

GeneMark

Version 2.5p (09.08.06)

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PROGRAM INFORMATION

Sequence : Mycobacteriophage Inca complete genome, 74070 bp including 9-base 3' overhang (CGCTTGTC),
Analysis Date : 1/18/18 at 18:46:40
Pages : 37
Sequence Length : 74070 bp
GC Content : 62.80%

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,

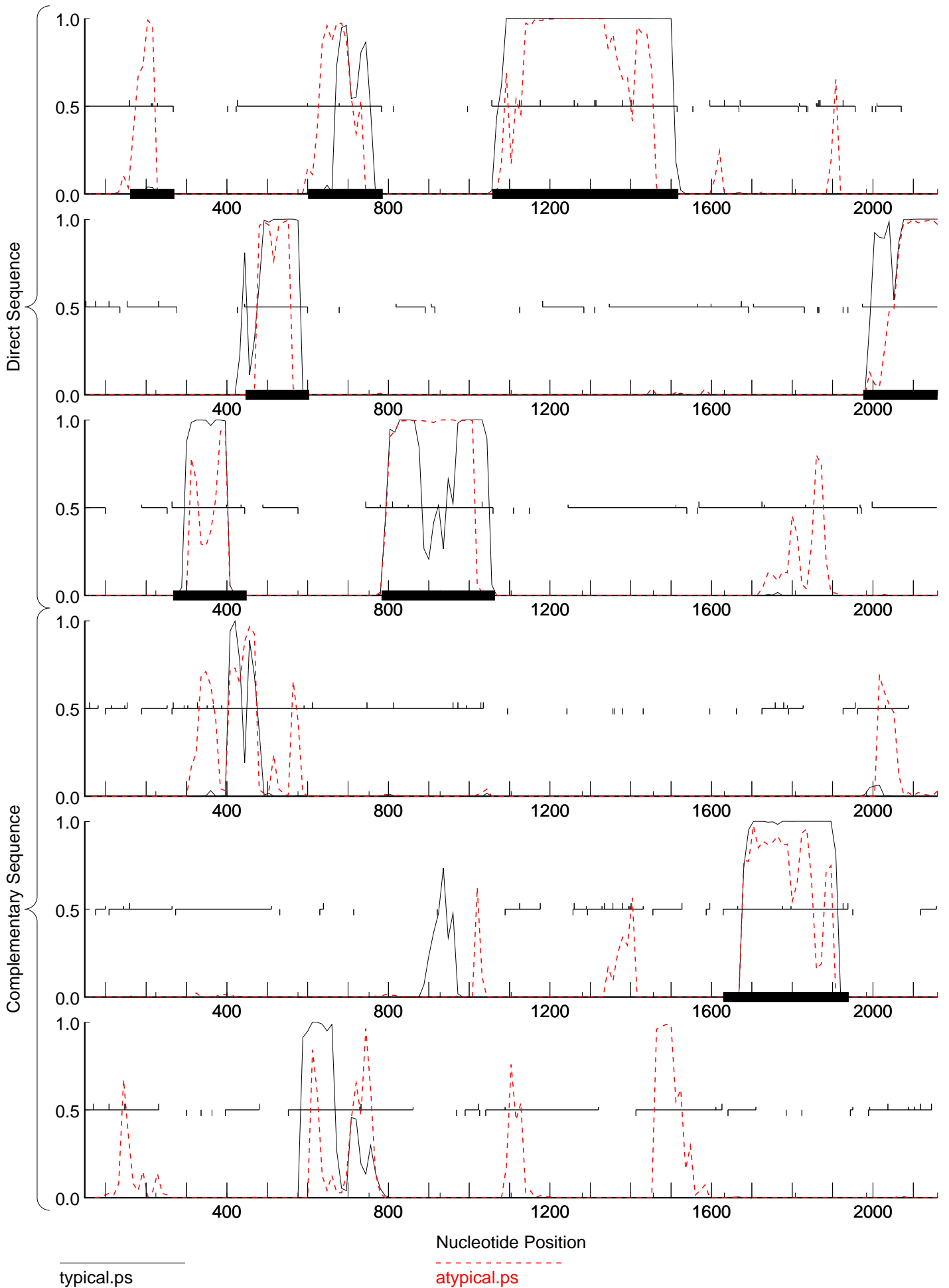
Matrix notes & comments

Training set derived by GeneMarkS, 4.30 October 2014
Thu Jan 18 18:46:40 2018

MATRIX INFORMATION

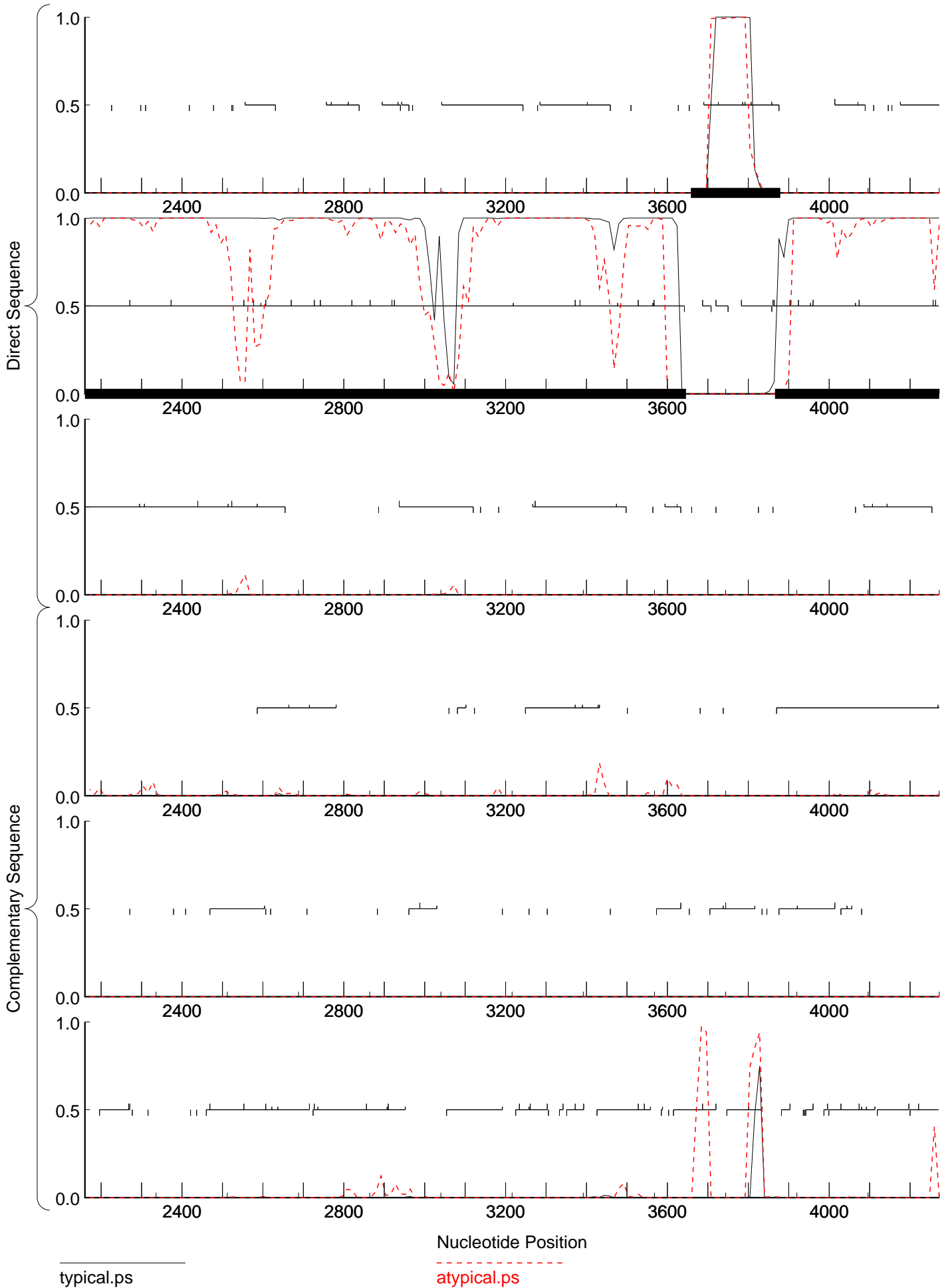
Matrix : -
Author : -
Order : 4

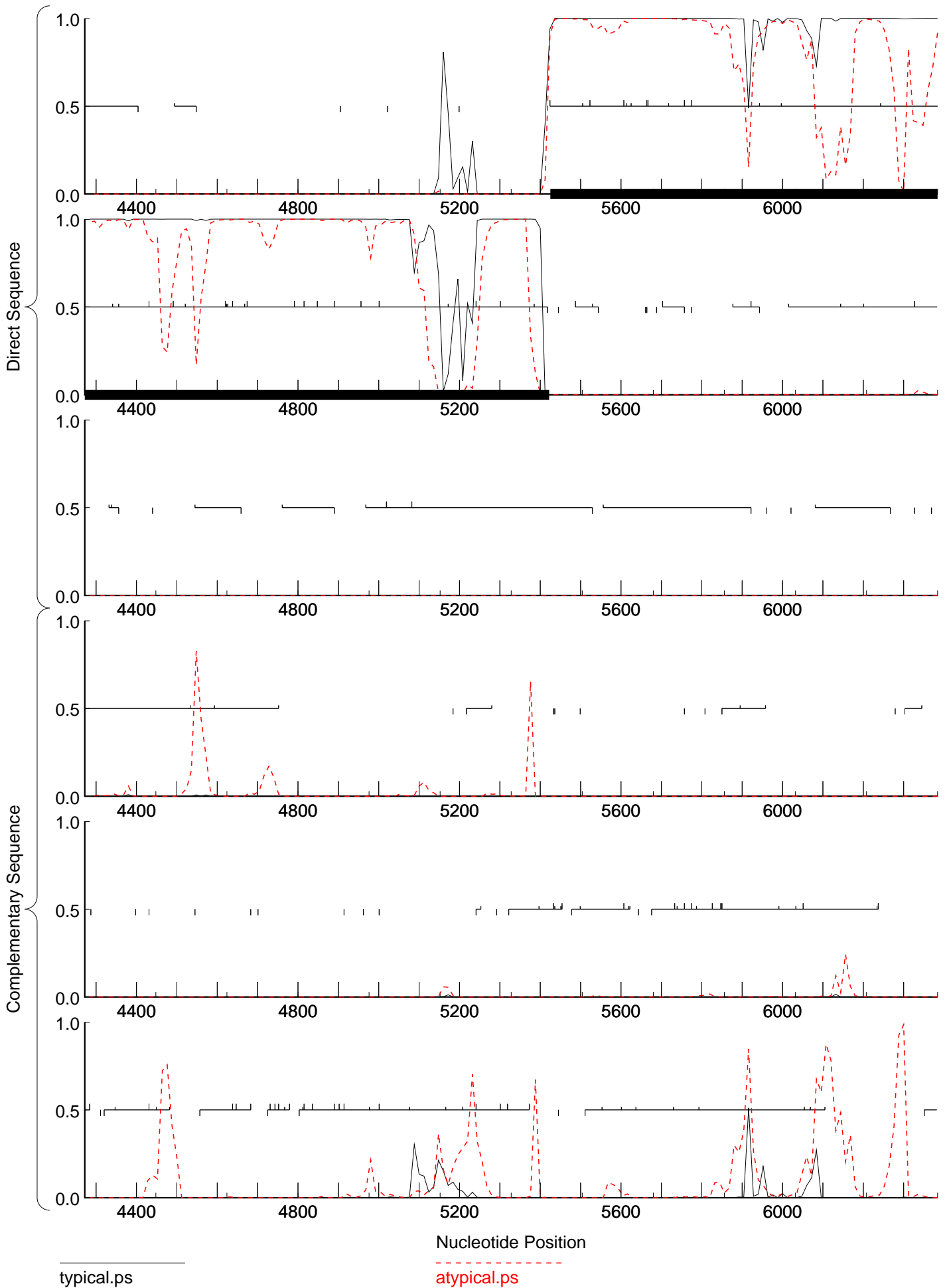
Send questions / comments to:
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Georgia Institute of Technology
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Atlanta, GA 30332-0230

