

GeneMark

Version 2.5p (09.08.06)

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PROGRAM INFORMATION

Sequence : Gordonia phage Mayweather complete sequence, 48382 bp including 13-base 3' overhang (CGGCGG
Analysis Date : 10/2/22 at 18:25:57
Pages : 24
Sequence Length : 48382 bp
GC Content : 60.63%

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,

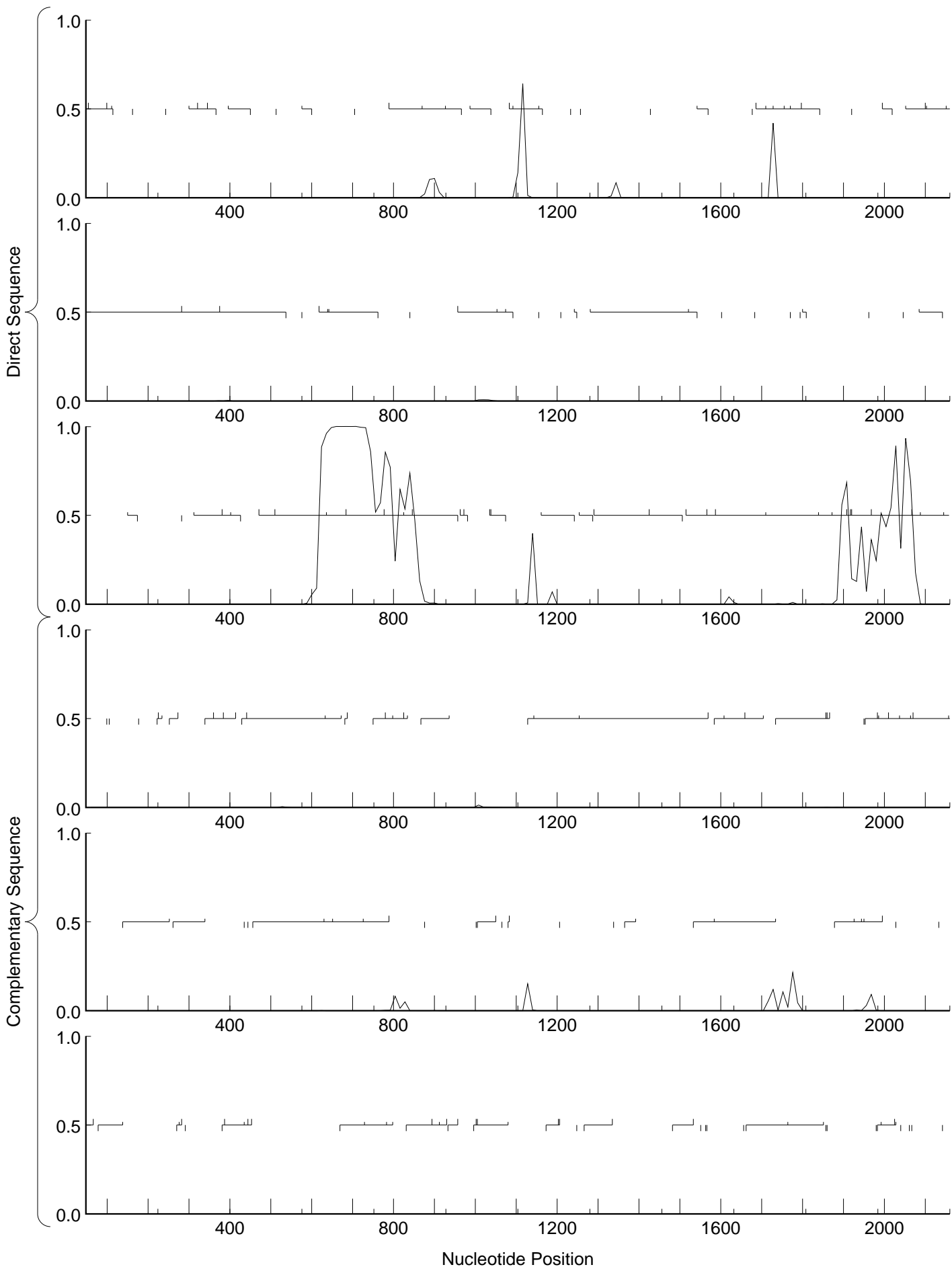
Matrix notes & comments

Training set derived by GeneMarkS, 4.30 October 2014
