

# GeneMark

Version 2.5p (09.08.06)

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## PROGRAM INFORMATION

Sequence : Mycobacterium phage Heffalump complete sequence, 53085 bp including 10-base 3' overhang (CG  
Analysis Date : 10/23/17 at 15:04:30  
Pages : 27  
Sequence Length : 53085 bp  
GC Content : 63.58%

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,

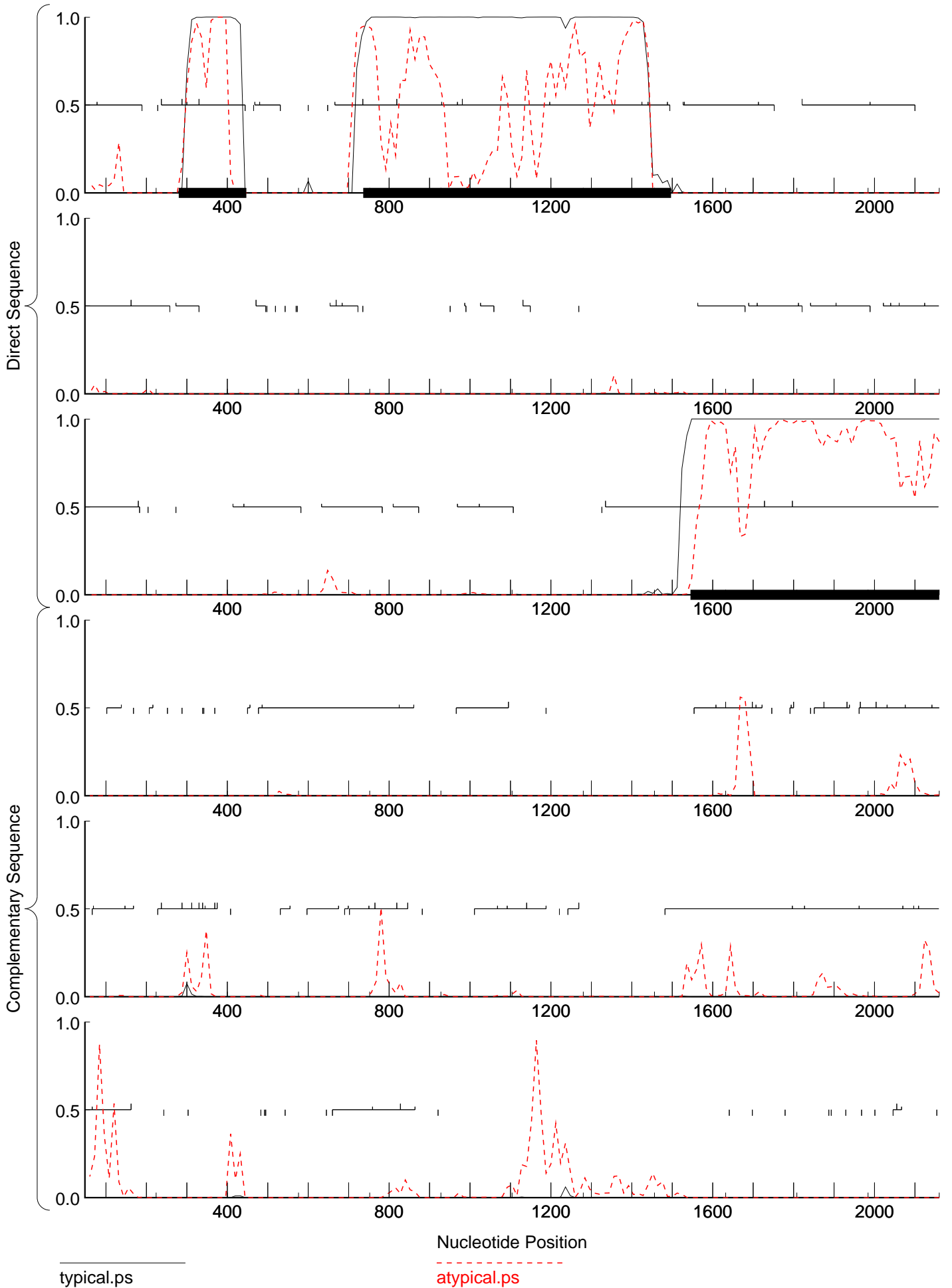
## Matrix notes & comments

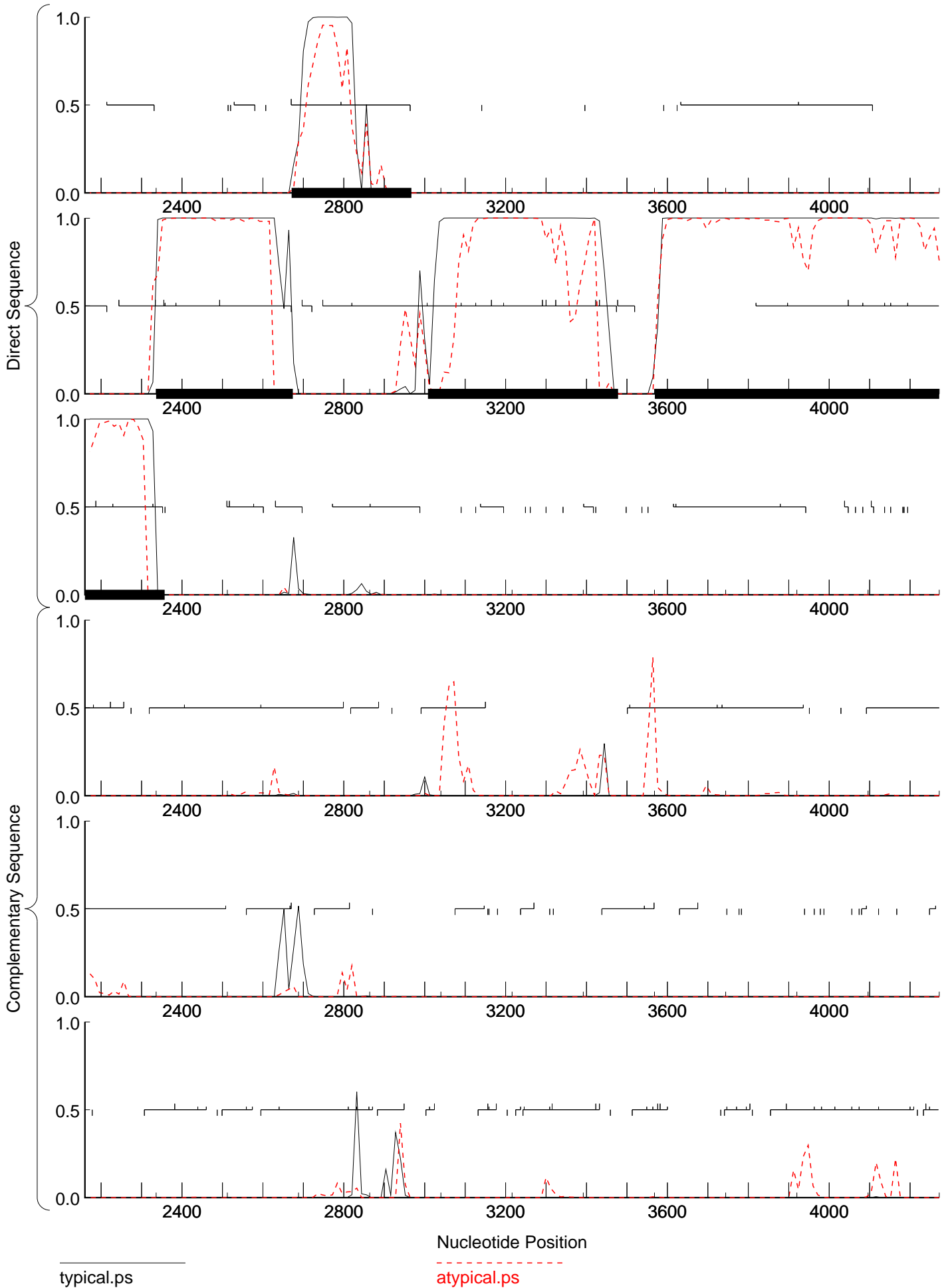
Training set derived by GeneMarkS, 4.30 October 2014  
Mon Oct 23 15:04:29 2017

