

GeneMark

Version 2.5p (09.08.06)

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PROGRAM INFORMATION

Sequence : Mycobacterium phage MilanaBonita complete sequence, 56251 bp including 10-base 3' overhang
Analysis Date : 2/27/21 at 17:29:20
Pages : 28
Sequence Length : 56251 bp
GC Content : 62.06%

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,

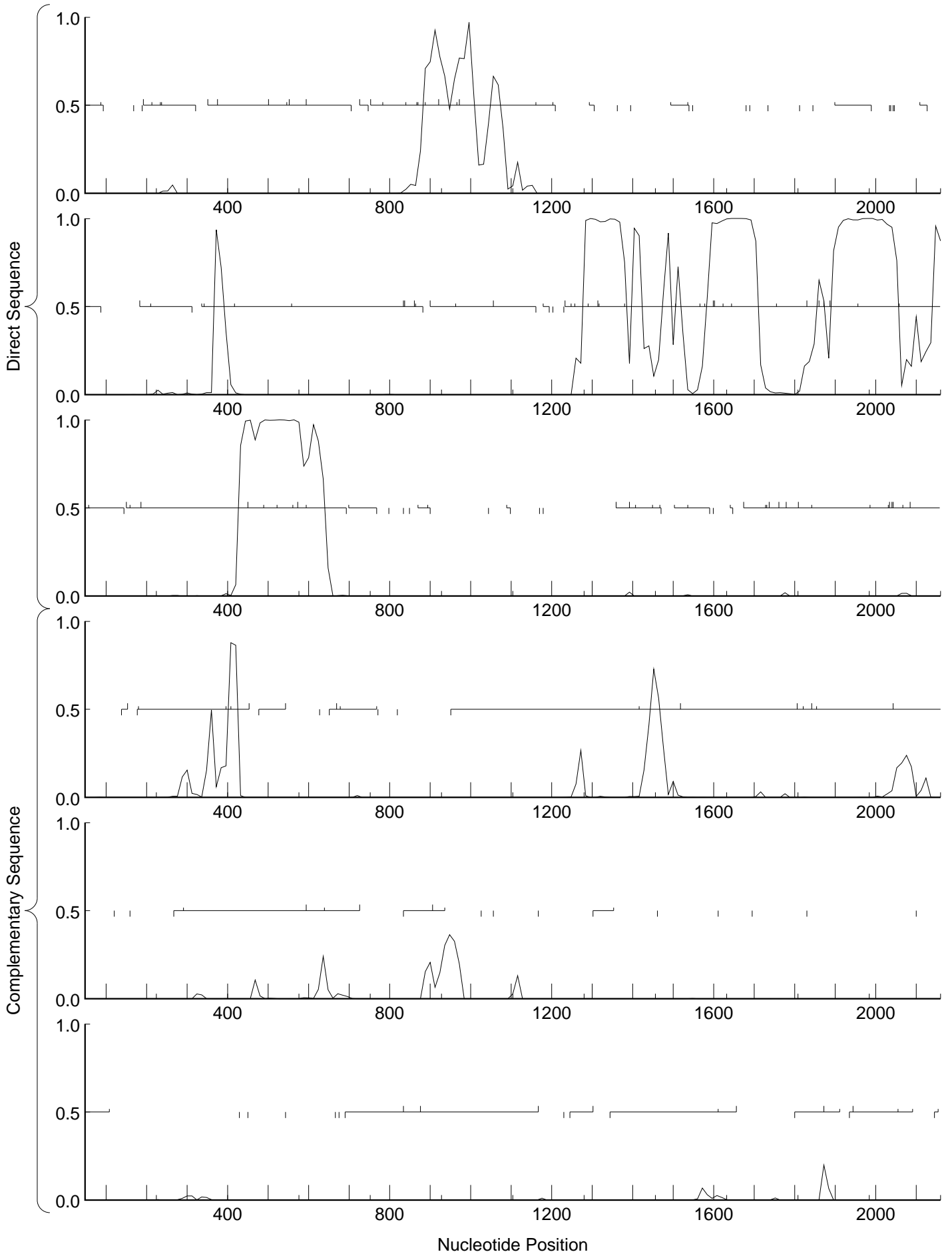
Matrix notes & comments

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

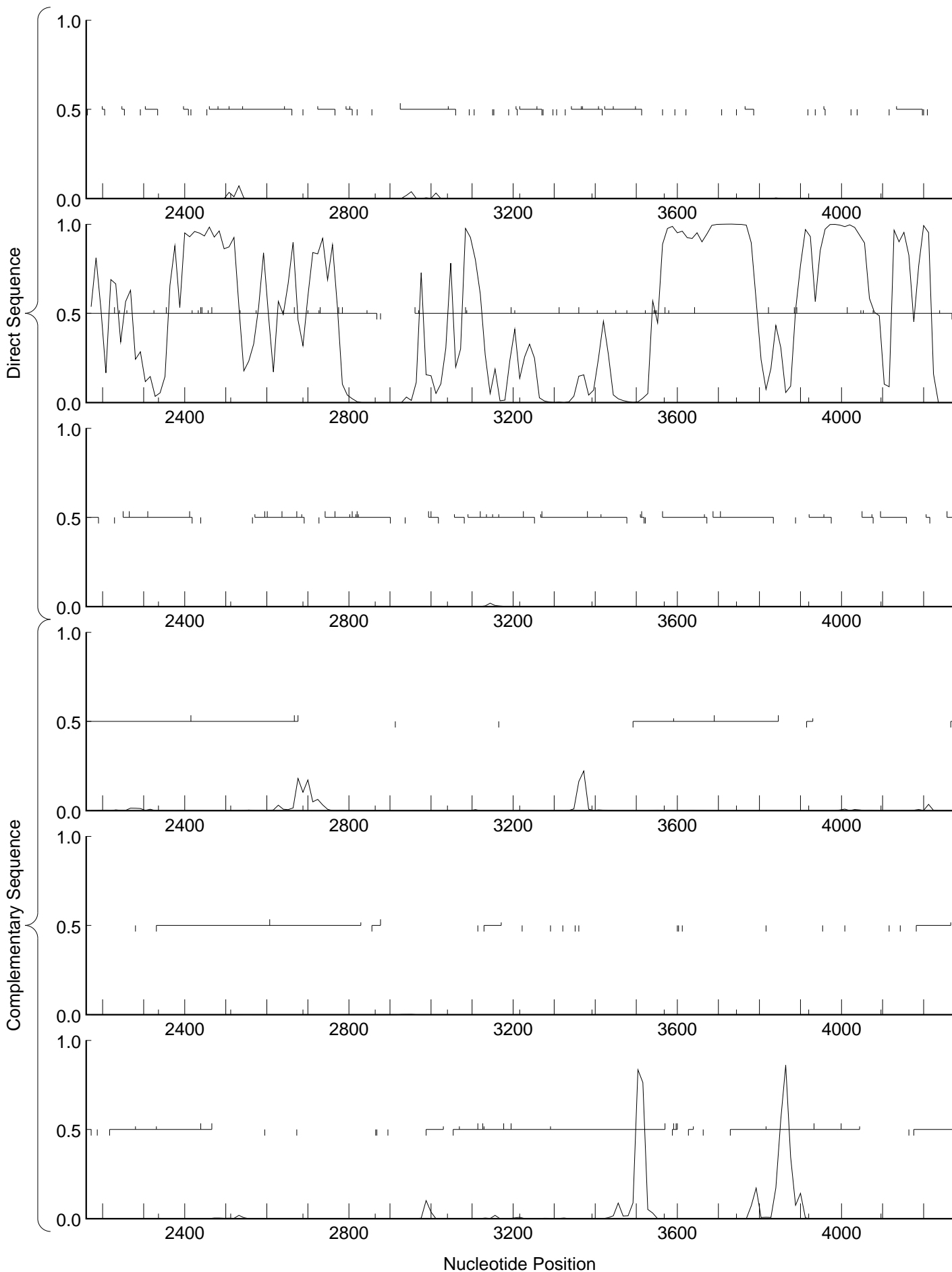
MATRIX INFORMATION

Matrix : Mycobacterium_tuberculosis_H37Rv
Author : -
Order : 4

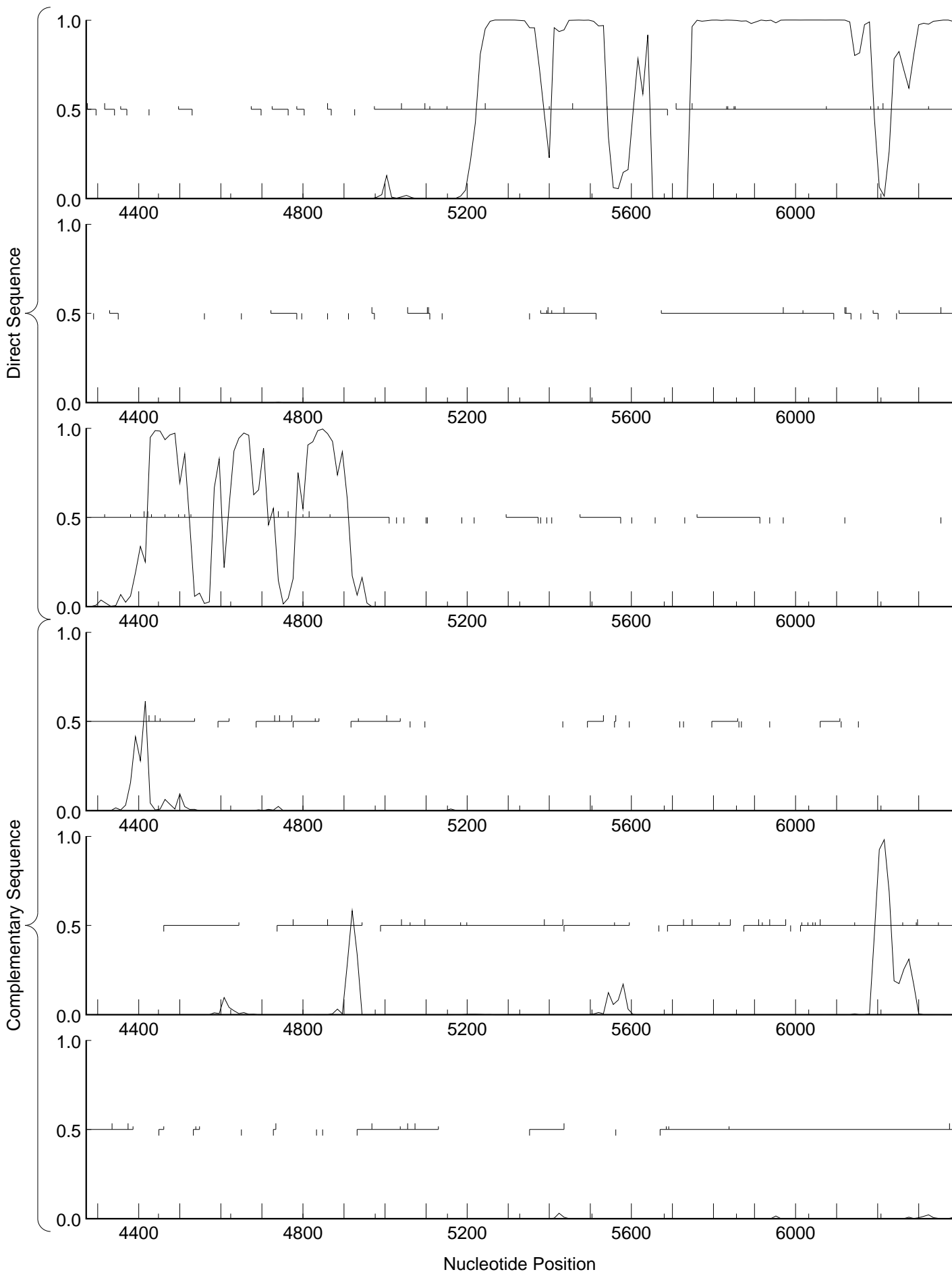
Send questions / comments to:
Dr. M. Borodovsky
Georgia Institute of Technology
School of Biology
Atlanta, GA 30332-0230

