

GeneMark

Version 2.5p (09.08.06)

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PROGRAM INFORMATION

Sequence : Mycobacterium phage Chuckly complete sequence, 55189 bp including 10-base 3' overhang (CGGT
Analysis Date : 3/2/21 at 11:49:51
Pages : 28
Sequence Length : 55189 bp
GC Content : 61.56%

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,

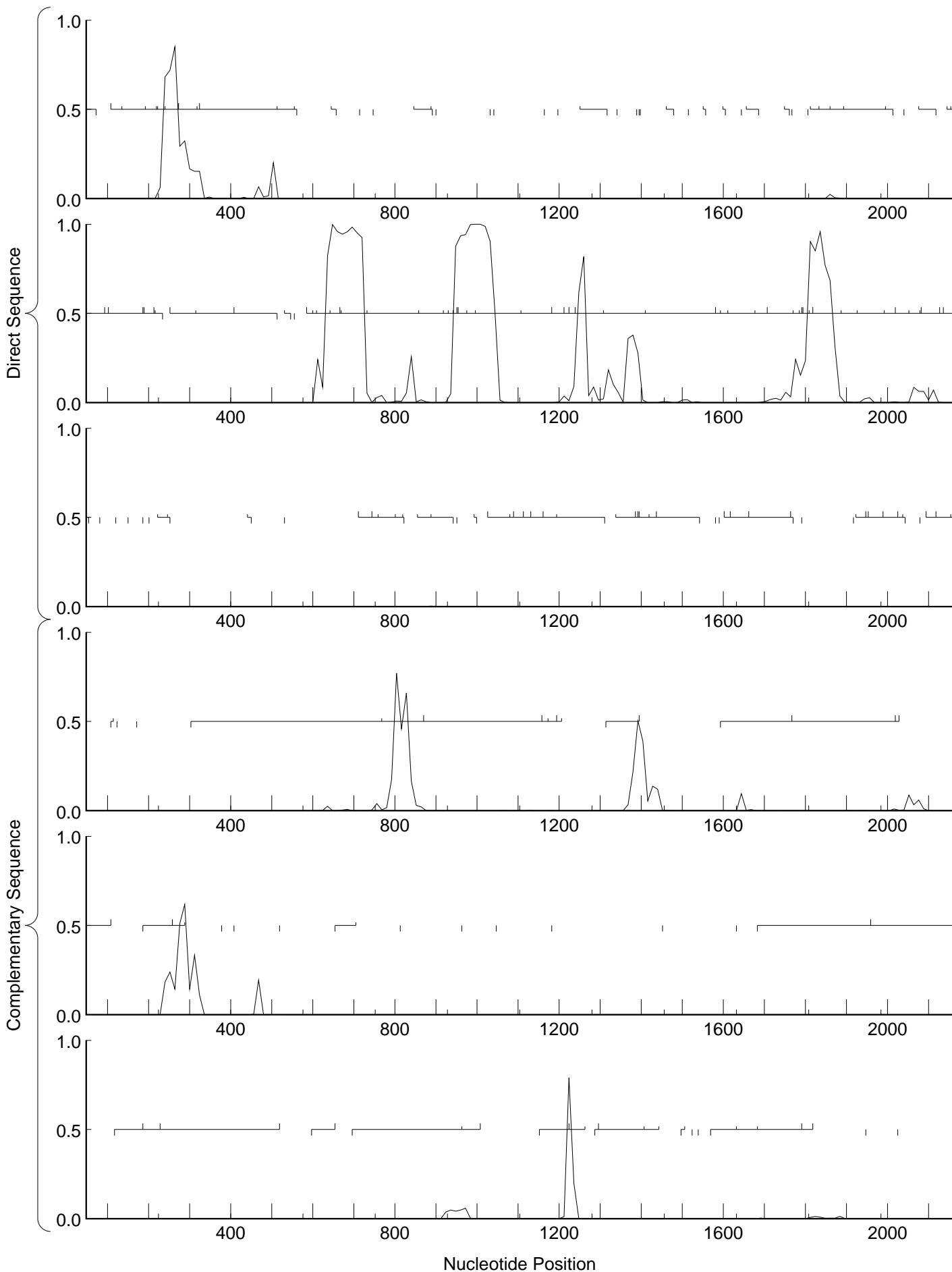
Matrix notes & comments

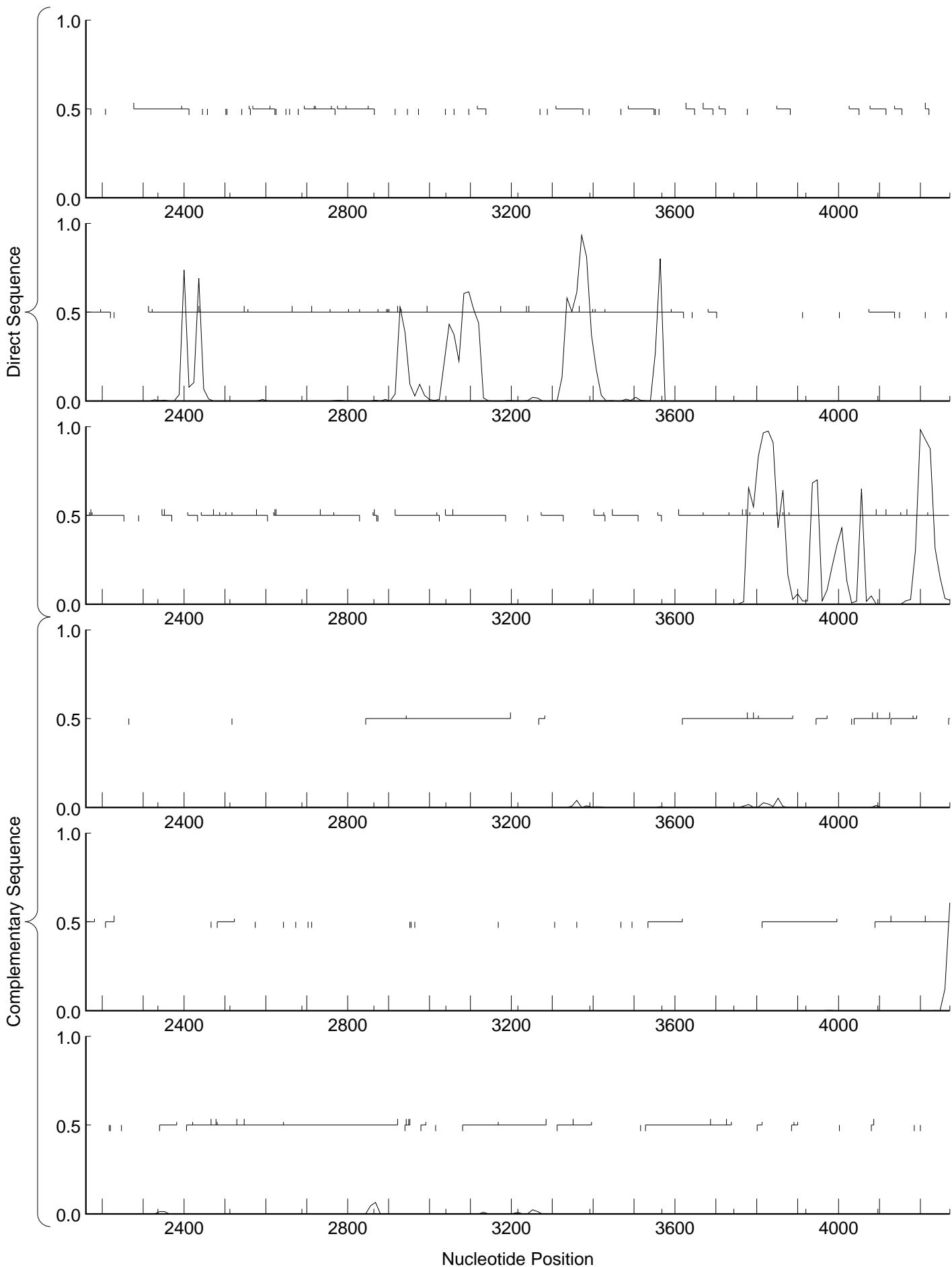
Training set derived by GeneMarkS, 4.27 September 2014
Tue Sep 23 15:23:03 2014

