

GeneMark

Version 2.5p (09.08.06)

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PROGRAM INFORMATION

Sequence : Gordonia phage Zirinka complete sequence, 52077 bp including 10-base 3' overhang (CGGCTGGGG
Analysis Date : 1/18/18 at 20:17:58
Pages : 26
Sequence Length : 52077 bp
GC Content : 66.67%

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,

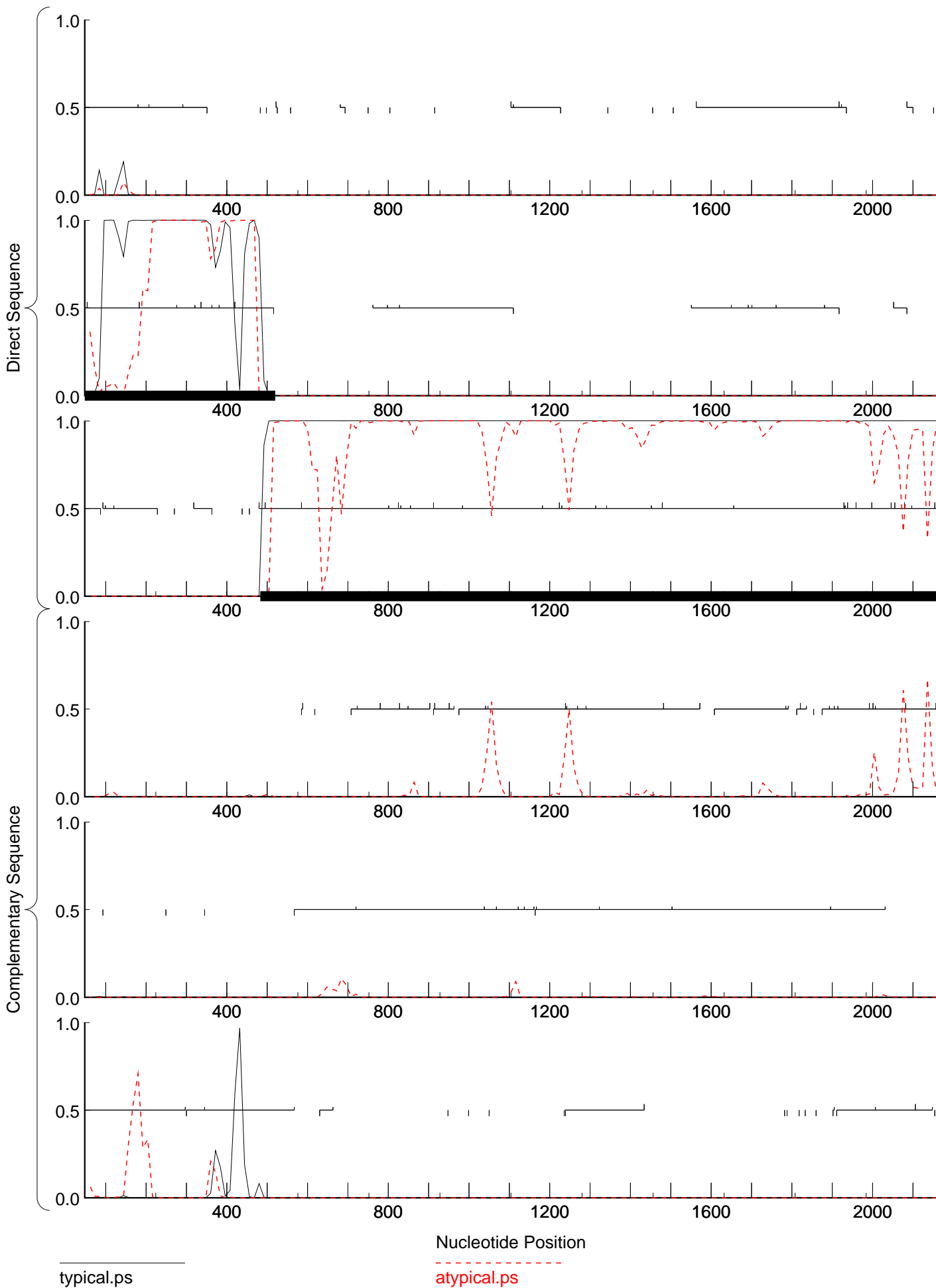
Matrix notes & comments

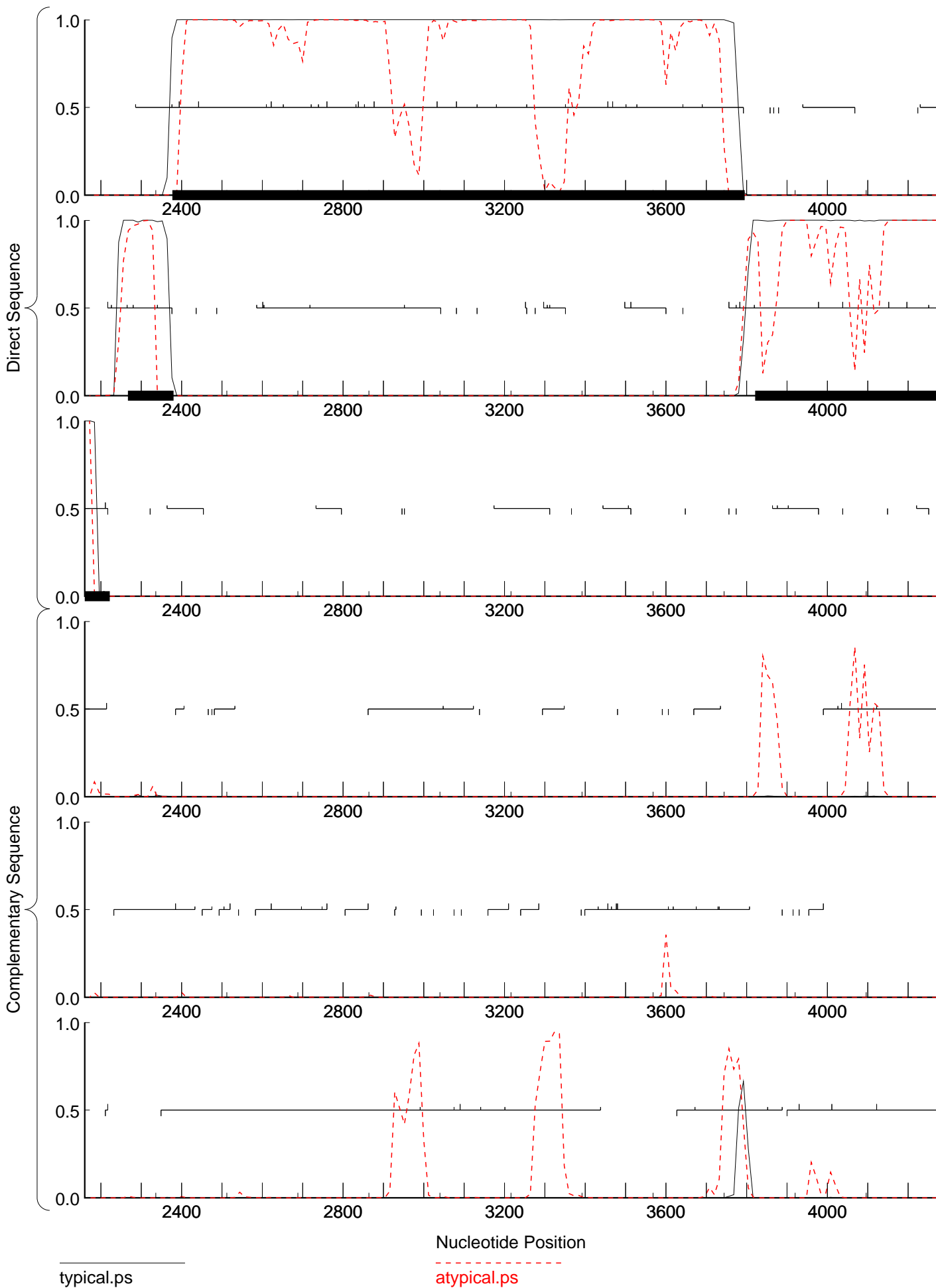
Training set derived by GeneMarkS, 4.30 October 2014
Thu Jan 18 20:17:58 2018

