

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 : 10/20/17 at 13:49:17
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,

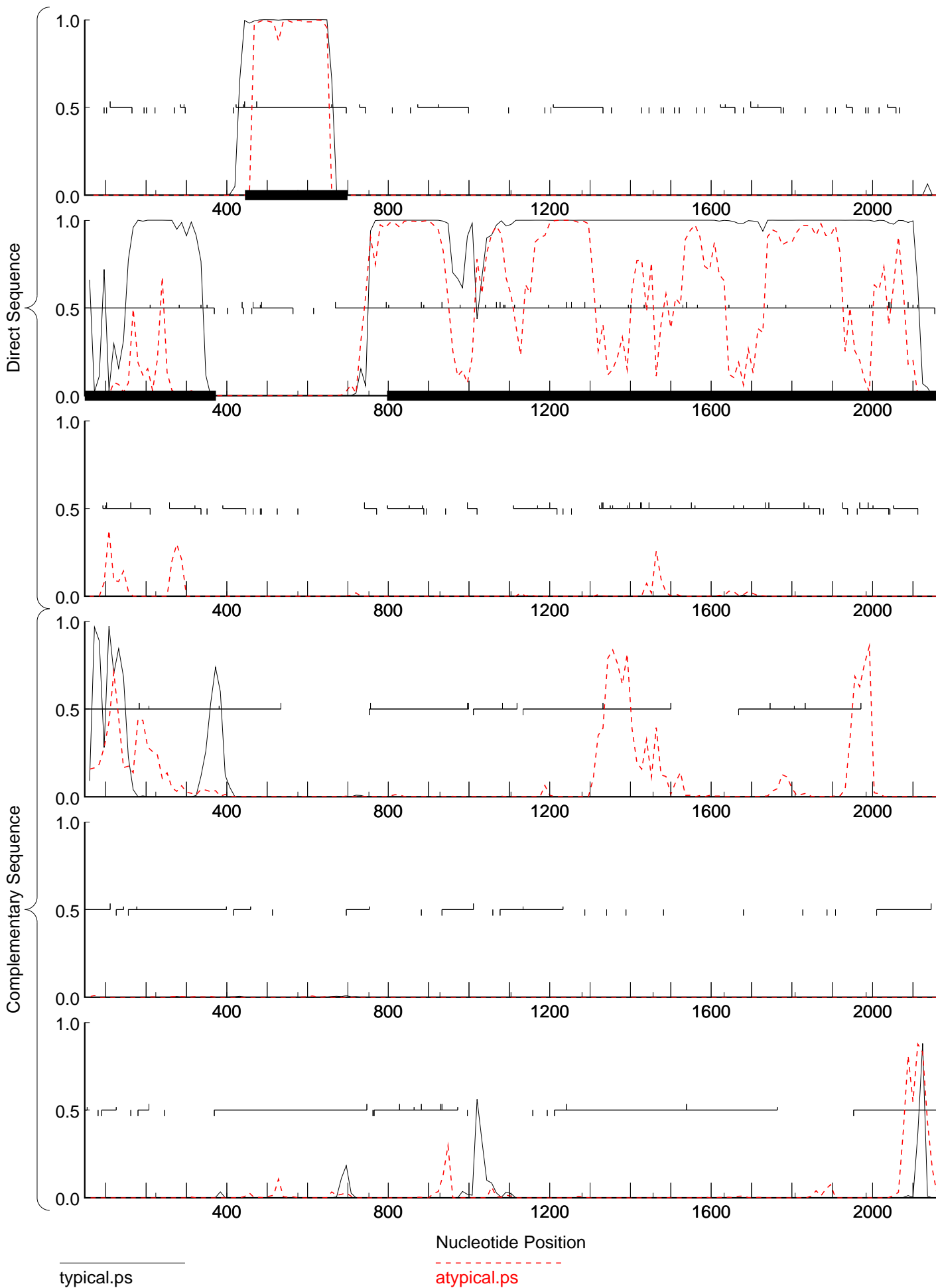
Matrix notes & comments

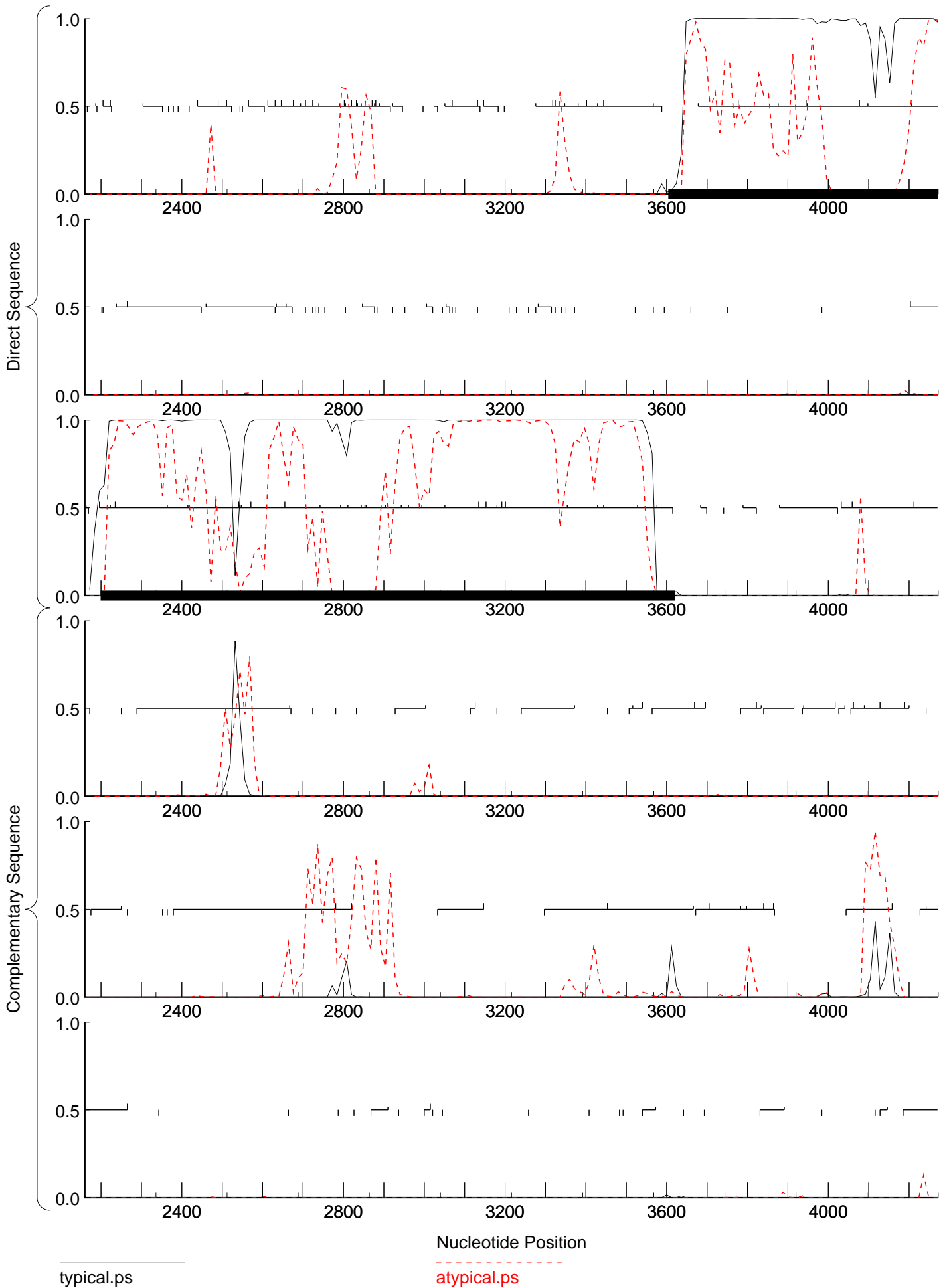
Training set derived by GeneMarkS, 4.30 October 2014
Fri Oct 20 13:49:17 2017

