GeneMark

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

PROGRAM INFORMATION

```
Sequence         :  Mycobacterium phage Scoot17C
Analysis Date    :  3/29/24 at 11:52:25
Pages            :  34
Sequence Length  :  68432 bp
GC Content       :   66.50%

Window Length    :  96 bp
Window Step      :  12 bp
Threshold Value  :  0.500

PS-Version       :  1.2

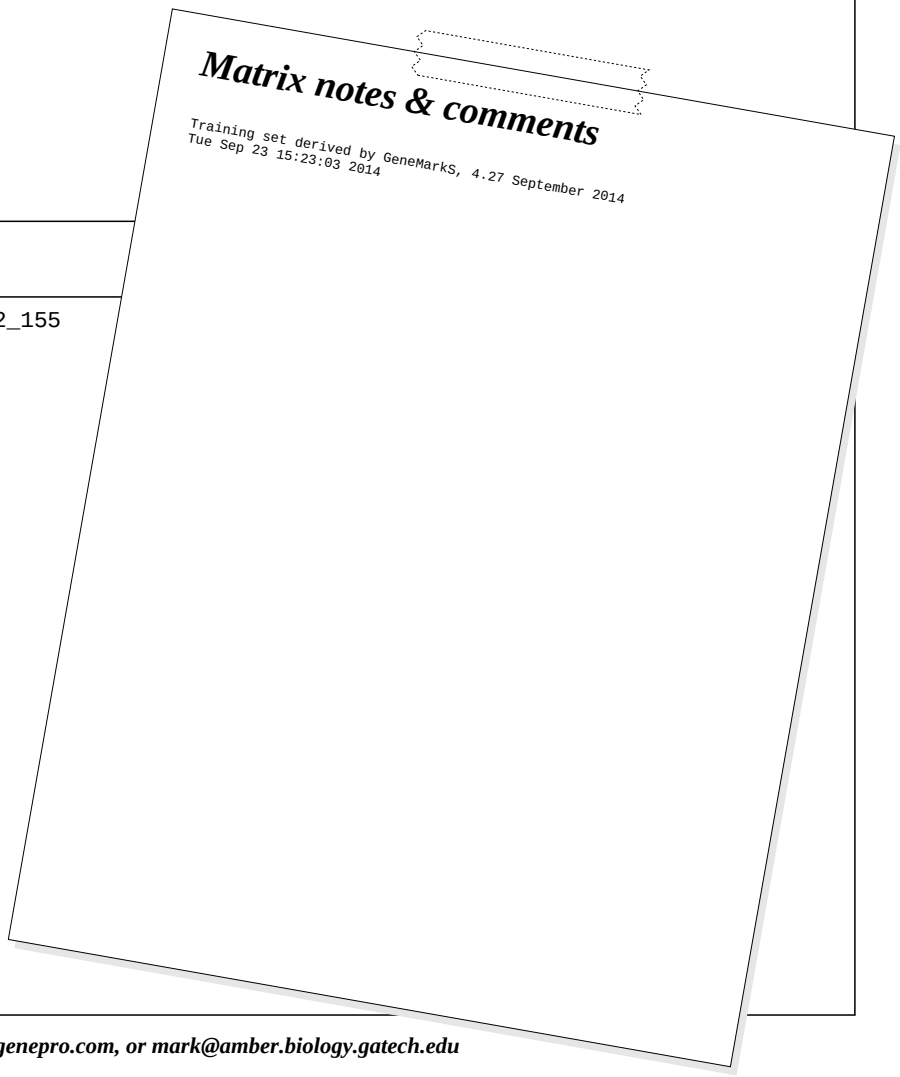
GeneMark Options :  PostScript graph,
                    Mark ORFs / splice sites,
                    List ORFs,
                    List regions and/or splice sites,
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MATRIX INFORMATION

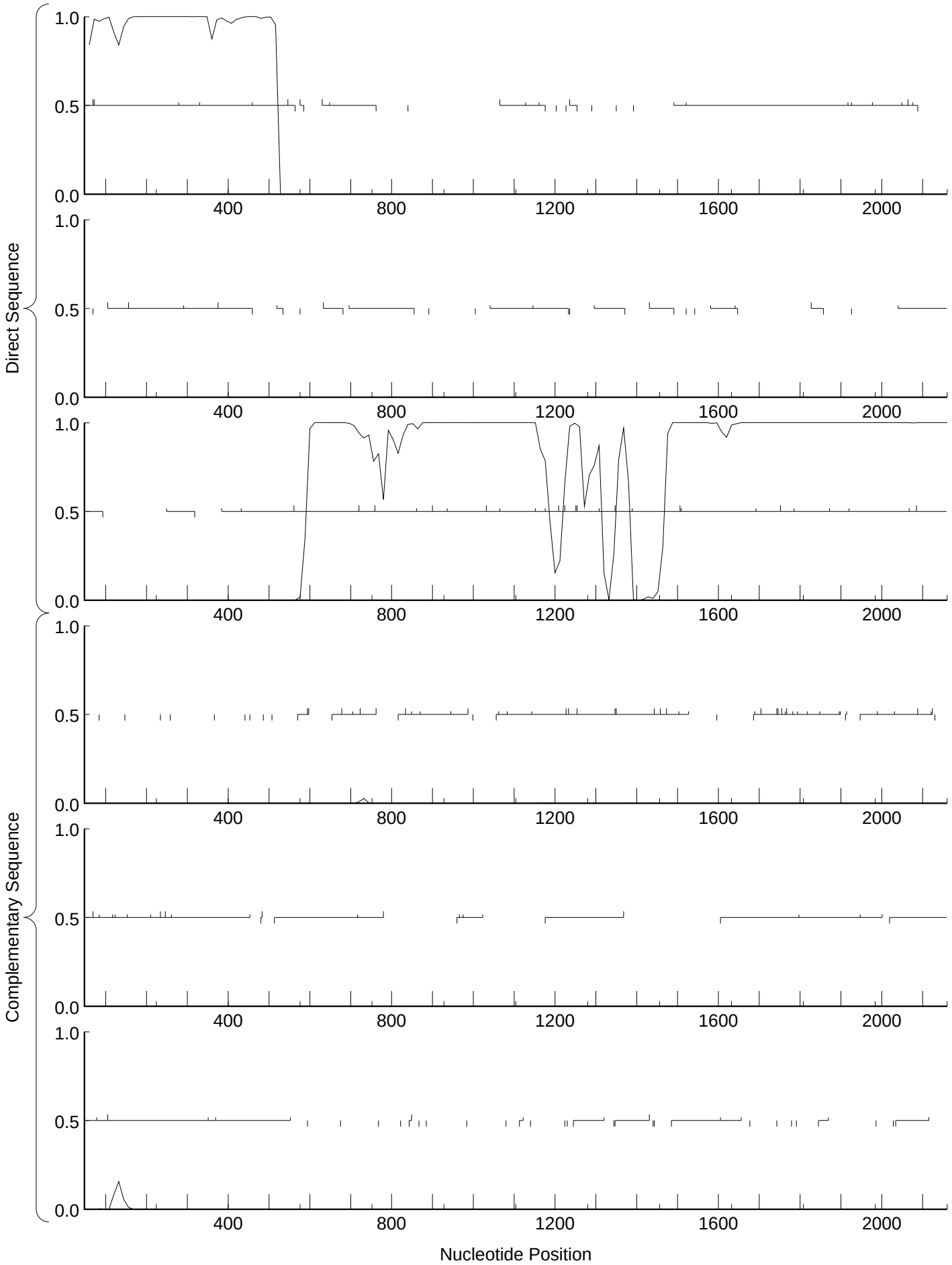
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Matrix     :  Mycobacterium_smegmatis_MC2_155
Author     :  -

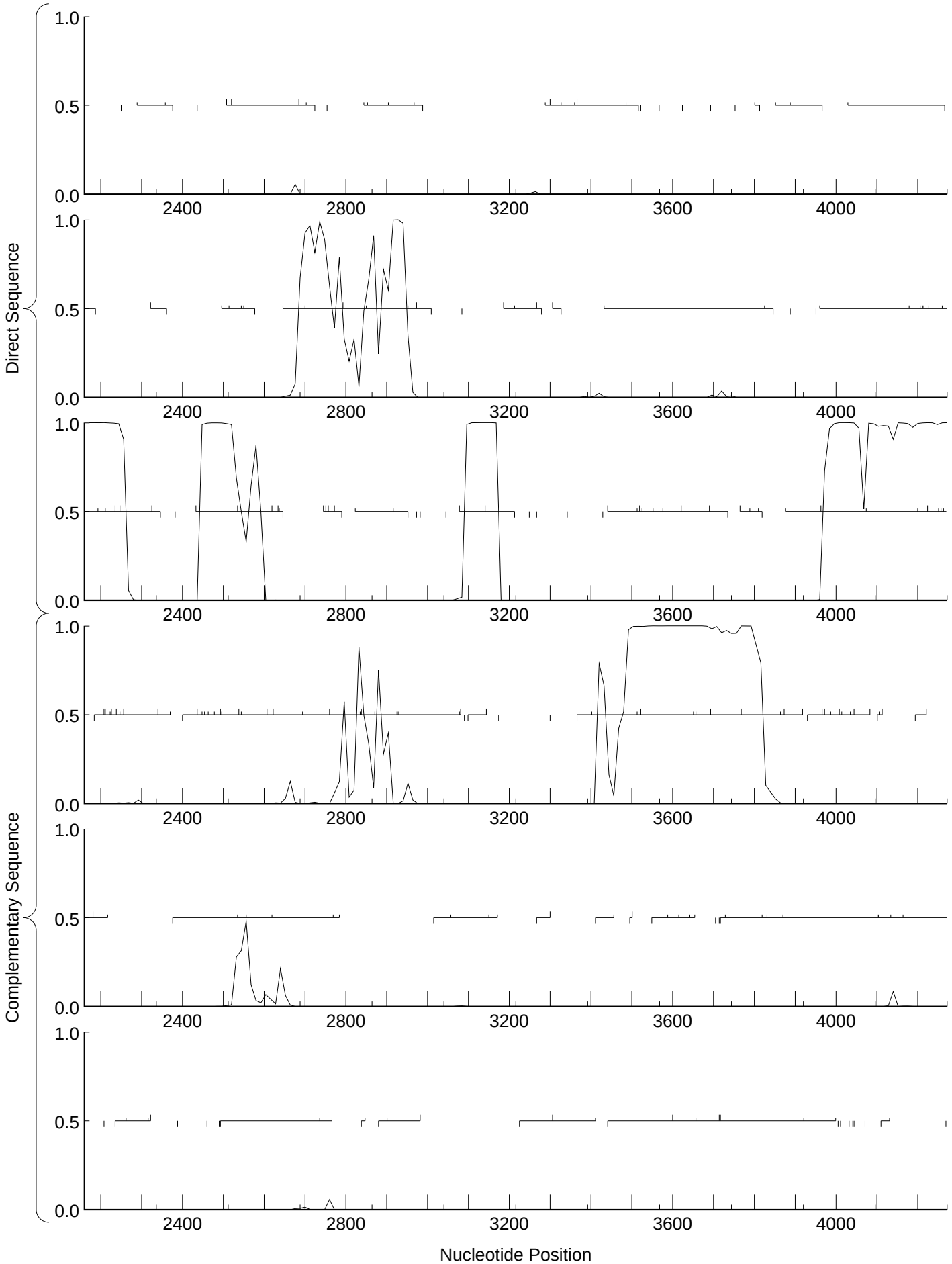
Order      :  4

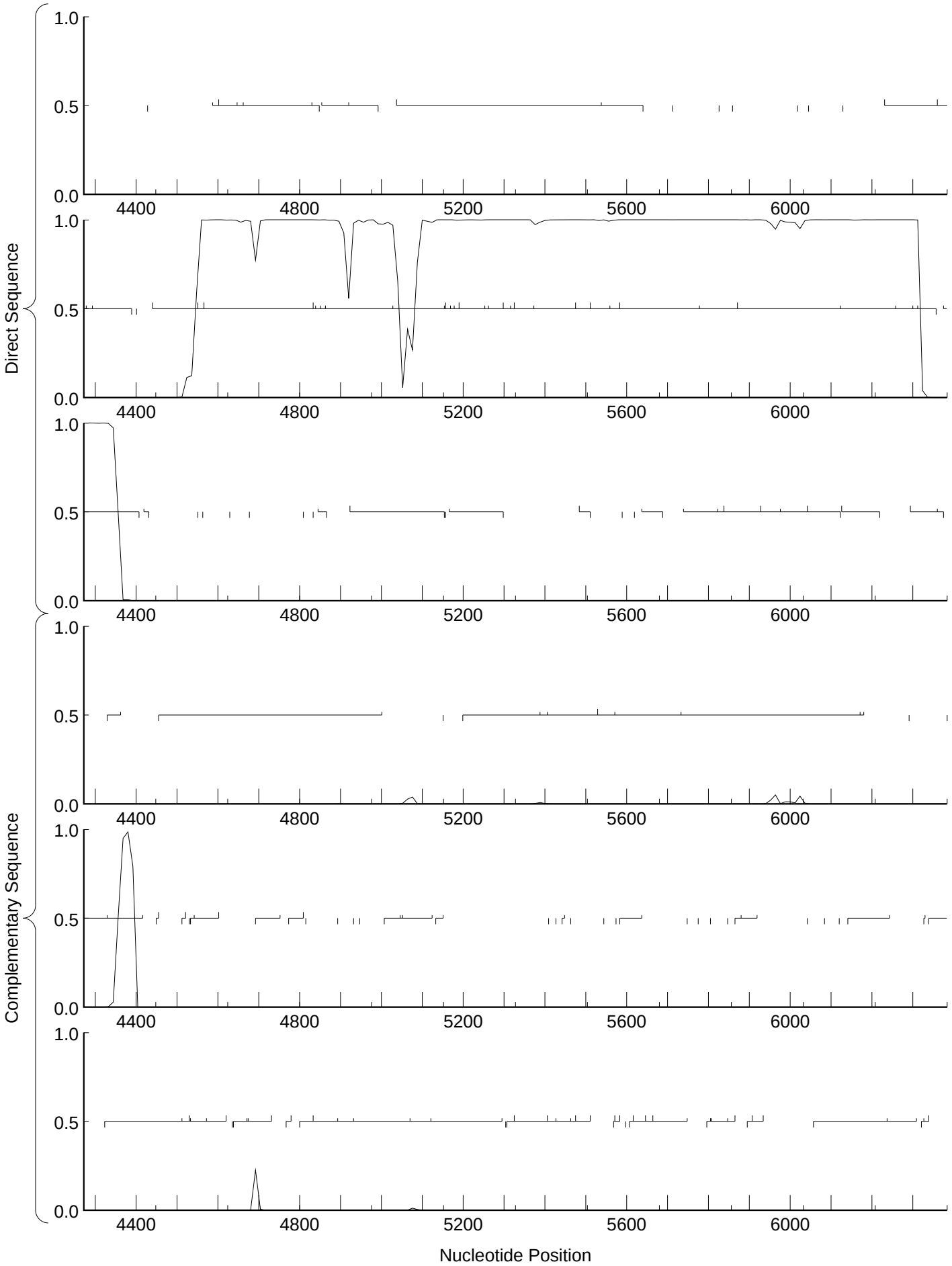

  Send questions / comments to:
      Dr. M. Borodovsky
      Georgia Institute of Technology
      School of Biology
      Atlanta, GA  30332-0230
```

