

GeneMark

Version 2.5p (09.08.06)

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PROGRAM INFORMATION

Sequence : Microbacterium phage TinyTimothy complete sequence, 53932 bp, circularly permuted
Analysis Date : 8/14/18 at 4:41:52
Pages : 27
Sequence Length : 53932 bp
GC Content : 56.75%

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,

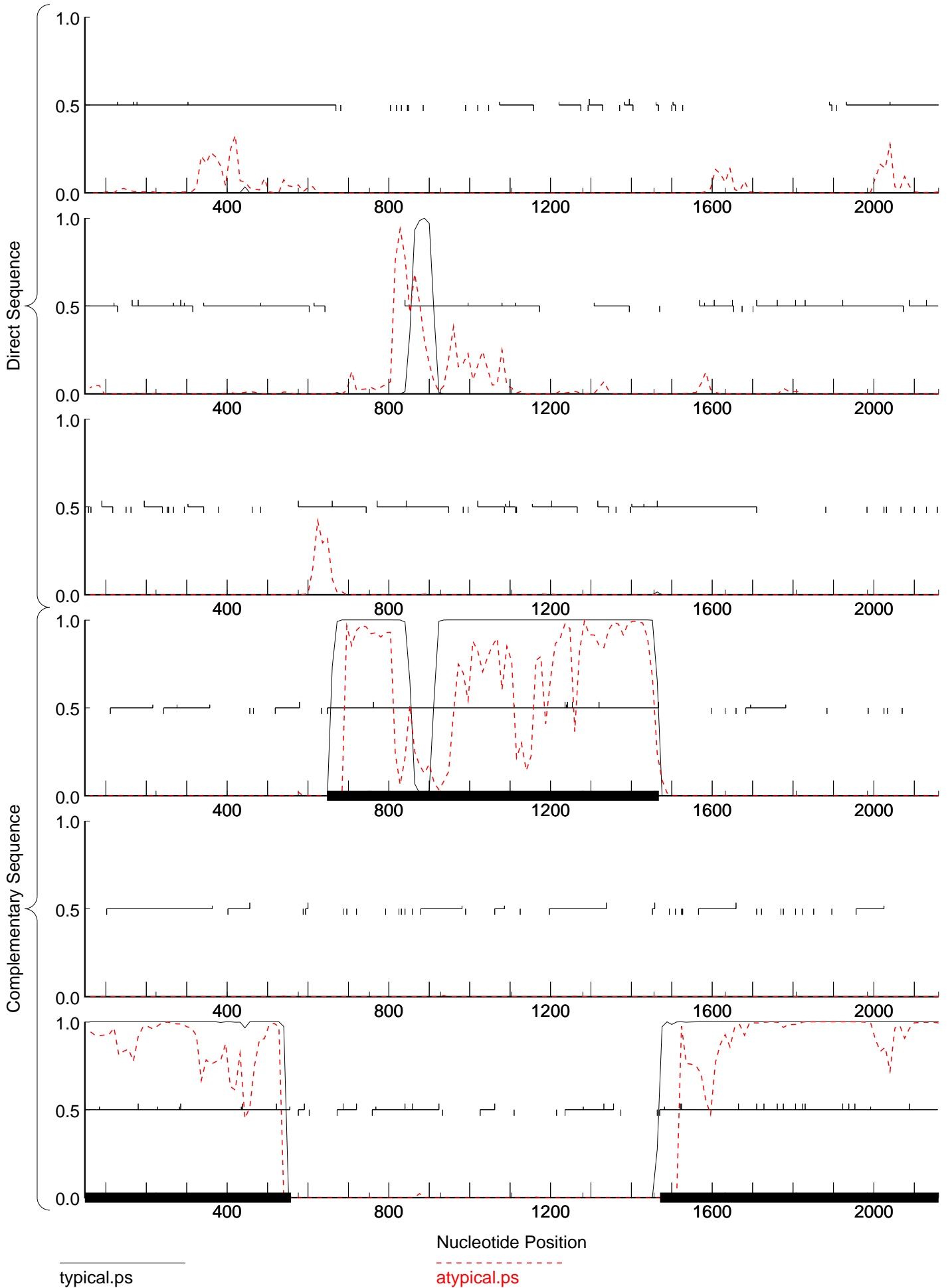
Matrix notes & comments

Training set derived by GeneMarkS, 4.30 October 2014
Tue Aug 14 04:41:51 2018

MATRIX INFORMATION

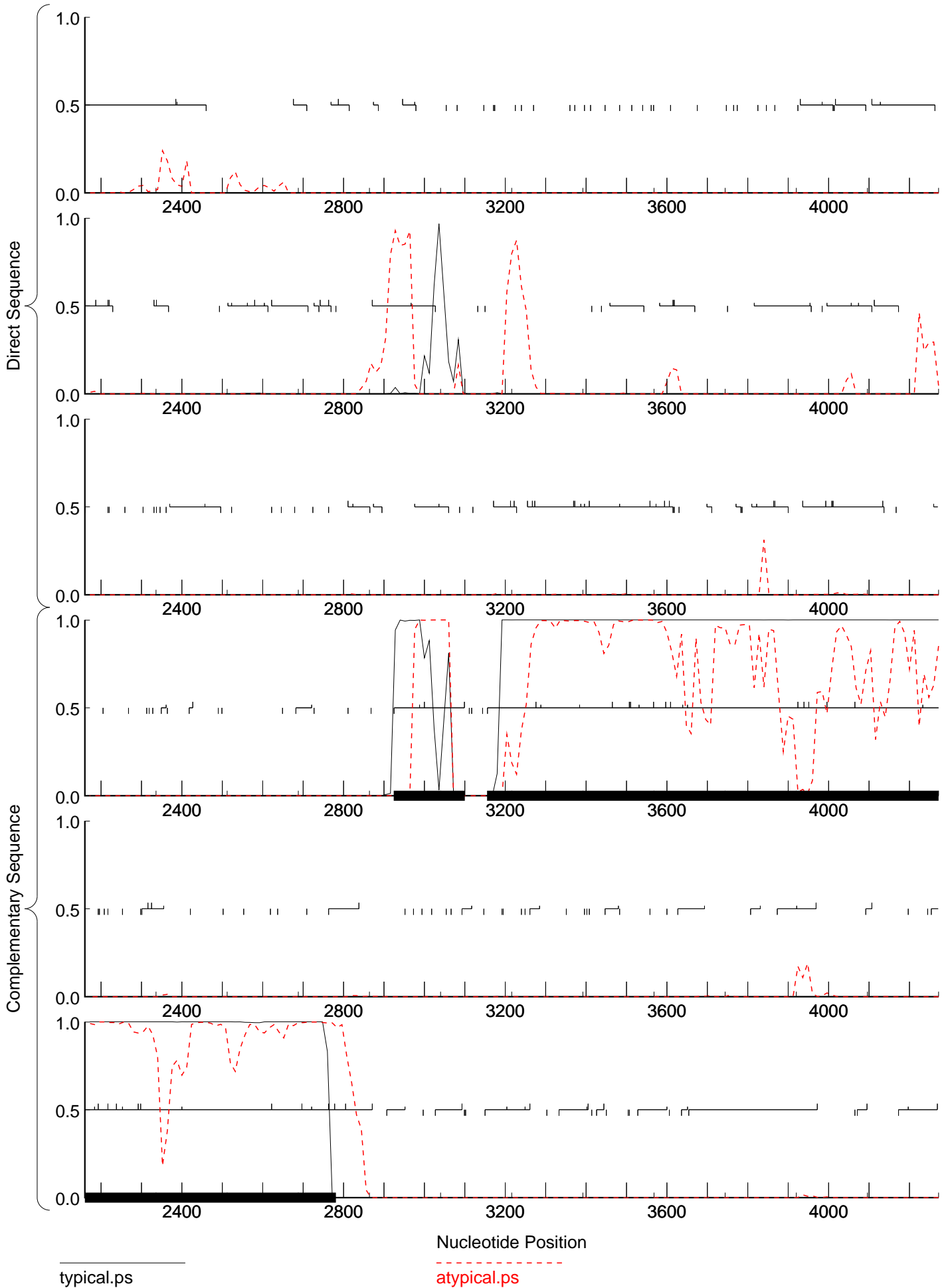
Matrix : -
Author : -
Order : 4

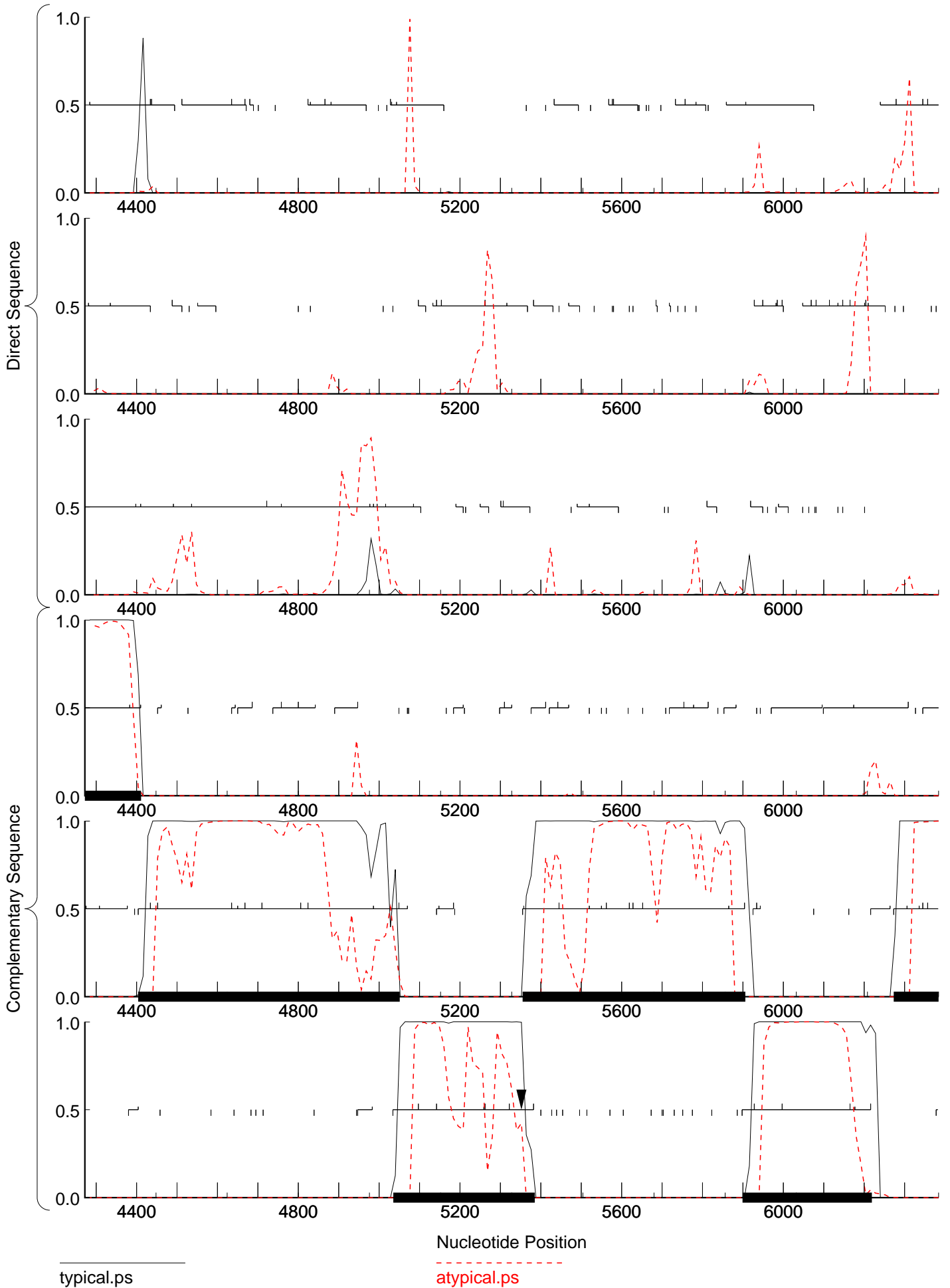
Send questions / comments to:
Dr. M. Borodovsky
Georgia Institute of Technology
School of Biology
Atlanta, GA 30332-0230

