

# GeneMark

Version 2.5p (09.08.06)

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## PROGRAM INFORMATION

Sequence : Gordonia phage Sombrero complete sequence, 76485 bp including 201 bp terminal repeat, Clust  
Analysis Date : 6/24/18 at 18:17:22  
Pages : 38  
Sequence Length : 76485 bp  
GC Content : 59.02%

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

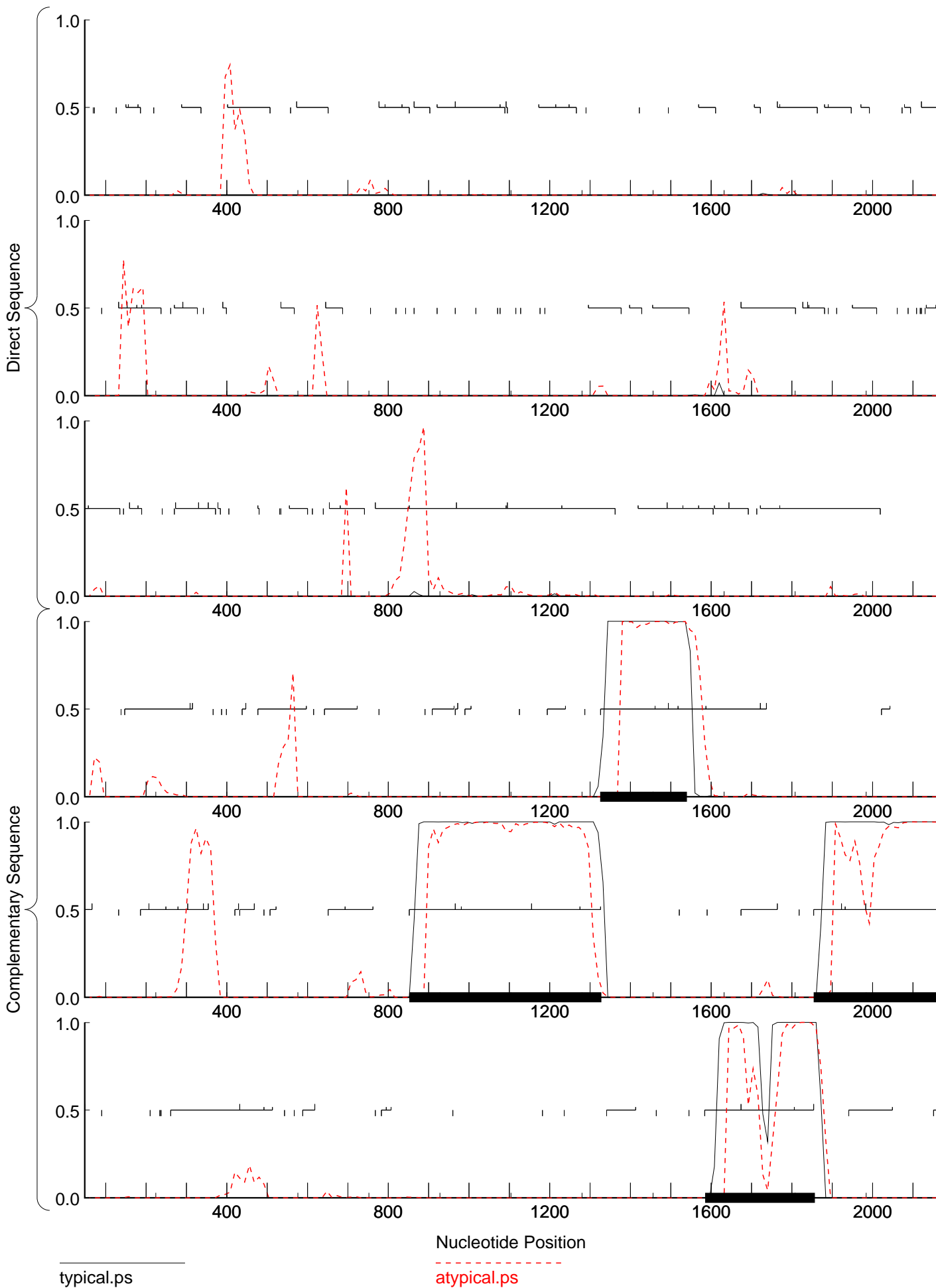
Training set derived by GeneMarkS, 4.30 October 2014  
Sun Jun 24 18:17:22 2018

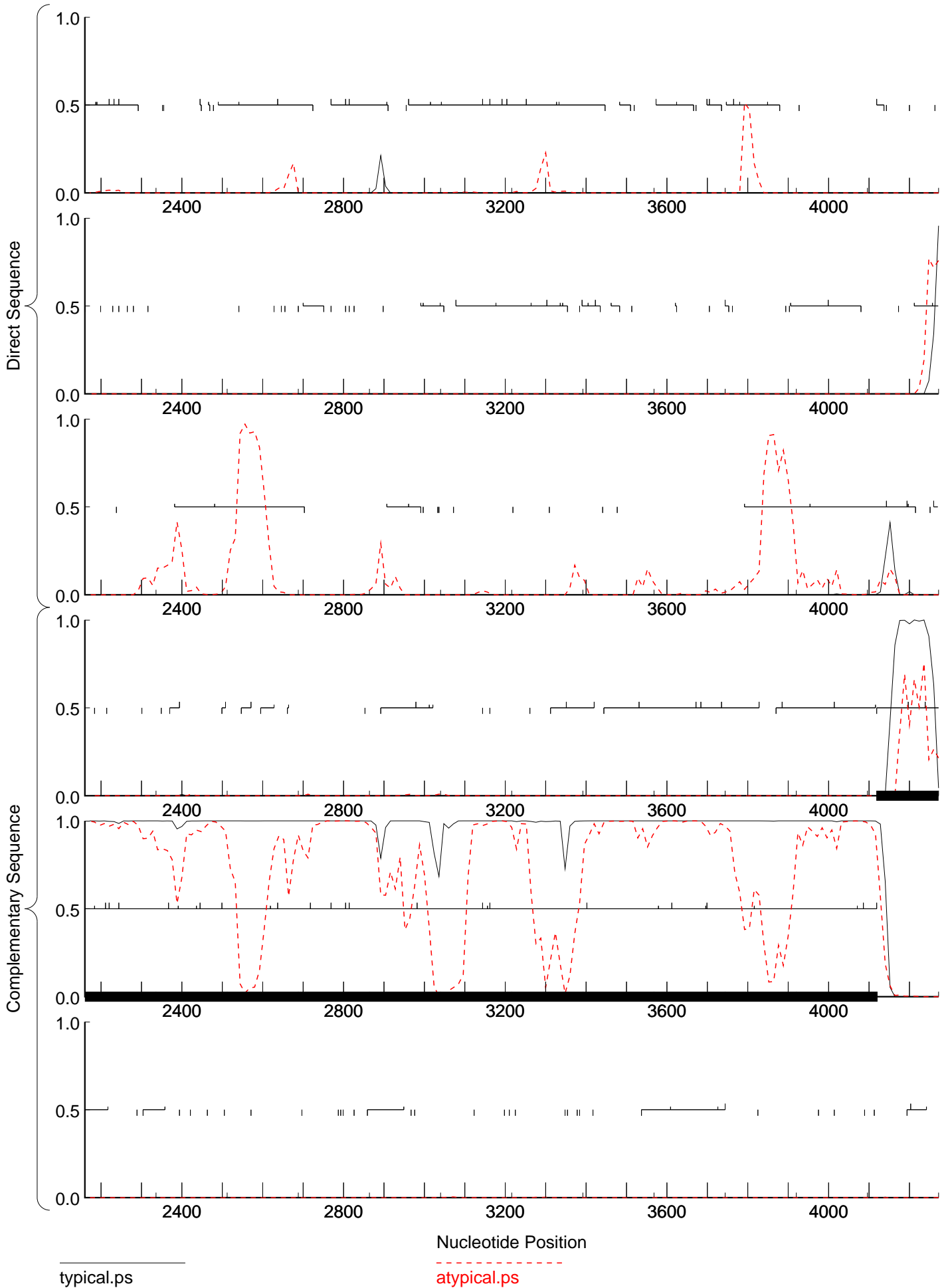
## MATRIX INFORMATION

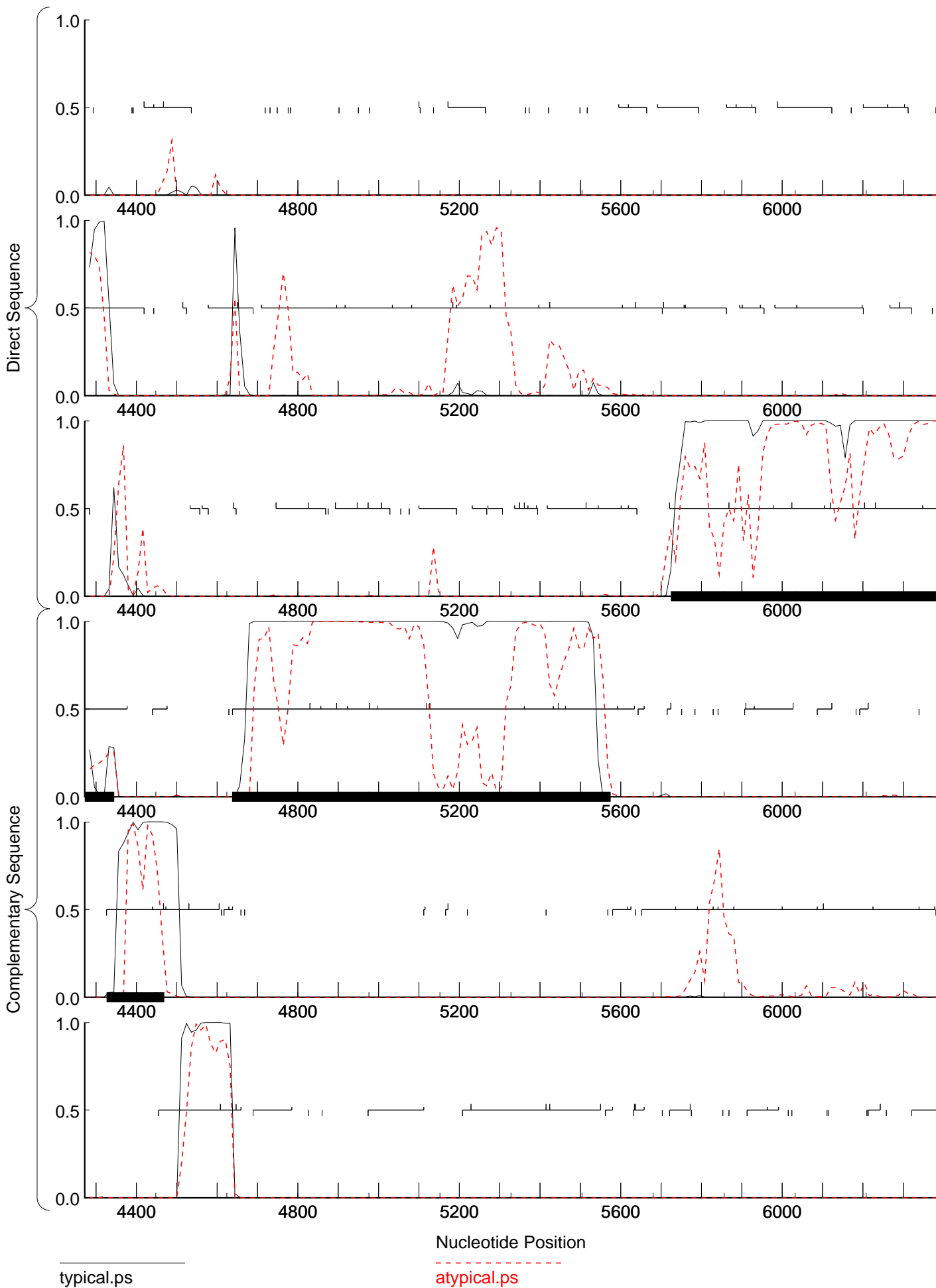
Matrix : -  
Author : -  
Order : 4

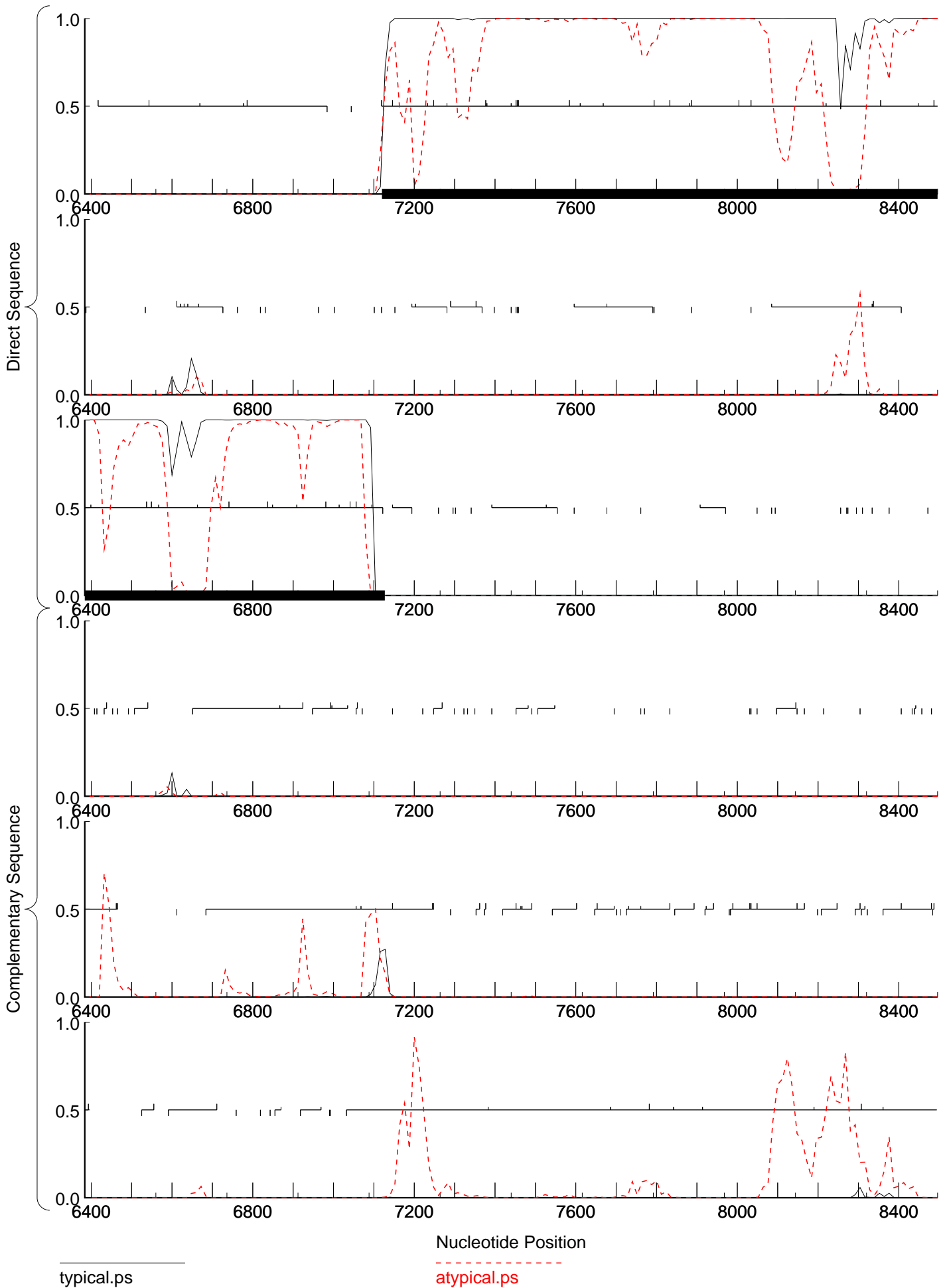
Send questions / comments to:  
Dr. M. Borodovsky  
Georgia Institute of Technology  
School of Biology  
Atlanta, GA 30332-0230

