GeneMark

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

PROGRAM INFORMATION

```
Sequence         :  Streptomyces phage Sham complete sequence, 130327 bp including 787 bp terminal repeat, Clus
Analysis Date    :  3/29/24 at 4:34:39
Pages            :  63
Sequence Length  :  130327 bp
GC Content       :   47.07%

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

PS-Version       :  1.2

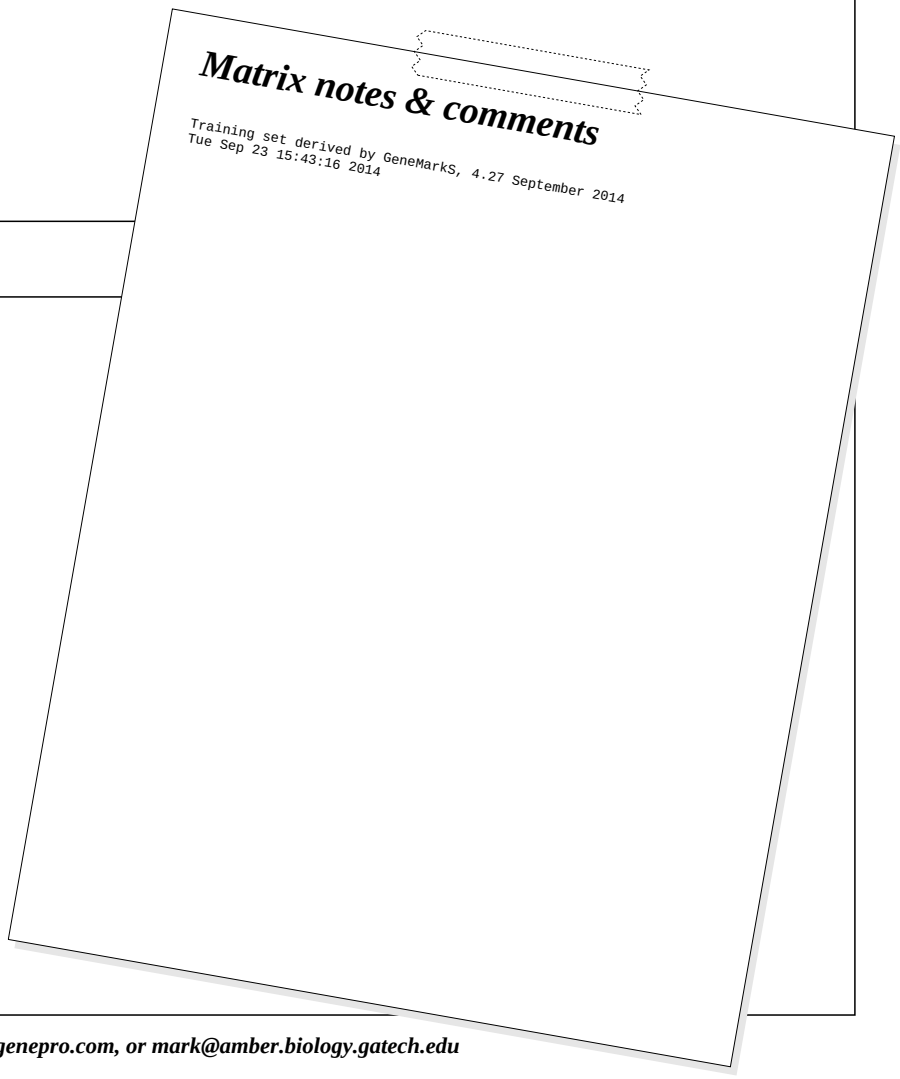
GeneMark Options :  PostScript graph,
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MATRIX INFORMATION

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Author     :  -

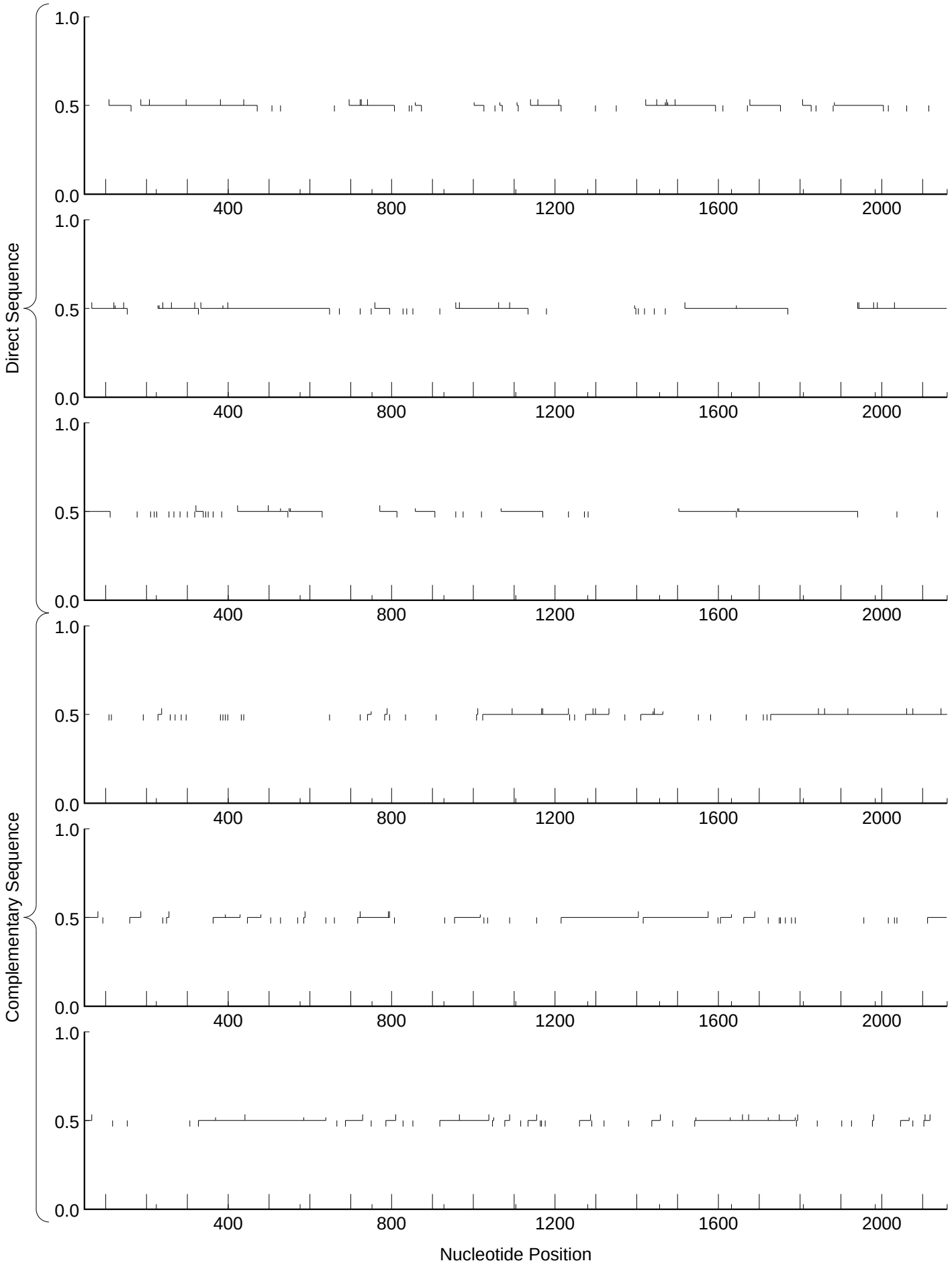
Order      :  4


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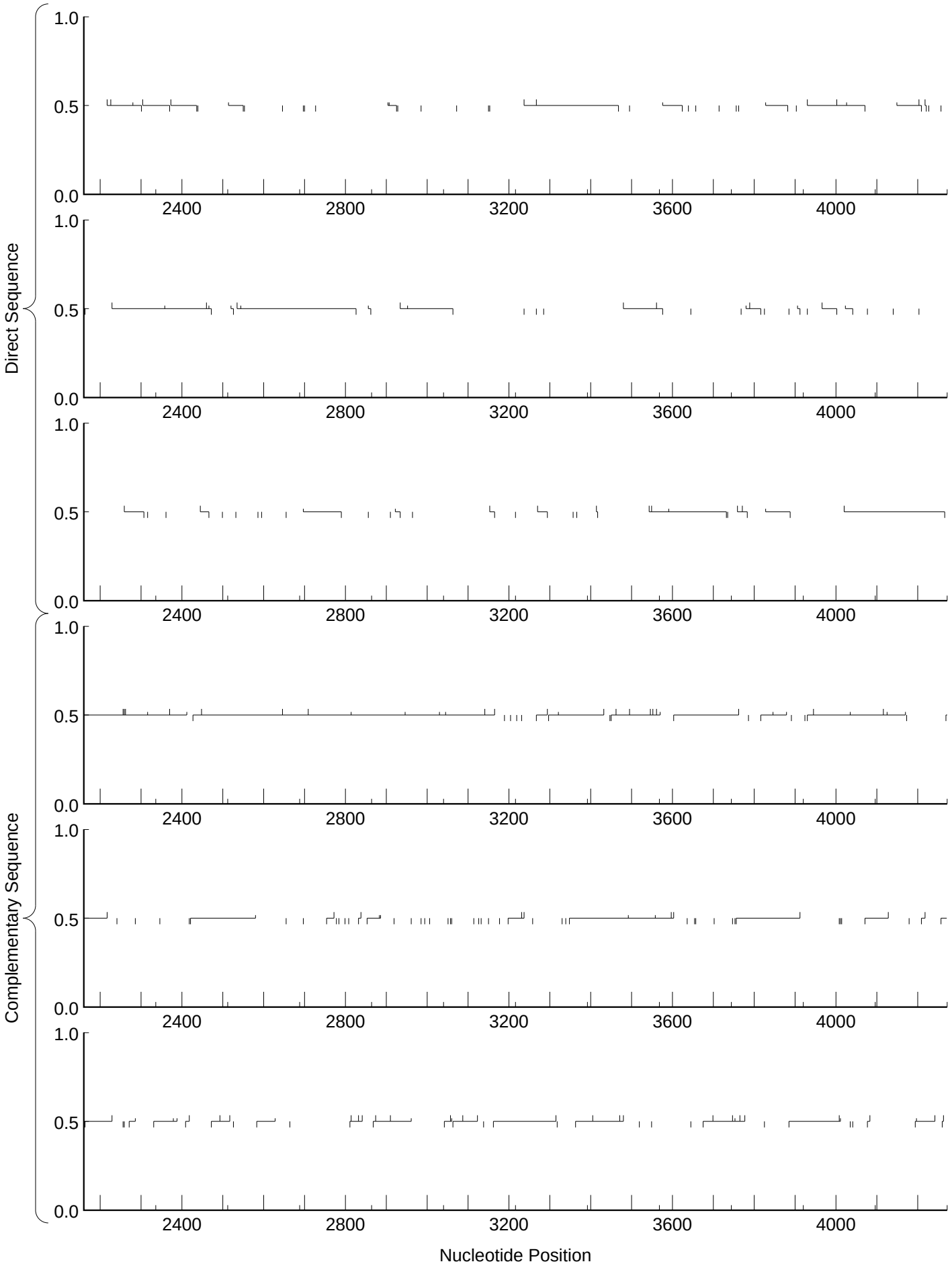


For help or information, send electronic mail to: custserv@genepro.com, or mark@amber.biology.gatech.edu

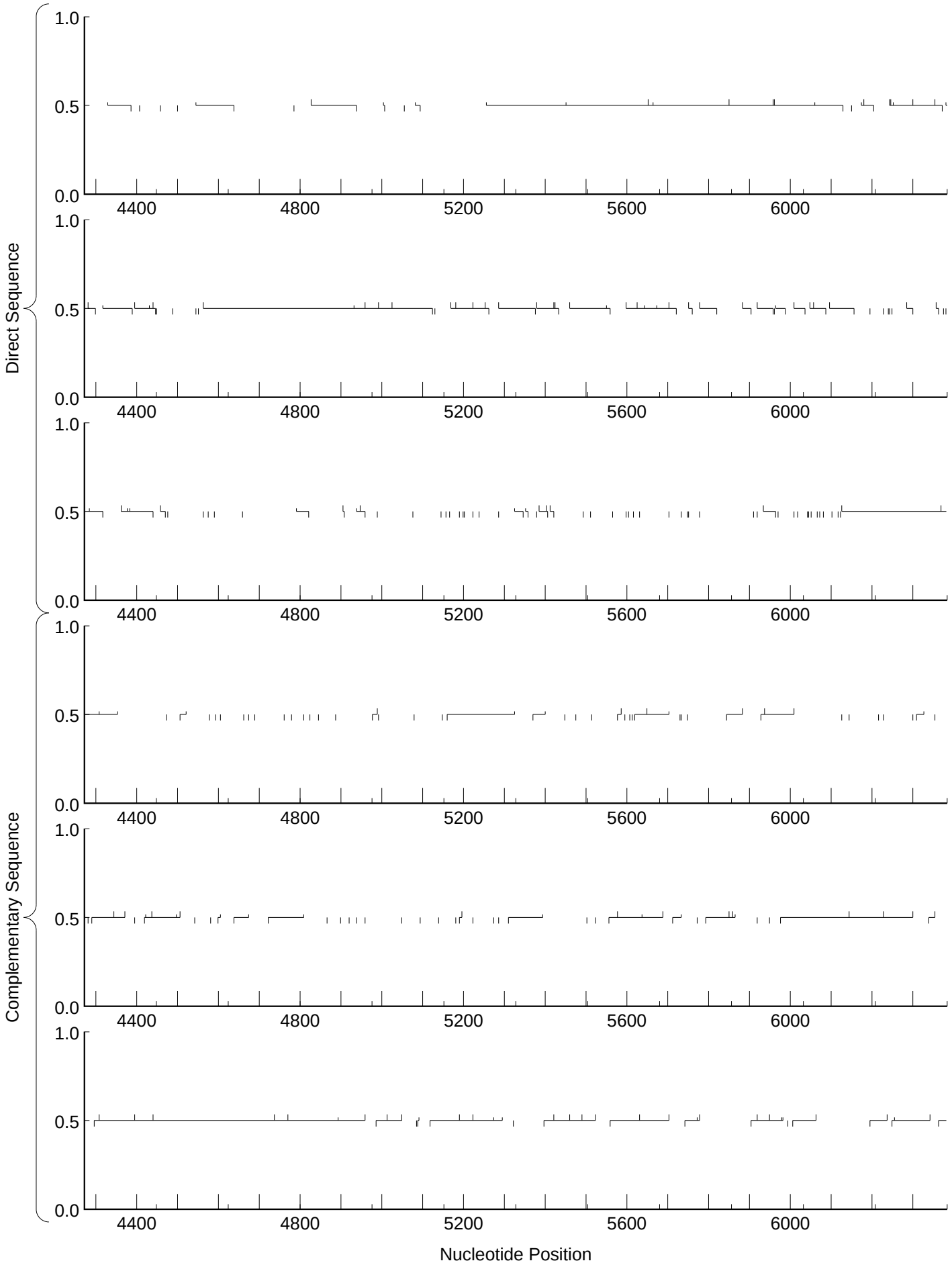