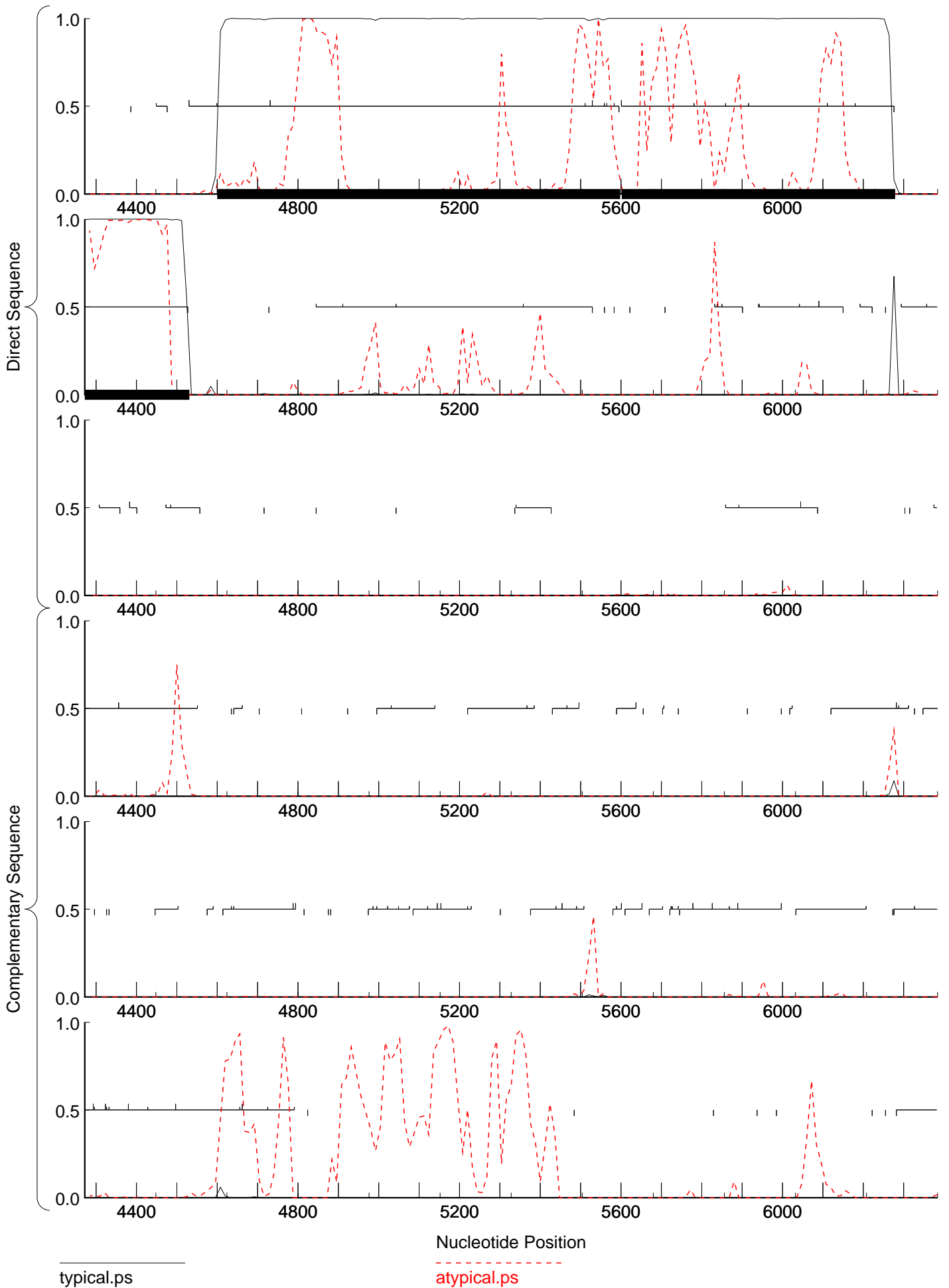
## MATRIX INFORMATION

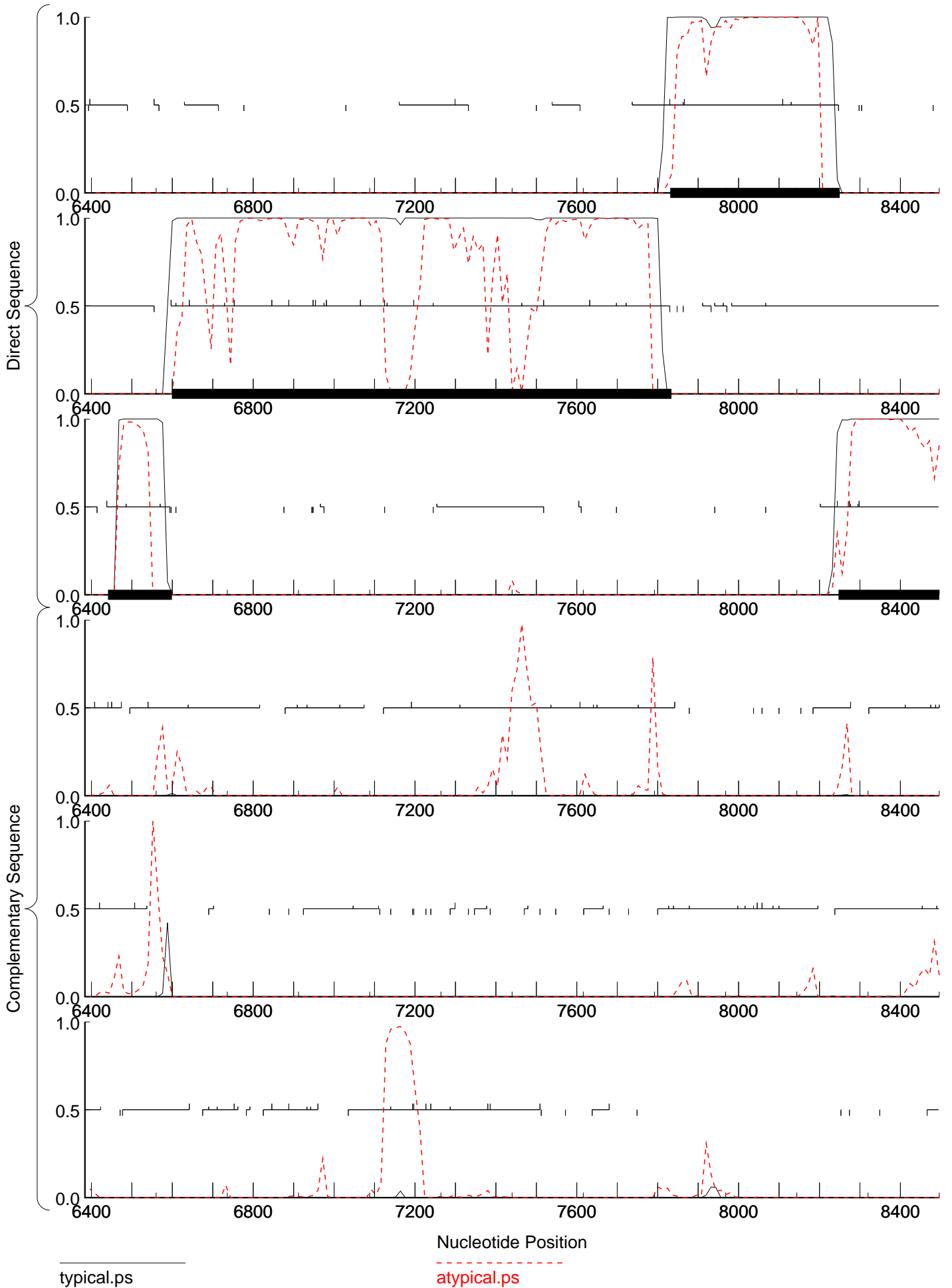
Matrix : -  
Author : -  
Order : 4

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



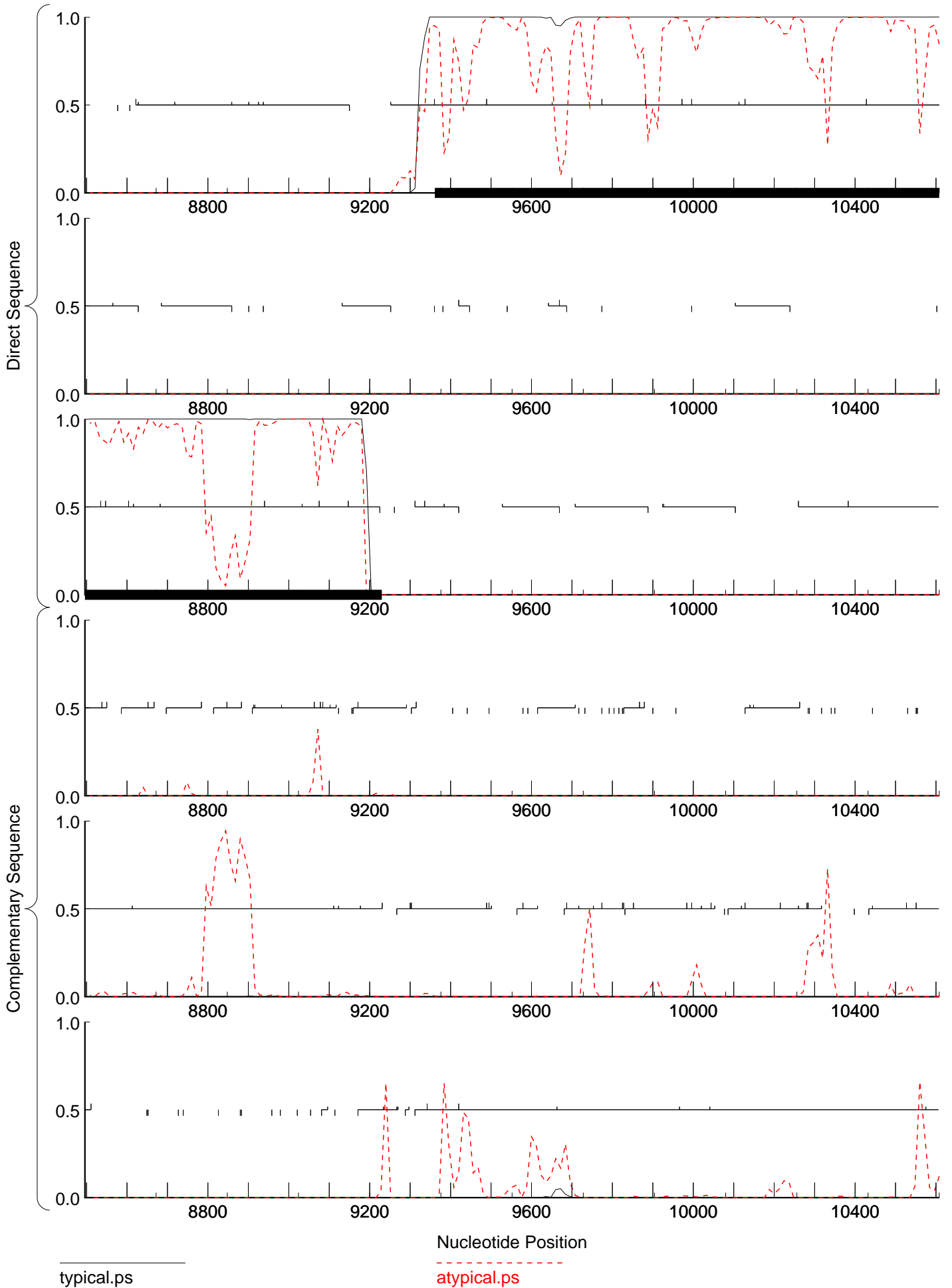


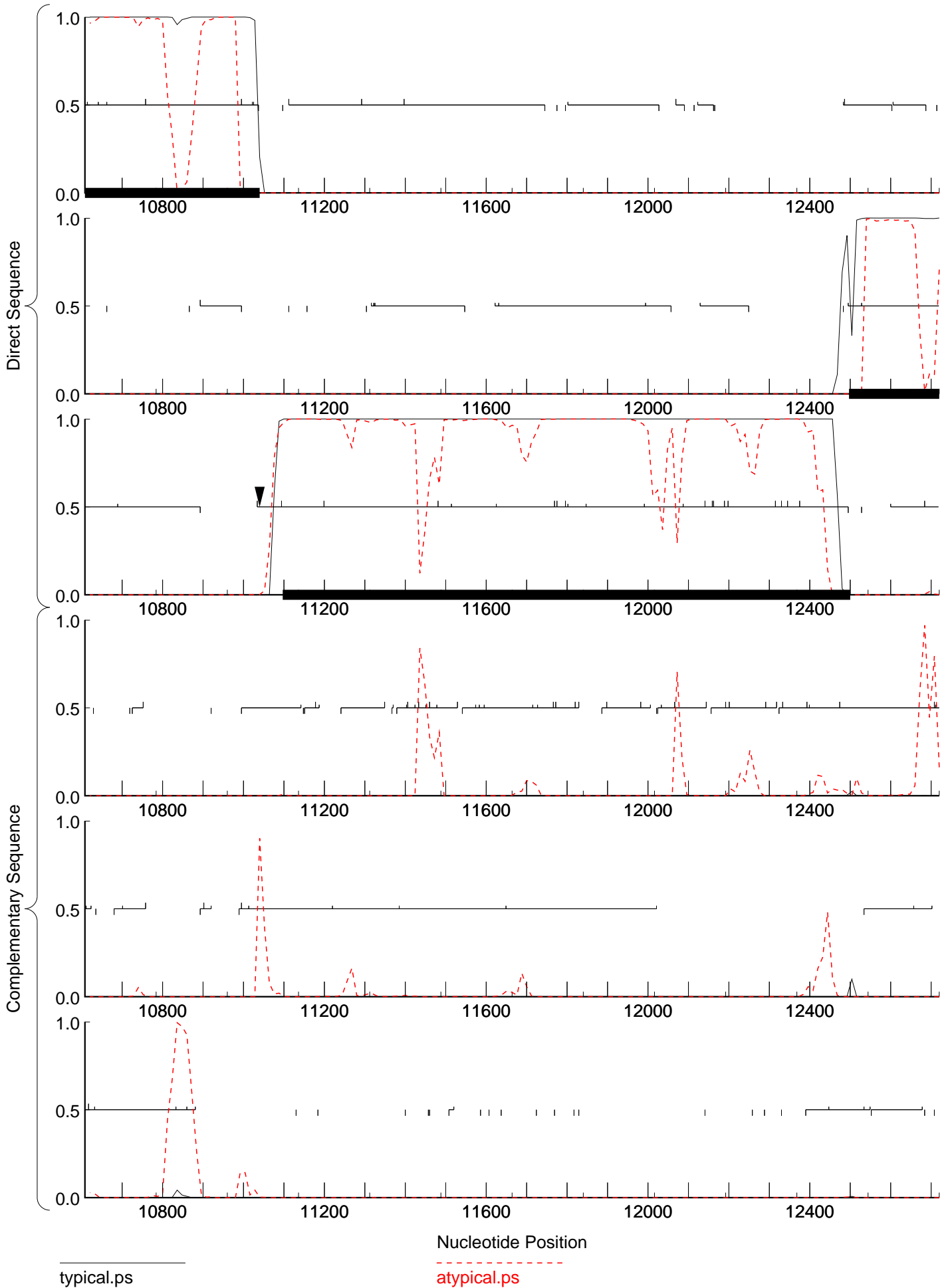




typical.ps

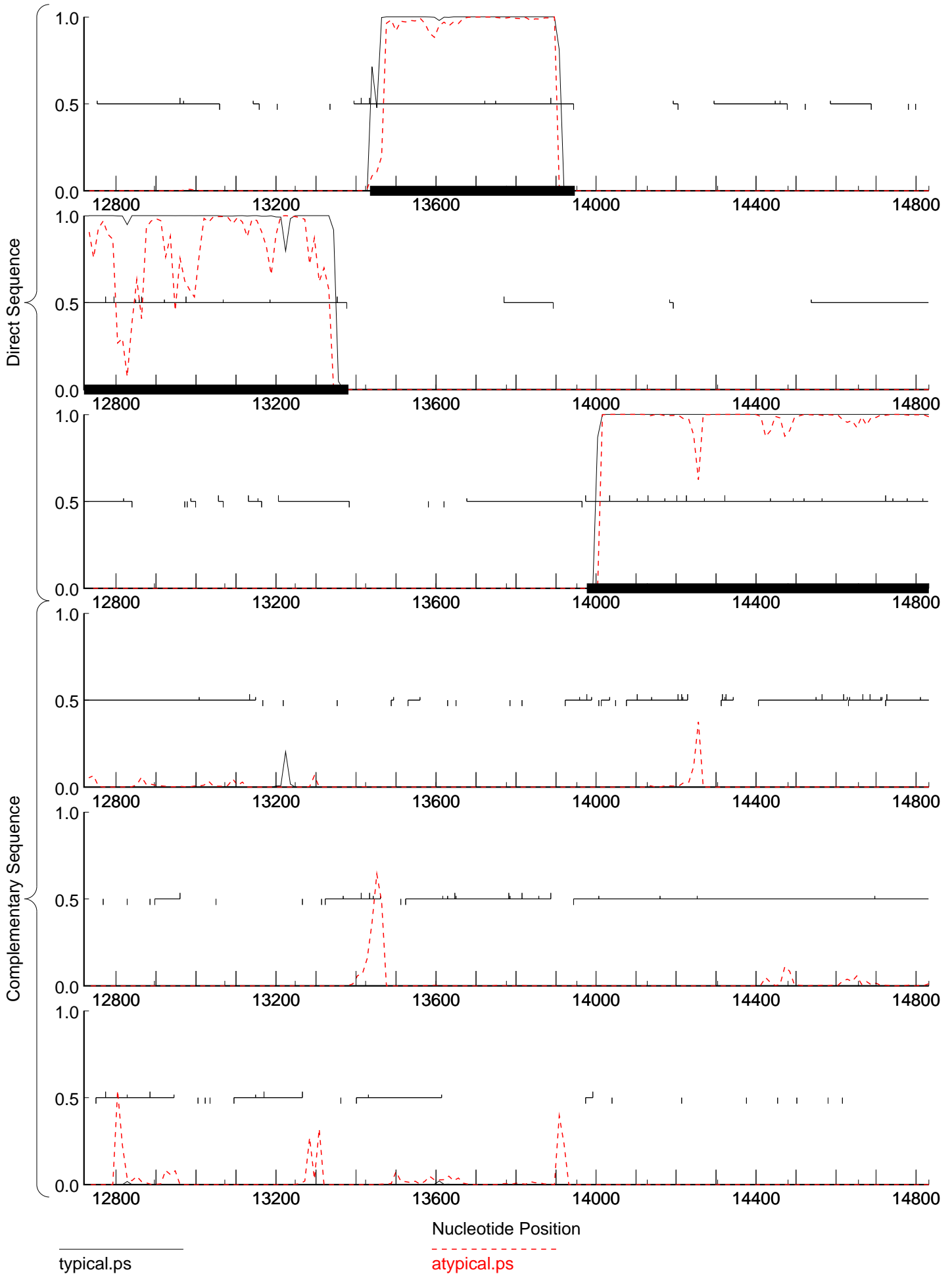