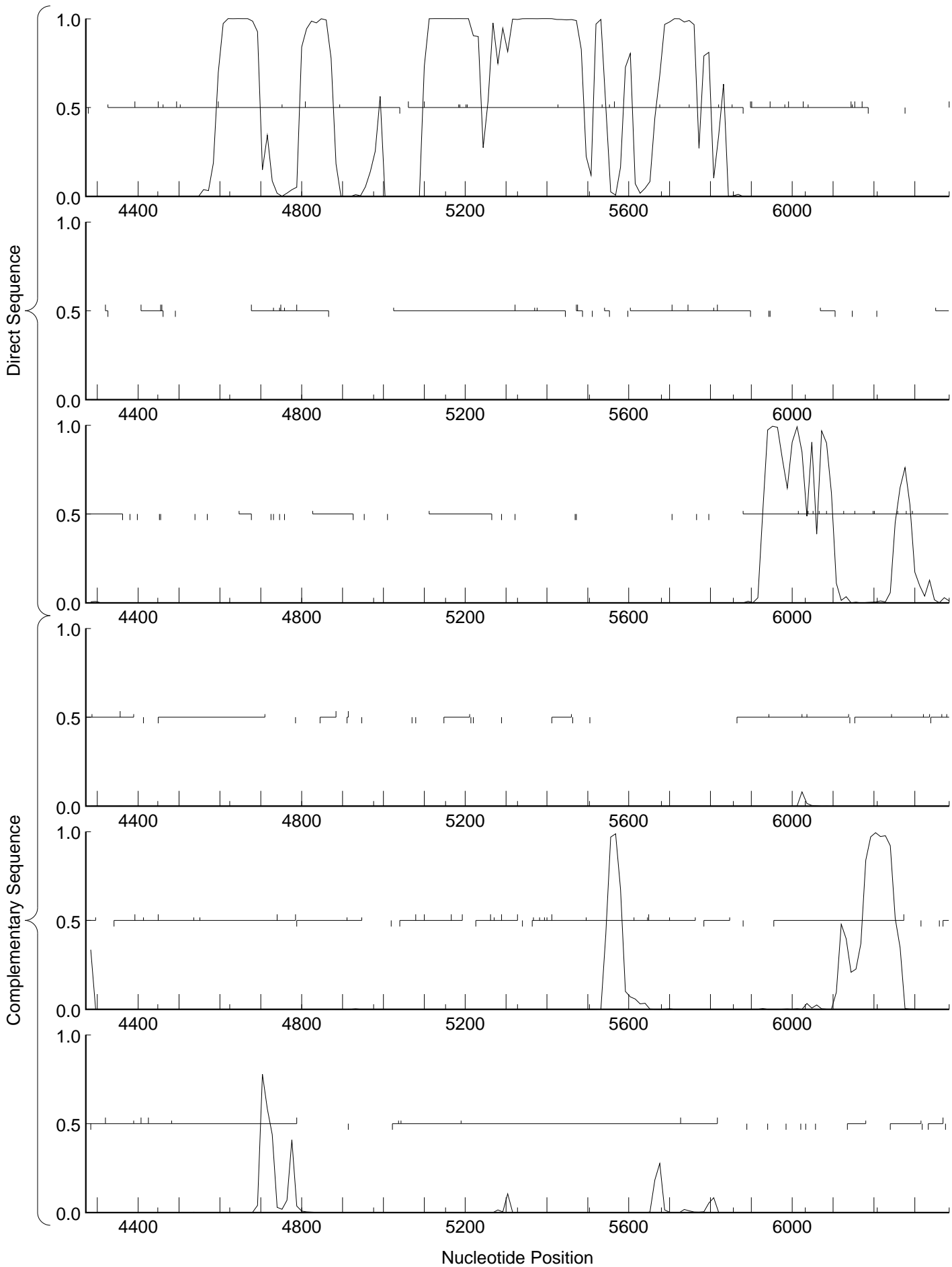
MATRIX INFORMATION

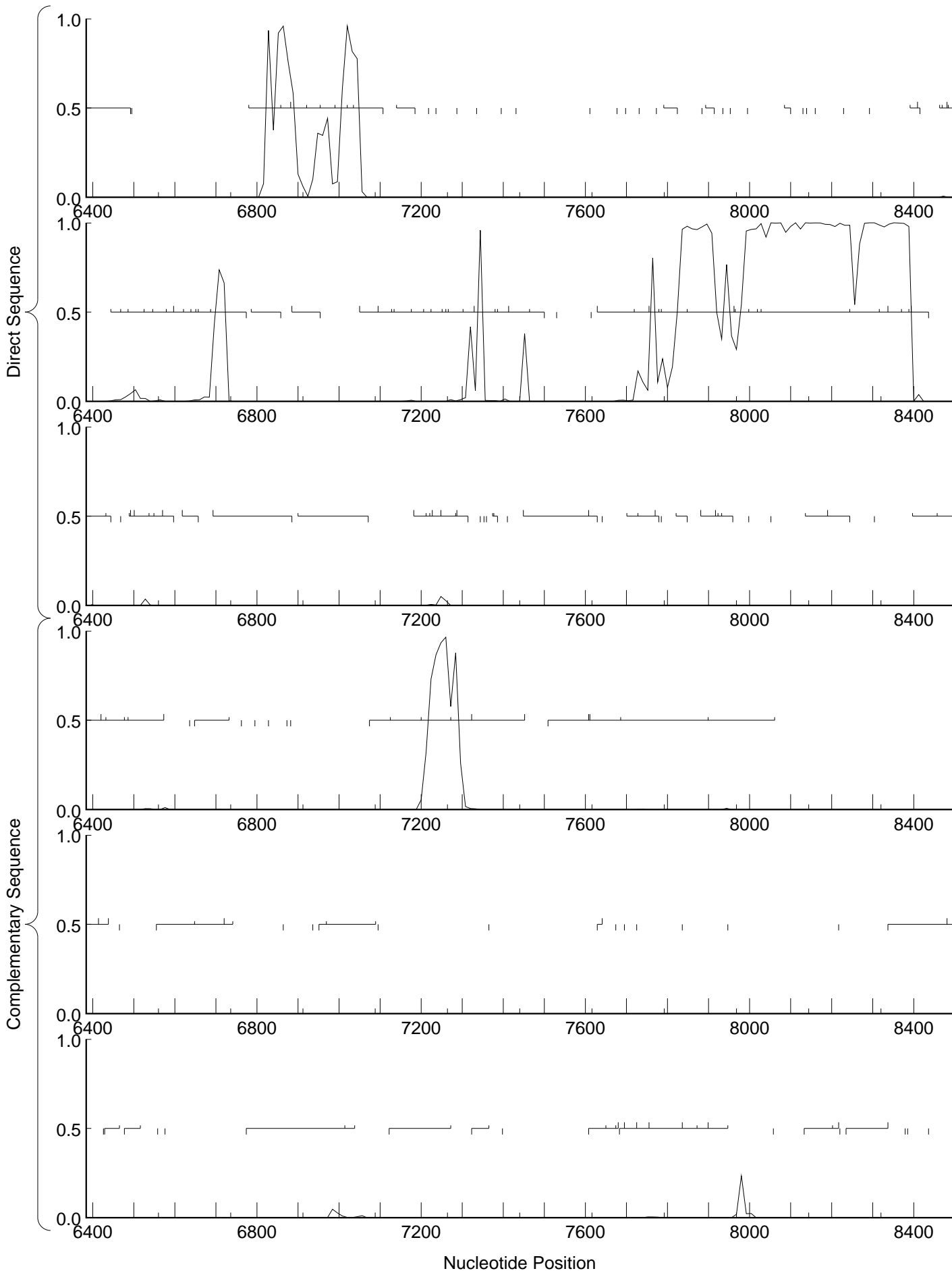
Matrix : Mycobacterium_smegmatis_MC2_155
Author : -
Order : 4

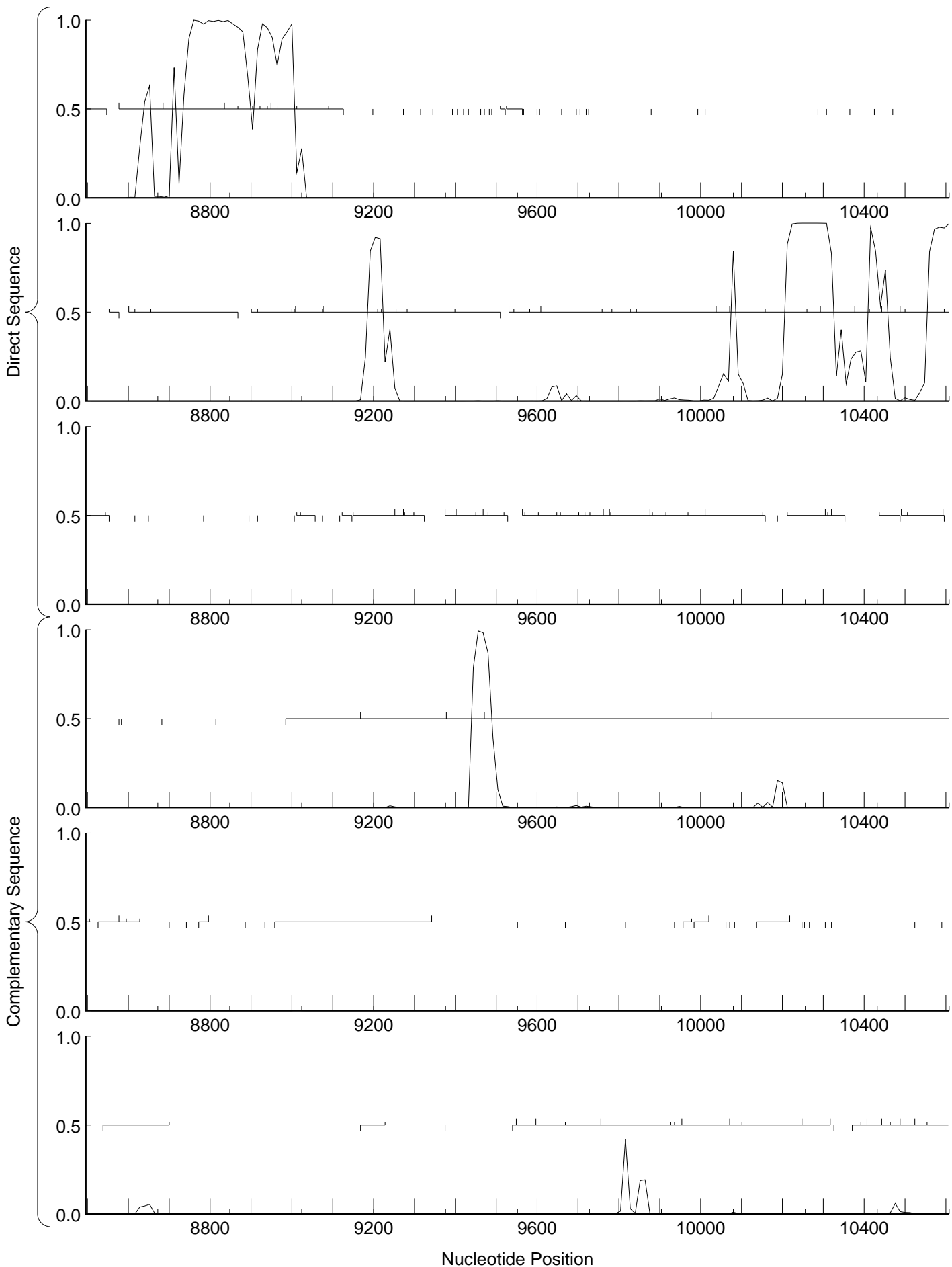
Send questions / comments to:
Dr. M. Borodovsky
Georgia Institute of Technology
School of Biology
Atlanta, GA 30332-0230

