

# GeneMark

Version 2.5p (09.08.06)

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## PROGRAM INFORMATION

Sequence : Mycobacterium phage AugsMagnumOpus complete sequence, 48545 bp including 10-base 3' overhang  
Analysis Date : 9/17/19 at 11:06:47  
Pages : 24  
Sequence Length : 48545 bp  
GC Content : 64.17%

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,

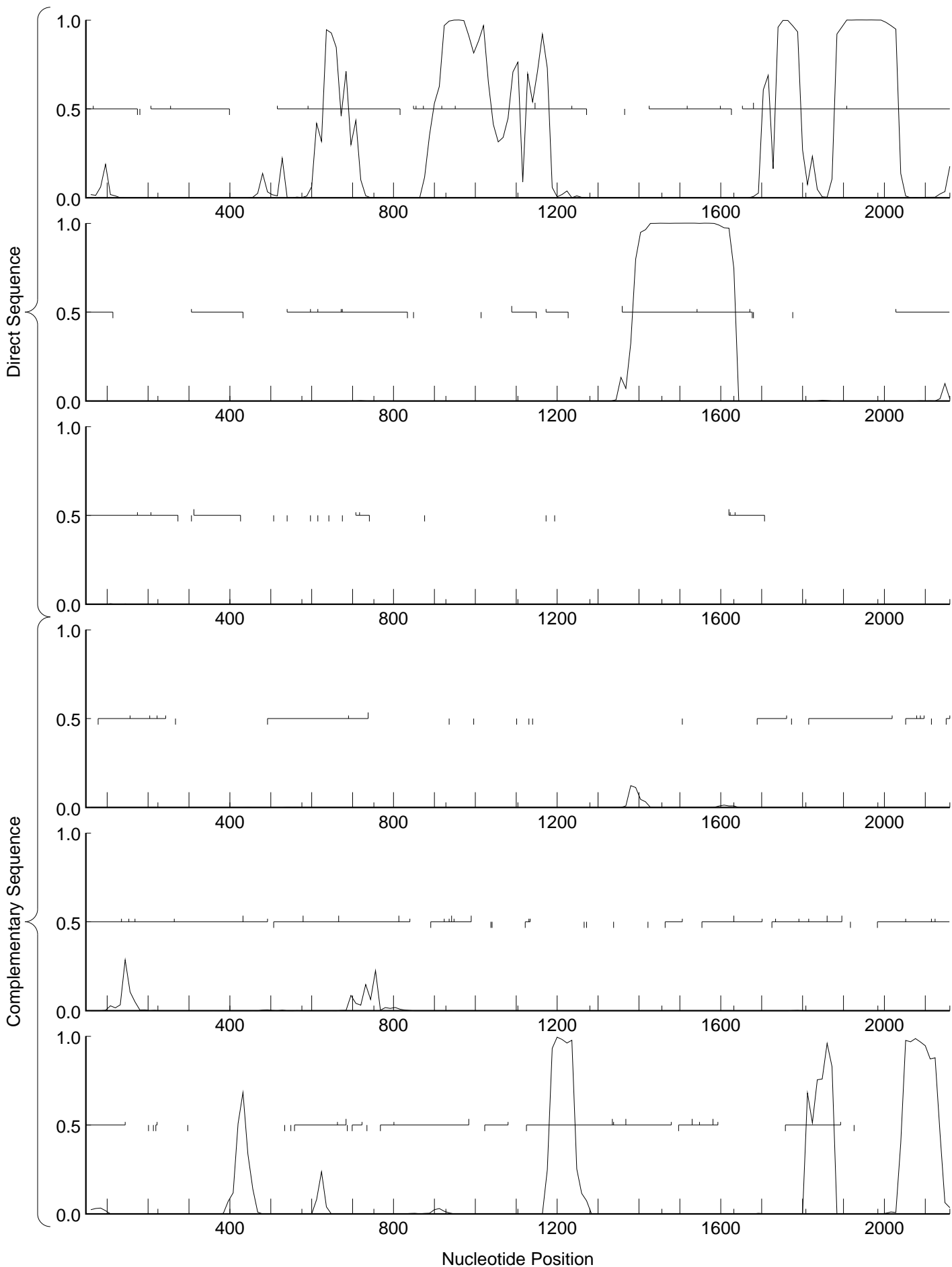
## Matrix notes & comments

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

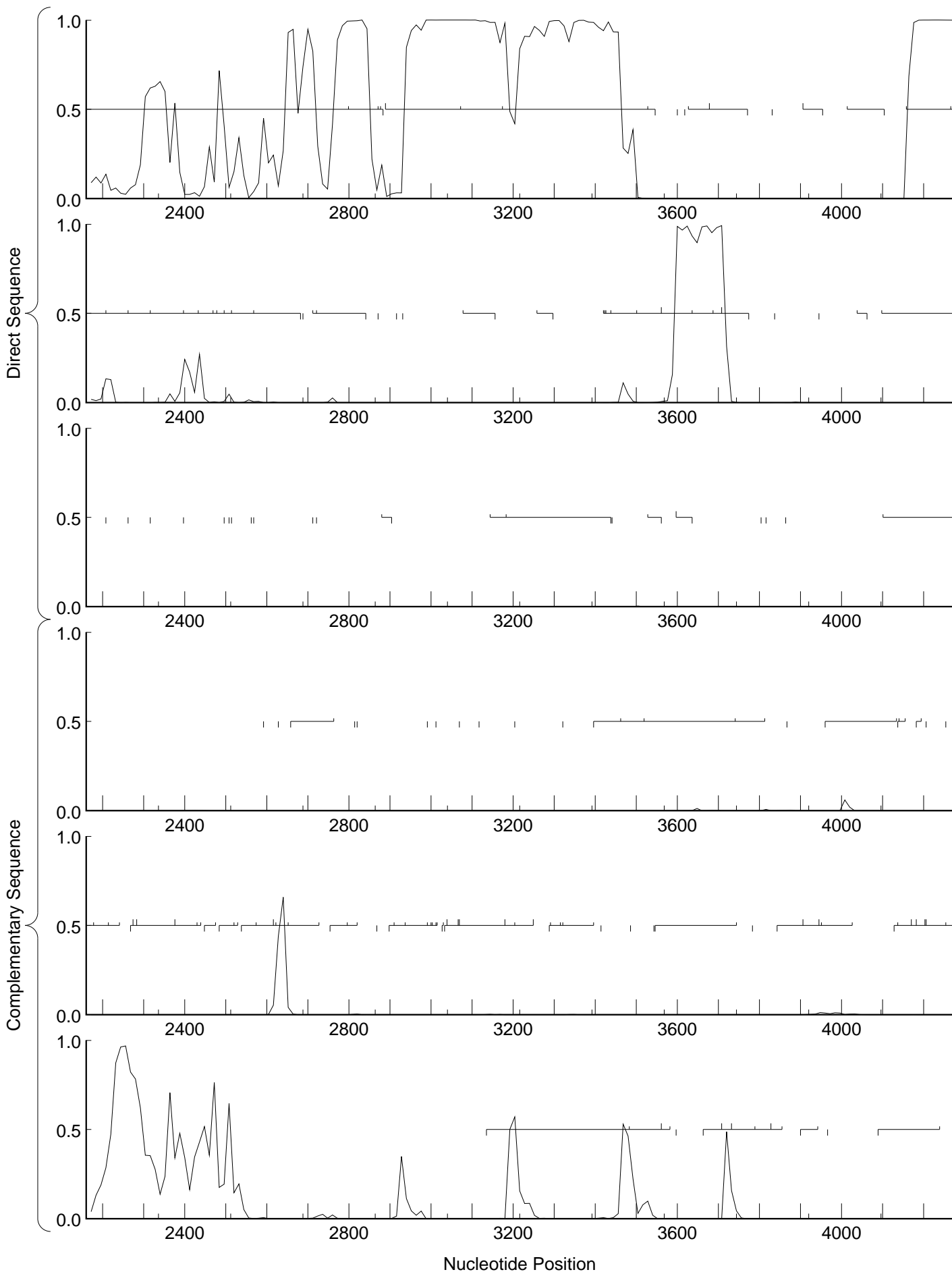
## MATRIX INFORMATION

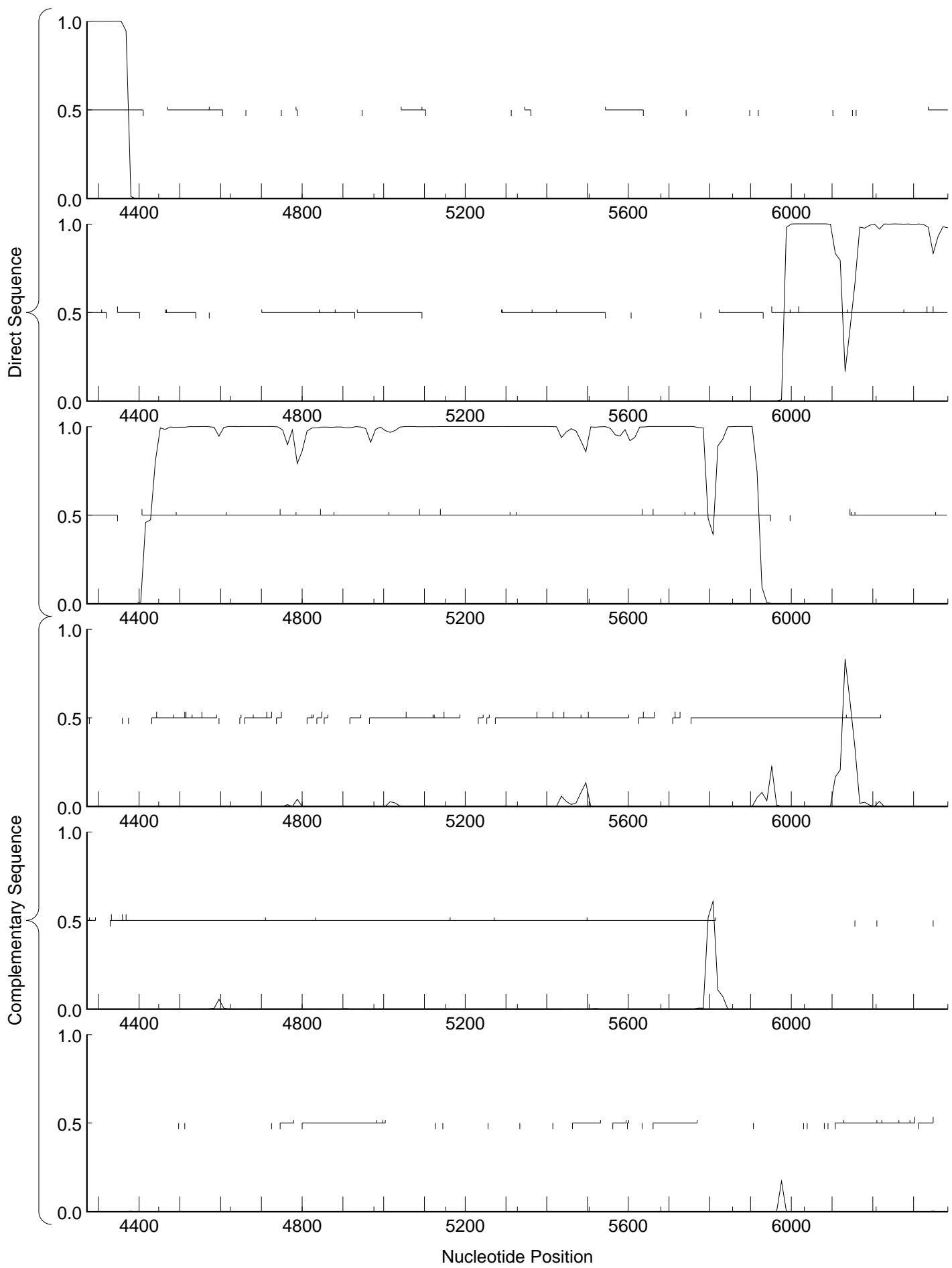
Matrix : Mycobacterium\_tuberculosis\_H37Rv  
Author : -  
Order : 4