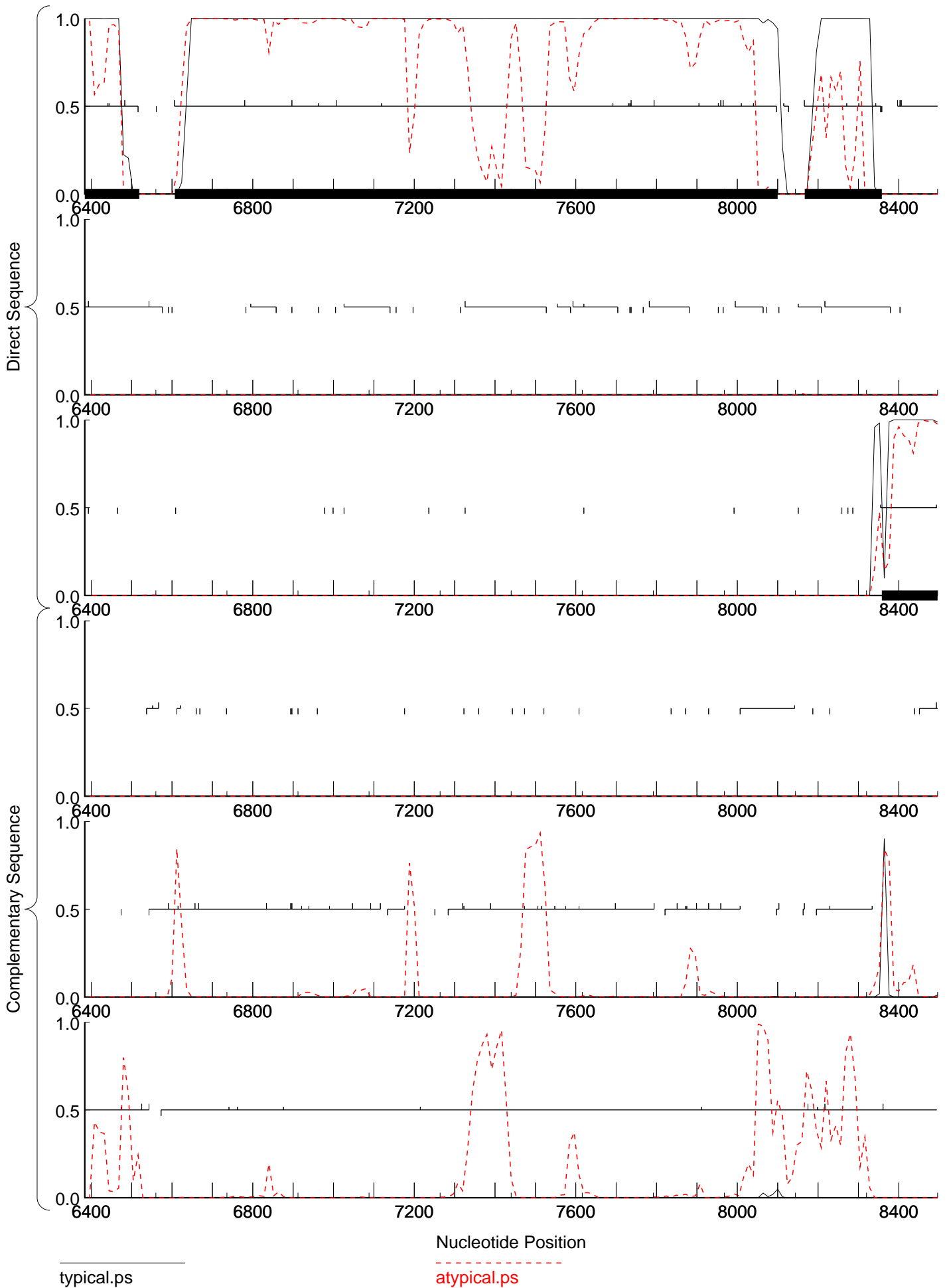


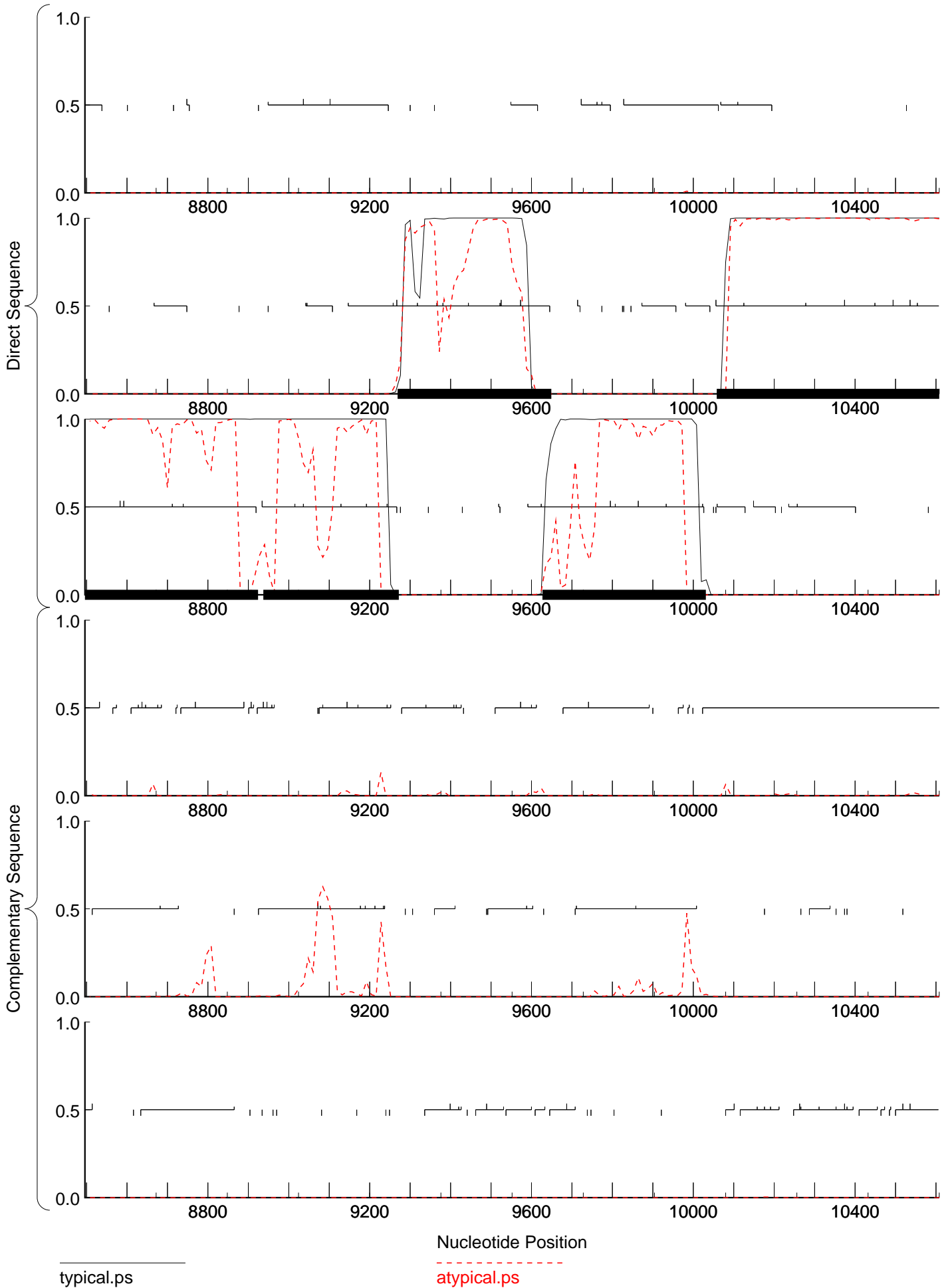
typical.ps