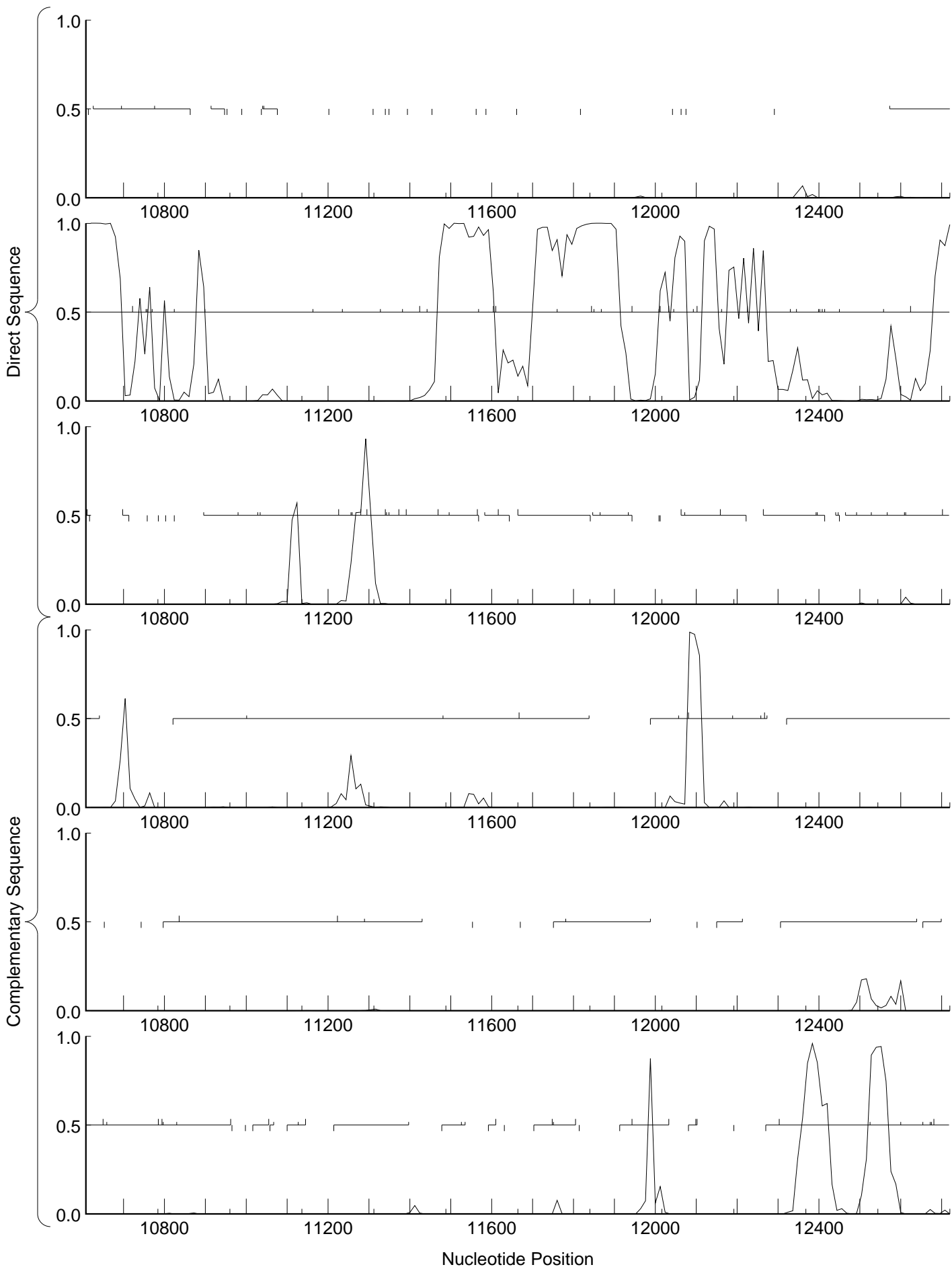


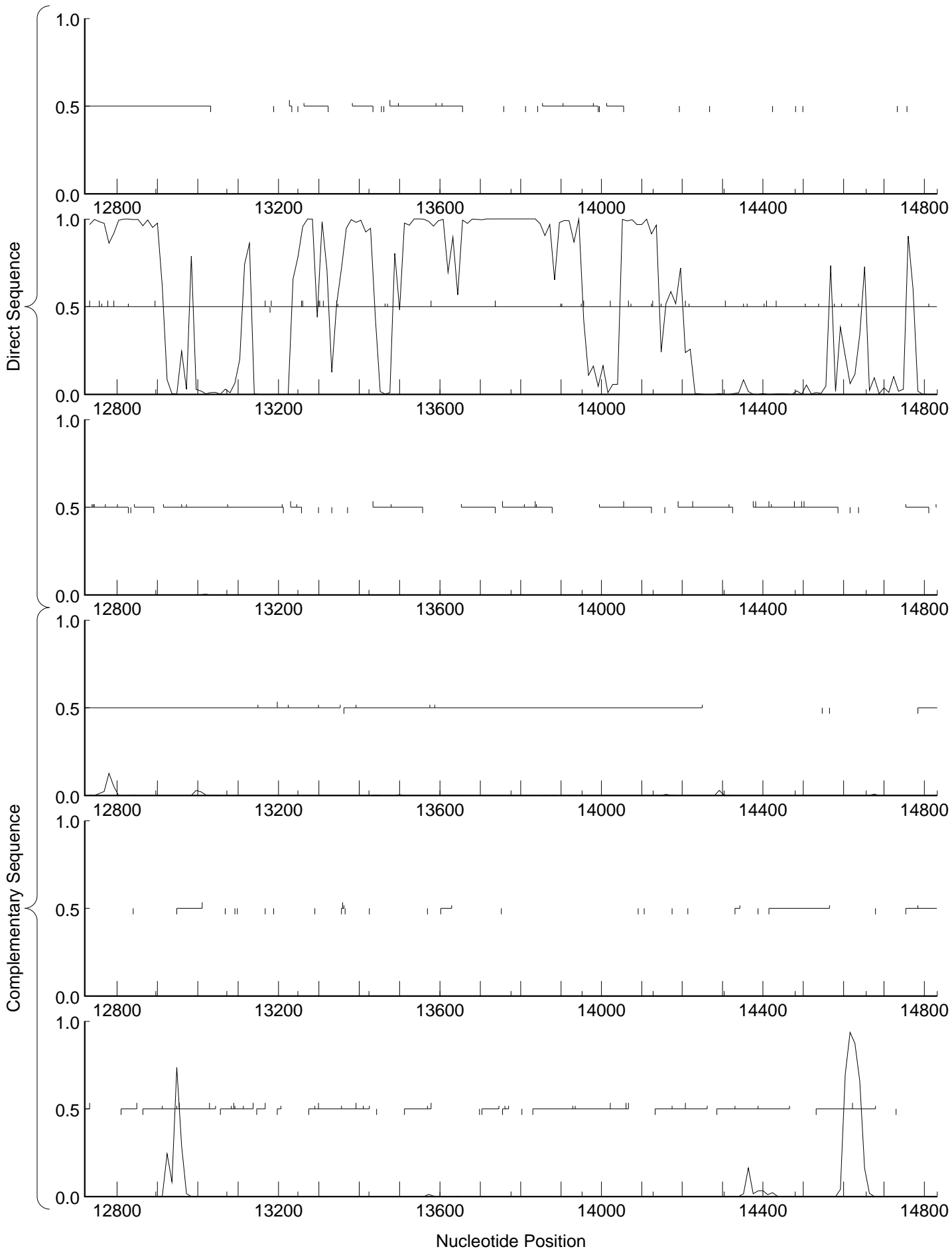


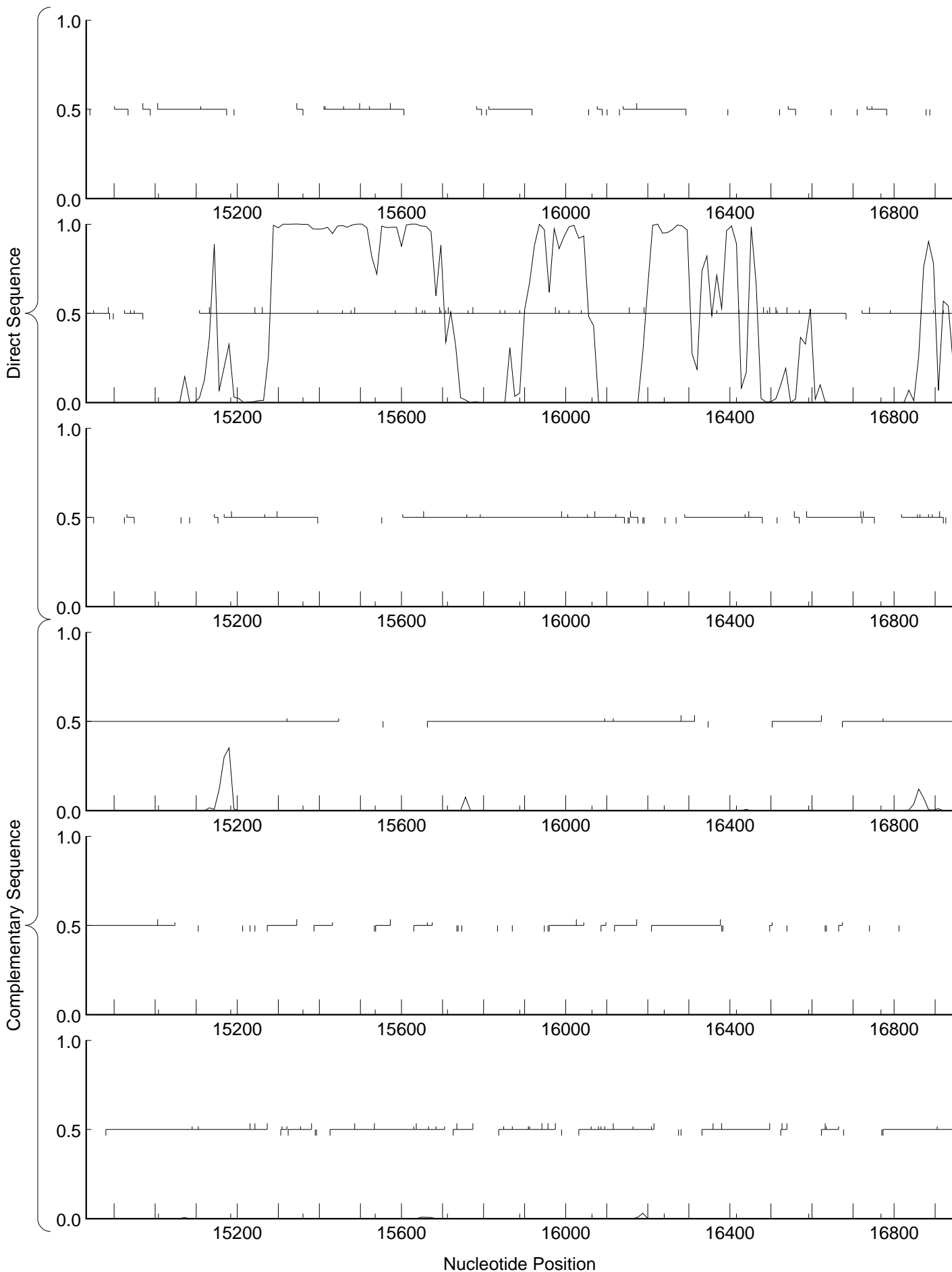




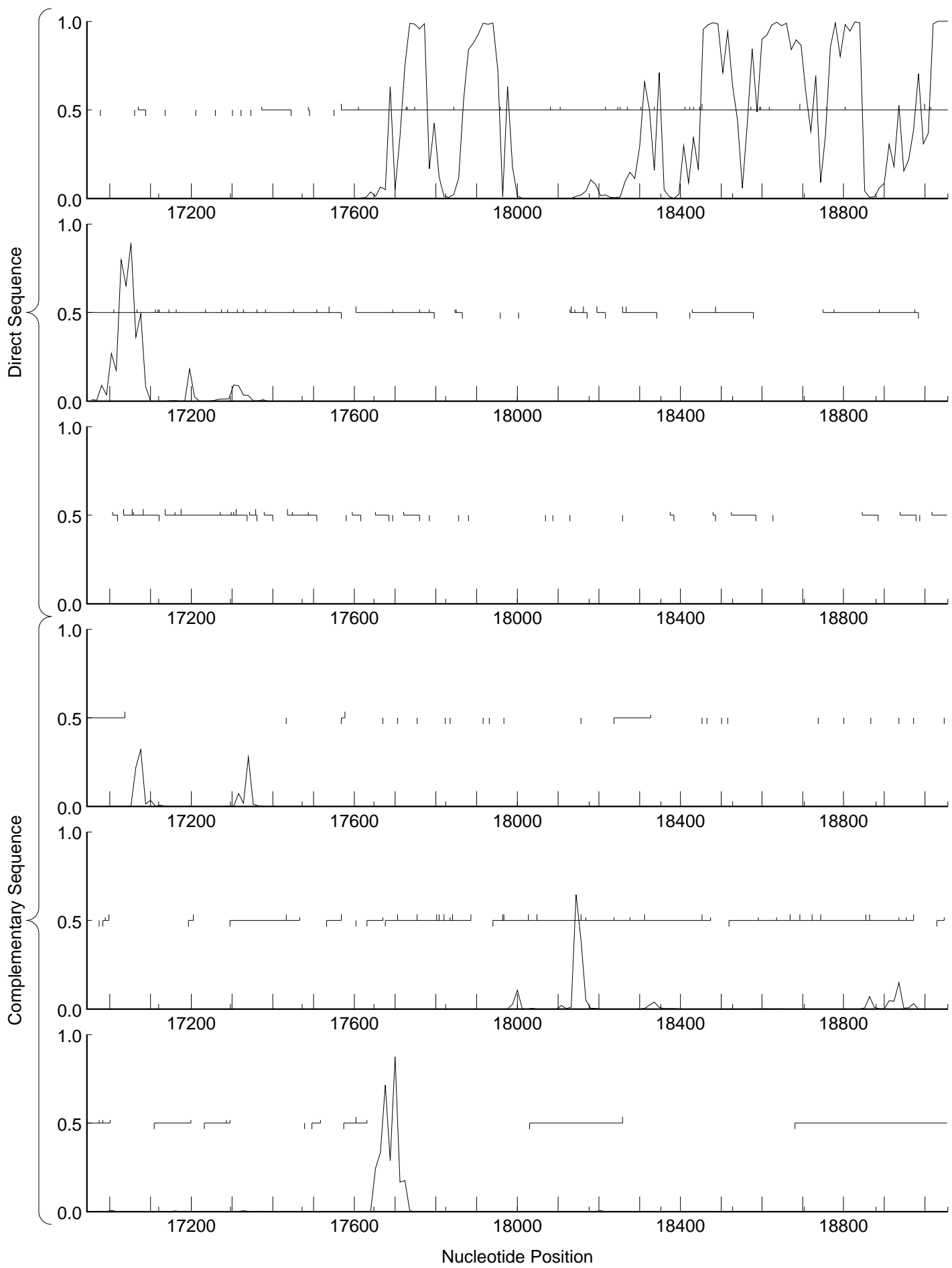