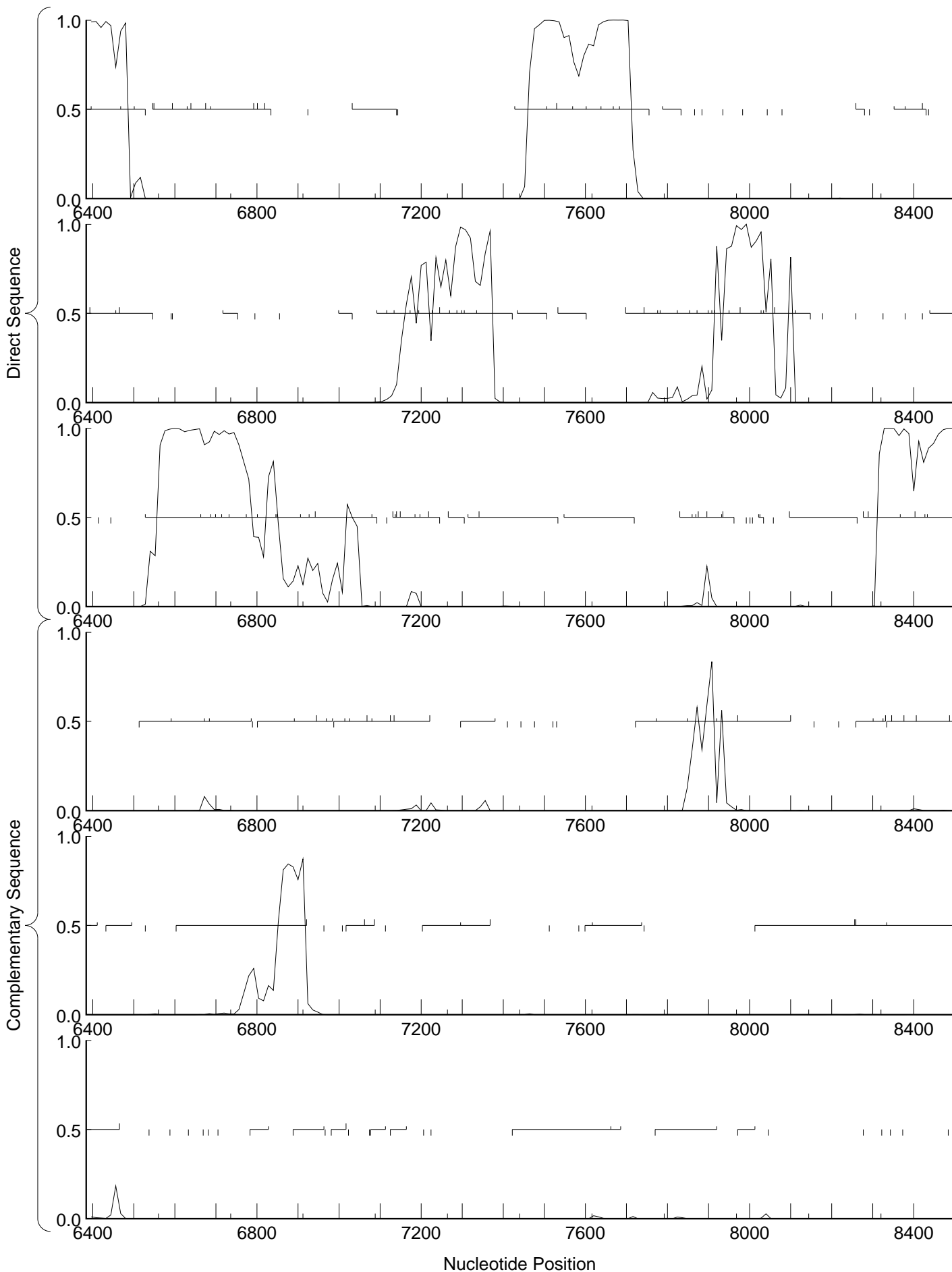


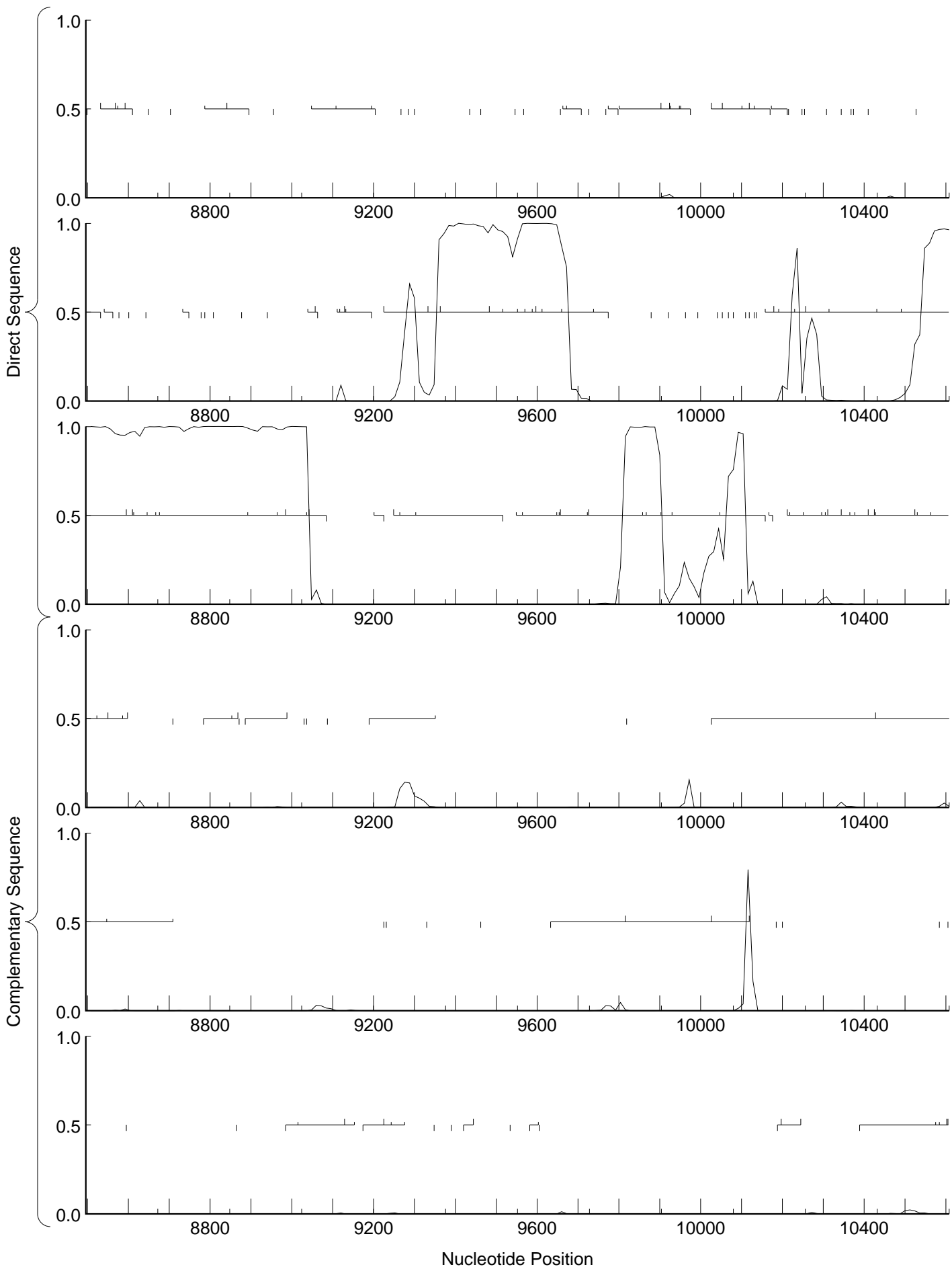
a complete sequence, 56251 bp including 10-base 3' overhang (CGGAAGGCGC), Cluster F1, Order 4, Window 96, Step 12, 3/28



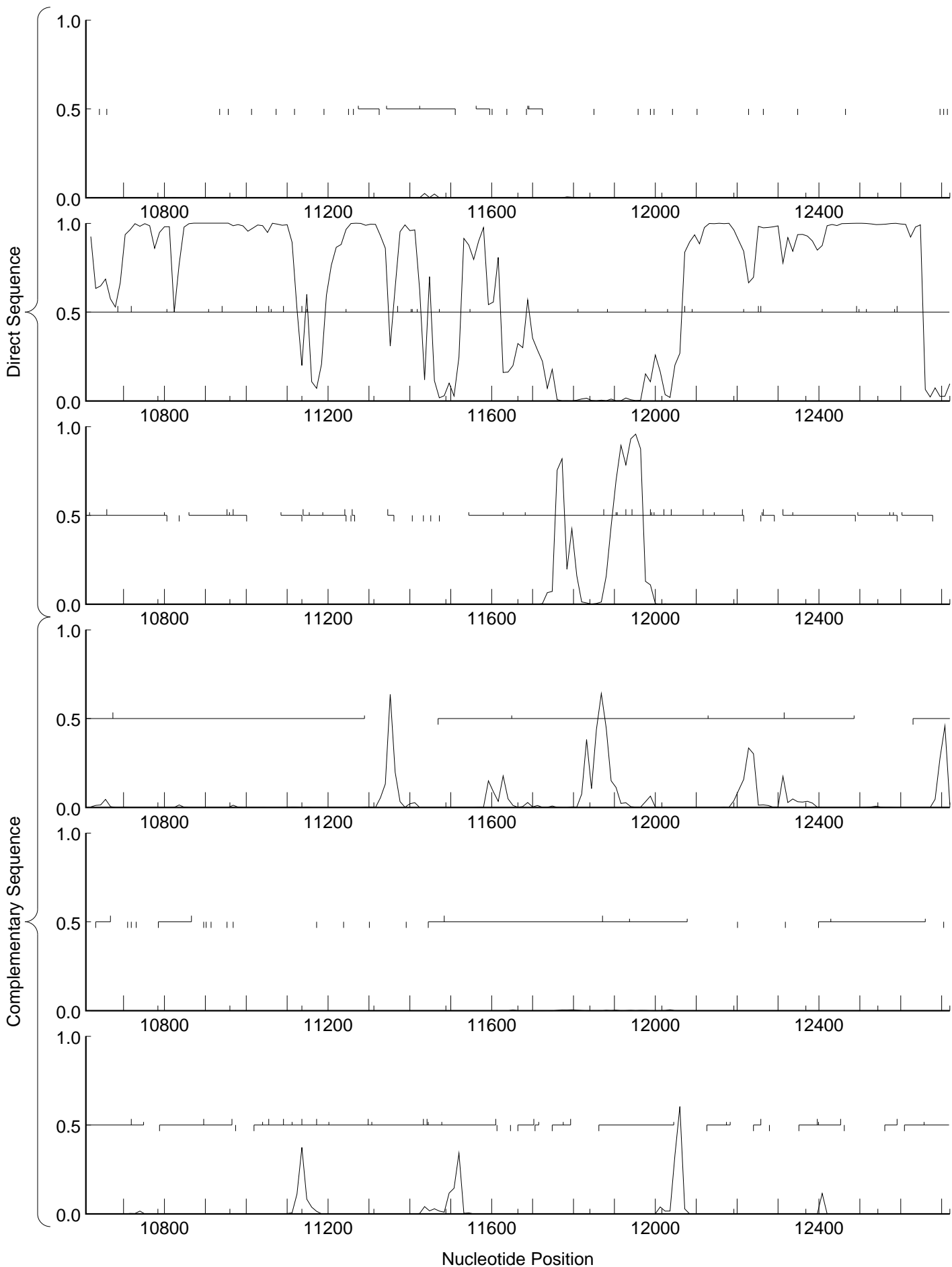
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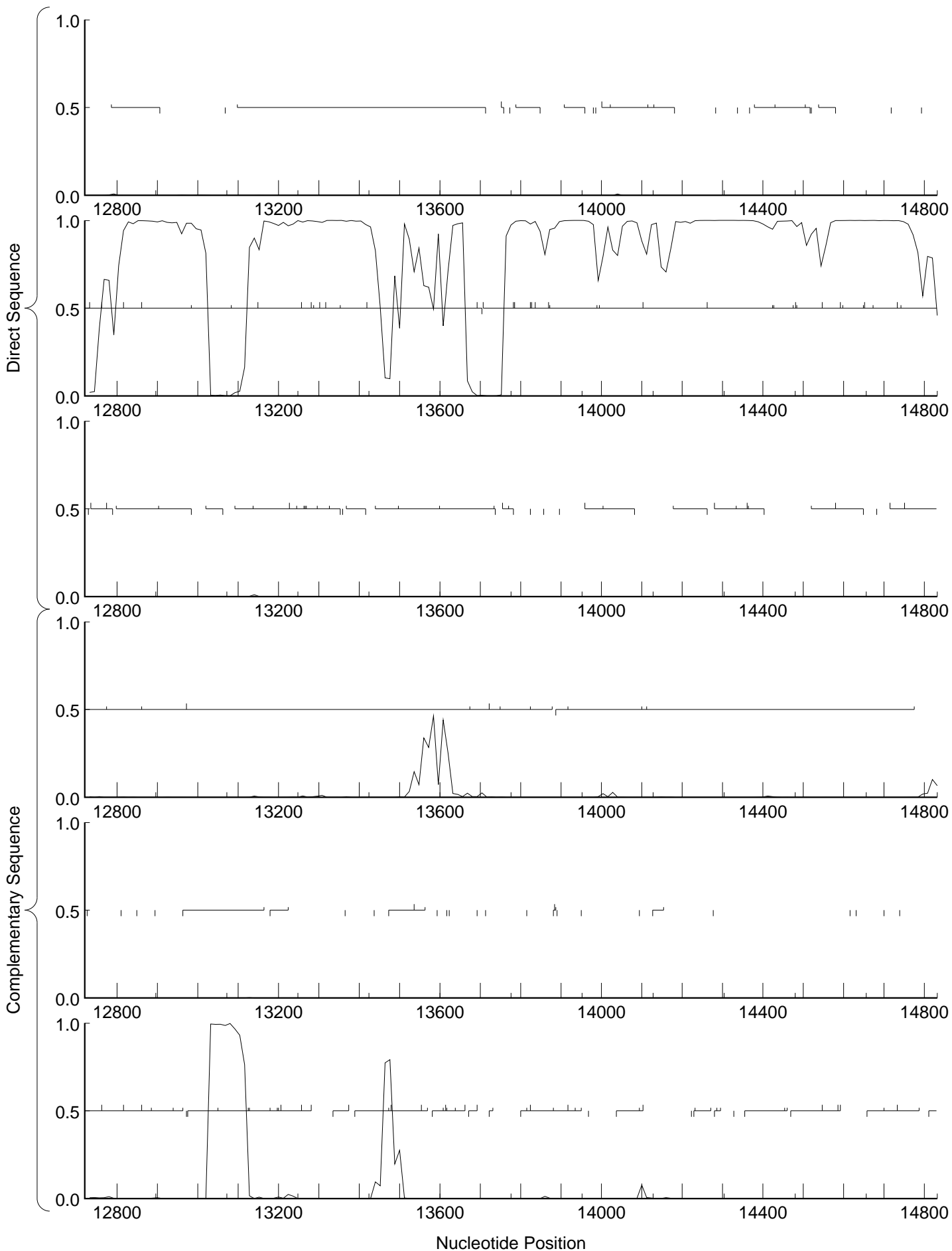