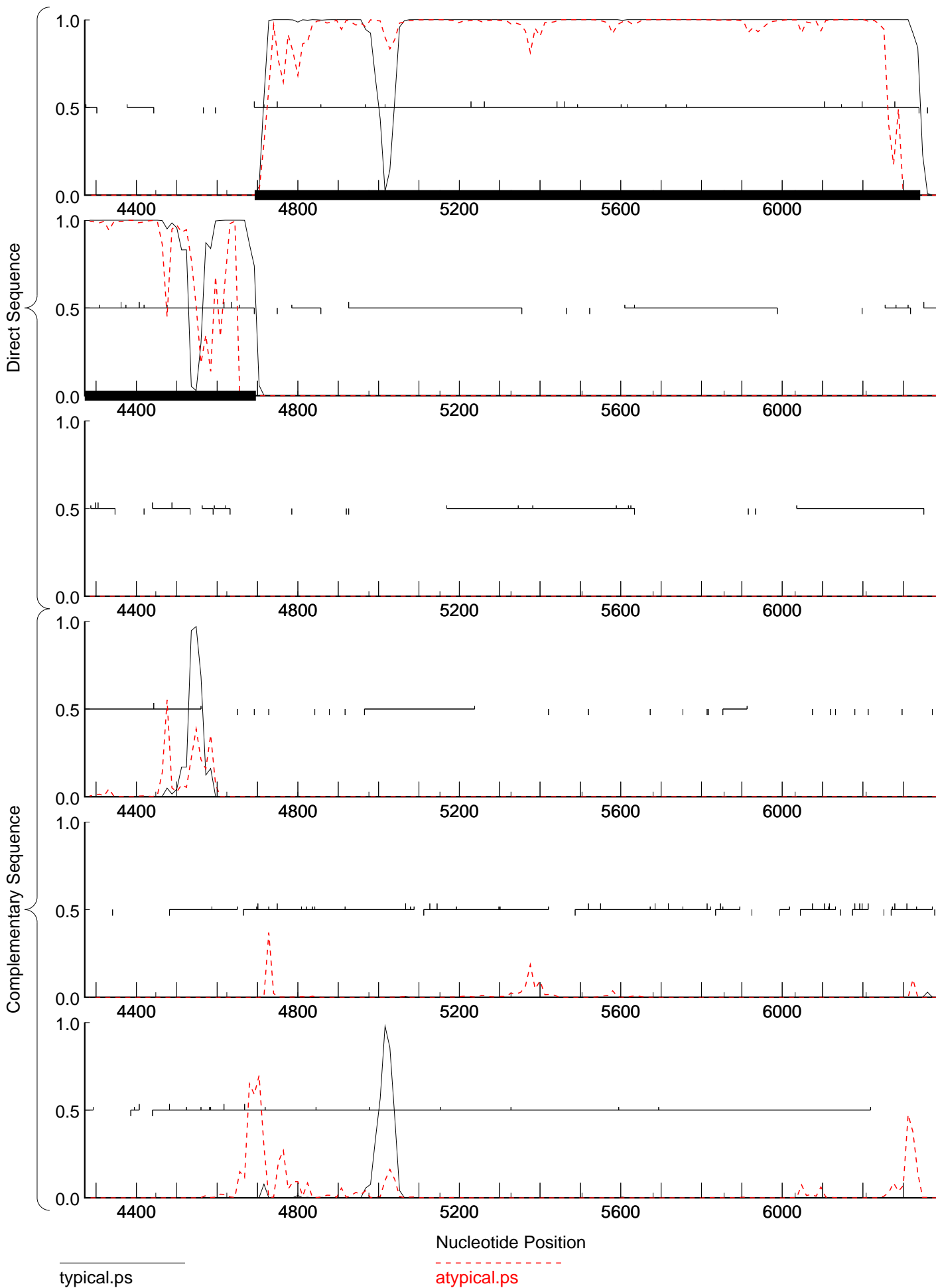
MATRIX INFORMATION

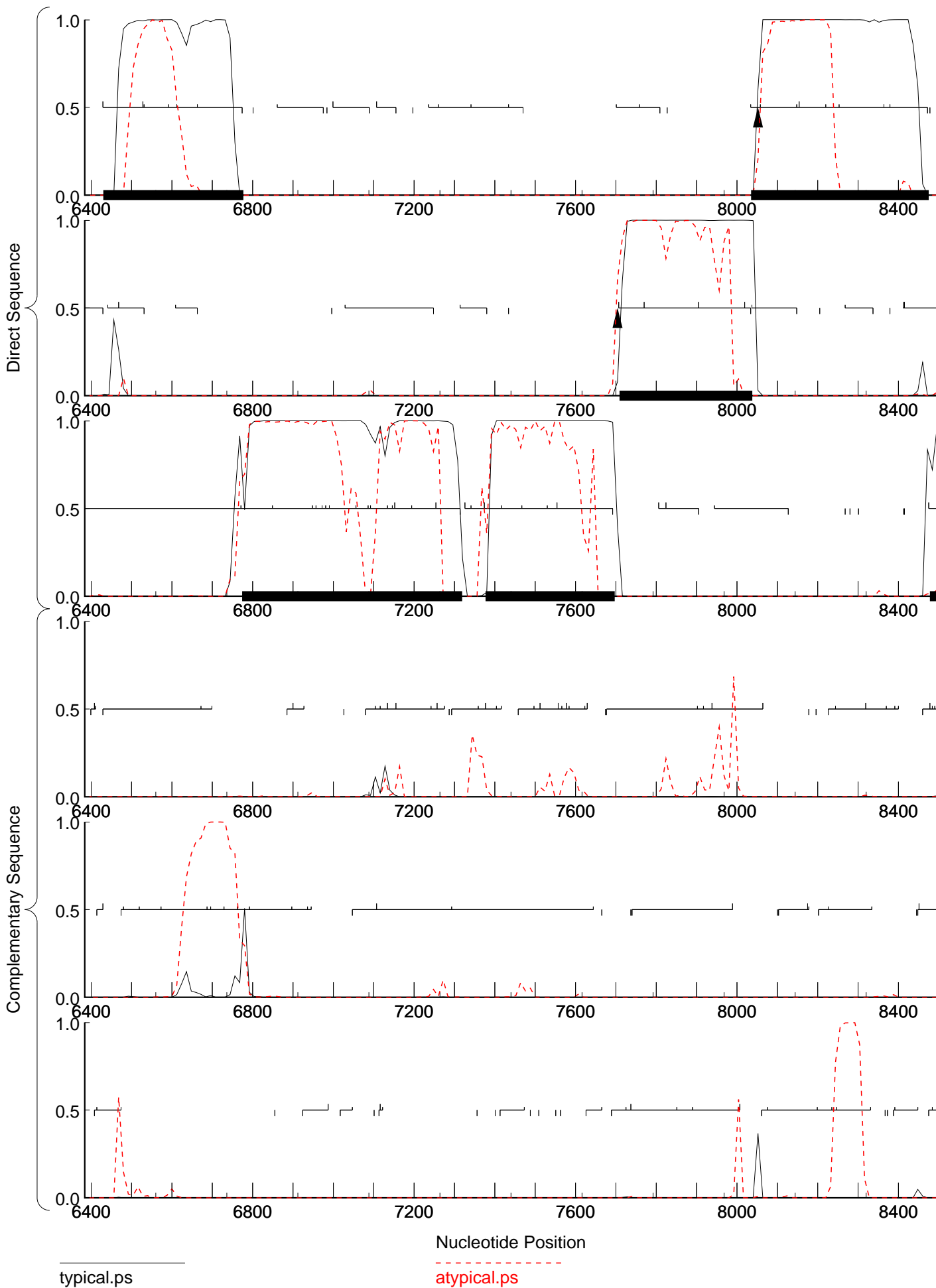
Matrix : -
Author : -
Order : 4

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