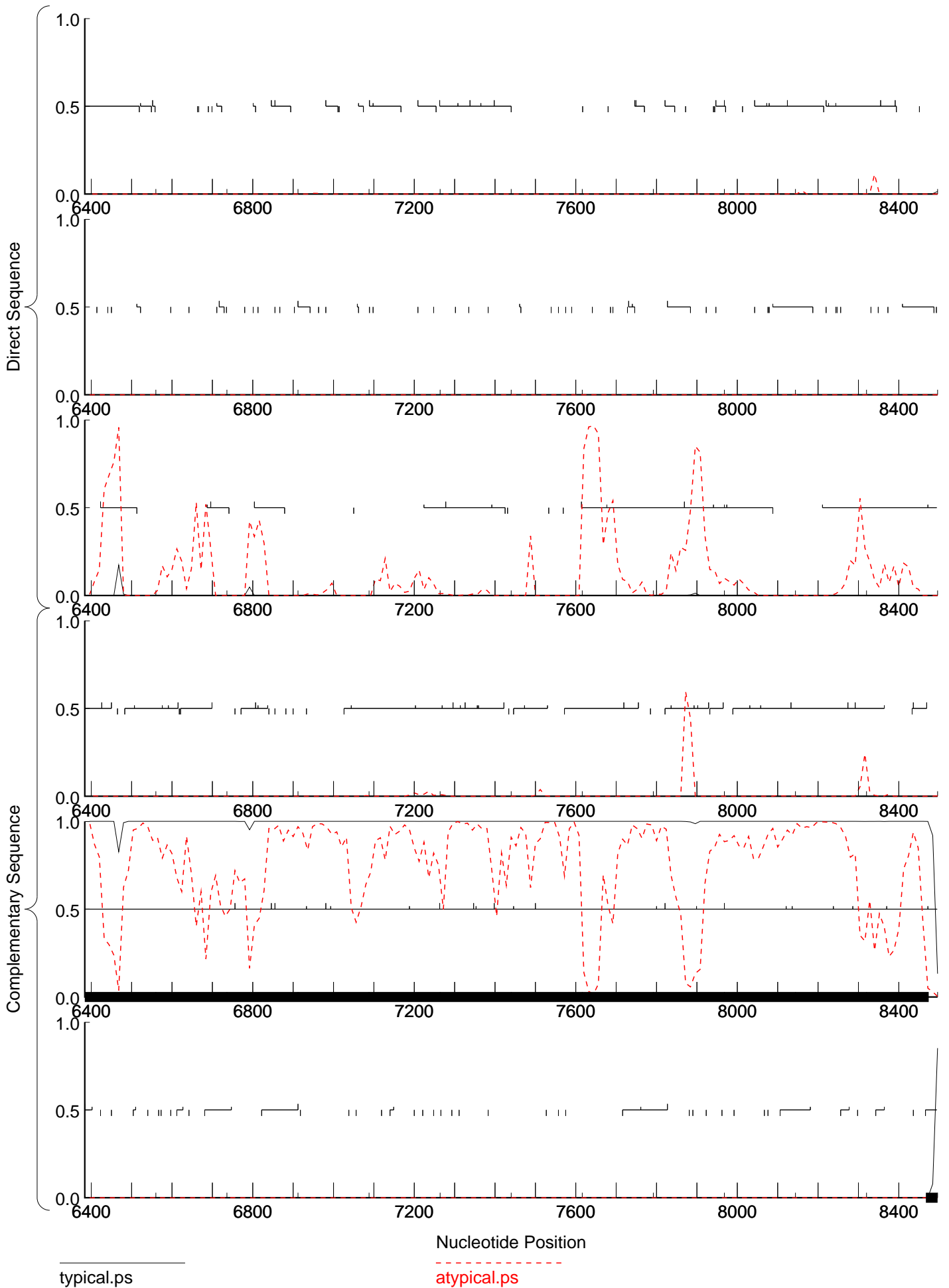


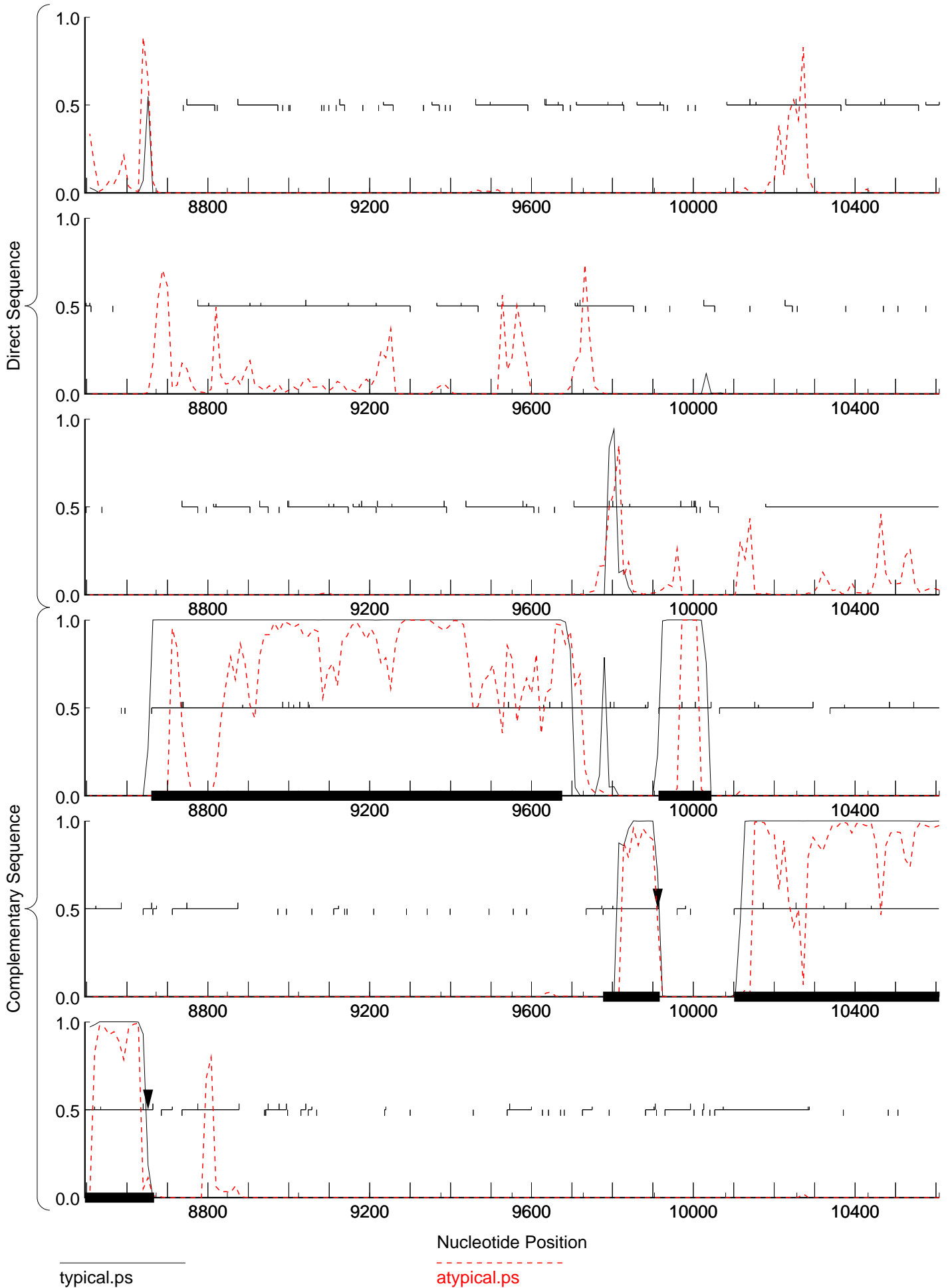
typical.ps

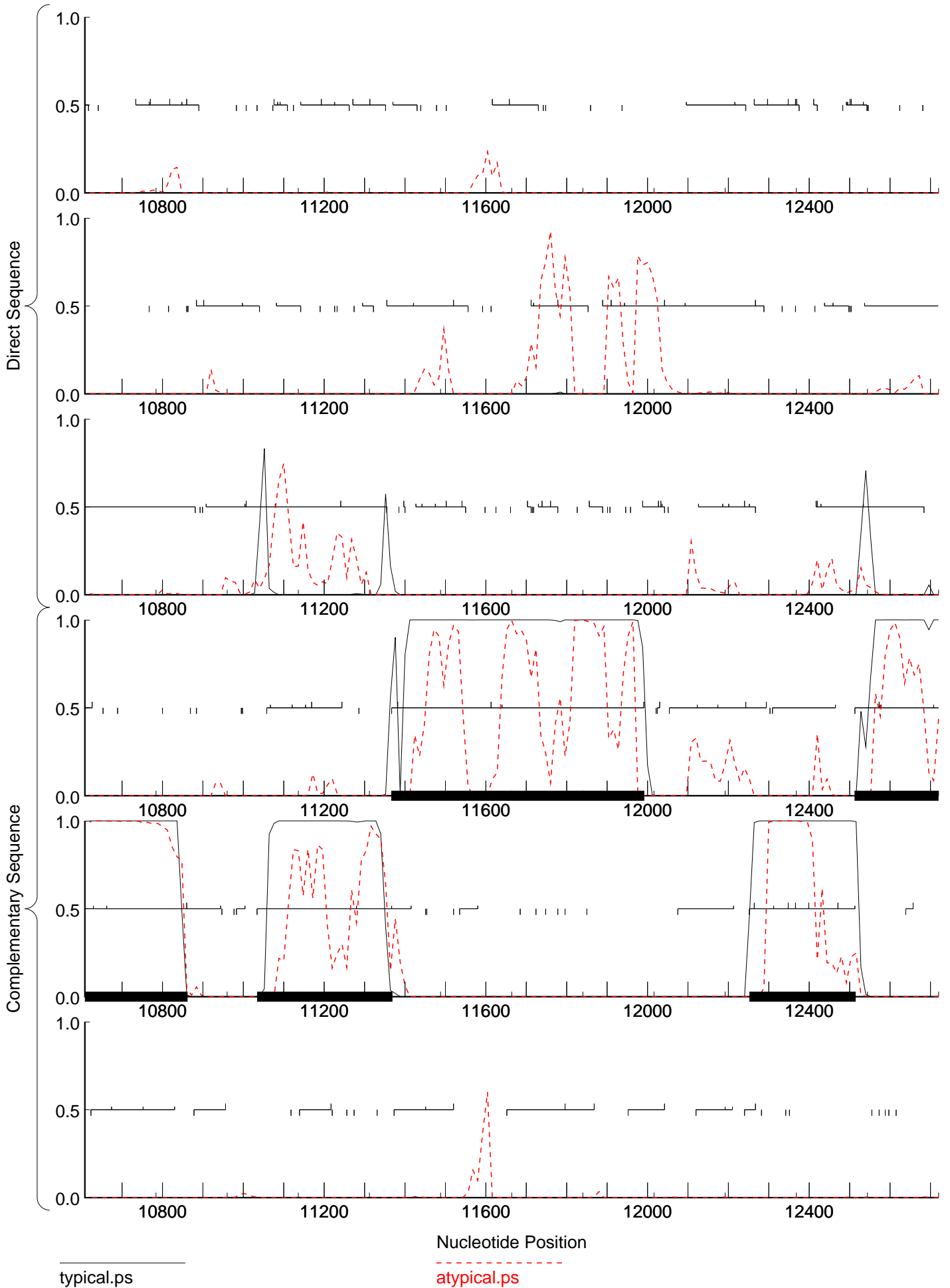
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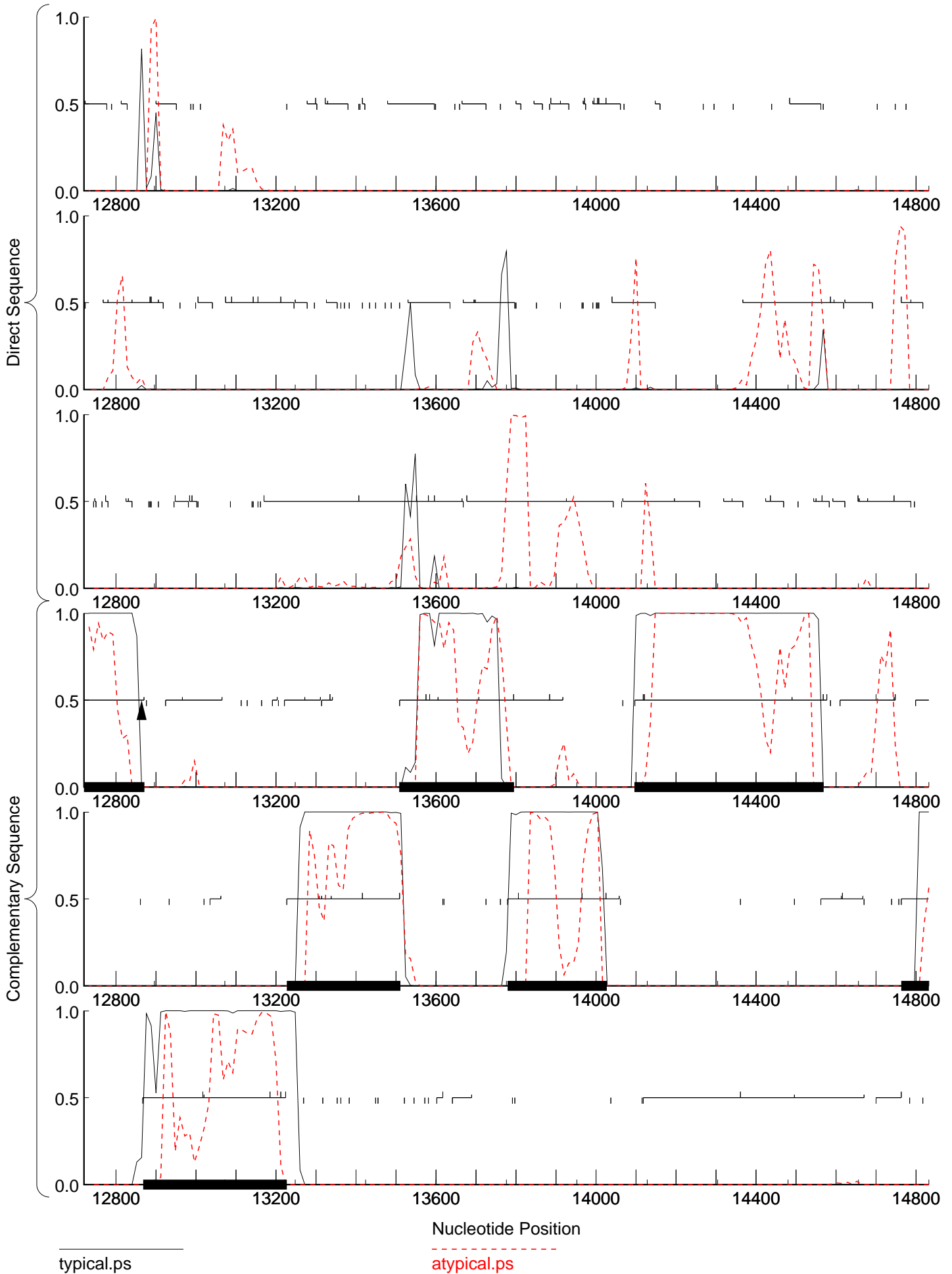


