GeneMark

Version 2.5p (09.08.06) *Copyright 1993 - M. Borodovsky, J. McIninch*

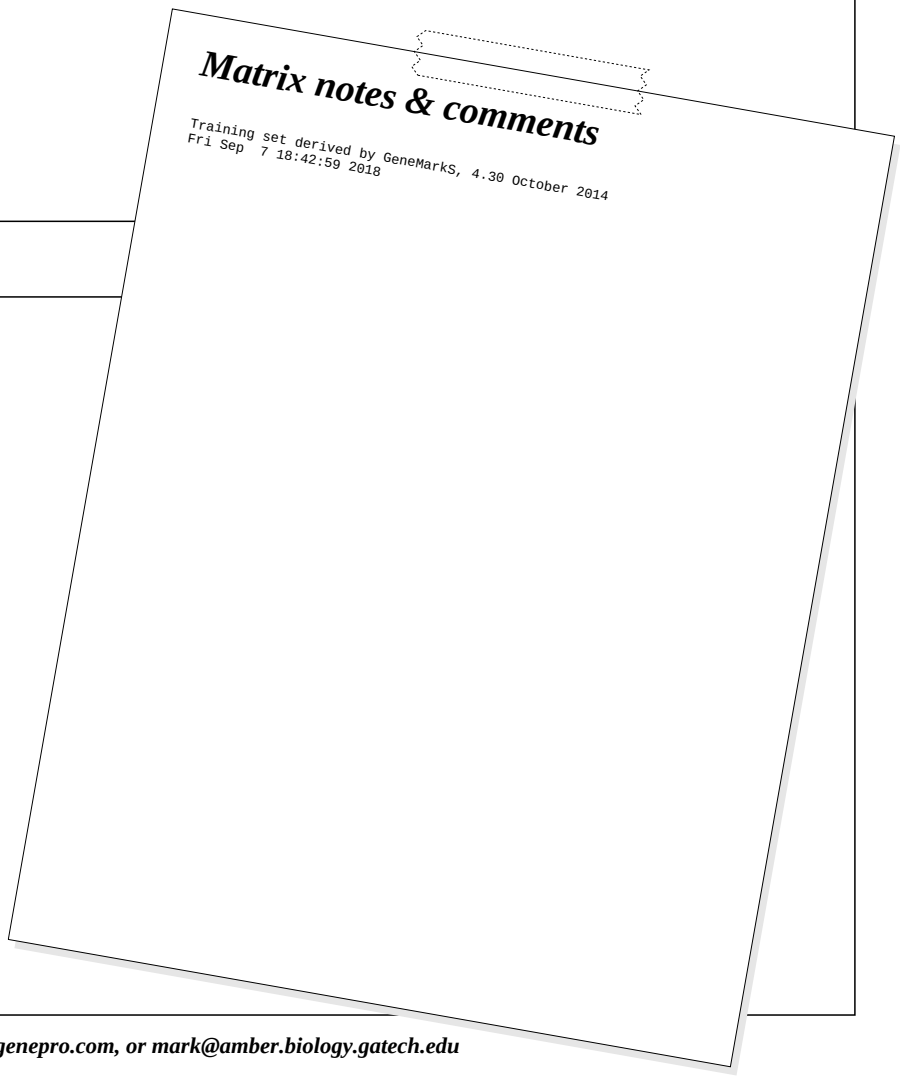
PROGRAM INFORMATION

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Sequence         :  Microbacterium phage OlinDD complete sequence, 63123 bp including 3158 bp direct terminal r
Analysis Date    :  3/29/24 at 5:32:47
Pages            :  31
Sequence Length  :  63123 bp
GC Content       :   64.60%

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,
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MATRIX INFORMATION

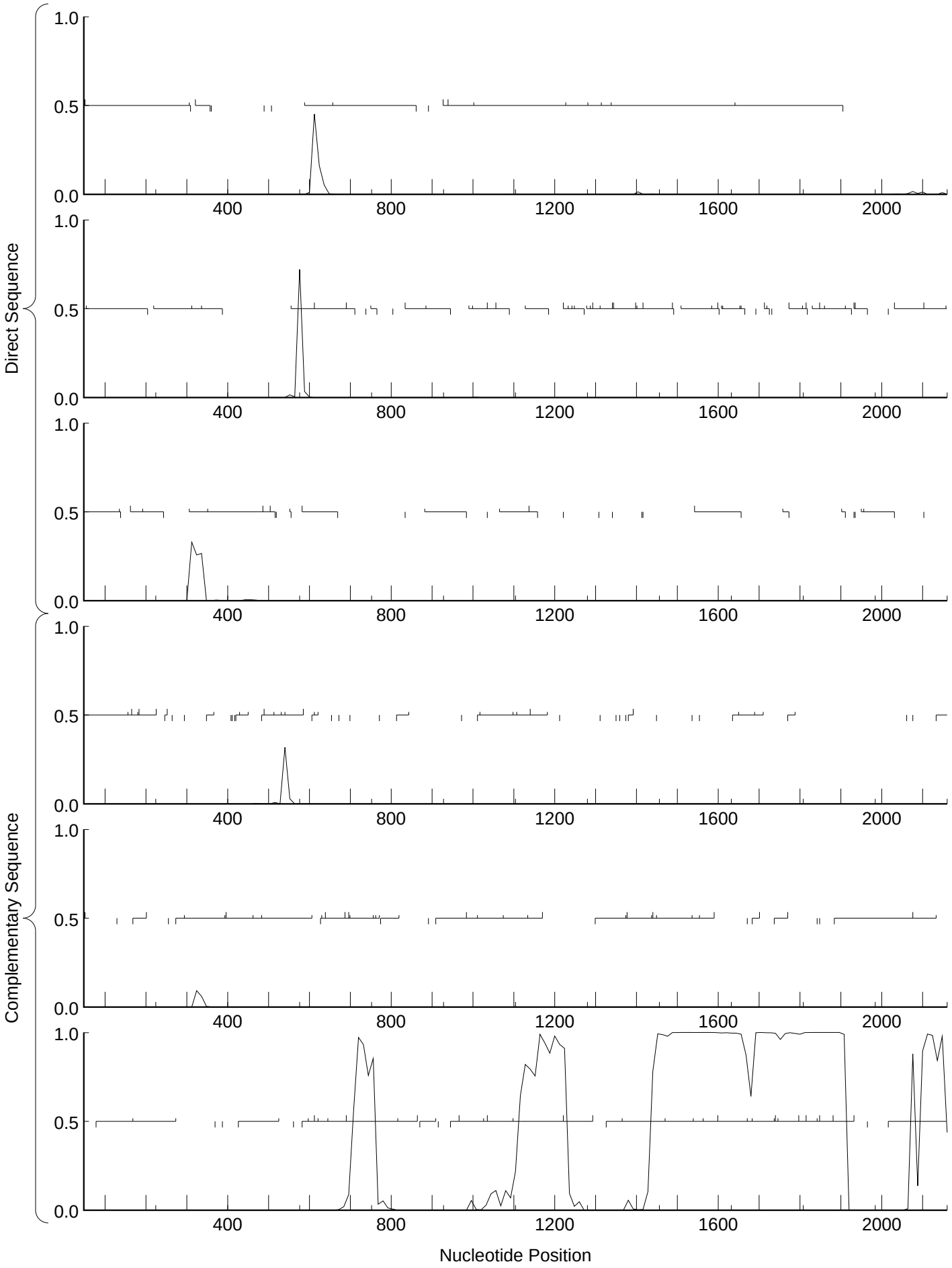
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Matrix     :  -
Author     :  -

Order      :  4
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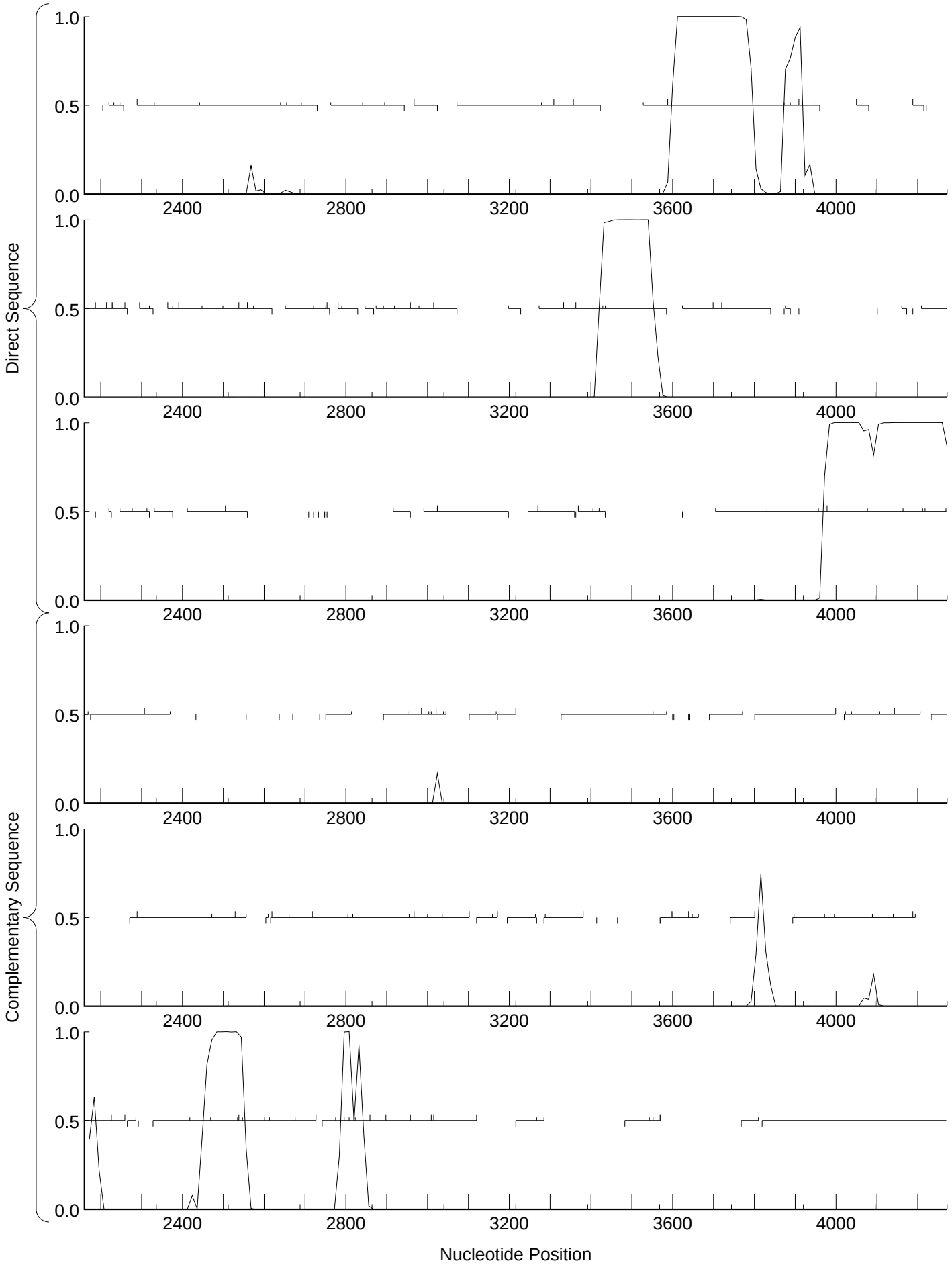

```
  Send questions / comments to:
      Dr. M. Borodovsky
      Georgia Institute of Technology
      School of Biology
      Atlanta, GA  30332-0230
```


For help or information, send electronic mail to: custserv@genepro.com, or mark@amber.biology.gatech.edu

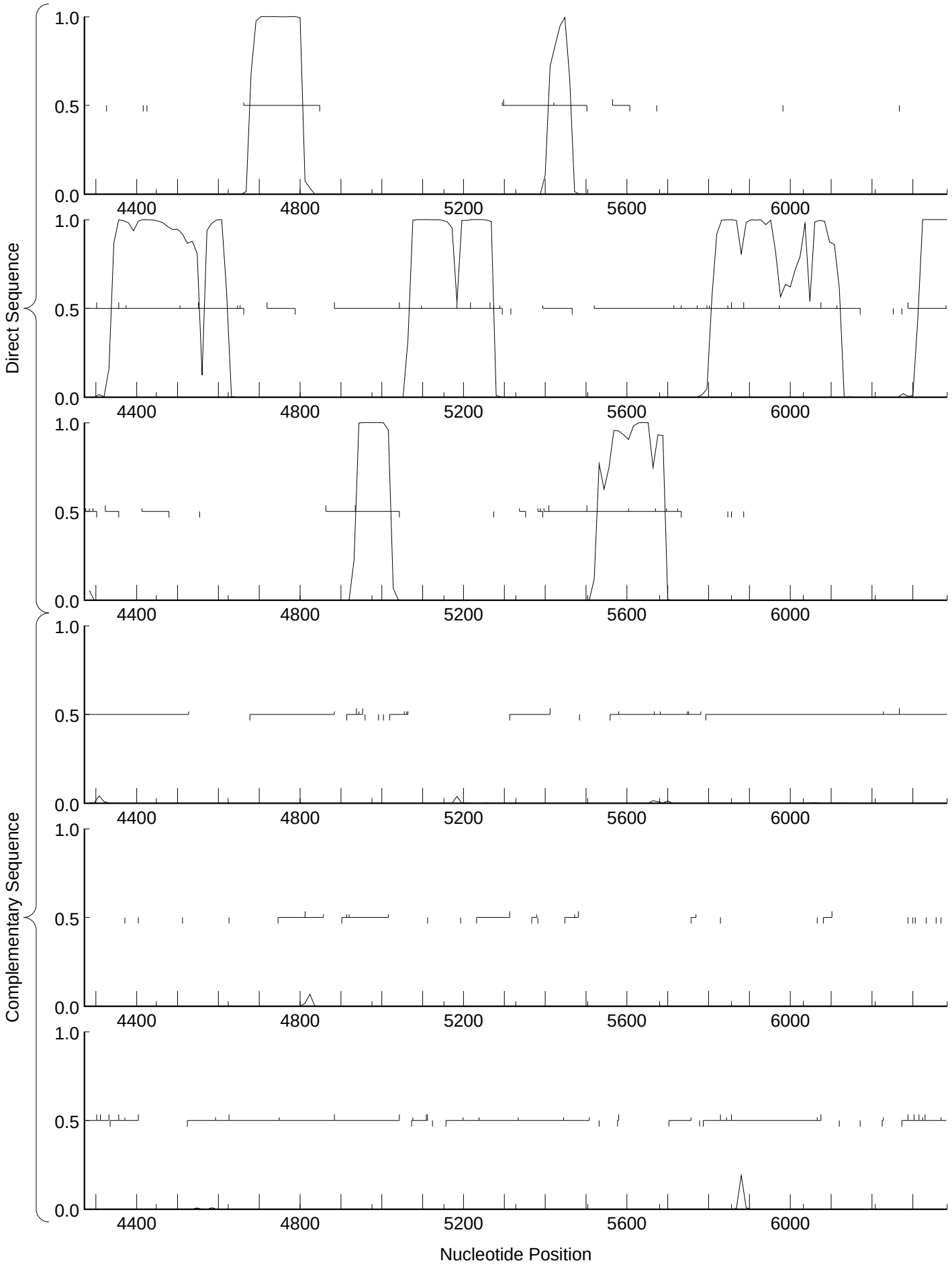
ge OlinDD complete sequence, 63123 bp including 3158 bp direct terminal repeat, Cluster ED, Order 4, Window 96, Step 12, 2/31

