

GeneMark

Version 2.5p (09.08.06)

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PROGRAM INFORMATION

Sequence : Gordonia phage Mossy complete sequence, 61183 including 9-base 3' overhang (CGCCGCCCT), Clu
Analysis Date : 4/27/25 at 22:24:44
Pages : 30
Sequence Length : 61183 bp
GC Content : 51.95%

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,

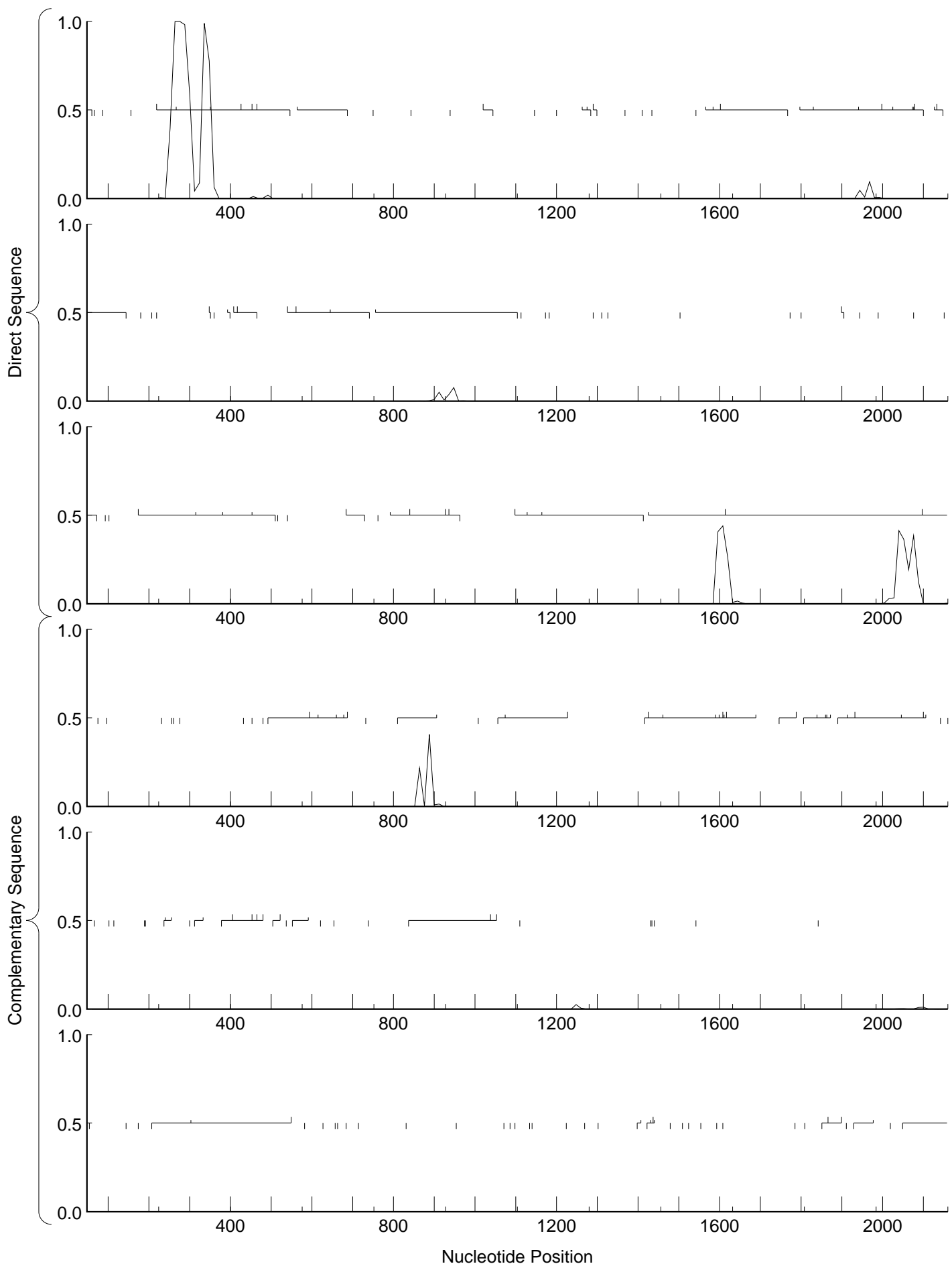
MATRIX INFORMATION

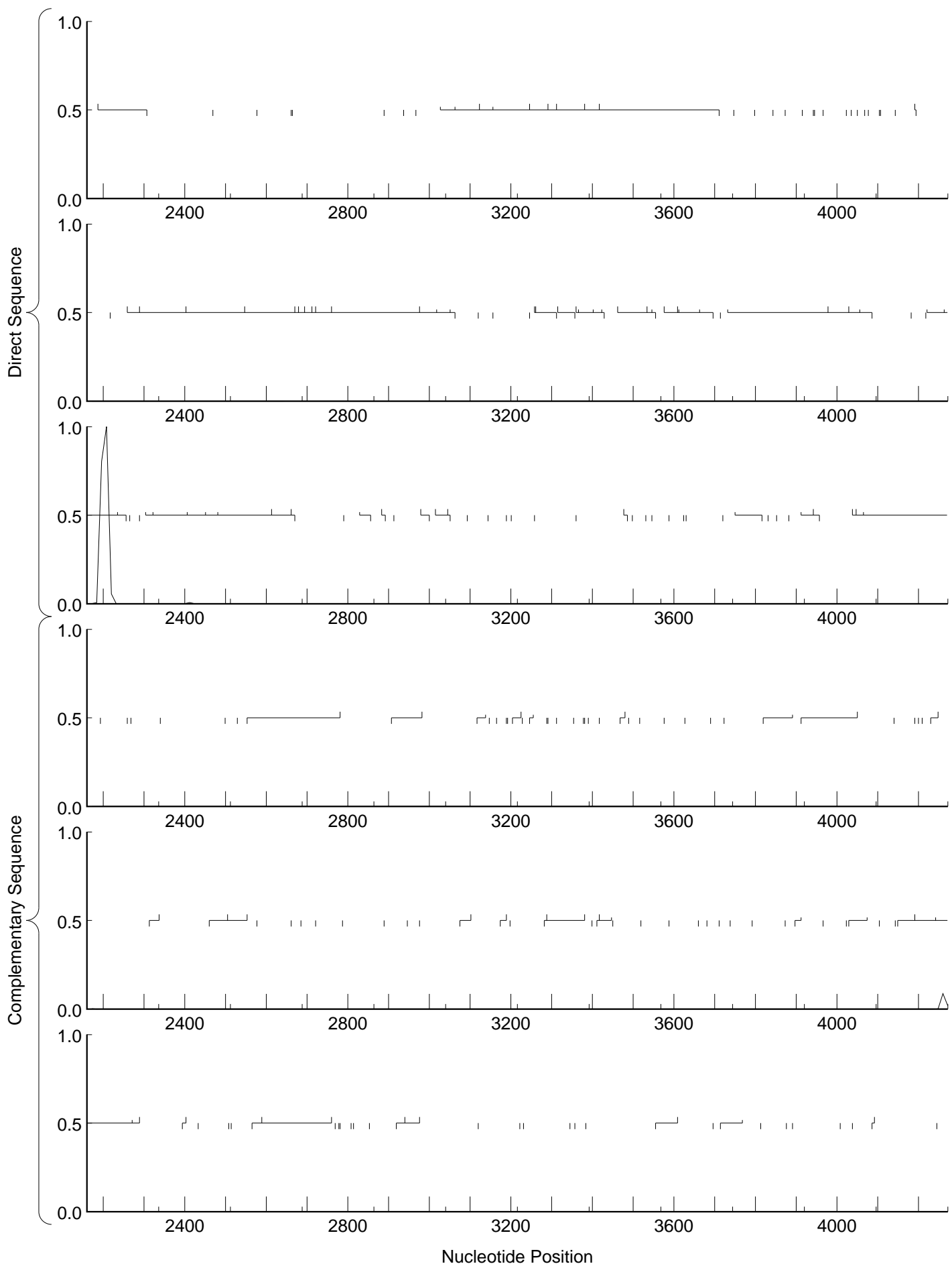
Matrix : -
Author : -
Order : 4

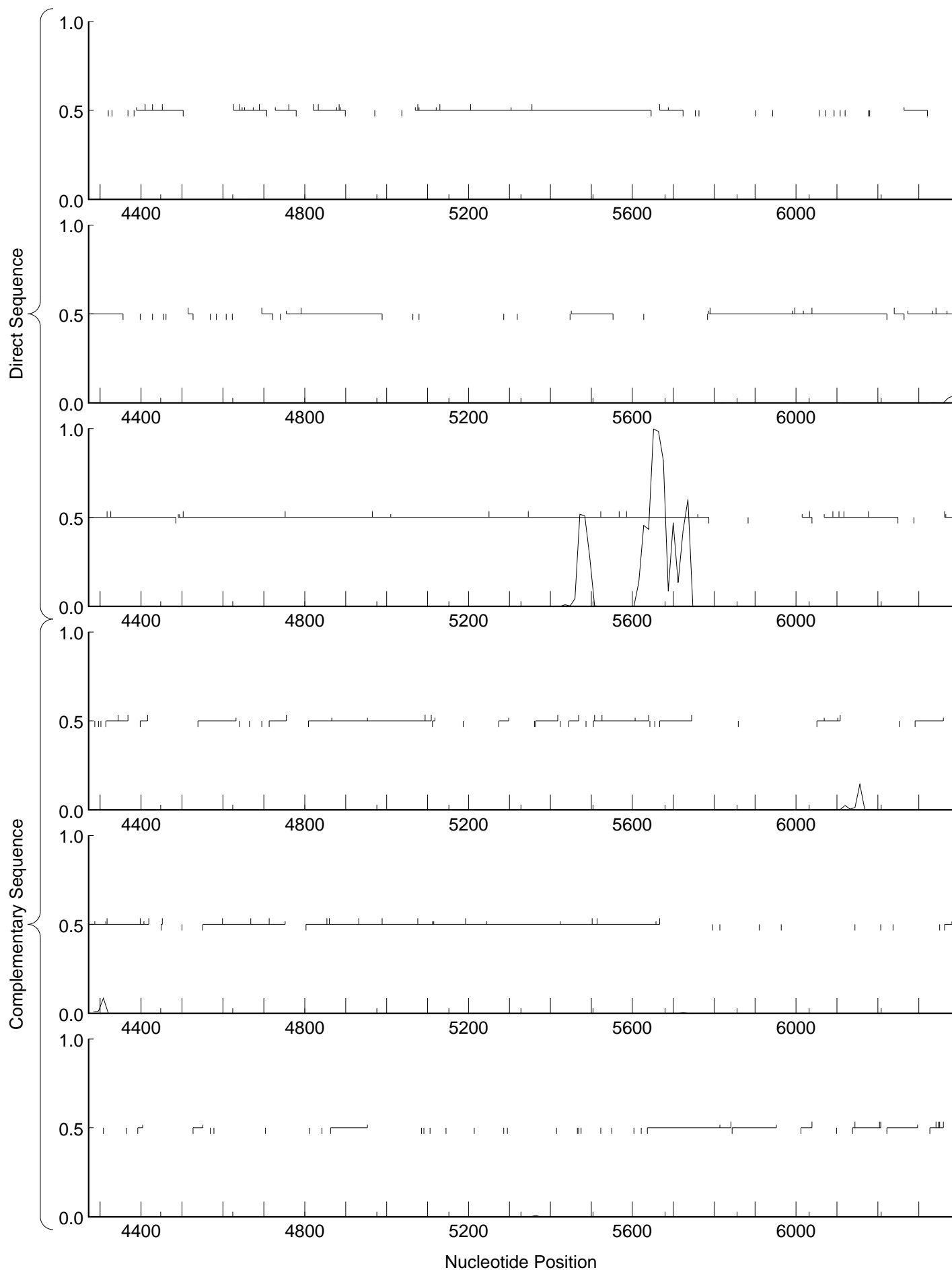
Send questions / comments to:
Dr. M. Borodovsky
Georgia Institute of Technology
School of Biology
Atlanta, GA 30332-0230

