

GeneMark

Version 2.5p (09.08.06)

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PROGRAM INFORMATION

Sequence : Mycobacterium phage BodEinwohner17 complete sequence, 60222 bp including 10-base 3' overhang
Analysis Date : 5/24/18 at 4:05:25
Pages : 30
Sequence Length : 60222 bp
GC Content : 61.34%

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,

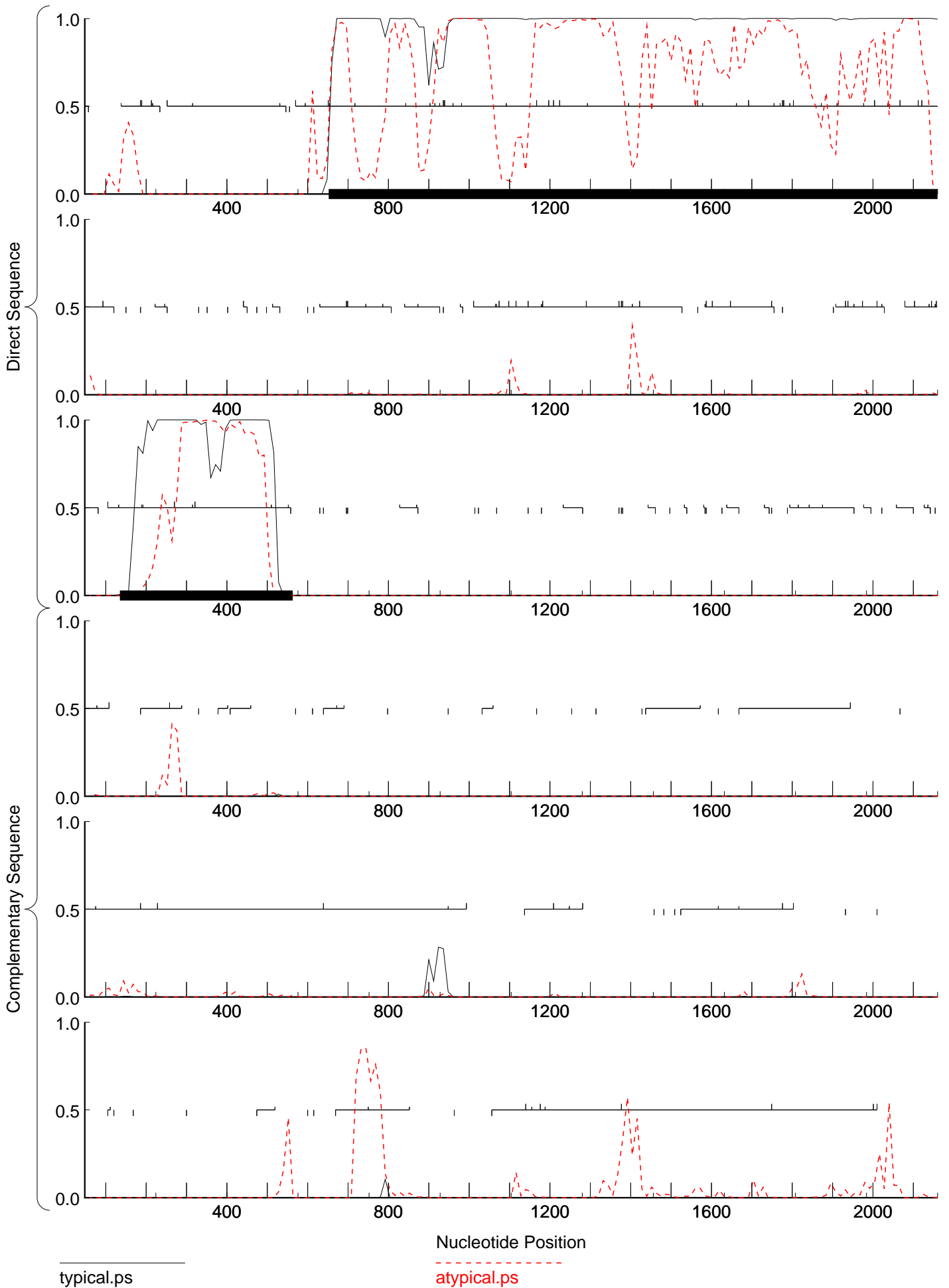
Matrix notes & comments

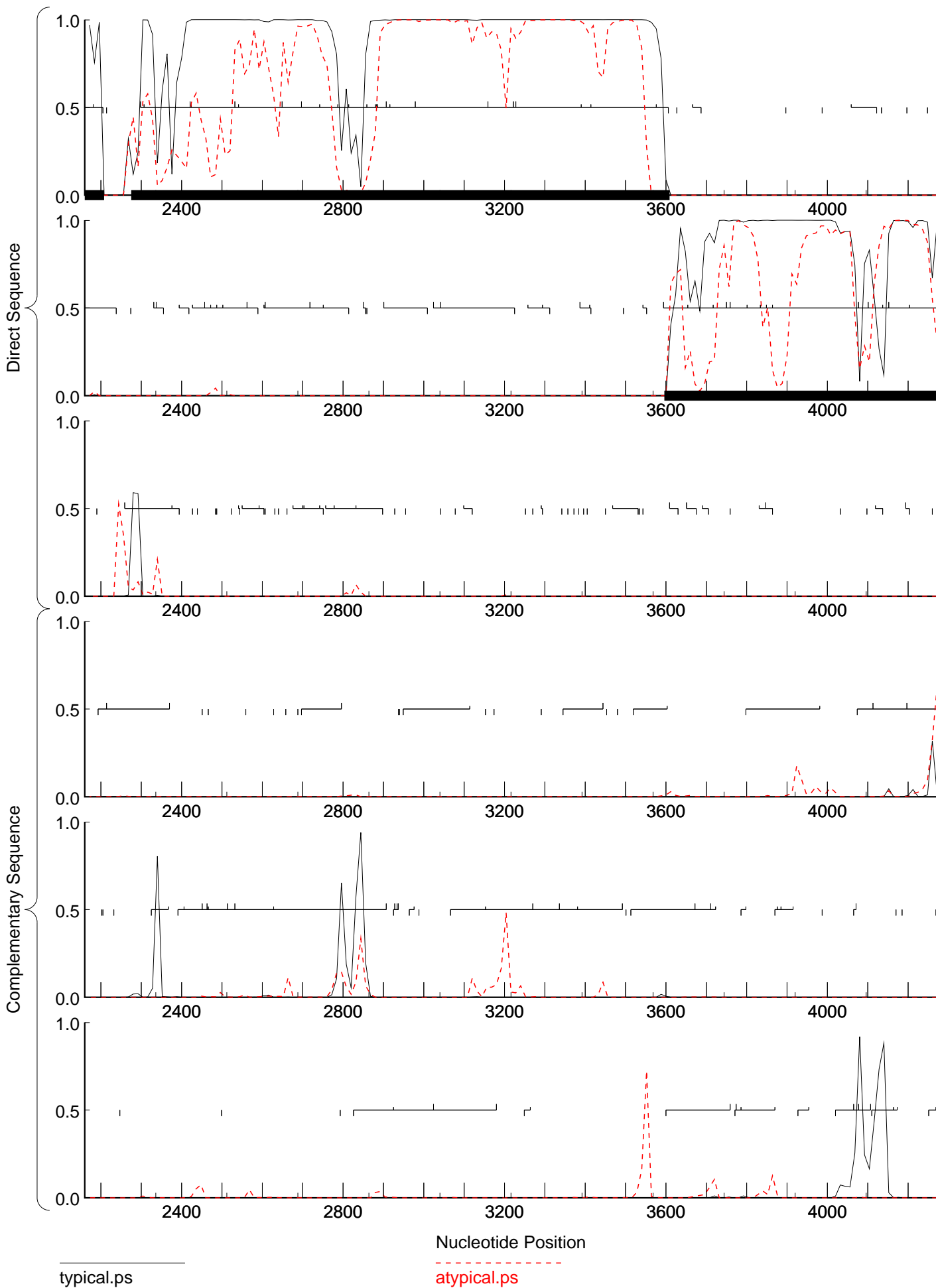
Training set derived by GeneMarkS, 4.30 October 2014
Thu May 24 04:05:25 2018