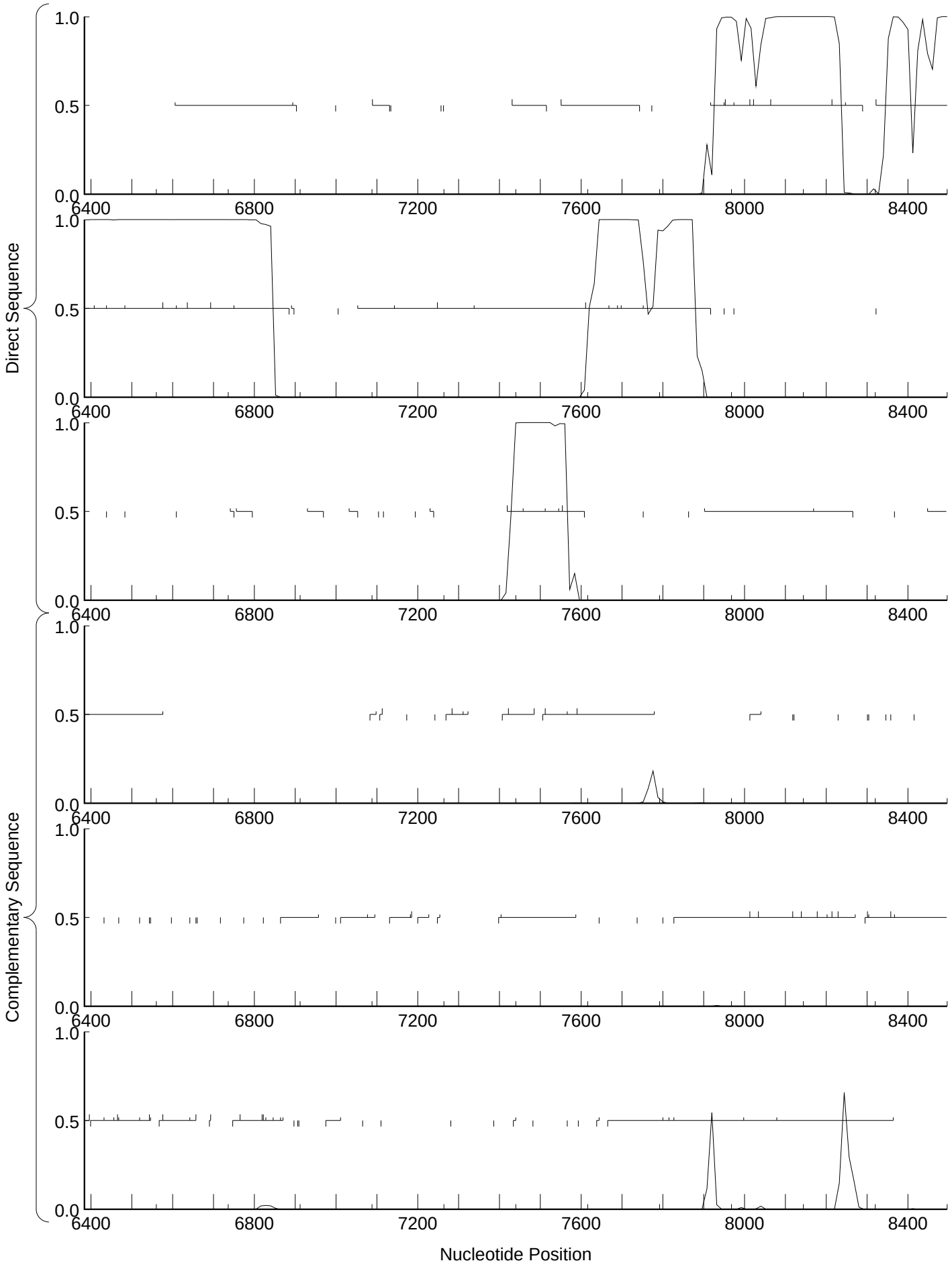


ge OlinDD complete sequence, 63123 bp including 3158 bp direct terminal repeat, Cluster ED, Order 4, Window 96, Step 12, 3/31

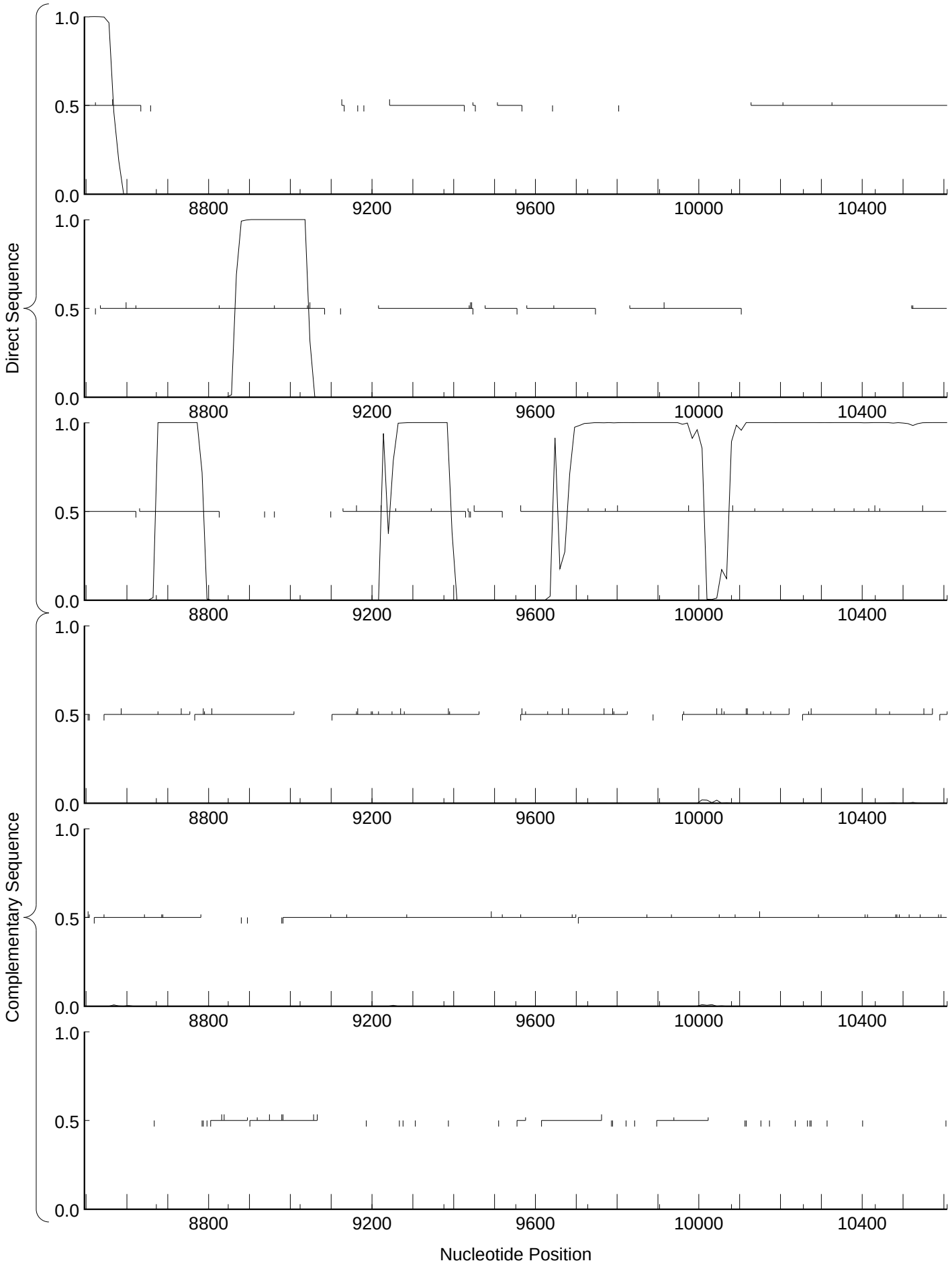


ge OlinDD complete sequence, 63123 bp including 3158 bp direct terminal repeat, Cluster ED, Order 4, Window 96, Step 12, 4/31

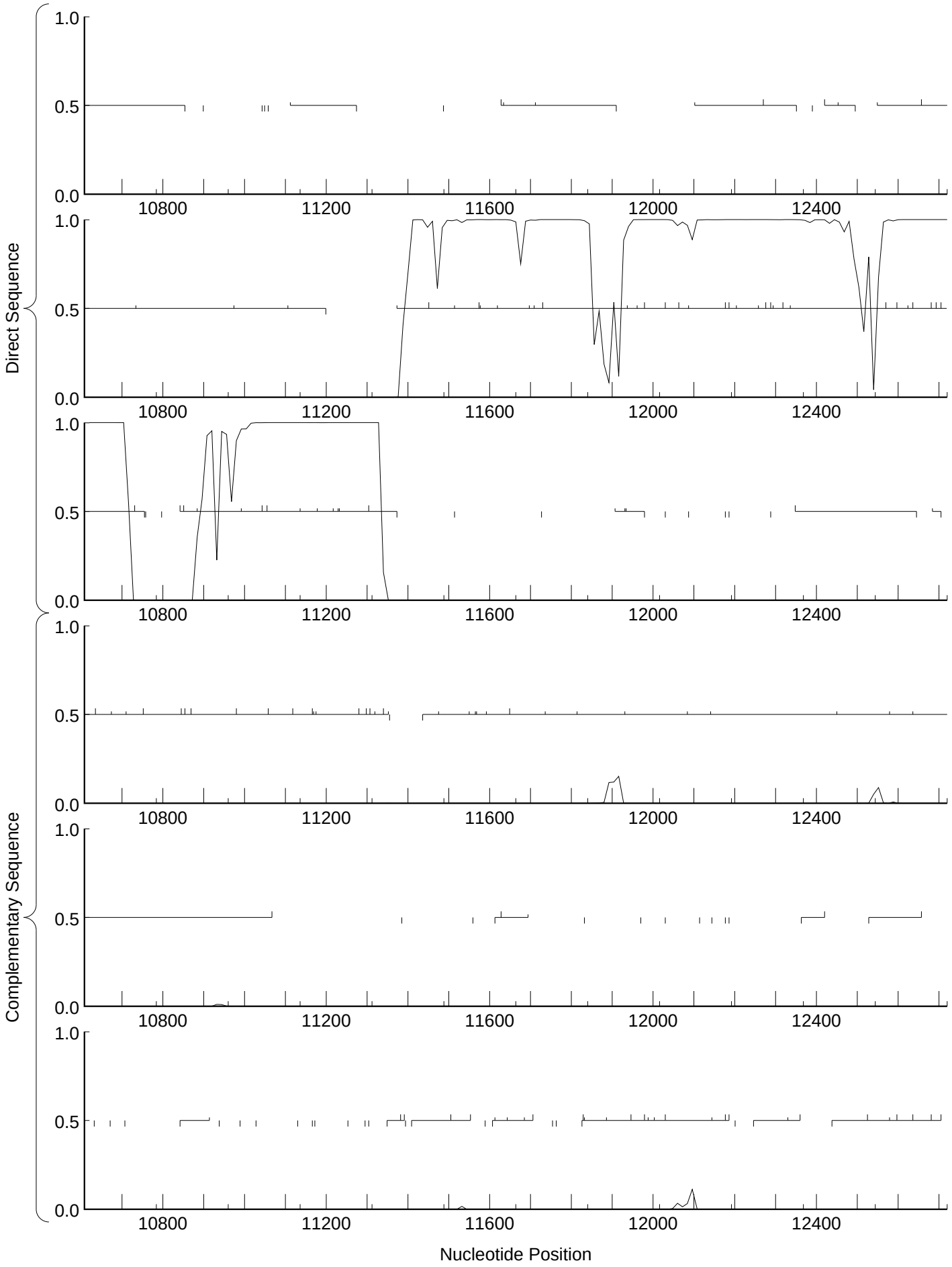


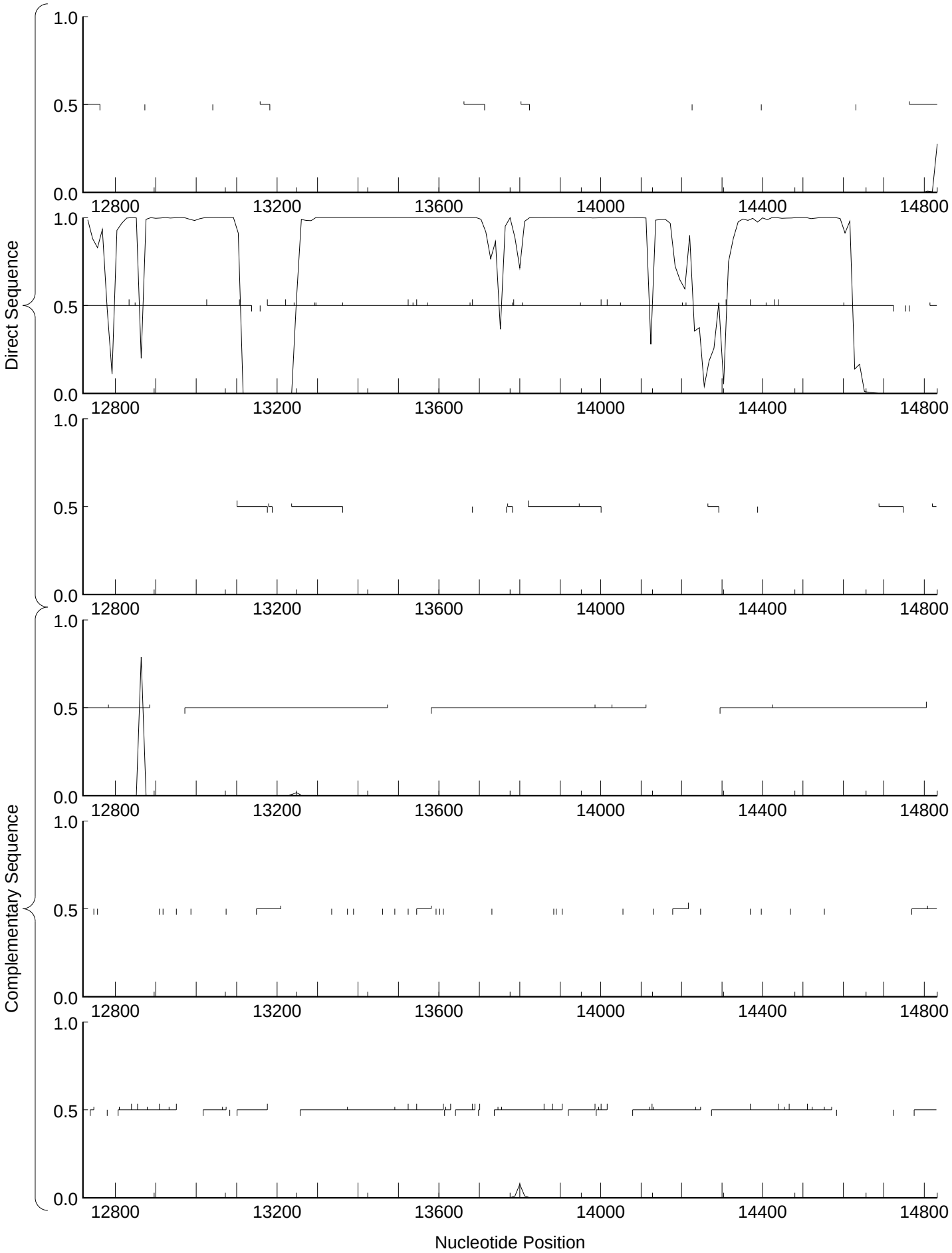

Direct Sequence
Complementary Sequence
1.0
0.5
0.0
6400
6800
7200
7600
8000
8400
Nucleotide Position

ge OlinDD complete sequence, 63123 bp including 3158 bp direct terminal repeat, Cluster ED, Order 4, Window 96, Step 12, 6/31

