

GeneMark

Version 2.5p (09.08.06)

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PROGRAM INFORMATION

Sequence : Mycobacterium phage BigMau complete sequence, 52632 bp including 10-base 3' overhang (CGGAT
Analysis Date : 6/24/18 at 5:13:00
Pages : 26
Sequence Length : 52632 bp
GC Content : 63.73%

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,

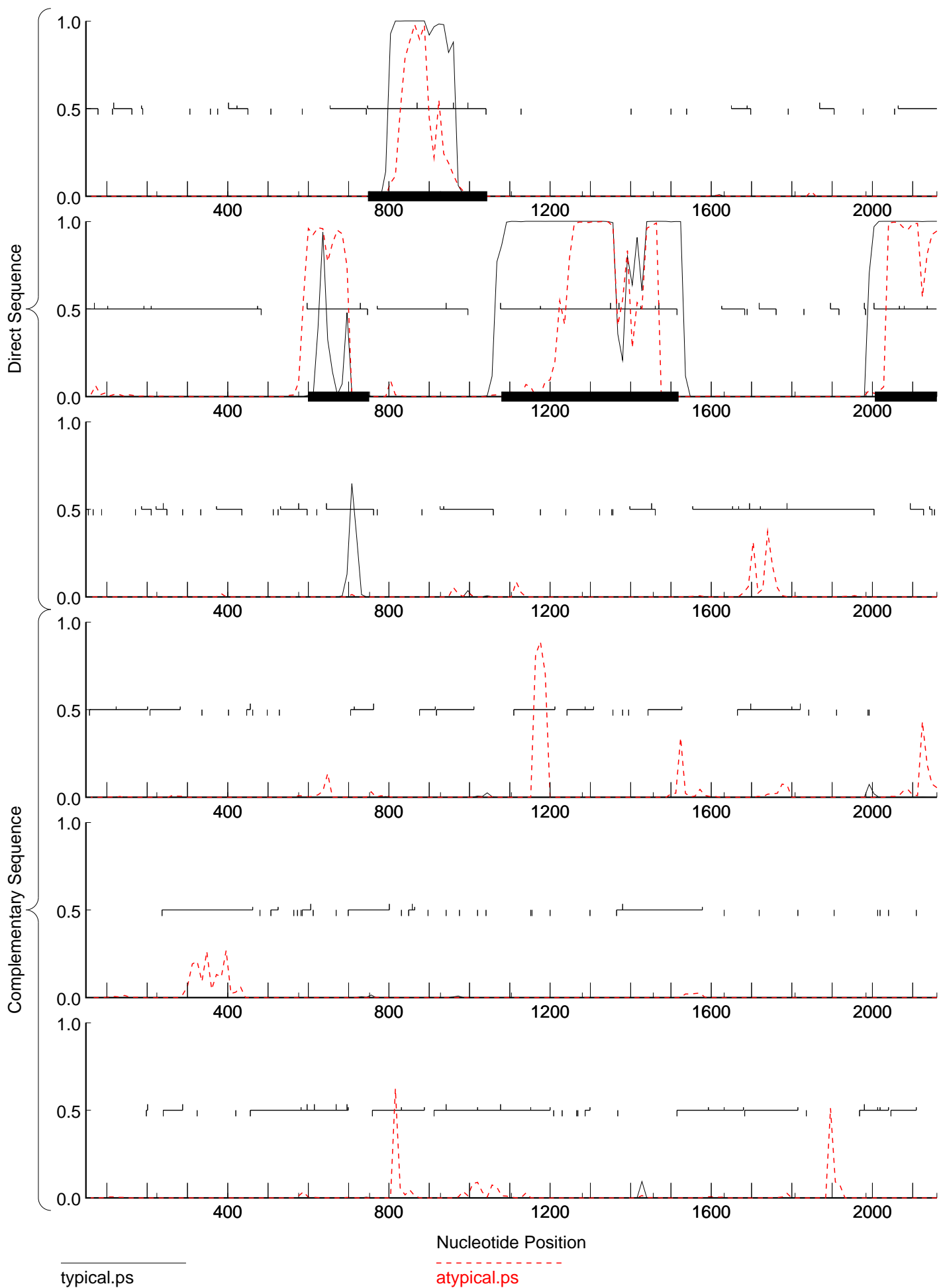
Matrix notes & comments

Training set derived by GeneMarkS, 4.30 October 2014
Sun Jun 24 05:13:00 2018

MATRIX INFORMATION

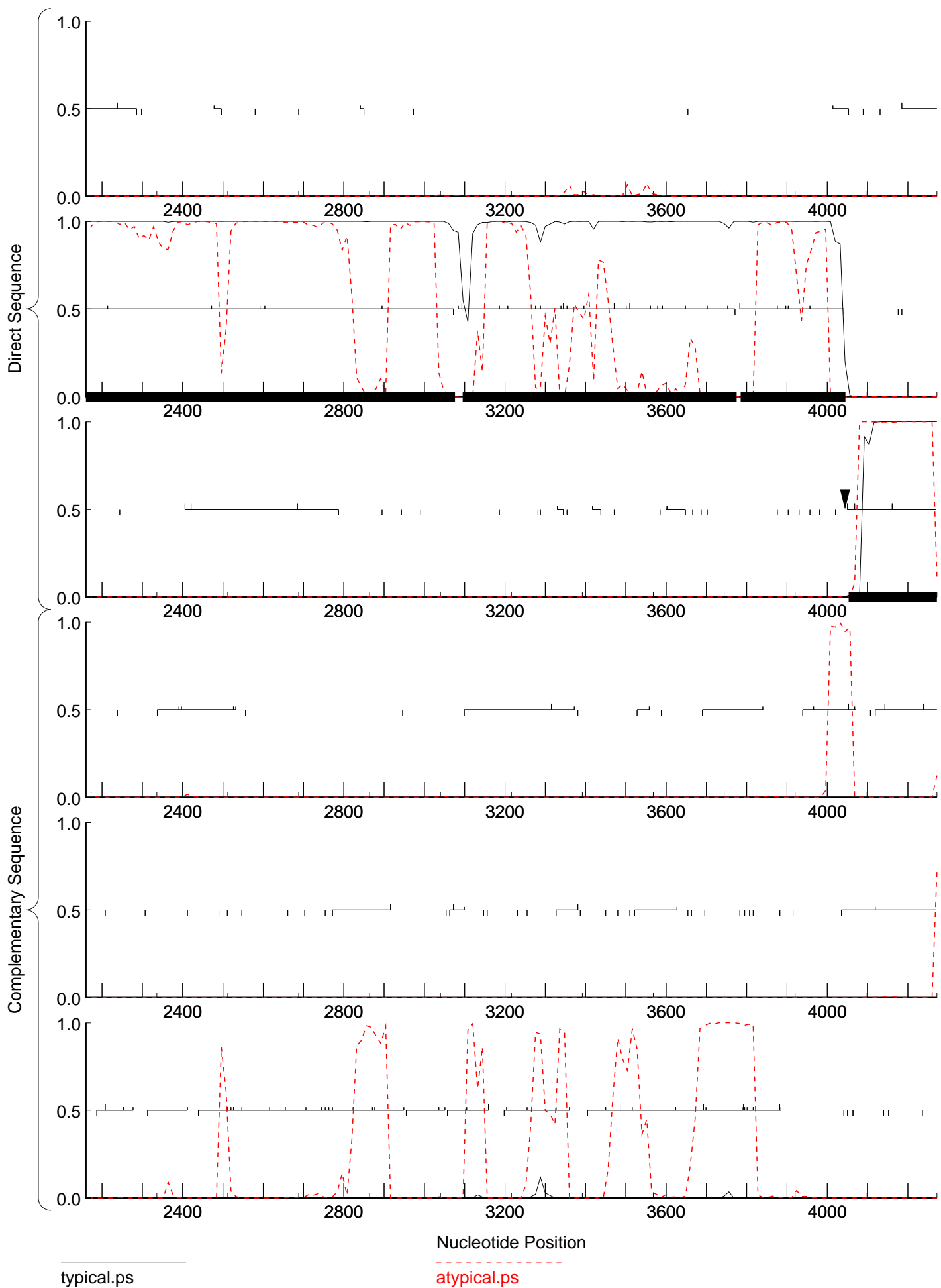
Matrix : -
Author : -
Order : 4

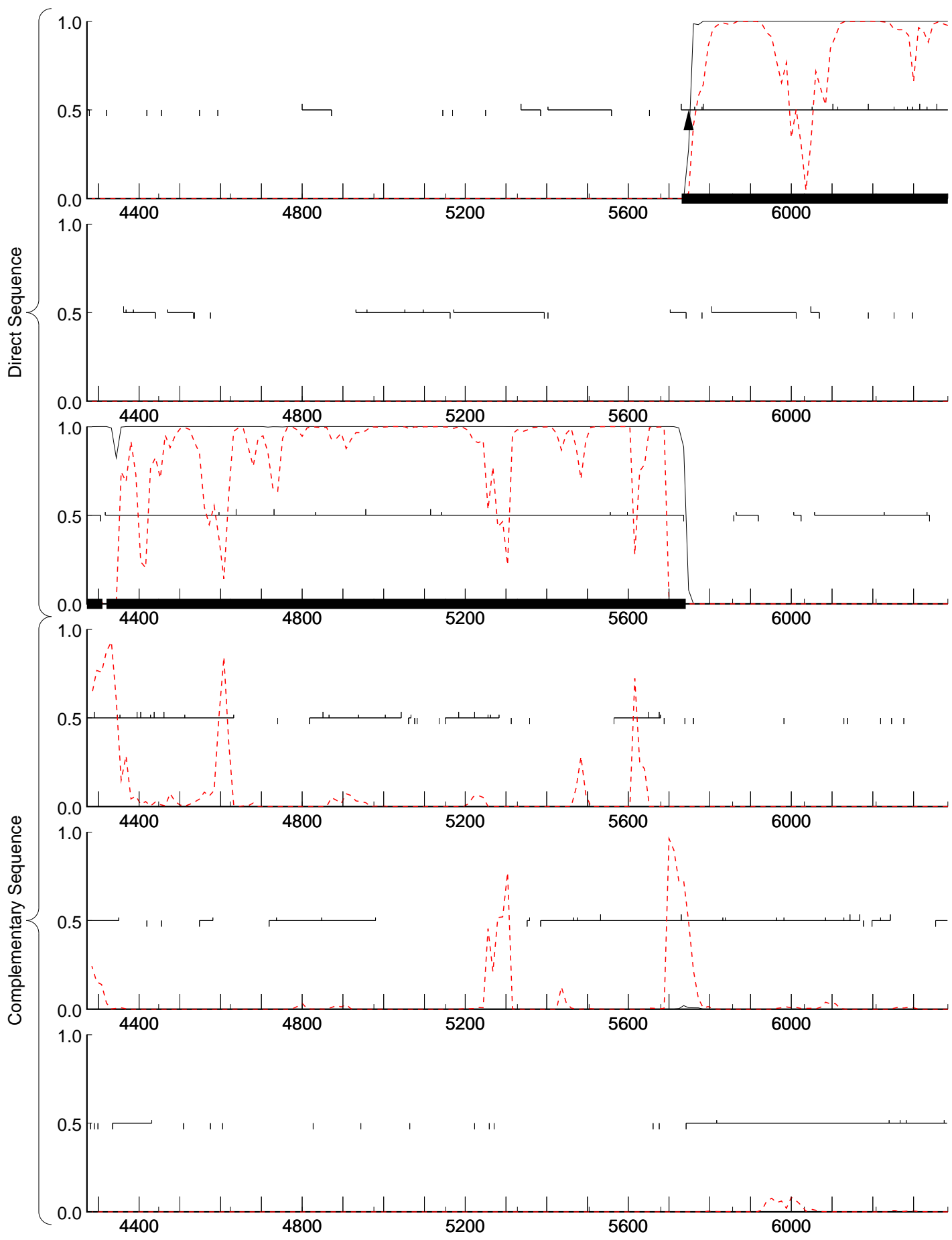
Send questions / comments to:
Dr. M. Borodovsky
Georgia Institute of Technology
School of Biology
Atlanta, GA 30332-0230