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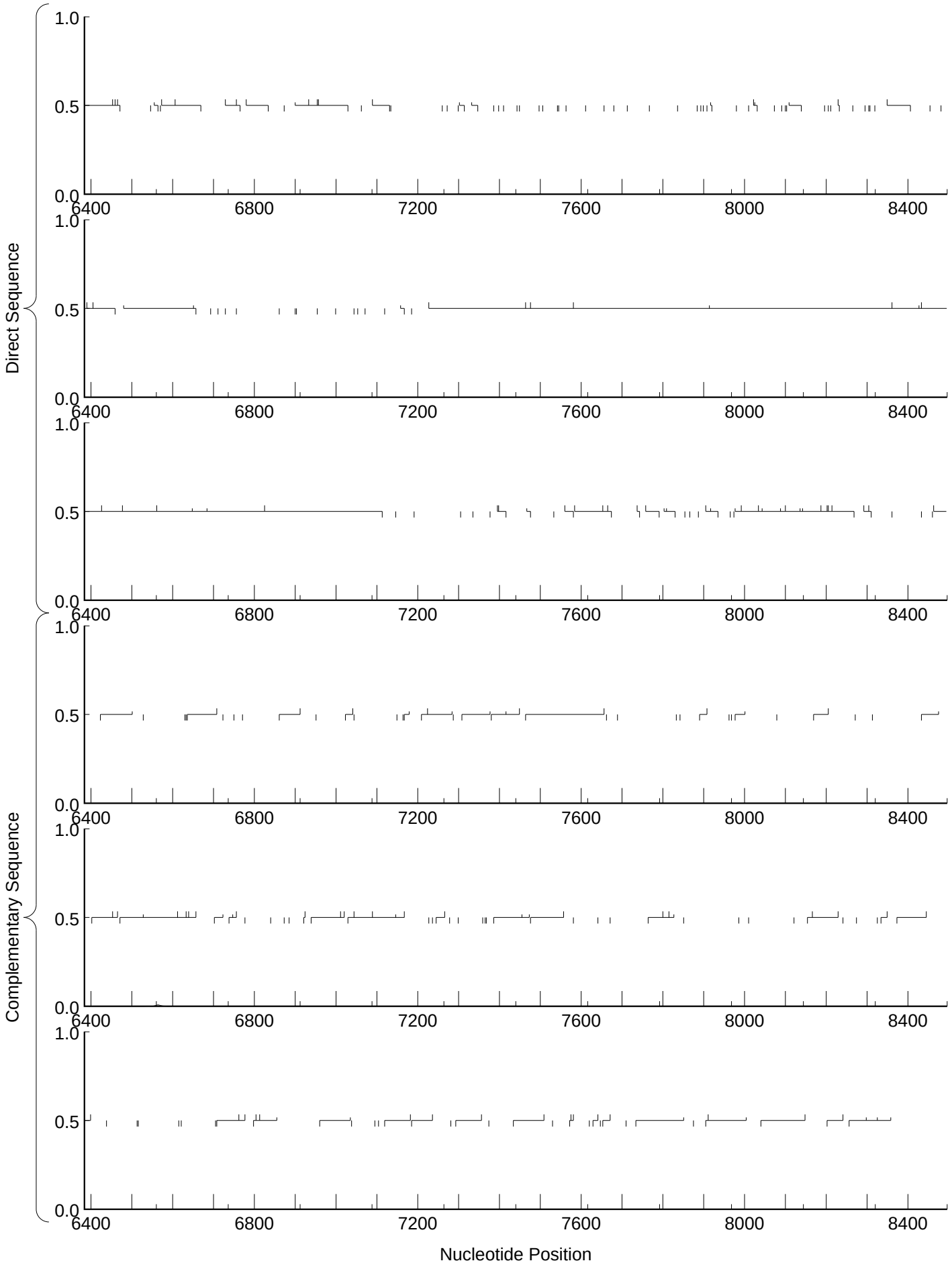
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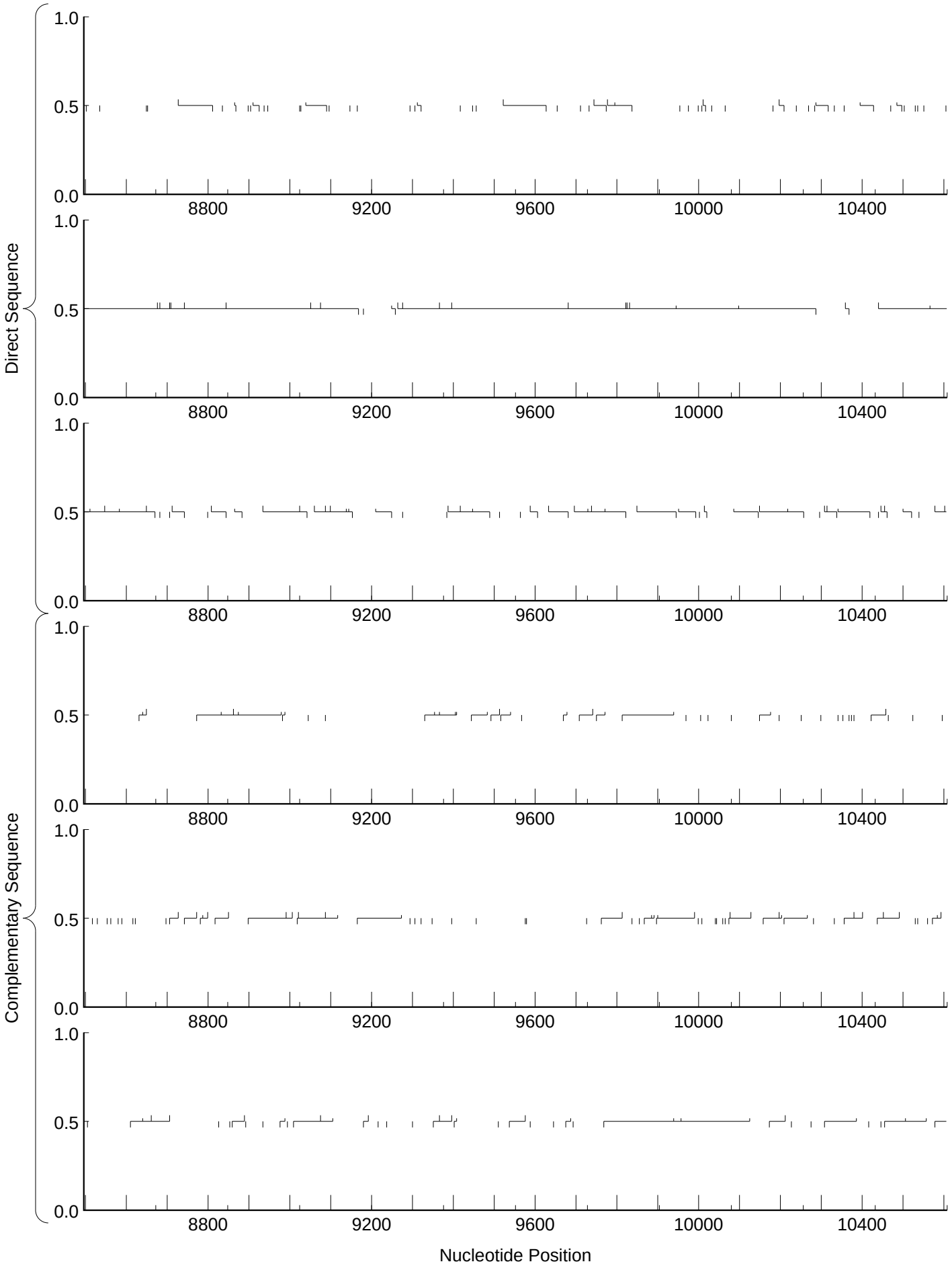
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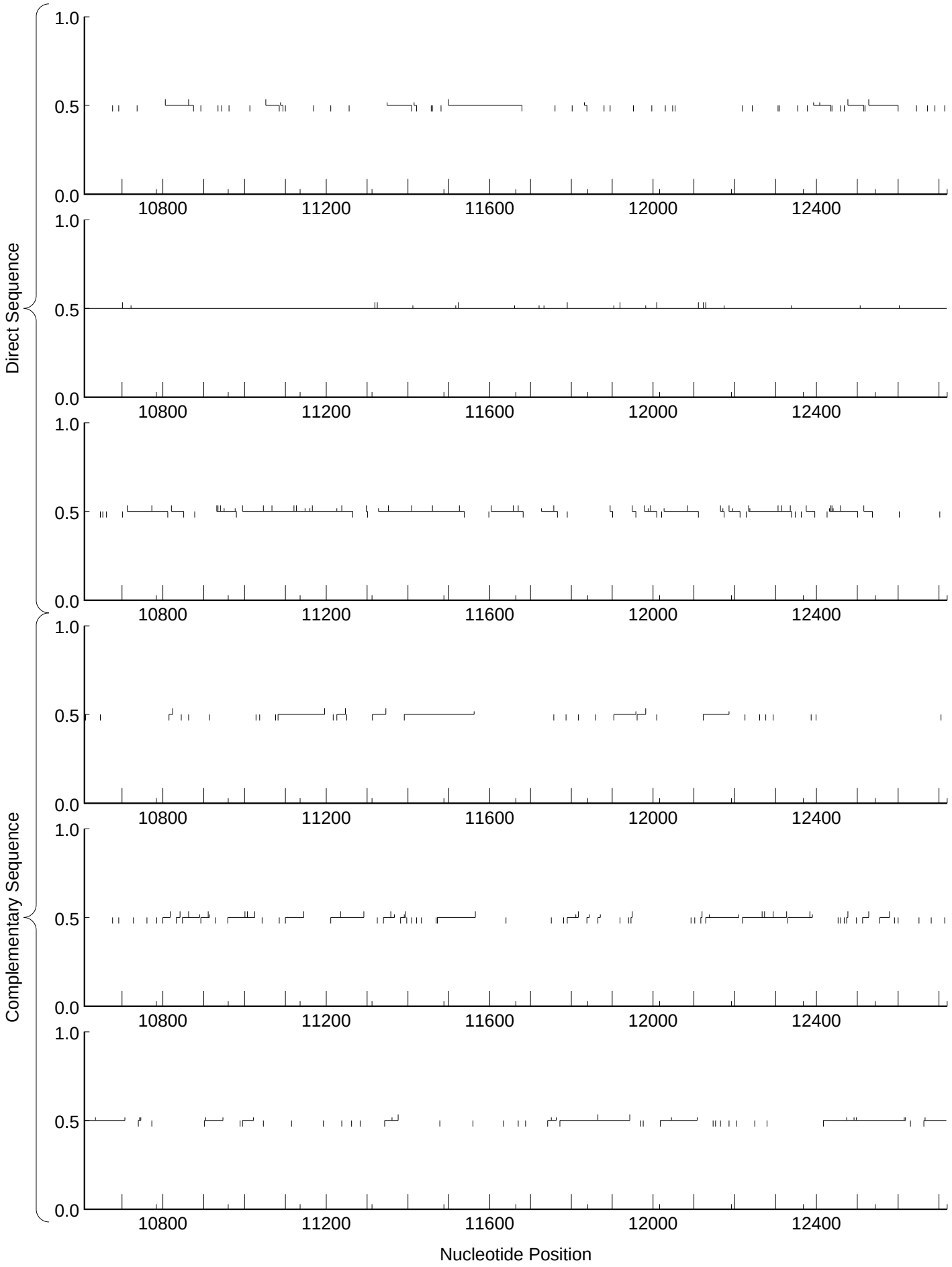
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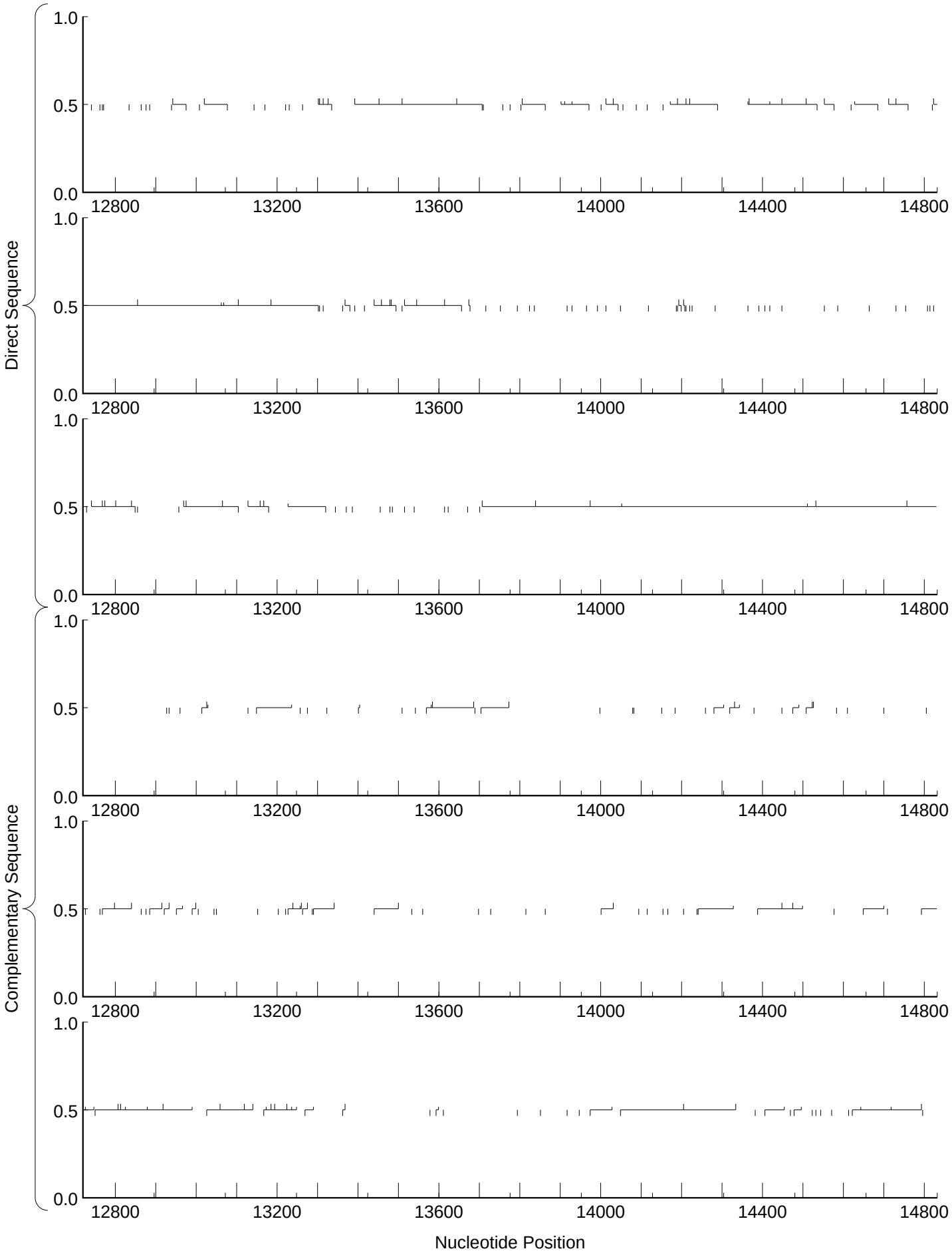
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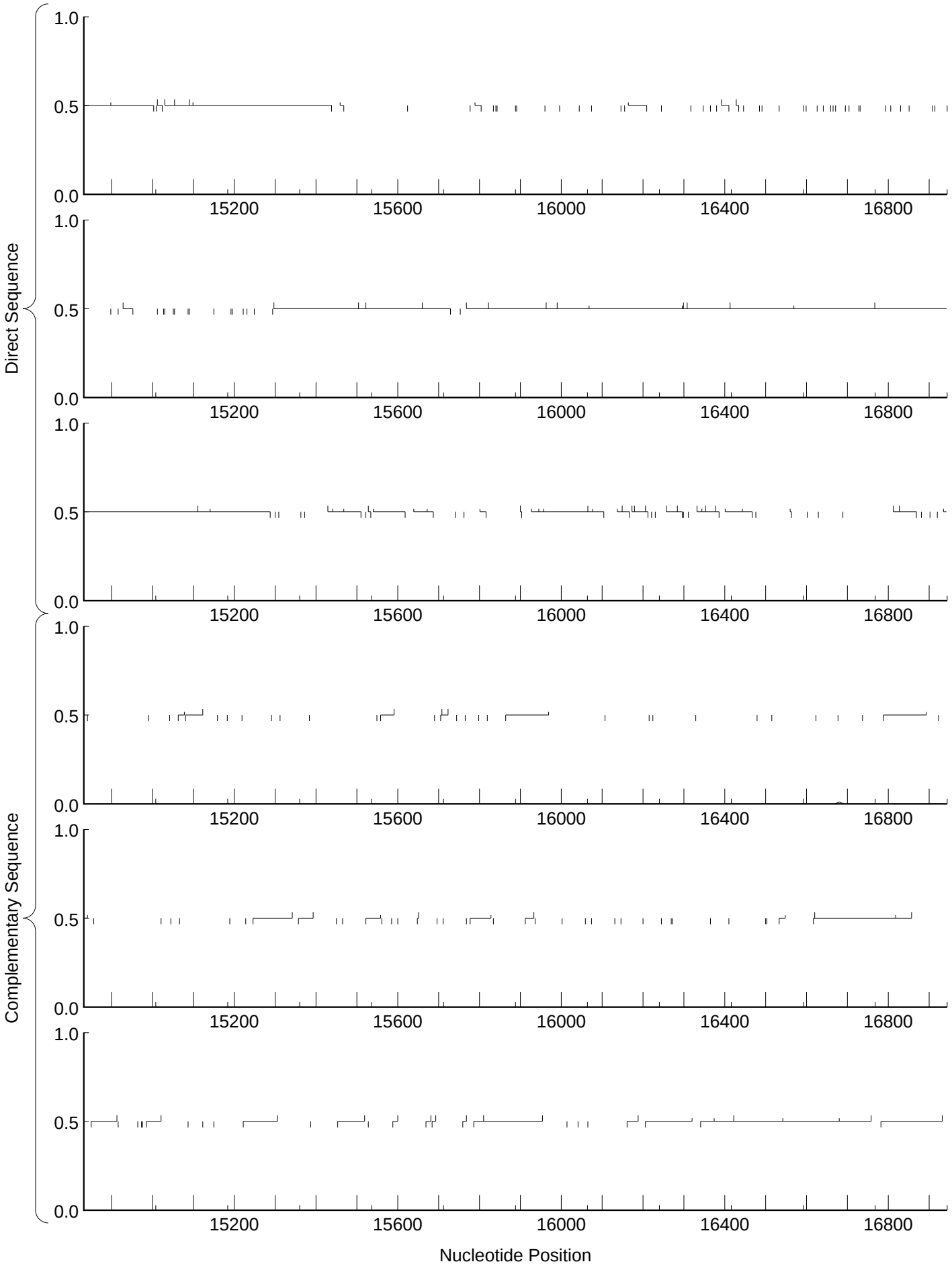
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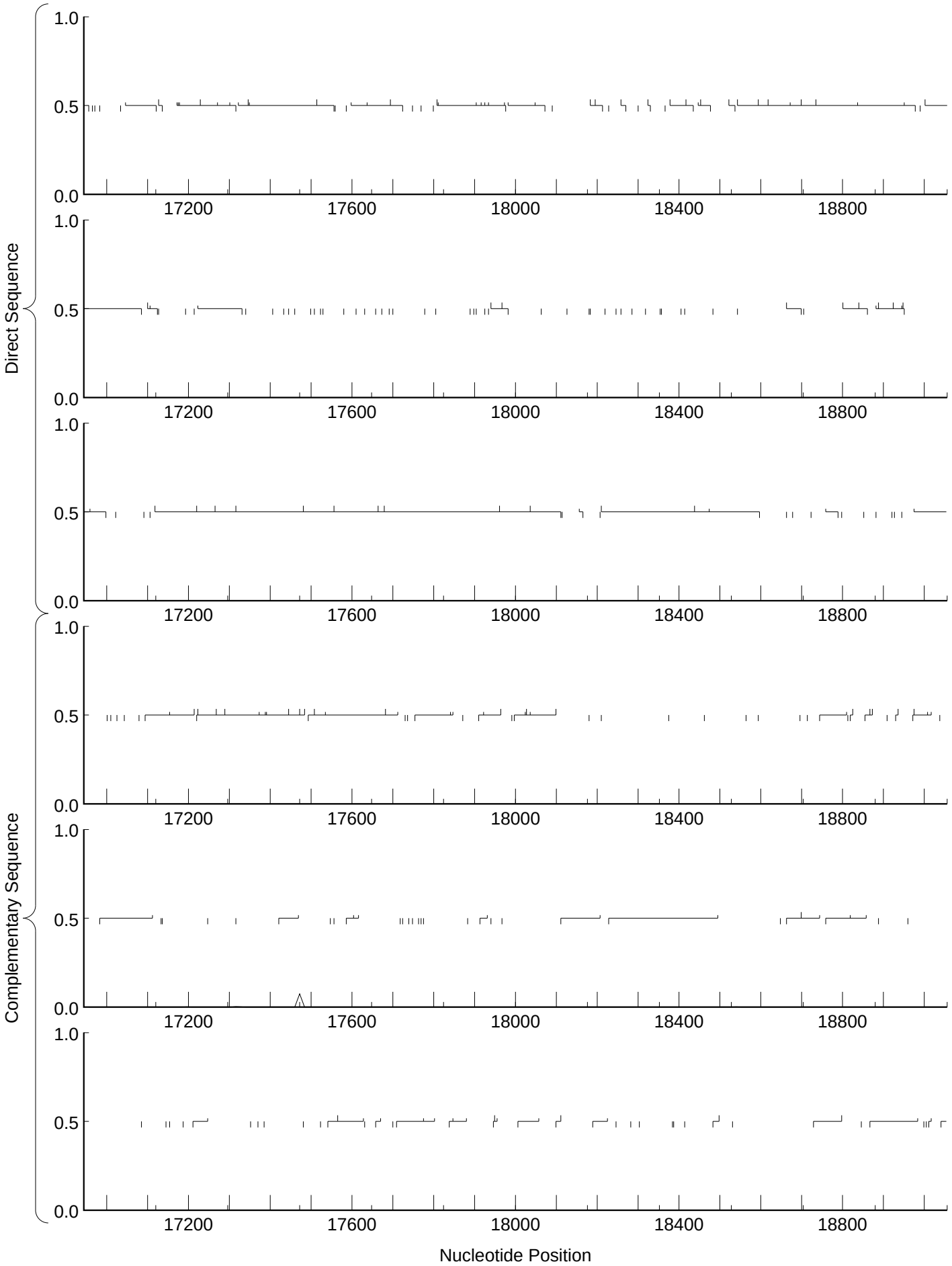
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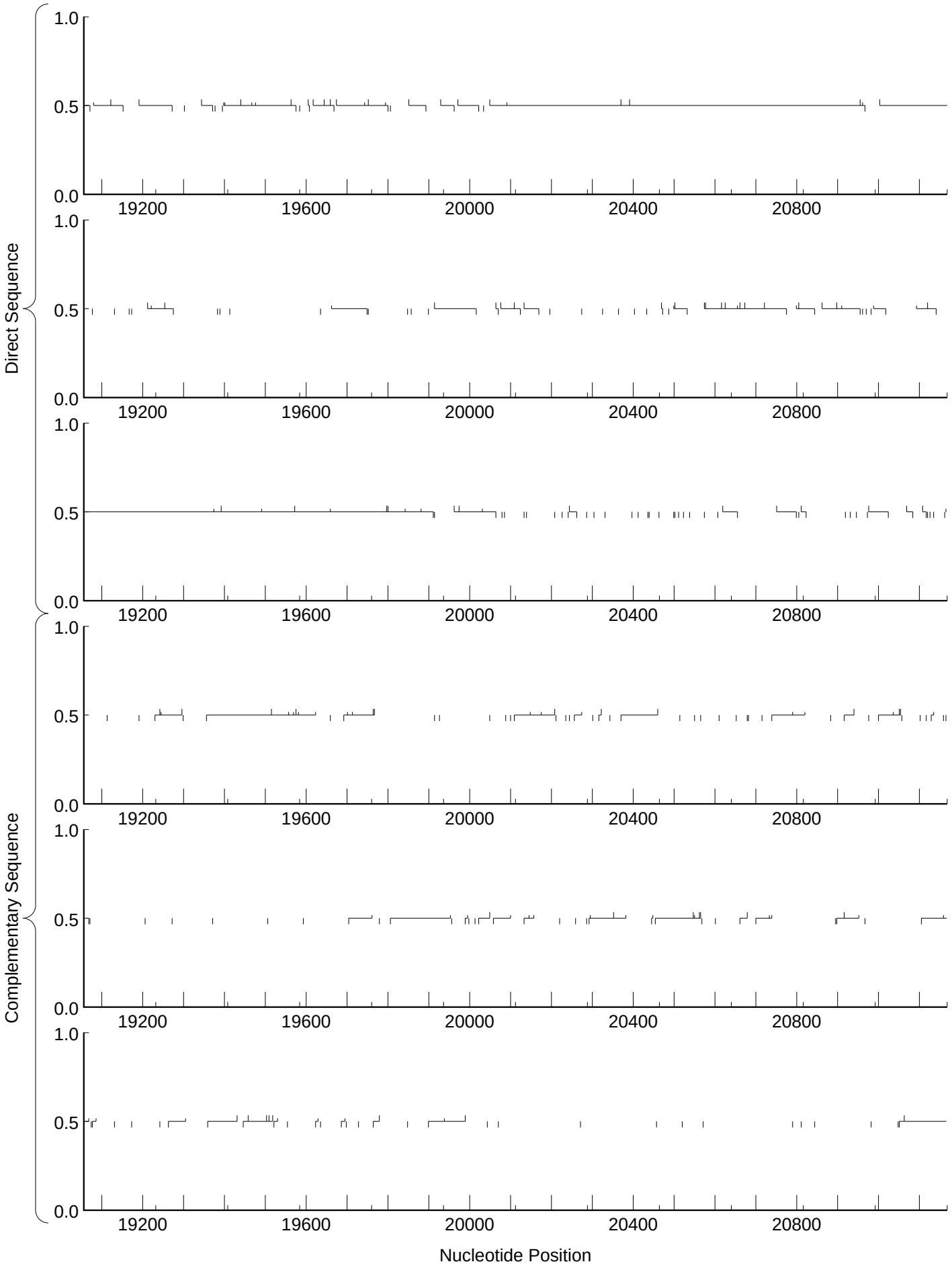
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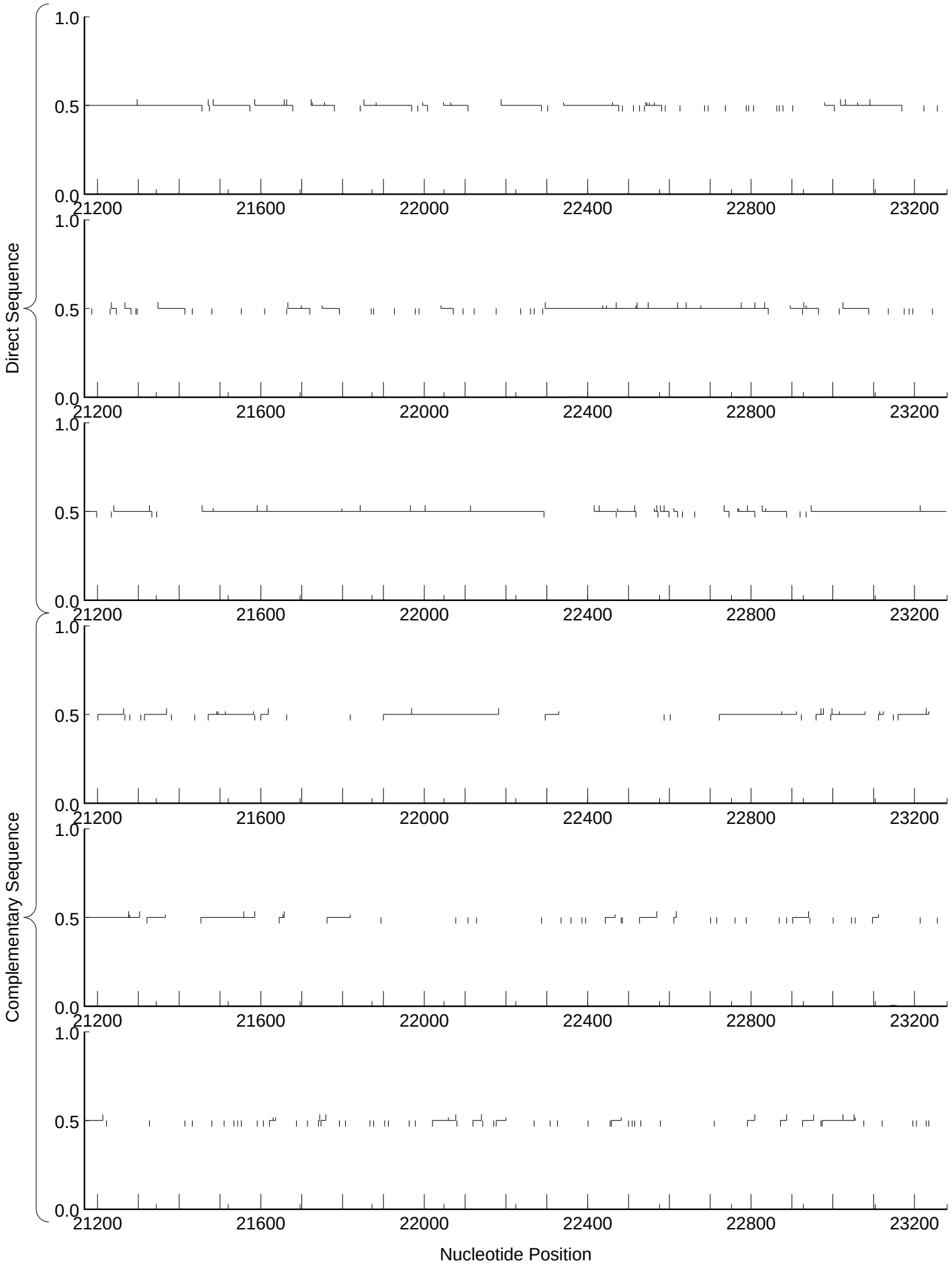
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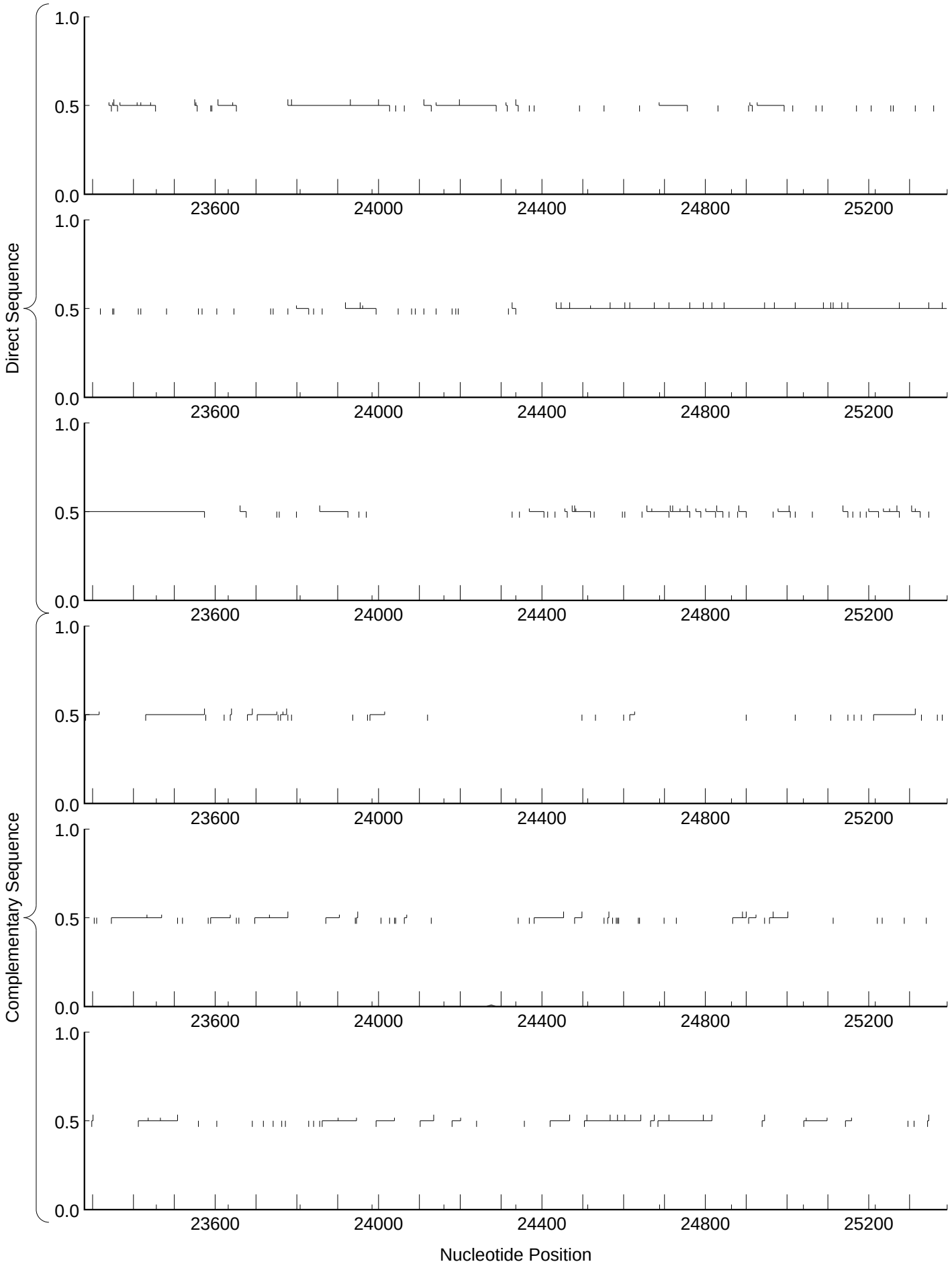
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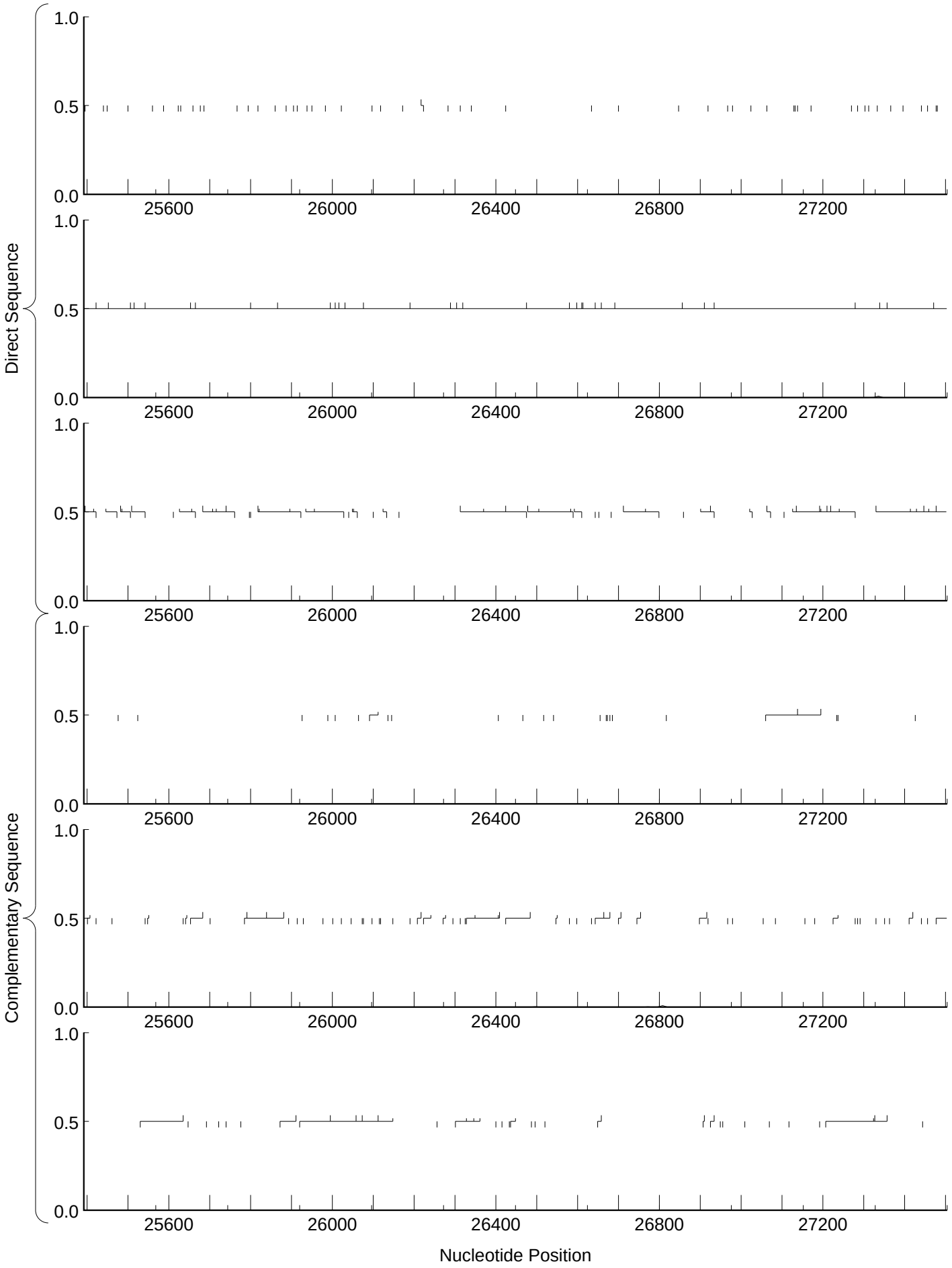
