

GeneMark

Version 2.5p (09.08.06)

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PROGRAM INFORMATION

Sequence : Gordonia phage Margaret complete sequence, 46950 bp including 10-base 3' overhang (CGGTAGGC
Analysis Date : 9/20/19 at 3:22:13
Pages : 24
Sequence Length : 46950 bp
GC Content : 62.78%

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,

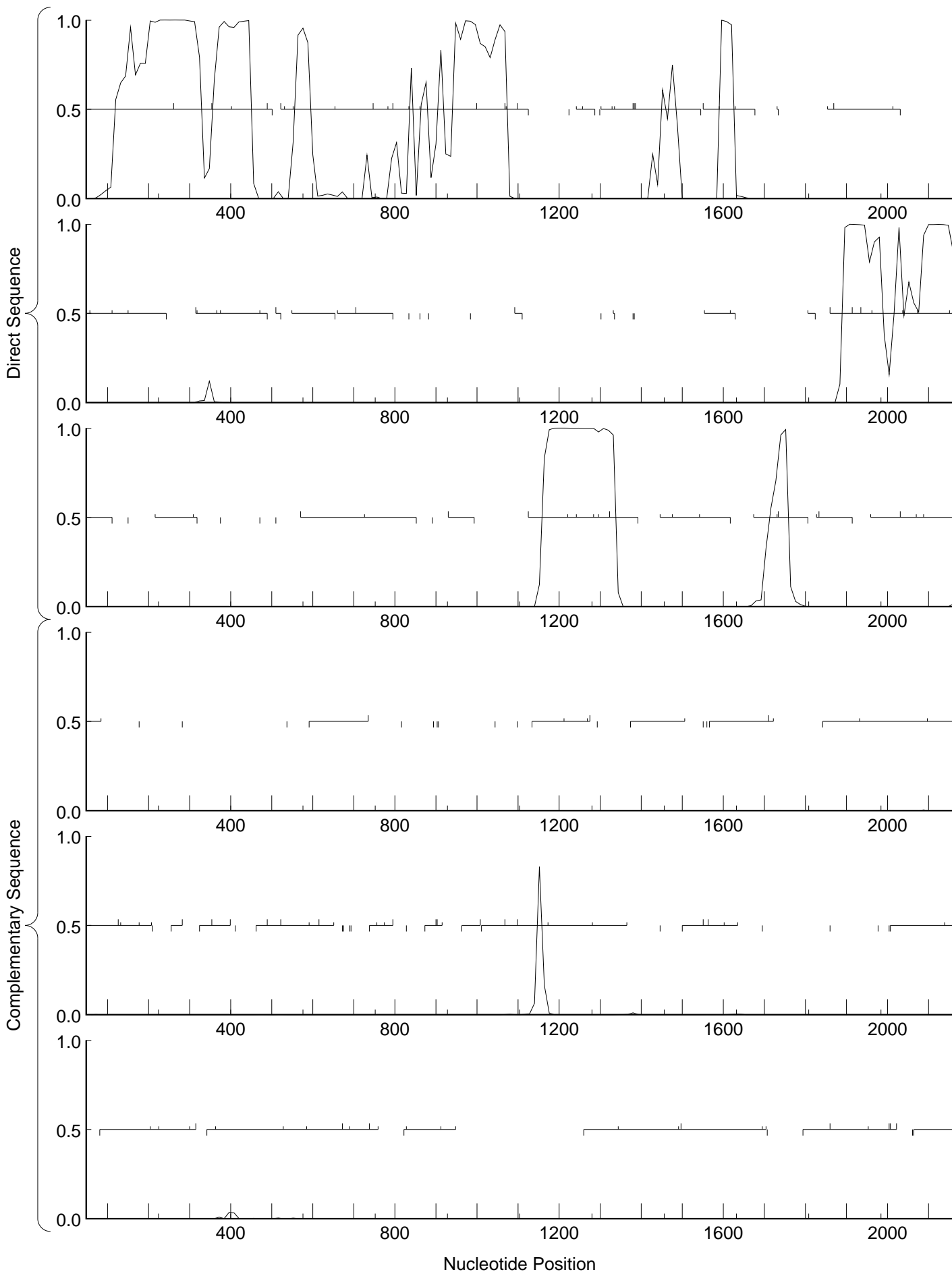
MATRIX INFORMATION

Matrix : -
Author : -
Order : 4

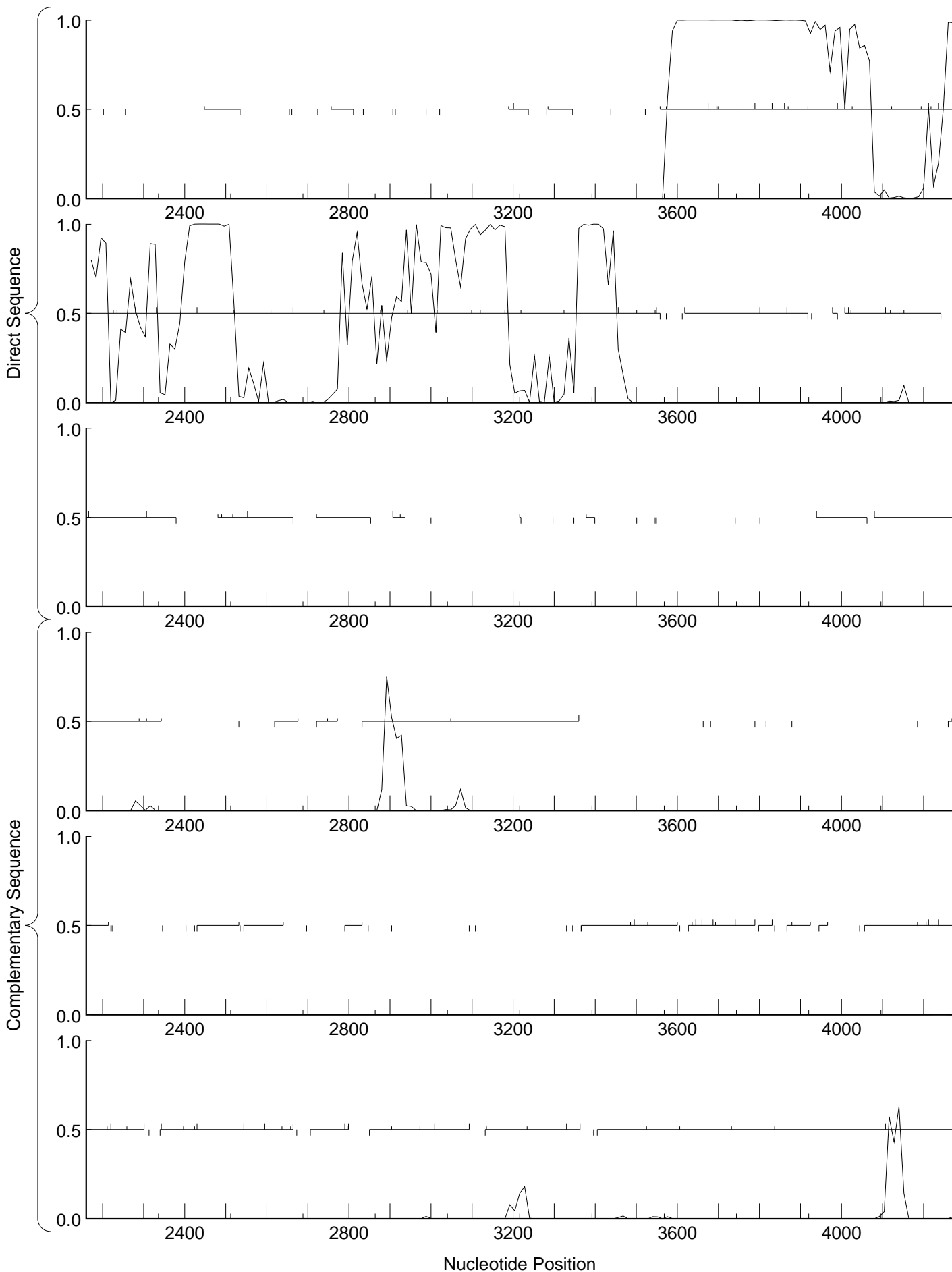
Send questions / comments to:
Dr. M. Borodovsky
Georgia Institute of Technology
School of Biology
Atlanta, GA 30332-0230

