

GeneMark

Version 2.5p (09.08.06)

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PROGRAM INFORMATION

Sequence : Streptomyces phage Hippo complete sequence, 51566 bp including 11-base 3' overhang (CGGCCAG
Analysis Date : 4/15/21 at 13:53:38
Pages : 26
Sequence Length : 51566 bp
GC Content : 65.65%

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,

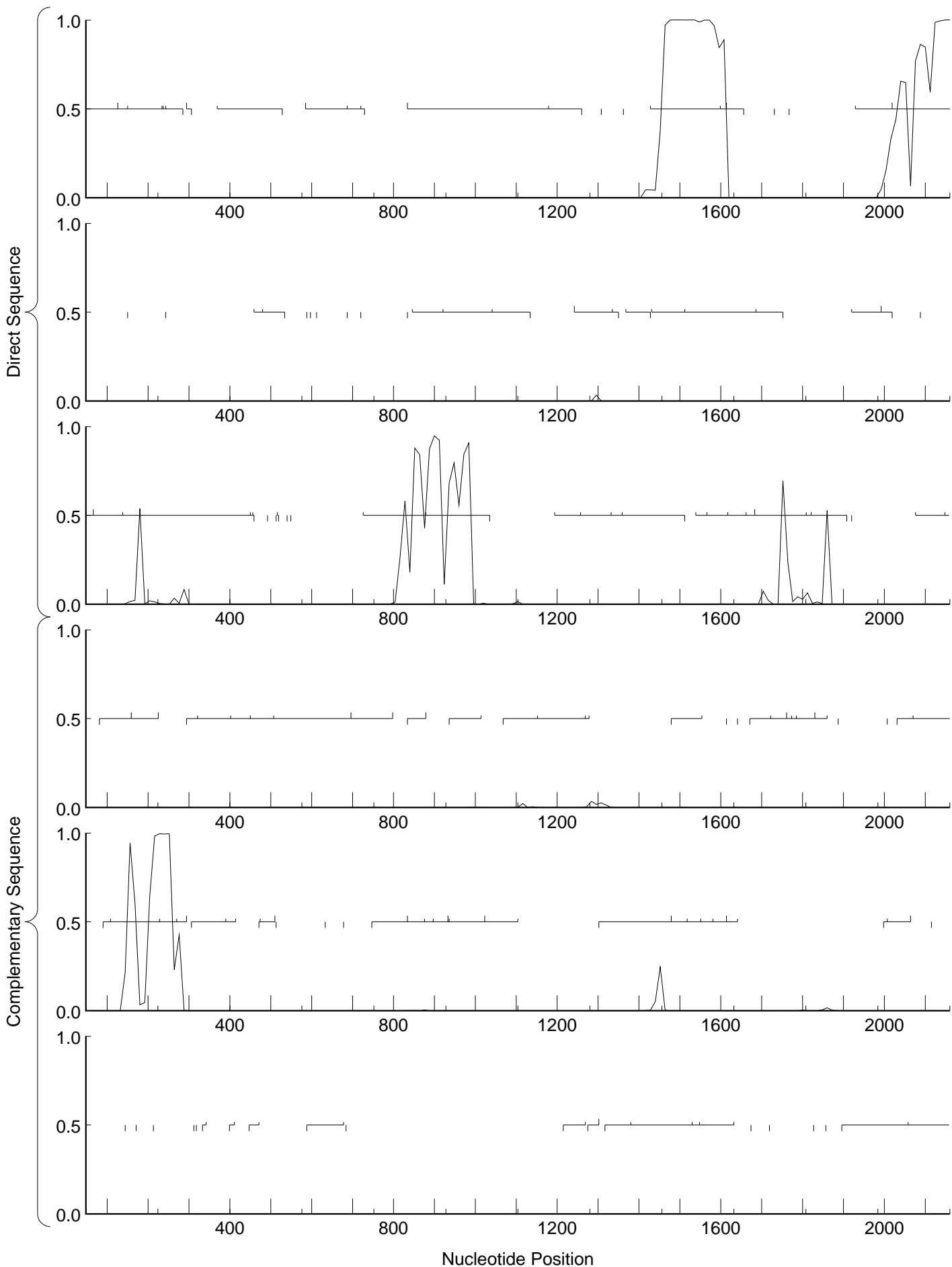
Matrix notes & comments

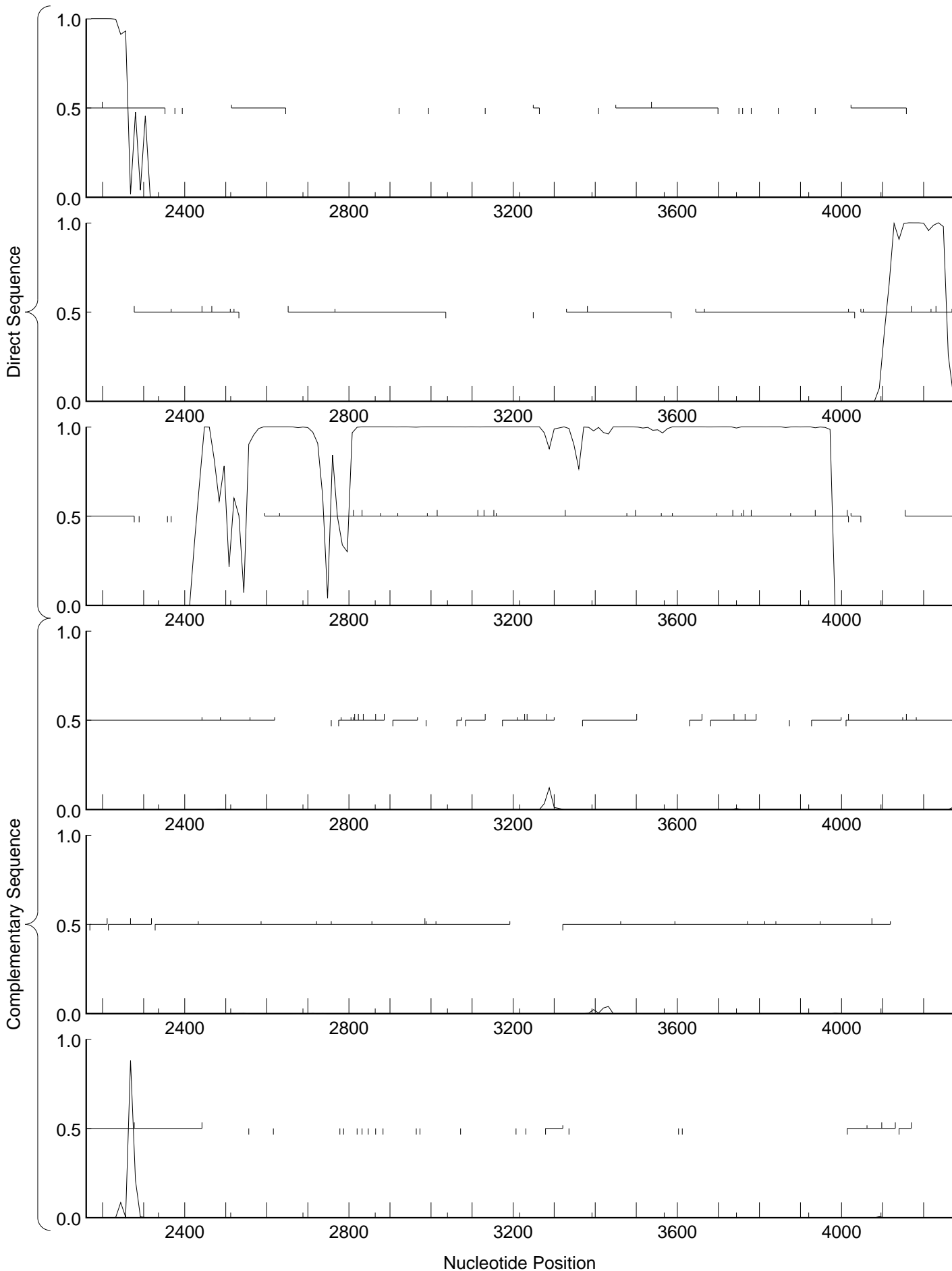
Training set derived by GeneMarkS, 4.27 September 2014
Tue Sep 23 15:43:16 2014