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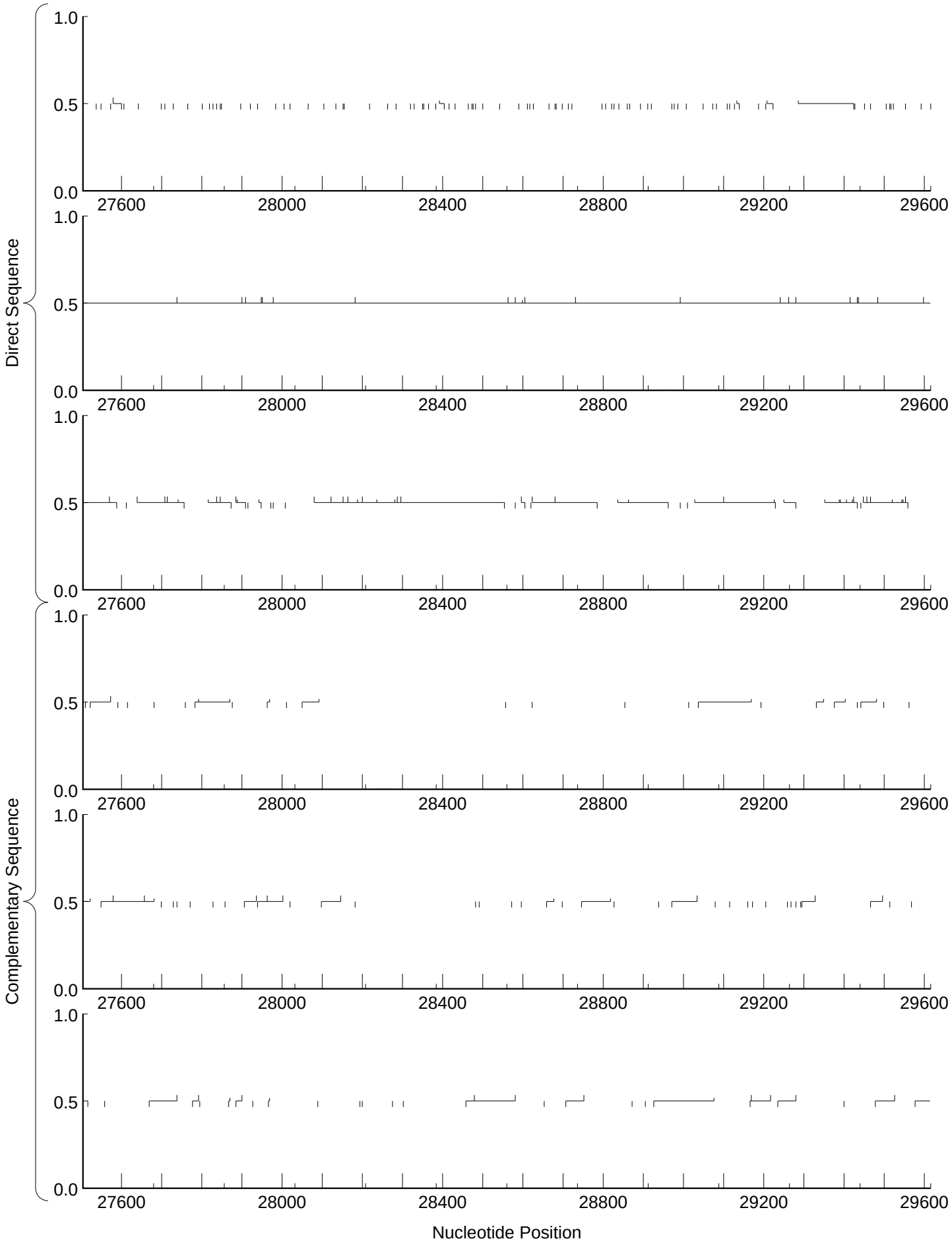
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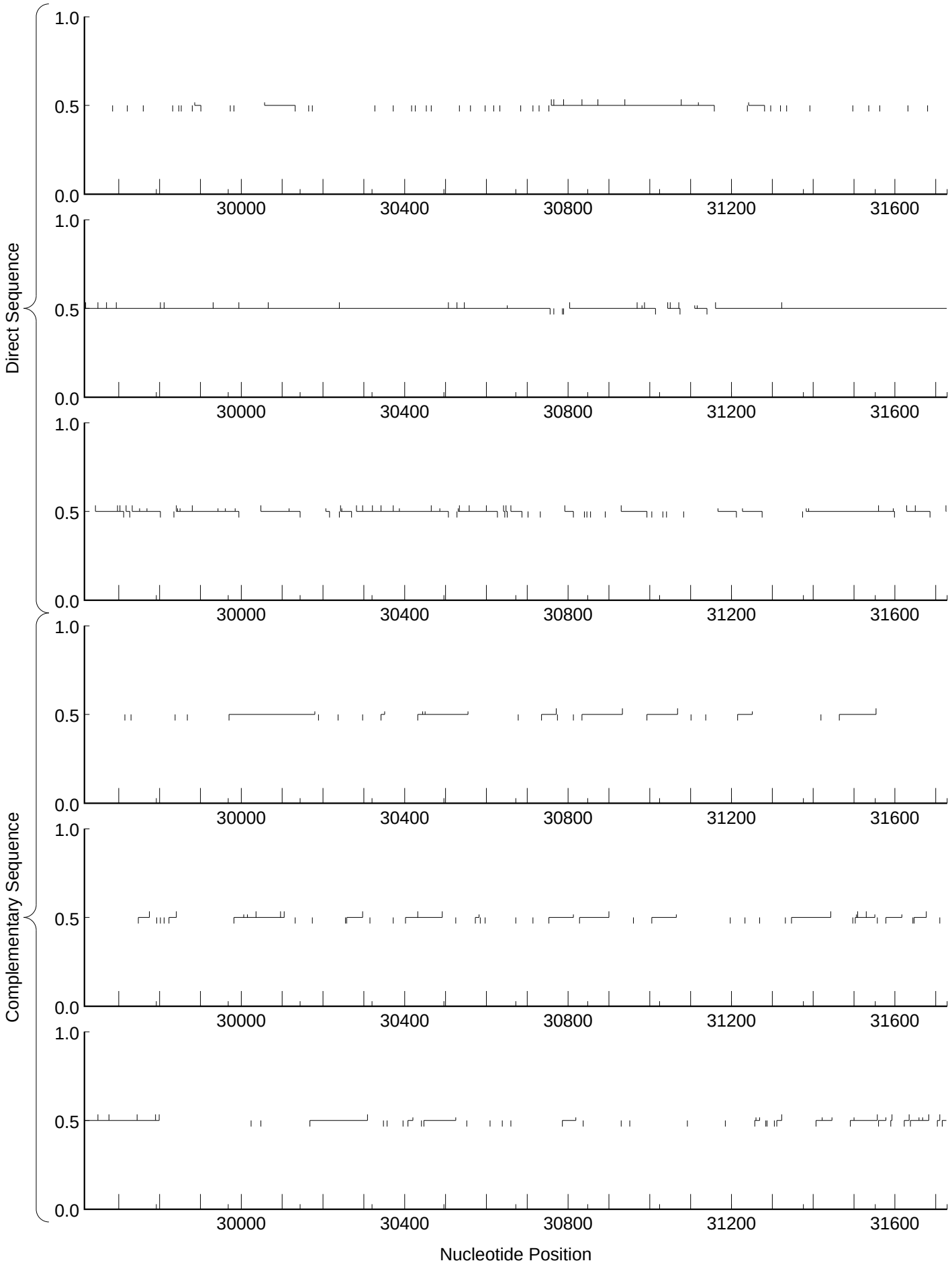
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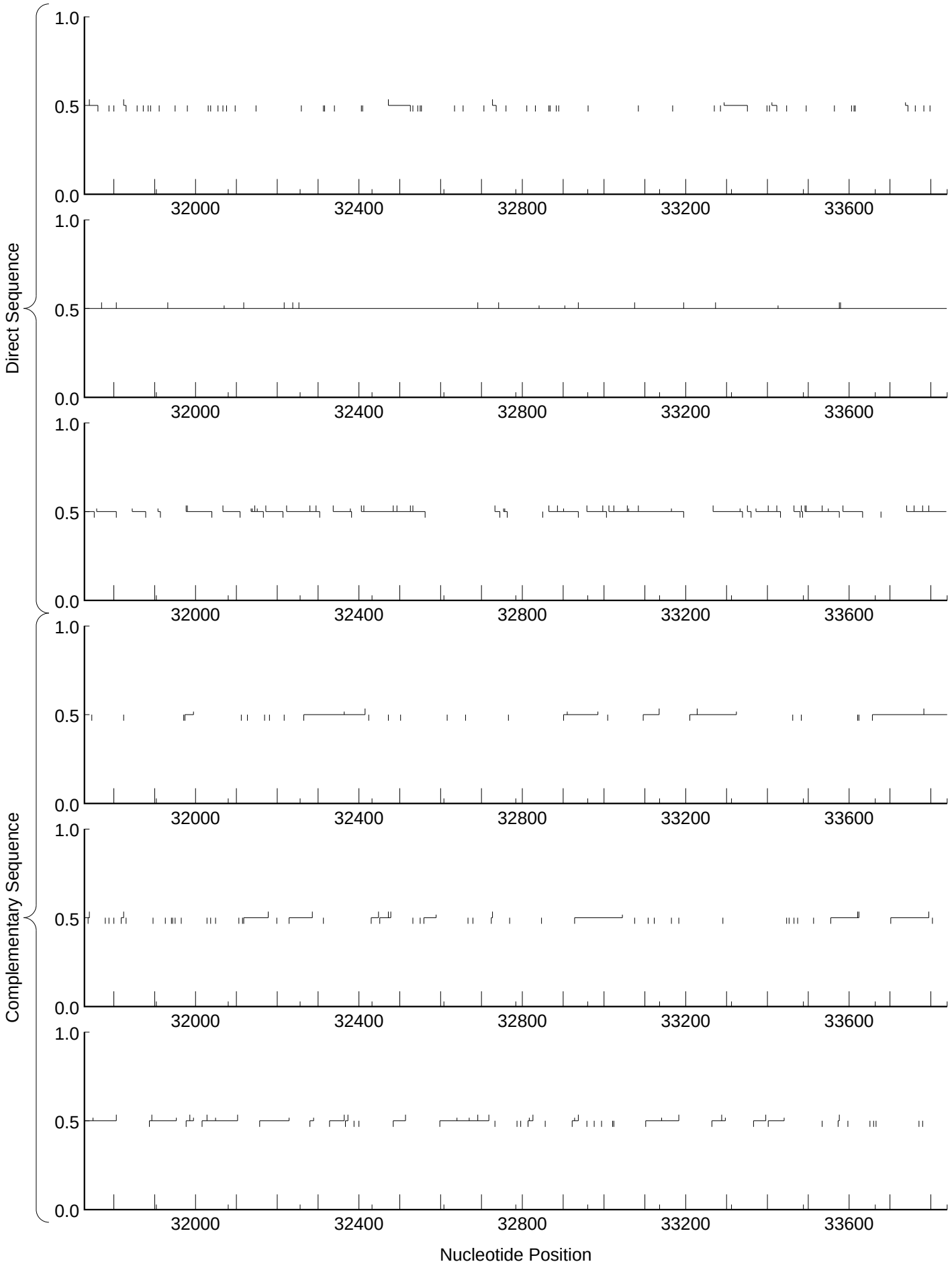
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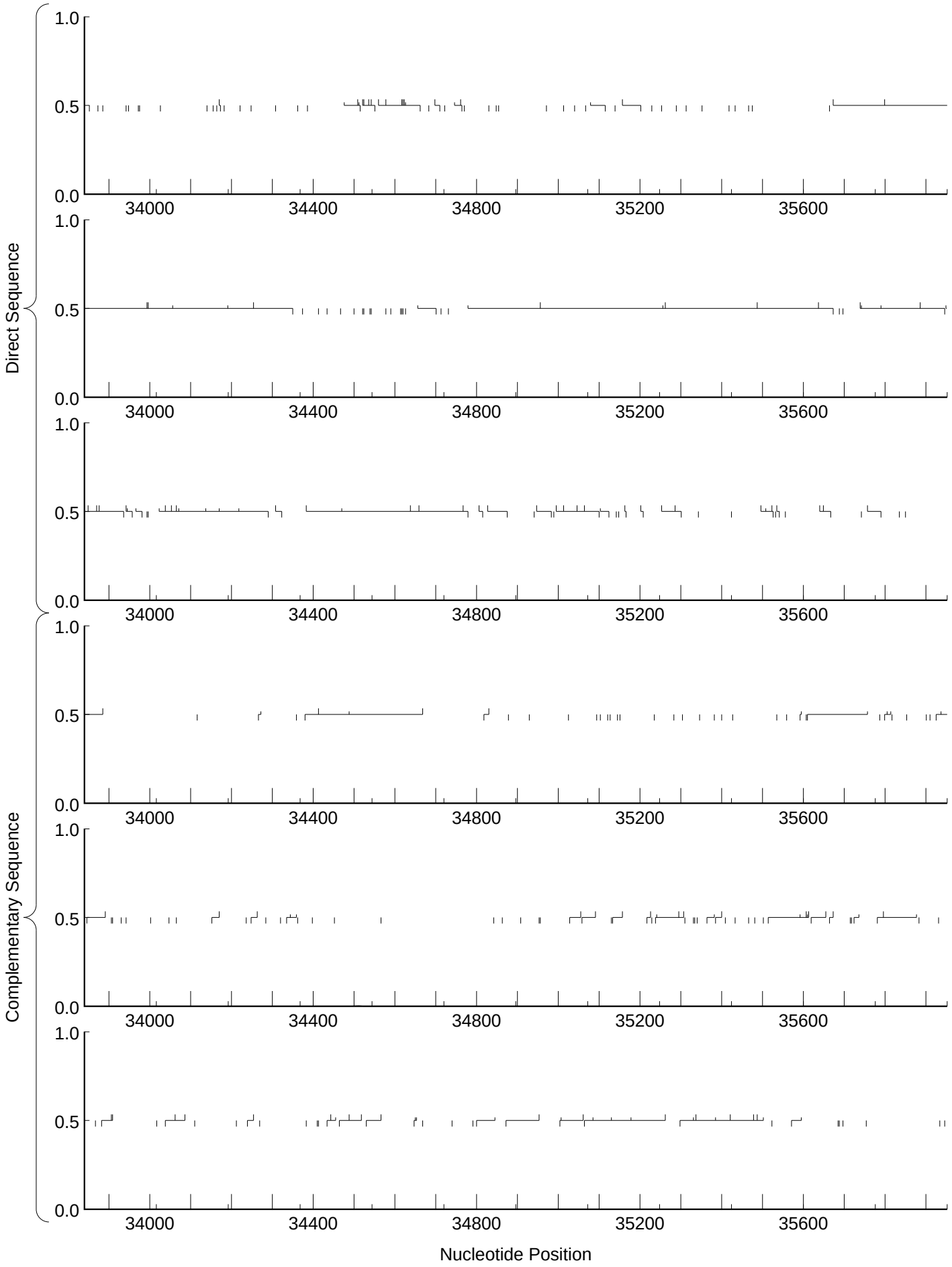
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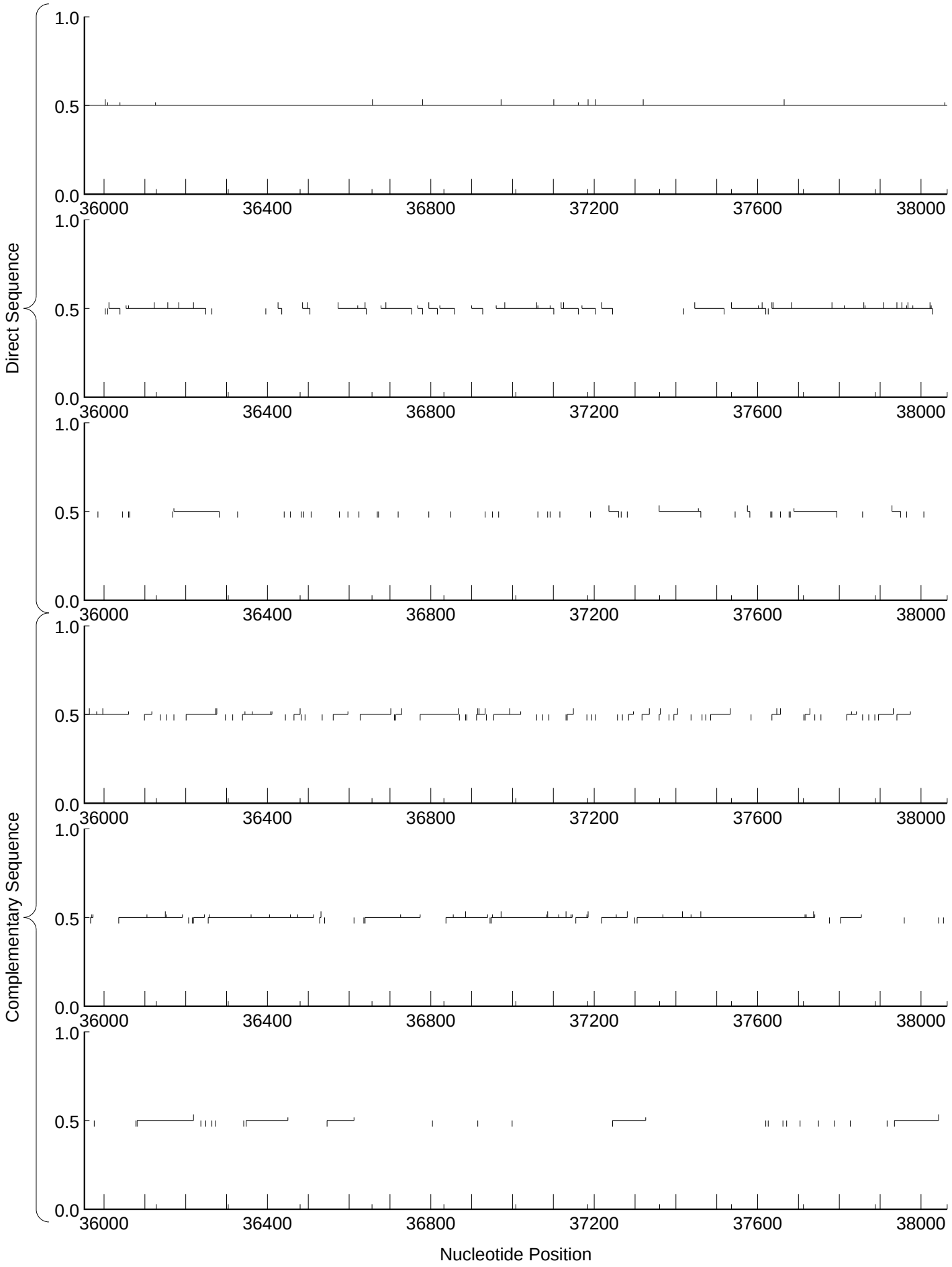
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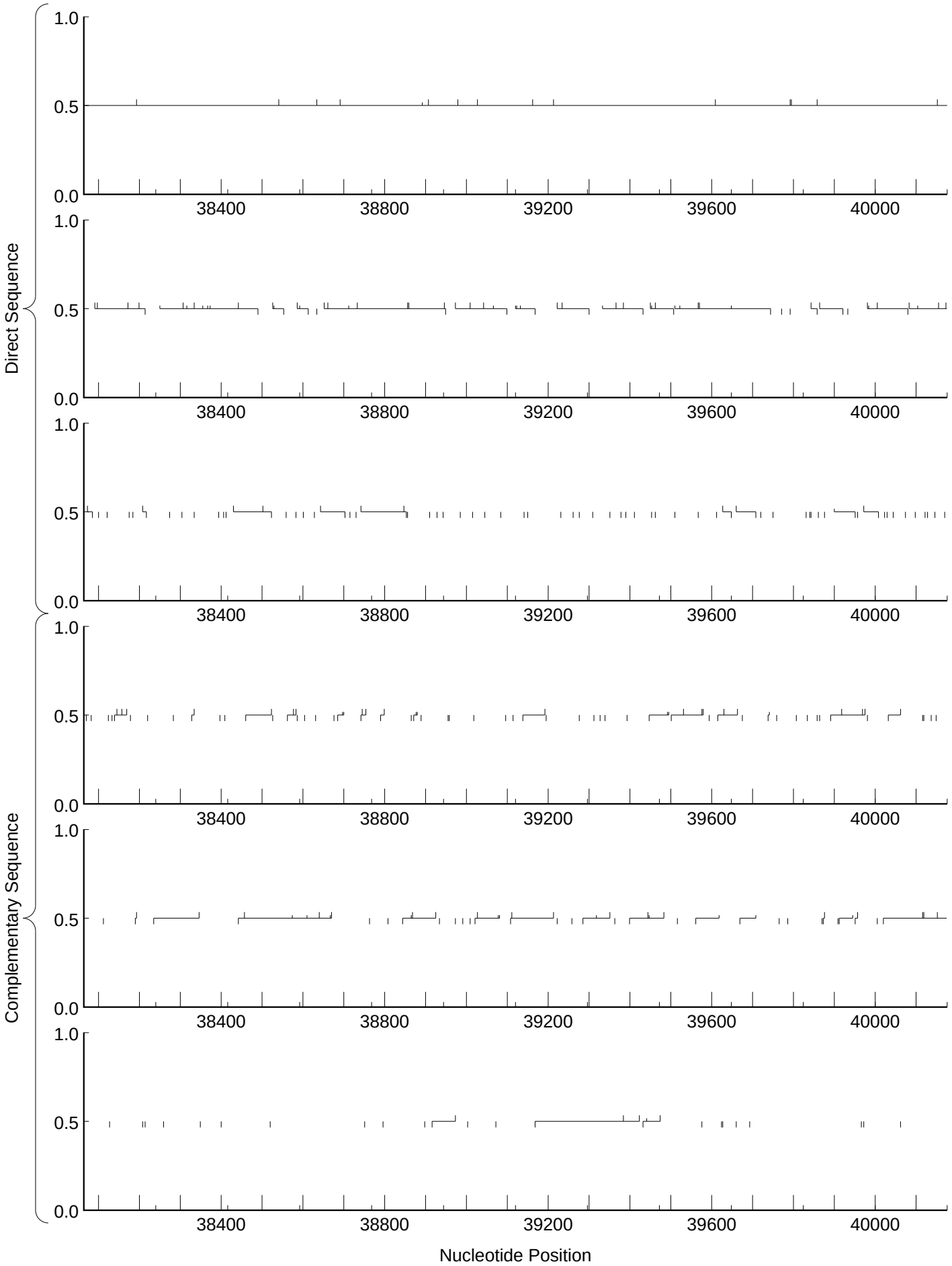
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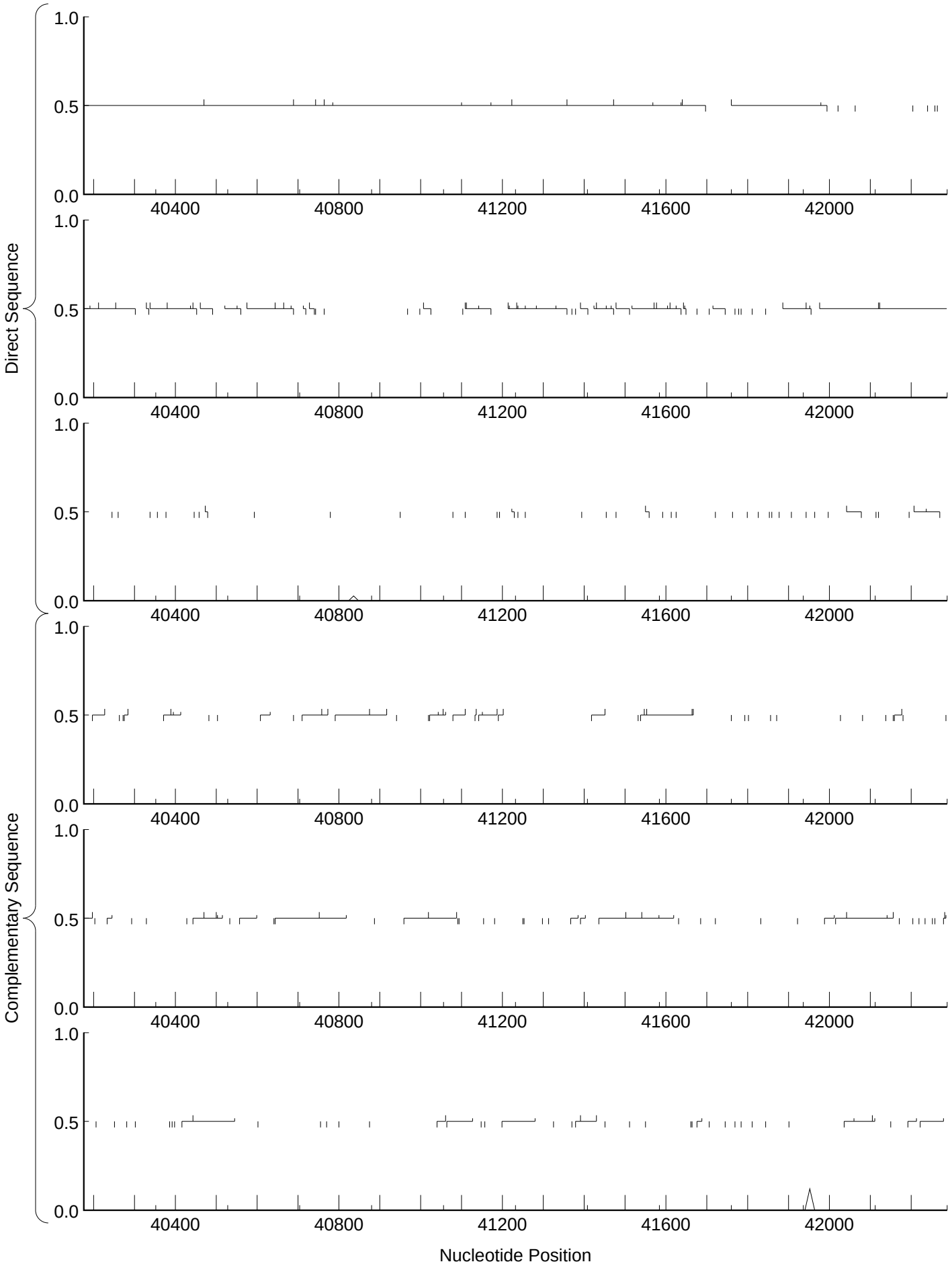
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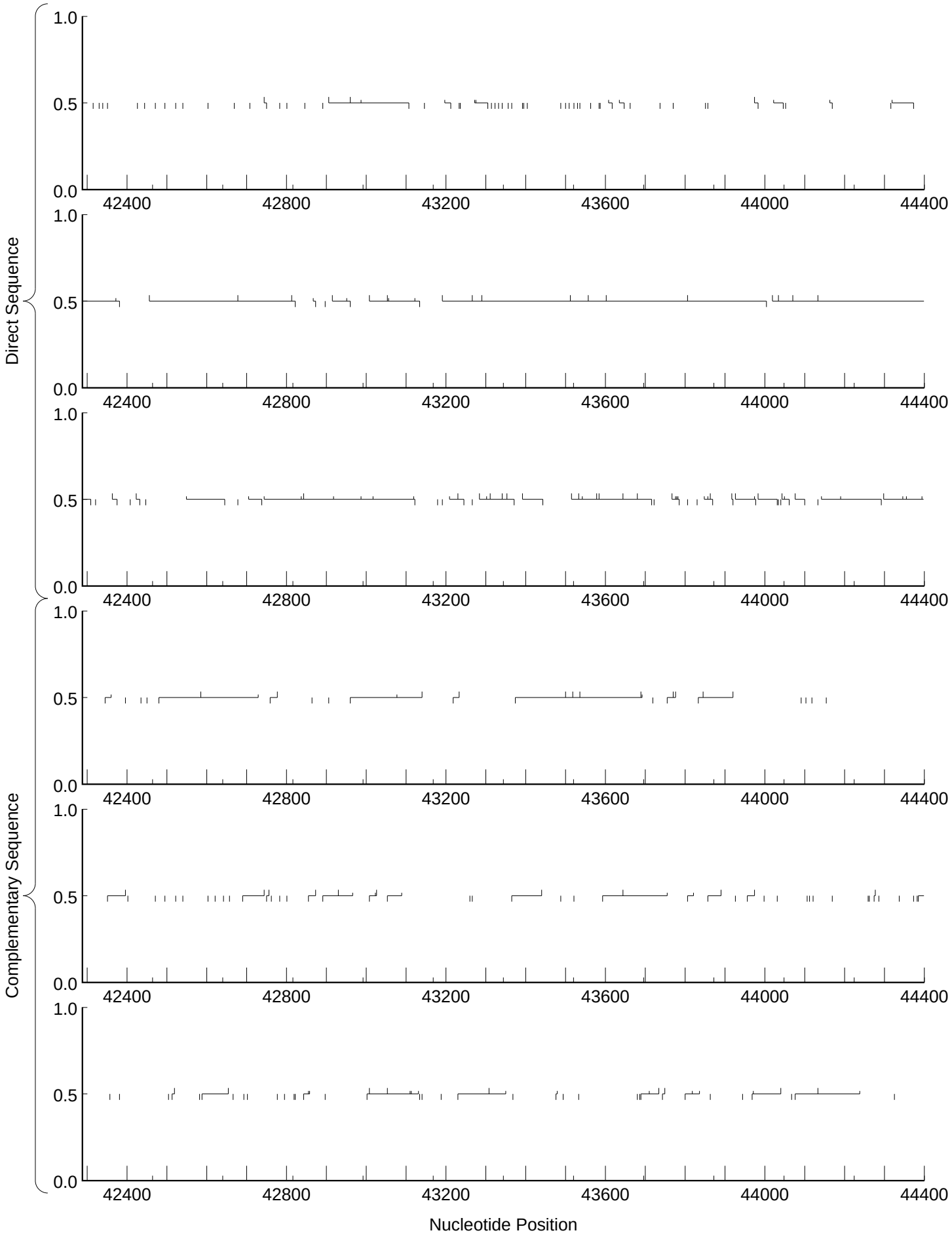
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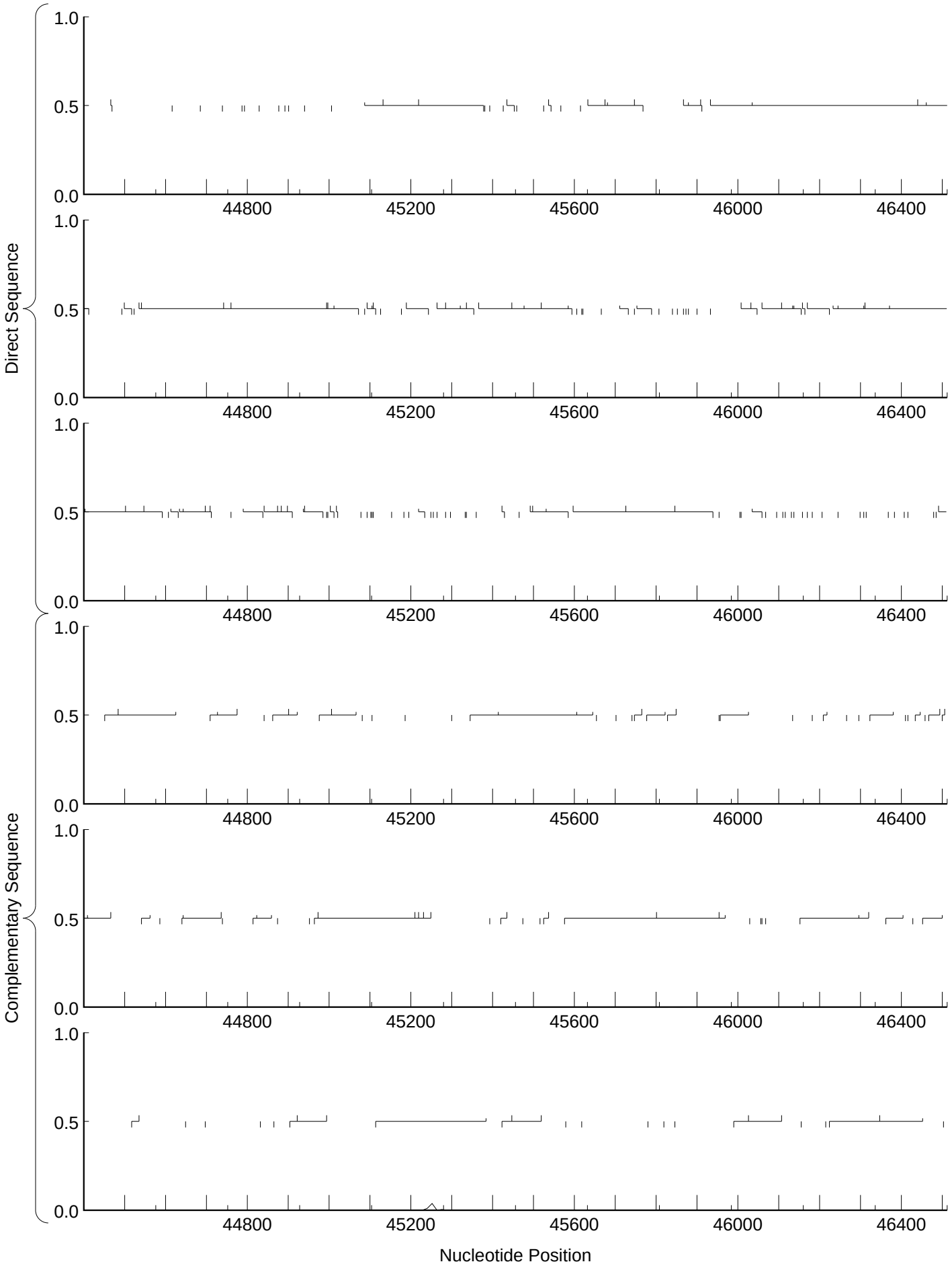