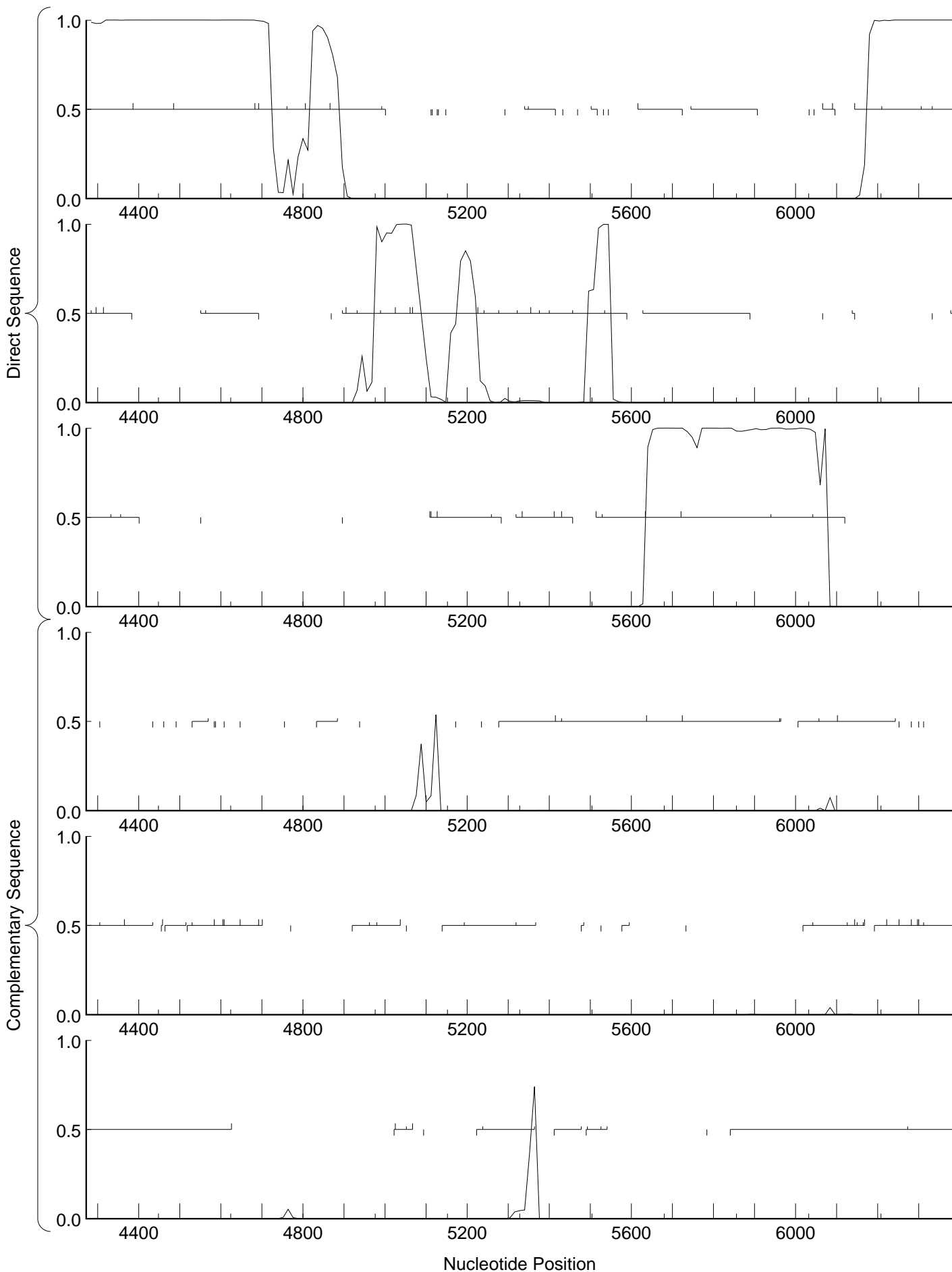
Matrix notes & comments

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

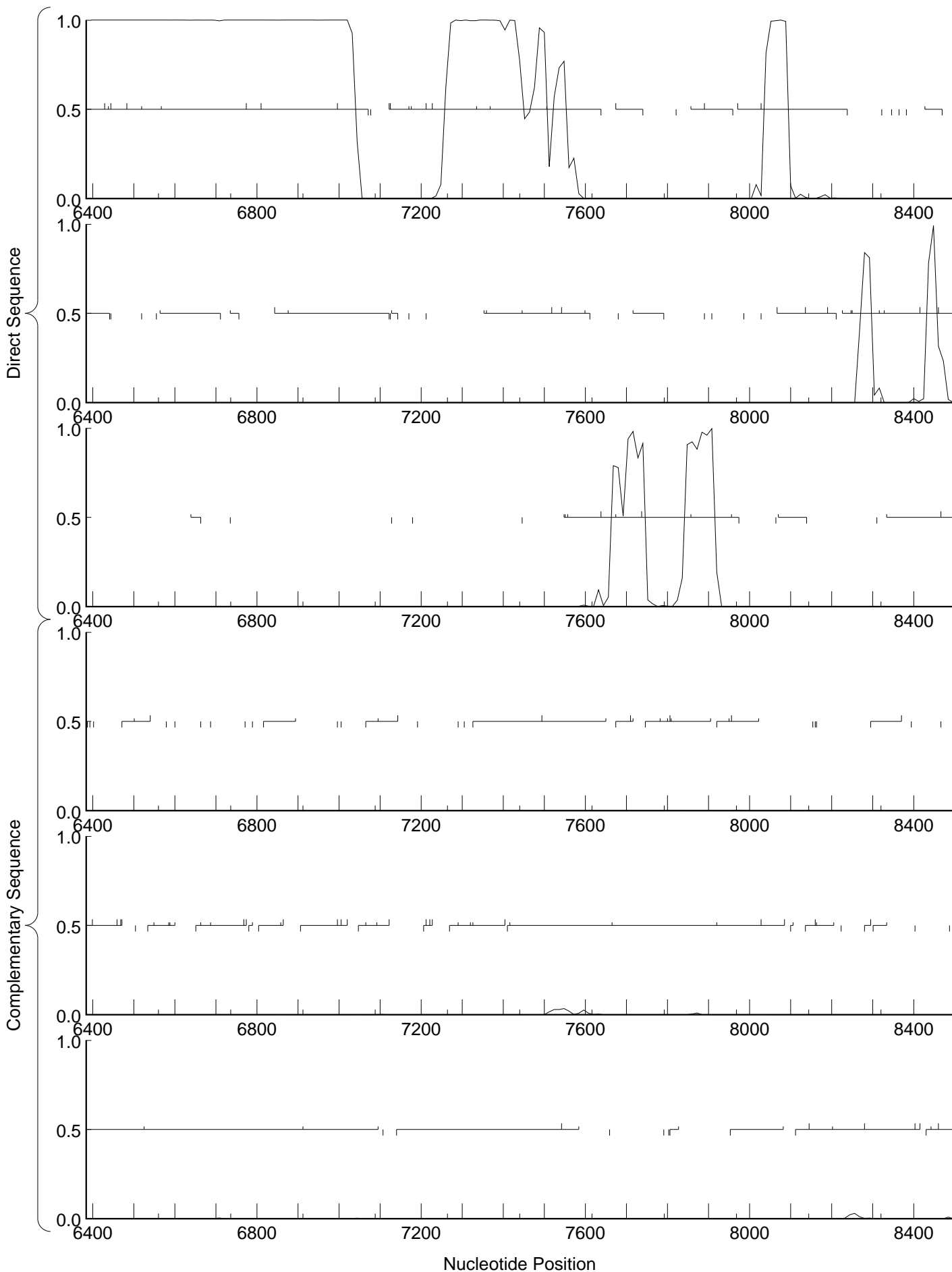


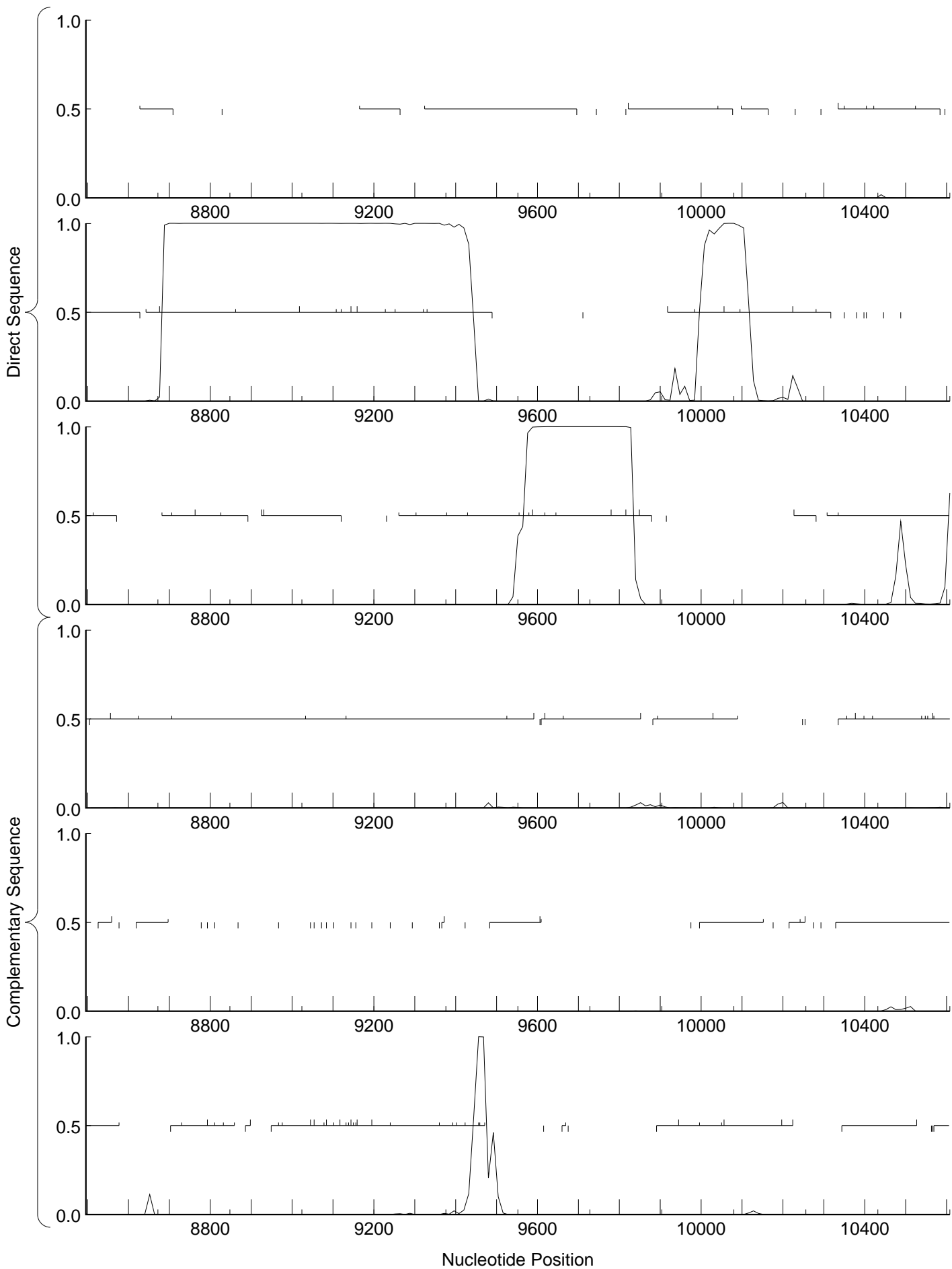
et complete sequence, 46950 bp including 10-base 3' overhang (CGGTAGGCAT), Cluster CT, Order 4, Window 96, Step 12, 3/24