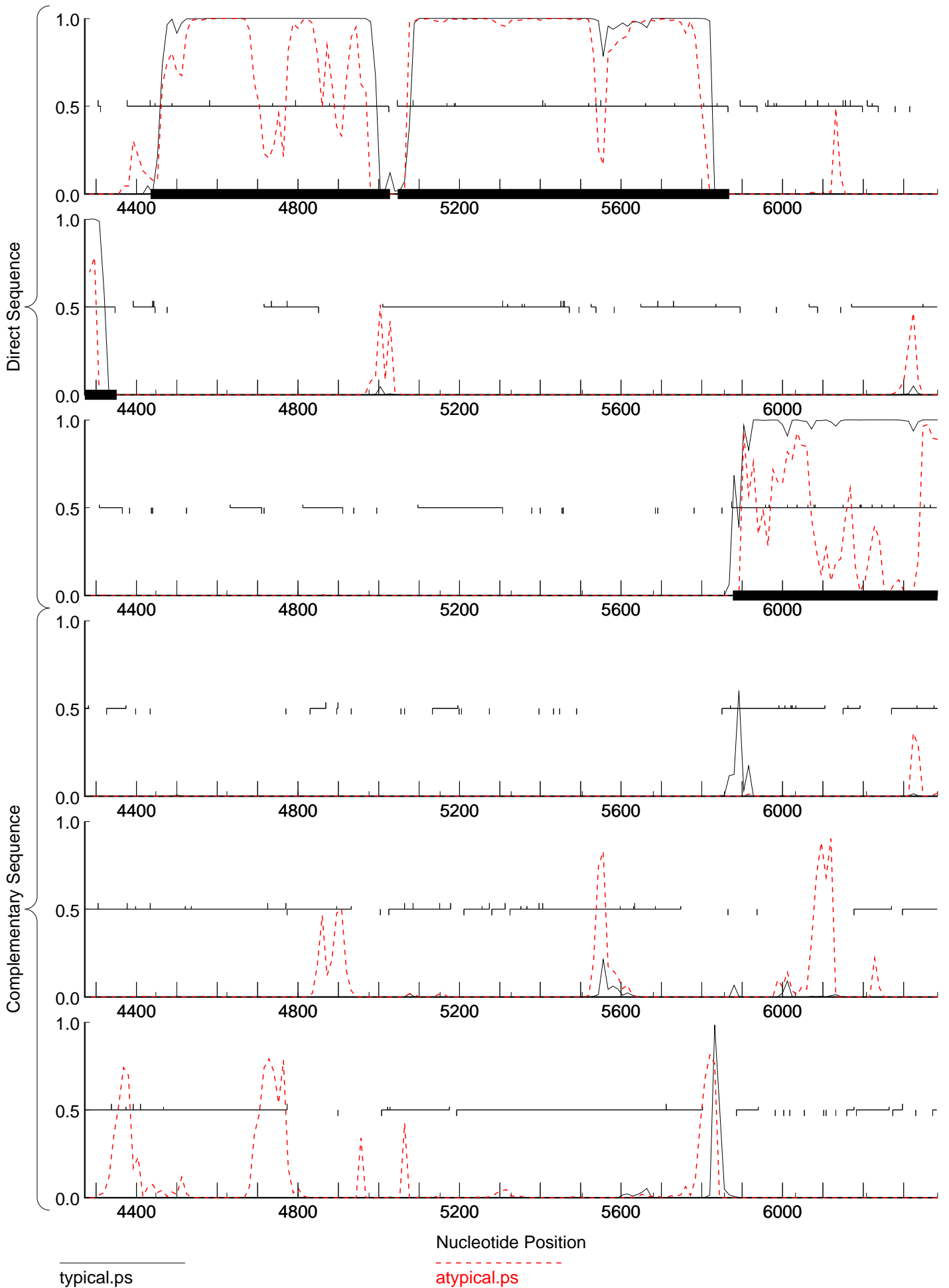
MATRIX INFORMATION

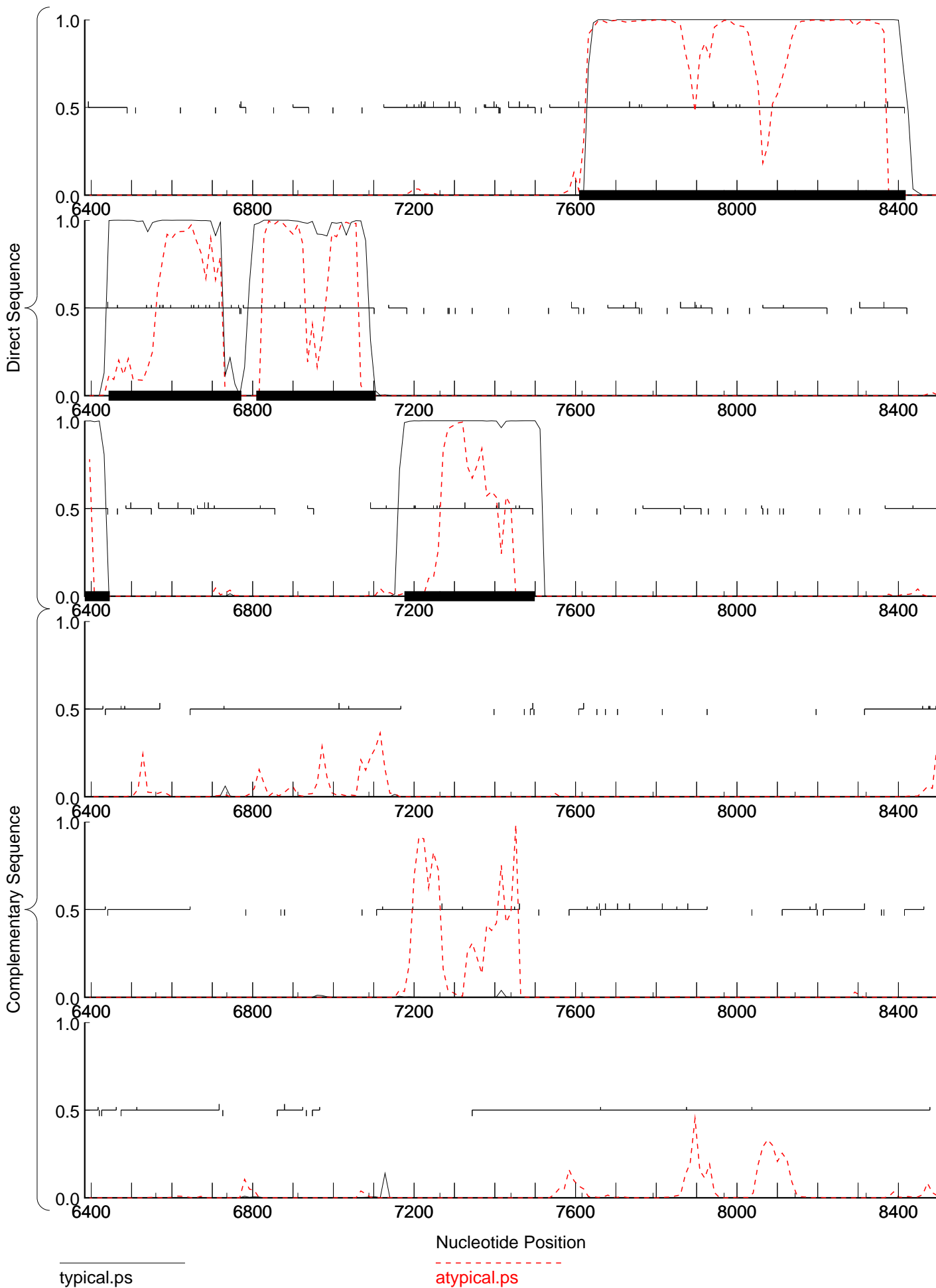
Matrix : -
Author : -
Order : 4

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



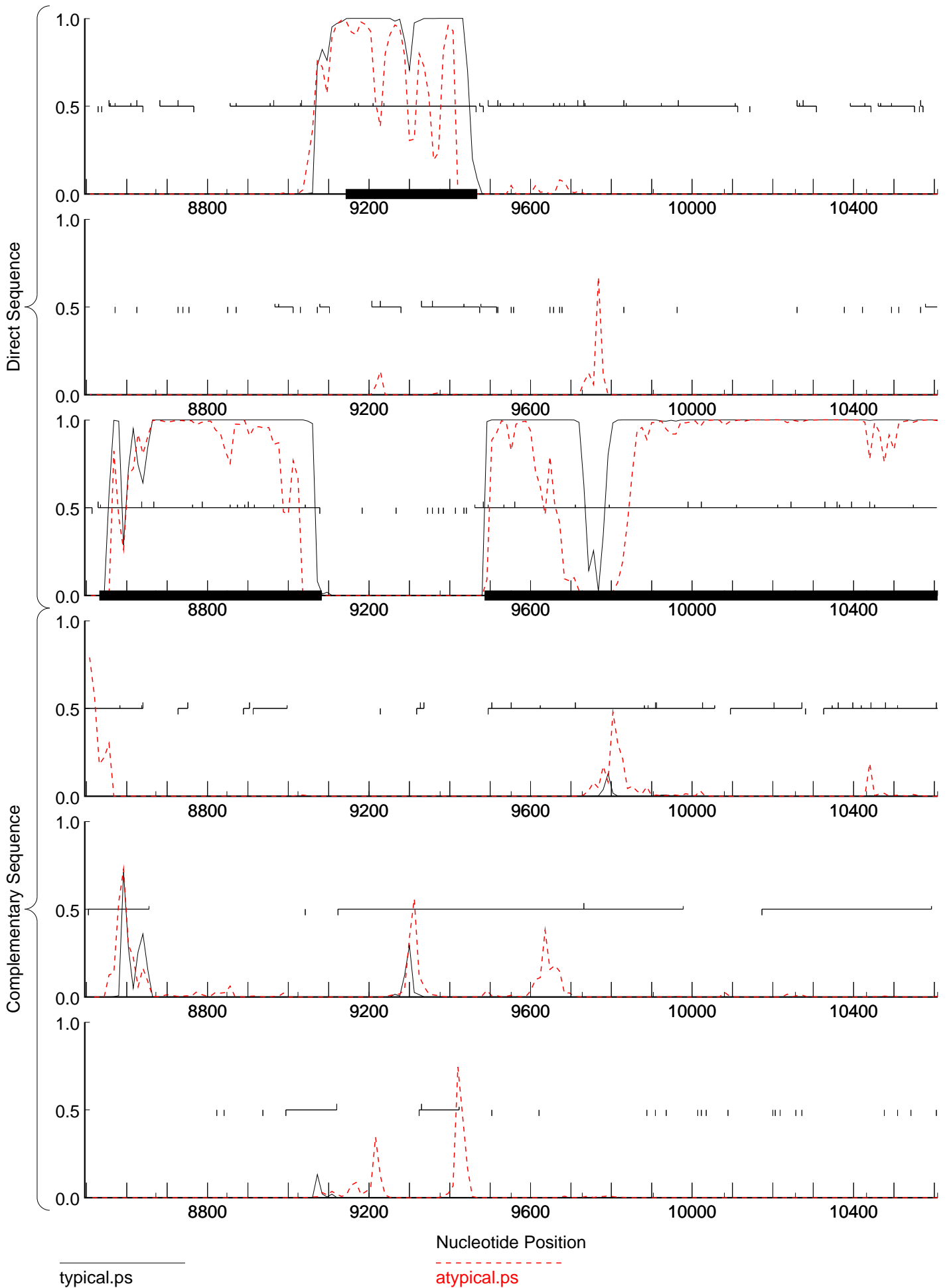


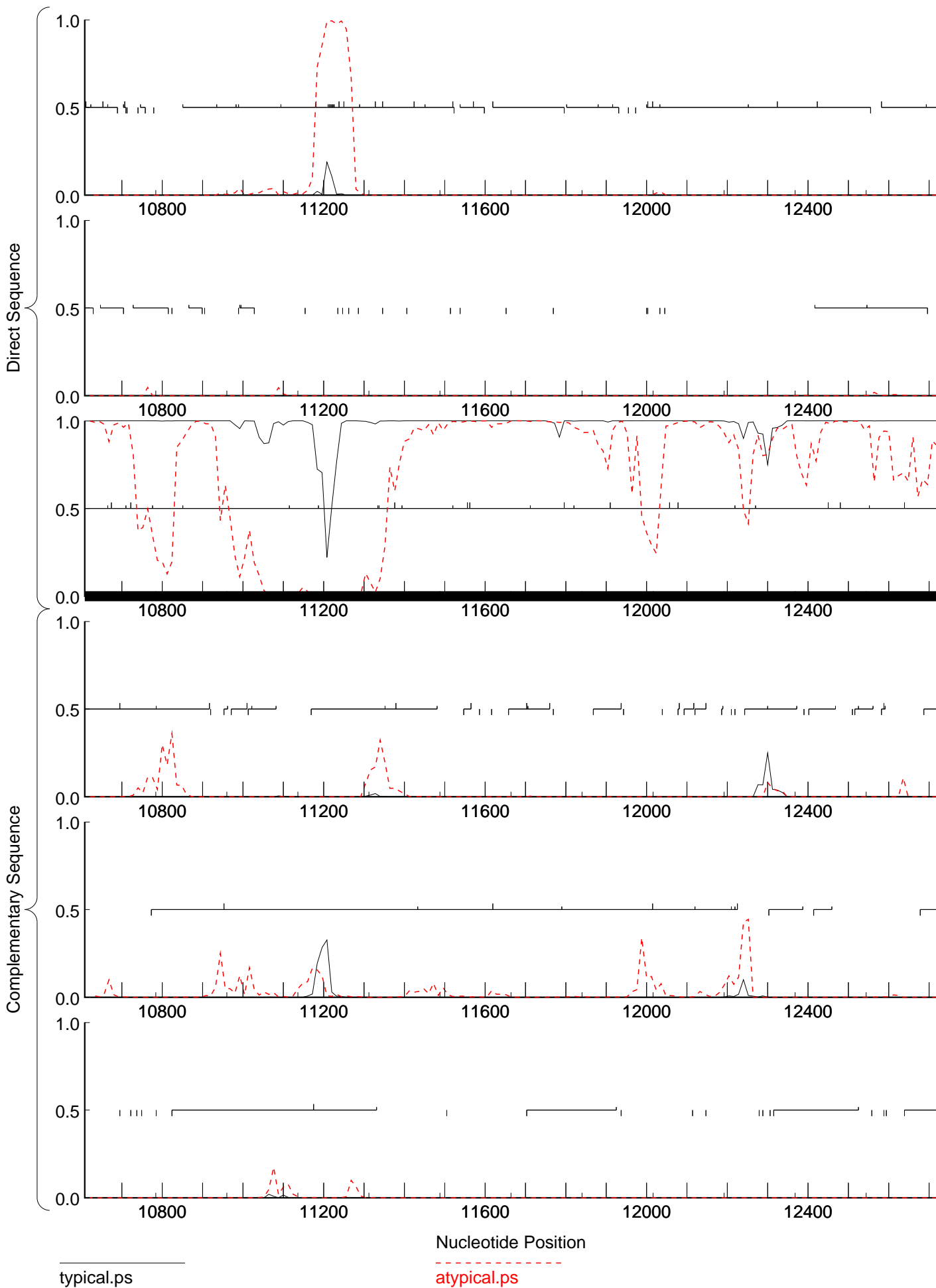


