

# GeneMark

Version 2.5p (09.08.06)

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## PROGRAM INFORMATION

Sequence : Mycobacterium phage Paola complete sequence, 61535 bp including 11-base 3' overhang (CTCAGT  
Analysis Date : 11/19/17 at 19:05:40  
Pages : 31  
Sequence Length : 61535 bp  
GC Content : 65.01%

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,

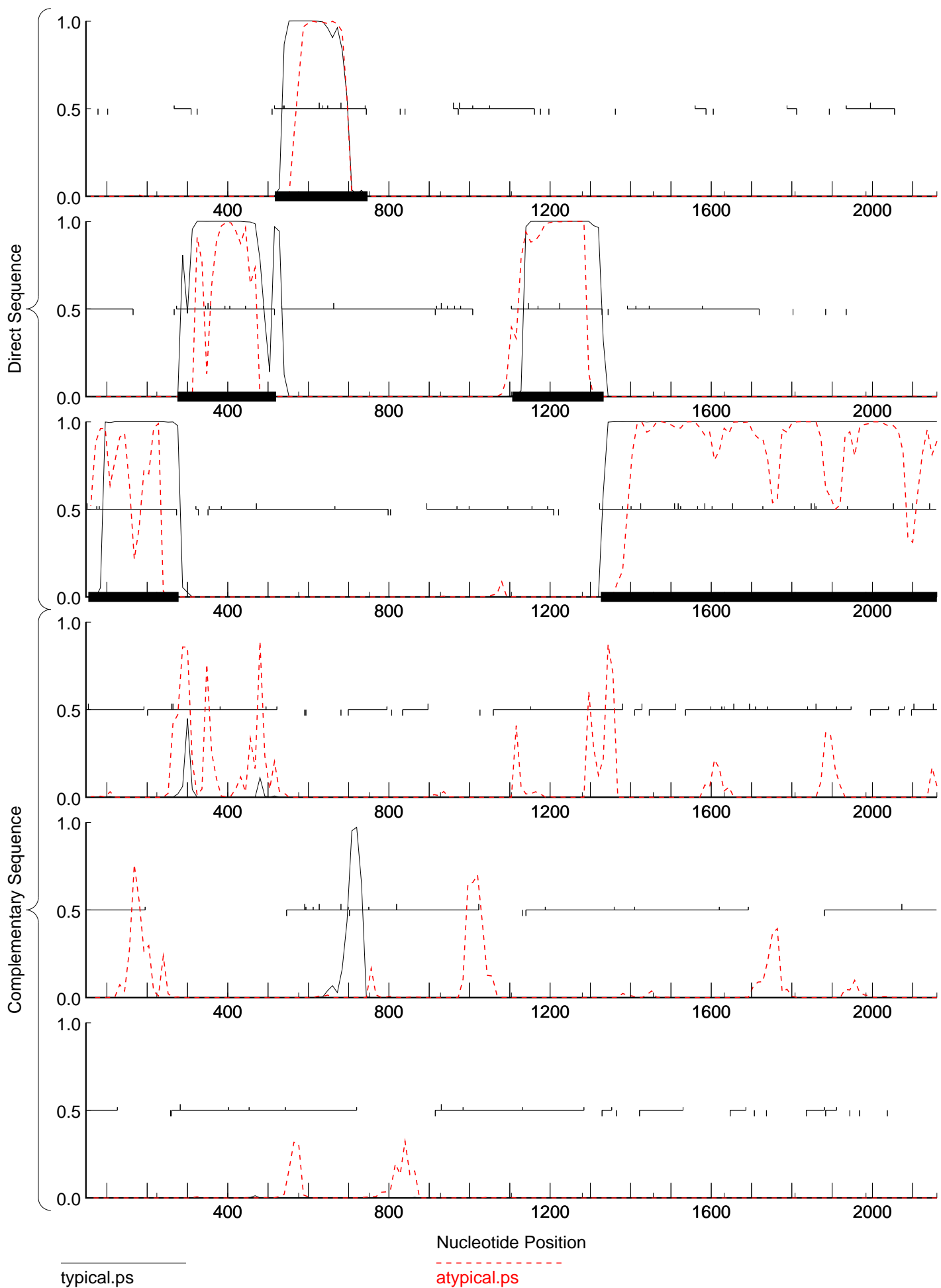
## MATRIX INFORMATION

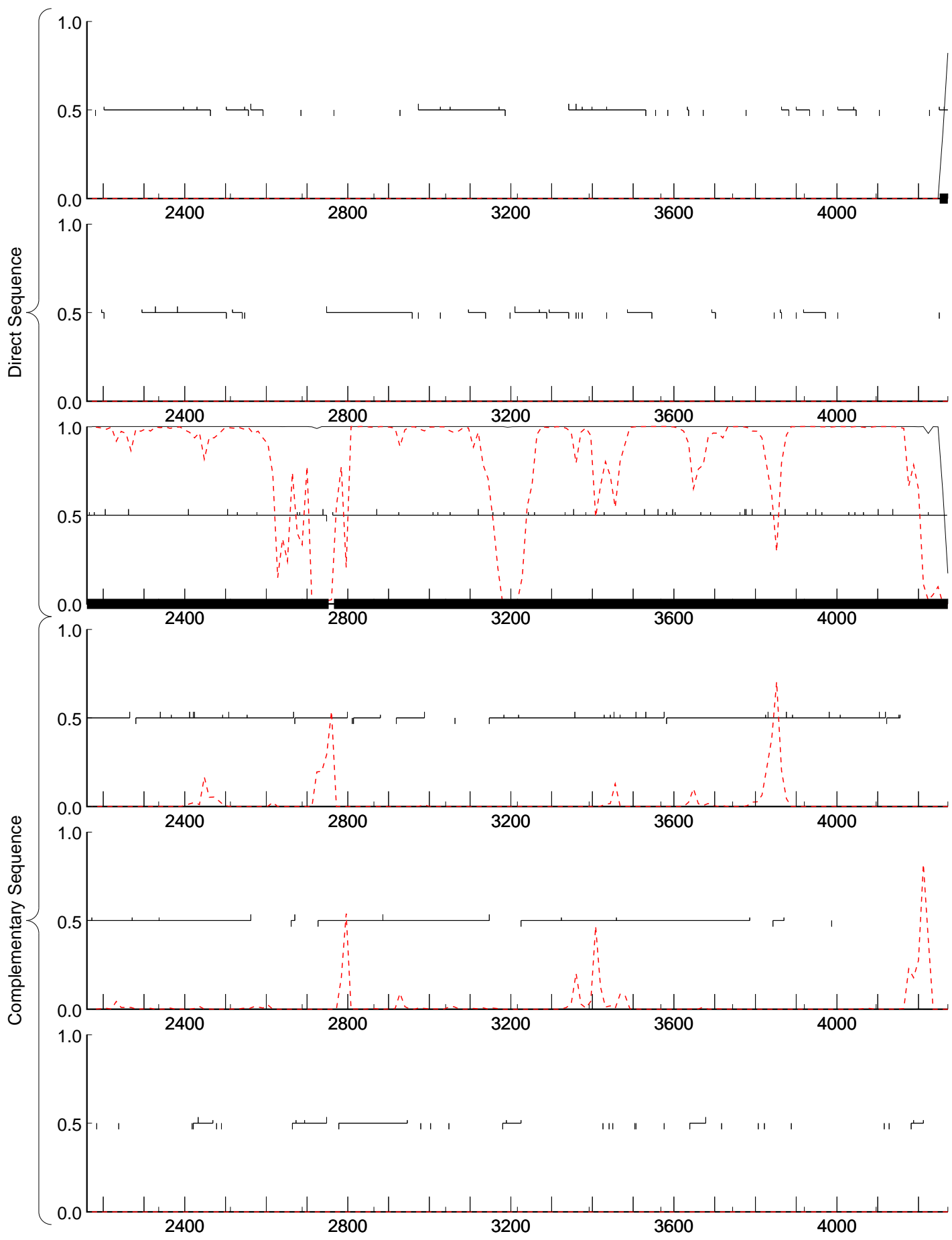
Matrix : -  
Author : -  
Order : 4

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

### Matrix notes & comments

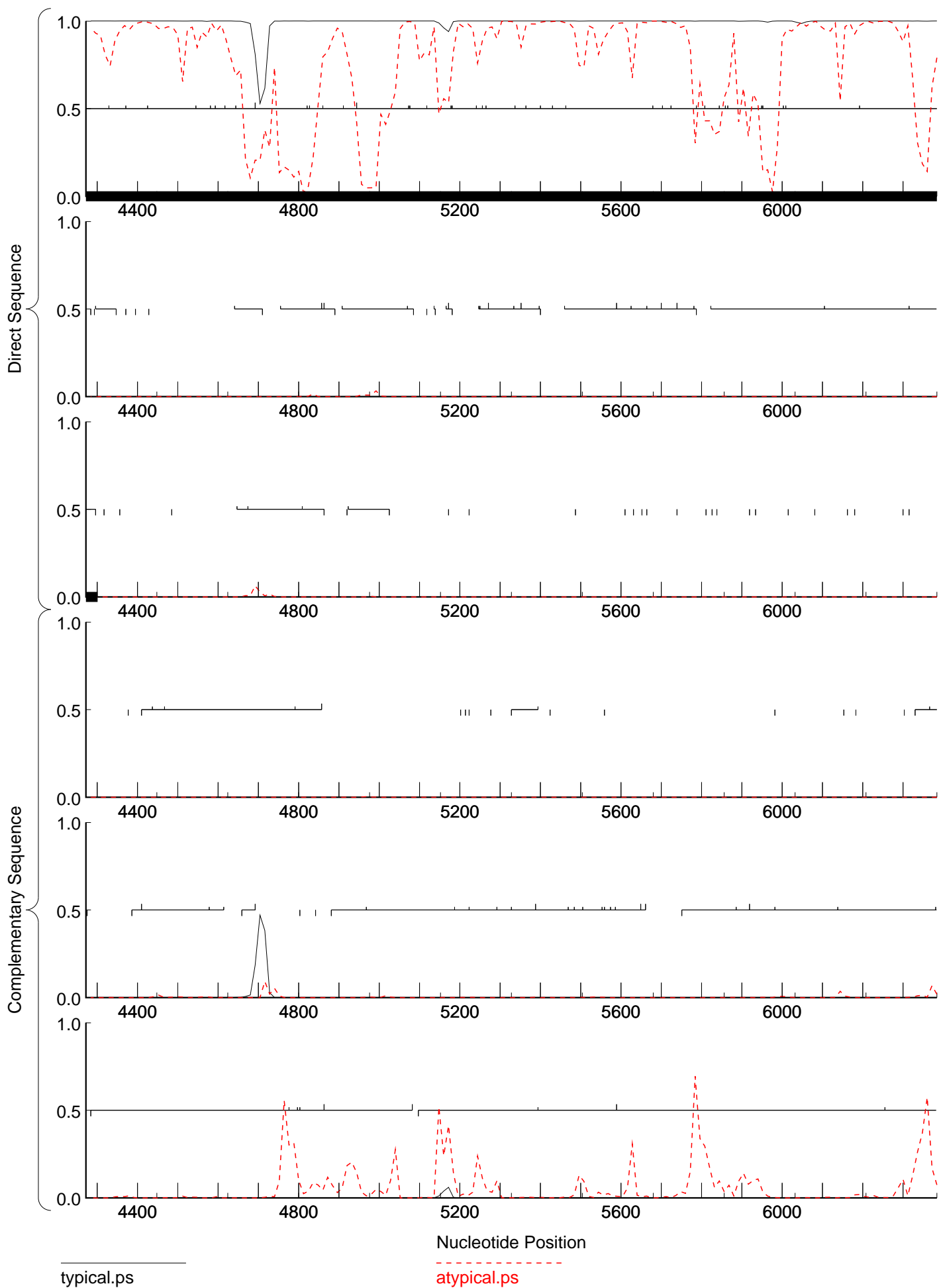
Training set derived by GeneMarkS, 4.30 October 2014  
Sun Nov 19 19:05:40 2017

