

GeneMark

Version 2.5p (09.08.06)

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PROGRAM INFORMATION

Sequence : Mycobacterium phage Zulu complete sequence, 52499 bp including 10-base 3' overhang (CGGTGGG
Analysis Date : 11/23/17 at 6:09:23
Pages : 26
Sequence Length : 52499 bp
GC Content : 61.42%

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,

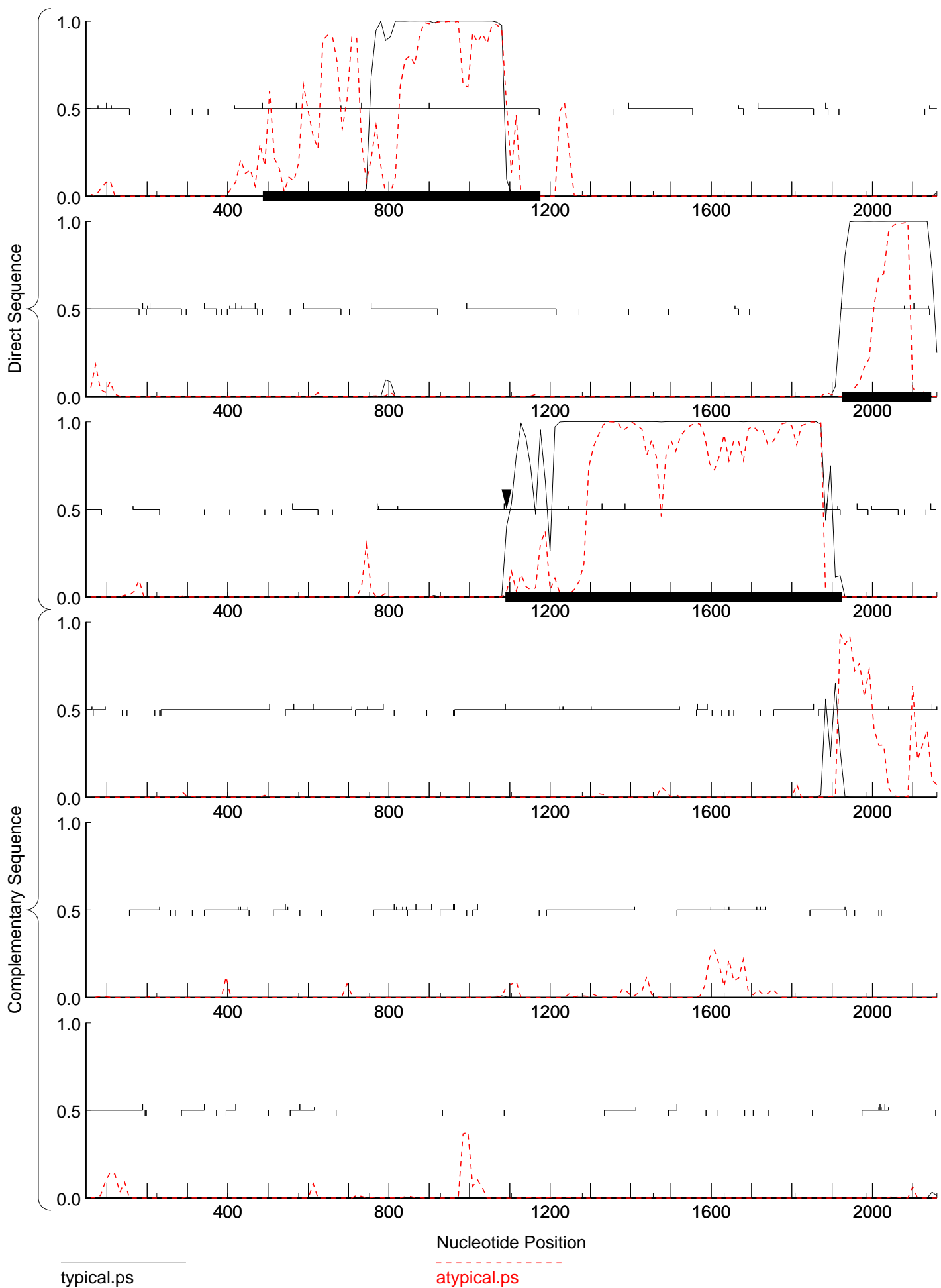
MATRIX INFORMATION

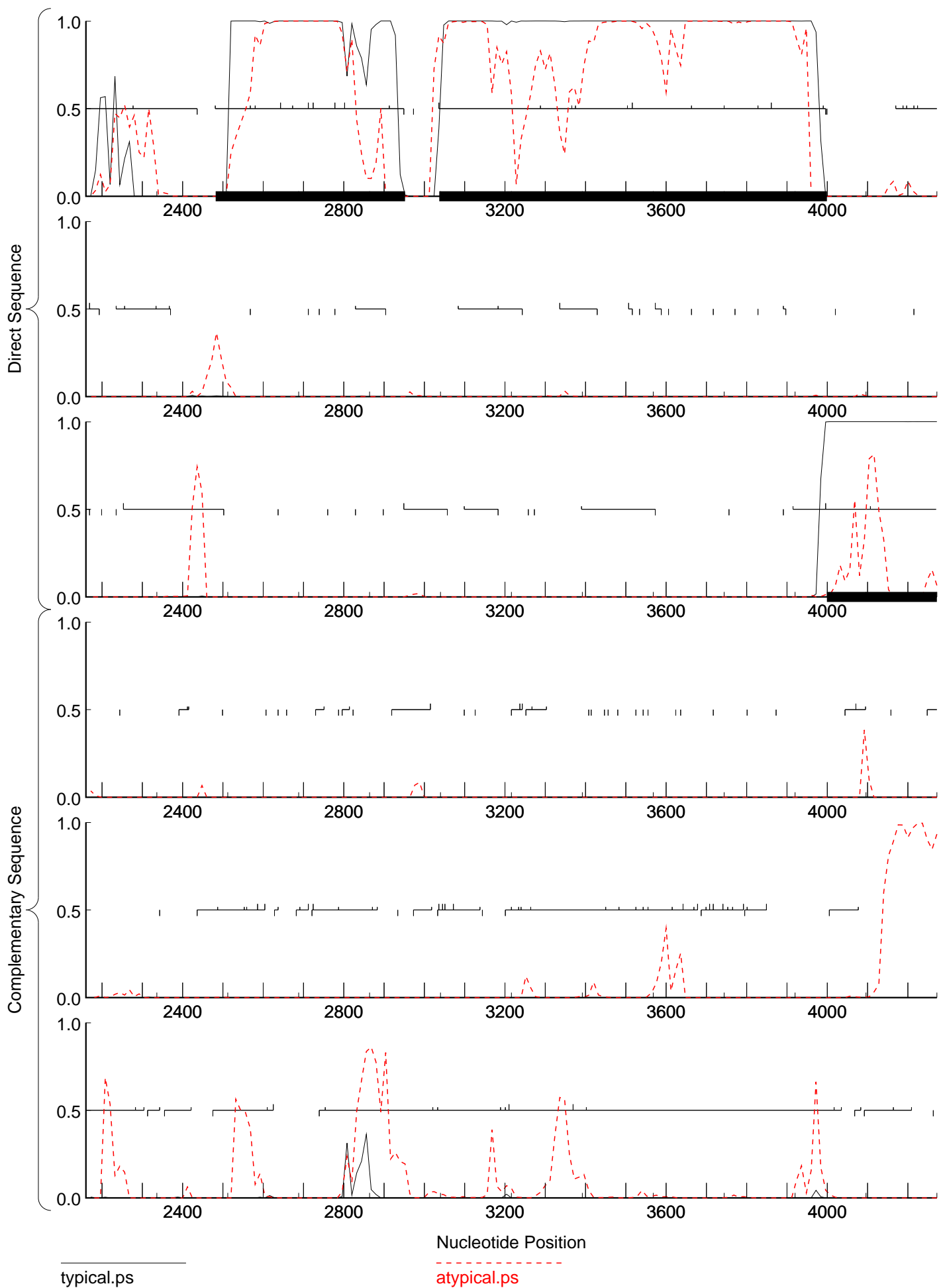
Matrix : -
Author : -
Order : 4

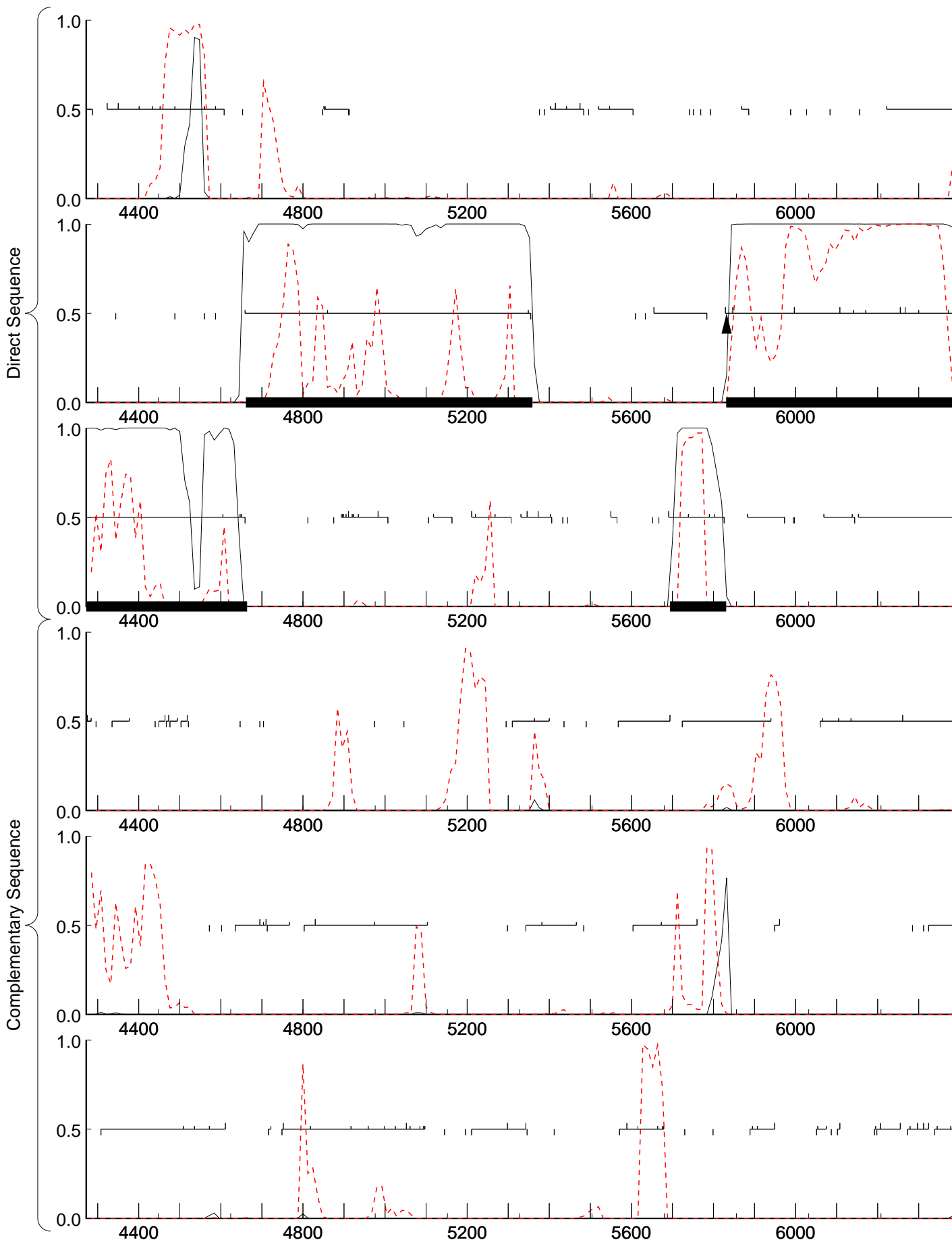
Send questions / comments to:
Dr. M. Borodovsky
Georgia Institute of Technology
School of Biology
Atlanta, GA 30332-0230

