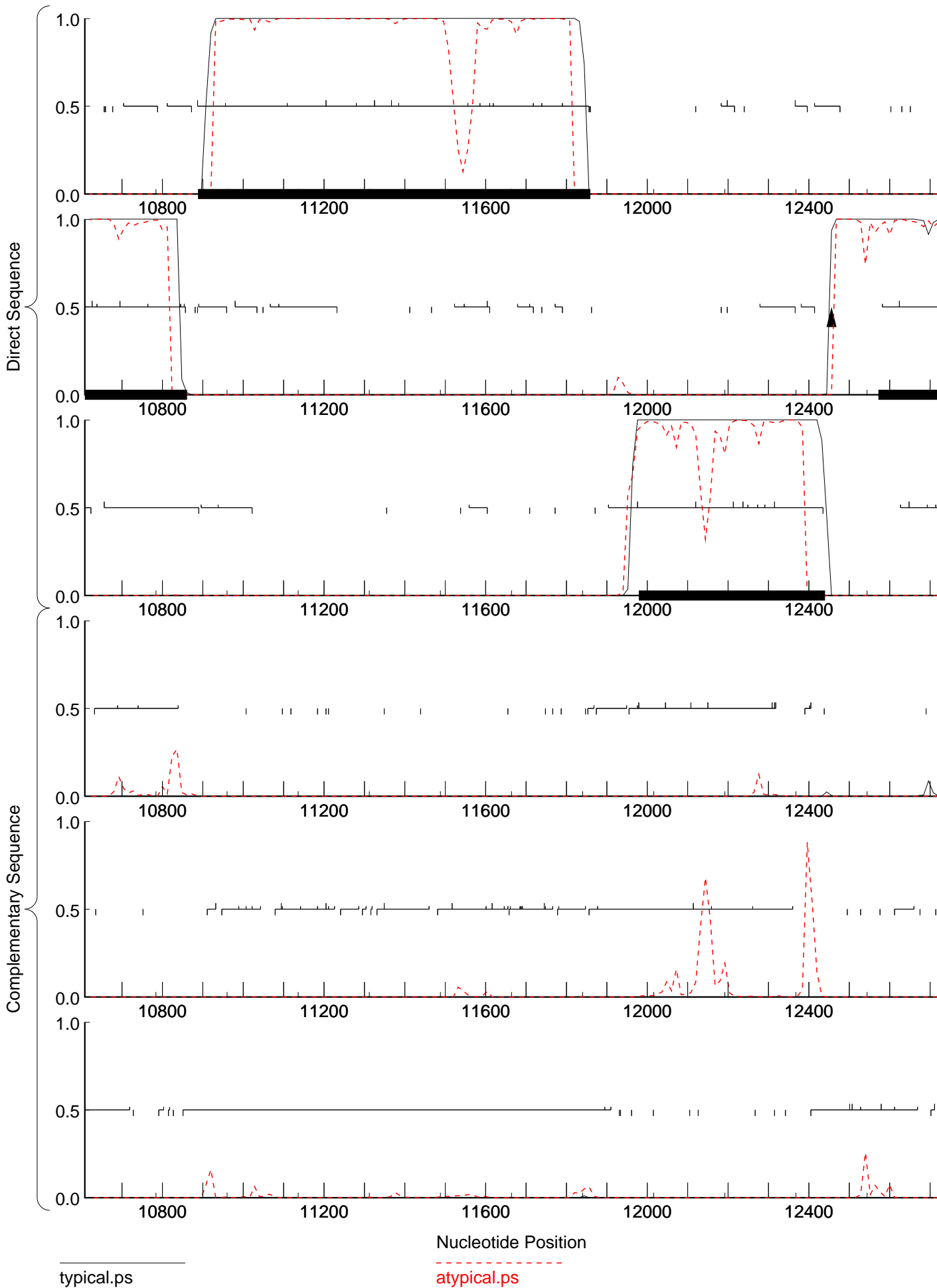
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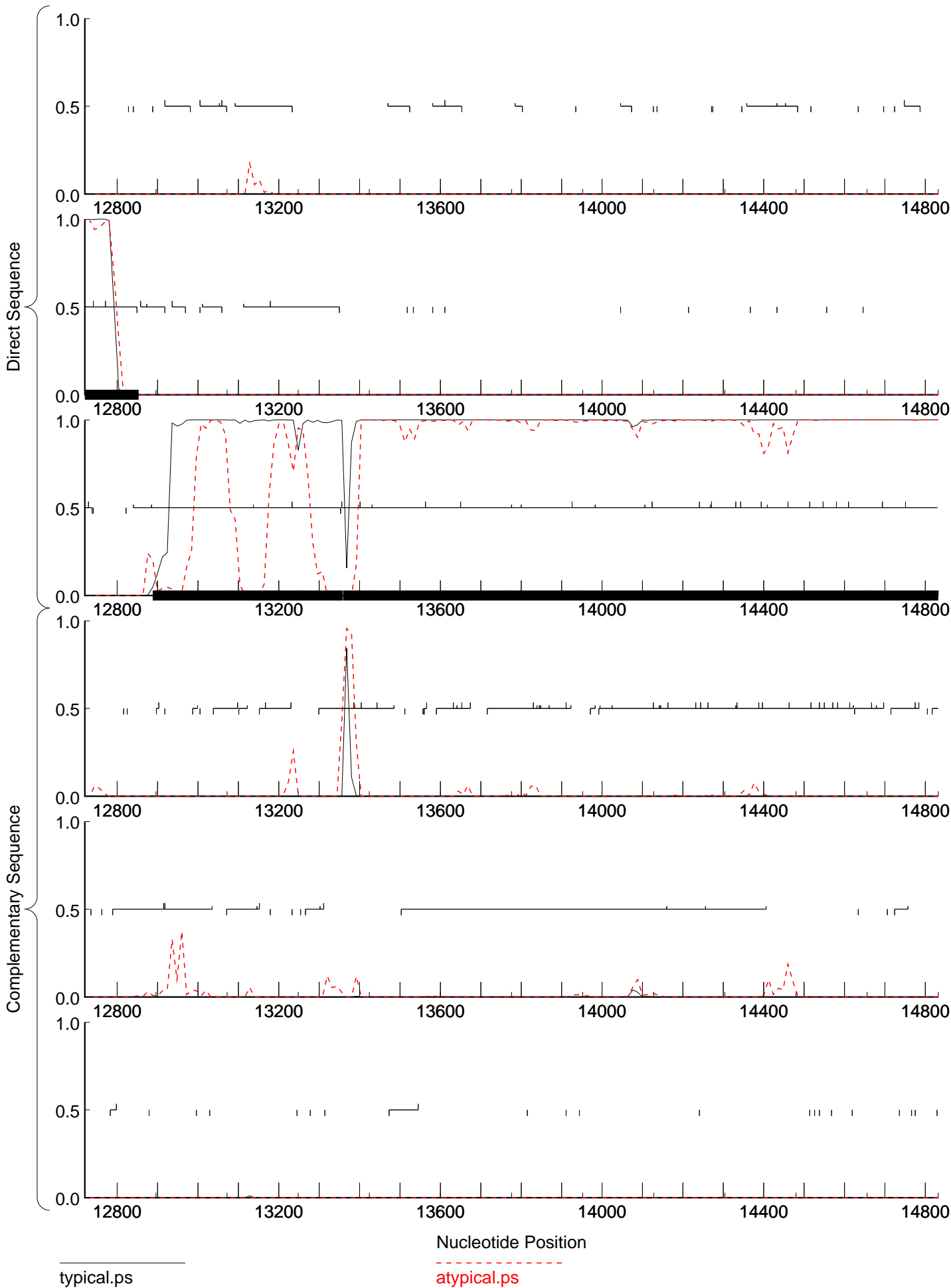