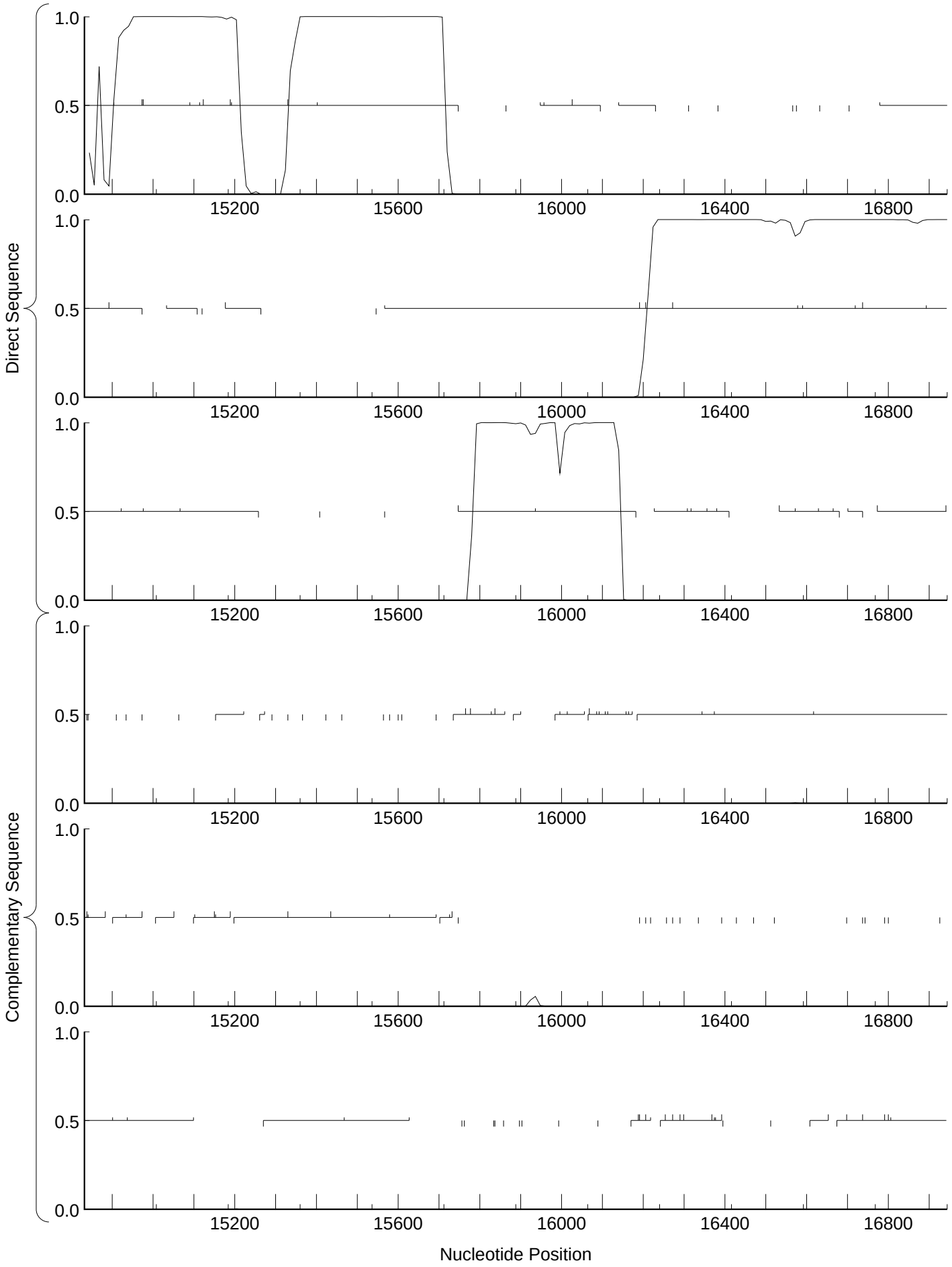


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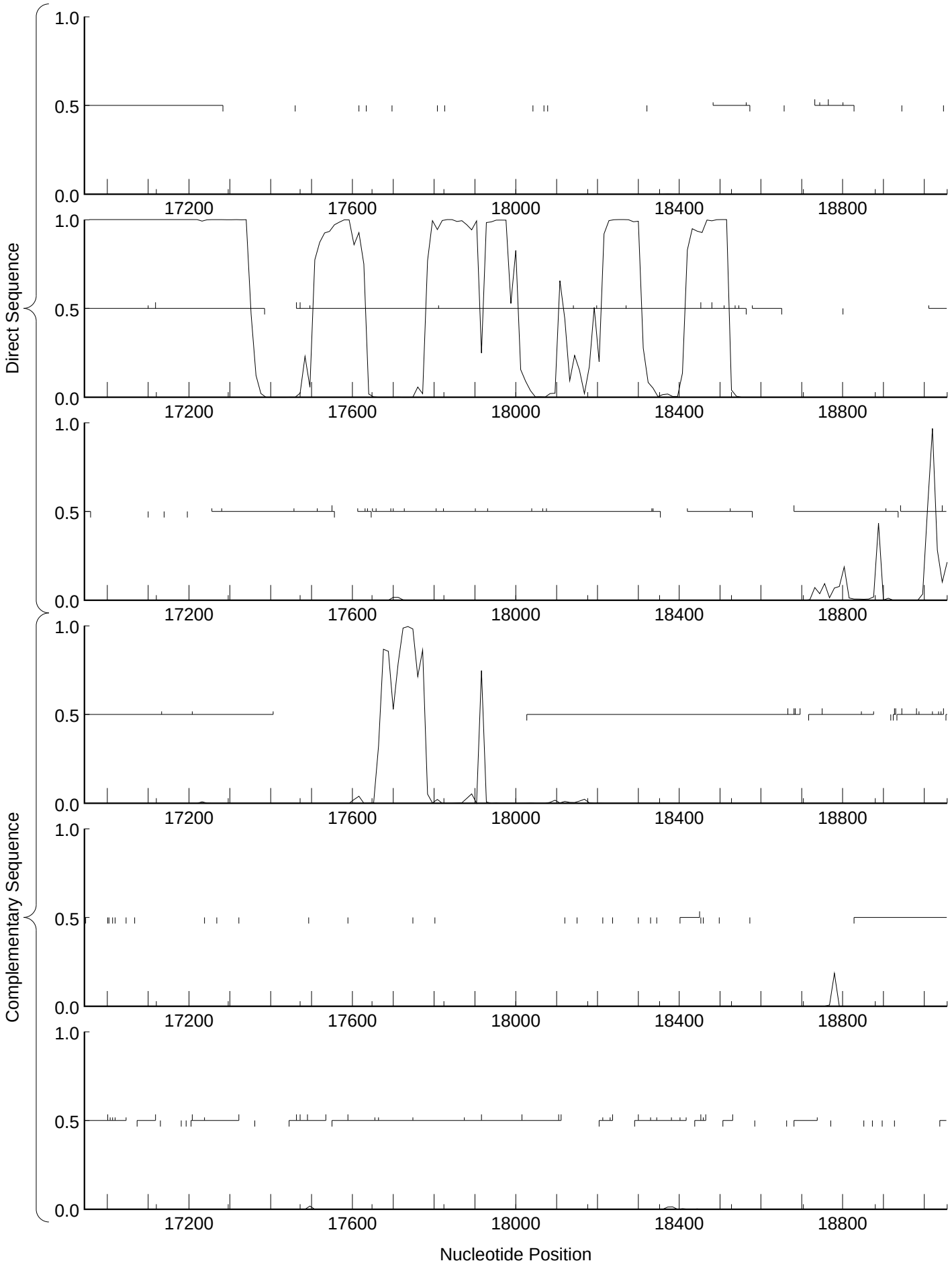



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Complementary Sequence
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0.5
0.0
12800
13200
13600
14000
14400
14800
Nucleotide Position

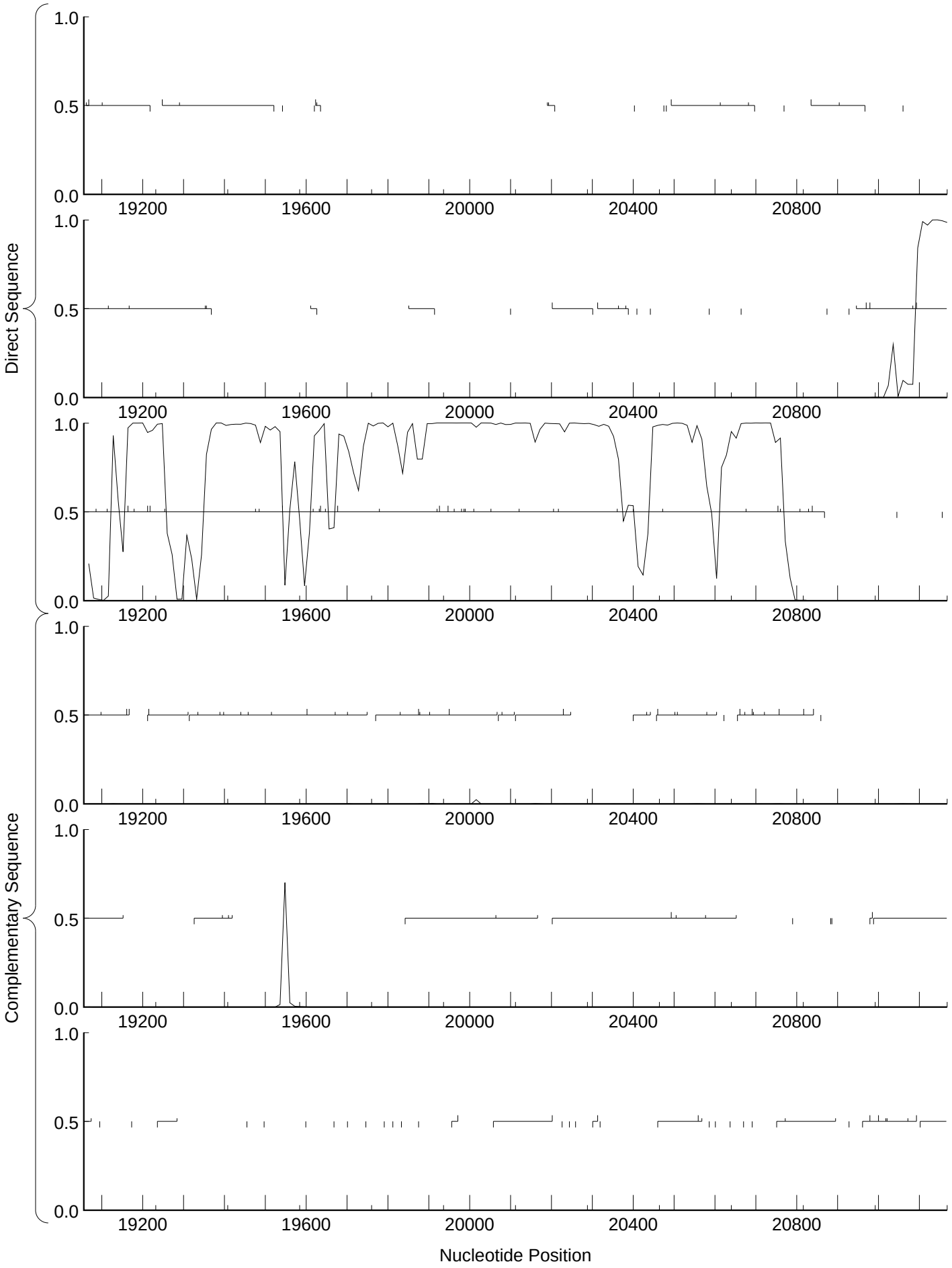
ge OlinDD complete sequence, 63123 bp including 3158 bp direct terminal repeat, Cluster ED, Order 4, Window 96, Step 12, 9/31



