

GeneMark

Version 2.5p (09.08.06)

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PROGRAM INFORMATION

Sequence : Rhodococcus phage Alpacados complete sequence, 46493 bp including 10-base 3' overhang (CGGC
Analysis Date : 5/24/18 at 13:33:47
Pages : 24
Sequence Length : 46493 bp
GC Content : 58.85%

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,

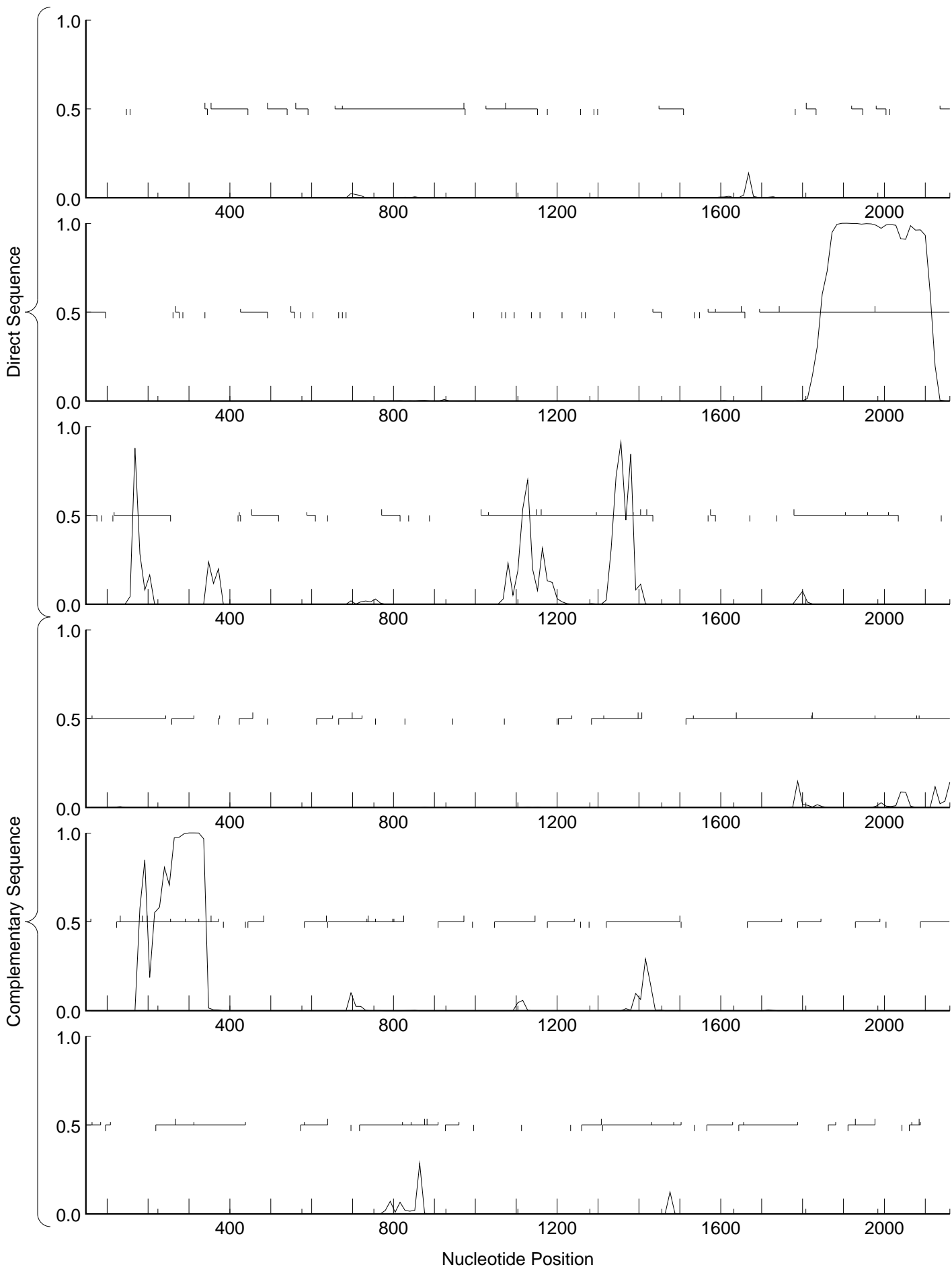
MATRIX INFORMATION

Matrix : Mycobacterium_tuberculosis_H37Rv
Author : -
Order : 4

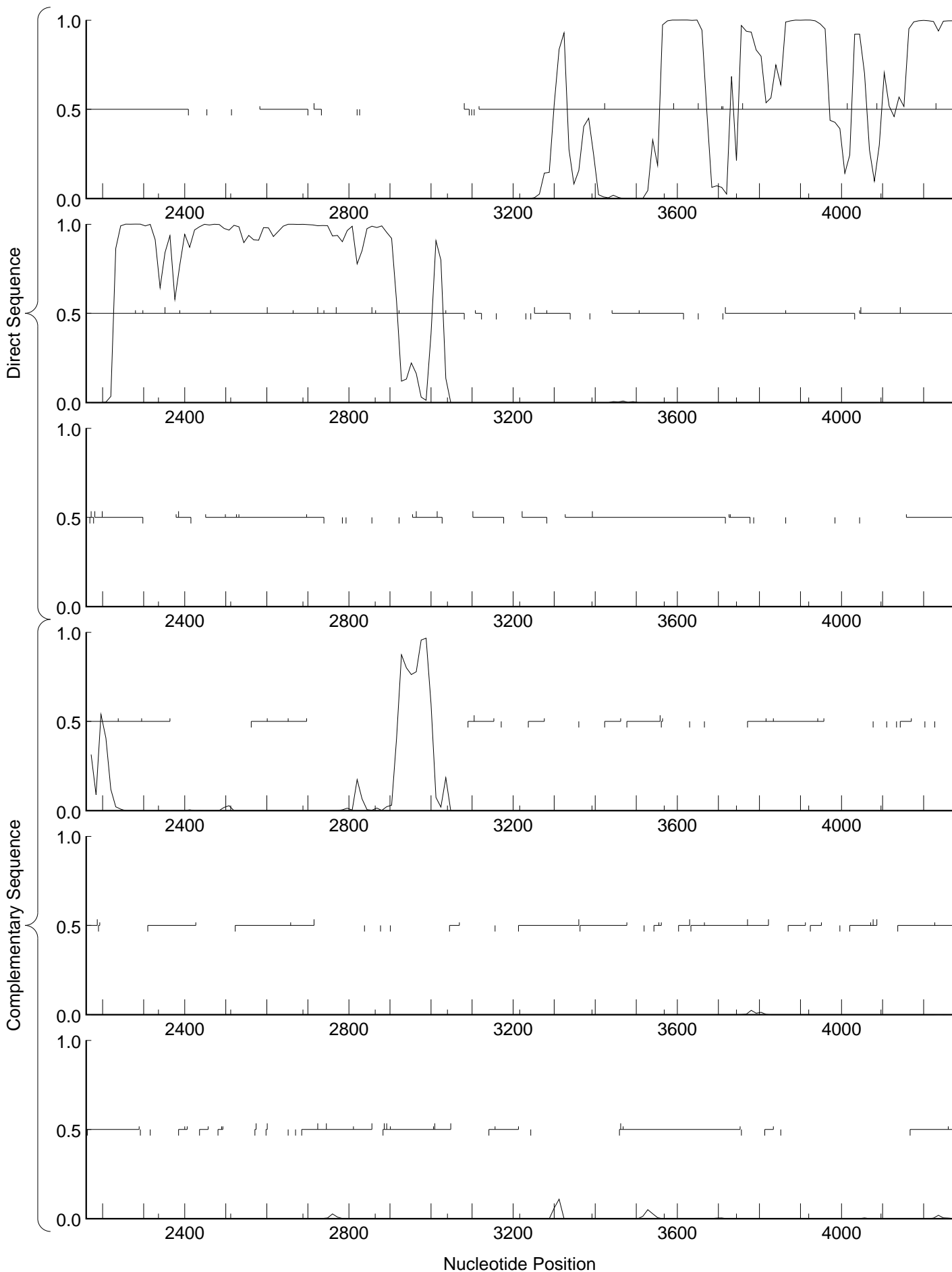
Send questions / comments to:
Dr. M. Borodovsky
Georgia Institute of Technology
School of Biology
Atlanta, GA 30332-0230

