

GeneMark

Version 2.5p (09.08.06)

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PROGRAM INFORMATION

Sequence : Streptomyces phage Warpy complete sequence, 132996 bp including 11458 bp terminal repeat, C
Analysis Date : 11/19/17 at 19:03:47
Pages : 64
Sequence Length : 132996 bp
GC Content : 49.56%

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,

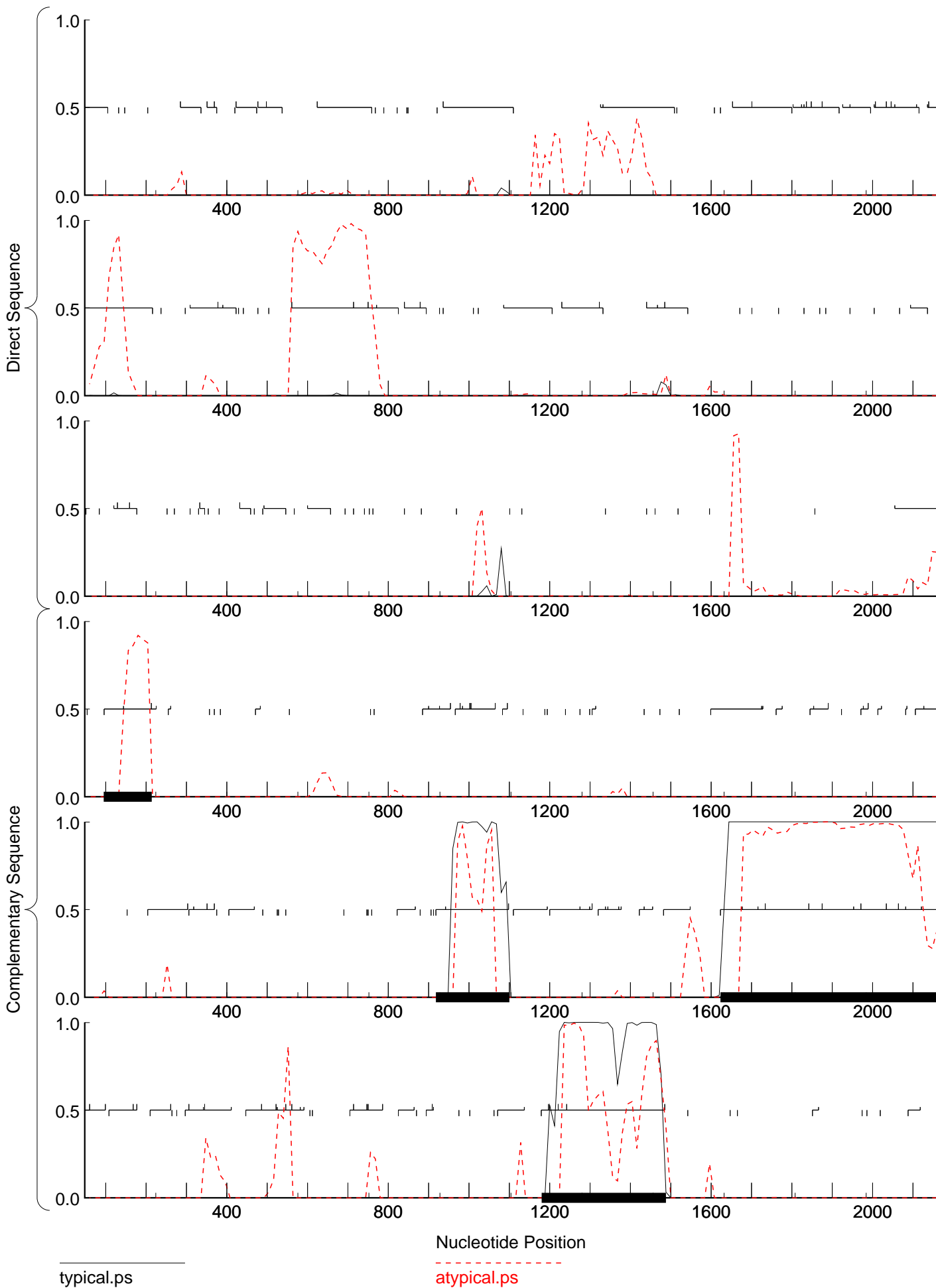
Matrix notes & comments

Training set derived by GeneMarkS, 4.30 October 2014
Sun Nov 19 19:03:47 2017

MATRIX INFORMATION

Matrix : -
Author : -
Order : 4

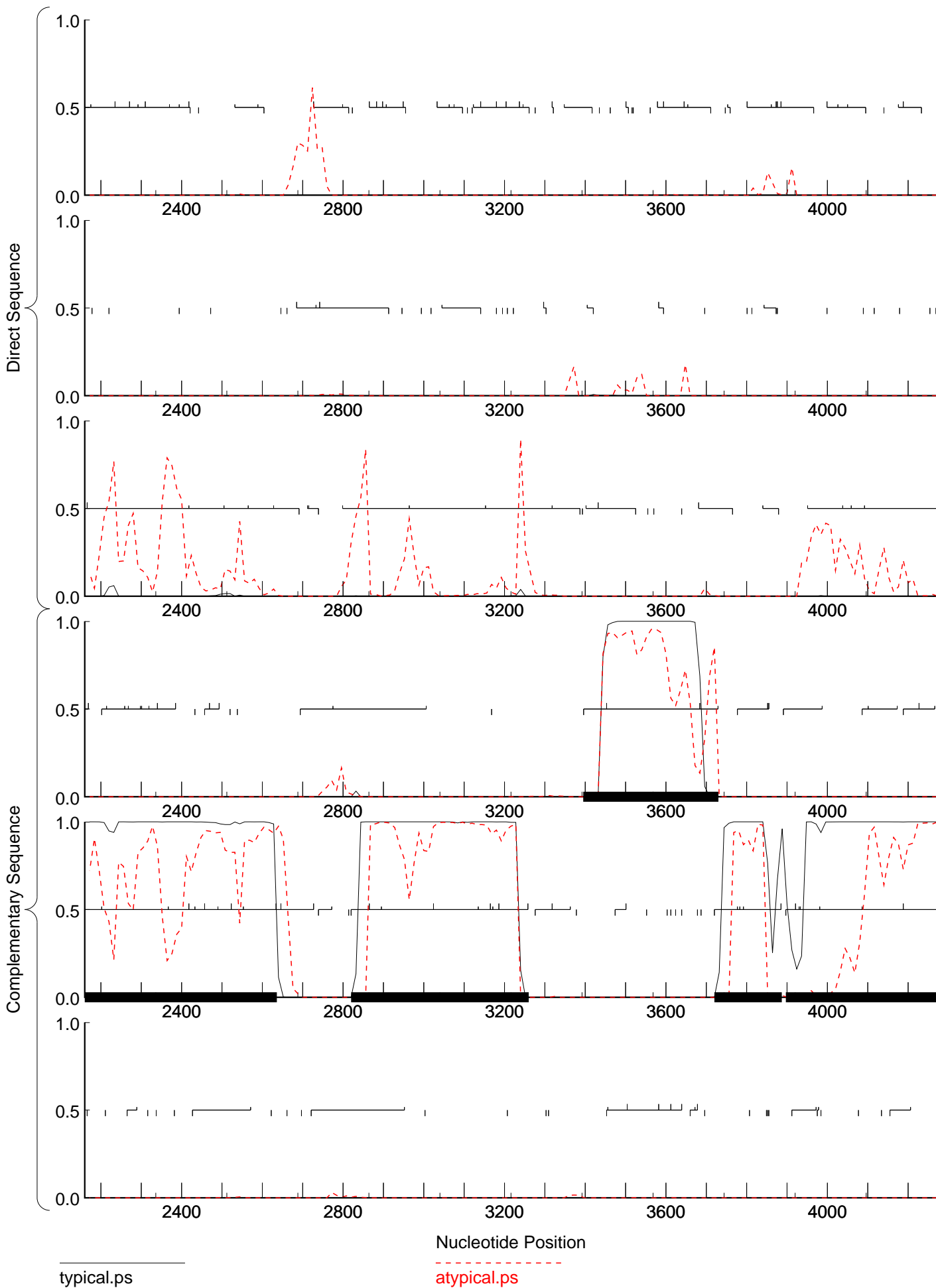
Send questions / comments to:
Dr. M. Borodovsky
Georgia Institute of Technology
School of Biology
Atlanta, GA 30332-0230



typical.ps

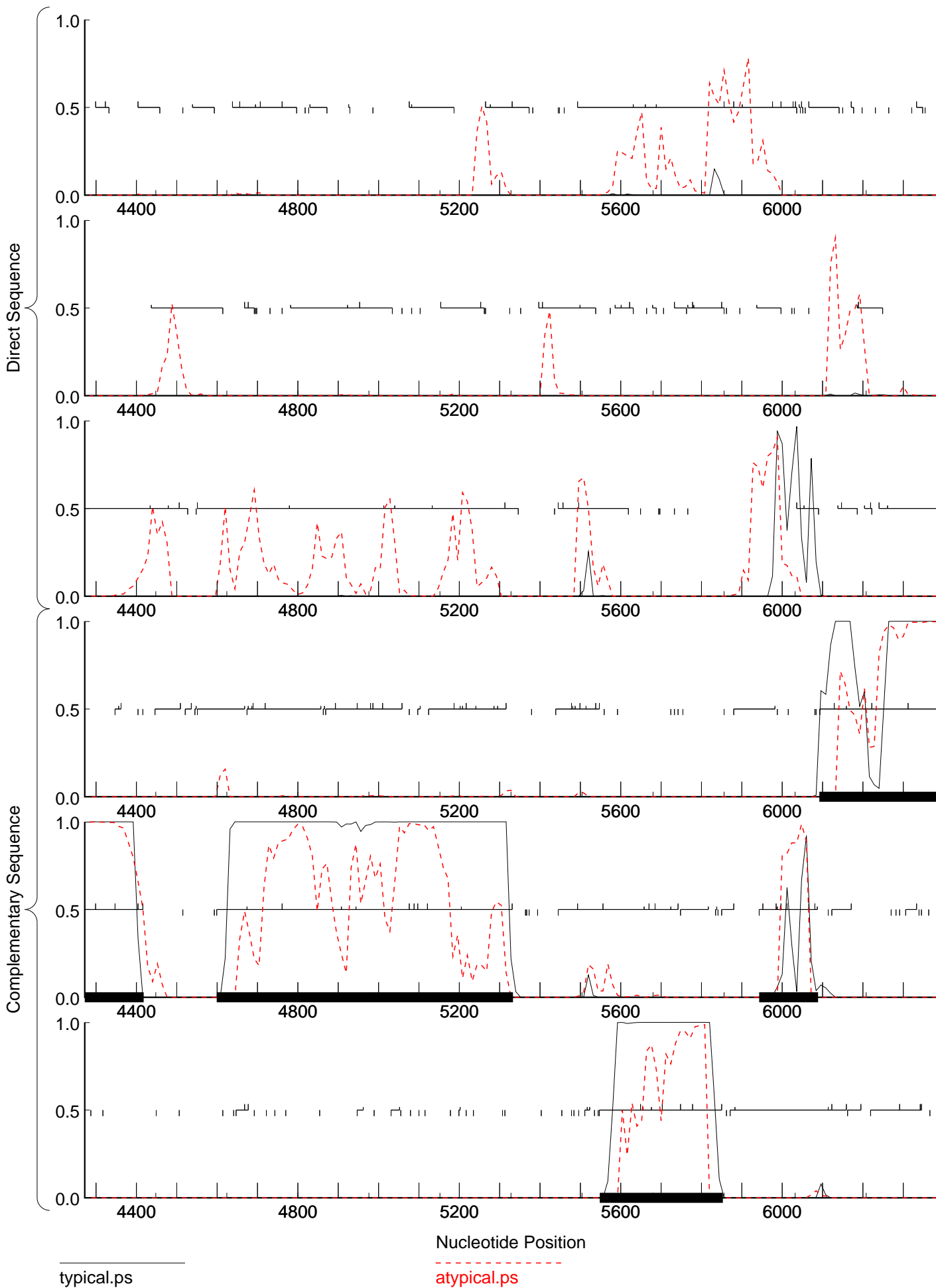
atypical.ps

hage Warty complete sequence, 132996 bp including 11458 bp terminal repeat, Cluster BE1, Order 4, Window 96, Step 12, 3/64
hage Warty complete sequence, 132996 bp including 11458 bp terminal repeat, Cluster BE1, Order 2, Window 96, Step 12, 3/64



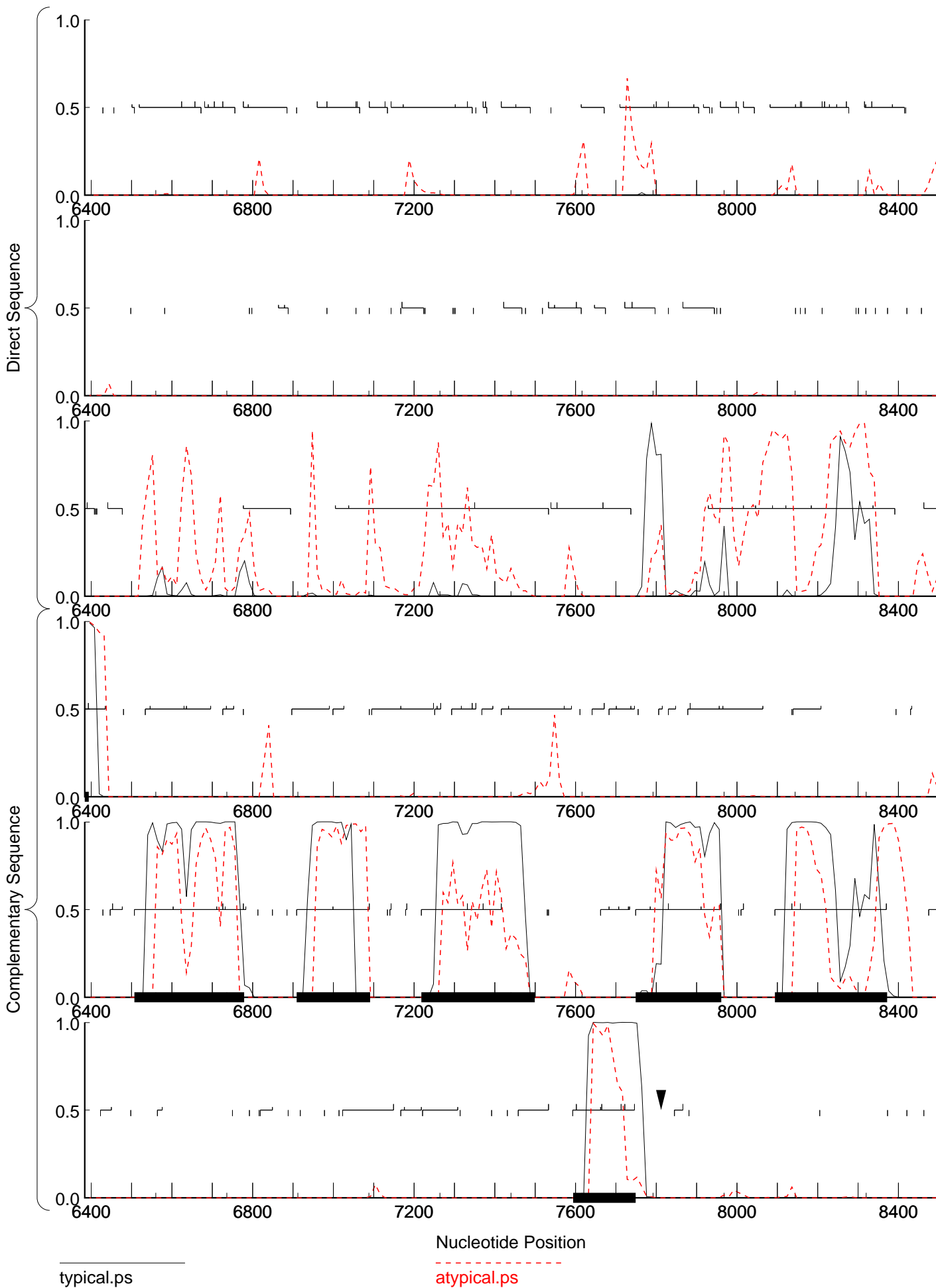
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atypical.ps



typical.ps

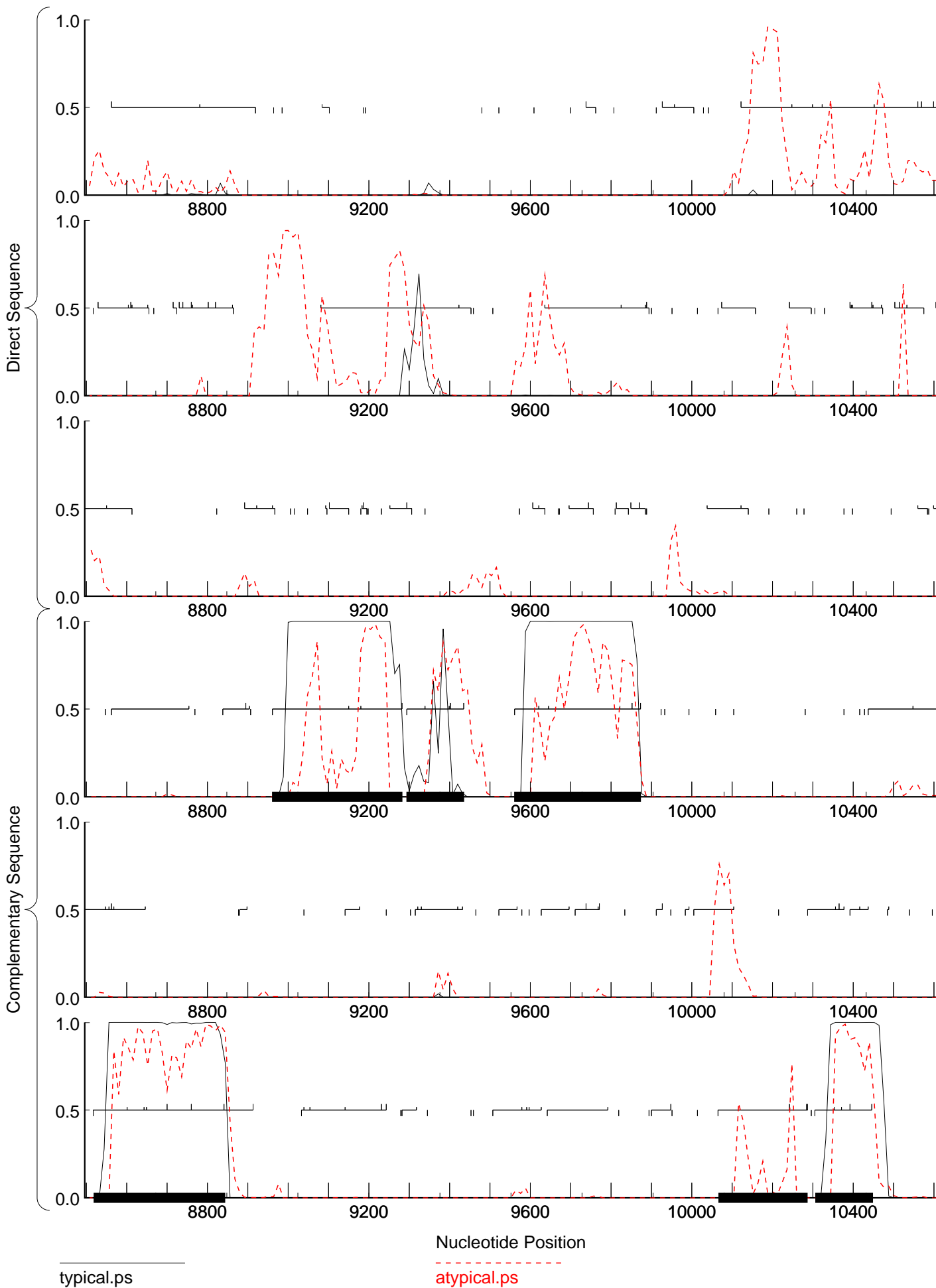
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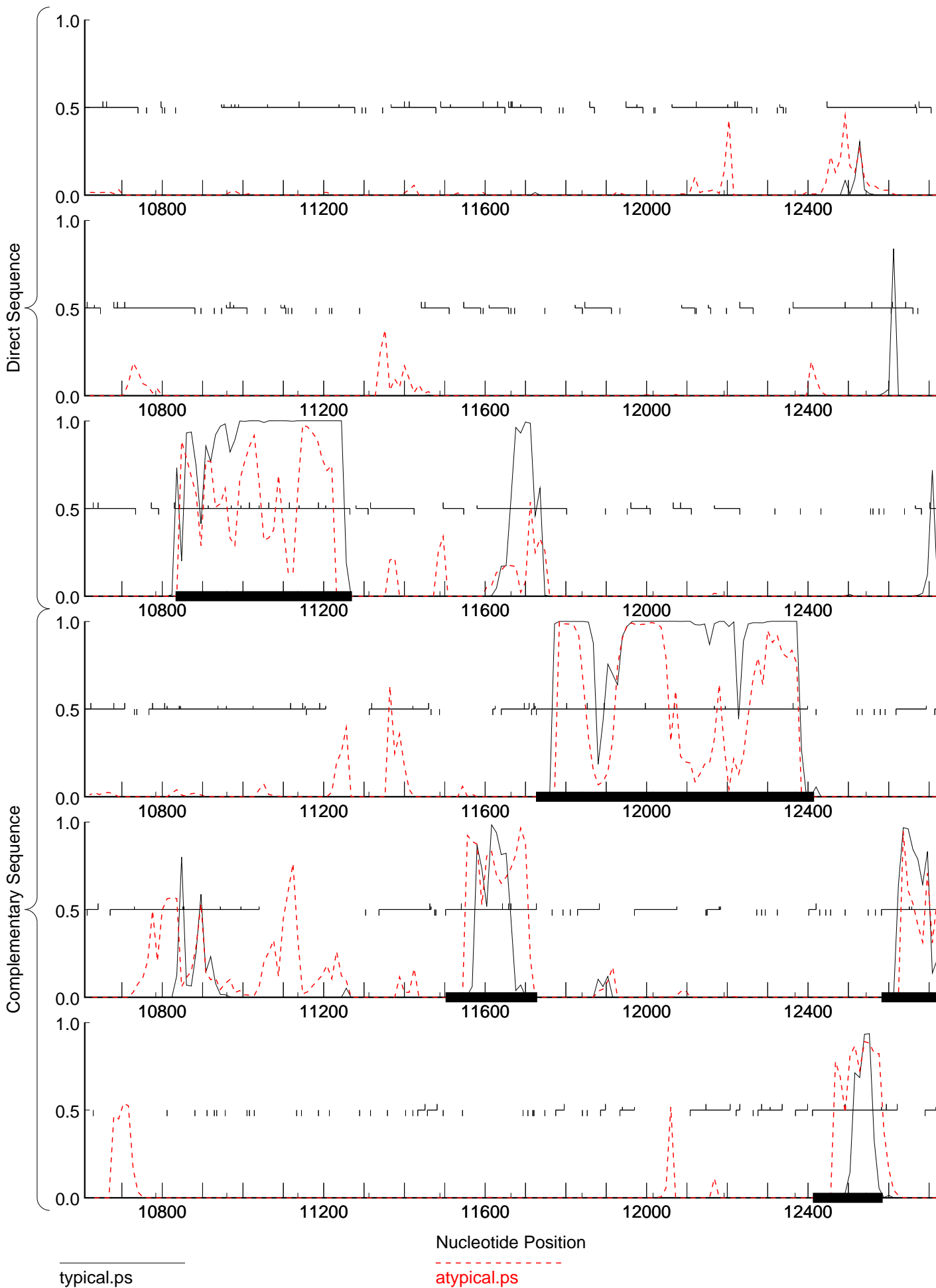


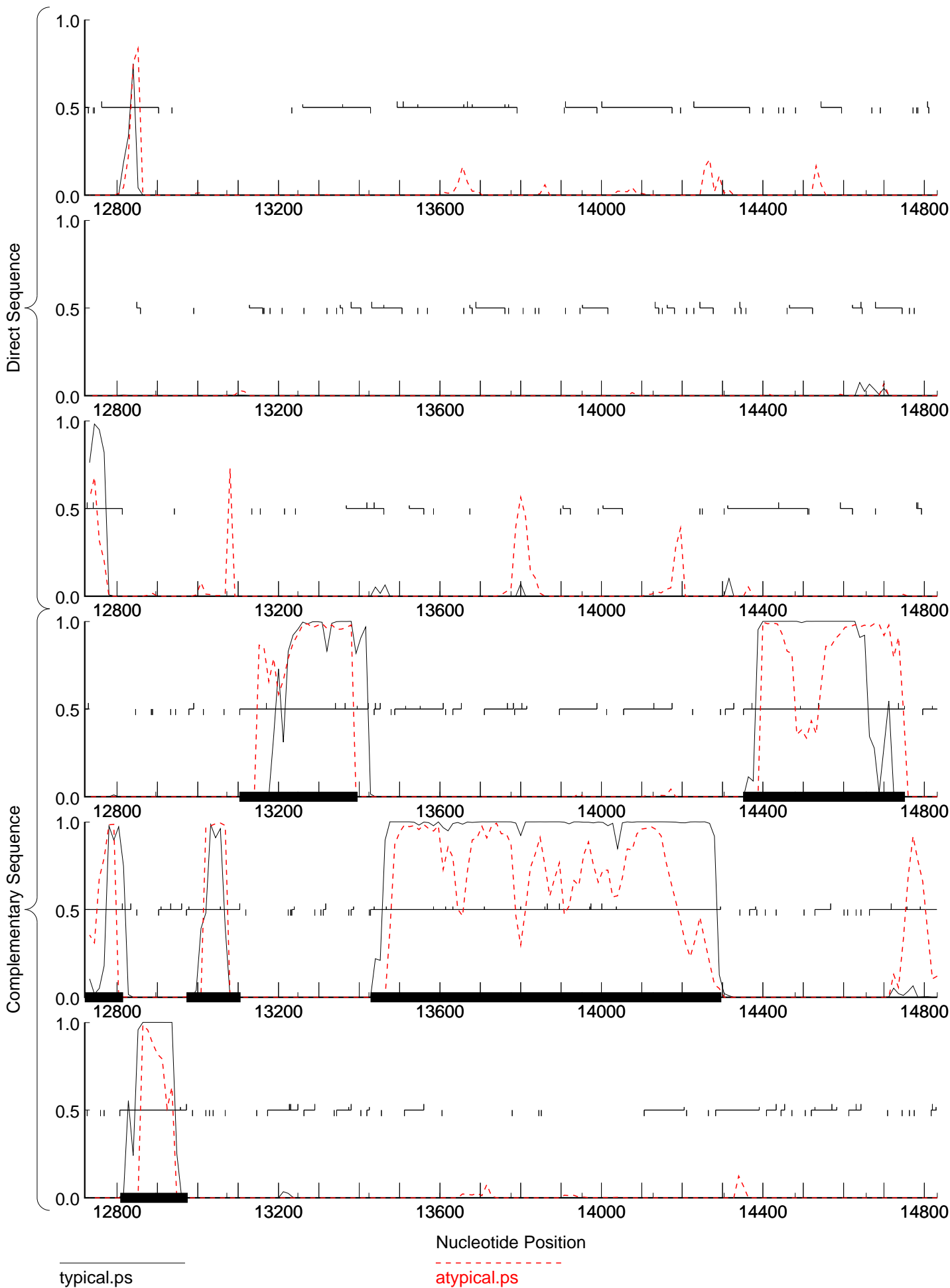
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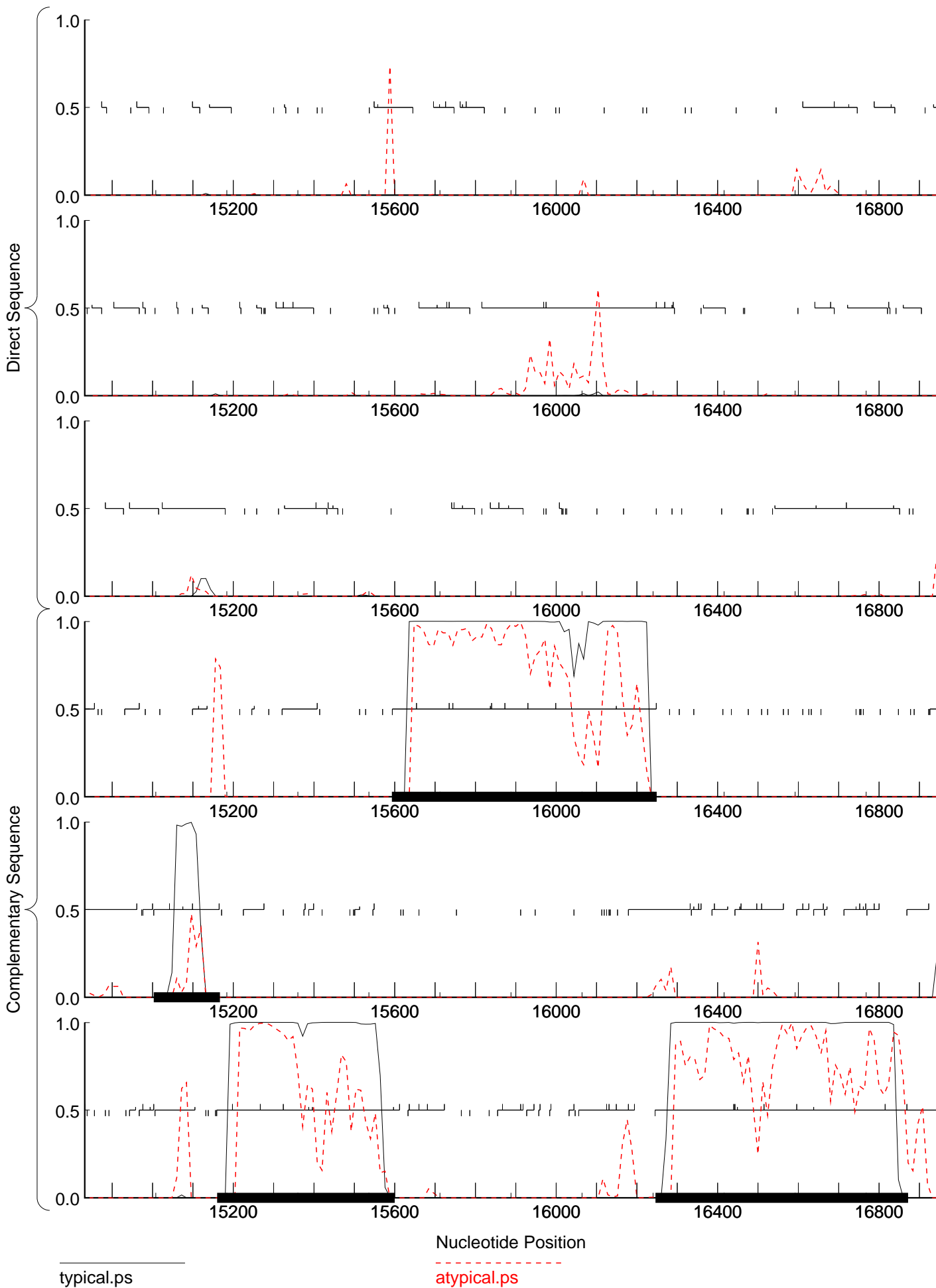
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hage Warpy complete sequence, 132996 bp including 11458 bp terminal repeat, Cluster BE1, Order 4, Window 96, Step 12, 6/64
hage Warpy complete sequence, 132996 bp including 11458 bp terminal repeat, Cluster BE1, Order 2, Window 96, Step 12, 6/64



