

GeneMark

Version 2.5p (09.08.06)

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PROGRAM INFORMATION

Sequence : Mycobacterium phage Loadrie complete sequence, 76492 bp including 10-base 3' overhang (TCGA)
Analysis Date : 2/3/23 at 5:04:22
Pages : 38
Sequence Length : 76492 bp
GC Content : 58.95%

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

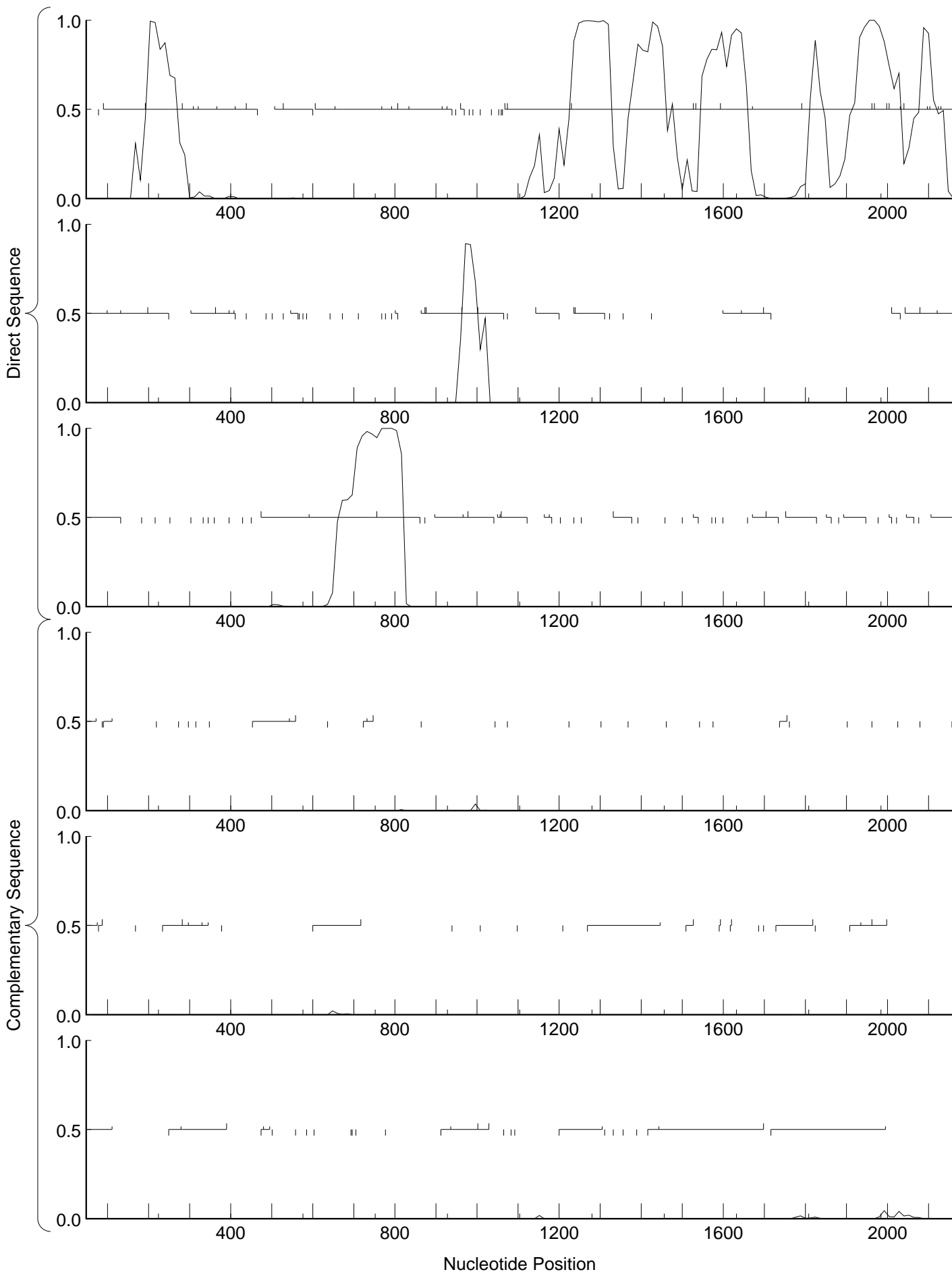
MATRIX INFORMATION

Matrix : Mycobacterium_tuberculosis_H37Rv
Author : -
Order : 4

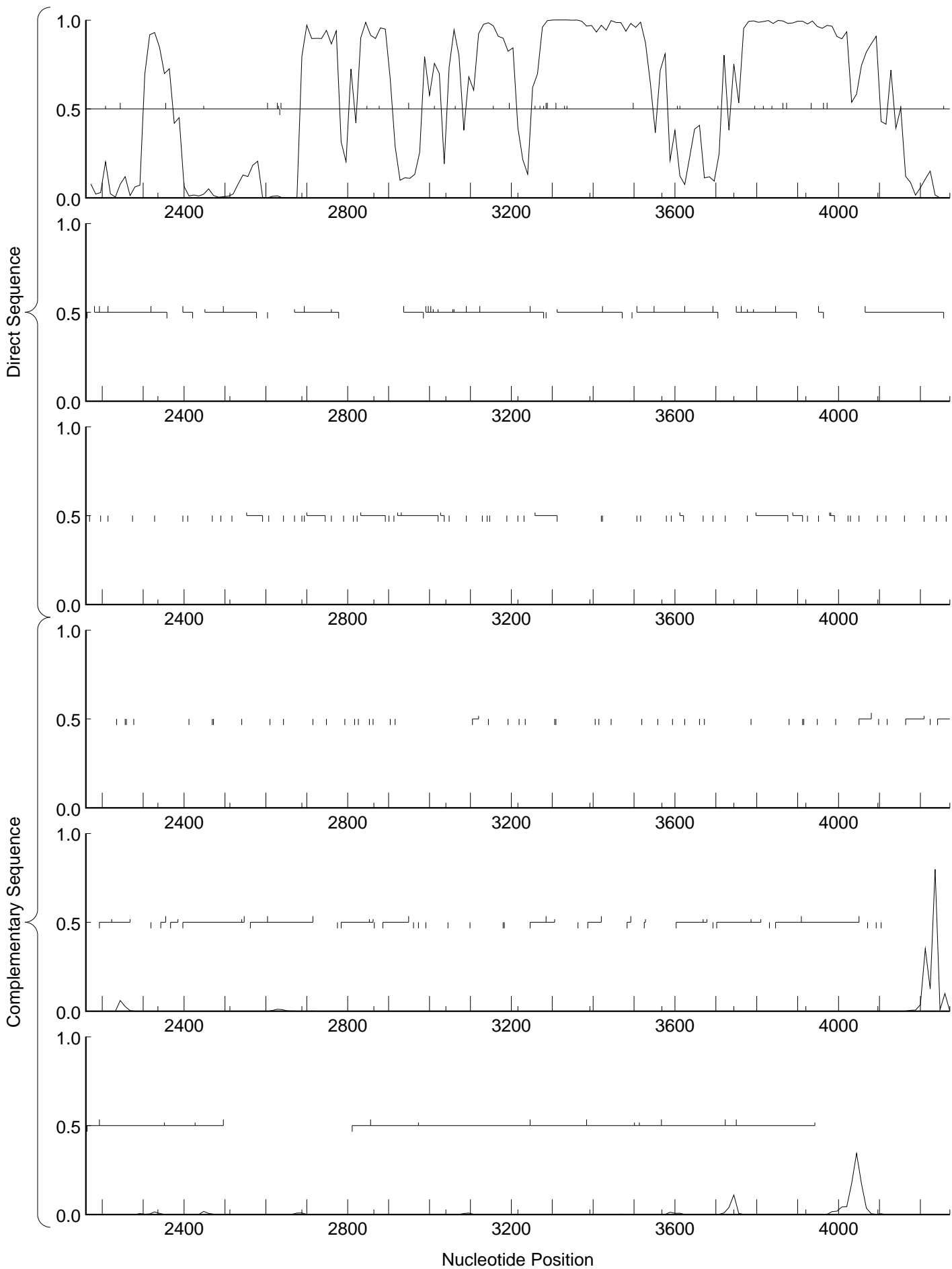
Send questions / comments to:
Dr. M. Borodovsky
Georgia Institute of Technology
School of Biology
Atlanta, GA 30332-0230

For help or information, send electronic mail to: custserv@genepro.com, or mark@amber.biology.gatech.edu

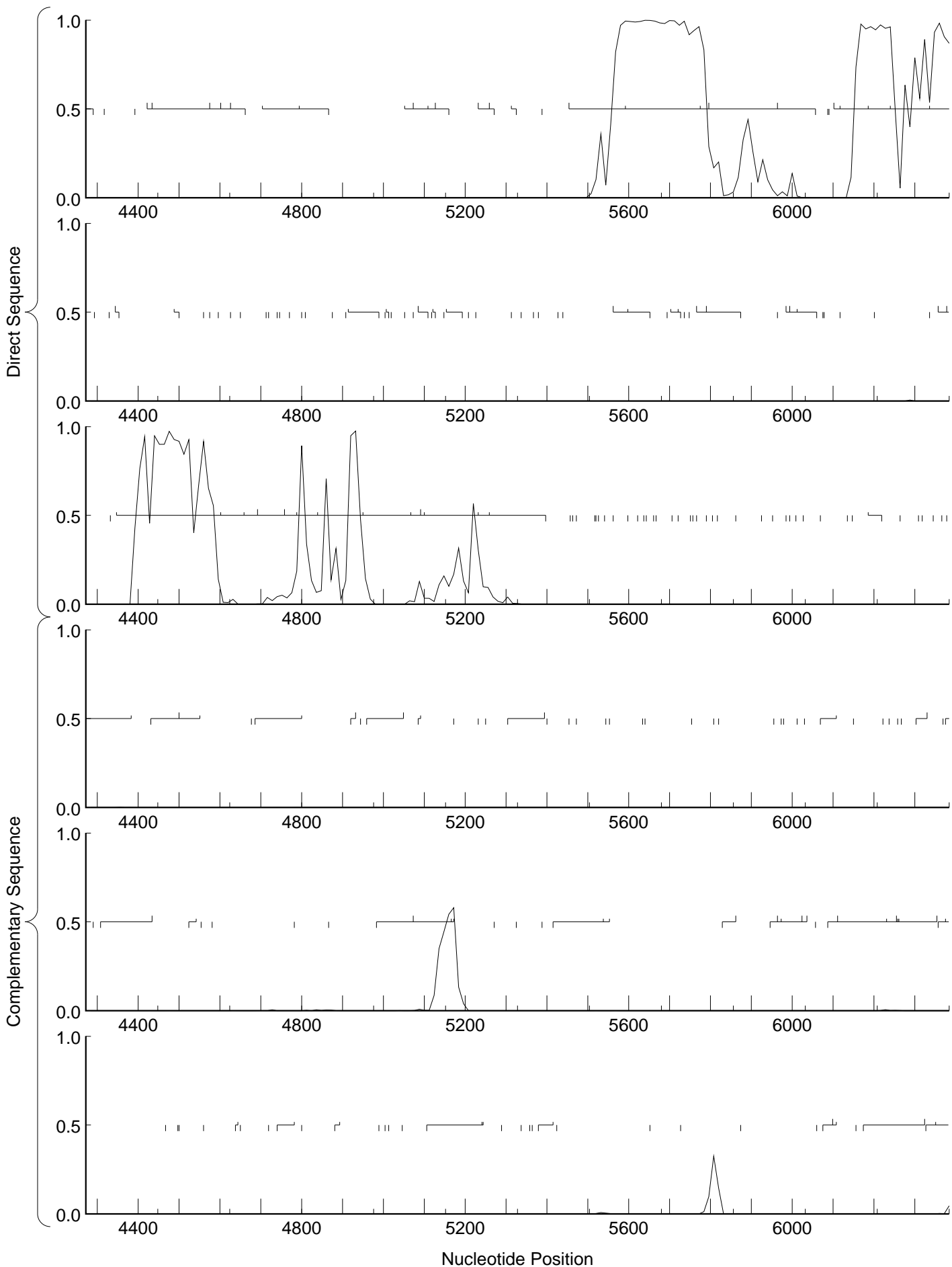
ie complete sequence, 76492 bp including 10-base 3' overhang (TCGATCAGCC), Cluster L2, Order 4, Window 96, Step 12, 2/38



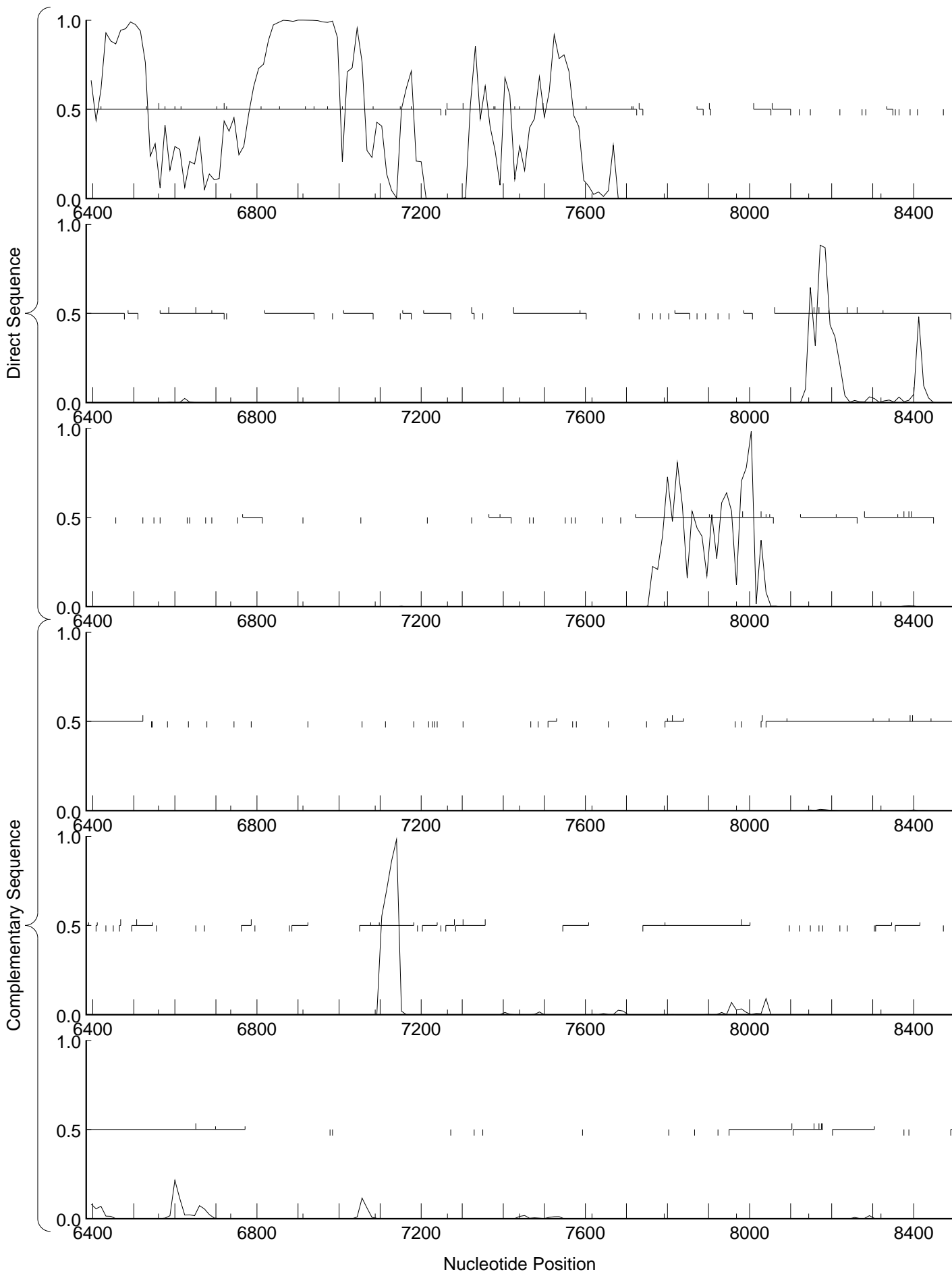
ie complete sequence, 76492 bp including 10-base 3' overhang (TCGATCAGCC), Cluster L2, Order 4, Window 96, Step 12, 3/38