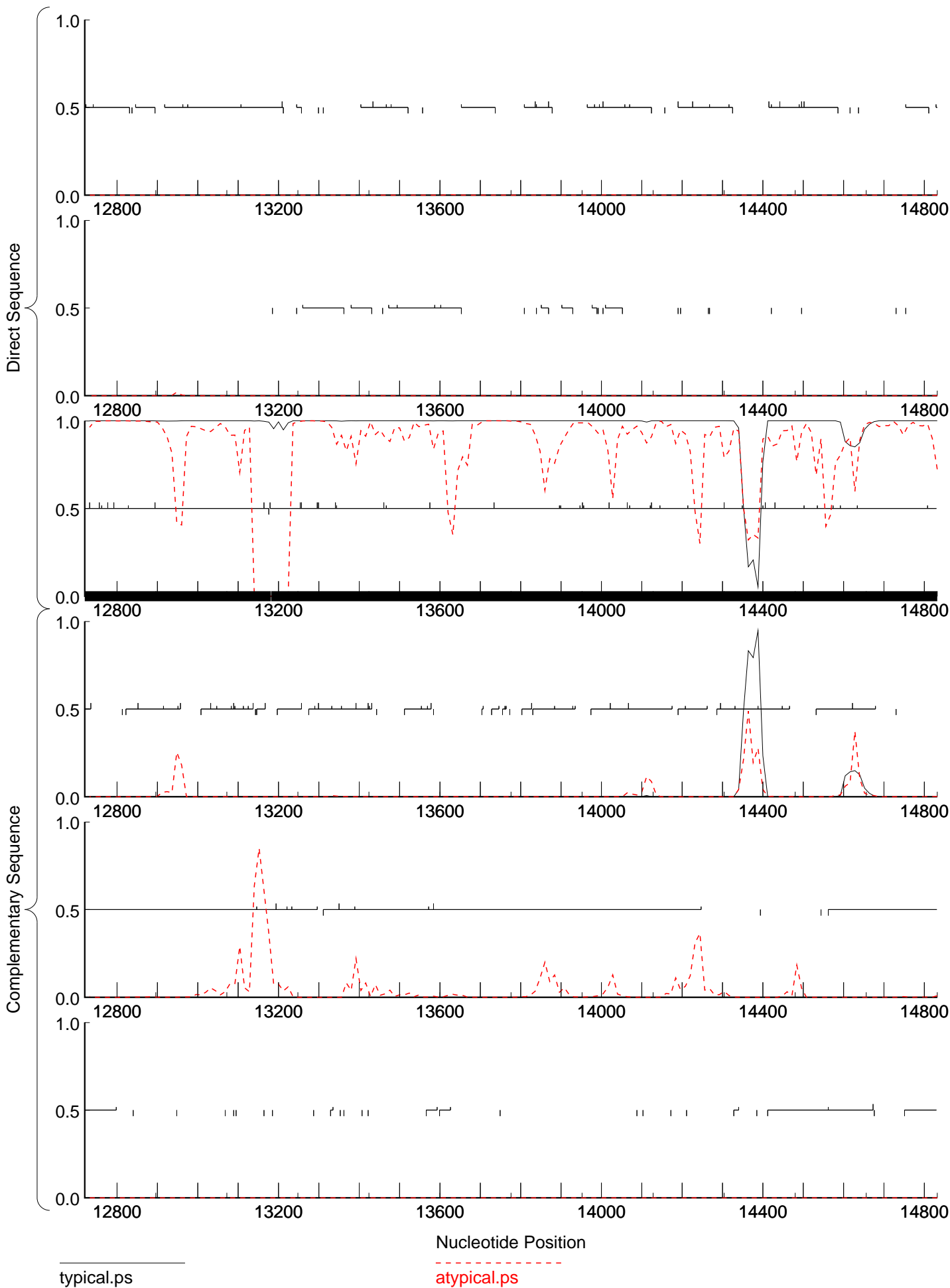


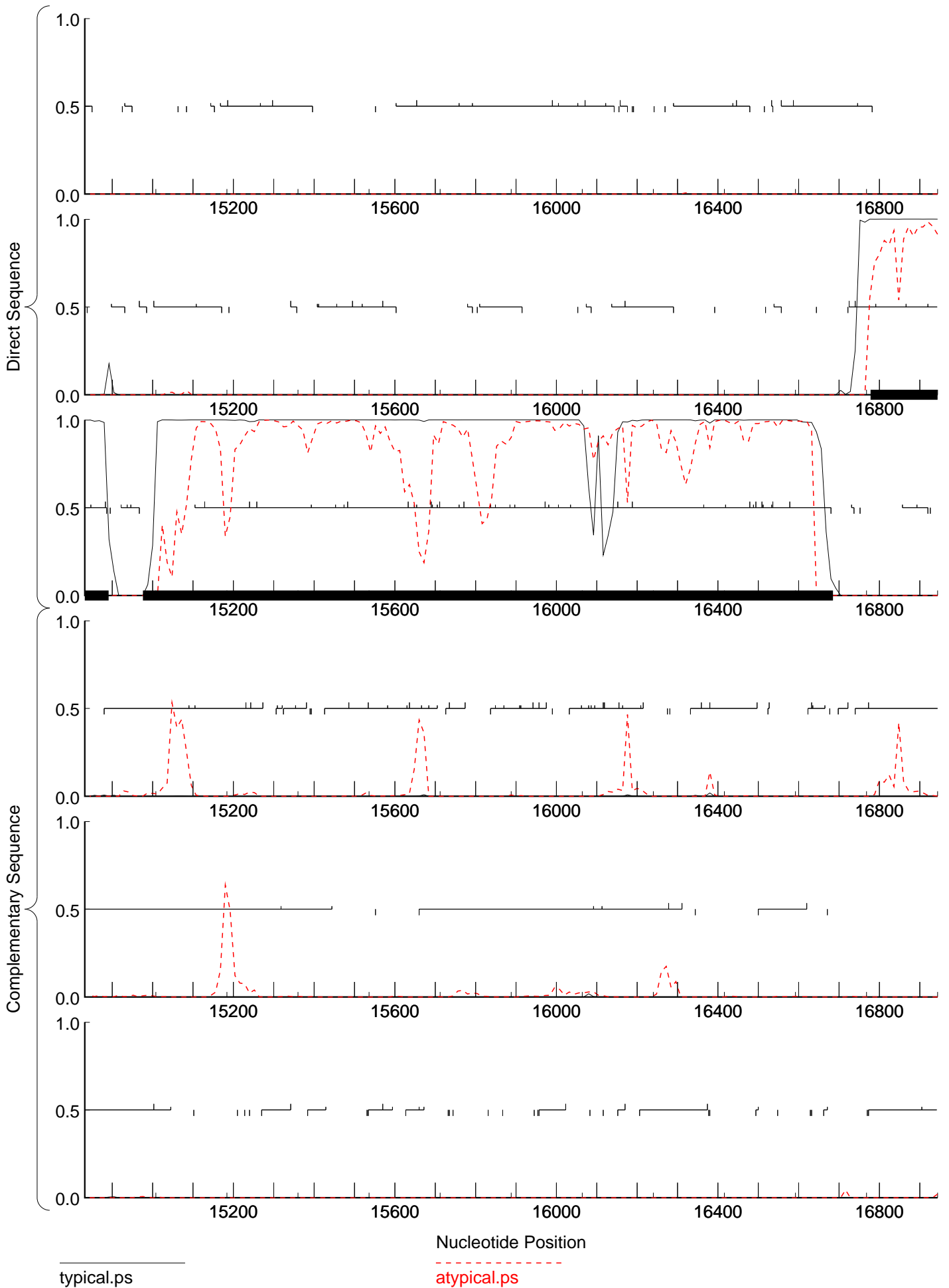
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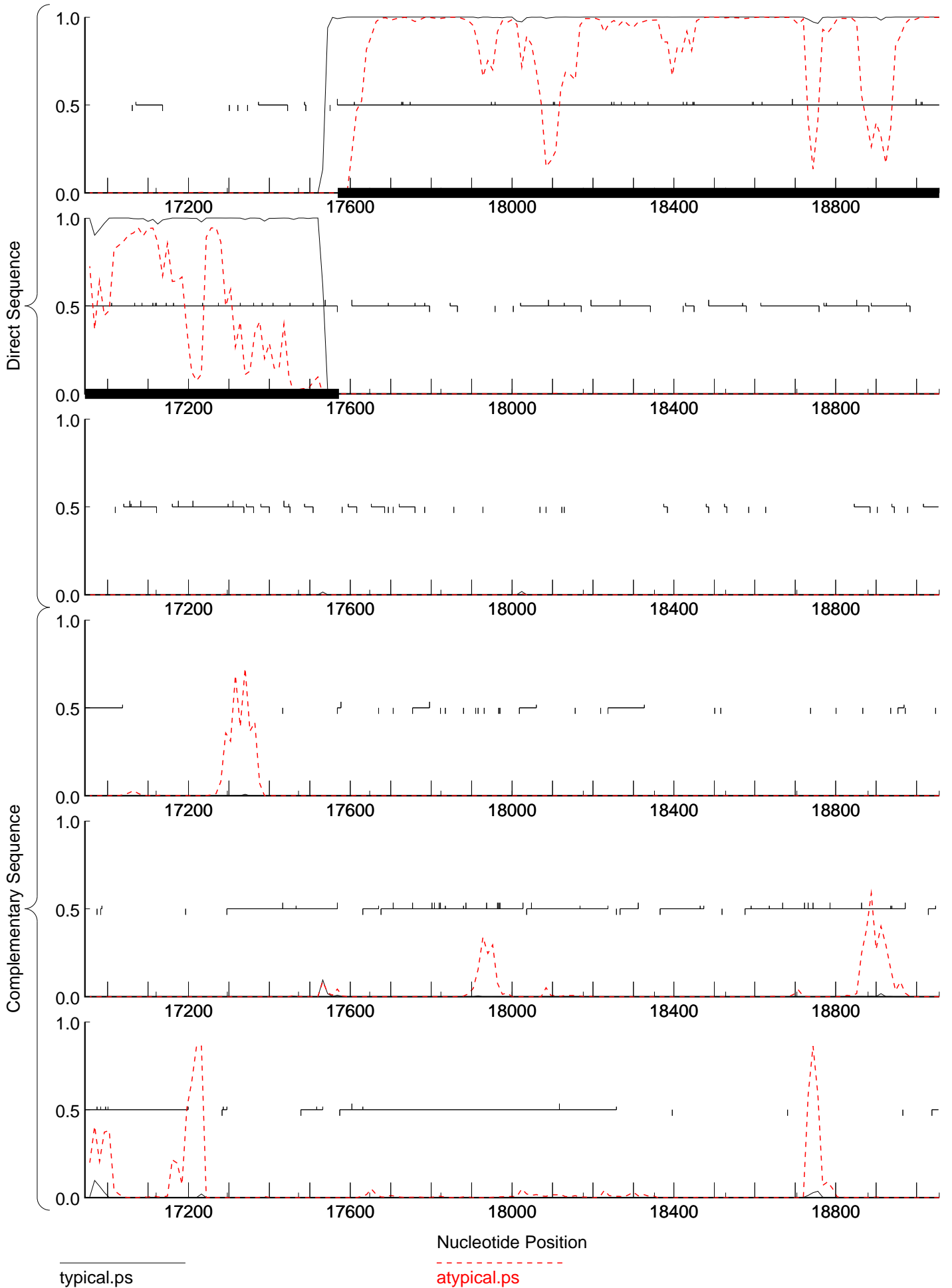
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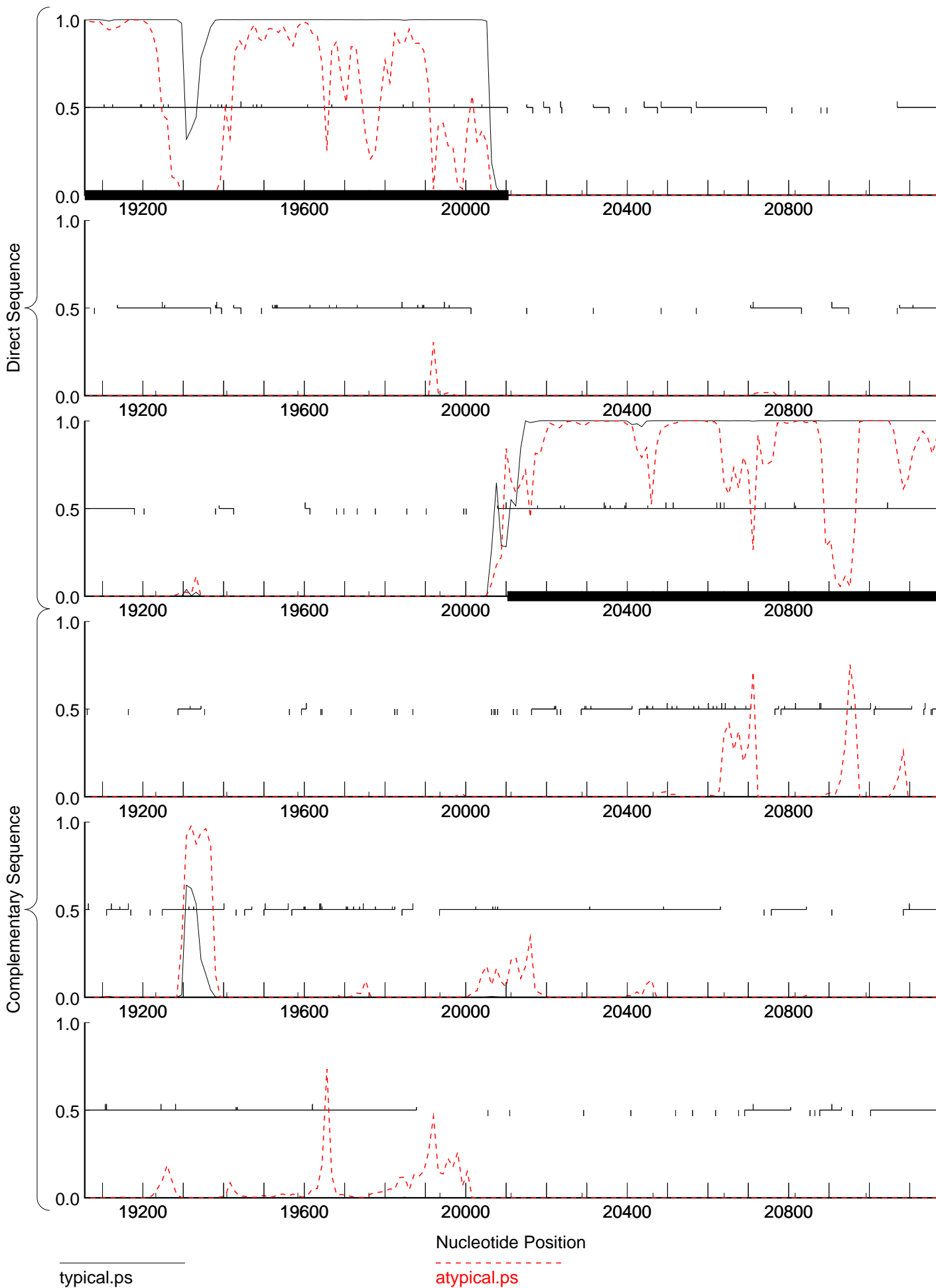


