

GeneMark

Version 2.5p (09.08.06)

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PROGRAM INFORMATION

Sequence : Mycobacterium phage Peeb complete sequence, 41876 bp including 11-base 3' overhang (CCCCATG
Analysis Date : 6/24/24 at 19:24:10
Pages : 21
Sequence Length : 41876 bp
GC Content : 66.58%

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,

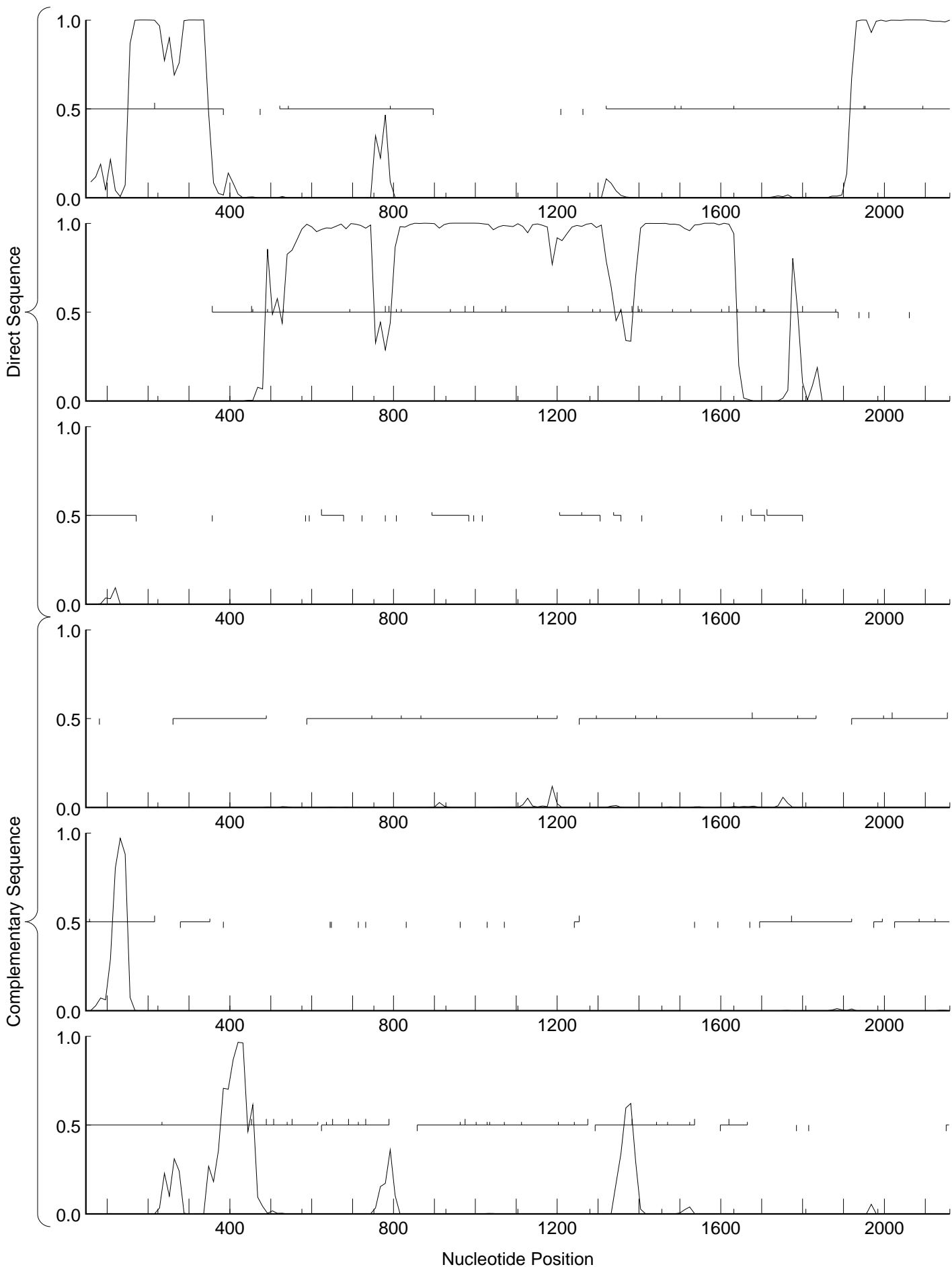