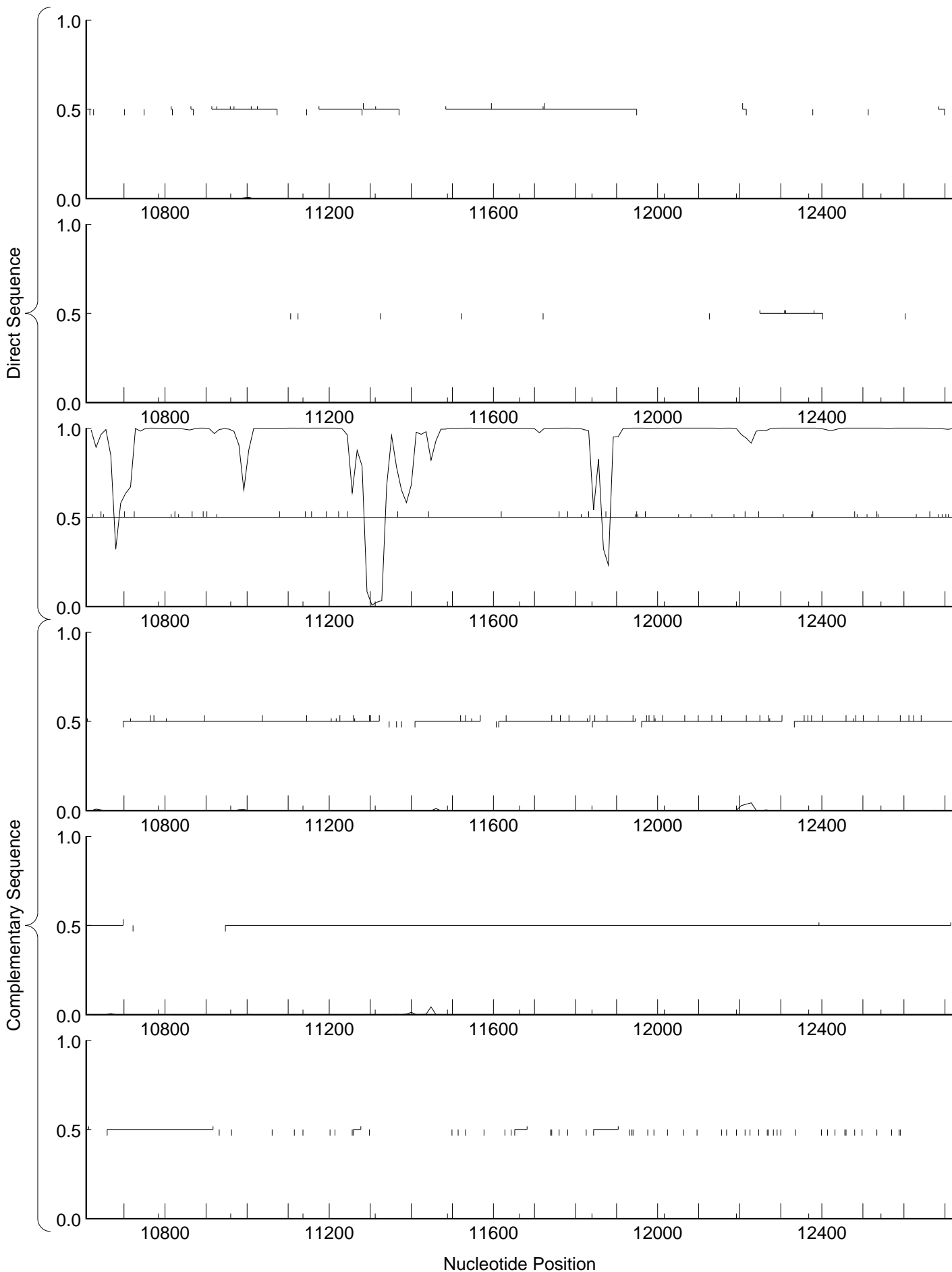


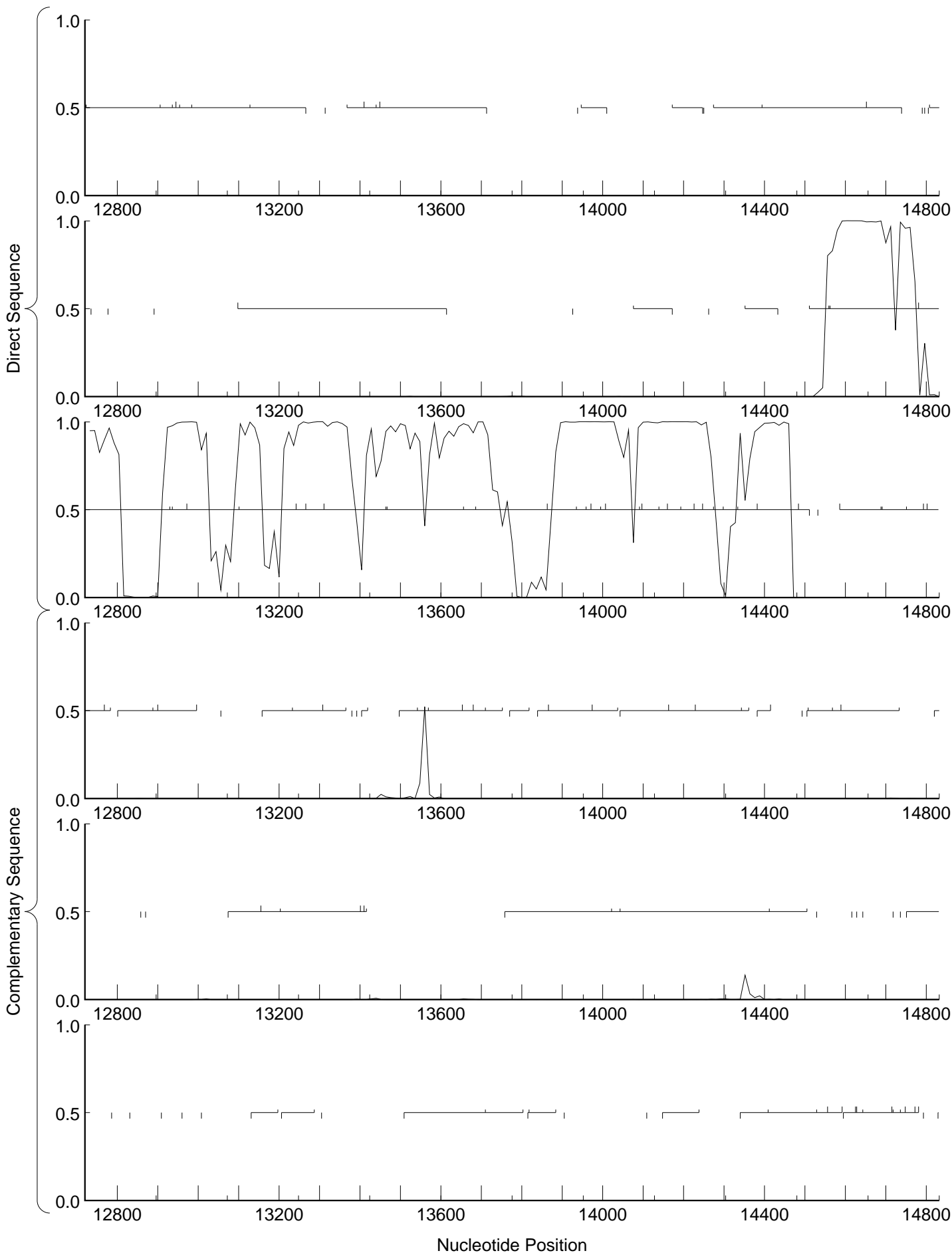


et complete sequence, 46950 bp including 10-base 3' overhang (CGGTAGGCAT), Cluster CT, Order 4, Window 96, Step 12, 5/24