Matrix notes & comments

Training set derived by GeneMarkS, 4.30 October 2014
Thu Nov 23 06:09:23 2017

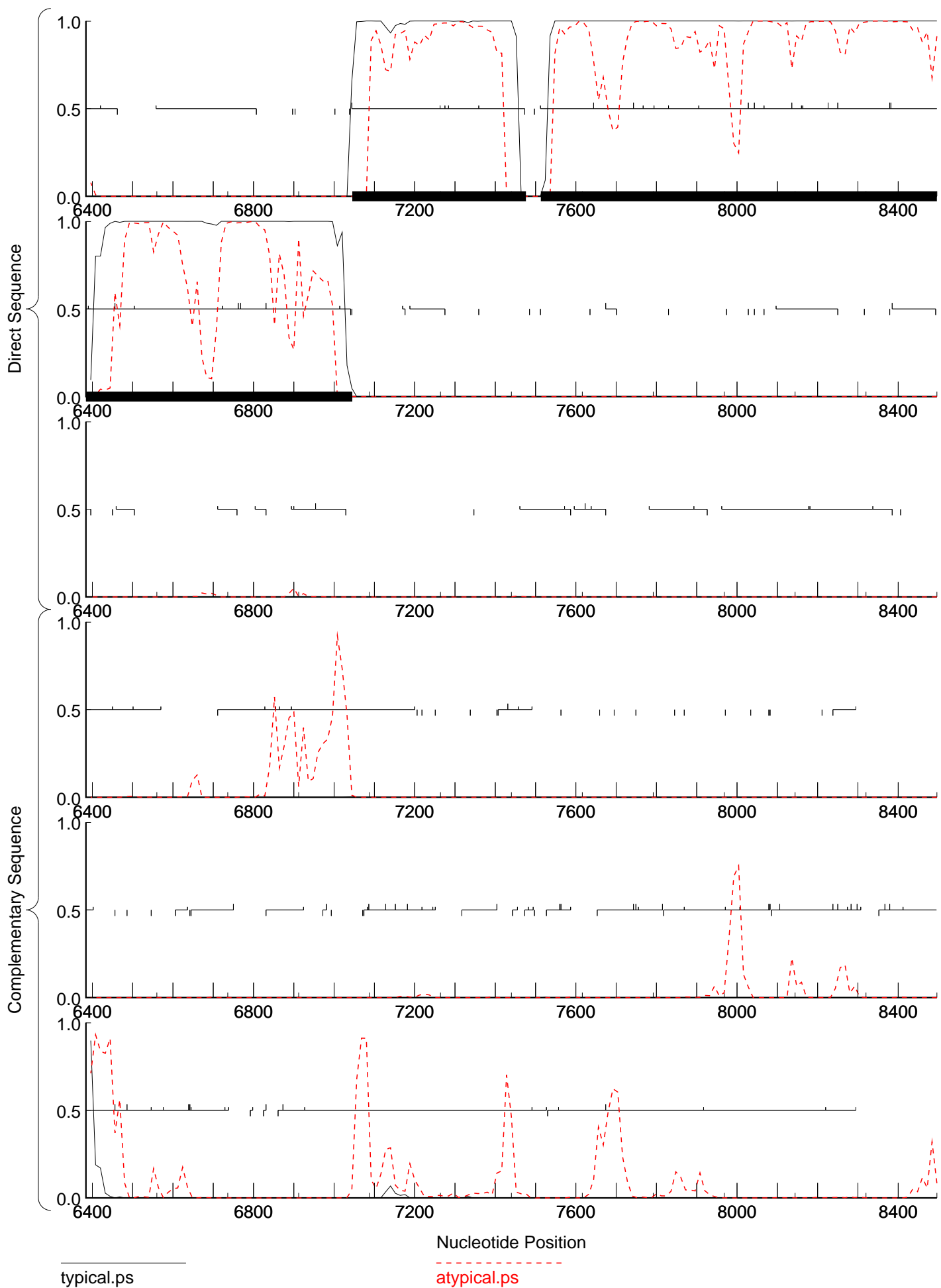


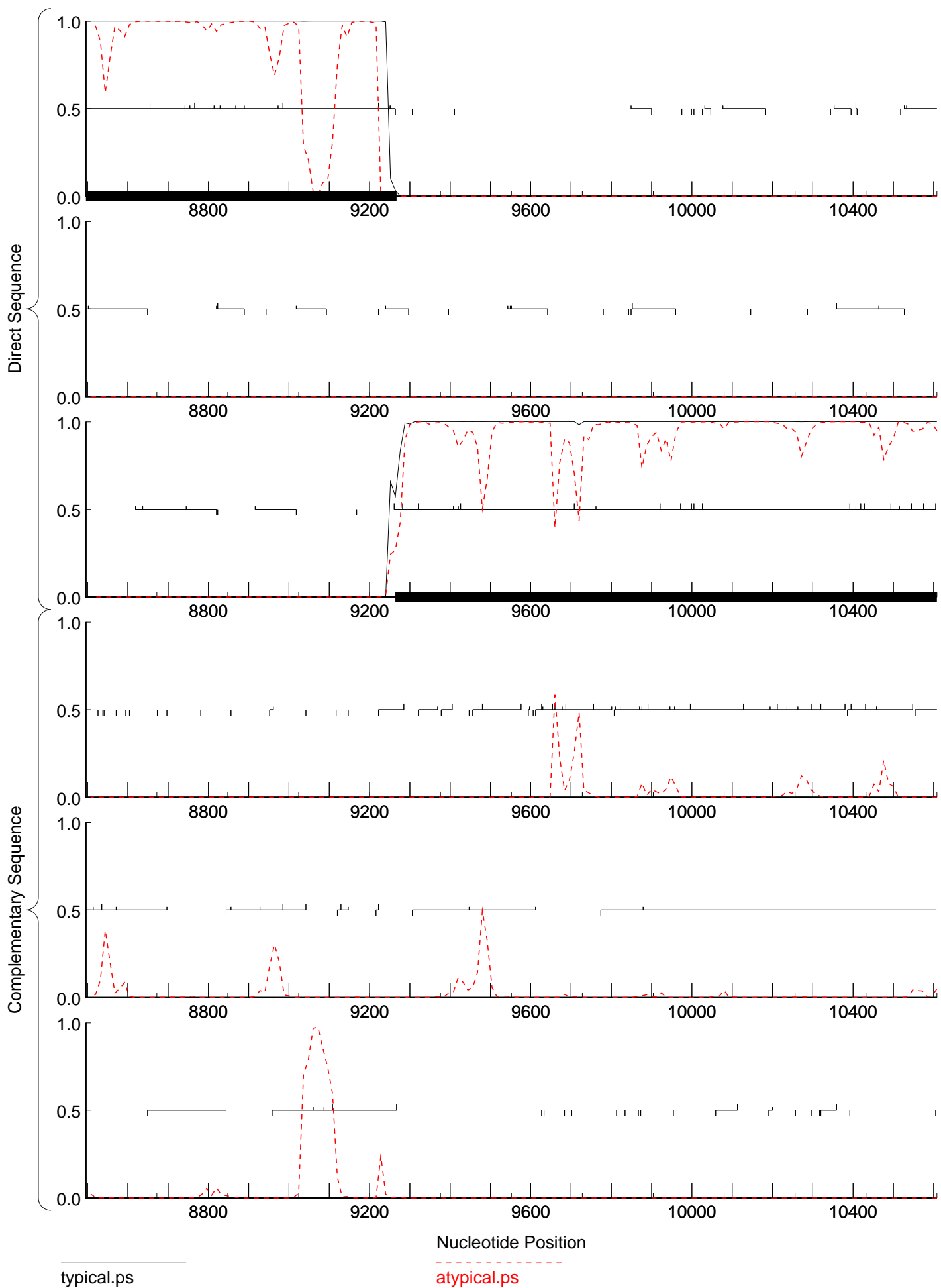


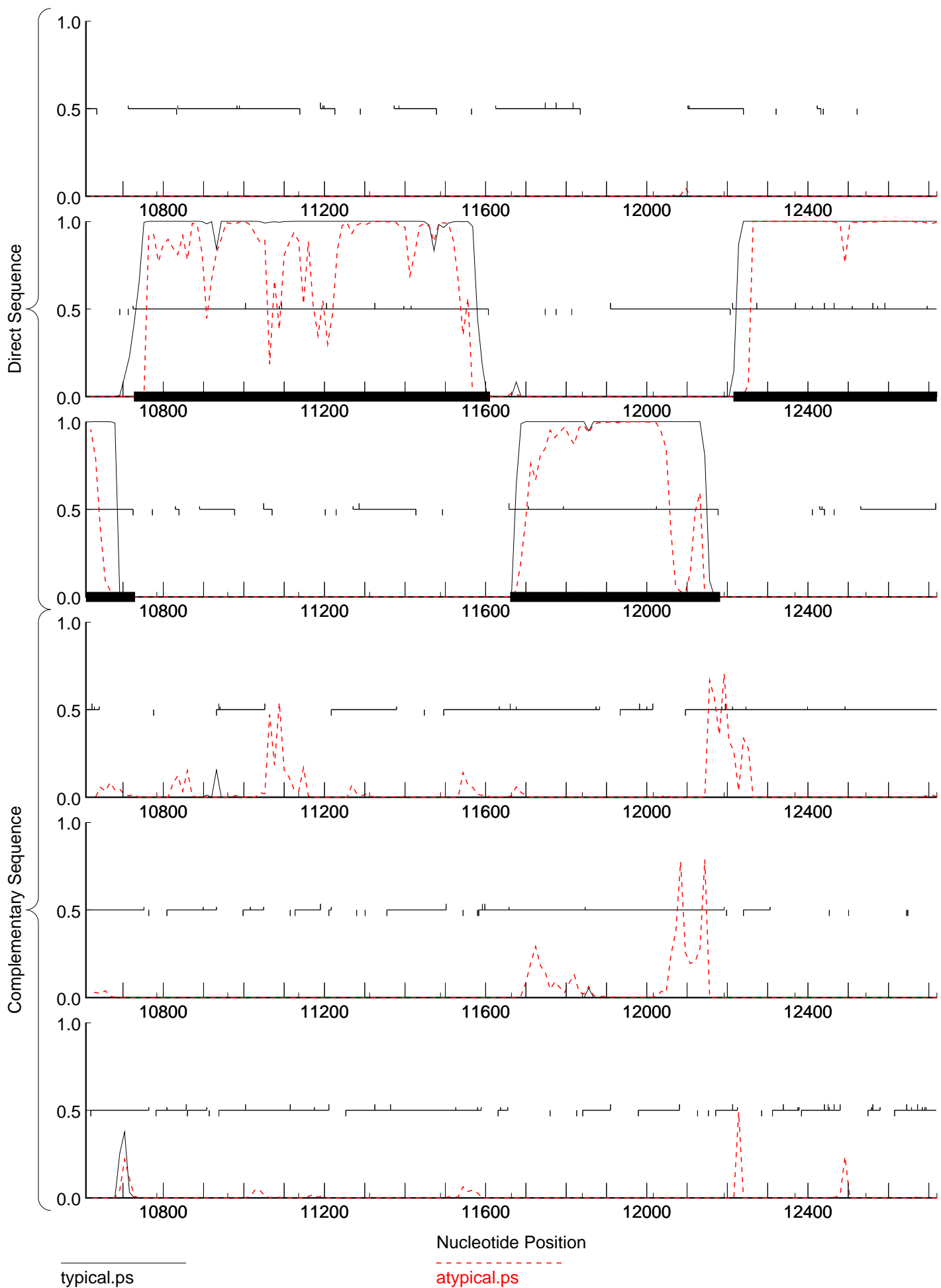