Matrix notes & comments

Training set derived by GeneMarkS, 4.27 September 2014
Tue Sep 23 15:14:37 2014

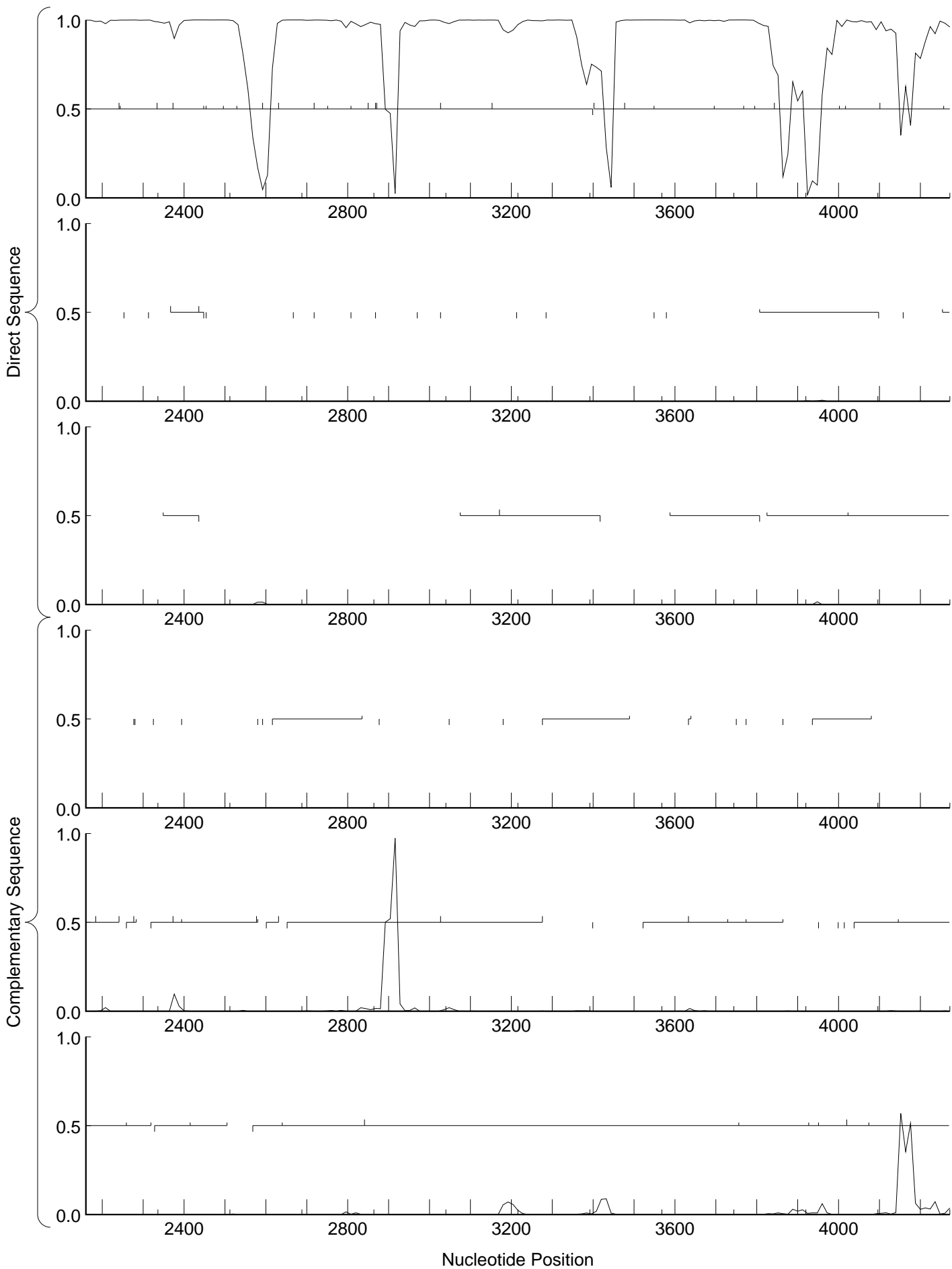
MATRIX INFORMATION

Matrix : Mycobacterium_tuberculosis_H37Rv
Author : -
Order : 4

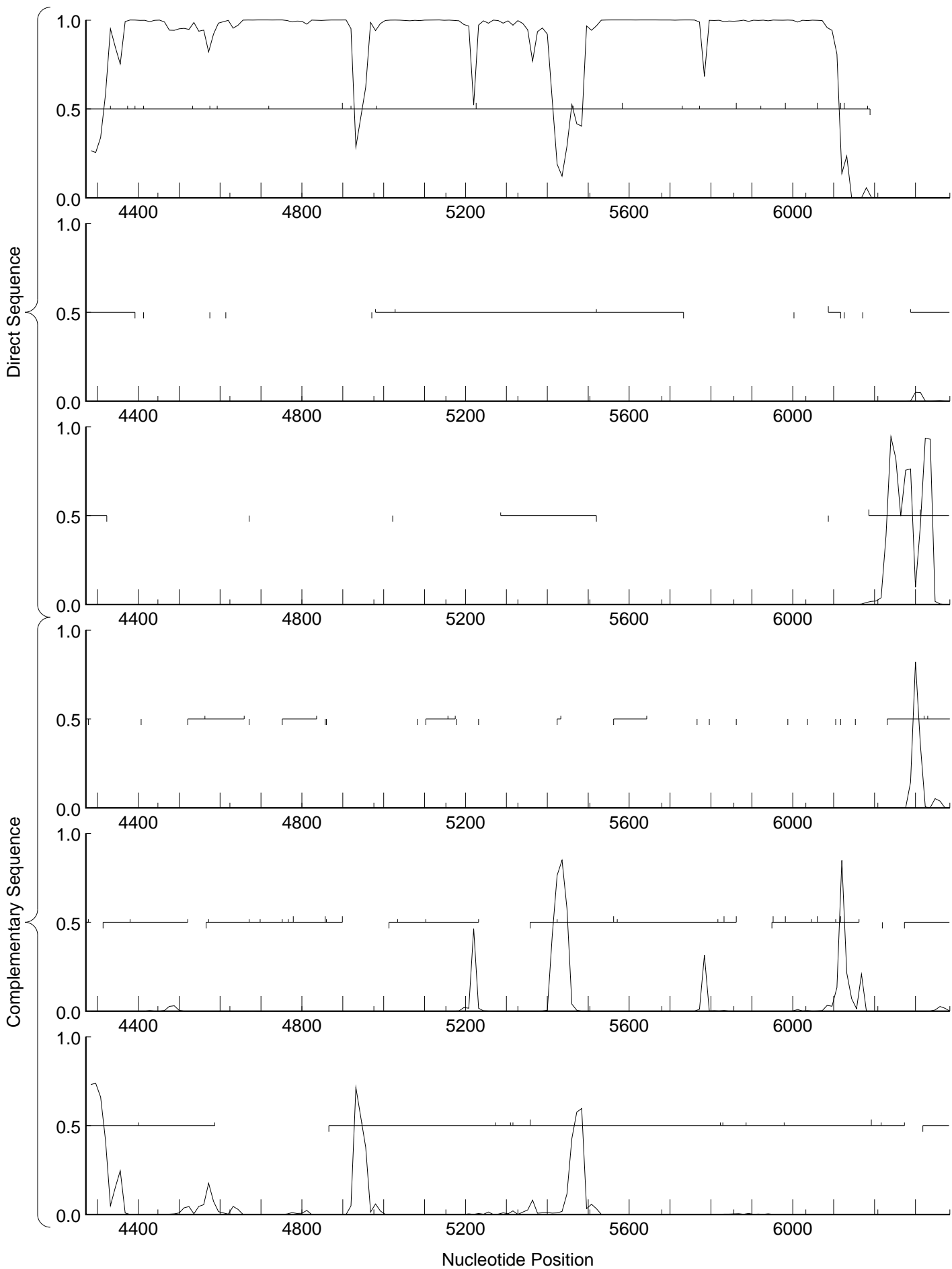
Send questions / comments to:
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Atlanta, GA 30332-0230

