

GeneMark

Version 2.5p (09.08.06)

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PROGRAM INFORMATION

Sequence : Mycobacterium phage Gengar complete sequence, 61626 bp including 11-base 3' overhang (CTCAG
Analysis Date : 2/7/25 at 13:34:10
Pages : 31
Sequence Length : 61626 bp
GC Content : 64.99%

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,

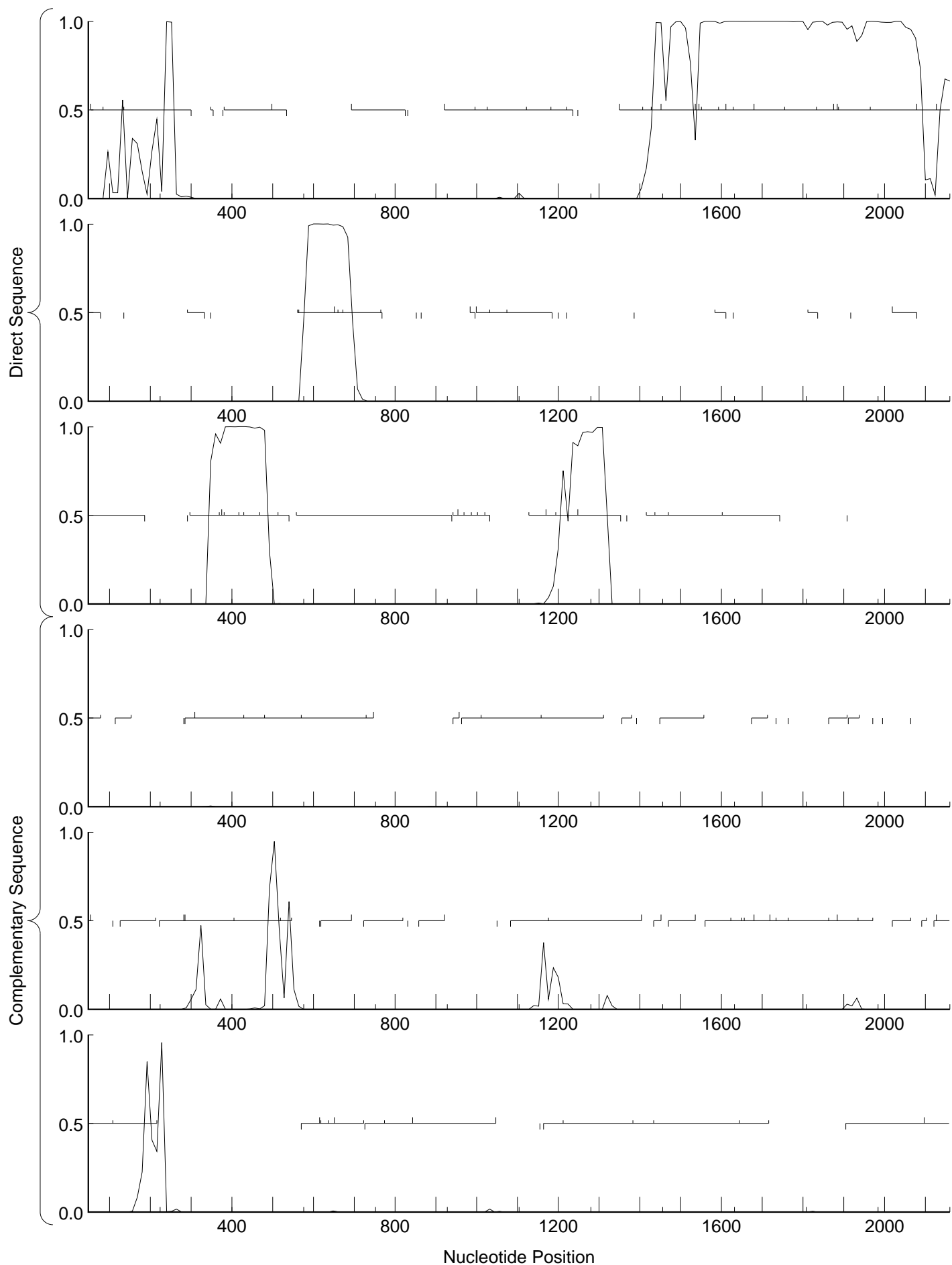
Matrix notes & comments

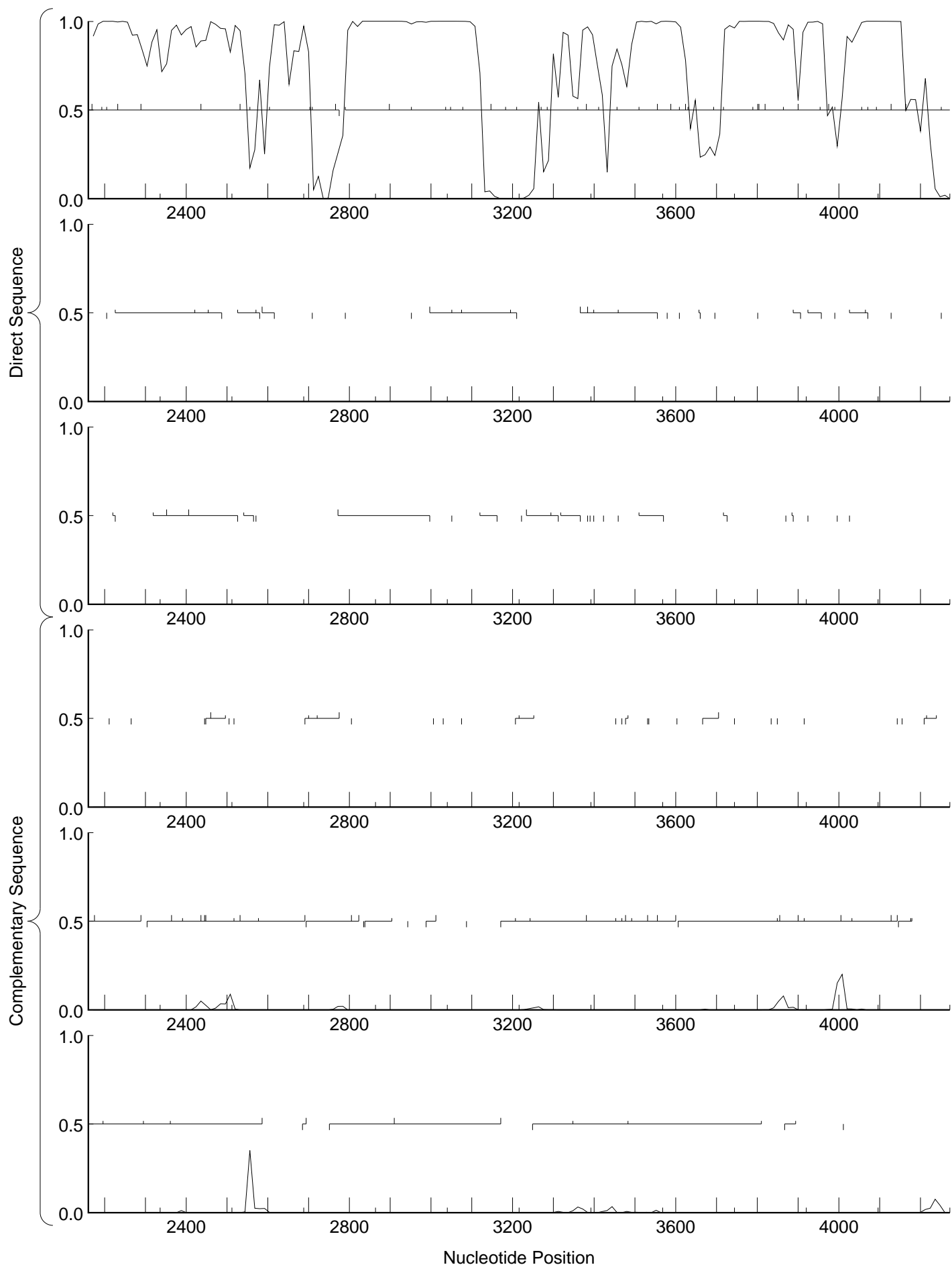
Training set derived by GeneMarkS, 4.27 September 2014
Tue Sep 23 15:23:03 2014

