

GeneMark

Version 2.5p (09.08.06)

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PROGRAM INFORMATION

Sequence : Gordonia phage Cucurbita complete sequence, 93686 bp including 10-base 3' overhang (CGCGACG
Analysis Date : 9/15/24 at 5:42:34
Pages : 46
Sequence Length : 93686 bp
GC Content : 61.98%

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,

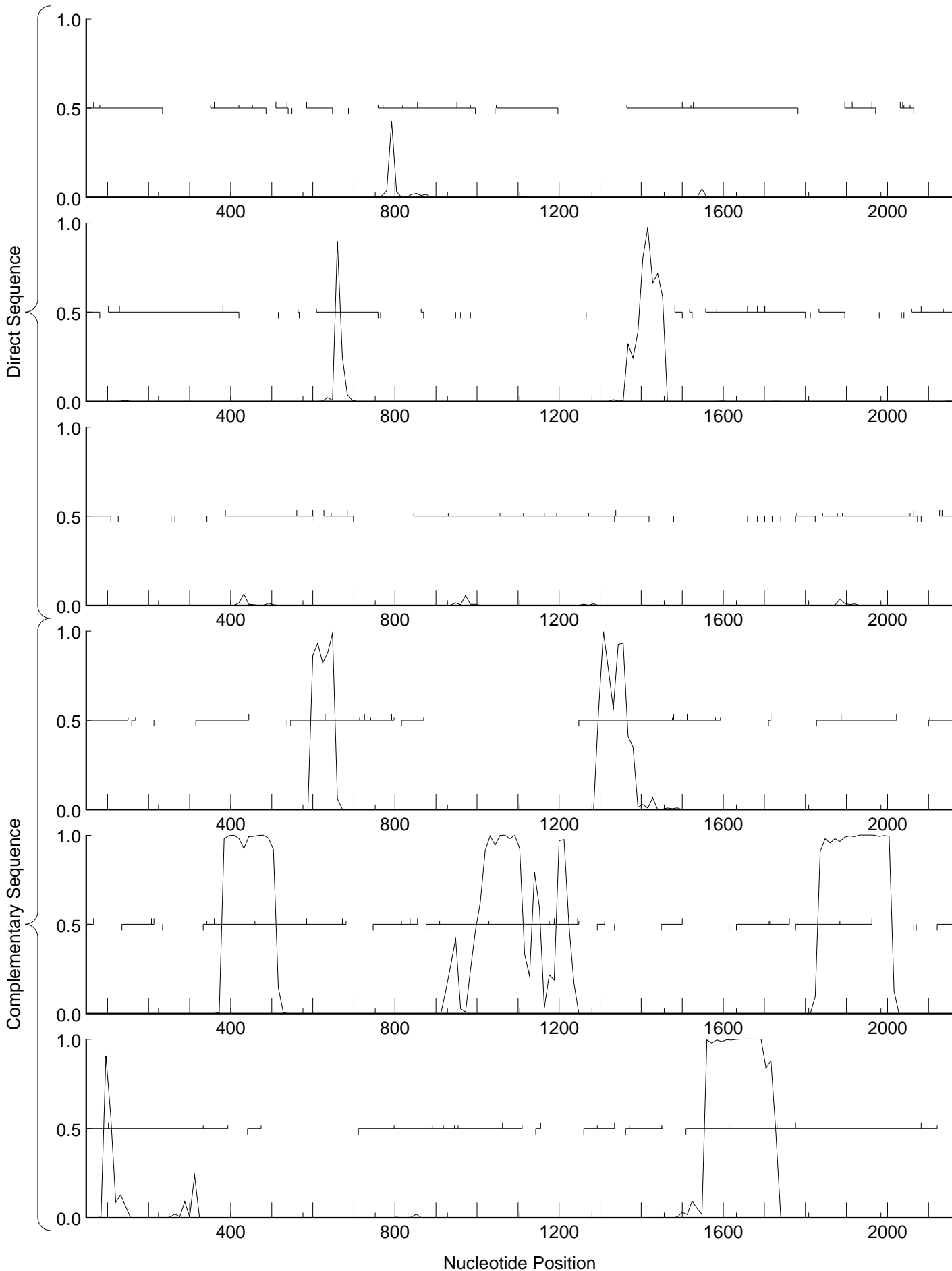
MATRIX INFORMATION

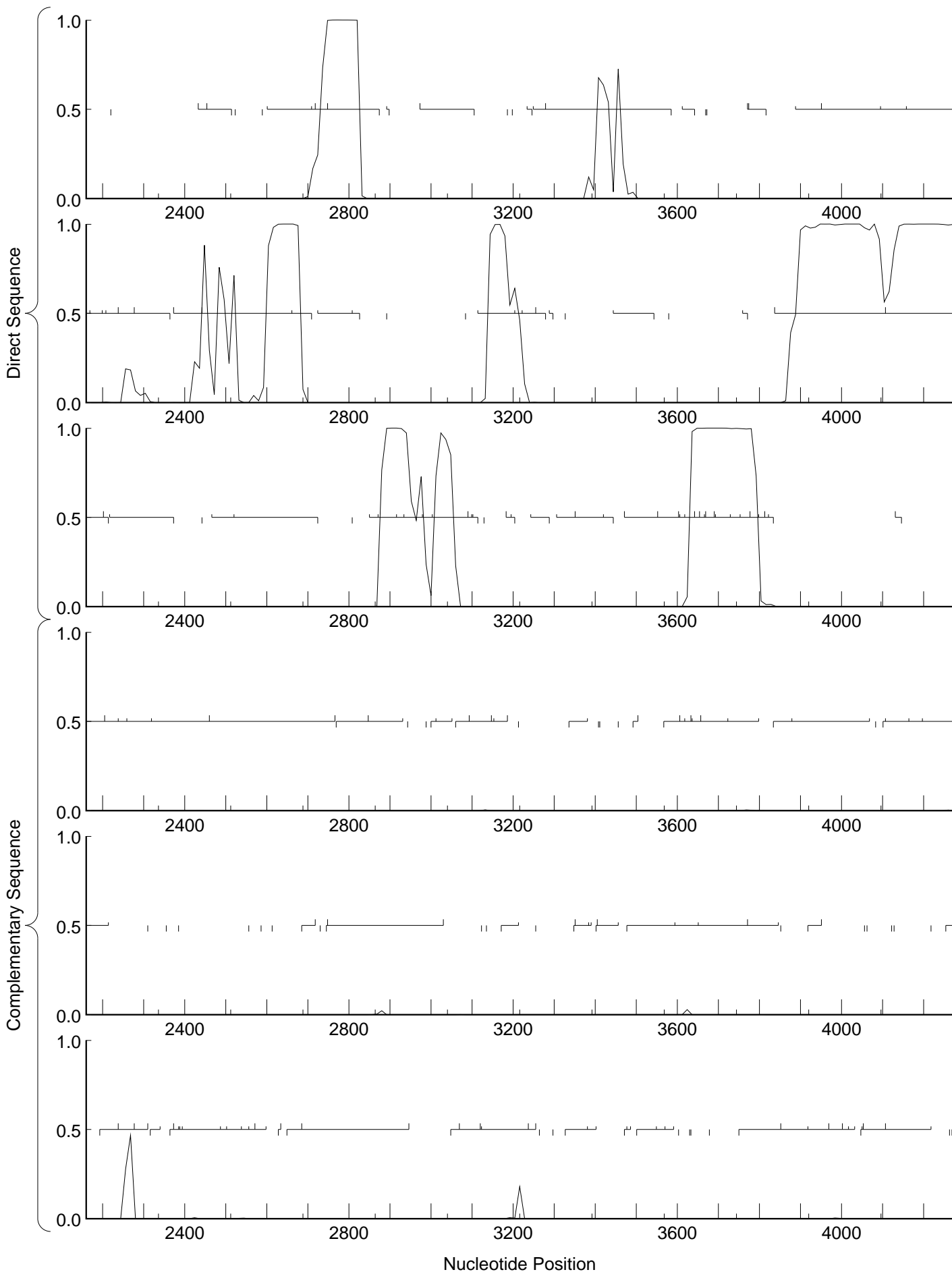
Matrix : -
Author : -
Order : 4

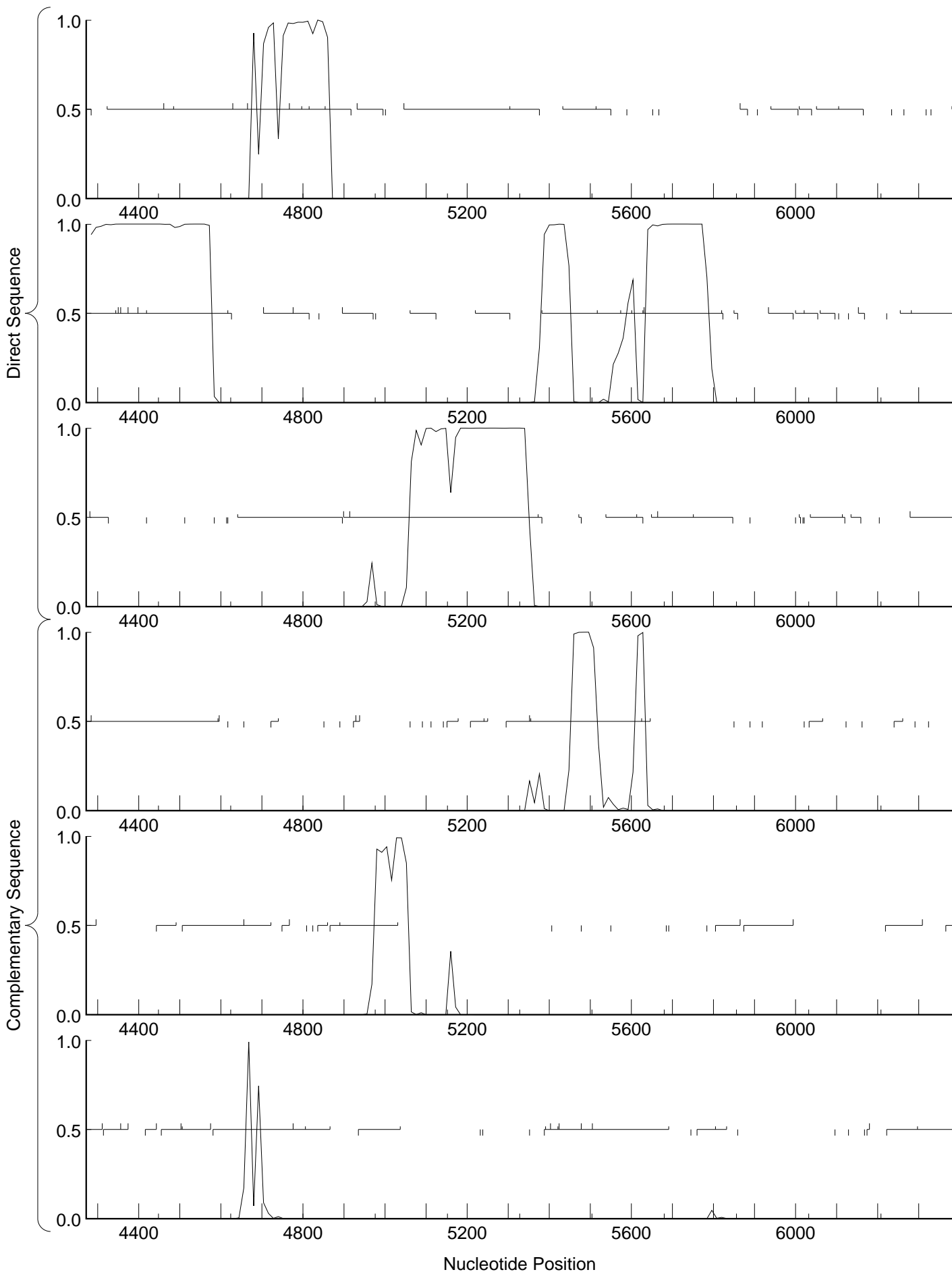
Send questions / comments to:
Dr. M. Borodovsky
Georgia Institute of Technology
School of Biology
Atlanta, GA 30332-0230