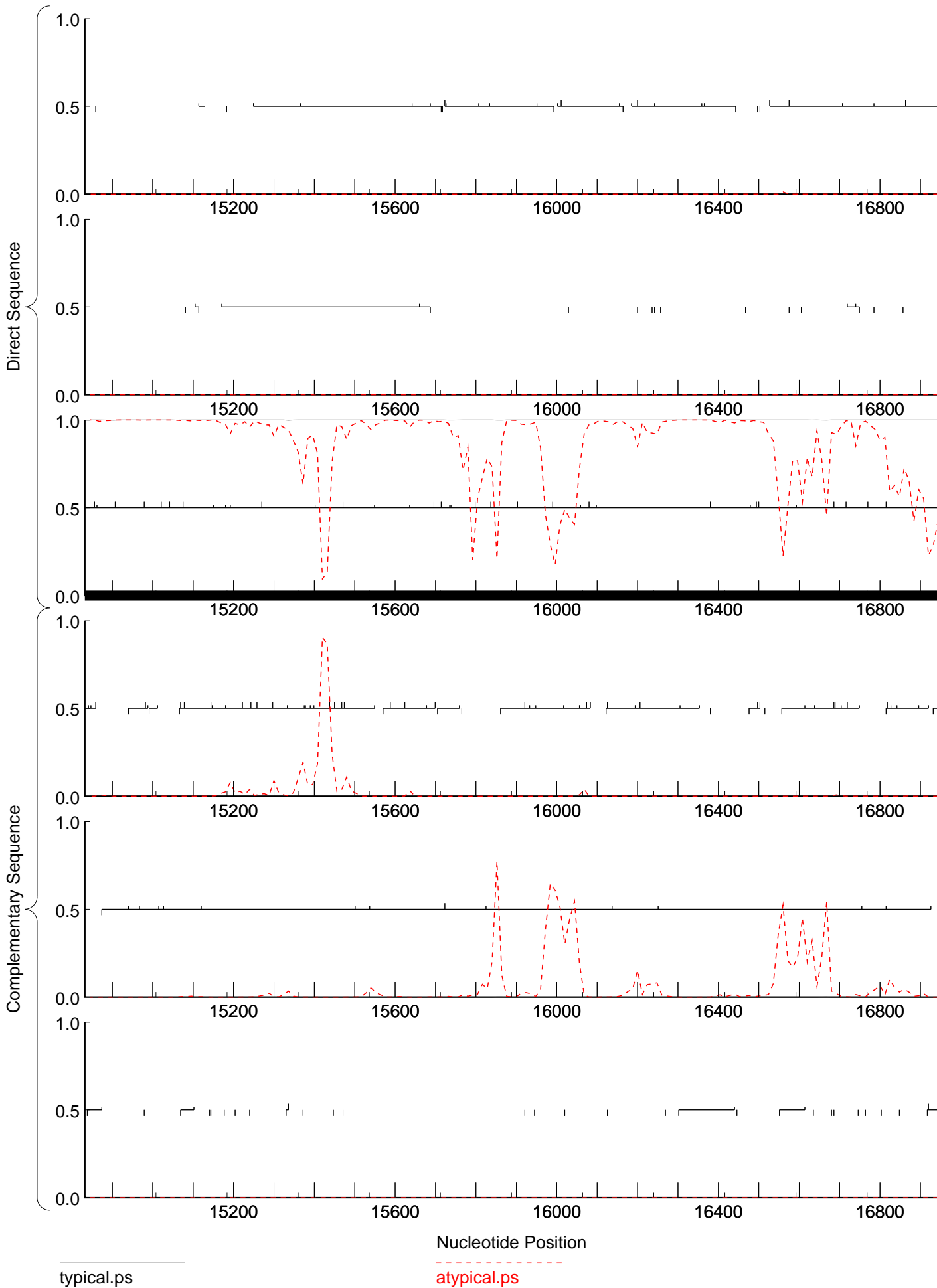






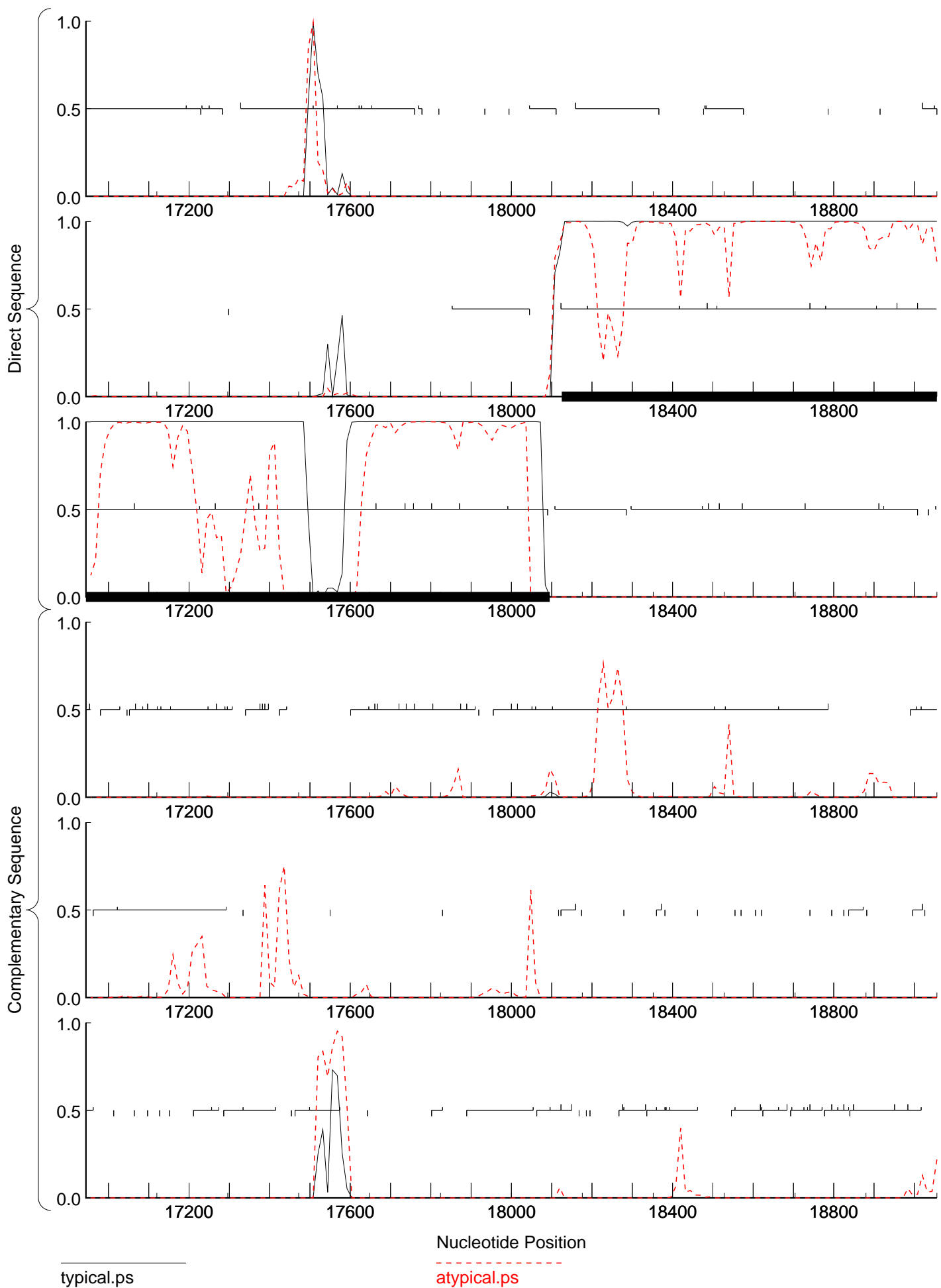






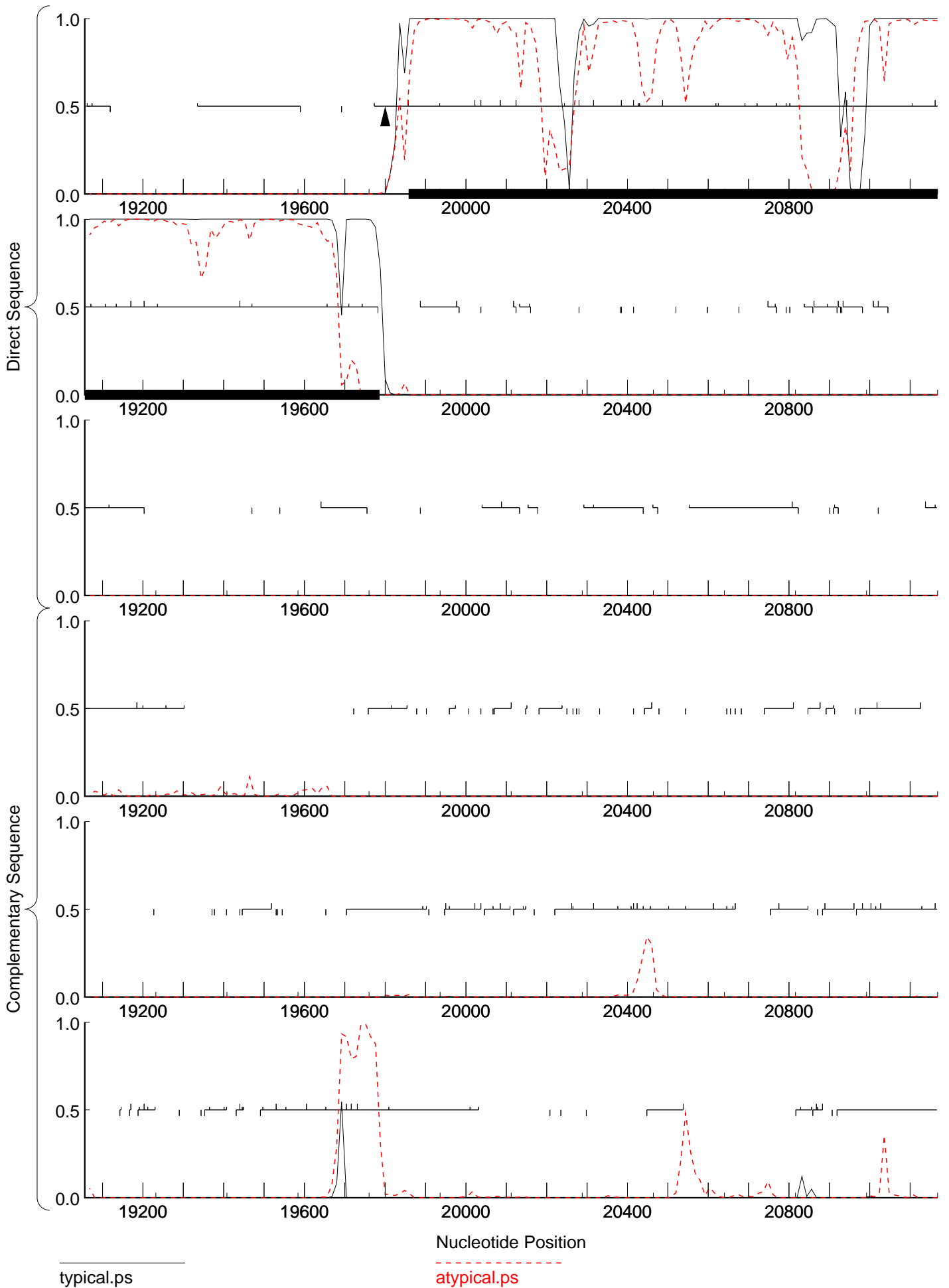
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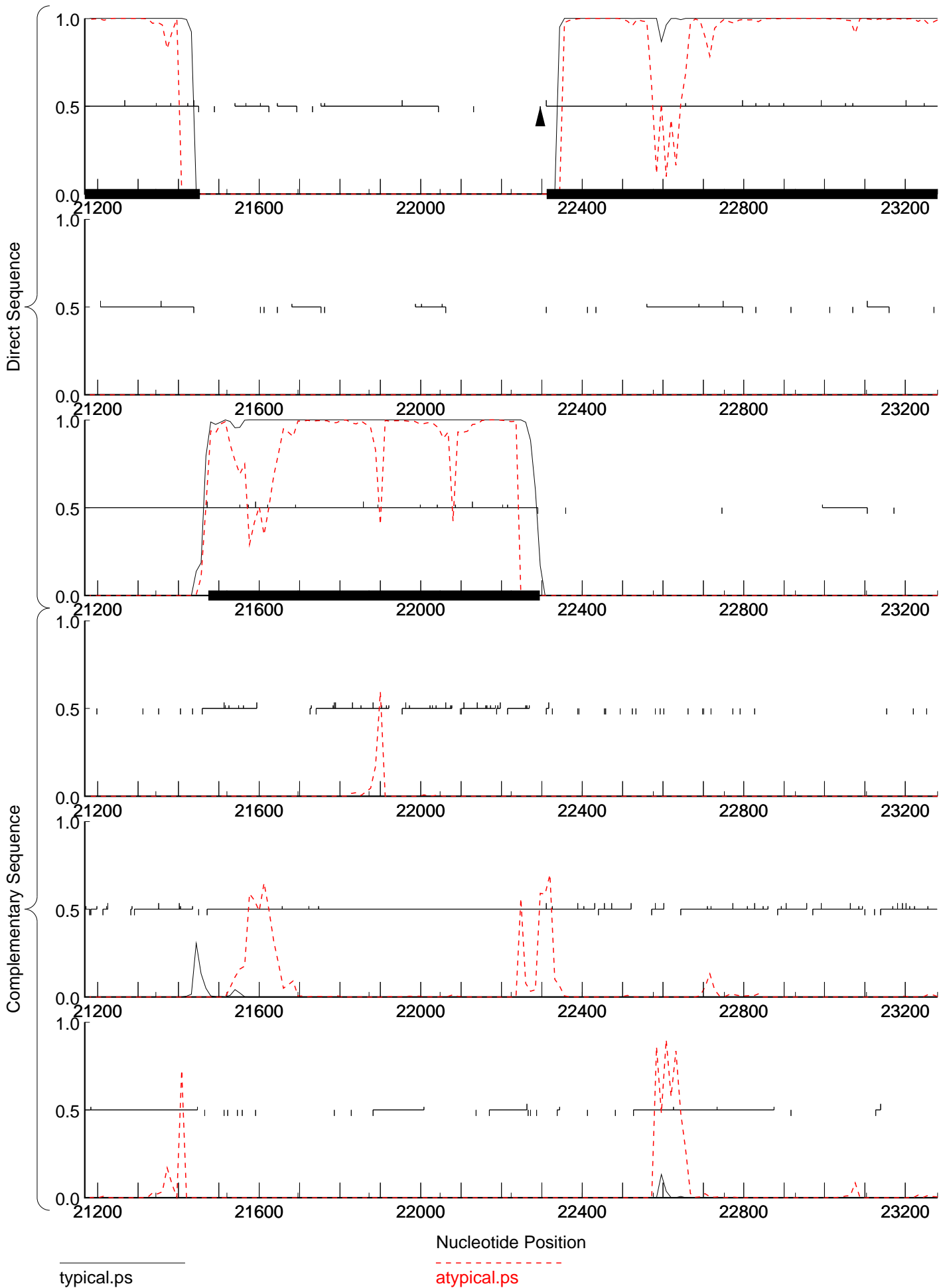
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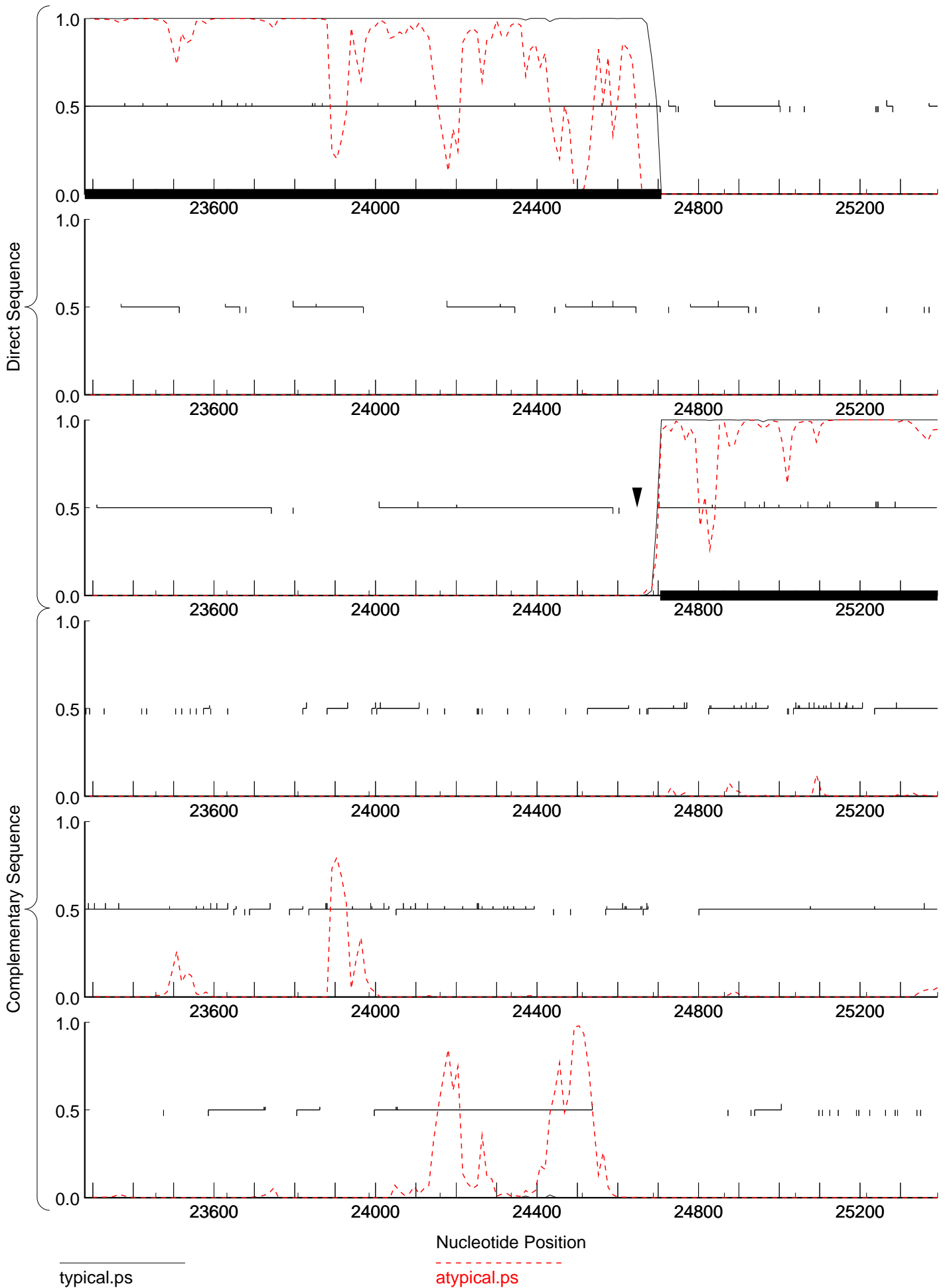