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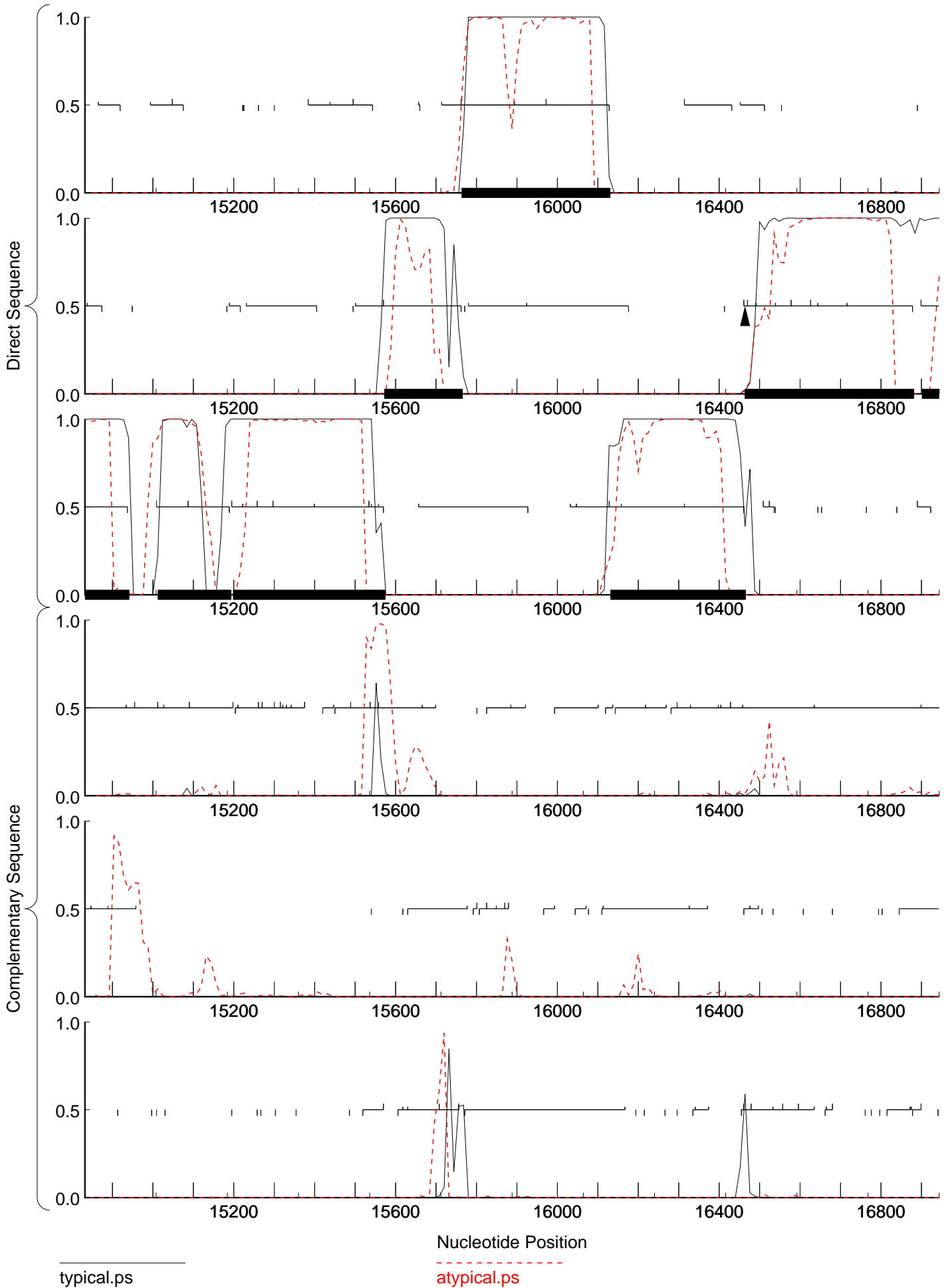


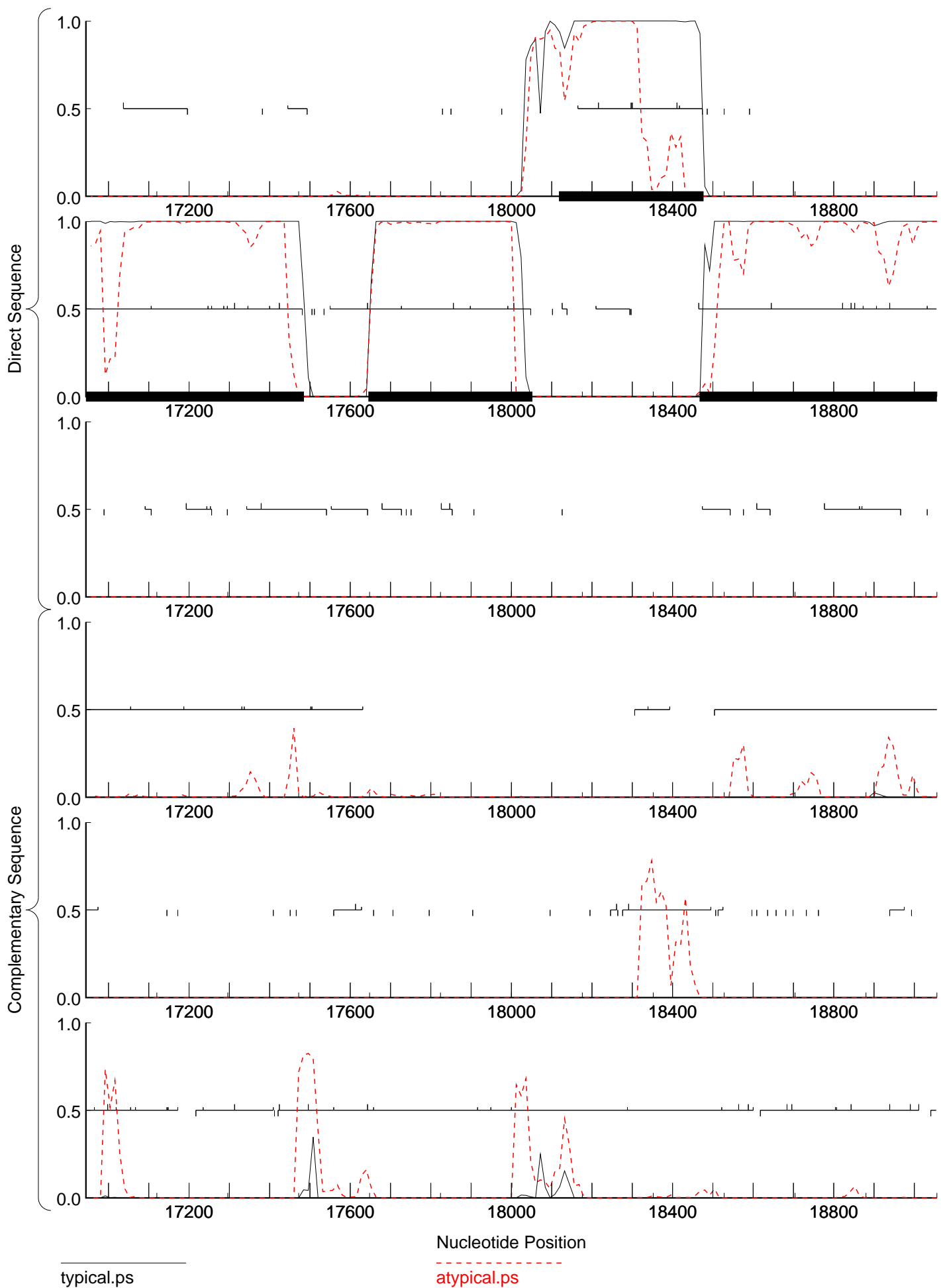
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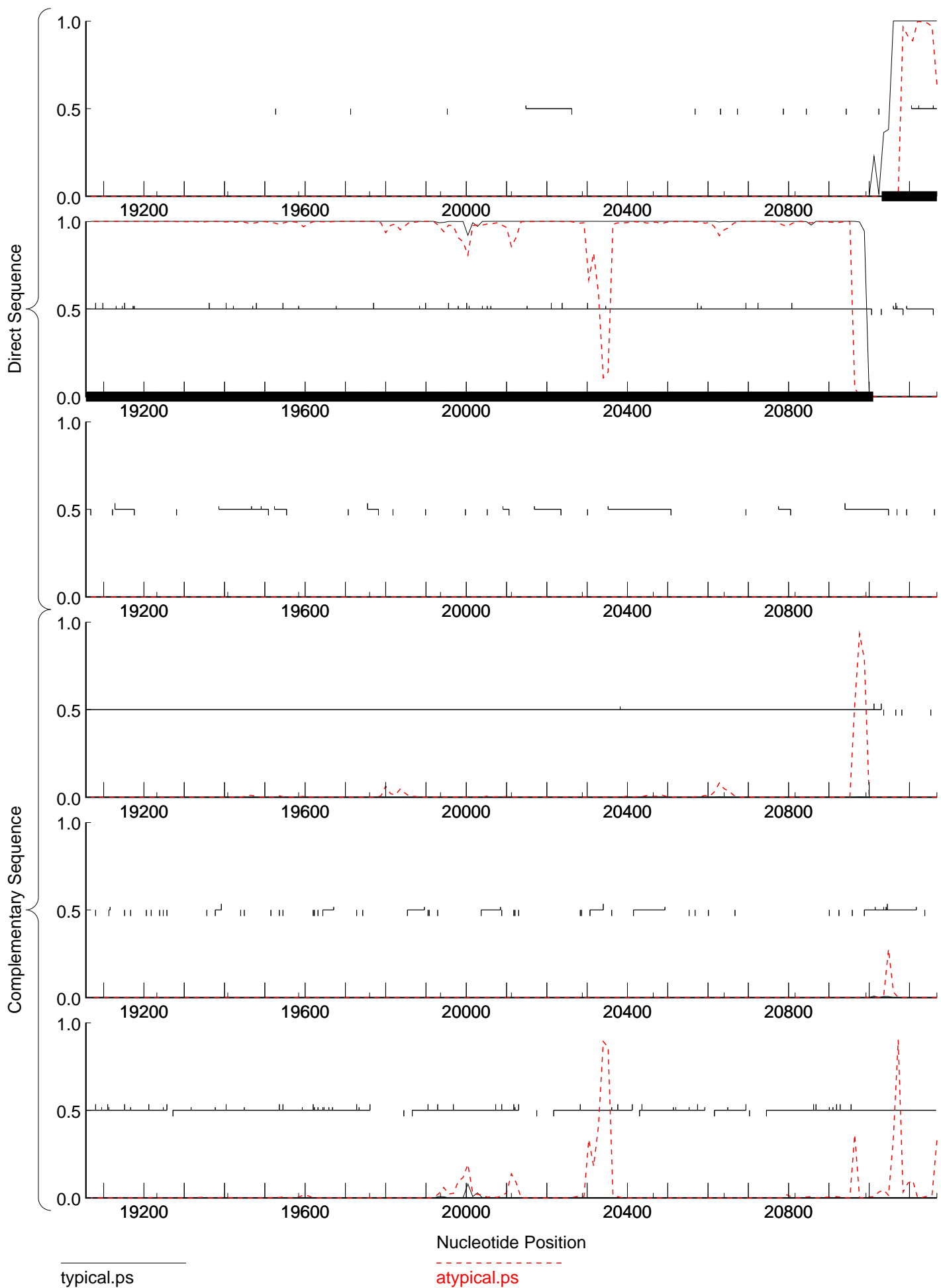
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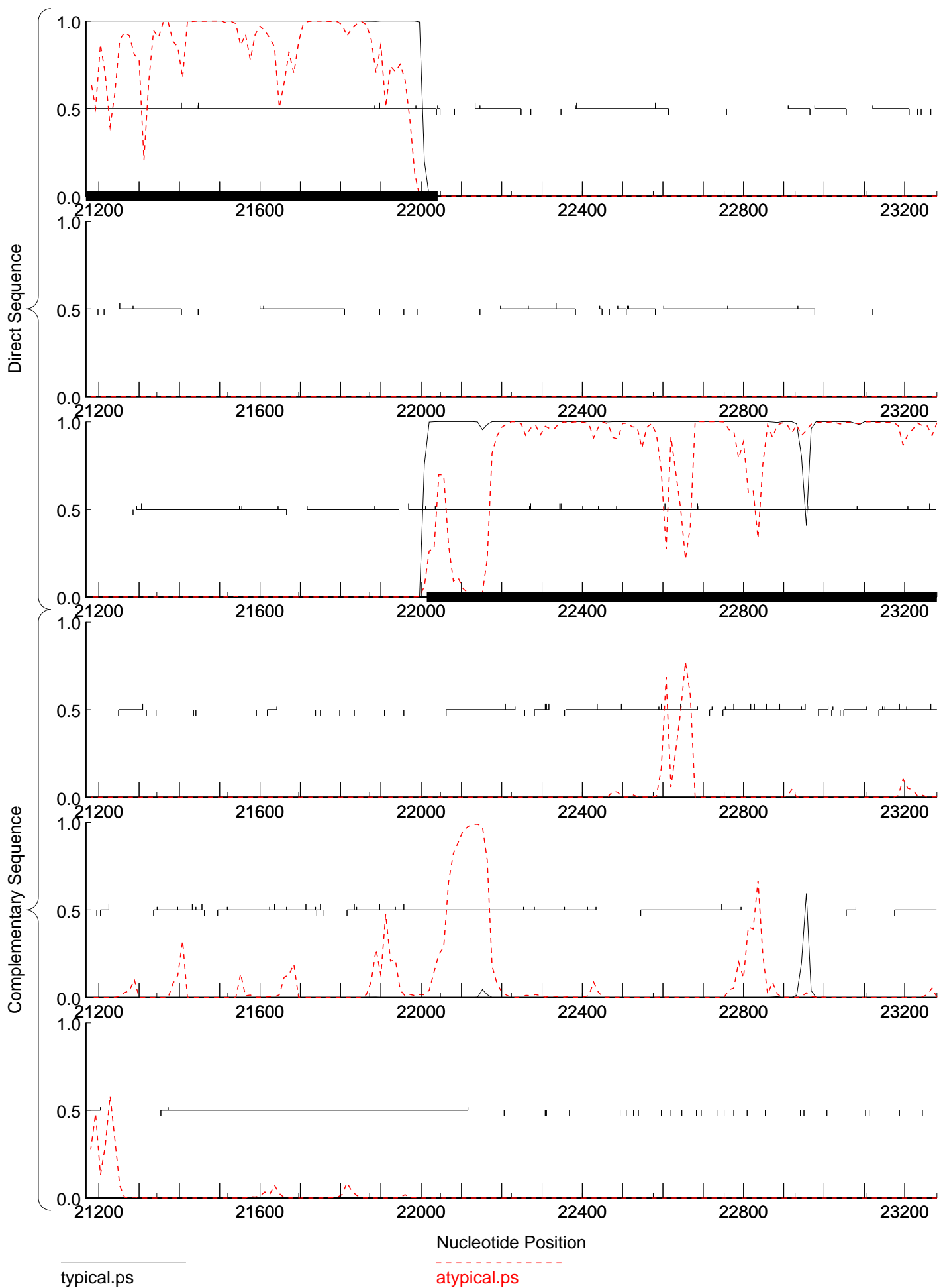


