

GeneMark

Version 2.5p (09.08.06)

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PROGRAM INFORMATION

Sequence : Arthrobacter phage Maureen complete sequence, 45802 bp including 11-base 3' overhang (CGGAG
Analysis Date : 8/25/19 at 18:17:39
Pages : 23
Sequence Length : 45802 bp
GC Content : 68.82%

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,

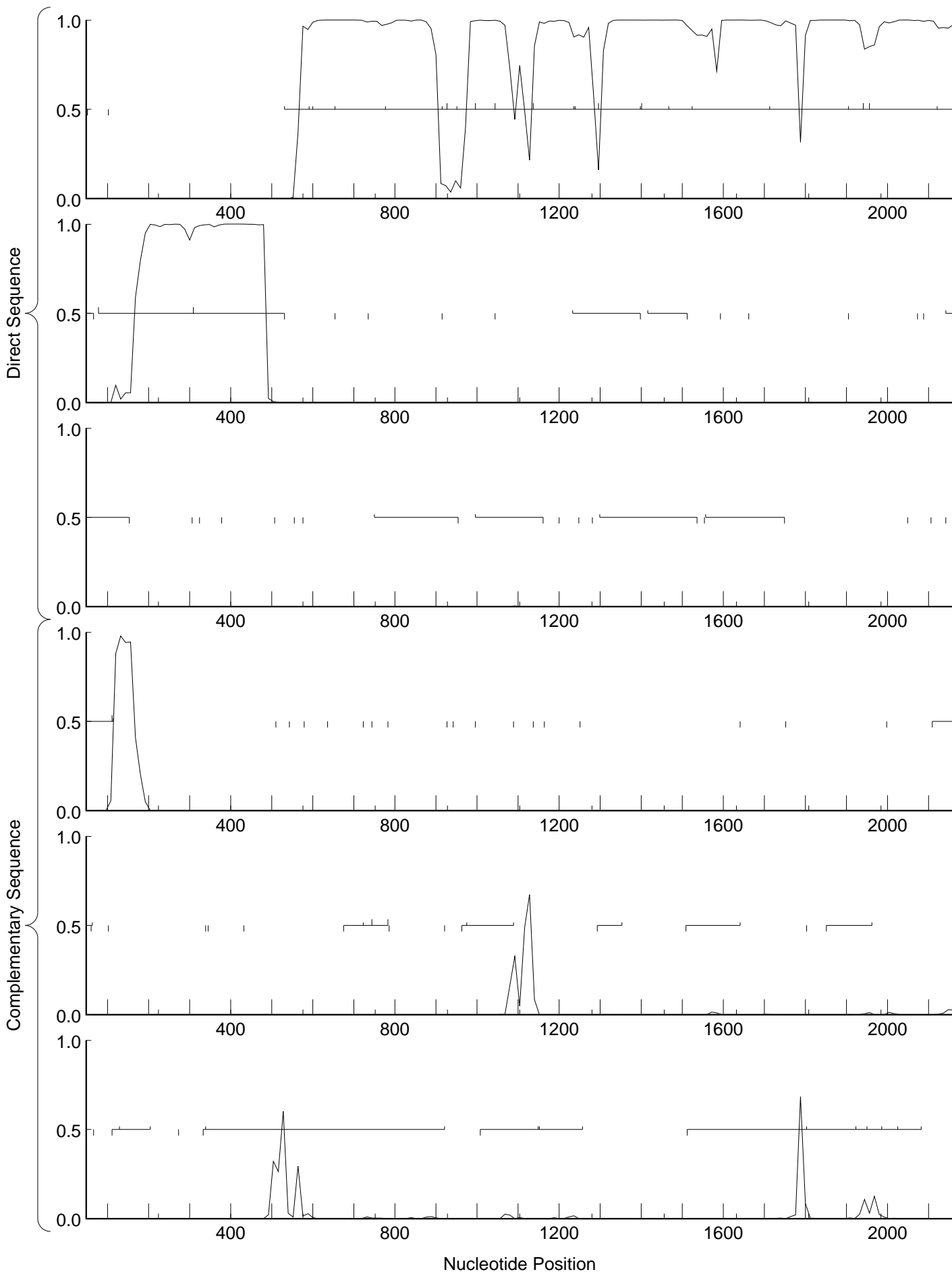
Matrix notes & comments

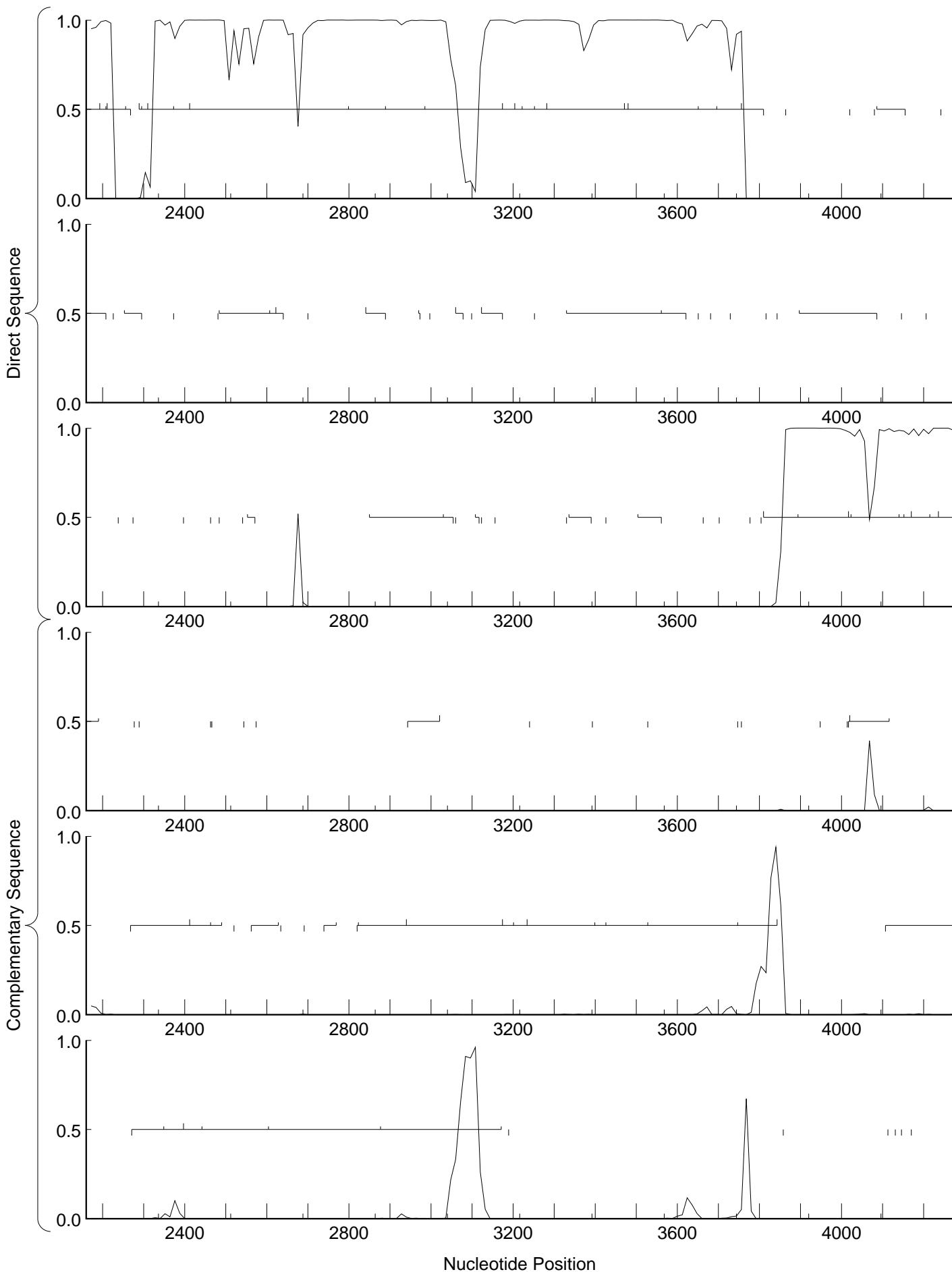
Training set derived by GeneMarkS, 4.30 October 2014
Sat Sep 8 00:22:00 2018