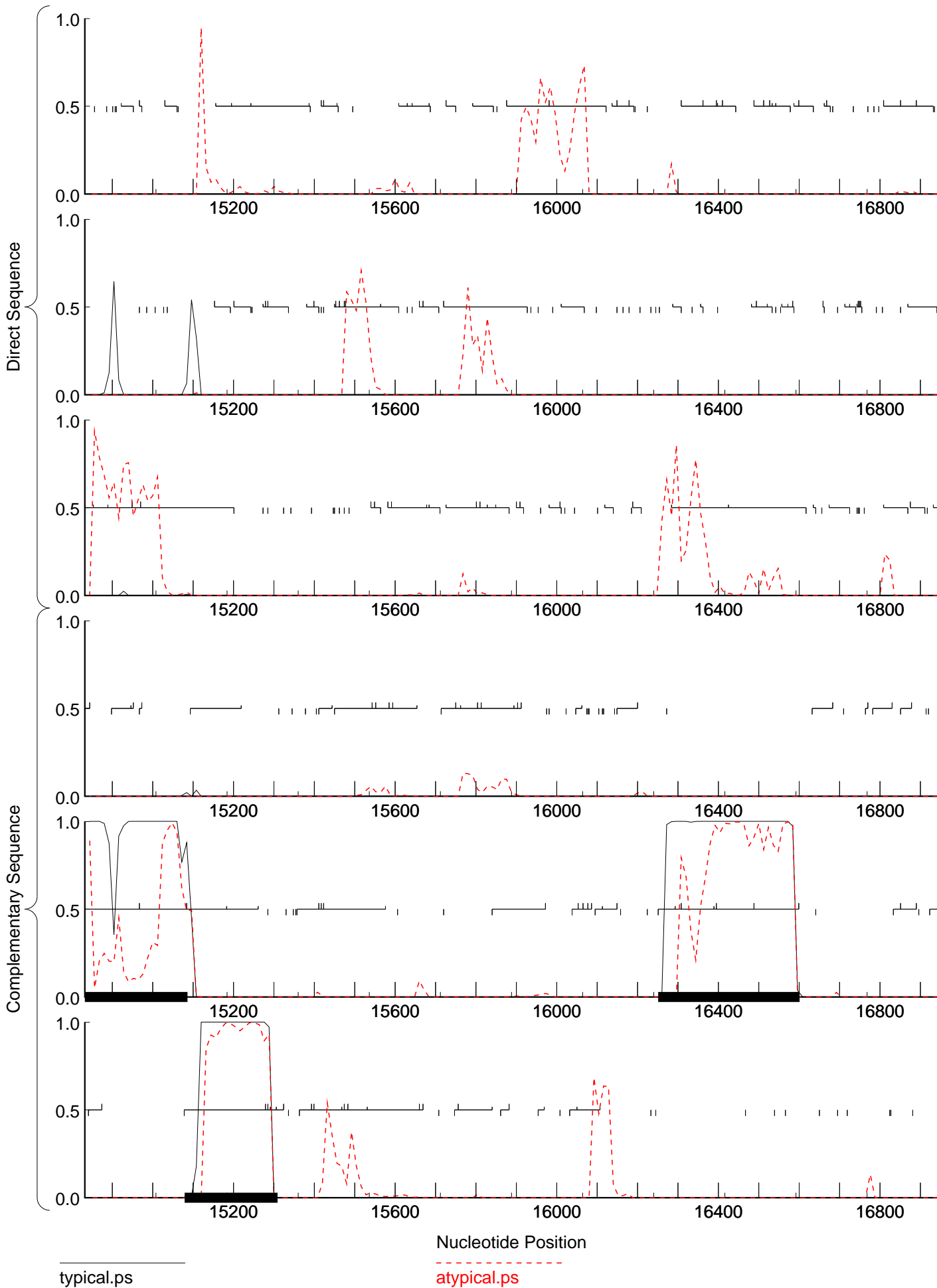






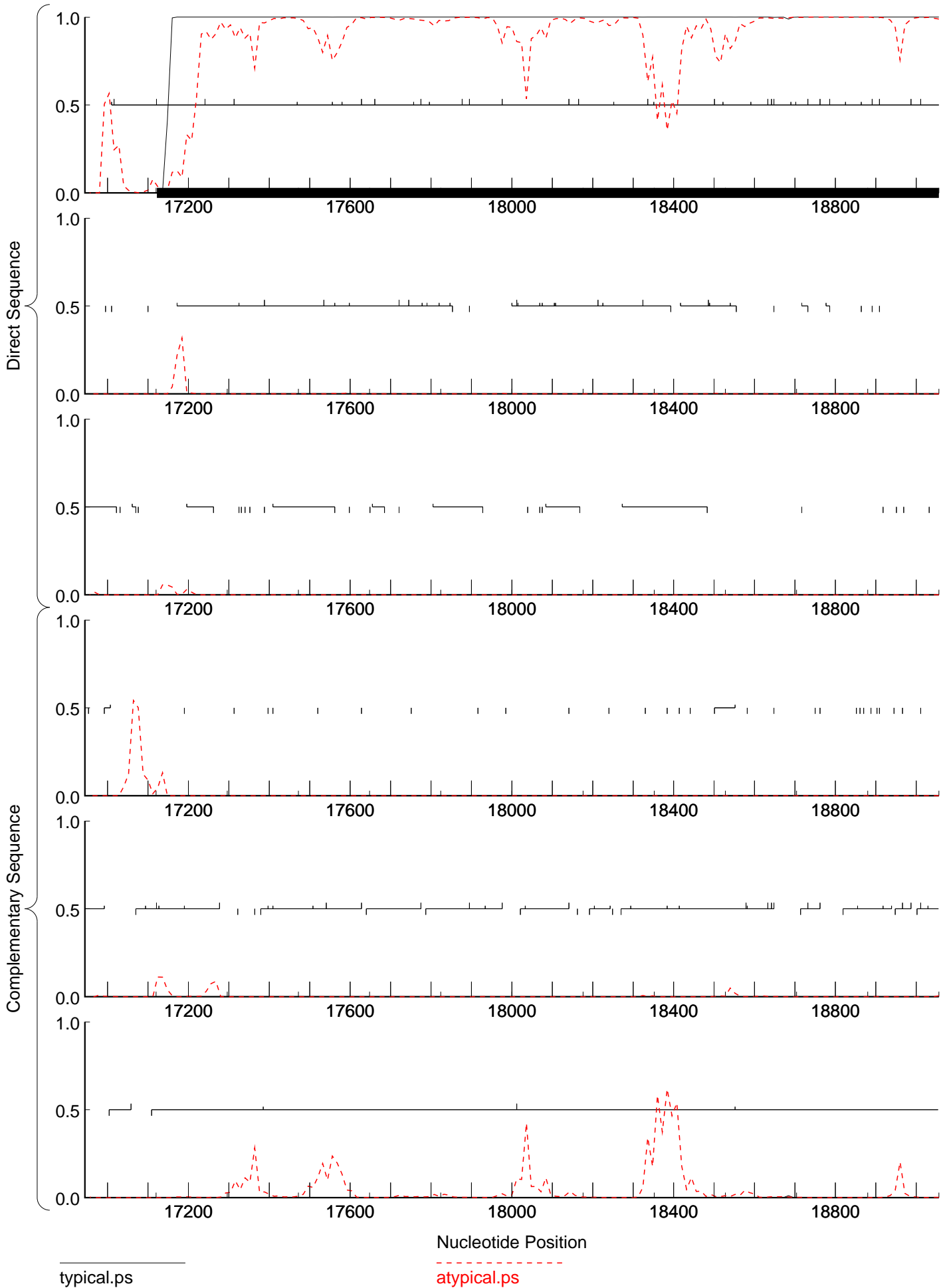


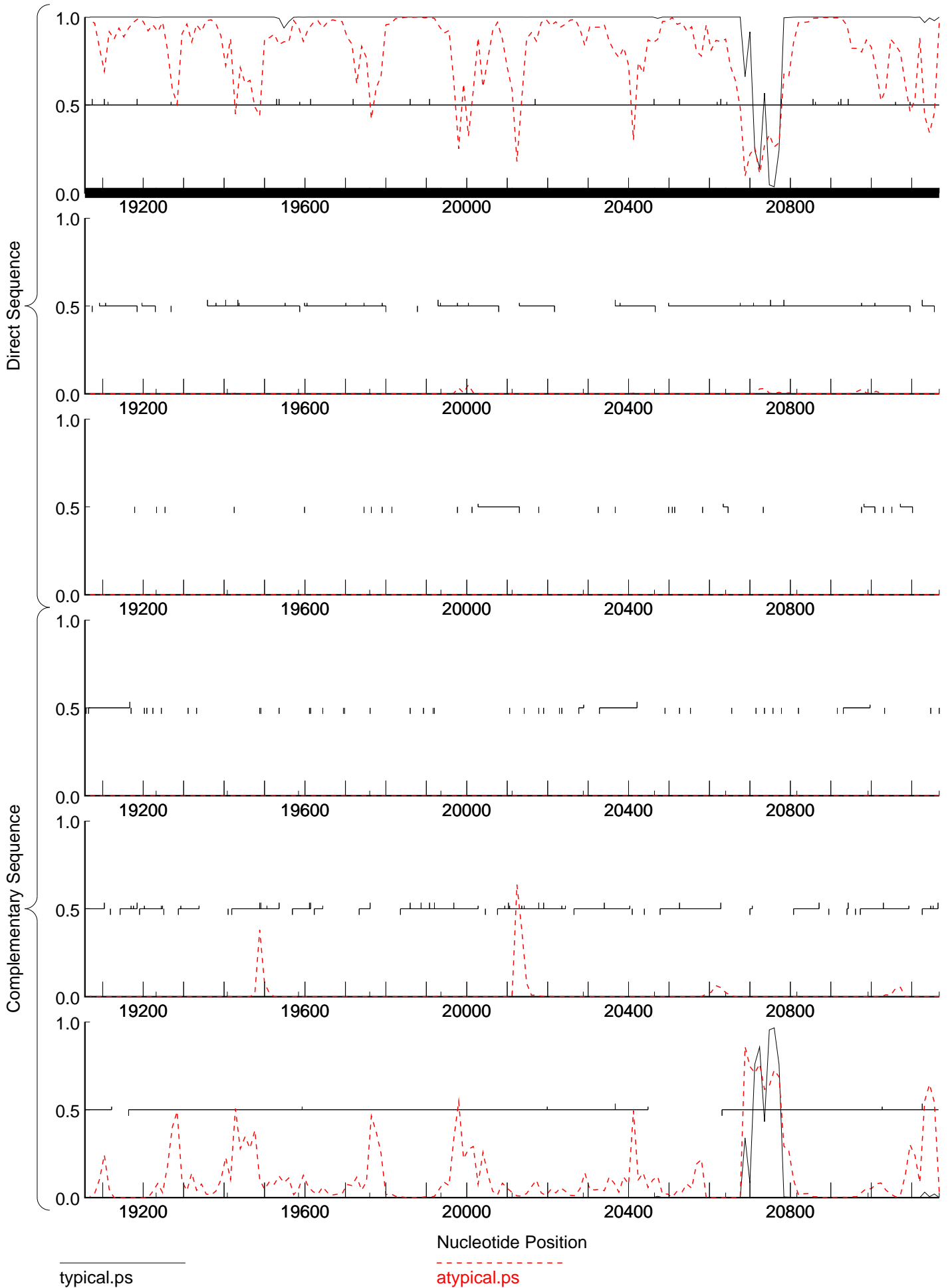


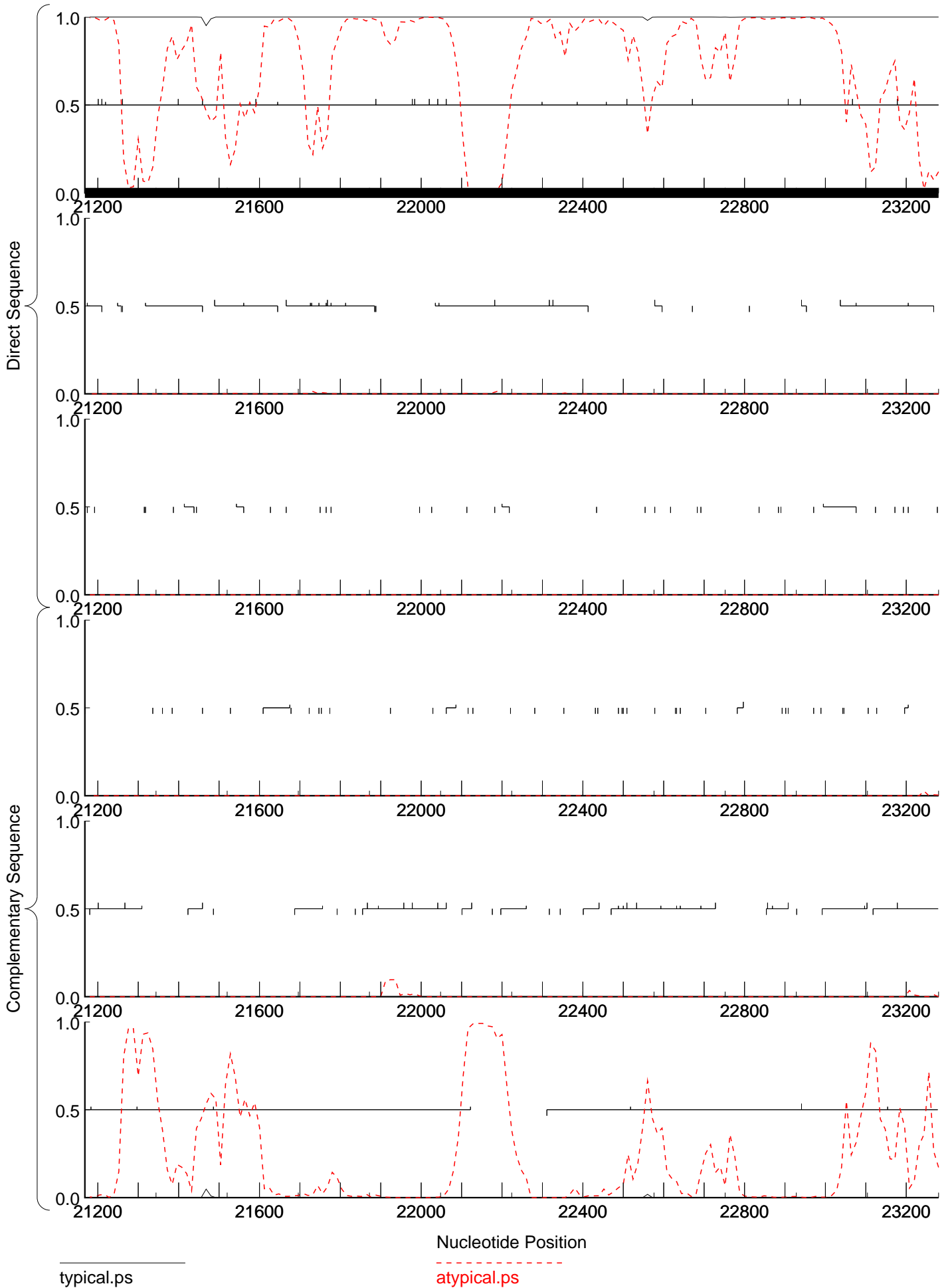


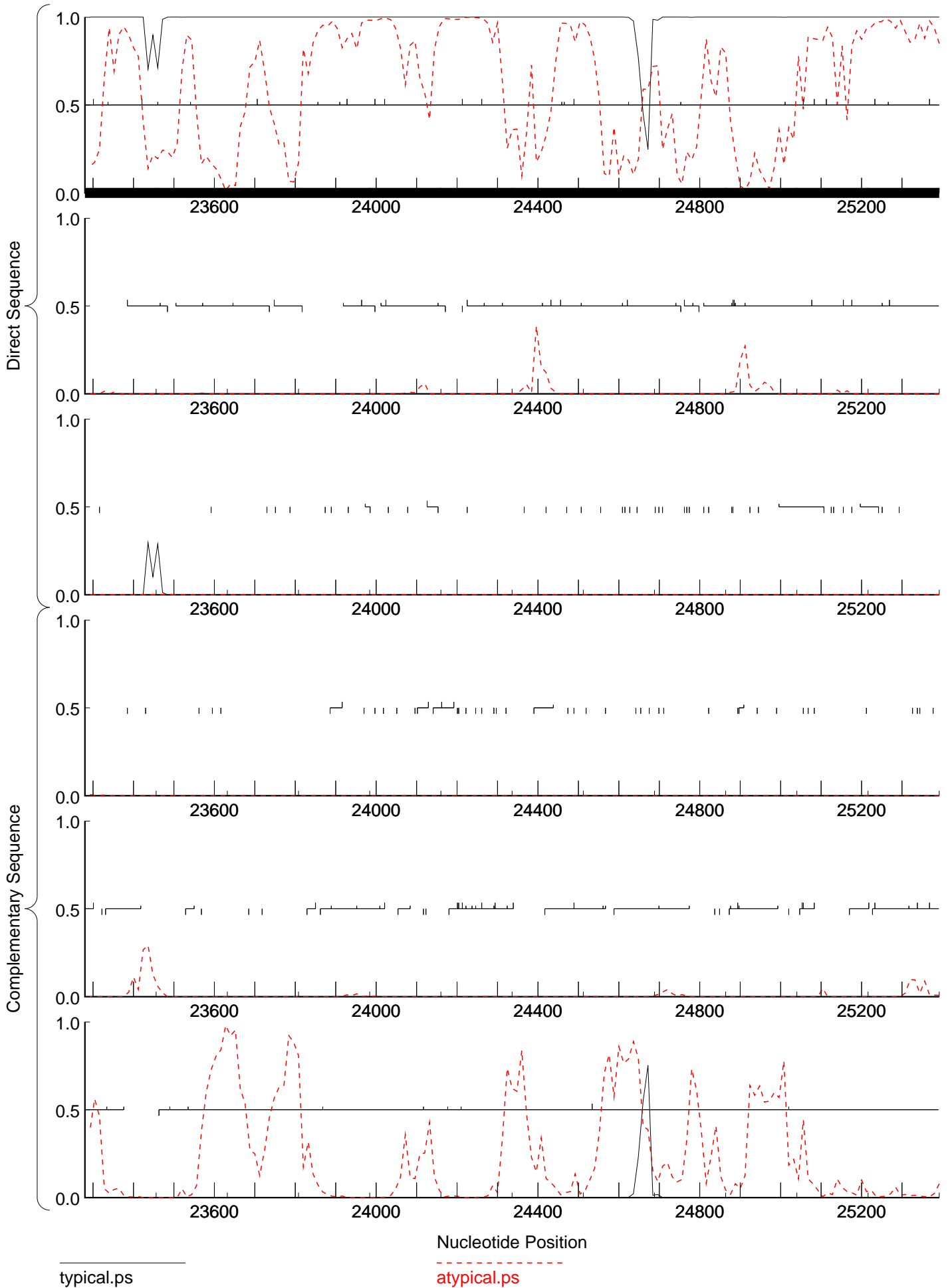
