GeneMark

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

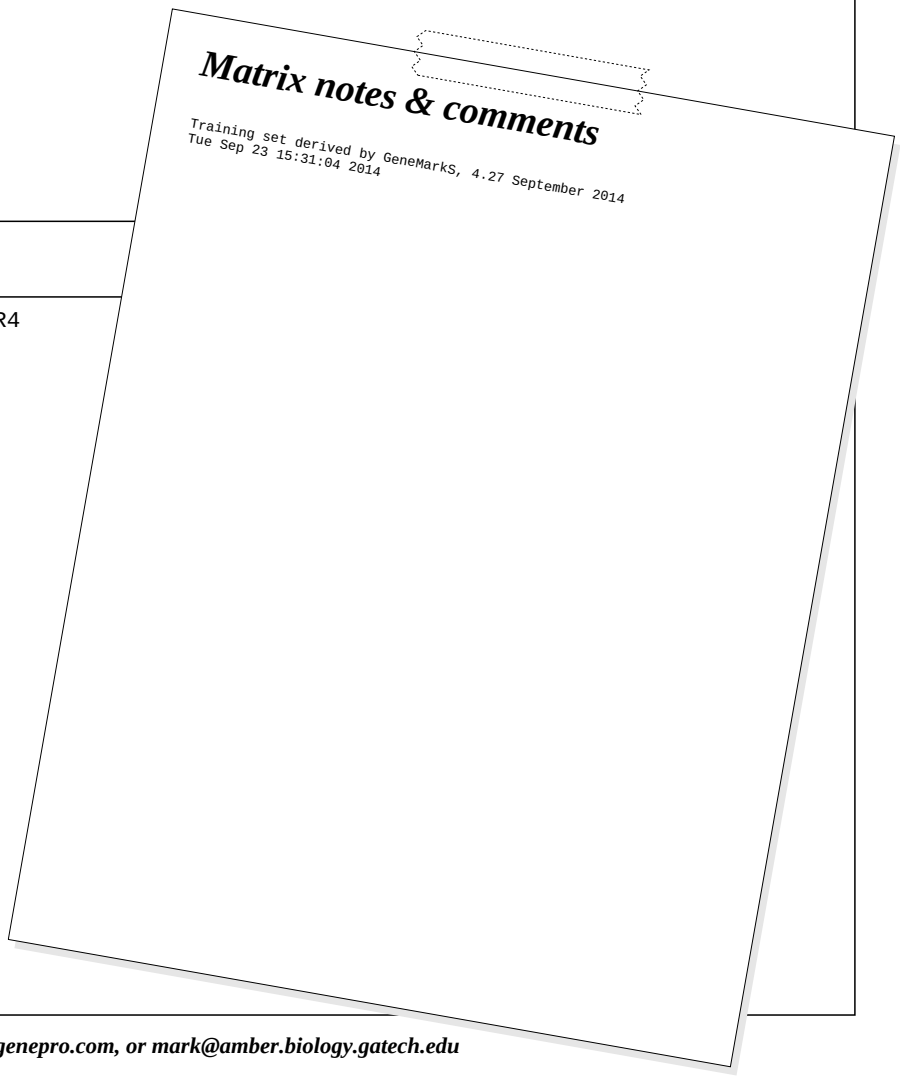
PROGRAM INFORMATION

```
Sequence         :  Rhodococcus phage Jflix2 complete sequence, 62017 bp including 6-base 3’ overhang (CCCGCT),
Analysis Date    :  3/29/24 at 1:43:23
Pages            :  31
Sequence Length  :  62017 bp
GC Content       :   66.55%

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 INFORMATION

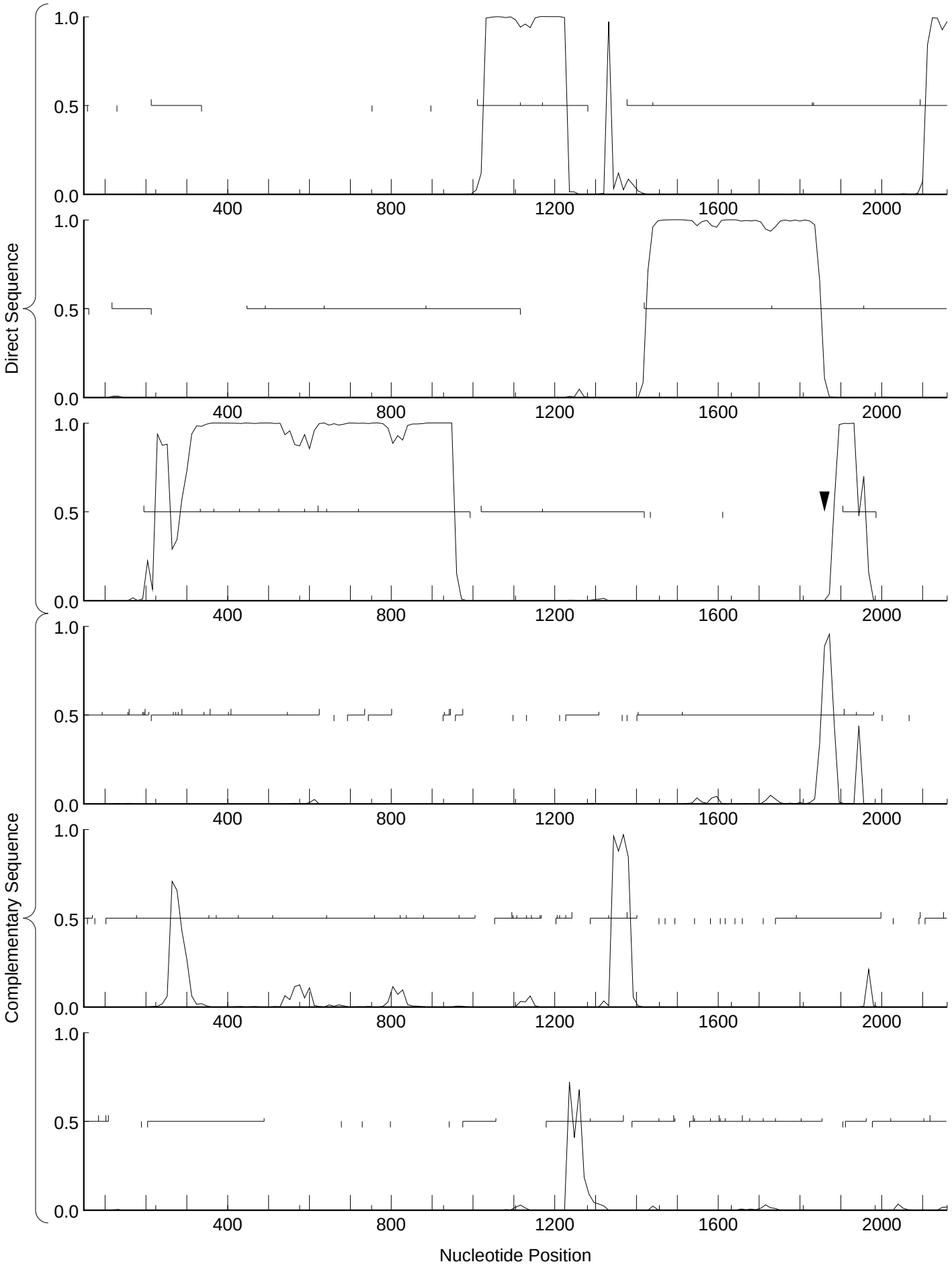
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Matrix     :  Rhodococcus_erythropolis_PR4
Author     :  -

