

GeneMark

Version 2.5p (09.08.06)

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PROGRAM INFORMATION

Sequence : Mycobacterium phage GardenSalsa complete sequence, 80309 bp including 11-base 3' overhang ()
Analysis Date : 7/2/22 at 1:40:12
Pages : 40
Sequence Length : 80309 bp
GC Content : 60.92%

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,

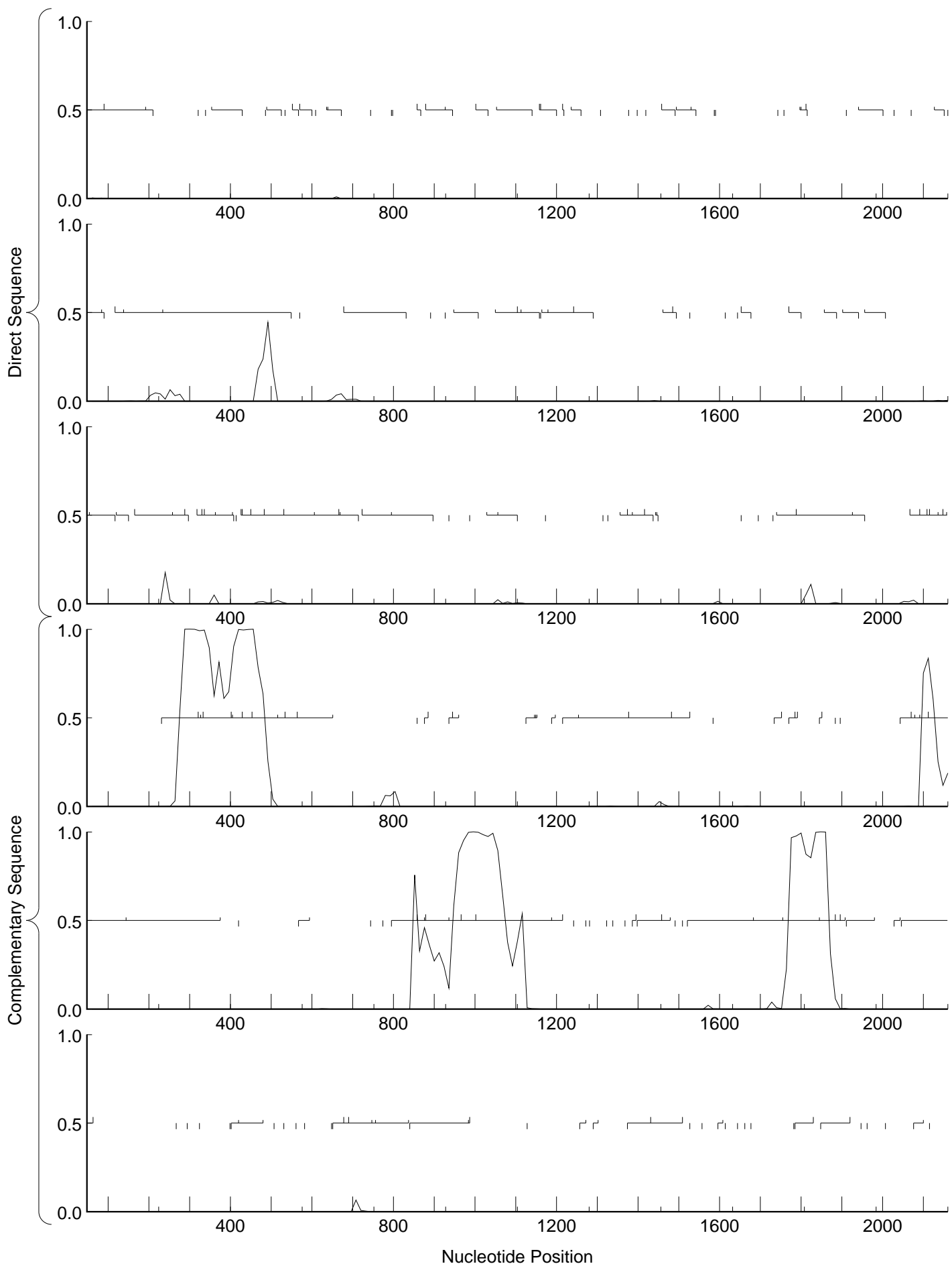
MATRIX INFORMATION

Matrix : Mycobacterium_tuberculosis_H37Rv
Author : -
Order : 4

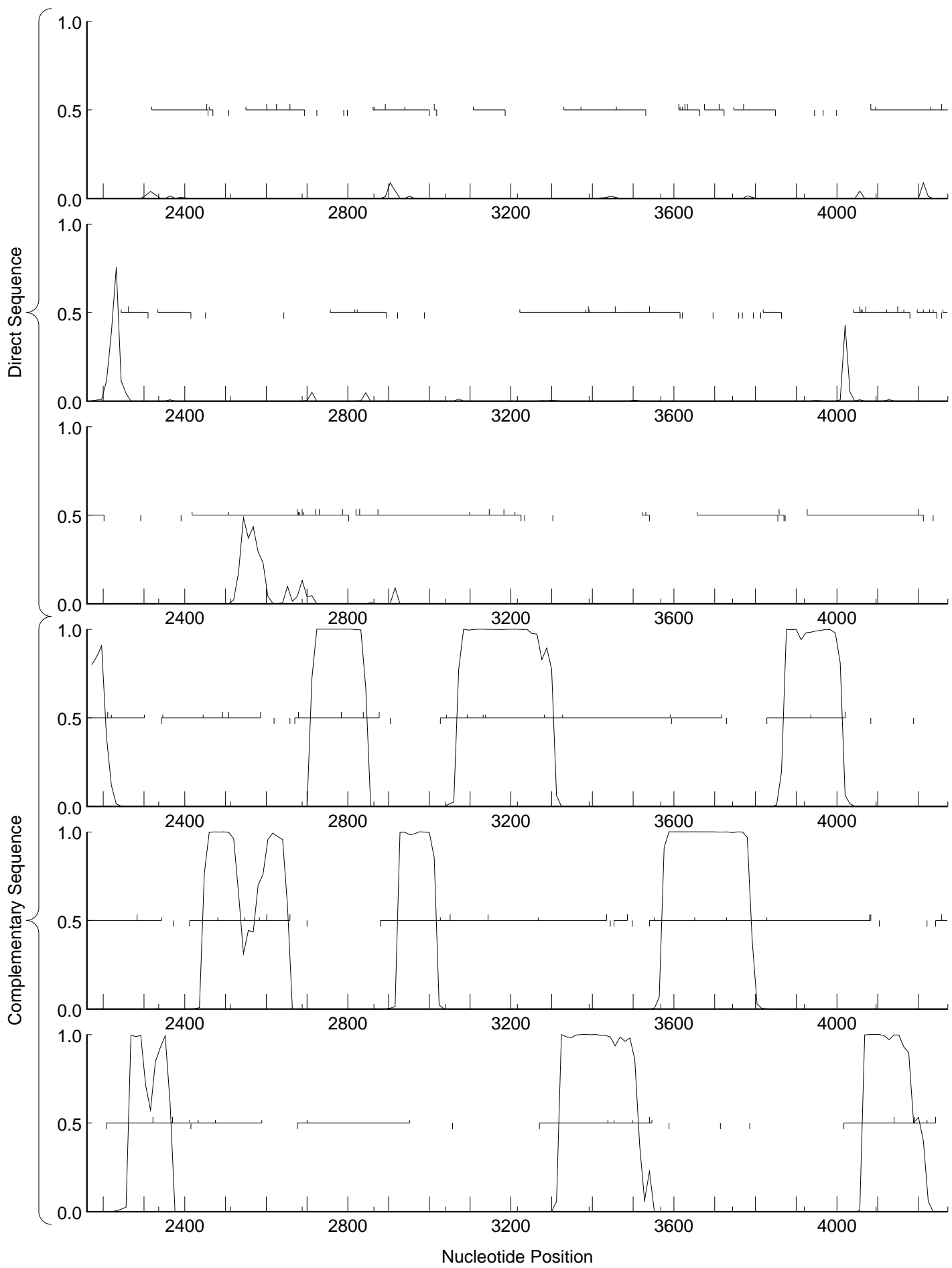
Send questions / comments to:
Dr. M. Borodovsky
Georgia Institute of Technology
School of Biology
Atlanta, GA 30332-0230

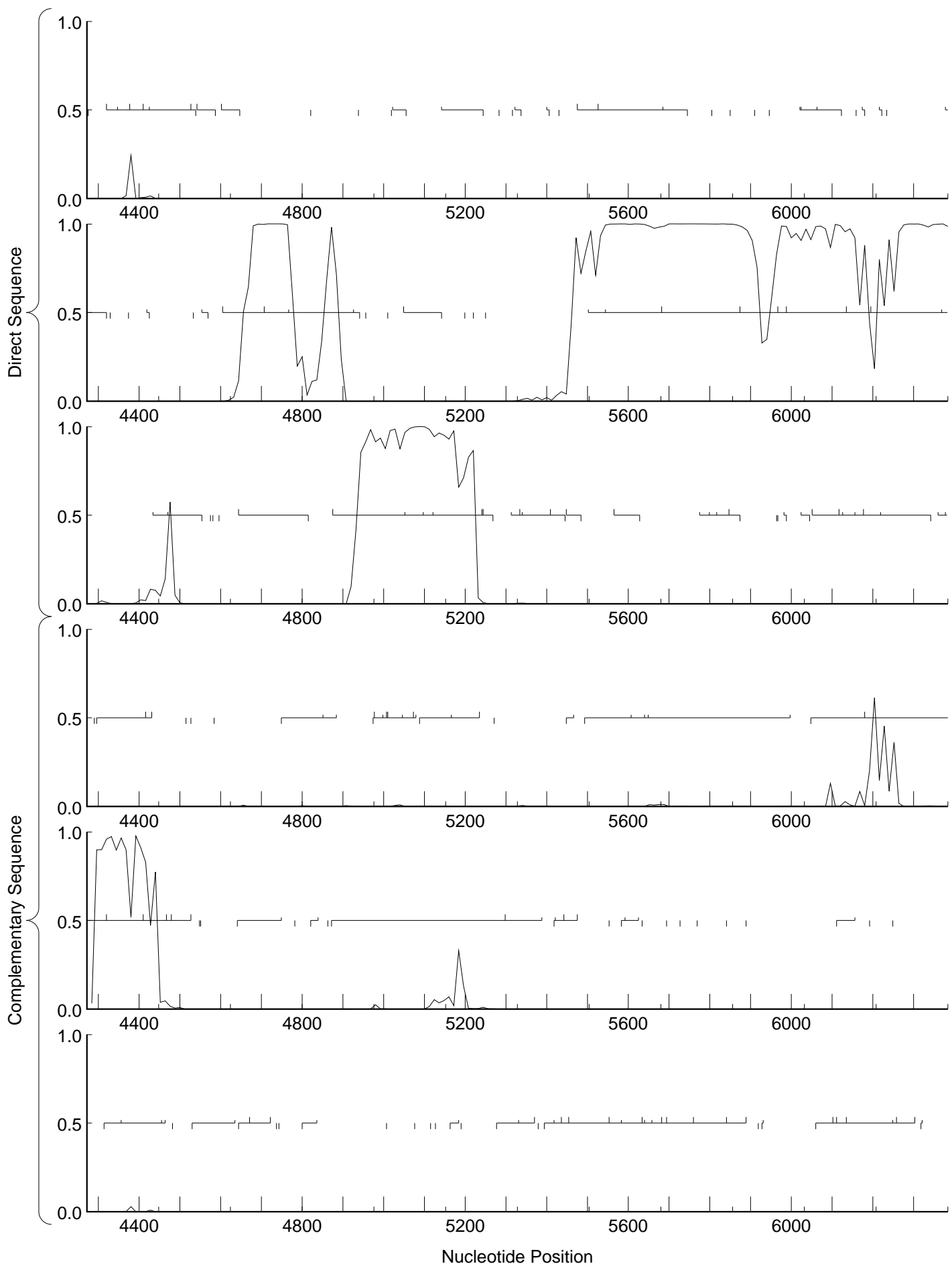
Matrix notes & comments

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

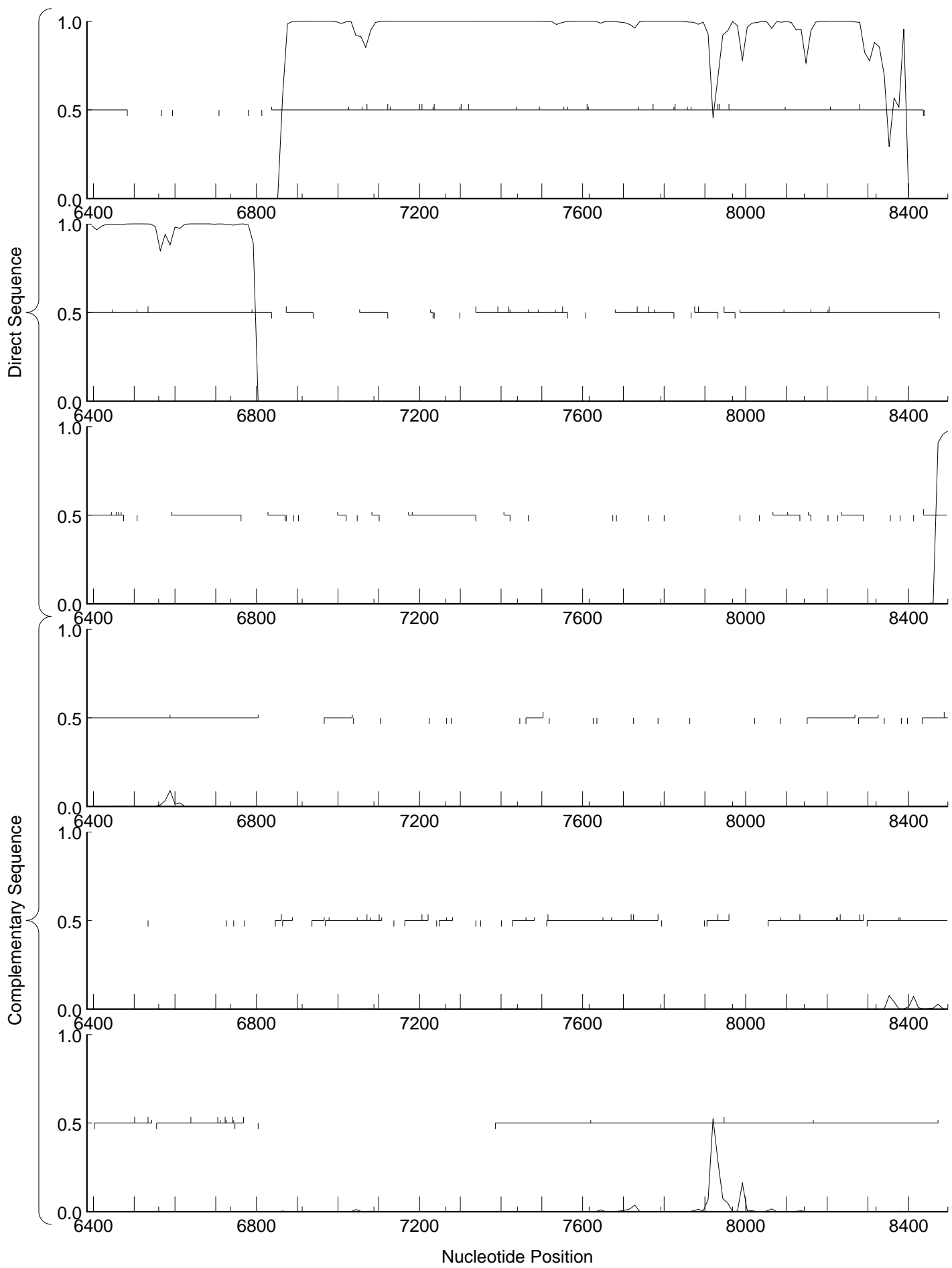


complete sequence, 80309 bp including 11-base 3' overhang (ACCCATGCAA), Cluster M2, Order 4, Window 96, Step 12, 3/40