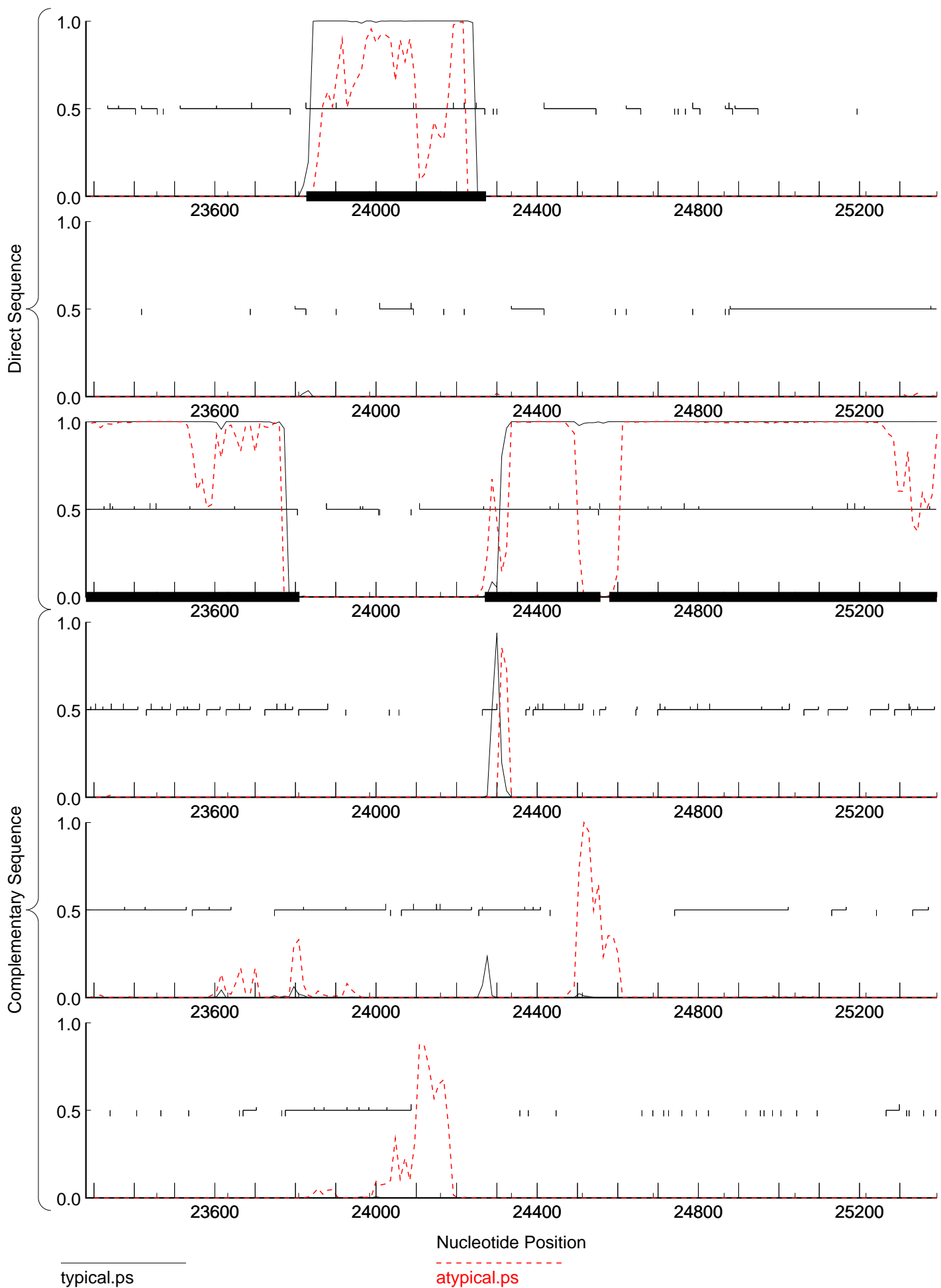


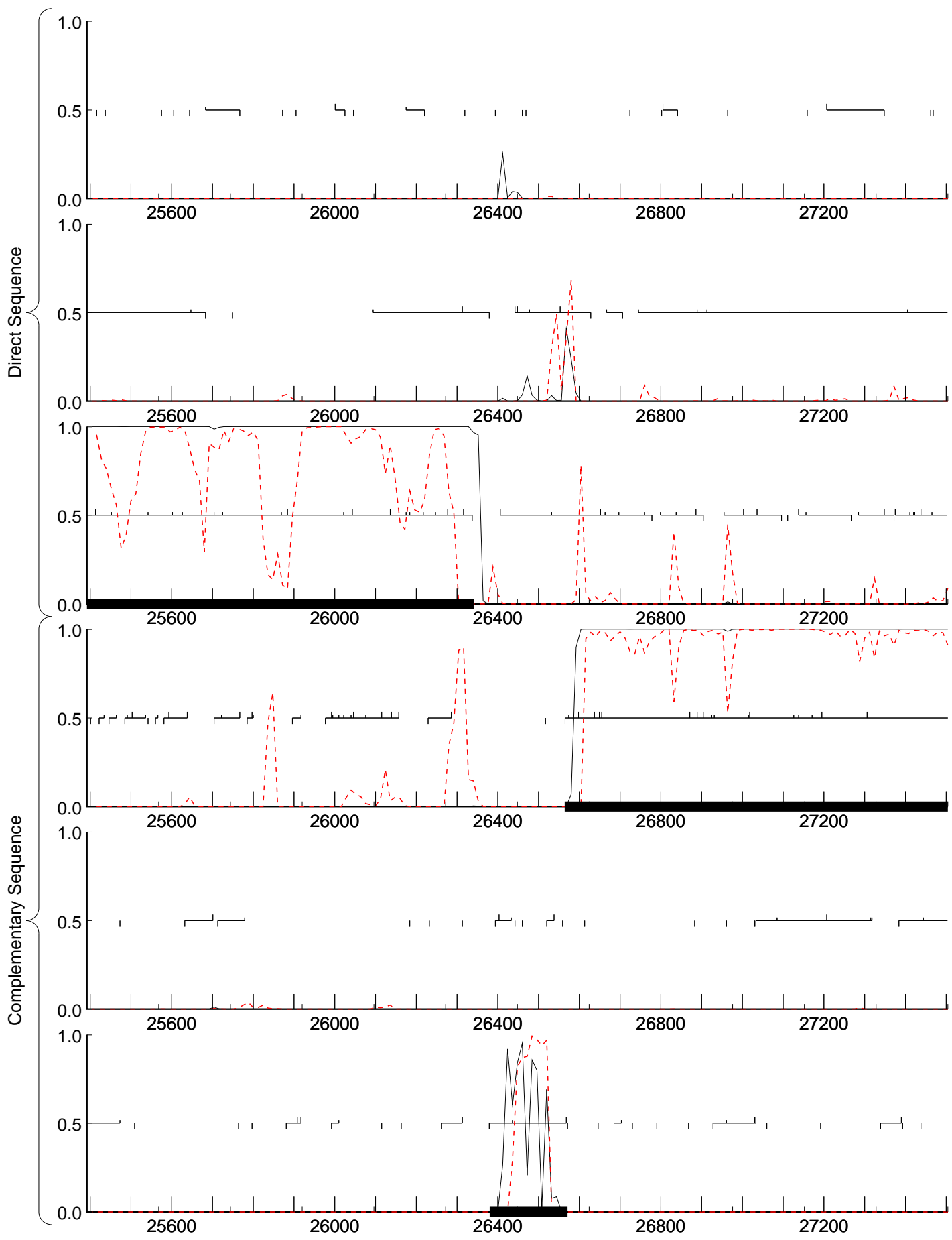




complete sequence, 53085 bp including 10-base 3' overhang (CGTTCGGTTA), Cluster A2, Order 4, Window 96, Step 12, 12/27  
complete sequence, 53085 bp including 10-base 3' overhang (CGTTCGGTTA), Cluster A2, Order 2, Window 96, Step 12, 12/27