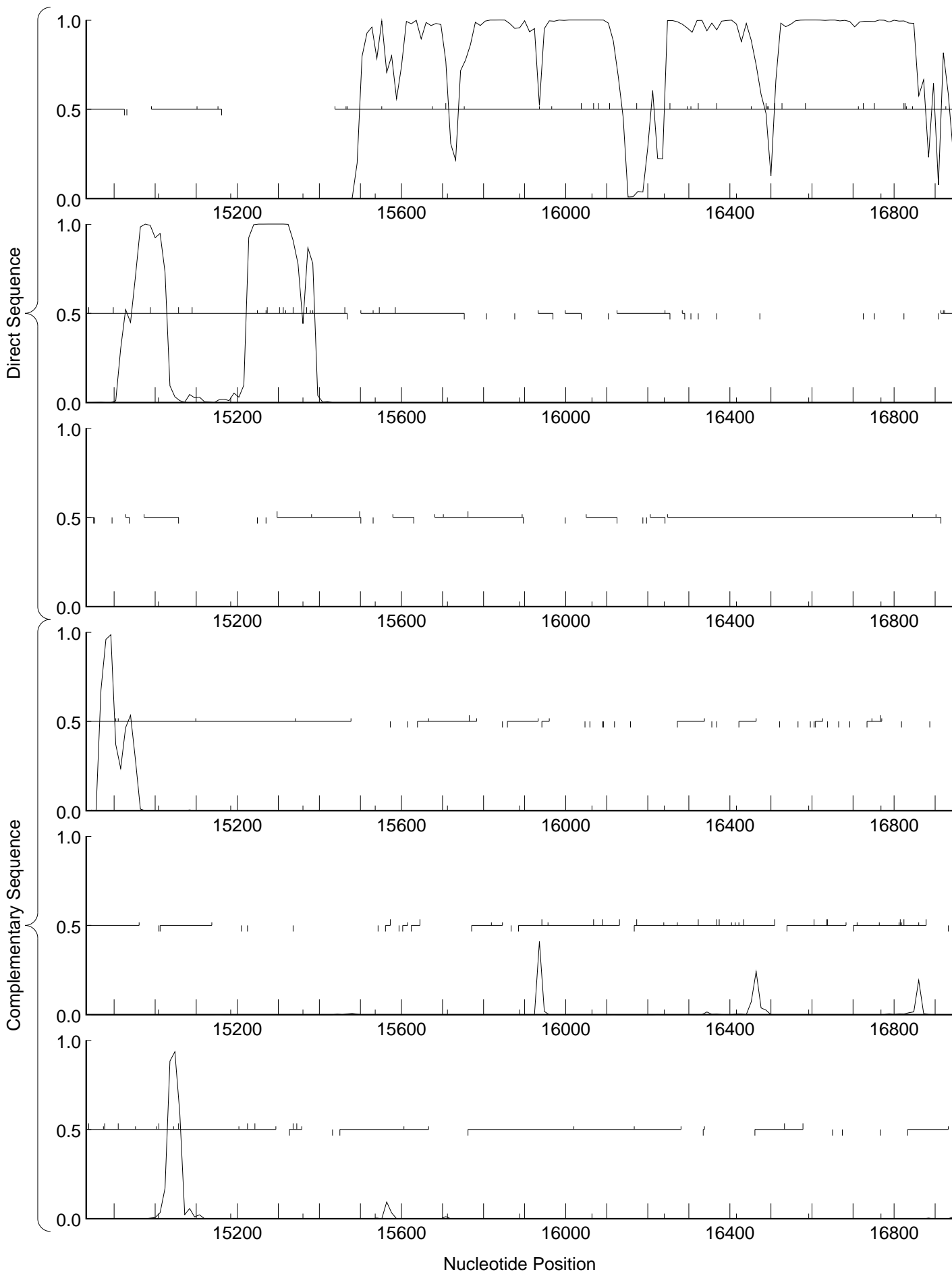




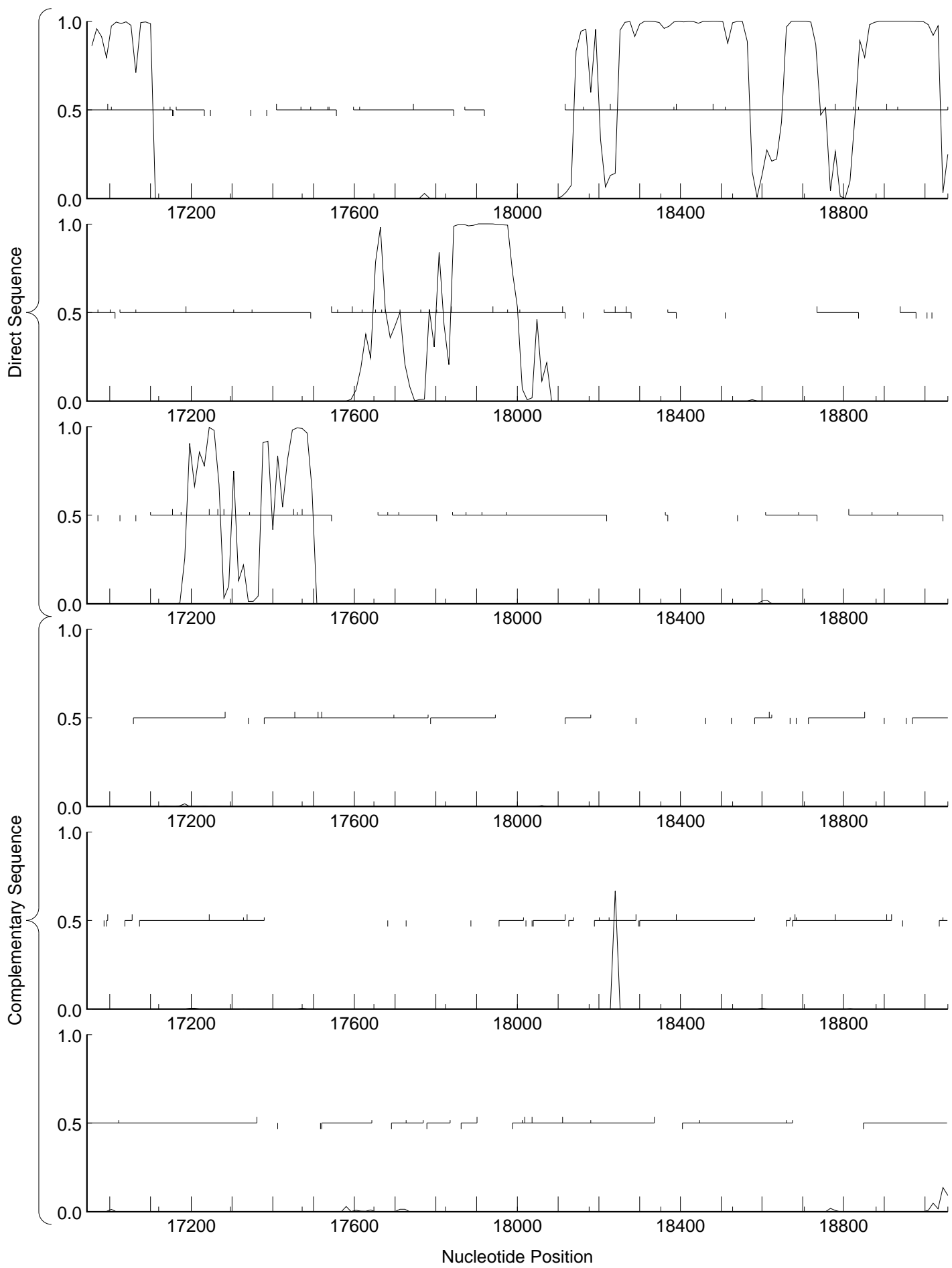




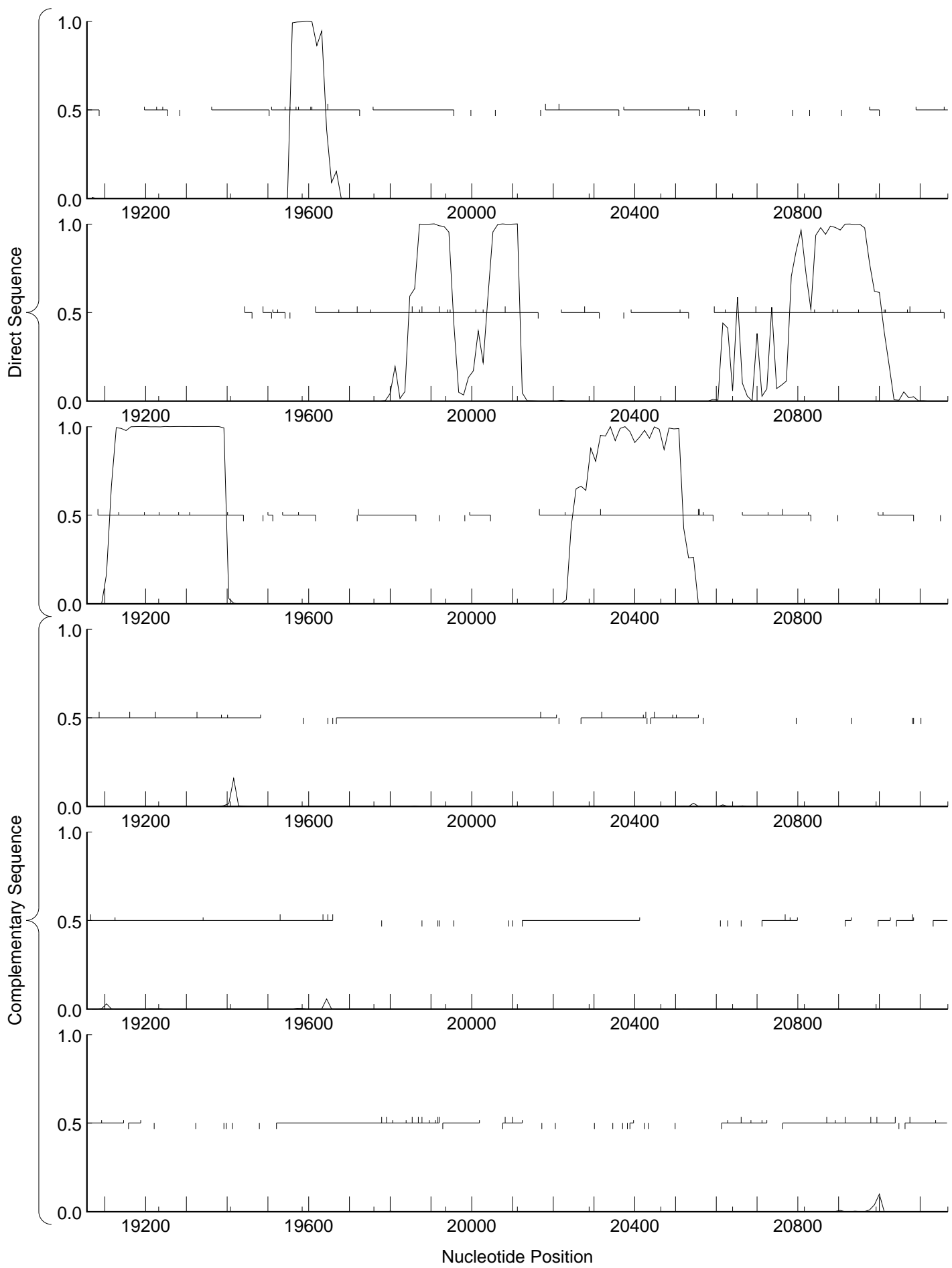
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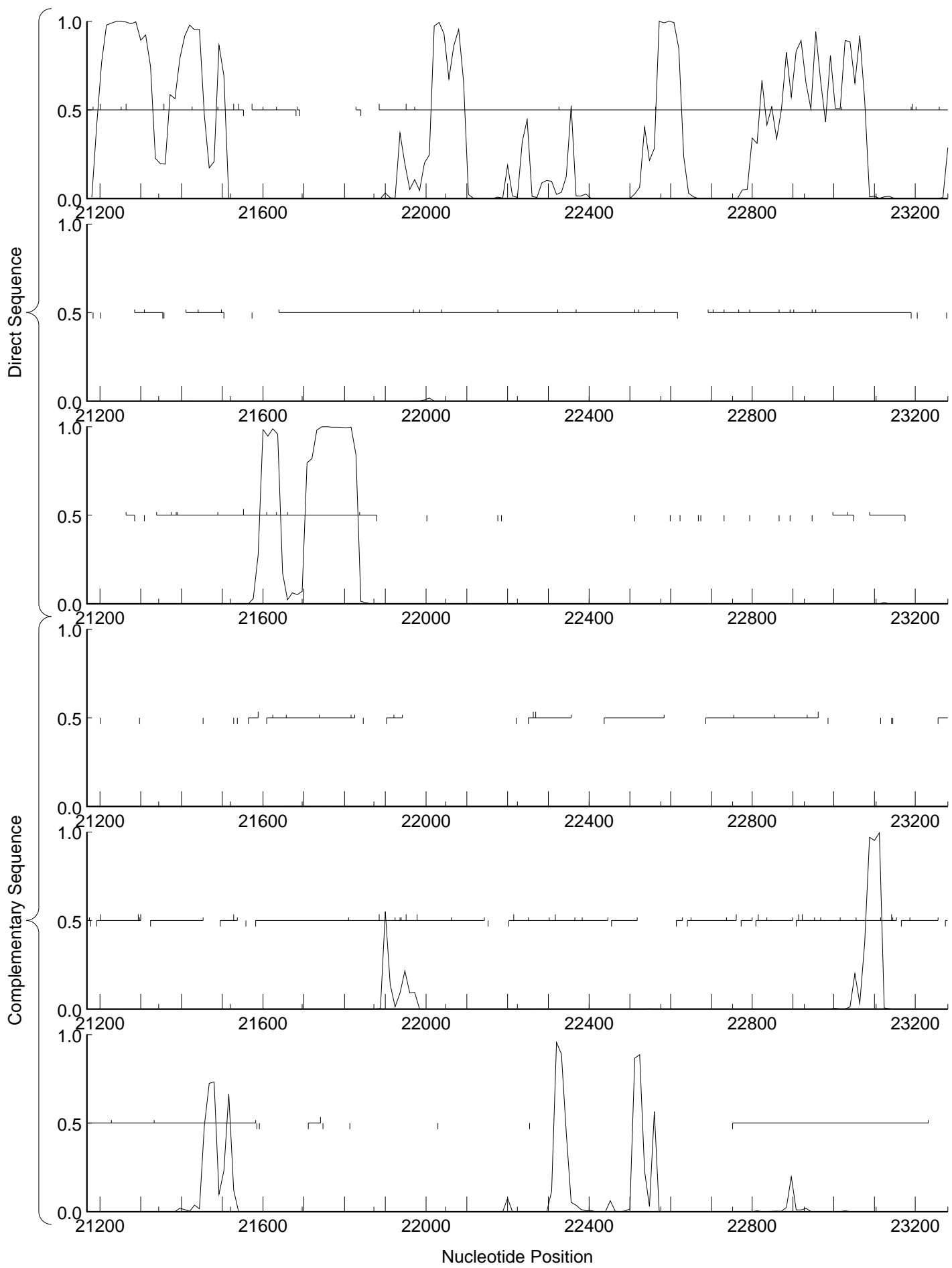
complete sequence, 46950 bp including 10-base 3' overhang (CGGTAGGCAT), Cluster CT, Order 4, Window 96, Step 12, 10/24