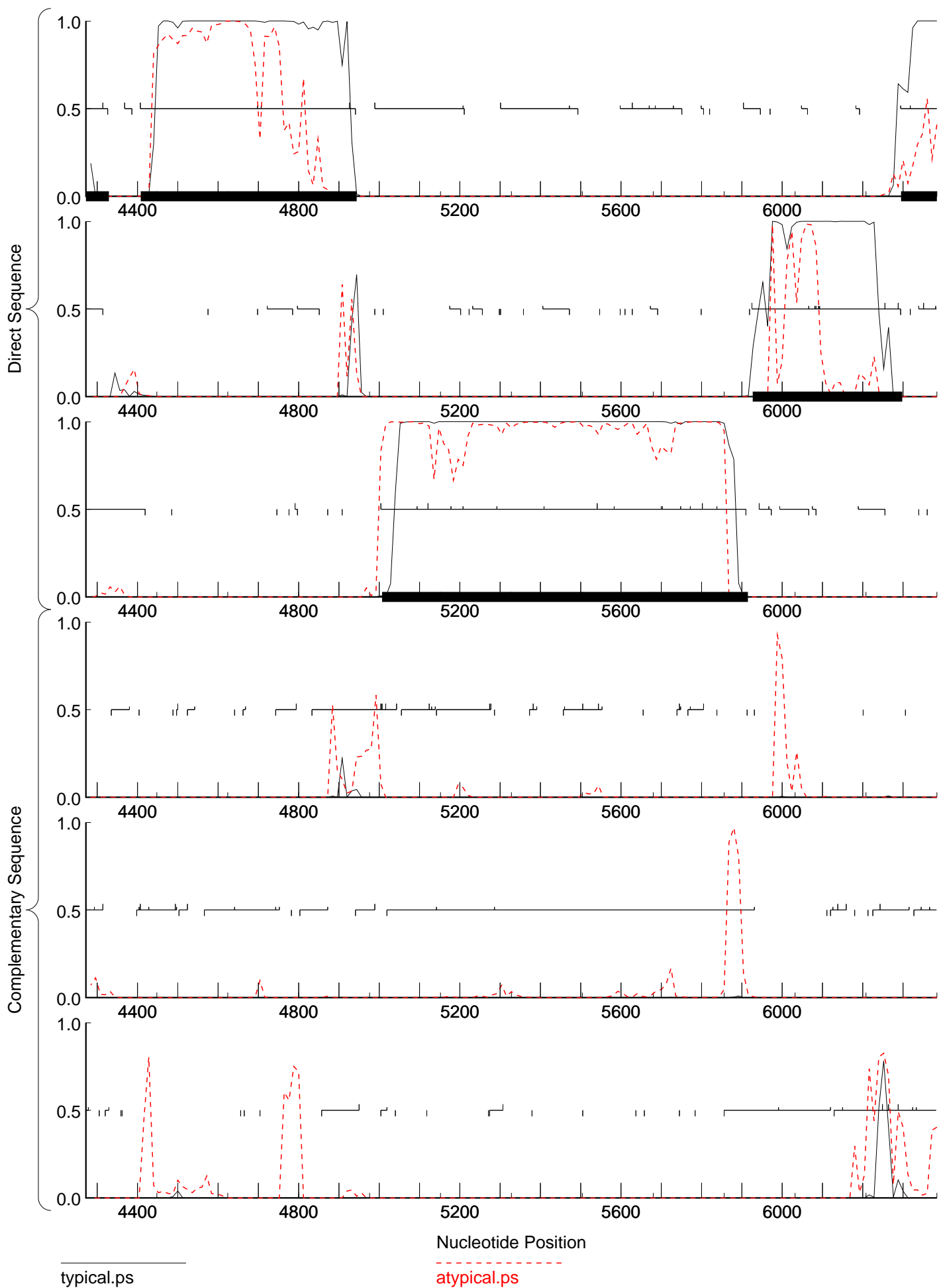
MATRIX INFORMATION

Matrix : -
Author : -
Order : 4

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

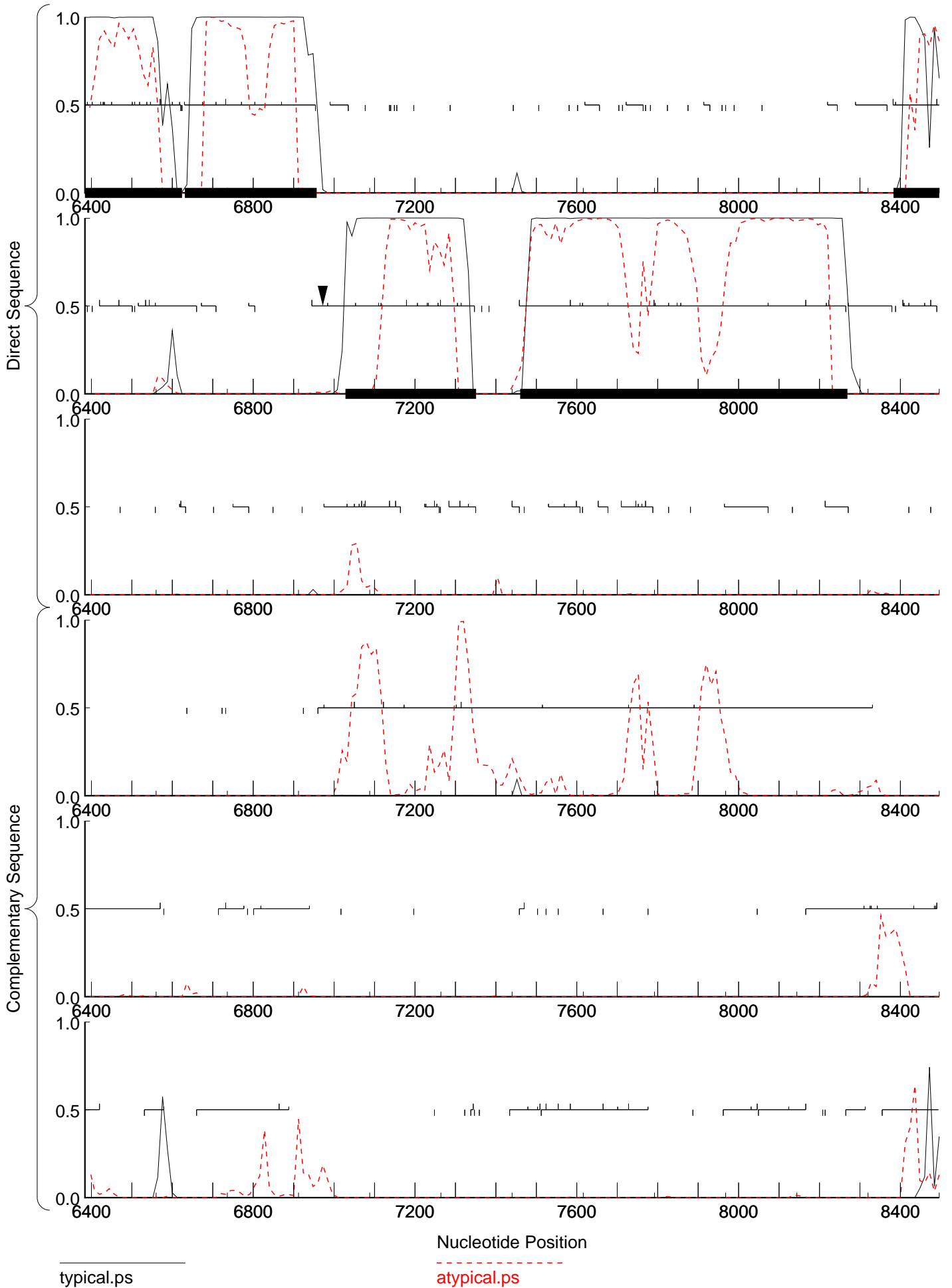