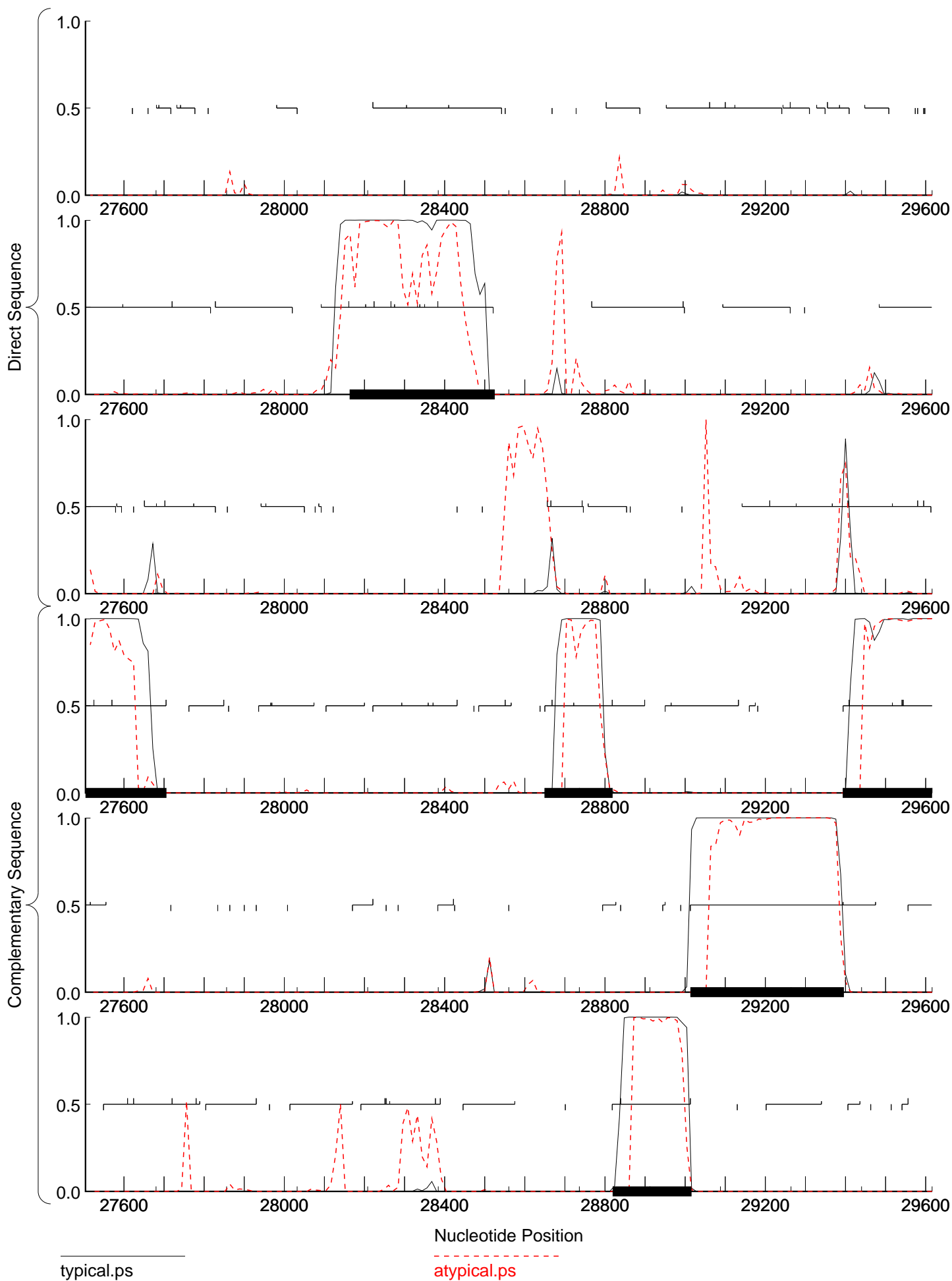


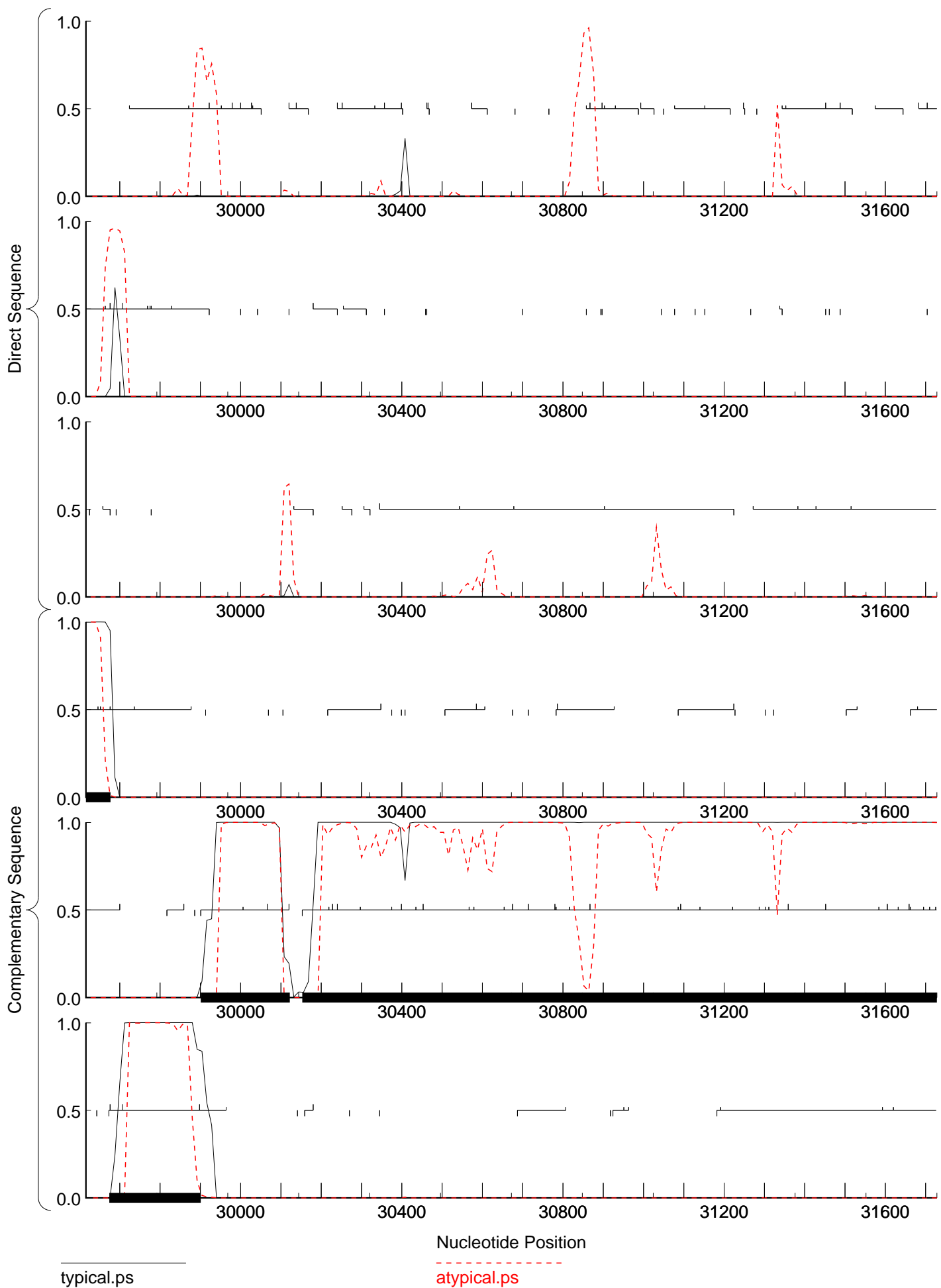




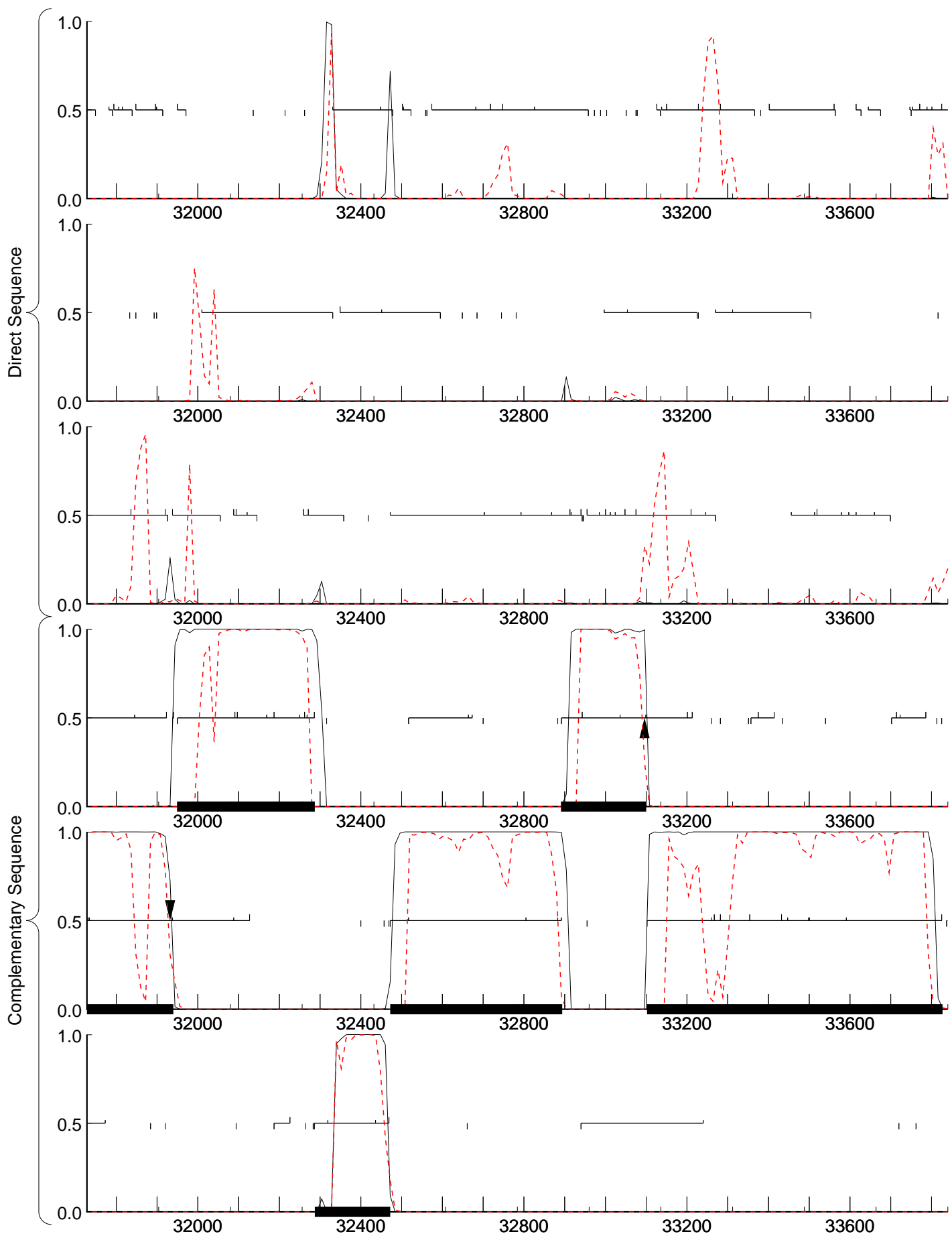
typical.ps

atypical.ps



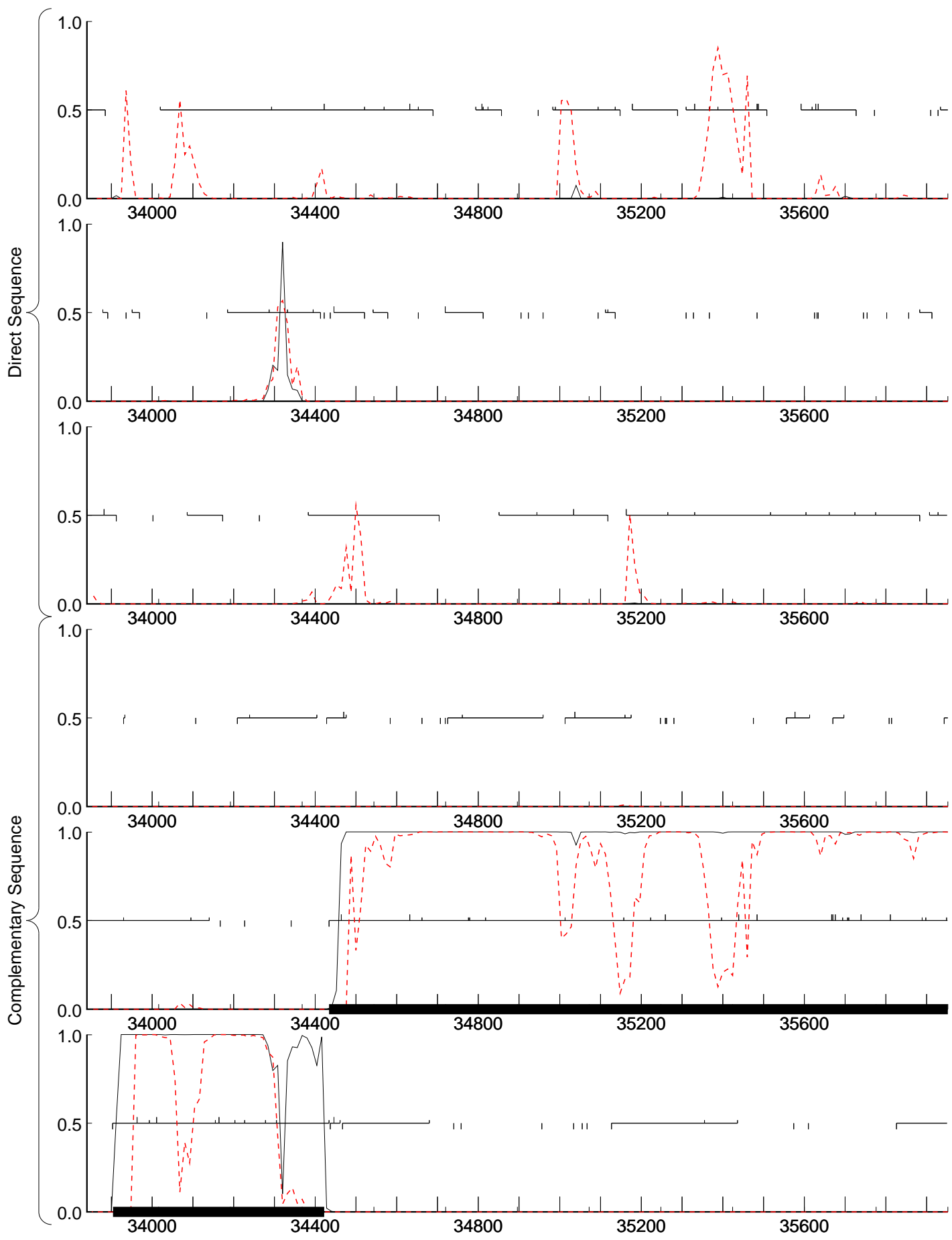