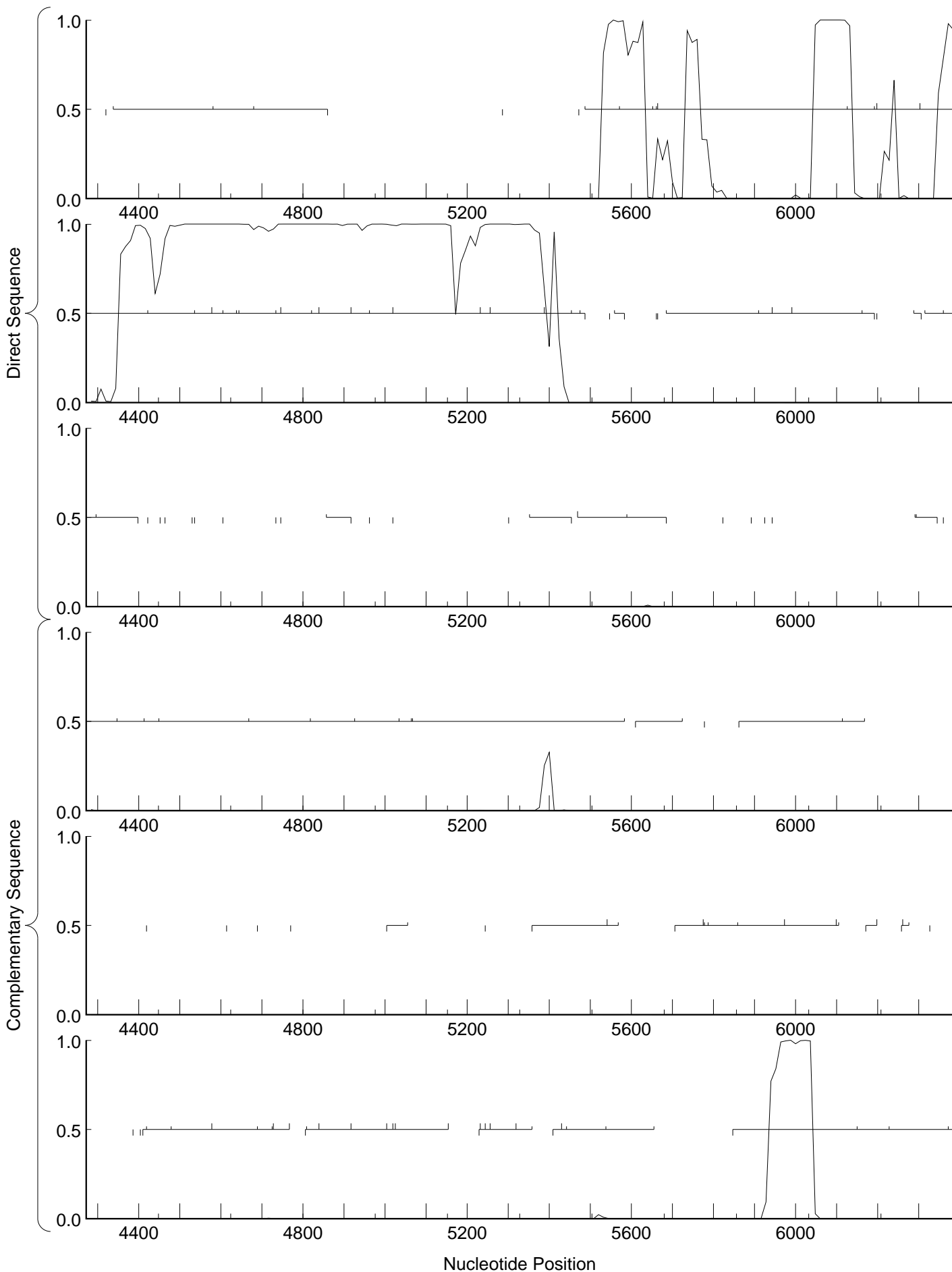
MATRIX INFORMATION

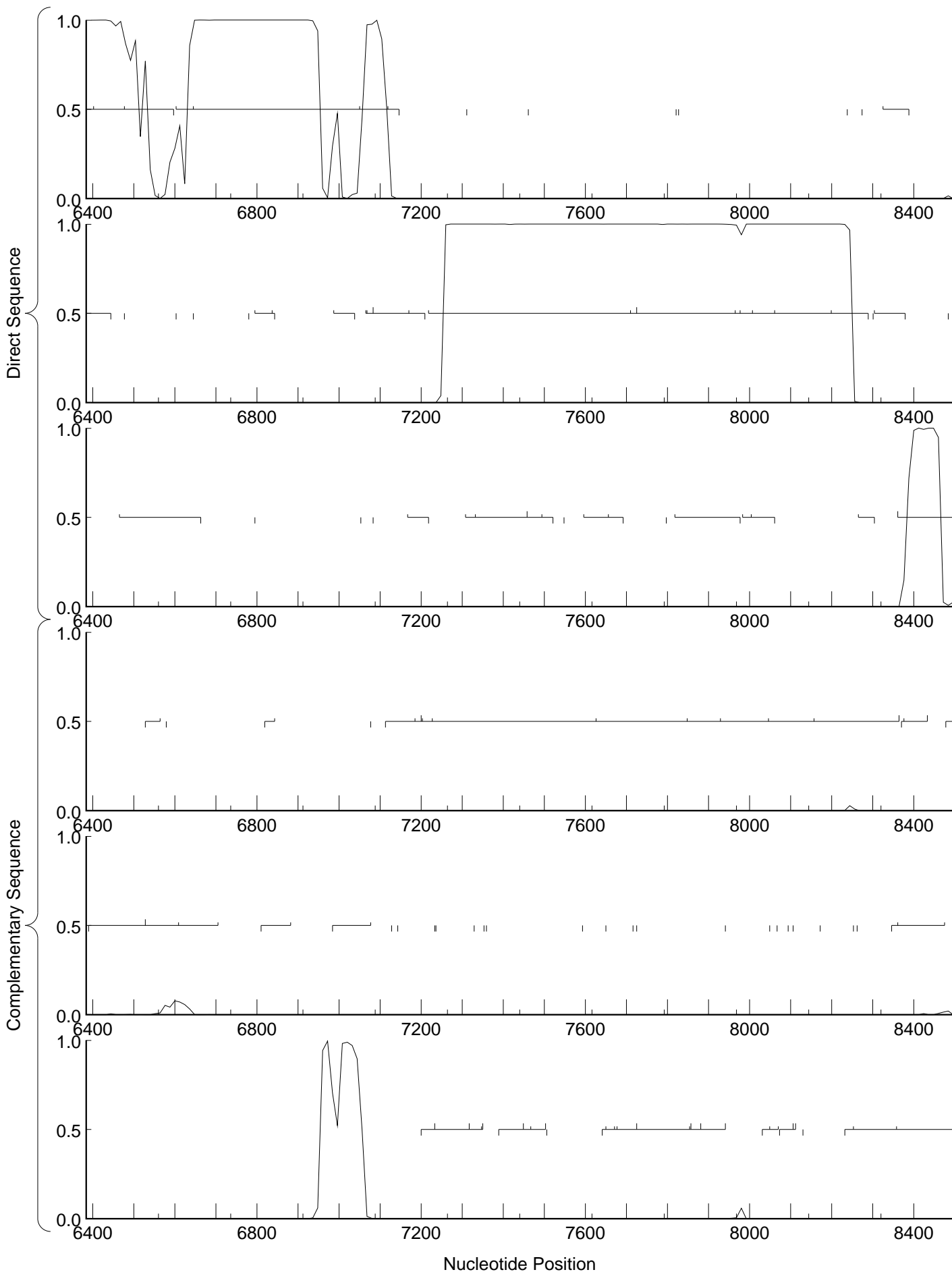
Matrix : Streptomyces_scabiei_87_22
Author : -
Order : 4