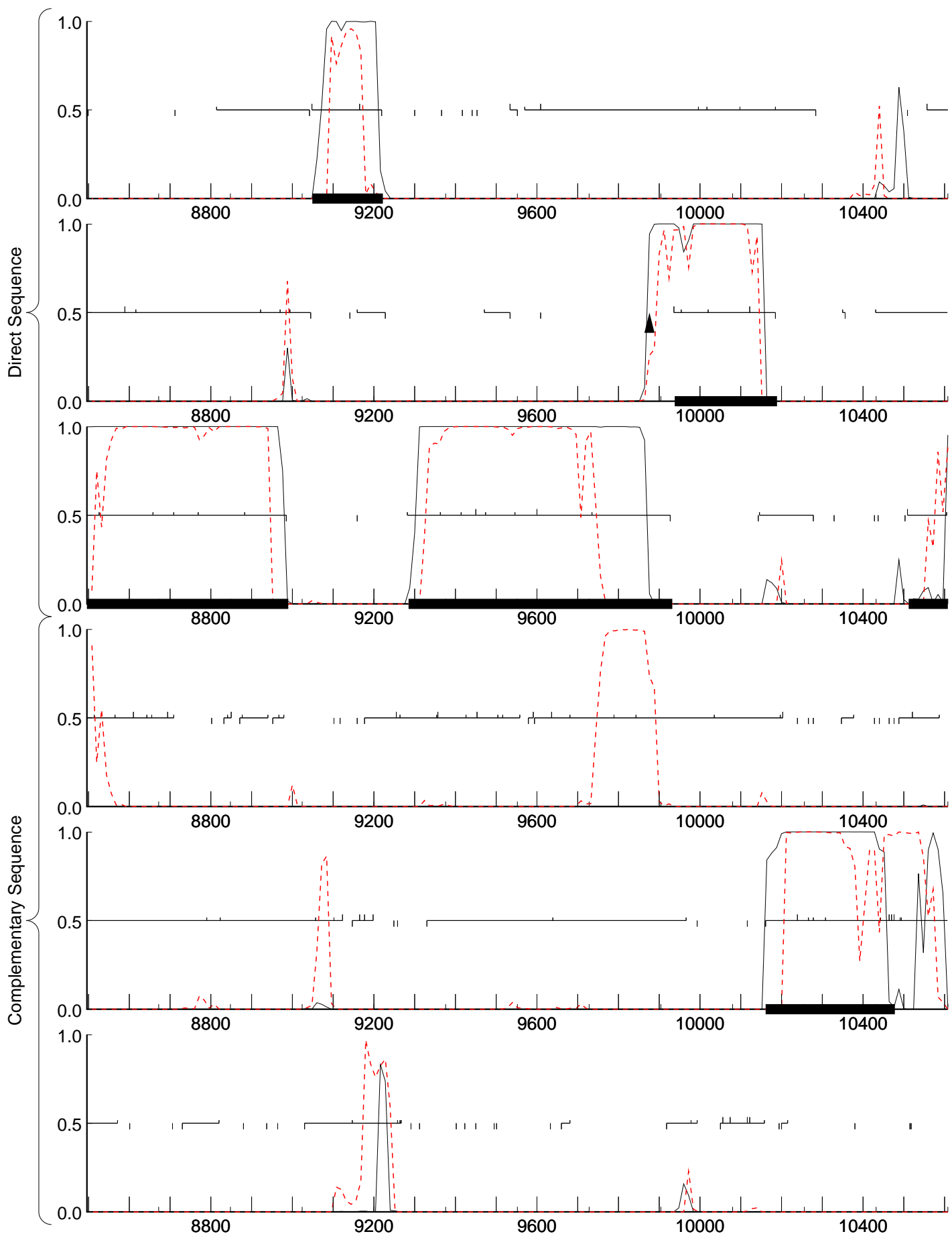






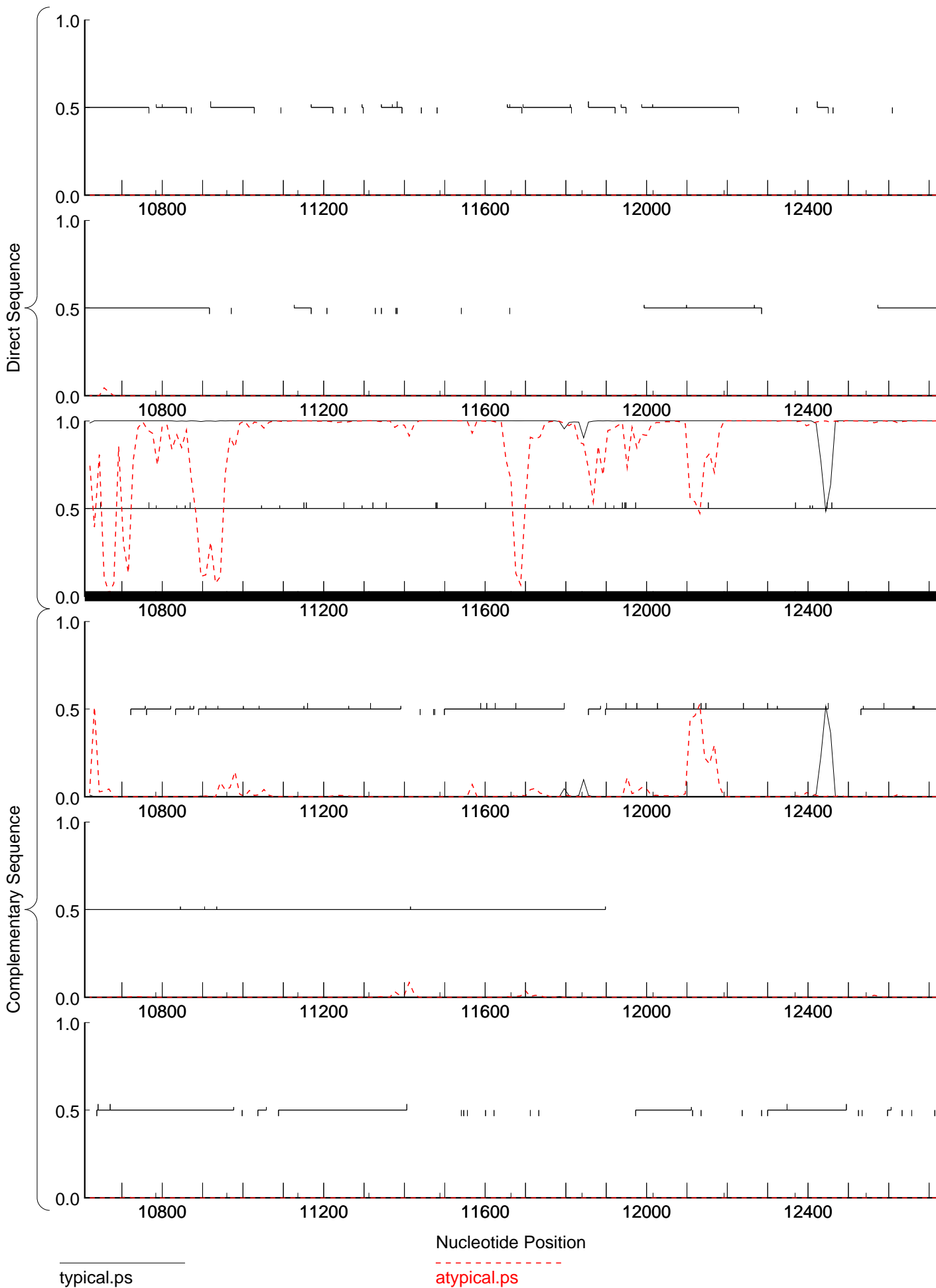


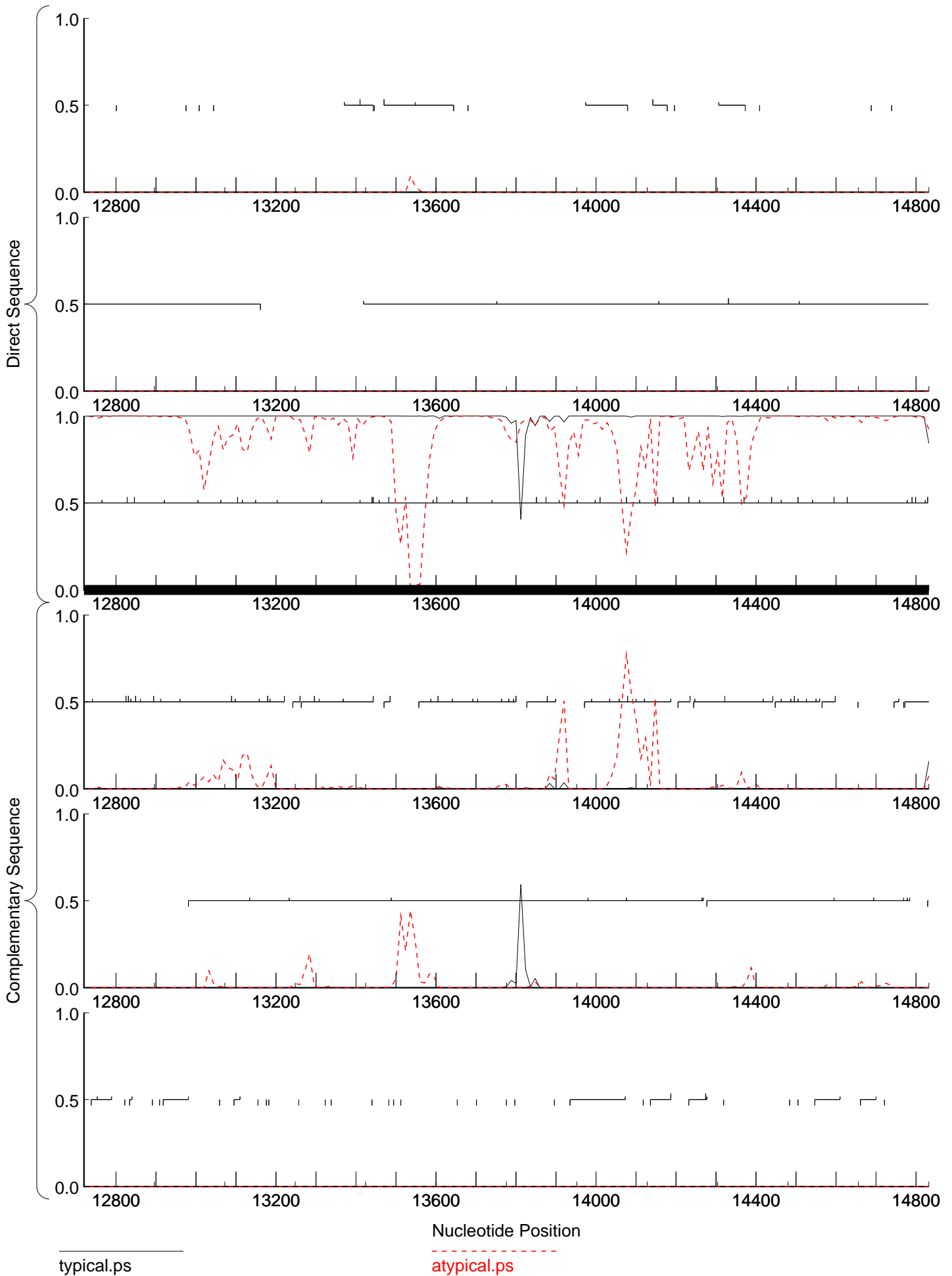
GeneMark.hmm prediction