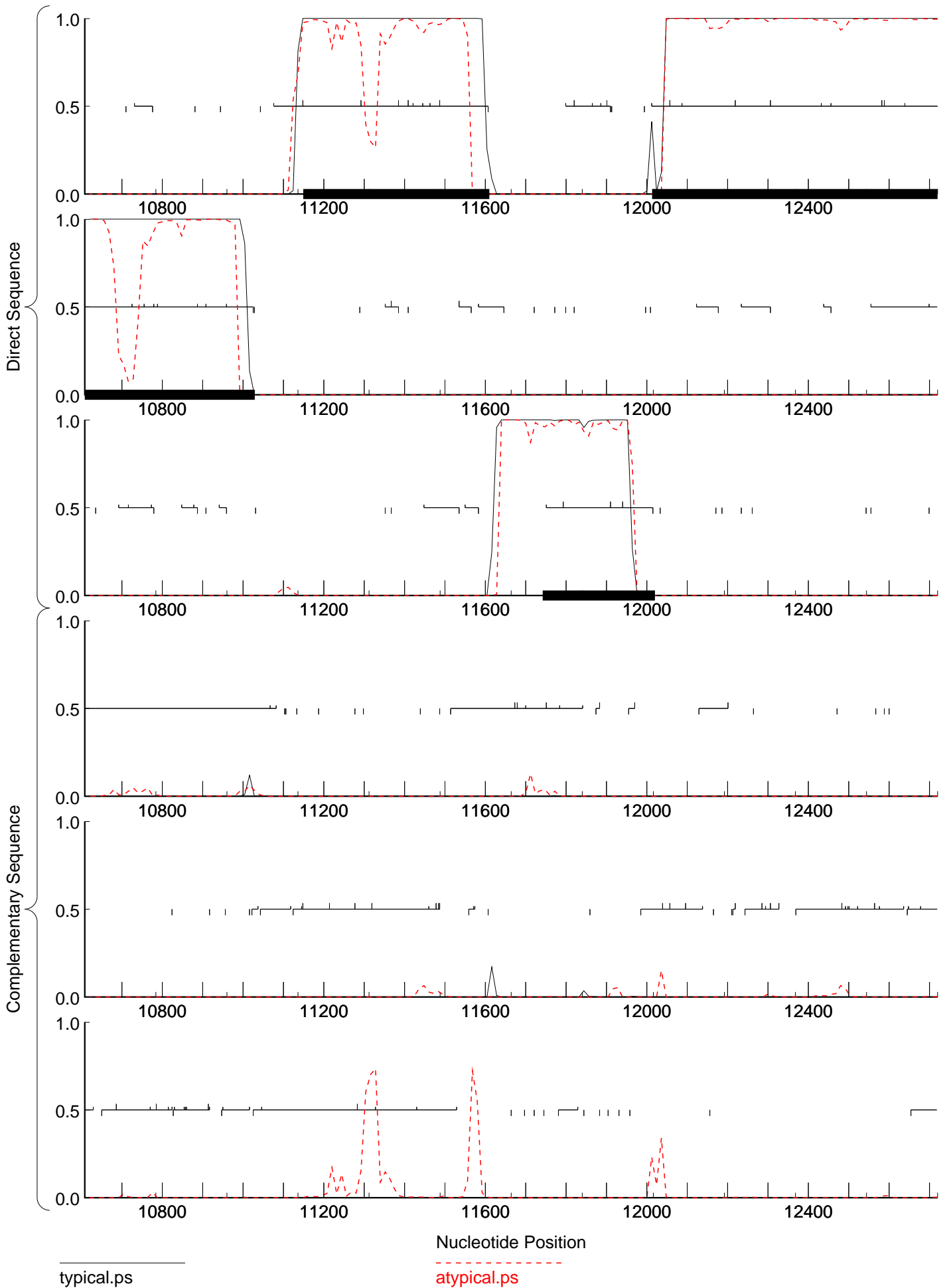
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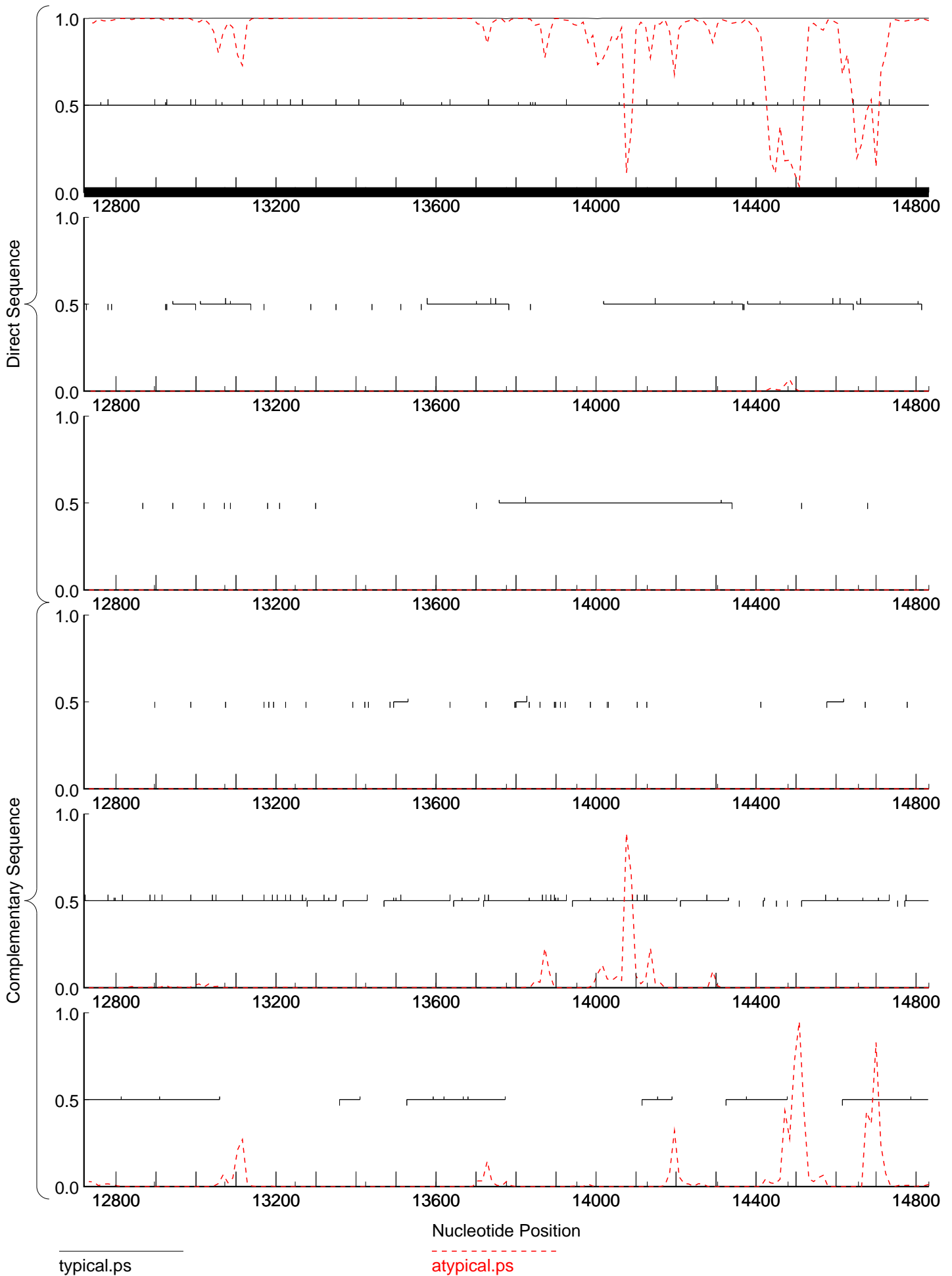