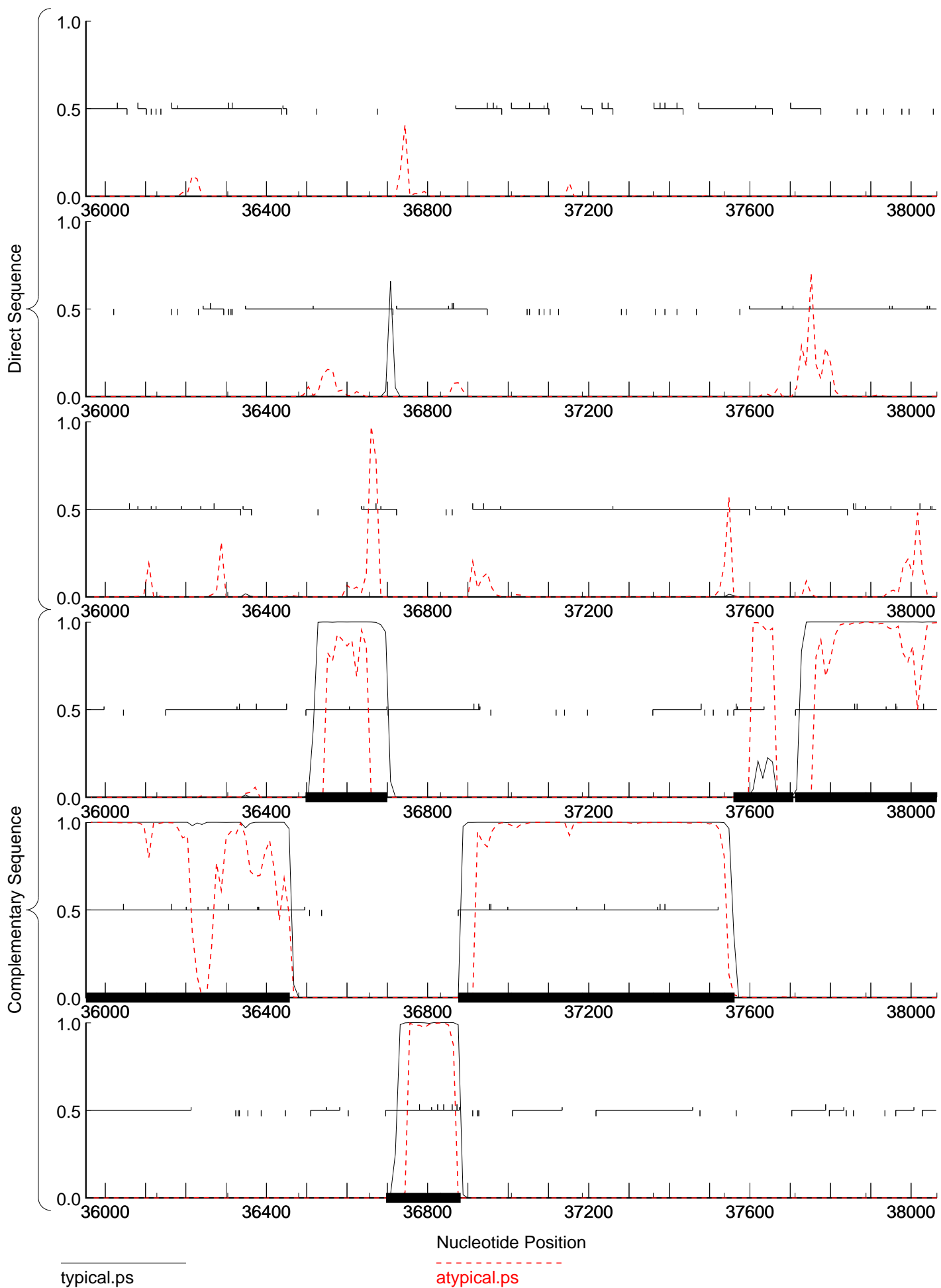
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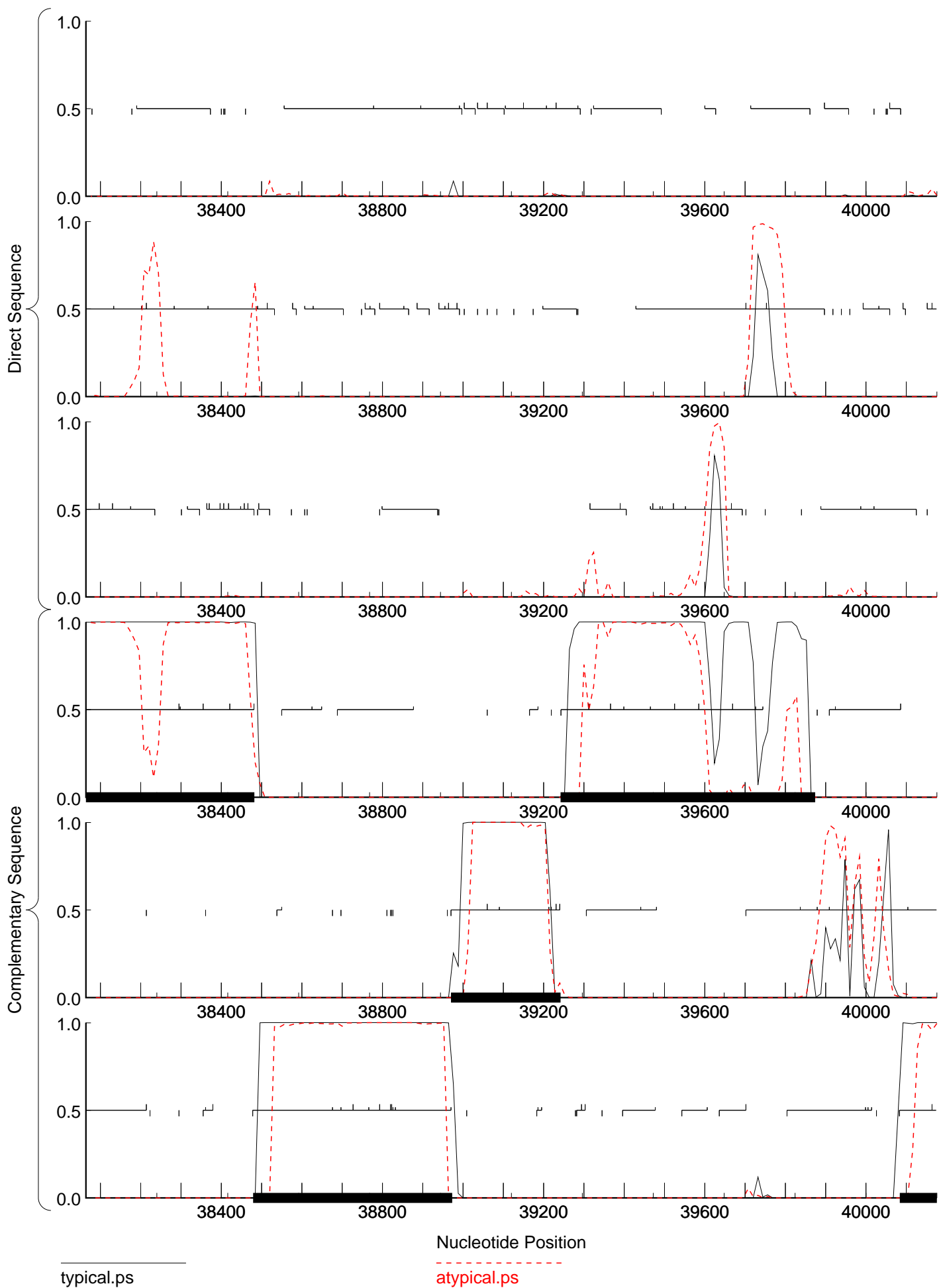
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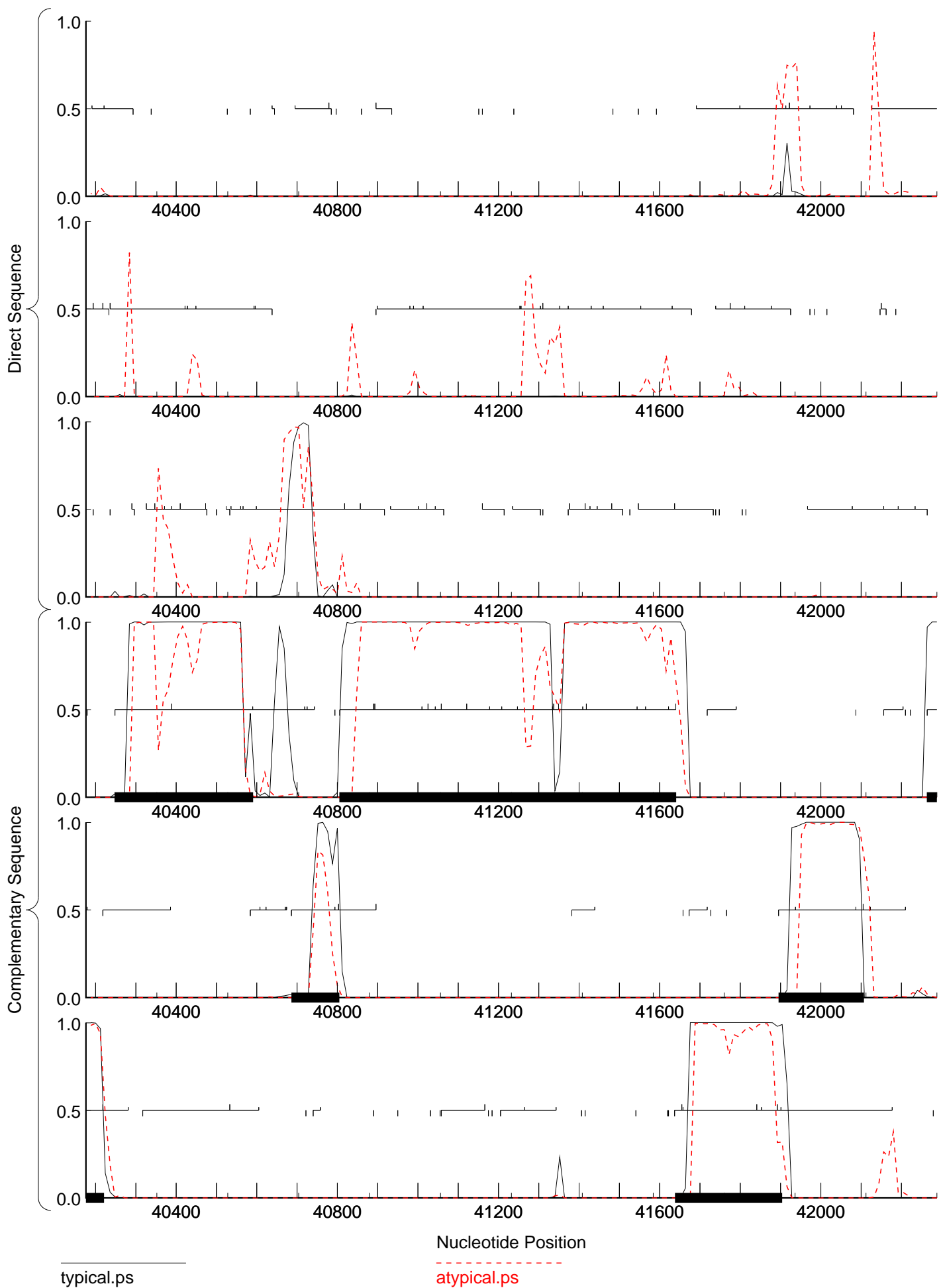


