

# GeneMark

Version 2.5p (09.08.06)

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## PROGRAM INFORMATION

Sequence : Mycobacterium phage Kingsley complete sequence, 60300 bp including 10-base 3' overhang (CCG  
Analysis Date : 2/23/18 at 17:22:10  
Pages : 30  
Sequence Length : 60300 bp  
GC Content : 61.38%

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

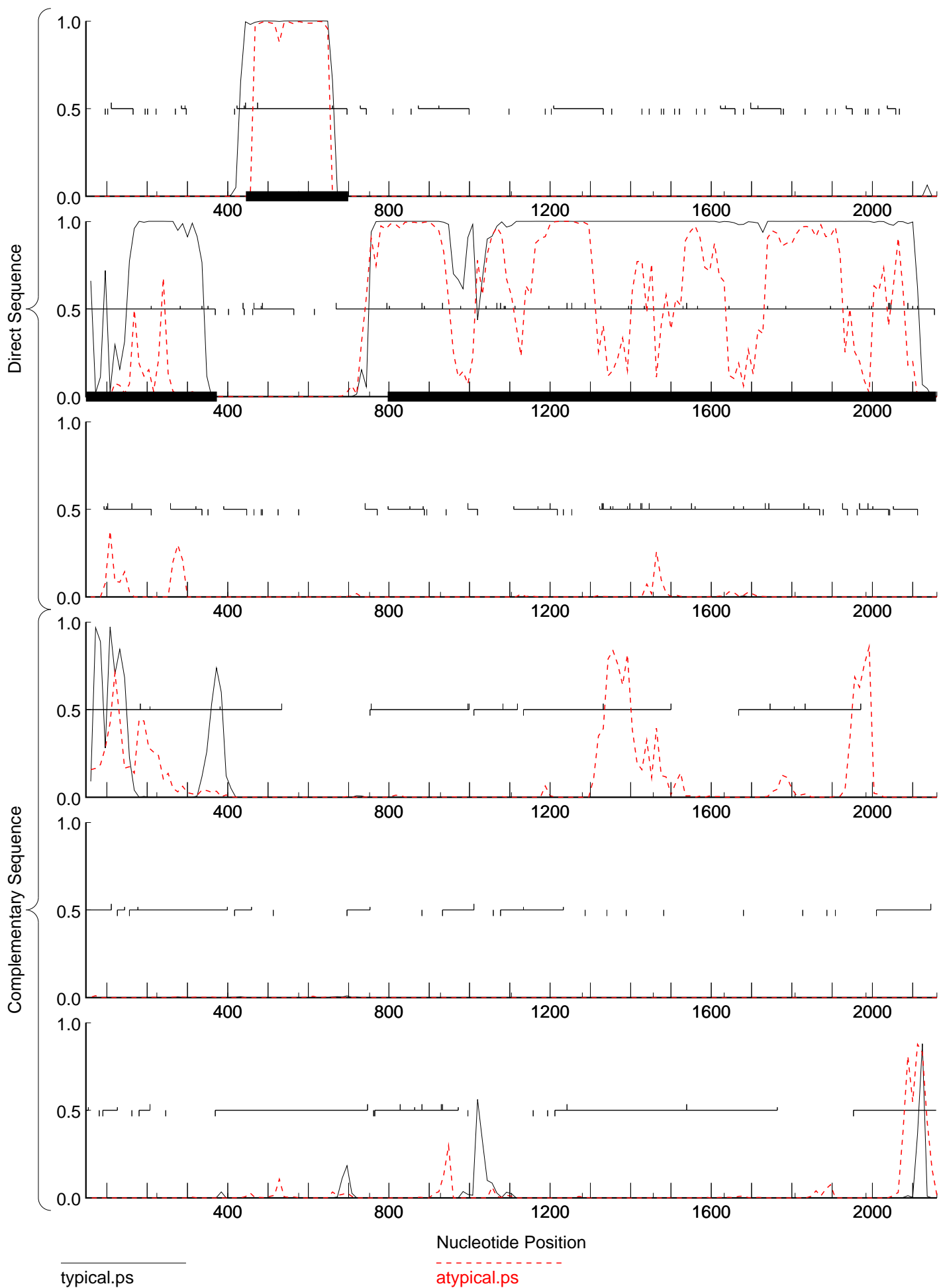
Training set derived by GeneMarkS, 4.30 October 2014  
Fri Feb 23 17:22:10 2018