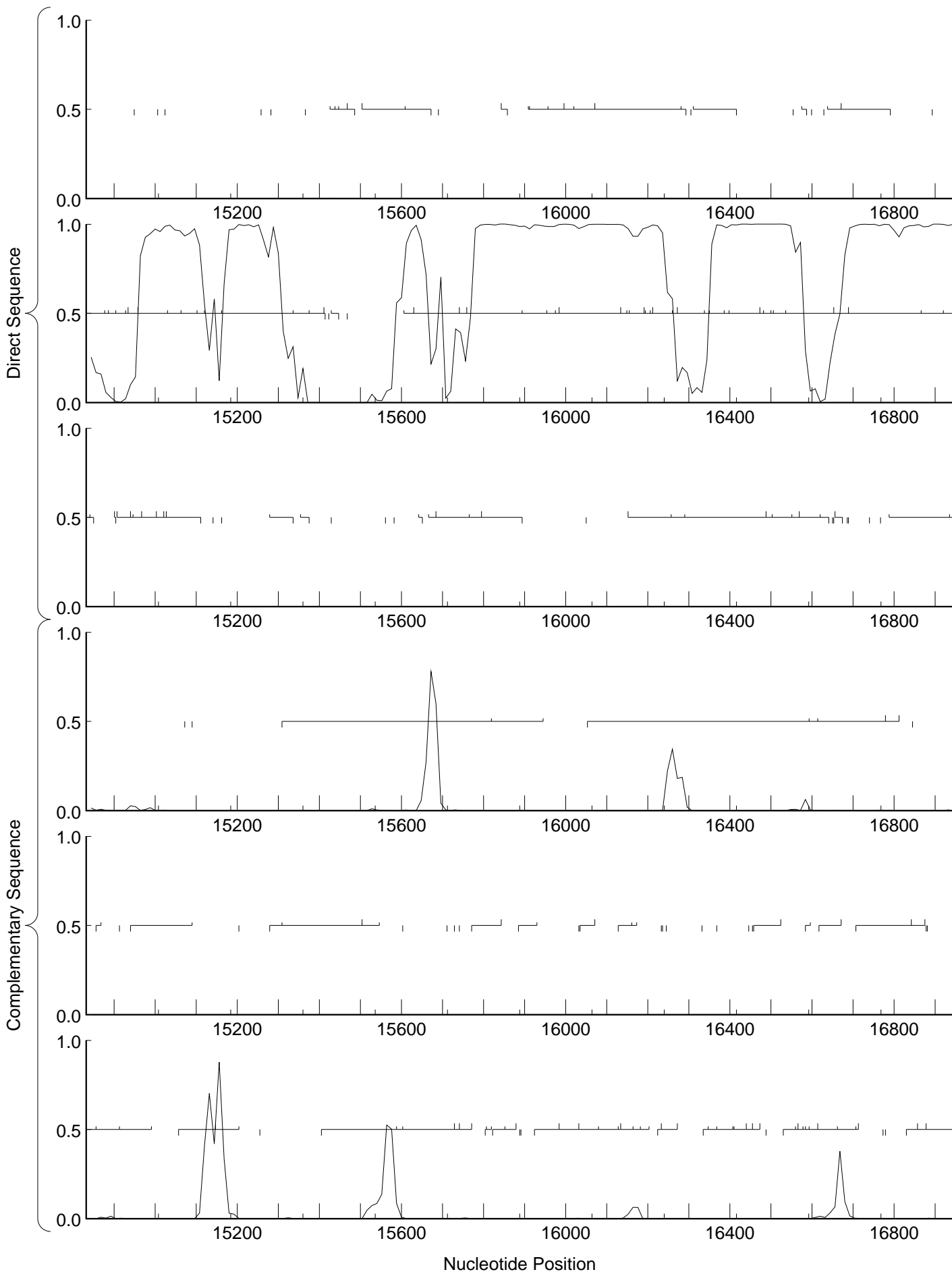




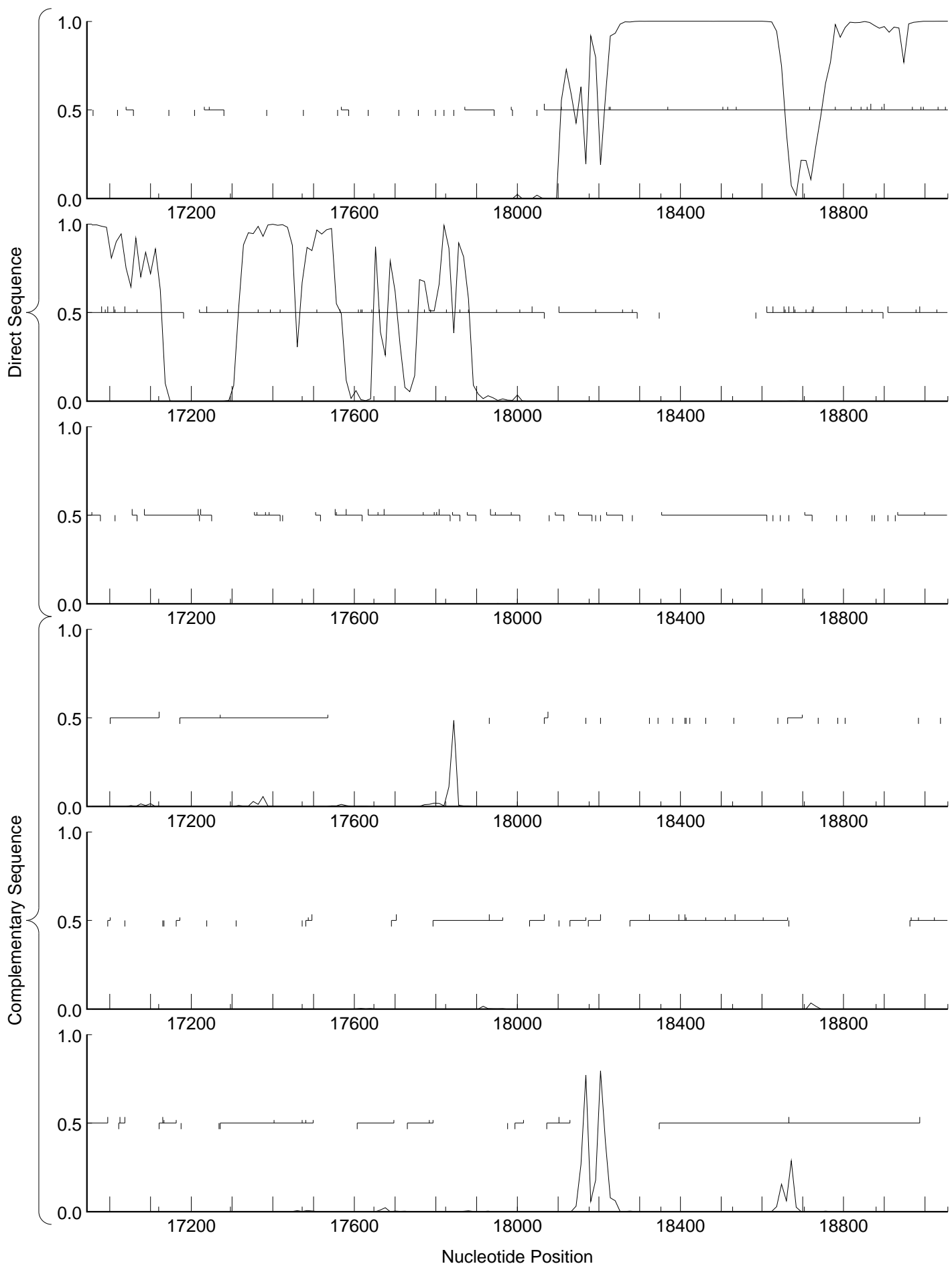
a complete sequence, 56251 bp including 10-base 3' overhang (CGGAAGGCGC), Cluster F1, Order 4, Window 96, Step 12, 7/28