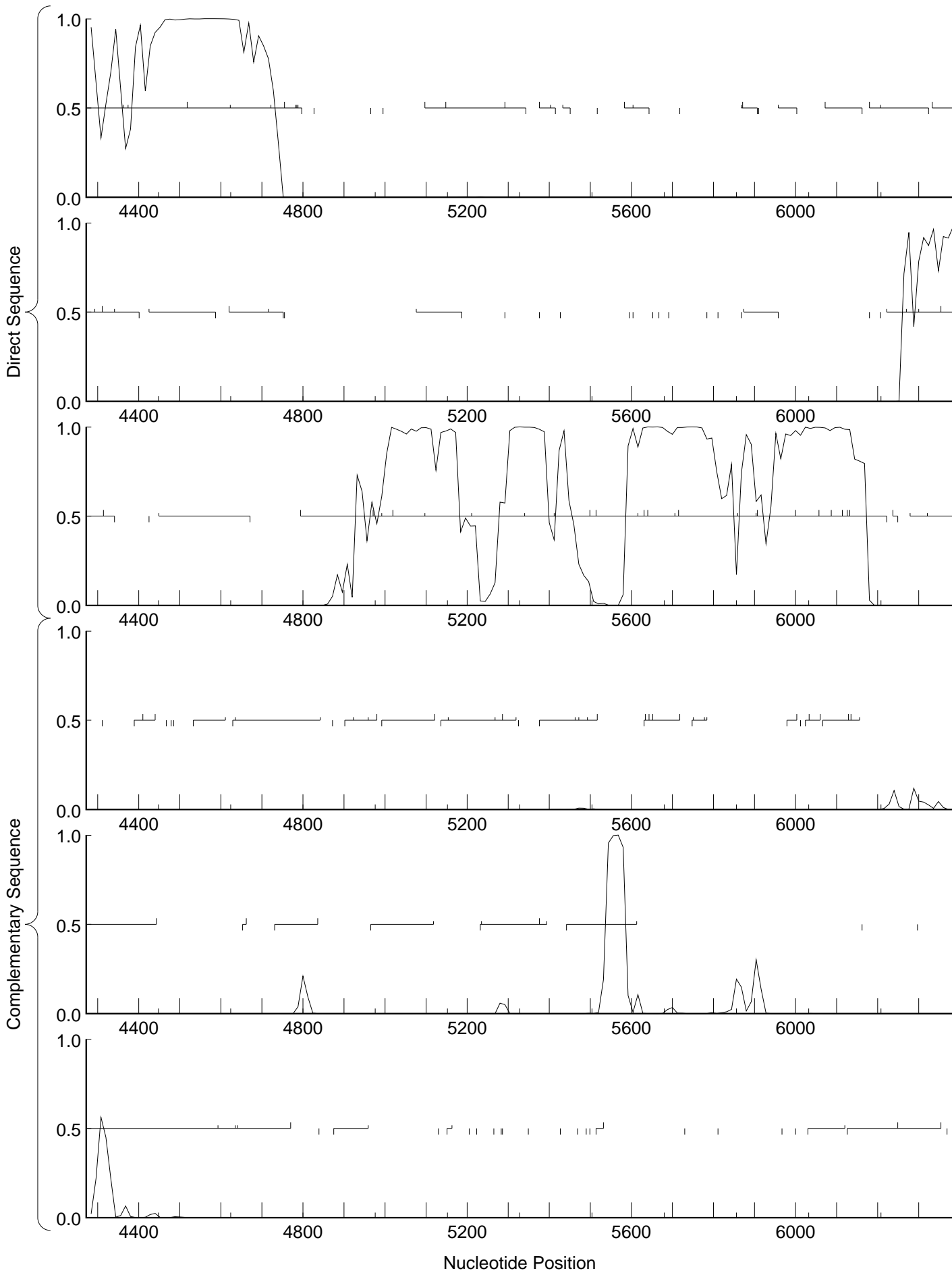
Matrix notes & comments

Training set derived by GeneMarkS, 4.27 September 2014
Tue Sep 23 15:14:37 2014

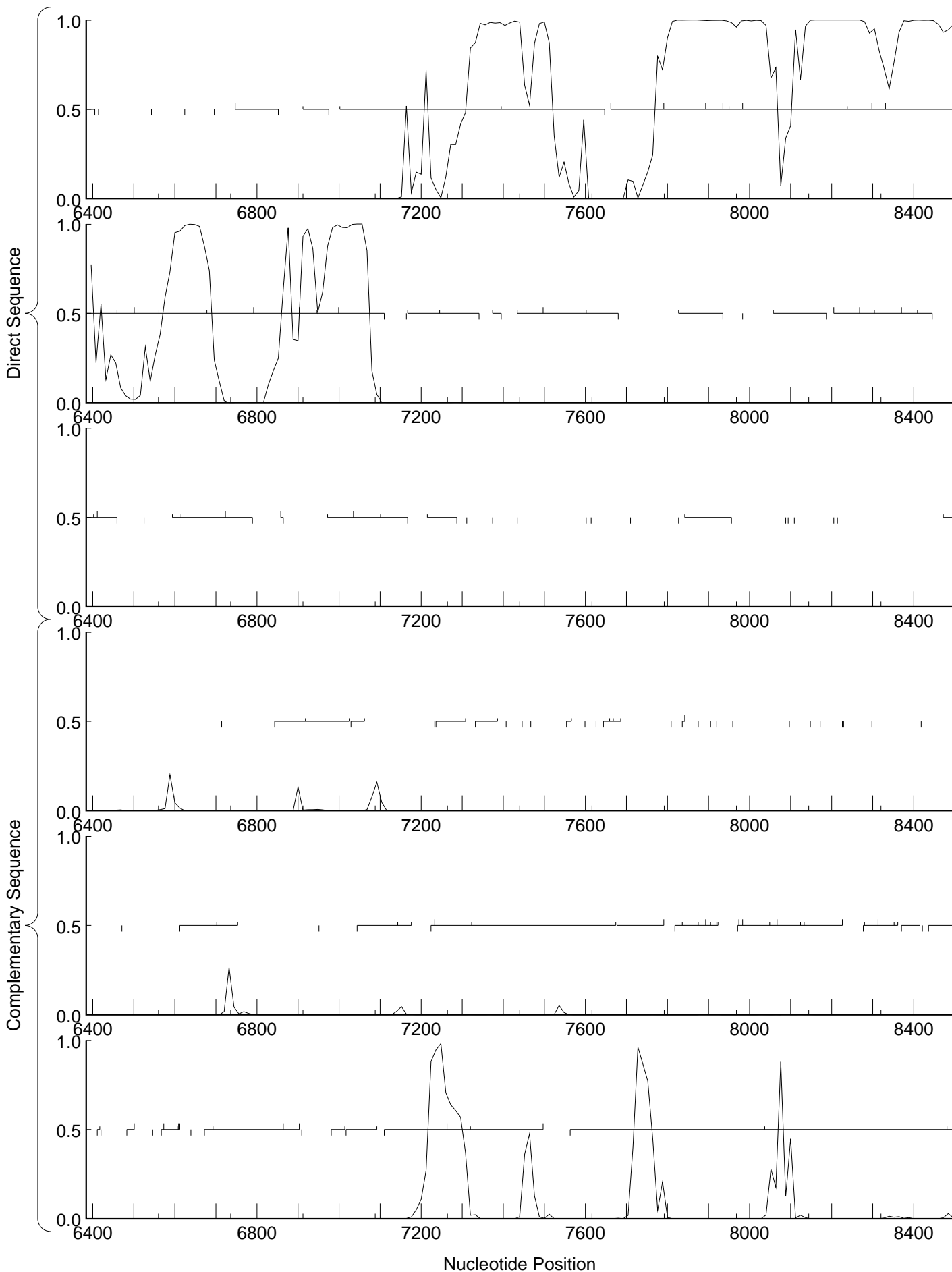


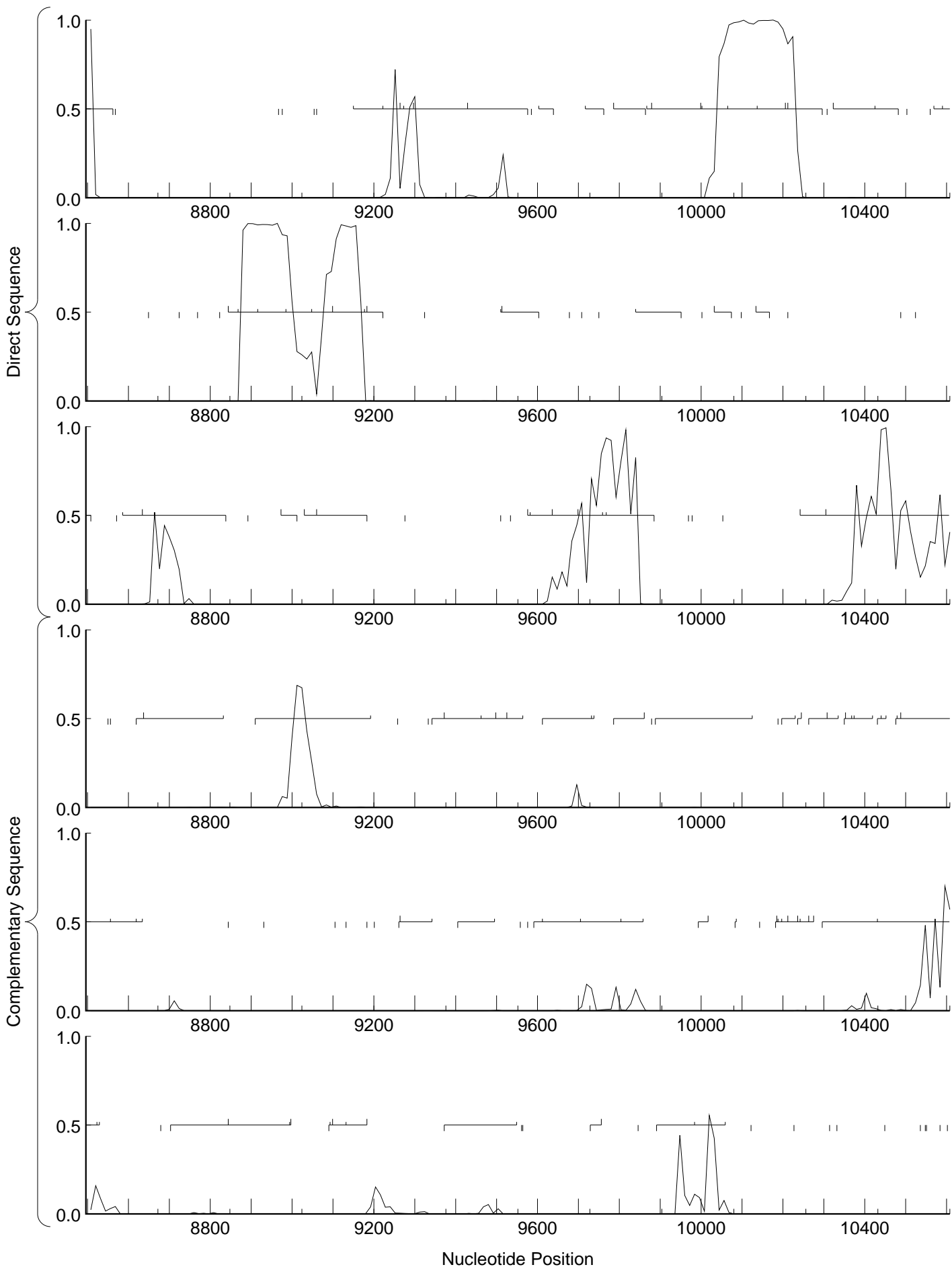
s complete sequence, 46493 bp including 10-base 3' overhang (CGGCCGTGAT), Cluster CA, Order 4, Window 96, Step 12, 3/24