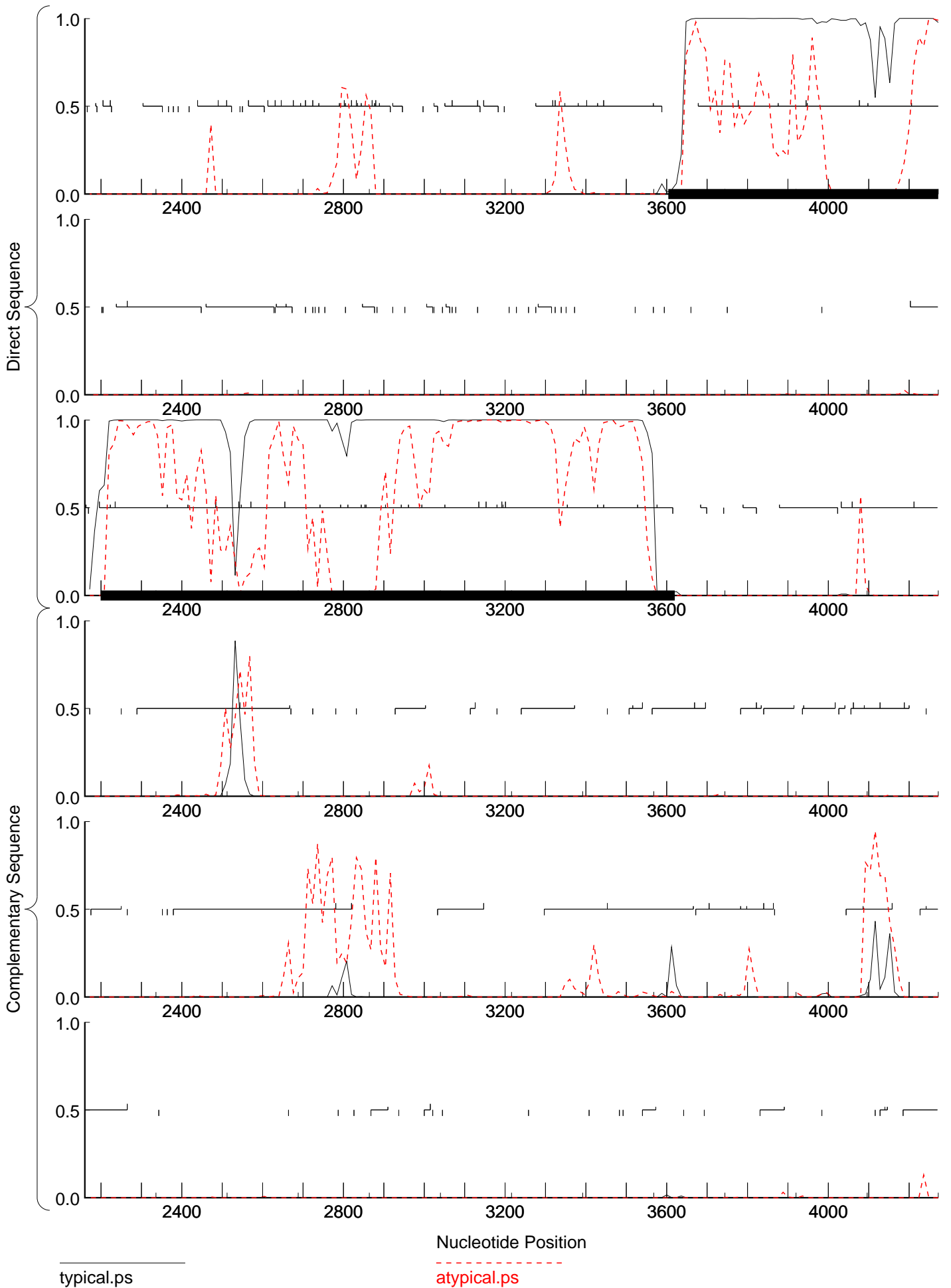
## MATRIX INFORMATION

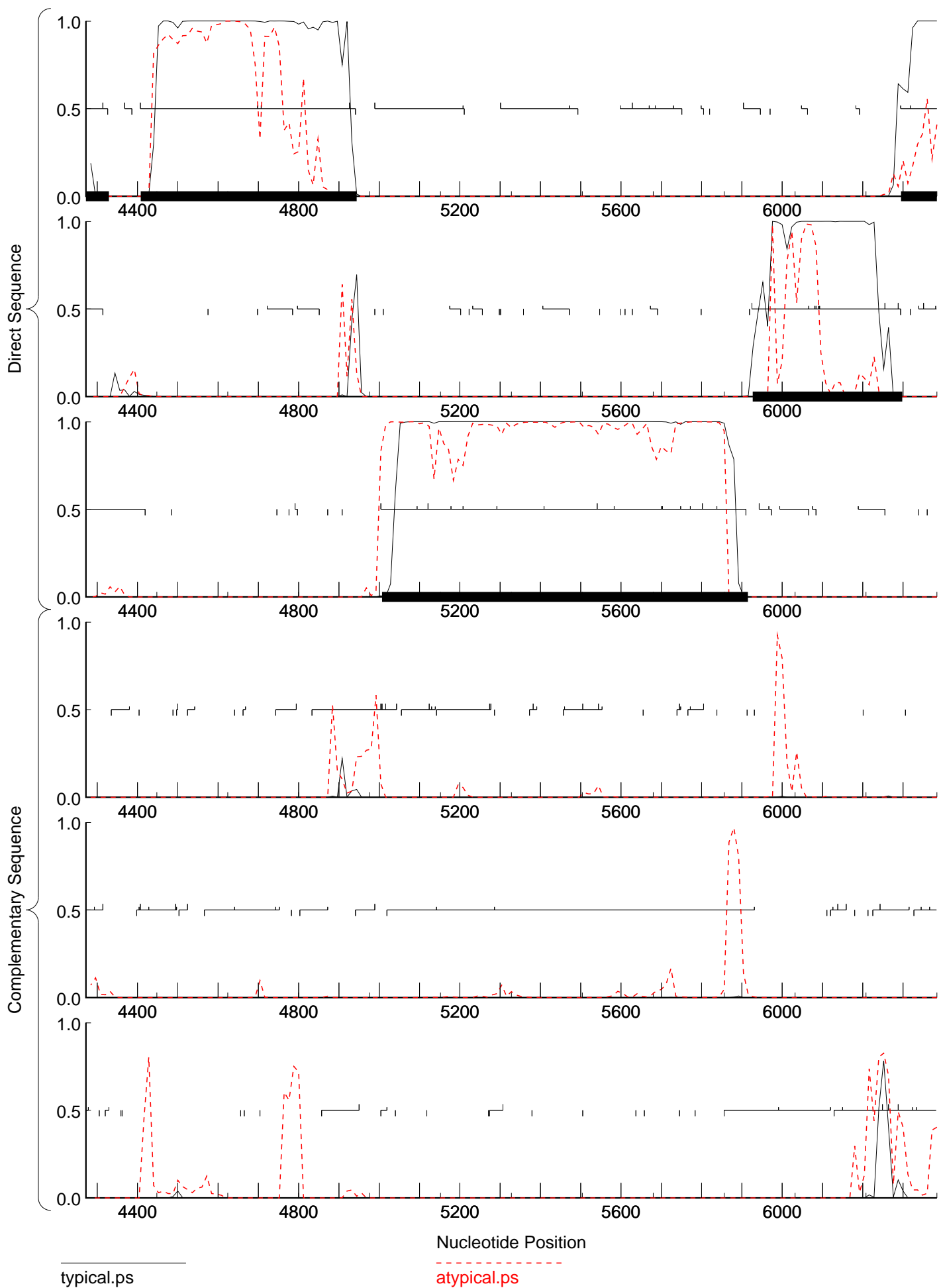
Matrix : -  
Author : -  
Order : 4

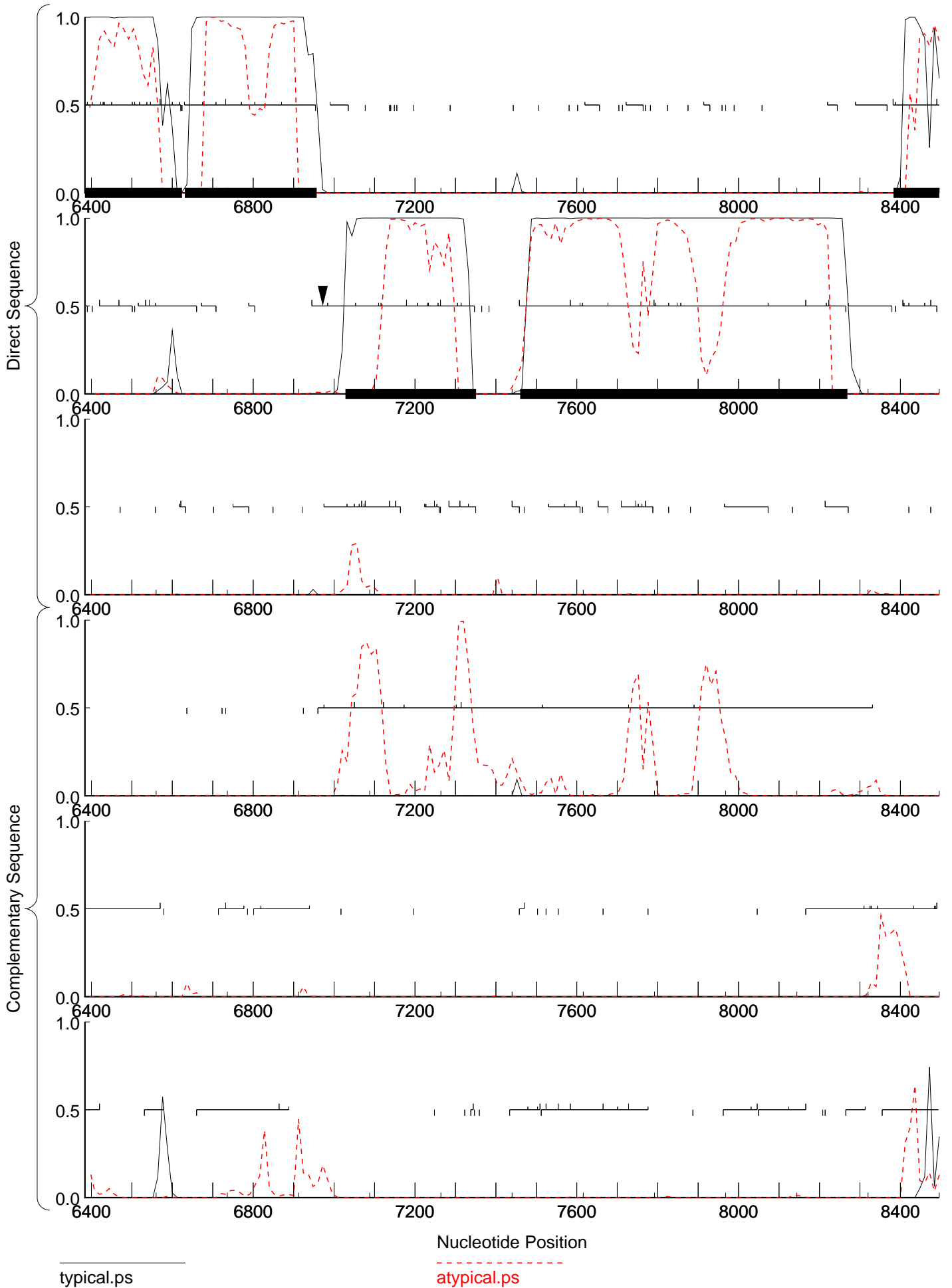
Send questions / comments to:  
Dr. M. Borodovsky  
Georgia Institute of Technology  
School of Biology  
Atlanta, GA 30332-0230

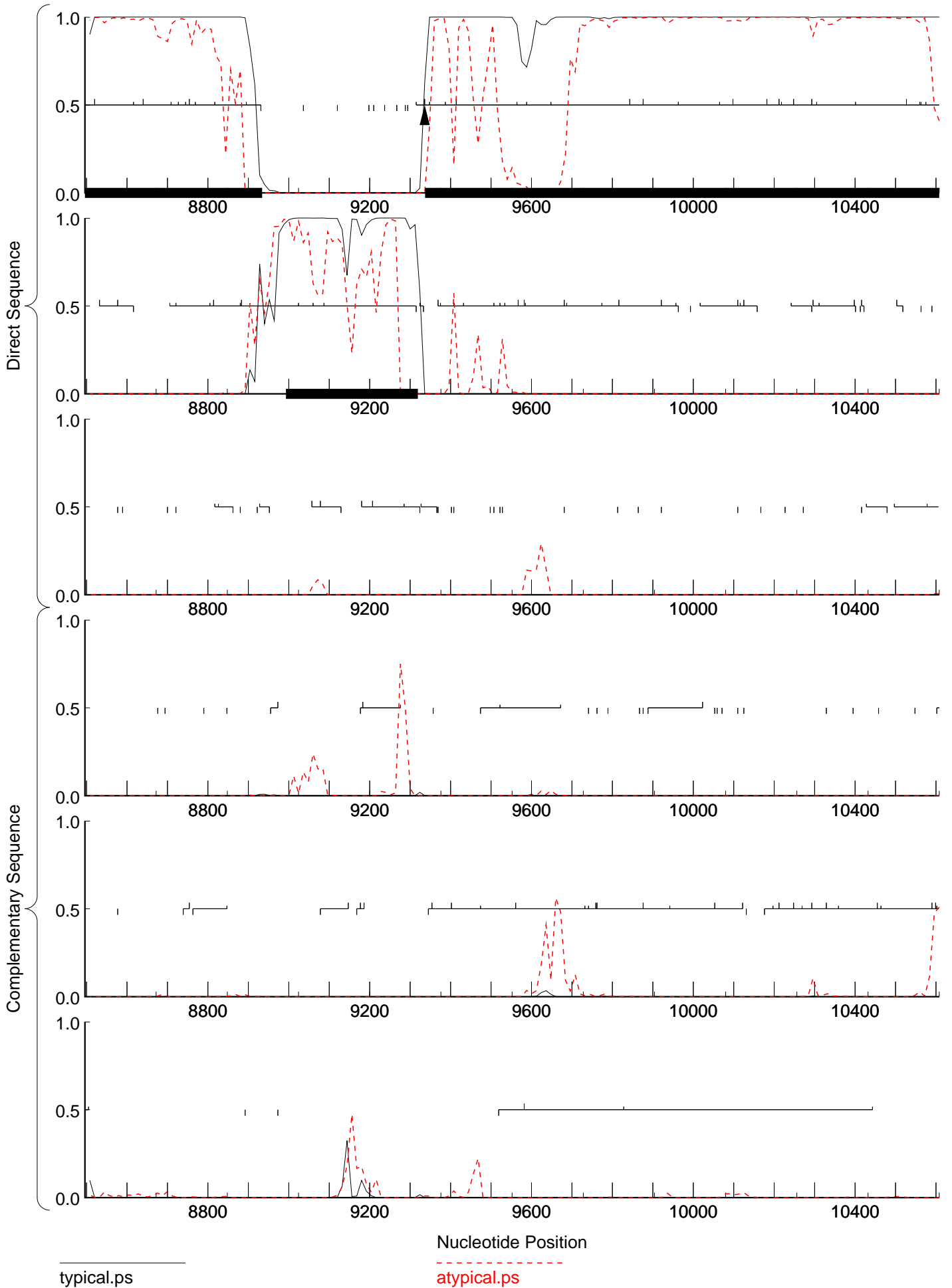
For help or information, send electronic mail to: [custserv@genepro.com](mailto:custserv@genepro.com), or [mark@amber.biology.gatech.edu](mailto:mark@amber.biology.gatech.edu)