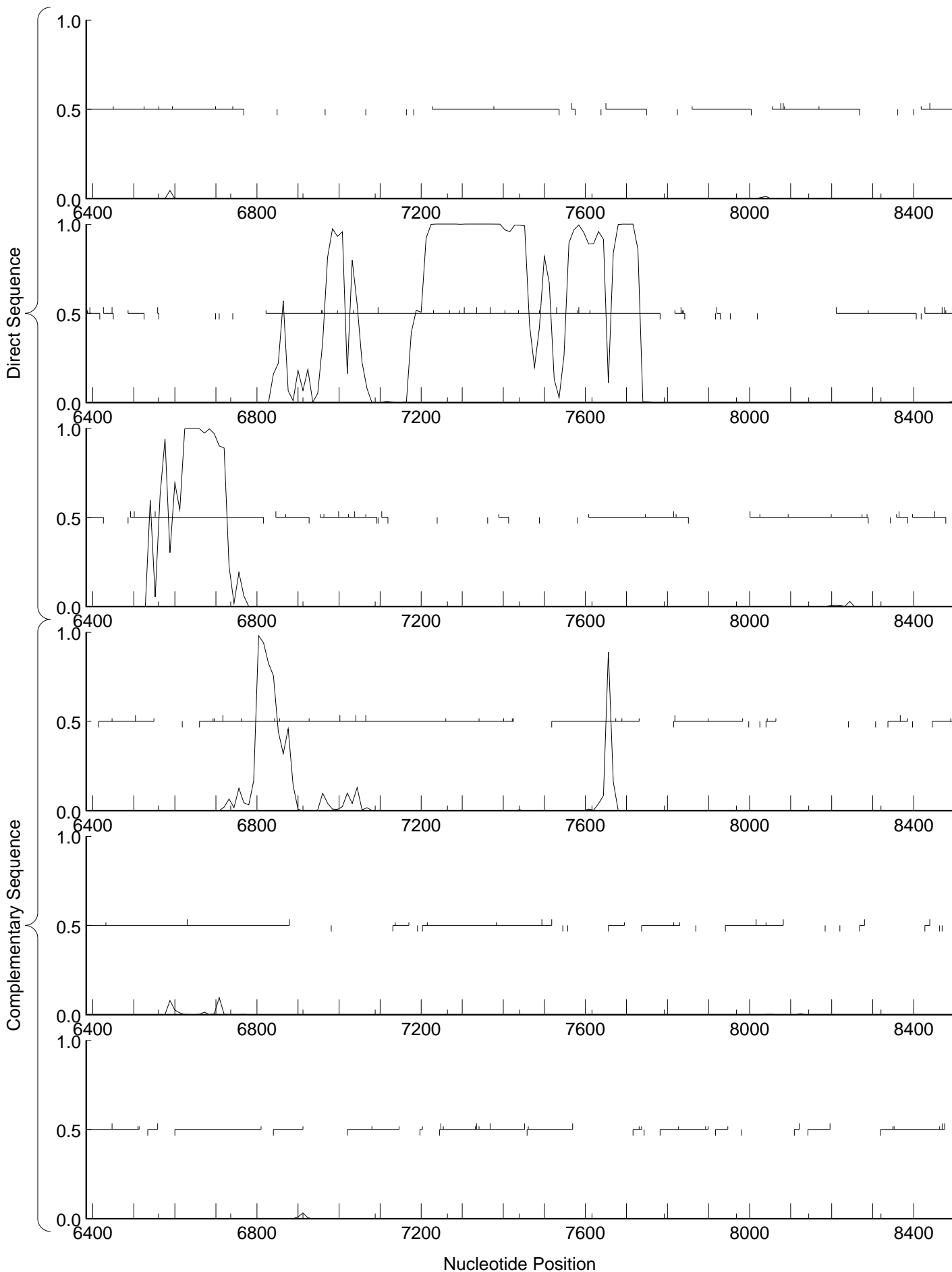
Matrix notes & comments

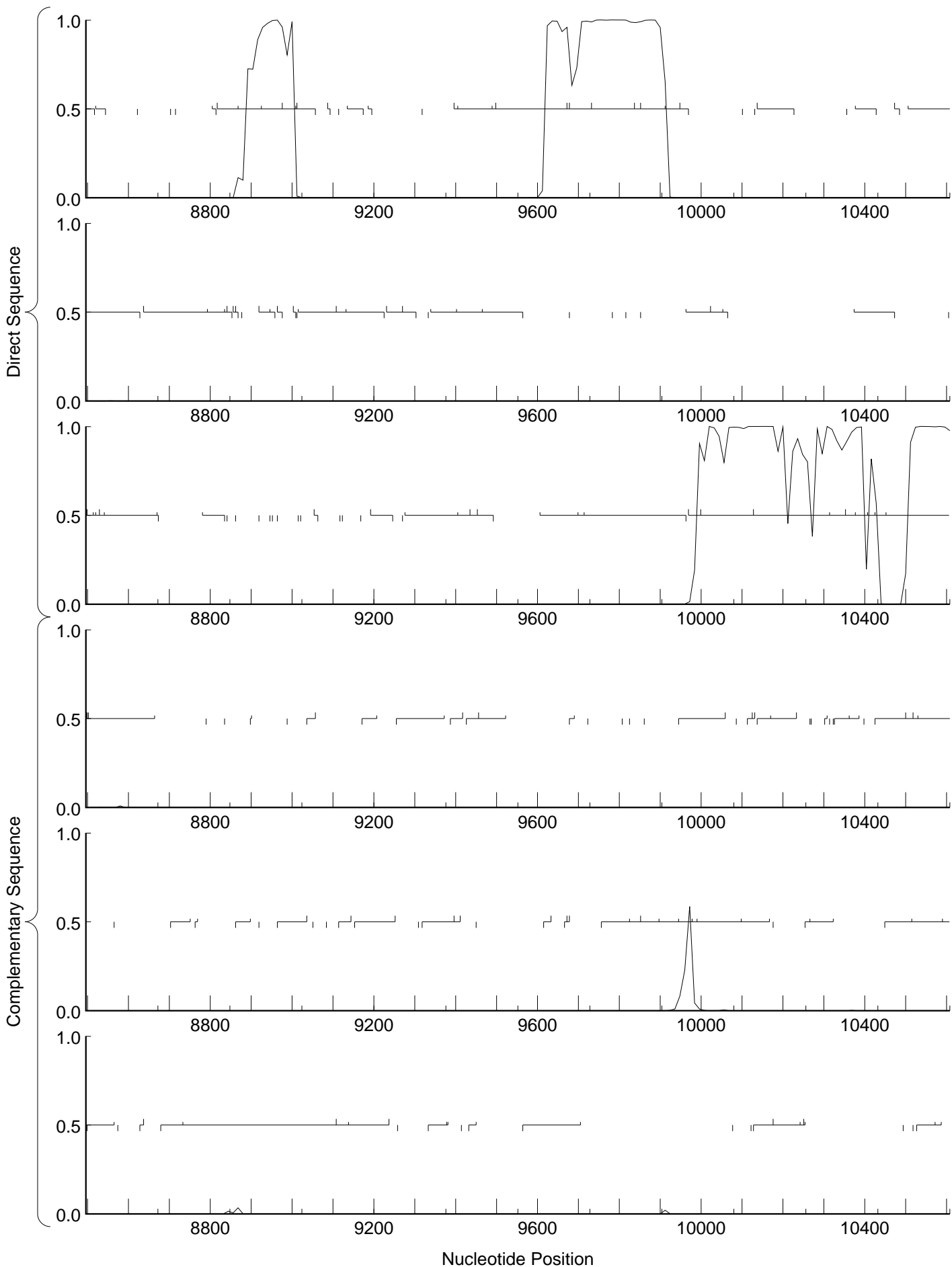
Training set derived by GeneMarkS, 4.30 October 2014
Sat Sep 8 00:36:14 2018