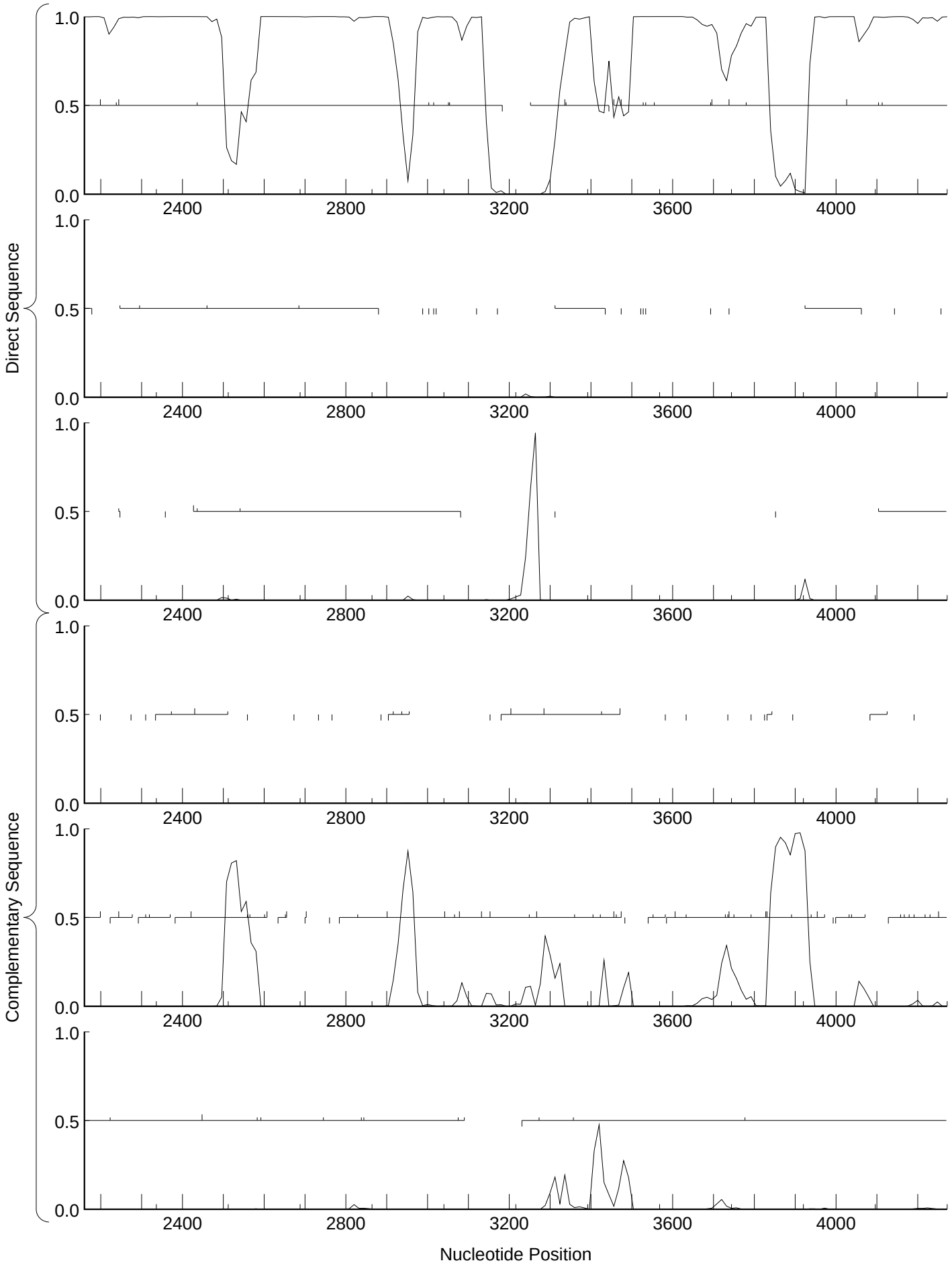
Order      :  4
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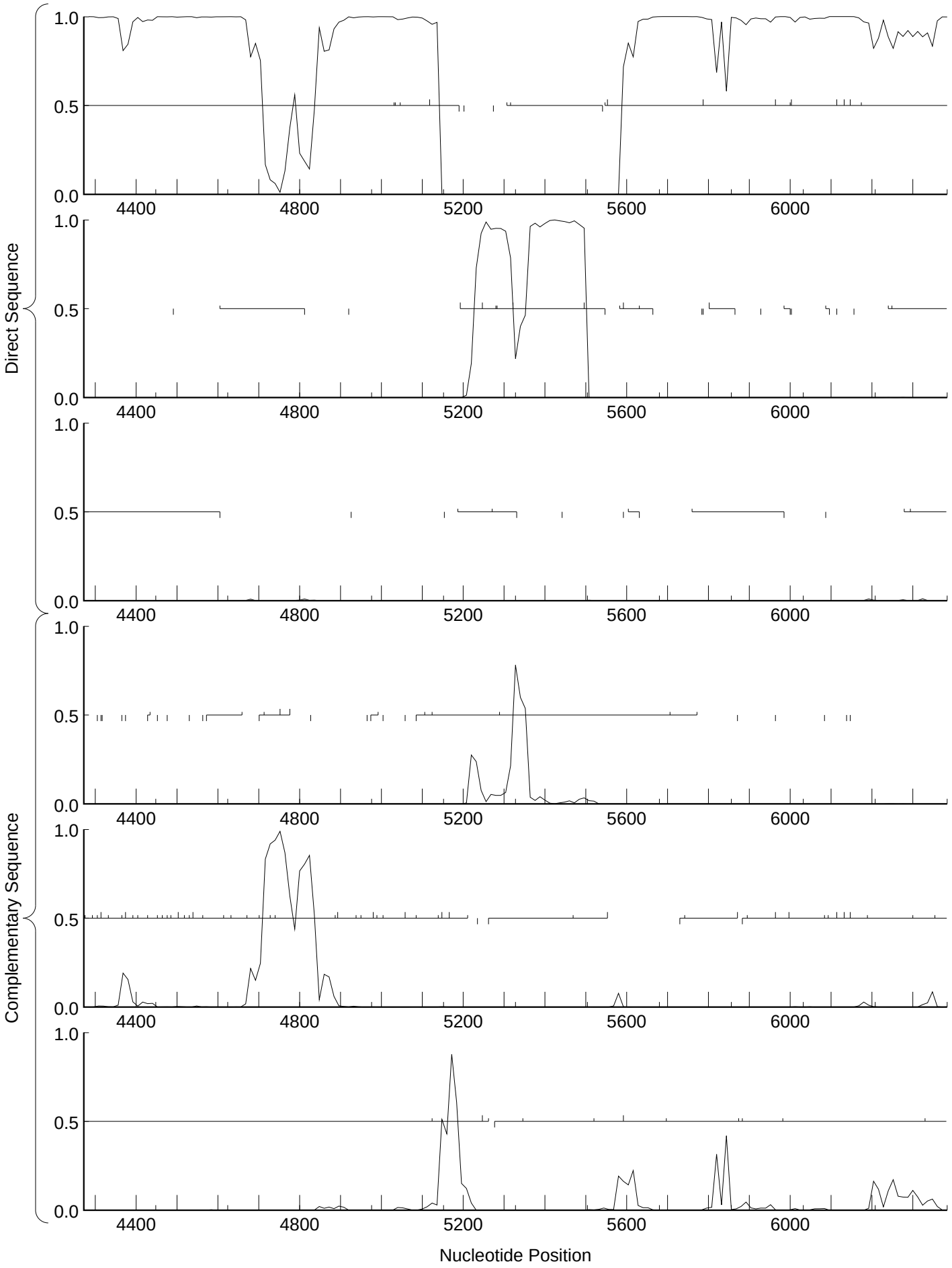

```
  Send questions / comments to:
      Dr. M. Borodovsky
      Georgia Institute of Technology
      School of Biology
      Atlanta, GA  30332-0230
```


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

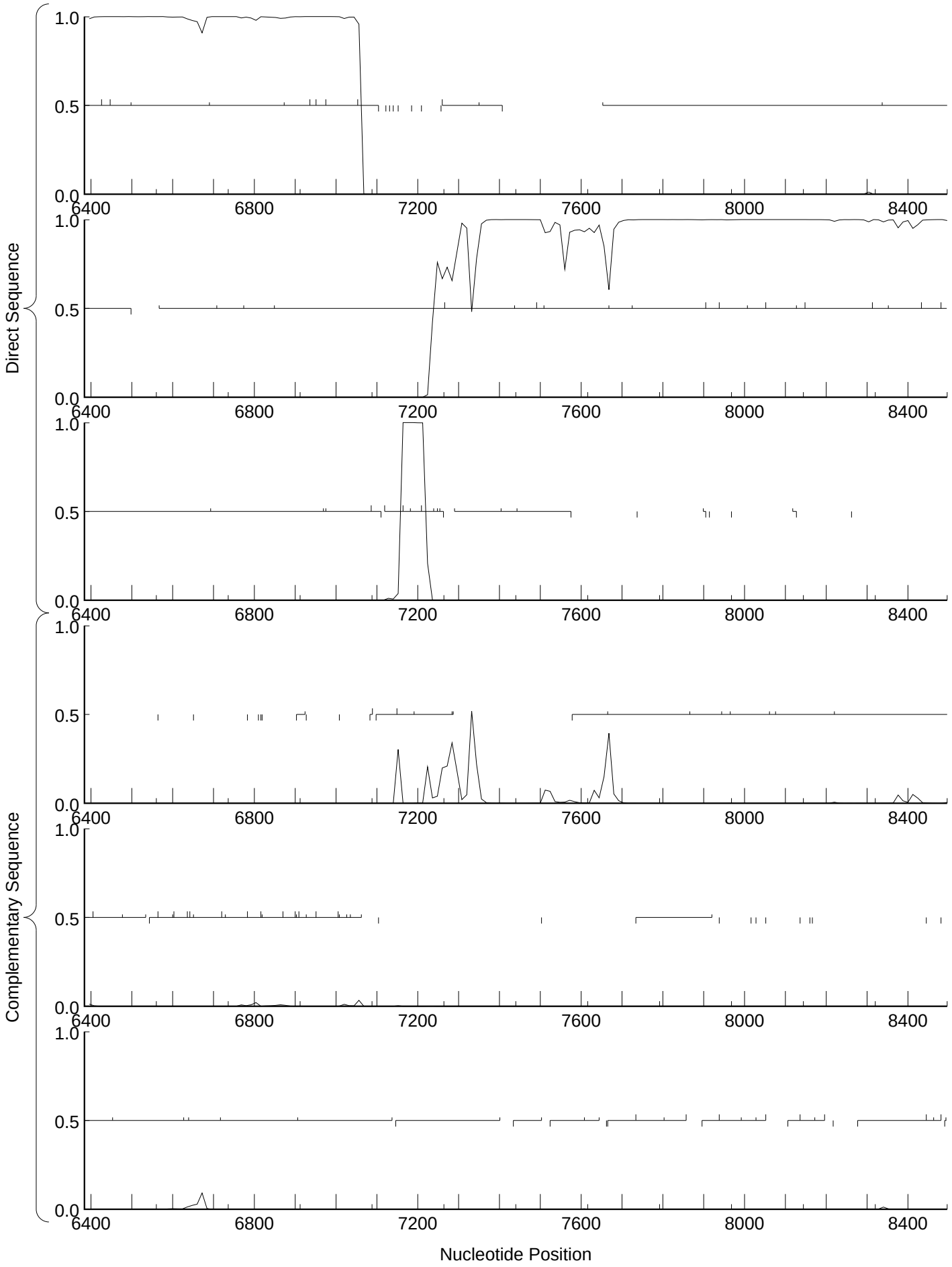
age Jflix2 complete sequence, 62017 bp including 6-base 3' overhang (CCCGCT), Cluster CF, Order 4, Window 96, Step 12, 2/31

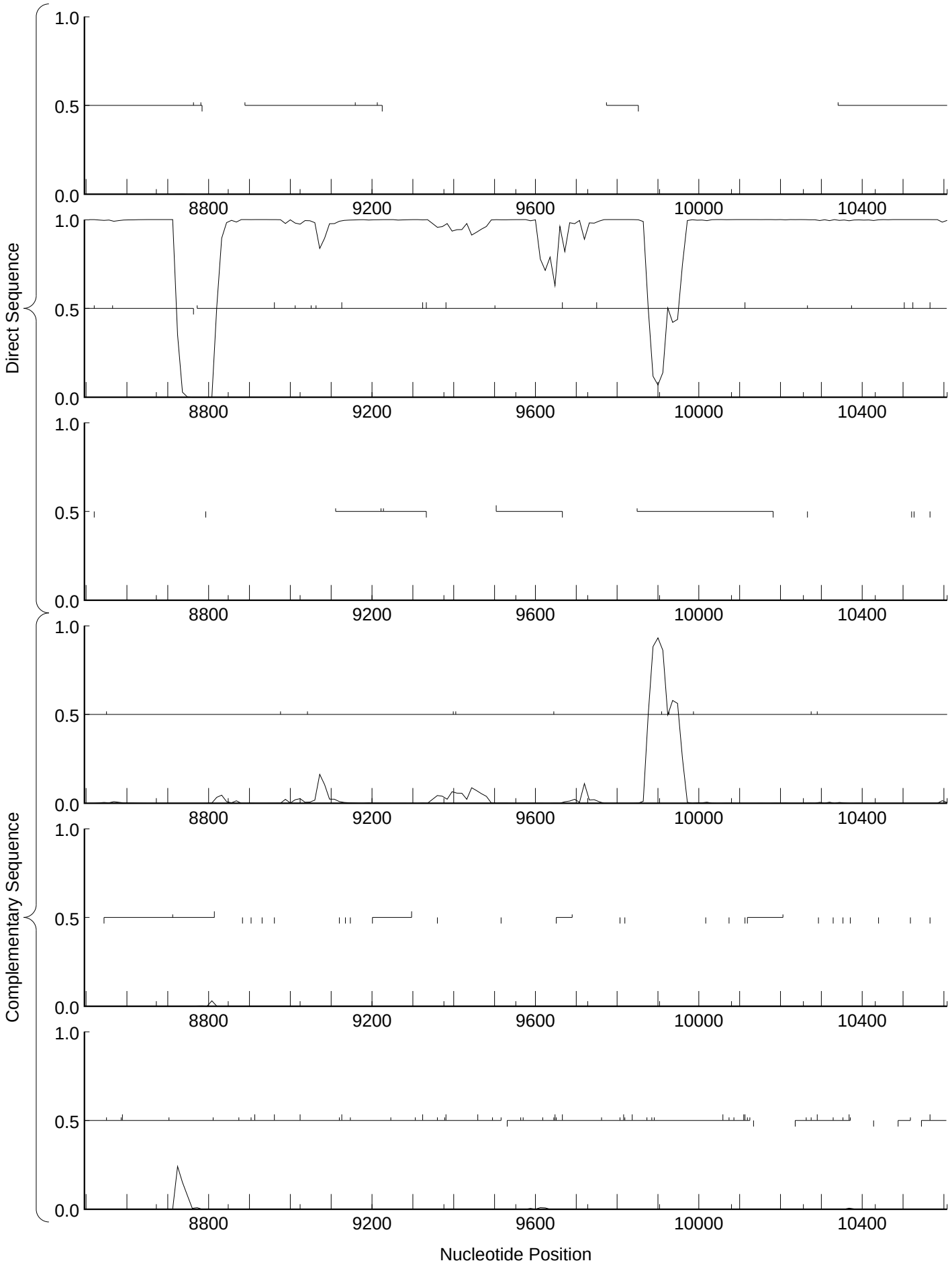


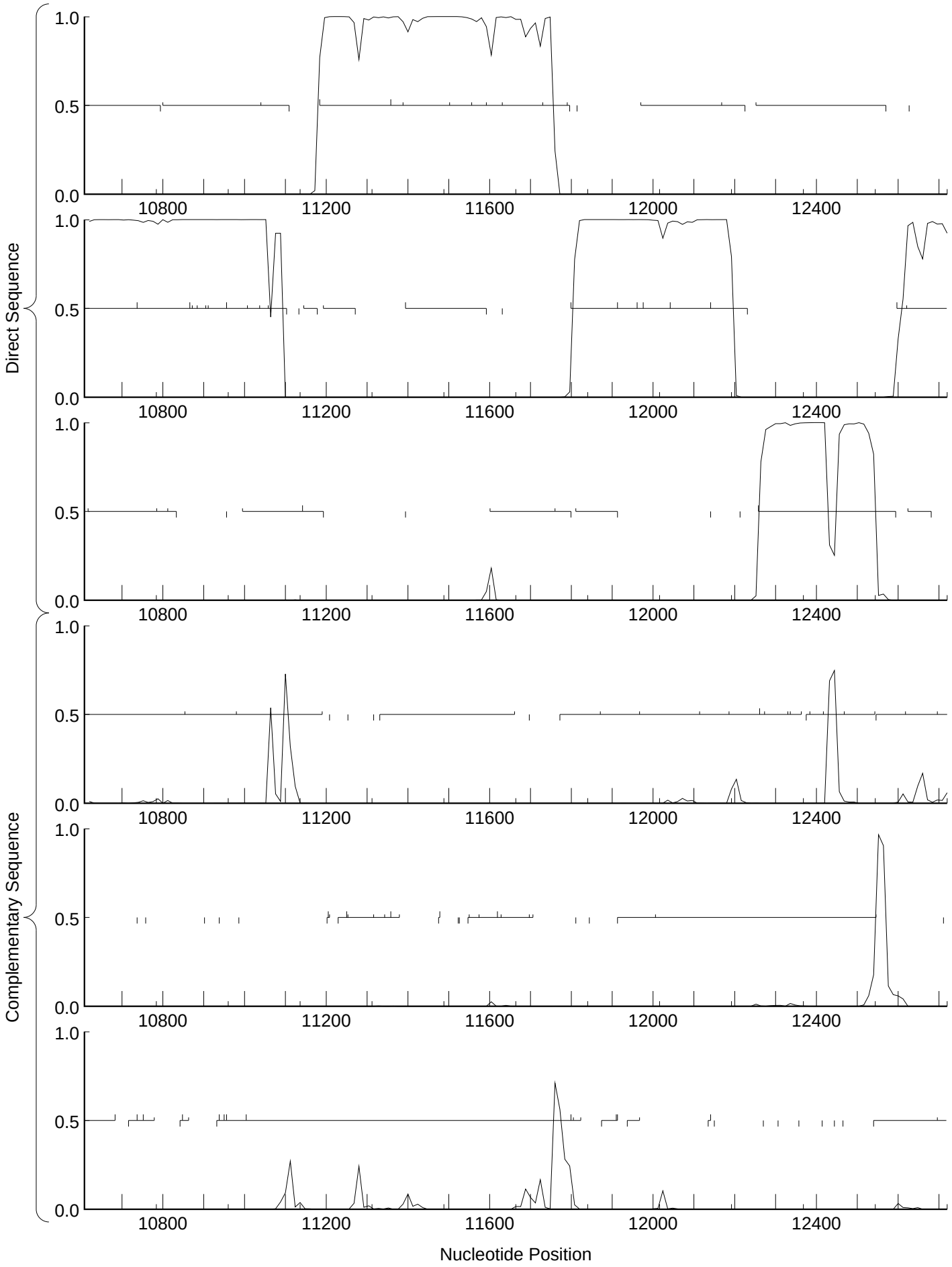