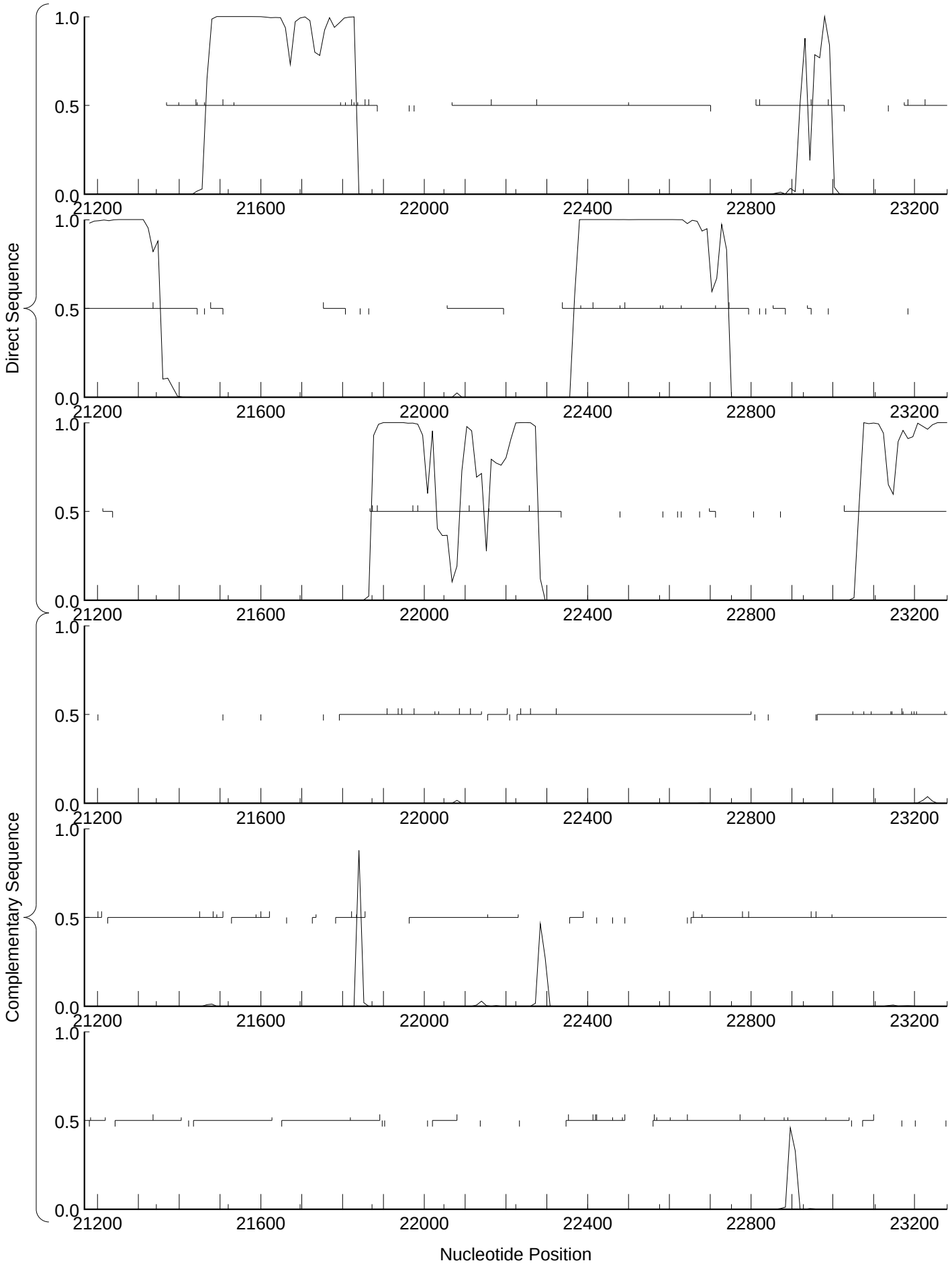
e OlinDD complete sequence, 63123 bp including 3158 bp direct terminal repeat, Cluster ED, Order 4, Window 96, Step 12, 10/31



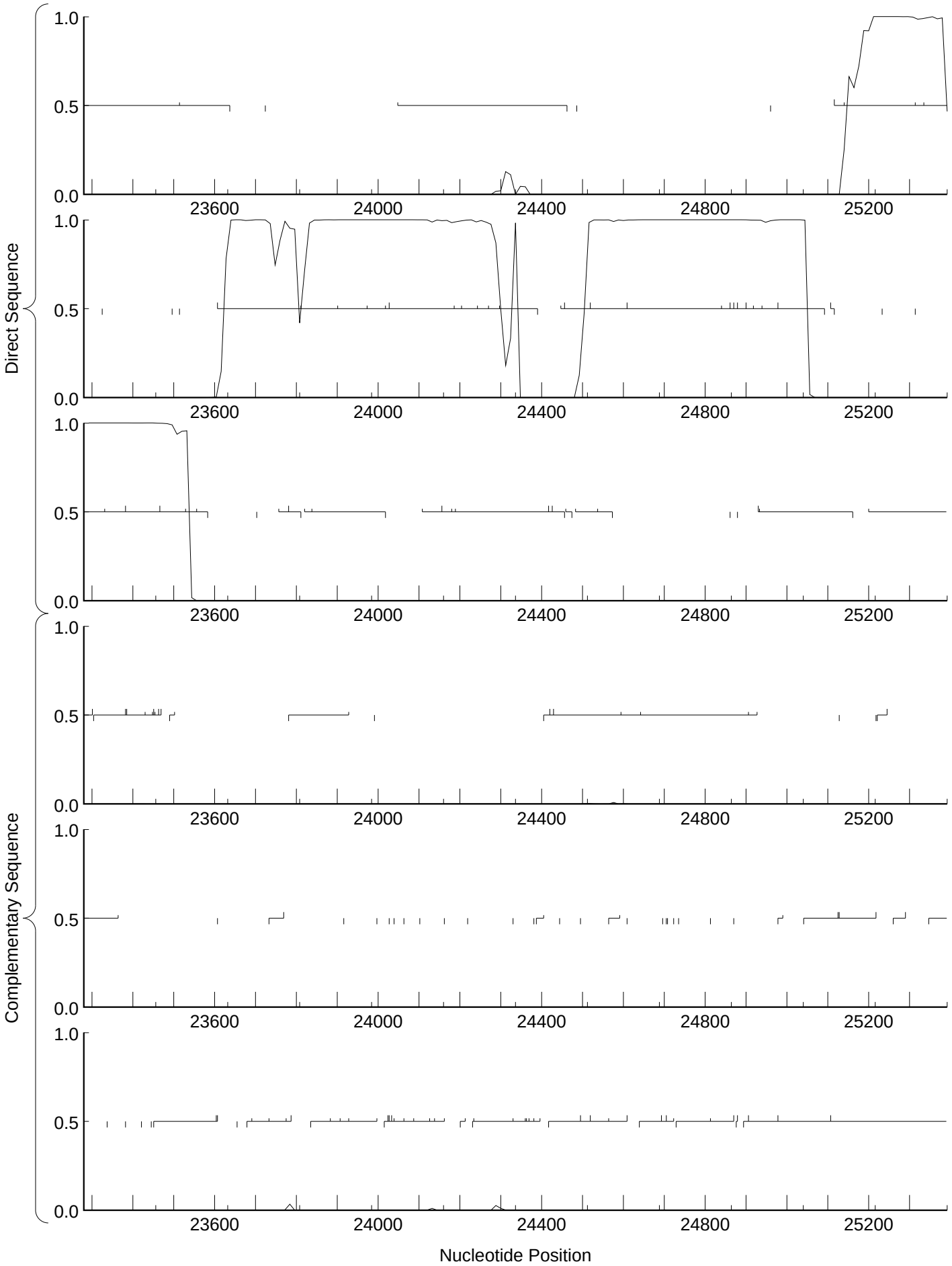
e OlinDD complete sequence, 63123 bp including 3158 bp direct terminal repeat, Cluster ED, Order 4, Window 96, Step 12, 11/31



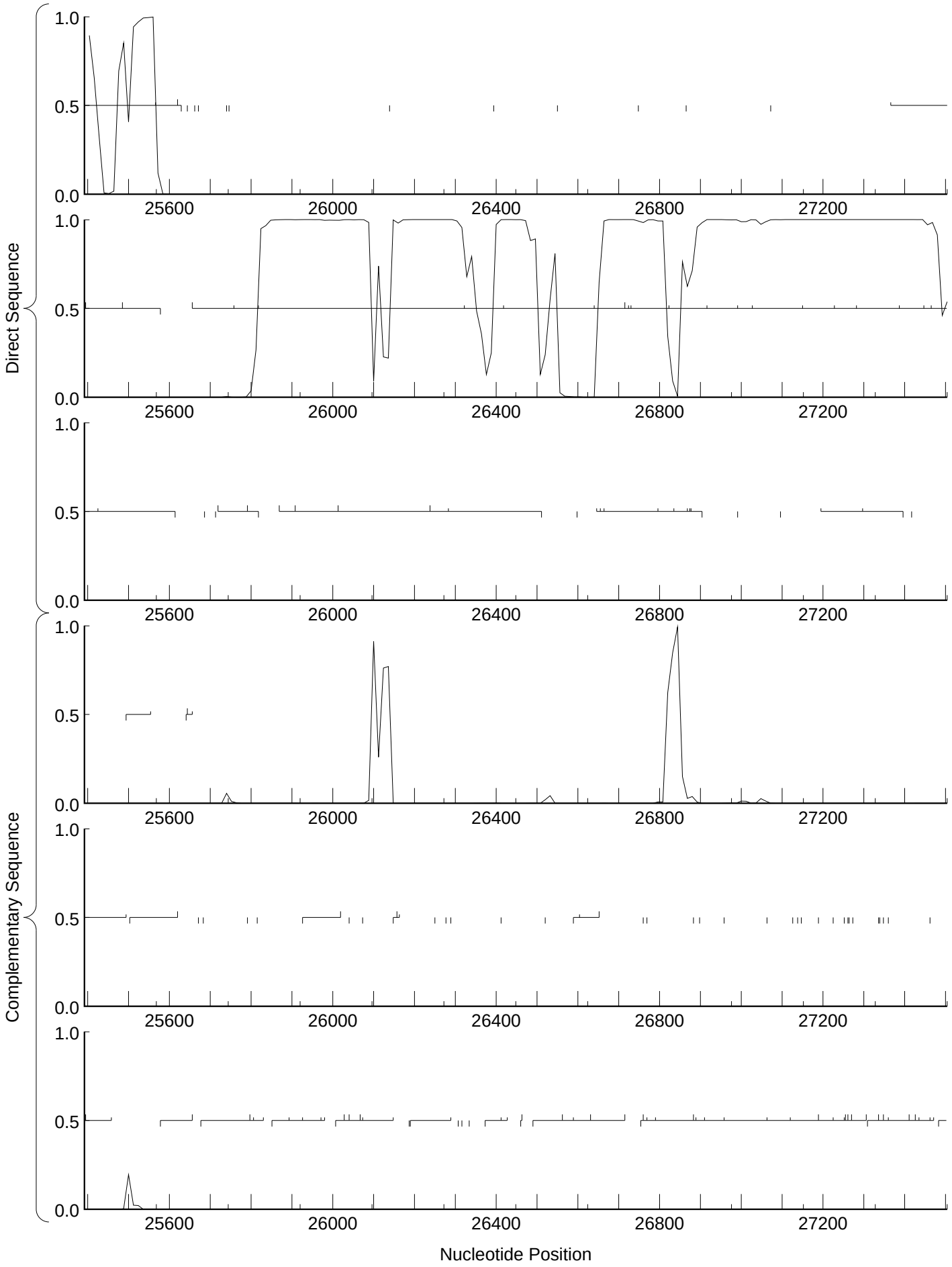
e OlinDD complete sequence, 63123 bp including 3158 bp direct terminal repeat, Cluster ED, Order 4, Window 96, Step 12, 12/31



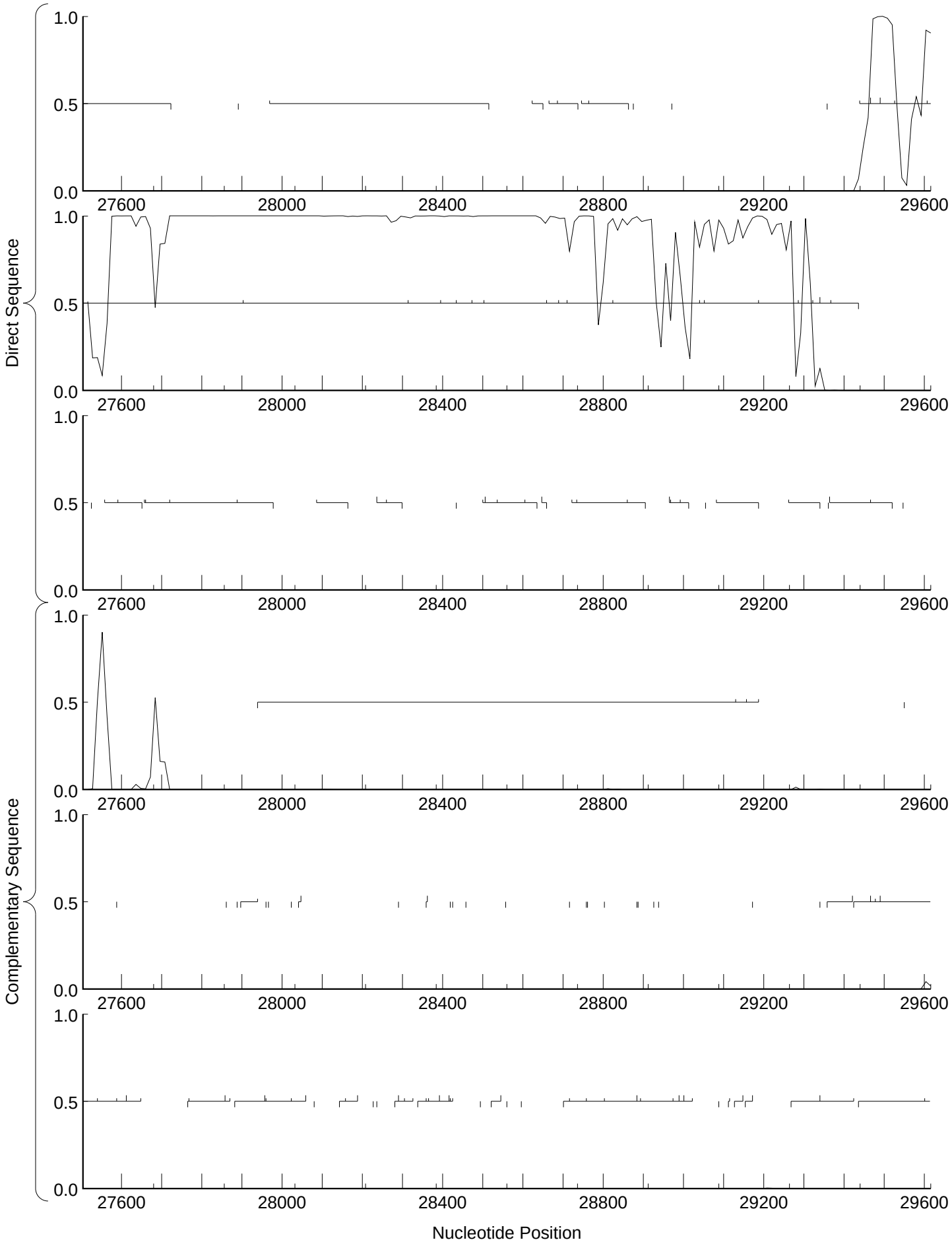
e OlinDD complete sequence, 63123 bp including 3158 bp direct terminal repeat, Cluster ED, Order 4, Window 96, Step 12, 13/31