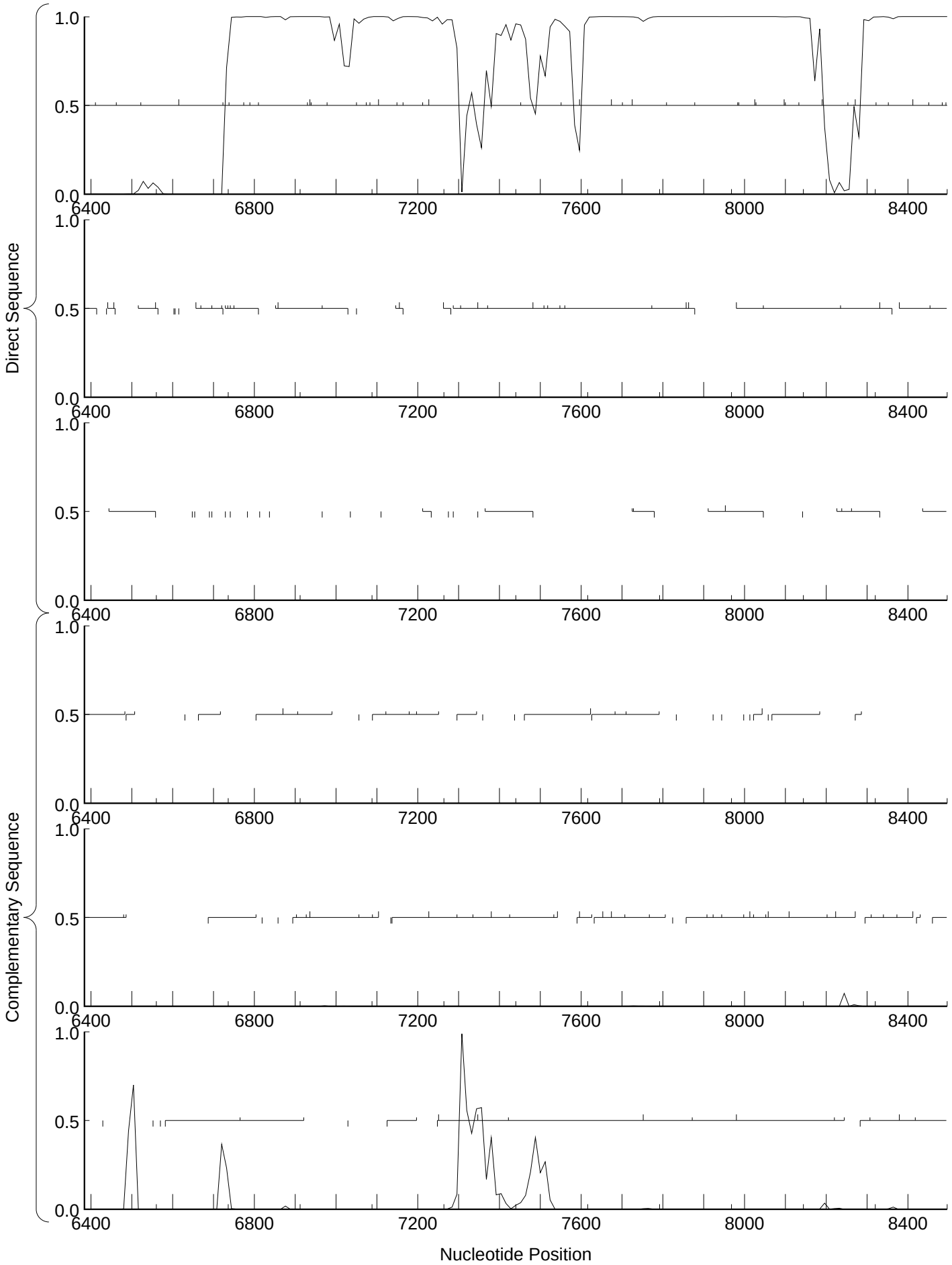


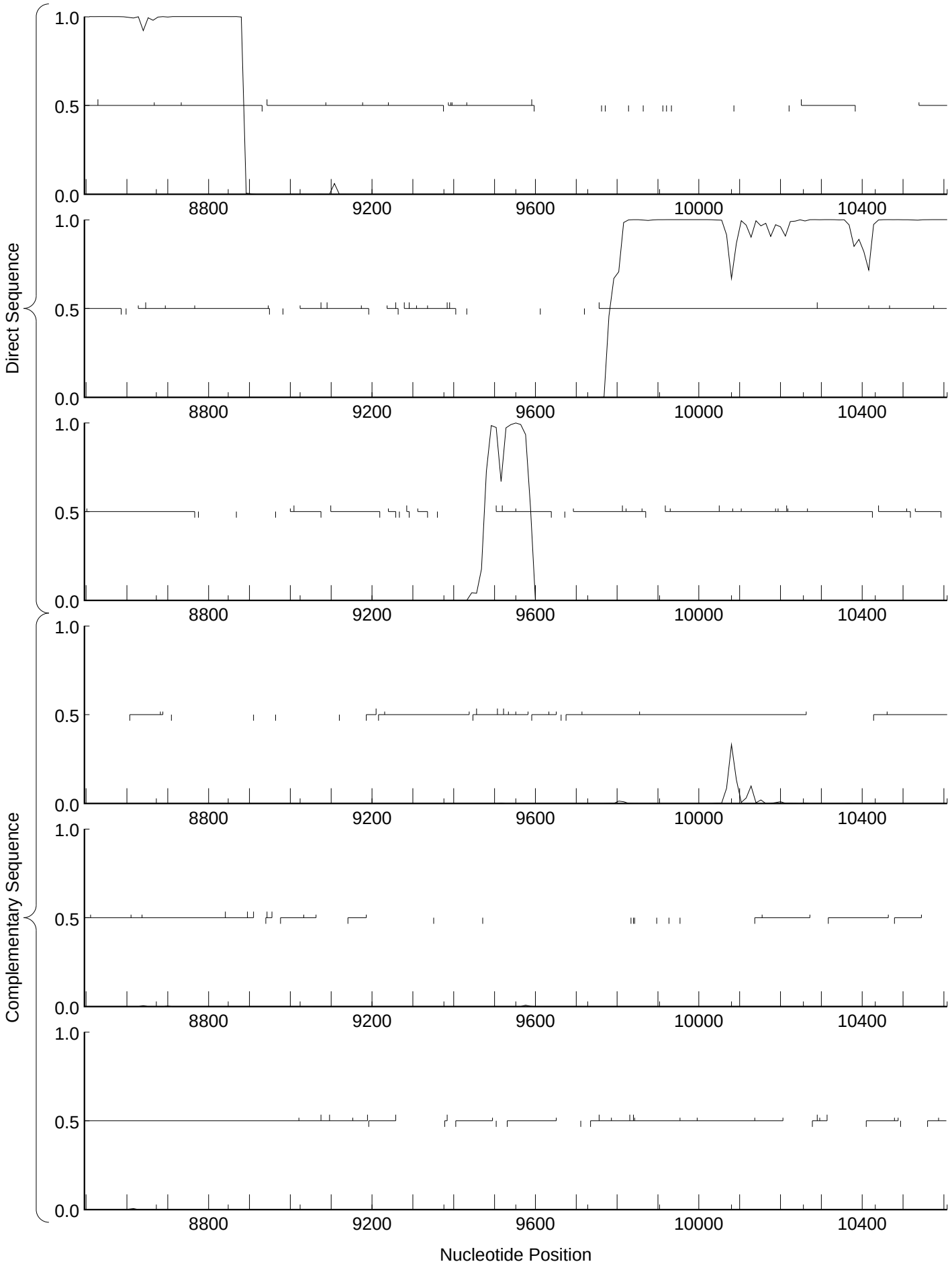
For help or information, send electronic mail to: custserv@genepro.com, or mark@amber.biology.gatech.edu

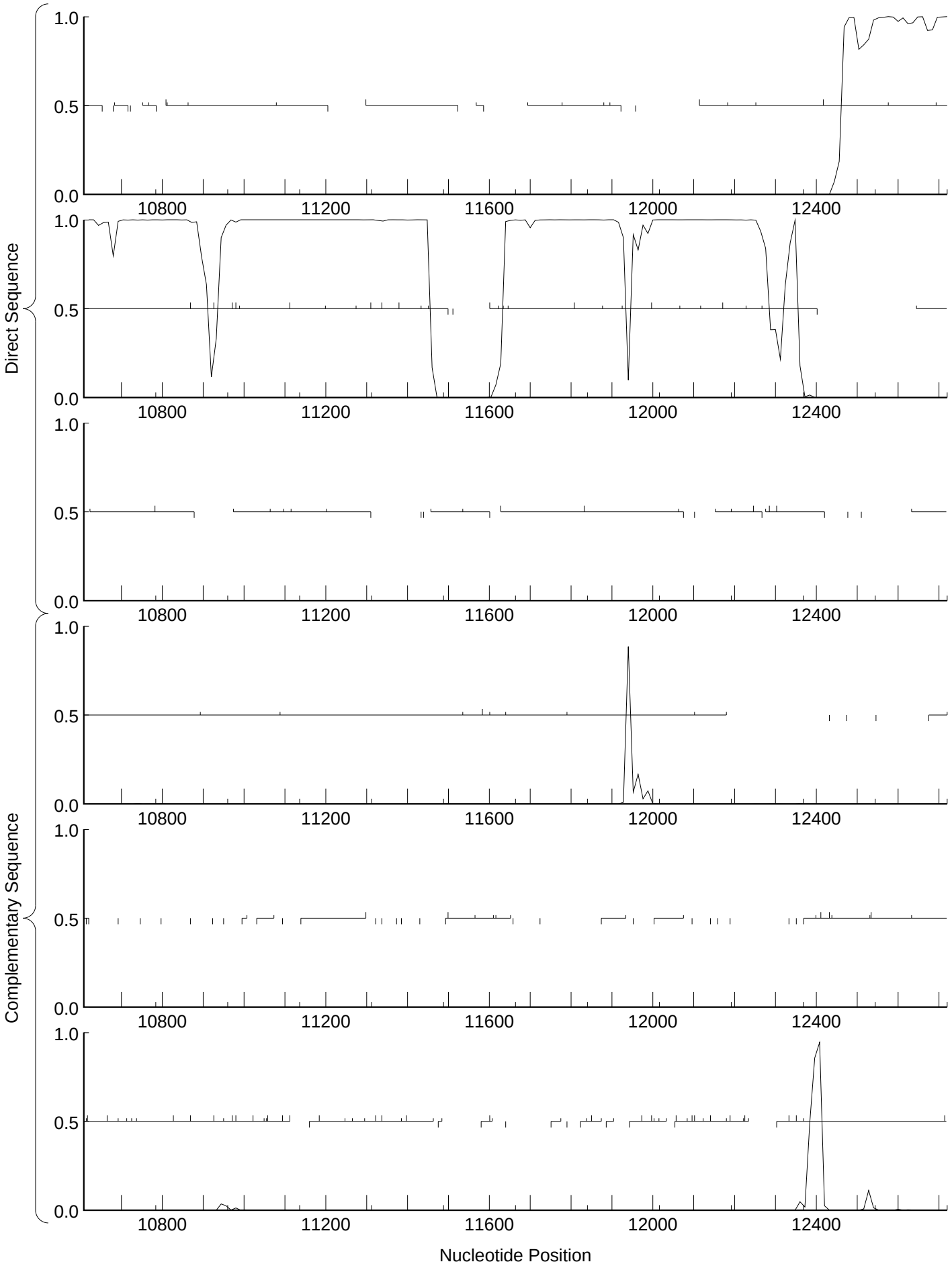

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Complementary Sequence
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1200
1600
2000
Nucleotide Position