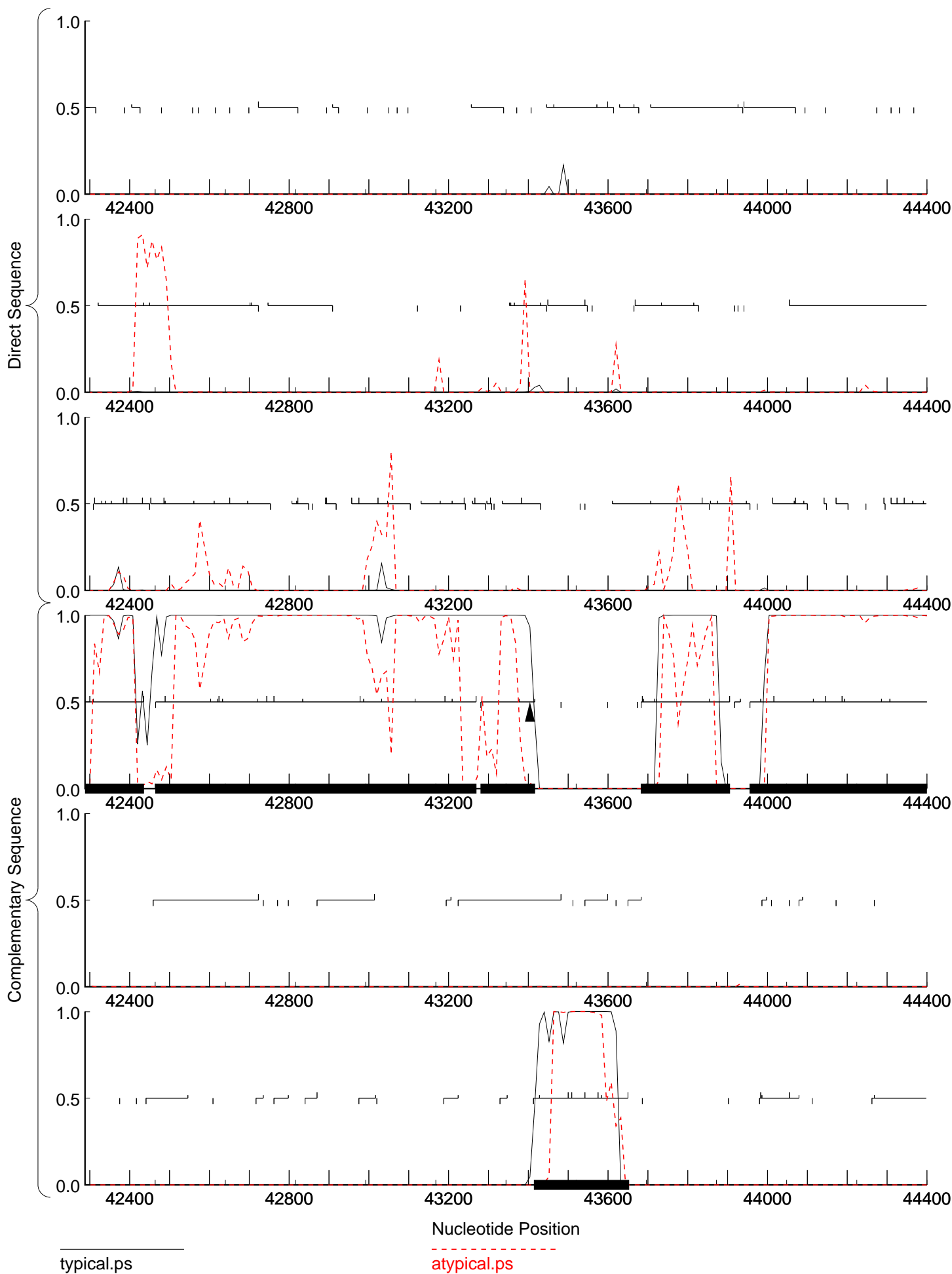
typical.ps

atypical.ps



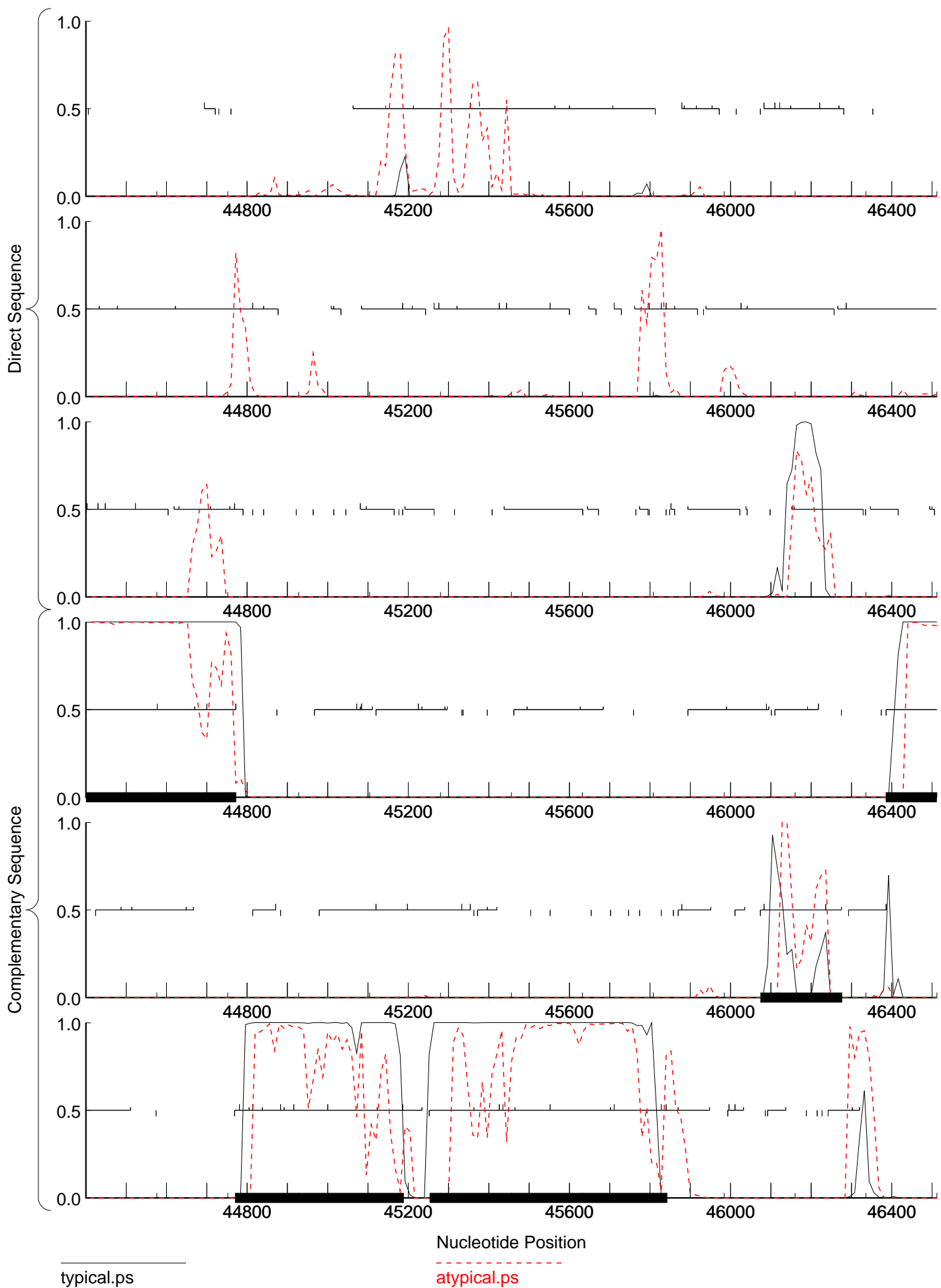


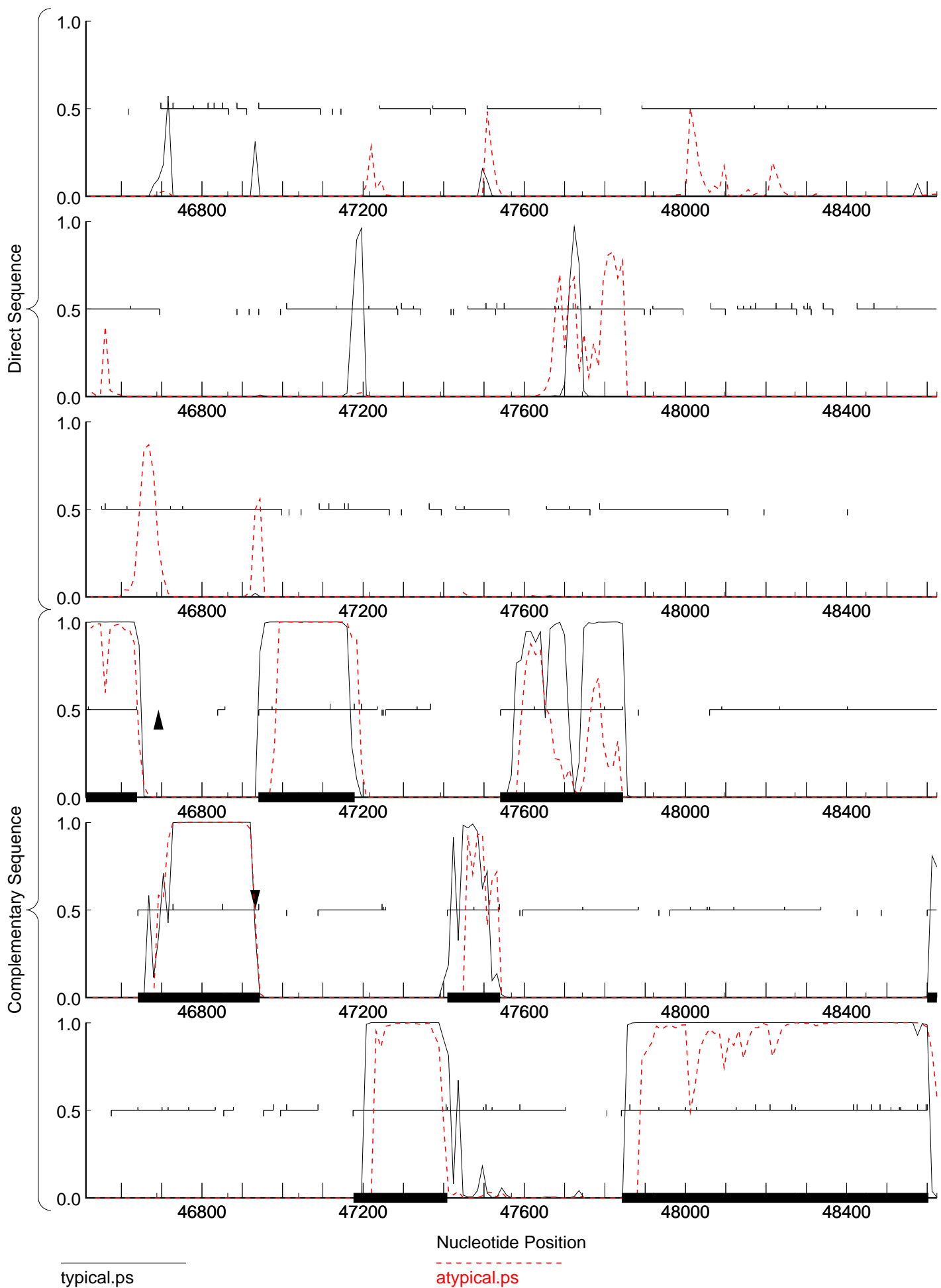




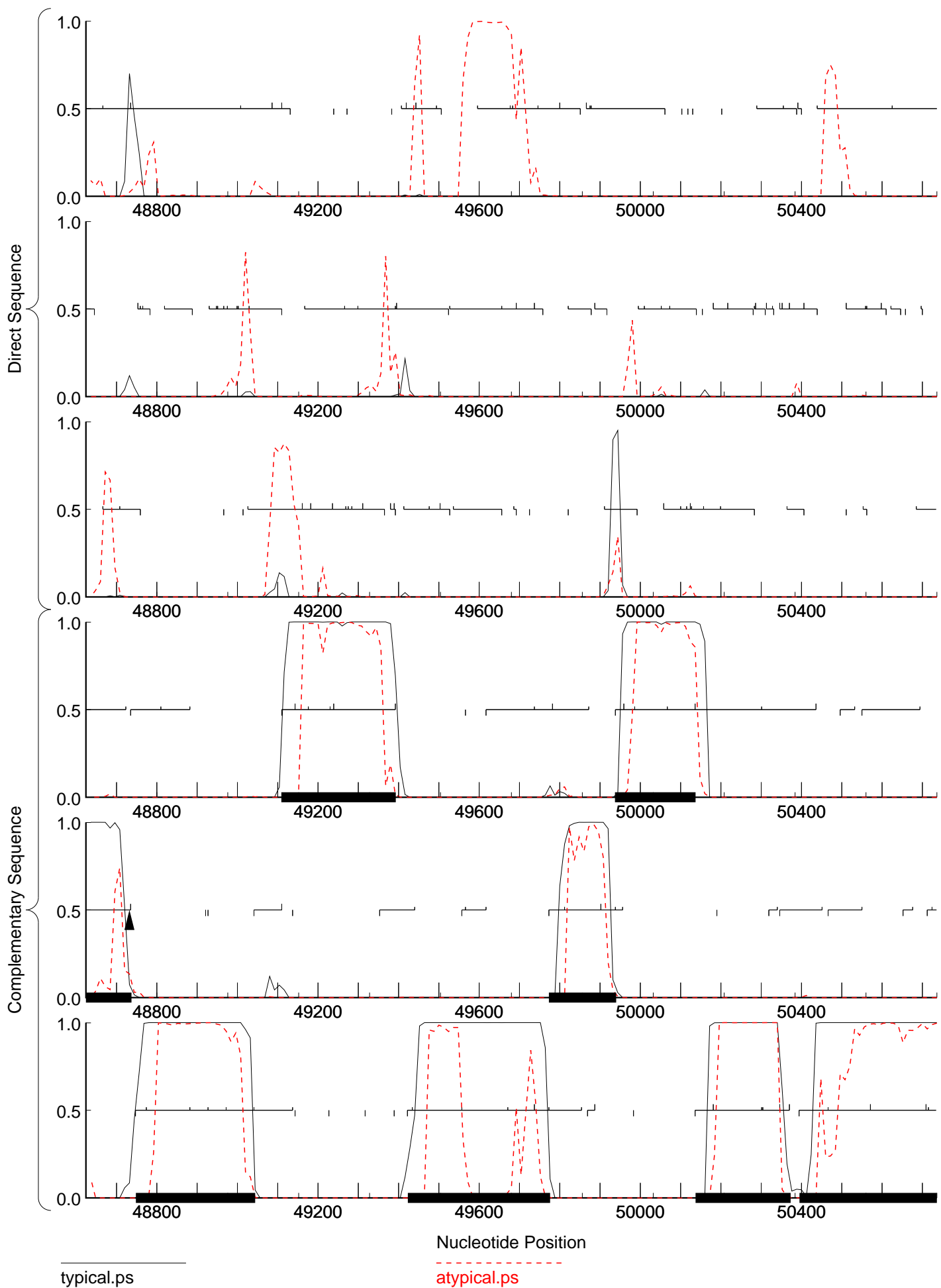
