

GeneMark

Version 2.5p (09.08.06)

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PROGRAM INFORMATION

Sequence : Mycobacterium phage DoctorDiddles complete sequence, 75070 bp including 9-base 3' overhang
Analysis Date : 4/20/18 at 0:32:52
Pages : 37
Sequence Length : 75070 bp
GC Content : 62.81%

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,

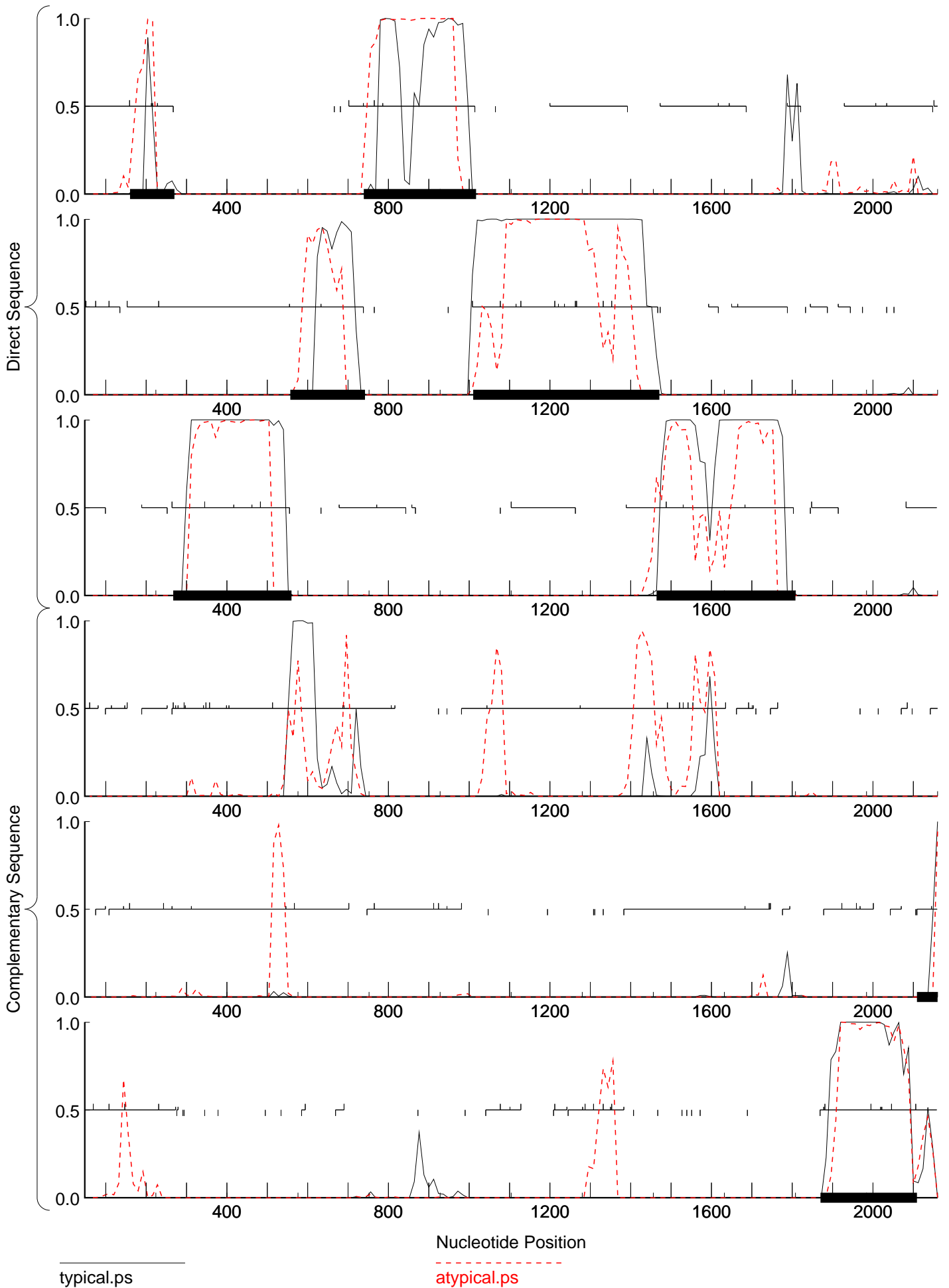
Matrix notes & comments

Training set derived by GeneMarkS, 4.30 October 2014
Fri Apr 20 00:32:52 2018

MATRIX INFORMATION

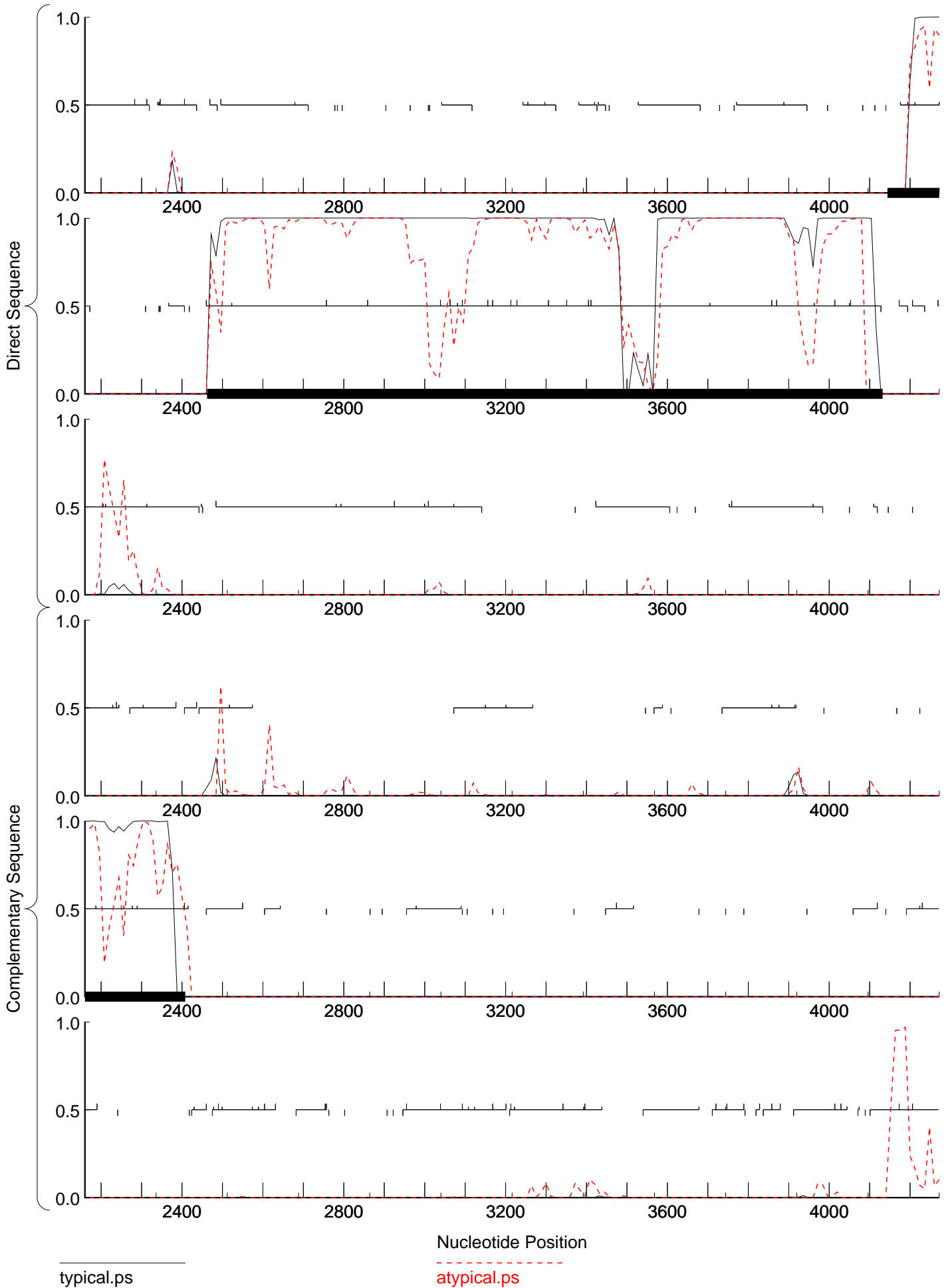
Matrix : -
Author : -
Order : 4

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

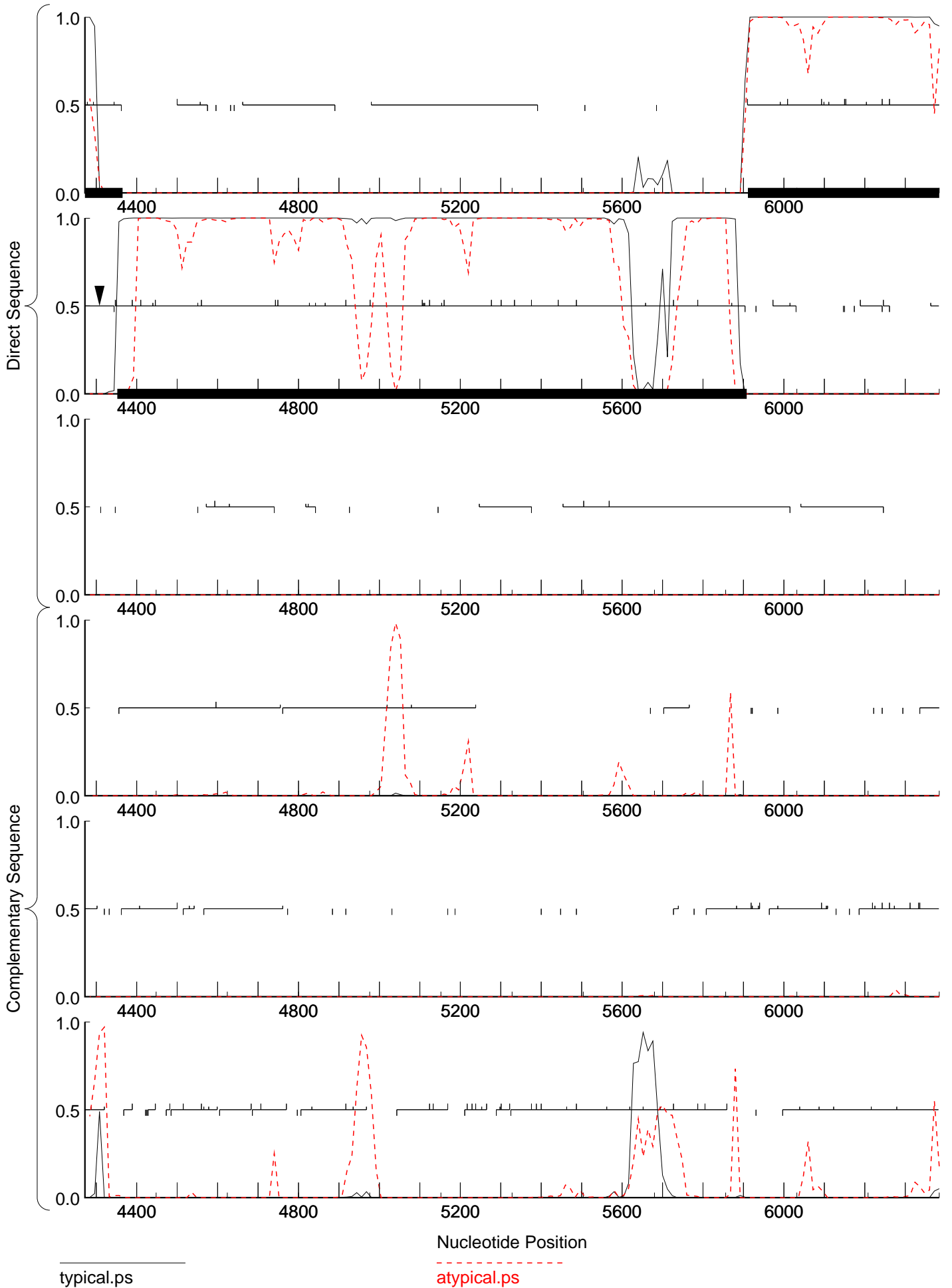


typical.ps

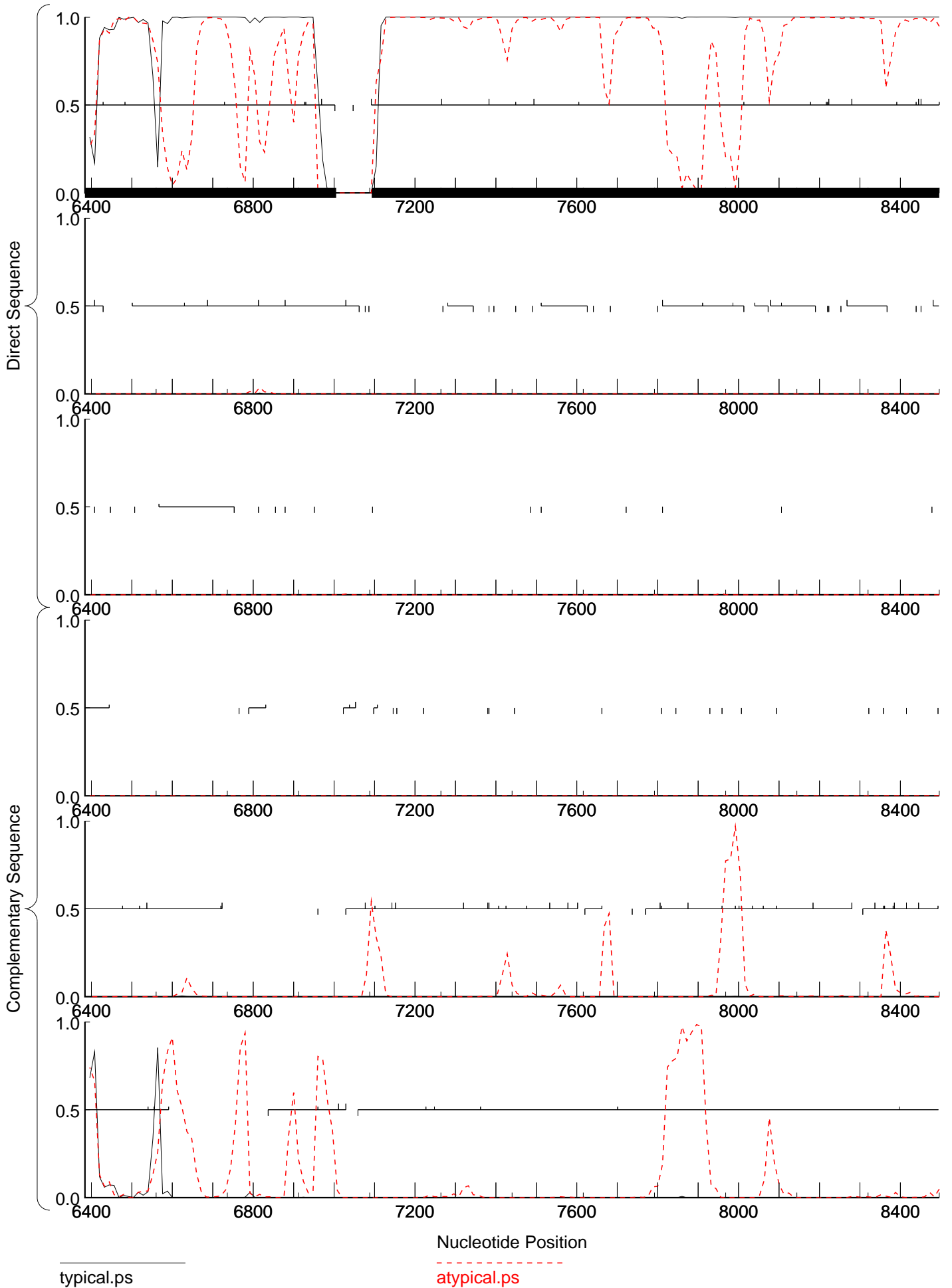
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...iddles complete sequence, 75070 bp including 9-base 3' overhang (CGCTTGTC), Cluster E, Order 4, Window 96, Step 12, 4/37
...iddles complete sequence, 75070 bp including 9-base 3' overhang (CGCTTGTC), Cluster E, Order 2, Window 96, Step 12, 4/37