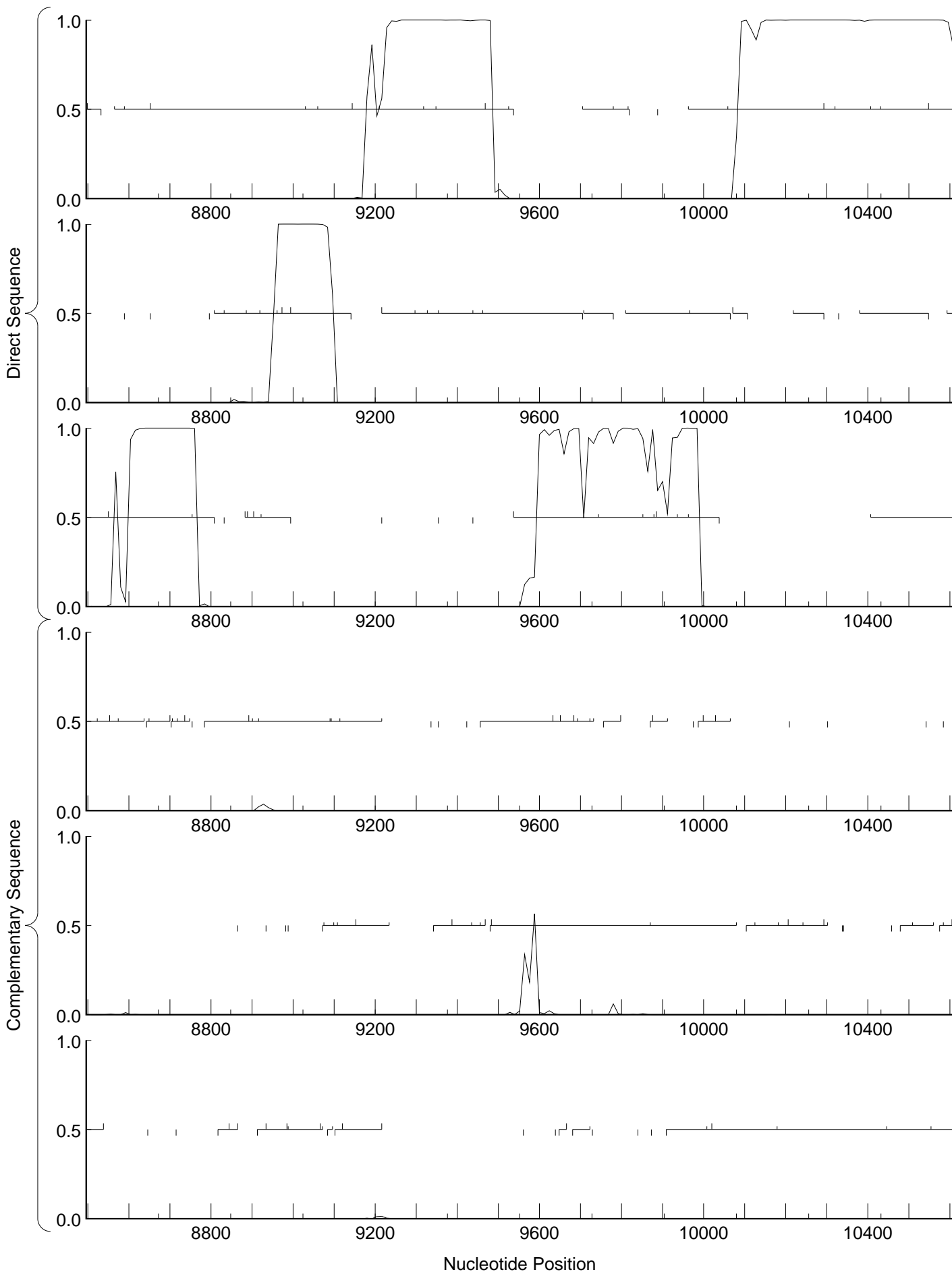
Send questions / comments to:
Dr. M. Borodovsky
Georgia Institute of Technology
School of Biology
Atlanta, GA 30332-0230