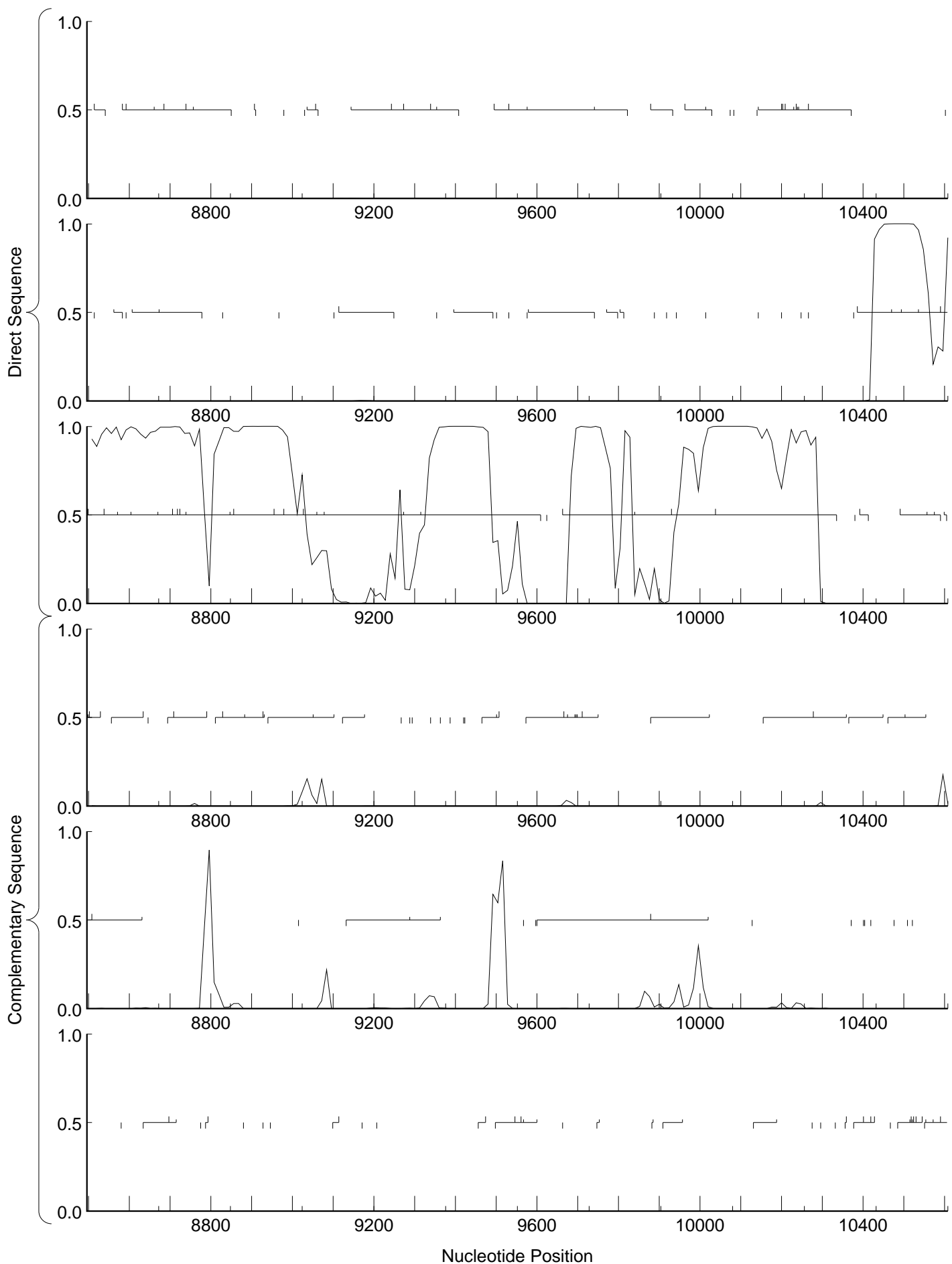




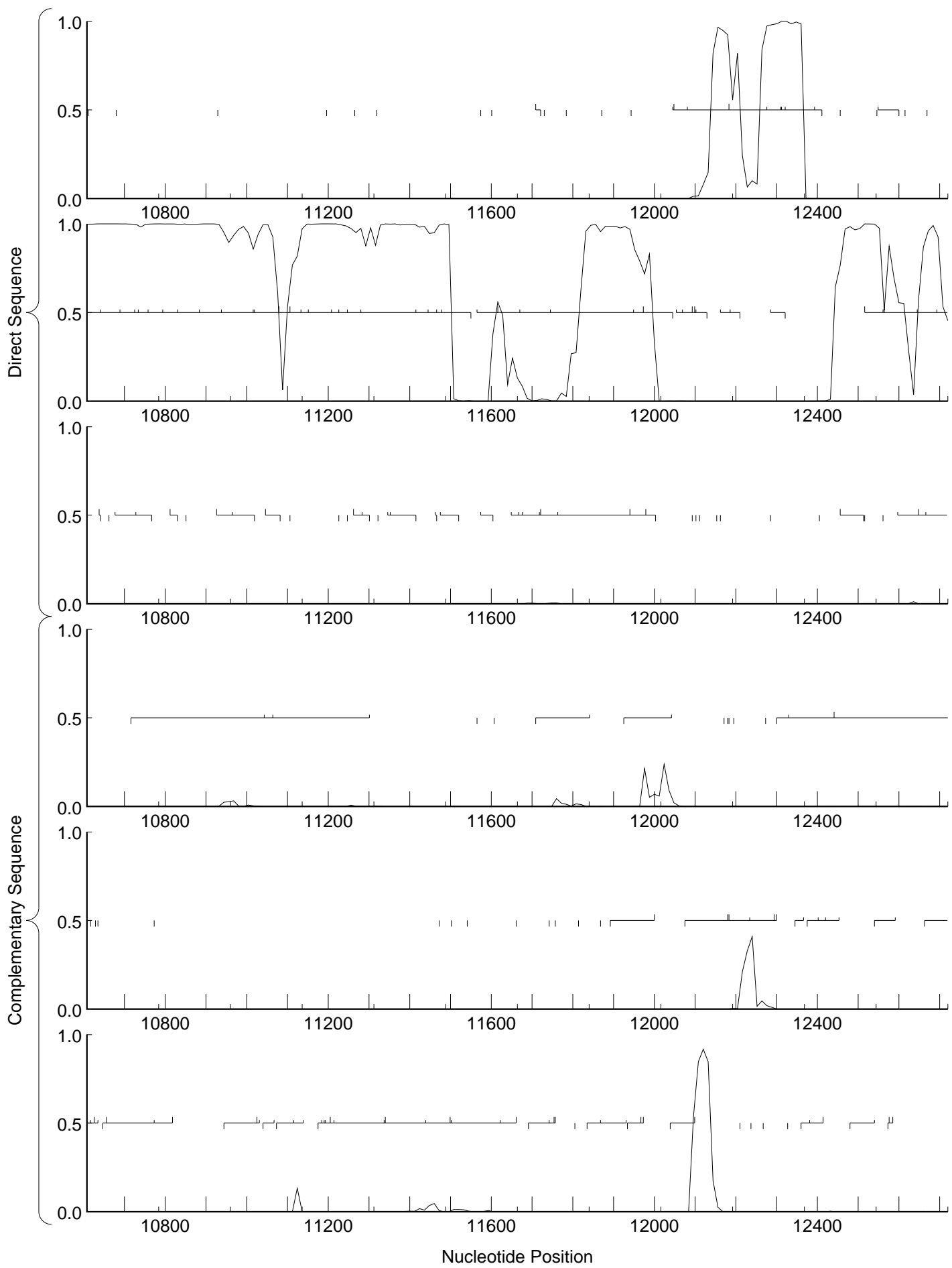
complete sequence, 80309 bp including 11-base 3' overhang (ACCCATGCAA), Cluster M2, Order 4, Window 96, Step 12, 5/40

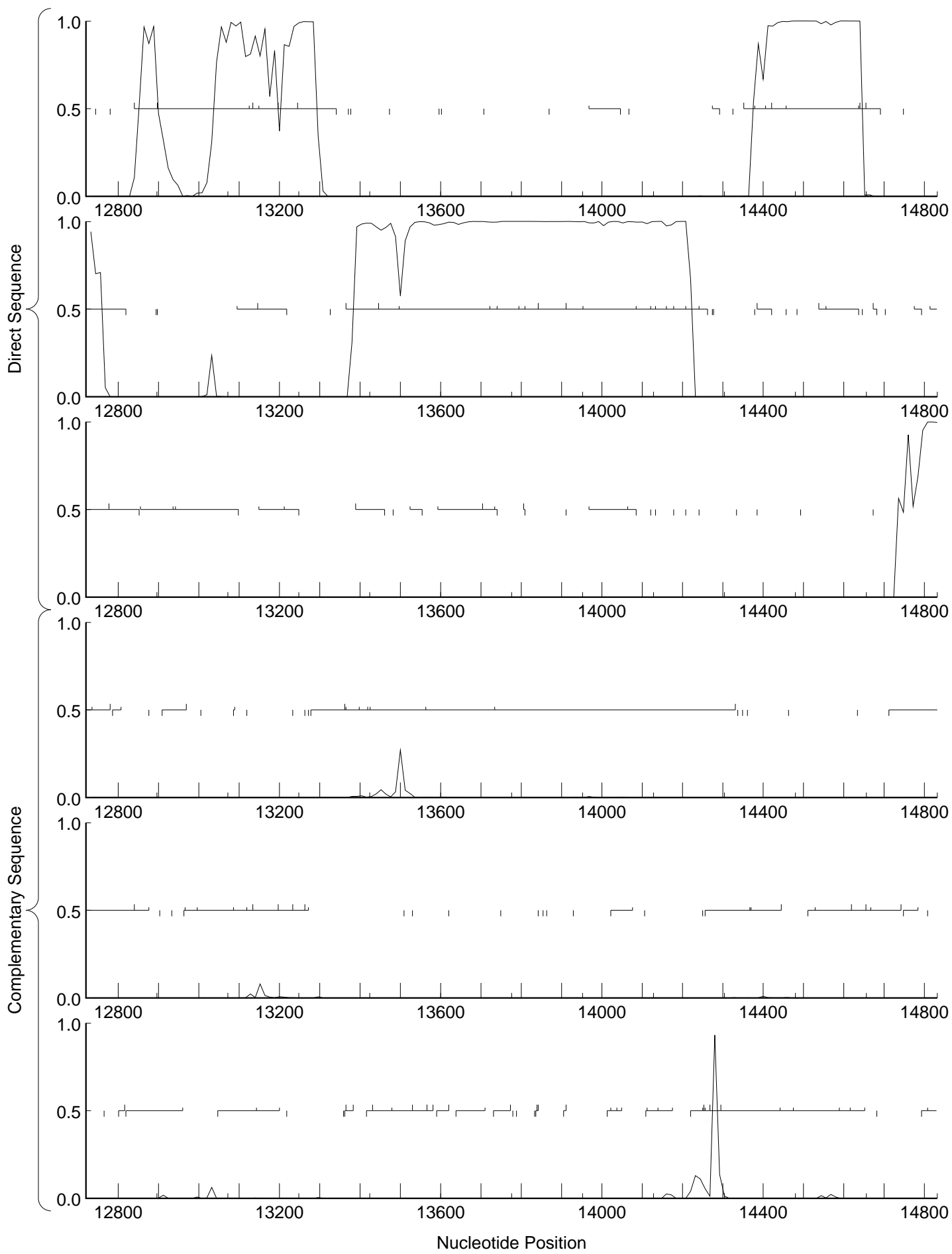


complete sequence, 80309 bp including 11-base 3' overhang (ACCCATGCAA), Cluster M2, Order 4, Window 96, Step 12, 6/40