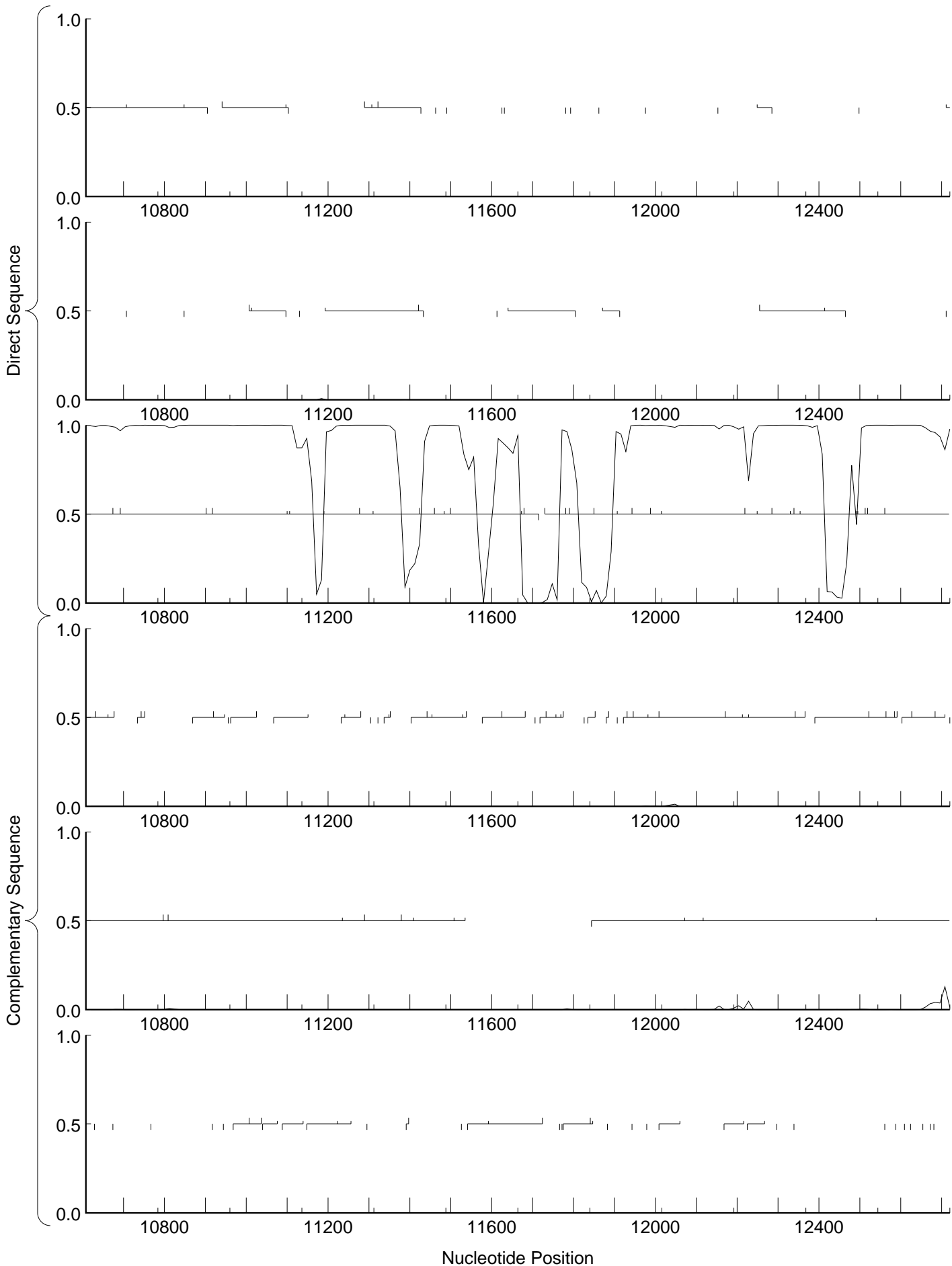




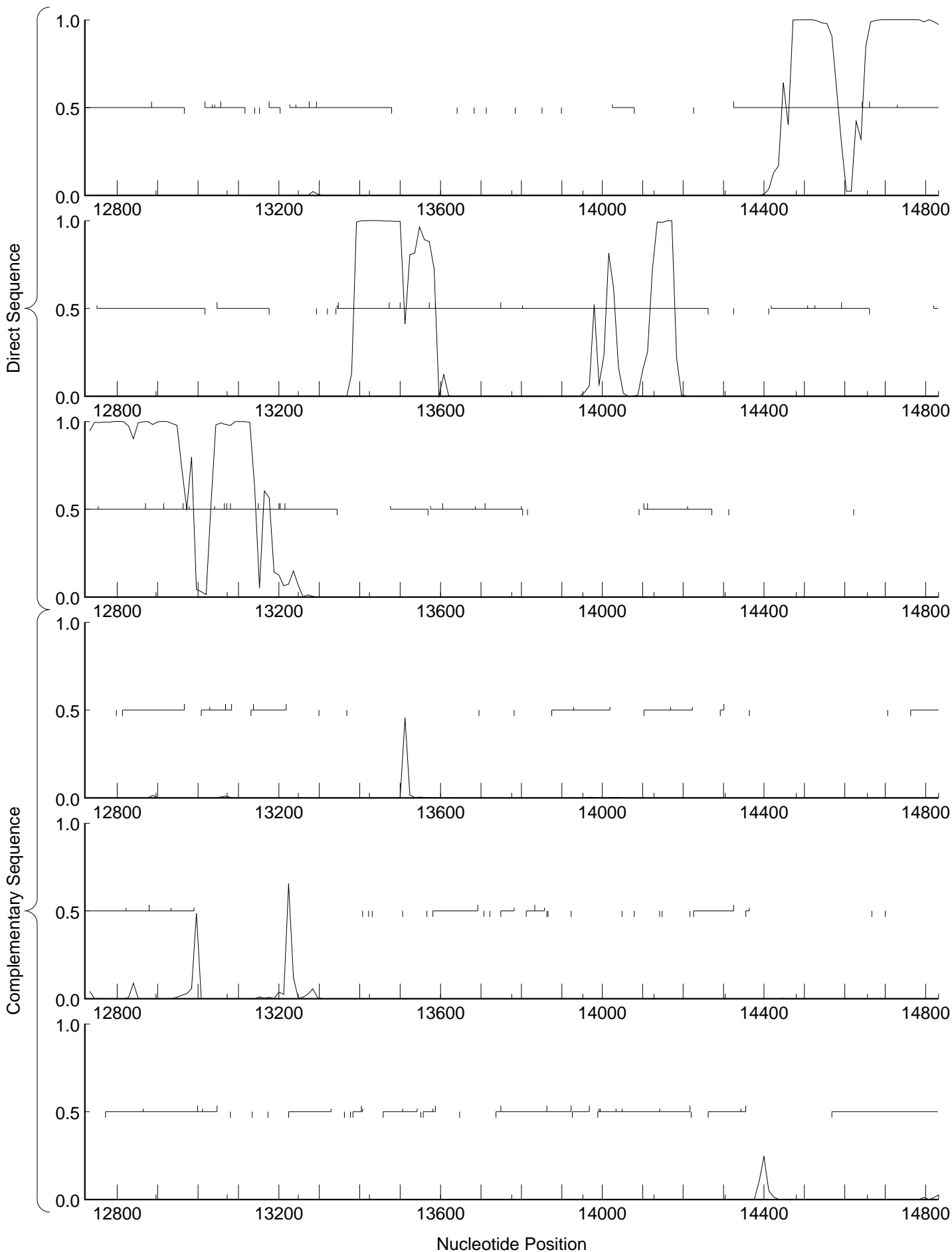




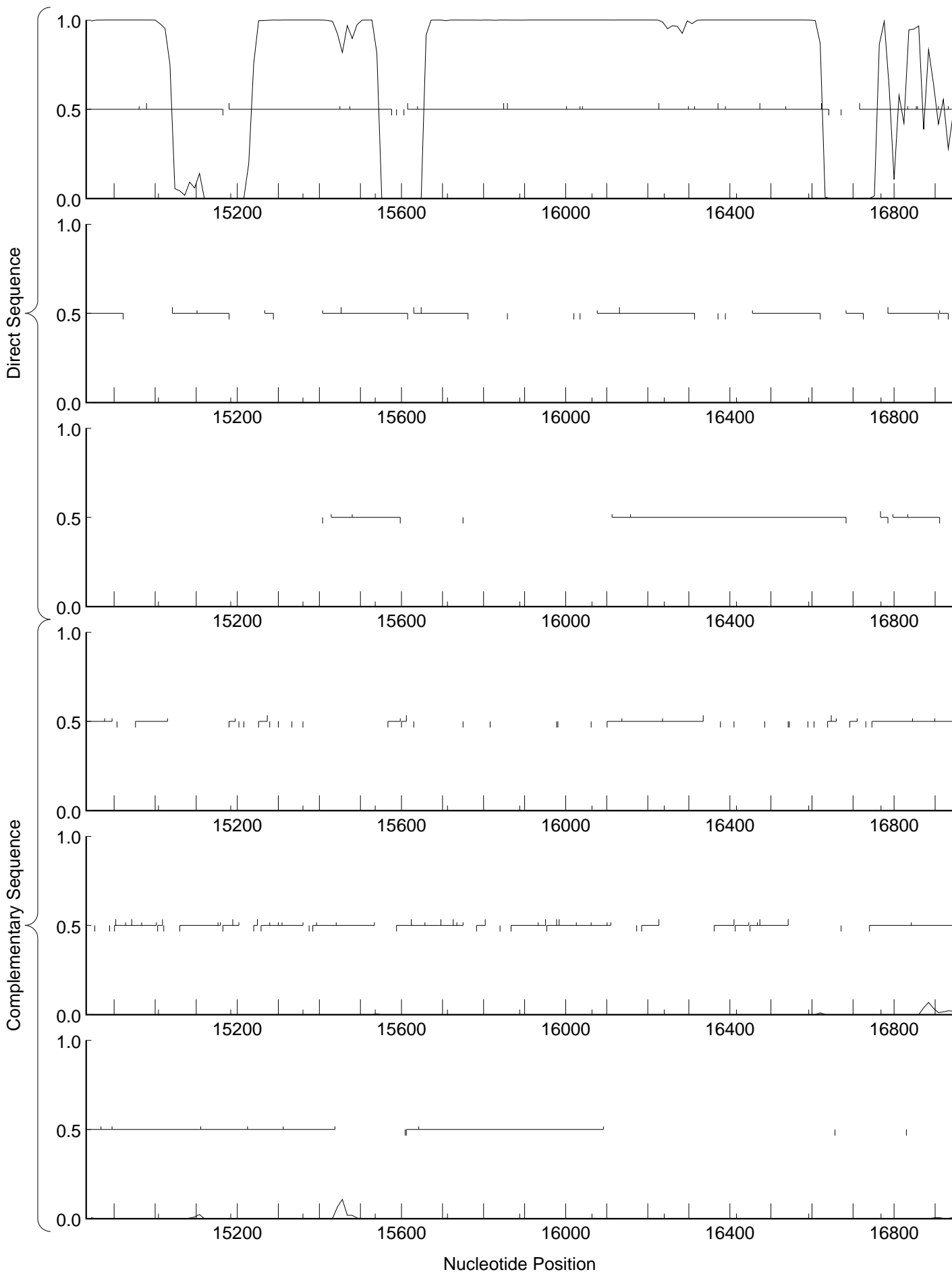


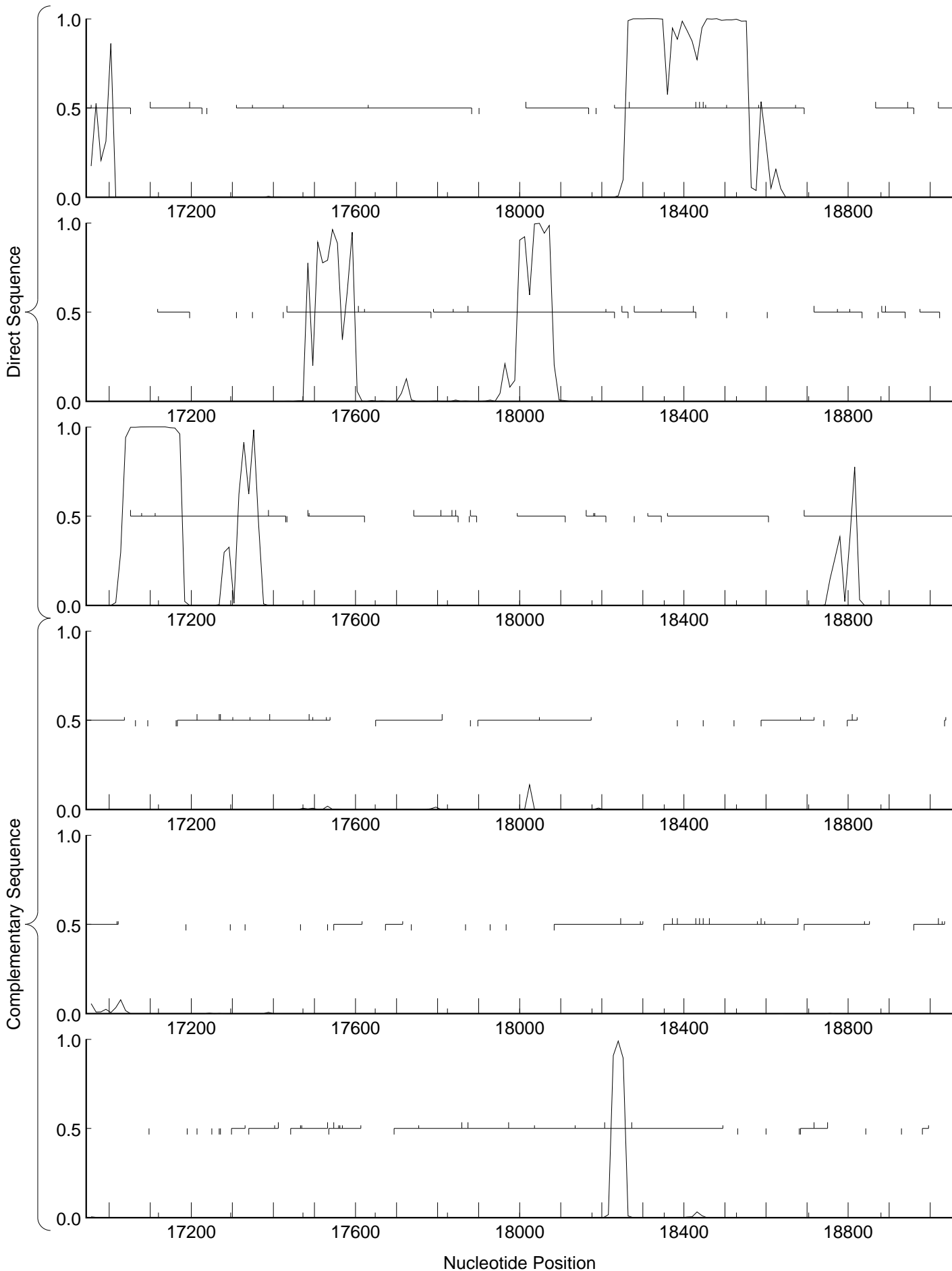


a complete sequence, 93686 bp including 10-base 3' overhang (CGCGACGCTC), Cluster CQ, Order 4, Window 96, Step 12, 8/46

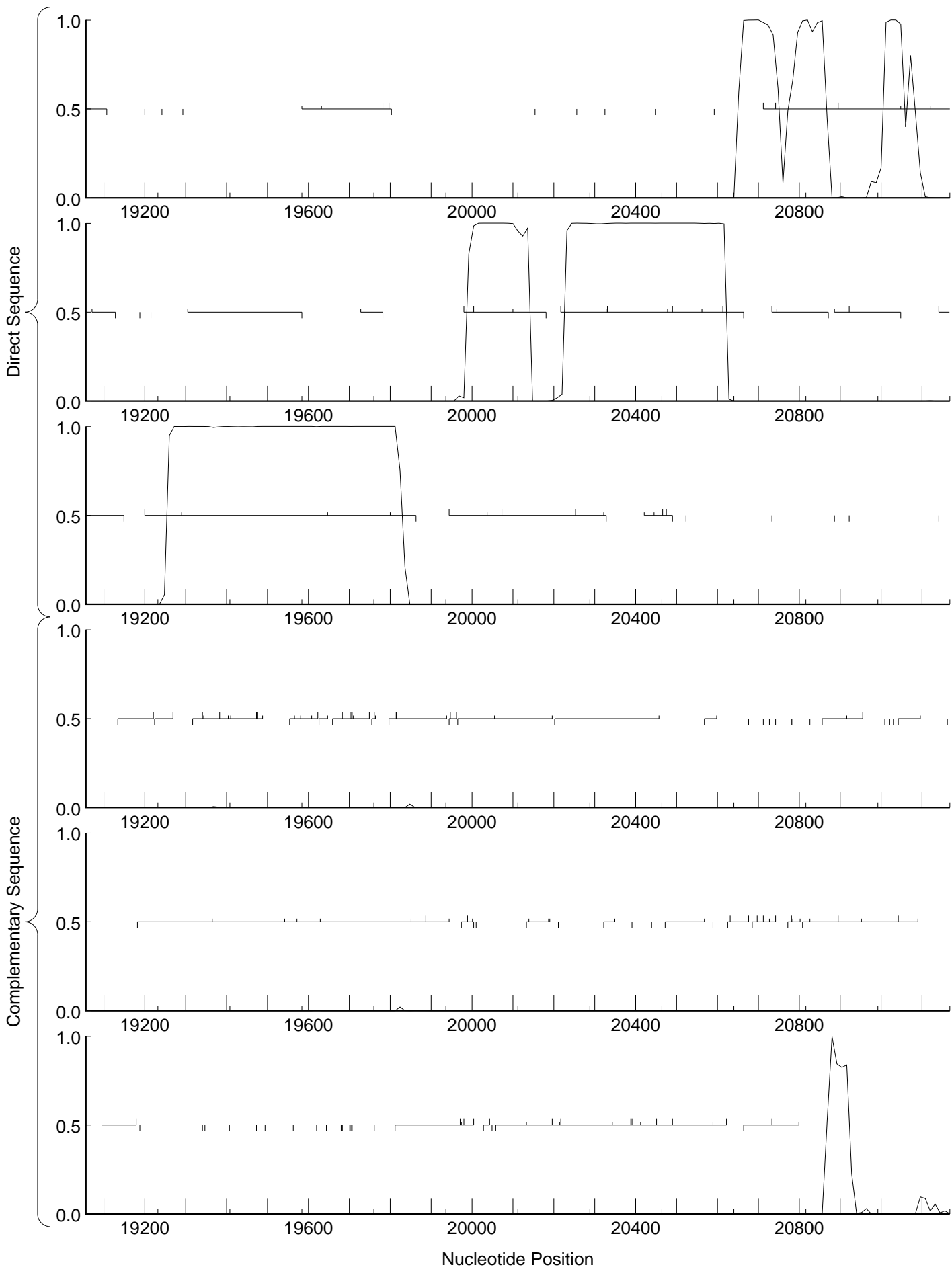


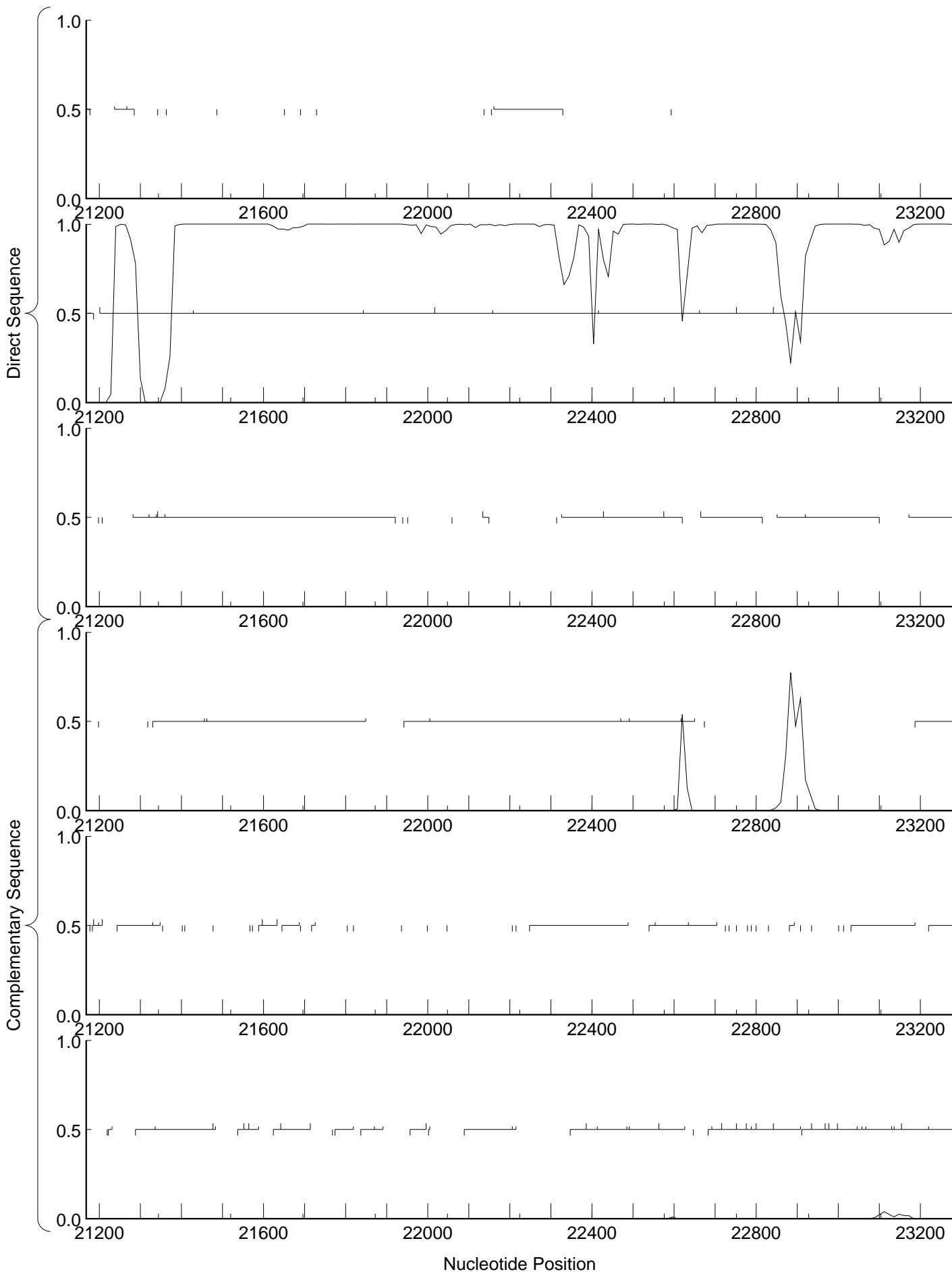
a complete sequence, 93686 bp including 10-base 3' overhang (CGCGACGCTC), Cluster CQ, Order 4, Window 96, Step 12, 9/46