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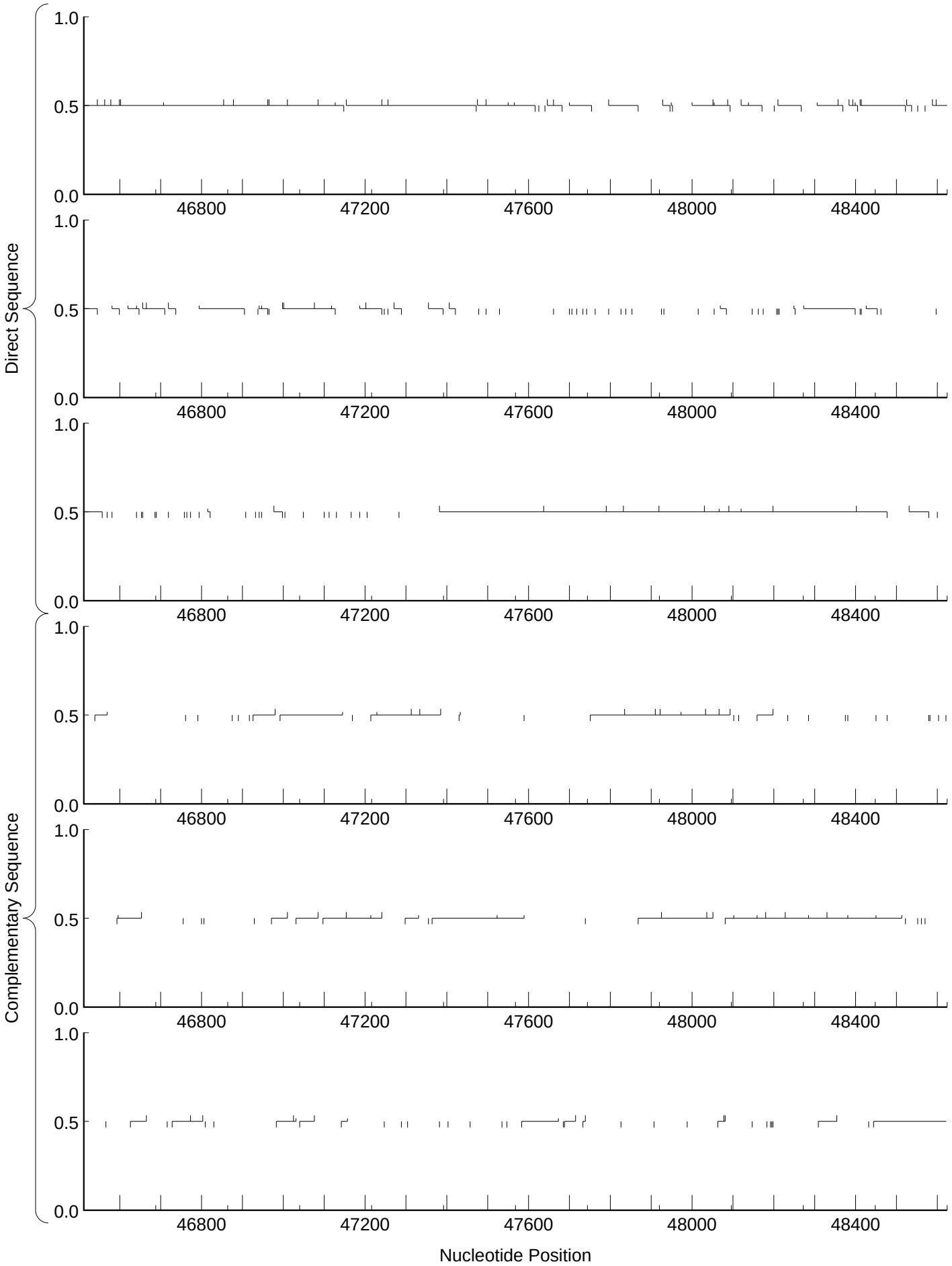
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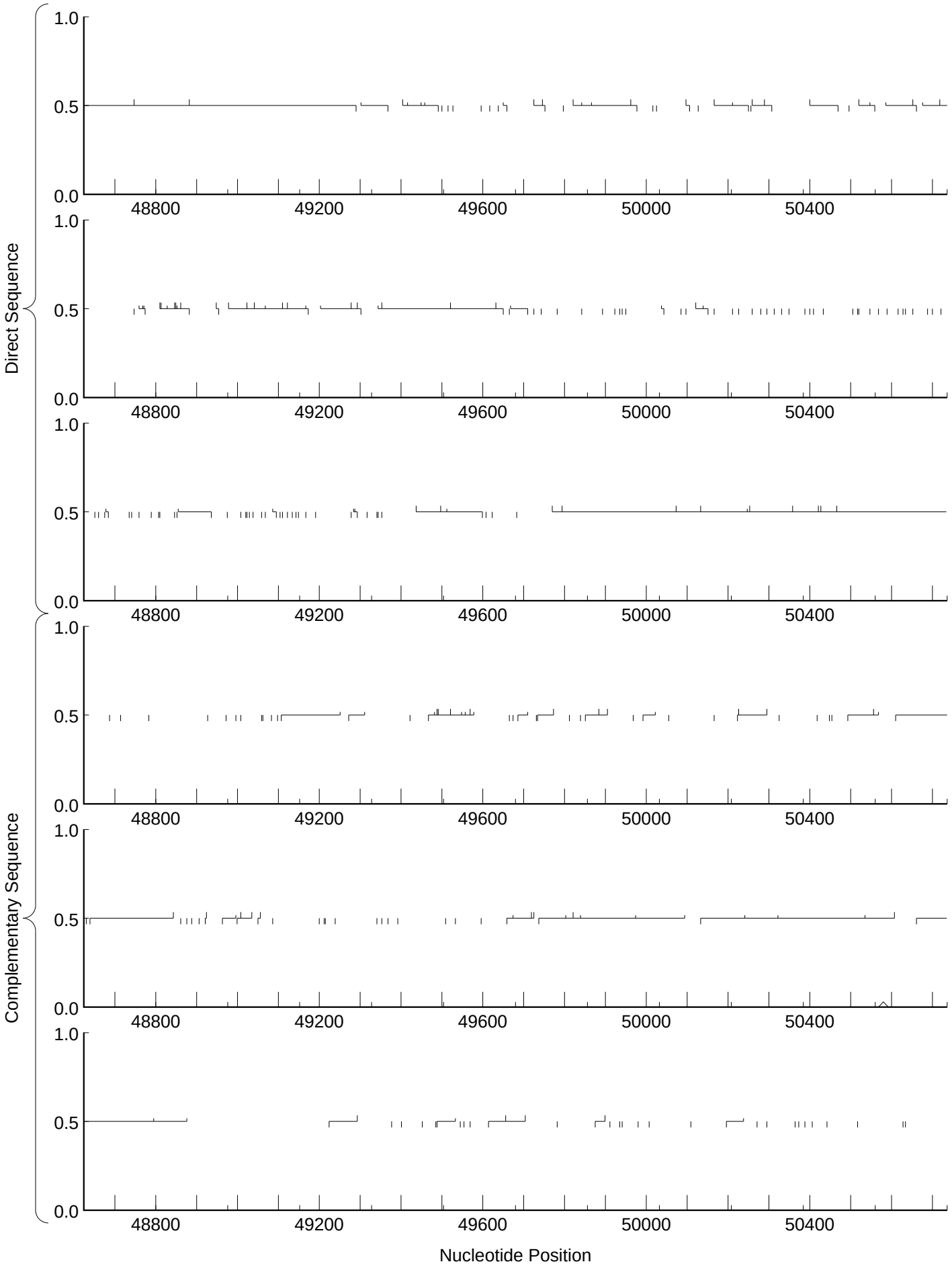
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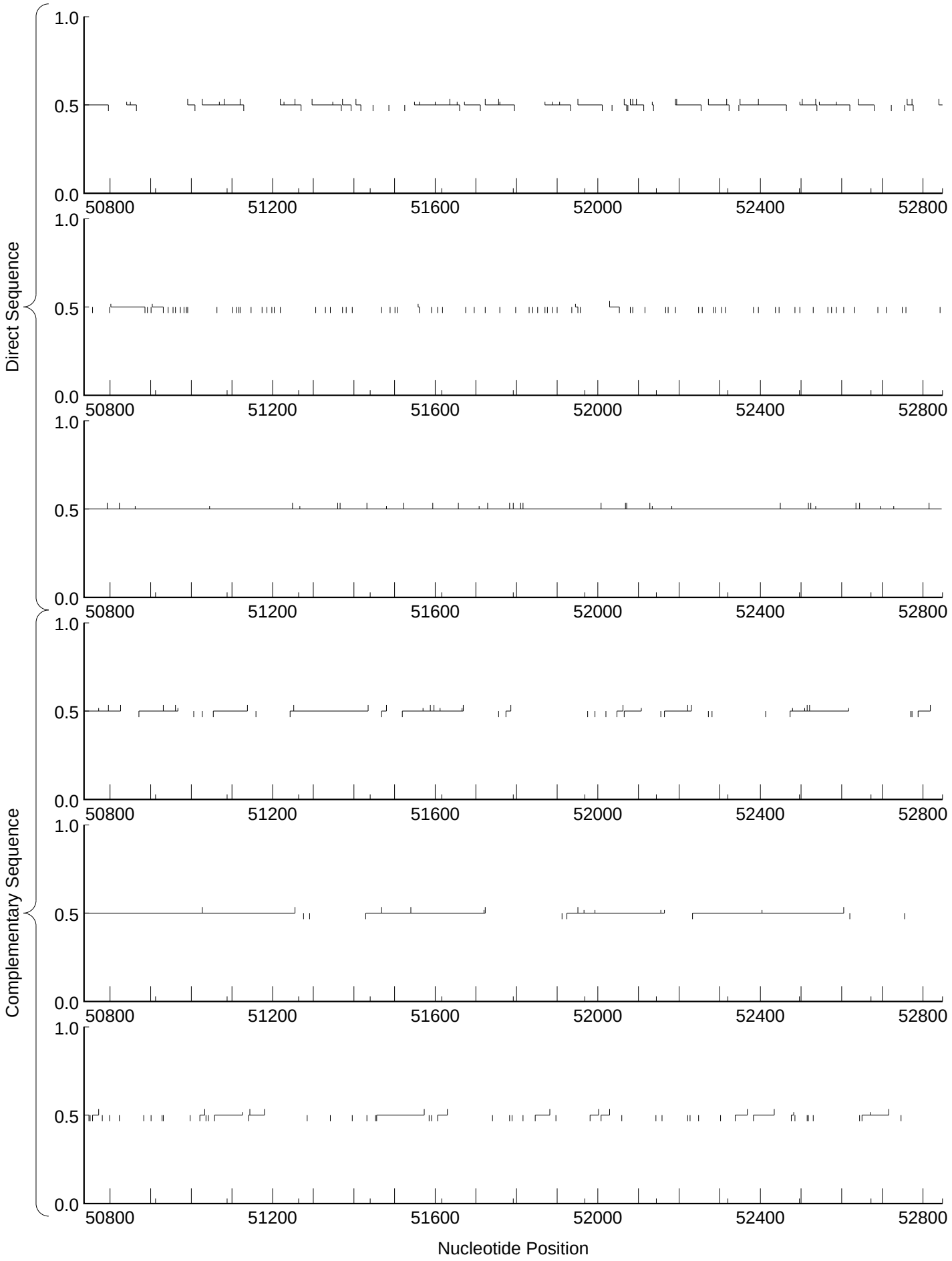
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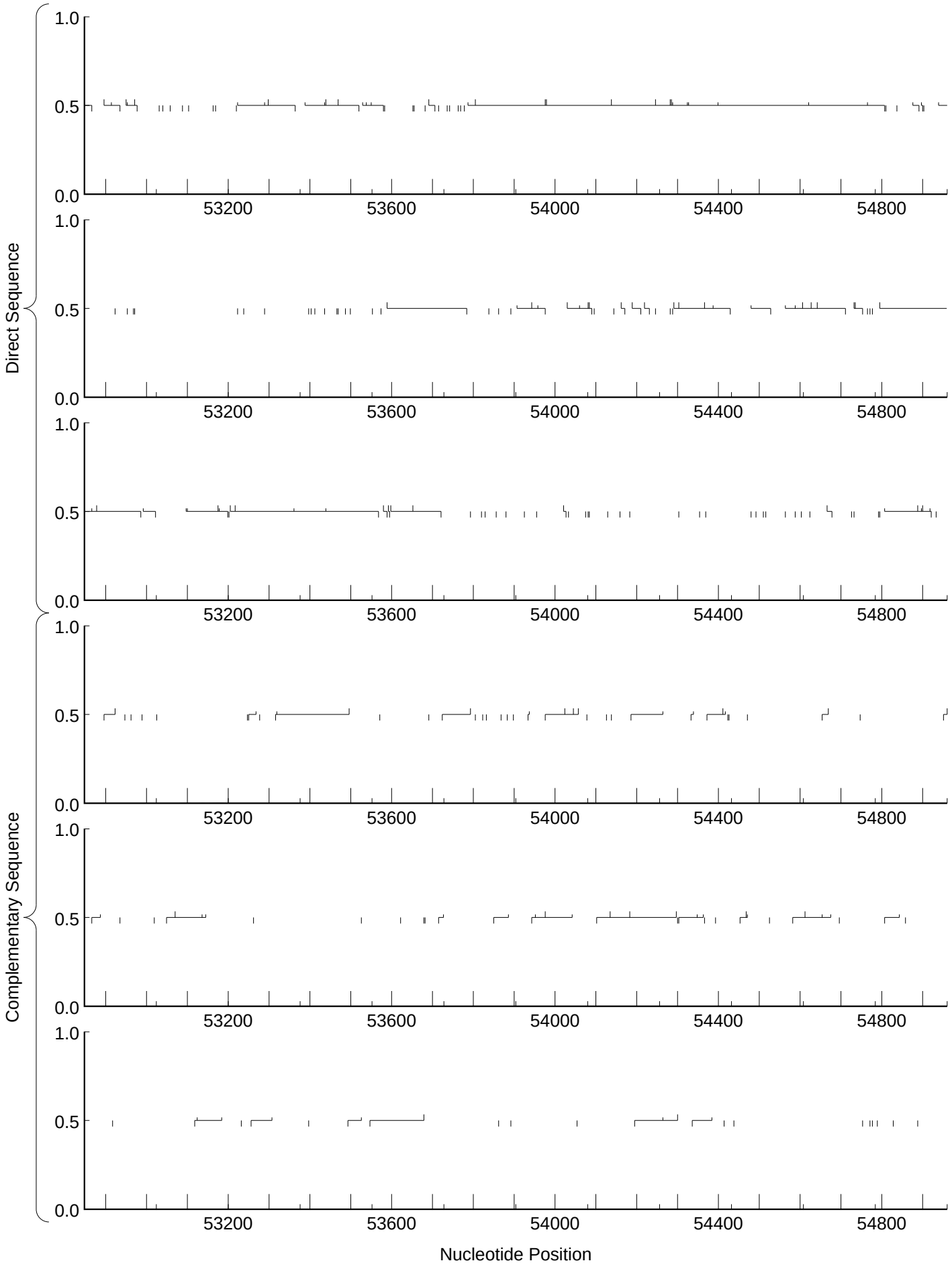
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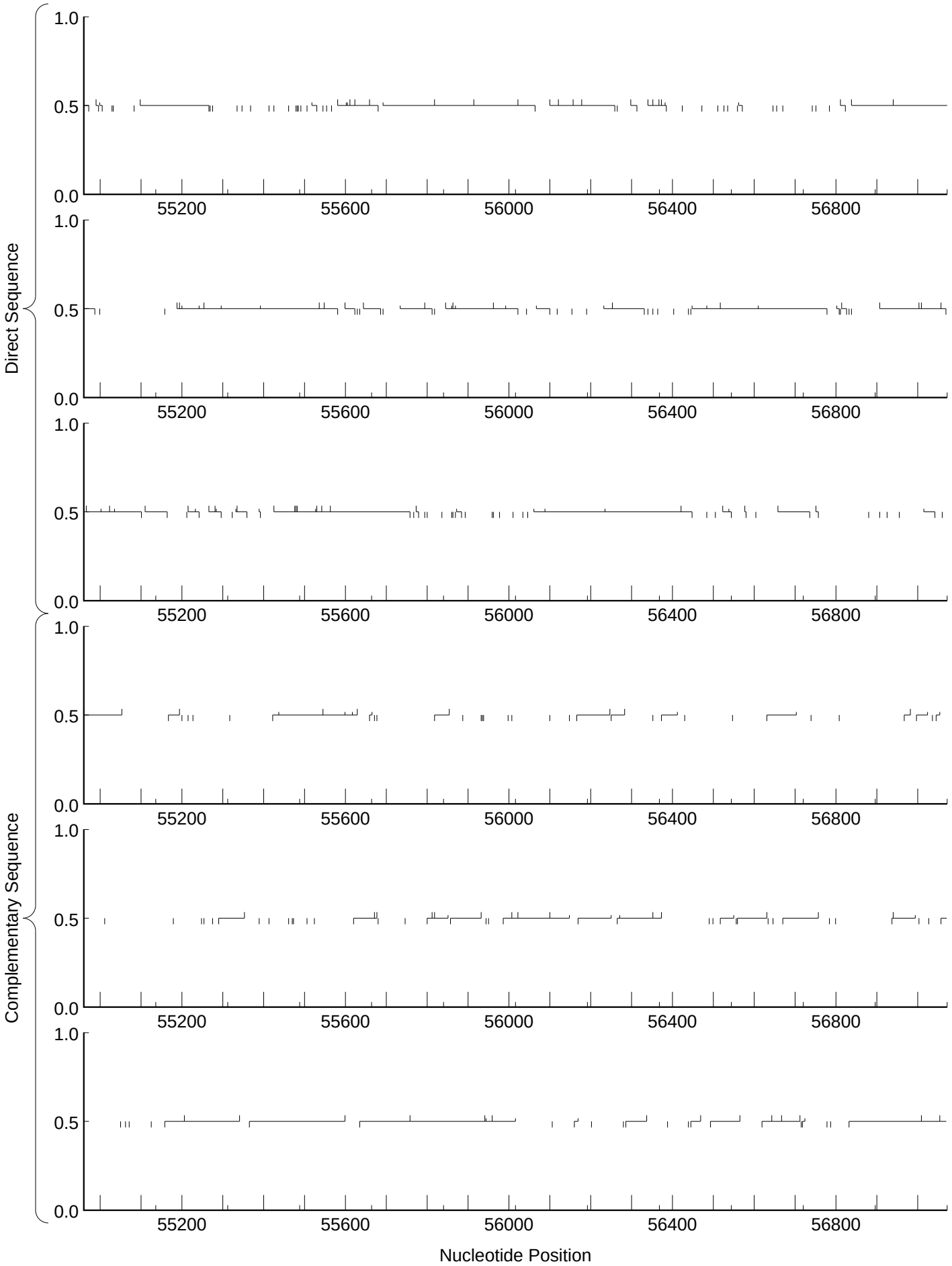
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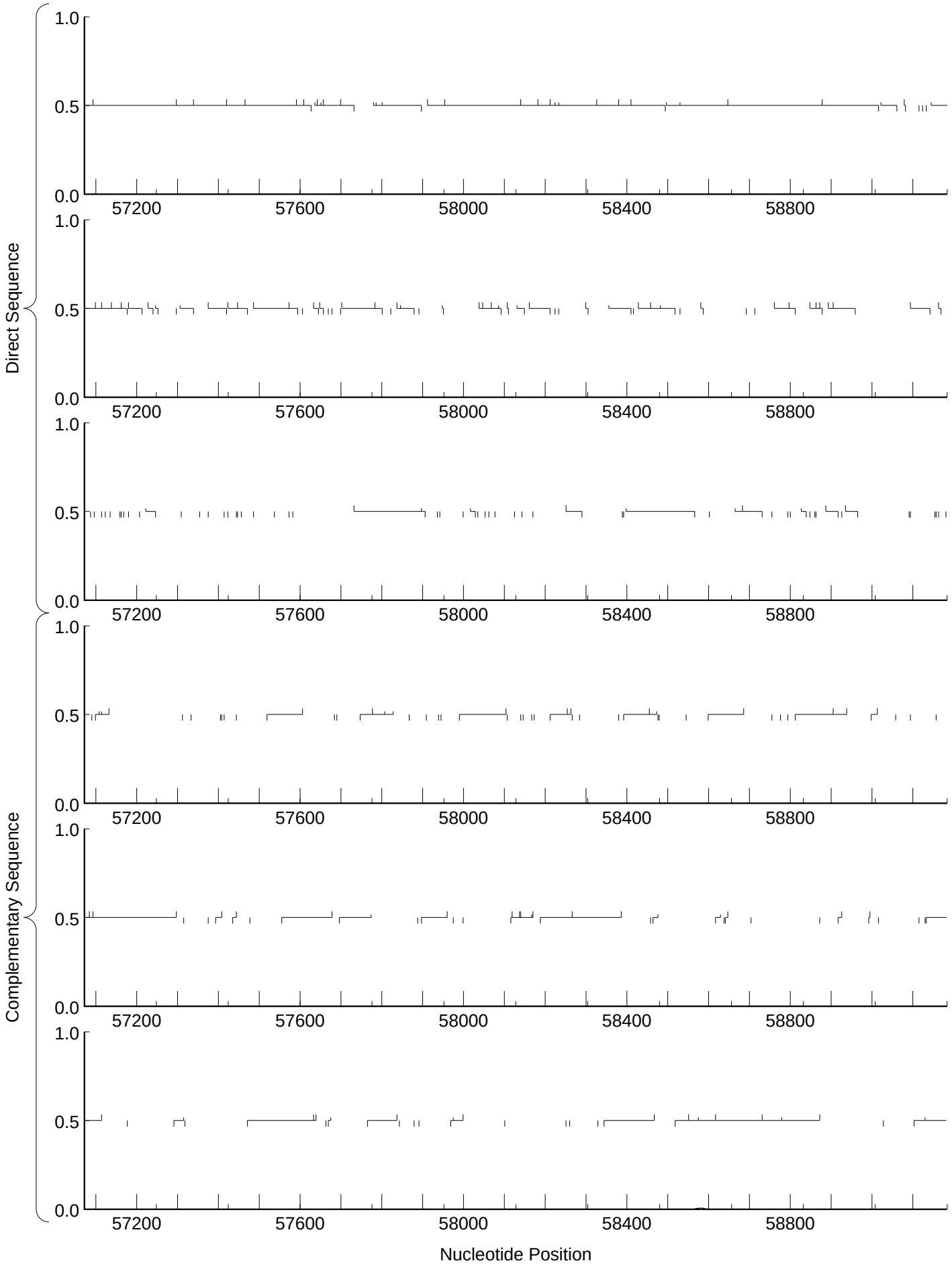
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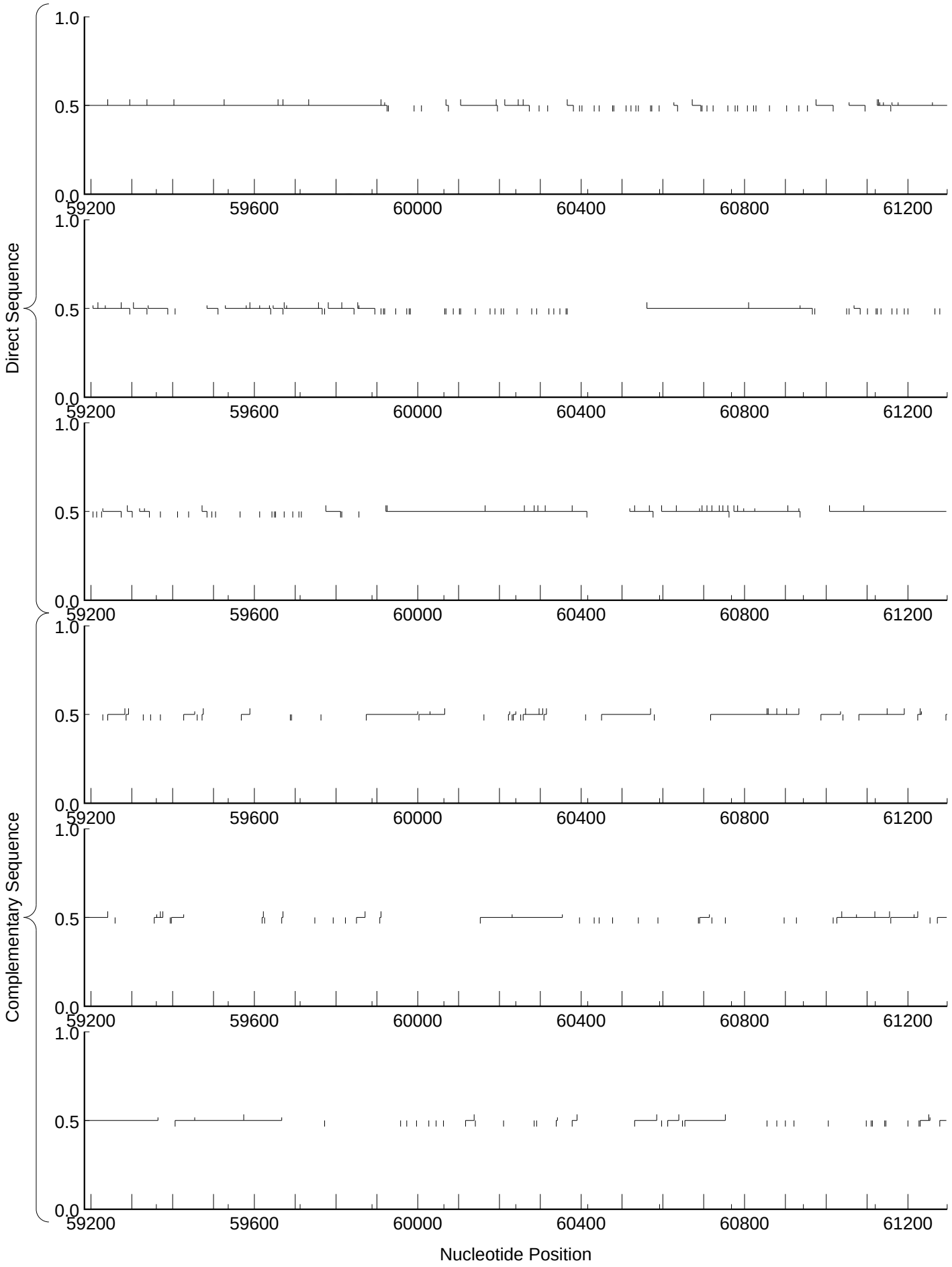
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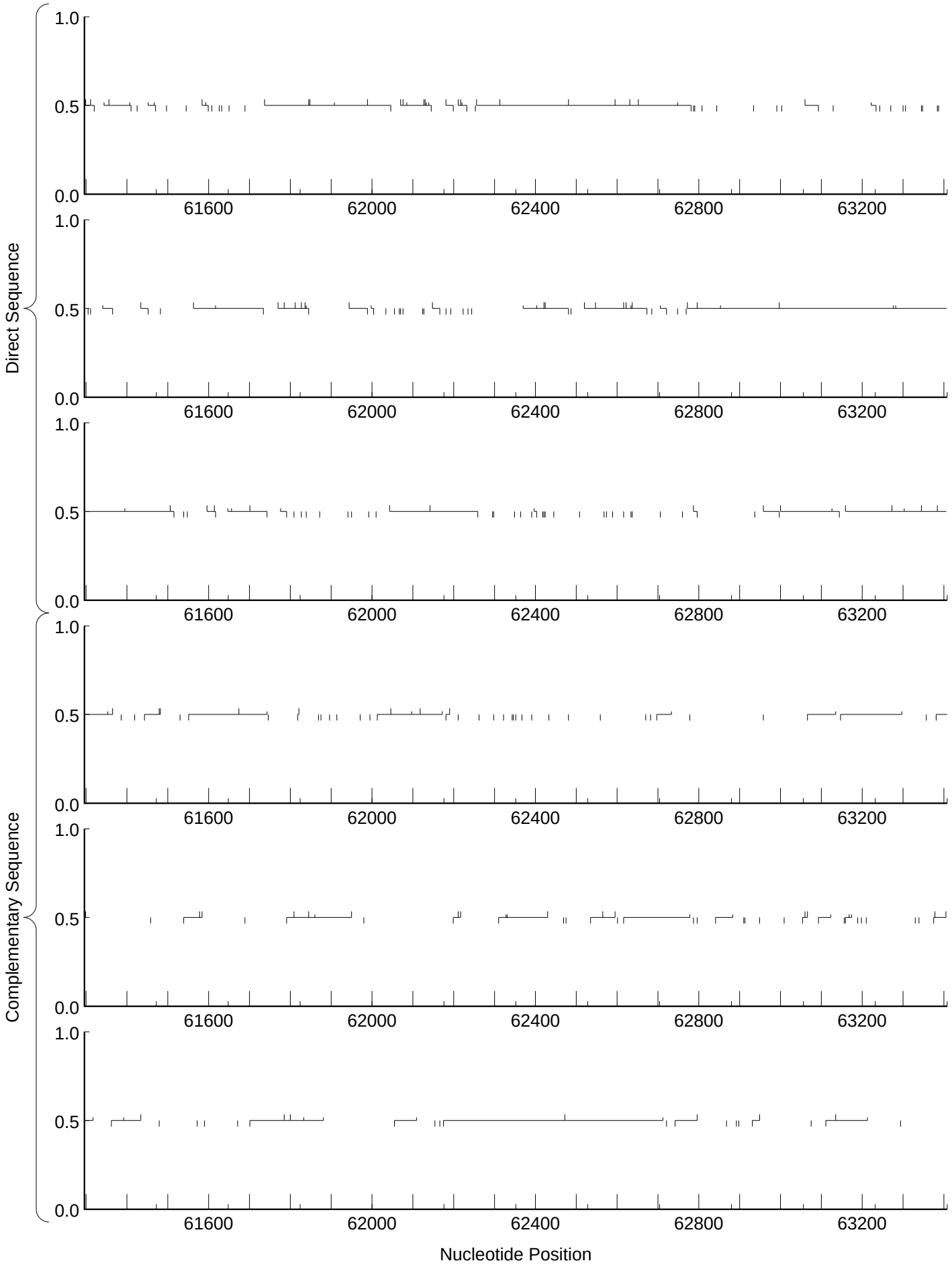
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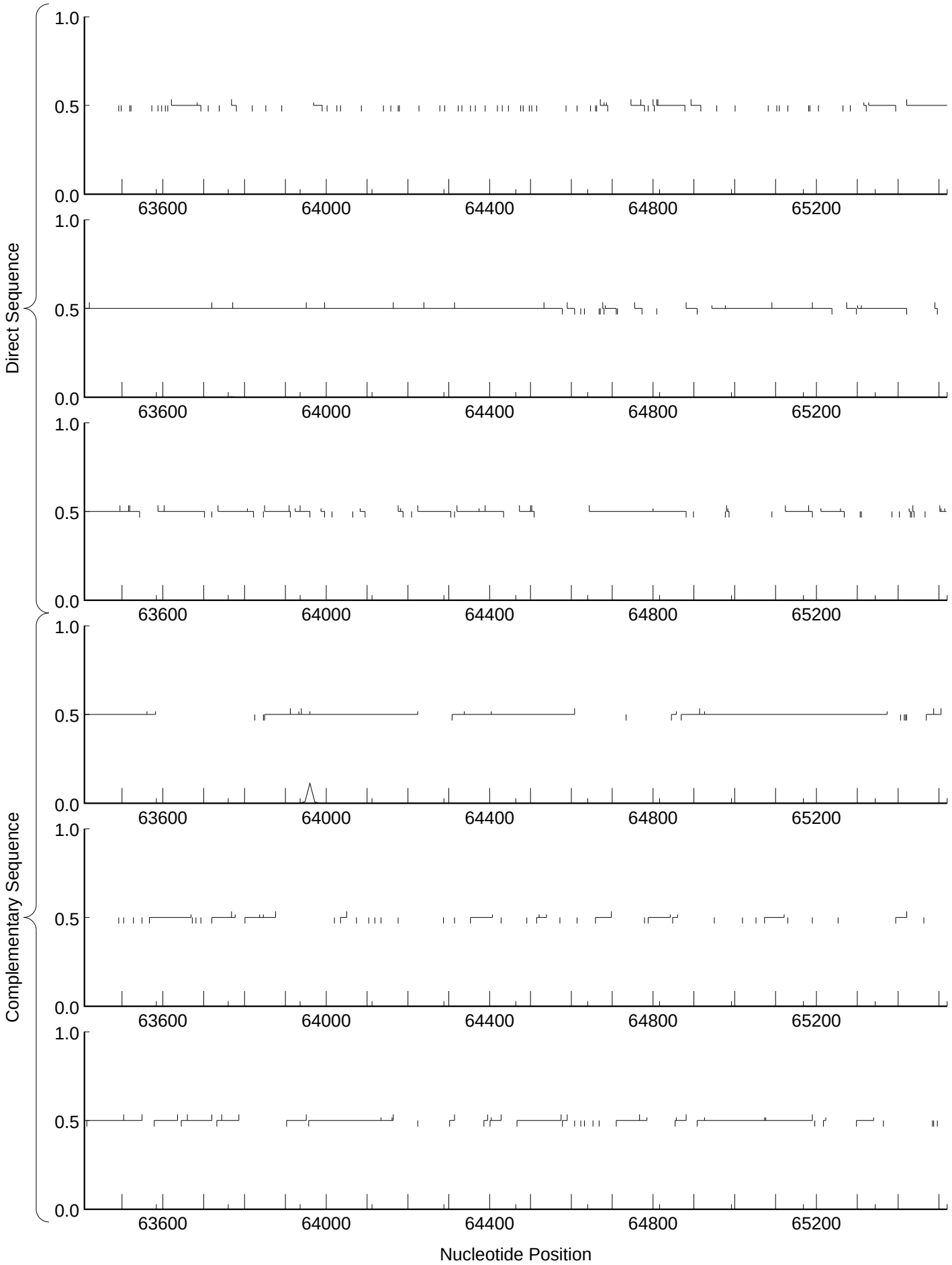
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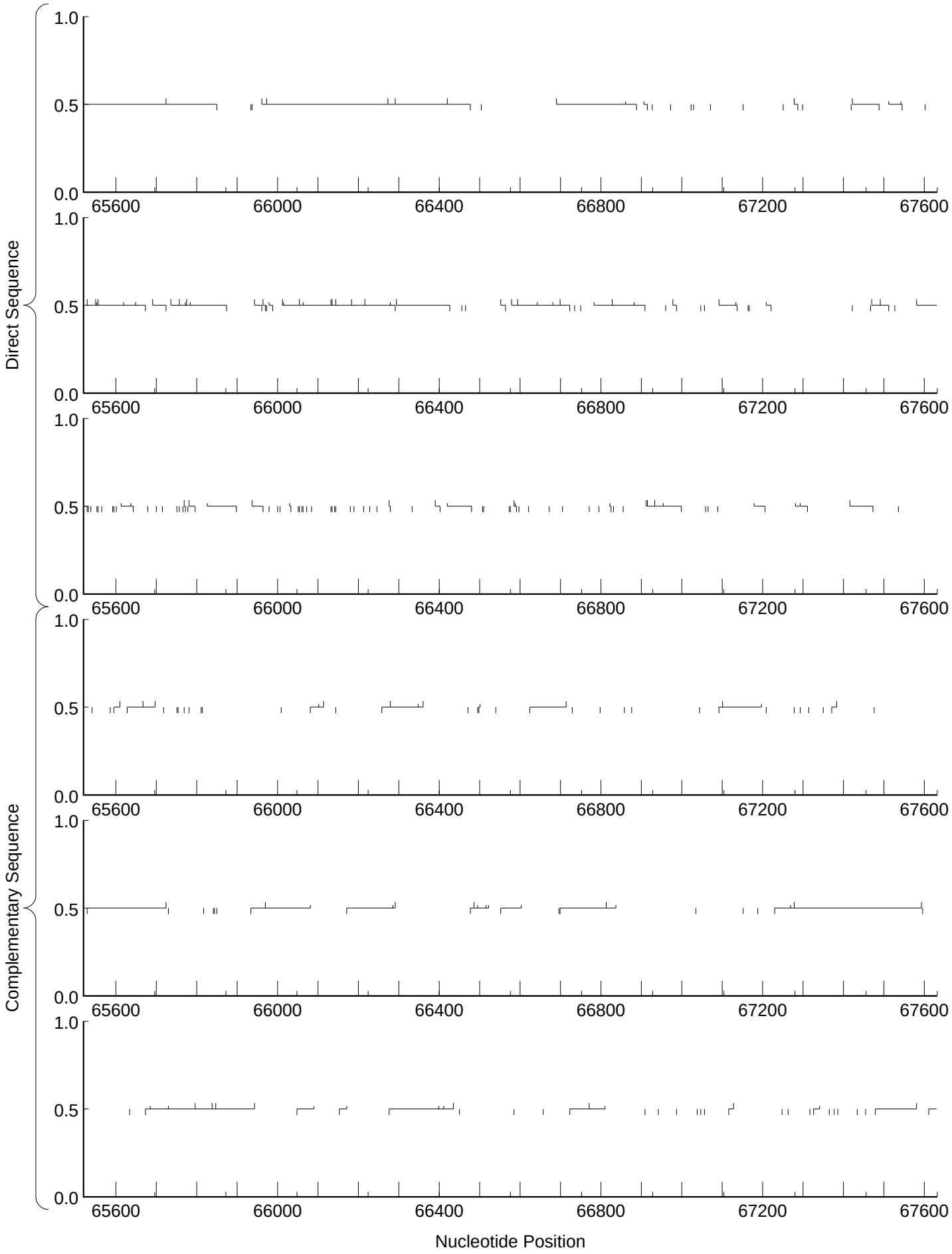