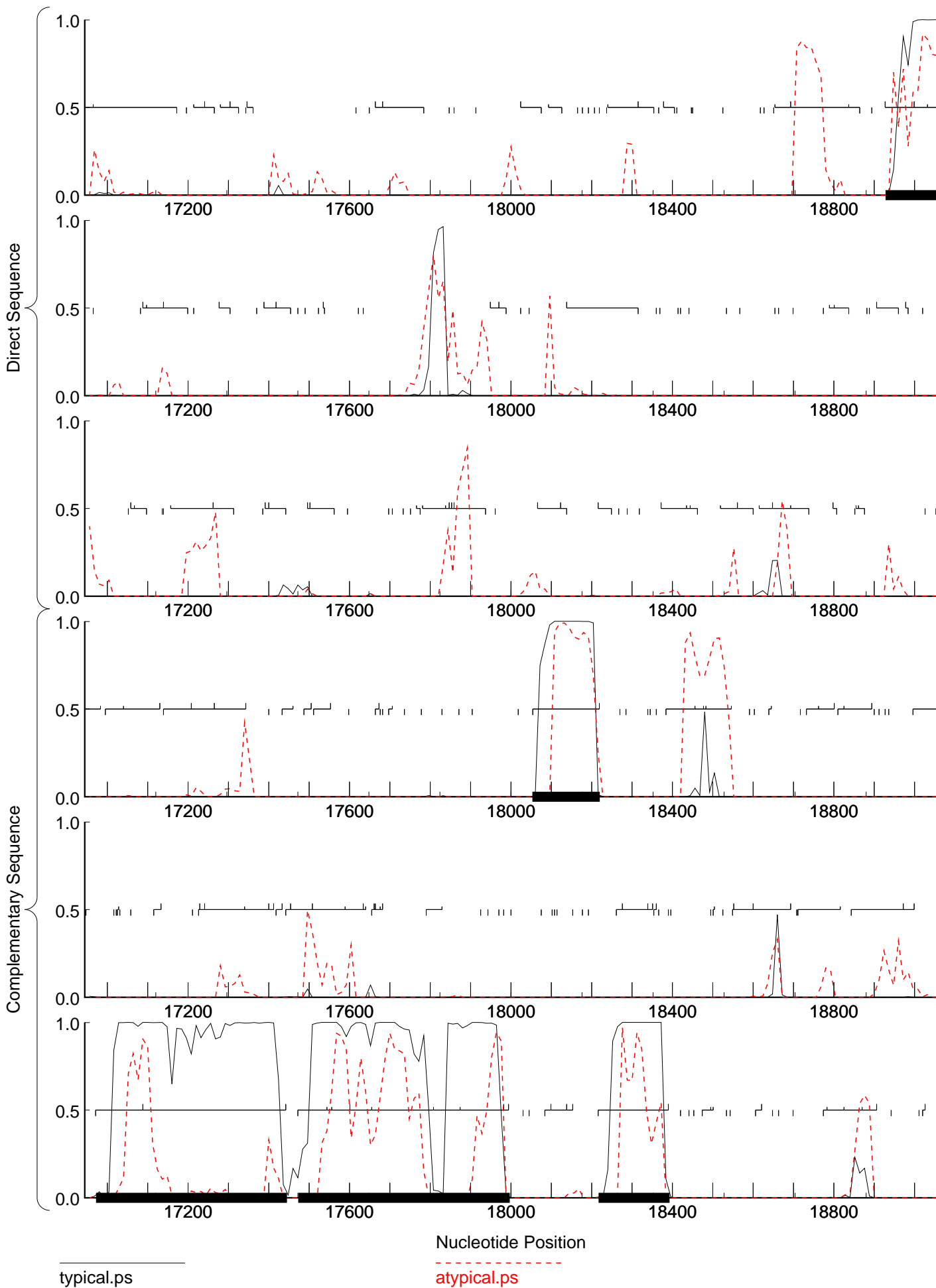


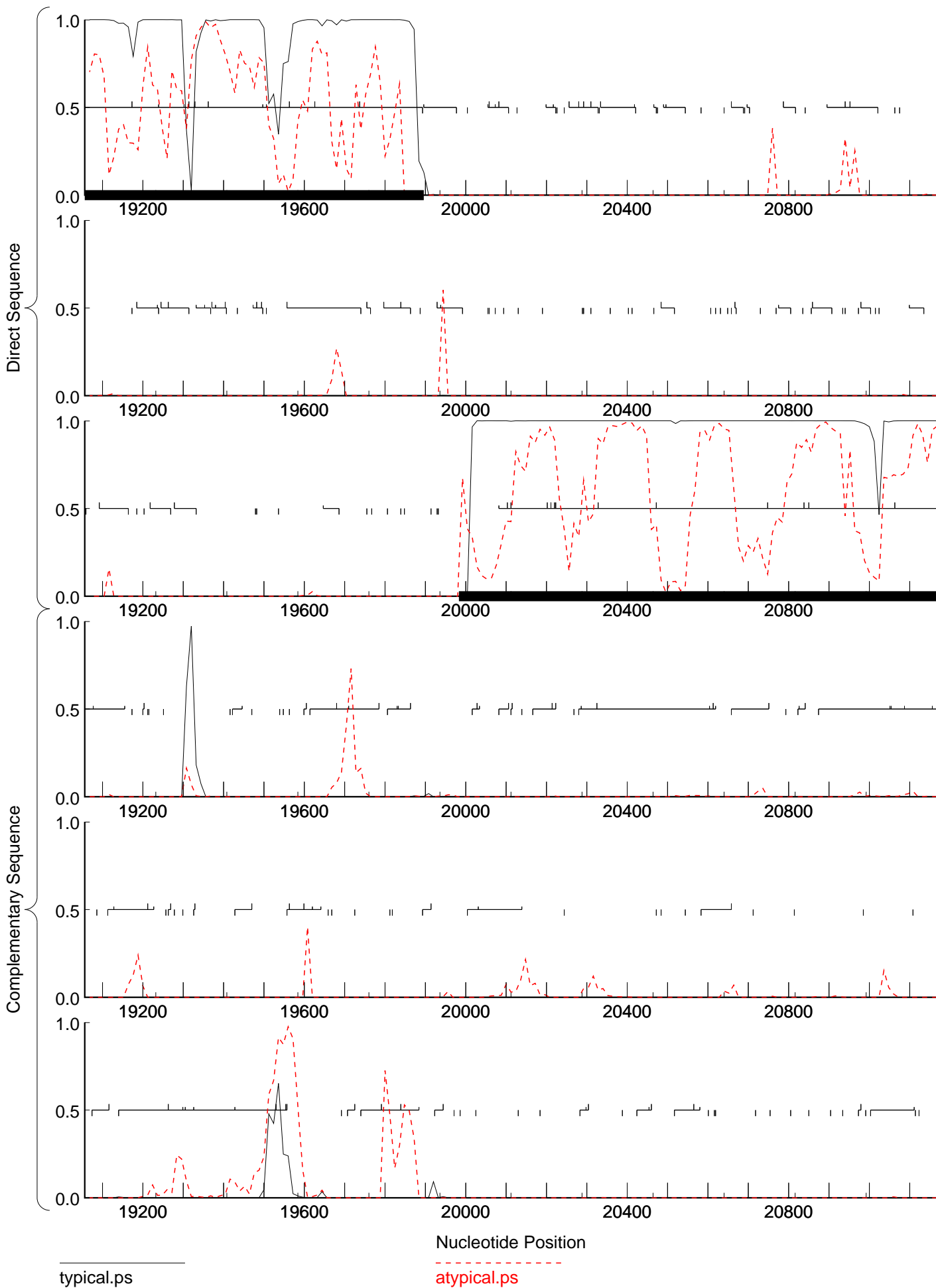


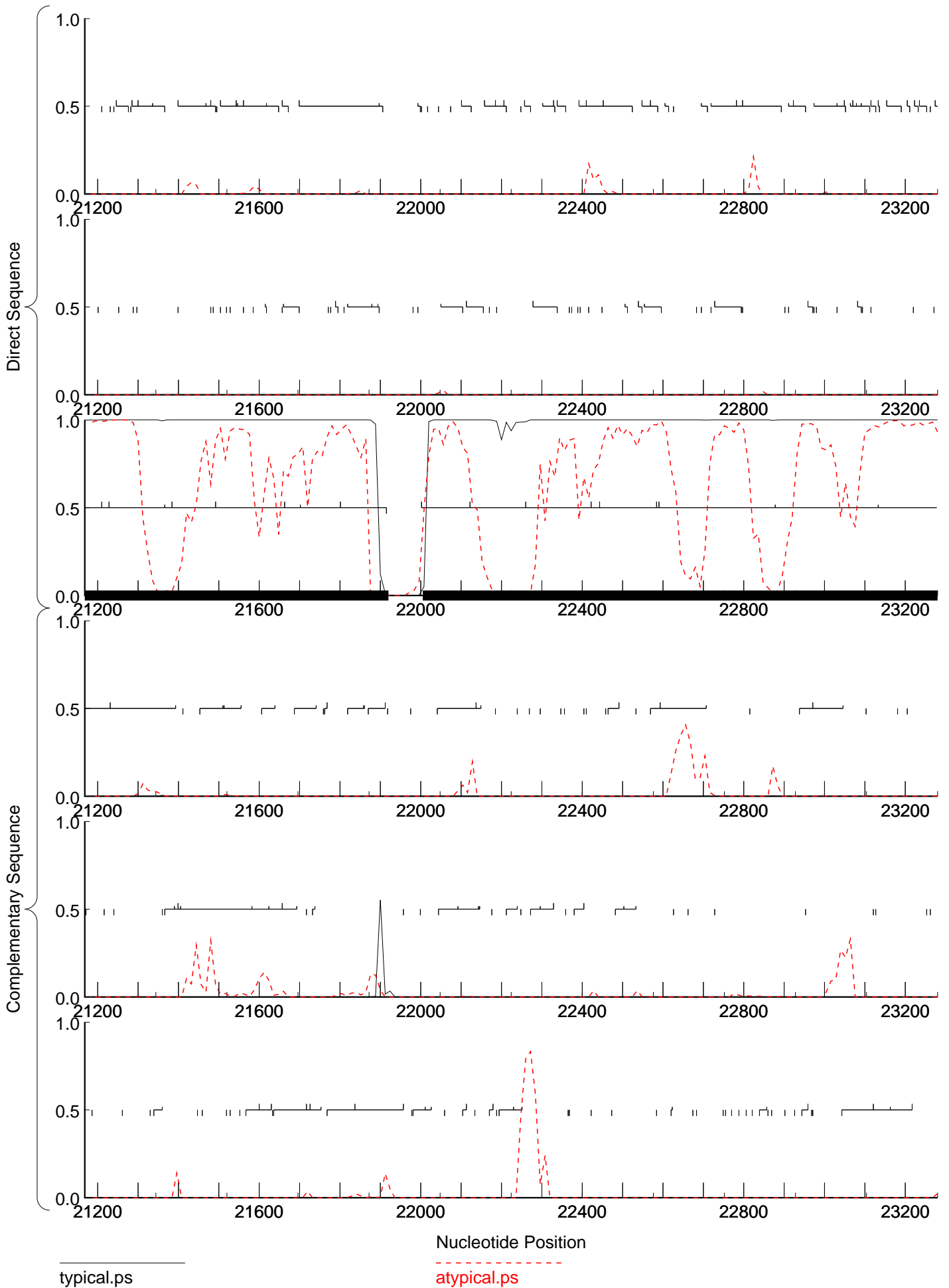


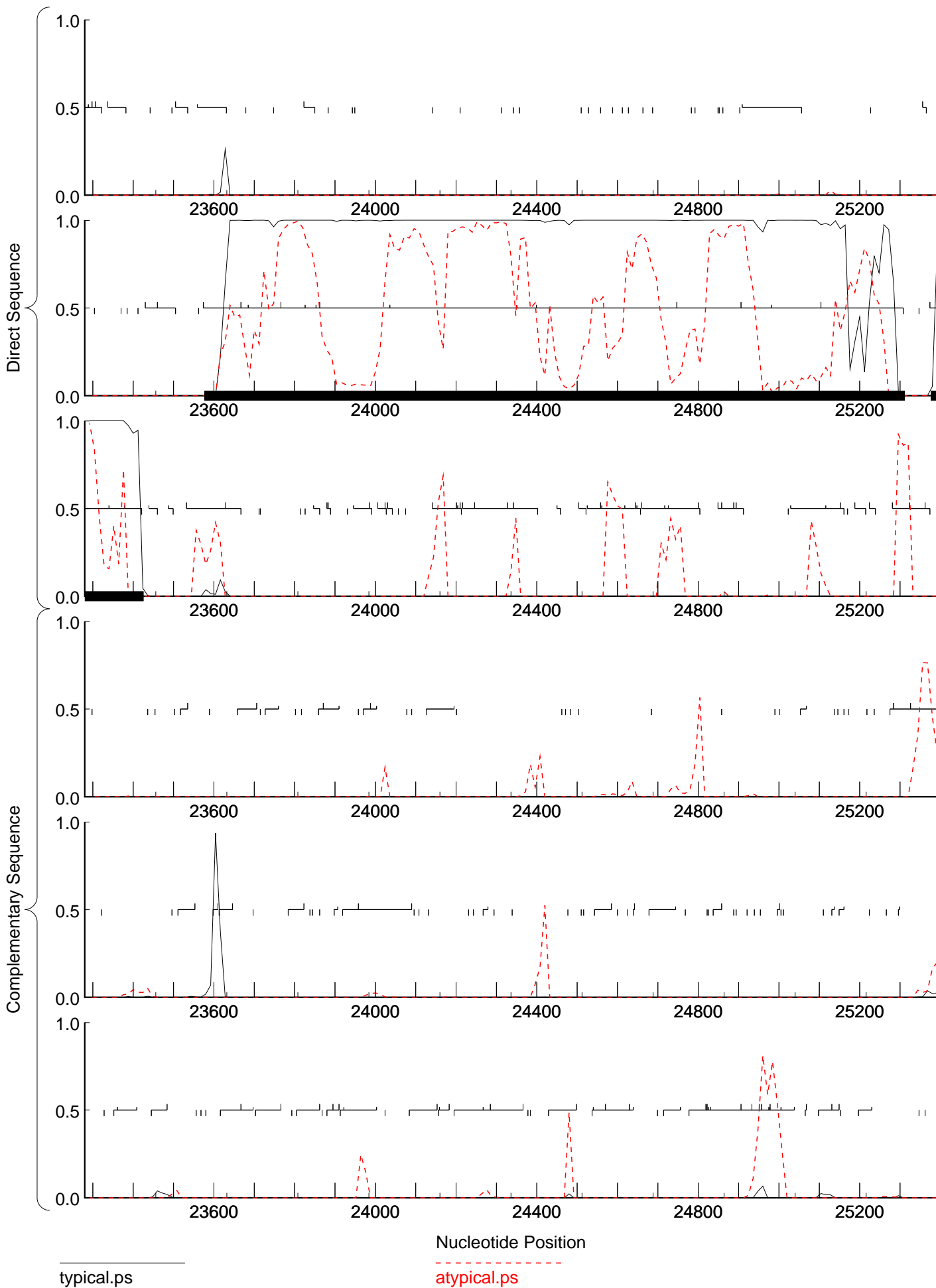
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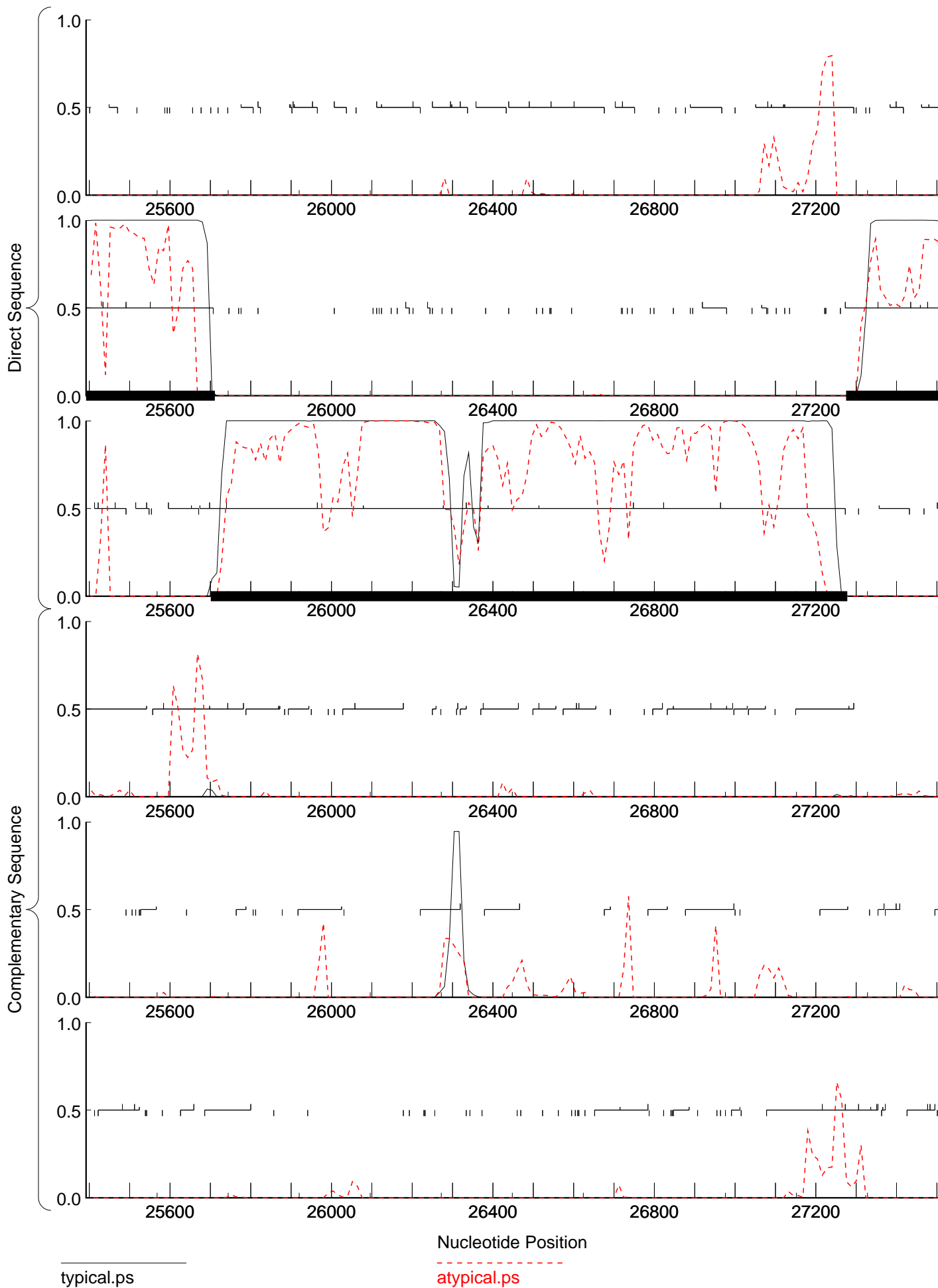
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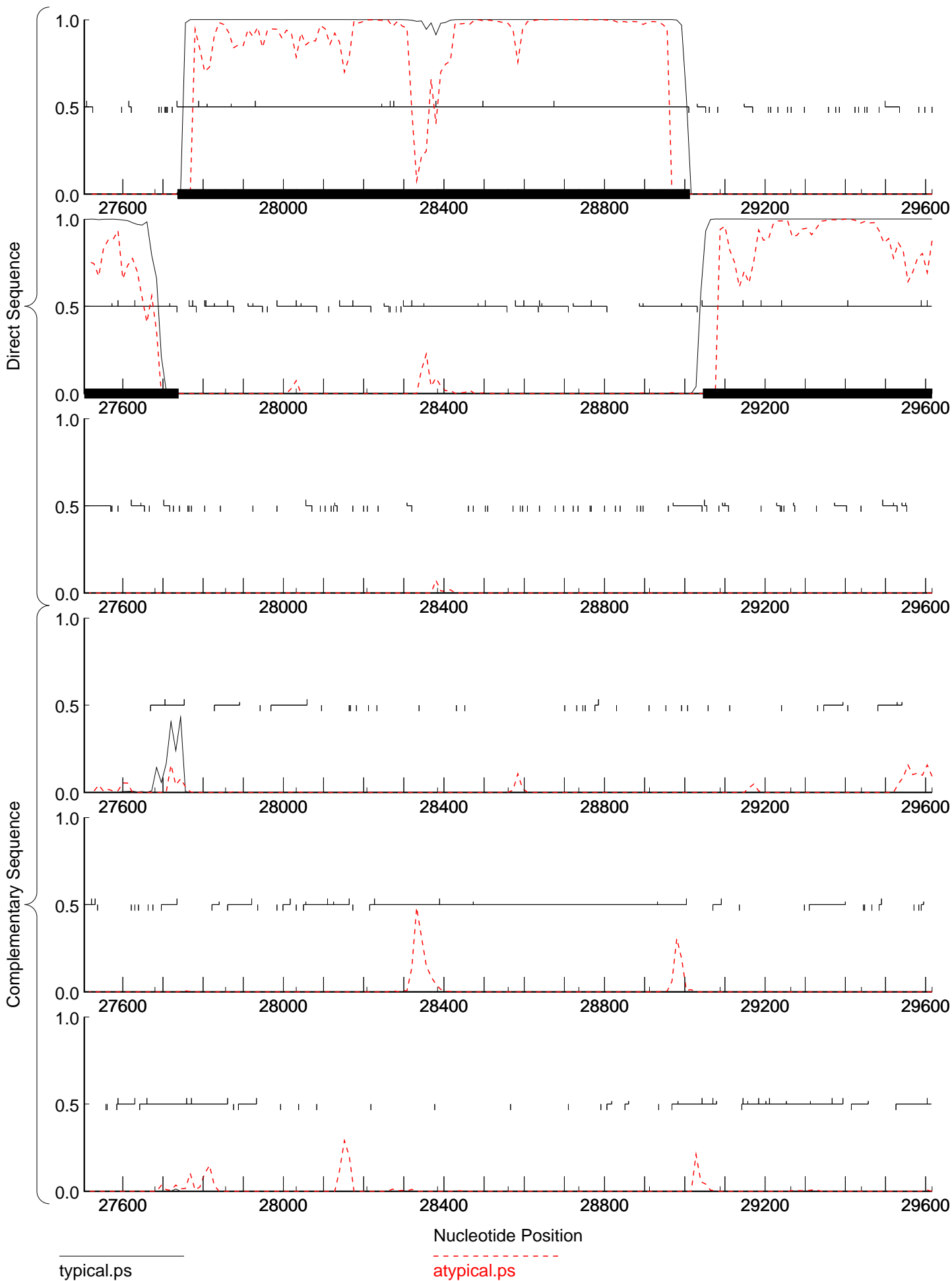