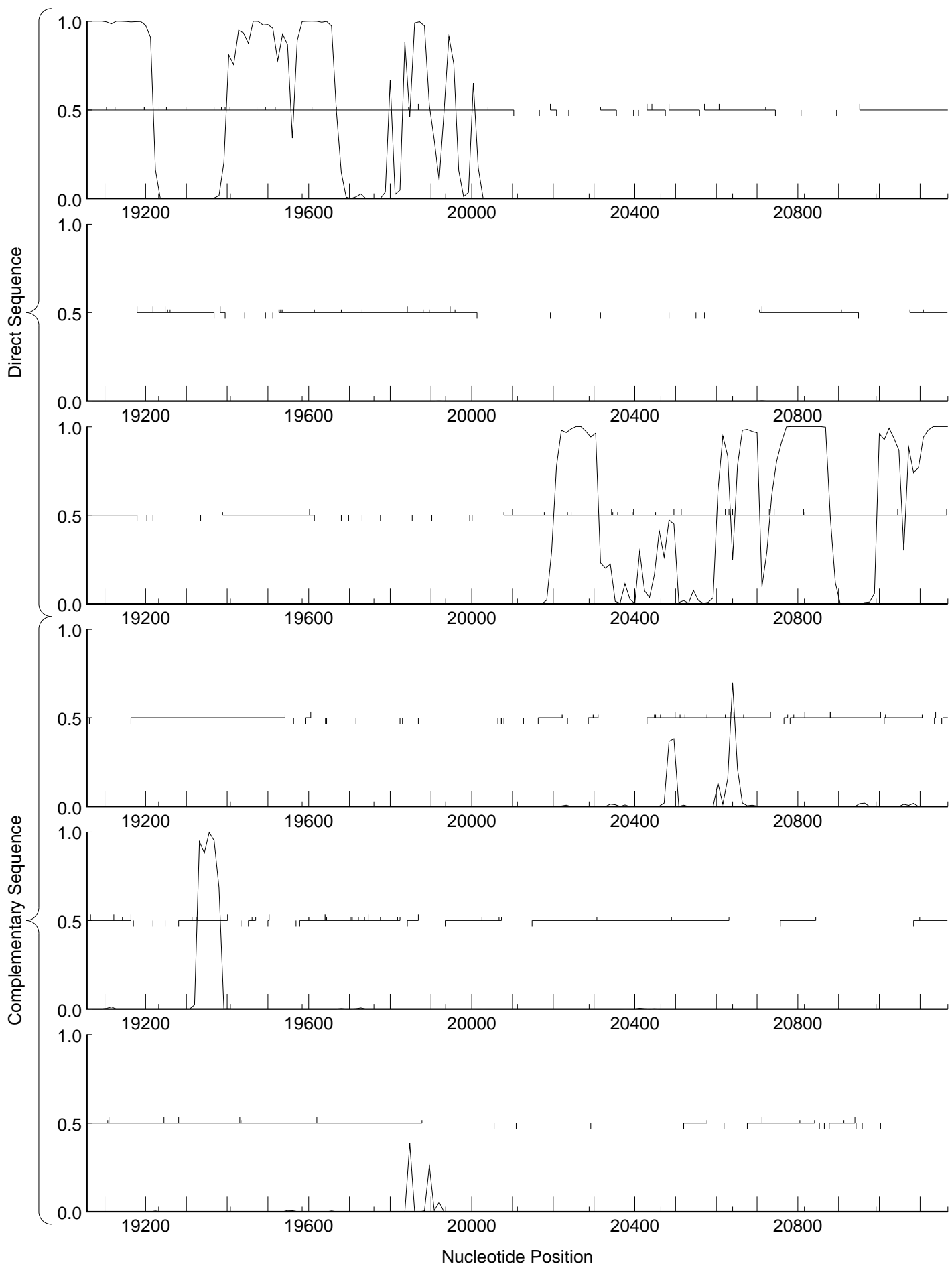


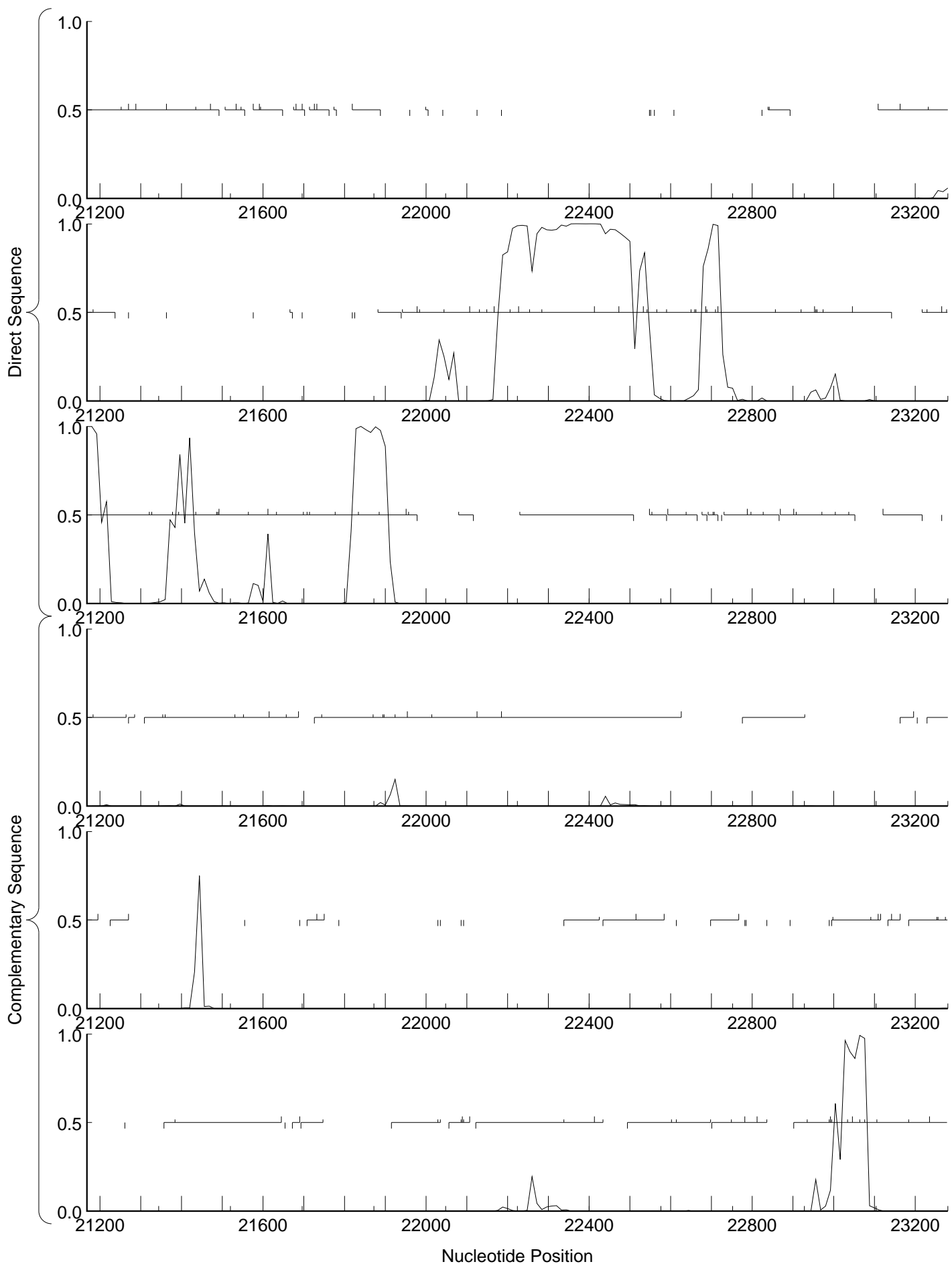


complete sequence, 55189 bp including 10-base 3' overhang (CGGTAGGCGC), Cluster F1, Order 4, Window 96, Step 12, 10/28