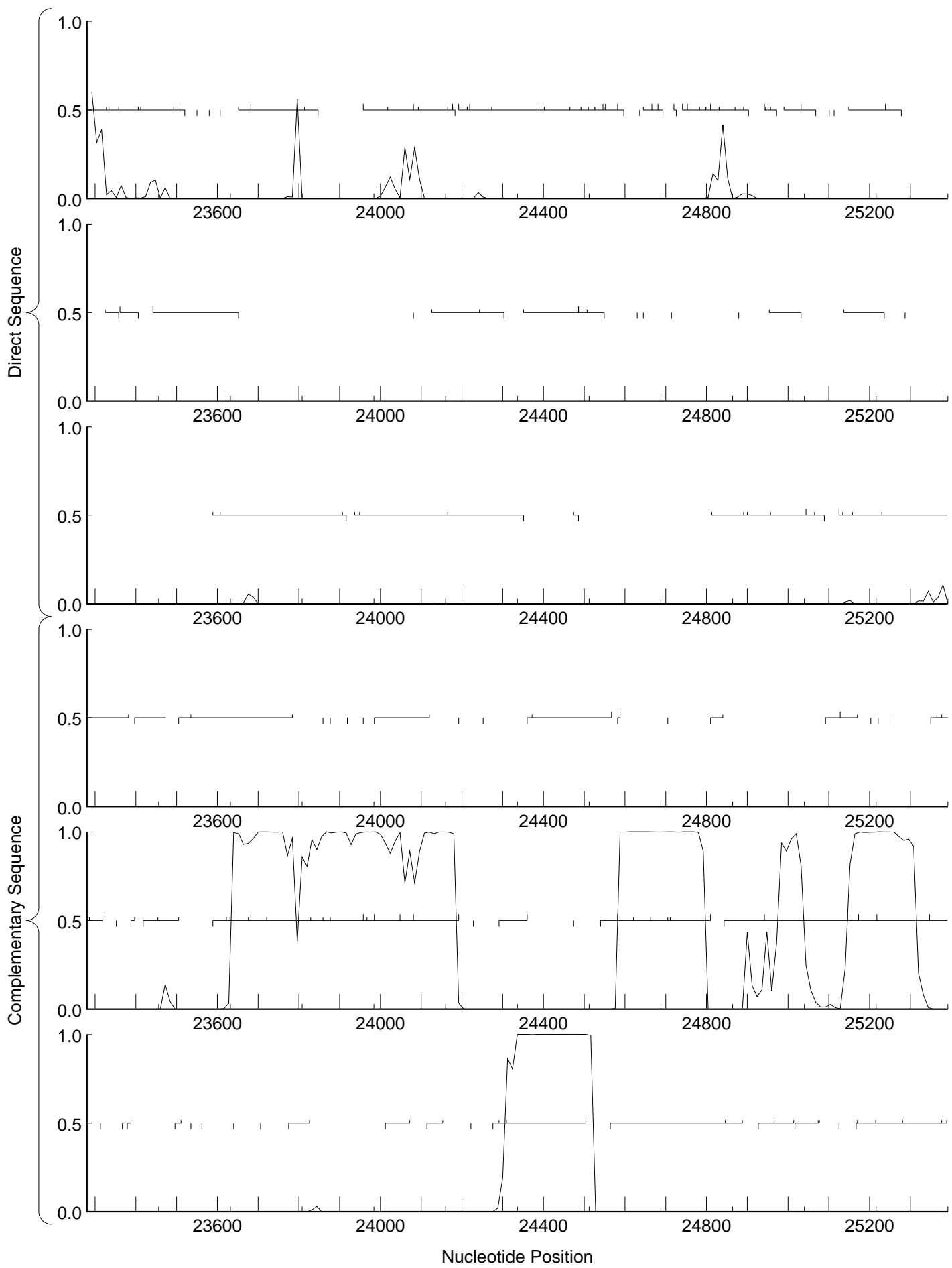
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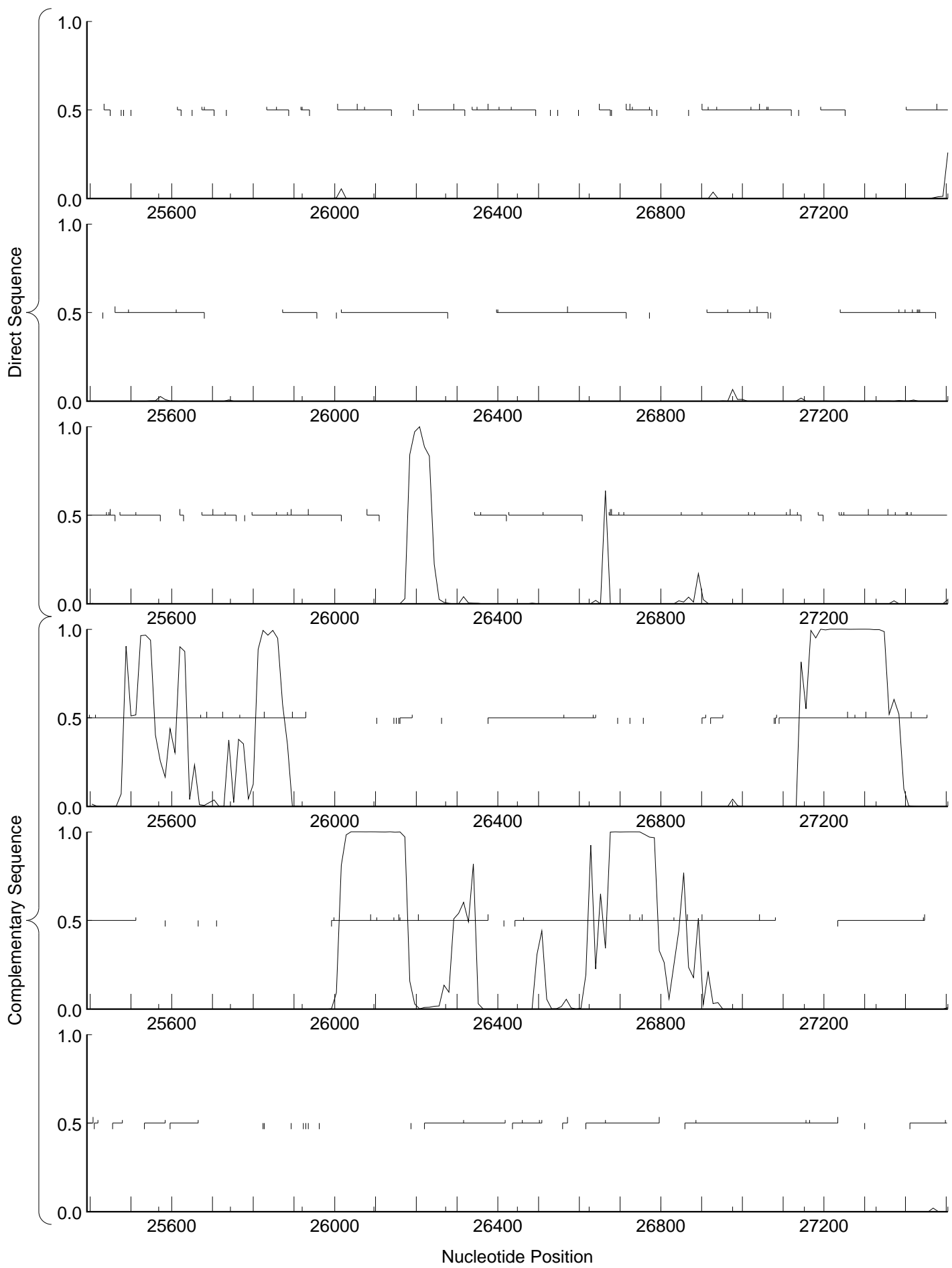
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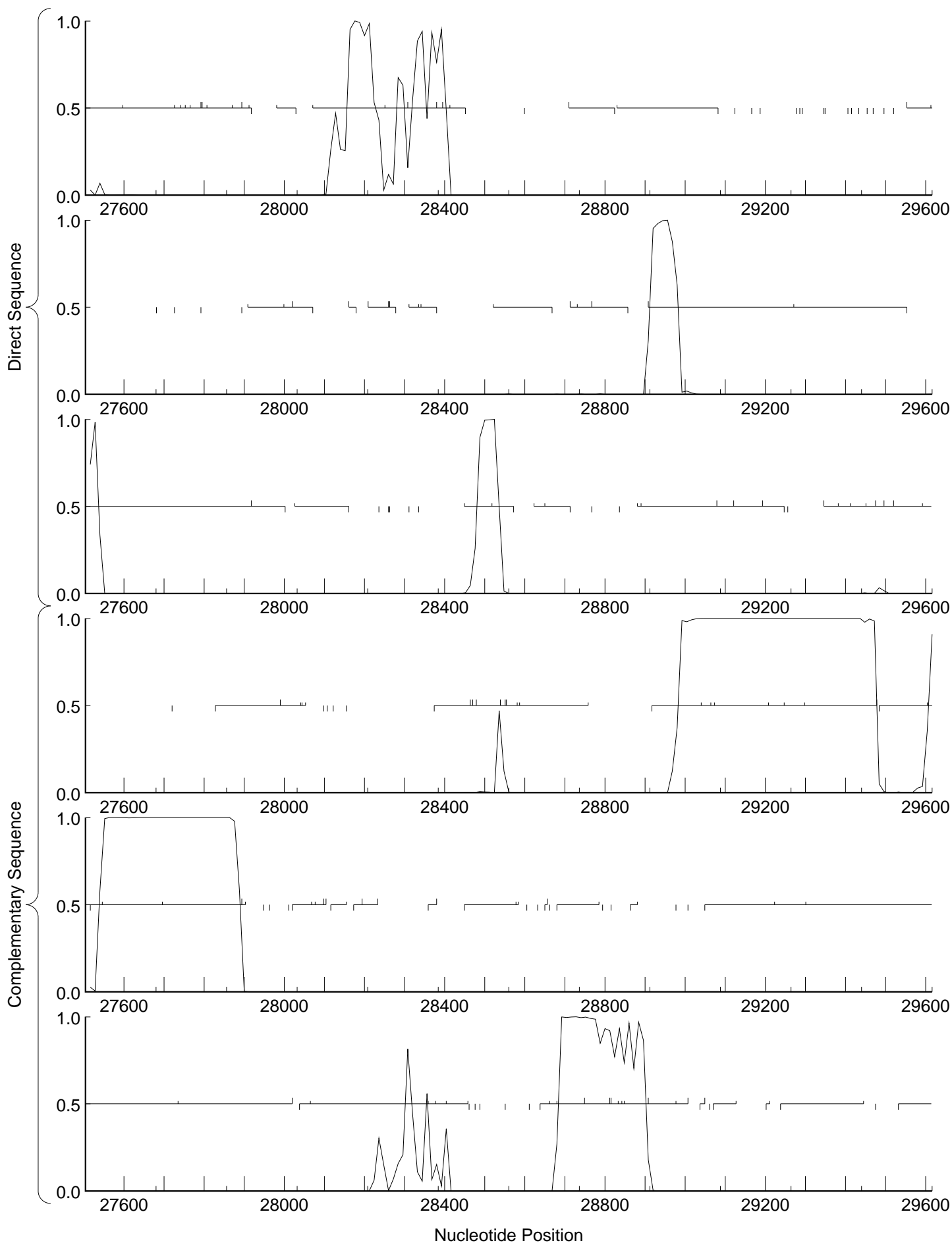