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

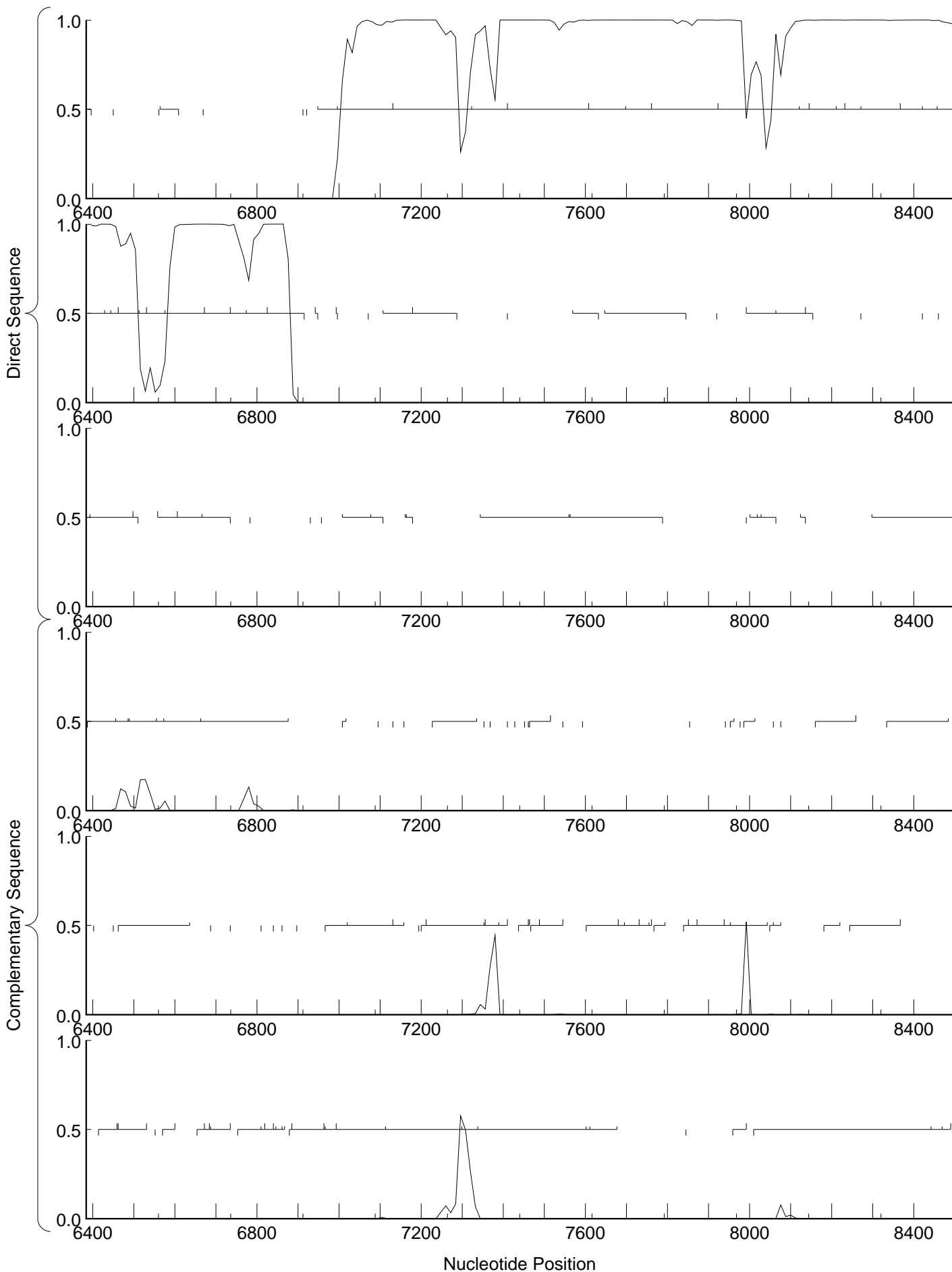


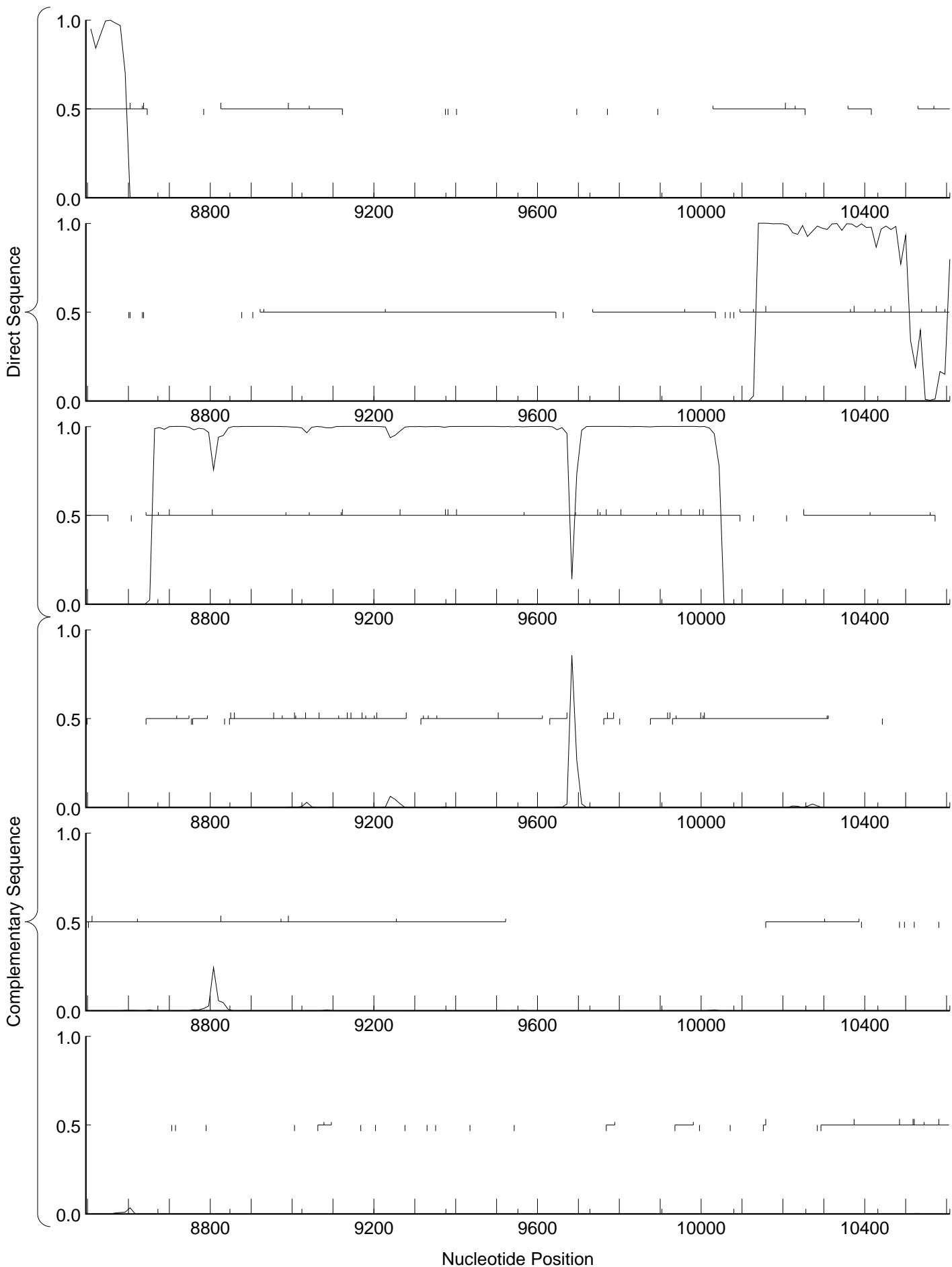
s complete sequence, 48545 bp including 10-base 3' overhang (CGGGTGGTAA), Cluster A3, Order 4, Window 96, Step 12, 3/24

