GeneMark

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

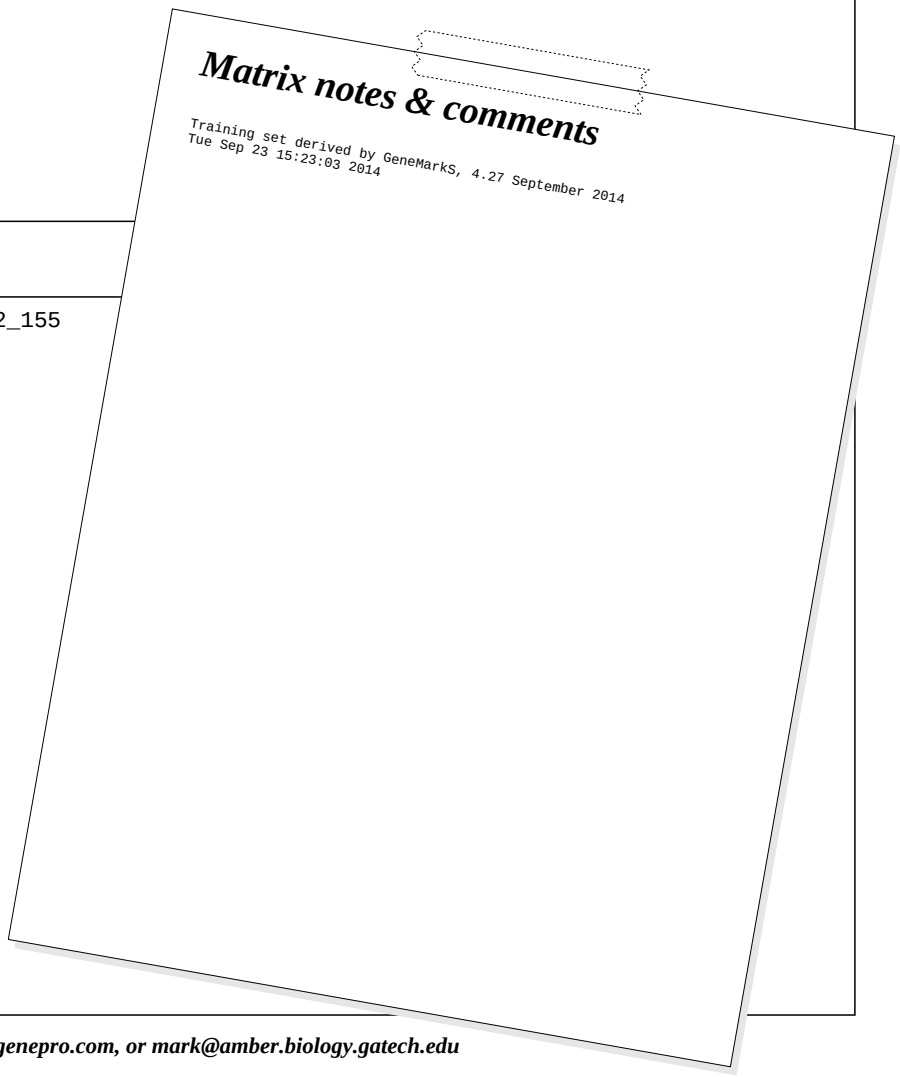
PROGRAM INFORMATION

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Sequence         :  Mycobacterium phage 244 complete genome
Analysis Date    :  3/29/24 at 8:50:54
Pages            :  37
Sequence Length  :  74483 bp
GC Content       :   62.94%

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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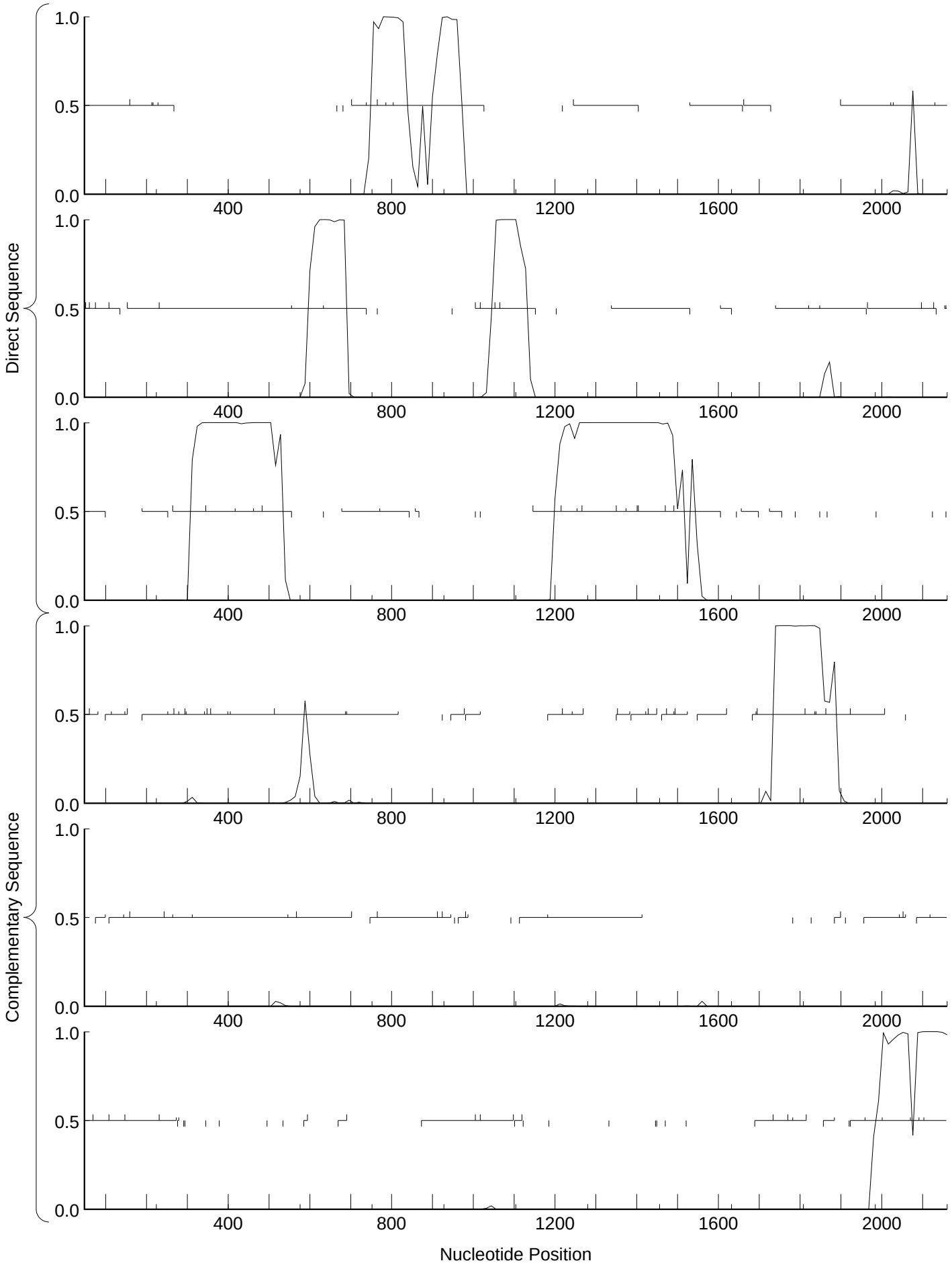
MATRIX INFORMATION

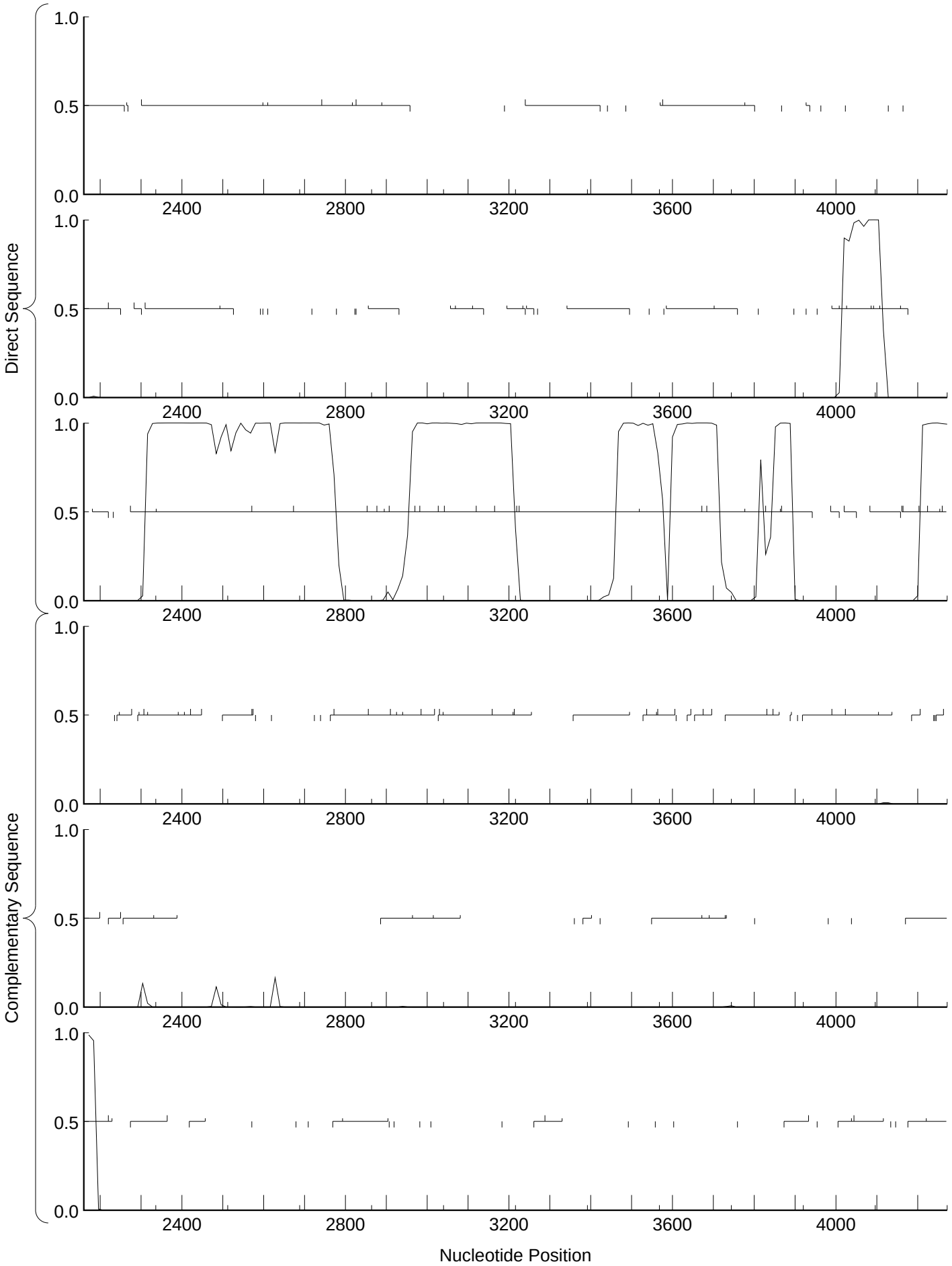