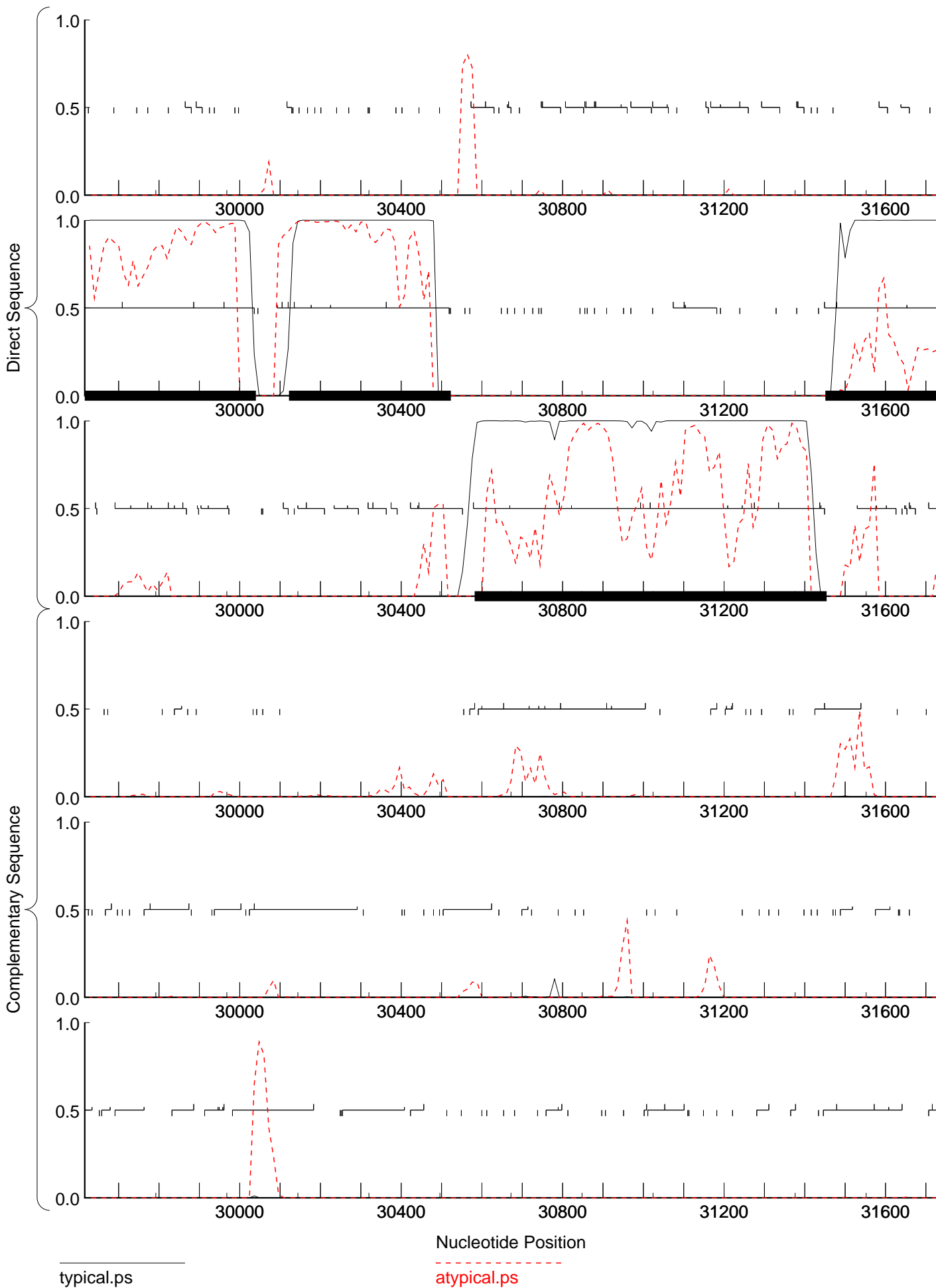






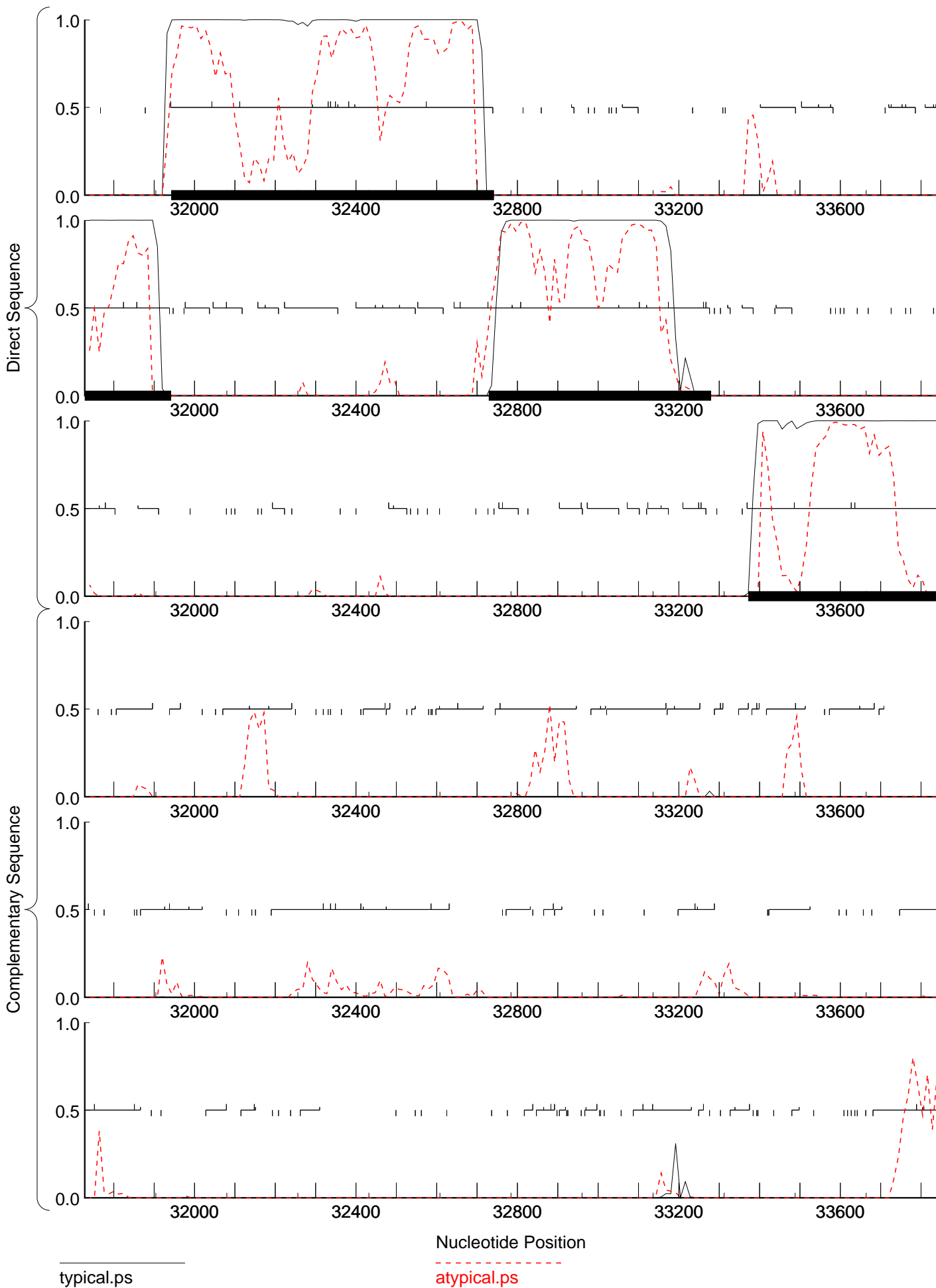
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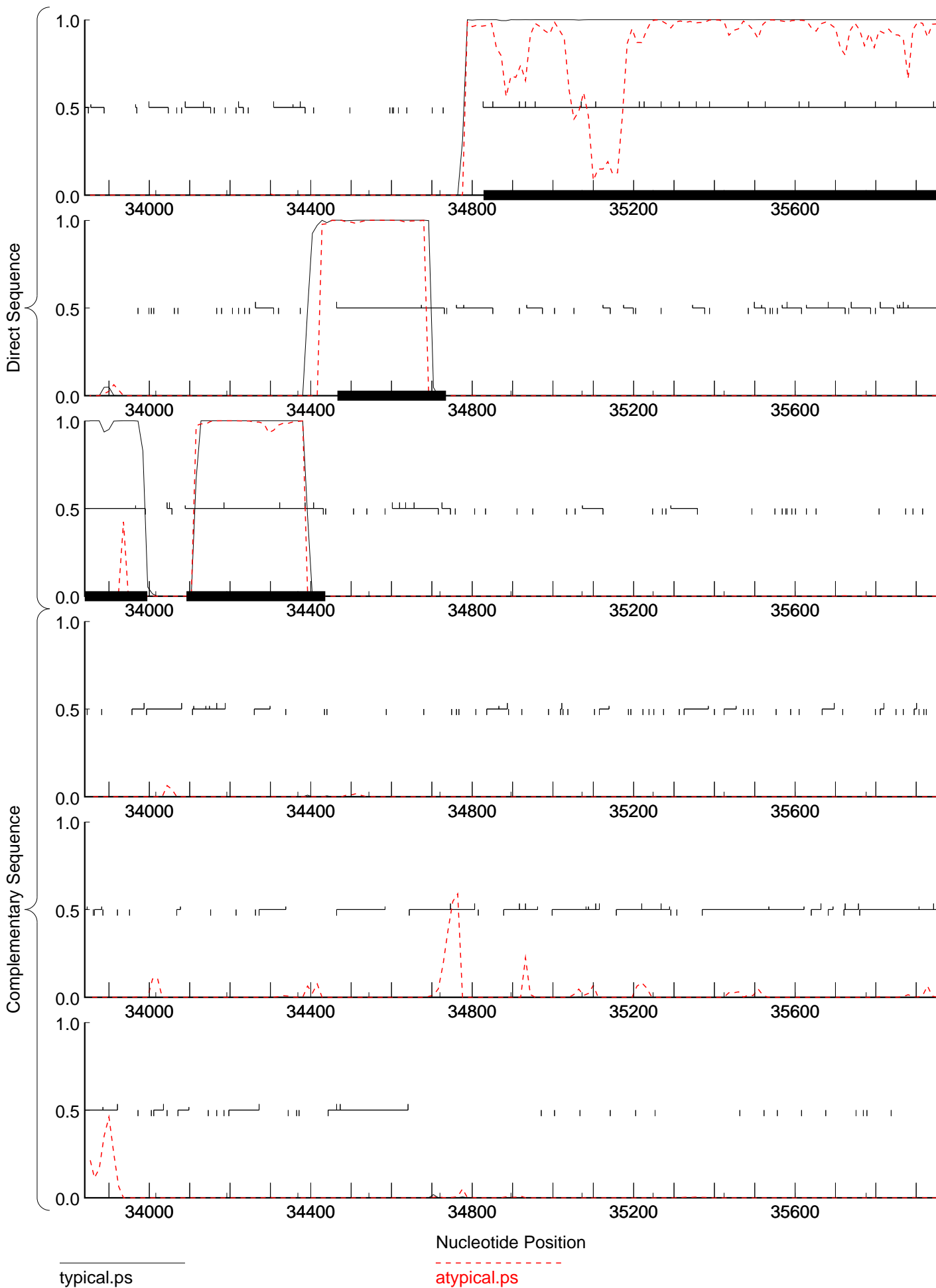
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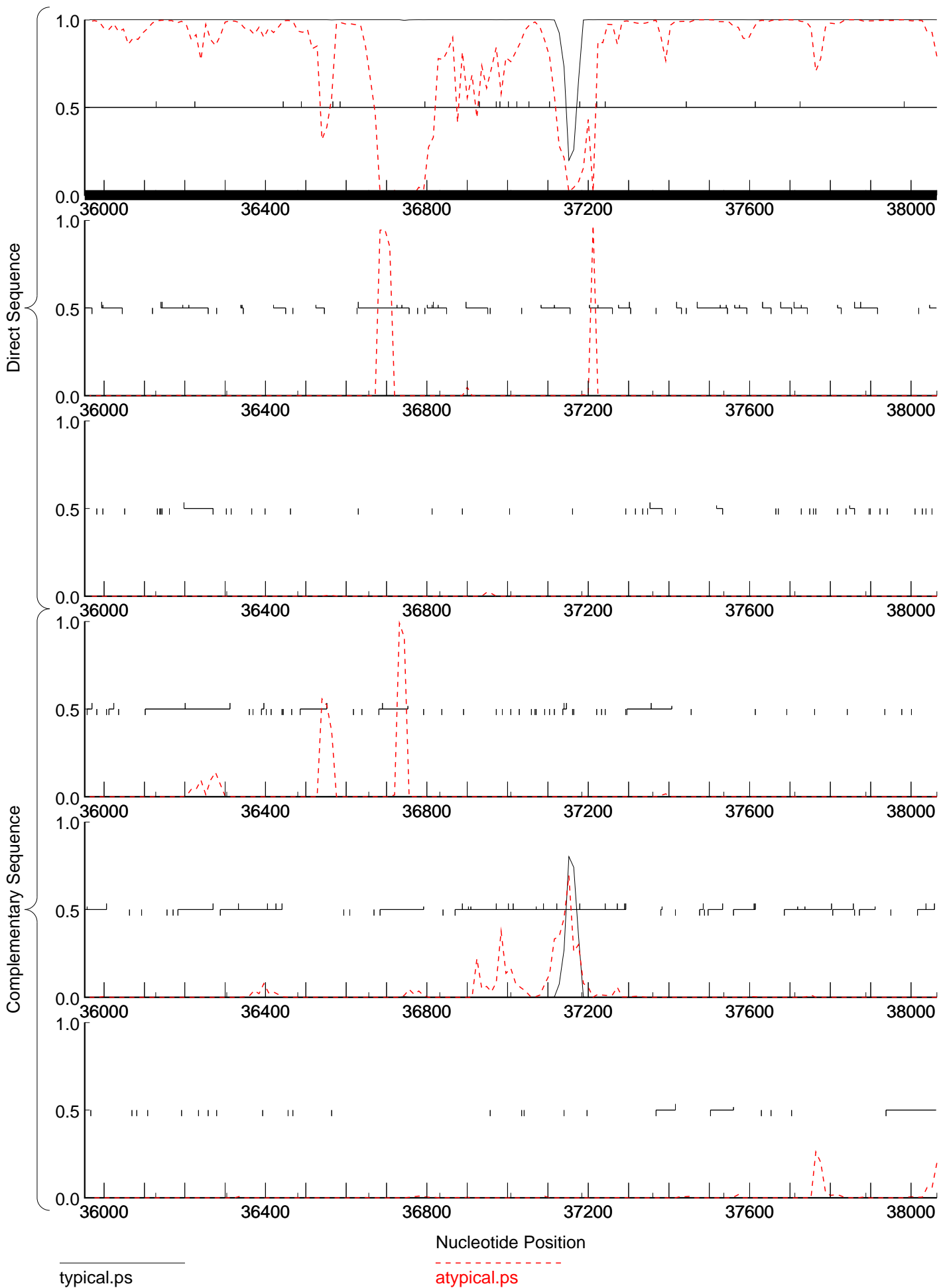
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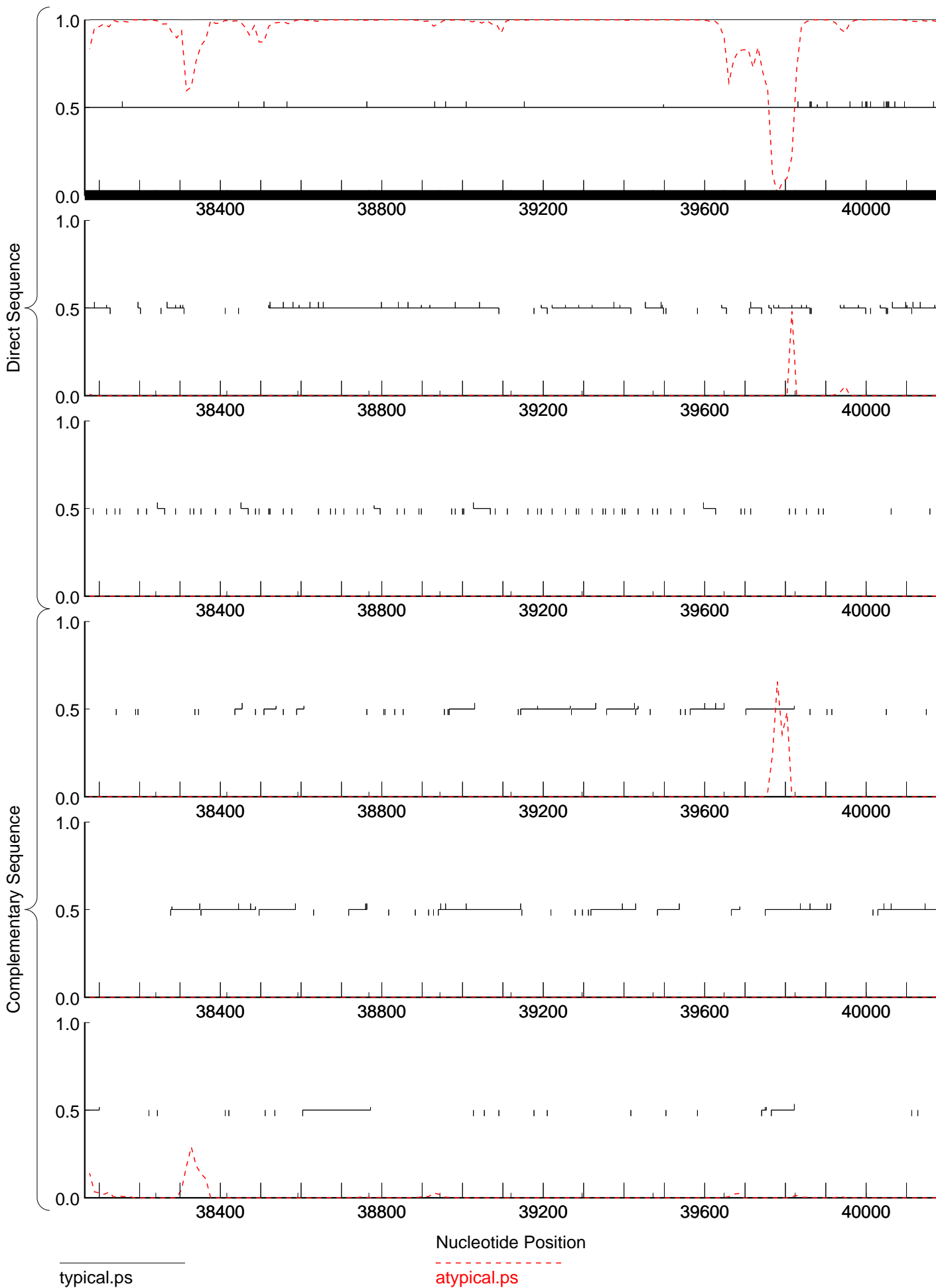


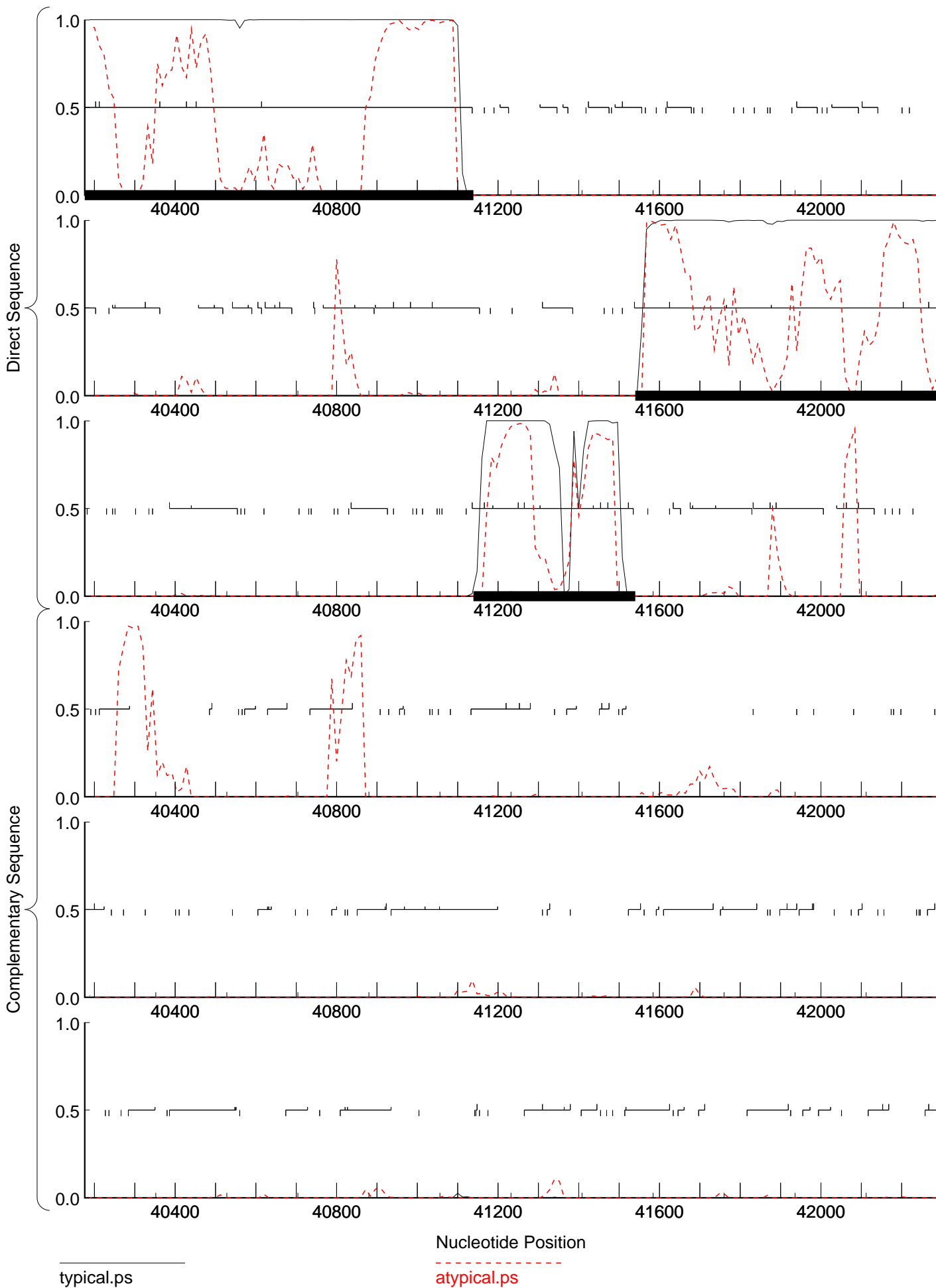
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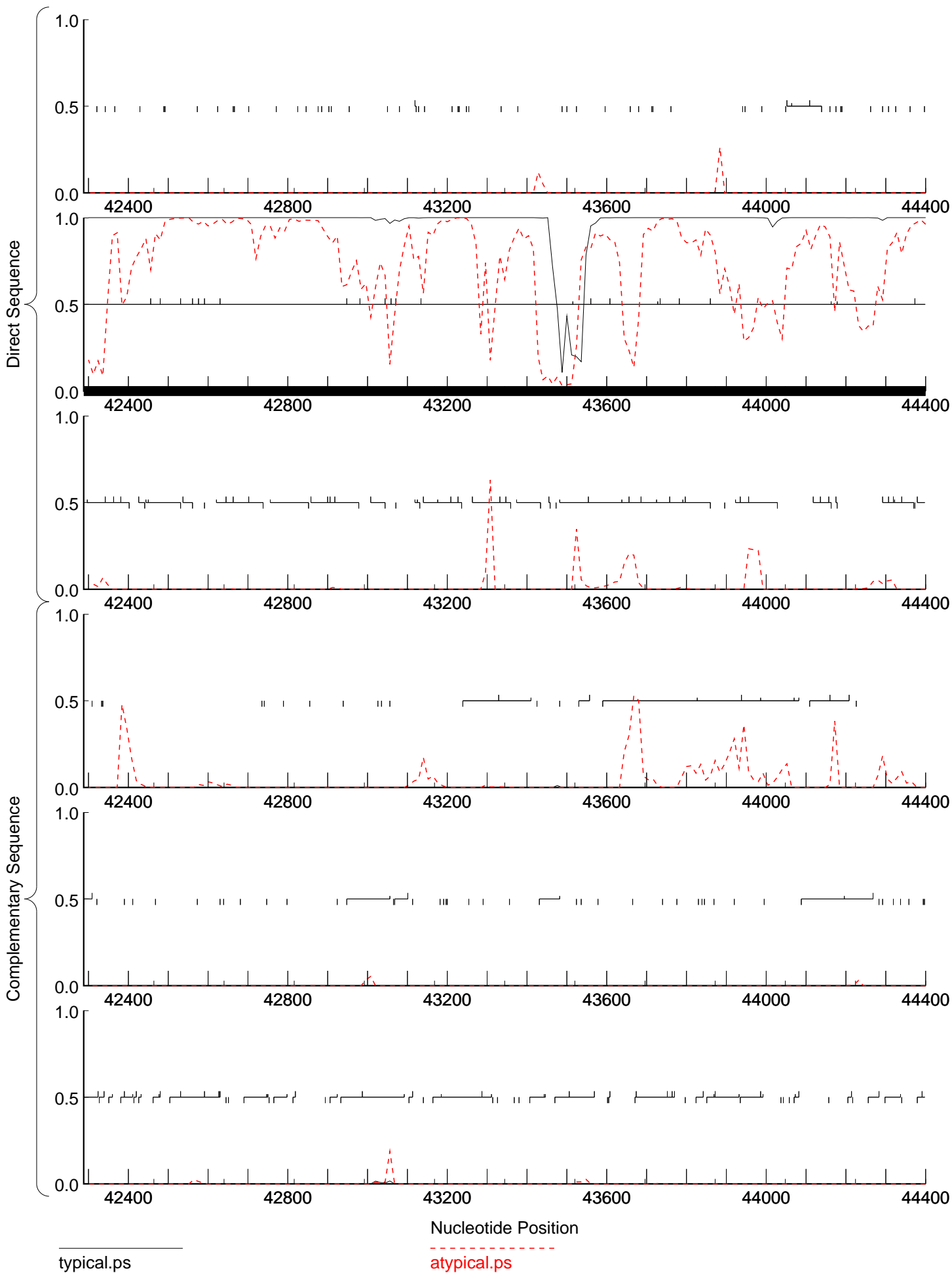
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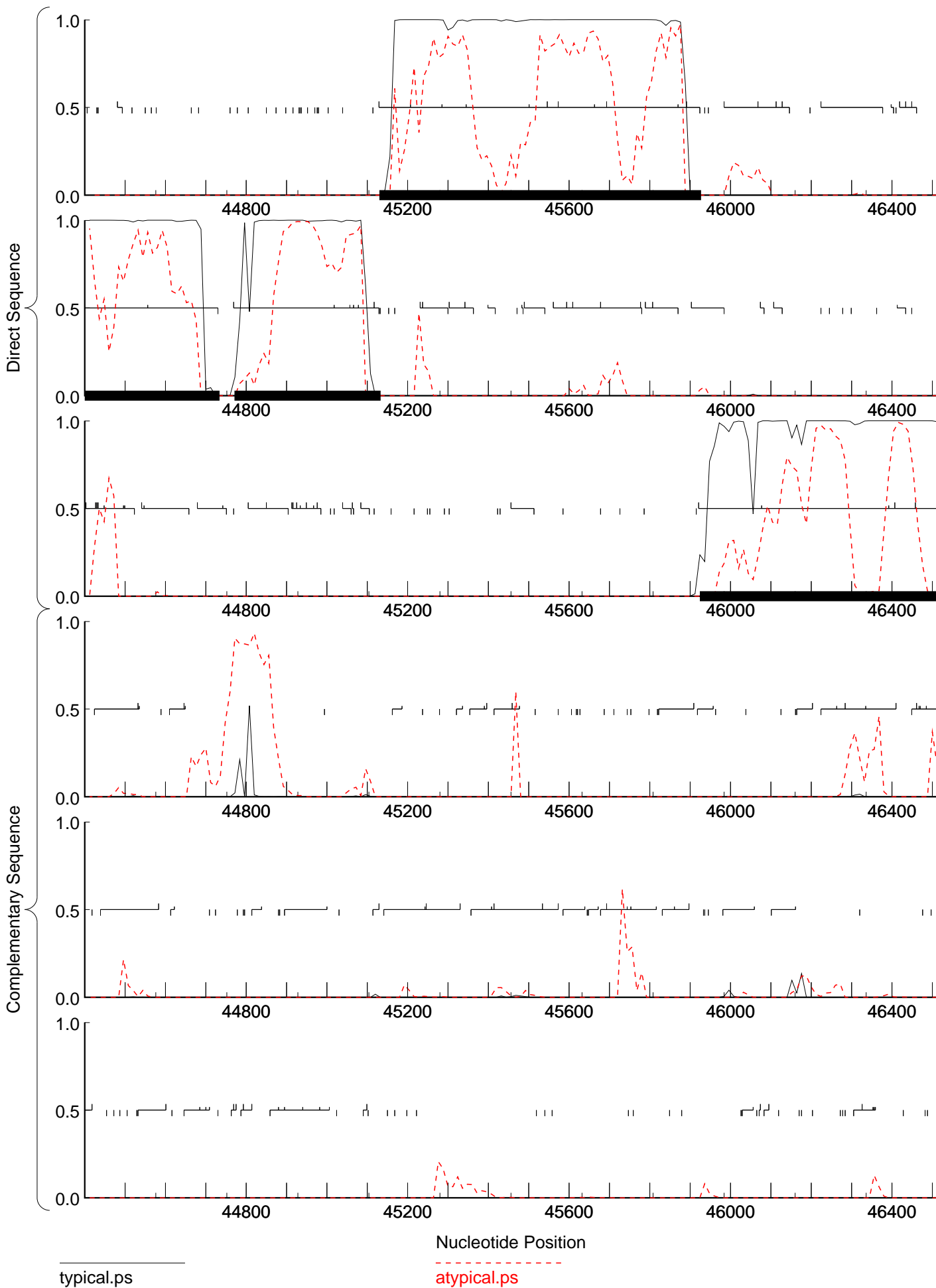