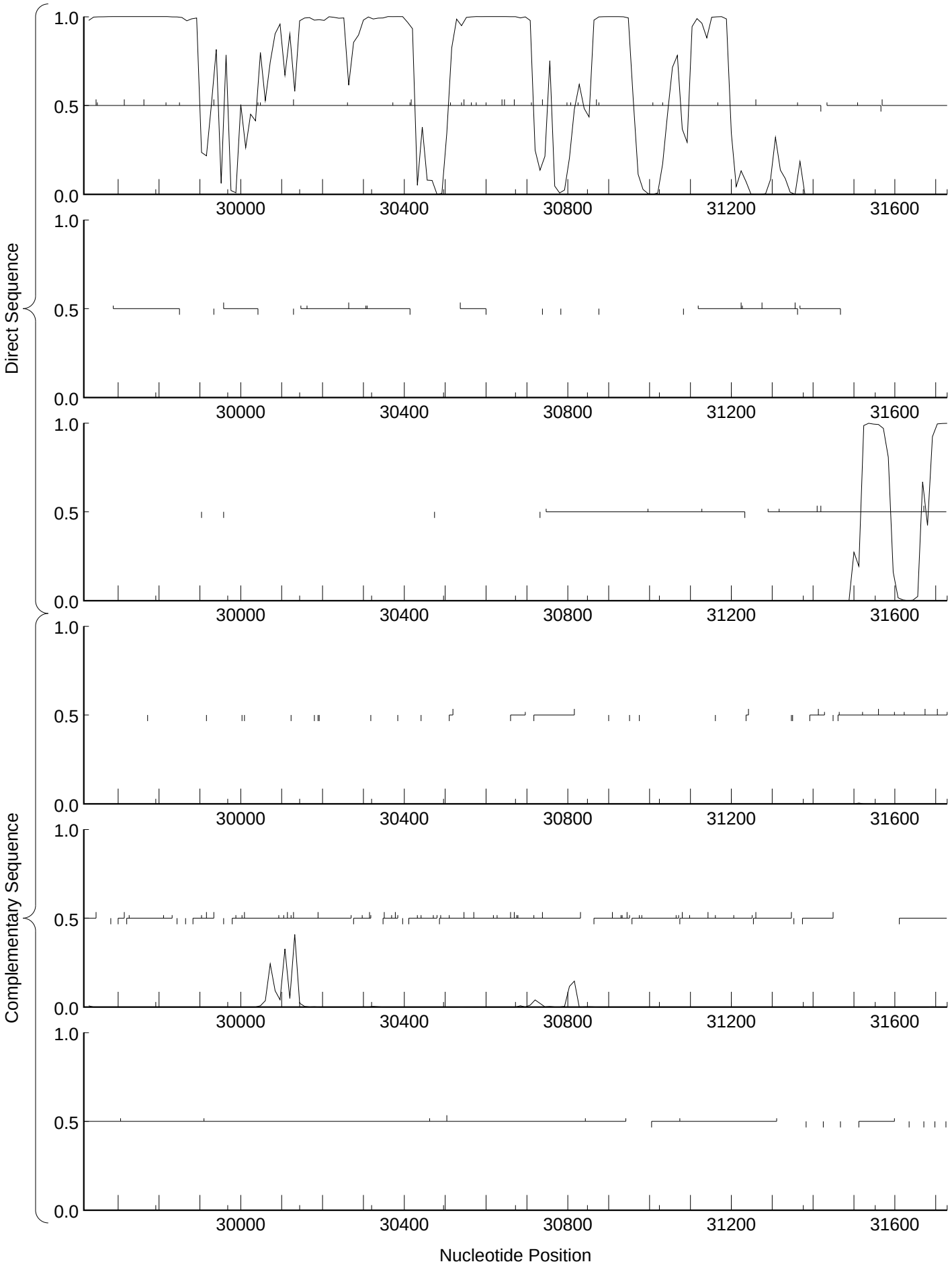
e OlinDD complete sequence, 63123 bp including 3158 bp direct terminal repeat, Cluster ED, Order 4, Window 96, Step 12, 14/31



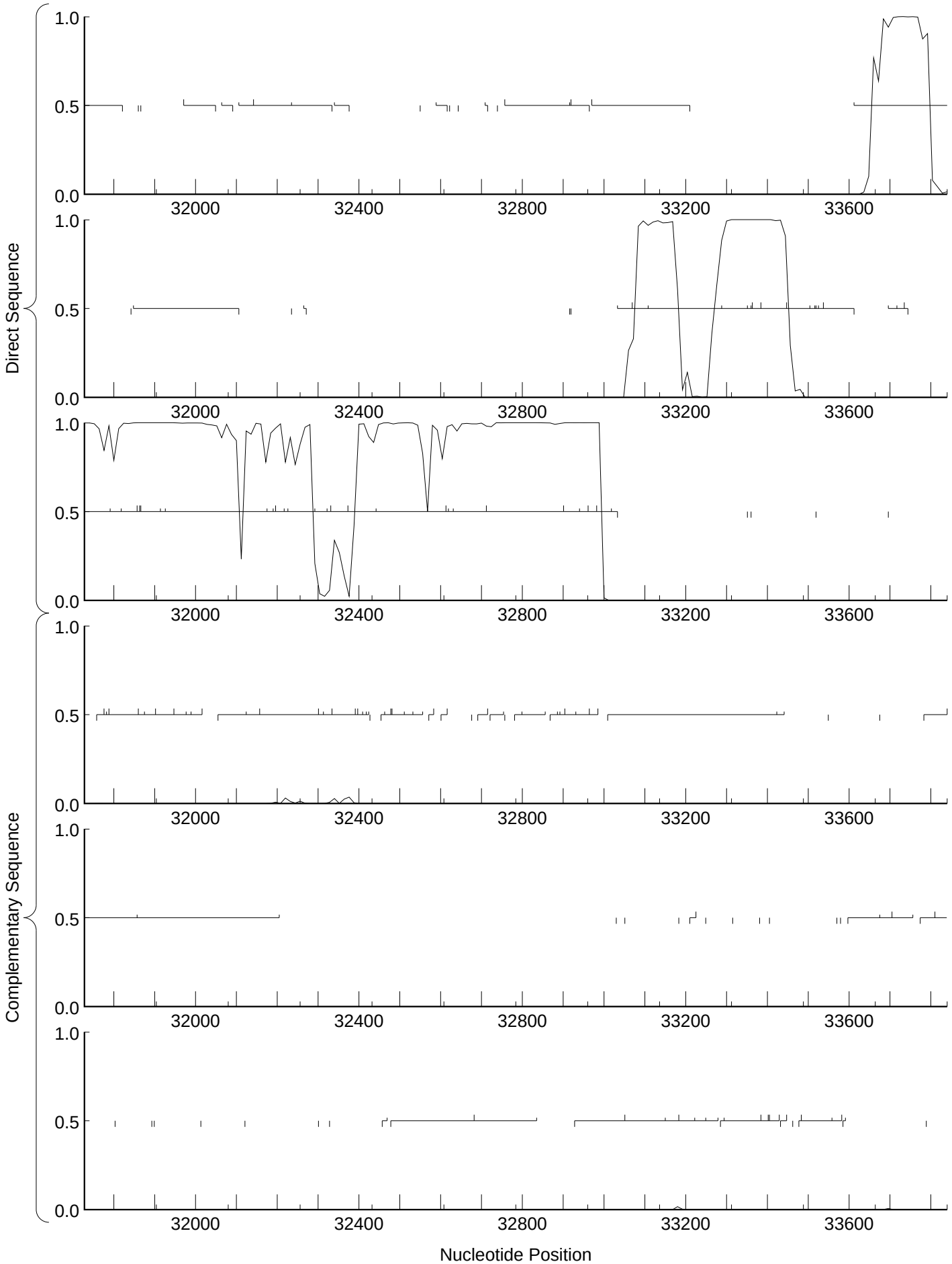
e OlinDD complete sequence, 63123 bp including 3158 bp direct terminal repeat, Cluster ED, Order 4, Window 96, Step 12, 15/31

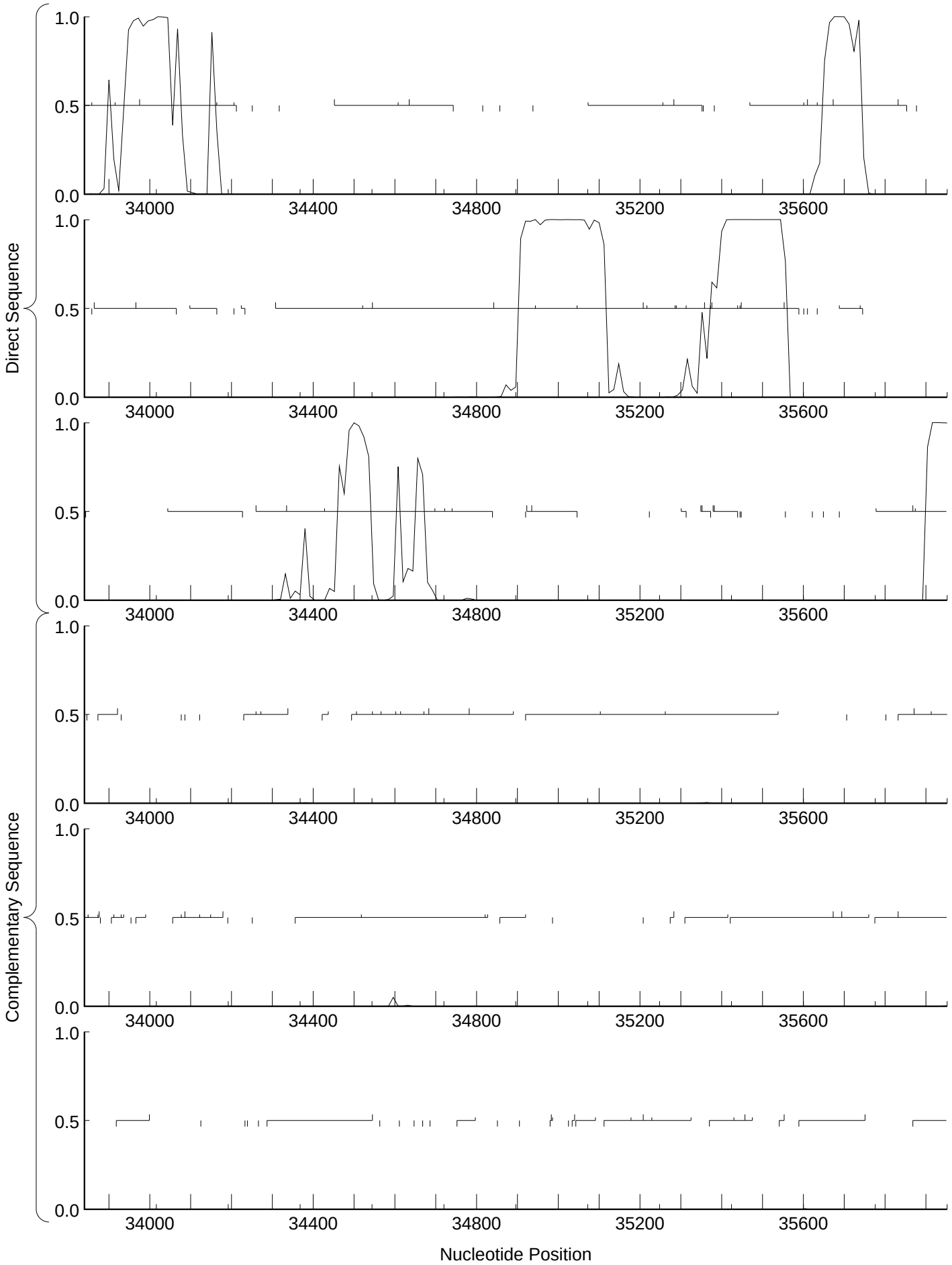


e OlinDD complete sequence, 63123 bp including 3158 bp direct terminal repeat, Cluster ED, Order 4, Window 96, Step 12, 16/31

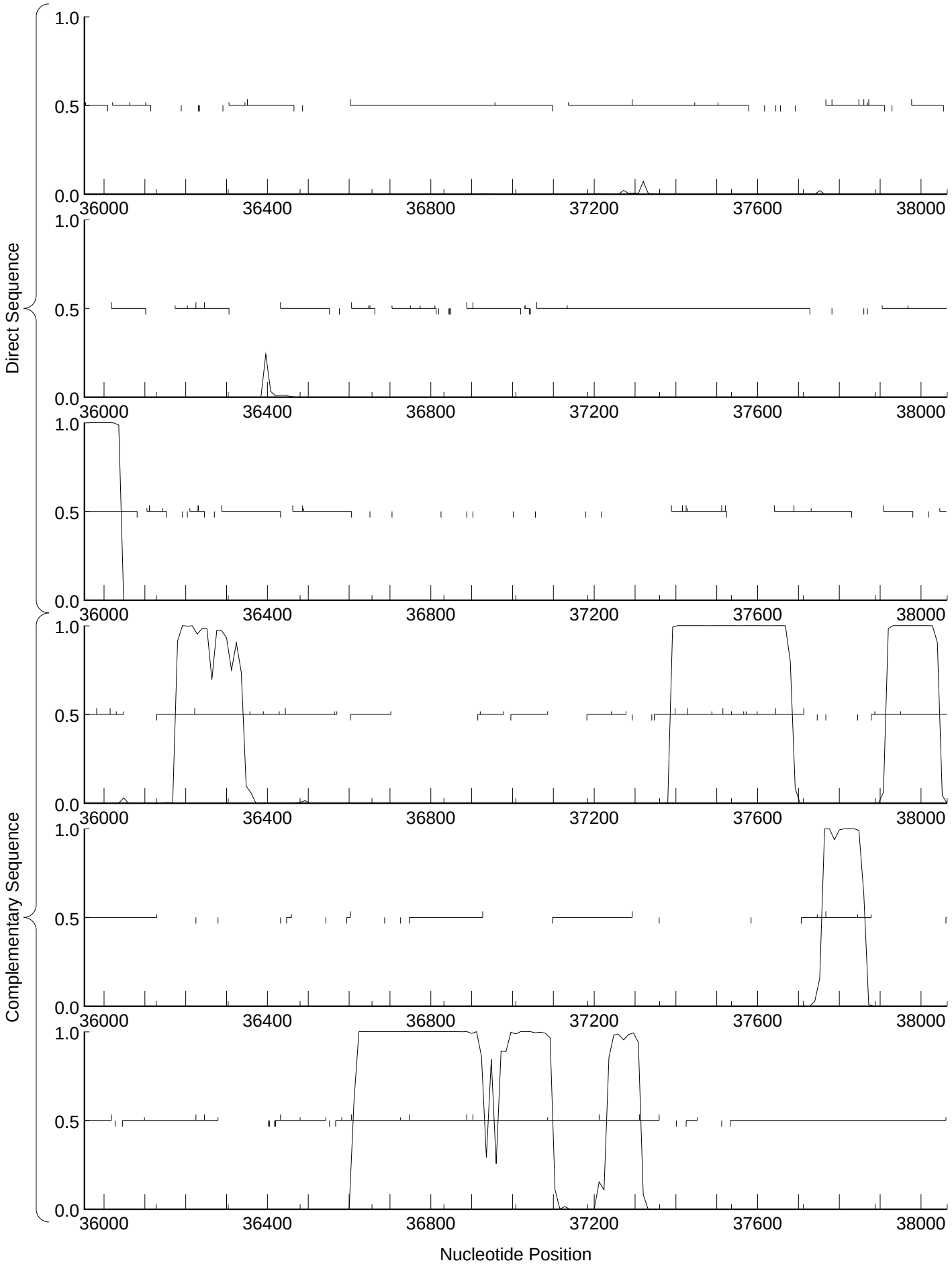


e OlinDD complete sequence, 63123 bp including 3158 bp direct terminal repeat, Cluster ED, Order 4, Window 96, Step 12, 17/31