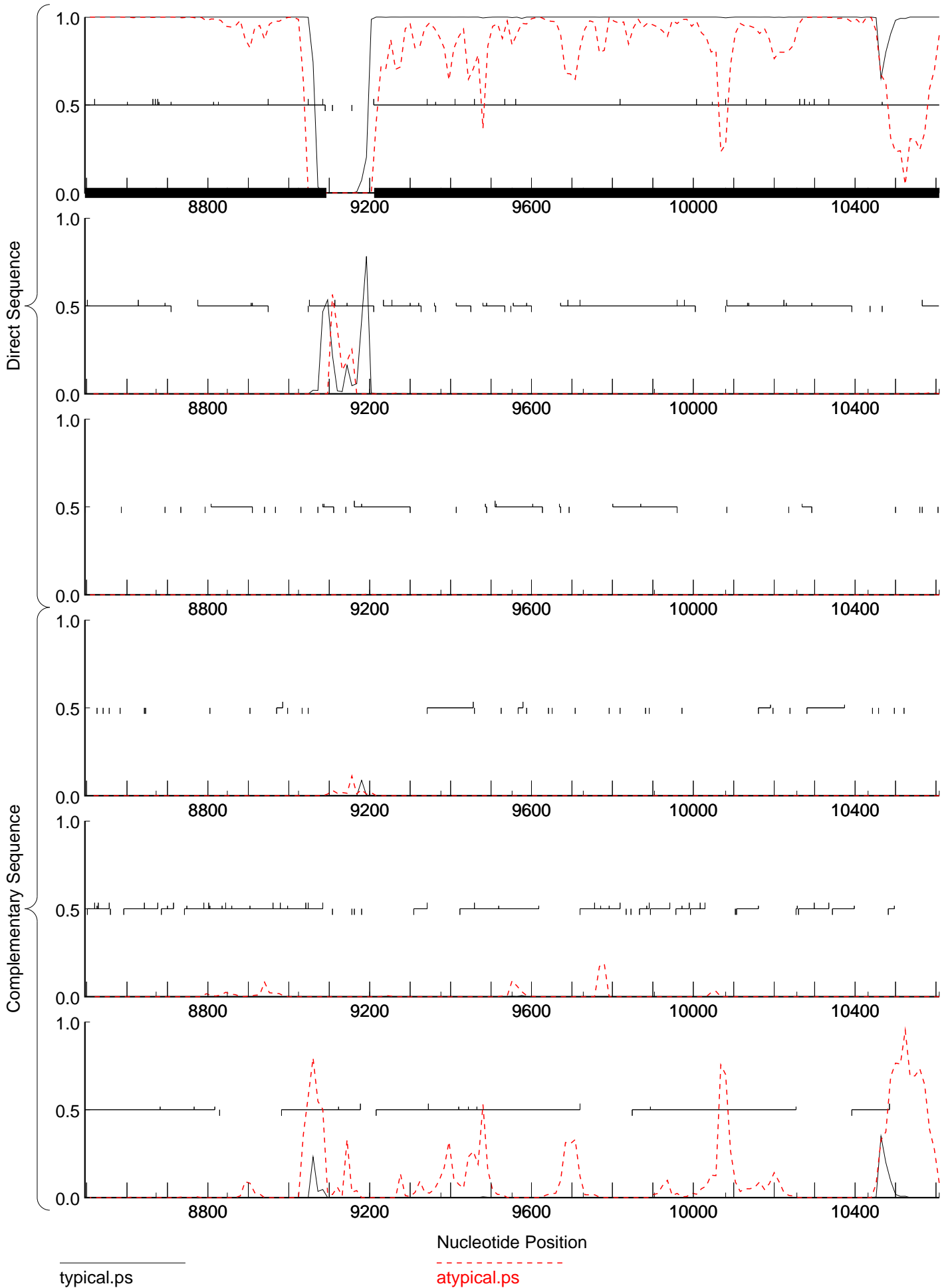
For help or information, send electronic mail to: [custserv@genepro.com](mailto:custserv@genepro.com), or [mark@amber.biology.gatech.edu](mailto:mark@amber.biology.gatech.edu)

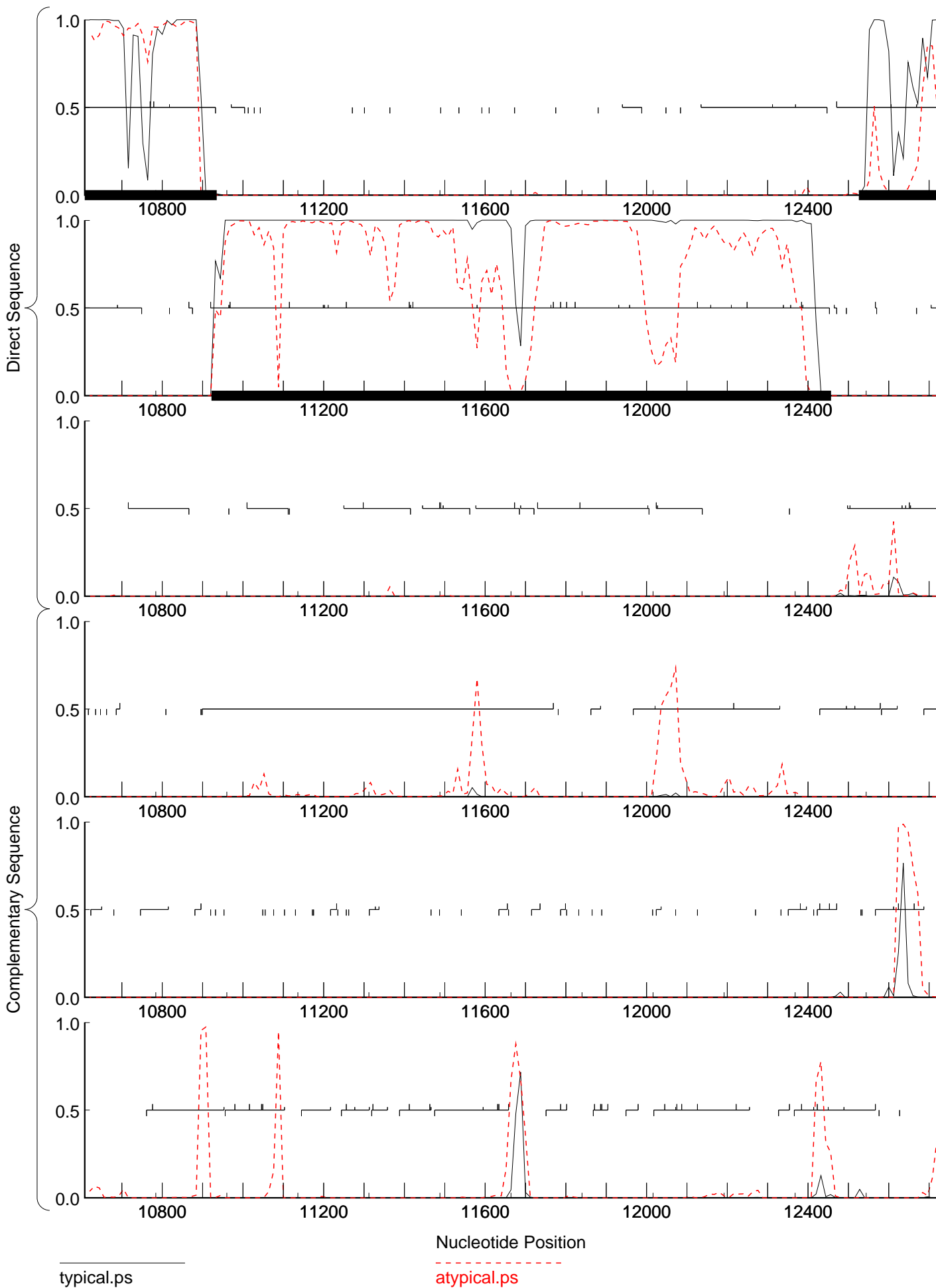






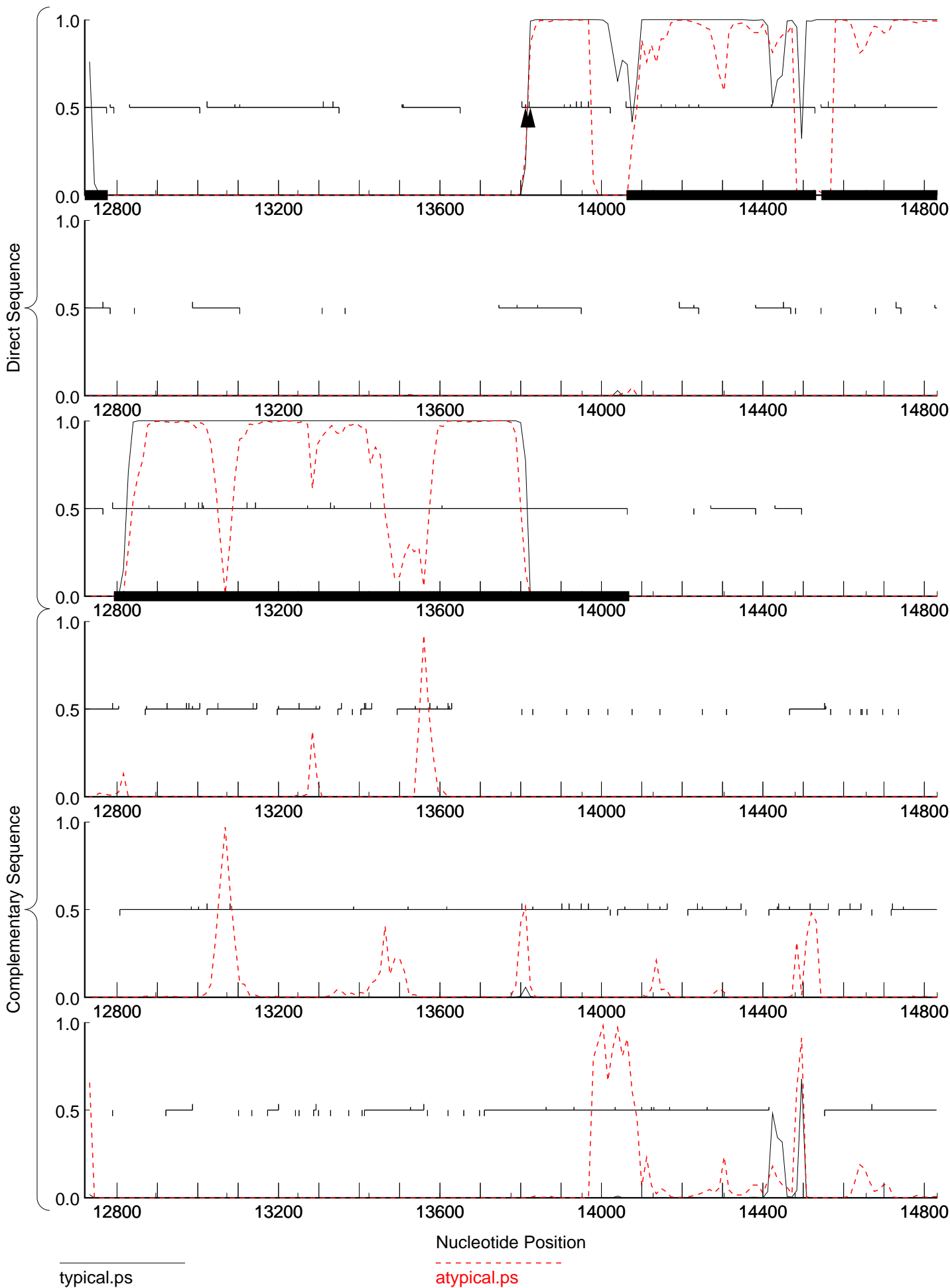






typical.ps

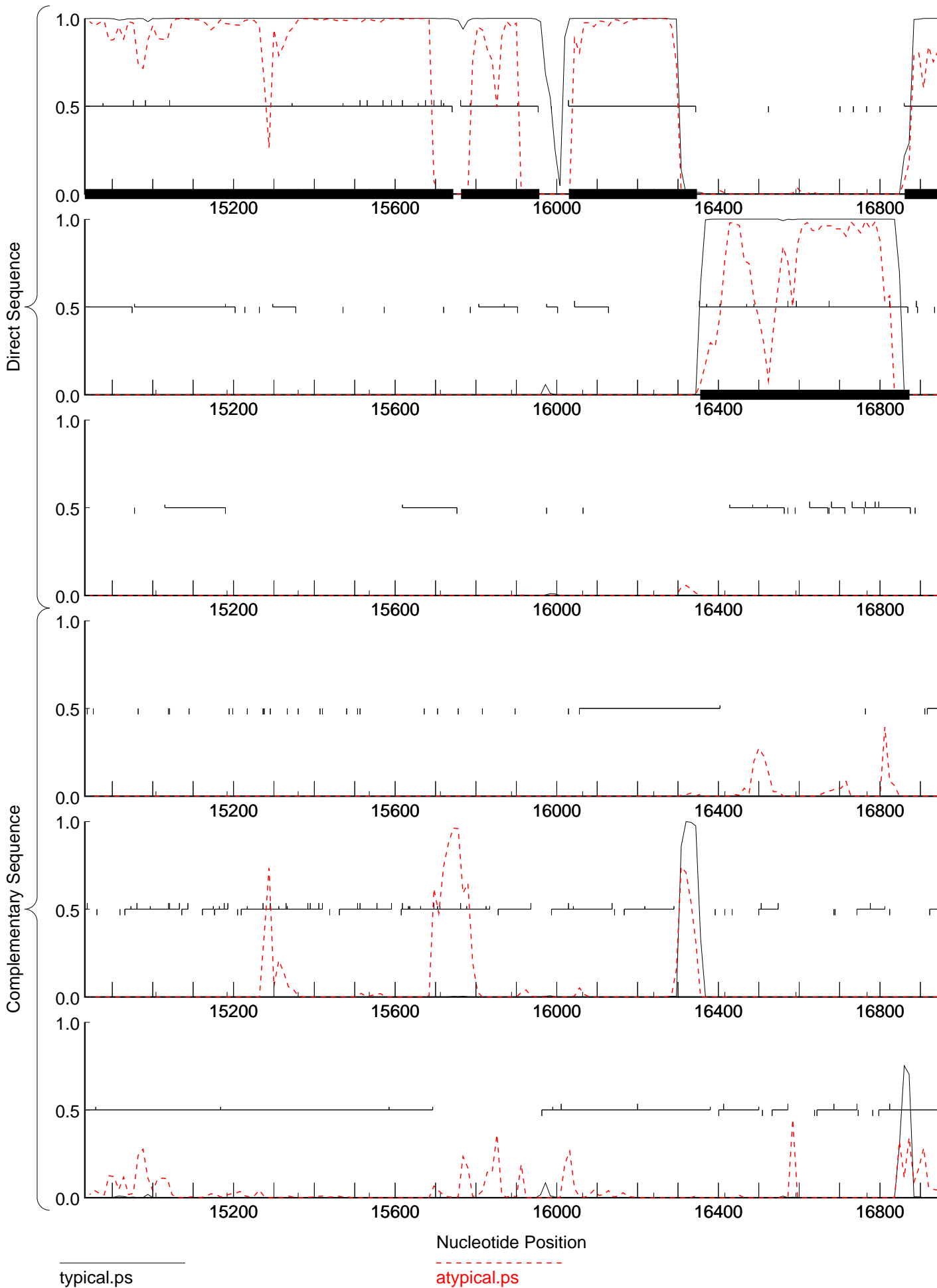
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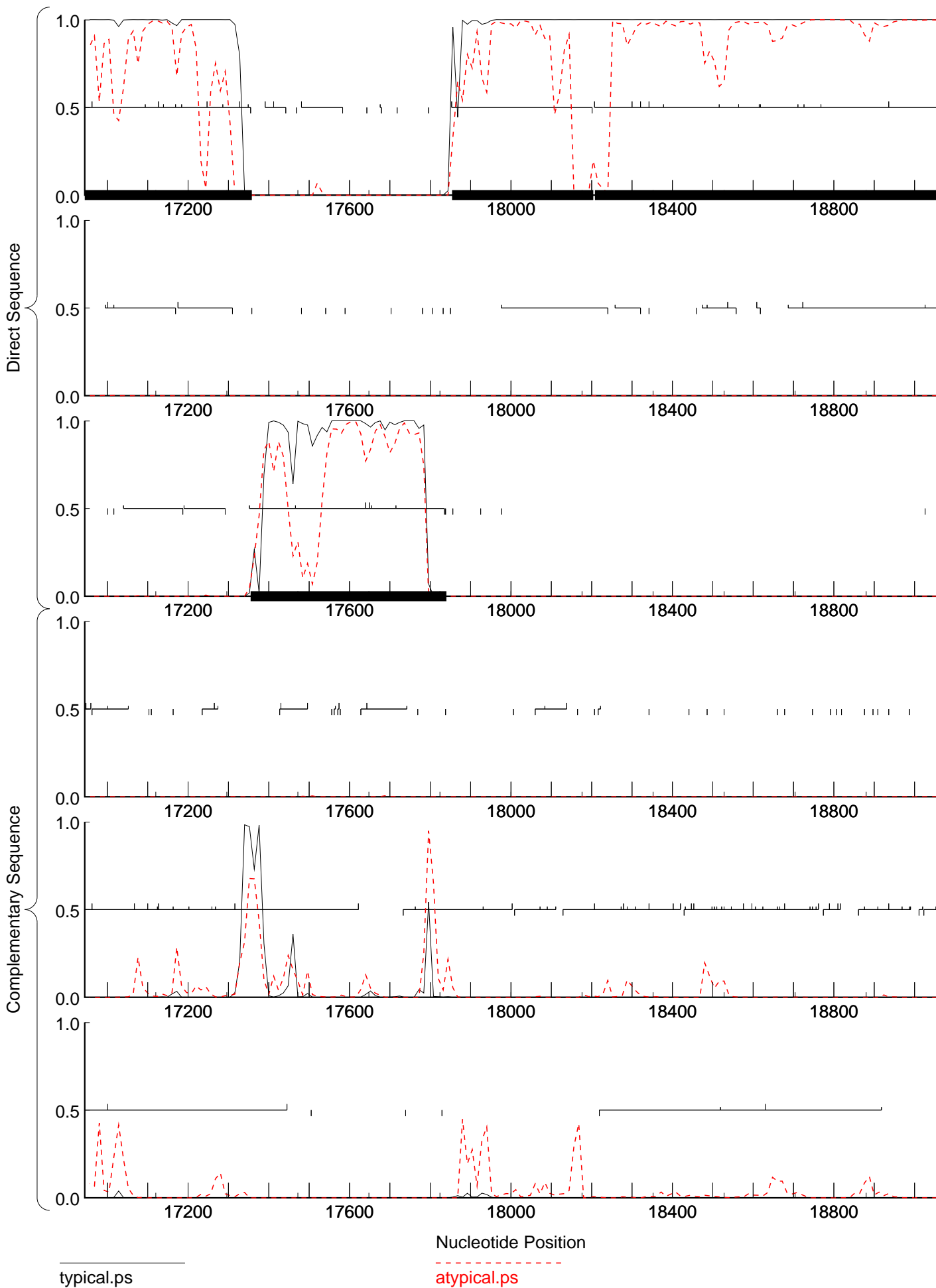