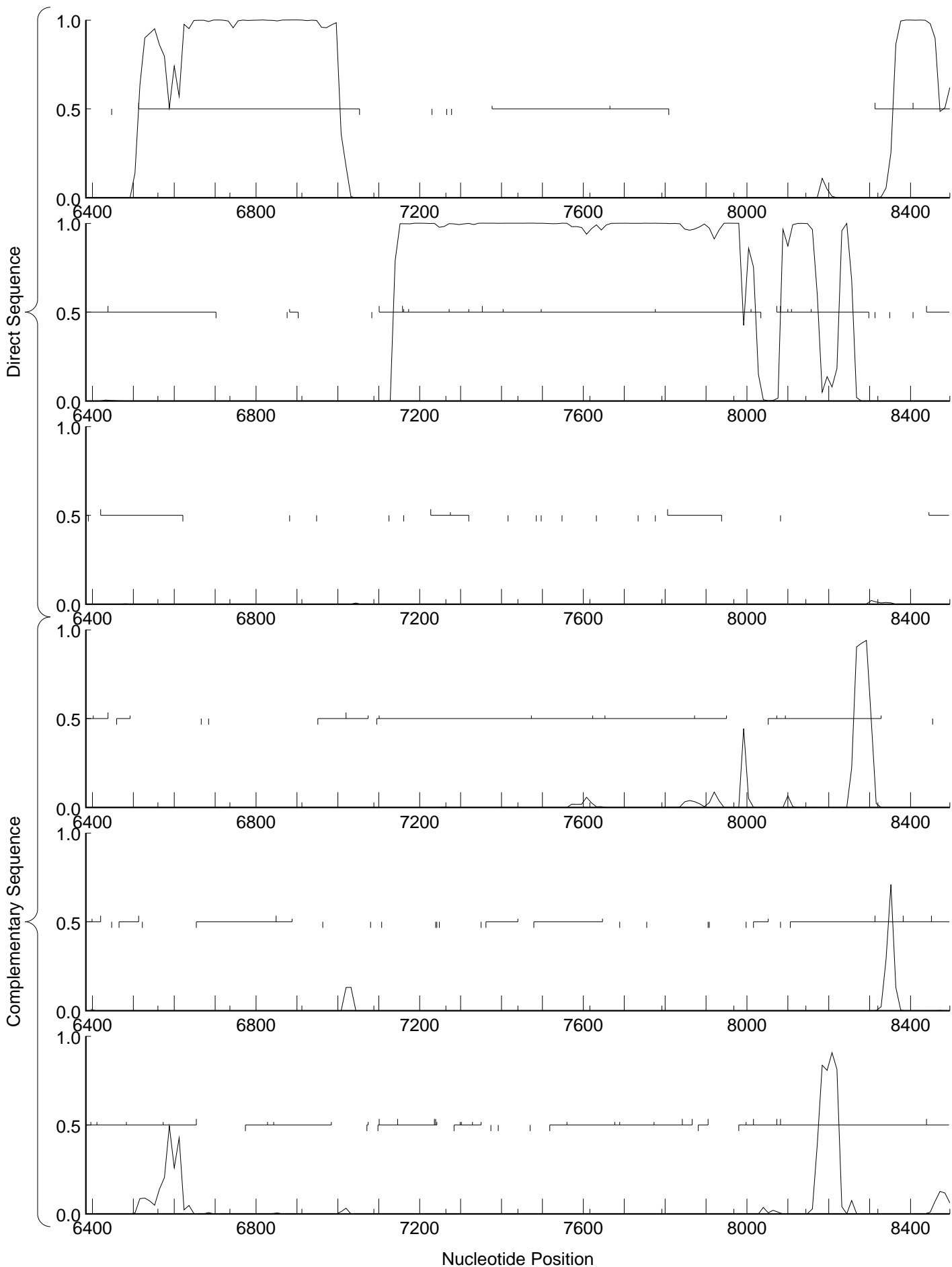


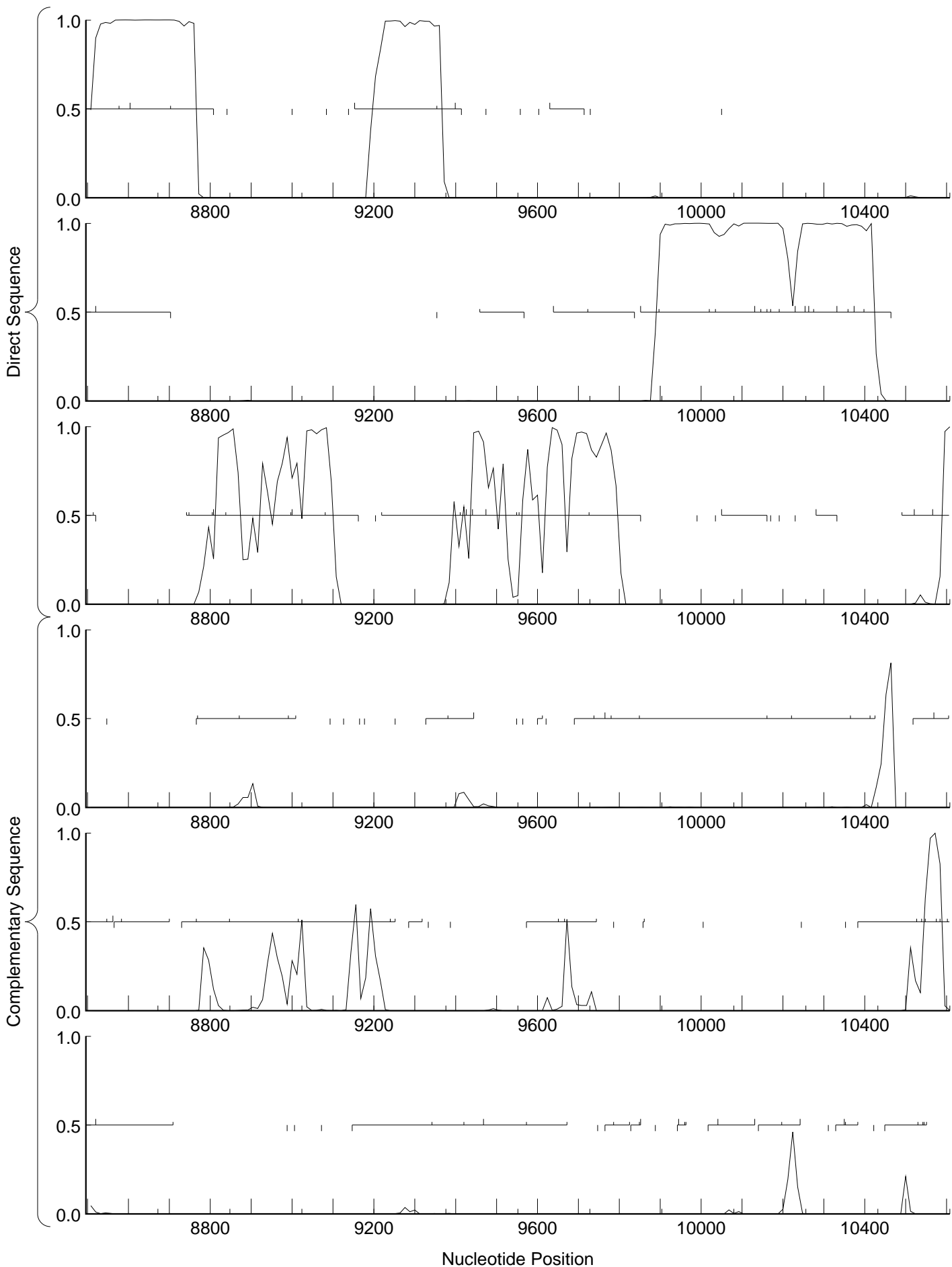
complete sequence, 41876 bp including 11-base 3' overhang (CCCCATGGCAT), Cluster G1, Order 4, Window 96, Step 12, 3/21