typical.ps

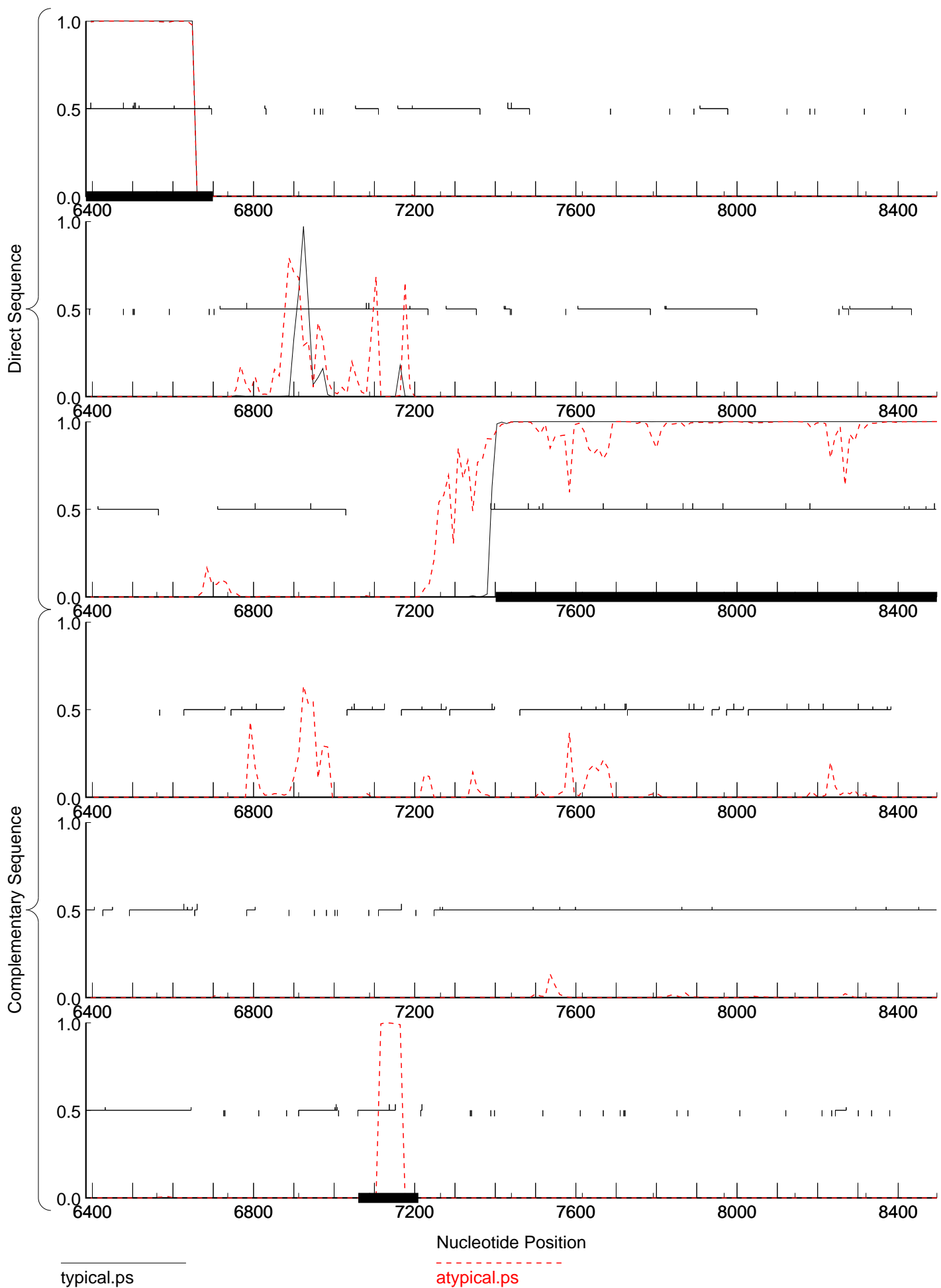
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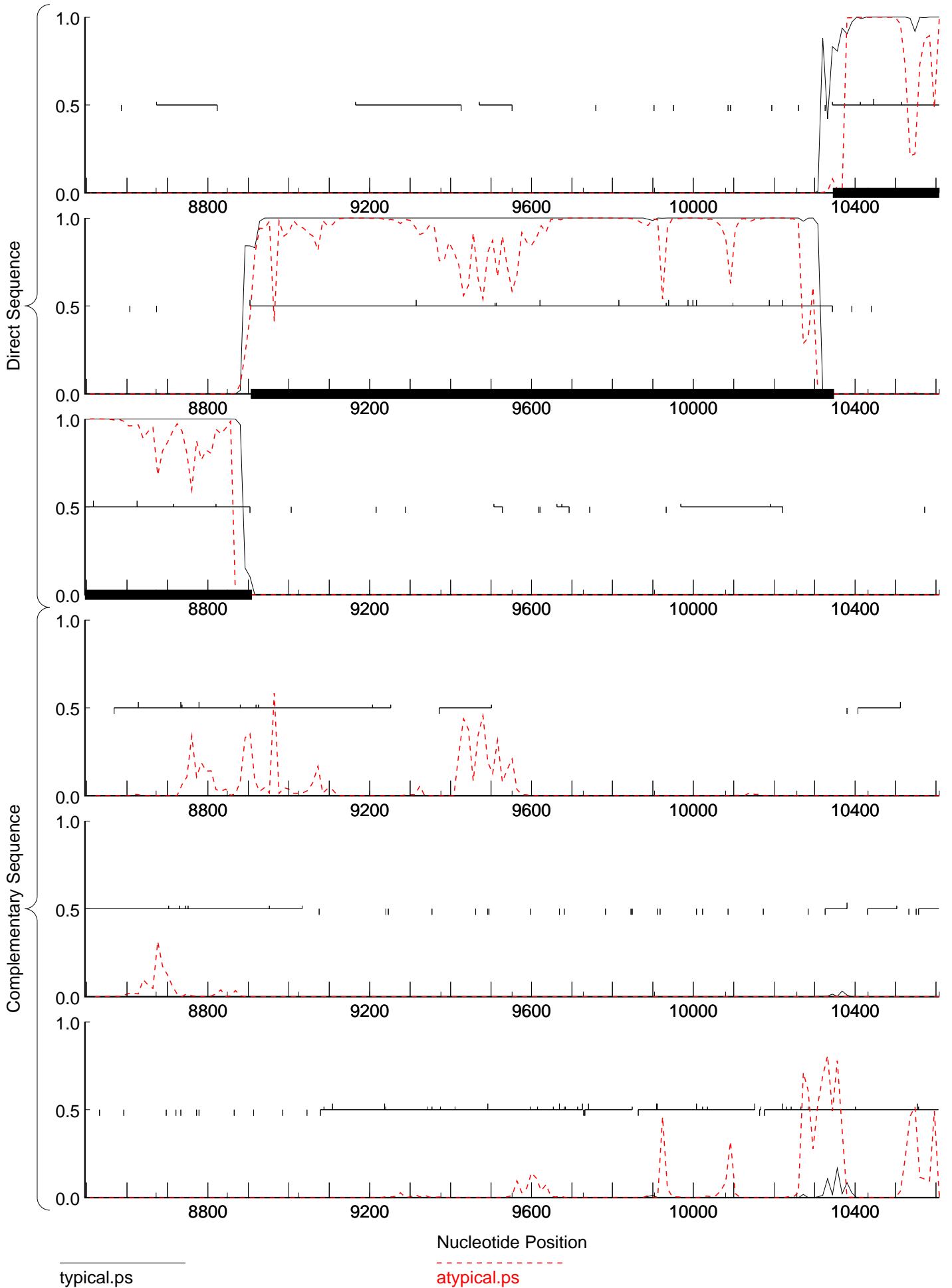