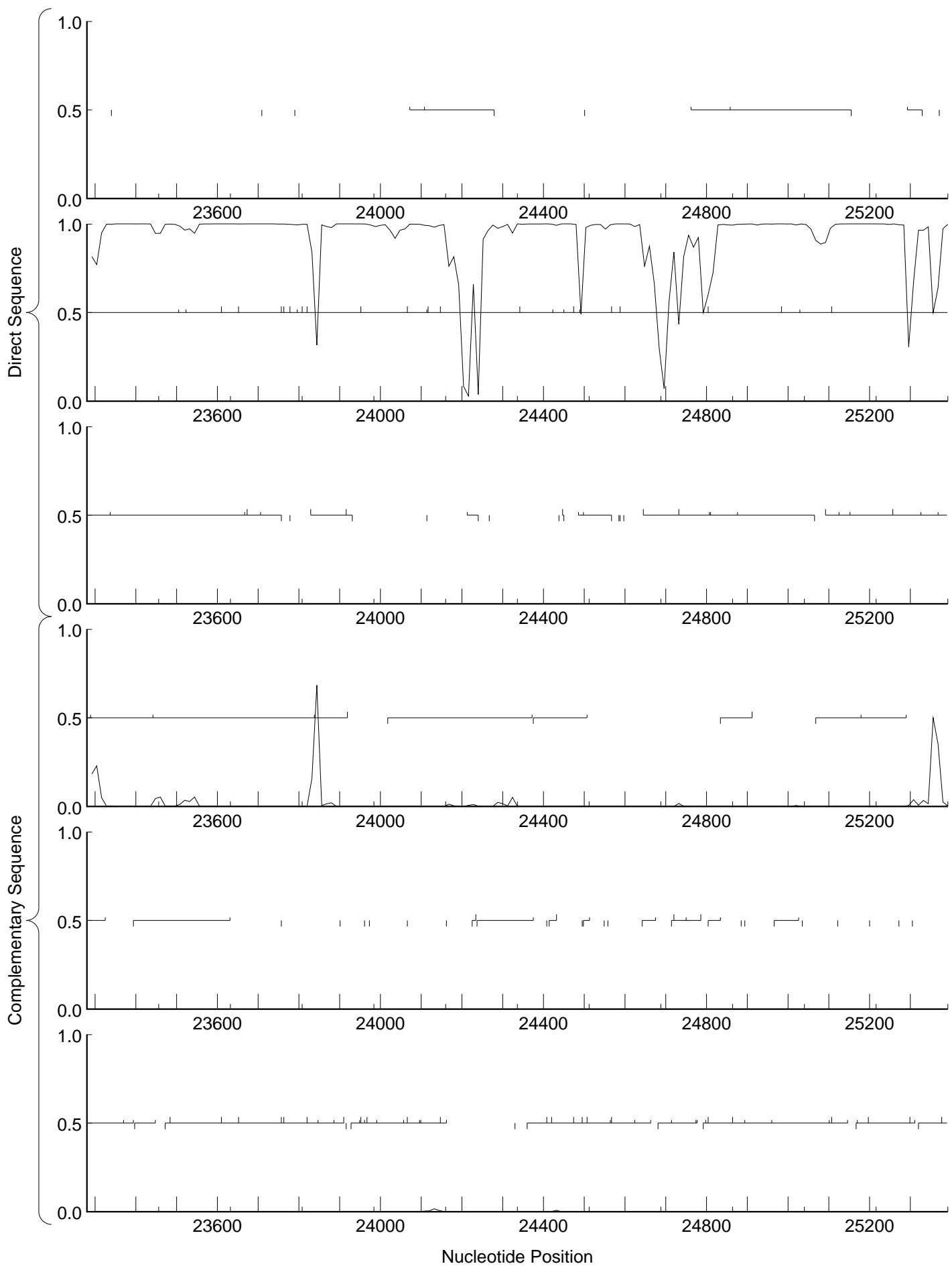


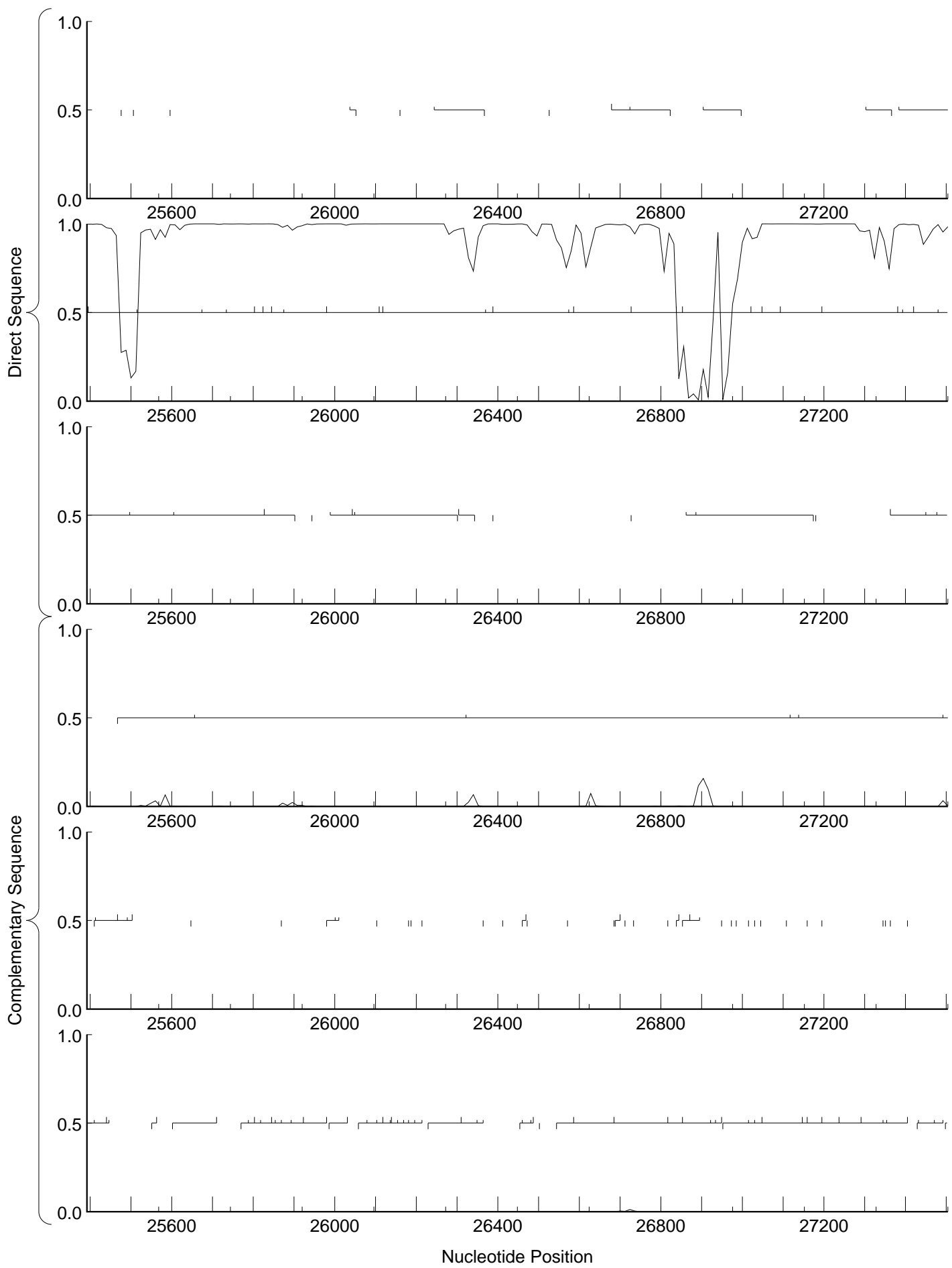


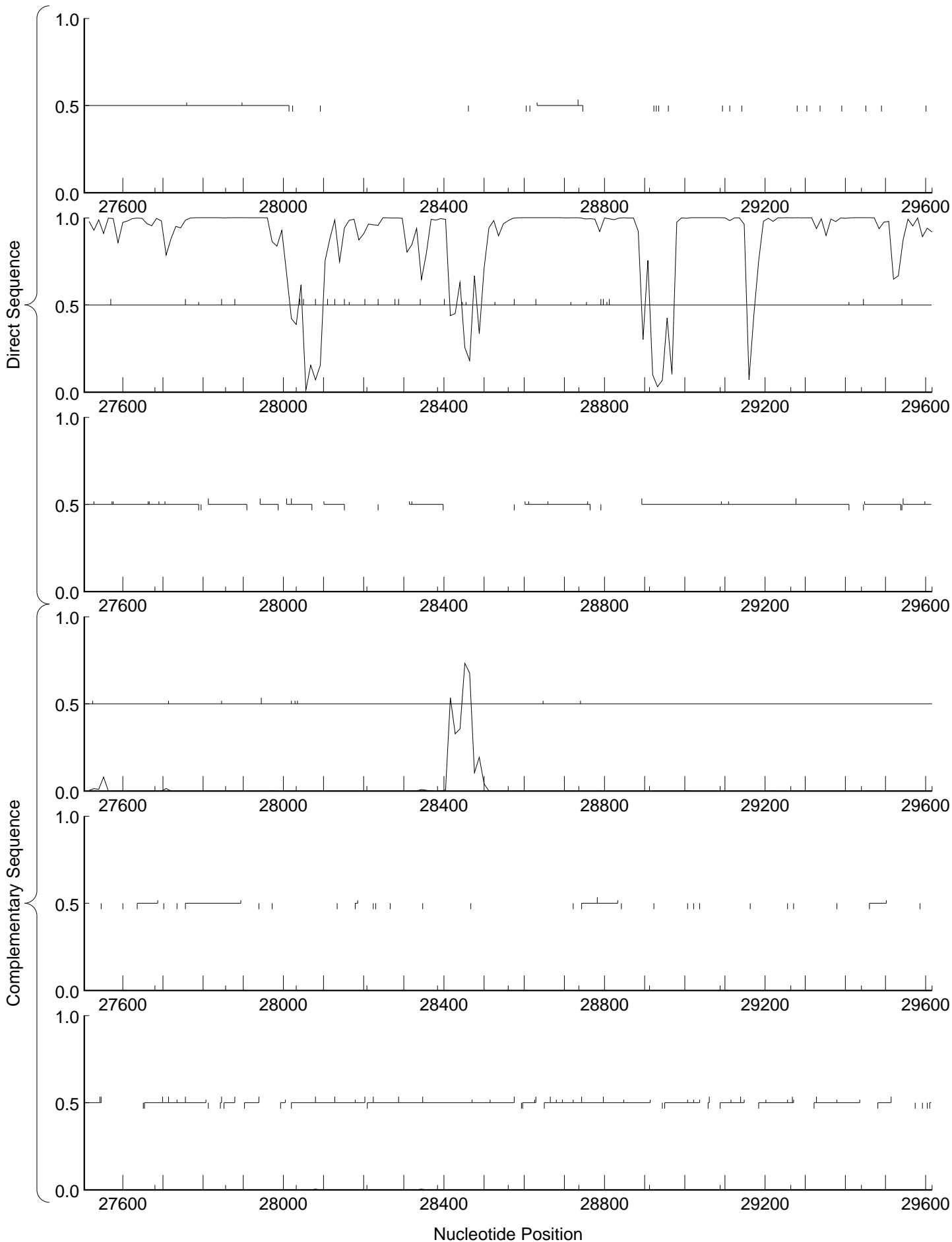
complete sequence, 93686 bp including 10-base 3' overhang (CGCGACGCTC), Cluster CQ, Order 4, Window 96, Step 12, 11/46