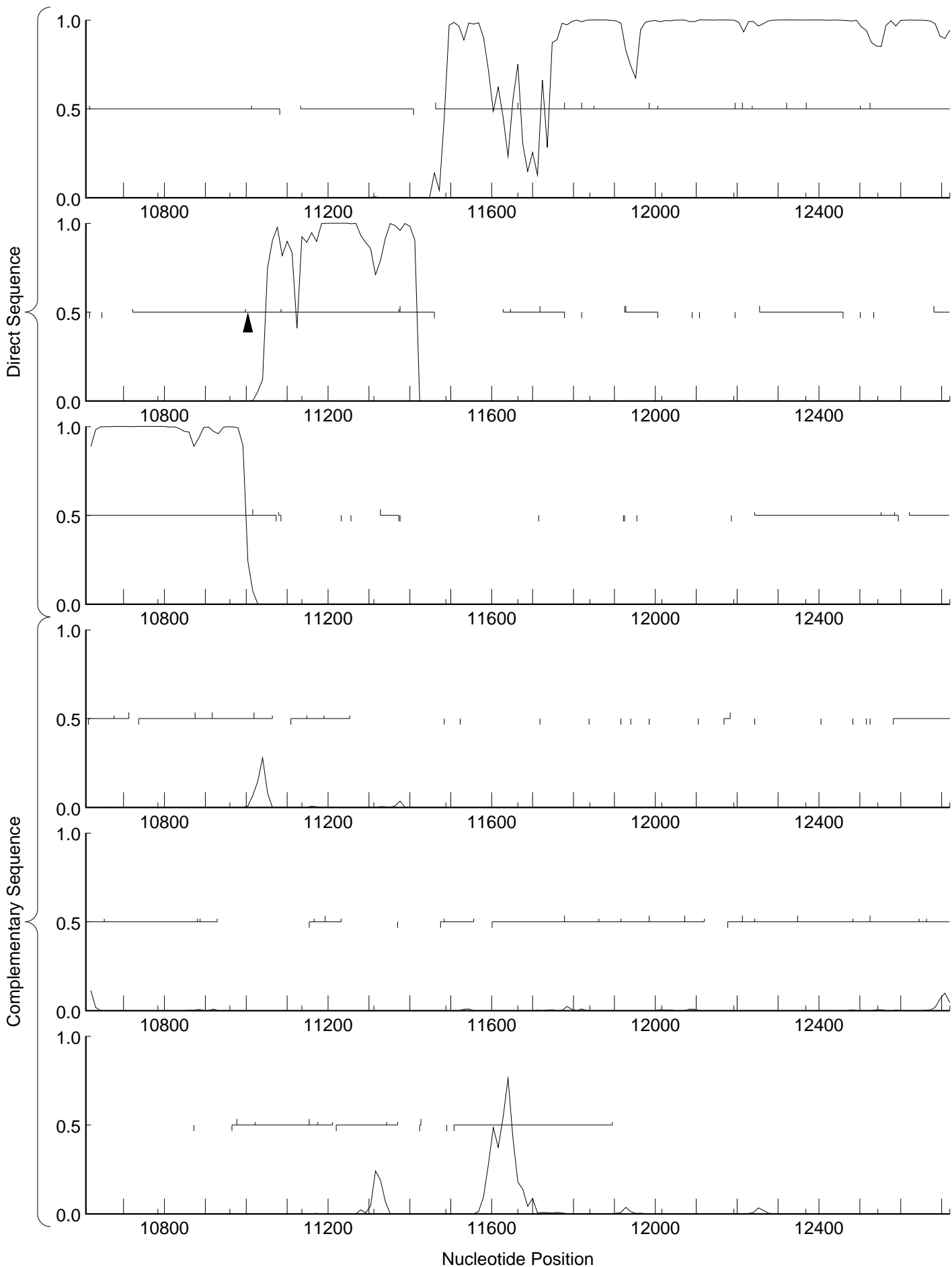
complete sequence, 41876 bp including 11-base 3' overhang (CCCCATGGCAT), Cluster G1, Order 4, Window 96, Step 12, 4/21