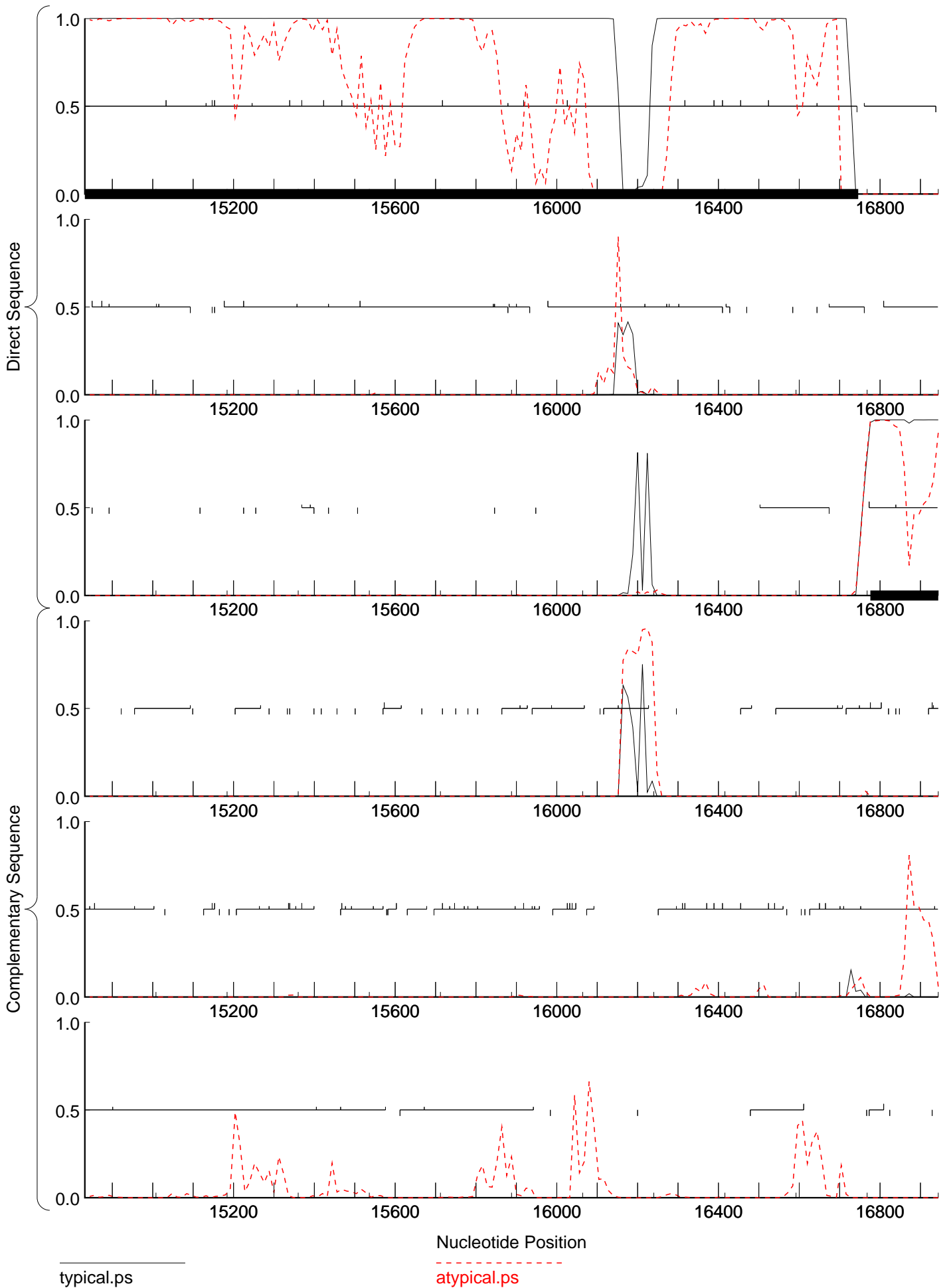


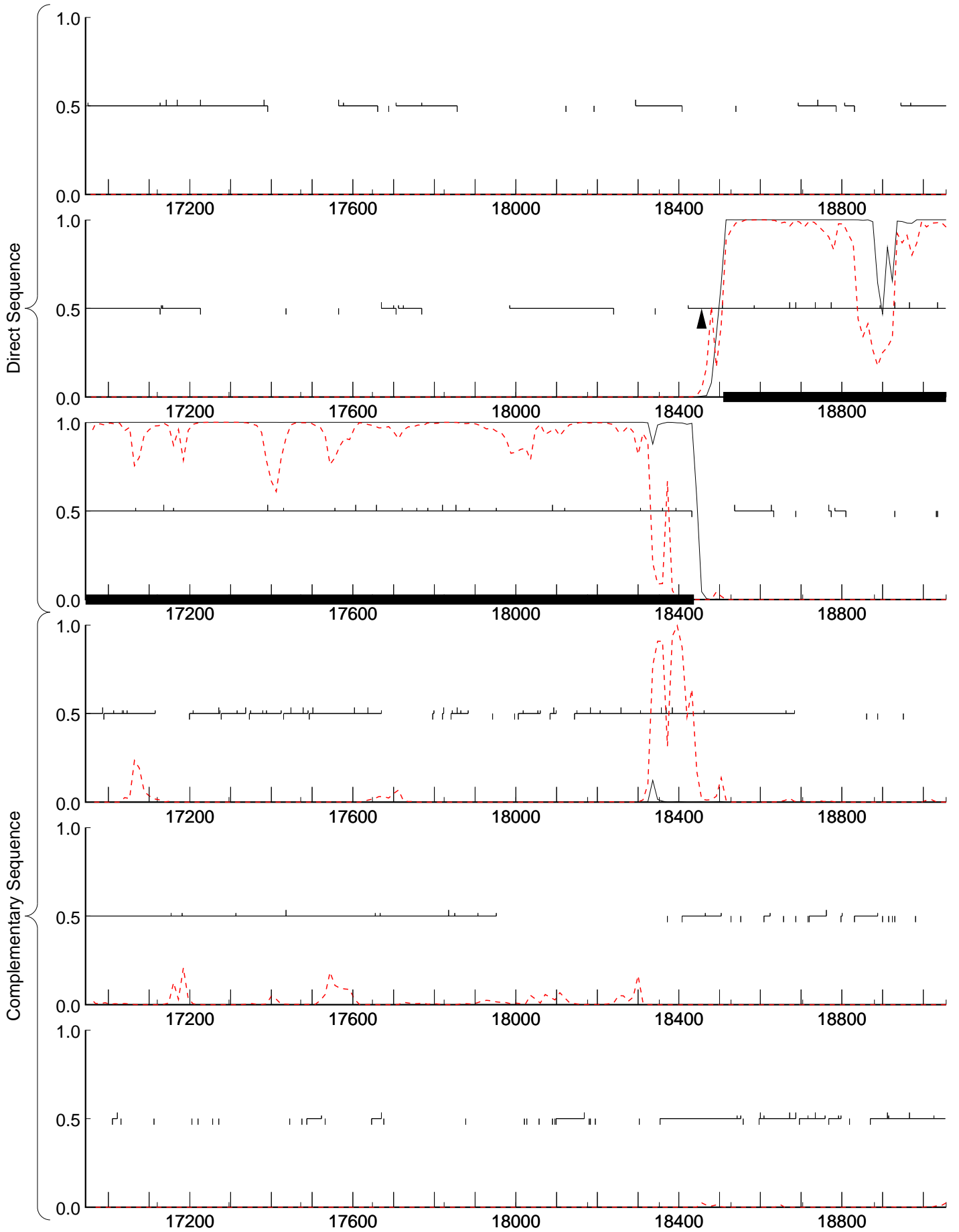








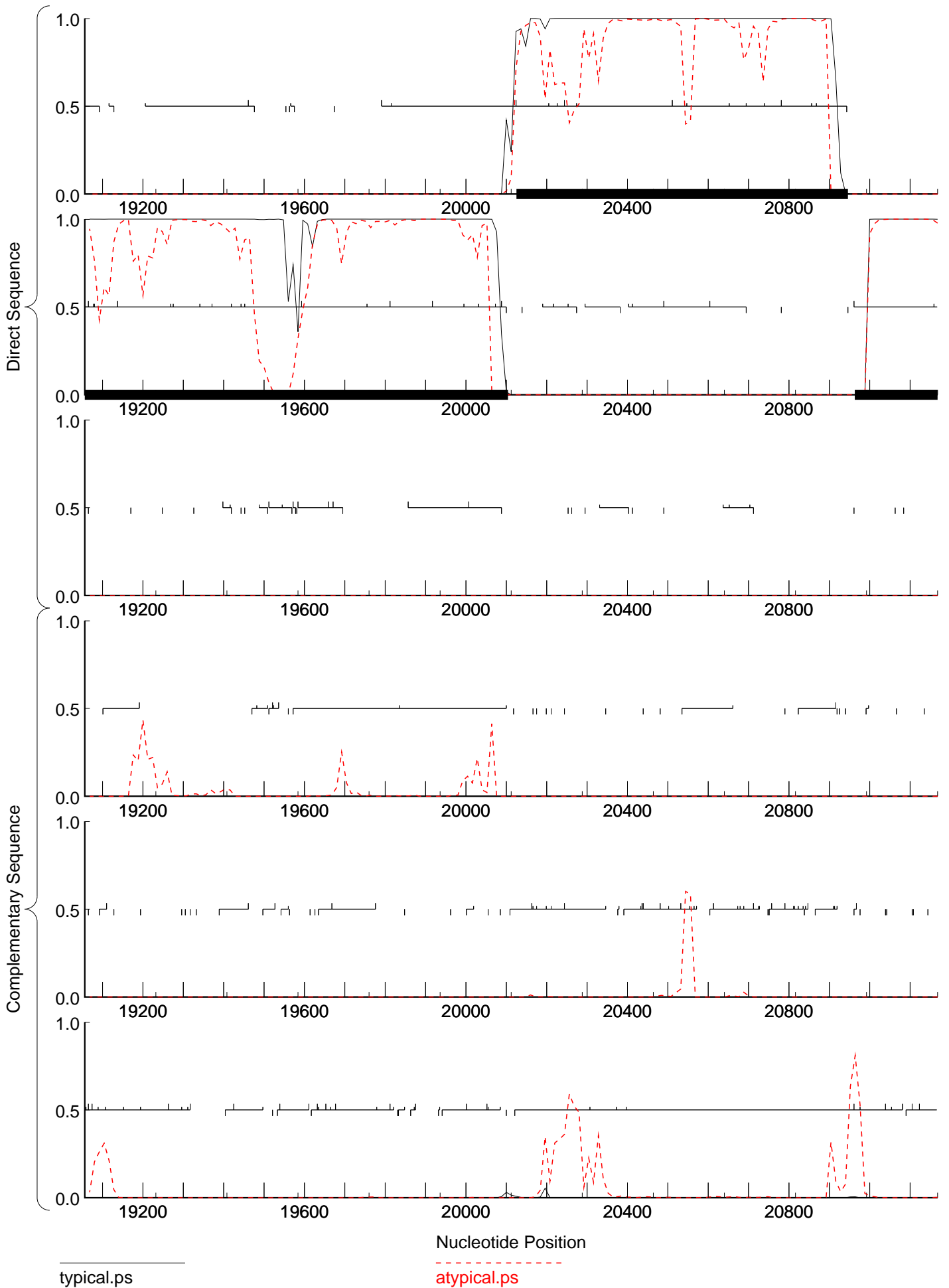


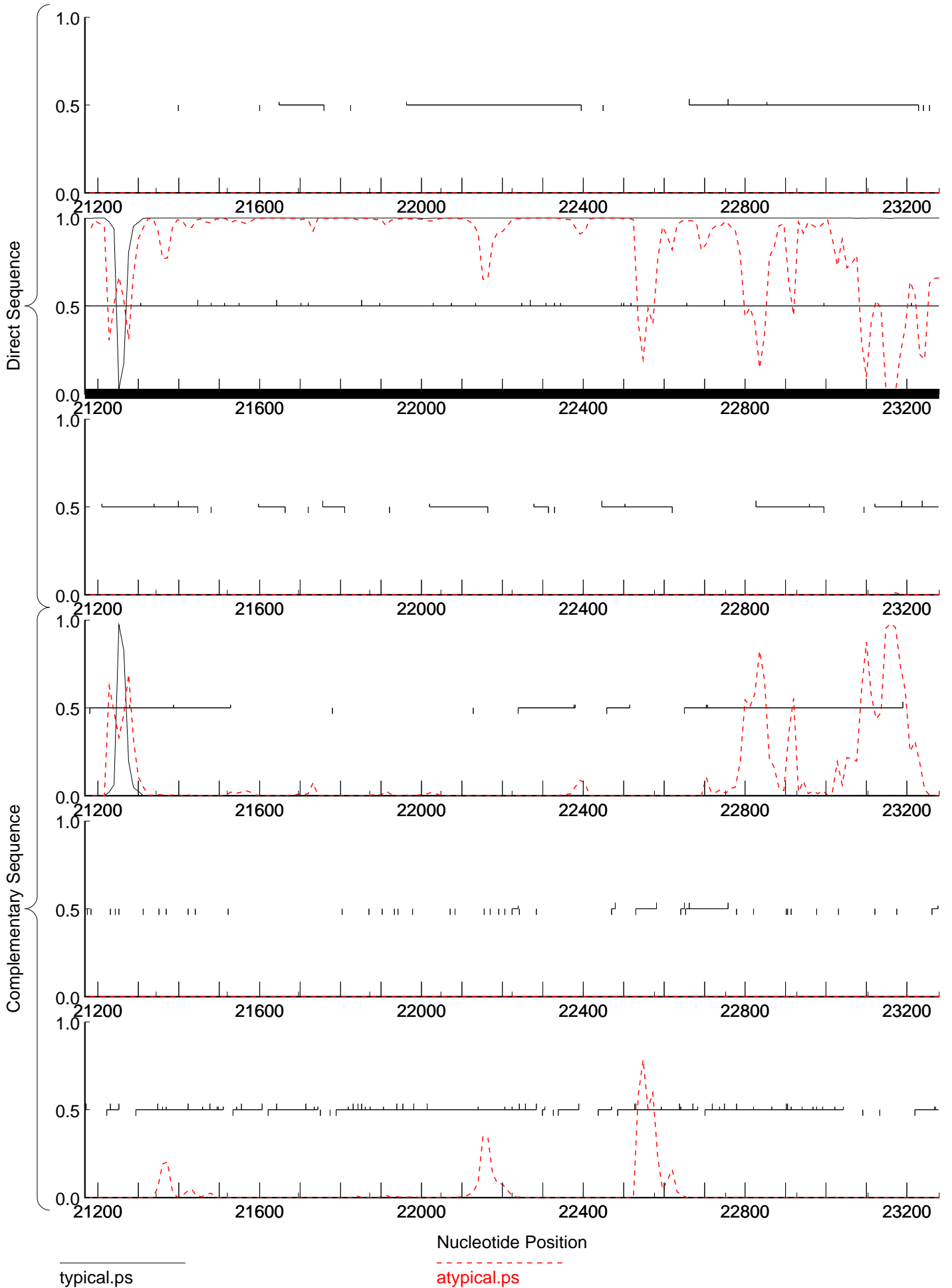