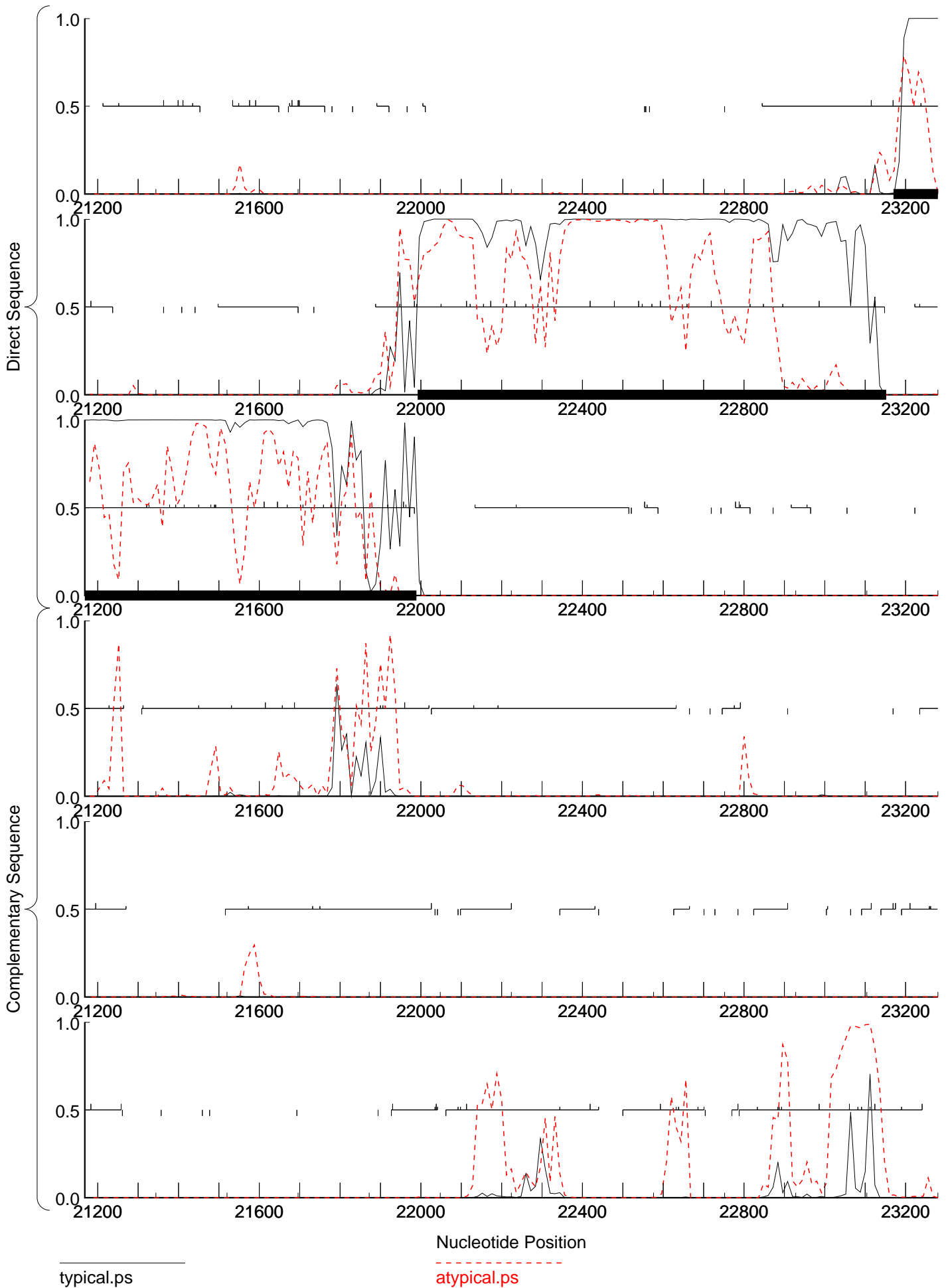


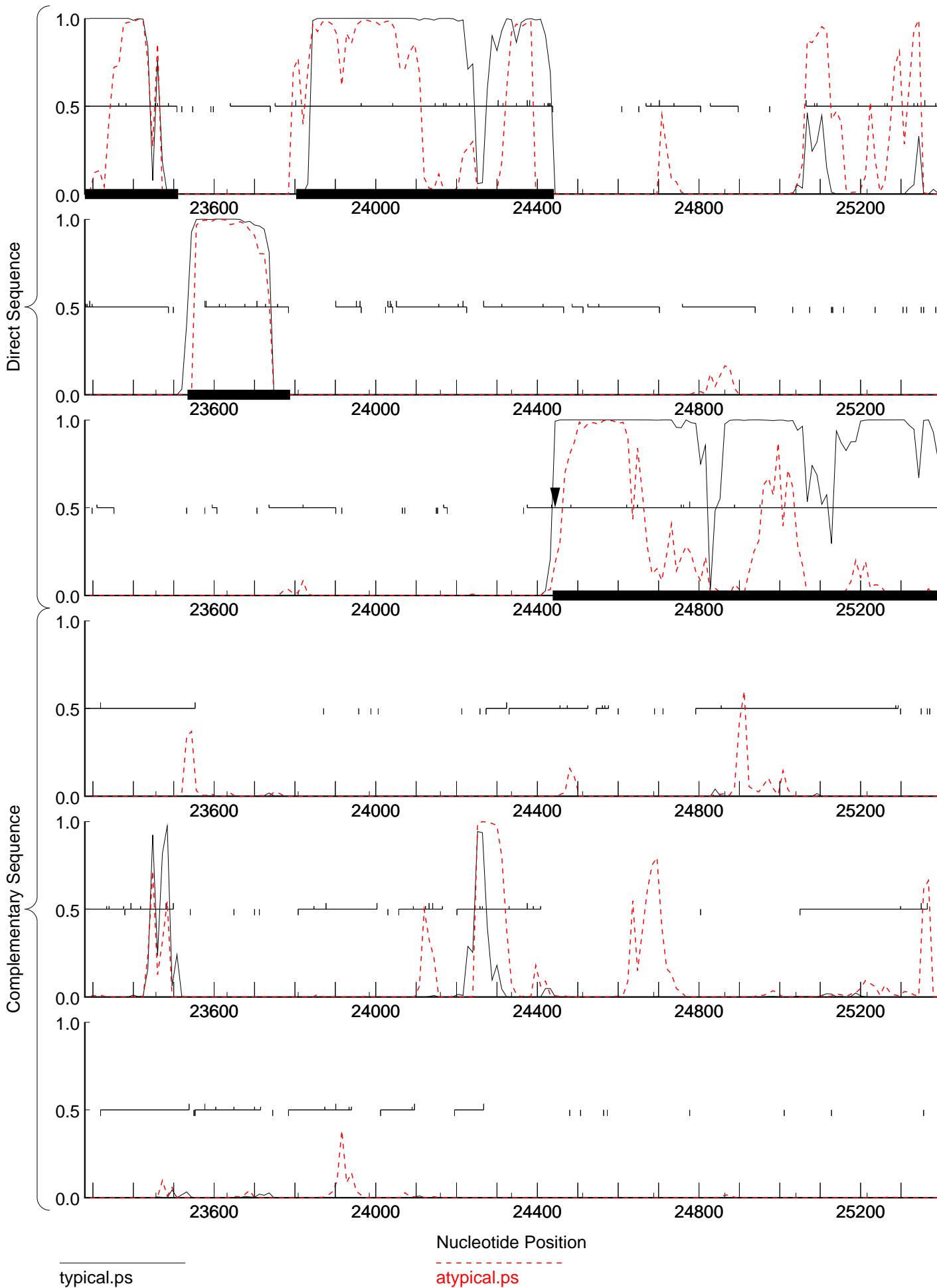






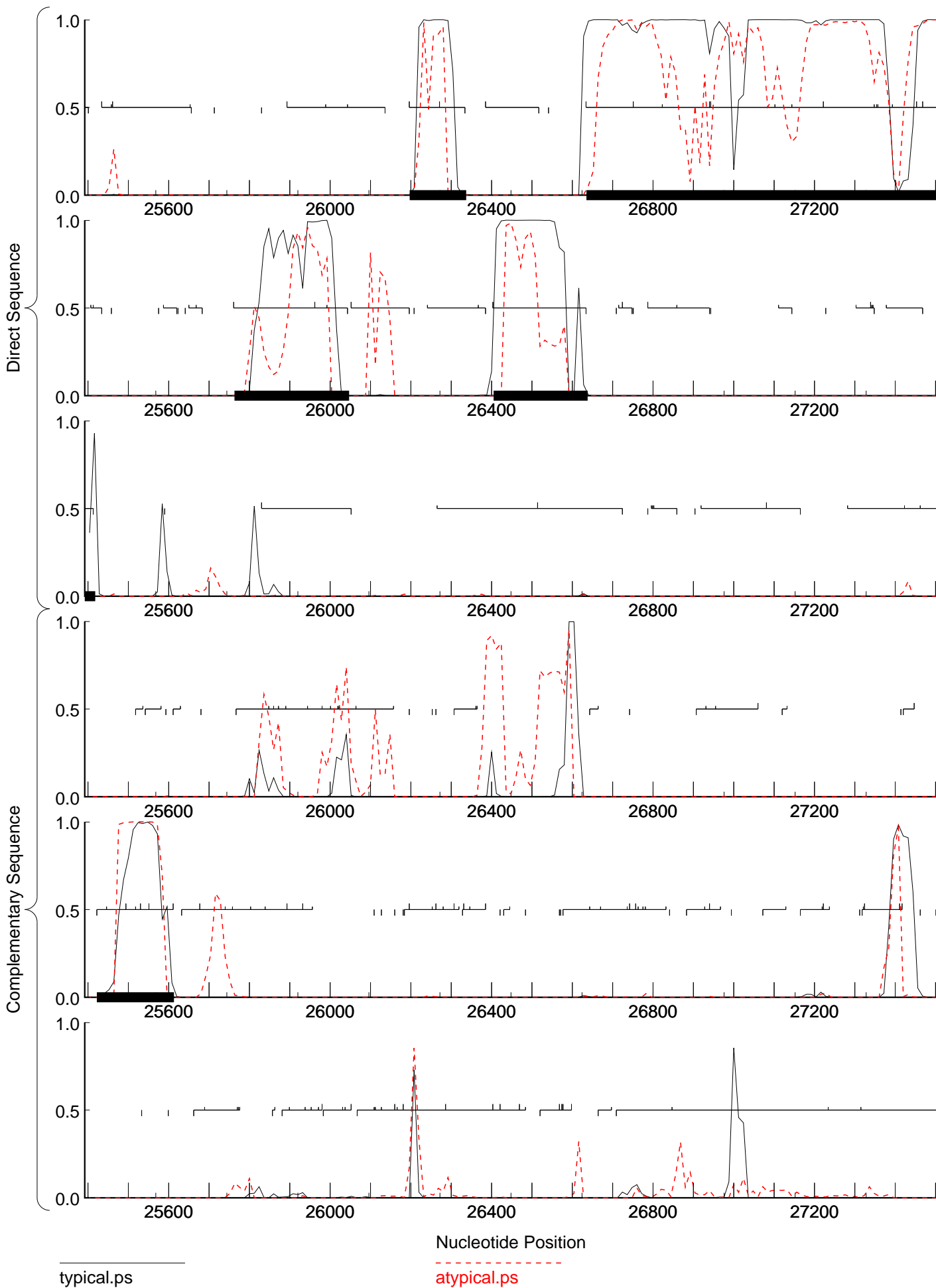


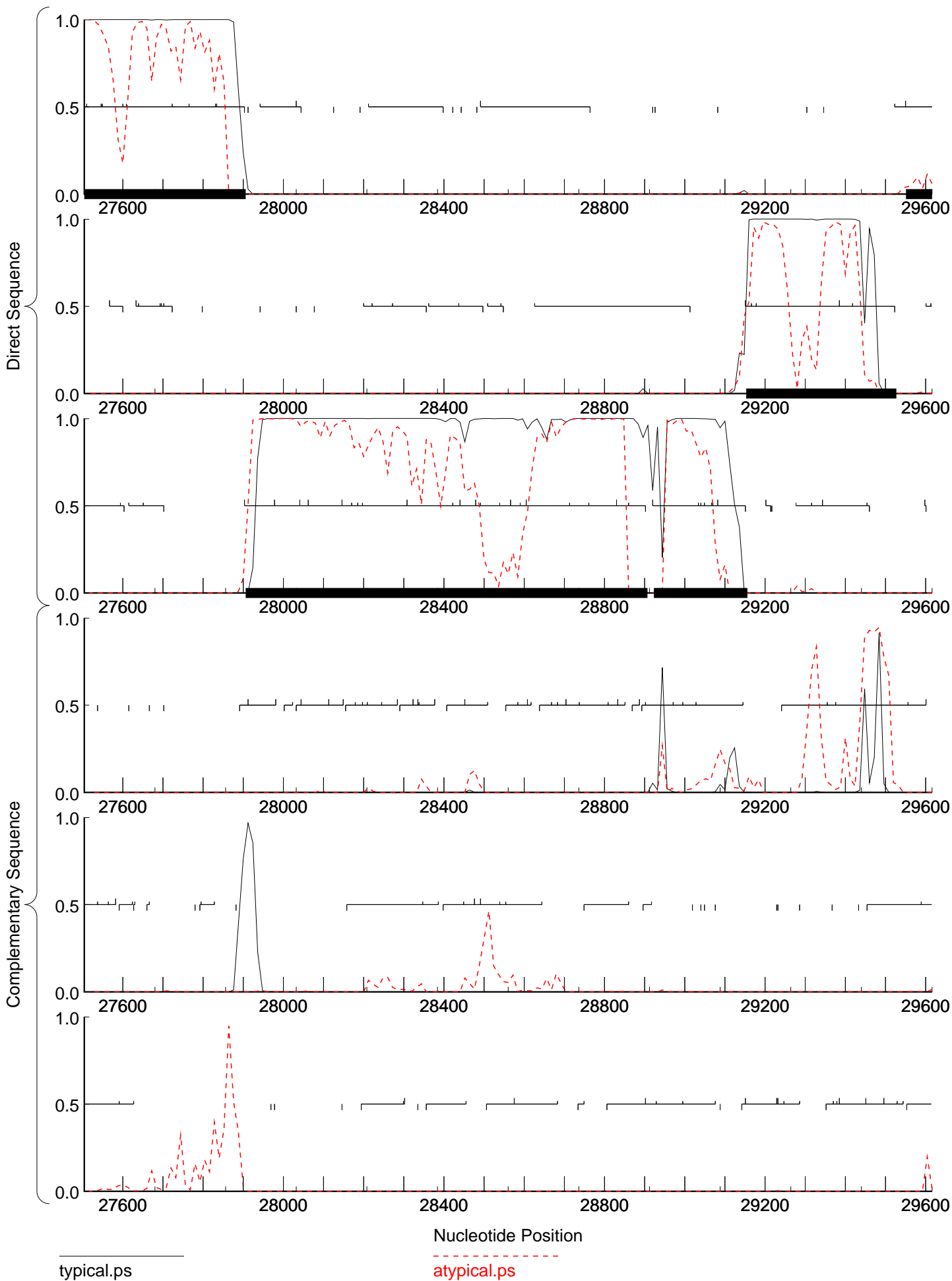


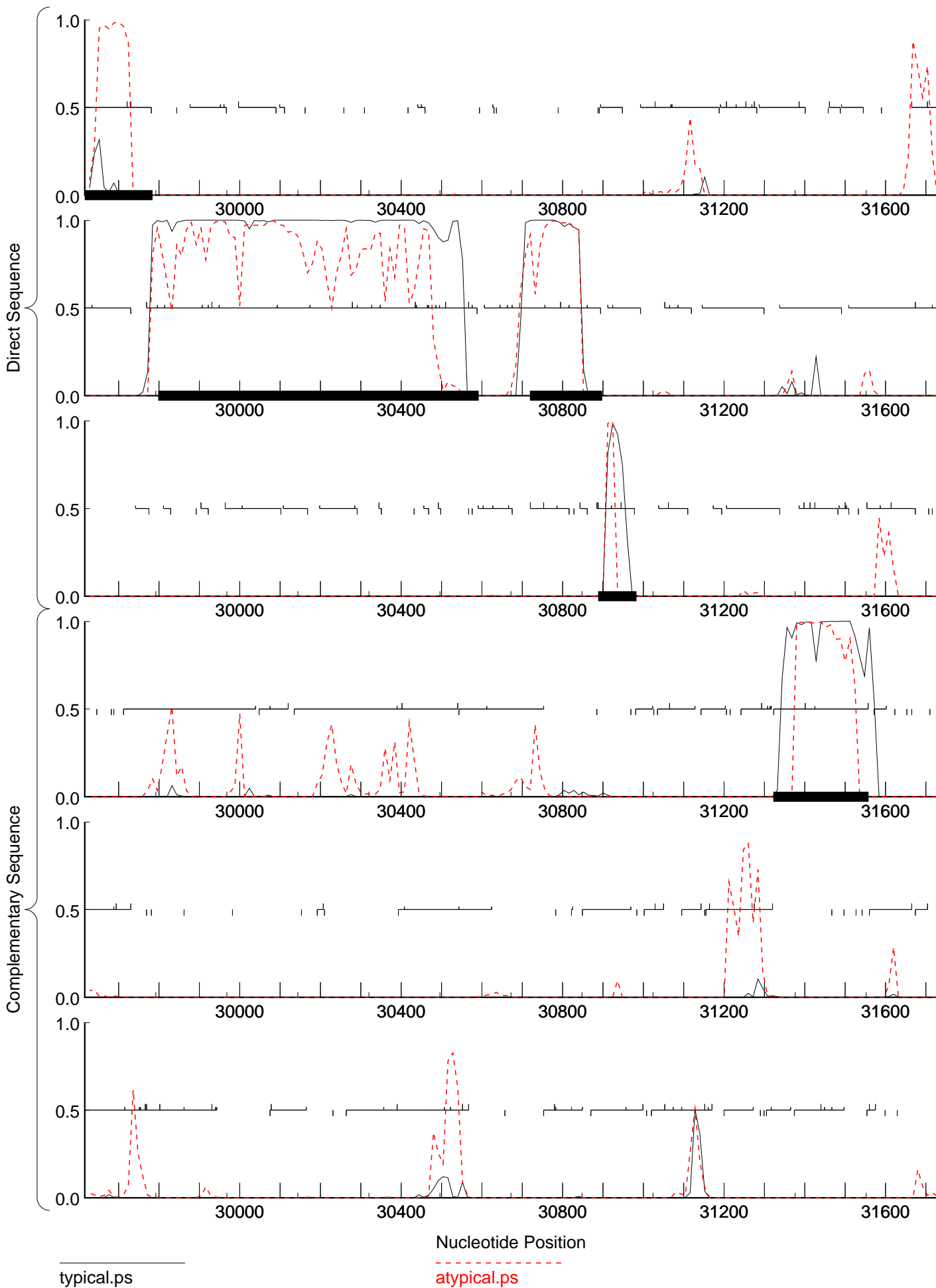