typical.ps

atypical.ps

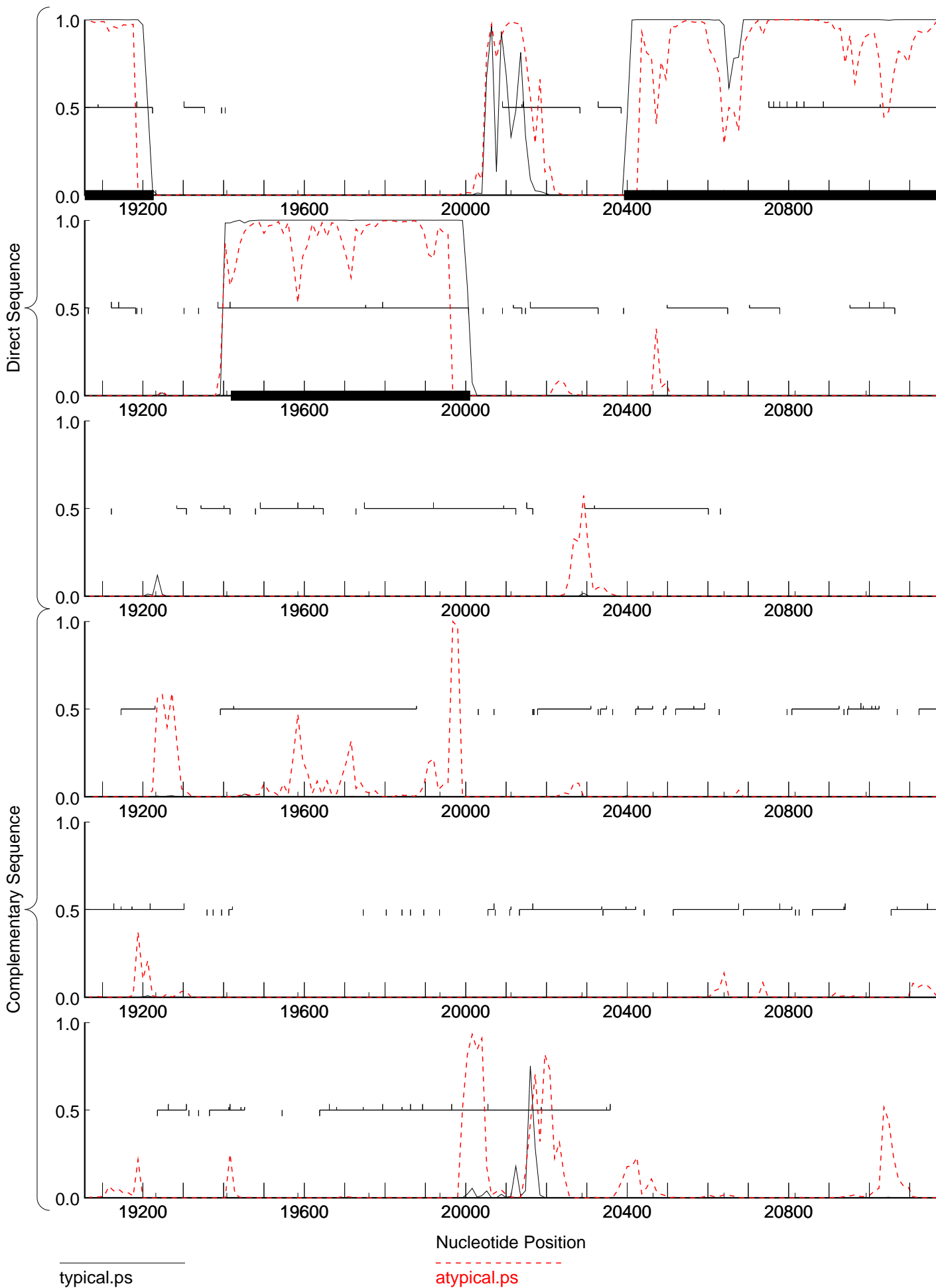


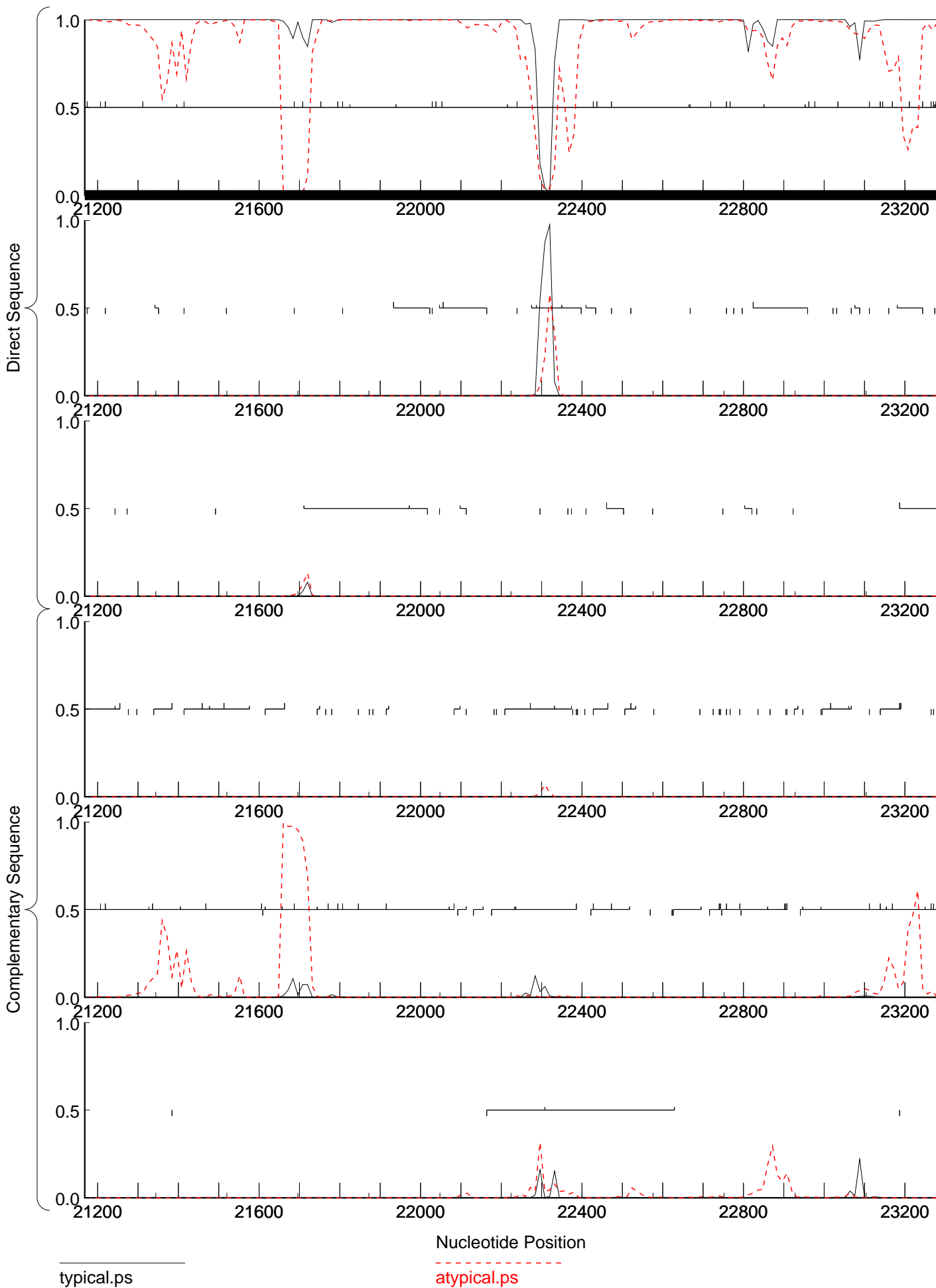




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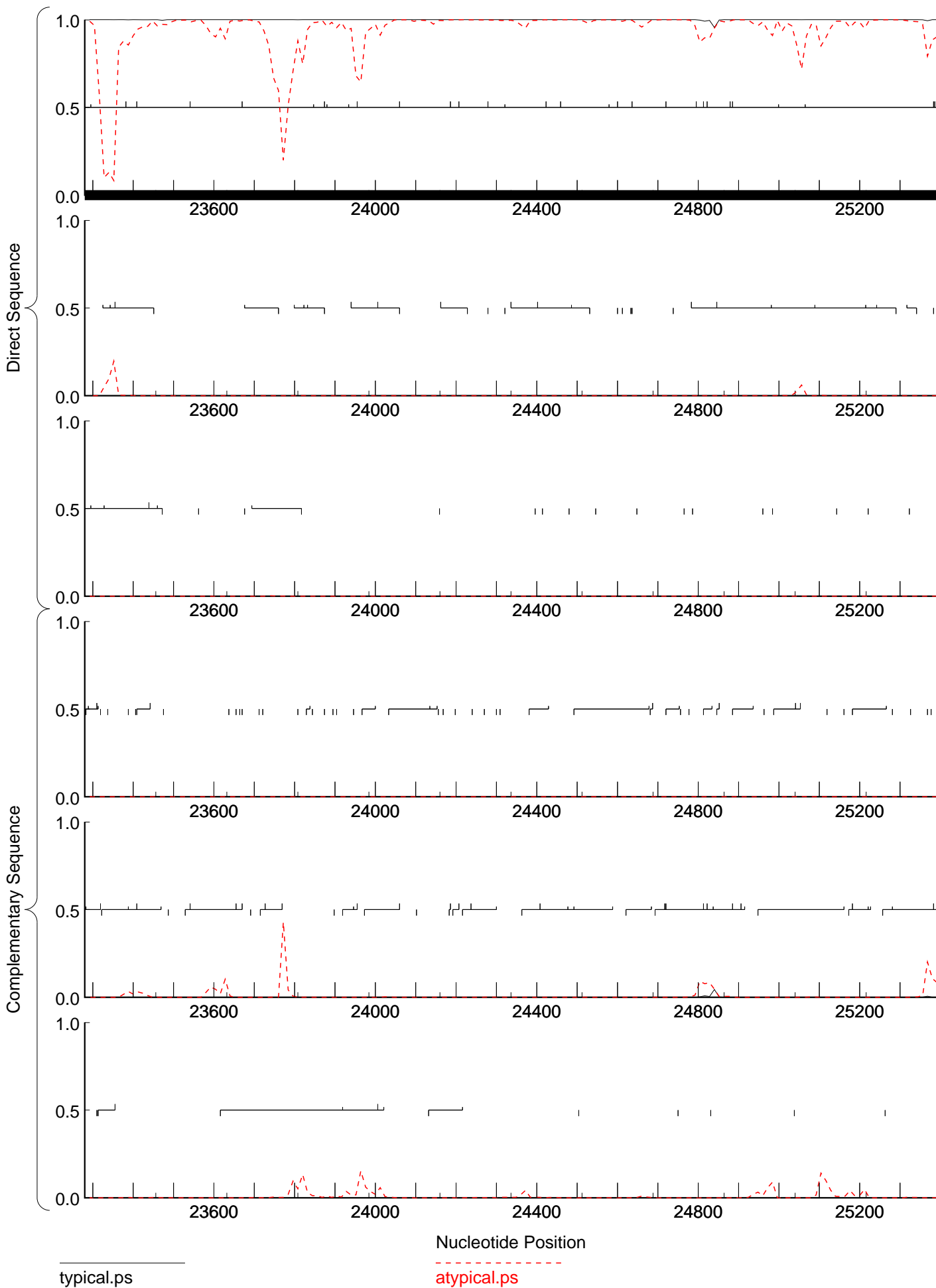
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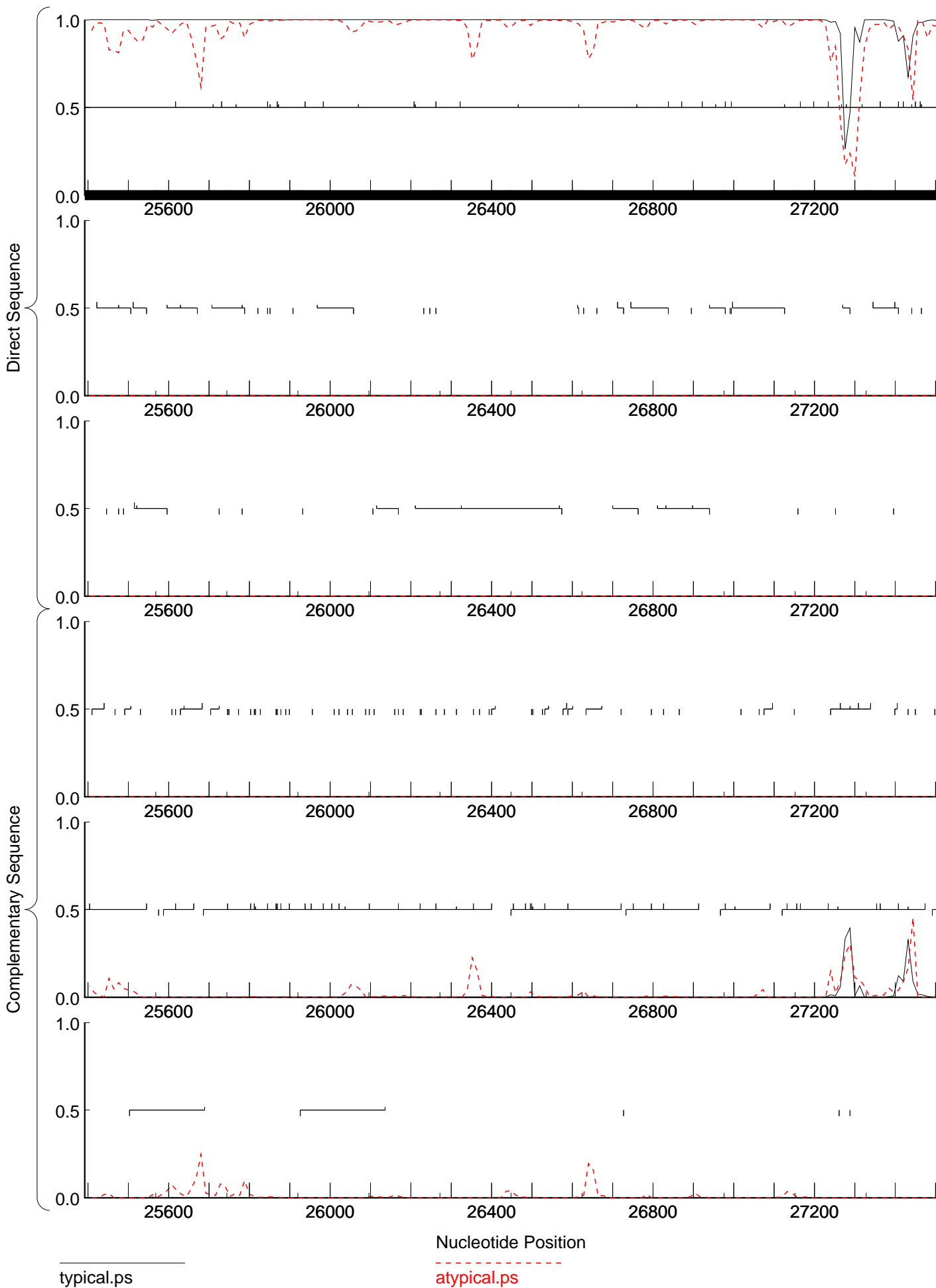