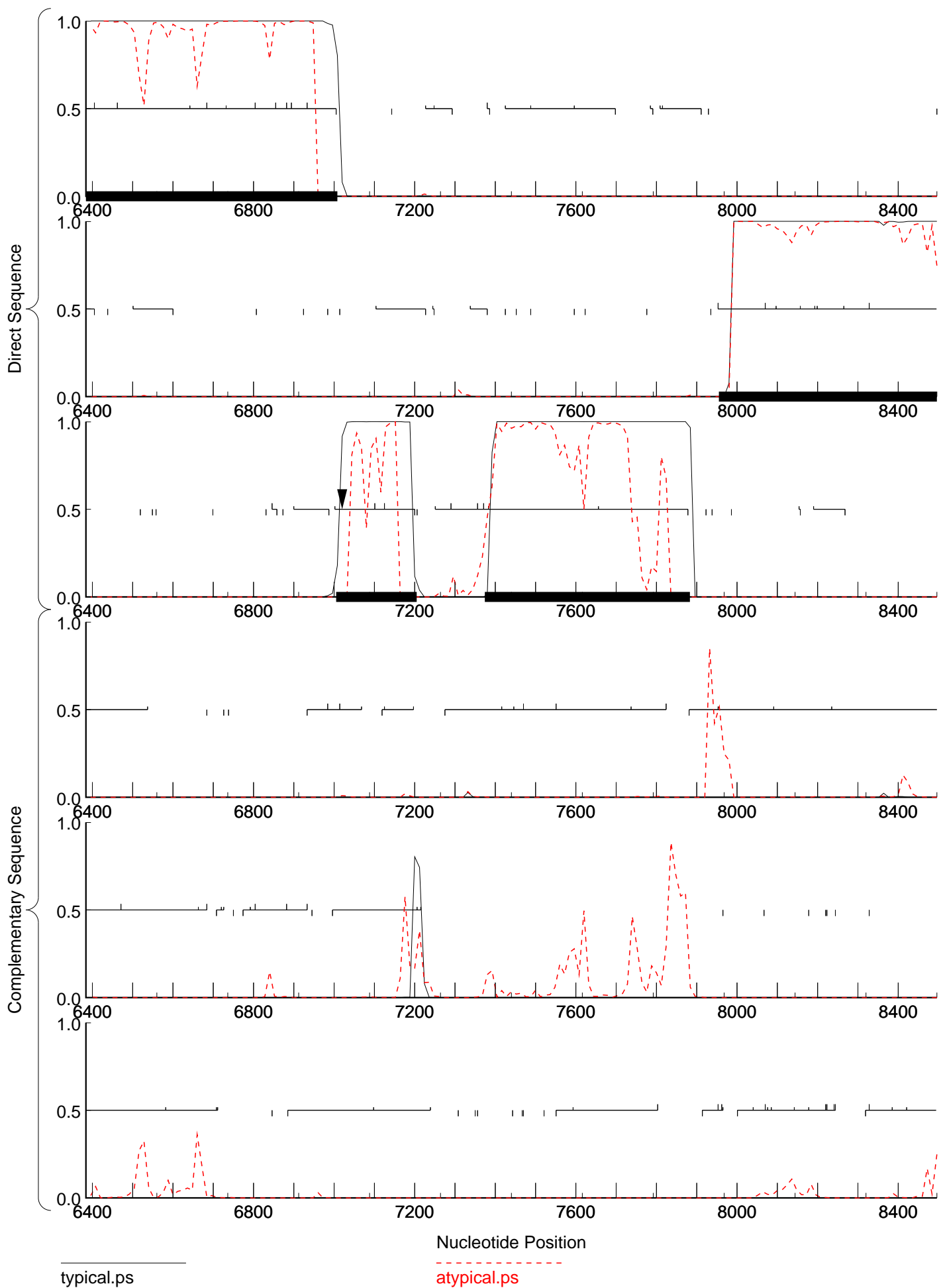




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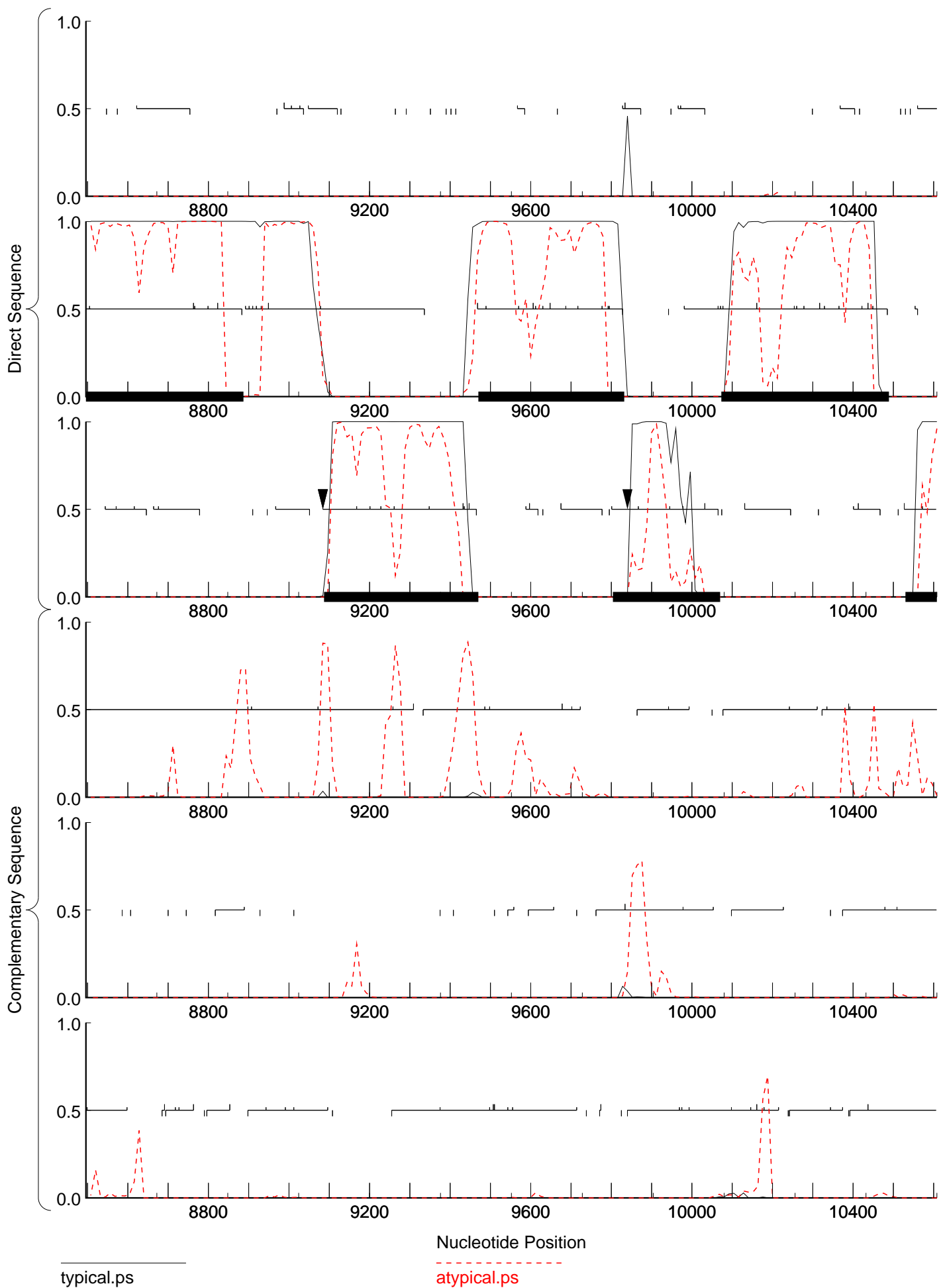
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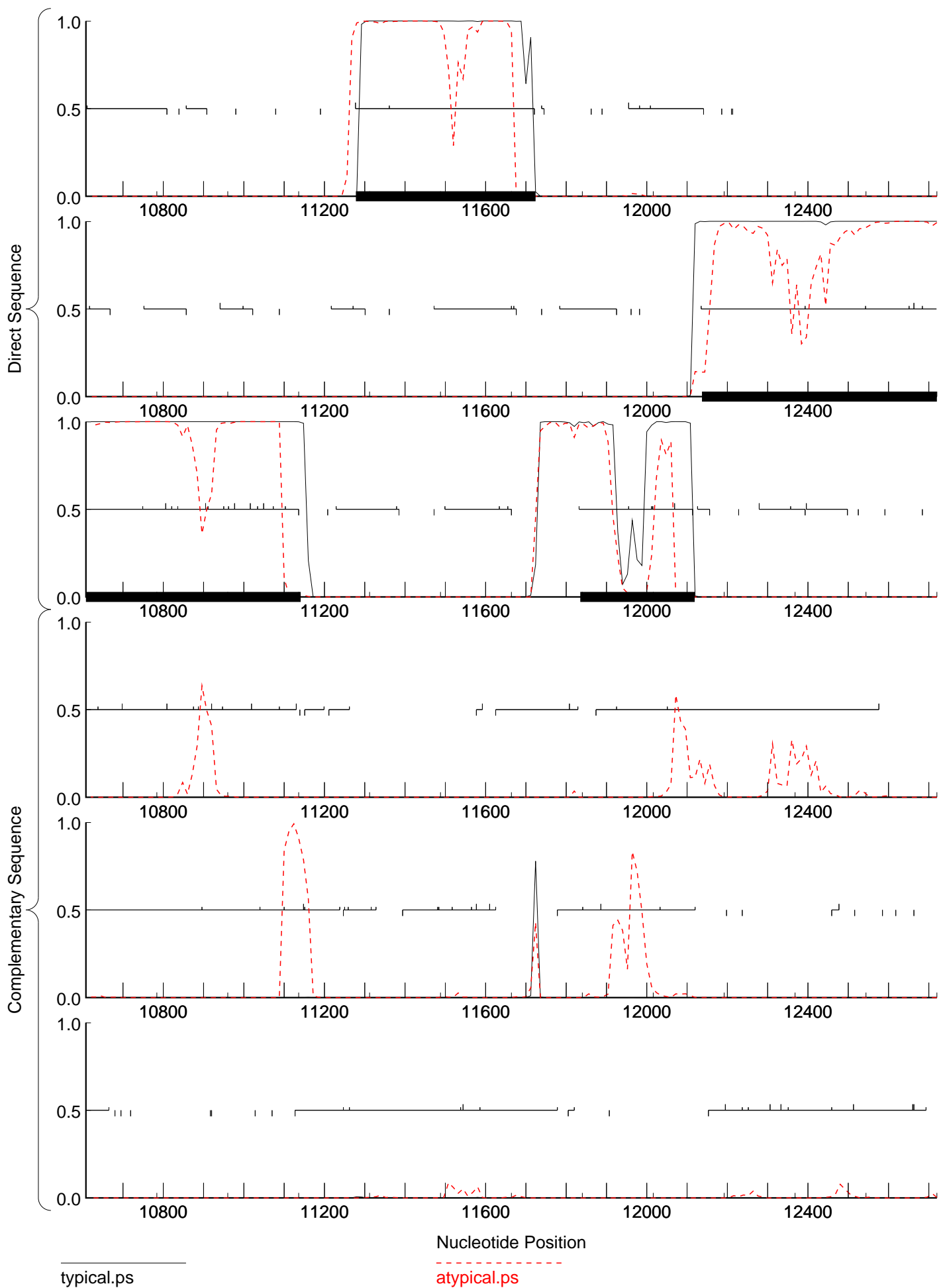