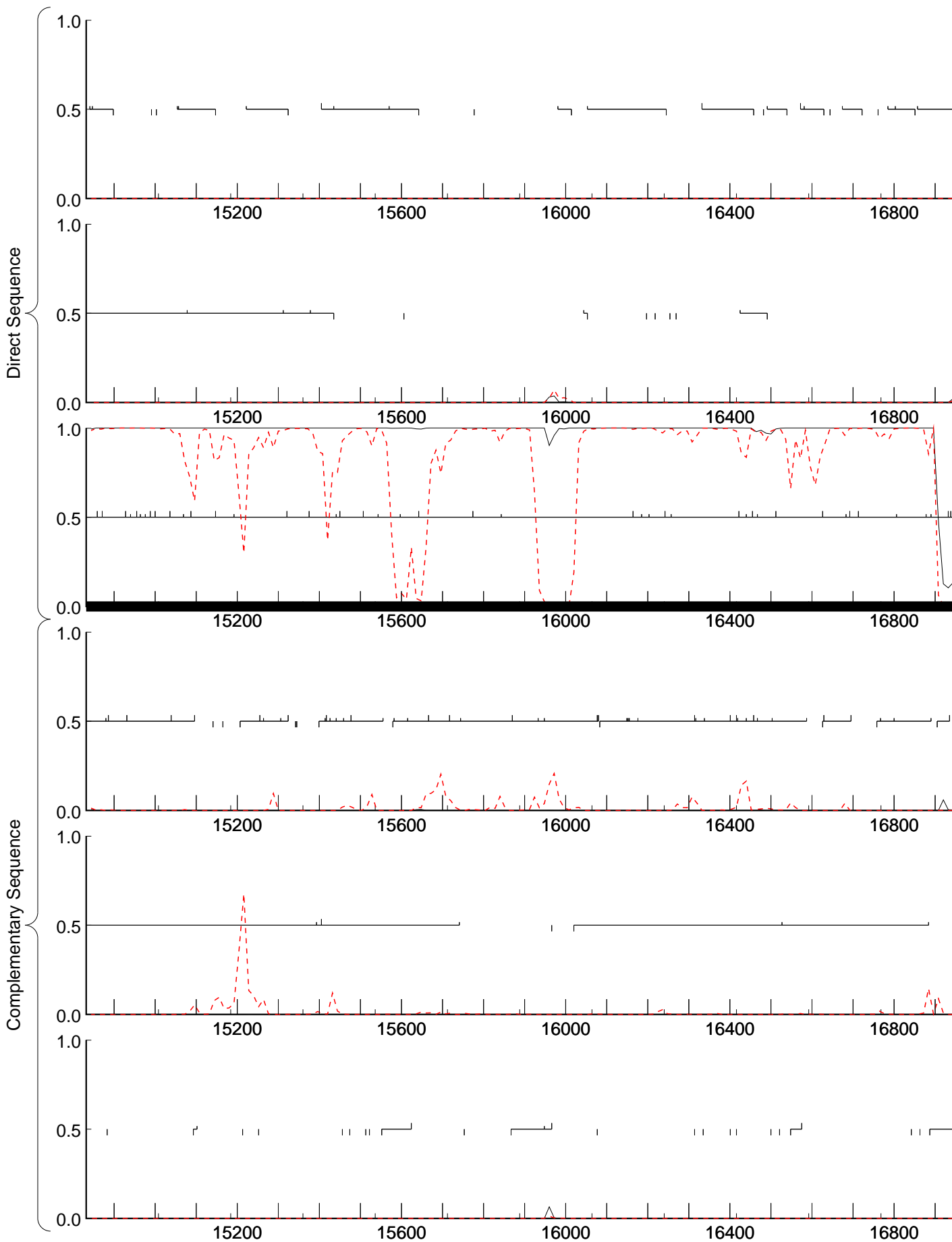


typical.ps

atypical.ps

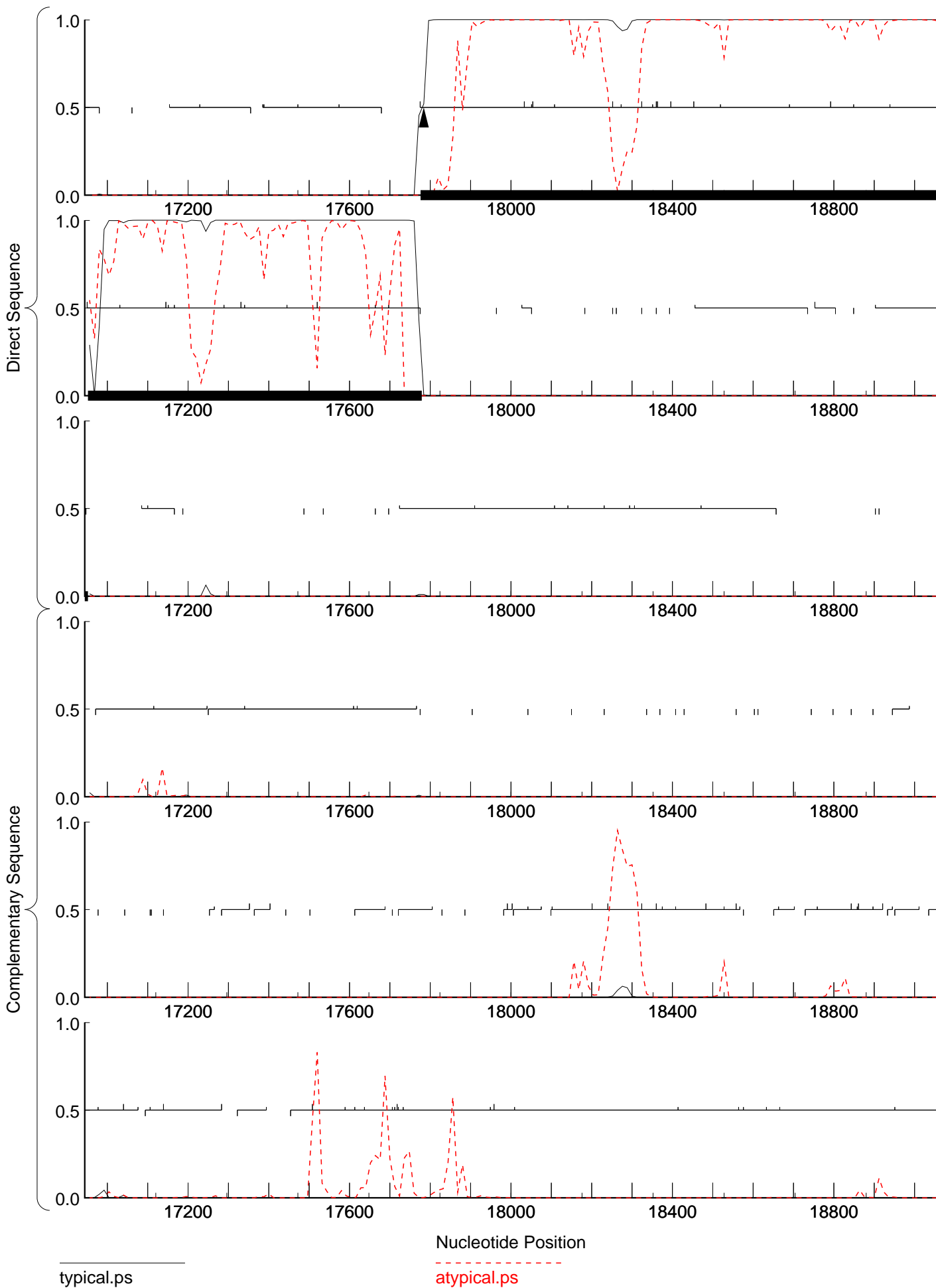


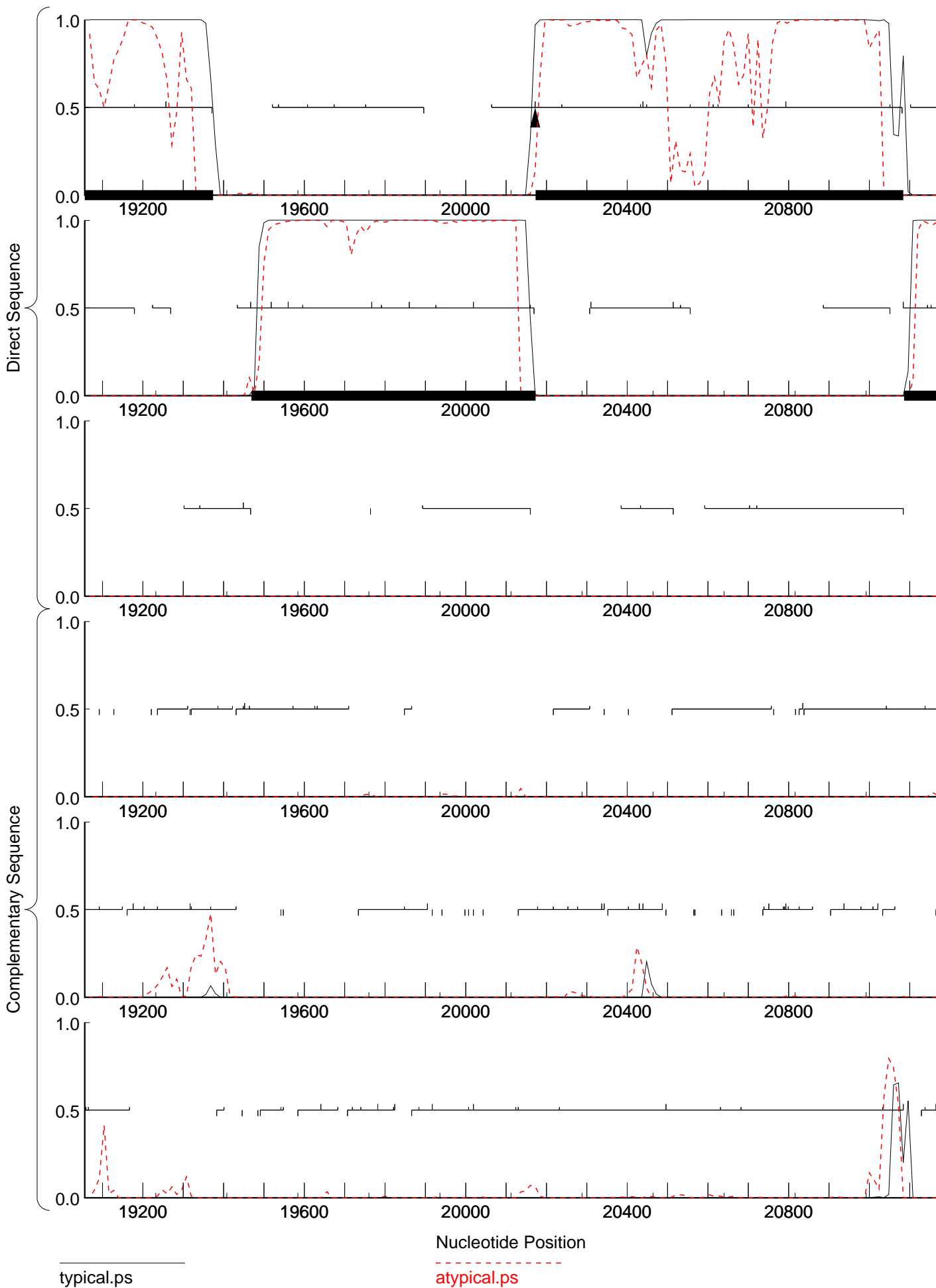