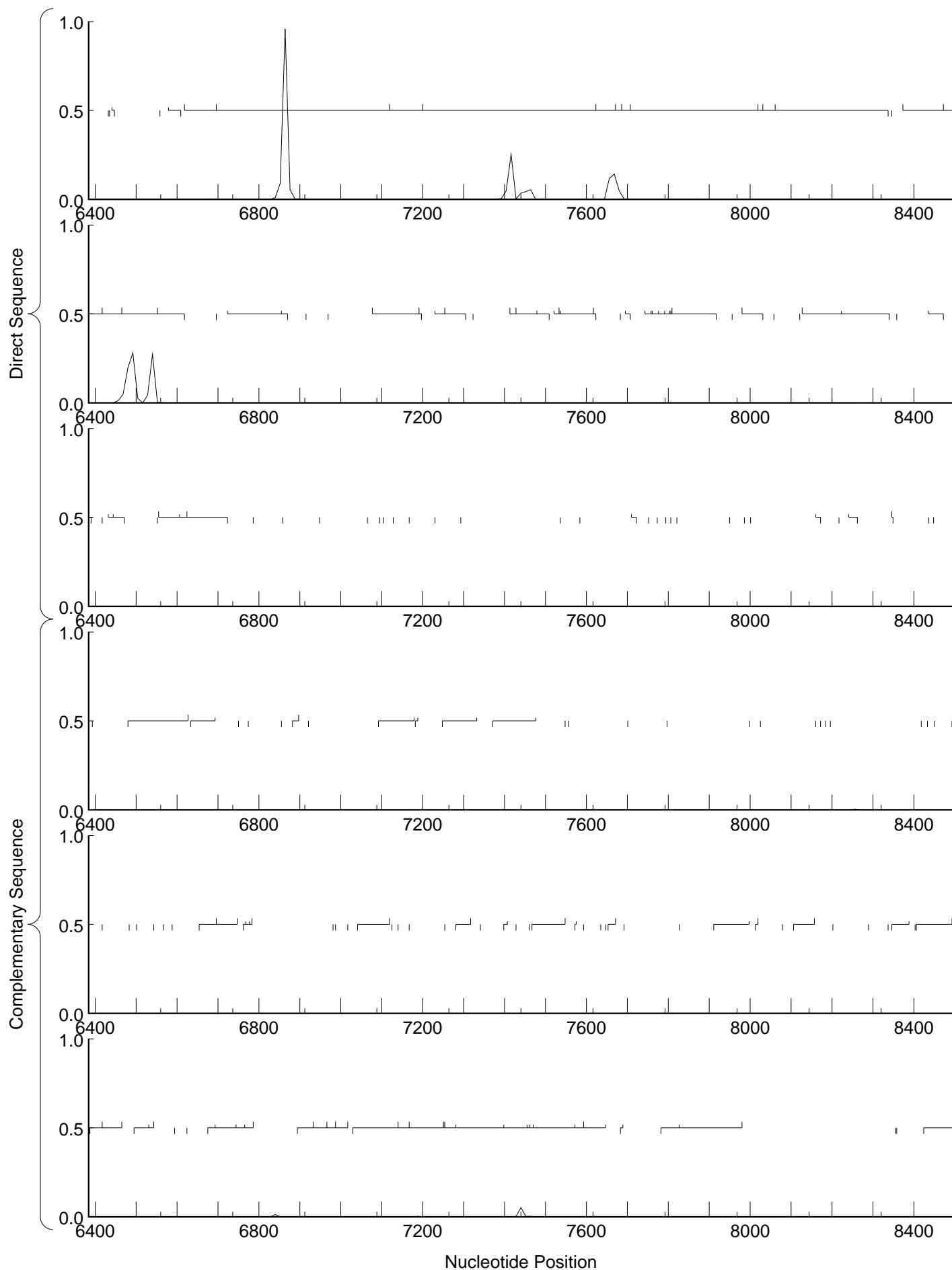
Matrix notes & comments

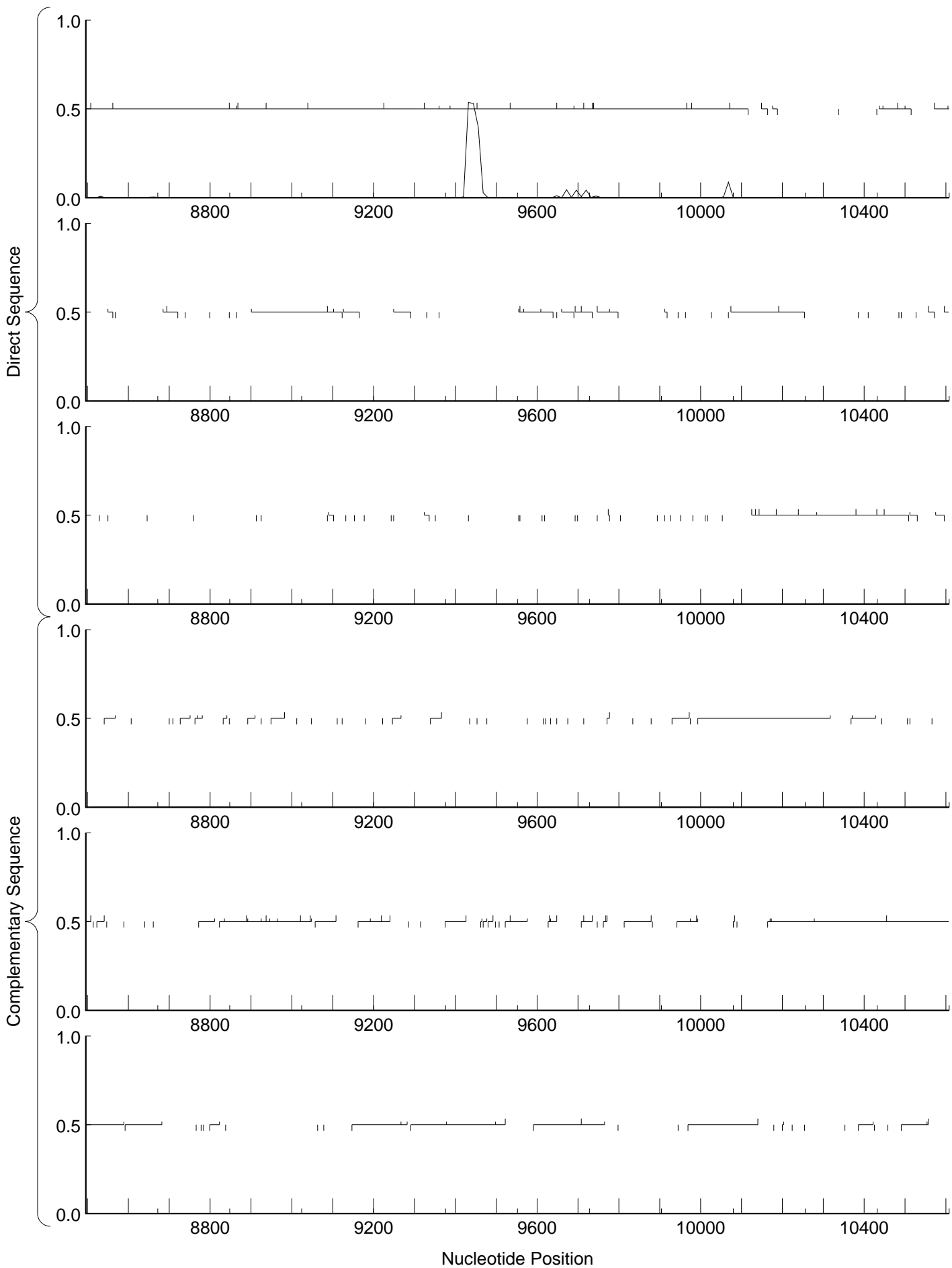
Training set derived by GeneMarkS, 4.30 October 2014
Sat Sep 8 00:36:14 2018