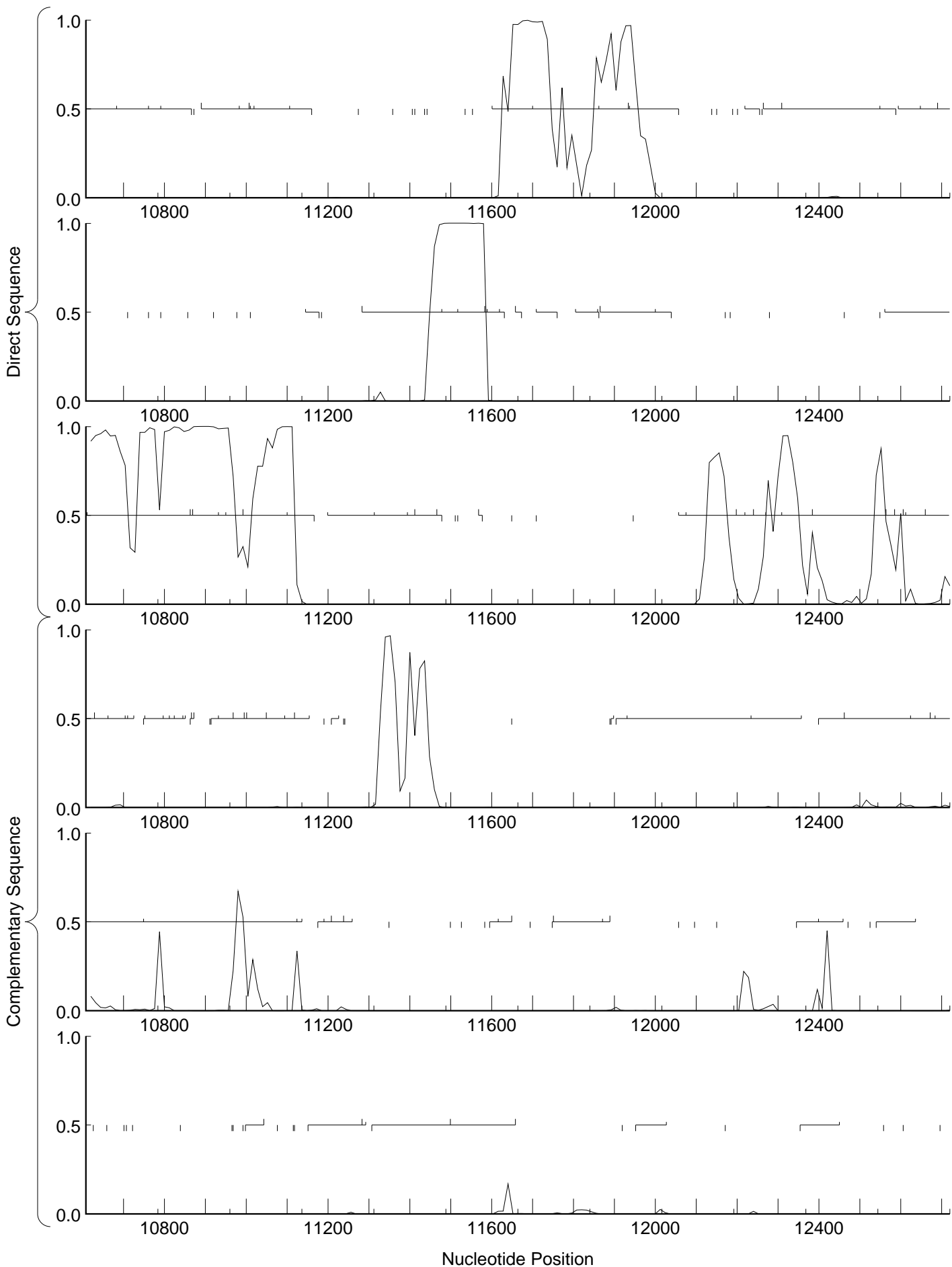


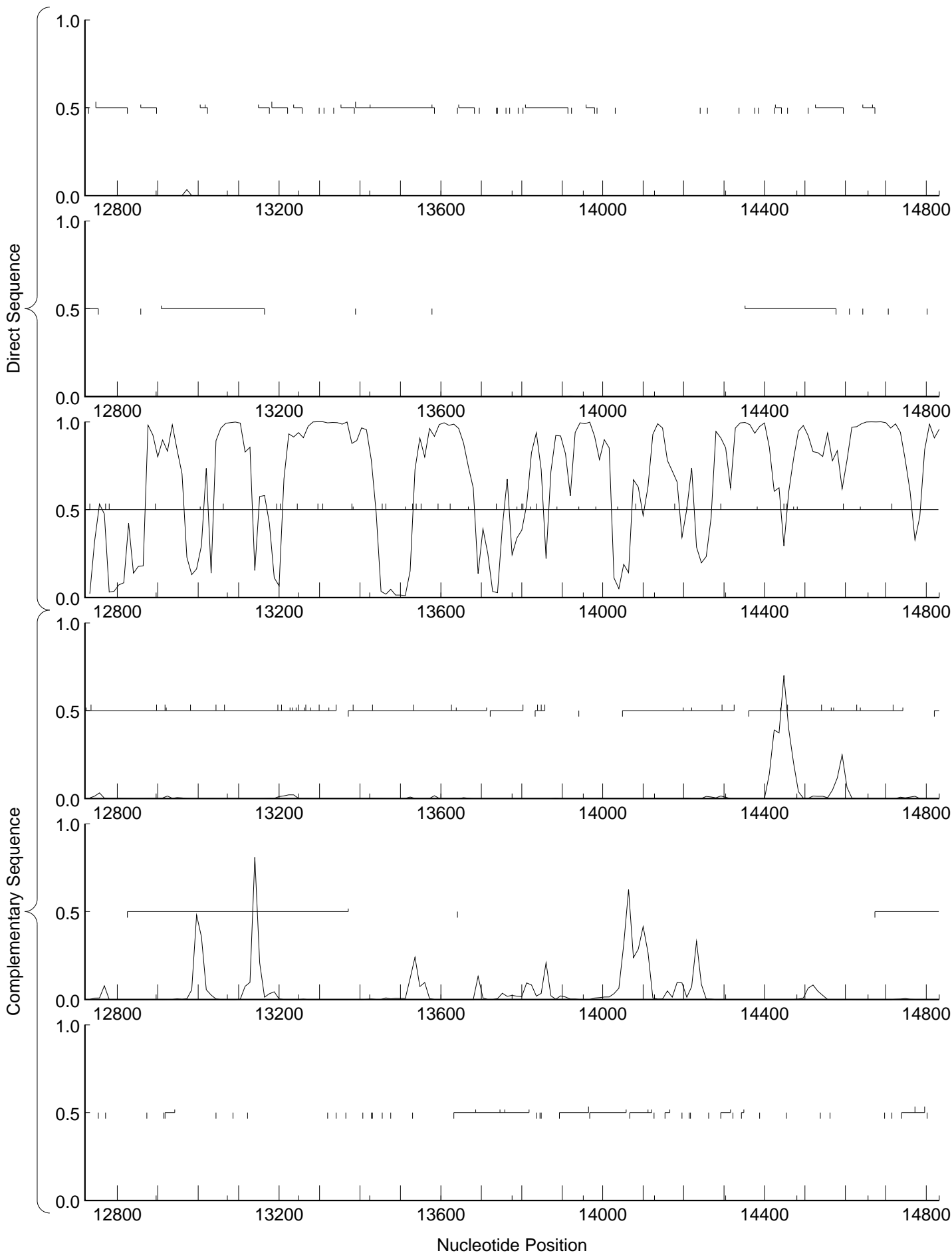
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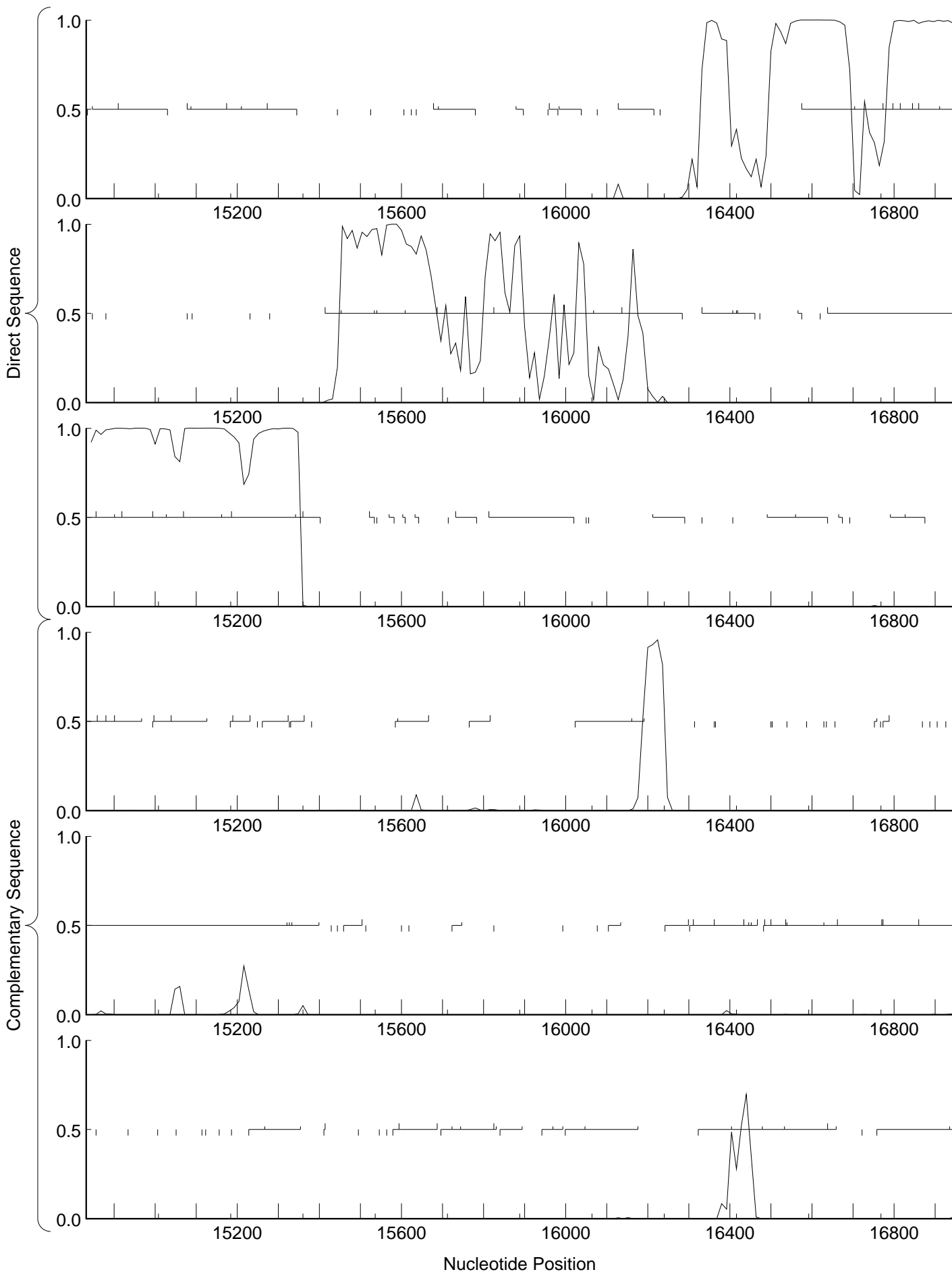