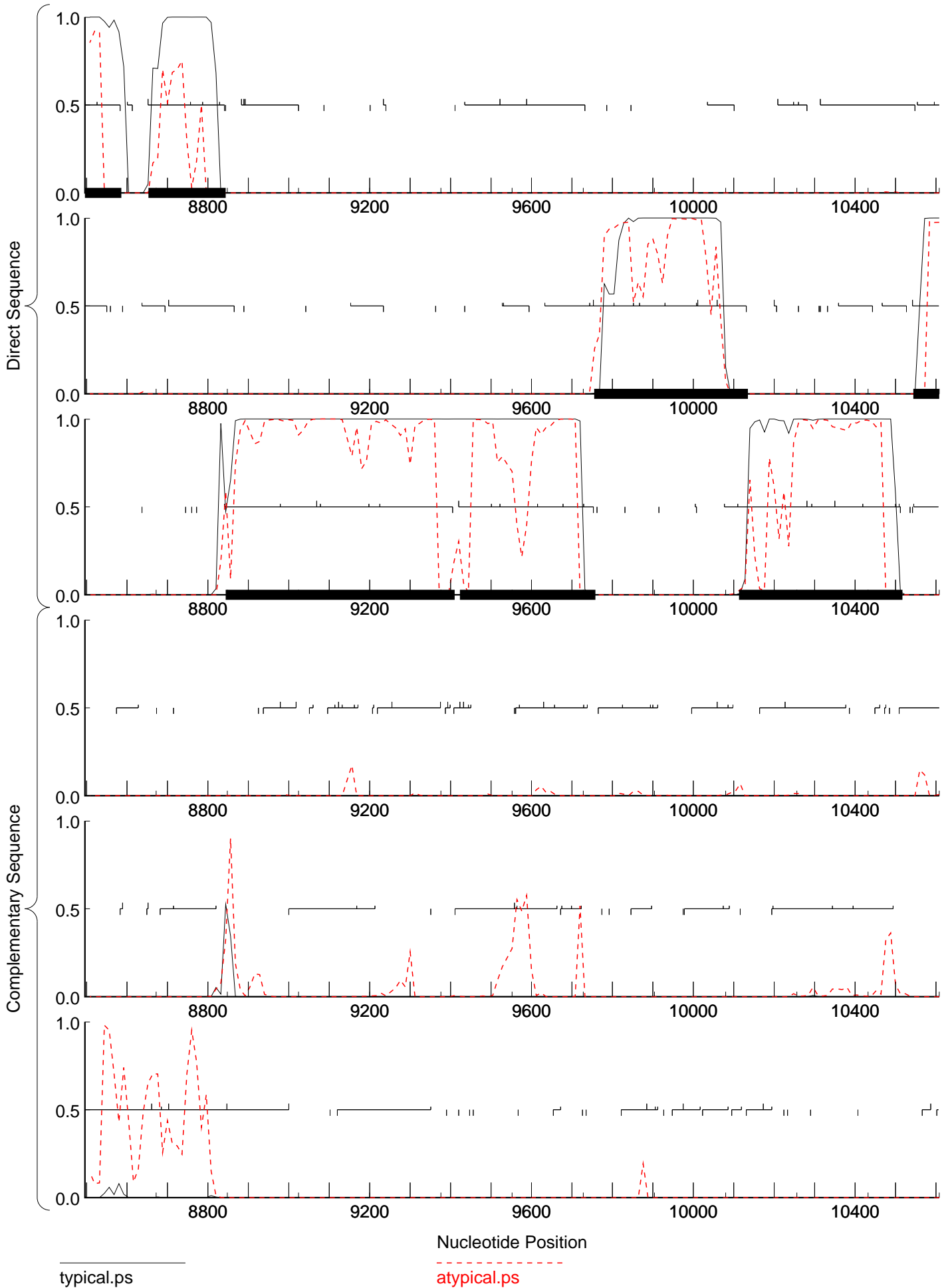


Genes complete sequence, 75070 bp including 9-base 3' overhang (CGCTTGTC), Cluster E, Order 4, Window 96, Step 12, 5/37
Genes complete sequence, 75070 bp including 9-base 3' overhang (CGCTTGTC), Cluster E, Order 2, Window 96, Step 12, 5/37



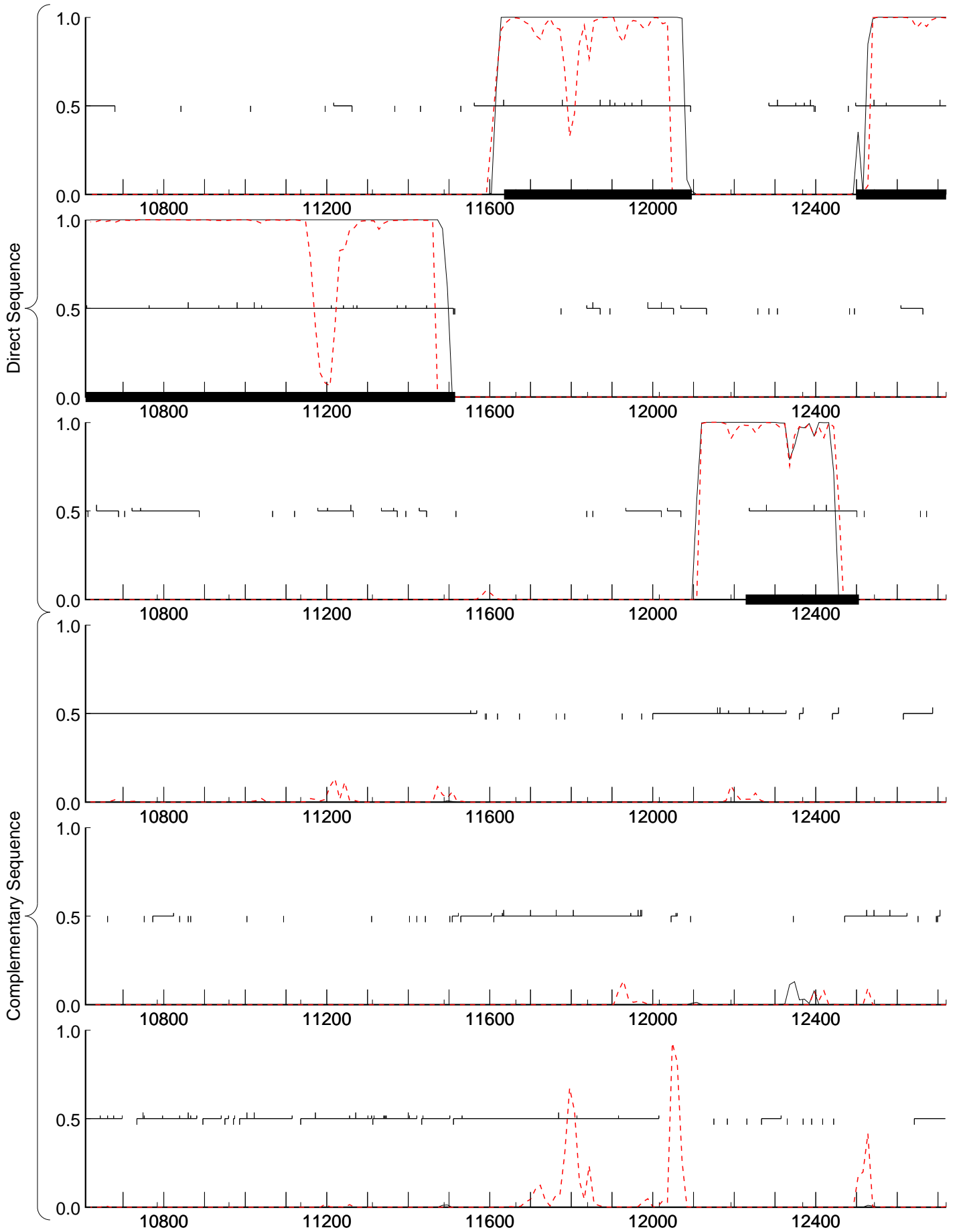
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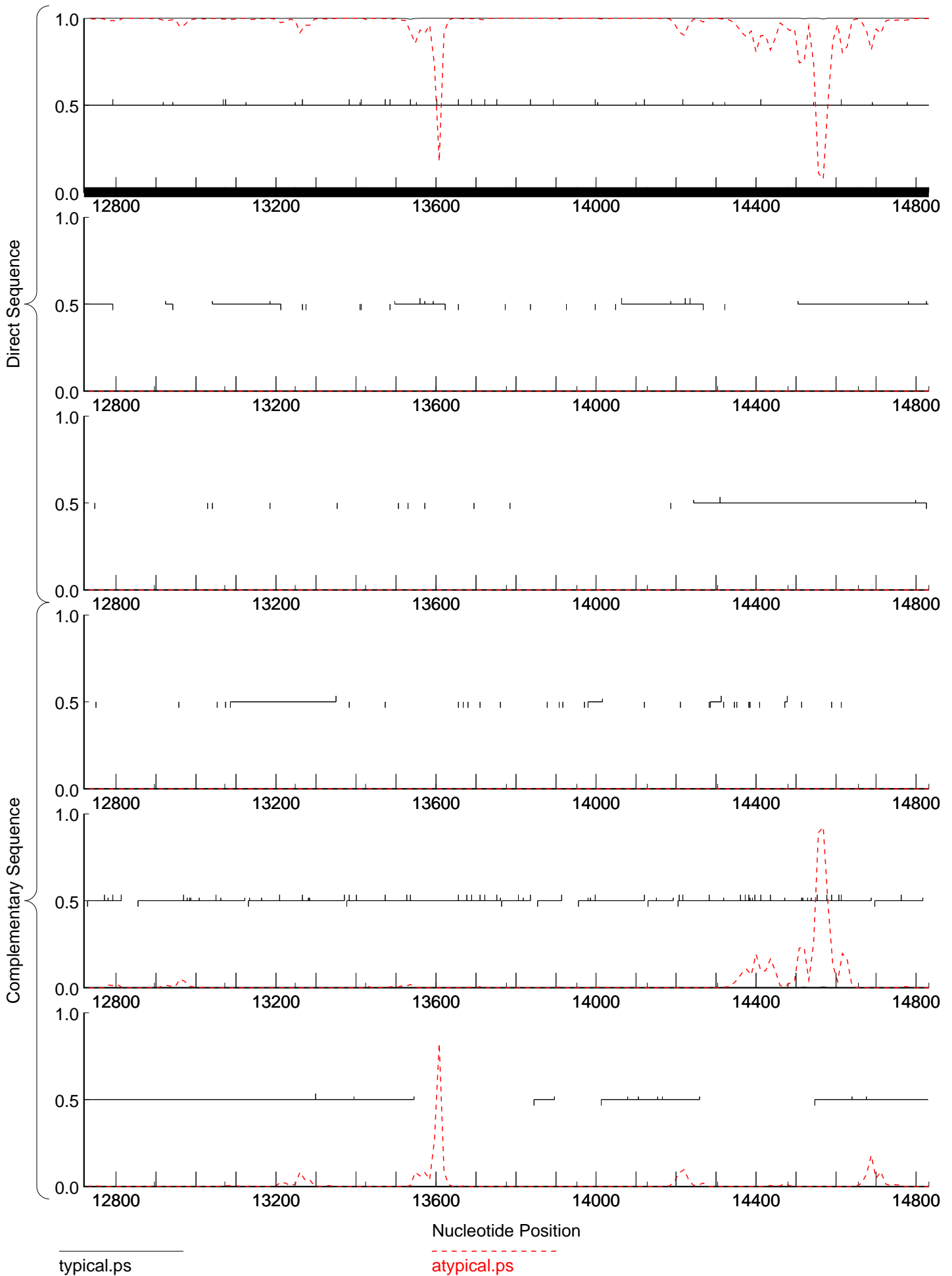
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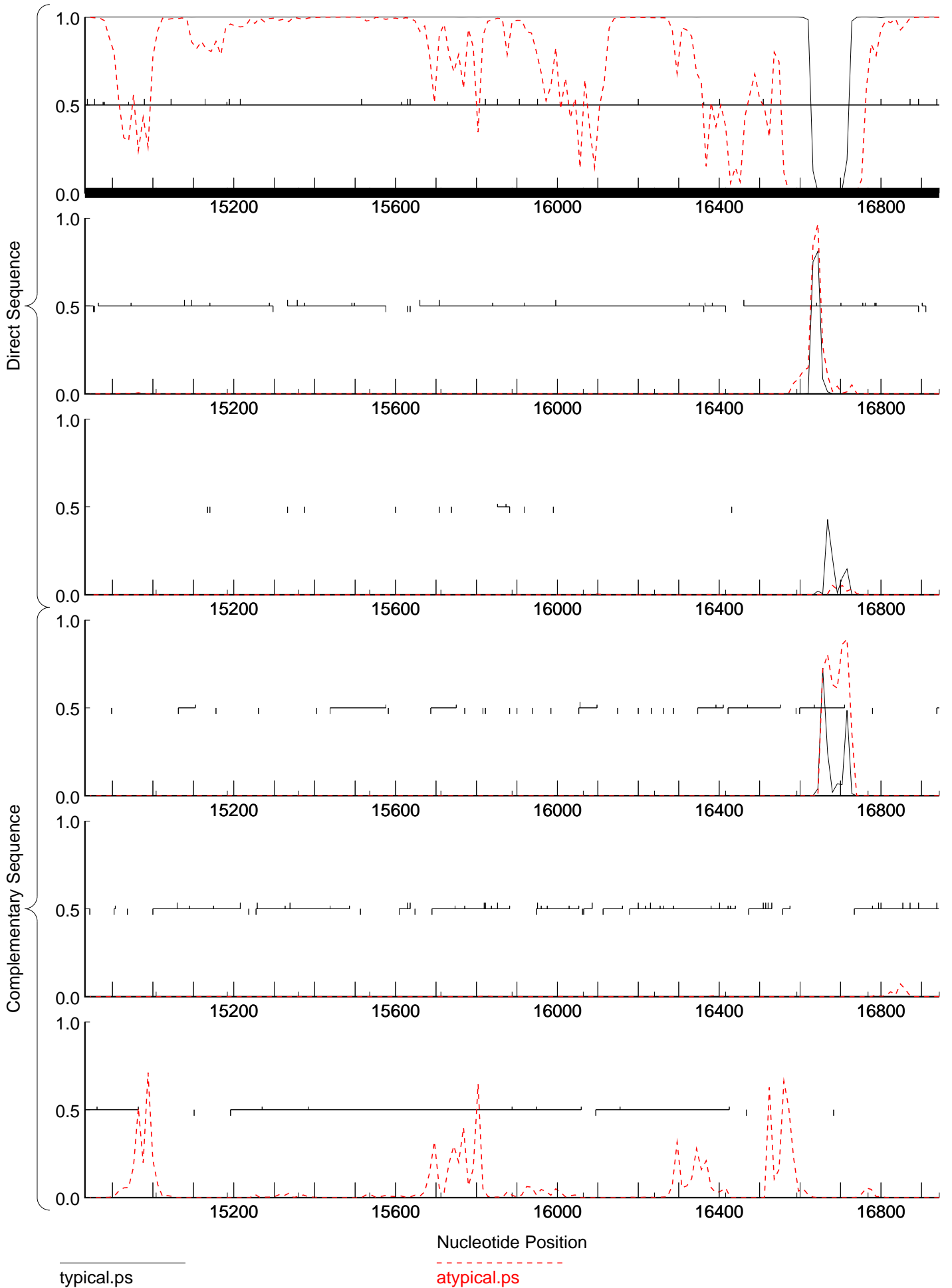
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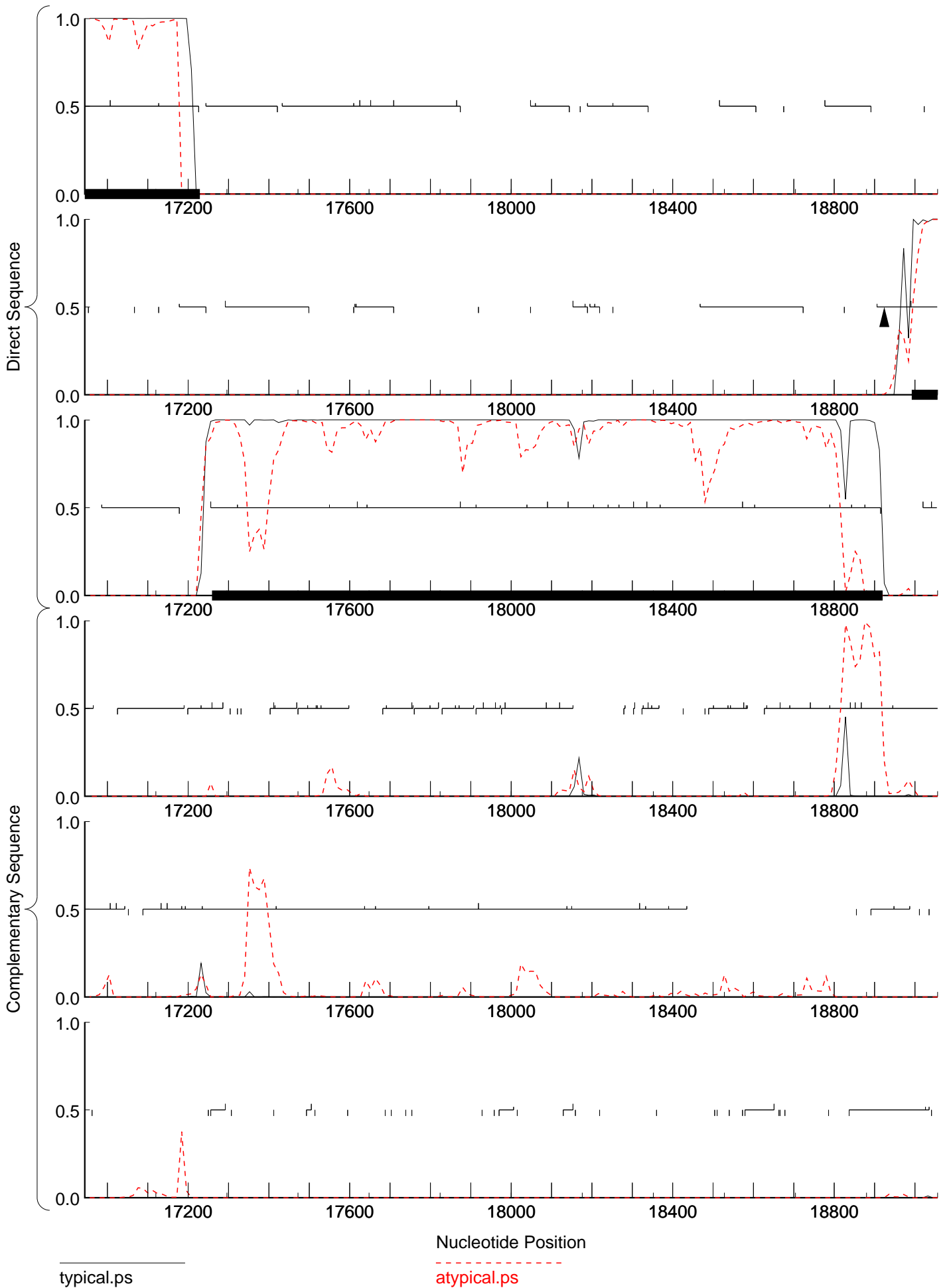
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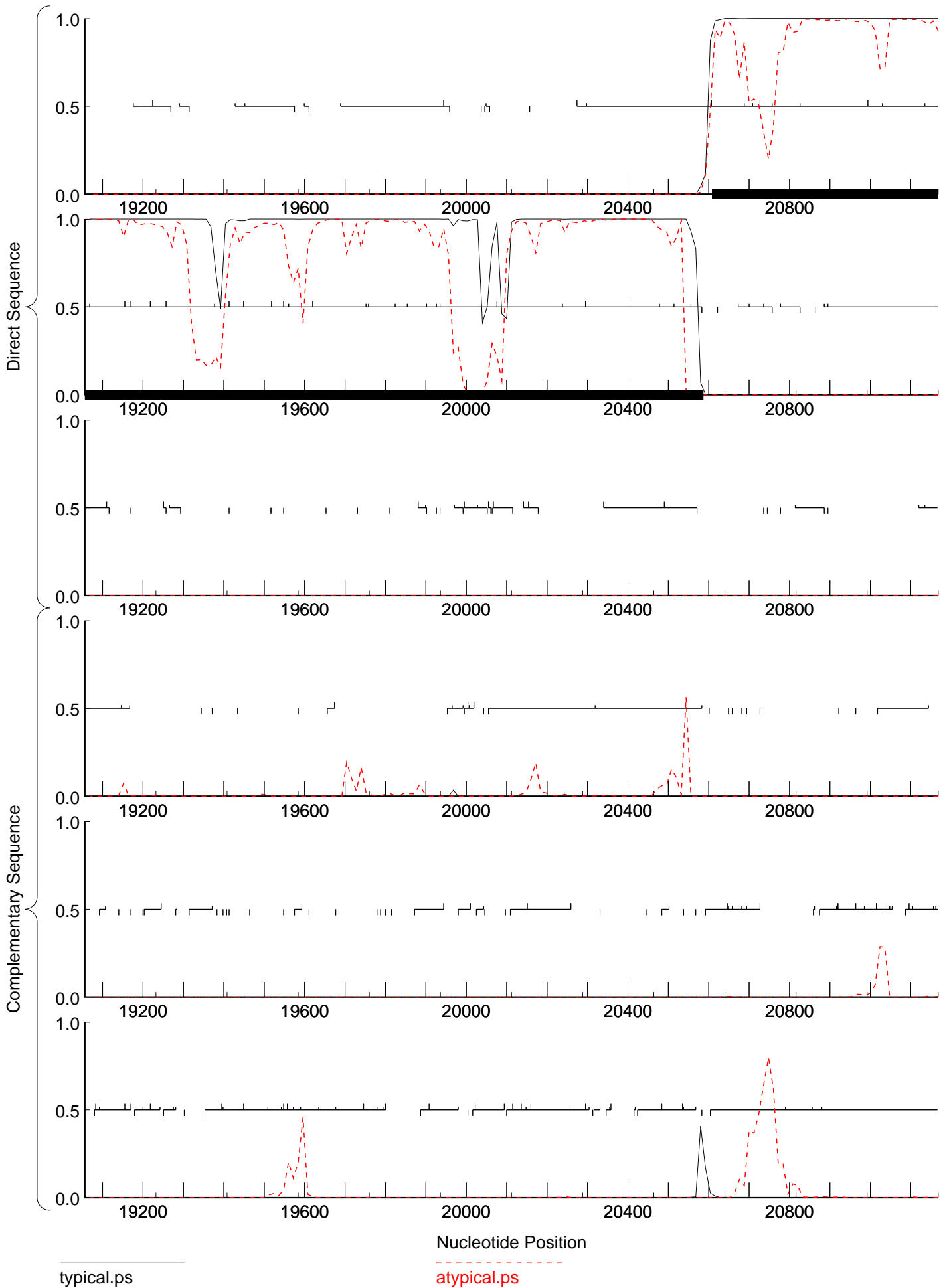