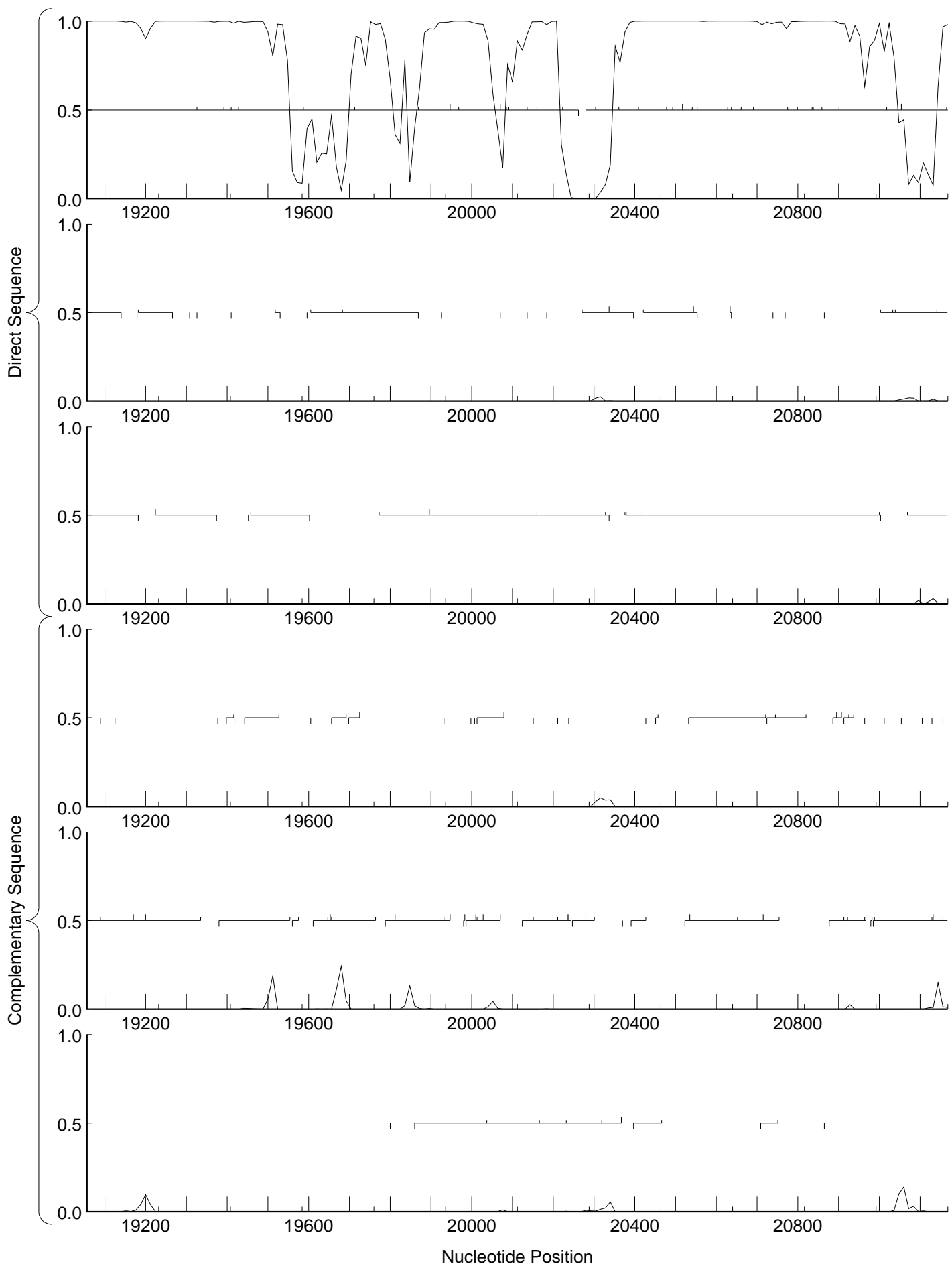


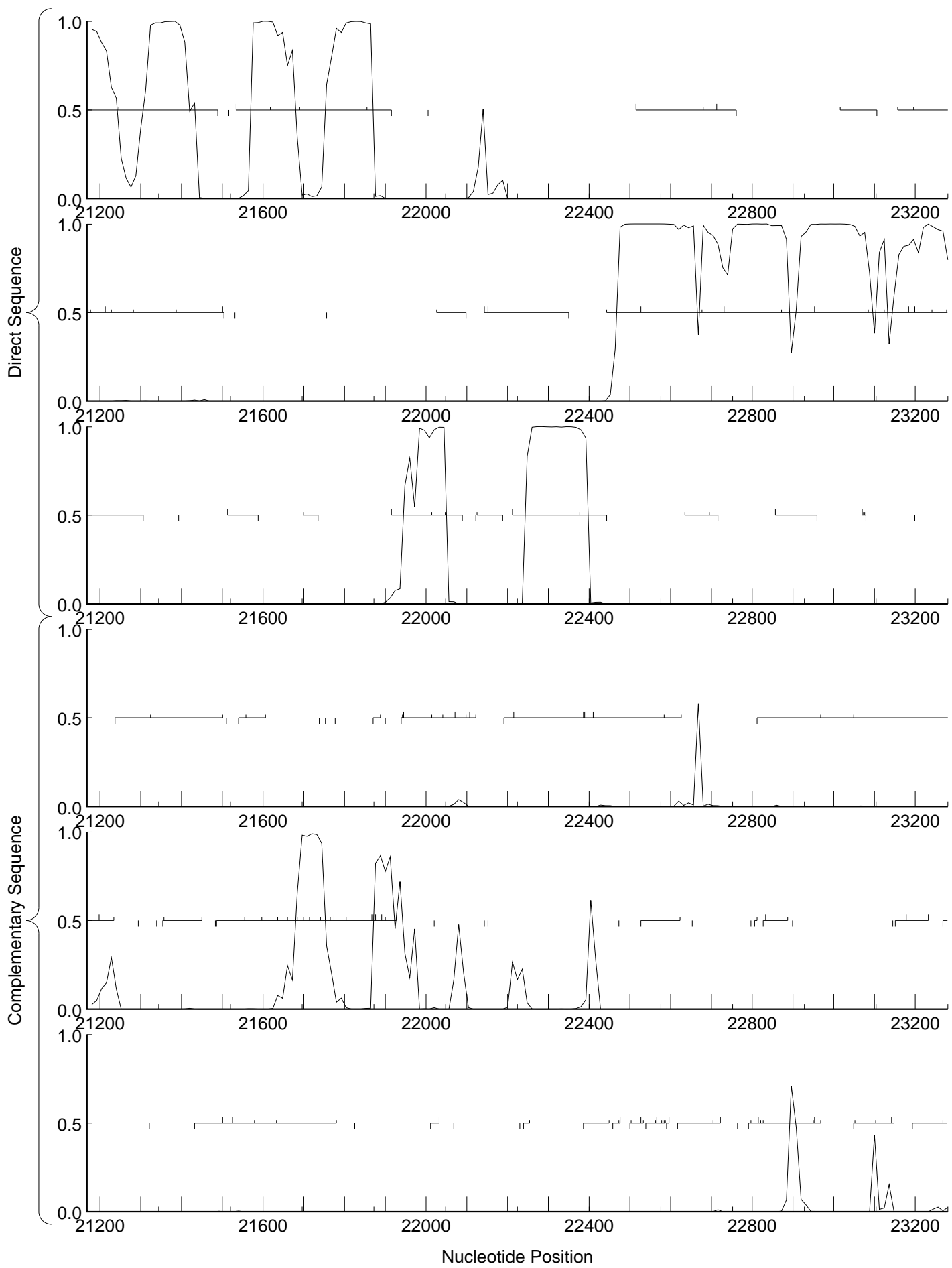


complete sequence, 56251 bp including 10-base 3' overhang (CGGAAGGCGC), Cluster F1, Order 4, Window 96, Step 12, 10/28

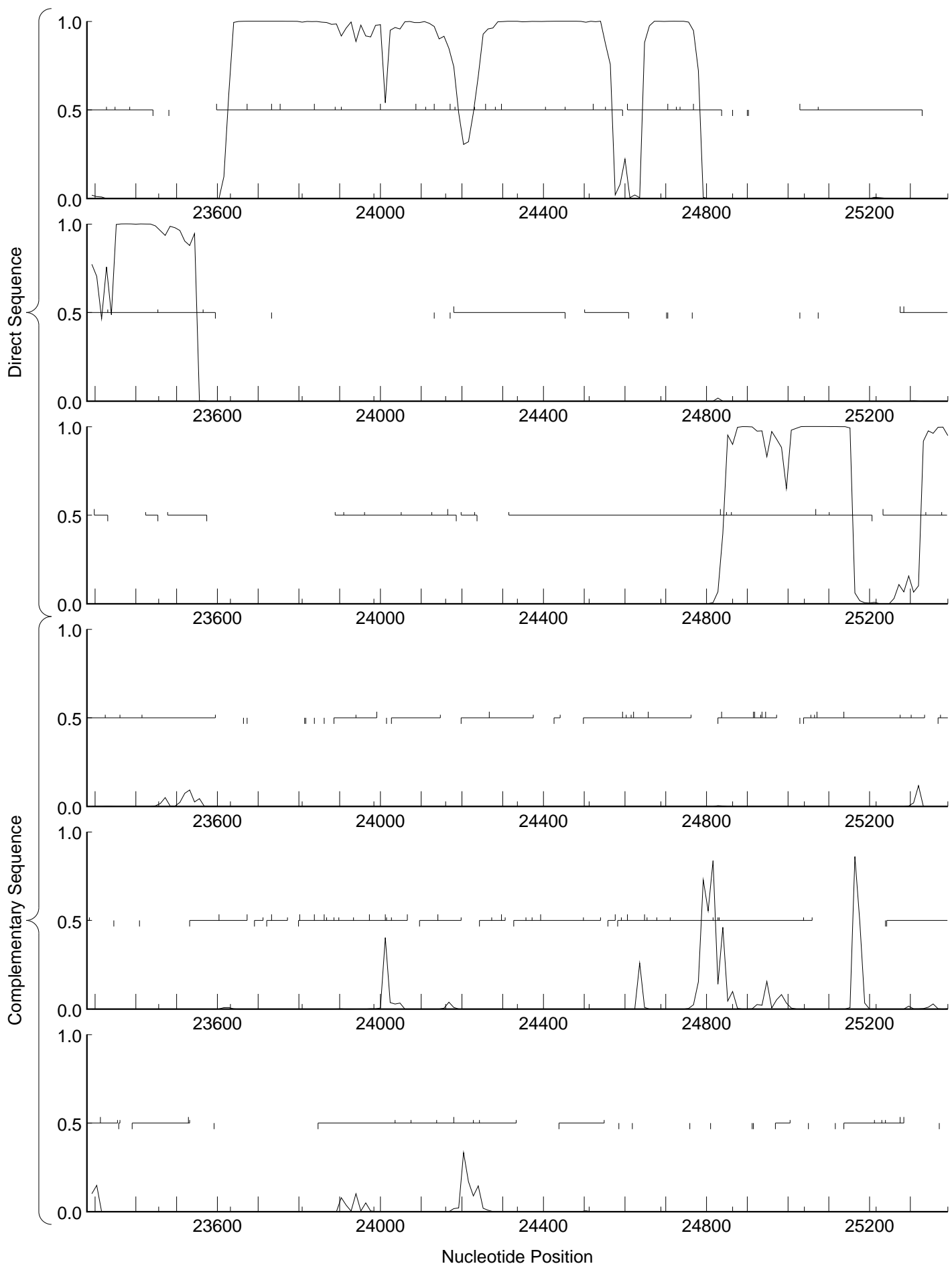


complete sequence, 56251 bp including 10-base 3' overhang (CGGAAGGCGC), Cluster F1, Order 4, Window 96, Step 12, 11/28