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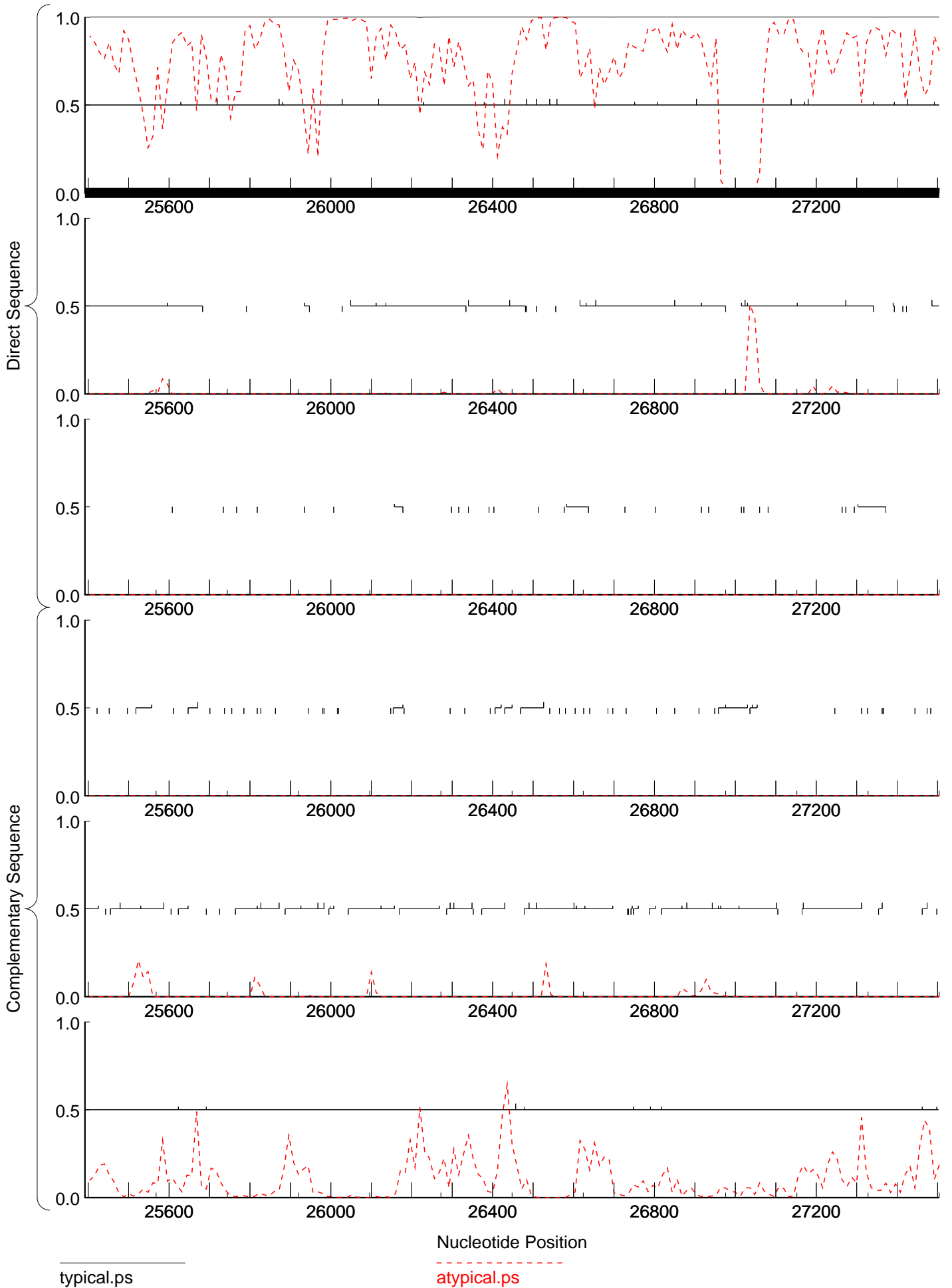
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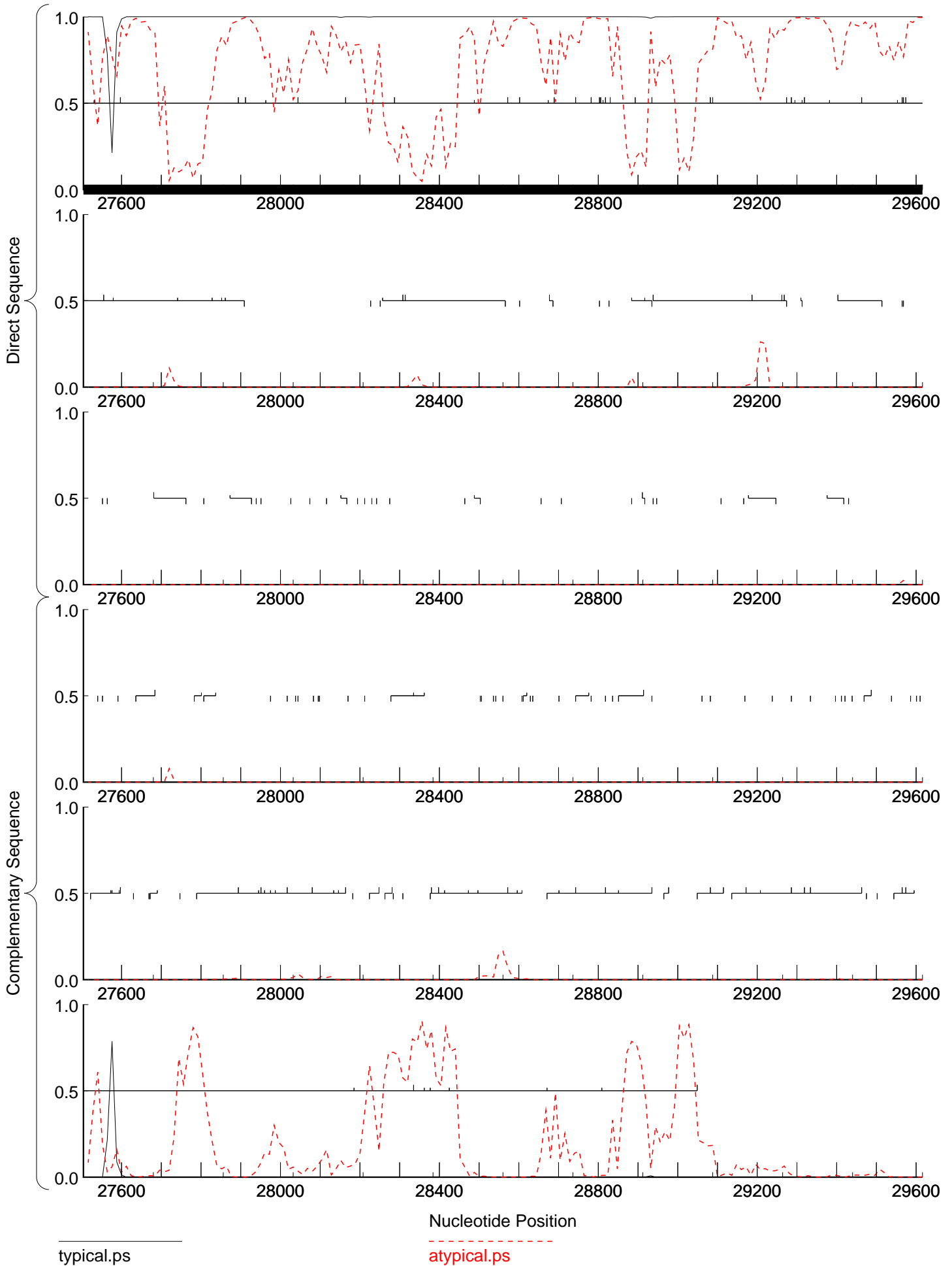