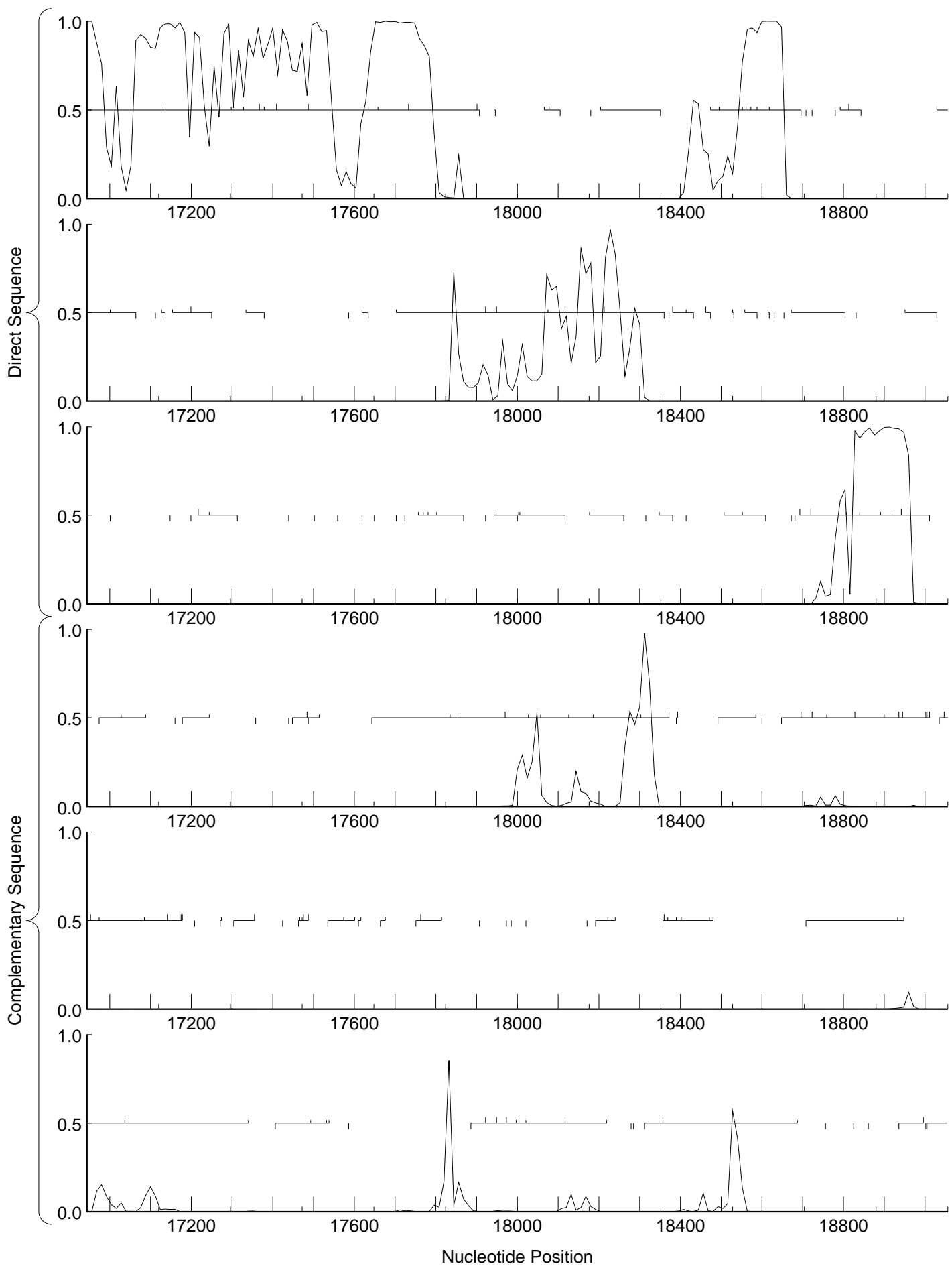
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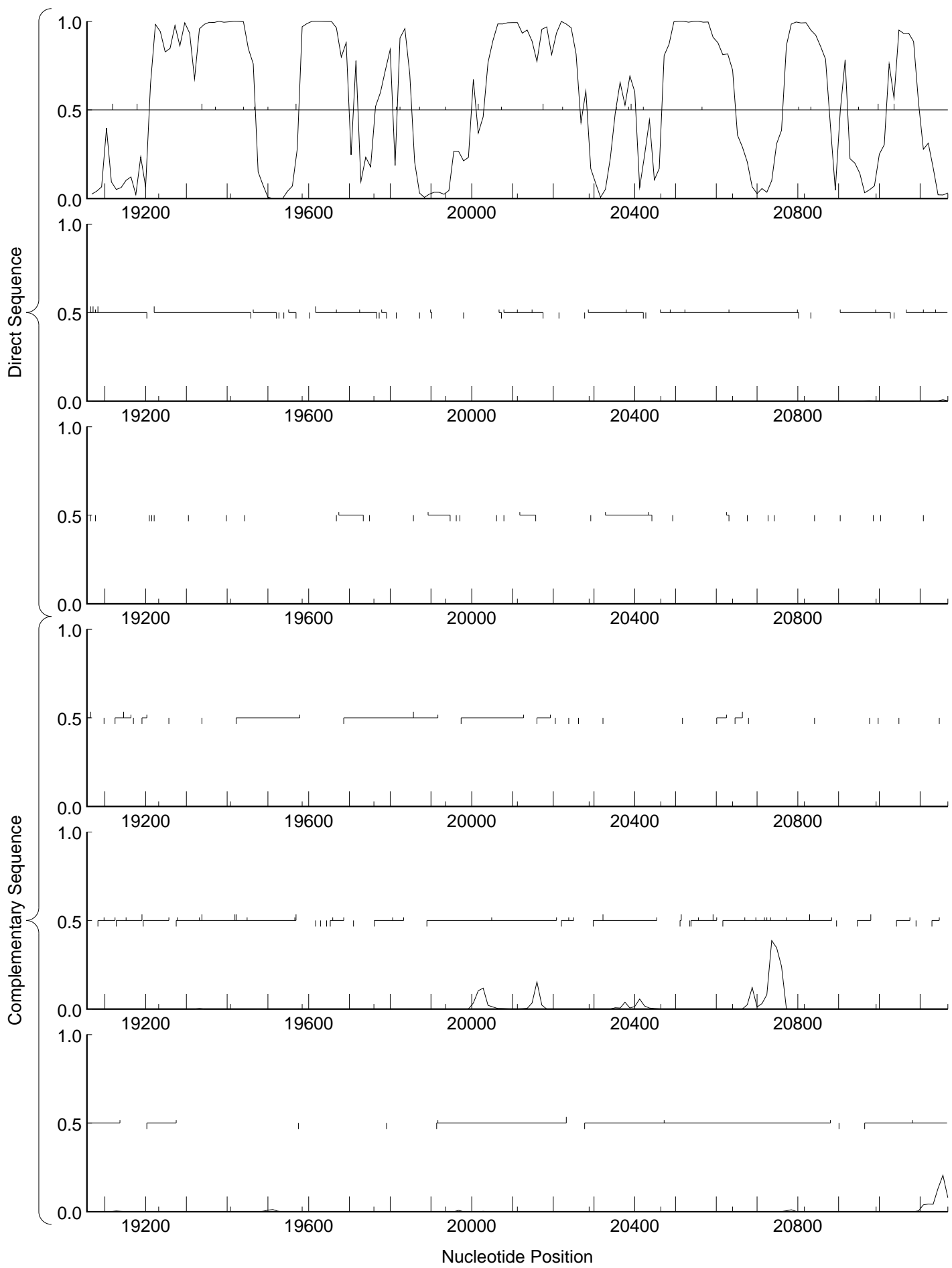