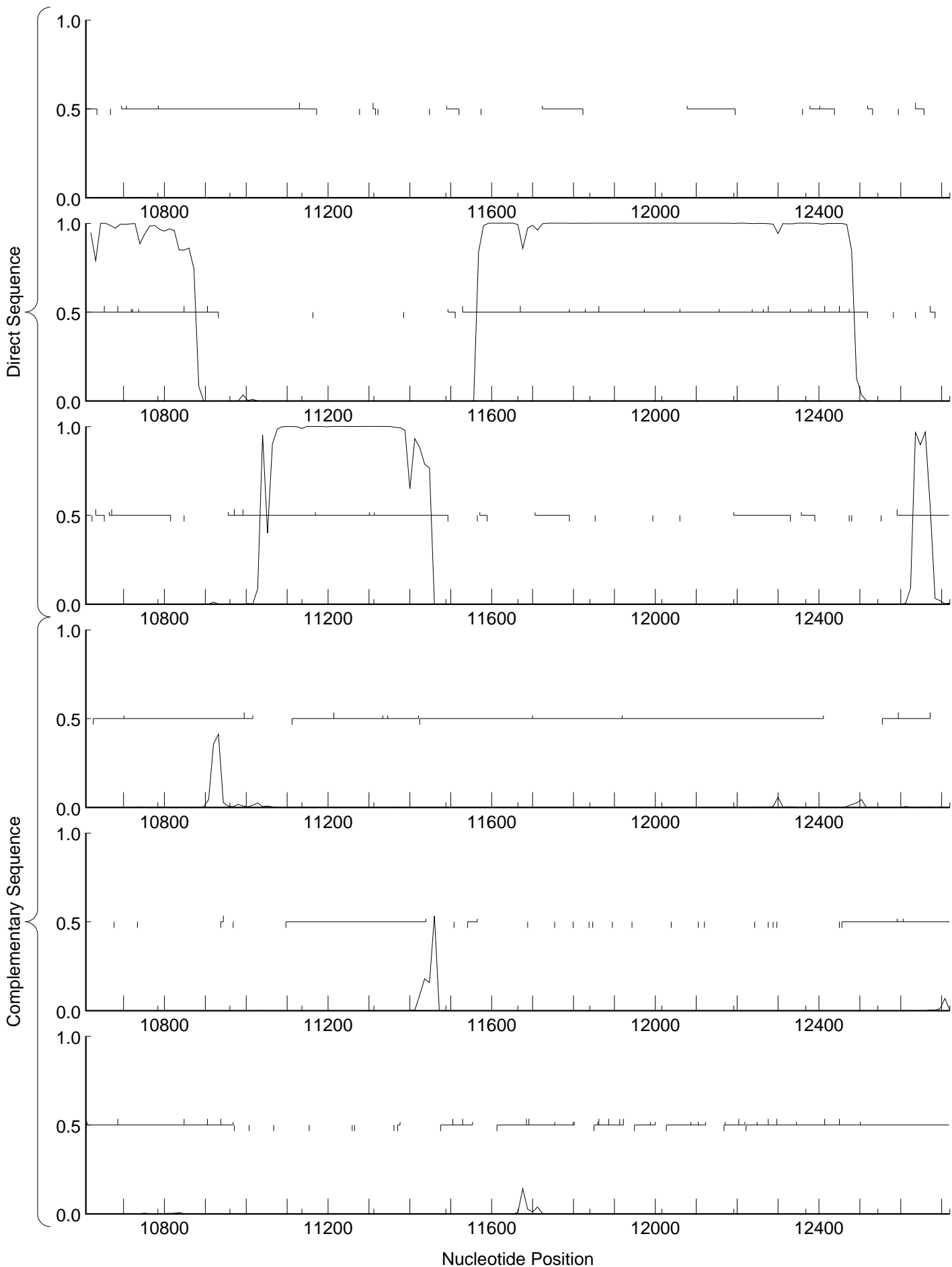


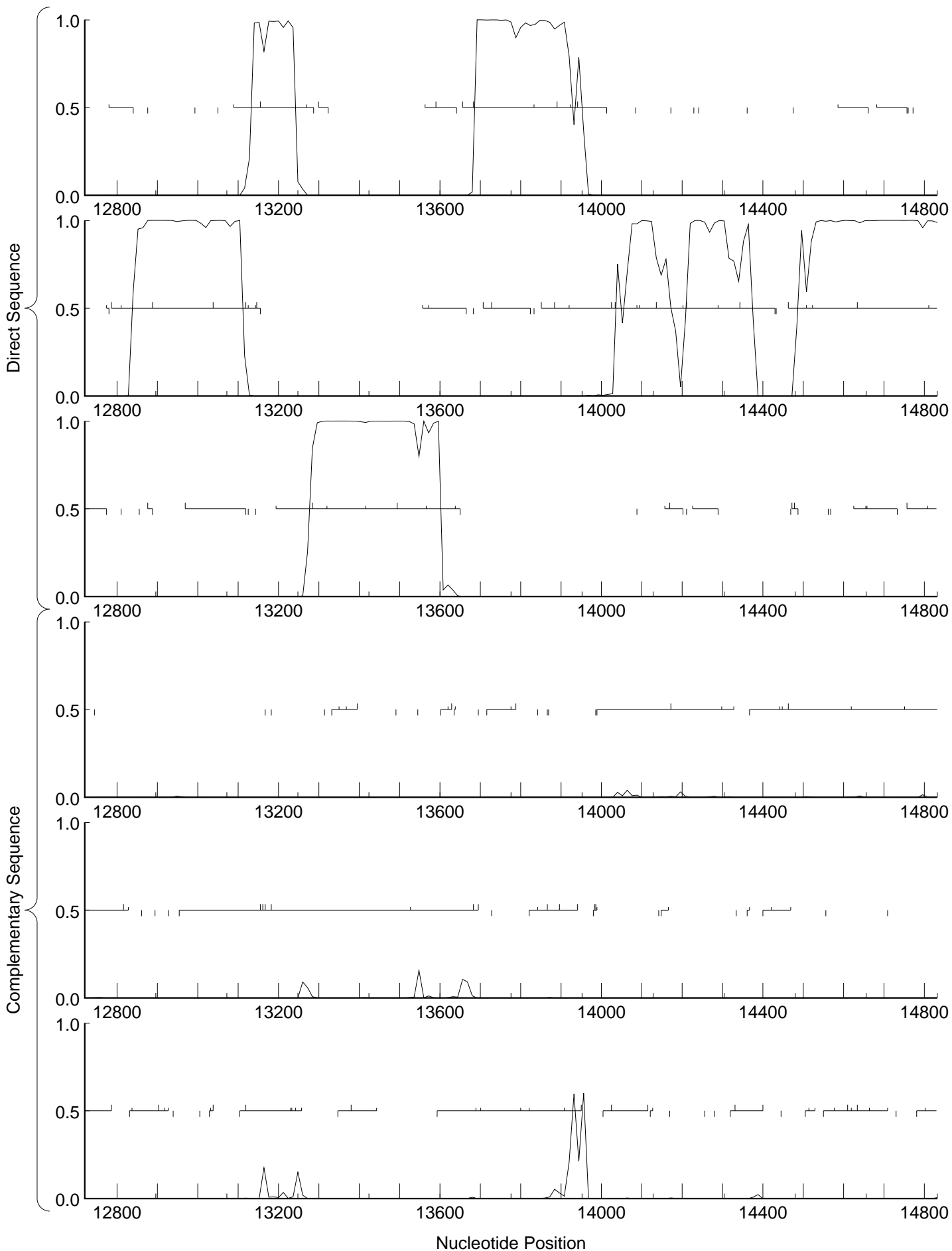


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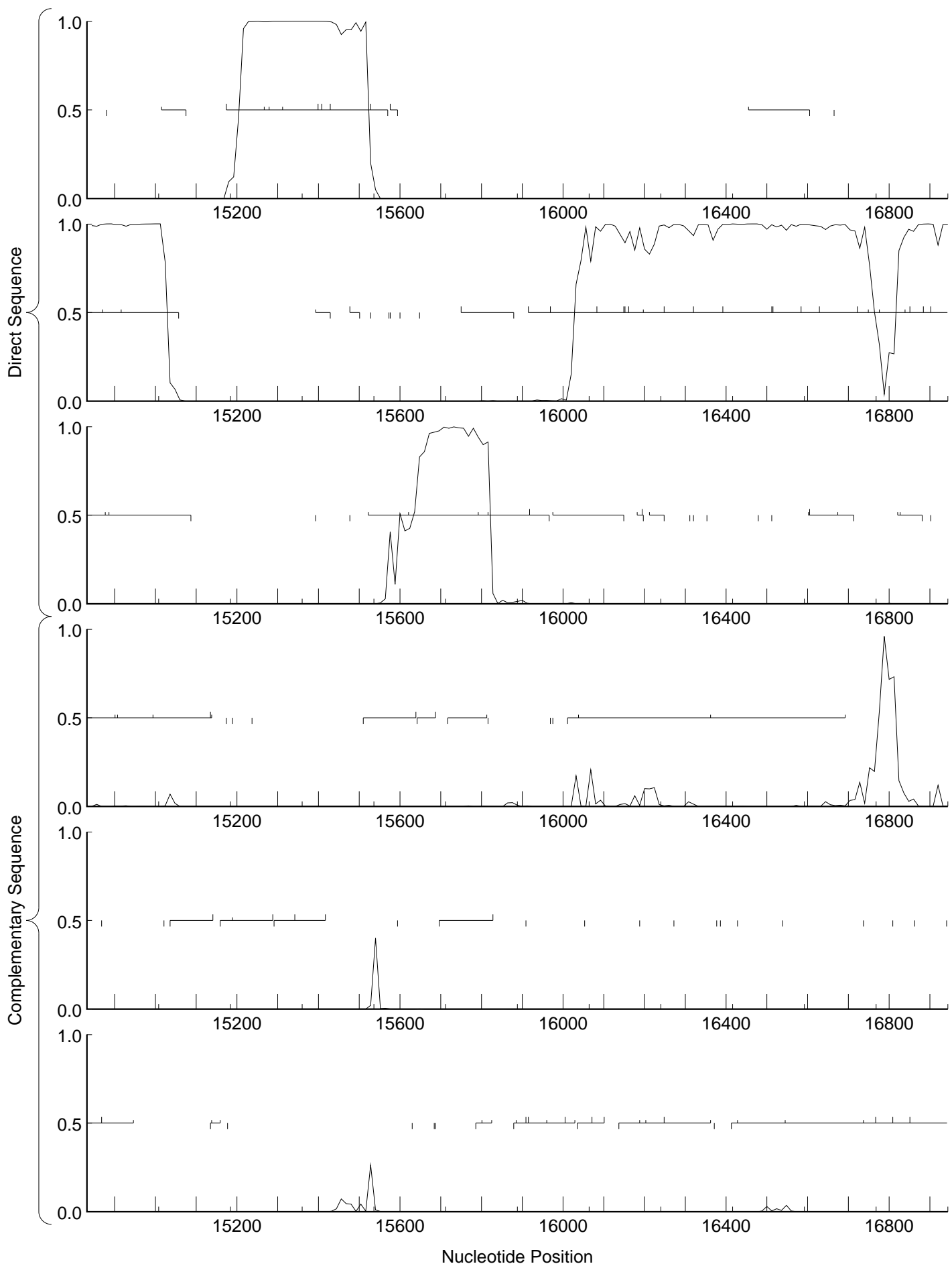