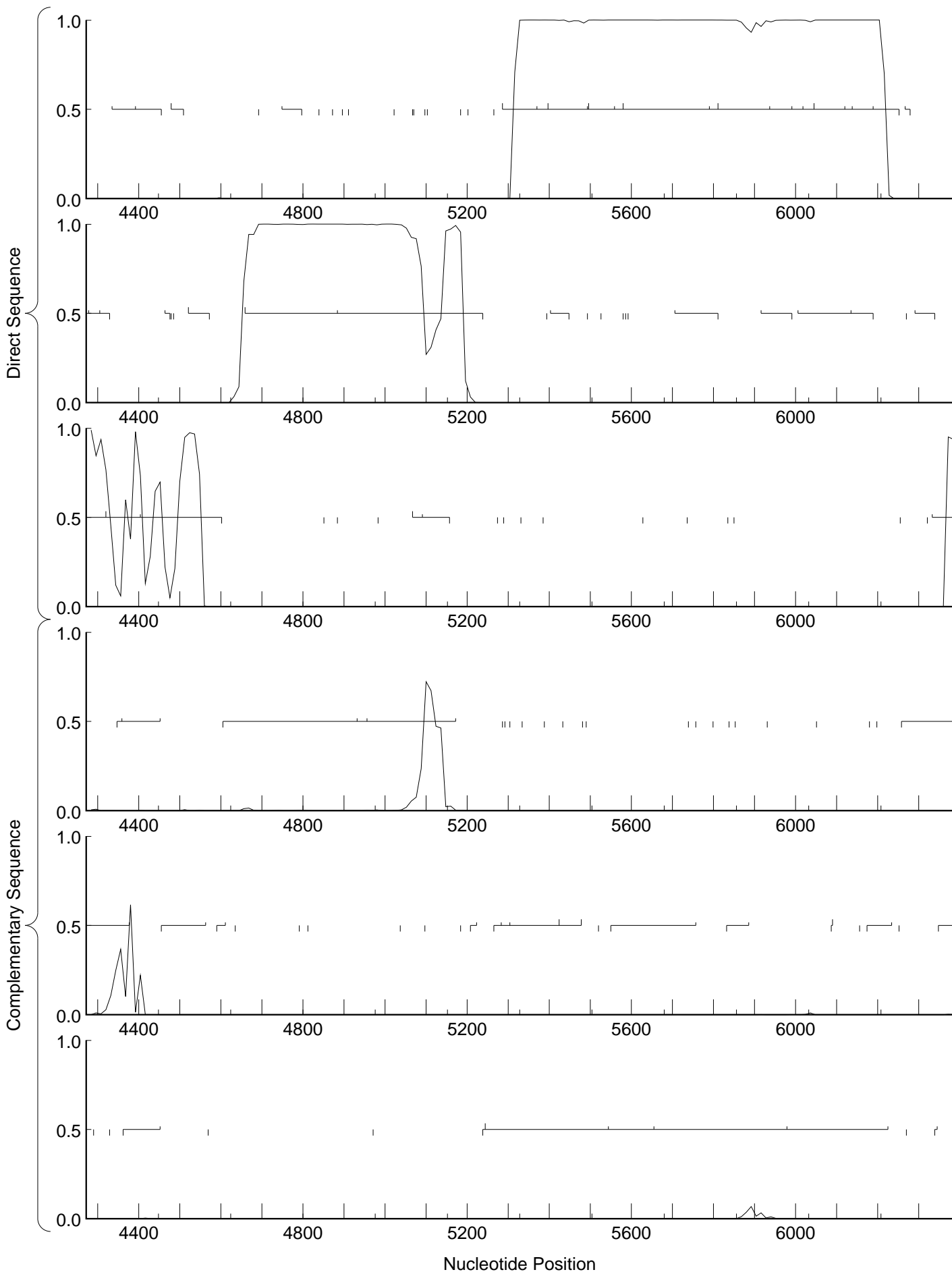
MATRIX INFORMATION

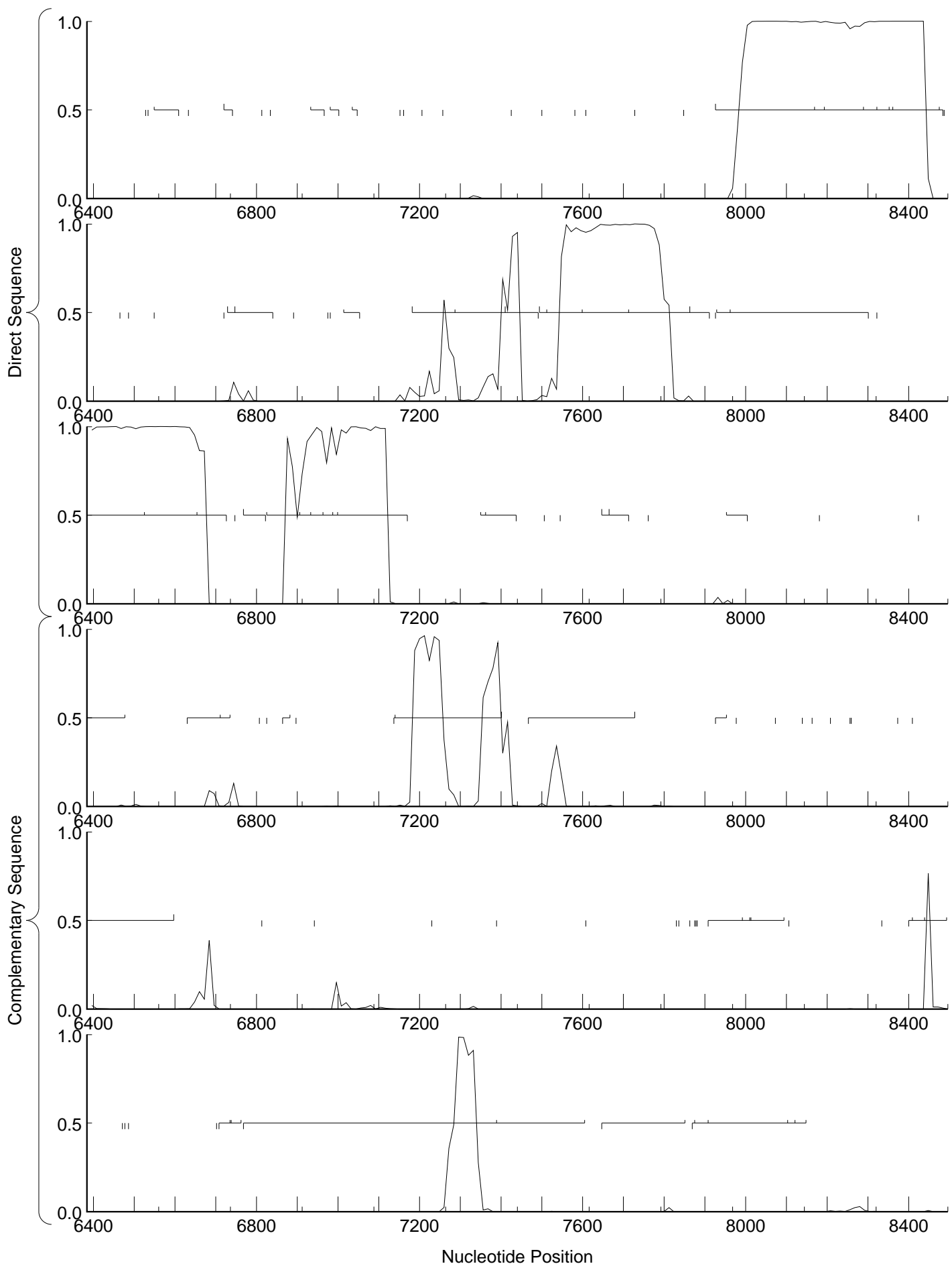
Matrix : -
Author : -
Order : 4

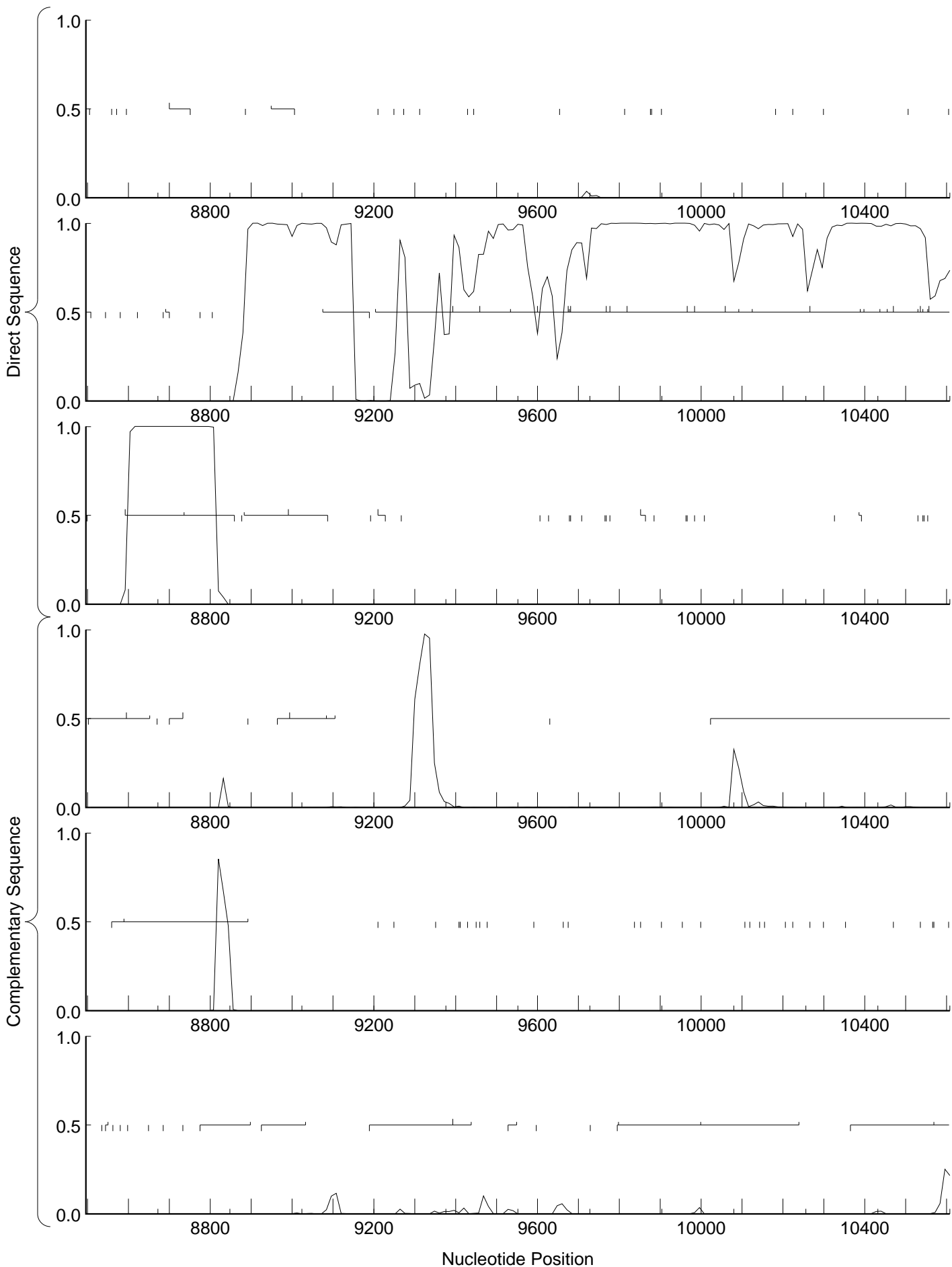
Send questions / comments to:
Dr. M. Borodovsky
Georgia Institute of Technology
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Atlanta, GA 30332-0230