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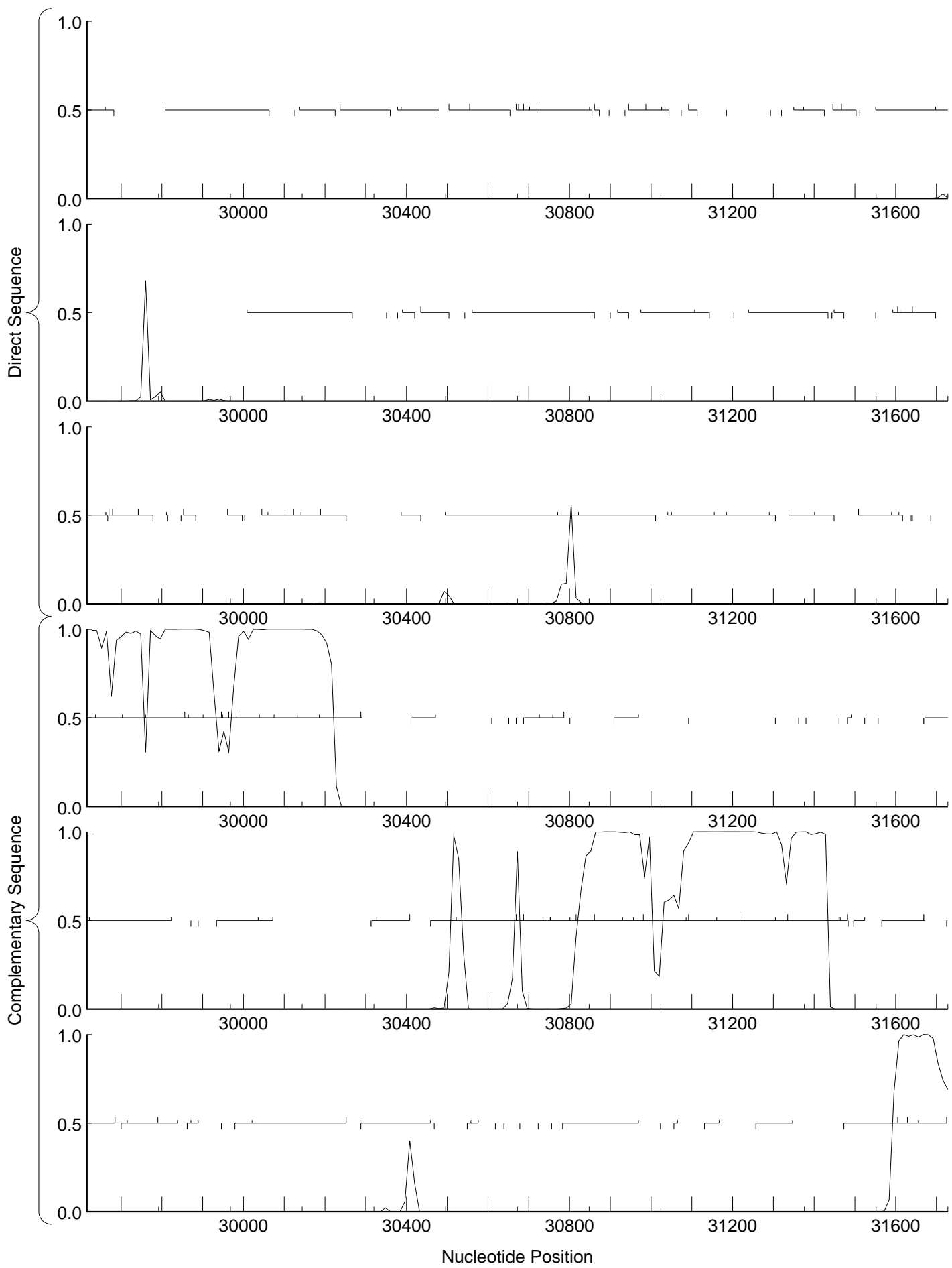
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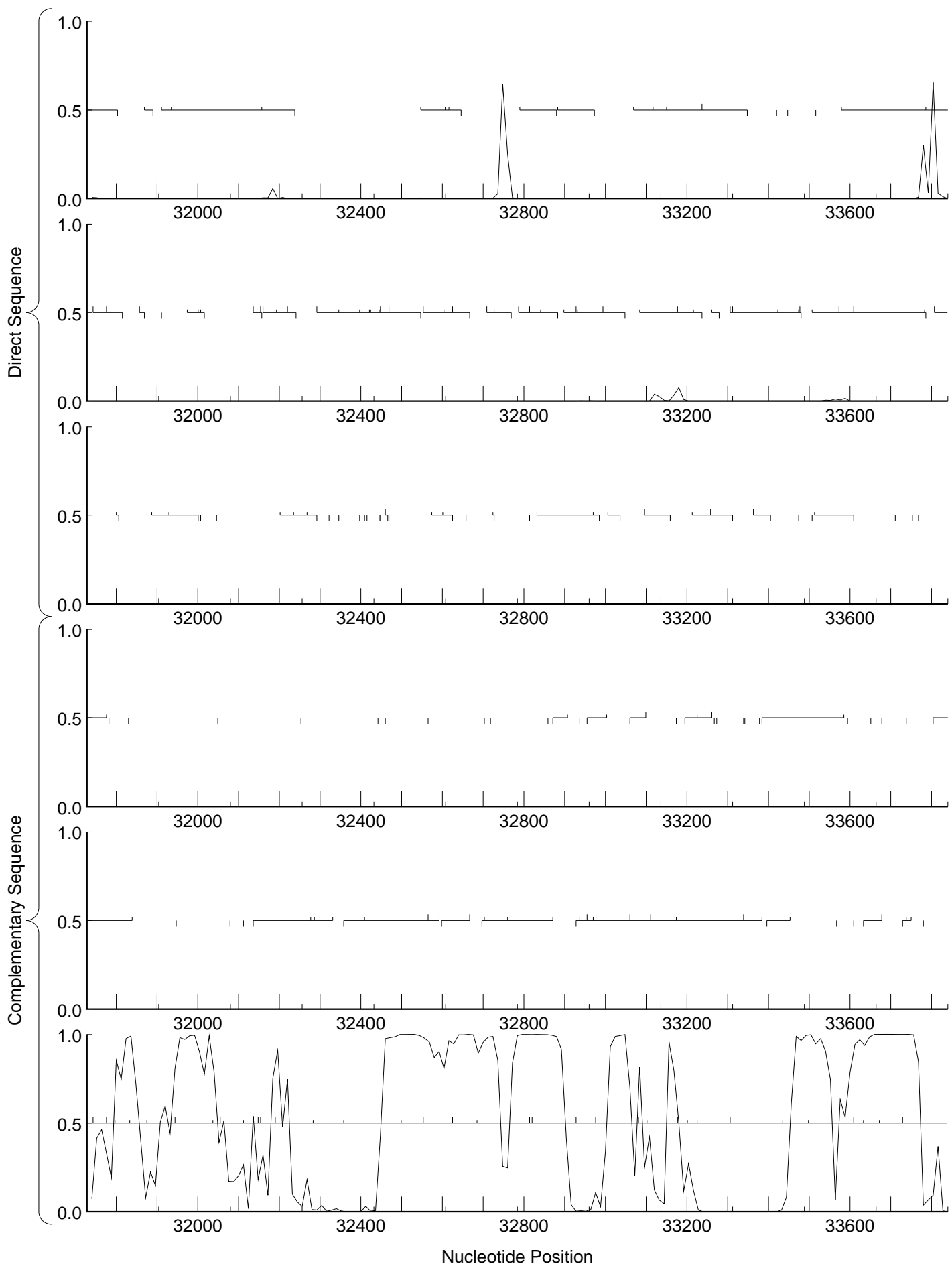
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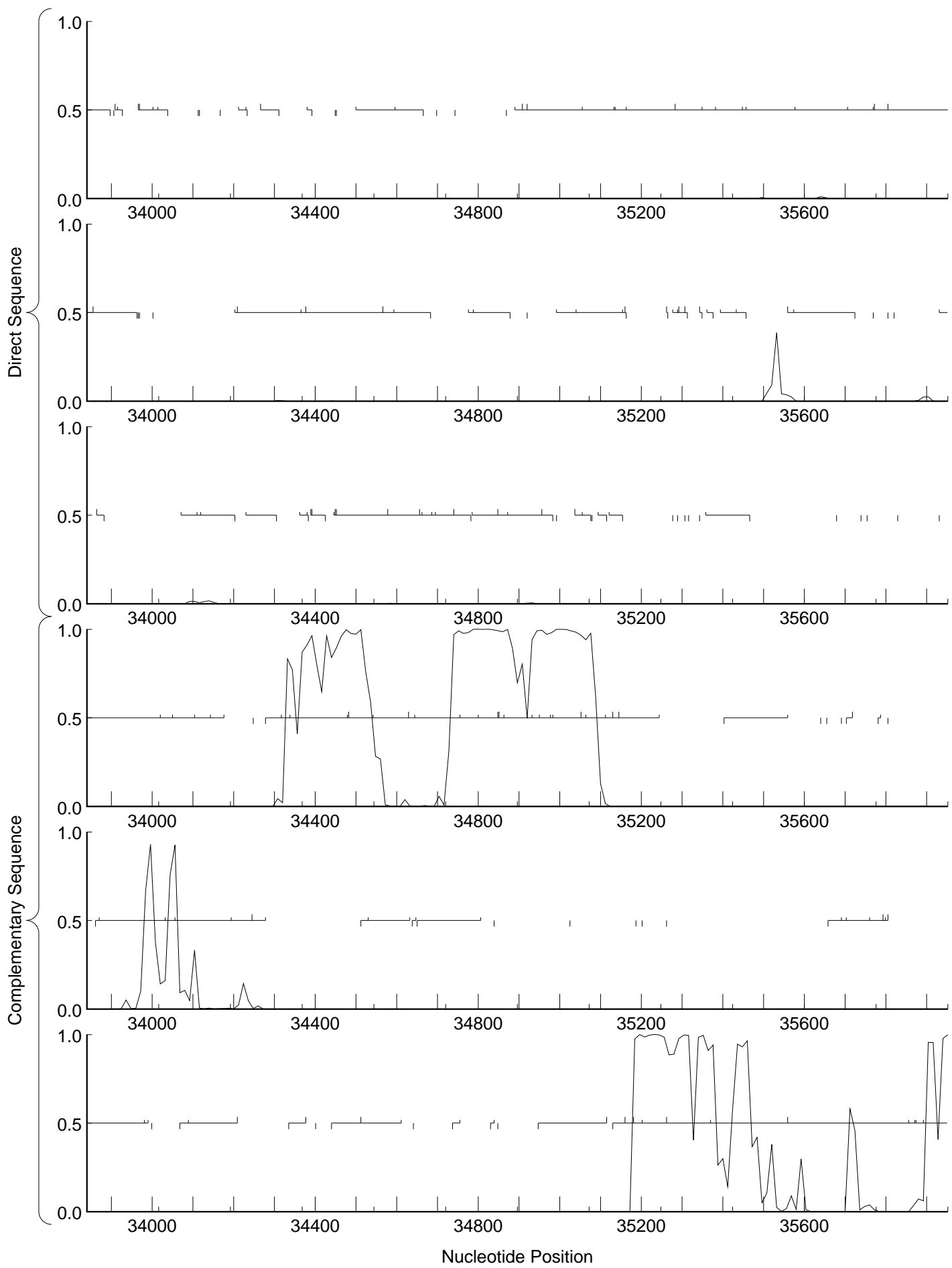