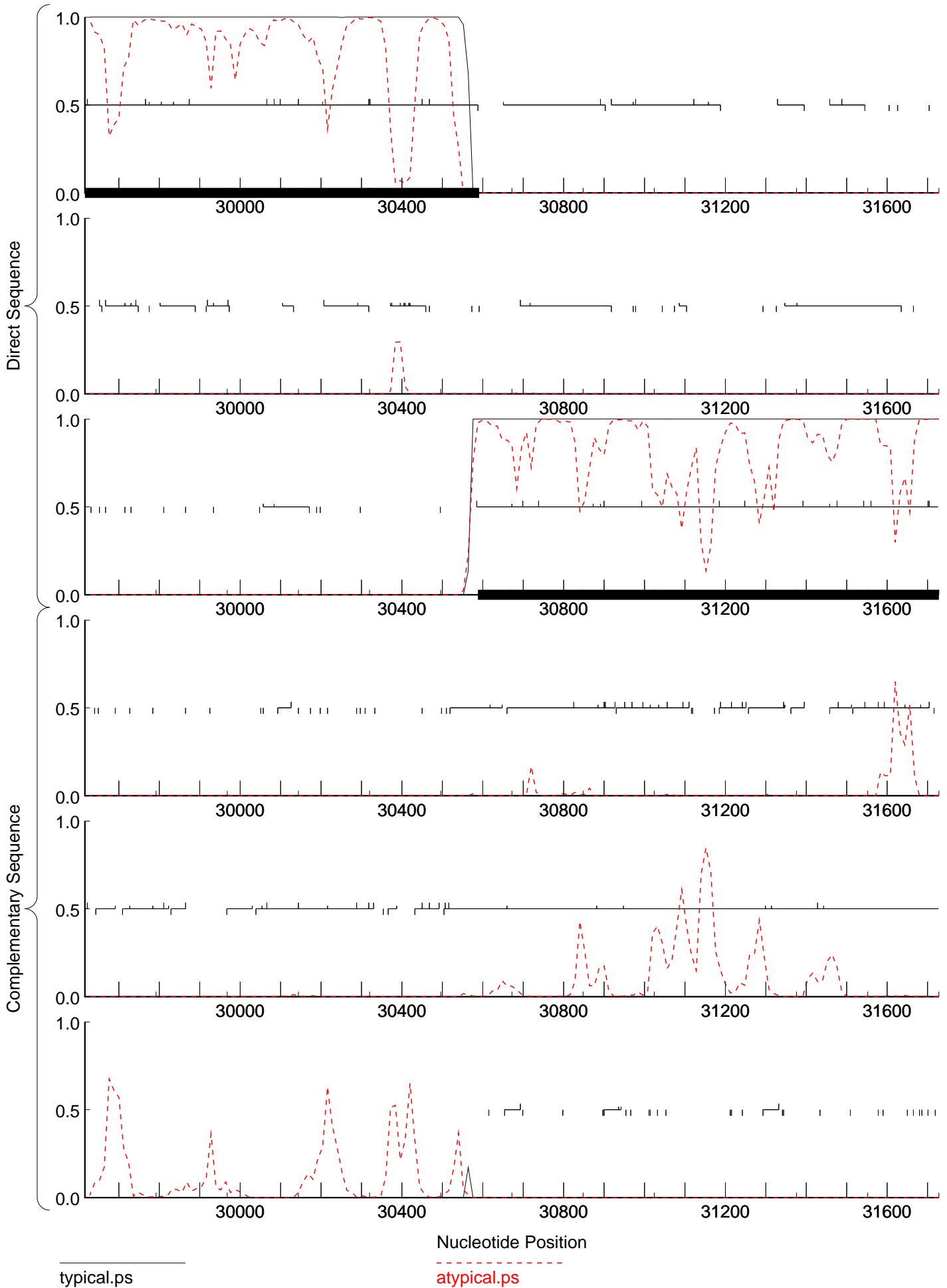


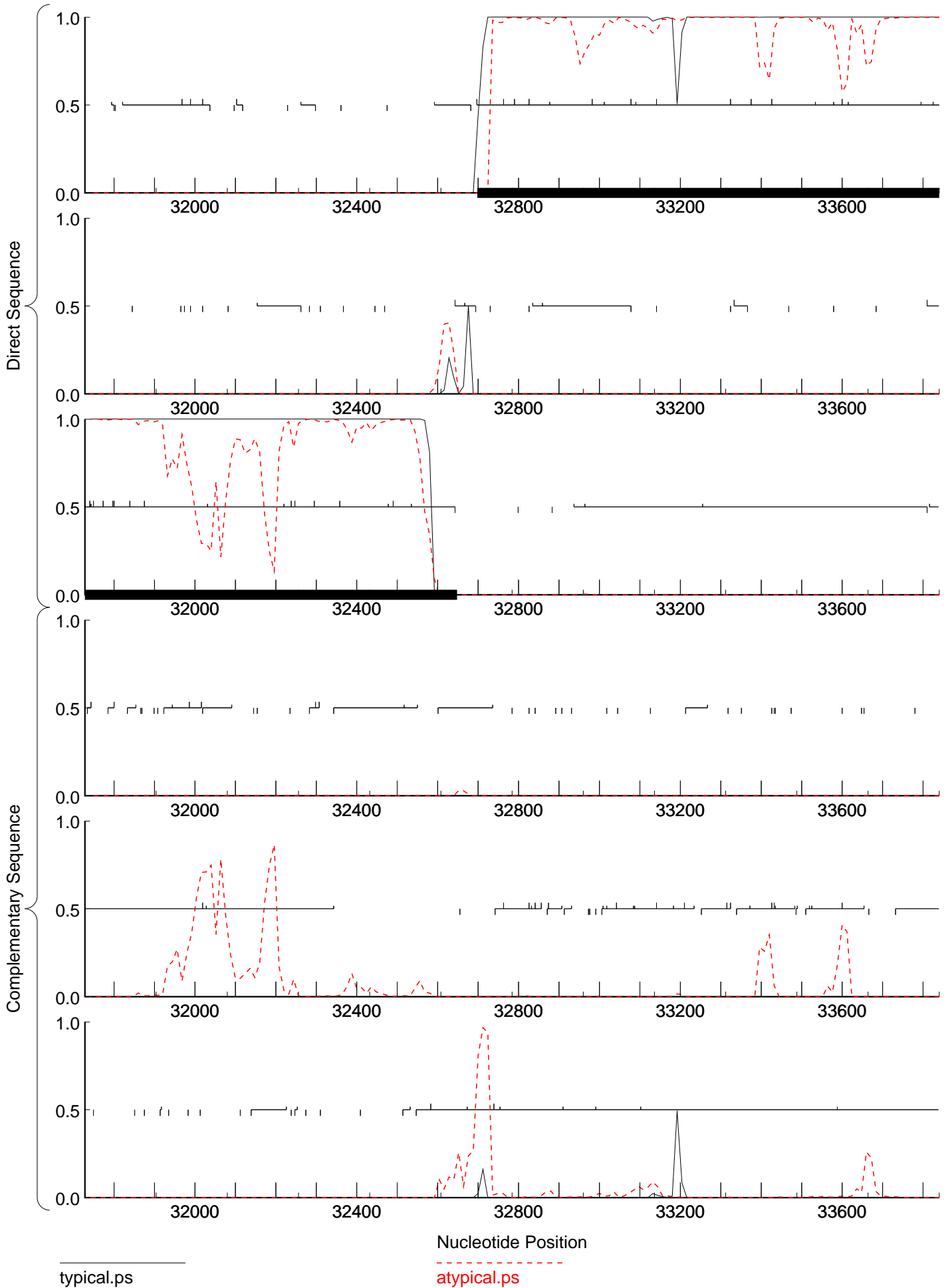


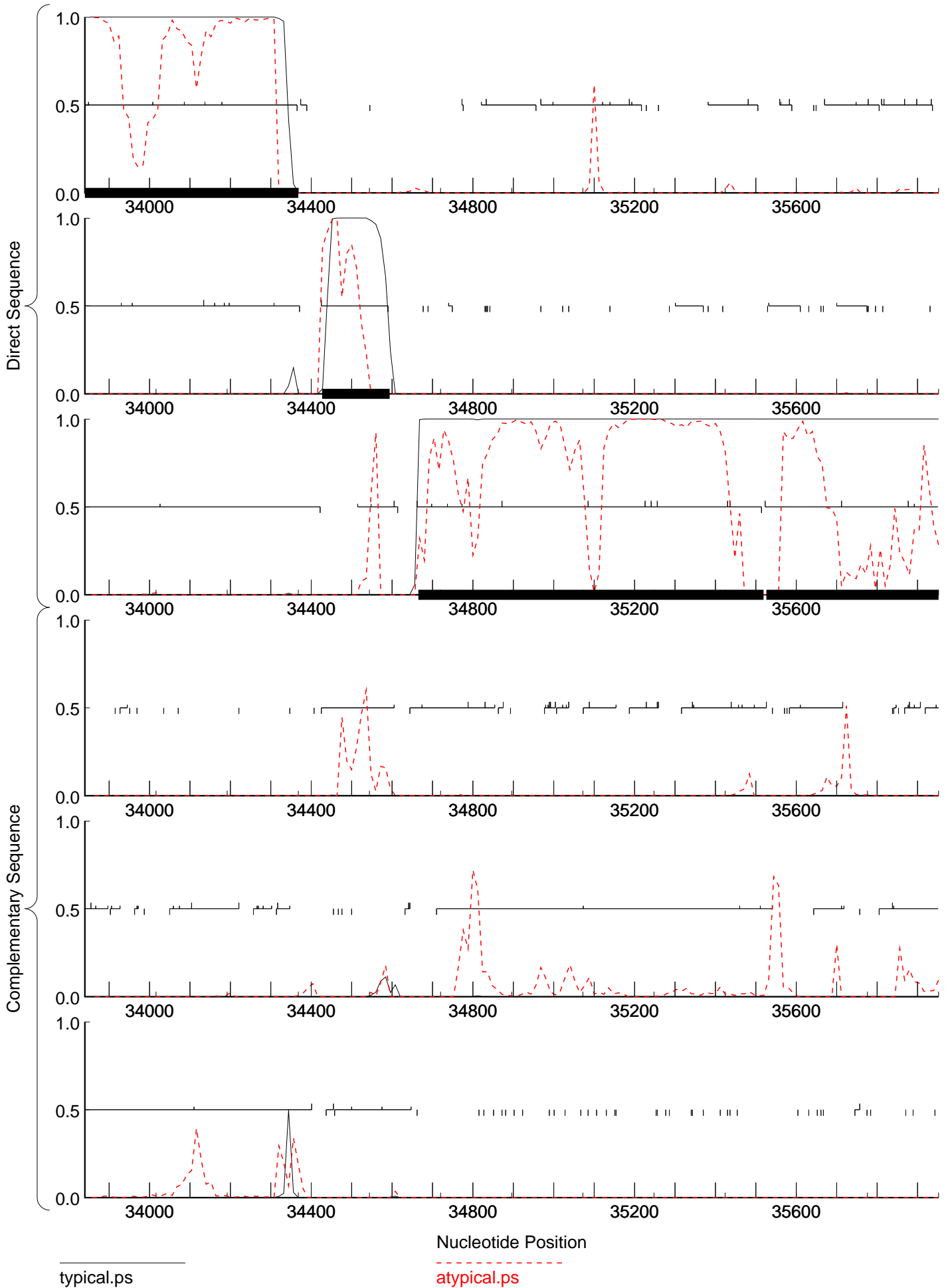