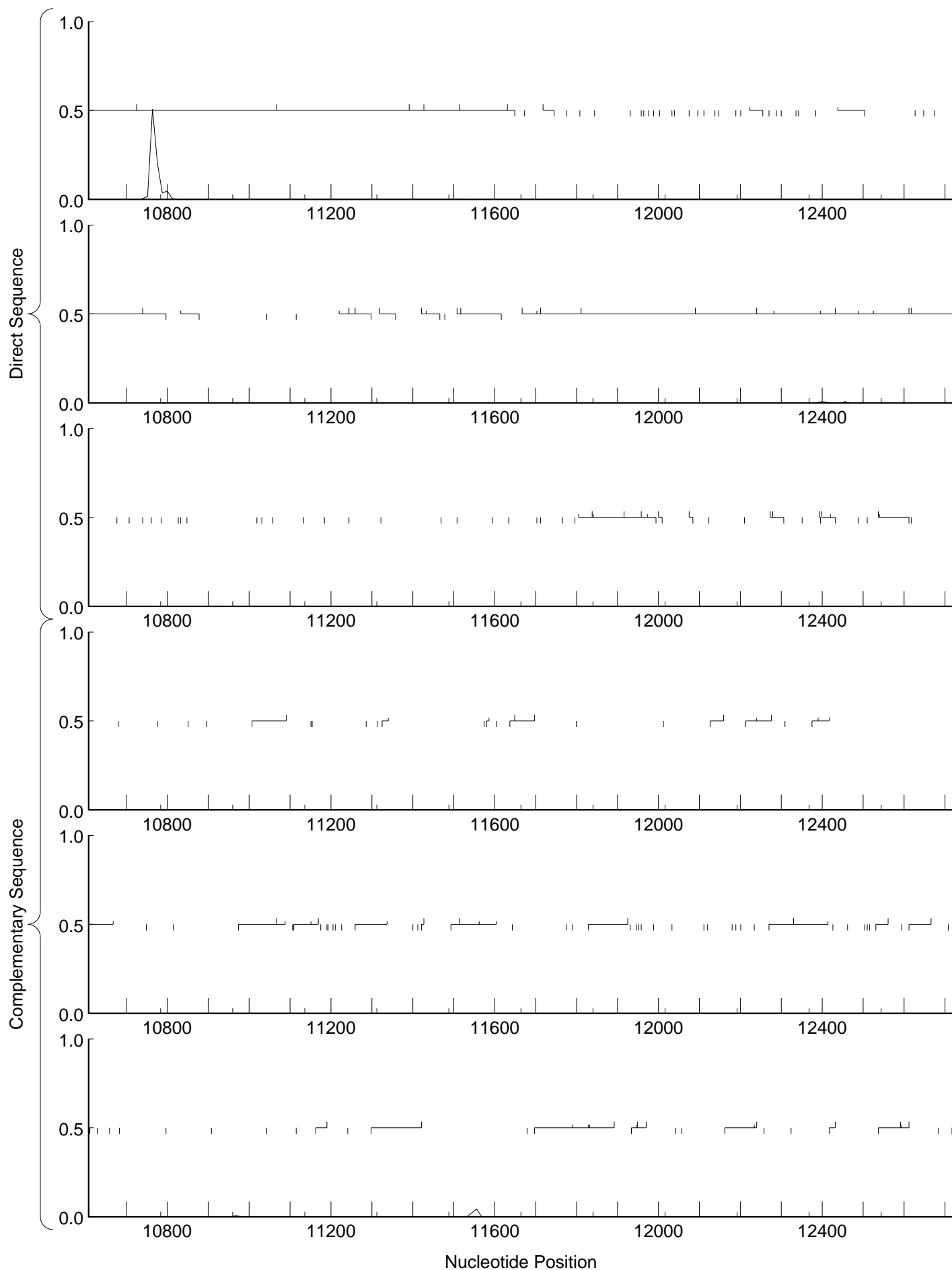


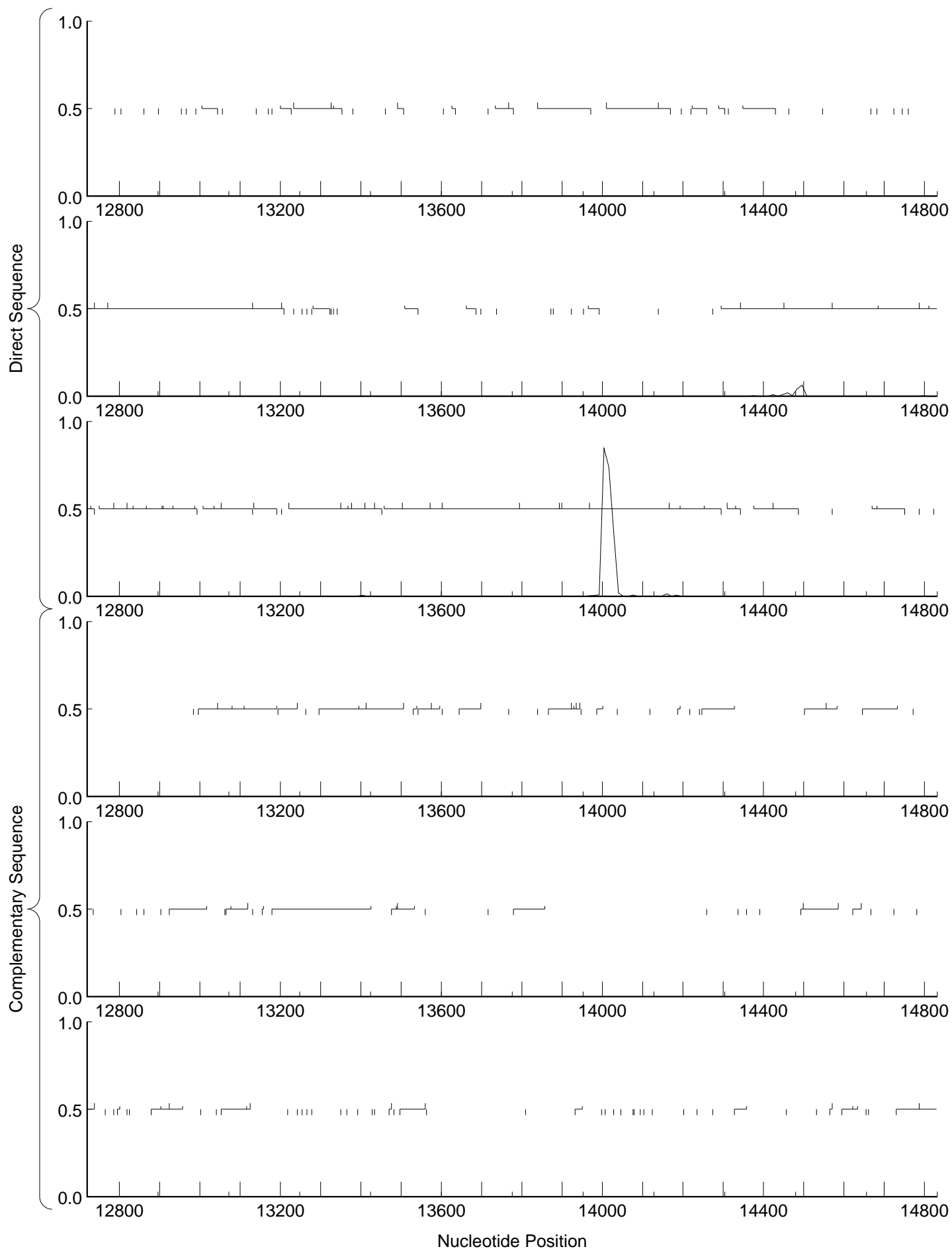


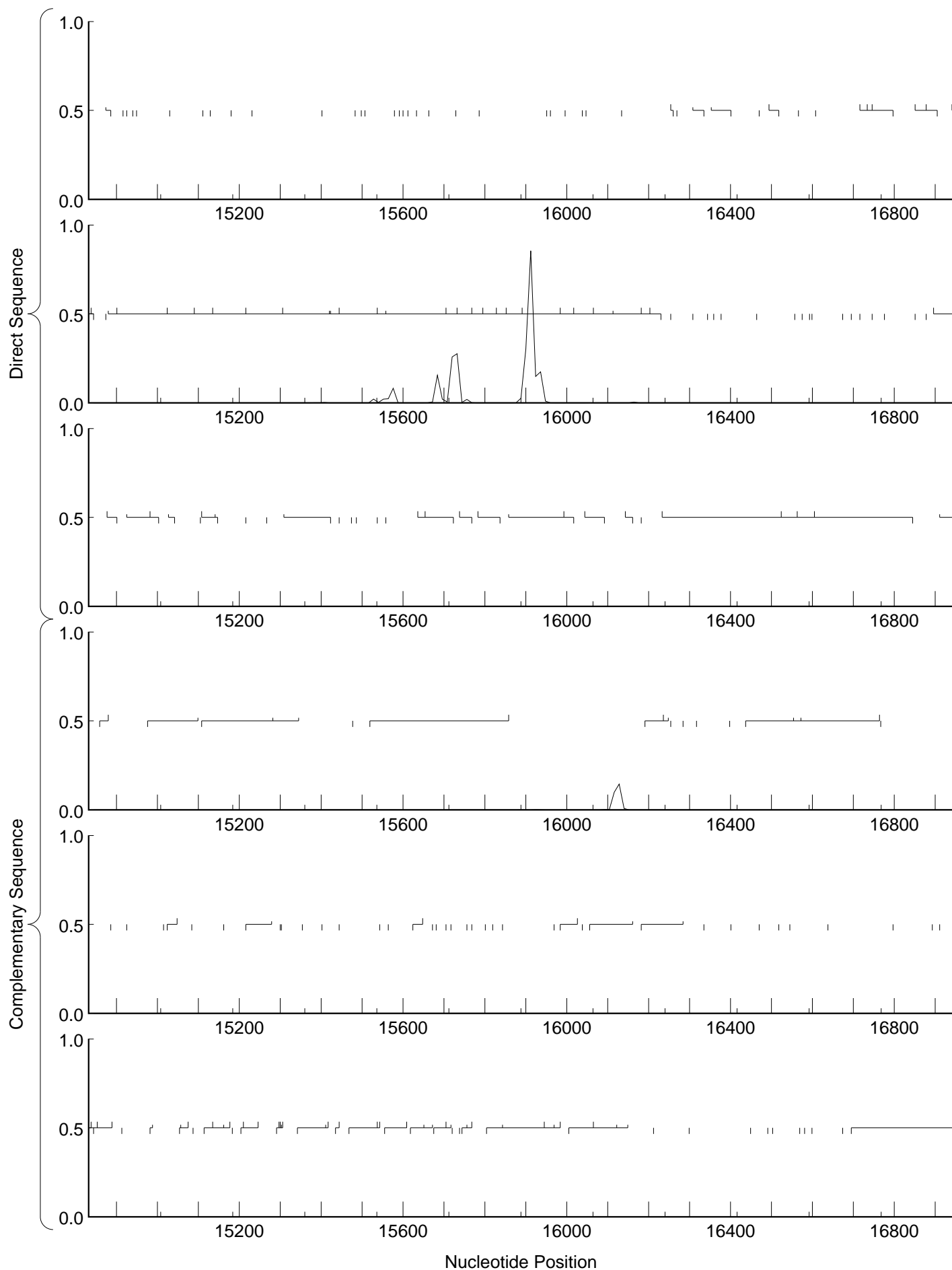


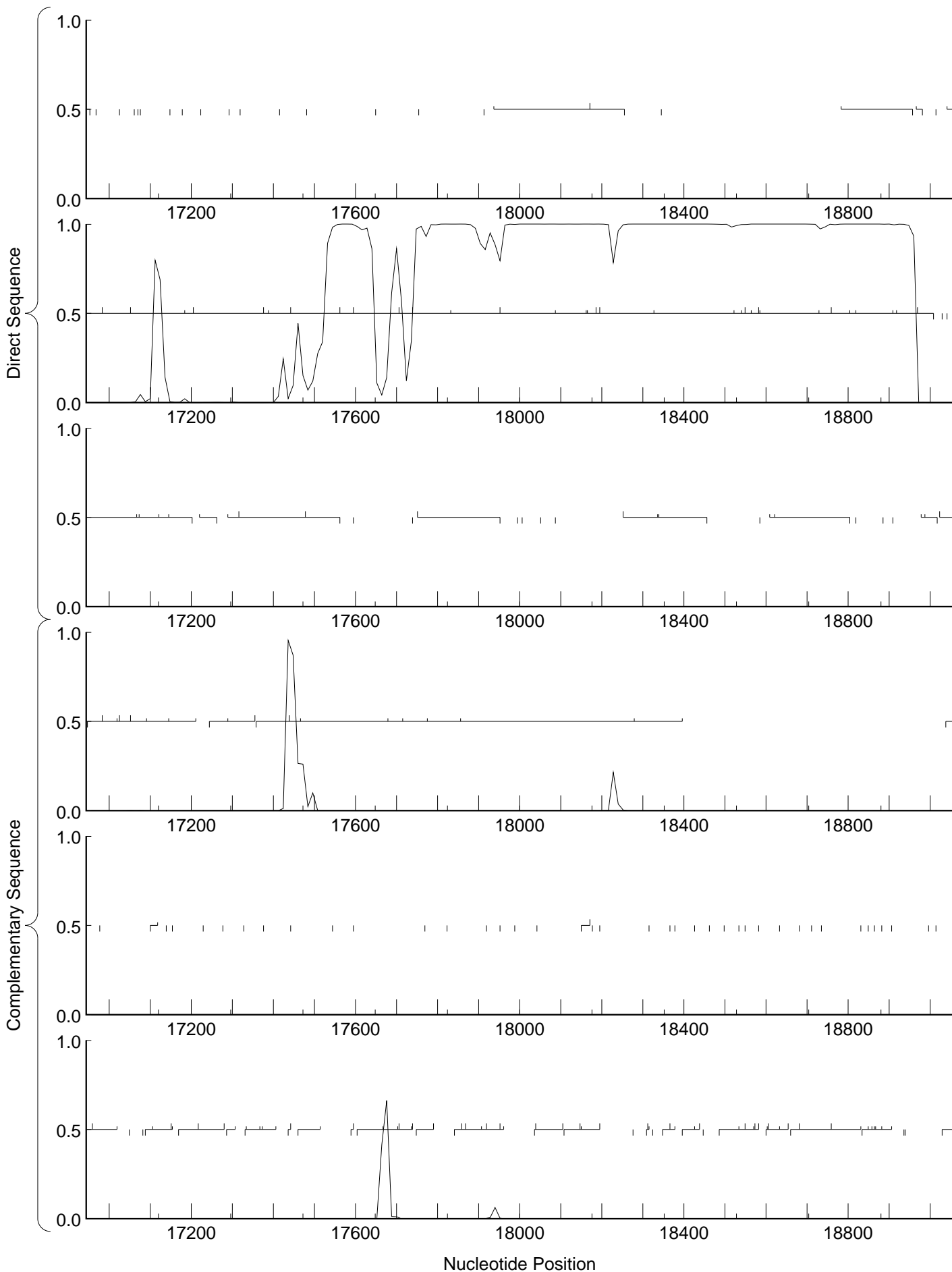