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Author     :  -

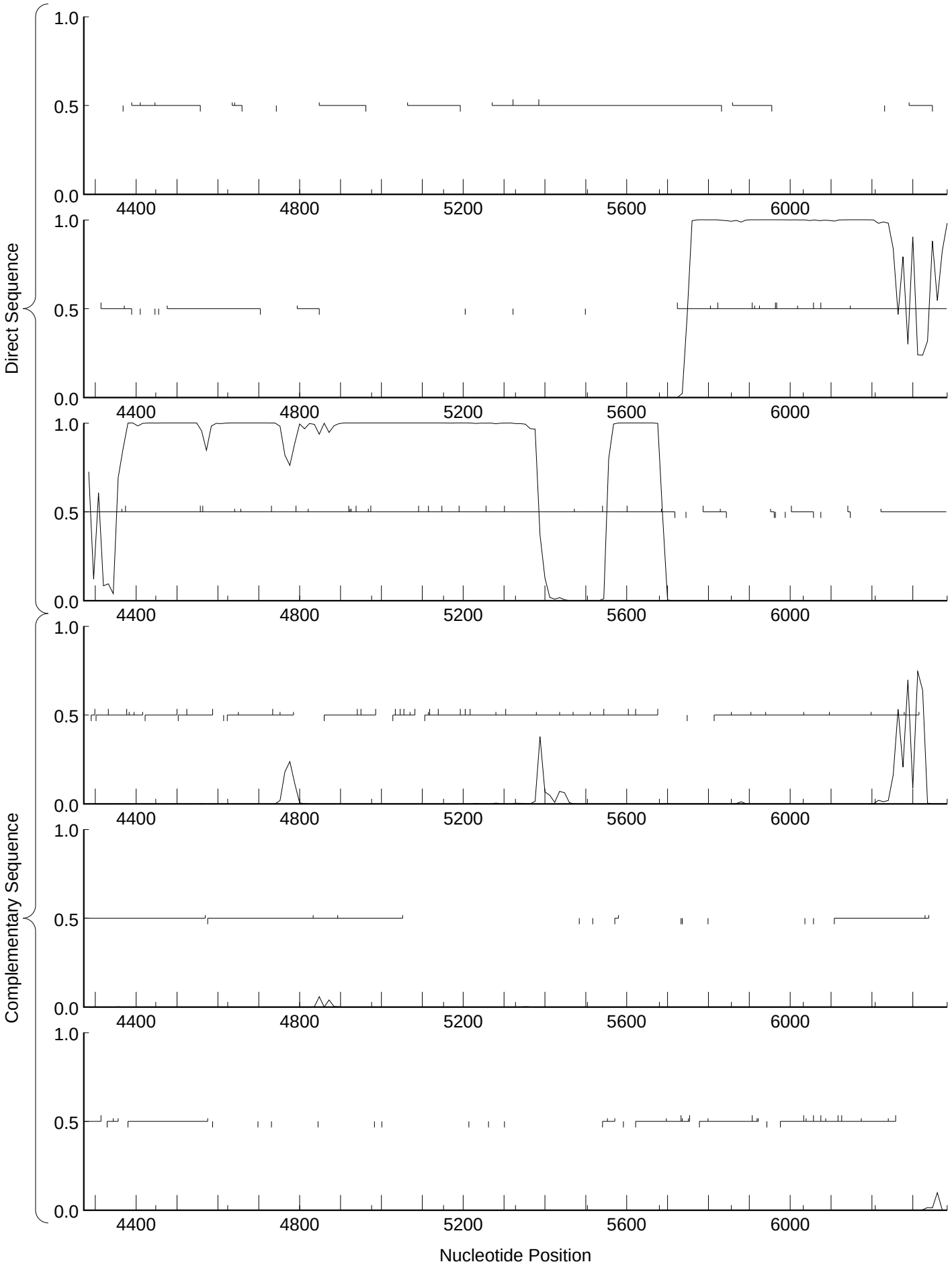
Order      :  4
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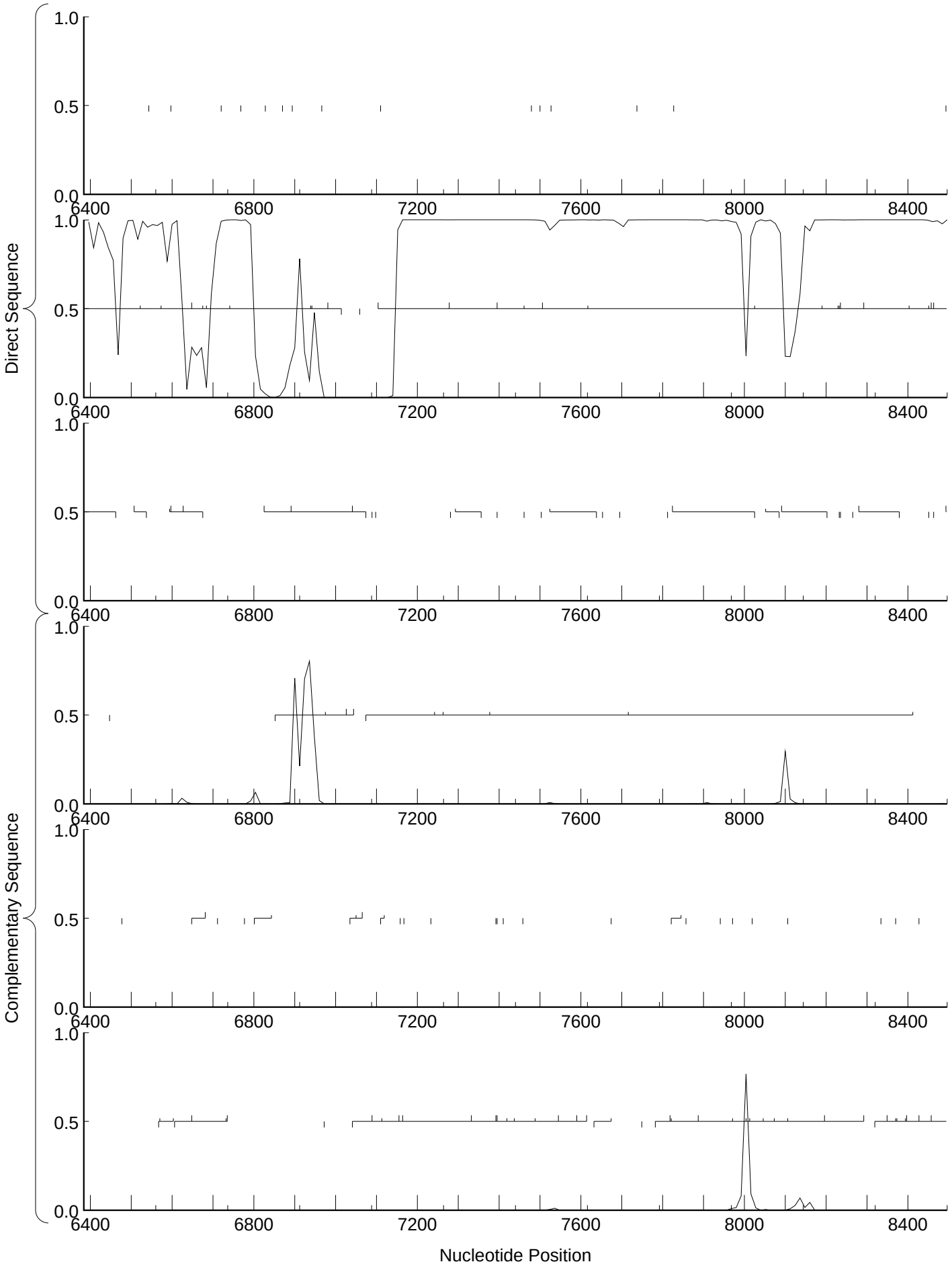

```
  Send questions / comments to:
      Dr. M. Borodovsky
      Georgia Institute of Technology
      School of Biology
      Atlanta, GA  30332-0230
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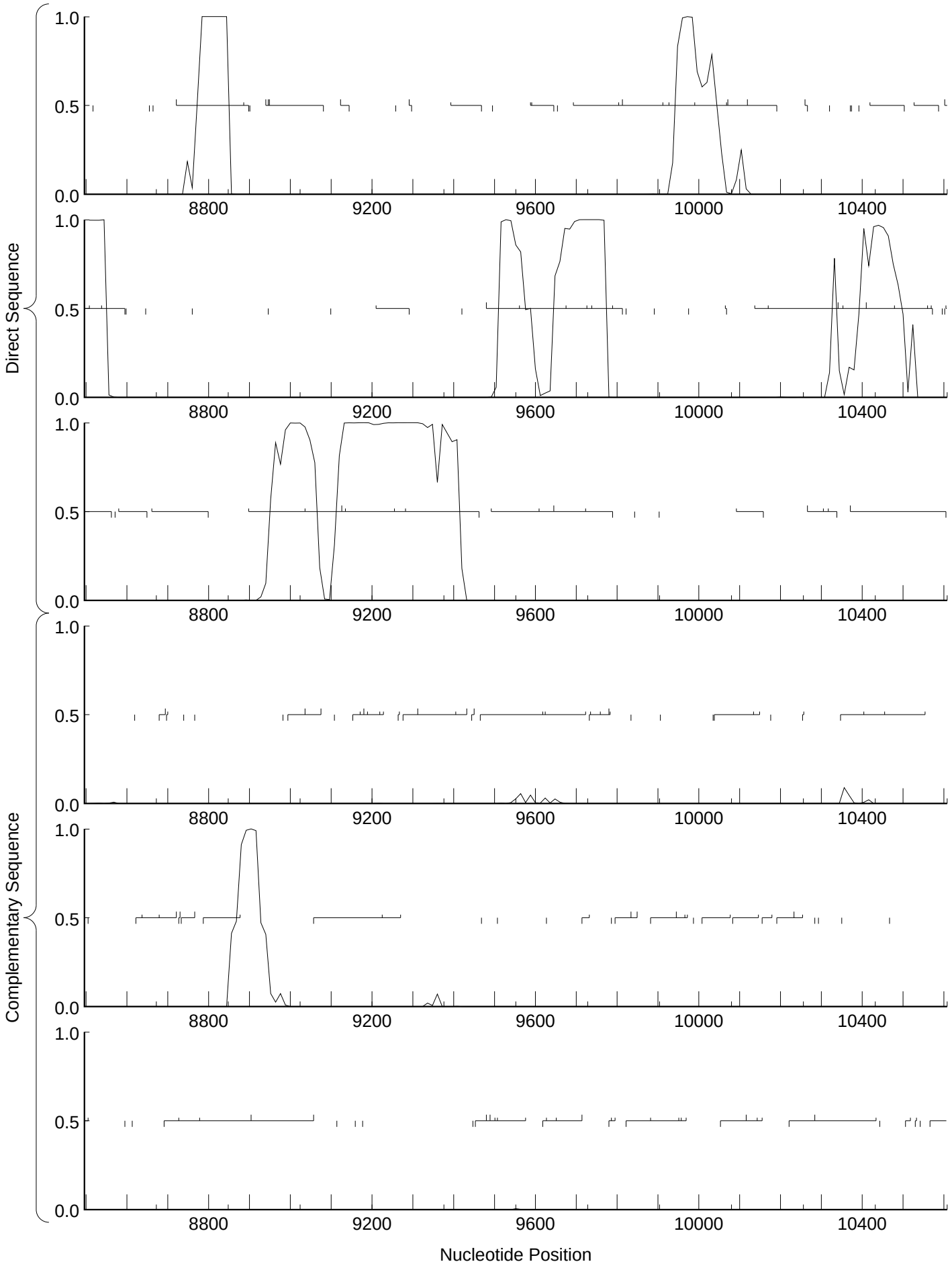

For help or information, send electronic mail to: custserv@genepro.com, or mark@amber.biology.gatech.edu

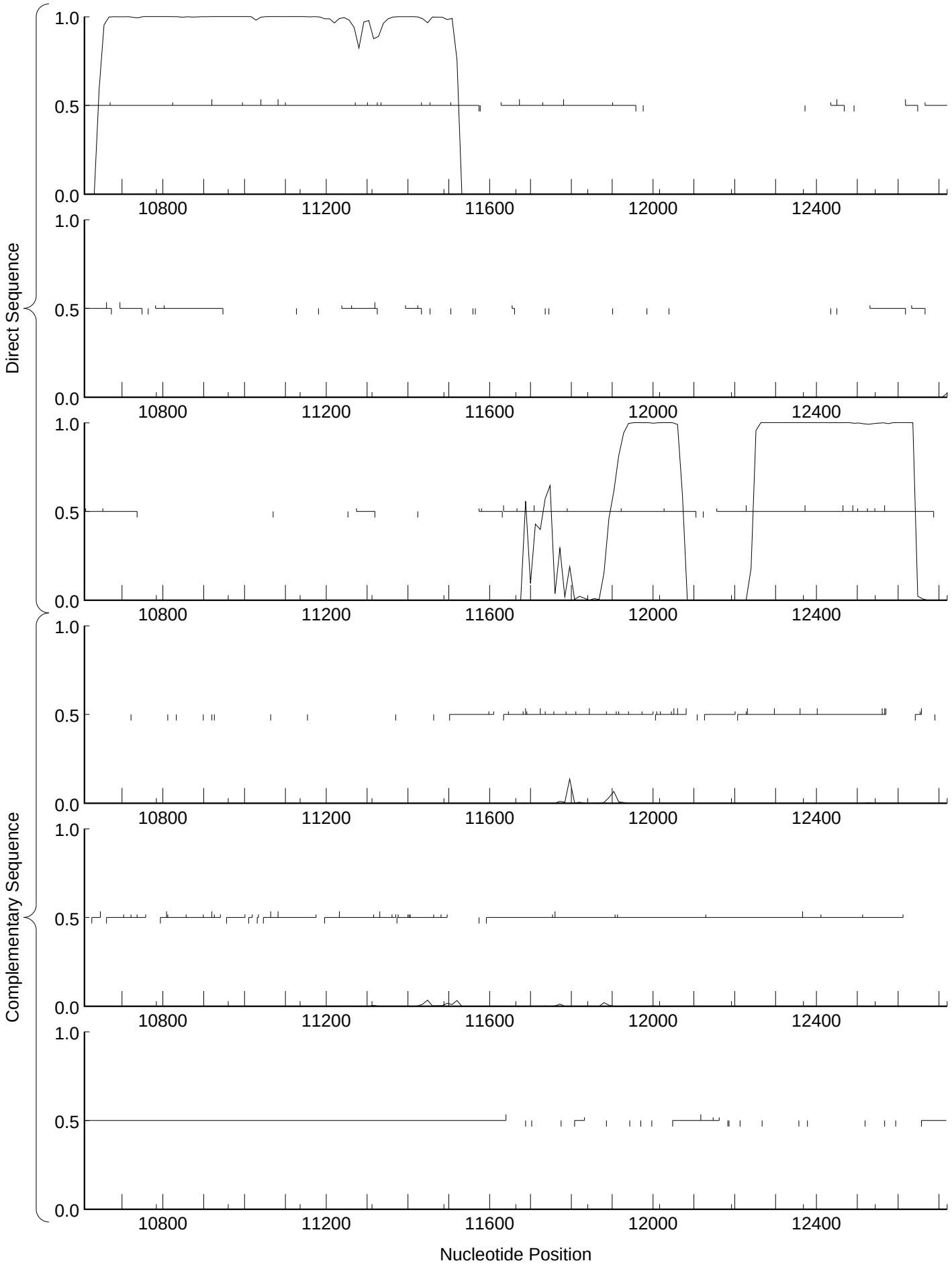