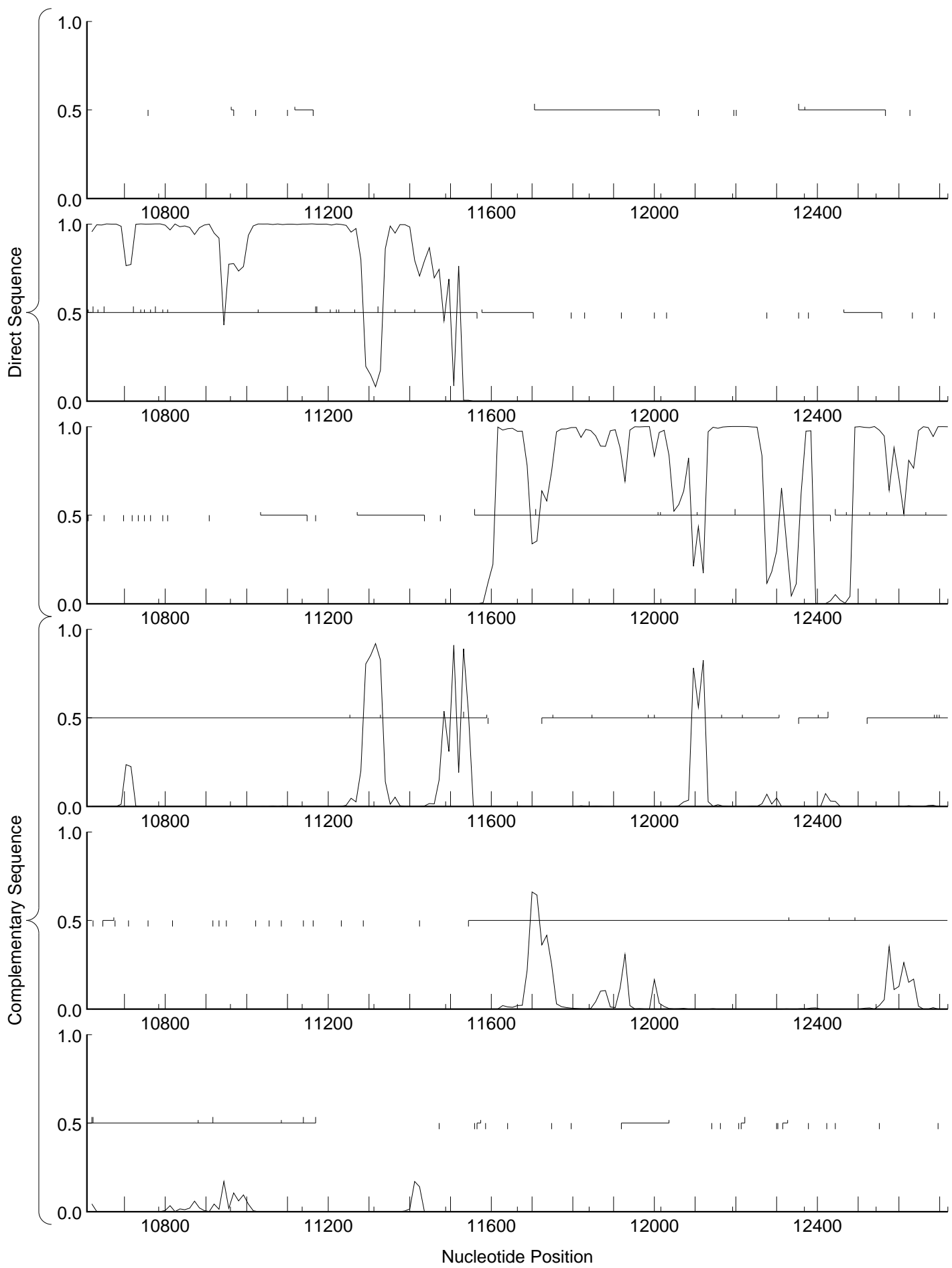


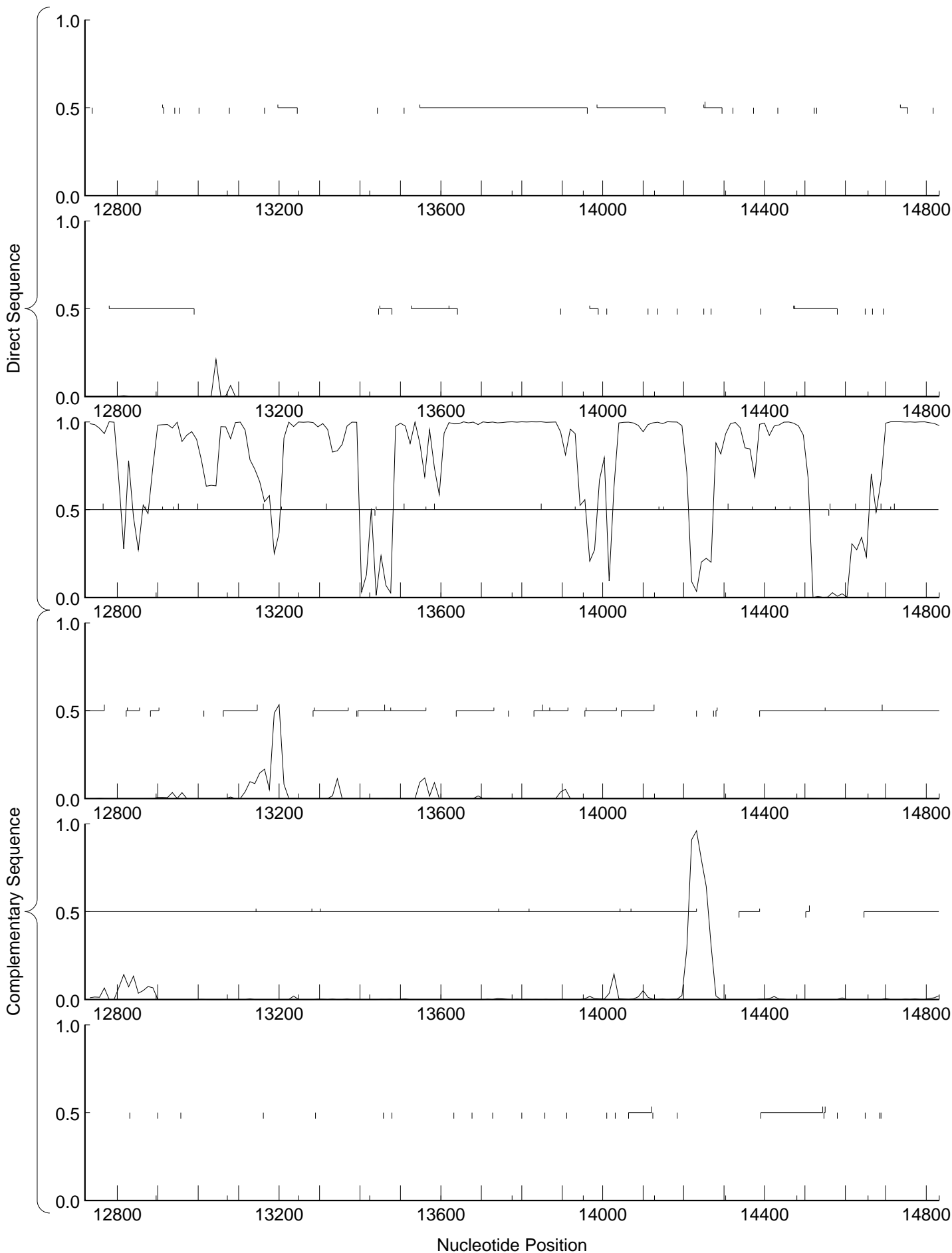