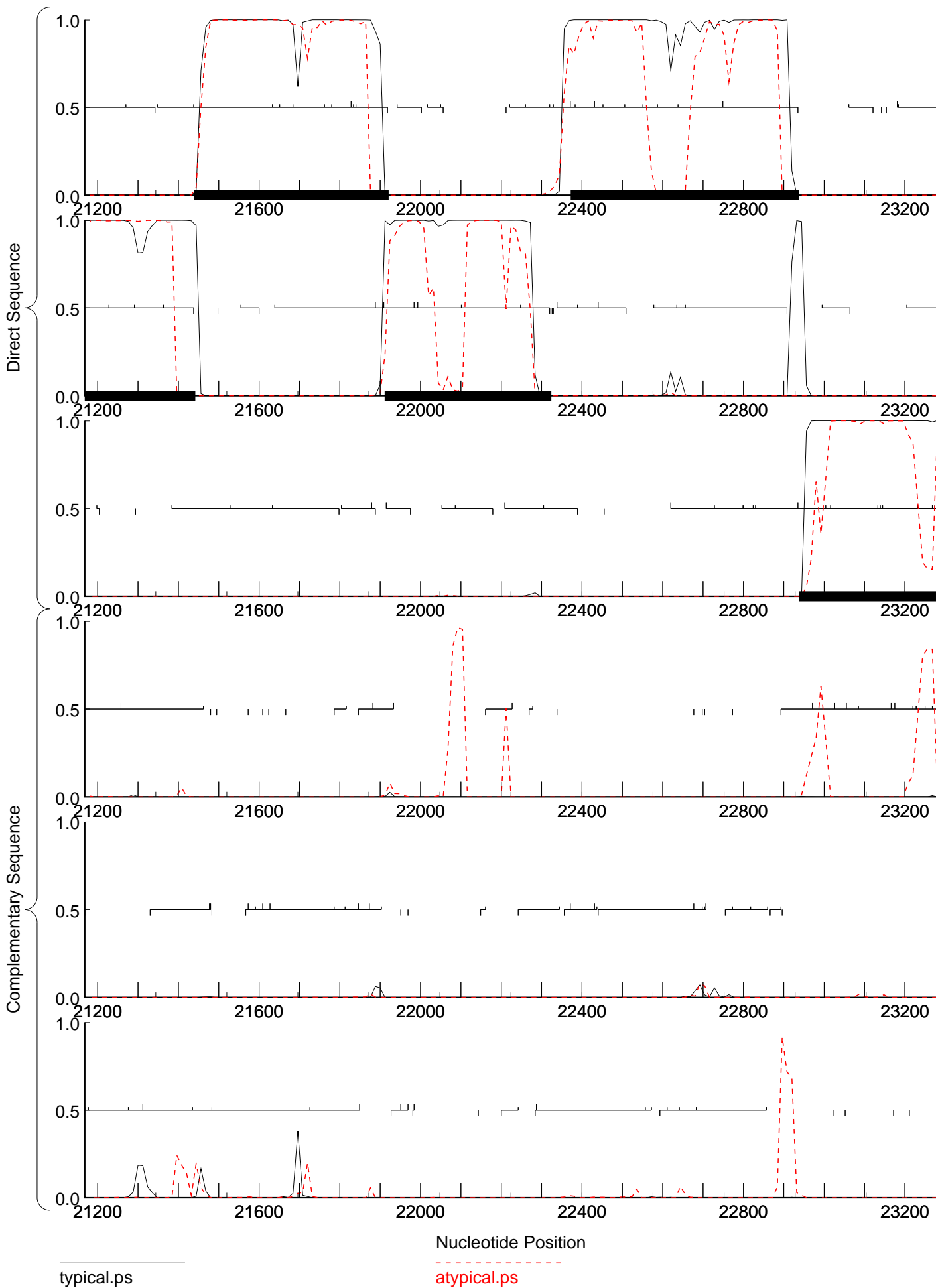


typical.ps

atypical.ps

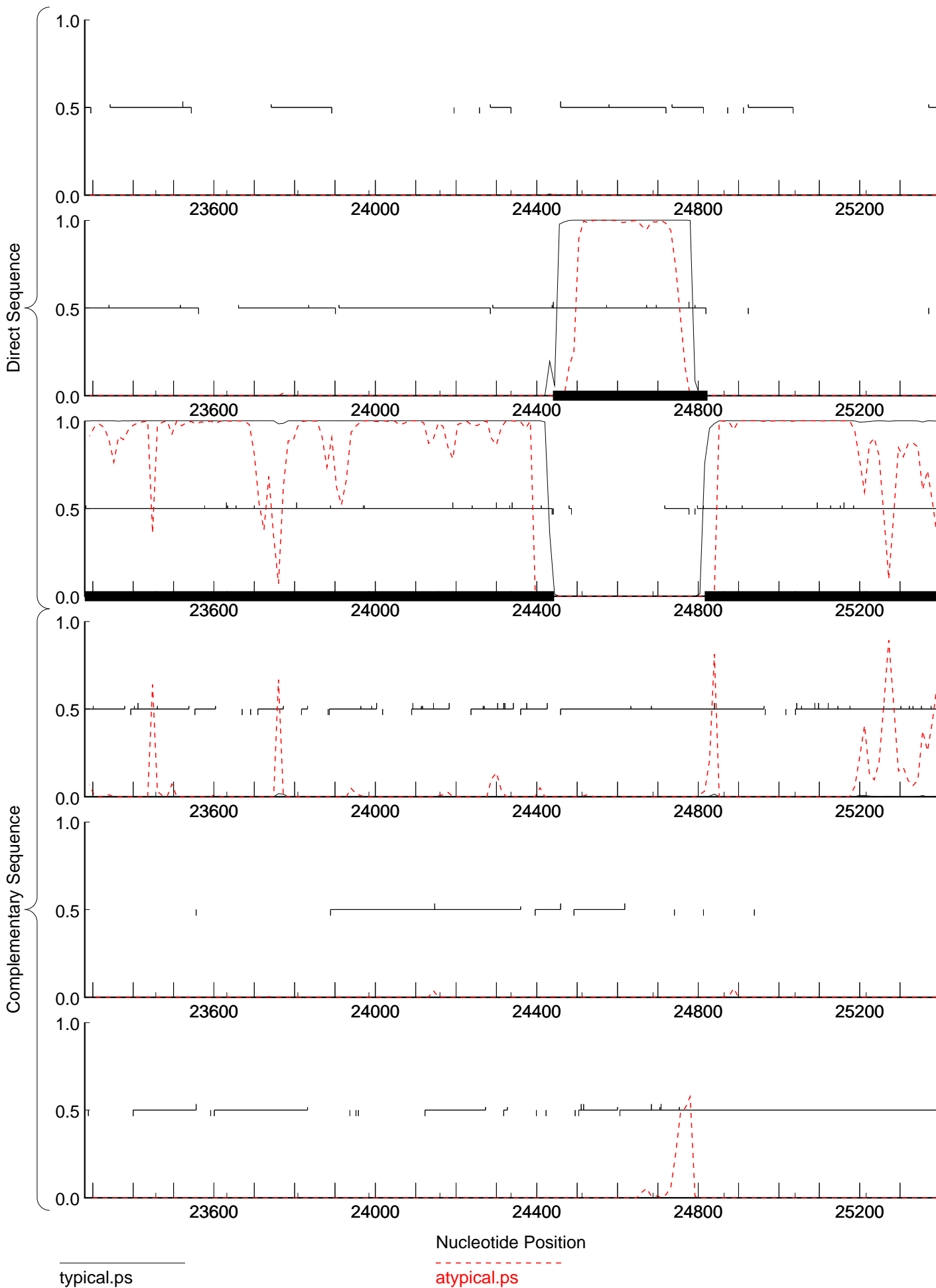


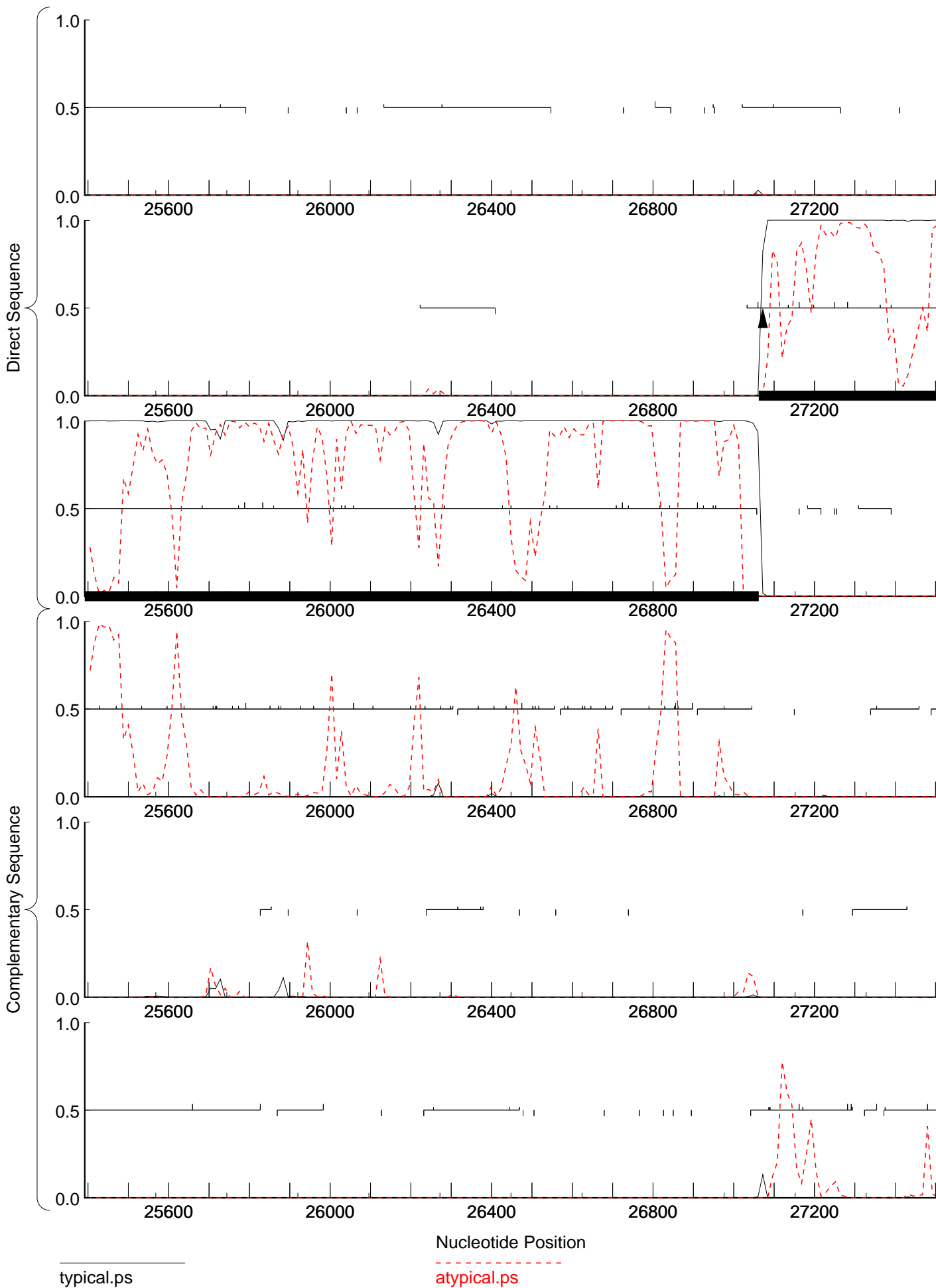


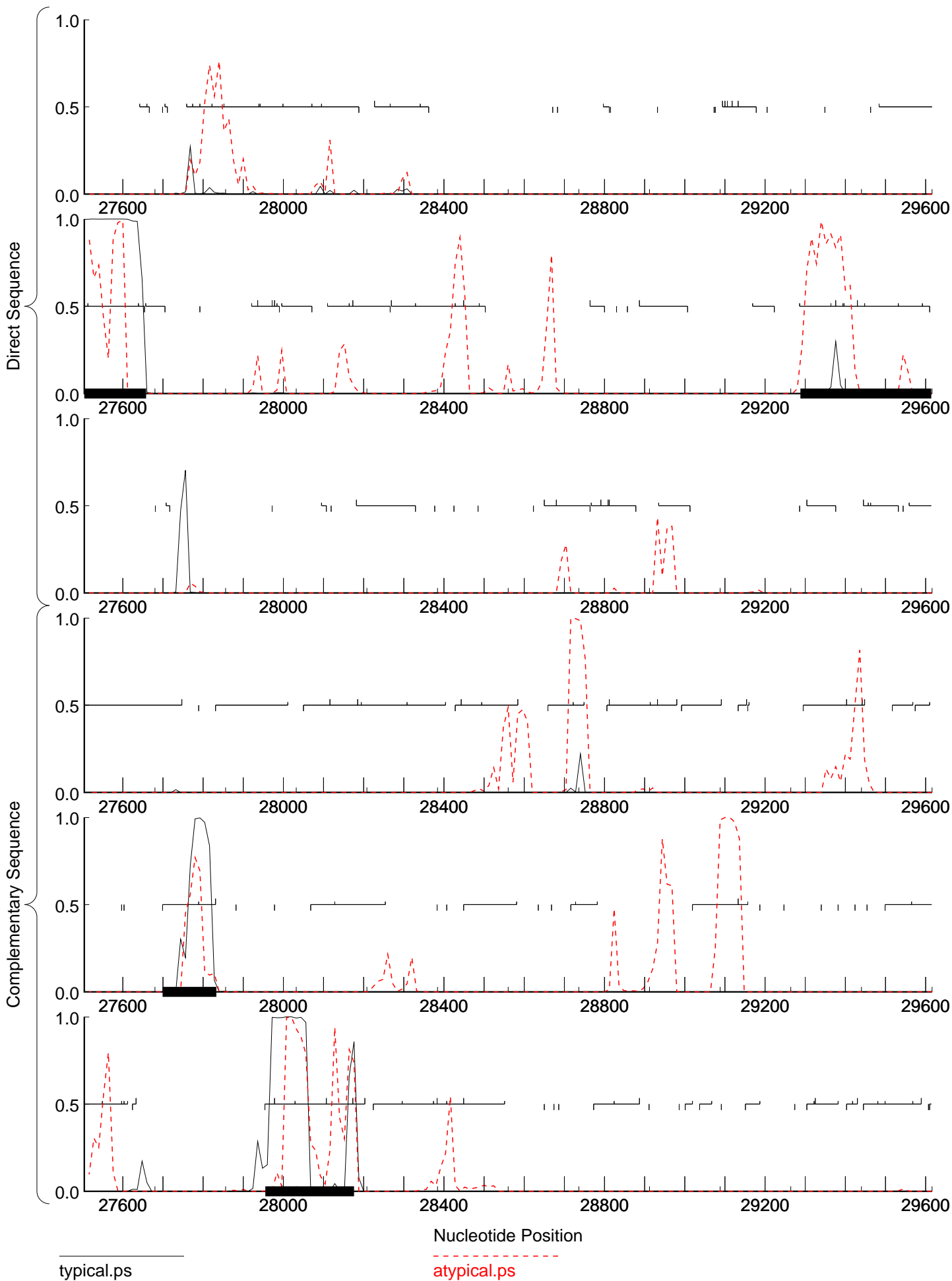