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Complementary Sequence
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0.0
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2000
Nucleotide Position

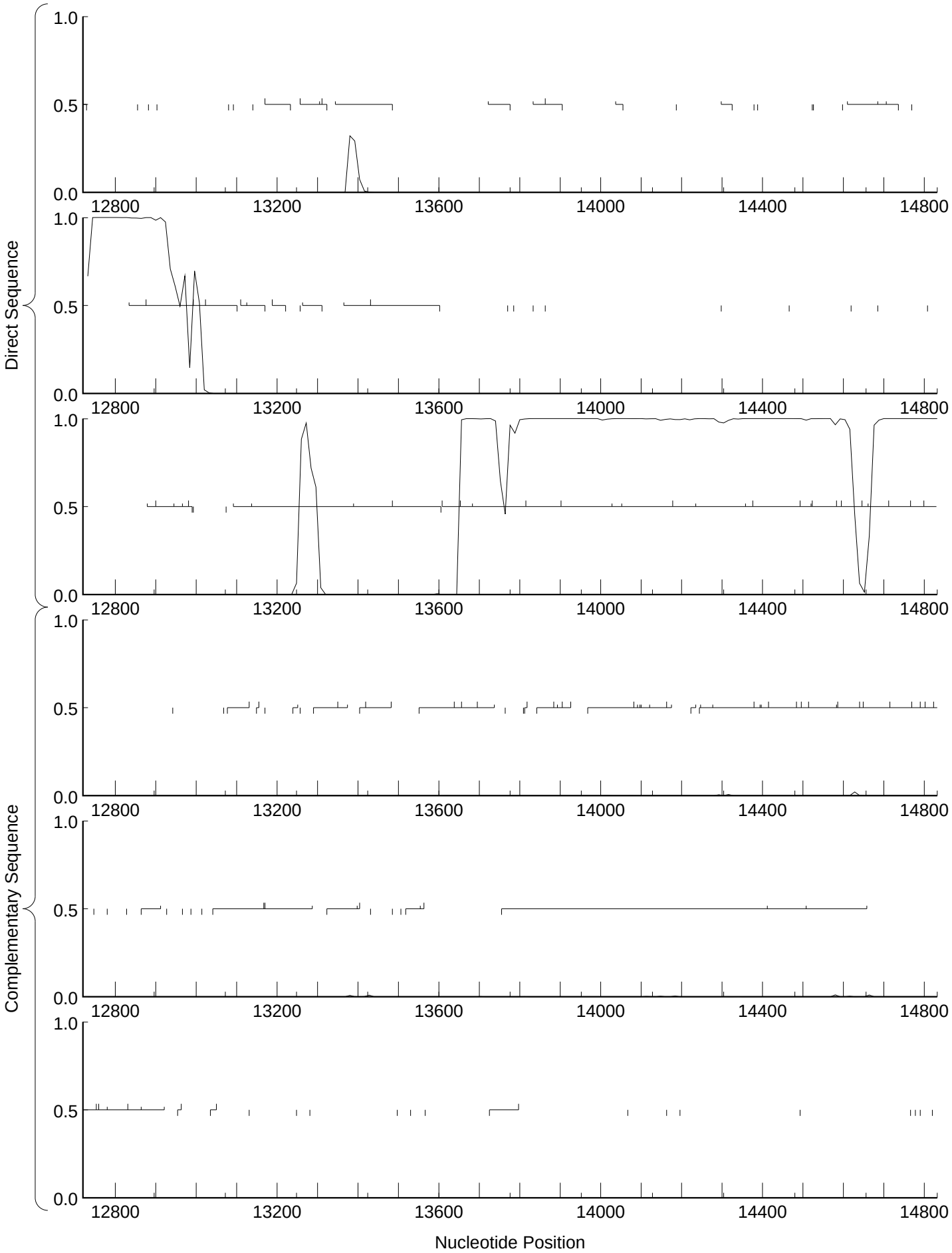

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Nucleotide Position

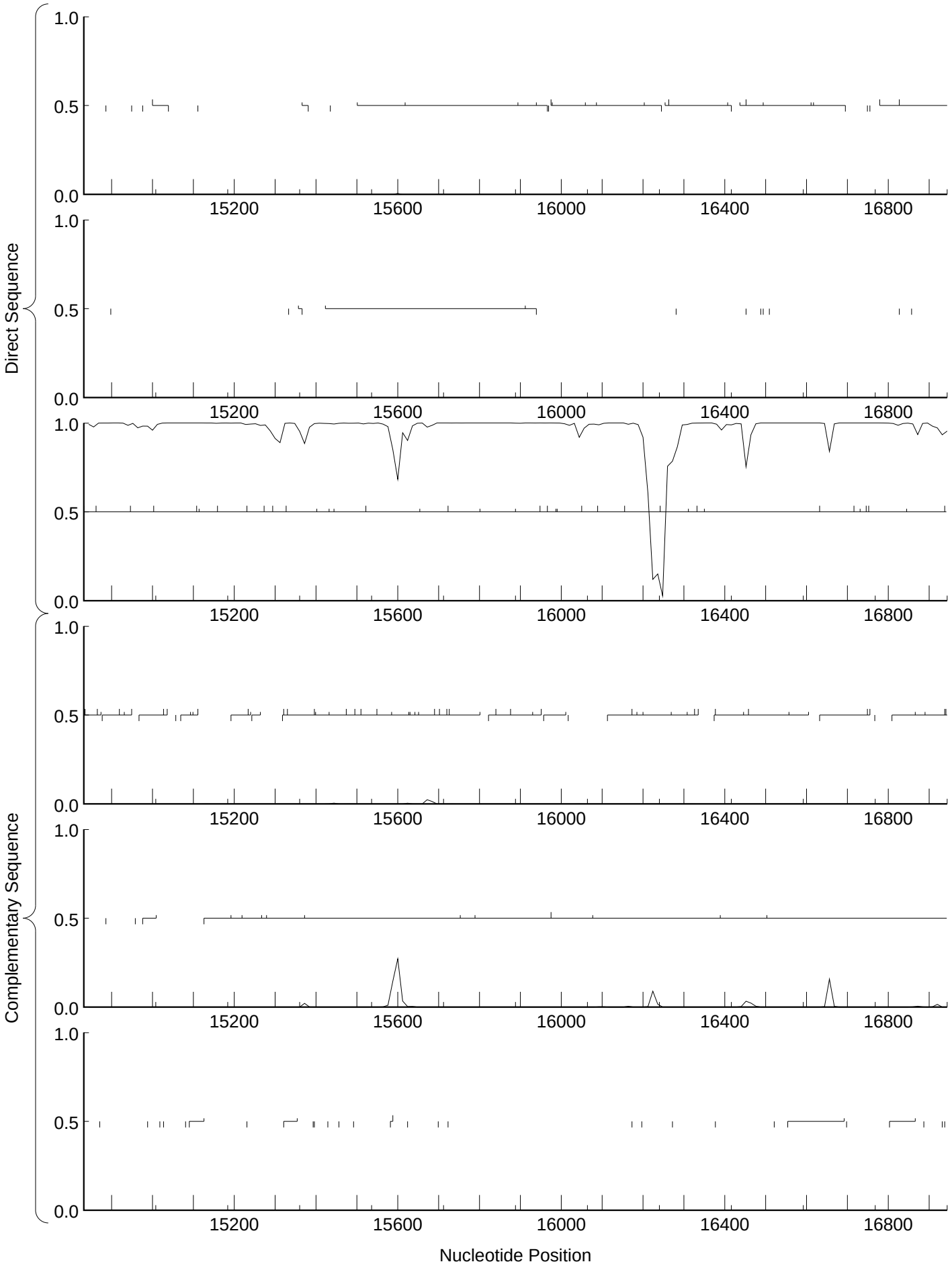

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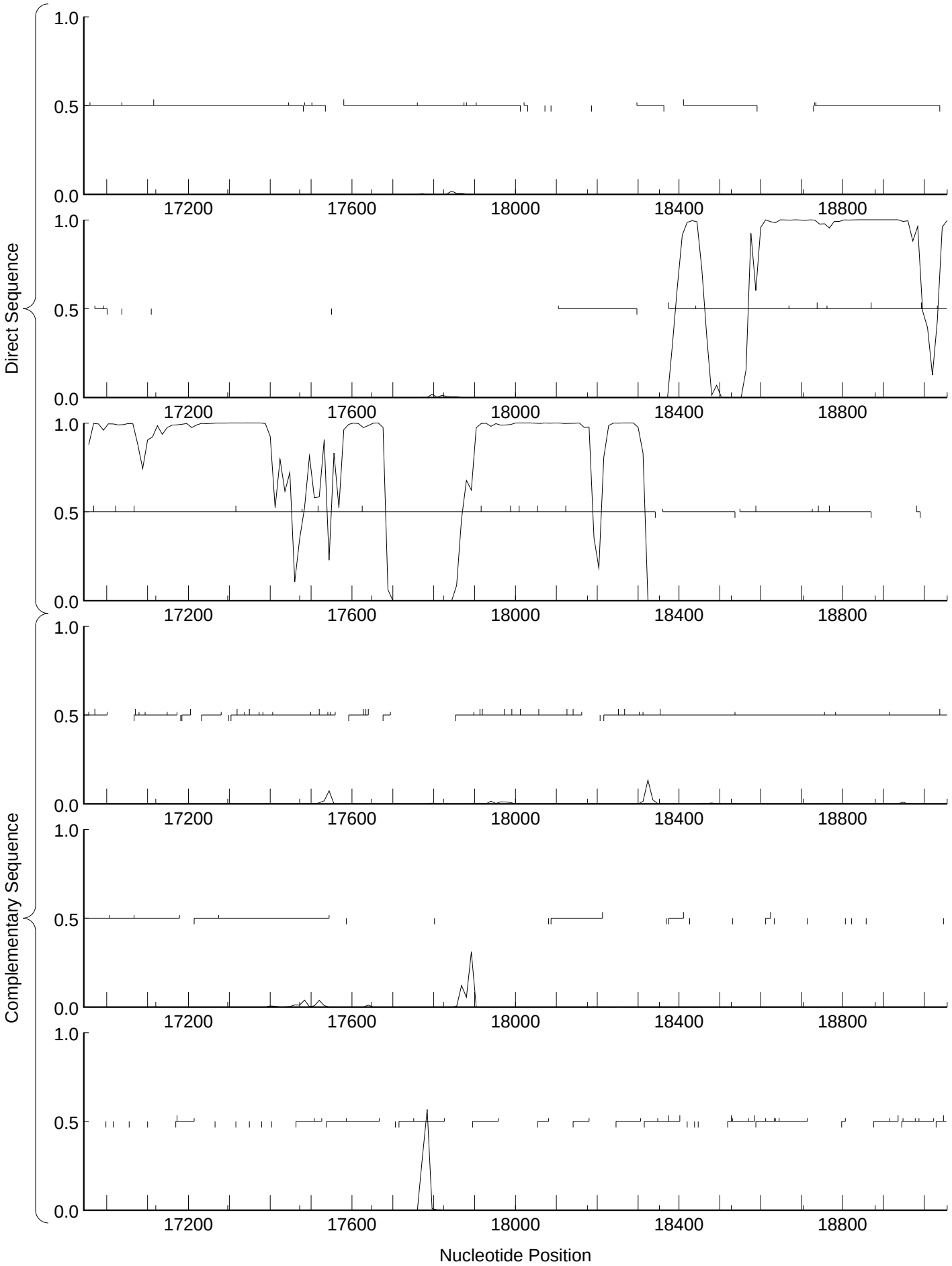

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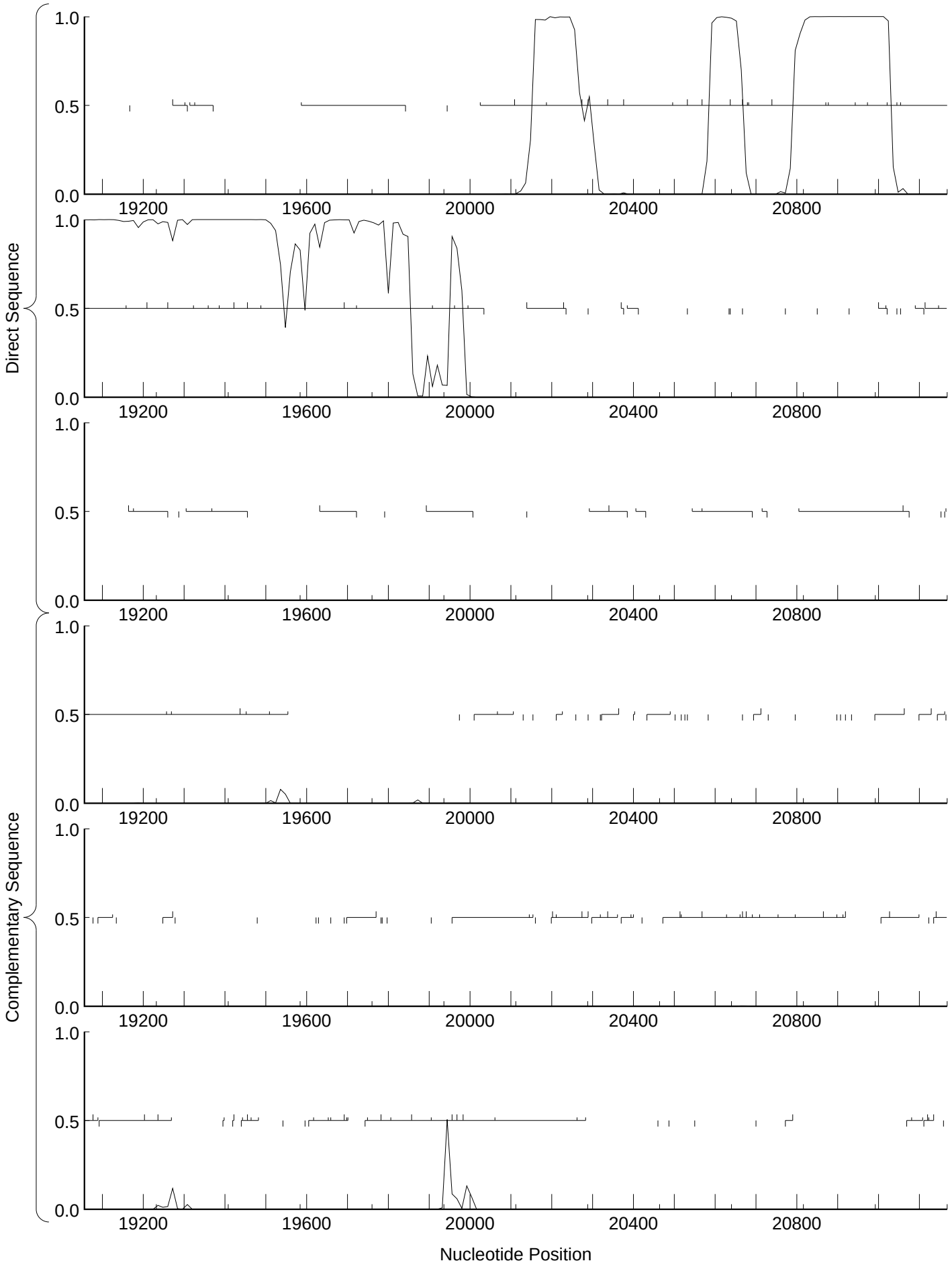