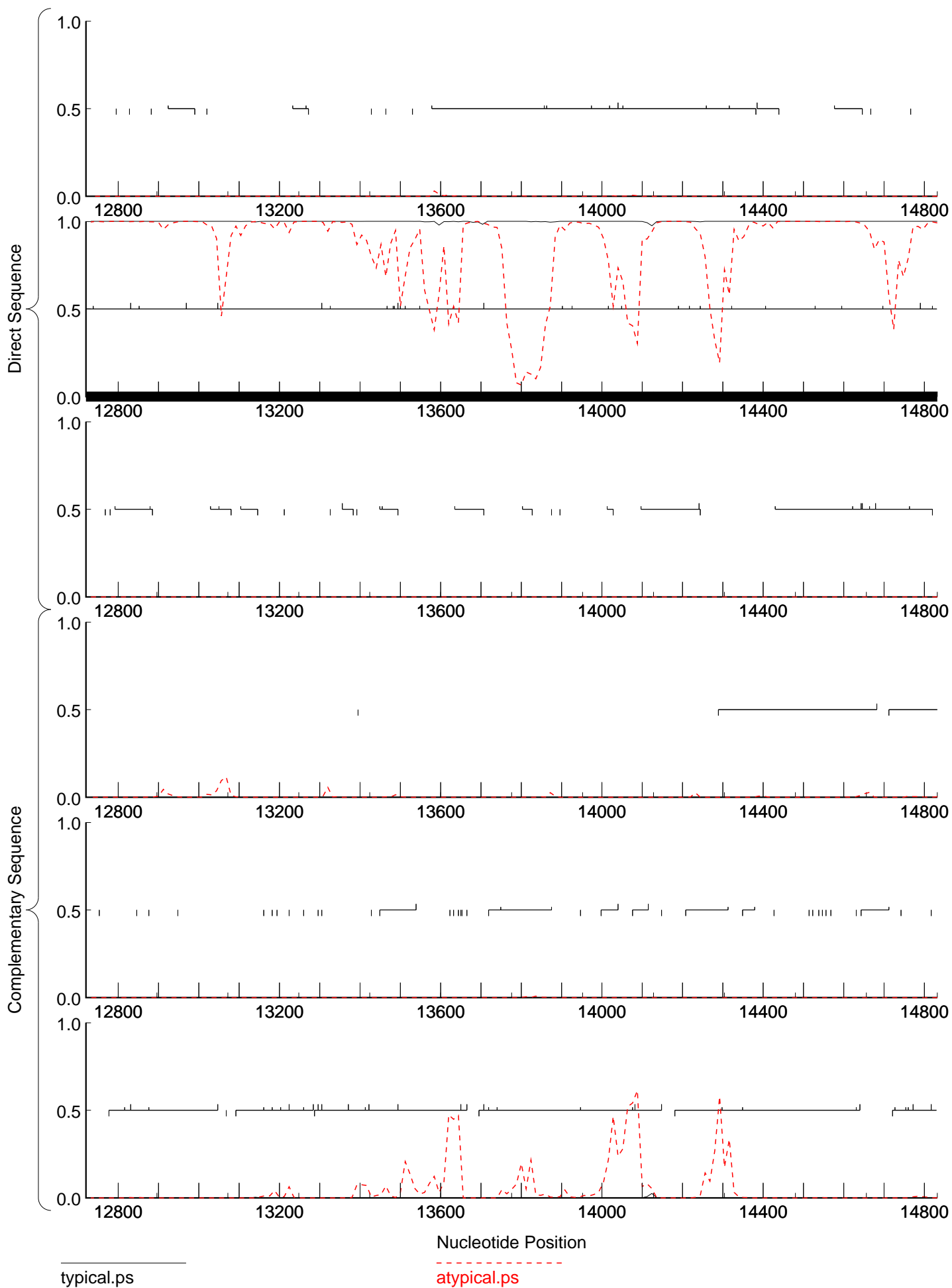


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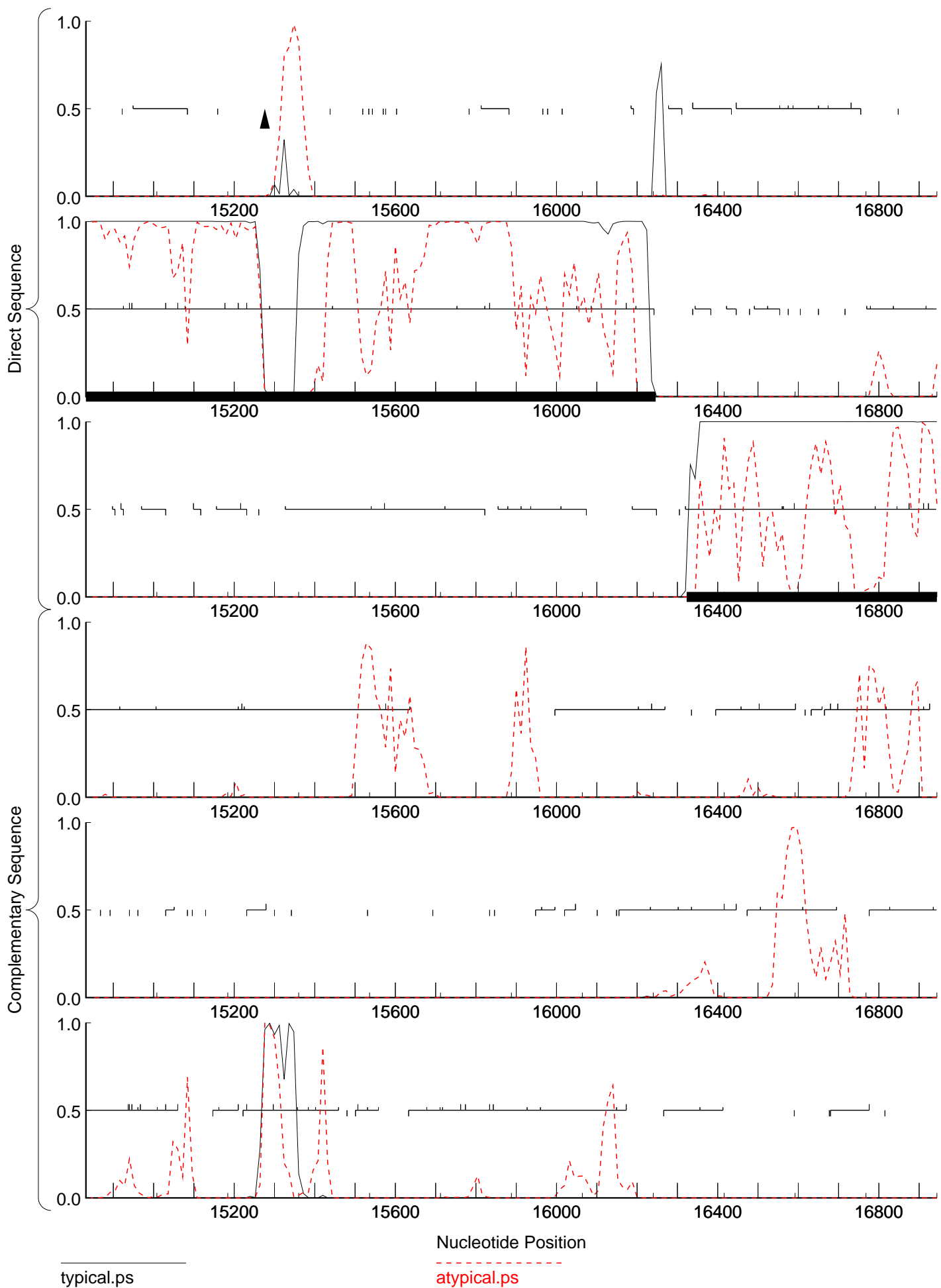
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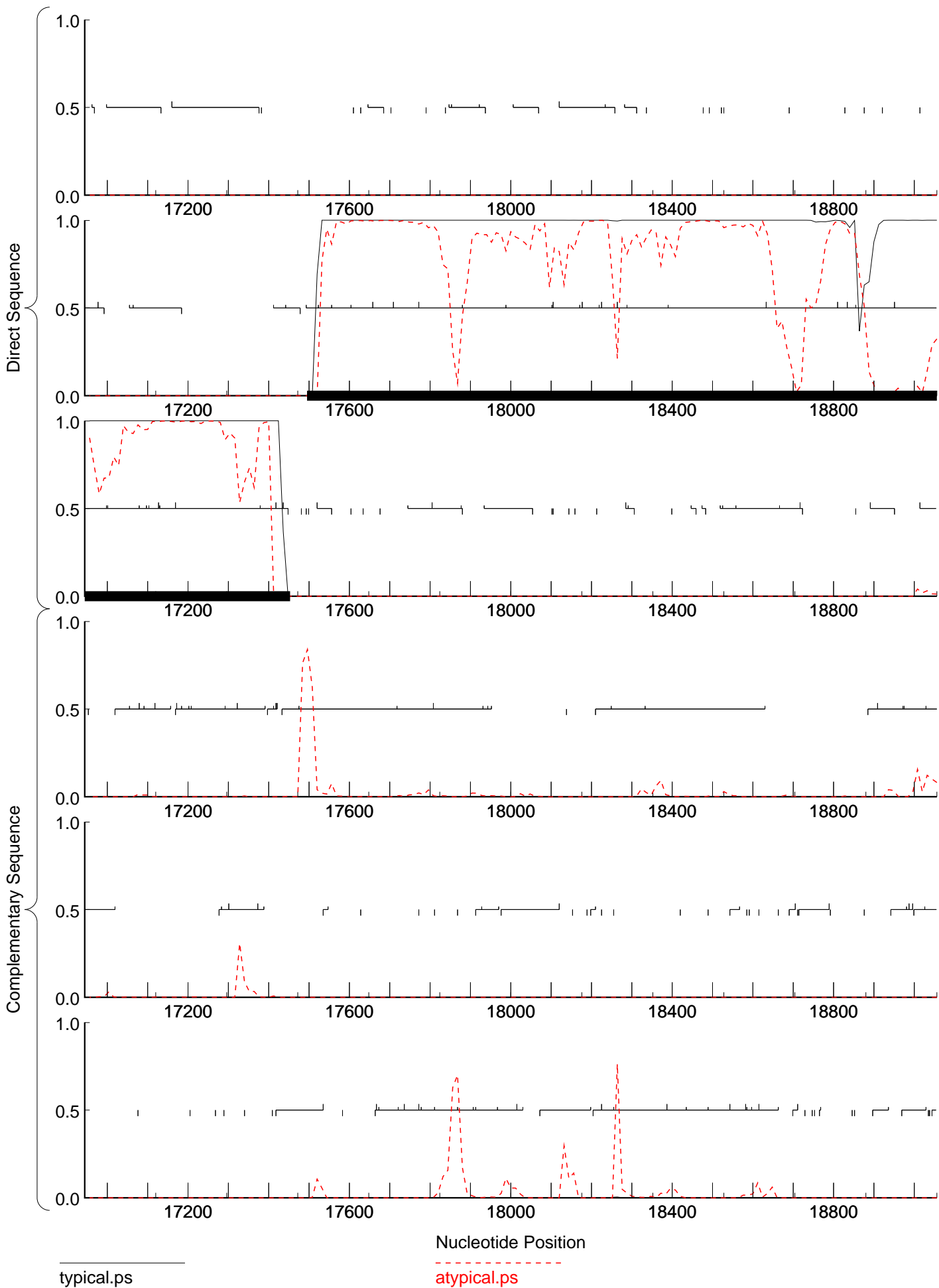