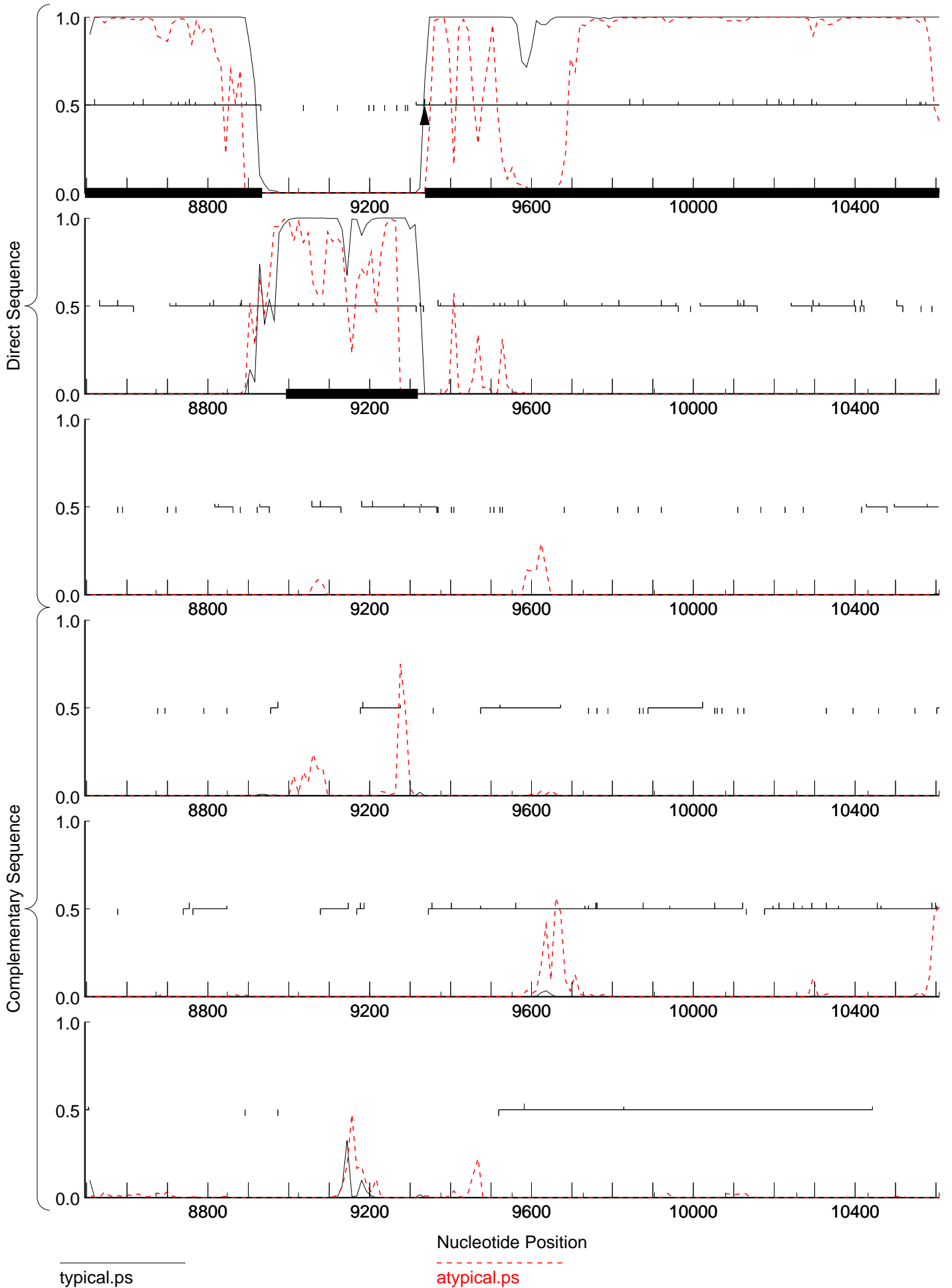




typical.ps

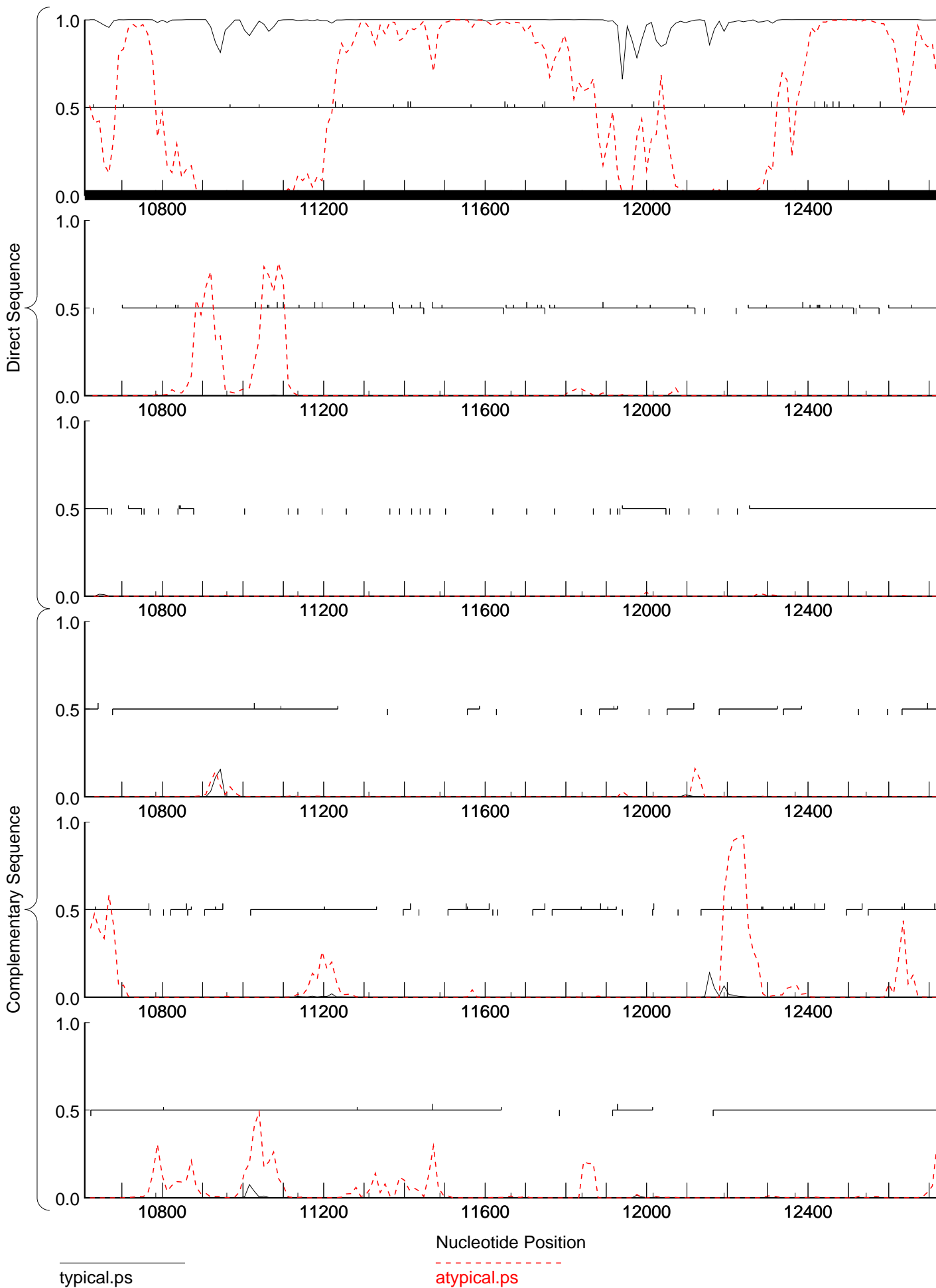
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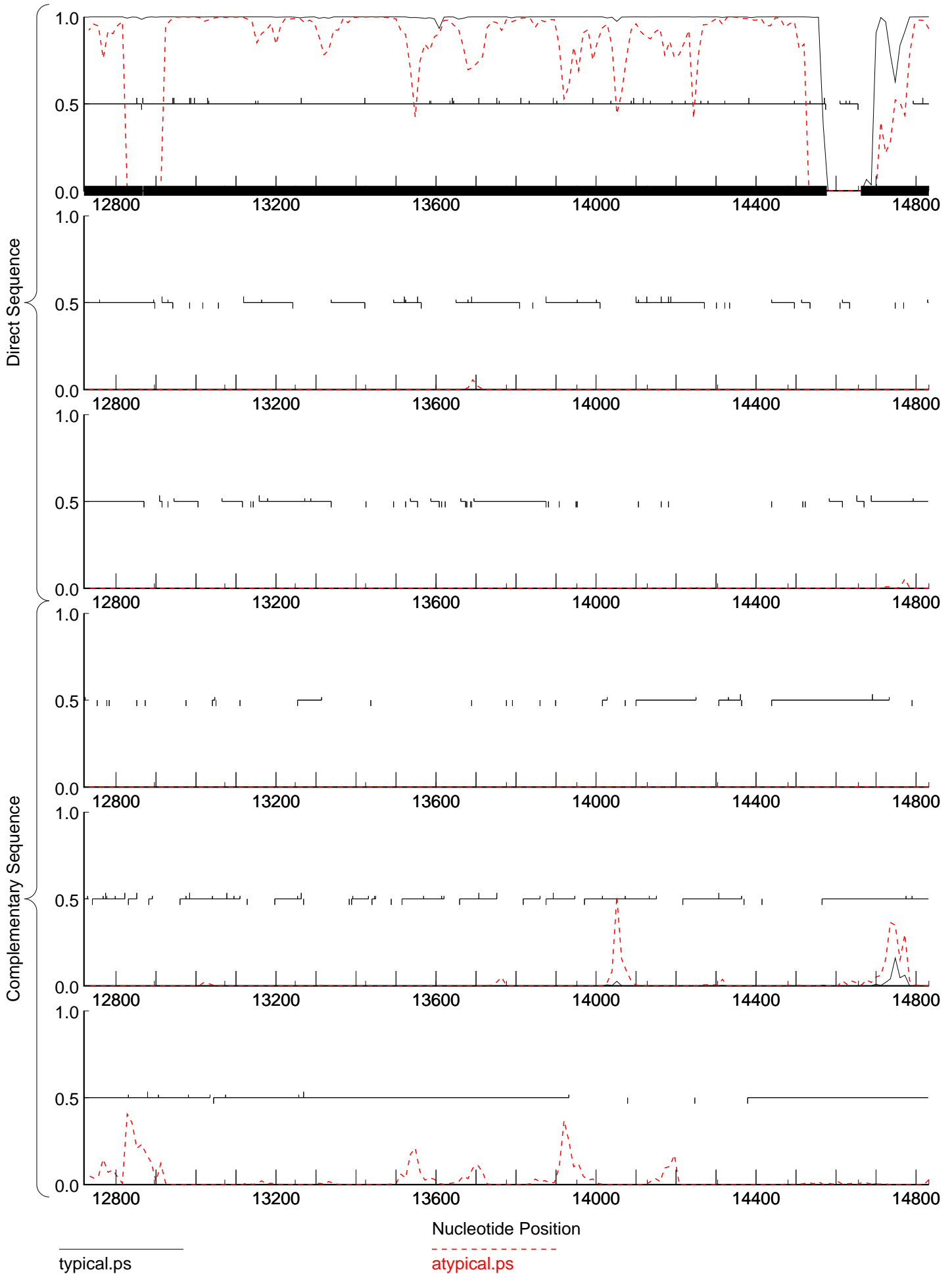