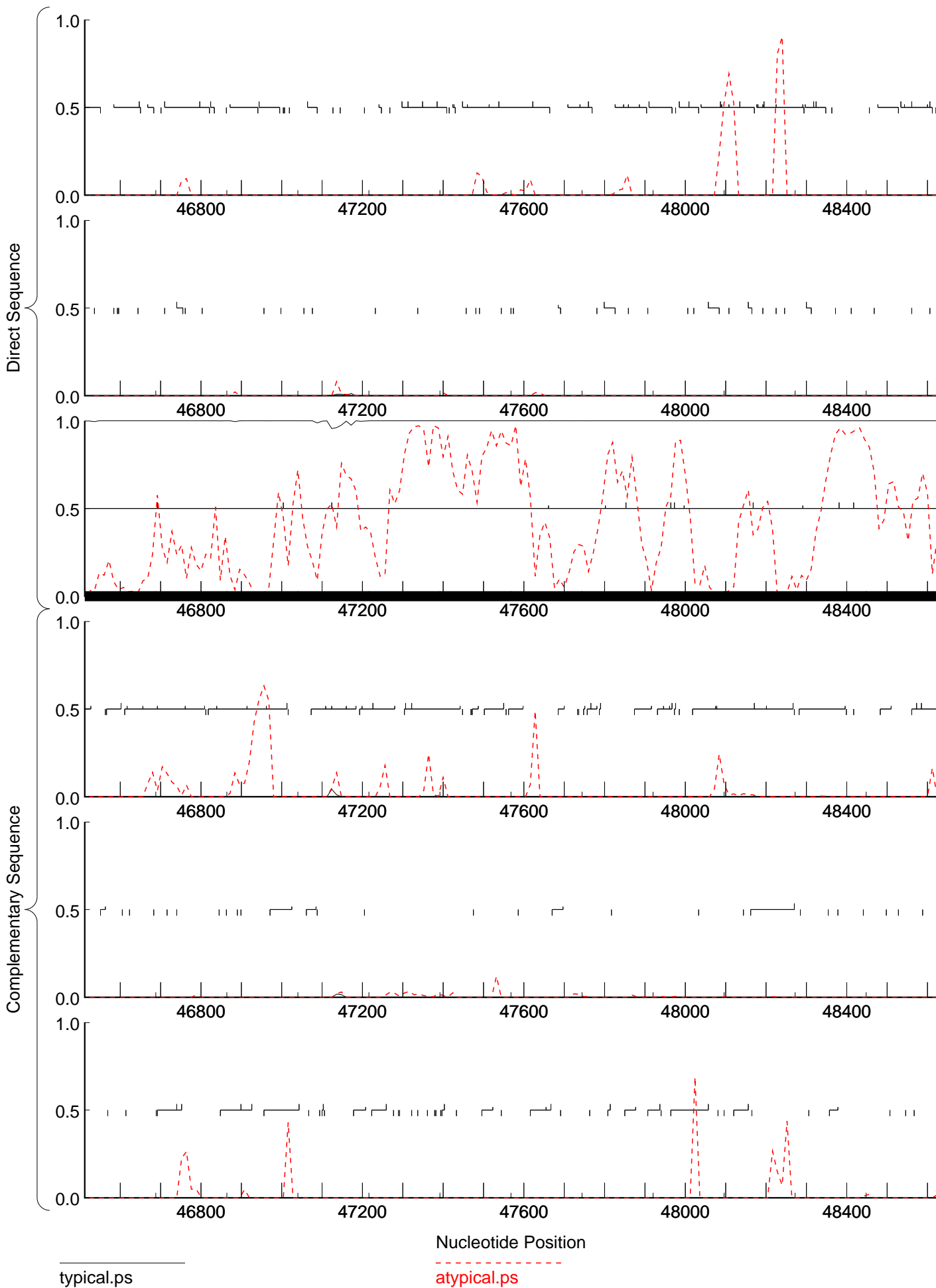


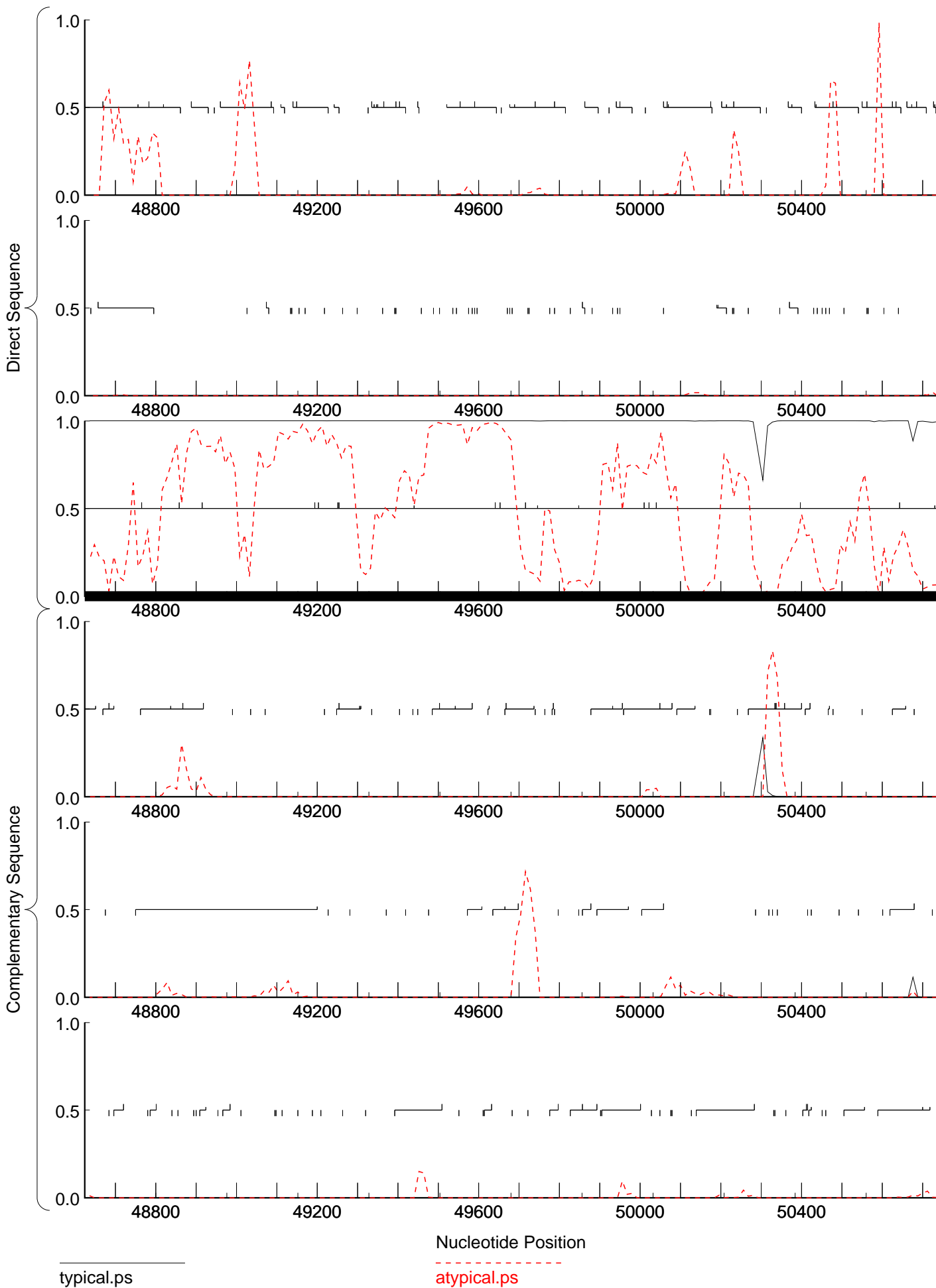


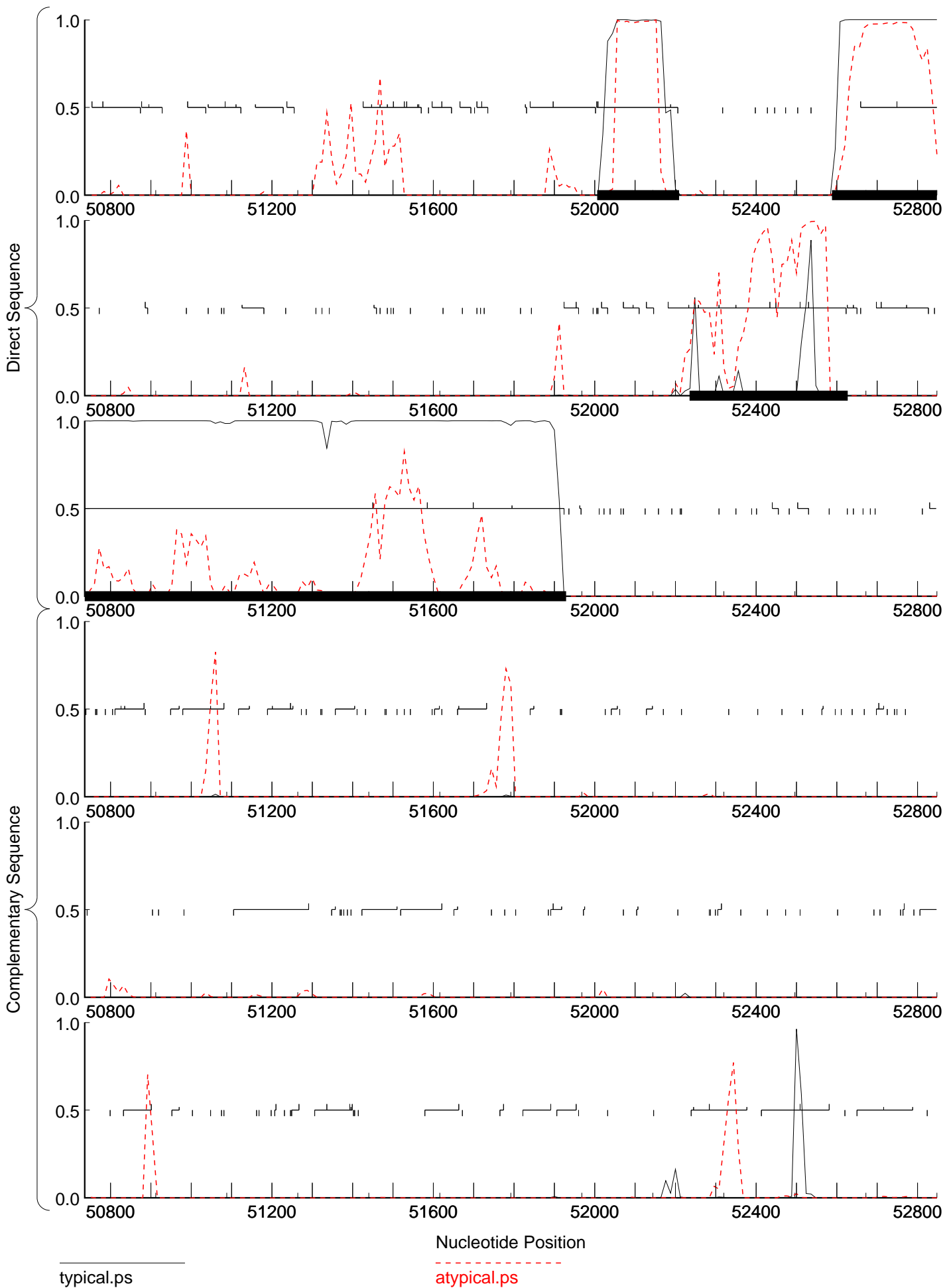


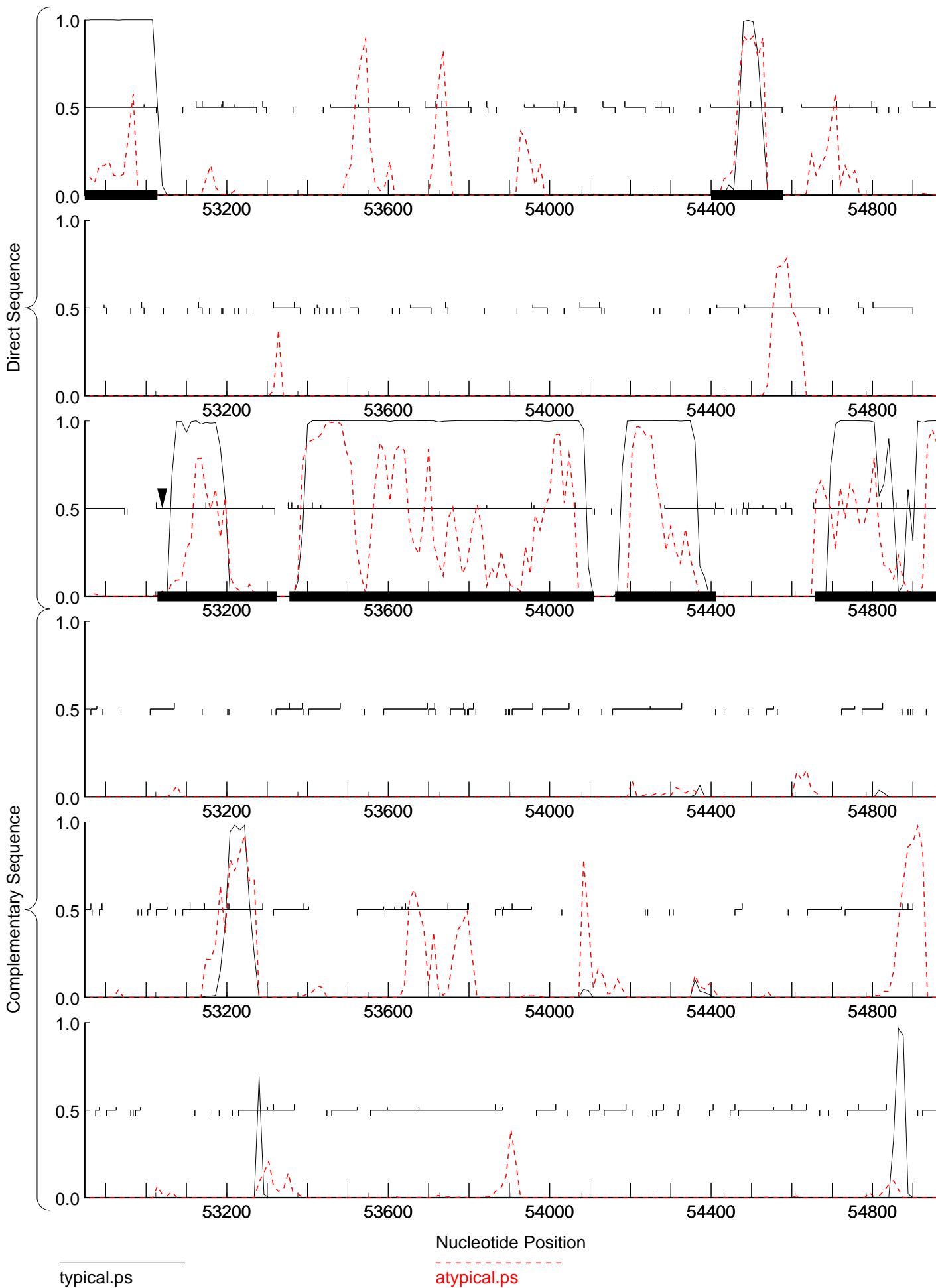
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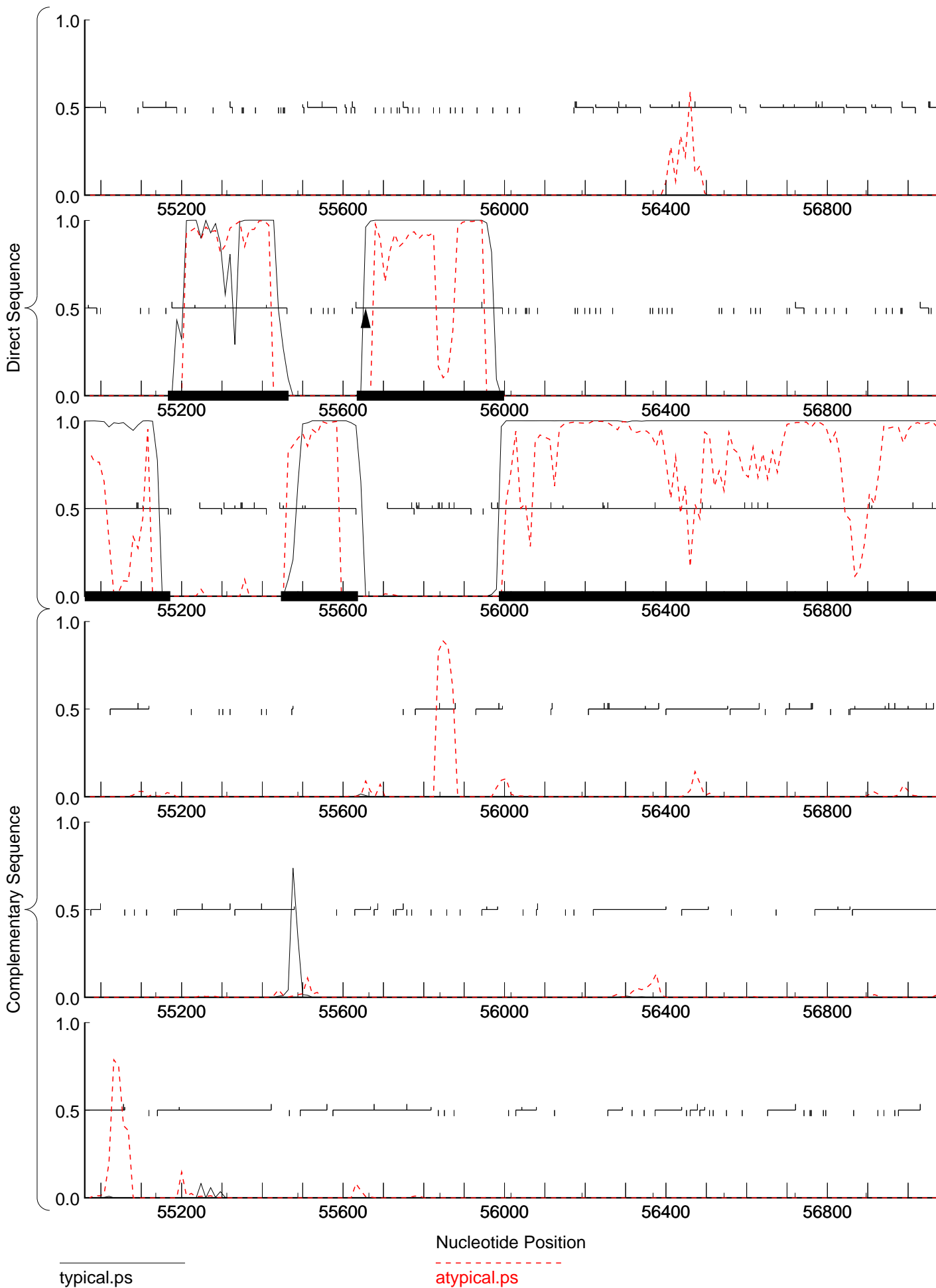


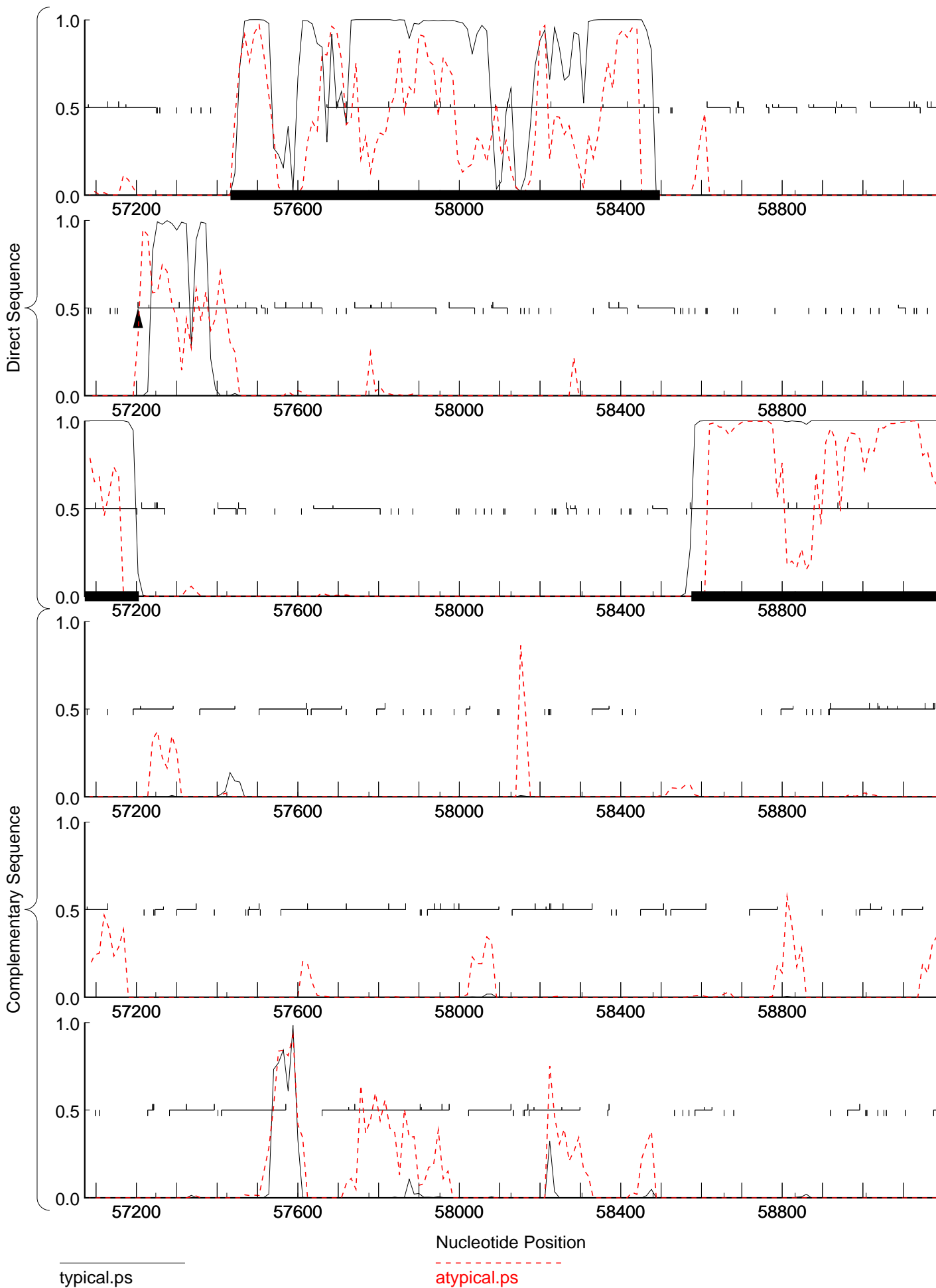




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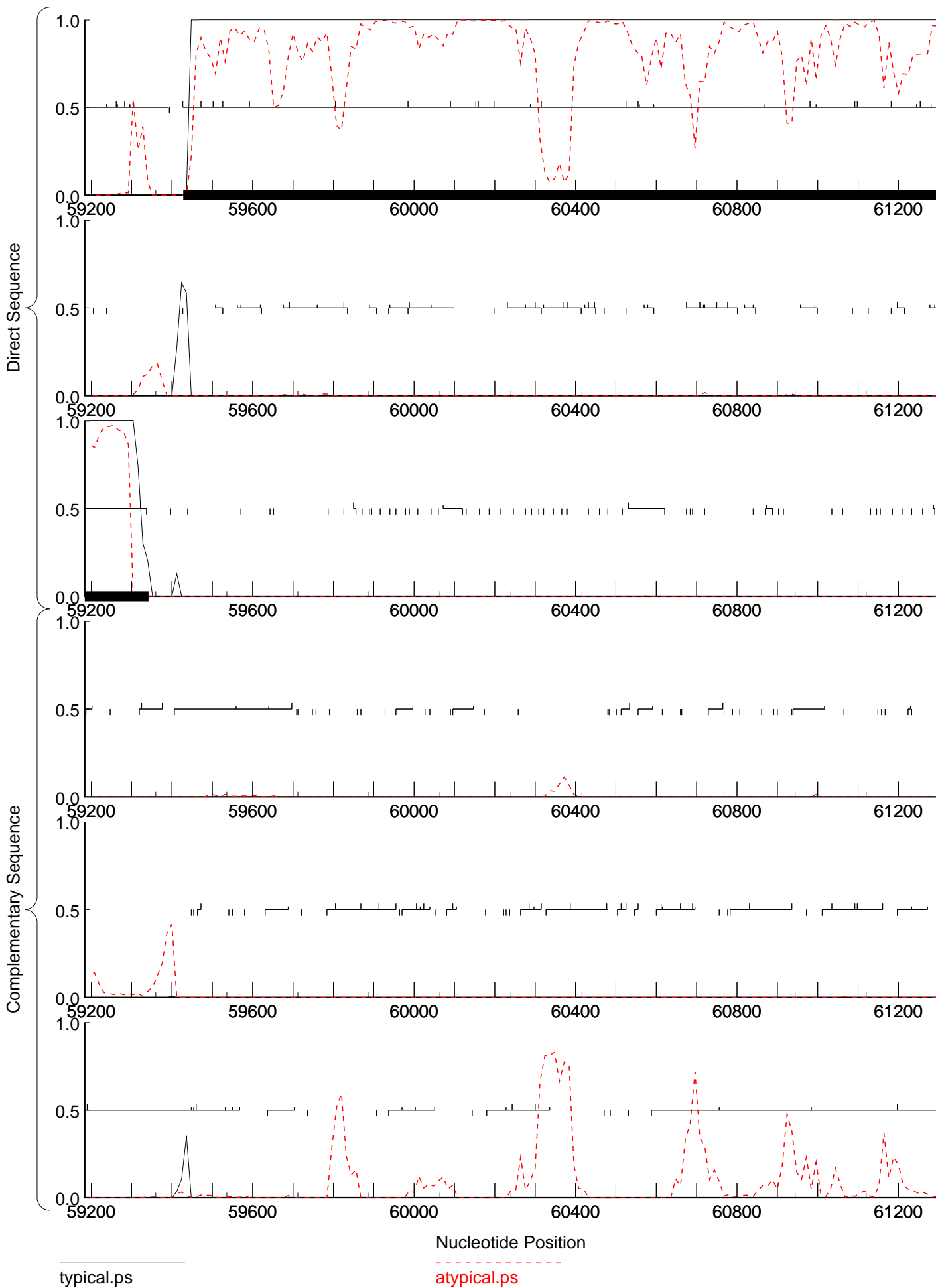
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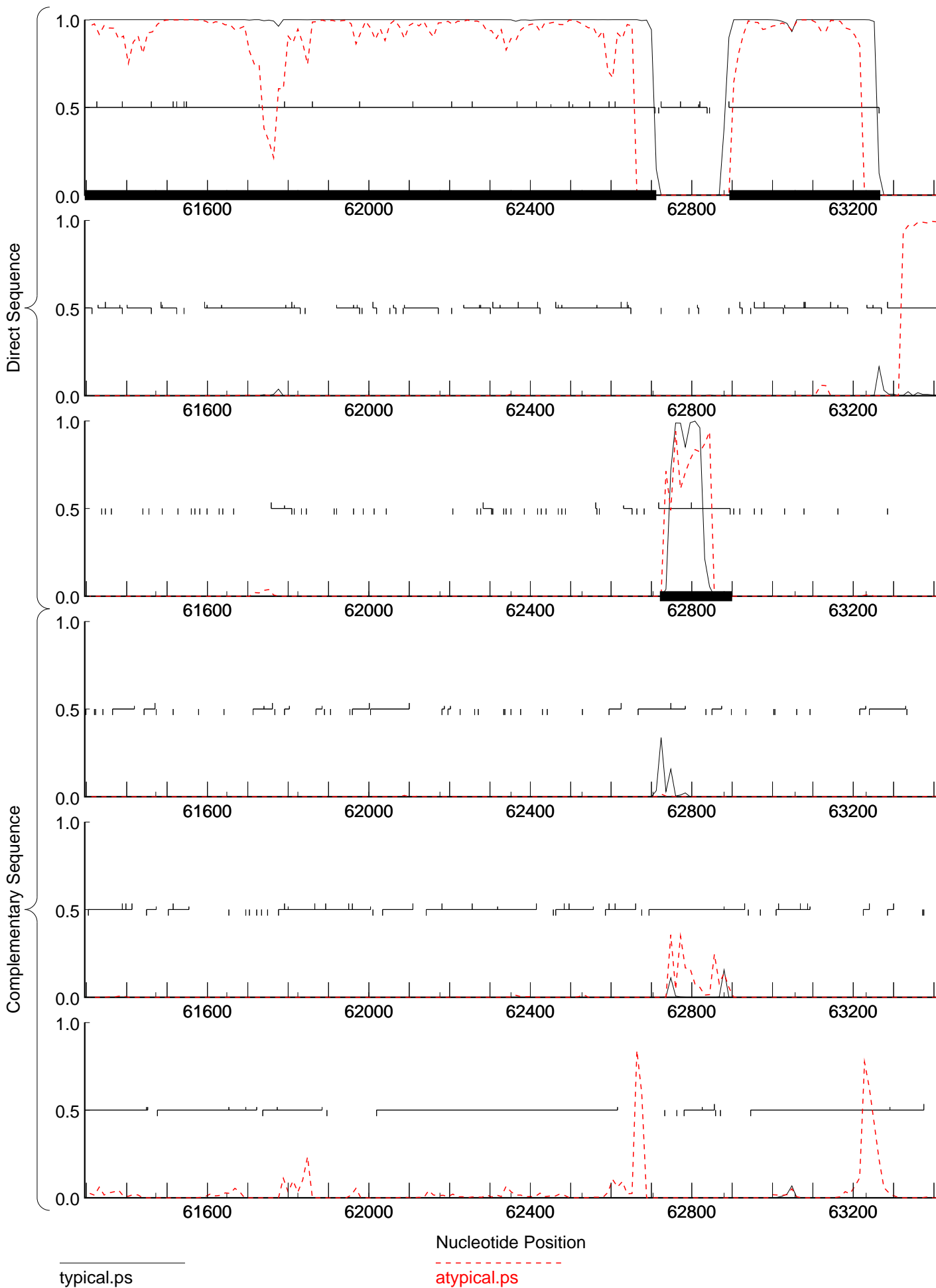




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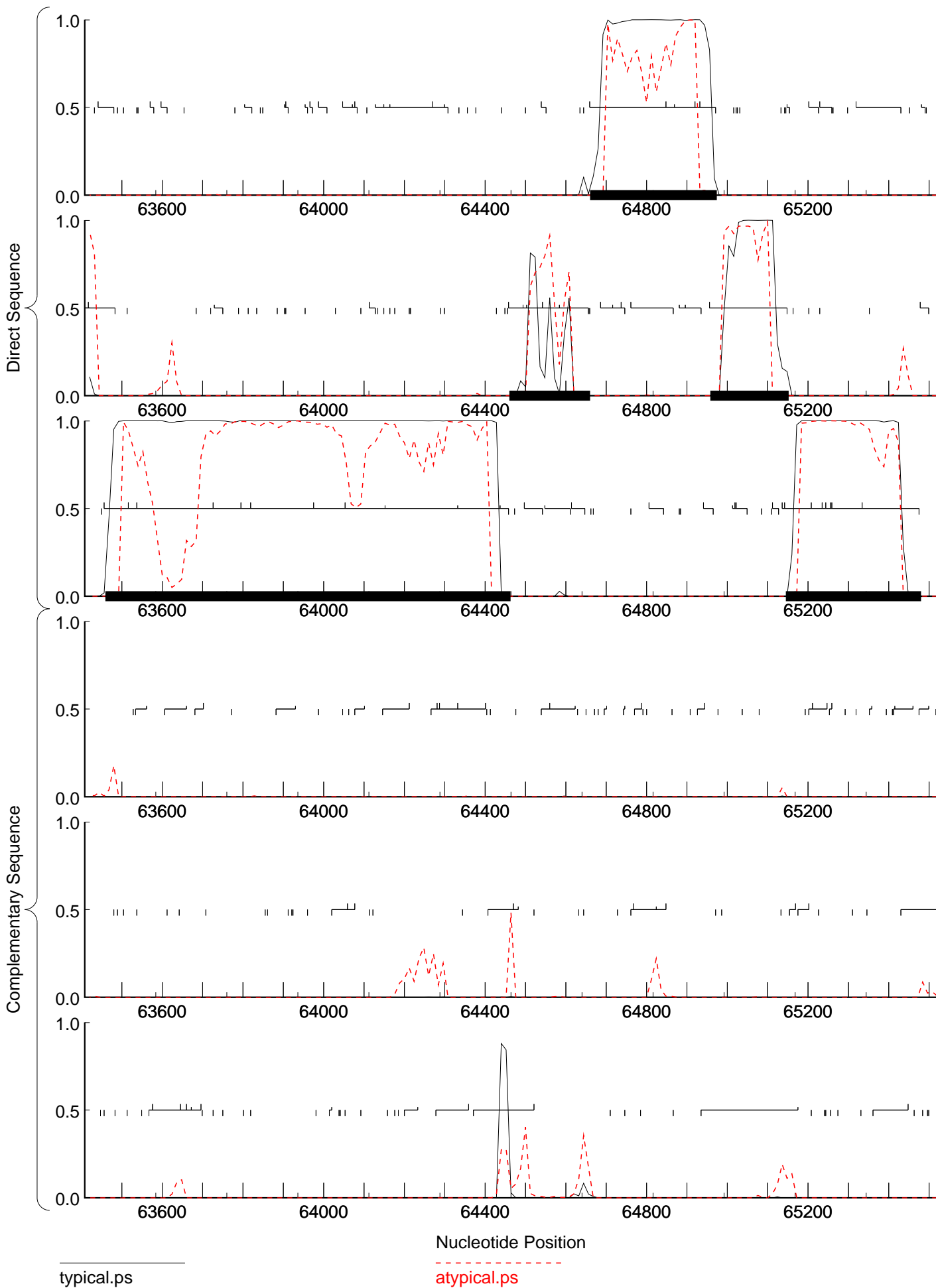
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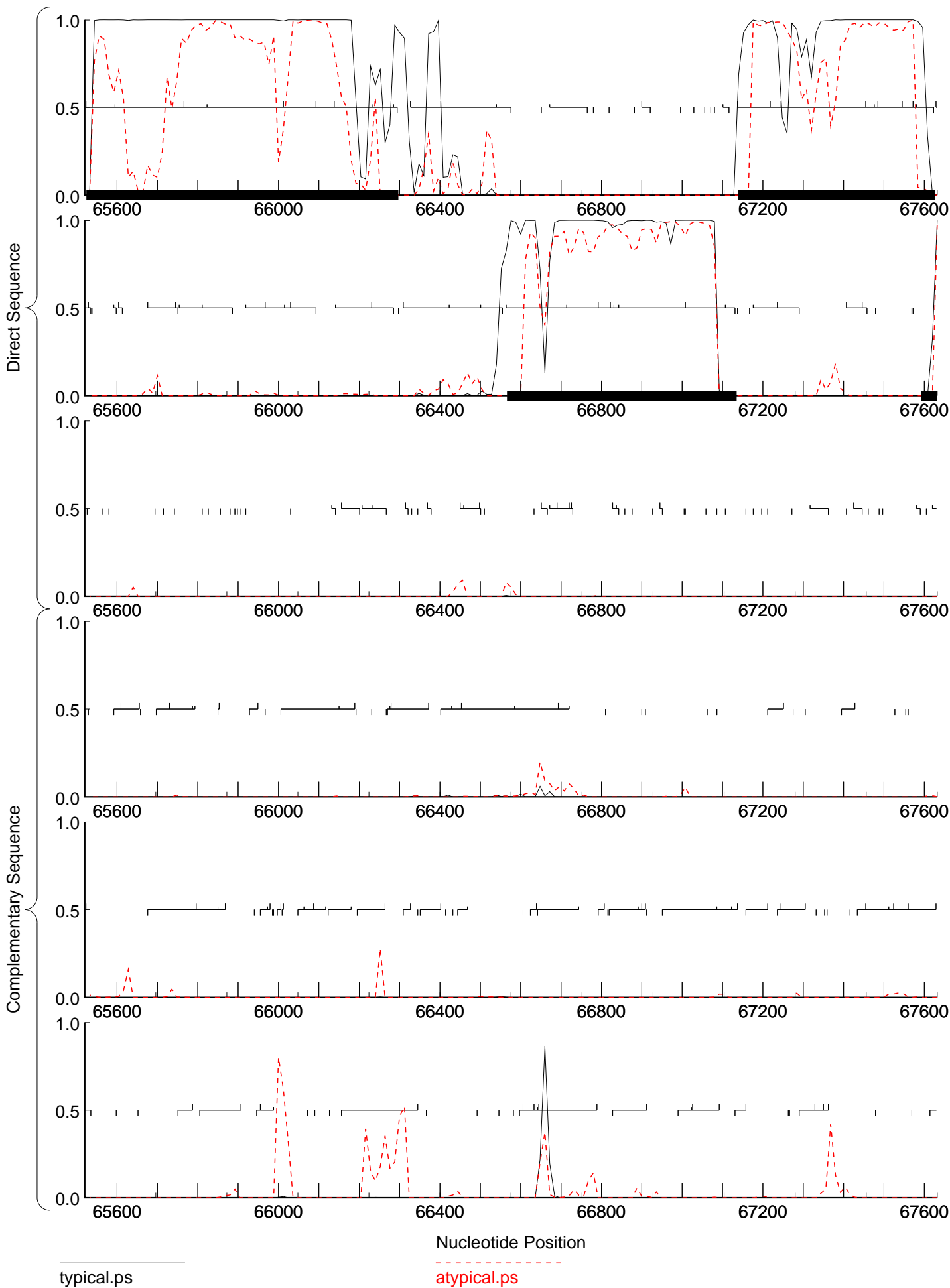