Sat Sep 8 00:36:14 2018

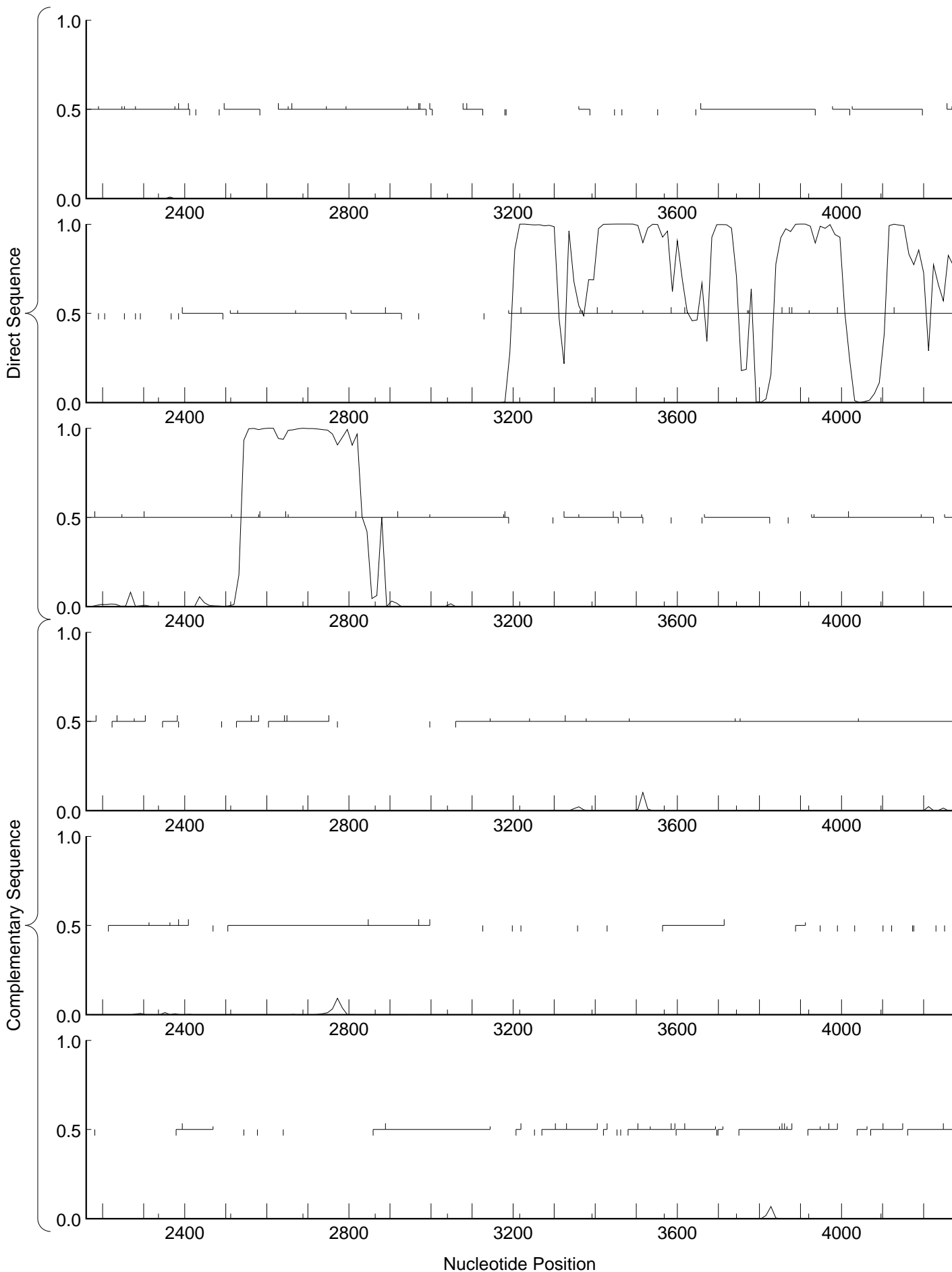
MATRIX INFORMATION

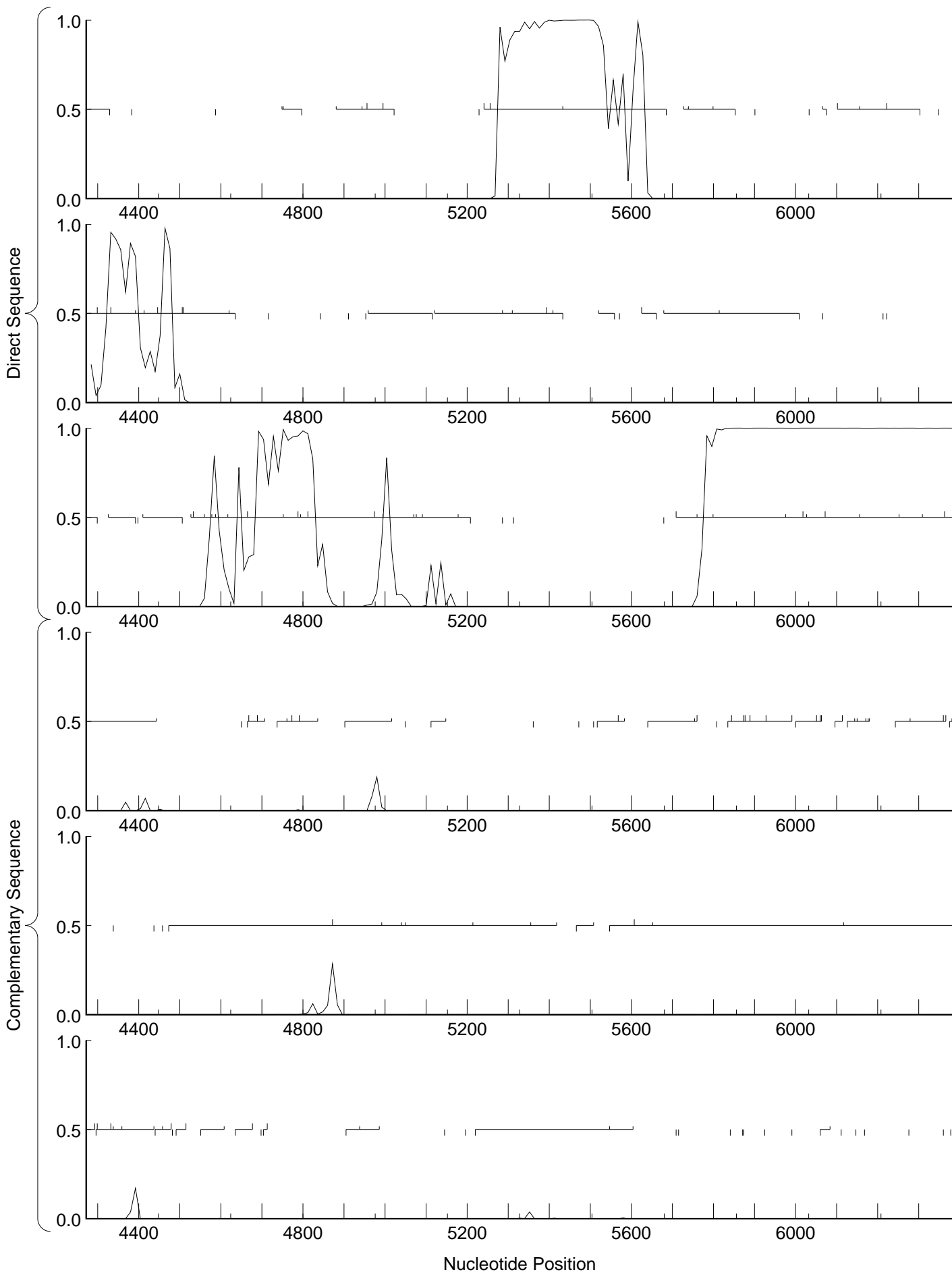
Matrix : -
Author : -
Order : 4

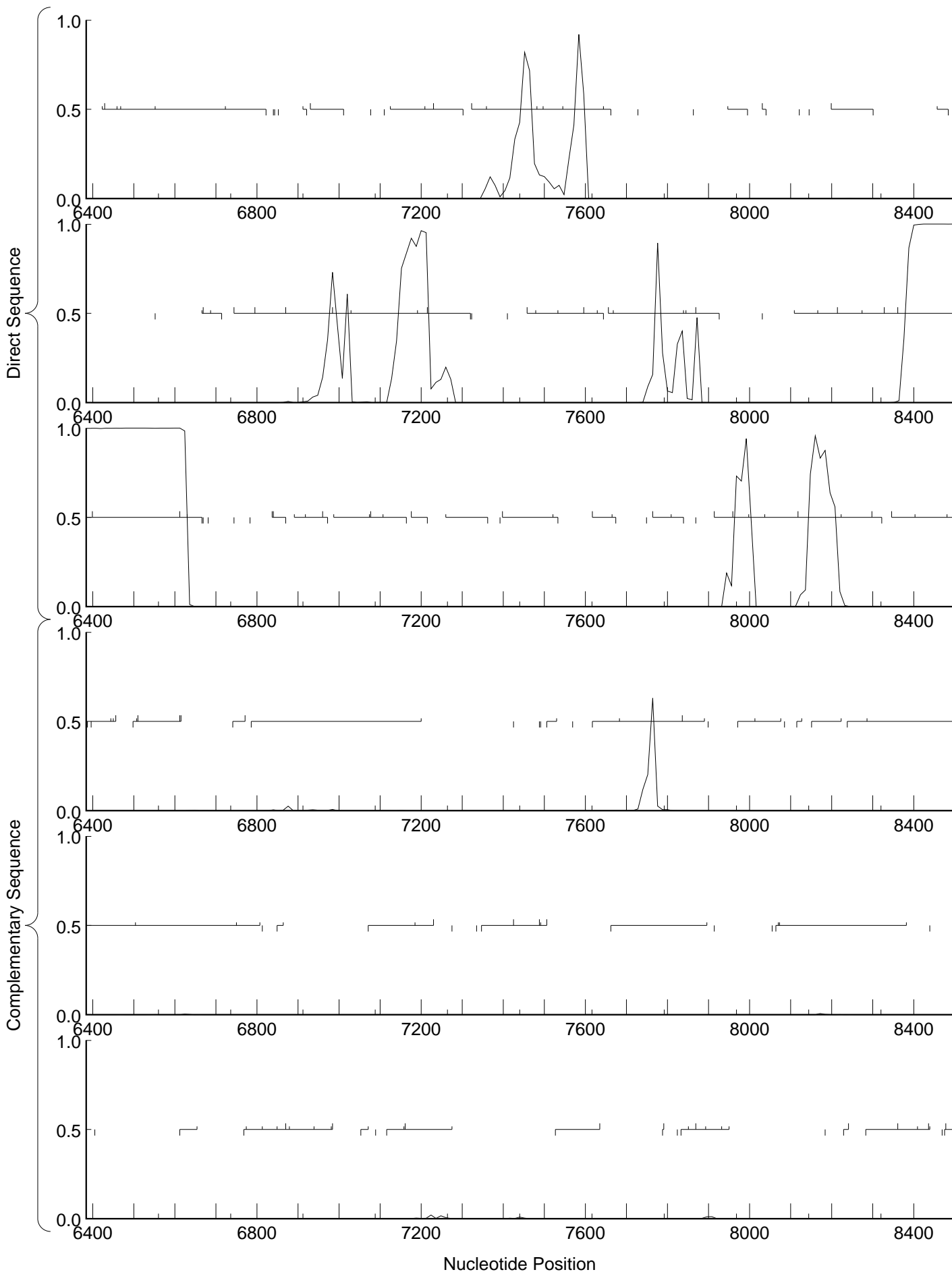
Send questions / comments to:
Dr. M. Borodovsky
Georgia Institute of Technology
School of Biology
Atlanta, GA 30332-0230

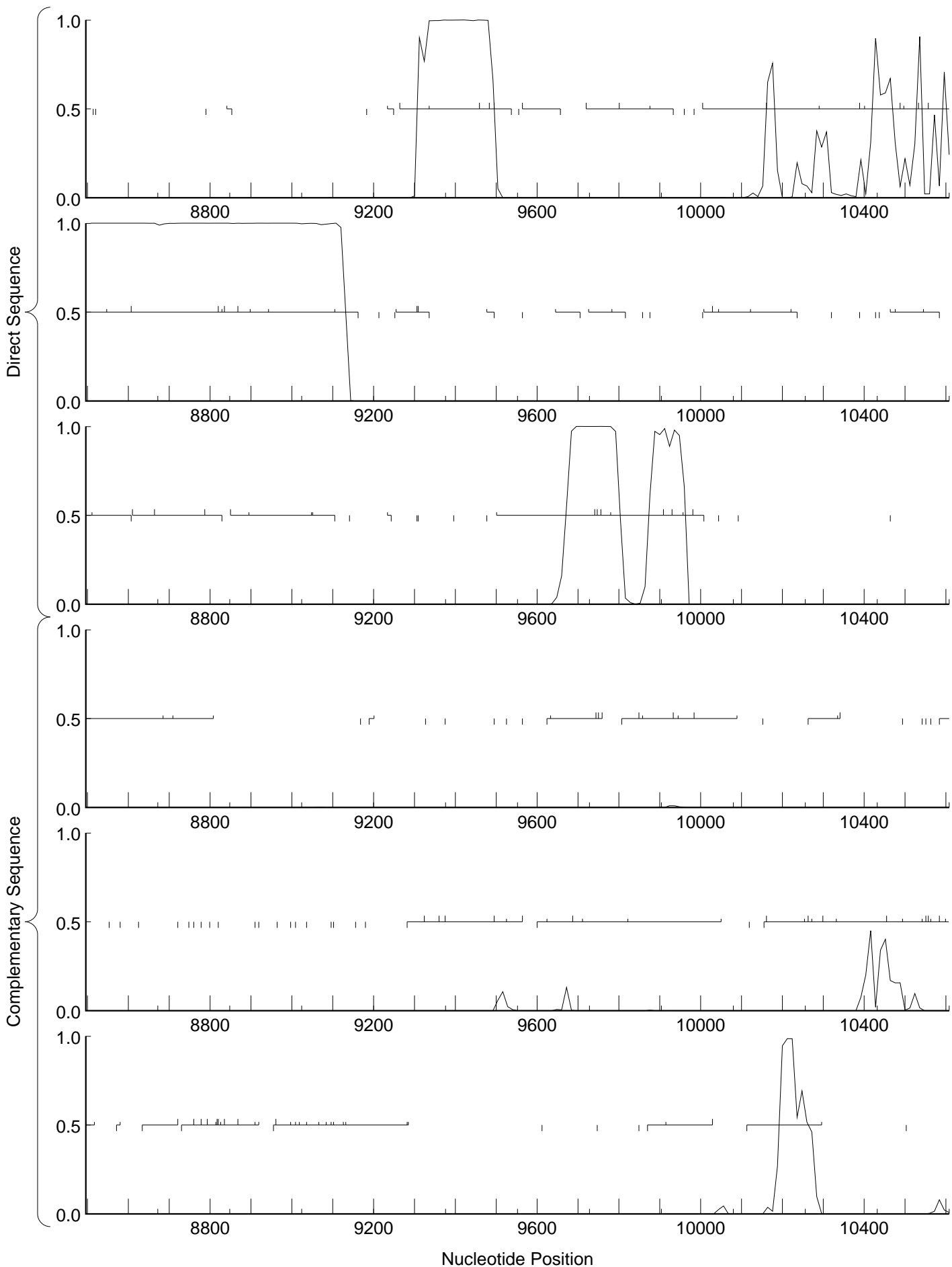


Complete sequence, 48382 bp including 13-base 3' overhang (CGGCGGTAGGCTT), Cluster CT, Order 4, Window 96, Step 12, 3/24