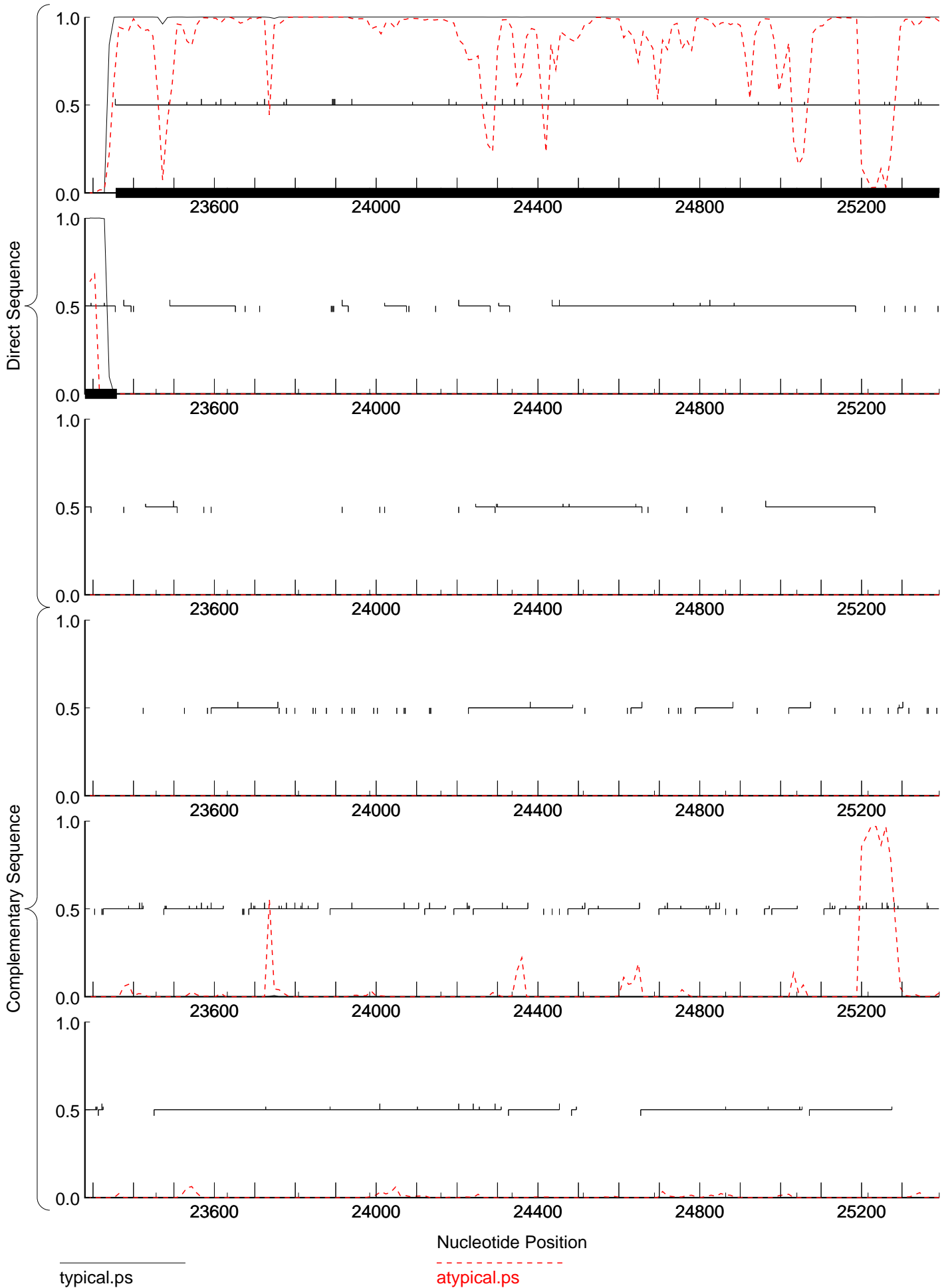
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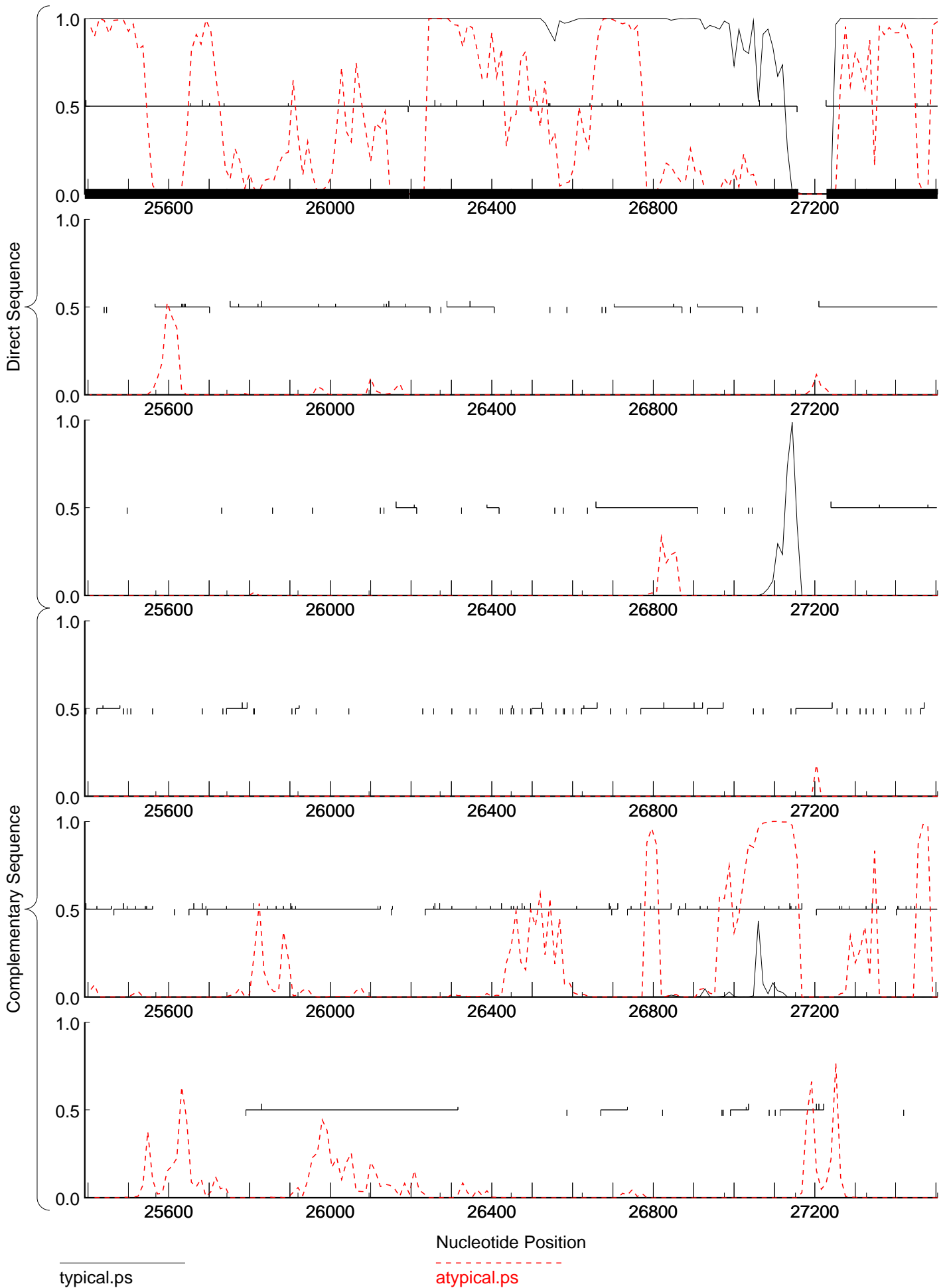