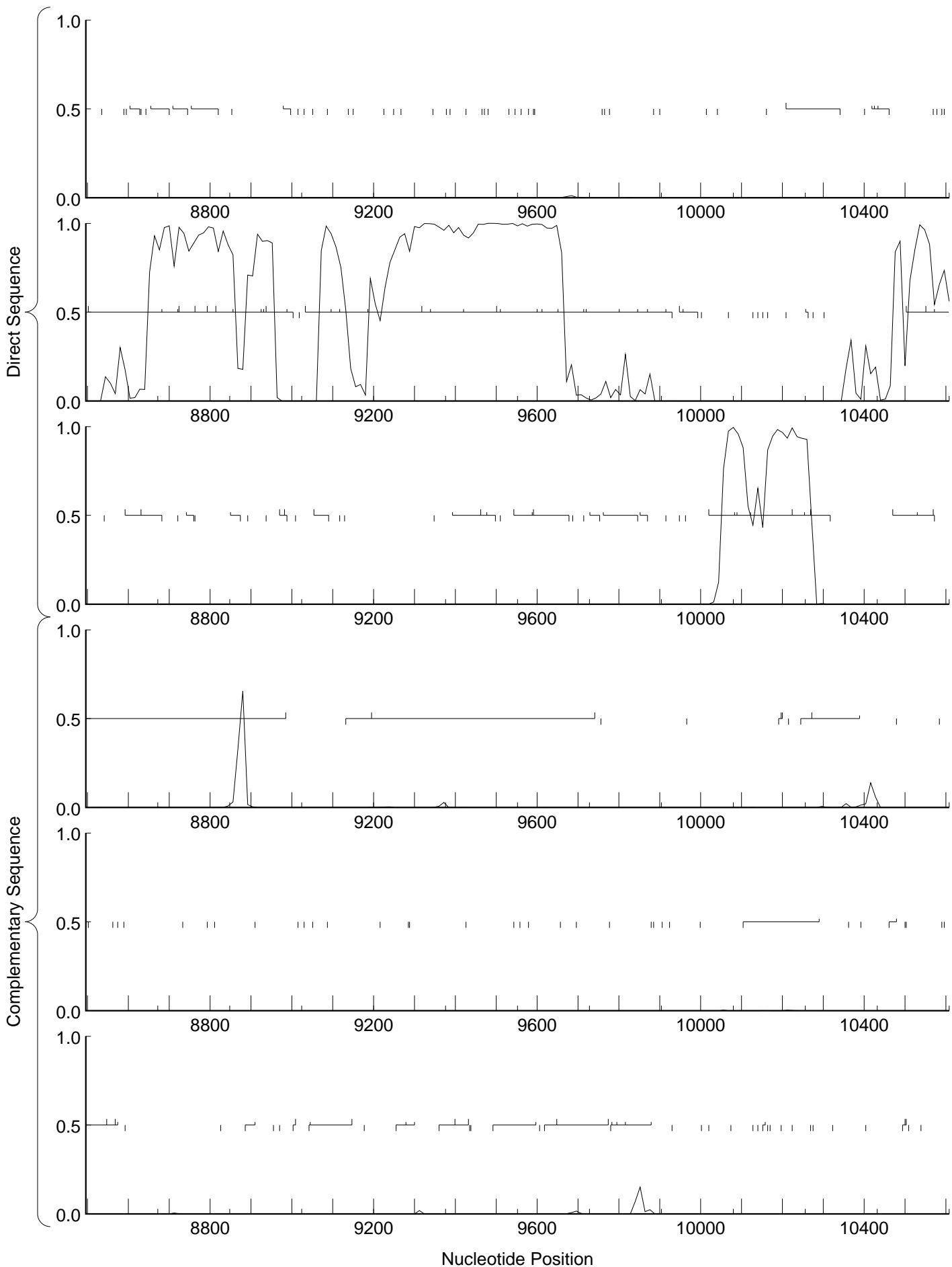


ie complete sequence, 76492 bp including 10-base 3' overhang (TCGATCAGCC), Cluster L2, Order 4, Window 96, Step 12, 4/38

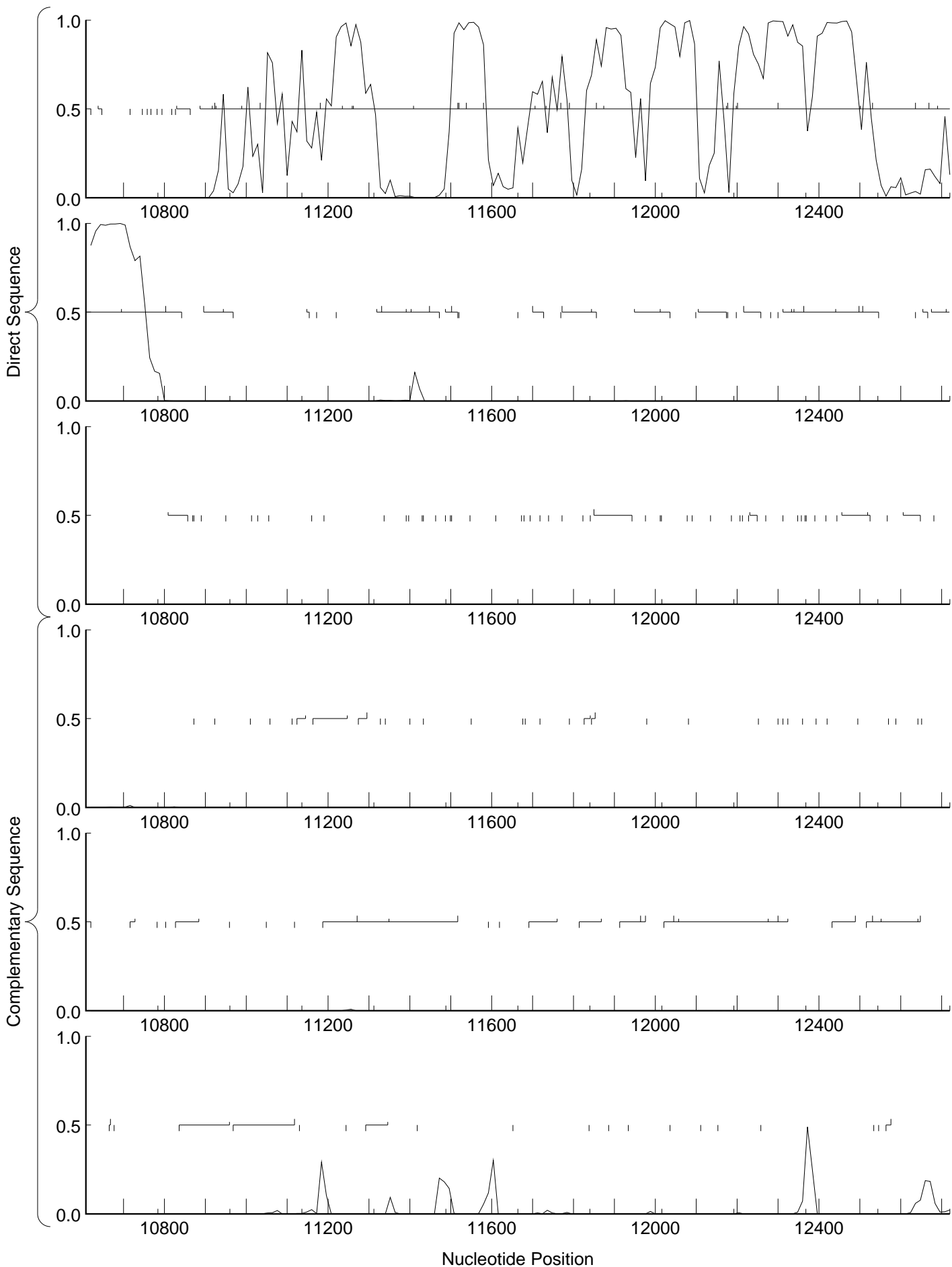


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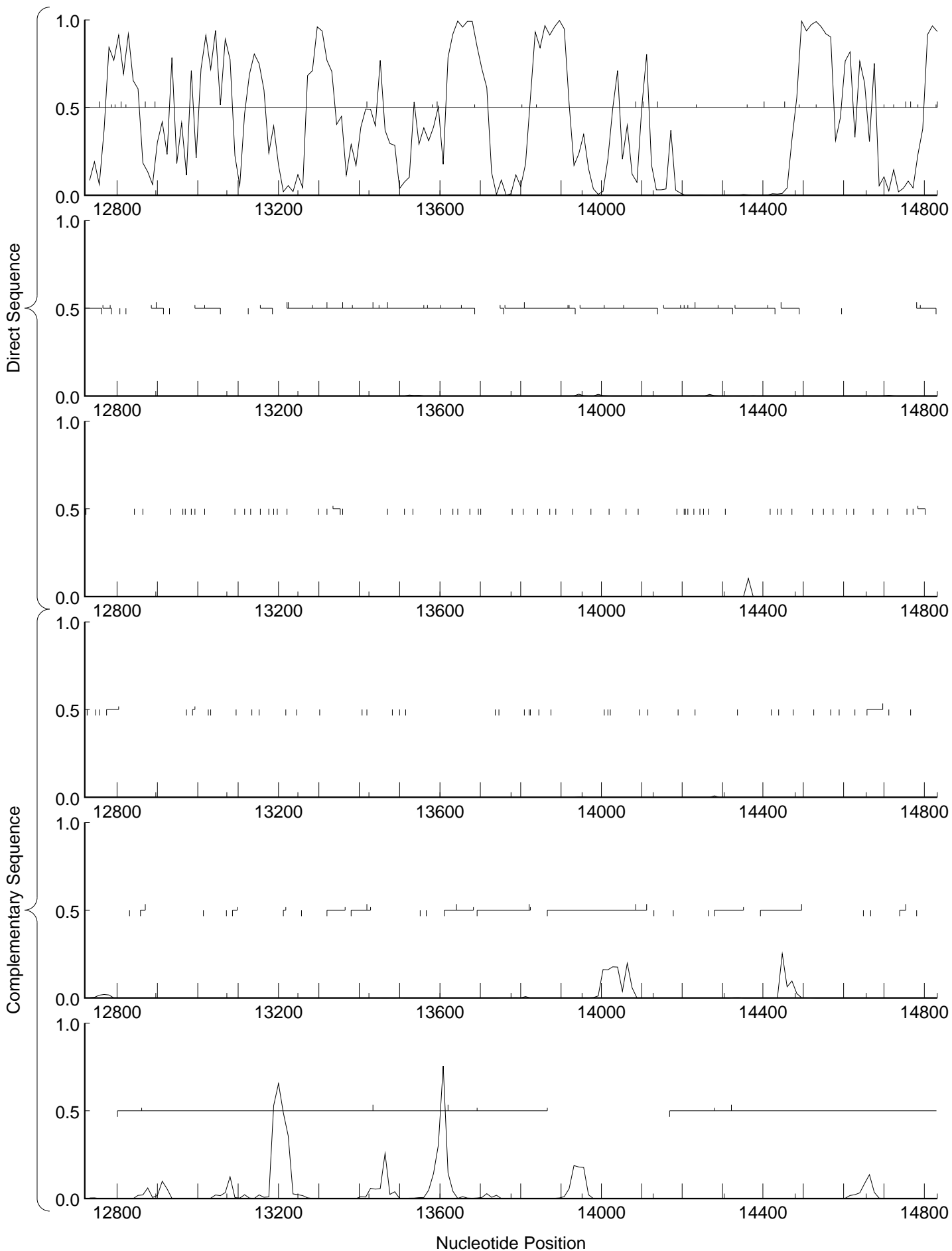