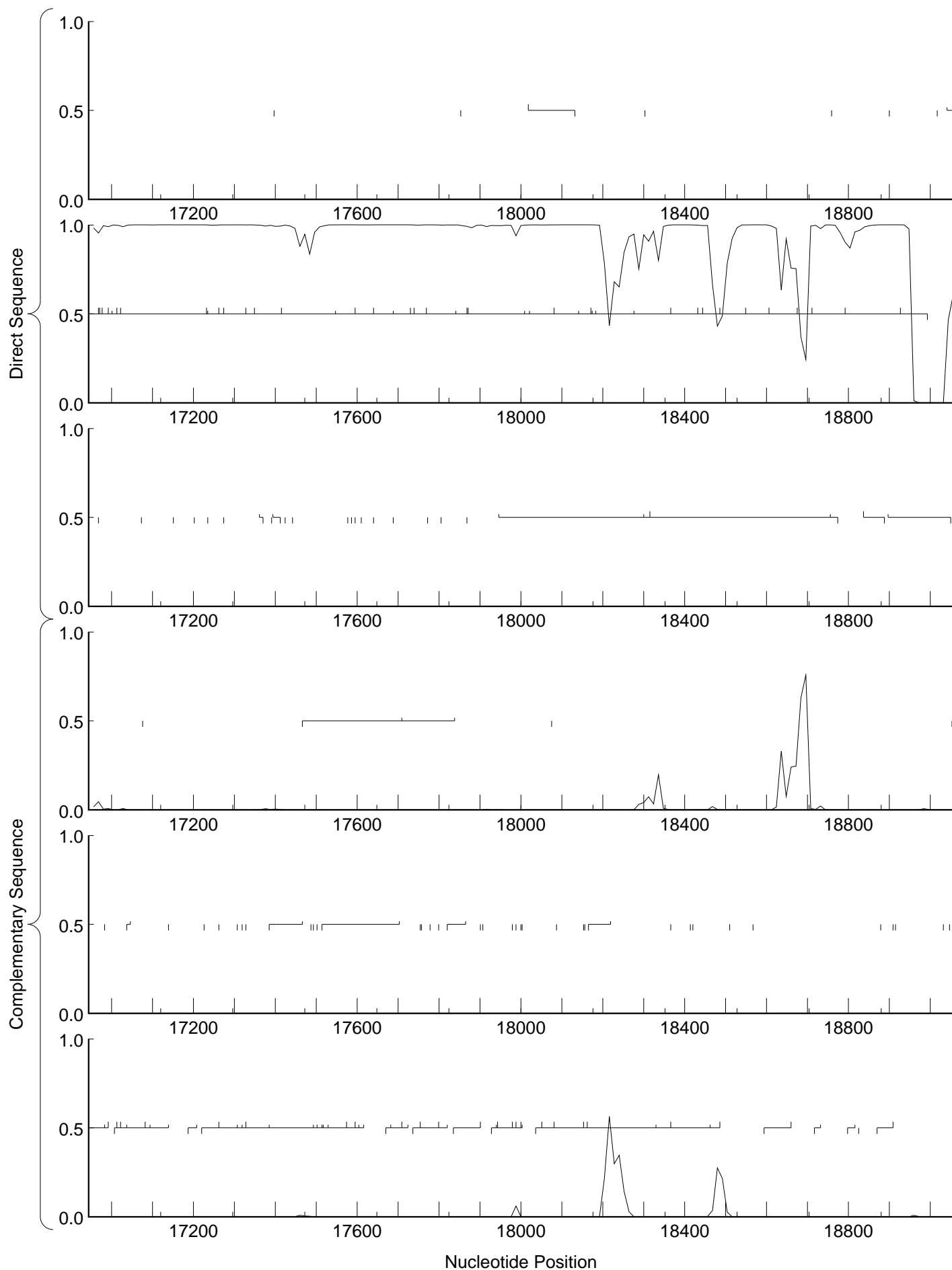




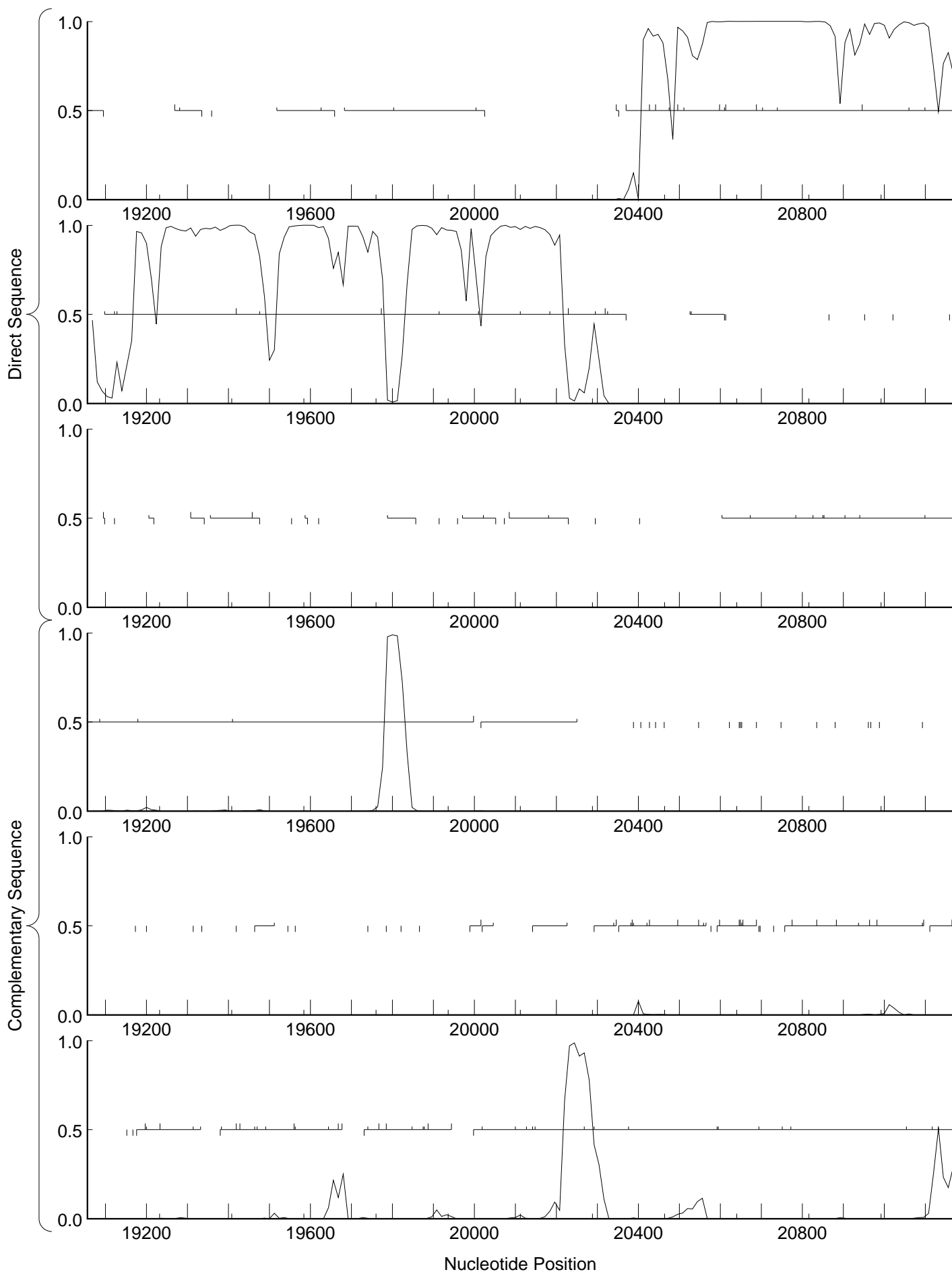


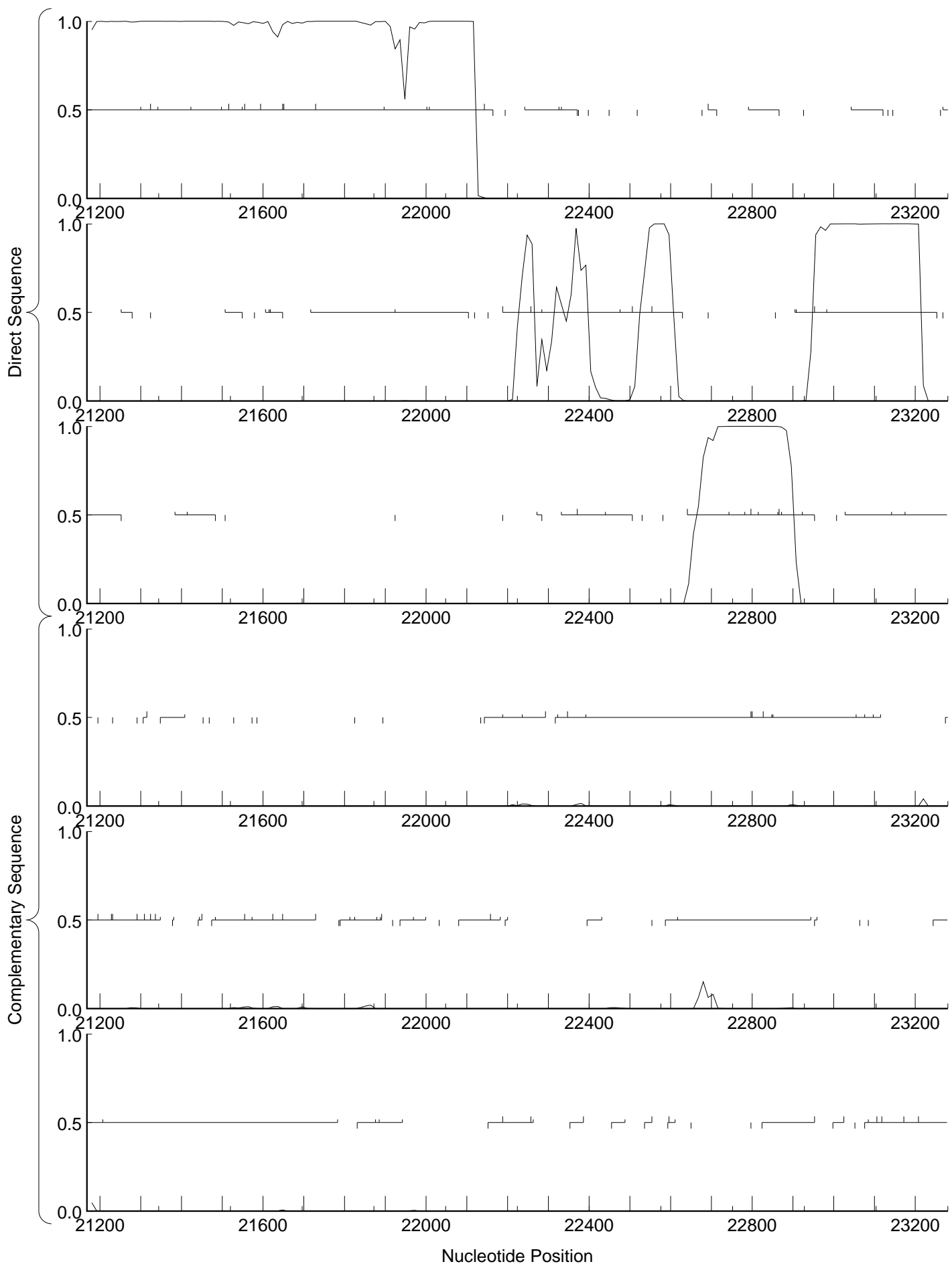




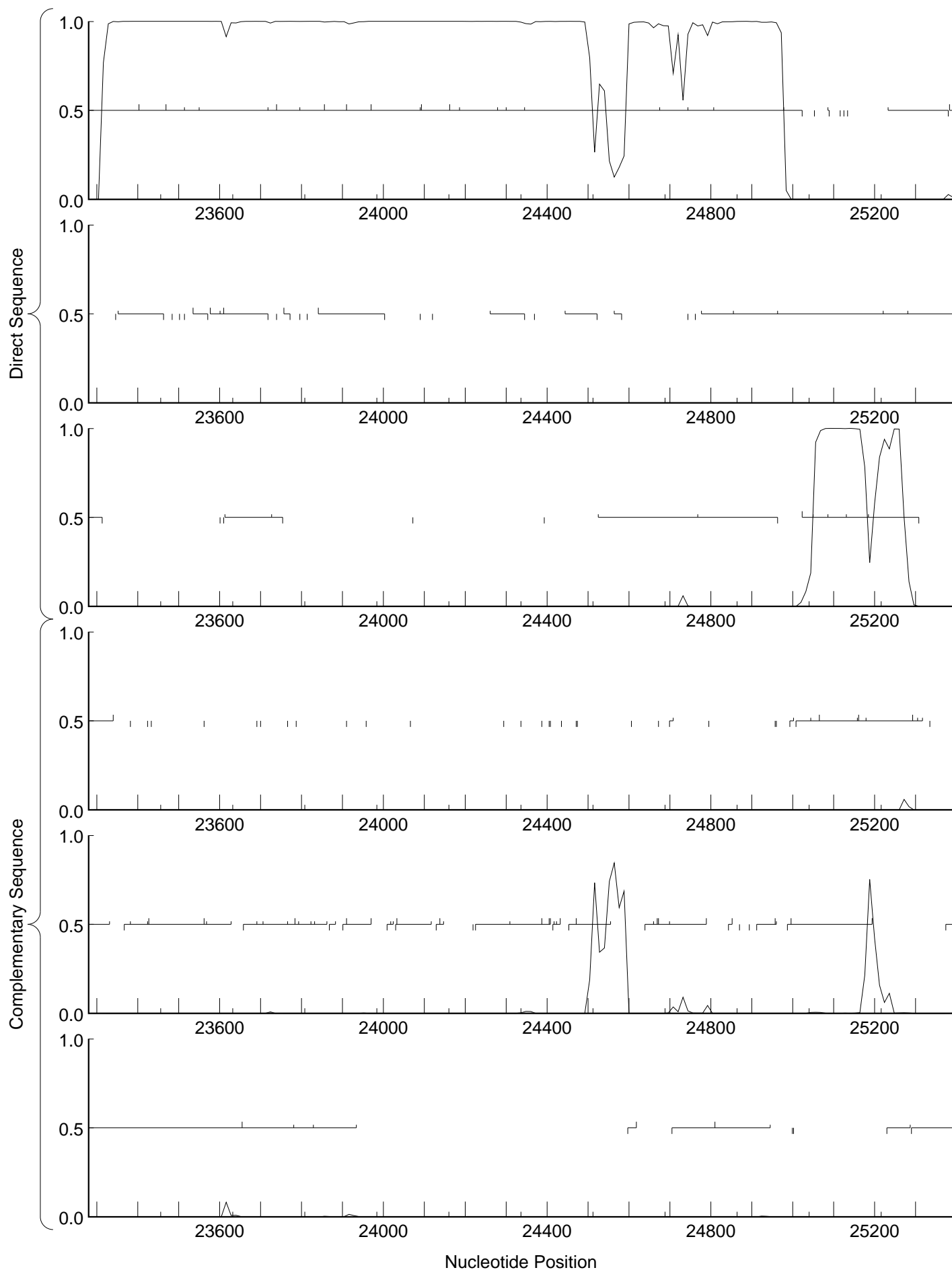