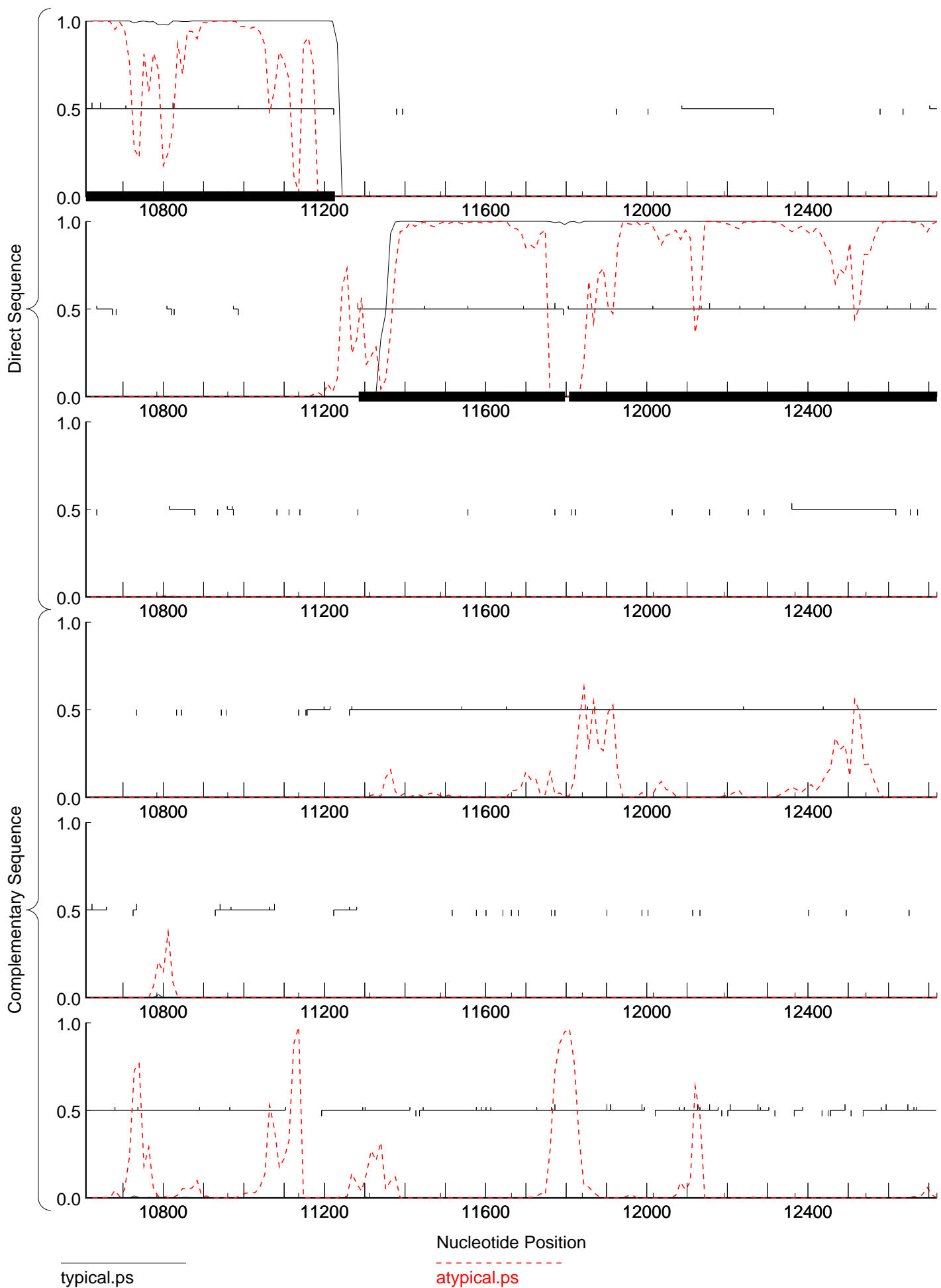


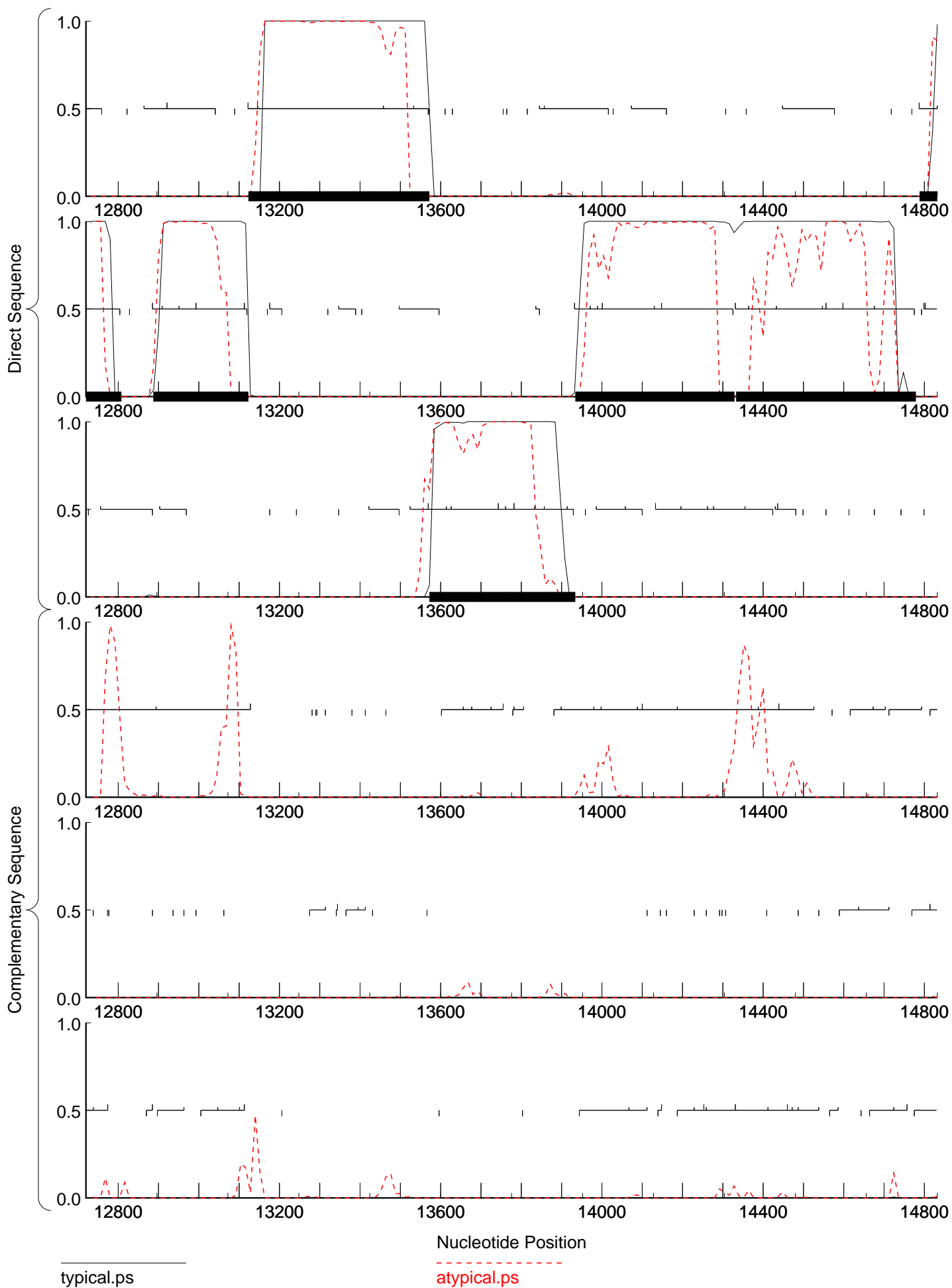
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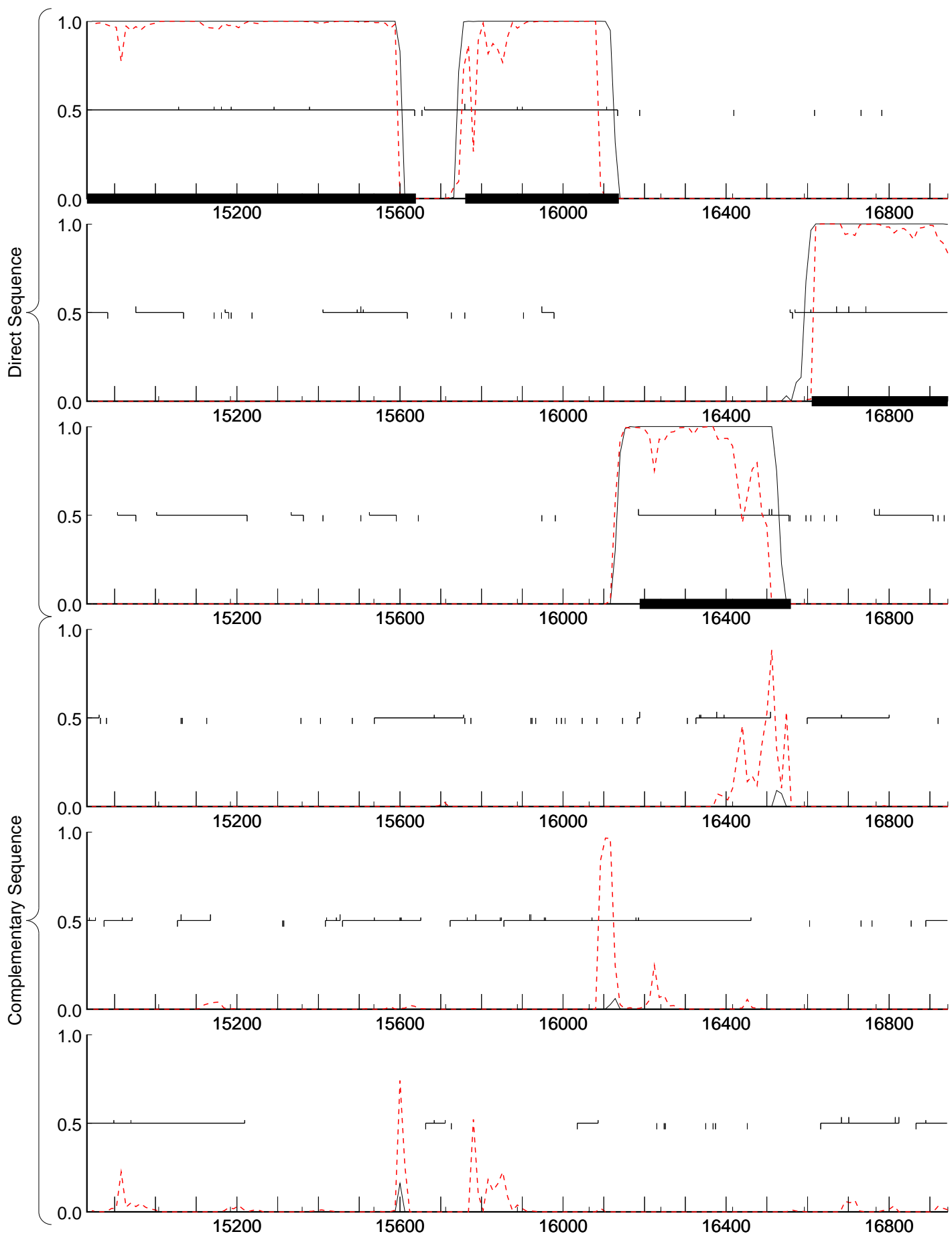
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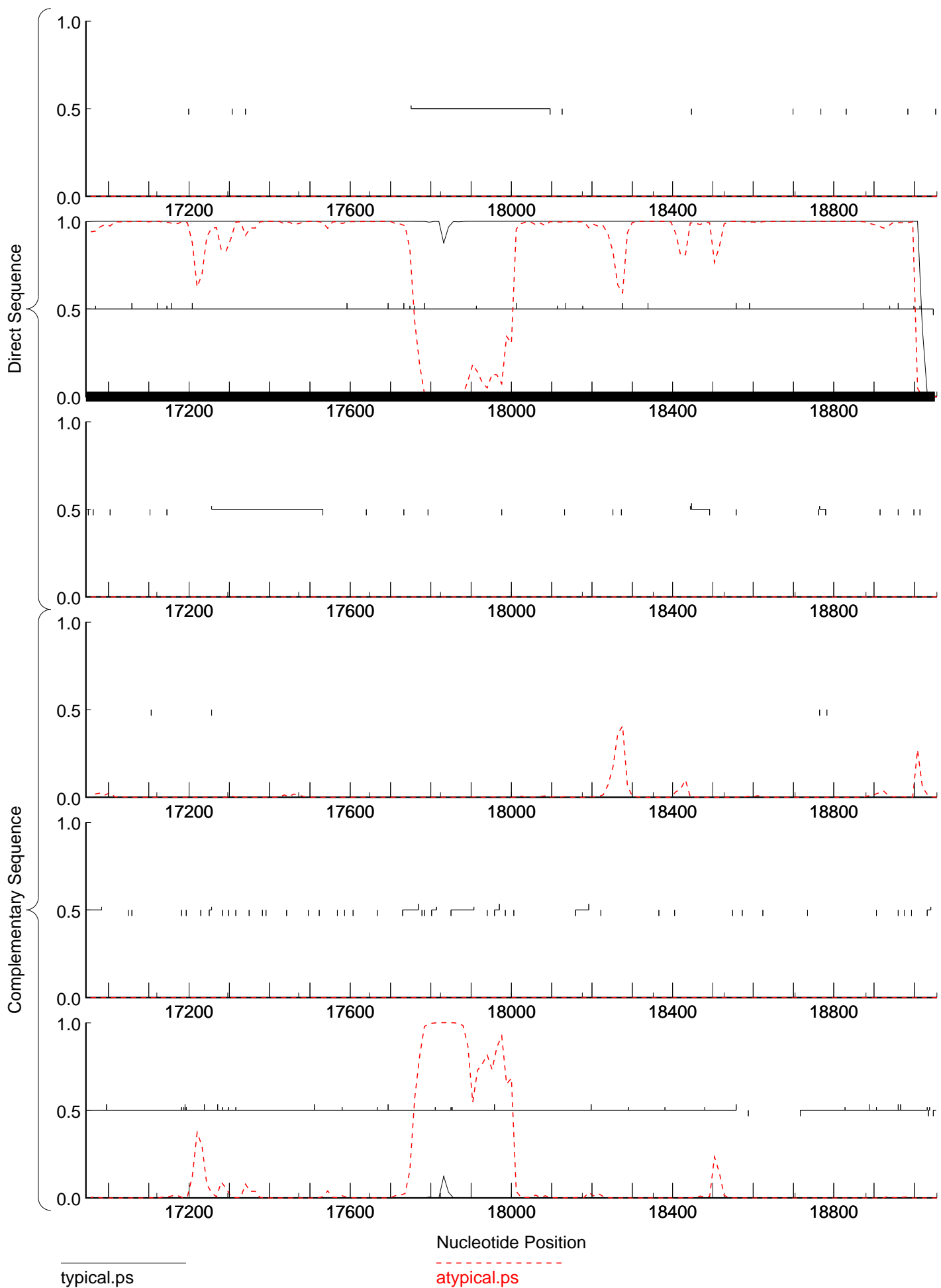