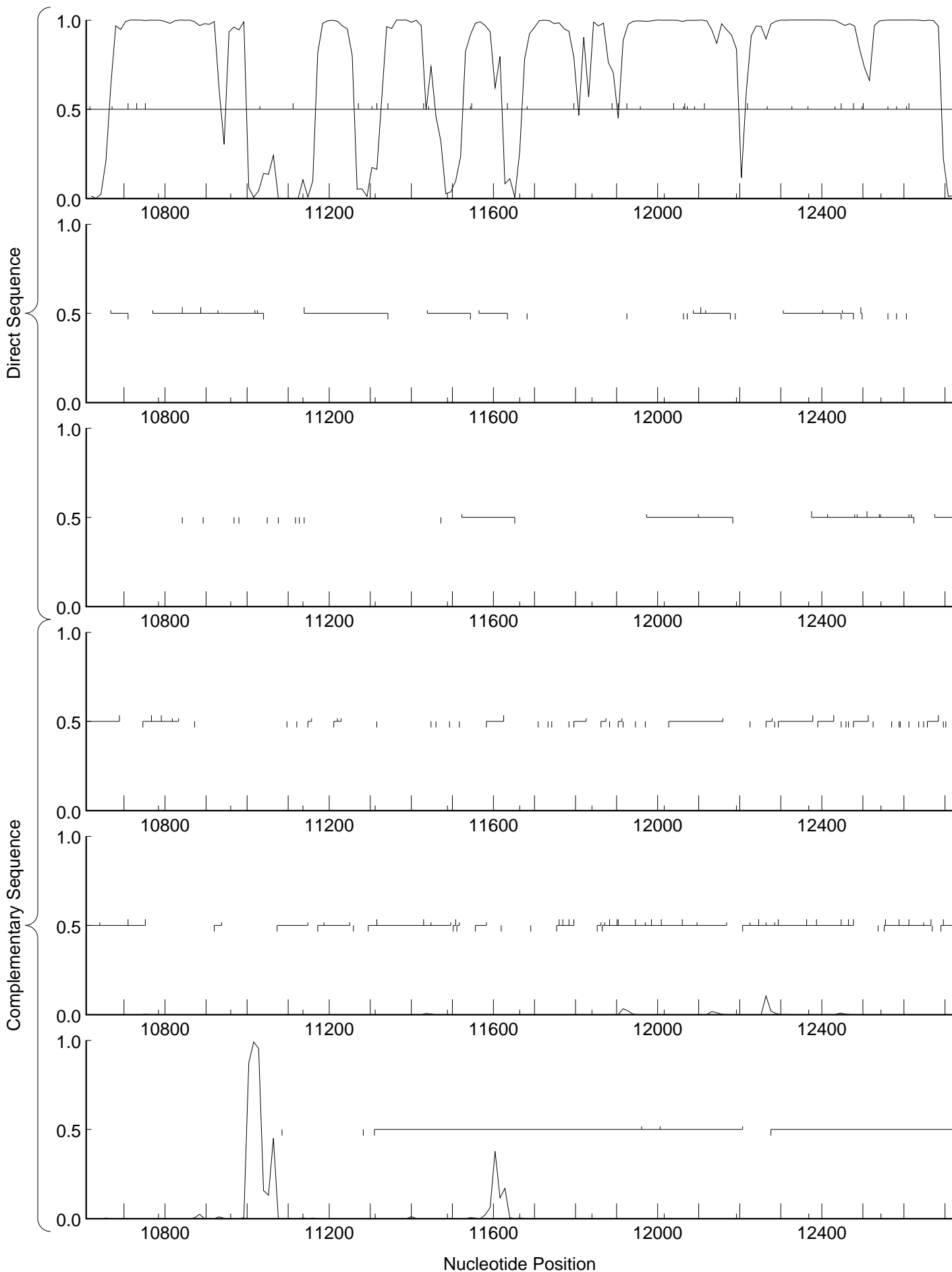




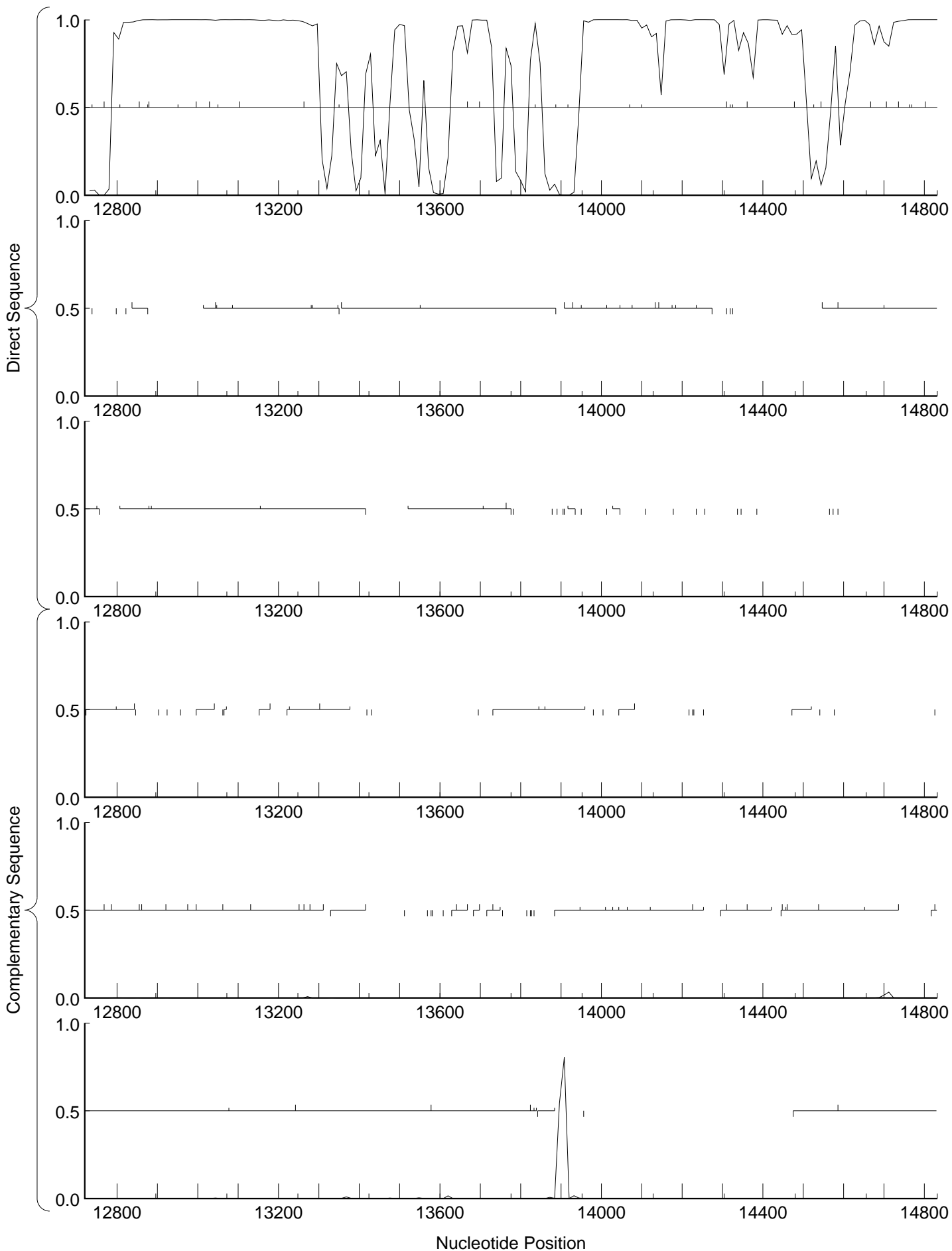


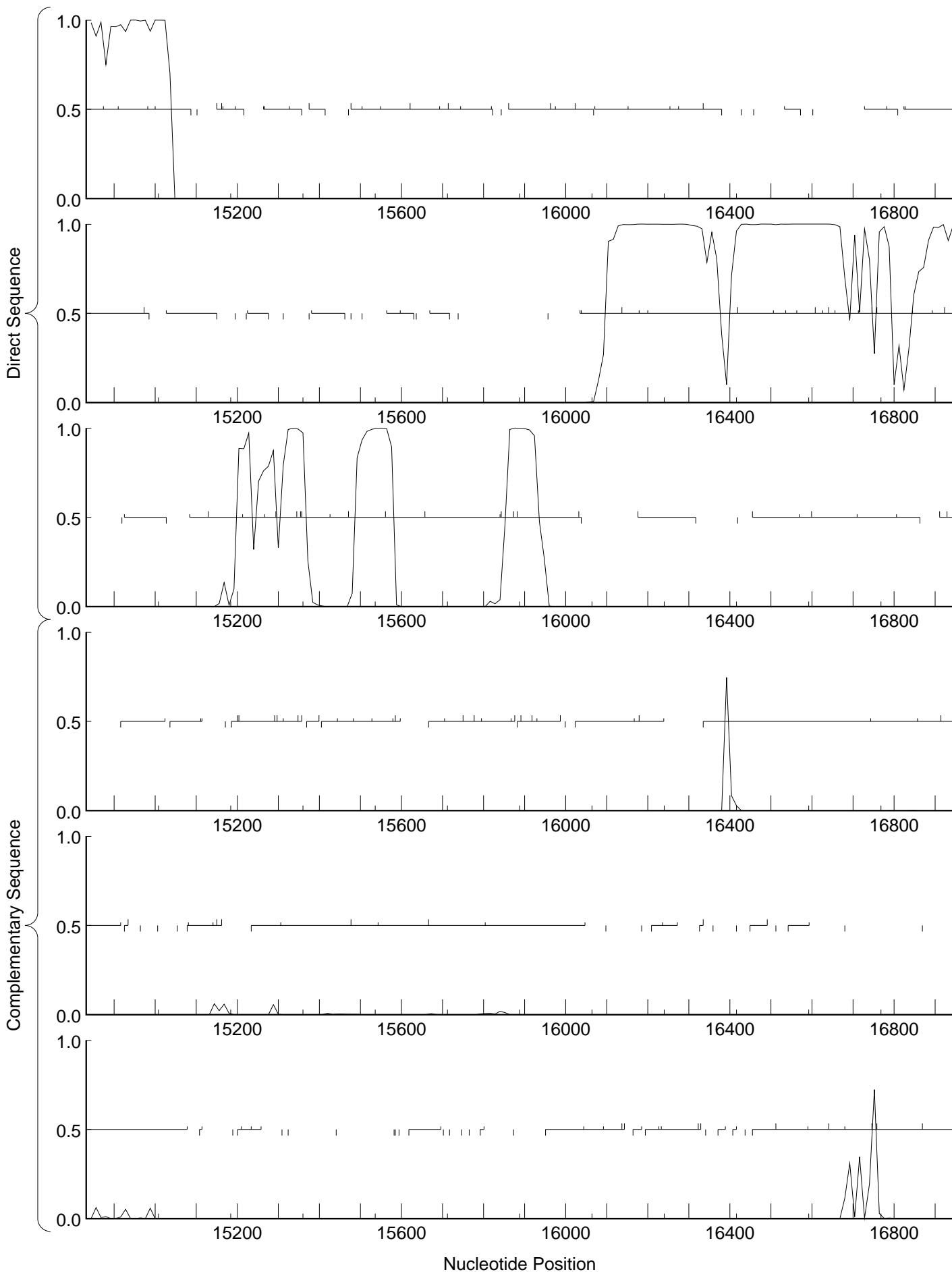


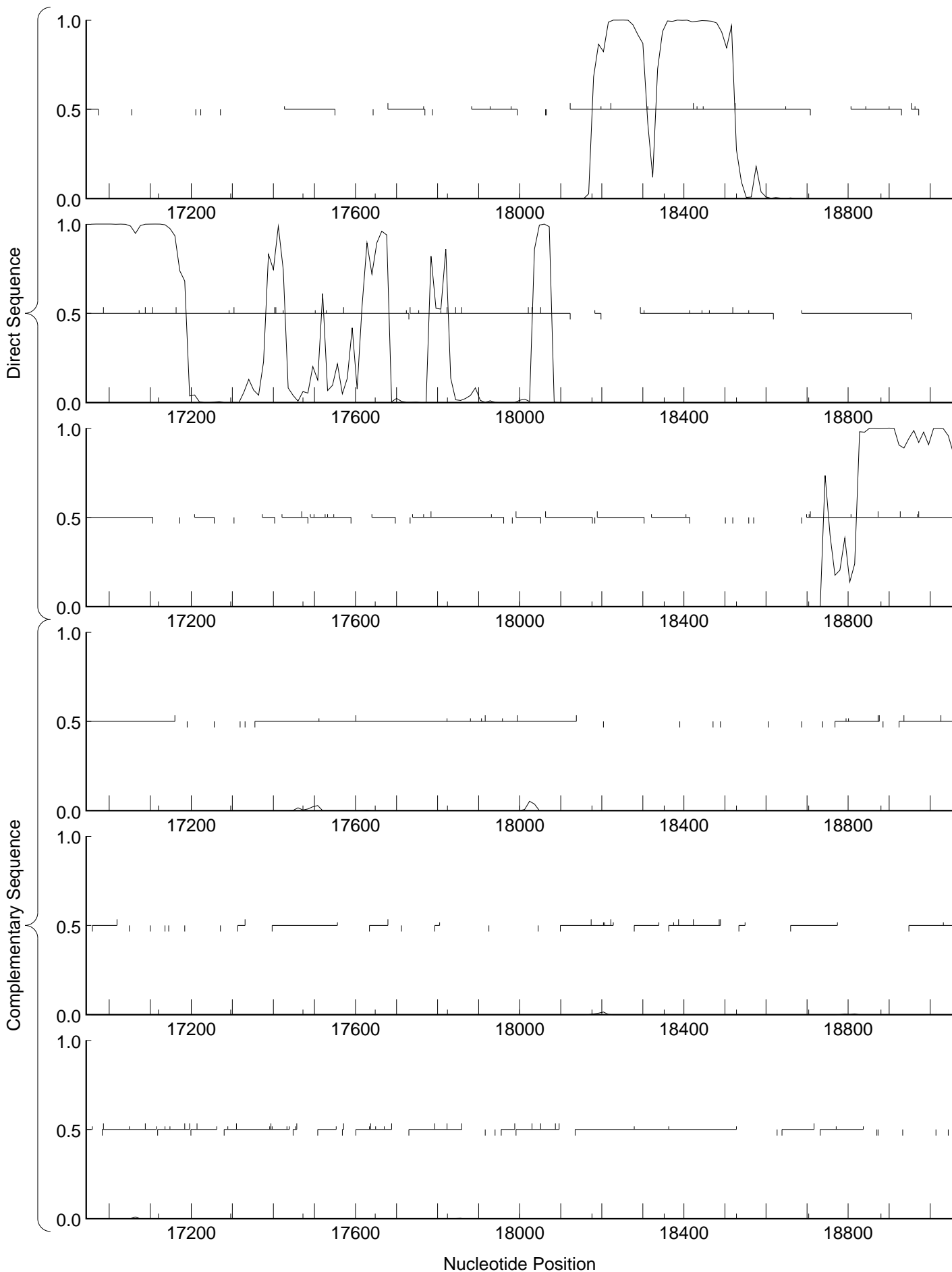
Complete sequence, 48382 bp including 13-base 3' overhang (CGGCGGTAGGCTT), Cluster CT, Order 4, Window 96, Step 12, 7/24