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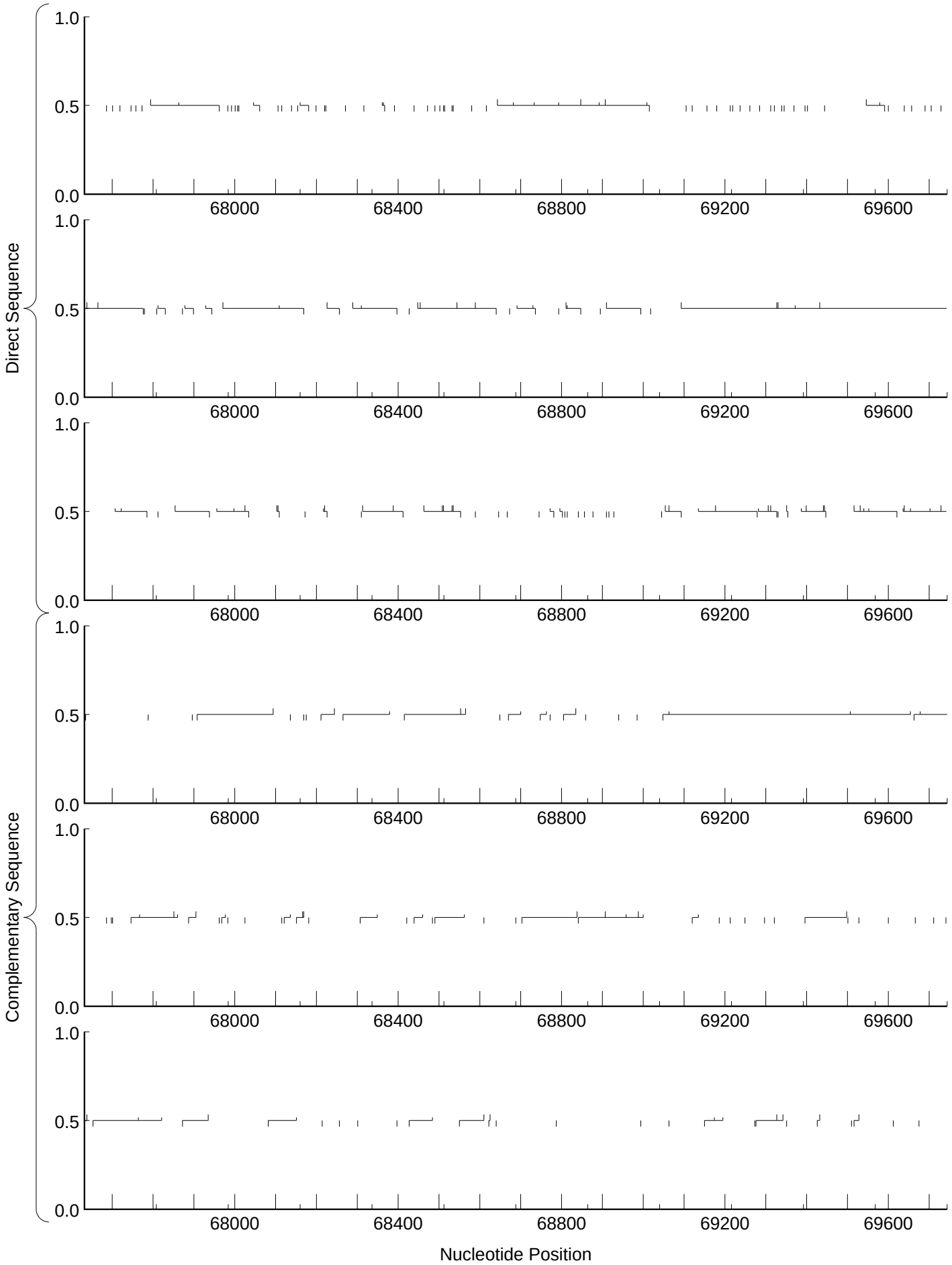
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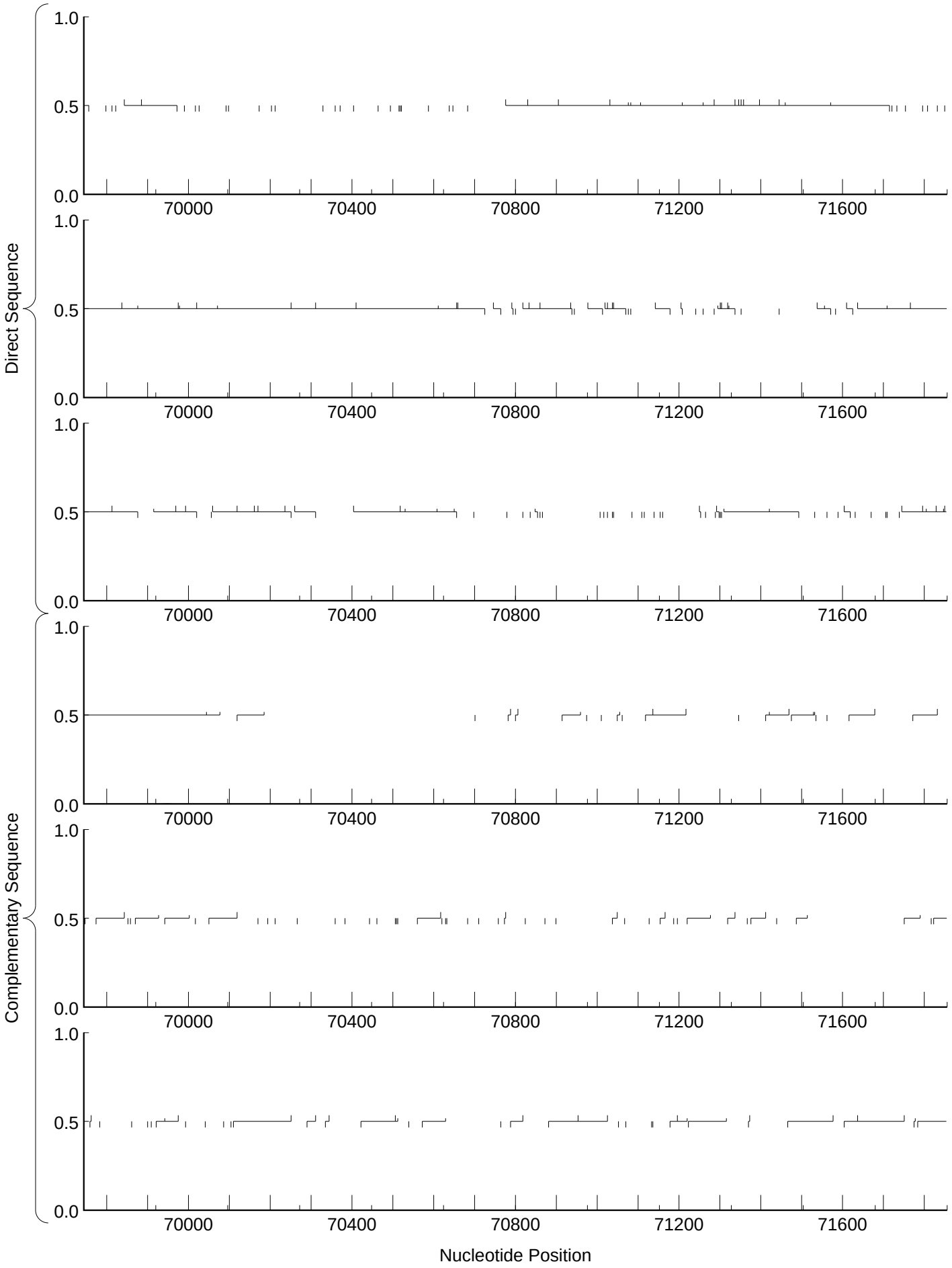
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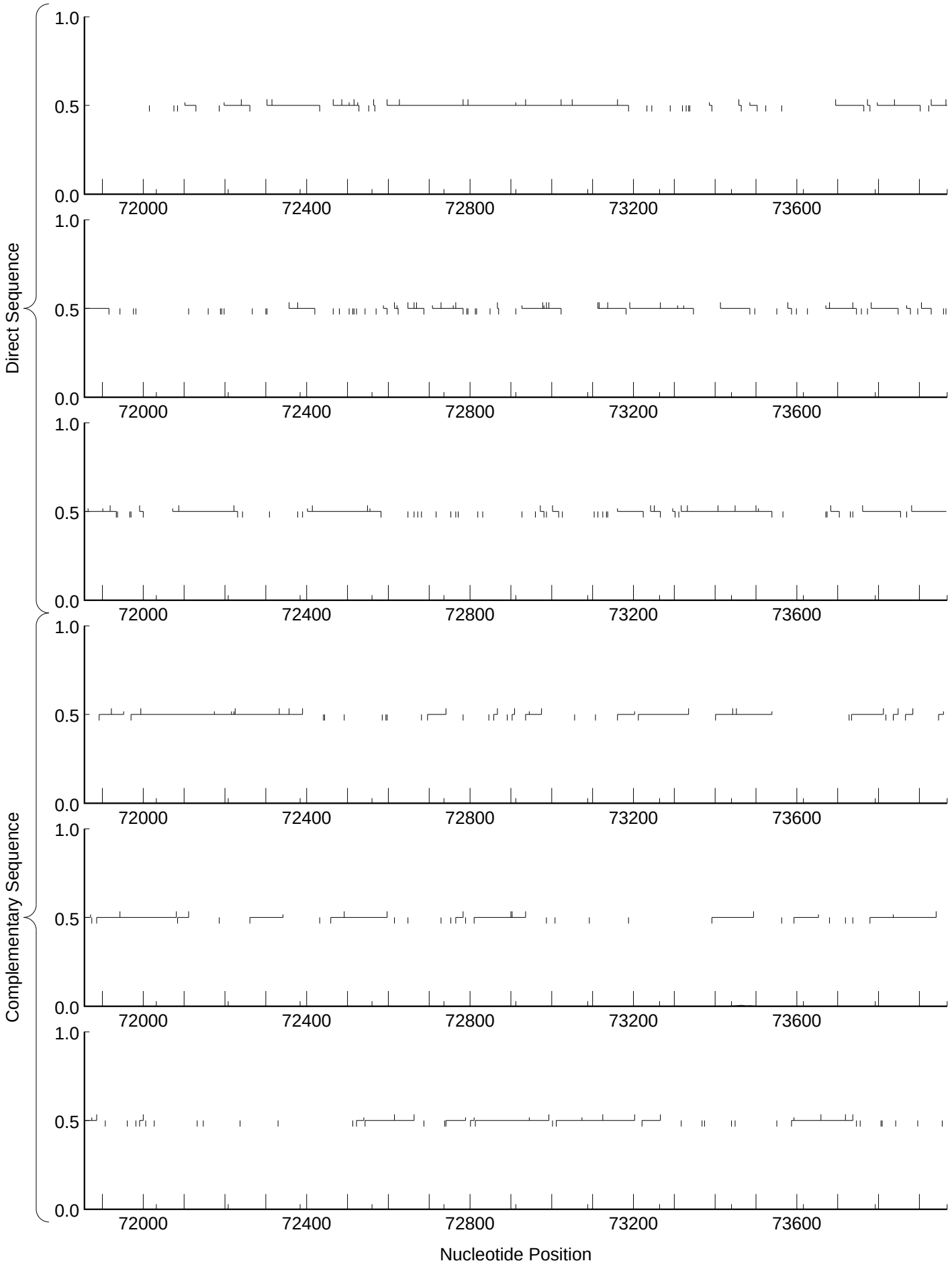
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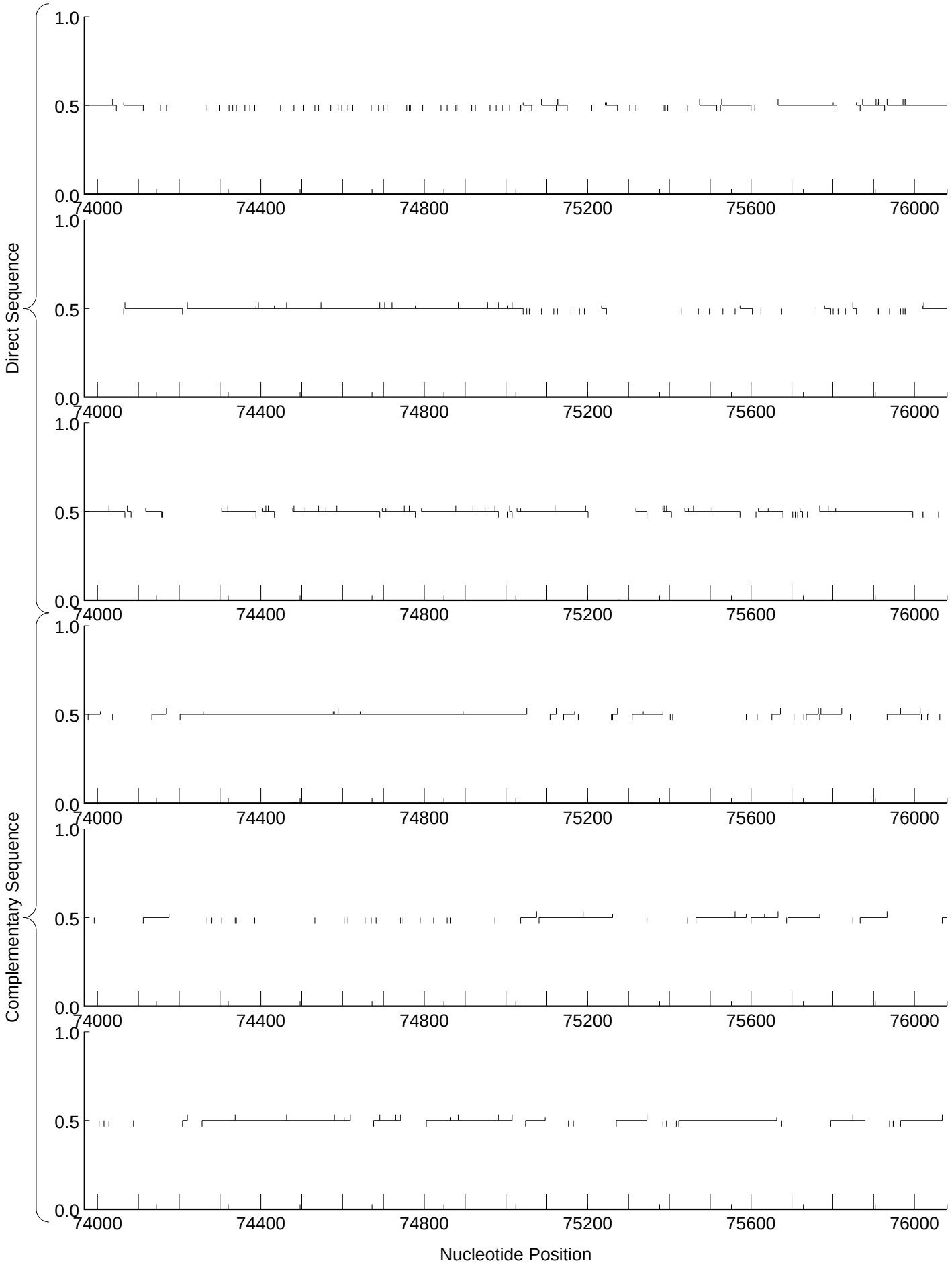
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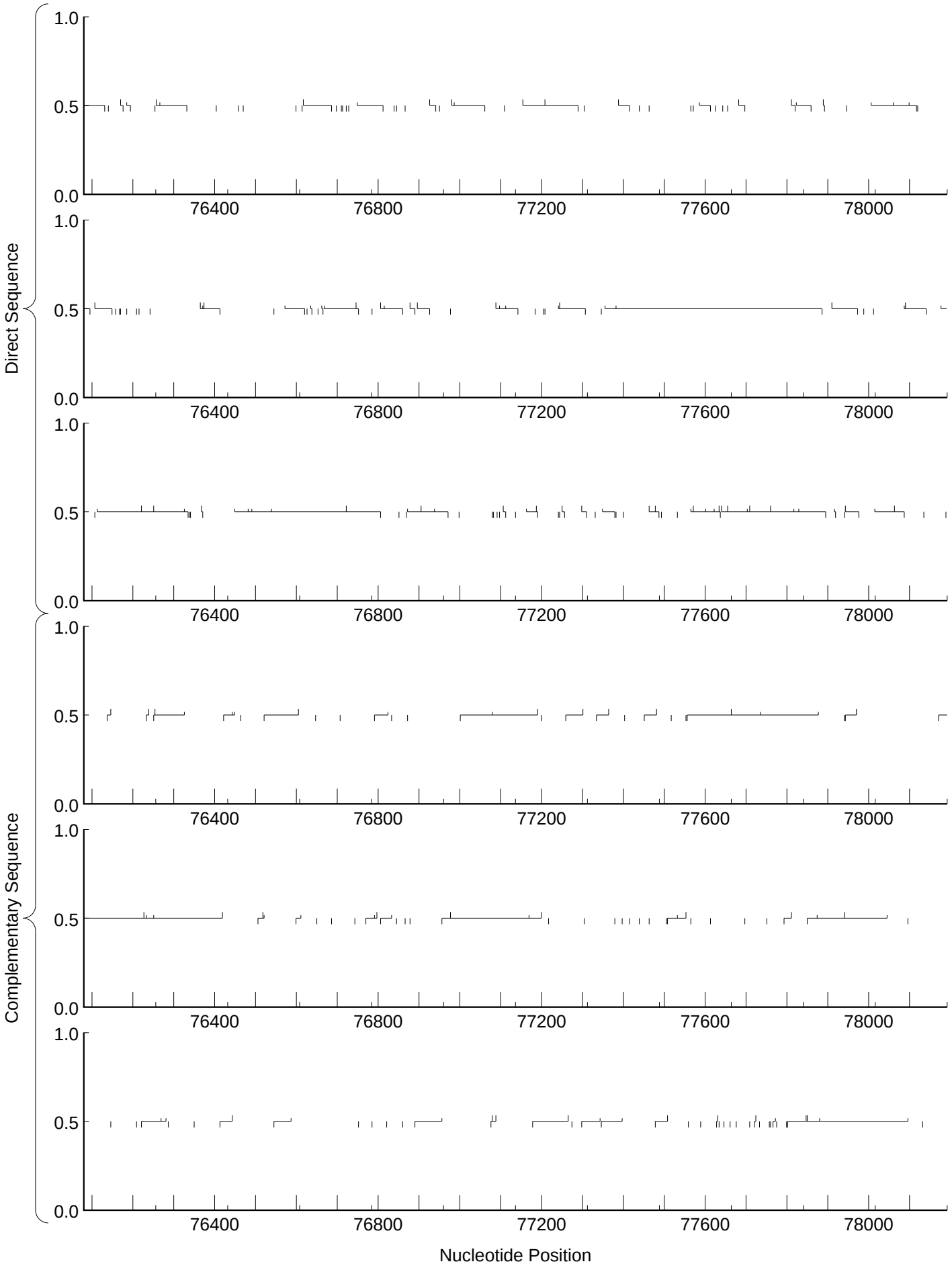
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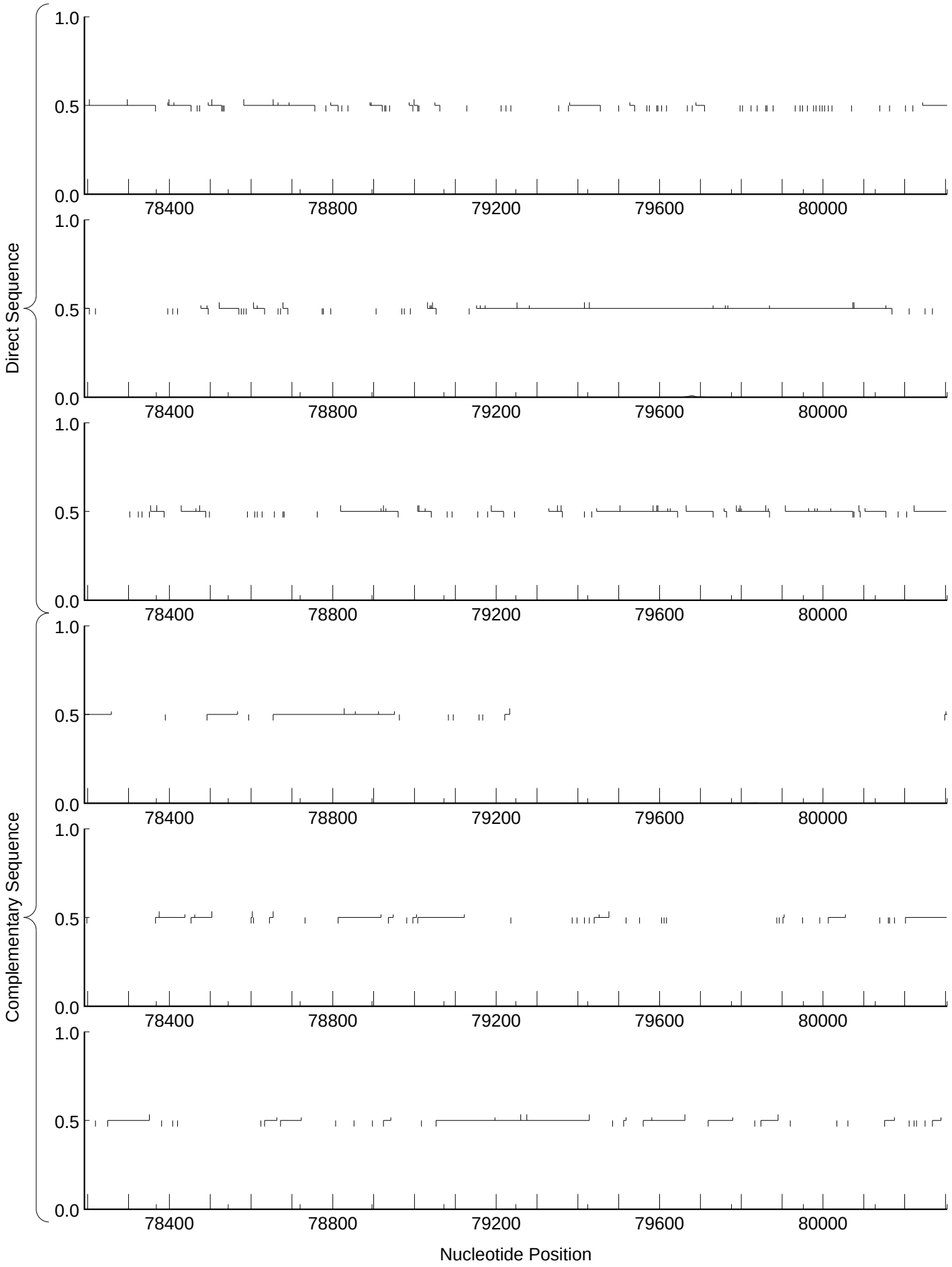
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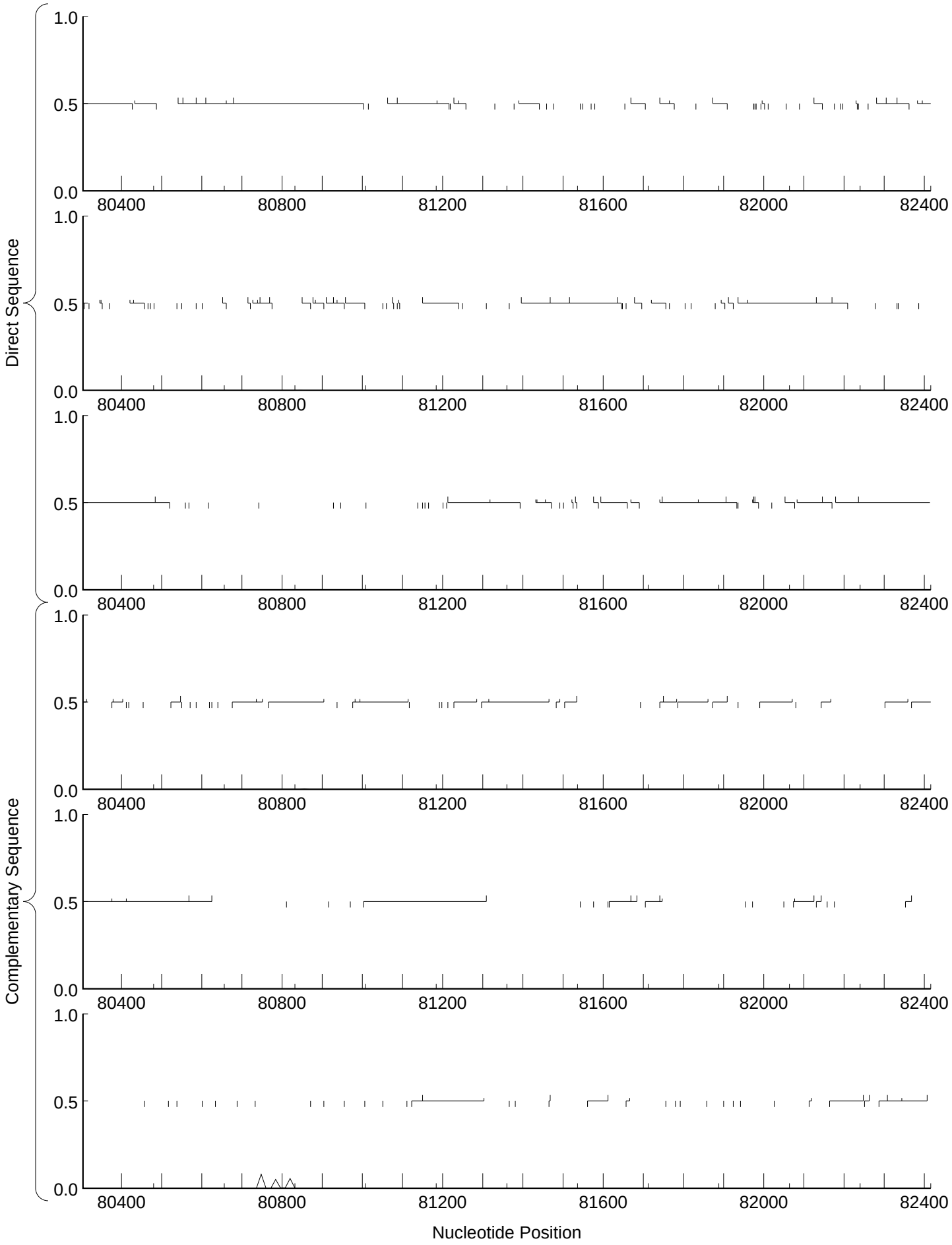
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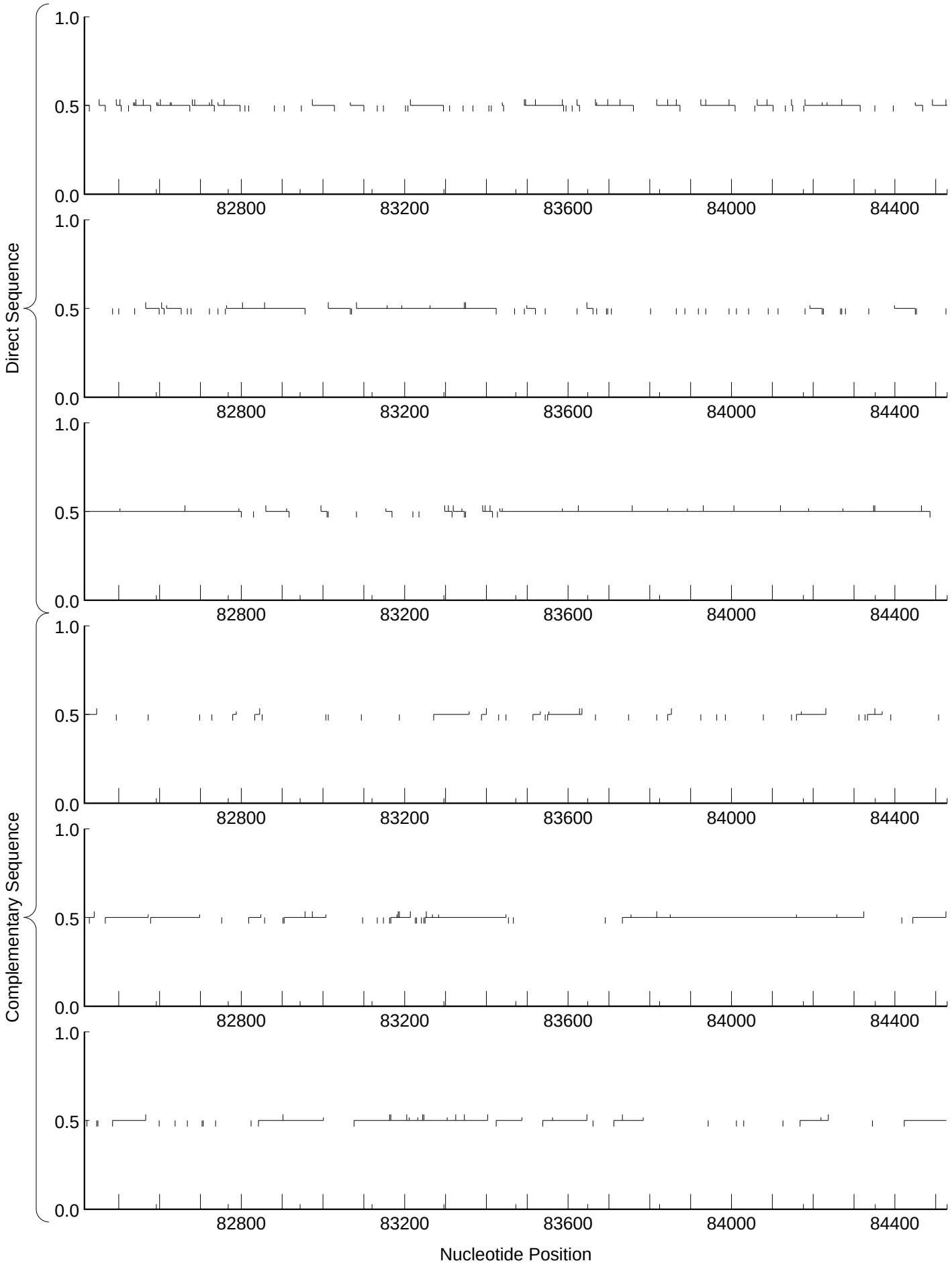
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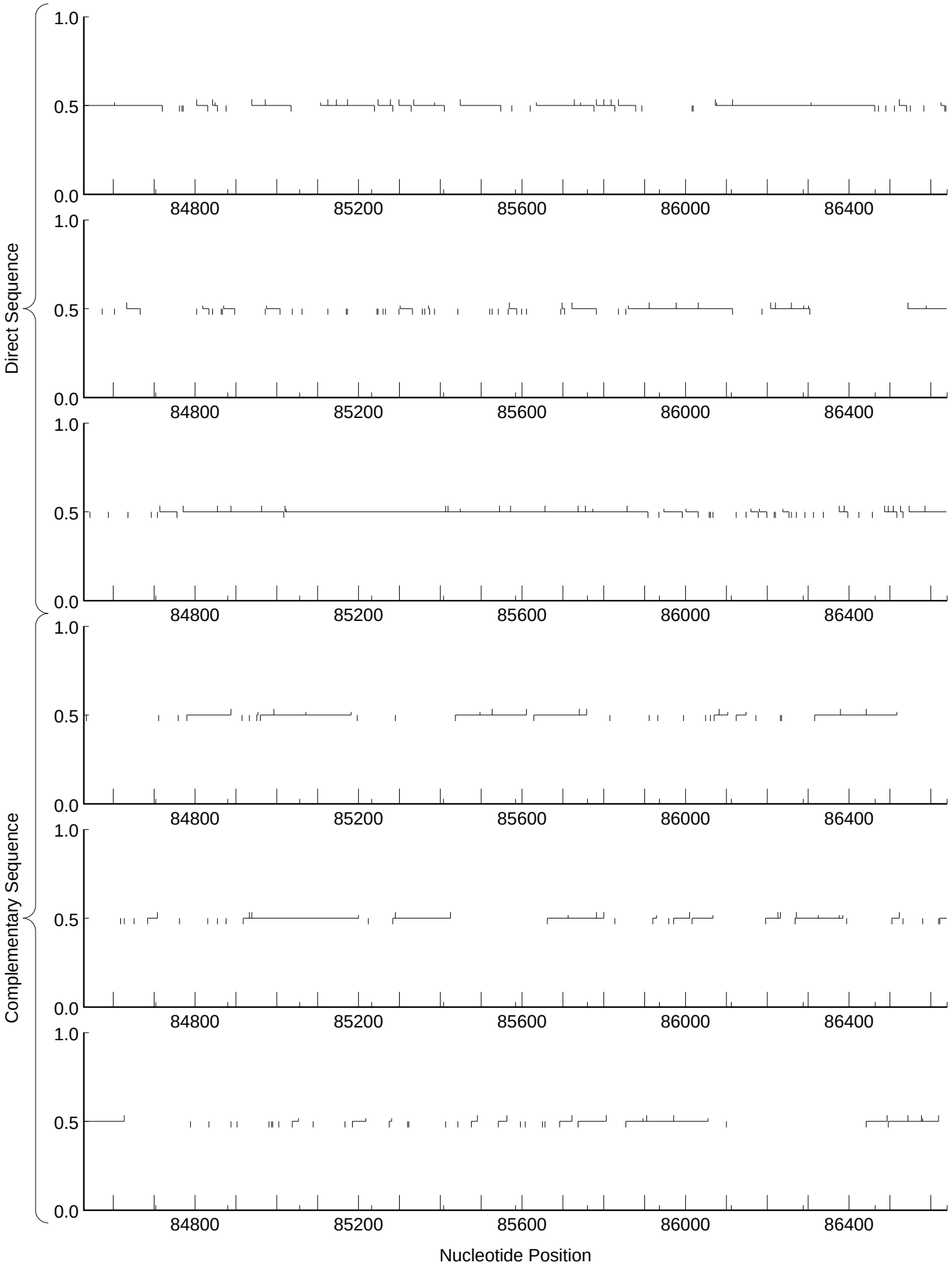
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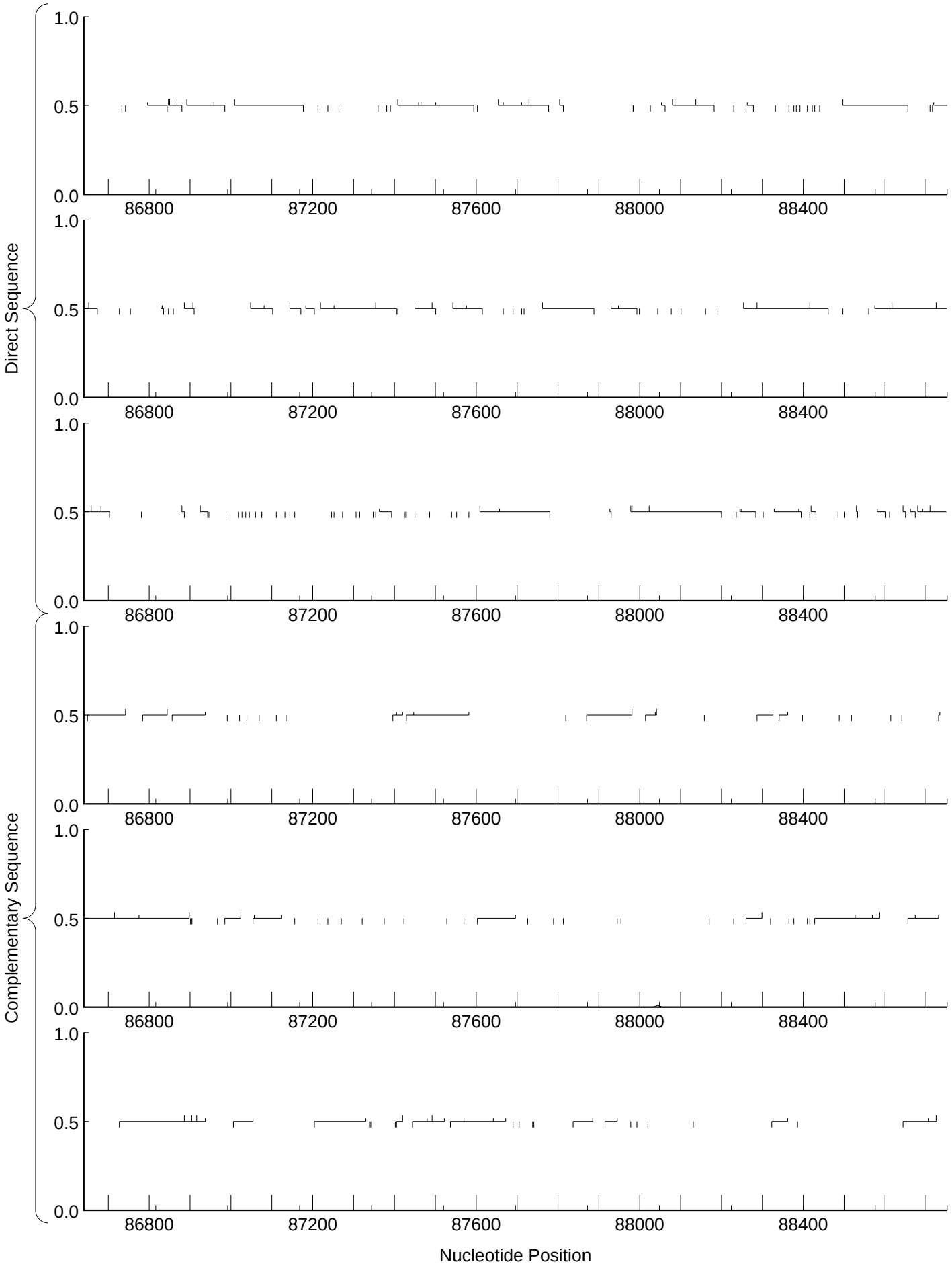