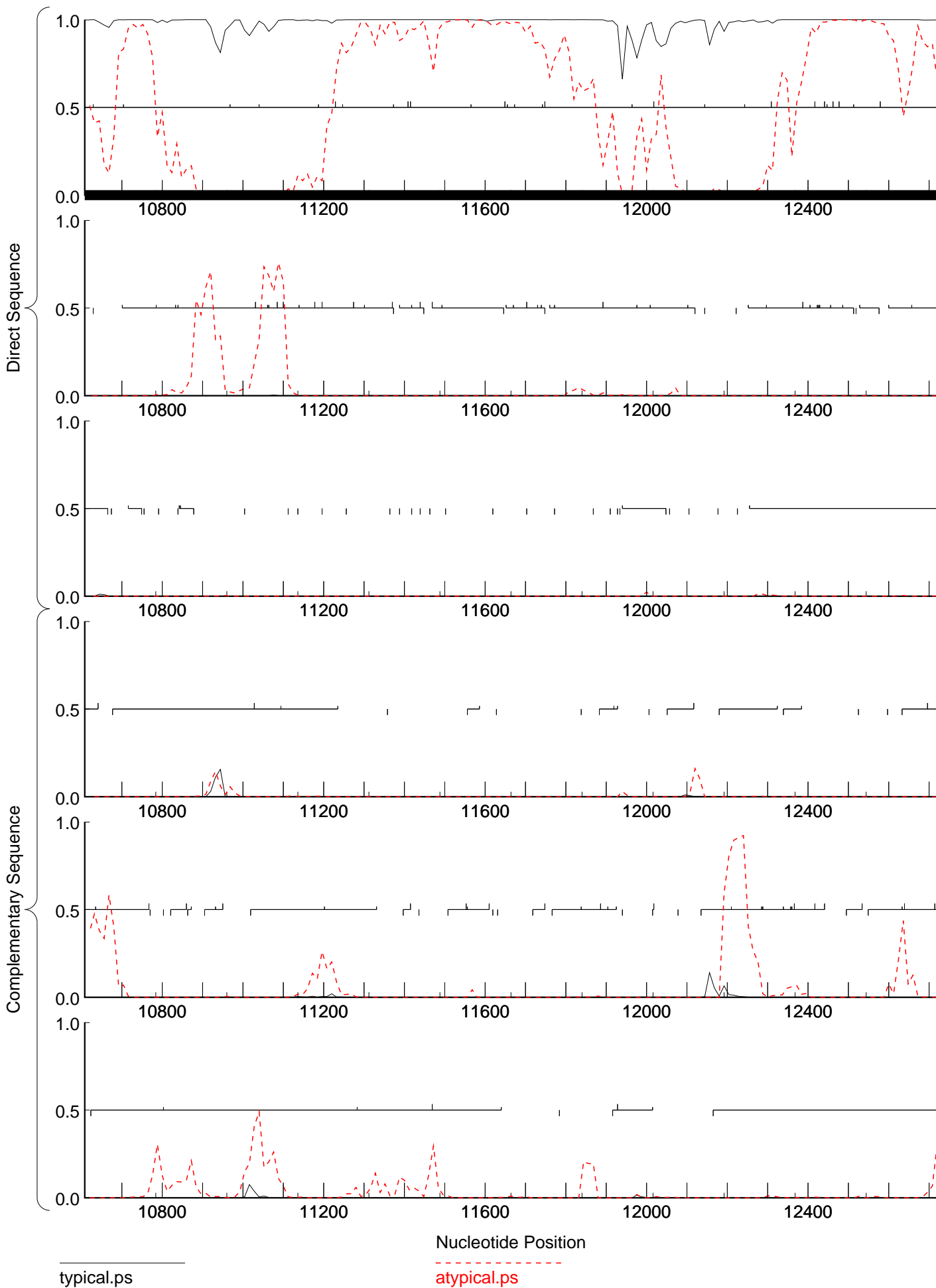






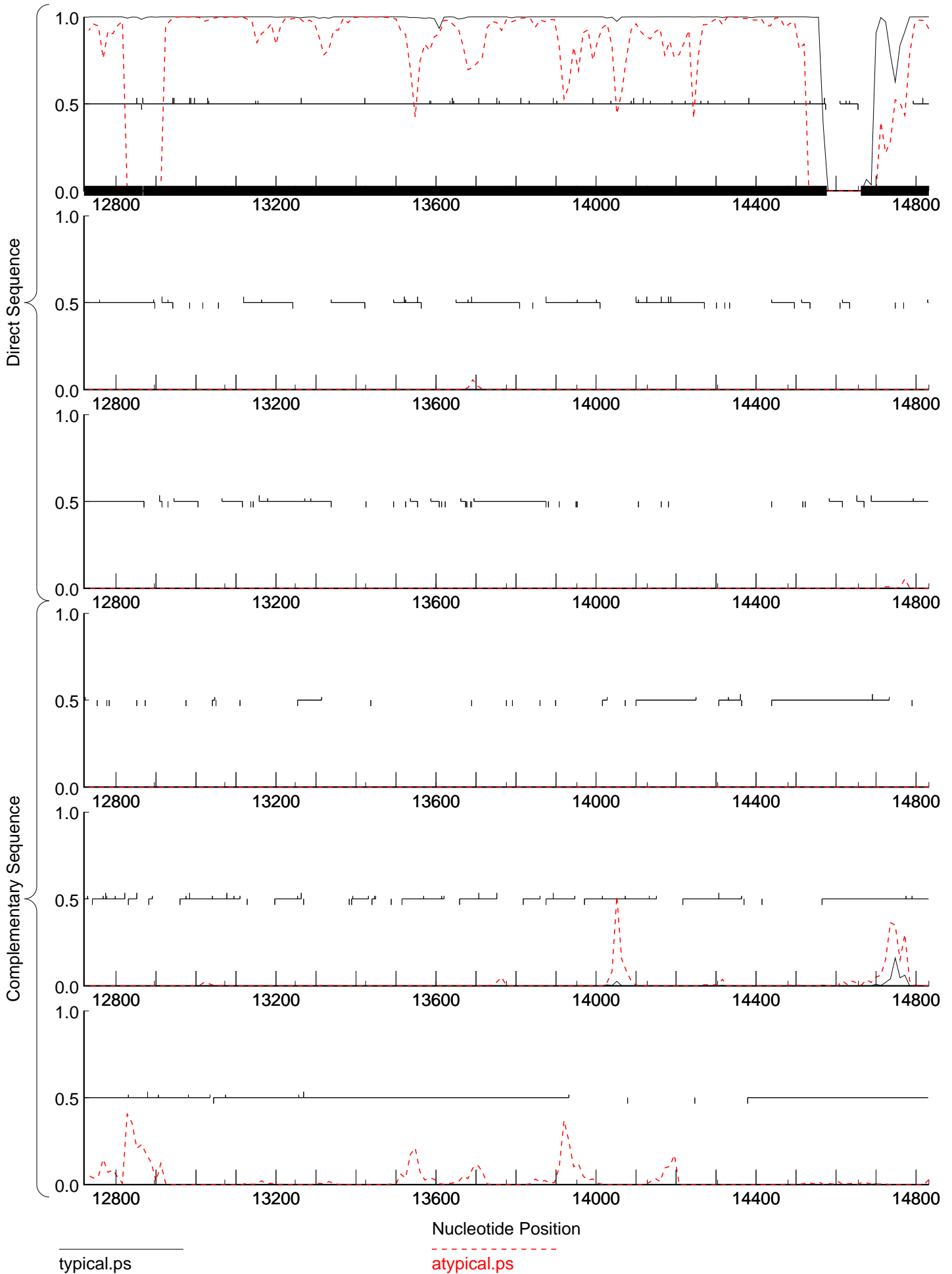
typical.ps

atypical.ps



typical.ps

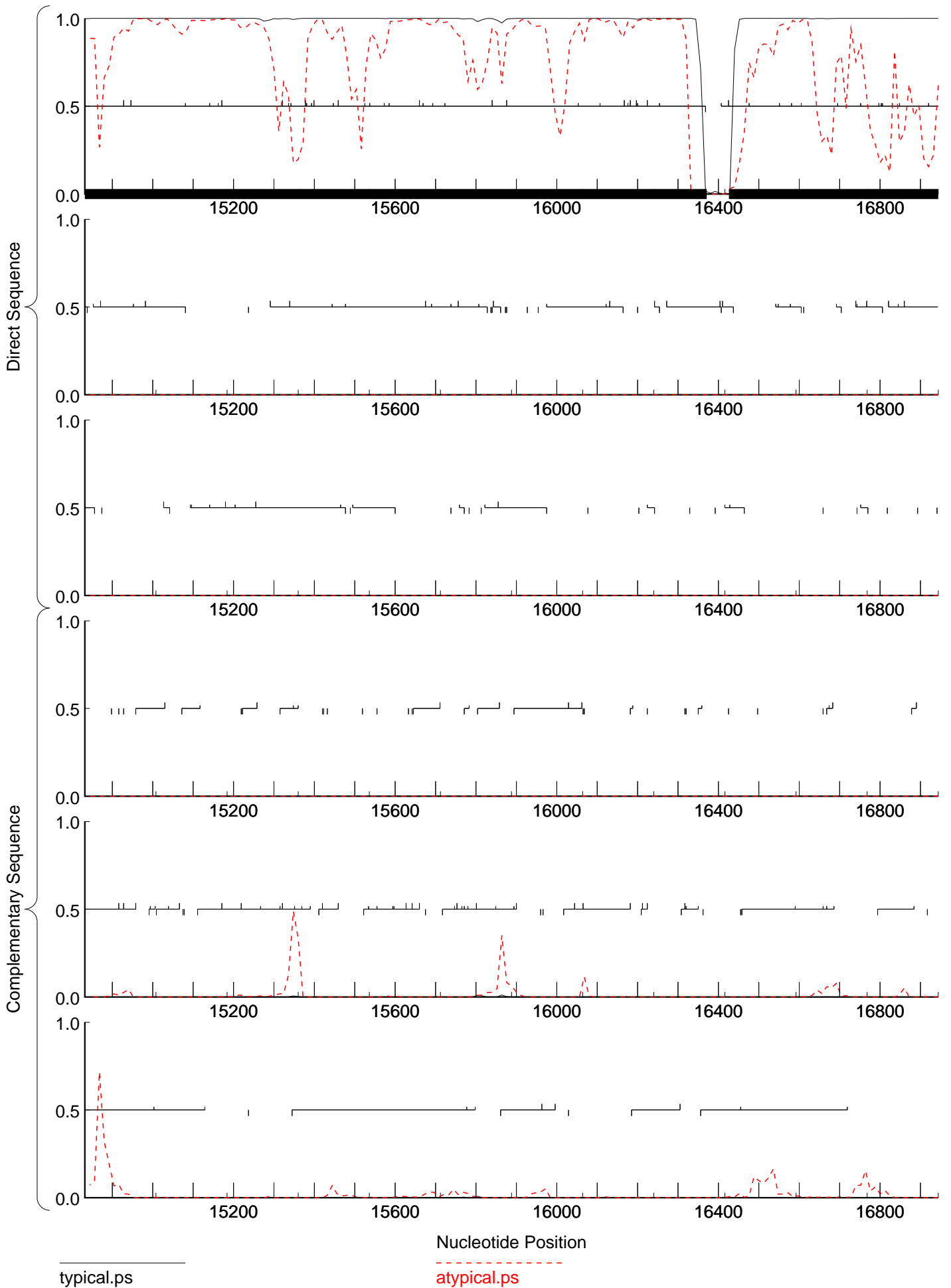
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