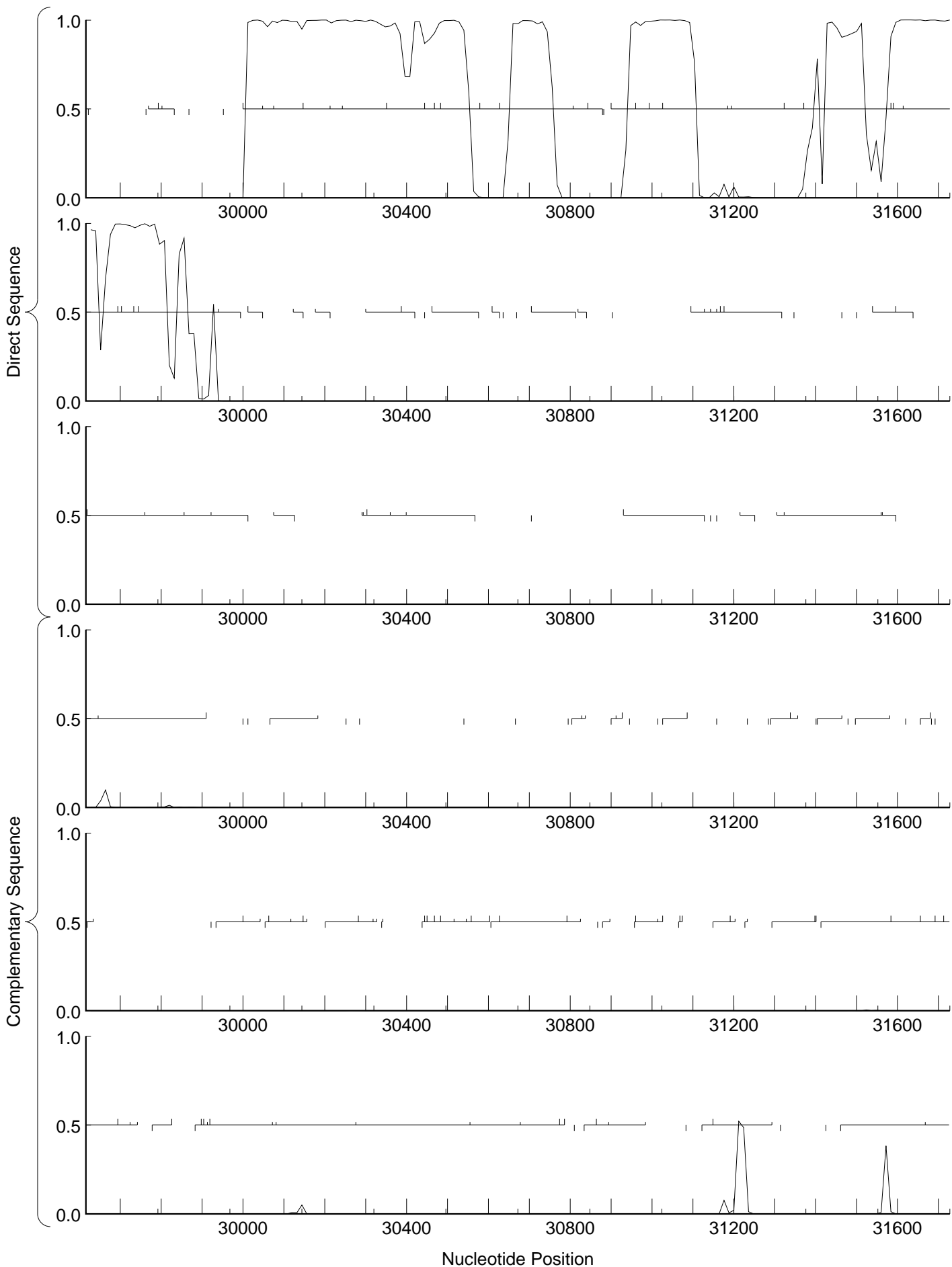


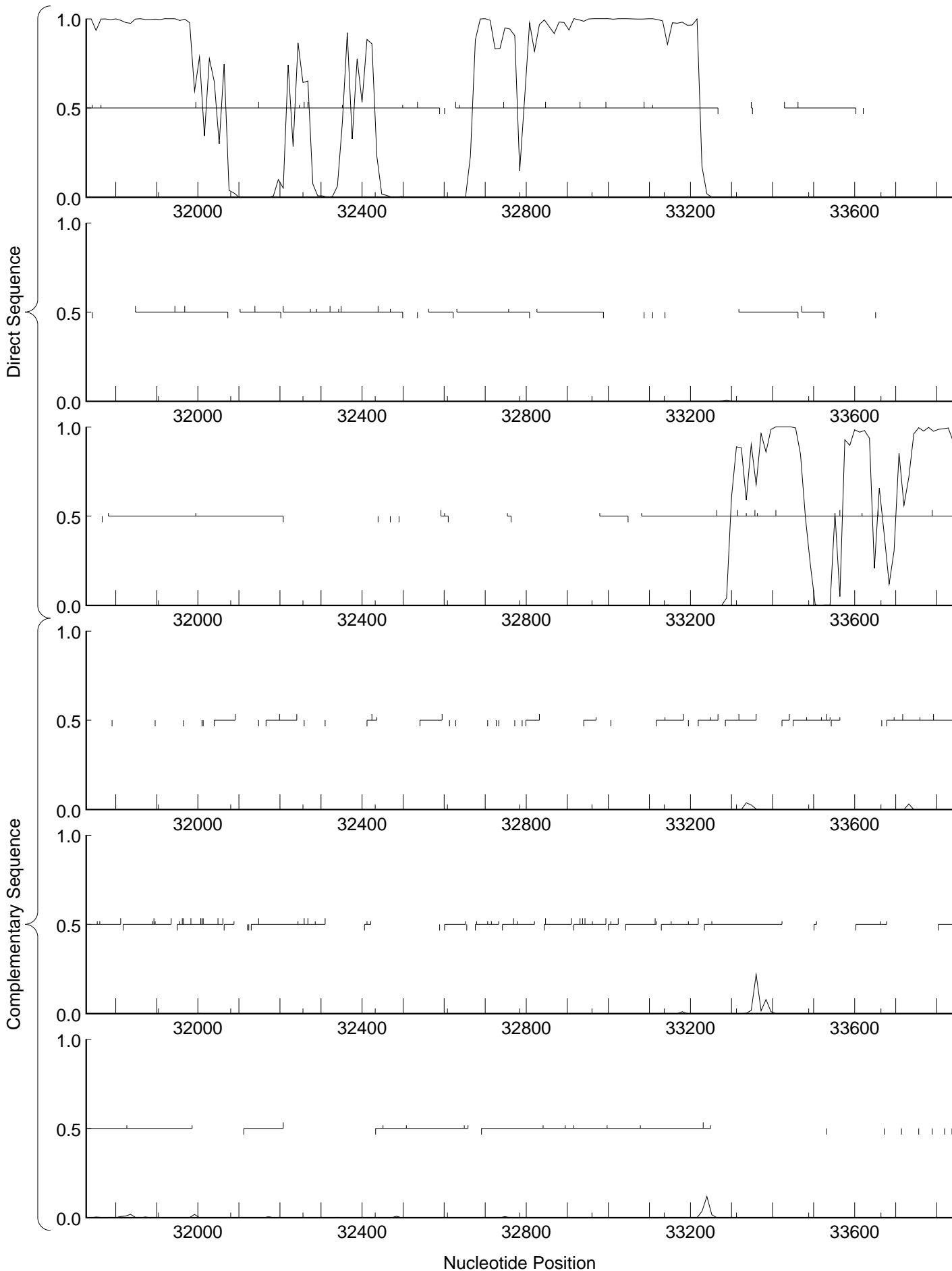


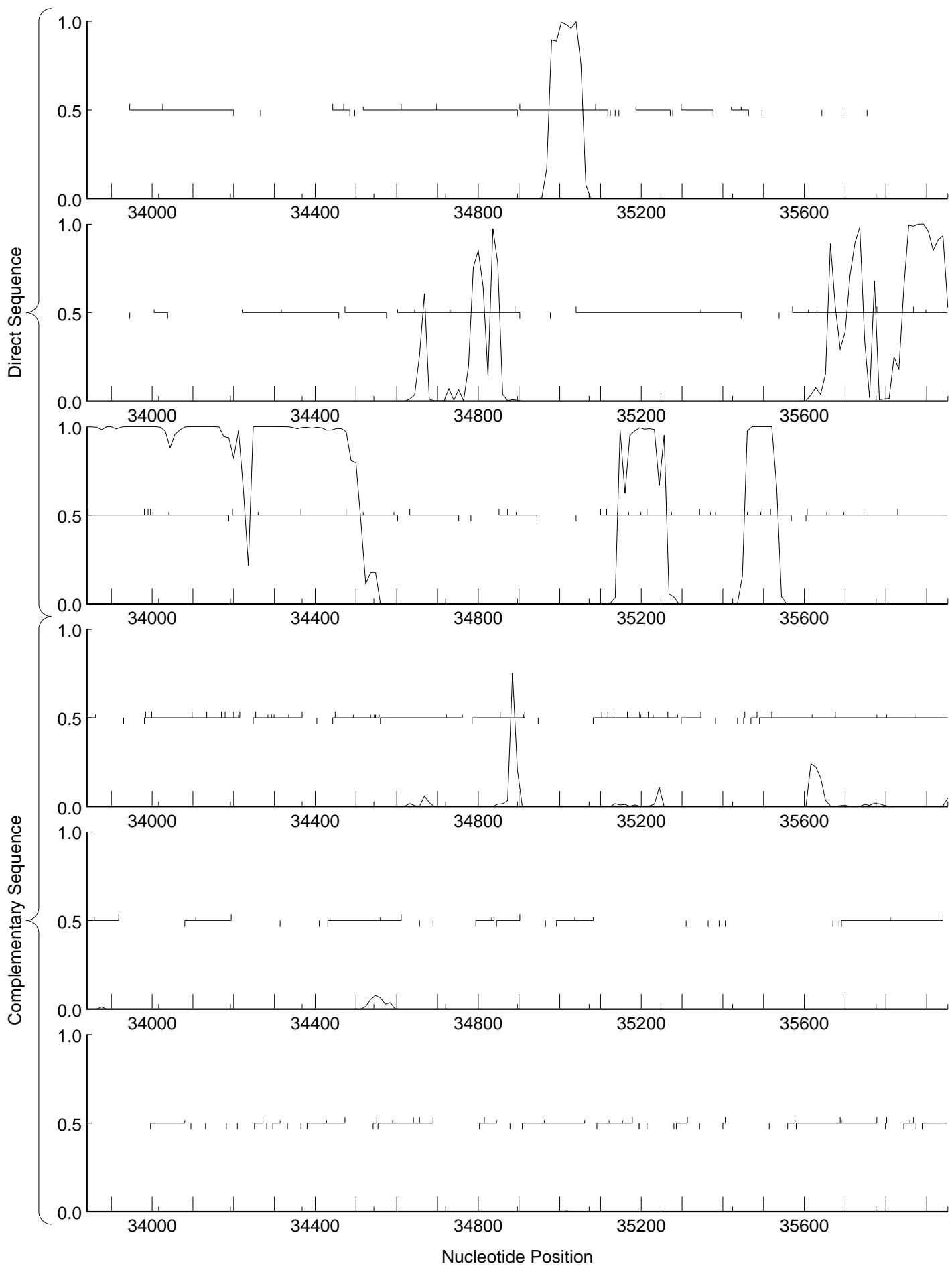




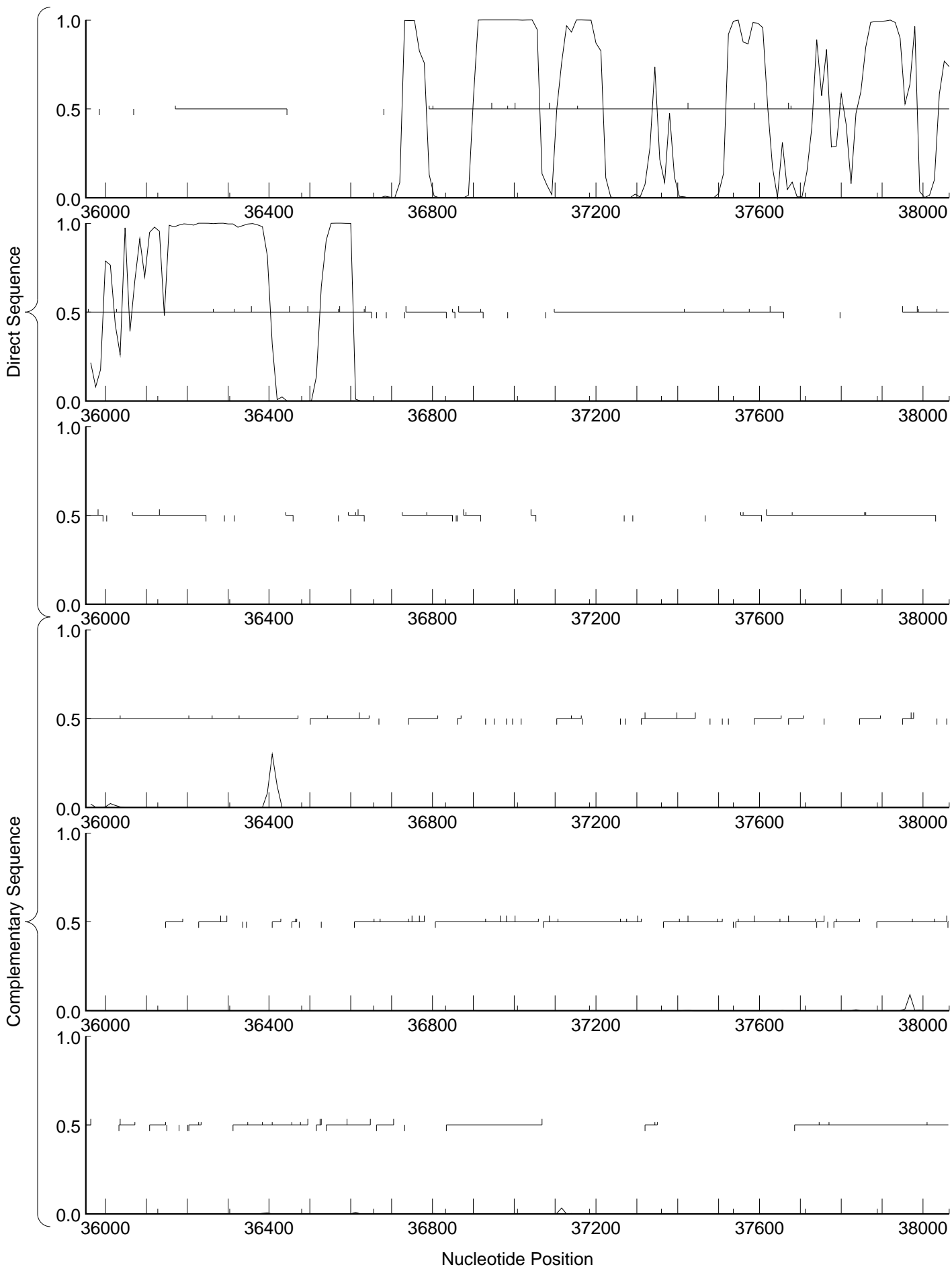
complete sequence, 93686 bp including 10-base 3' overhang (CGCGACGCTC), Cluster CQ, Order 4, Window 96, Step 12, 16/46



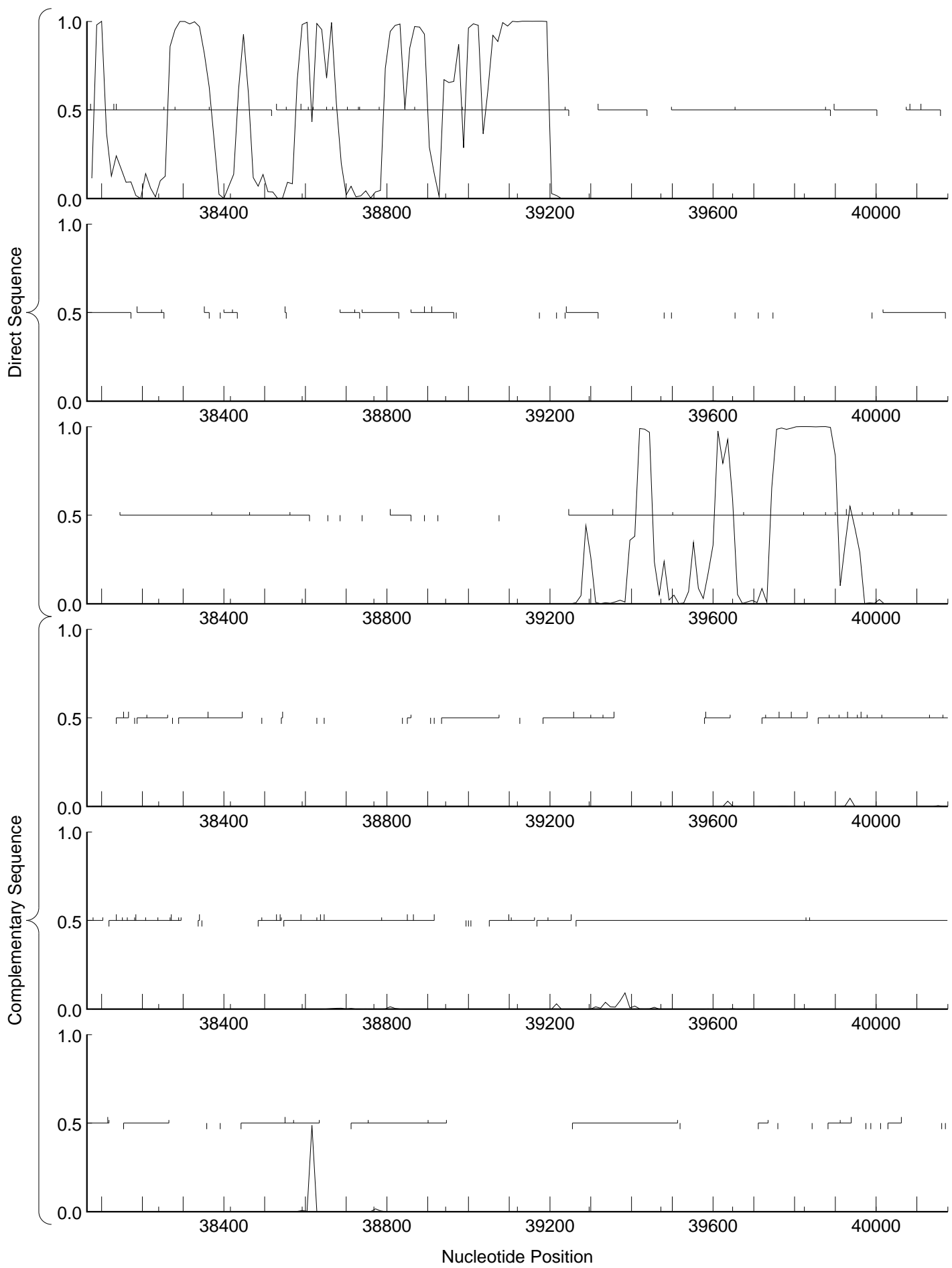


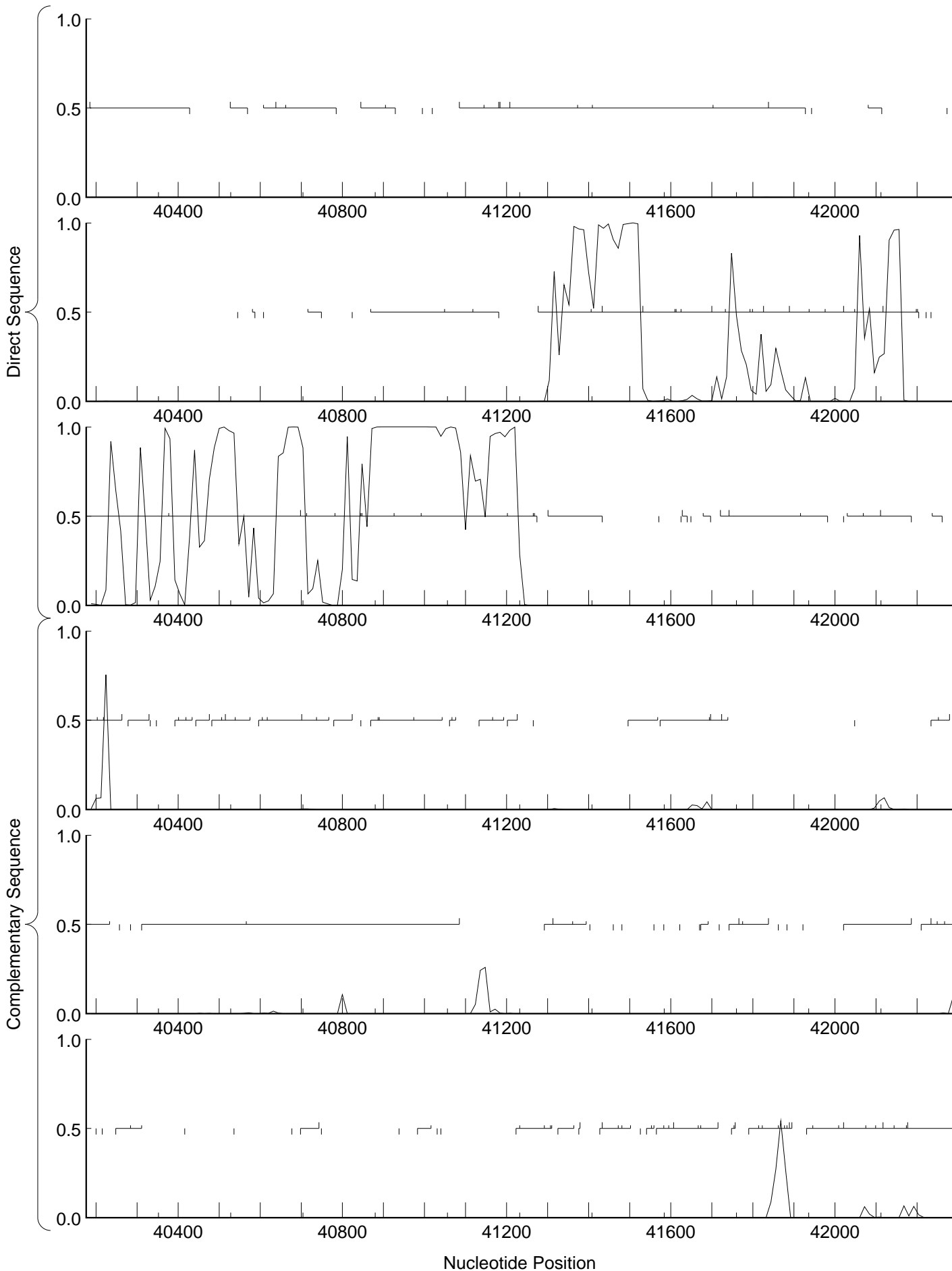


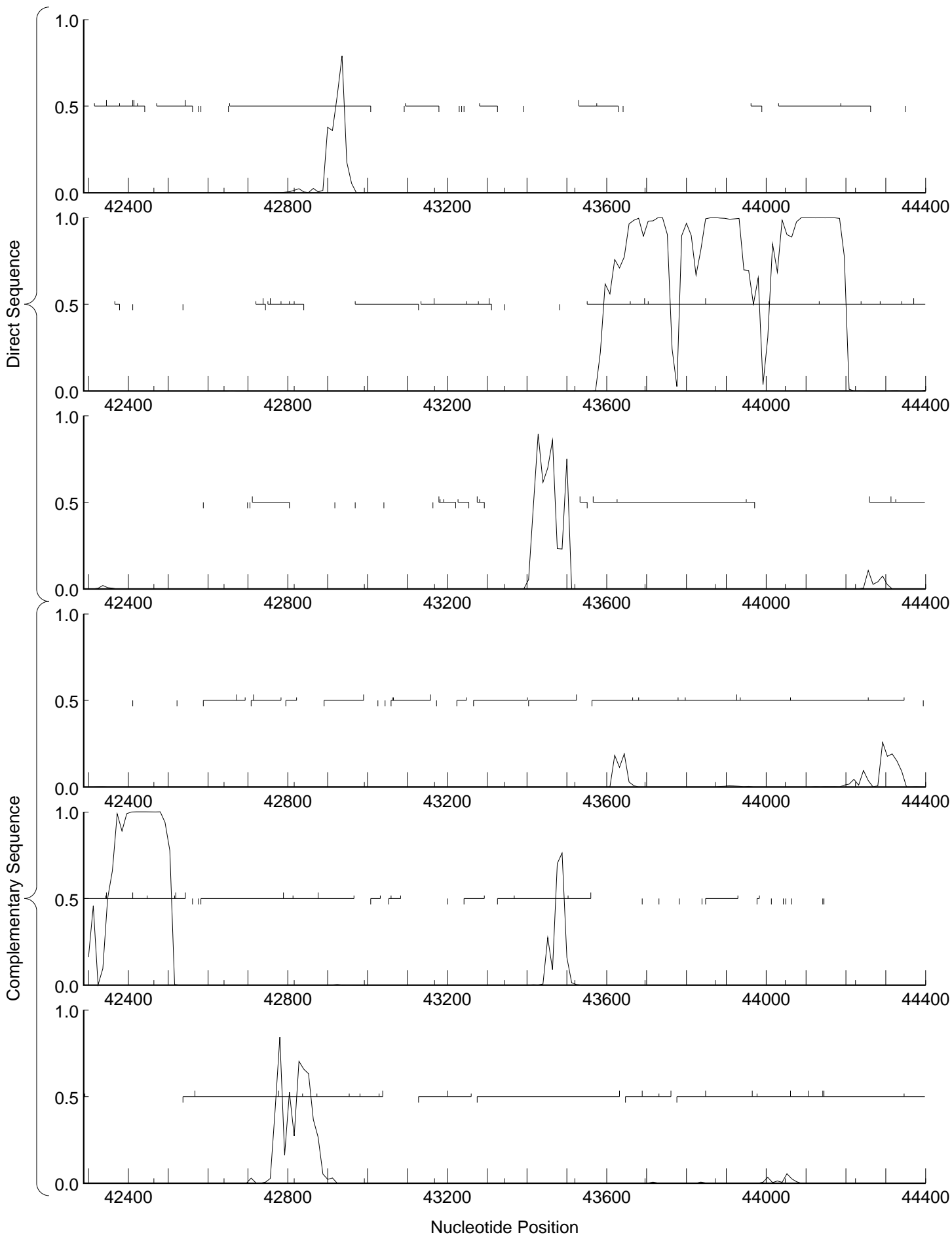
complete sequence, 93686 bp including 10-base 3' overhang (CGCGACGCTC), Cluster CQ, Order 4, Window 96, Step 12, 19/46