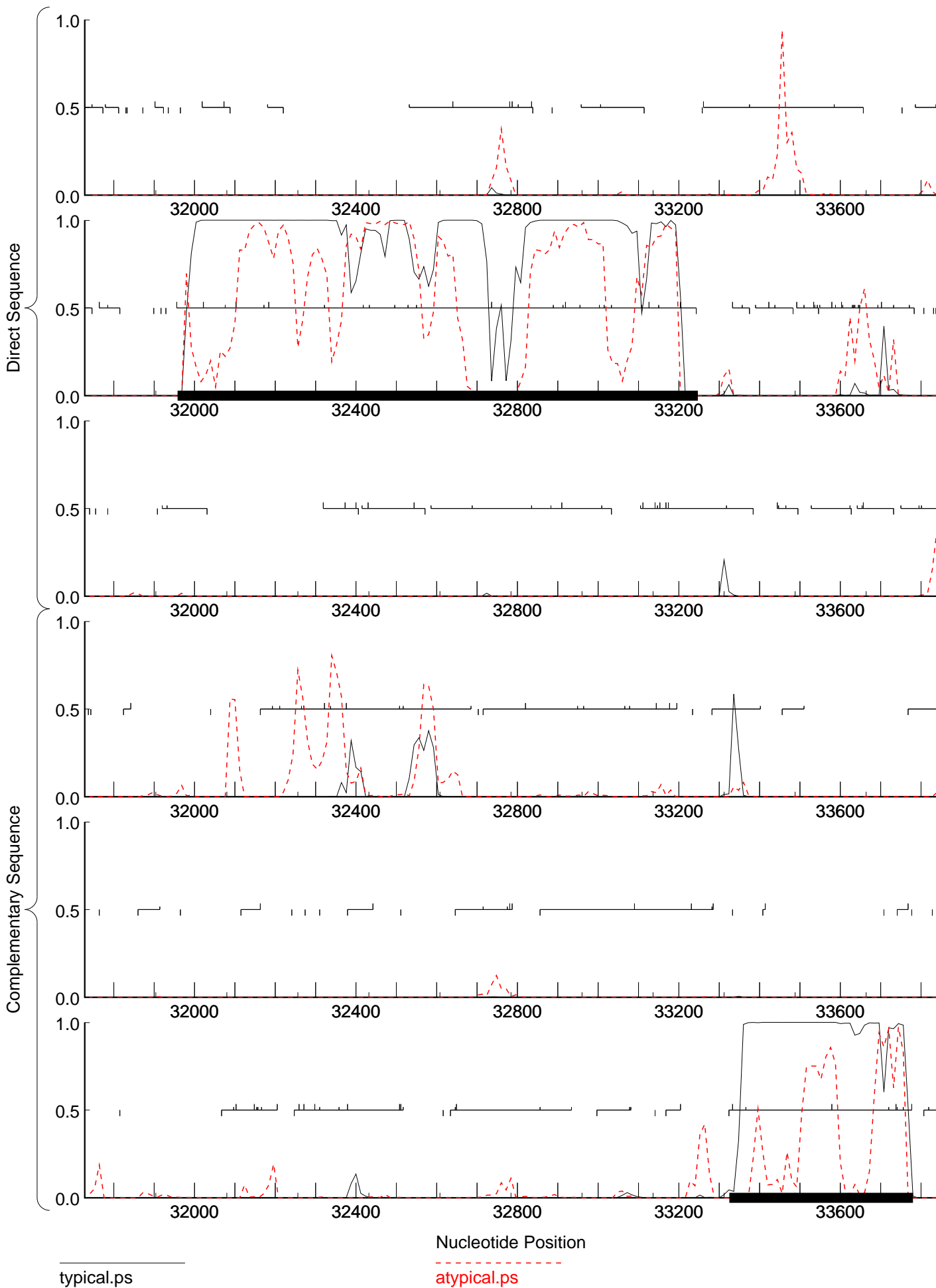
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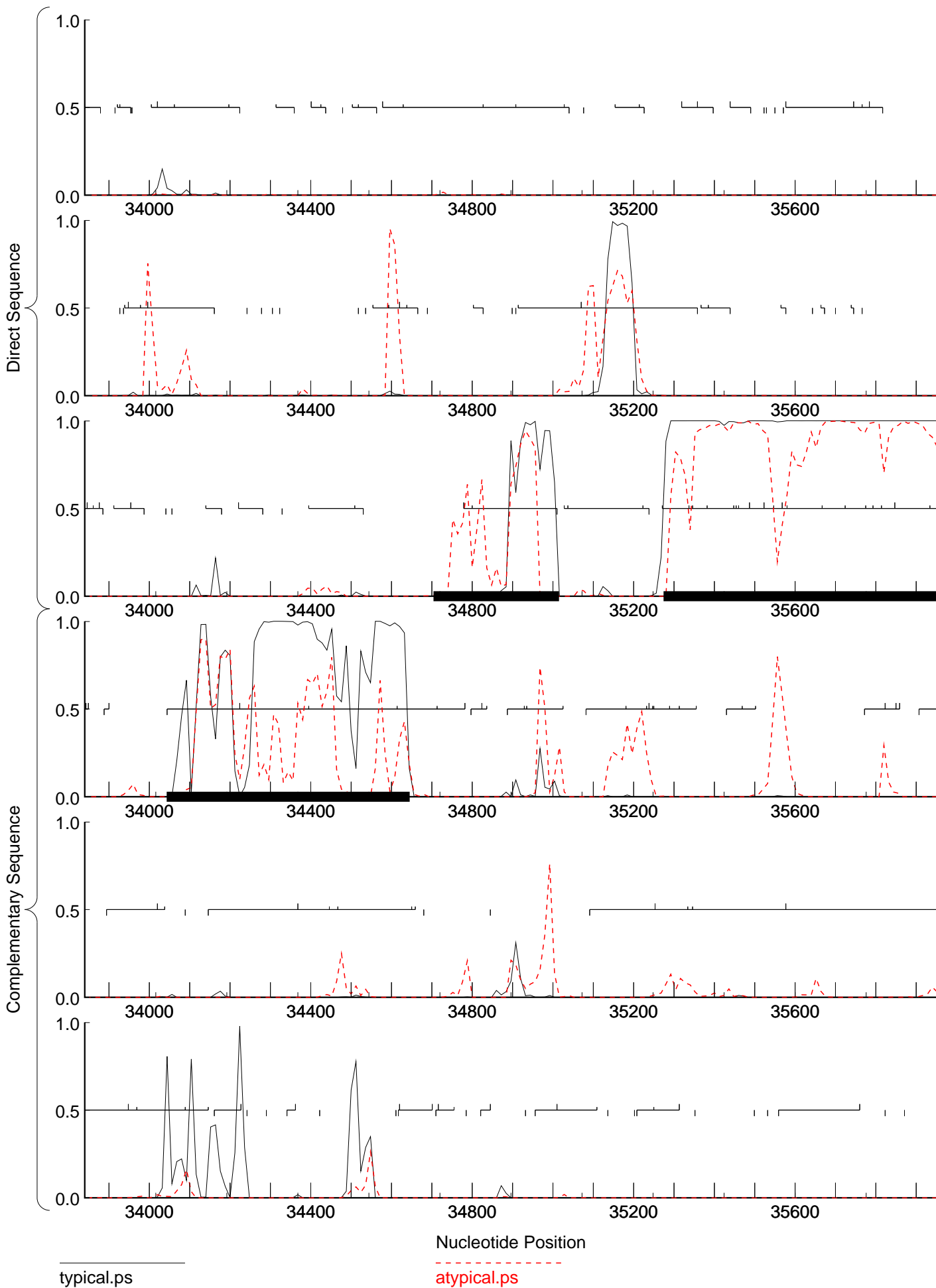
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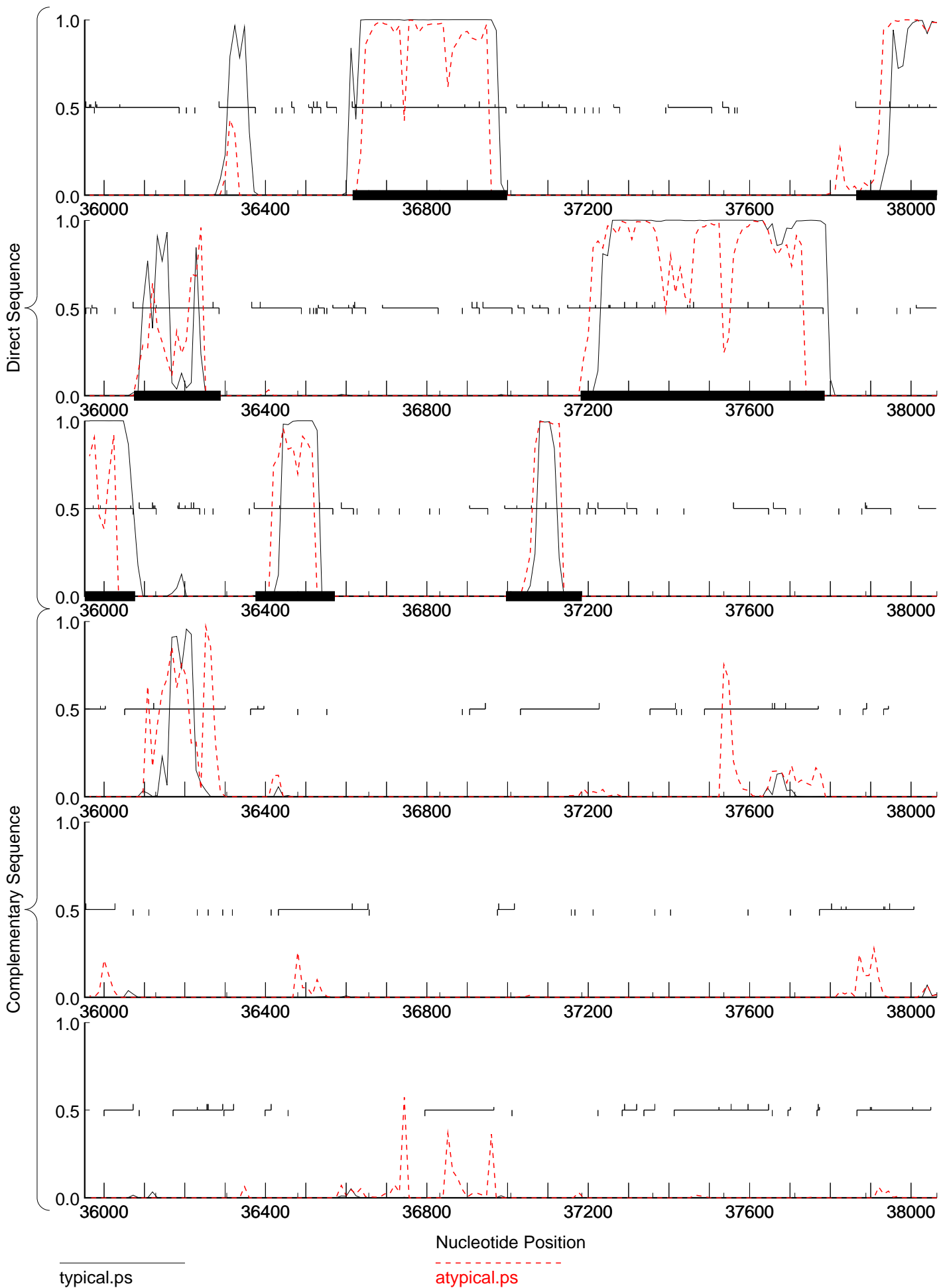