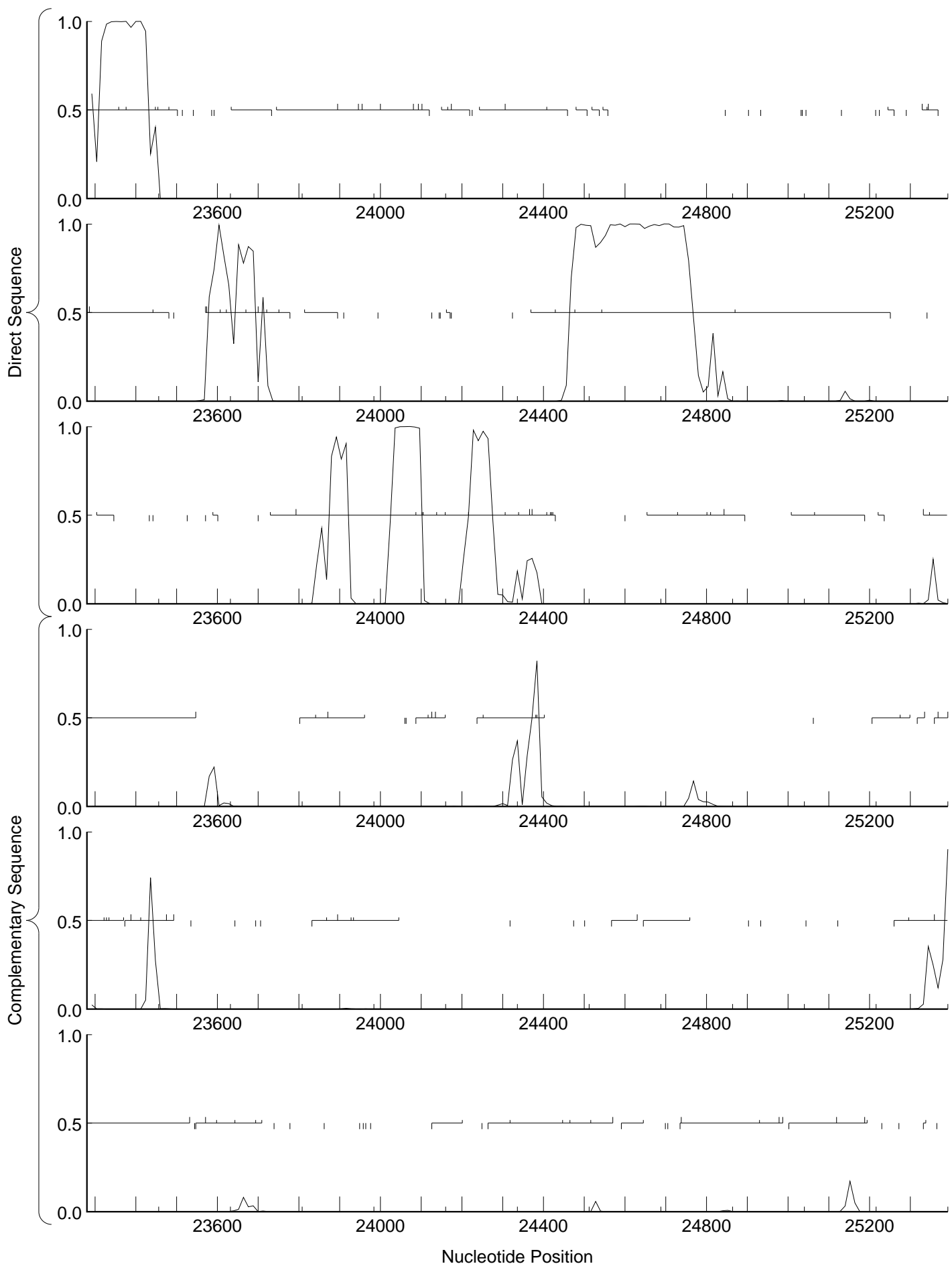


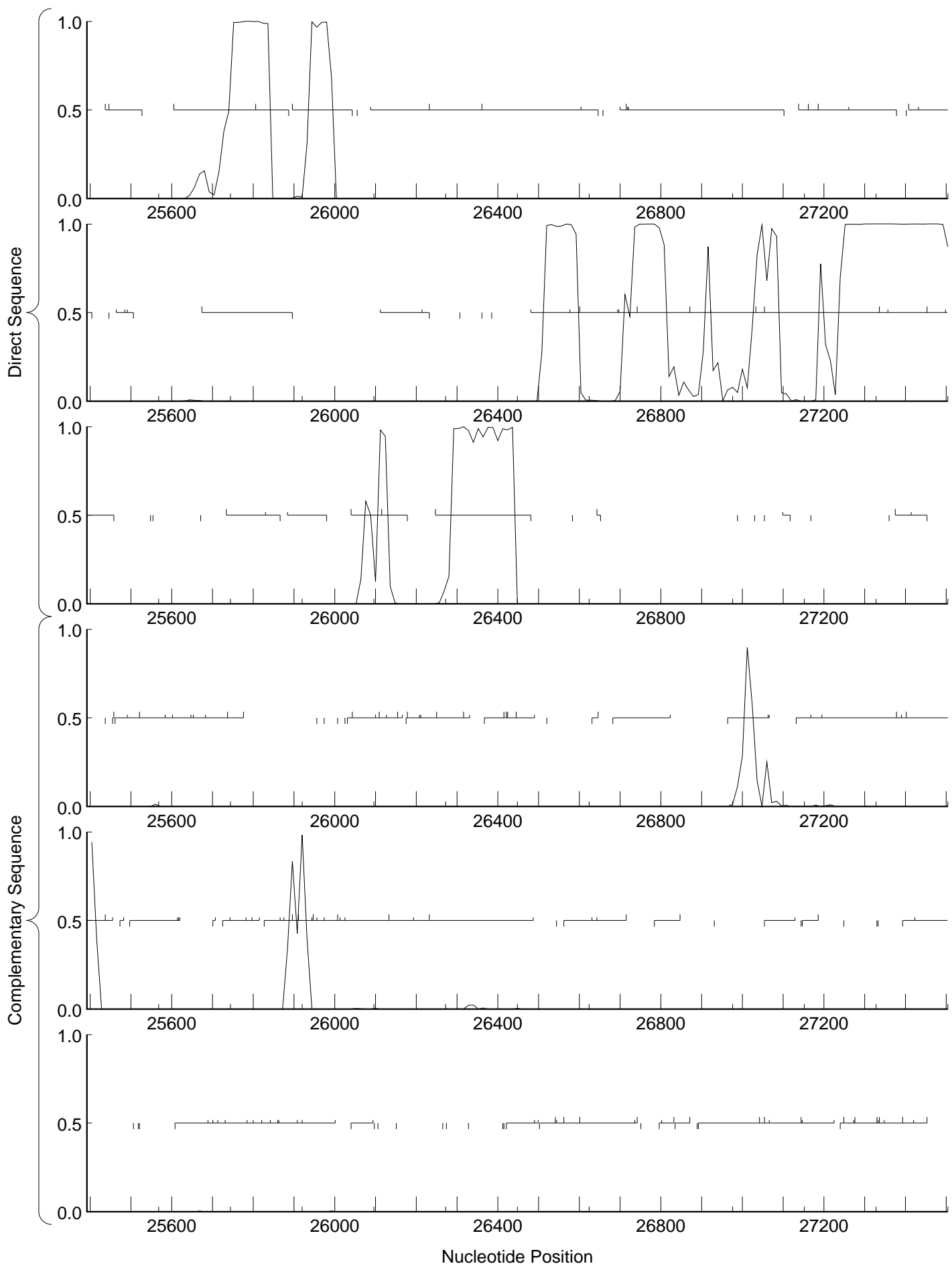
complete sequence, 55189 bp including 10-base 3' overhang (CGGTAGGCGC), Cluster F1, Order 4, Window 96, Step 12, 11/28



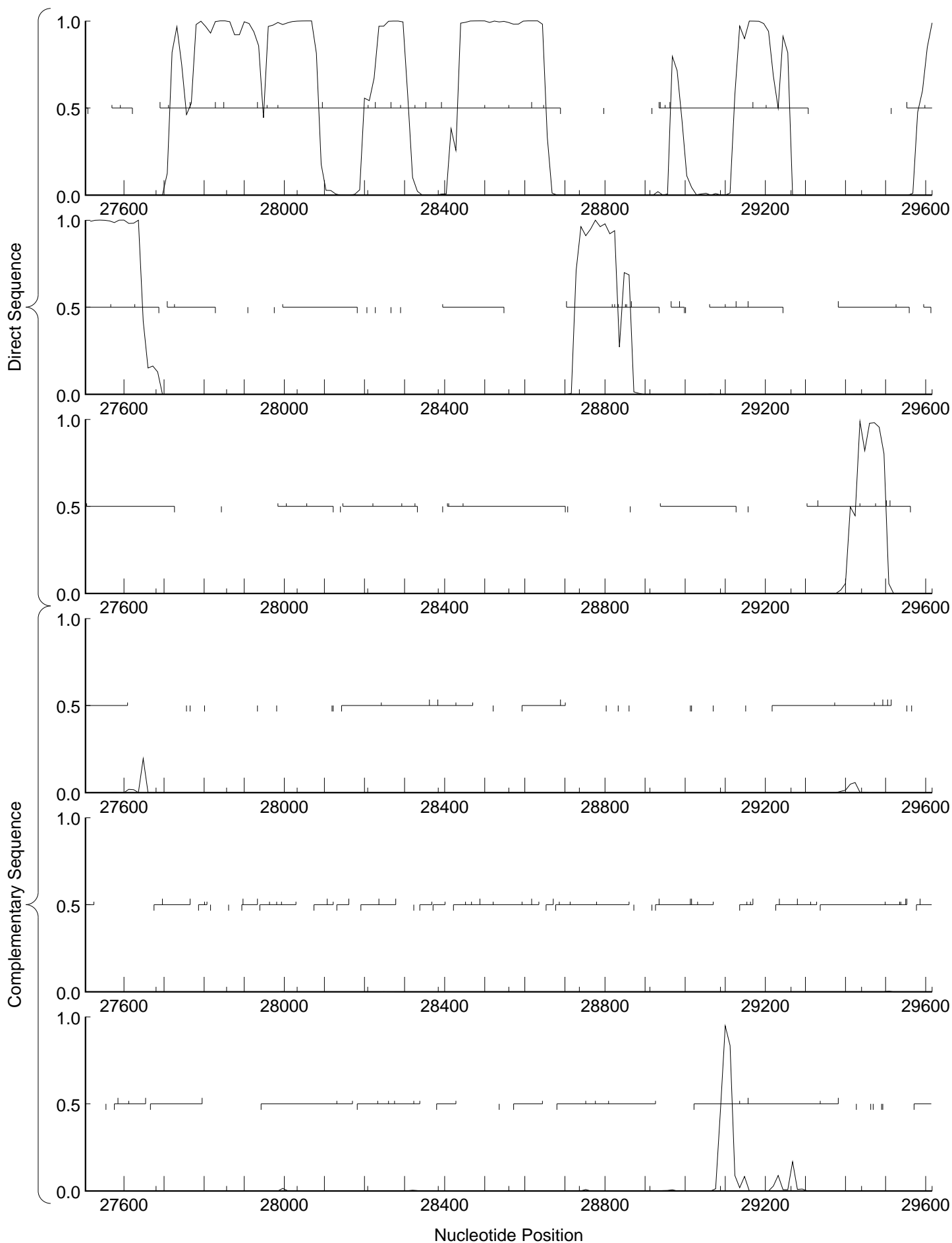


complete sequence, 55189 bp including 10-base 3' overhang (CGGTAGGCGC), Cluster F1, Order 4, Window 96, Step 12, 13/28

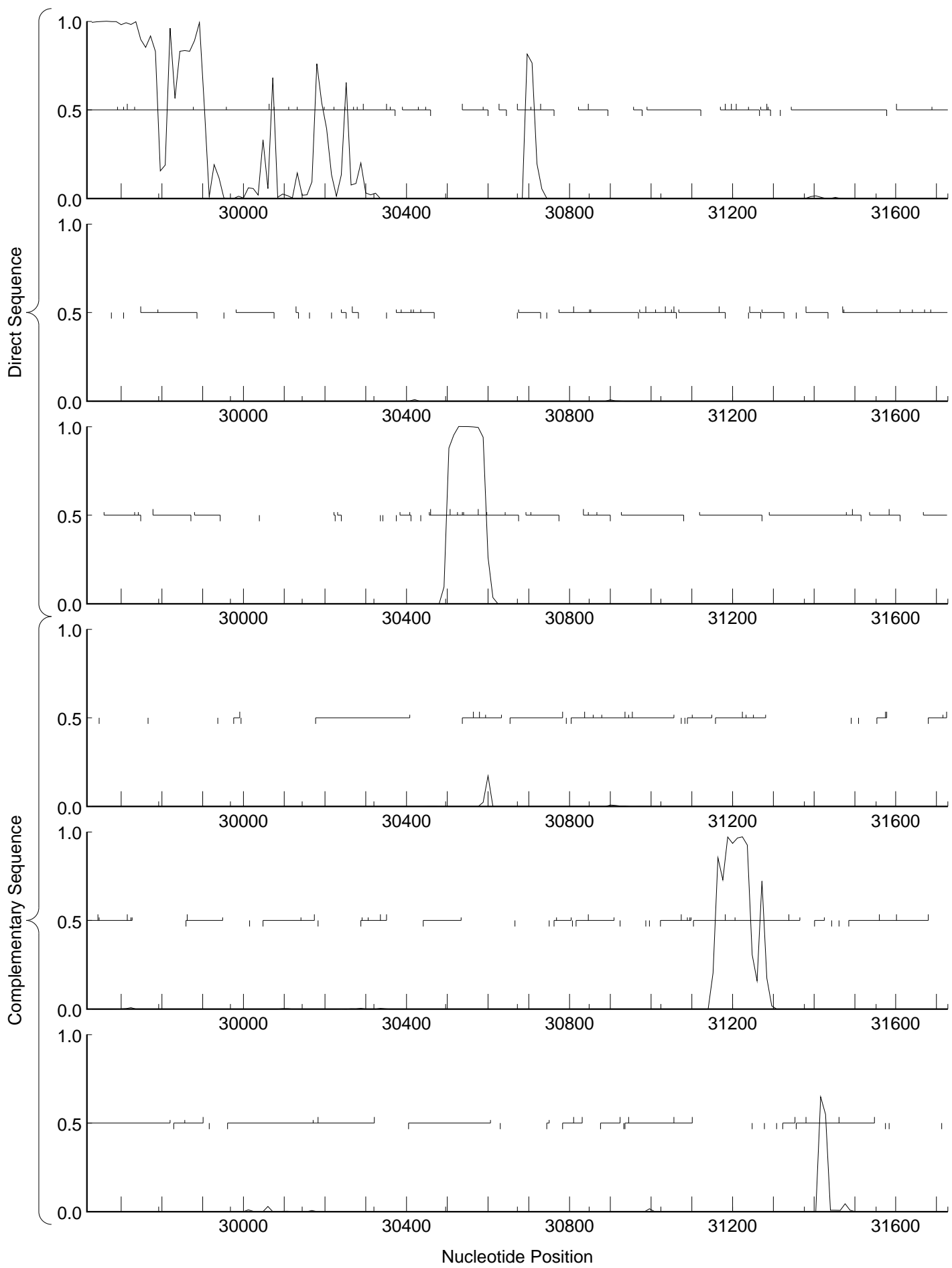




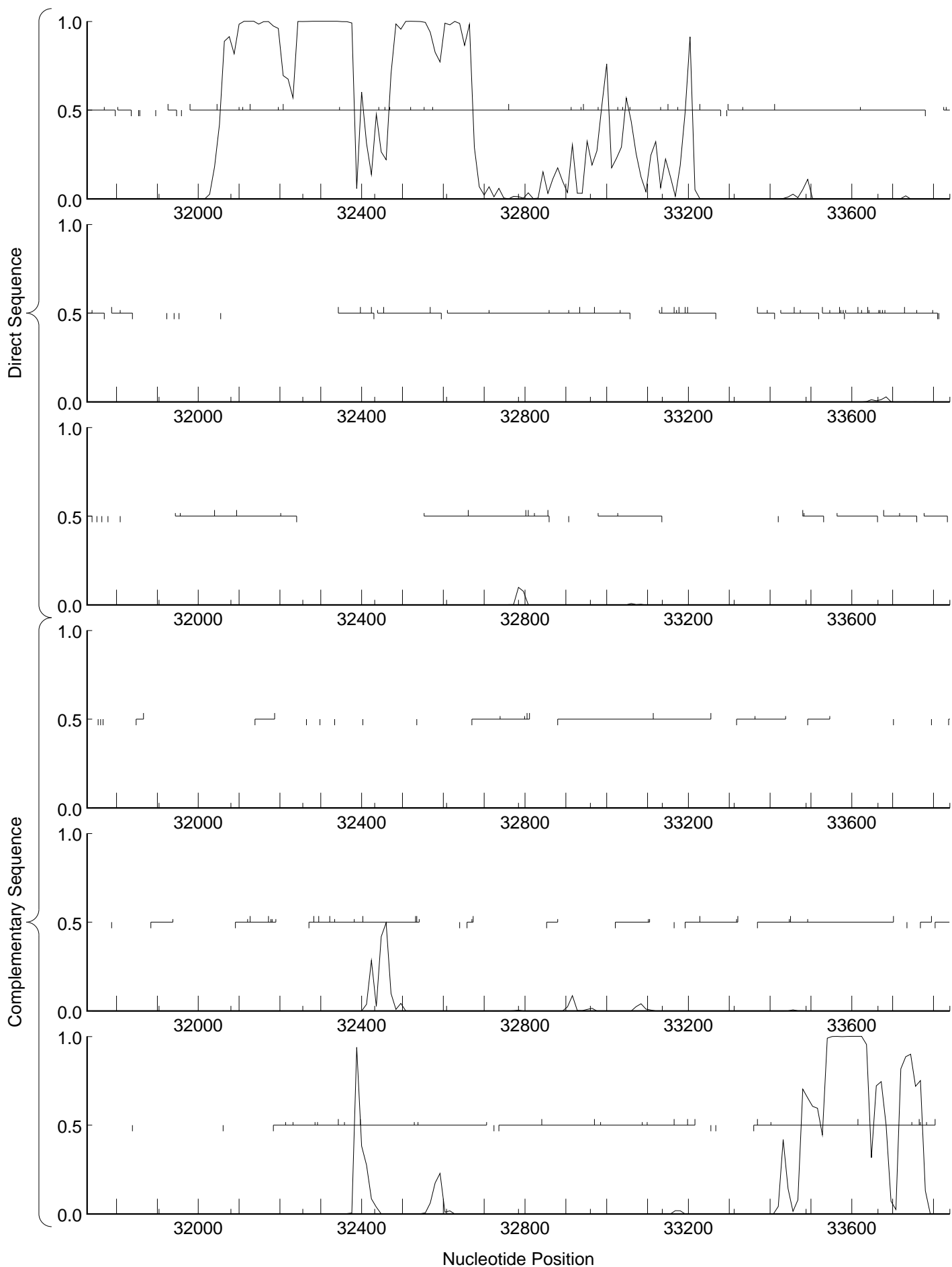
complete sequence, 55189 bp including 10-base 3' overhang (CGGTAGGCGC), Cluster F1, Order 4, Window 96, Step 12, 15/28