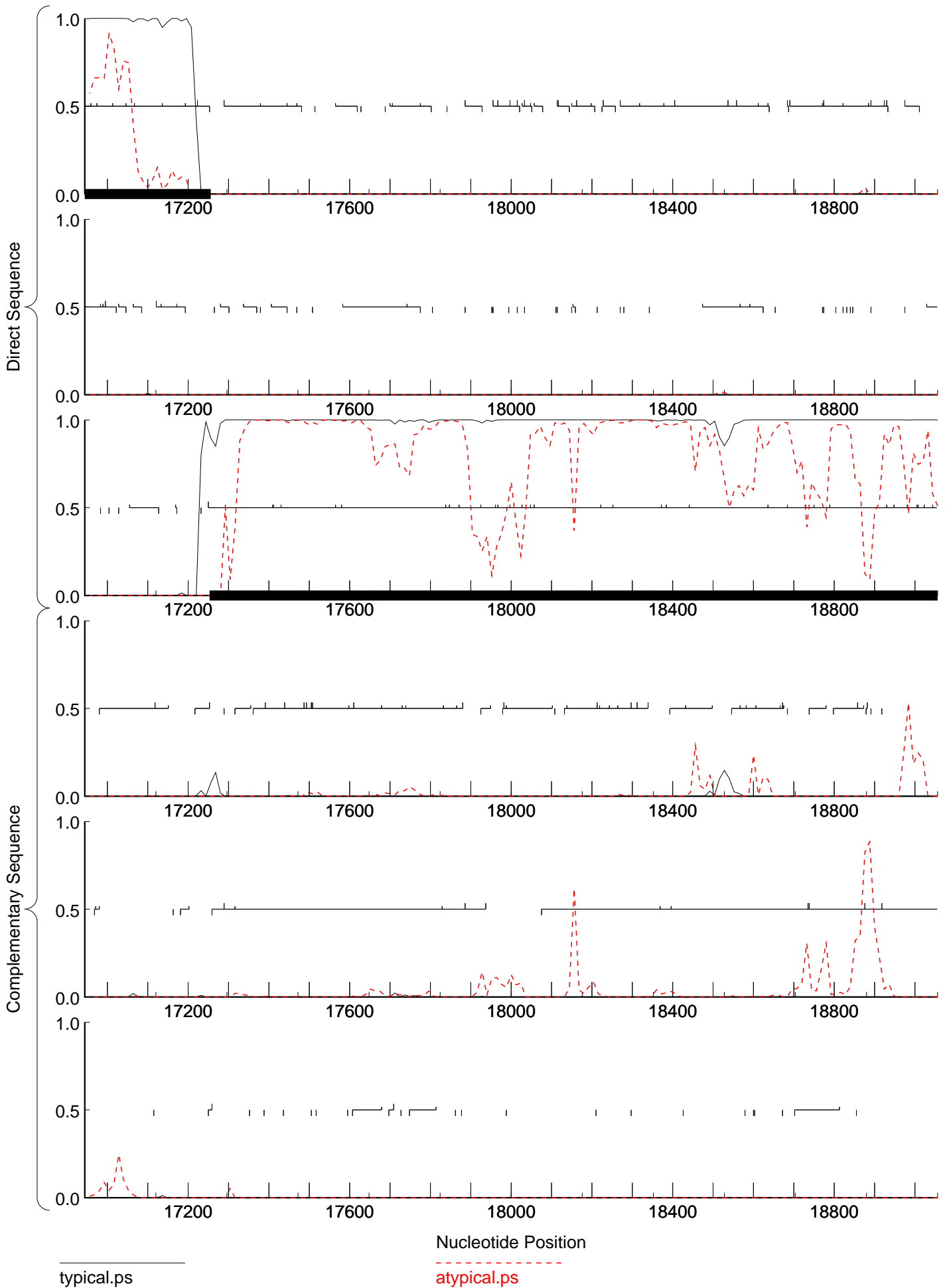


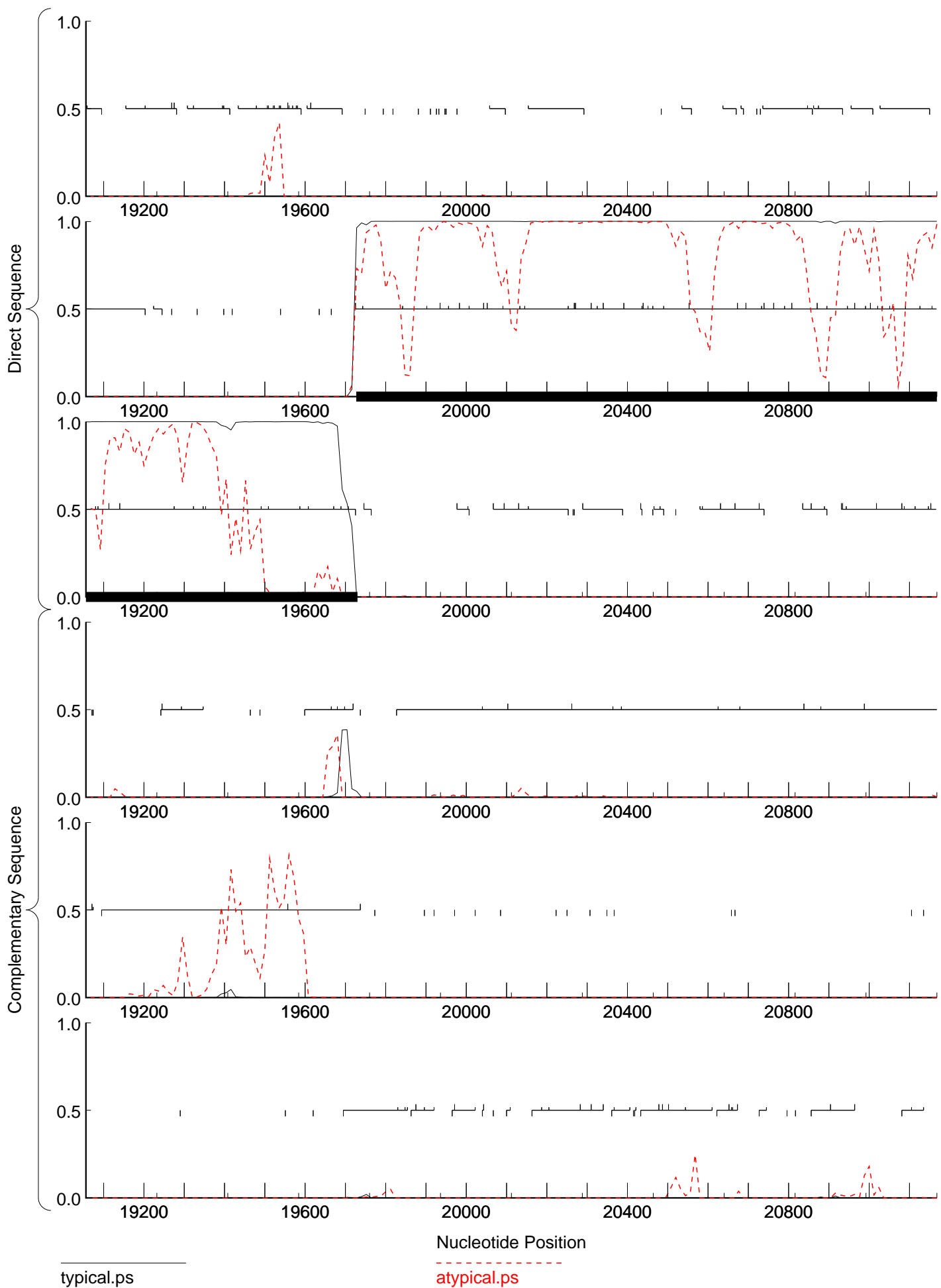
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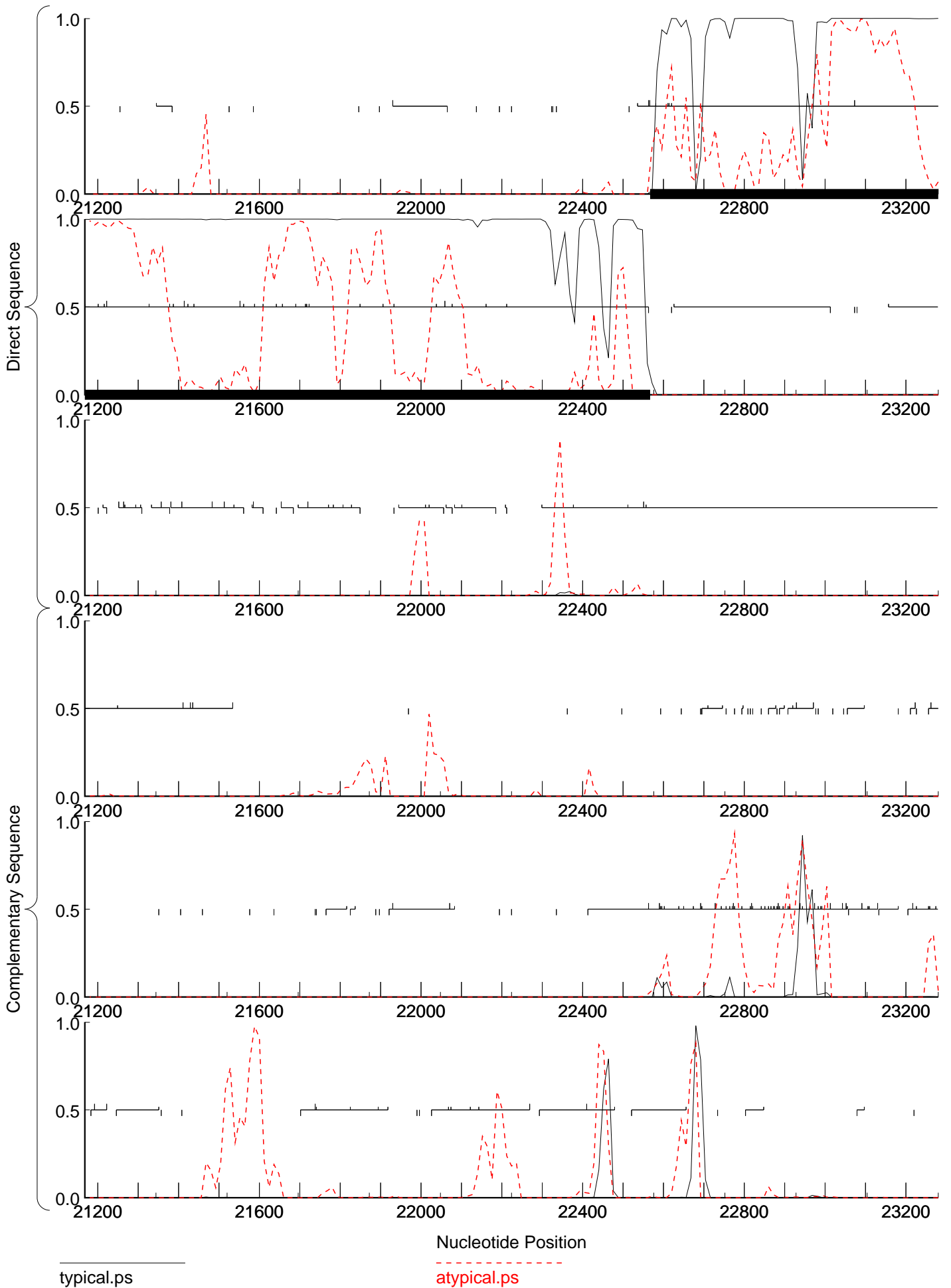