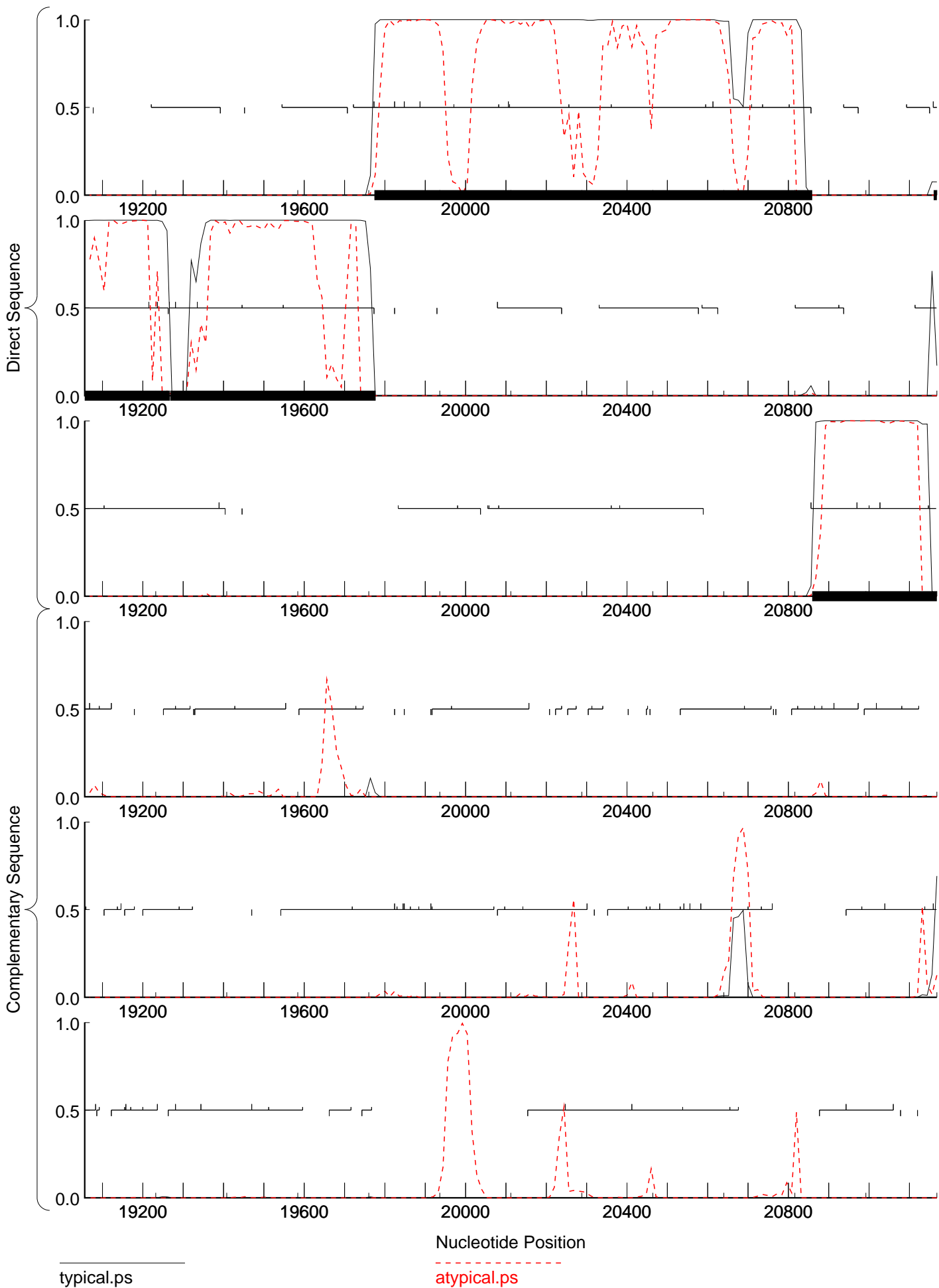


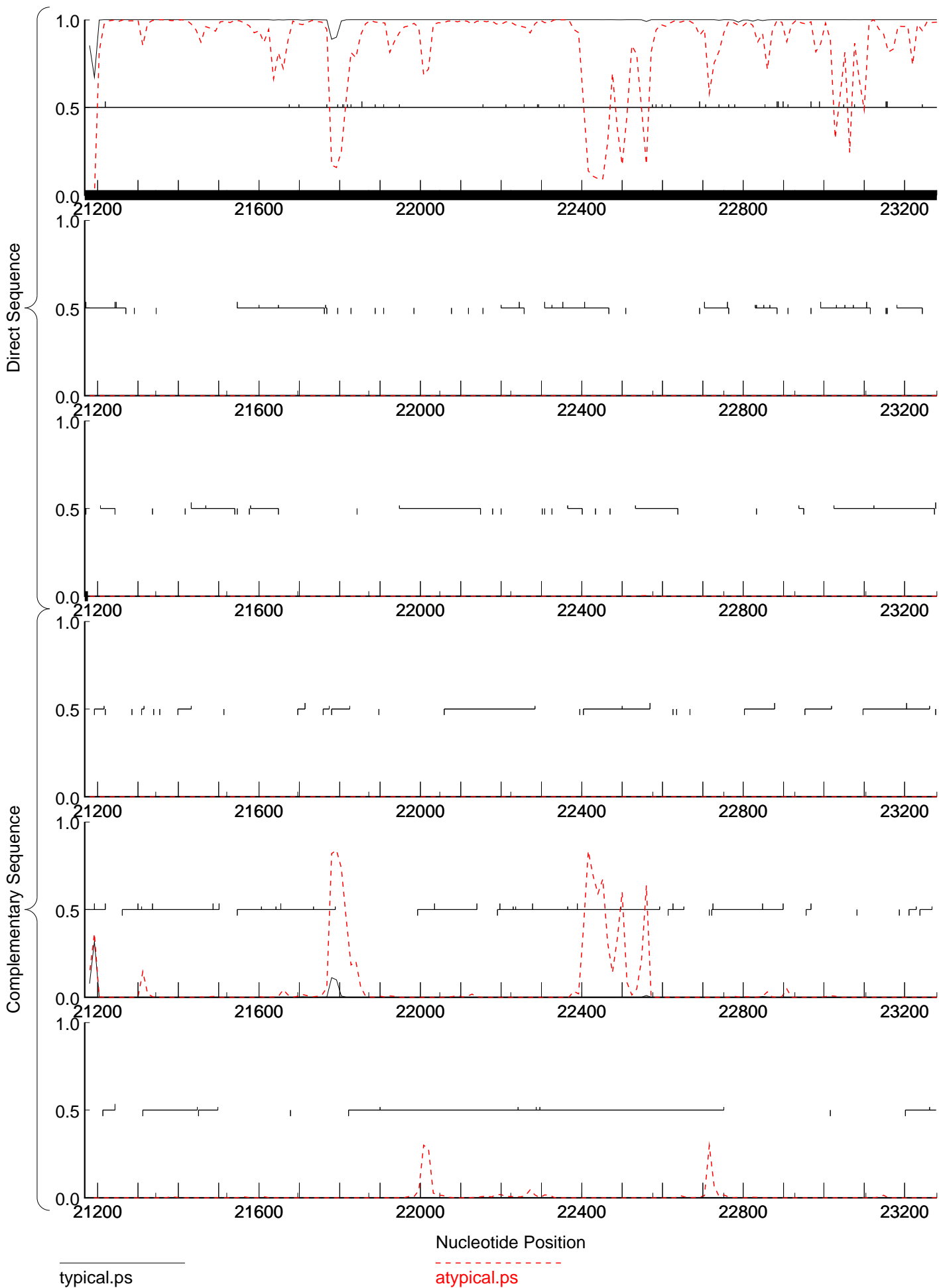


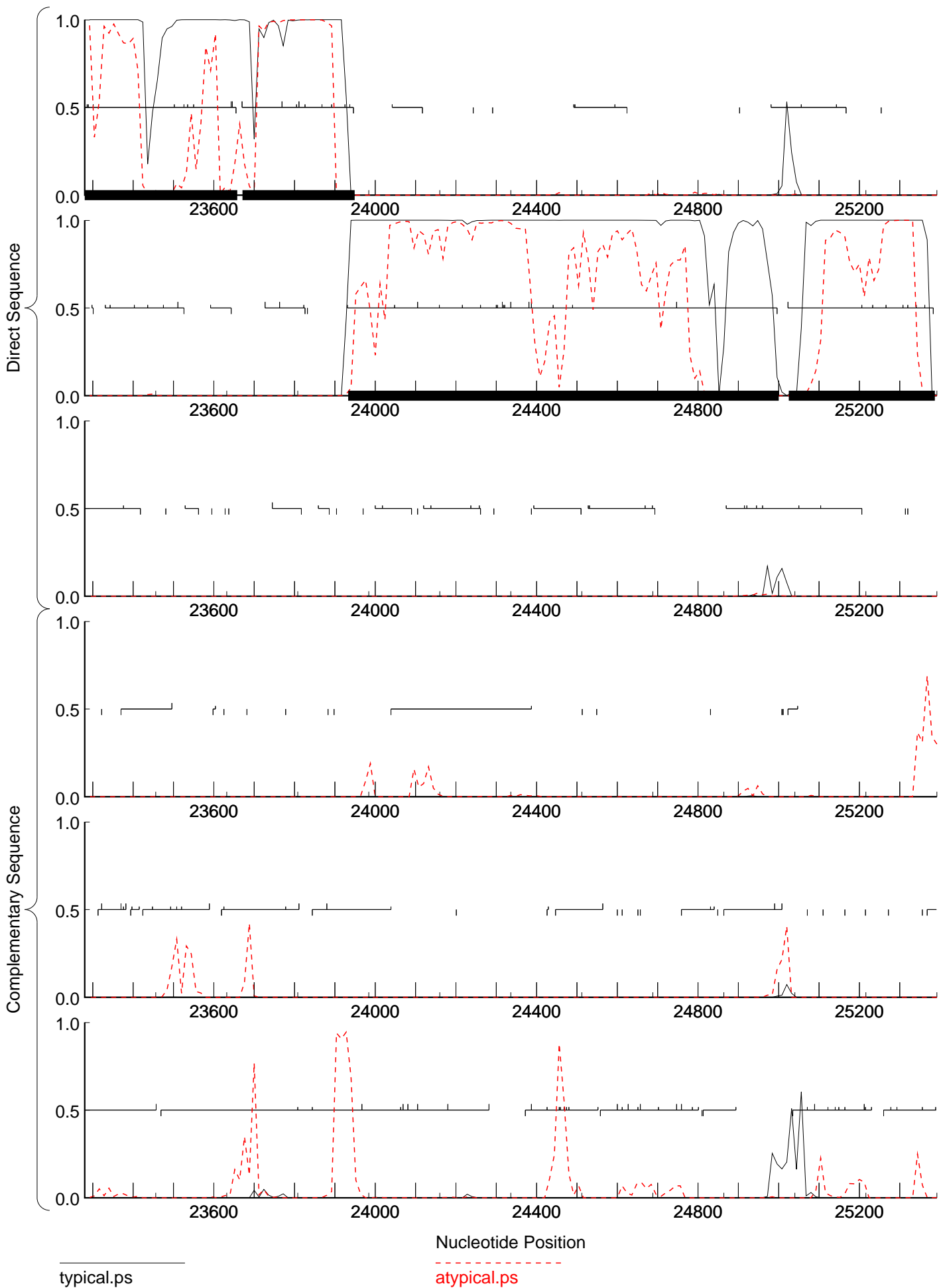


typical.ps

atypical.ps

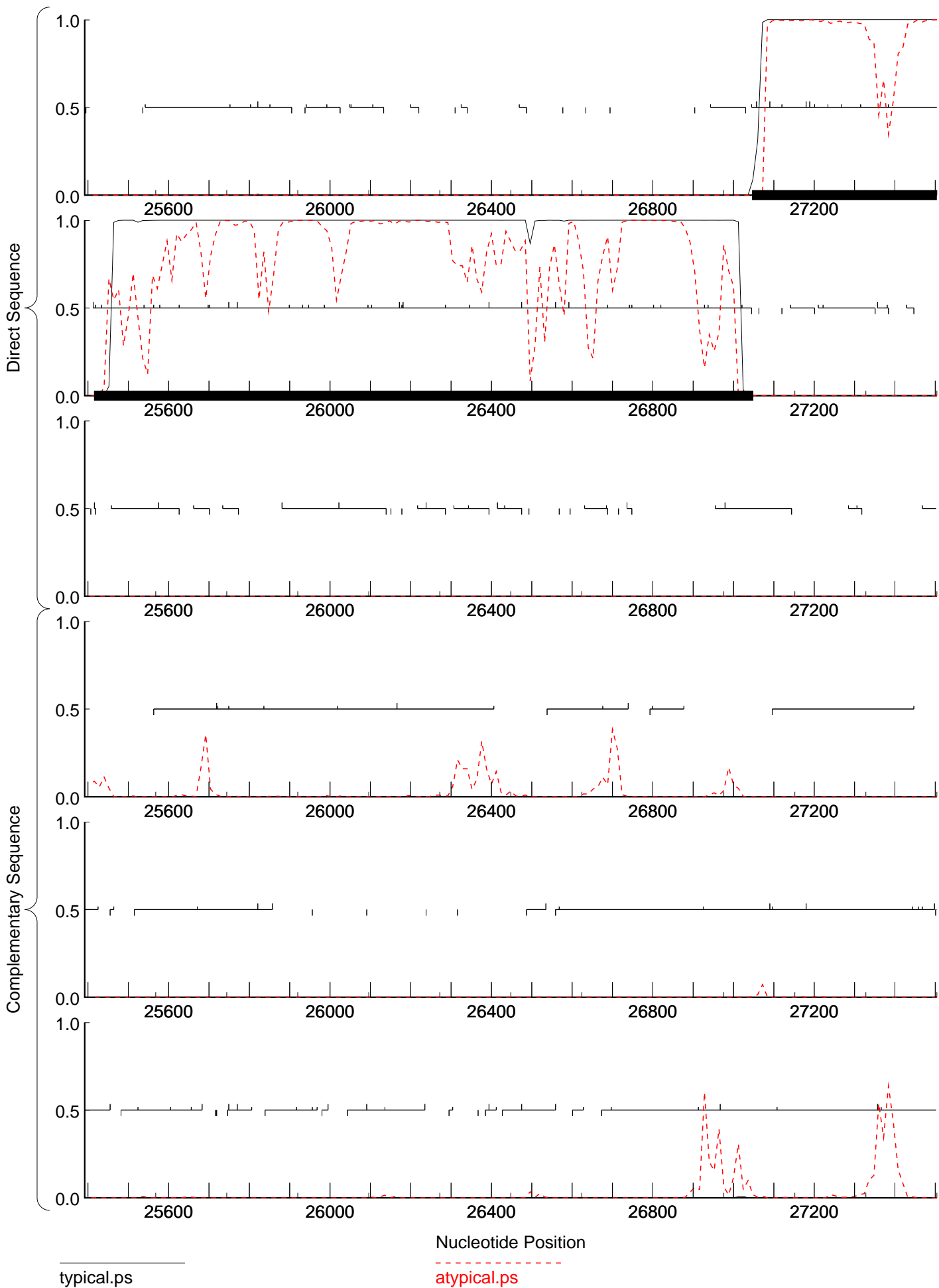


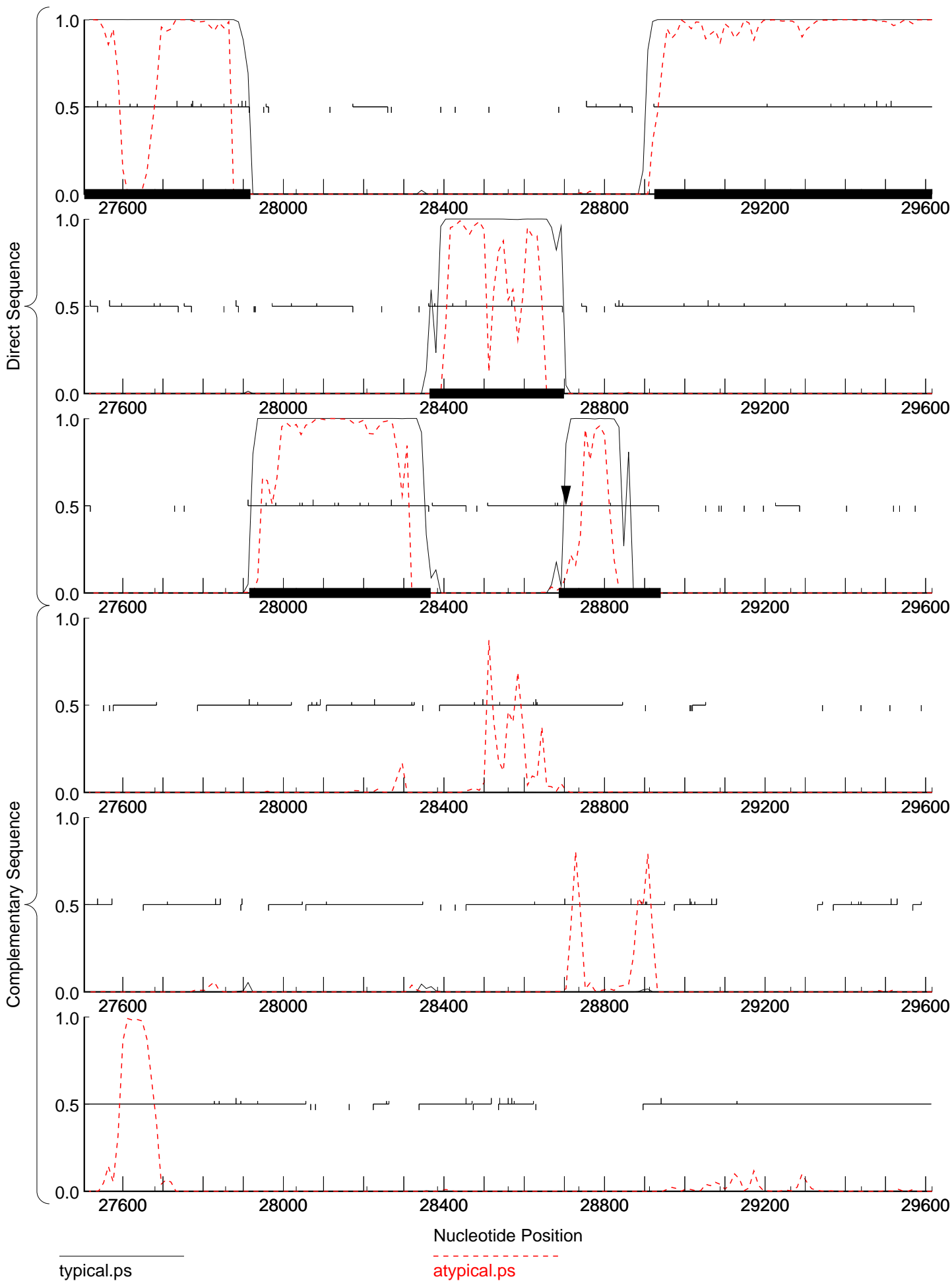




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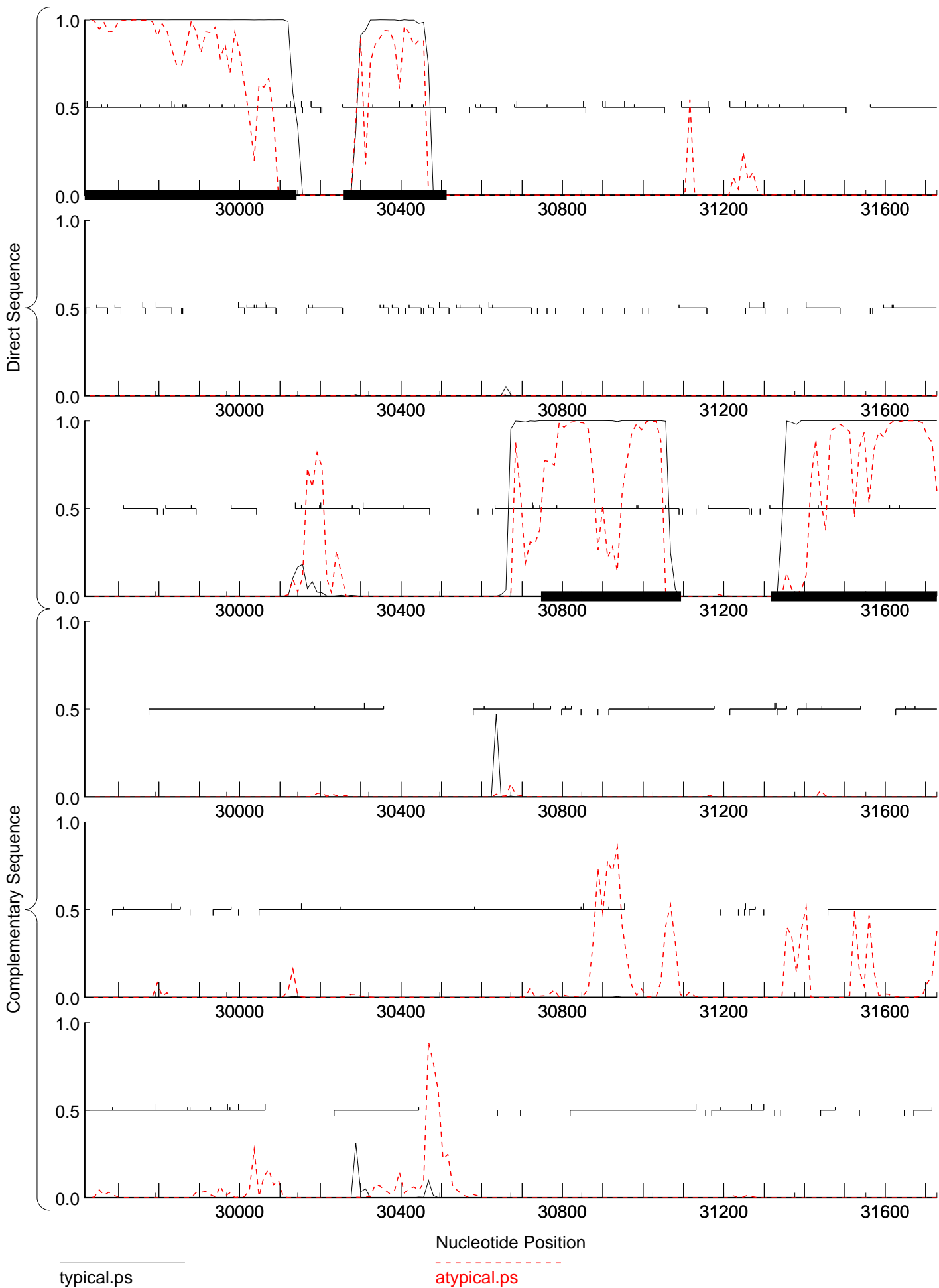
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typical.ps