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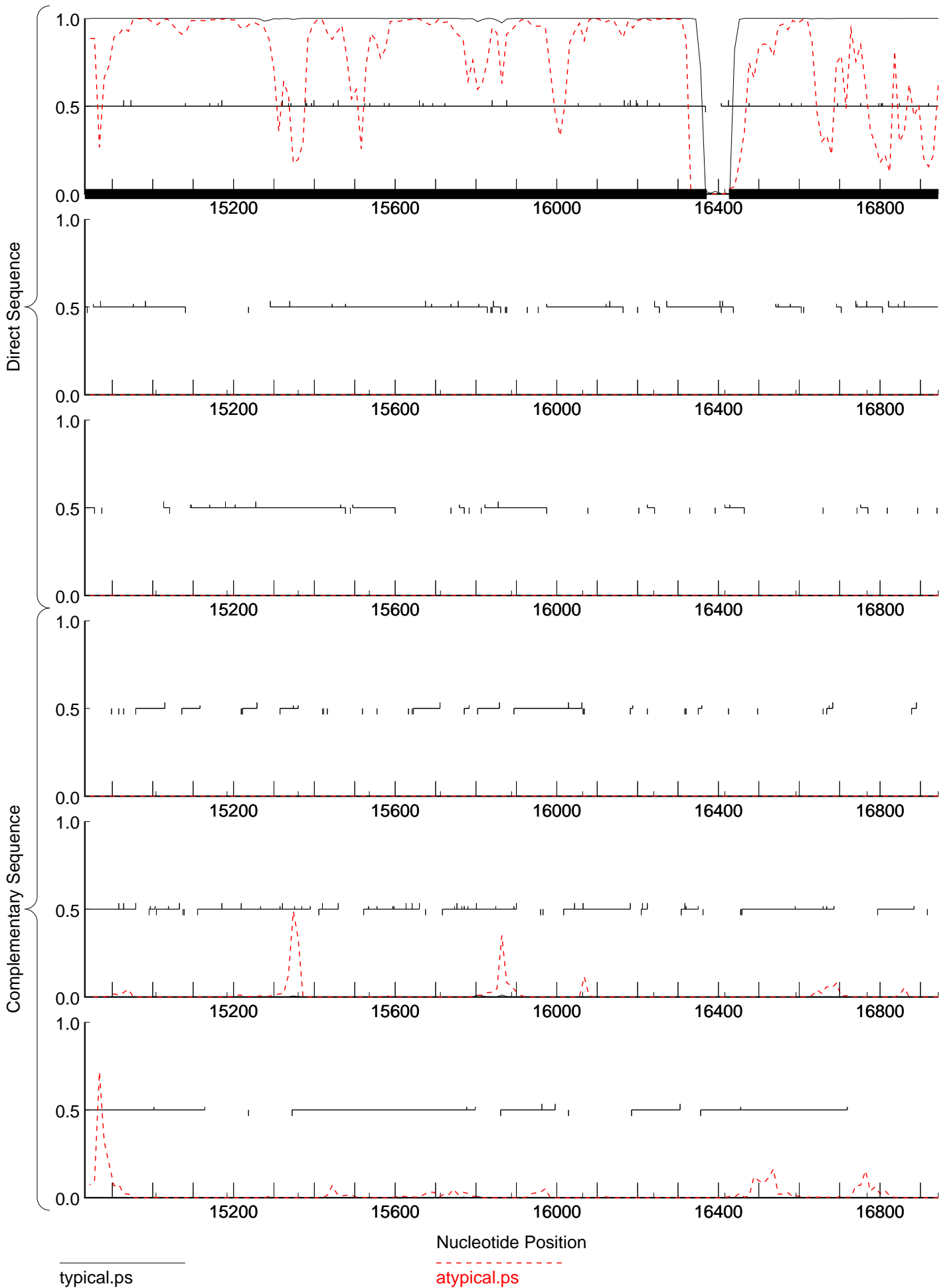
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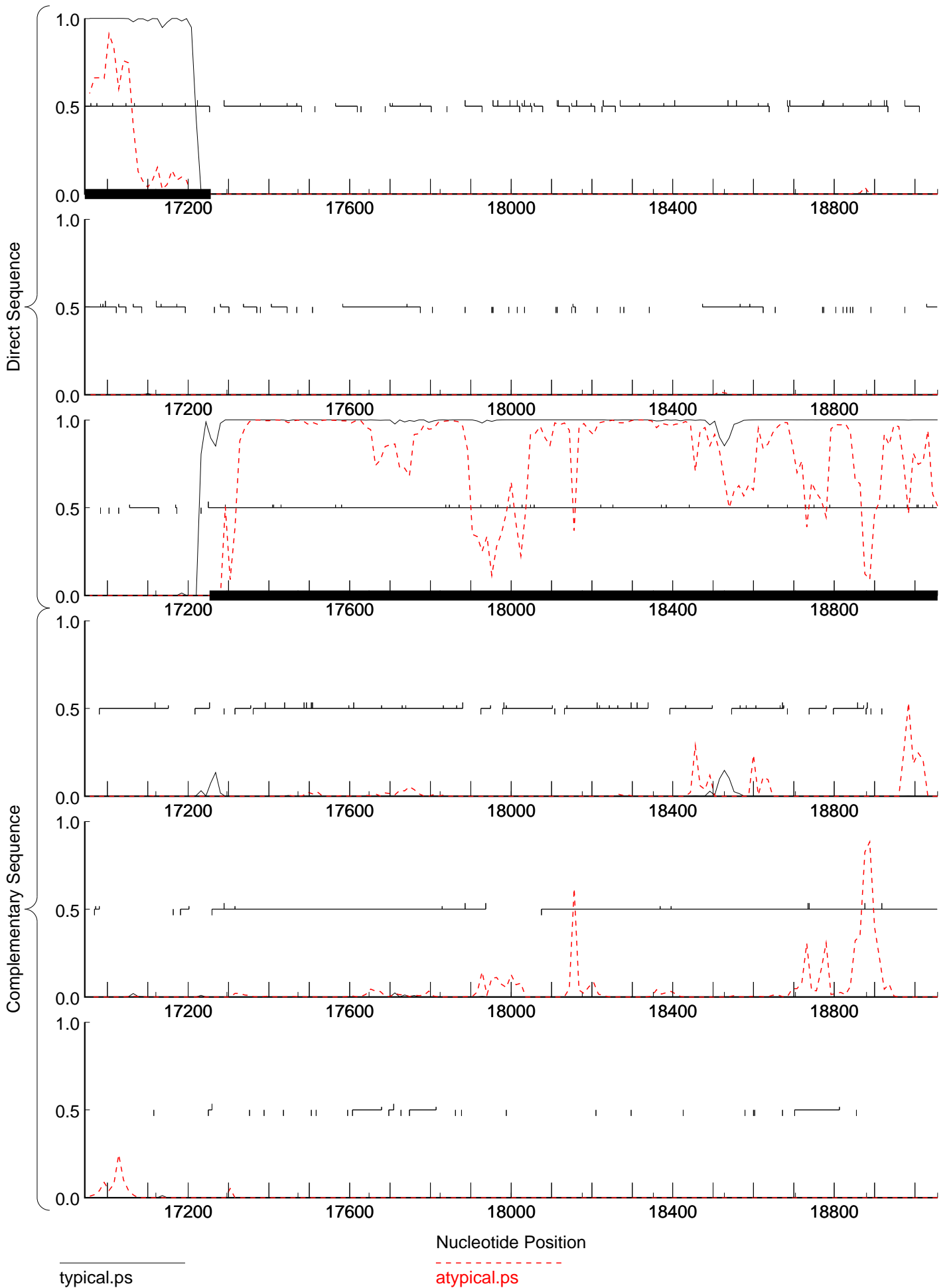


