

GeneMark

Version 2.5p (09.08.06)

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PROGRAM INFORMATION

Sequence : Microbacterium phage Stoor complete sequence, 41814 bp including 10-base 3' overhang (TCTCC
Analysis Date : 6/24/24 at 18:28:12
Pages : 21
Sequence Length : 41814 bp
GC Content : 69.11%

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,

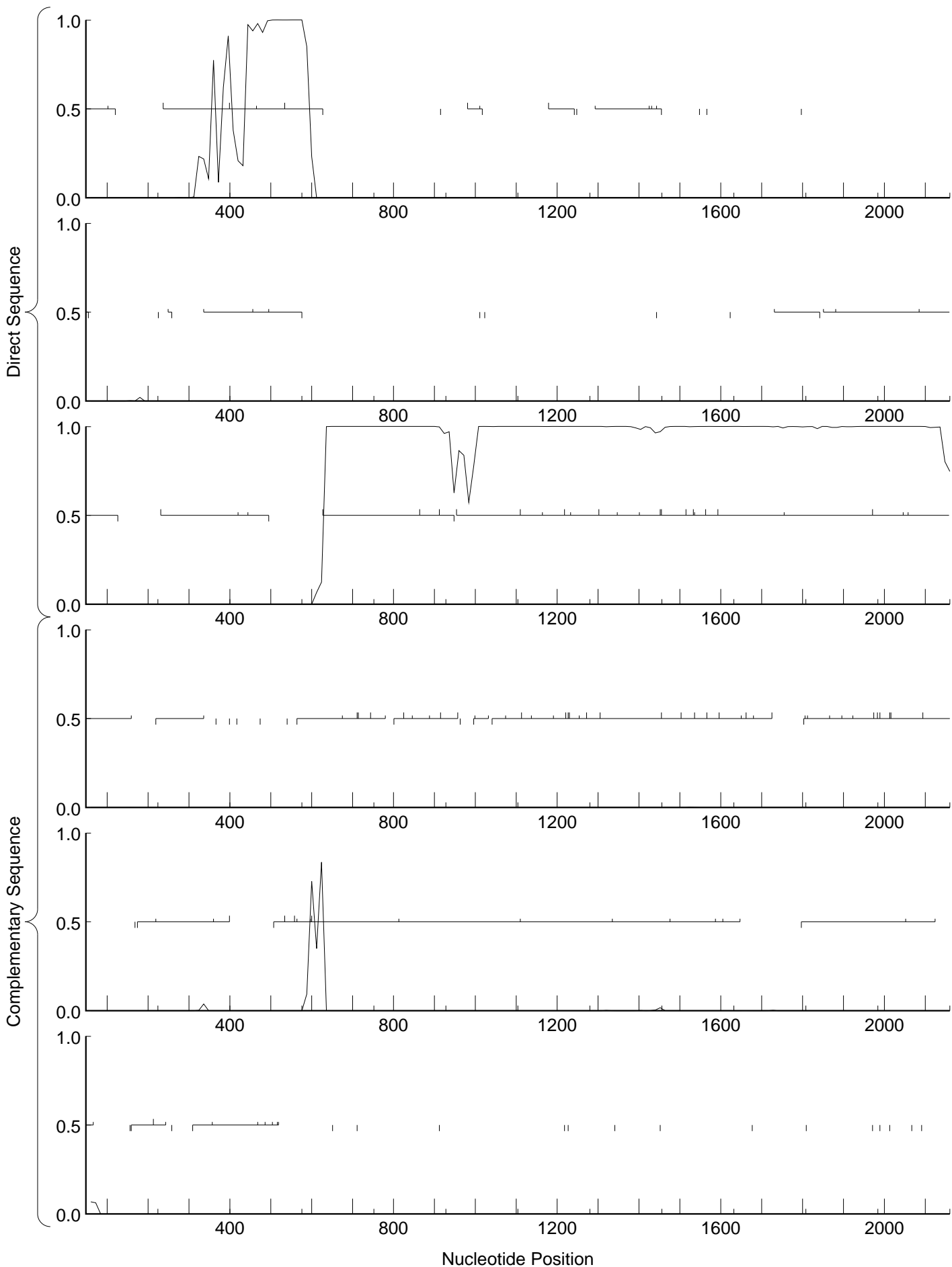
Matrix notes & comments

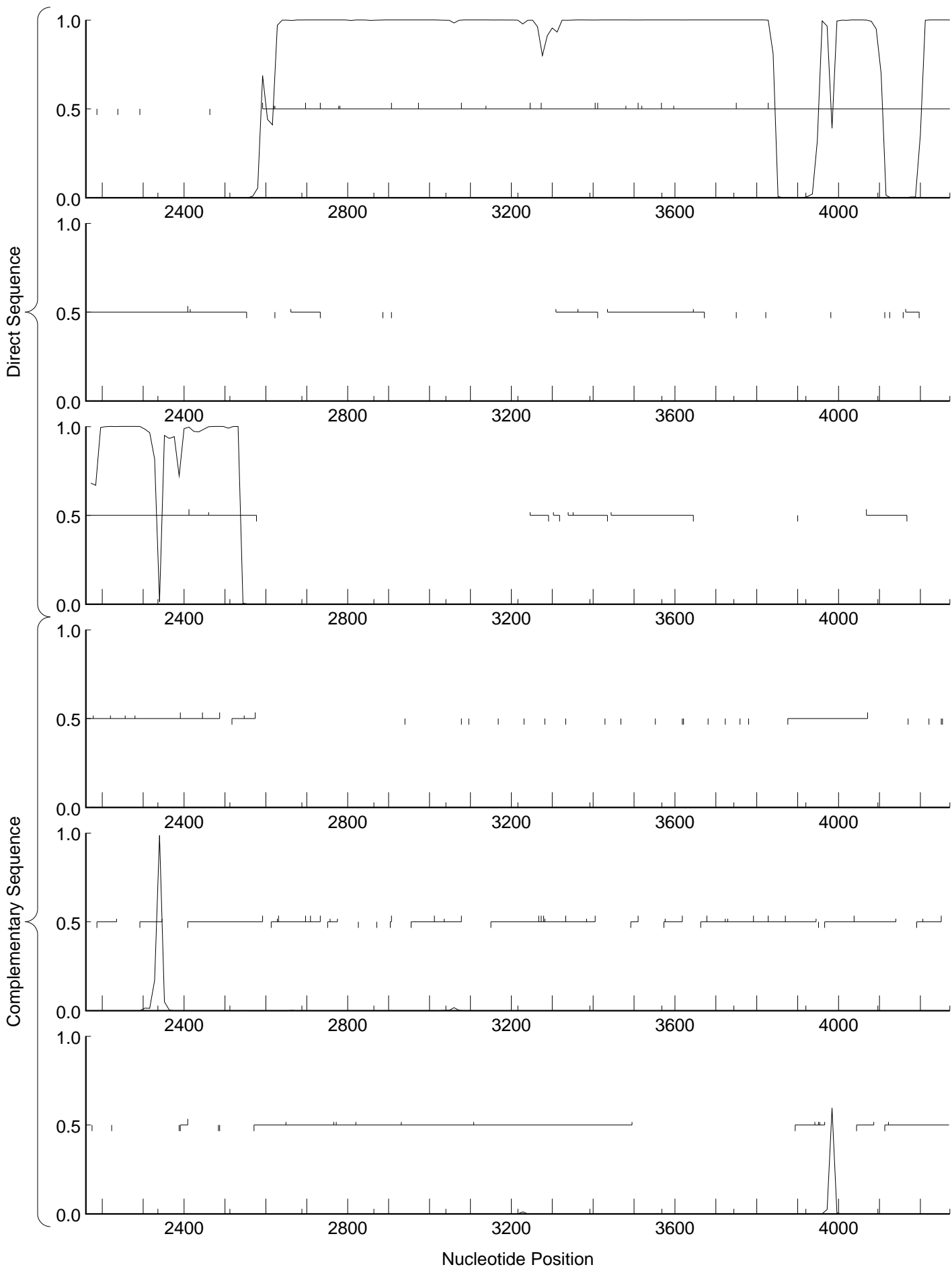
Training set derived by GeneMarkS, 4.30 October 2014
Fri Sep 7 18:42:59 2018