typical.ps

atypical.ps

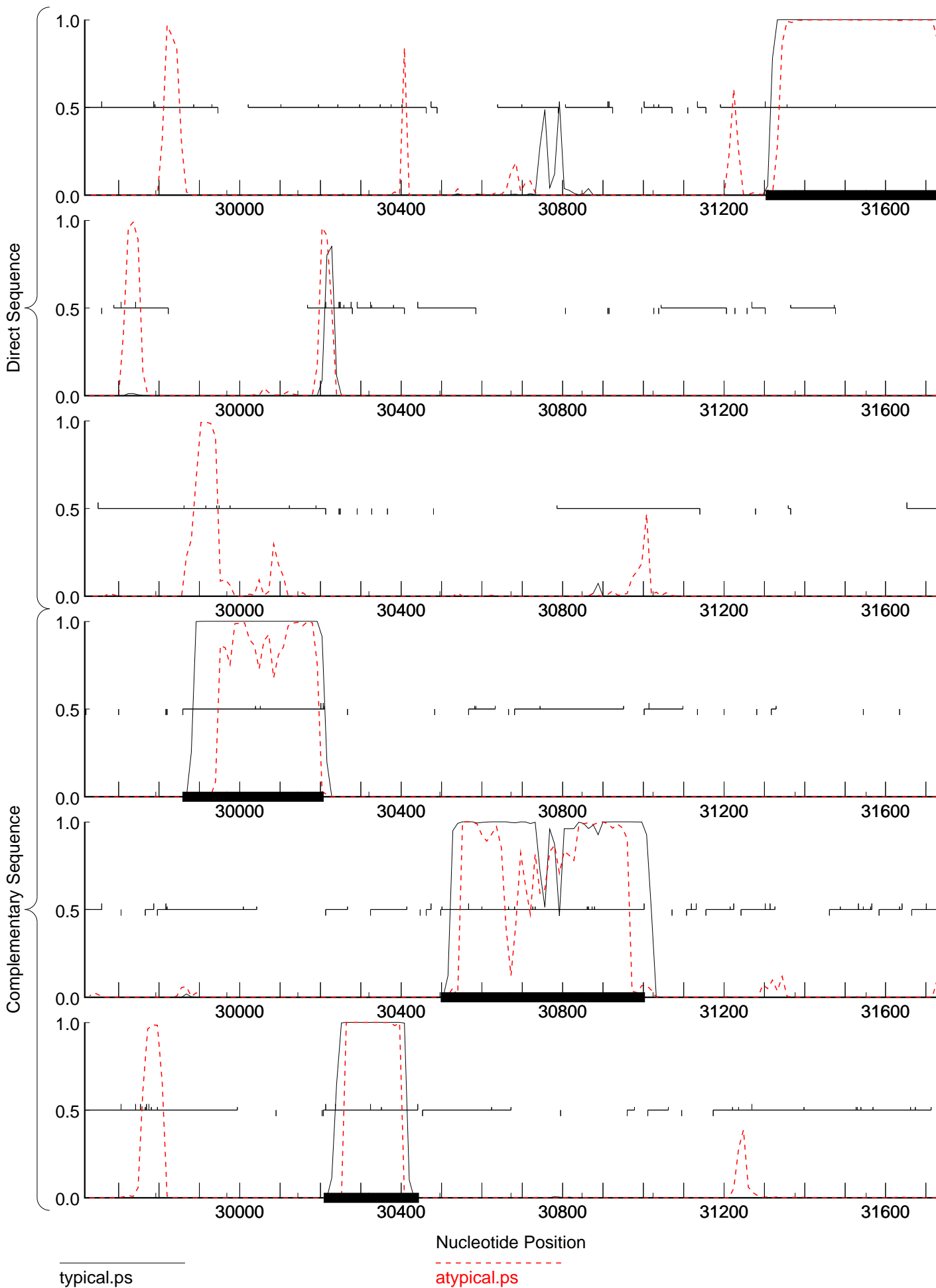






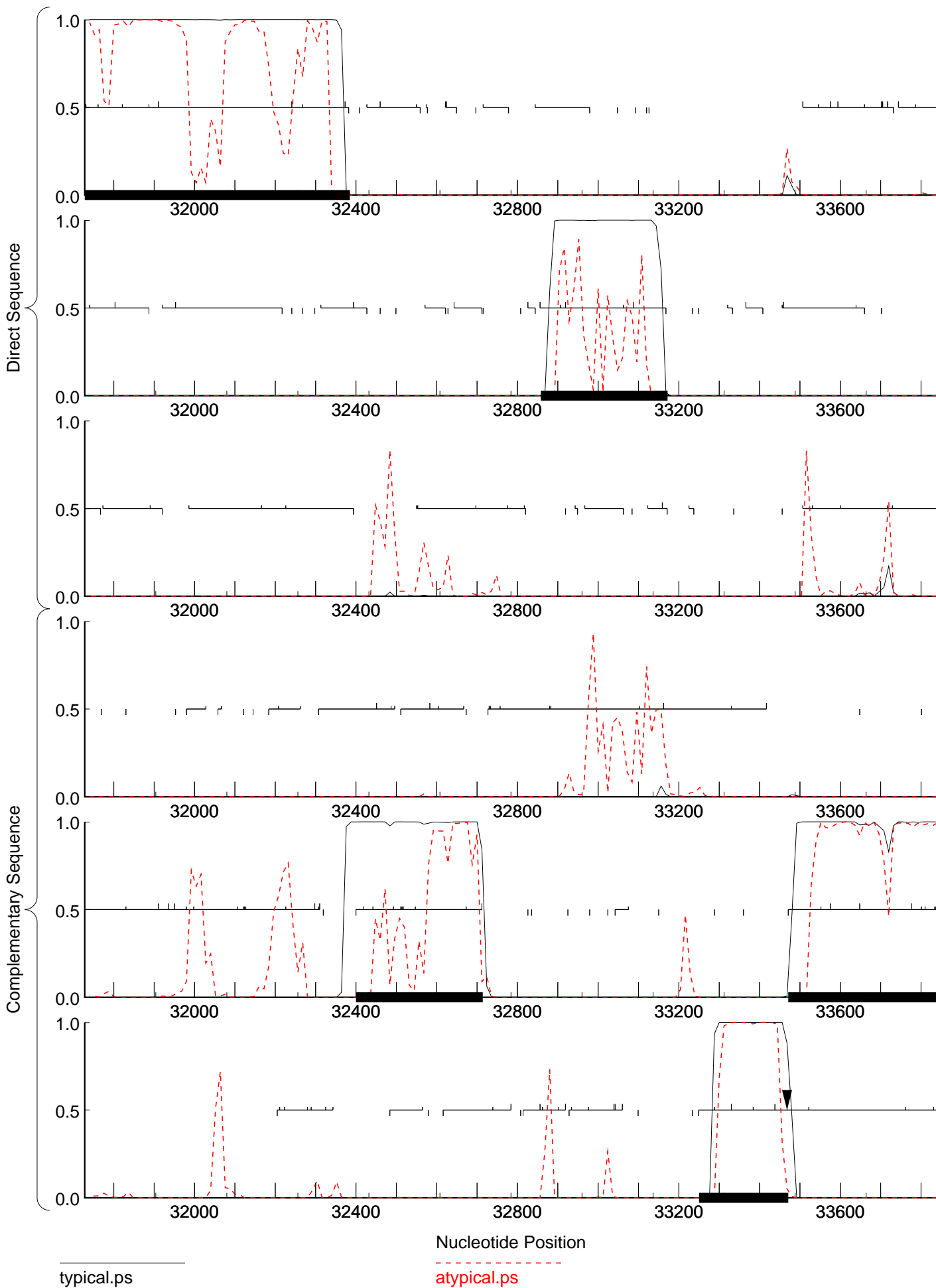
typical.ps

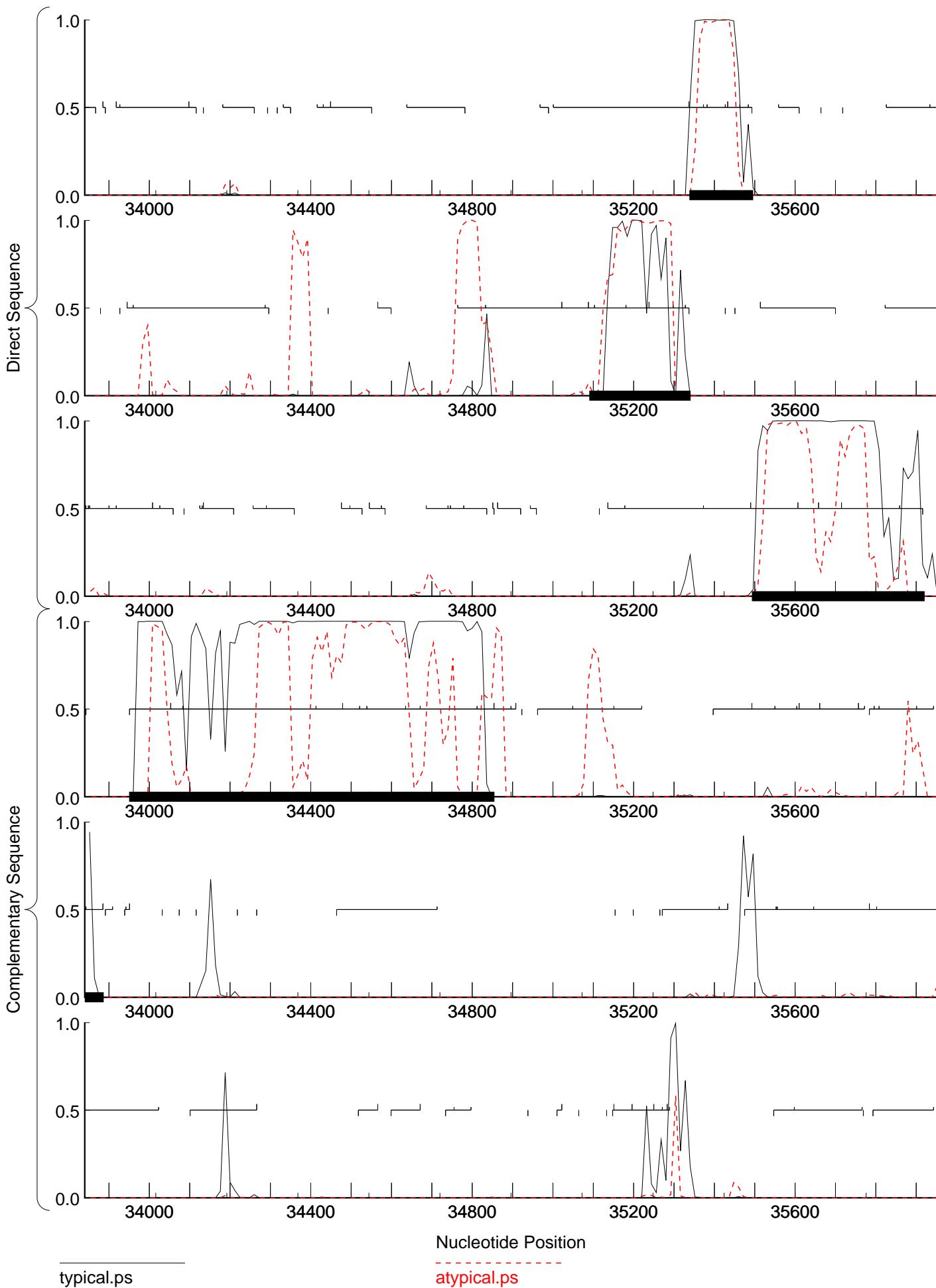