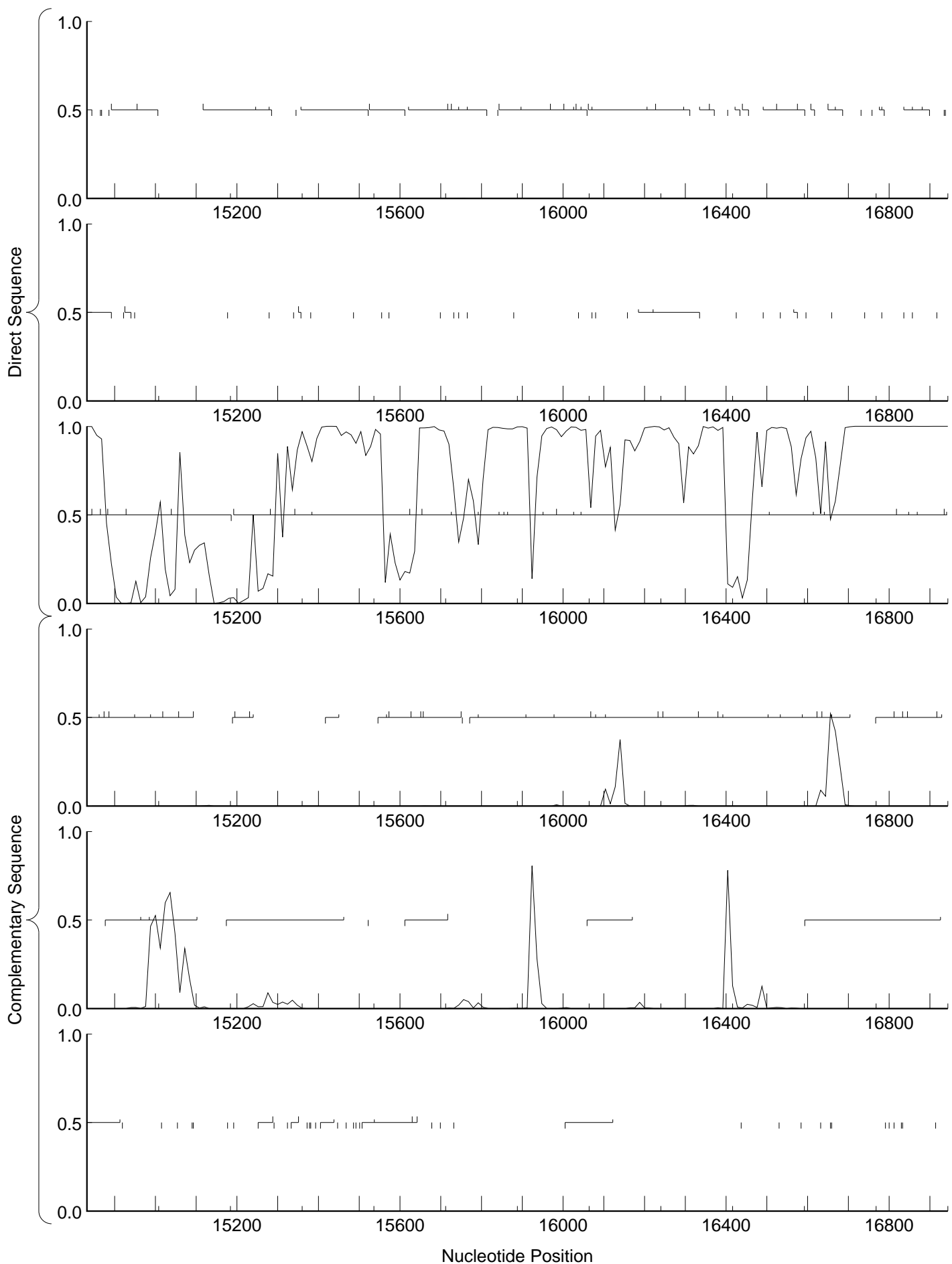


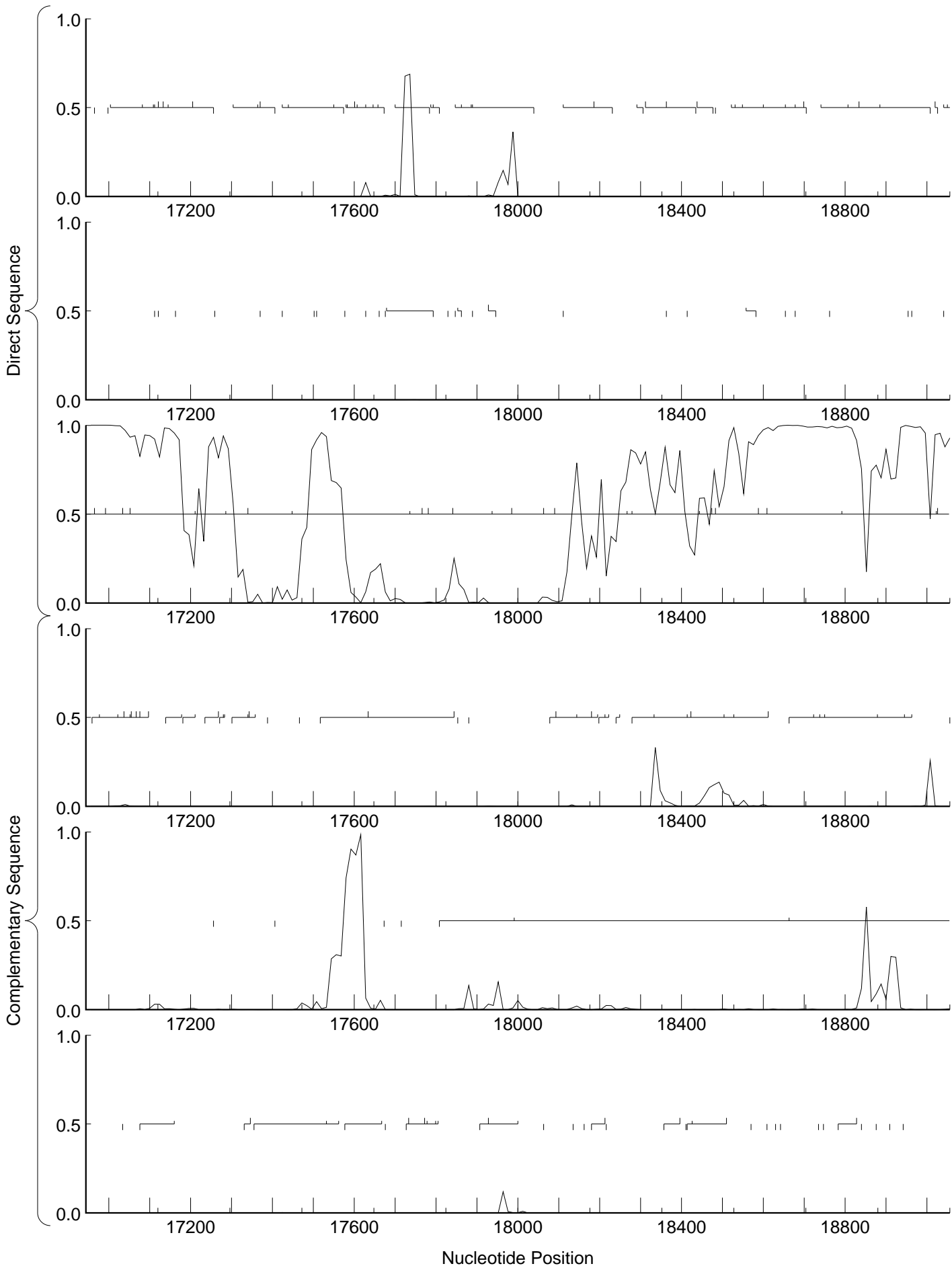
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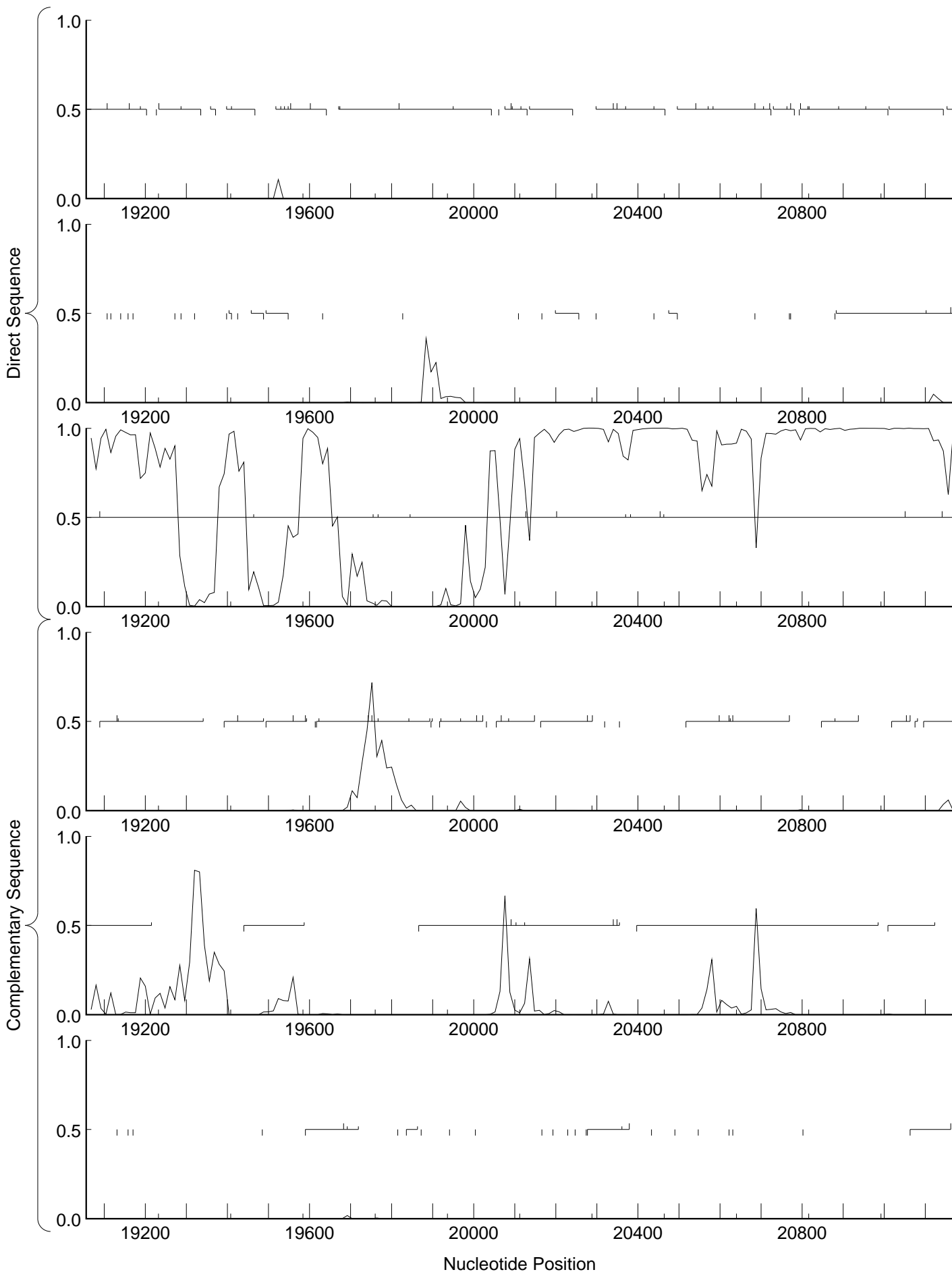


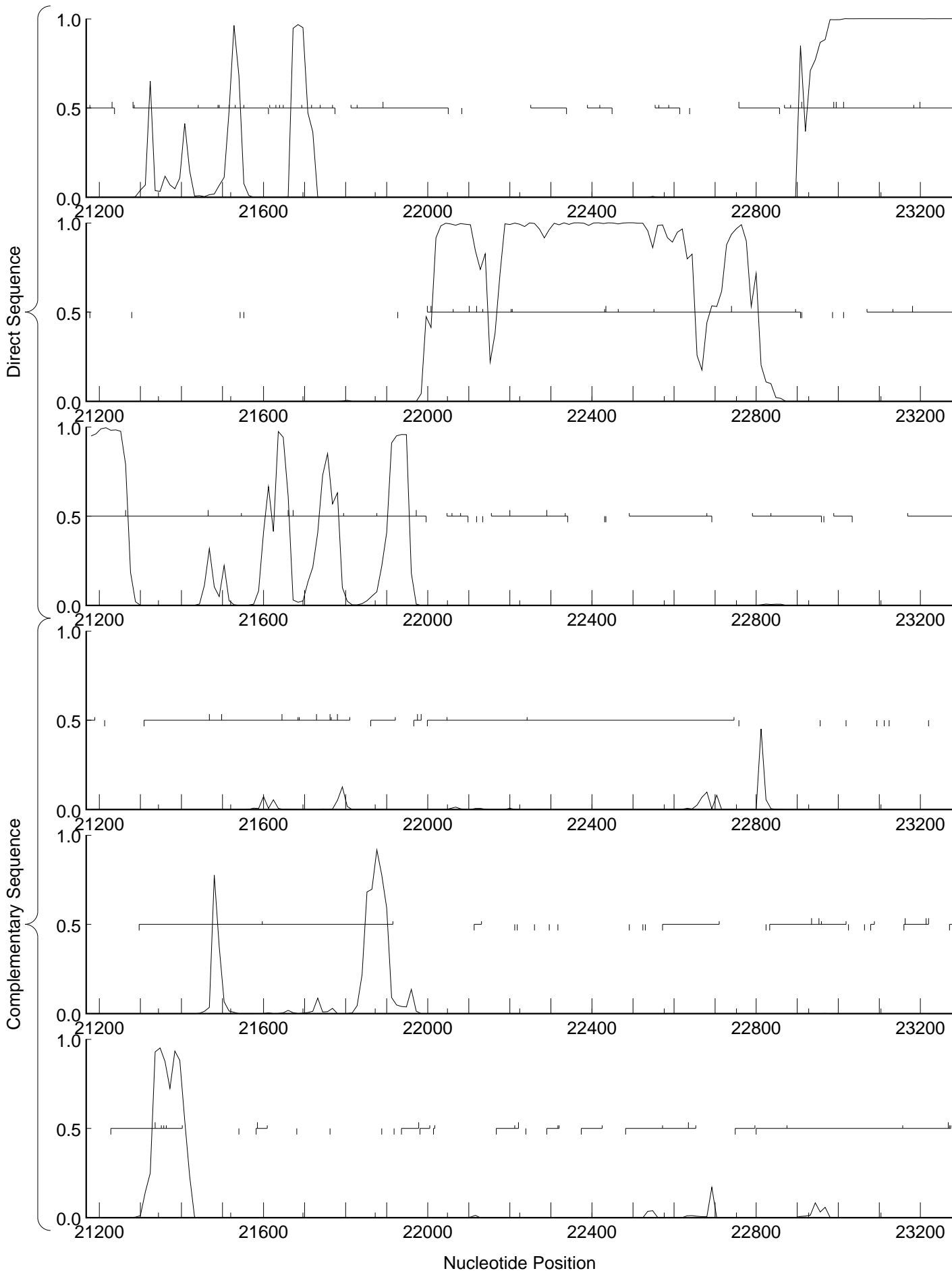
complete sequence, 80309 bp including 11-base 3' overhang (ACCCATGCAA), Cluster M2, Order 4, Window 96, Step 12, 9/40