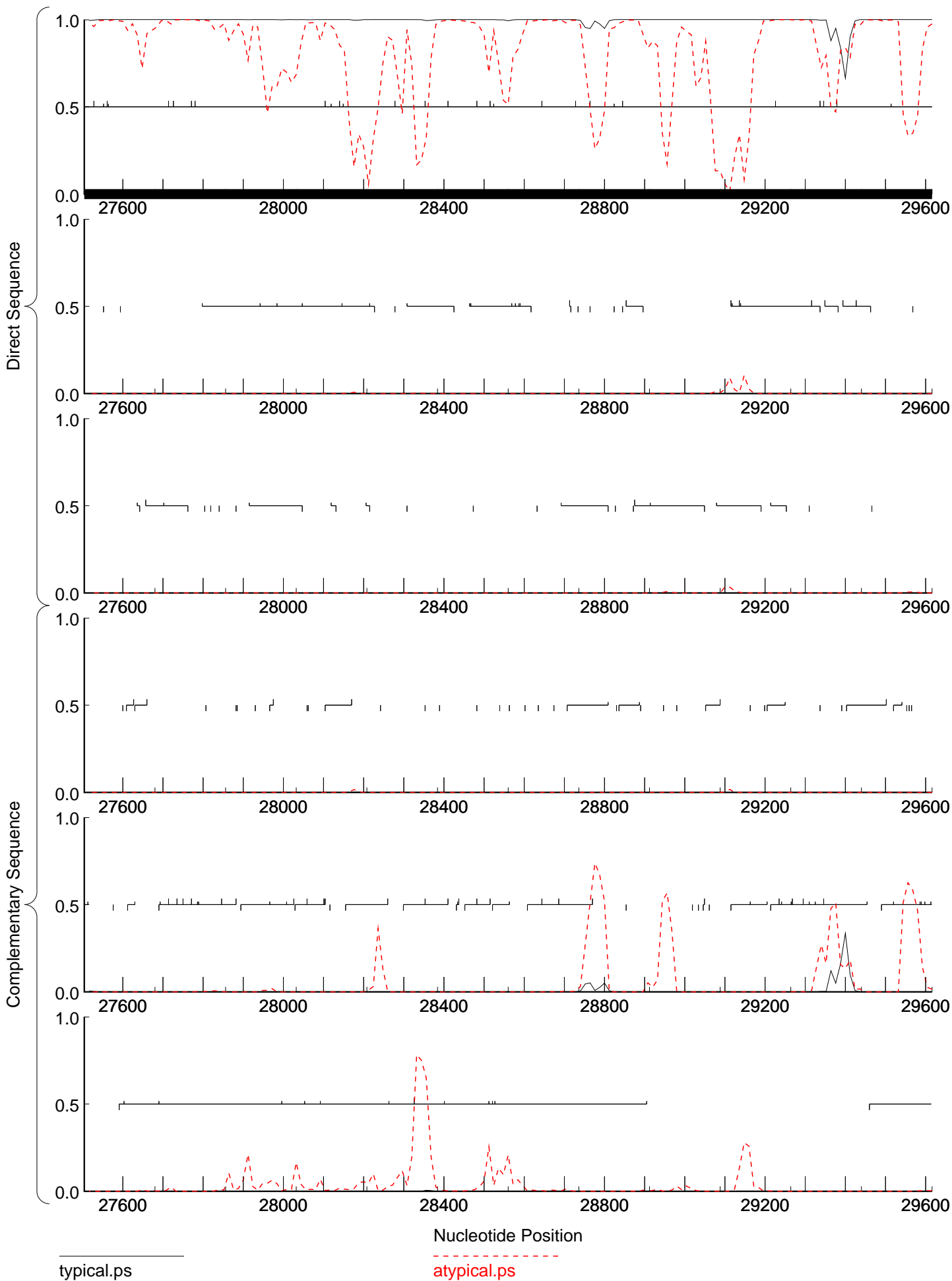


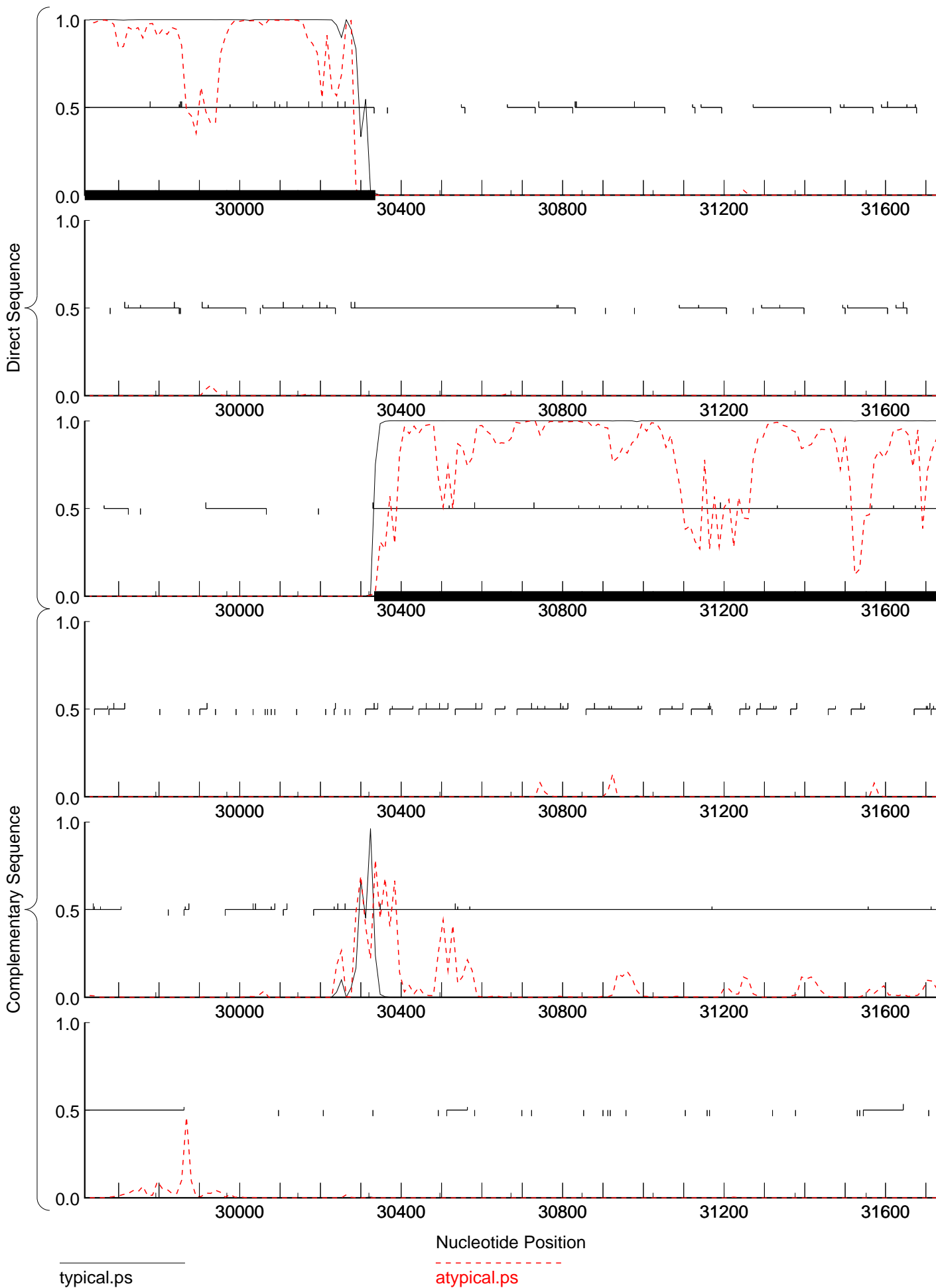
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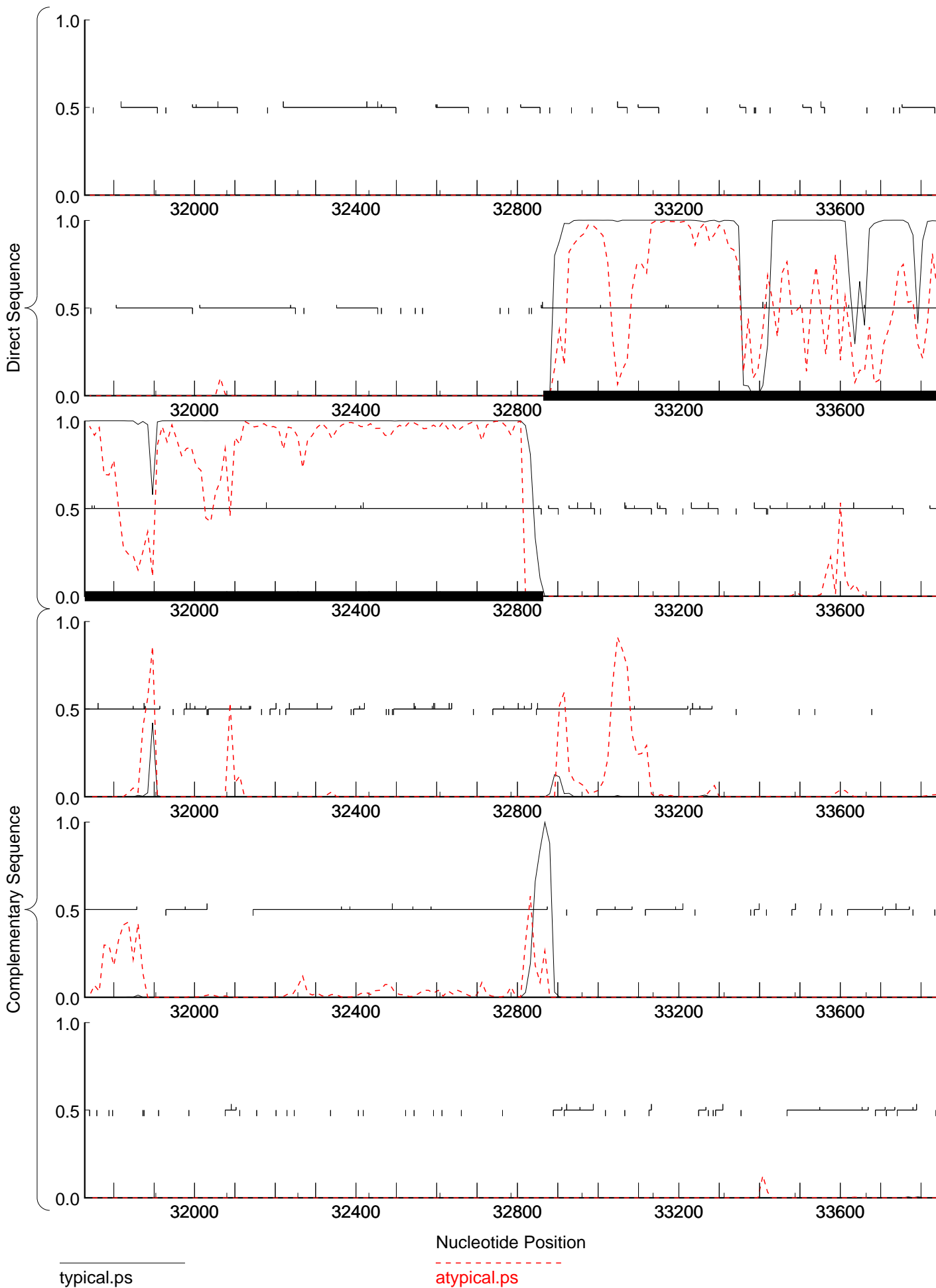