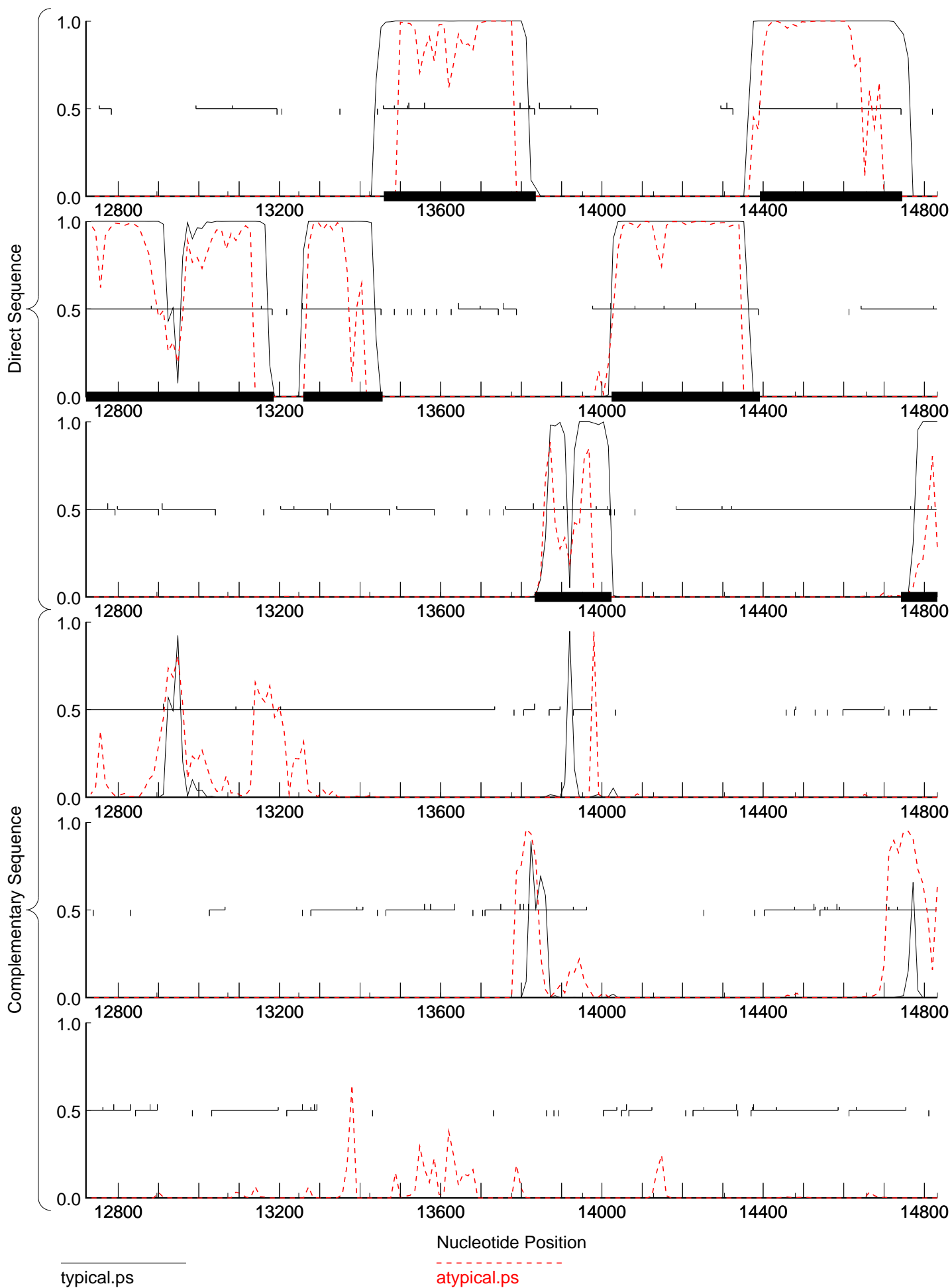
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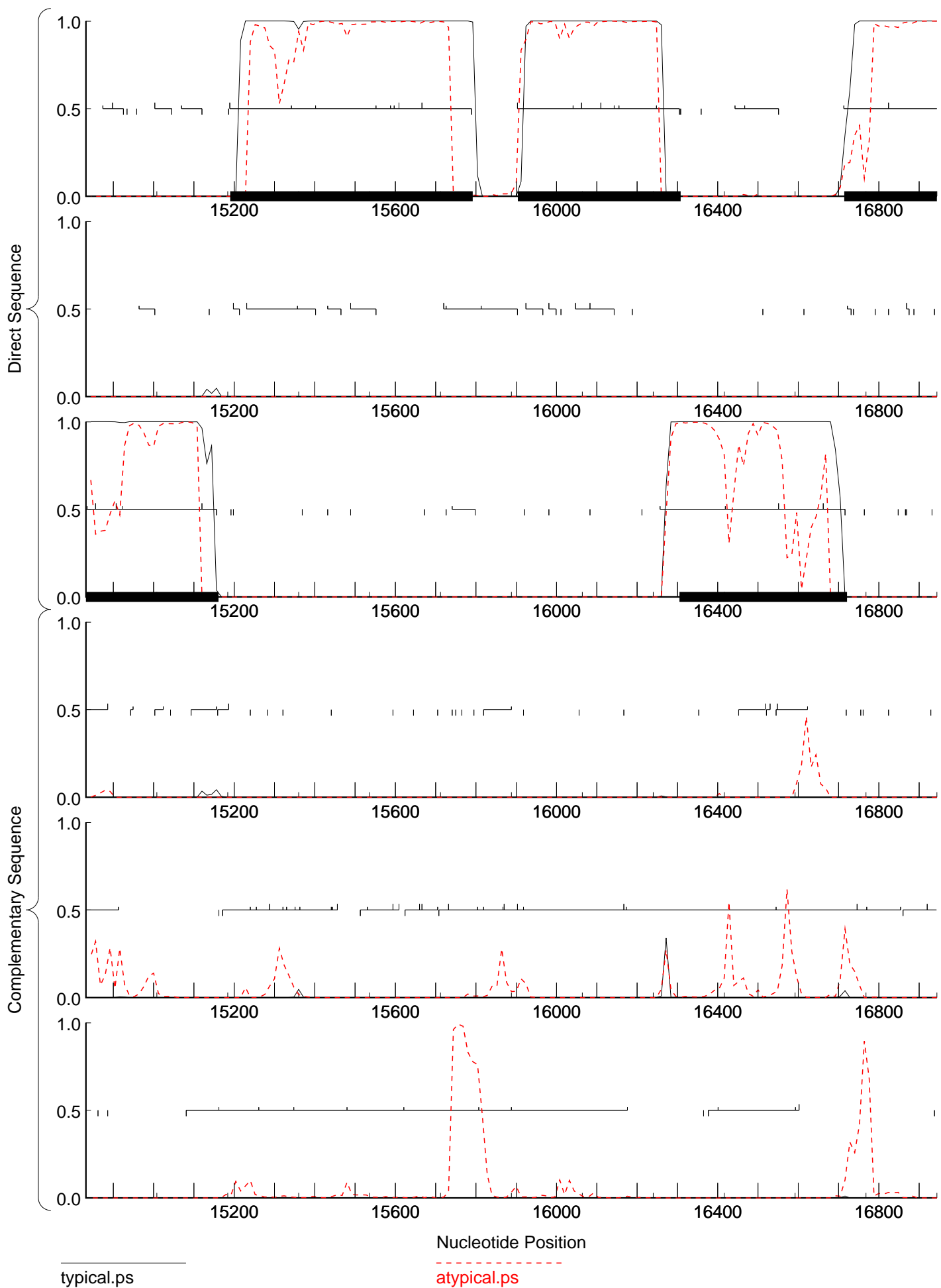
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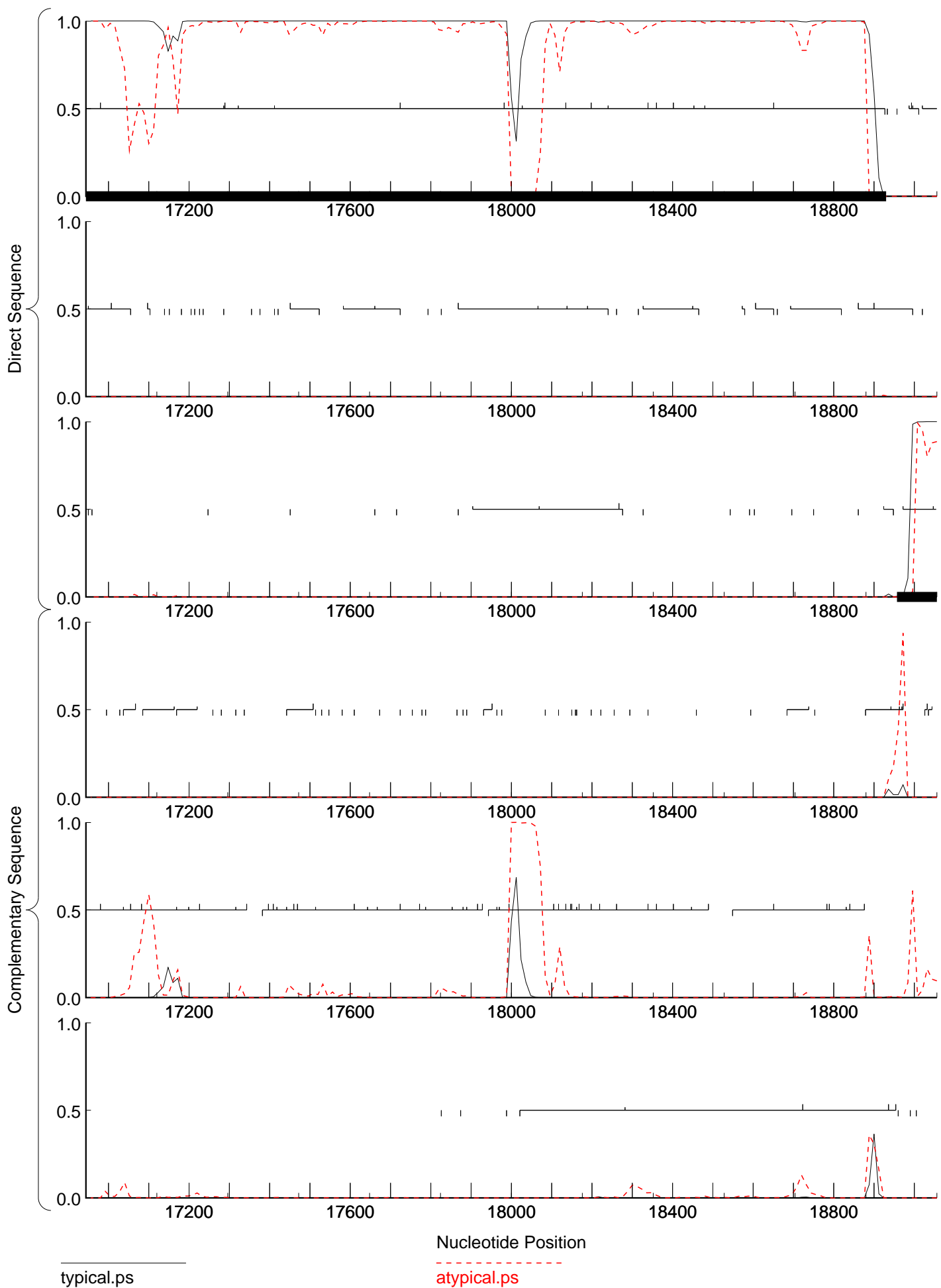