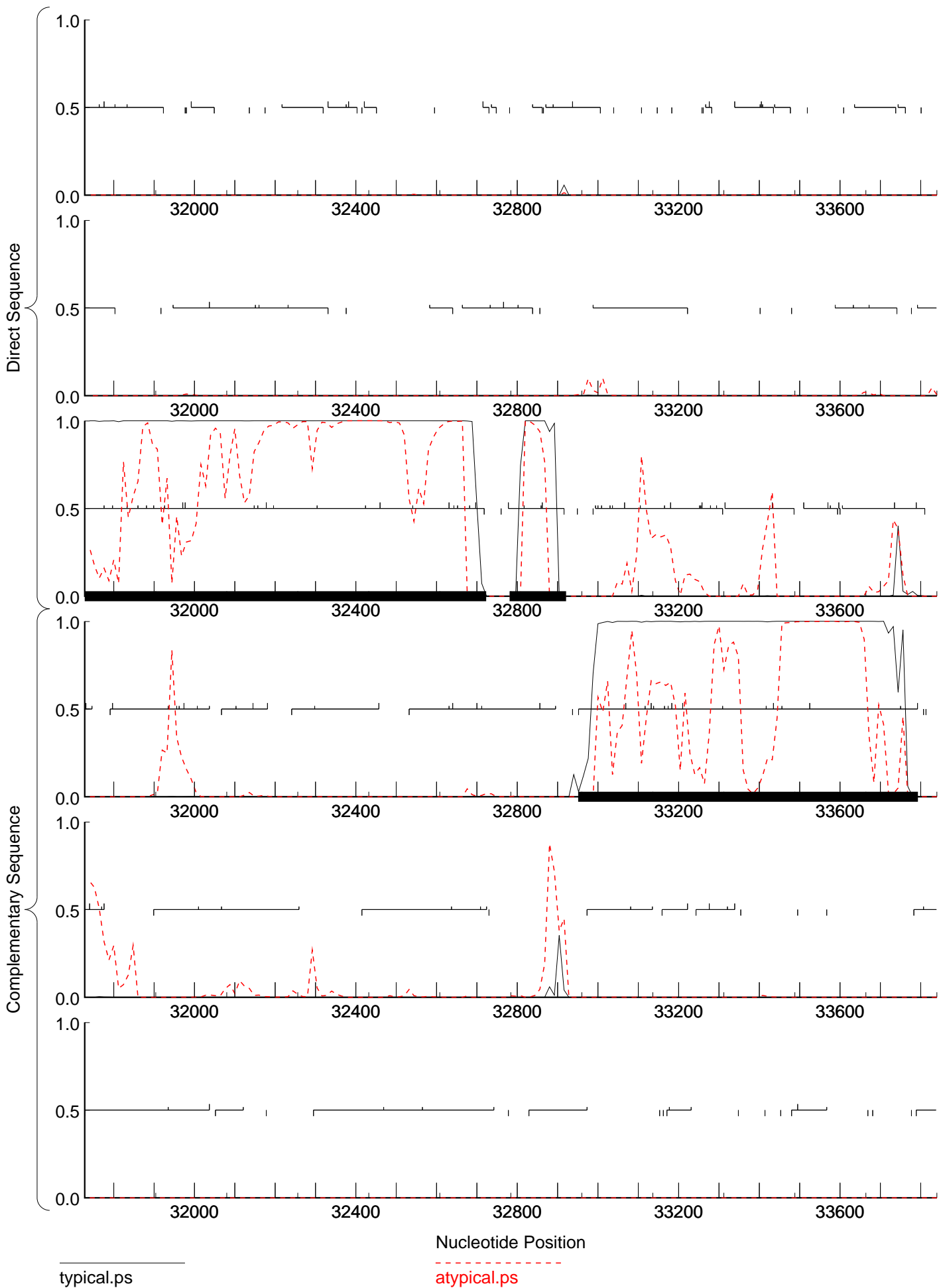
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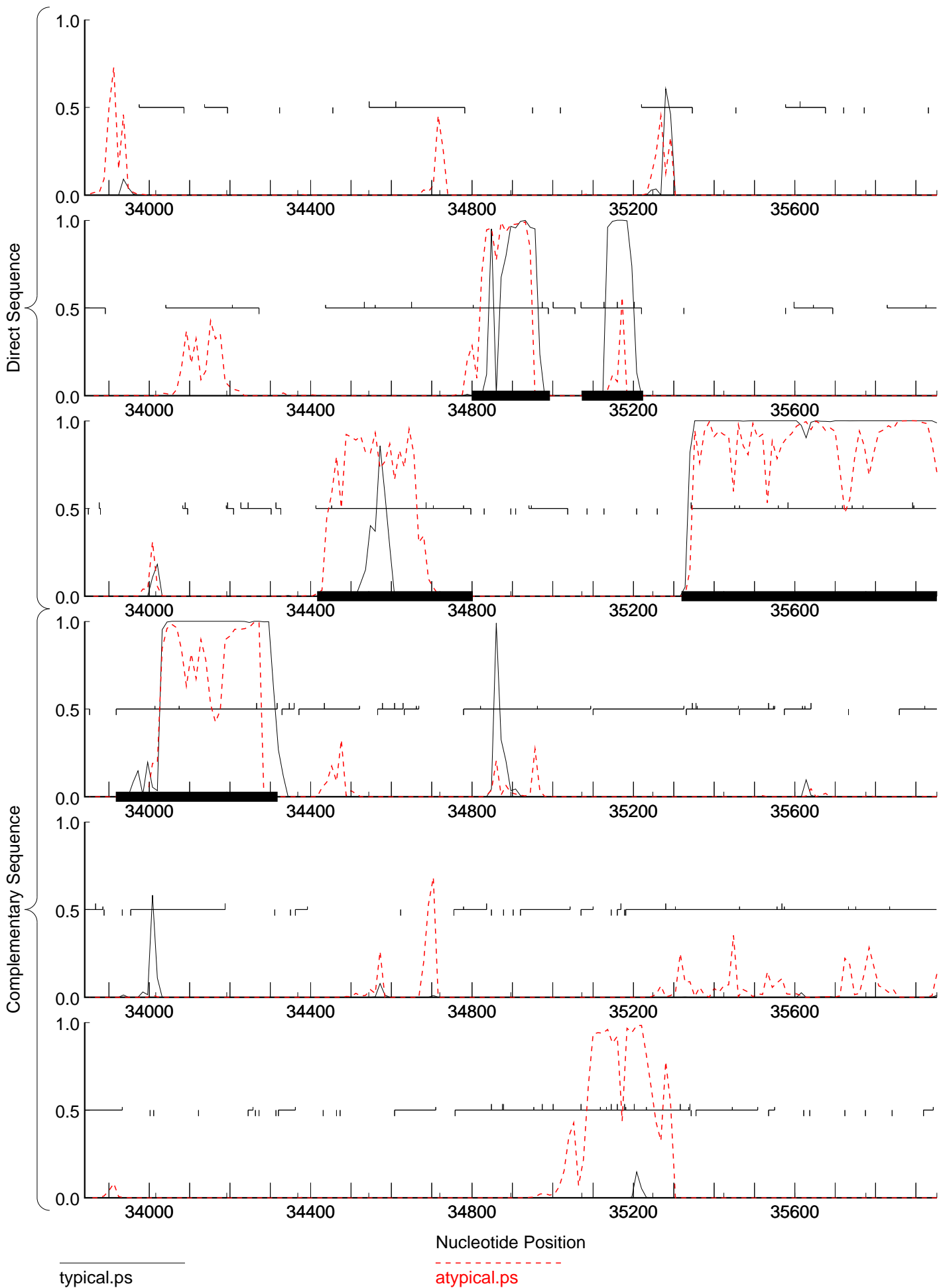


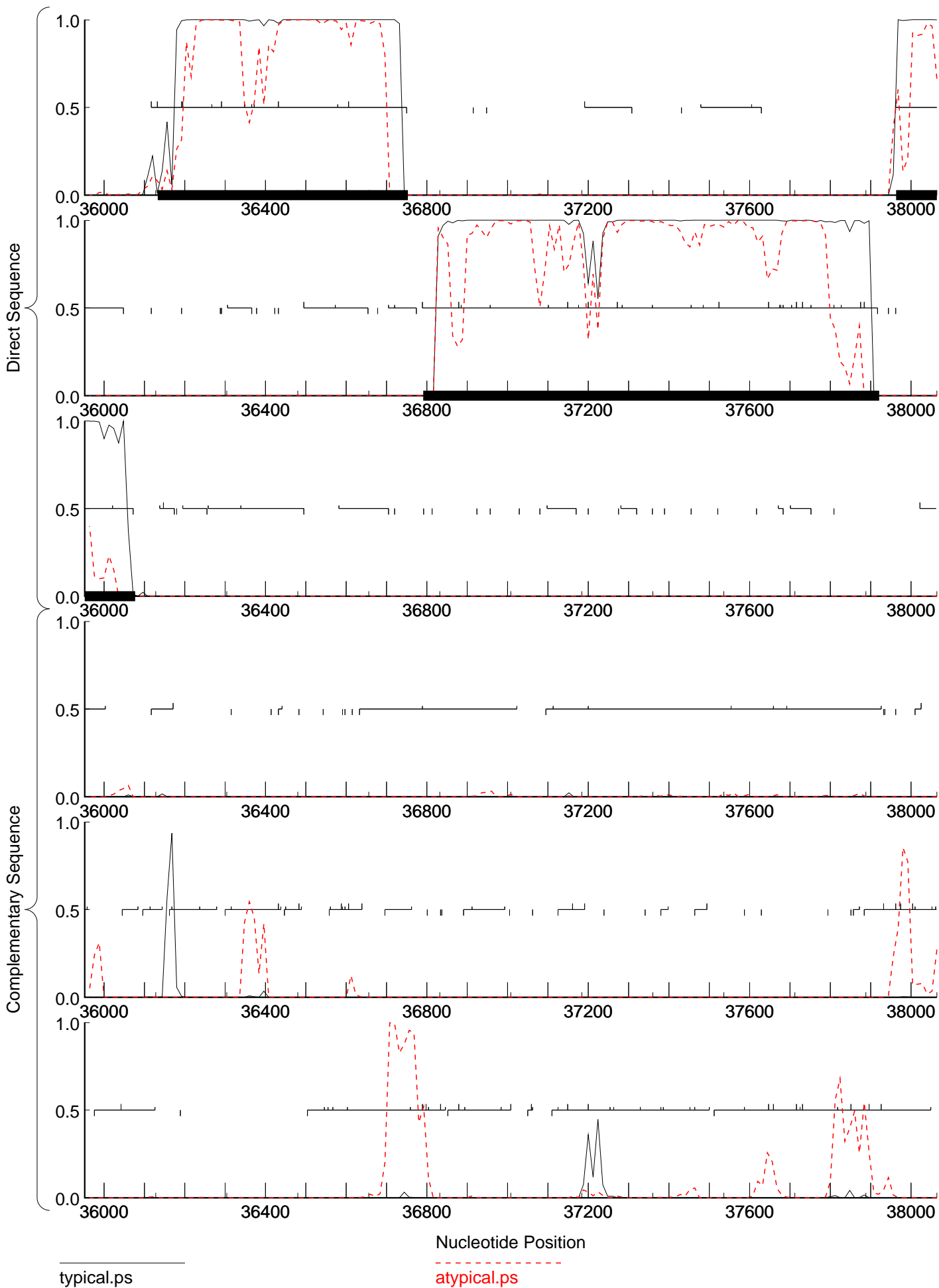
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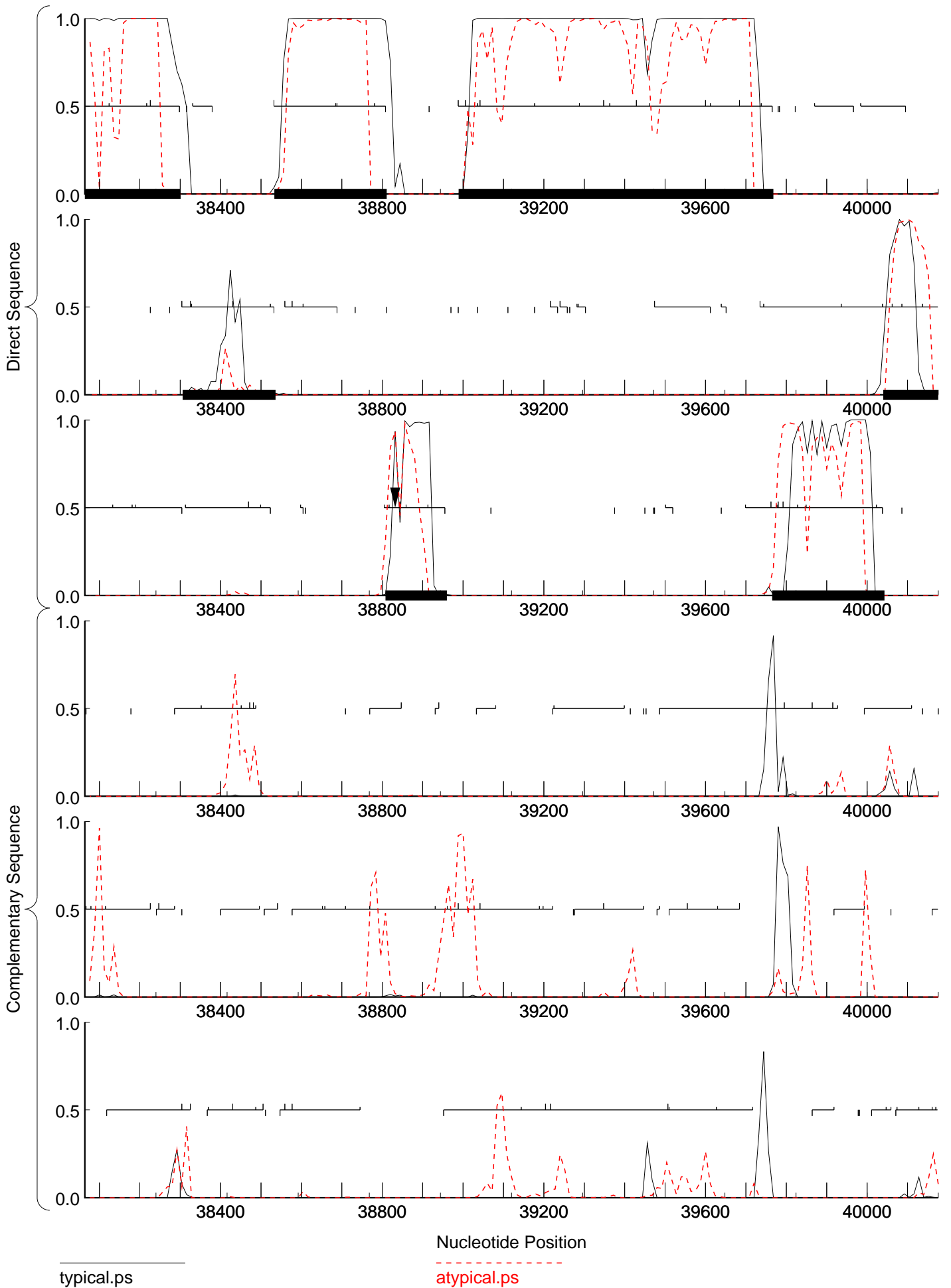
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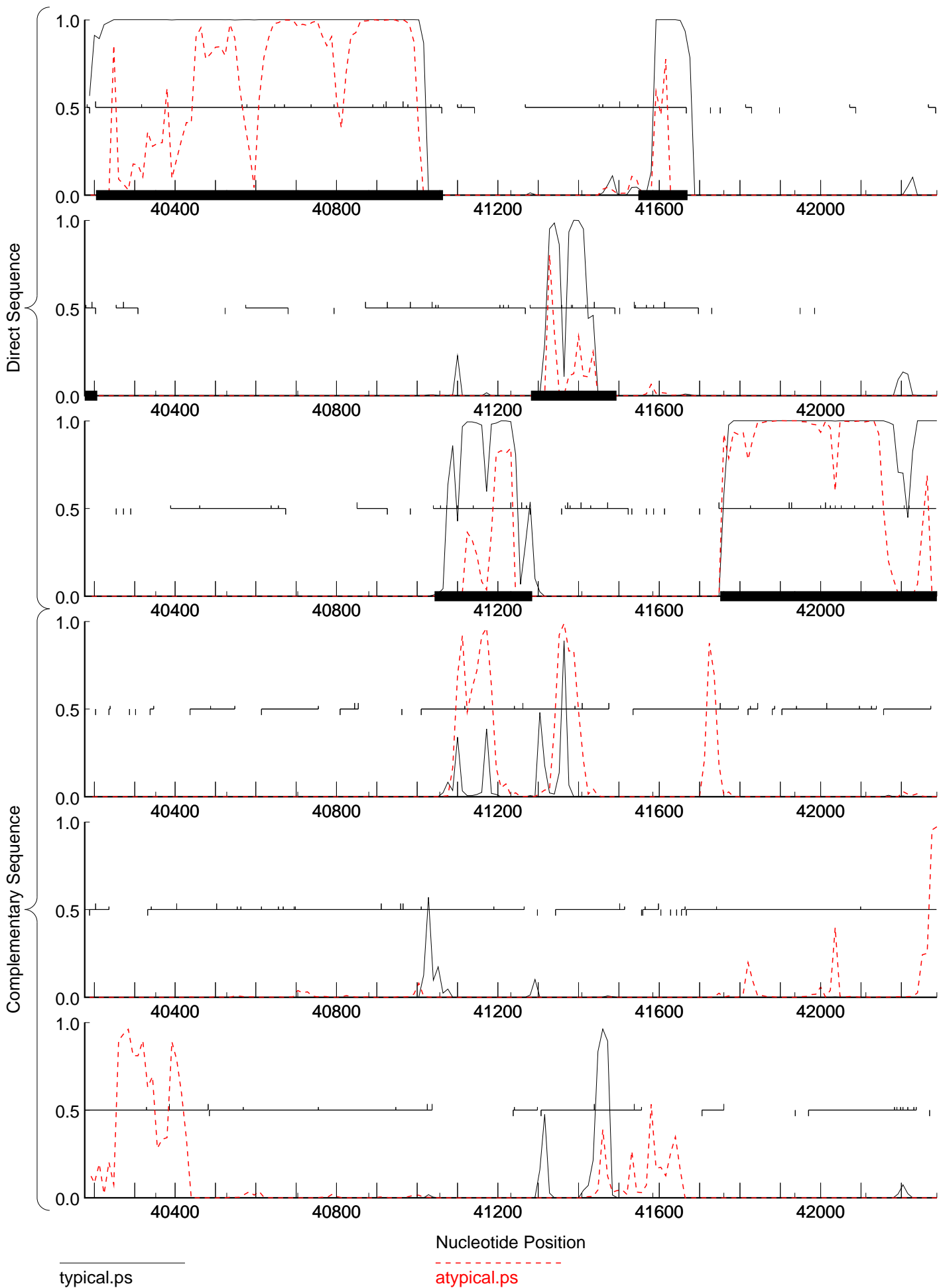