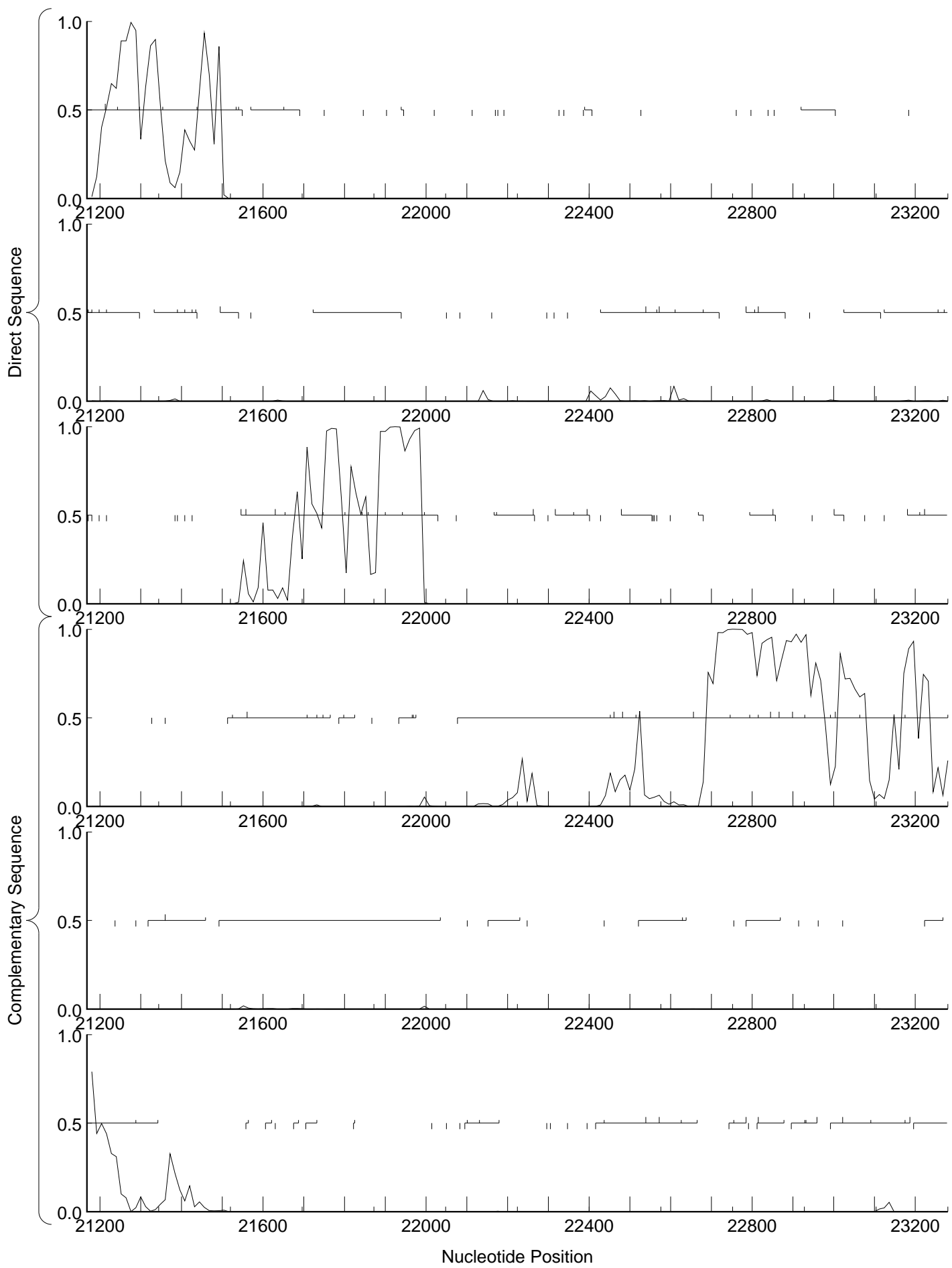


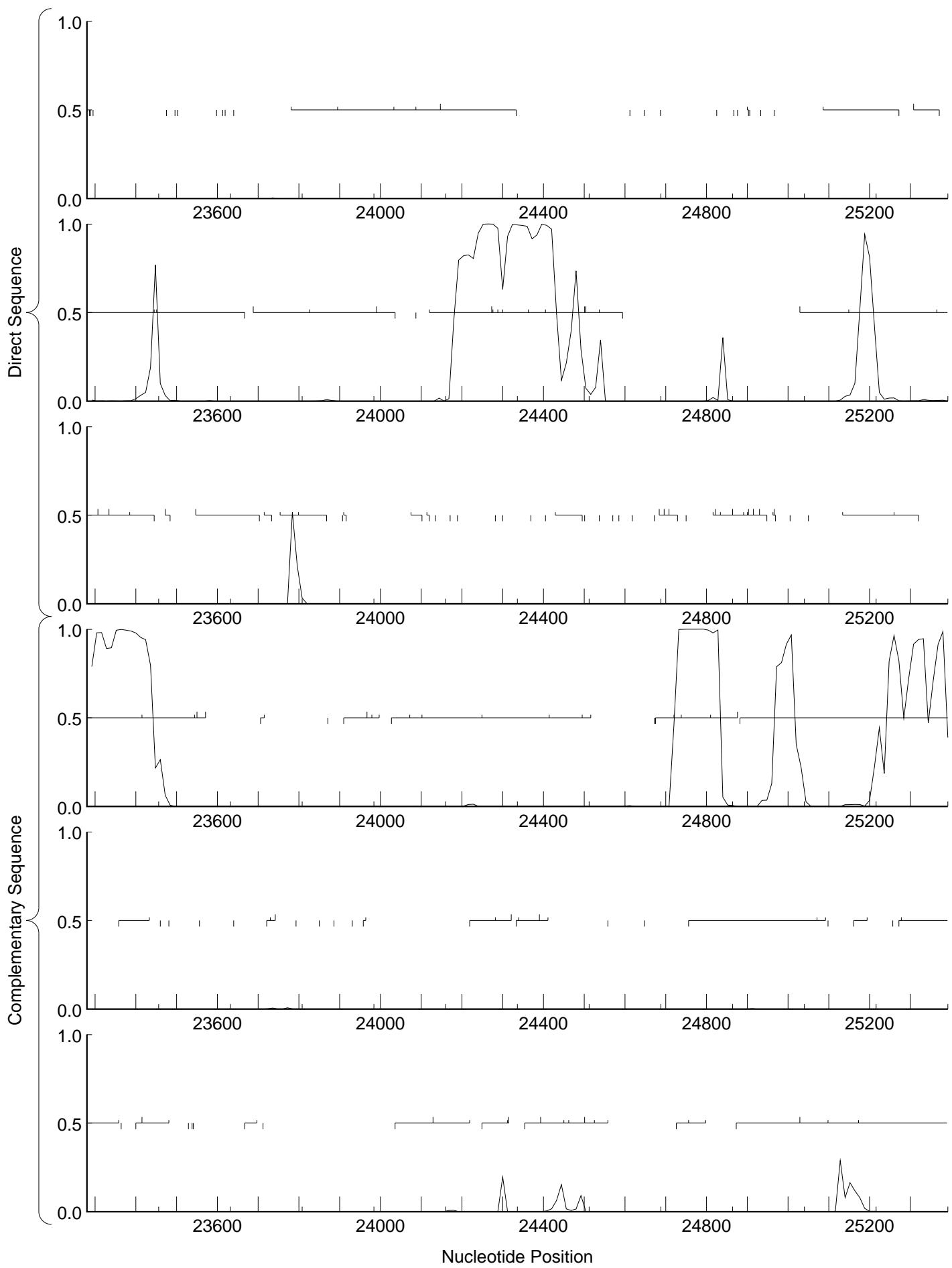
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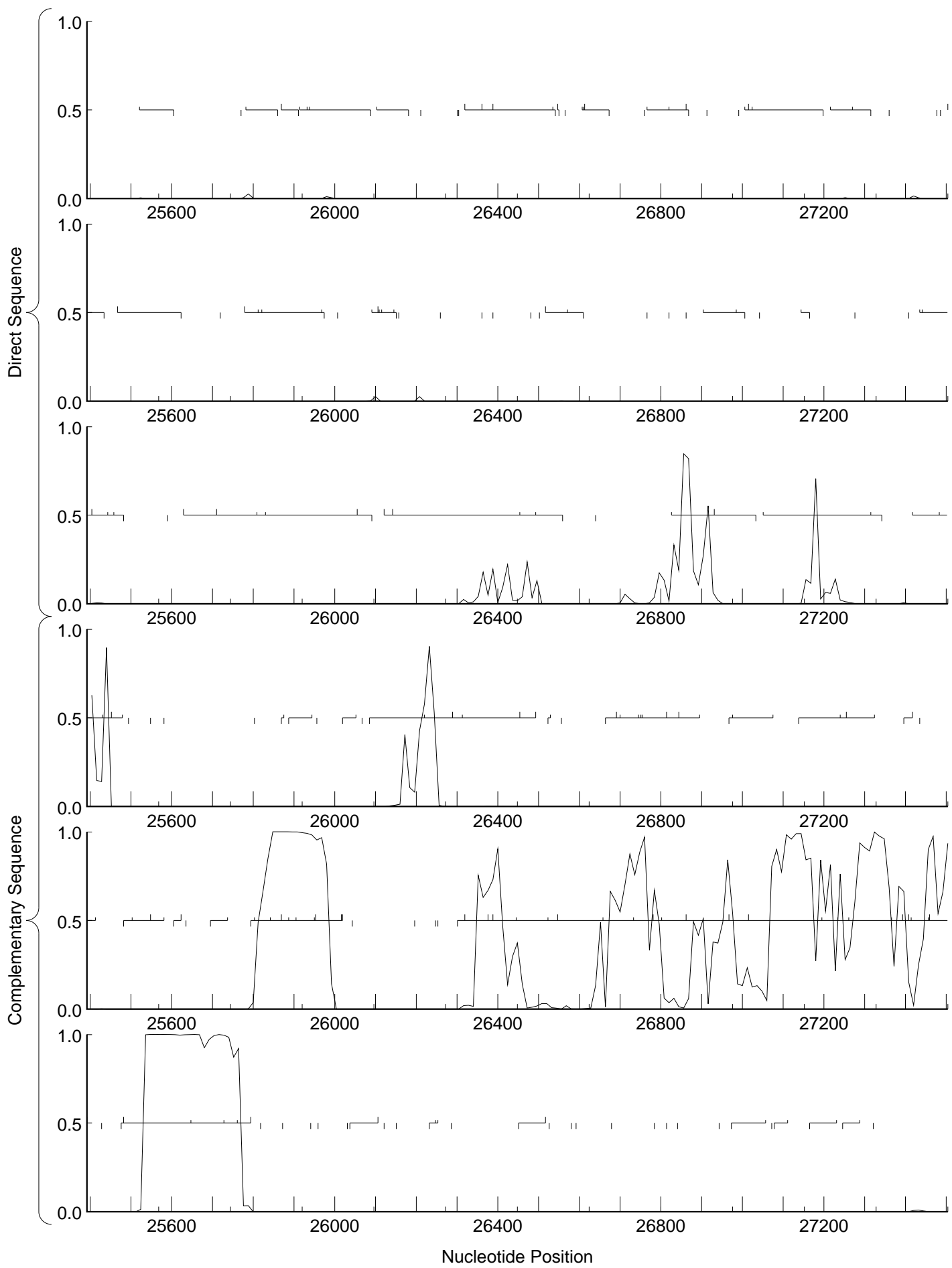


complete sequence, 46493 bp including 10-base 3' overhang (CGGCCGTGAT), Cluster CA, Order 4, Window 96, Step 12, 11/24