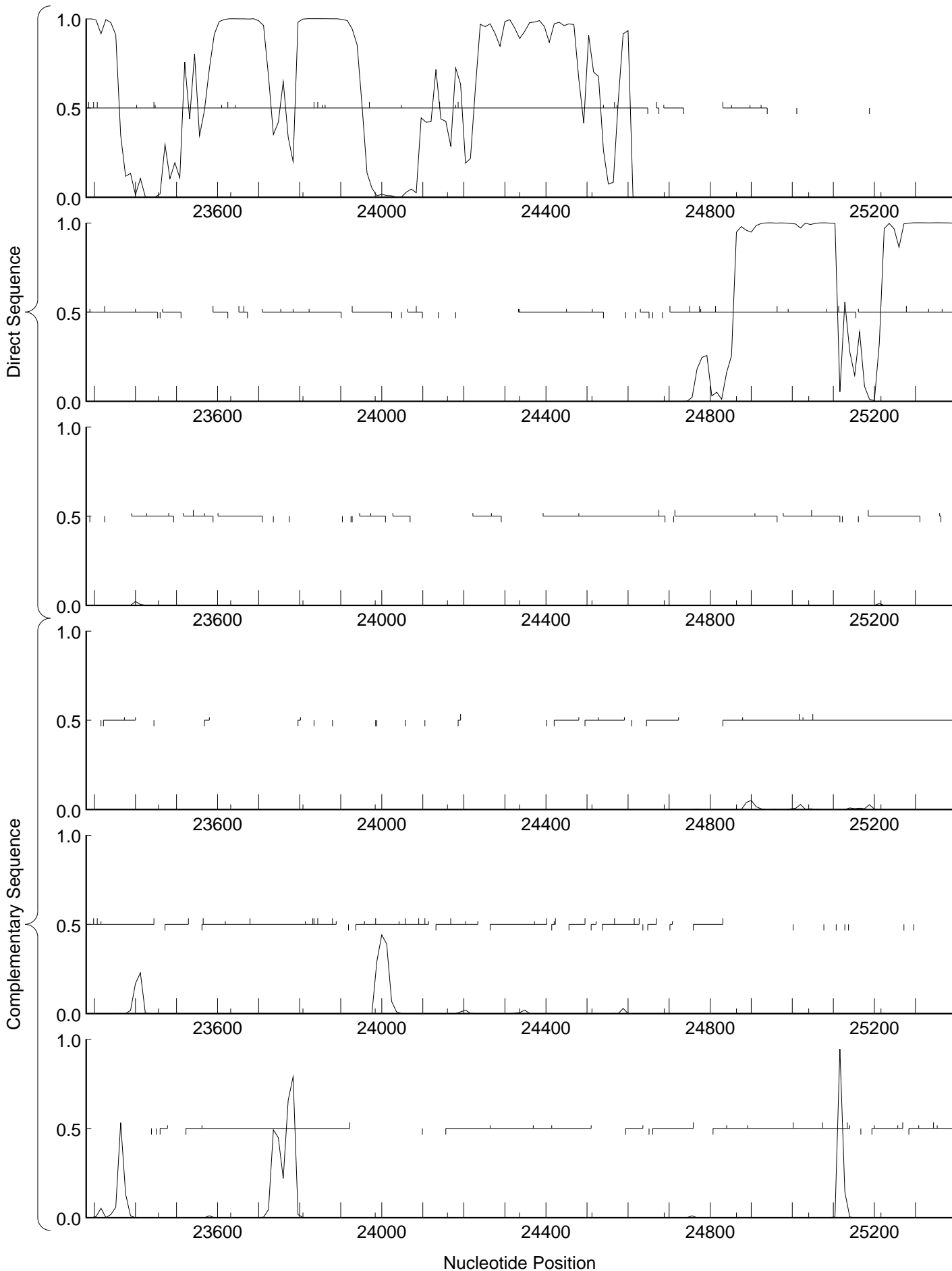


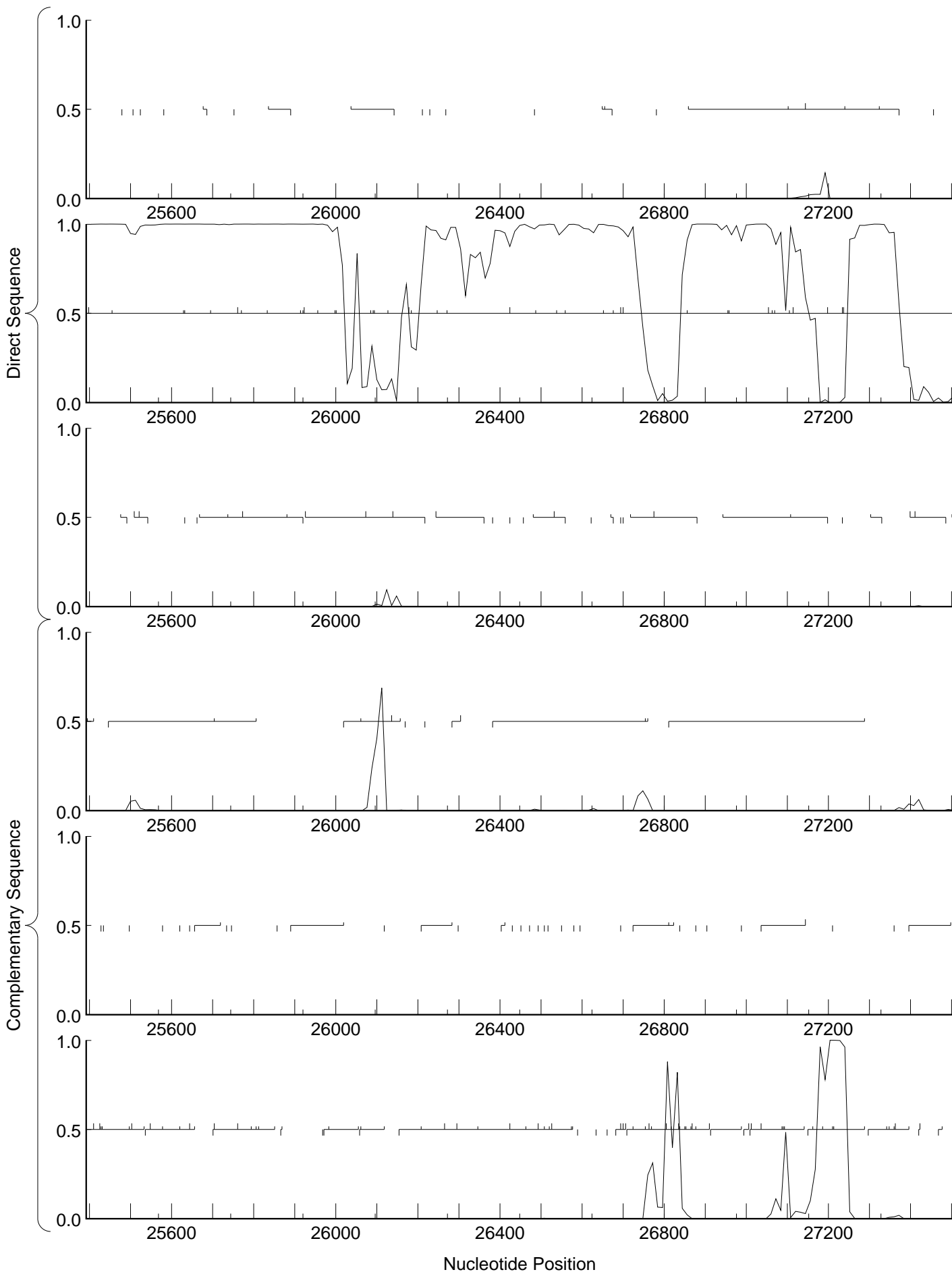
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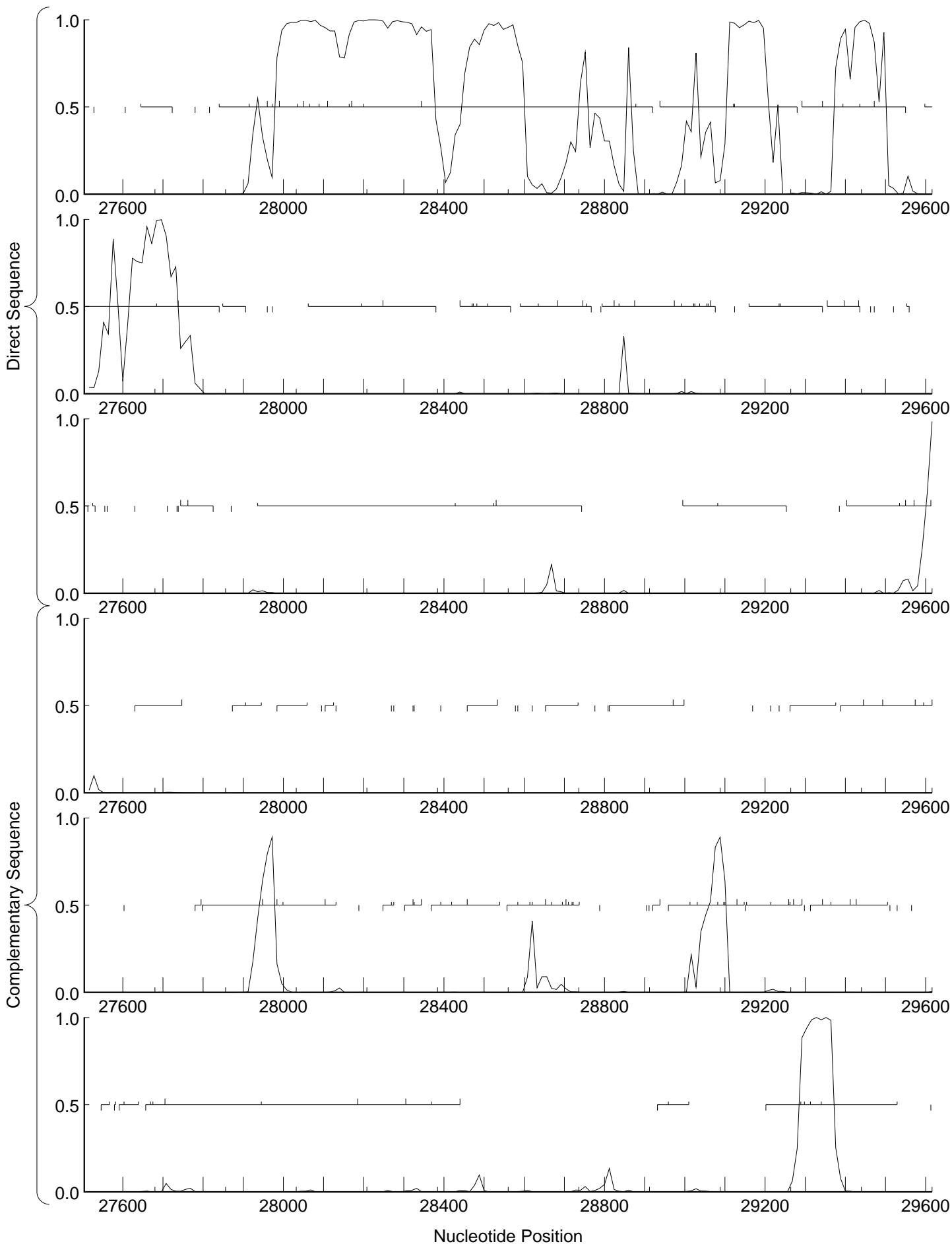


complete sequence, 80309 bp including 11-base 3' overhang (ACCCATGCAA), Cluster M2, Order 4, Window 96, Step 12, 13/40

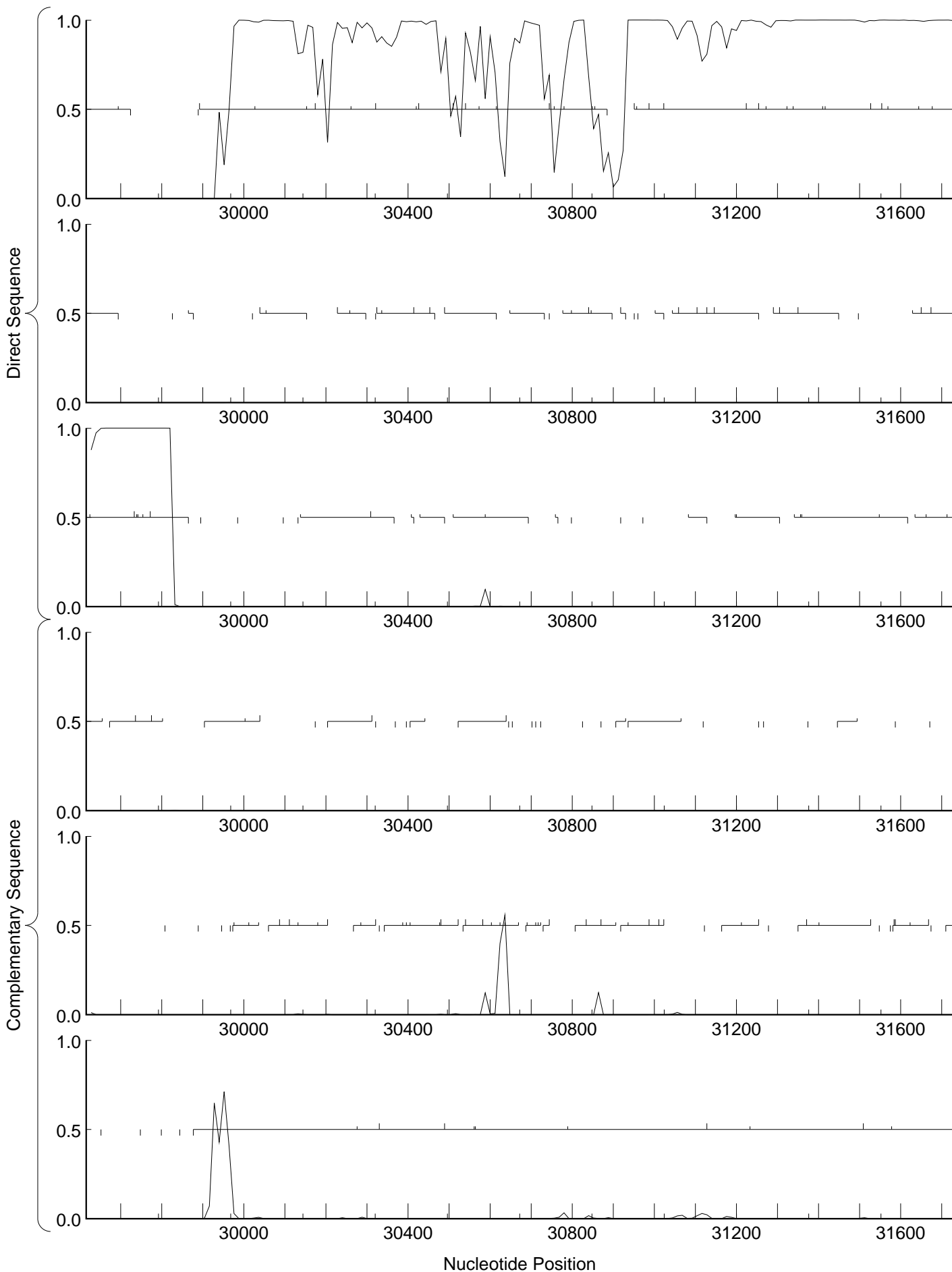




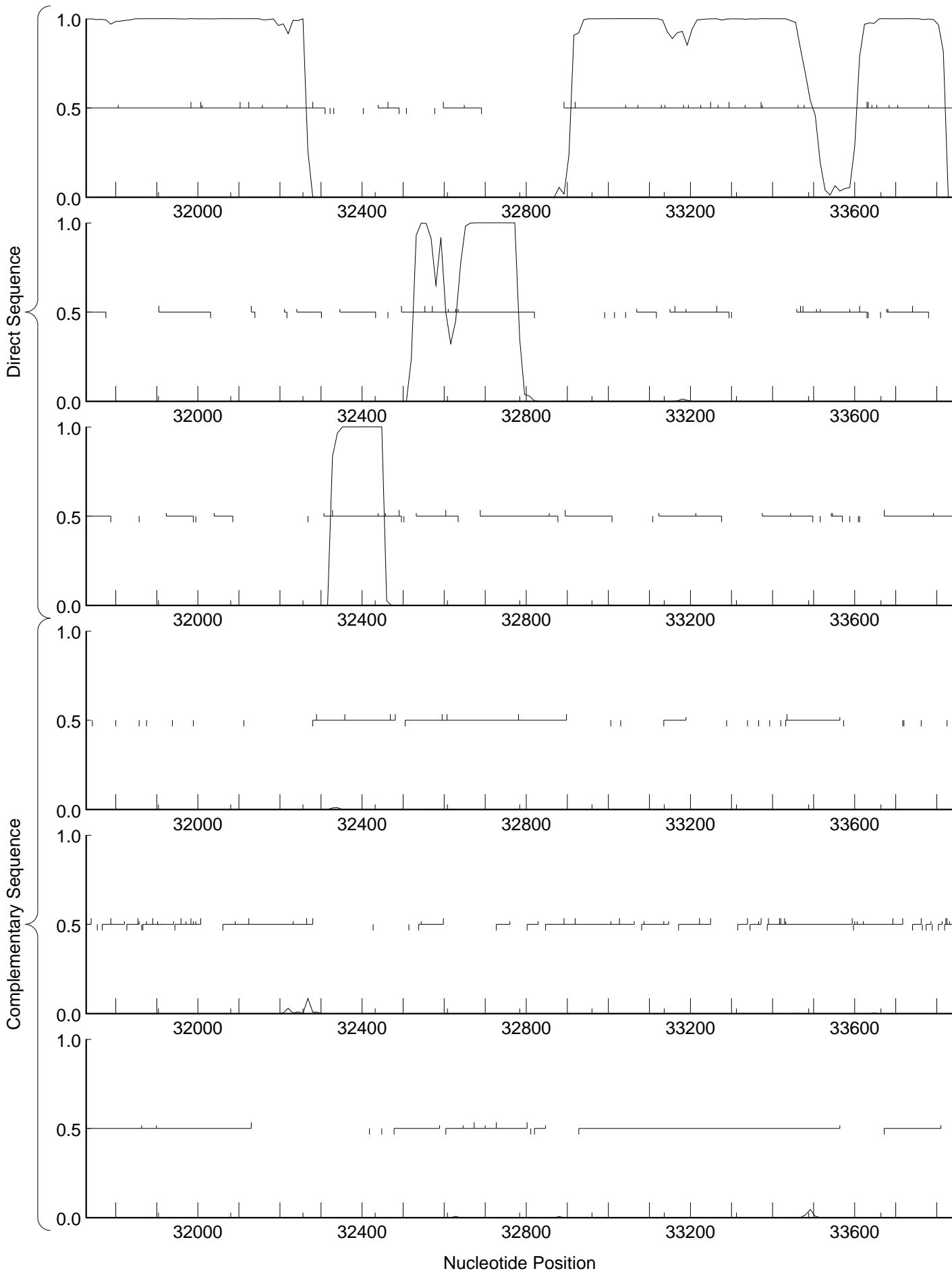
complete sequence, 80309 bp including 11-base 3' overhang (ACCCCATGCAA), Cluster M2, Order 4, Window 96, Step 12, 15/40