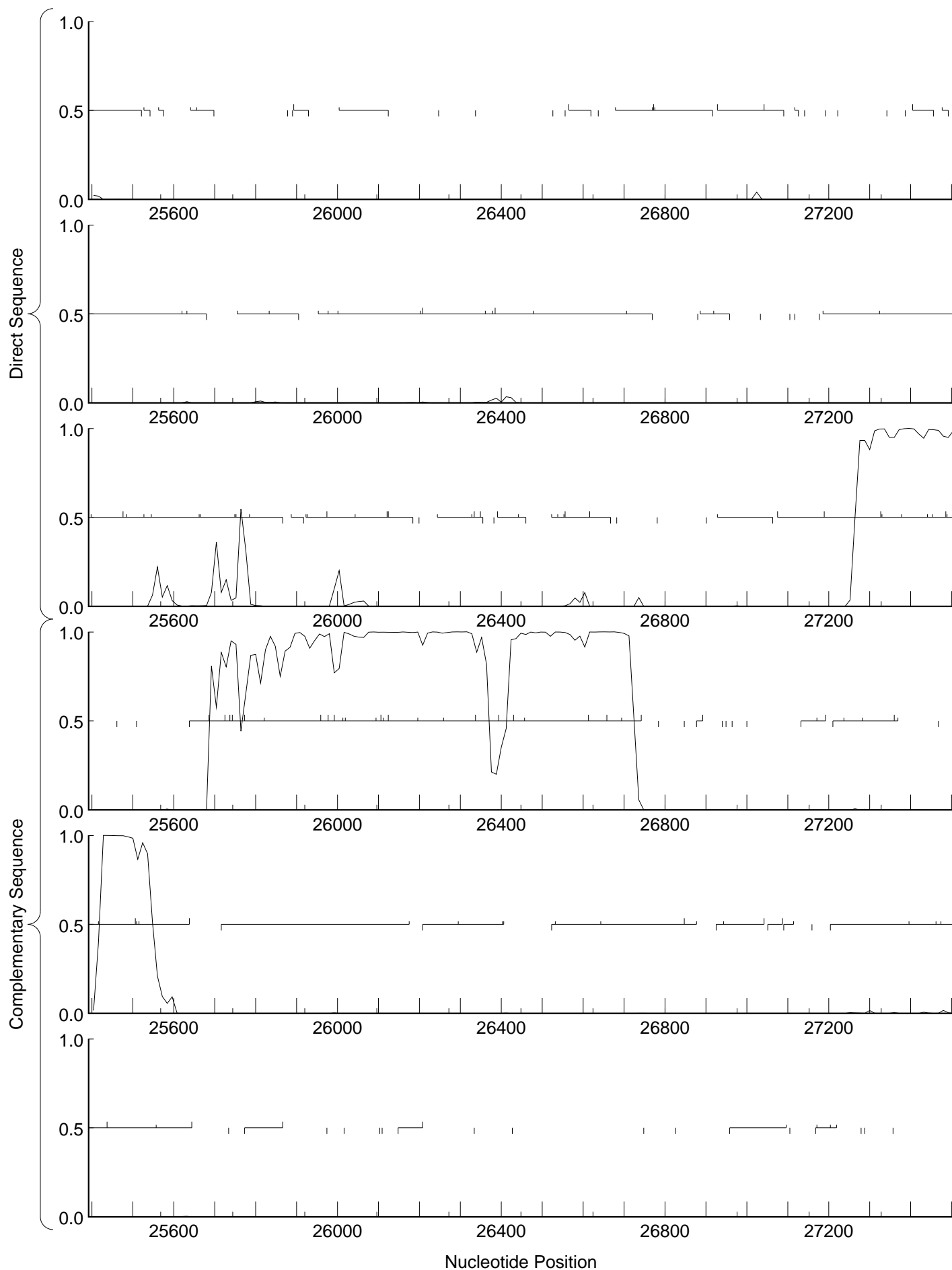
complete sequence, 48545 bp including 10-base 3' overhang (CGGGTGGTAA), Cluster A3, Order 4, Window 96, Step 12, 11/24