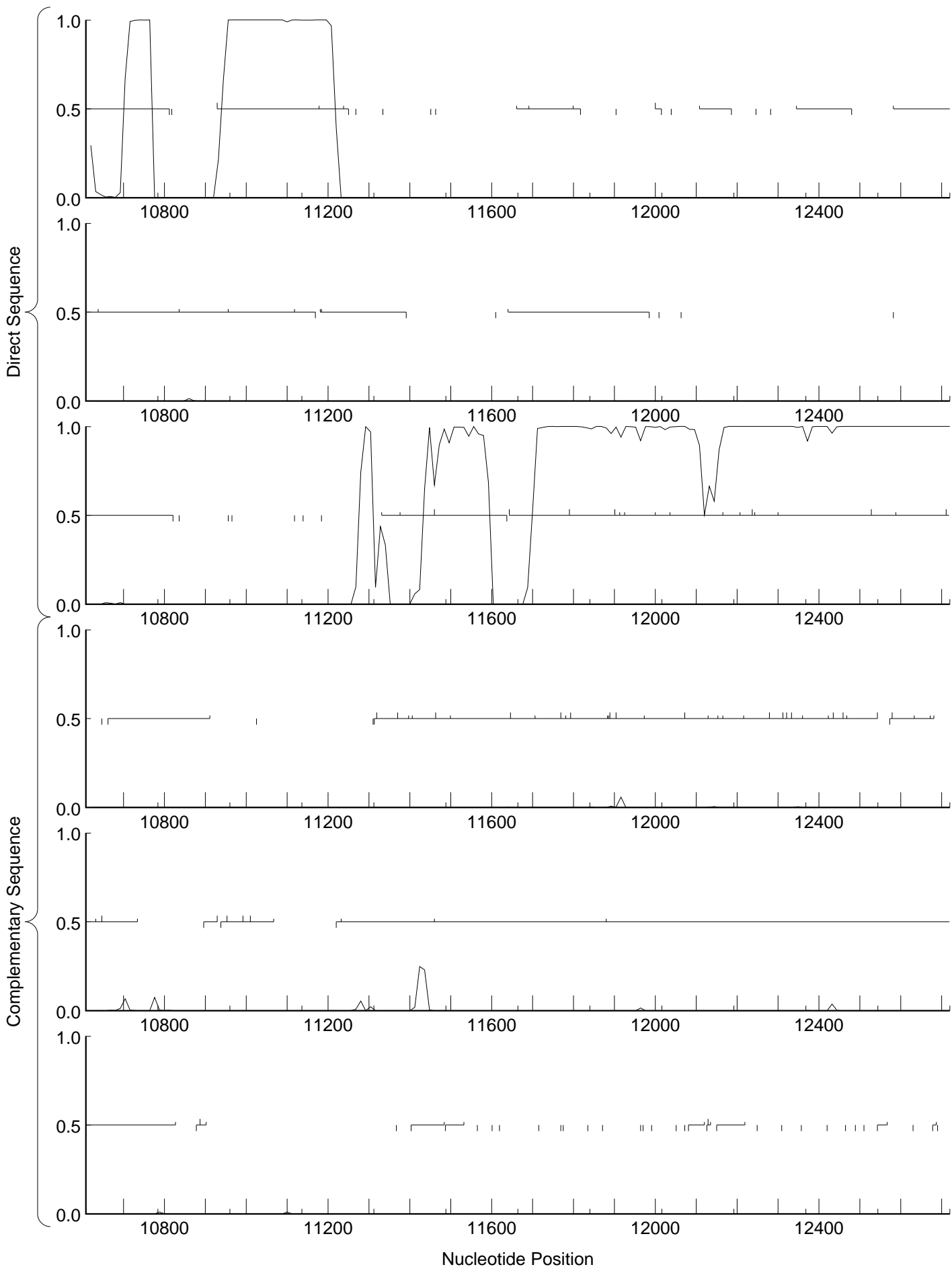


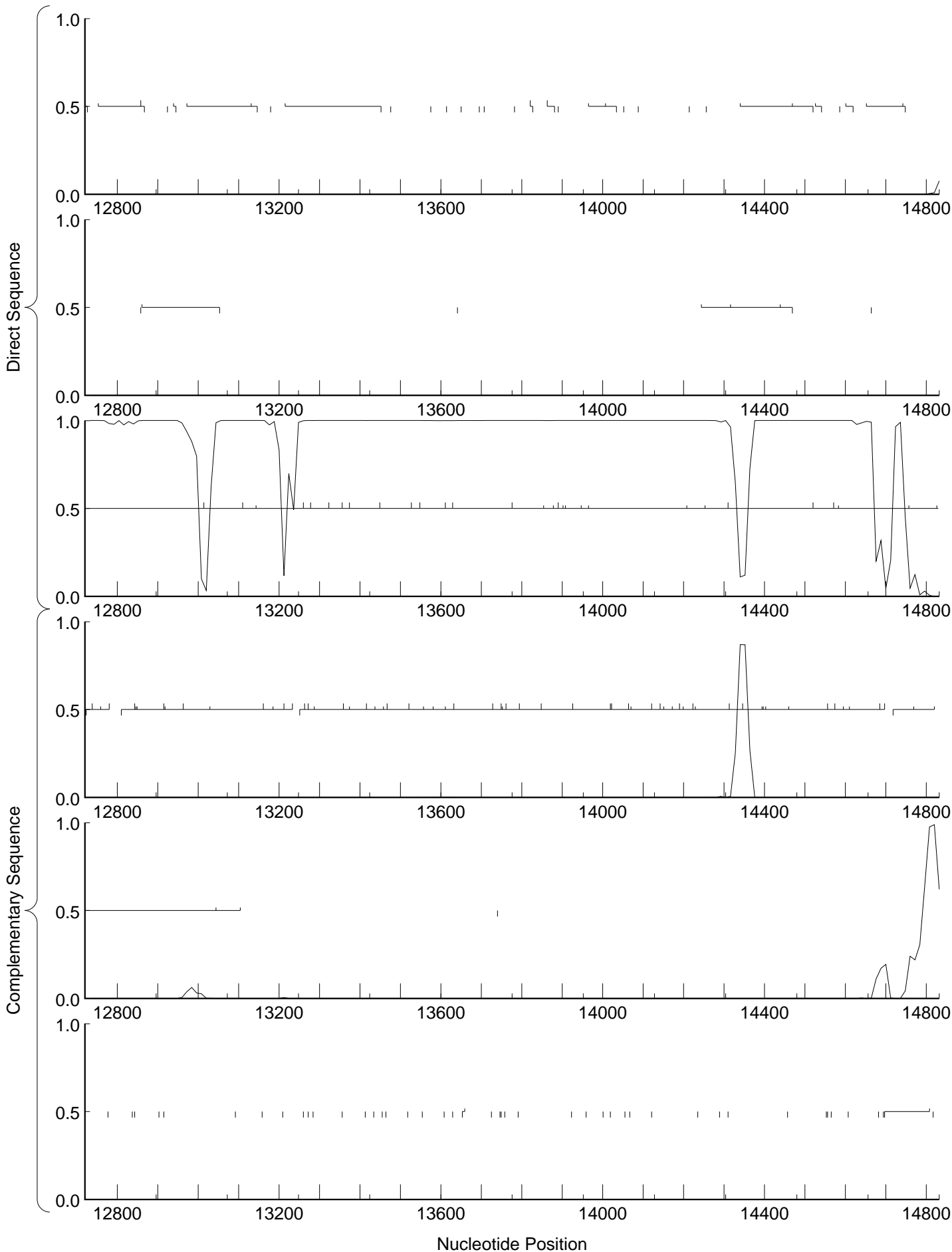


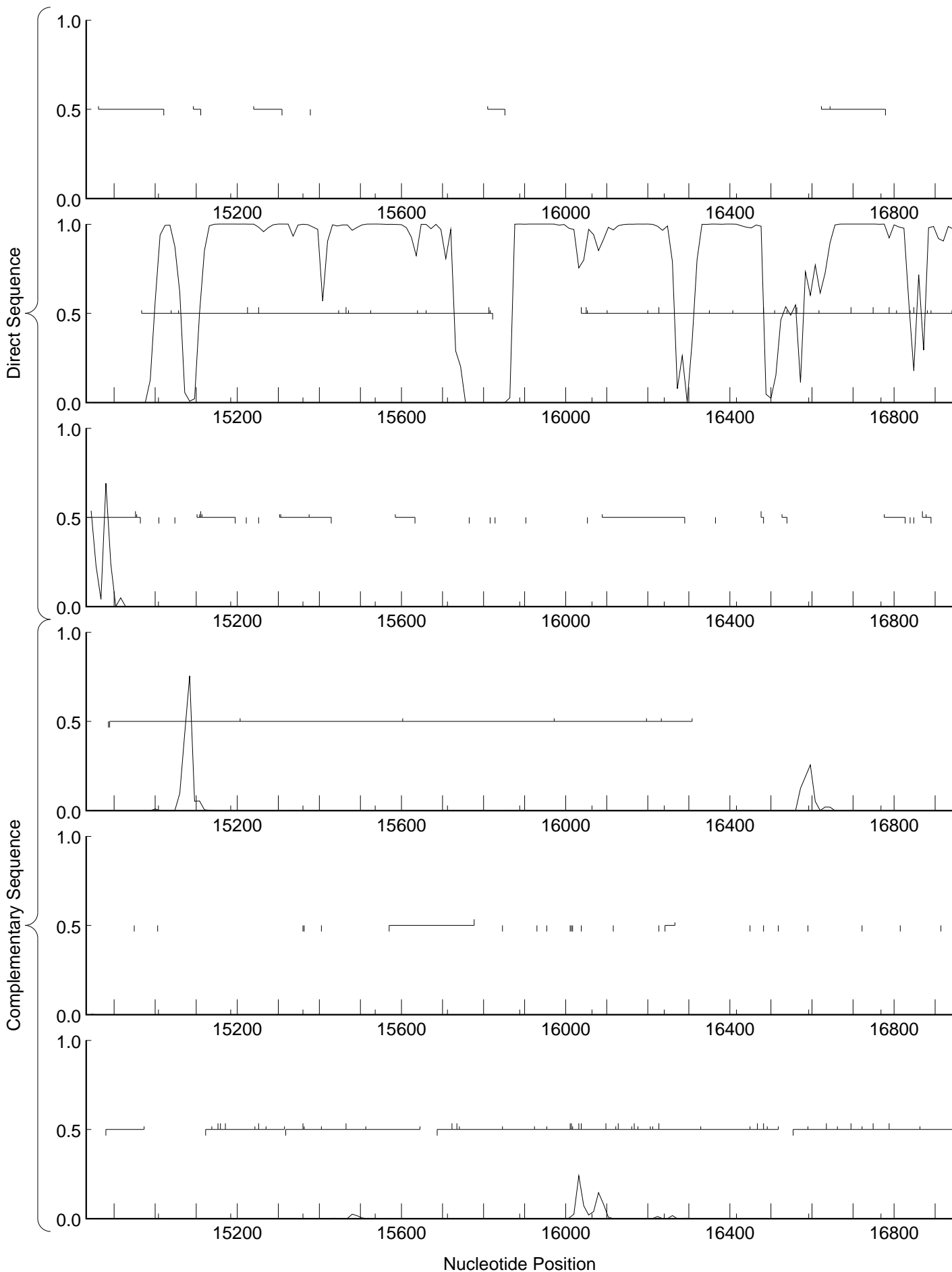