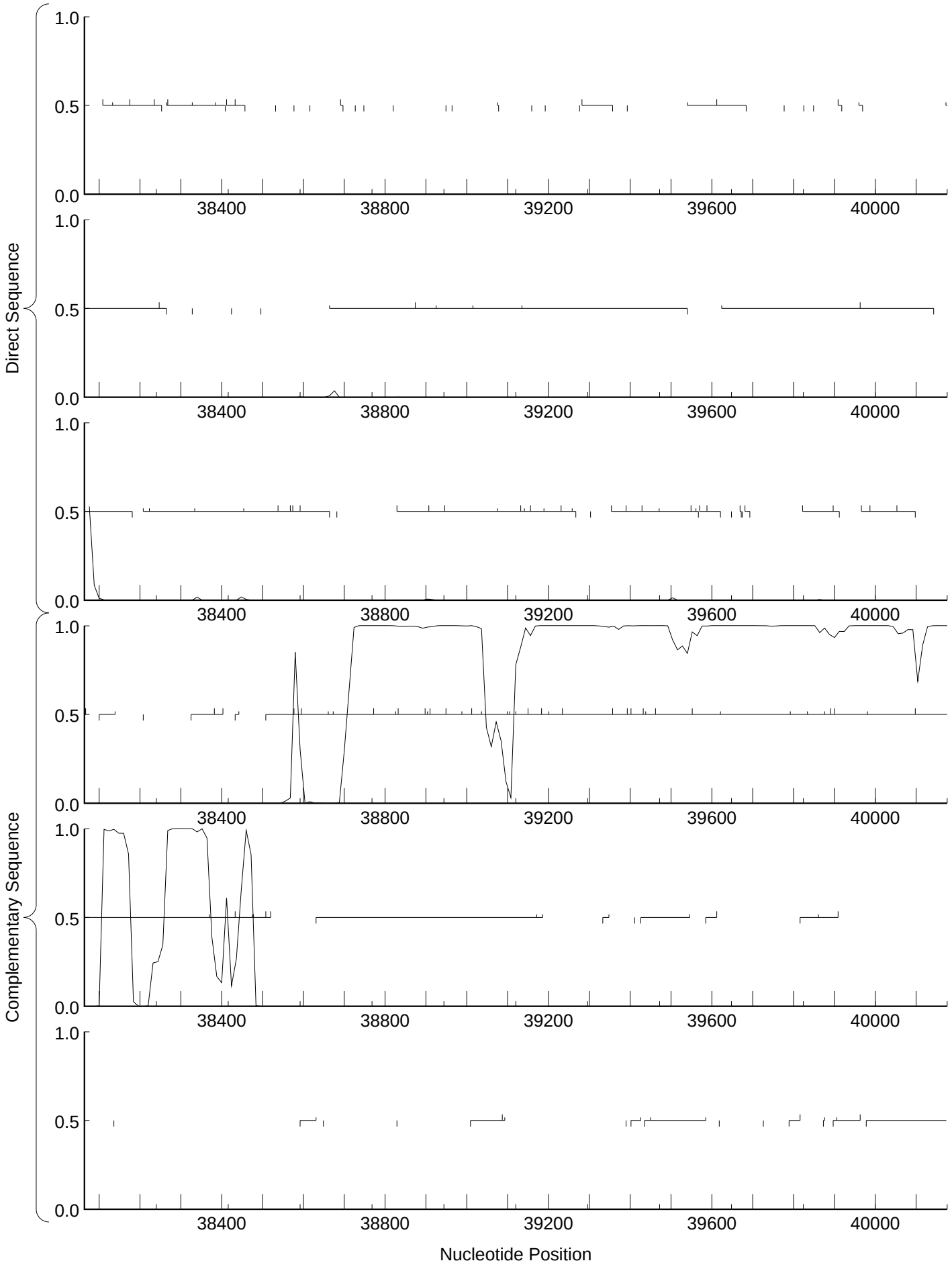


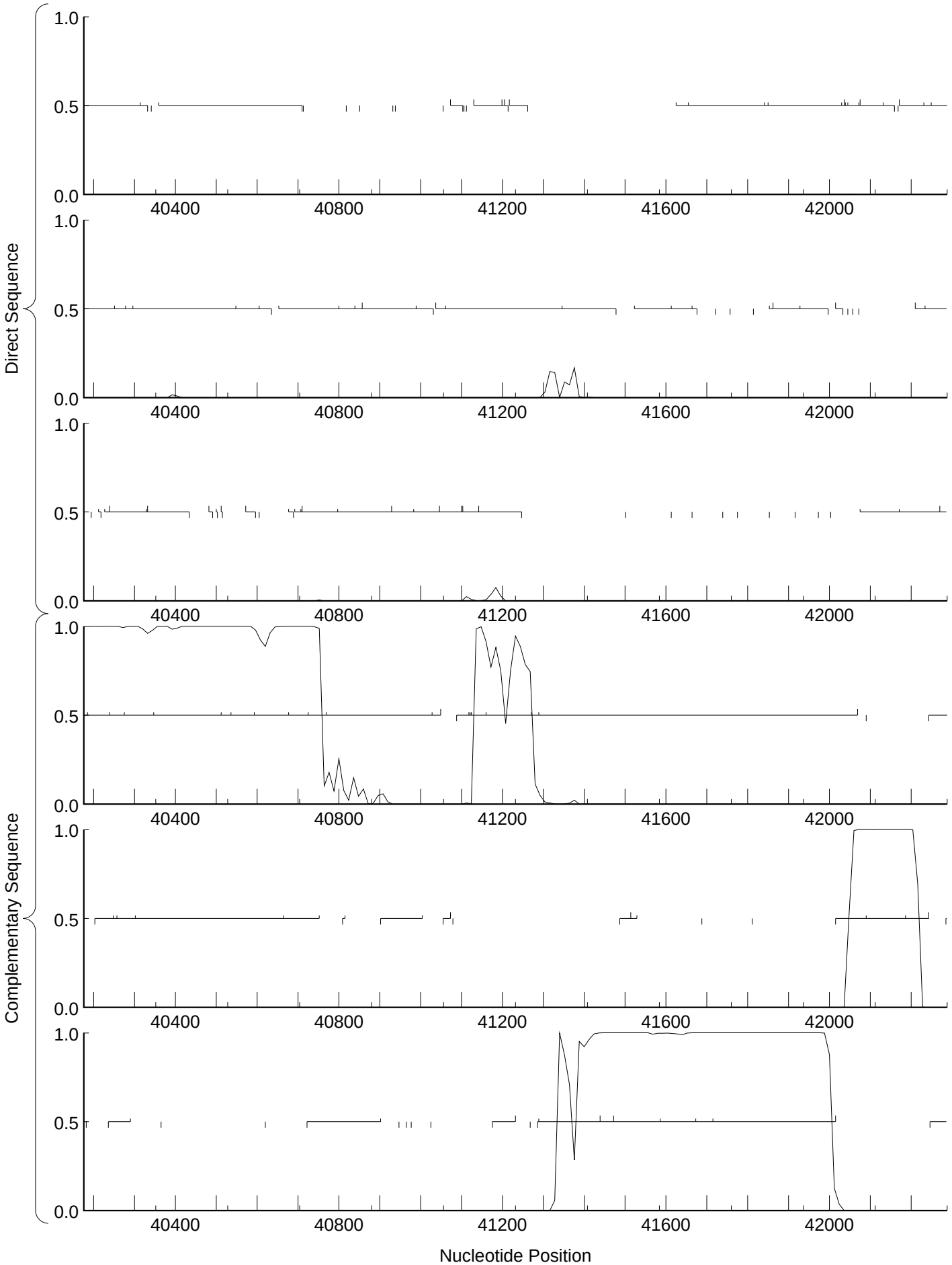

Direct Sequence
Complementary Sequence
1.0
0.5
0.0
34000
34400
34800
35200
35600
Nucleotide Position

e OlinDD complete sequence, 63123 bp including 3158 bp direct terminal repeat, Cluster ED, Order 4, Window 96, Step 12, 19/31

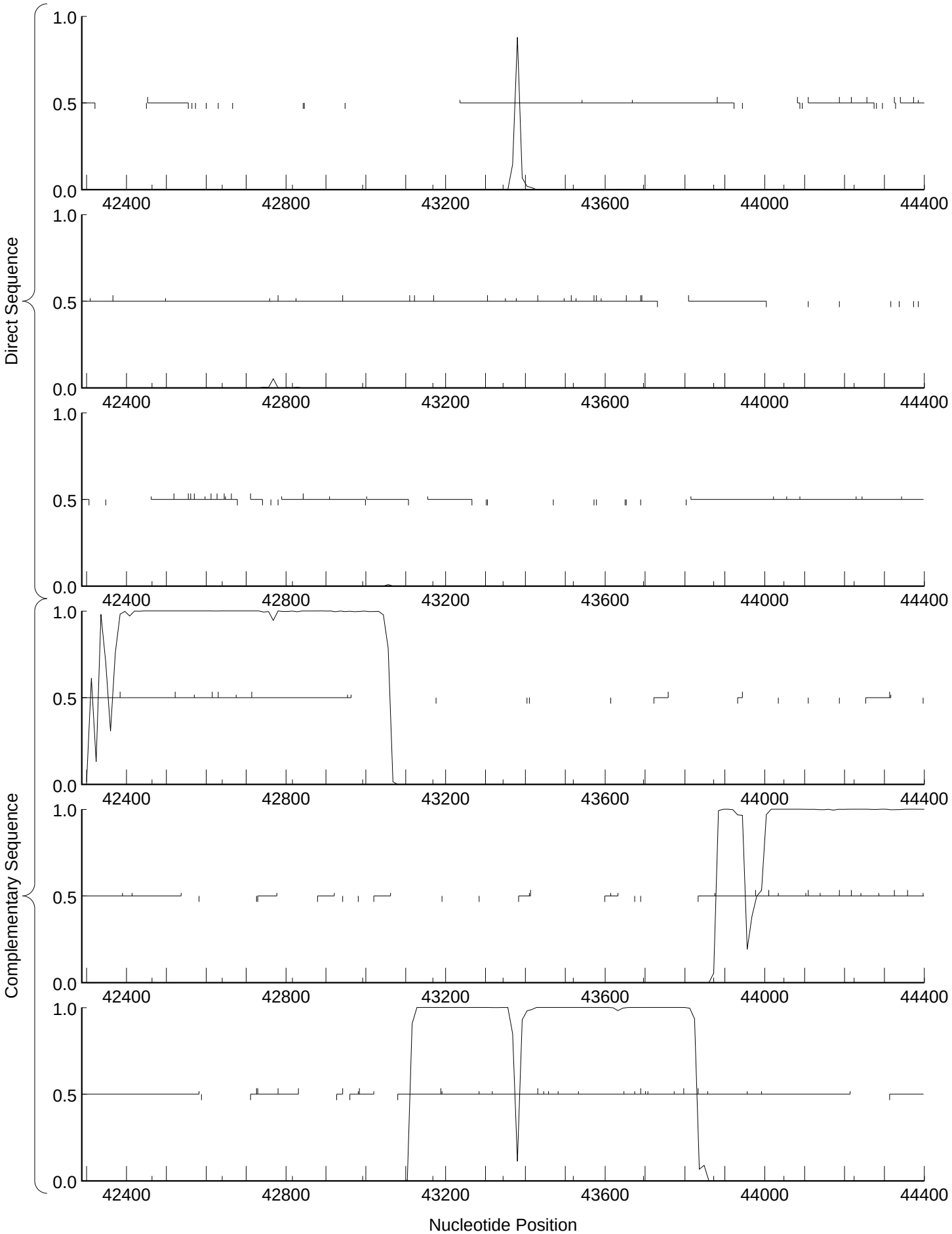


e OlinDD complete sequence, 63123 bp including 3158 bp direct terminal repeat, Cluster ED, Order 4, Window 96, Step 12, 20/31

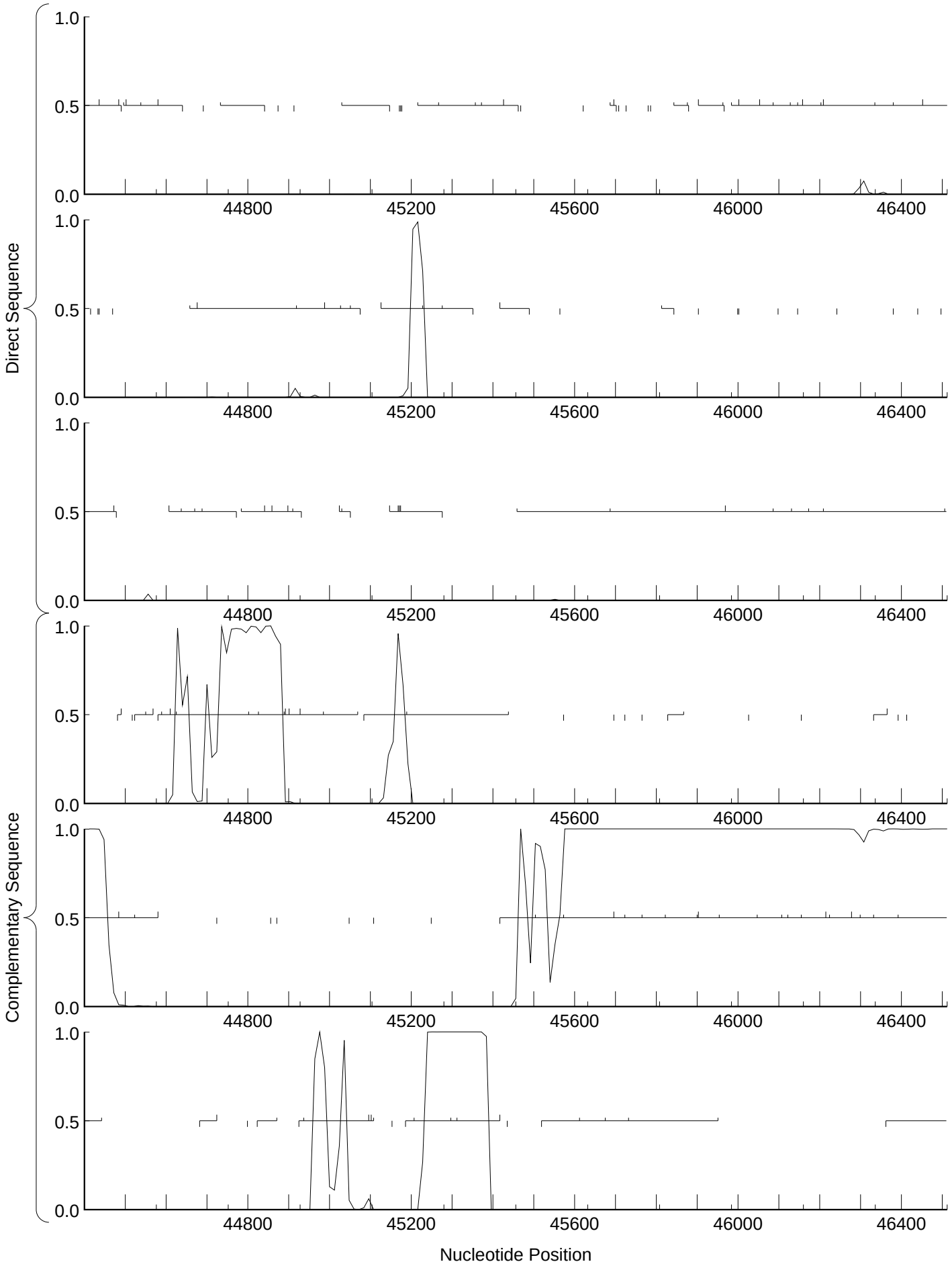



Direct Sequence
Complementary Sequence
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0.5
0.0
40400
40800
41200
41600
42000
Nucleotide Position