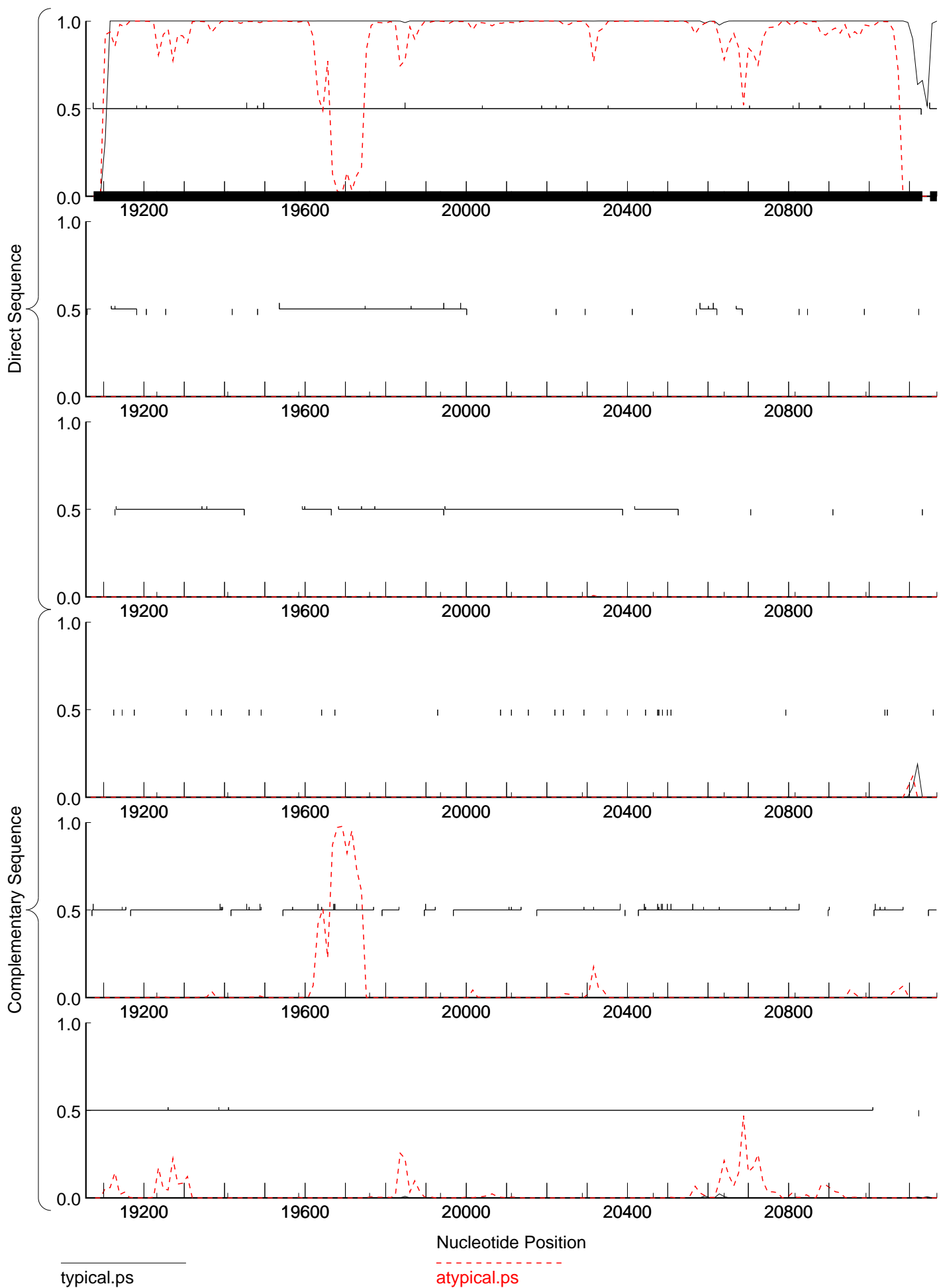




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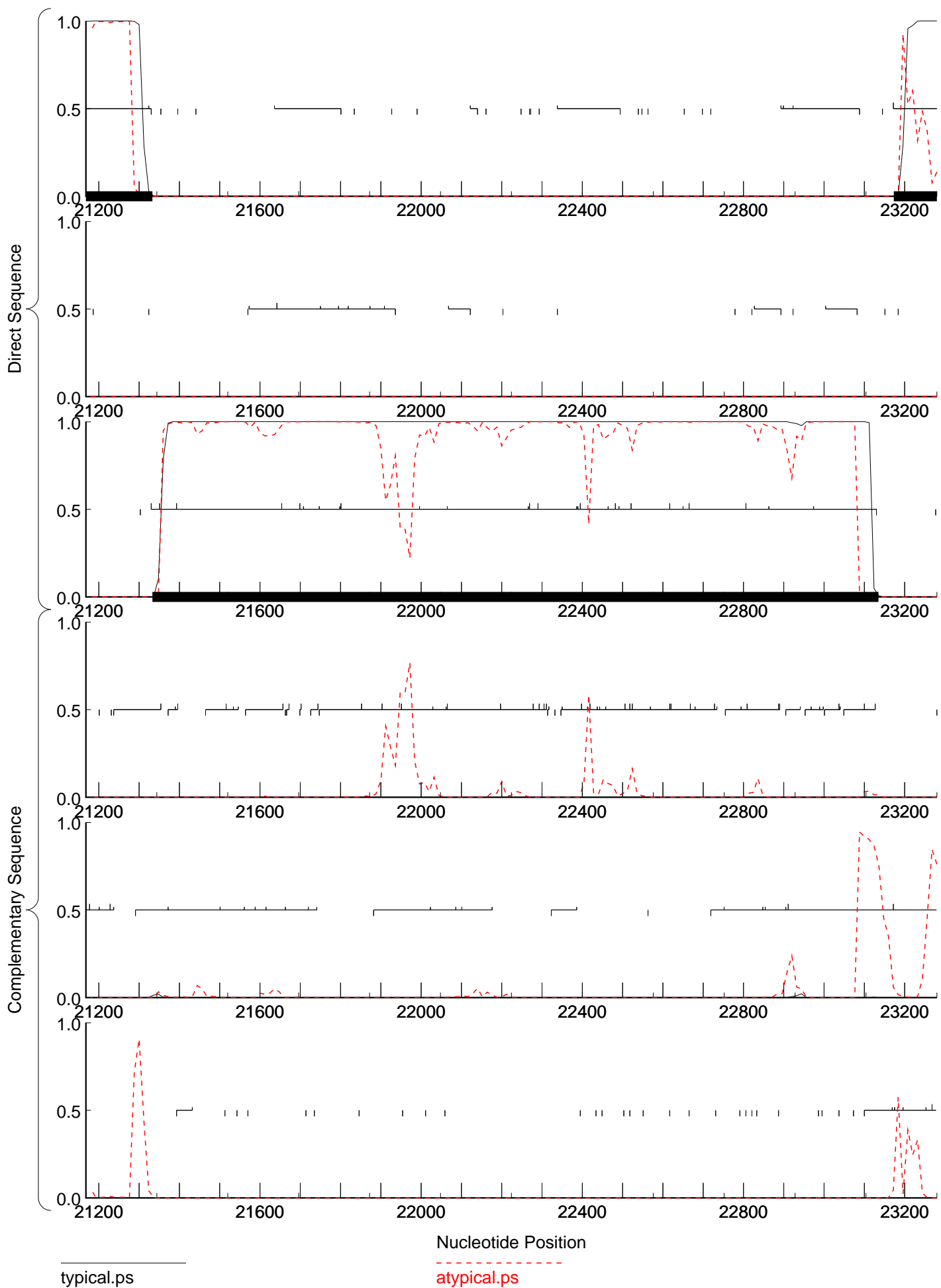
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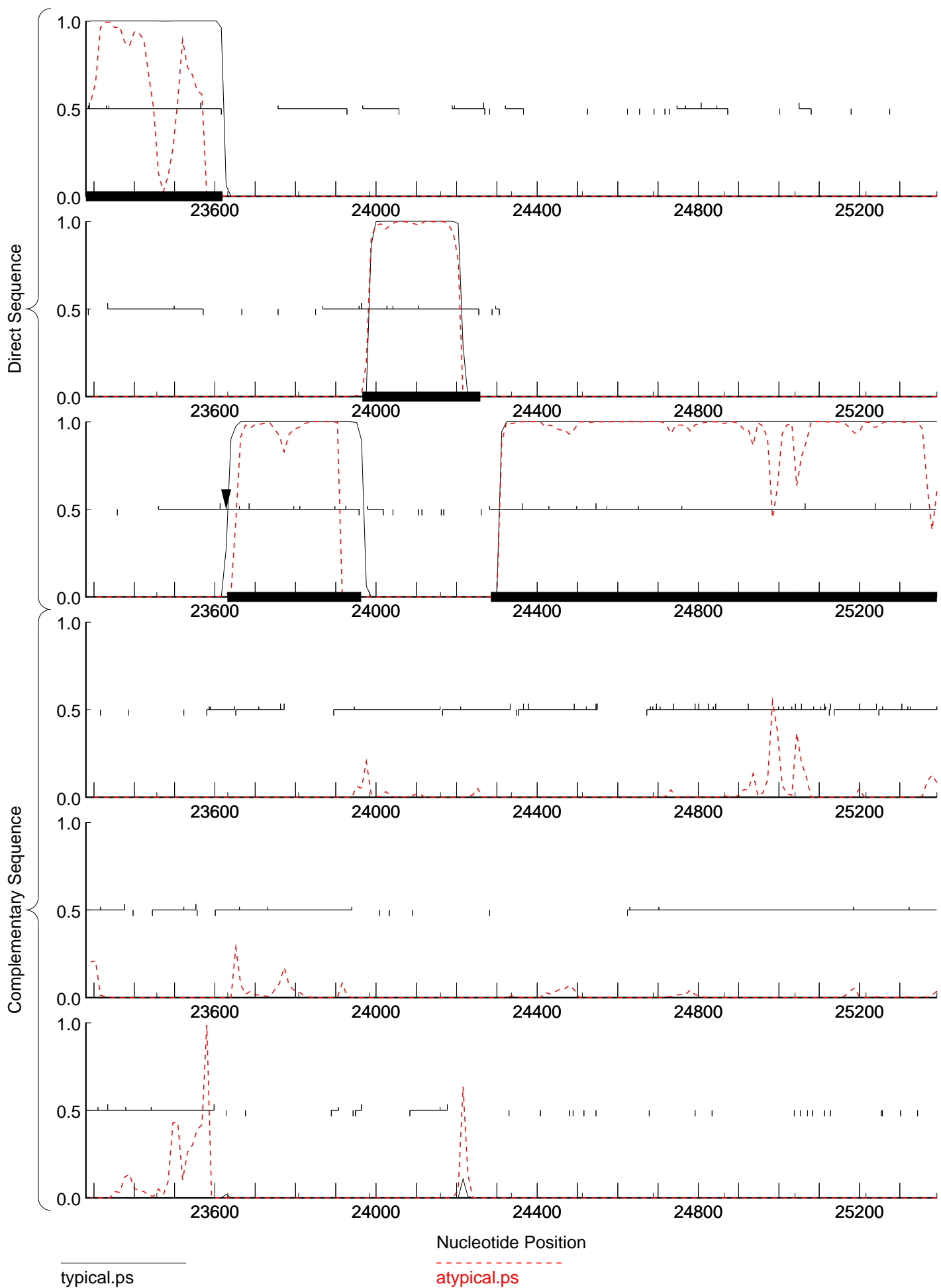