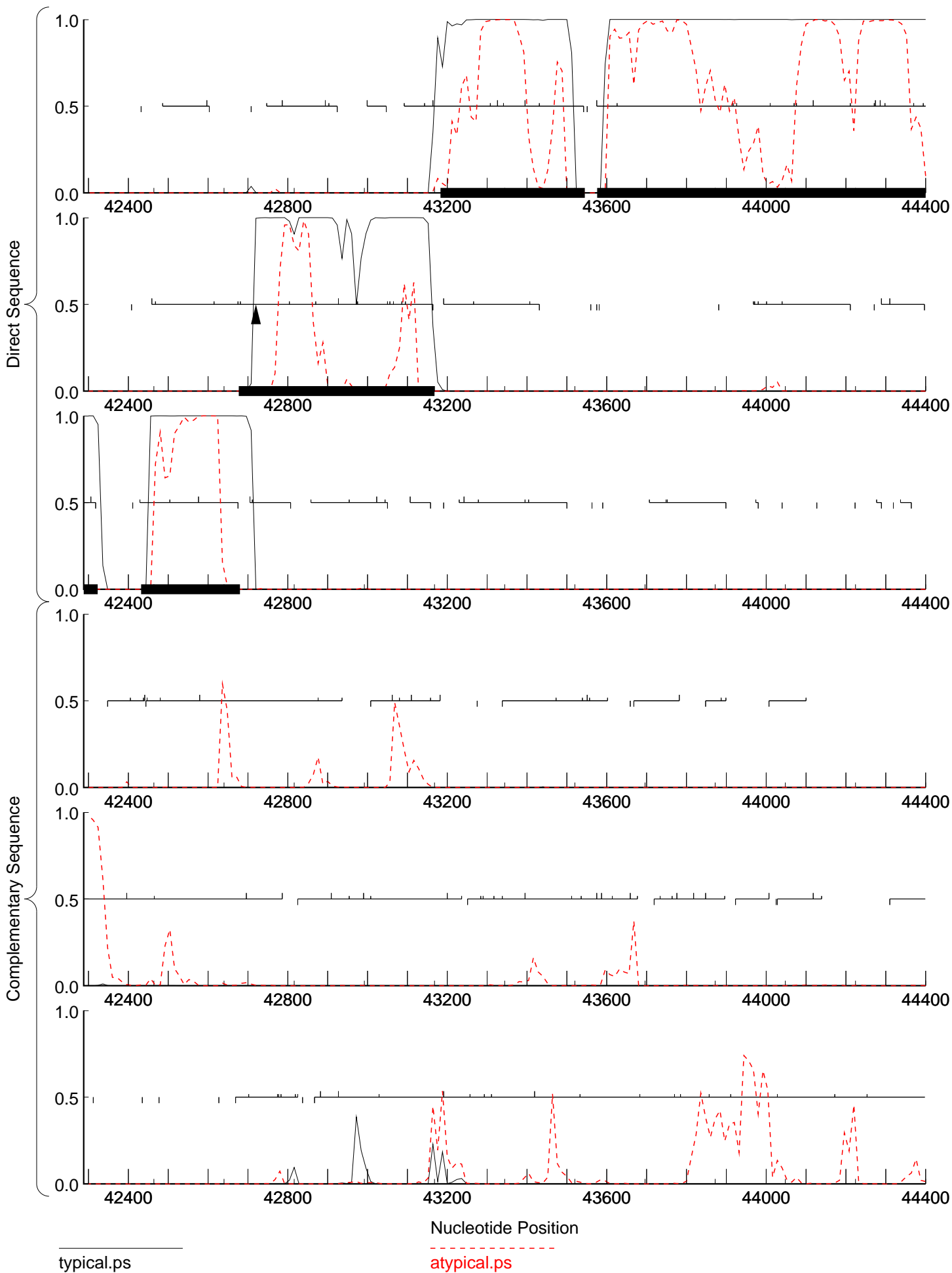


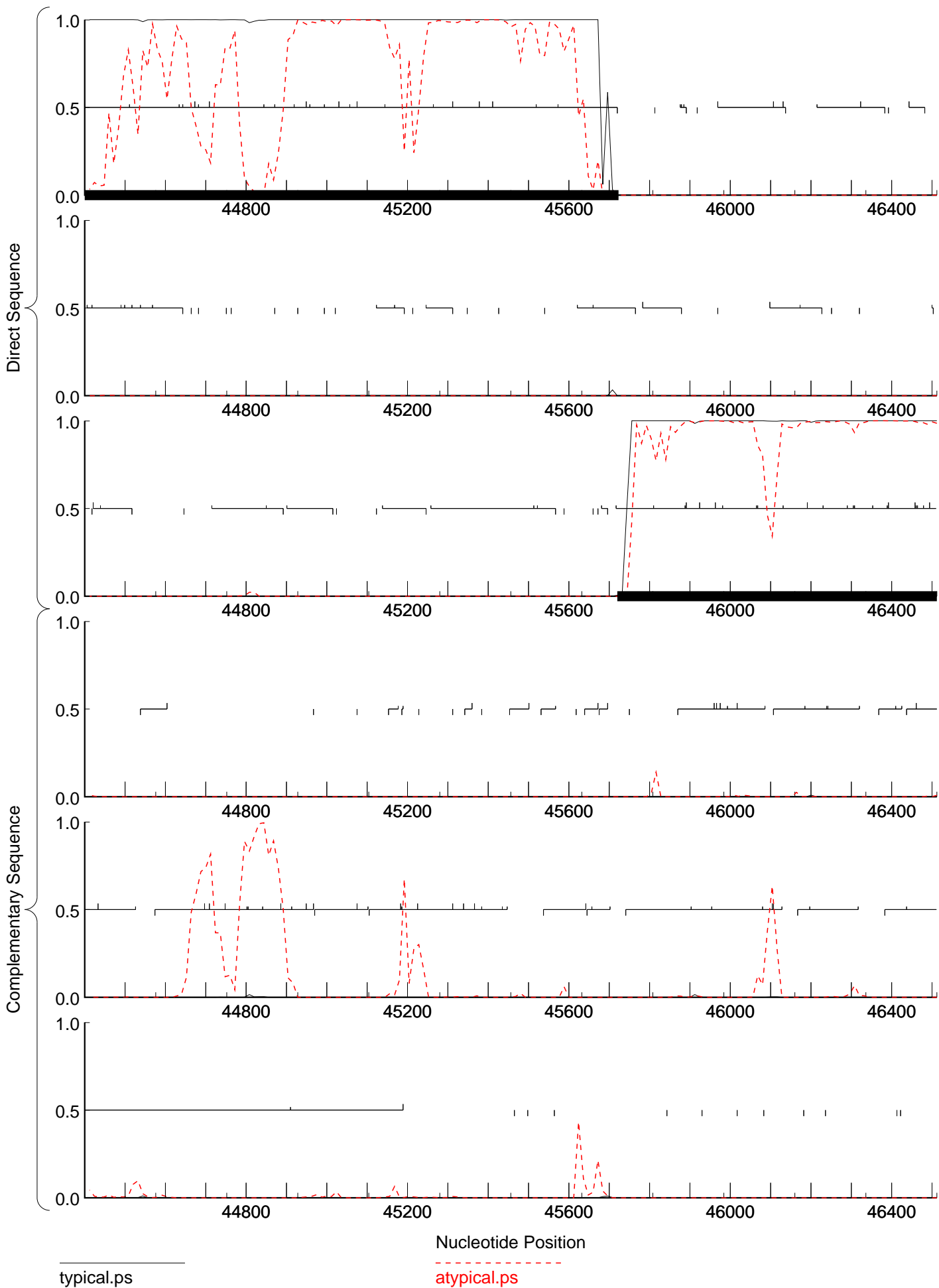


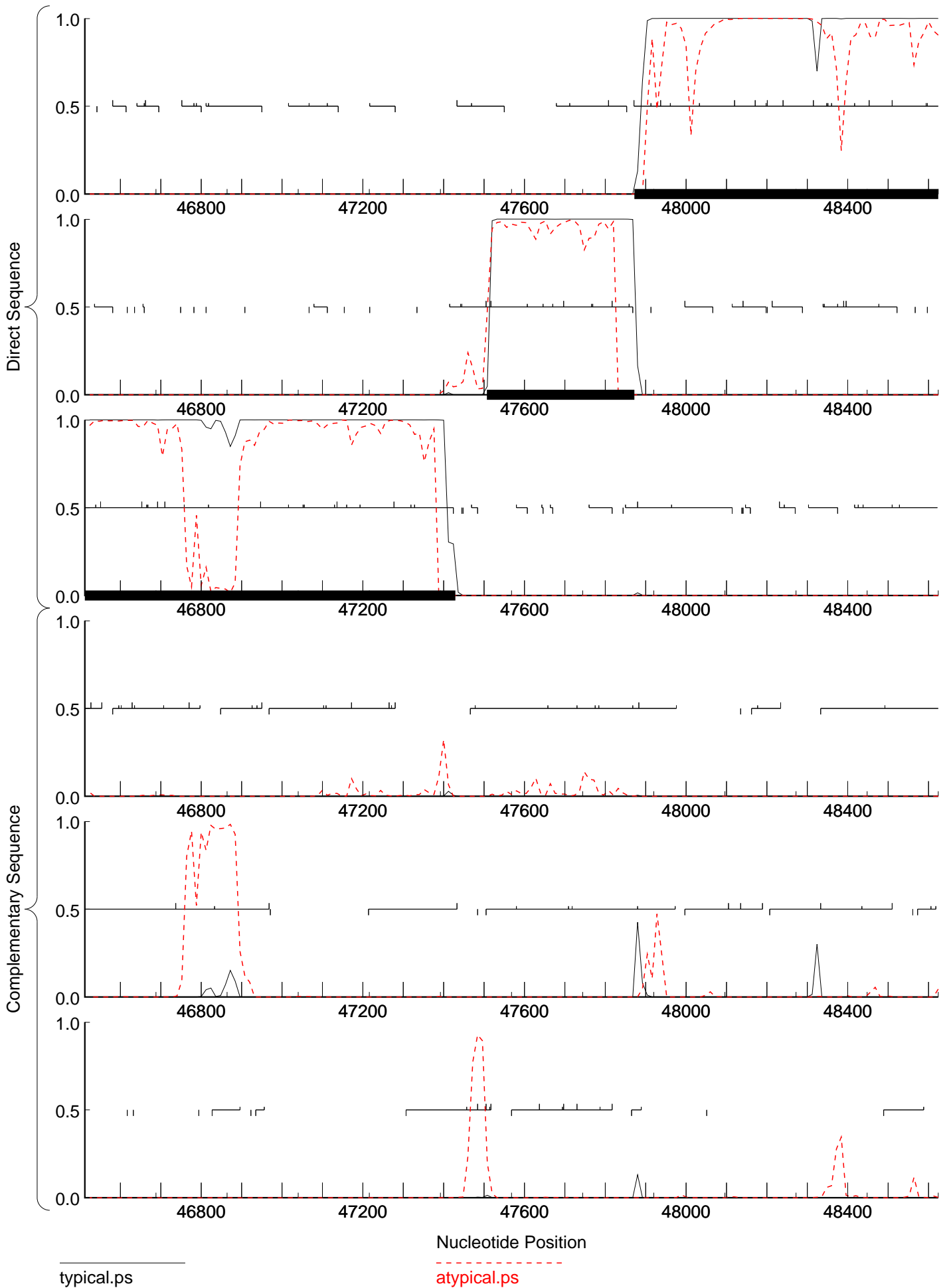
typical.ps

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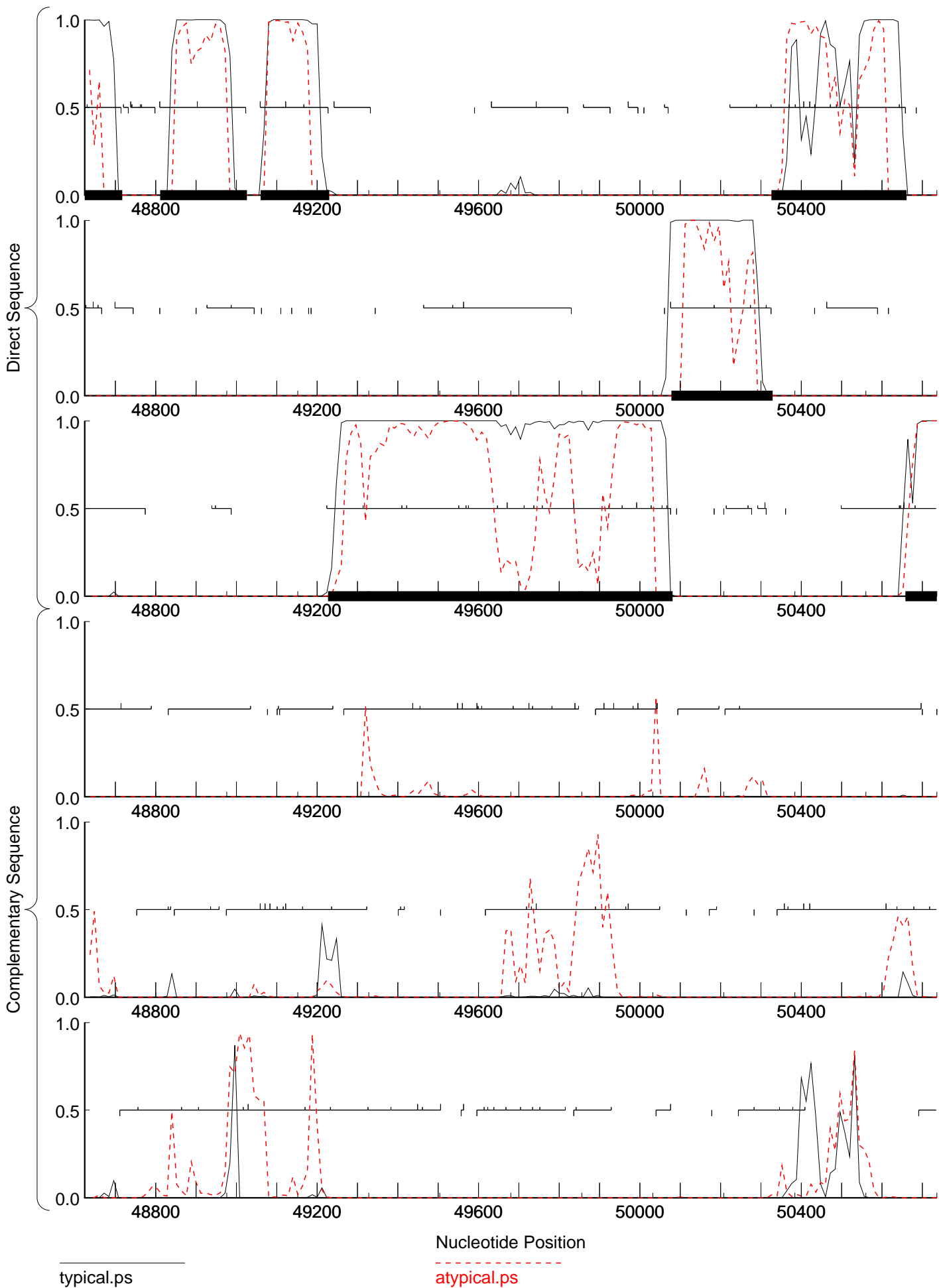