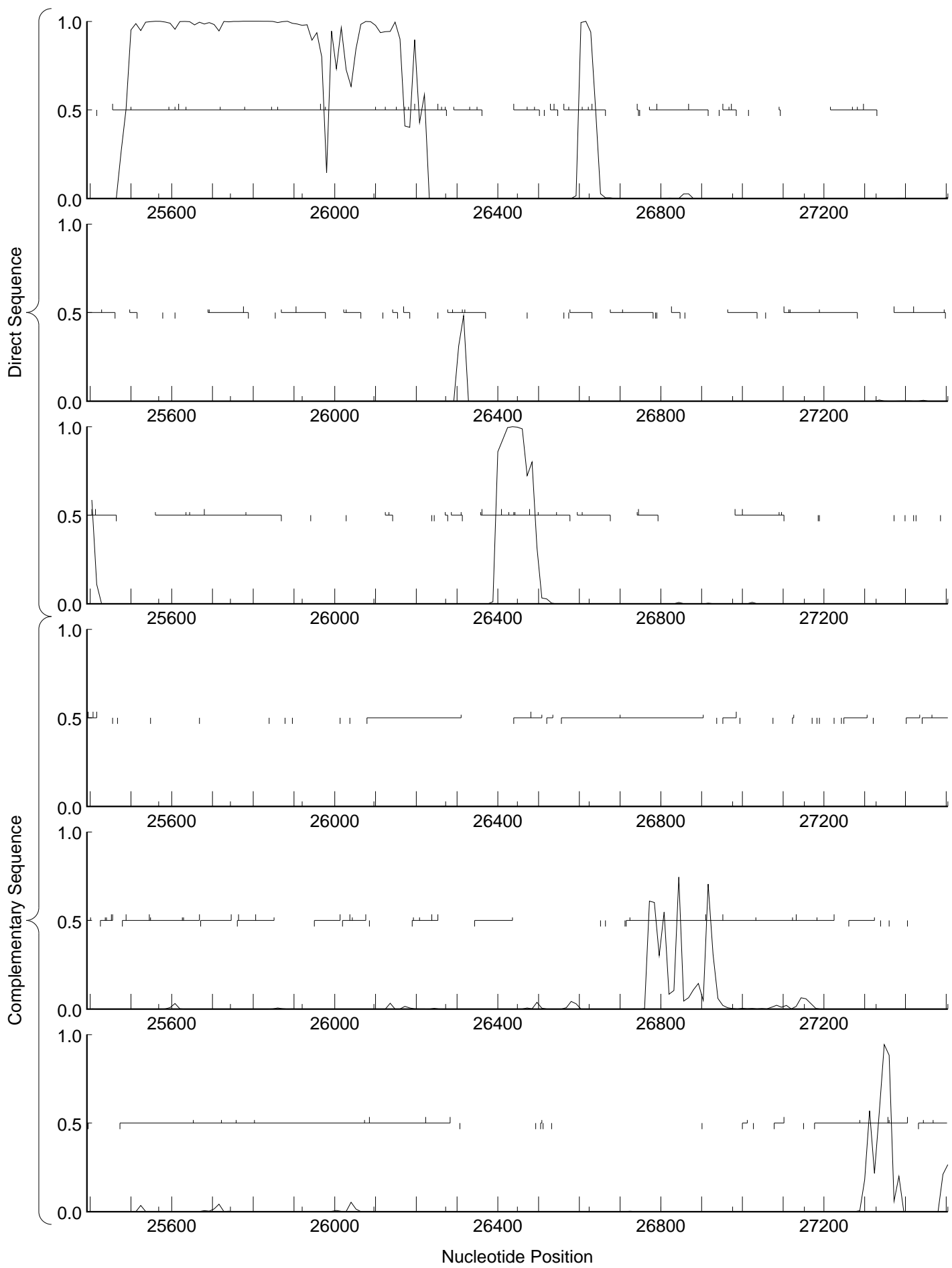




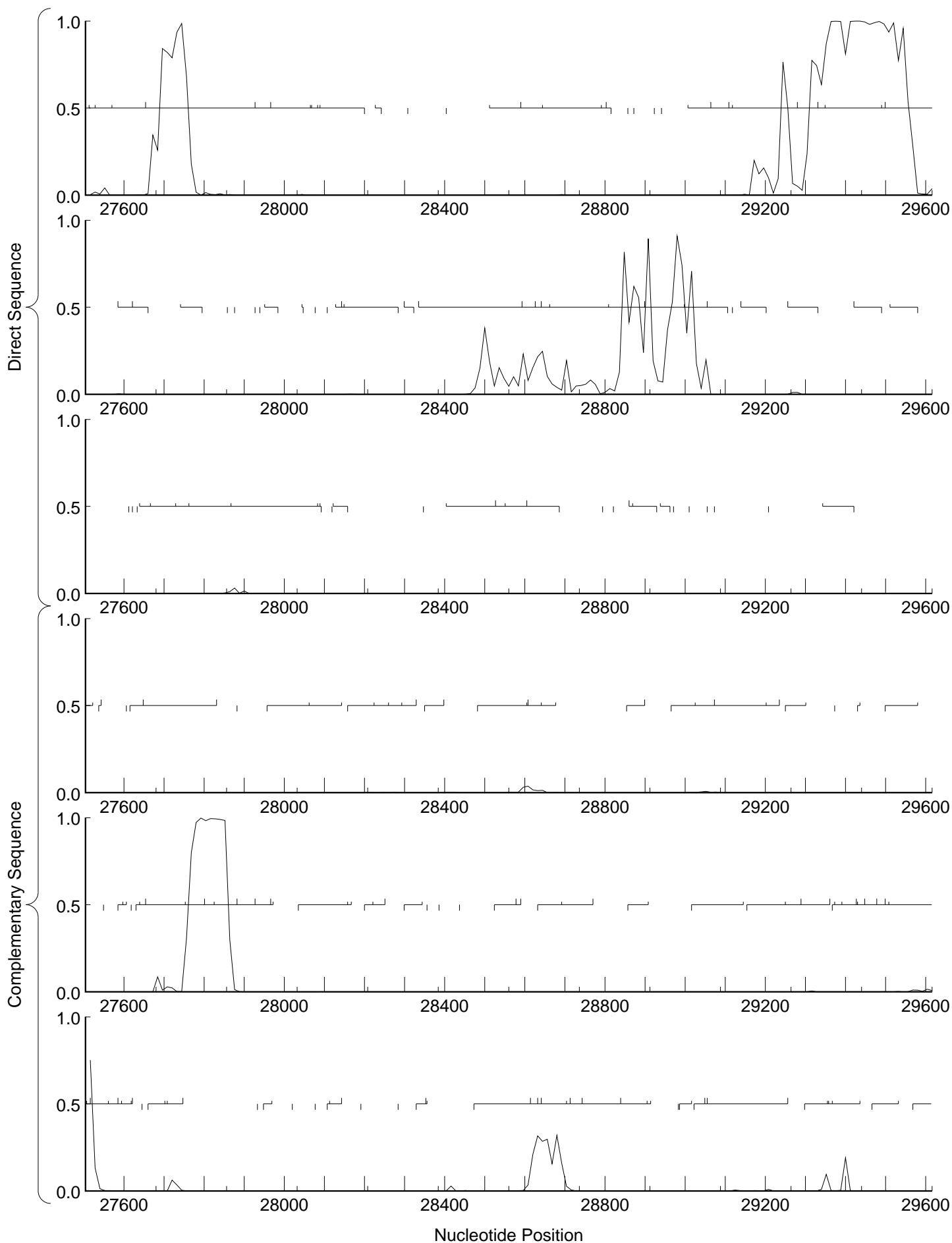
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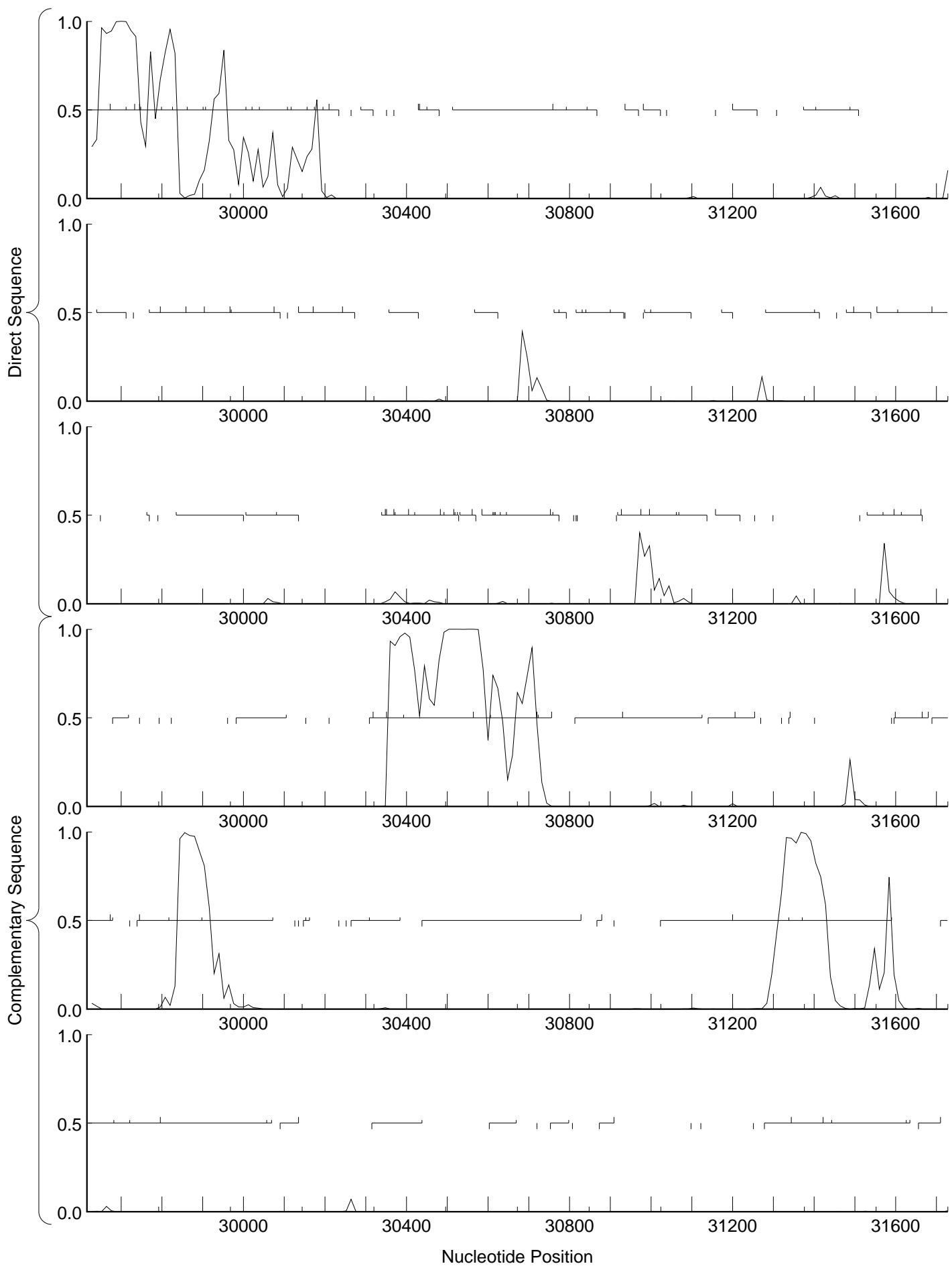


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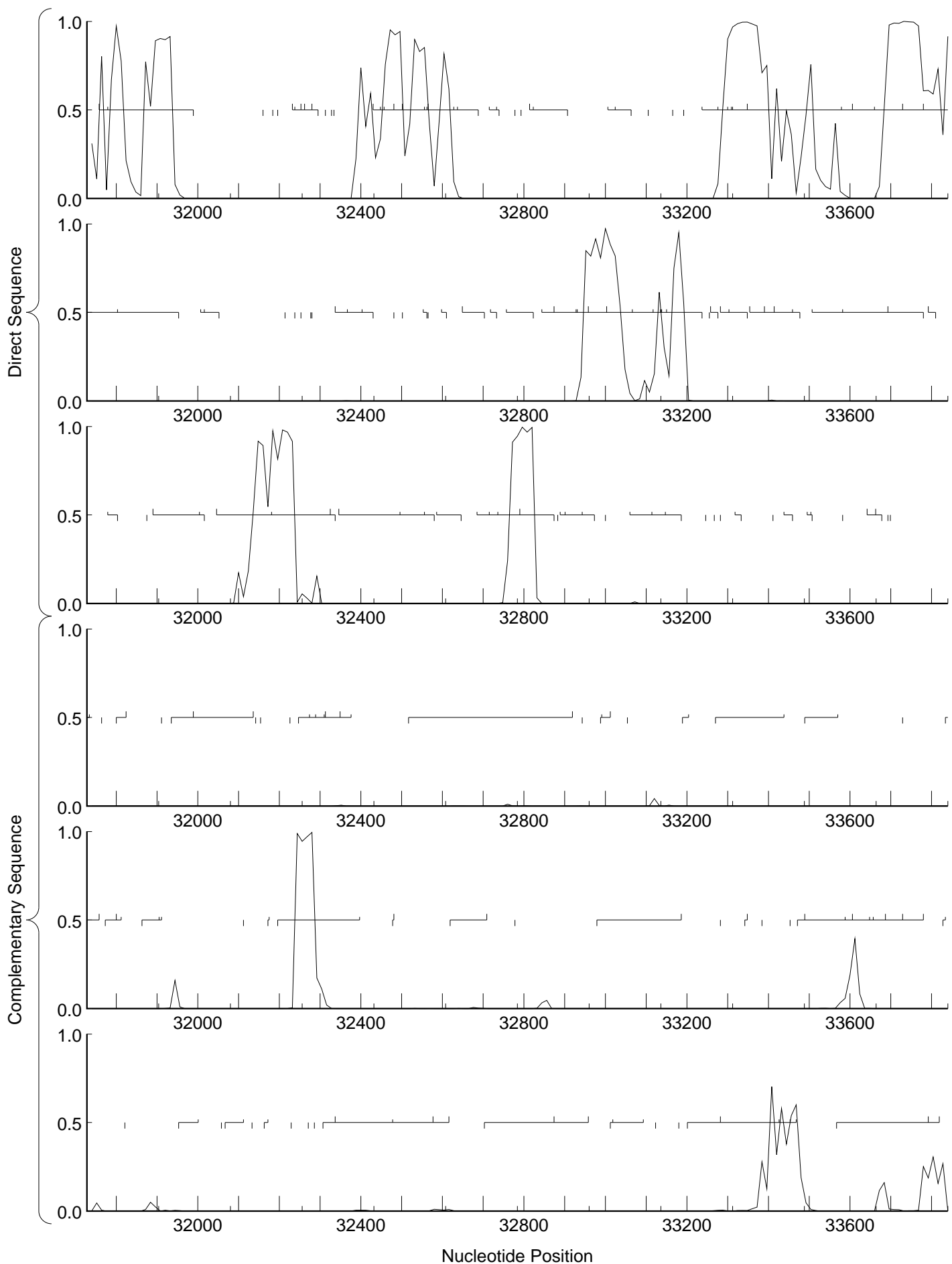