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Nucleotide Position

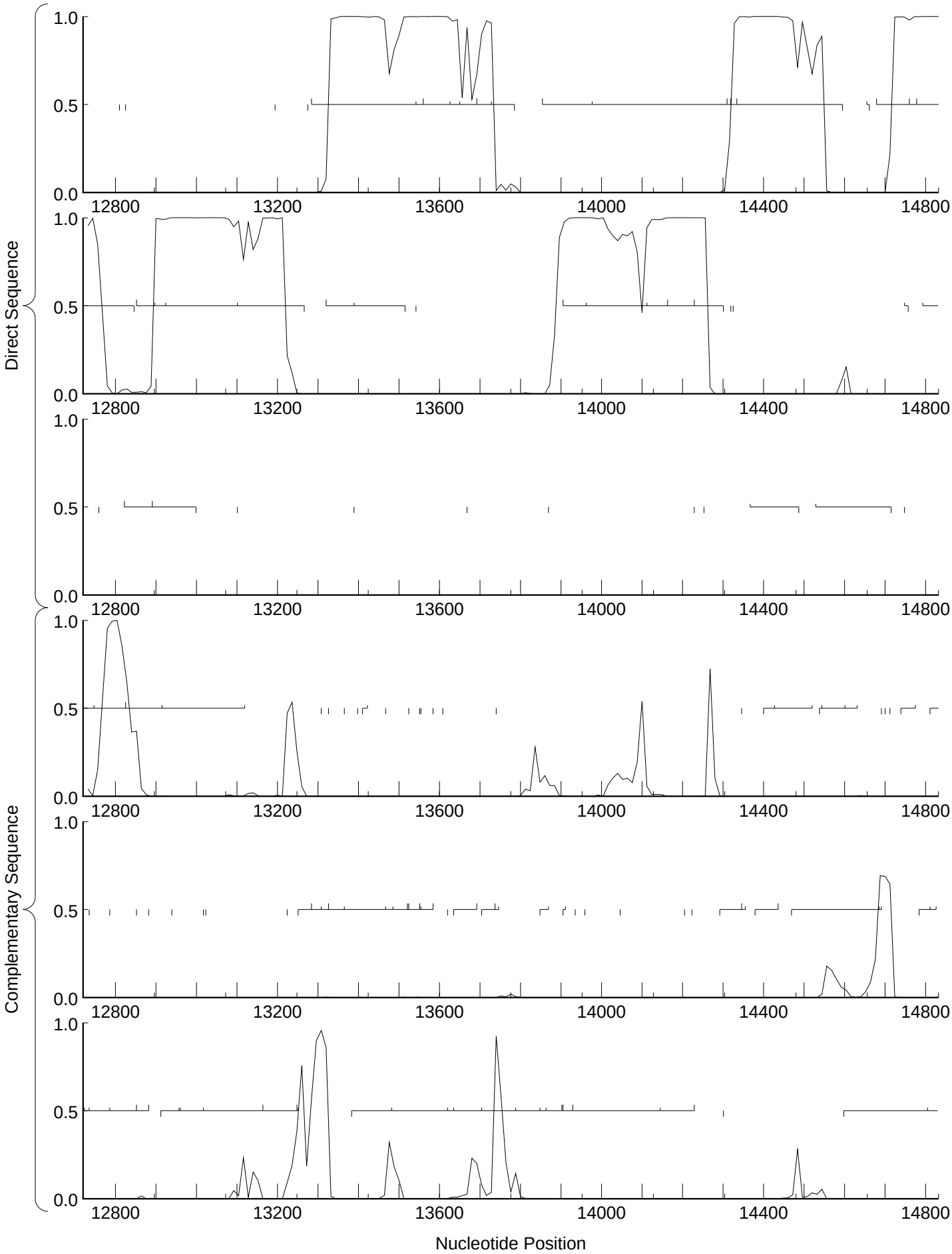

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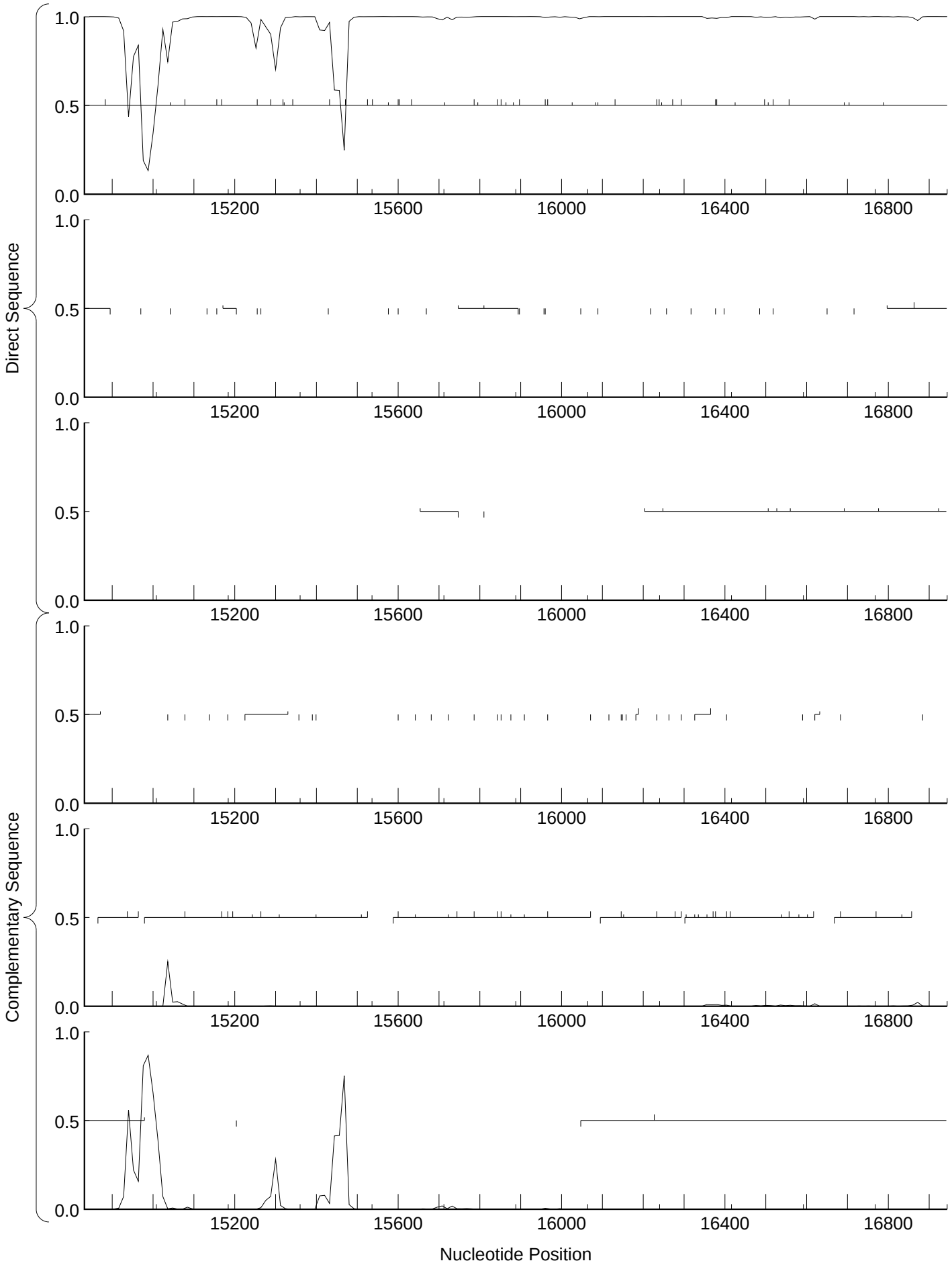
age Jflix2 complete sequence, 62017 bp including 6-base 3’ overhang (CCCGCT), Cluster CF, Order 4, Window 96, Step 12, 5/31



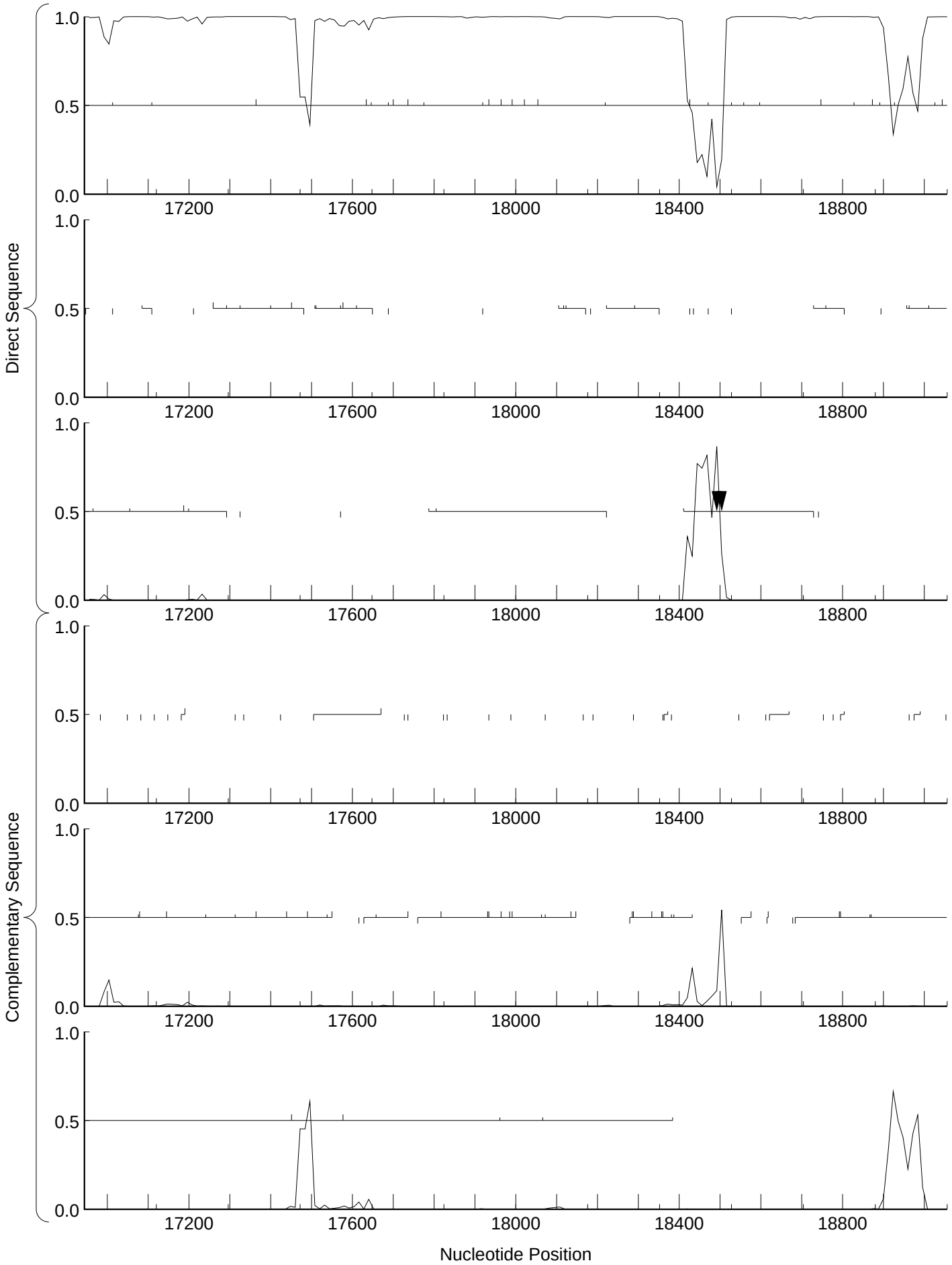

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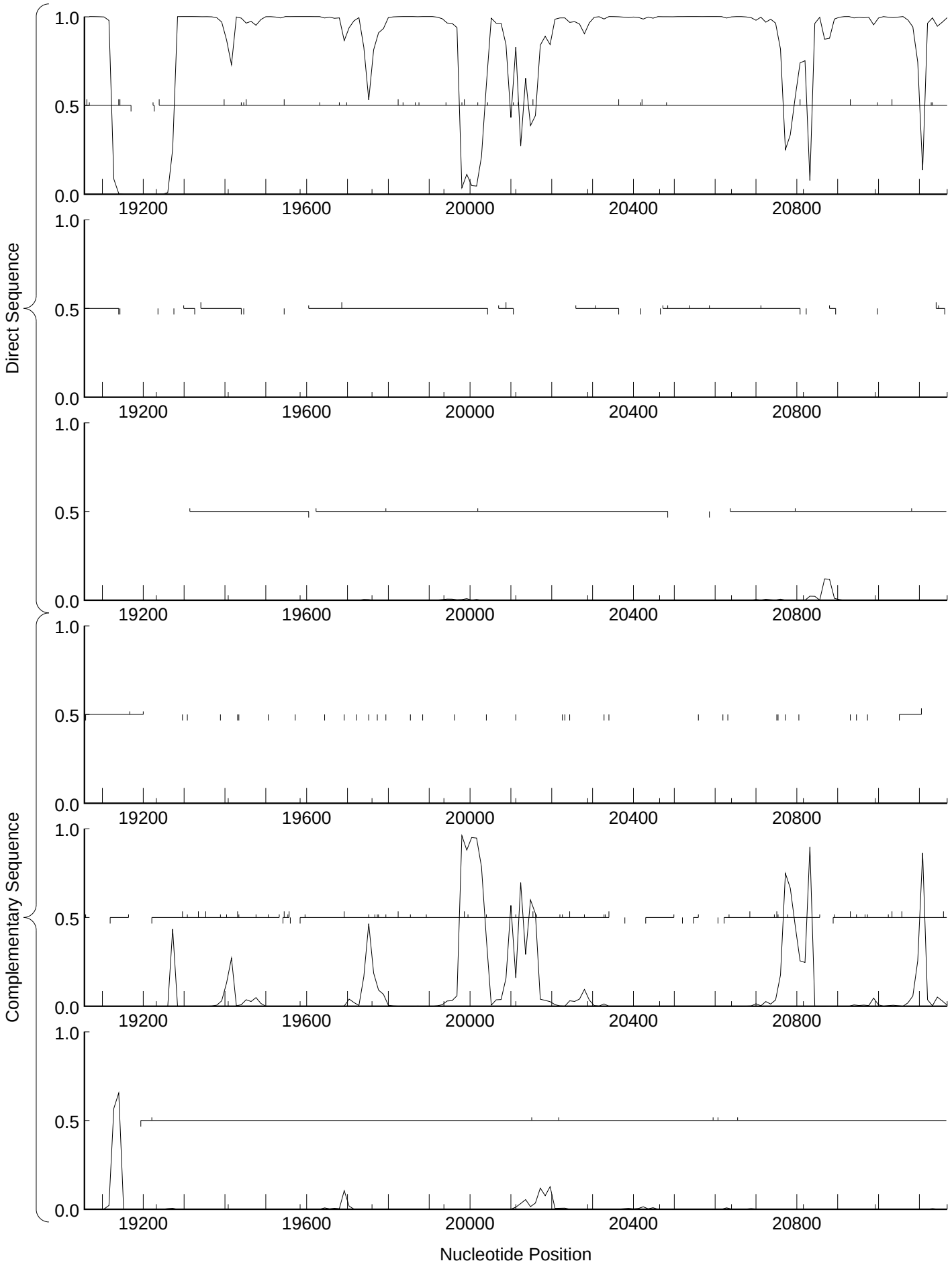

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Nucleotide Position

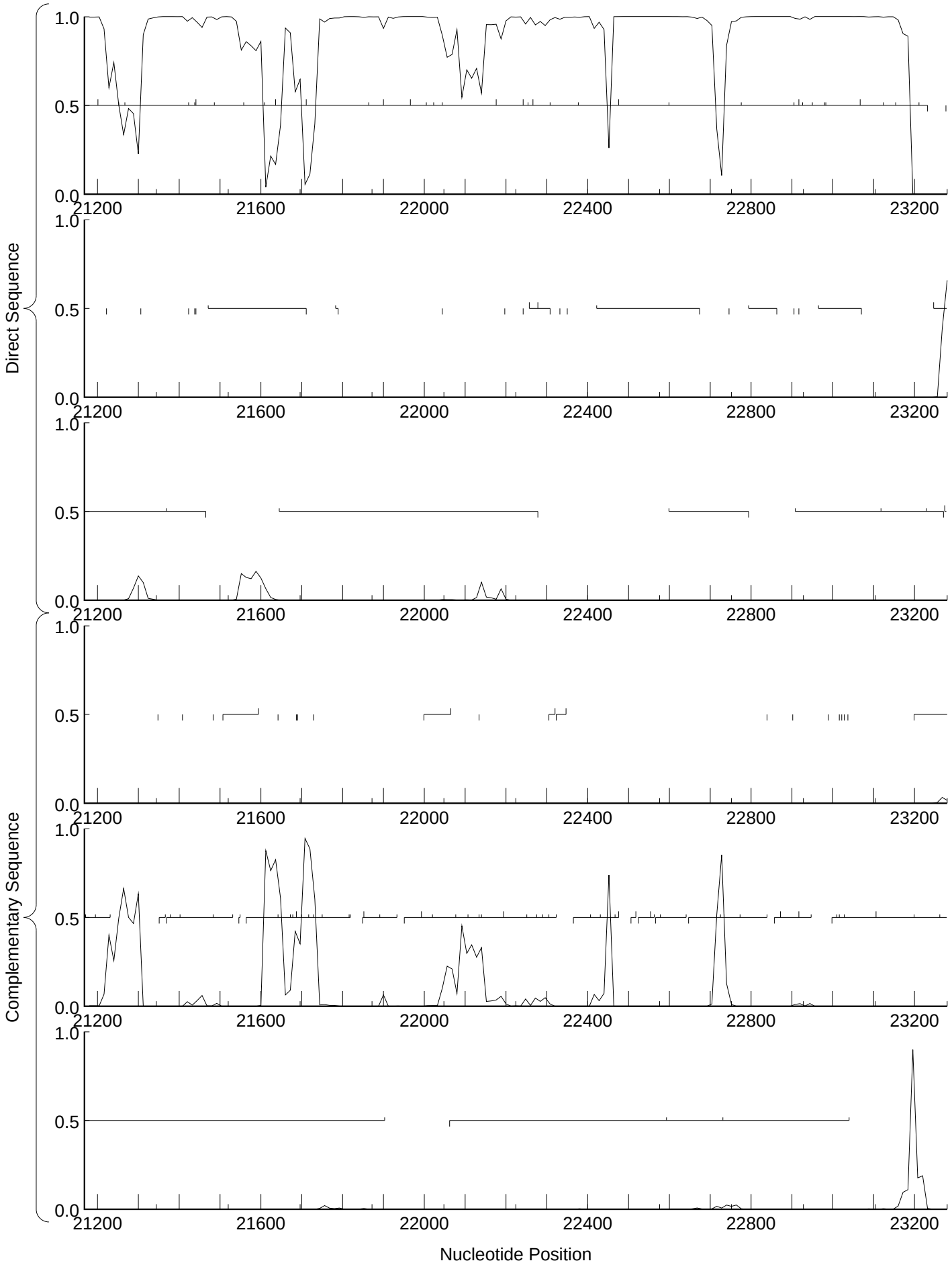