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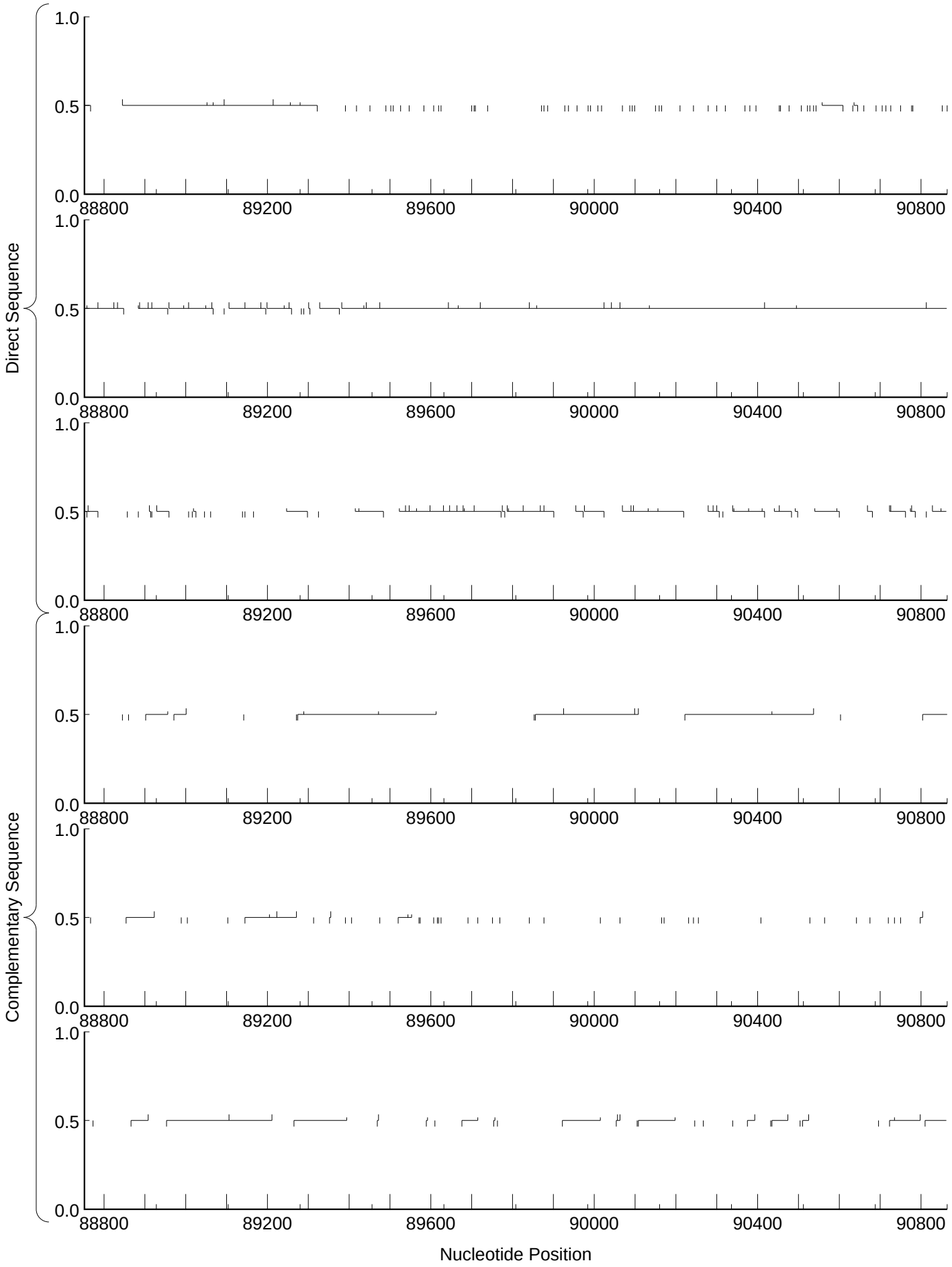
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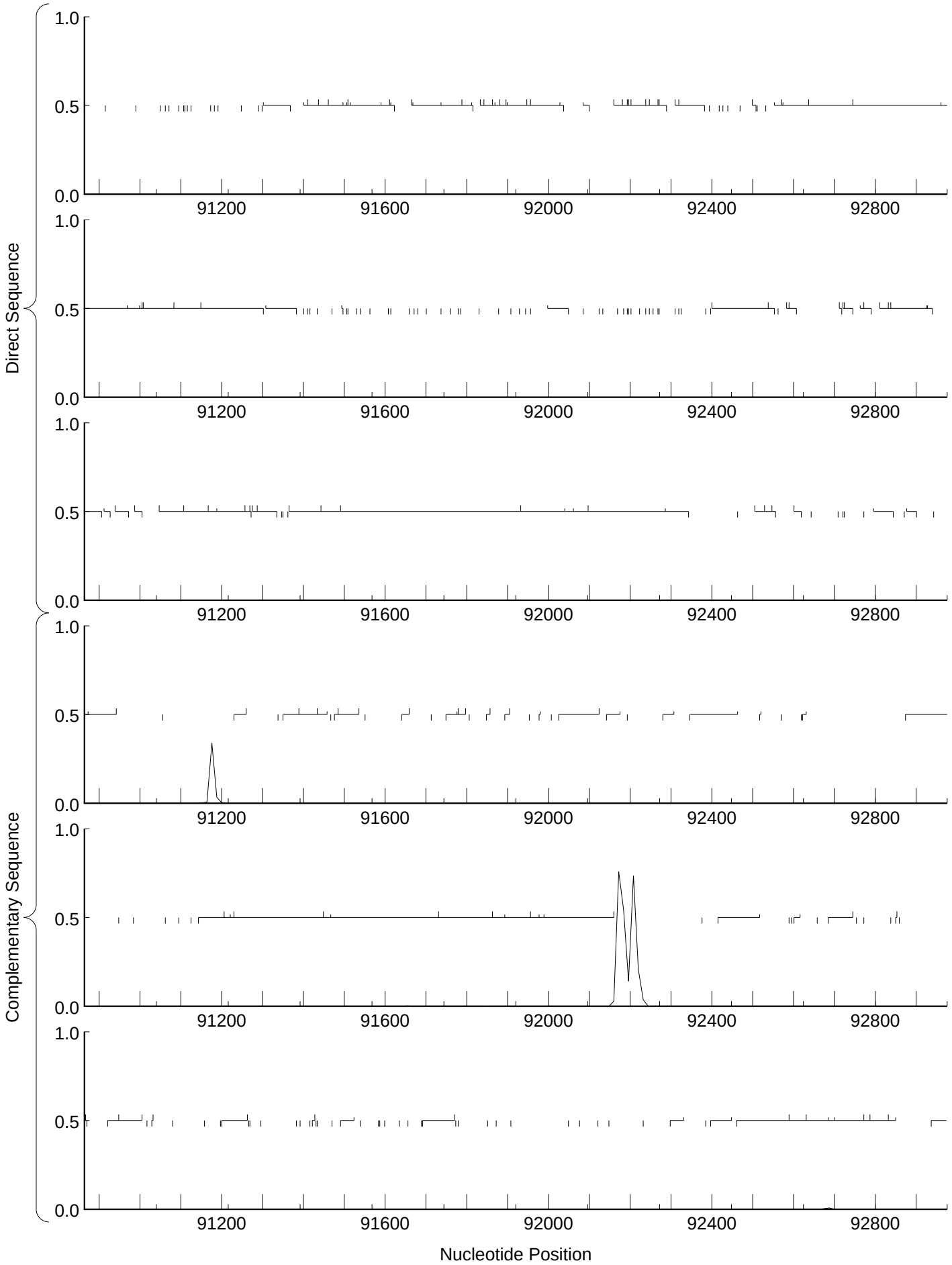
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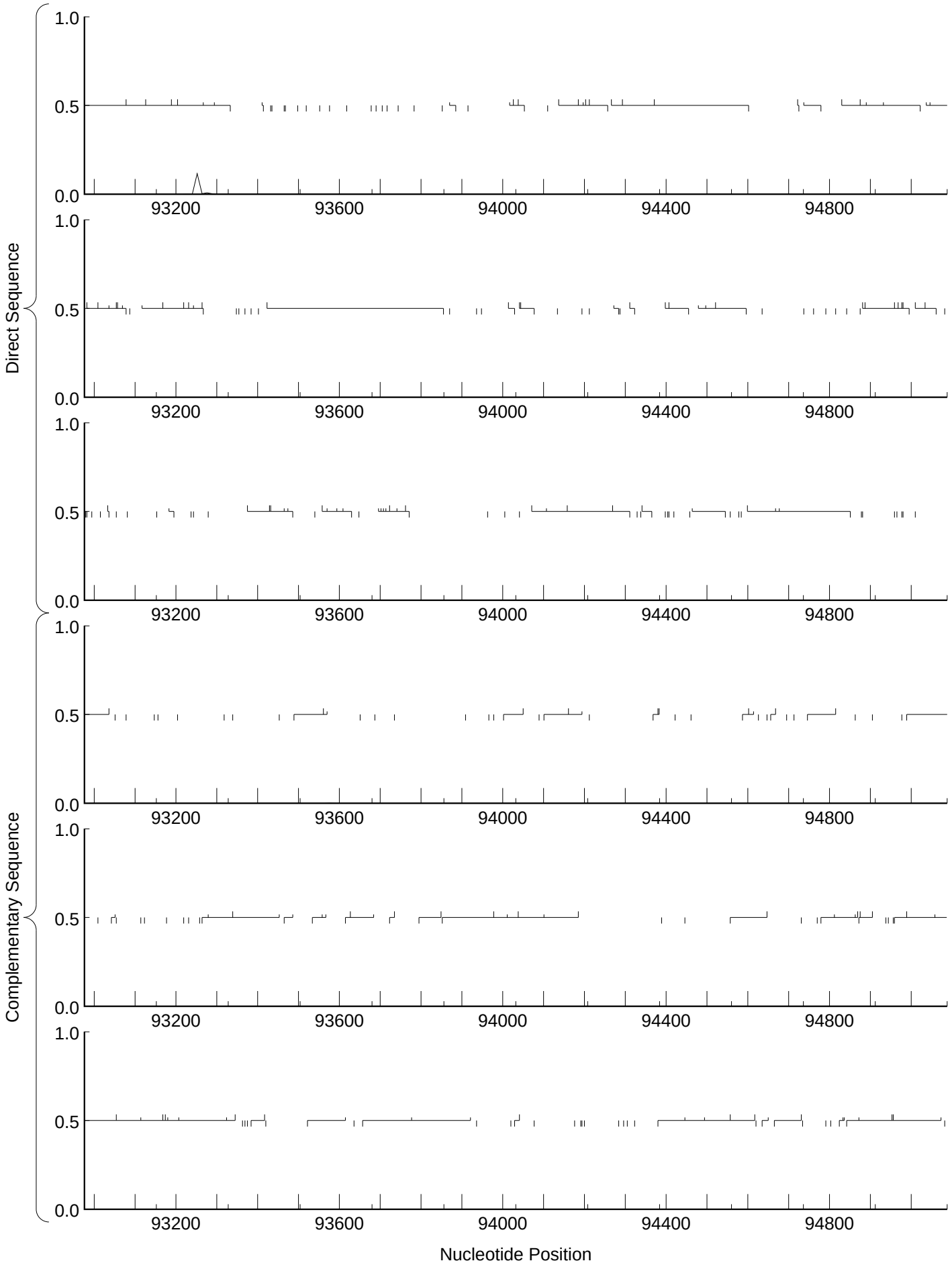
phage Sham complete sequence, 130327 bp including 787 bp terminal repeat, Cluster BK1 , Order 4, Window 96, Step 12, 44/63



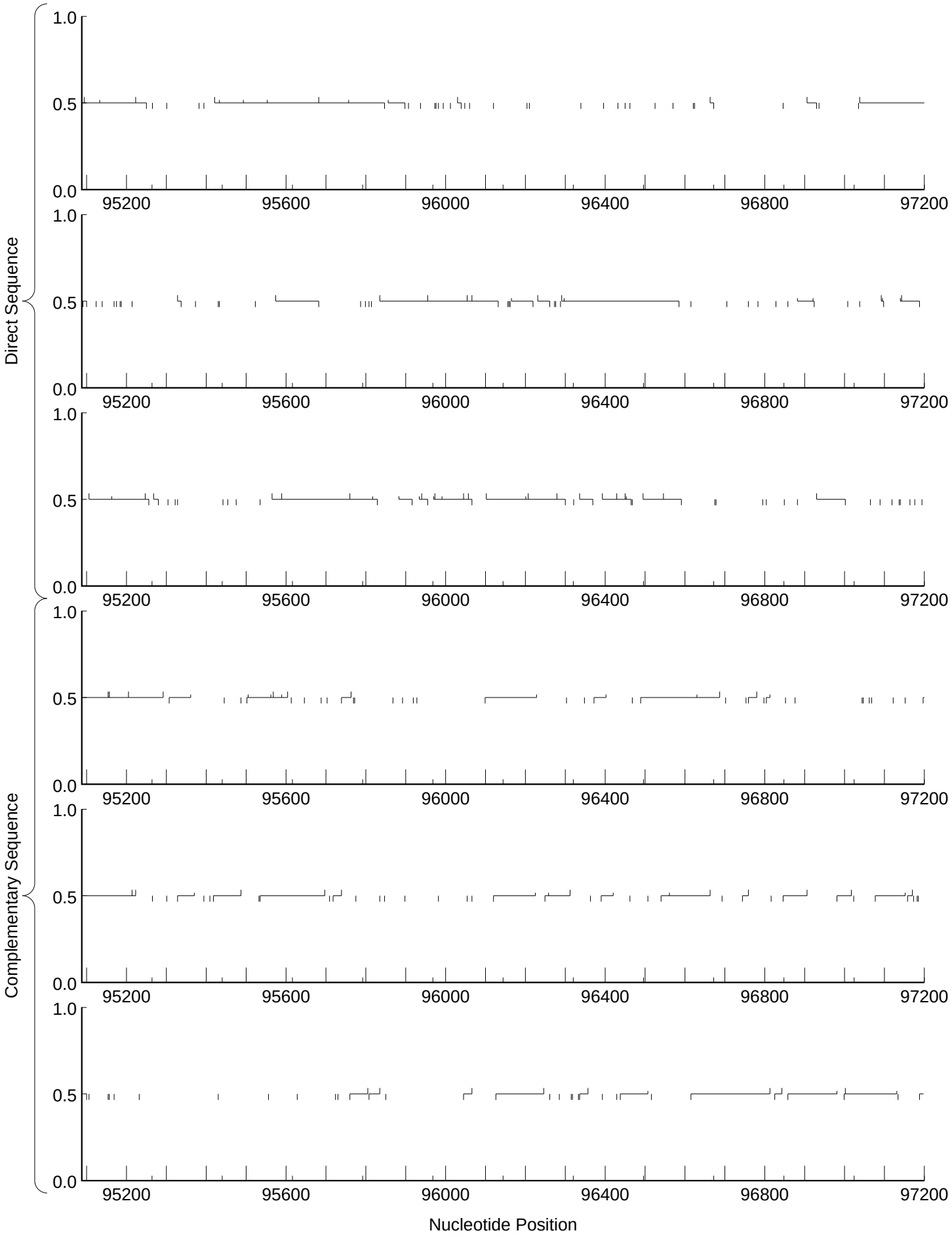
phage Sham complete sequence, 130327 bp including 787 bp terminal repeat, Cluster BK1 , Order 4, Window 96, Step 12, 45/63



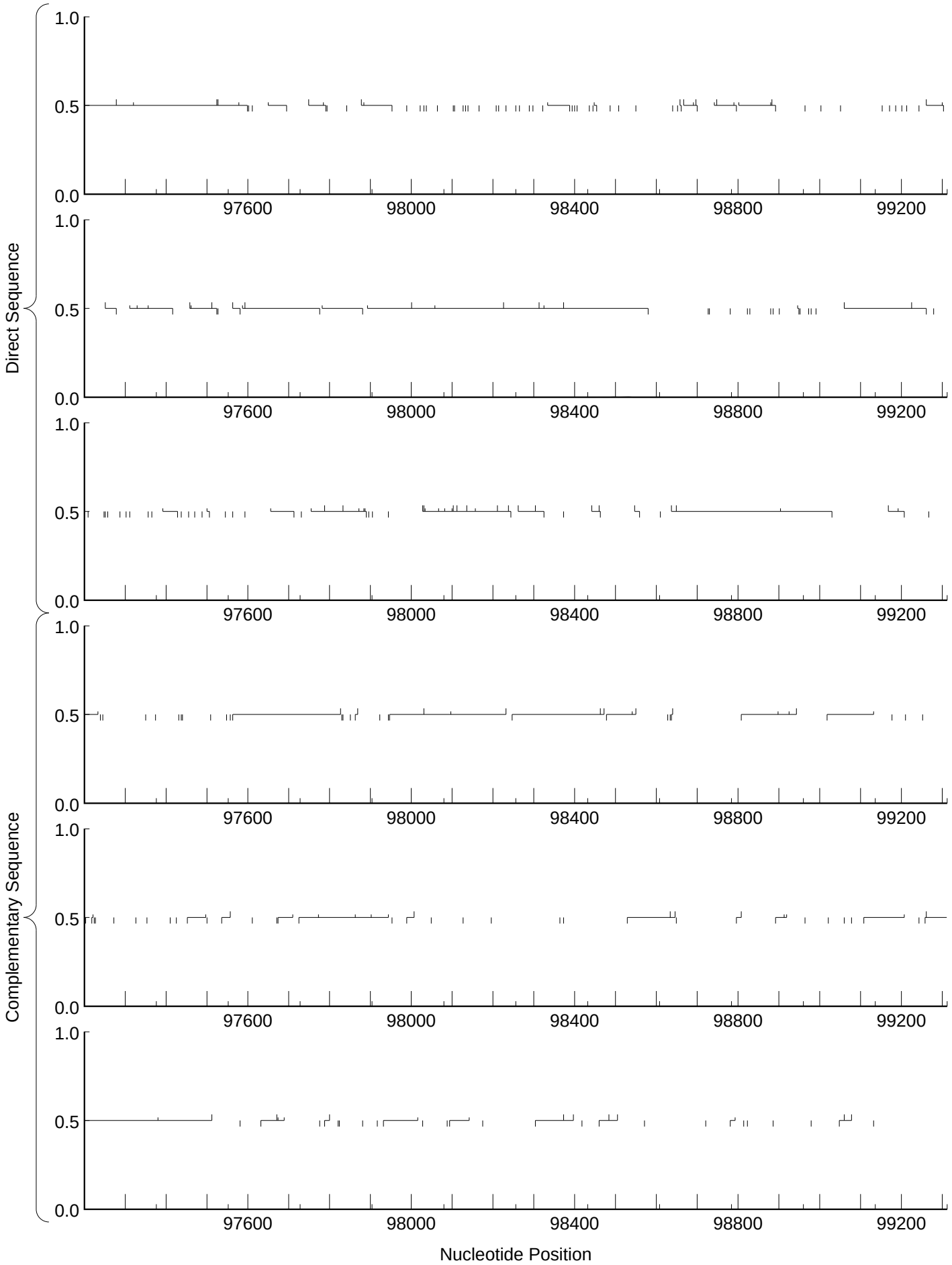
phage Sham complete sequence, 130327 bp including 787 bp terminal repeat, Cluster BK1 , Order 4, Window 96, Step 12, 46/63



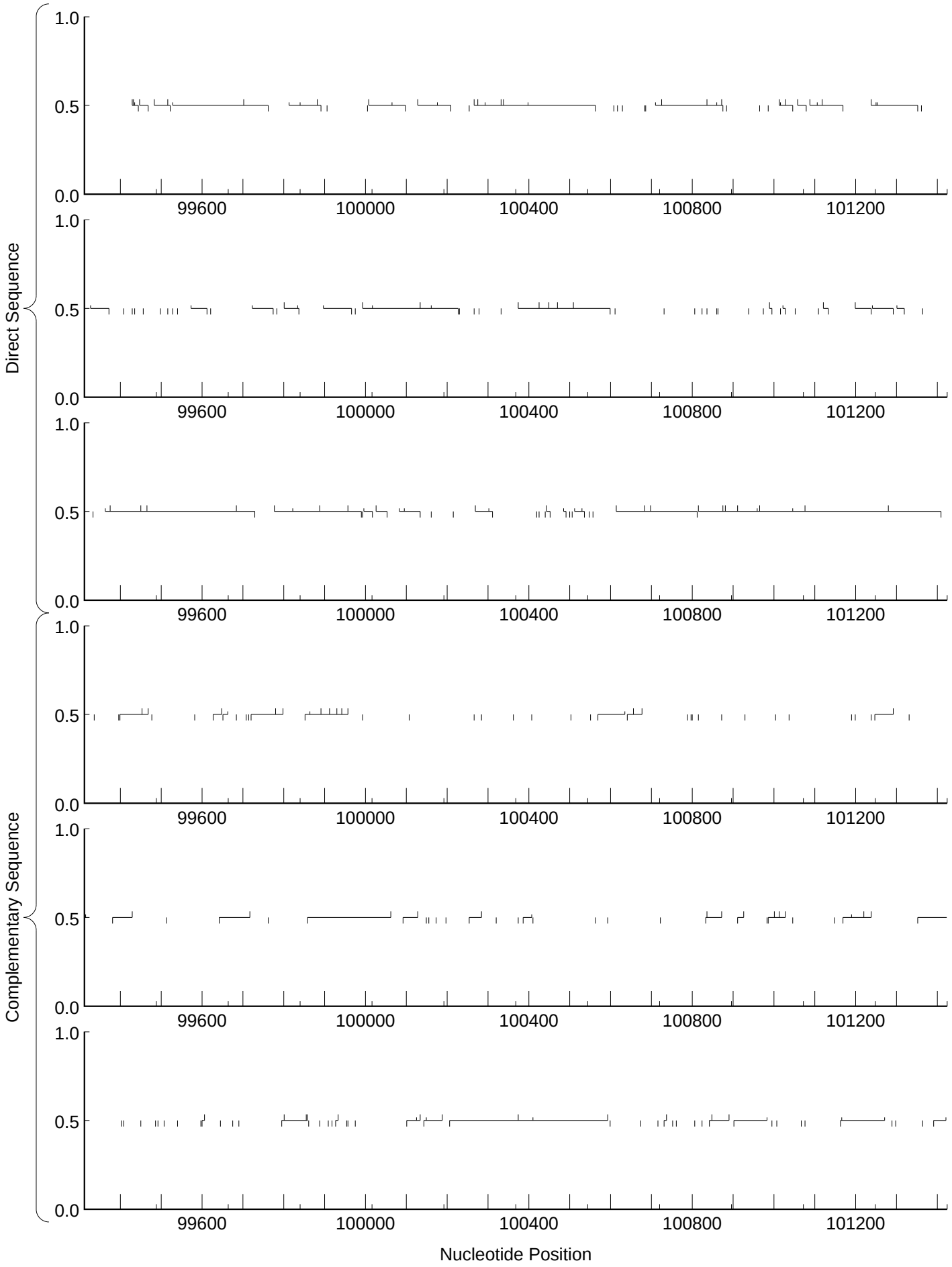
phage Sham complete sequence, 130327 bp including 787 bp terminal repeat, Cluster BK1 , Order 4, Window 96, Step 12, 47/63