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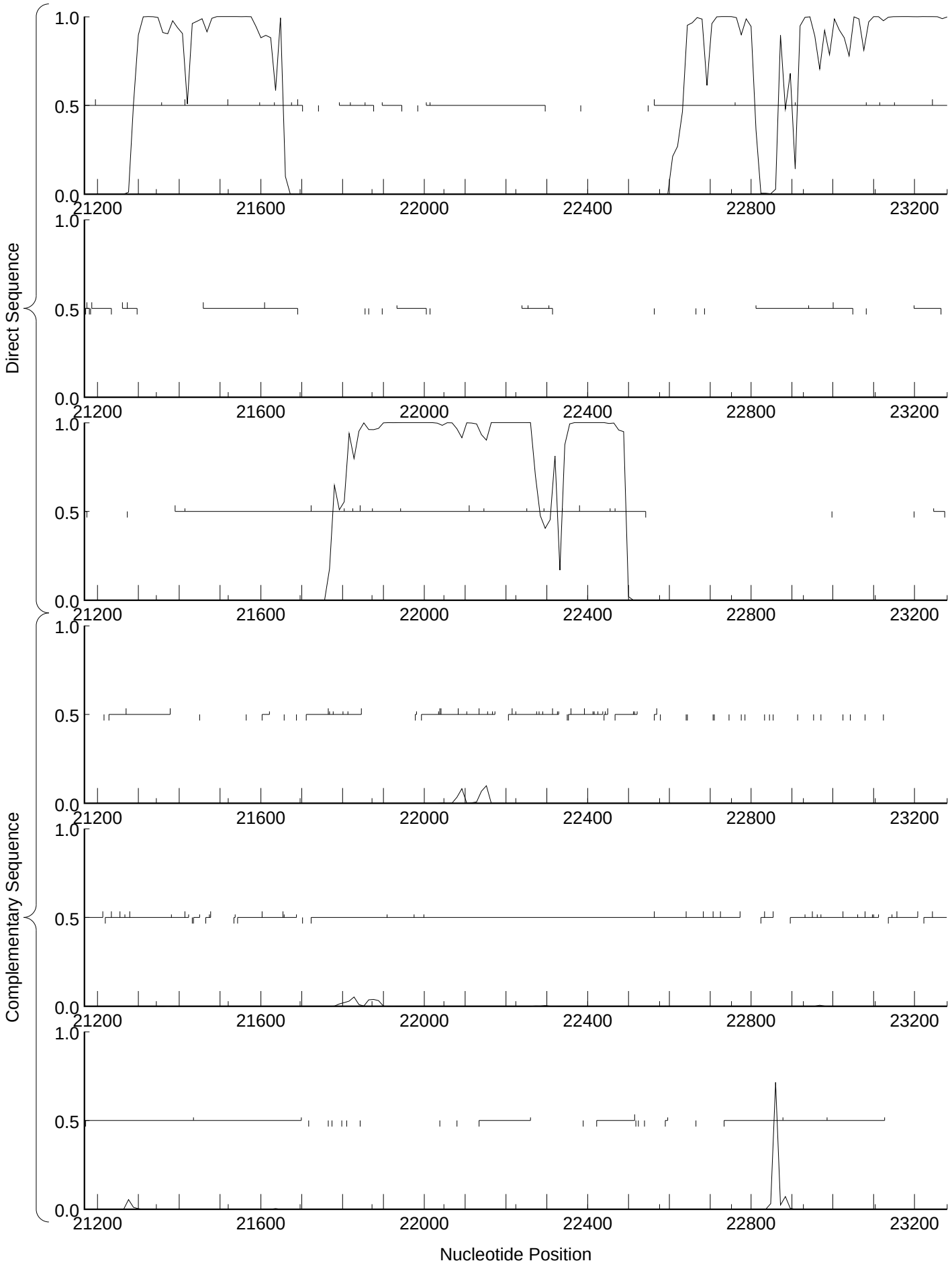

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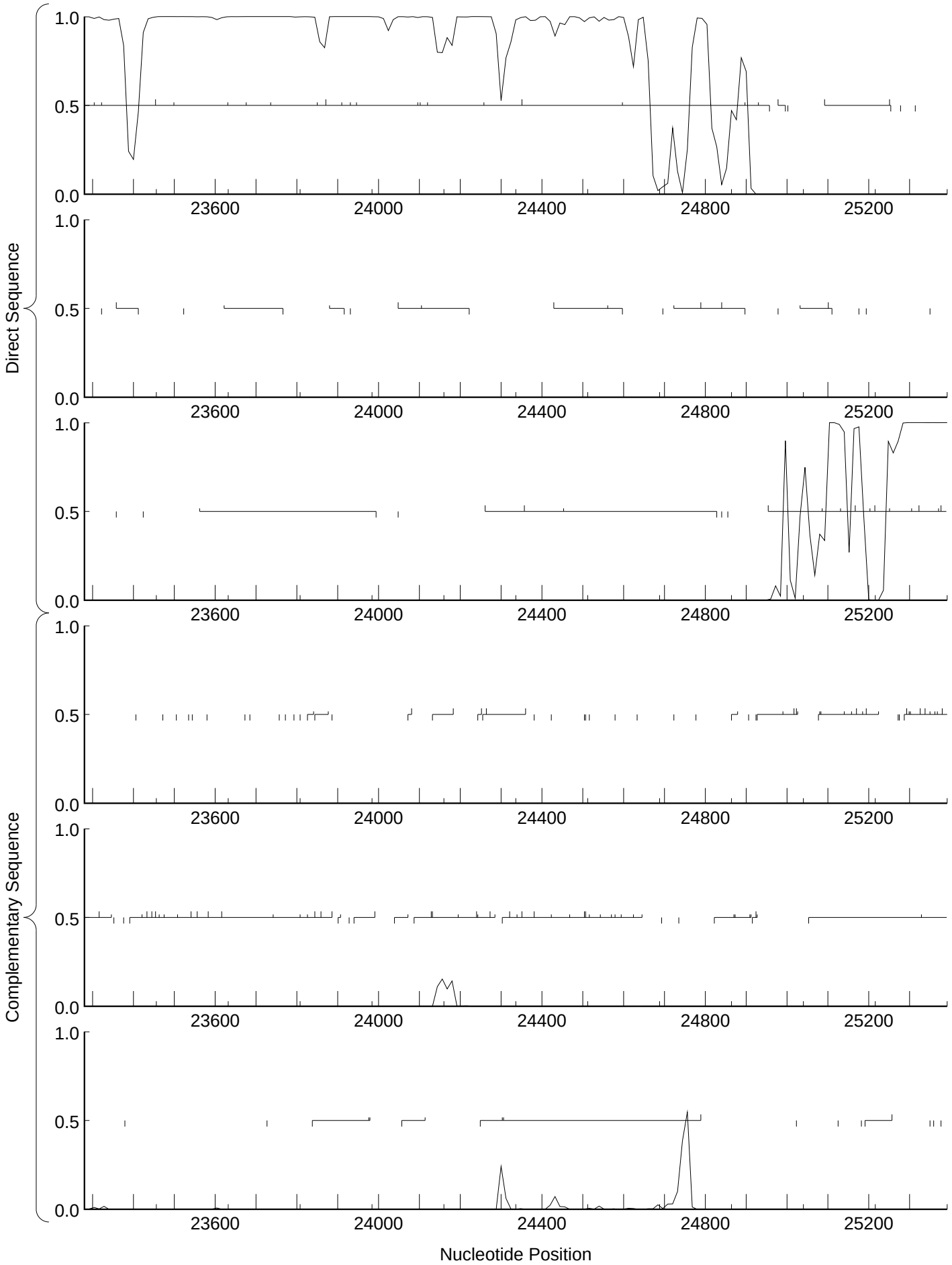

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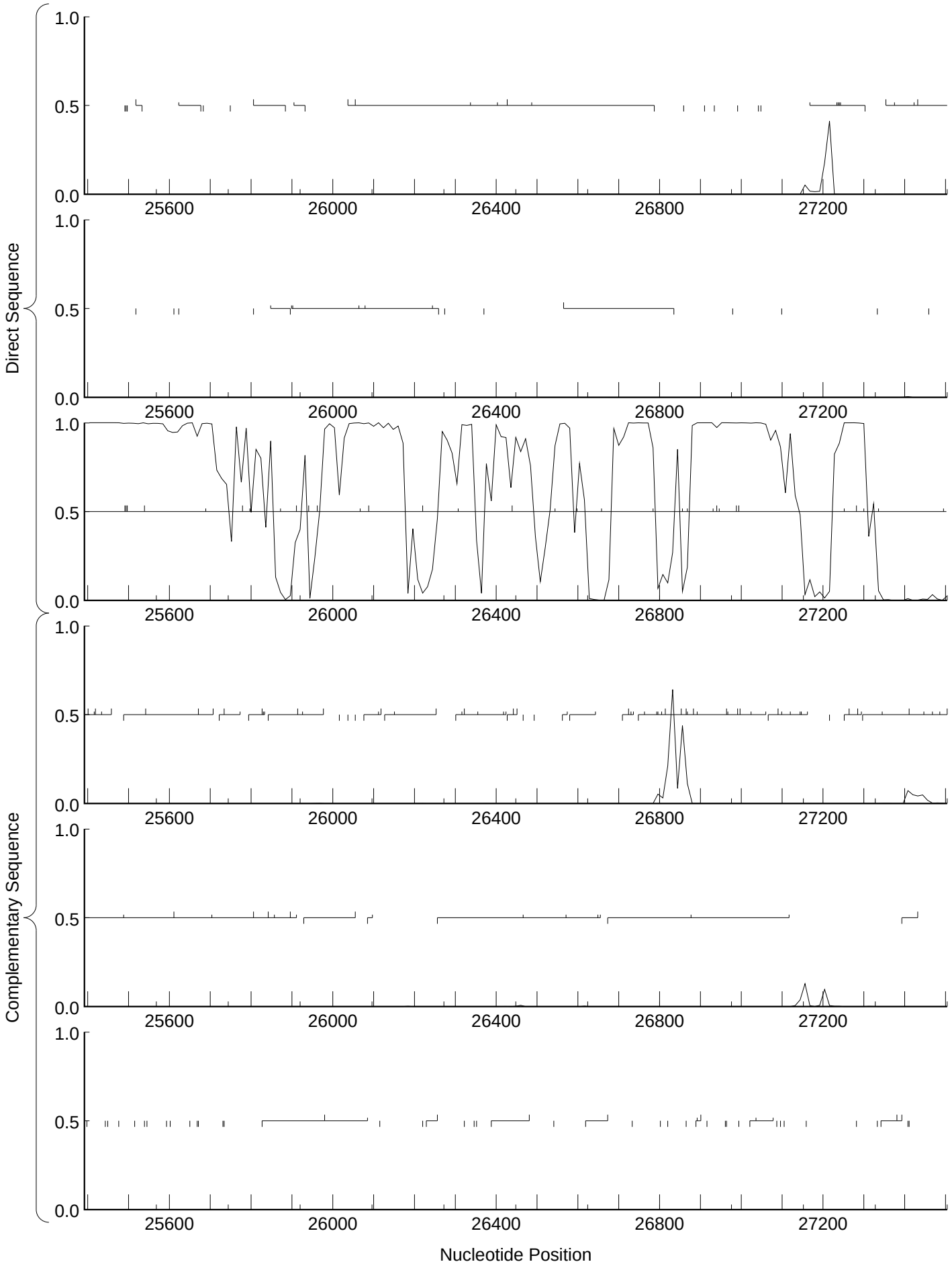

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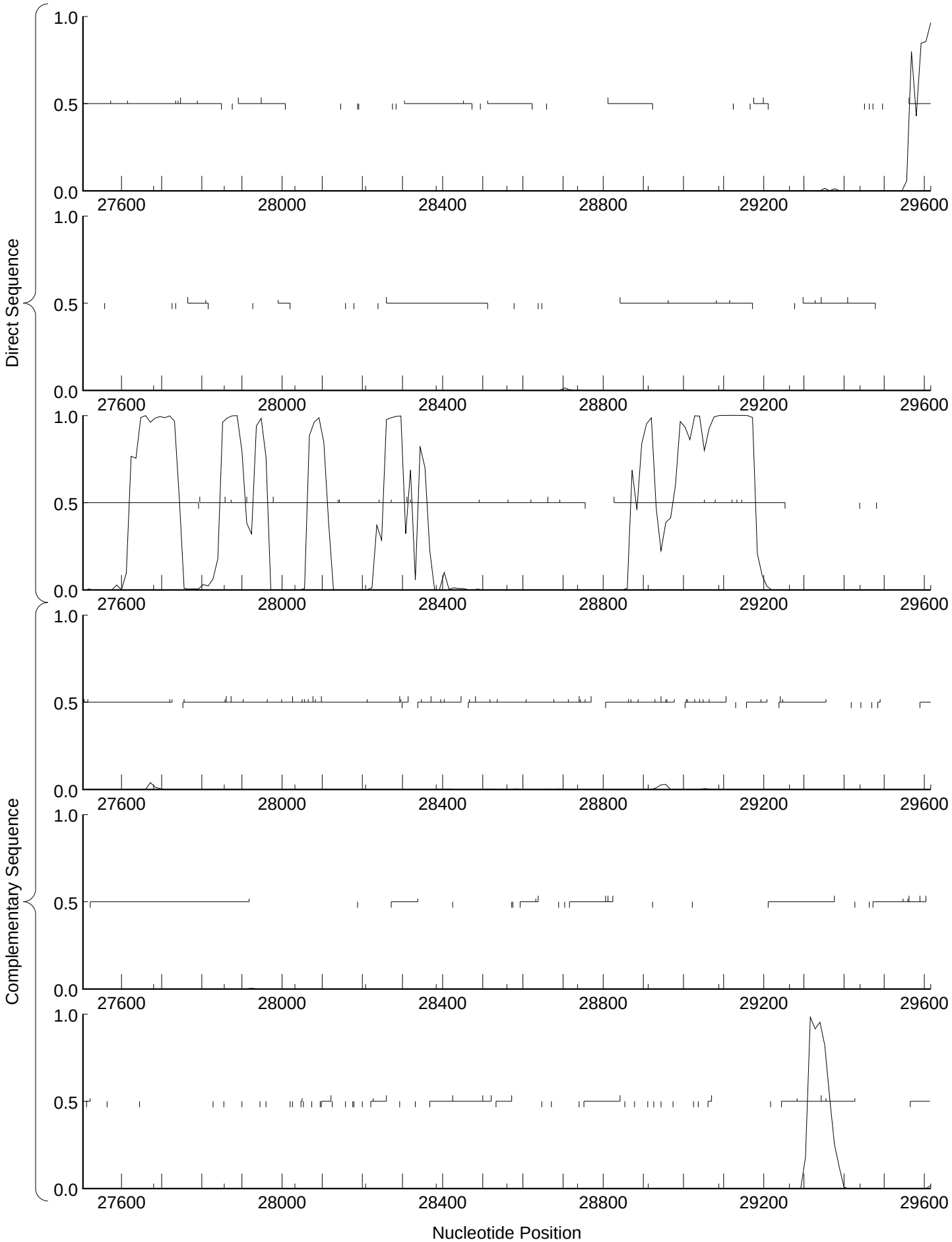

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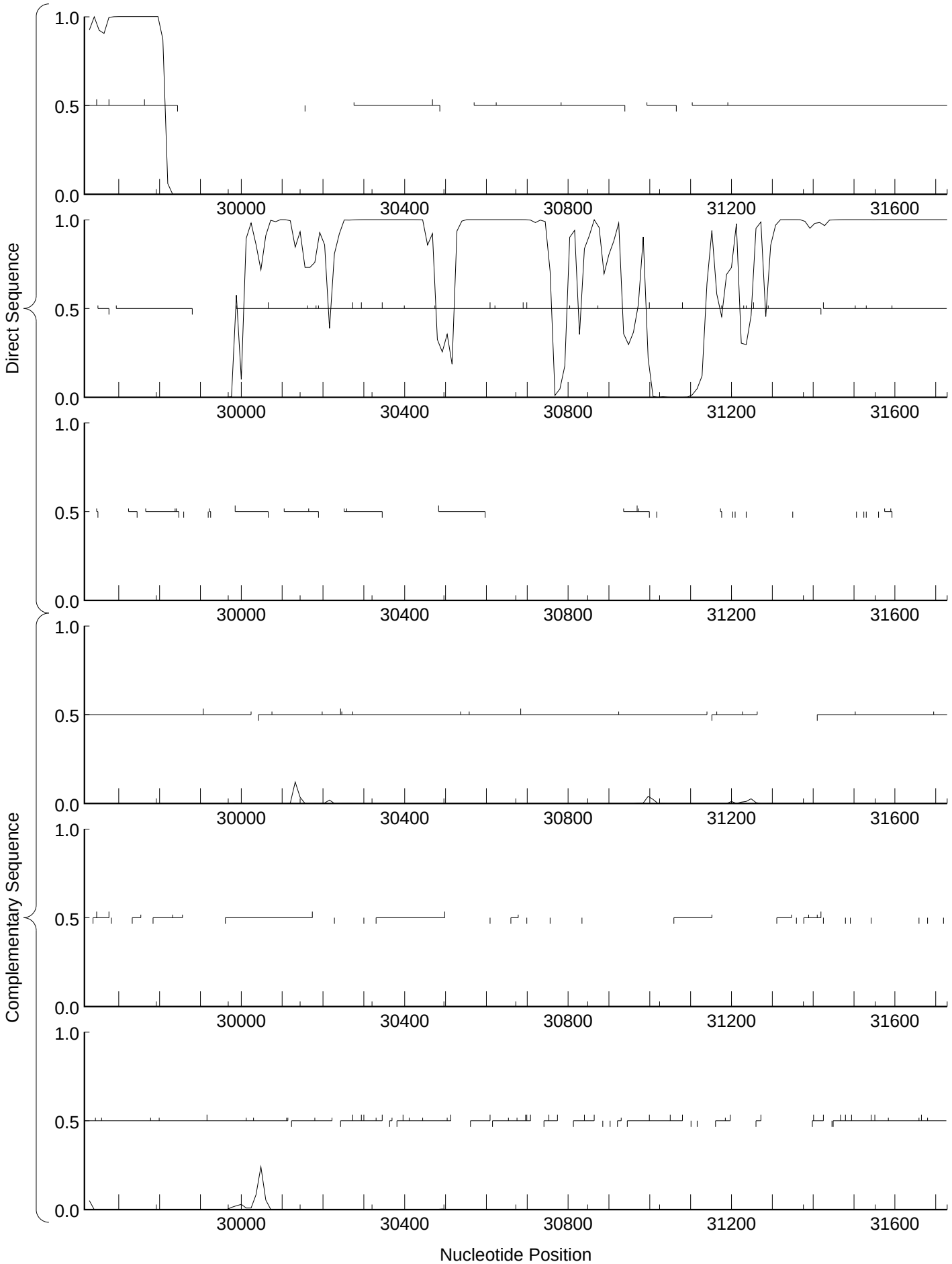

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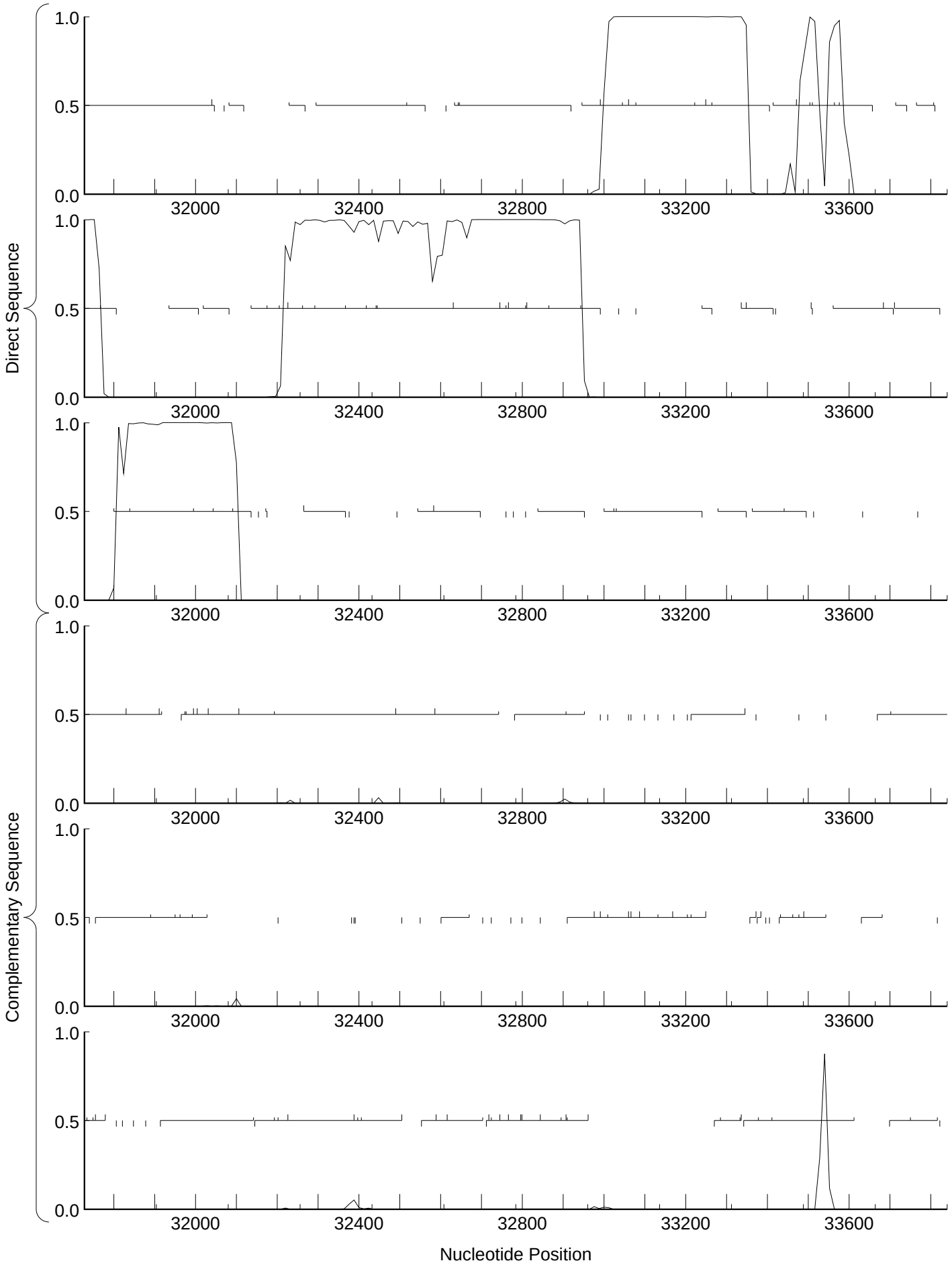

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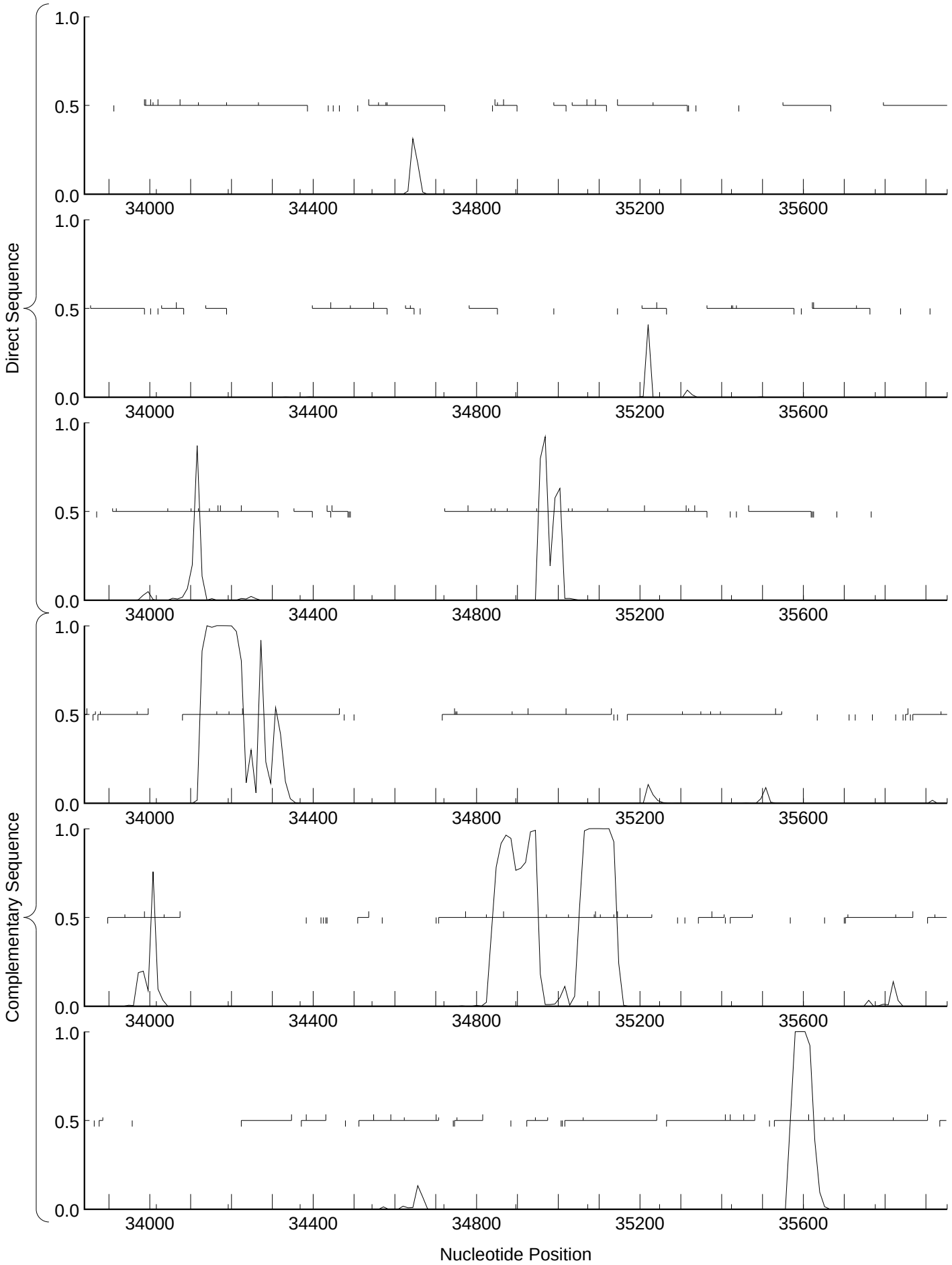