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Complementary Sequence
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14000
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Nucleotide Position


Direct Sequence
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Nucleotide Position

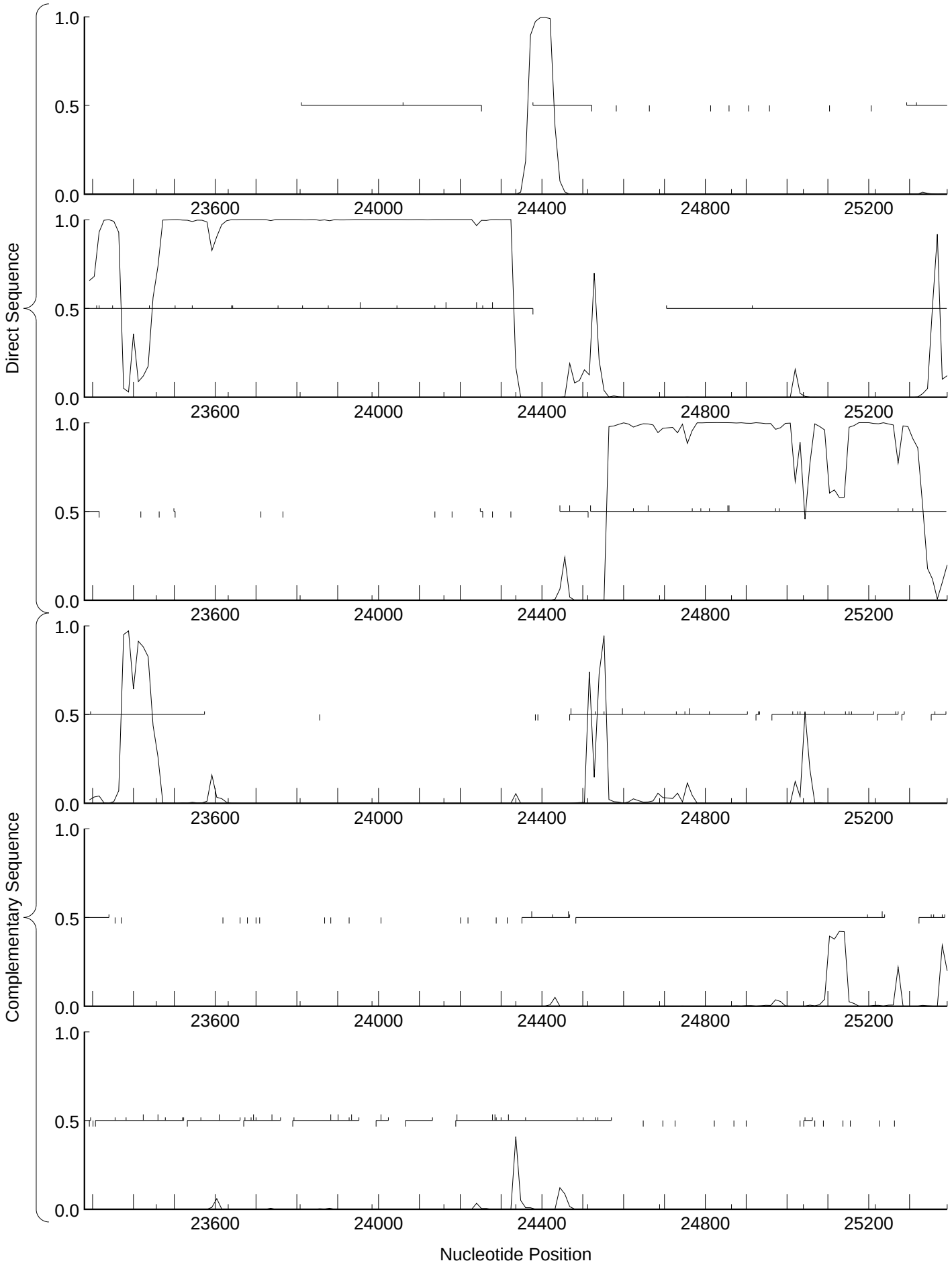
ge Jflix2 complete sequence, 62017 bp including 6-base 3’ overhang (CCCGCT), Cluster CF, Order 4, Window 96, Step 12, 10/31

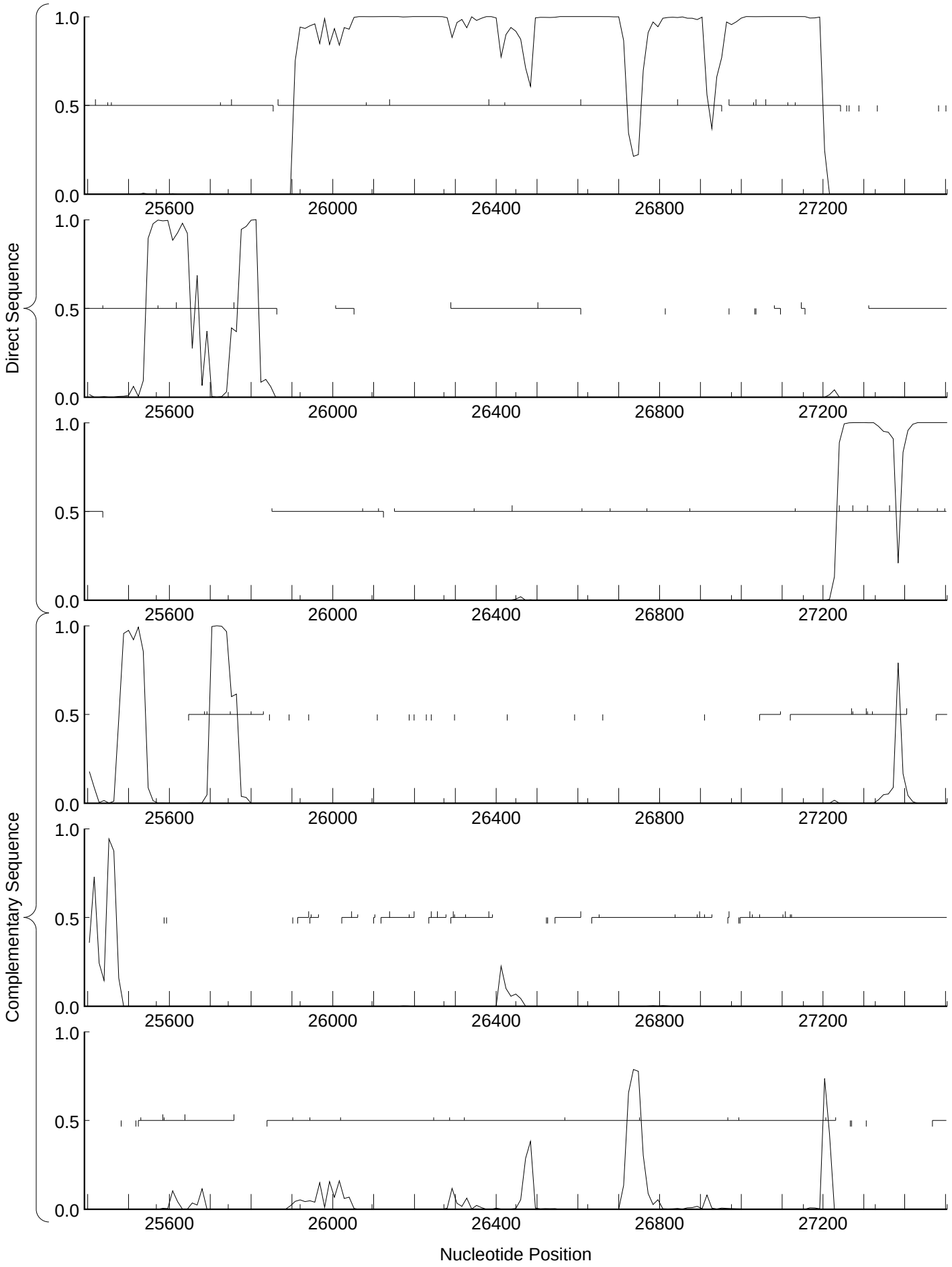



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Nucleotide Position

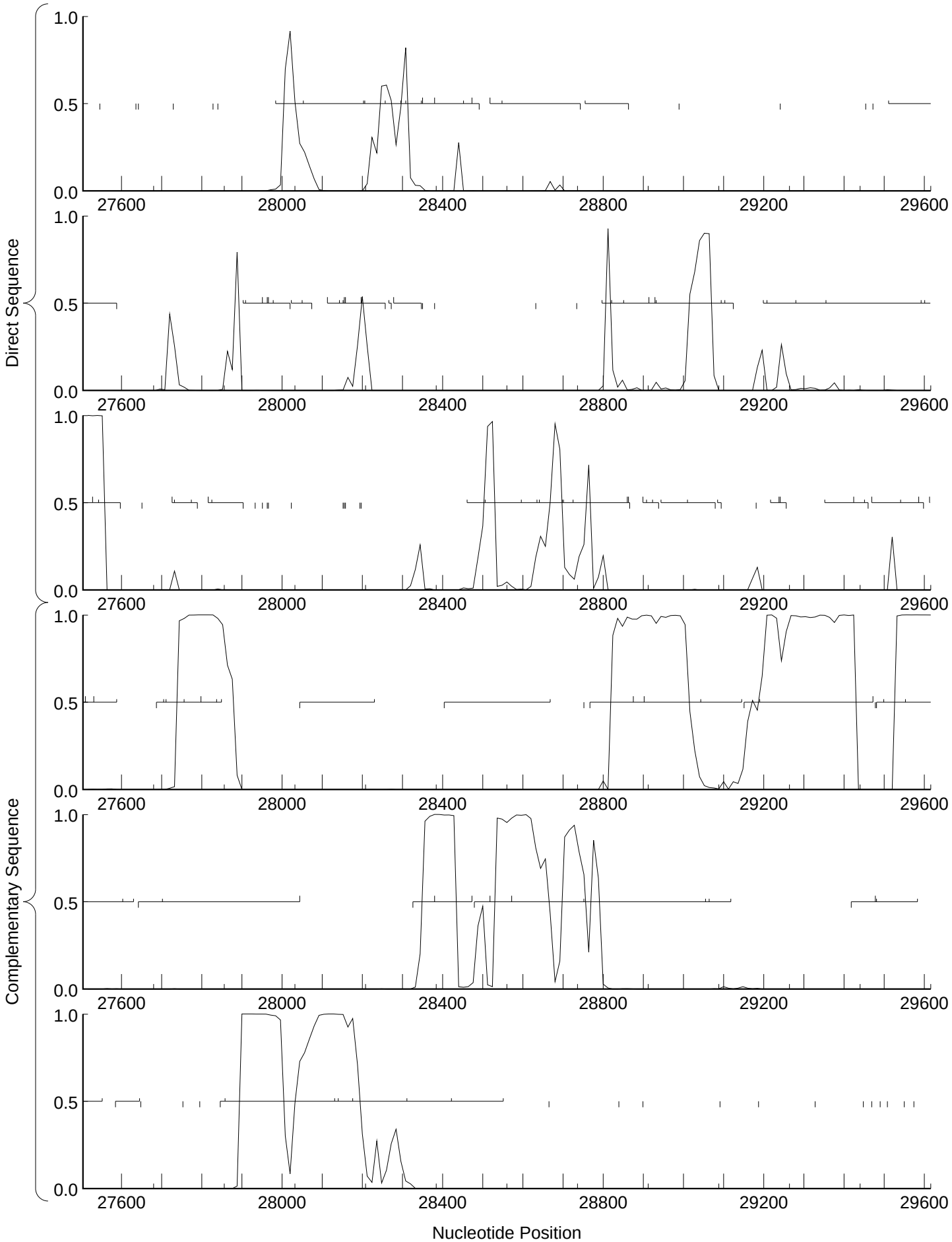