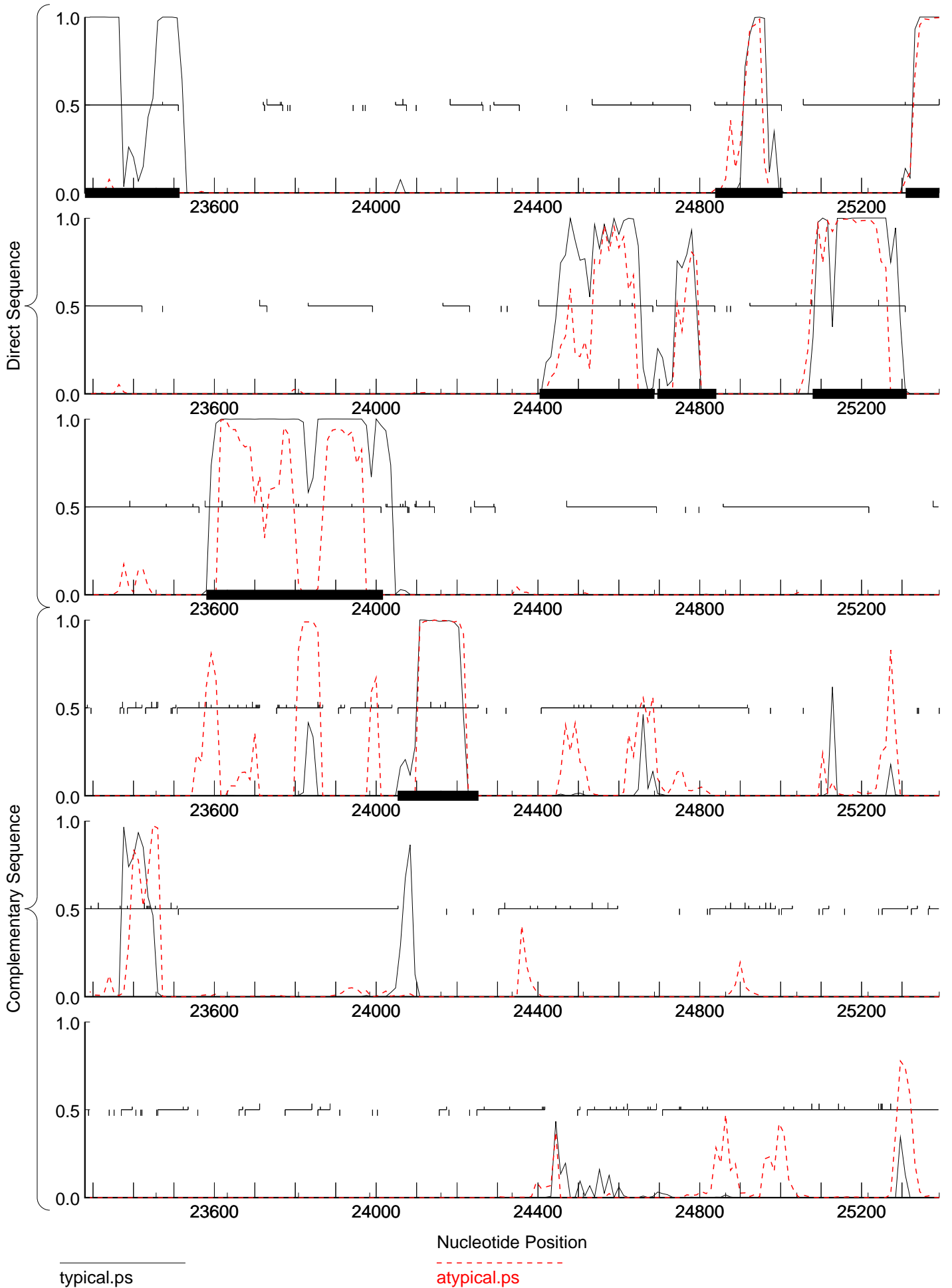


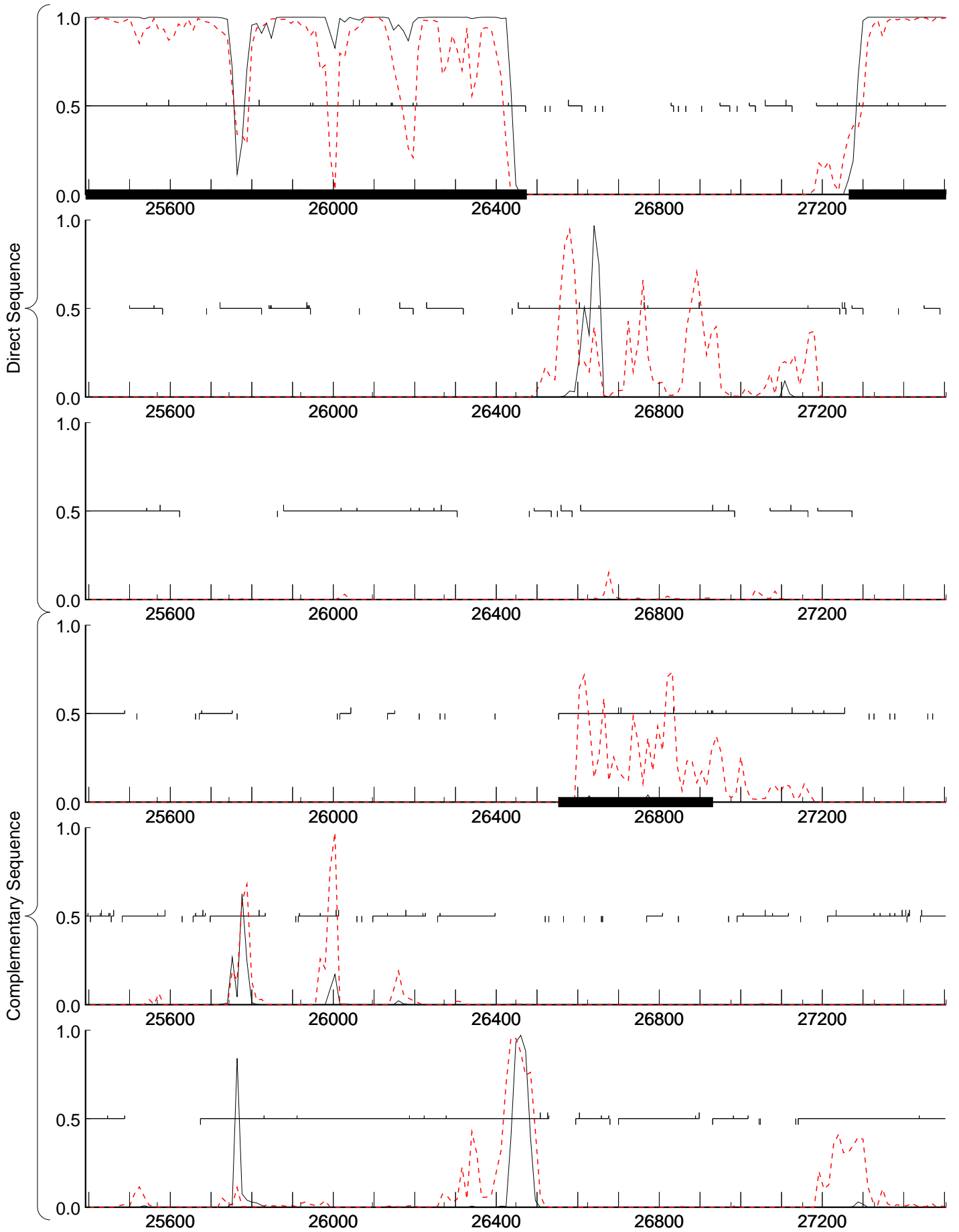


typical.ps

atypical.ps

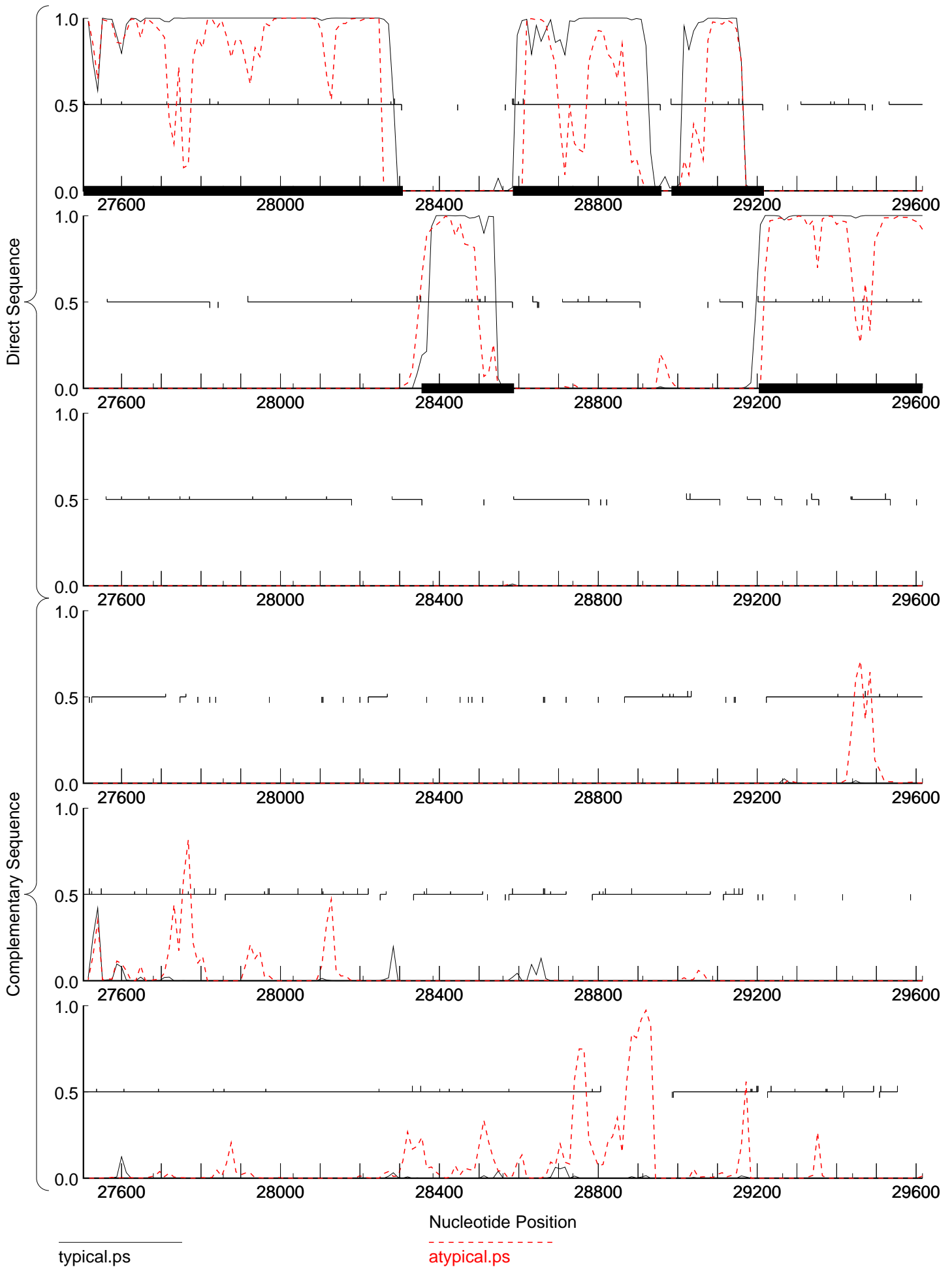


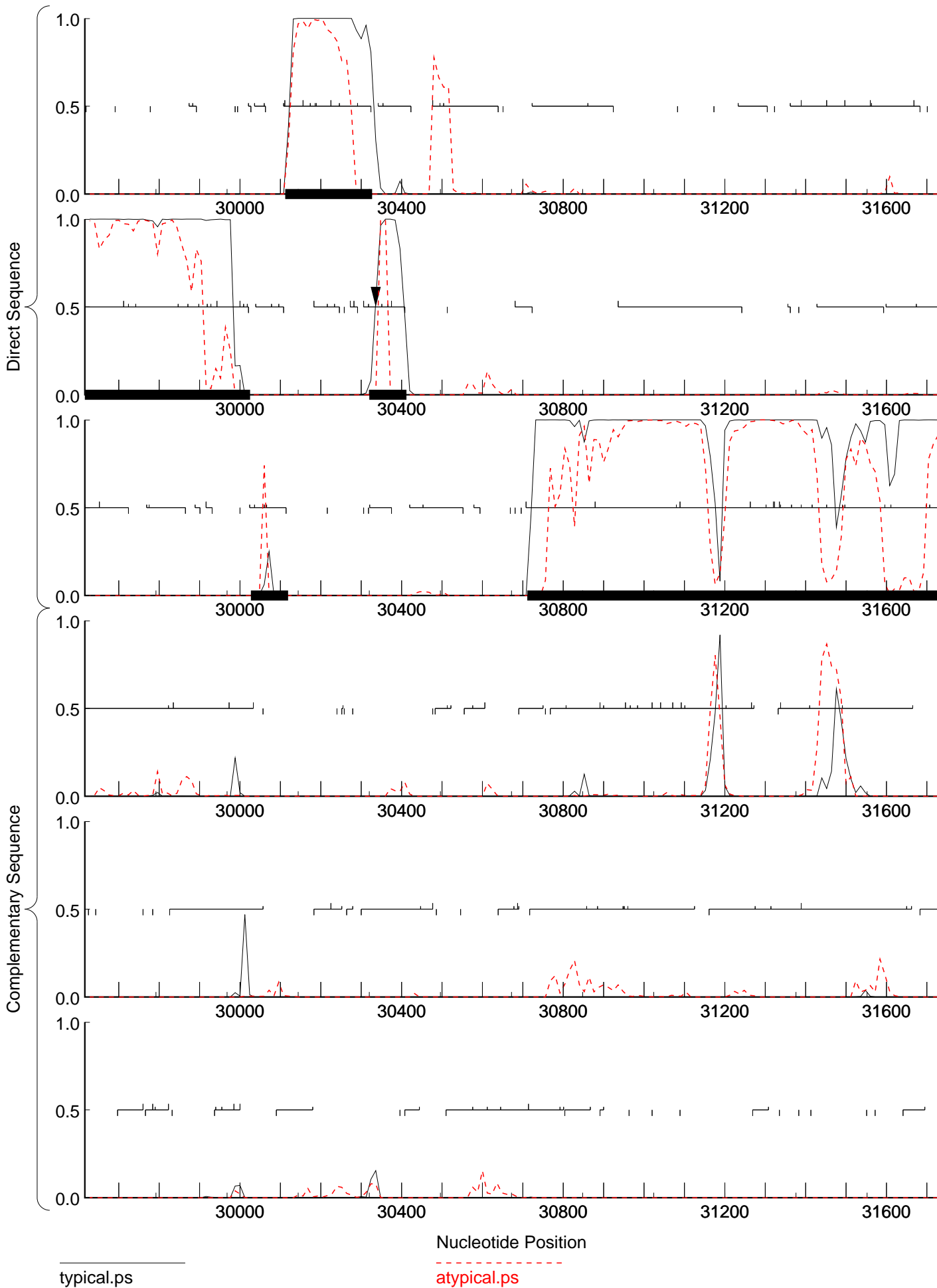