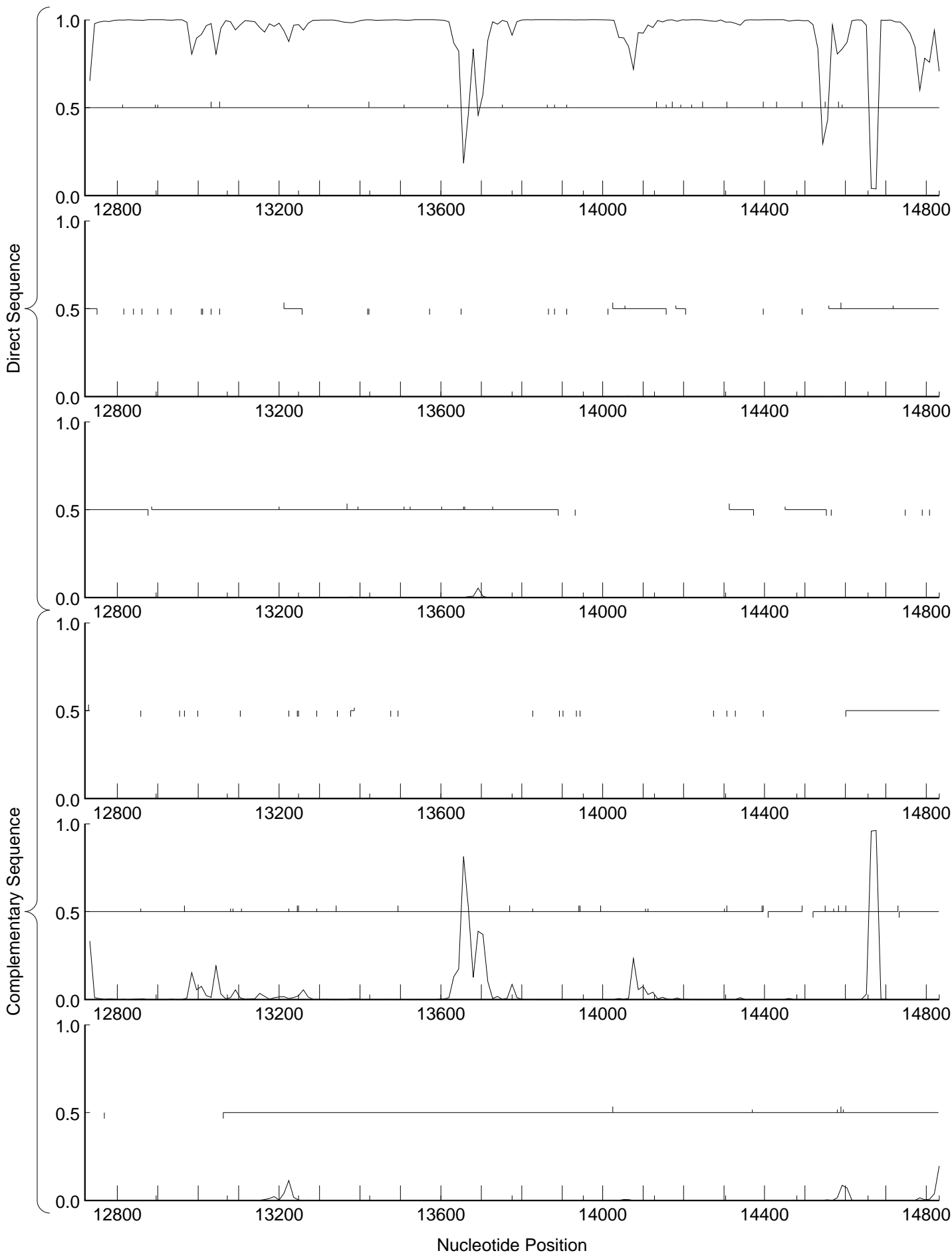


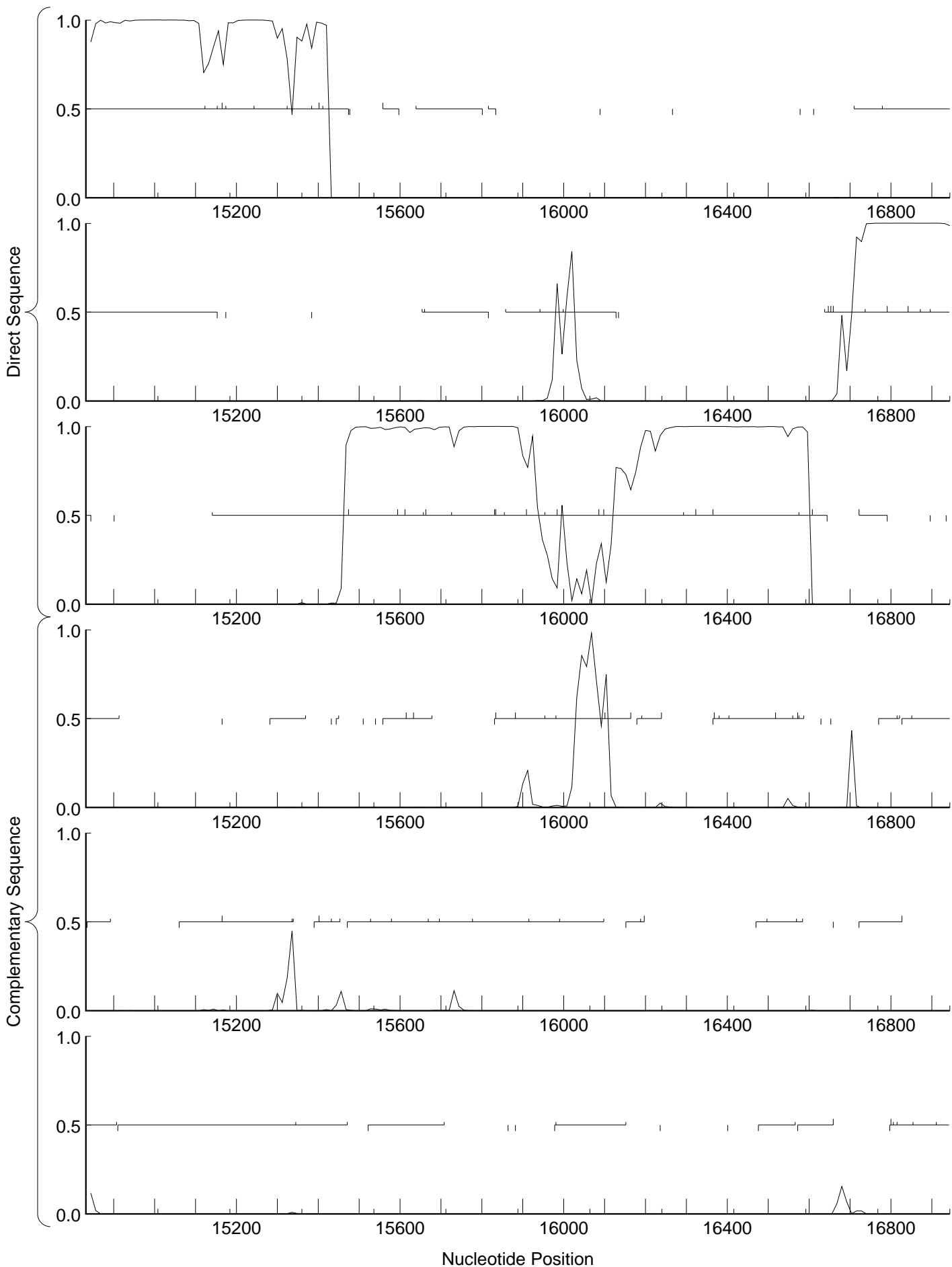


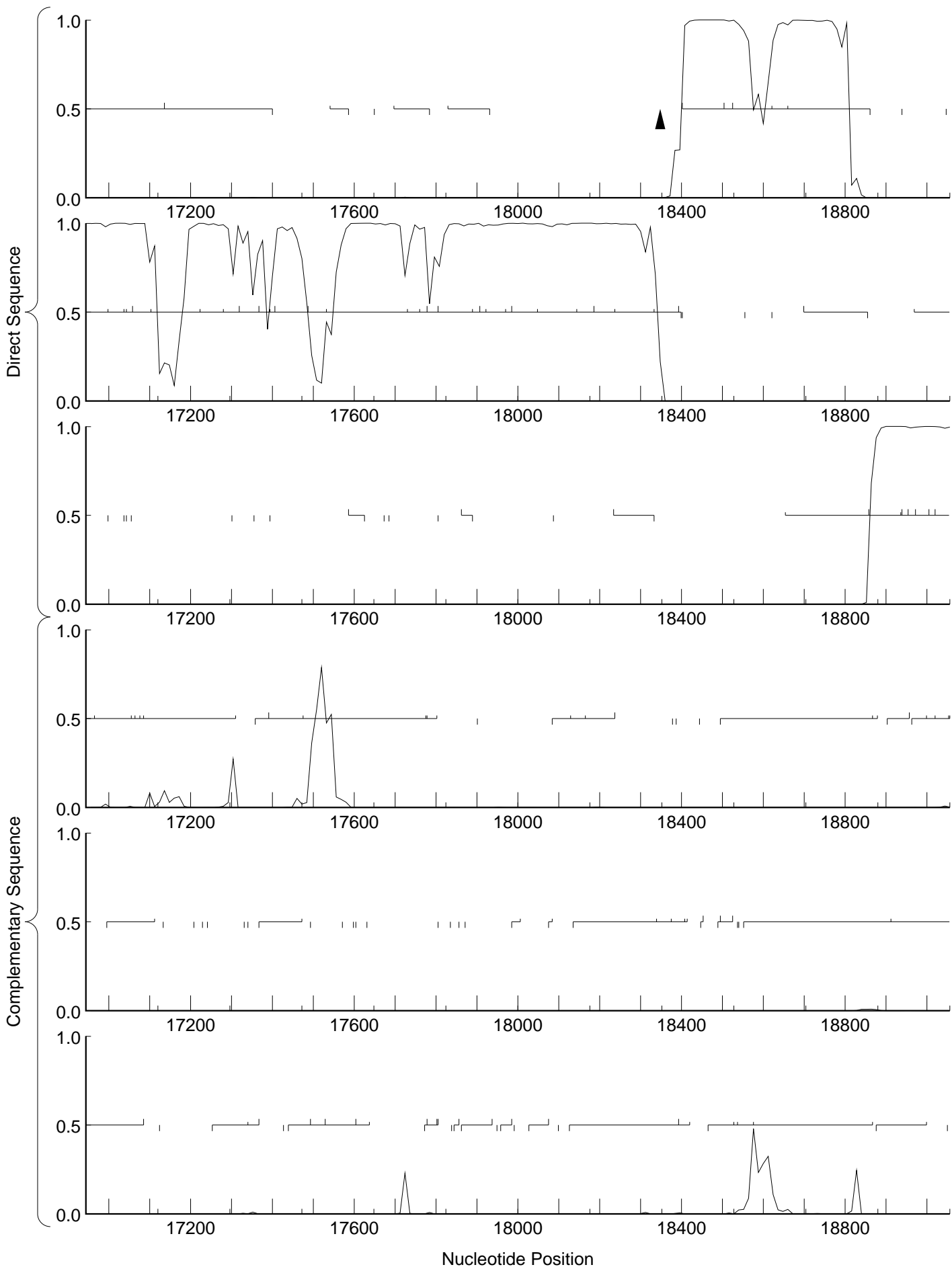
complete sequence, 41876 bp including 11-base 3' overhang (CCCCATGGGCAT), Cluster G1, Order 4, Window 96, Step 12, 7/21