typical.ps

atypical.ps

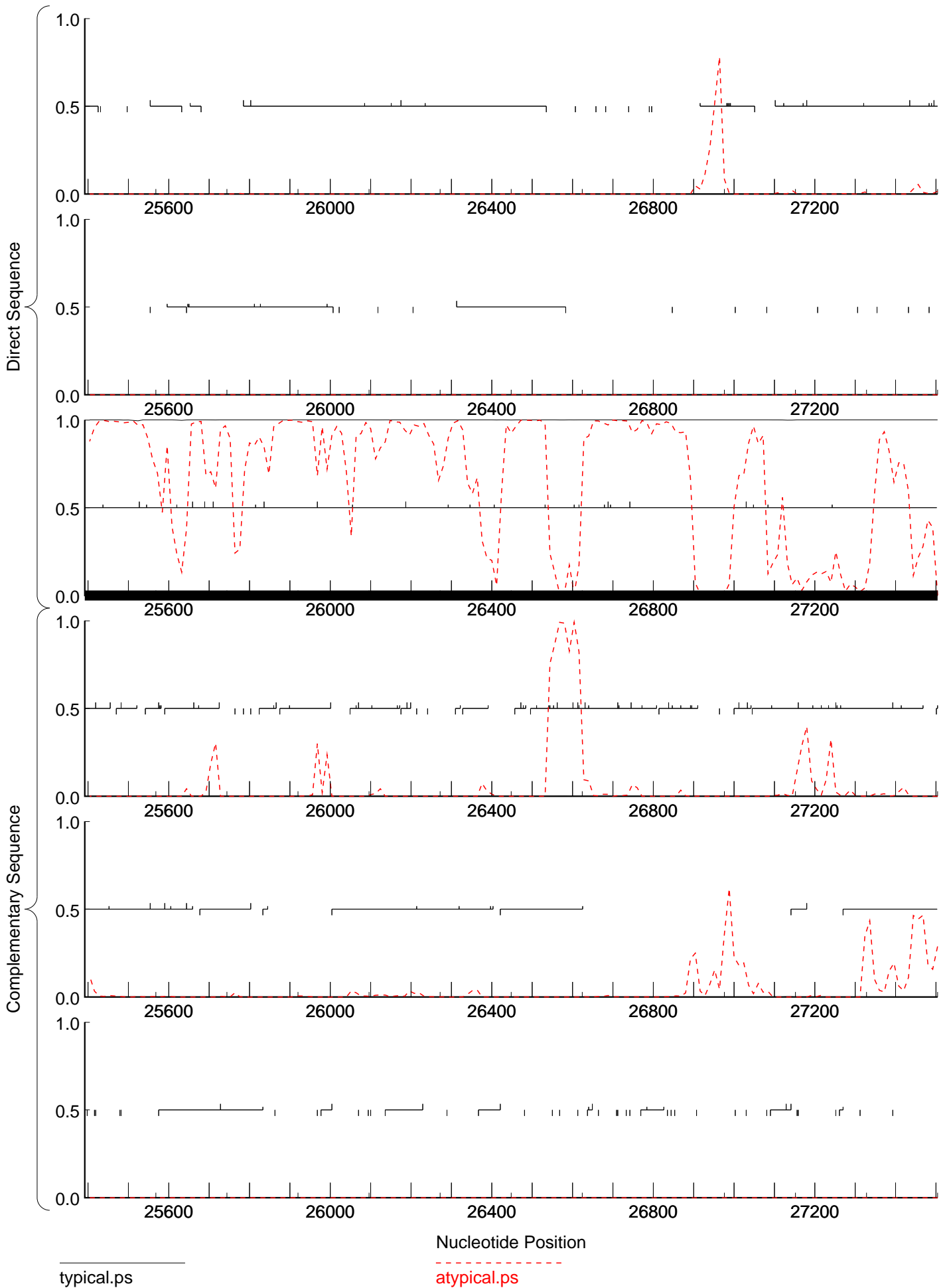






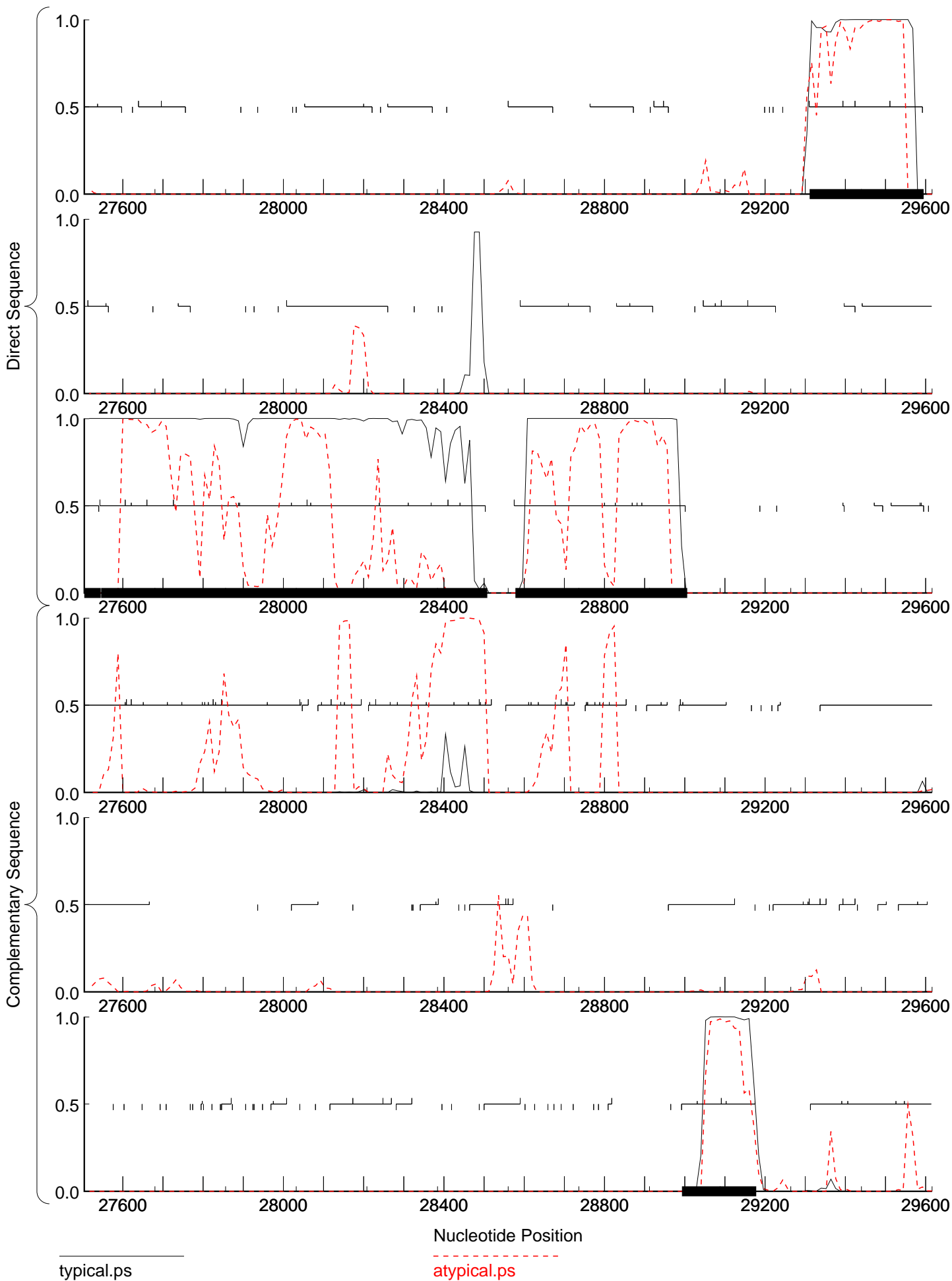
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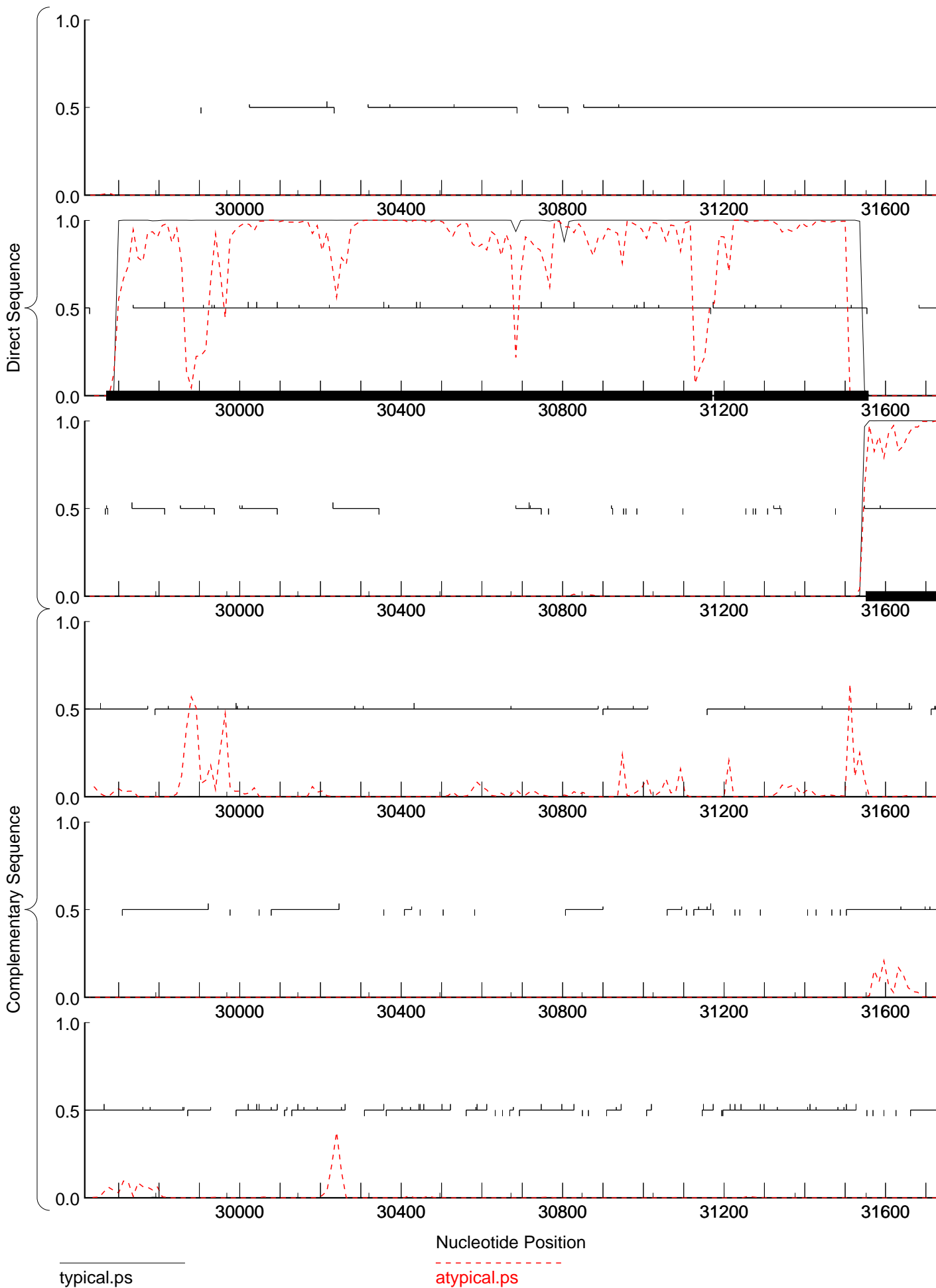
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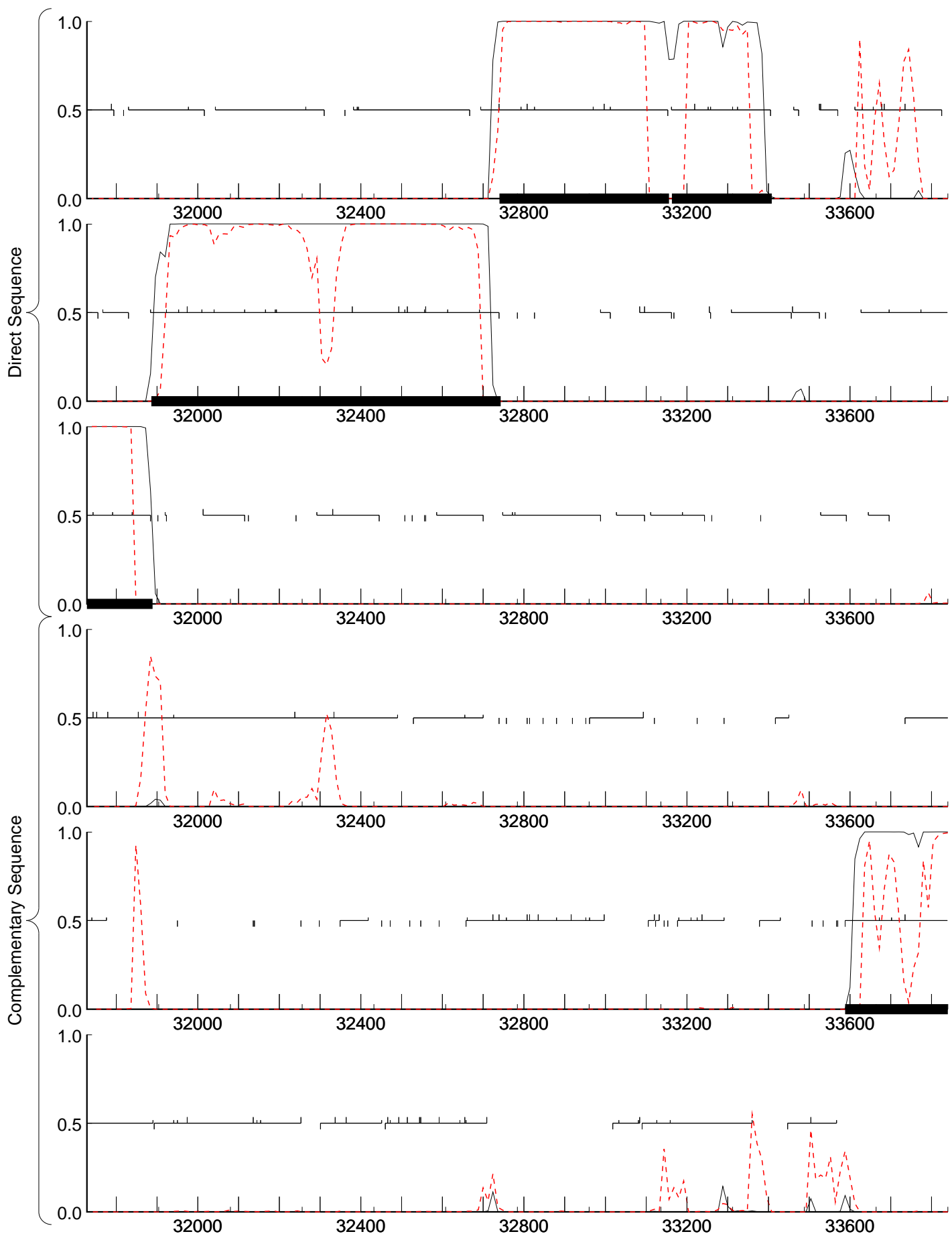