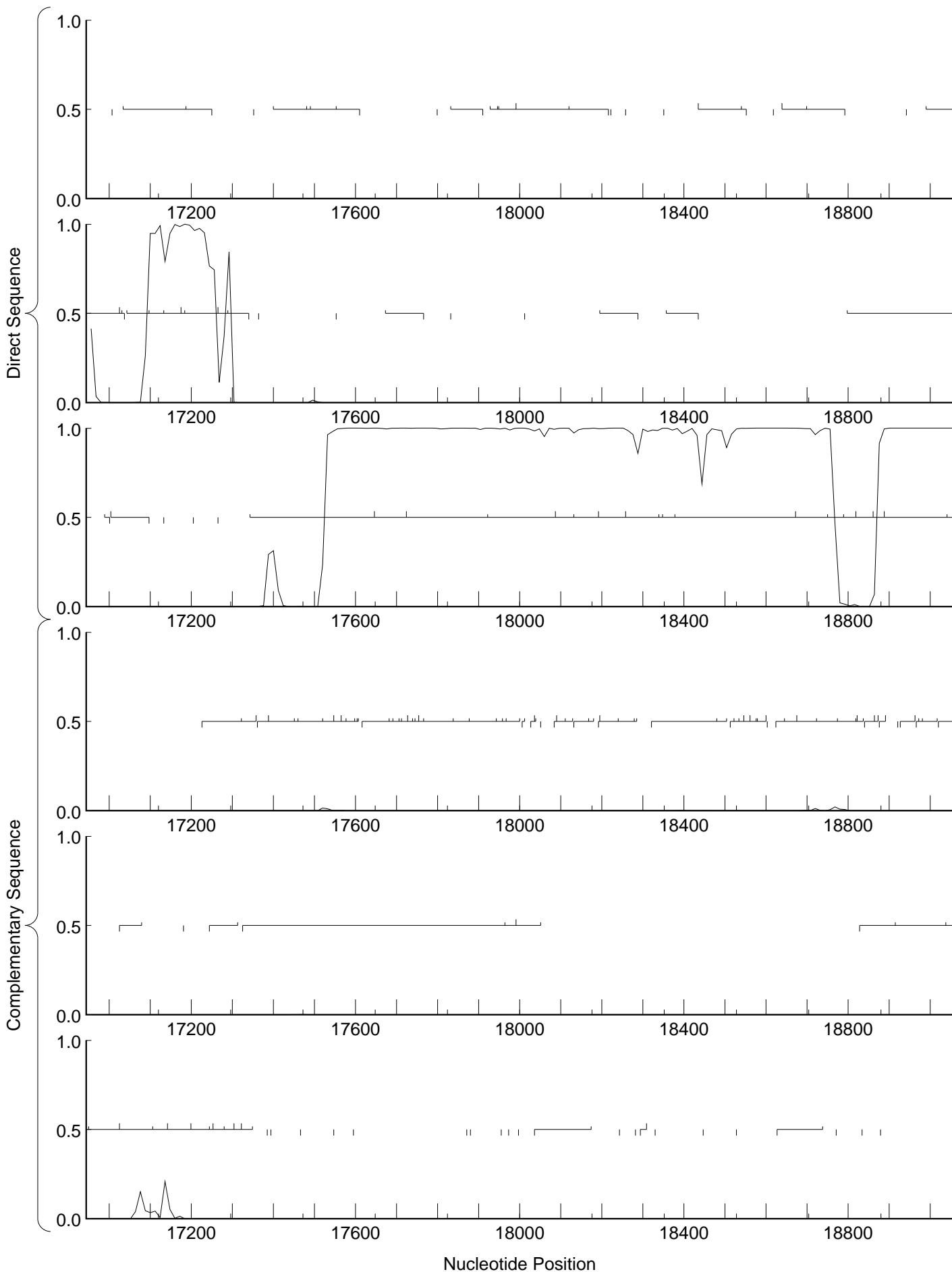


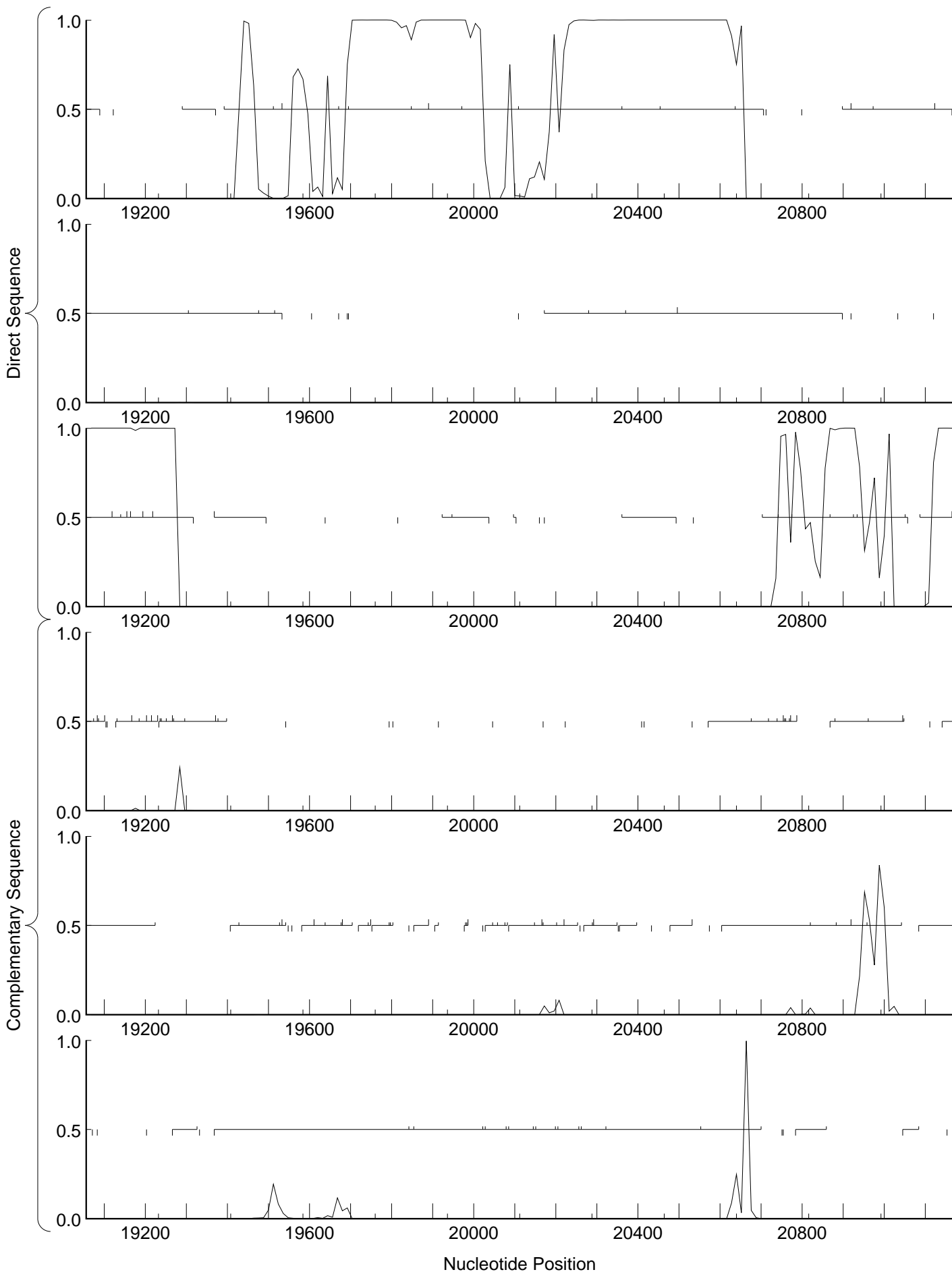