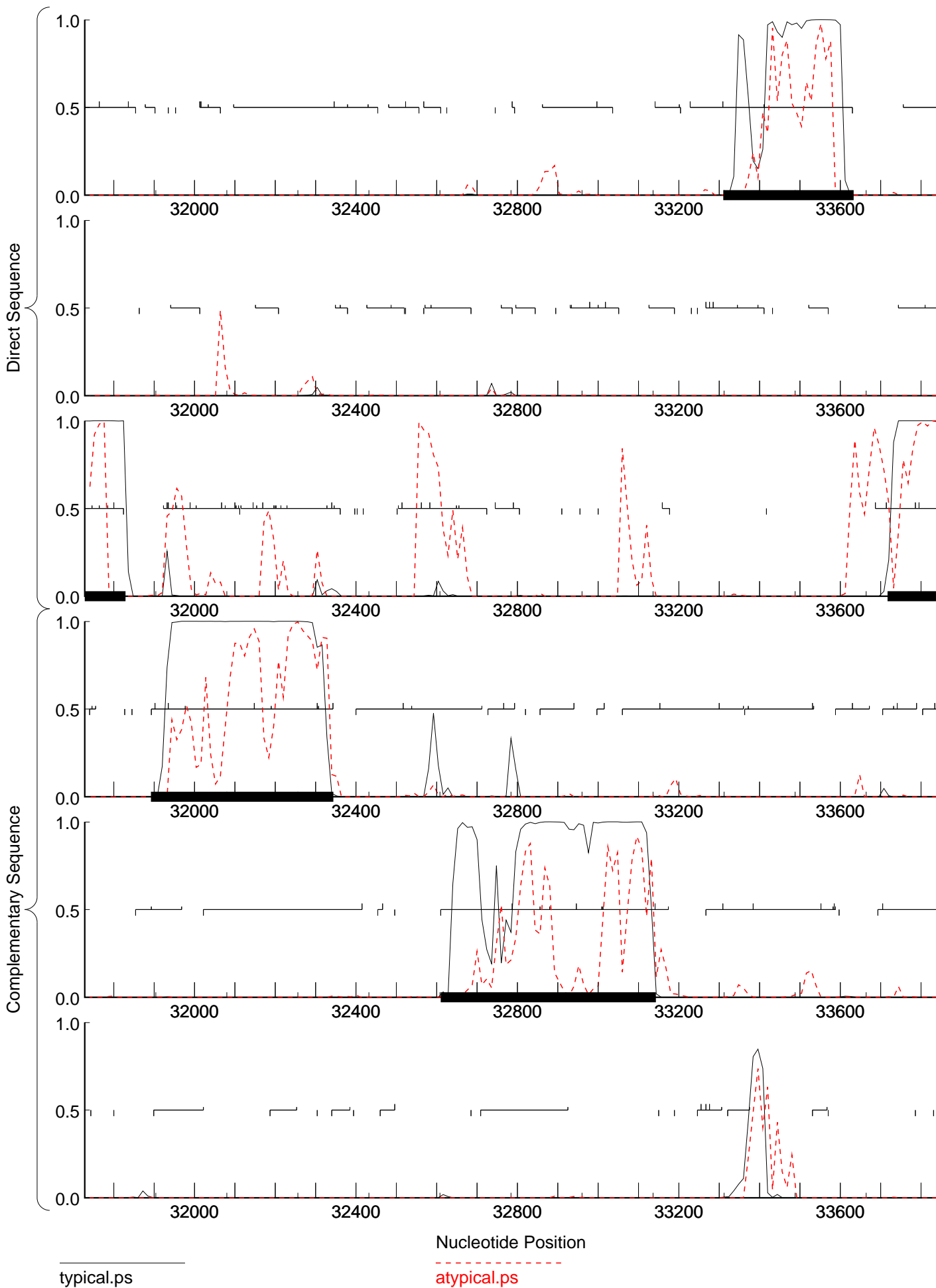
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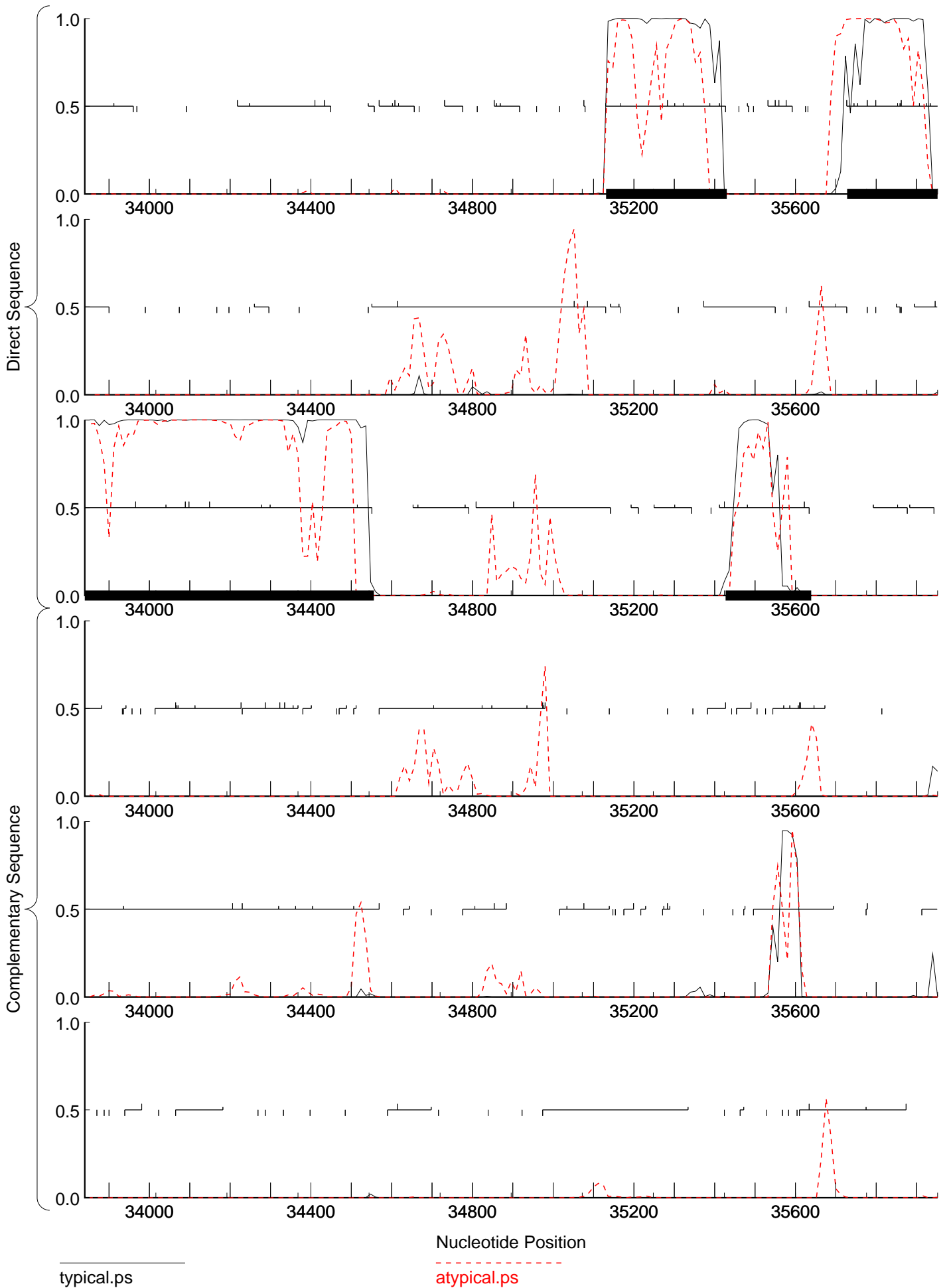
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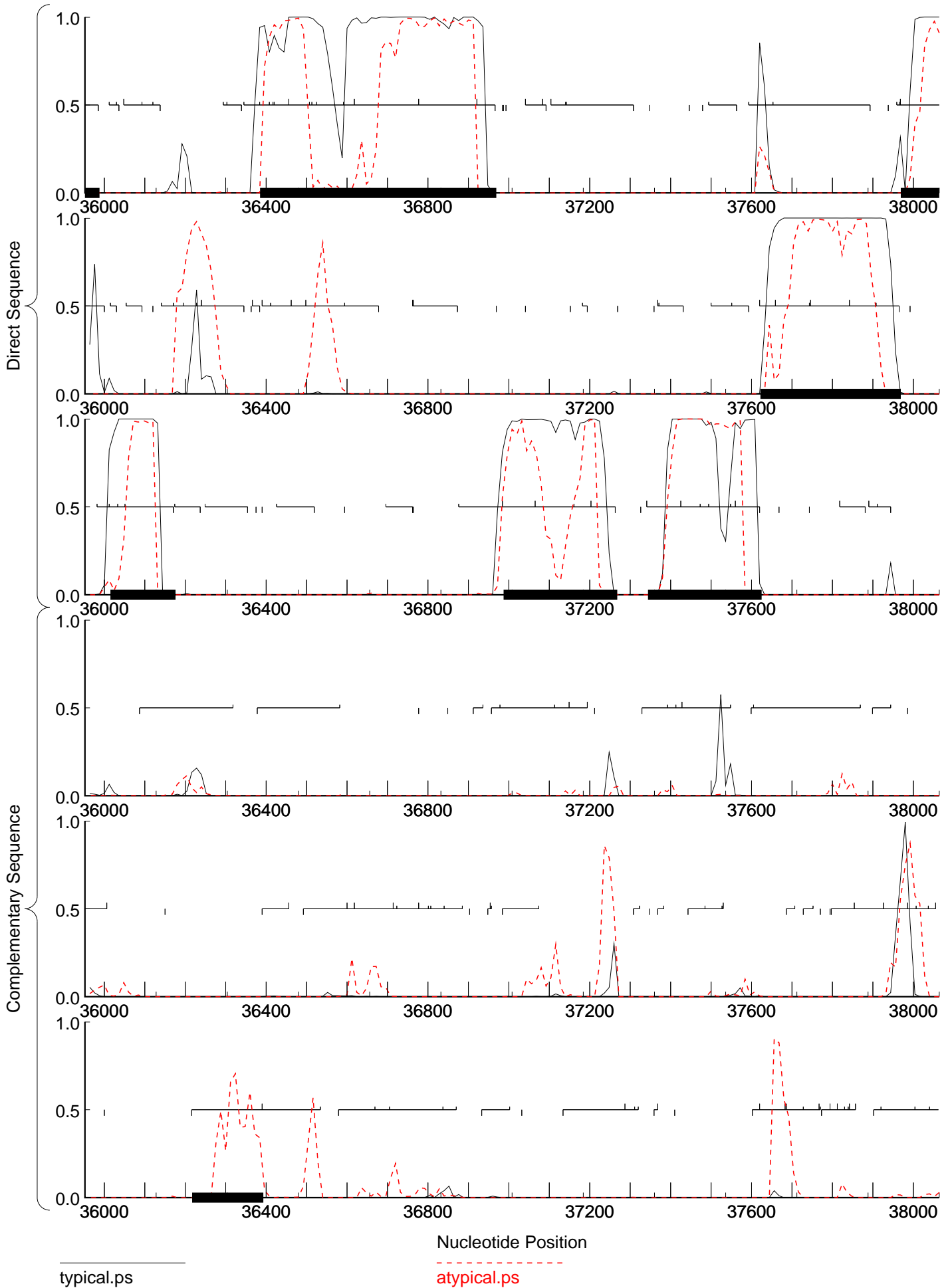