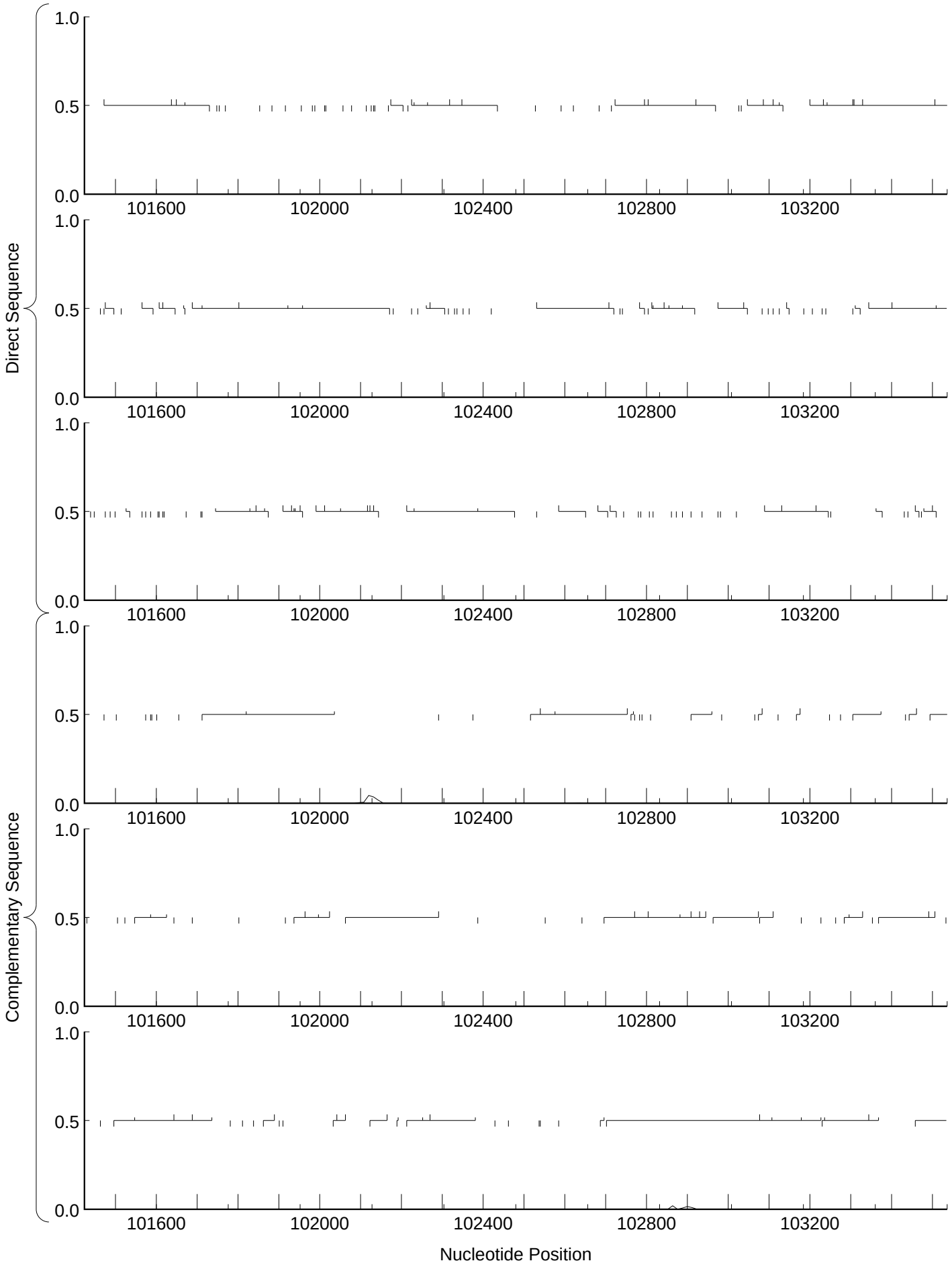
phage Sham complete sequence, 130327 bp including 787 bp terminal repeat, Cluster BK1 , Order 4, Window 96, Step 12, 48/63



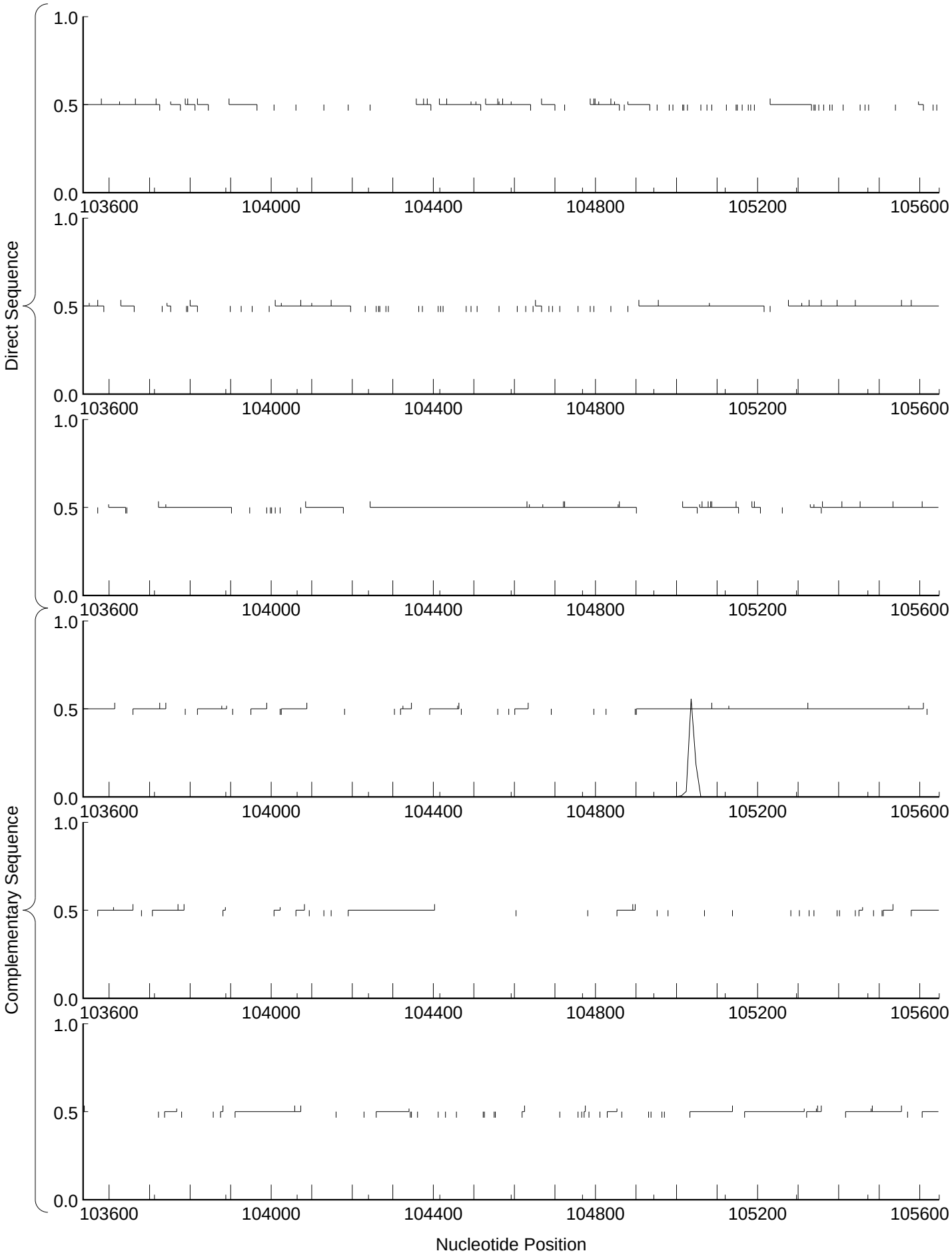
phage Sham complete sequence, 130327 bp including 787 bp terminal repeat, Cluster BK1 , Order 4, Window 96, Step 12, 49/63



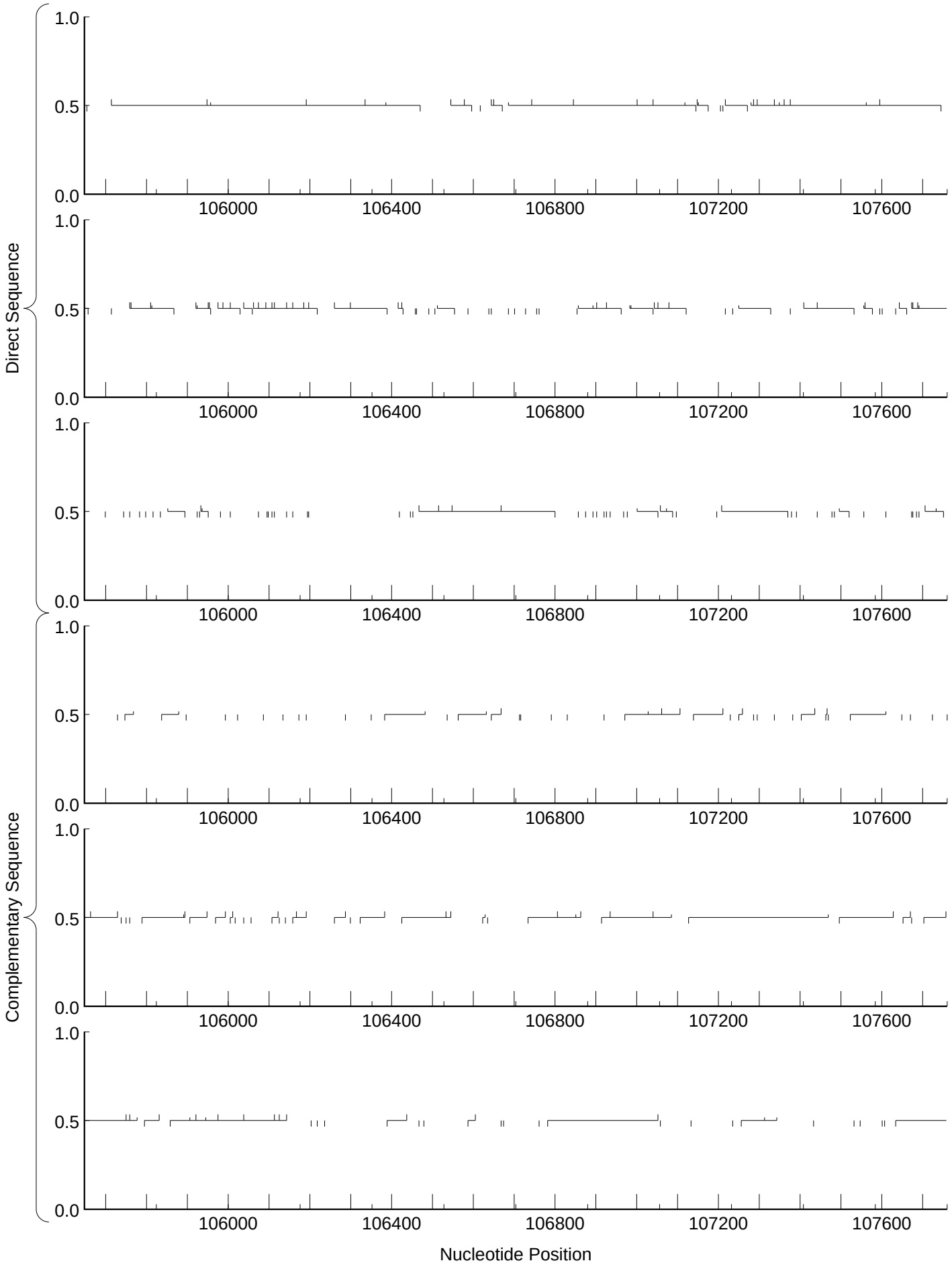
phage Sham complete sequence, 130327 bp including 787 bp terminal repeat, Cluster BK1 , Order 4, Window 96, Step 12, 50/63



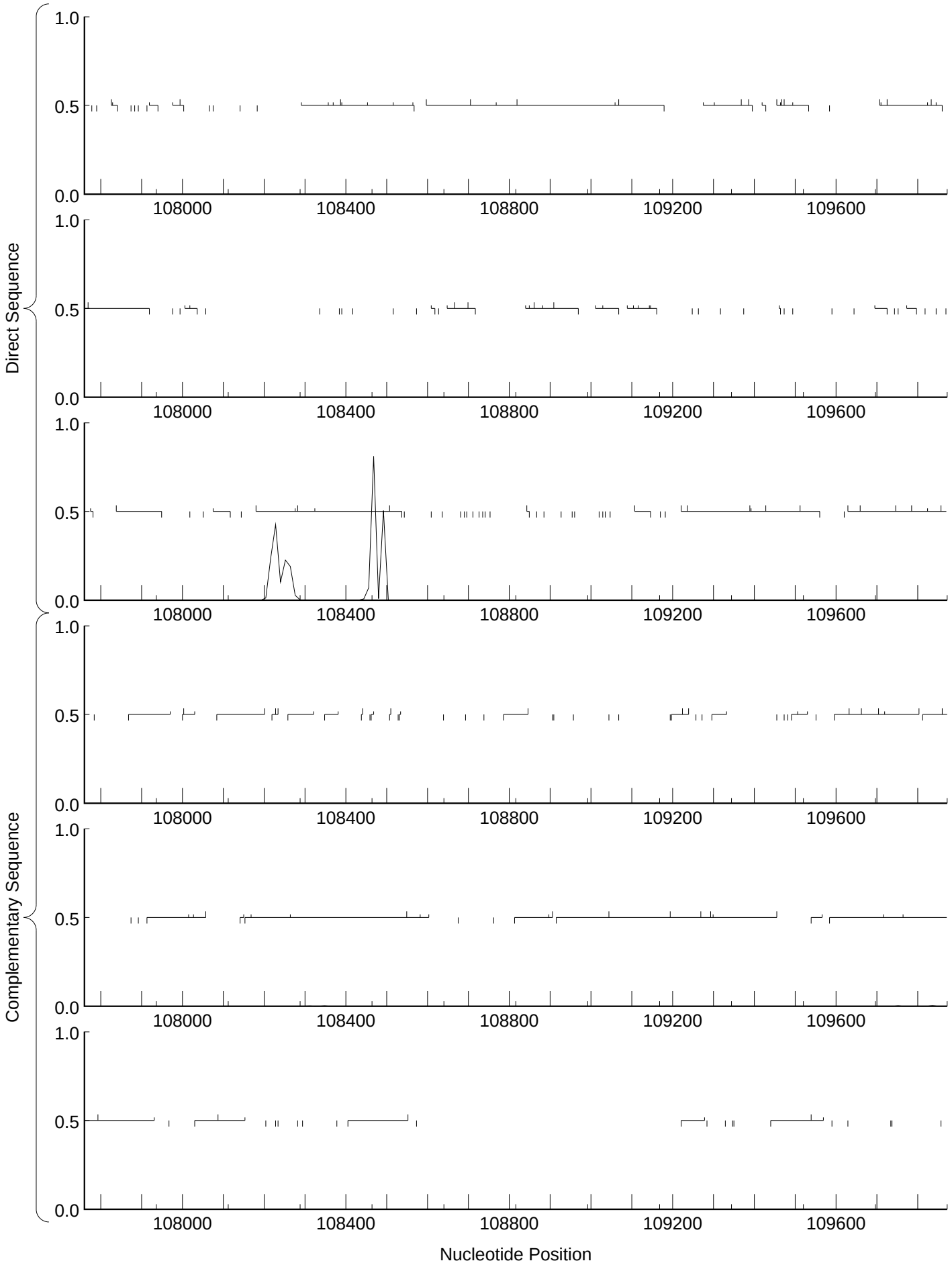
phage Sham complete sequence, 130327 bp including 787 bp terminal repeat, Cluster BK1 , Order 4, Window 96, Step 12, 51/63



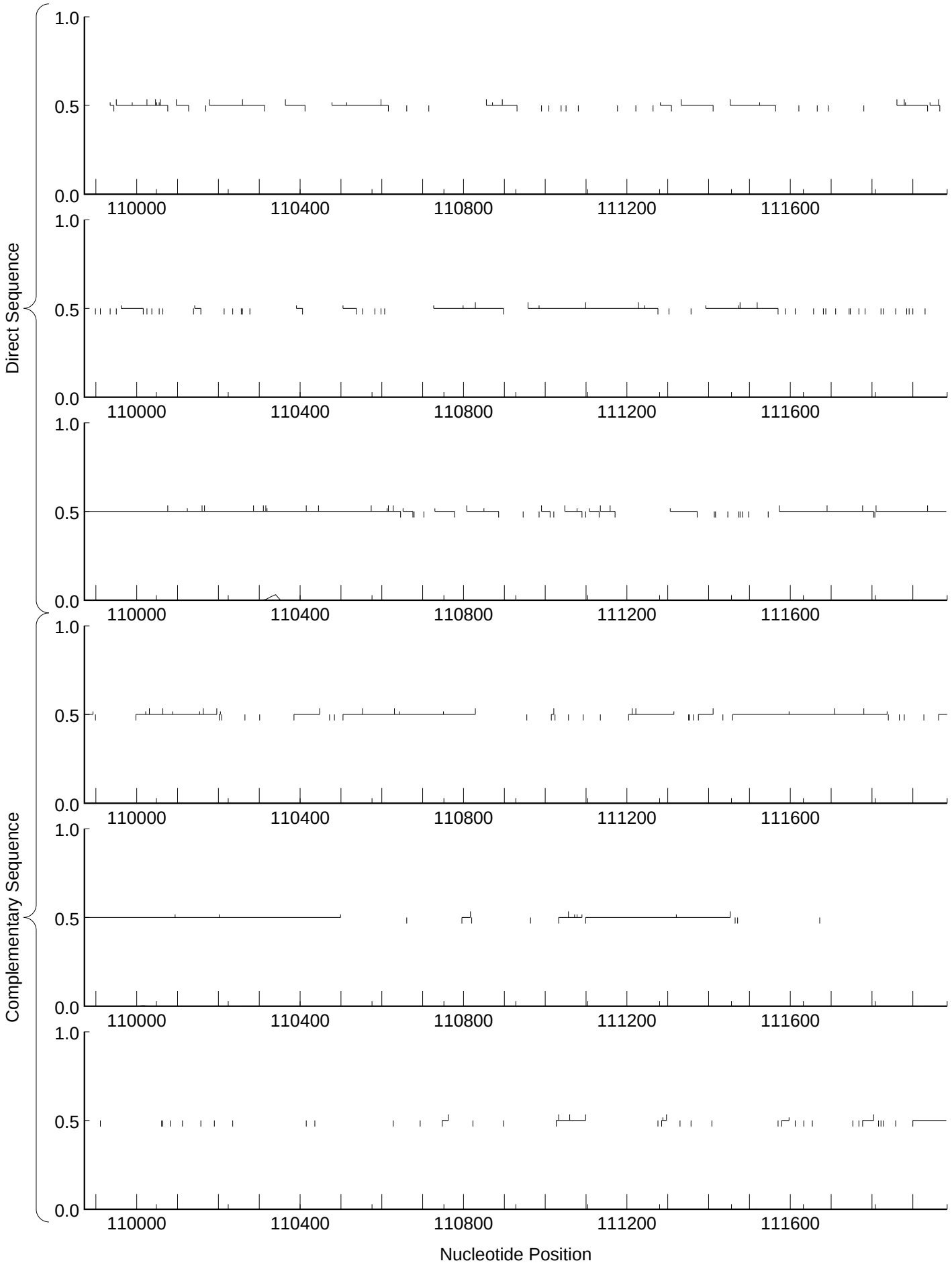
phage Sham complete sequence, 130327 bp including 787 bp terminal repeat, Cluster BK1 , Order 4, Window 96, Step 12, 52/63



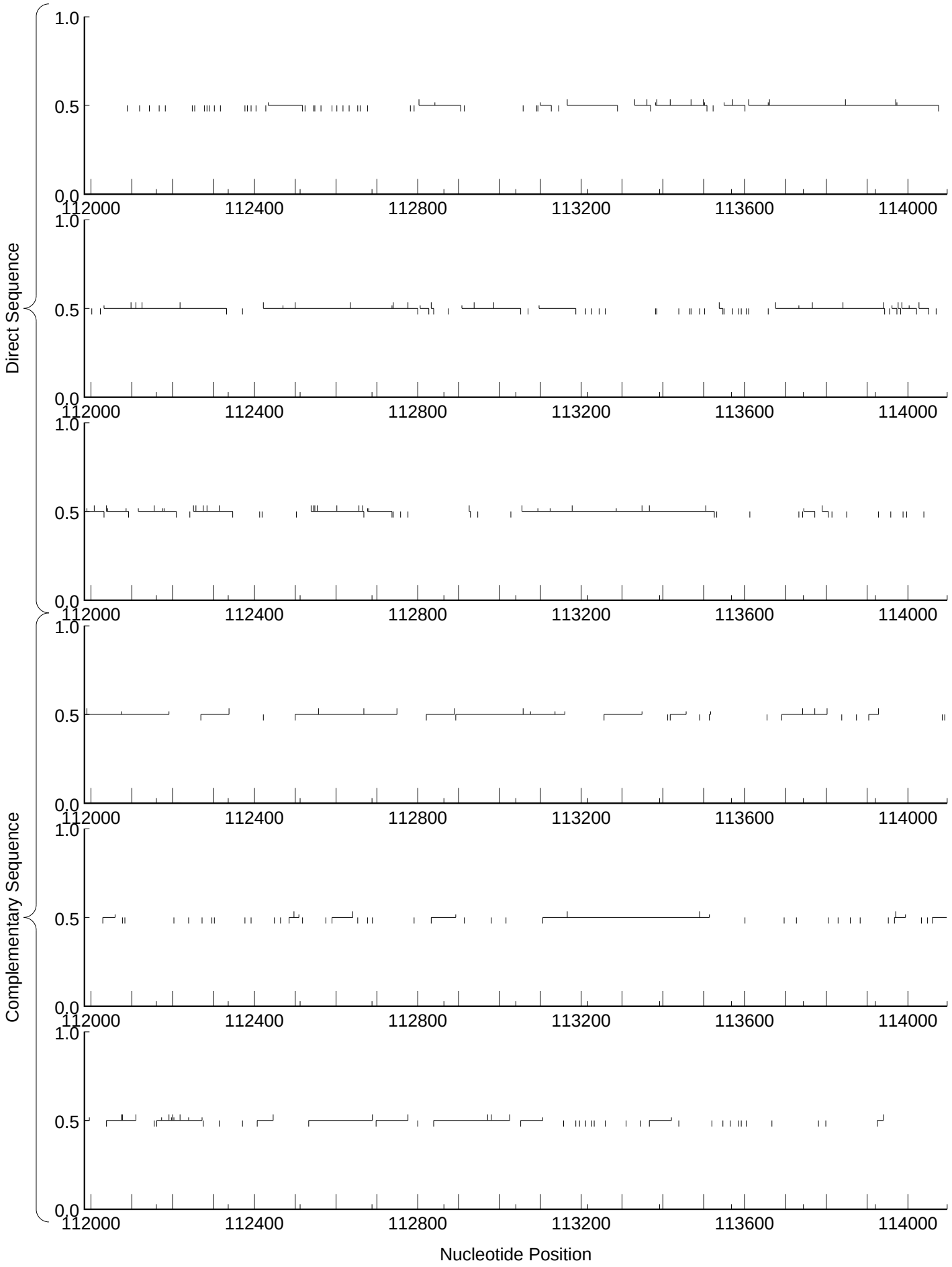