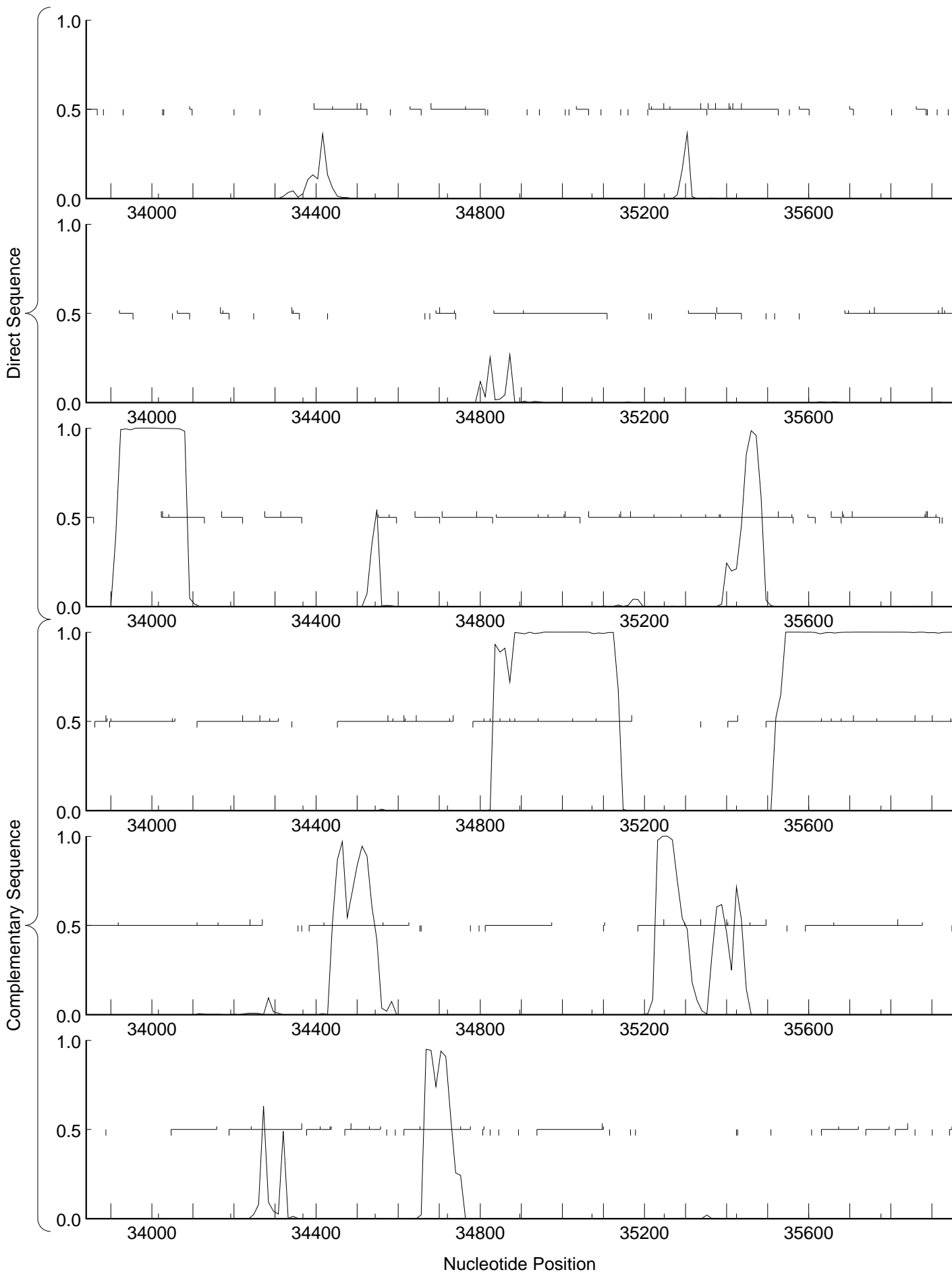


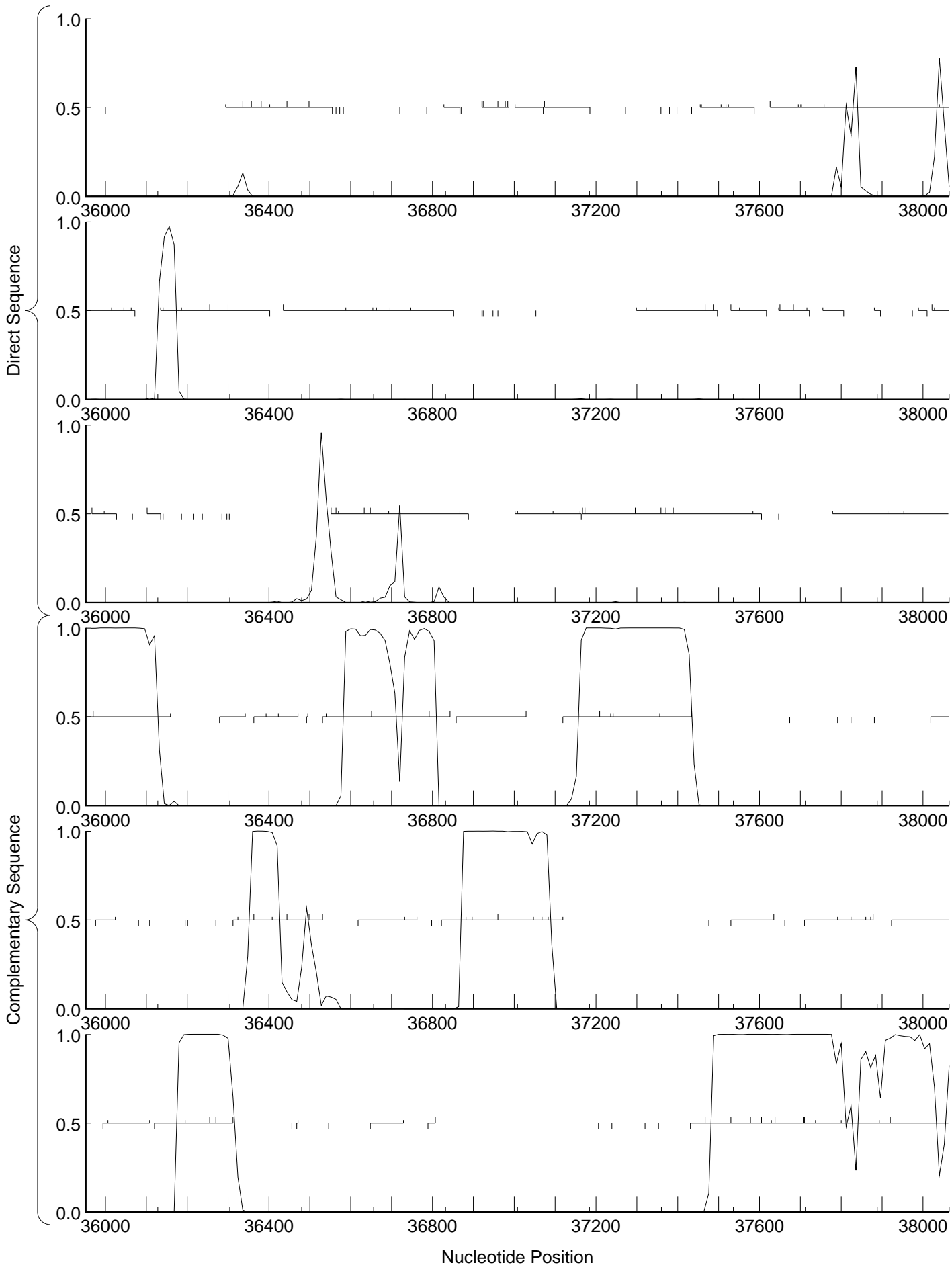
complete sequence, 80309 bp including 11-base 3' overhang (ACCCATGCAA), Cluster M2, Order 4, Window 96, Step 12, 16/40

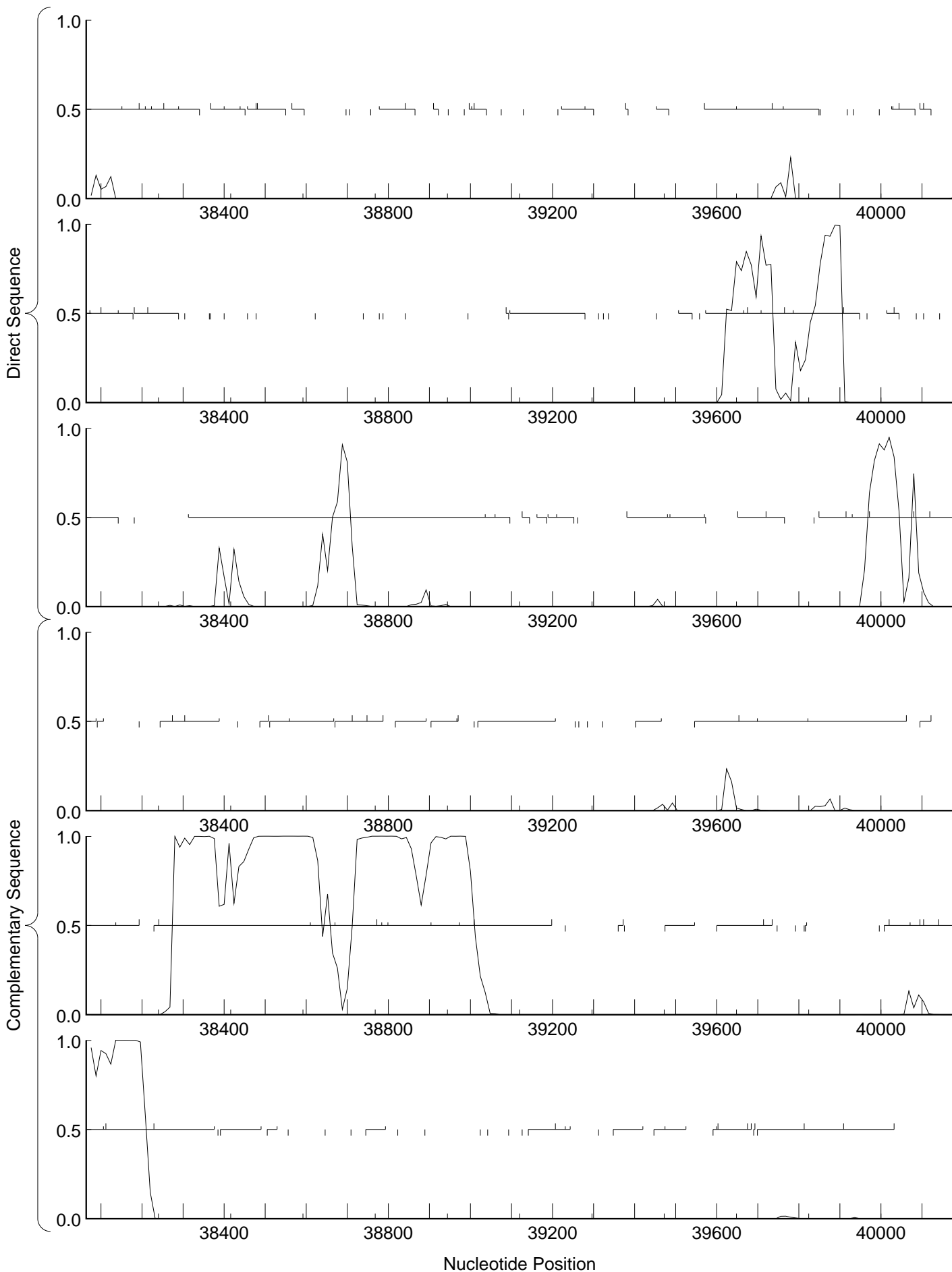


complete sequence, 80309 bp including 11-base 3' overhang (ACCCATGCAA), Cluster M2, Order 4, Window 96, Step 12, 17/40