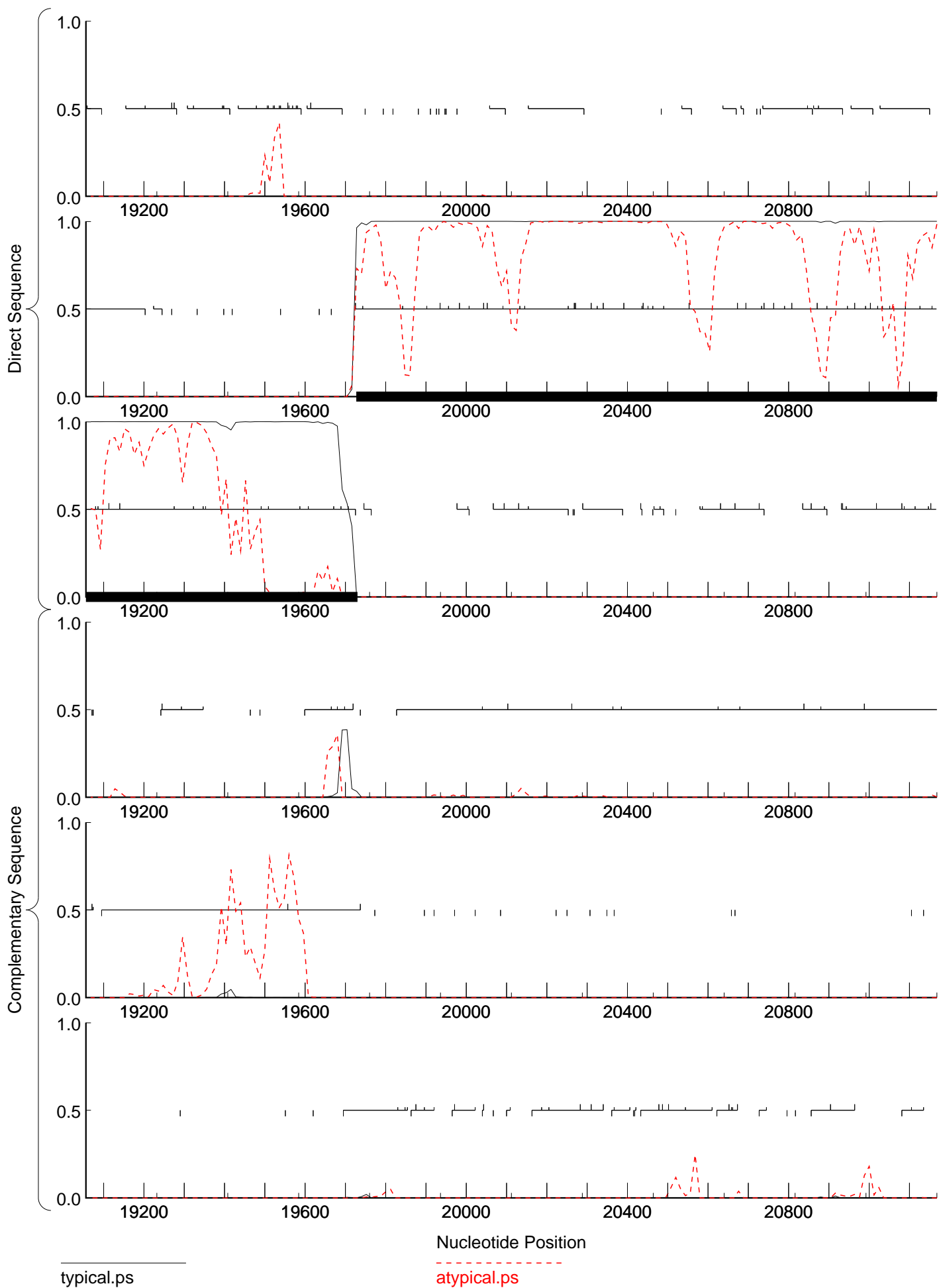


typical.ps

atypical.ps

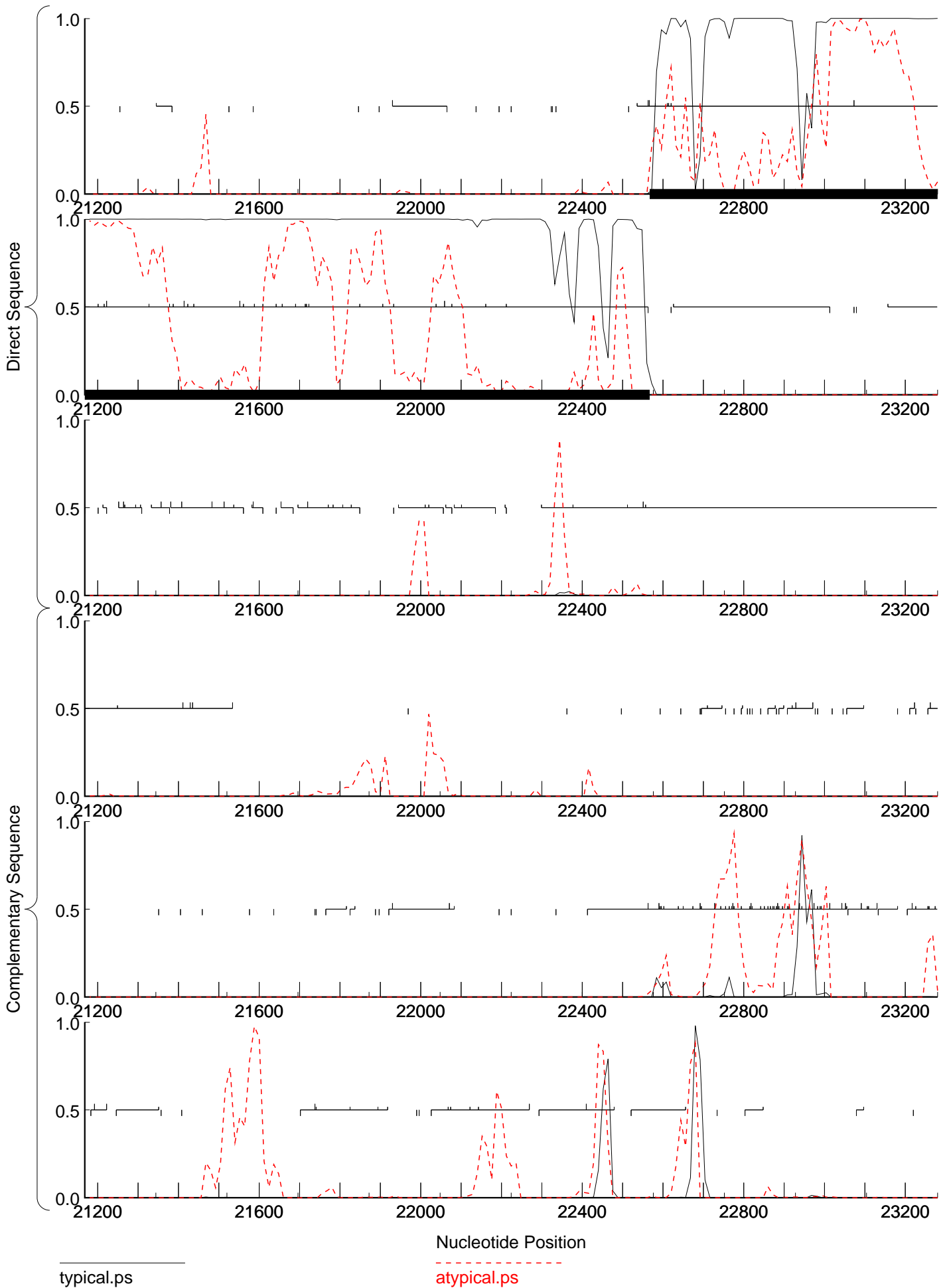


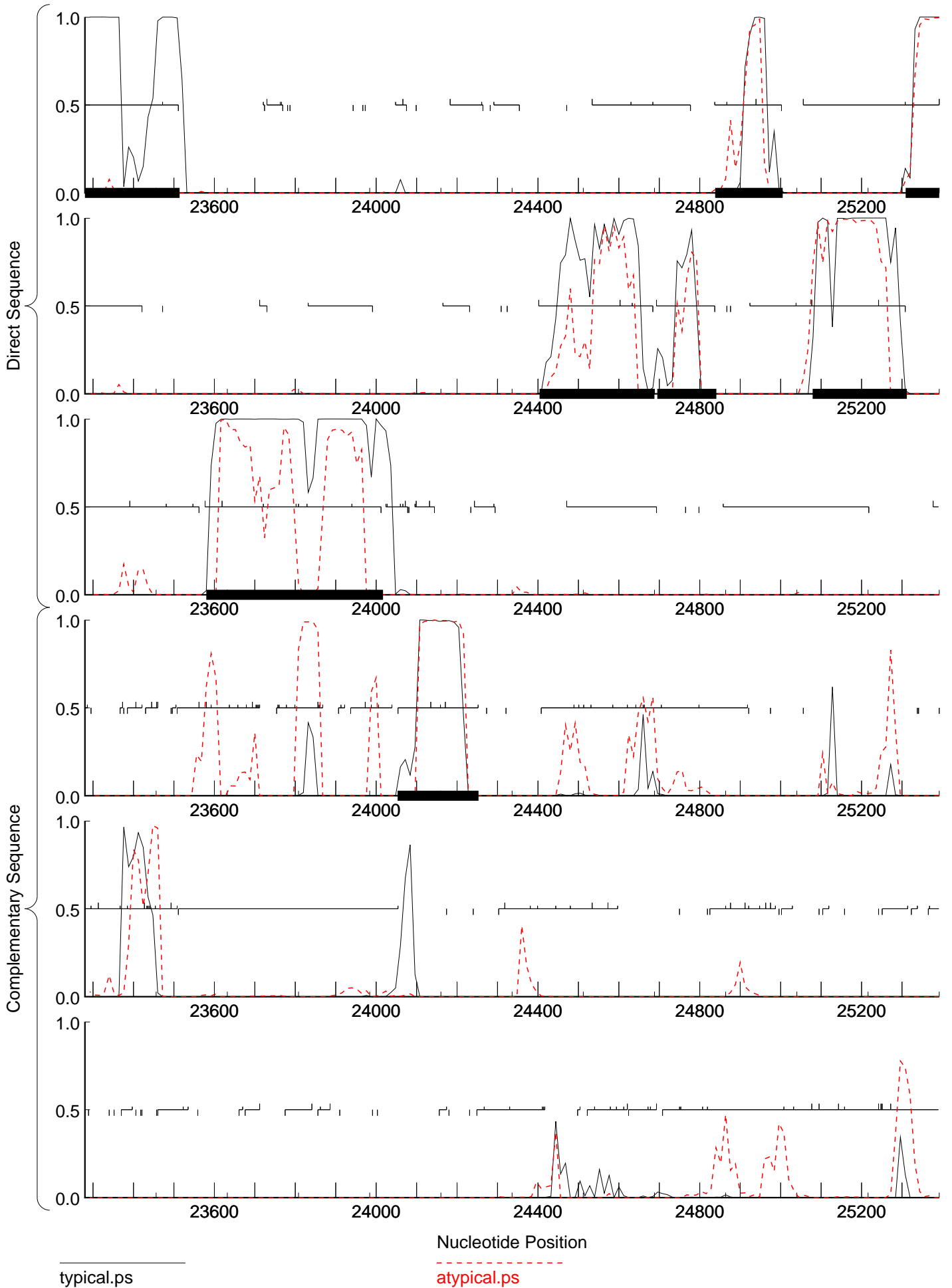


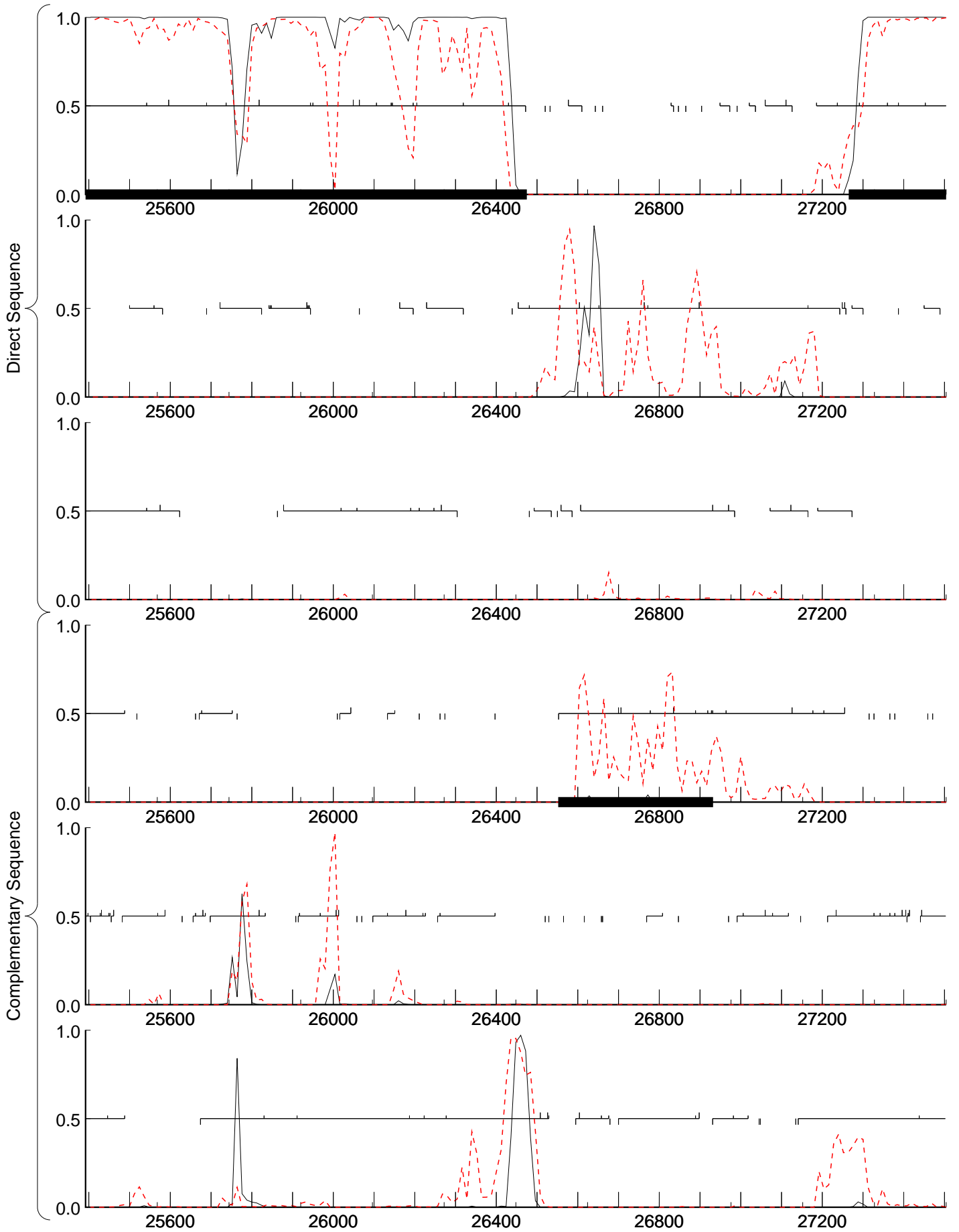


typical.ps

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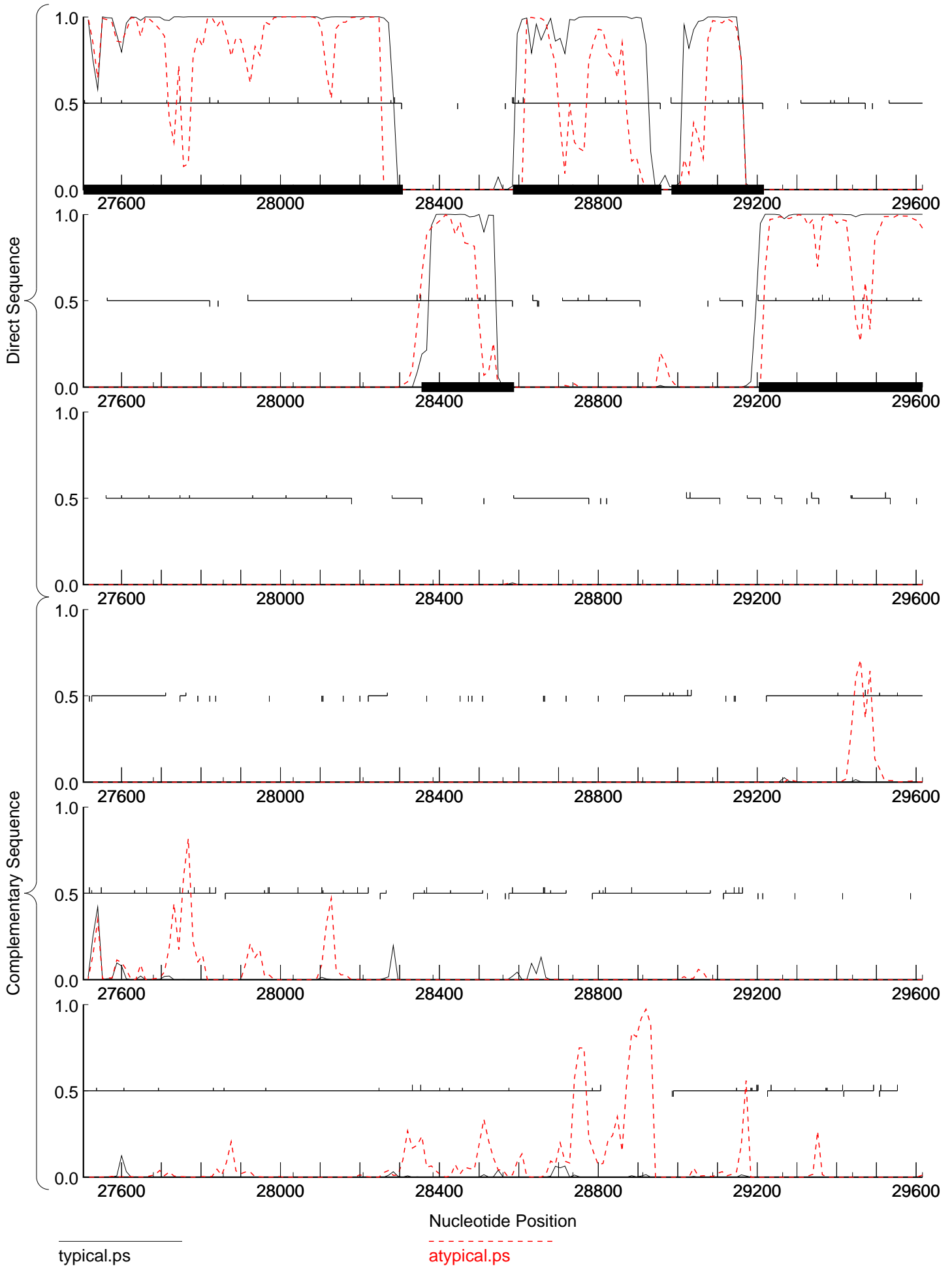