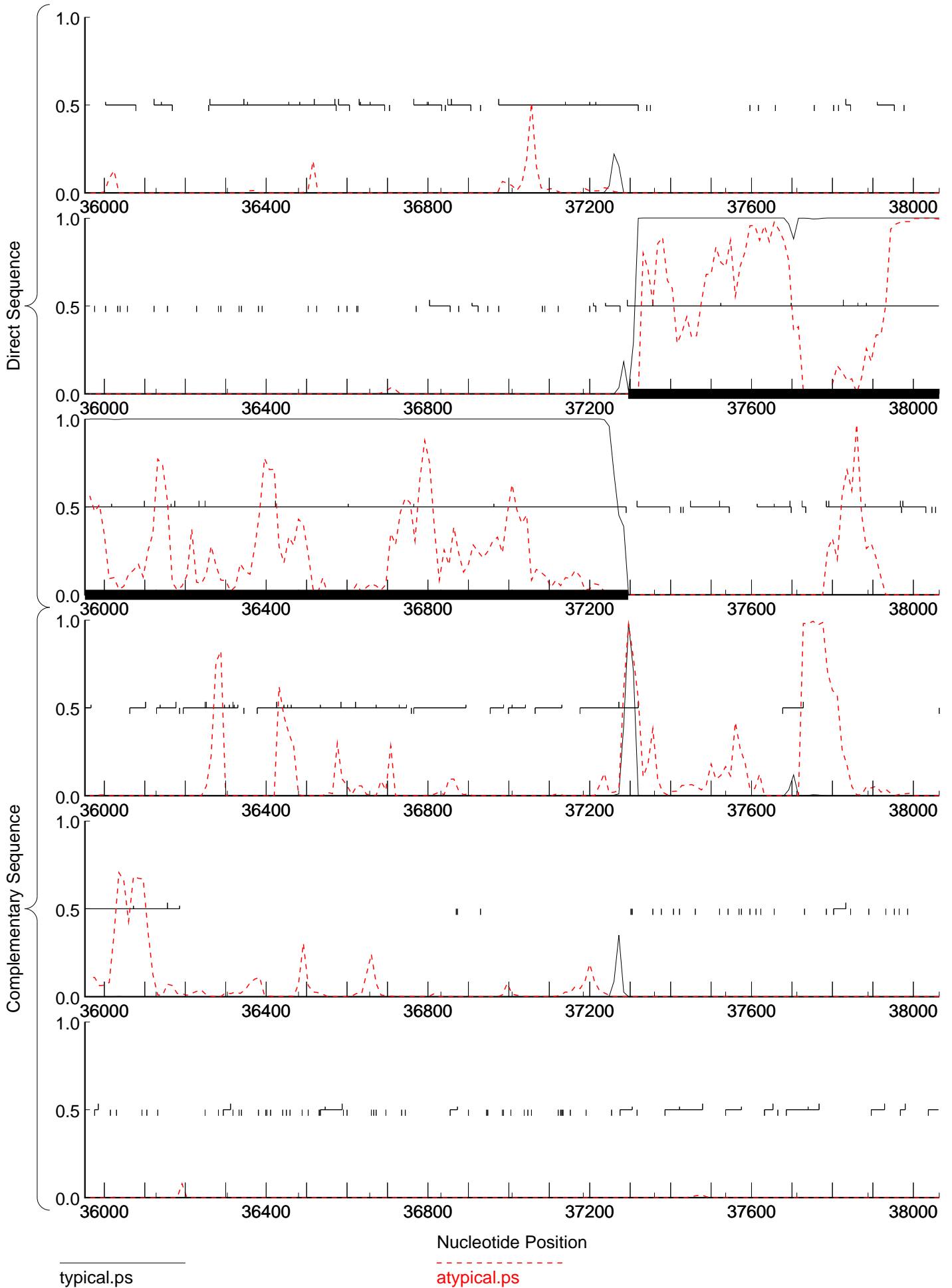


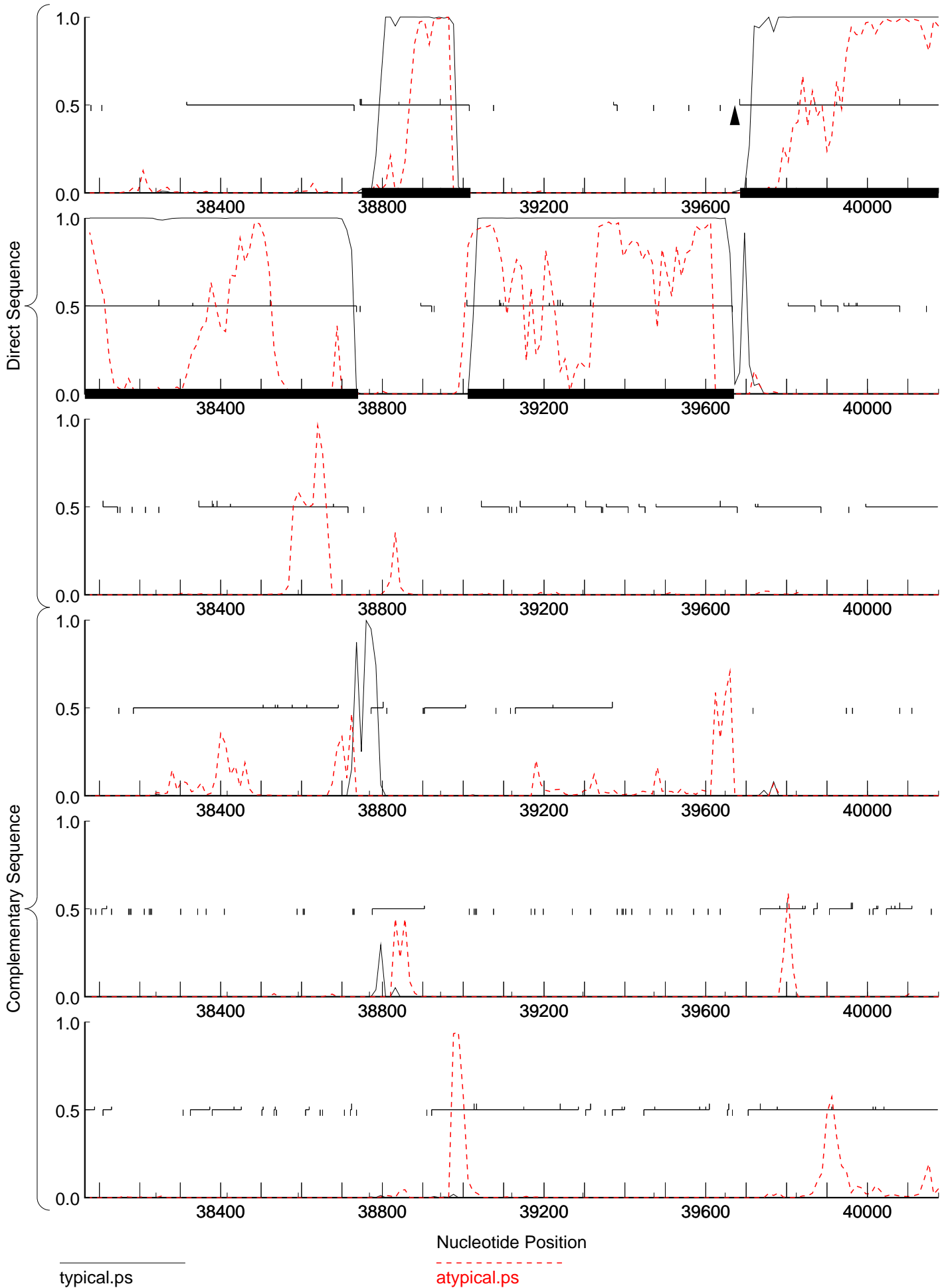


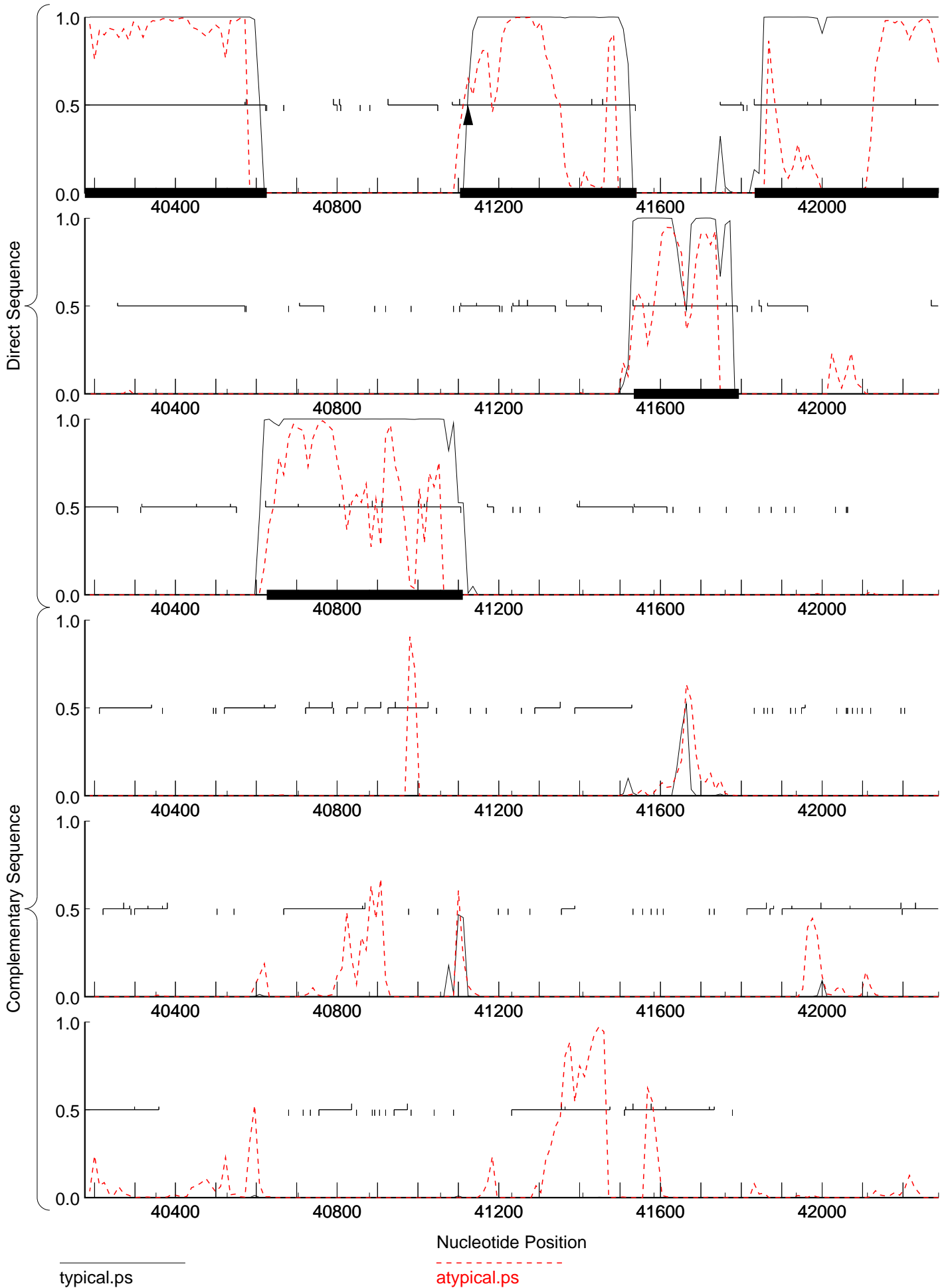