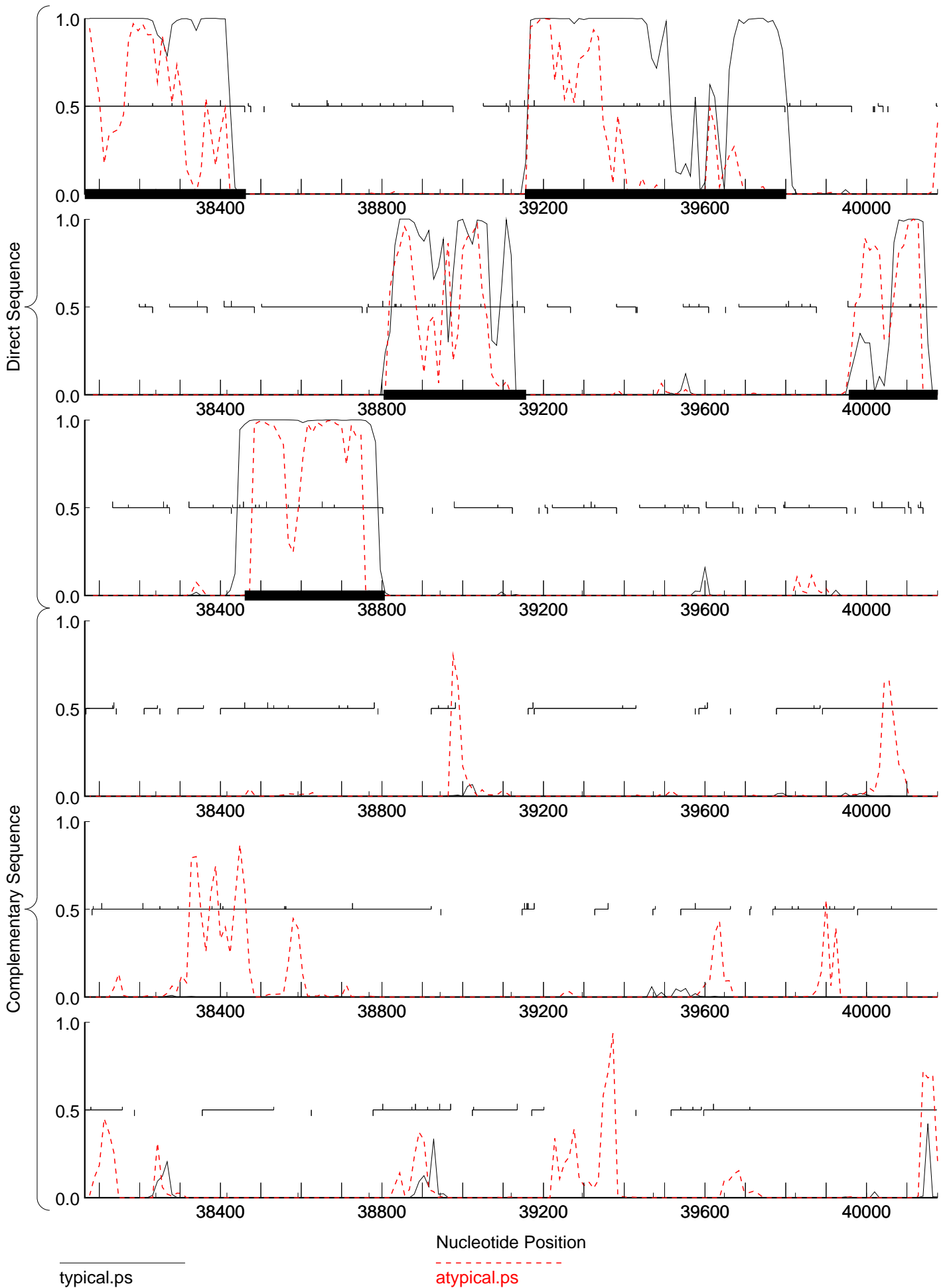


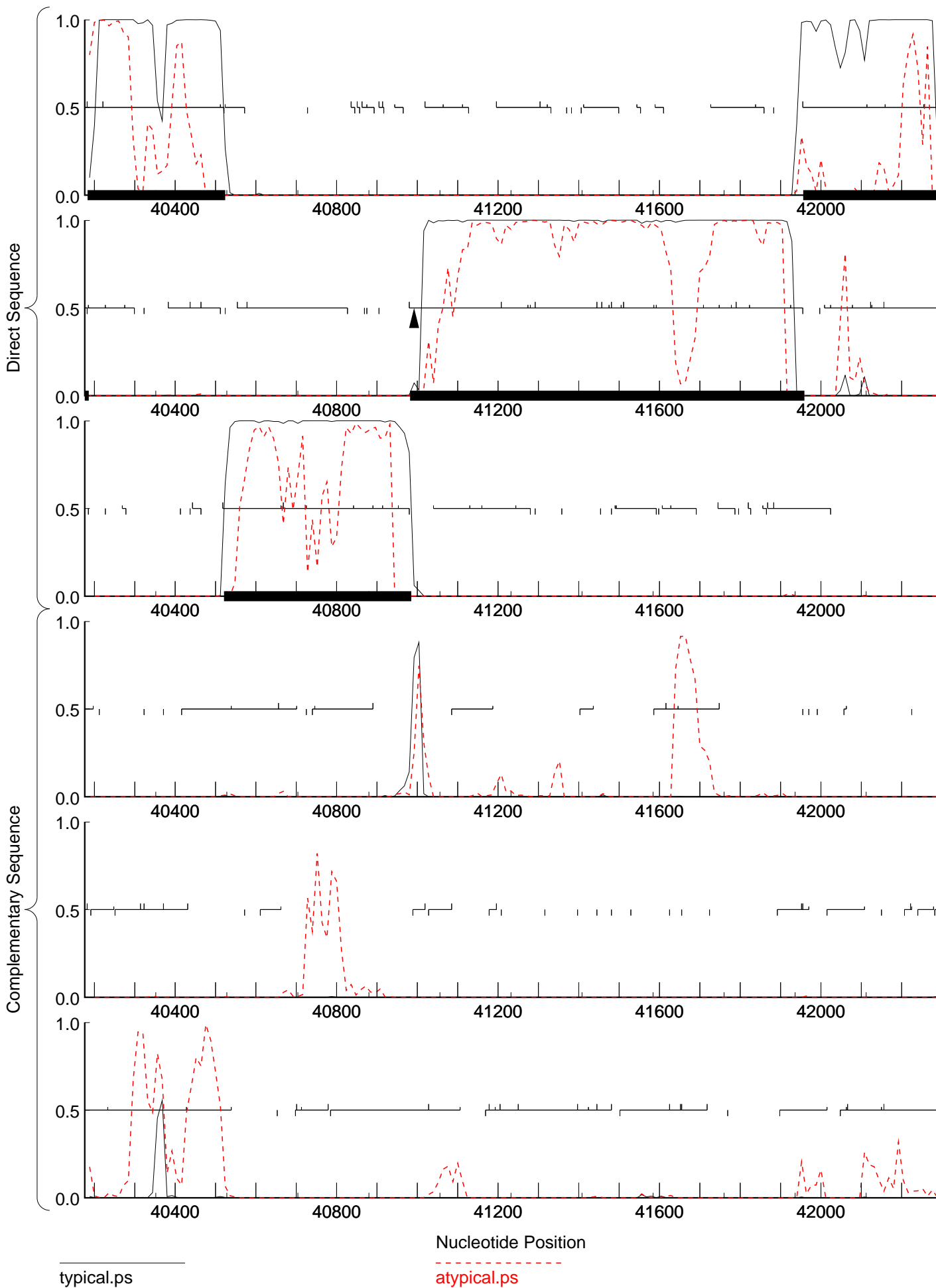


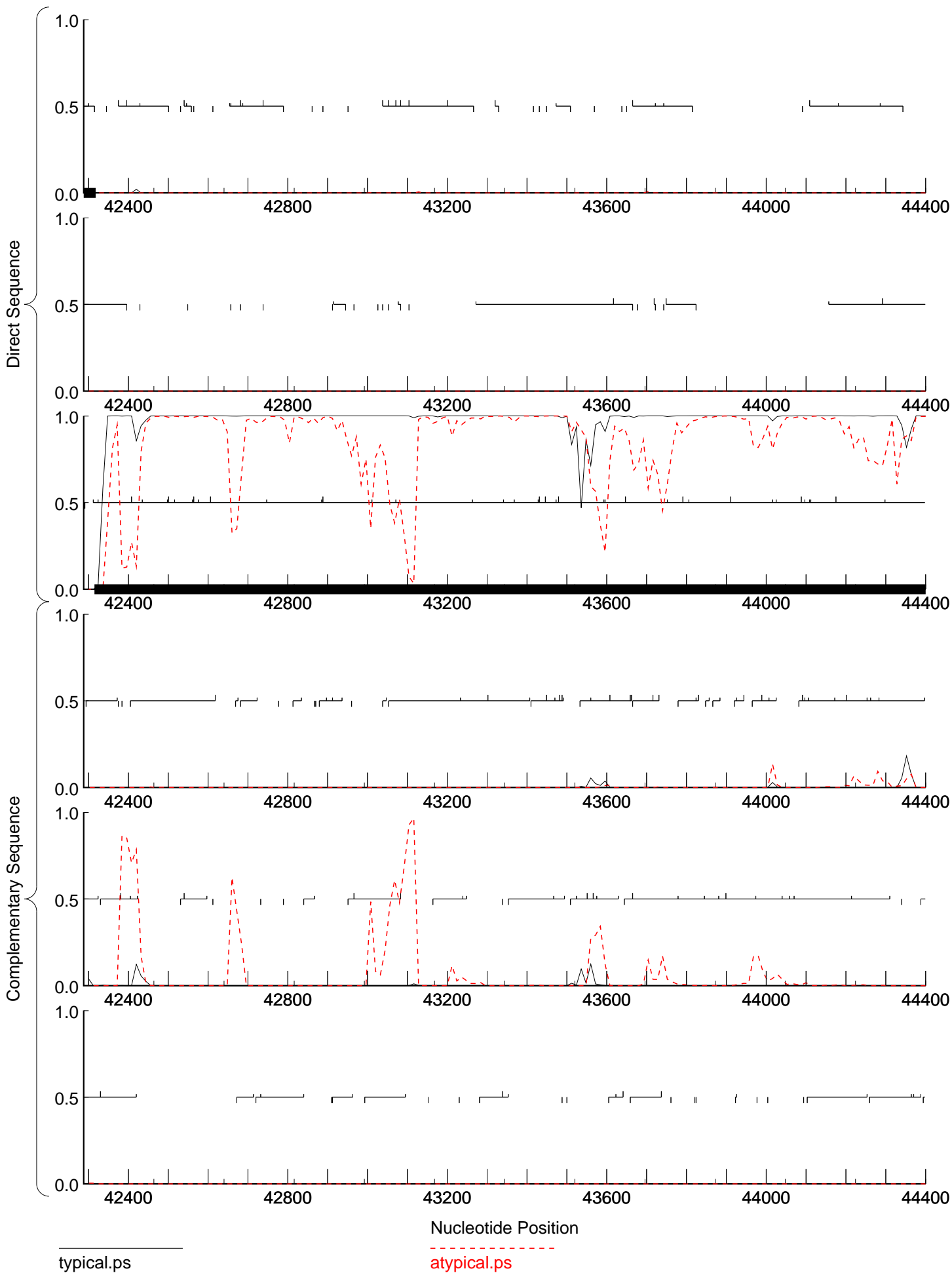


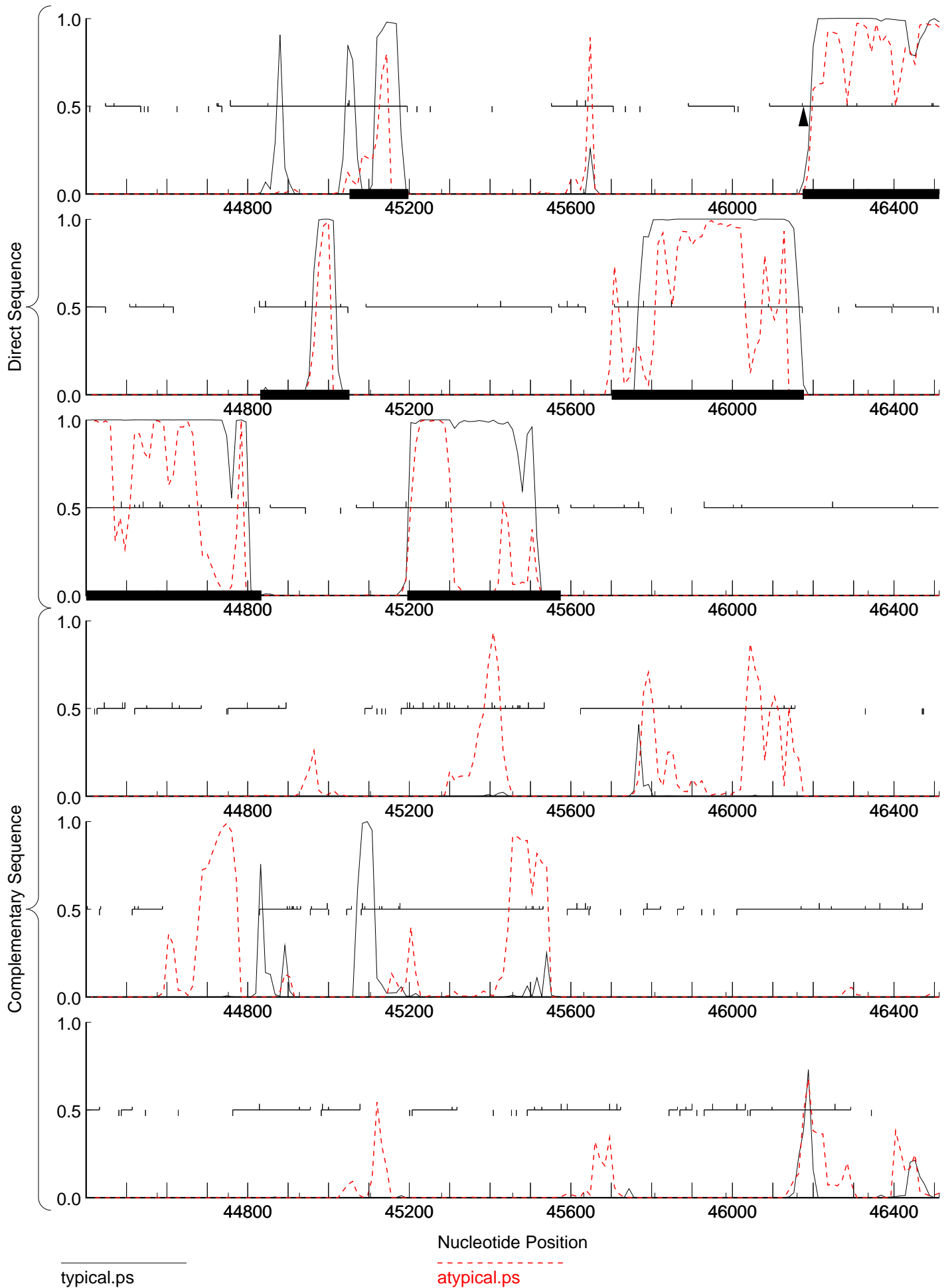
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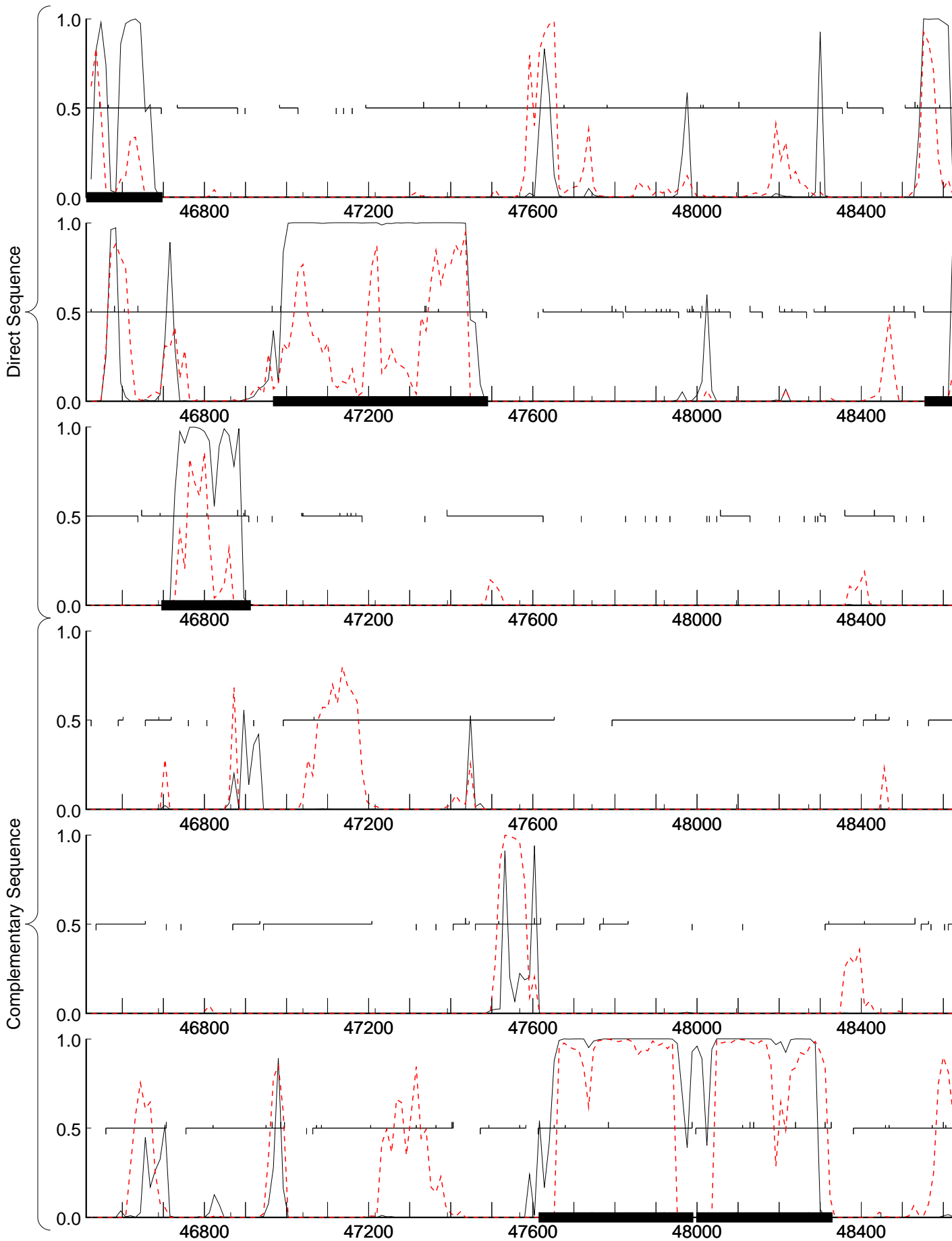
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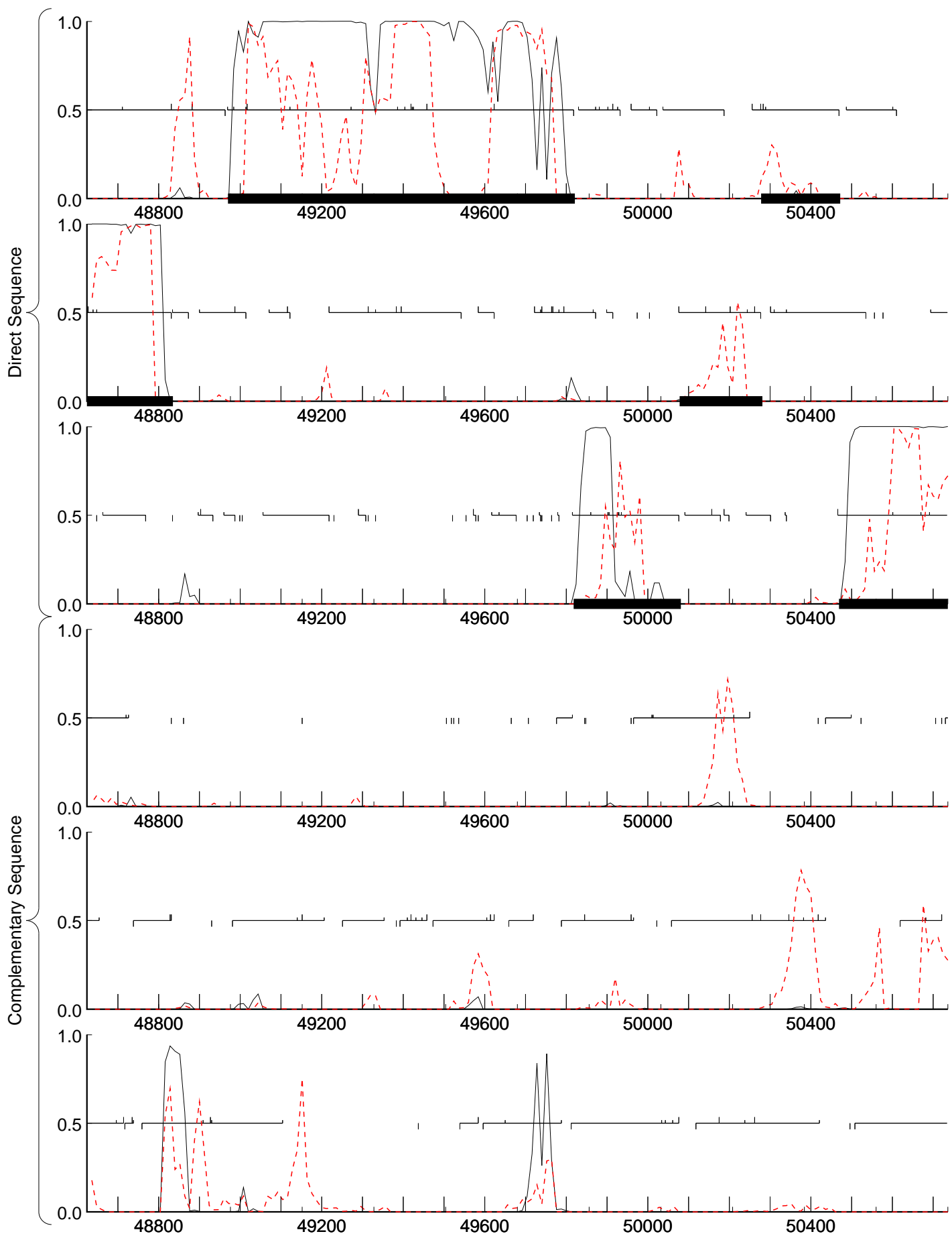