Nucleotide Position

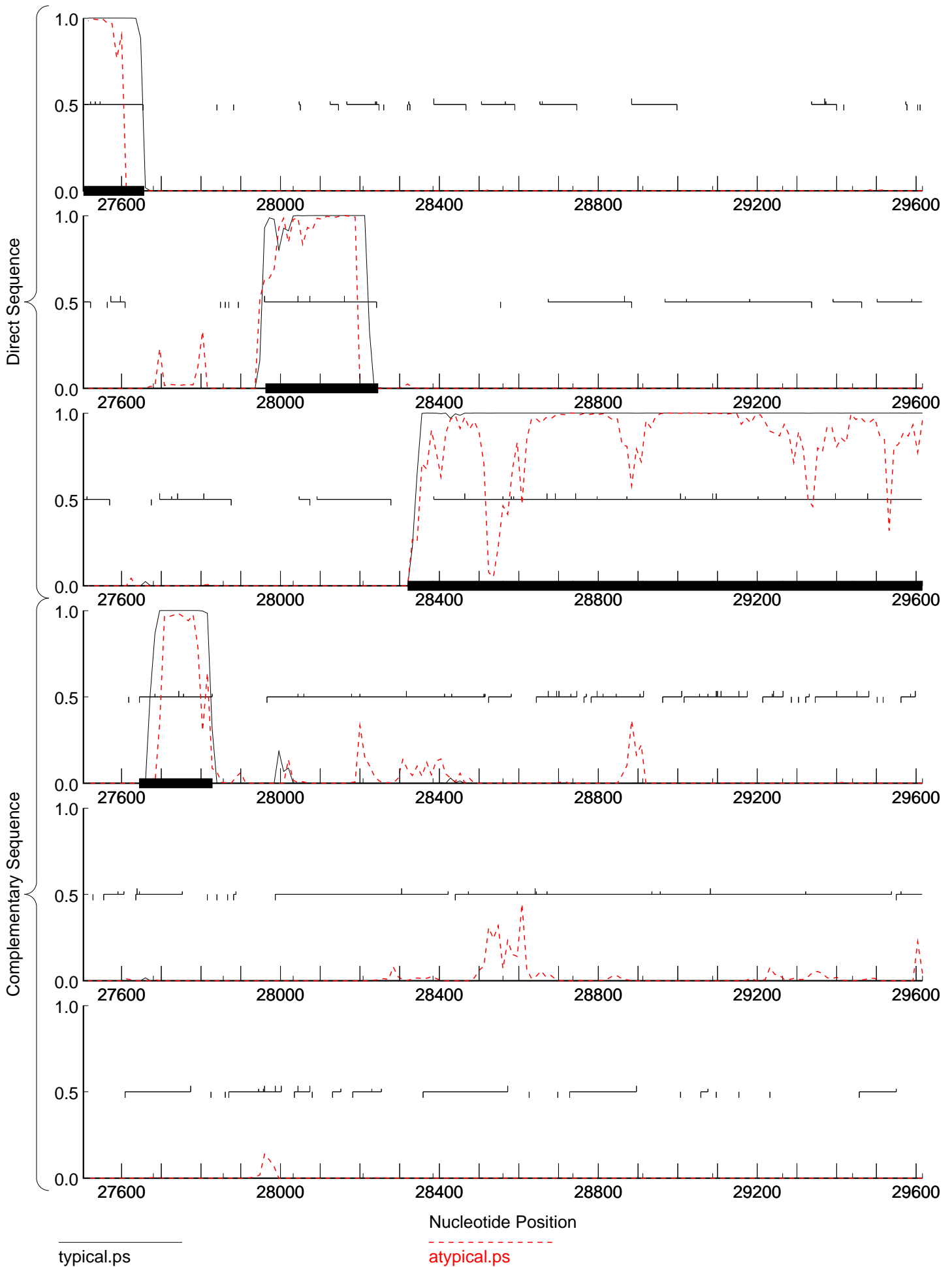
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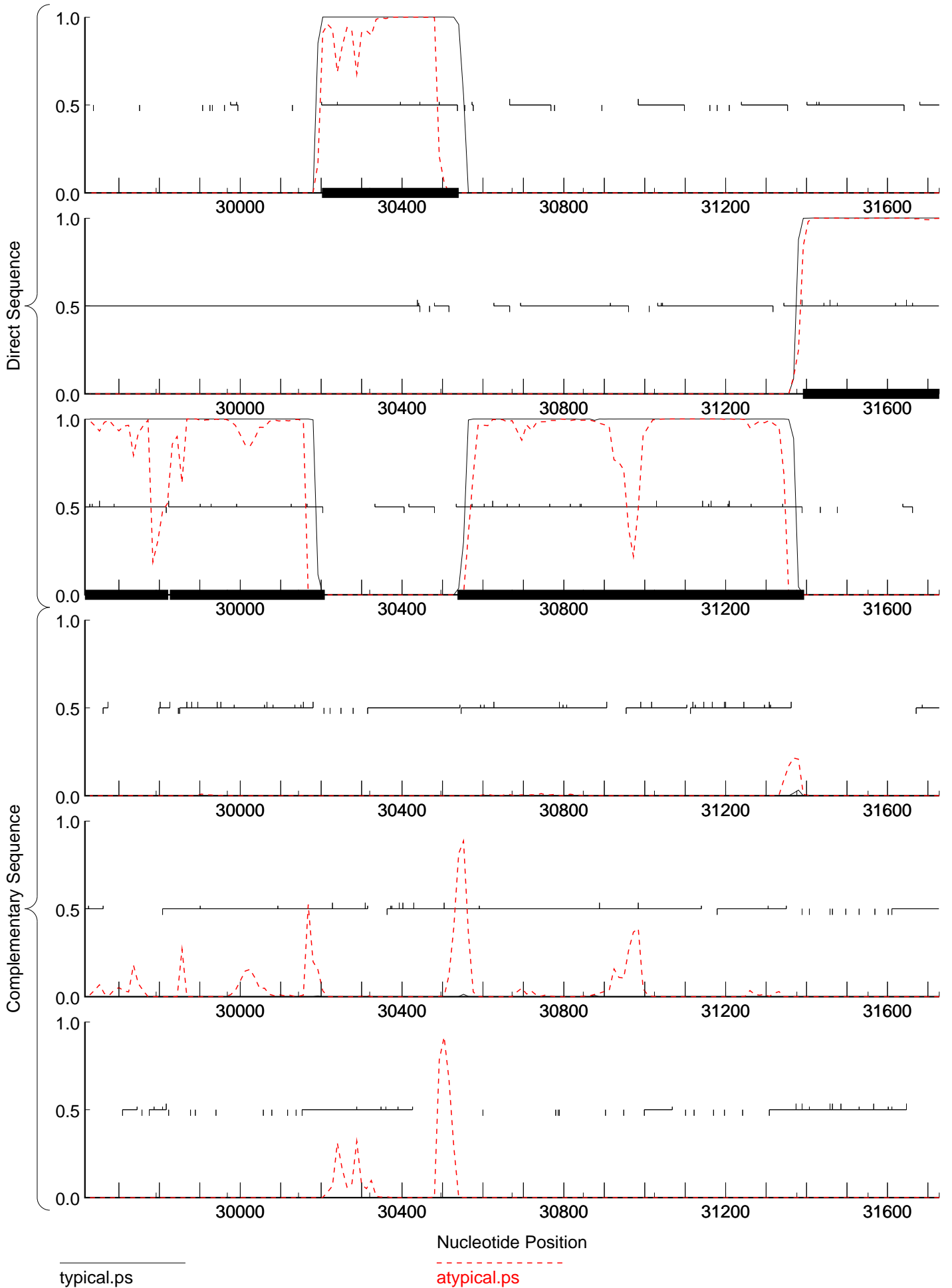