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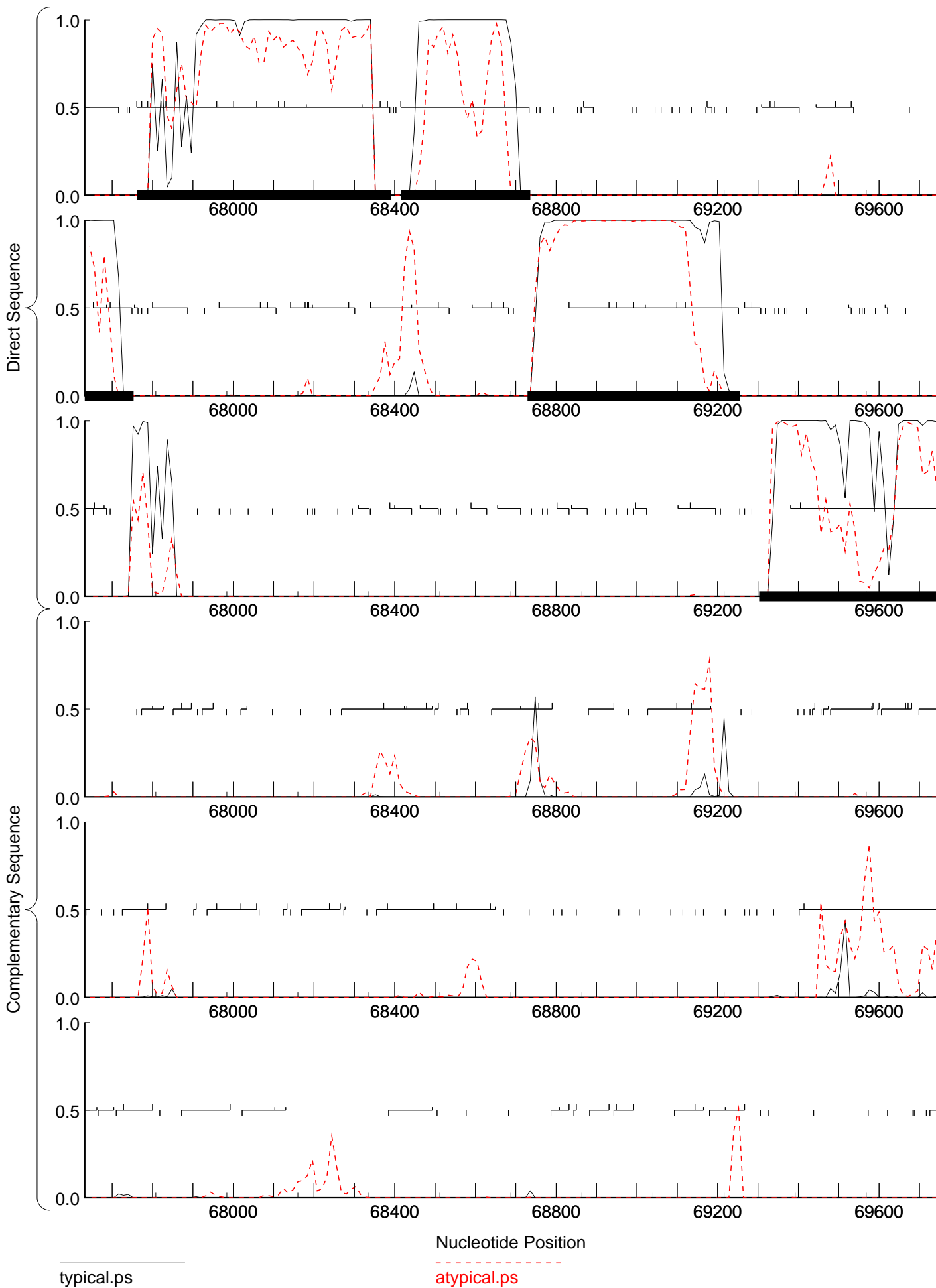
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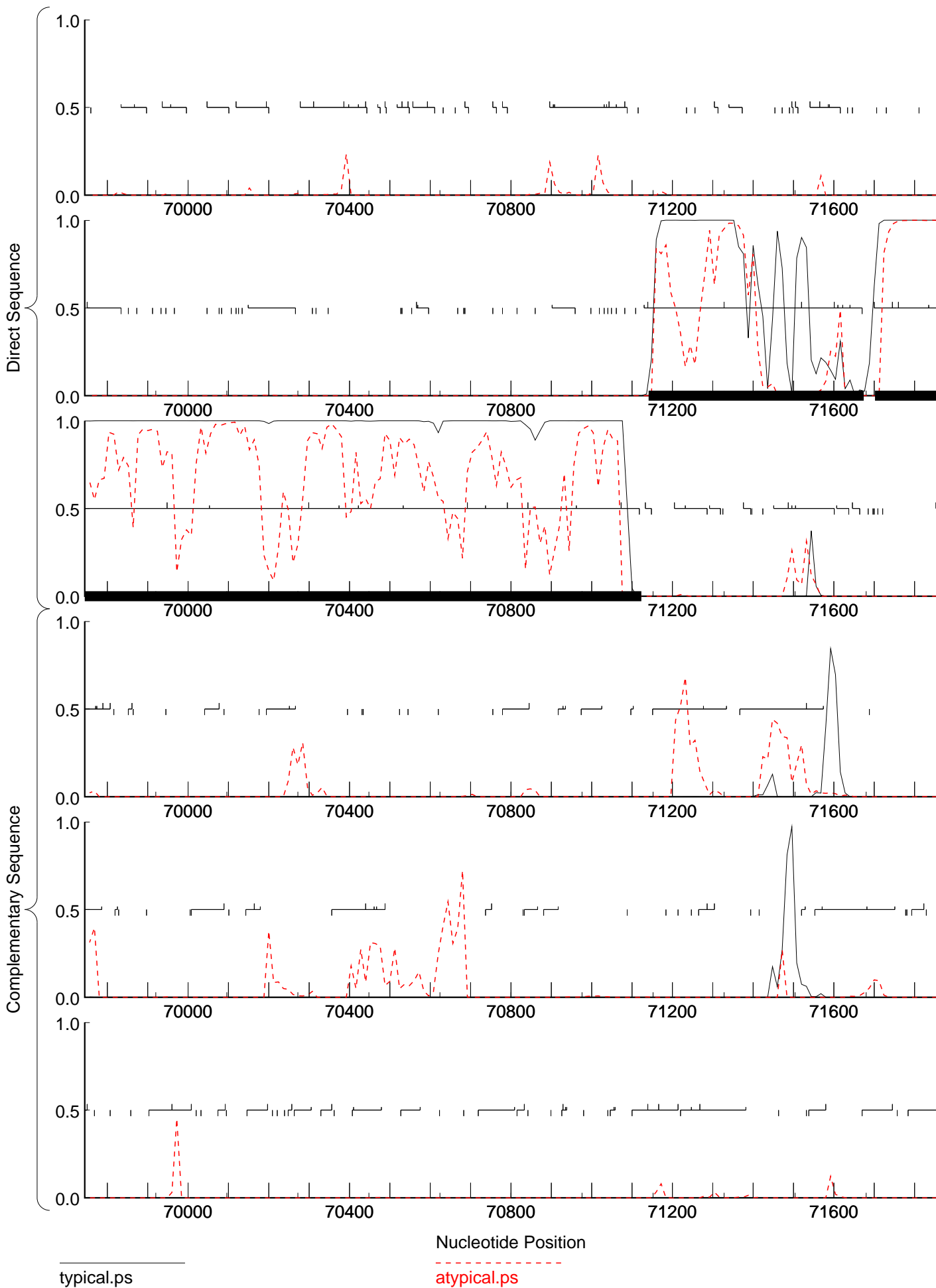
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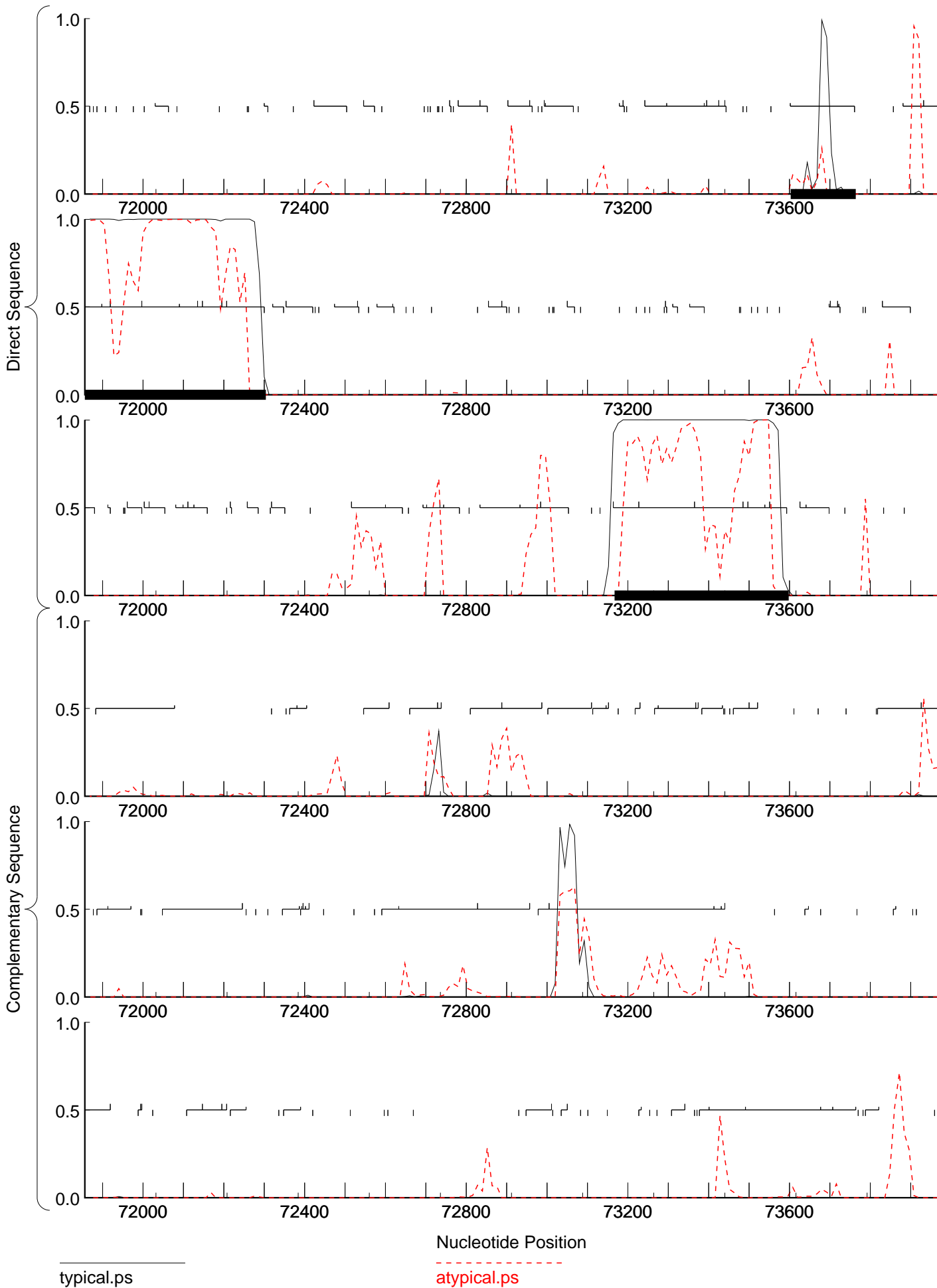
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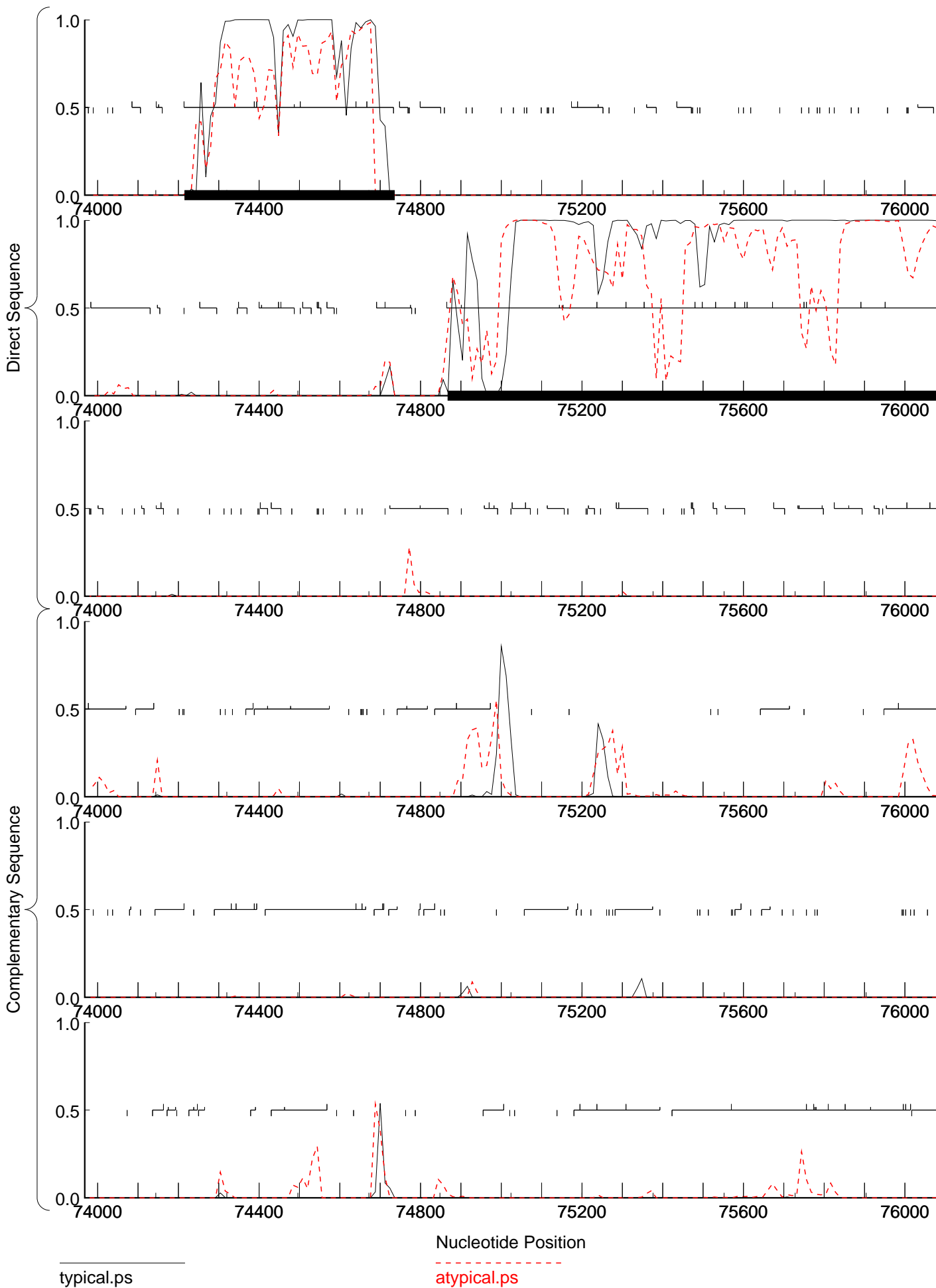


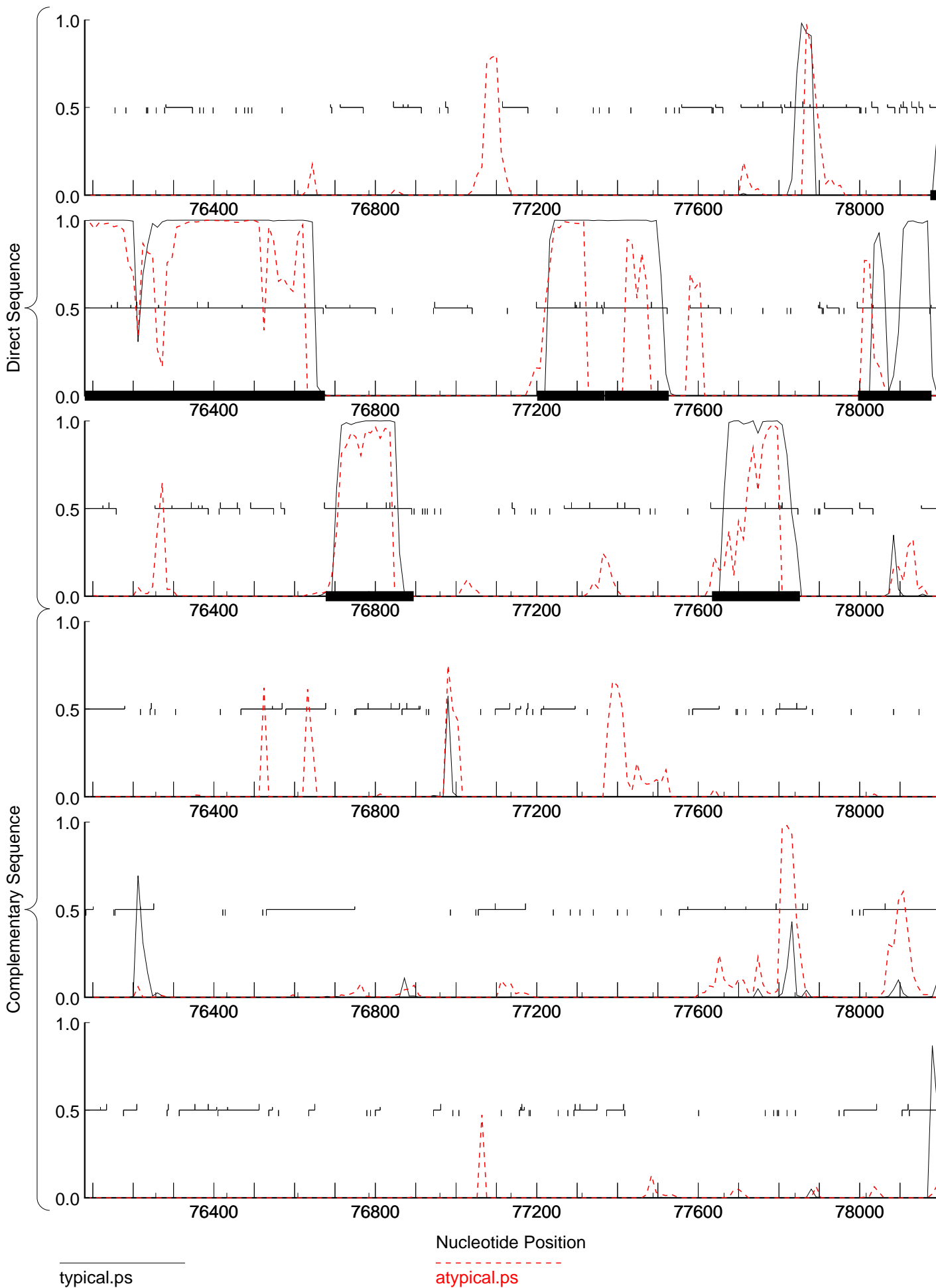
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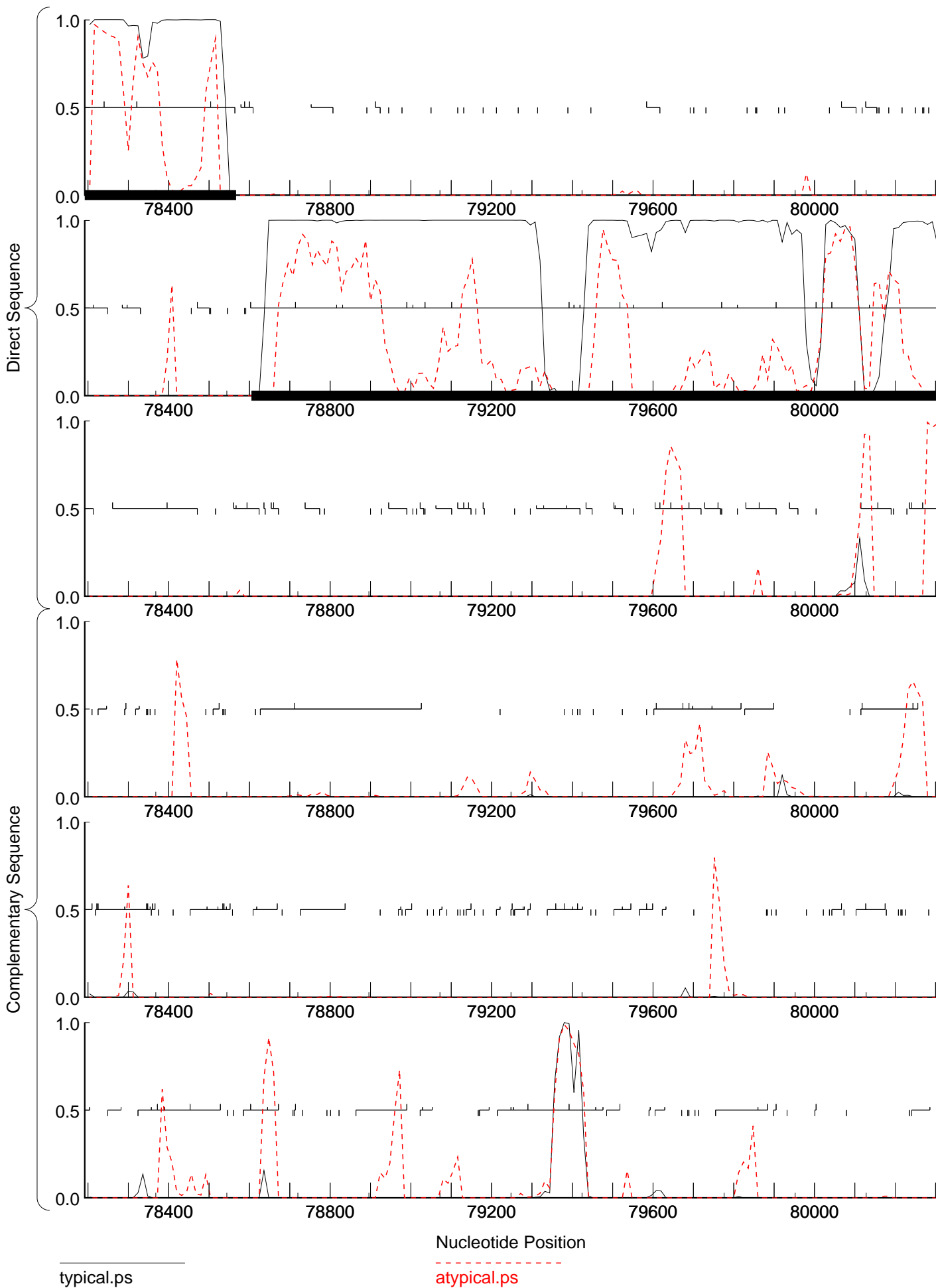
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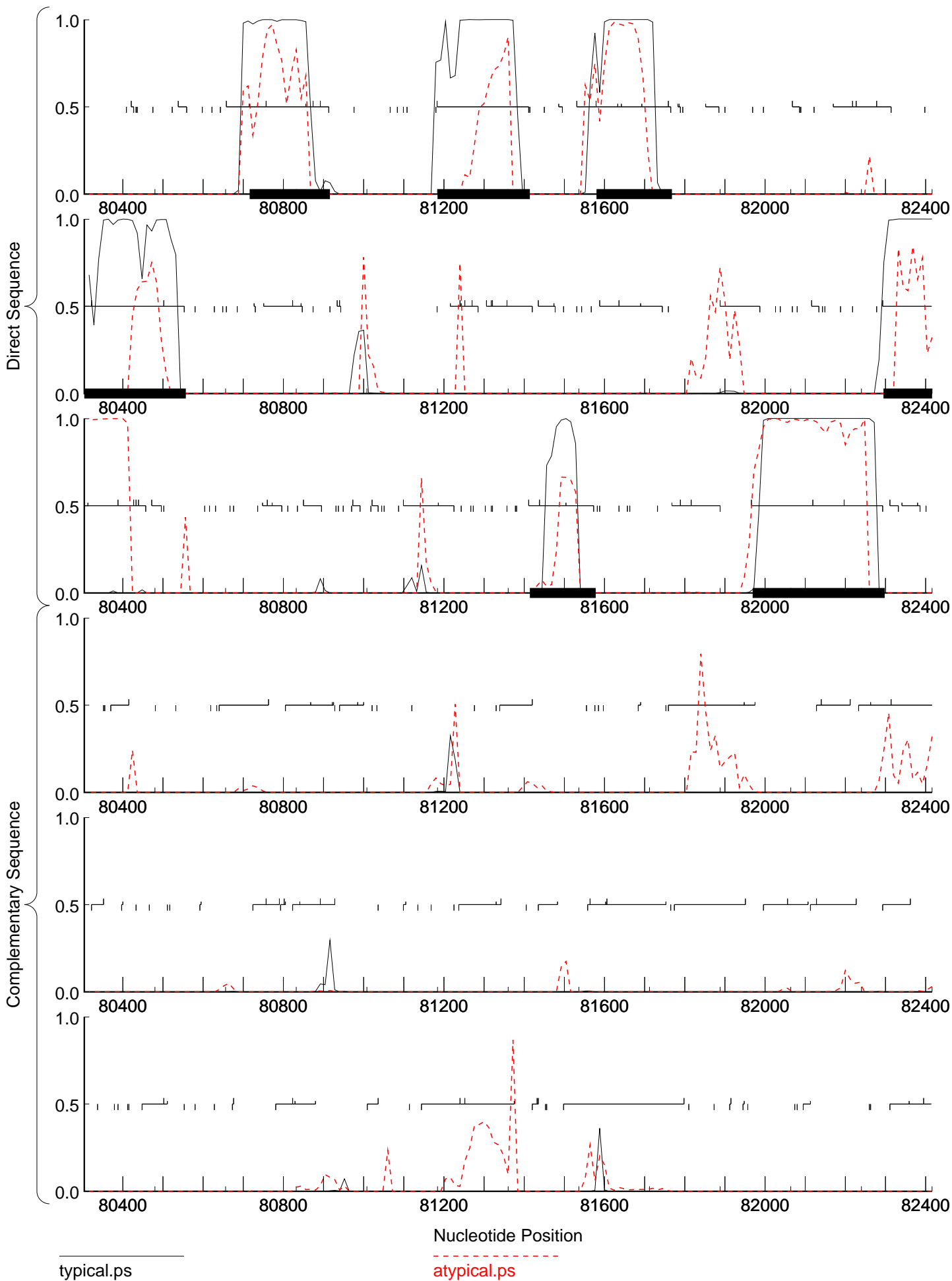