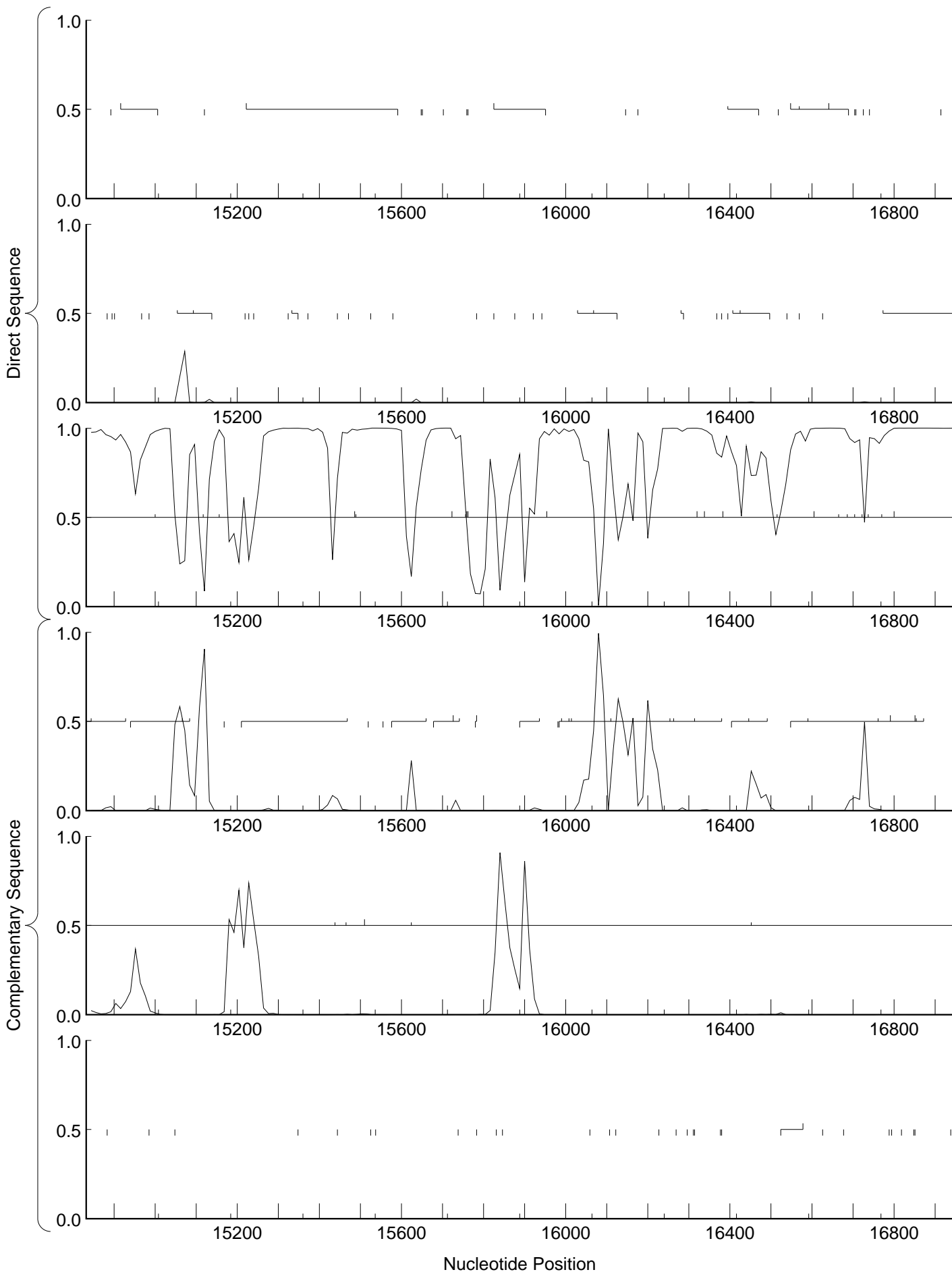


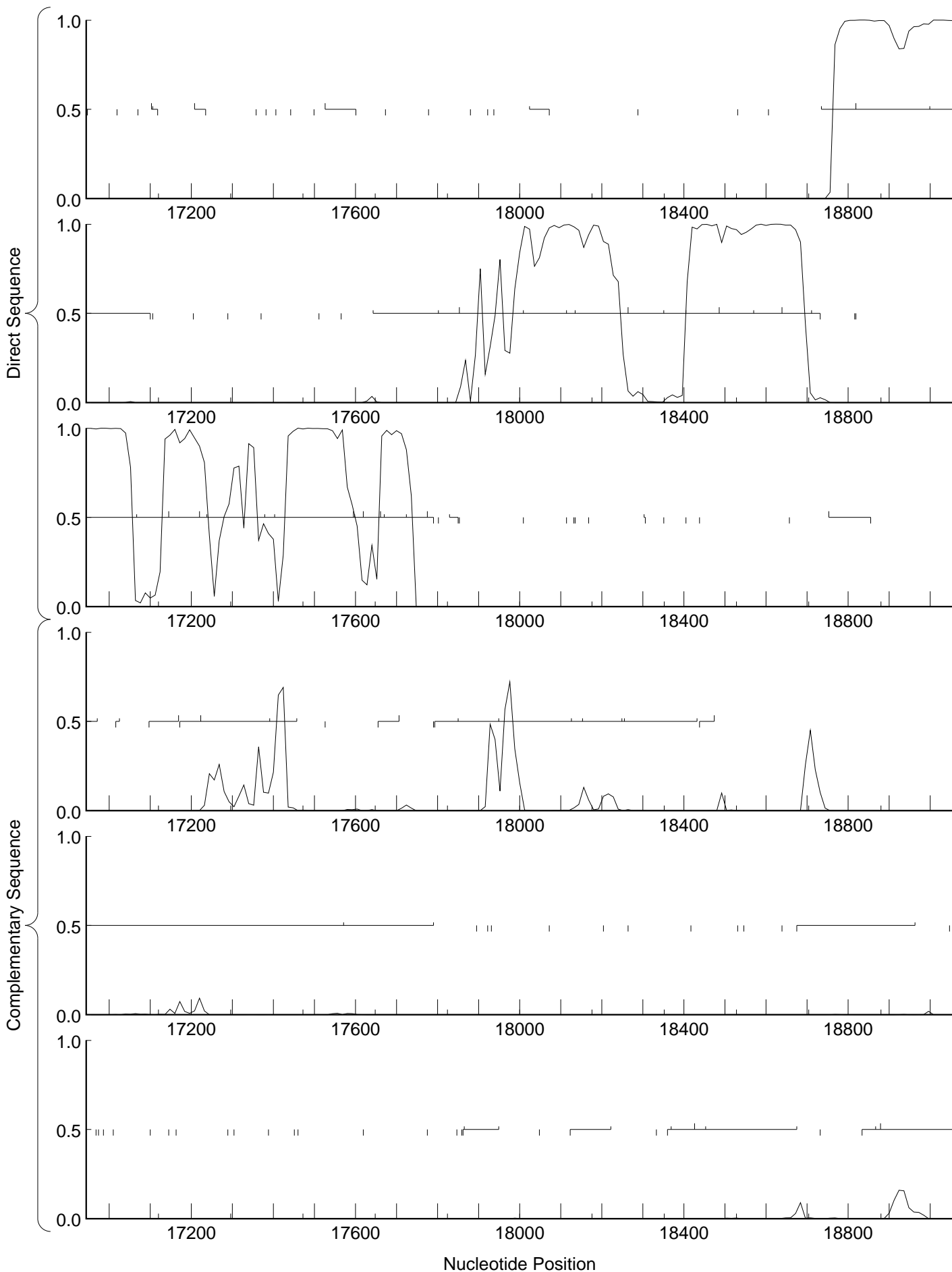