atypical.ps



typical.ps

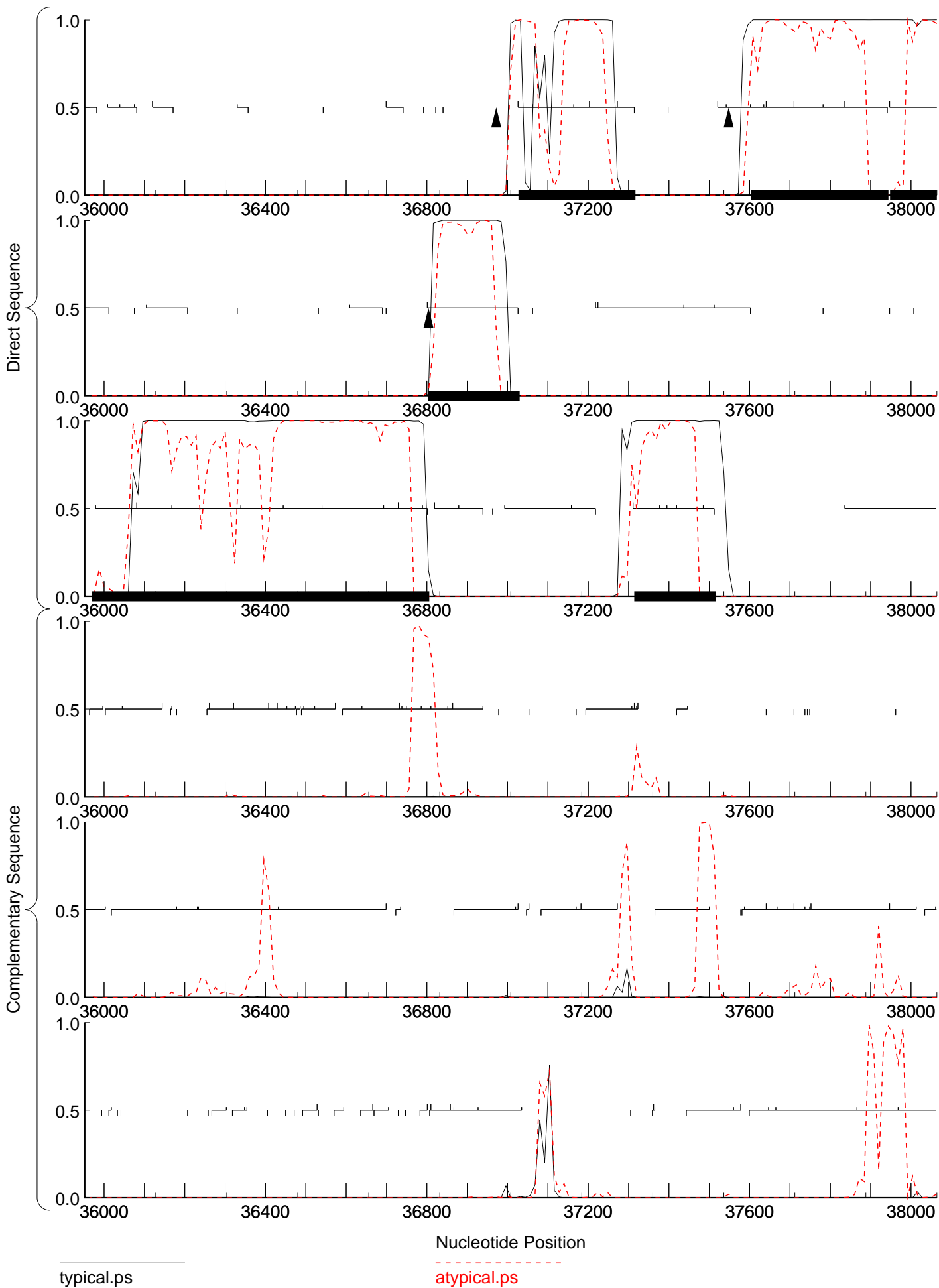
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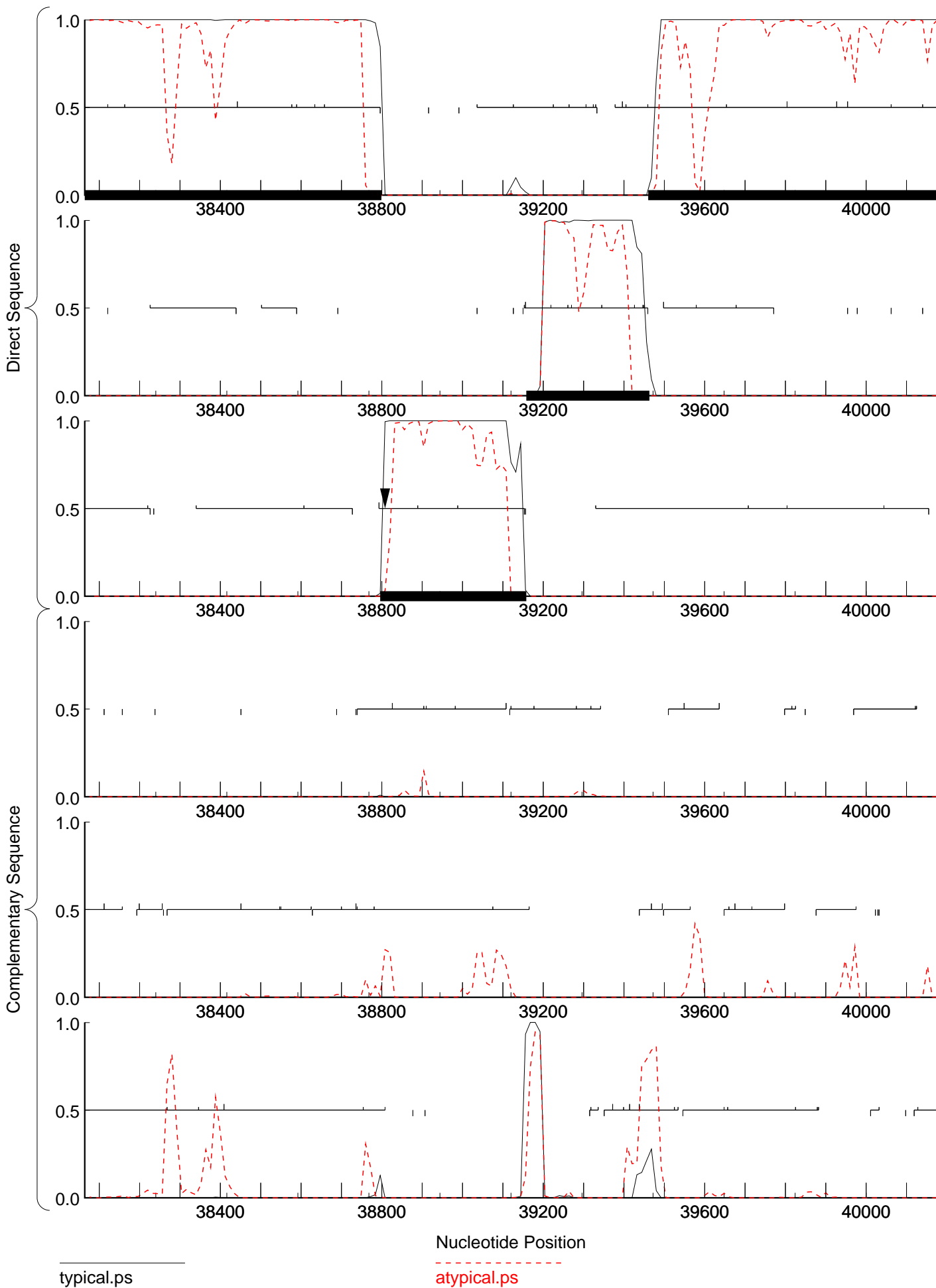