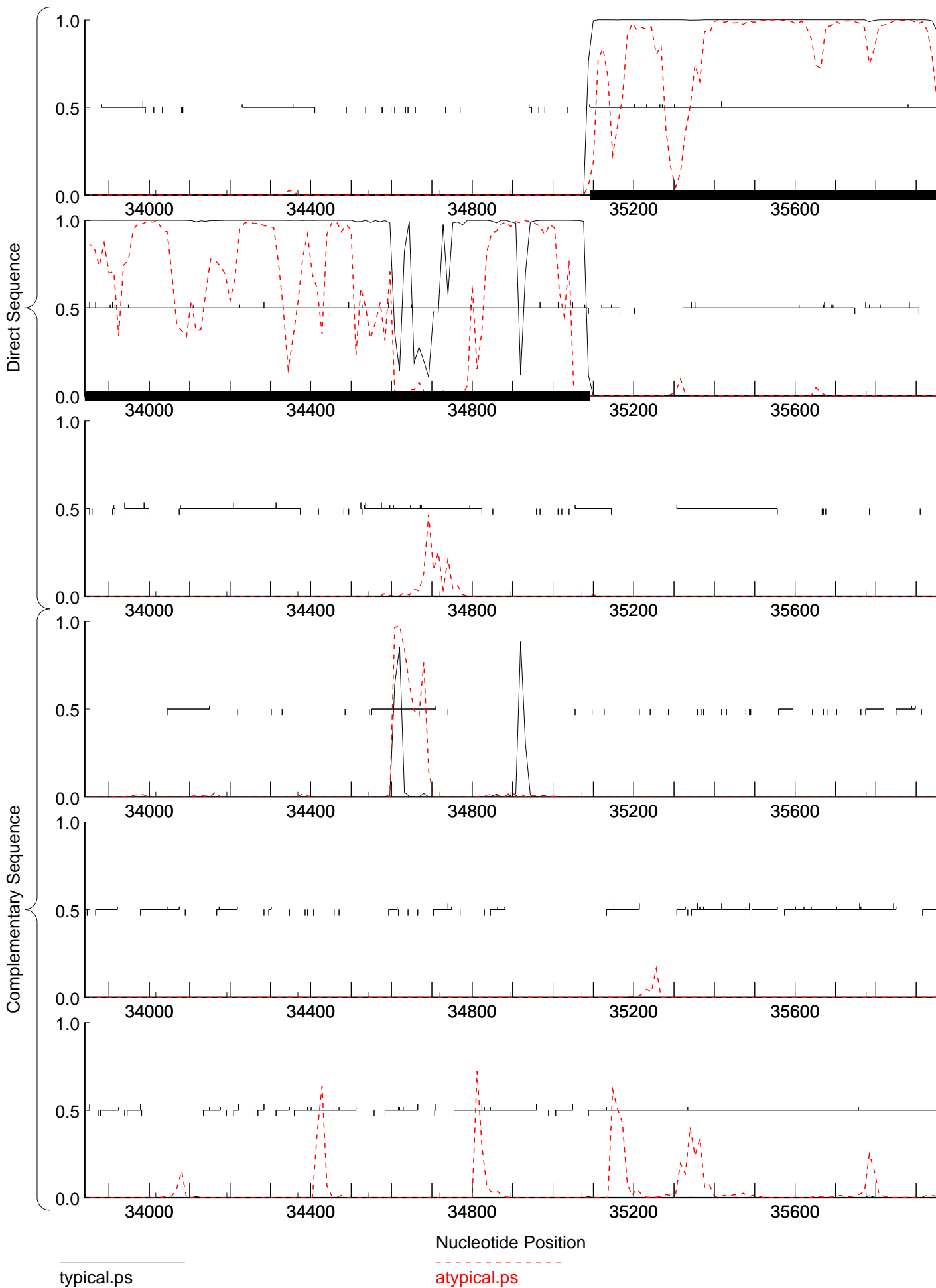


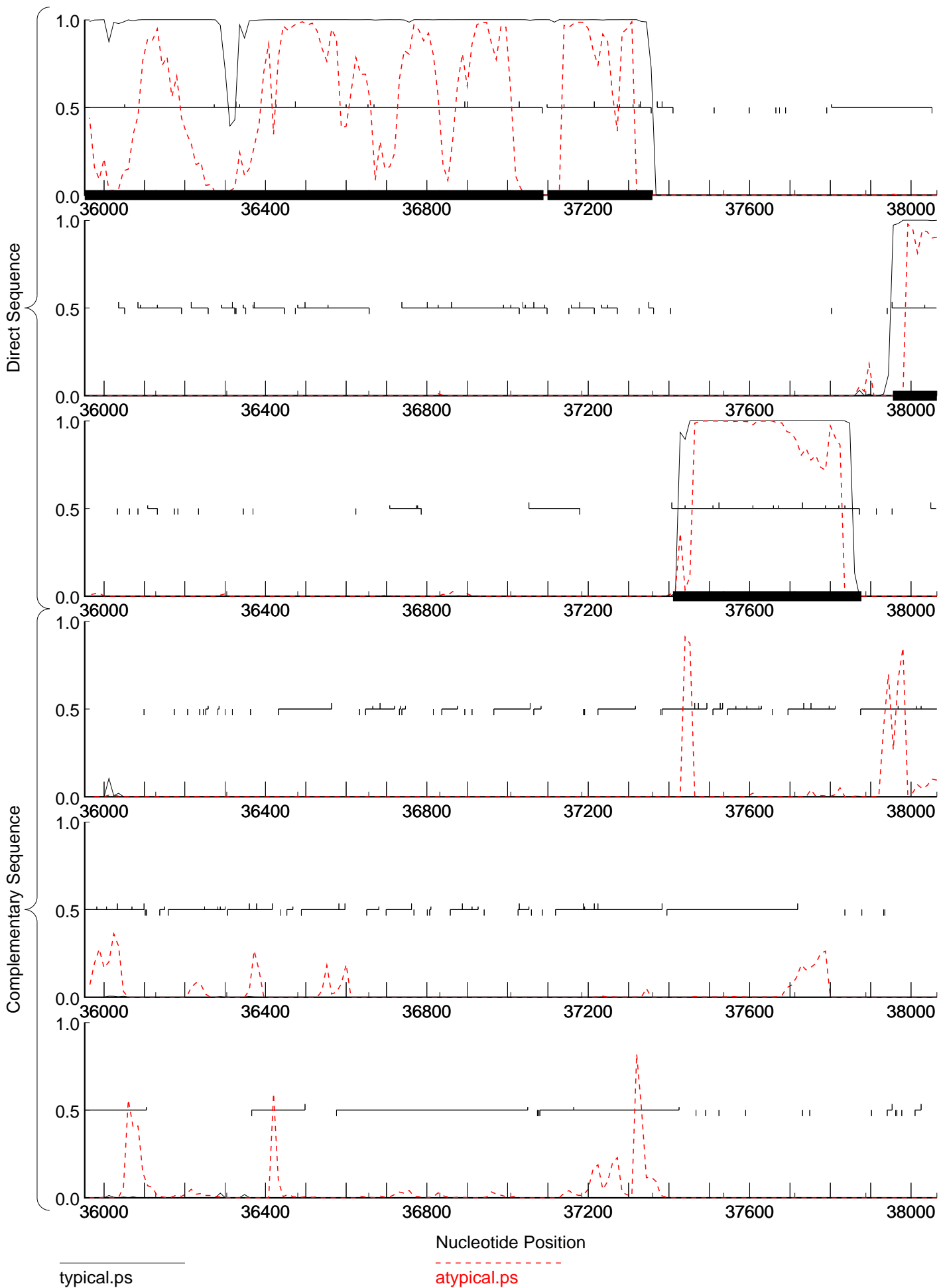






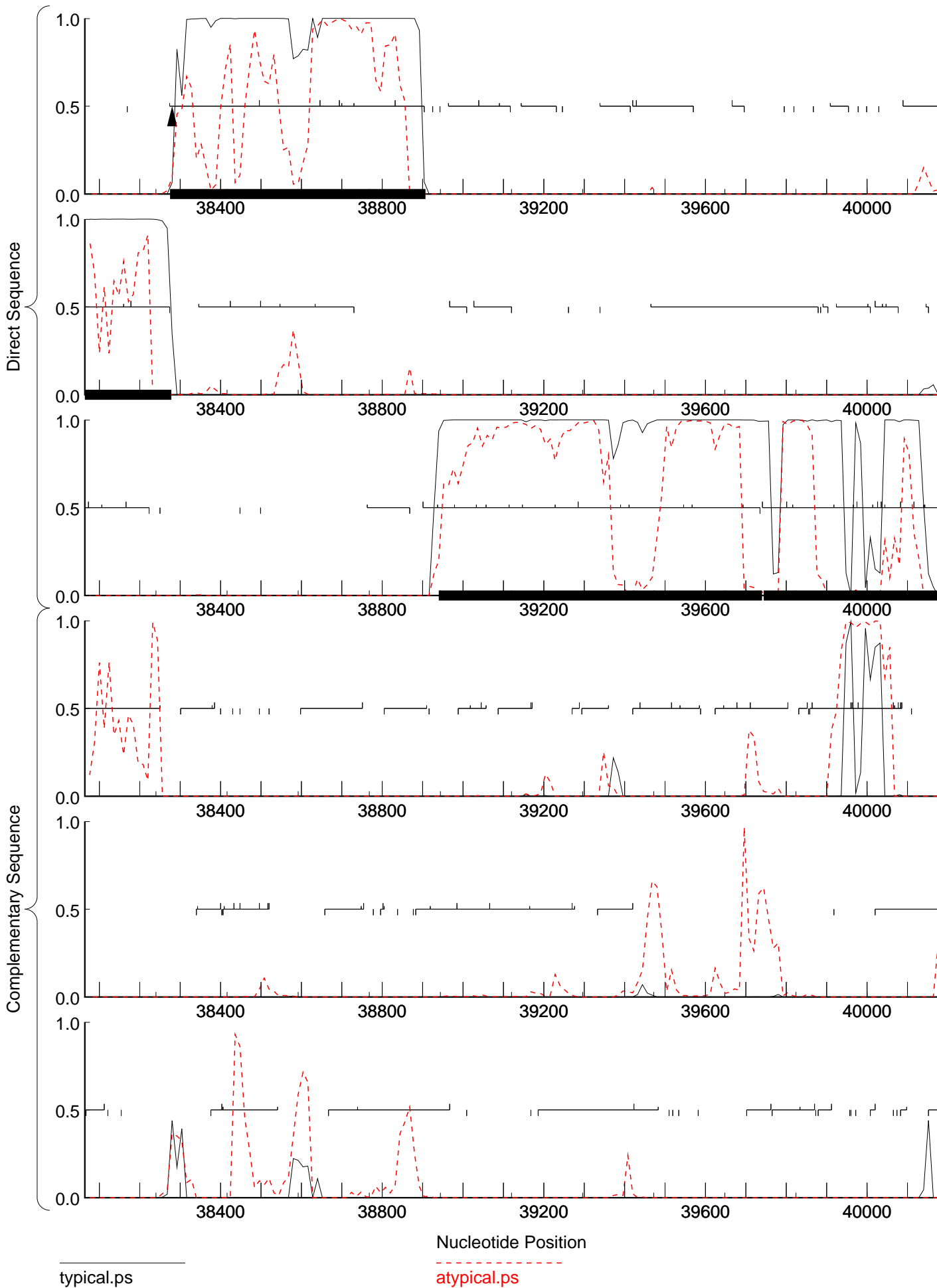


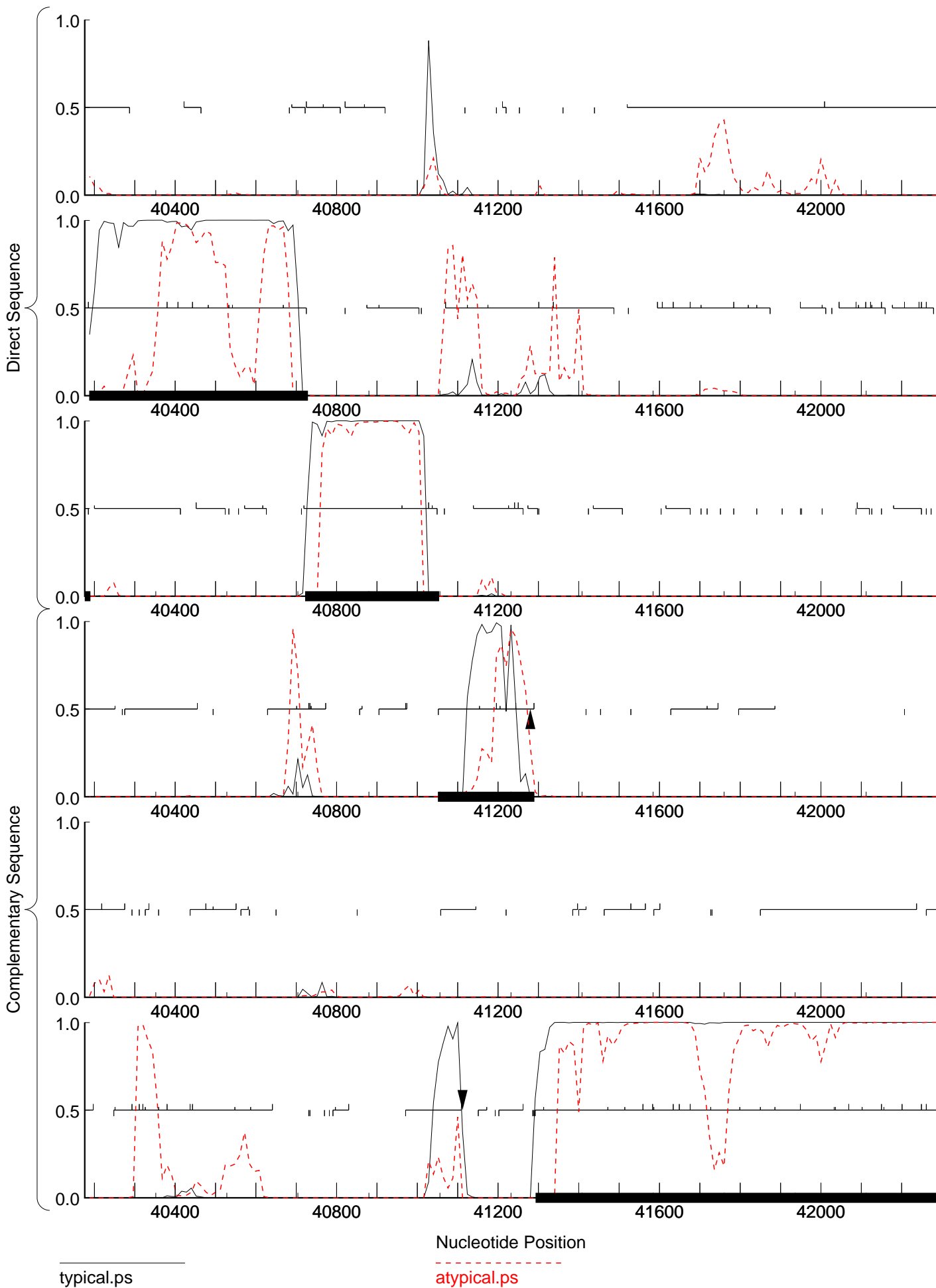


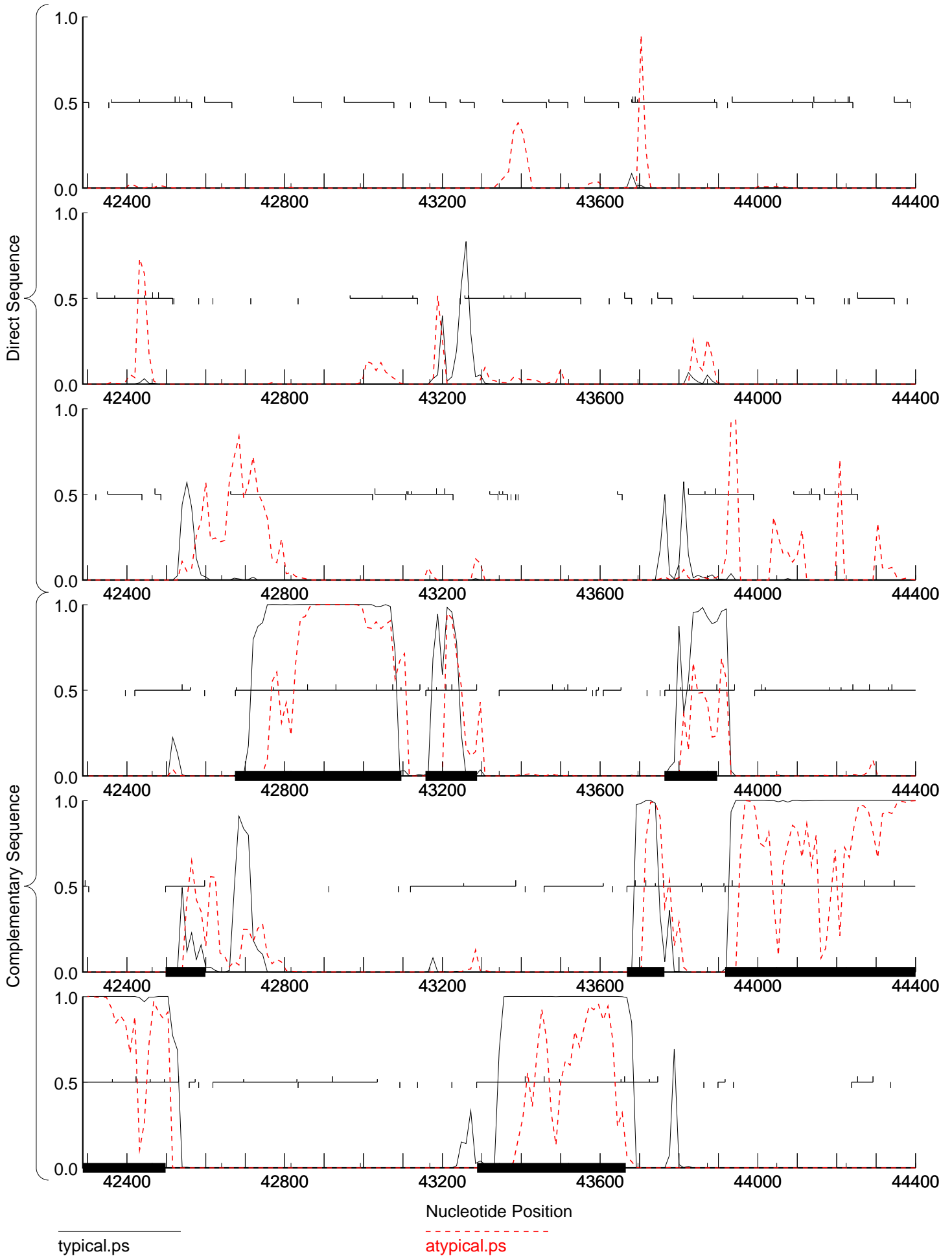