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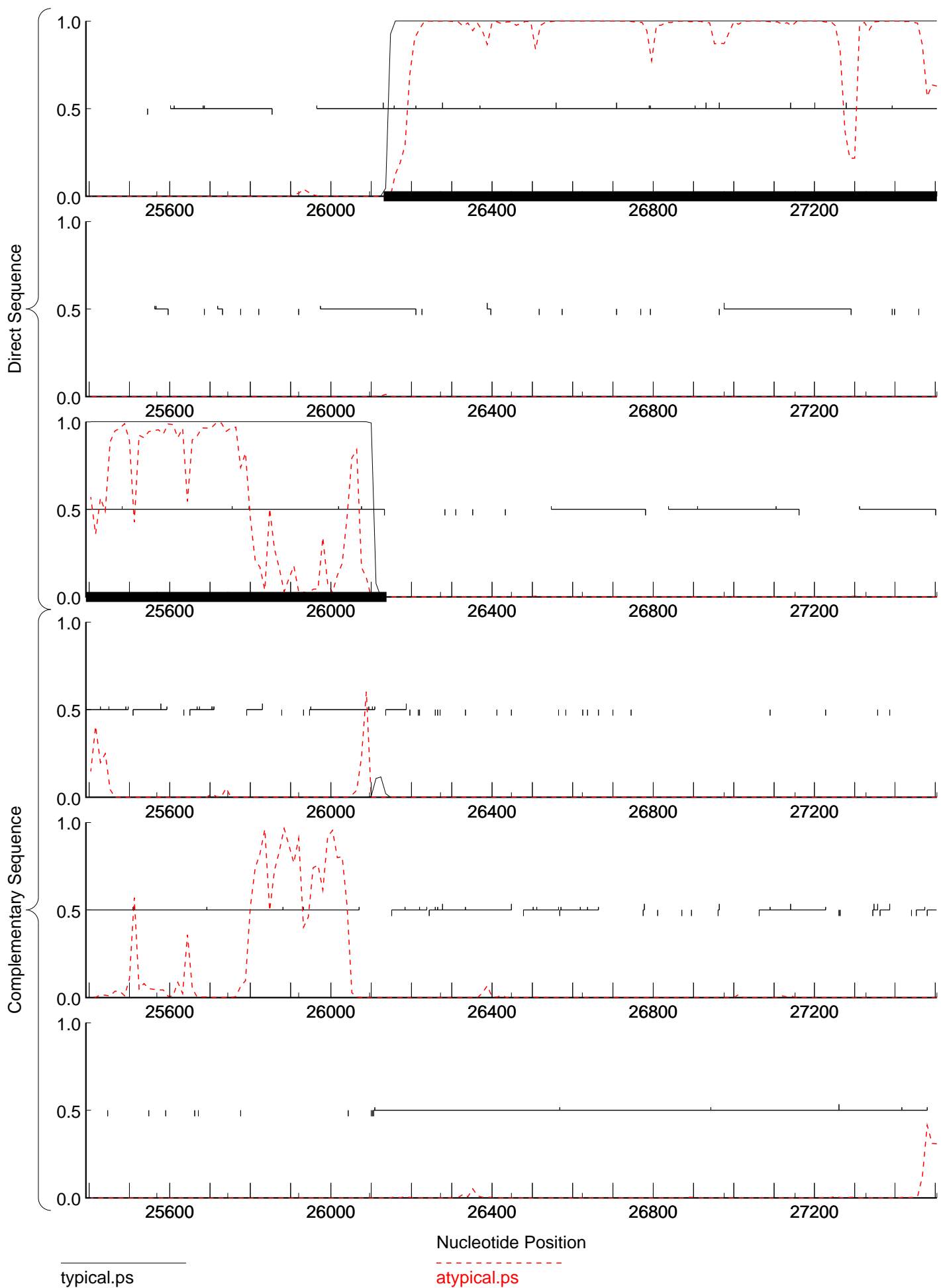
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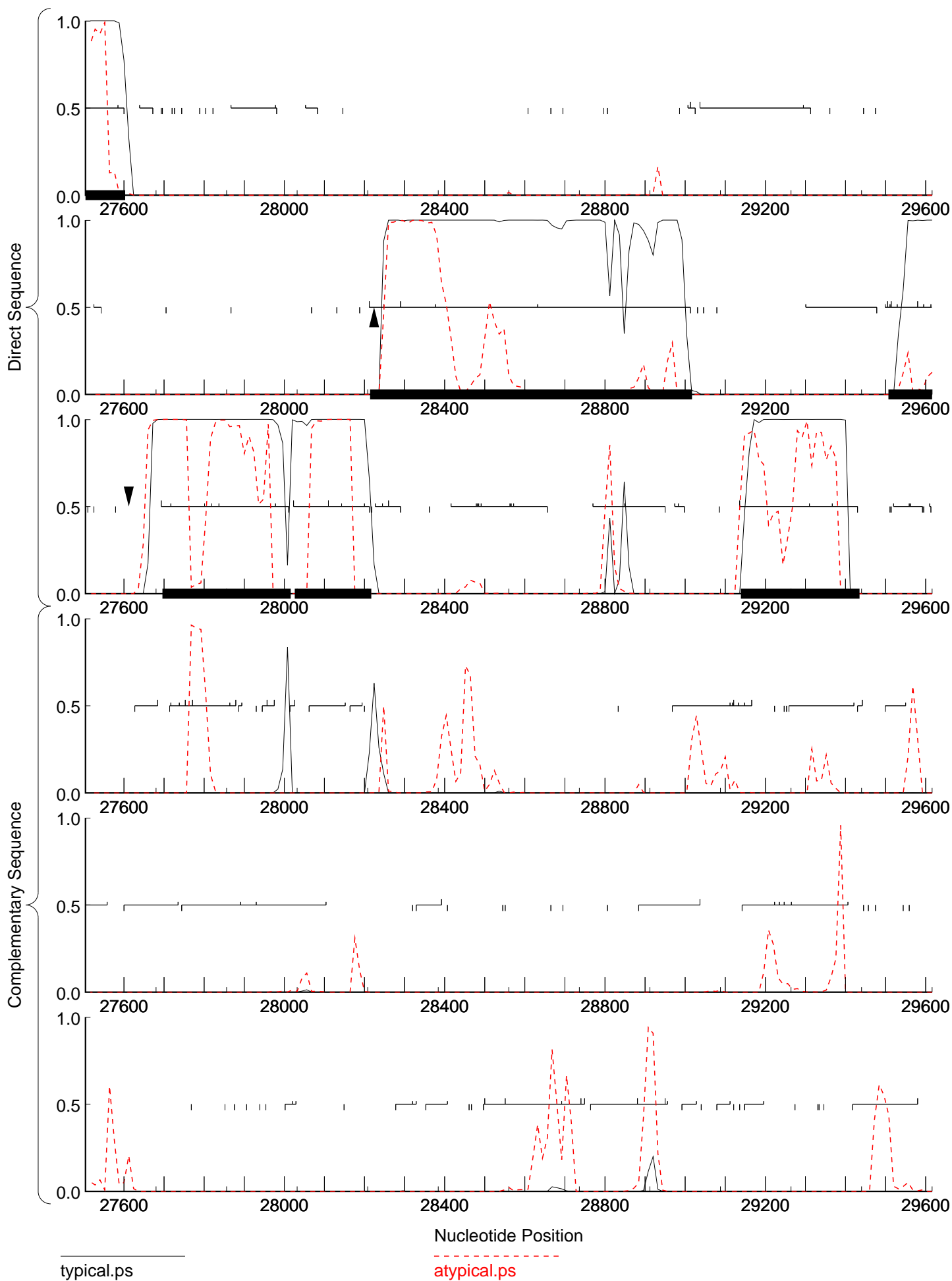


typical.ps

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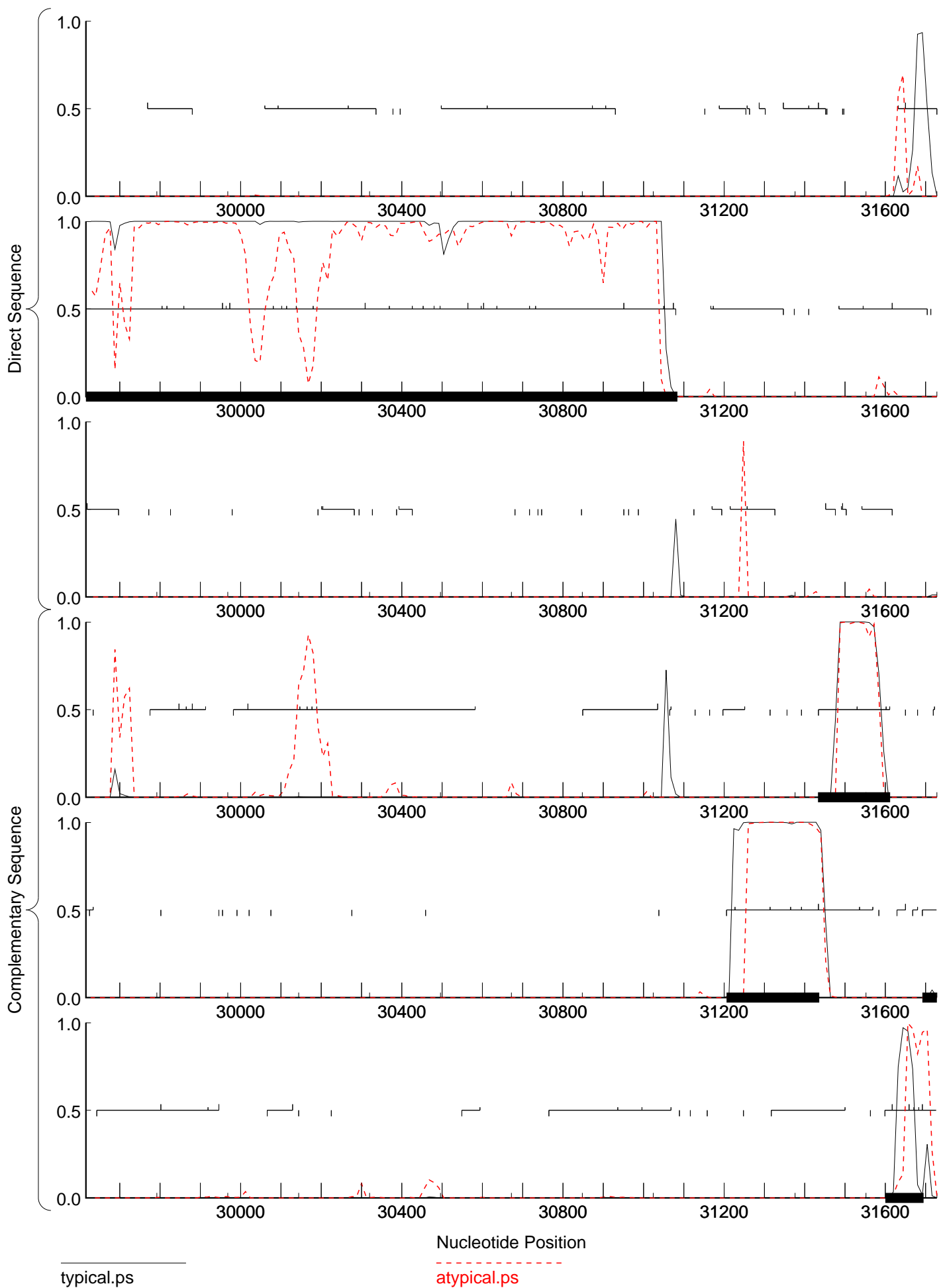