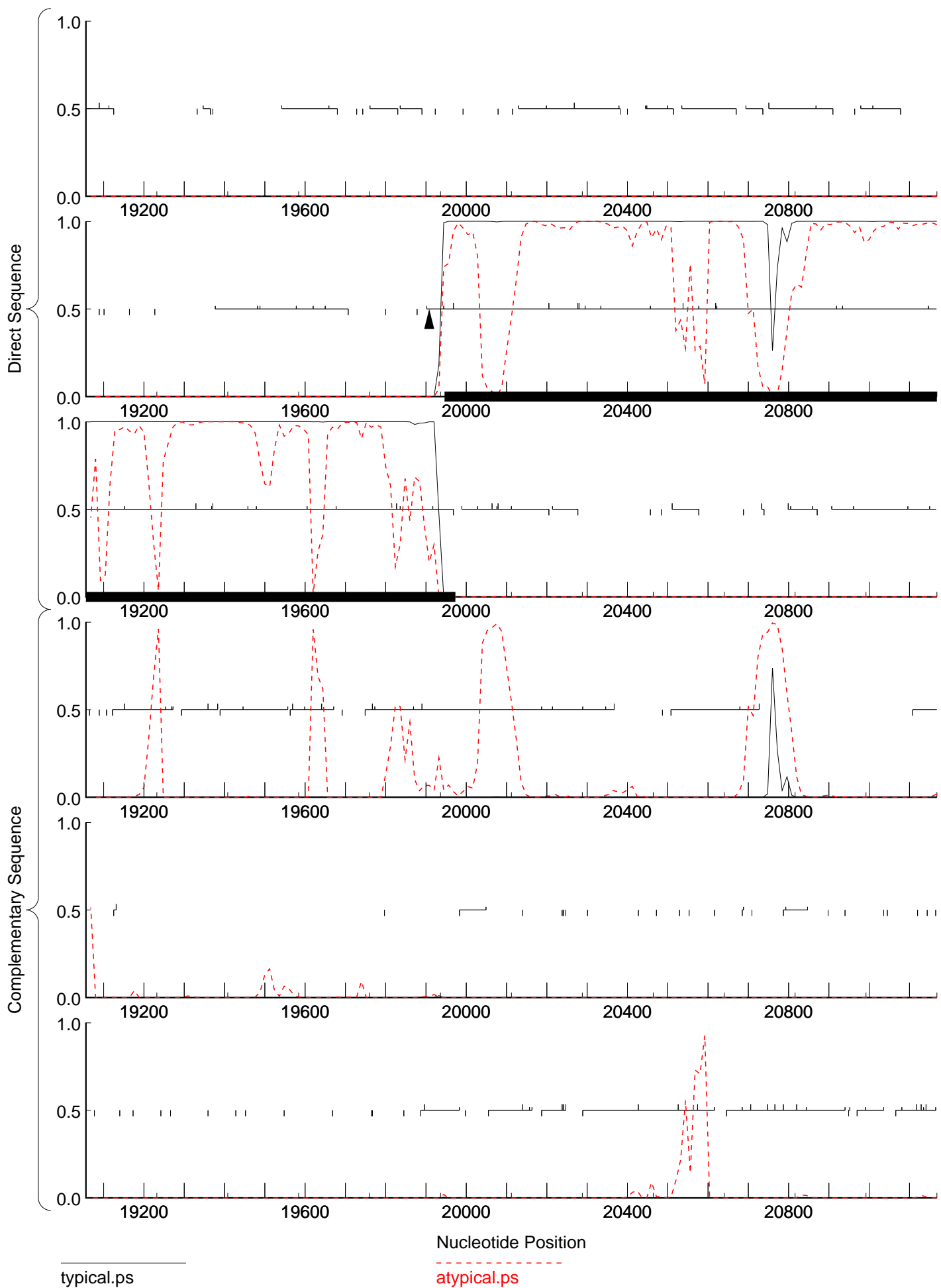
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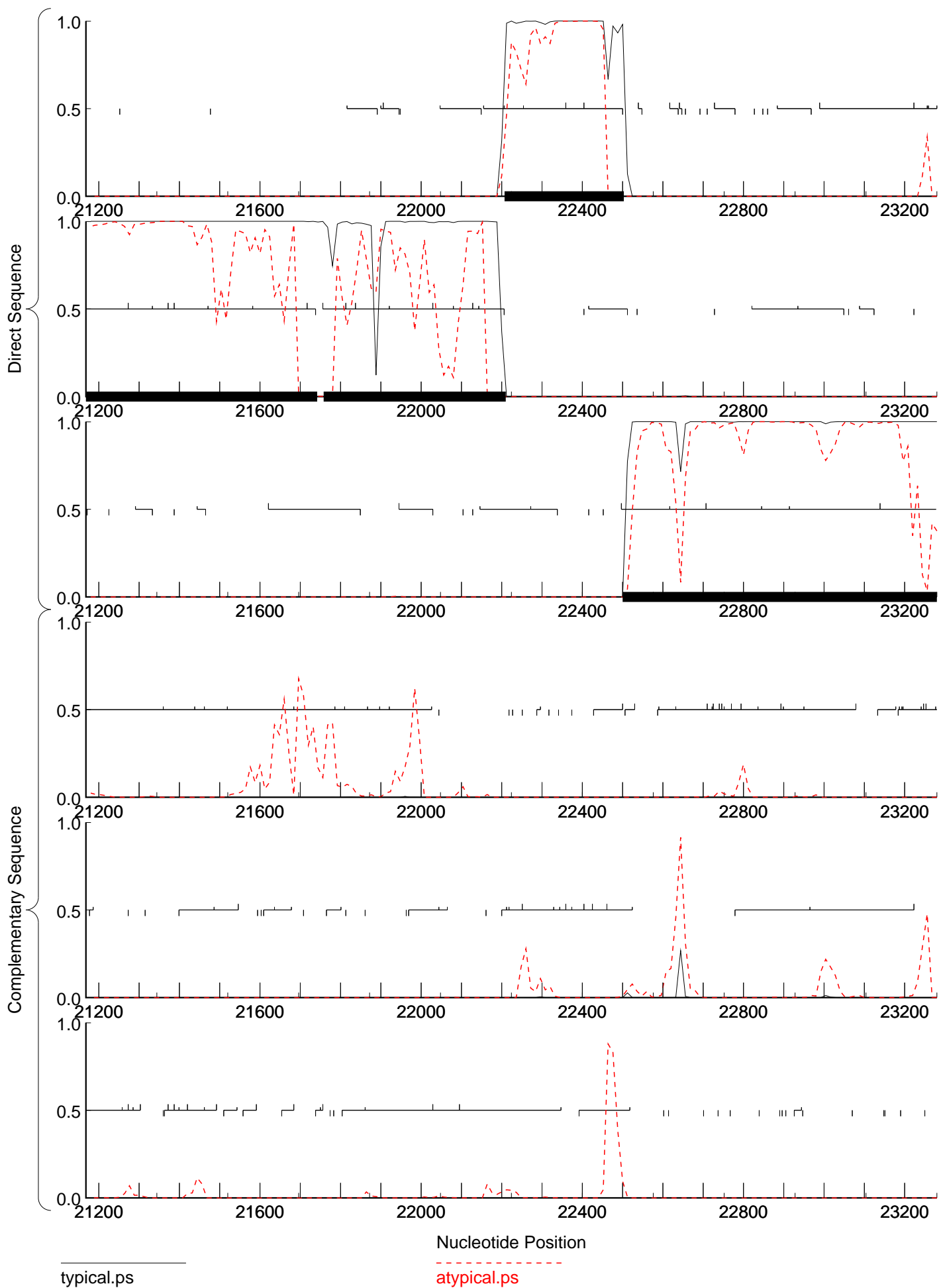
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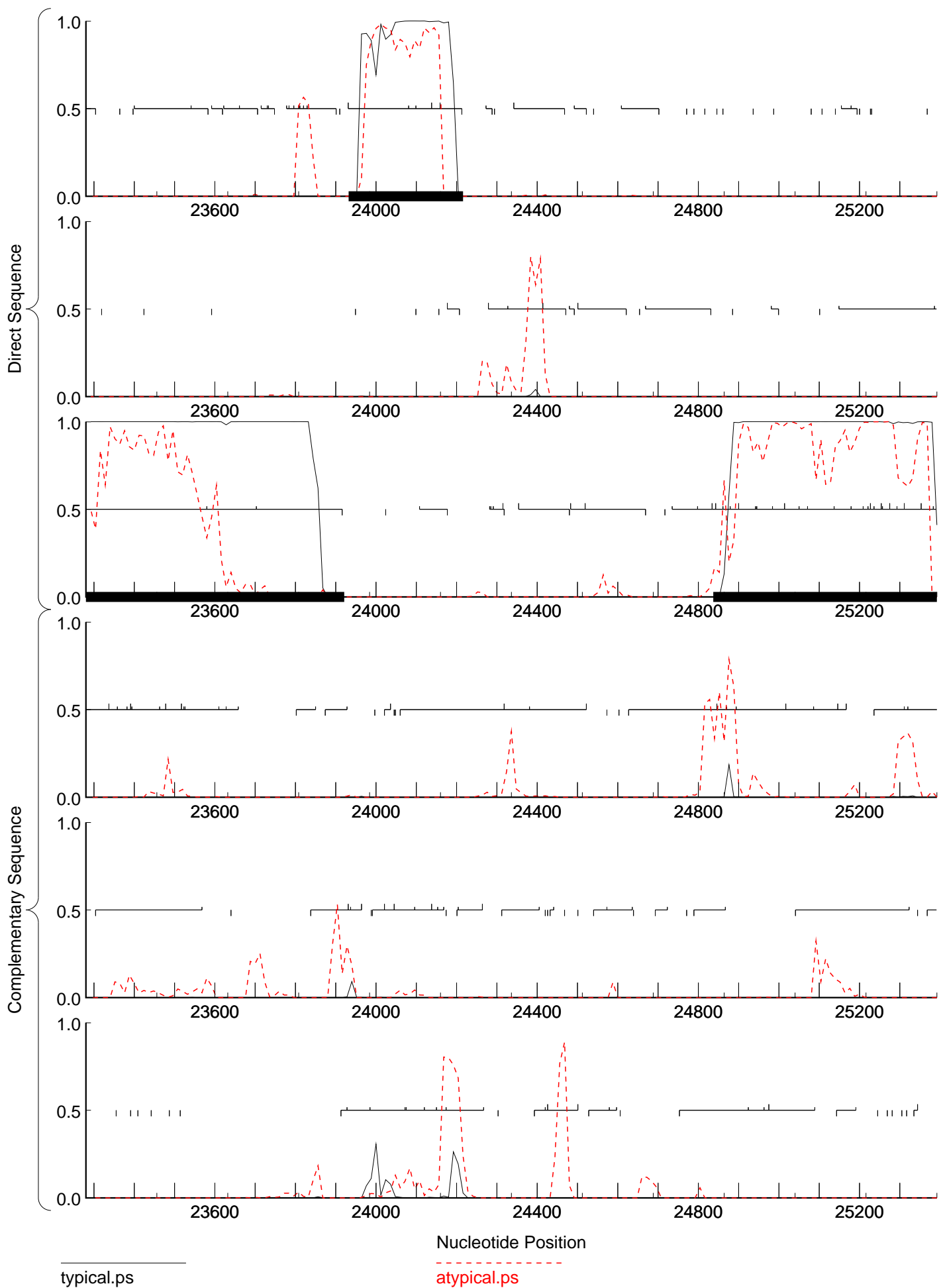