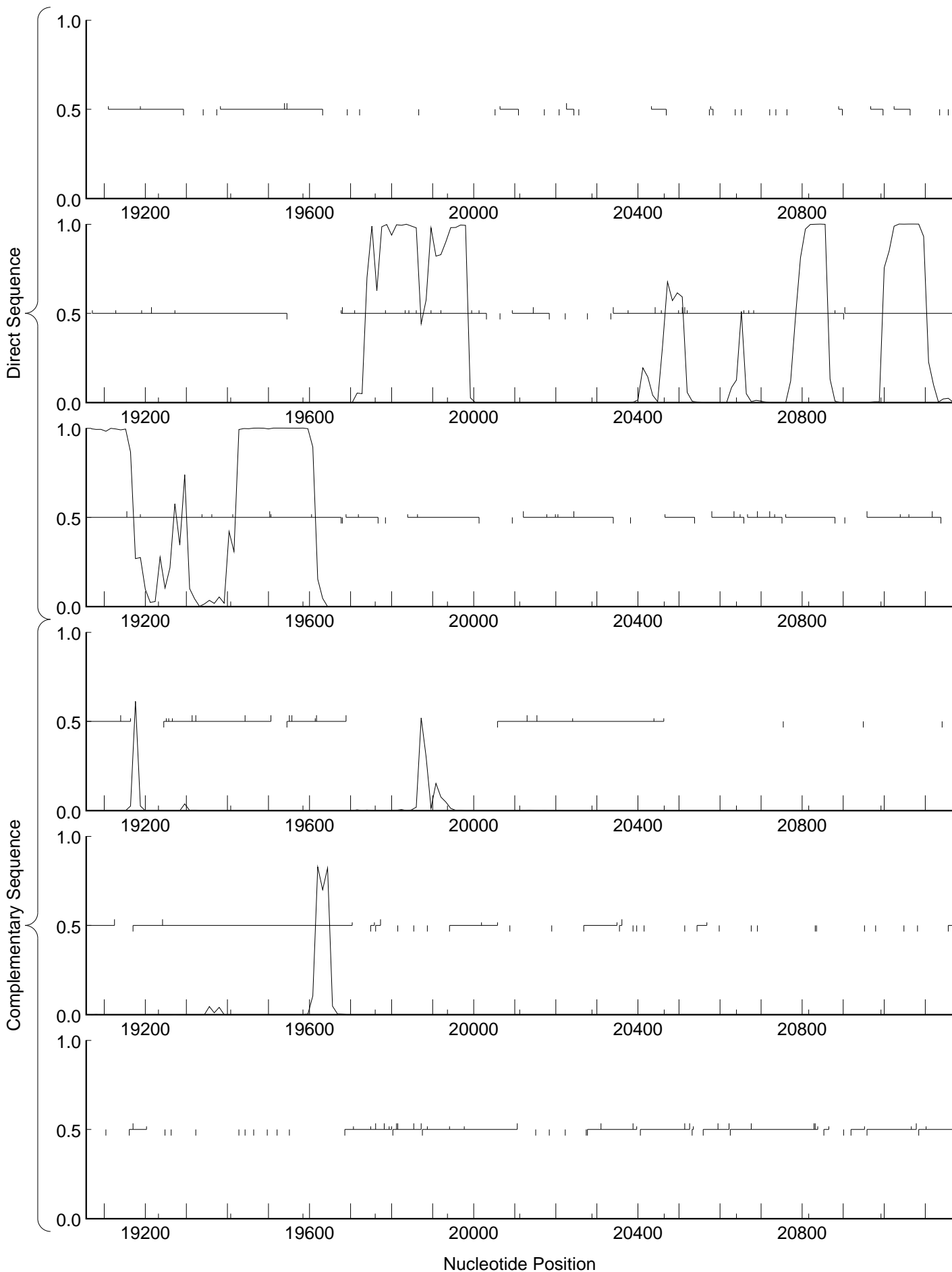


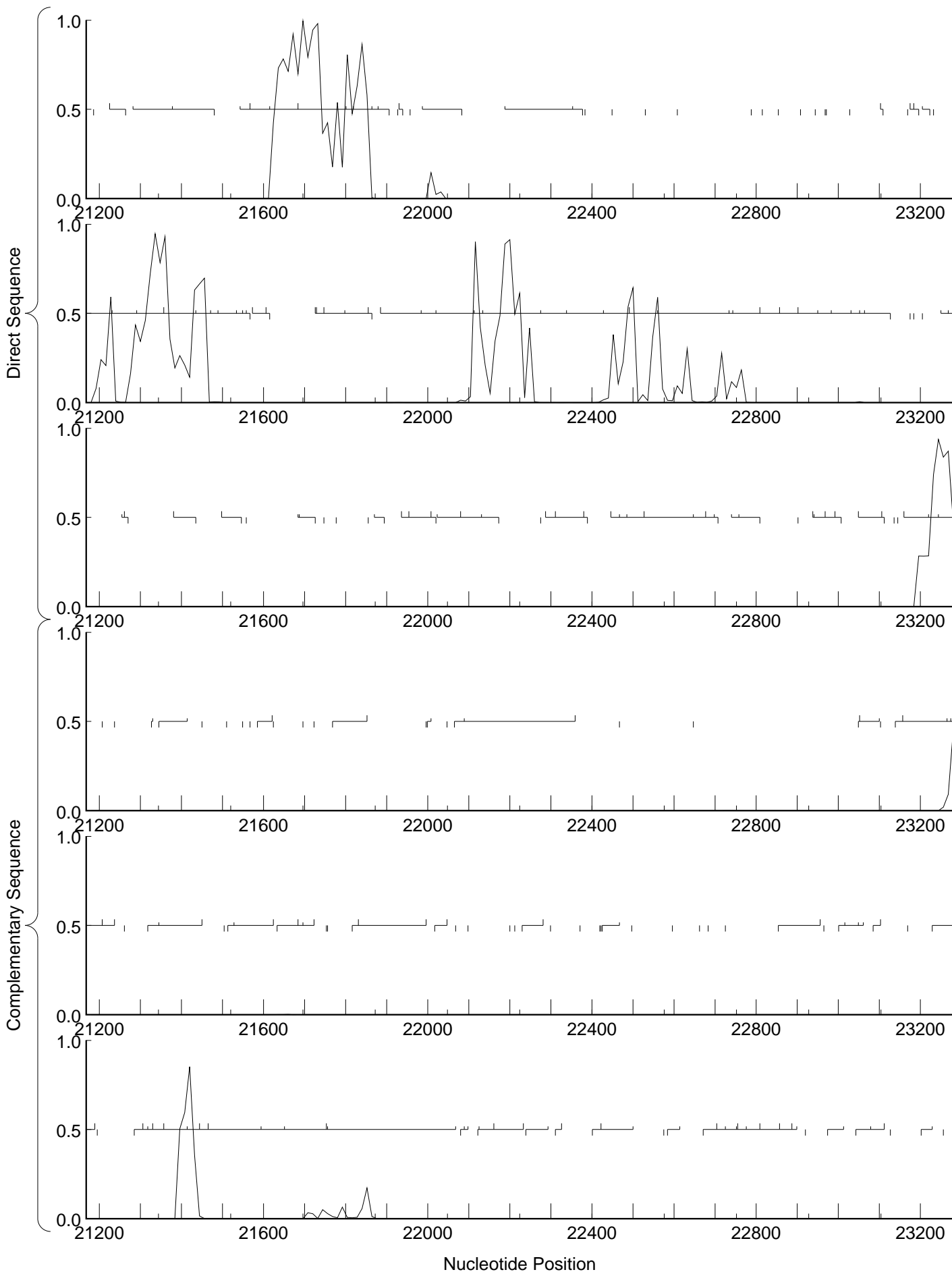
Complete sequence, 48382 bp including 13-base 3' overhang (CGGCGGTAGGCTT), Cluster CT, Order 4, Window 96, Step 12, 8/24