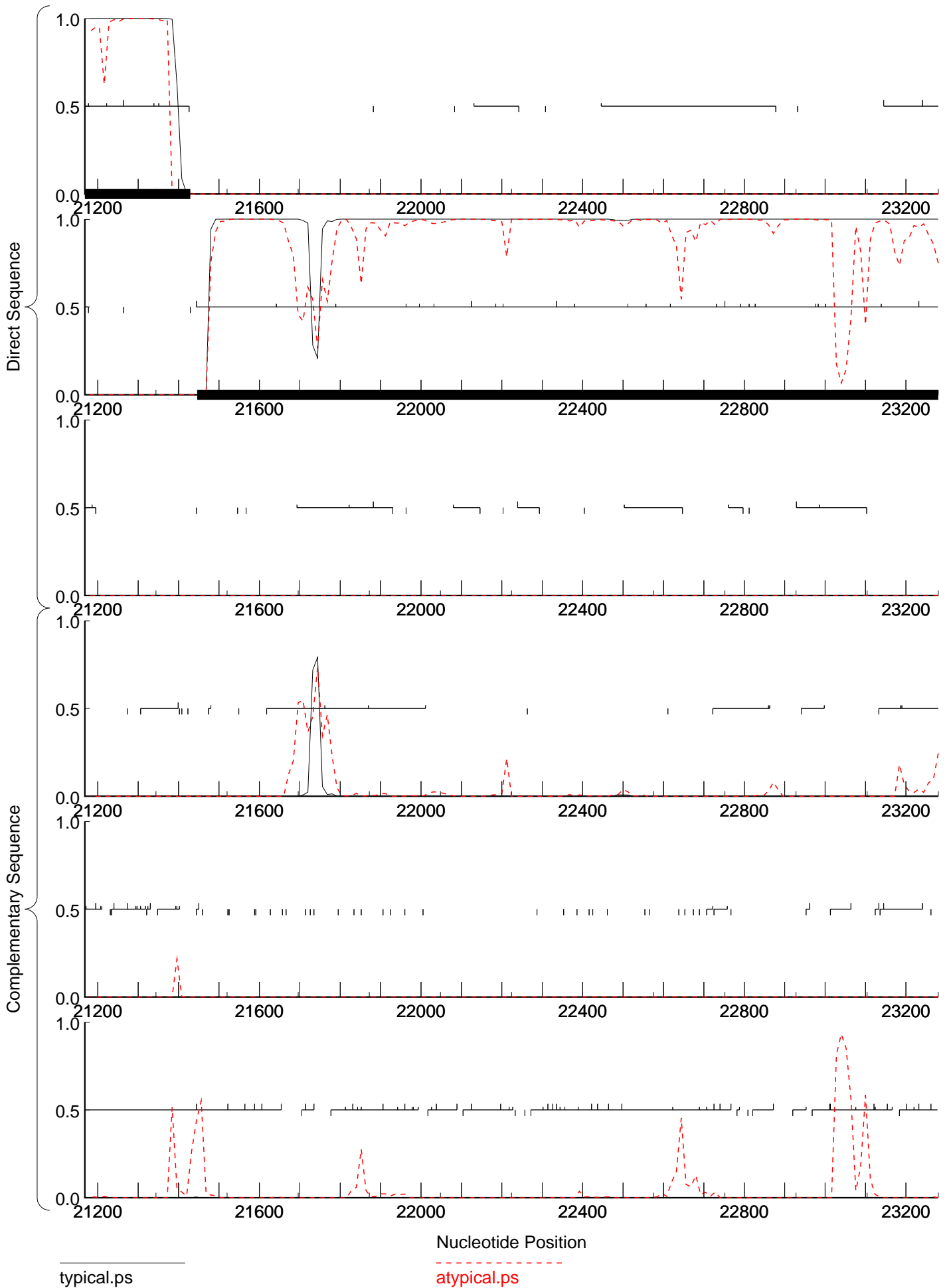
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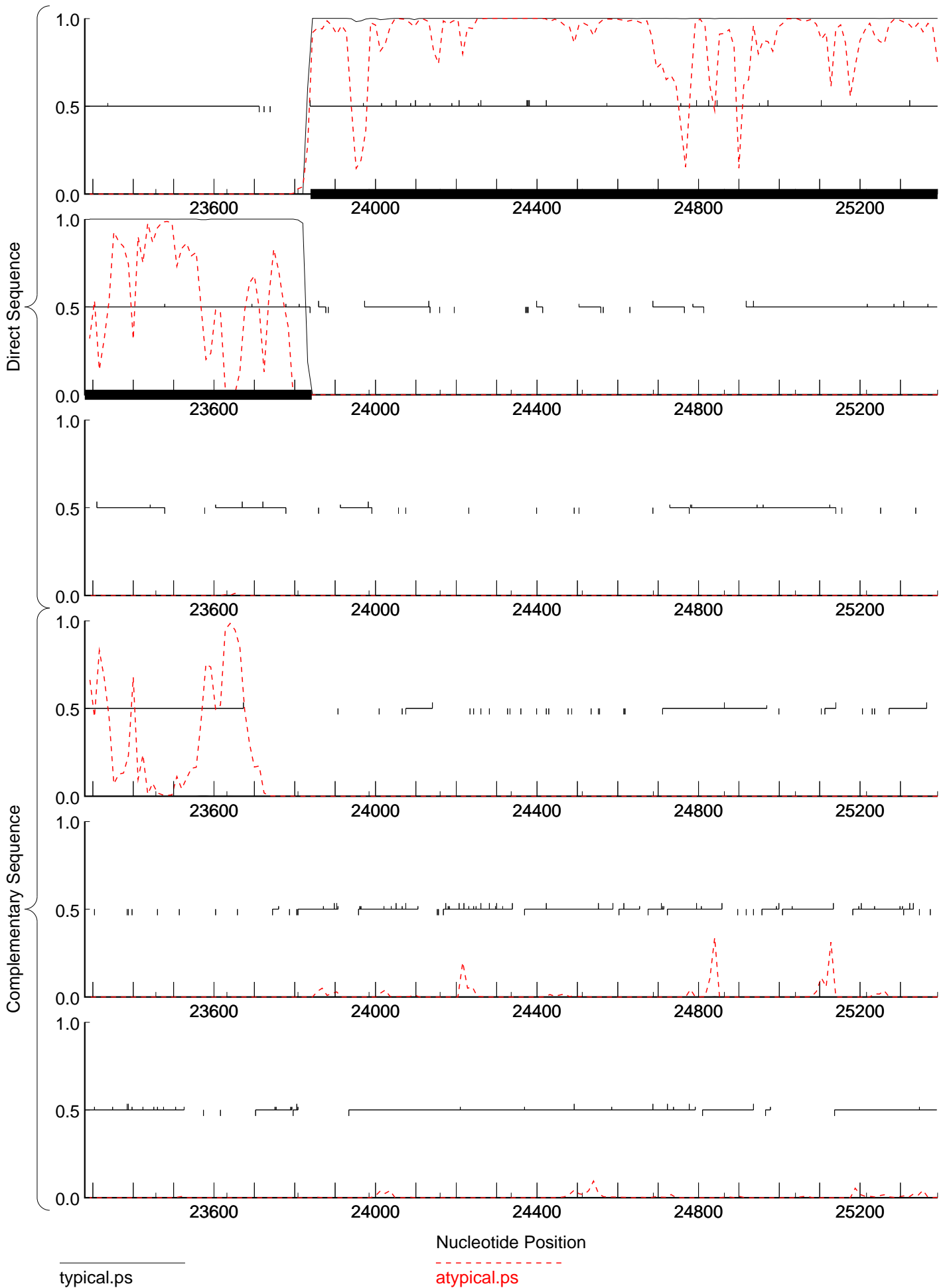
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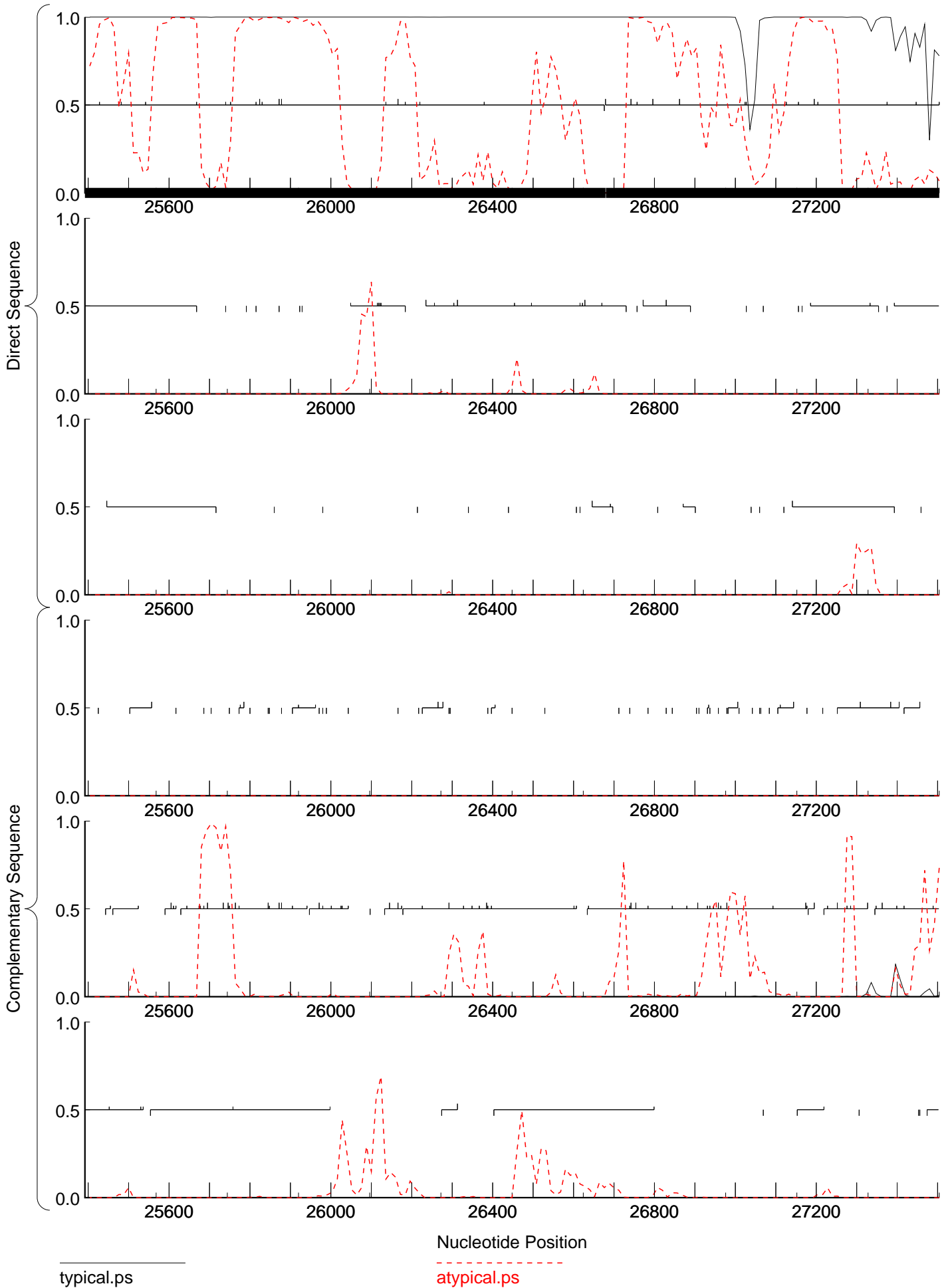