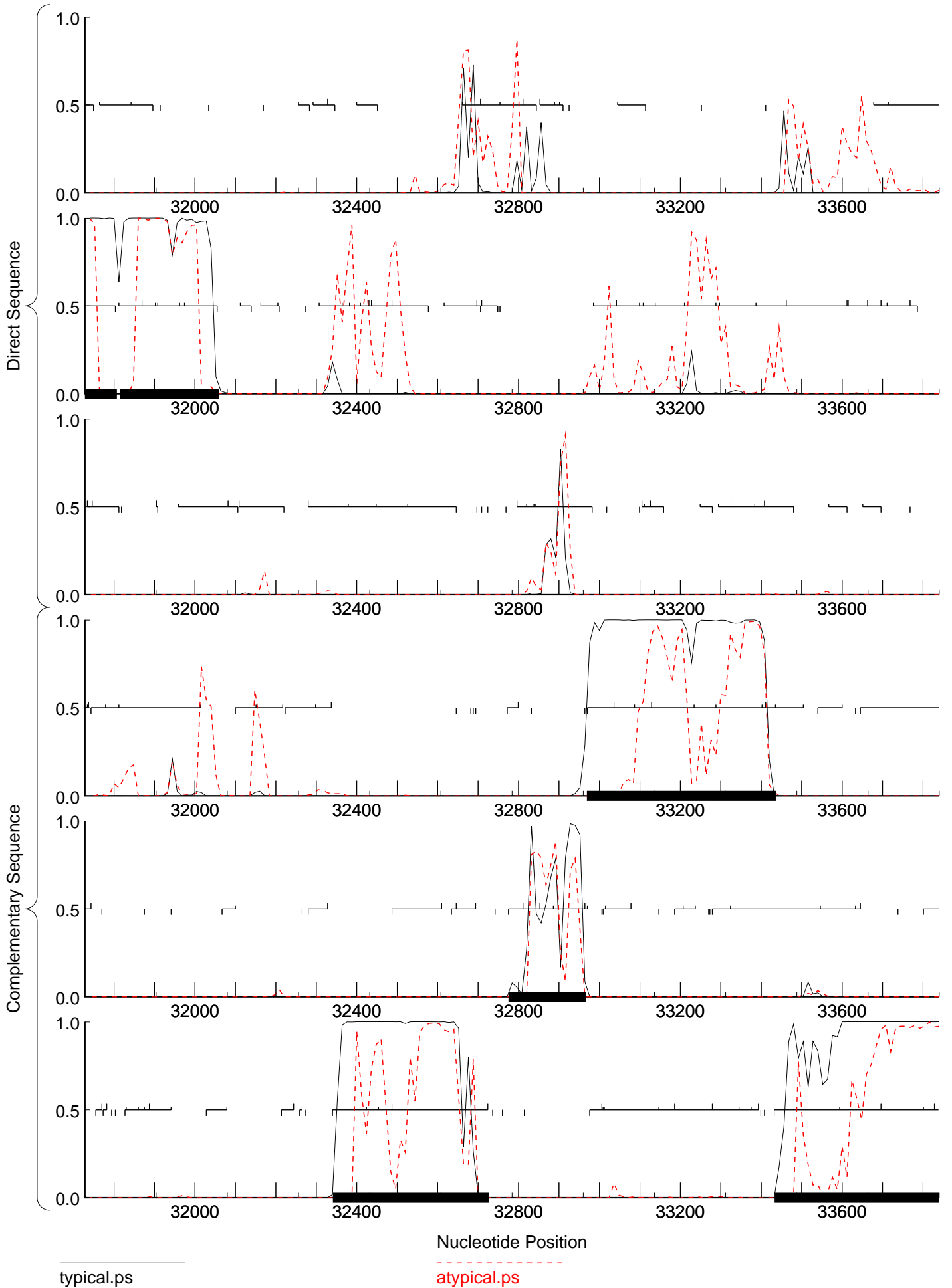




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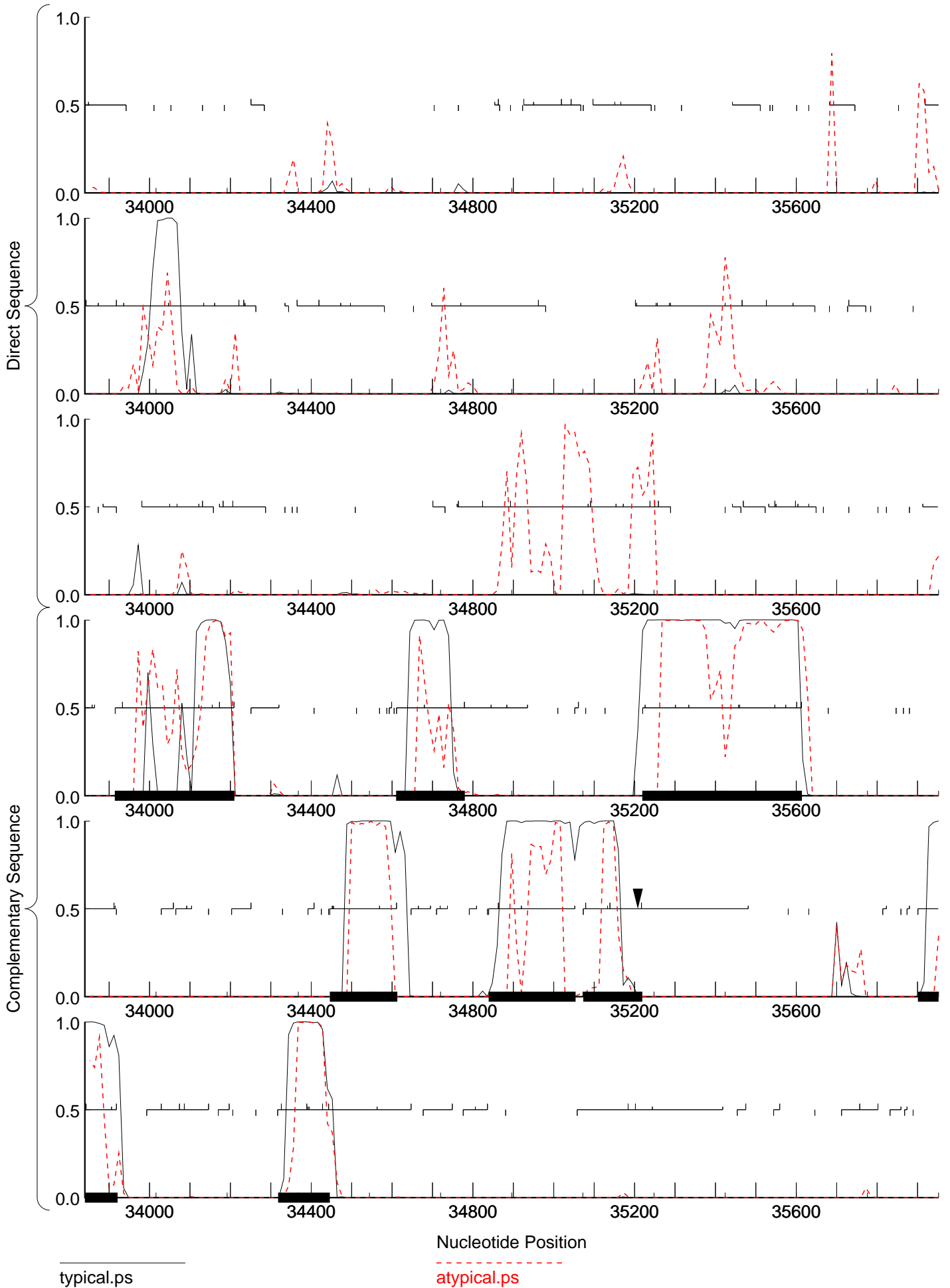
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