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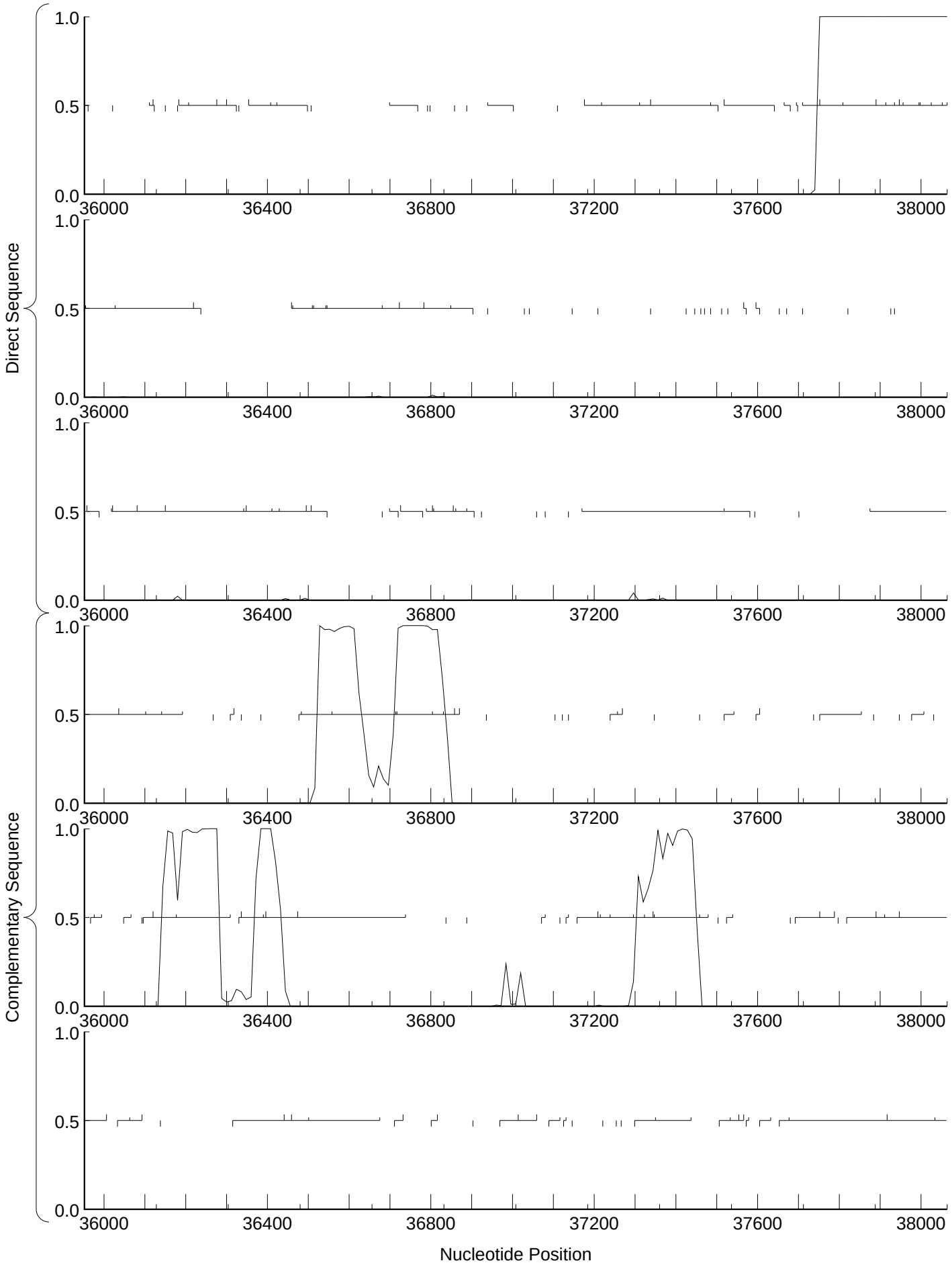

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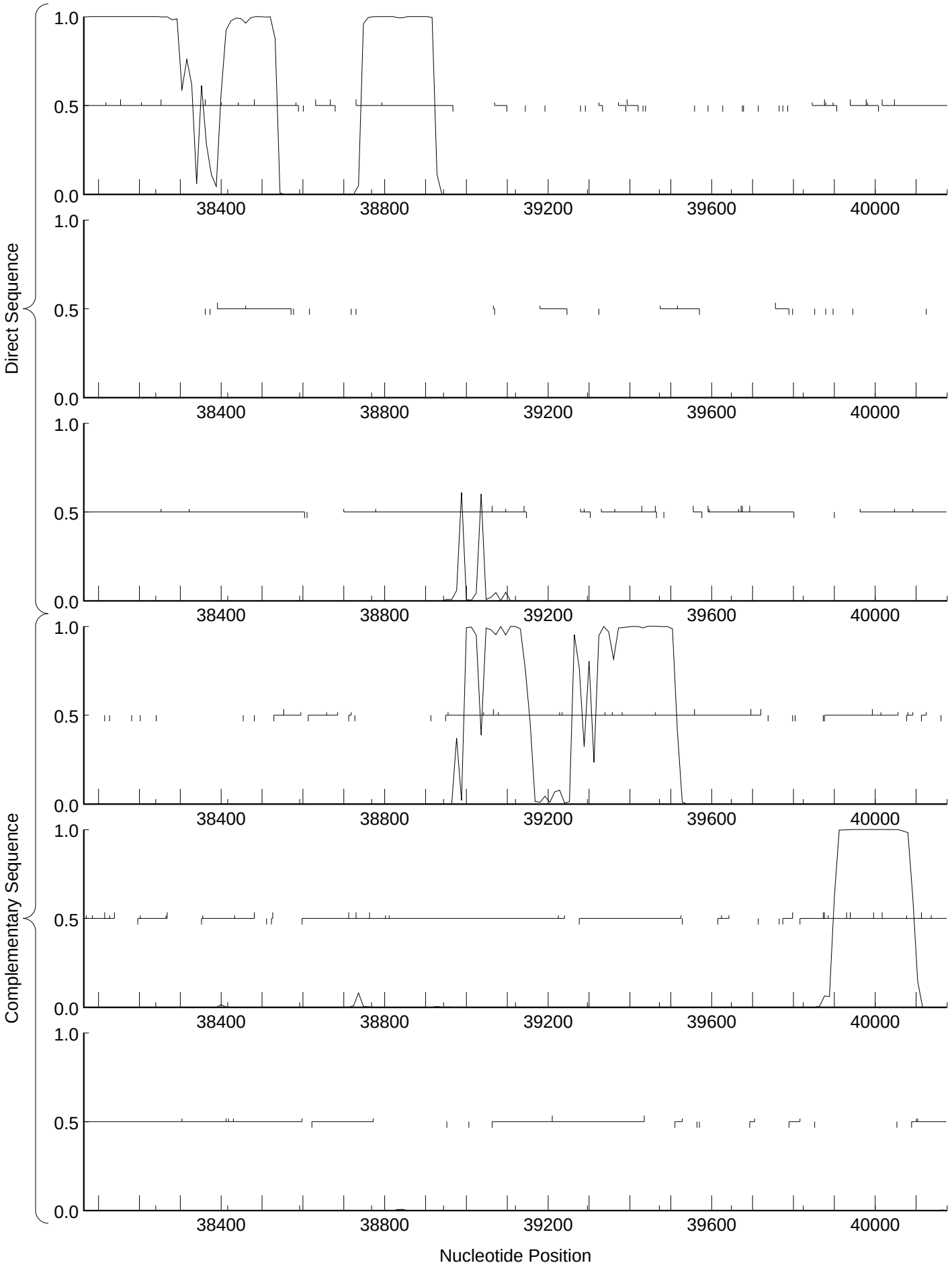

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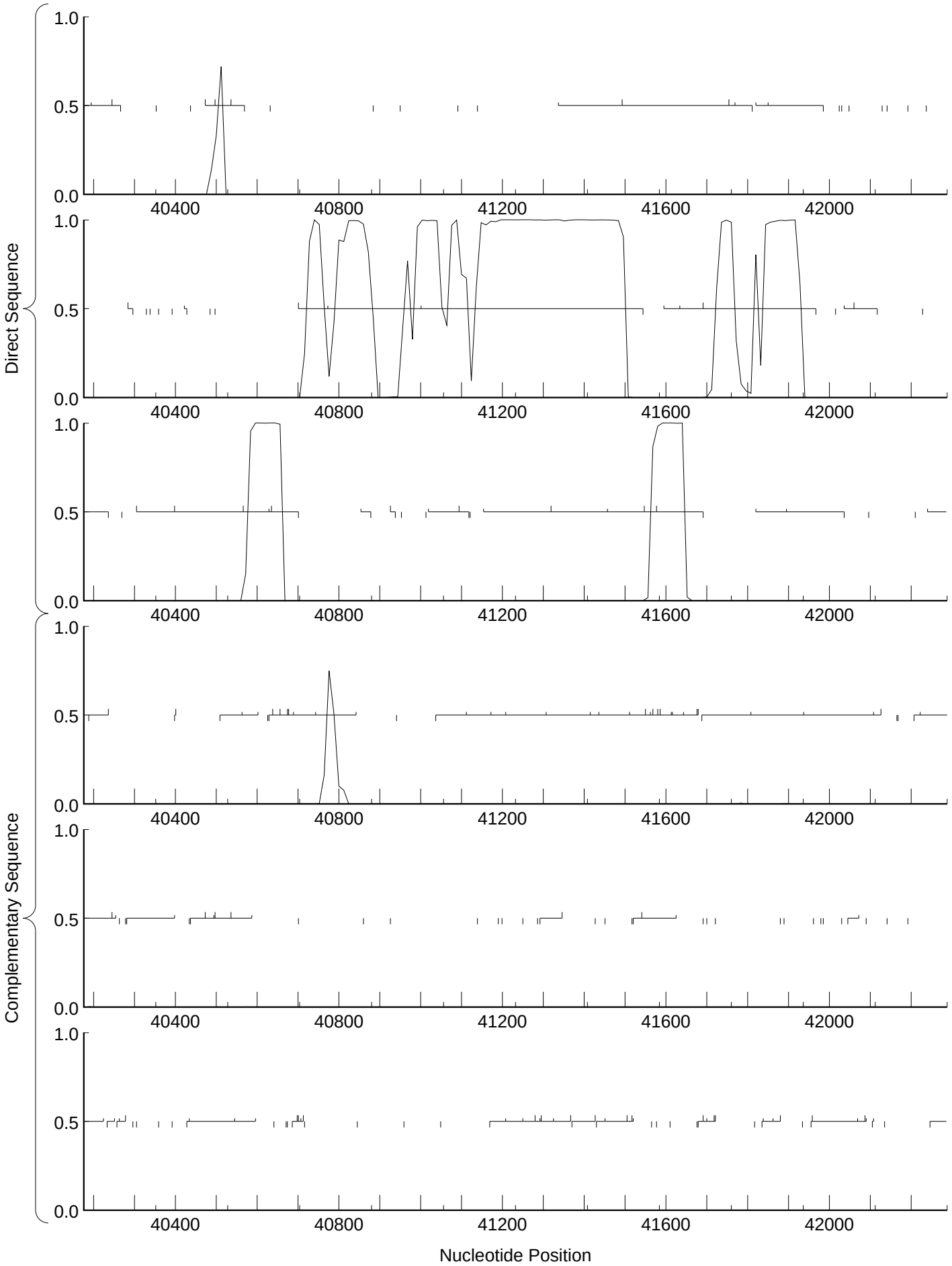

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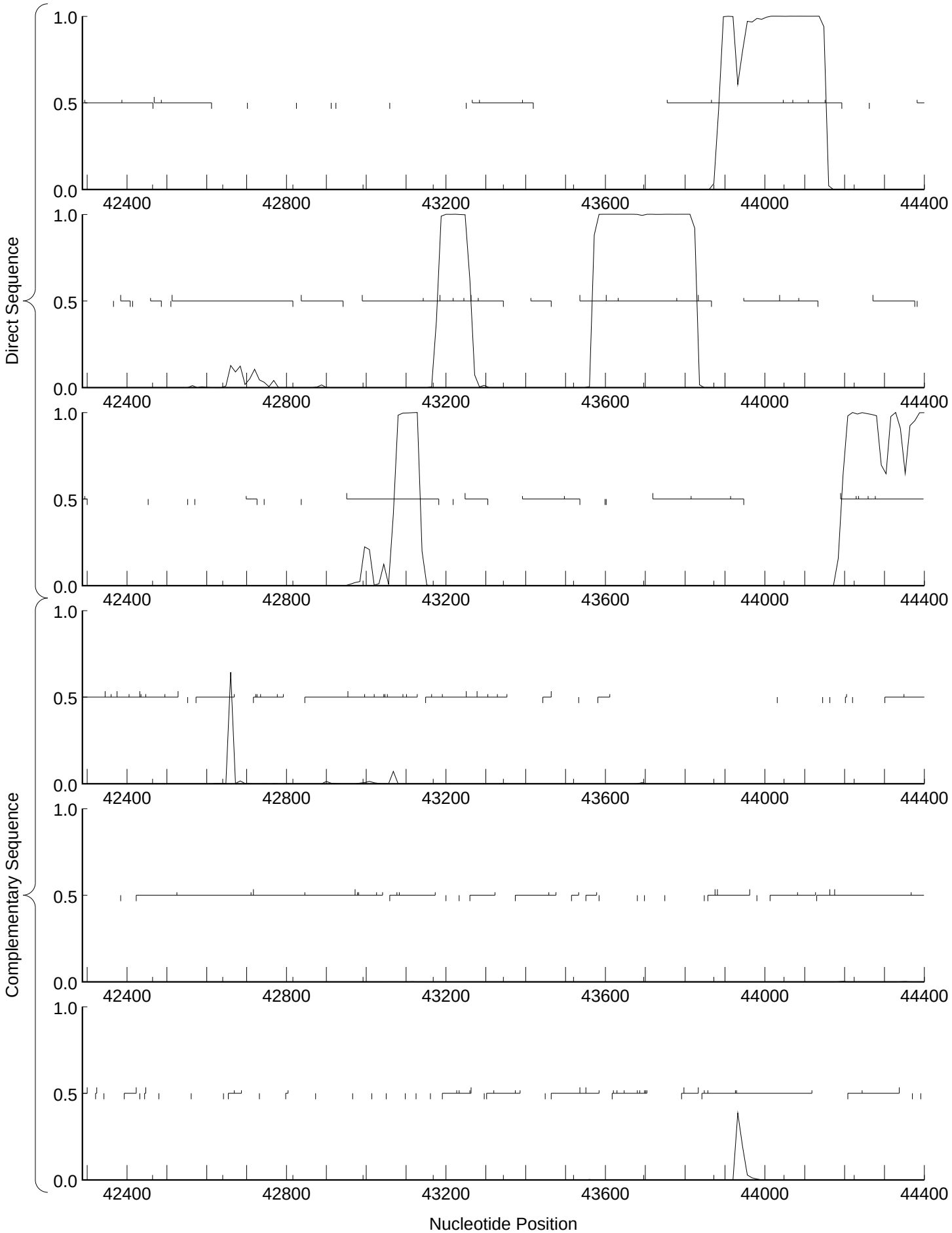

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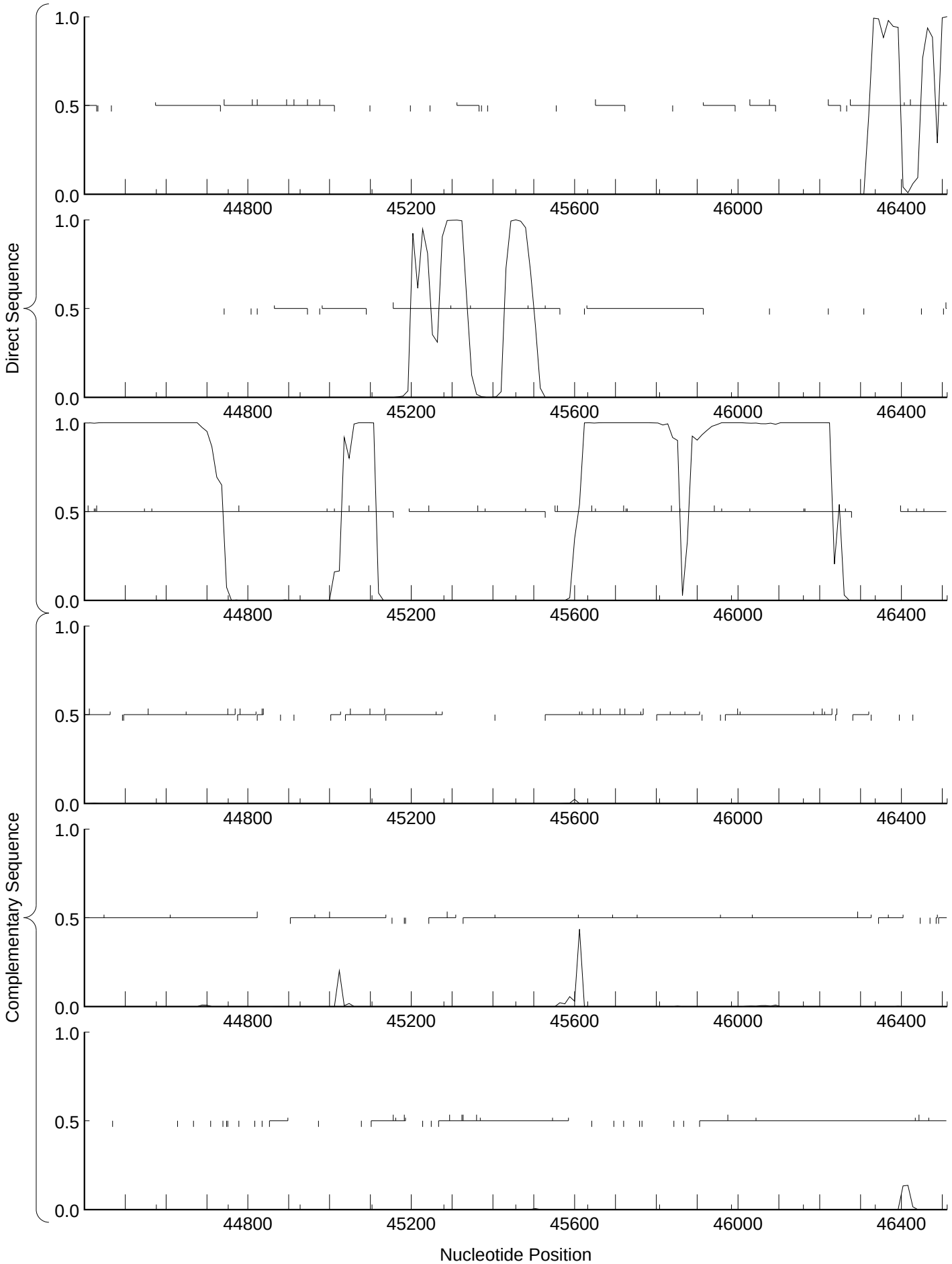

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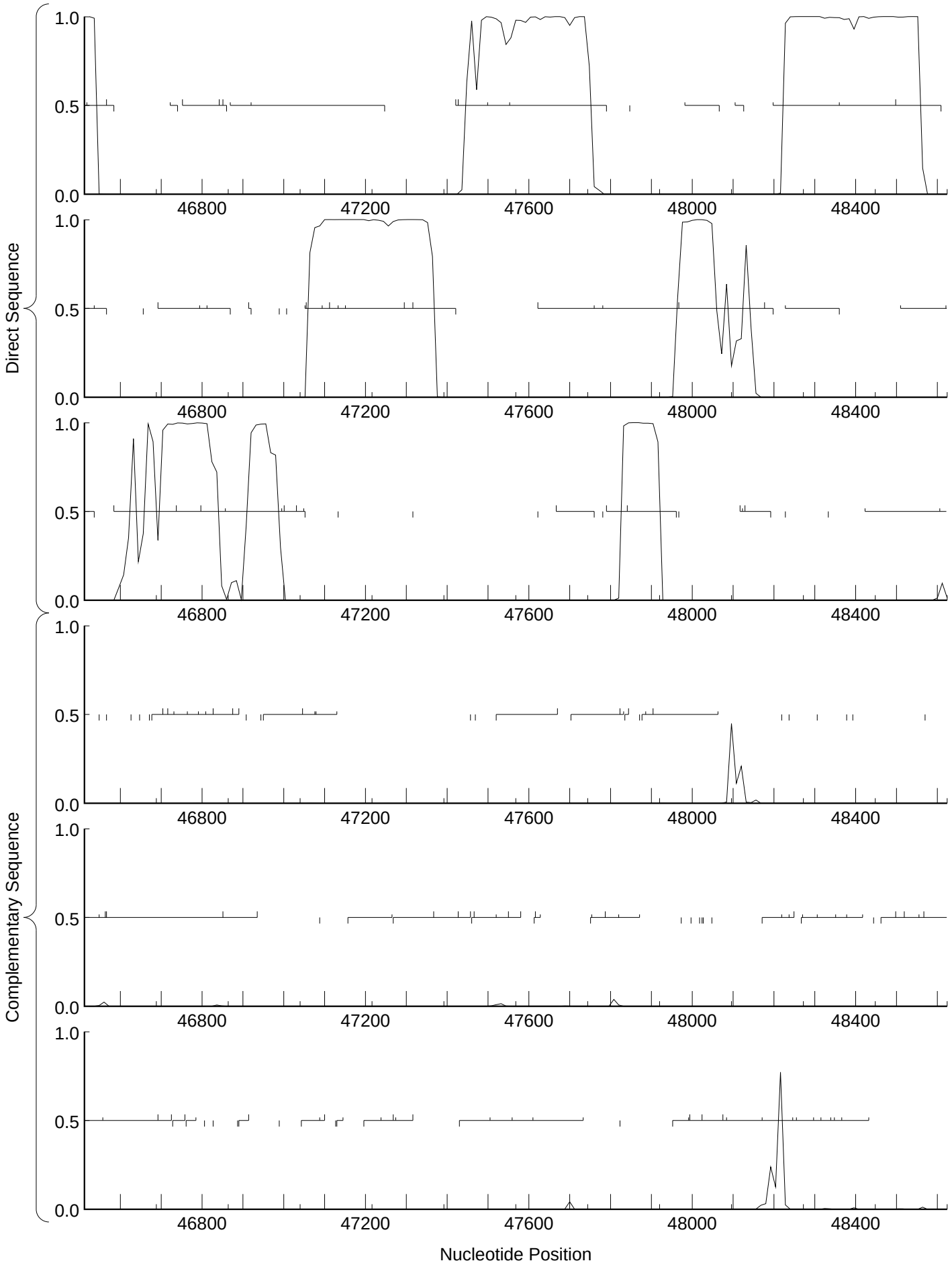