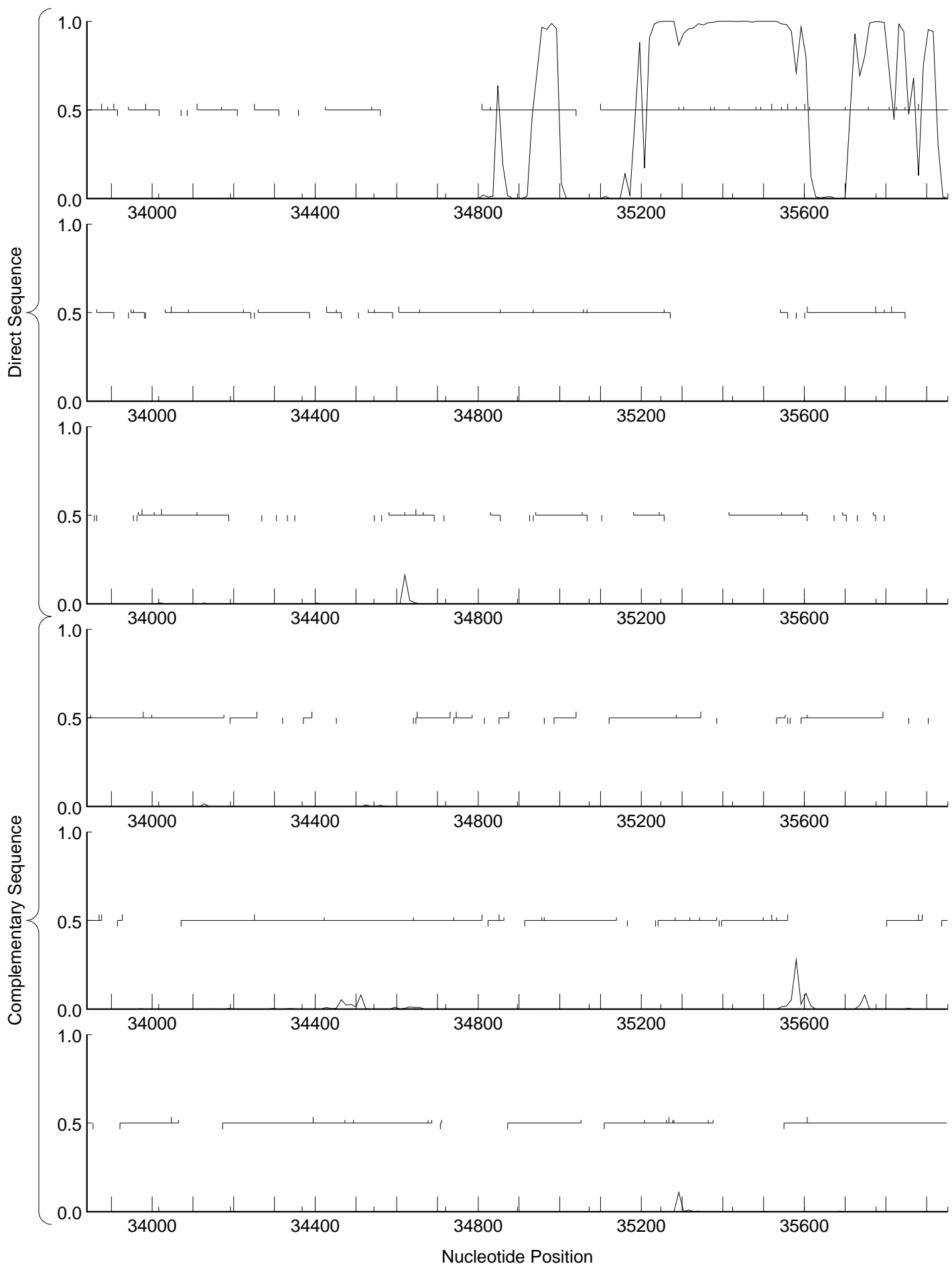


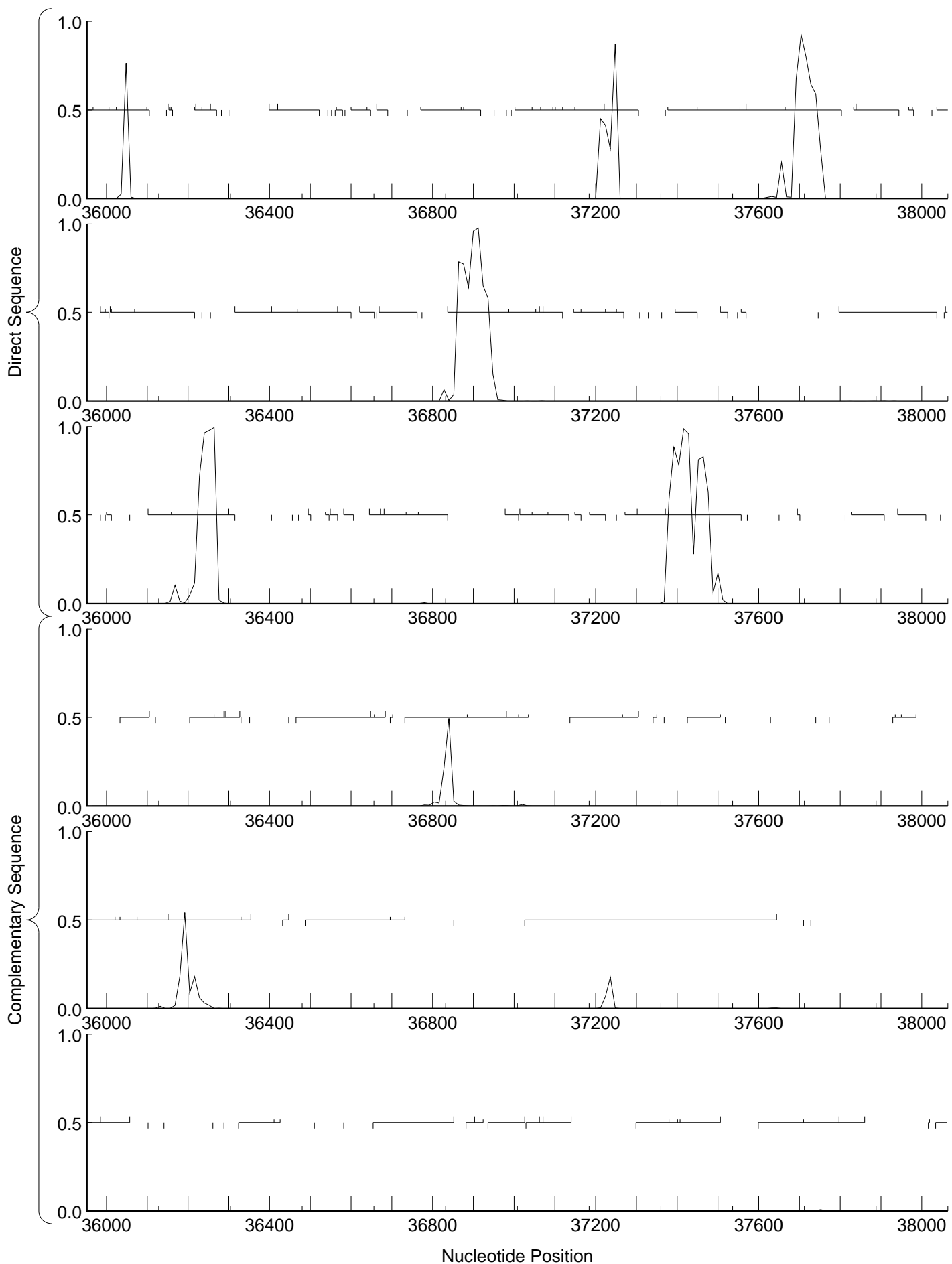
complete sequence, 55189 bp including 10-base 3' overhang (CGGTAGGCGC), Cluster F1, Order 4, Window 96, Step 12, 16/28