typical.ps

atypical.ps

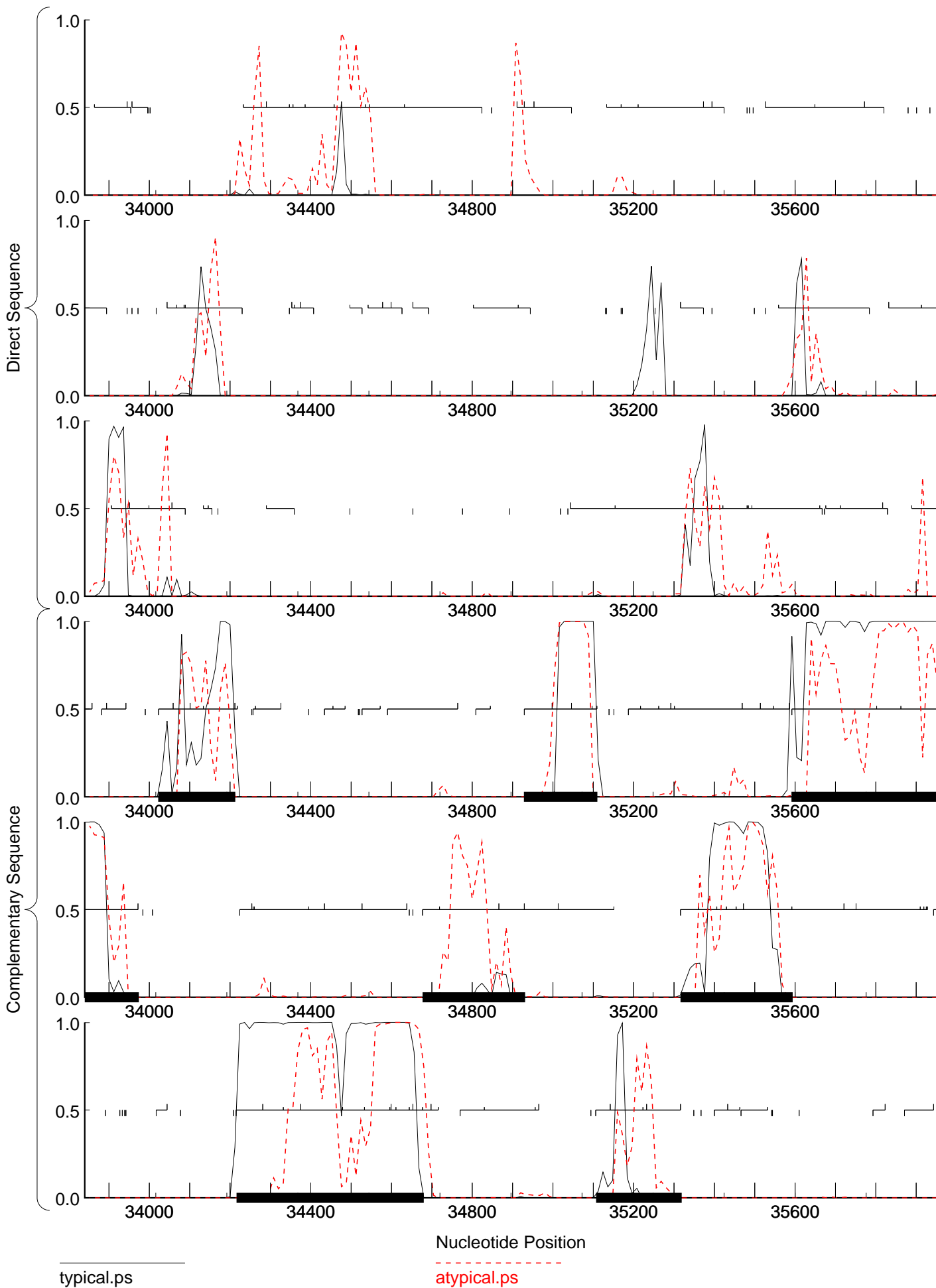


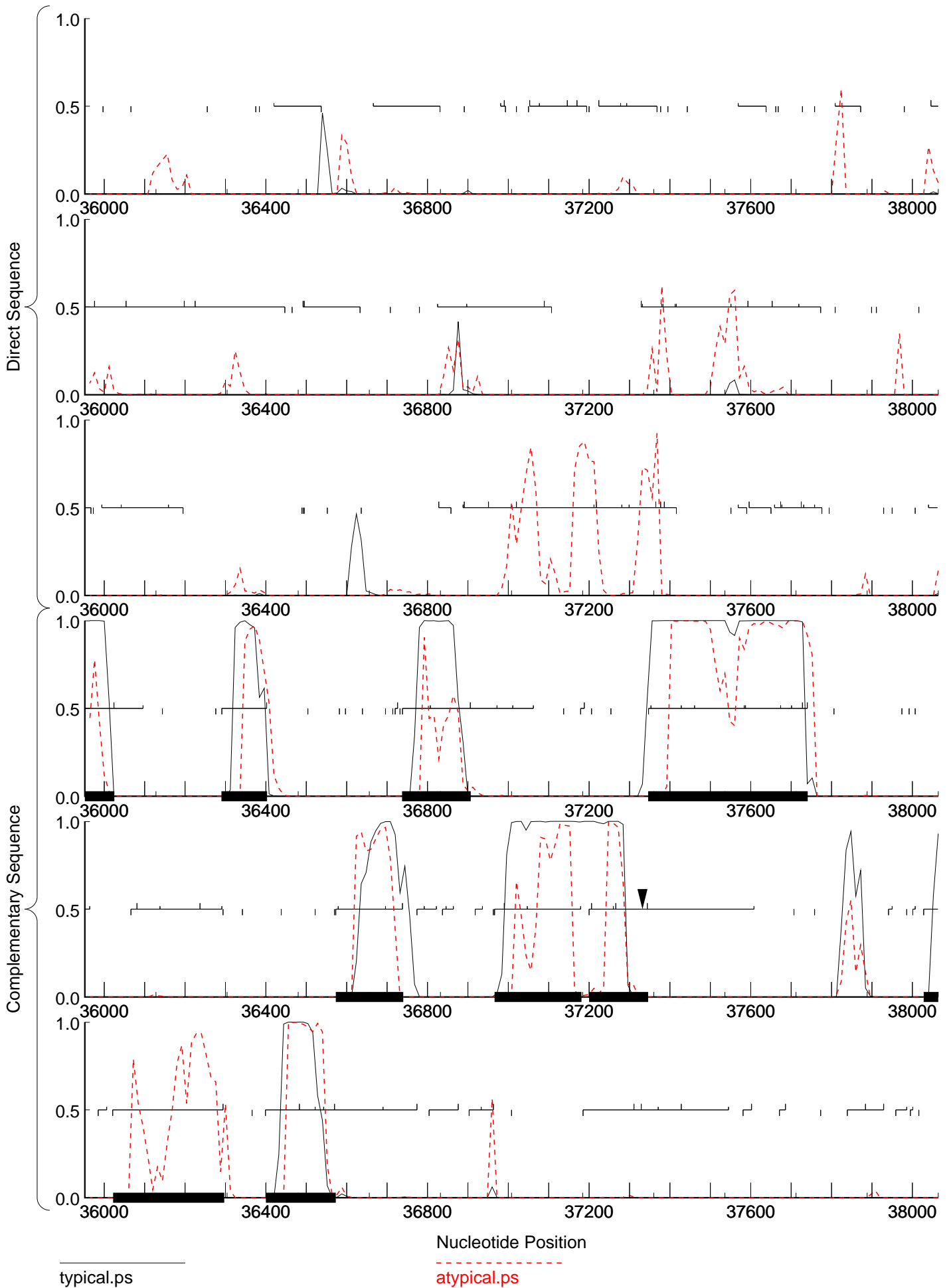


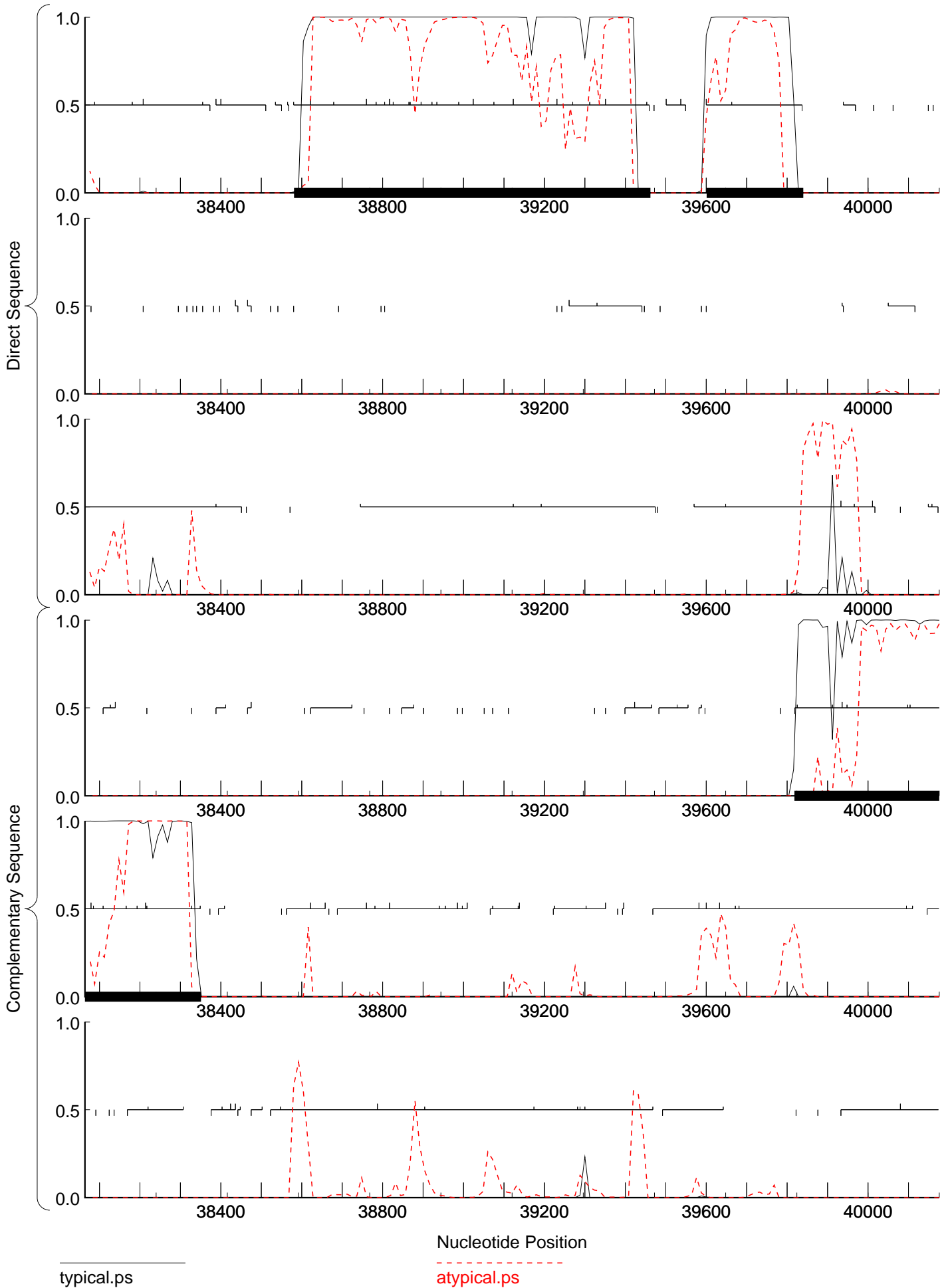


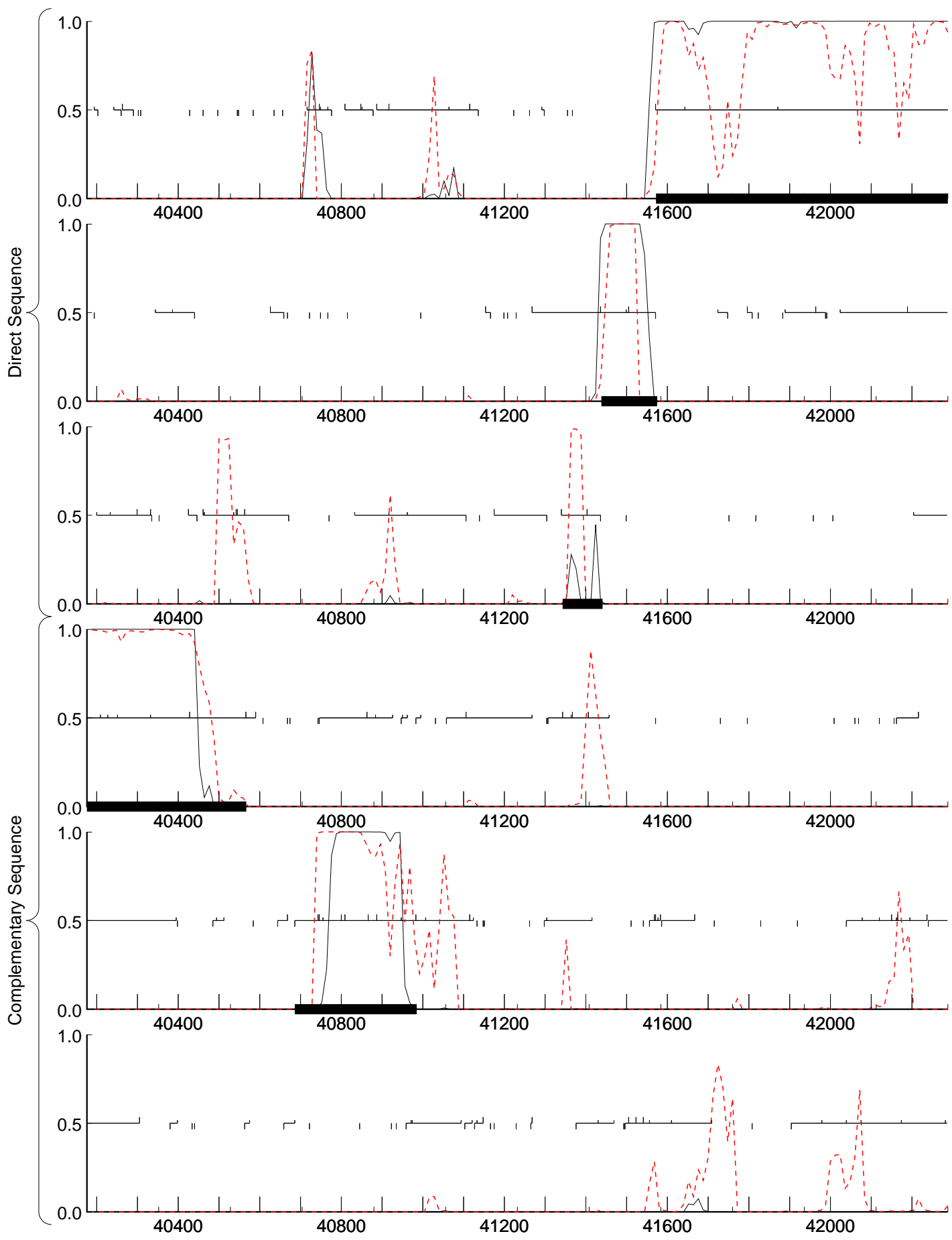
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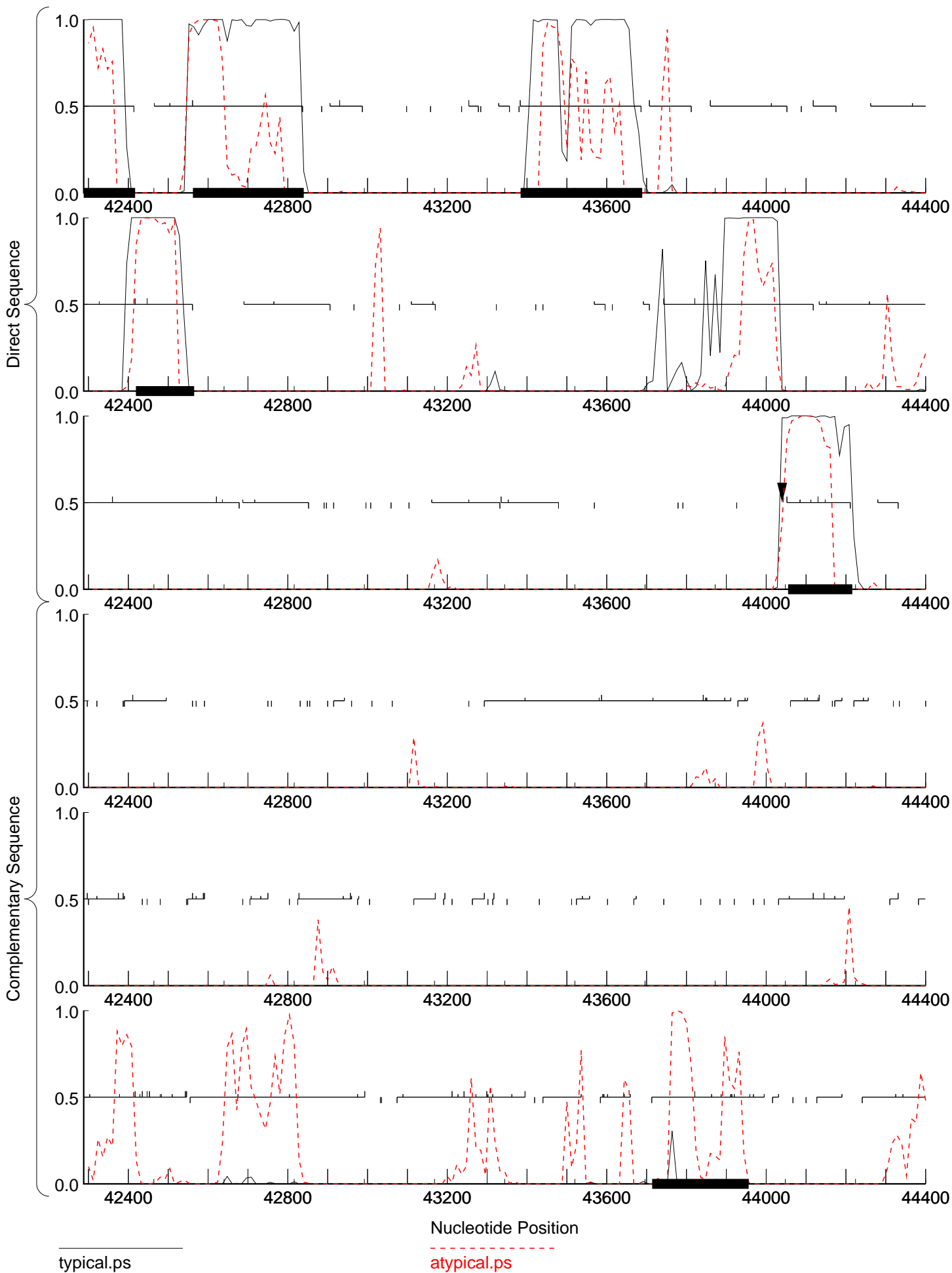