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Nucleotide Position

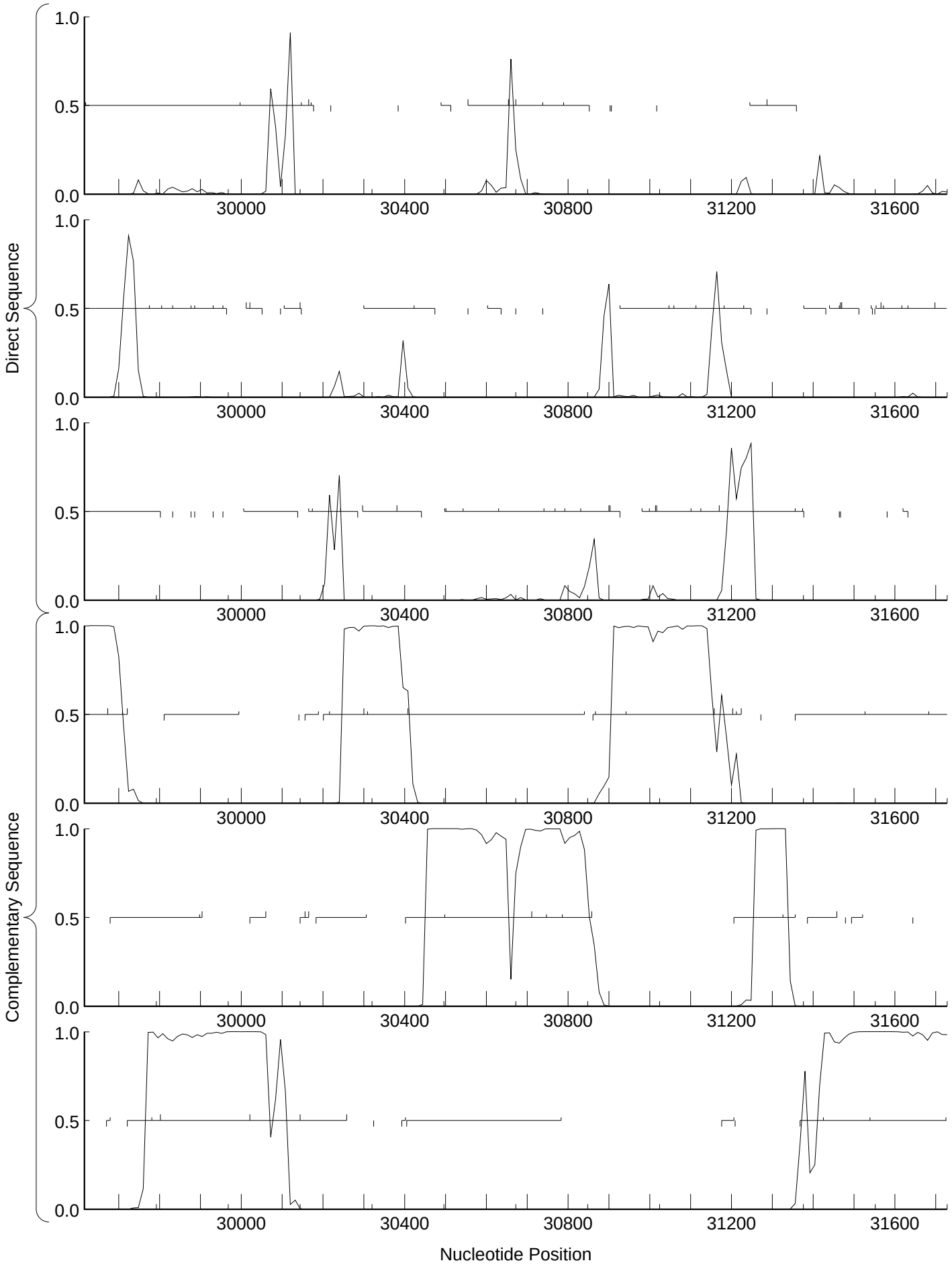
ge Jflix2 complete sequence, 62017 bp including 6-base 3’ overhang (CCCGCT), Cluster CF, Order 4, Window 96, Step 12, 13/31

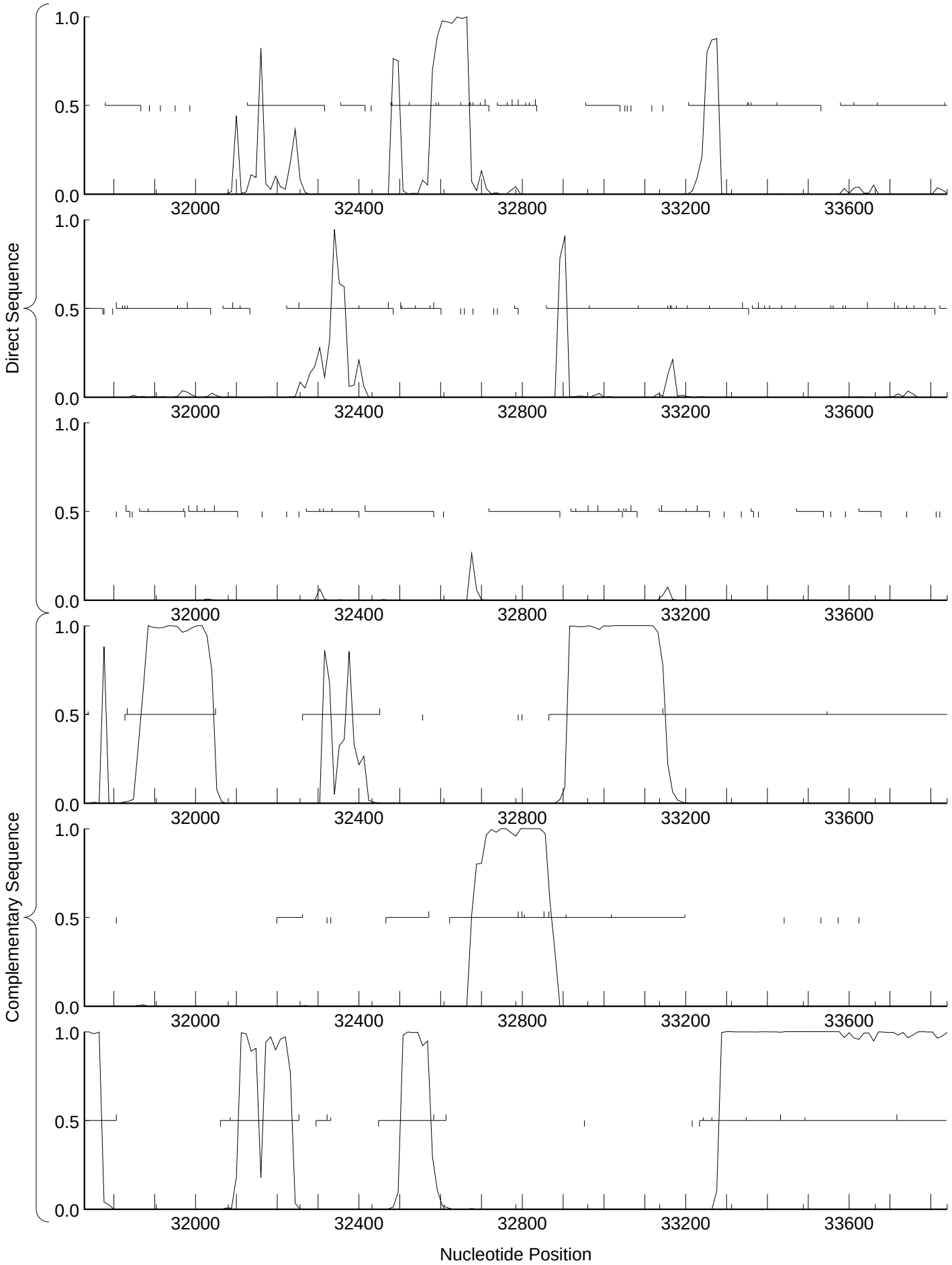


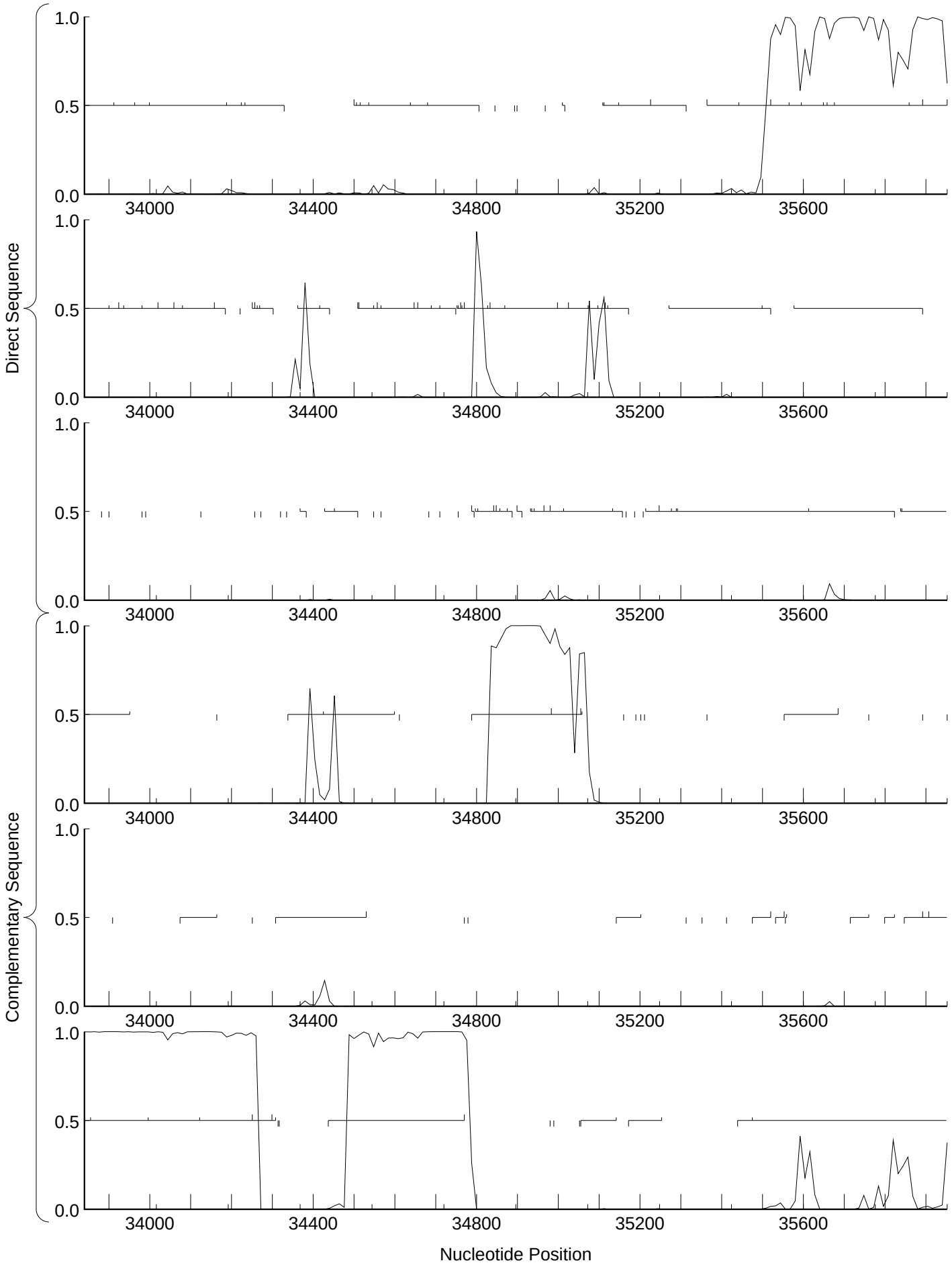

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Complementary Sequence
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Nucleotide Position

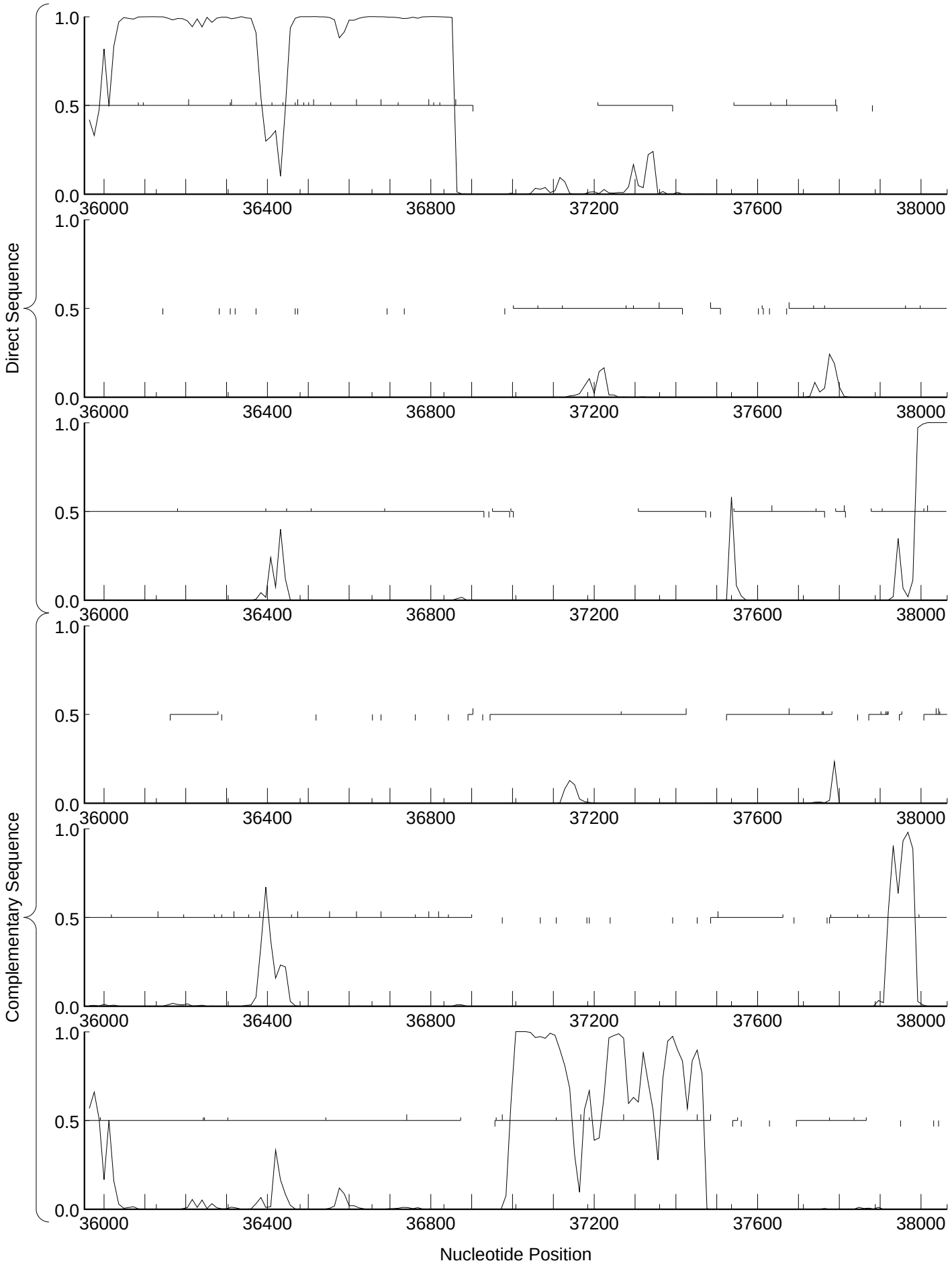
ge Jflix2 complete sequence, 62017 bp including 6-base 3’ overhang (CCCGCT), Cluster CF, Order 4, Window 96, Step 12, 15/31