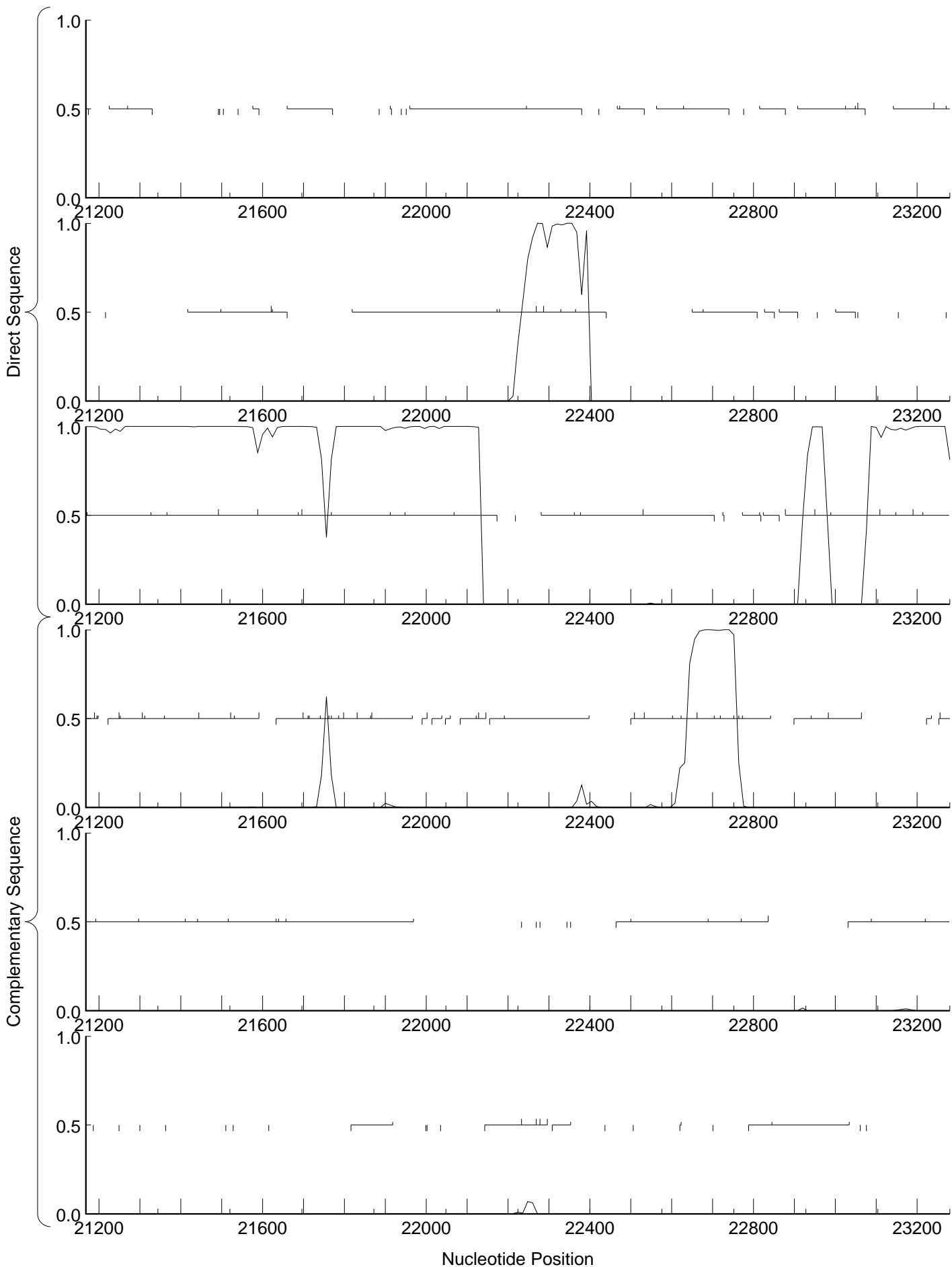


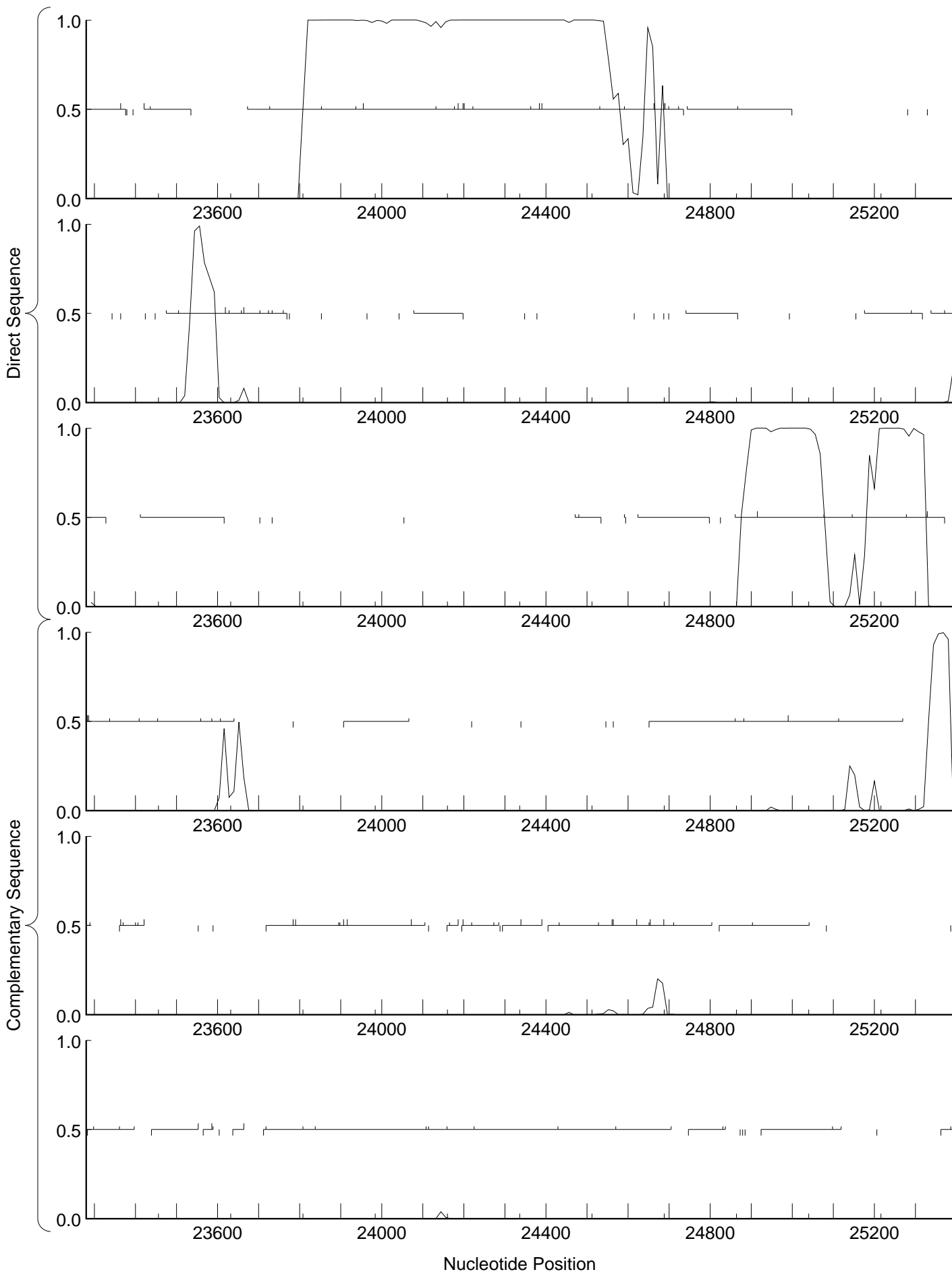


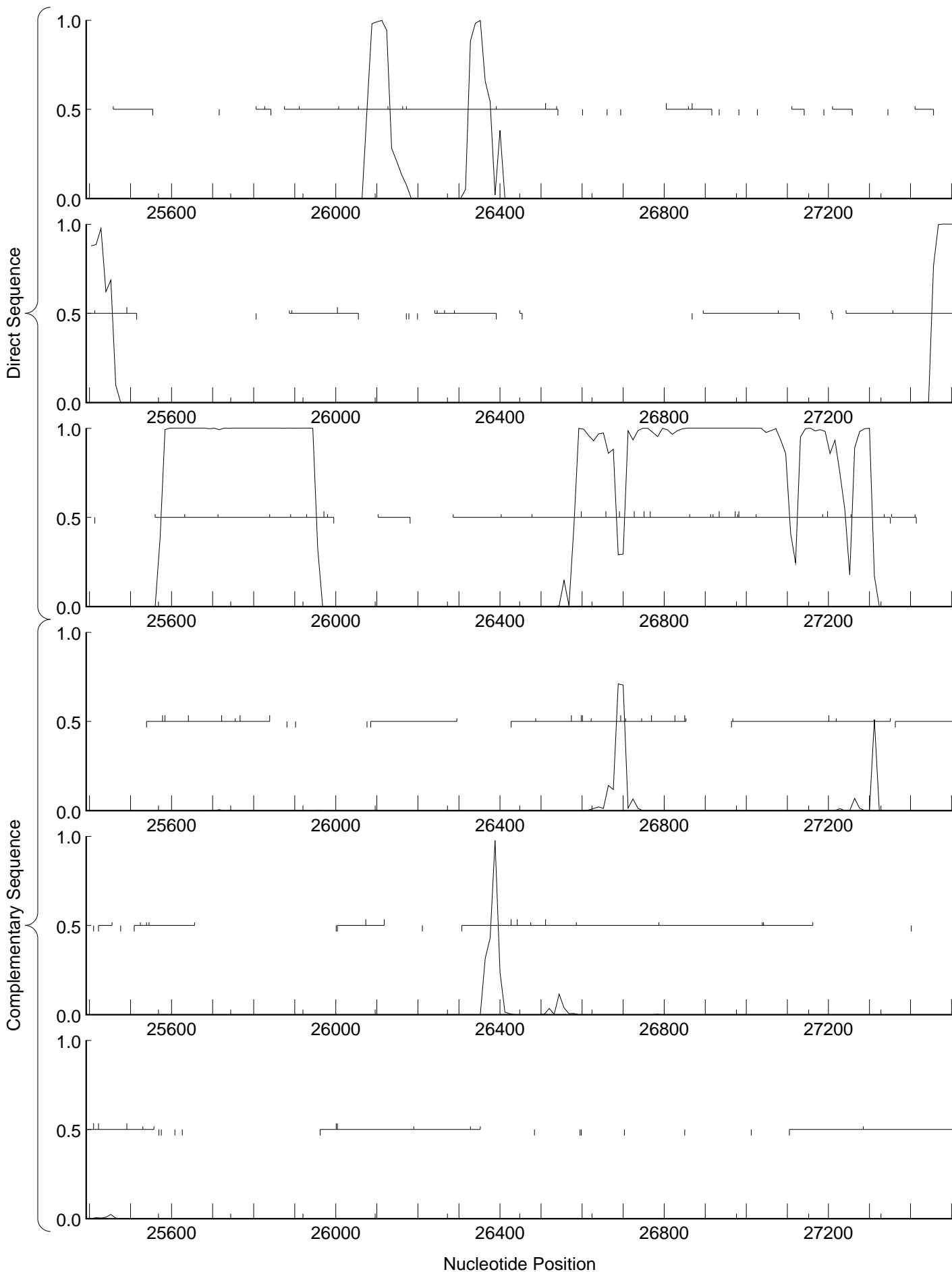