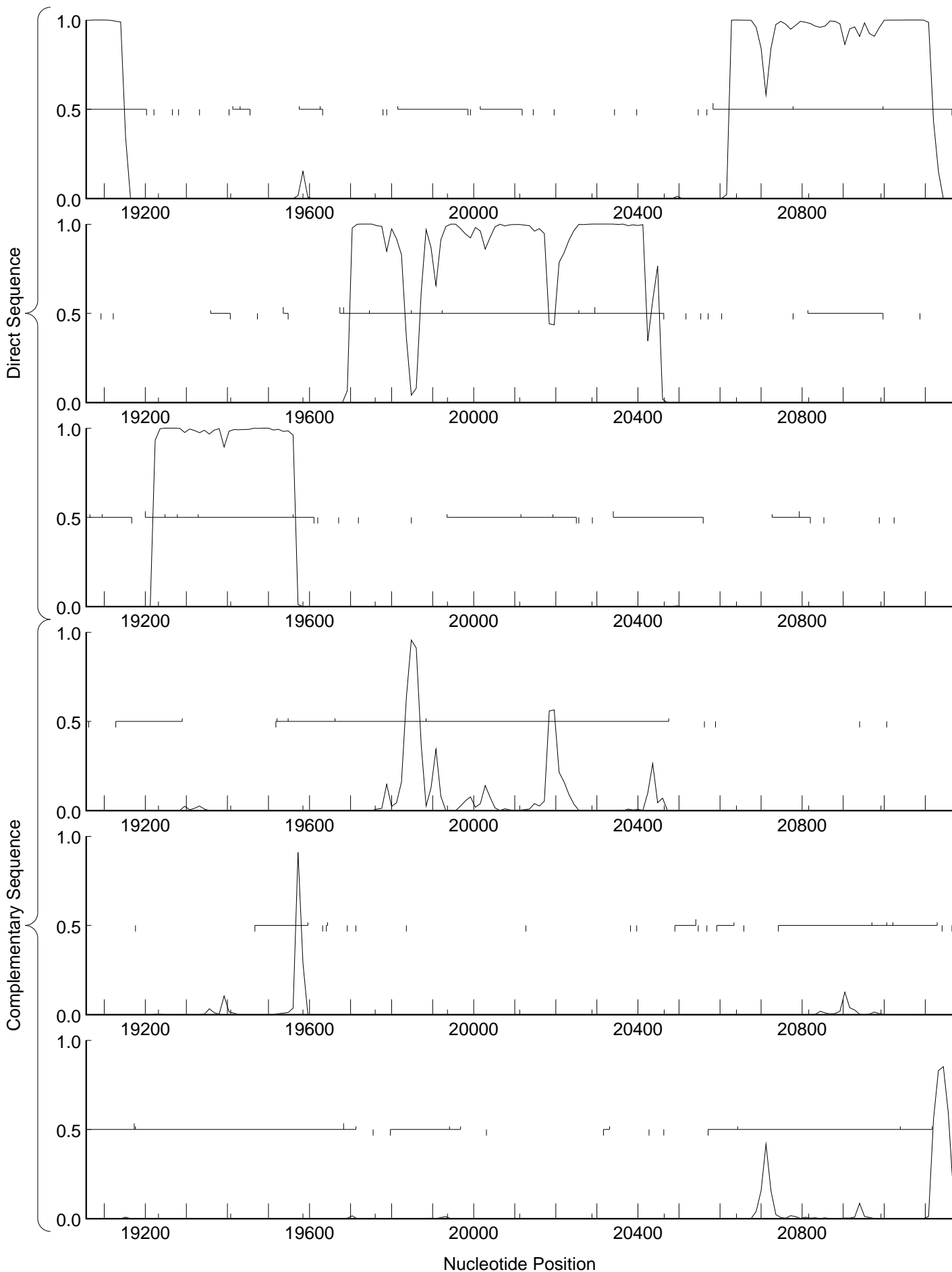


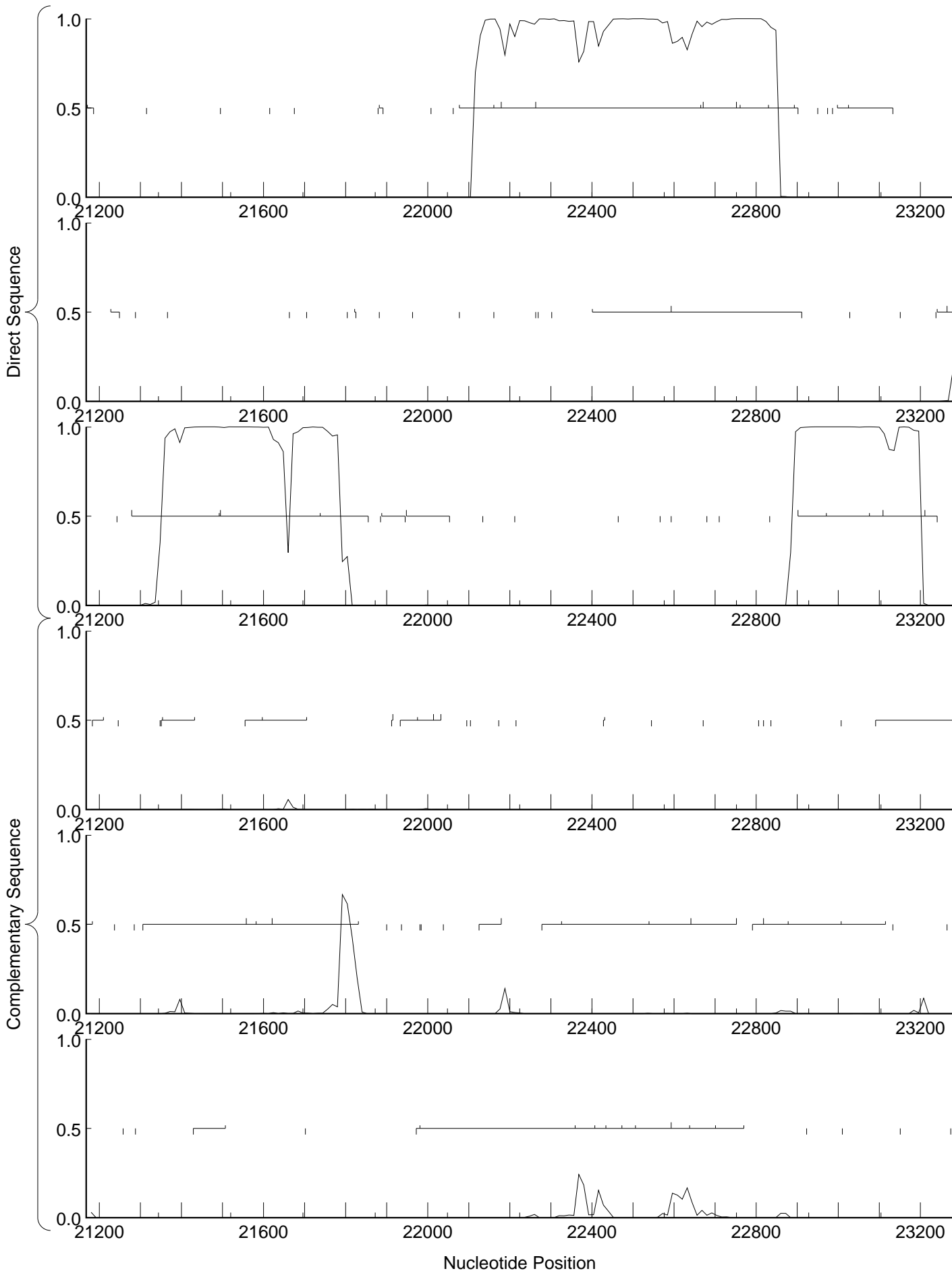


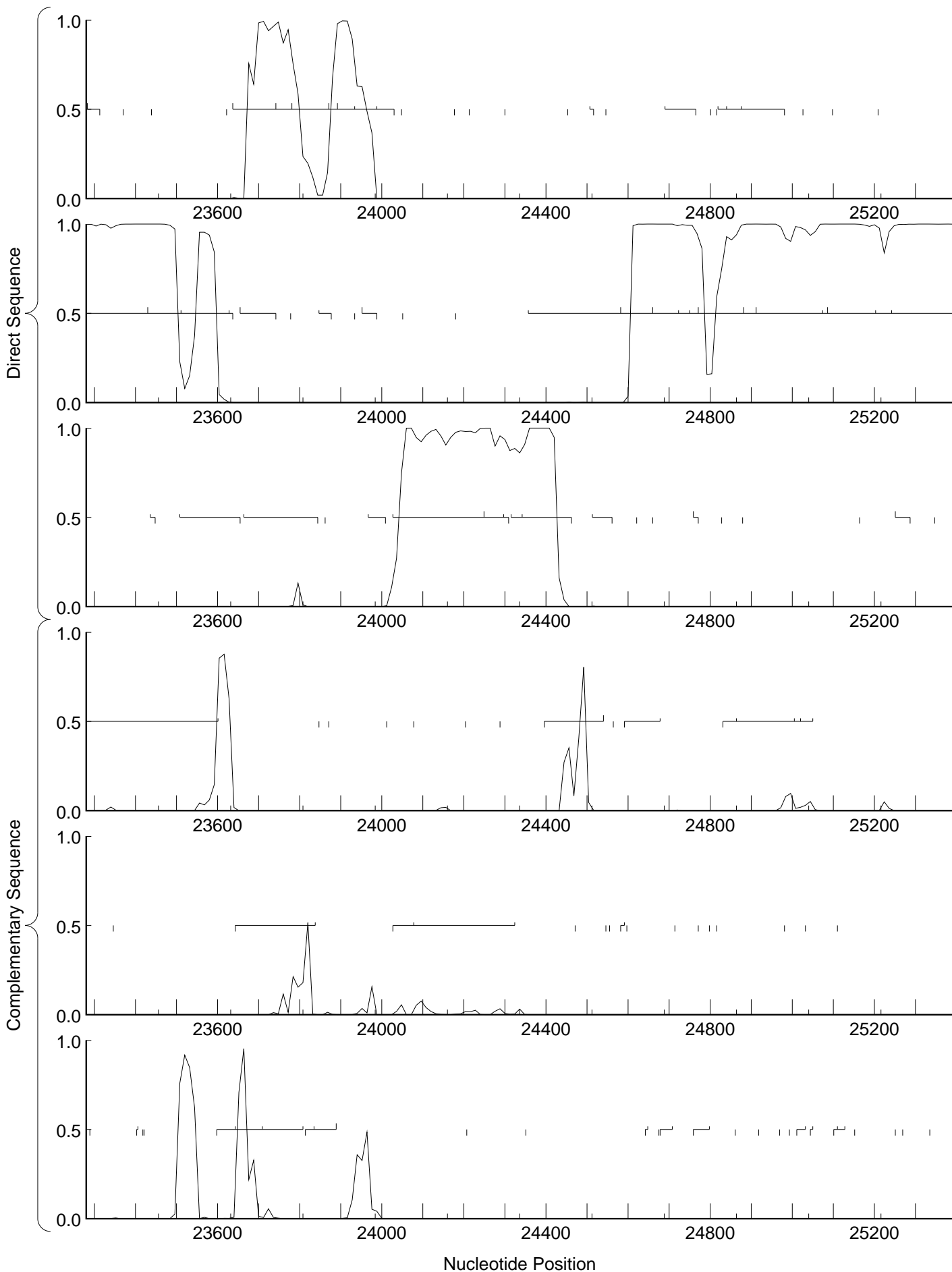