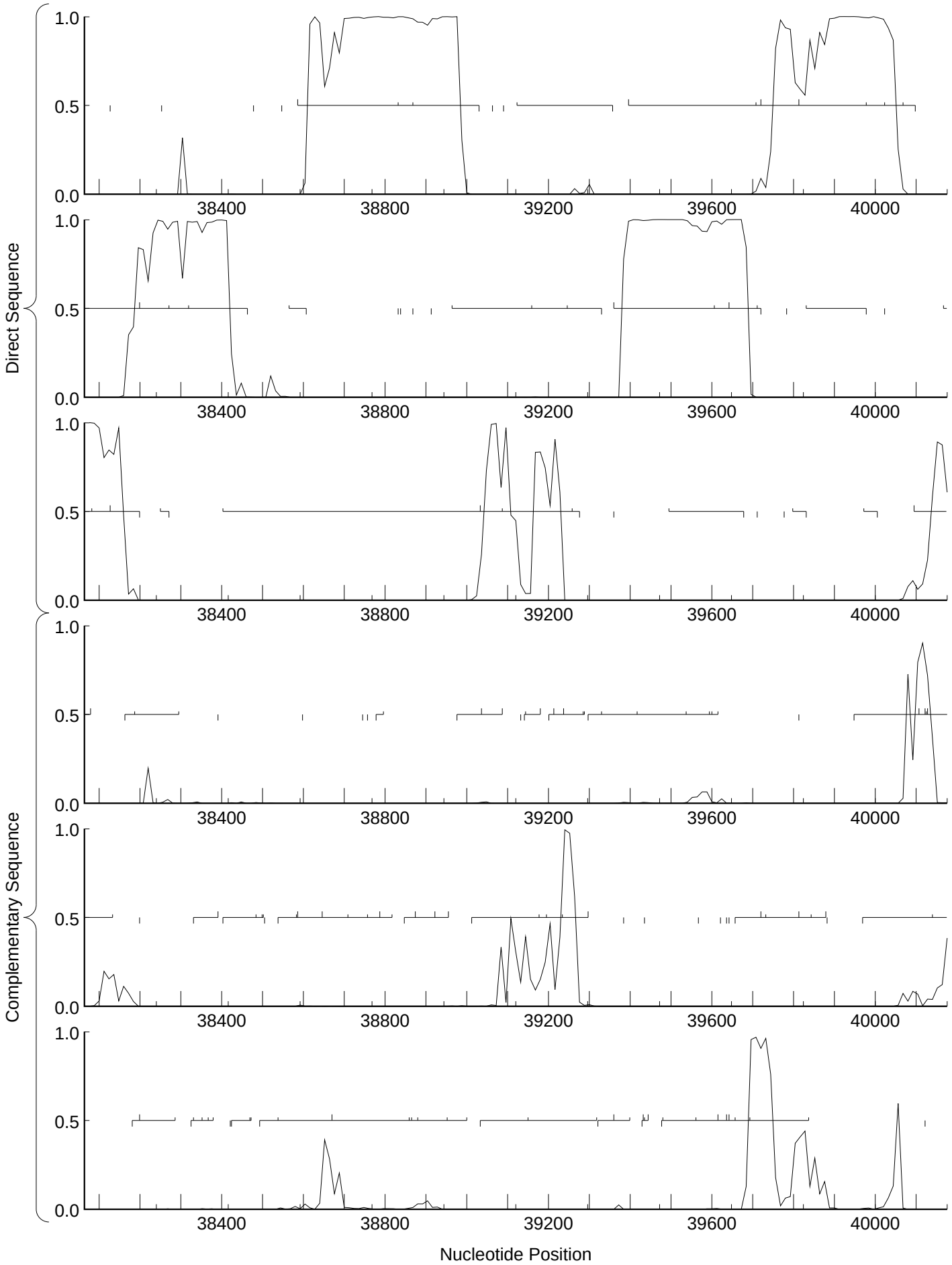



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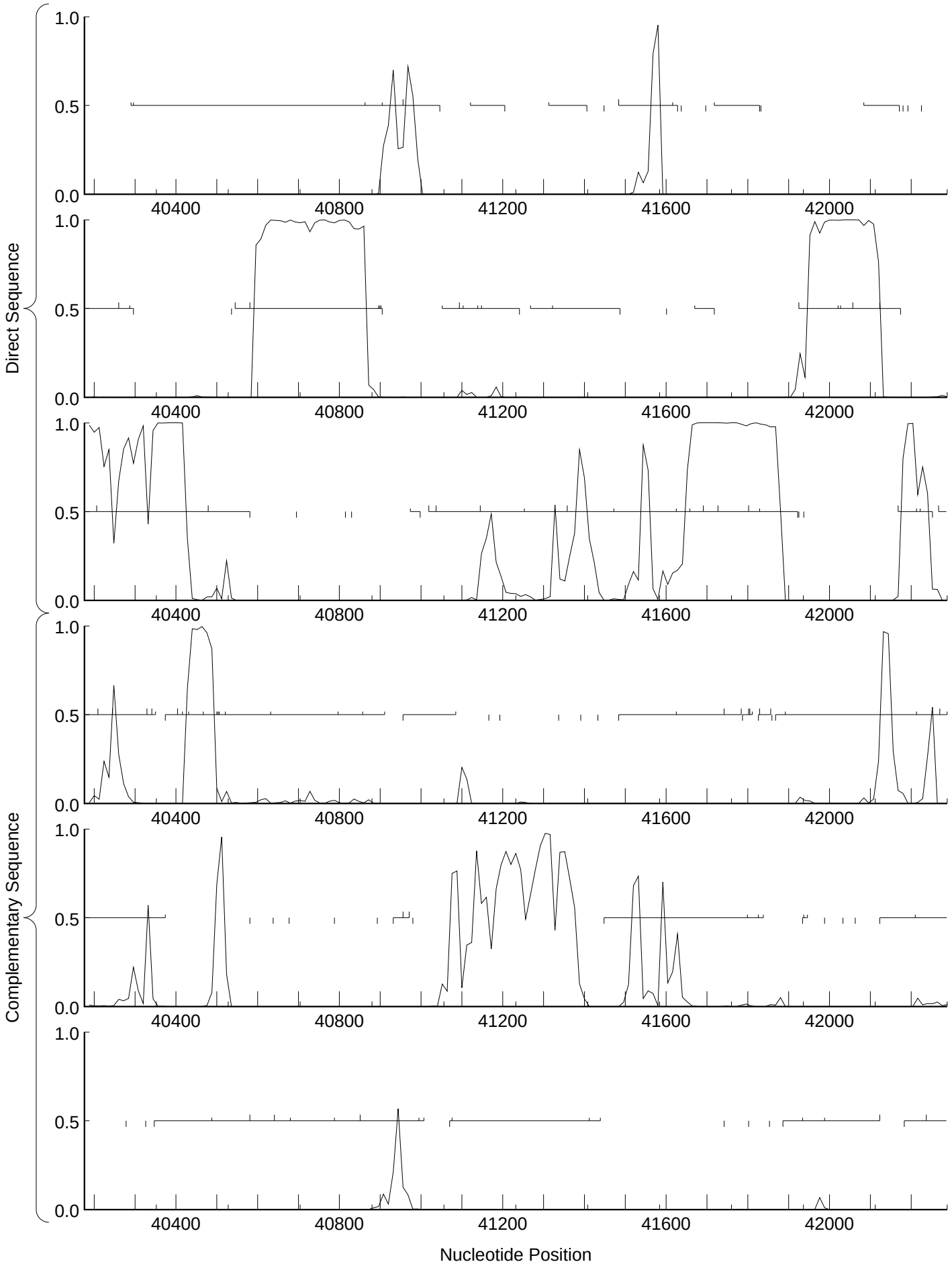

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Nucleotide Position

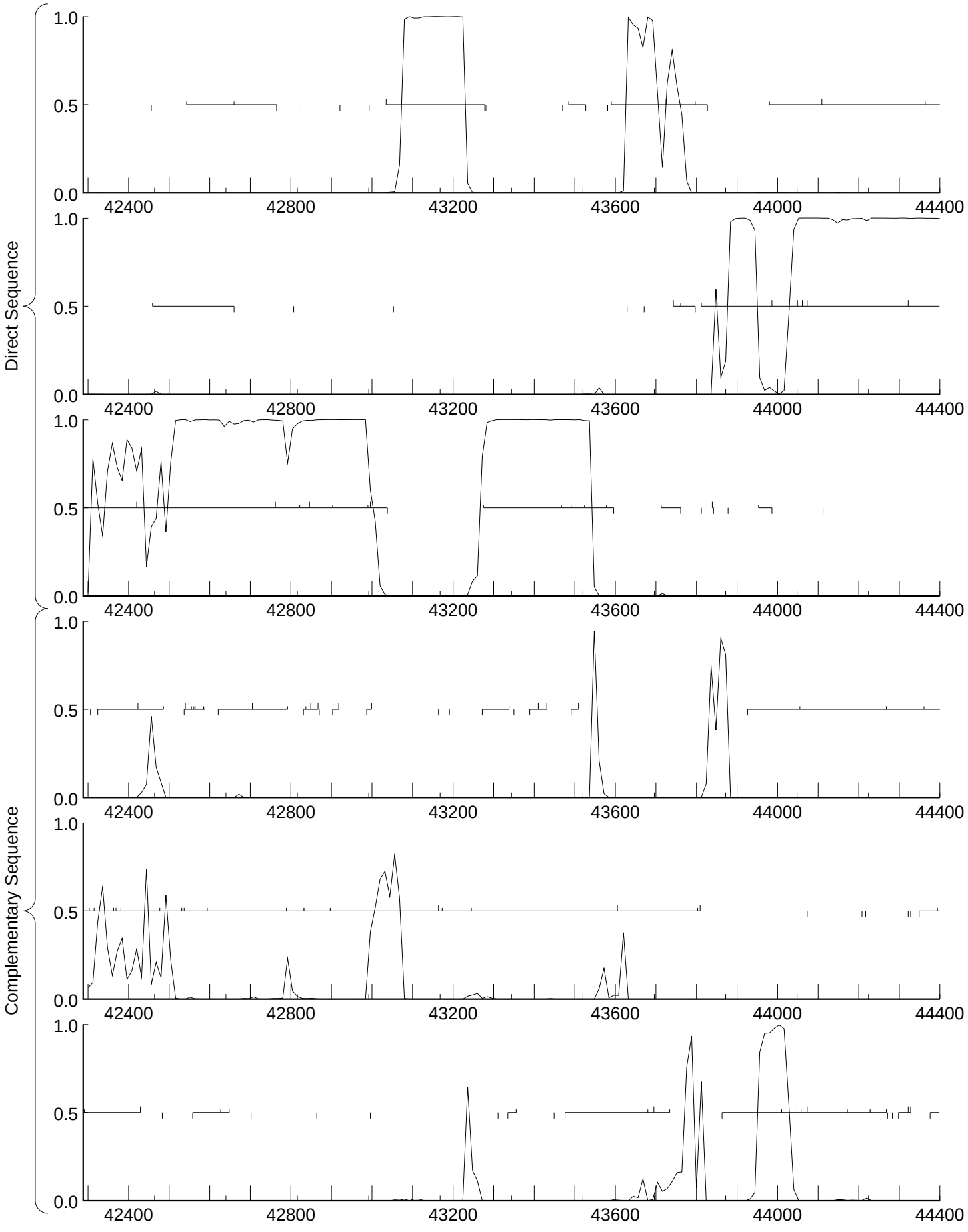

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Nucleotide Position

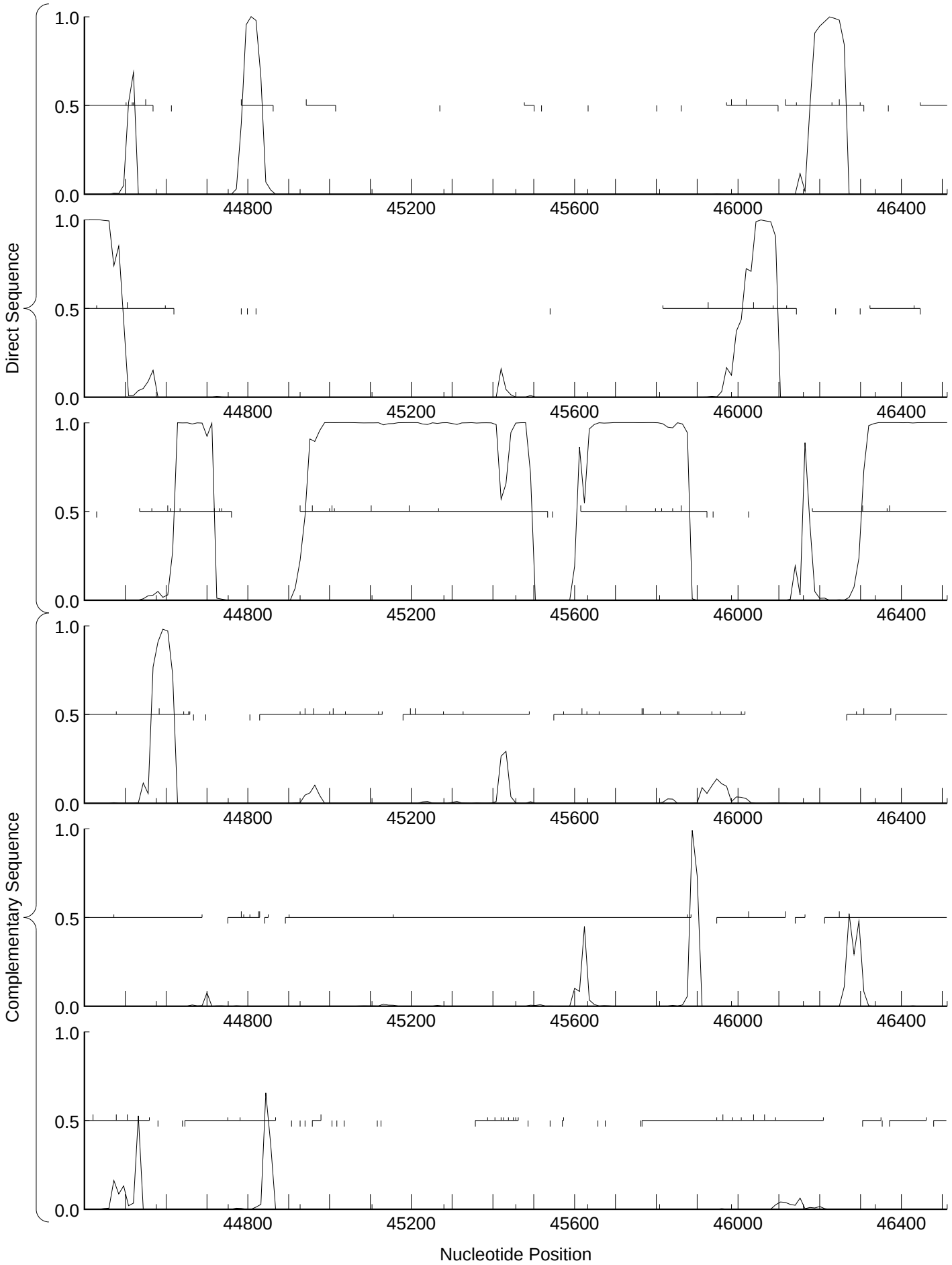

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40000
Nucleotide Position

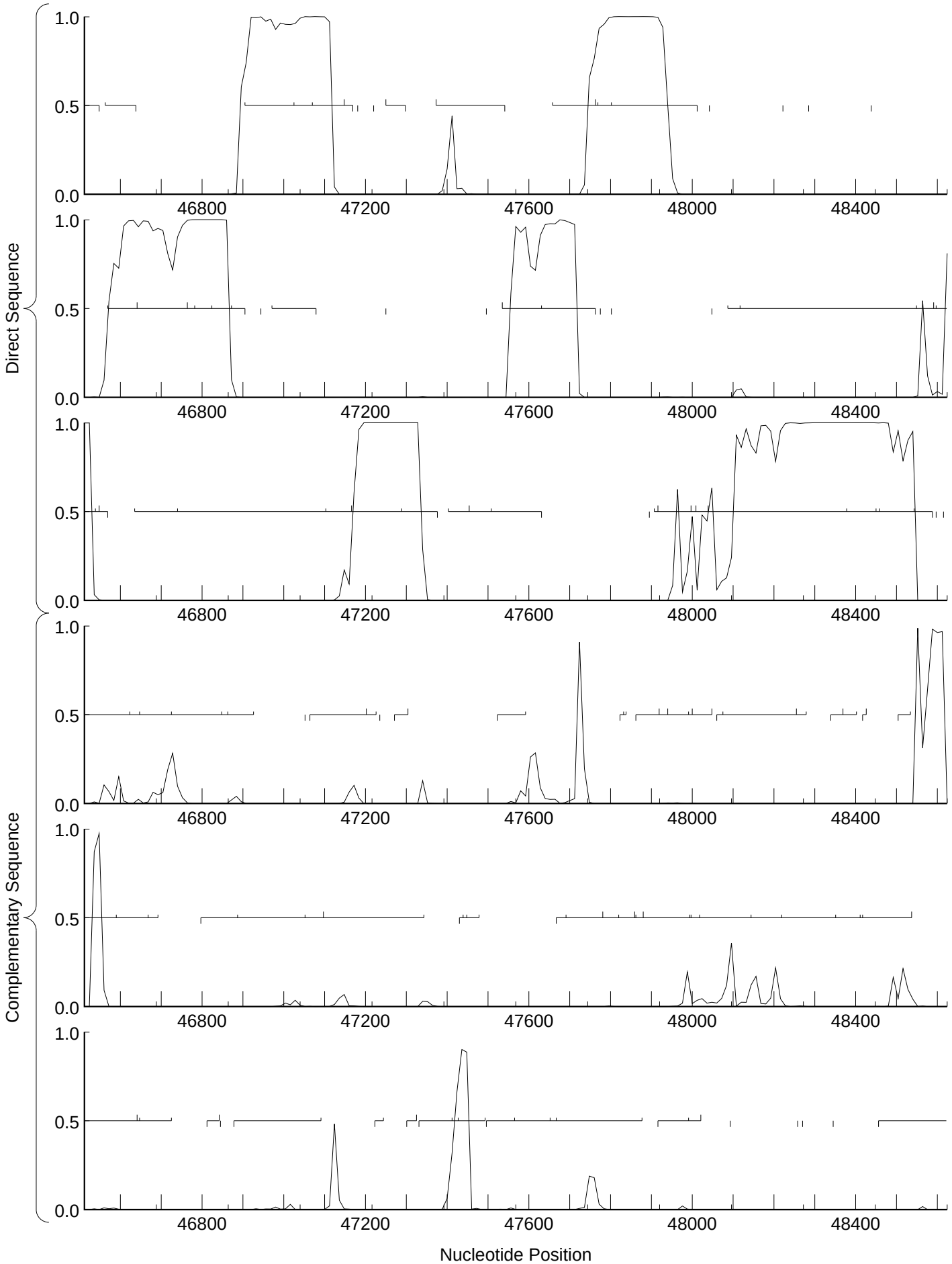