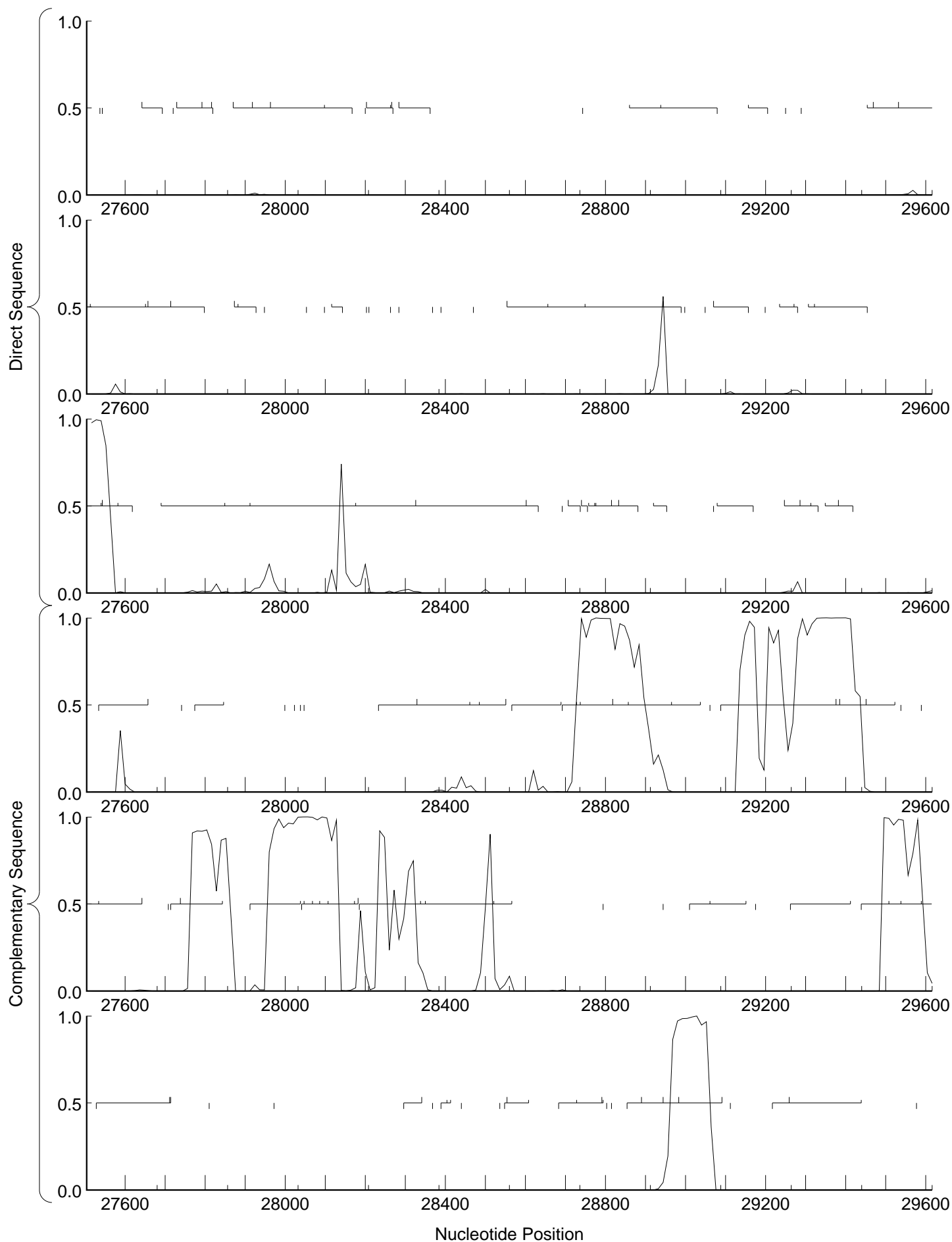


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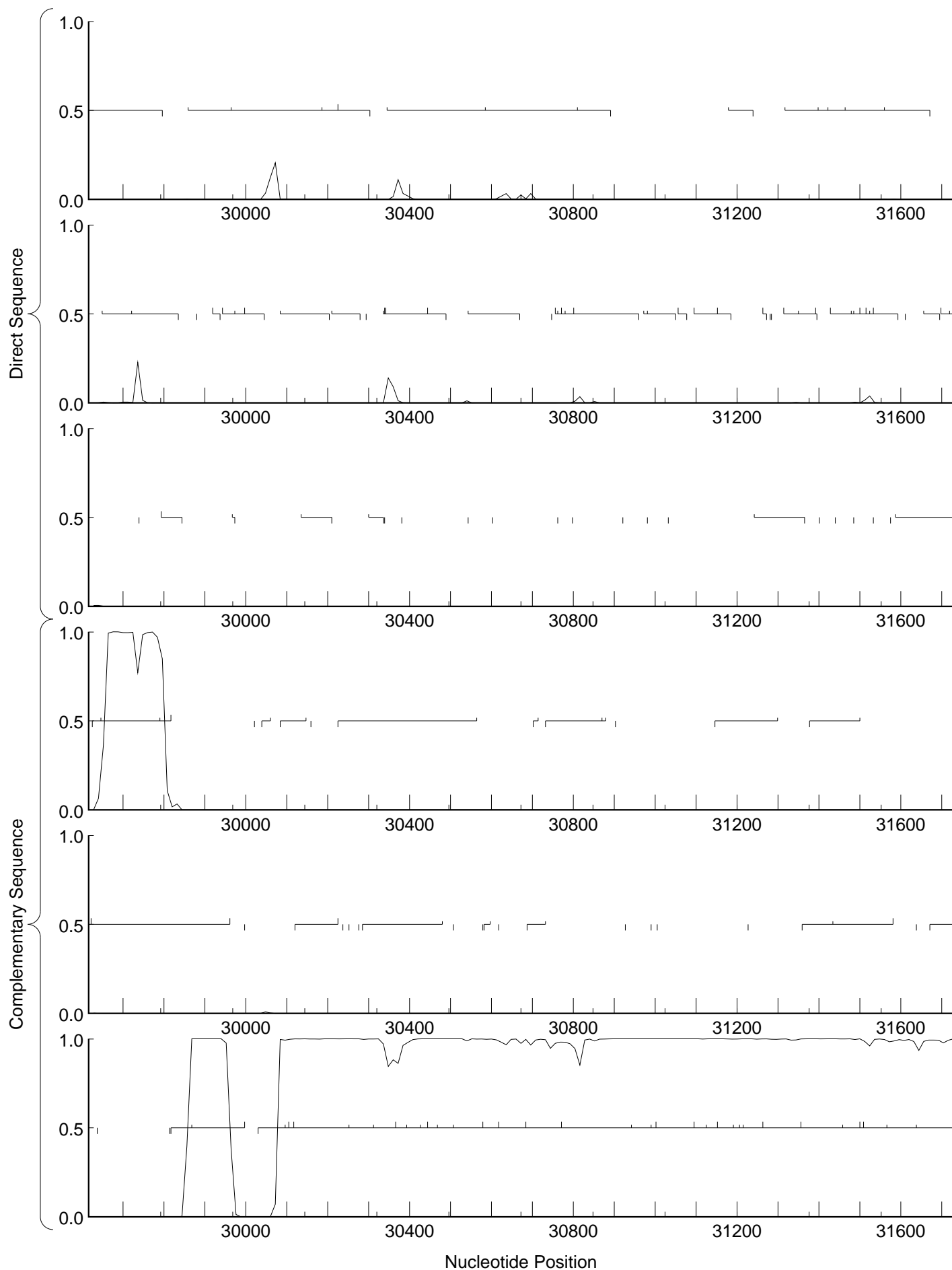




complete sequence, 48545 bp including 10-base 3' overhang (CGGGTGGTAA), Cluster A3, Order 4, Window 96, Step 12, 15/24

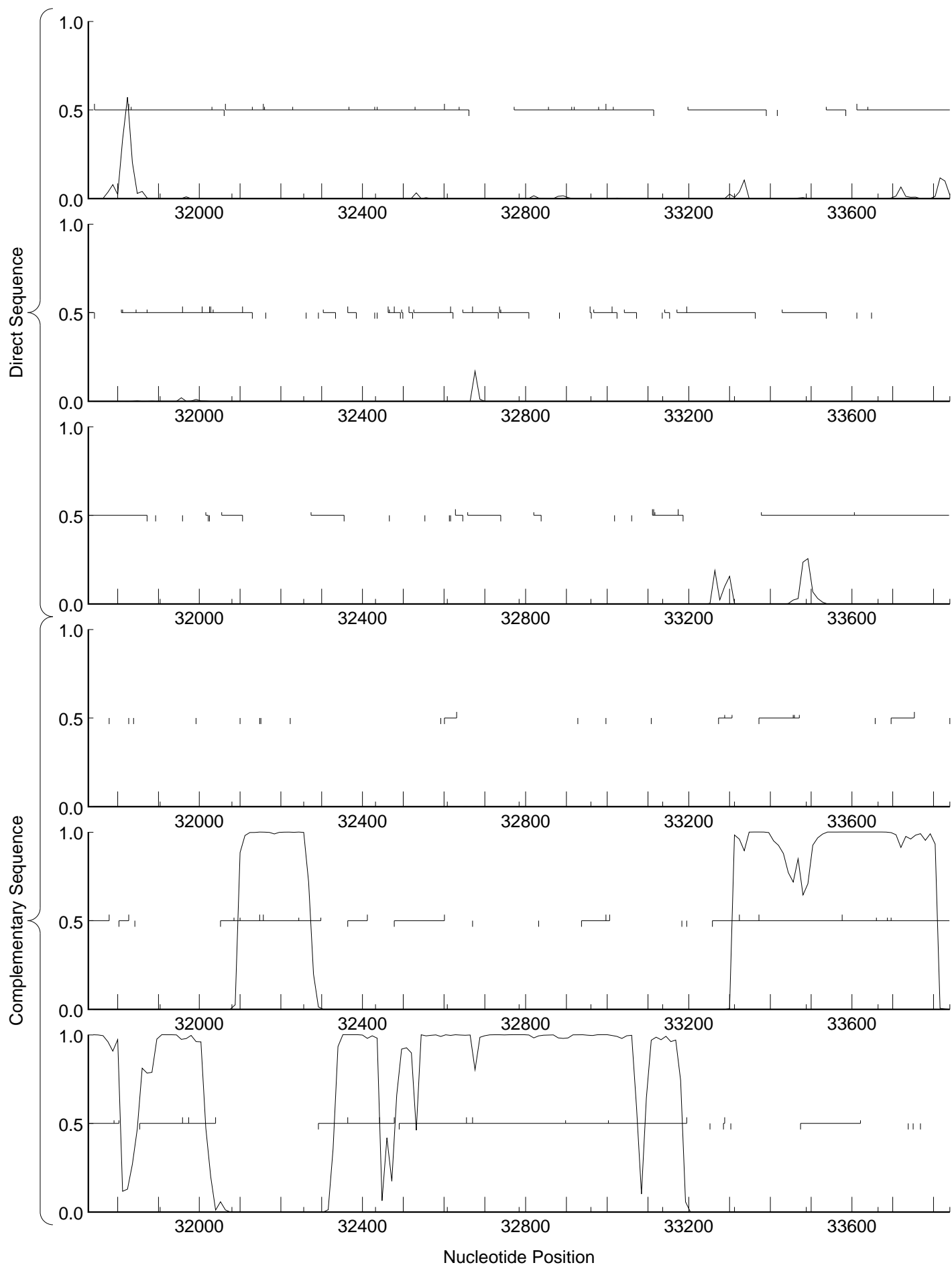


complete sequence, 48545 bp including 10-base 3' overhang (CGGGTGGTAA), Cluster A3, Order 4, Window 96, Step 12, 16/24