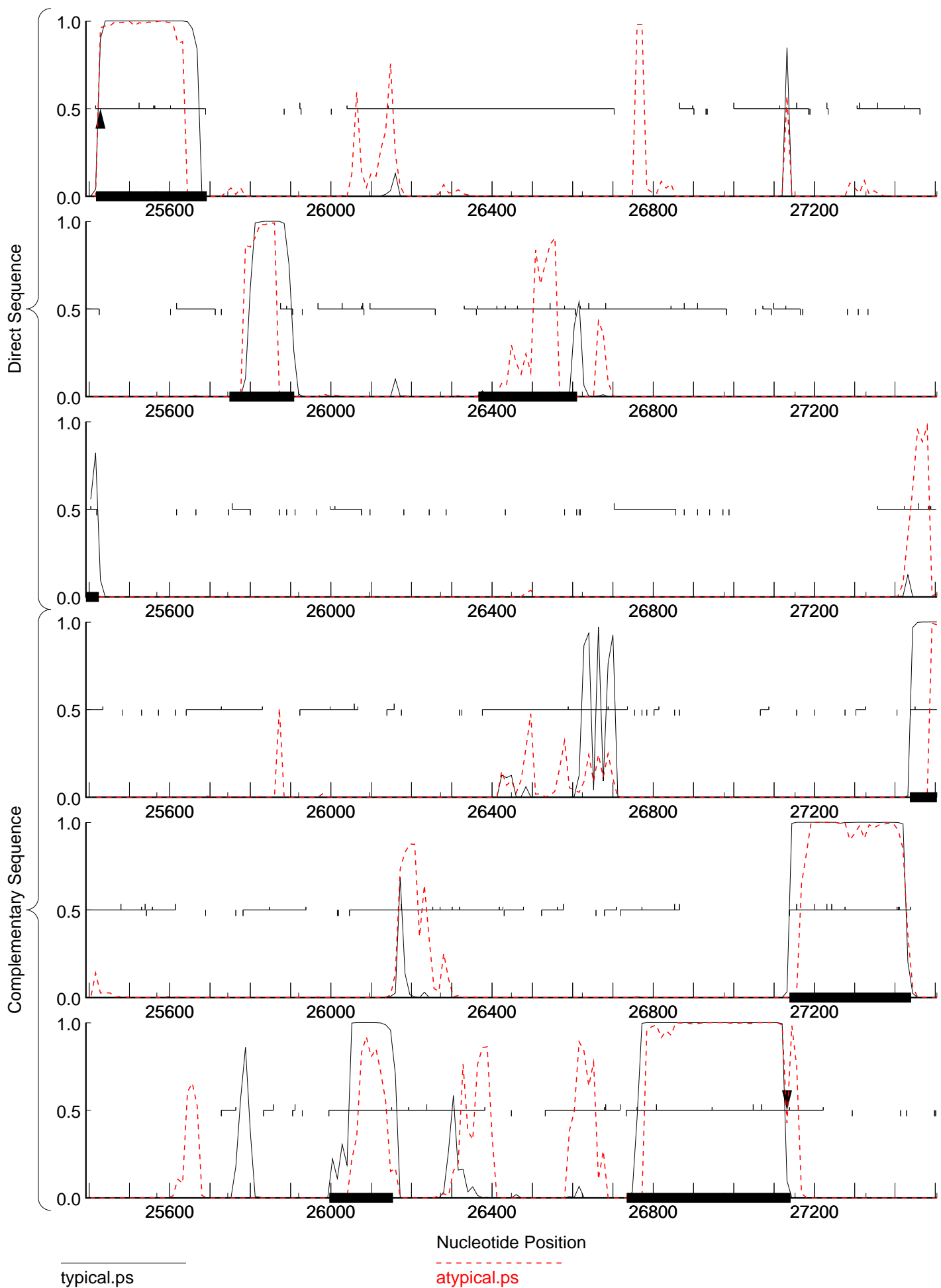
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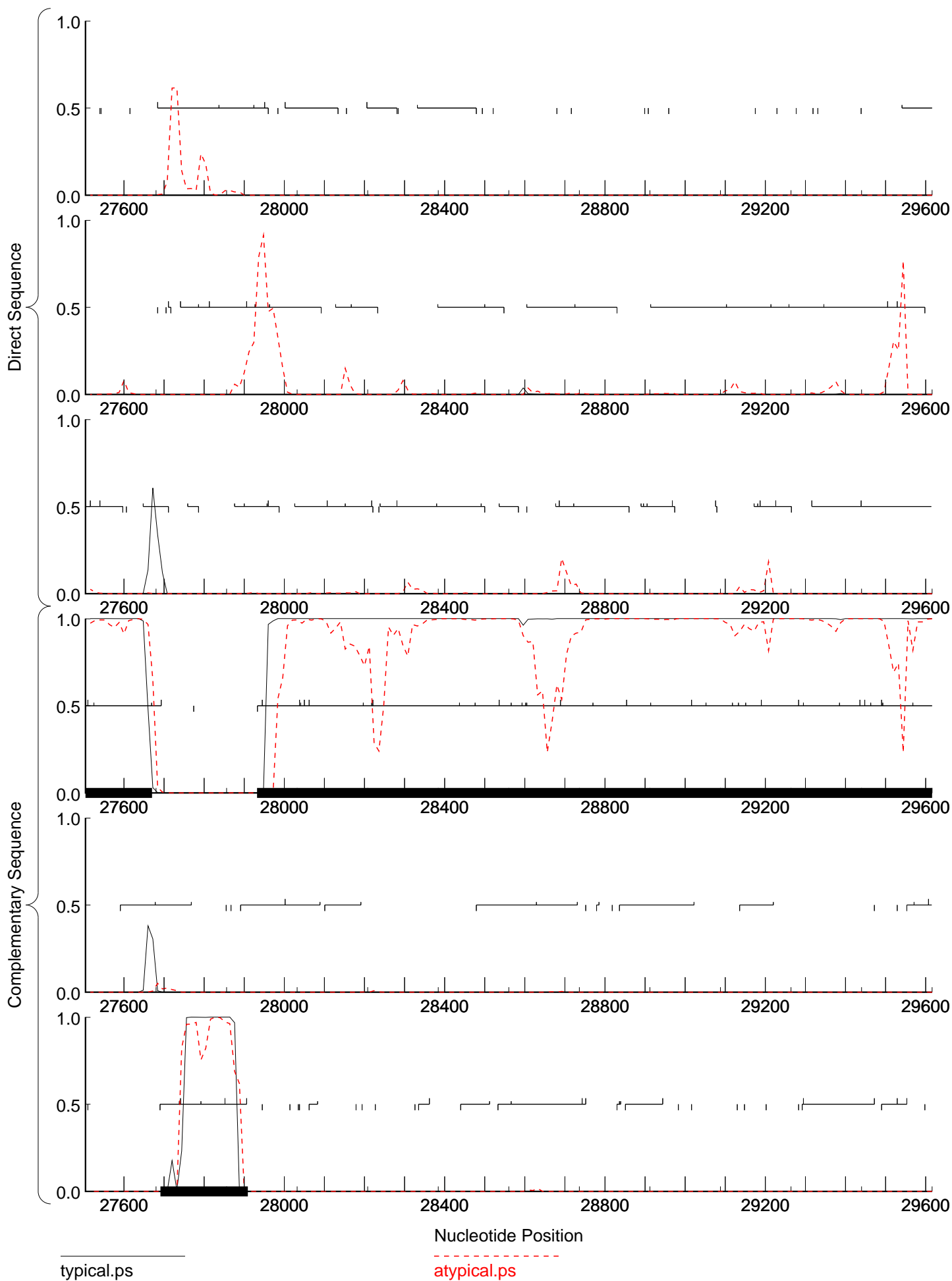
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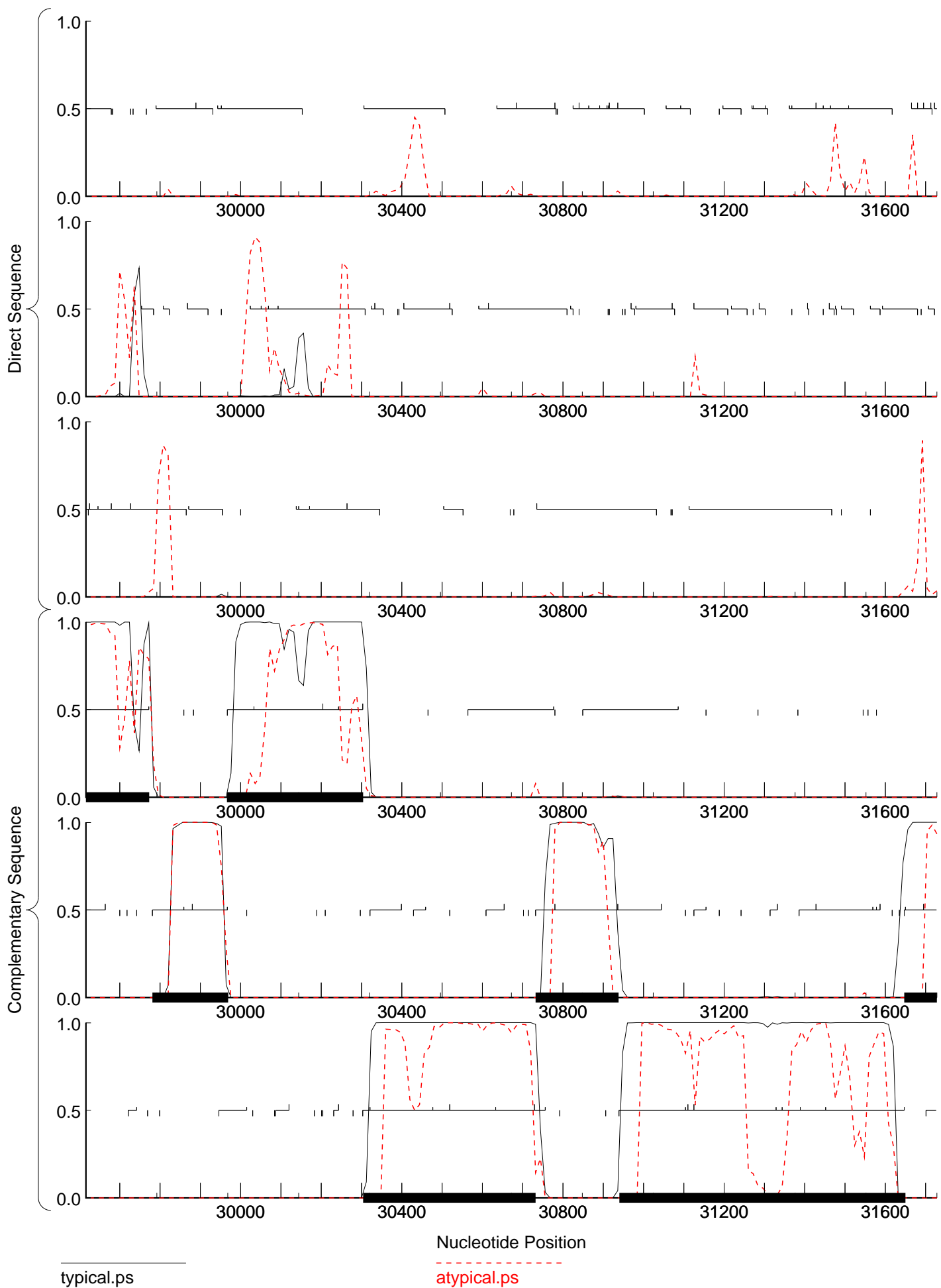