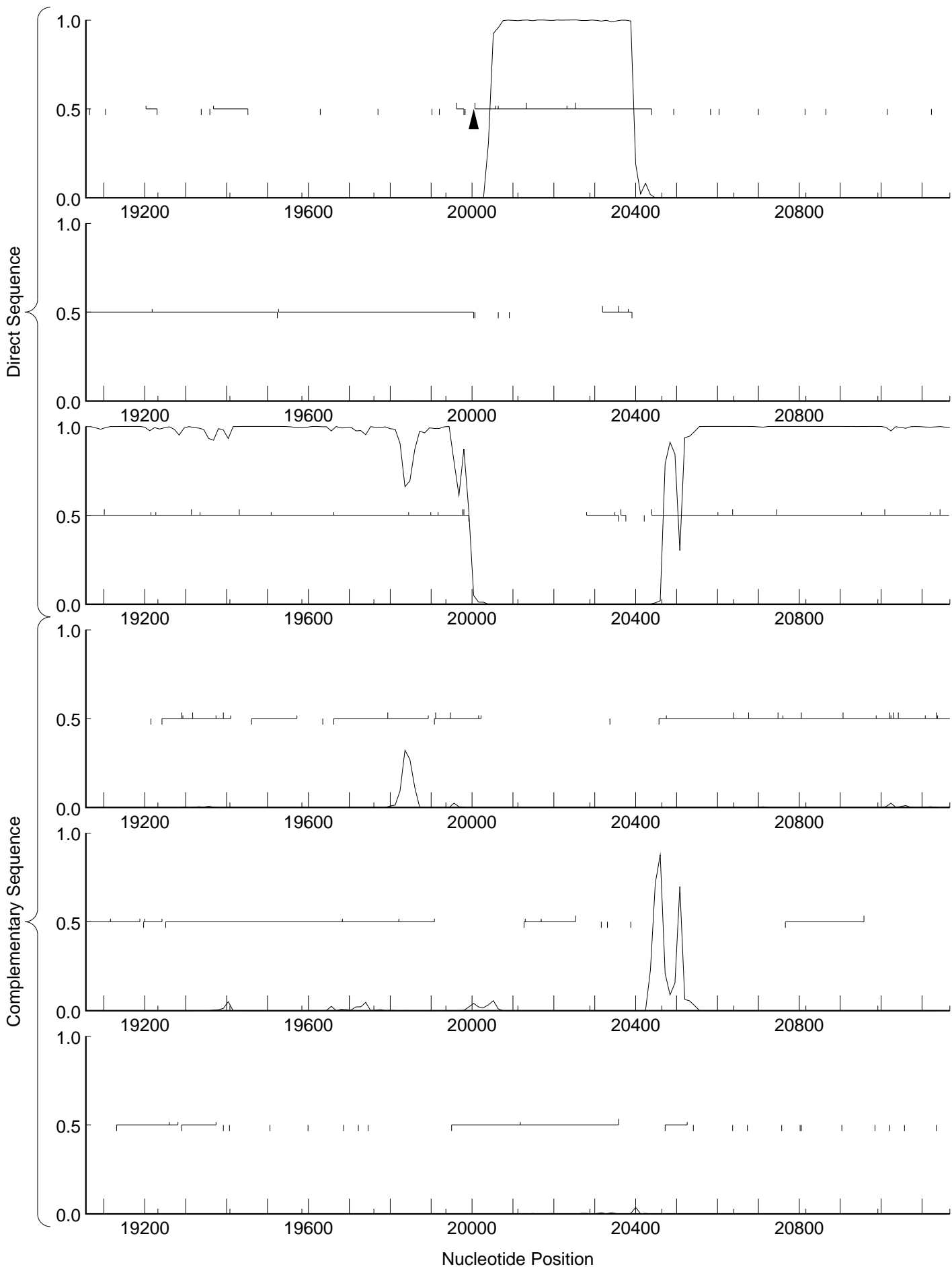


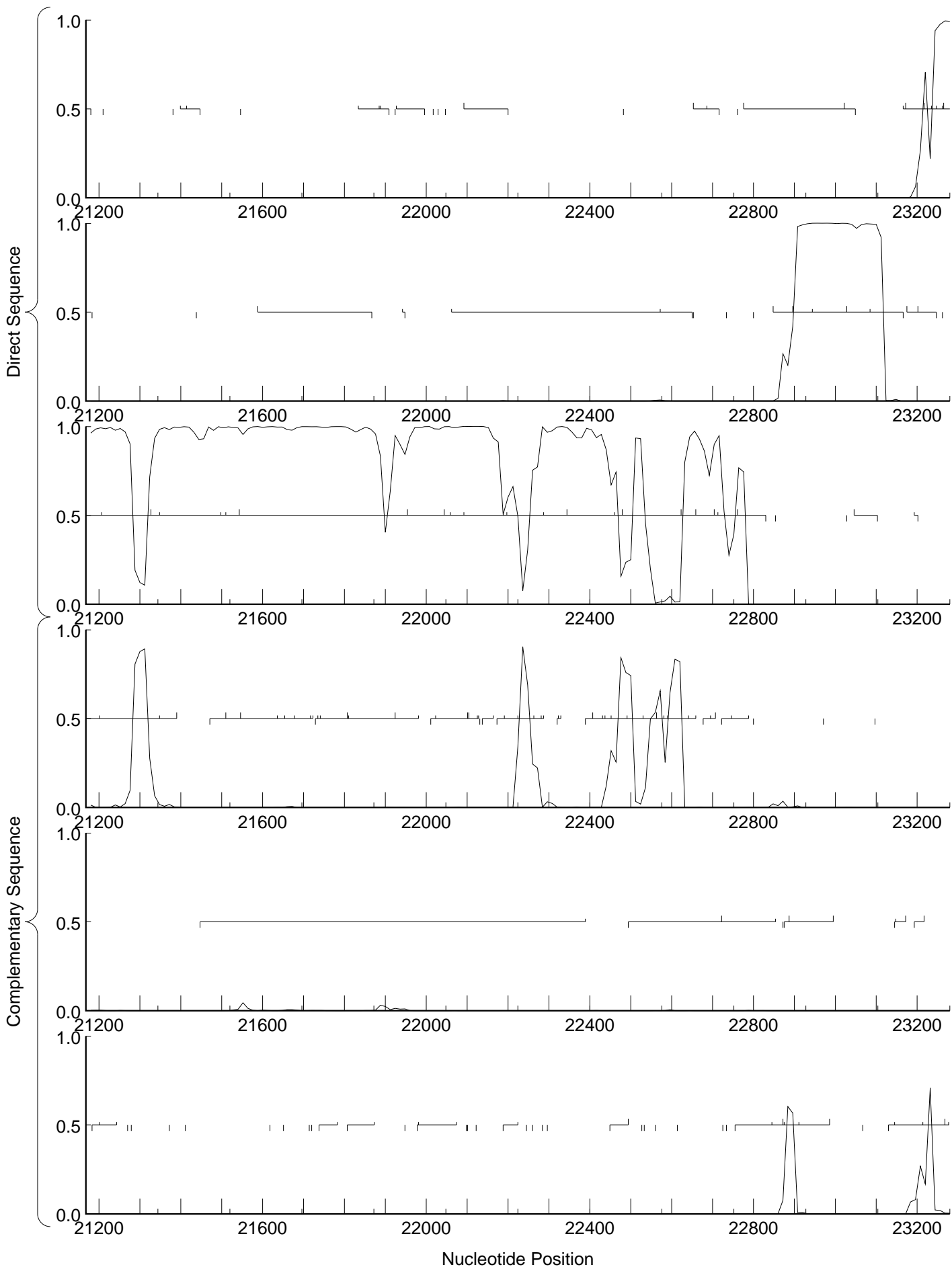
complete sequence, 41876 bp including 11-base 3' overhang (CCCCATGGGCAT), Cluster G1, Order 4, Window 96, Step 12, 8/21