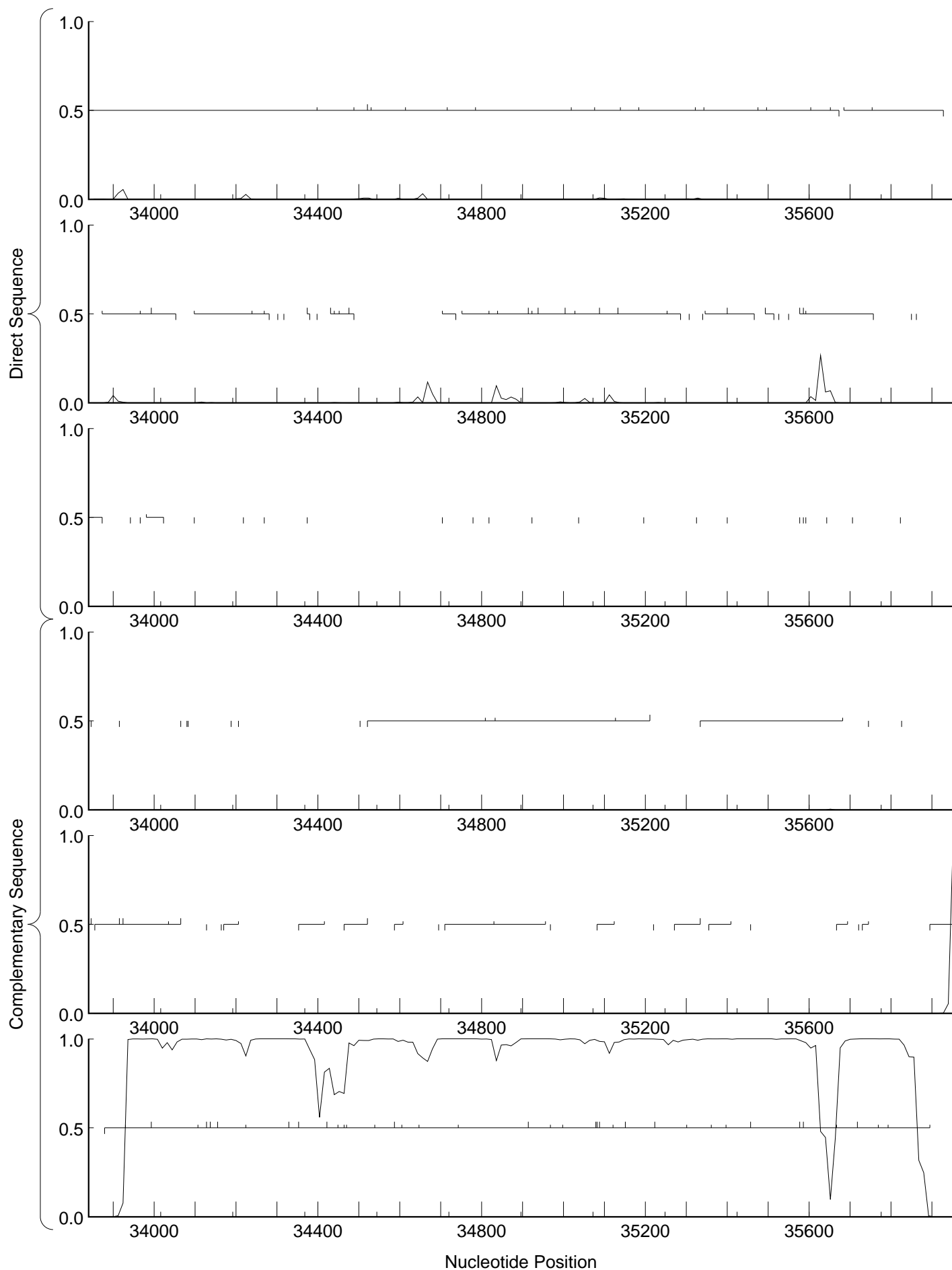




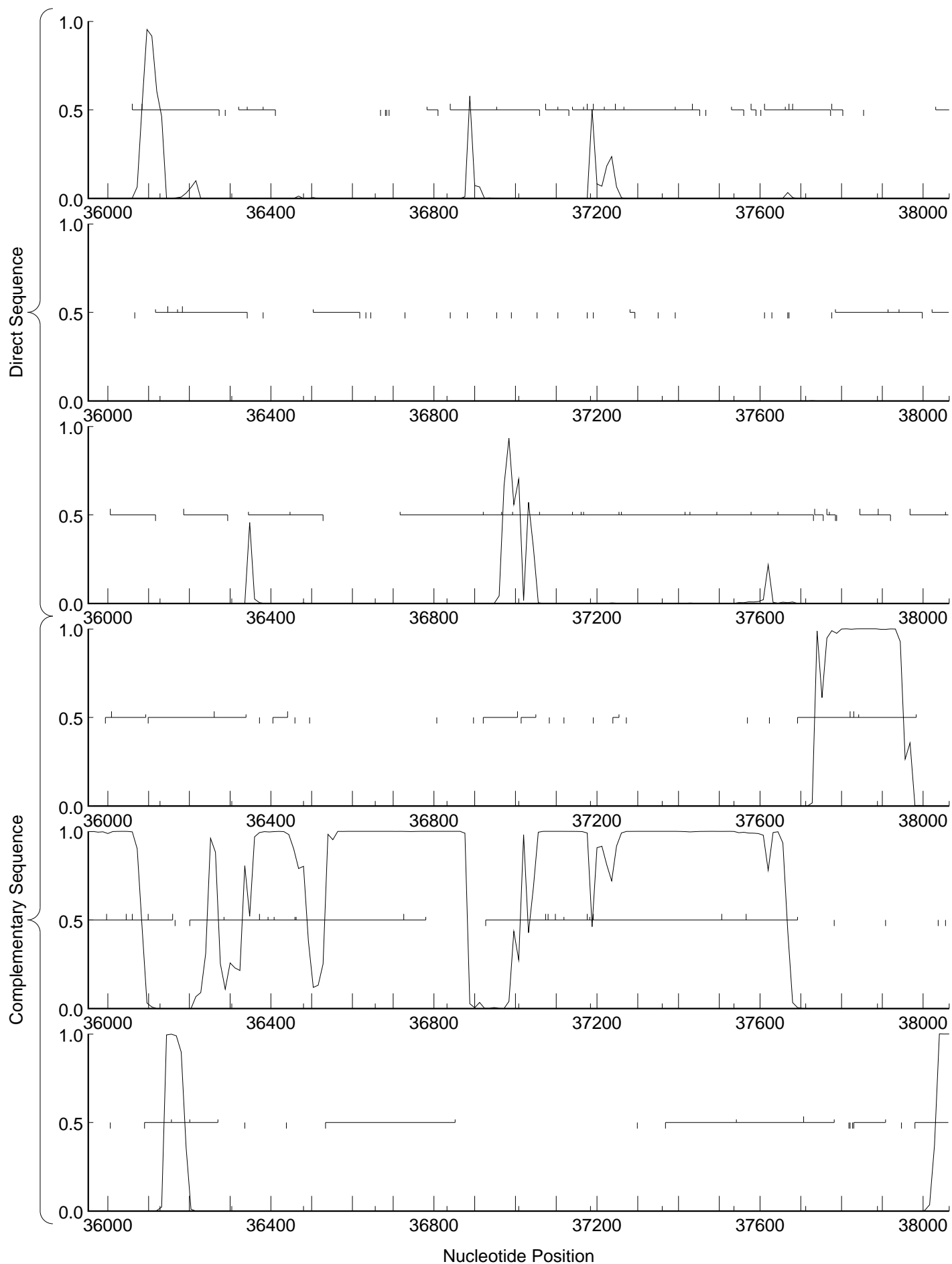
complete sequence, 48545 bp including 10-base 3' overhang (CGGGTGGTAA), Cluster A3, Order 4, Window 96, Step 12, 17/24