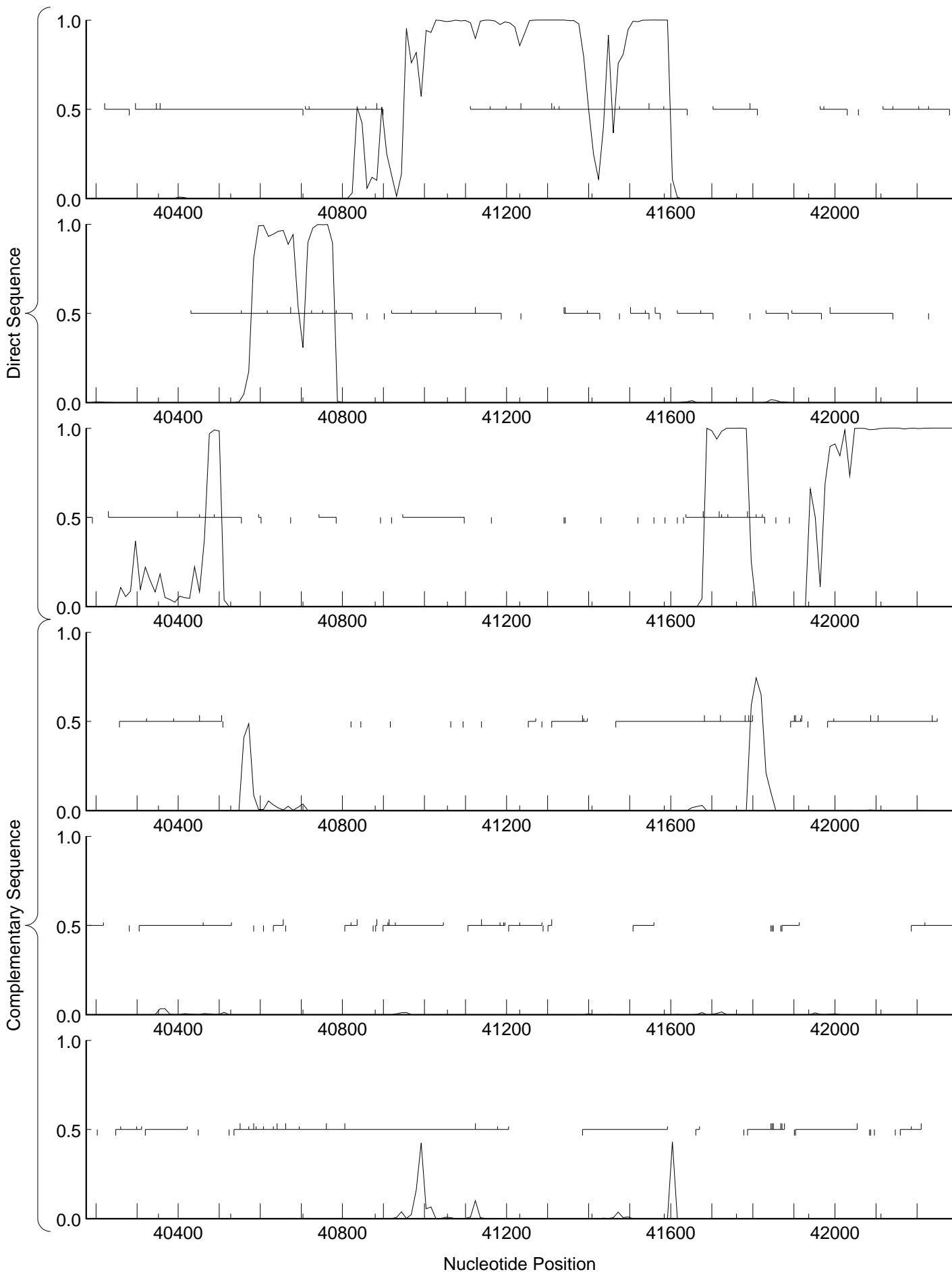


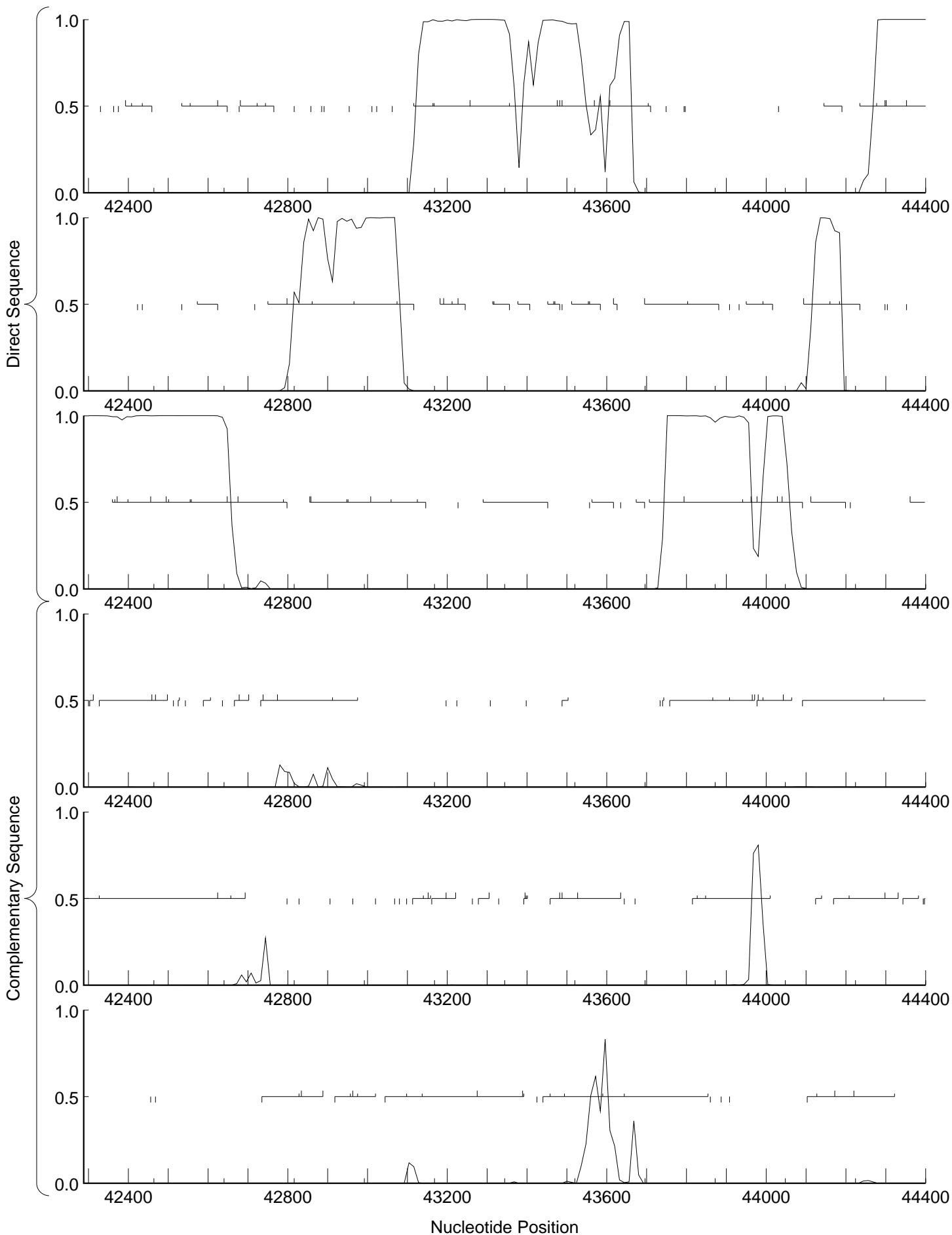


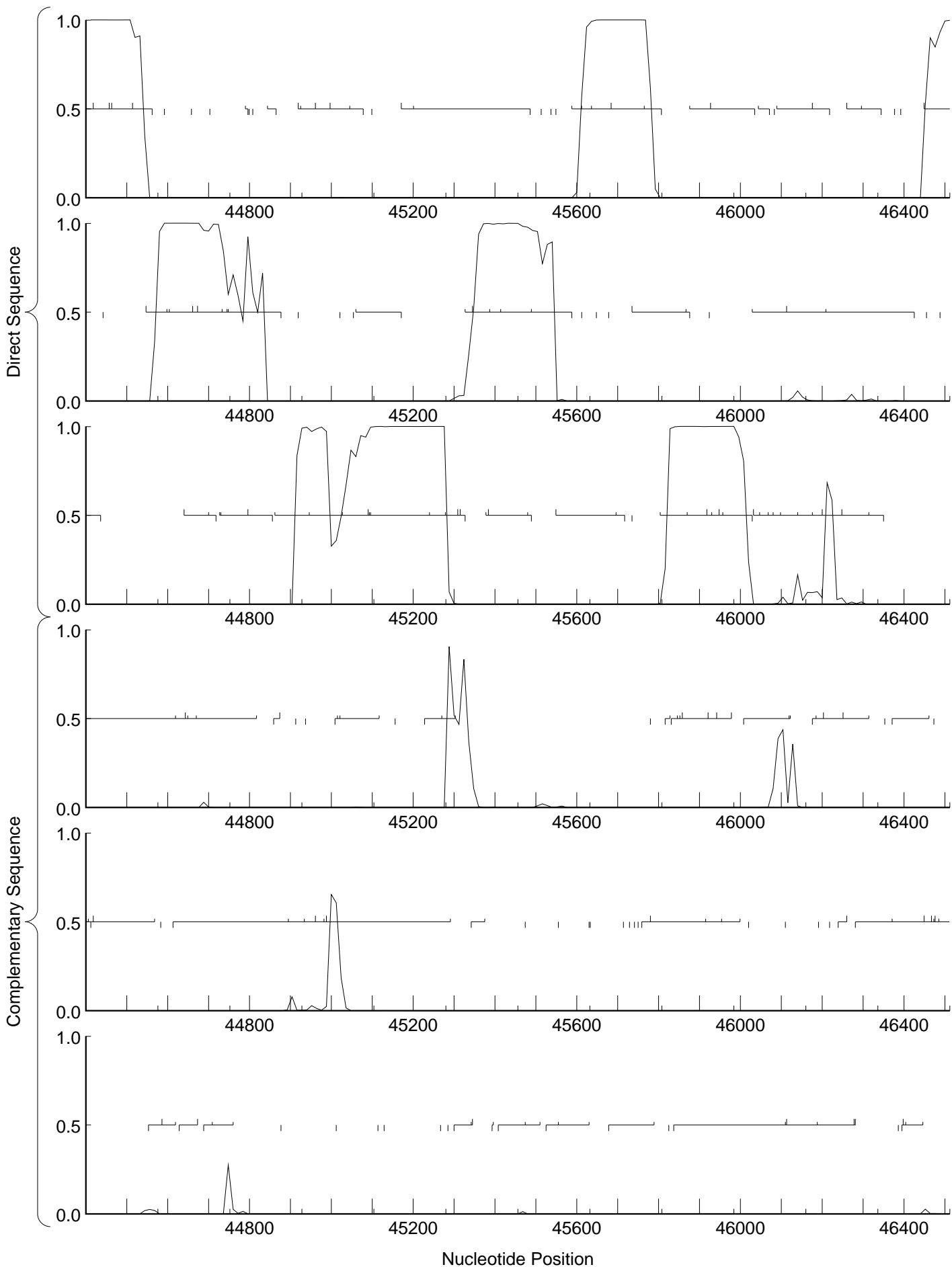


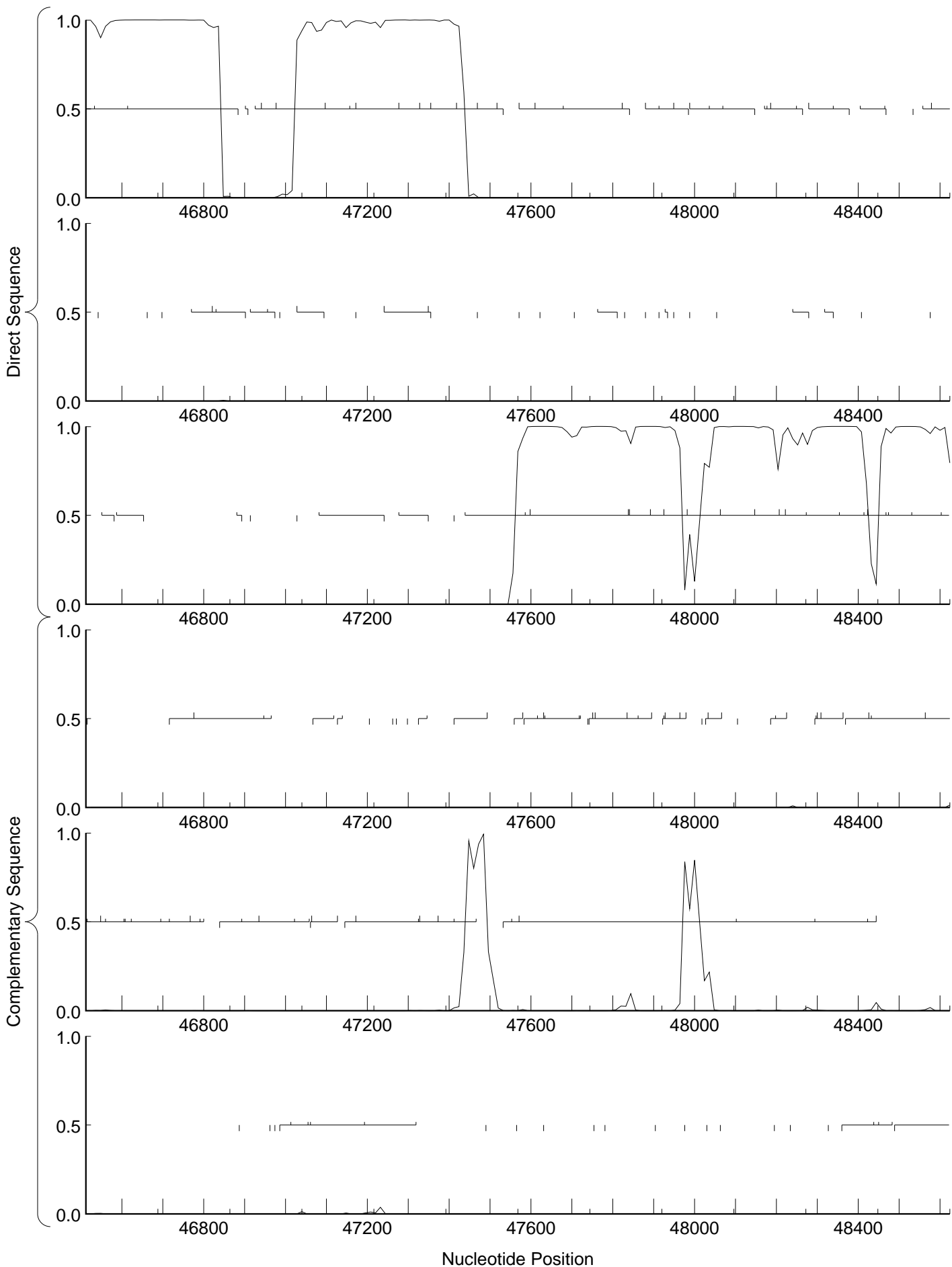


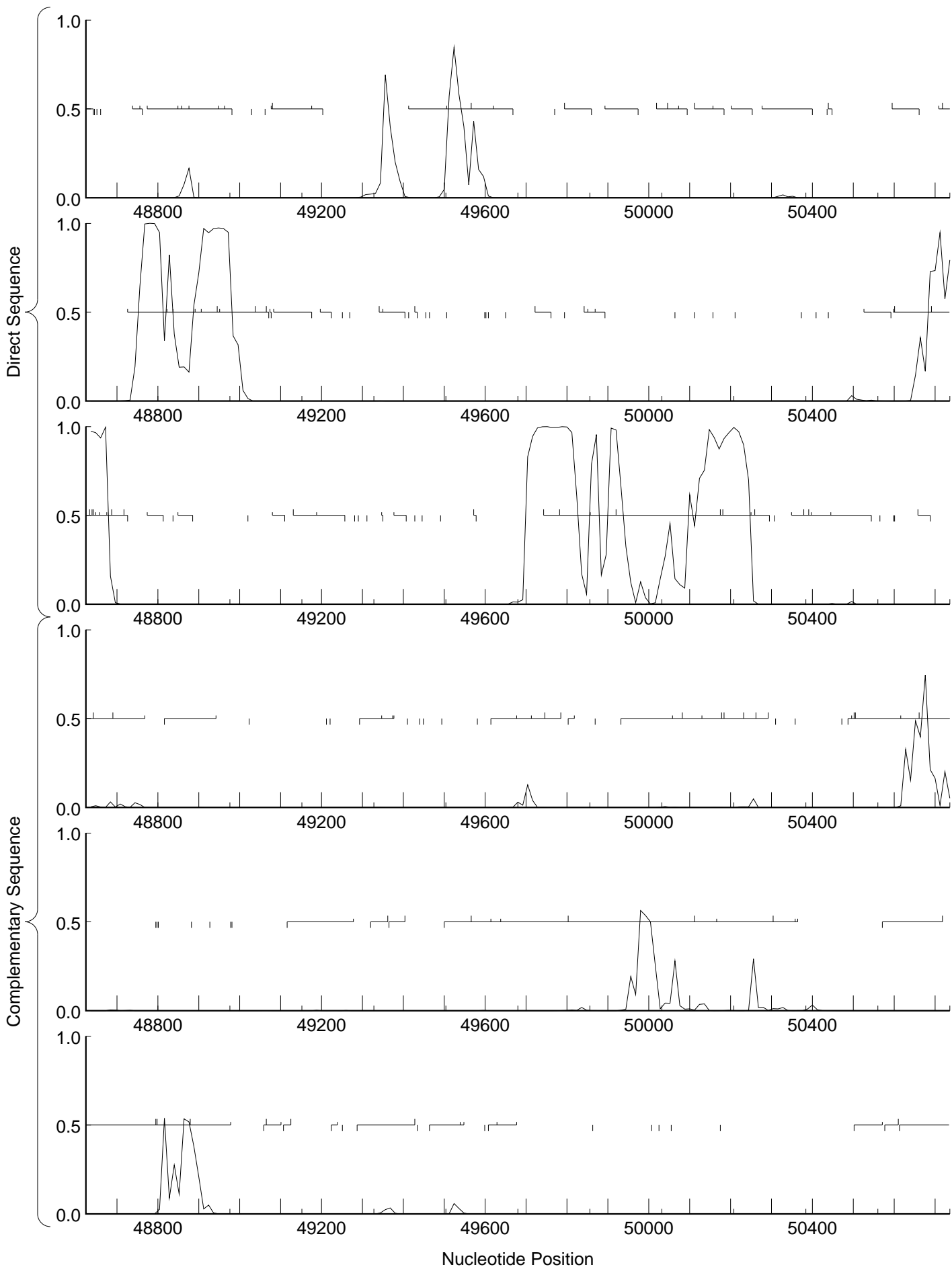
complete sequence, 80309 bp including 11-base 3' overhang (ACCCCATGCAA), Cluster M2, Order 4, Window 96, Step 12, 21/40