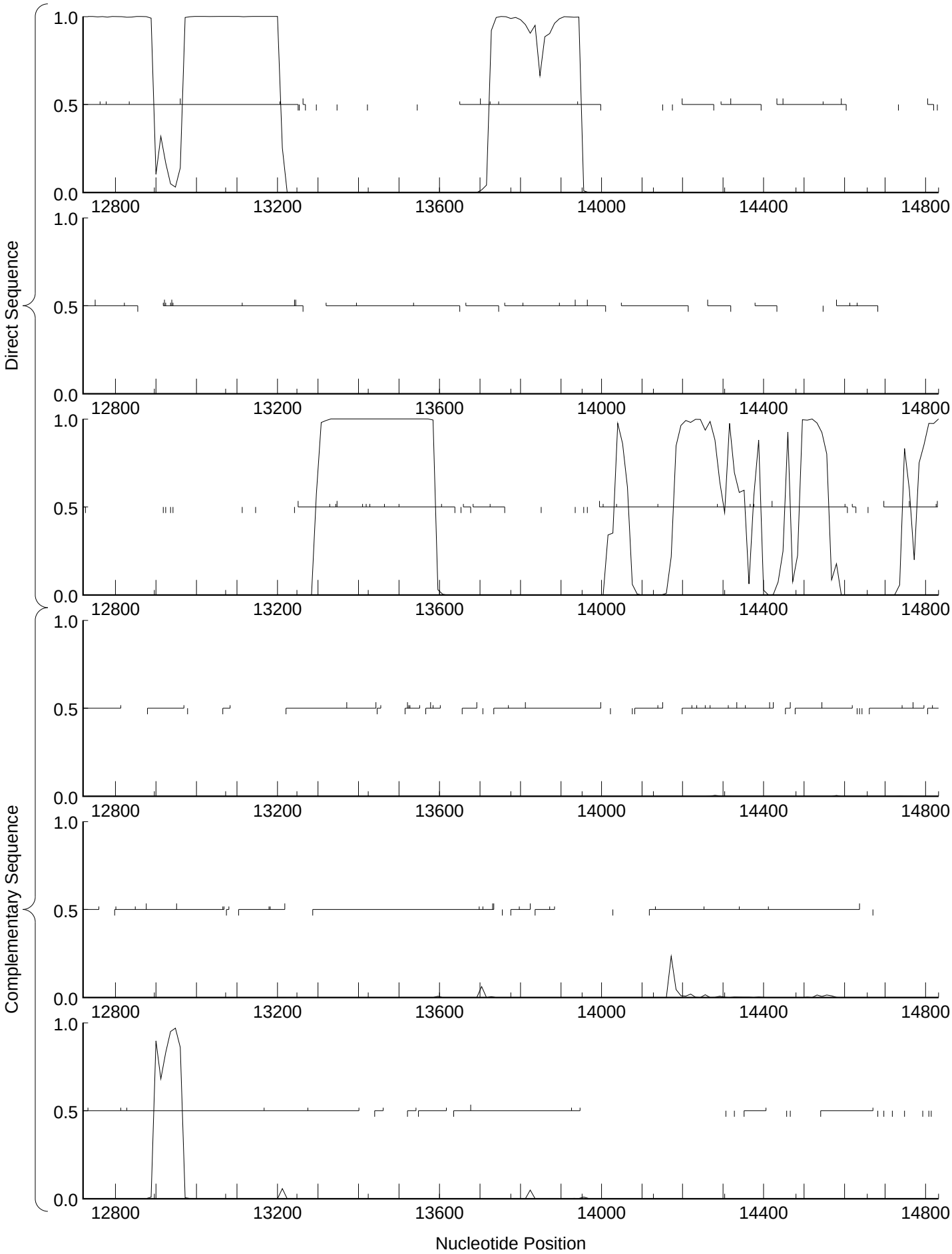

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Nucleotide Position

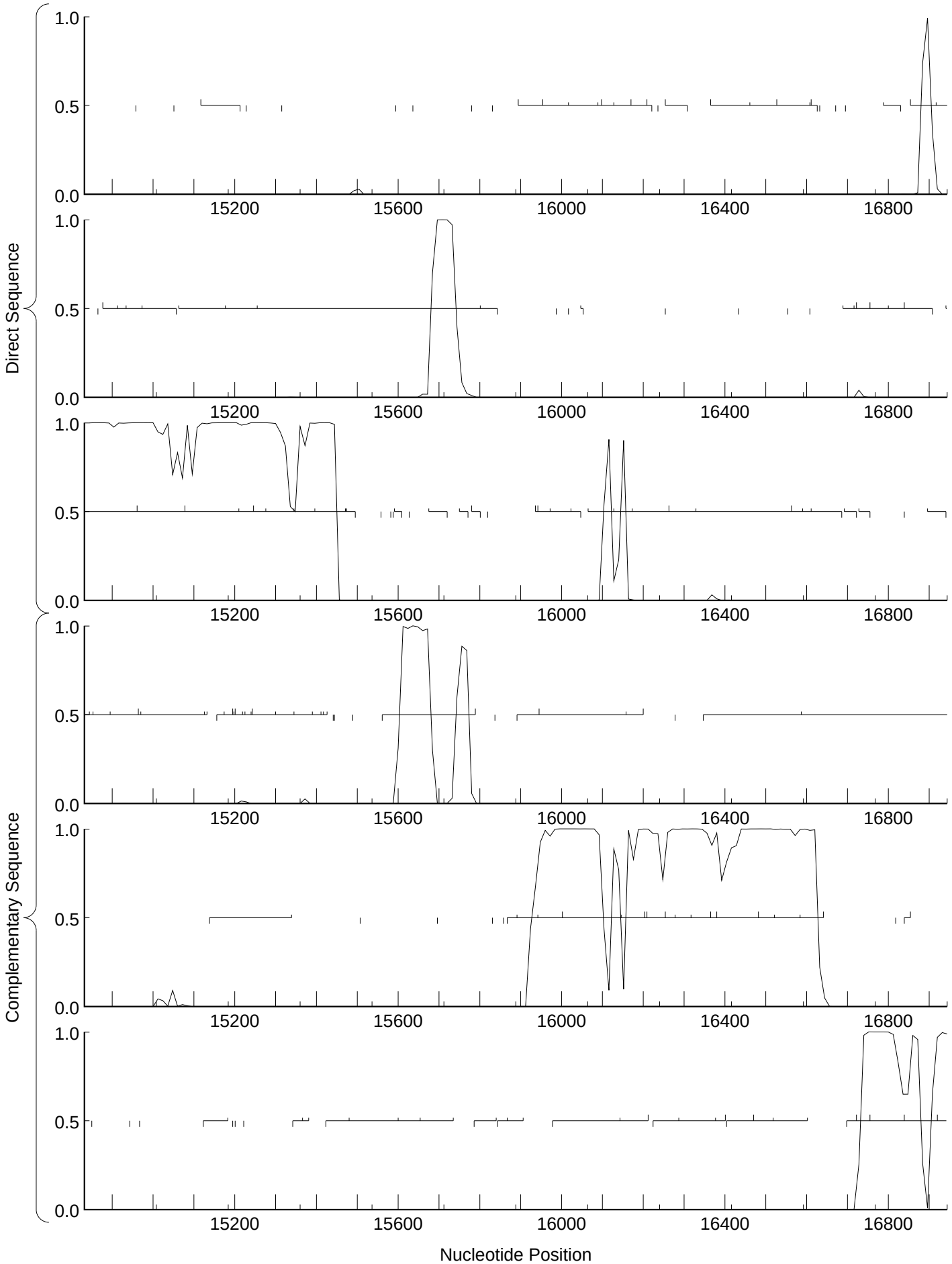

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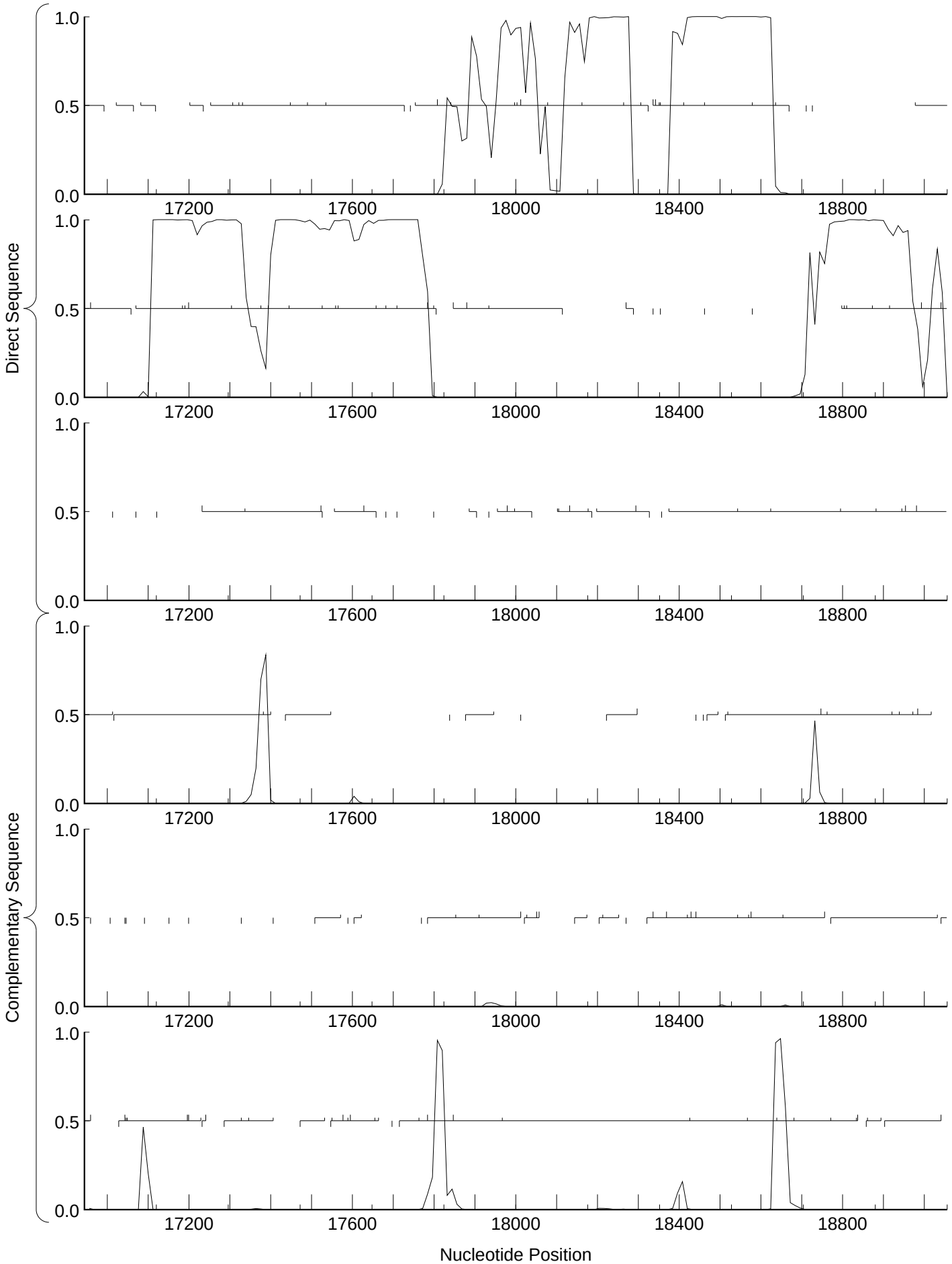

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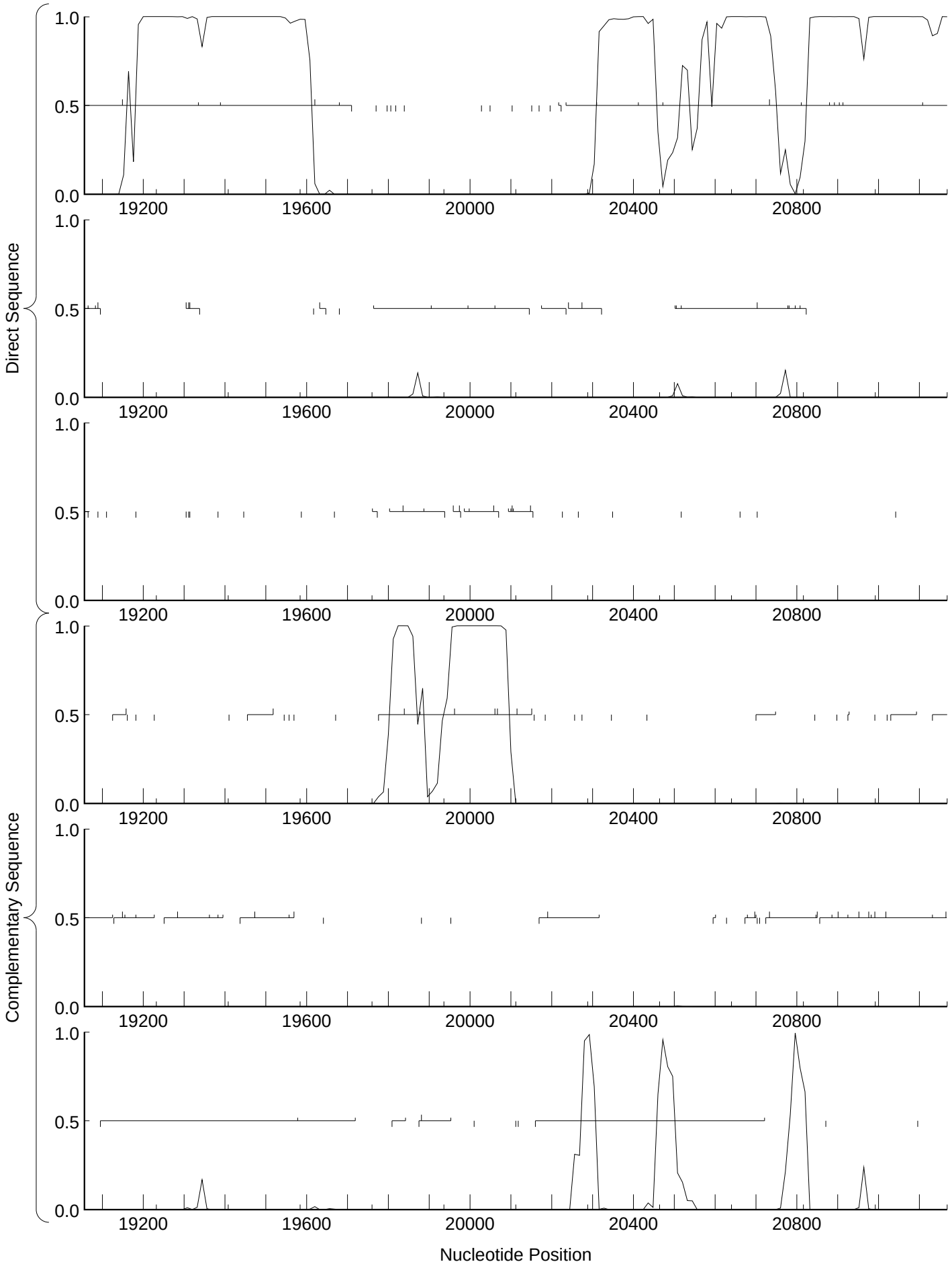

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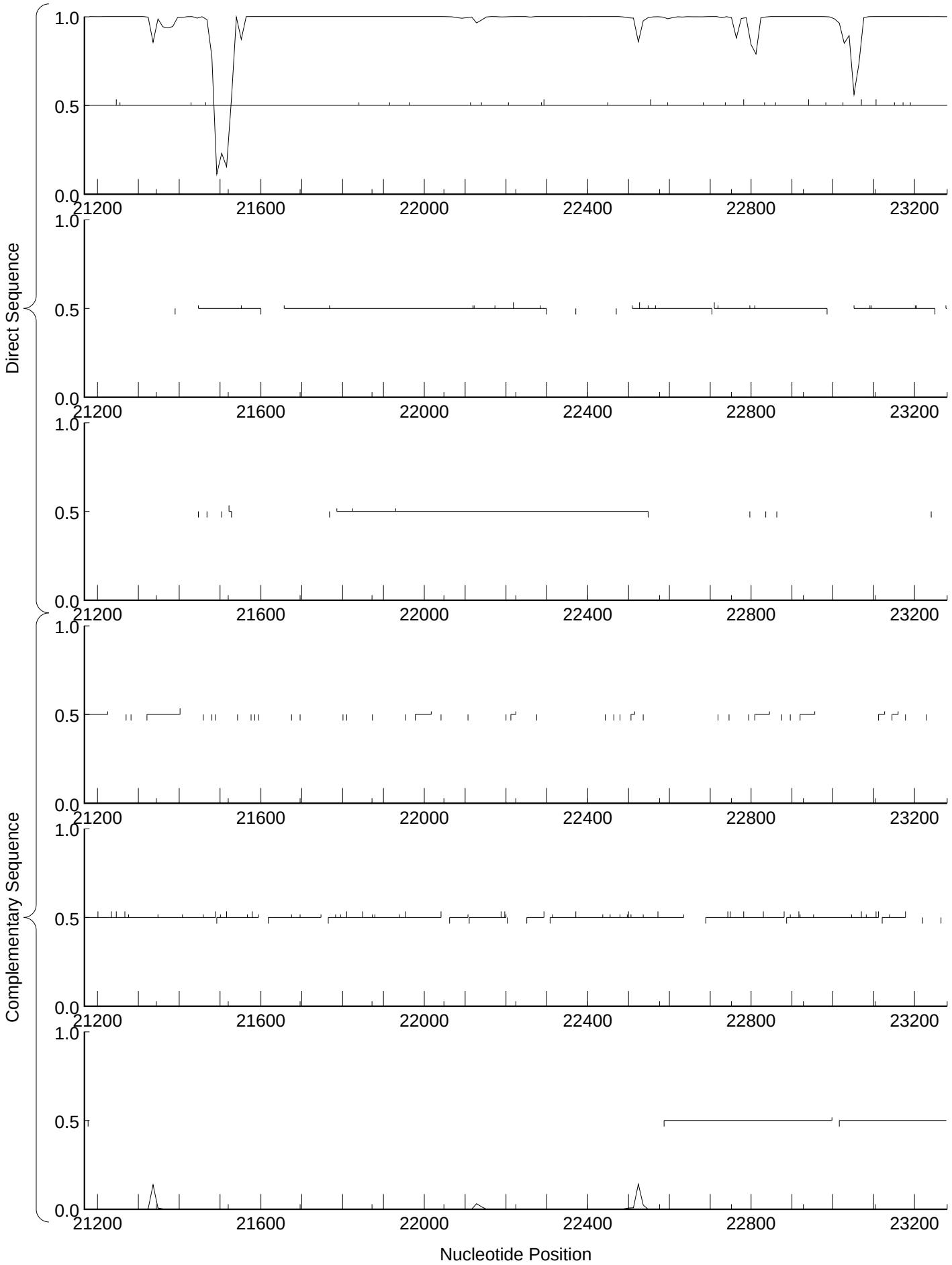

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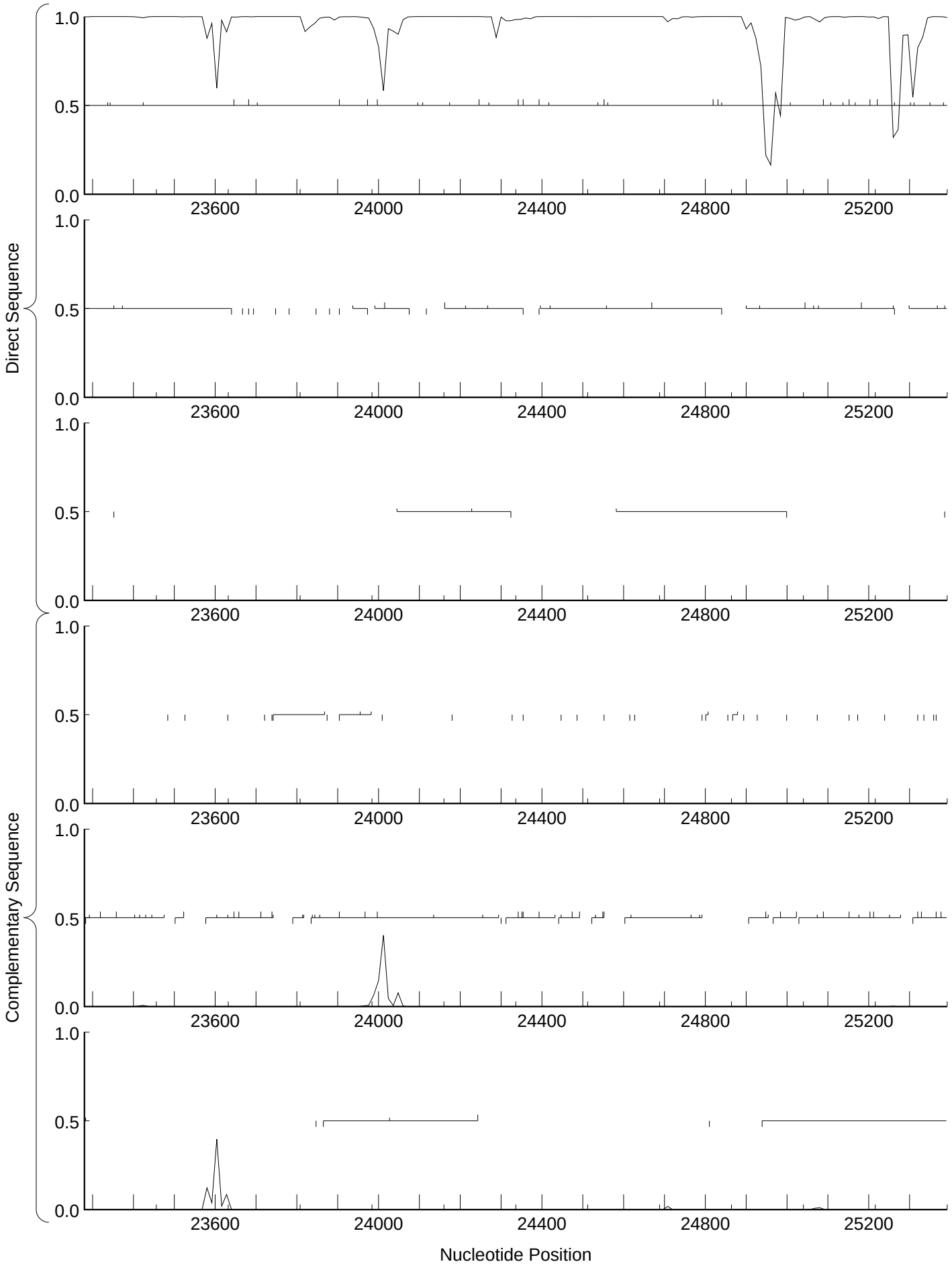