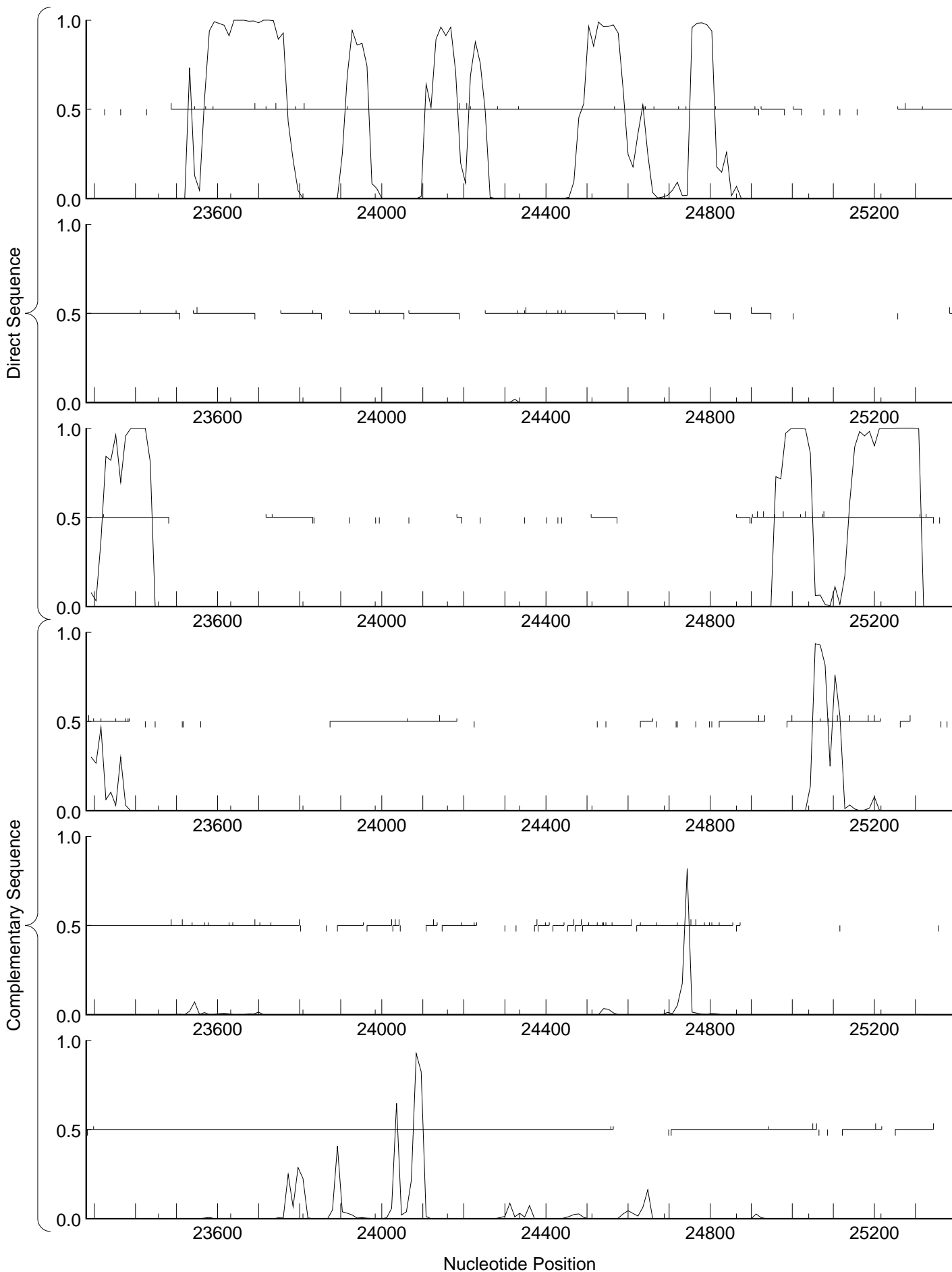


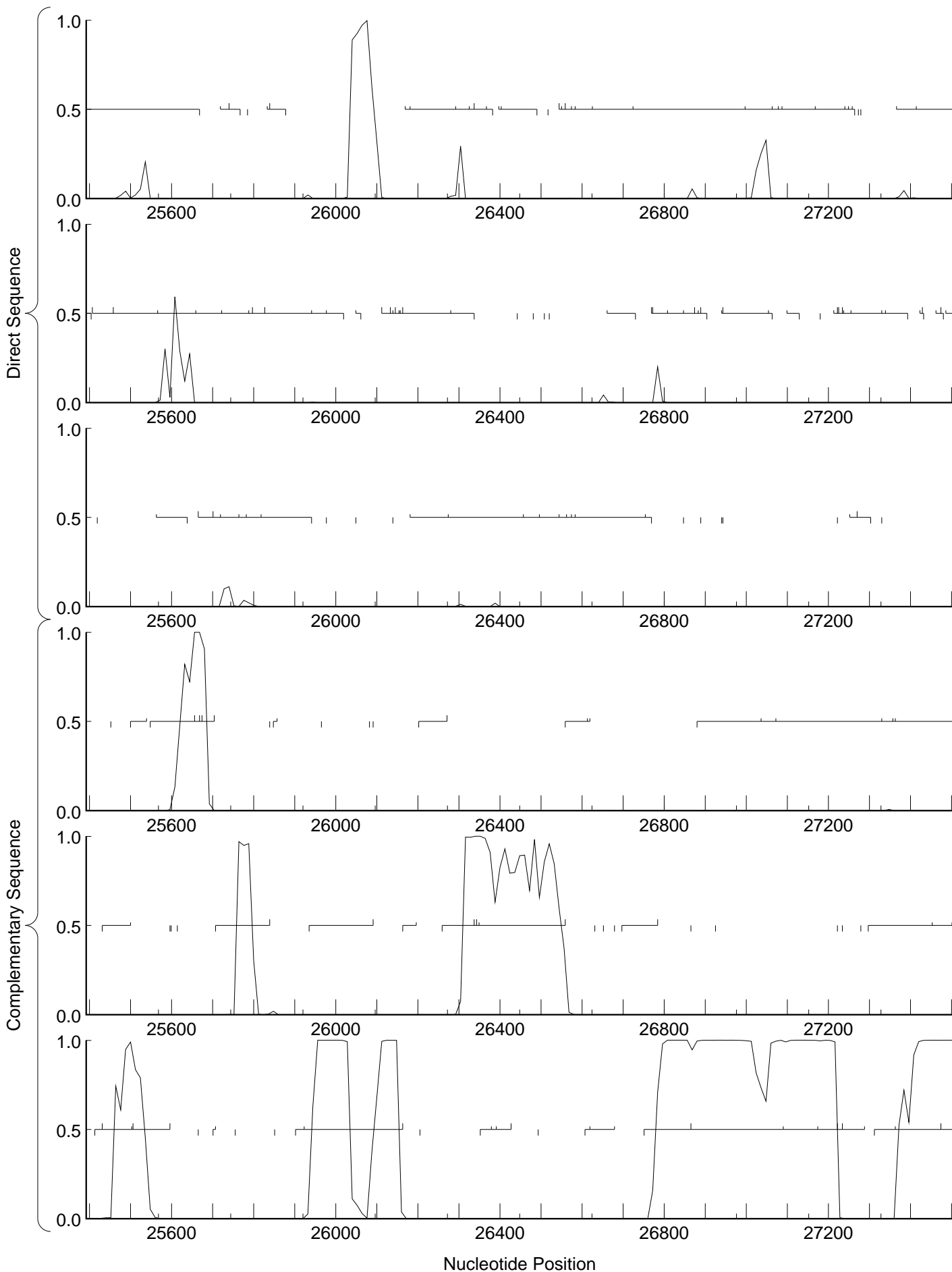




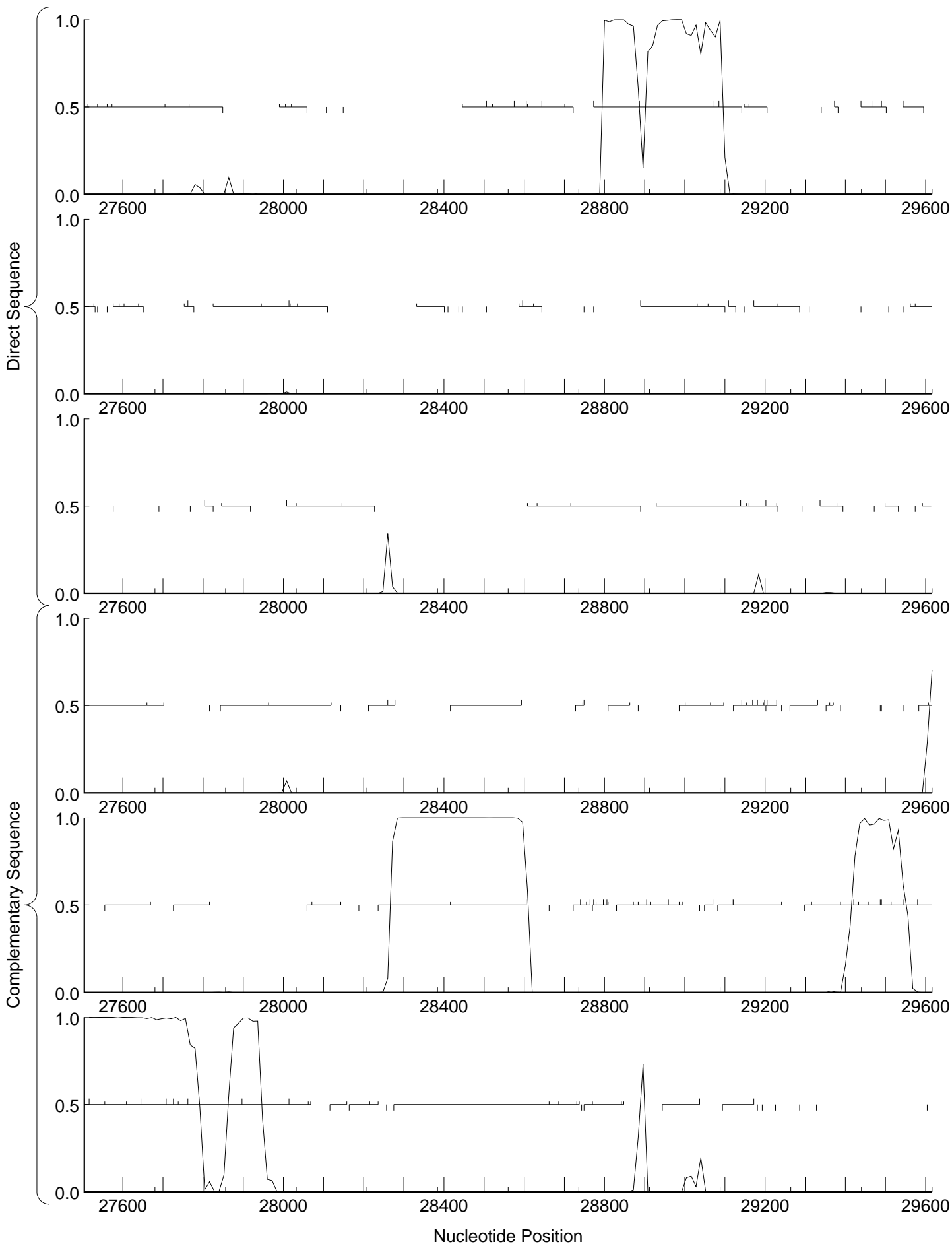