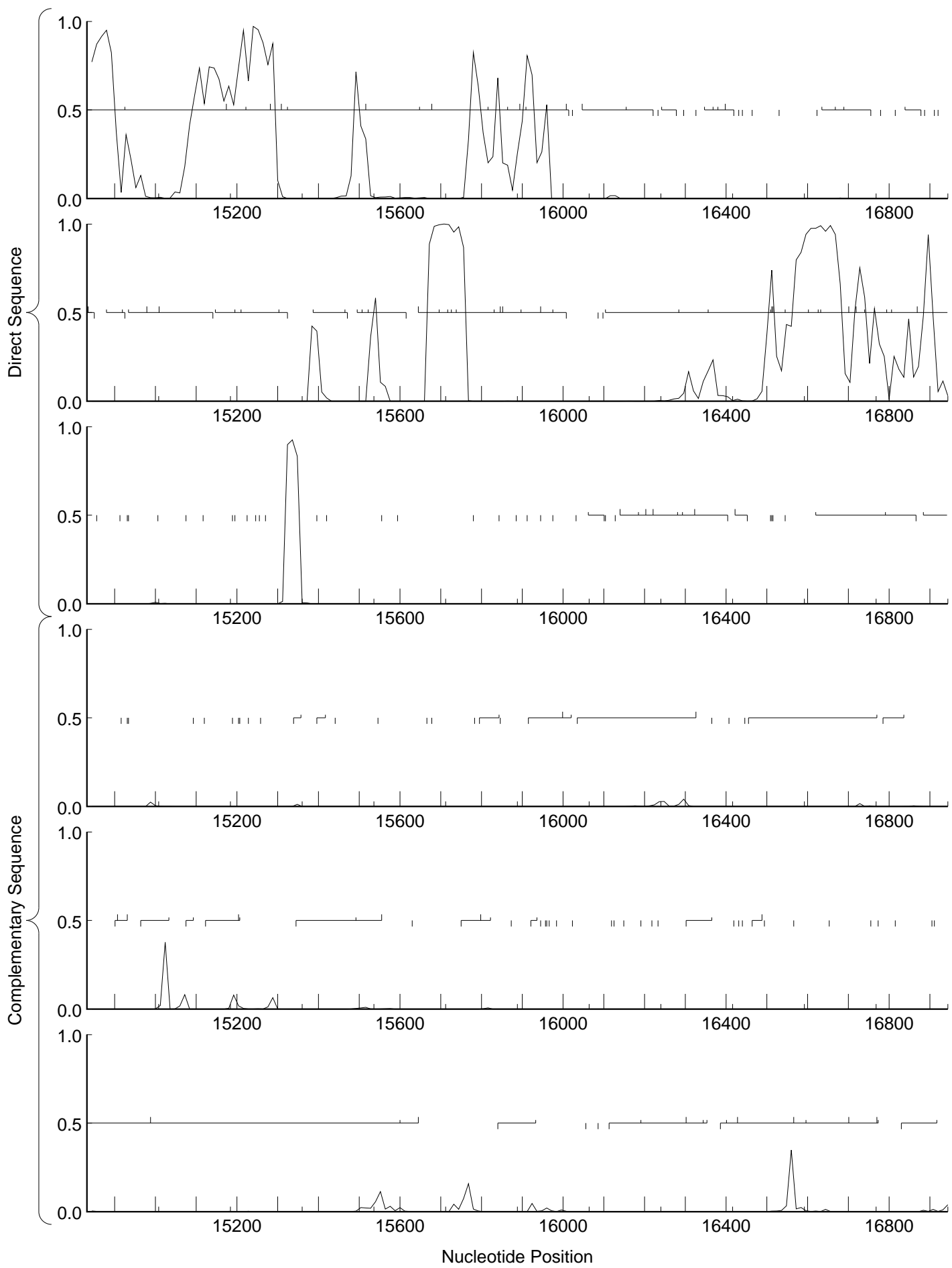
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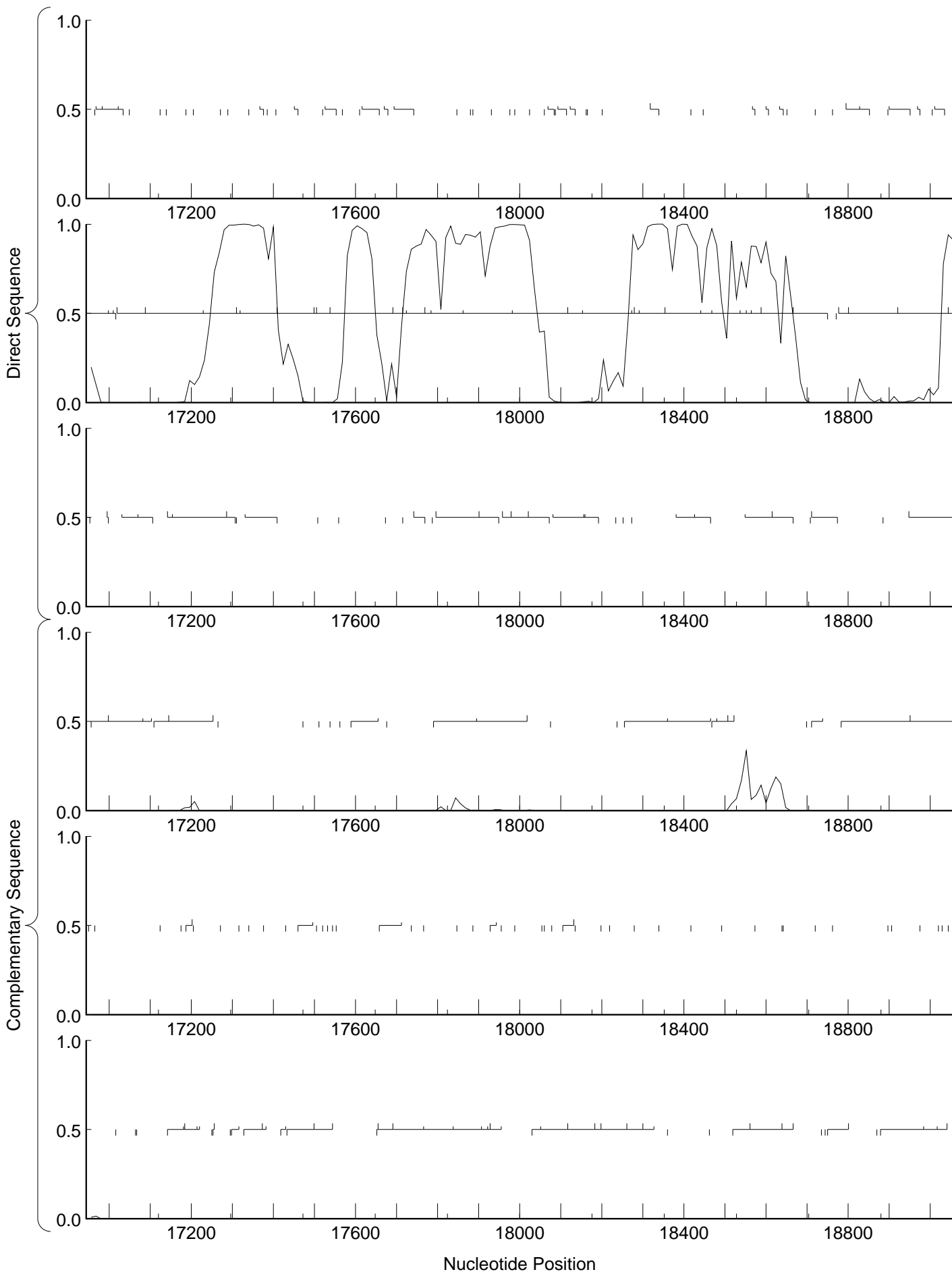


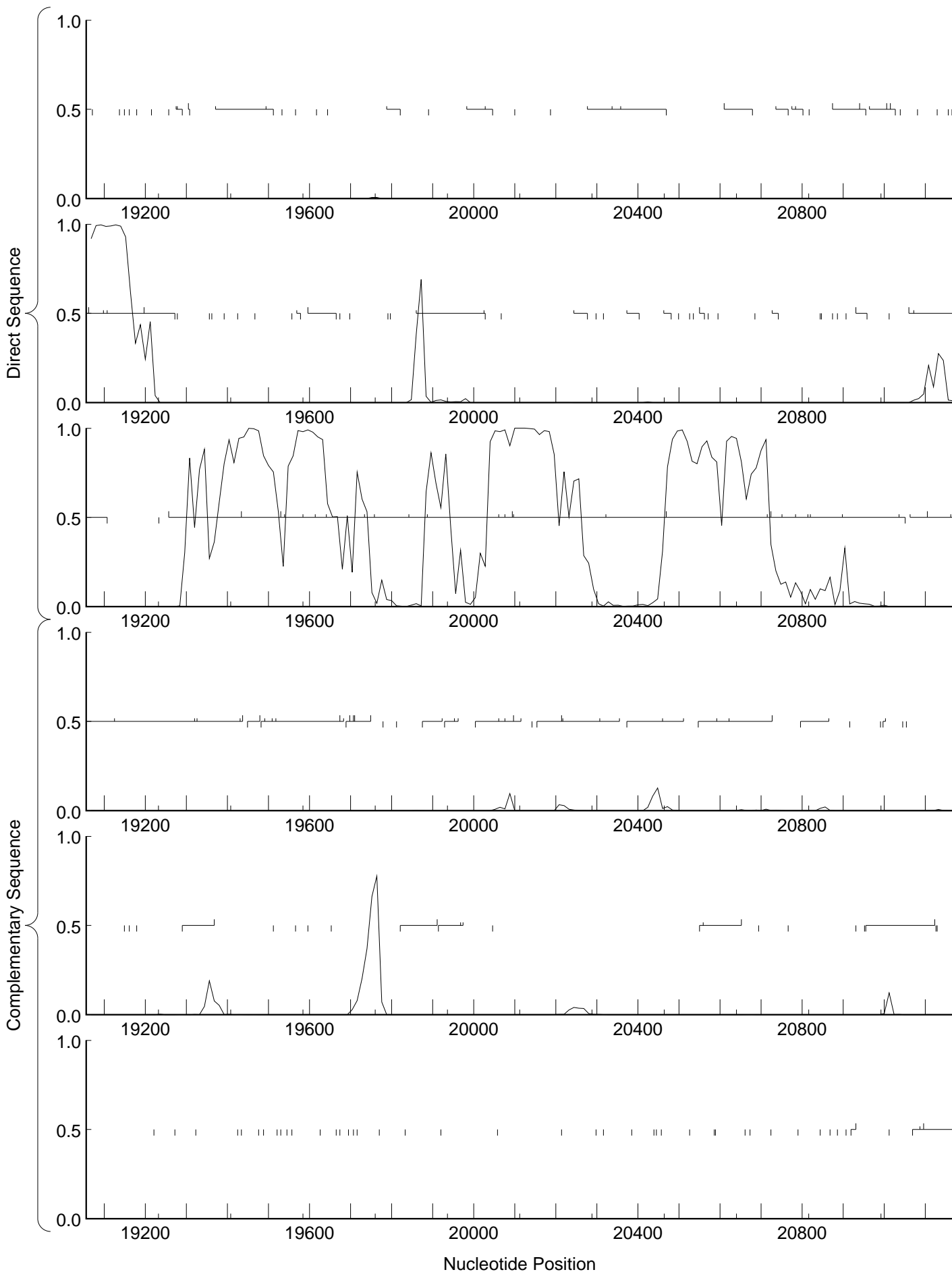
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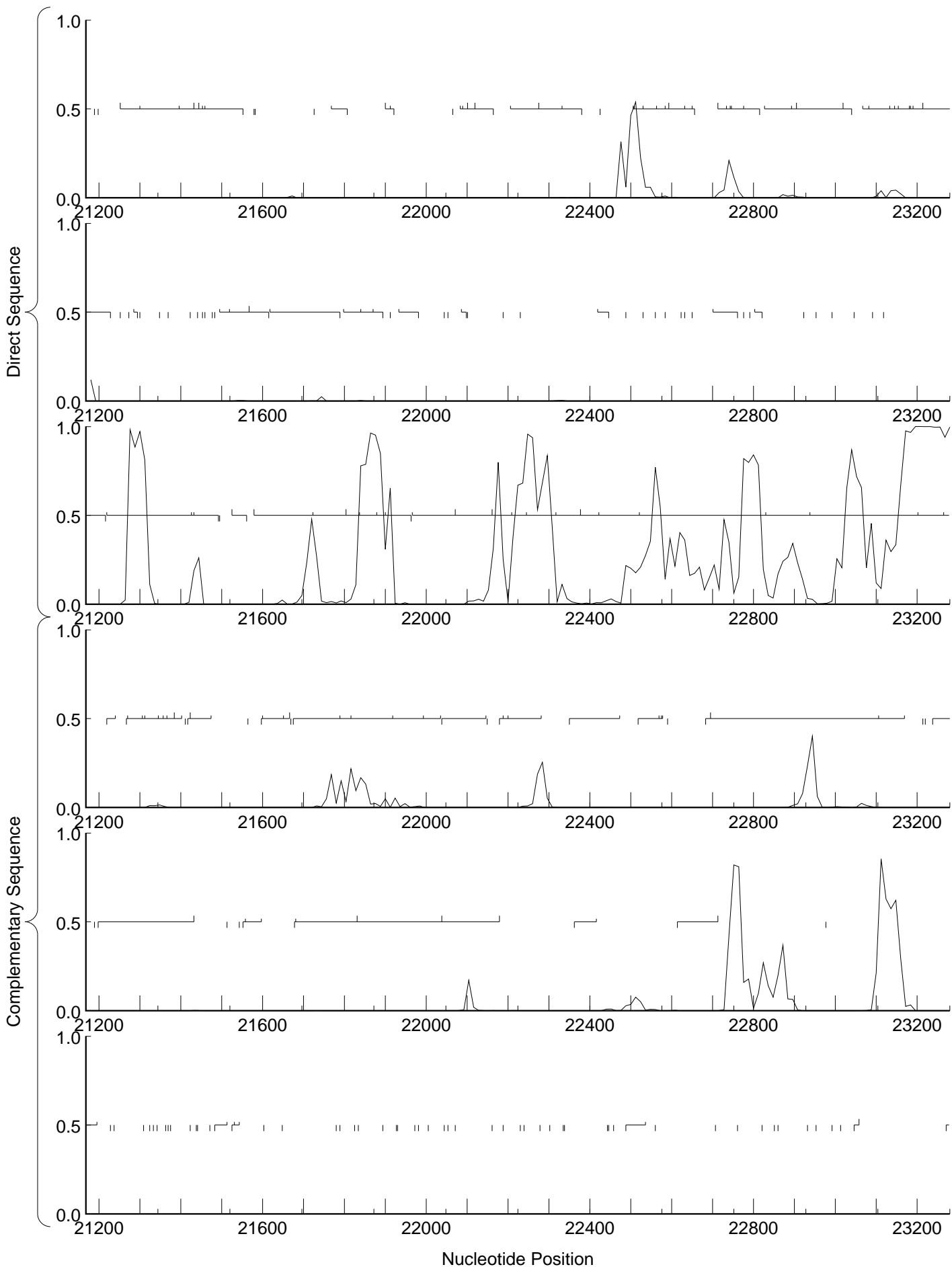