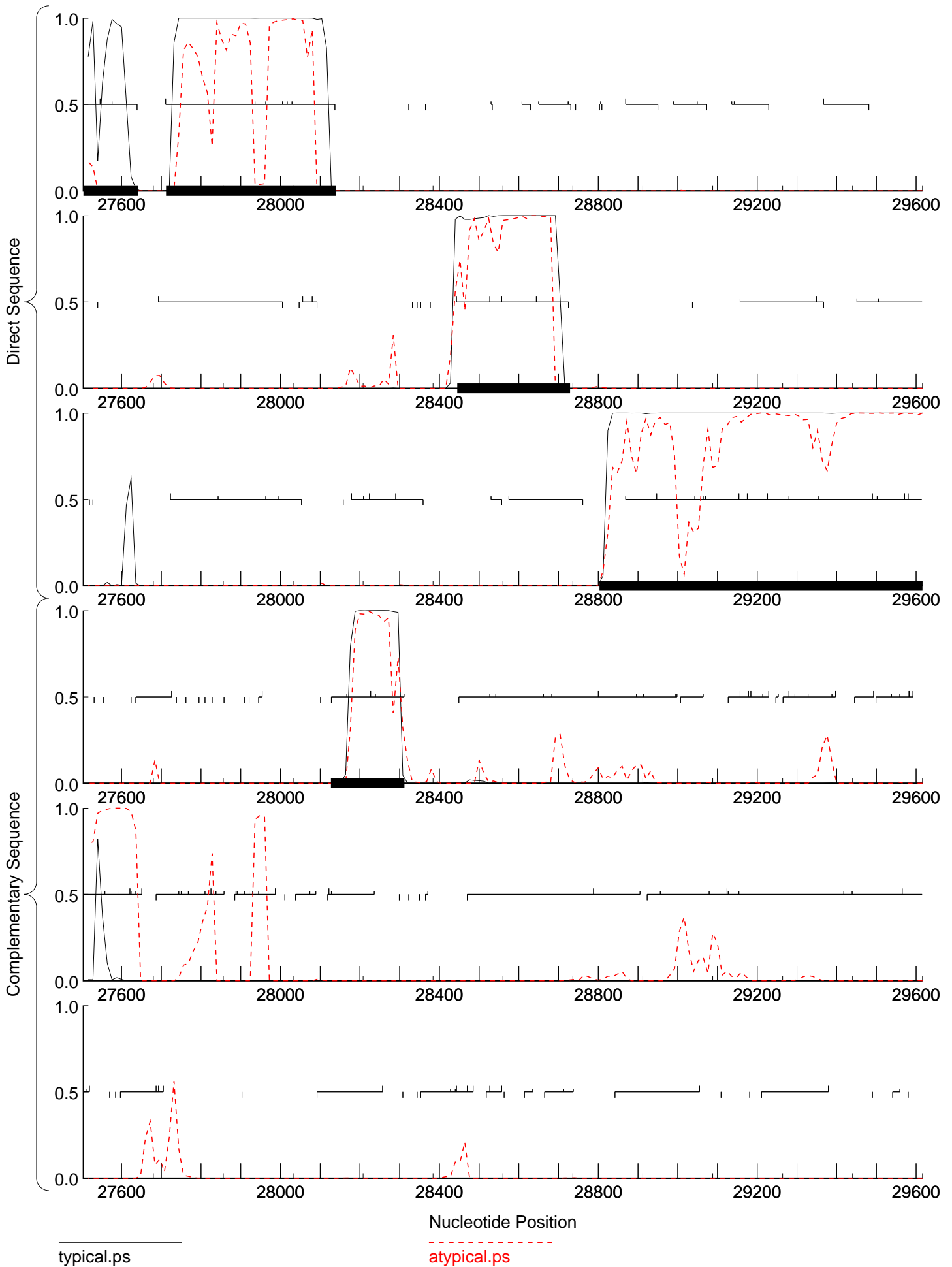


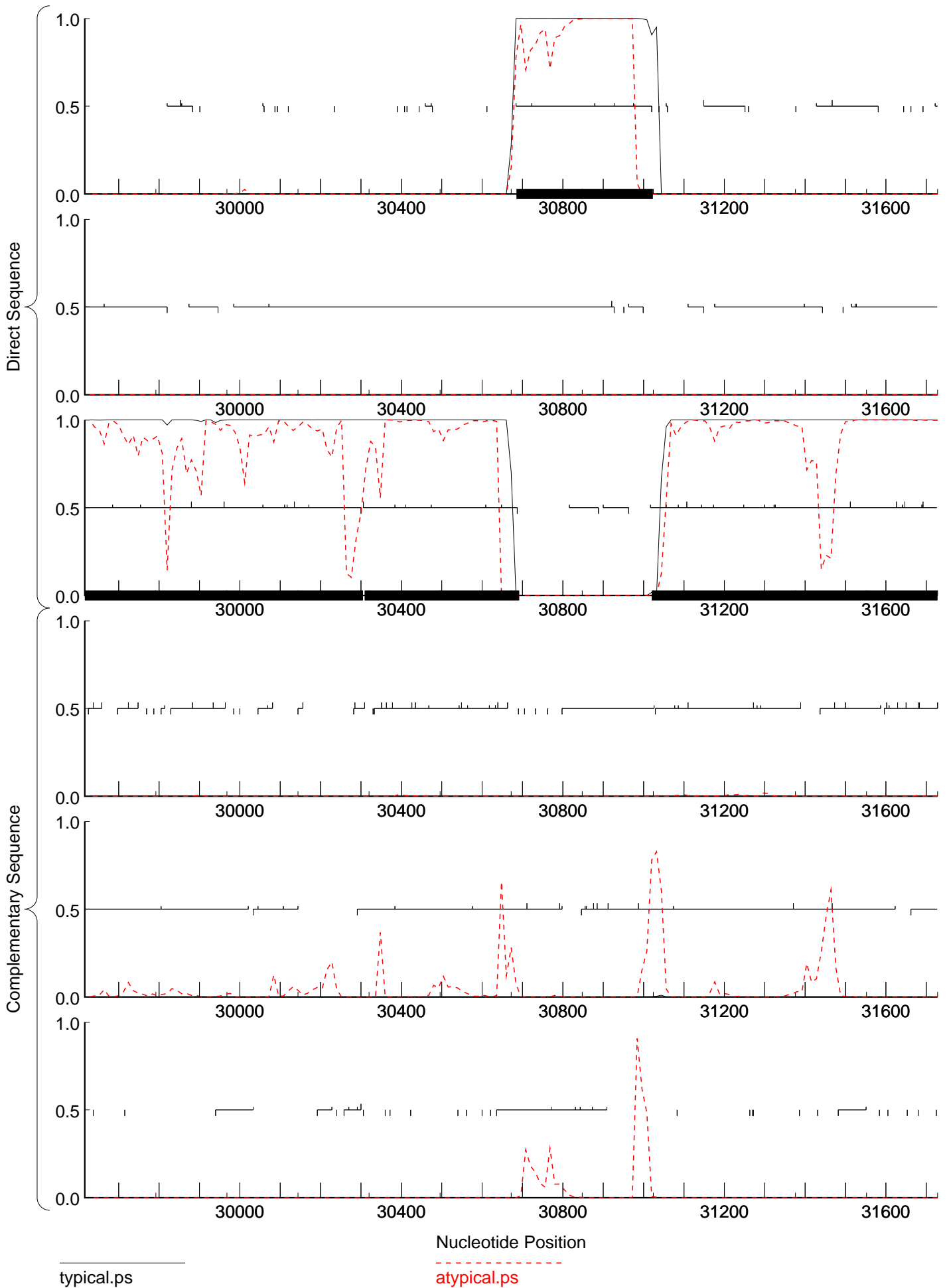


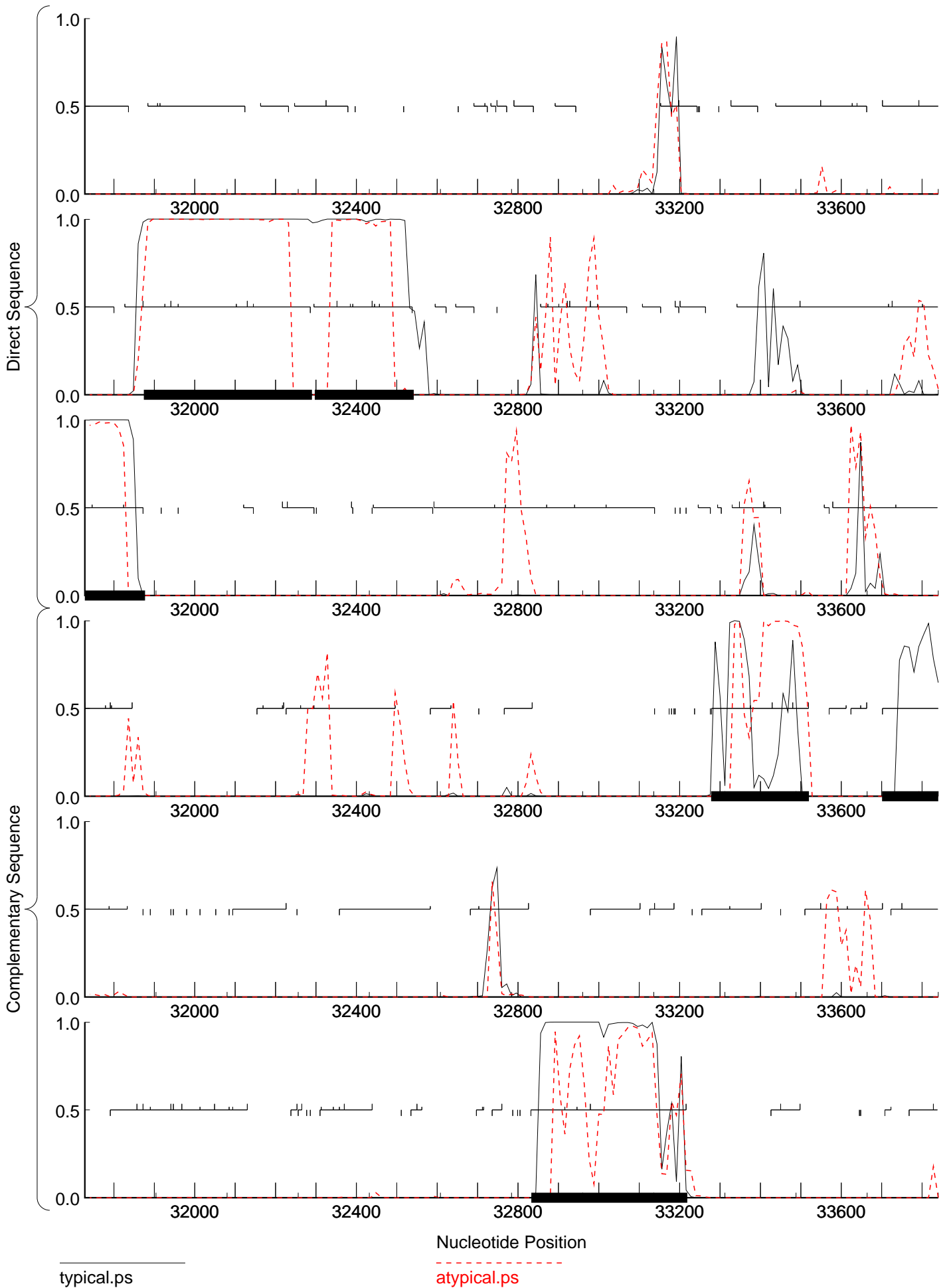
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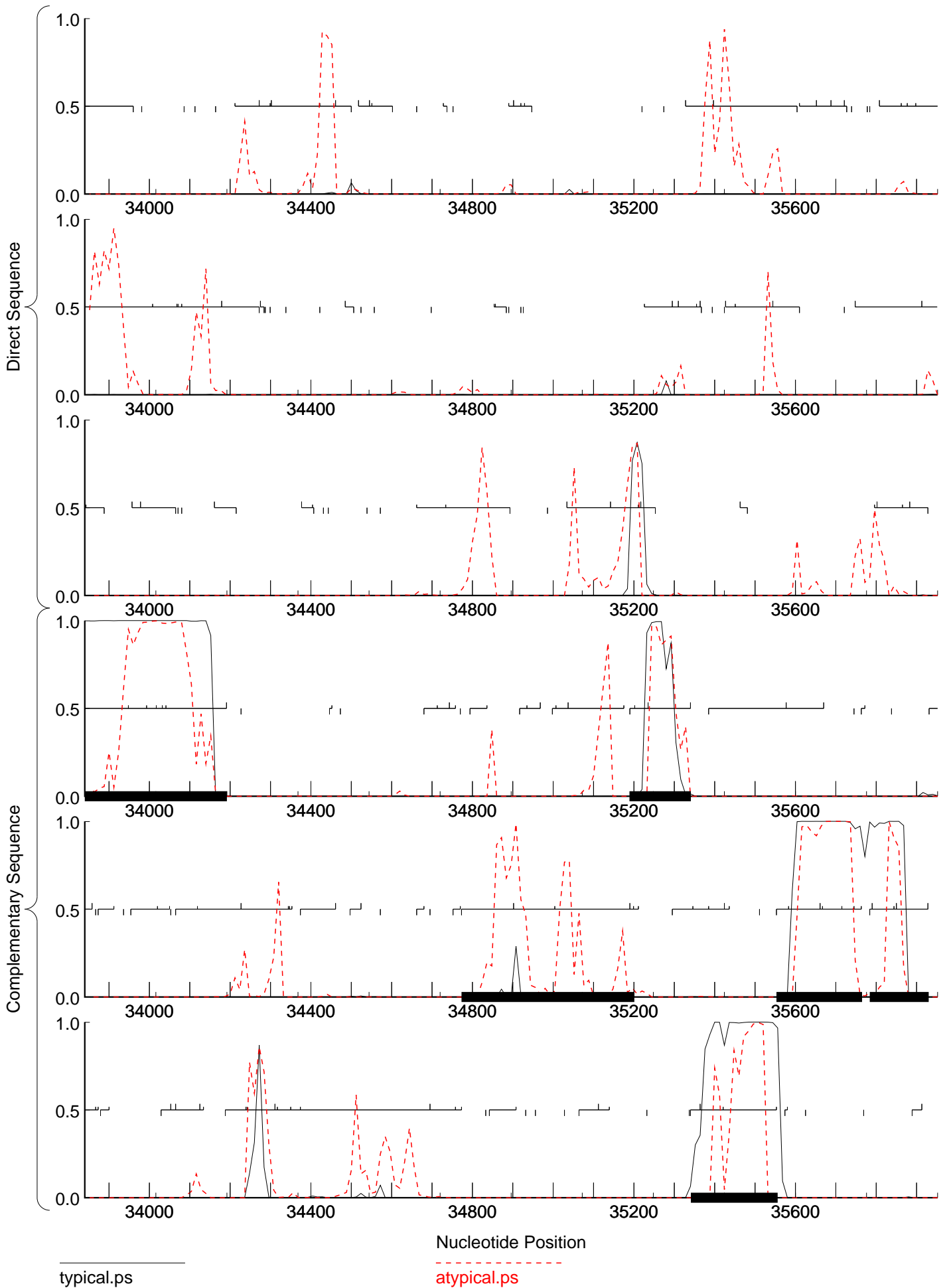
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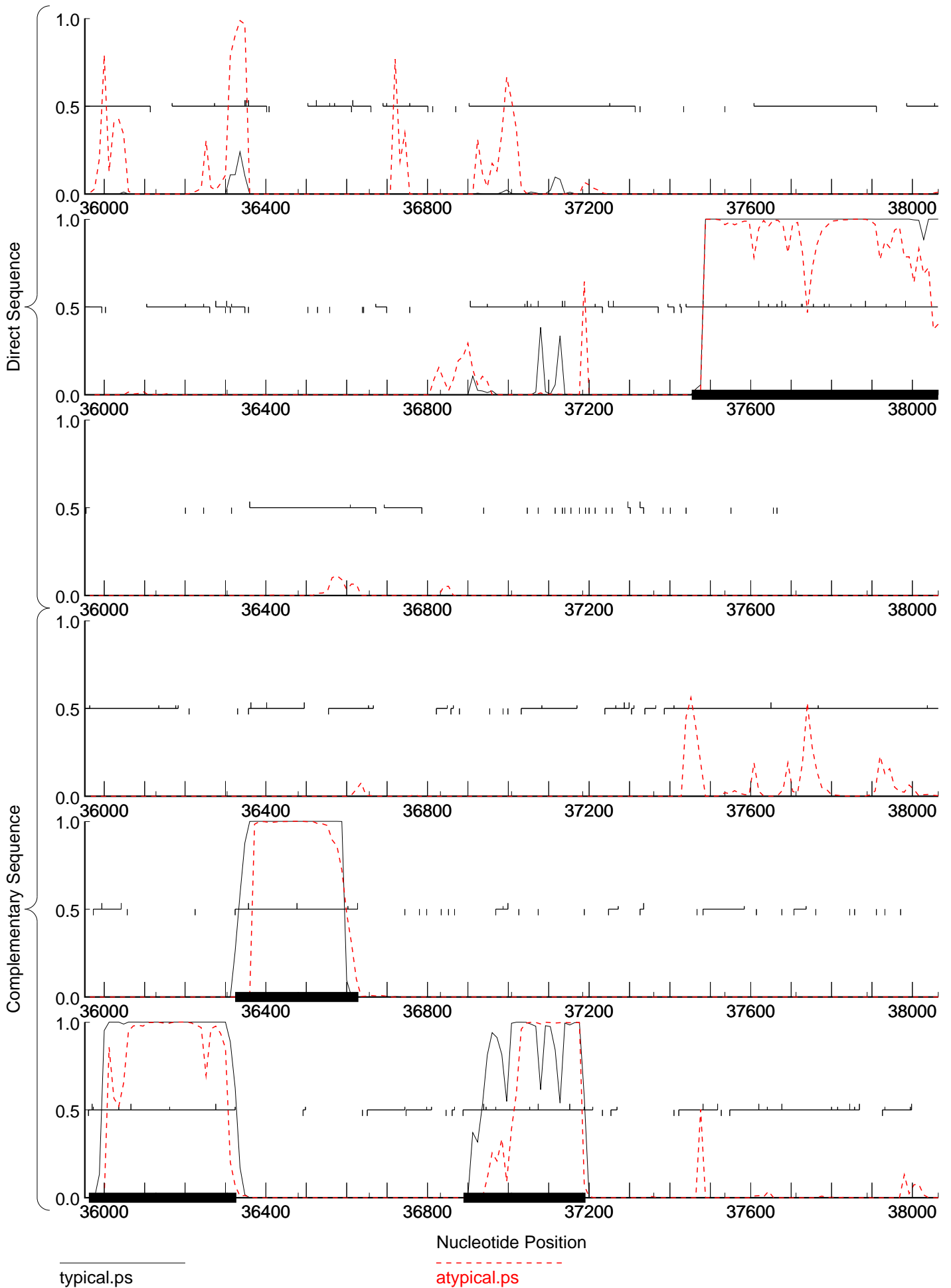