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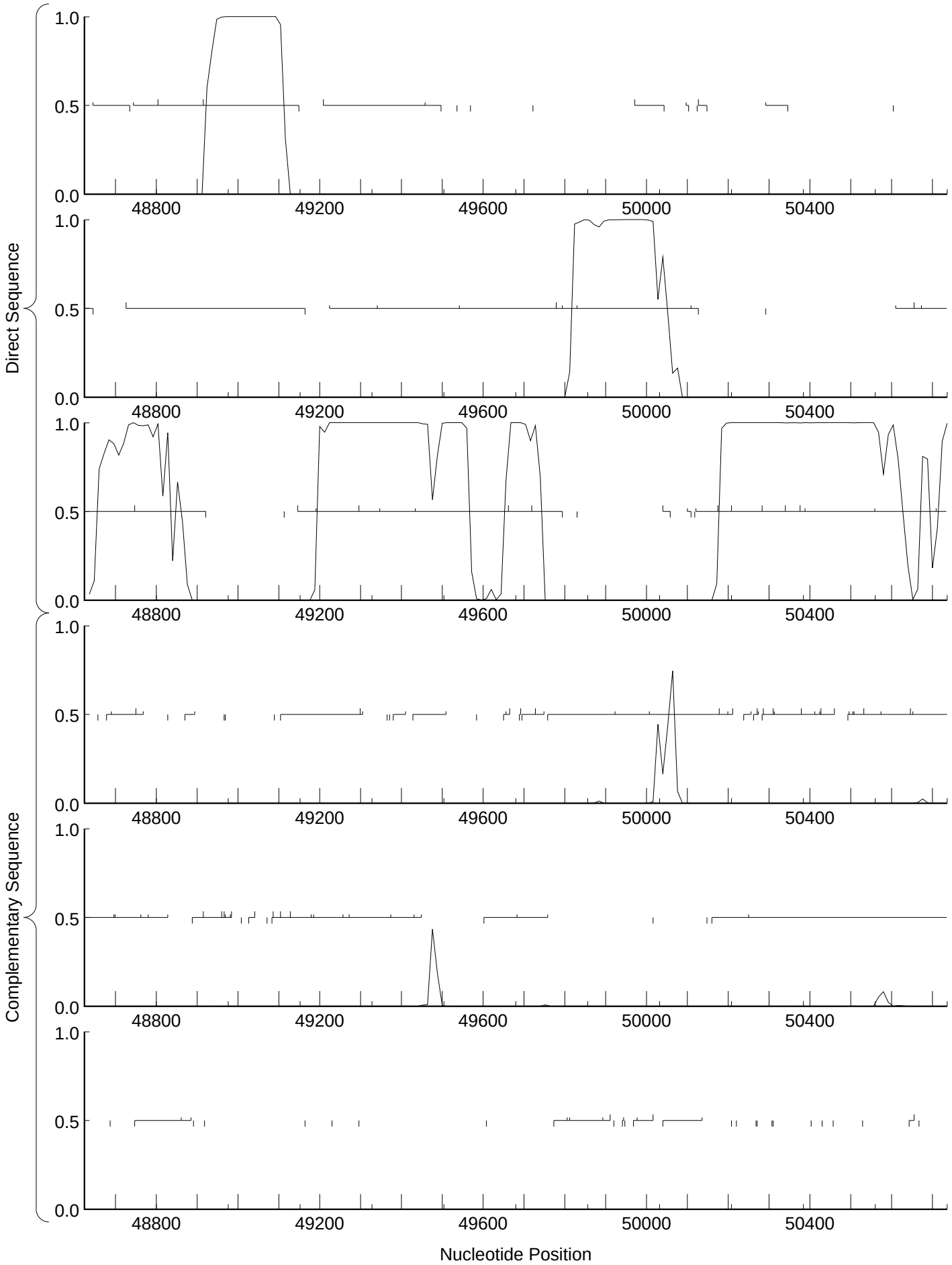

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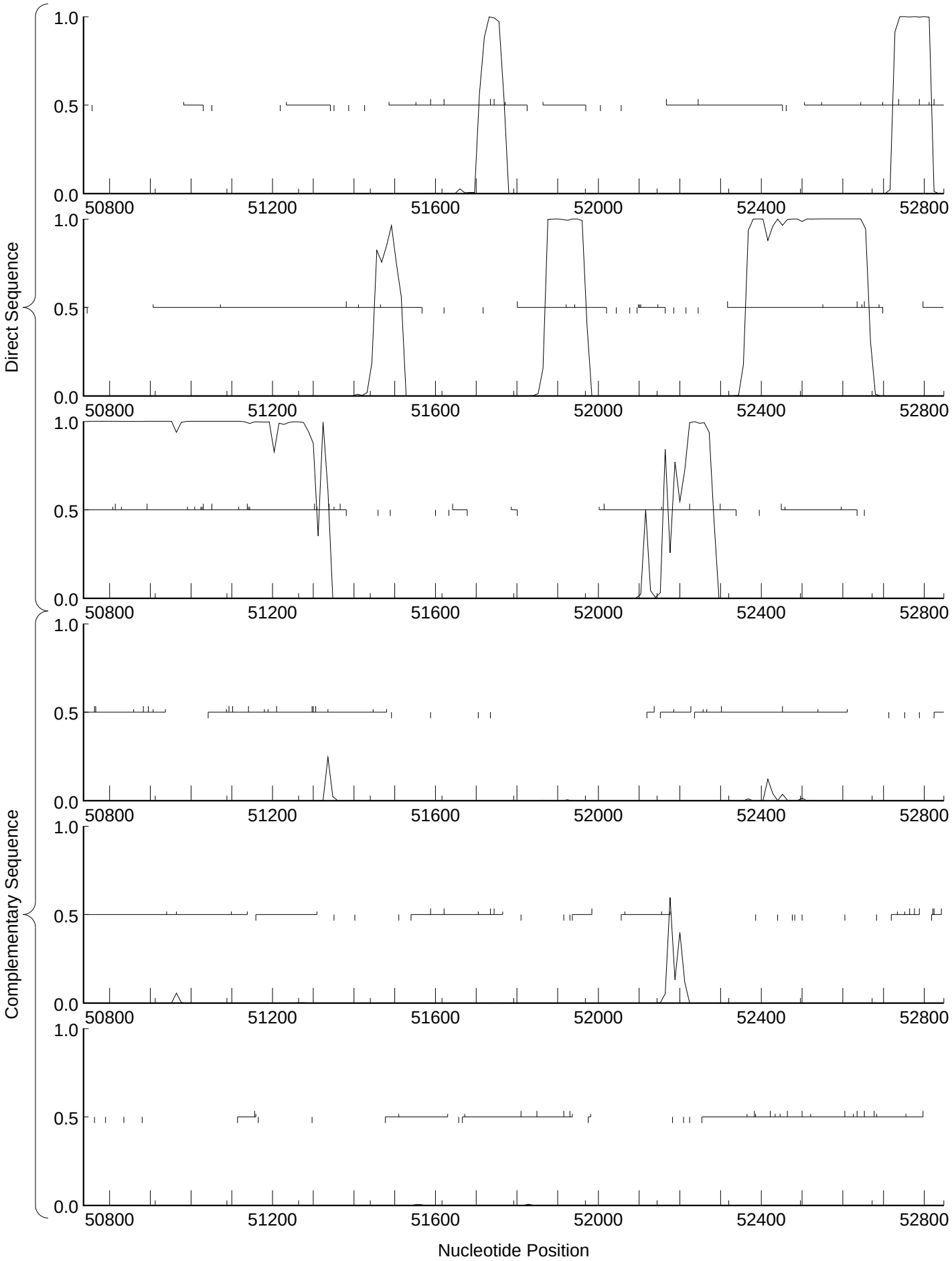

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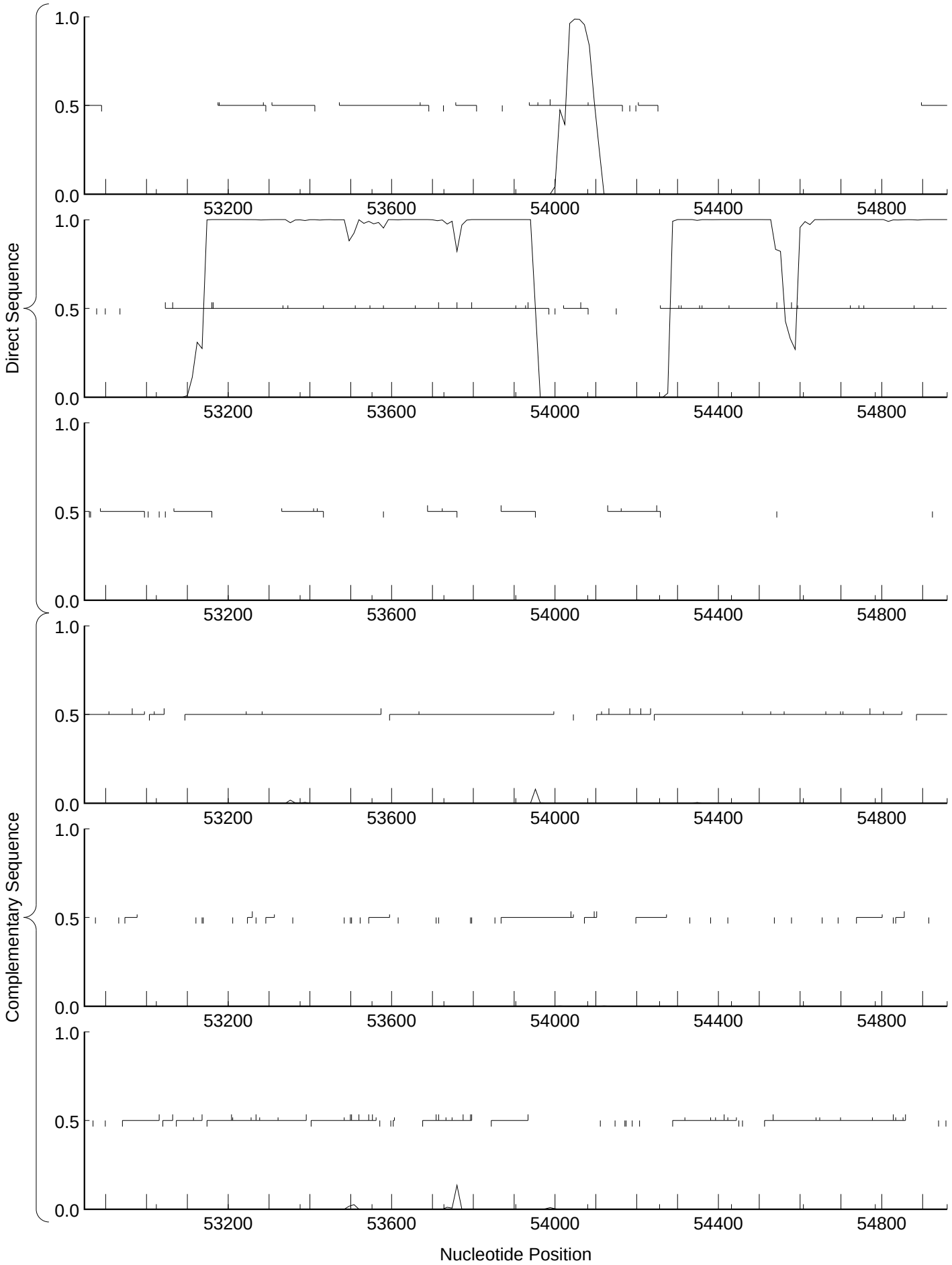

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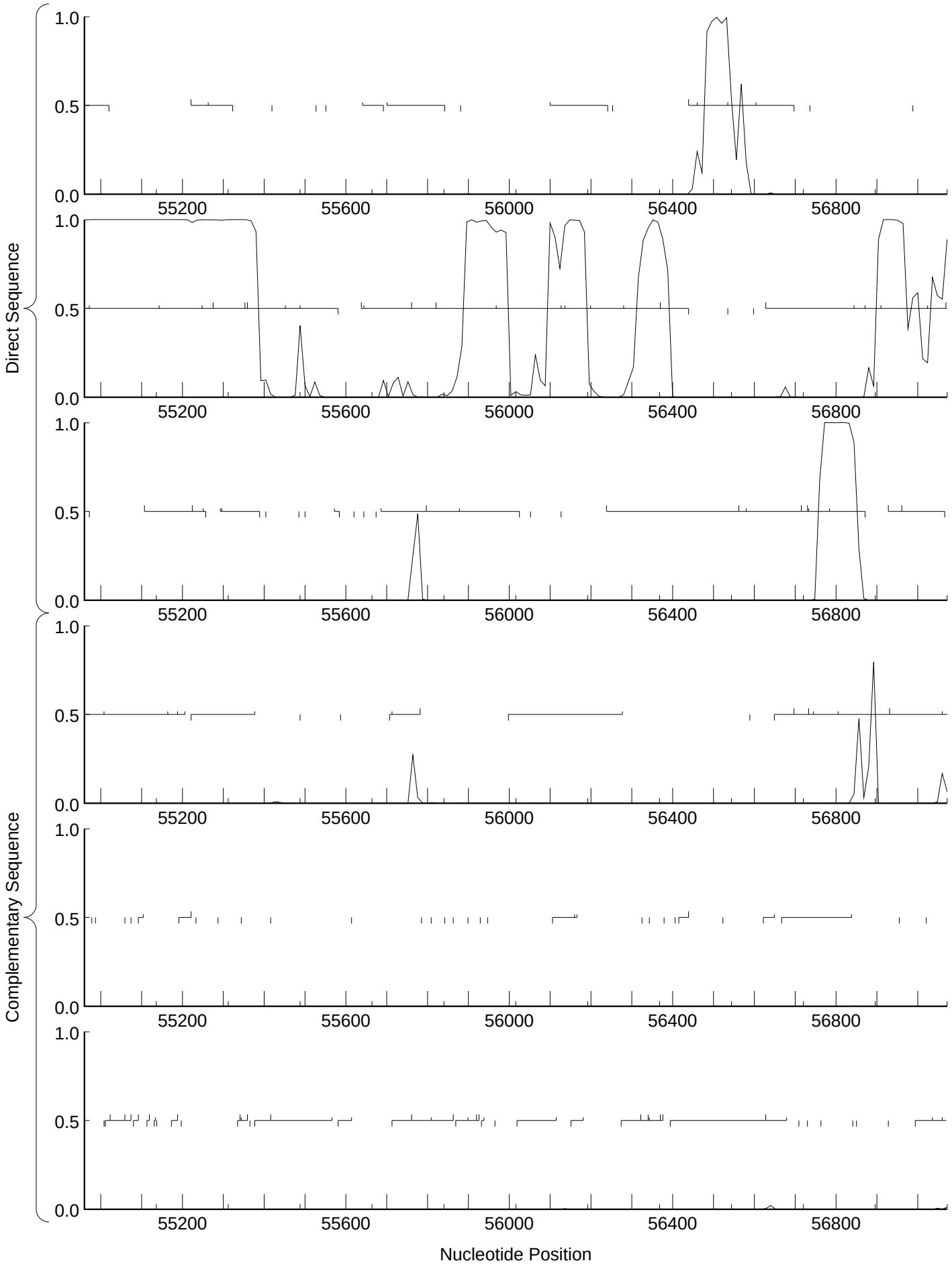

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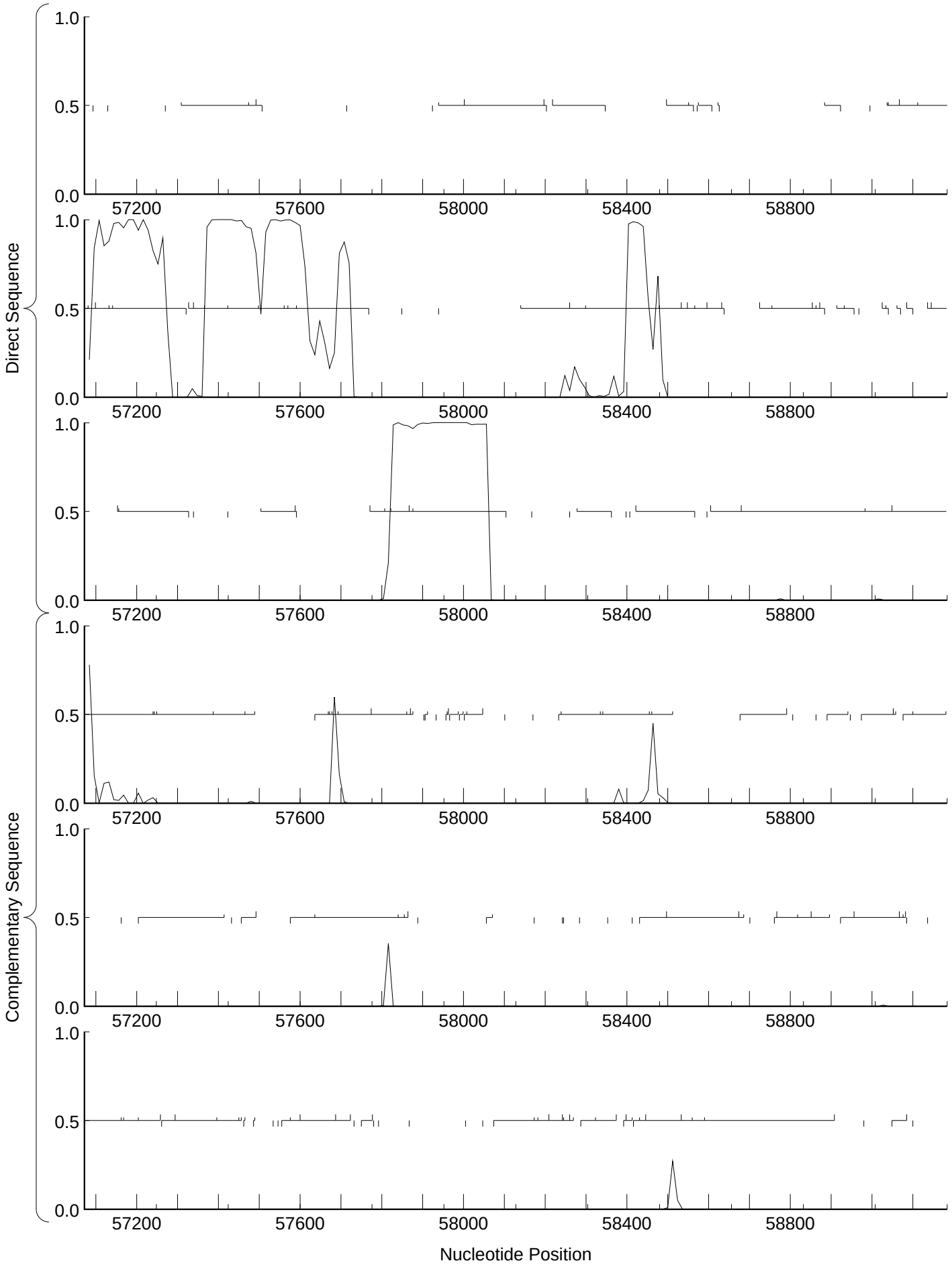

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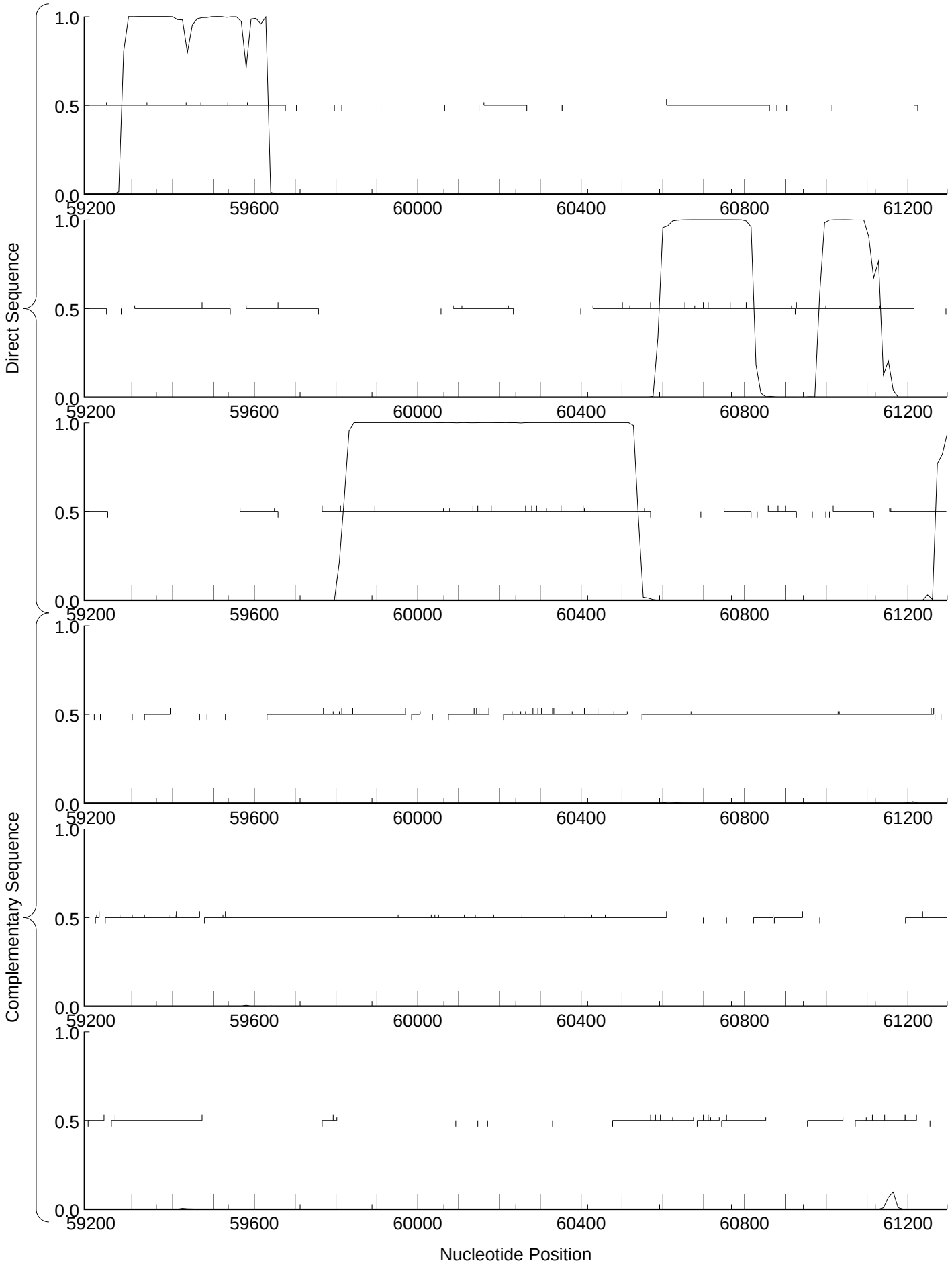