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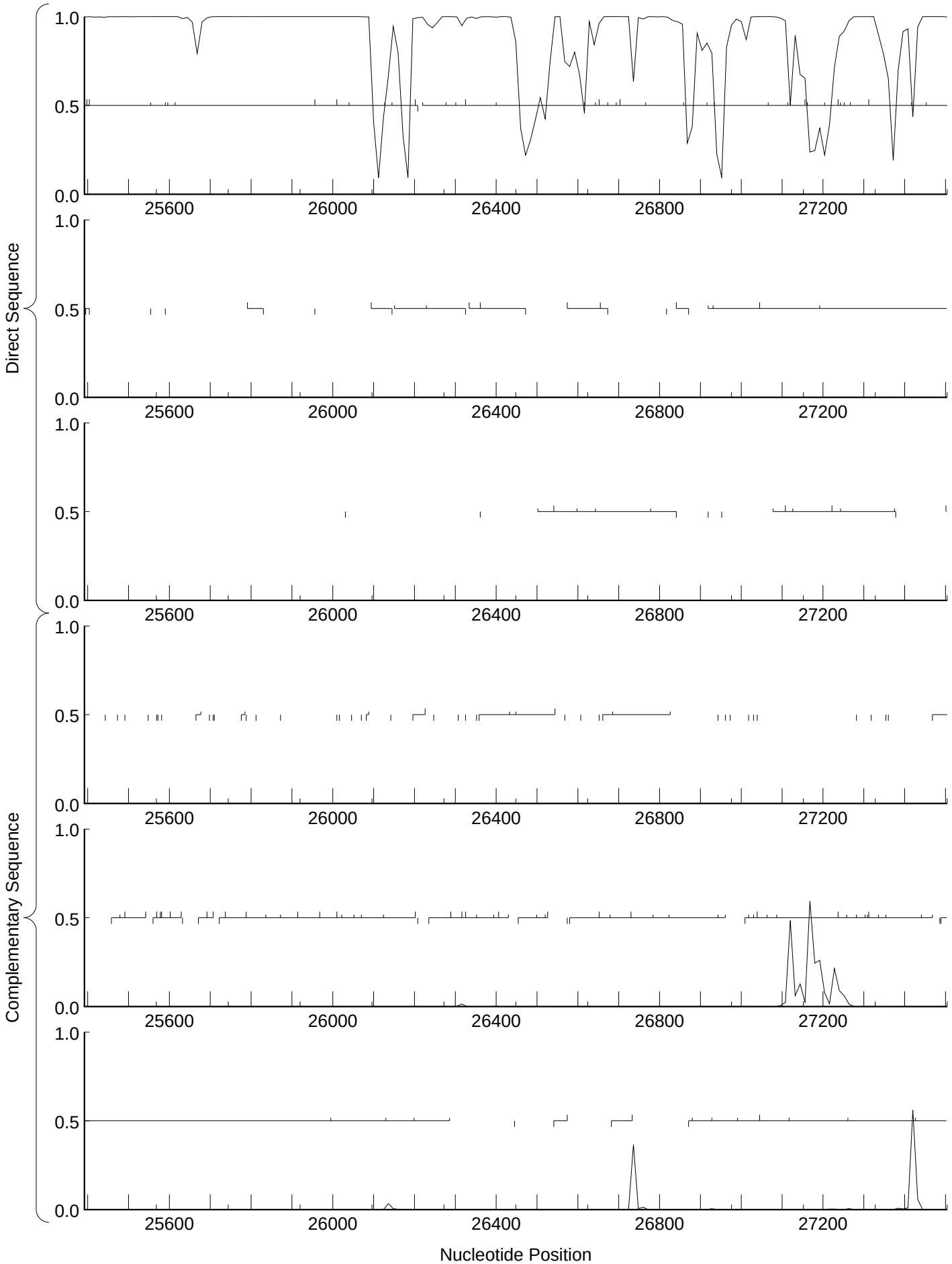

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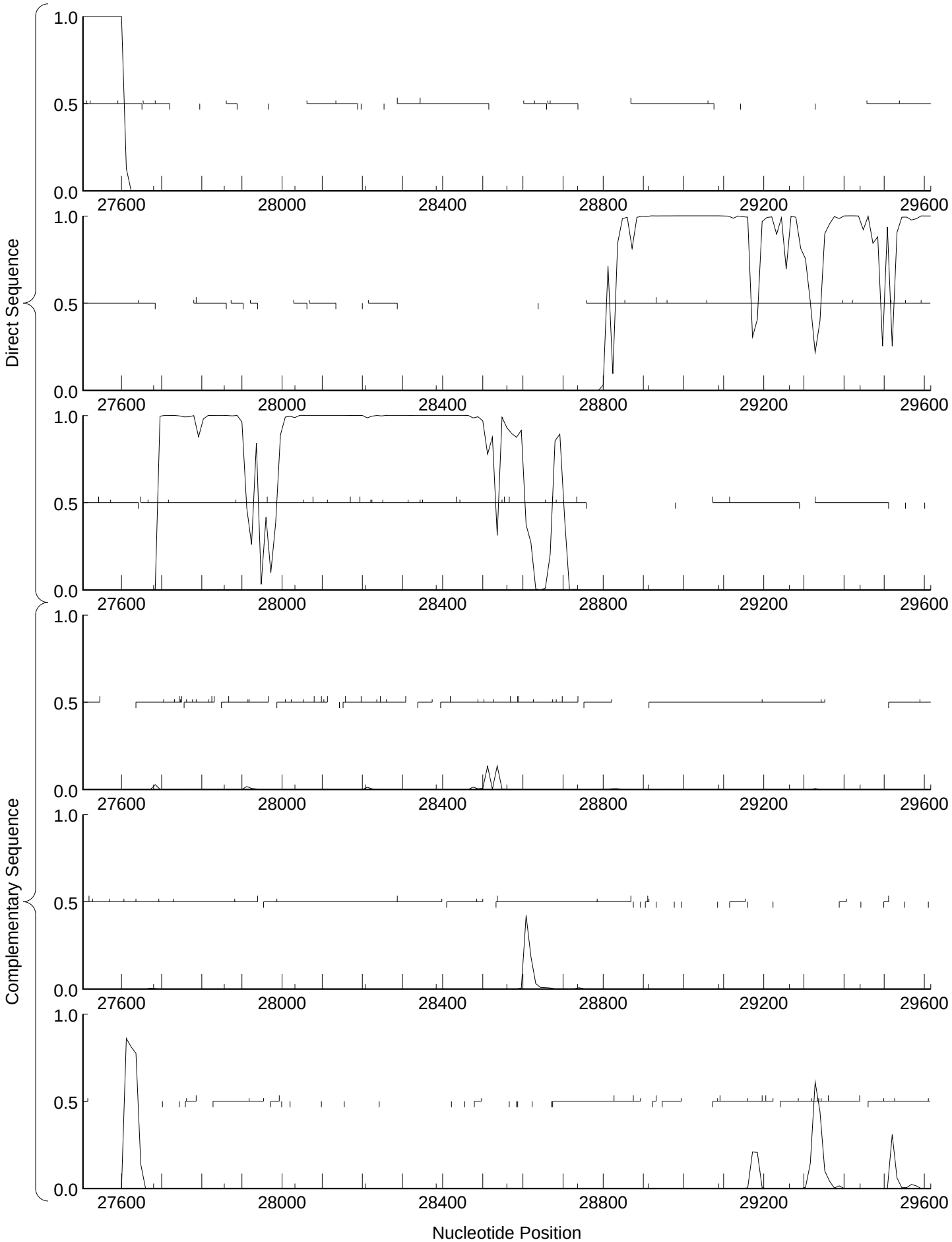

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Nucleotide Position

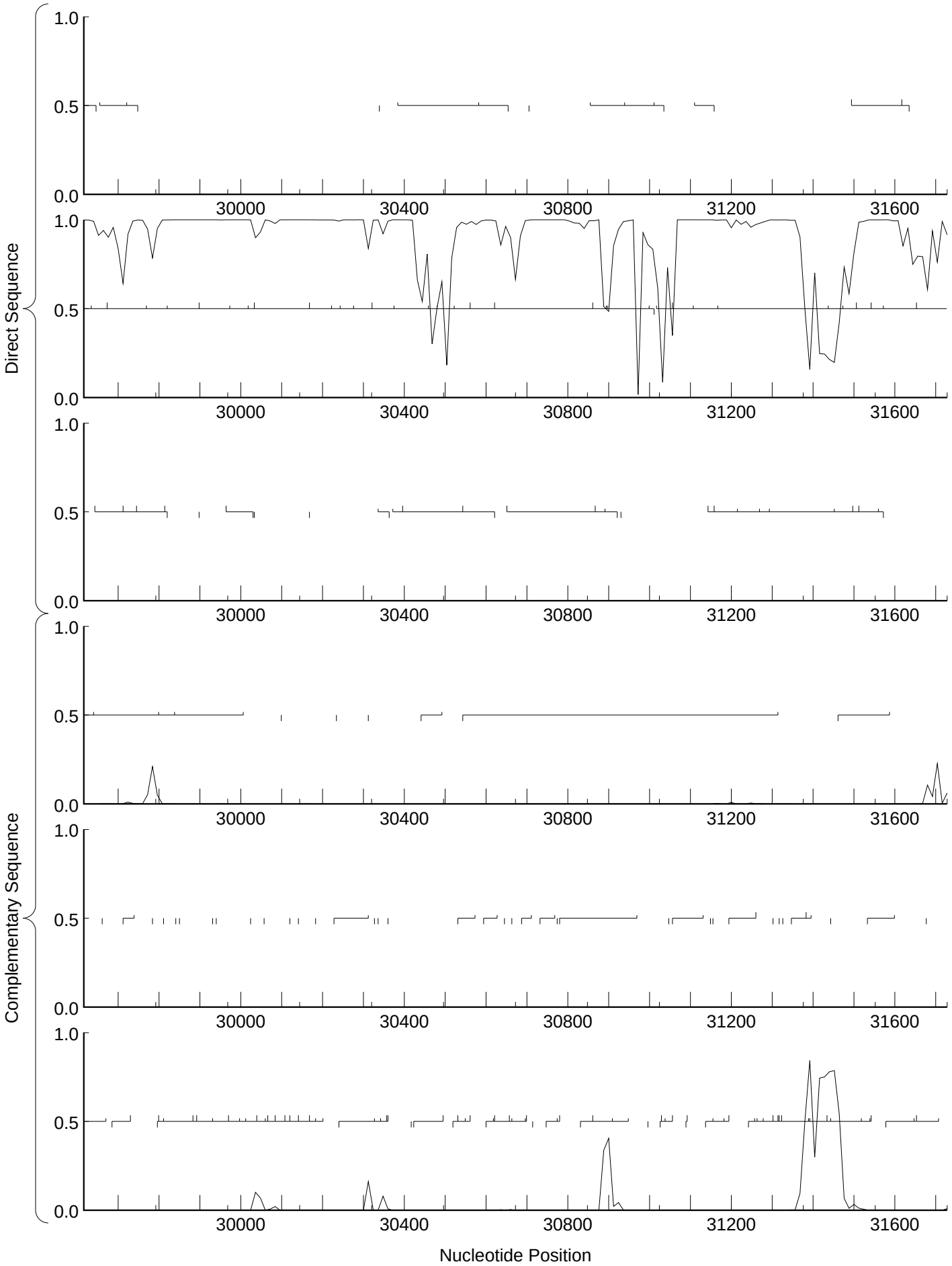

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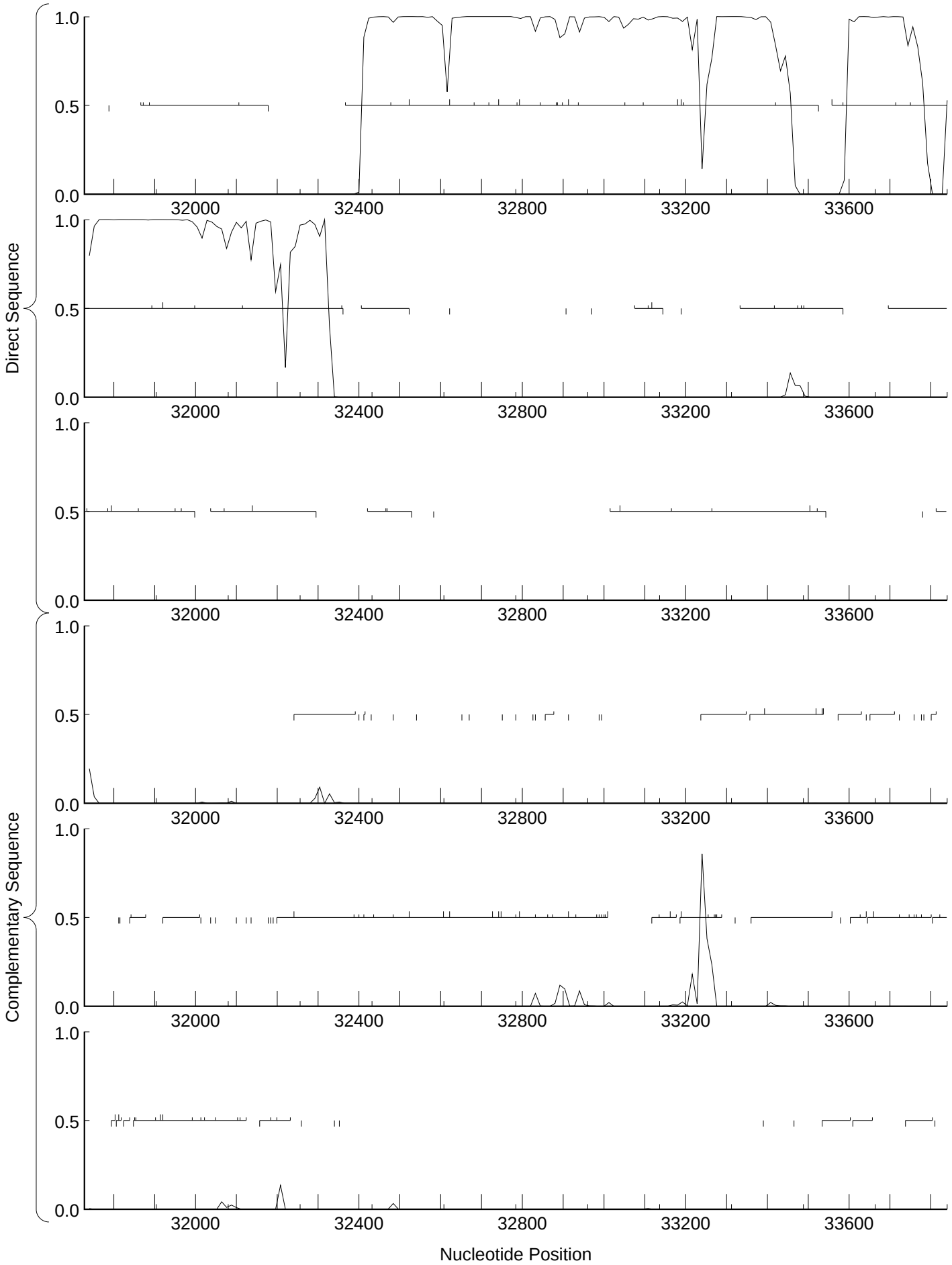