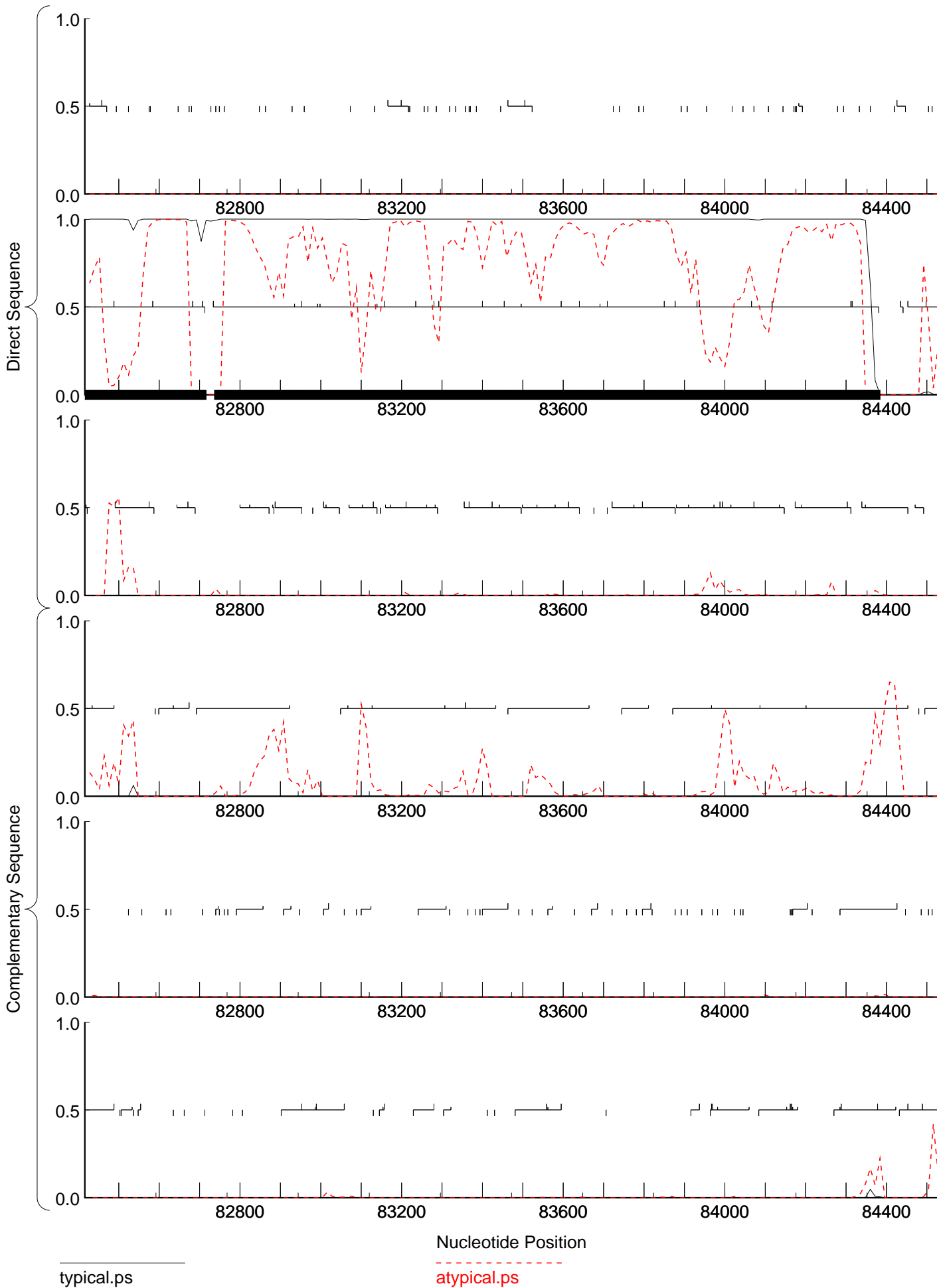
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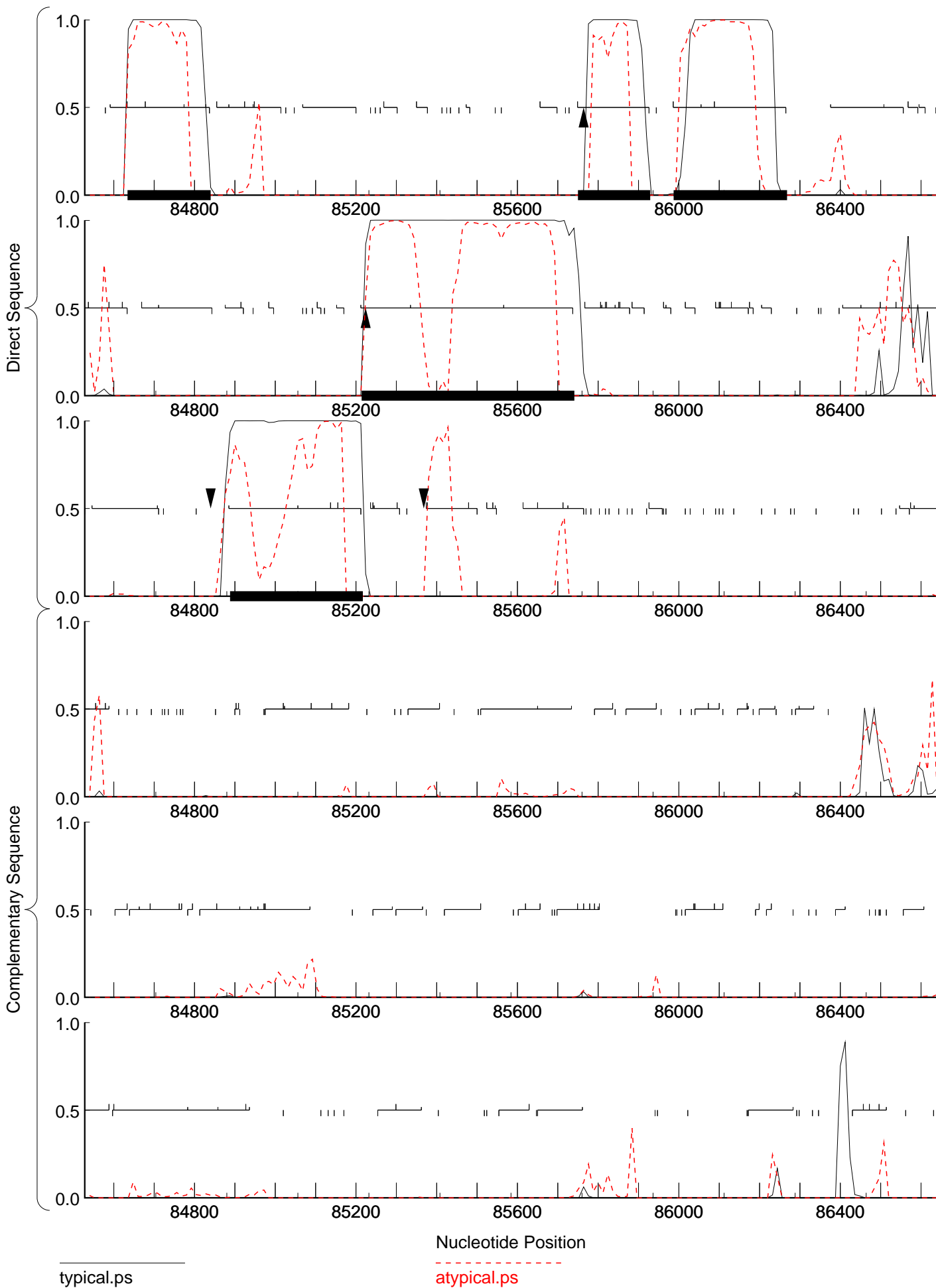
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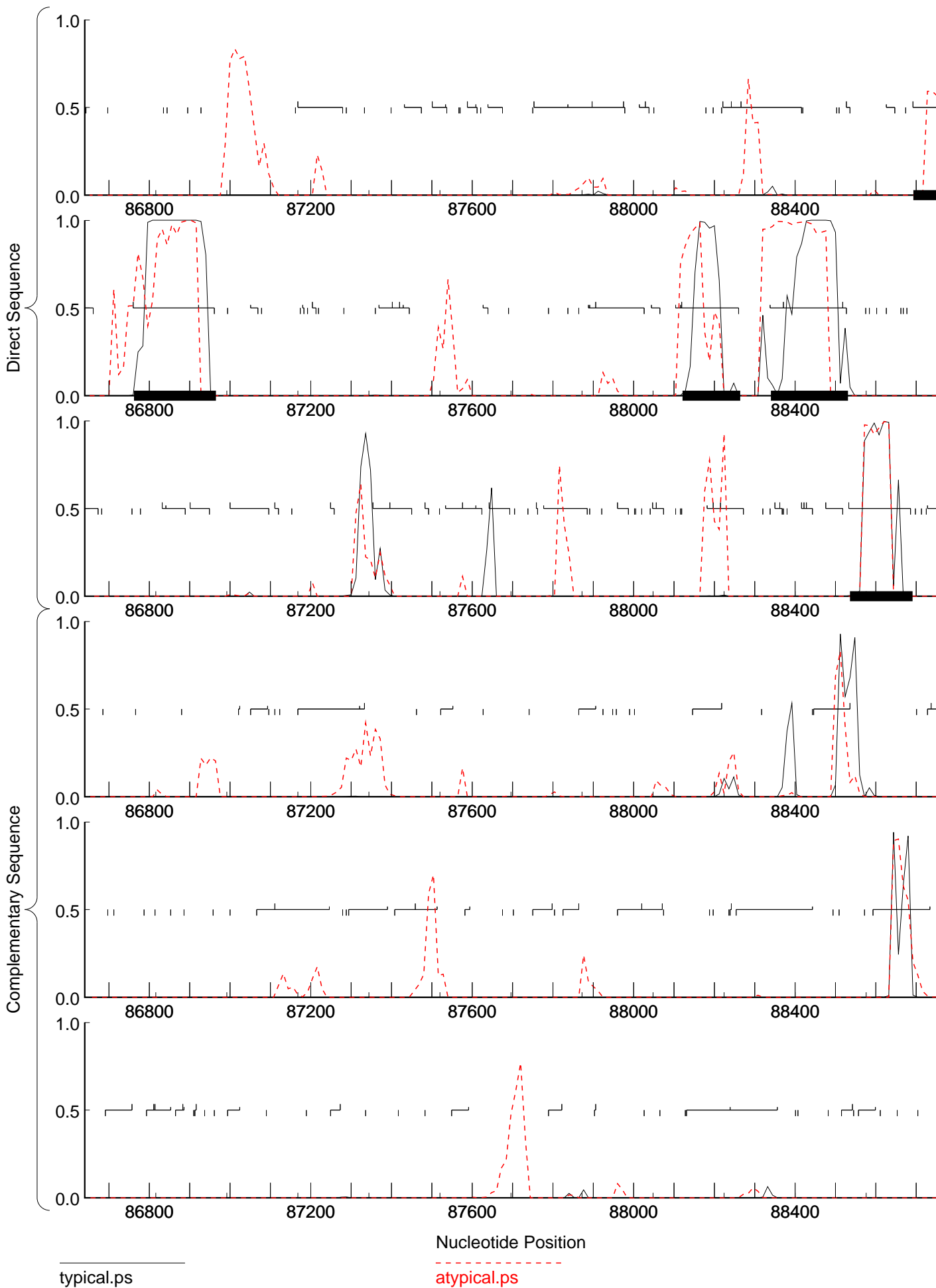


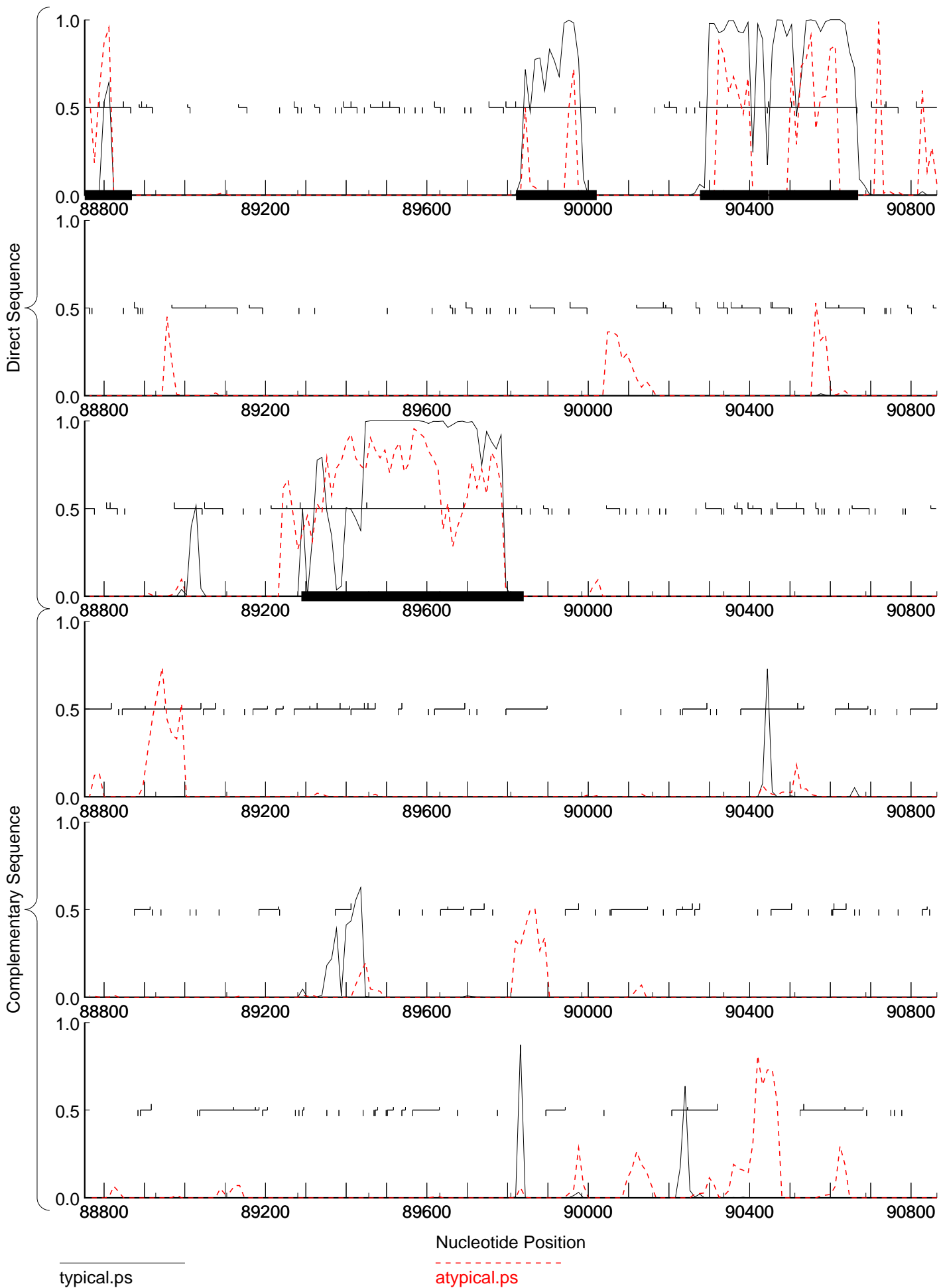
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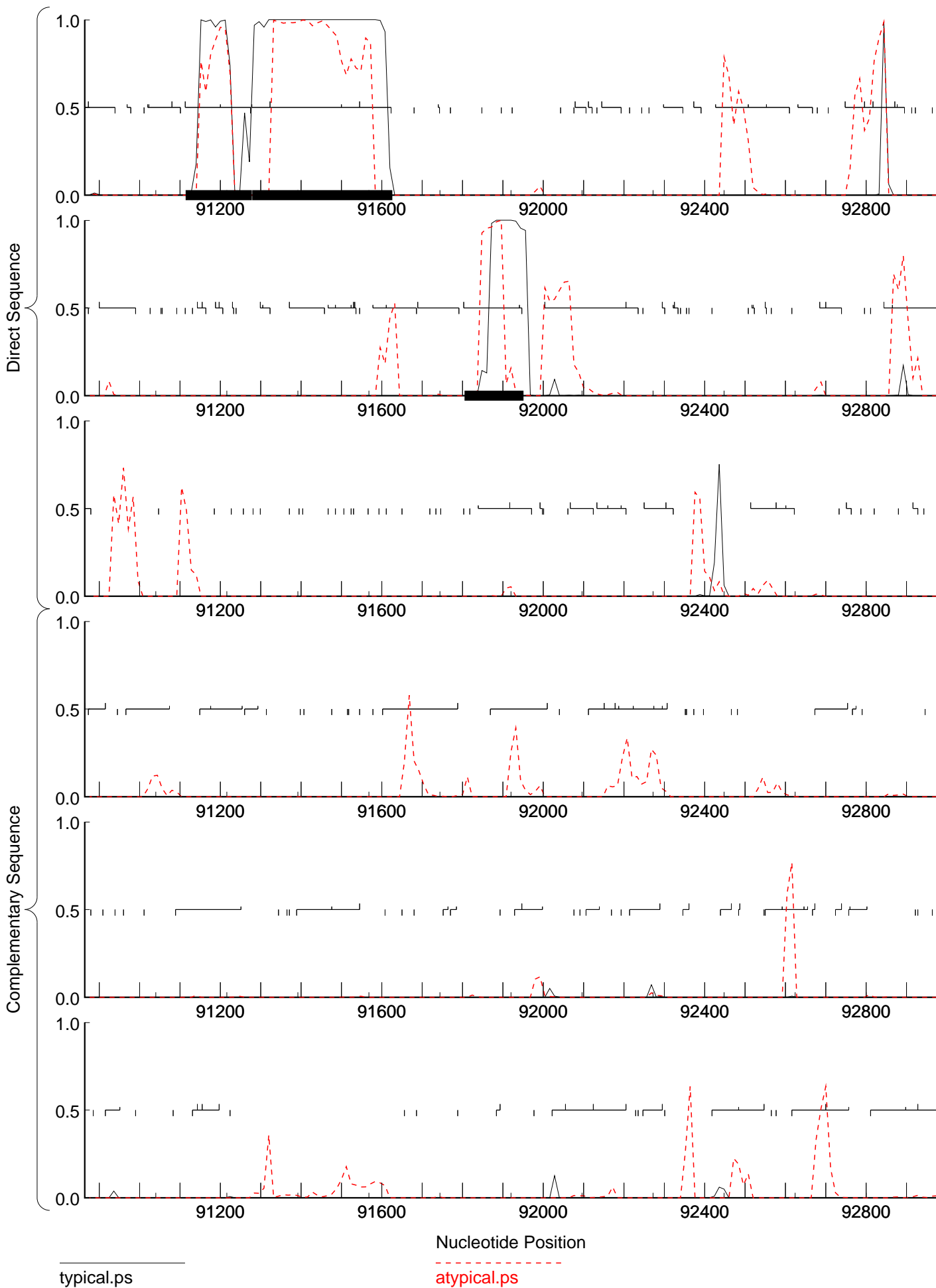






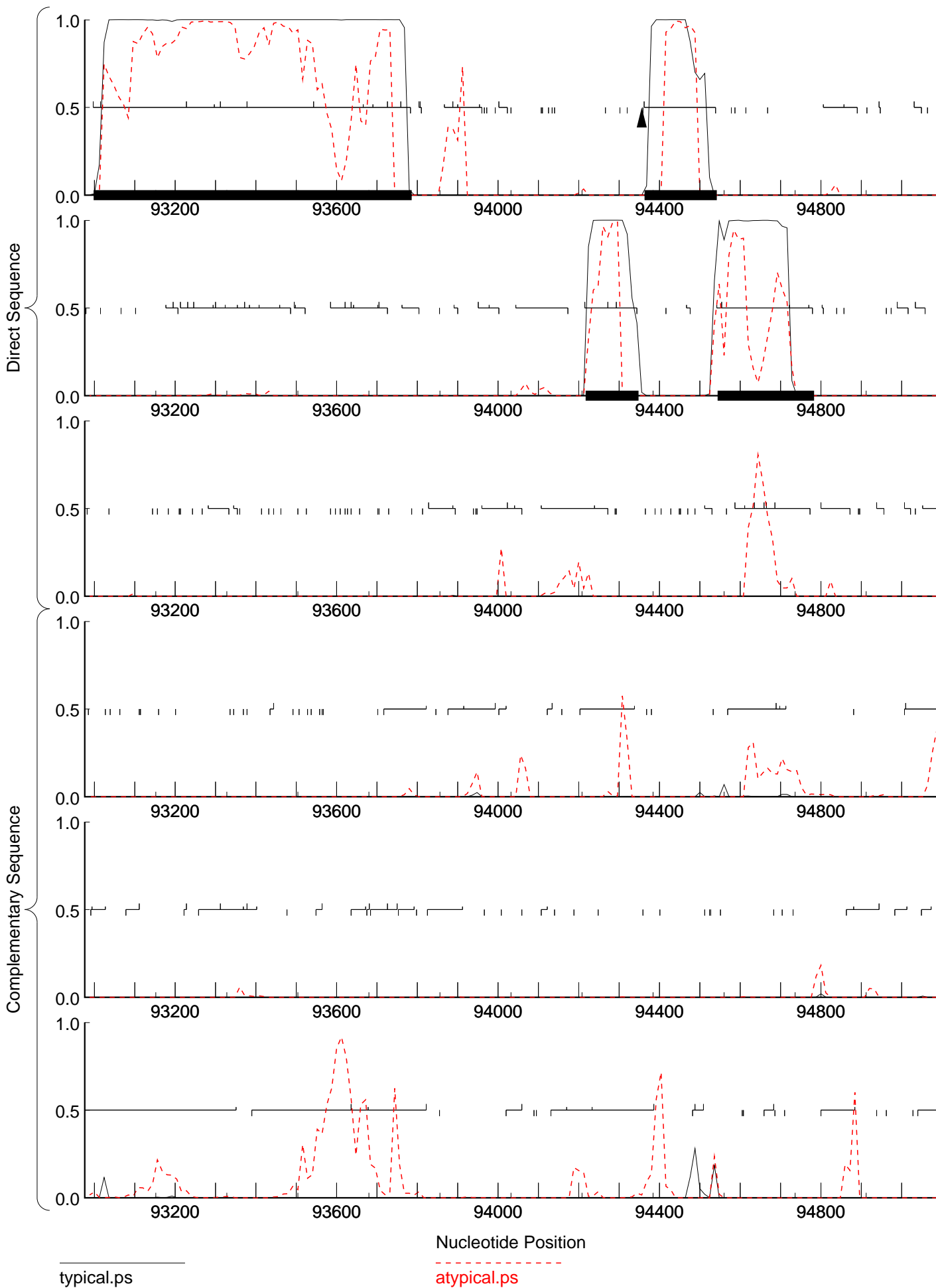
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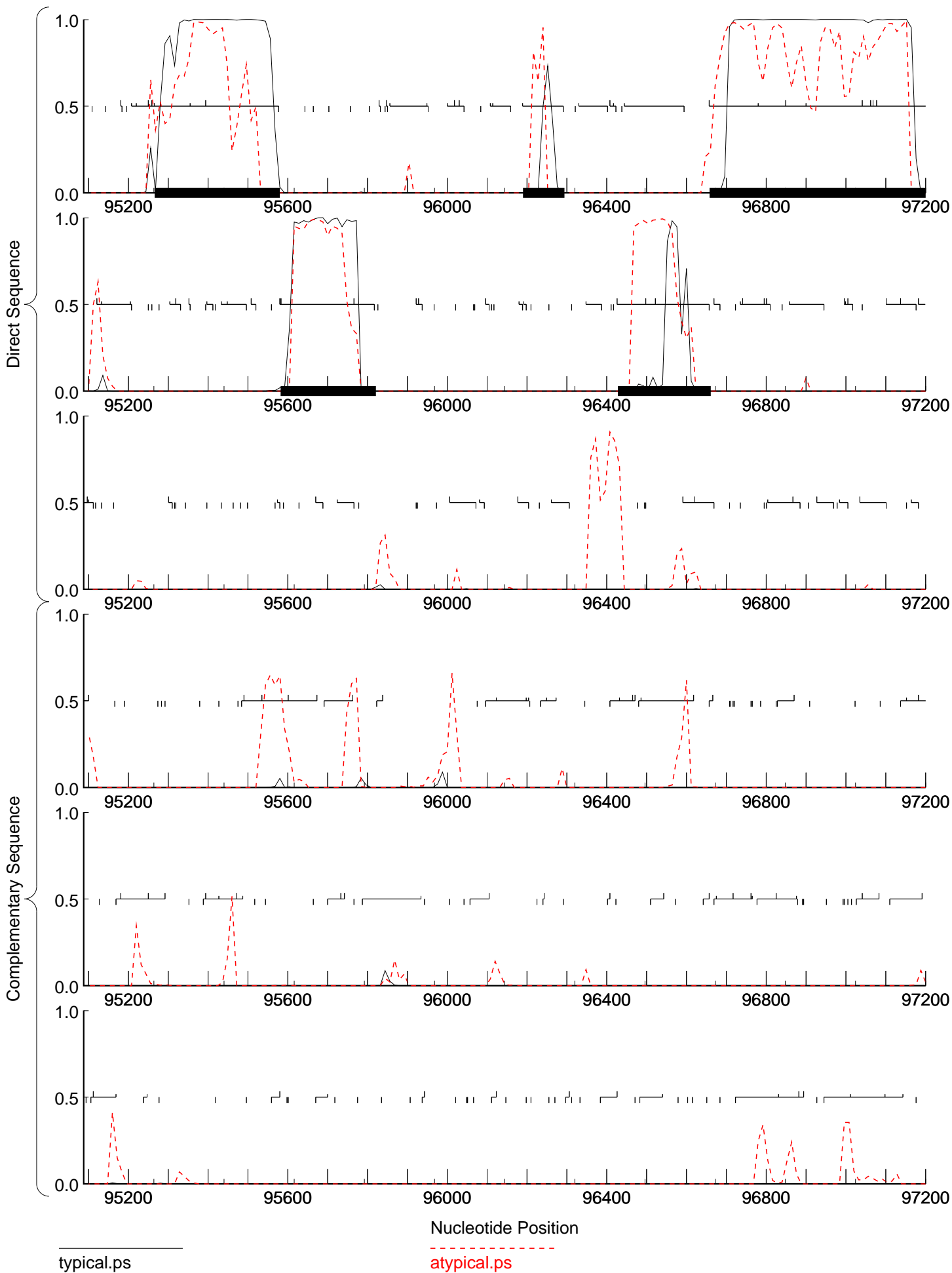
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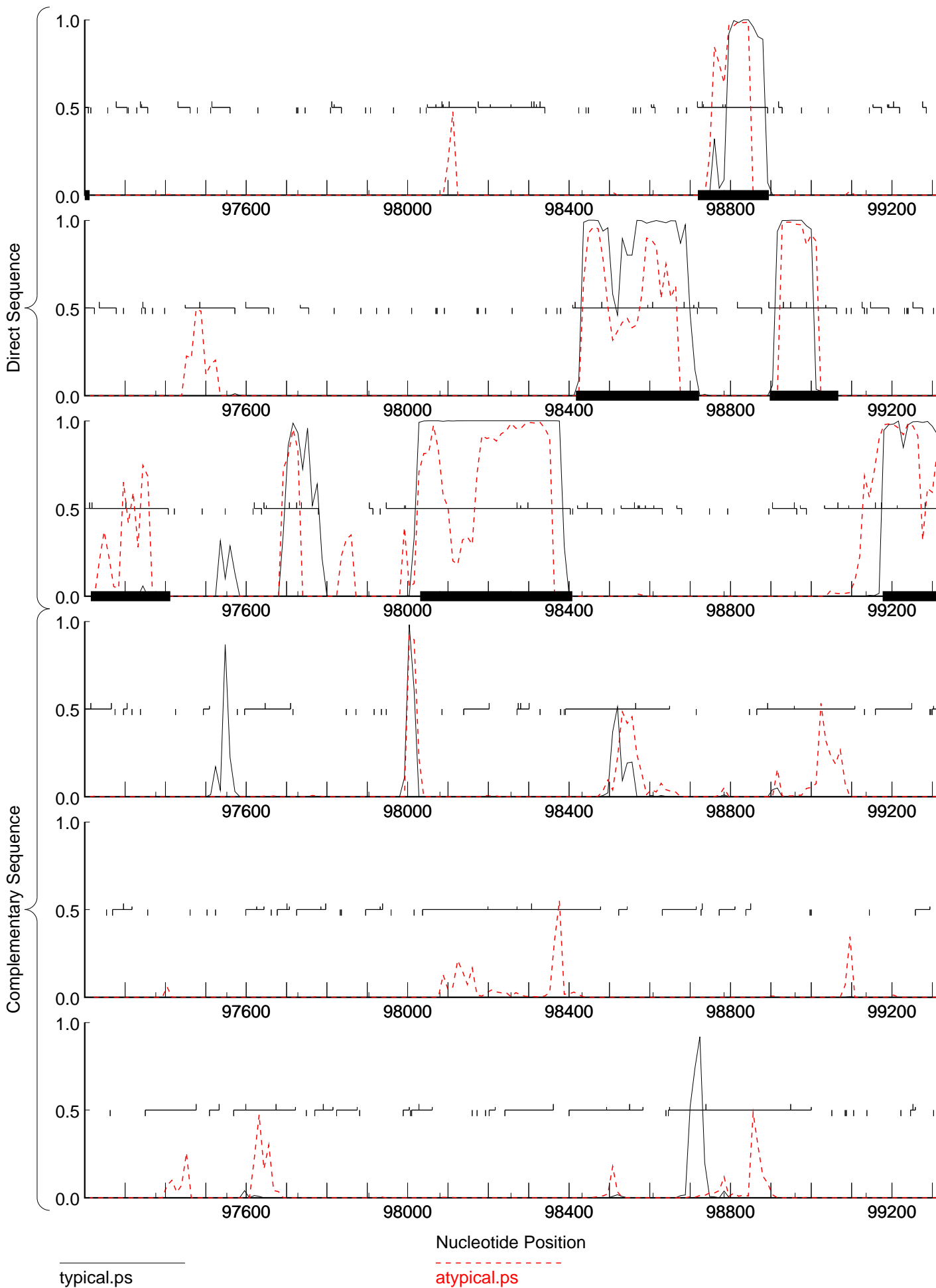