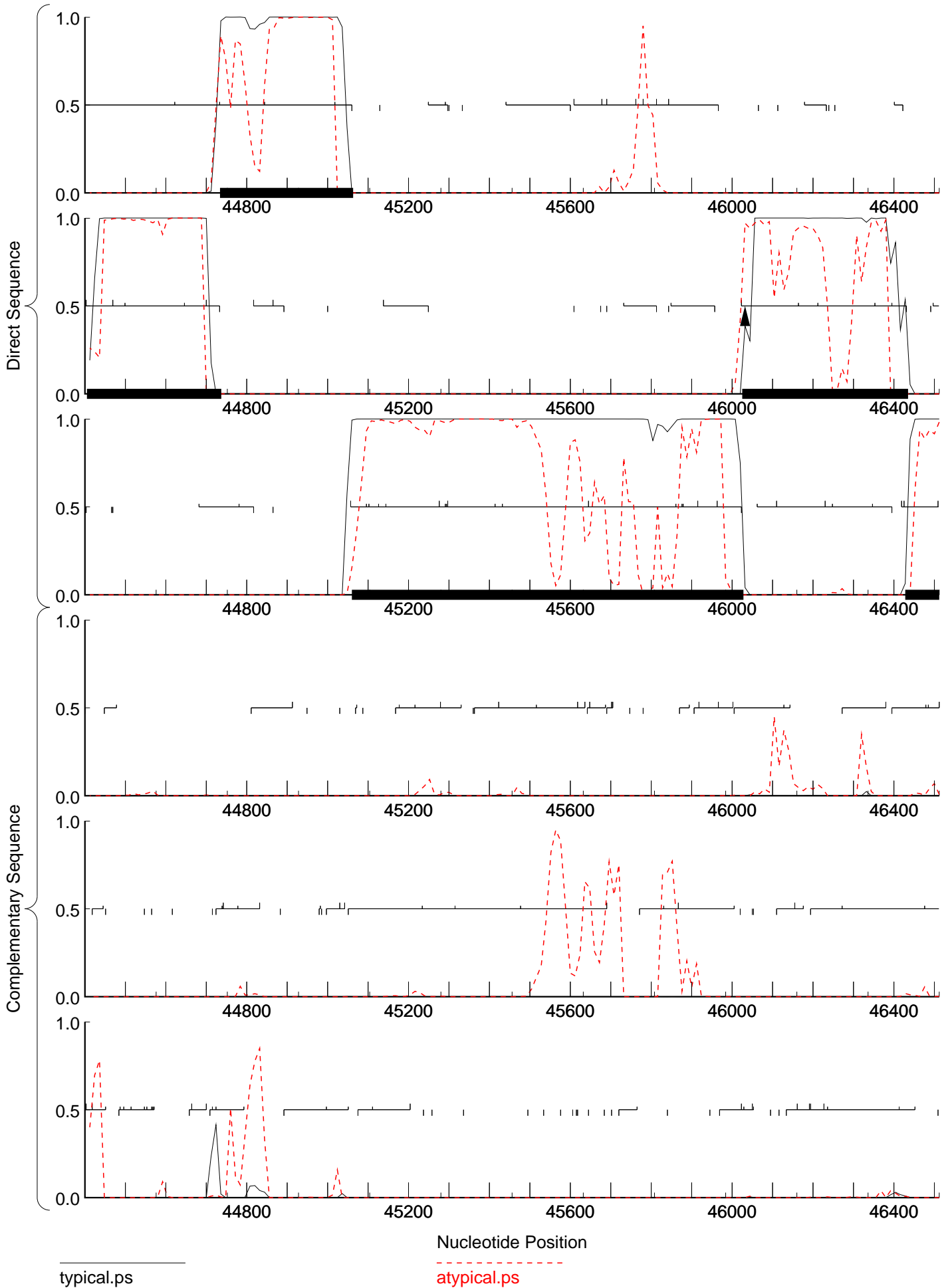
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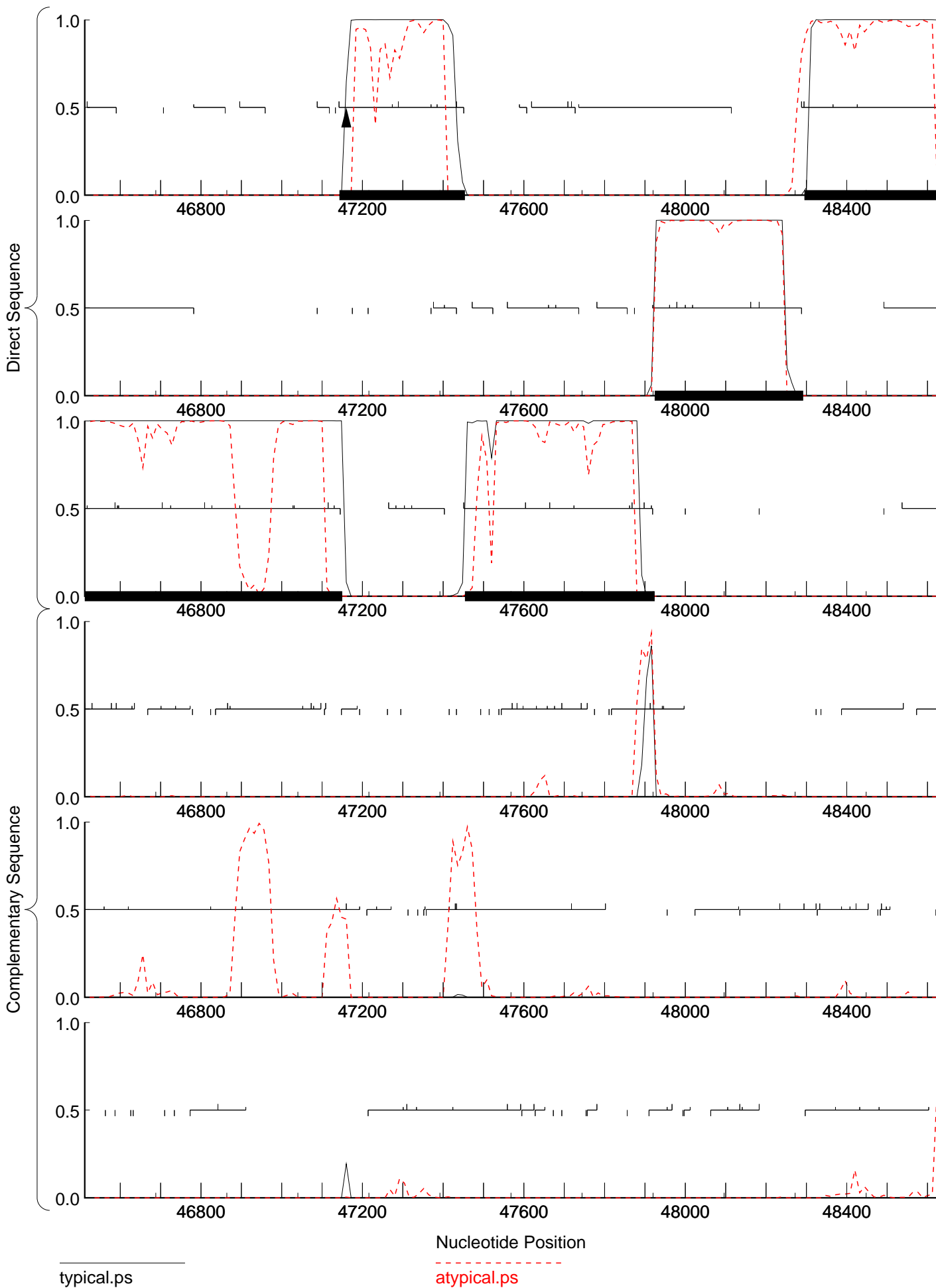
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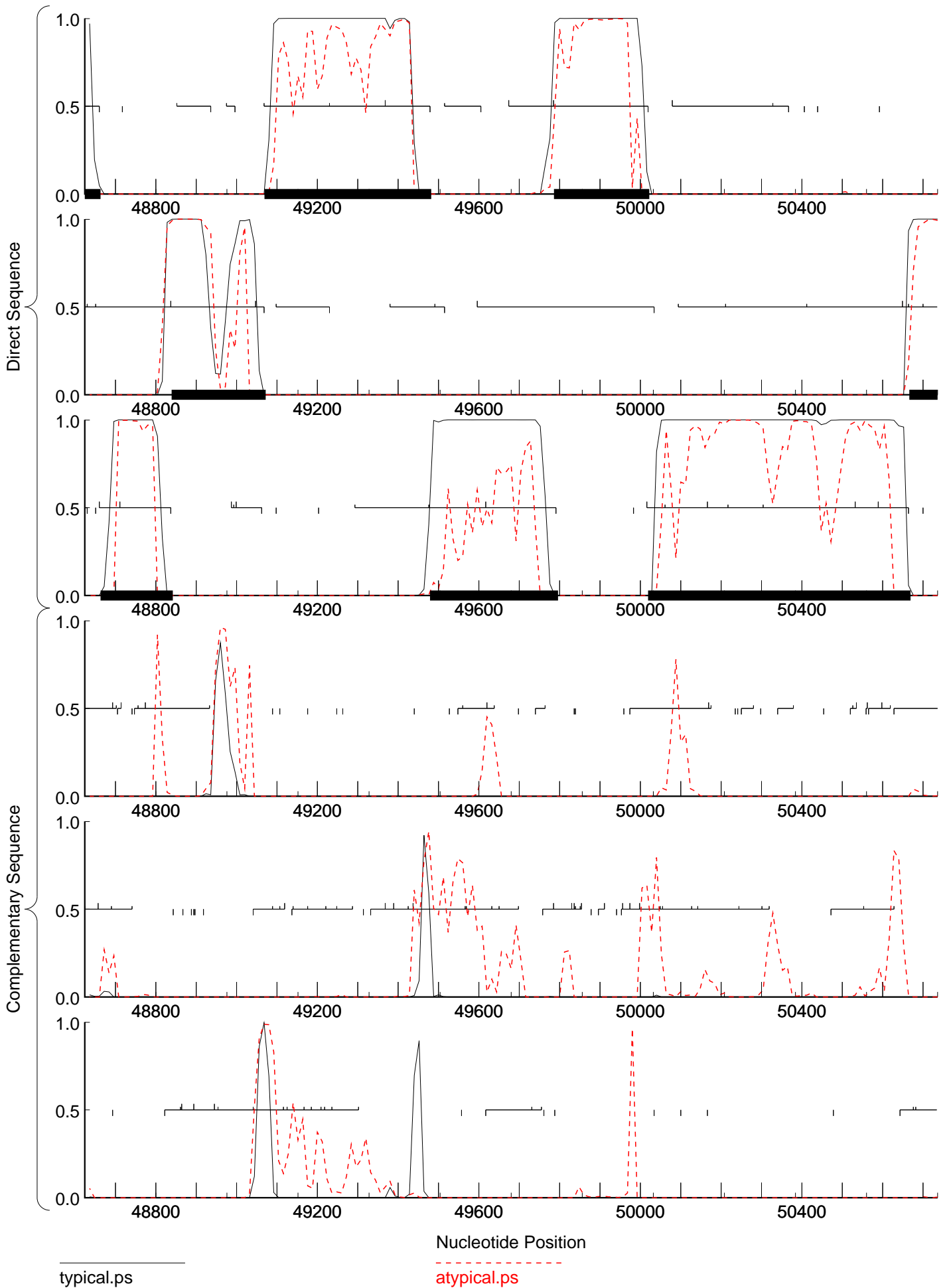
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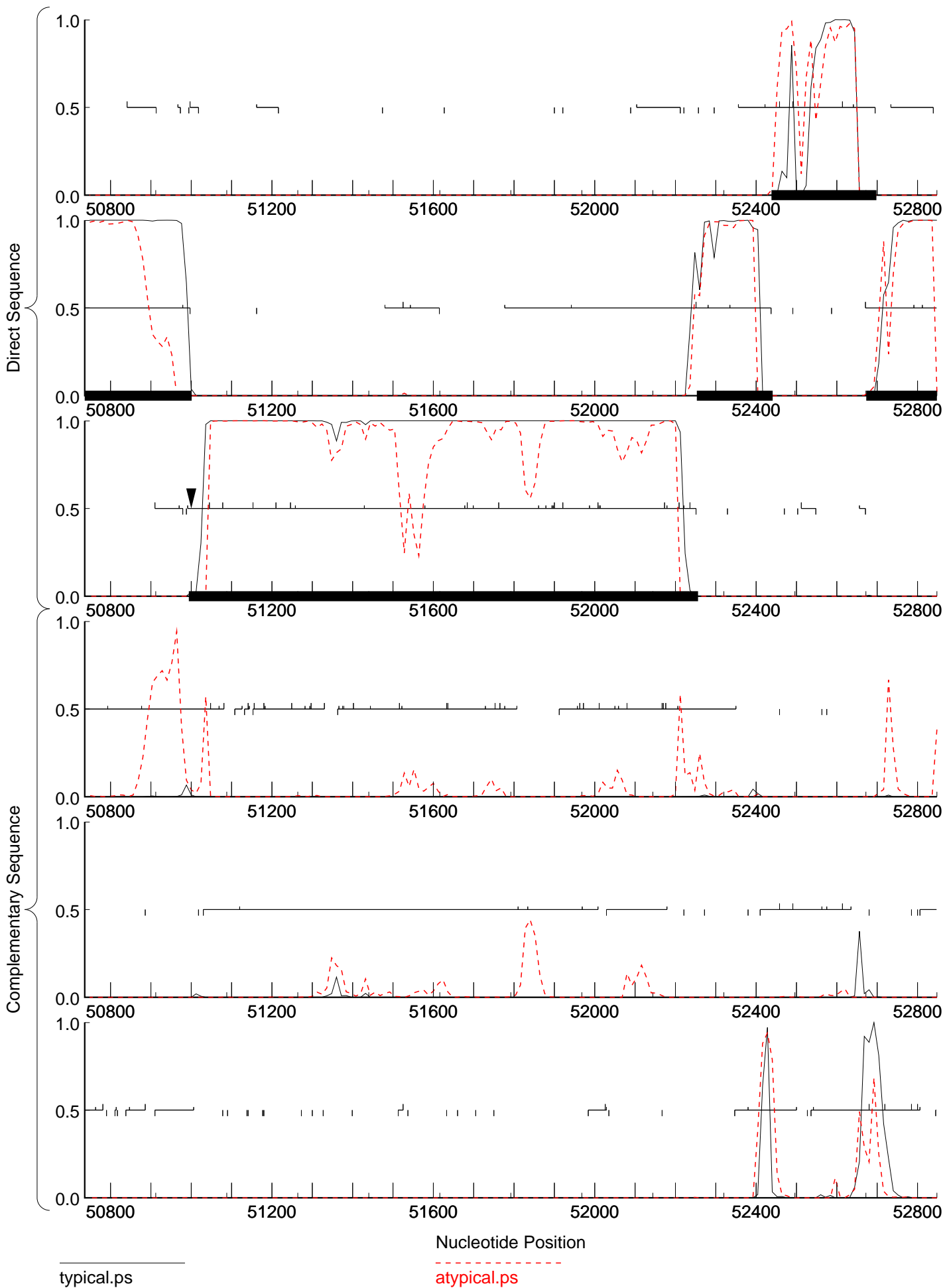