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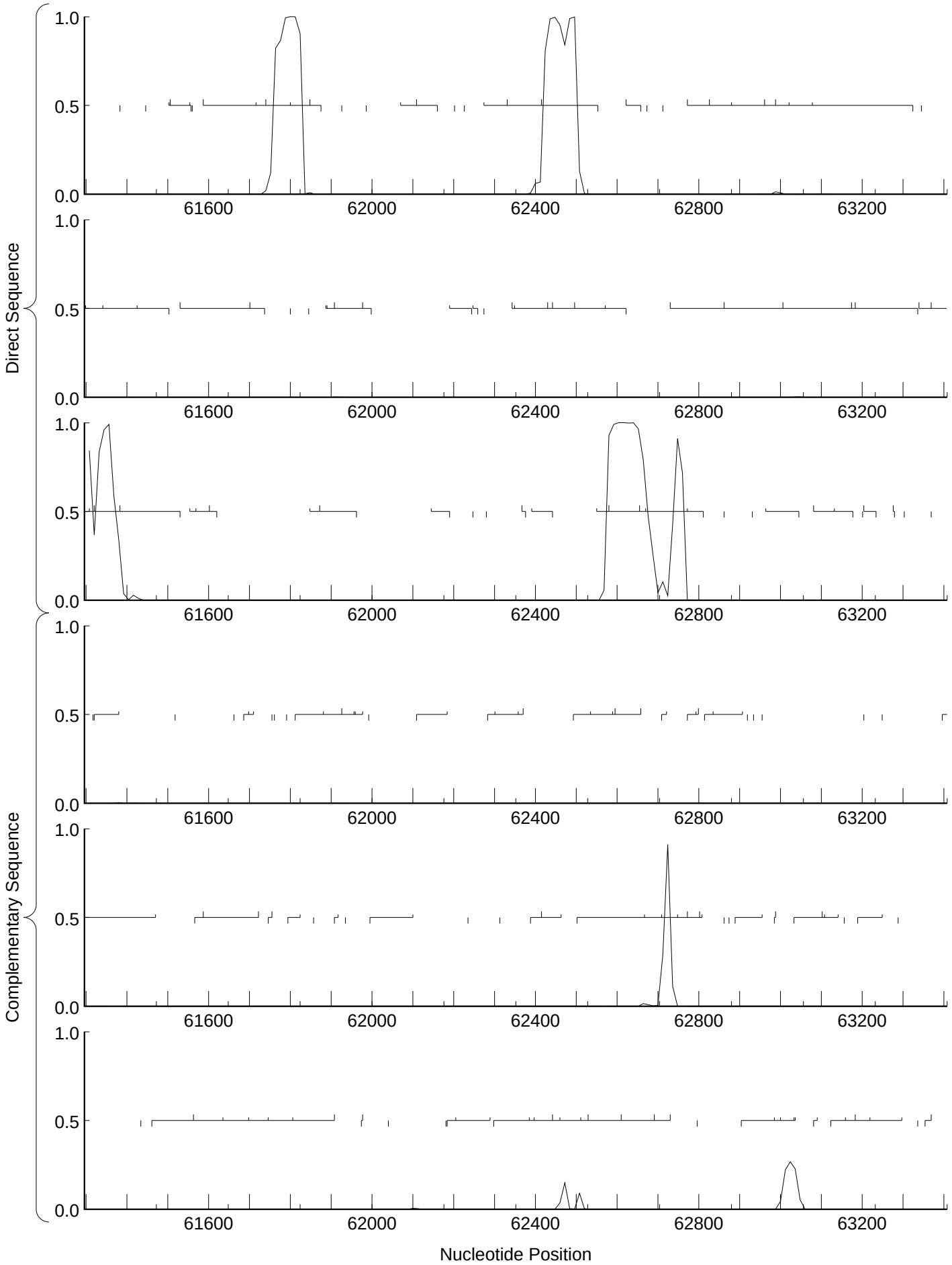

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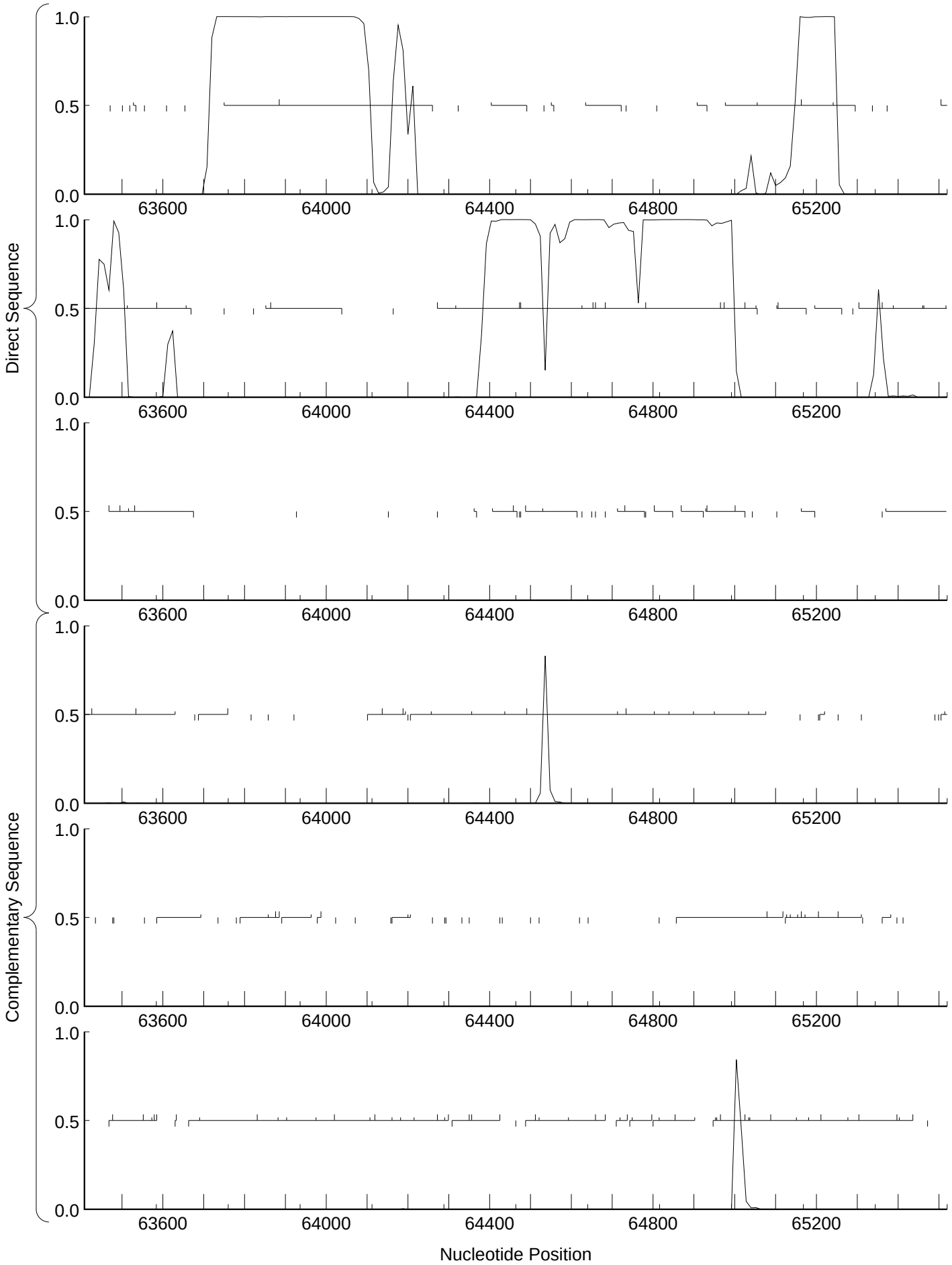

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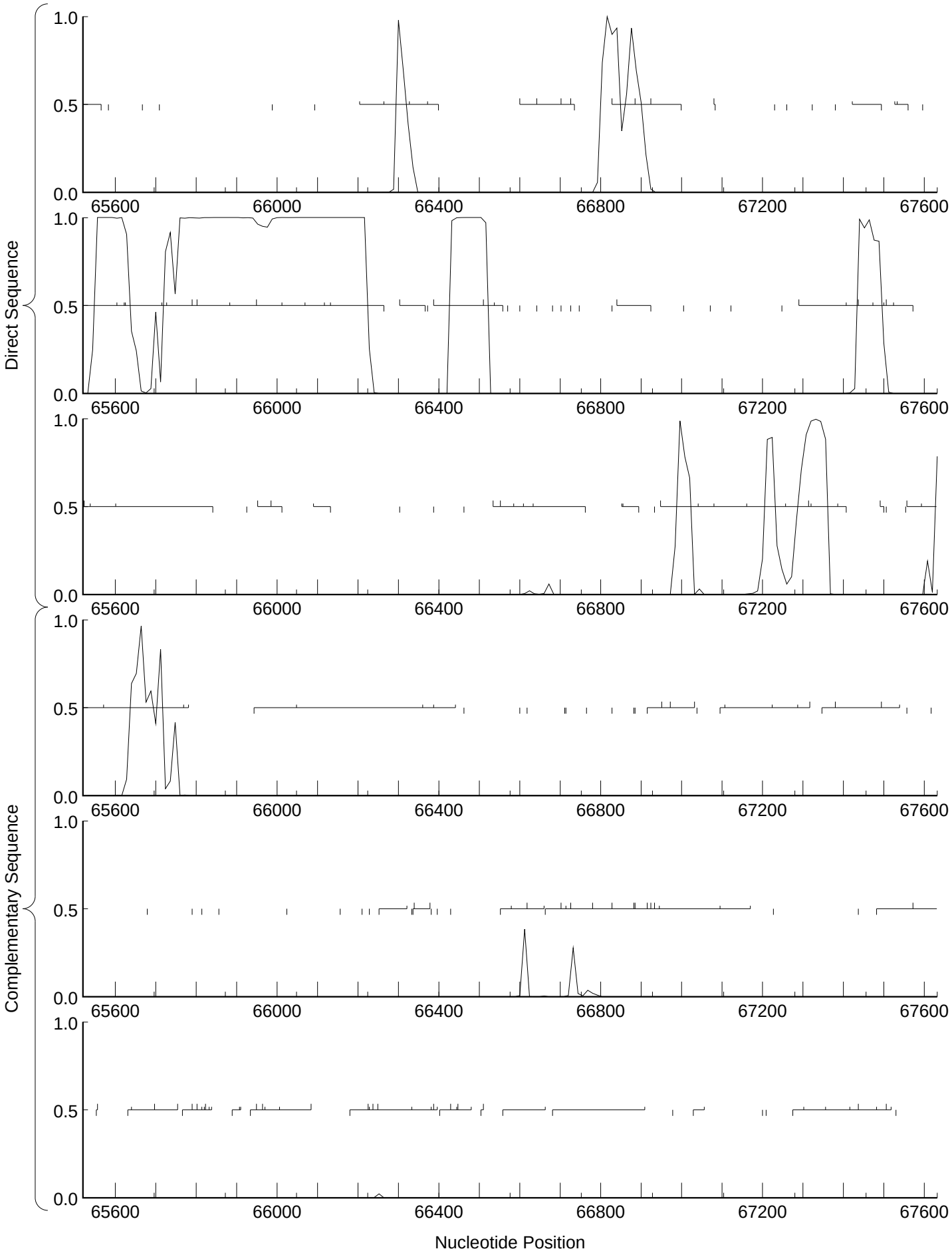

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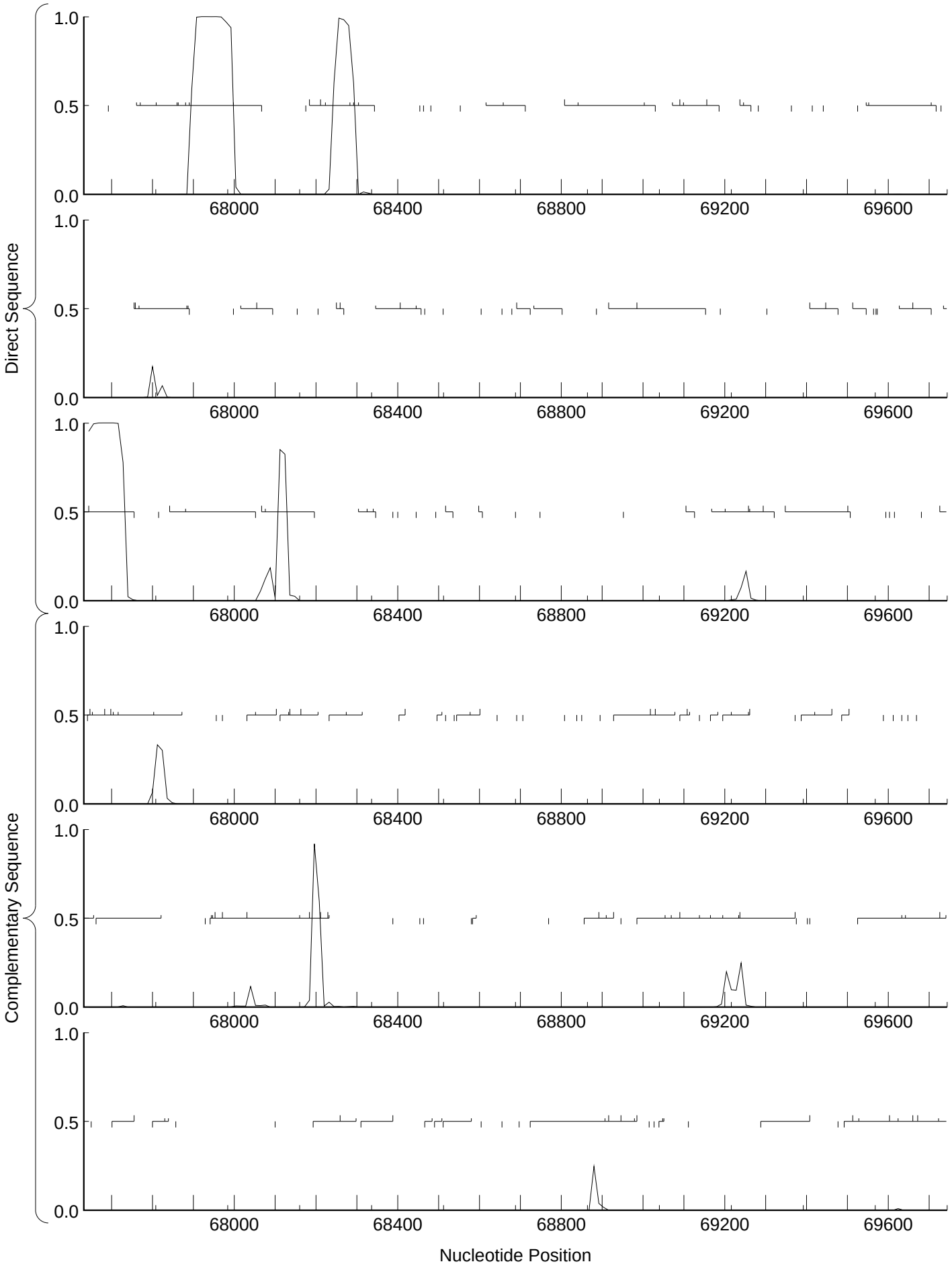

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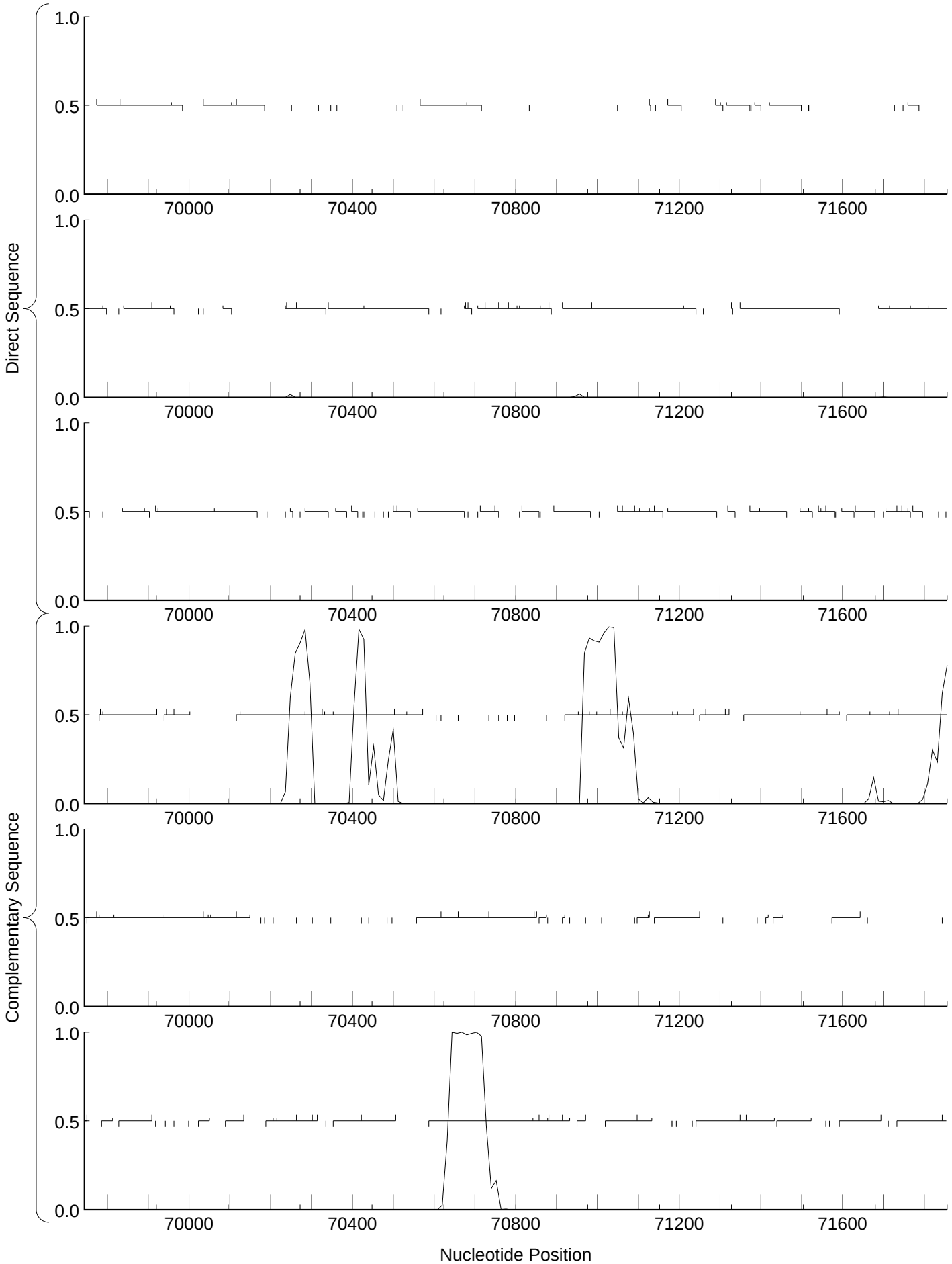

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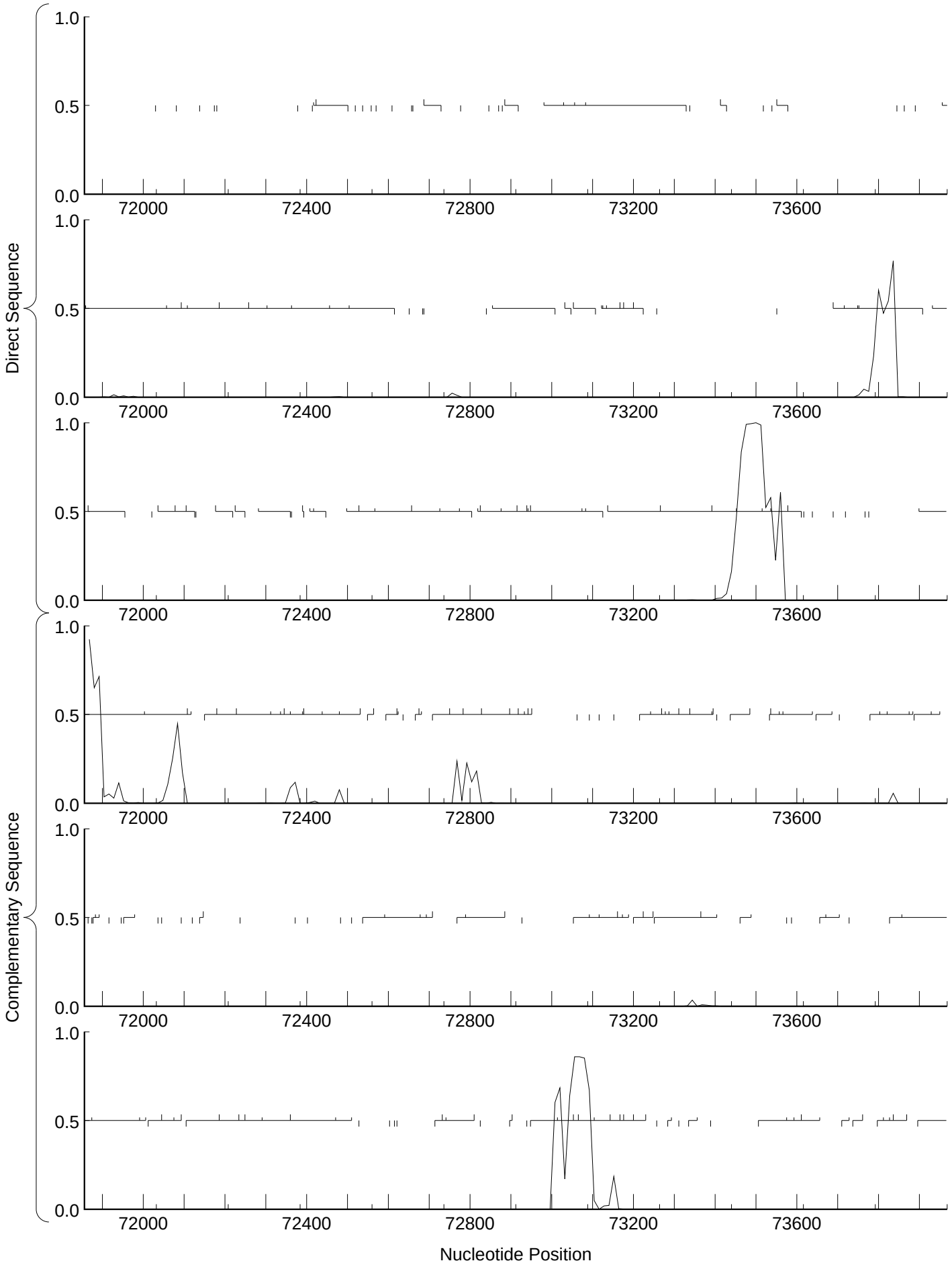