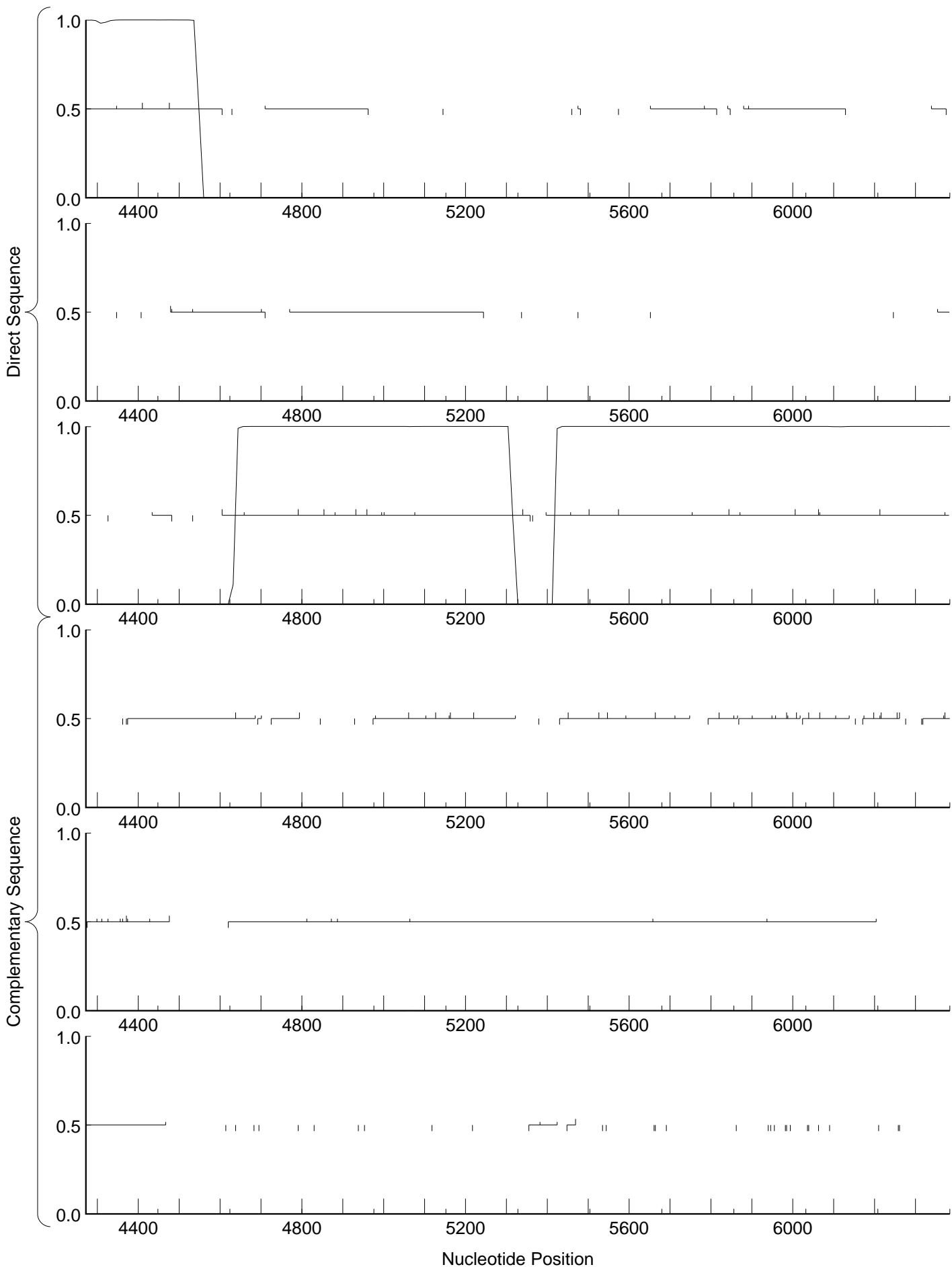
MATRIX INFORMATION

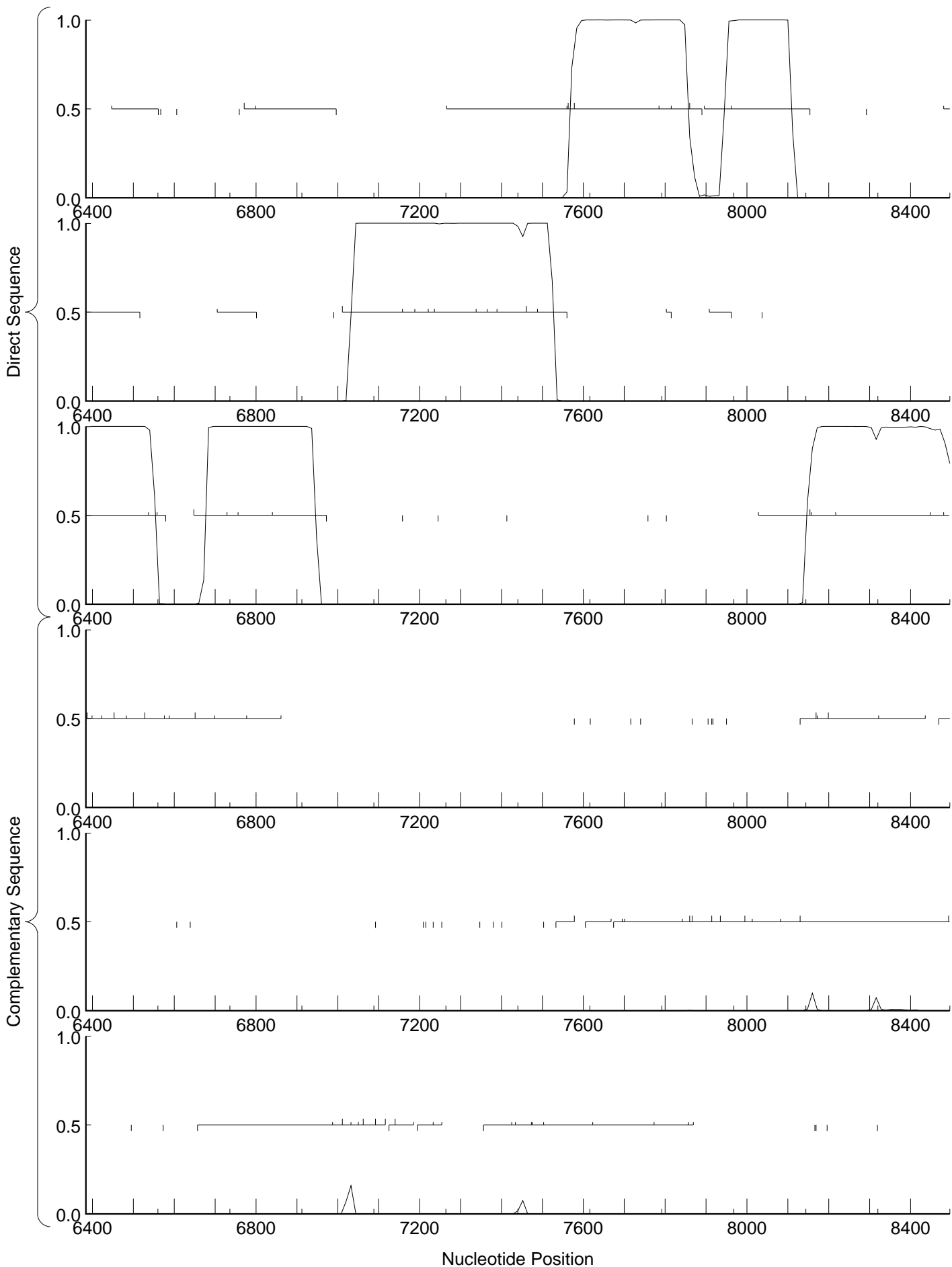
Matrix : -
Author : -
Order : 4

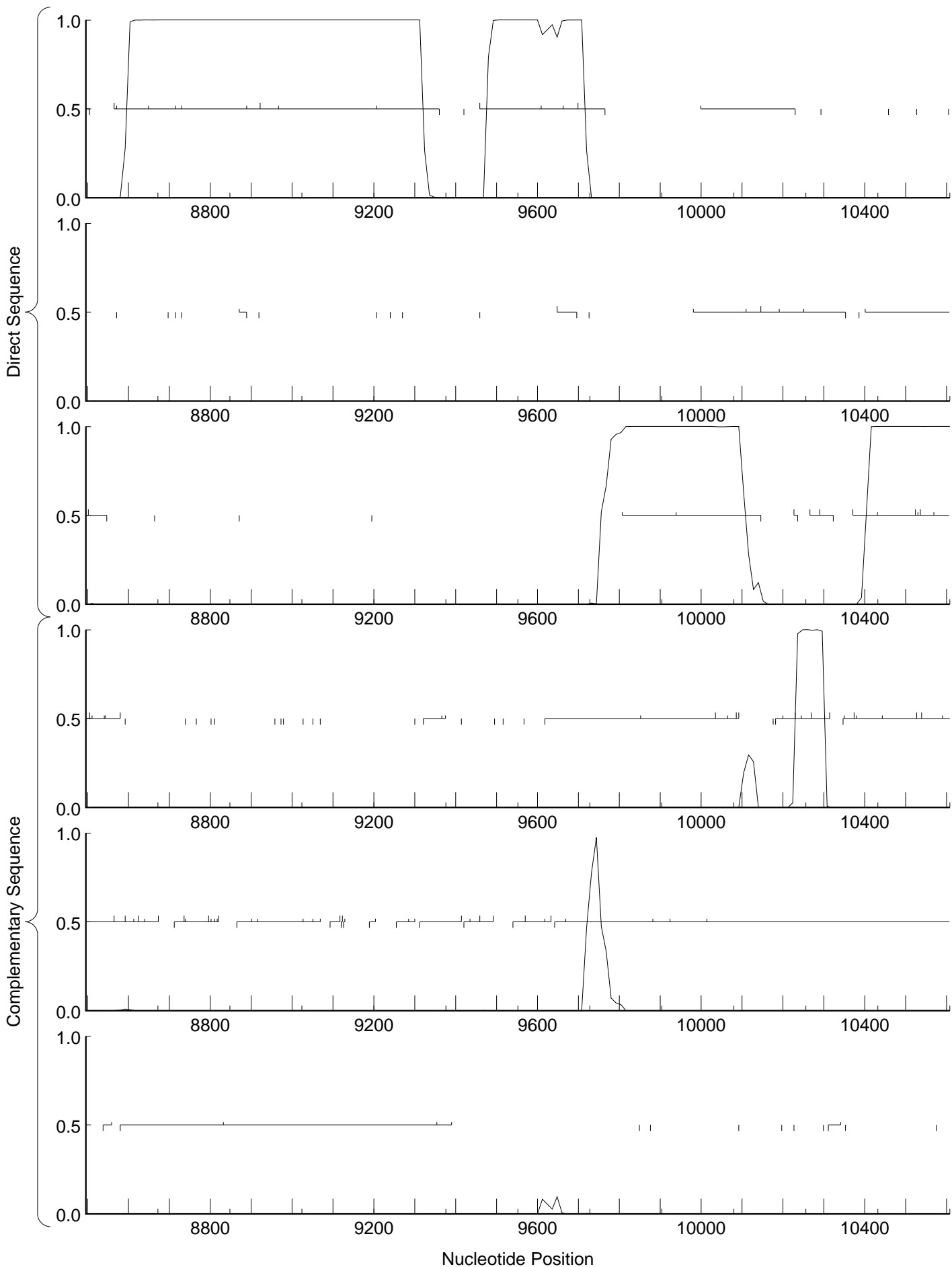
Send questions / comments to:
Dr. M. Borodovsky
Georgia Institute of Technology
School of Biology
Atlanta, GA 30332-0230