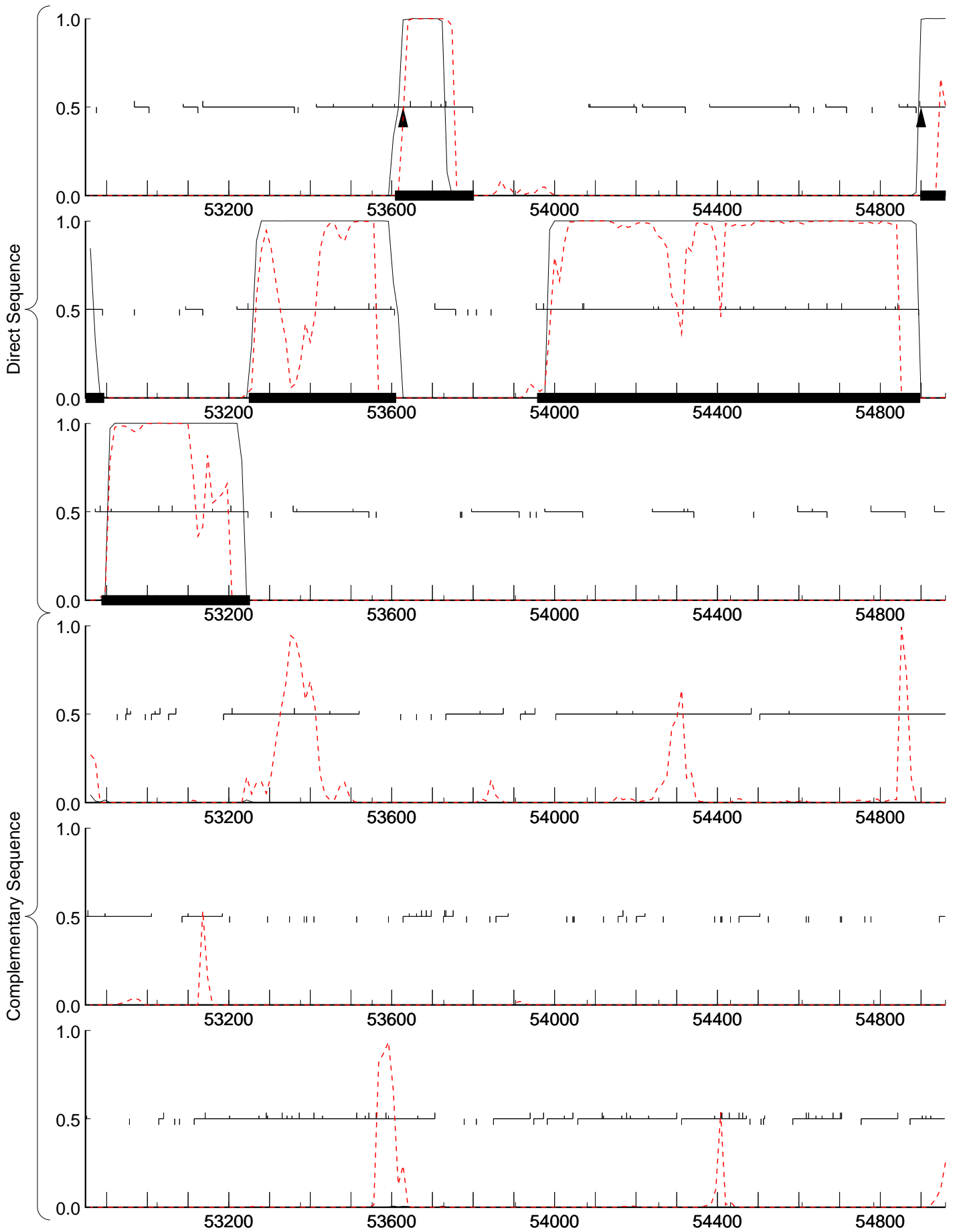
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atypical.ps