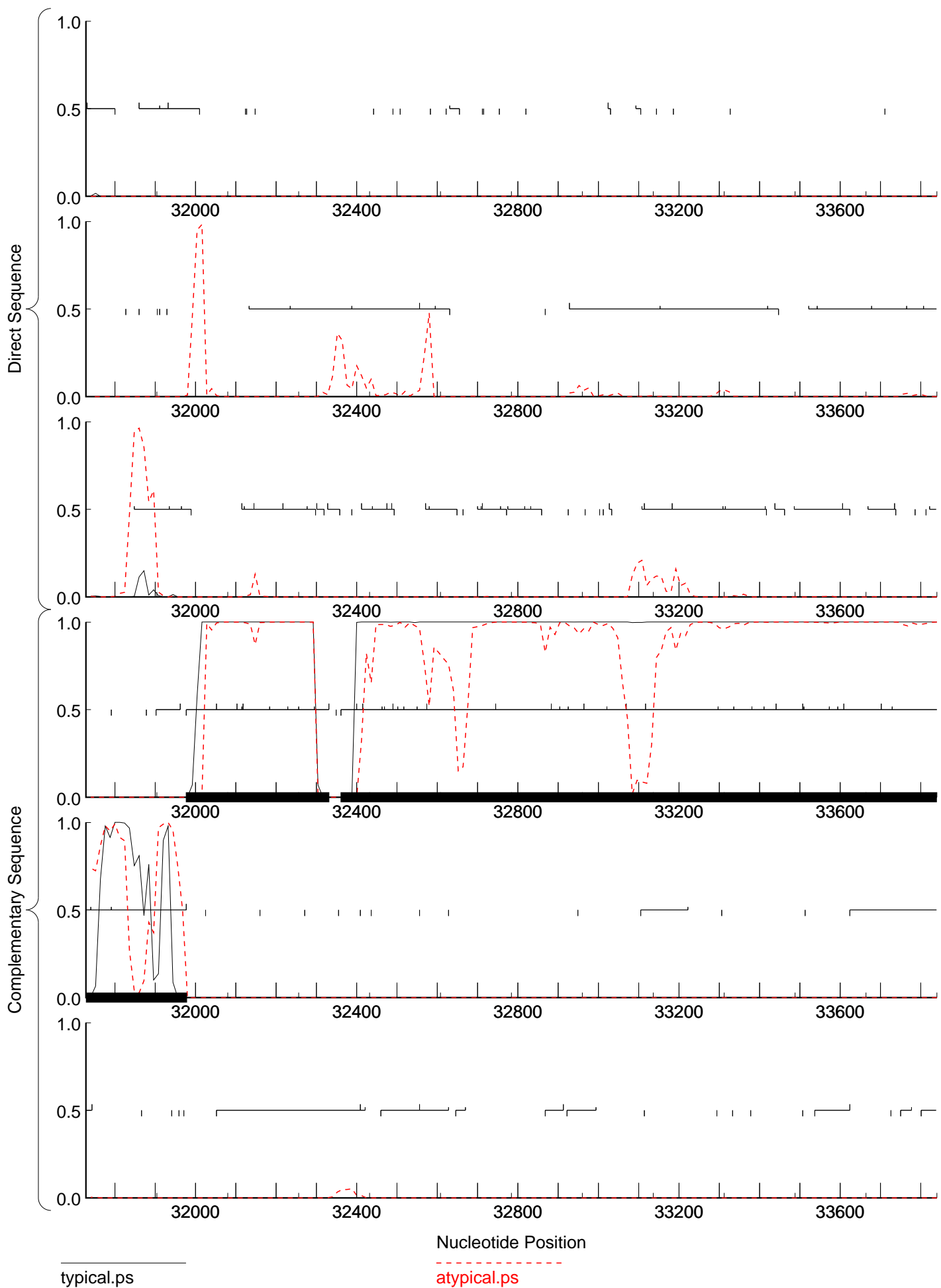


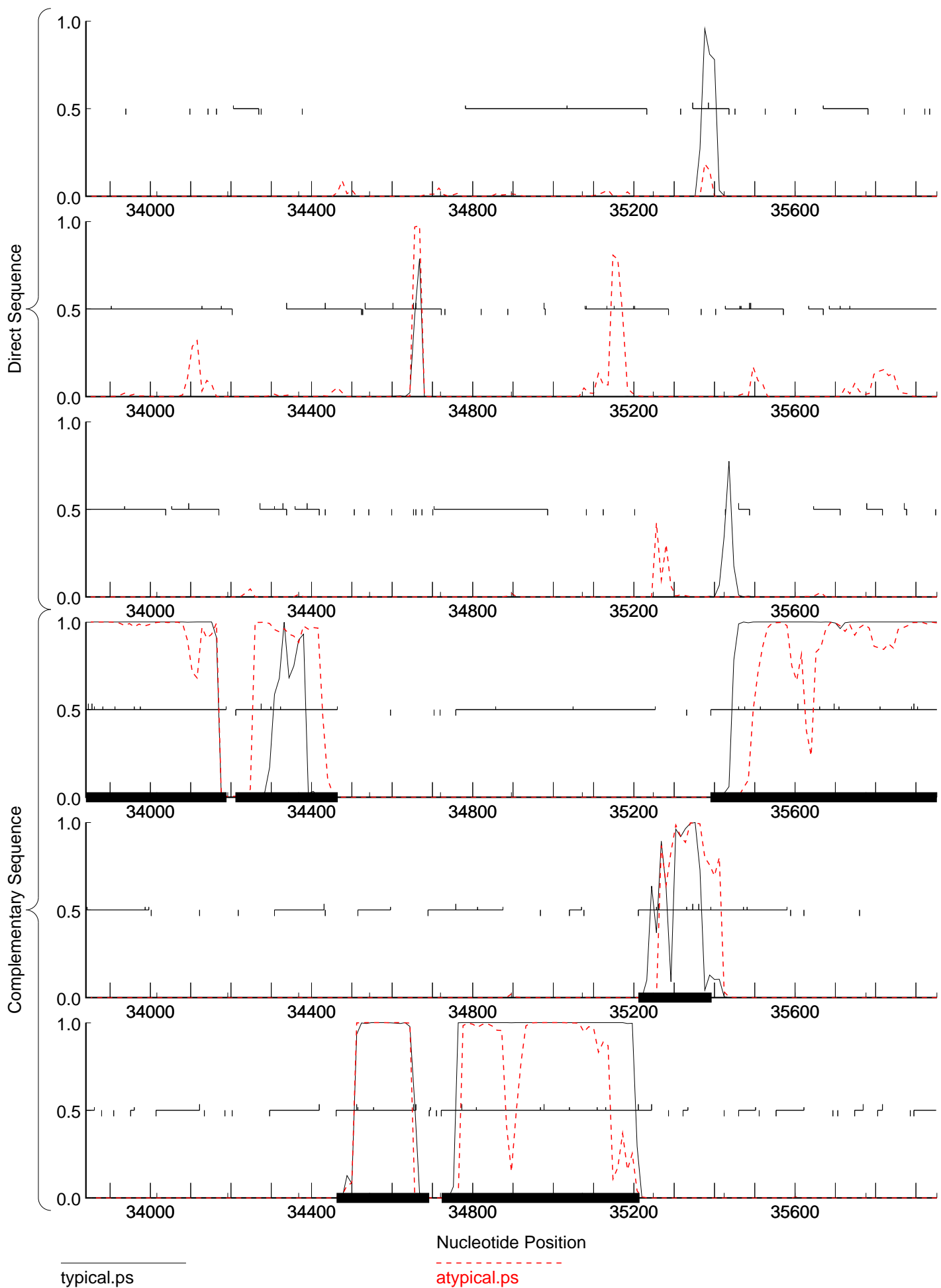


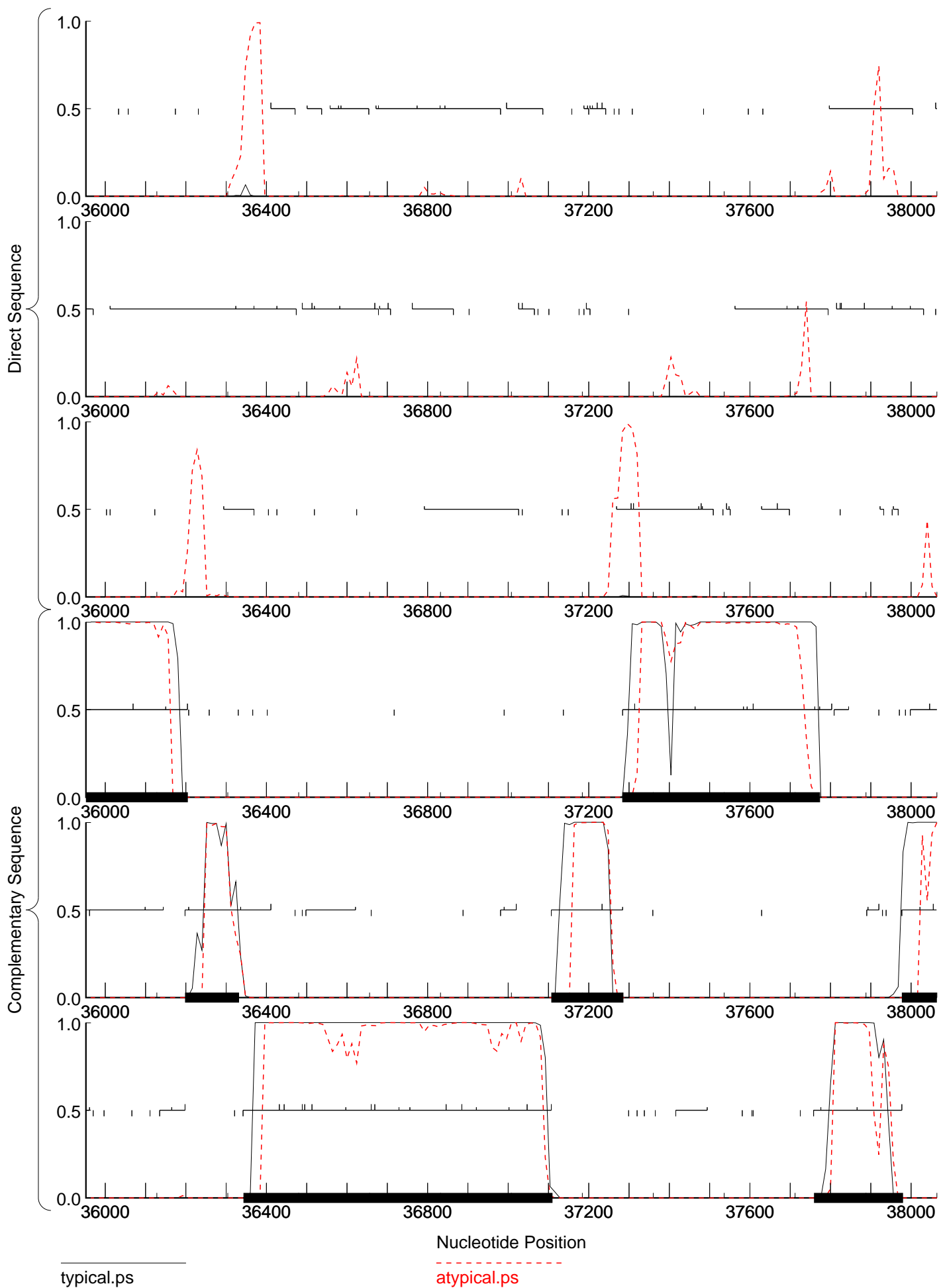
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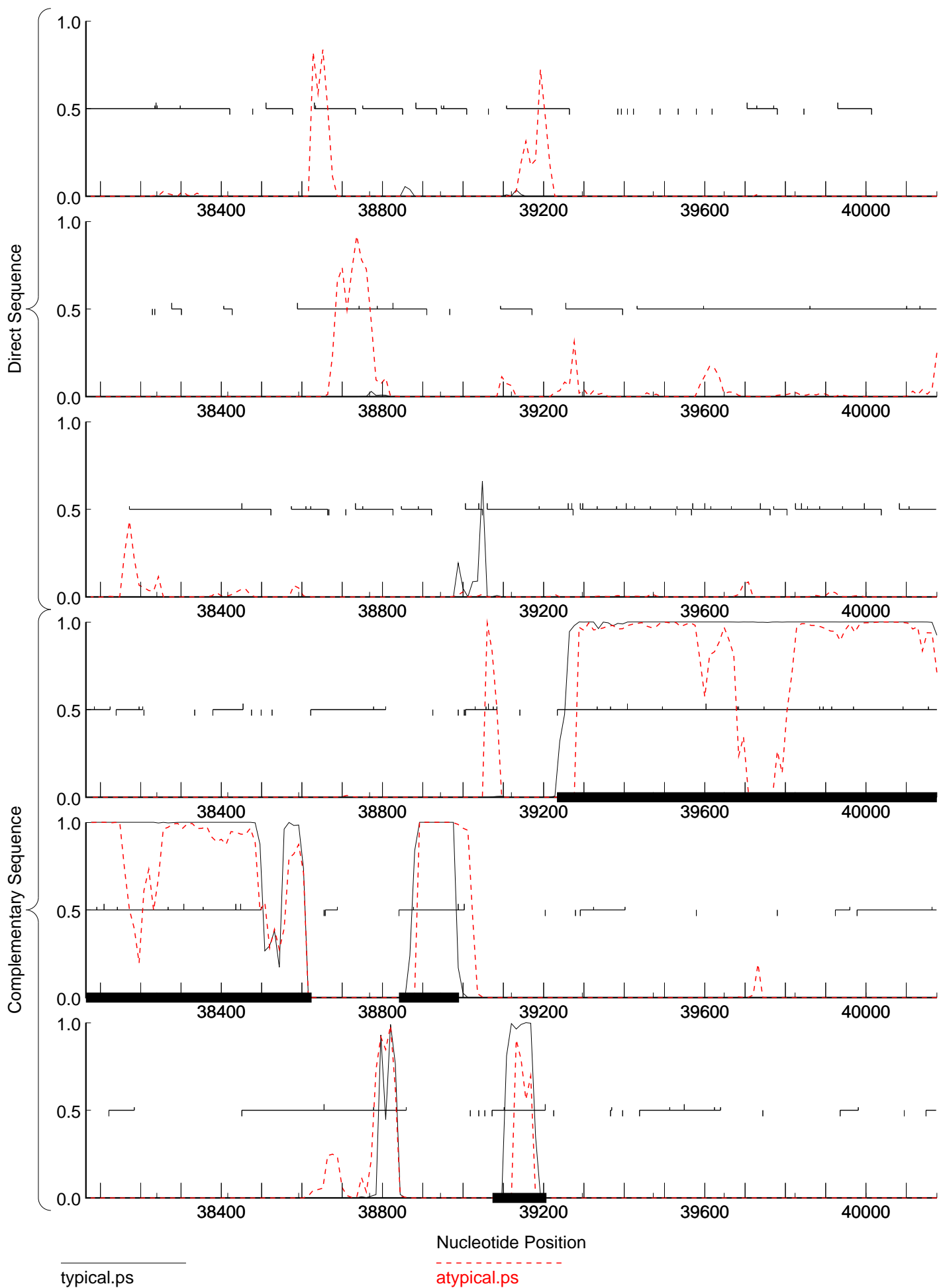