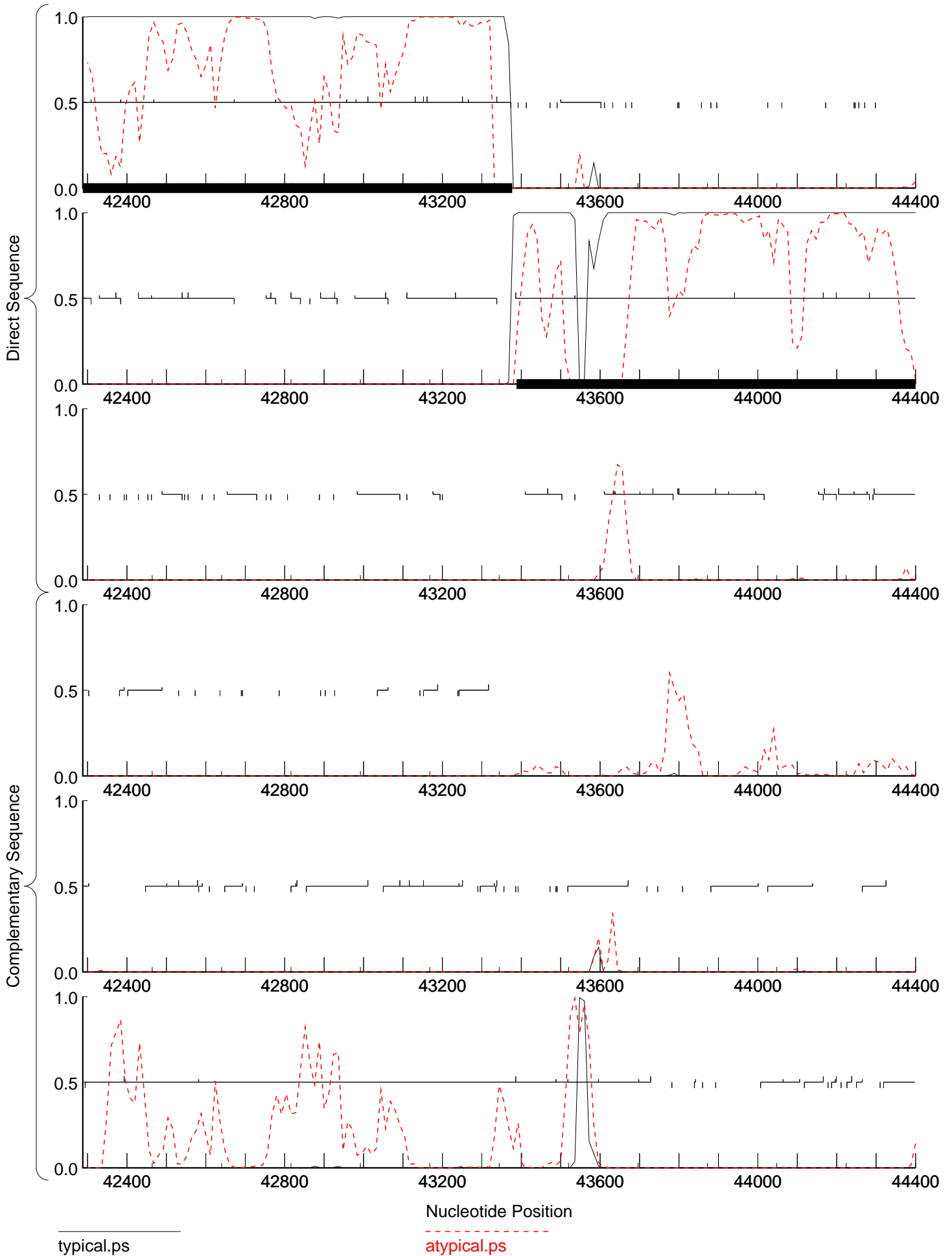


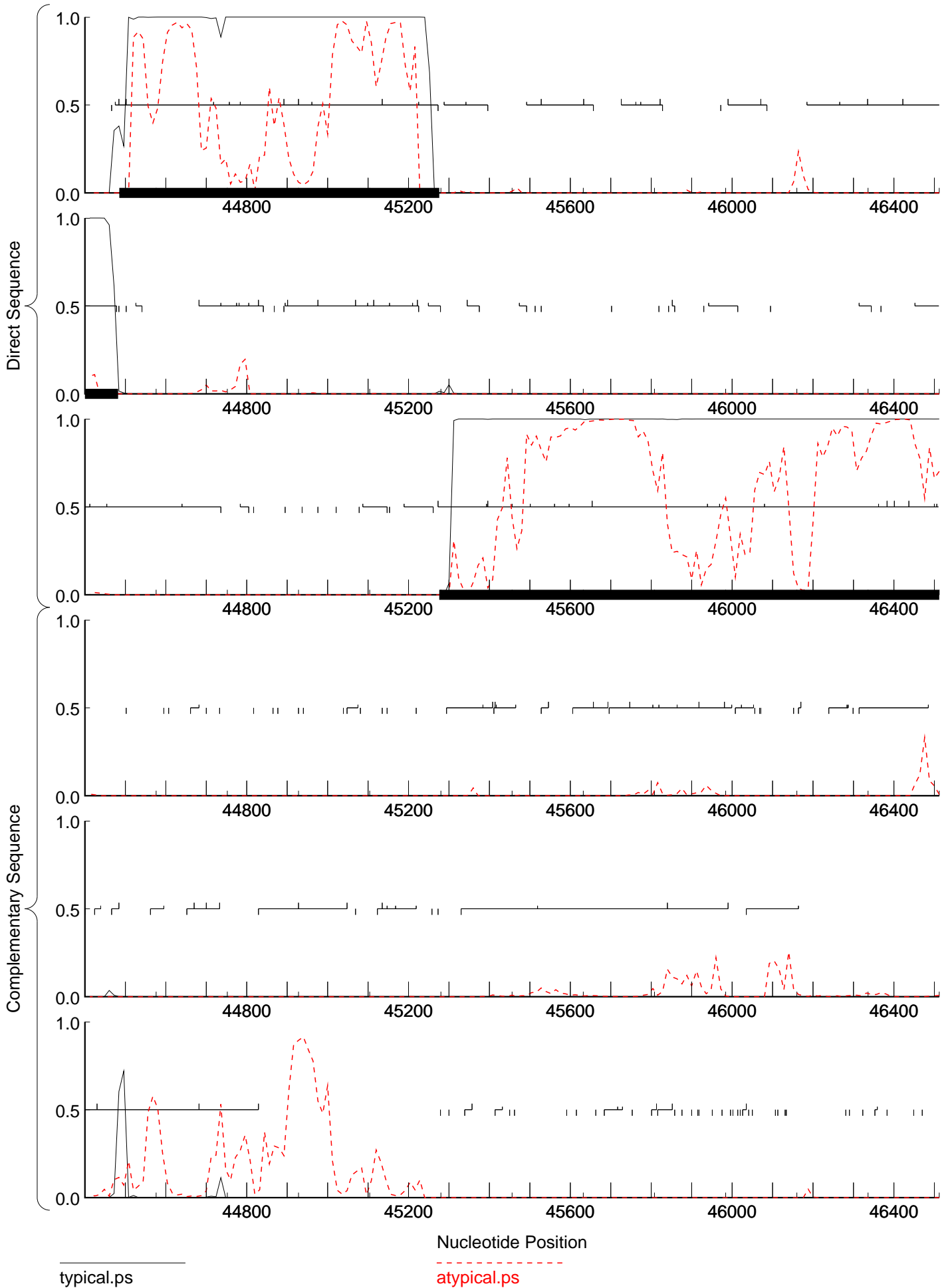


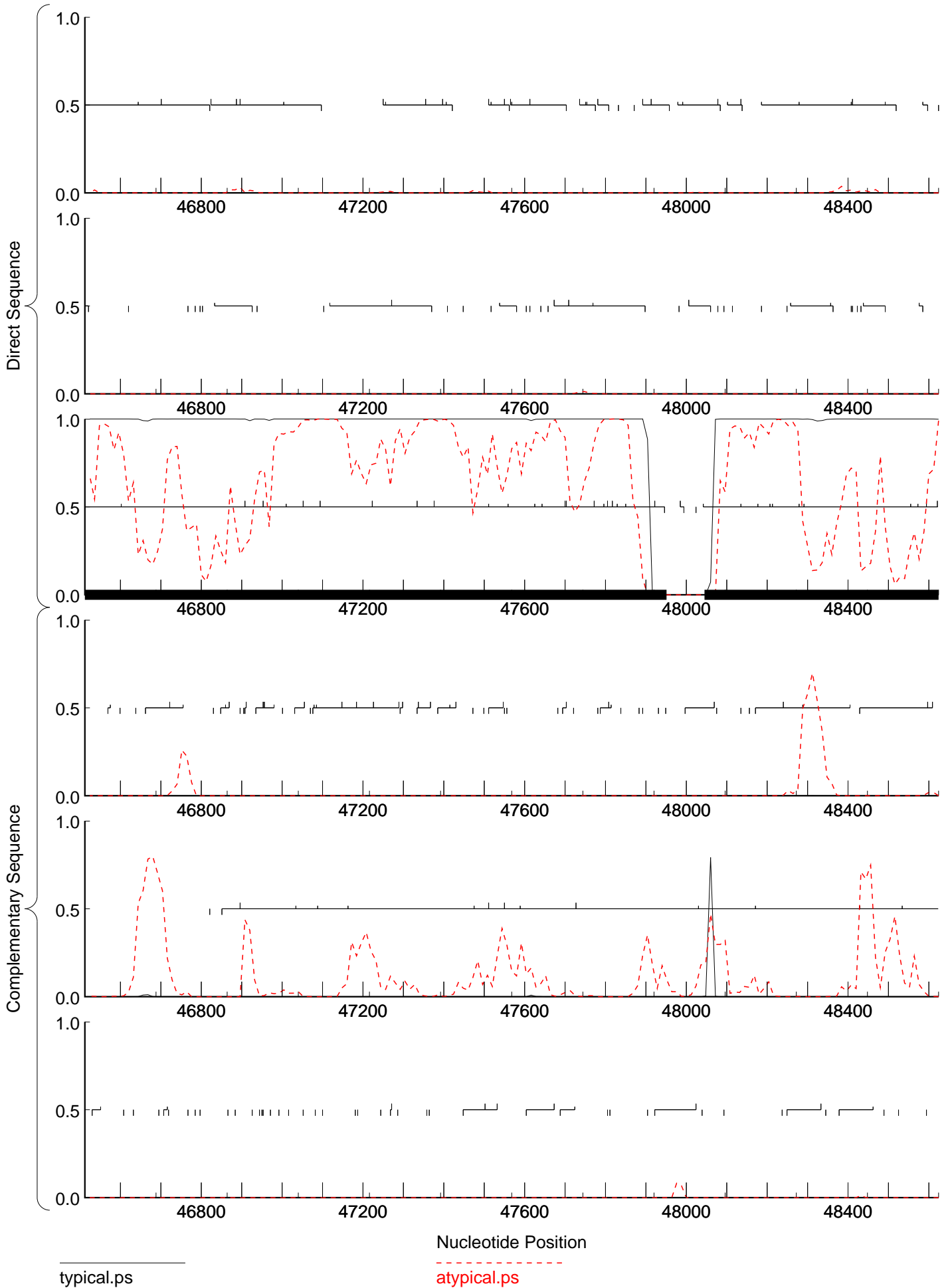


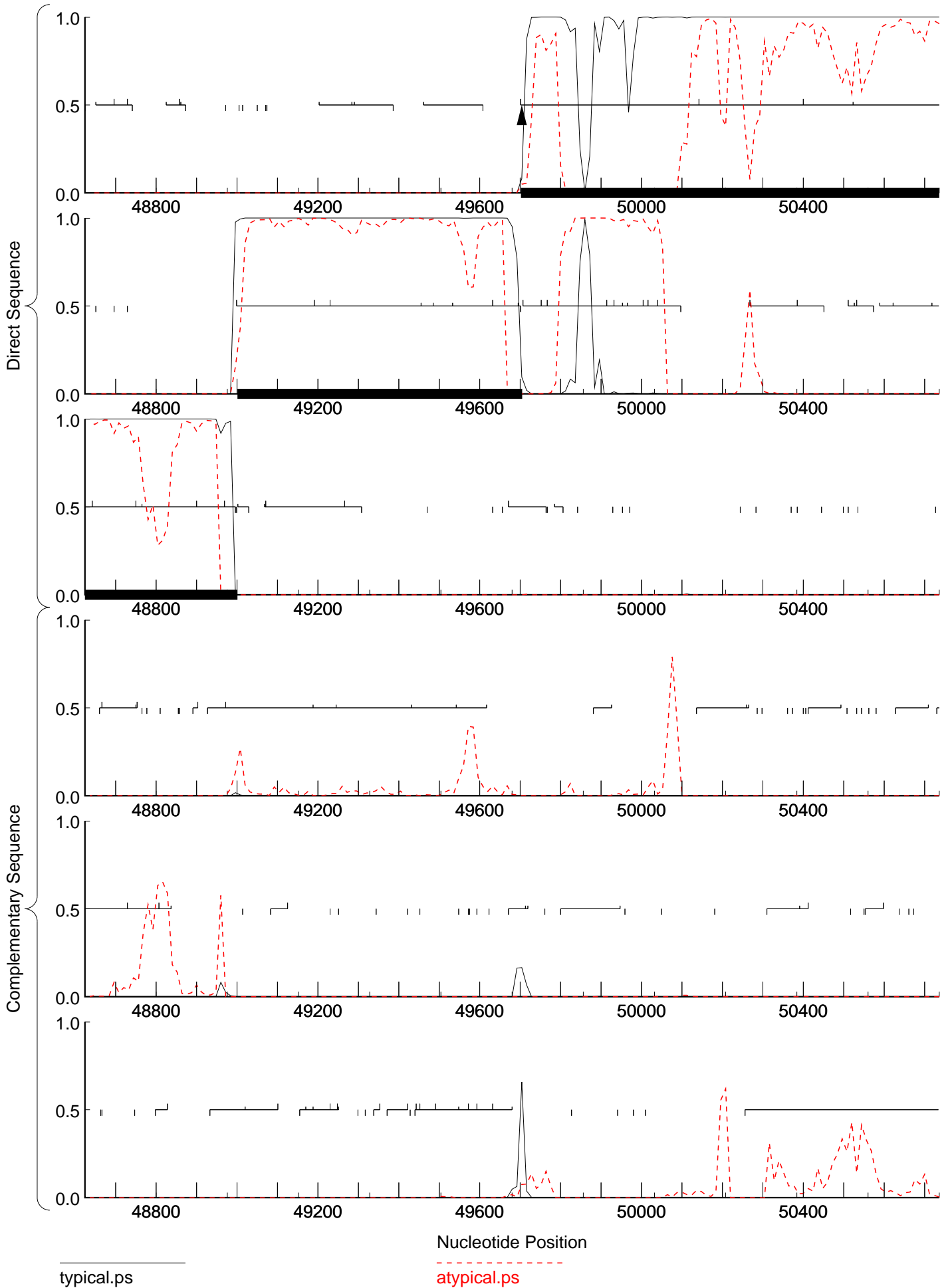