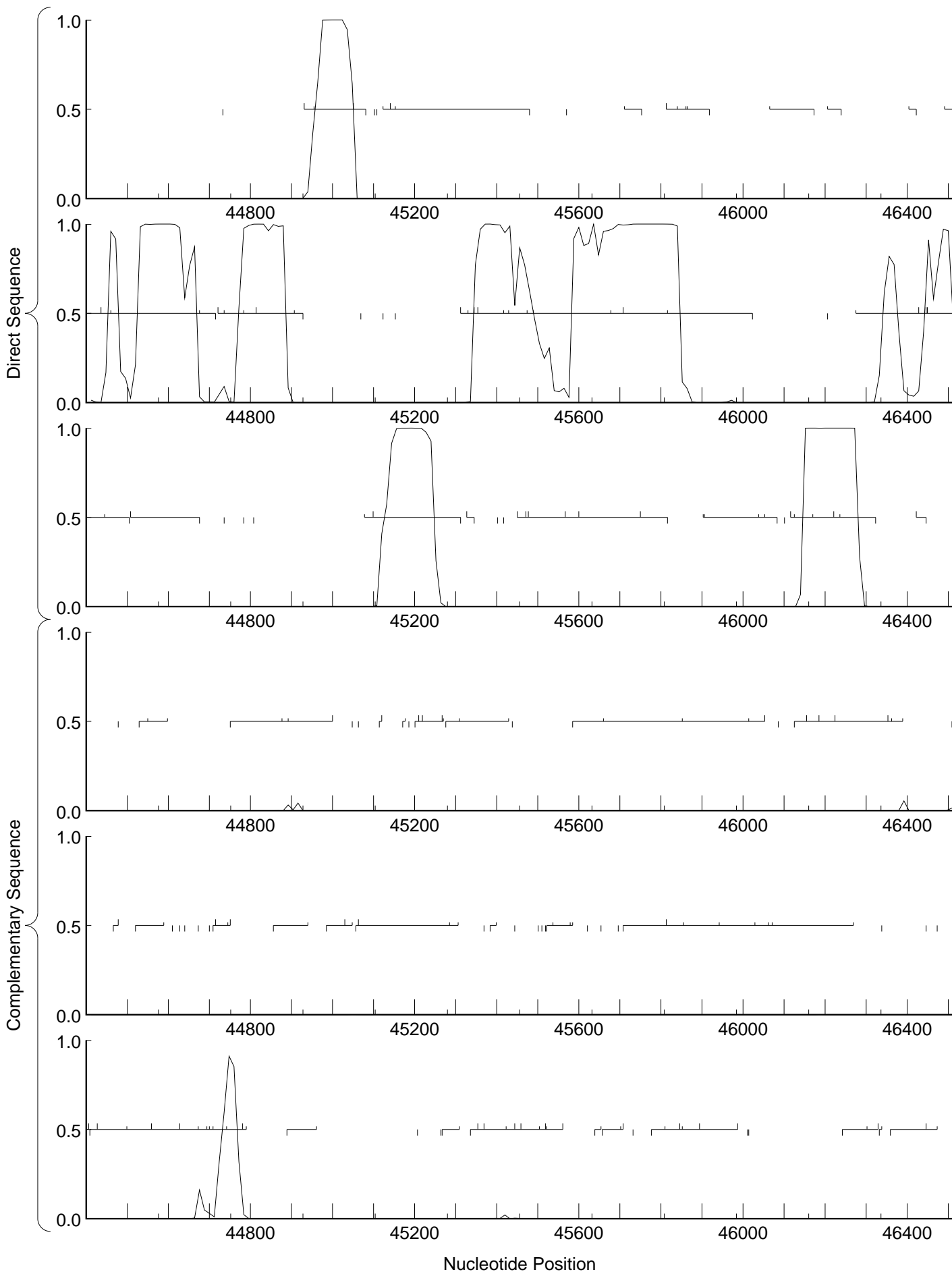


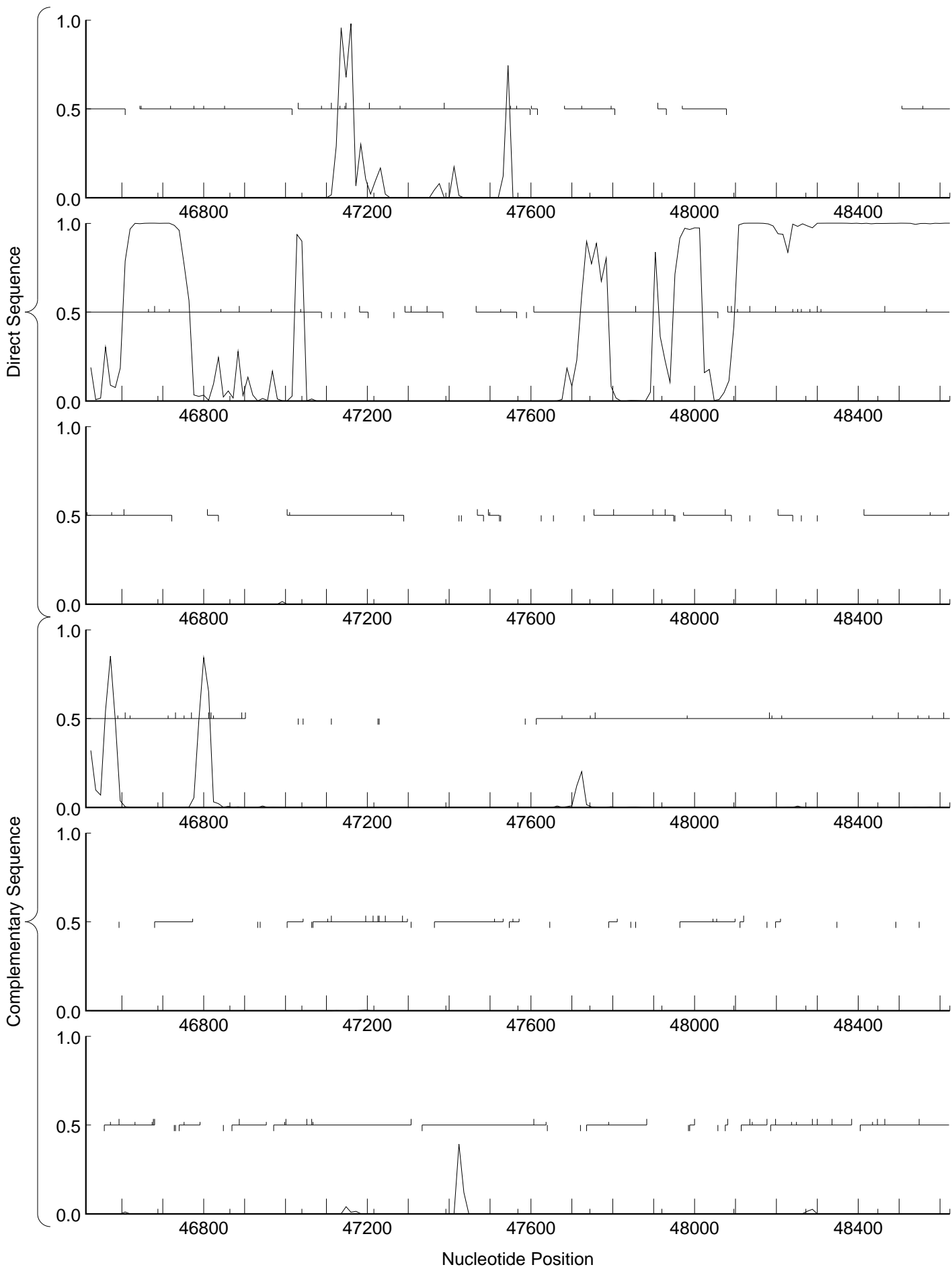
complete sequence, 93686 bp including 10-base 3' overhang (CGCGACGCTC), Cluster CQ, Order 4, Window 96, Step 12, 20/46