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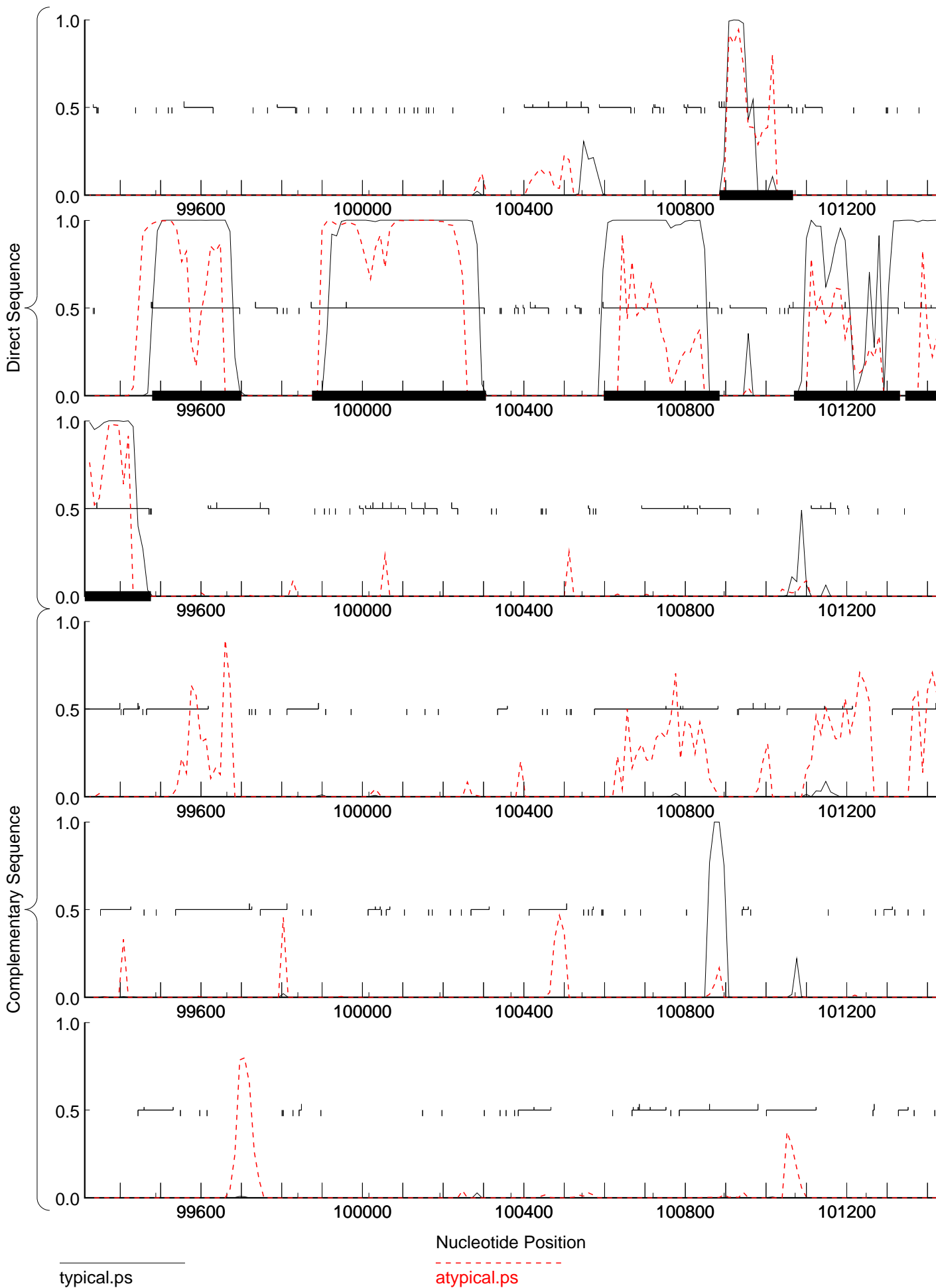


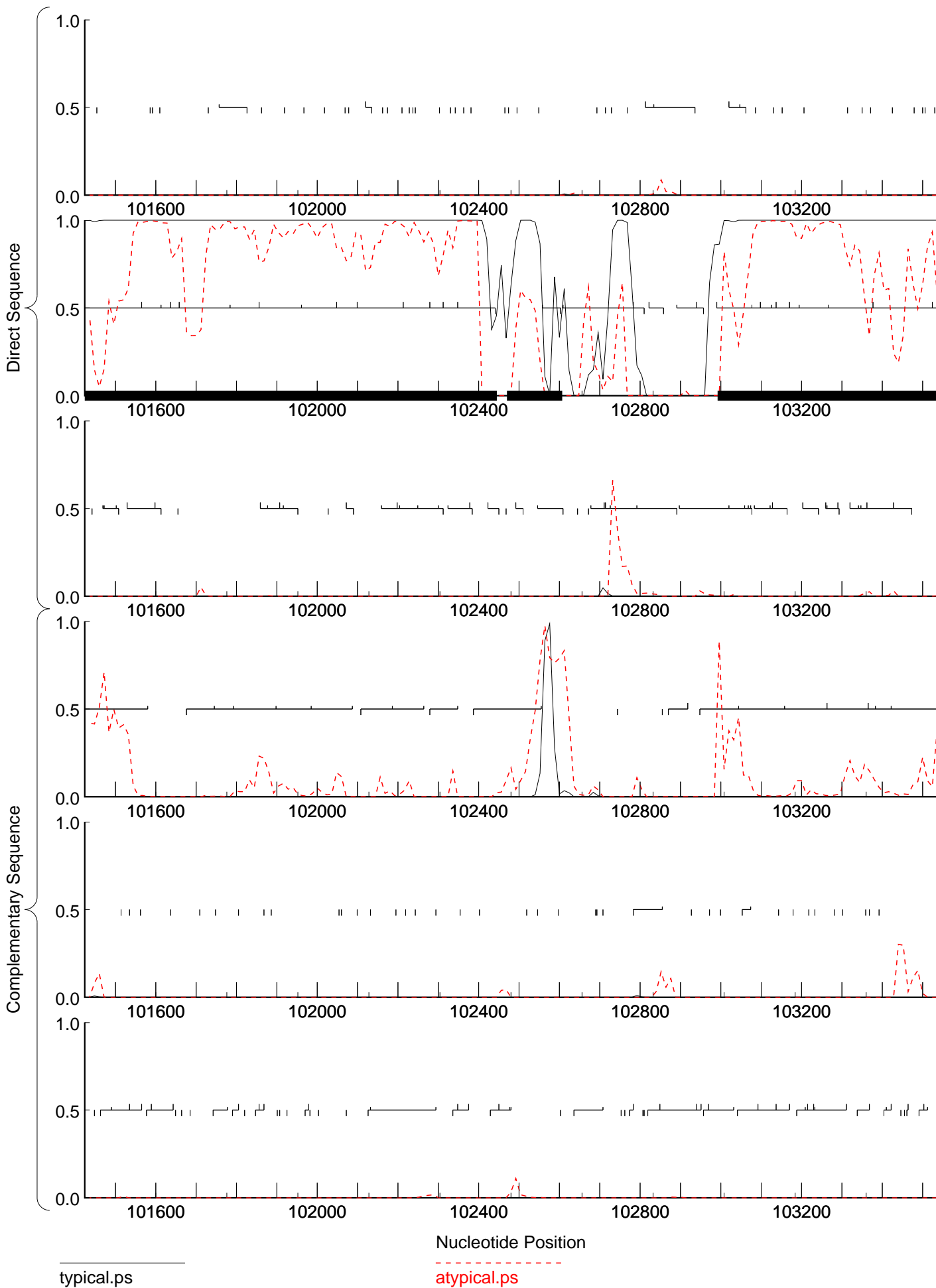


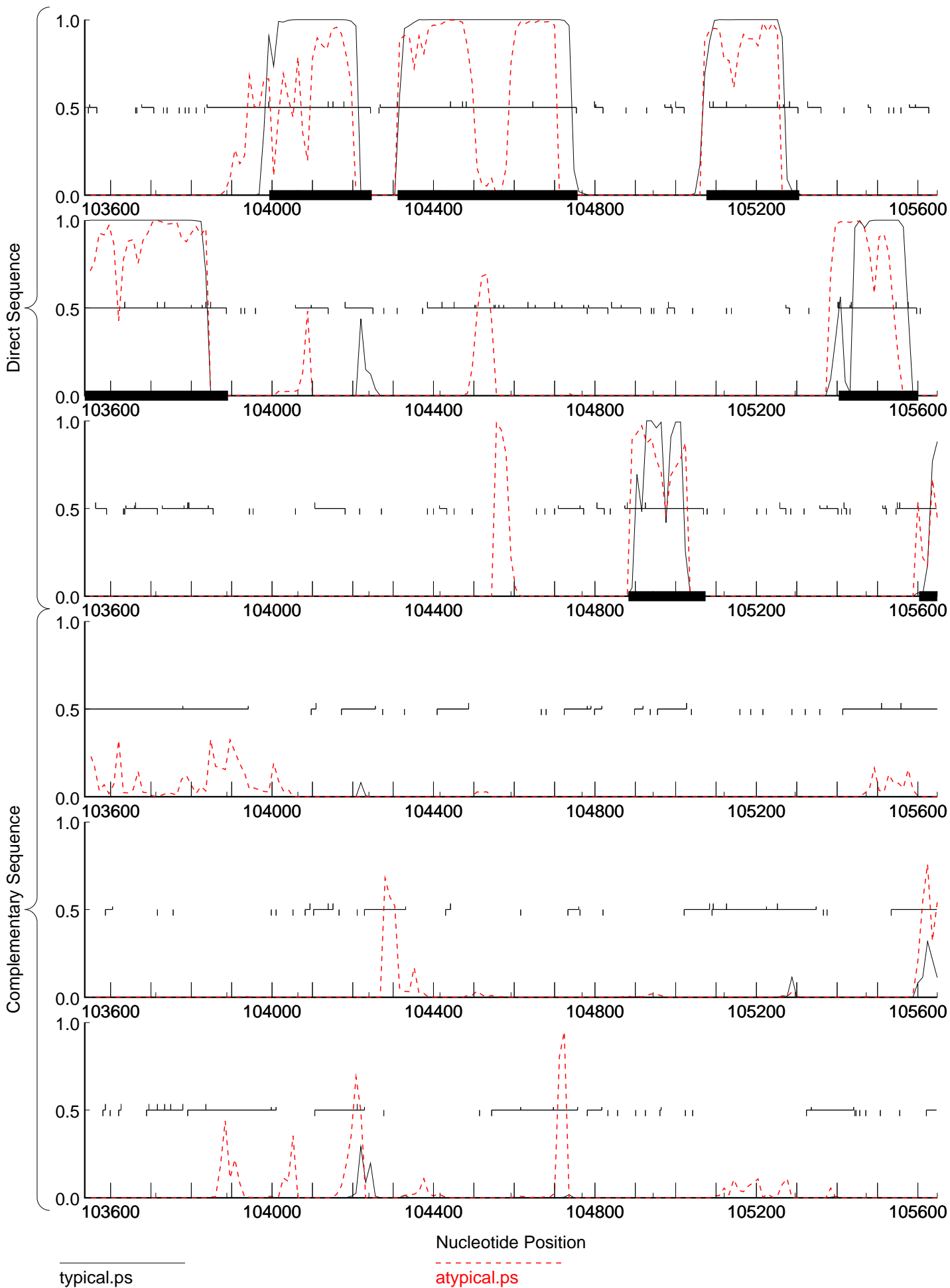


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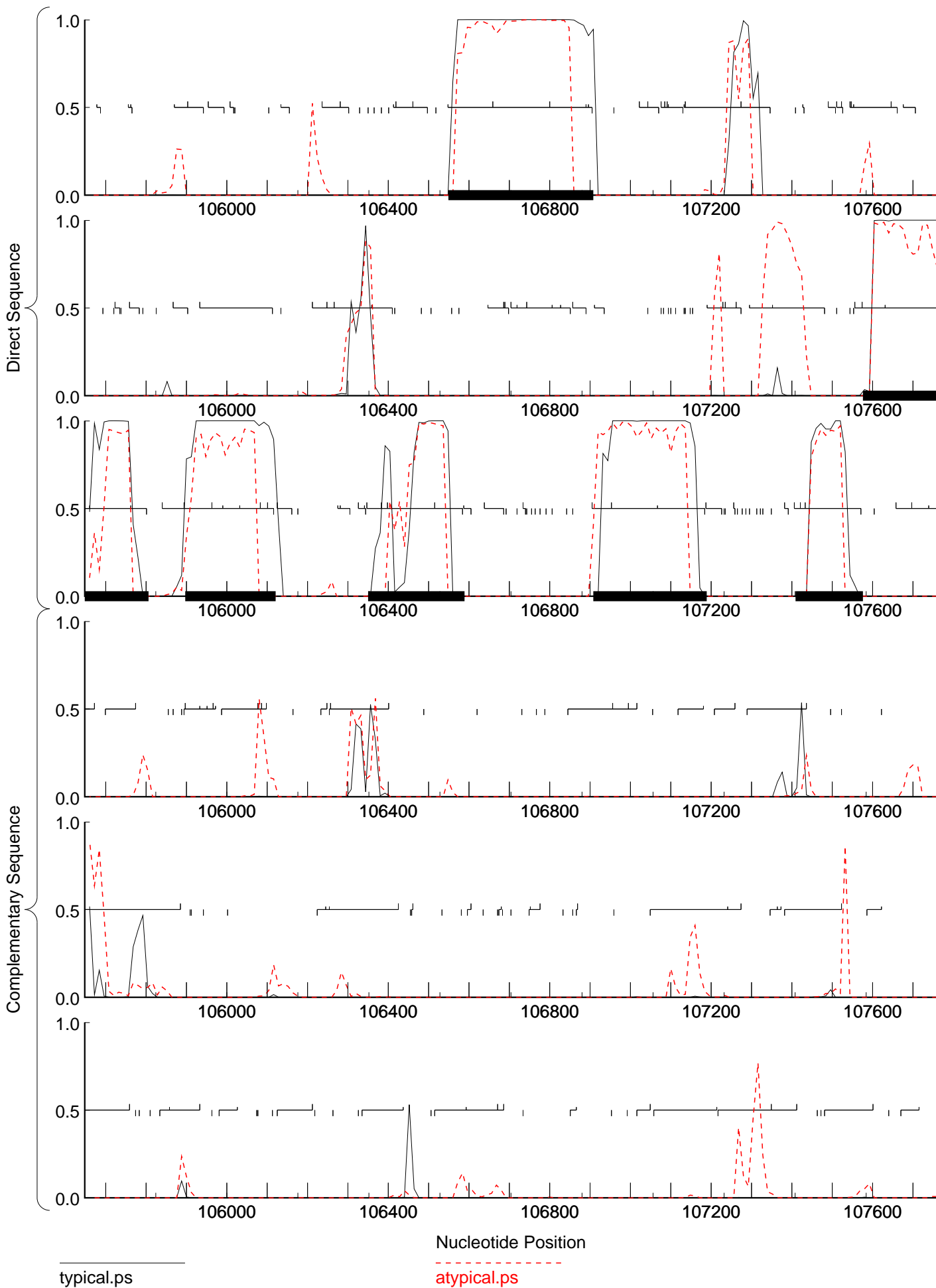






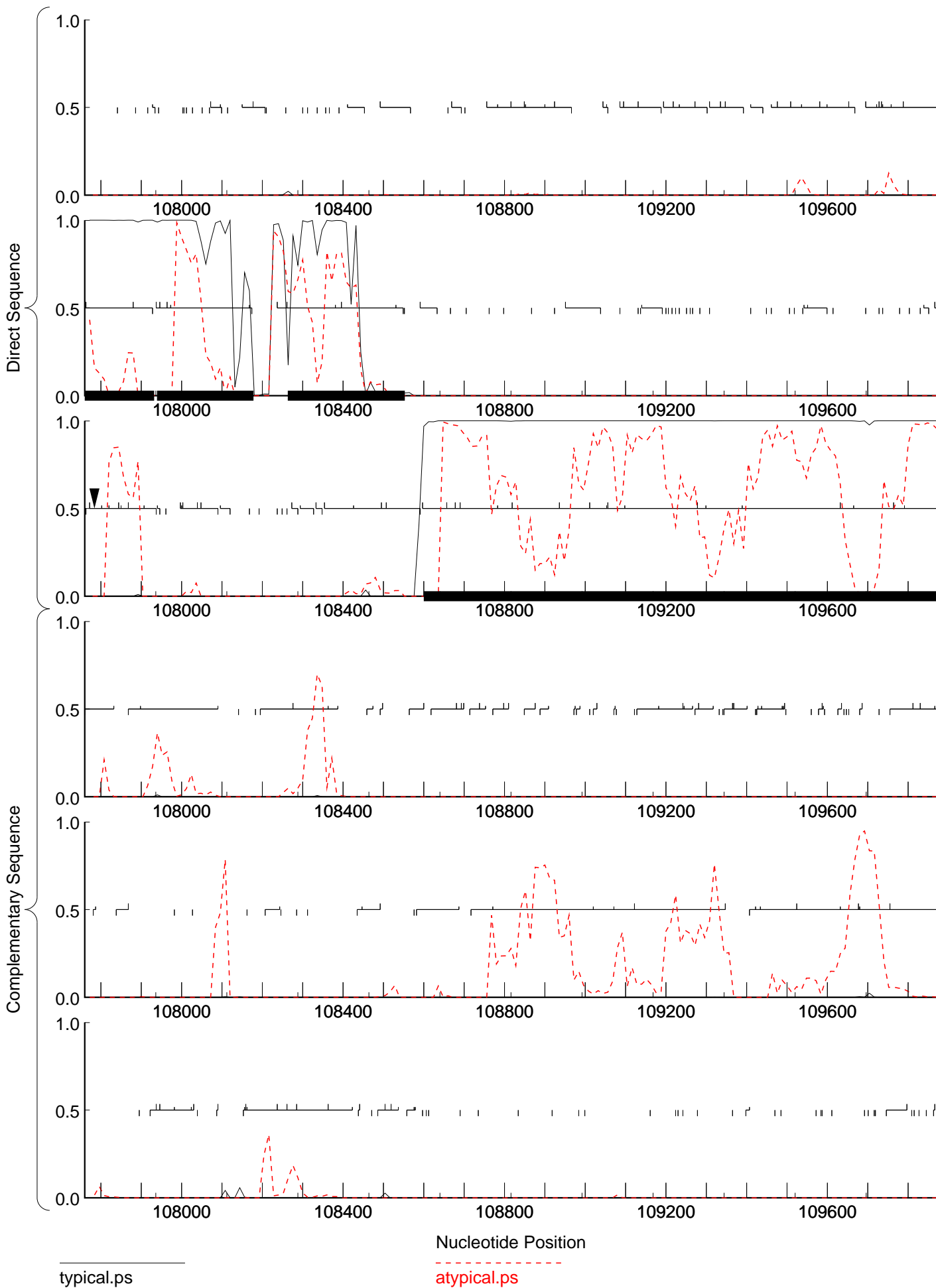
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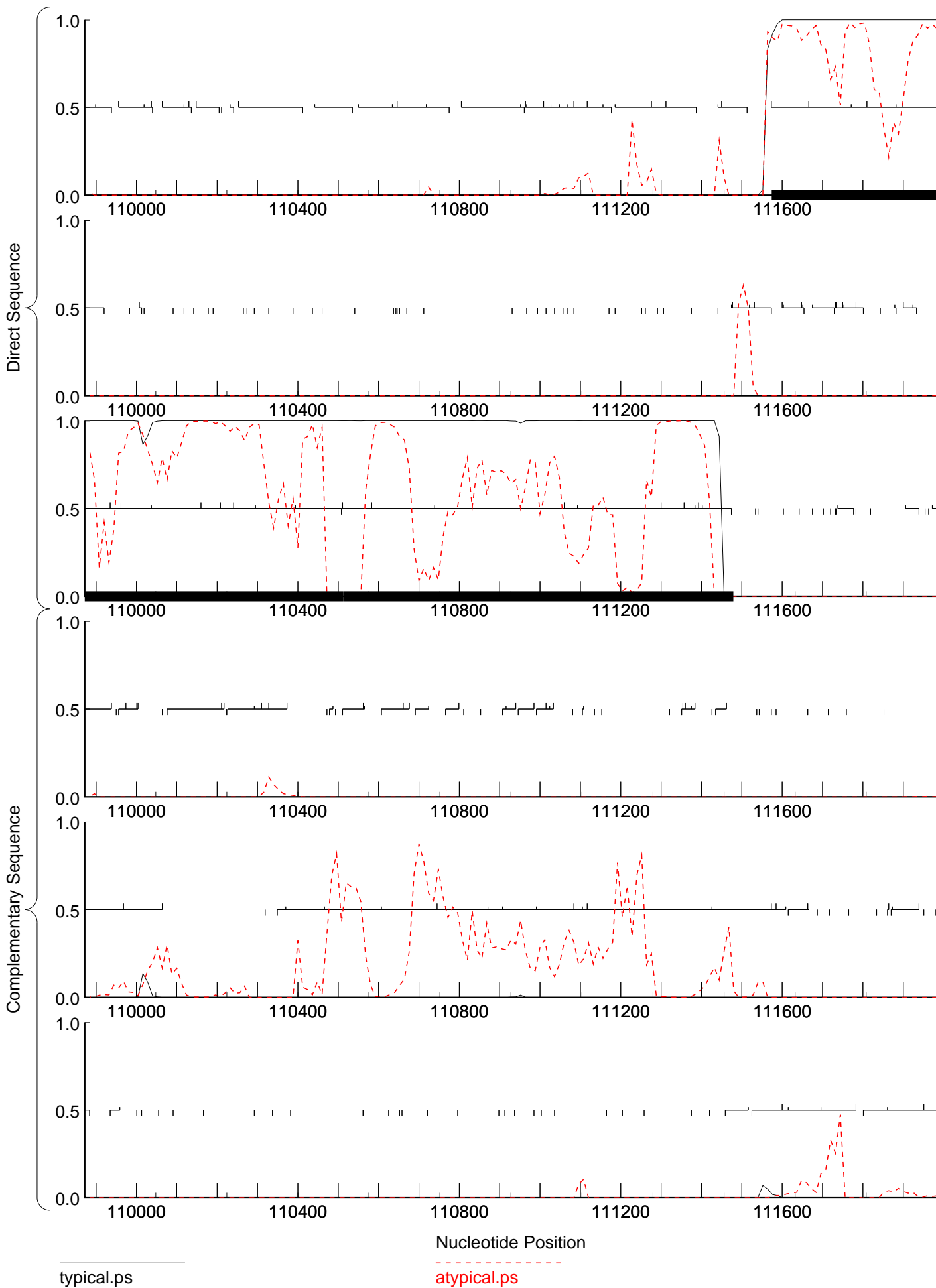
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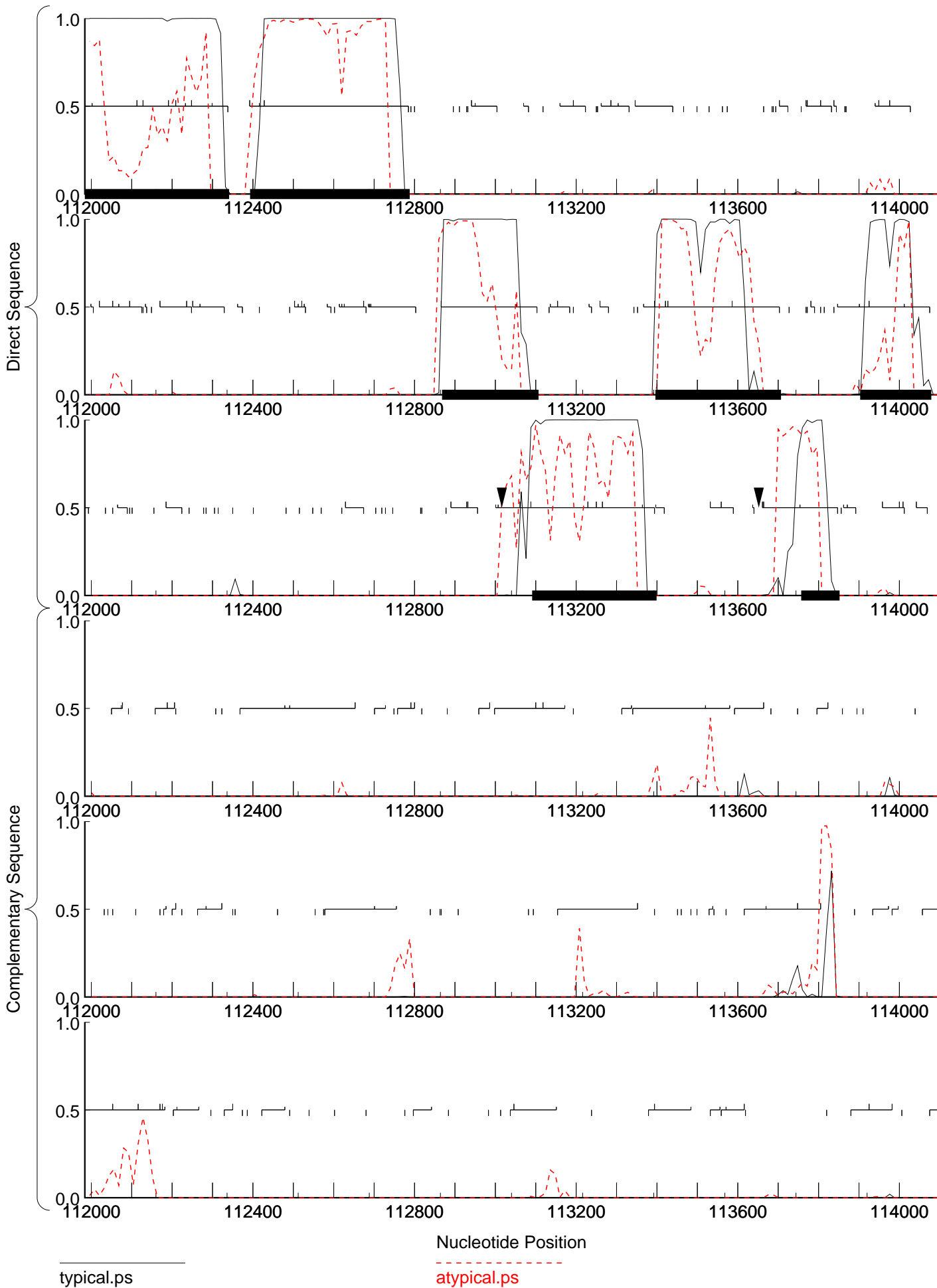


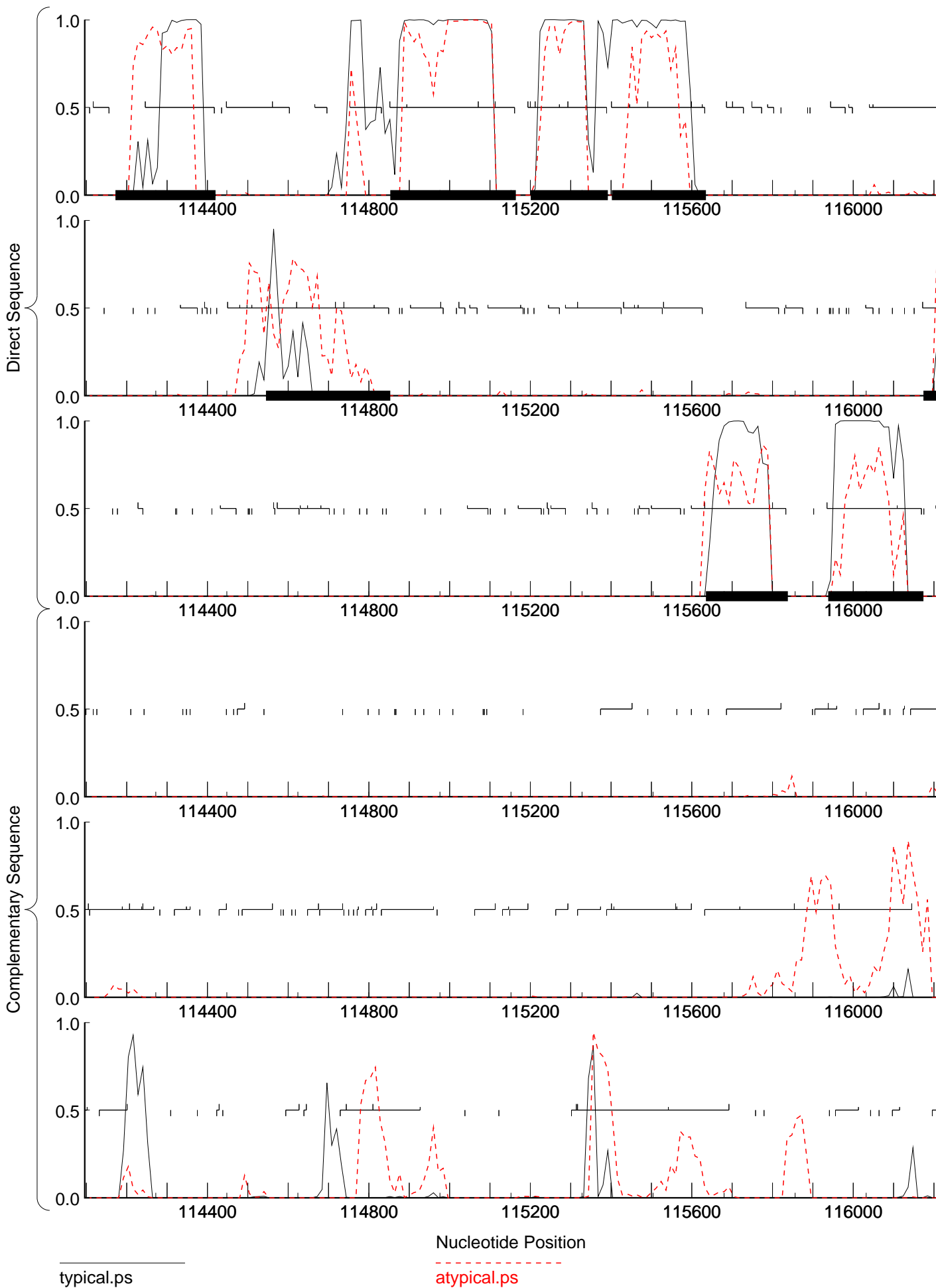
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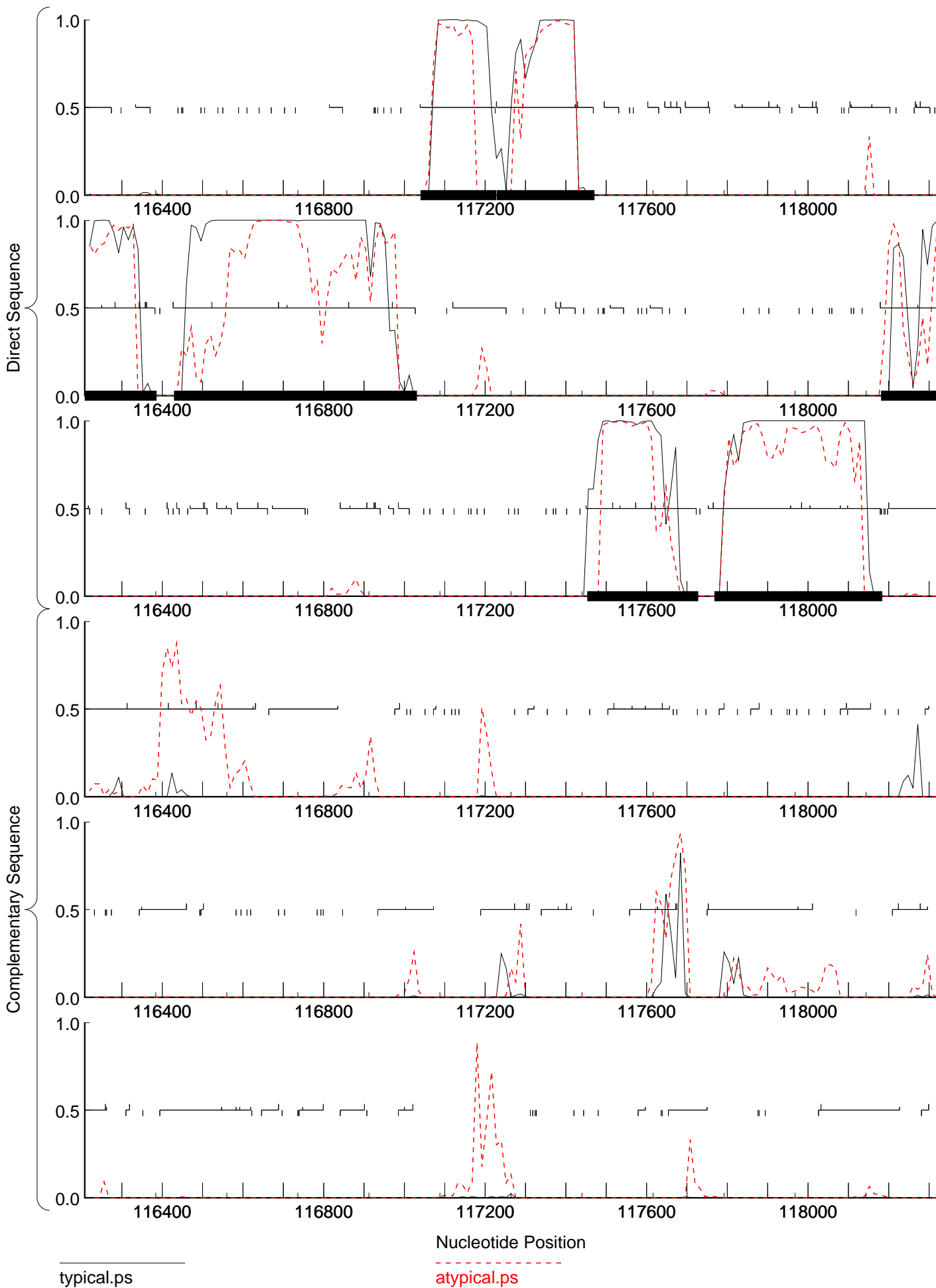