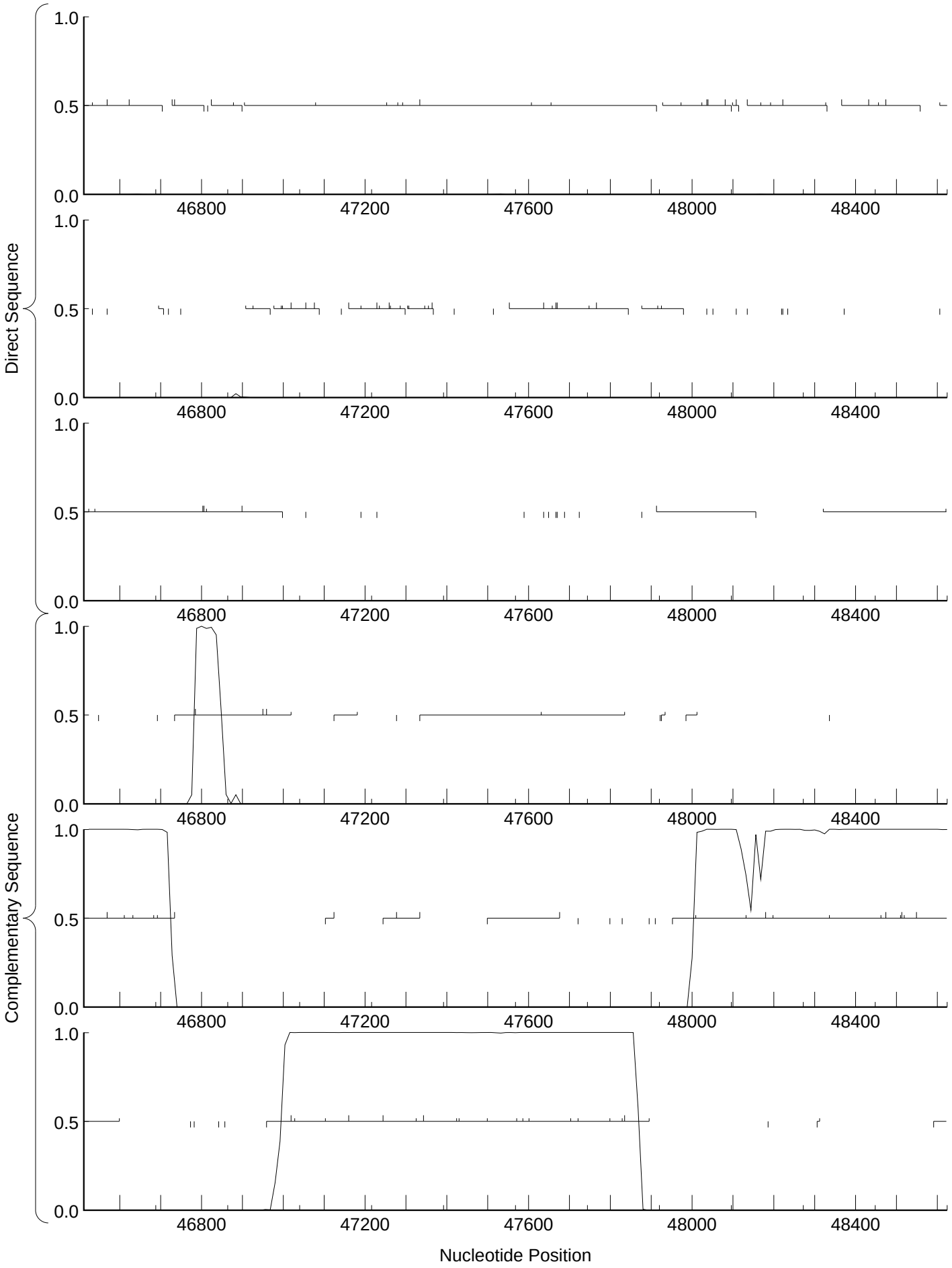
e OlinDD complete sequence, 63123 bp including 3158 bp direct terminal repeat, Cluster ED, Order 4, Window 96, Step 12, 22/31



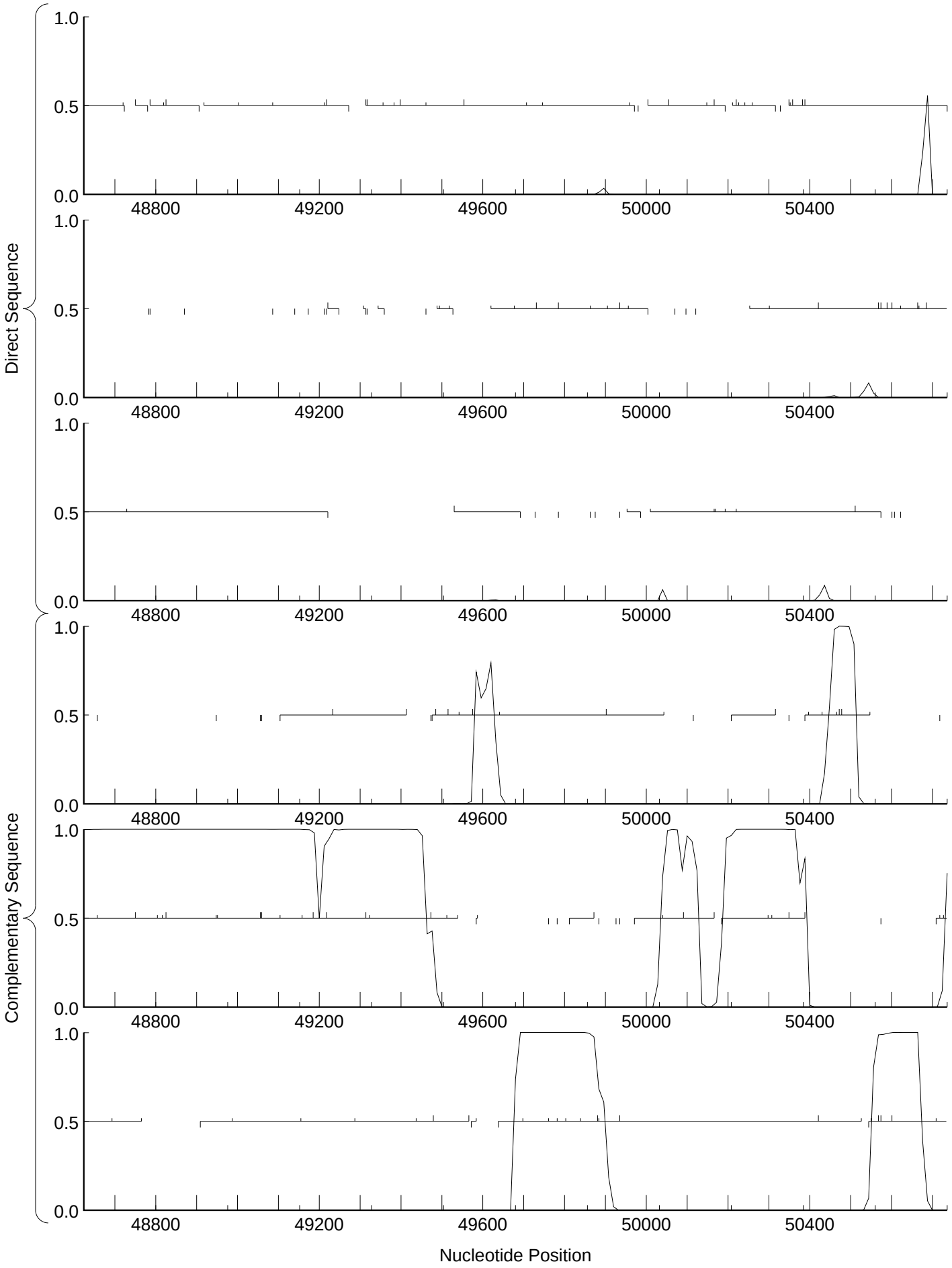
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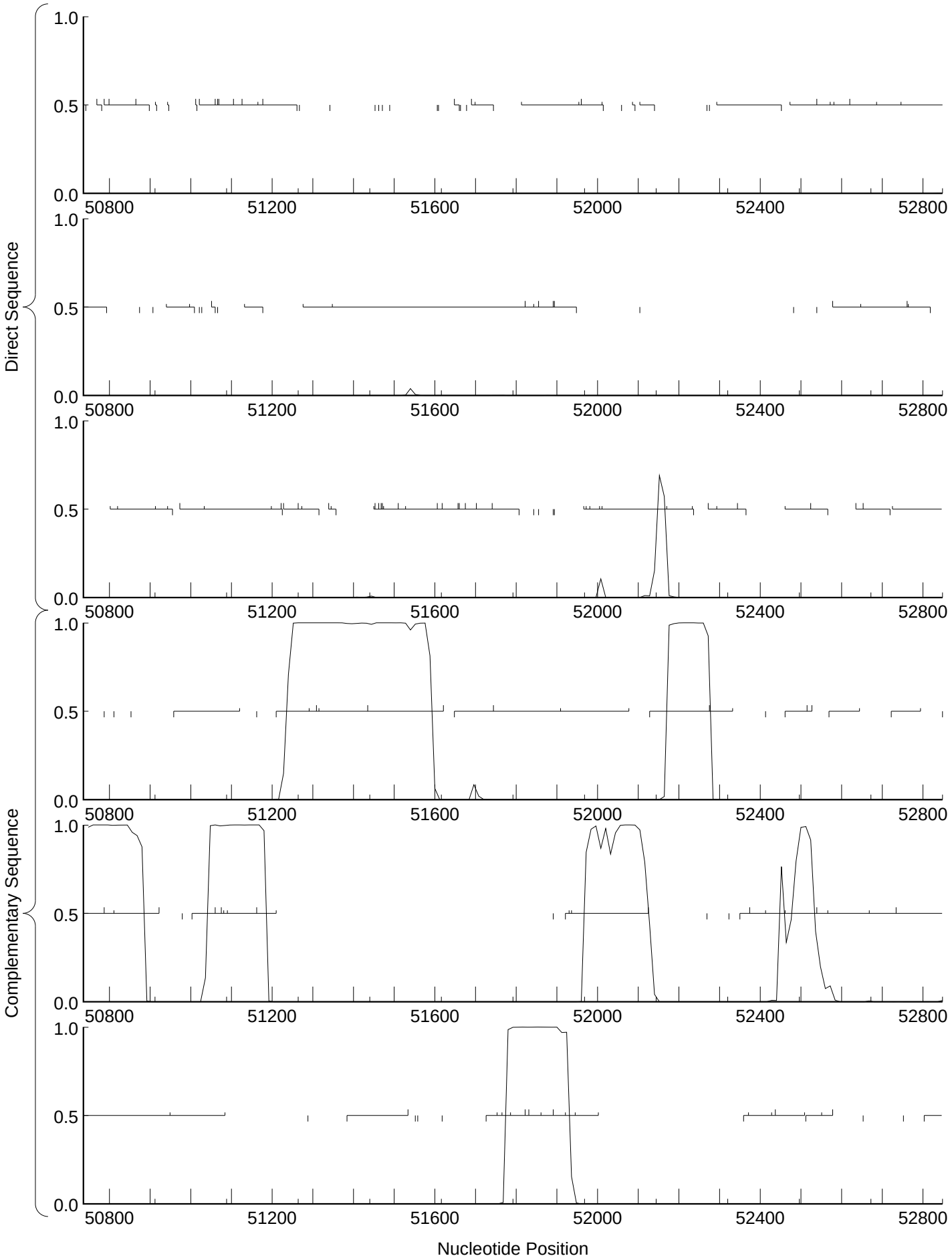


e OlinDD complete sequence, 63123 bp including 3158 bp direct terminal repeat, Cluster ED, Order 4, Window 96, Step 12, 24/31