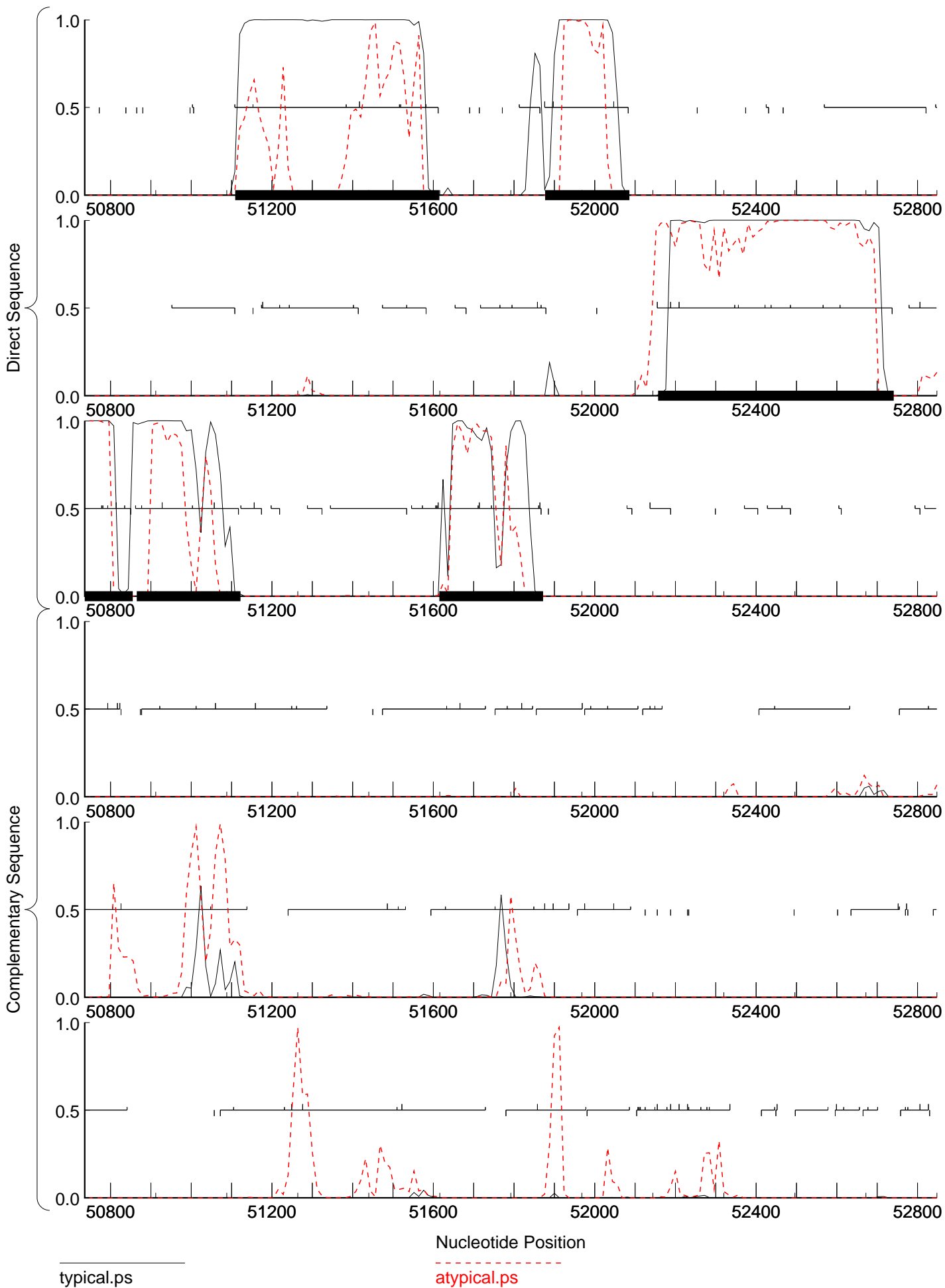


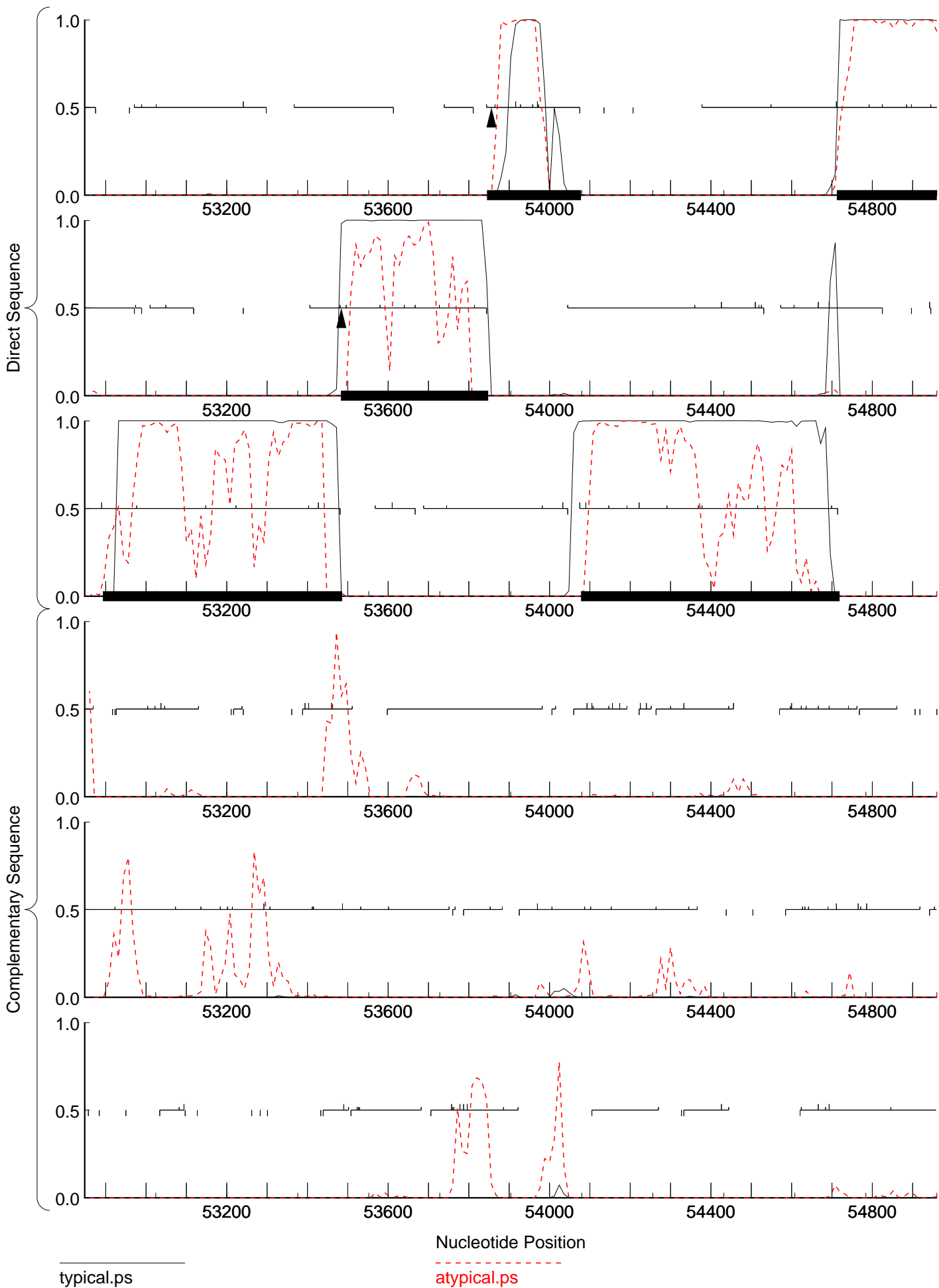




typical.ps

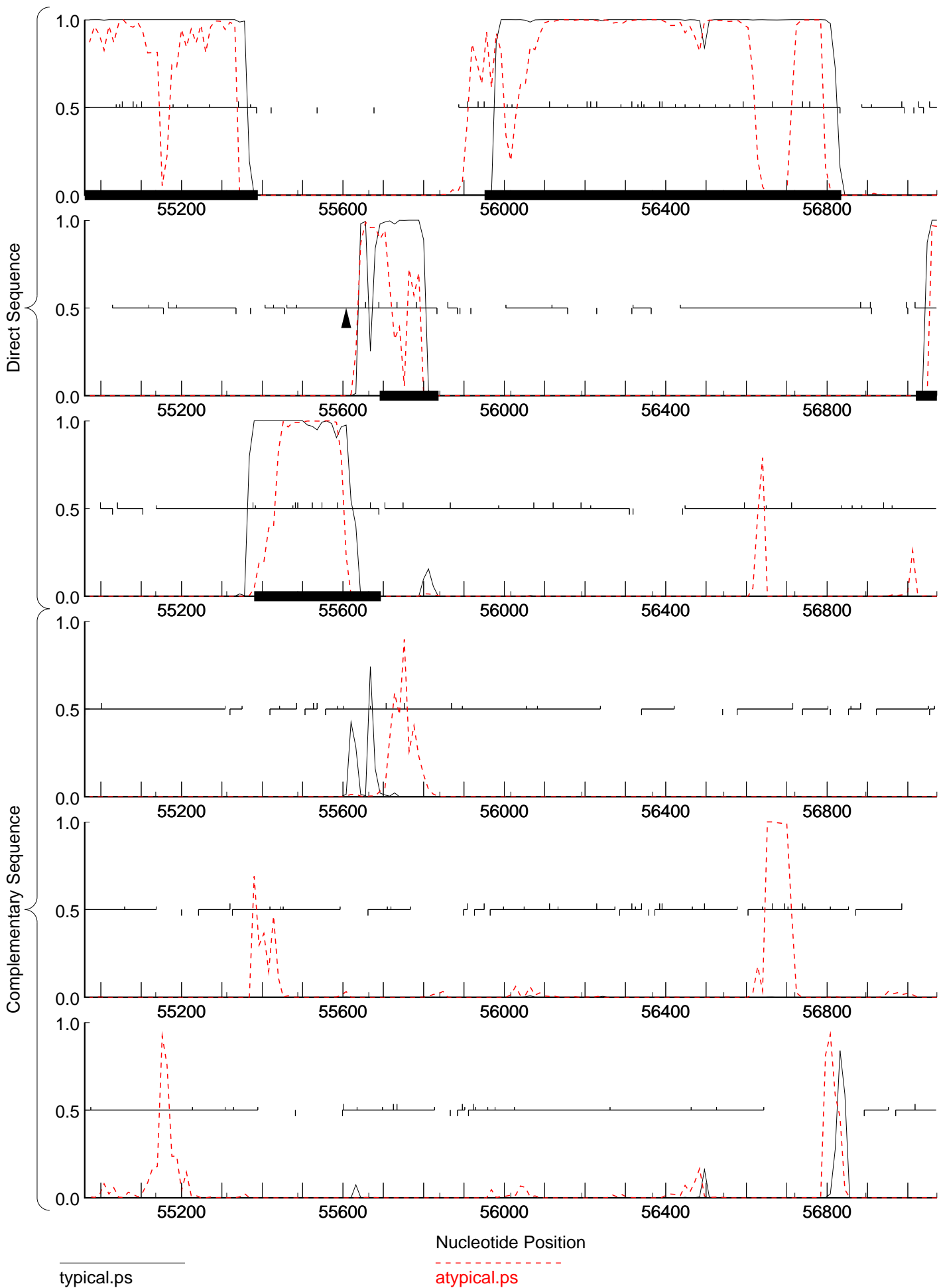
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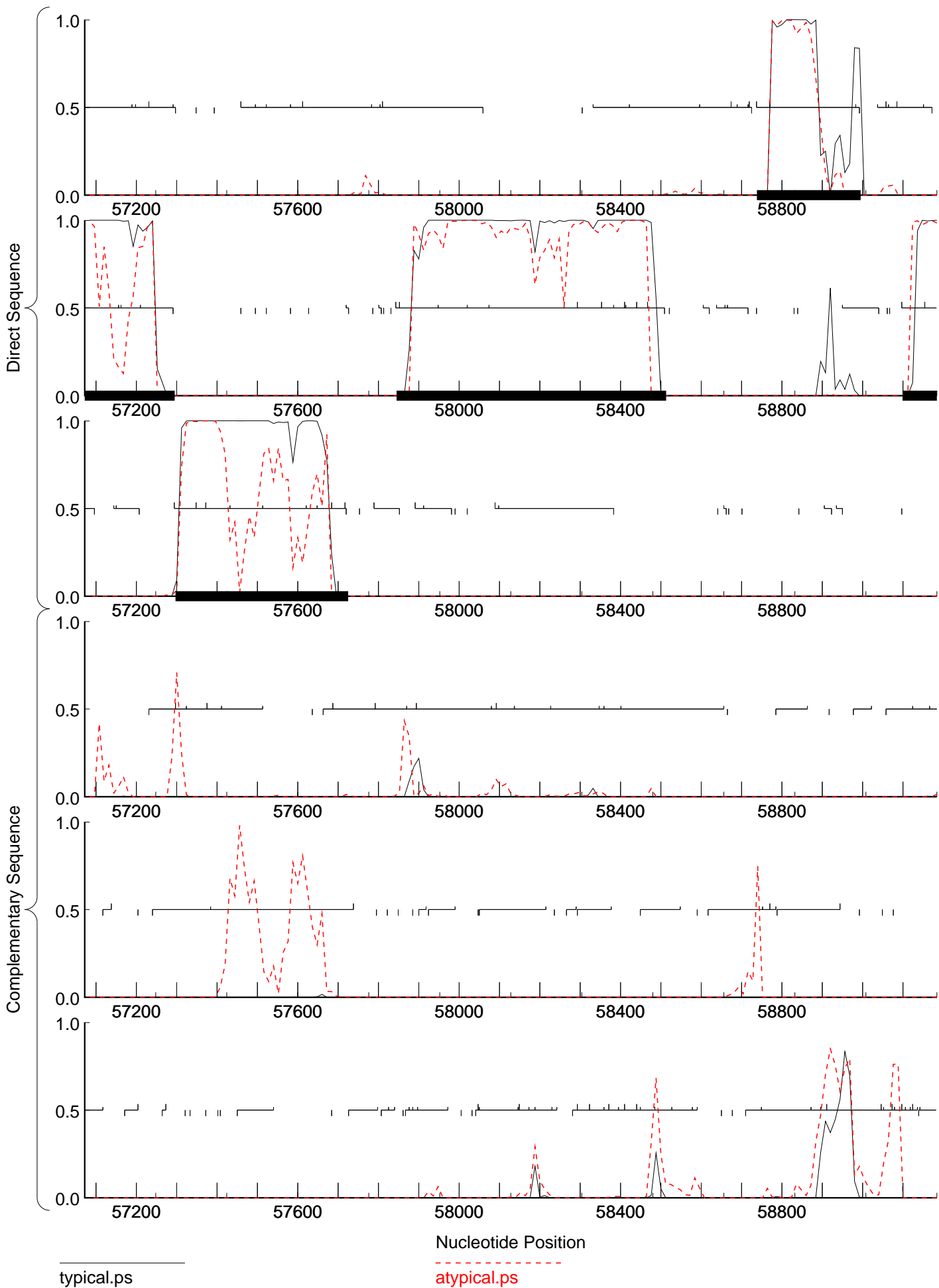
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