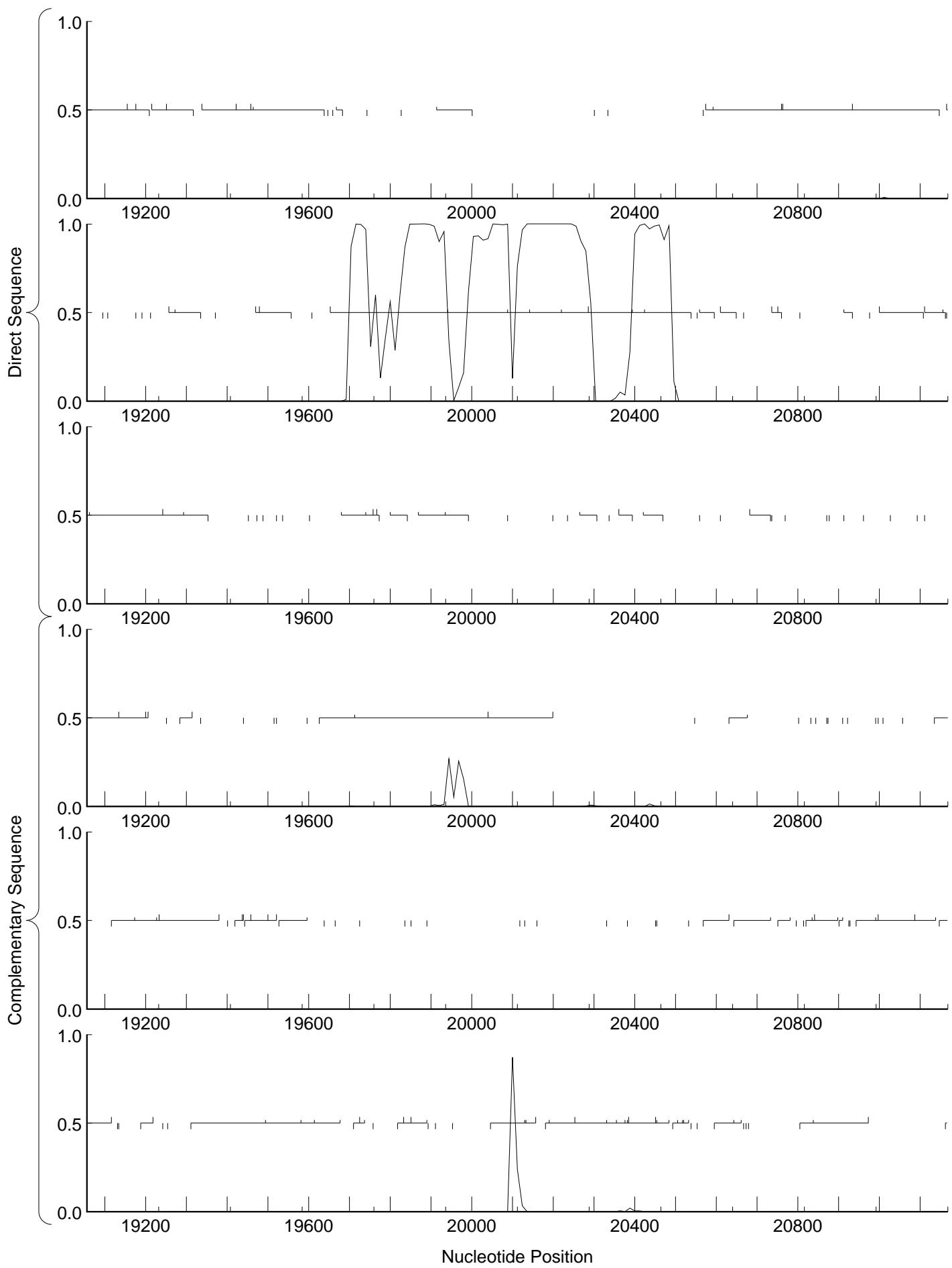


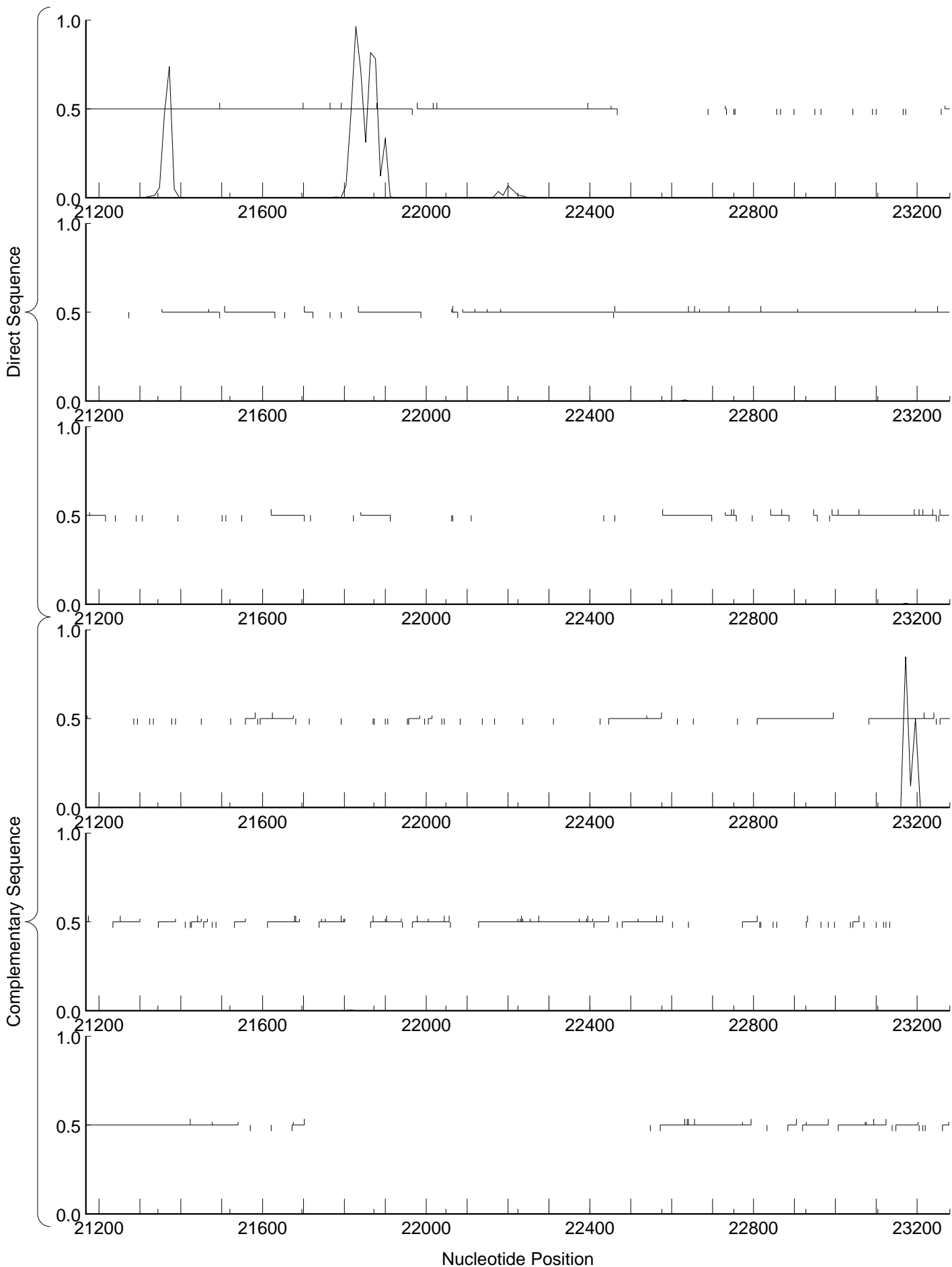


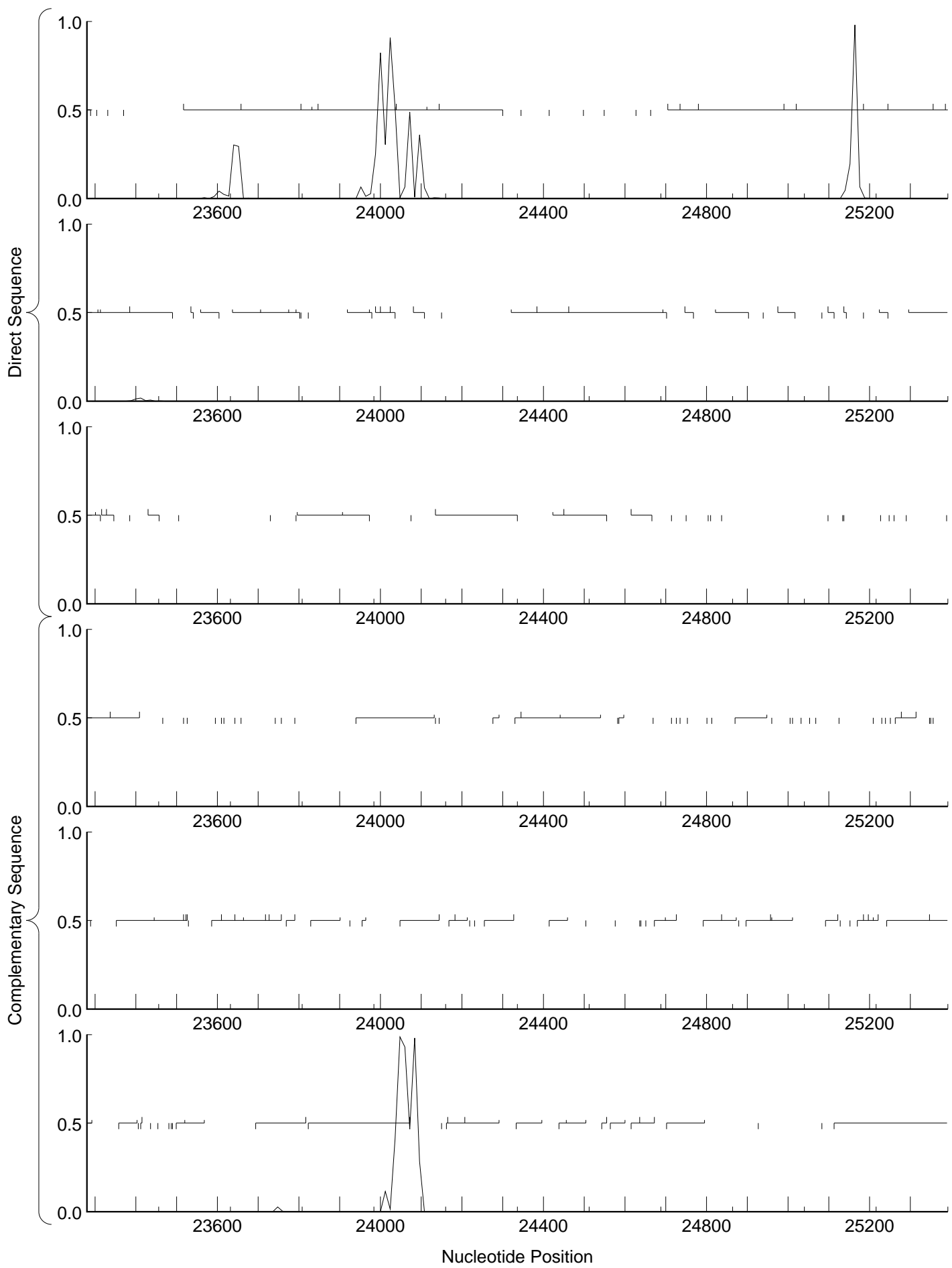