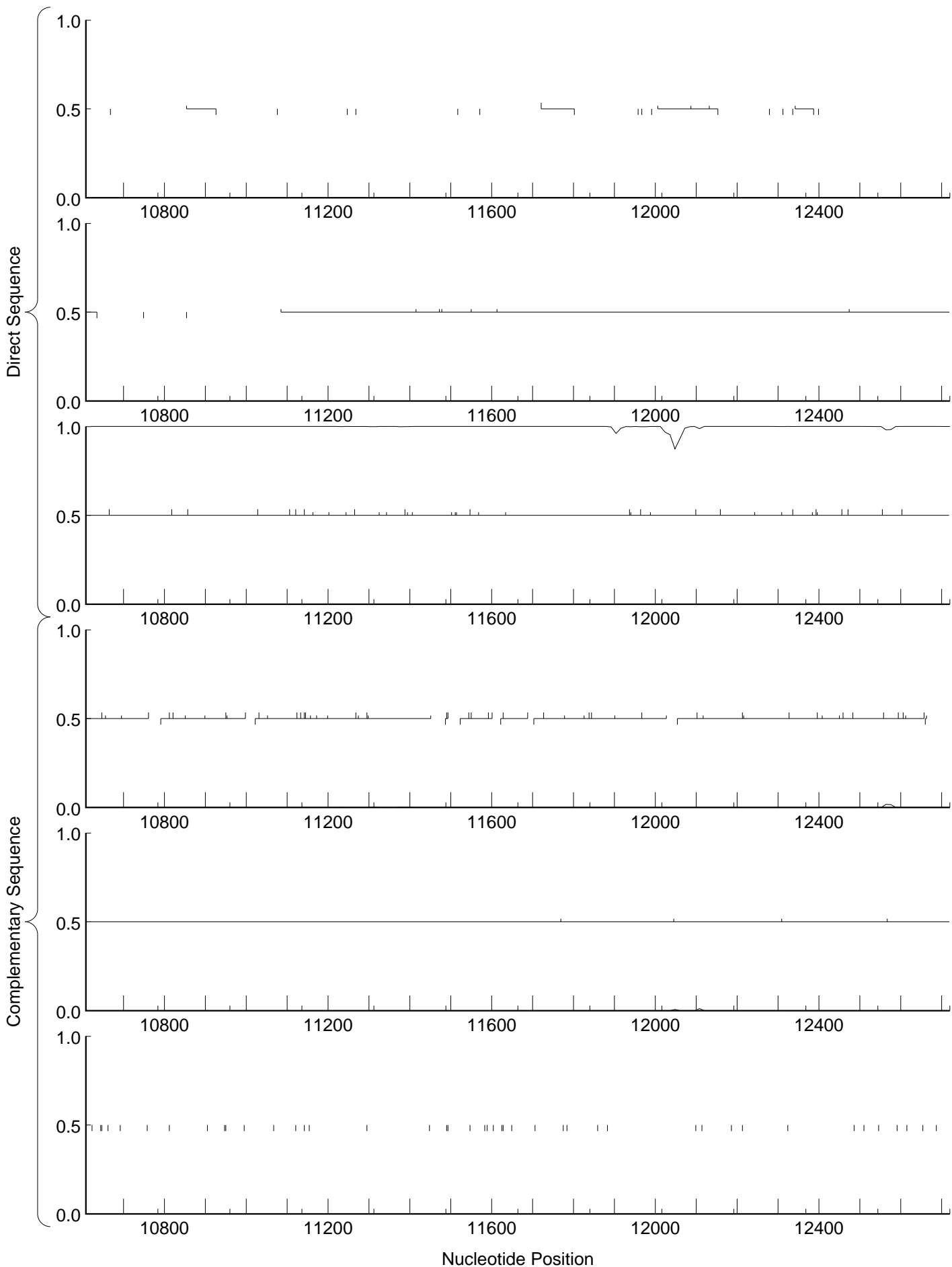


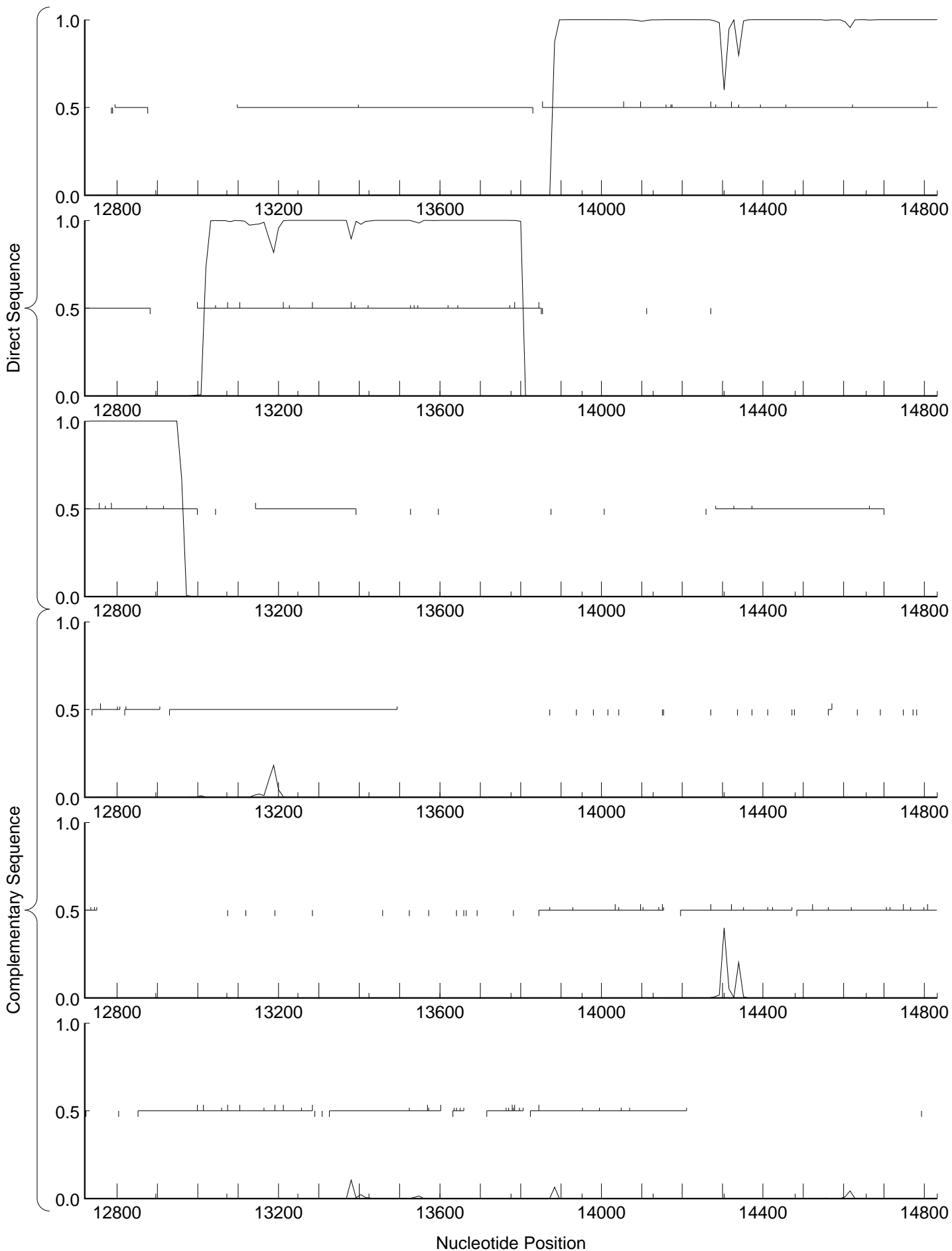


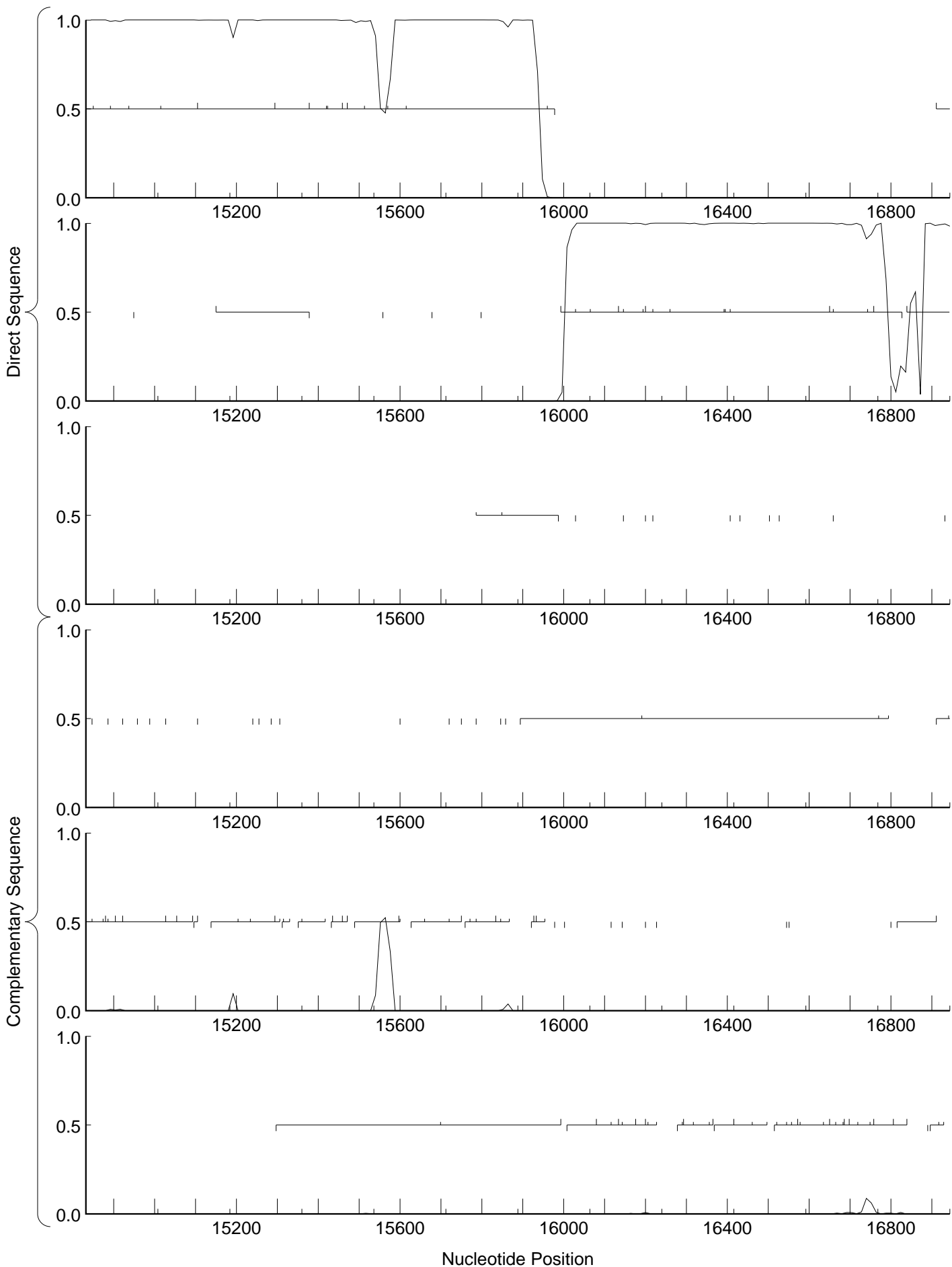