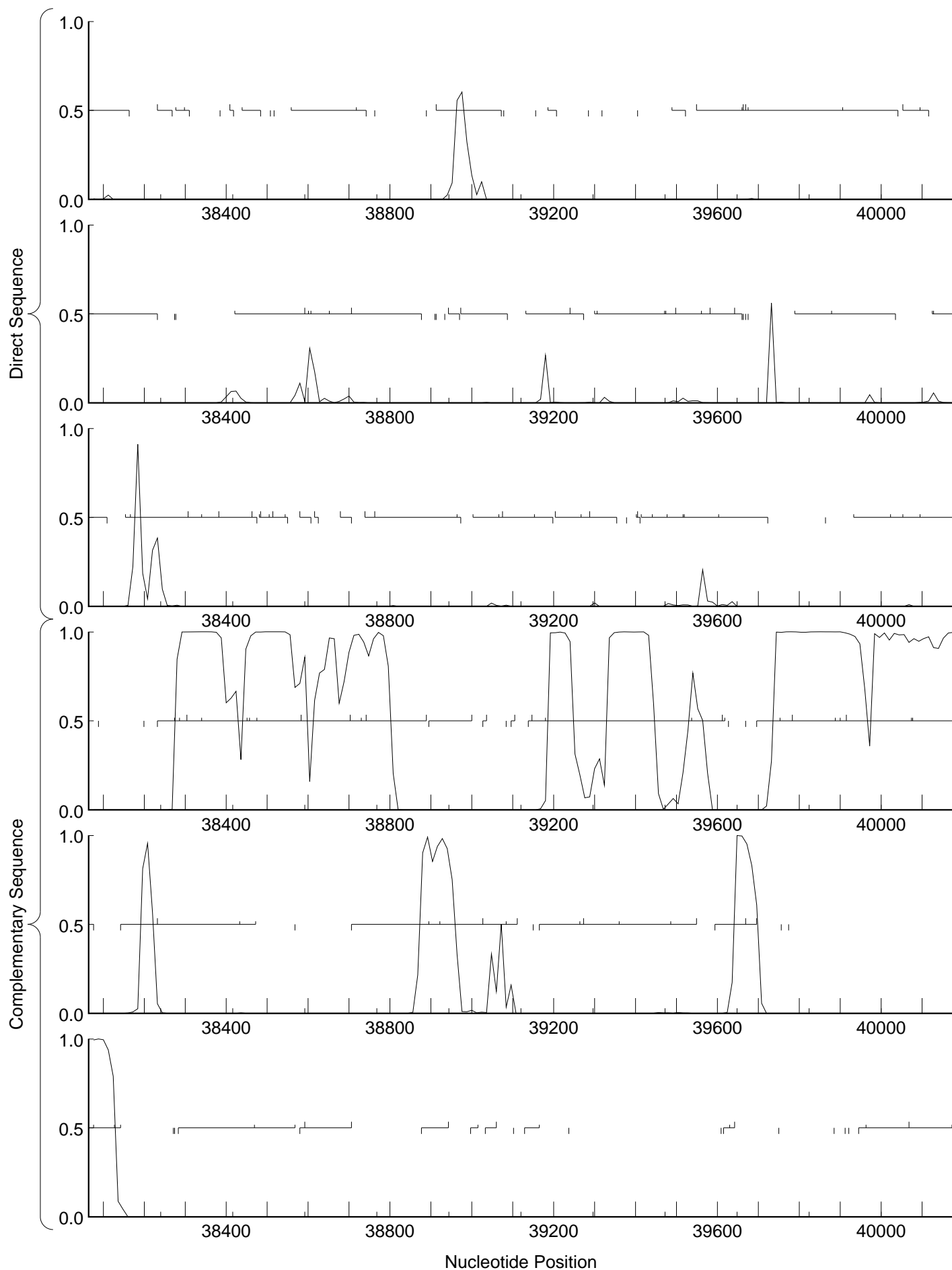


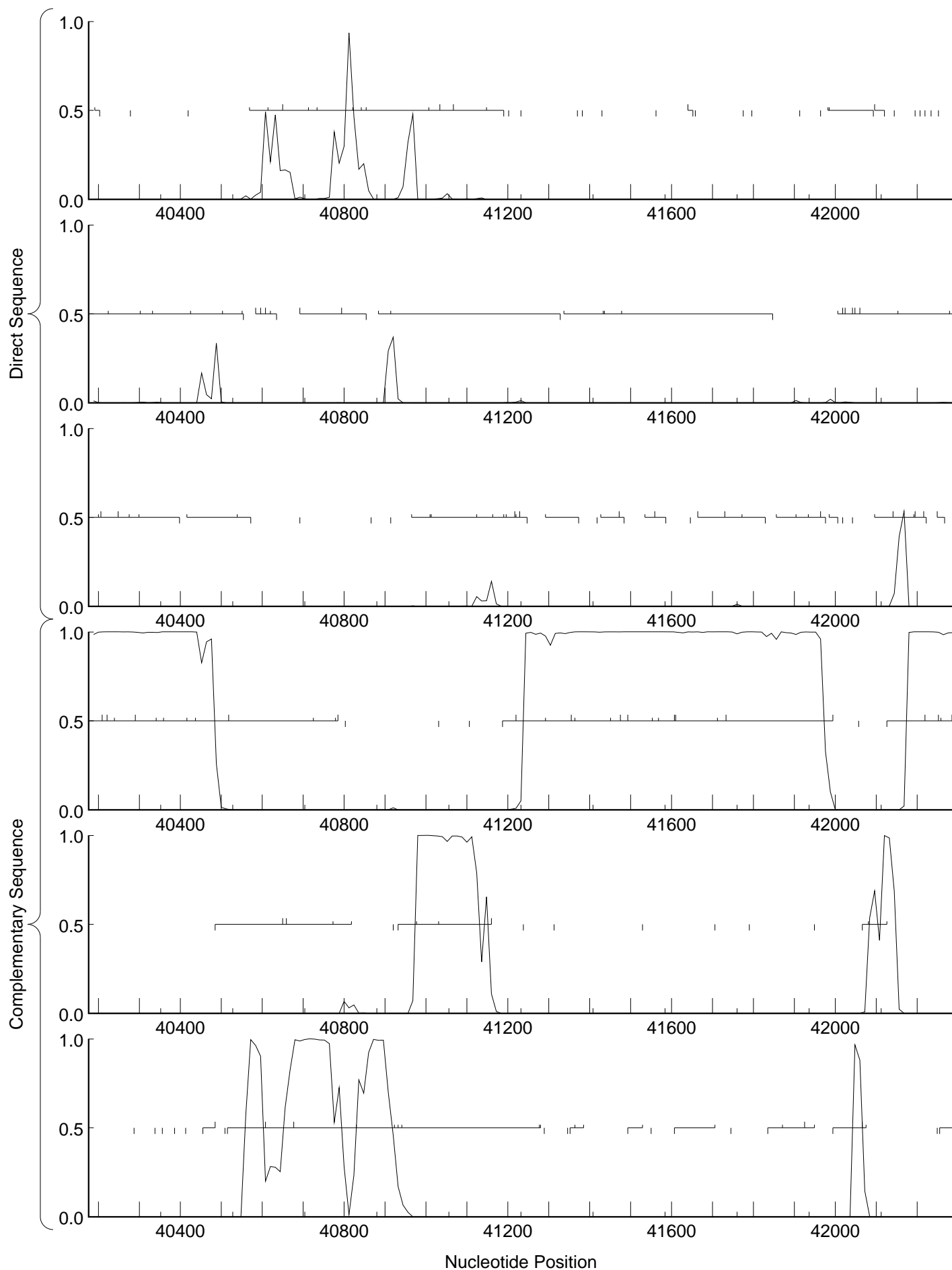
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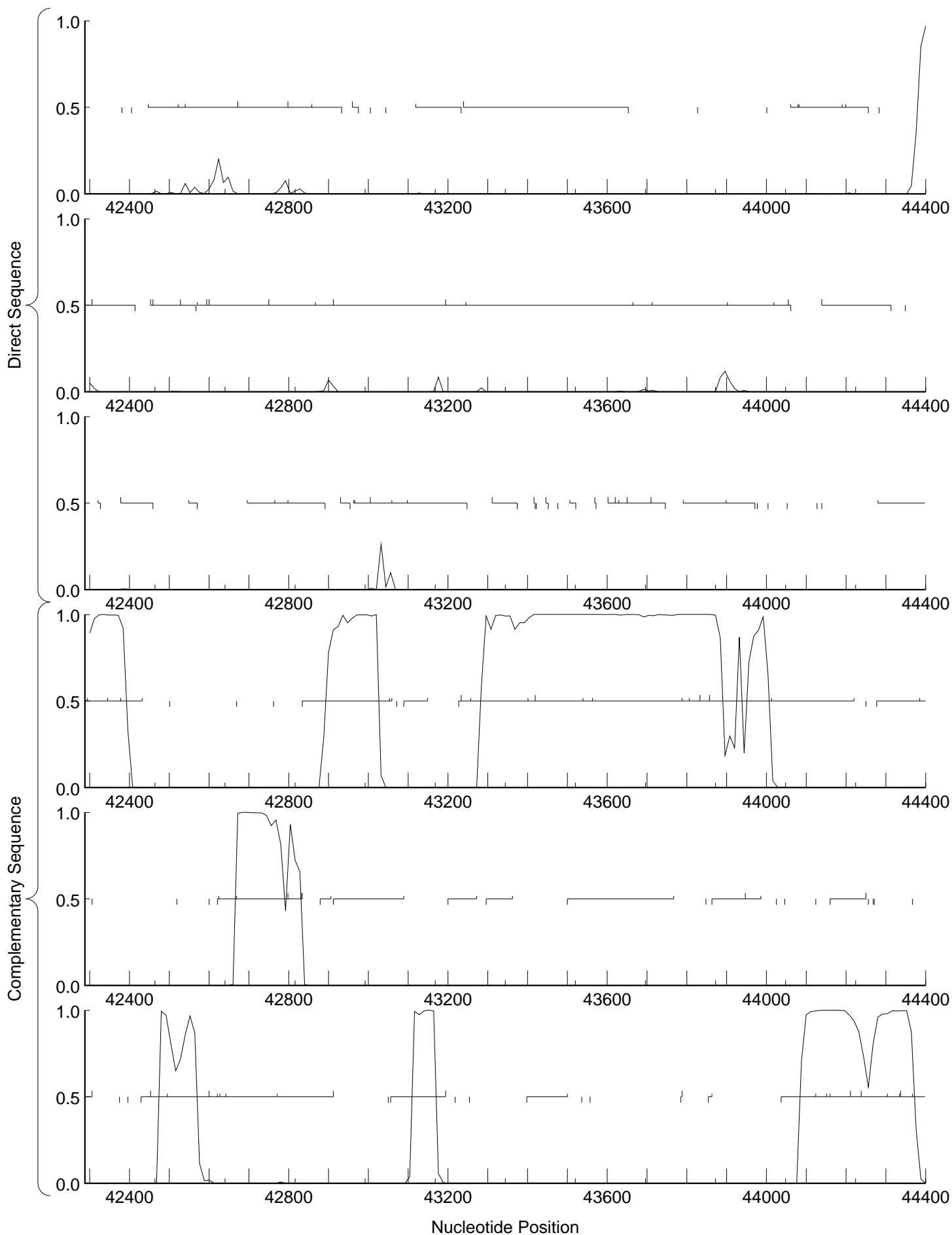


complete sequence, 48545 bp including 10-base 3' overhang (CGGGTGGTAA), Cluster A3, Order 4, Window 96, Step 12, 19/24









complete sequence, 48545 bp including 10-base 3' overhang (CGGGTGGTAA), Cluster A3, Order 4, Window 96, Step 12, 23/24