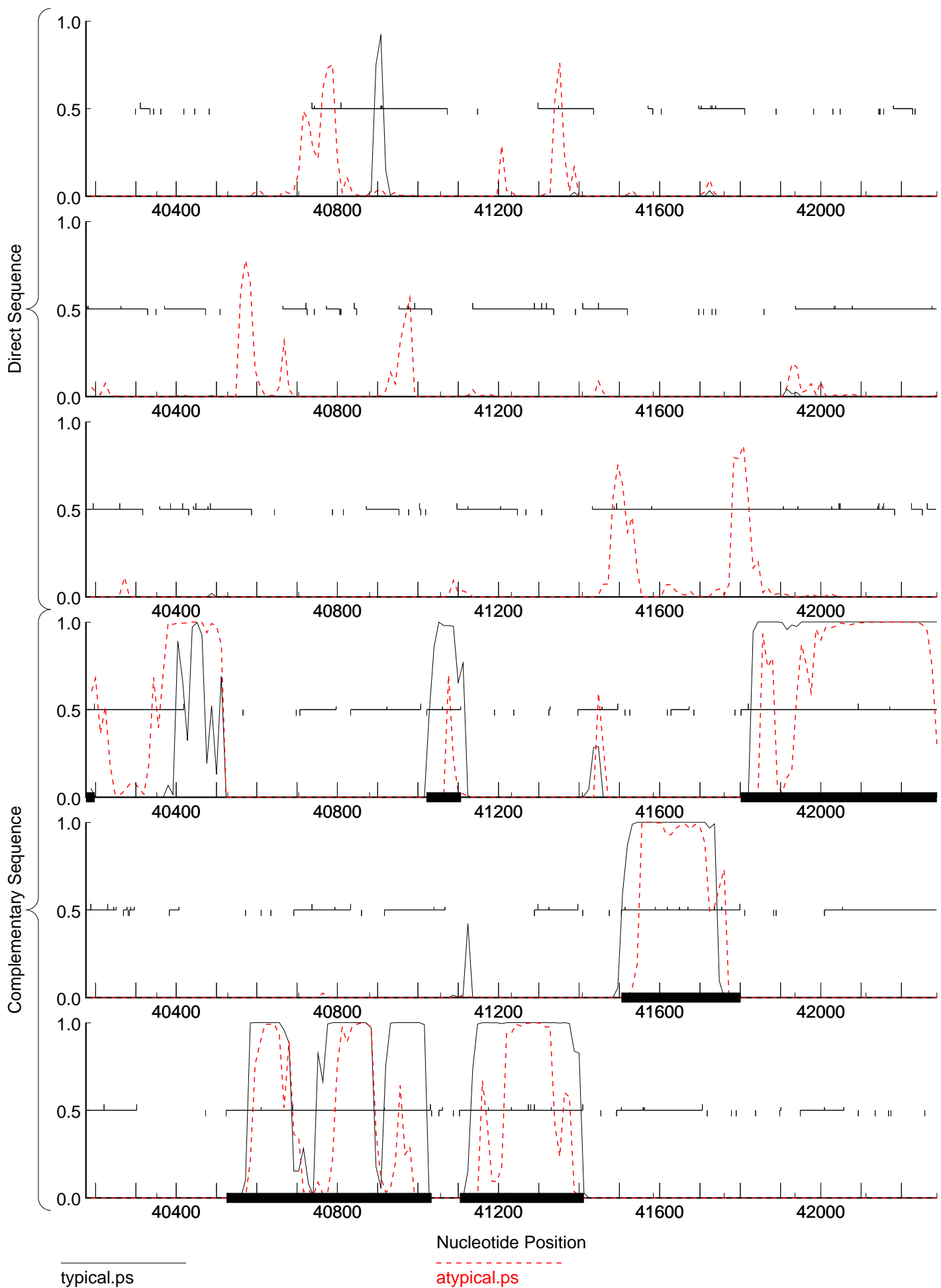


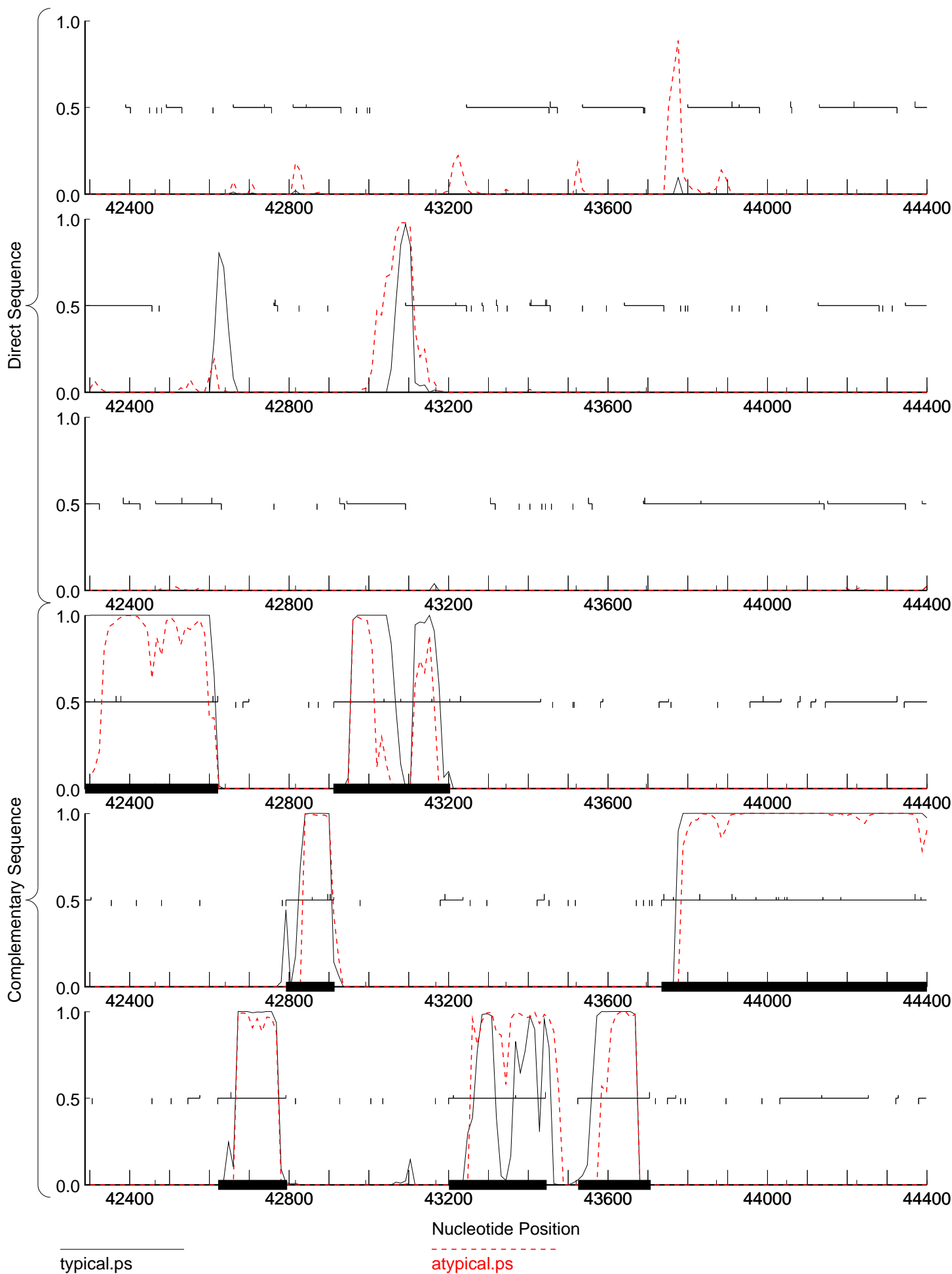


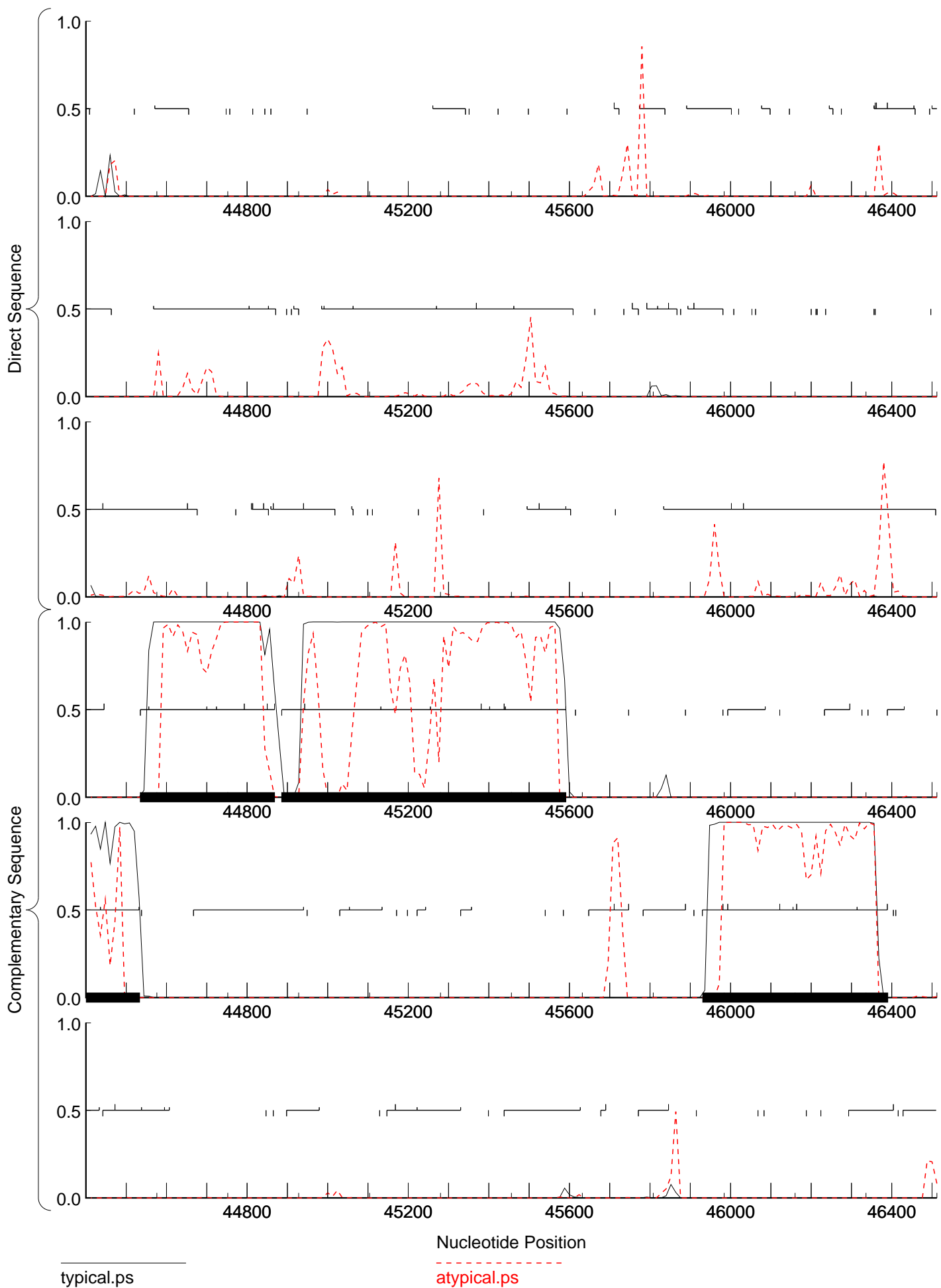
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