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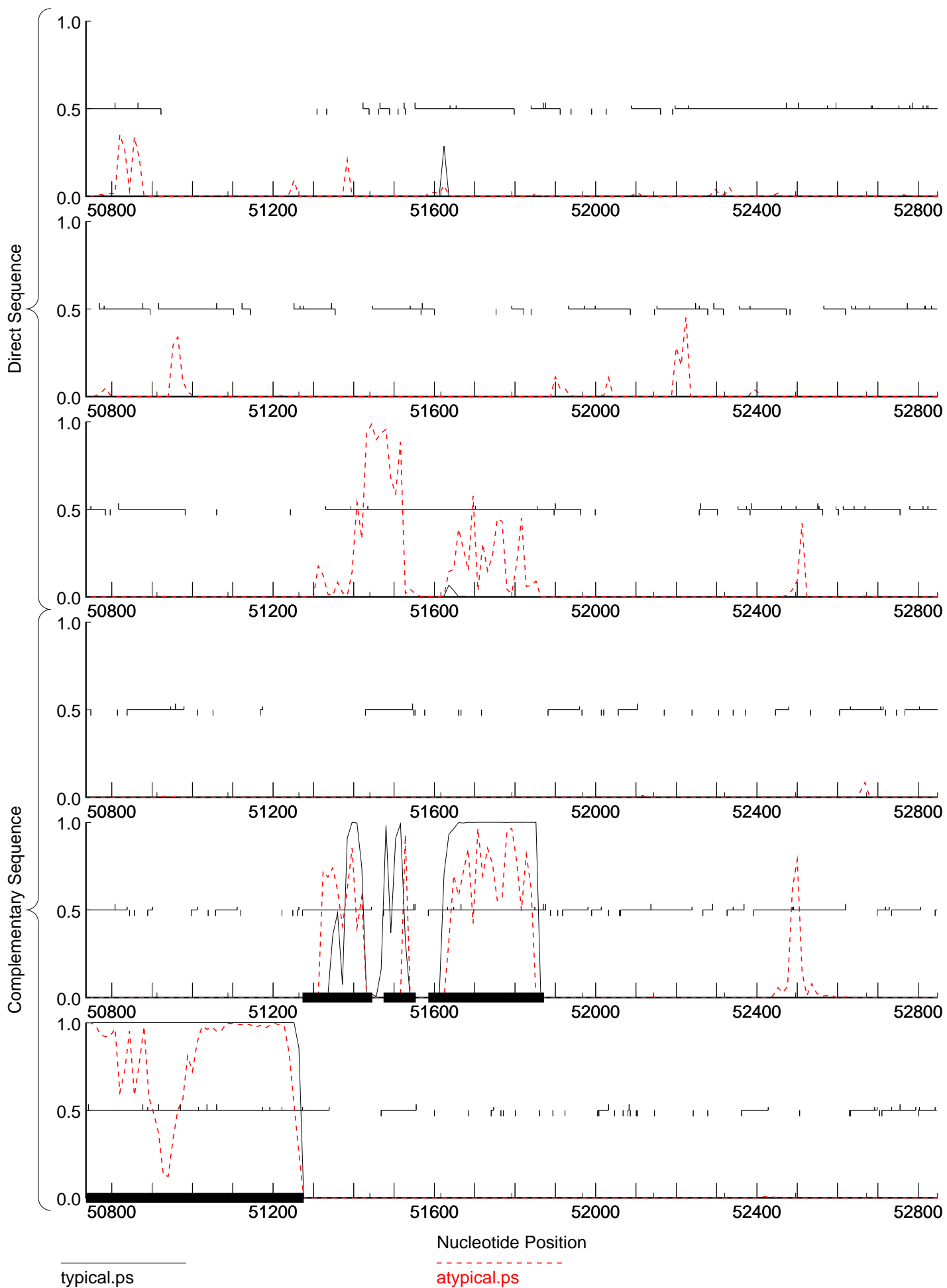
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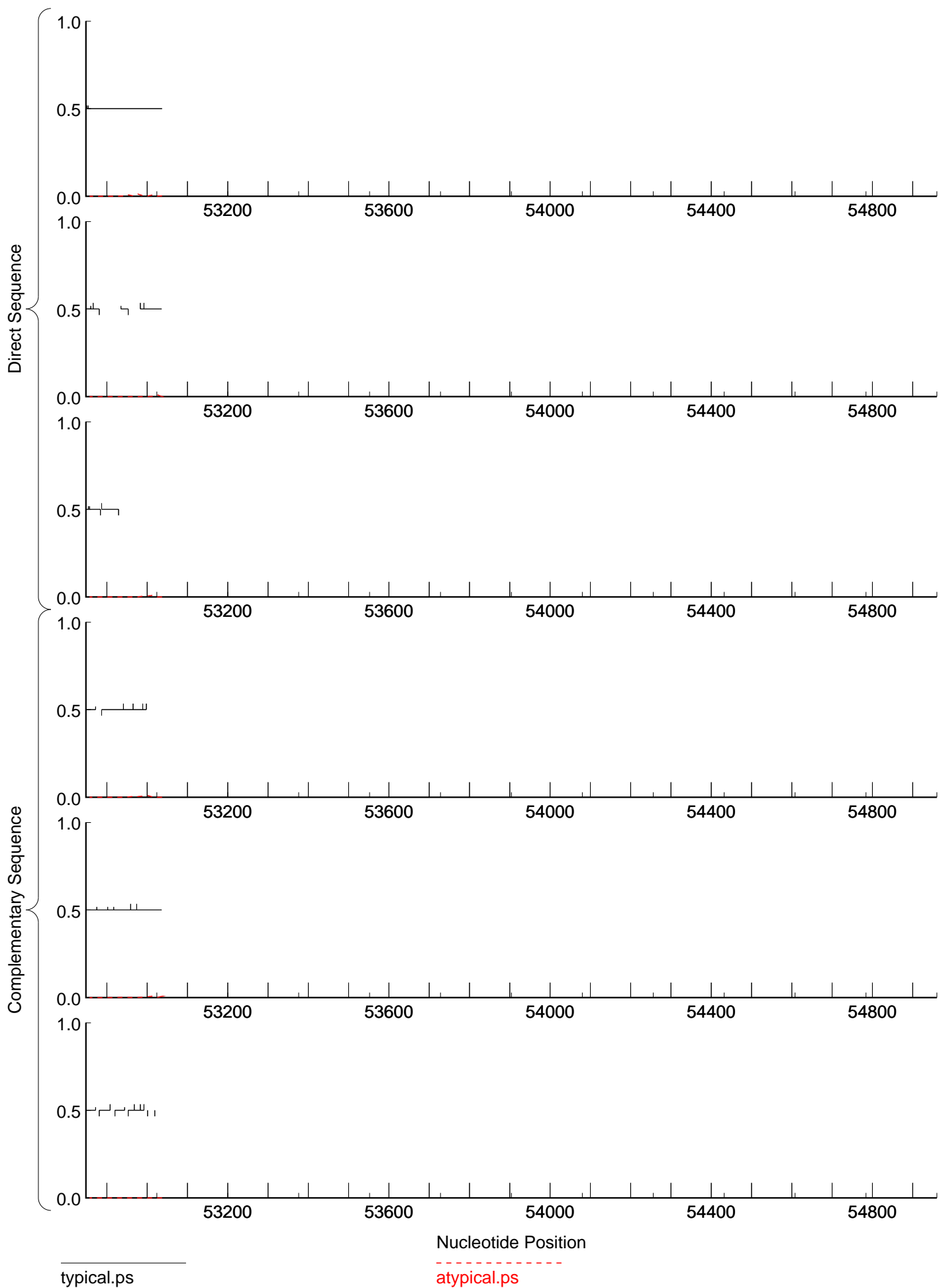












typical.ps

atypical.ps