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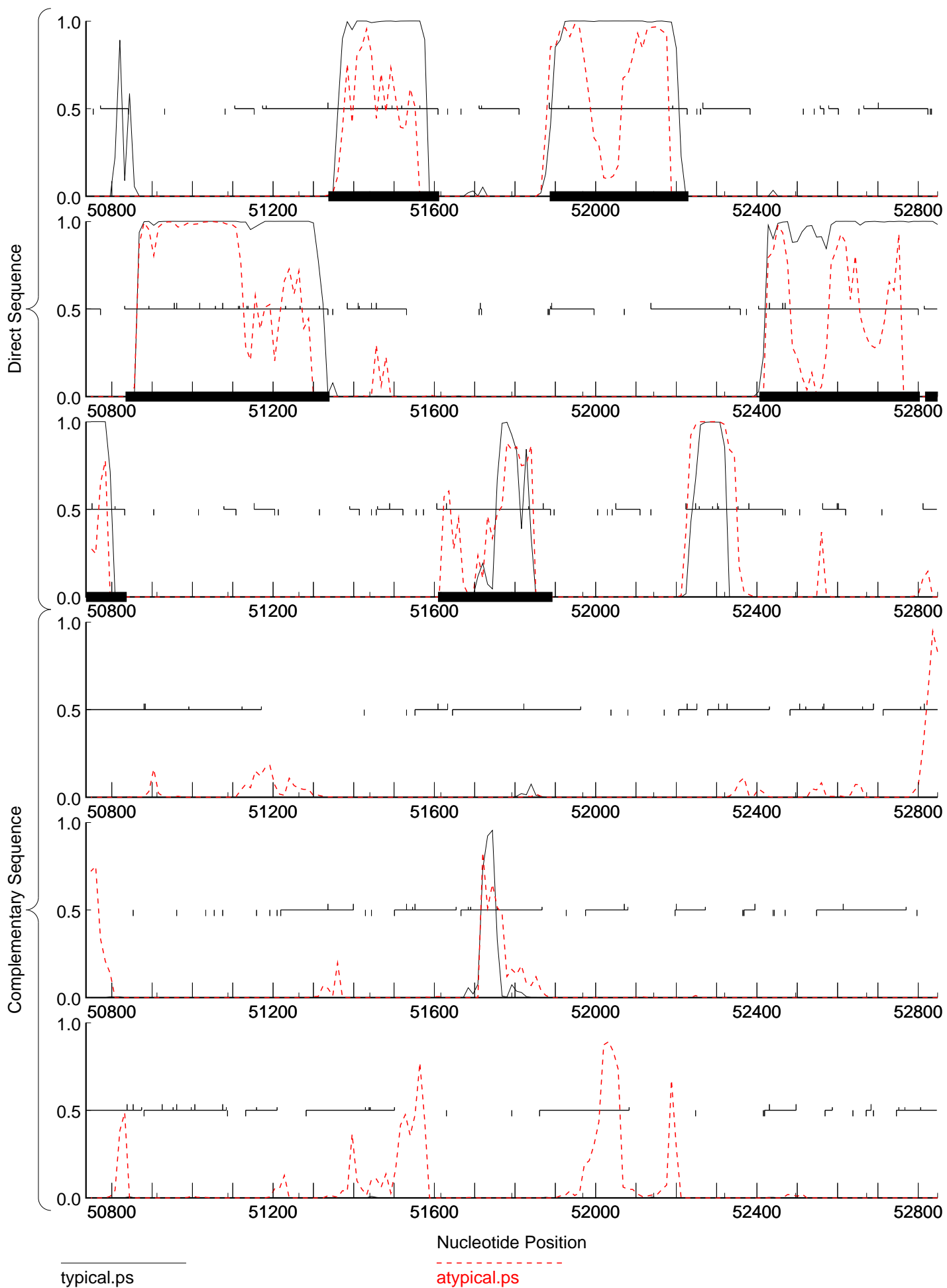
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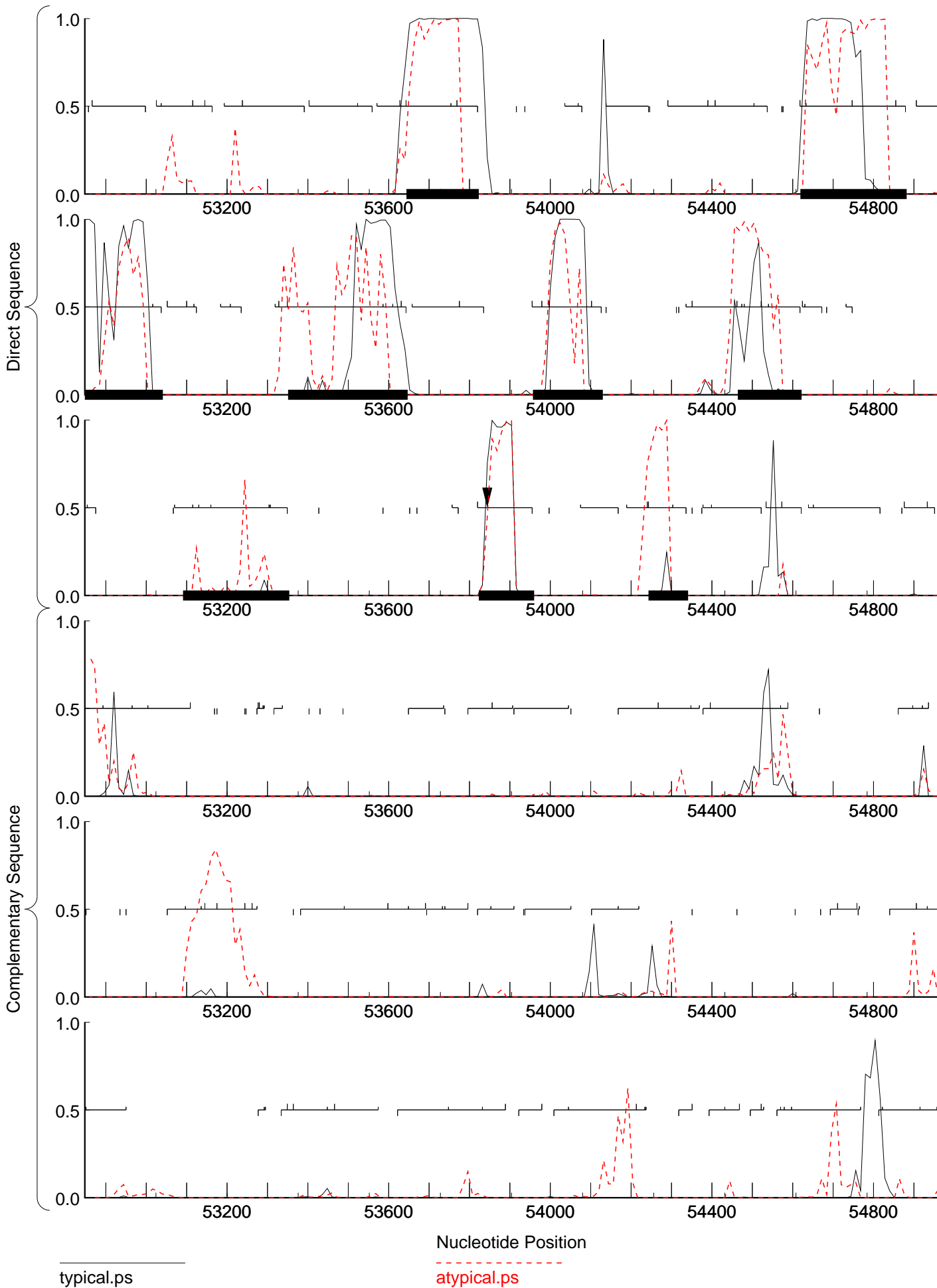


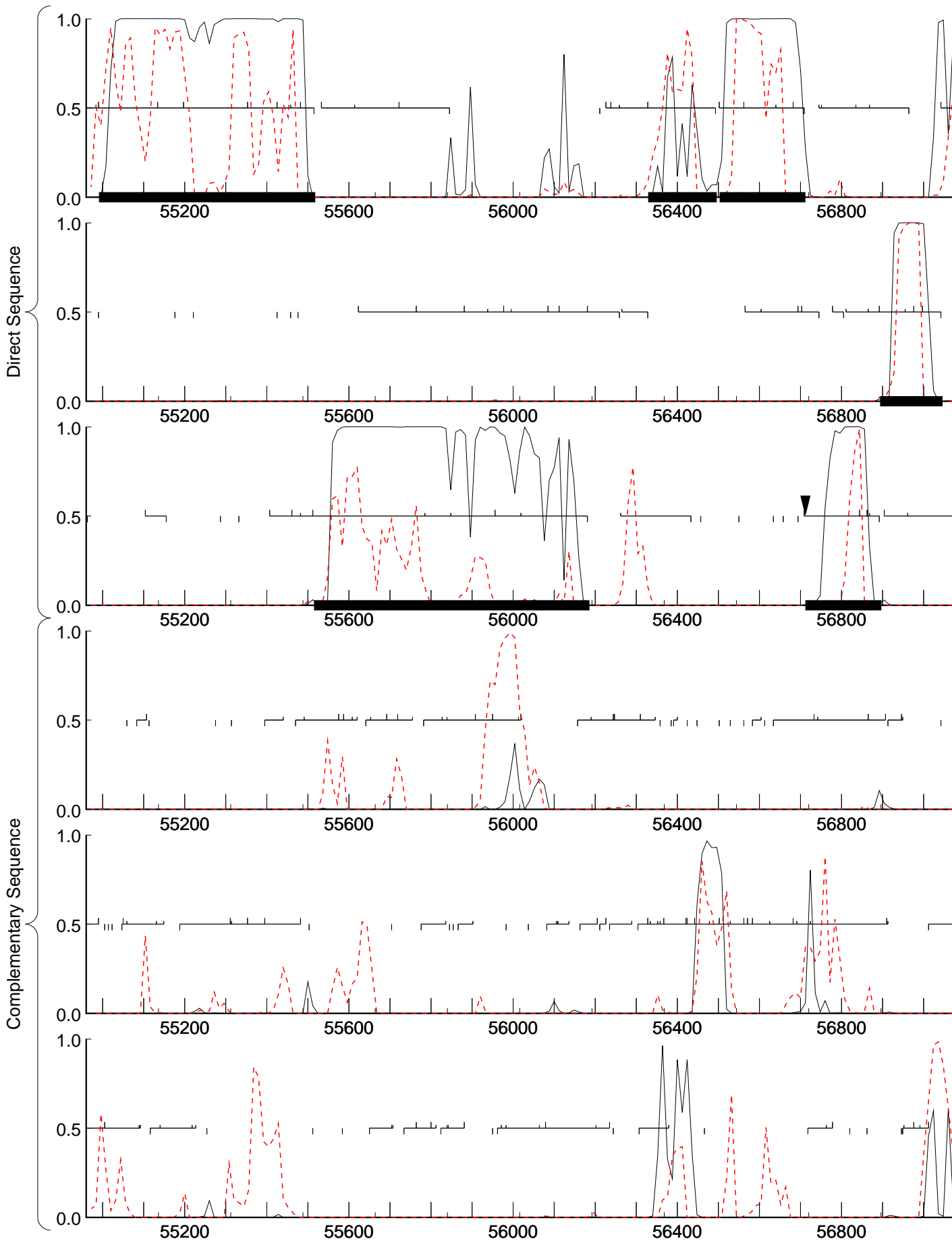


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atypical.ps

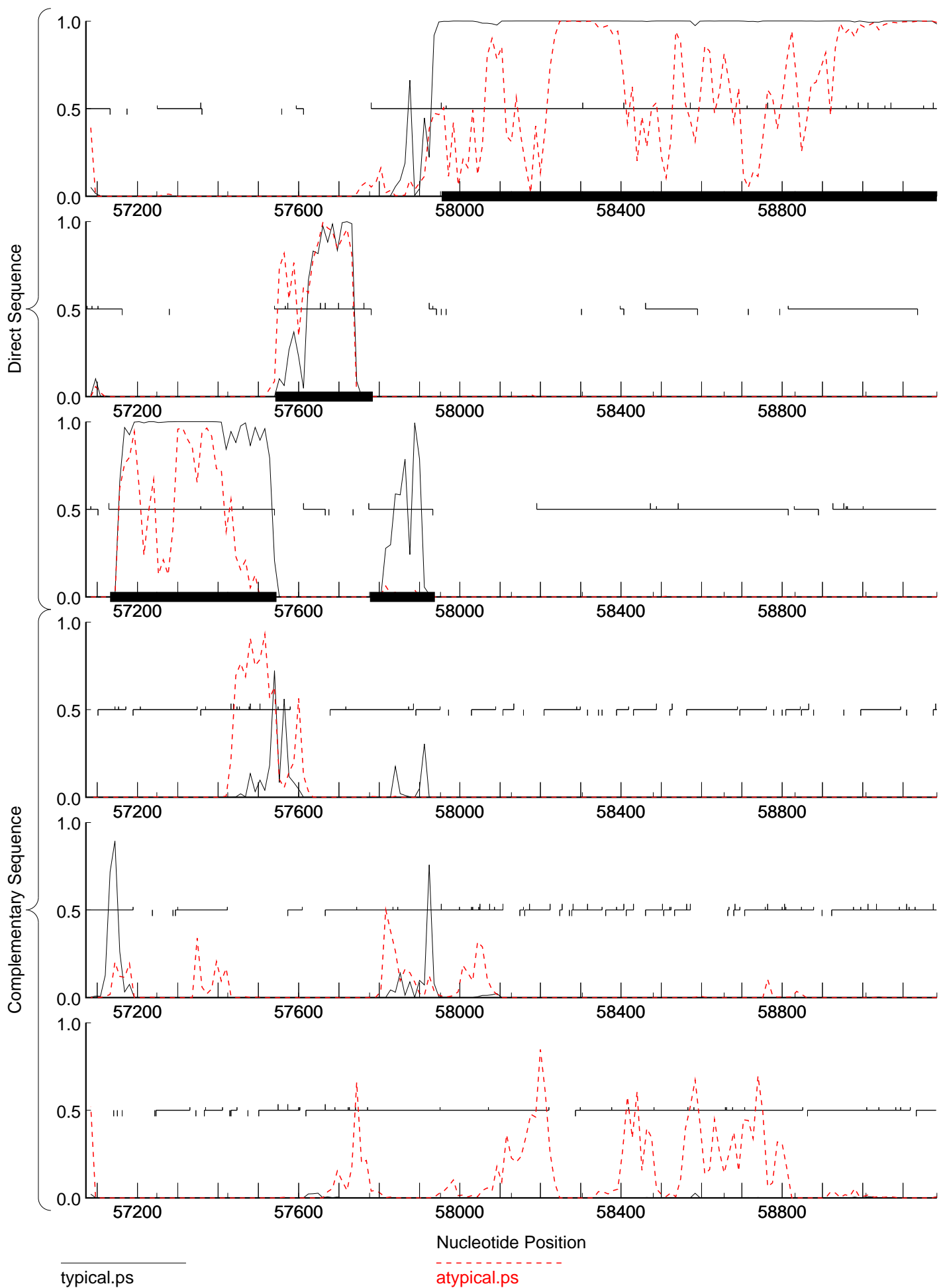






typical.ps

atypical.ps



typical.ps

atypical.ps