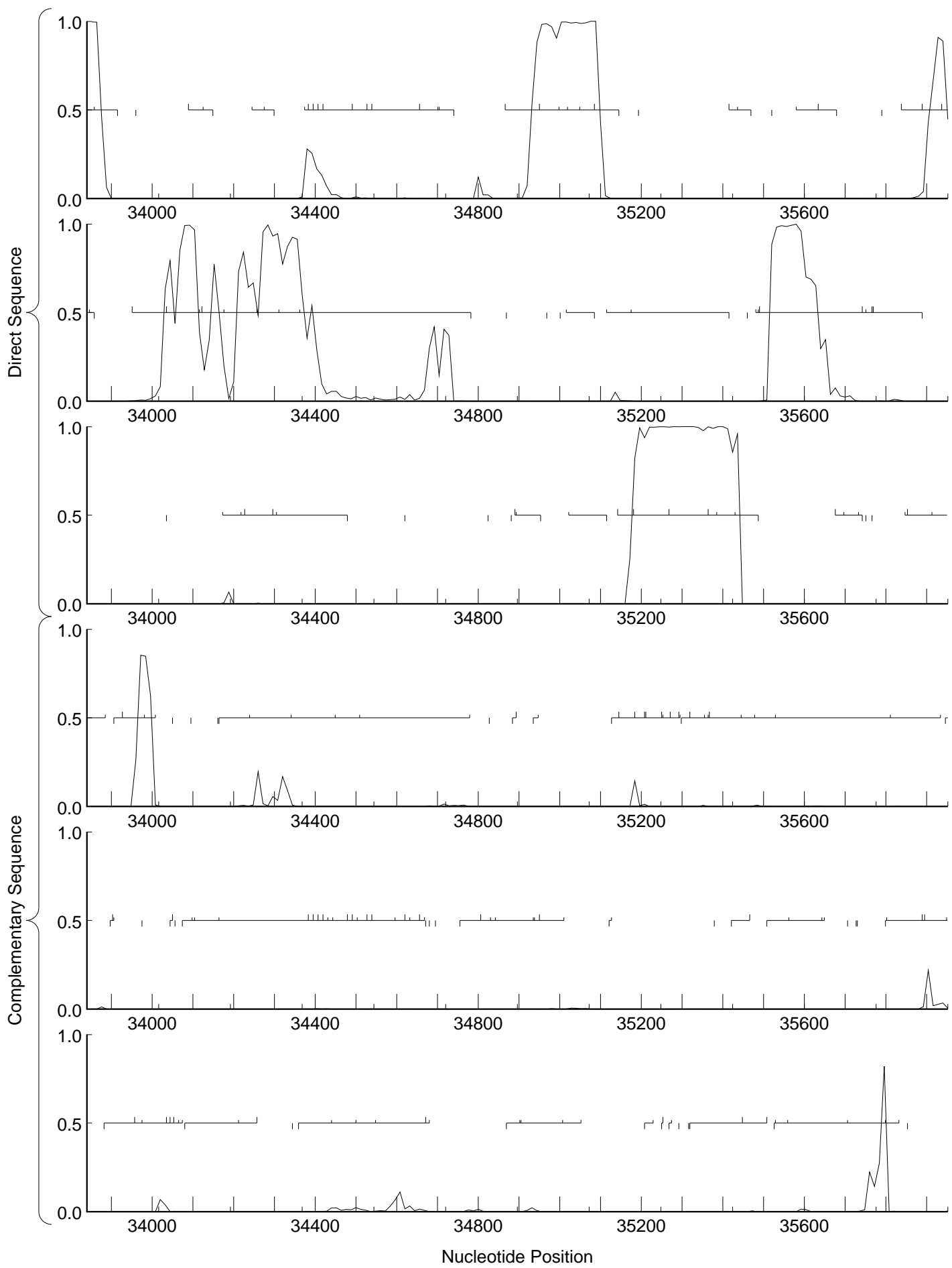
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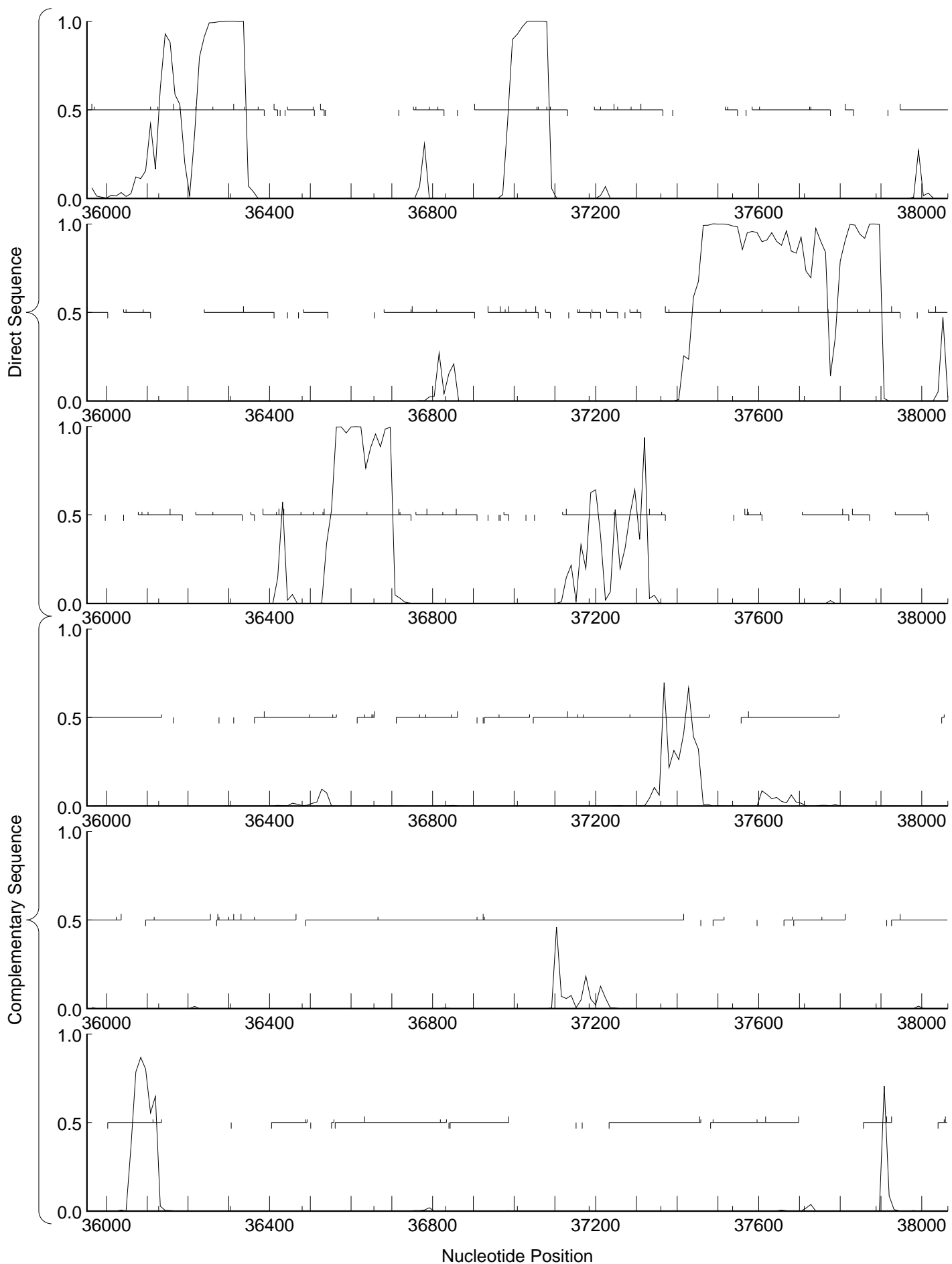


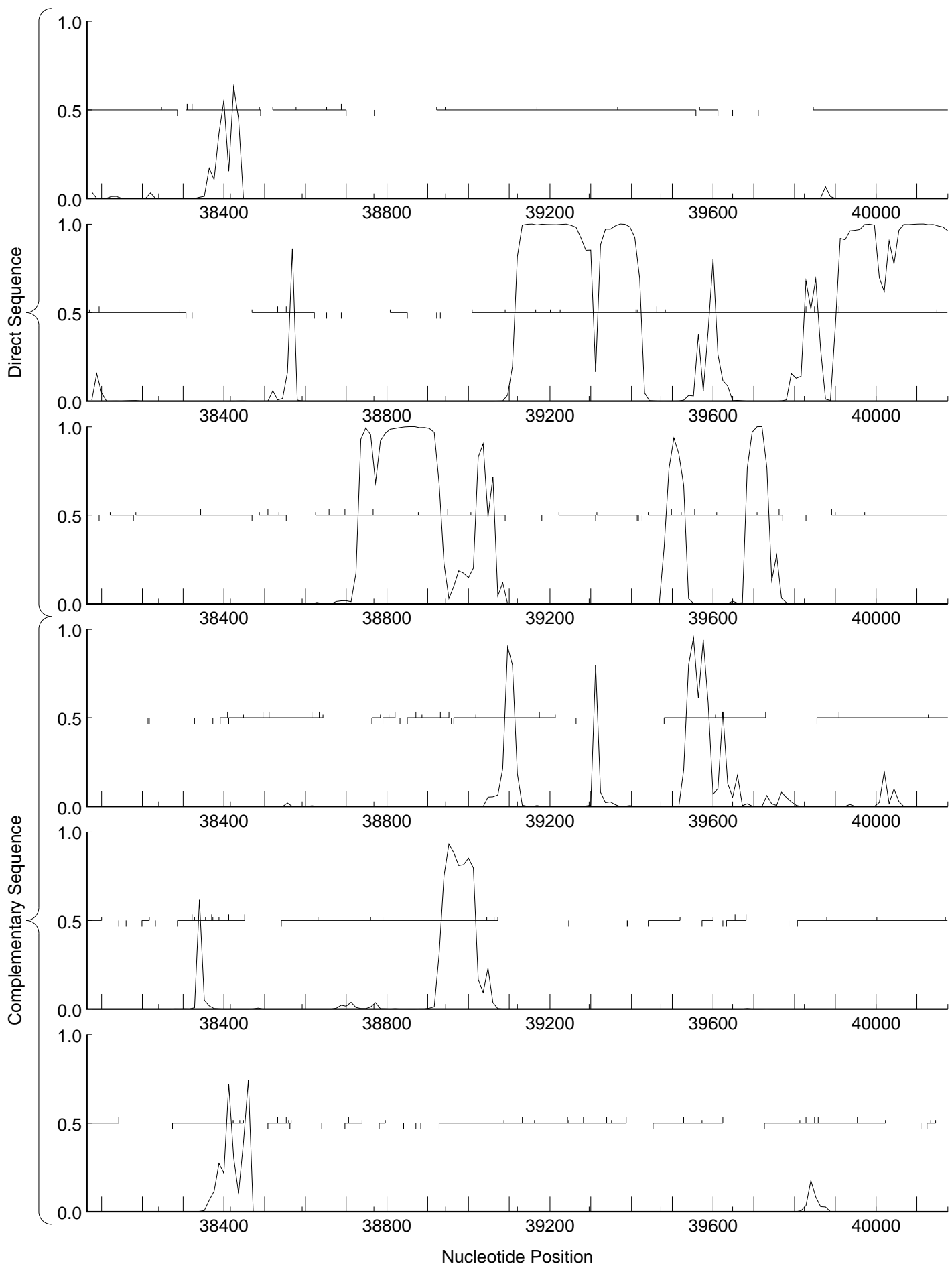
complete sequence, 56251 bp including 10-base 3' overhang (CGGAAGGCGC), Cluster F1, Order 4, Window 96, Step 12, 17/28