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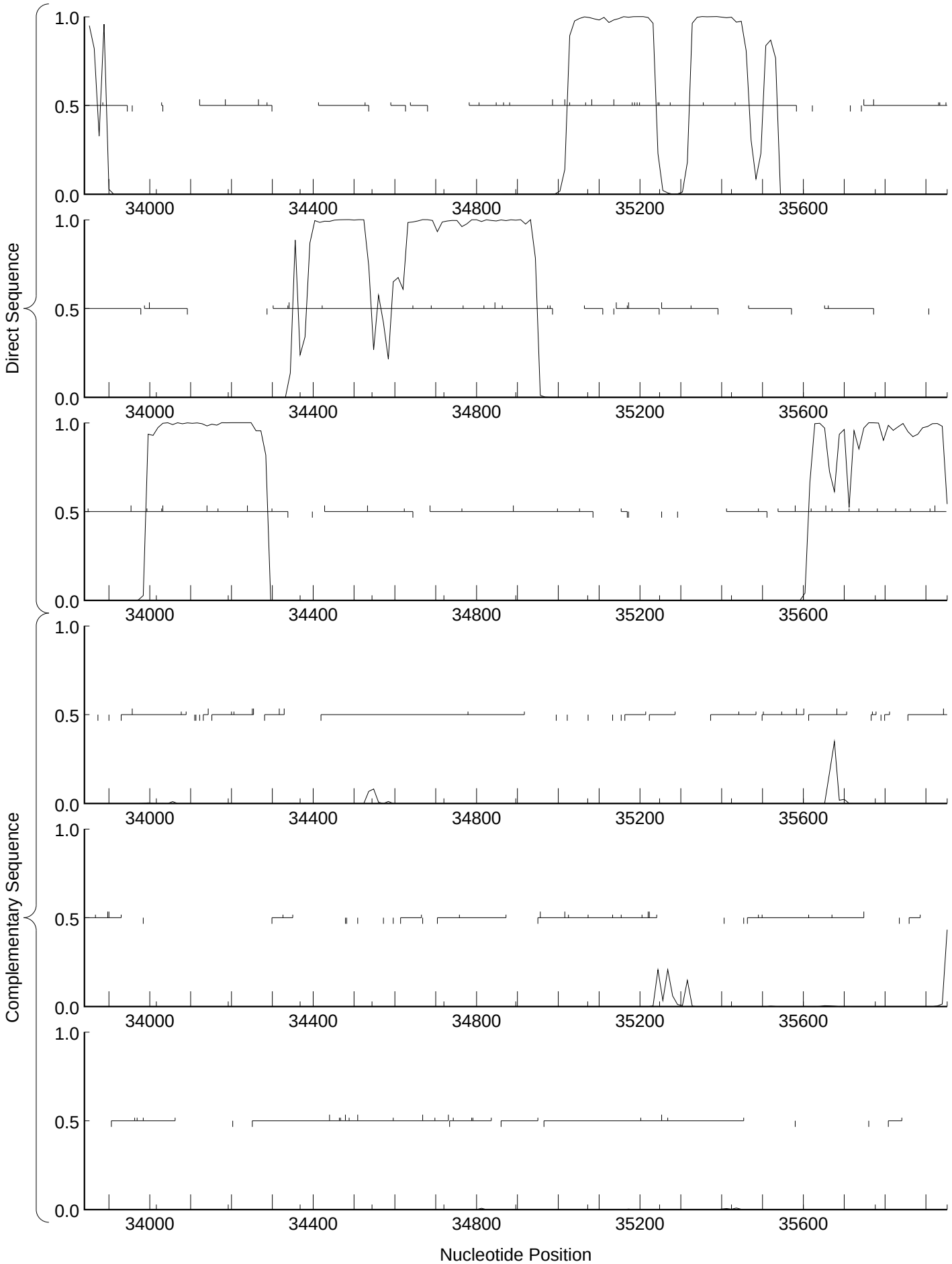

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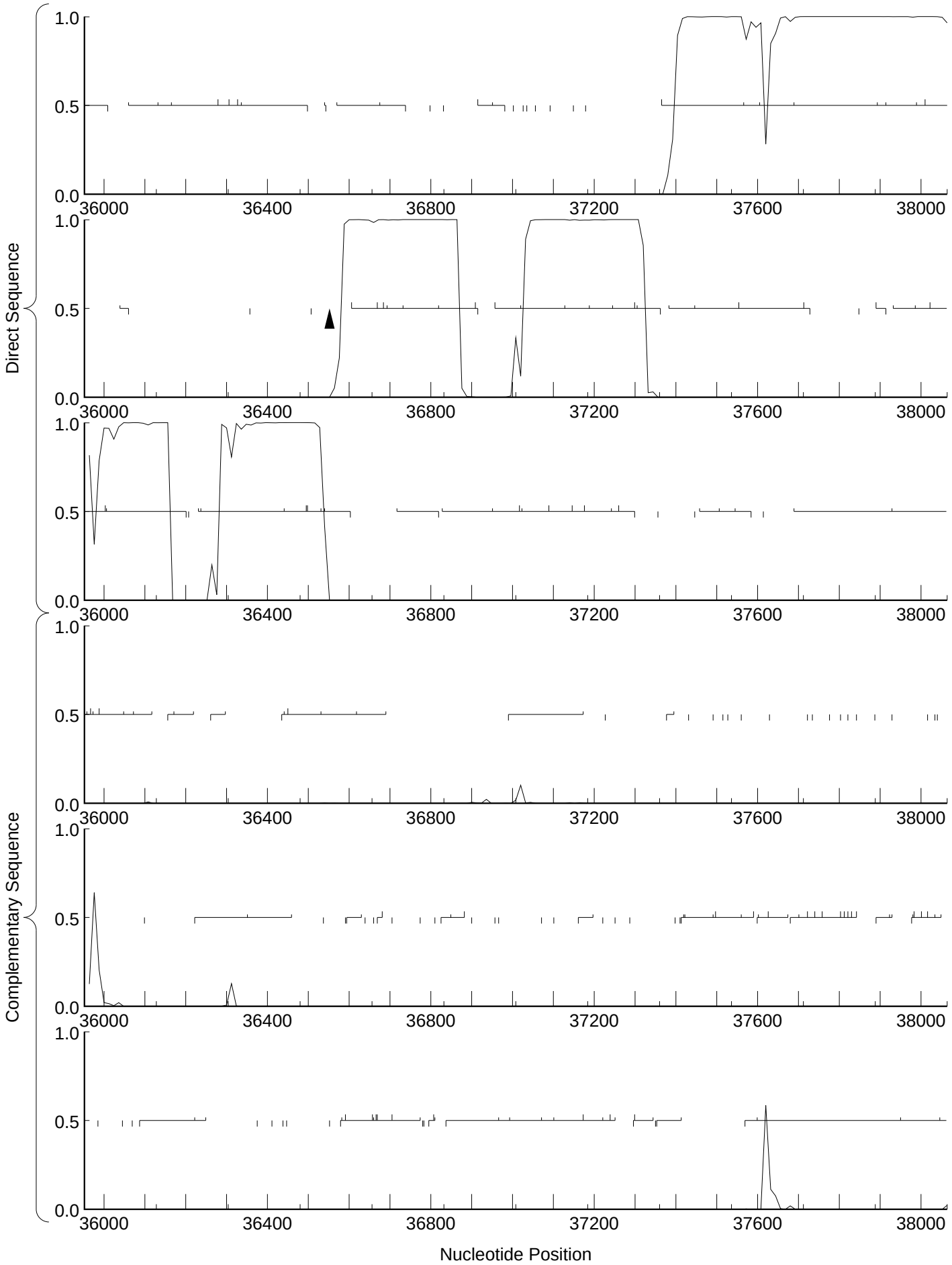

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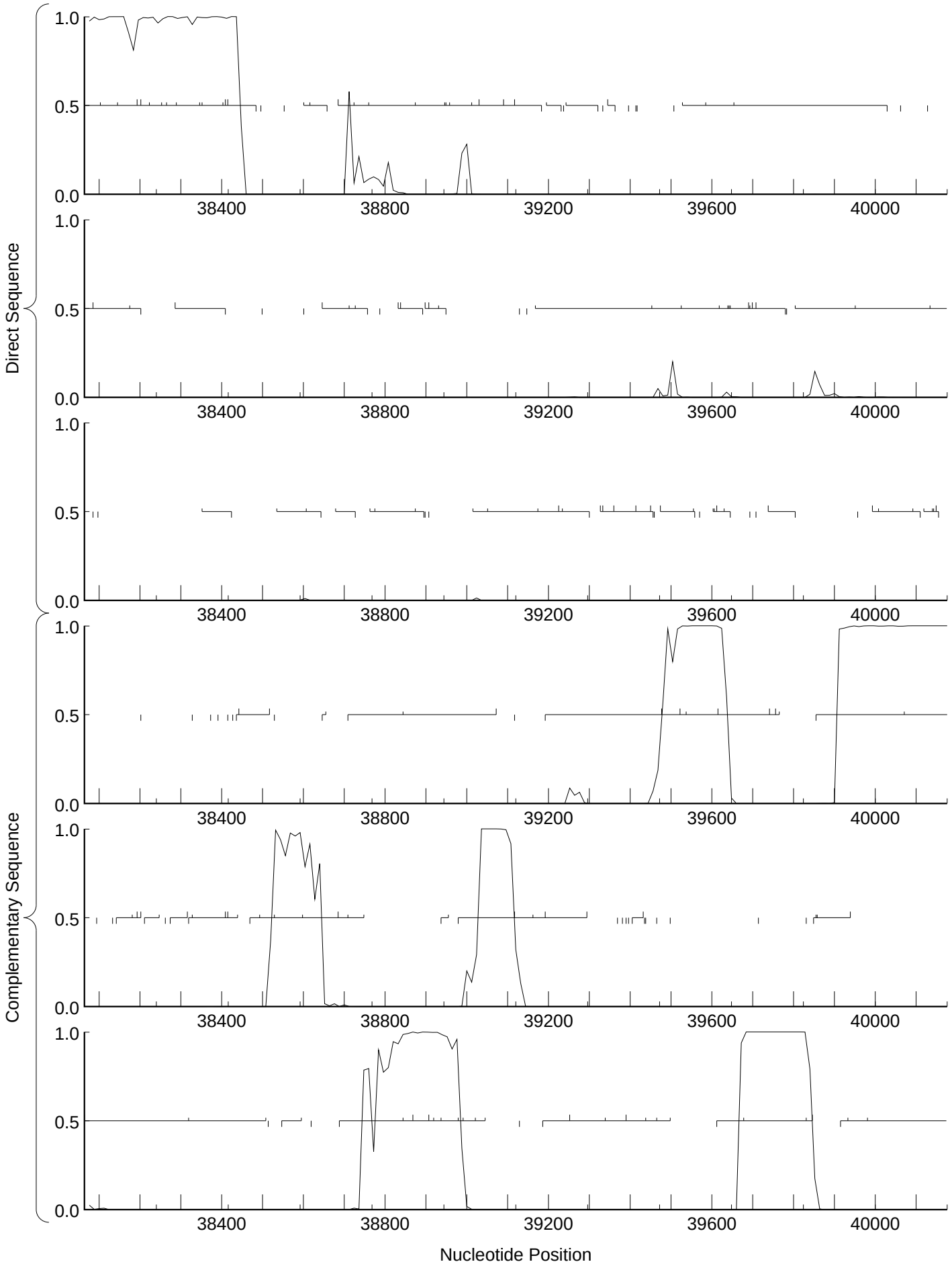

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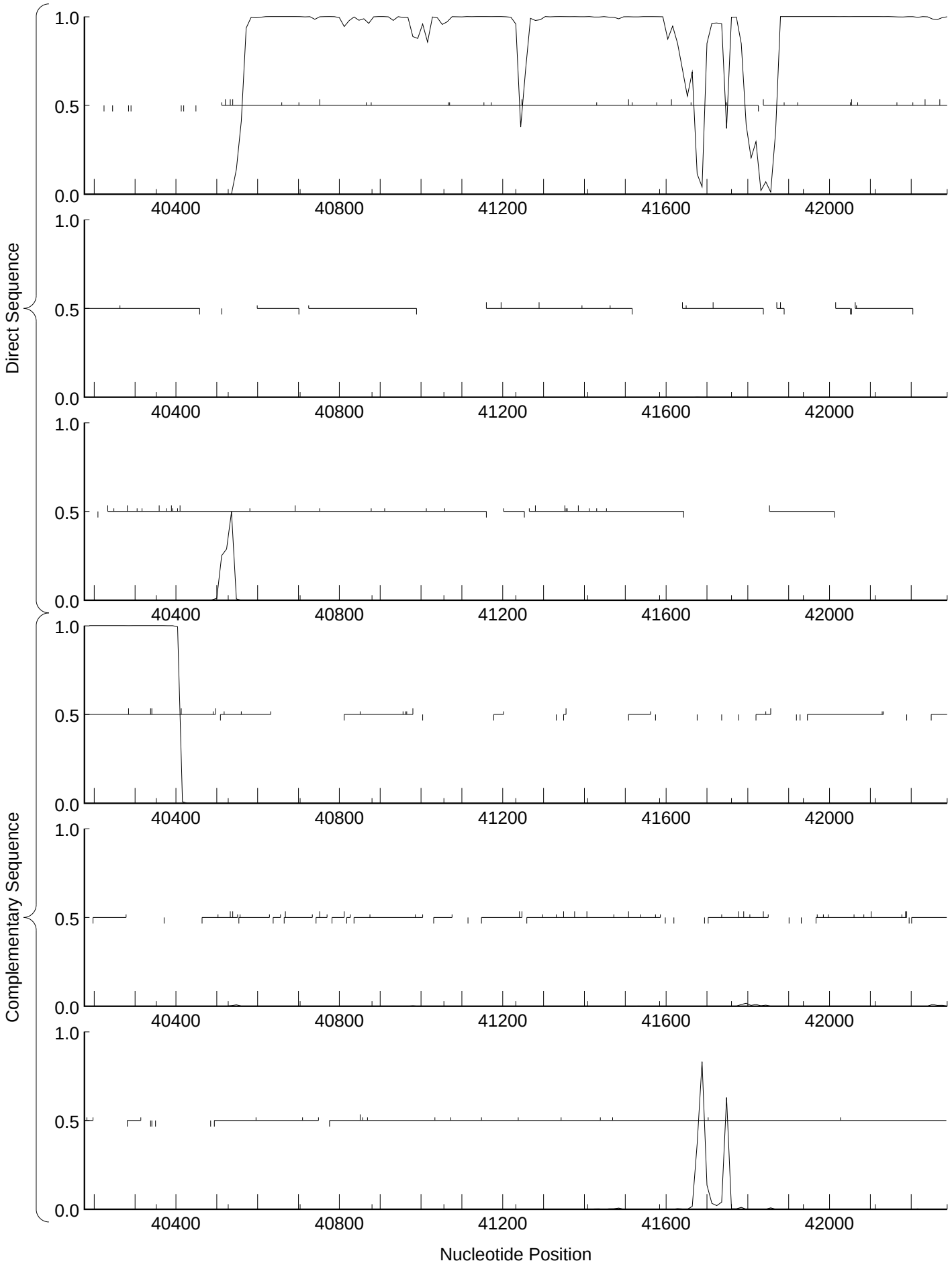