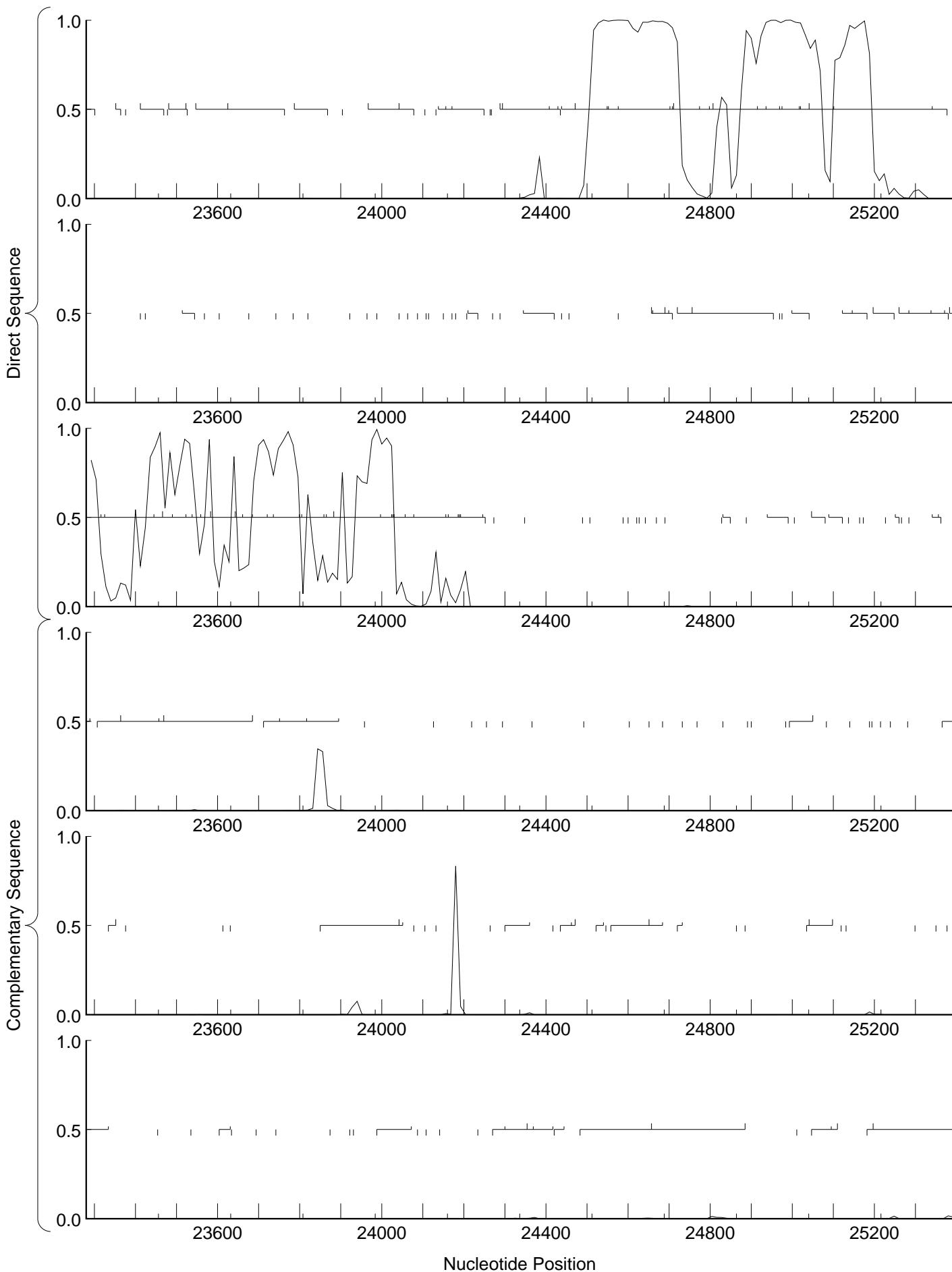
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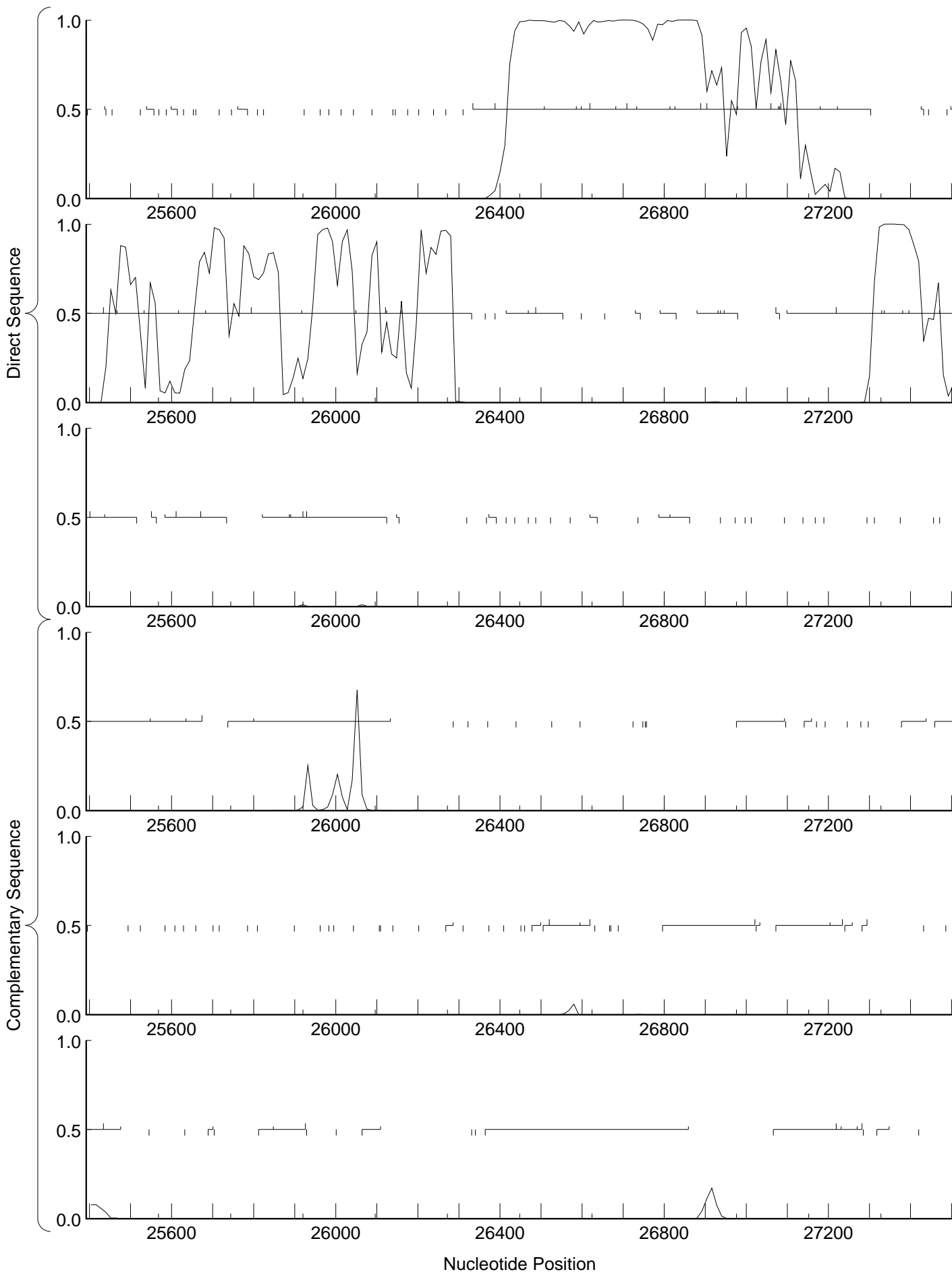