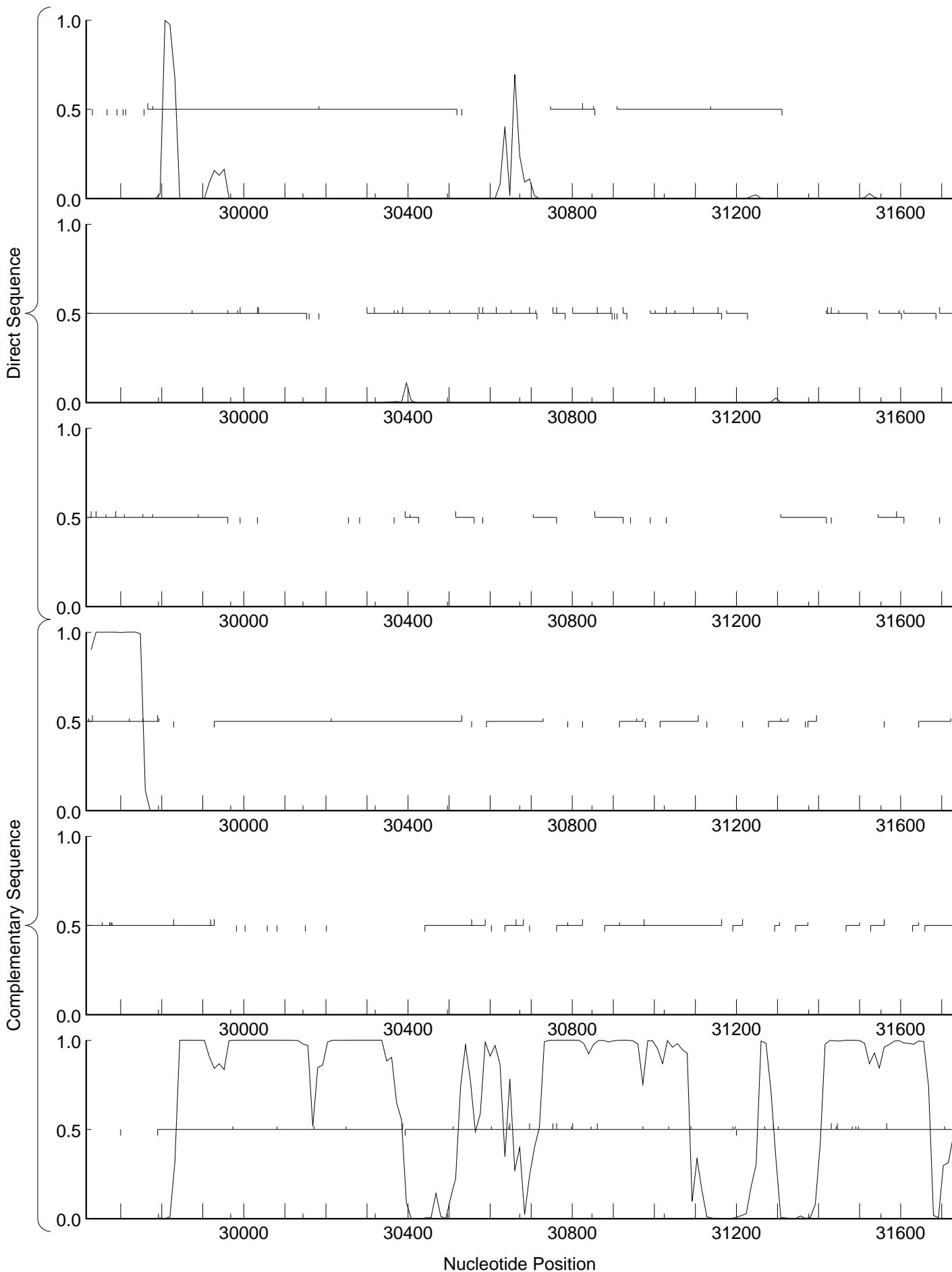
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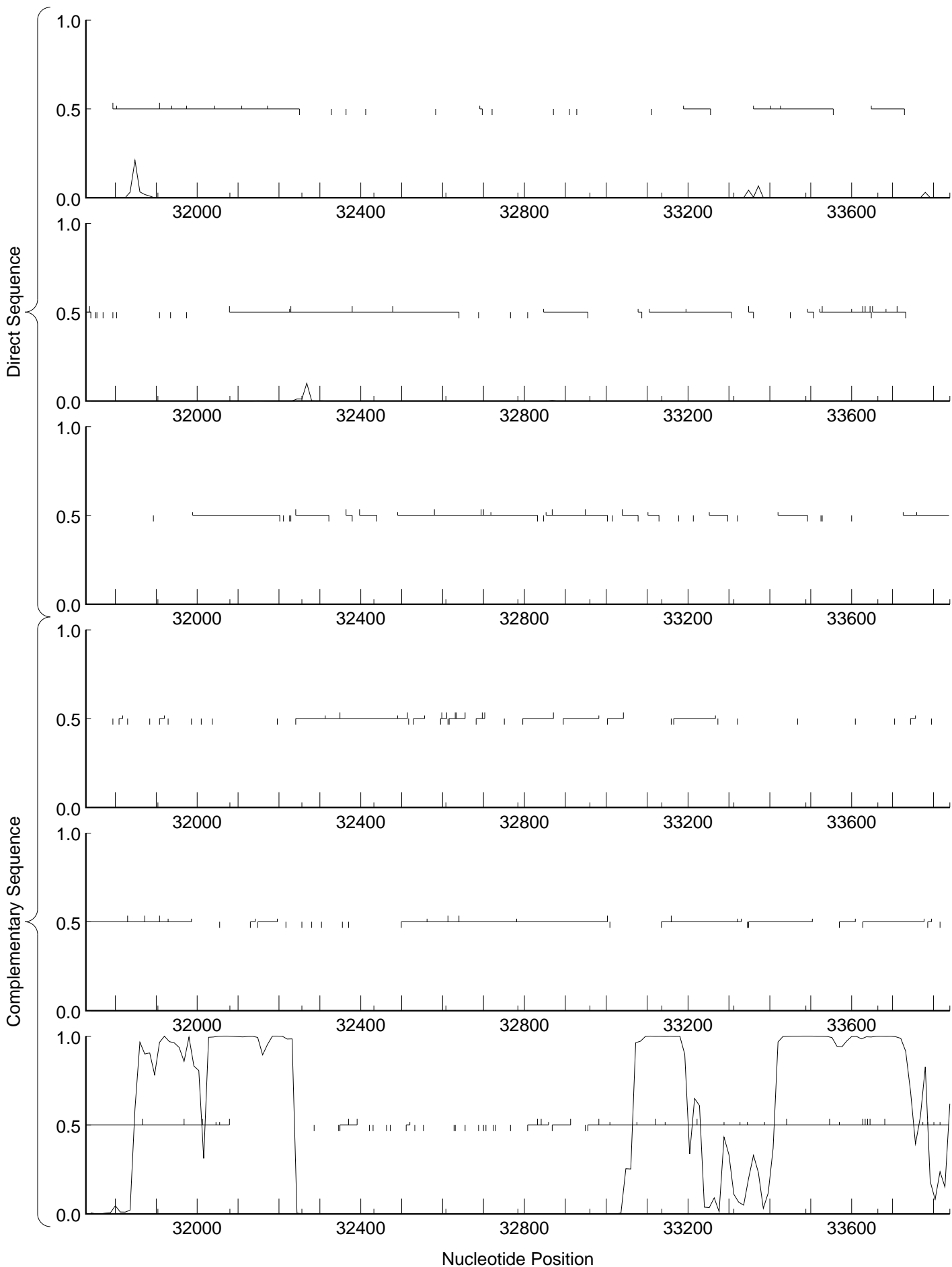