MATRIX INFORMATION

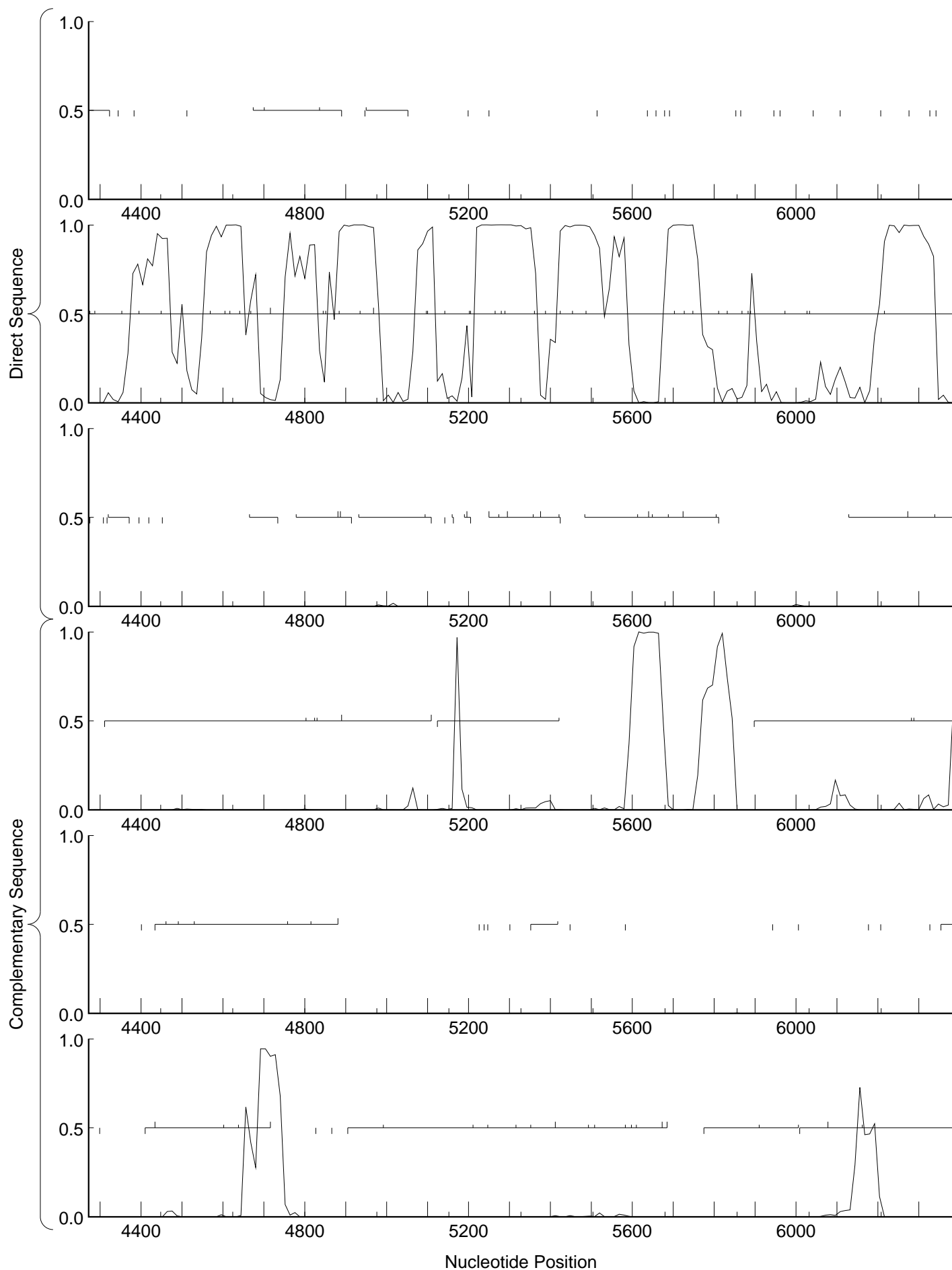
Matrix : Mycobacterium_smegmatis_MC2_155
Author : -
Order : 4