complete sequence, 46950 bp including 10-base 3' overhang (CGGTAGGCAT), Cluster CT, Order 4, Window 96, Step 12, 16/24

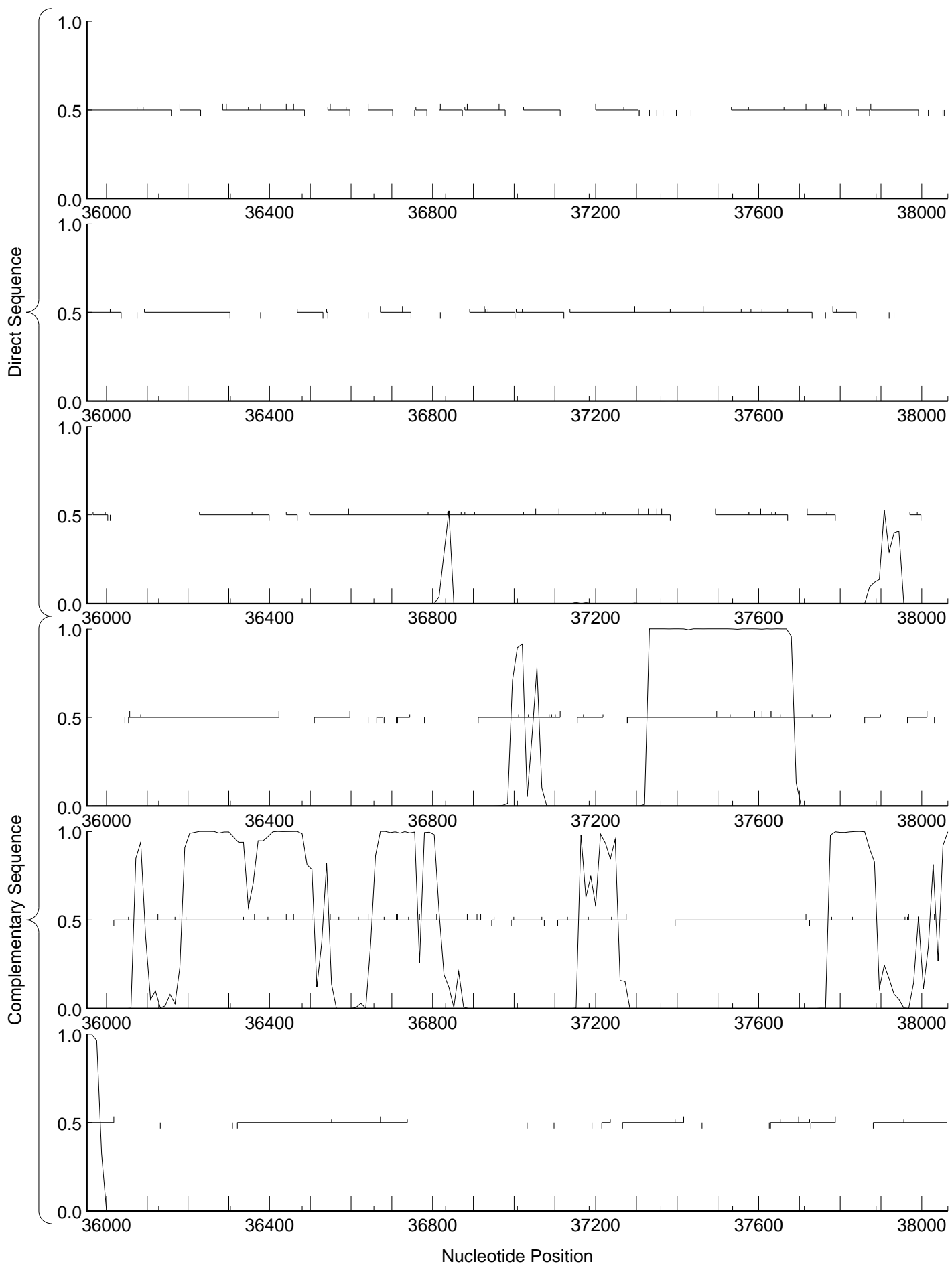


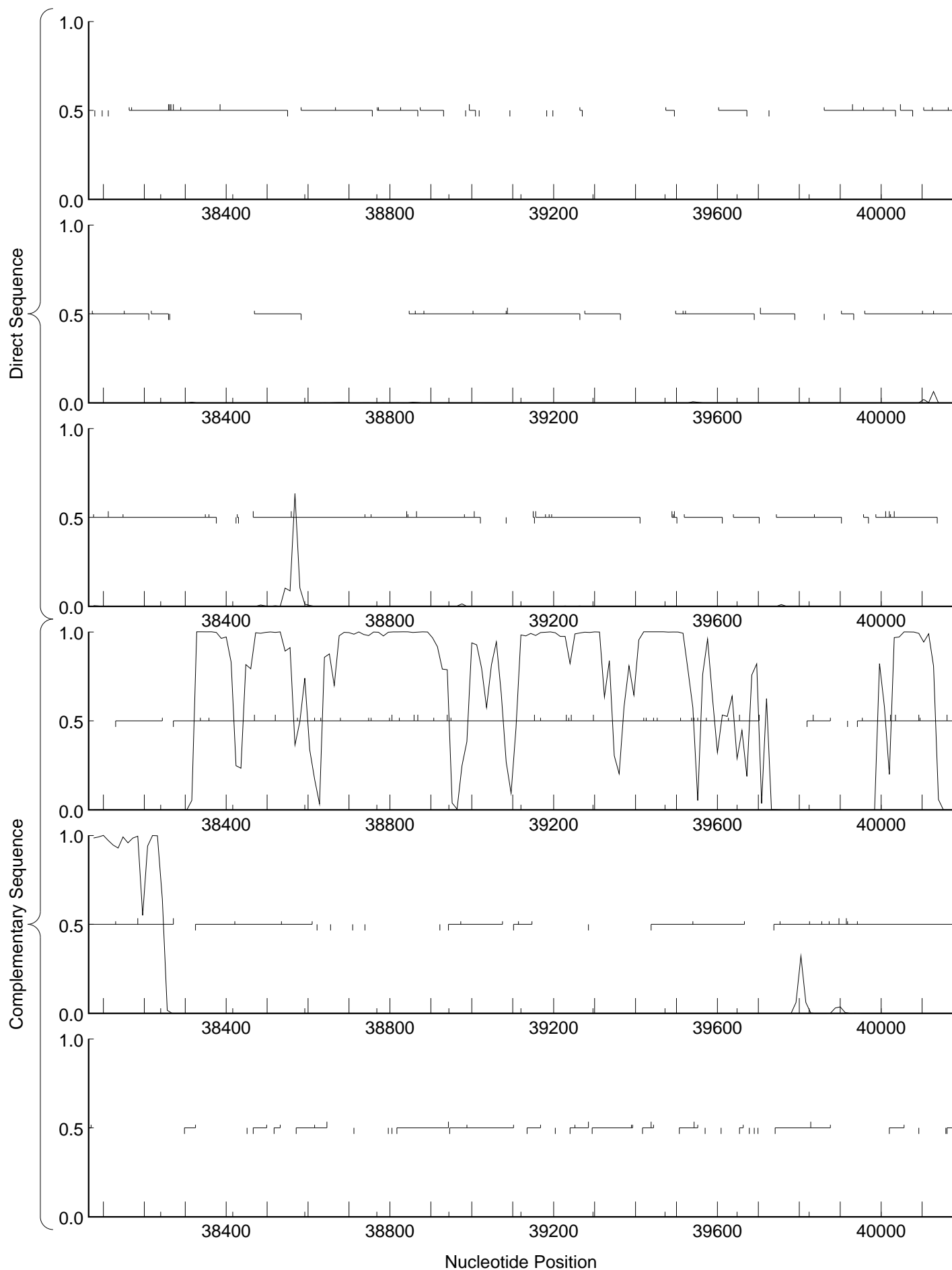
complete sequence, 46950 bp including 10-base 3' overhang (CGGTAGGCAT), Cluster CT, Order 4, Window 96, Step 12, 17/24