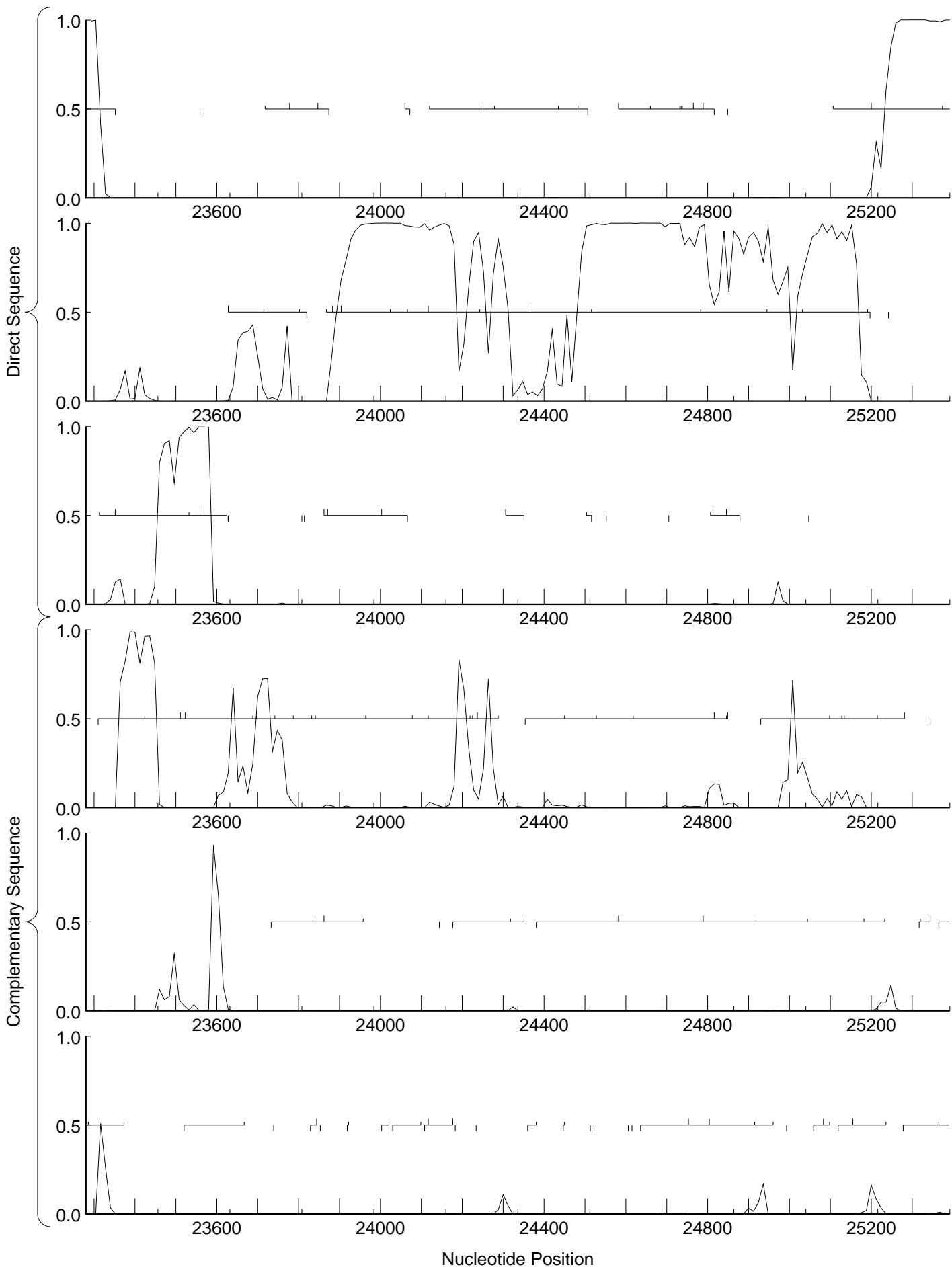


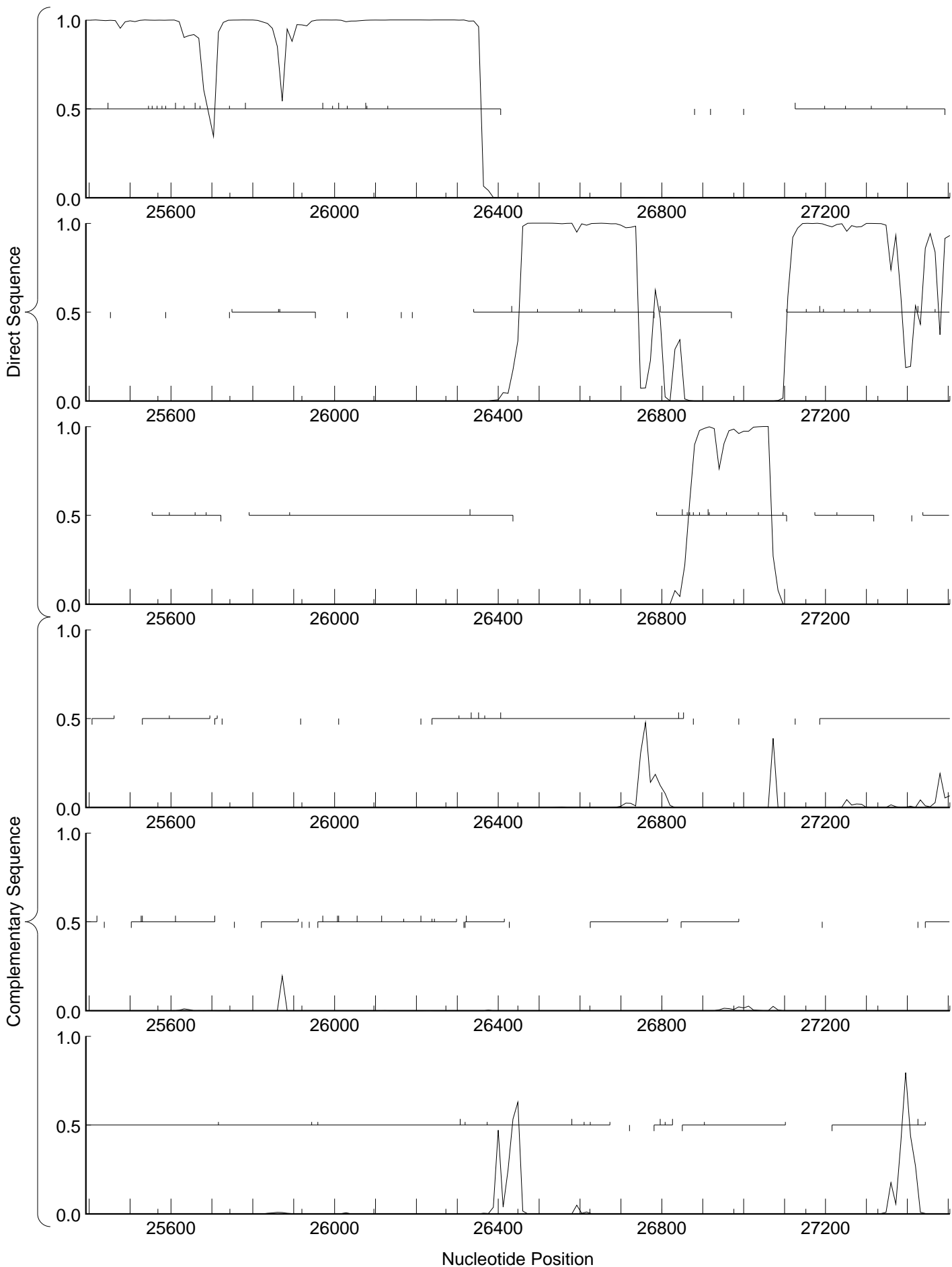












complete sequence, 41876 bp including 11-base 3' overhang (CCCCATGGCAT), Cluster G1, Order 4, Window 96, Step 12, 15/21