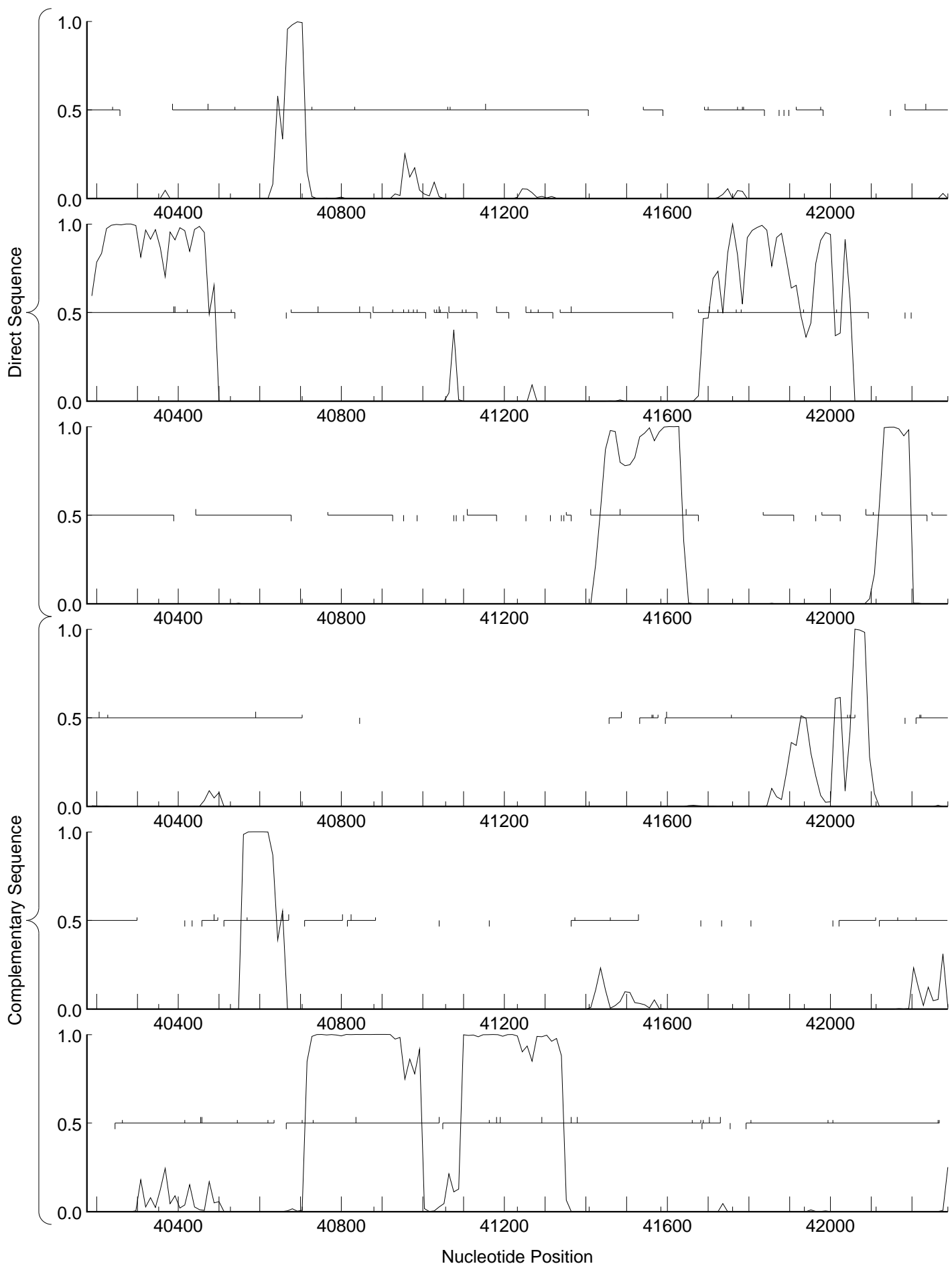


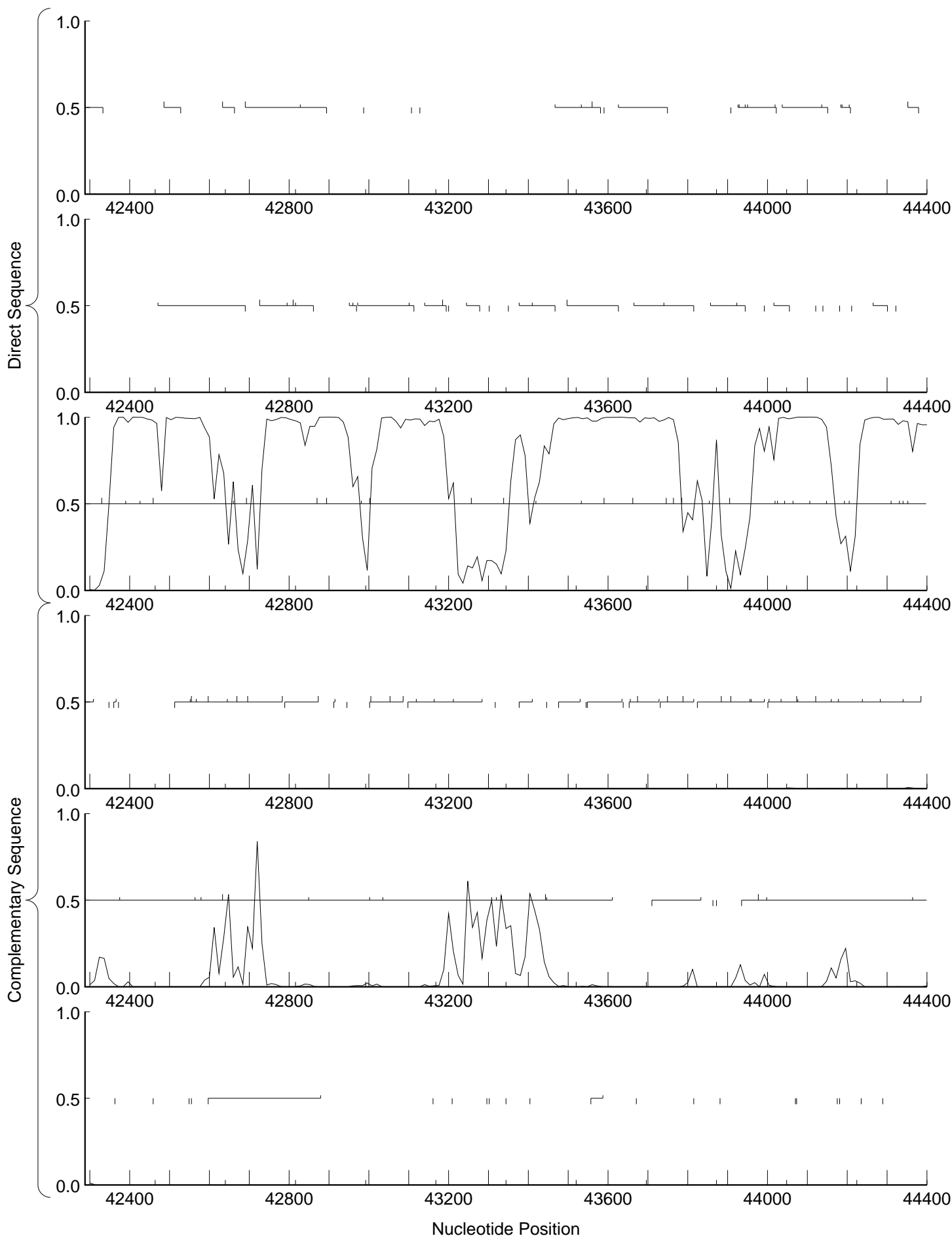


complete sequence, 56251 bp including 10-base 3' overhang (CGGAAGGCGC), Cluster F1, Order 4, Window 96, Step 12, 19/28