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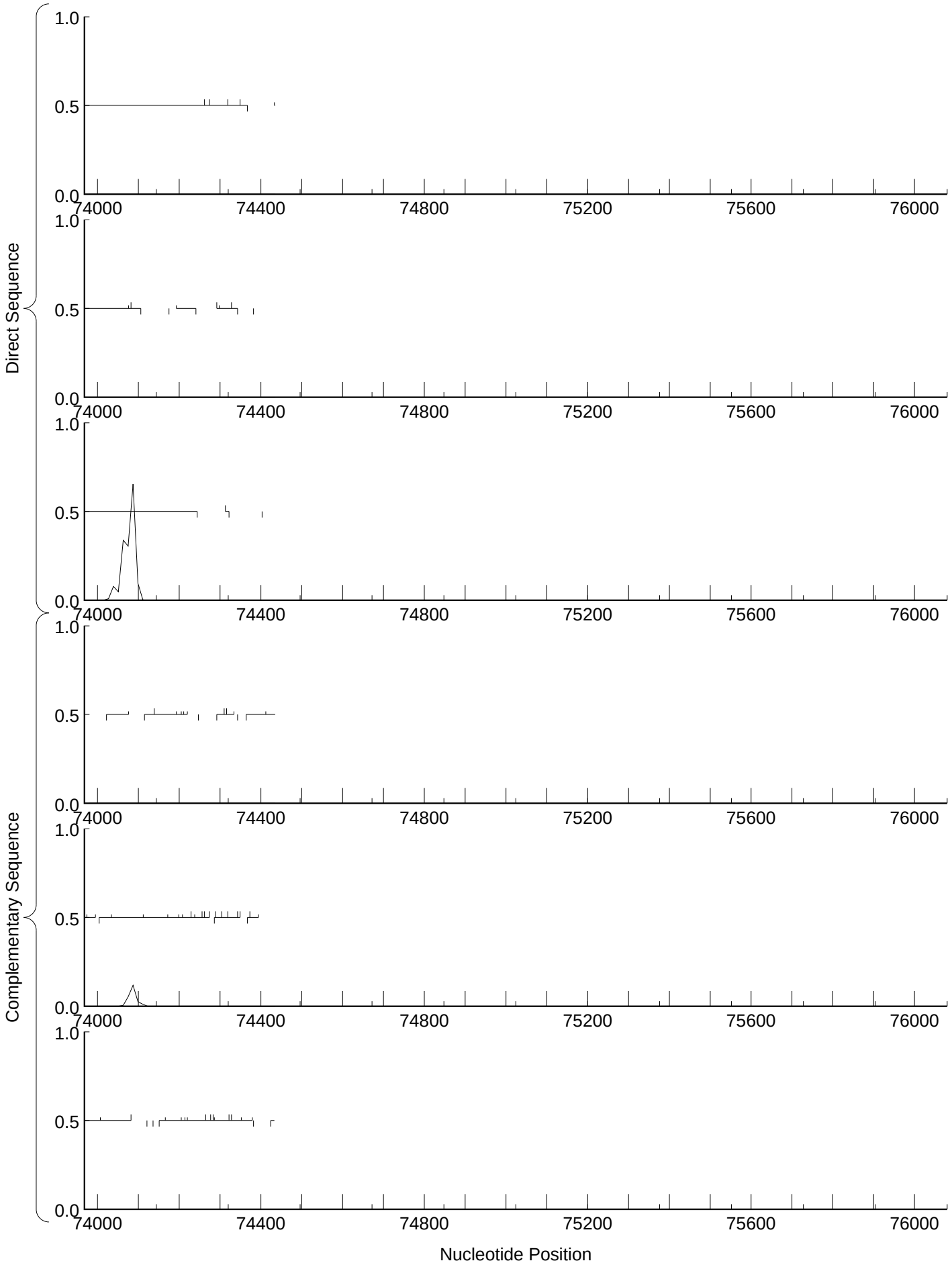

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Nucleotide Position


Direct Sequence
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Nucleotide Position


Direct Sequence
Complementary Sequence
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70800
71200
71600
Nucleotide Position


Direct Sequence
Complementary Sequence
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73600
Nucleotide Position


Direct Sequence
Complementary Sequence
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Nucleotide Position