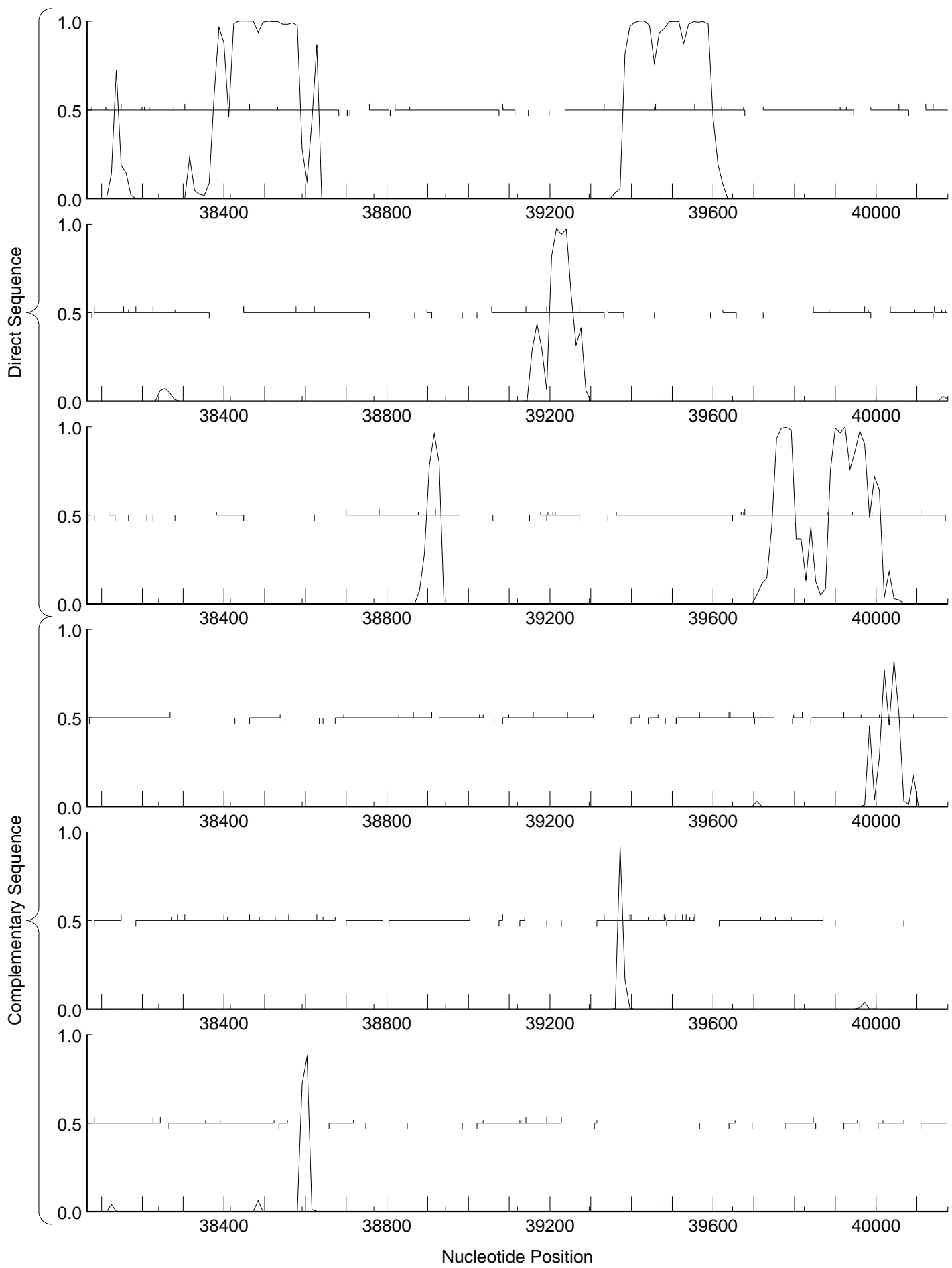


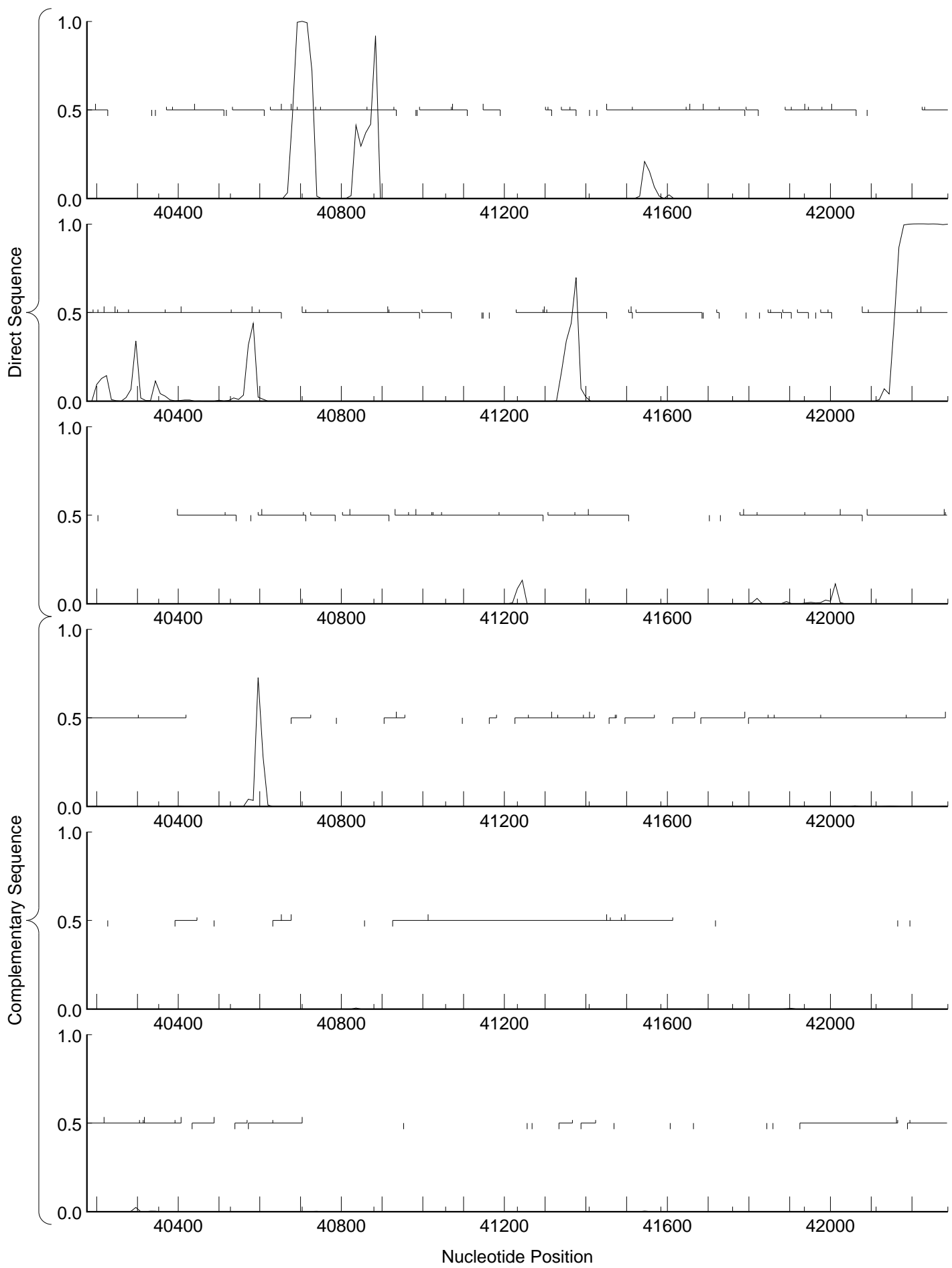
complete sequence, 55189 bp including 10-base 3' overhang (CGGTAGGCGC), Cluster F1, Order 4, Window 96, Step 12, 17/28