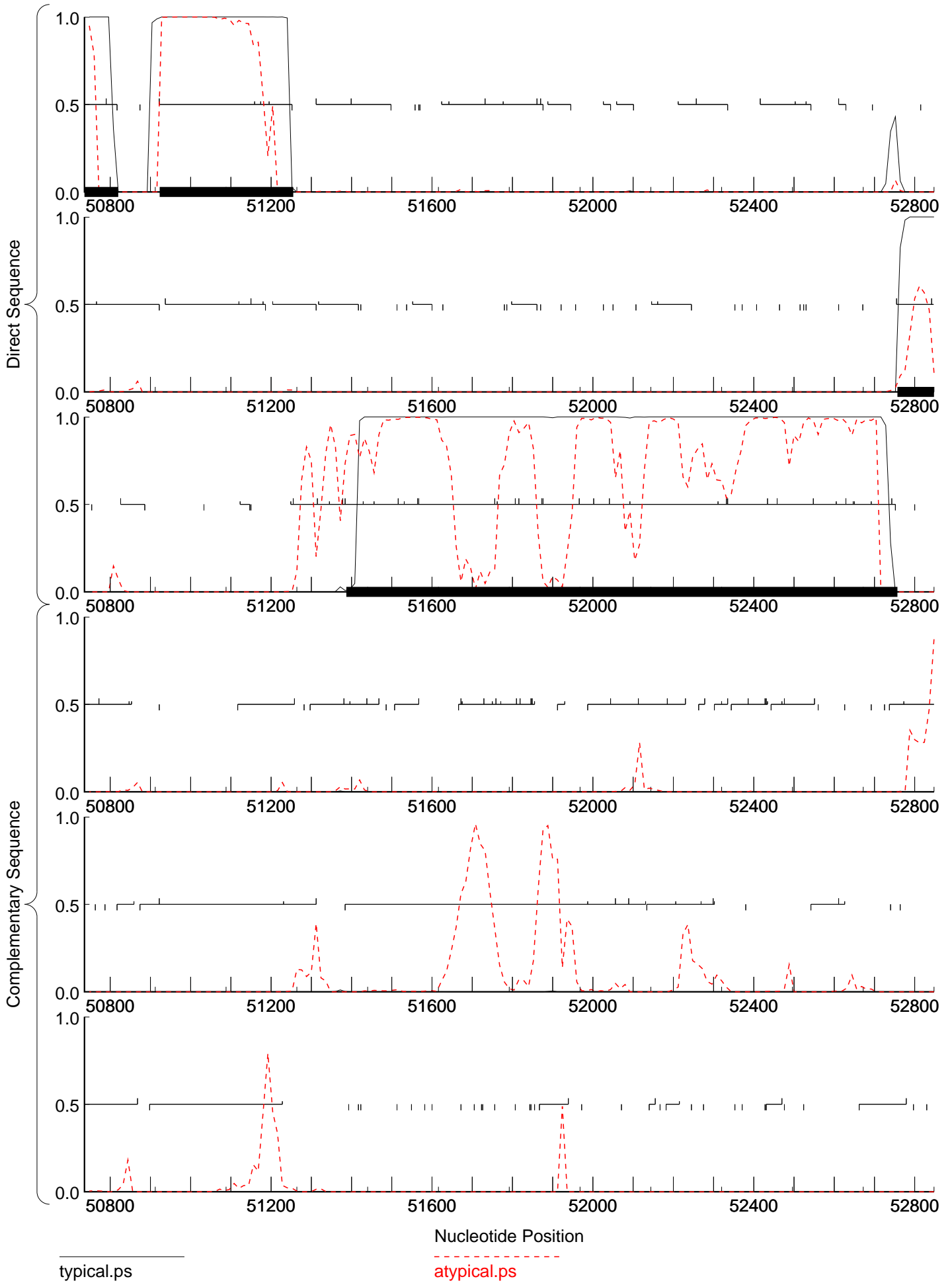


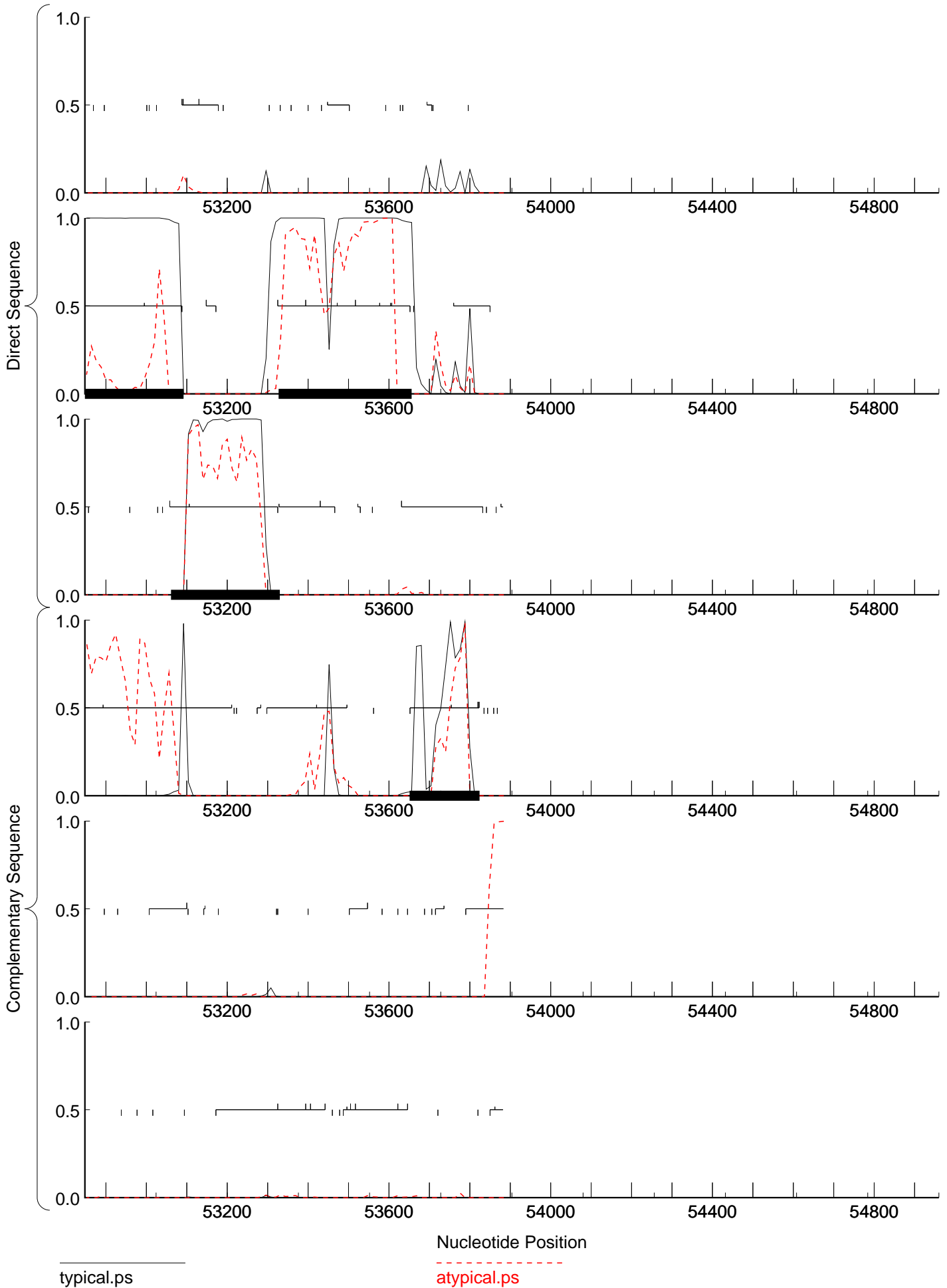












typical.ps

atypical.ps