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

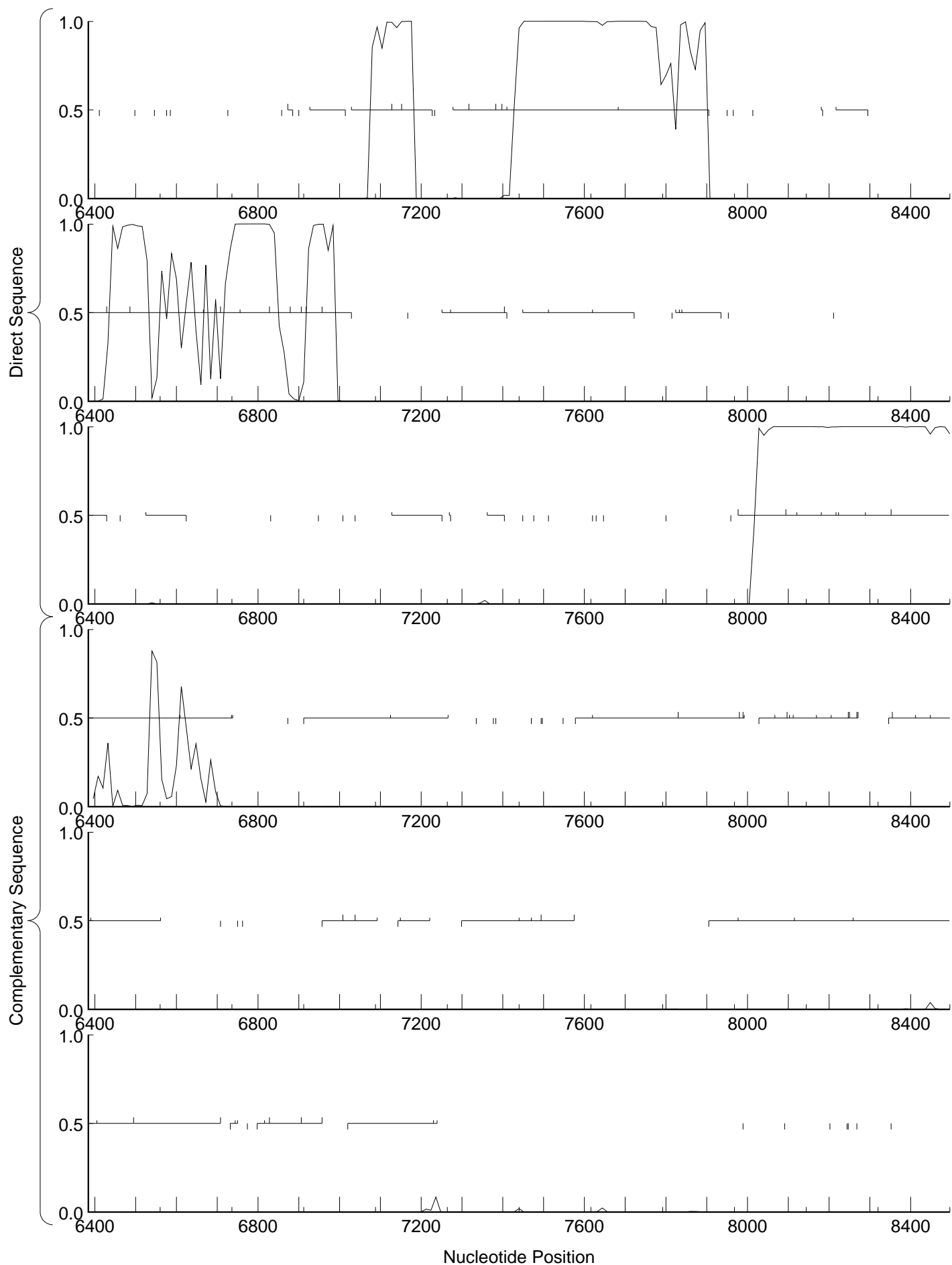


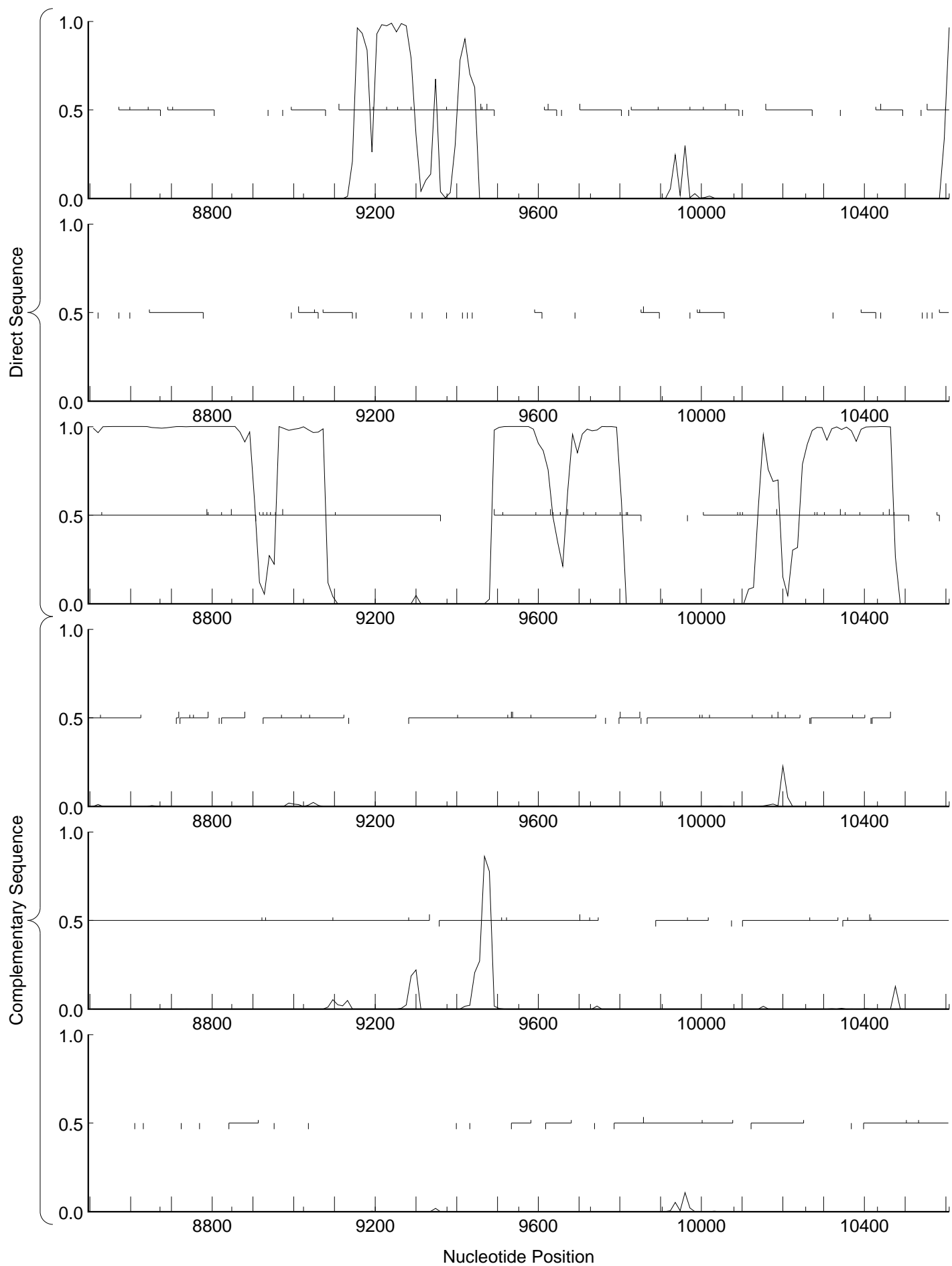


complete sequence, 61626 bp including 11-base 3' overhang (CTCAGTGGCAT), Cluster K5, Order 4, Window 96, Step 12, 4/31