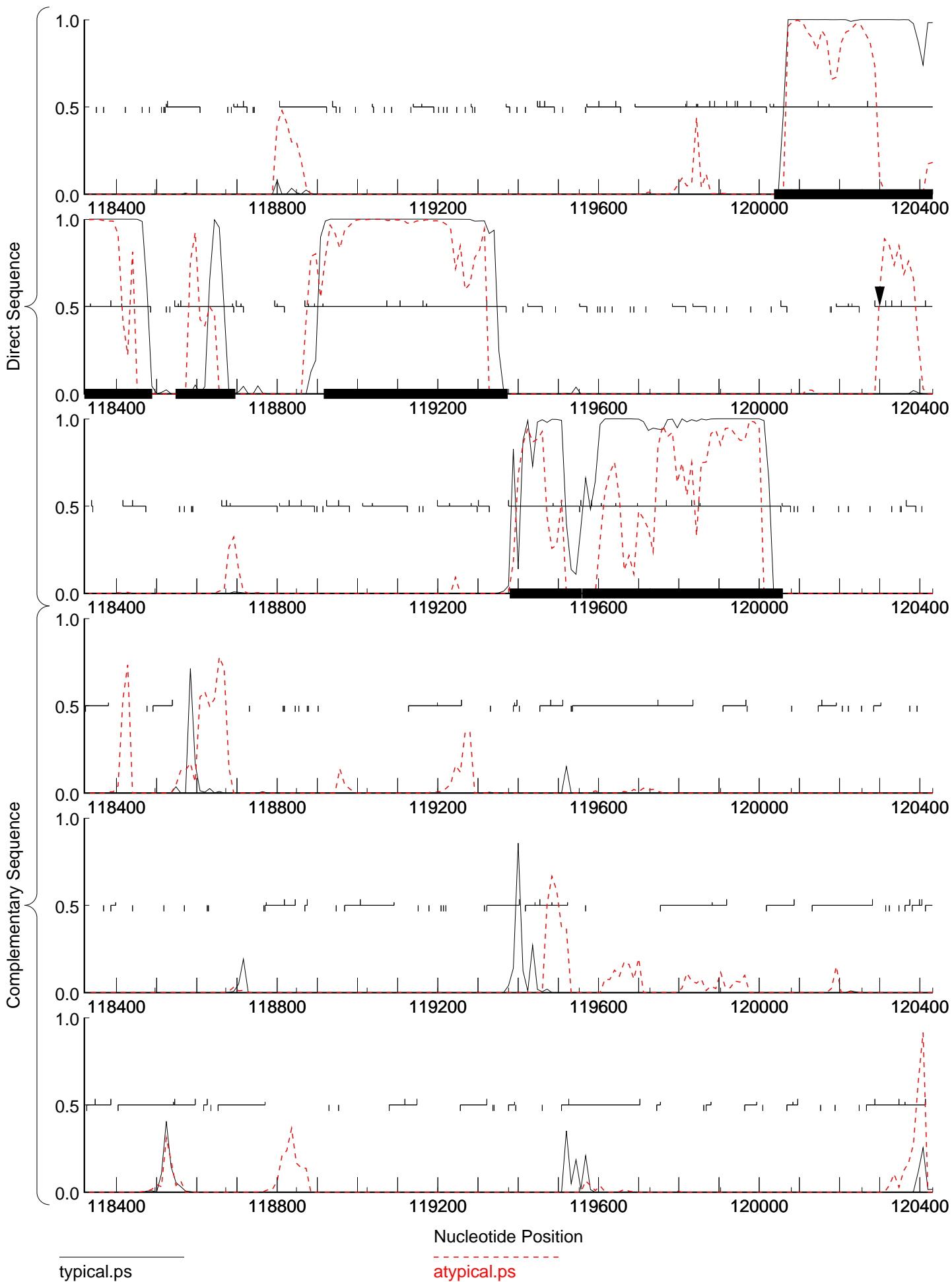
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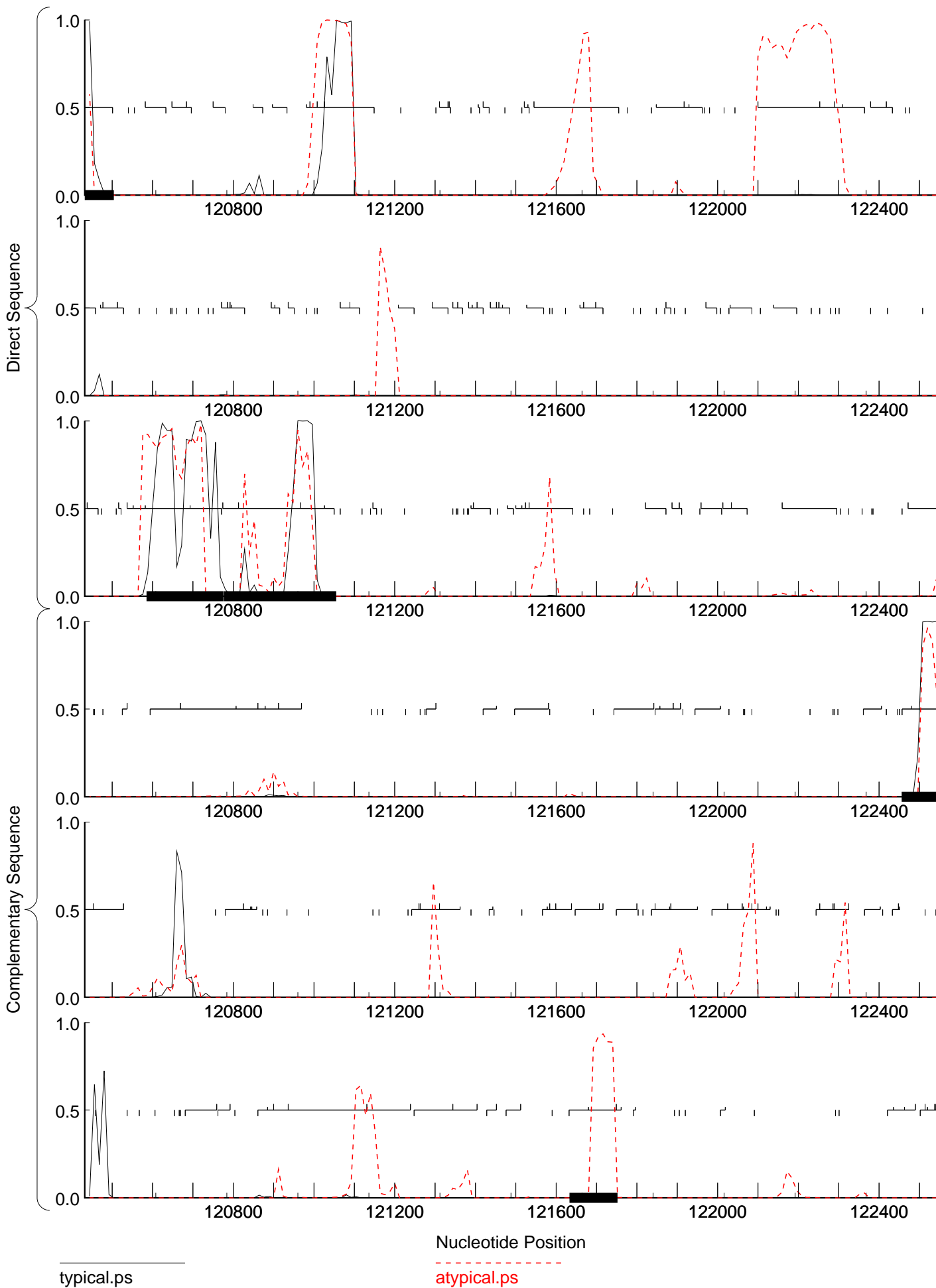
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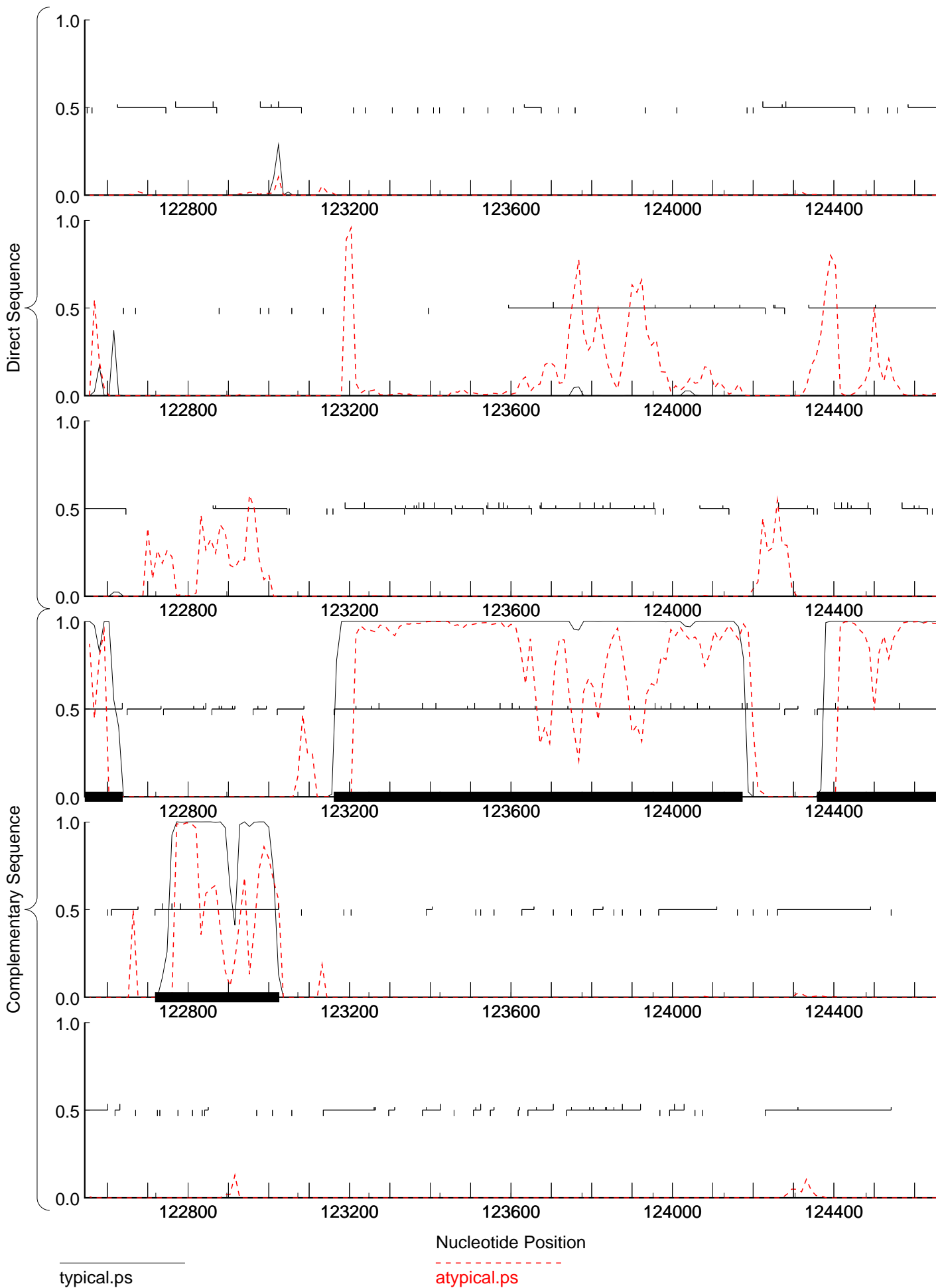


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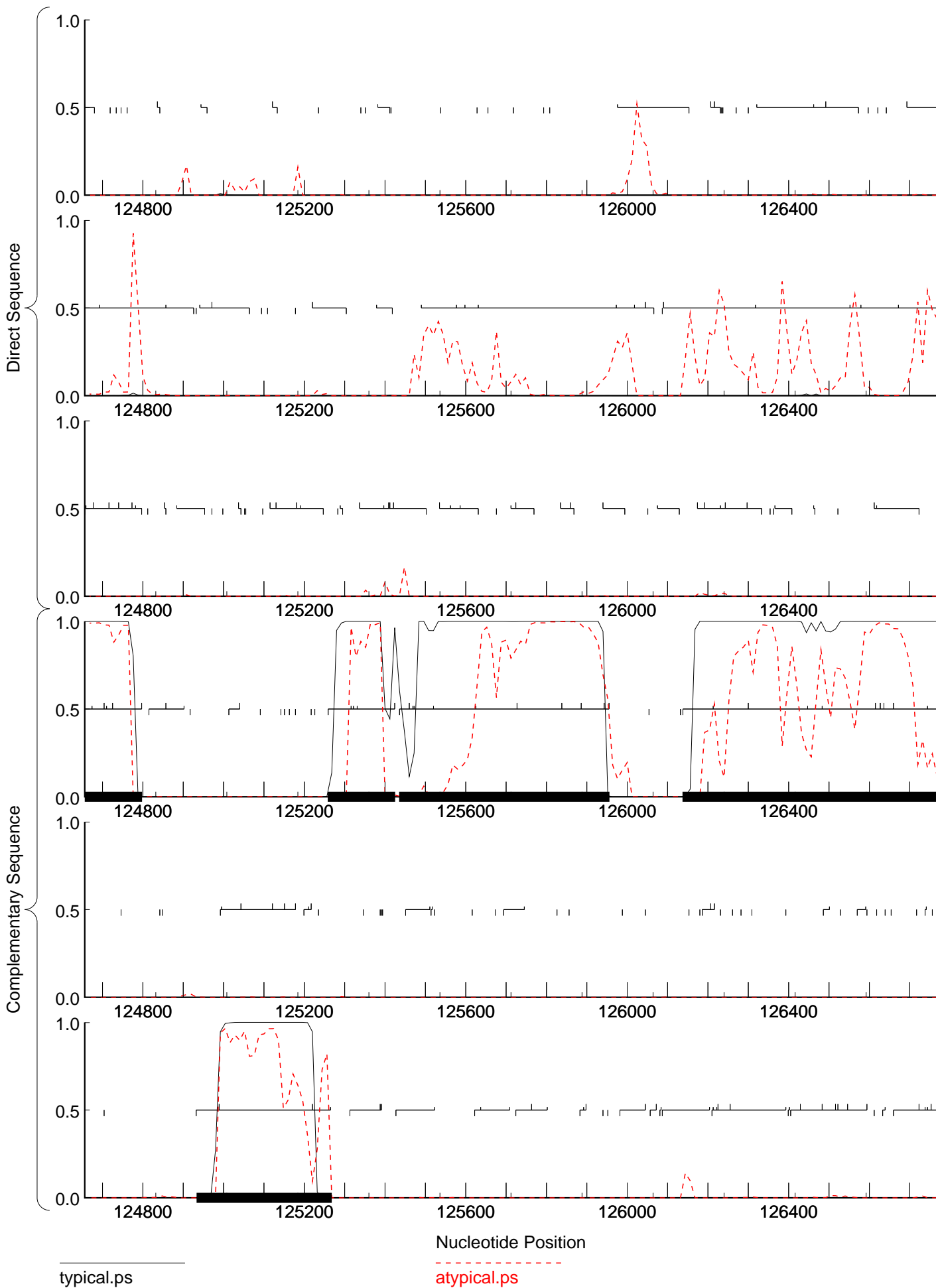




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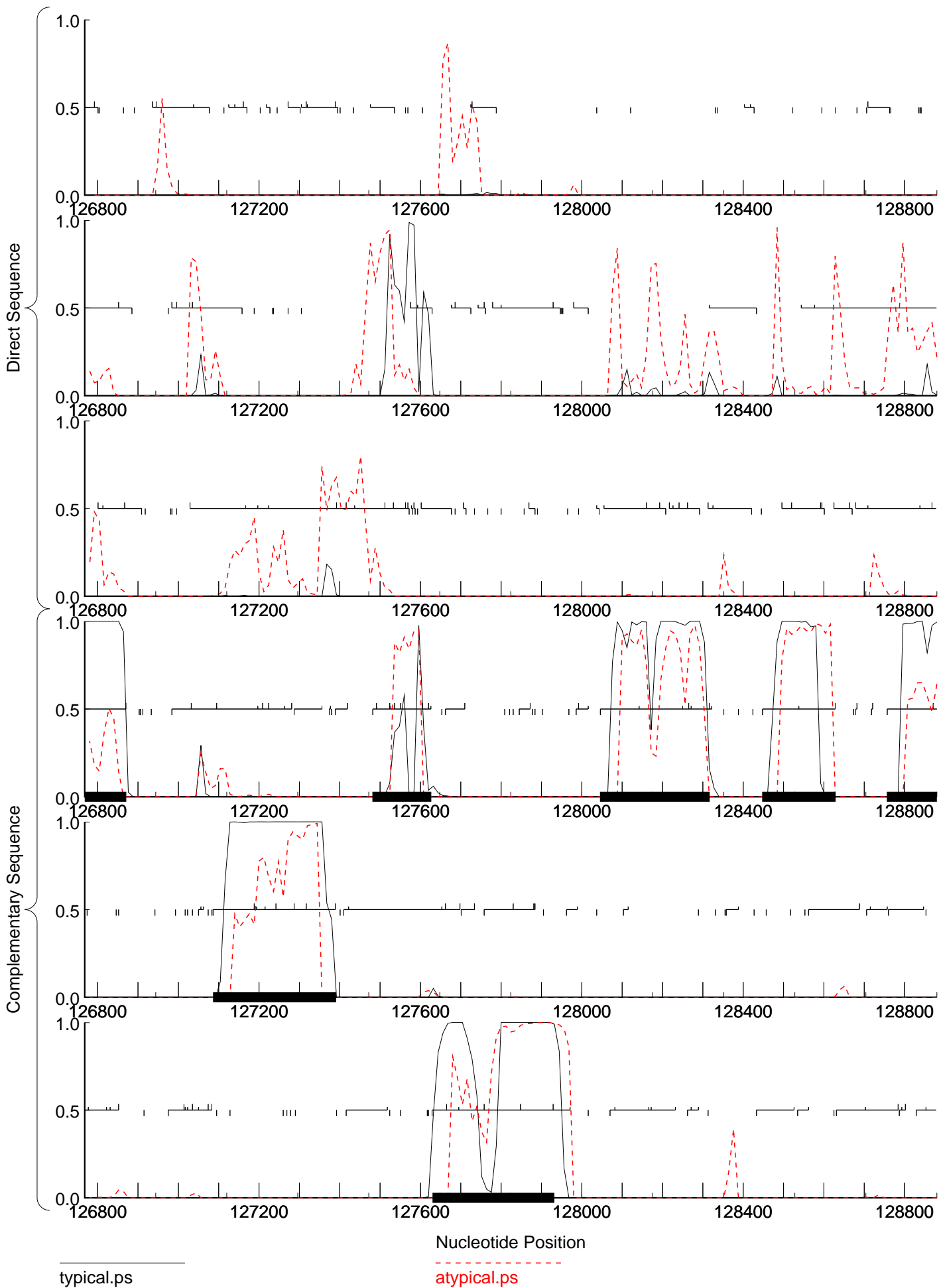
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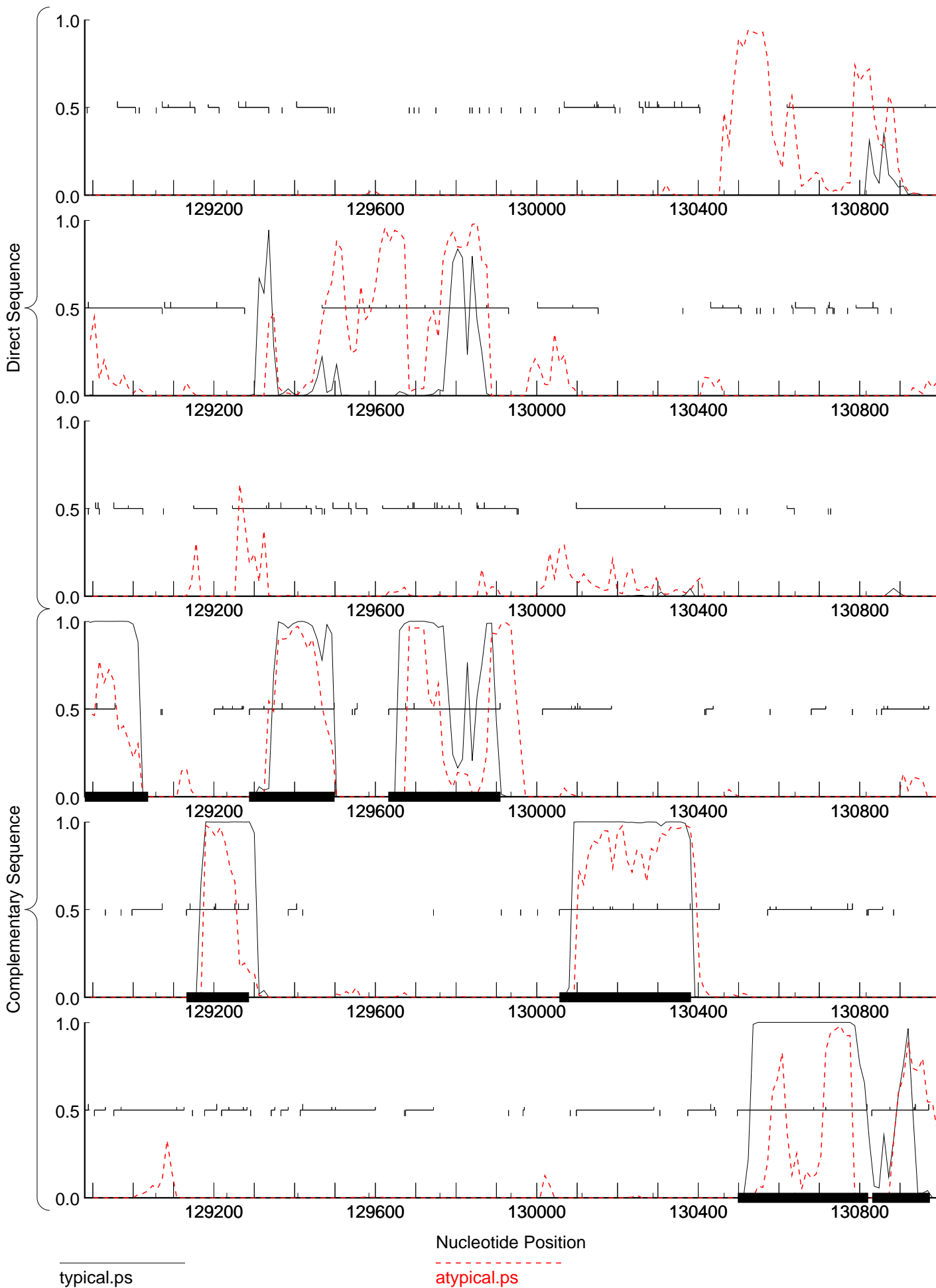
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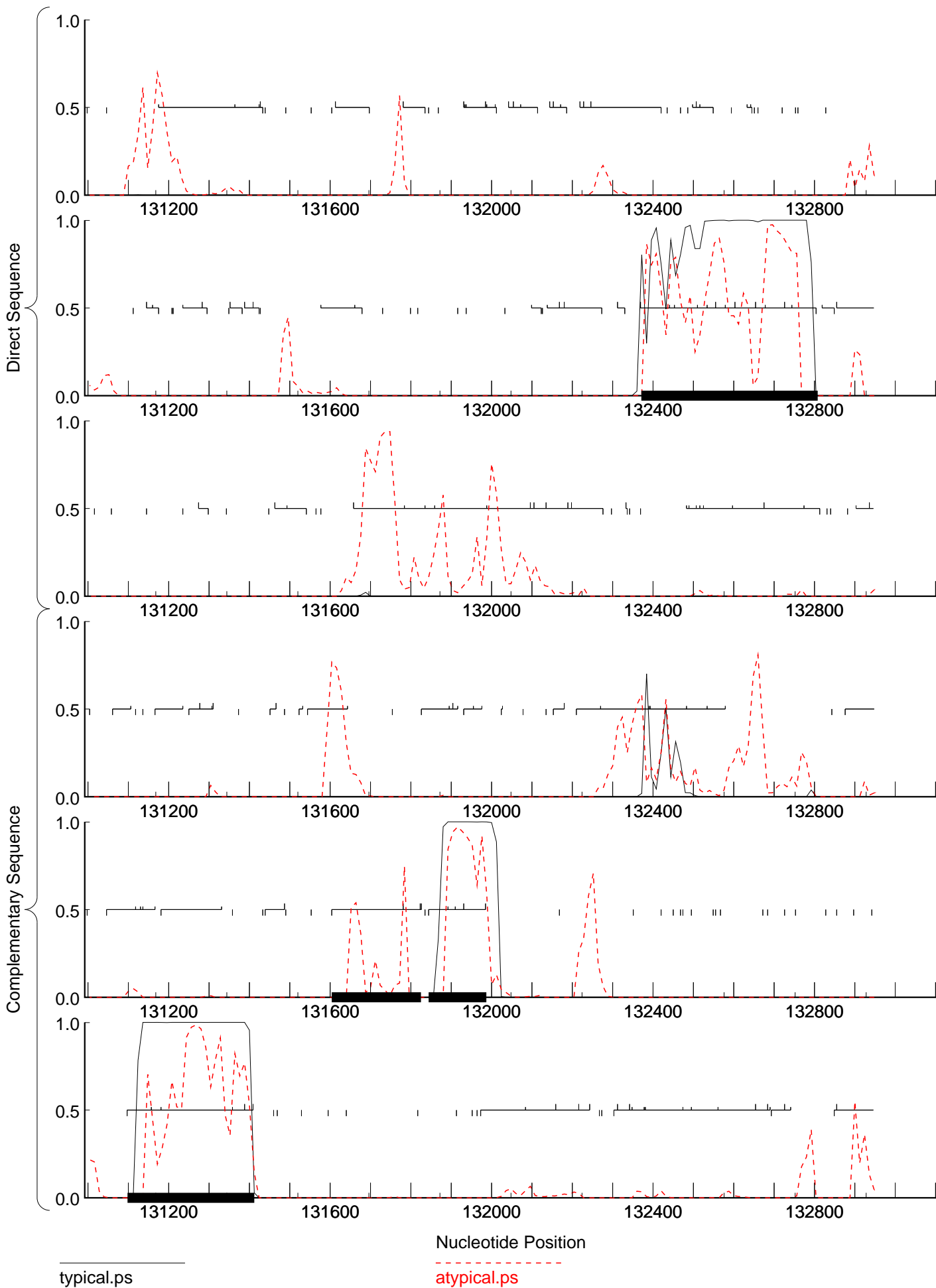


typical.ps

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