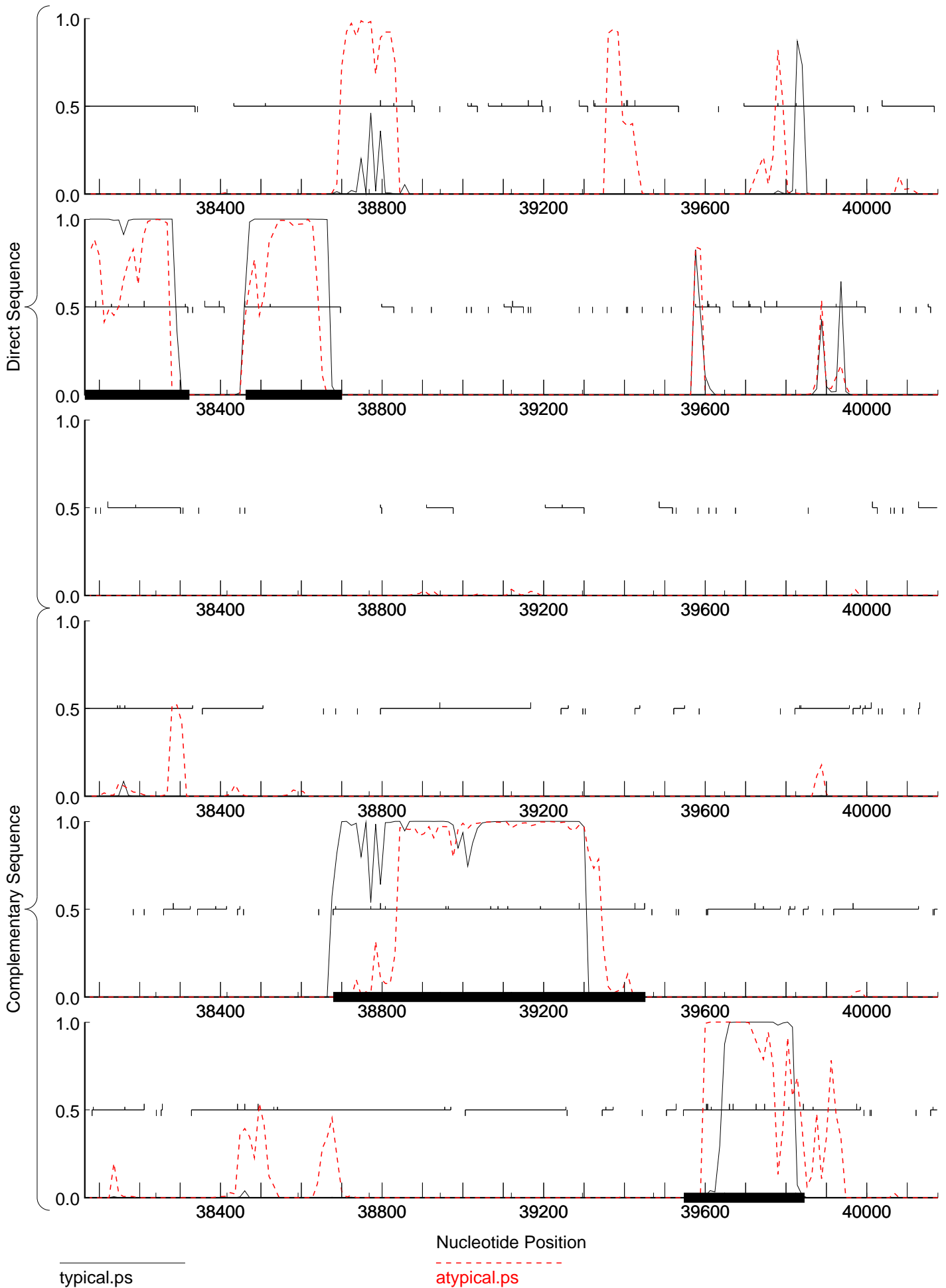


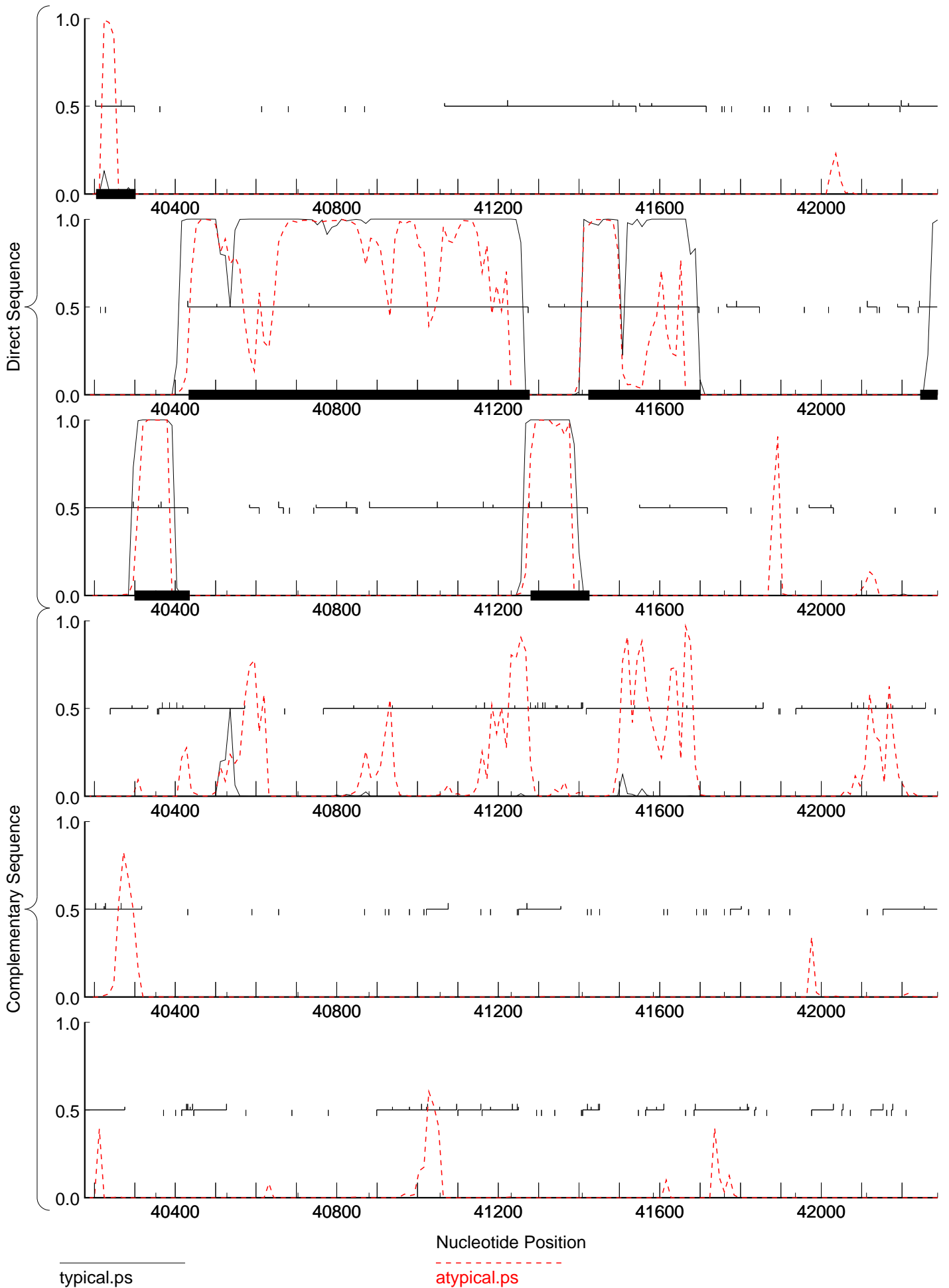


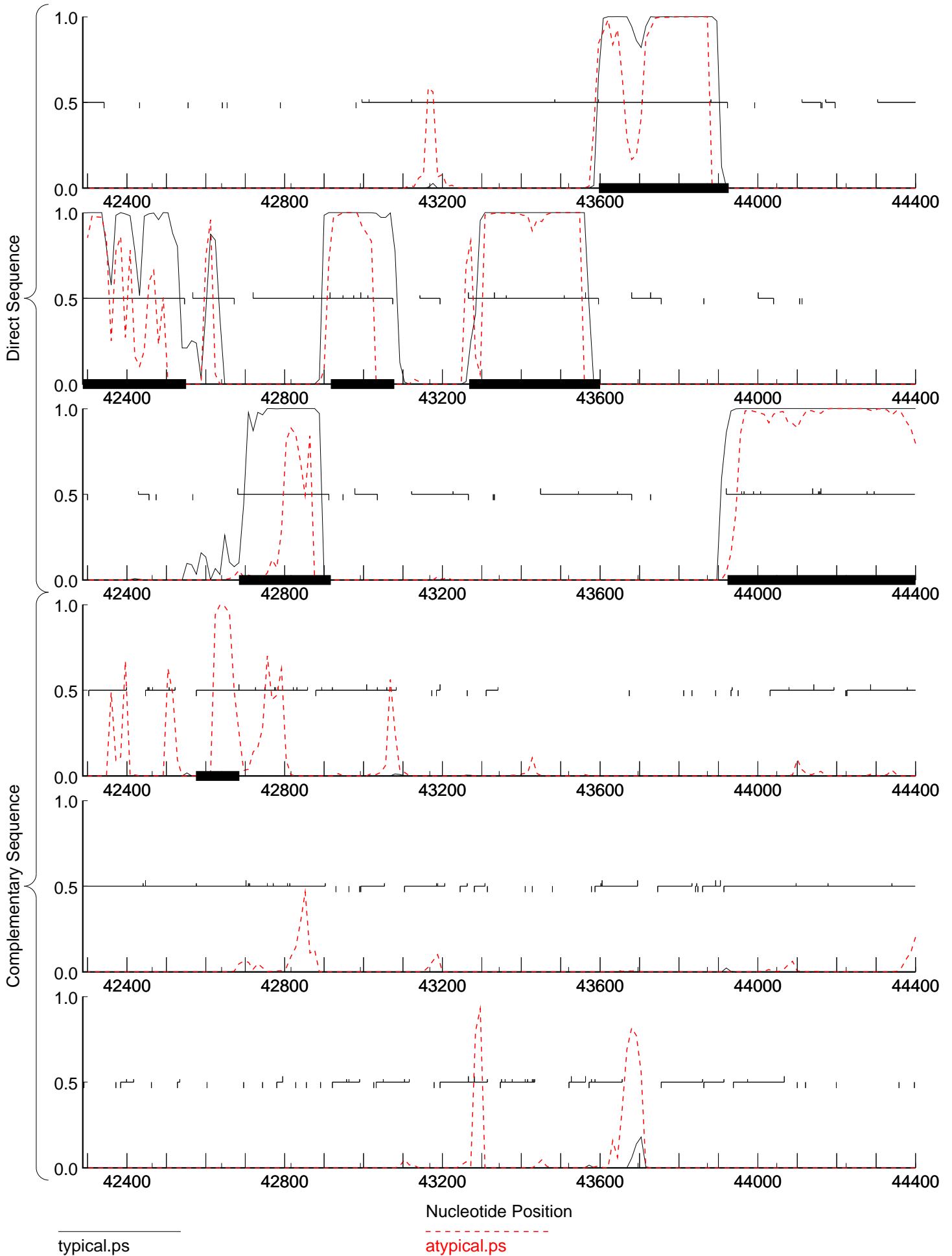


typical.ps

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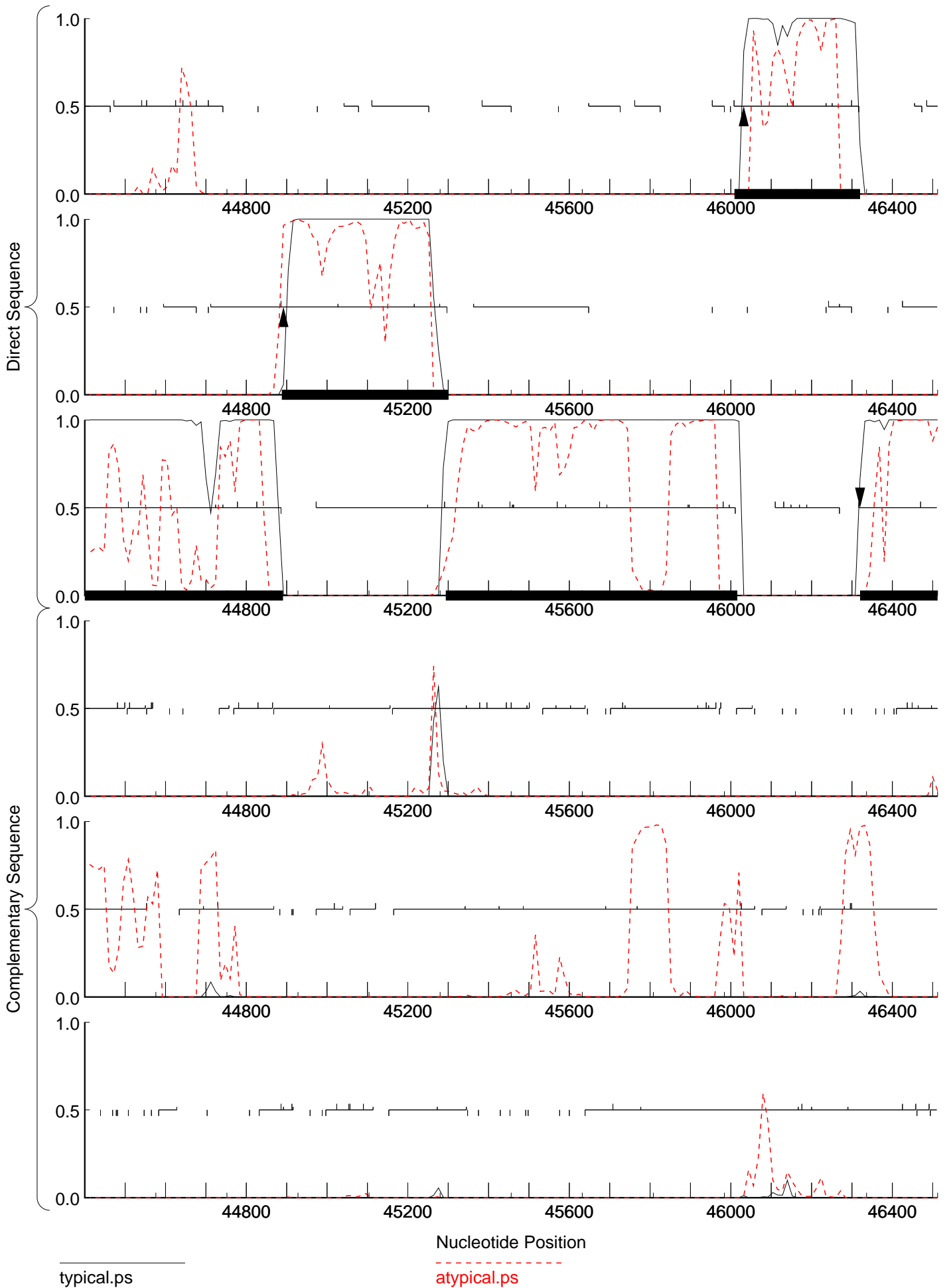