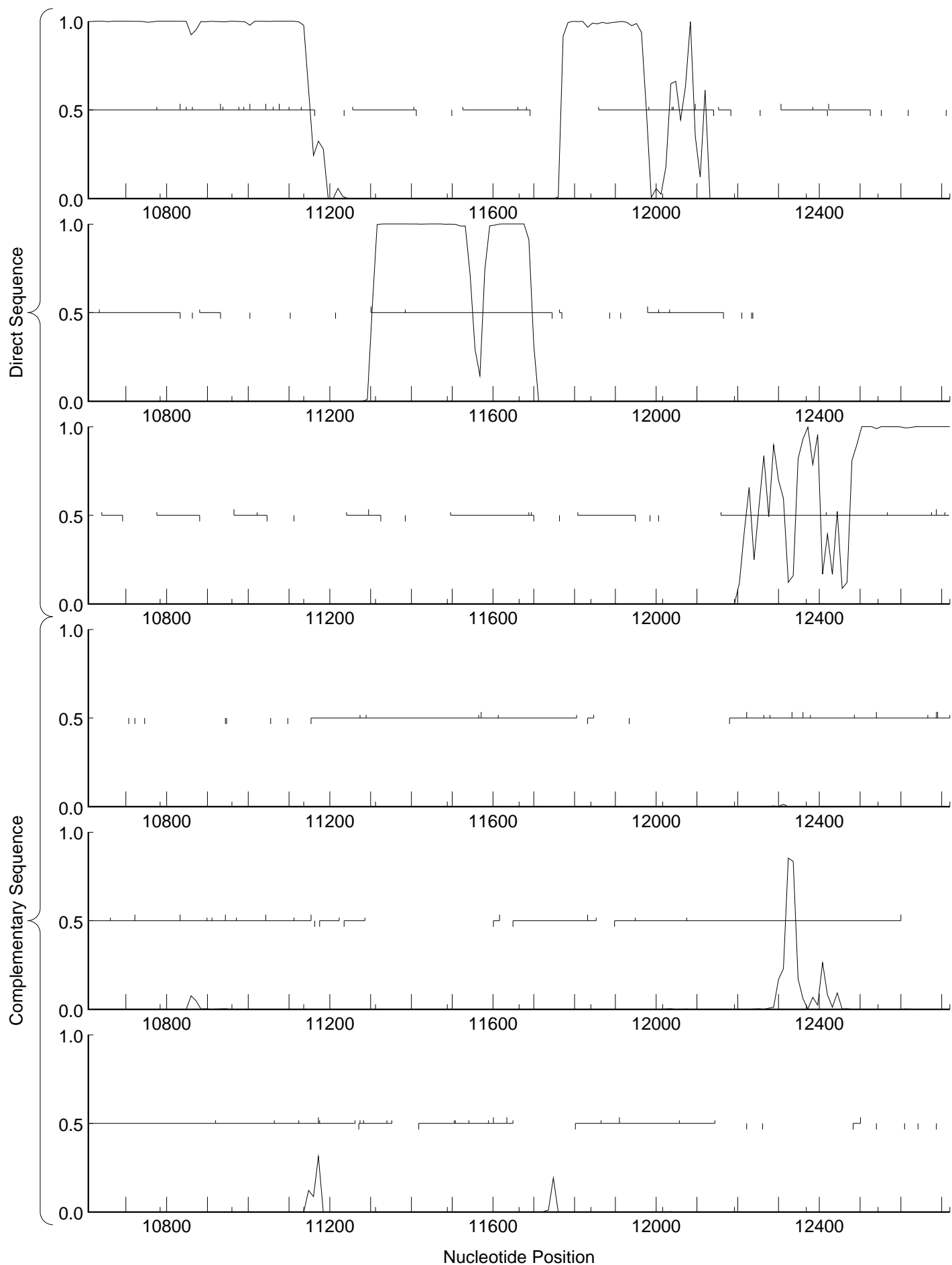


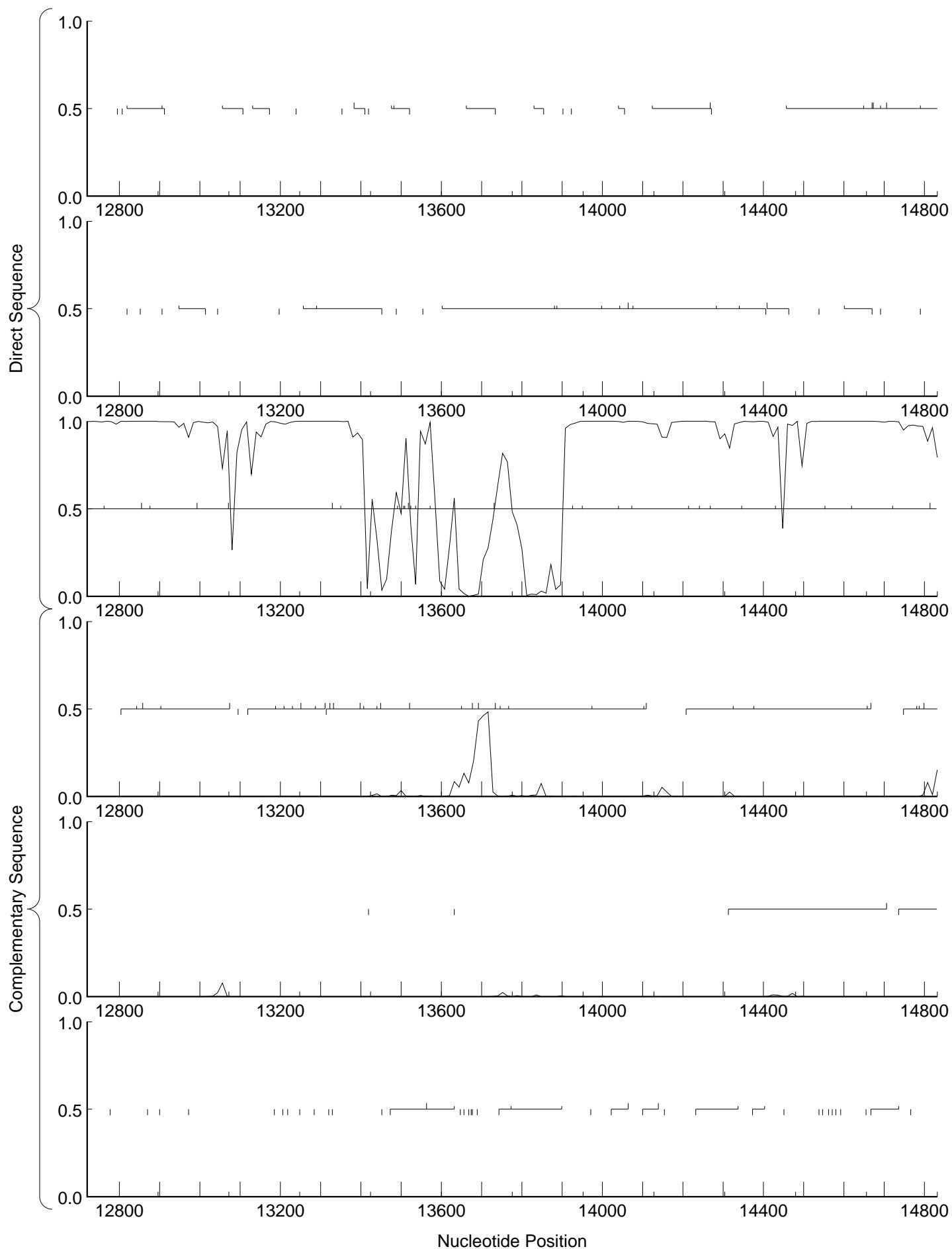
complete sequence, 61626 bp including 11-base 3' overhang (CTCAGTGGCAT), Cluster K5, Order 4, Window 96, Step 12, 5/31





complete sequence, 61626 bp including 11-base 3' overhang (CTCAGTGGCAT), Cluster K5, Order 4, Window 96, Step 12, 7/31





complete sequence, 61626 bp including 11-base 3' overhang (CTCAGTGGCAT), Cluster K5, Order 4, Window 96, Step 12, 9/31