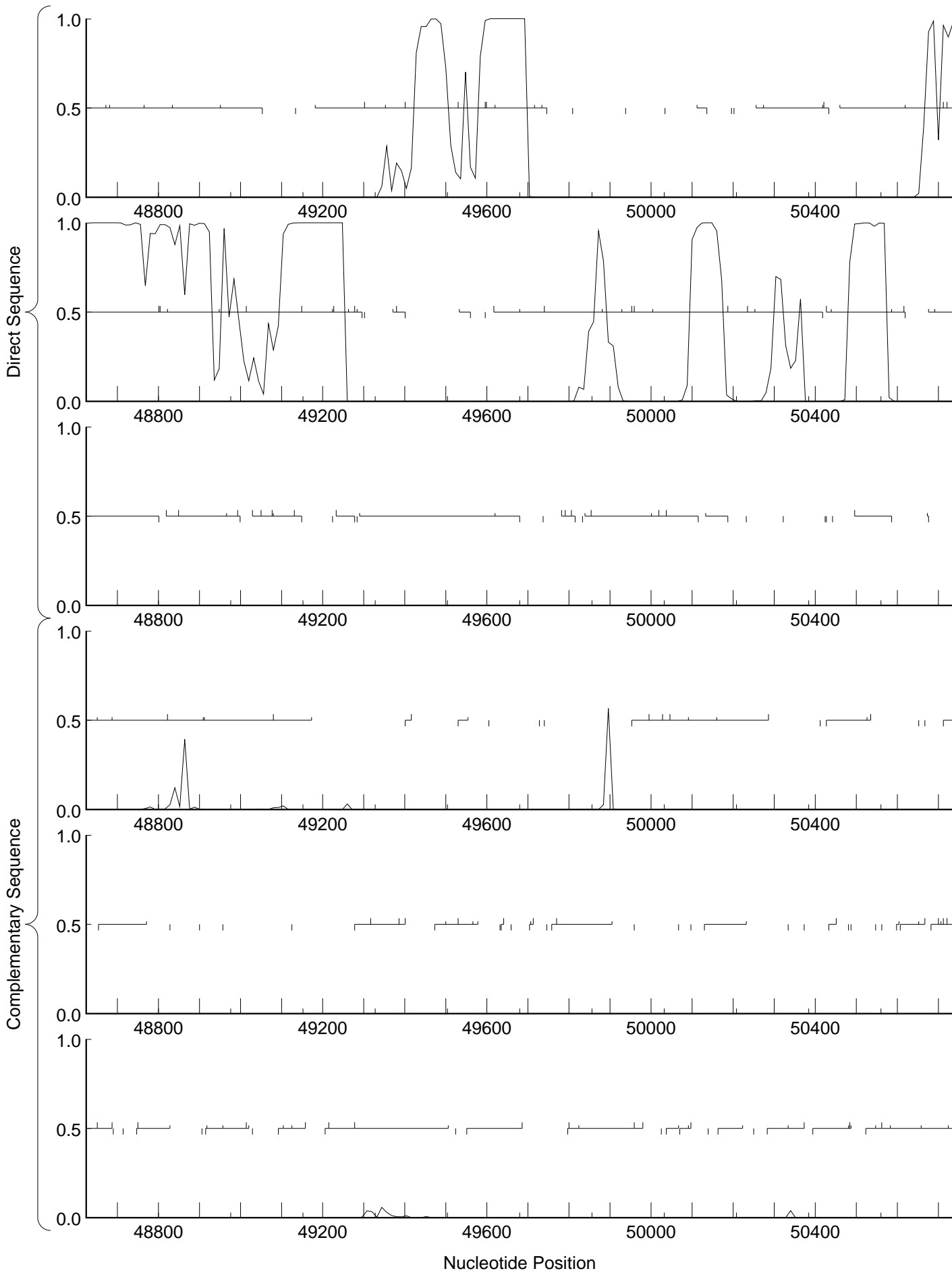


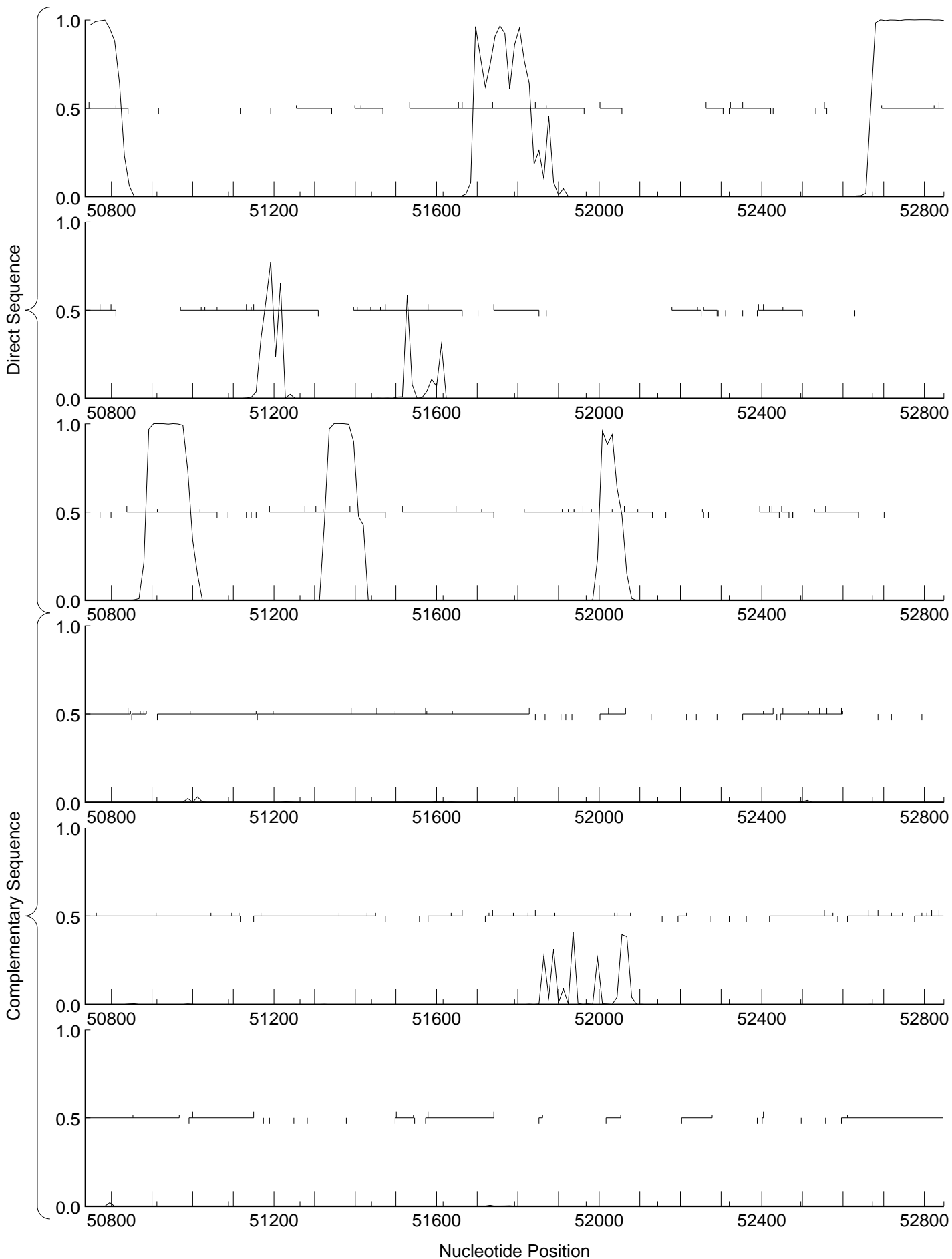


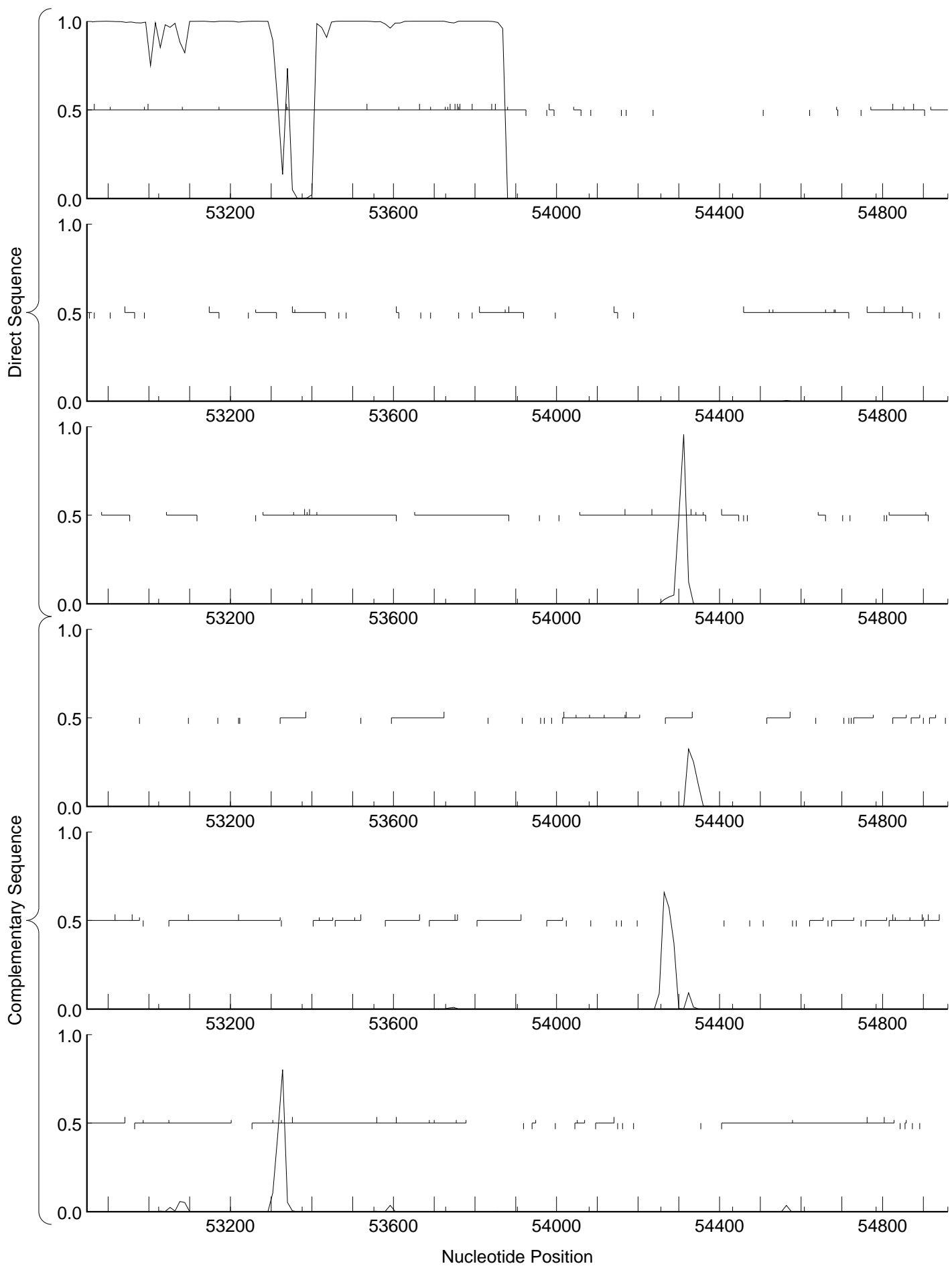


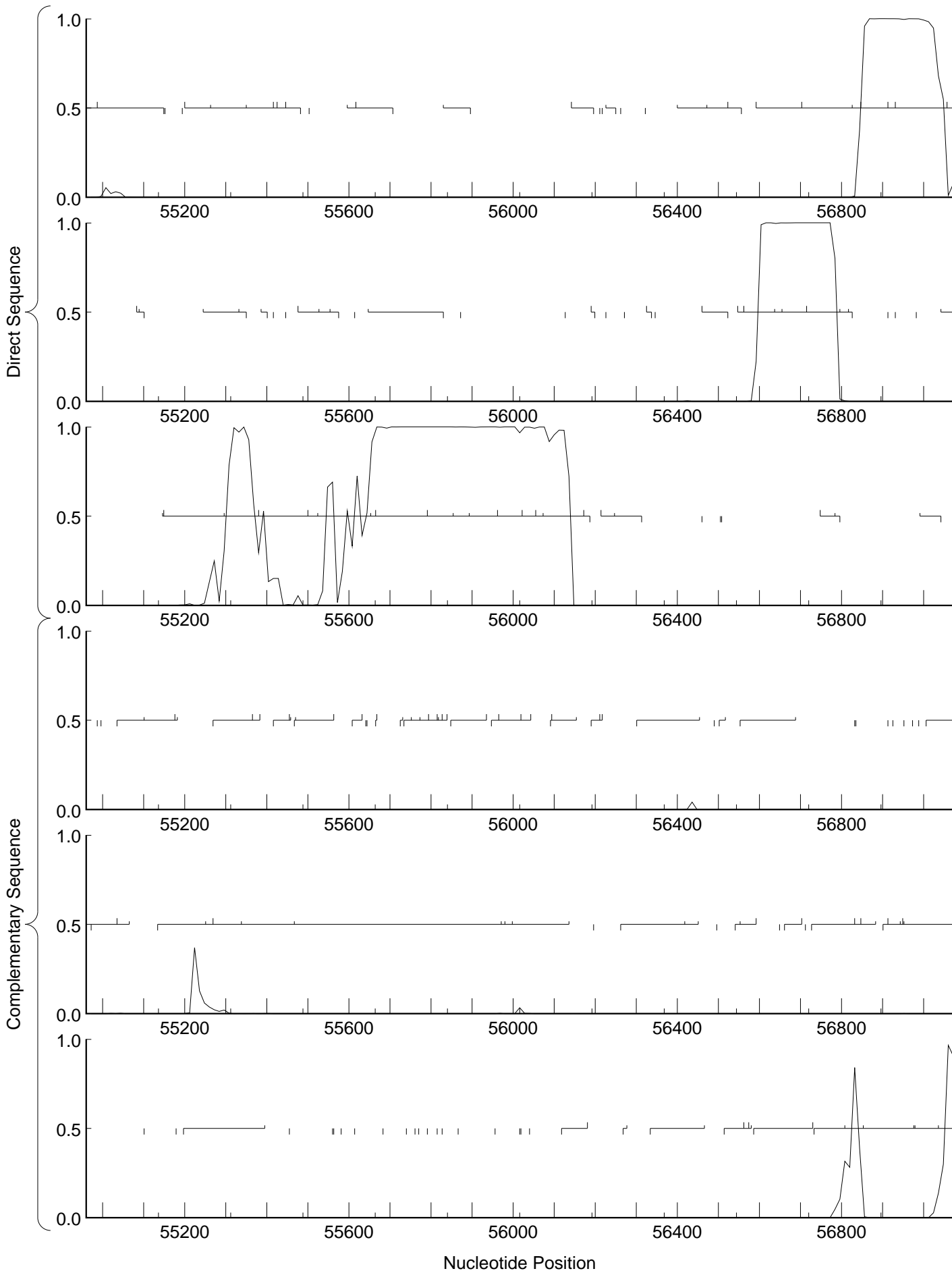


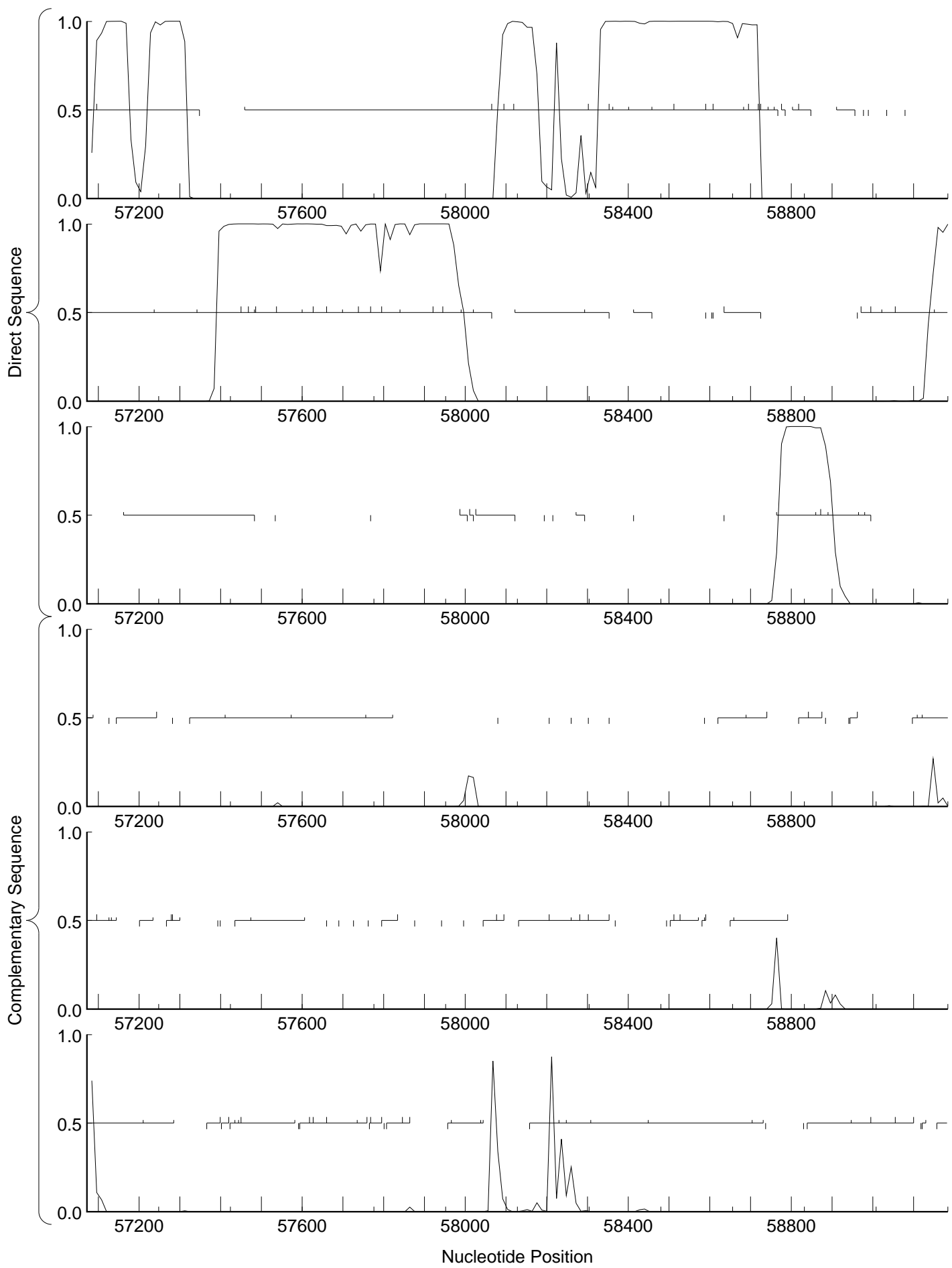


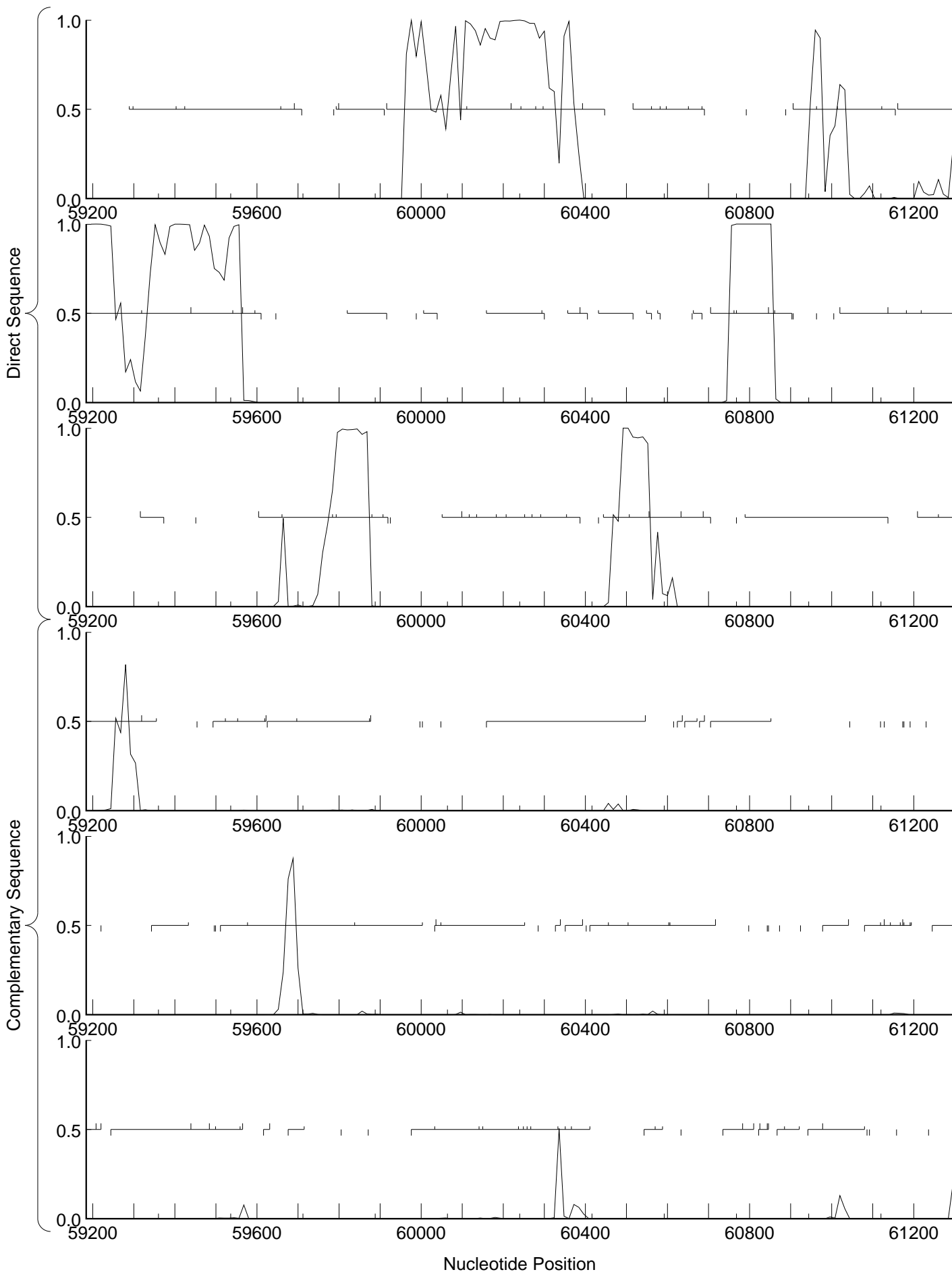


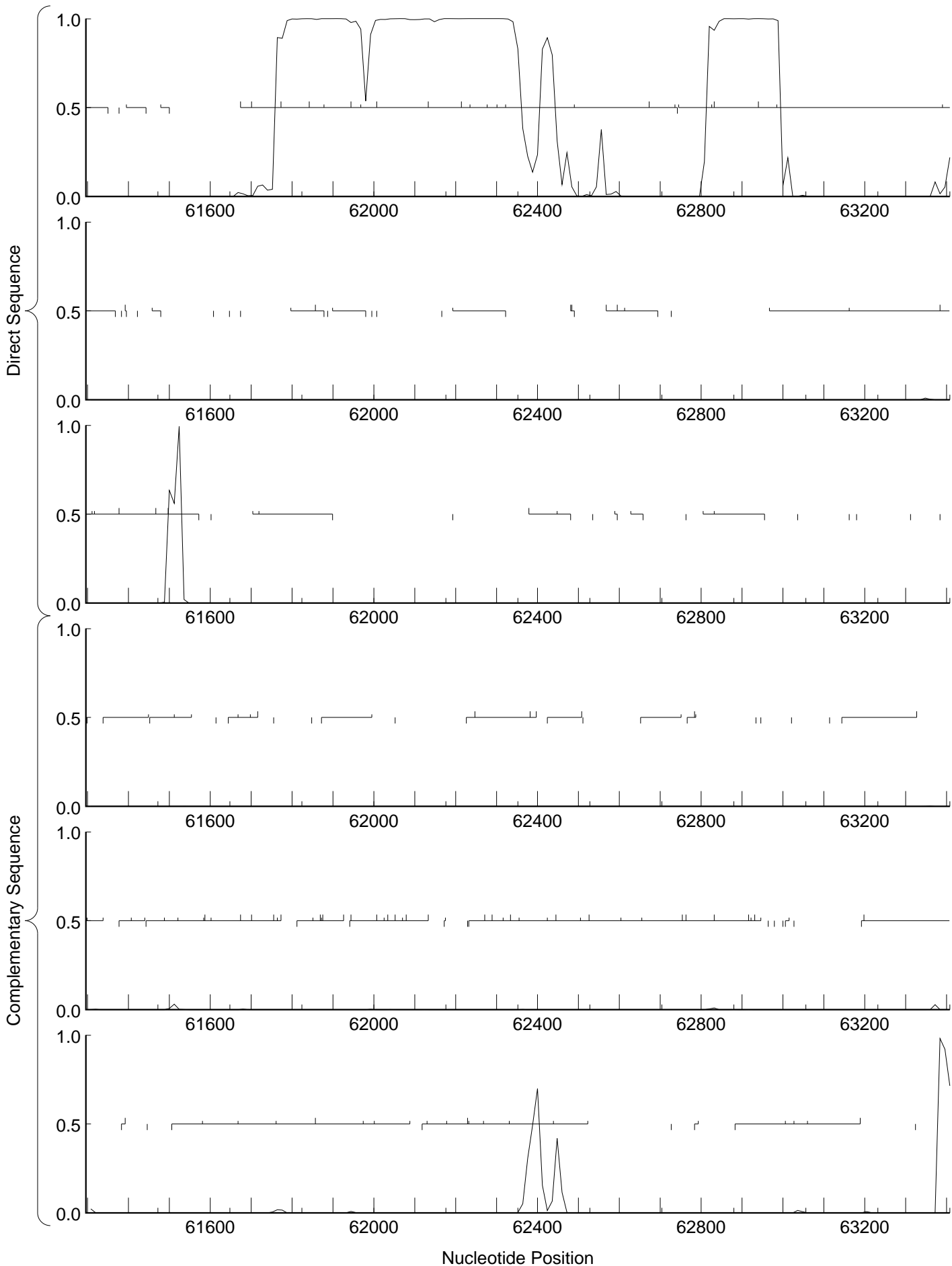