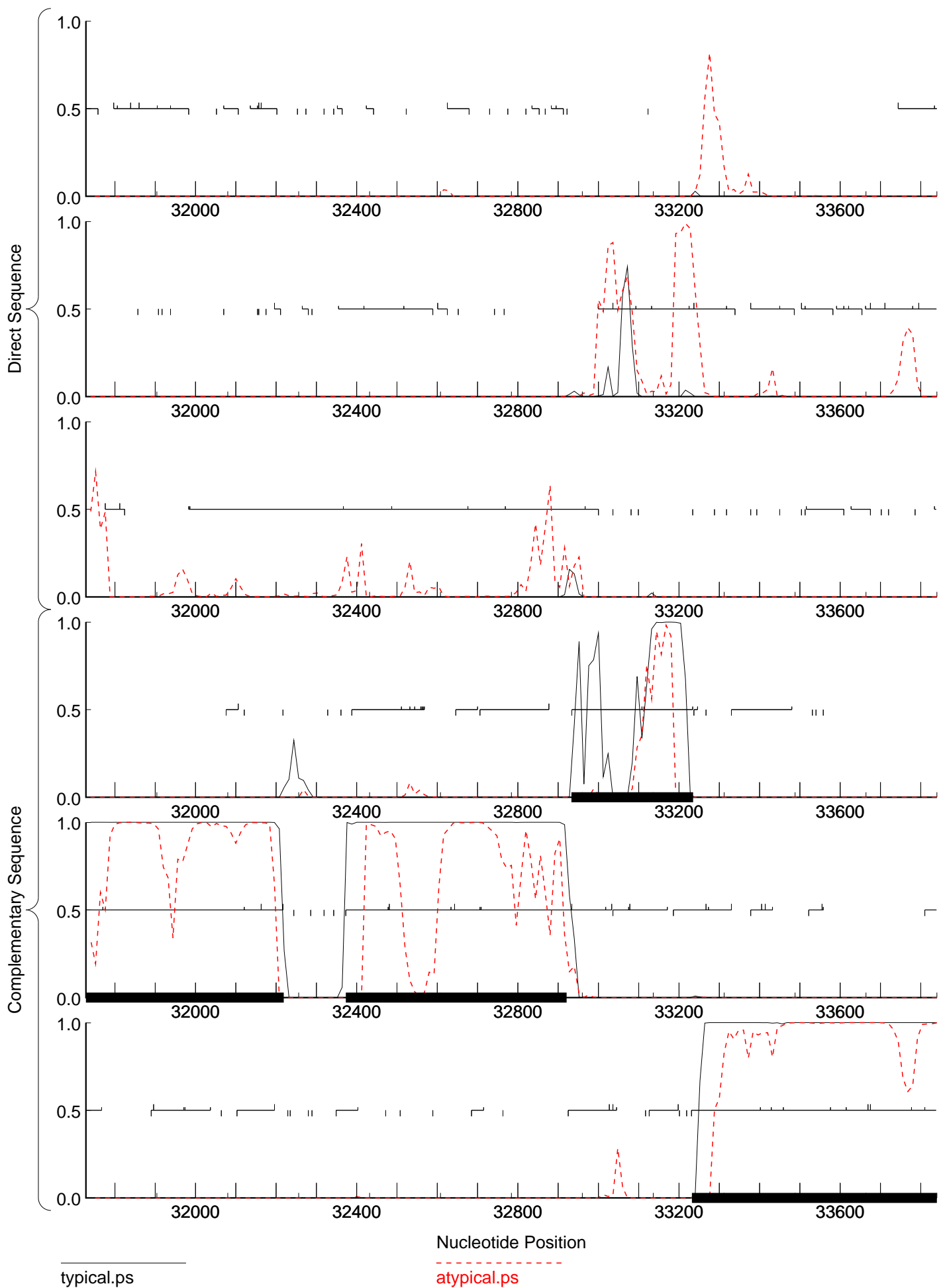




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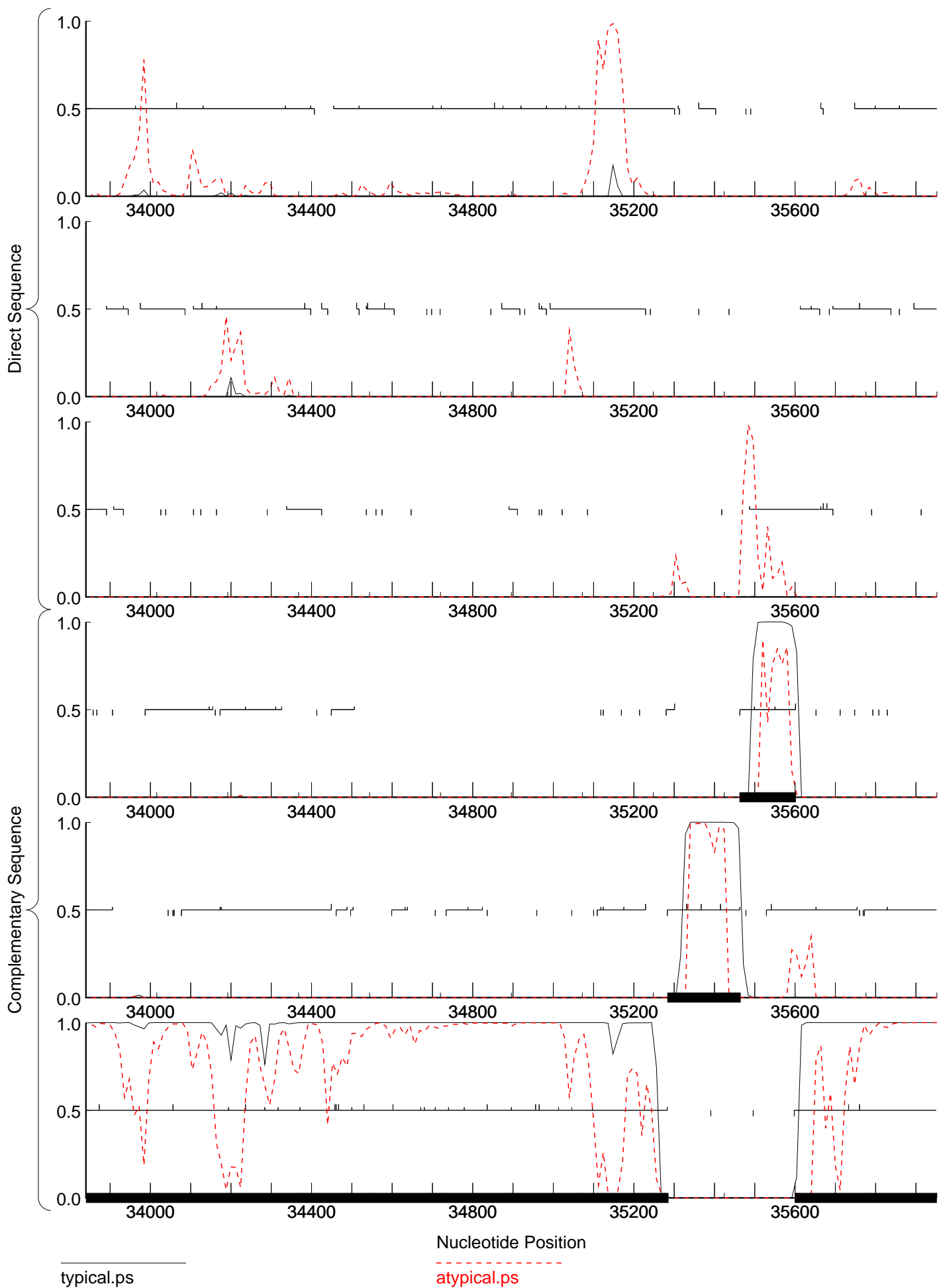
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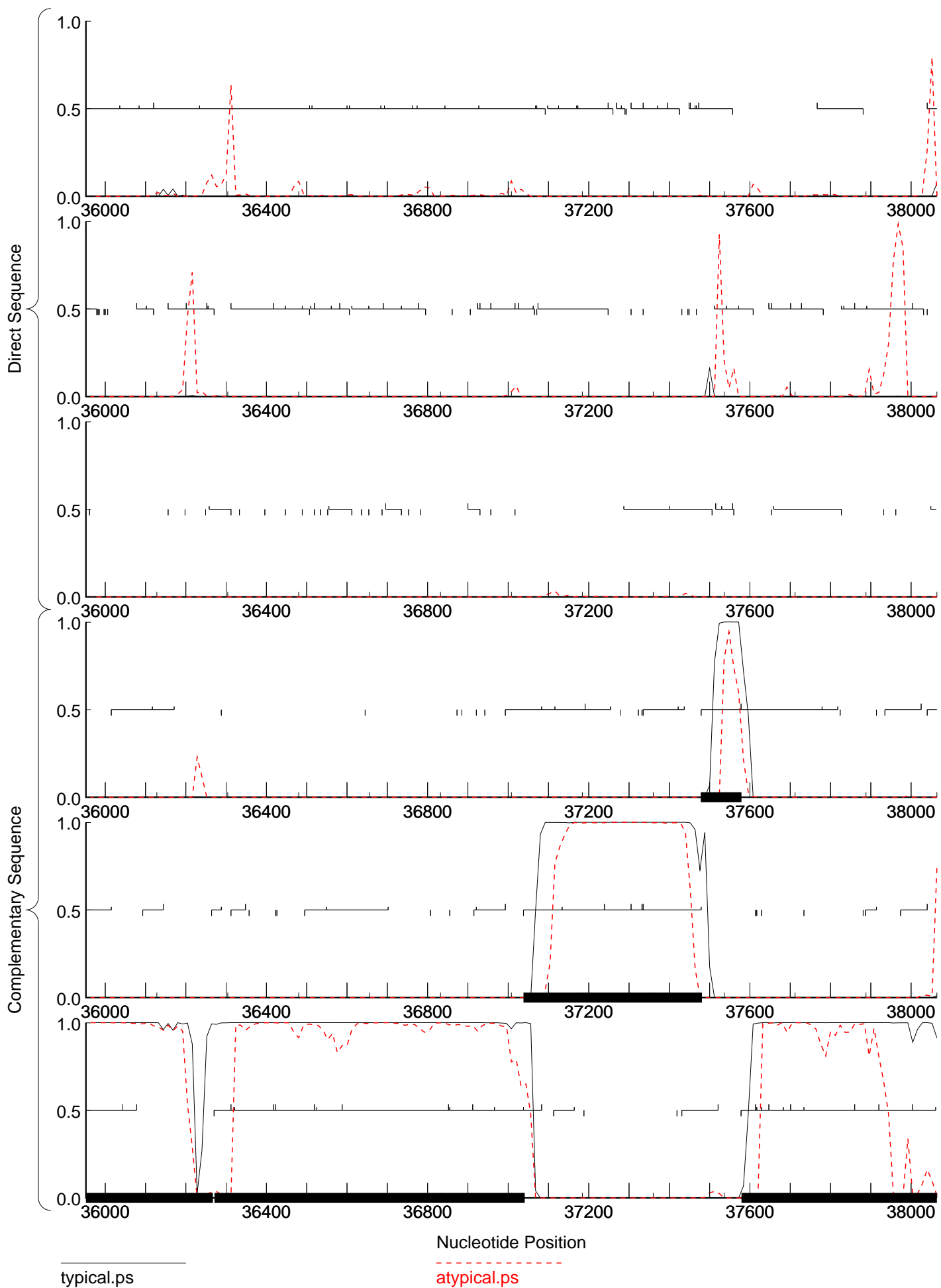