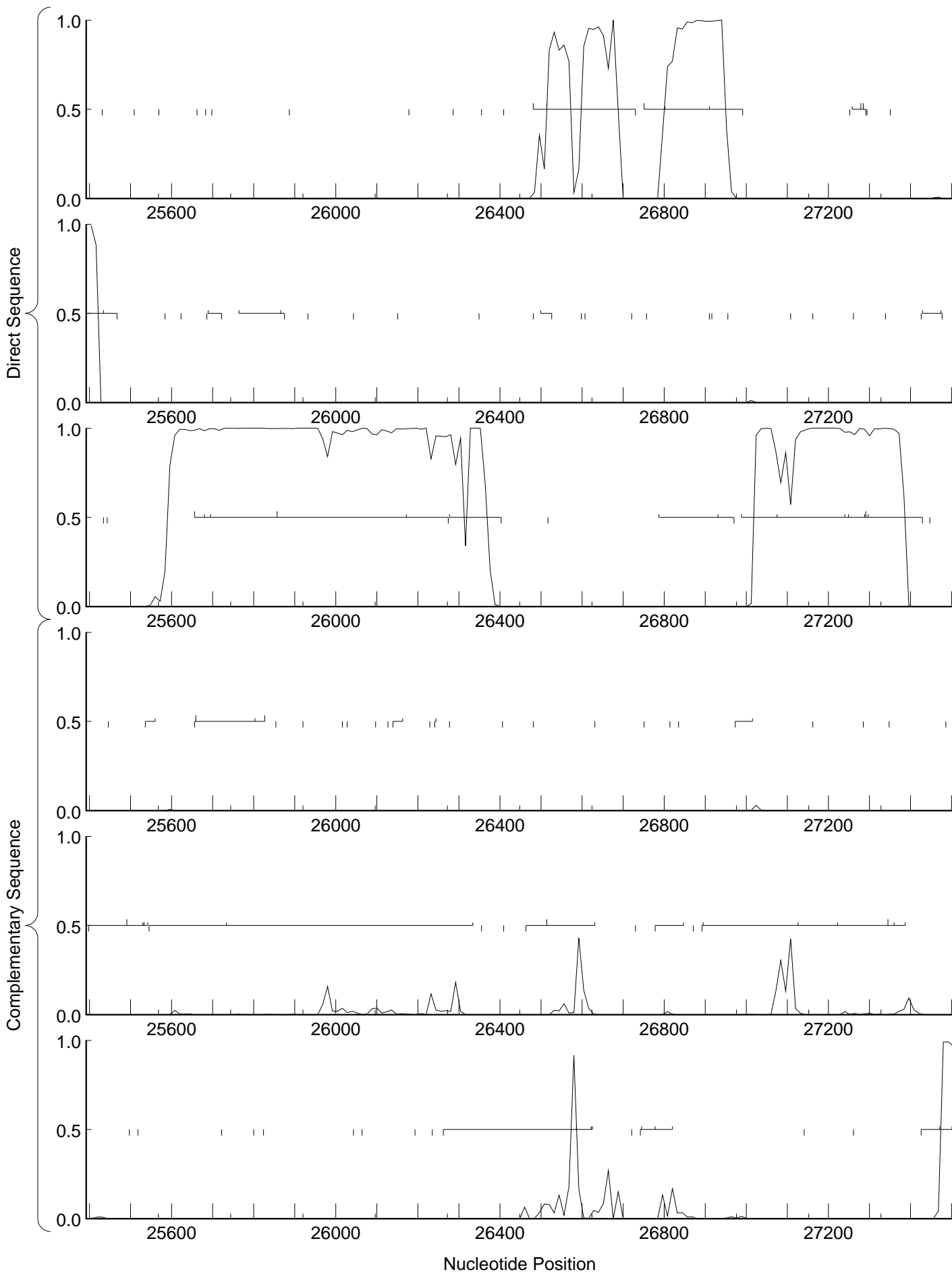












complete sequence, 45802 bp including 11-base 3' overhang (CGGAGAGGCAT), Cluster AZ, Order 4, Window 96, Step 12, 15/23