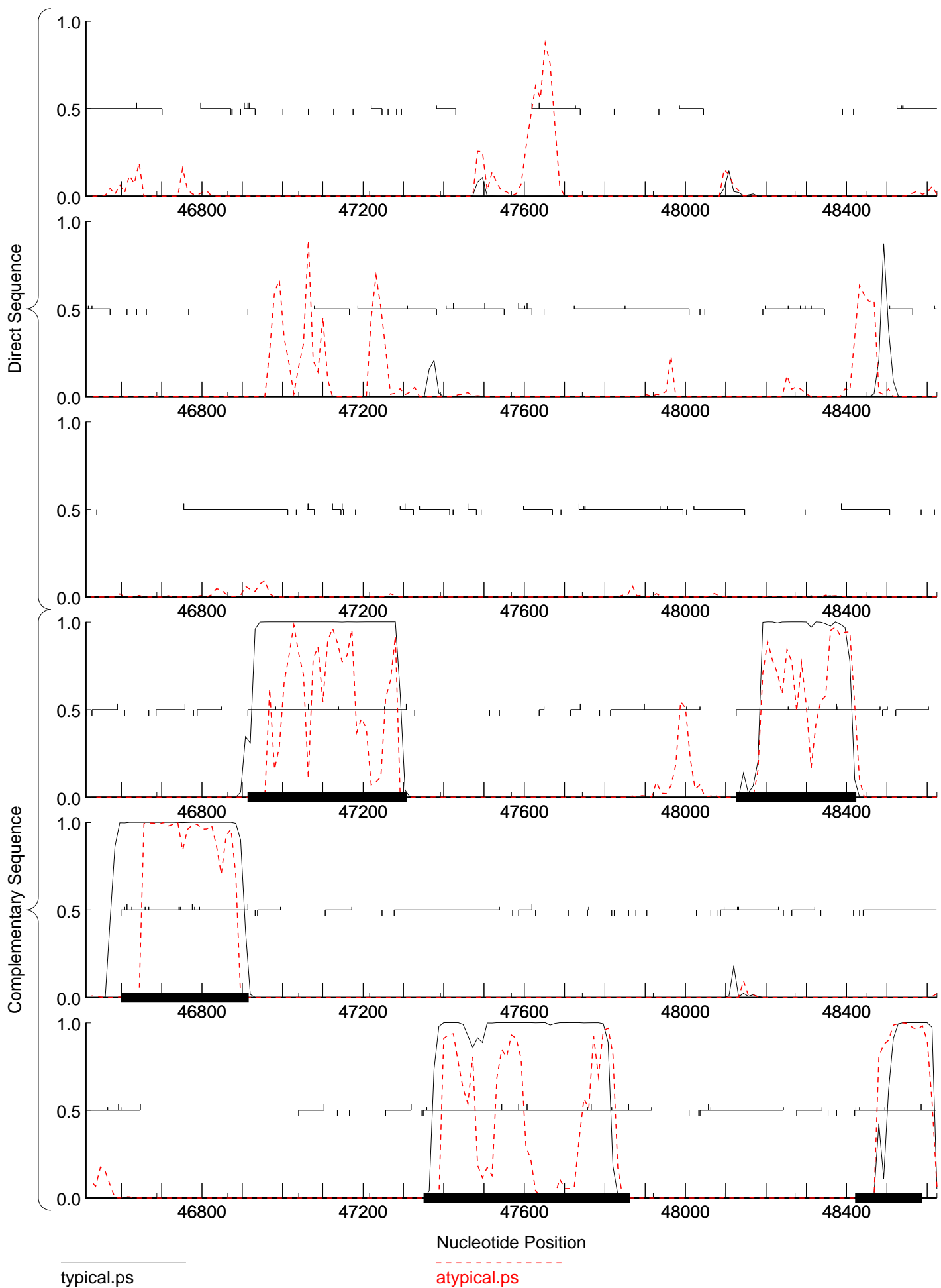
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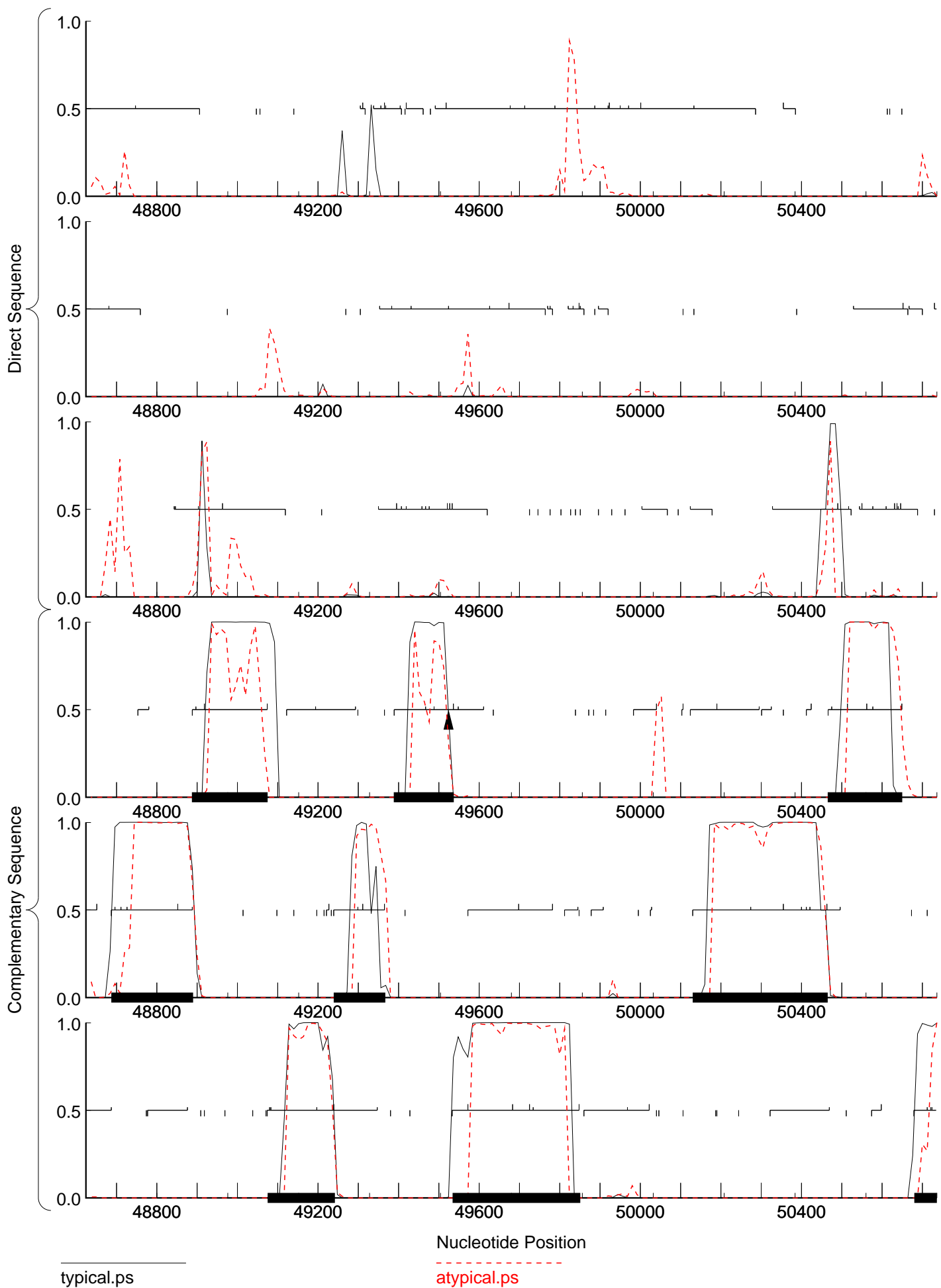












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