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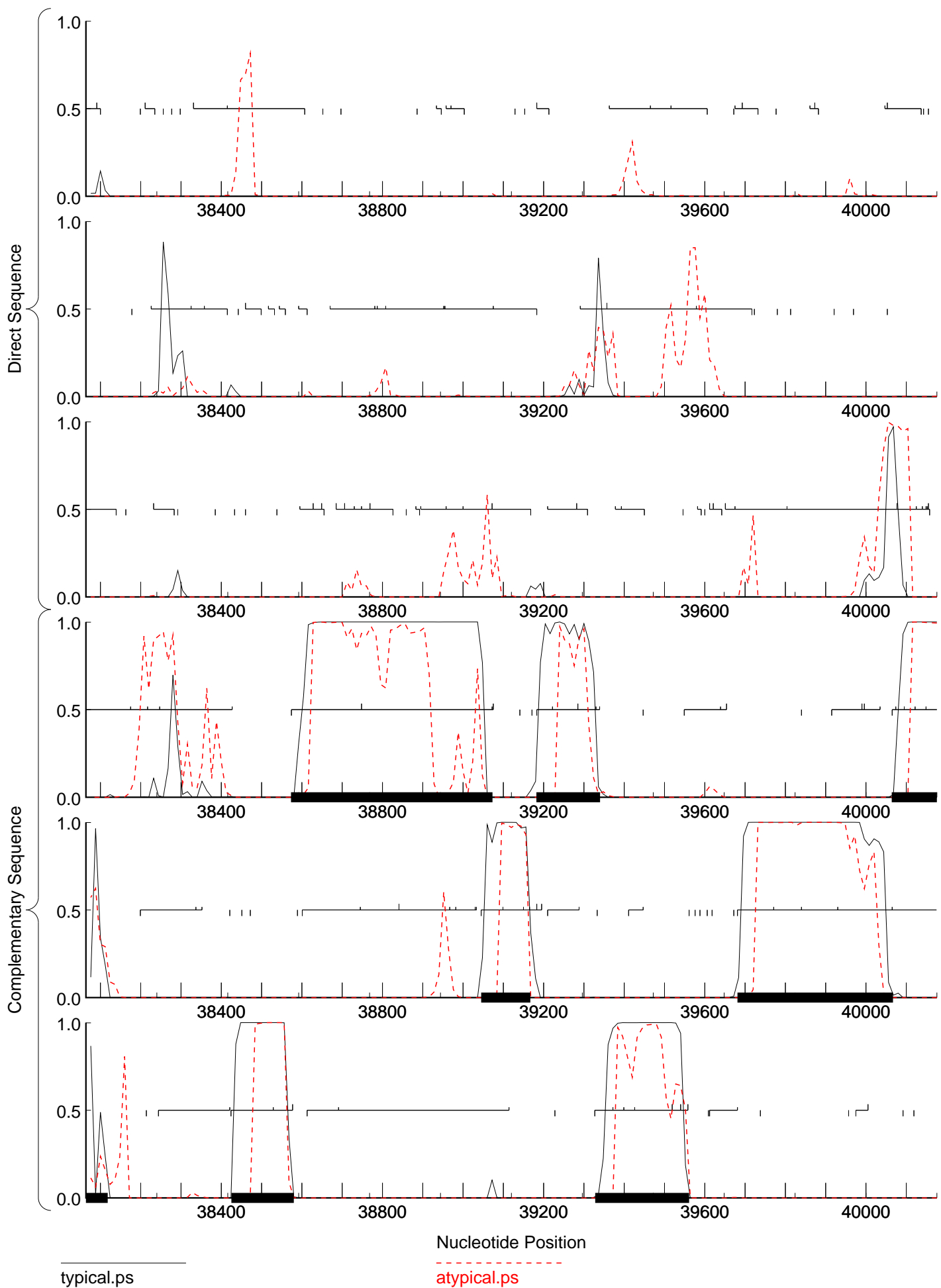
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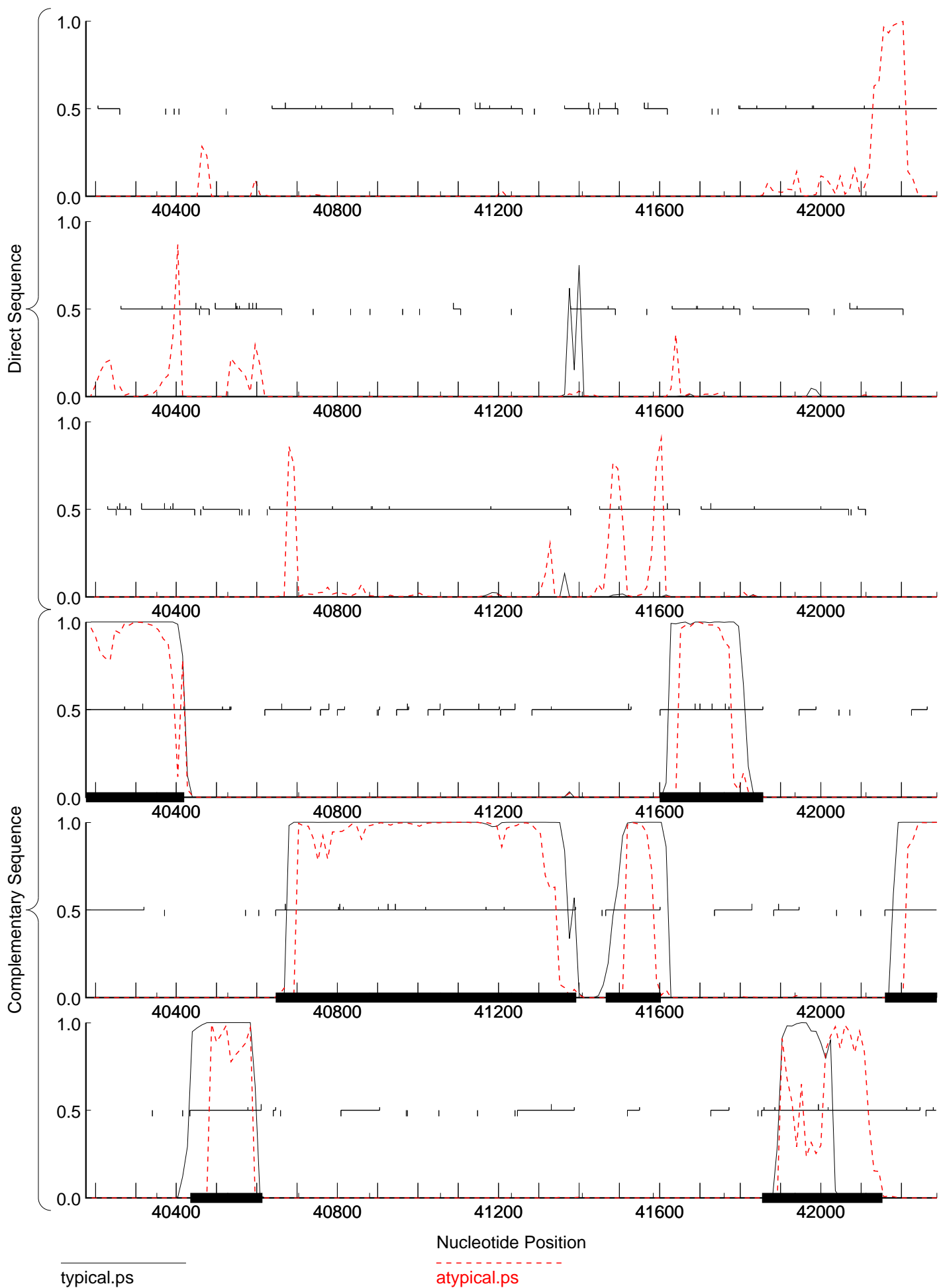
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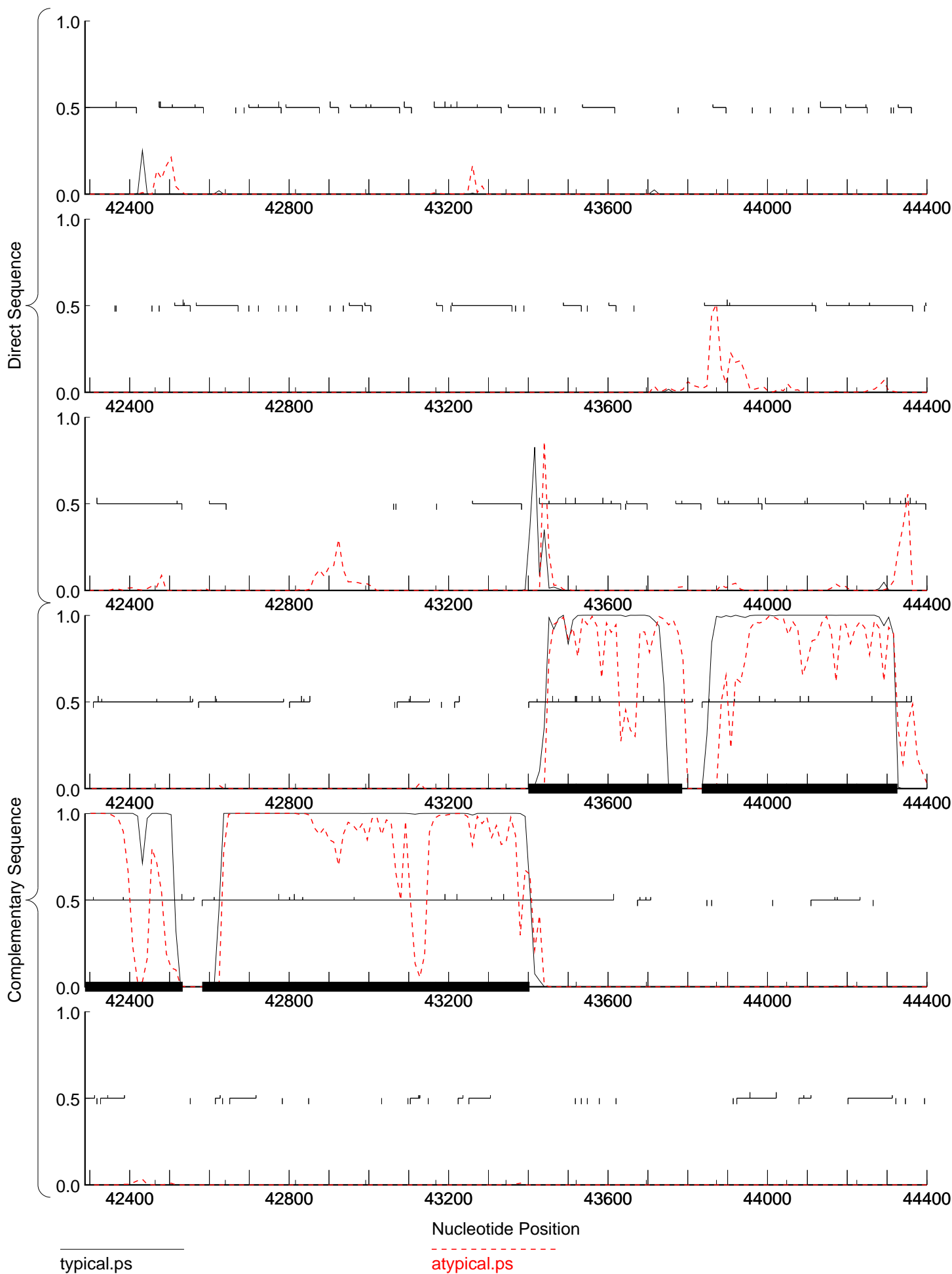