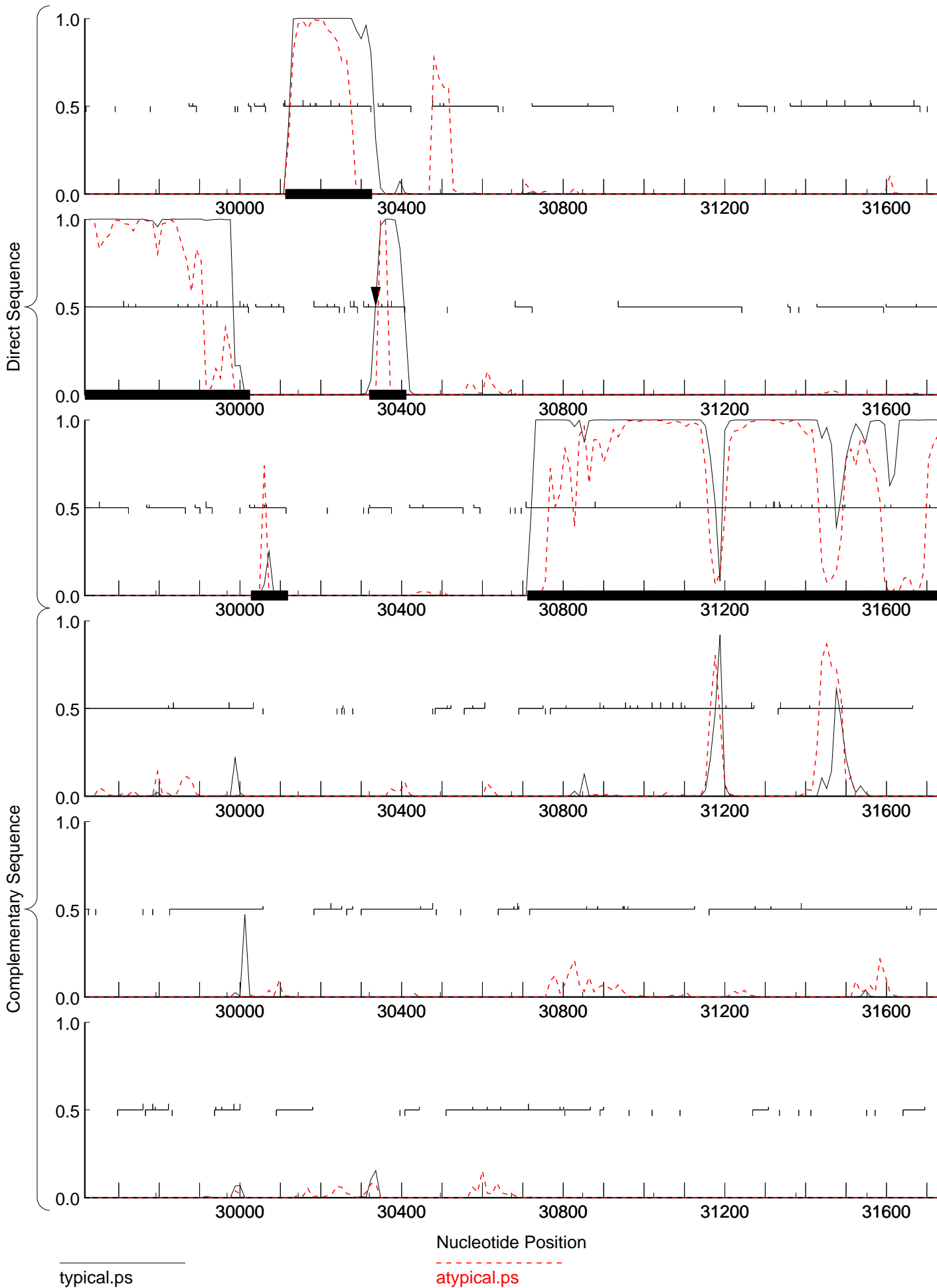


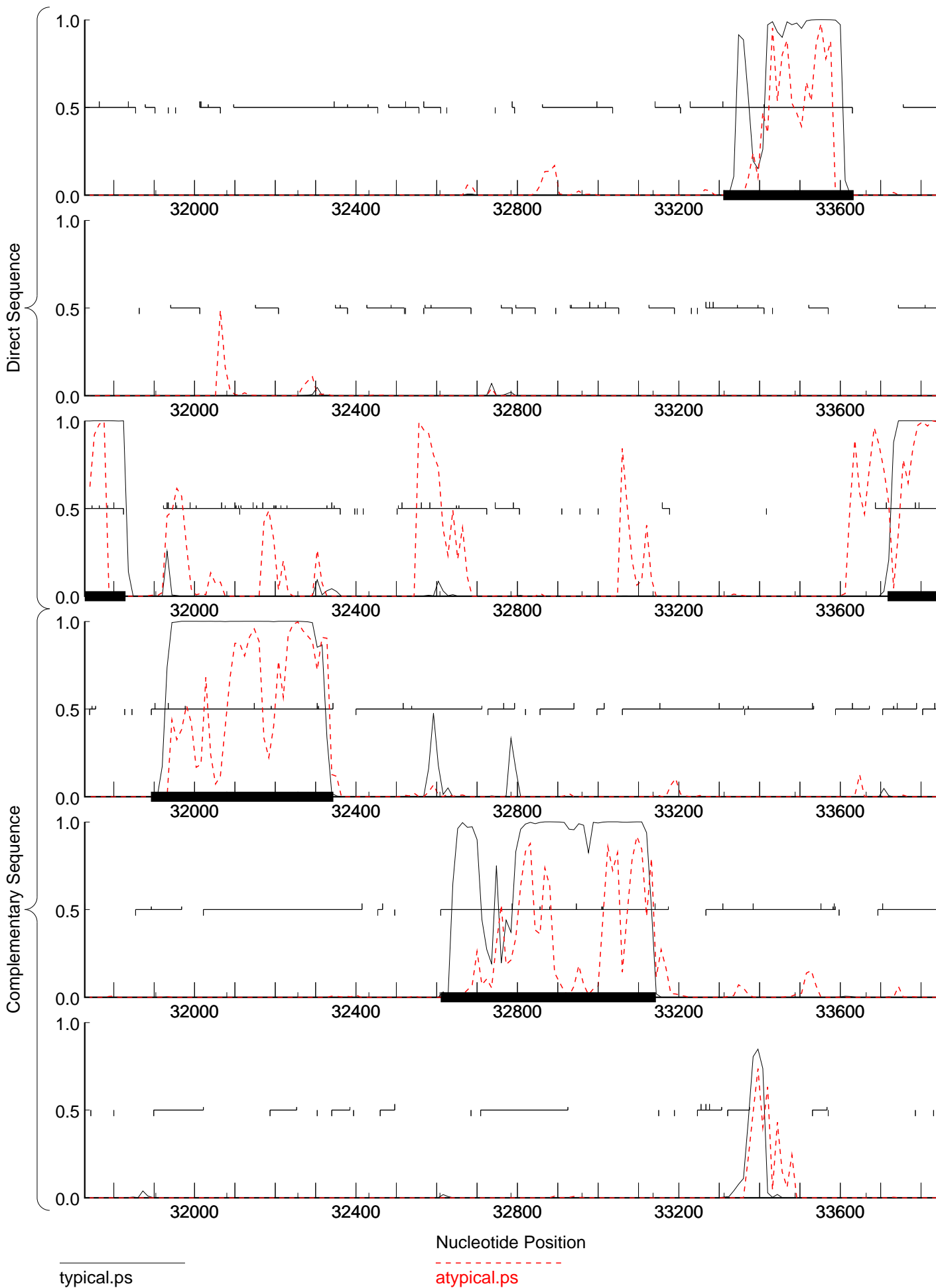


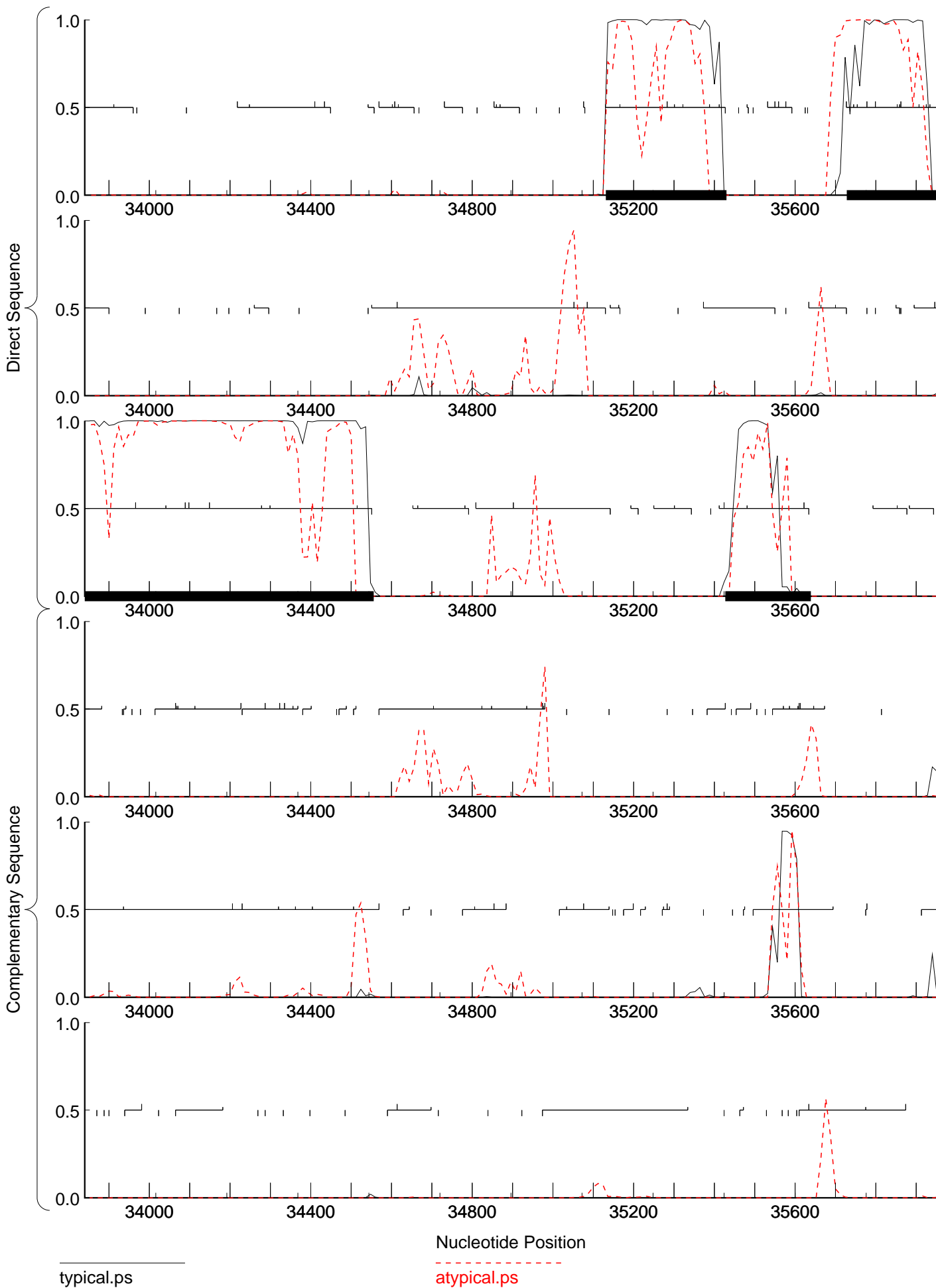
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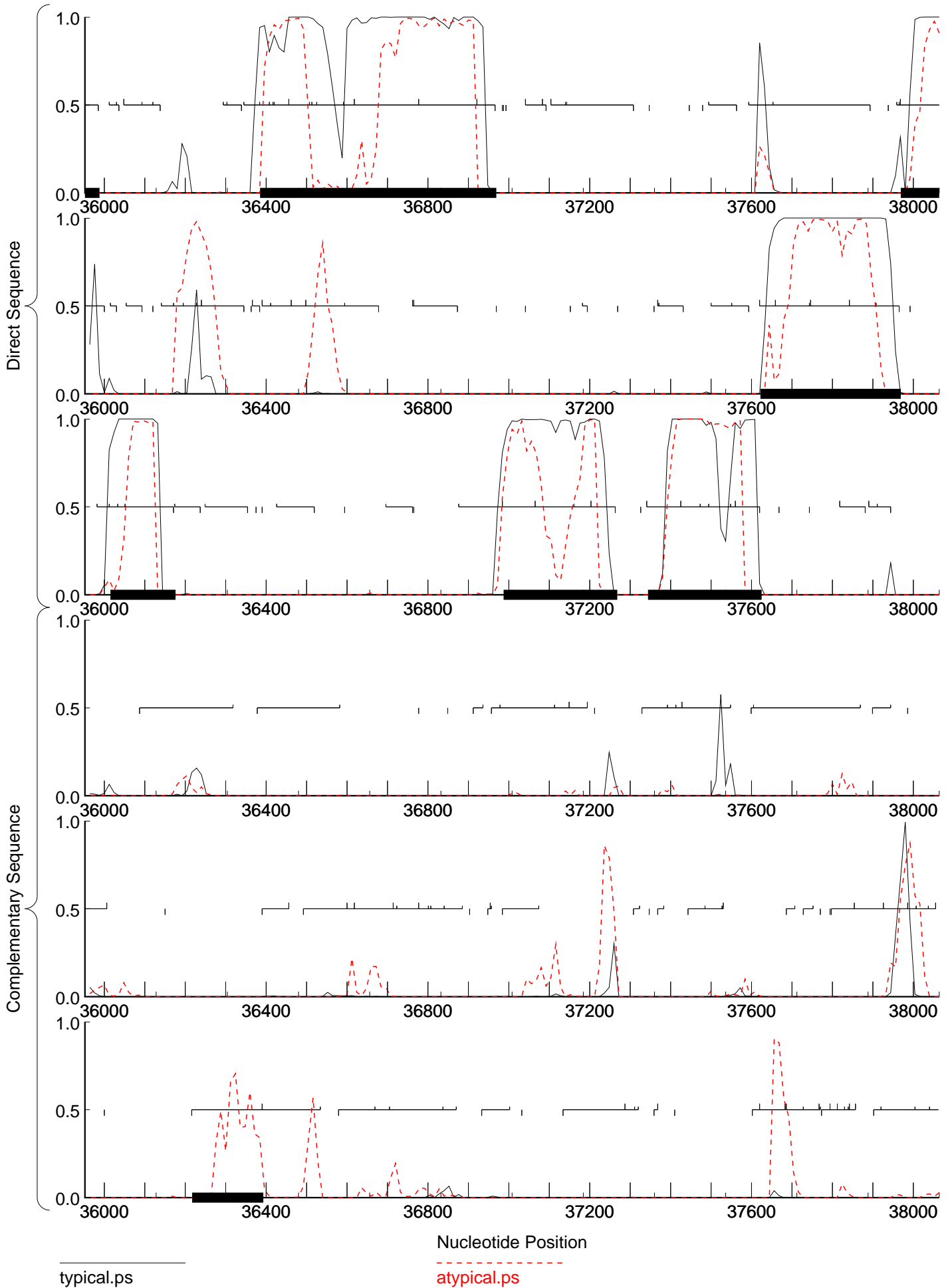
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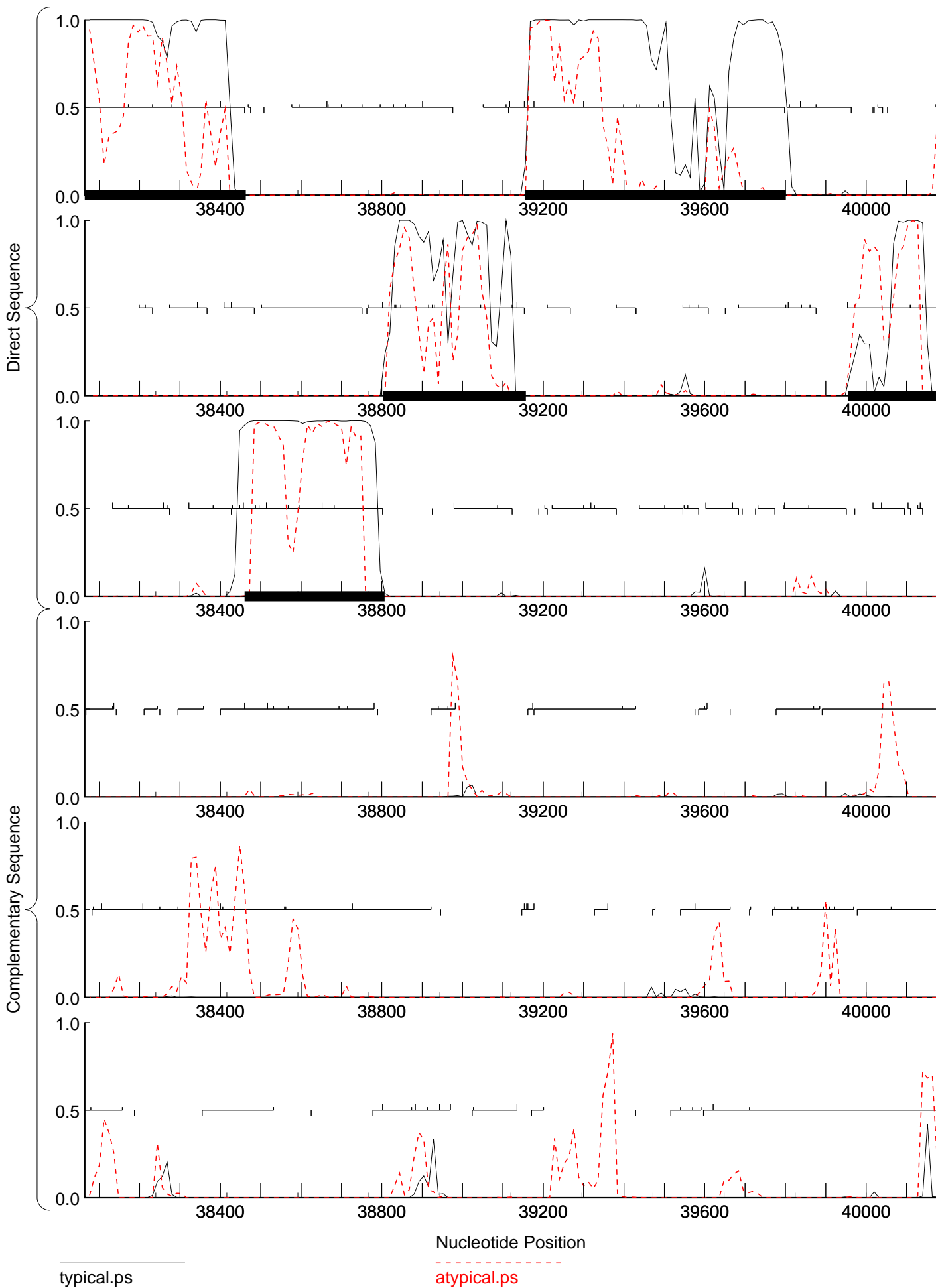