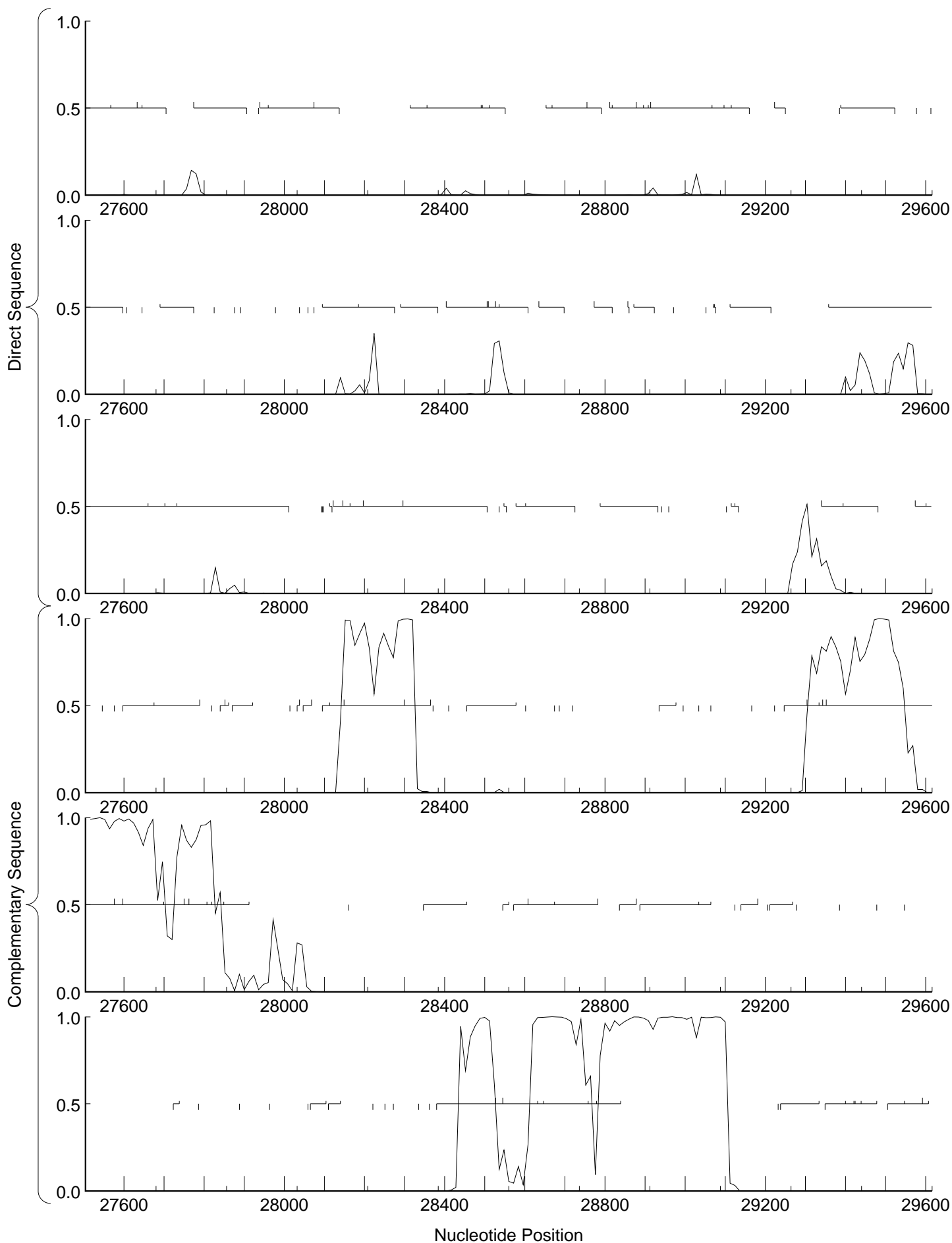




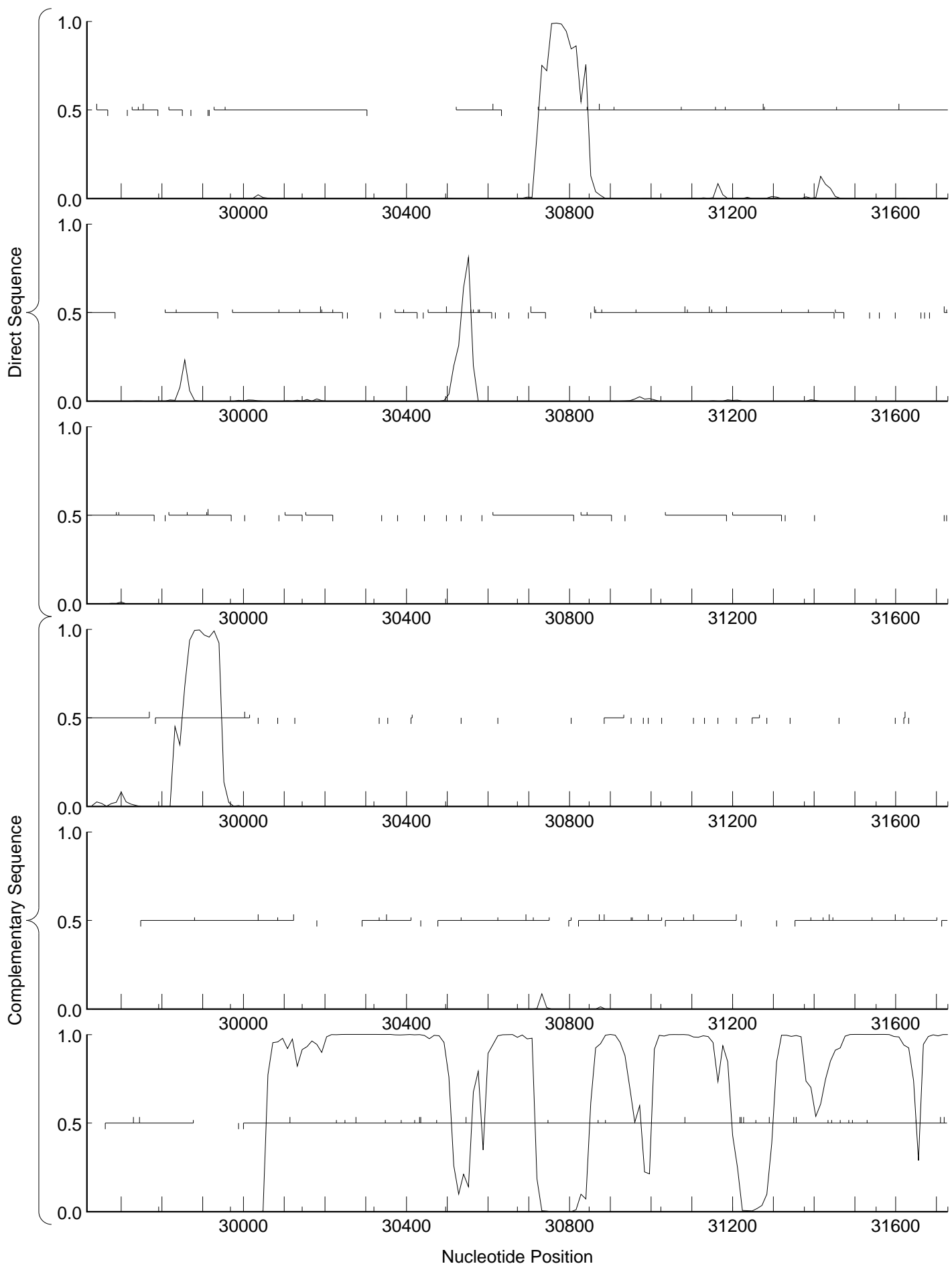




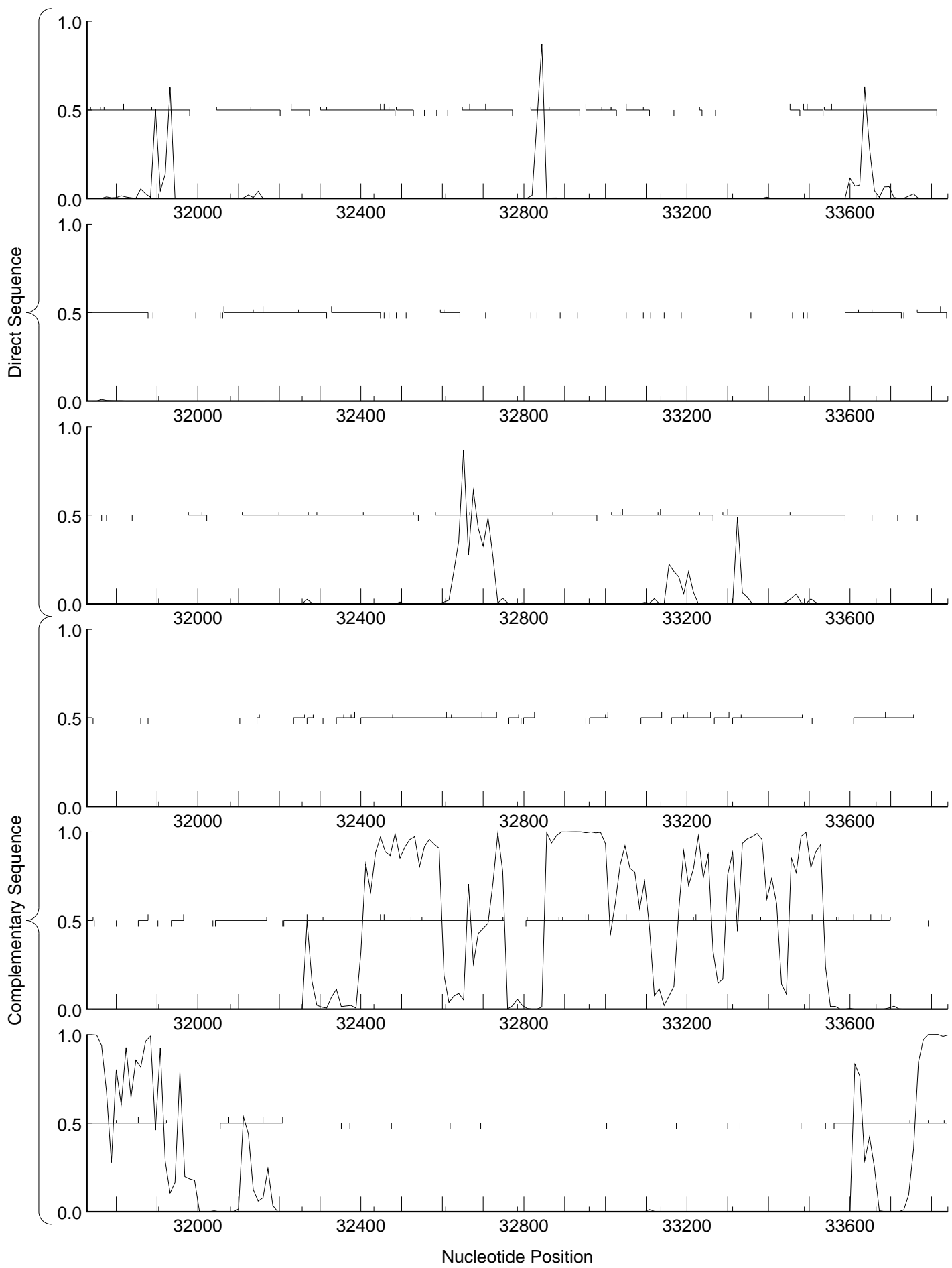
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