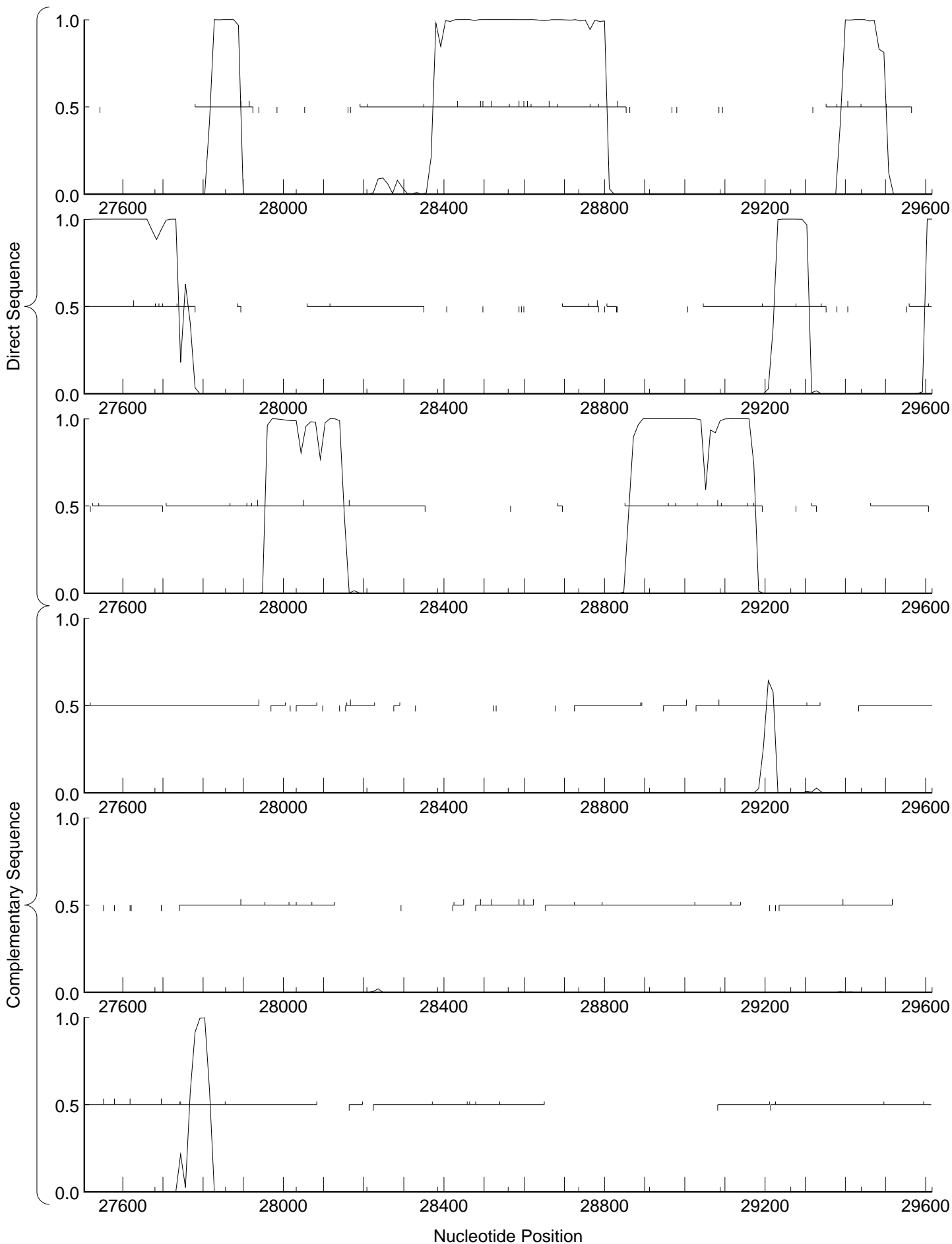


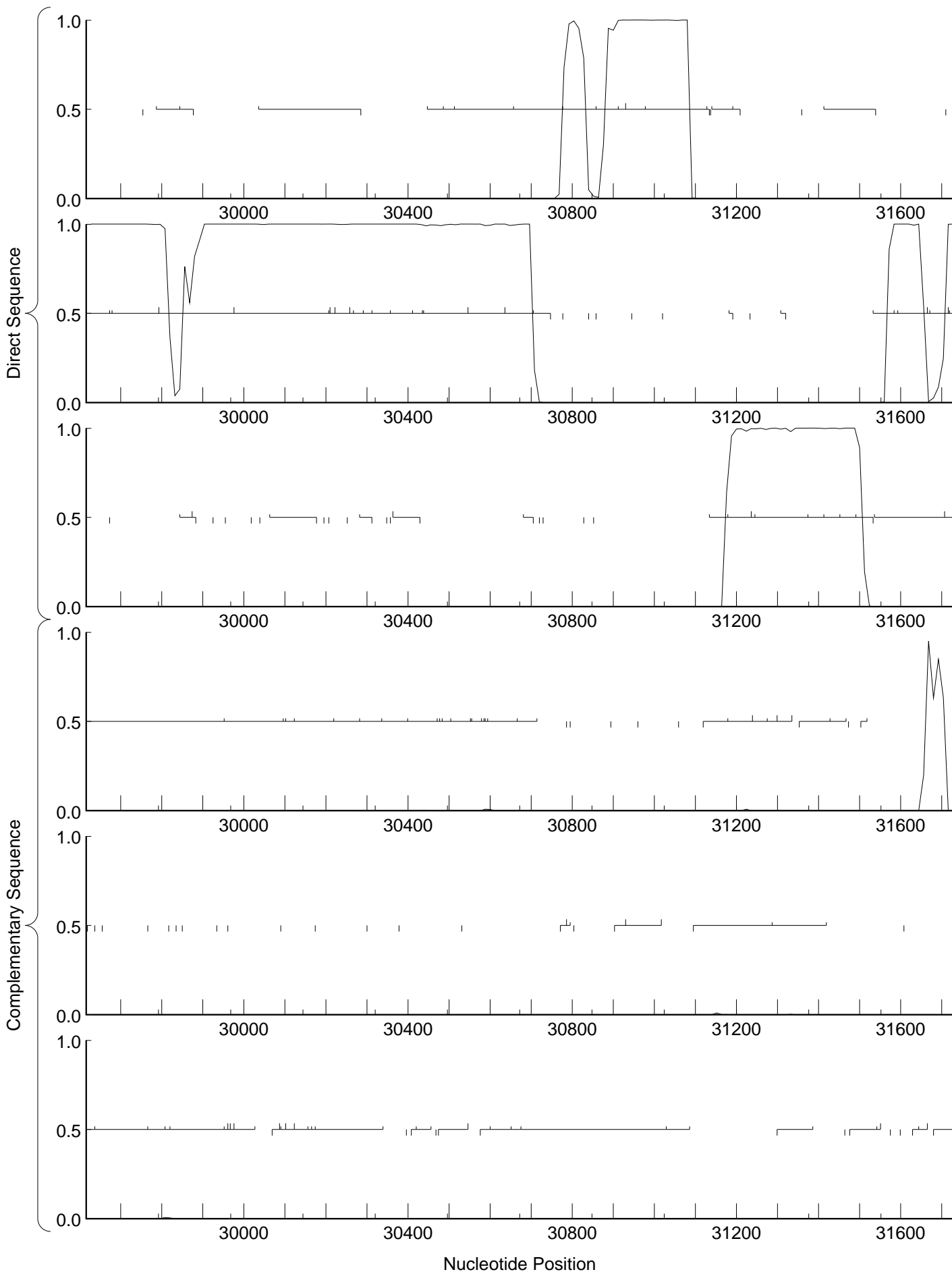












Complete sequence, 51566 bp including 11-base 3' overhang (CGGCCAGTGAT), Cluster BD1, Order 4, Window 96, Step 12, 17/26