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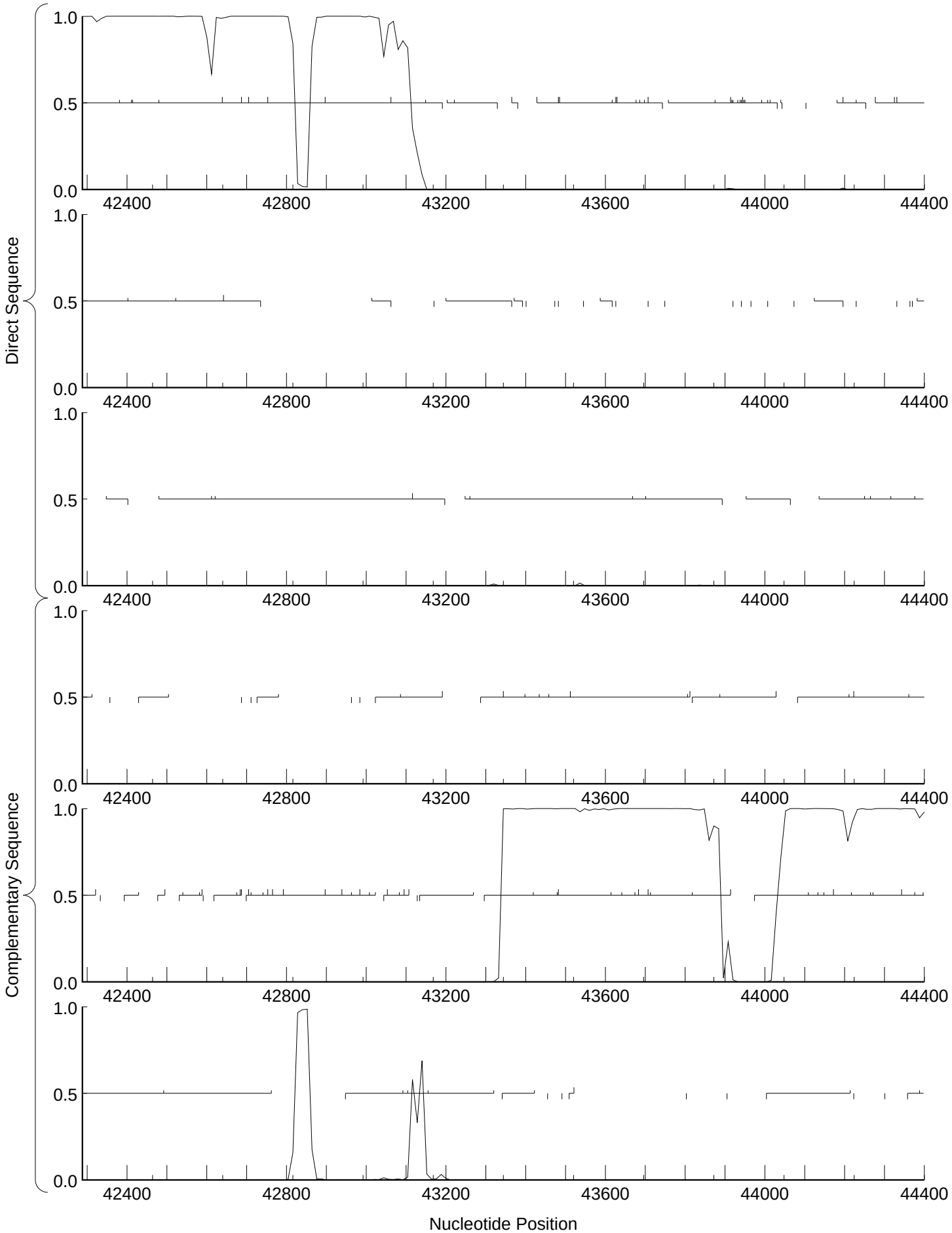

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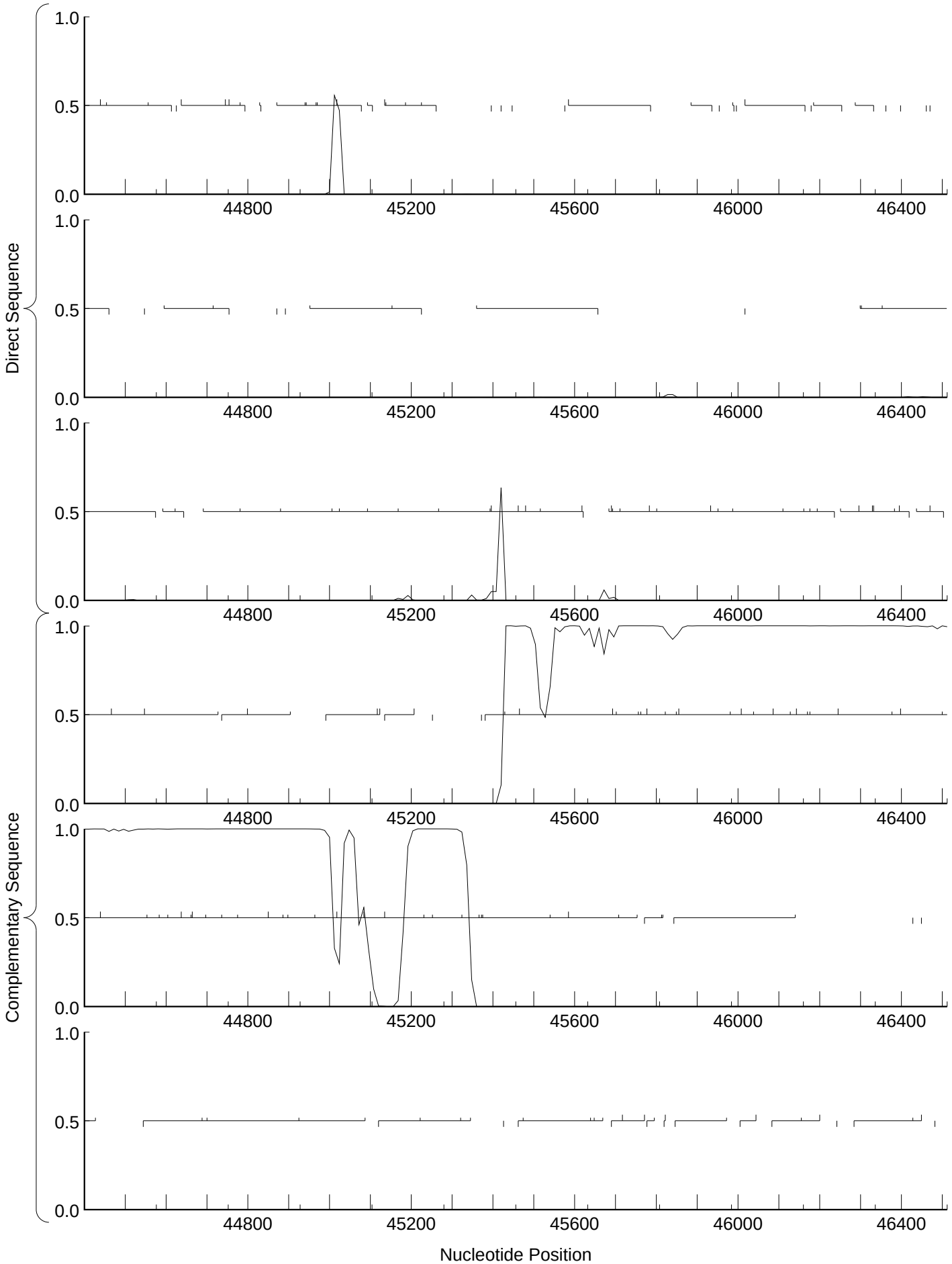

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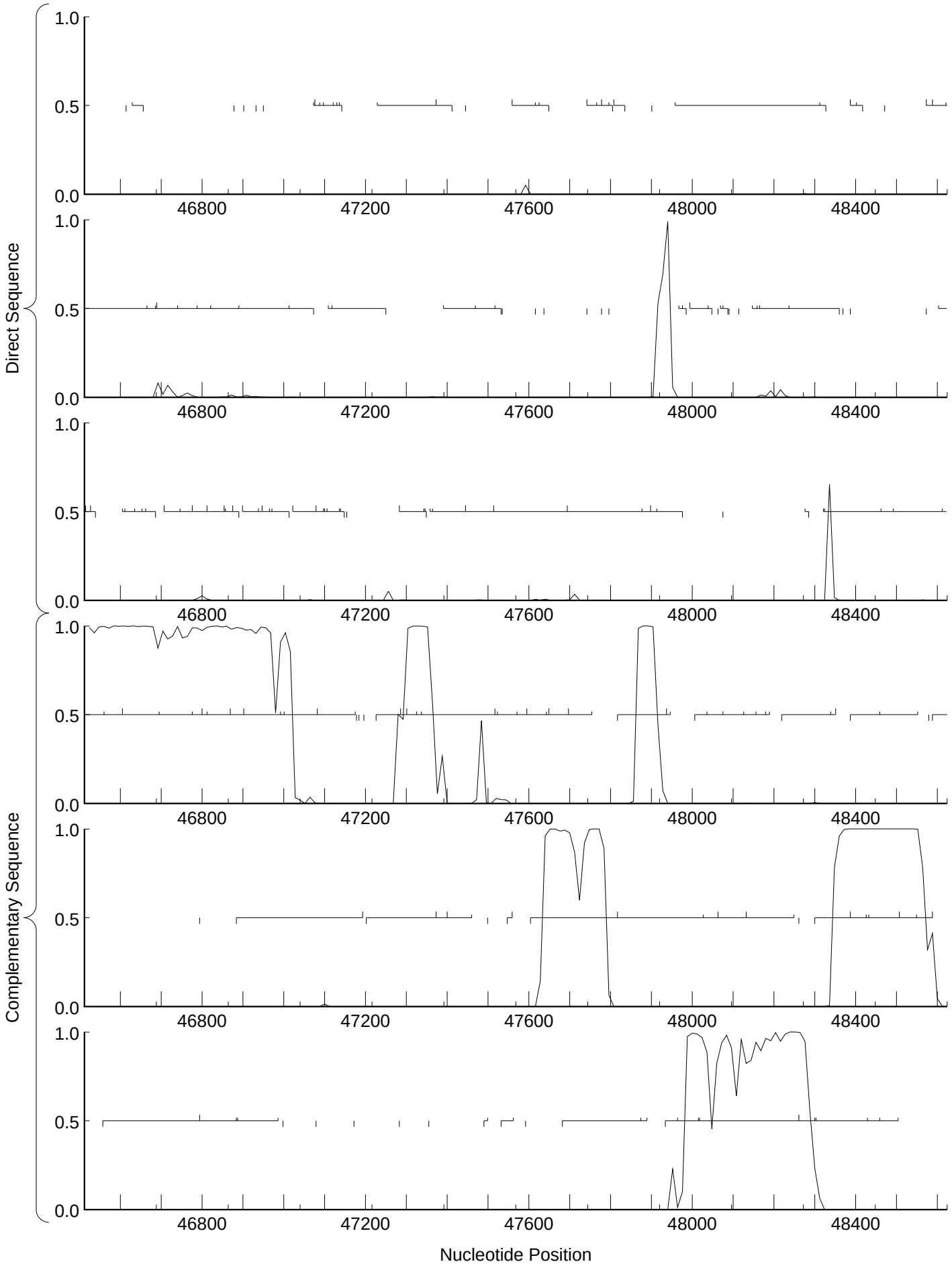

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Nucleotide Position

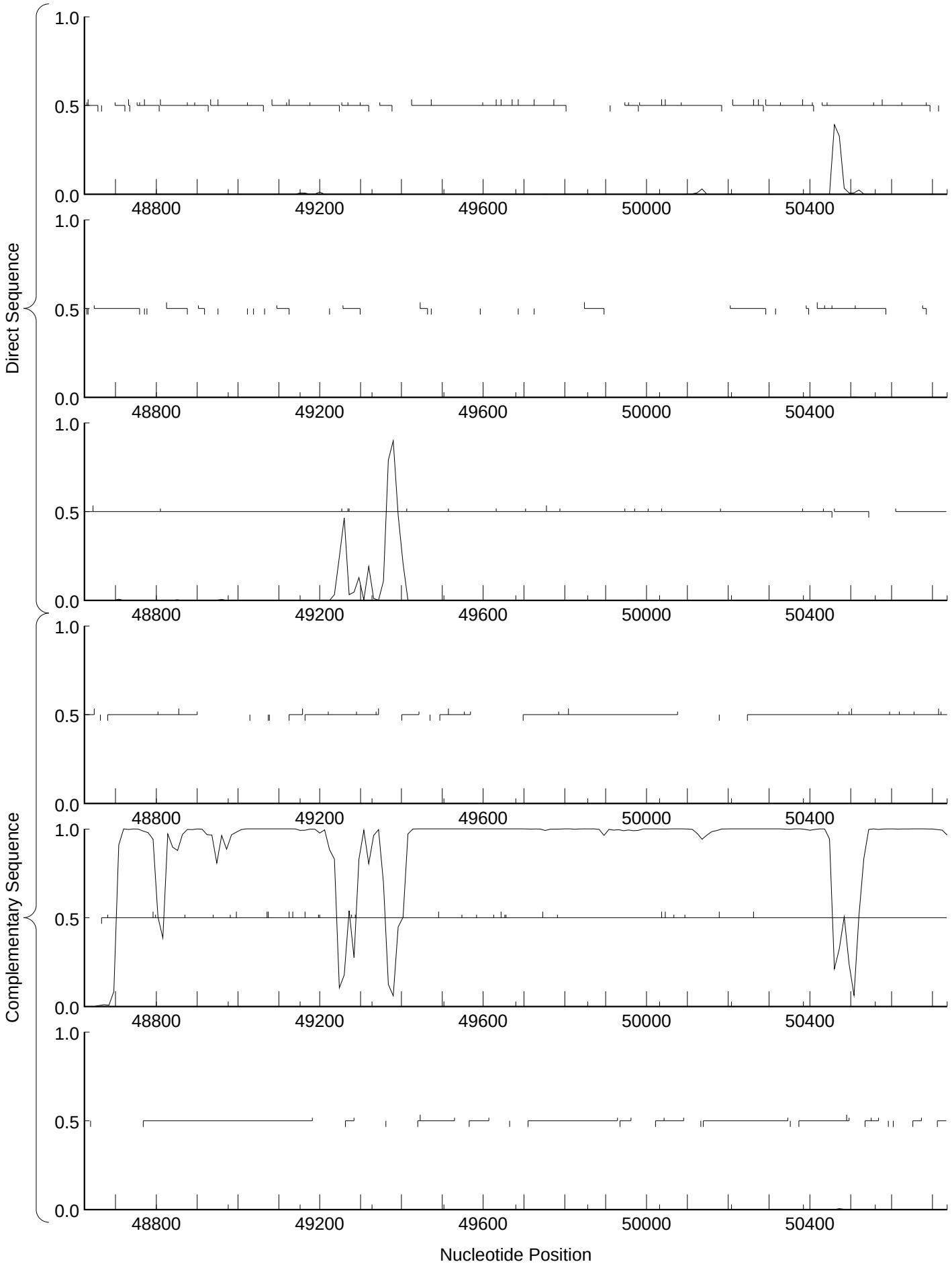

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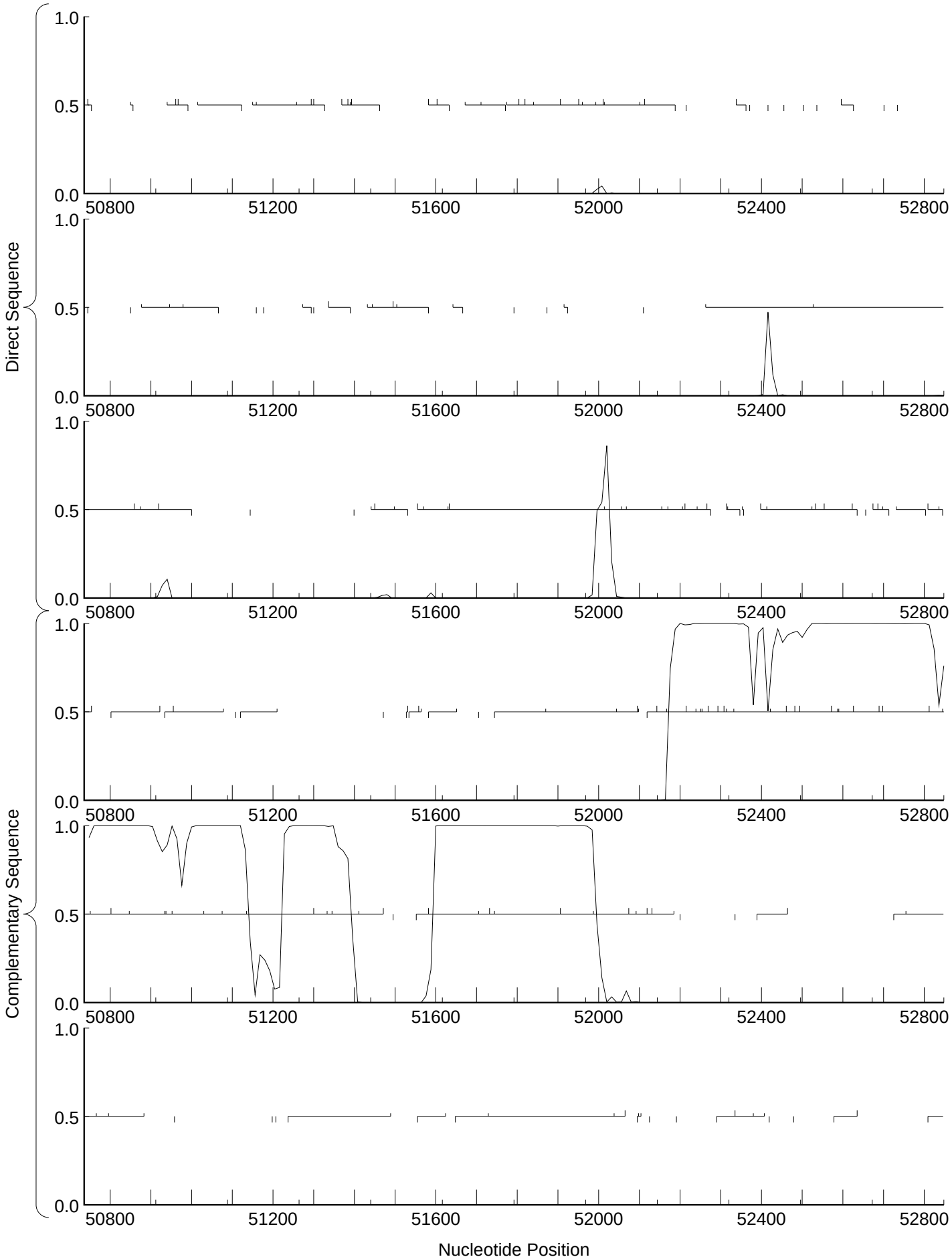