typical.ps

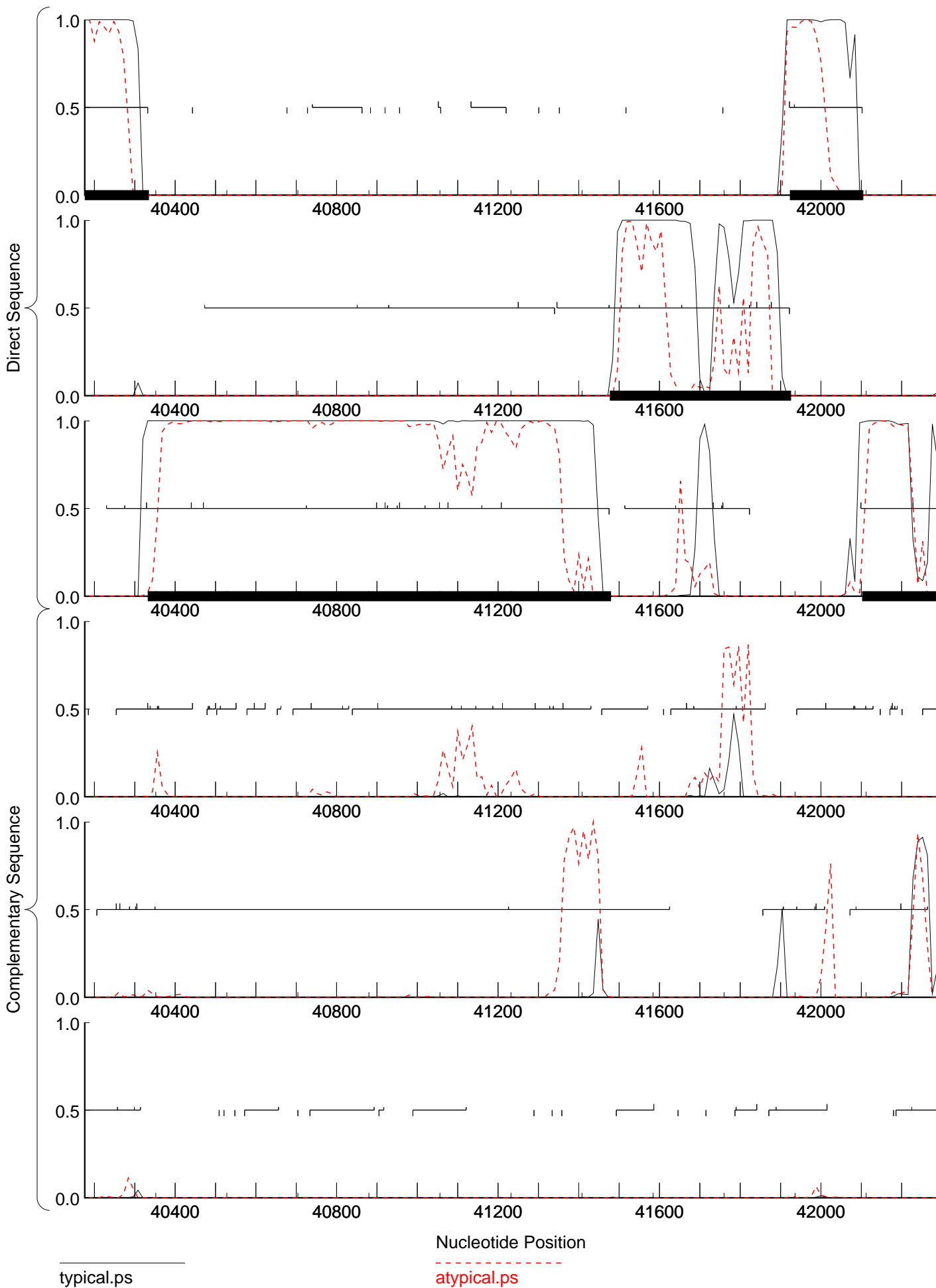
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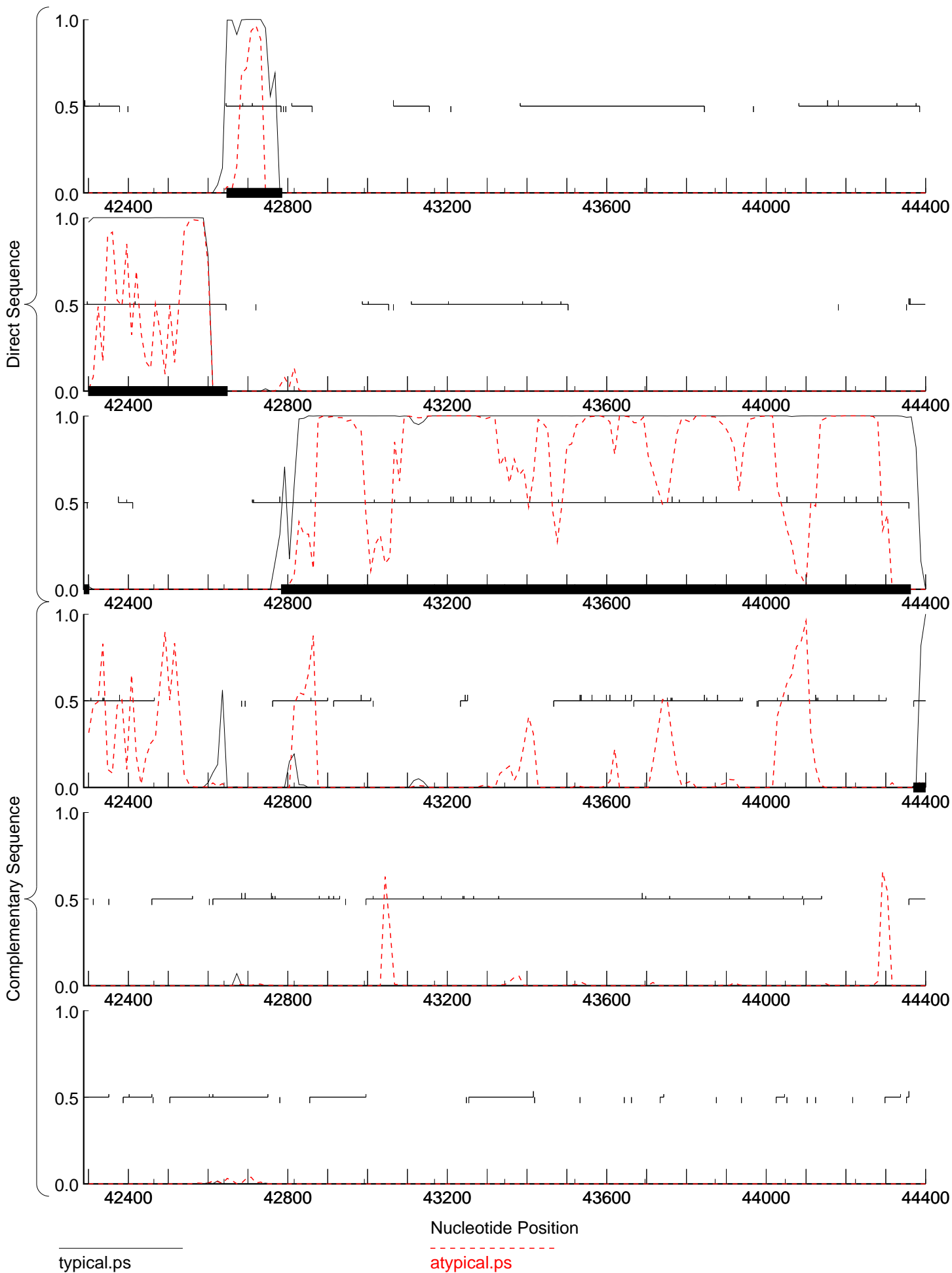


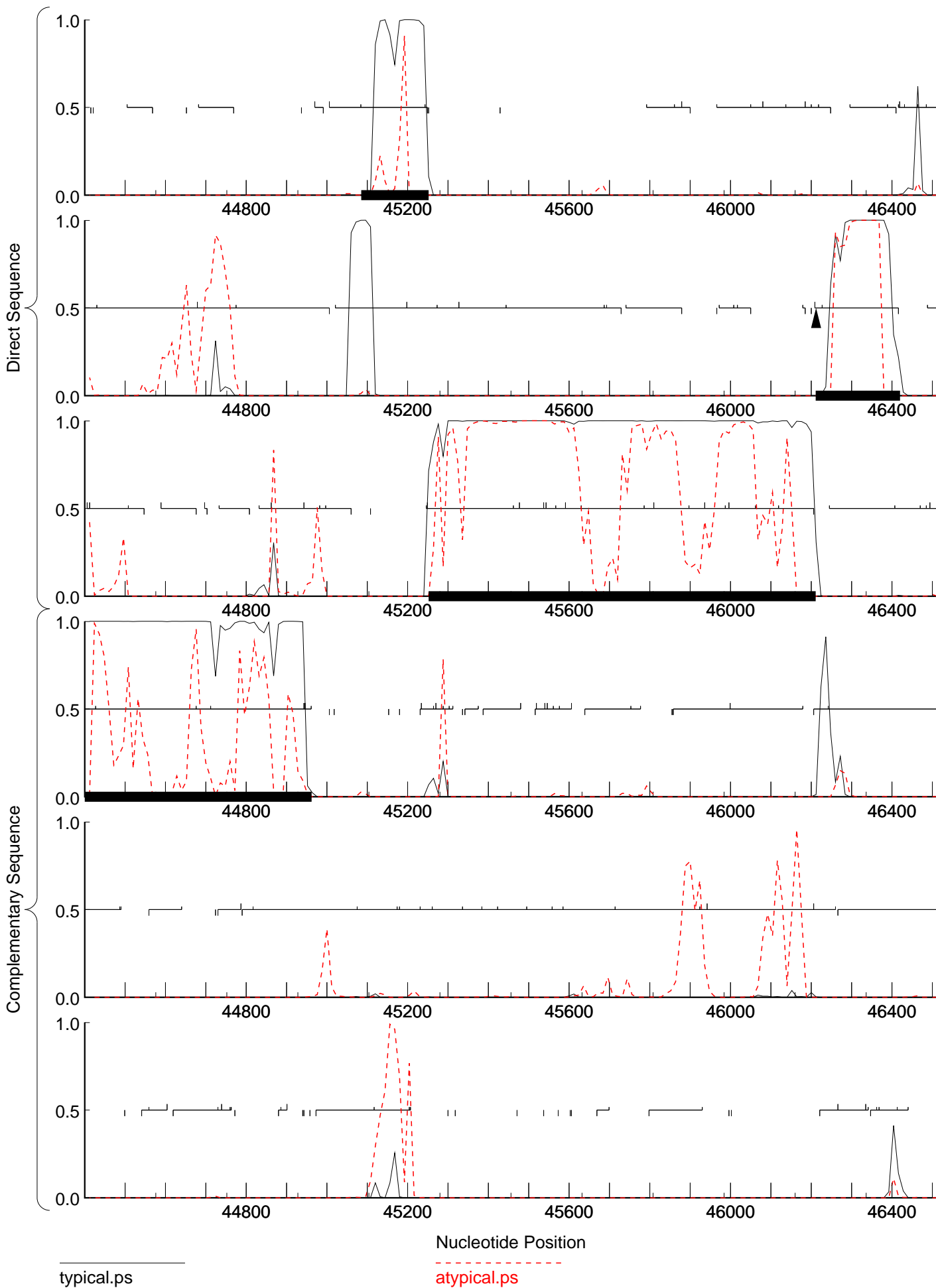


typical.ps

atypical.ps







typical.ps

atypical.ps