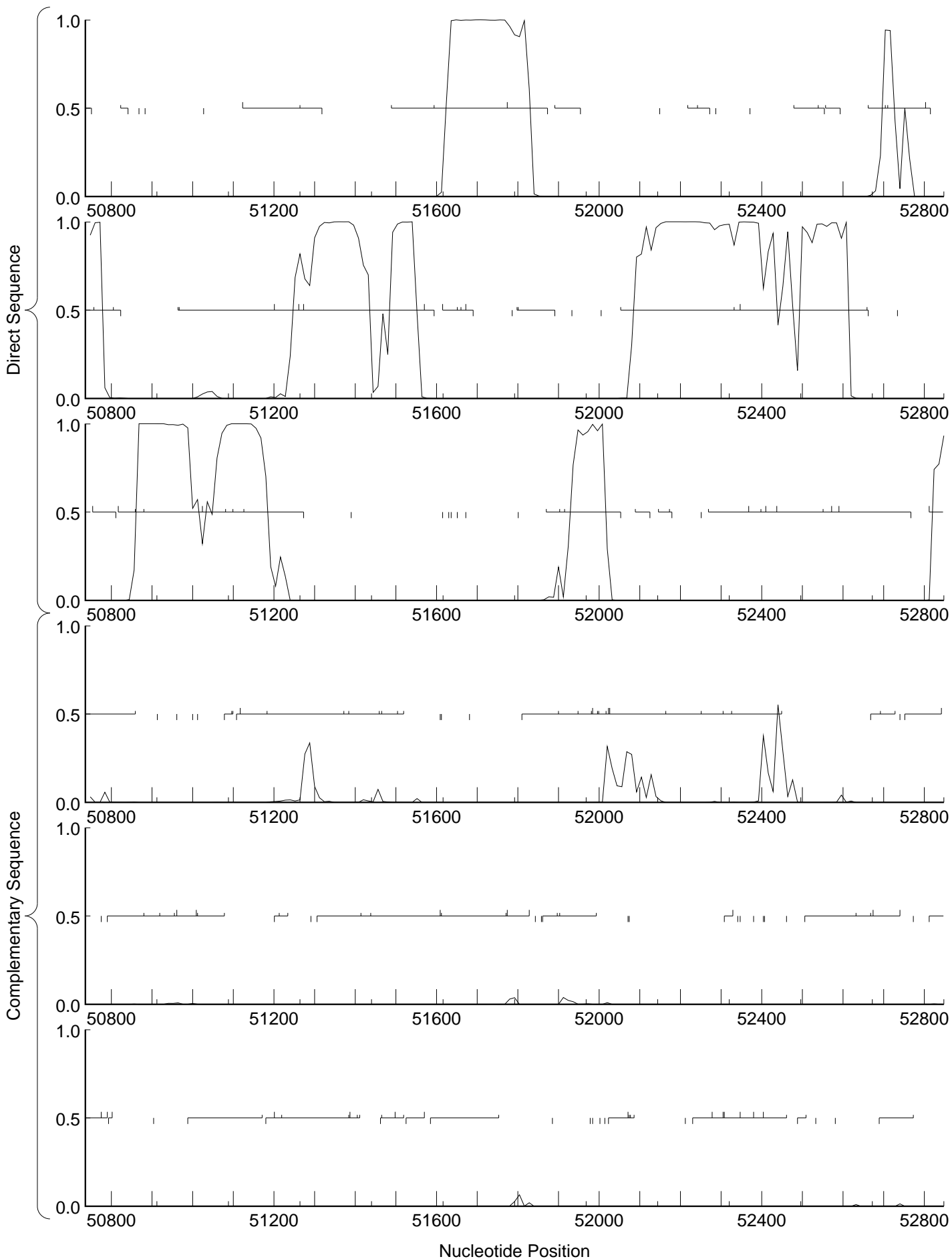


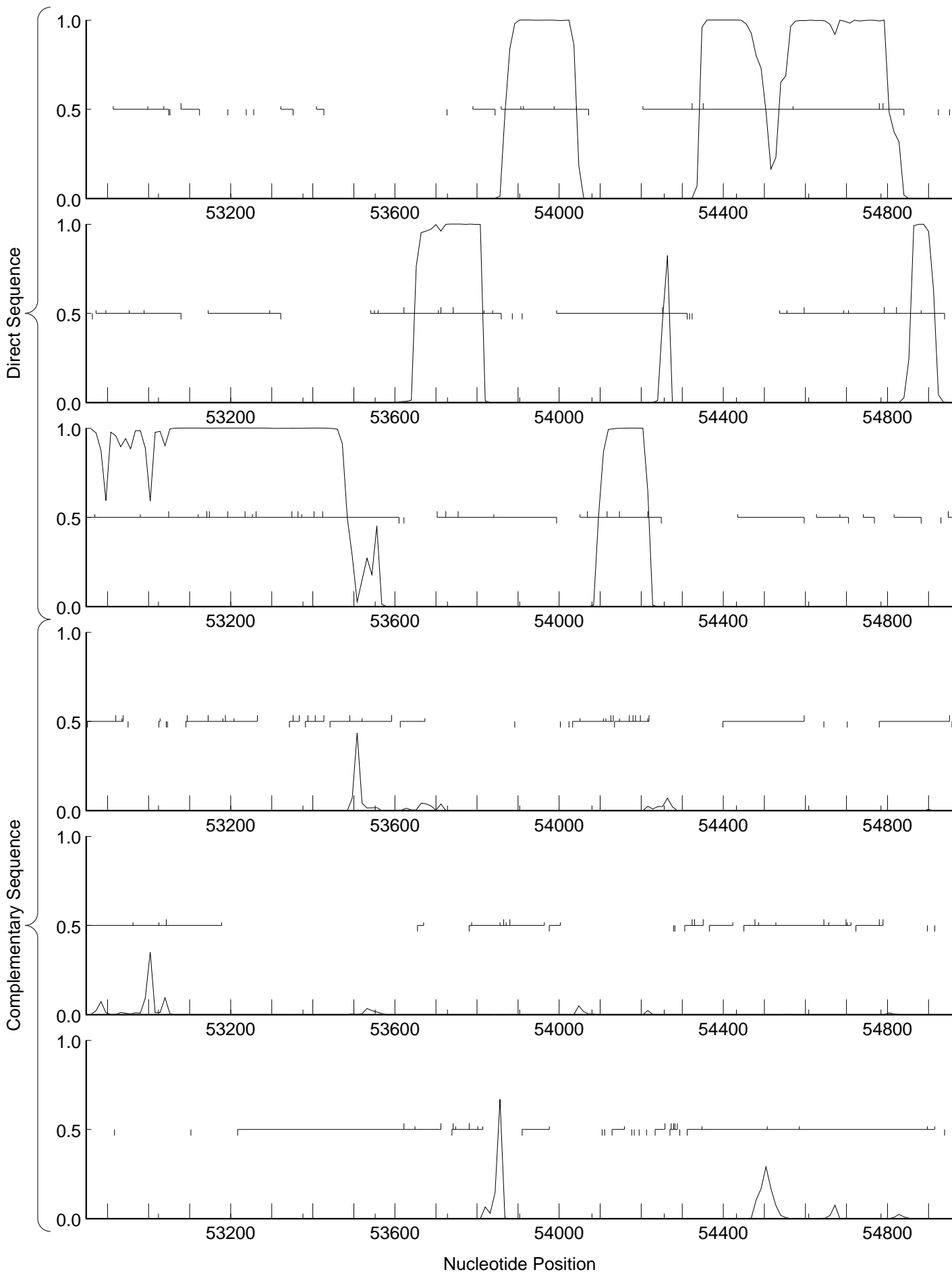


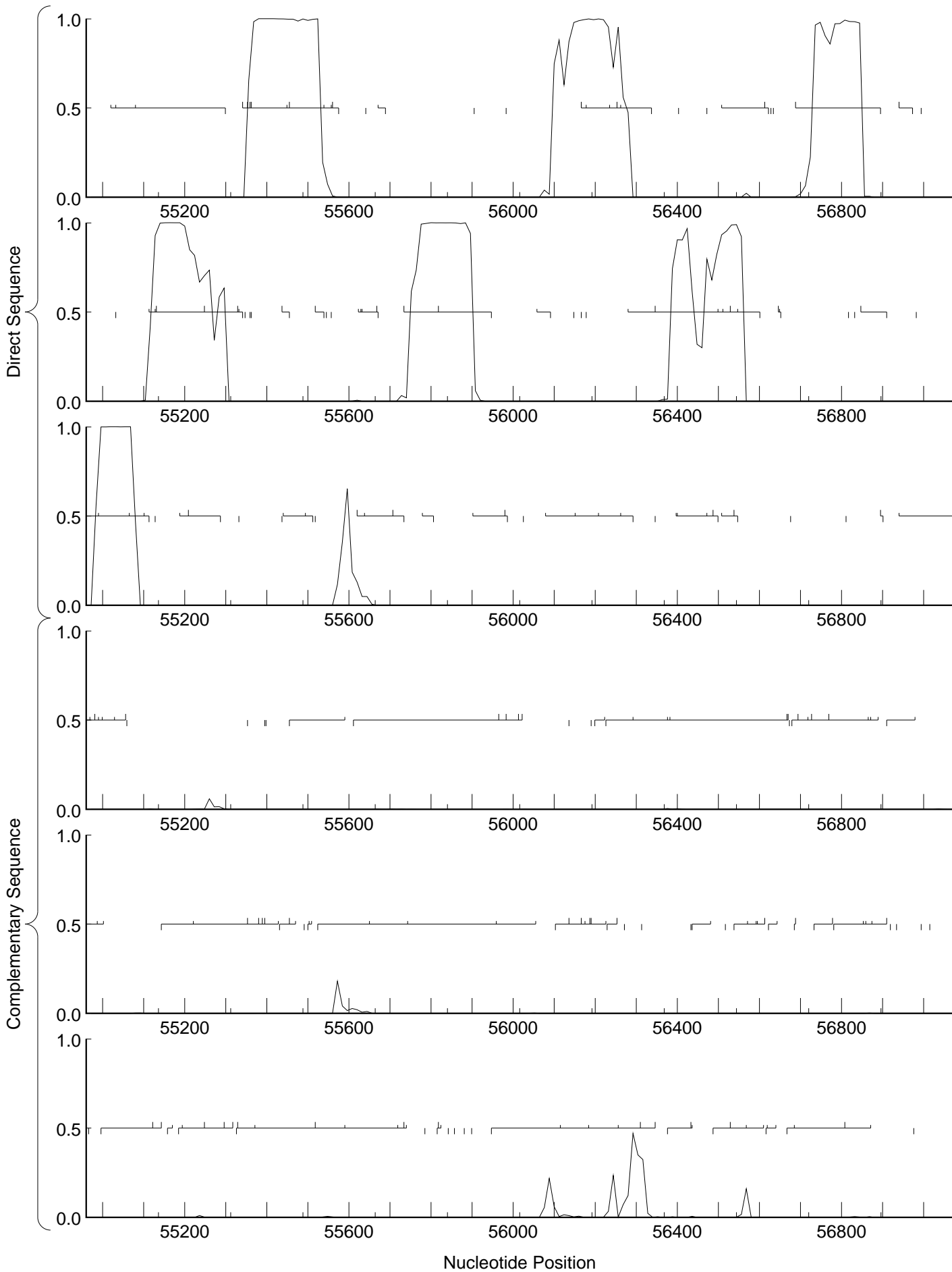


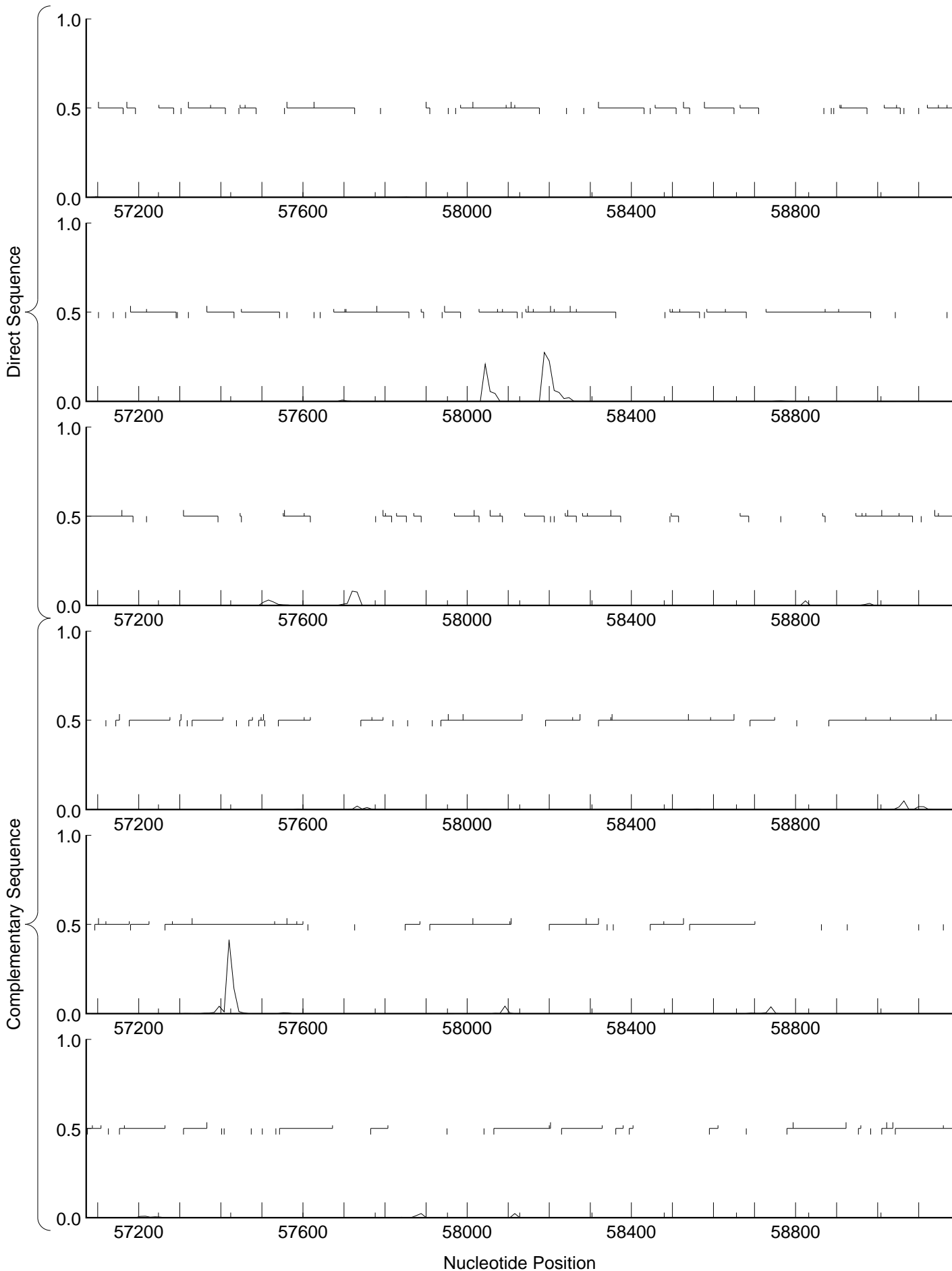


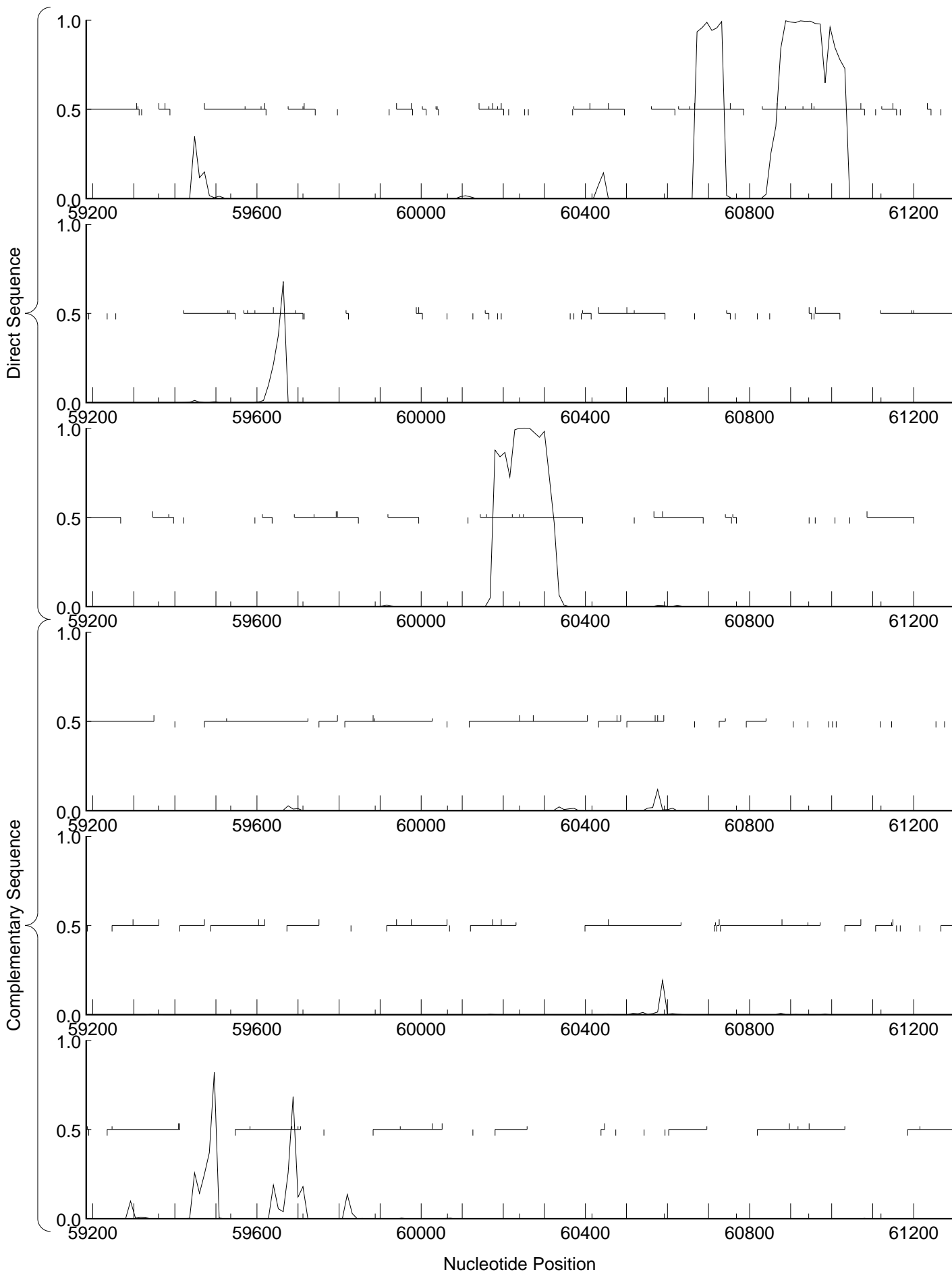
complete sequence, 80309 bp including 11-base 3' overhang (ACCCATGCAA), Cluster M2, Order 4, Window 96, Step 12, 26/40