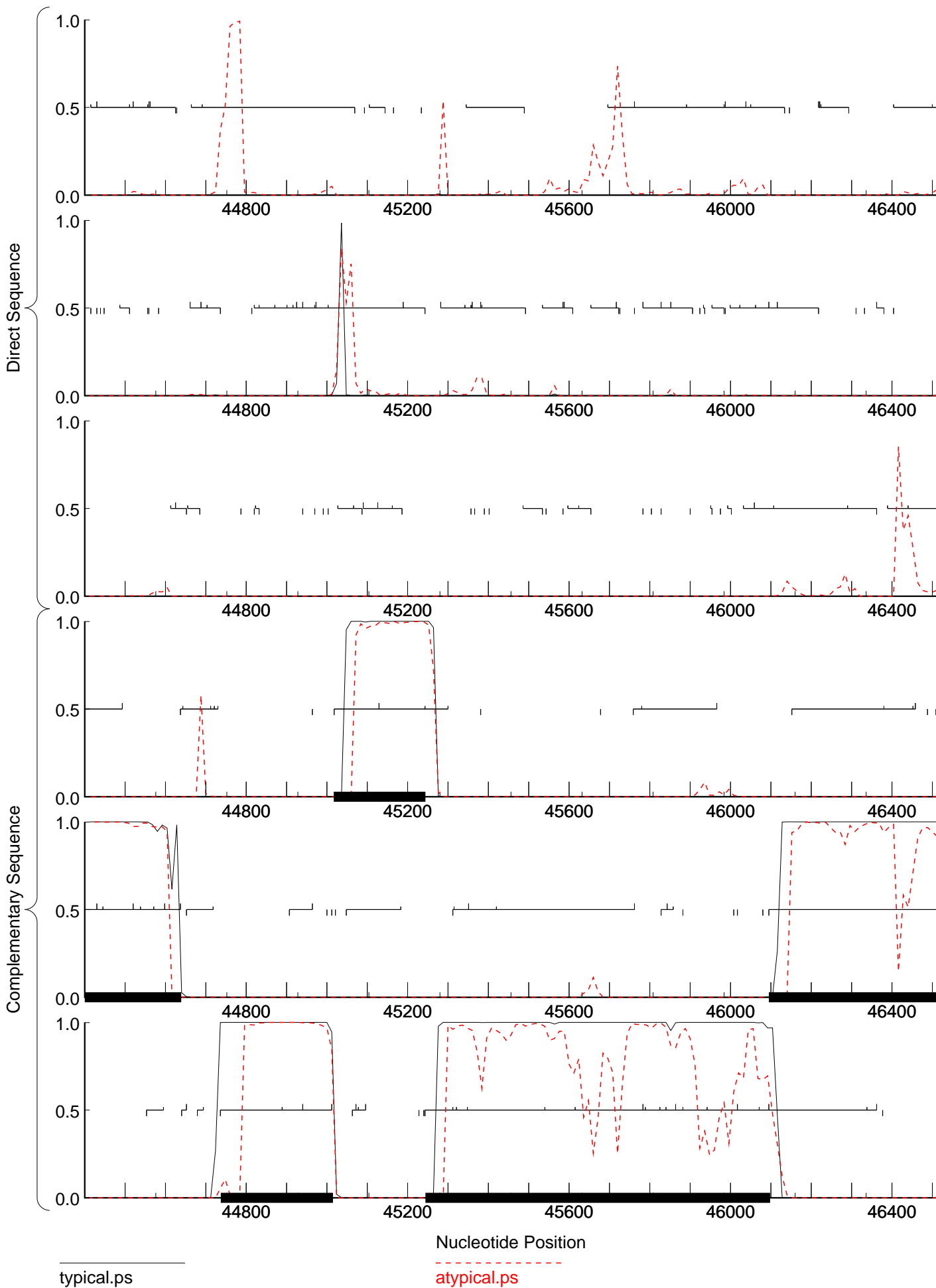
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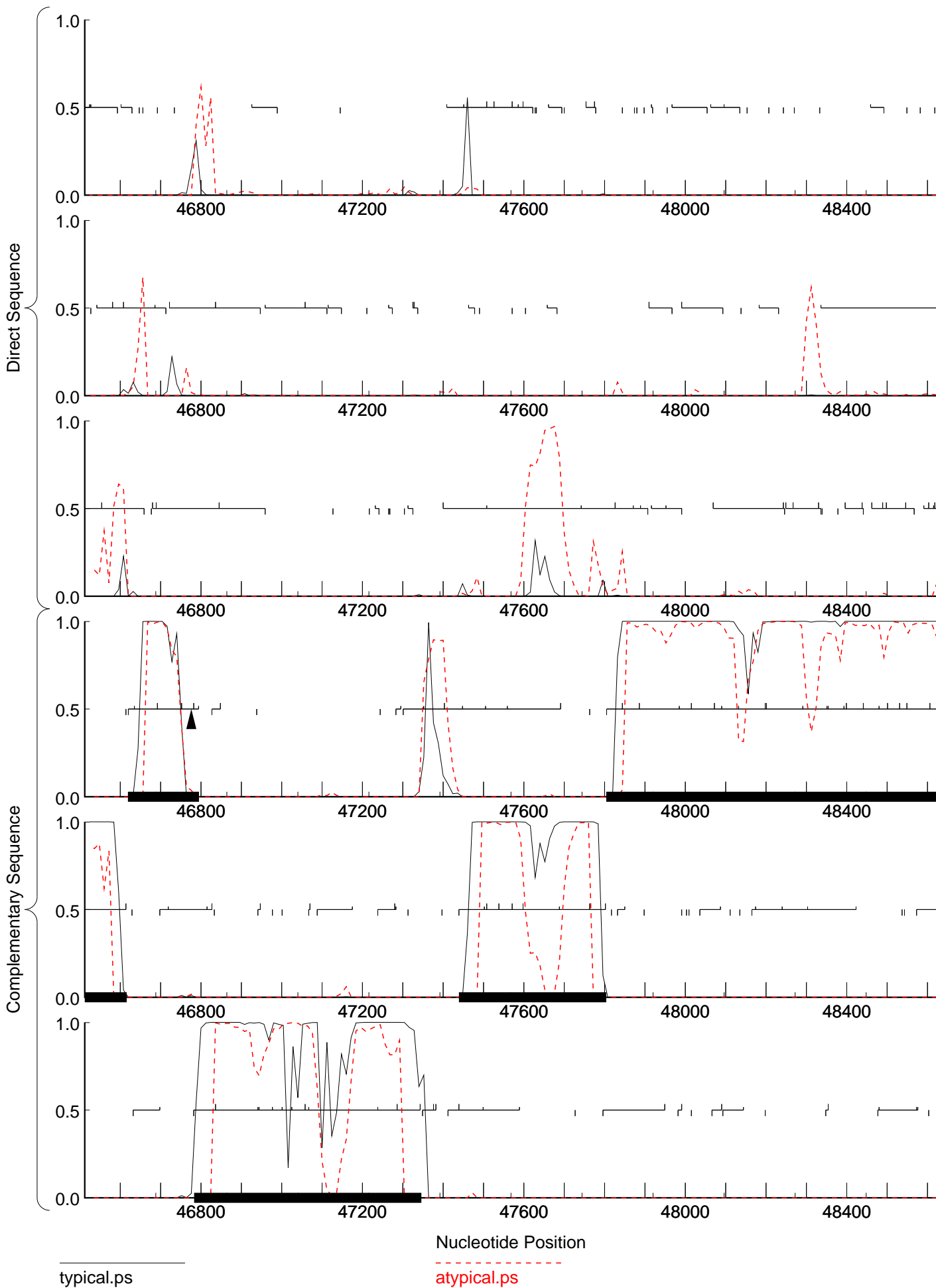
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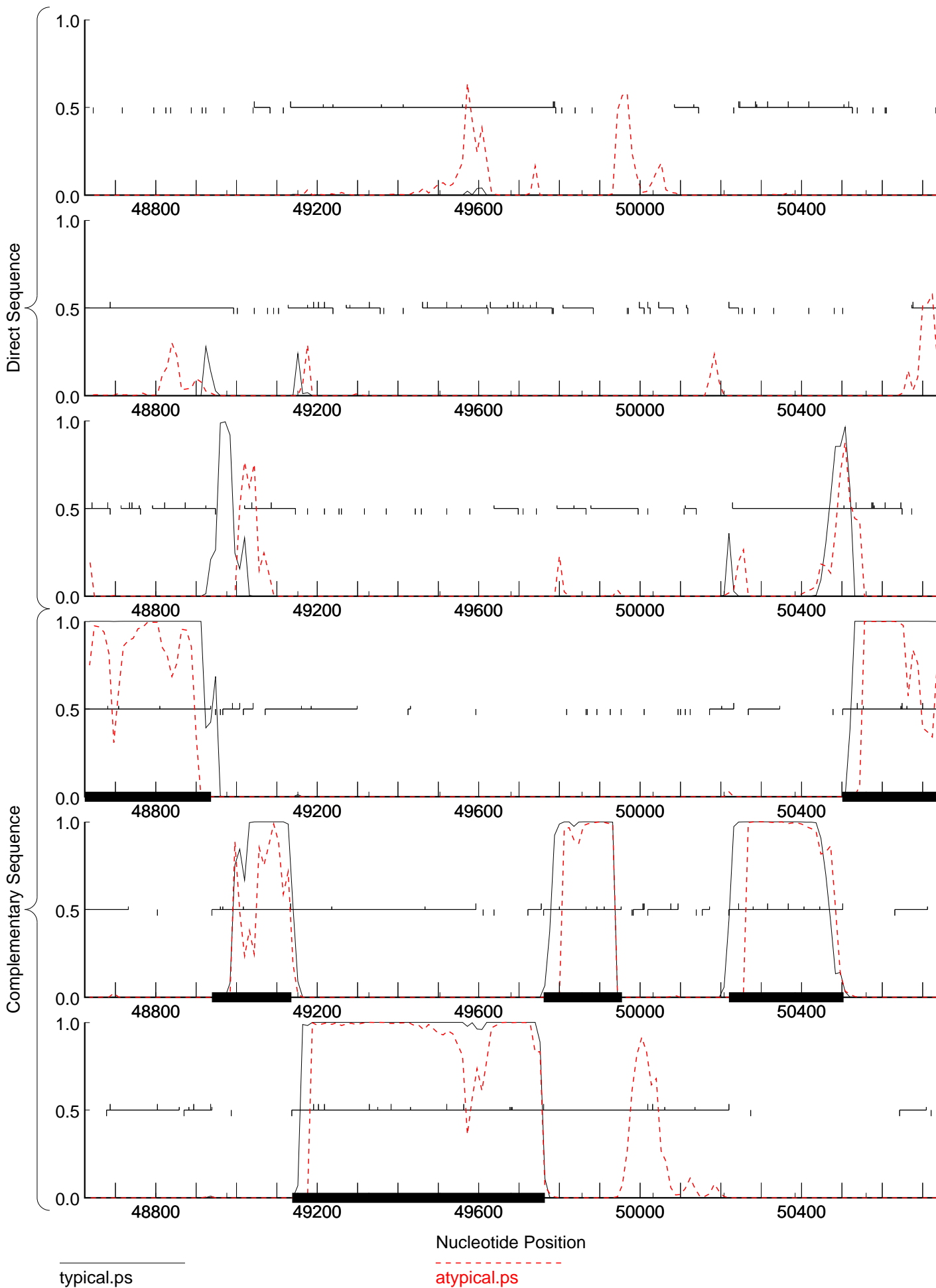