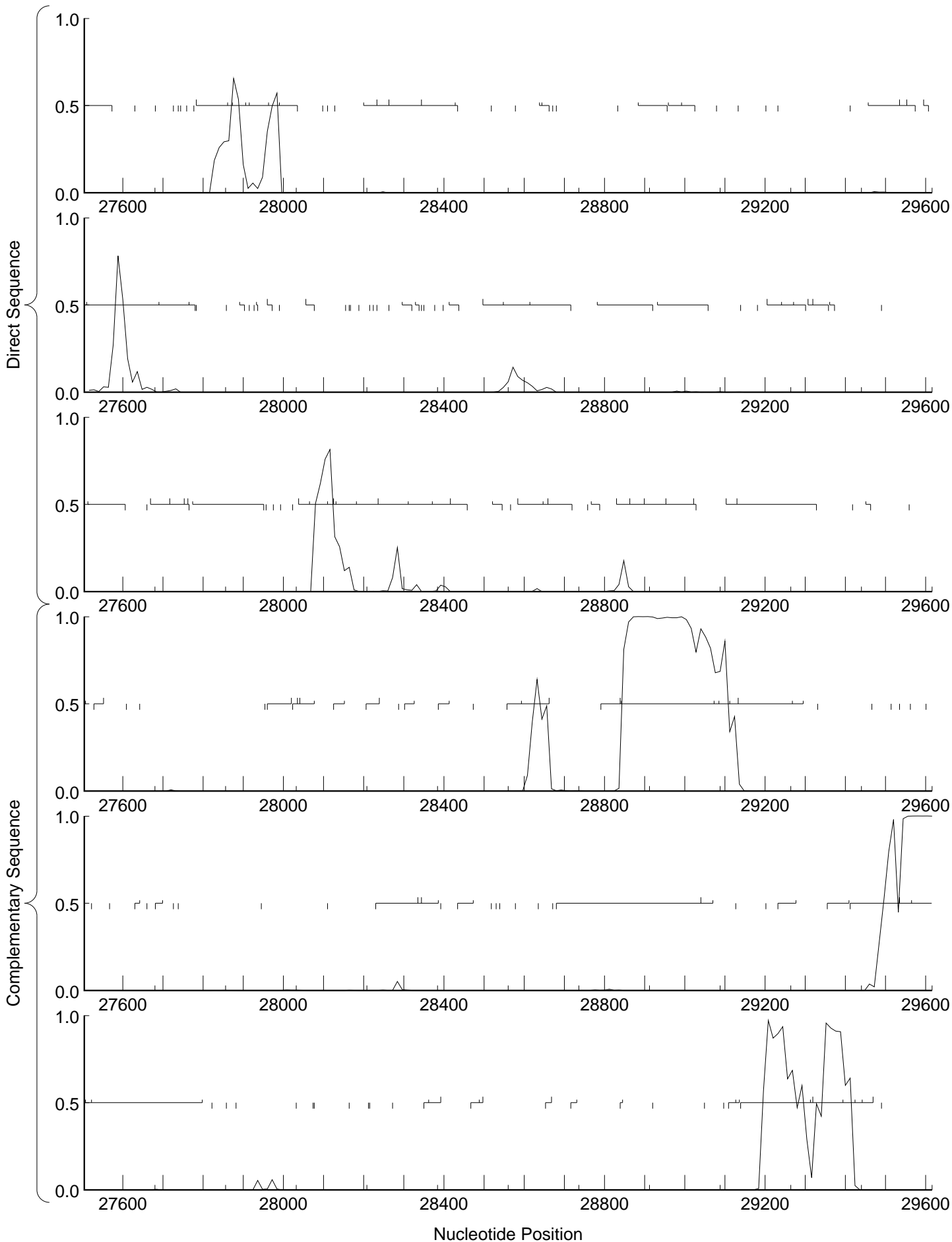


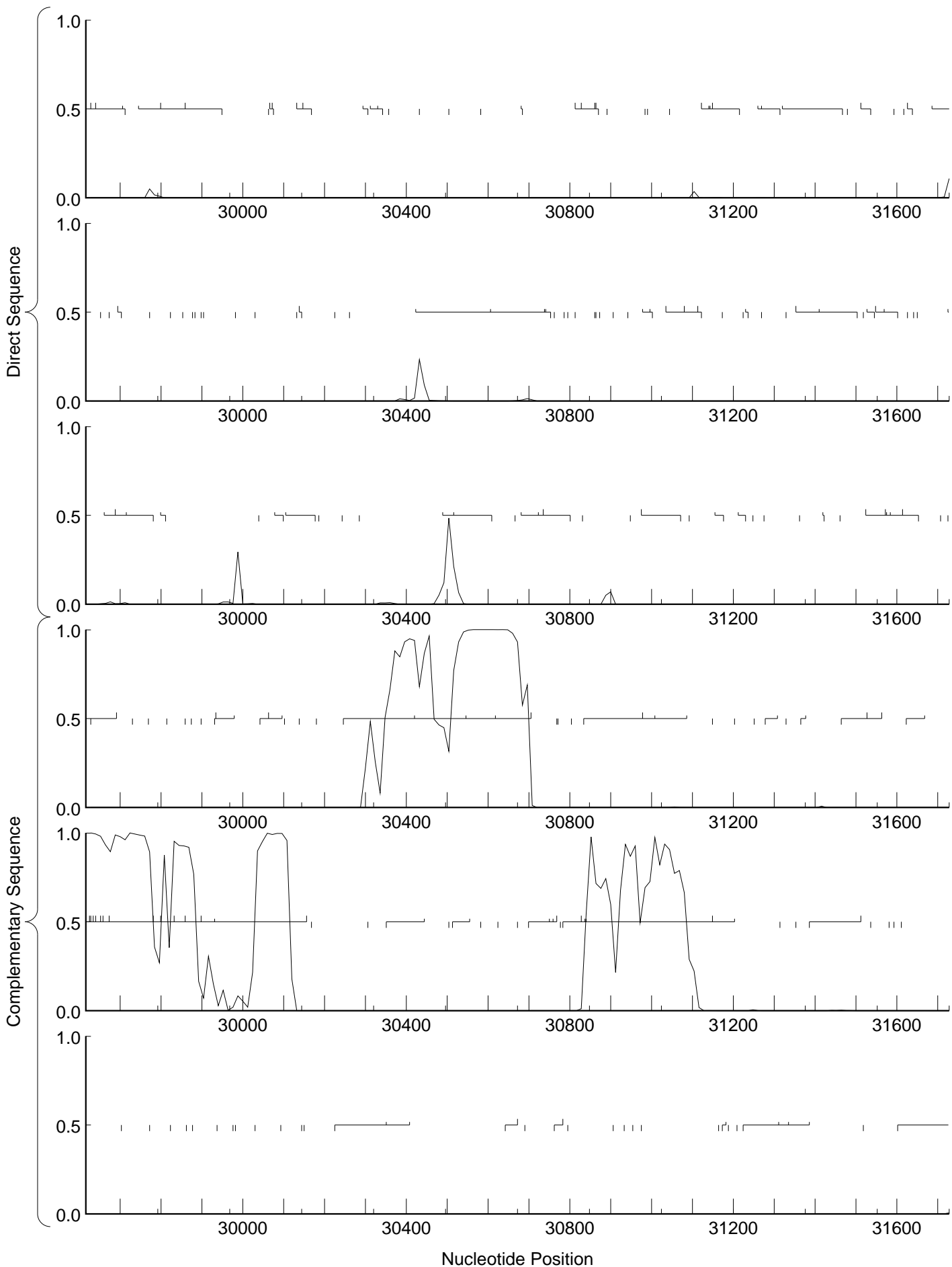


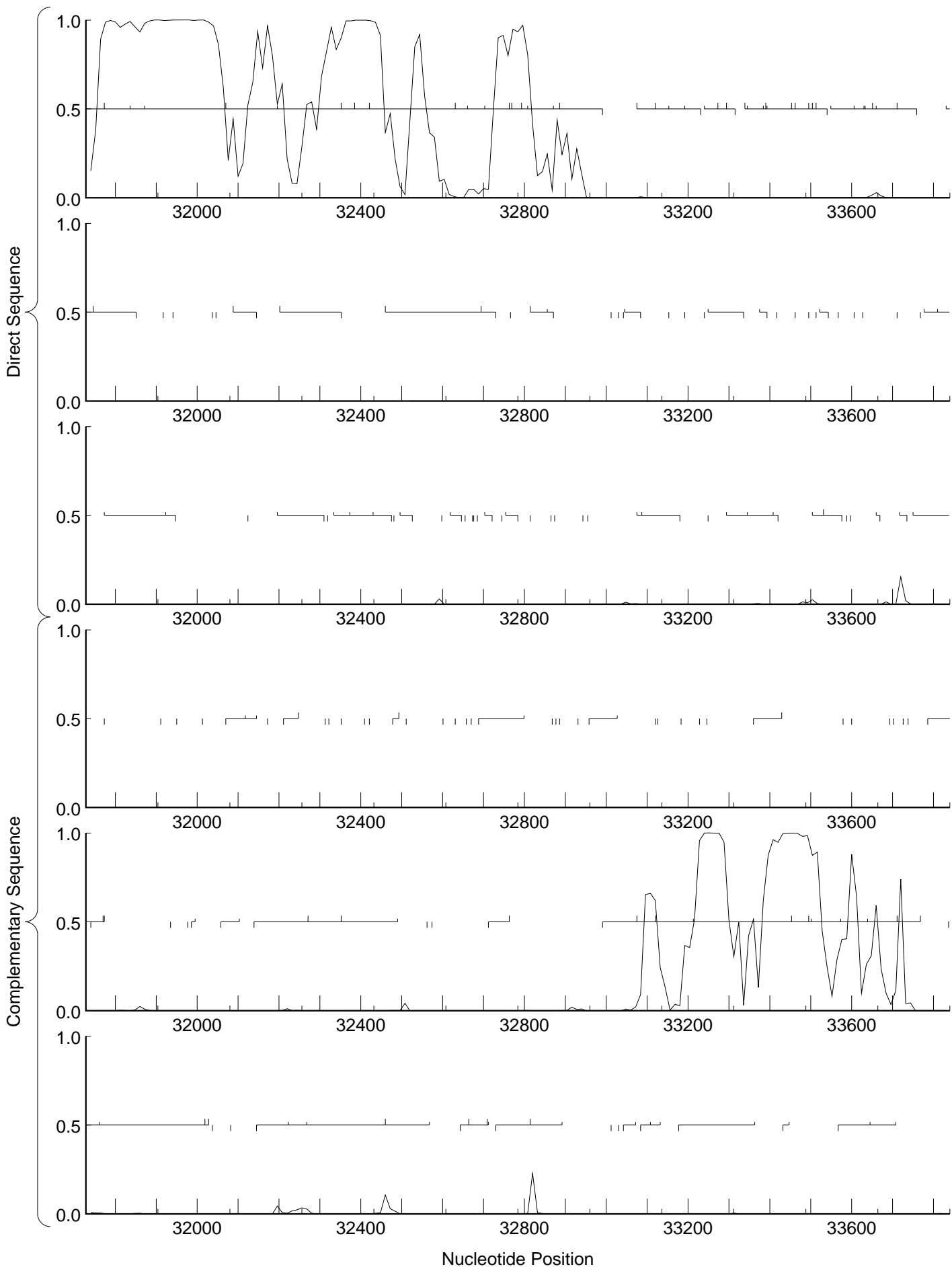


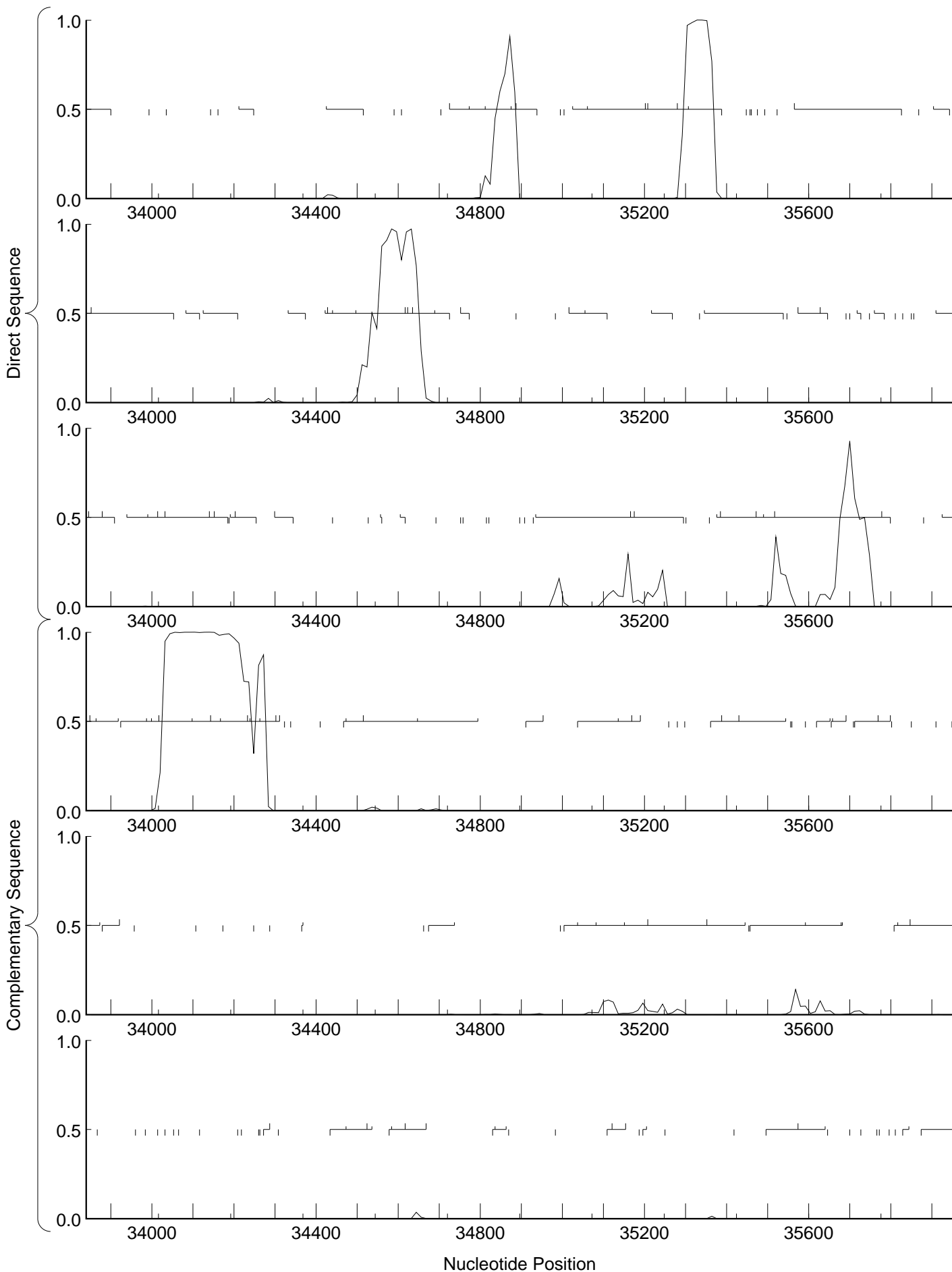


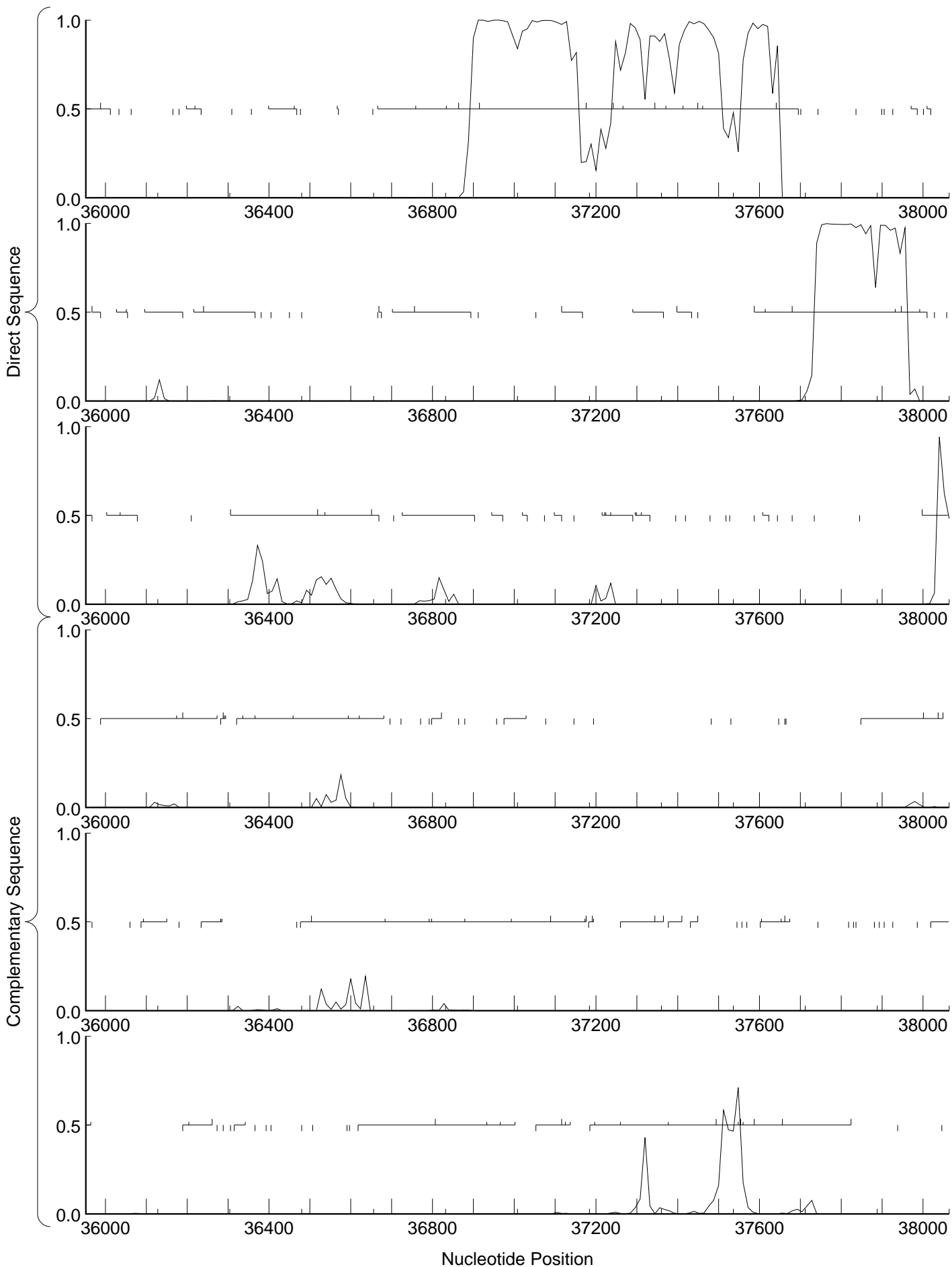


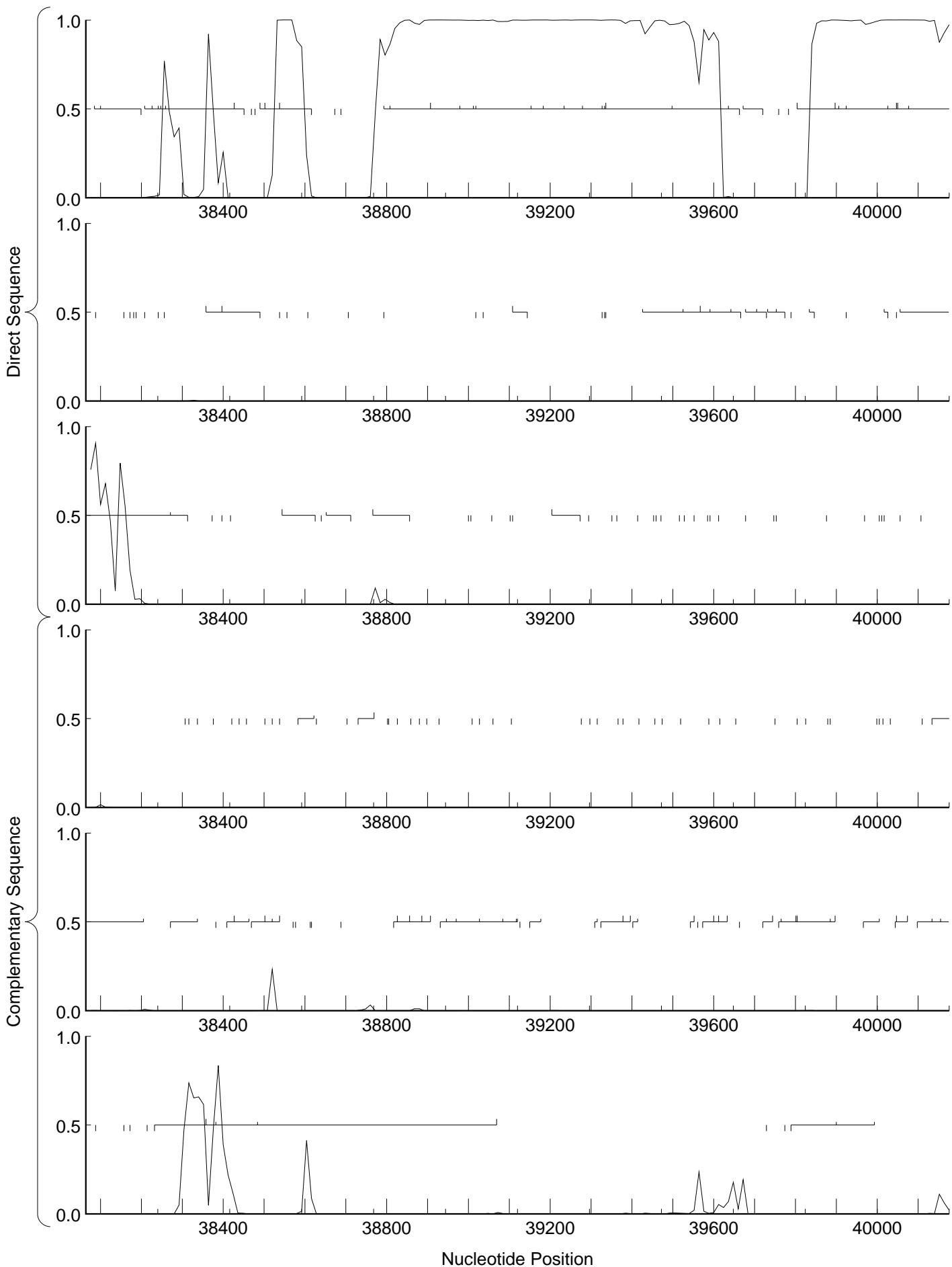