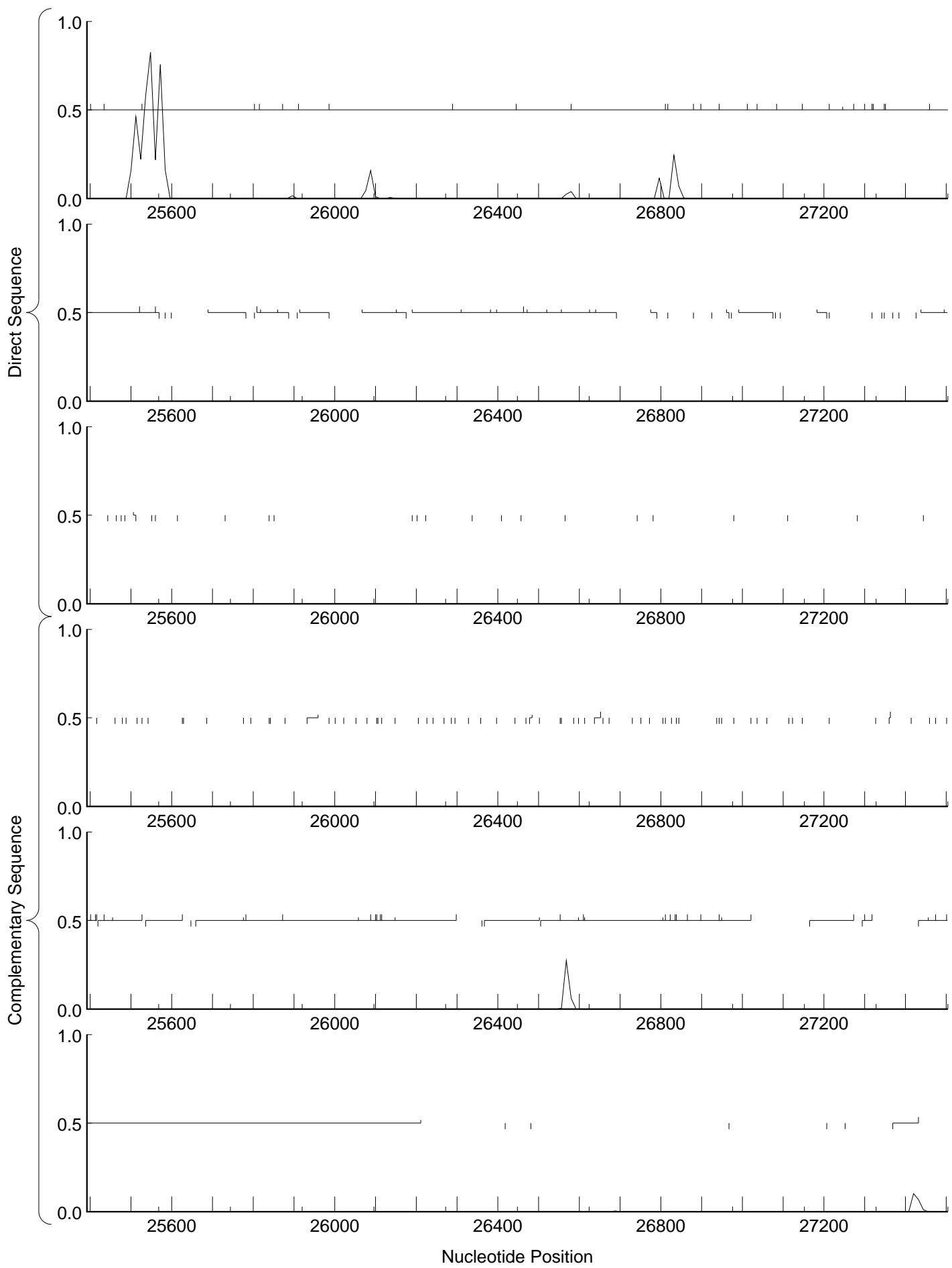




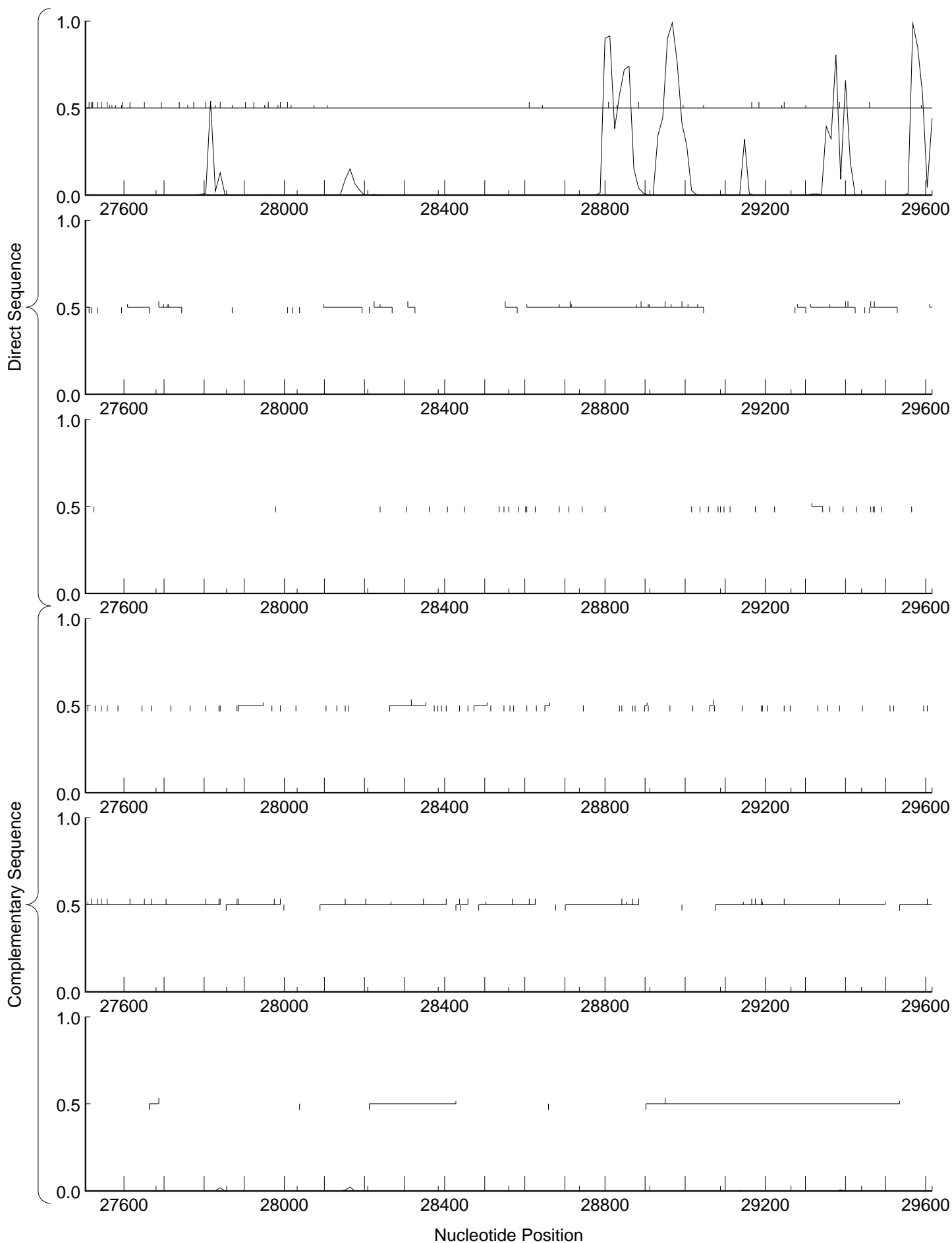


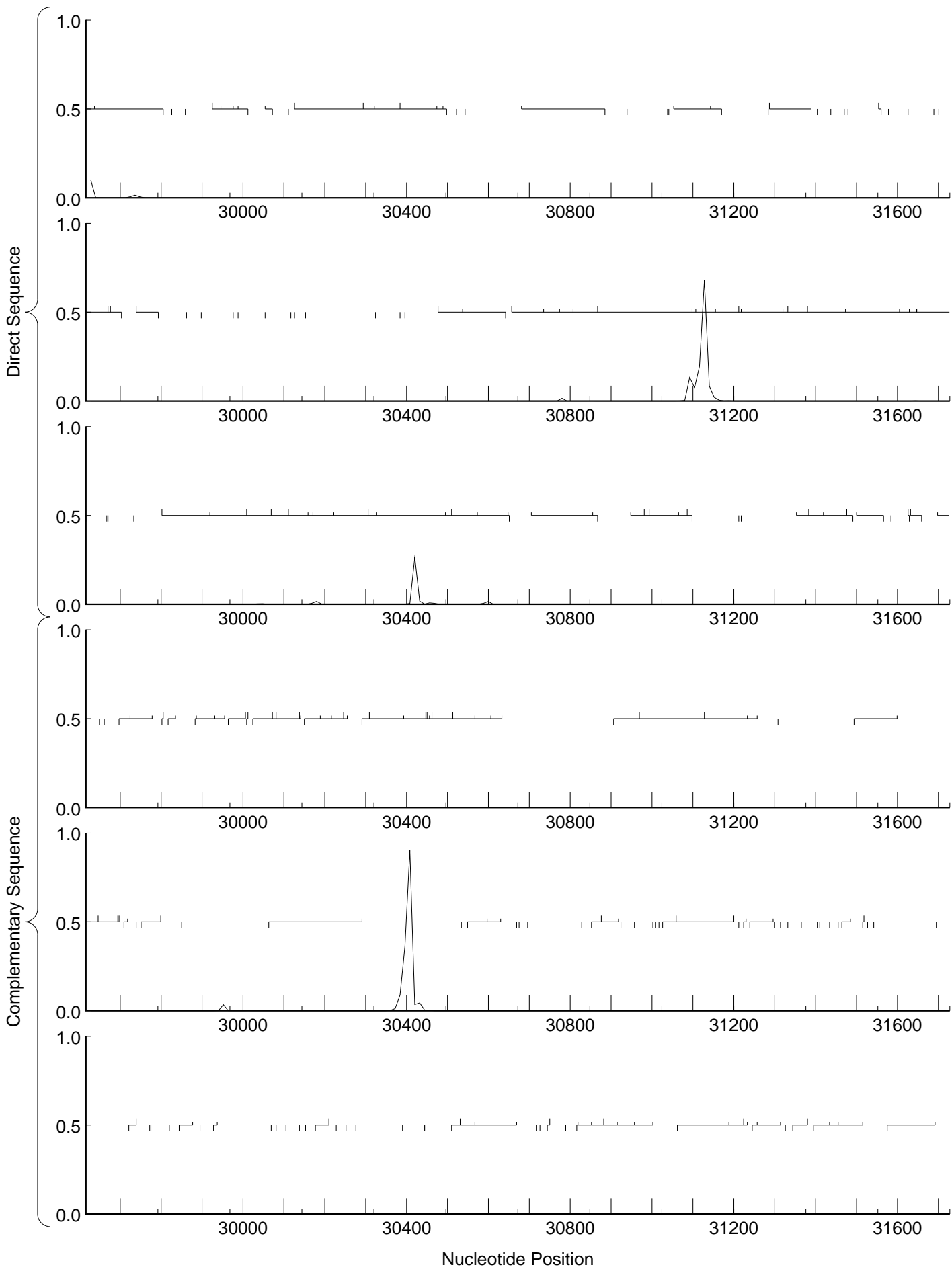


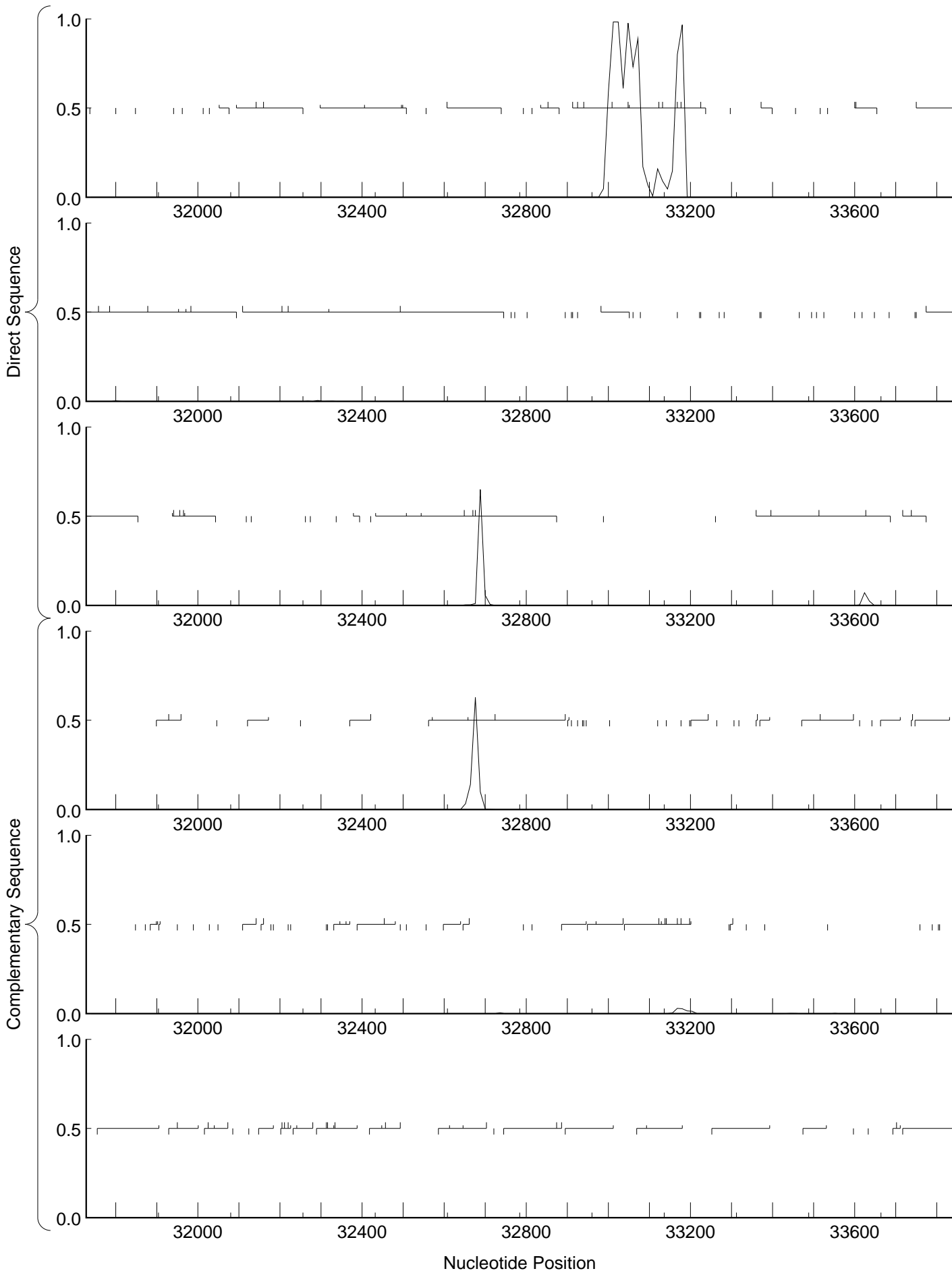