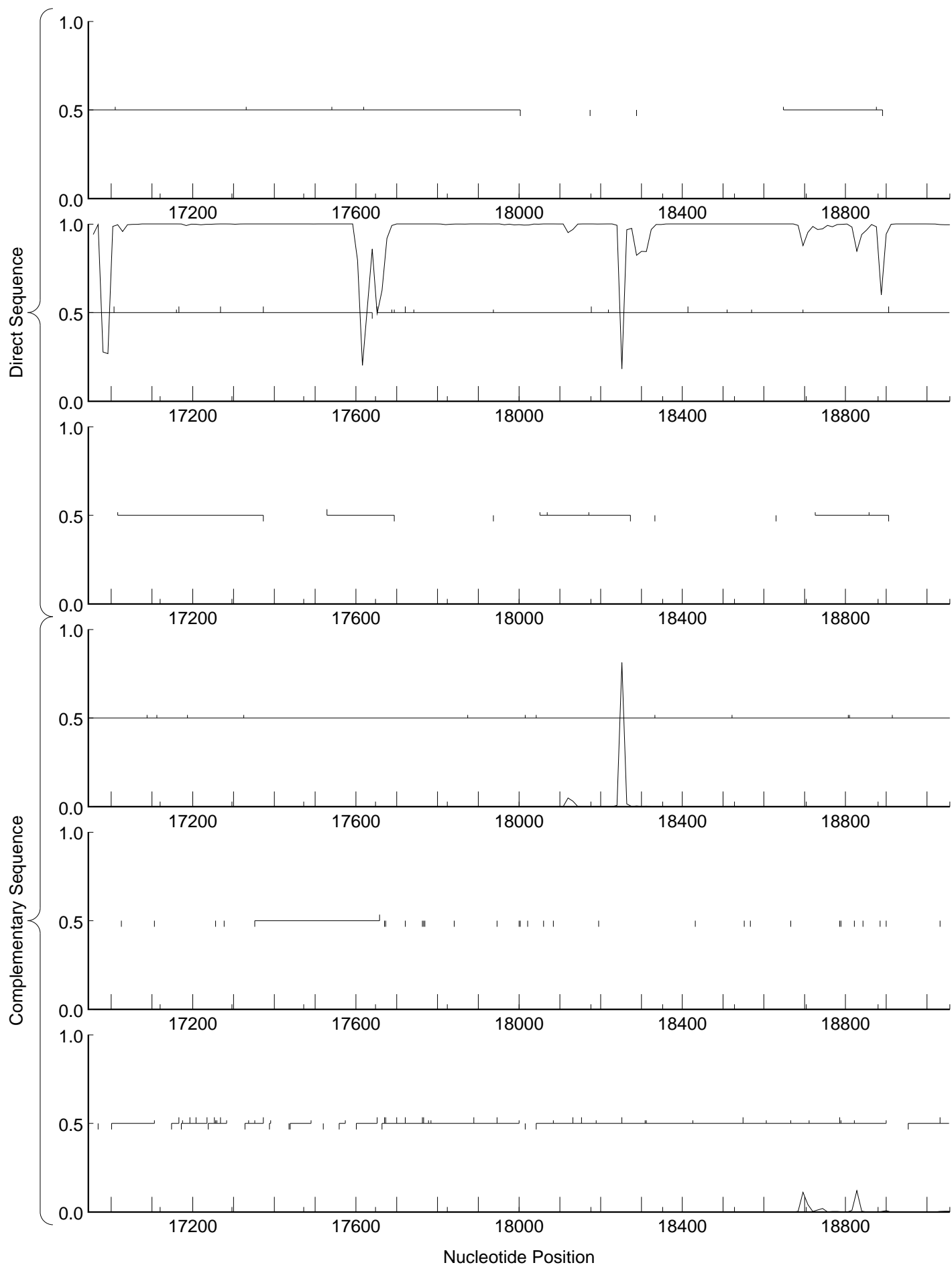




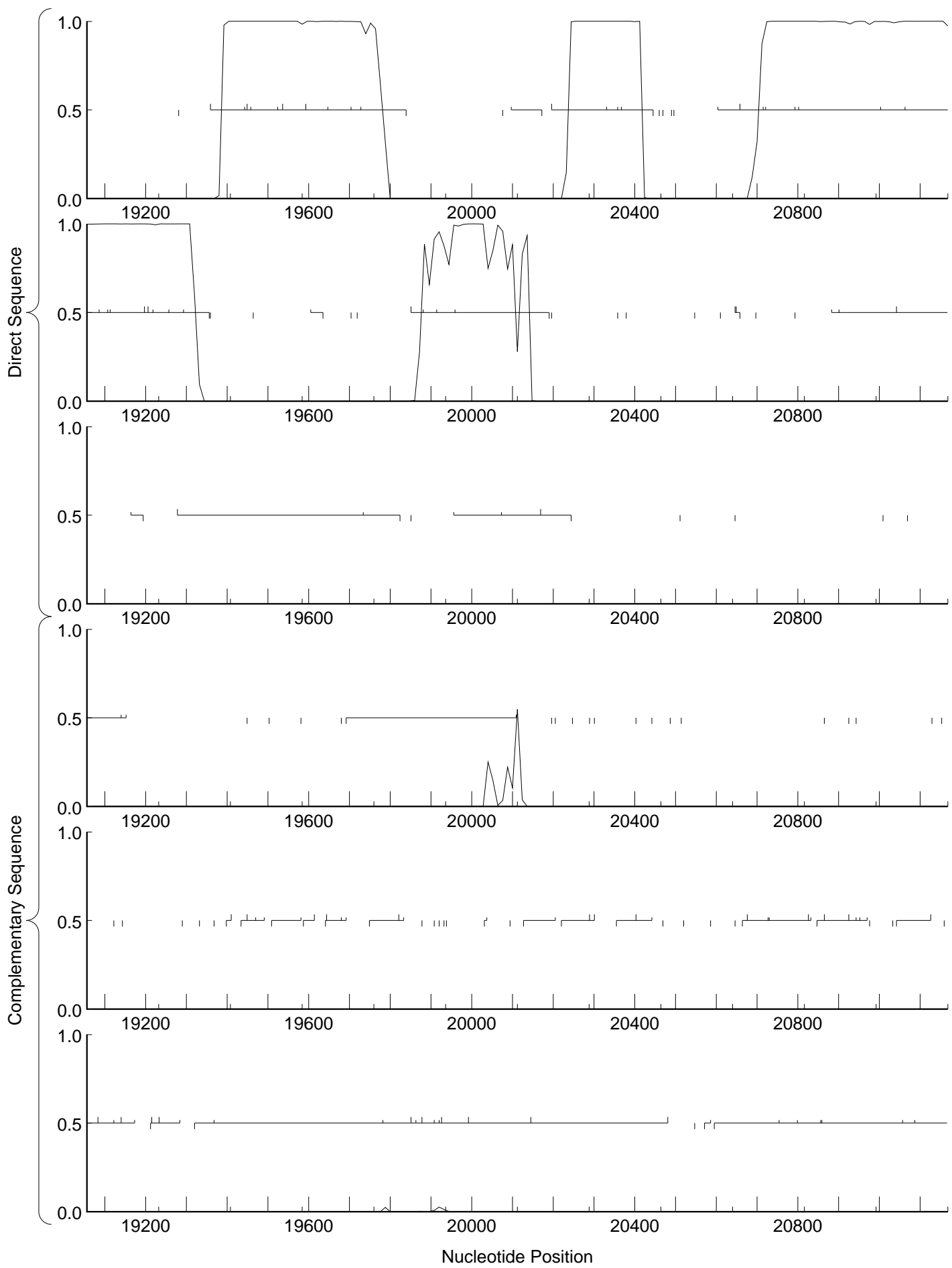


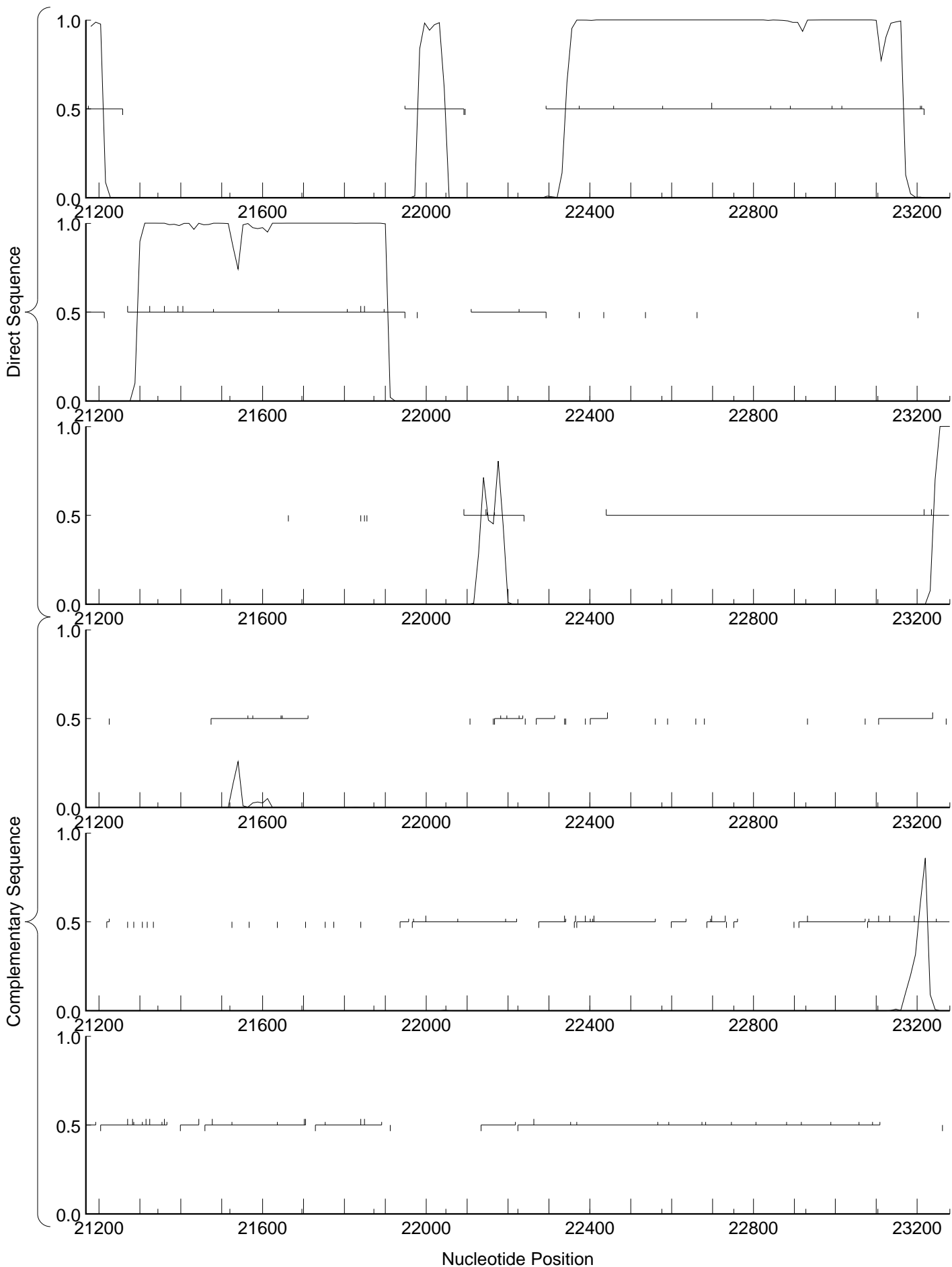