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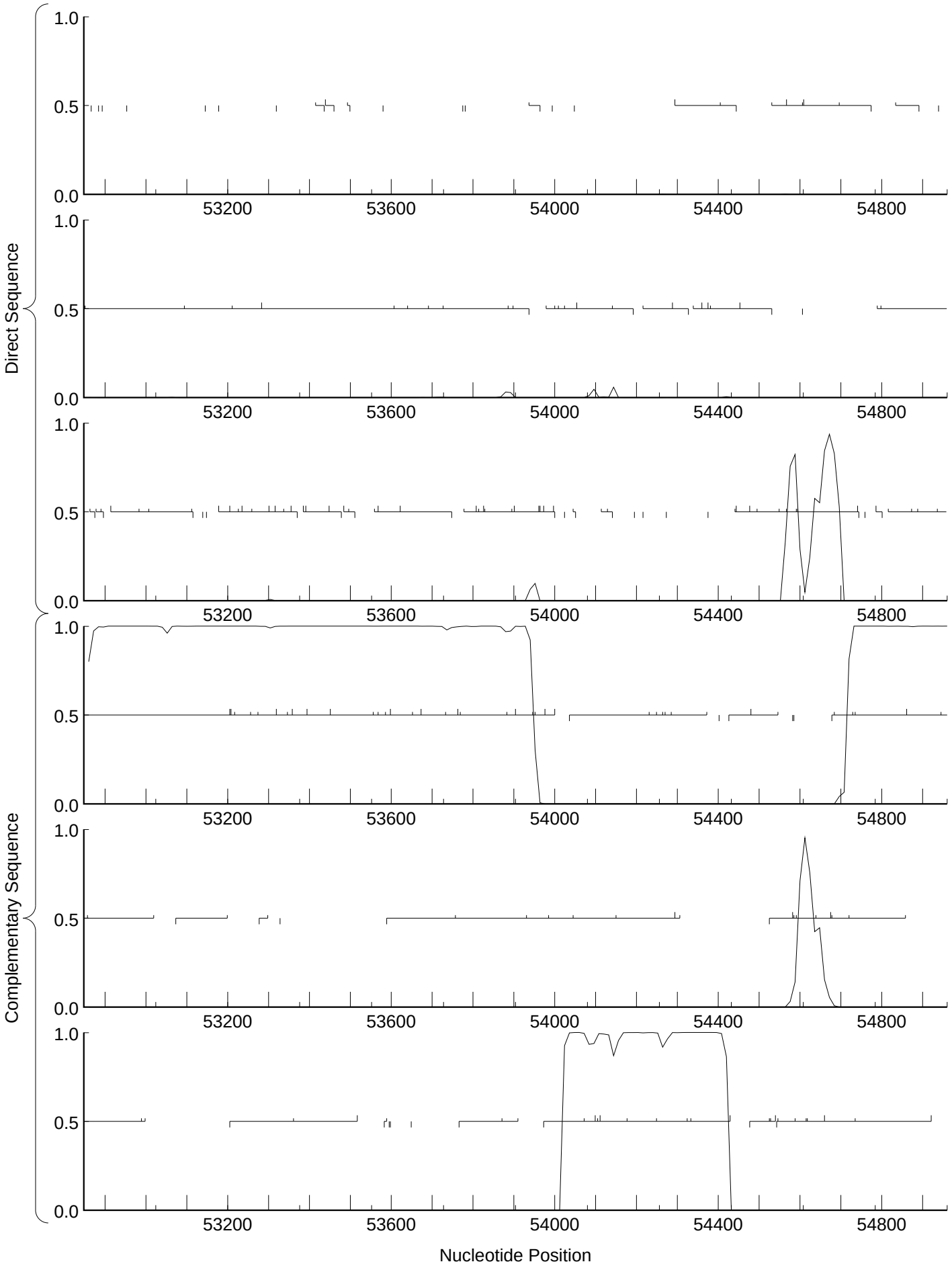

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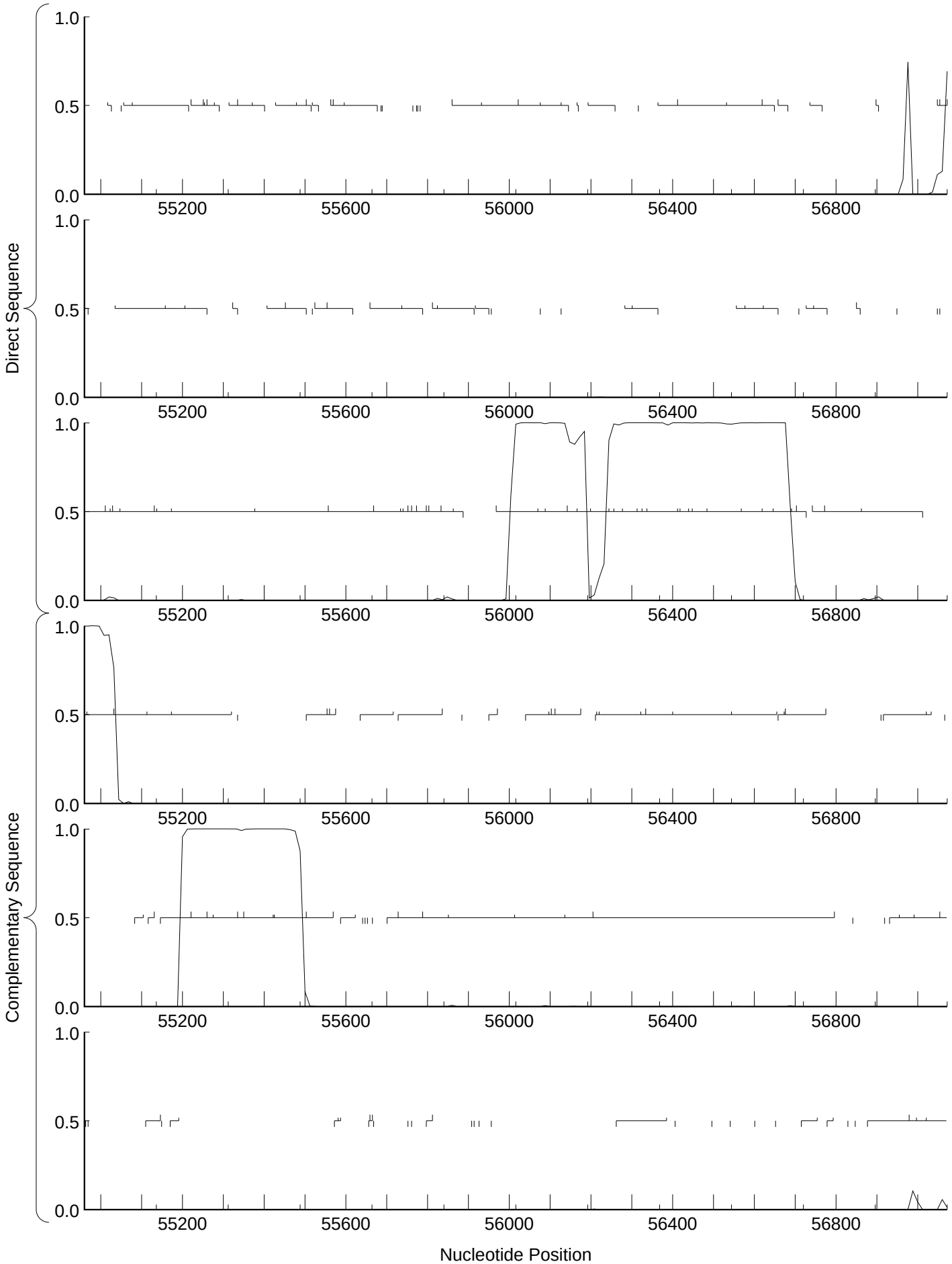

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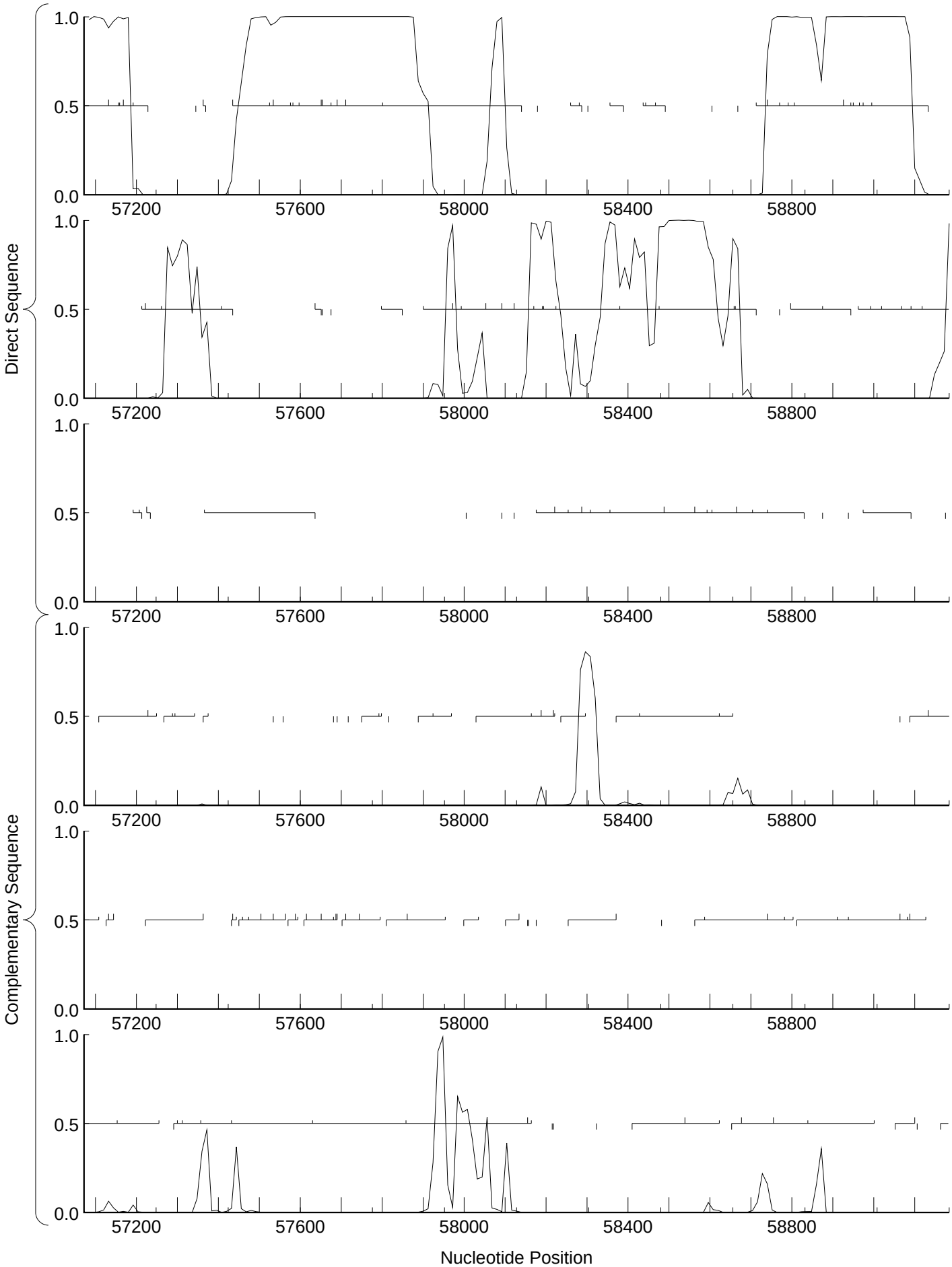

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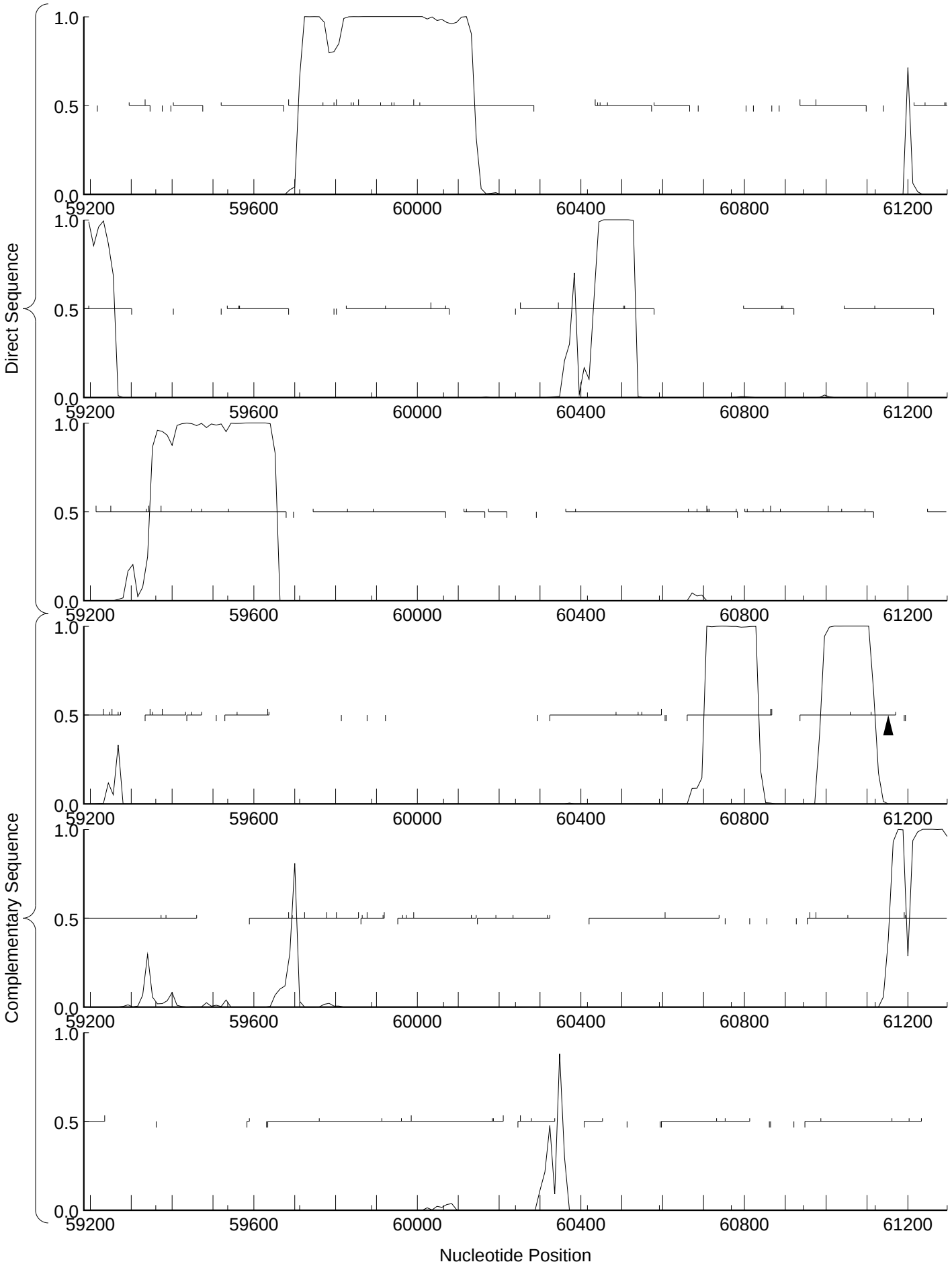