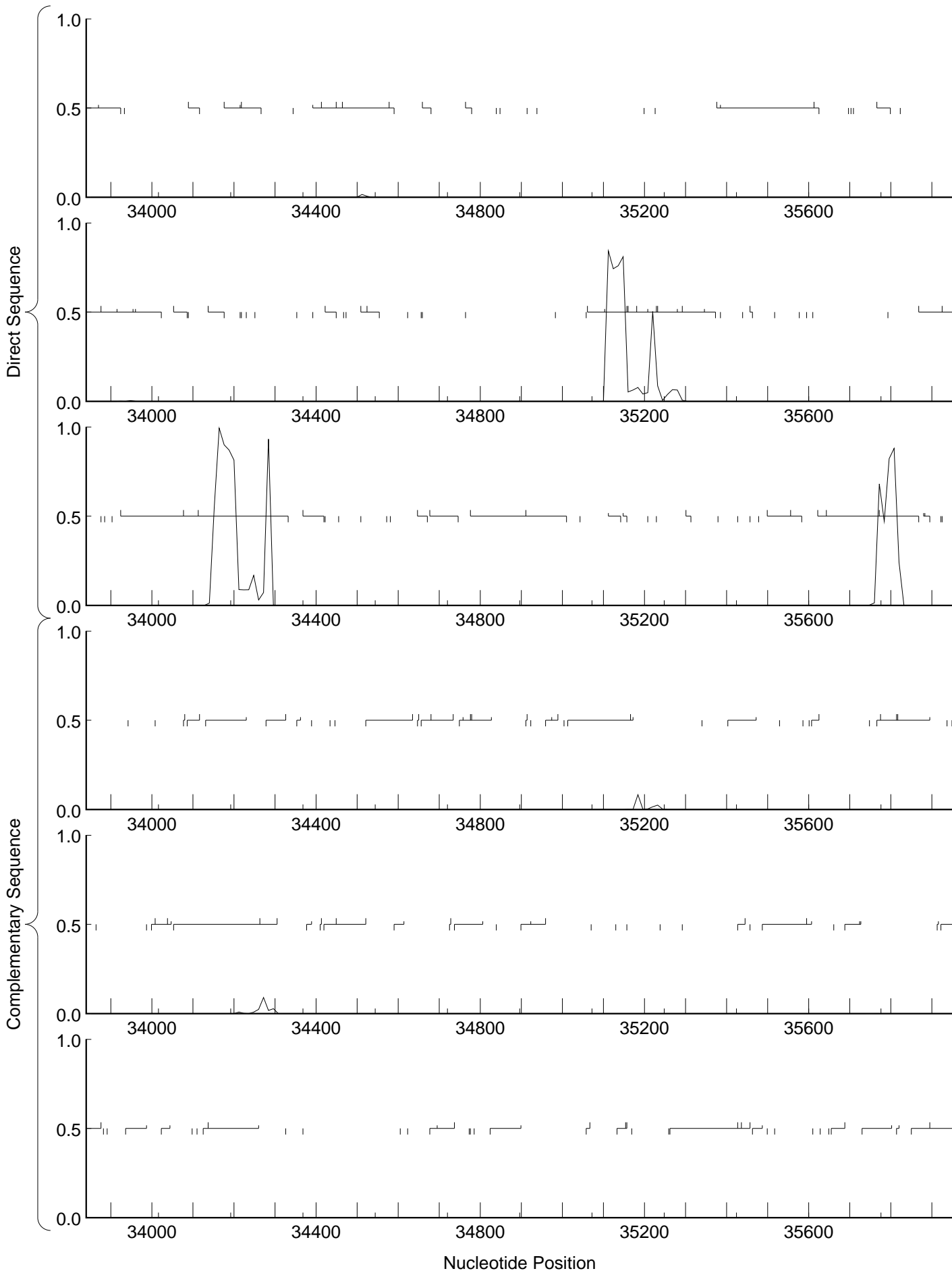


Mossy complete sequence, 61183 including 9-base 3' overhang (CGCCGCCCT), Cluster DJ, Order 4, Window 96, Step 12, 15/30