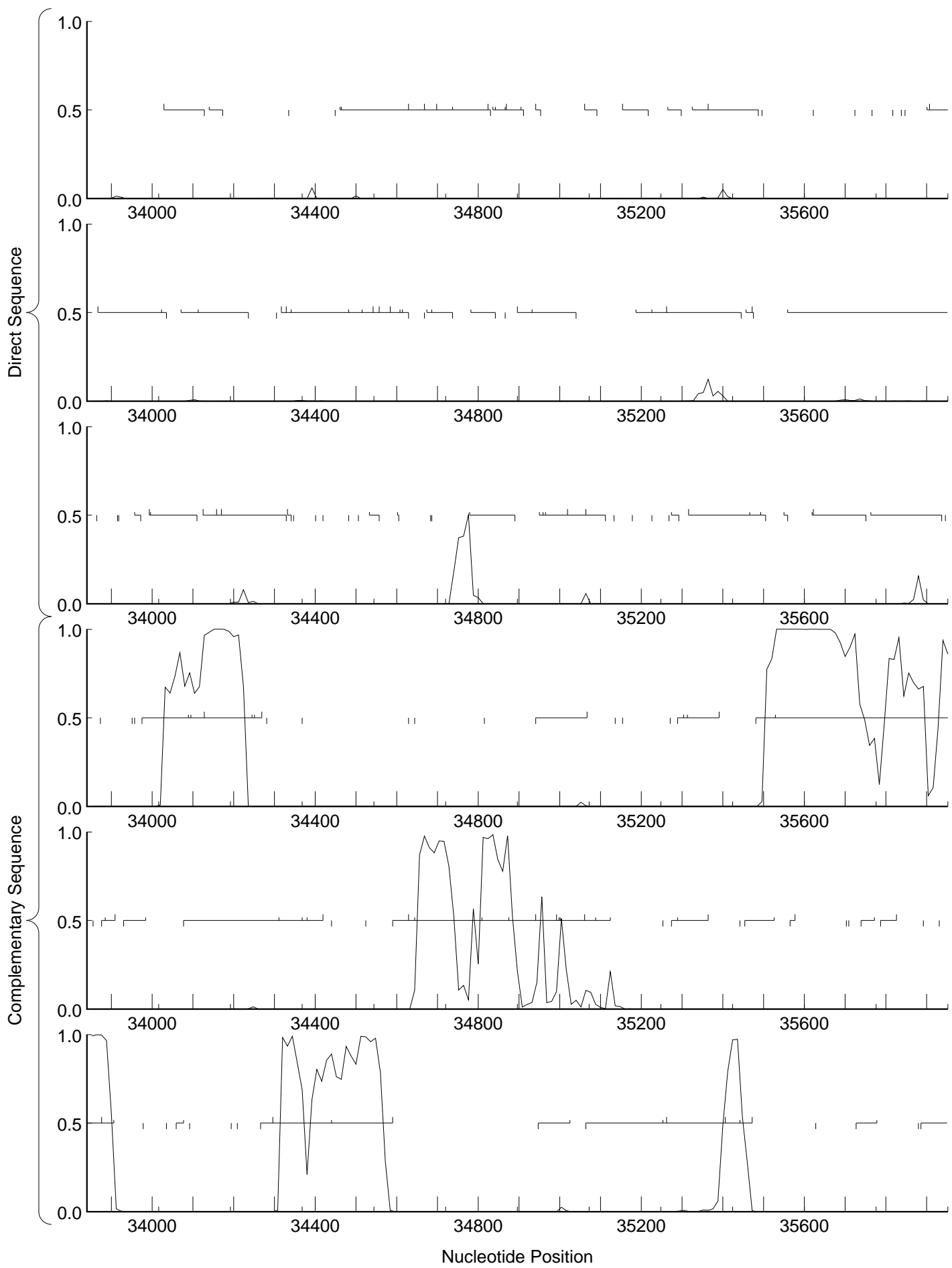
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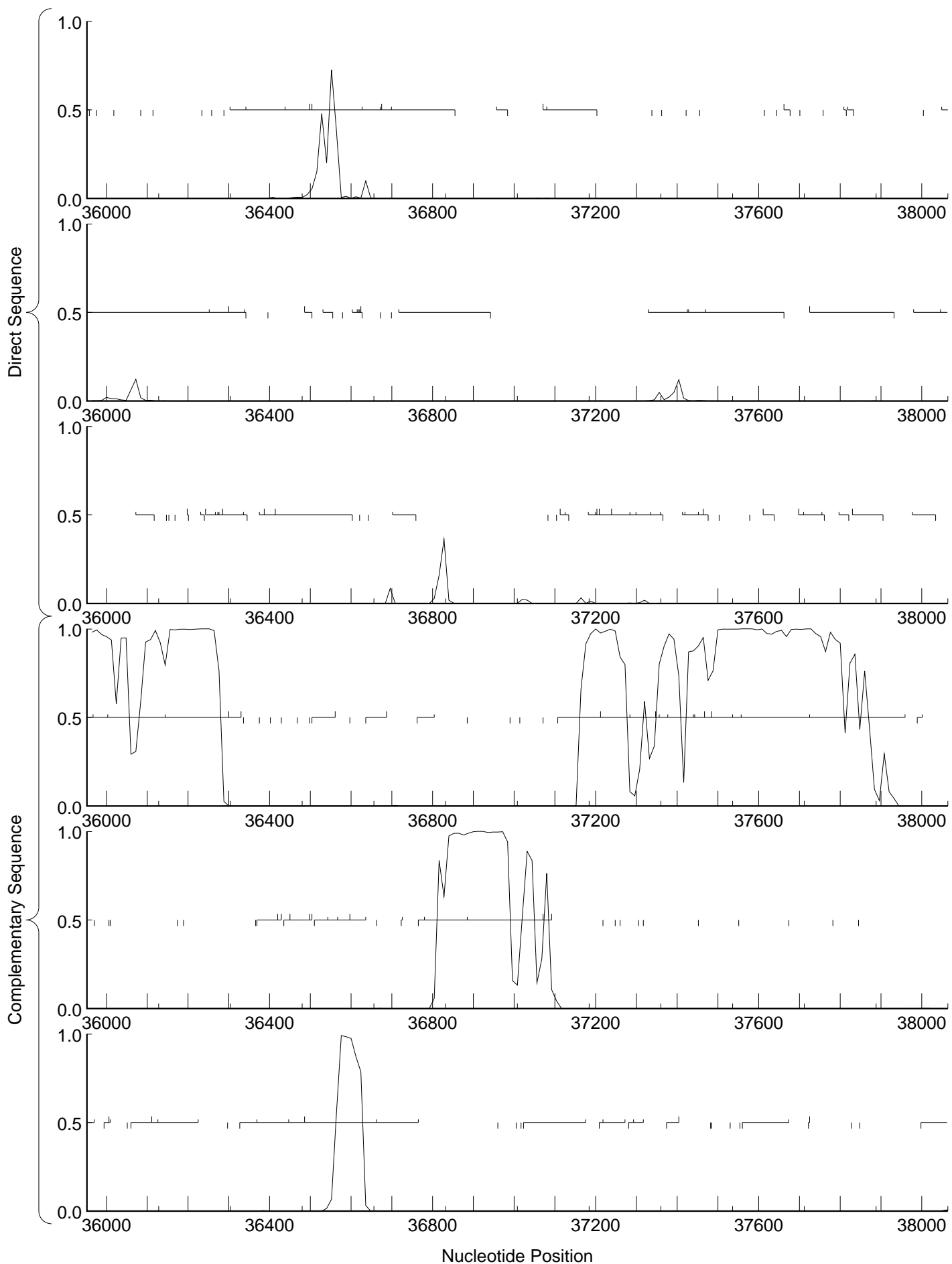