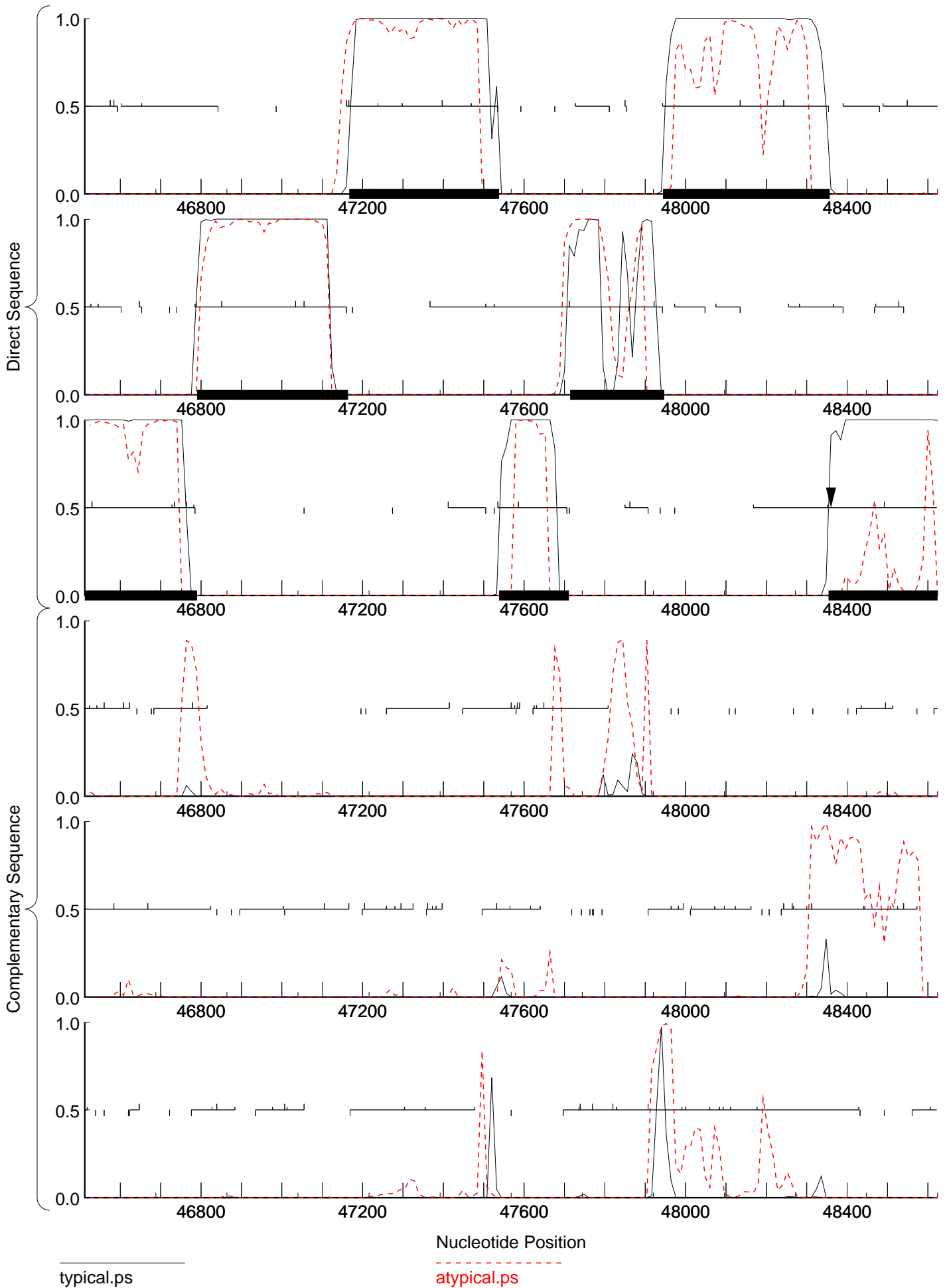
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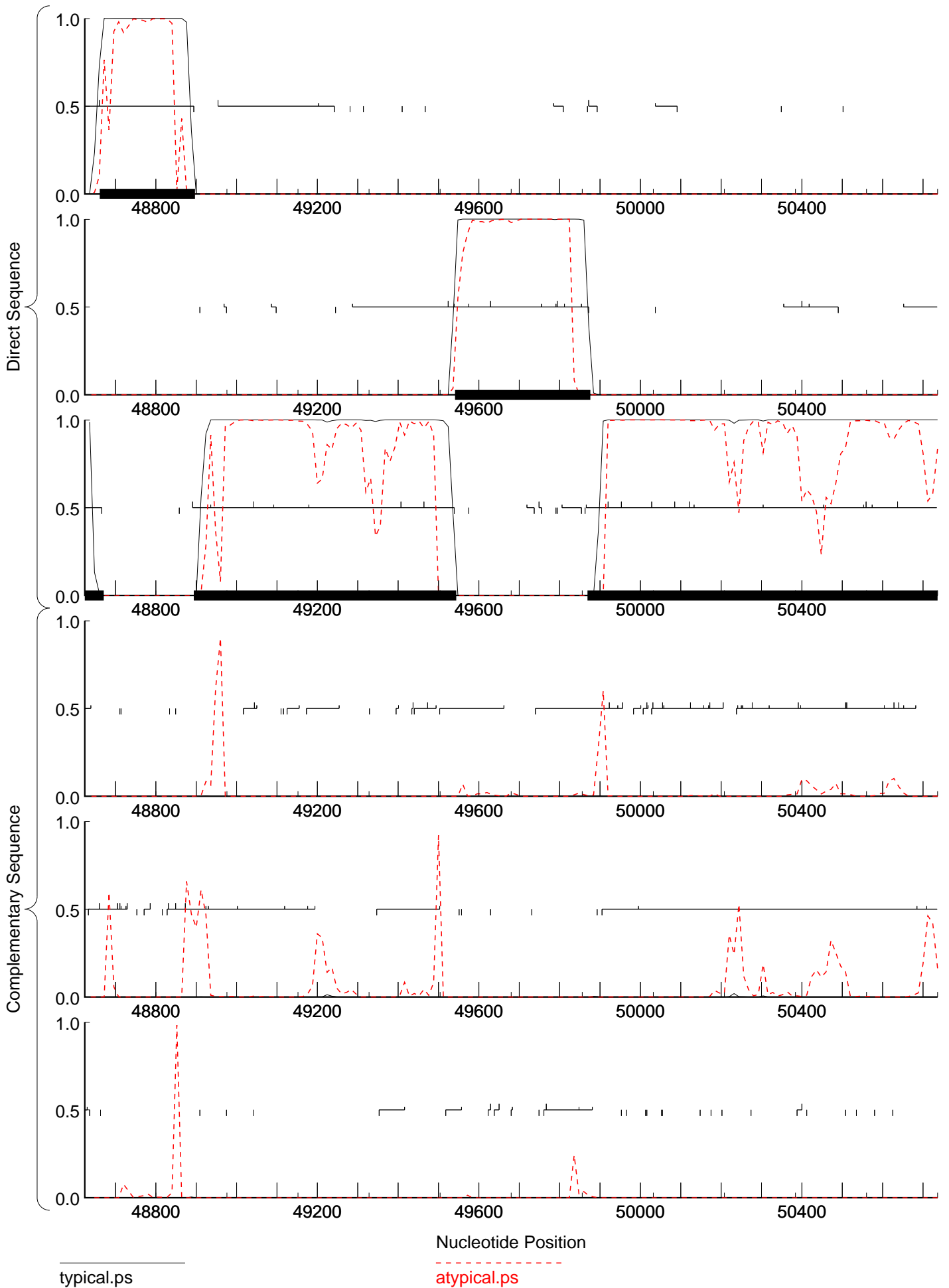
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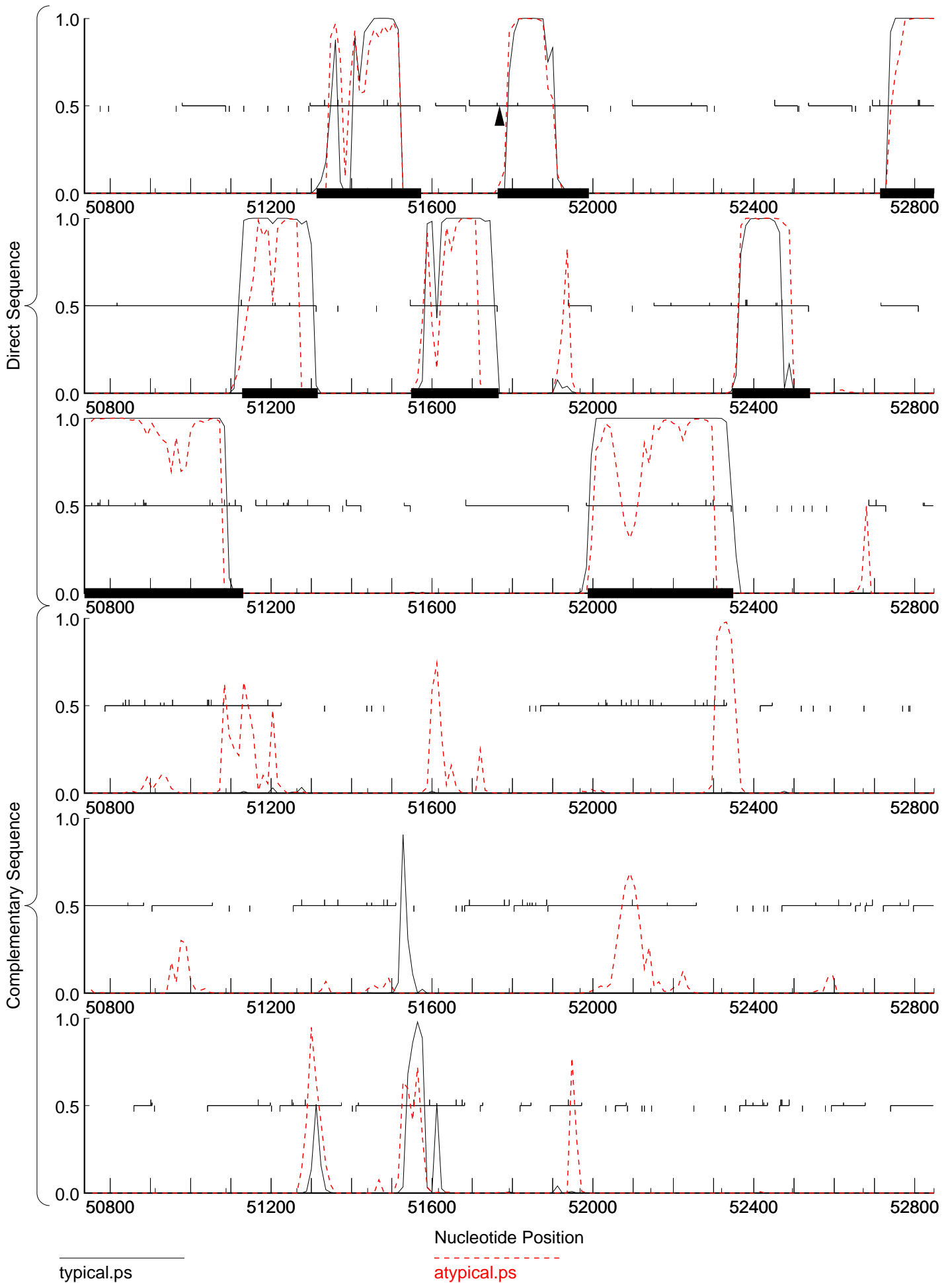


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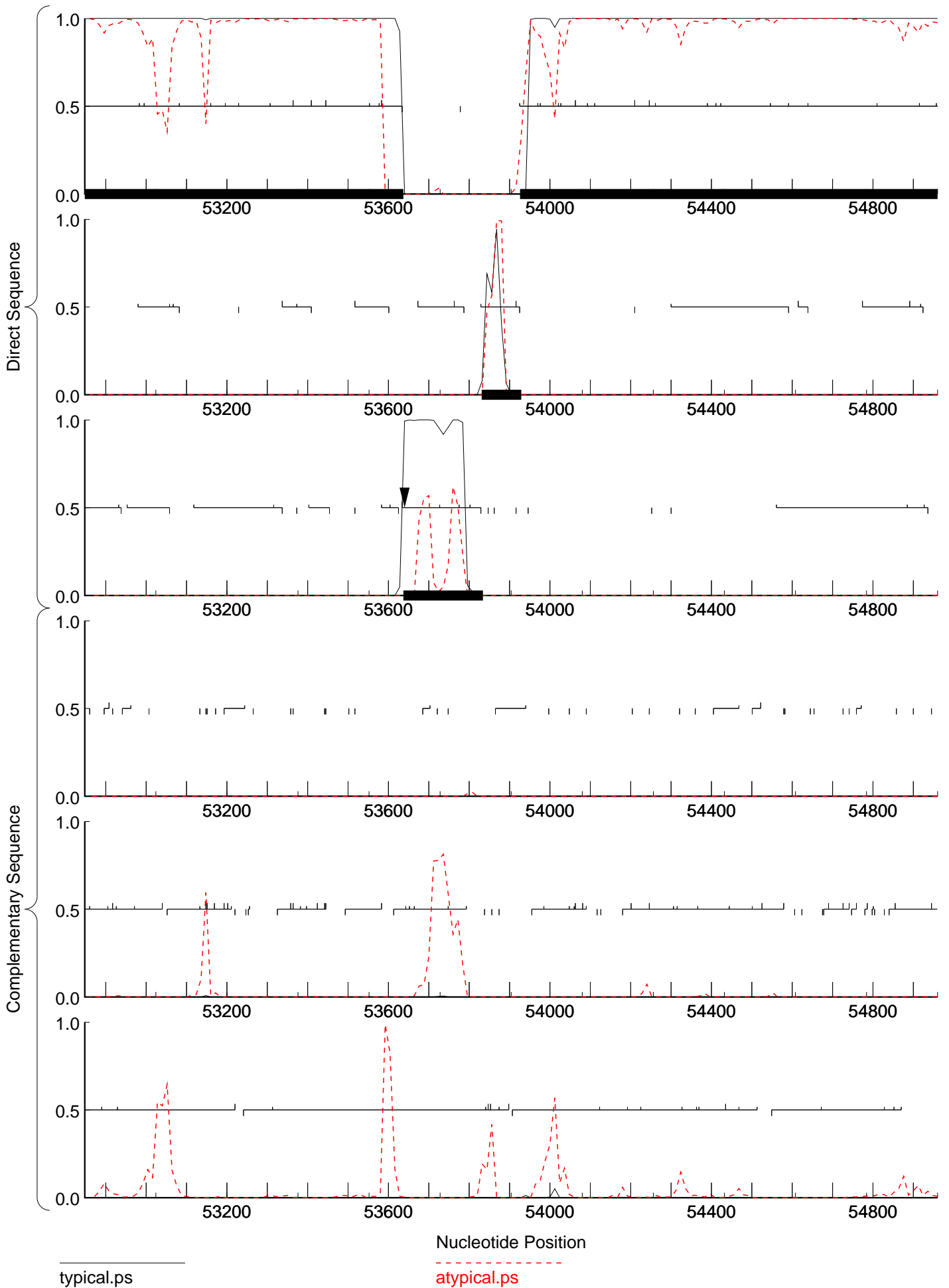
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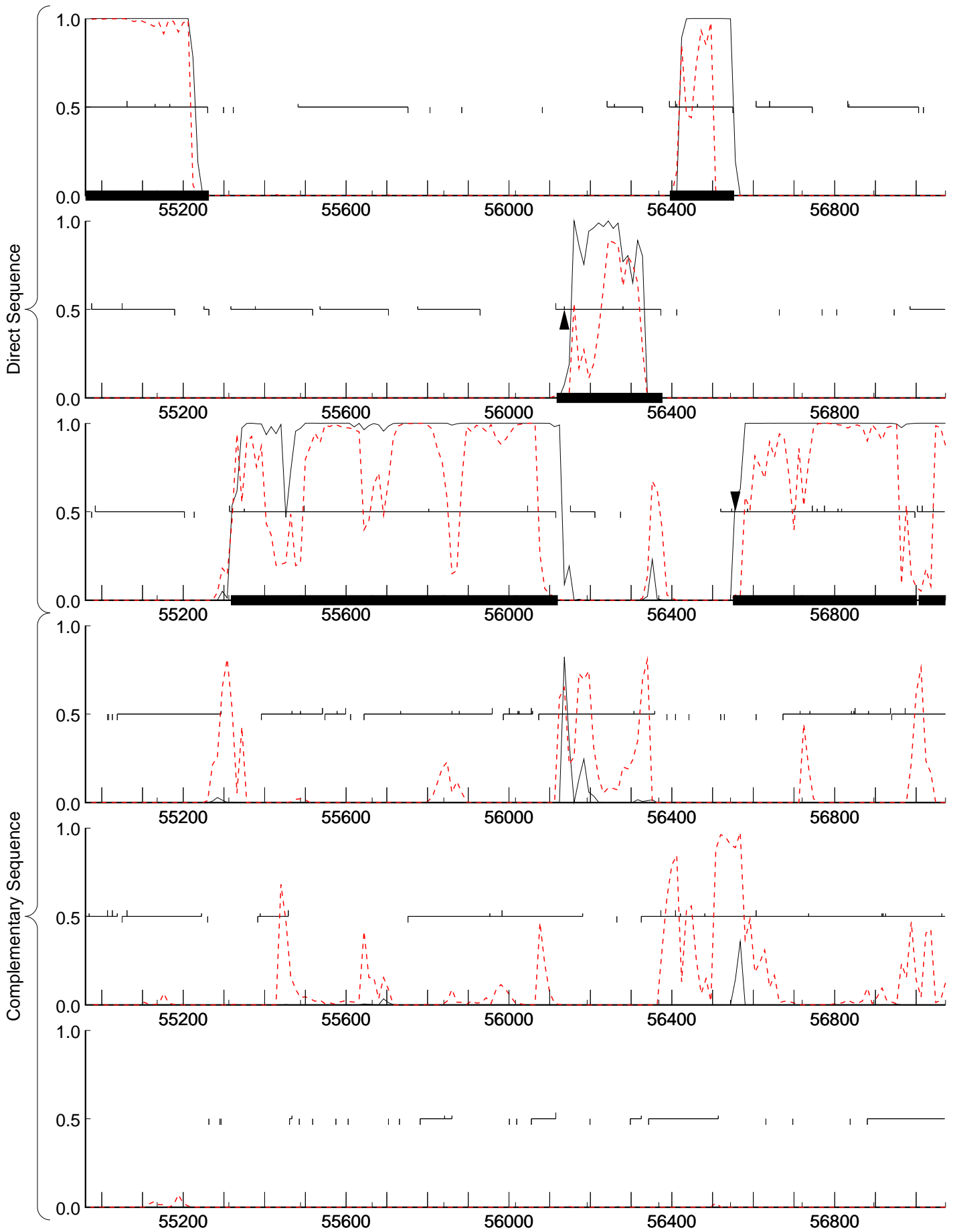