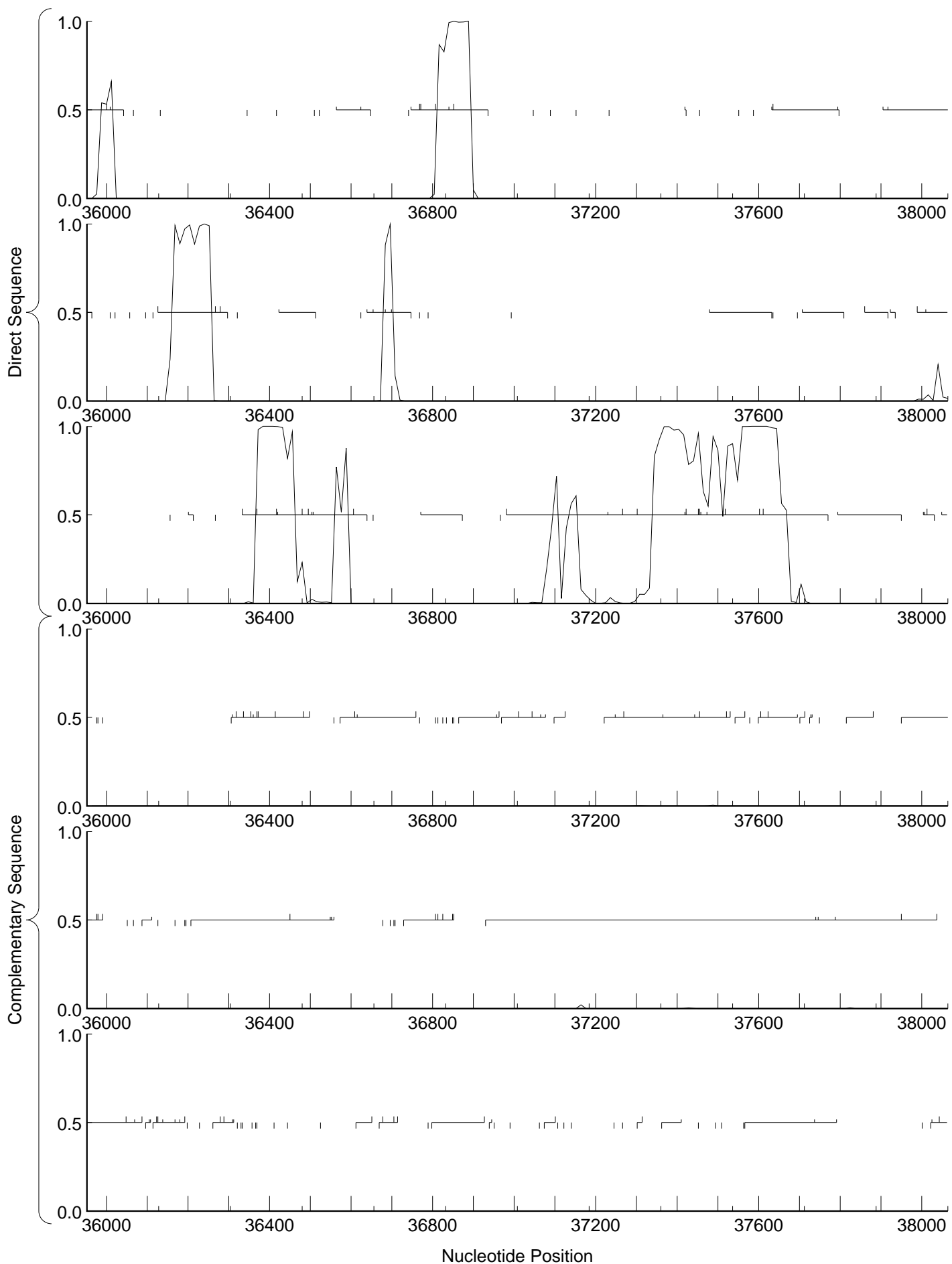


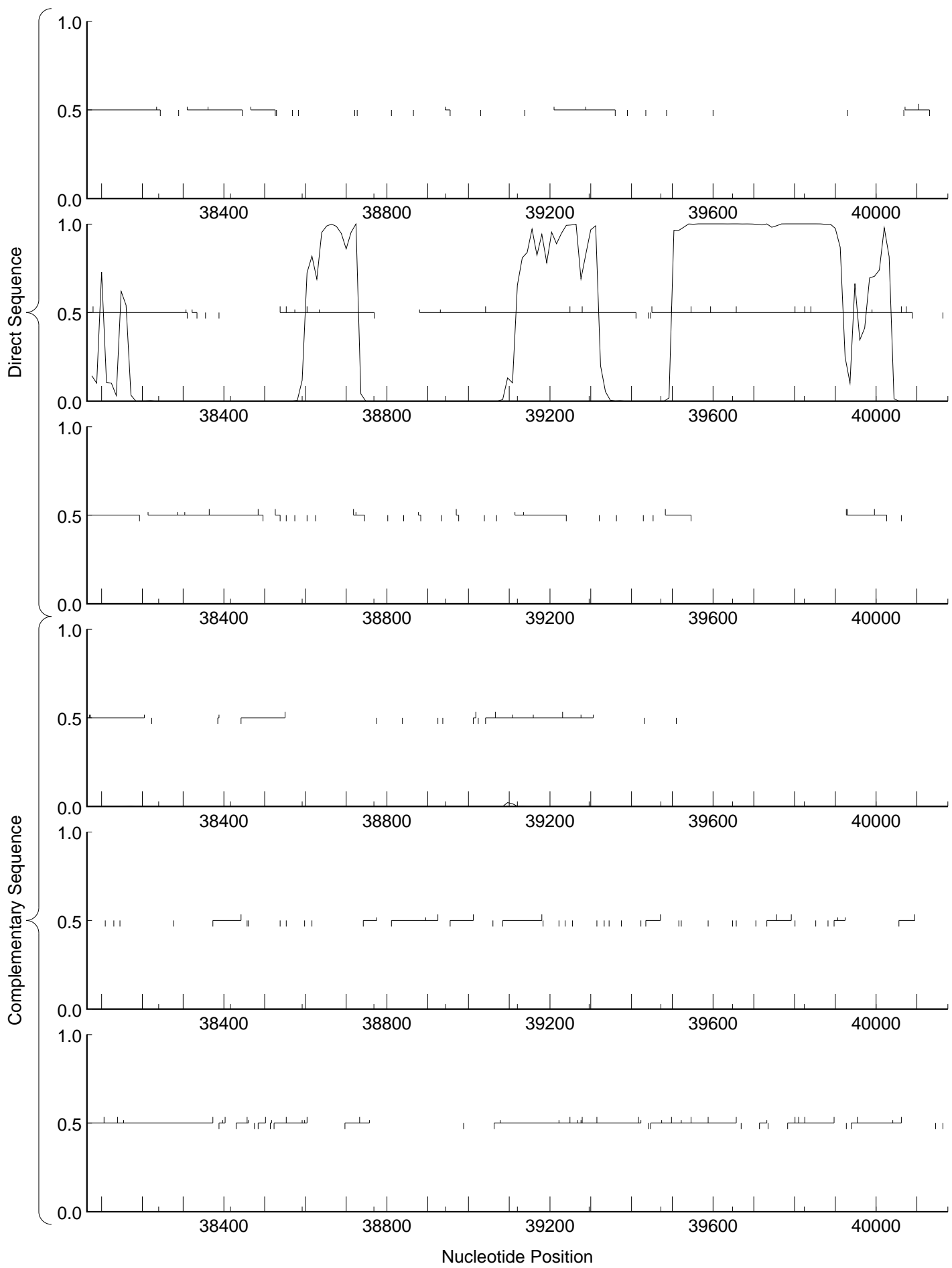


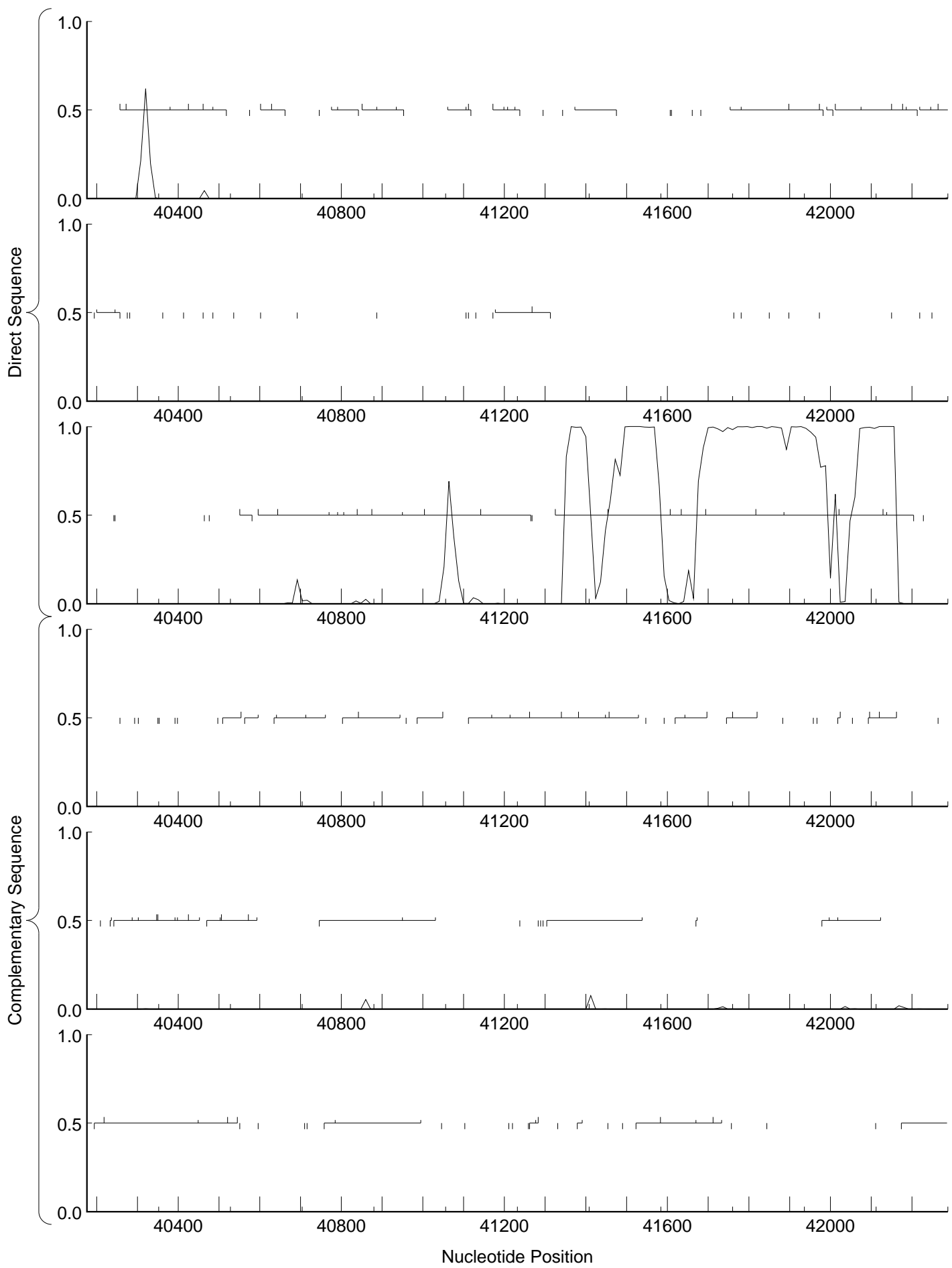


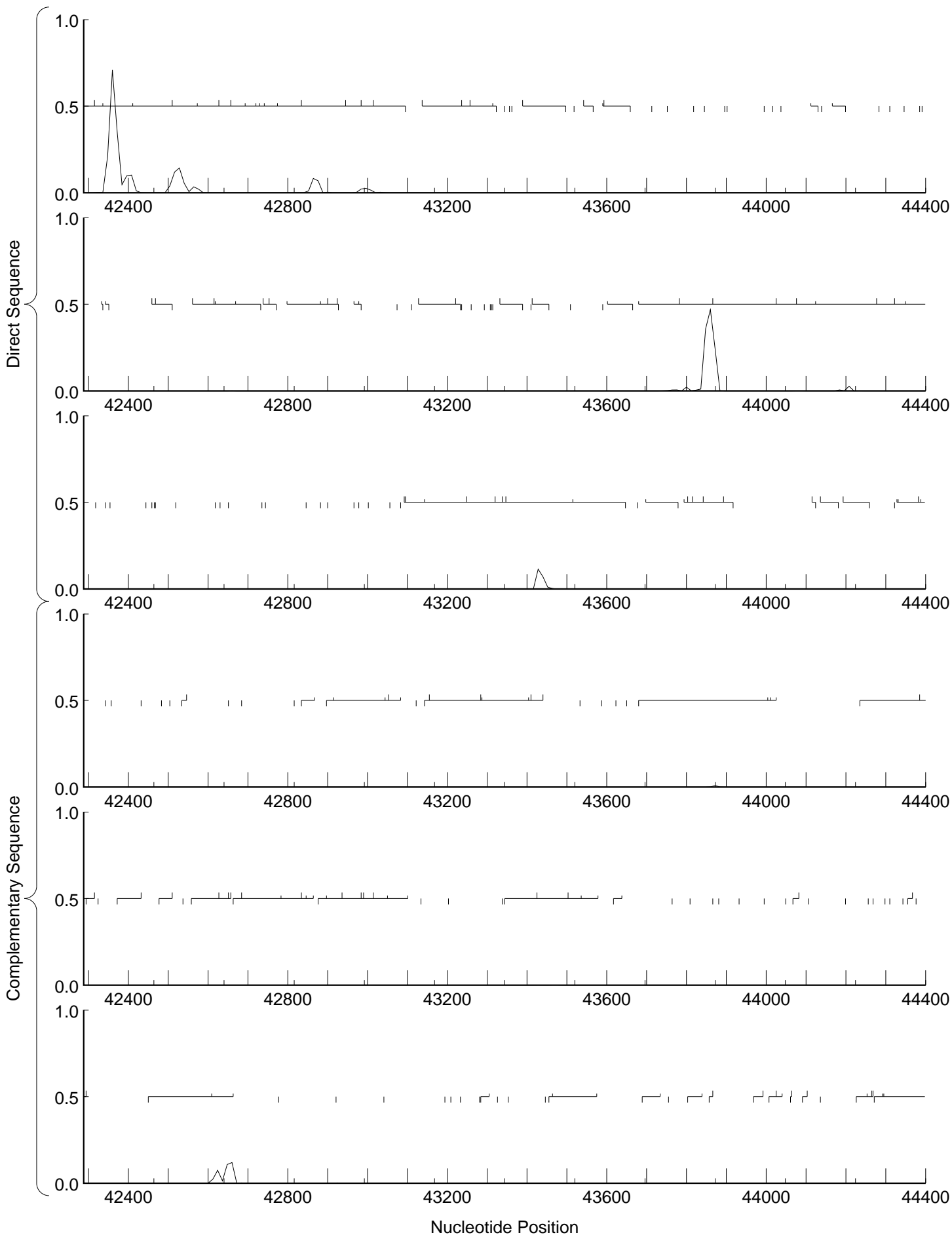


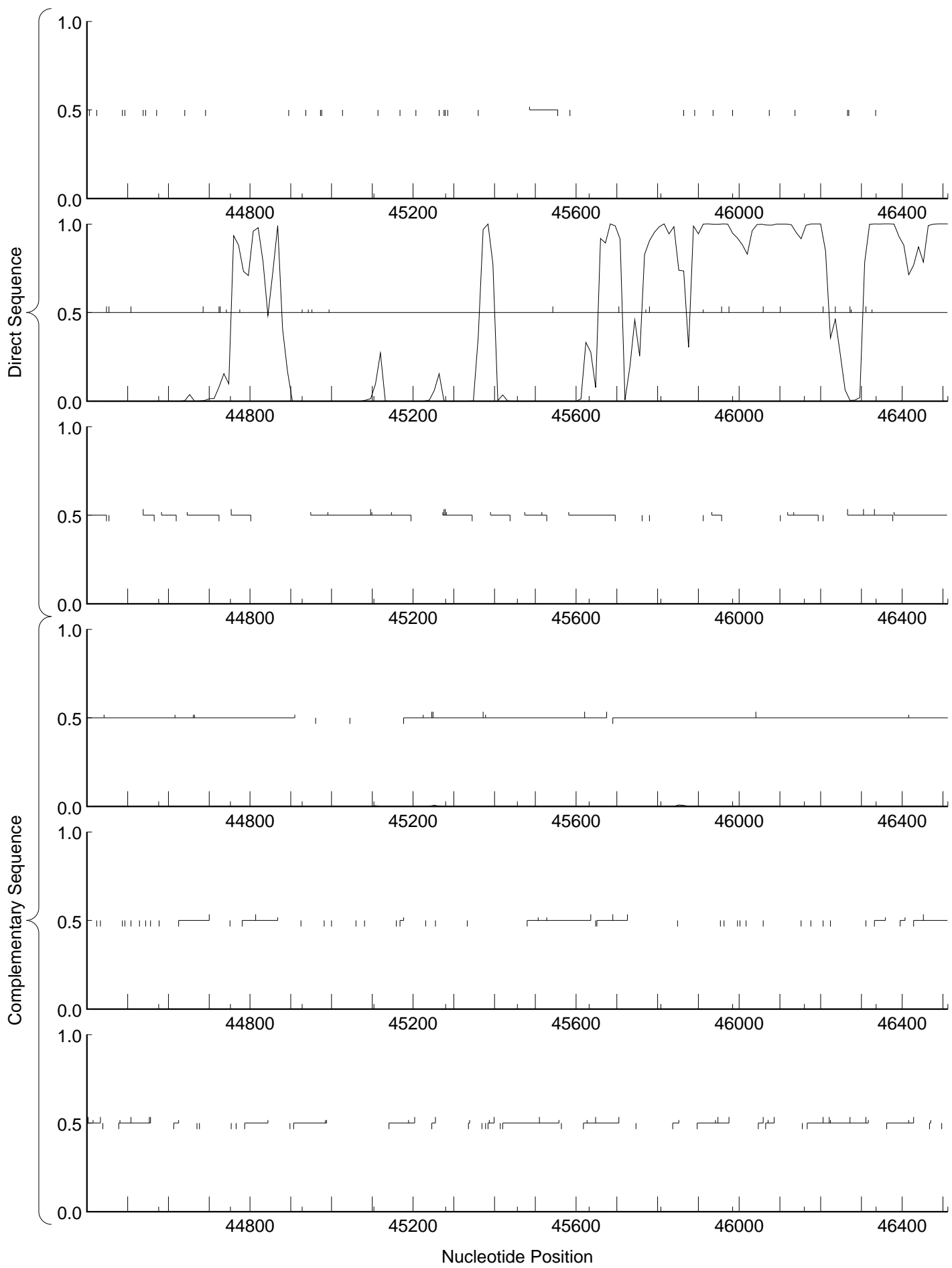
Mossy complete sequence, 61183 including 