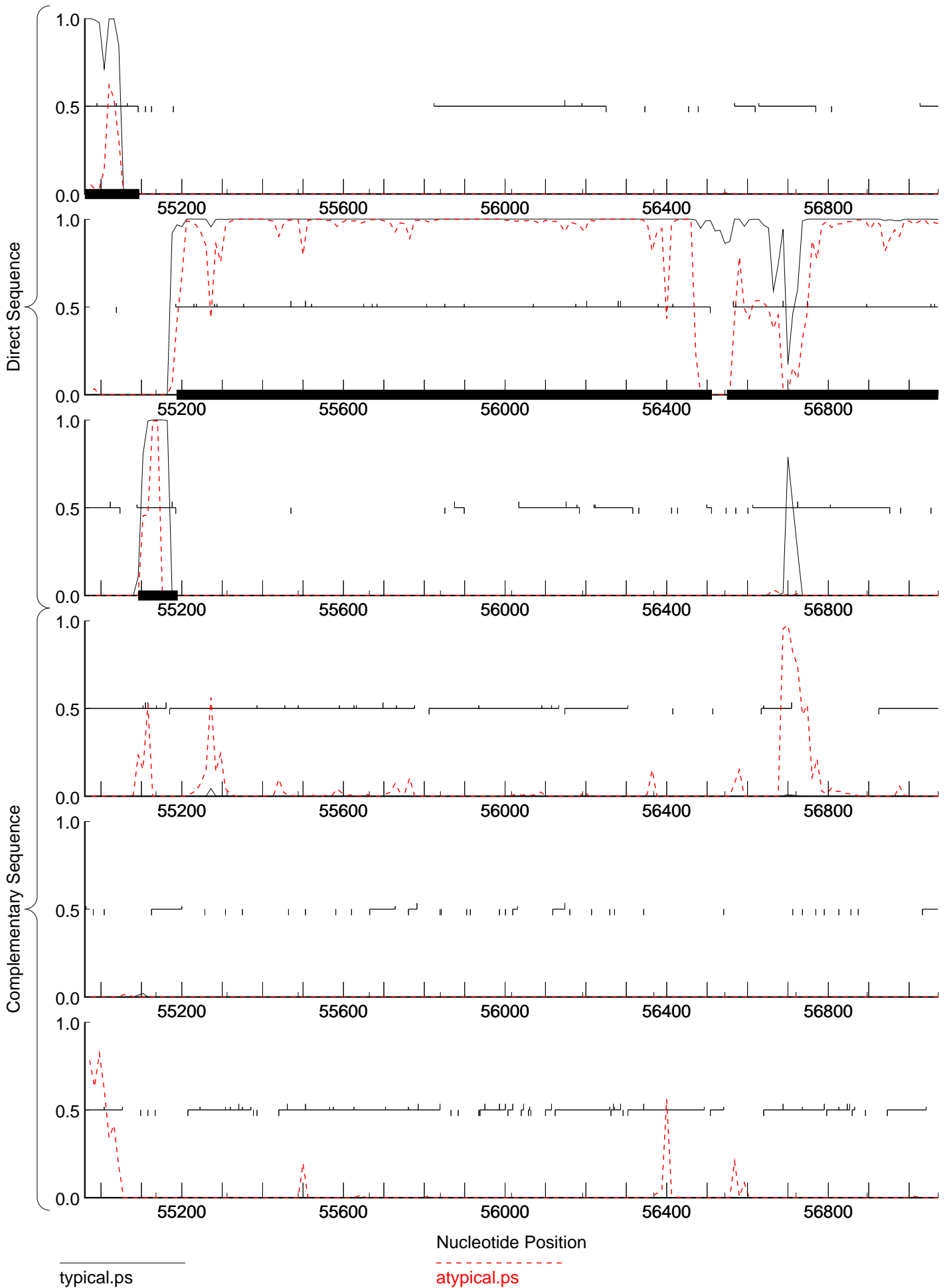






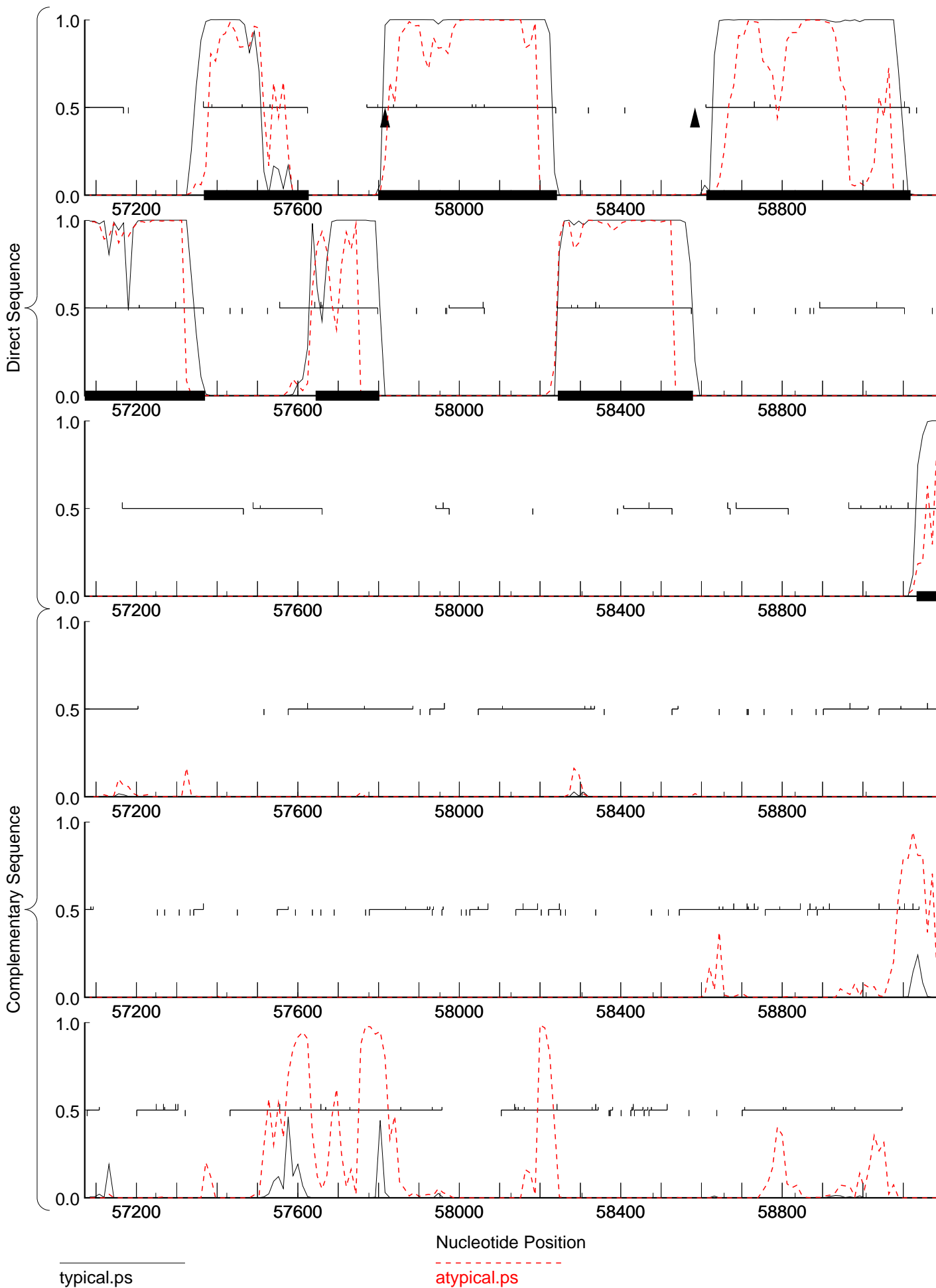
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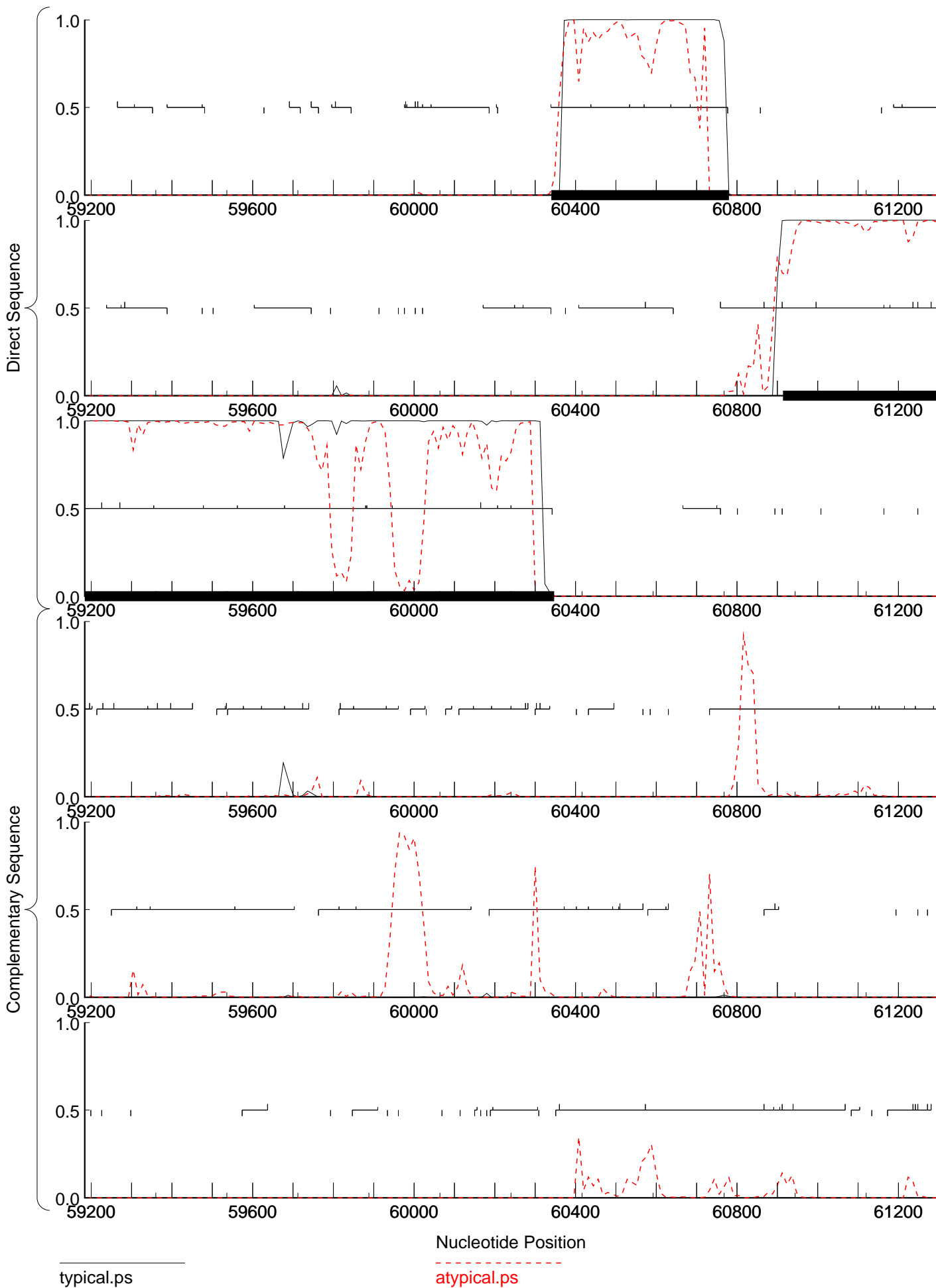
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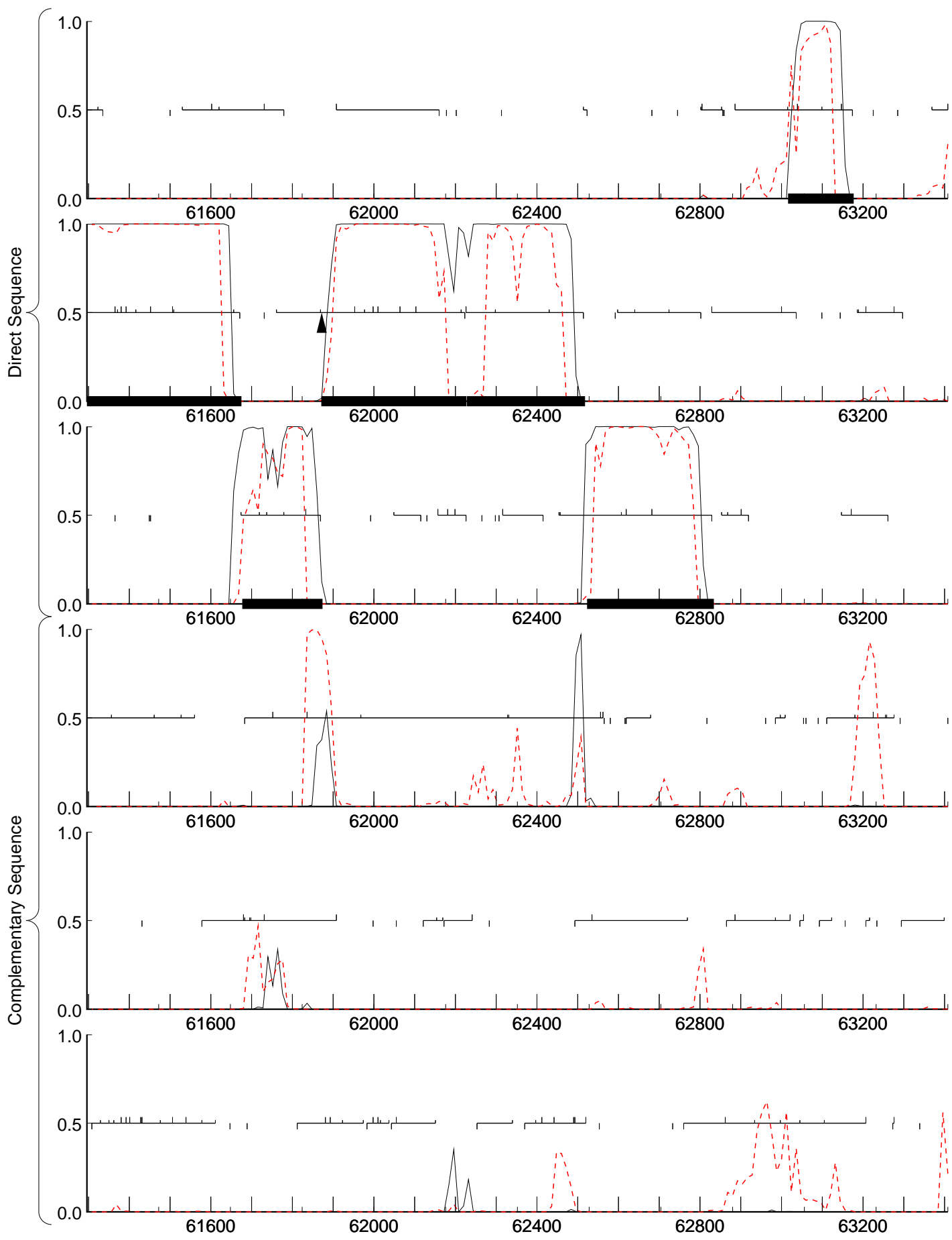


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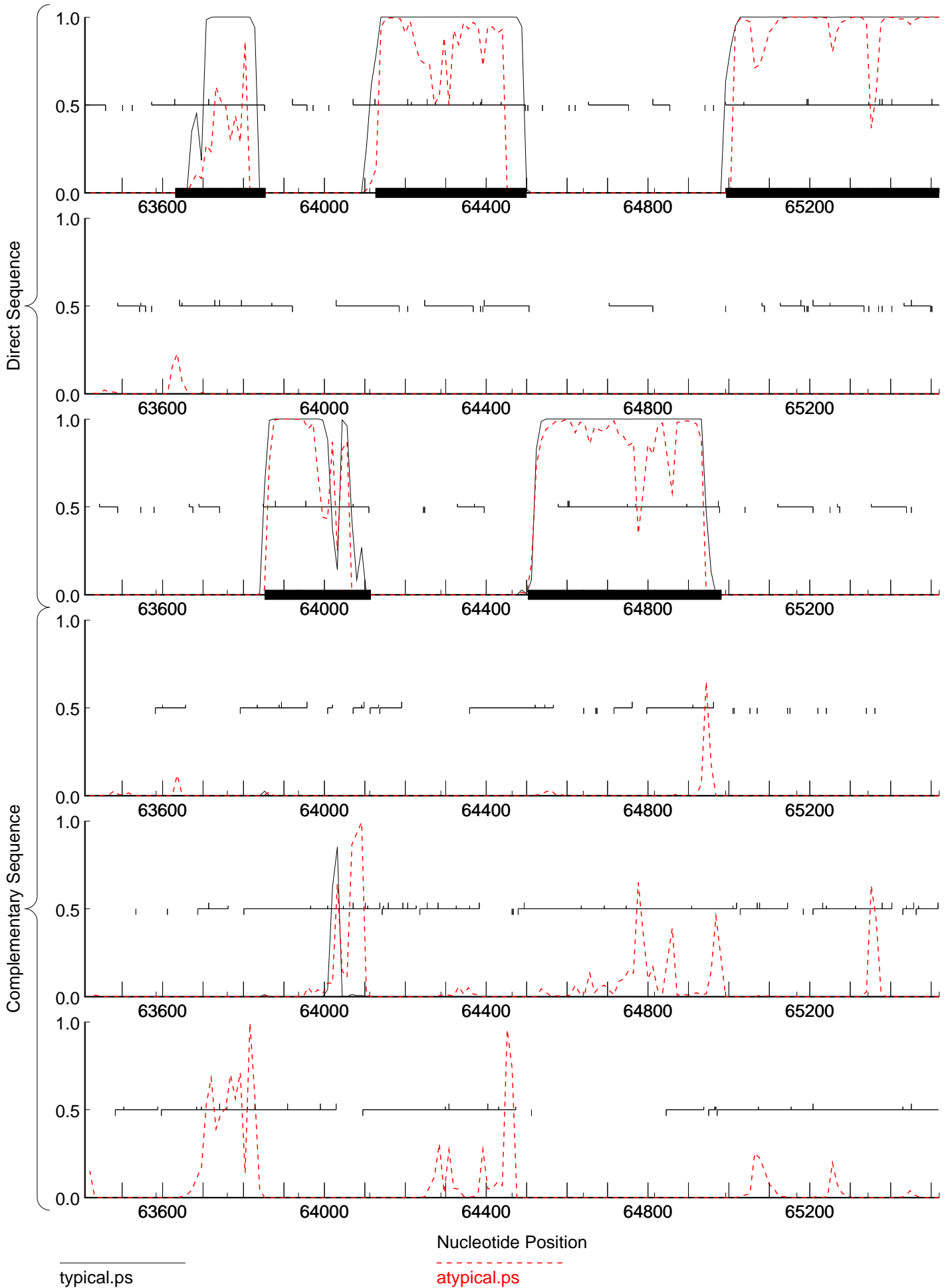