...dles complete sequence, 75070 bp including 9-base 3' overhang (CGCTTGTC), Cluster E, Order 4, Window 96, Step 12, 27/37
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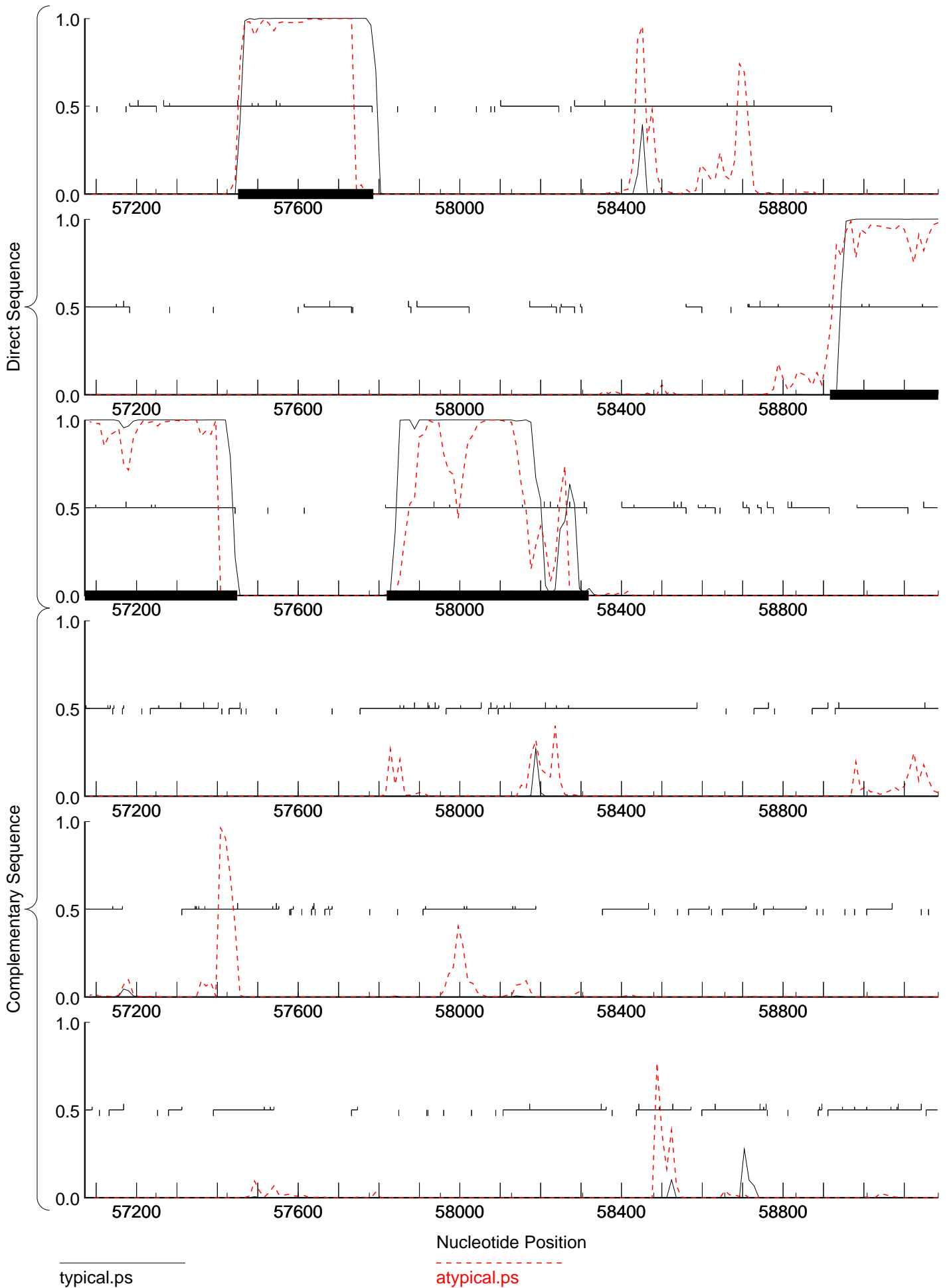


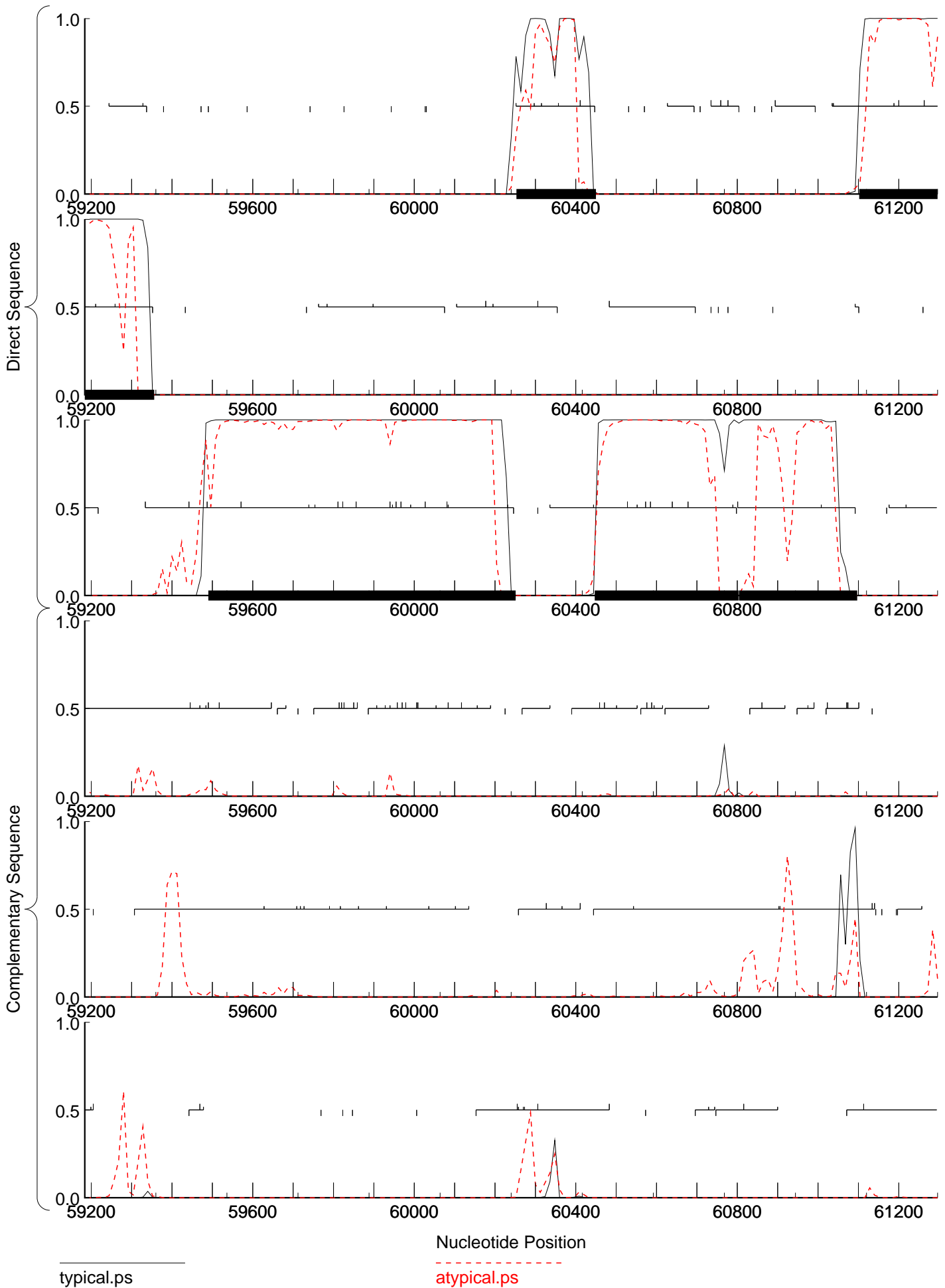


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Nucleotide Position

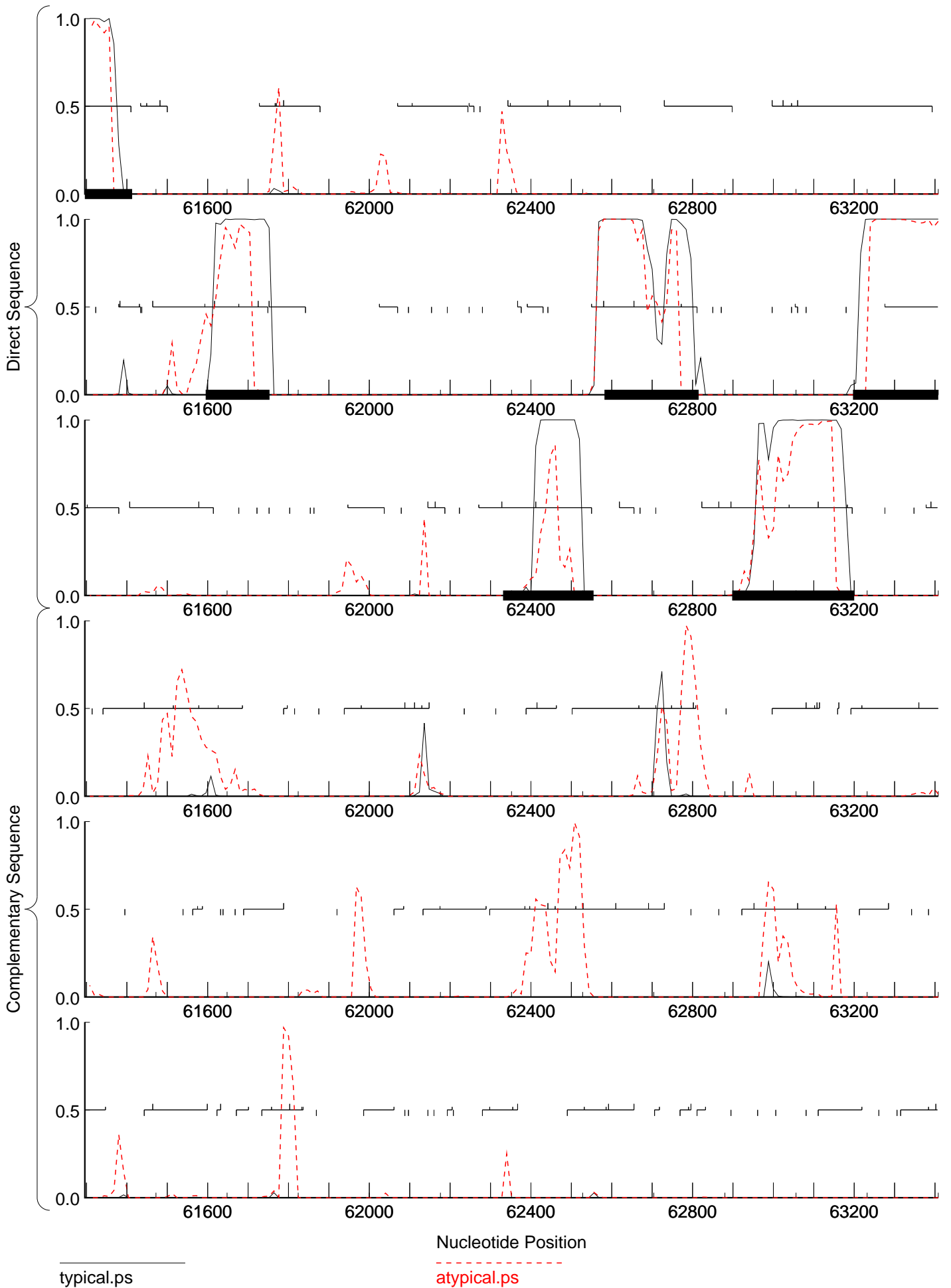
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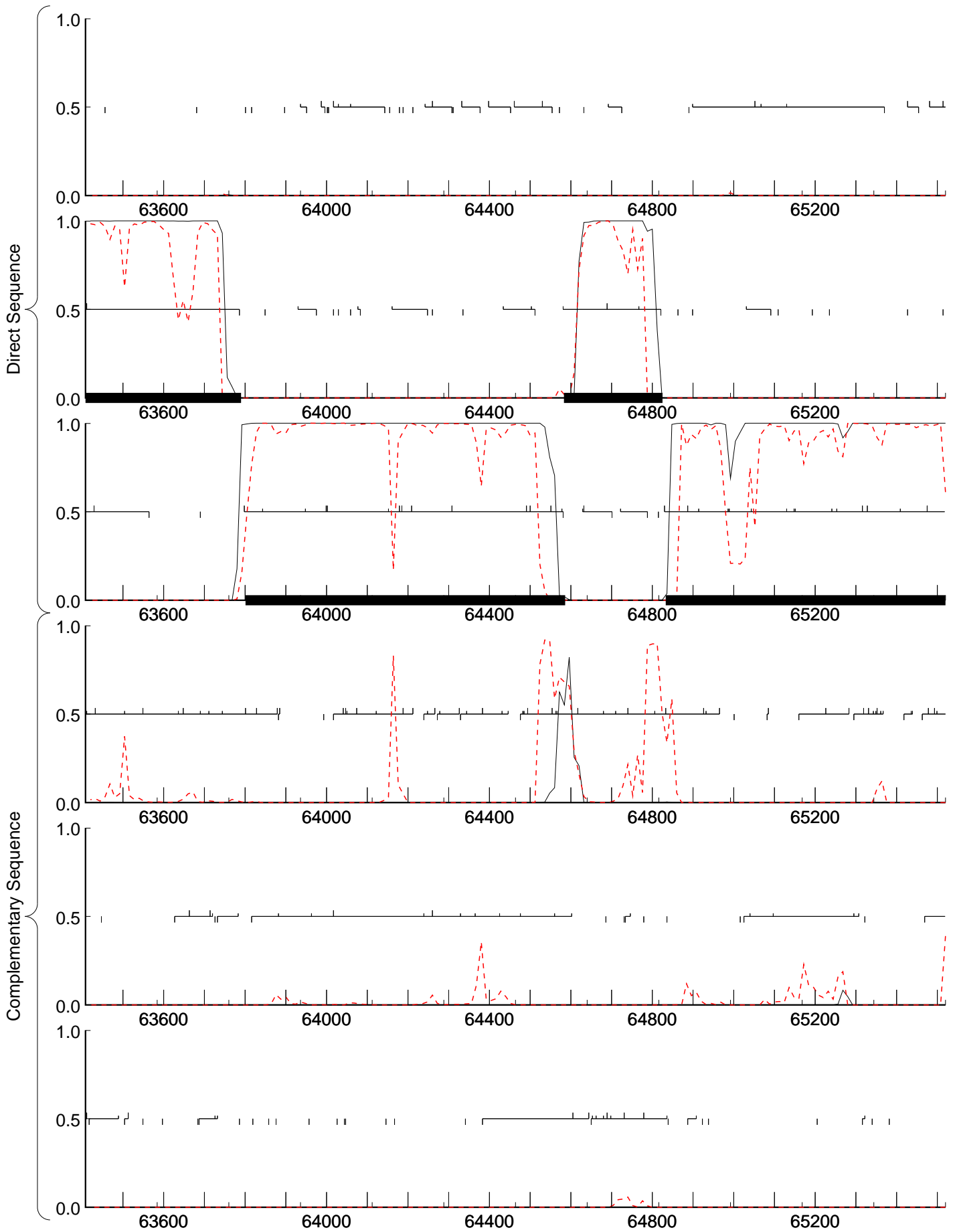