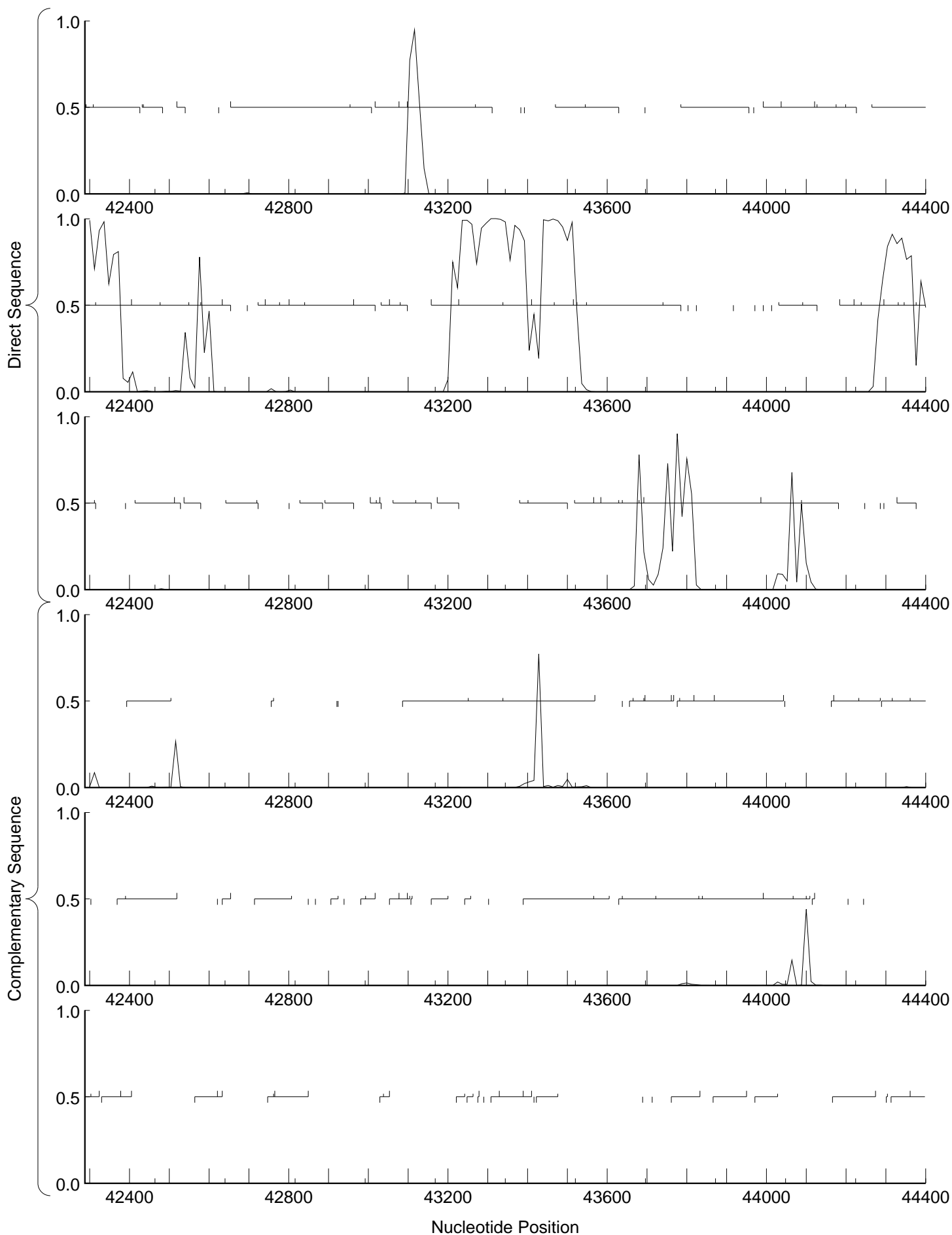




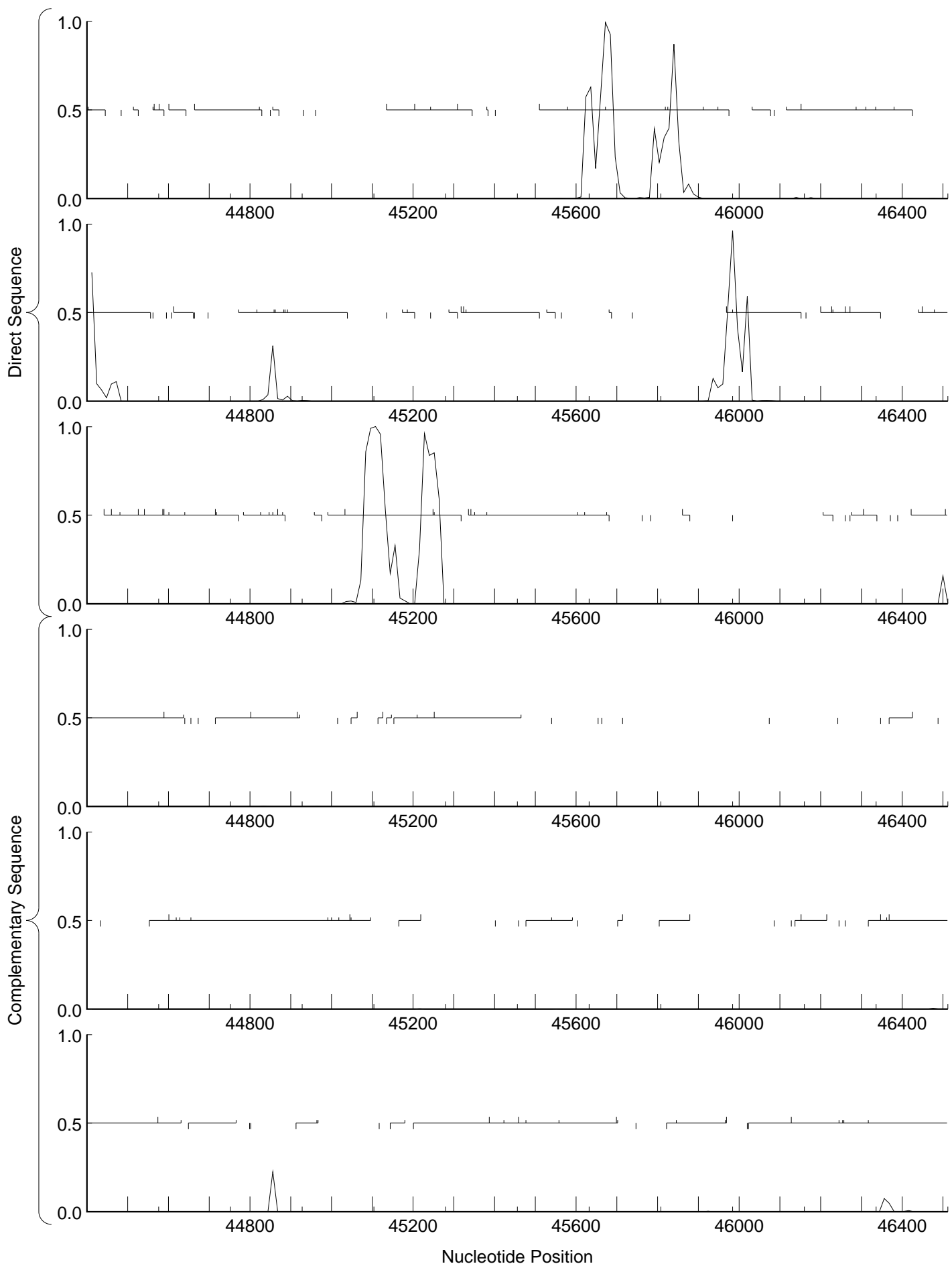


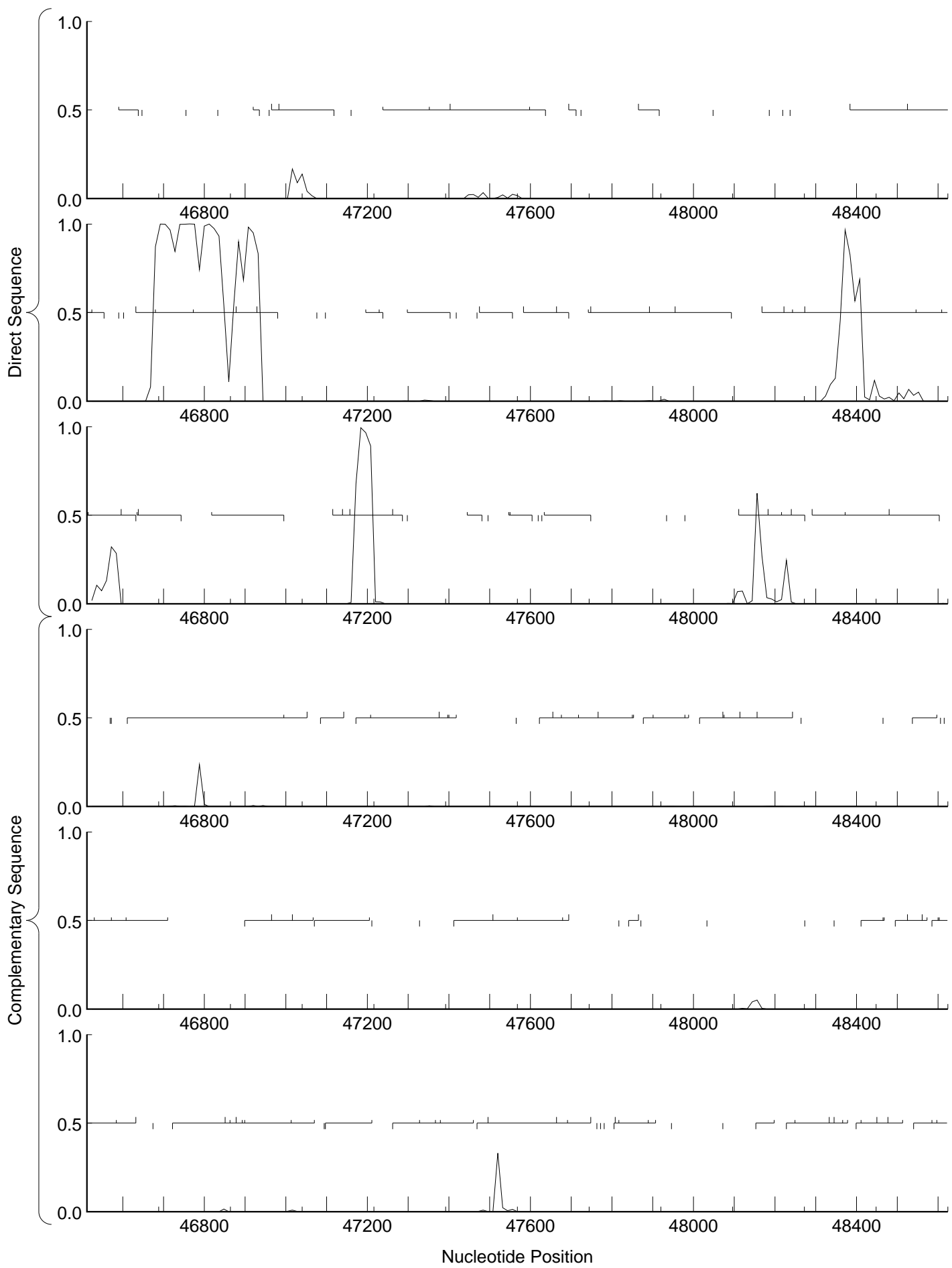


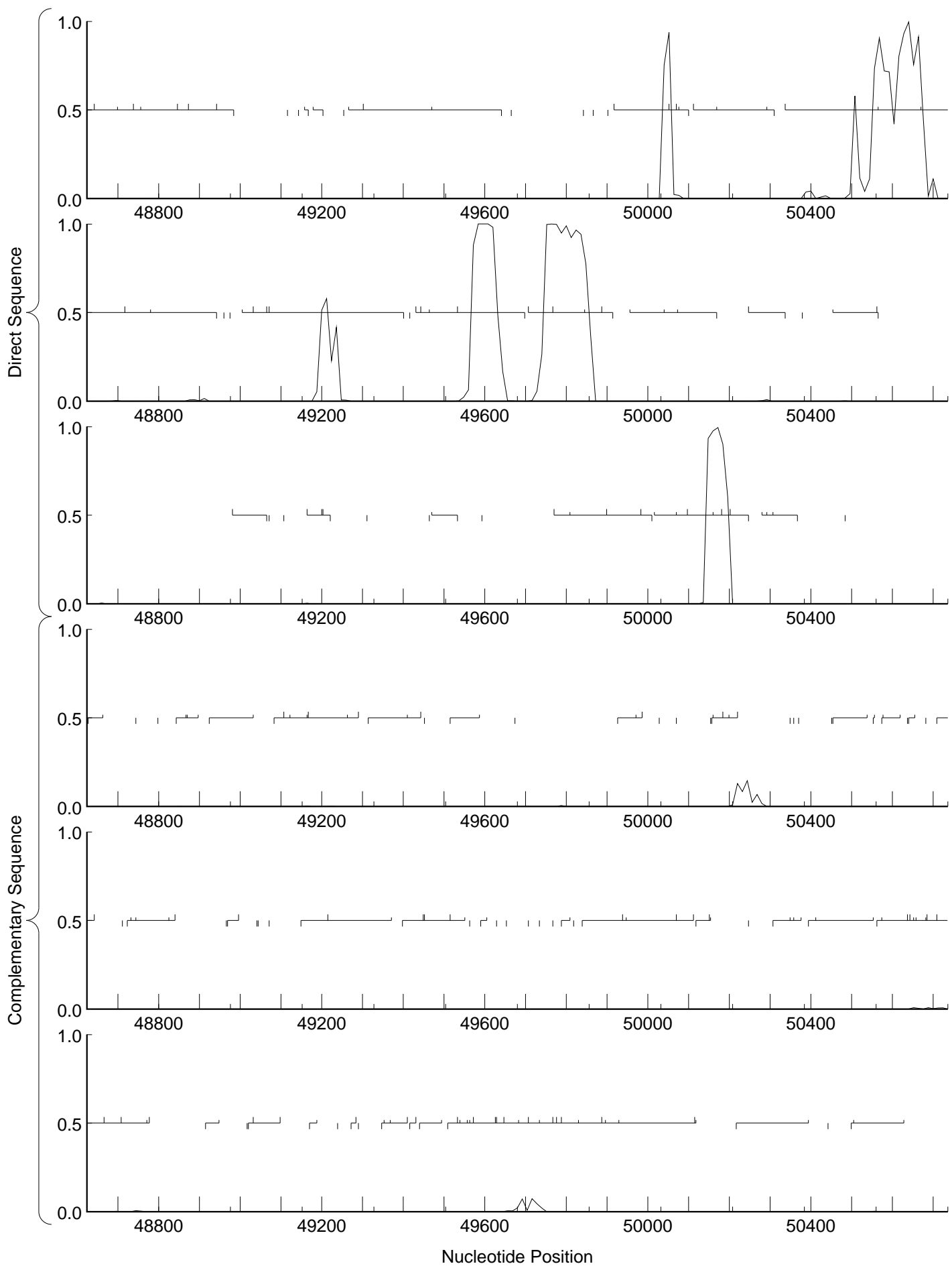


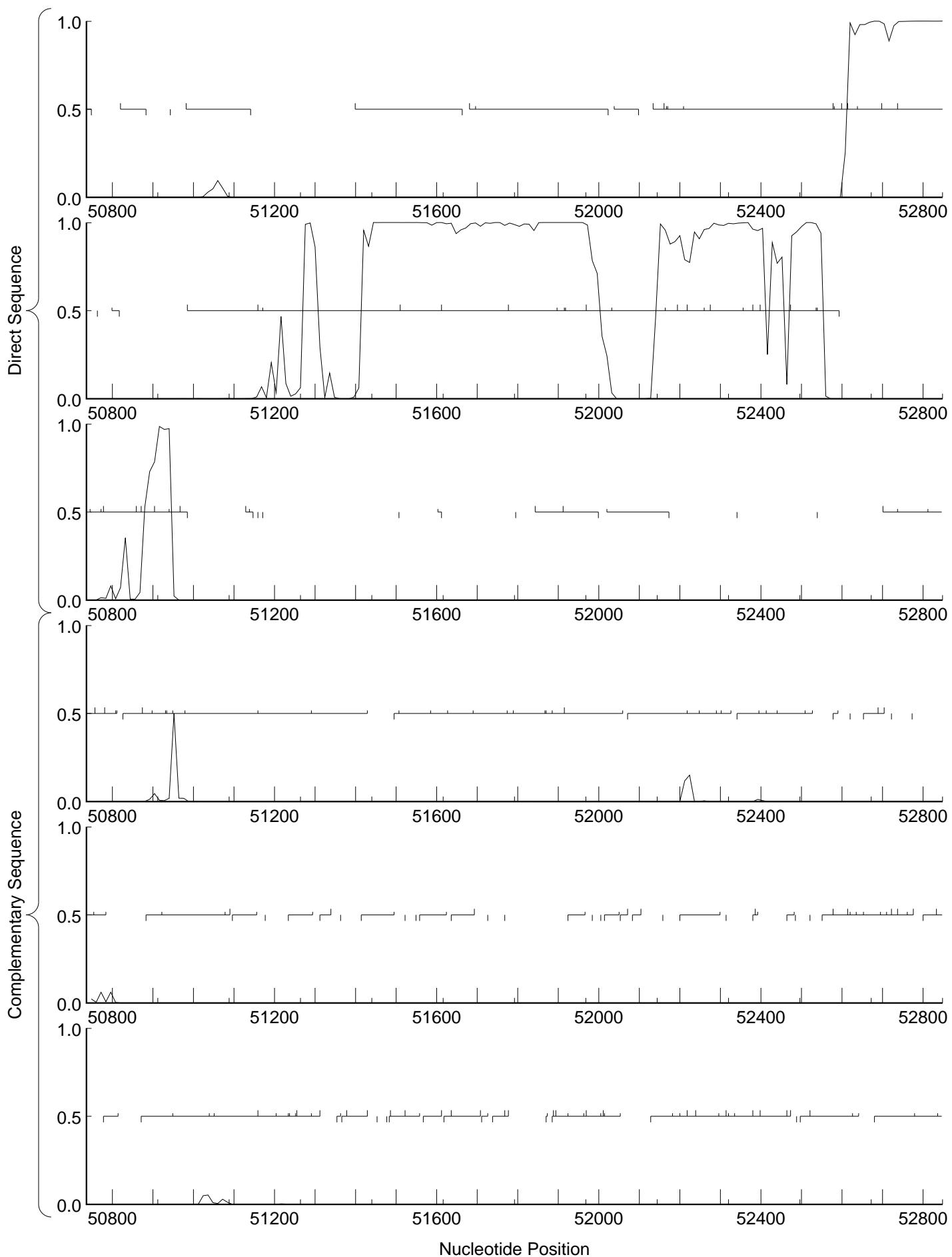


complete sequence, 55189 bp including 10-base 3' overhang (CGGTAGGCGC), Cluster F1, Order 4, Window 96, Step 12, 23/28









complete sequence, 55189 bp including 10-base 3' overhang (CGGTAGGCGC), Cluster F1, Order 4, Window 96, Step 12, 27/28