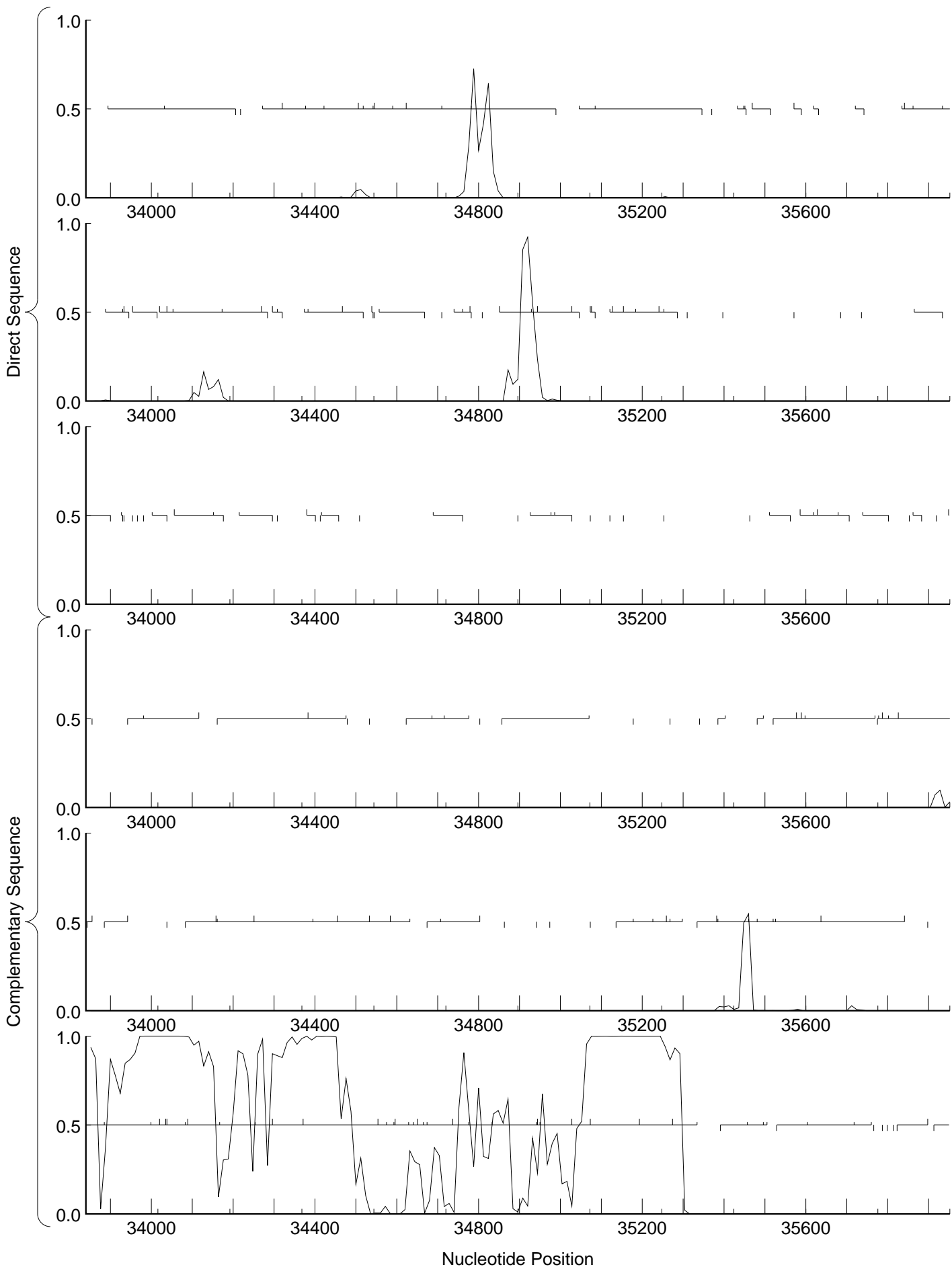
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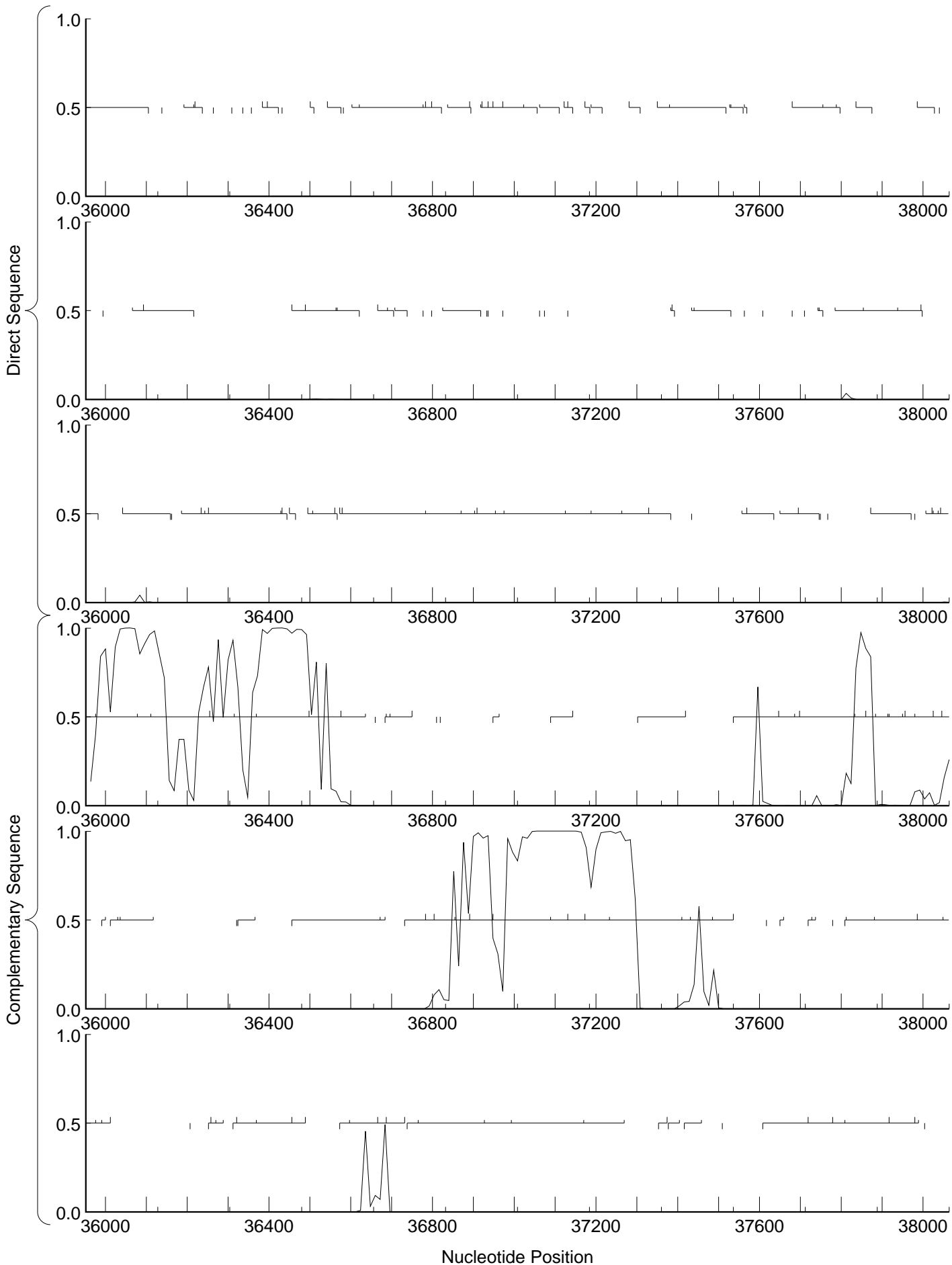