9-base 3' overhang (CGCCGCCCT), Cluster DJ, Order 4, Window 96, Step 12, 19/30

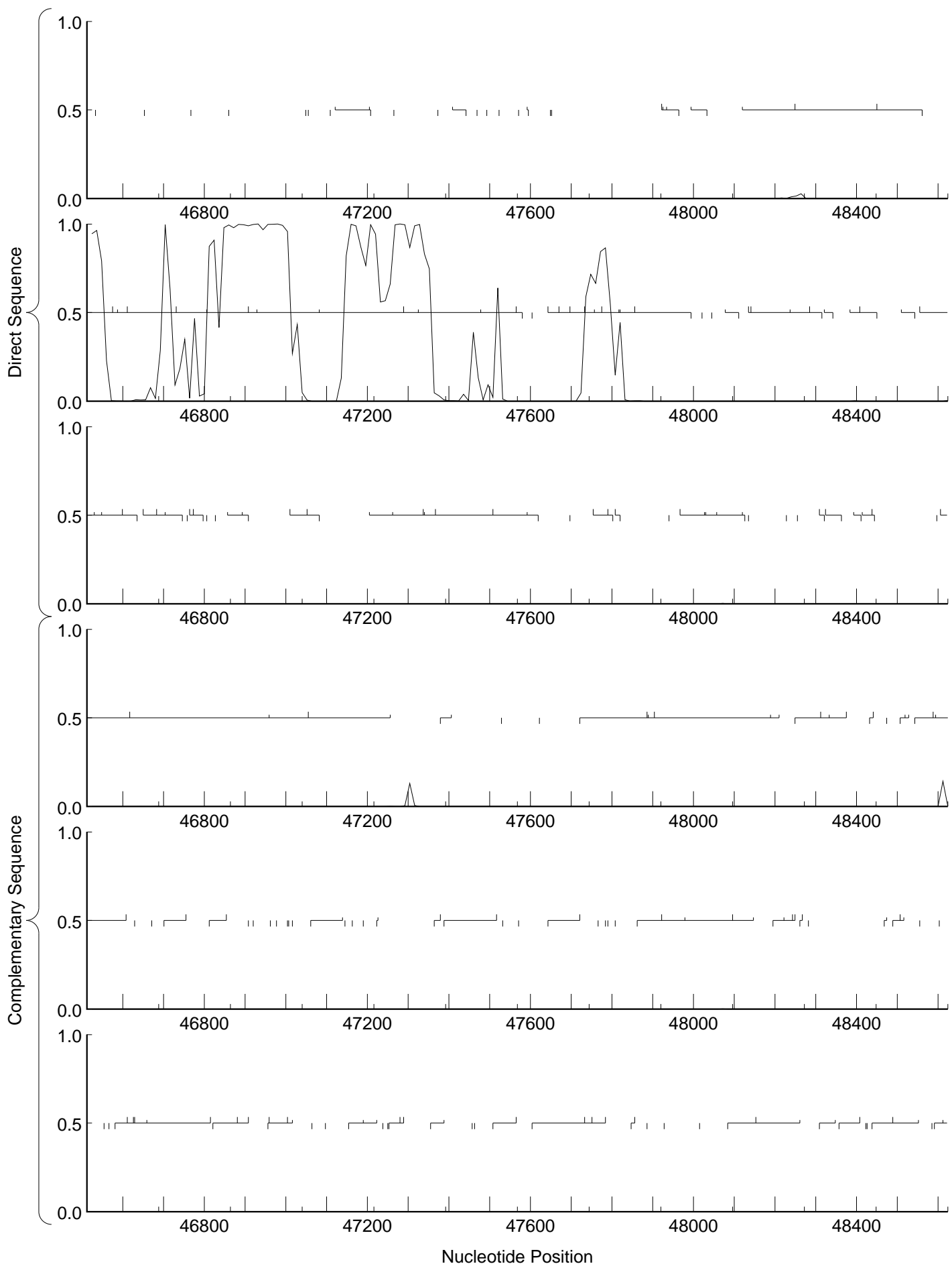












Mossy complete sequence, 61183 including 9-base 3' overhang (CGCCGCCCT), Cluster DJ, Order 4, Window 96, Step 12, 25/30