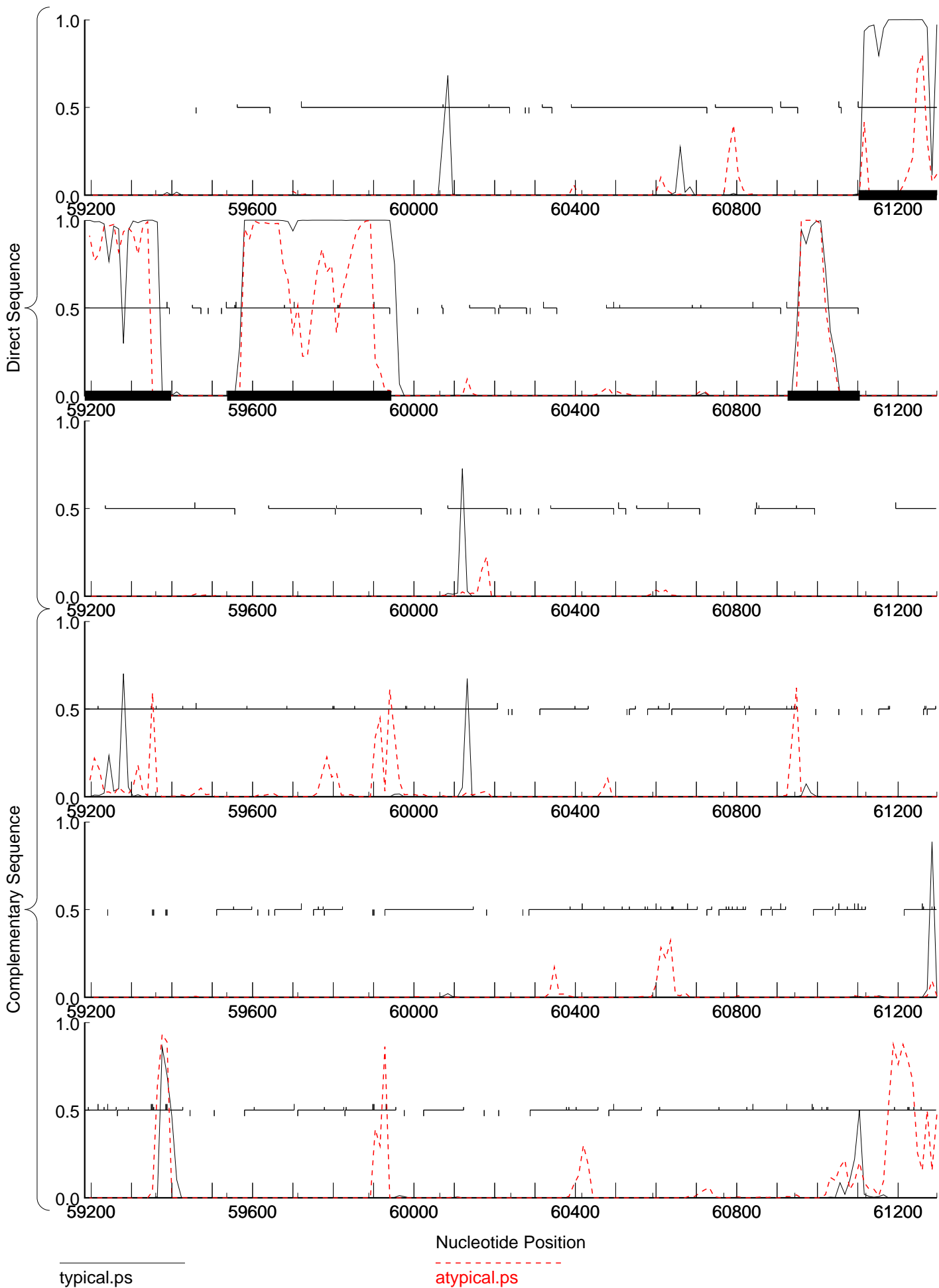
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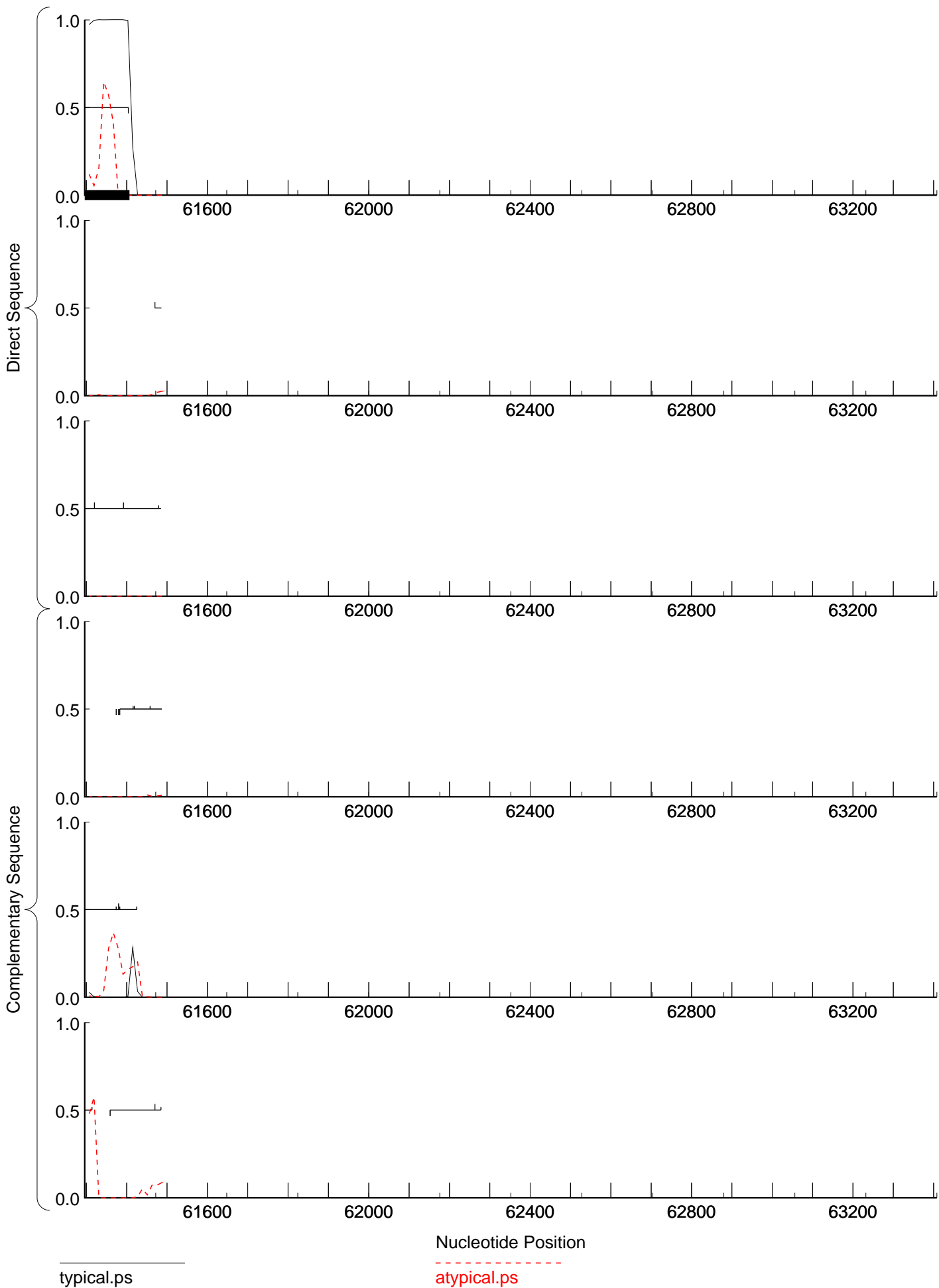
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typical.ps

atypical.ps





typical.ps

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