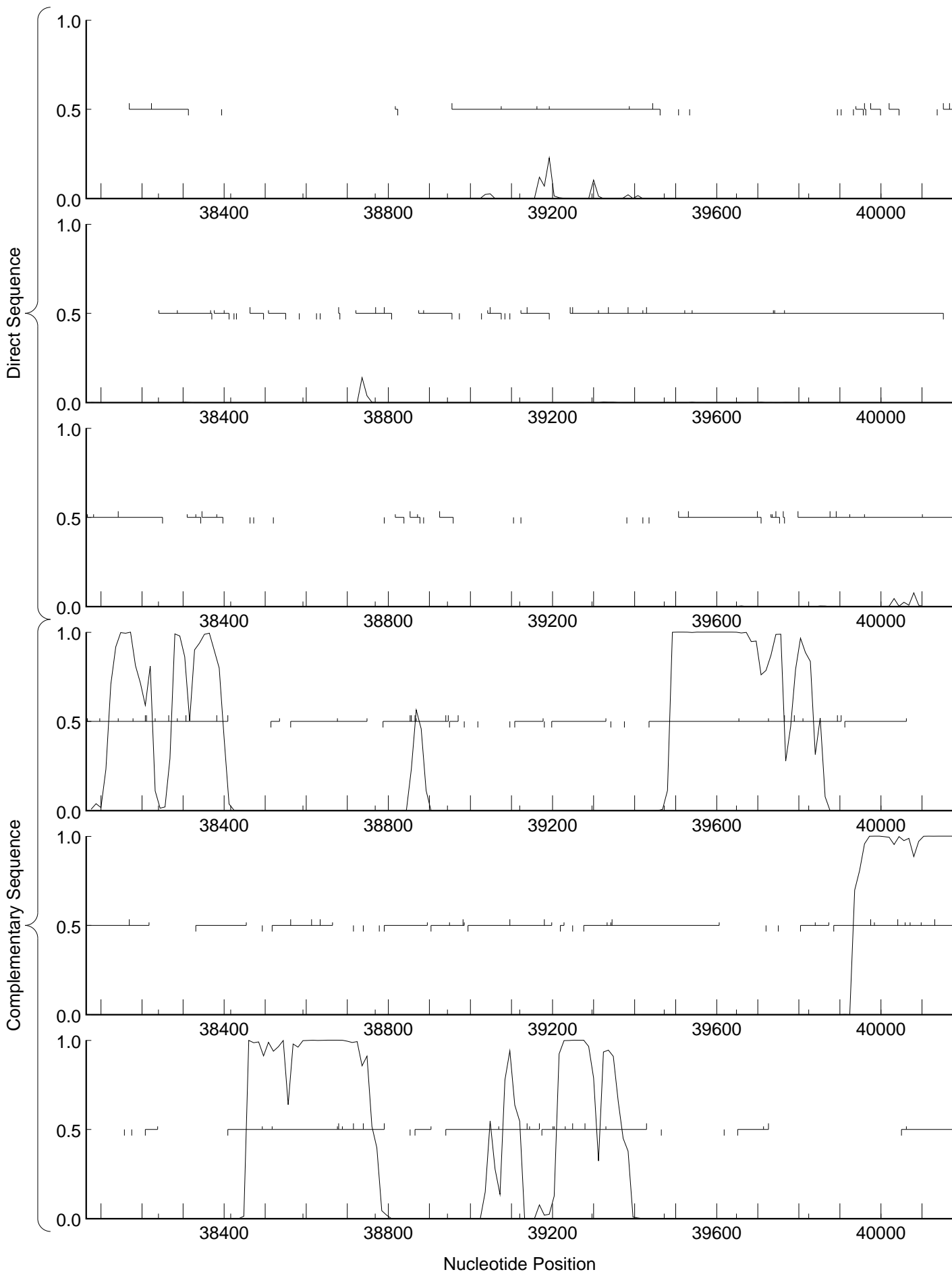


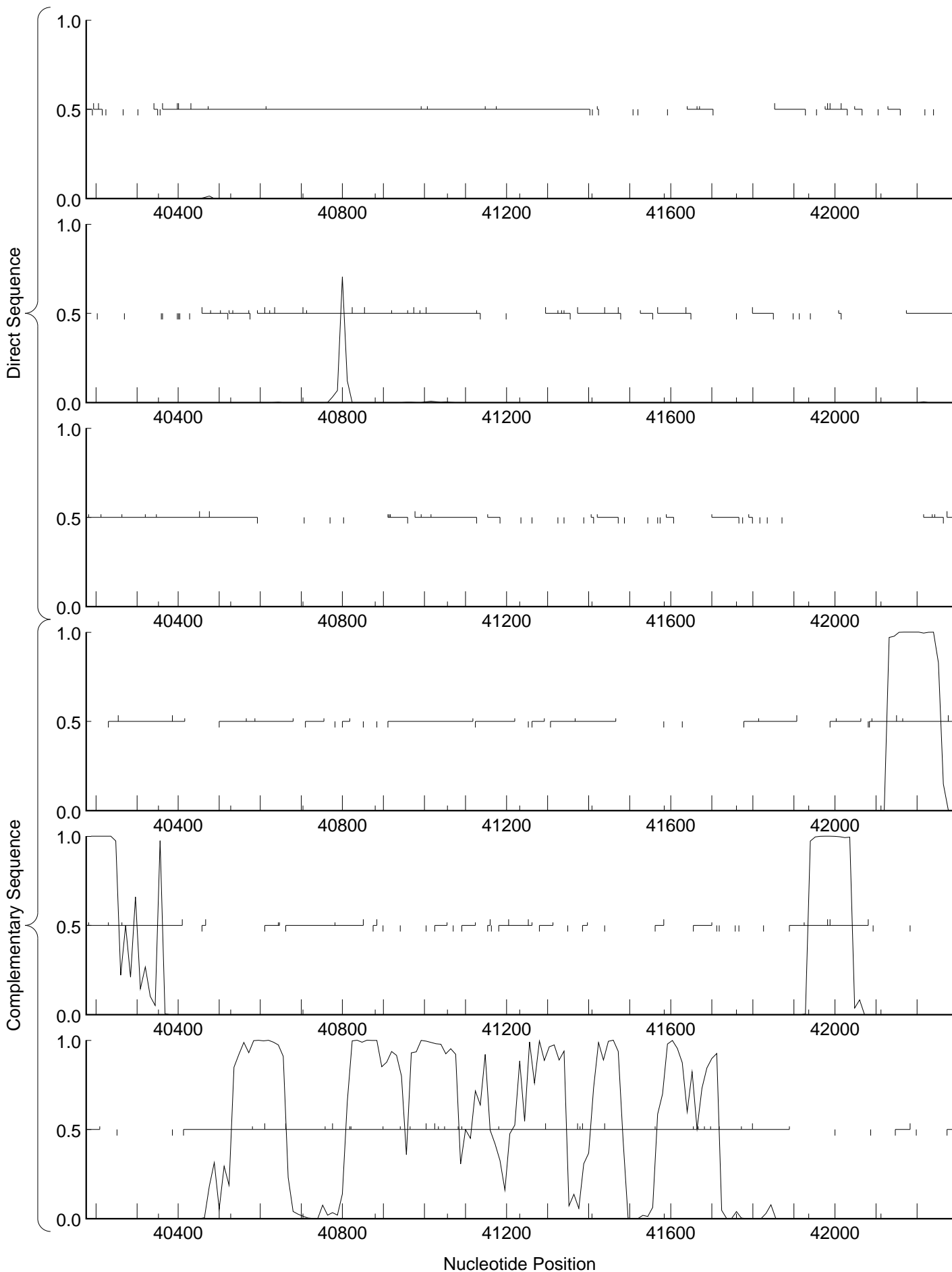
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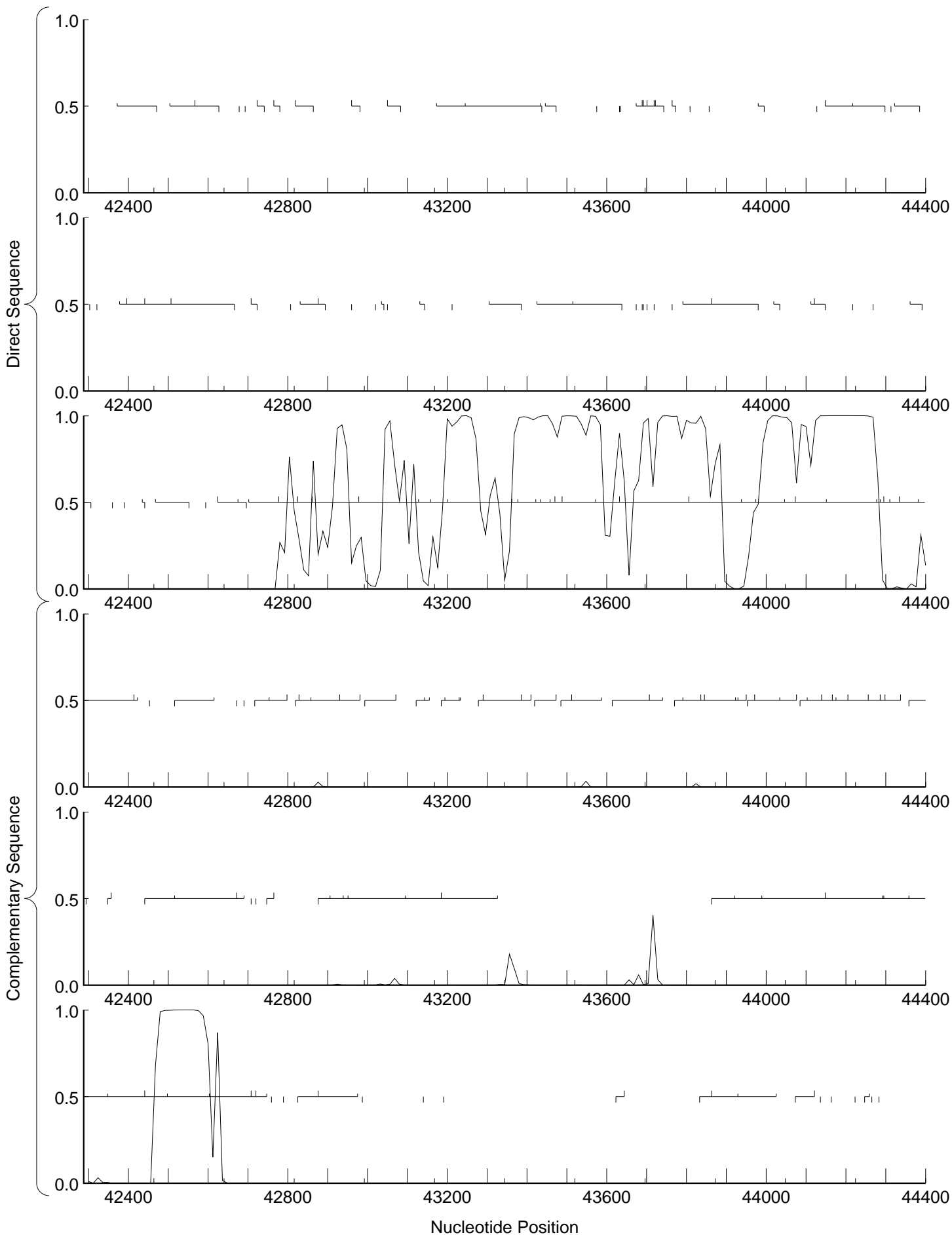








plete sequence, 48382 bp including 13-base 3' overhang (CGGCGGTAGGCTT), Cluster CT, Order 4, Window 96, Step 12, 22/24



plete sequence, 48382 bp including 13-base 3' overhang (CGGCGGTAGGCTT), Cluster CT, Order 4, Window 96, Step 12, 23/24