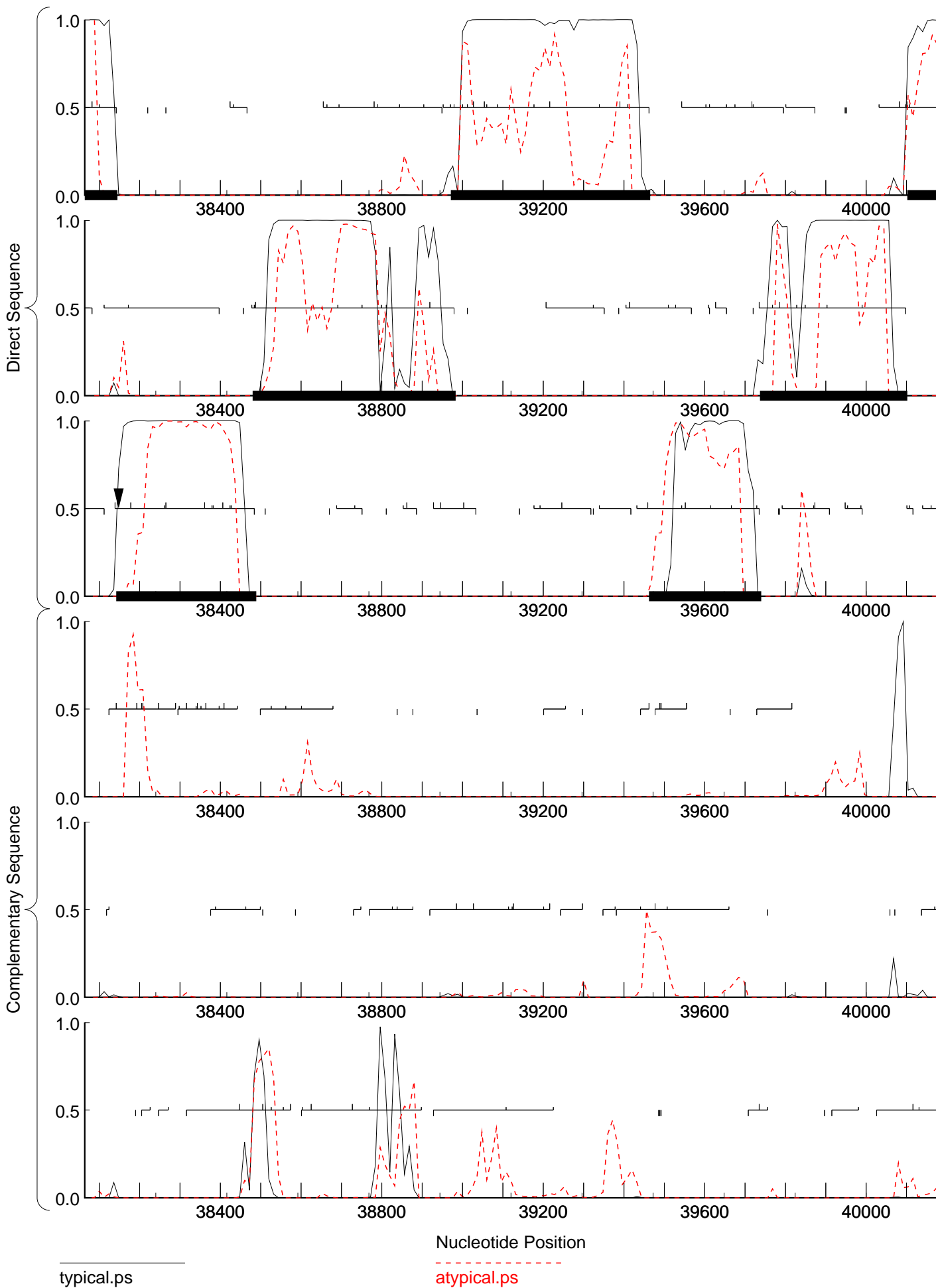


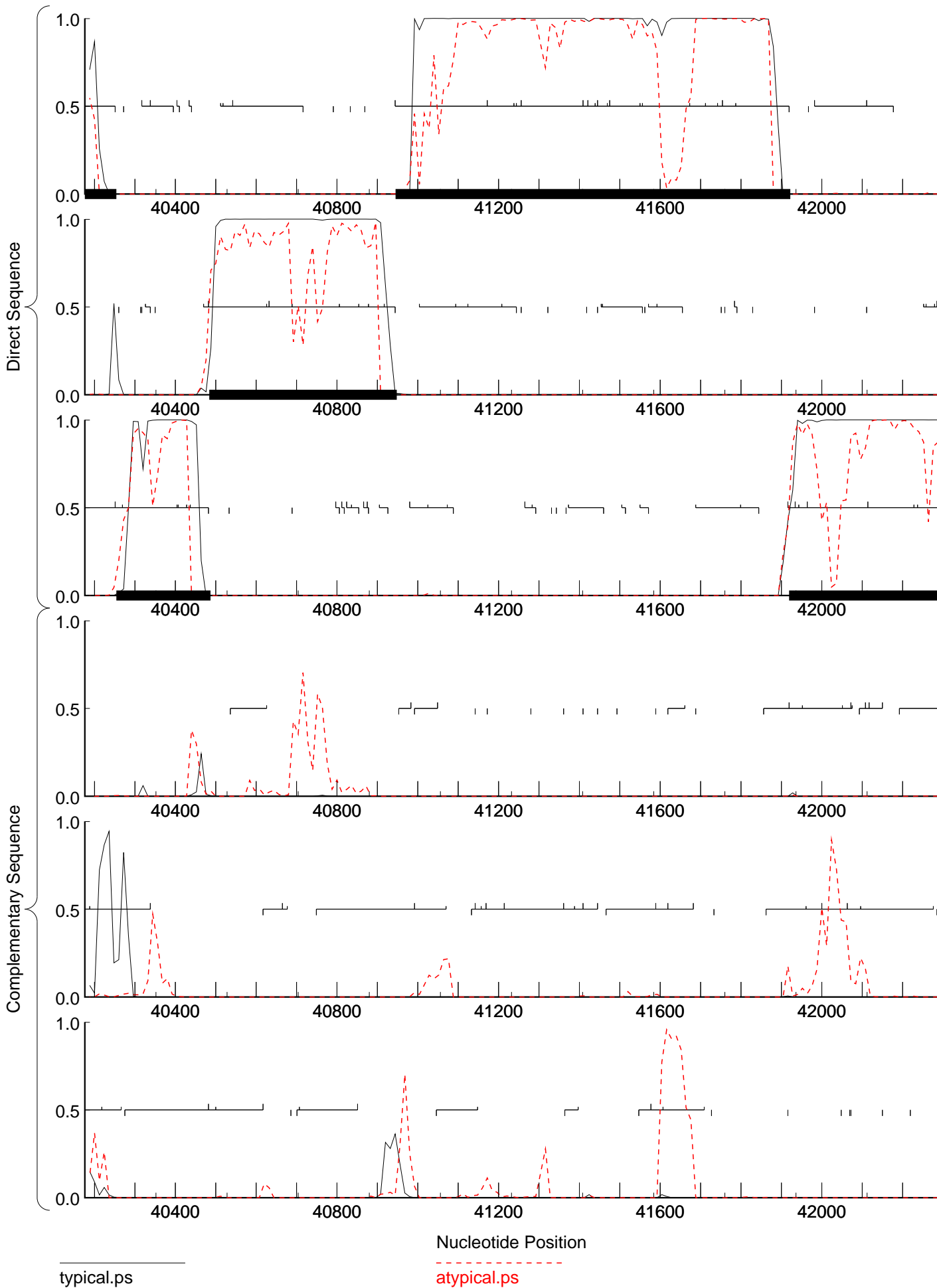


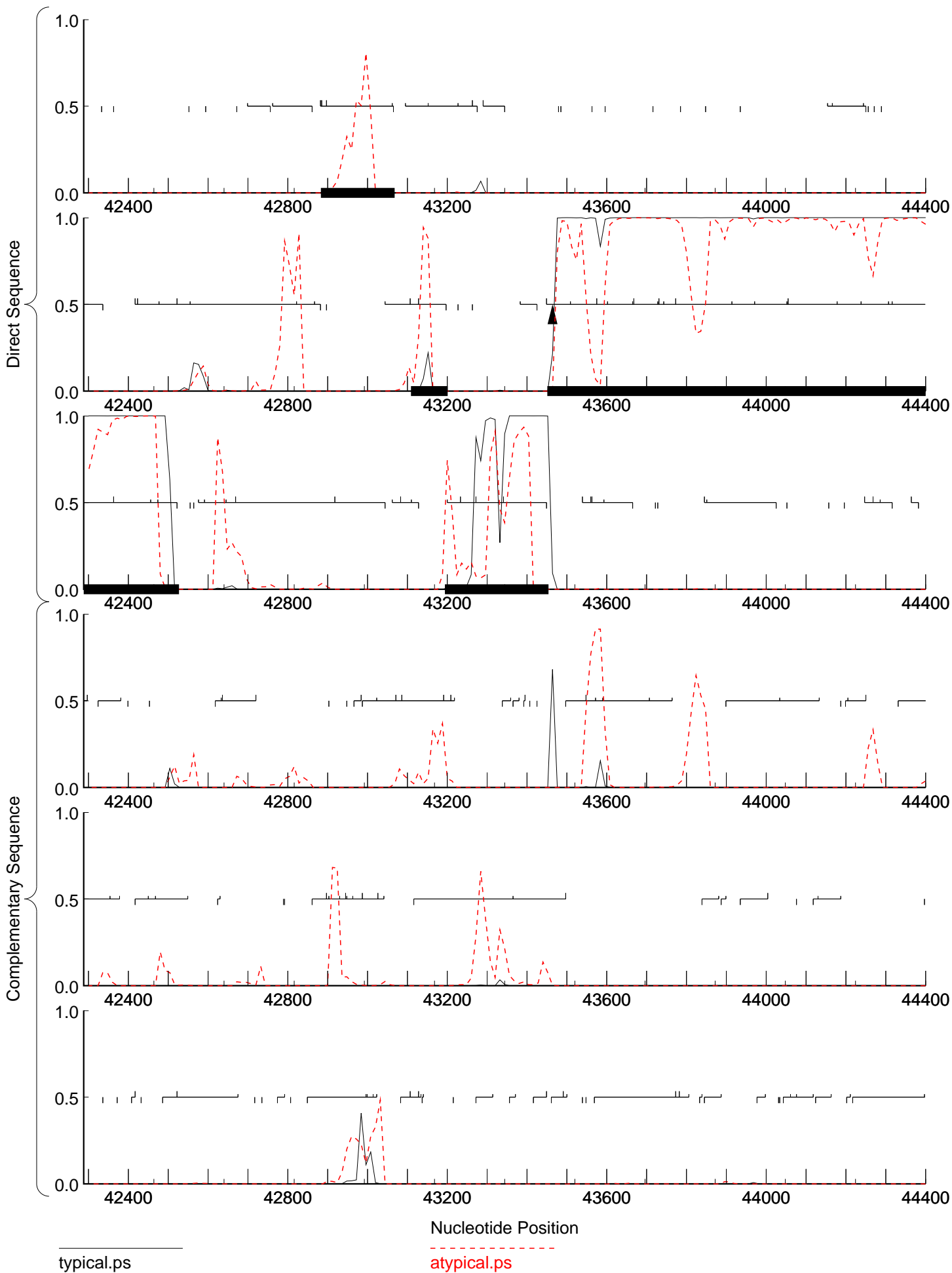


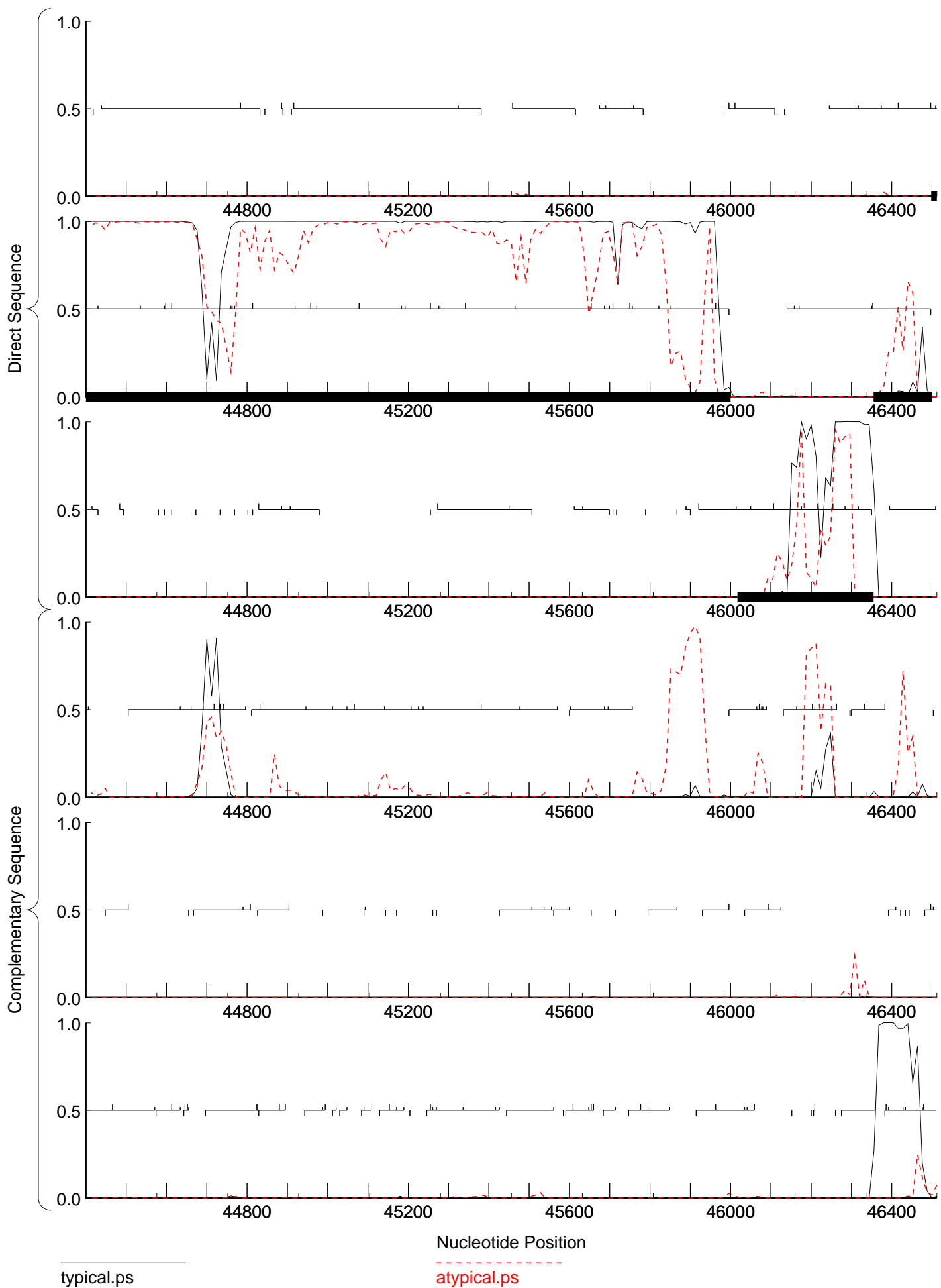