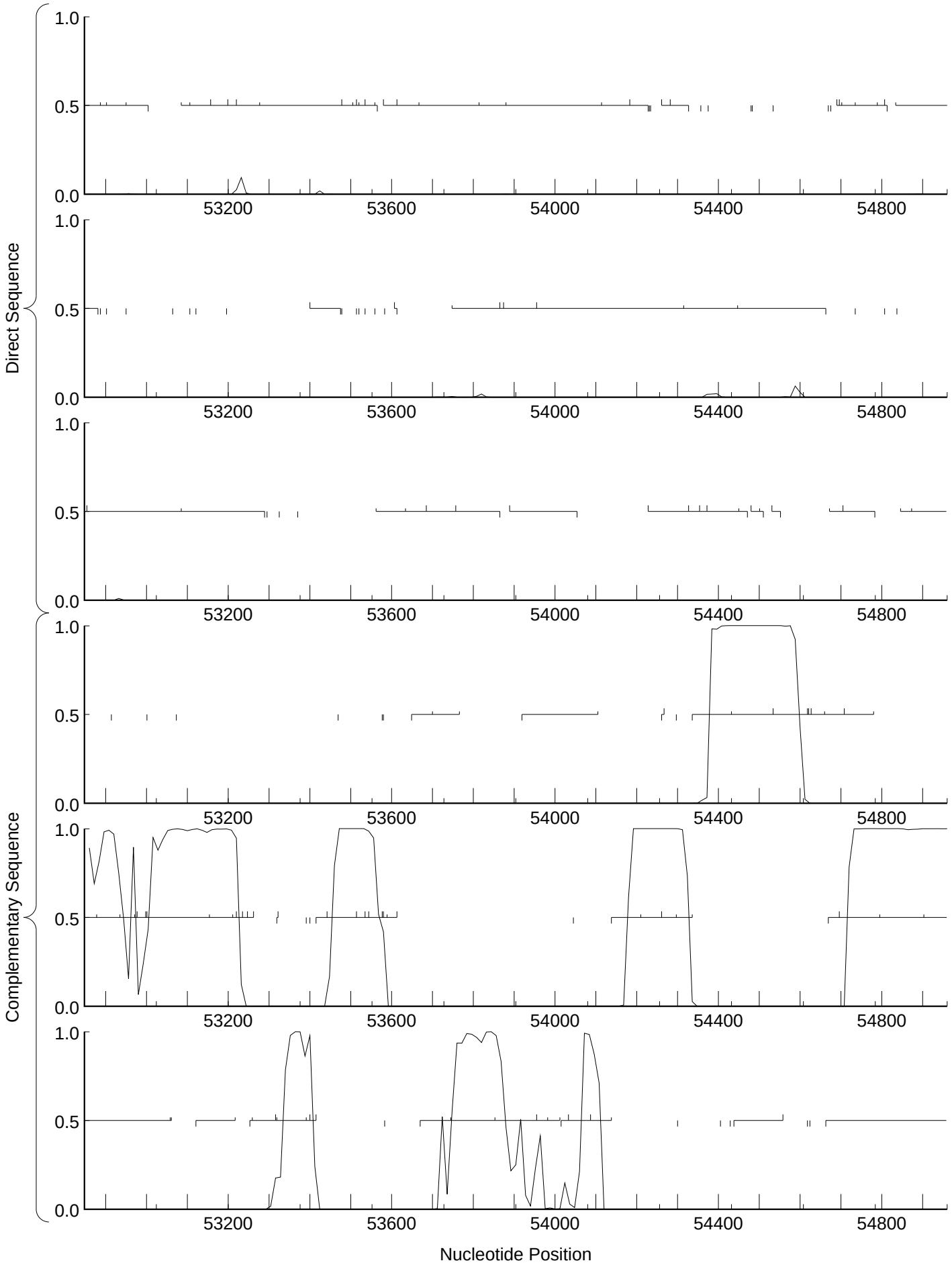


e OlinDD complete sequence, 63123 bp including 3158 bp direct terminal repeat, Cluster ED, Order 4, Window 96, Step 12, 25/31

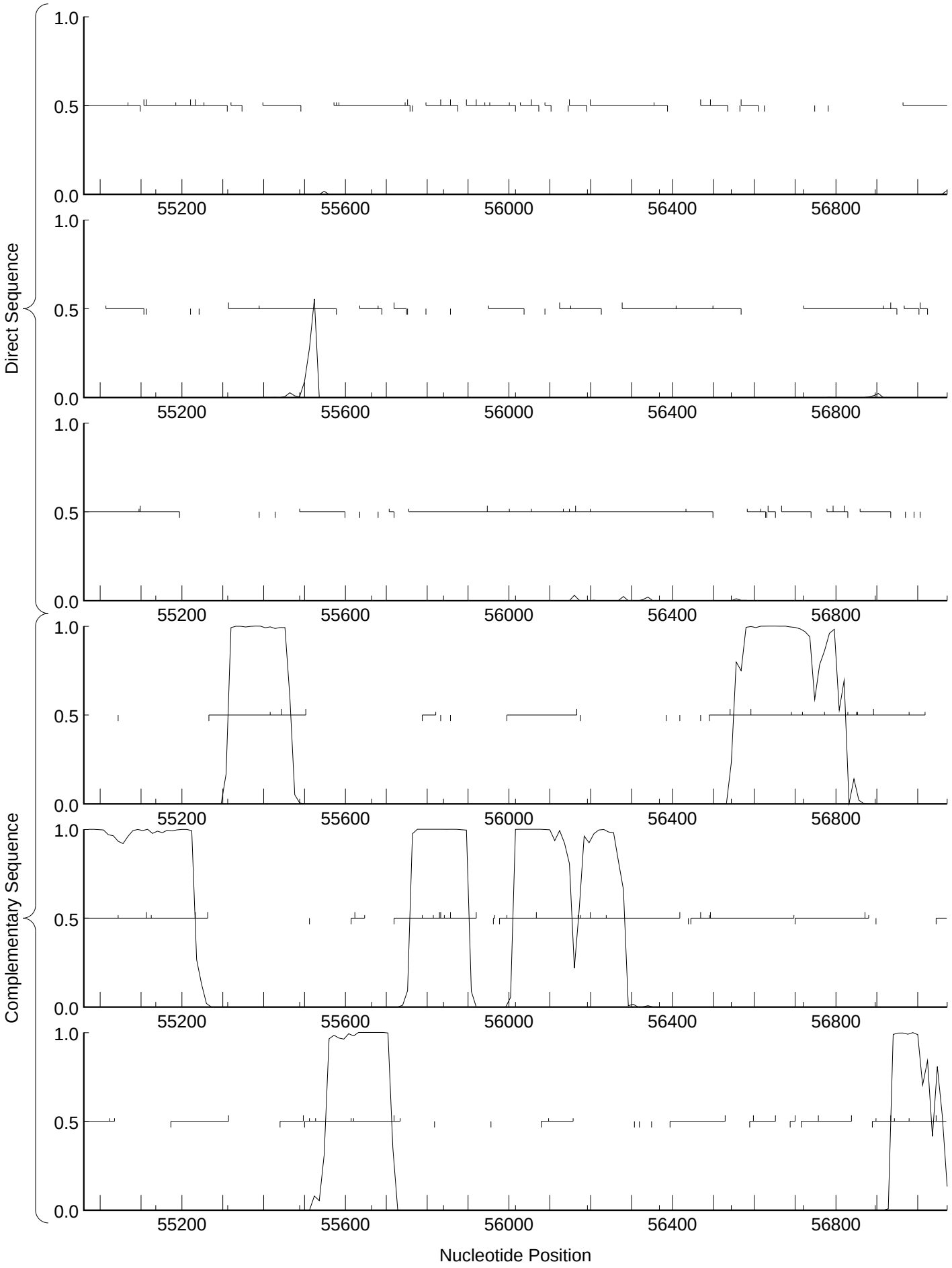



Direct Sequence
Complementary Sequence
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0.5
0.0
50800
51200
51600
52000
52400
52800
Nucleotide Position

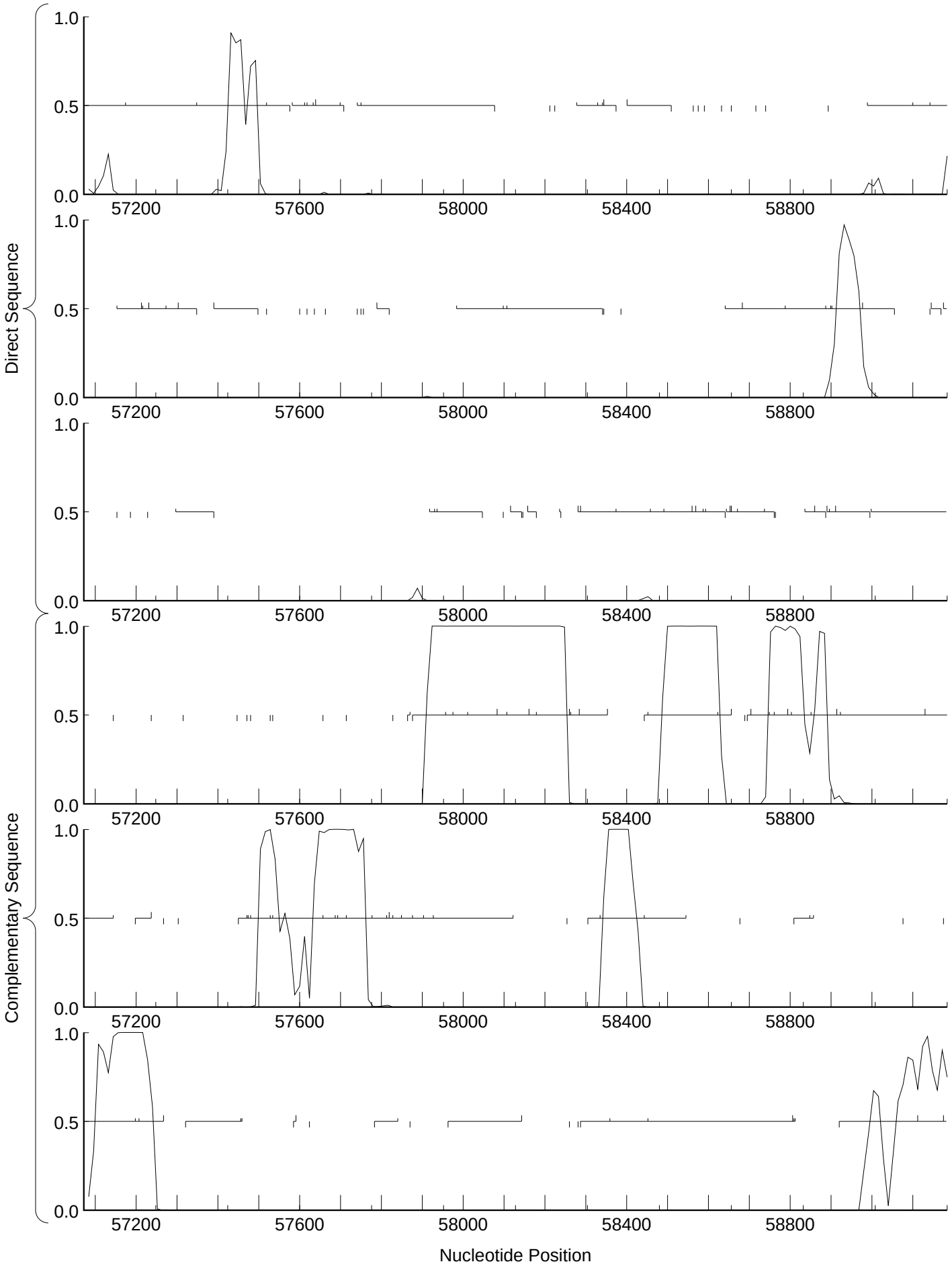
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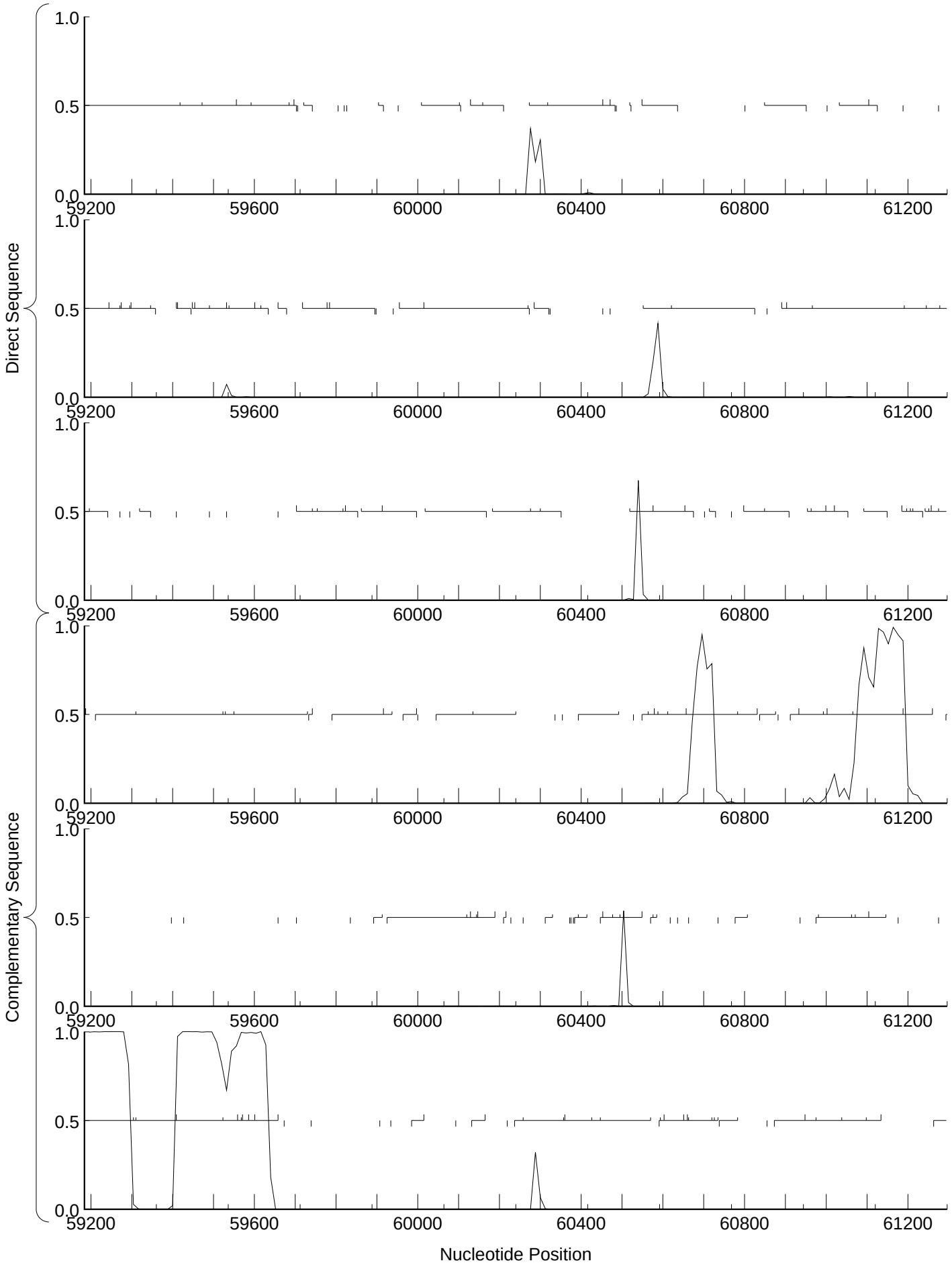
e OlinDD complete sequence, 63123 bp including 3158 bp direct terminal repeat, Cluster ED, Order 4, Window 96, Step 12, 28/31



e OlinDD complete sequence, 63123 bp including 3158 bp direct terminal repeat, Cluster ED, Order 4, Window 96, Step 12, 29/31



e OlinDD complete sequence, 63123 bp including 3158 bp direct terminal repeat, Cluster ED, Order 4, Window 96, Step 12, 30/31



e OlinDD complete sequence, 63123 bp including 3158 bp direct terminal repeat, Cluster ED, Order 4, Window 96, Step 12, 31/31