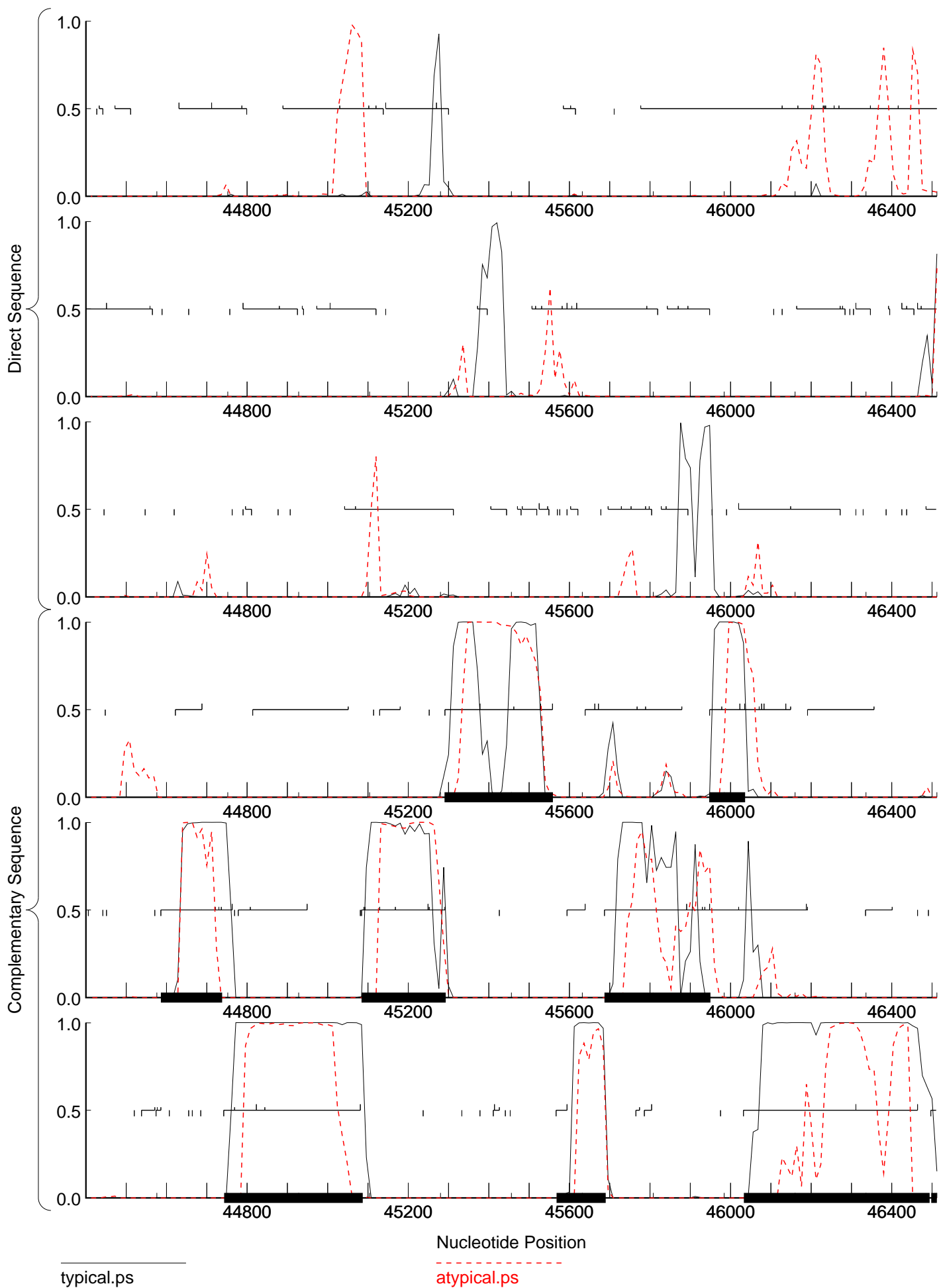


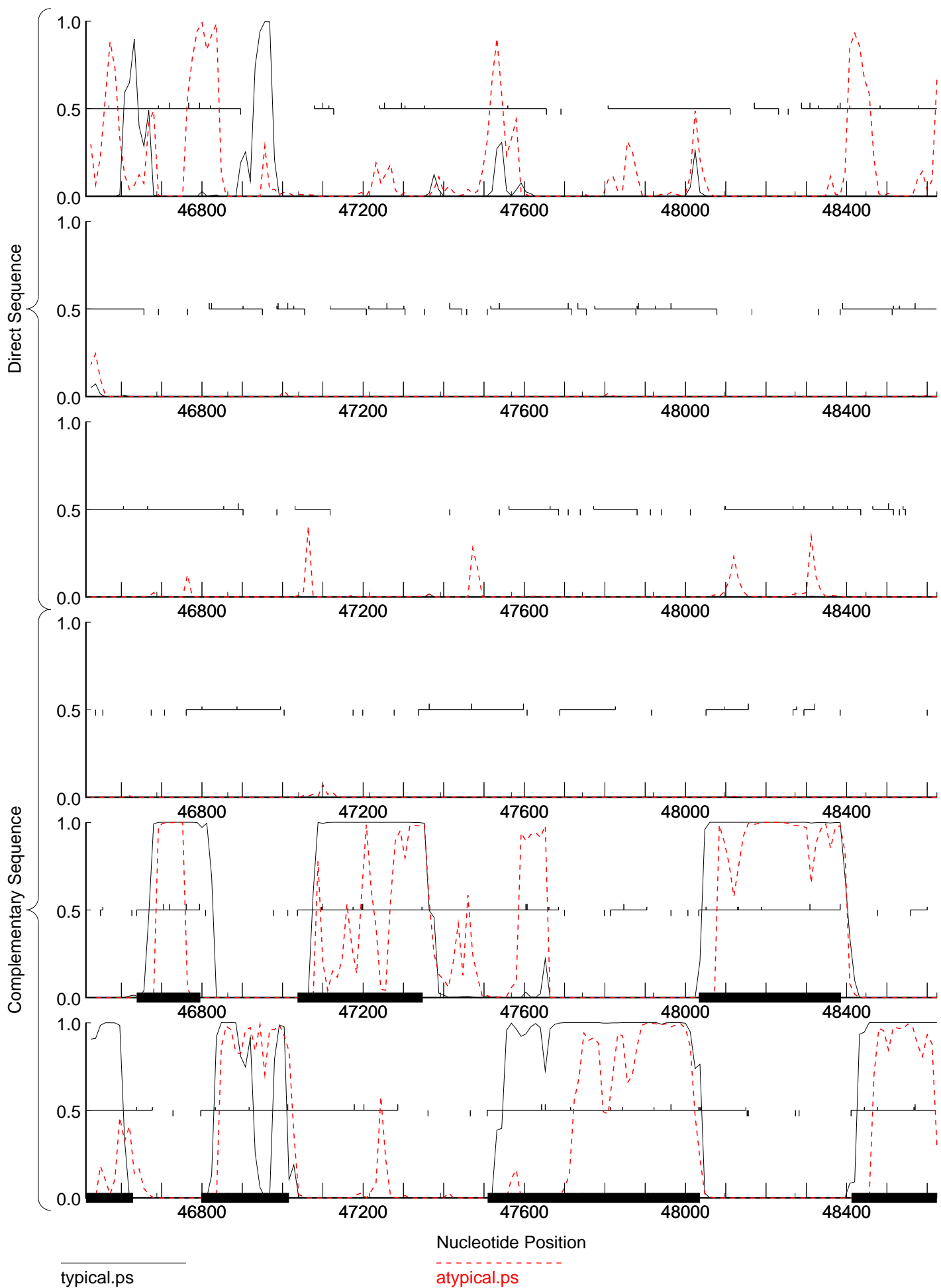
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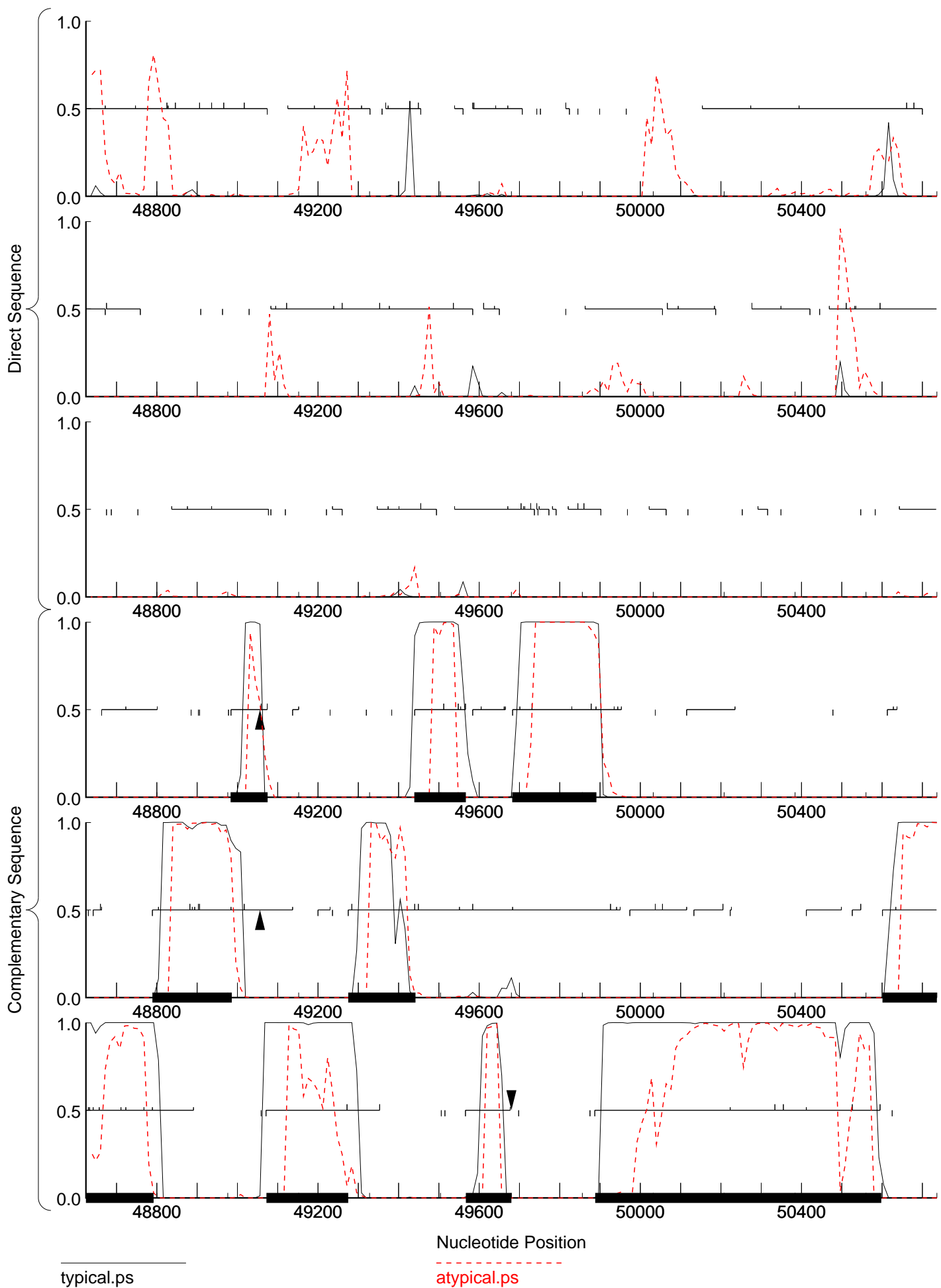


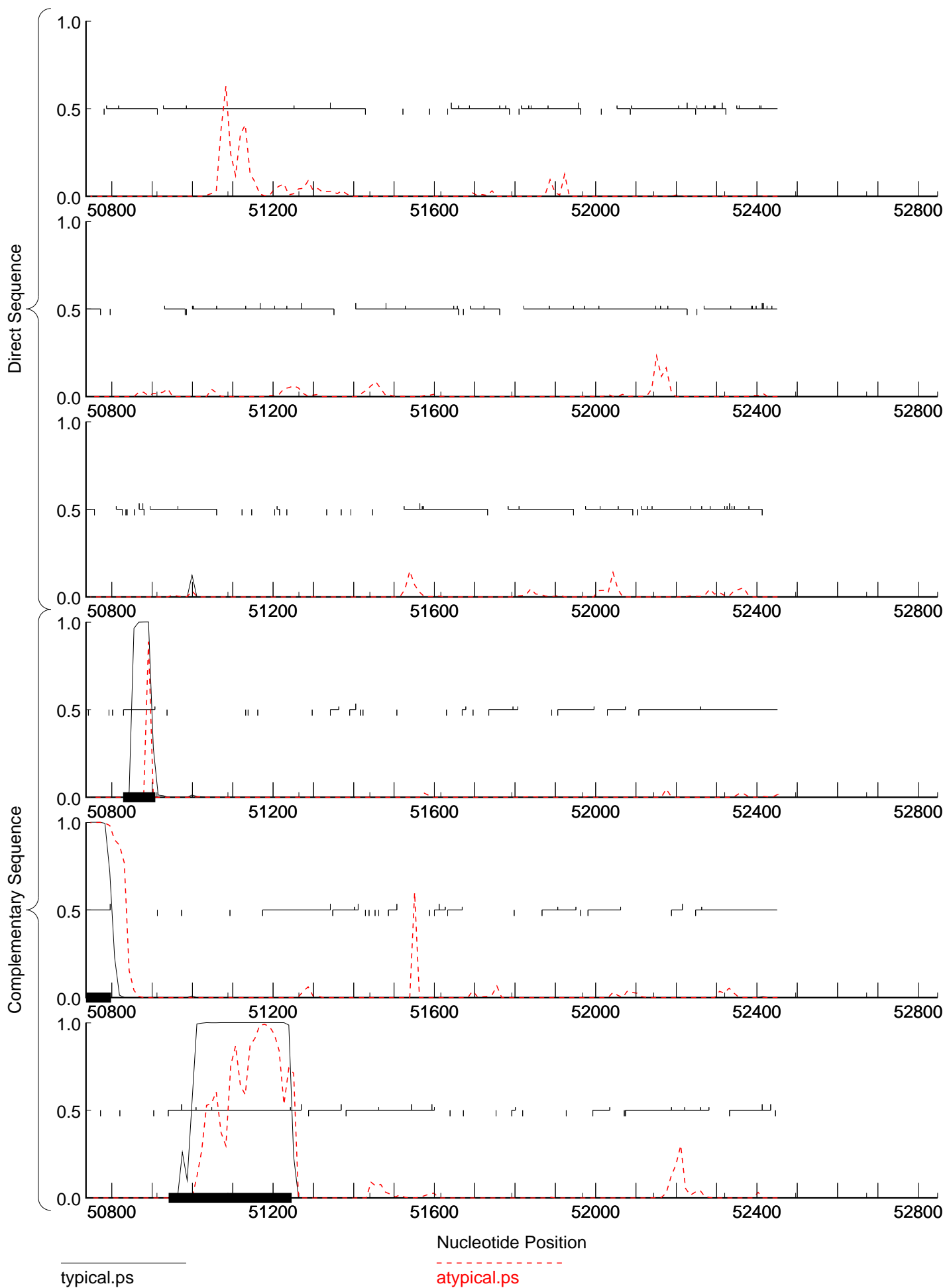




typical.ps

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typical.ps

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