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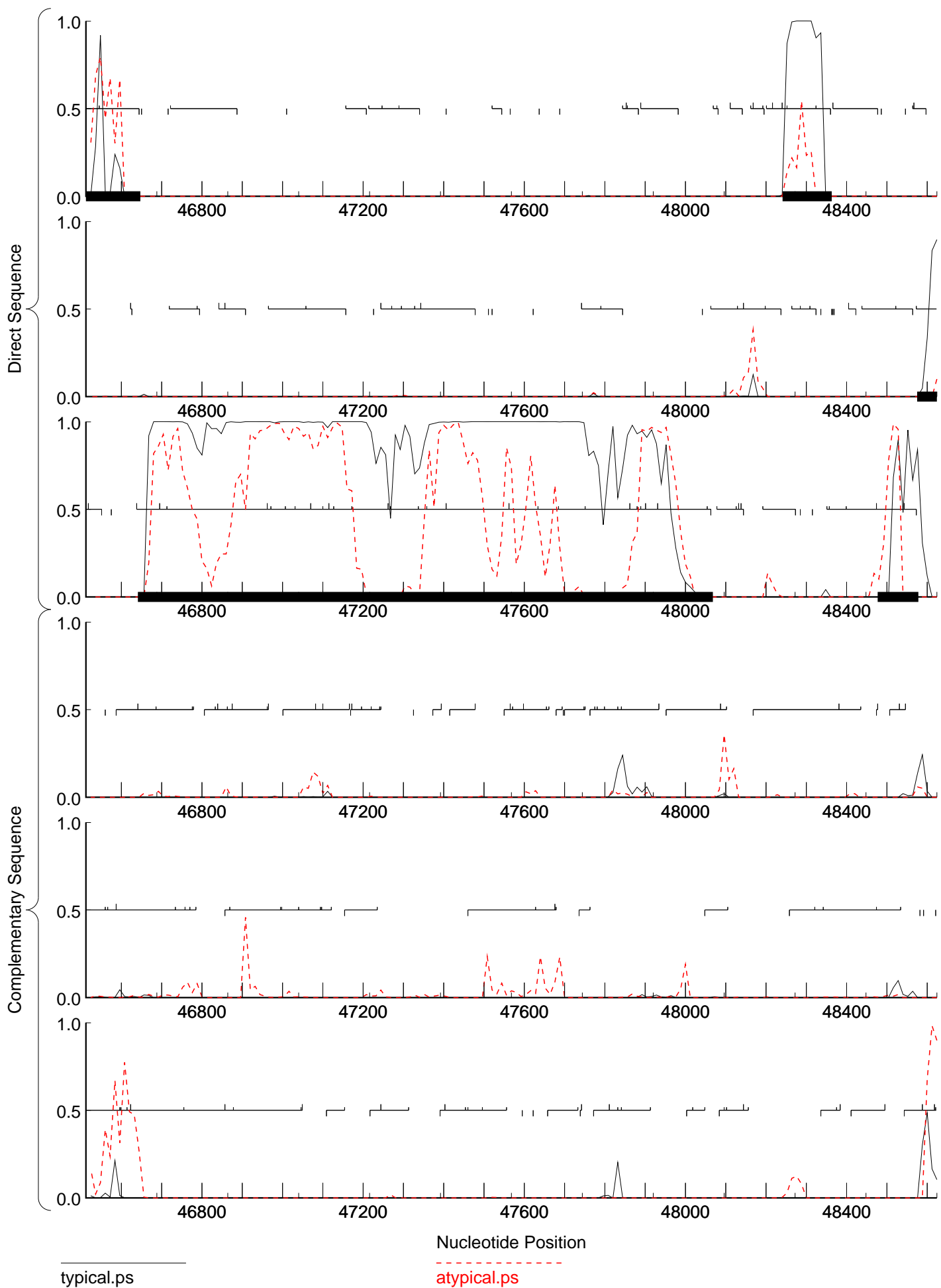
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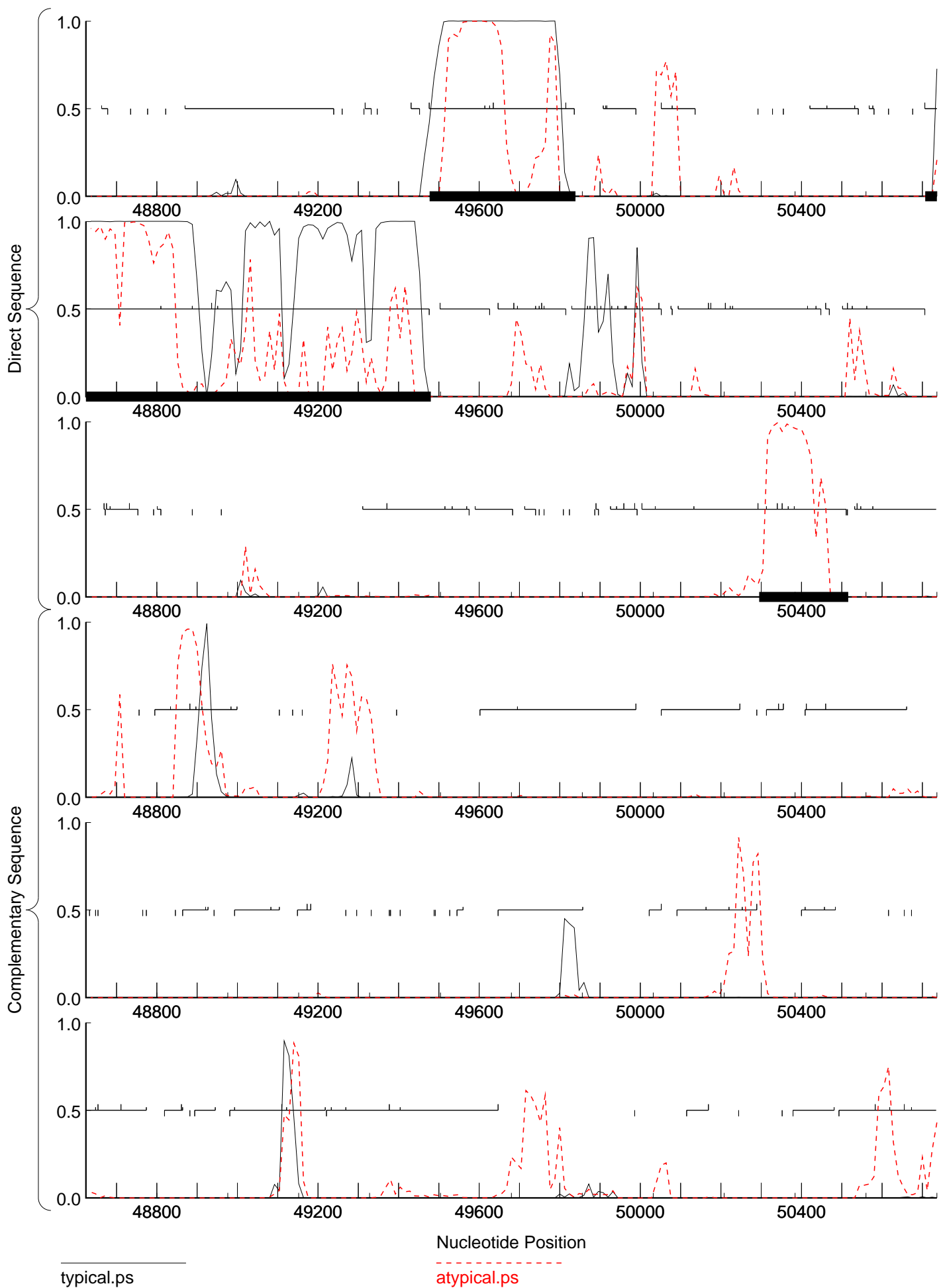