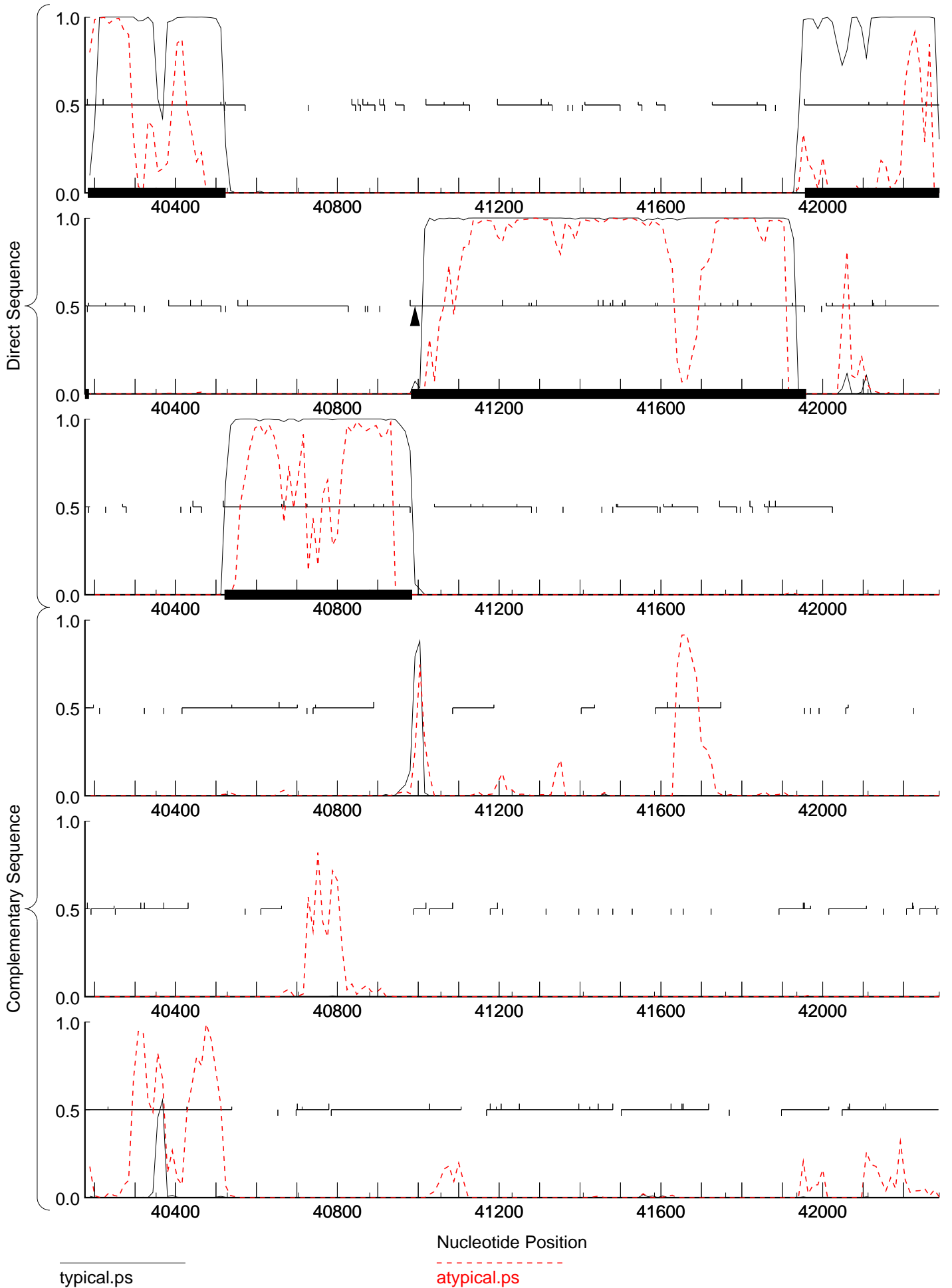


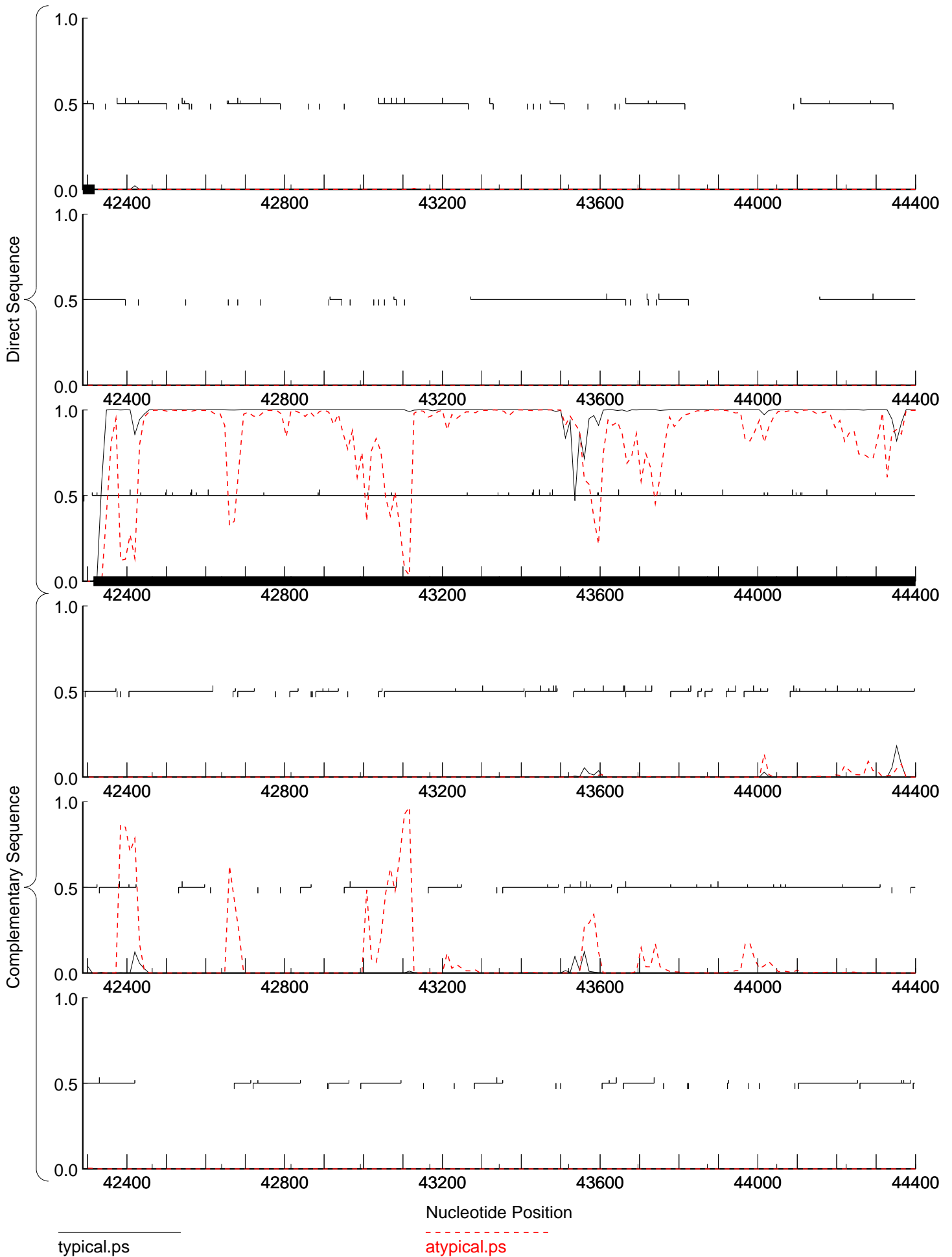


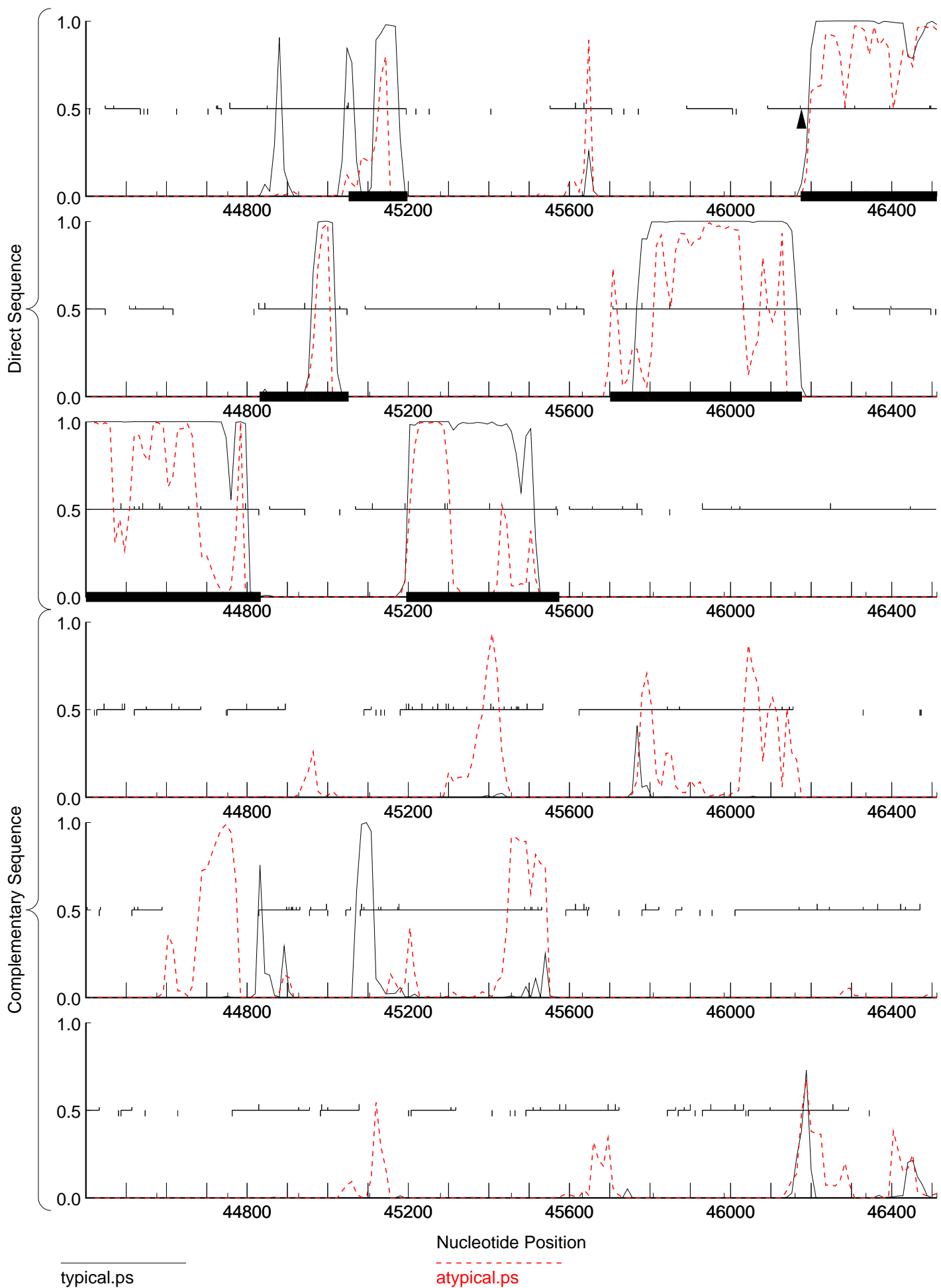


typical.ps

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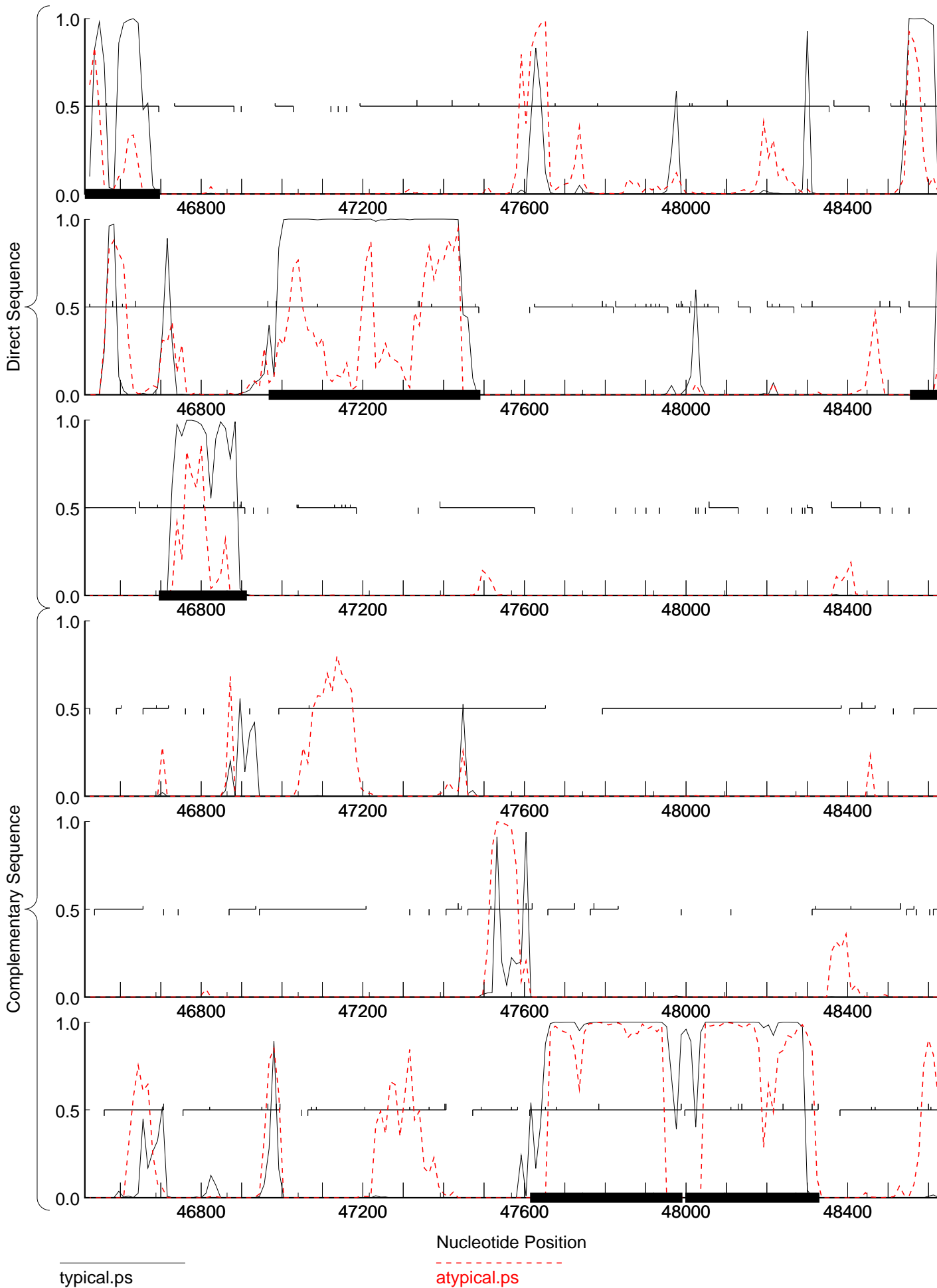