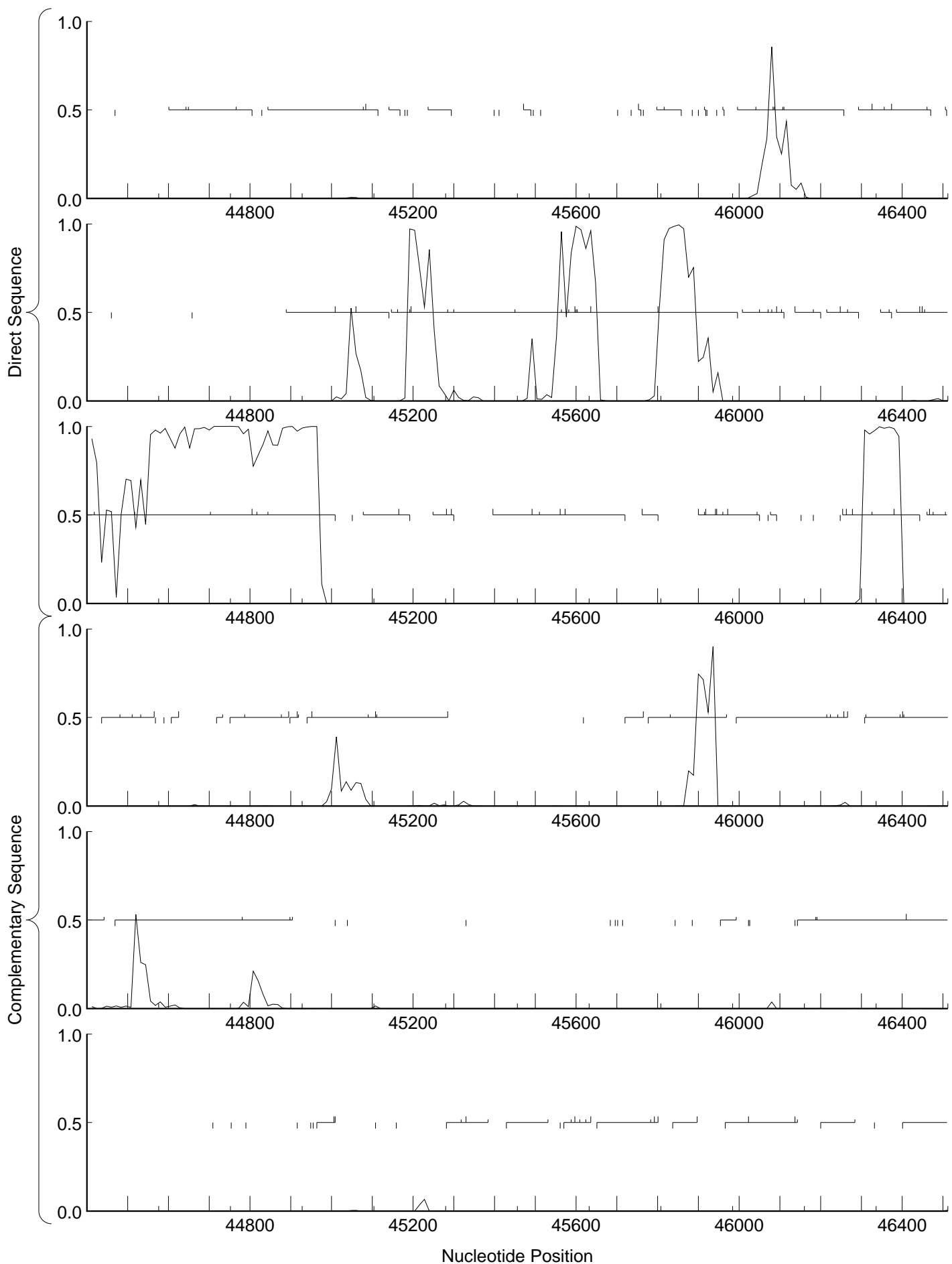


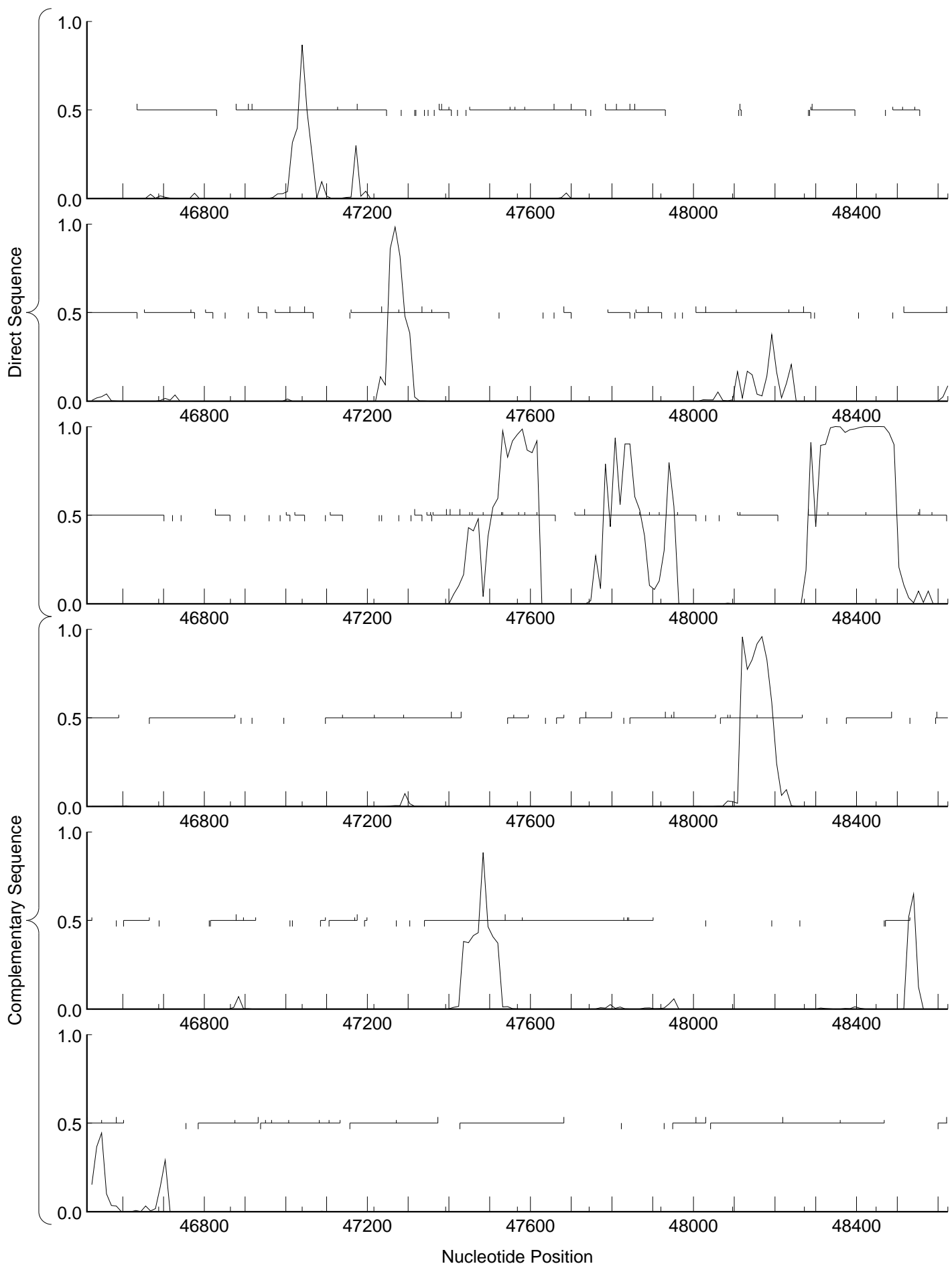


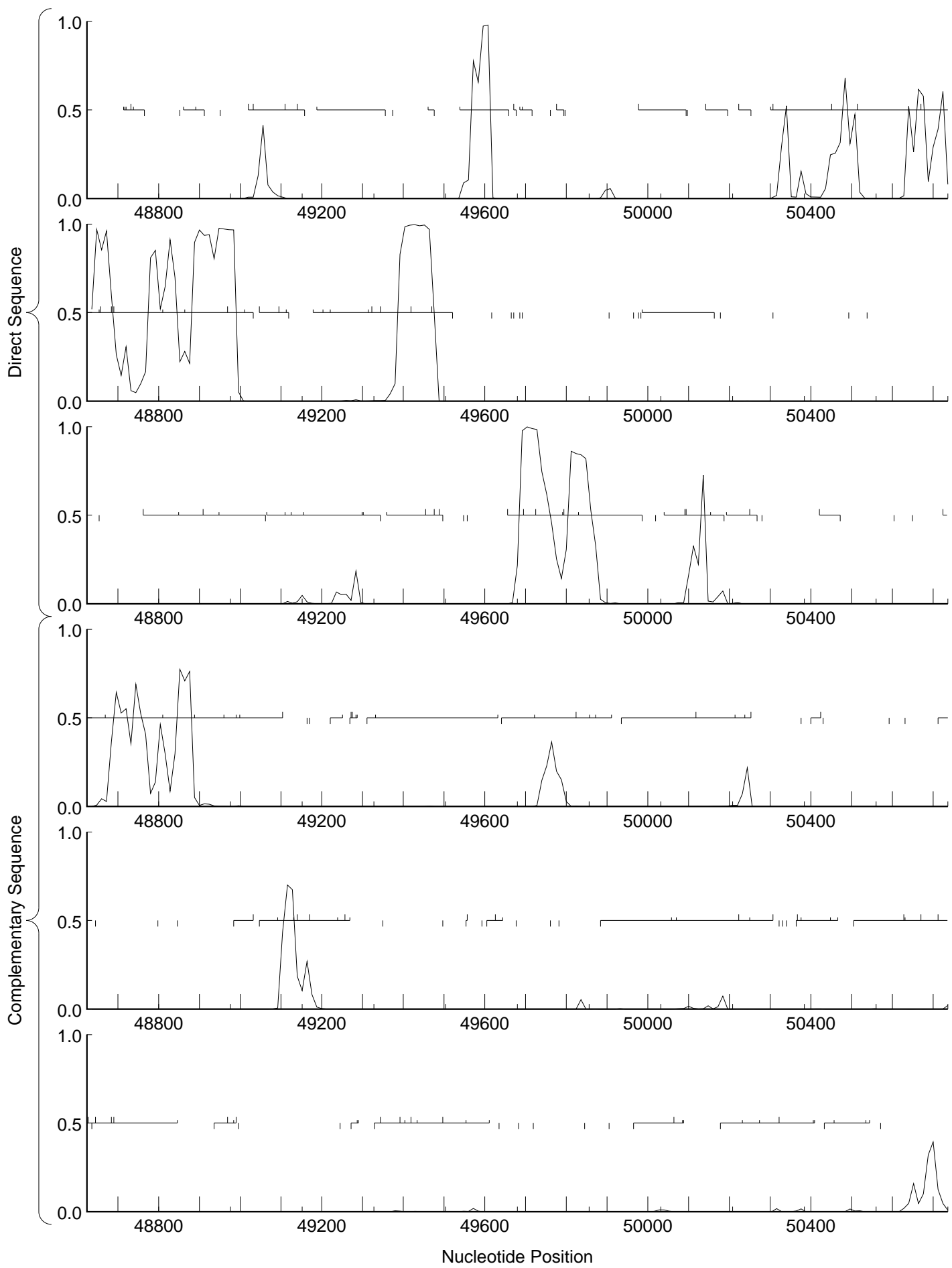


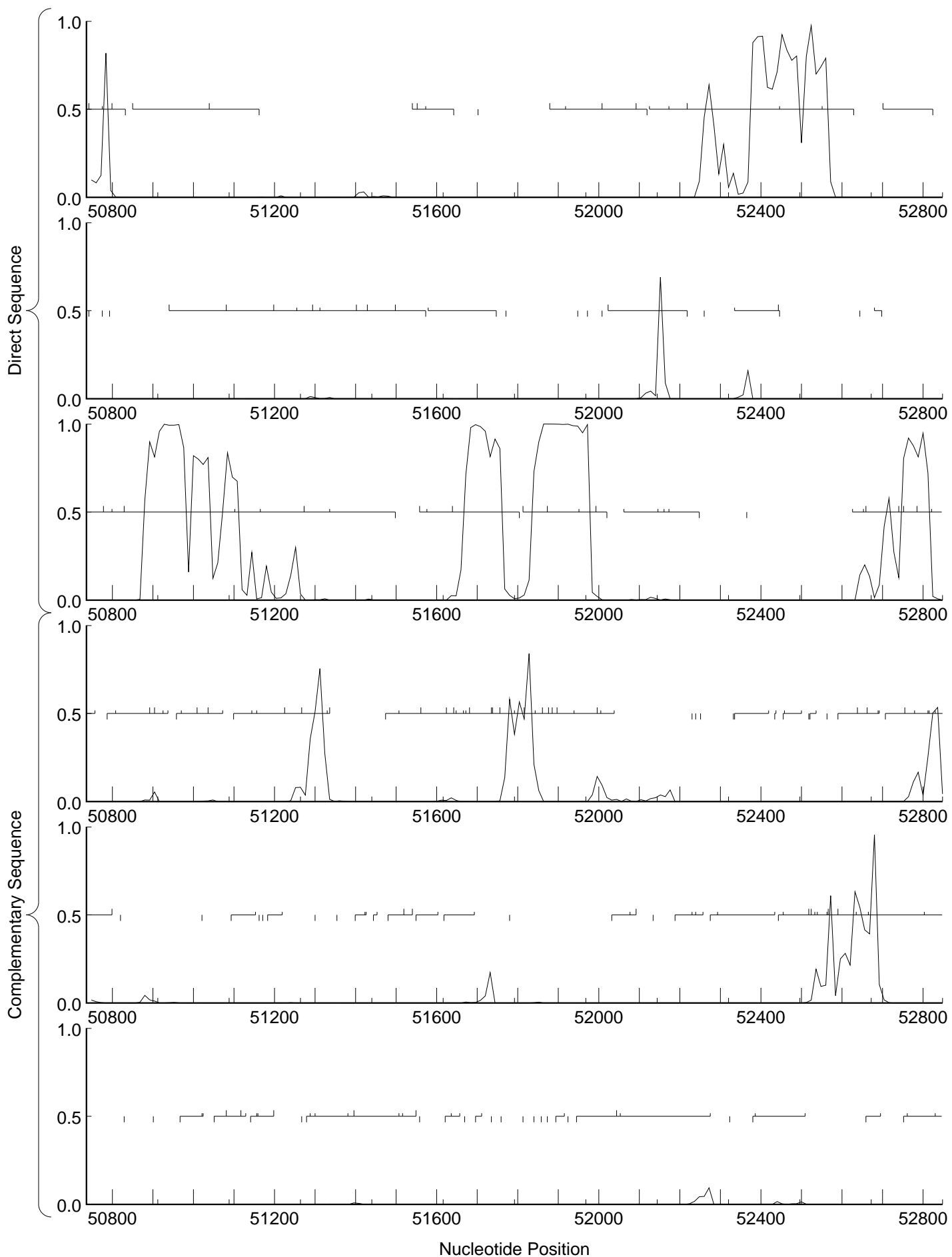


complete sequence, 56251 bp including 10-base 3' overhang (CGGAAGGCGC), Cluster F1, Order 4, Window 96, Step 12, 23/28









complete sequence, 56251 bp including 10-base 3' overhang (CGGAAGGCGC), Cluster F1, Order 4, Window 96, Step 12, 27/28