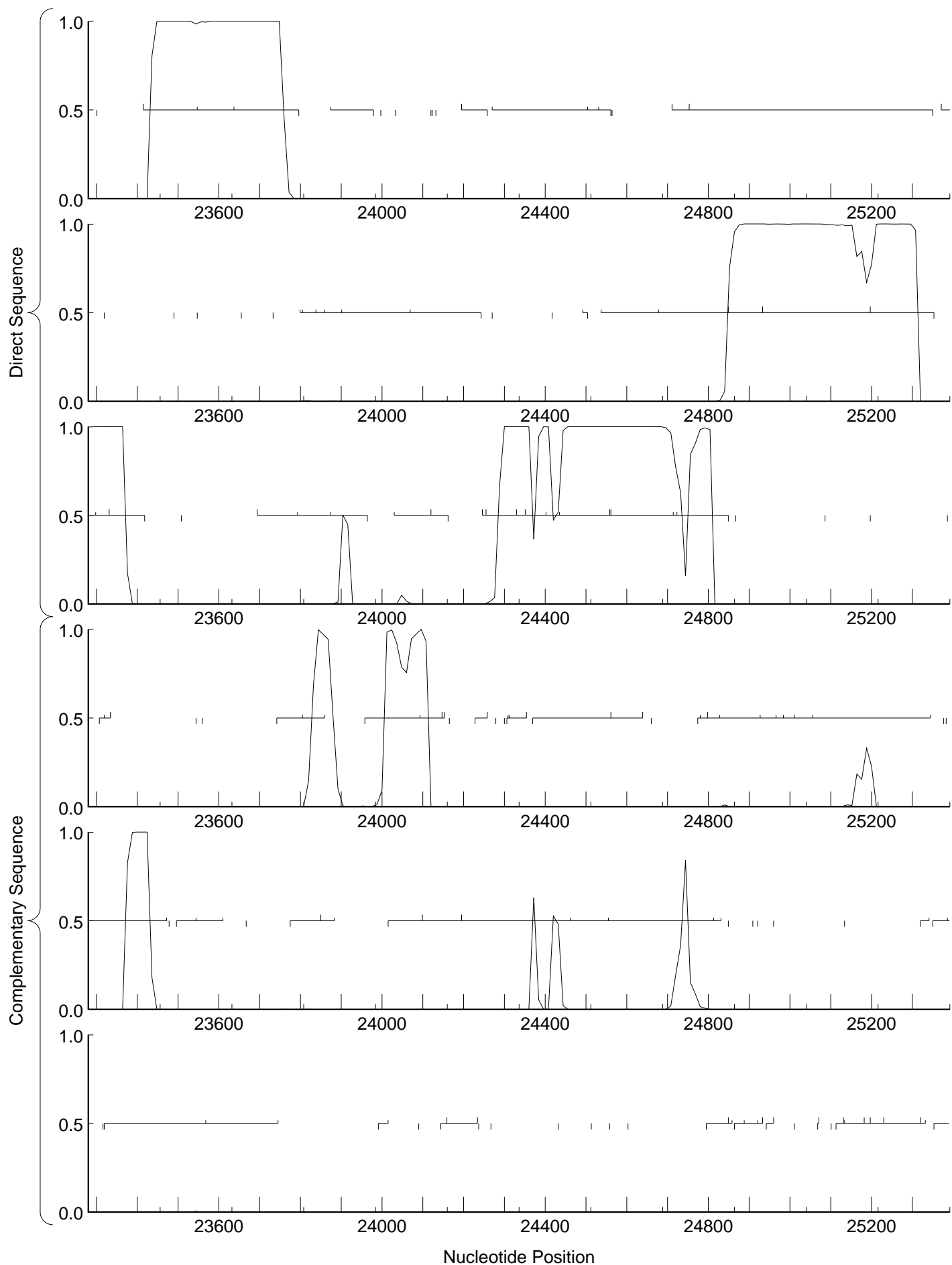


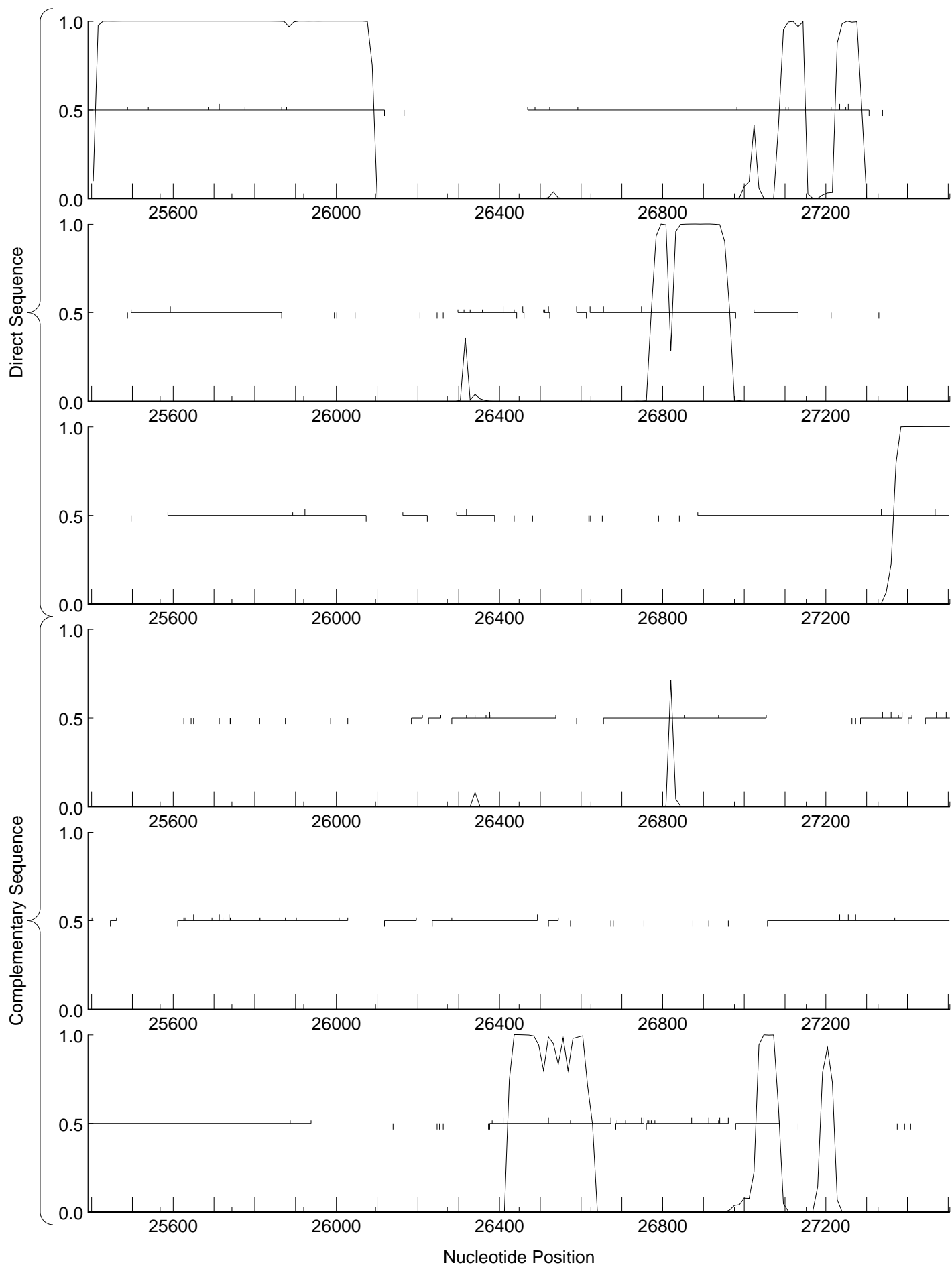


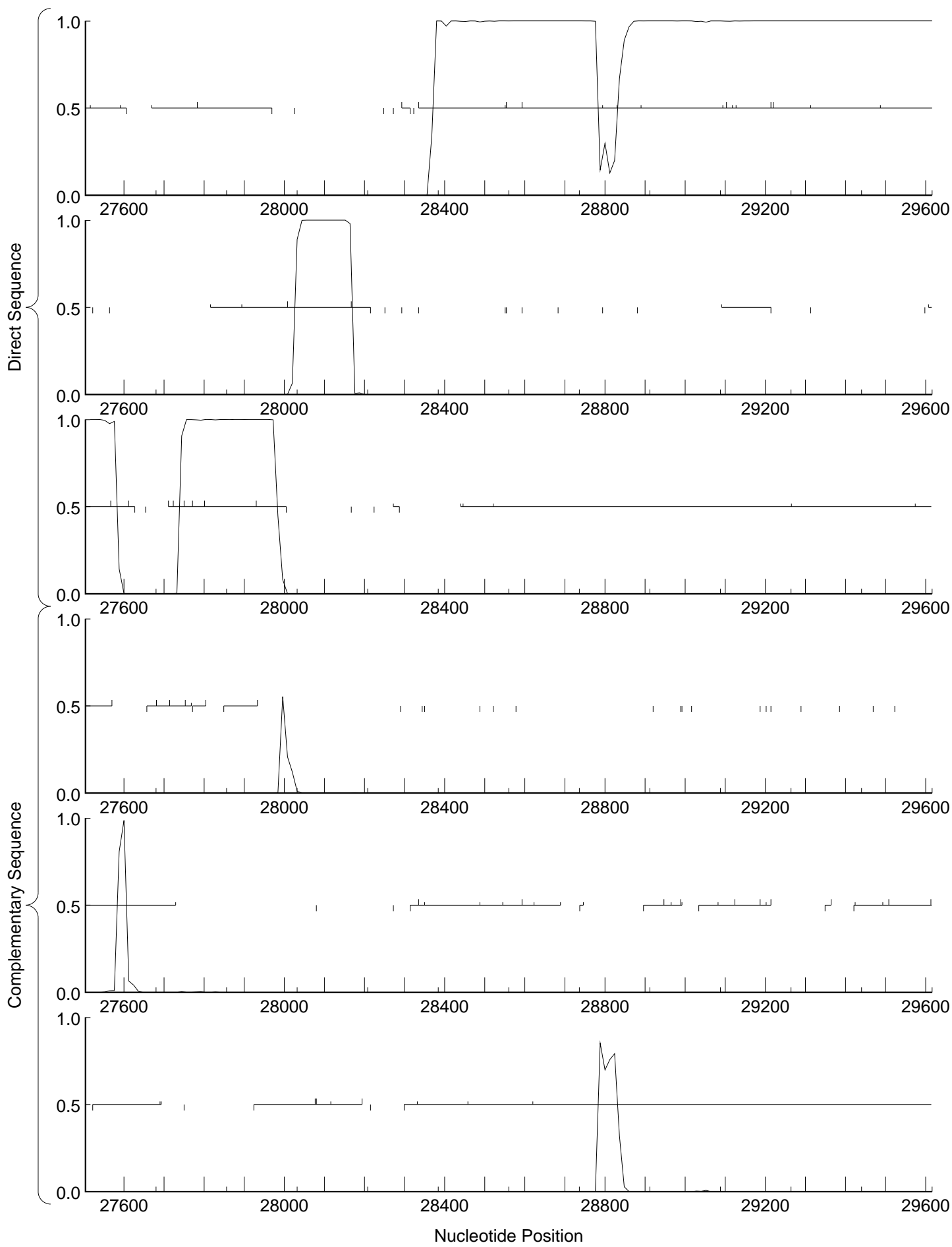
complete sequence, 41814 bp including 10-base 3' overhang (TCTCCCGGCA), Cluster EB, Order 4, Window 96, Step 12, 11/21