ge Jflix2 complete sequence, 62017 bp including 6-base 3’ overhang (CCCGCT), Cluster CF, Order 4, Window 96, Step 12, 21/31

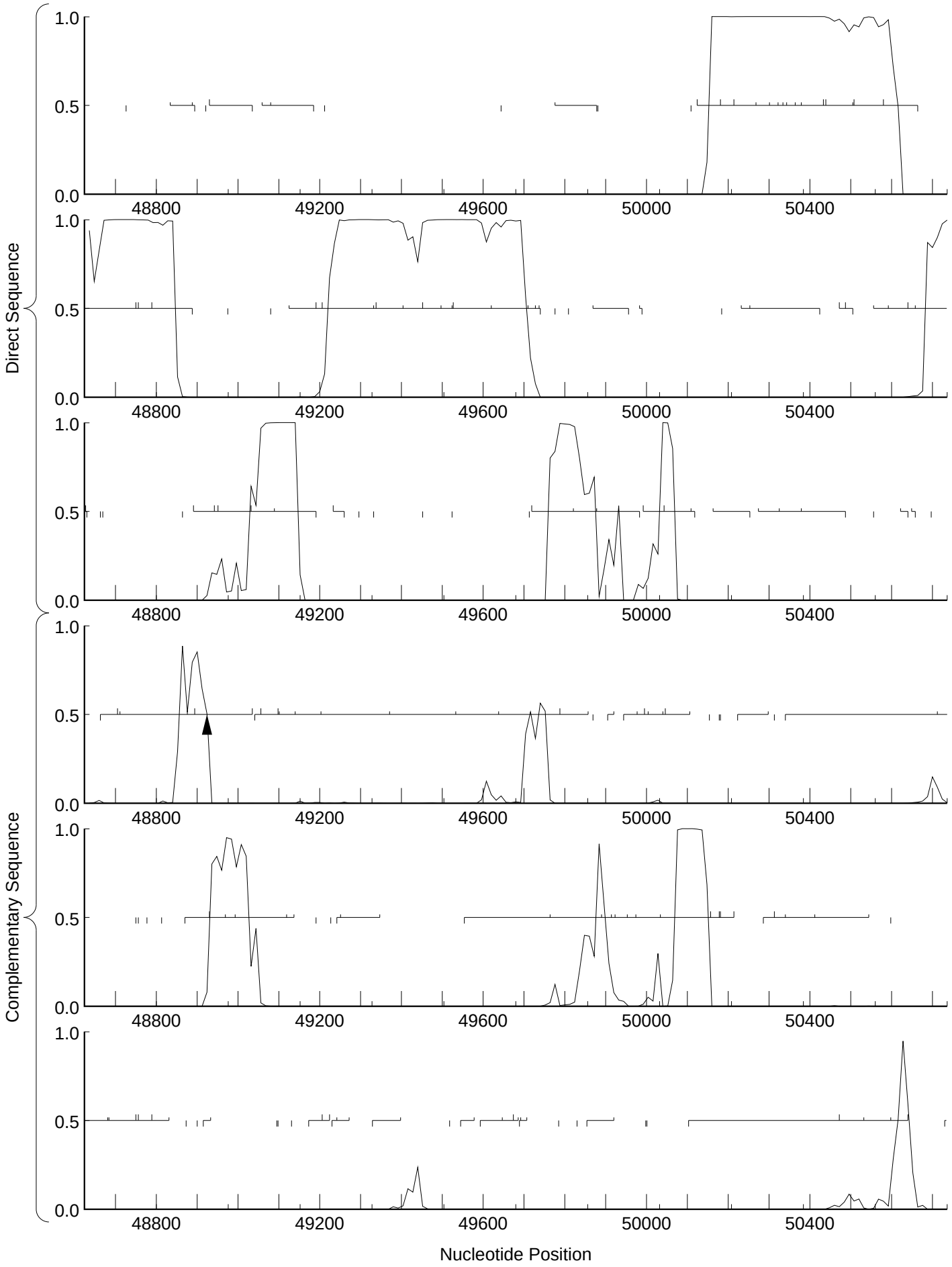


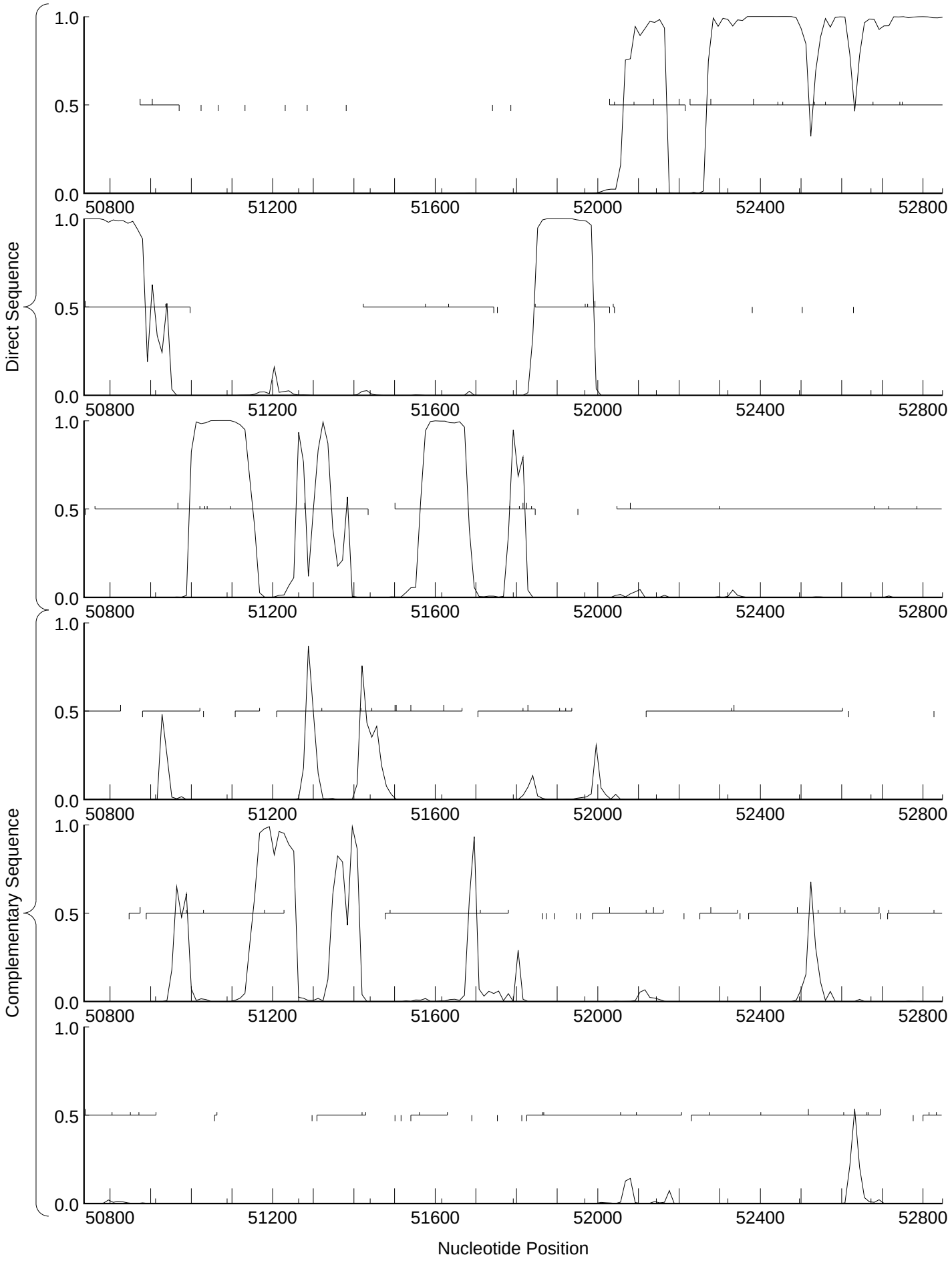

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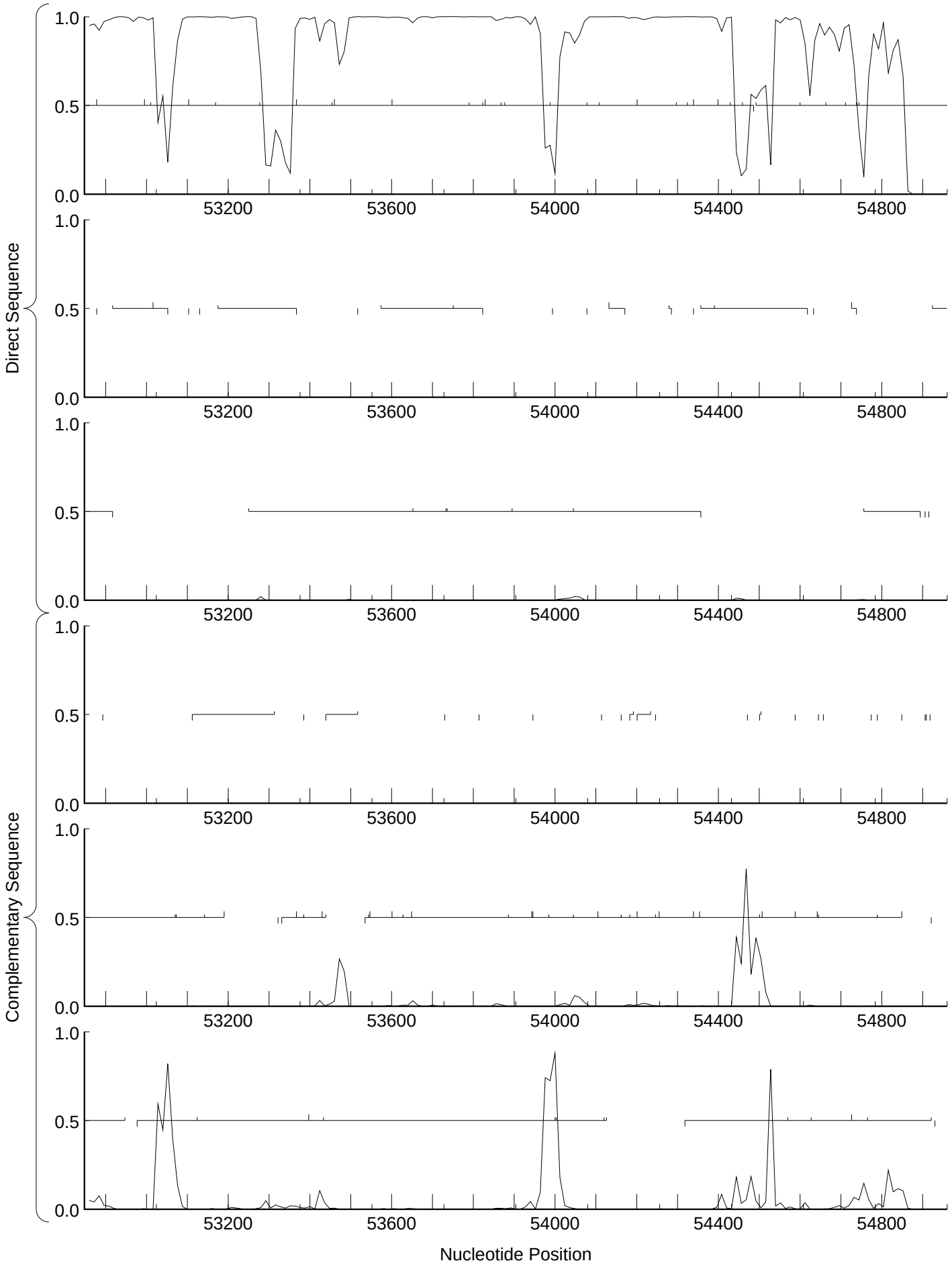

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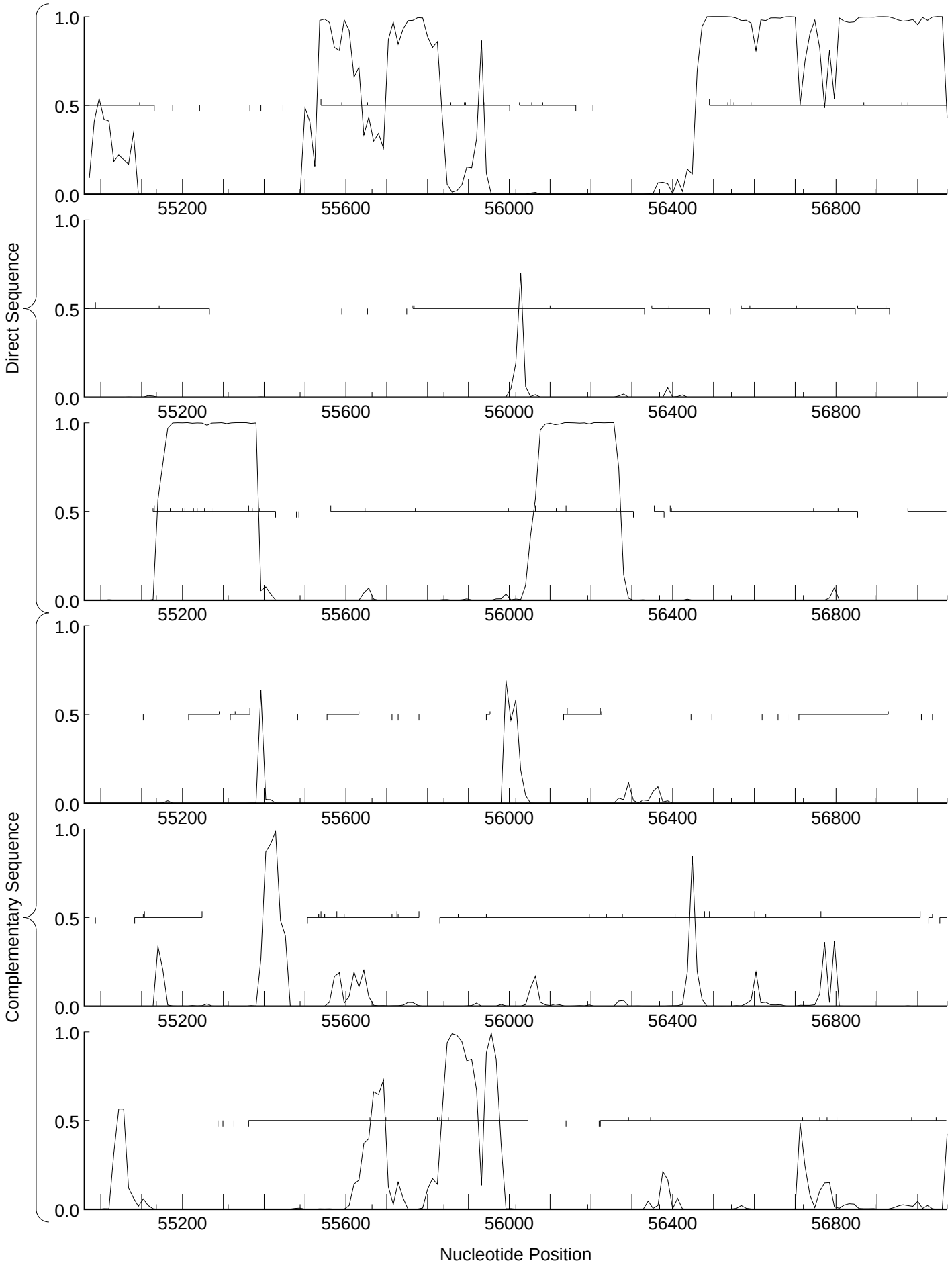
ge Jflix2 complete sequence, 62017 bp including 6-base 3’ overhang (CCCGCT), Cluster CF, Order 4, Window 96, Step 12, 24/31



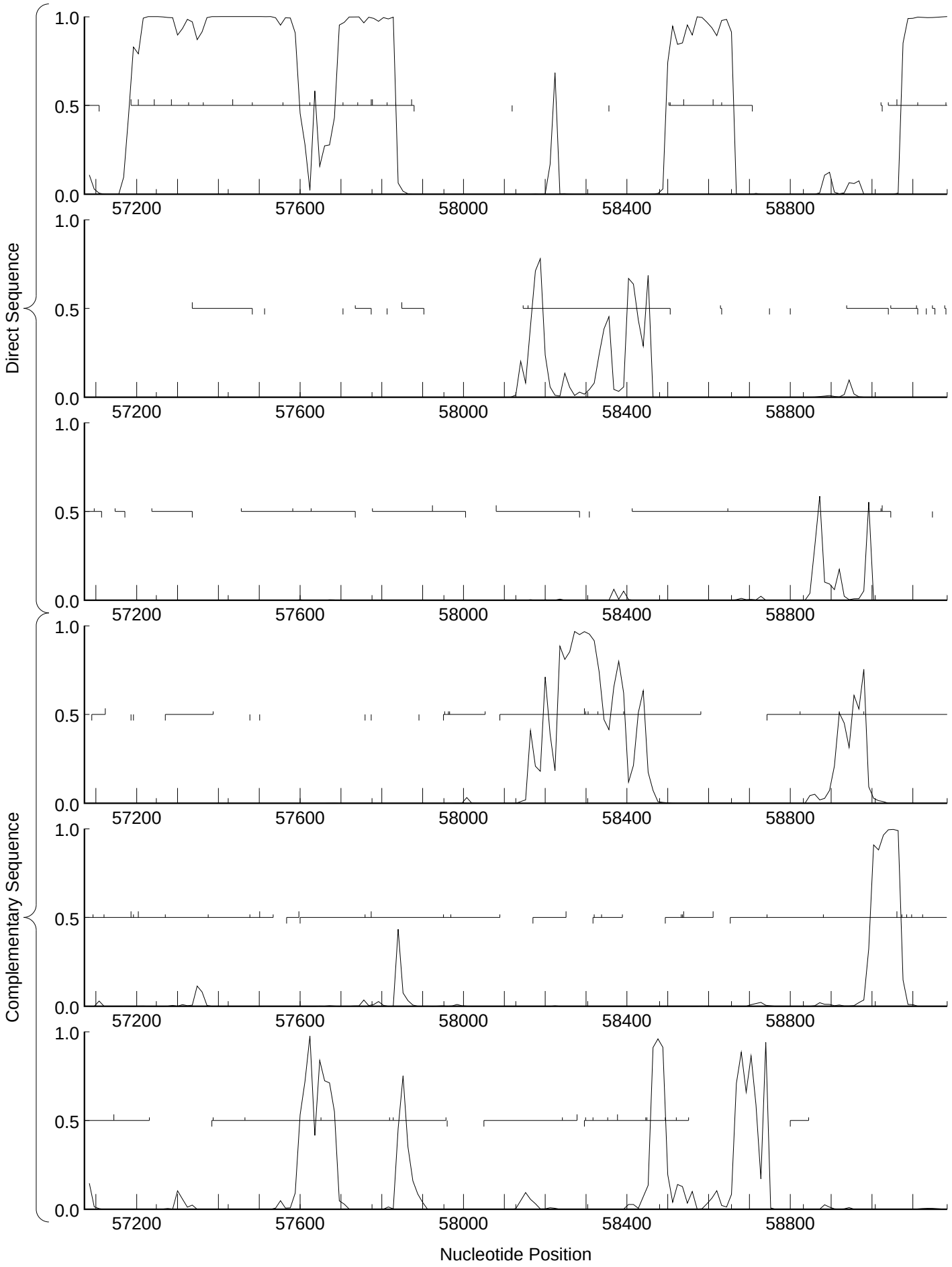

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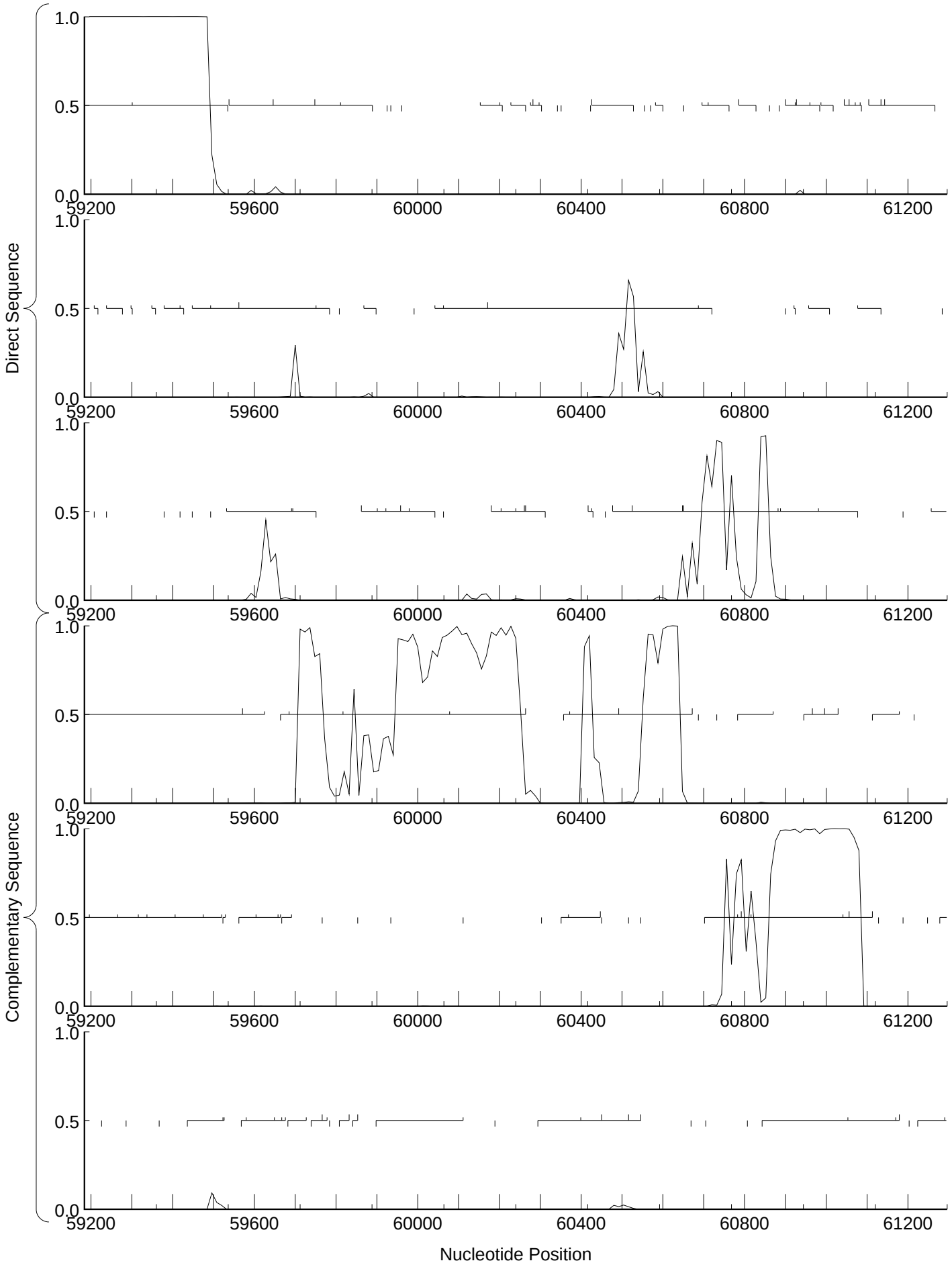

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Nucleotide Position


Direct Sequence
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0.5
0.0
55200
55600
56000
56400
56800
Nucleotide Position

ge Jflix2 complete sequence, 62017 bp including 6-base 3’ overhang (CCCGCT), Cluster CF, Order 4, Window 96, Step 12, 29/31



ge Jflix2 complete sequence, 62017 bp including 6-base 3’ overhang (CCCGCT), Cluster CF, Order 4, Window 96, Step 12, 30/31



ge Jflix2 complete sequence, 62017 bp including 6-base 3’ overhang (CCCGCT), Cluster CF, Order 4, Window 96, Step 12, 31/31