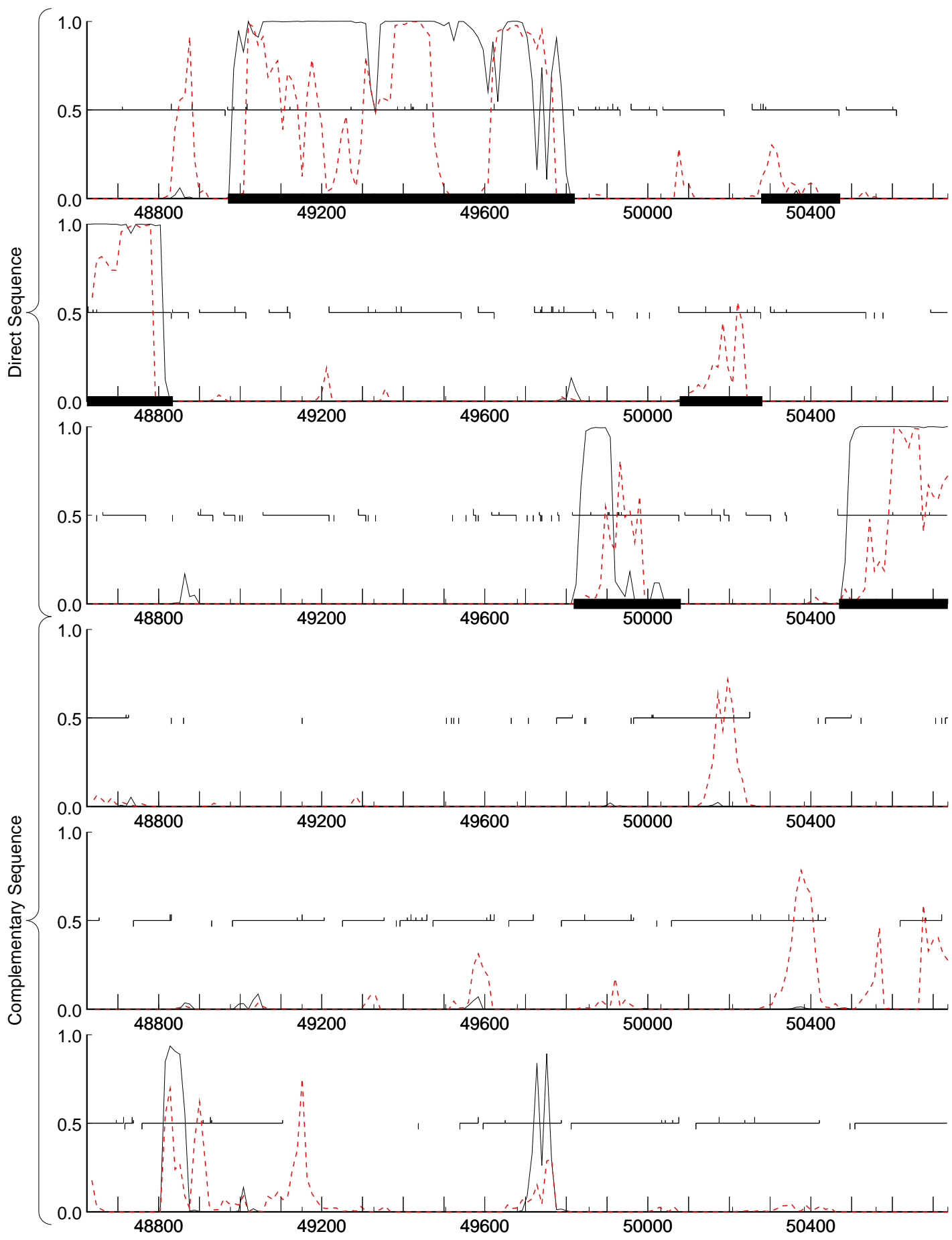
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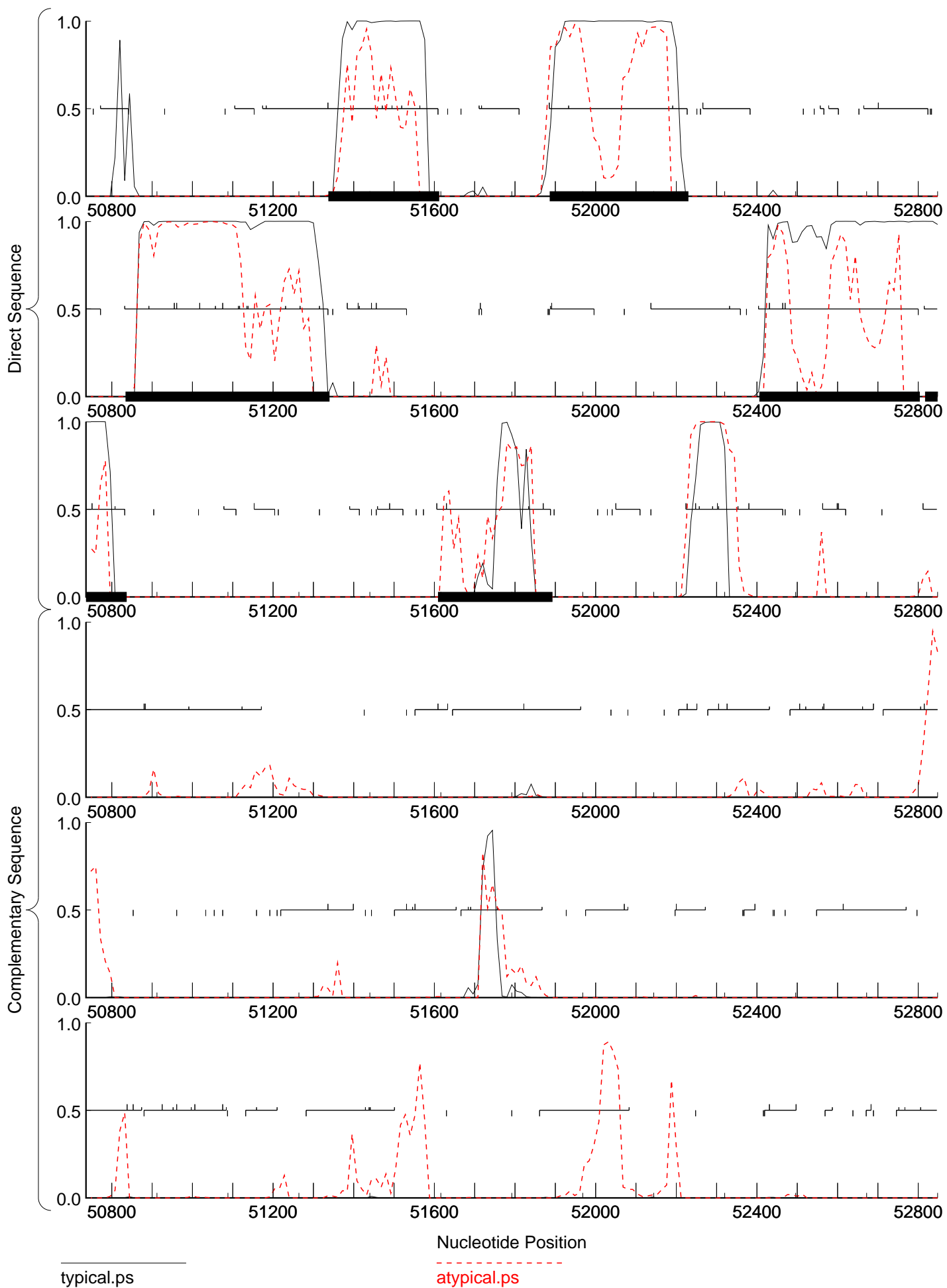
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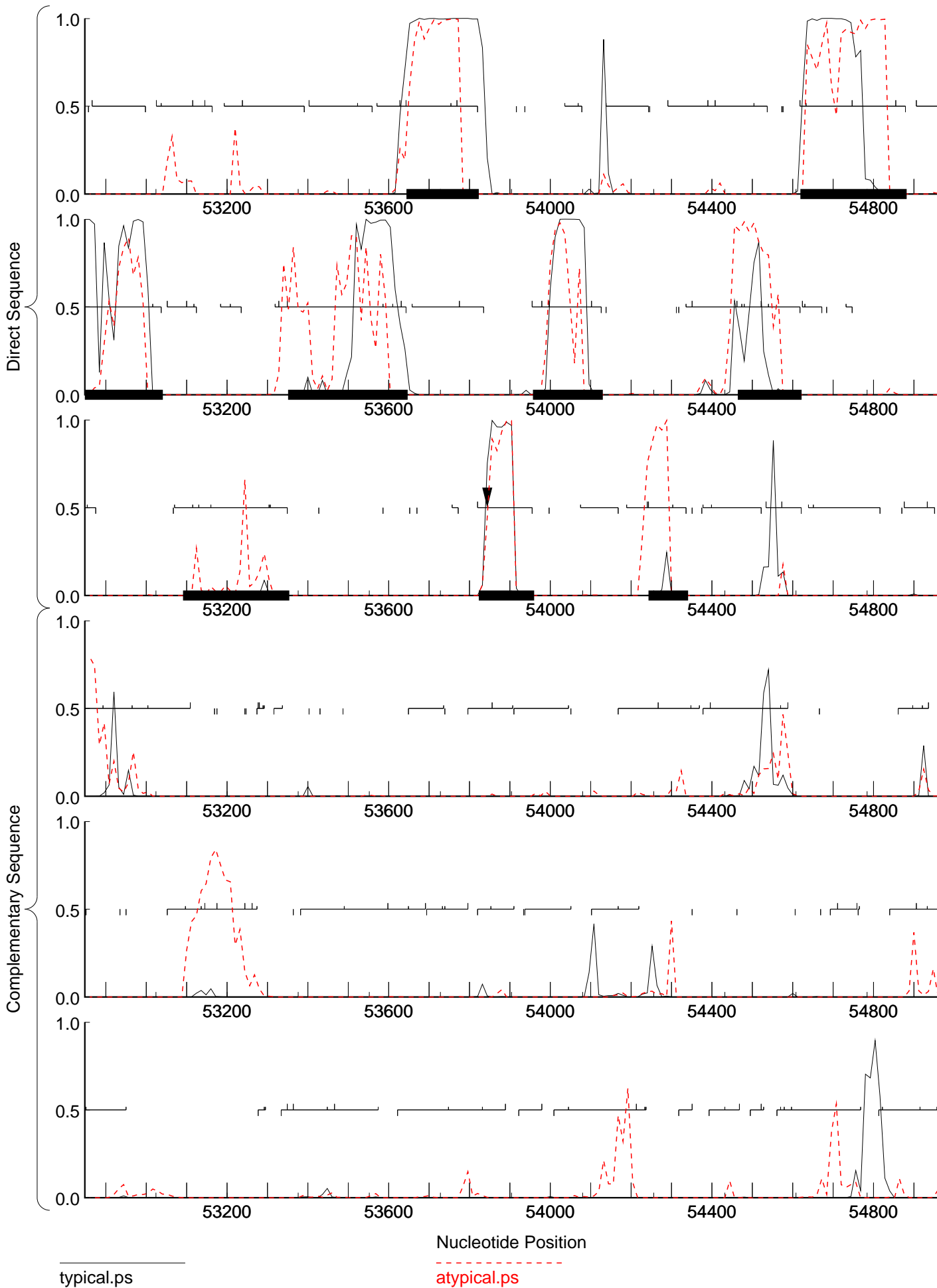
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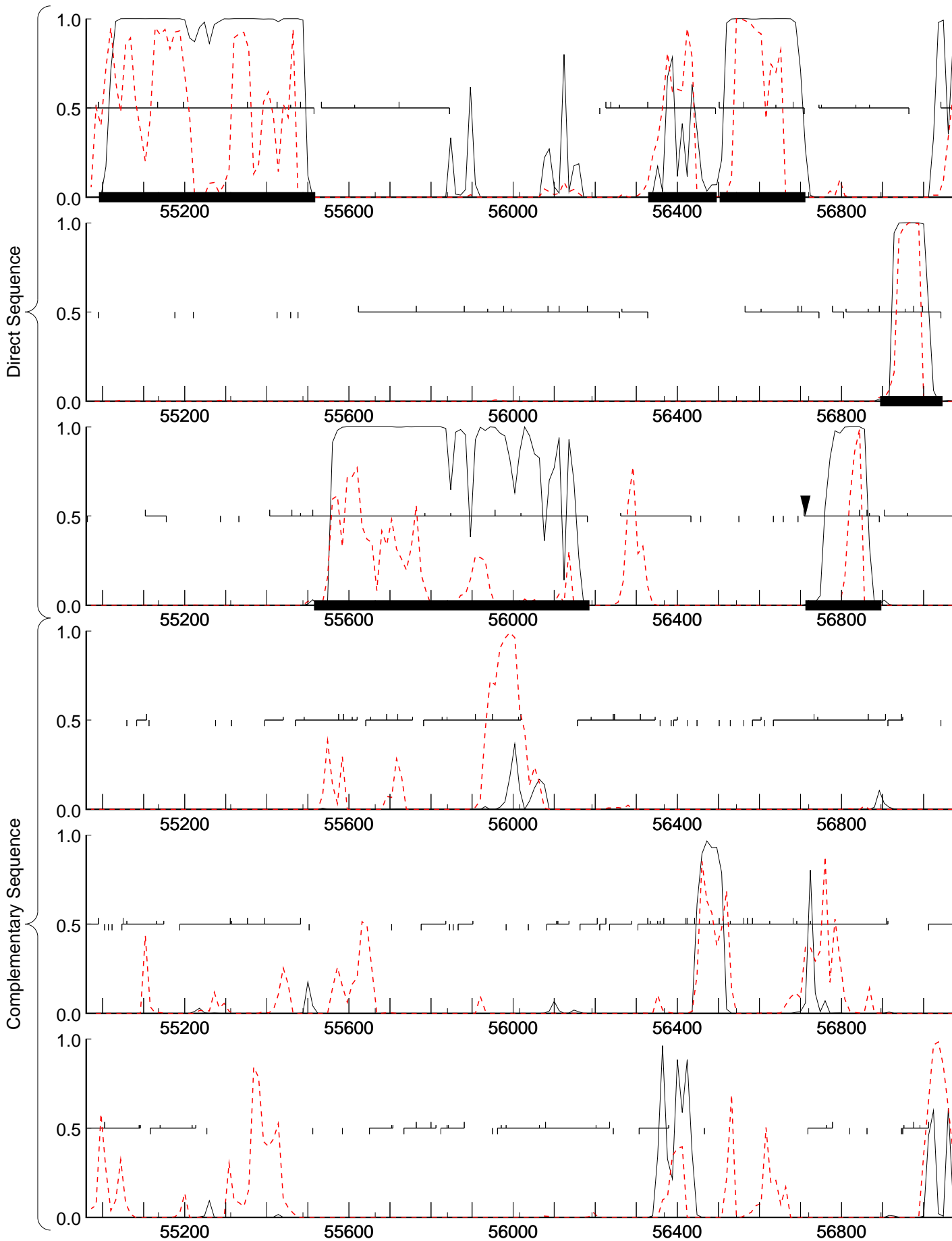


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