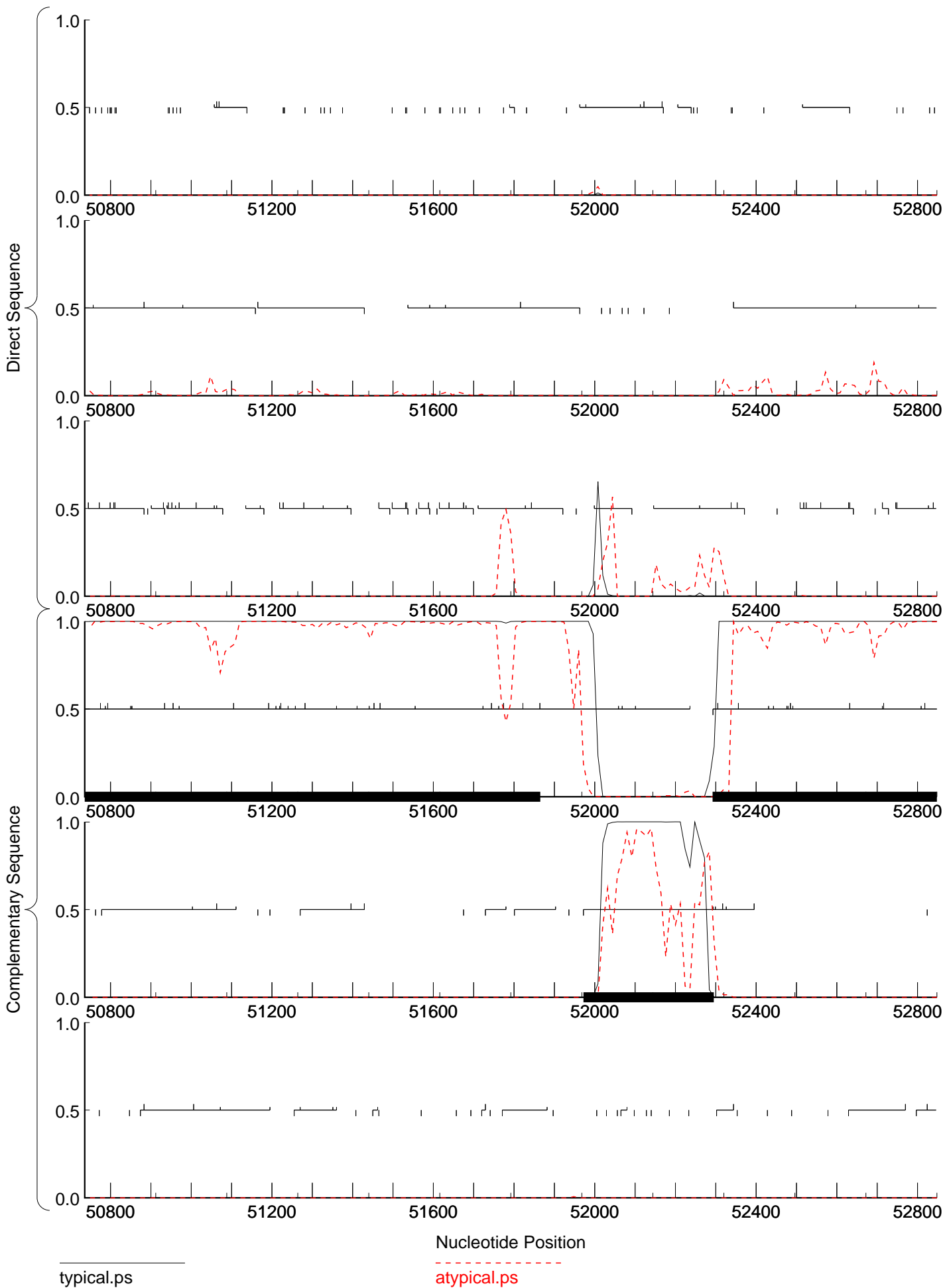


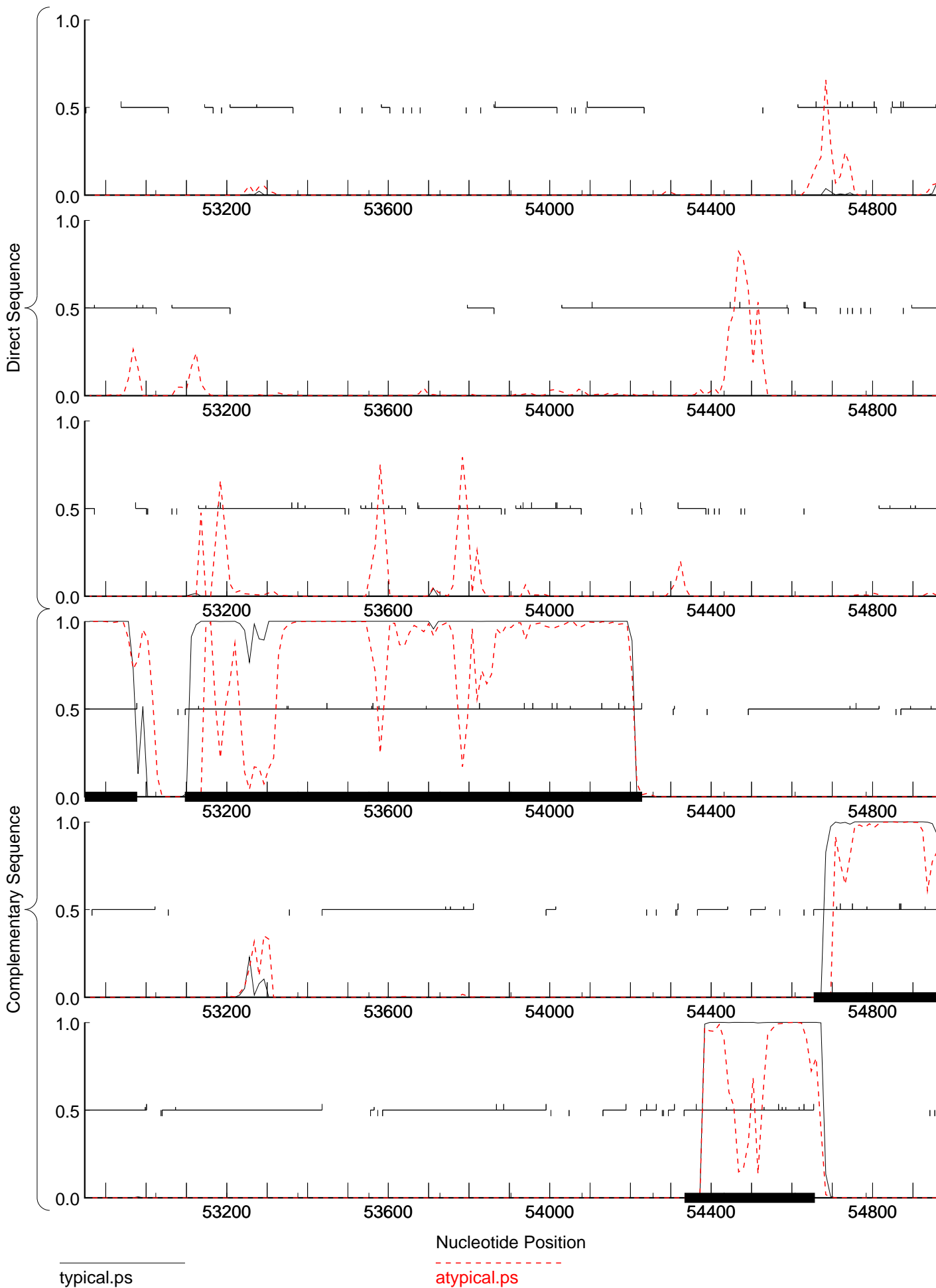






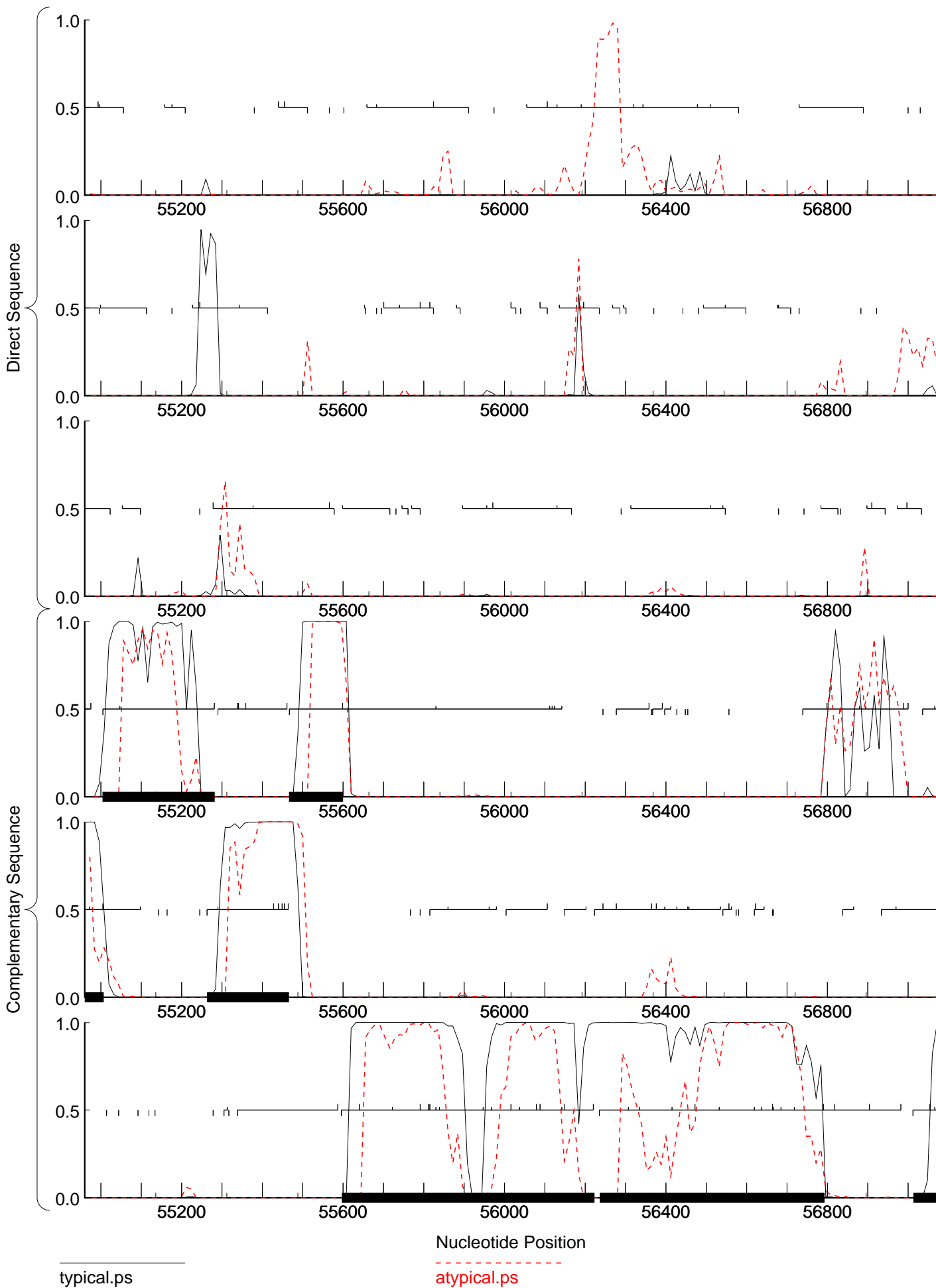






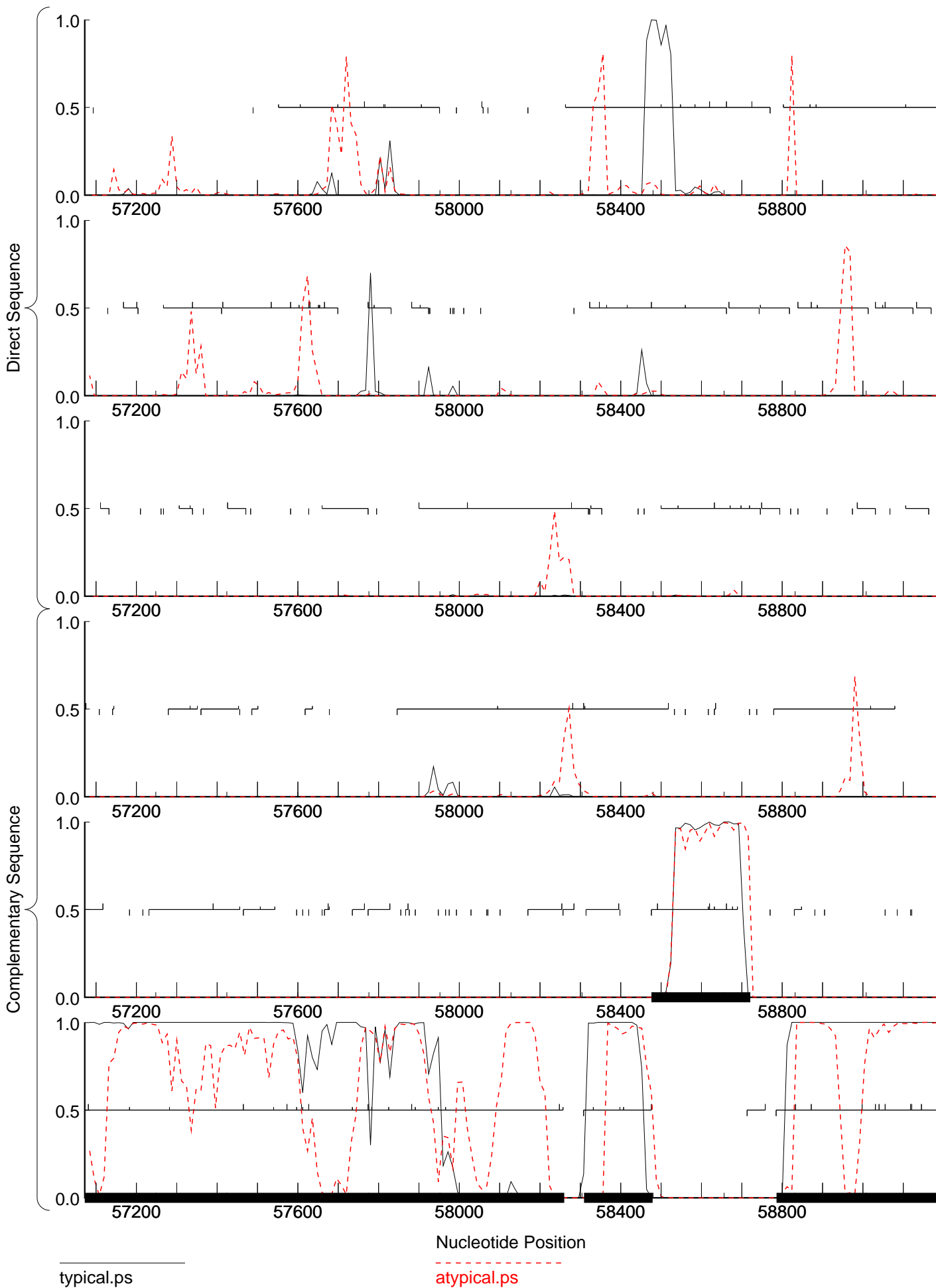
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atypical.ps



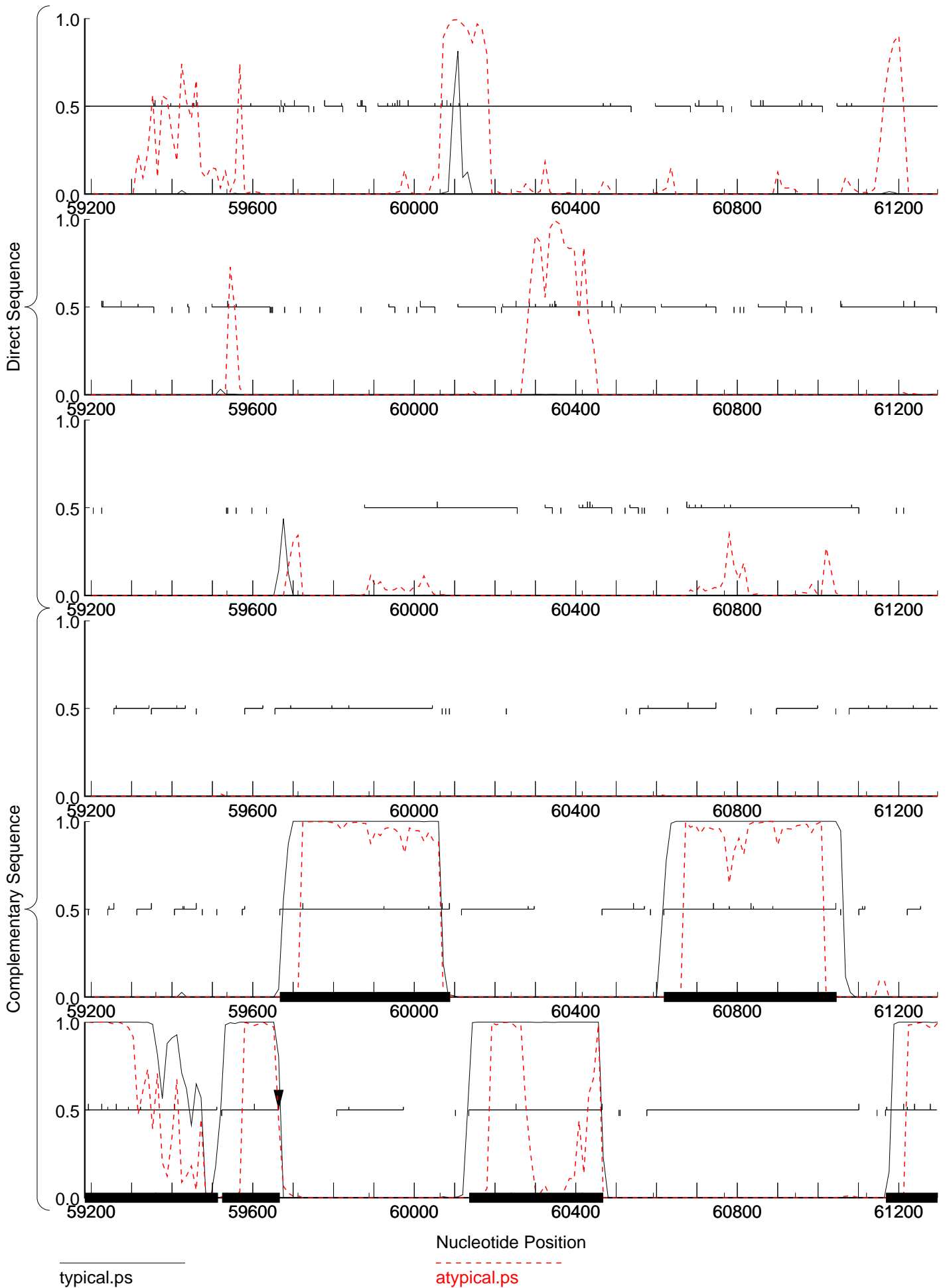
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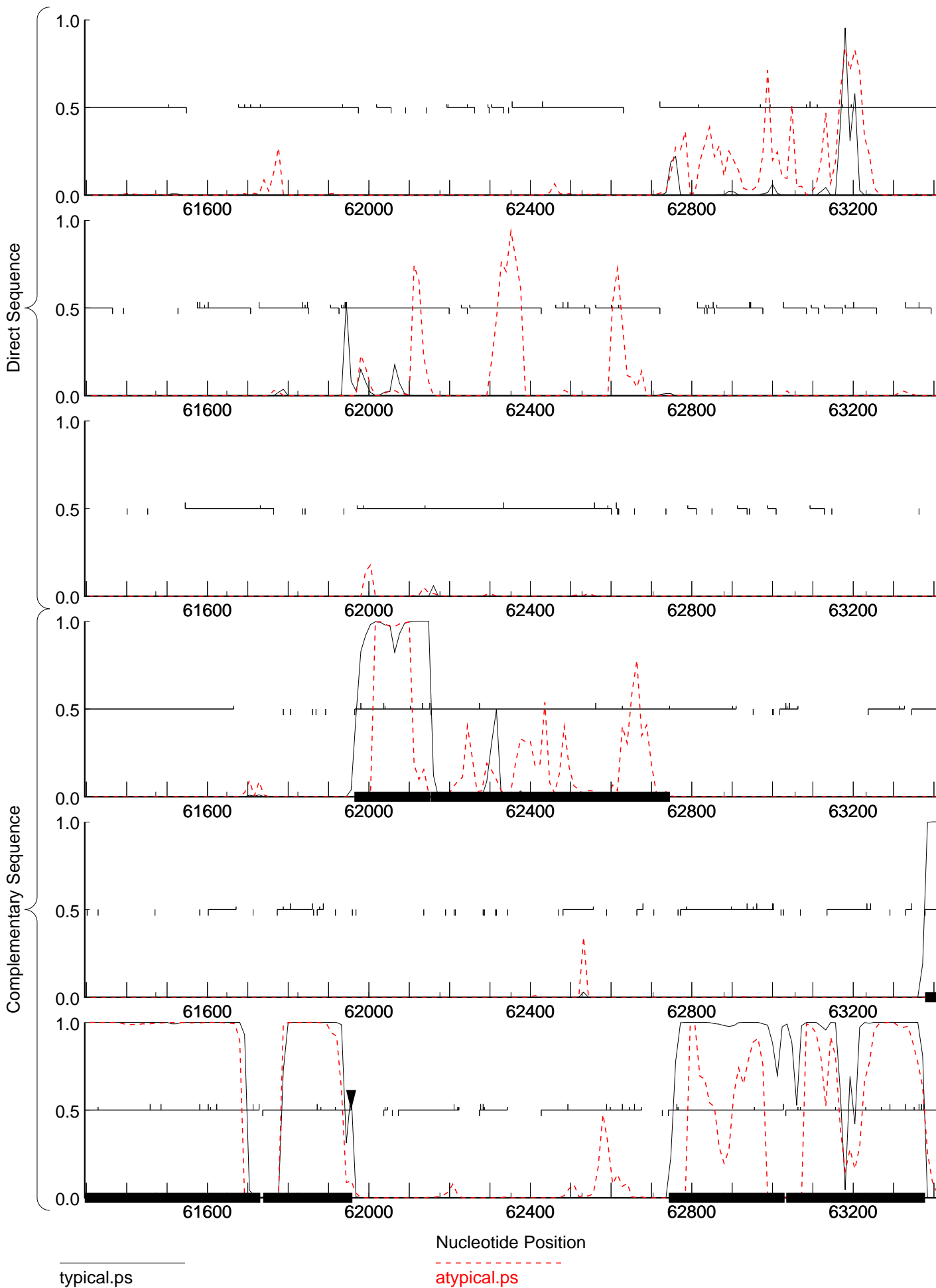
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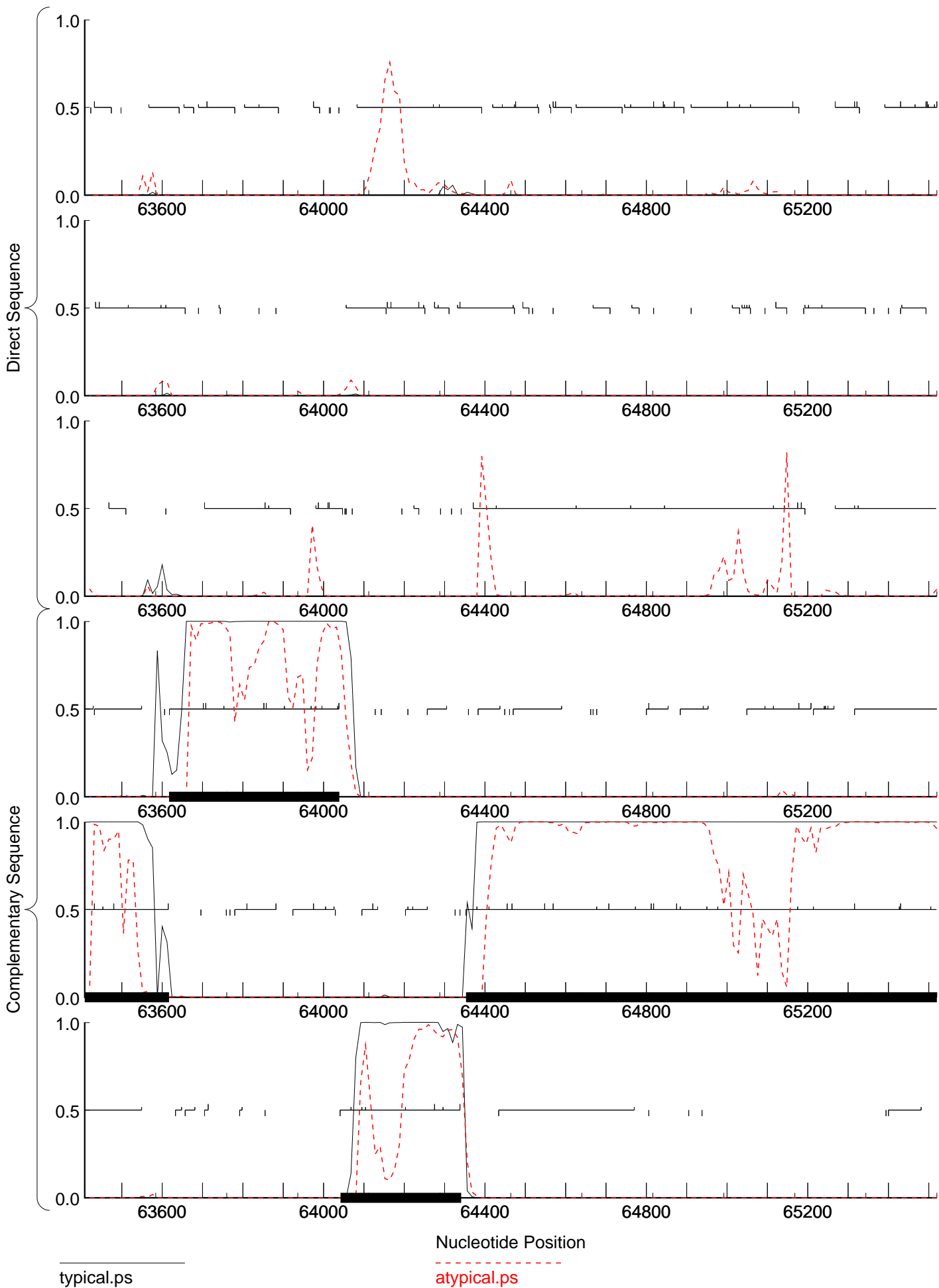