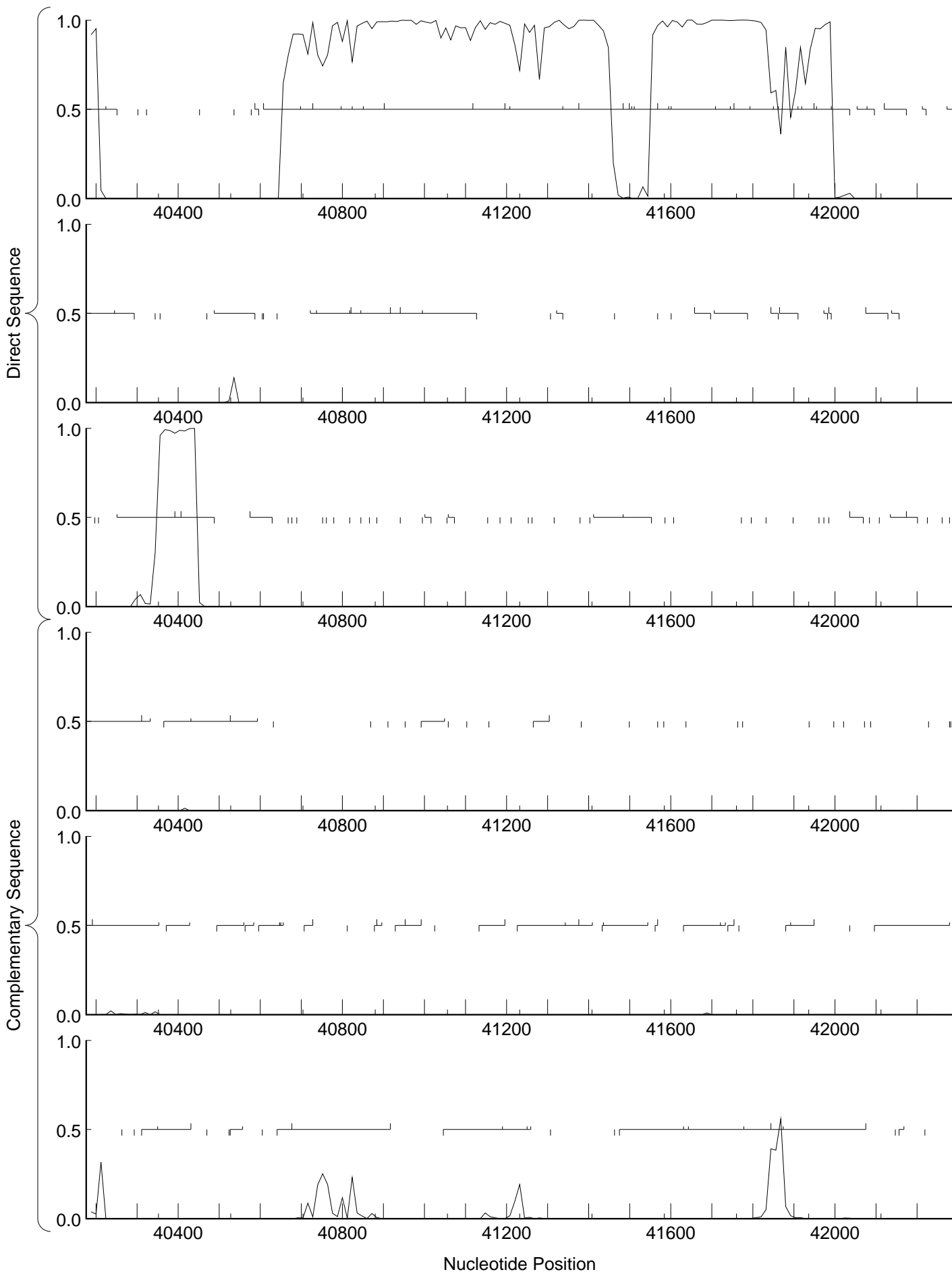


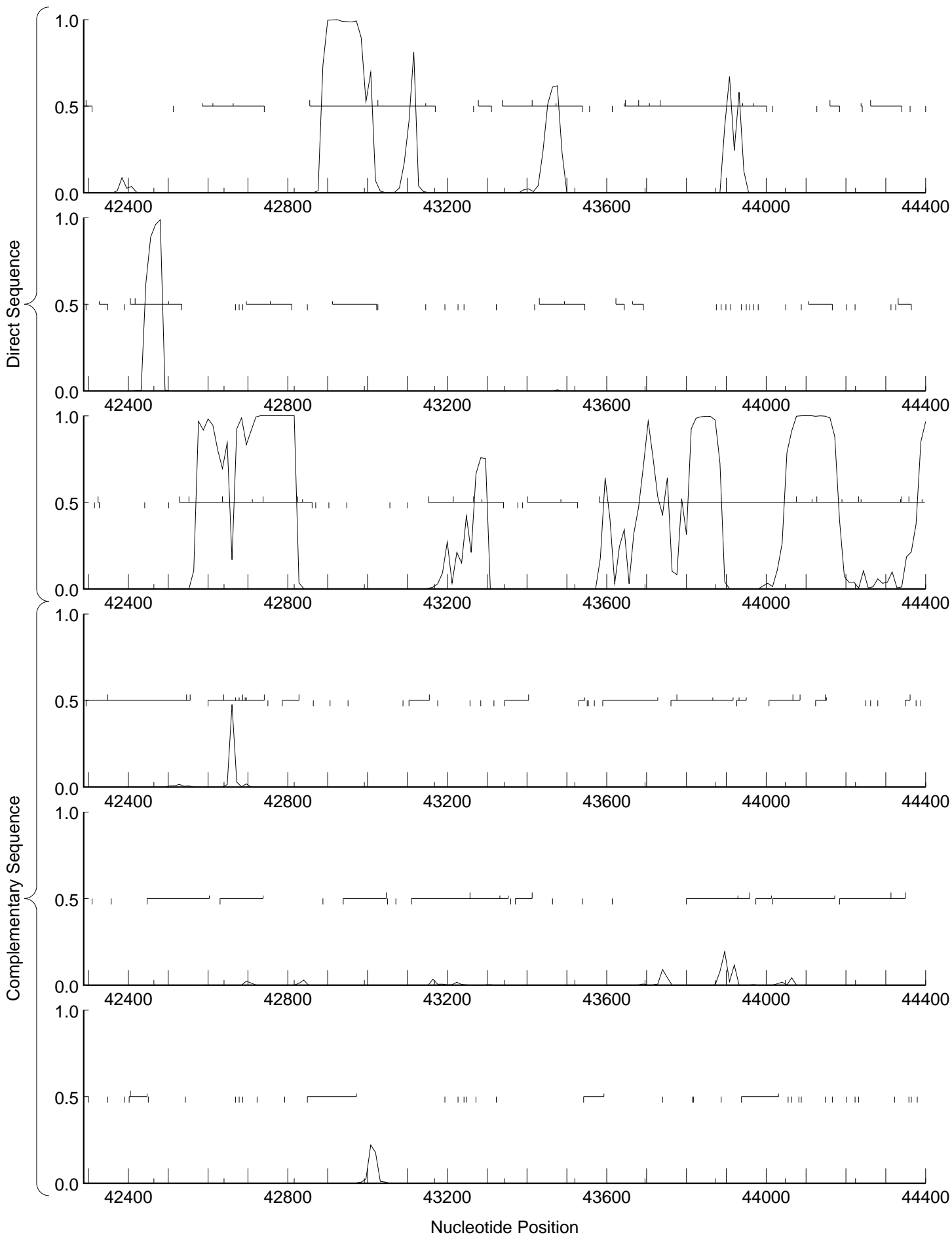


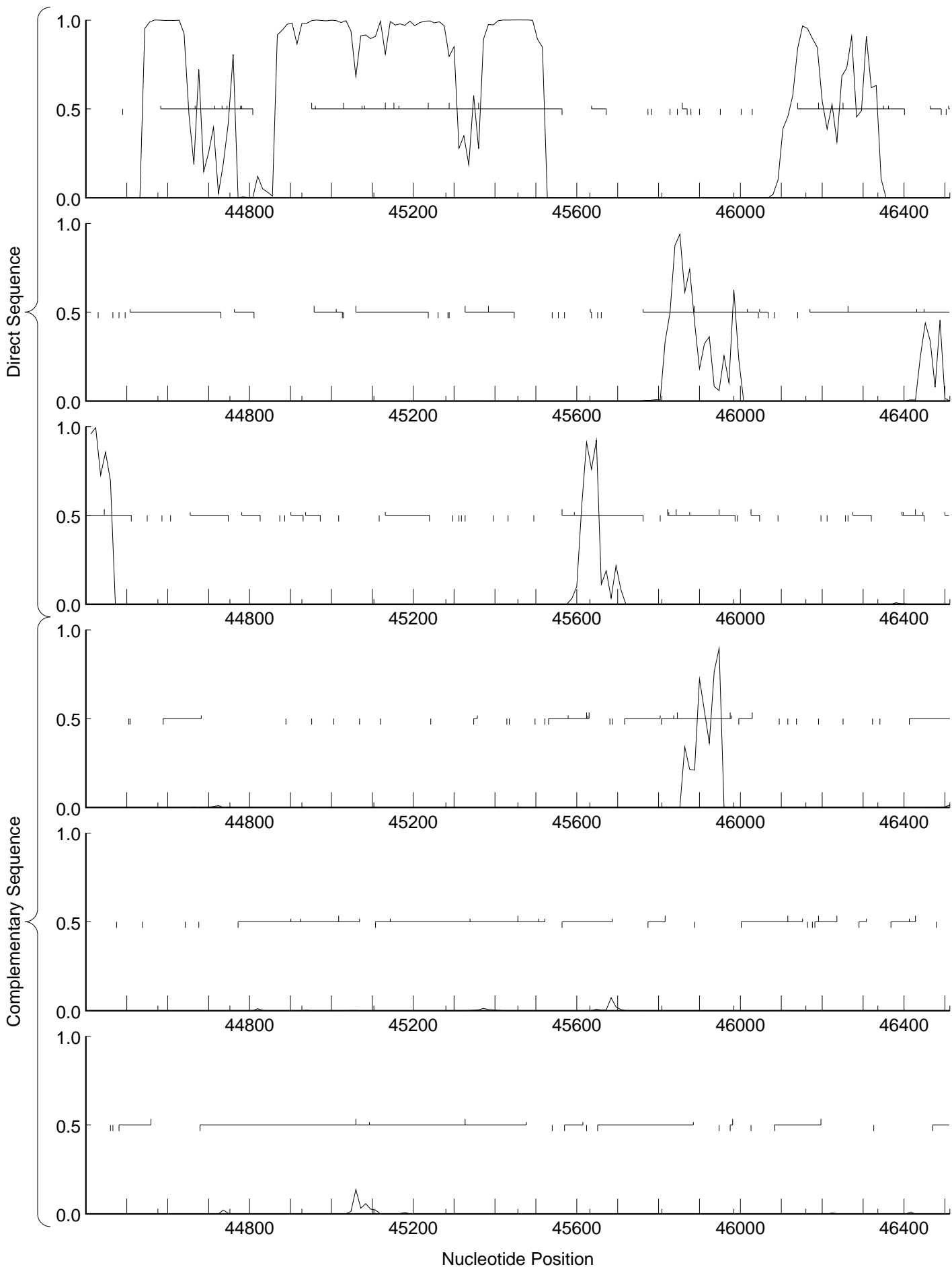


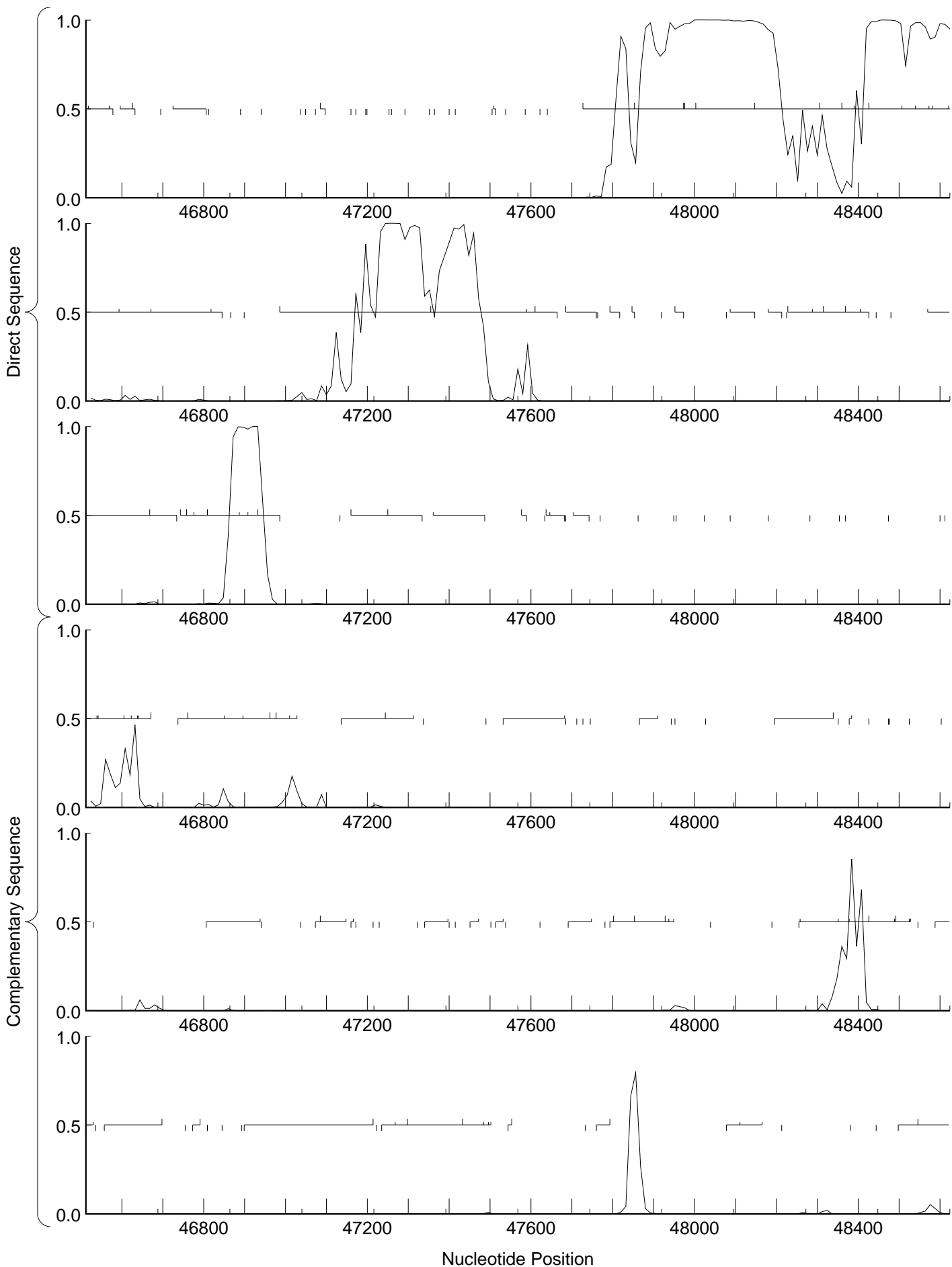


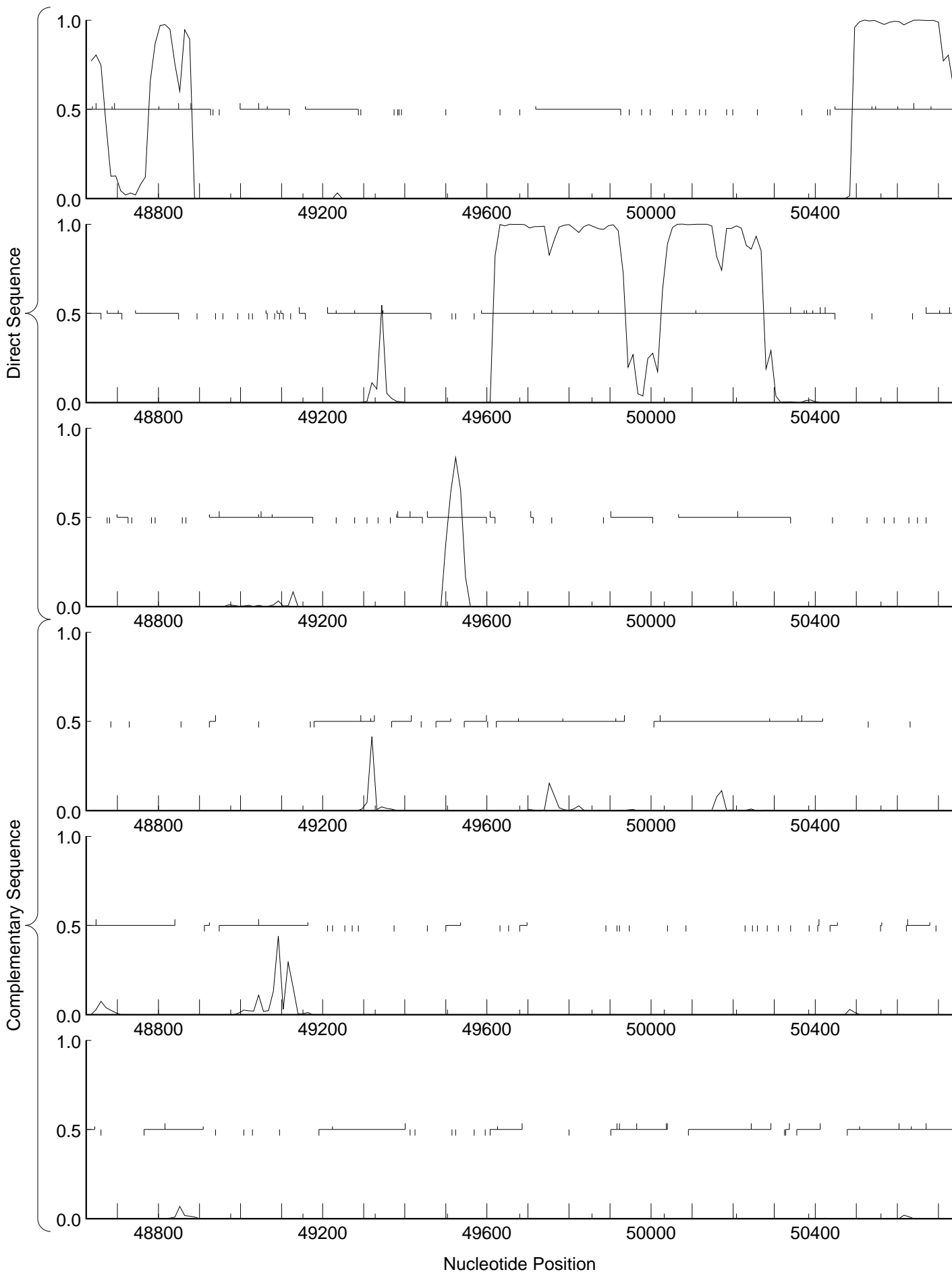