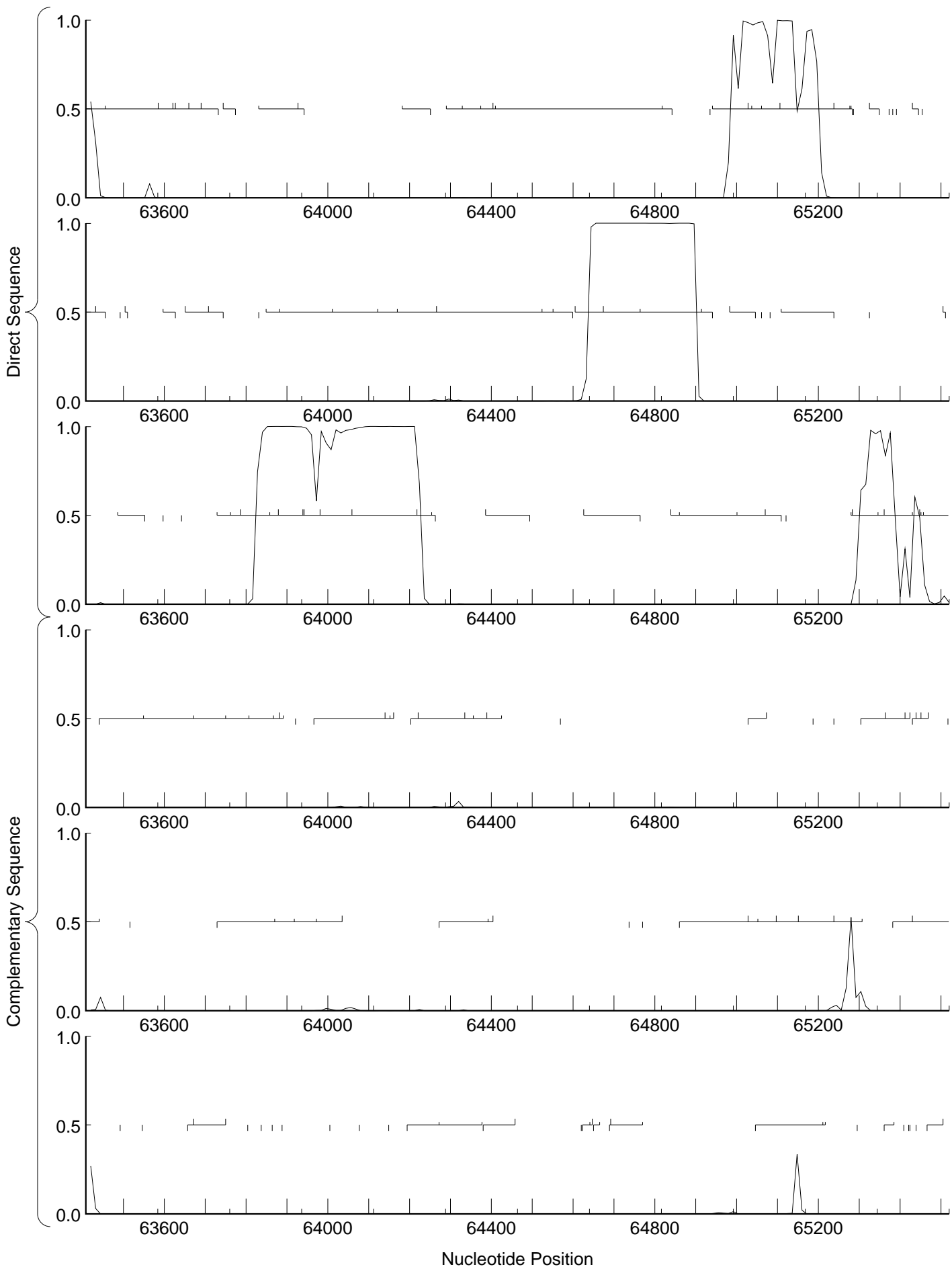


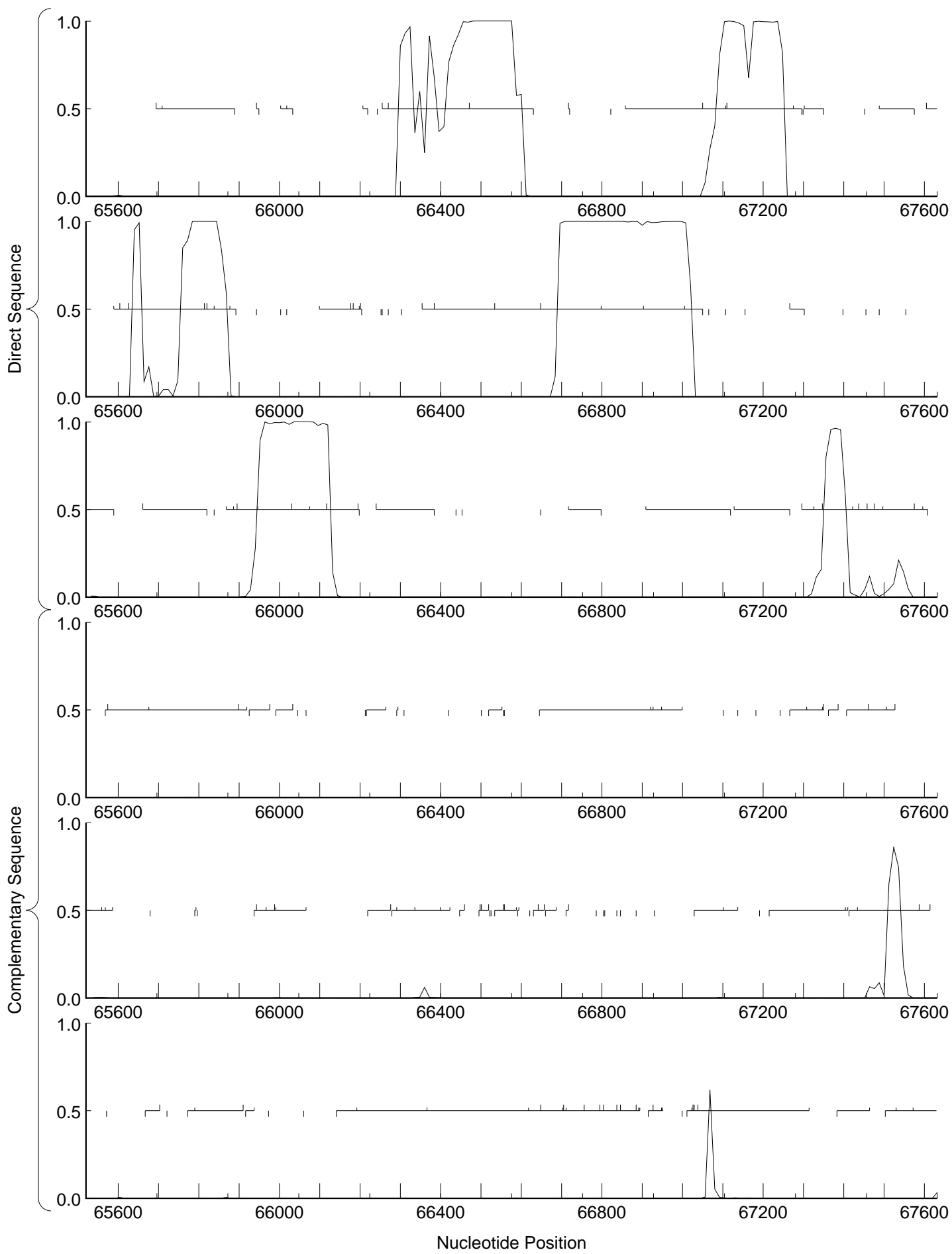


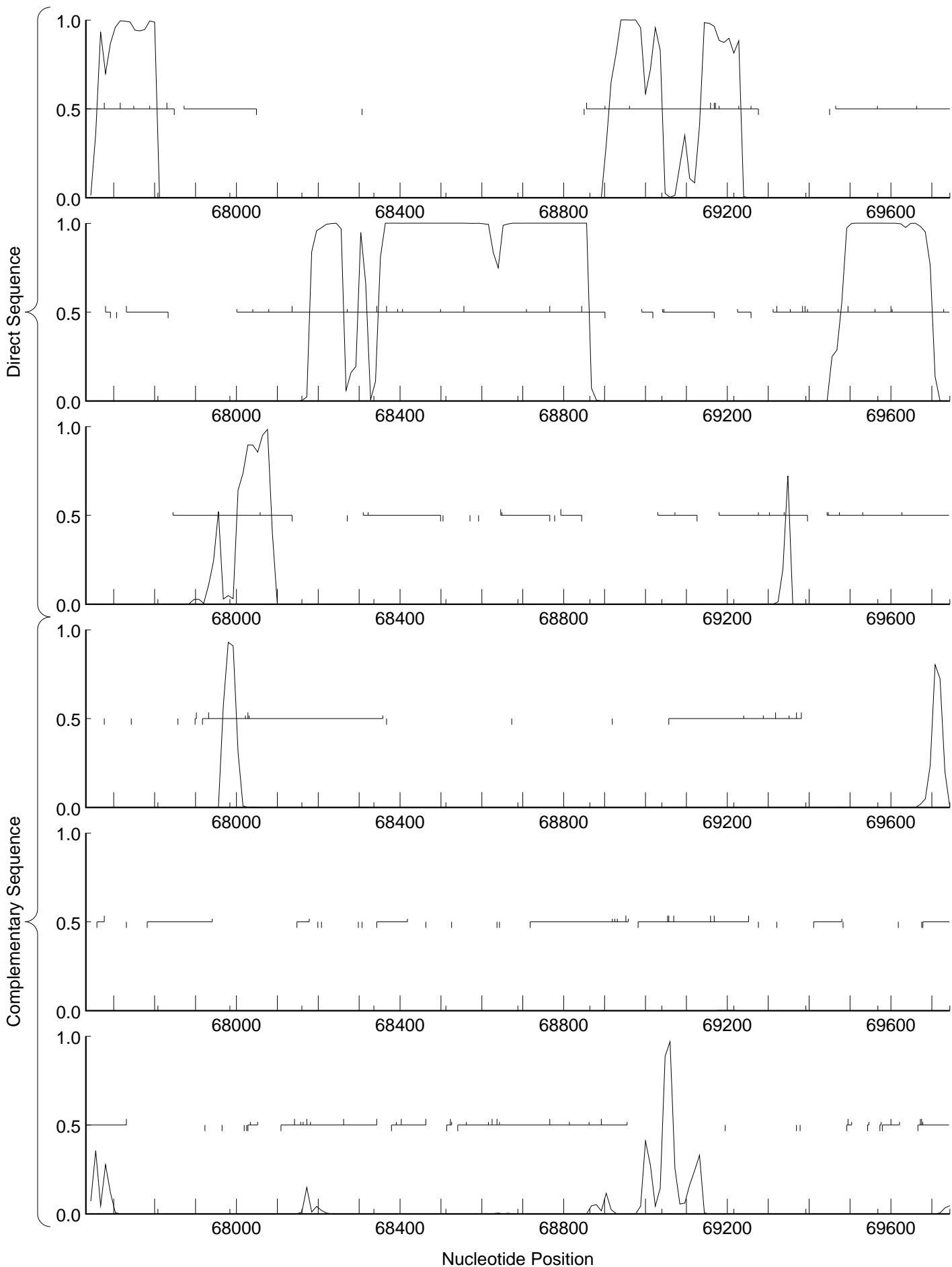












complete sequence, 93686 bp including 10-base 3' overhang (CGCGACGCTC), Cluster CQ, Order 4, Window 96, Step 12, 35/46