typical.ps

atypical.ps



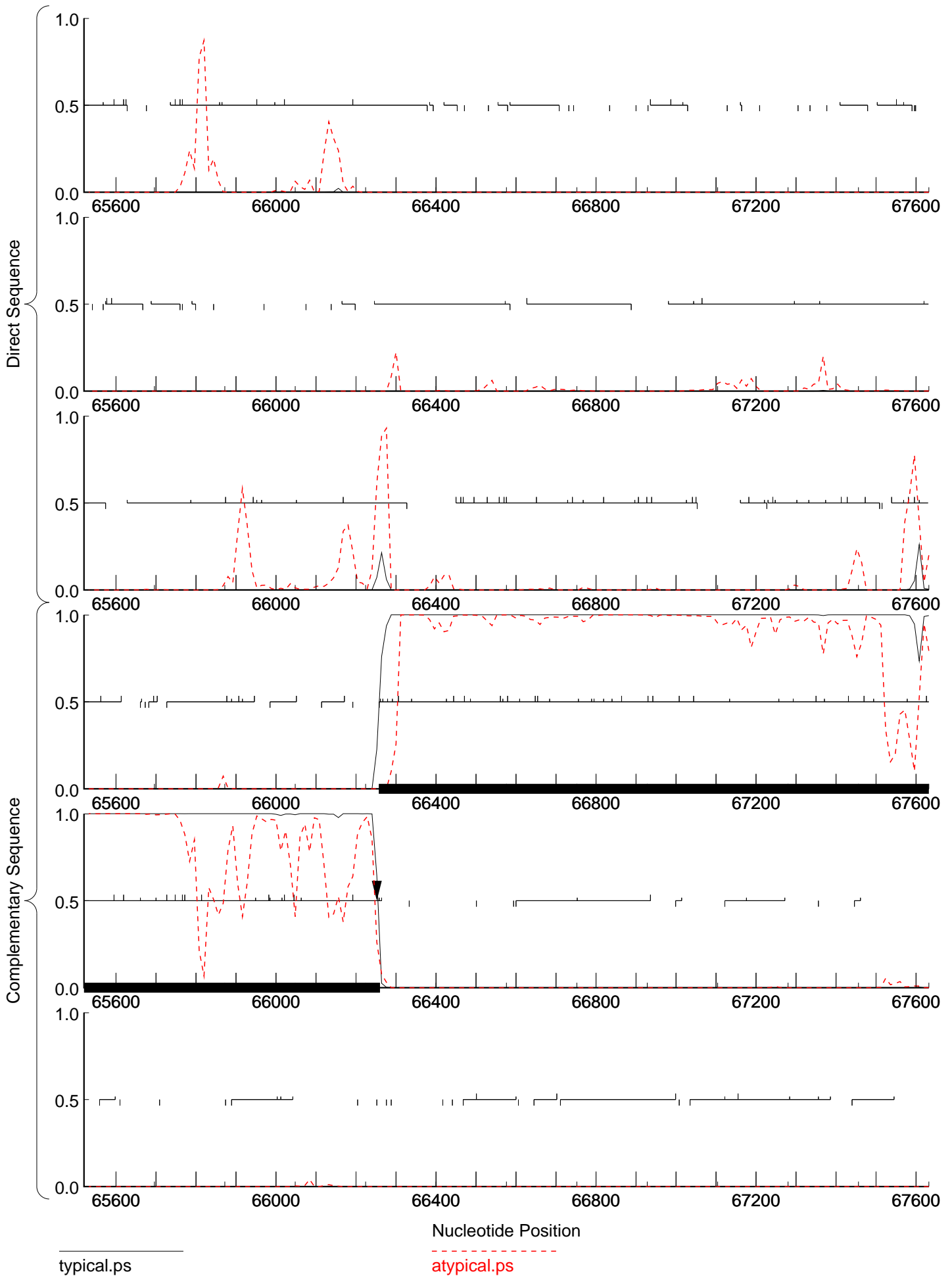




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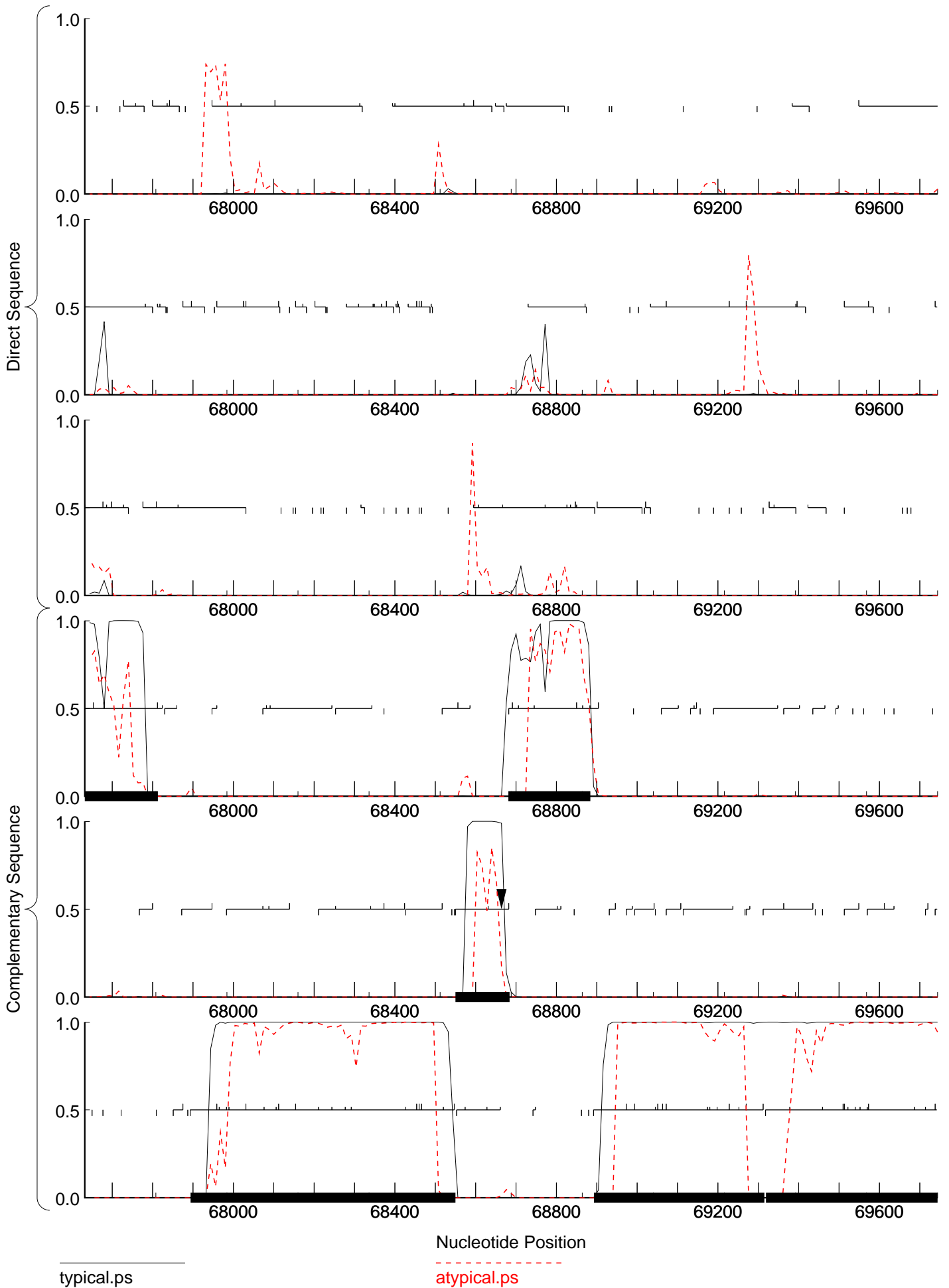
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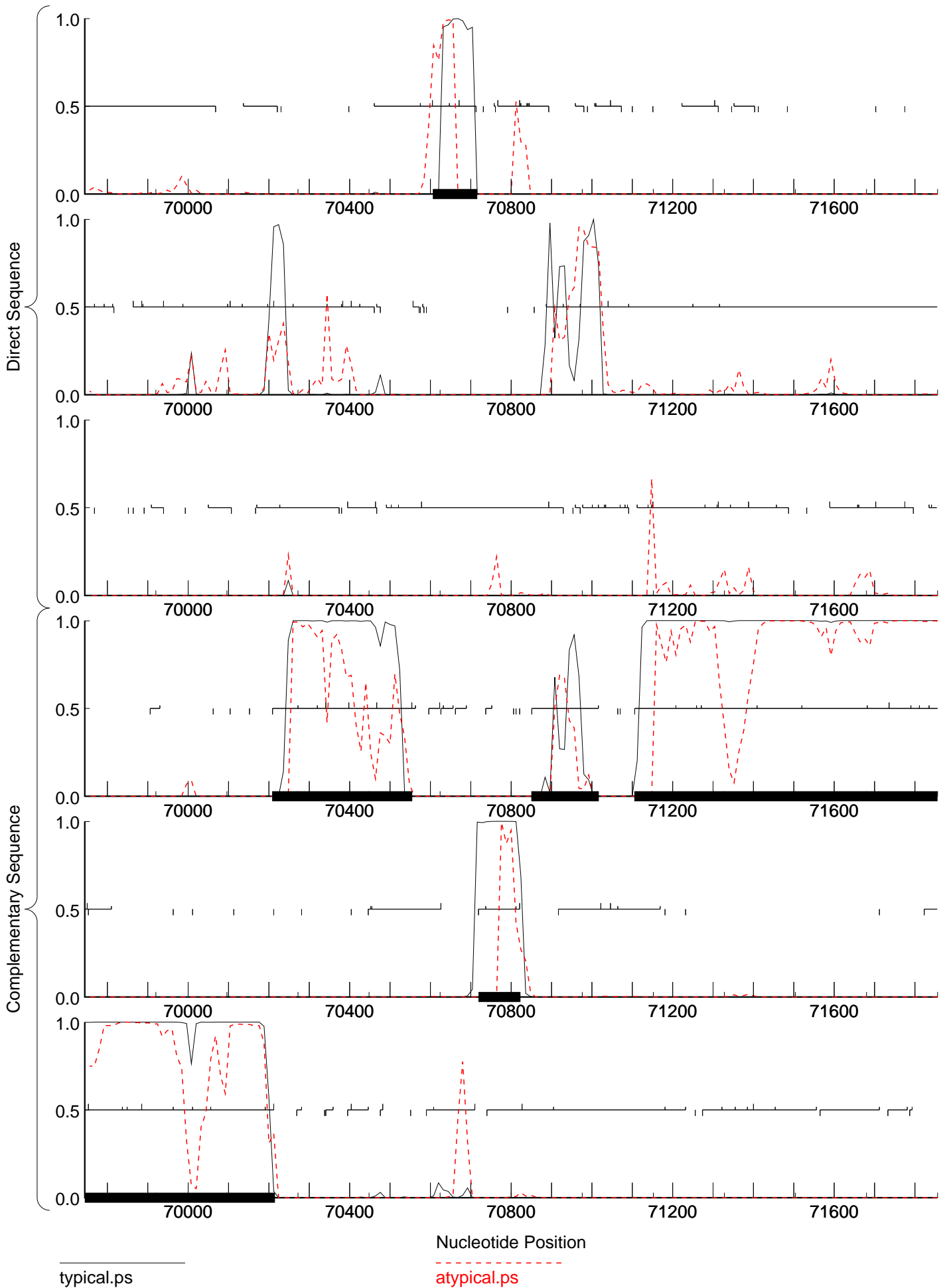
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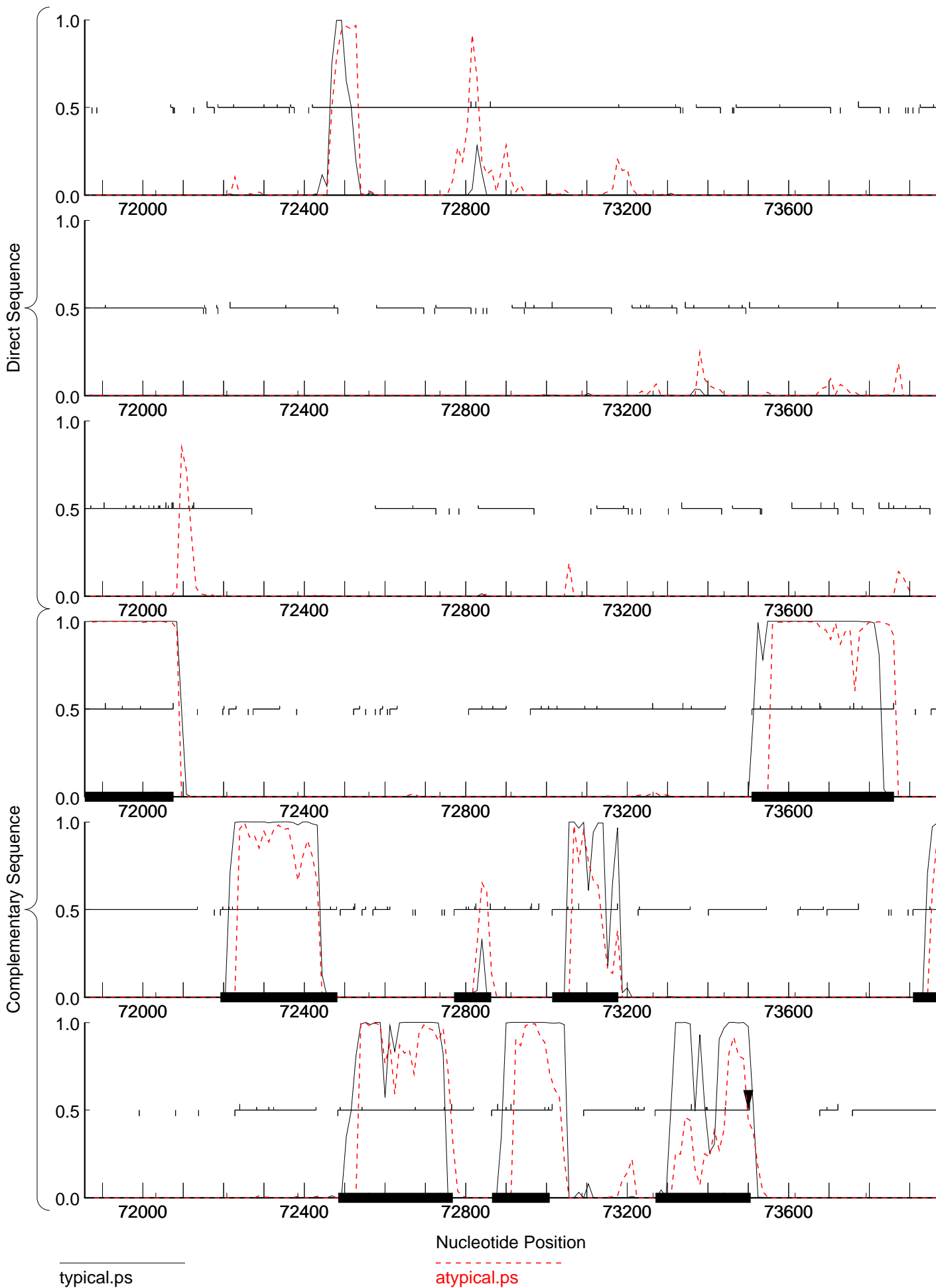
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typical.ps

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typical.ps

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