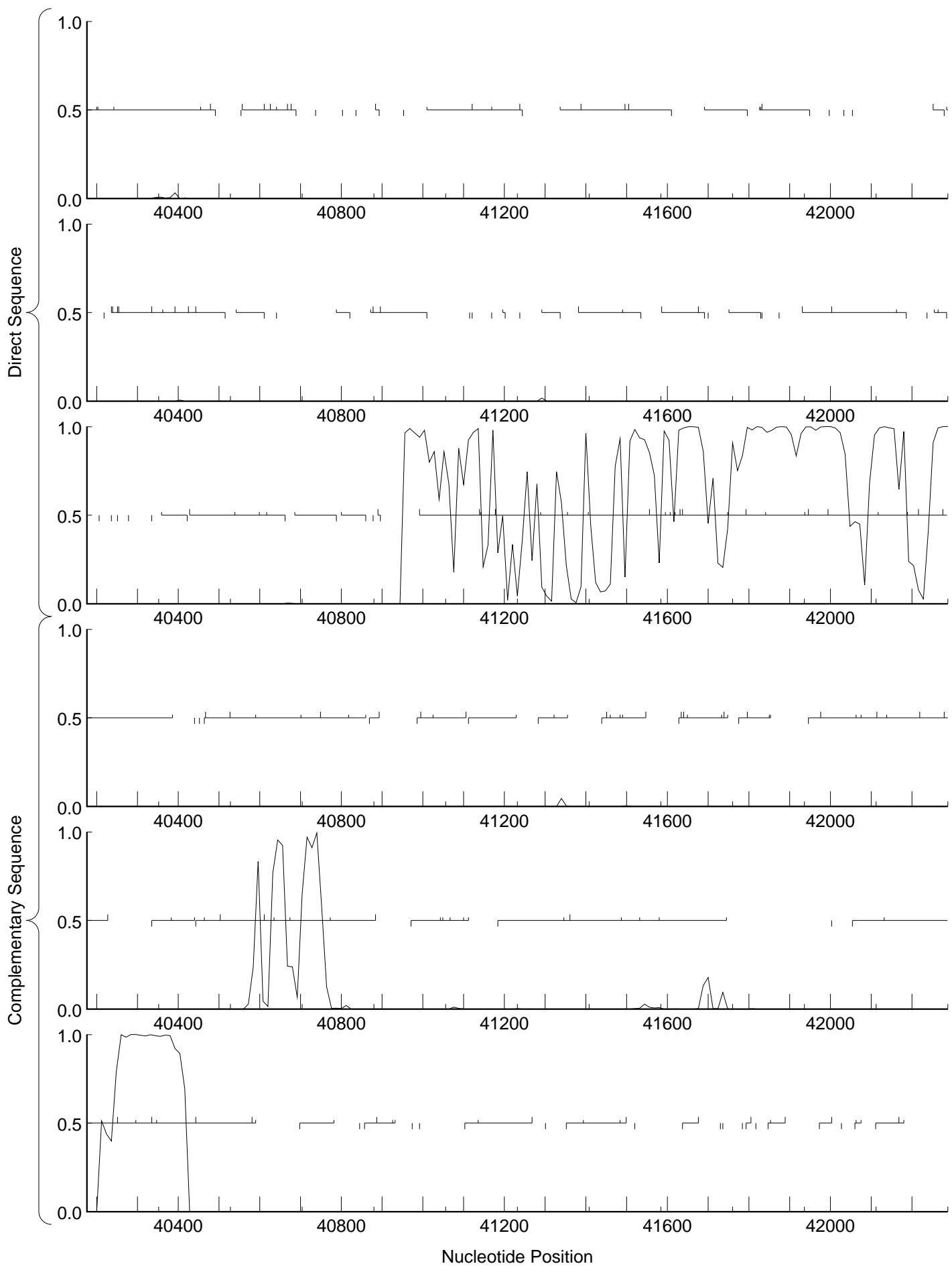


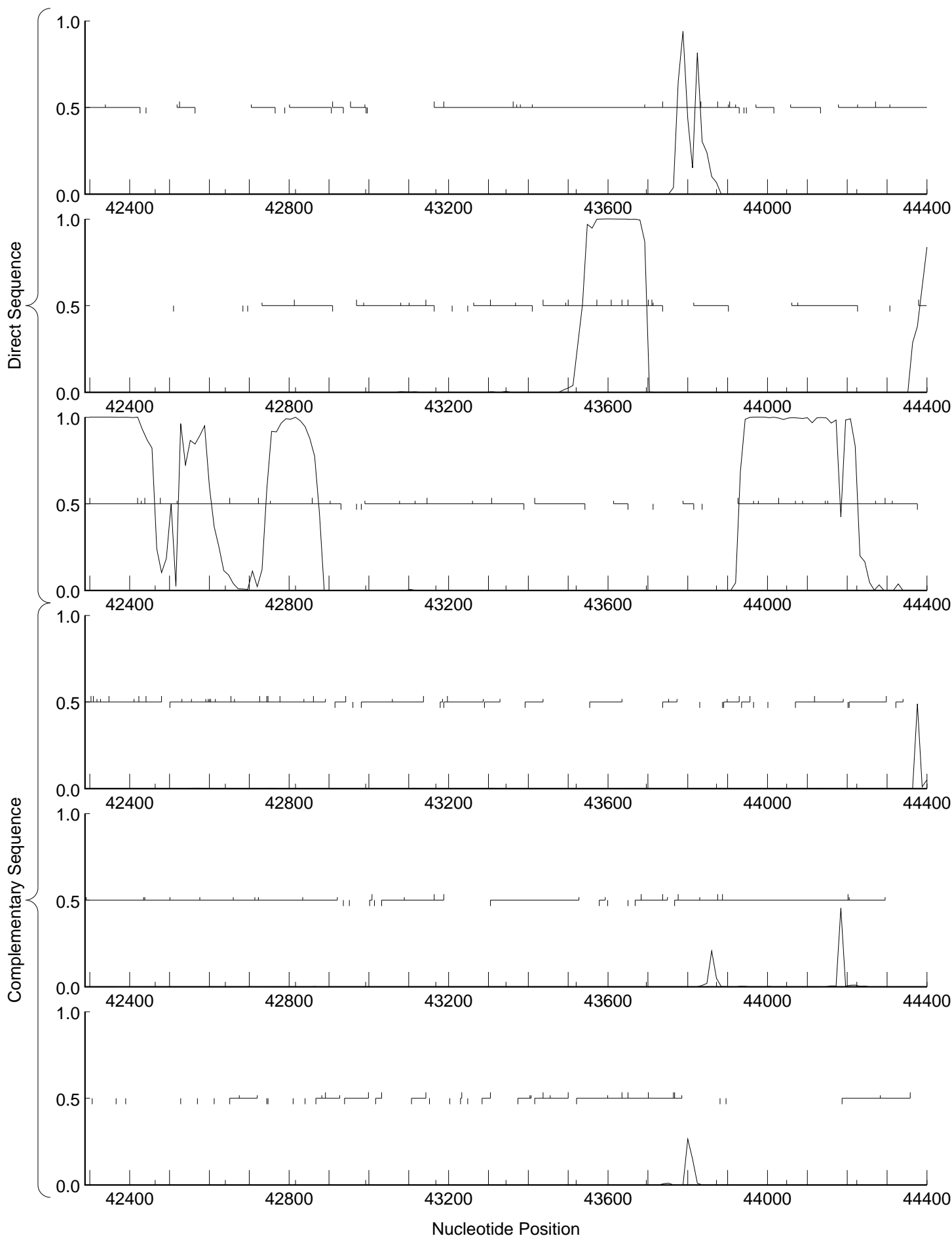


complete sequence, 46950 bp including 10-base 3' overhang (CGGTAGGCAT), Cluster CT, Order 4, Window 96, Step 12, 19/24









complete sequence, 46950 bp including 10-base 3' overhang (CGGTAGGCAT), Cluster CT, Order 4, Window 96, Step 12, 23/24