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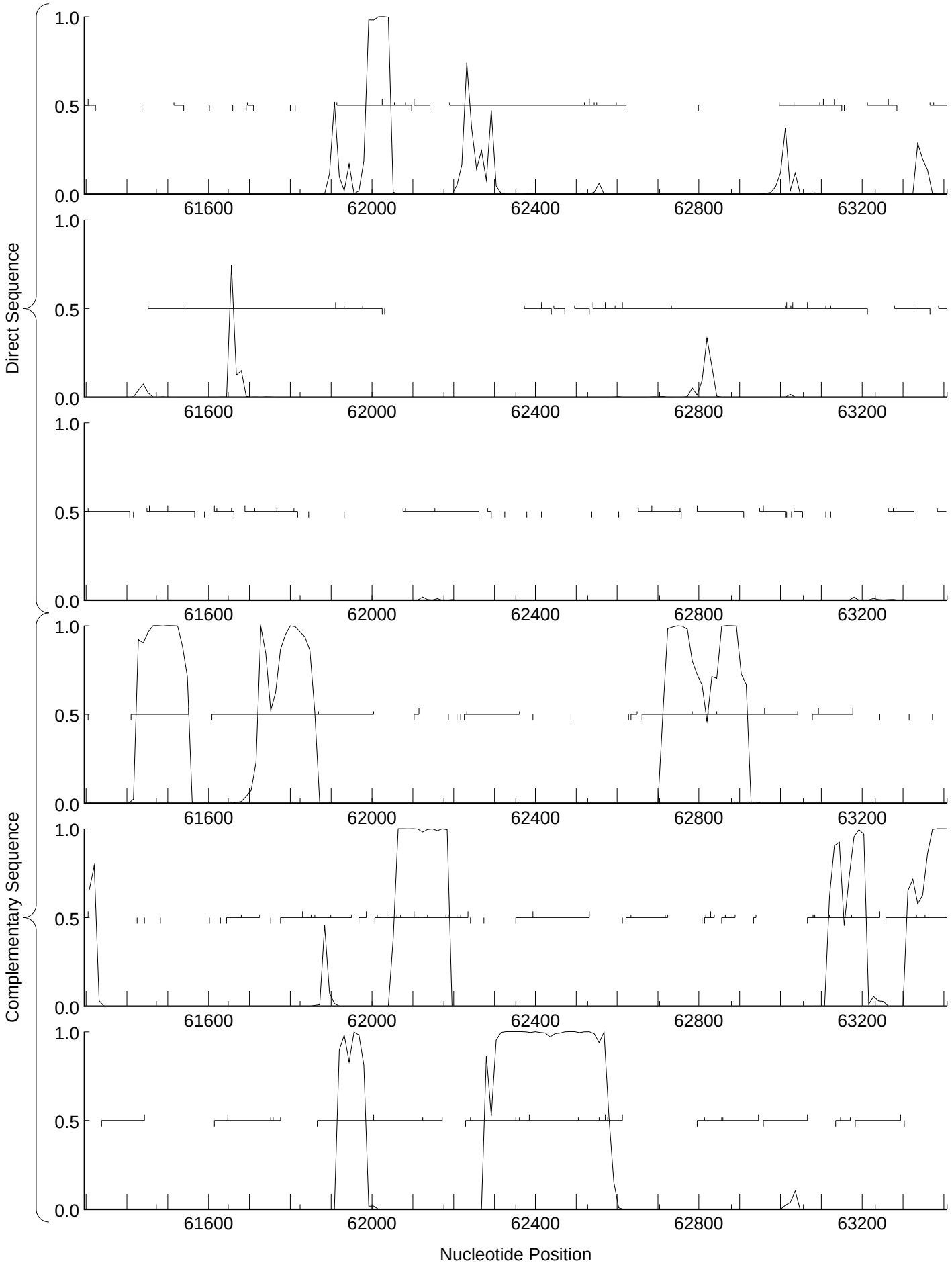

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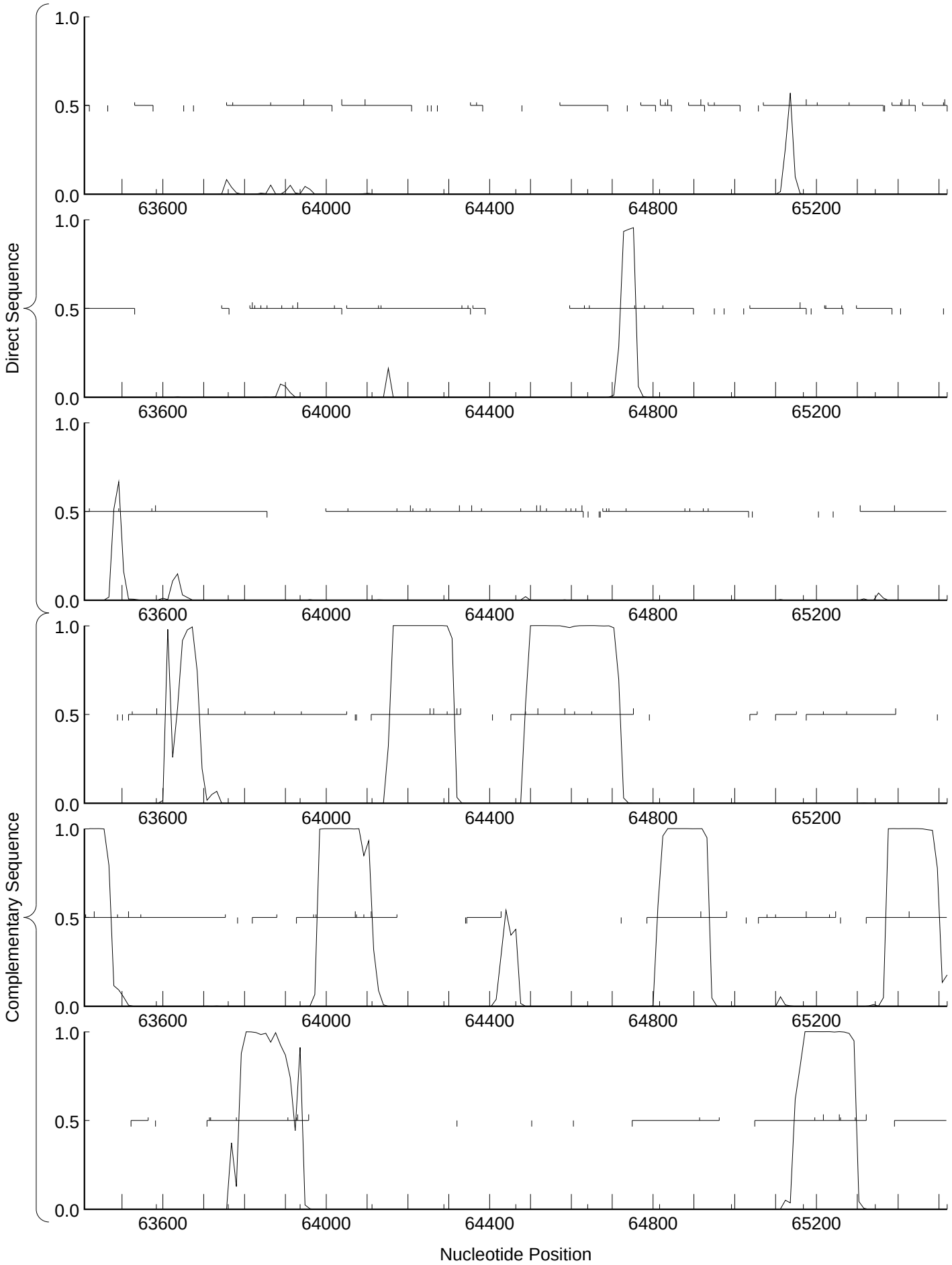

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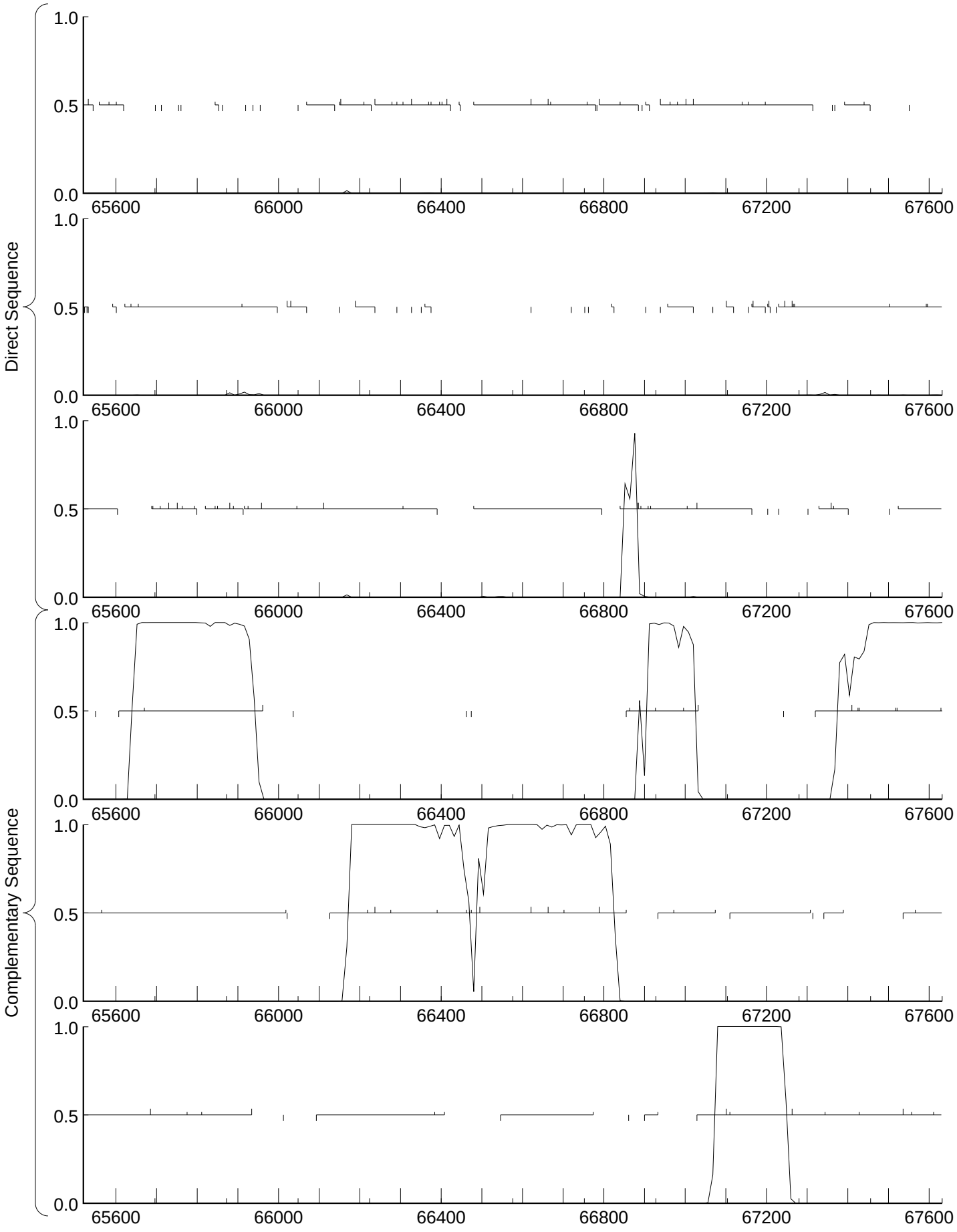

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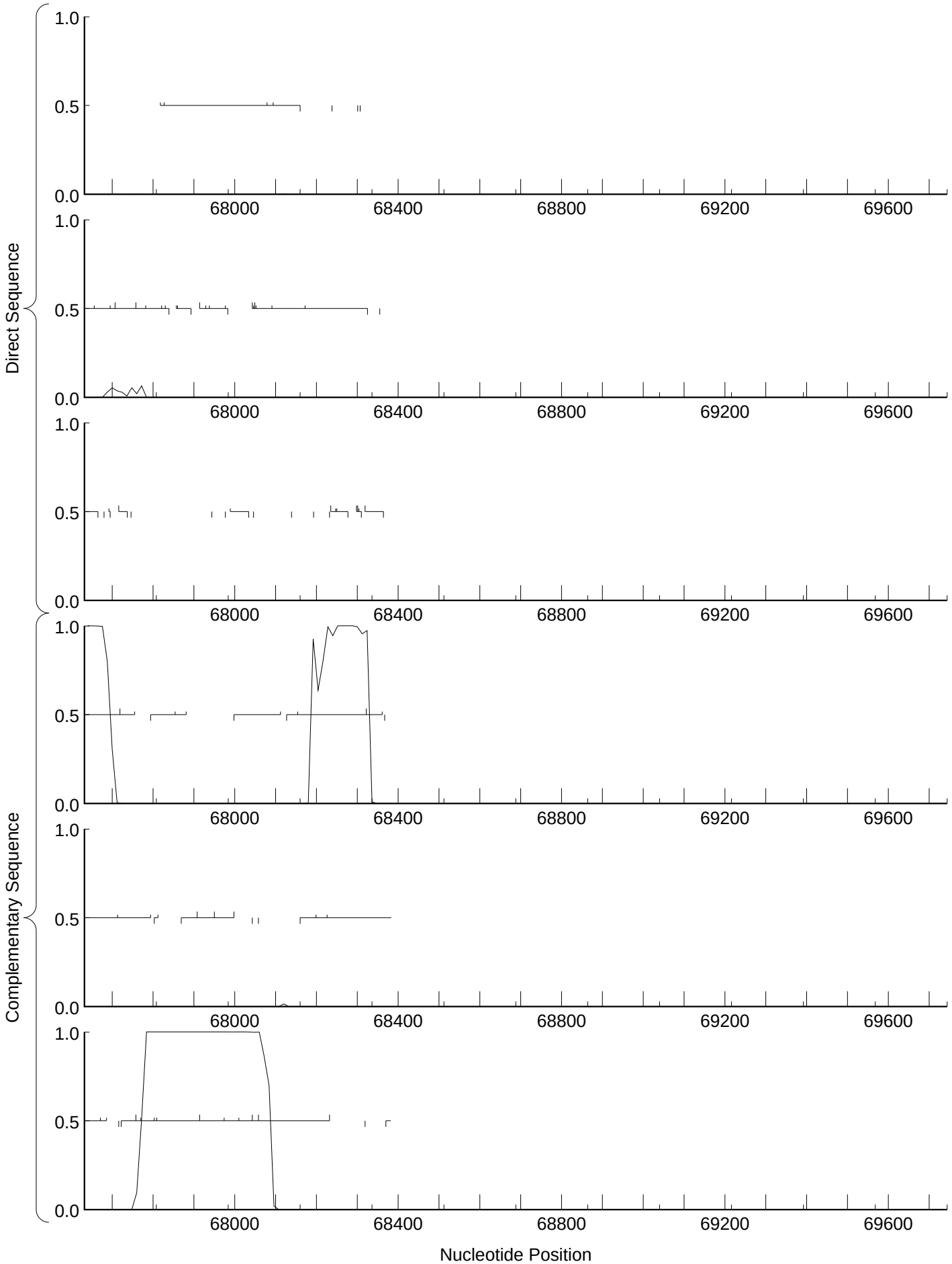

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Complementary Sequence
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Nucleotide Position


Direct Sequence
Complementary Sequence
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59600
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60800
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Nucleotide Position


Direct Sequence
Complementary Sequence
1.0
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63200
Nucleotide Position


Direct Sequence
Complementary Sequence
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64800
65200
Nucleotide Position


Direct Sequence
Complementary Sequence
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67600
Nucleotide Position


Direct Sequence
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