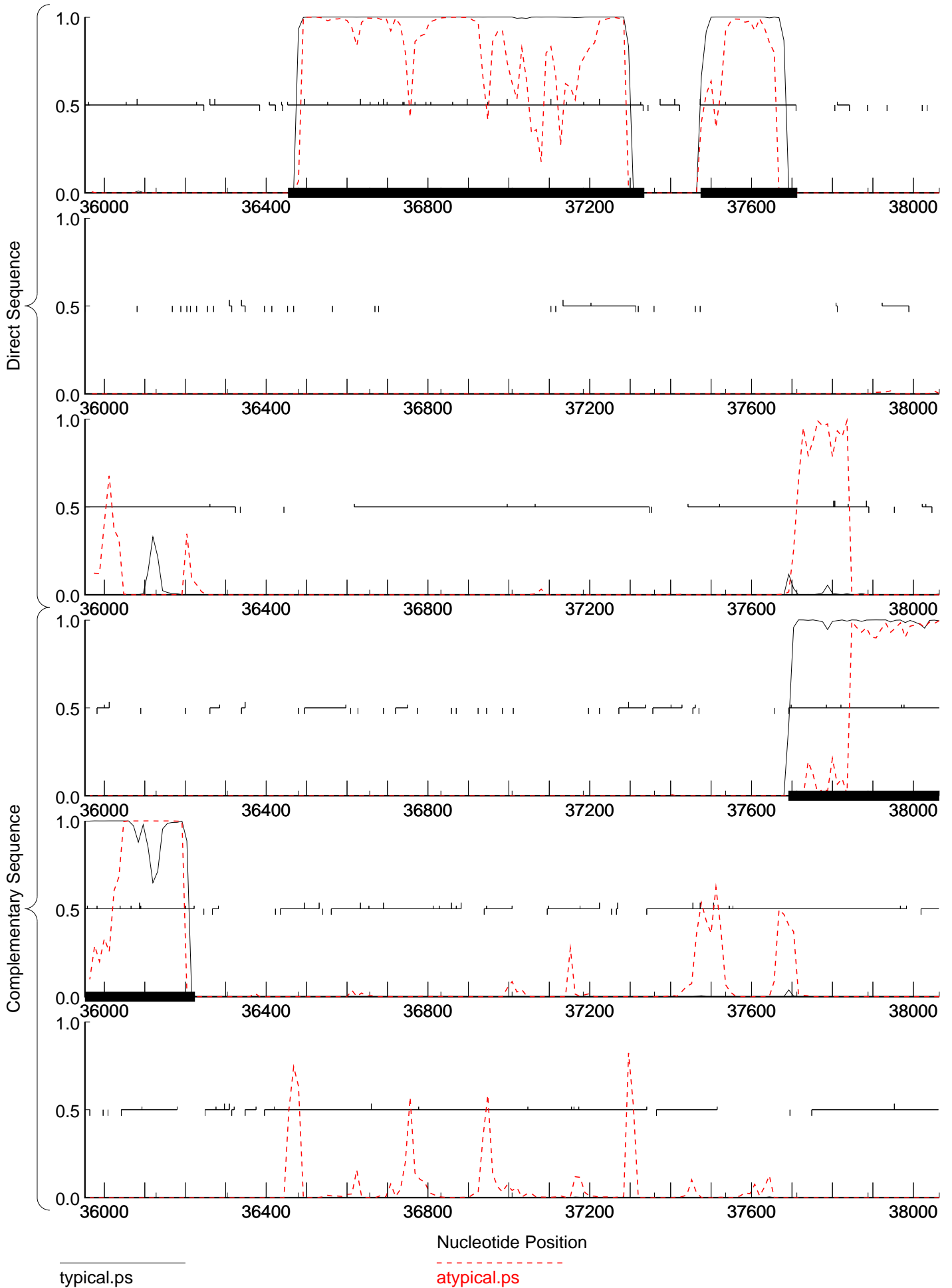


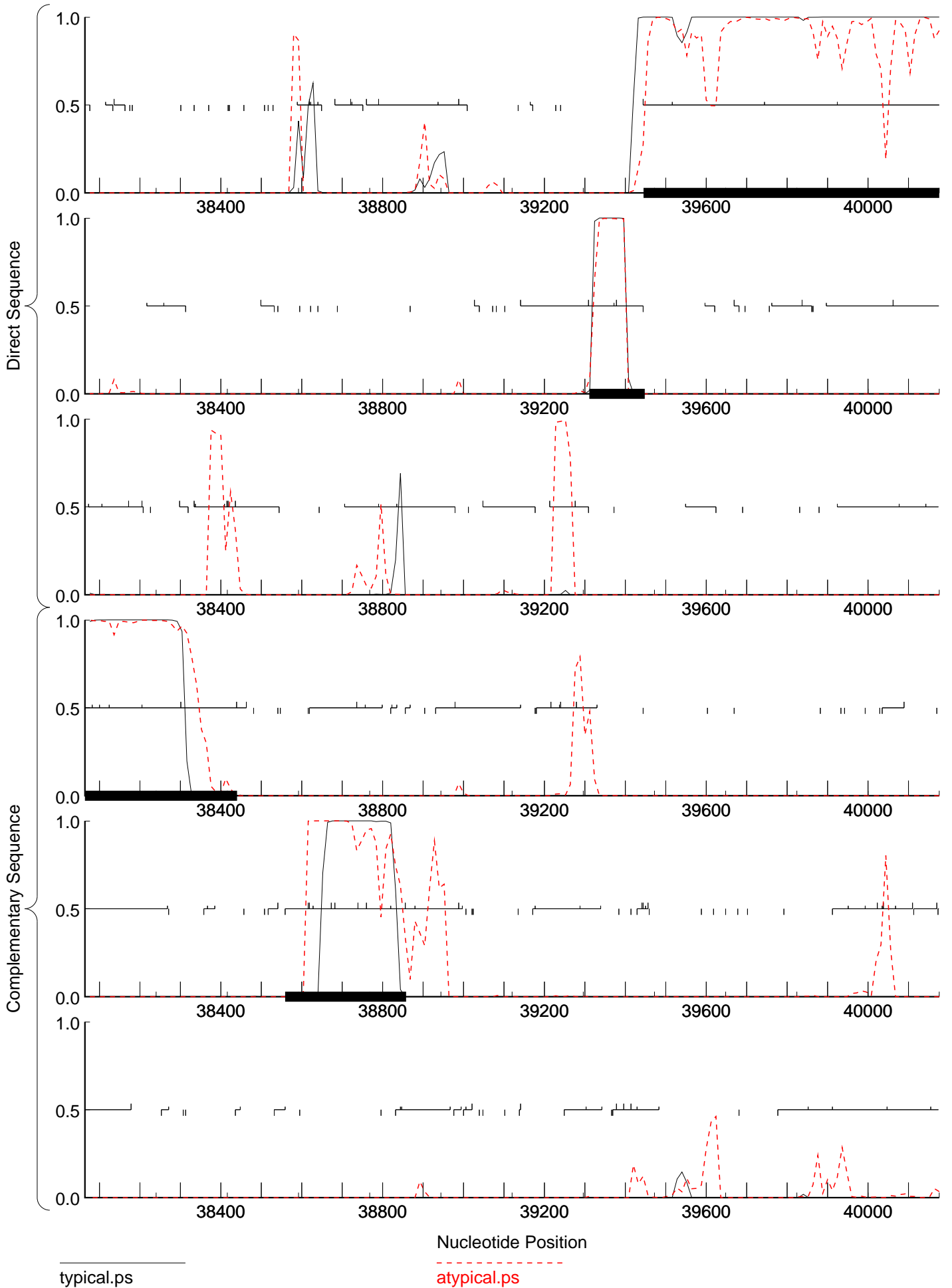


typical.ps

atypical.ps







typical.ps

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