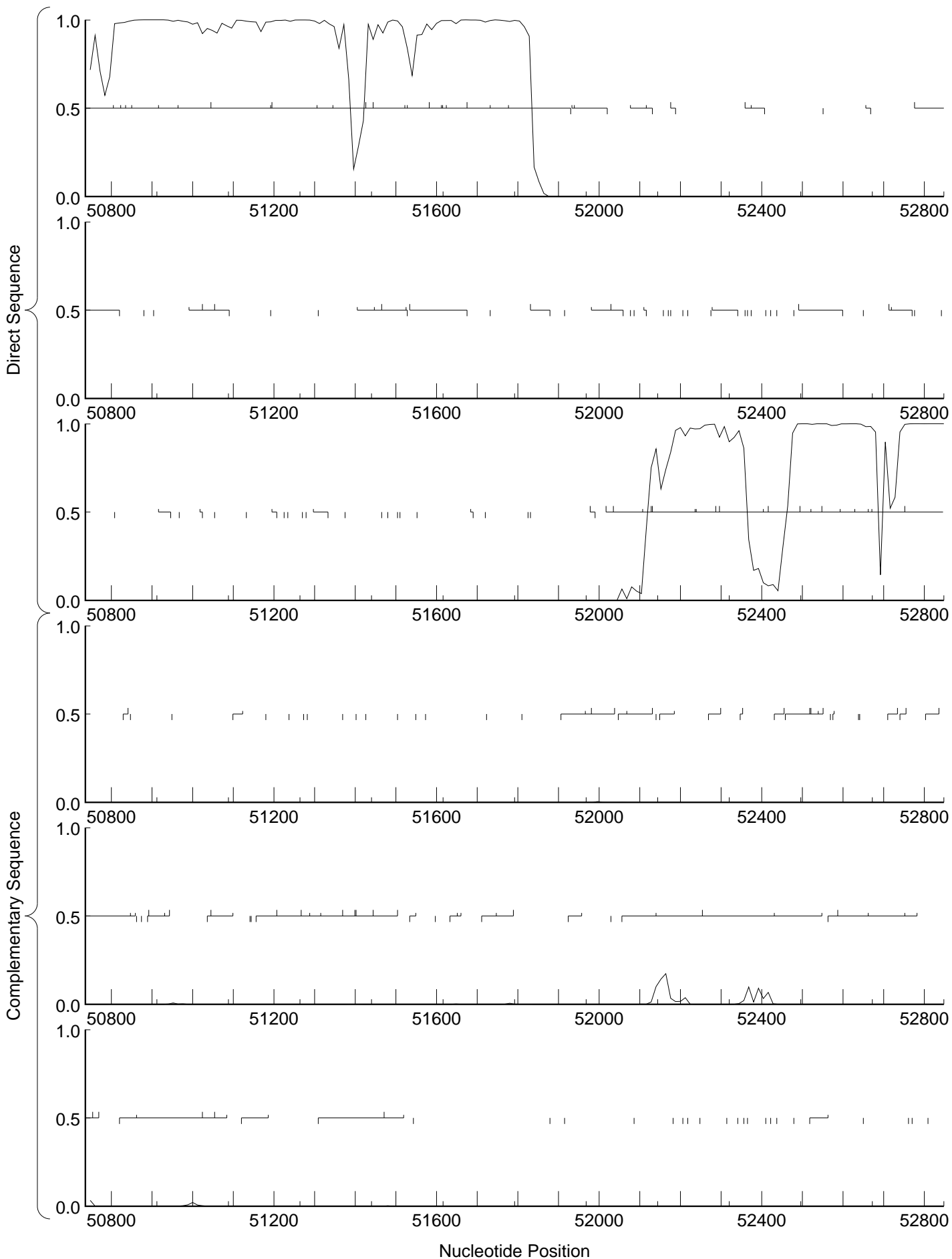


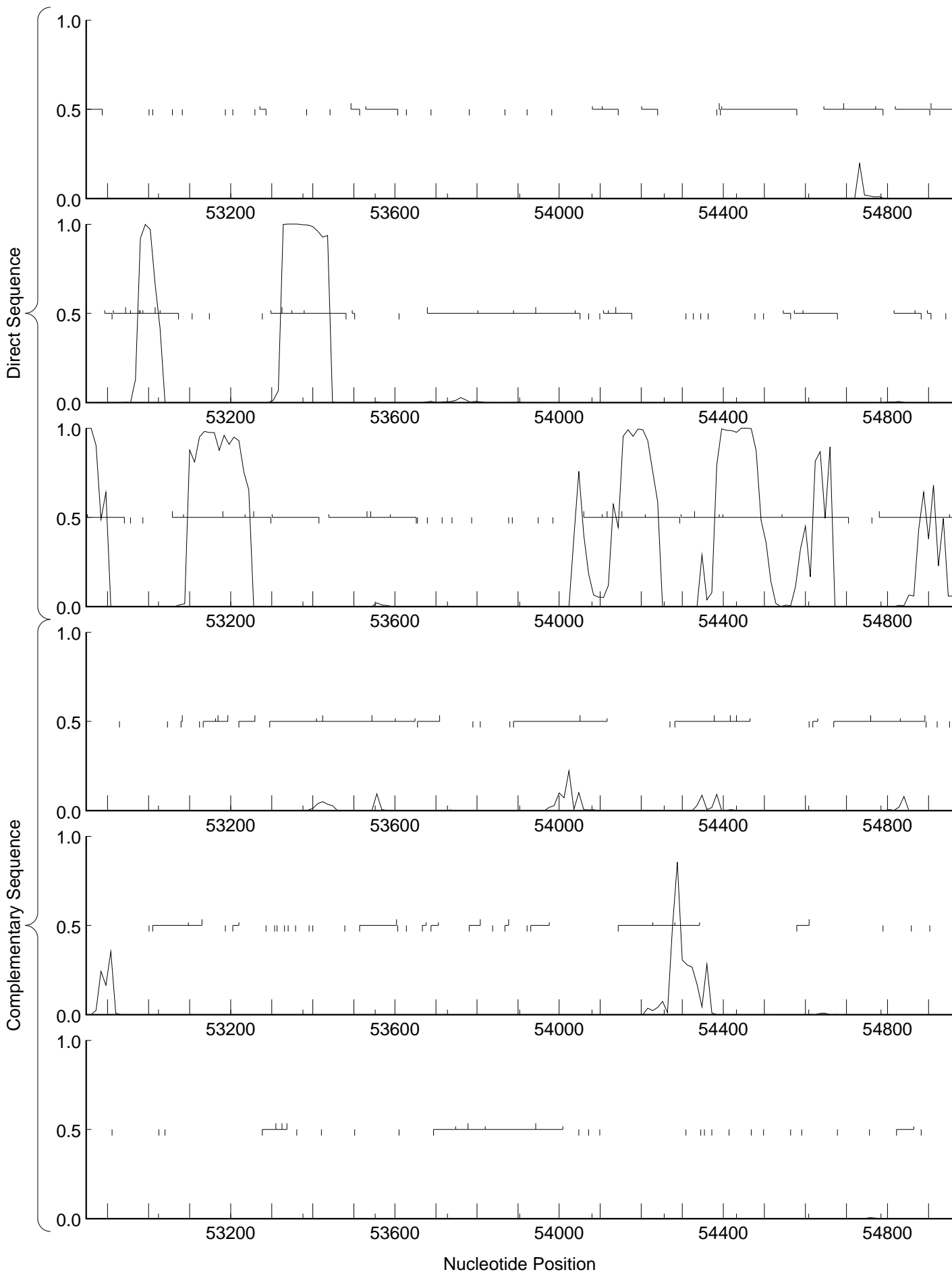


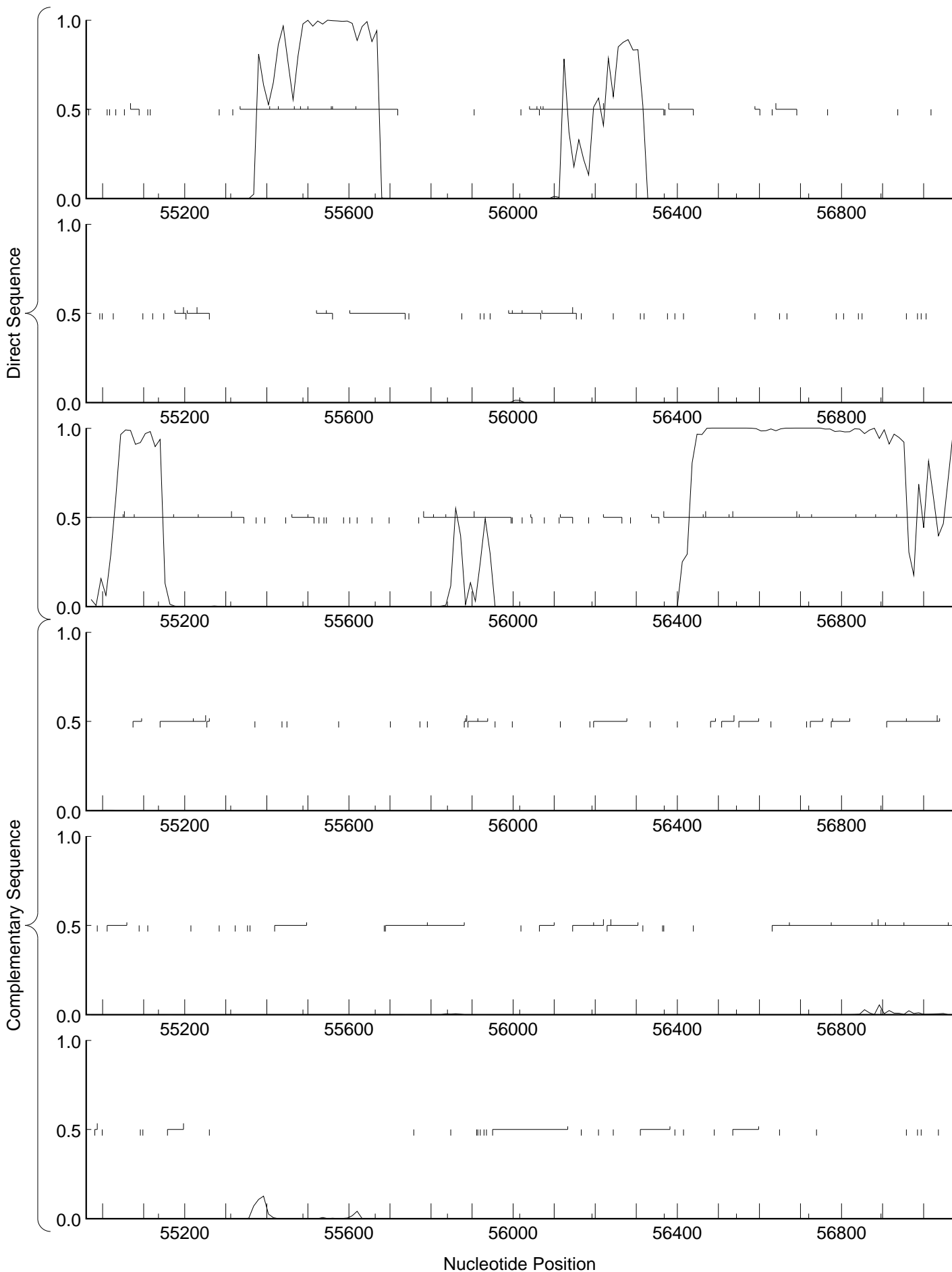


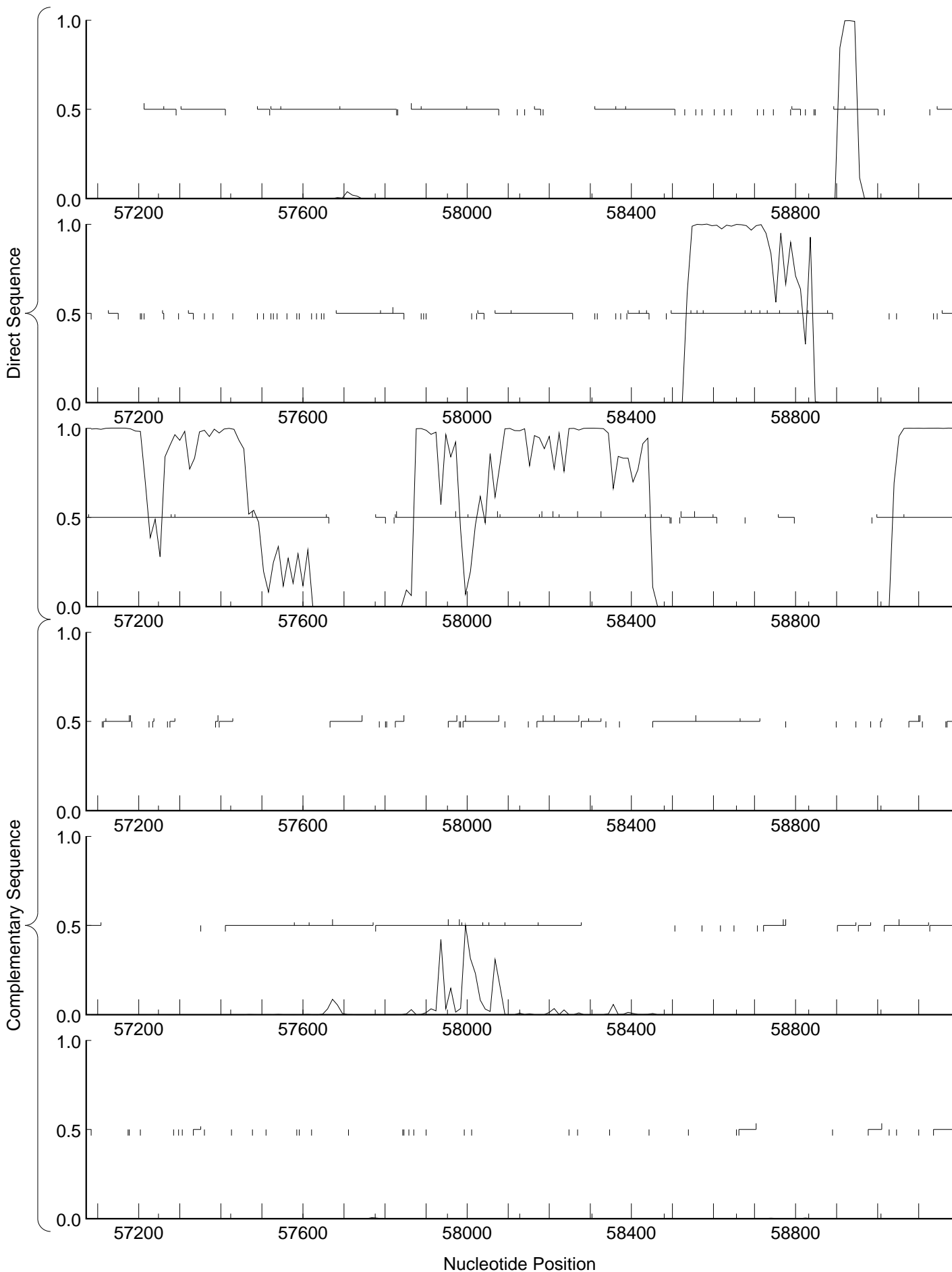




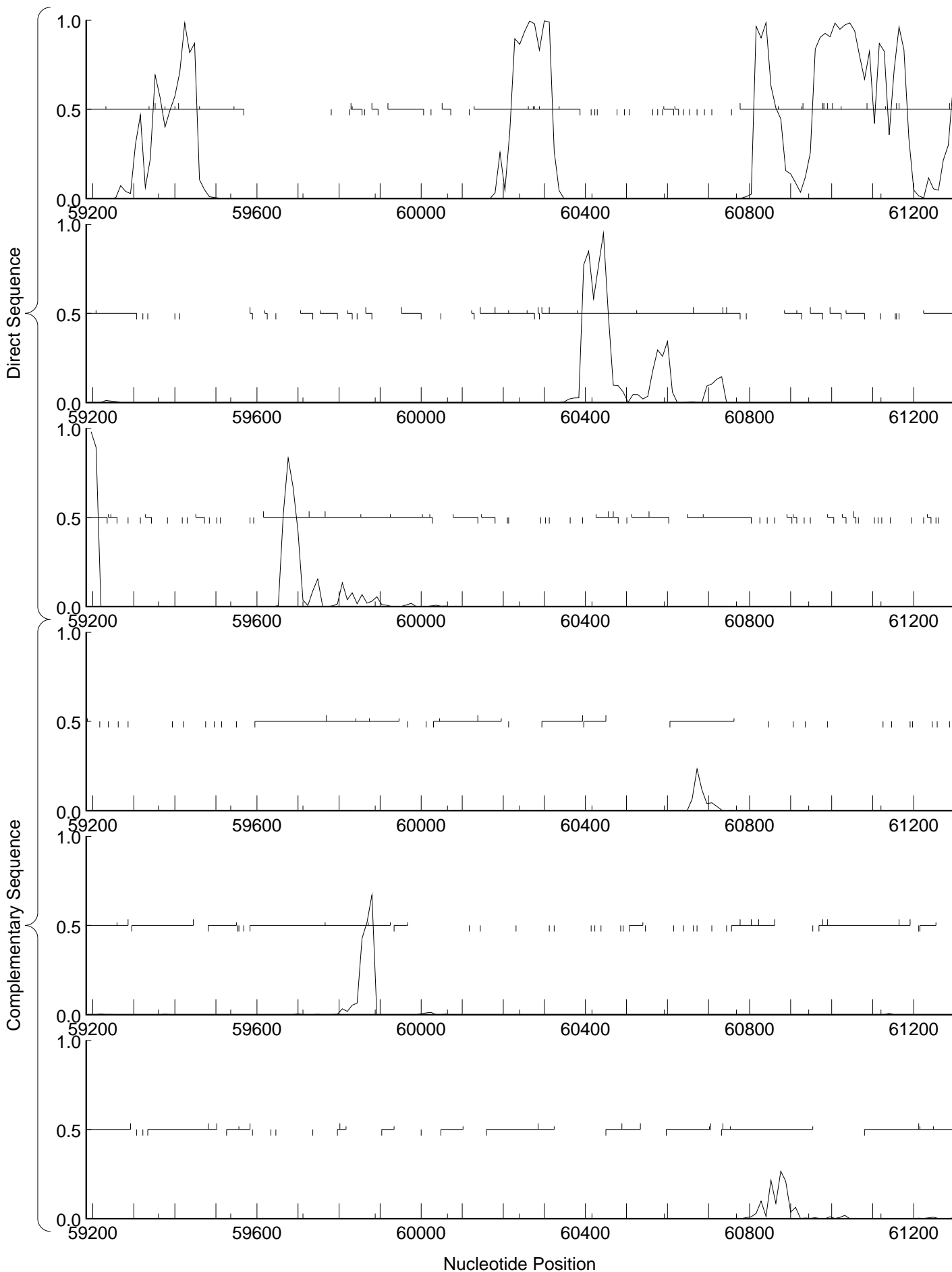


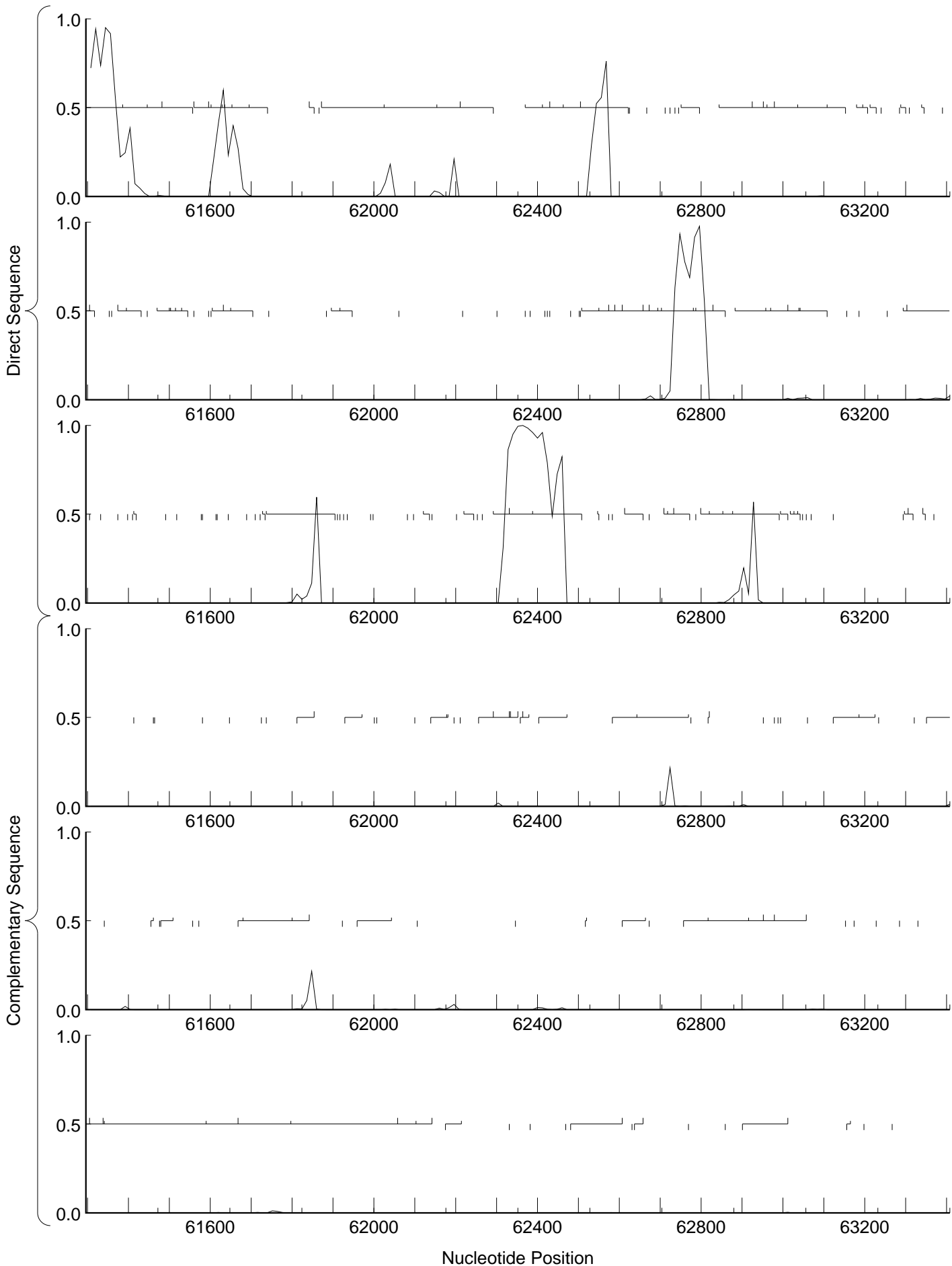






the complete sequence, 76492 bp including 10-base 3' overhang (TCGATCAGCC), Cluster L2, Order 4, Window 96, Step 12, 30/38





the complete sequence, 76492 bp including 10-base 3' overhang (TCGATCAGCC), Cluster L2, Order 4, Window 96, Step 12, 32/38