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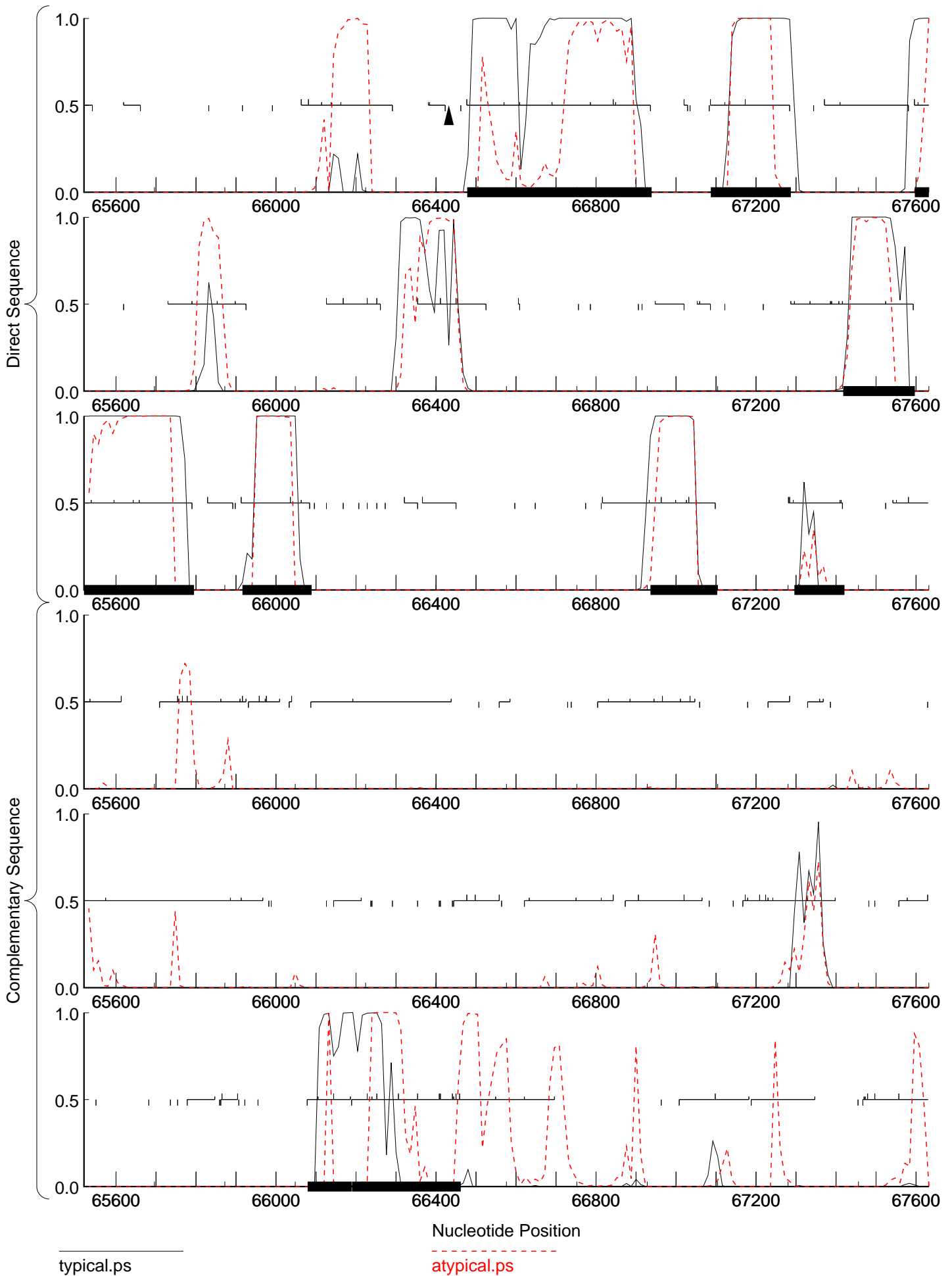
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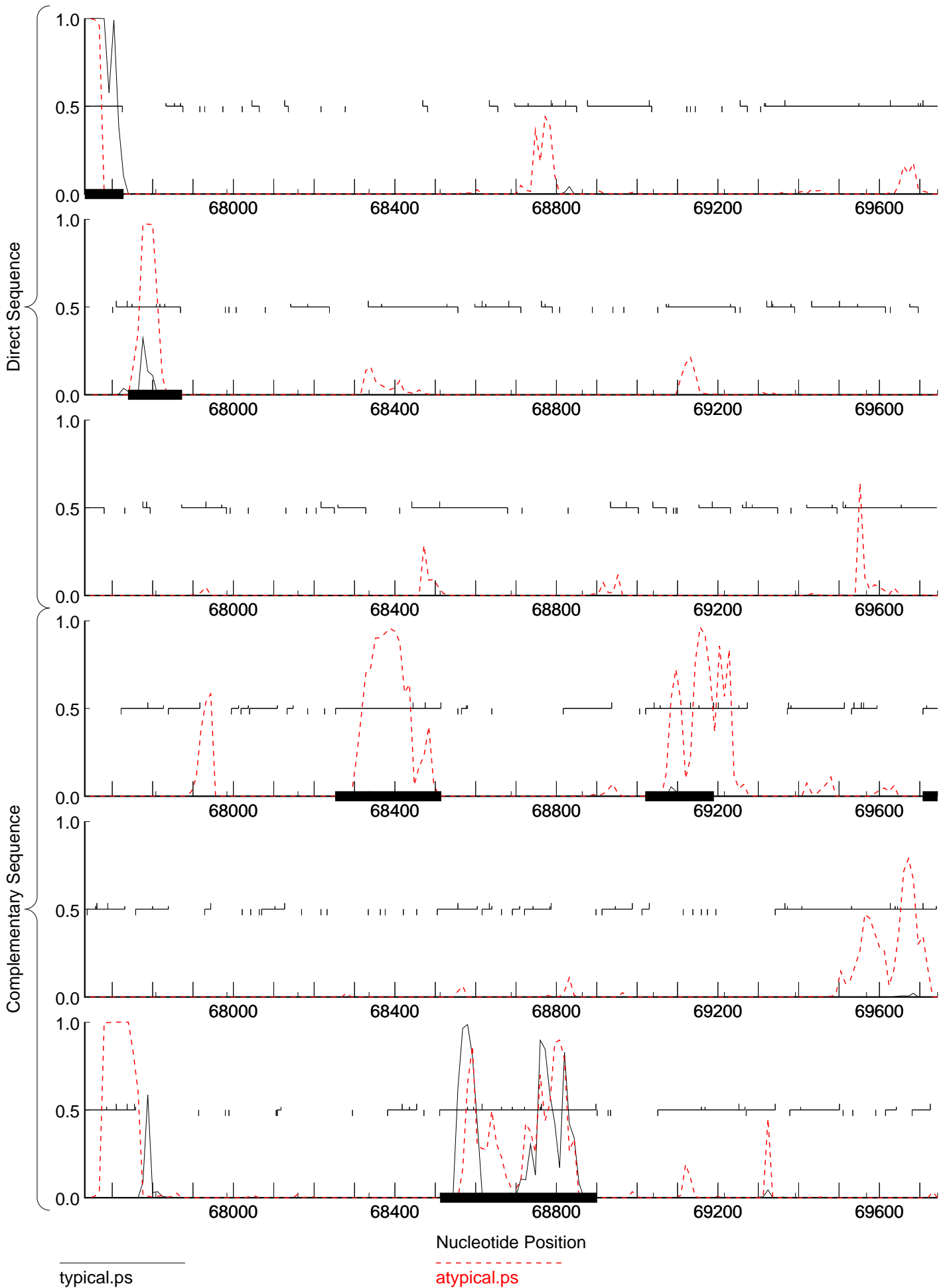
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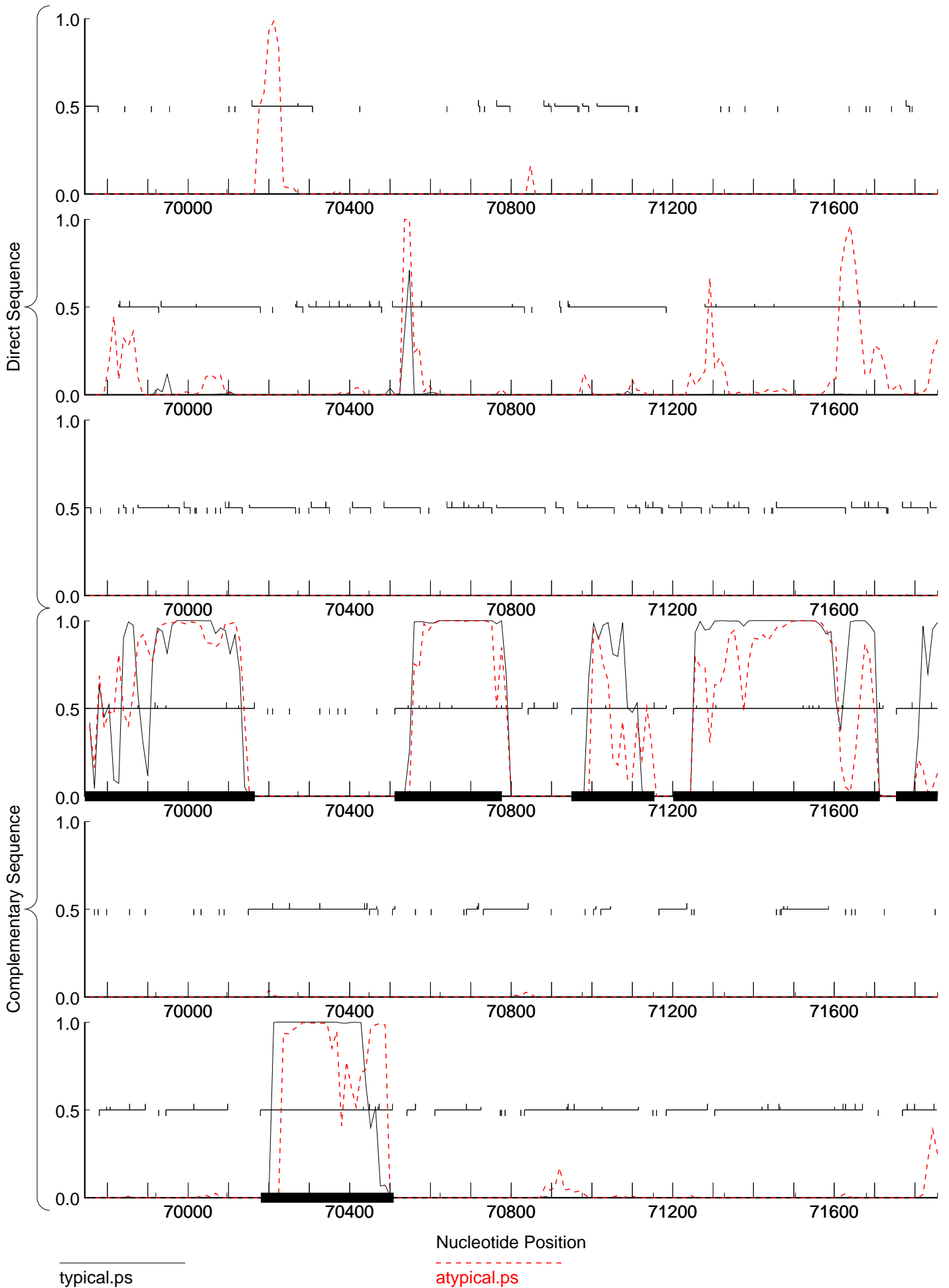
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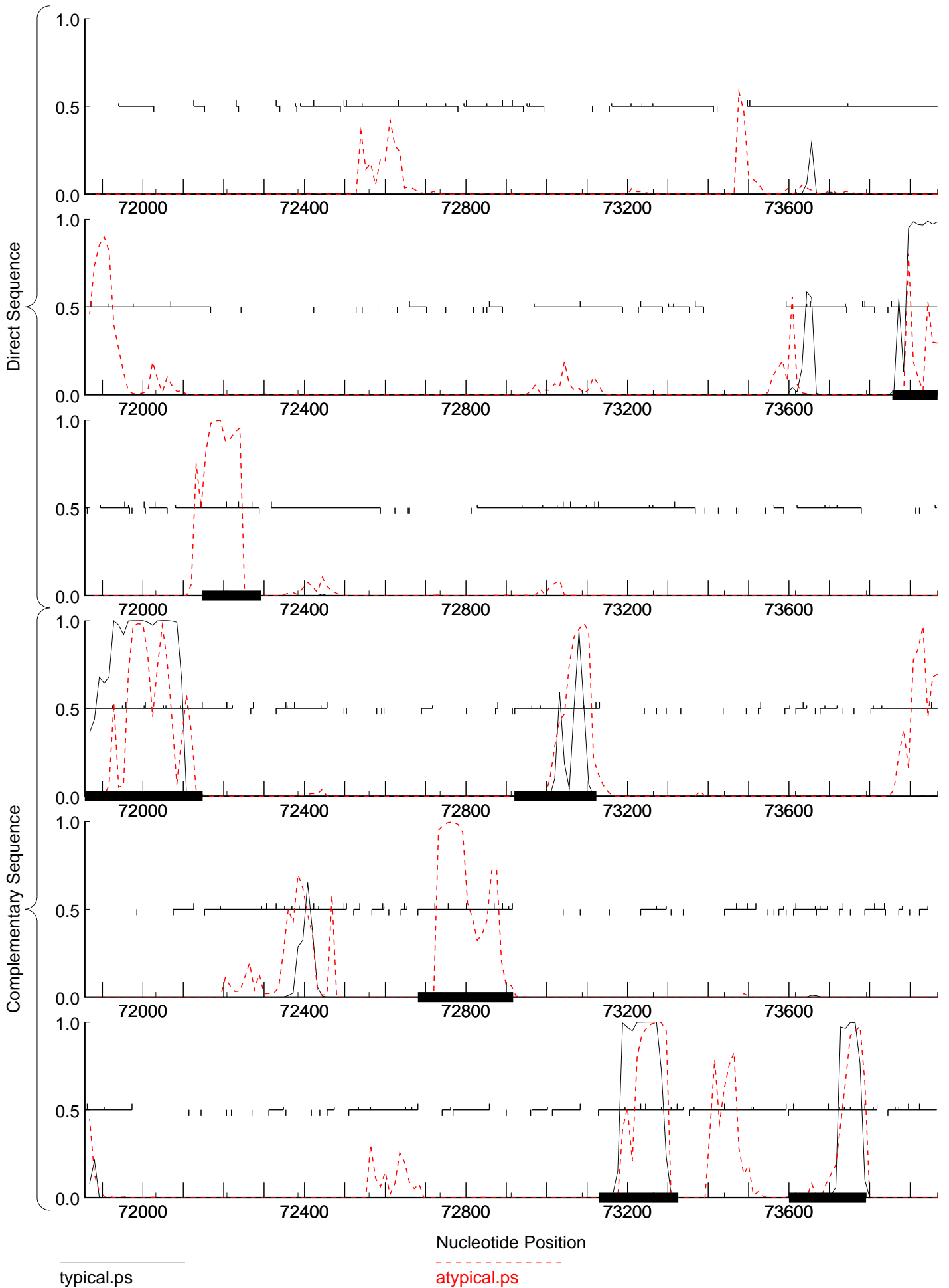


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