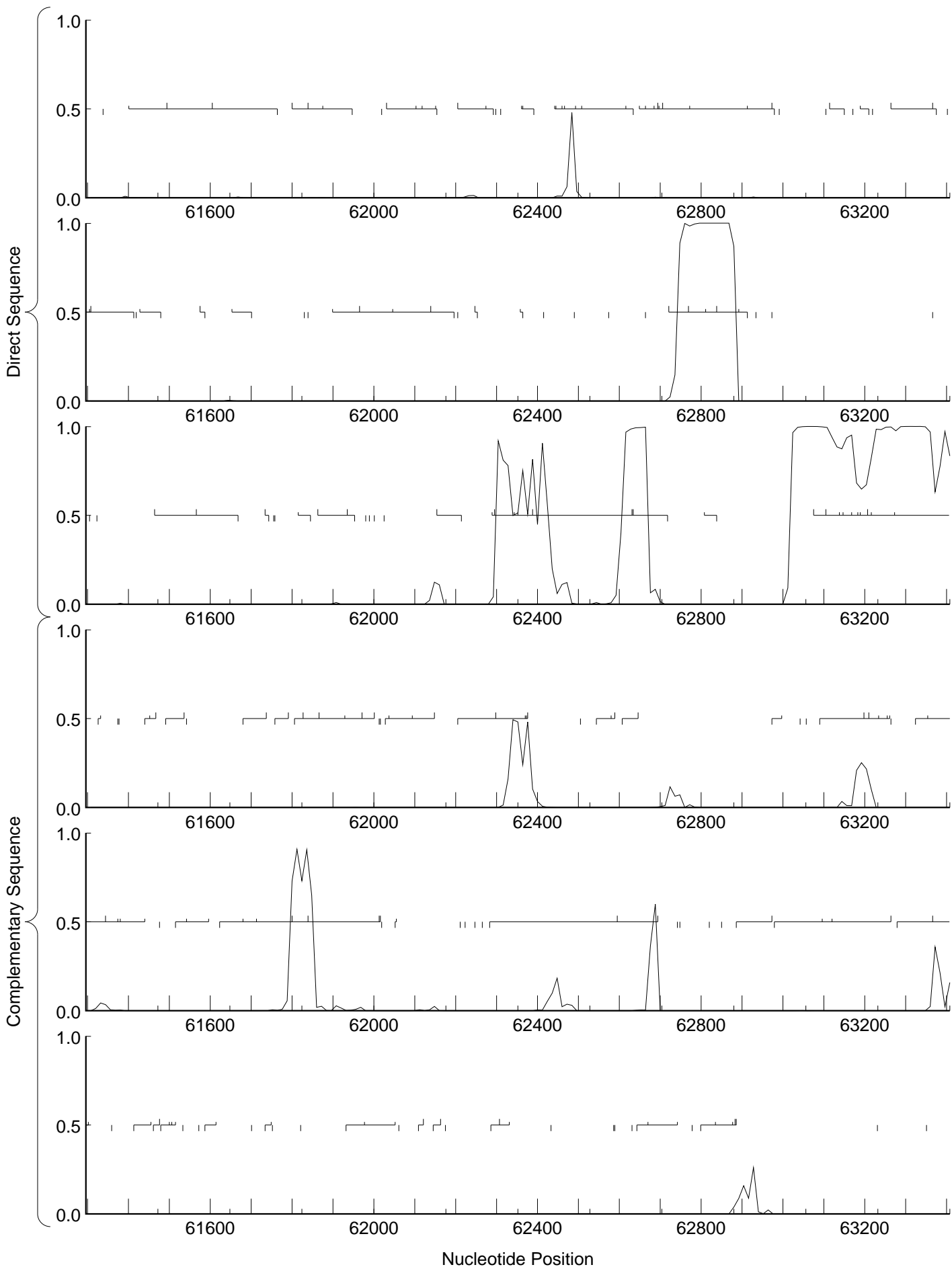


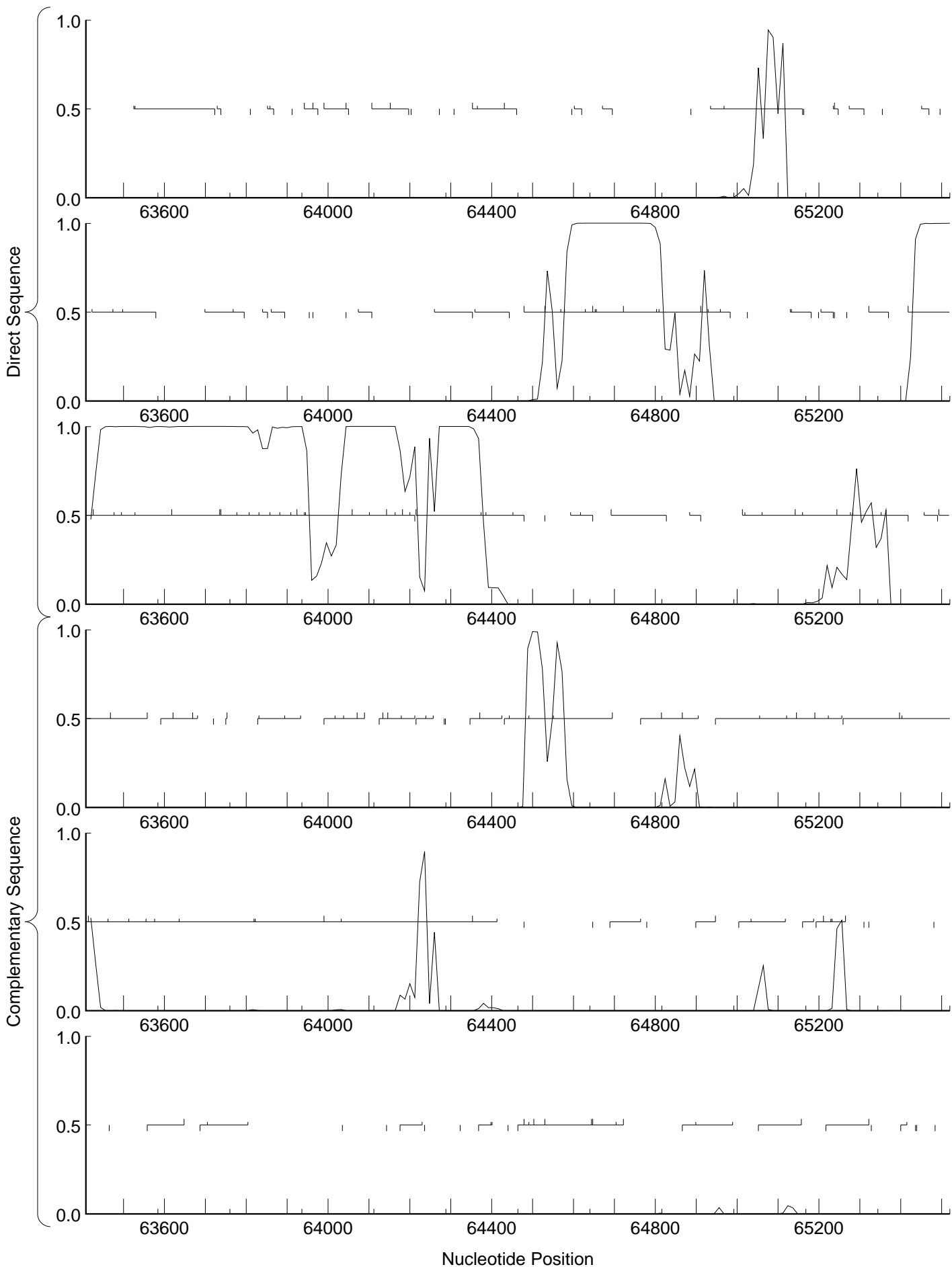




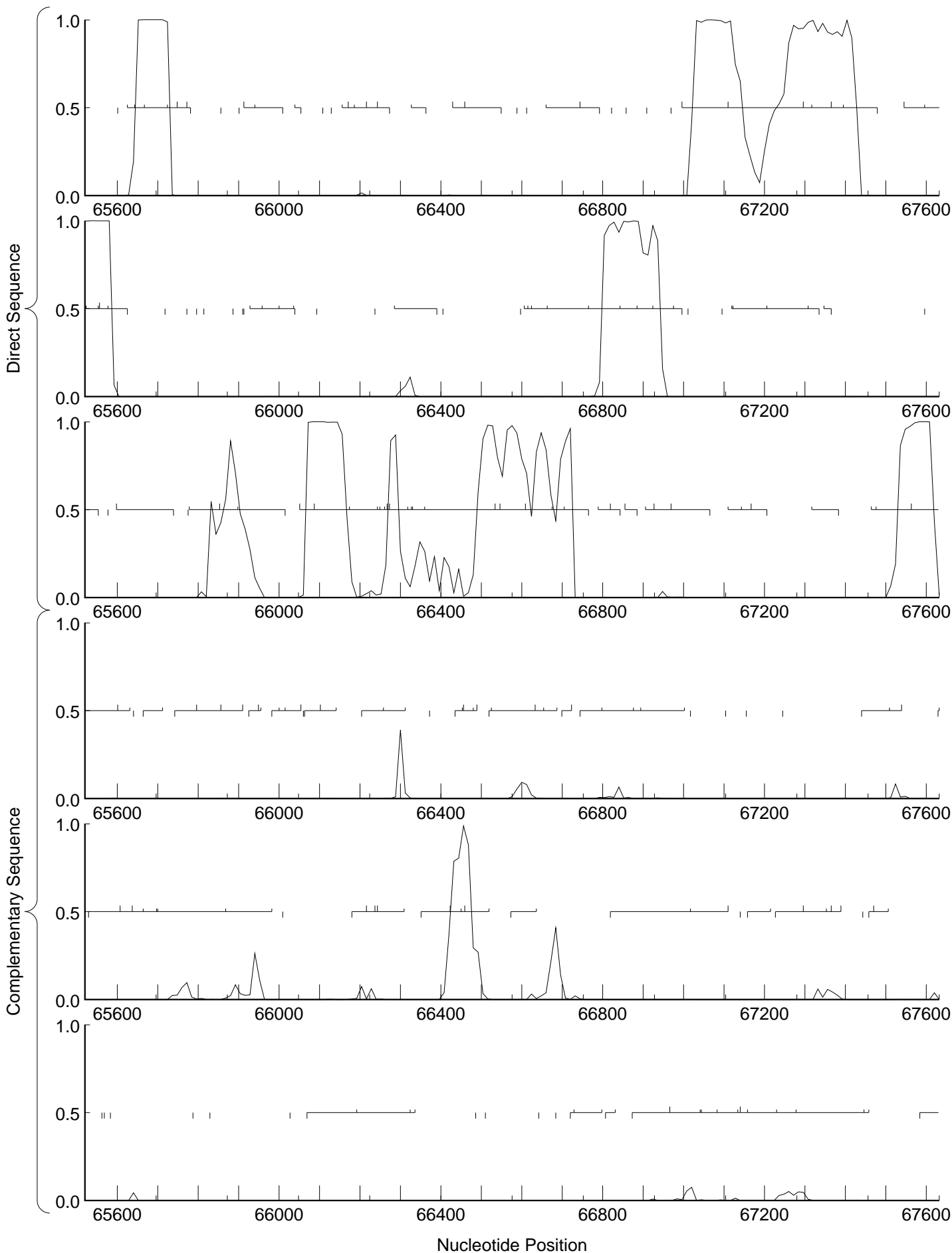


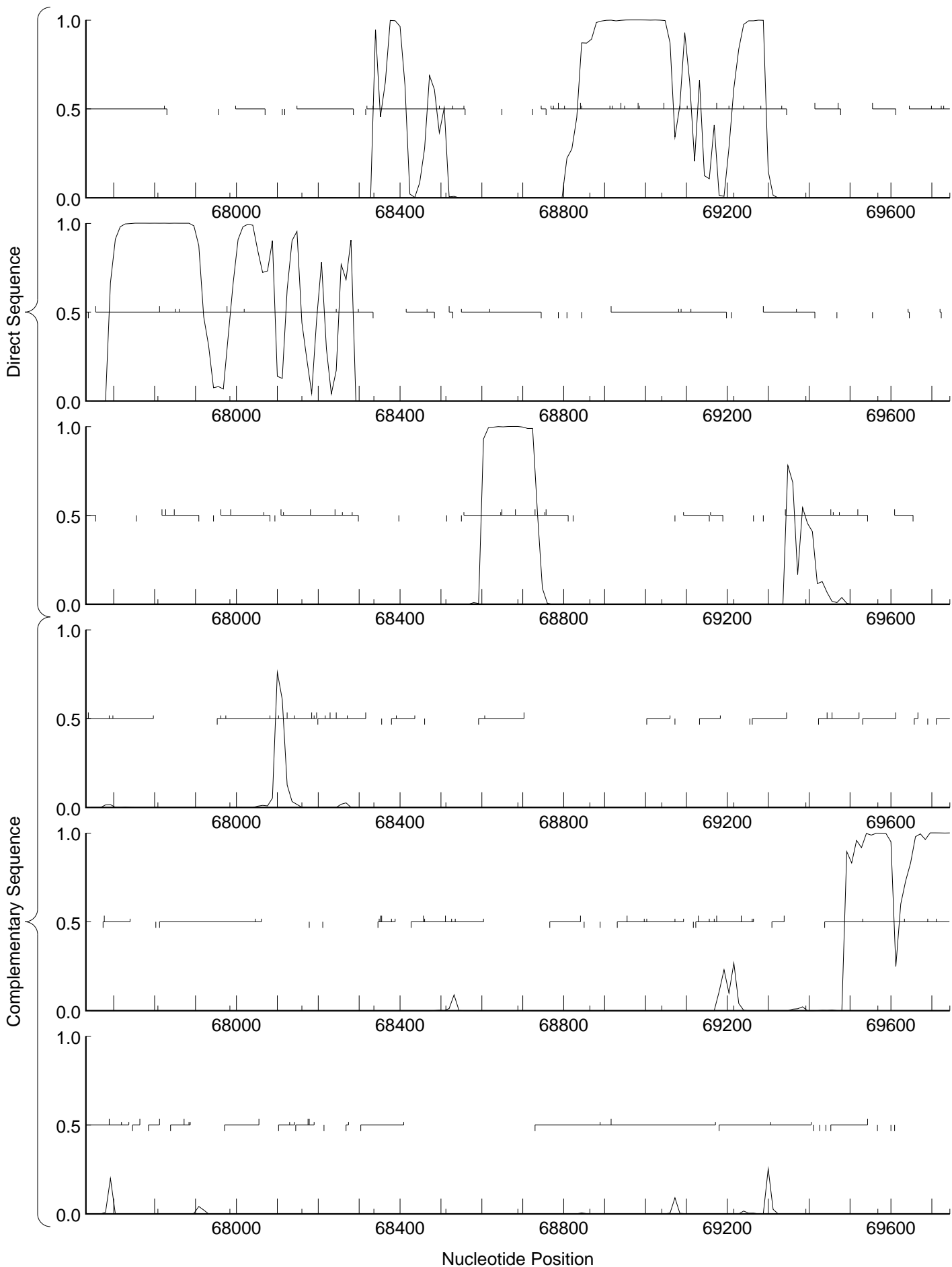






complete sequence, 80309 bp including 11-base 3' overhang (ACCCATGCAA), Cluster M2, Order 4, Window 96, Step 12, 33/40





complete sequence, 80309 bp including 11-base 3' overhang (ACCCCATGCAA), Cluster M2, Order 4, Window 96, Step 12, 35/40