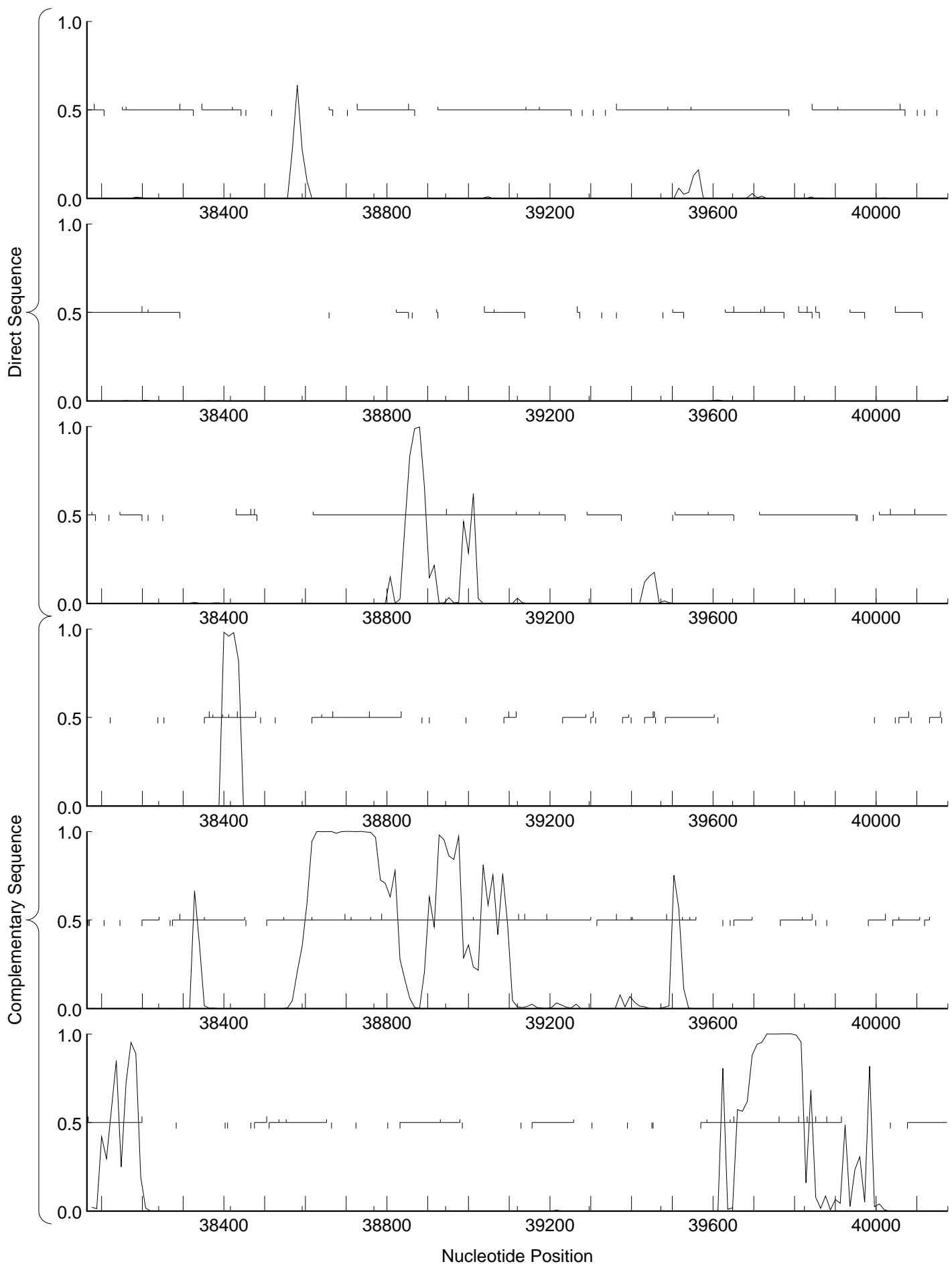
complete sequence, 46493 bp including 10-base 3' overhang (CGGCCGTGAT), Cluster CA, Order 4, Window 96, Step 12, 17/24

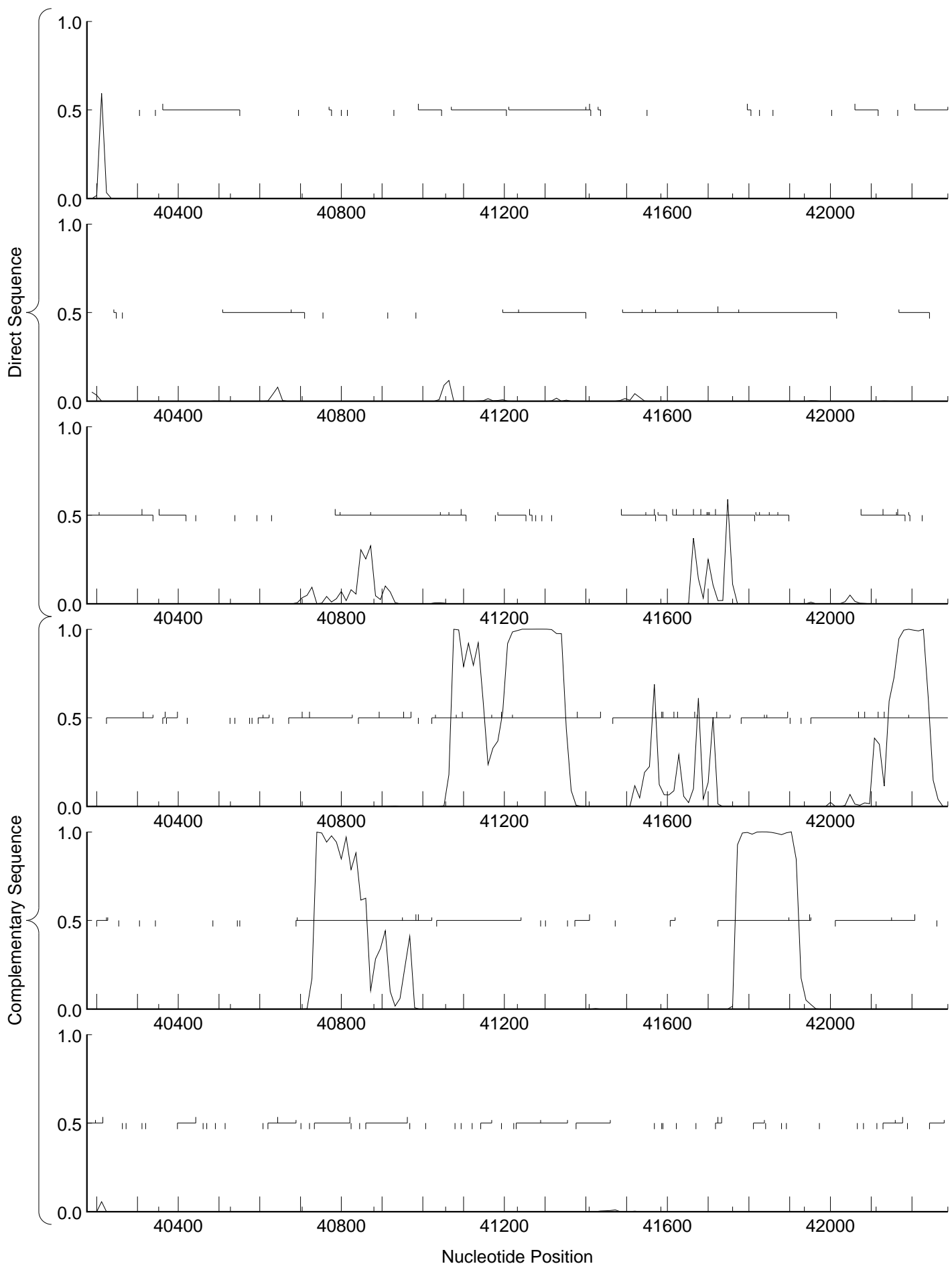


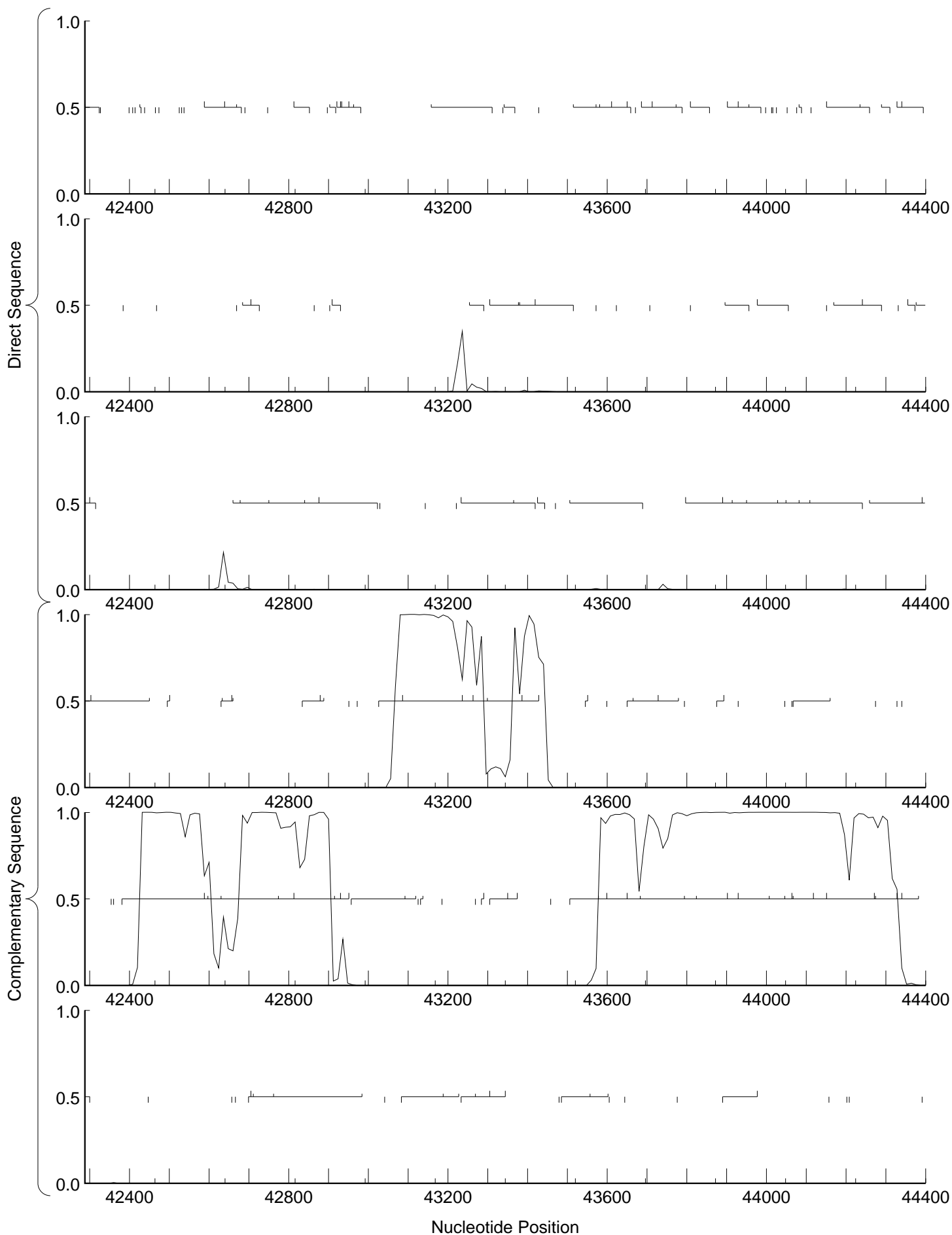
complete sequence, 46493 bp including 10-base 3' overhang (CGGCCGTGAT), Cluster CA, Order 4, Window 96, Step 12, 18/24











complete sequence, 46493 bp including 10-base 3' overhang (CGGCCGTGAT), Cluster CA, Order 4, Window 96, Step 12, 23/24