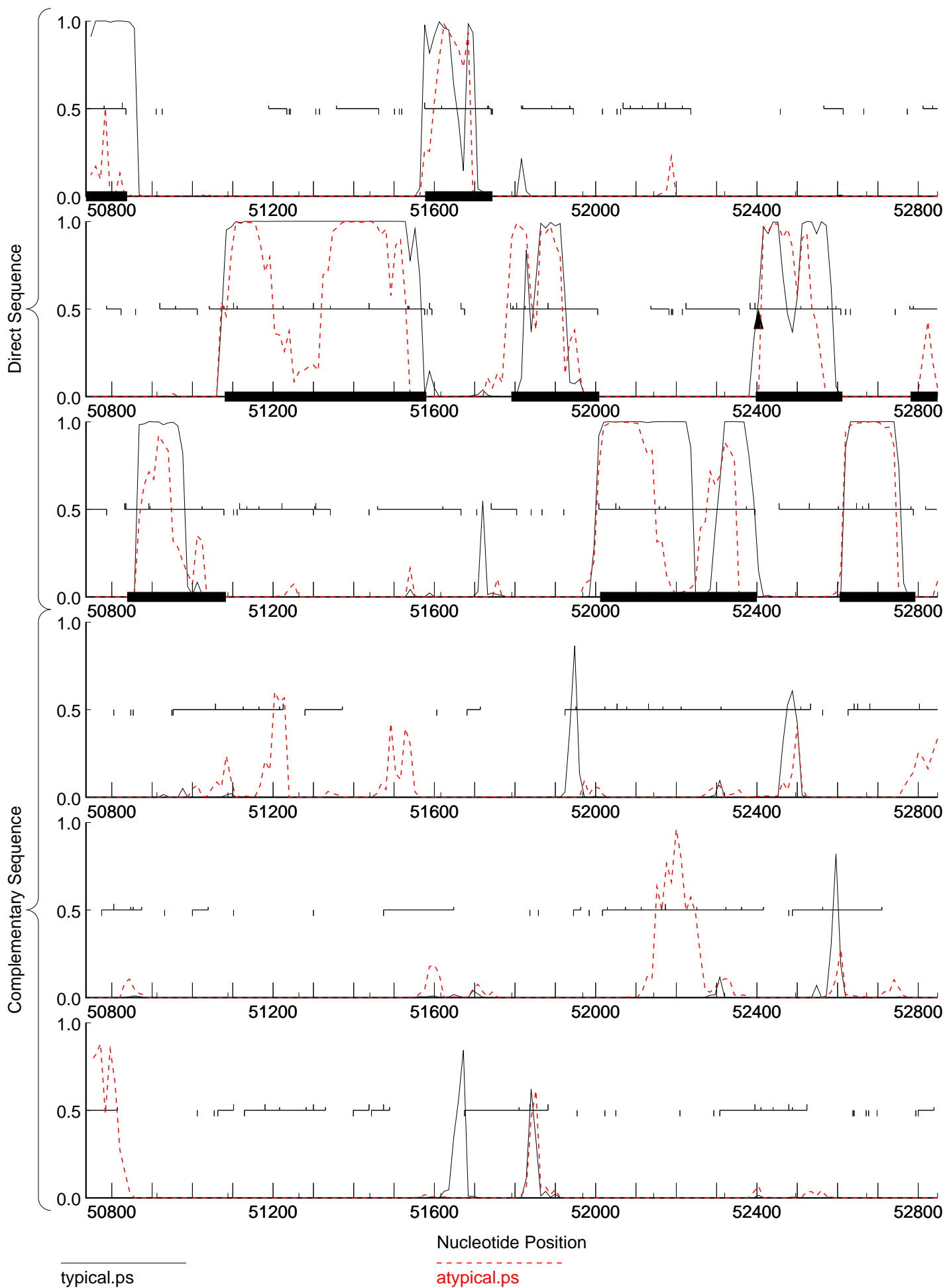


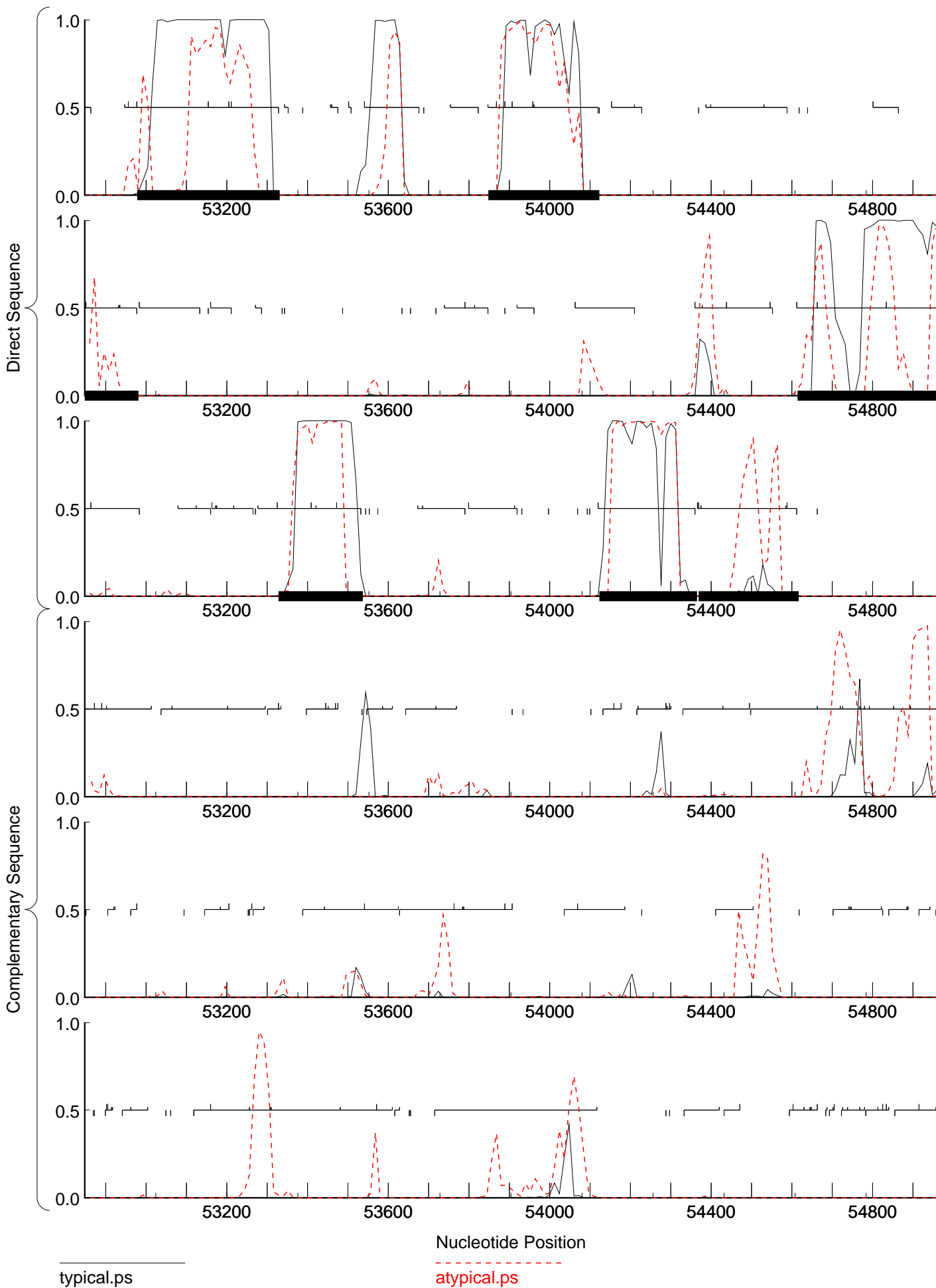


typical.ps

atypical.ps

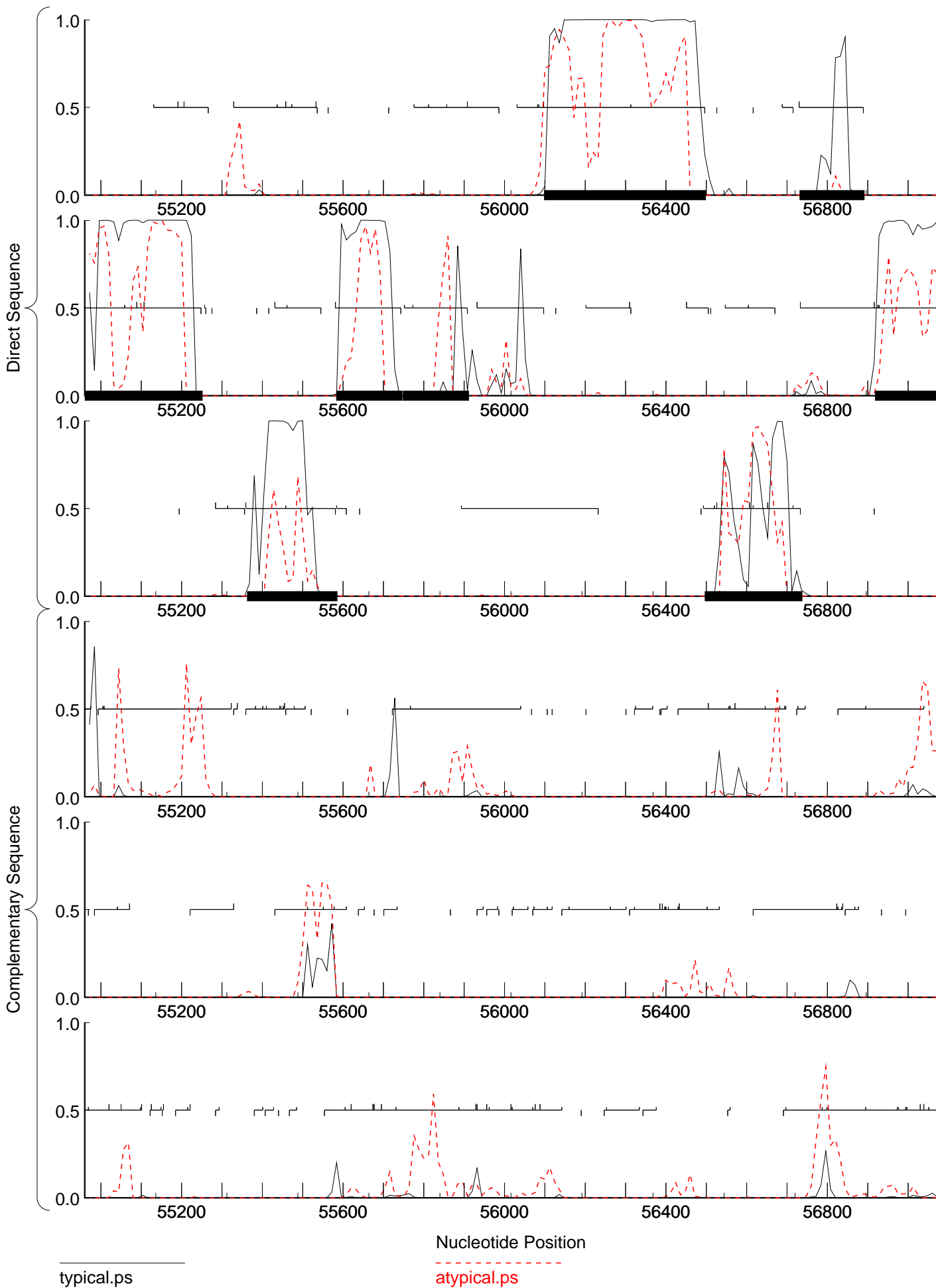


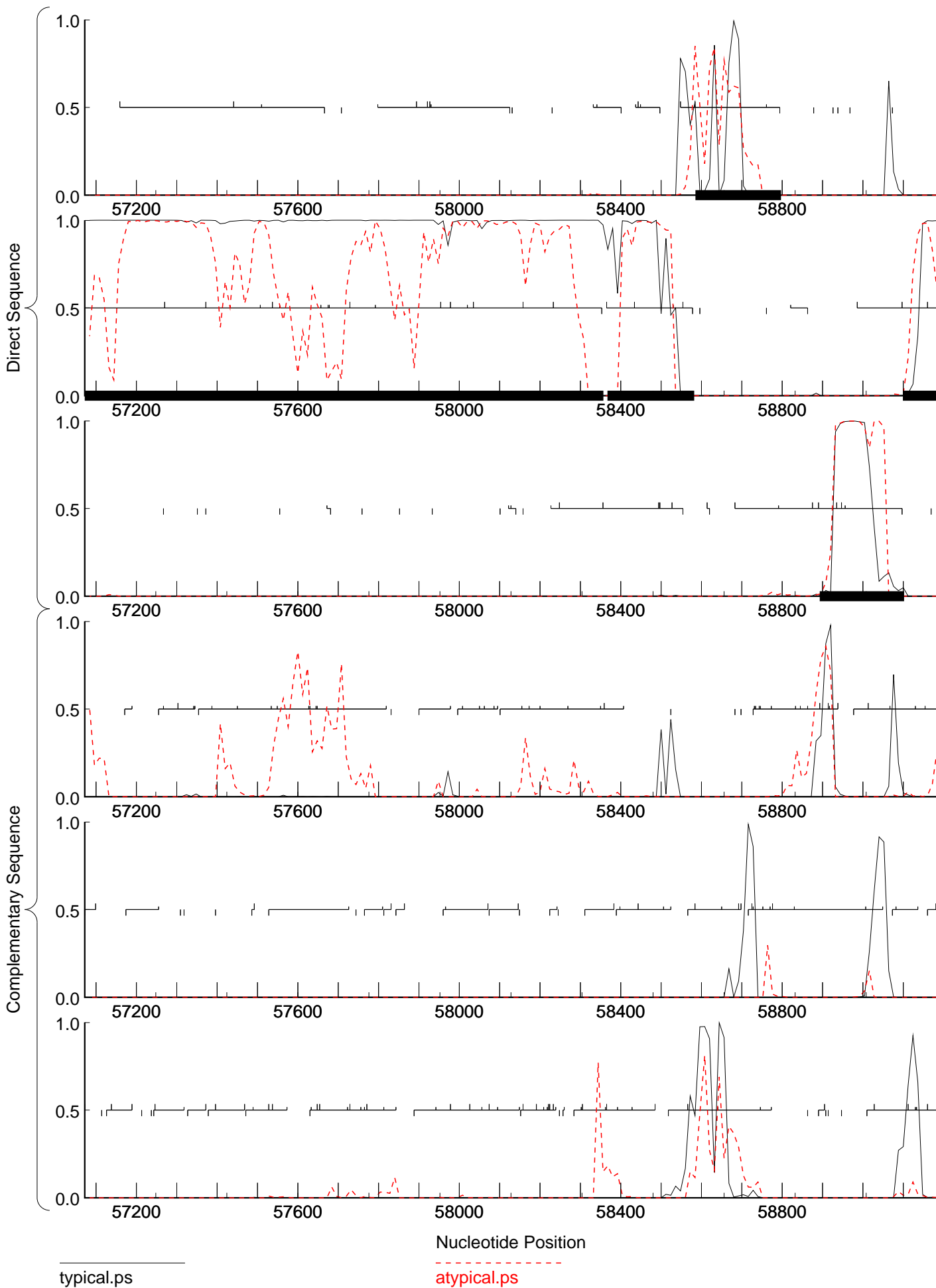


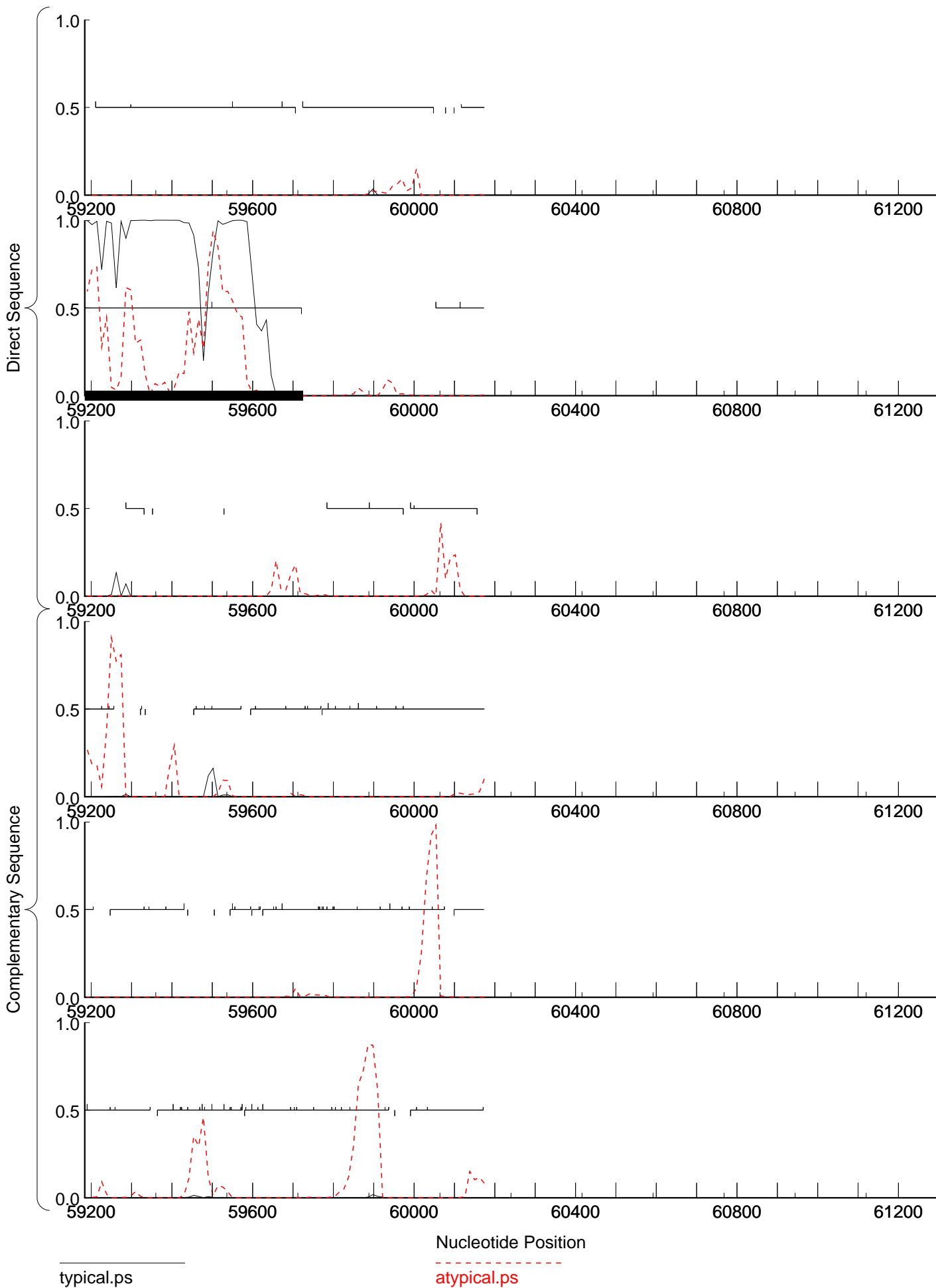


typical.ps

atypical.ps







typical.ps

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