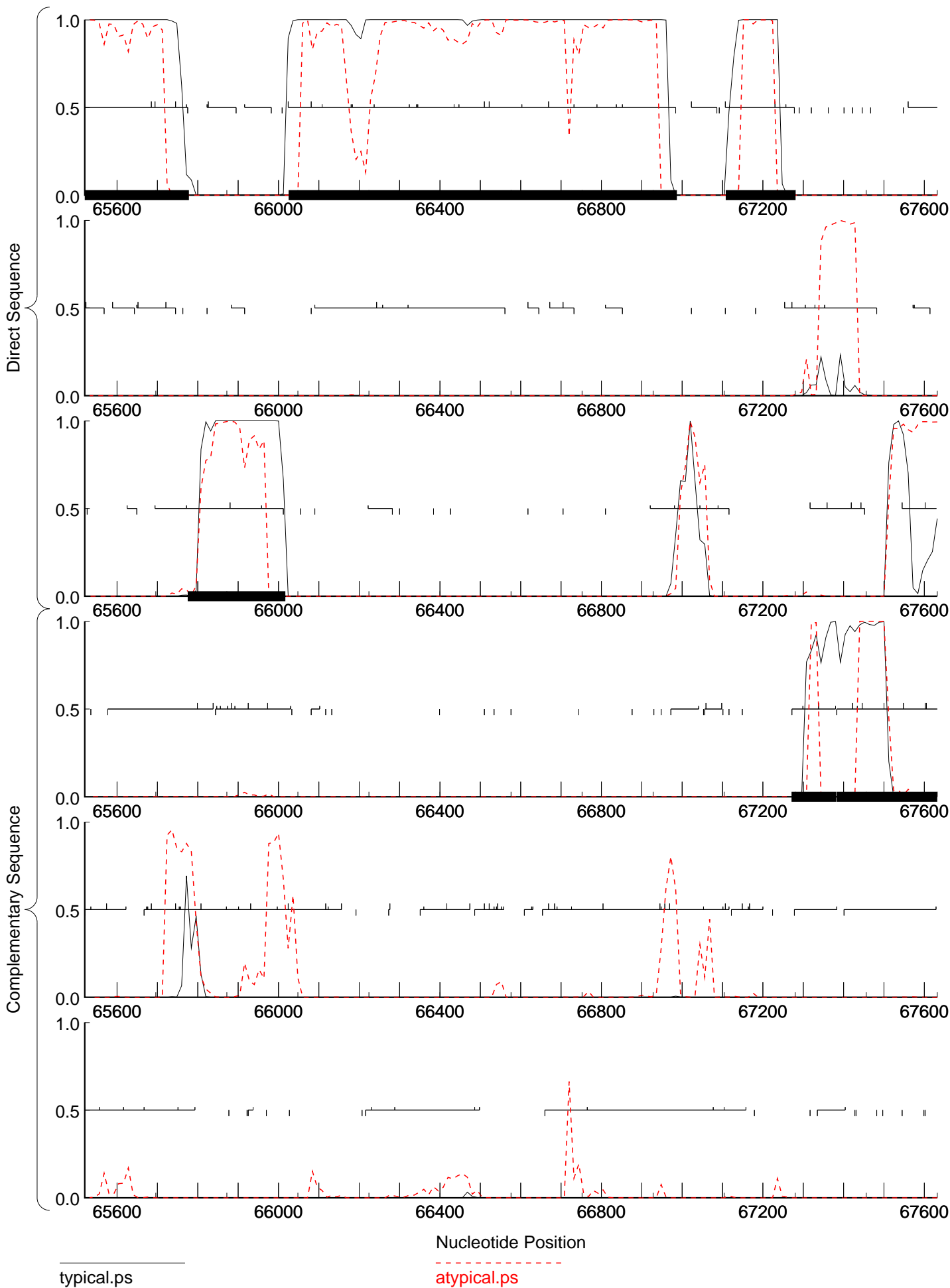


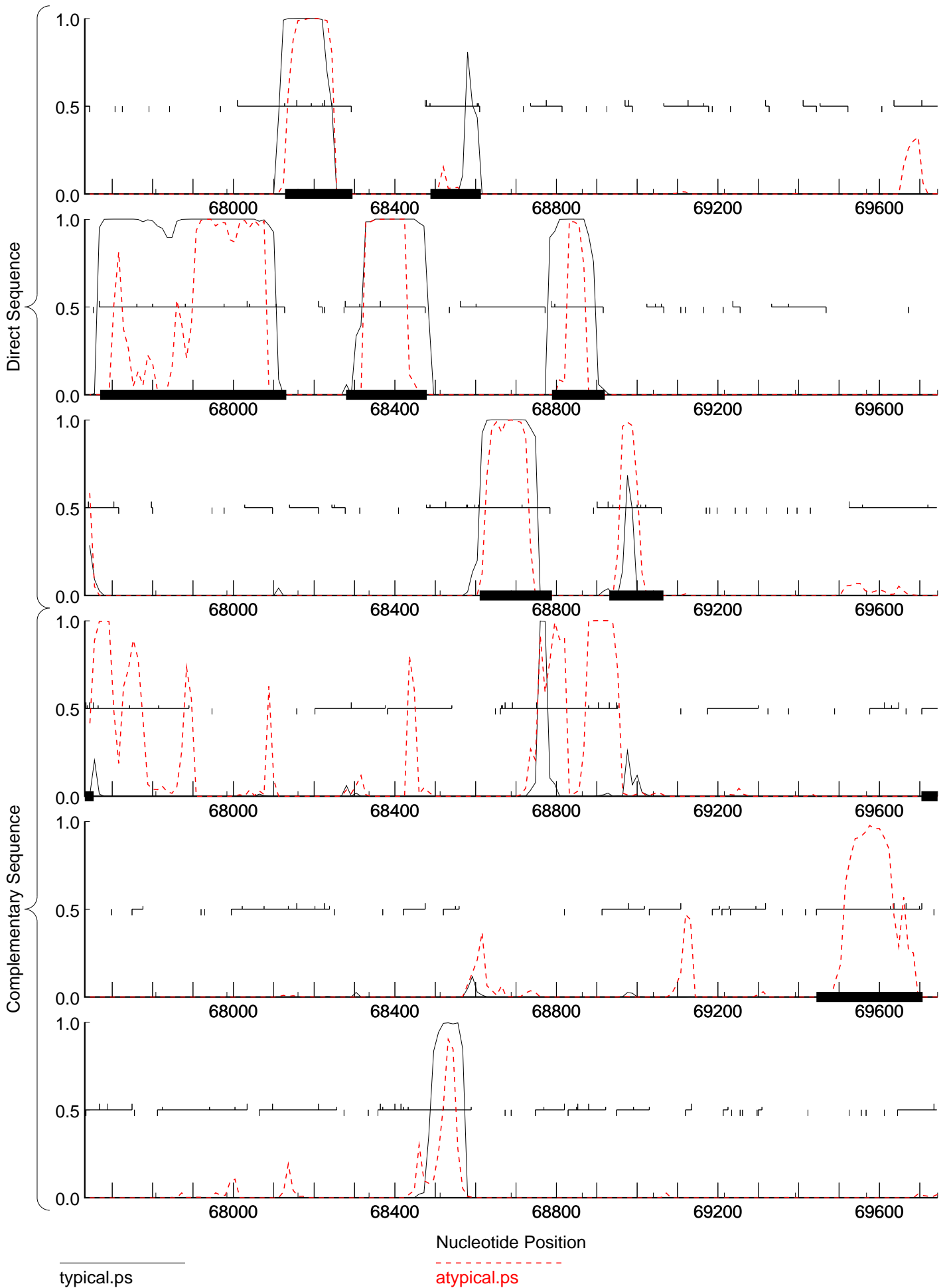


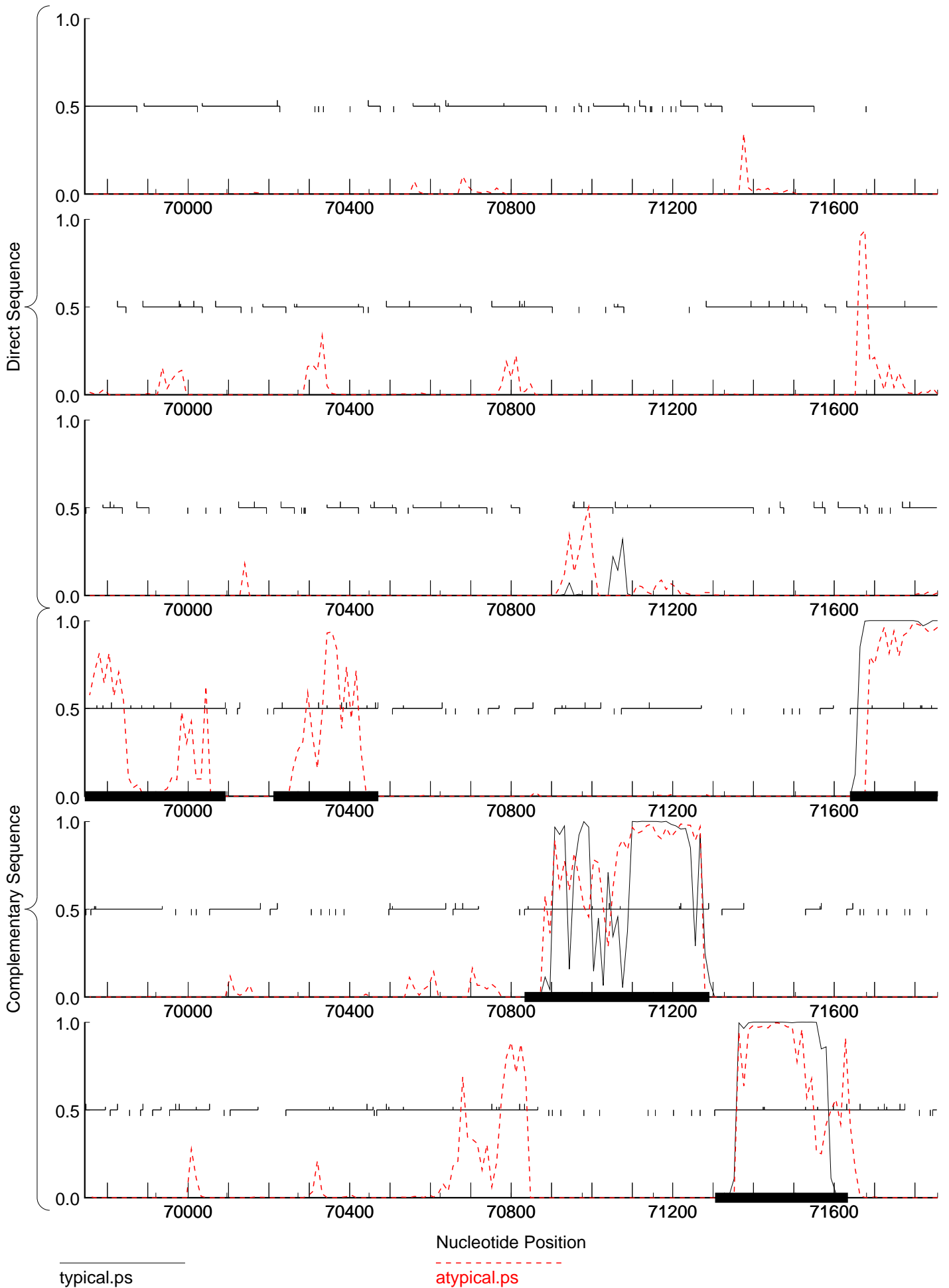
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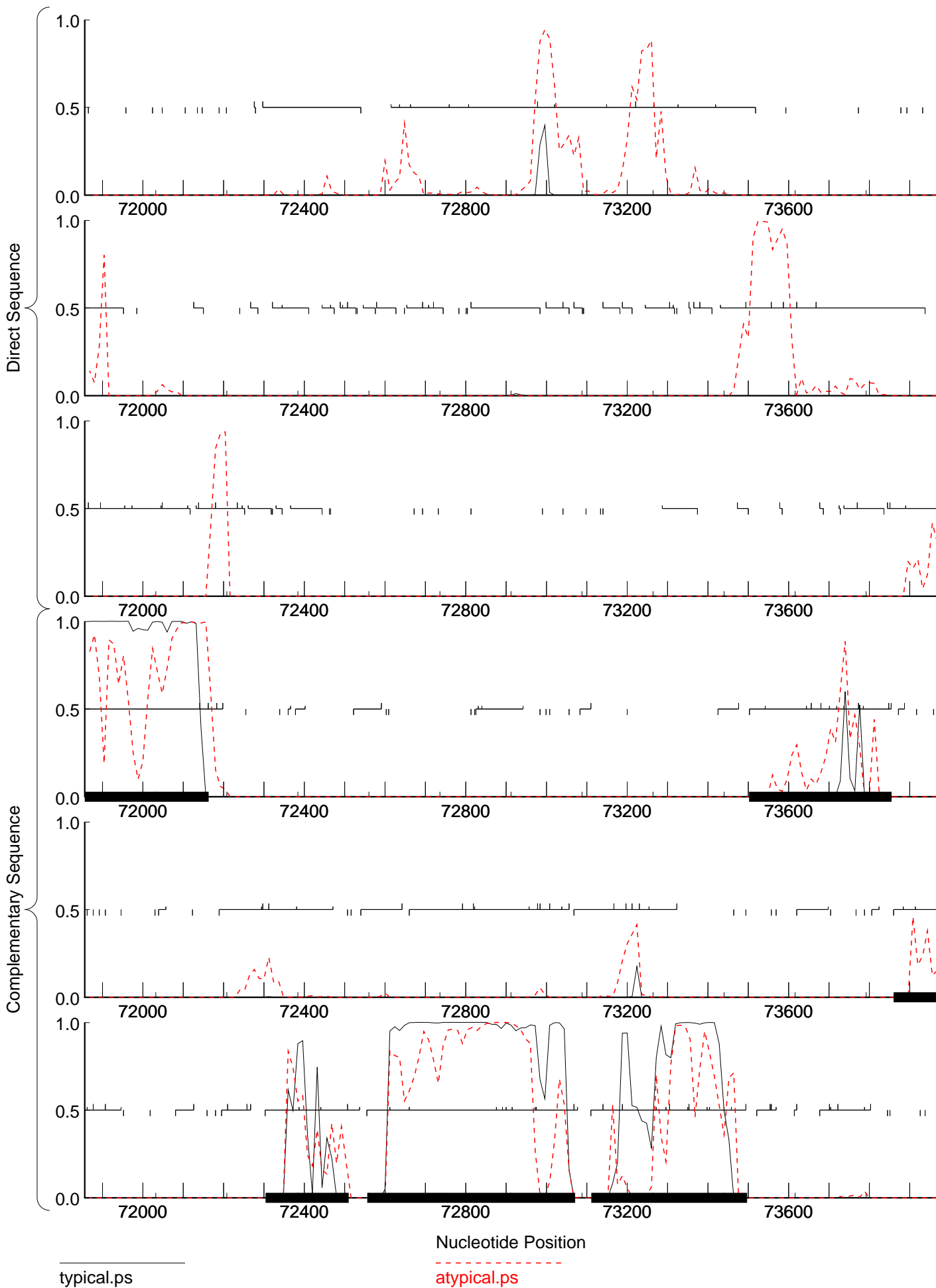
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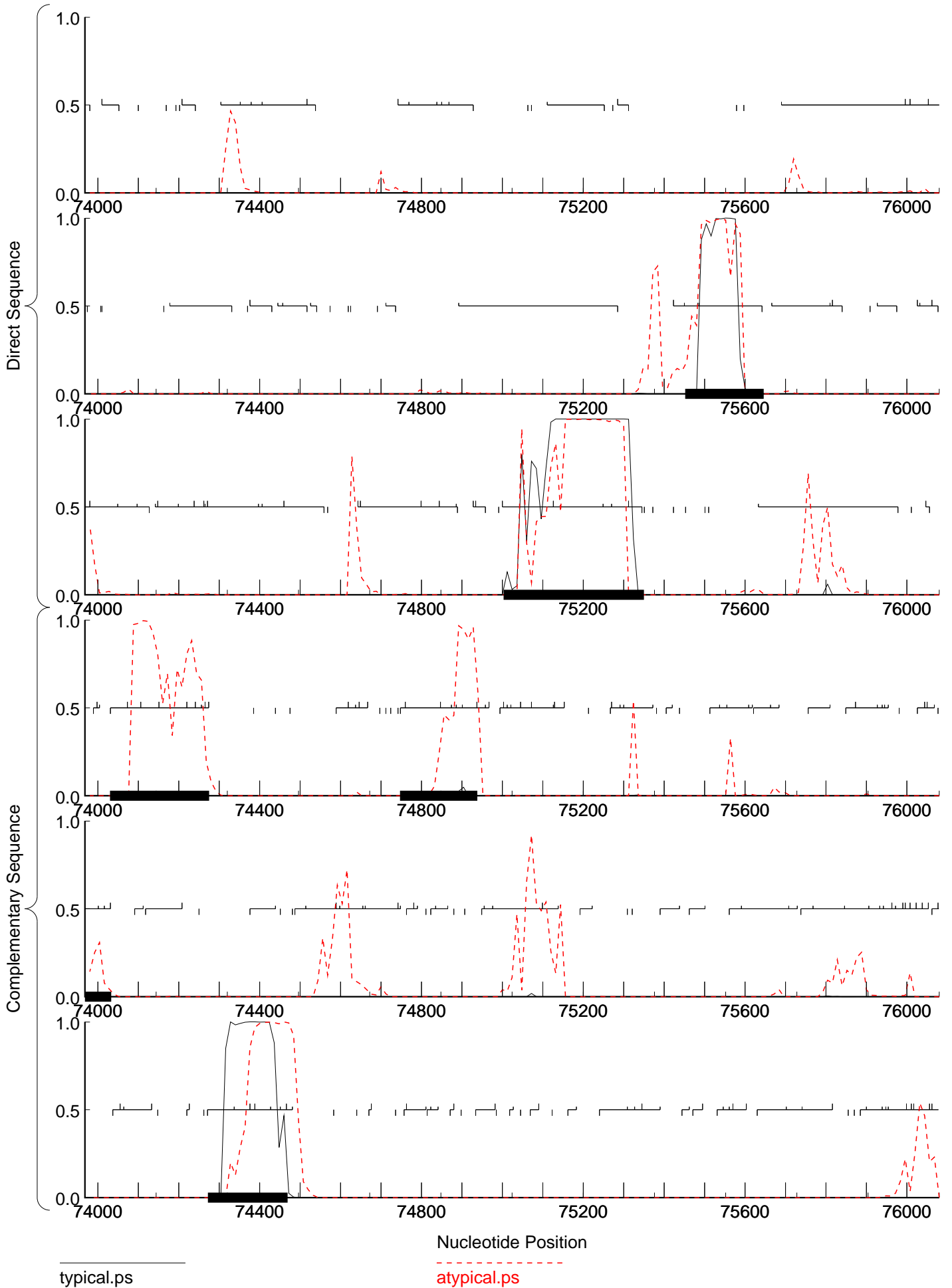






typical.ps

atypical.ps



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