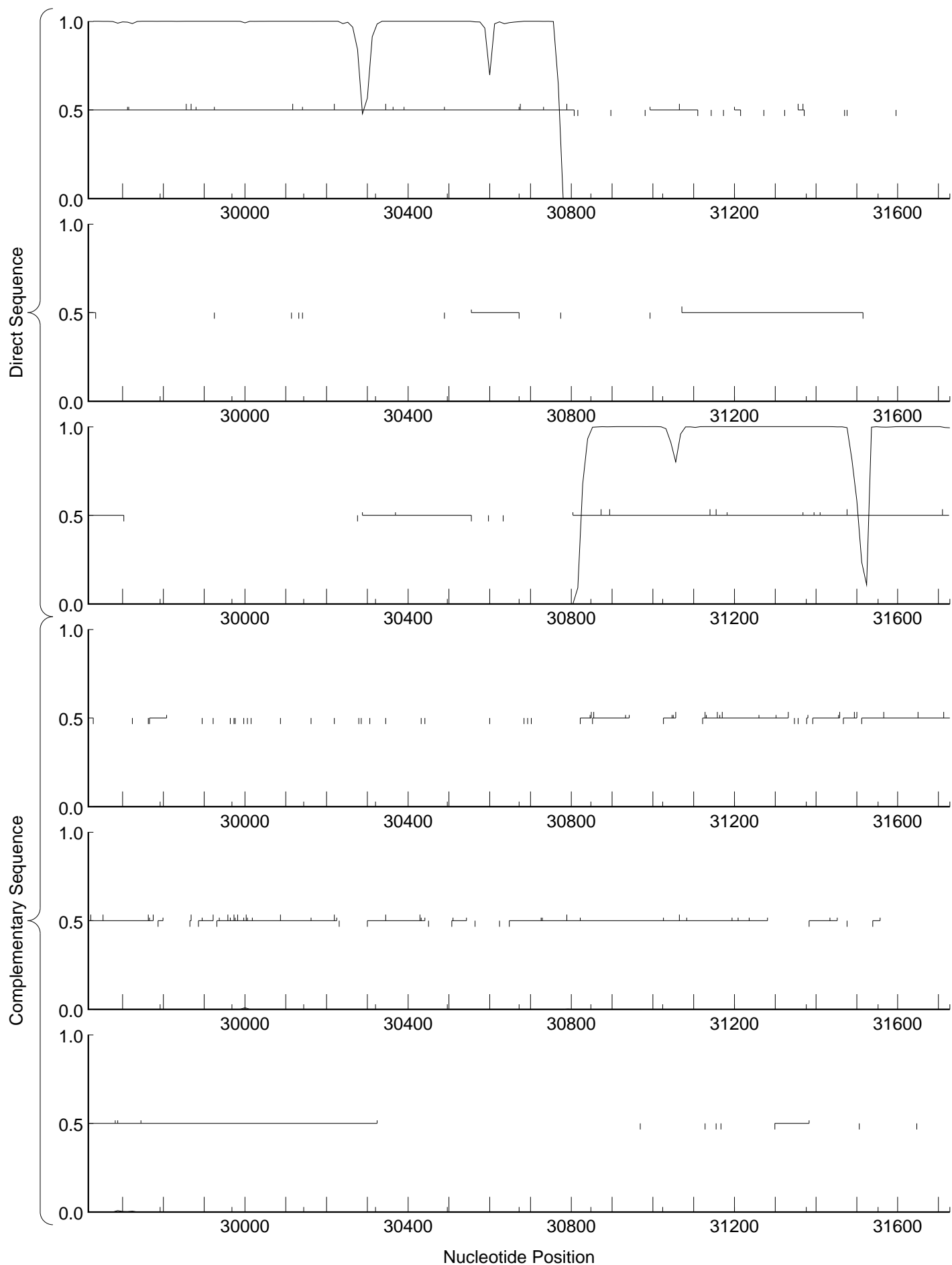


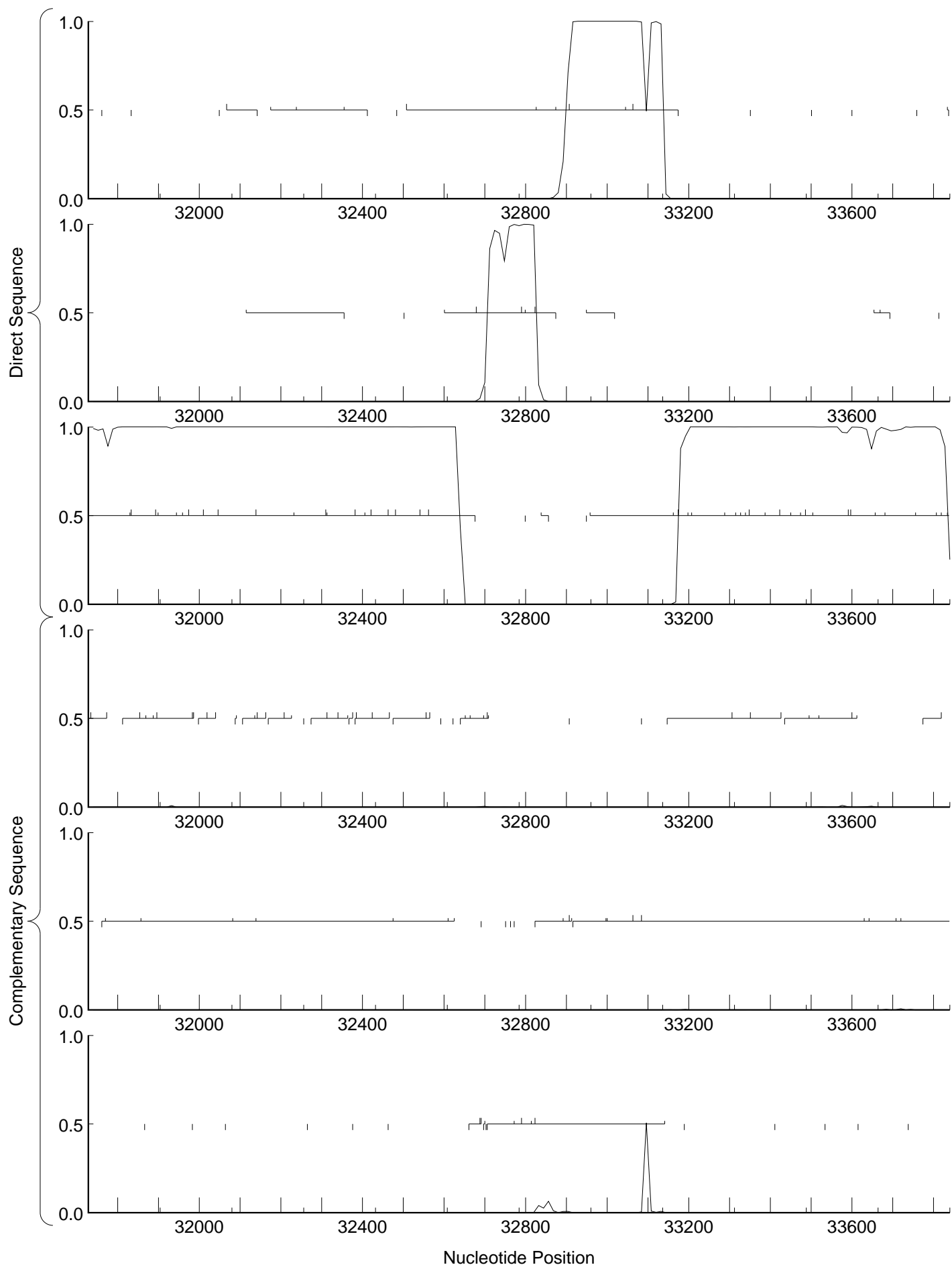


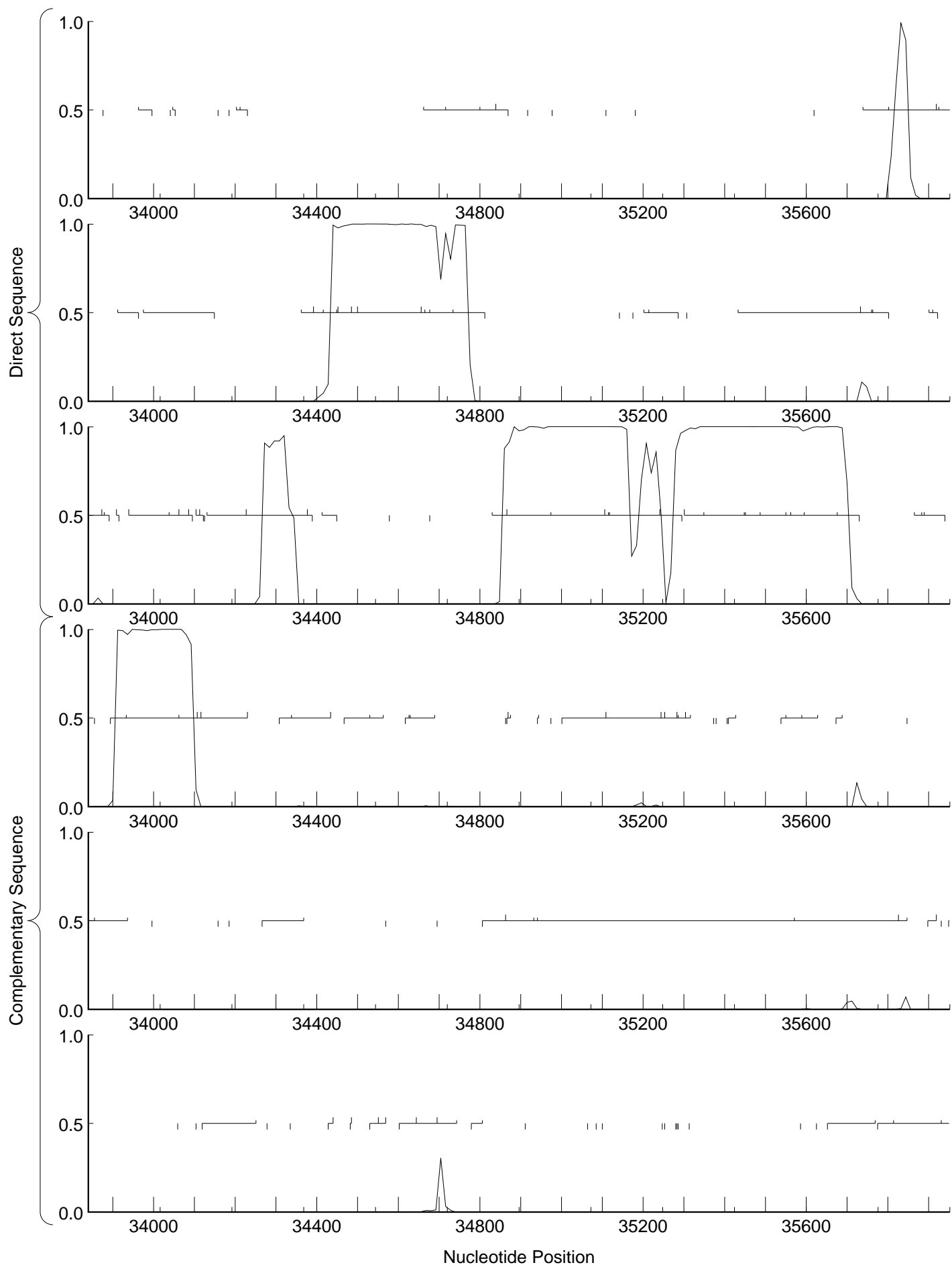


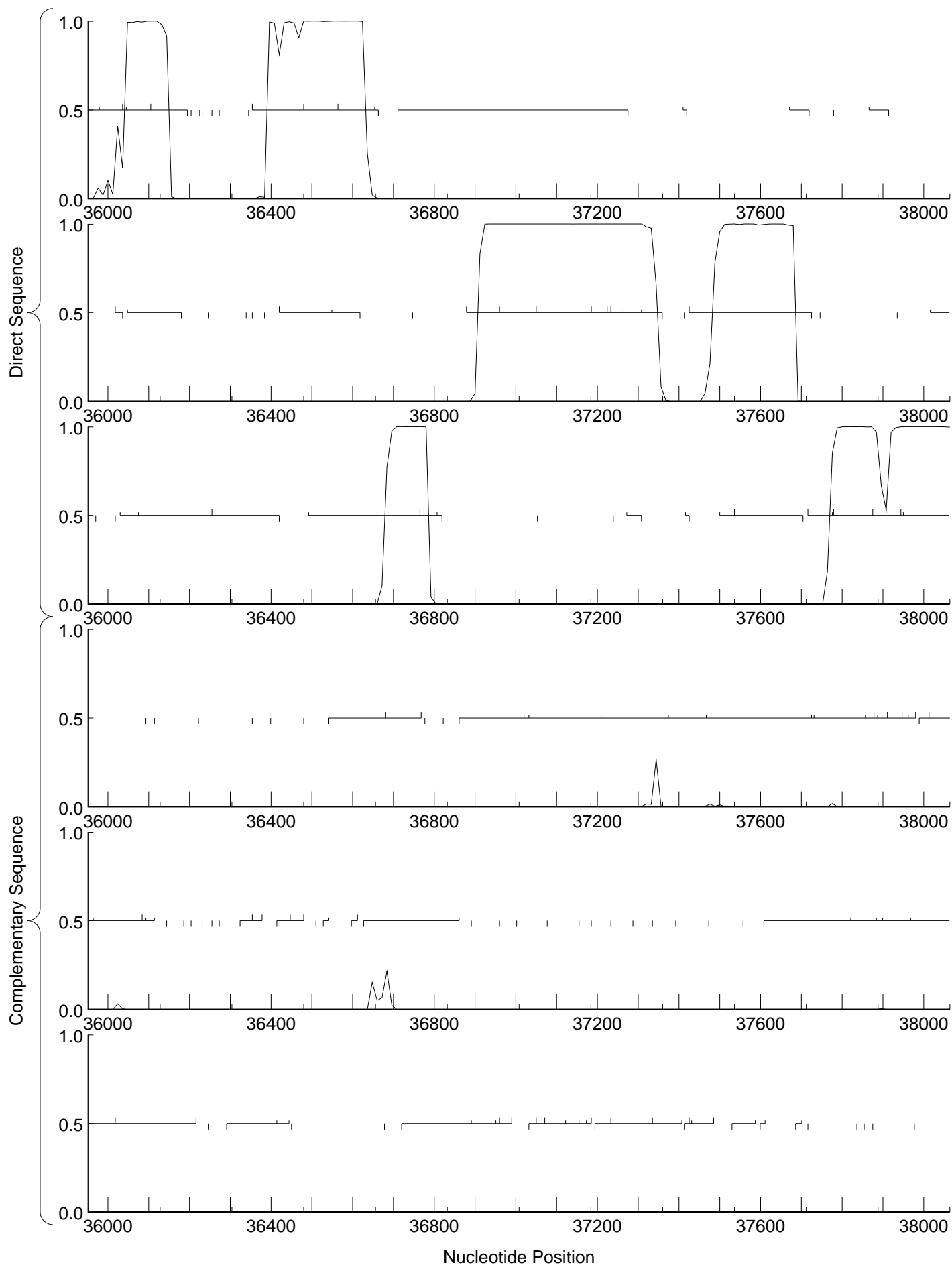


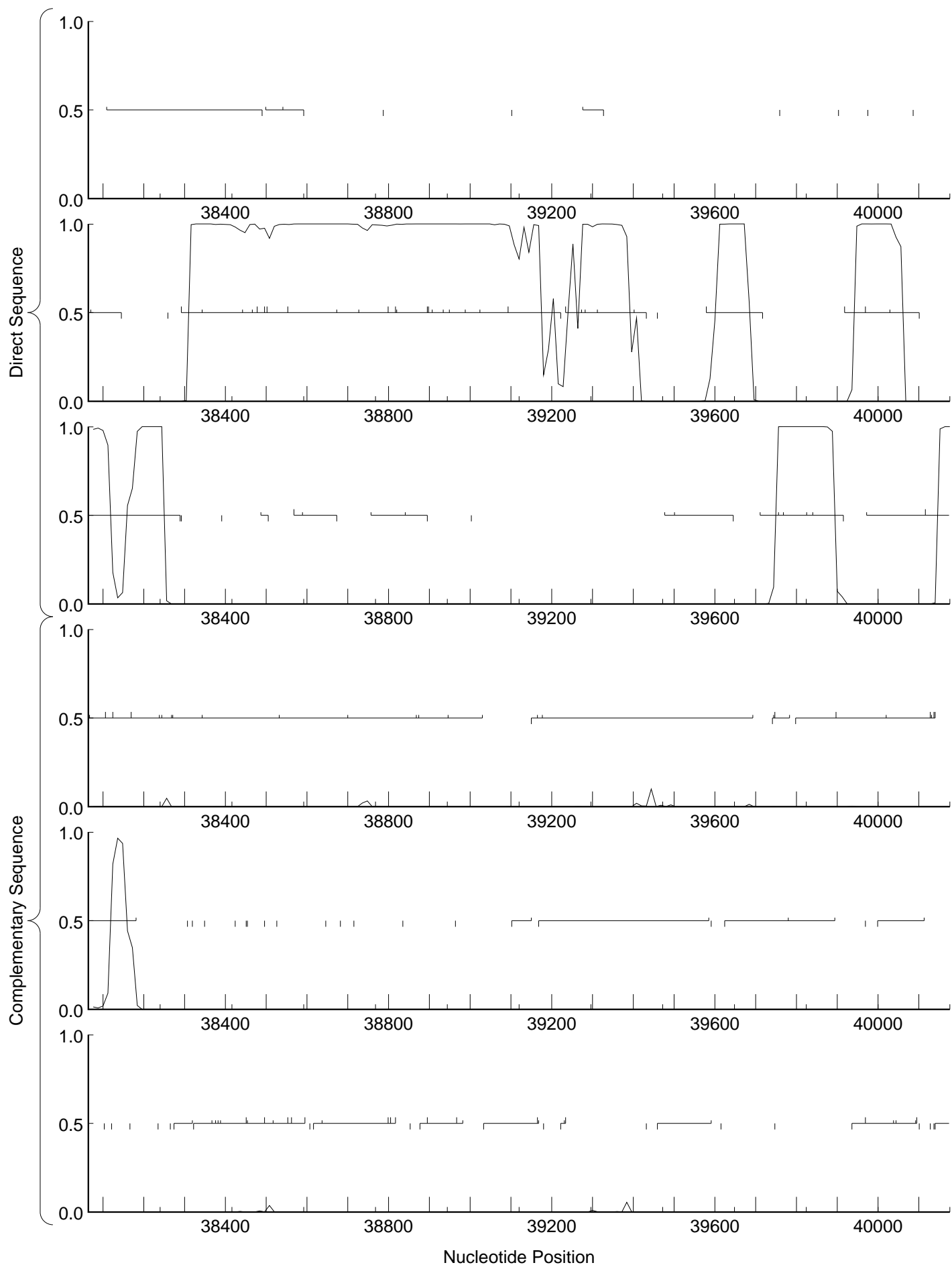
complete sequence, 41814 bp including 10-base 3' overhang (TCTCCCGGCA), Cluster EB, Order 4, Window 96, Step 12, 16/21











complete sequence, 41814 bp including 10-base 3' overhang (TCTCCCGGCA), Cluster EB, Order 4, Window 96, Step 12, 21/21