phage Sham complete sequence, 130327 bp including 787 bp terminal repeat, Cluster BK1 , Order 4, Window 96, Step 12, 53/63



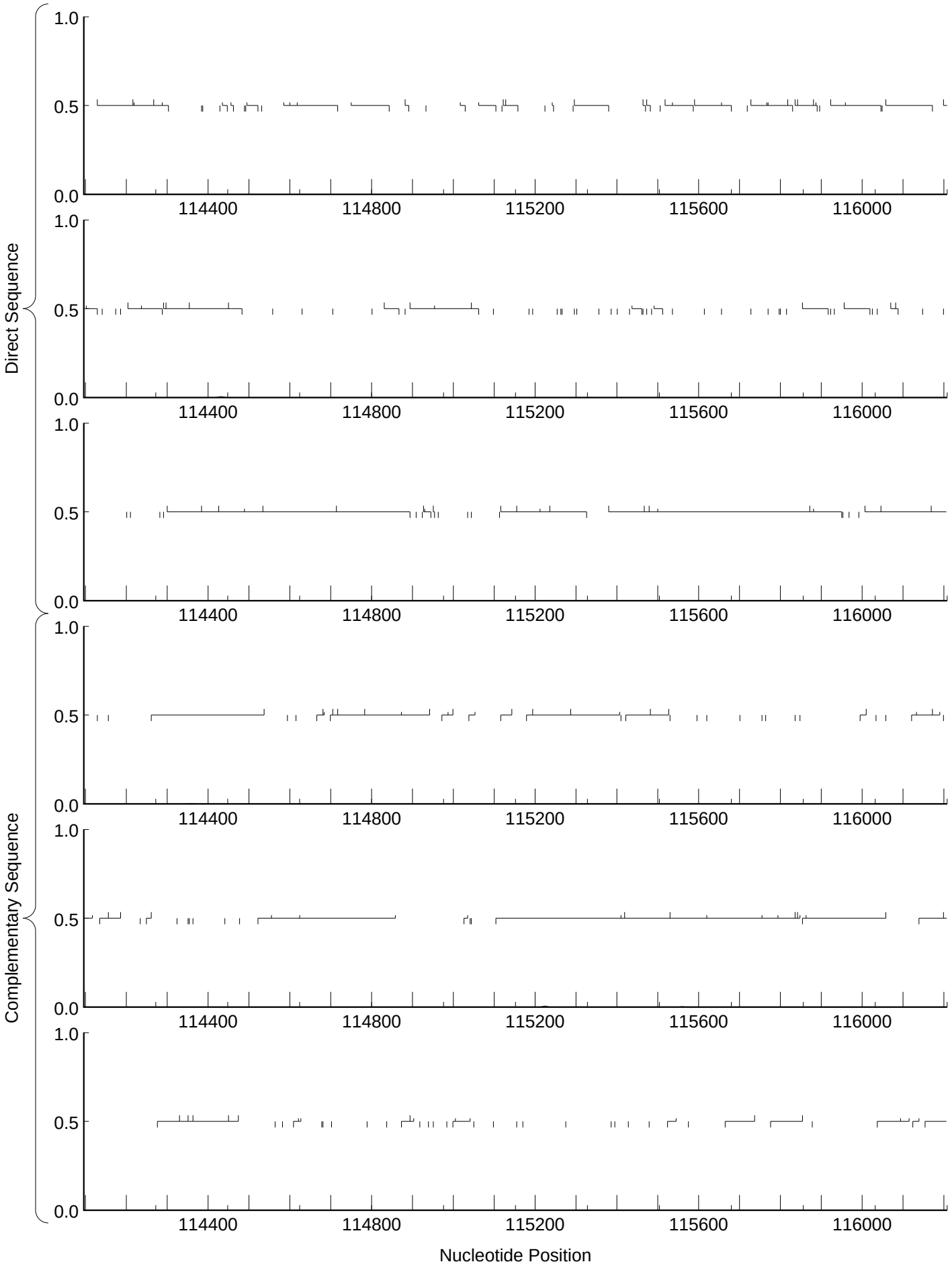
phage Sham complete sequence, 130327 bp including 787 bp terminal repeat, Cluster BK1 , Order 4, Window 96, Step 12, 54/63



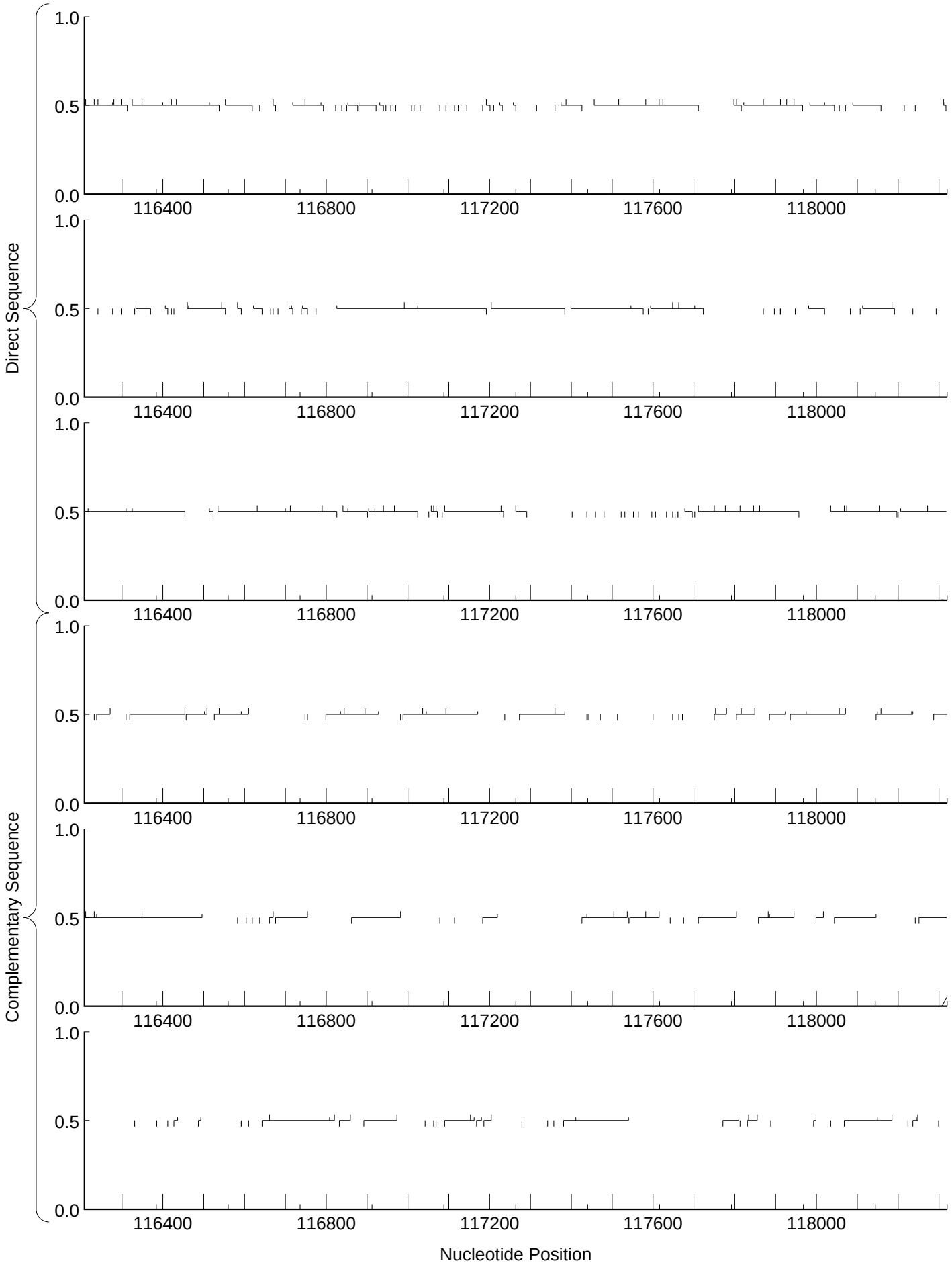
phage Sham complete sequence, 130327 bp including 787 bp terminal repeat, Cluster BK1 , Order 4, Window 96, Step 12, 55/63



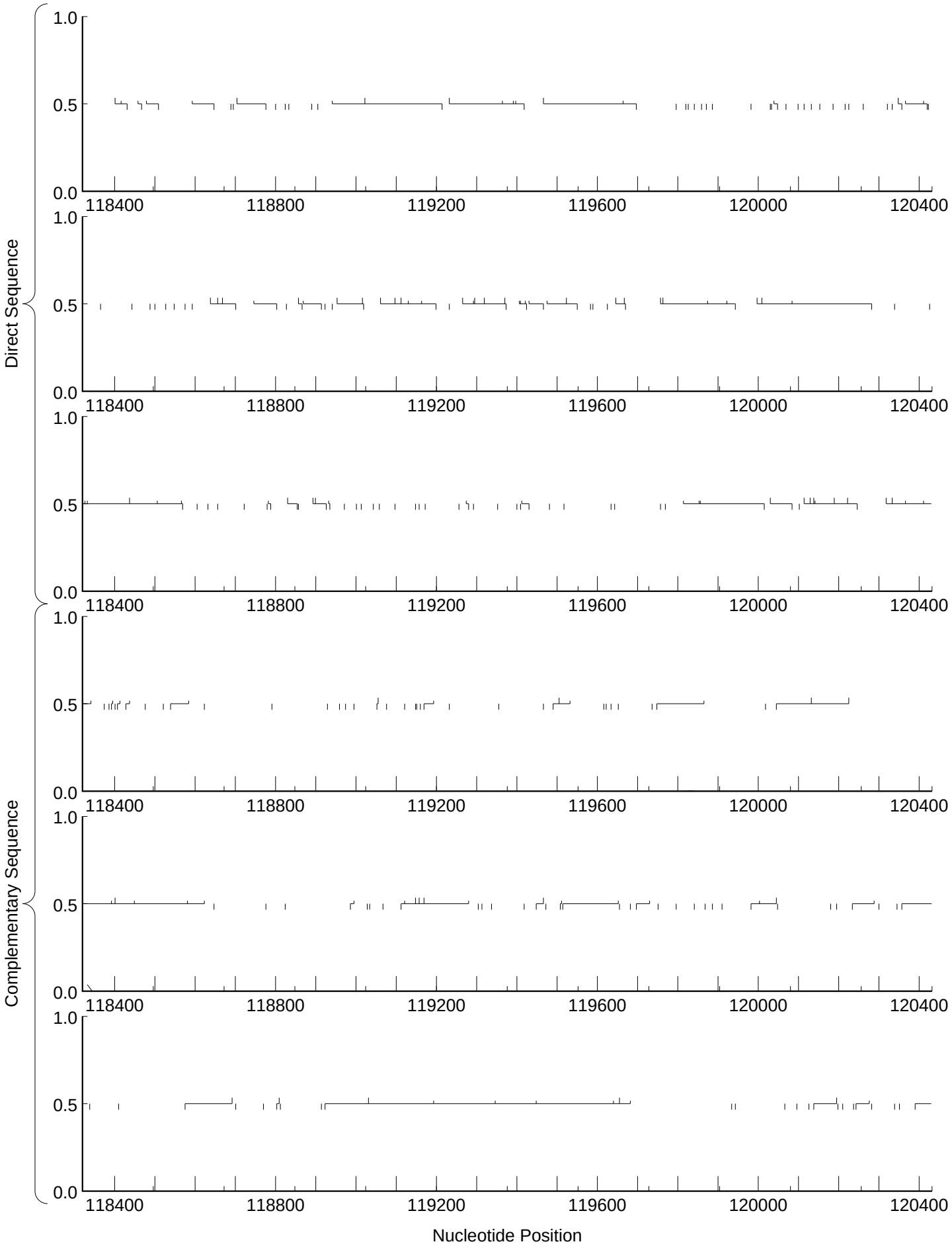
phage Sham complete sequence, 130327 bp including 787 bp terminal repeat, Cluster BK1 , Order 4, Window 96, Step 12, 56/63



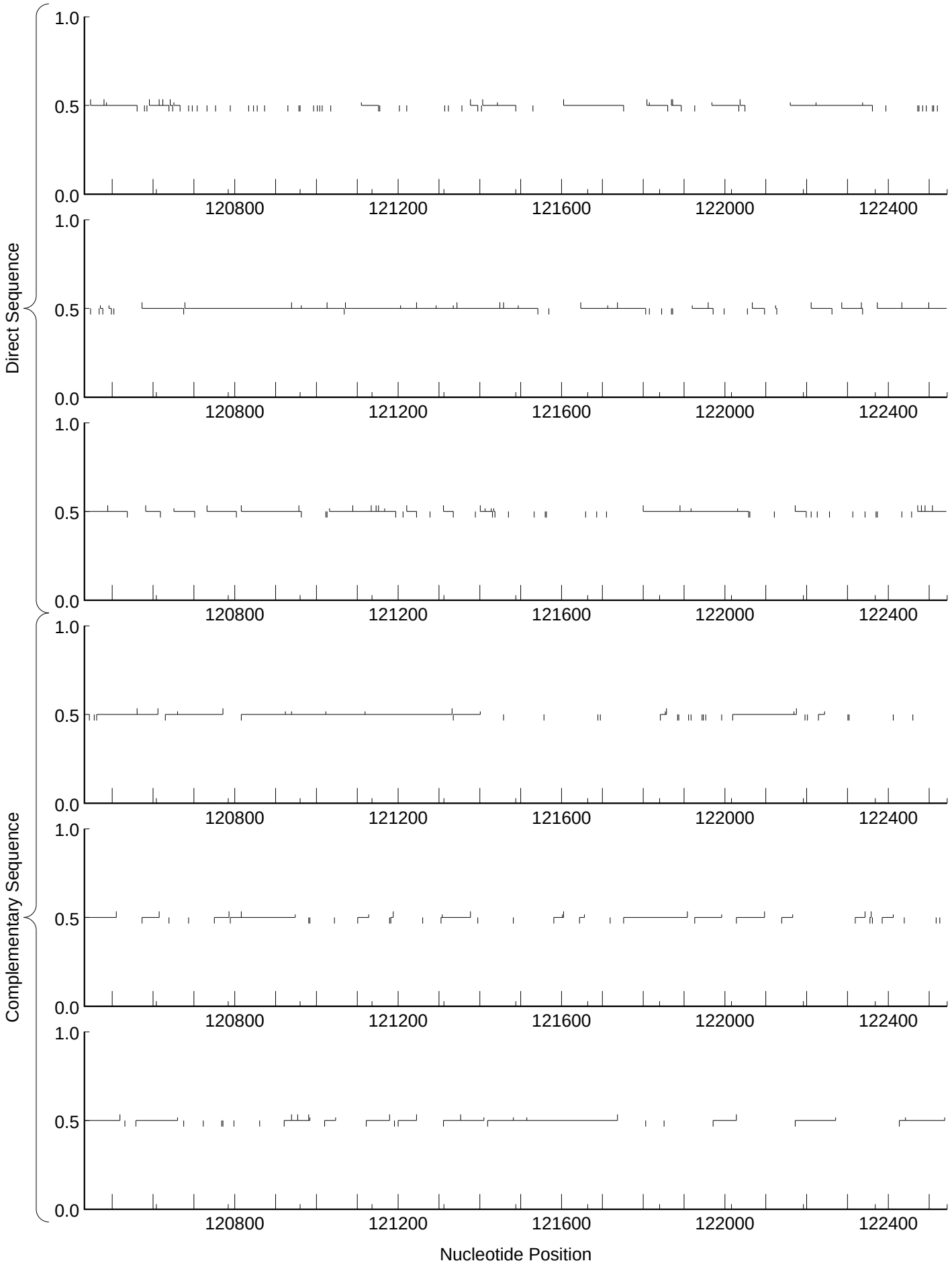
phage Sham complete sequence, 130327 bp including 787 bp terminal repeat, Cluster BK1 , Order 4, Window 96, Step 12, 57/63



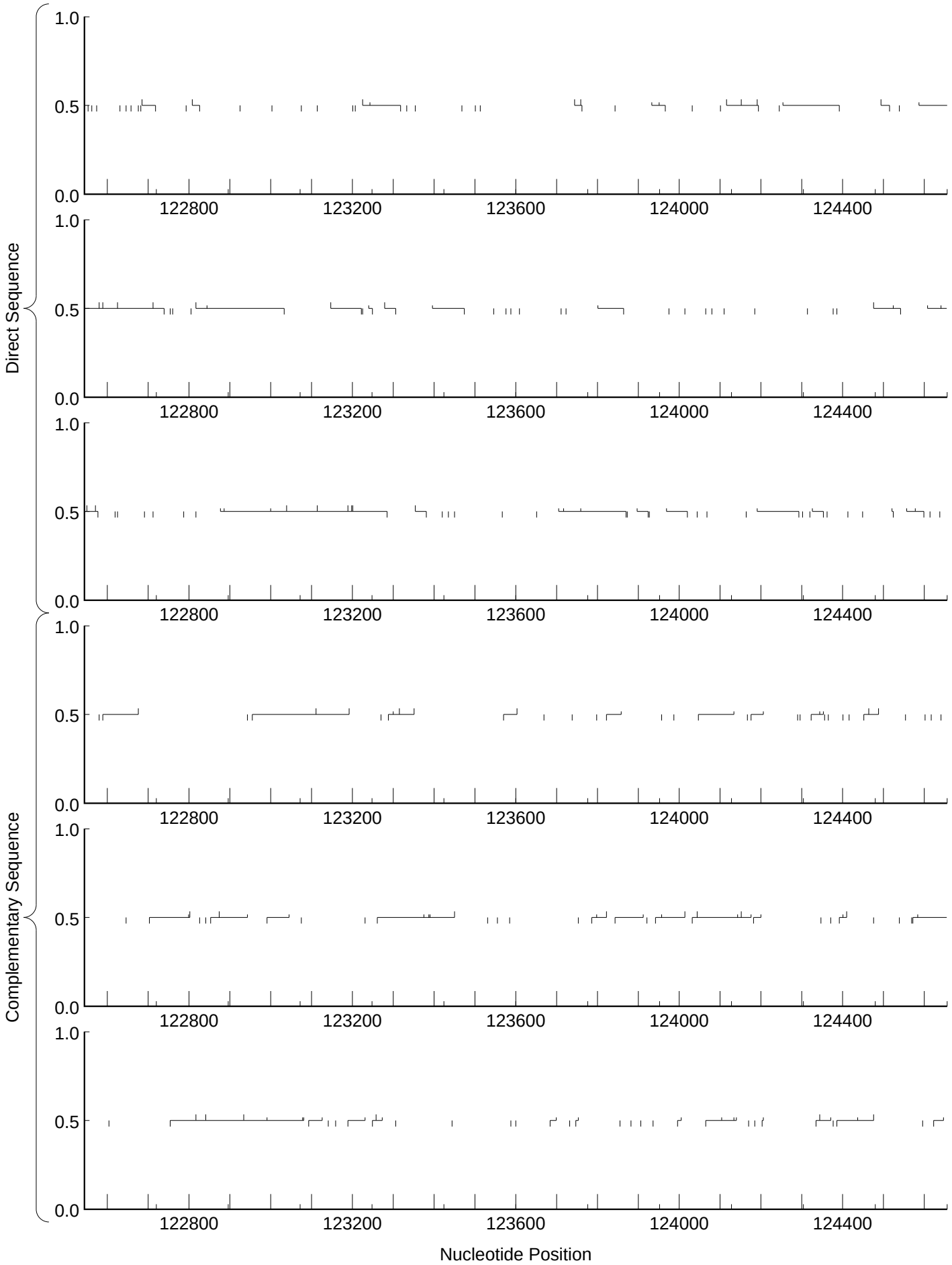
phage Sham complete sequence, 130327 bp including 787 bp terminal repeat, Cluster BK1 , Order 4, Window 96, Step 12, 58/63



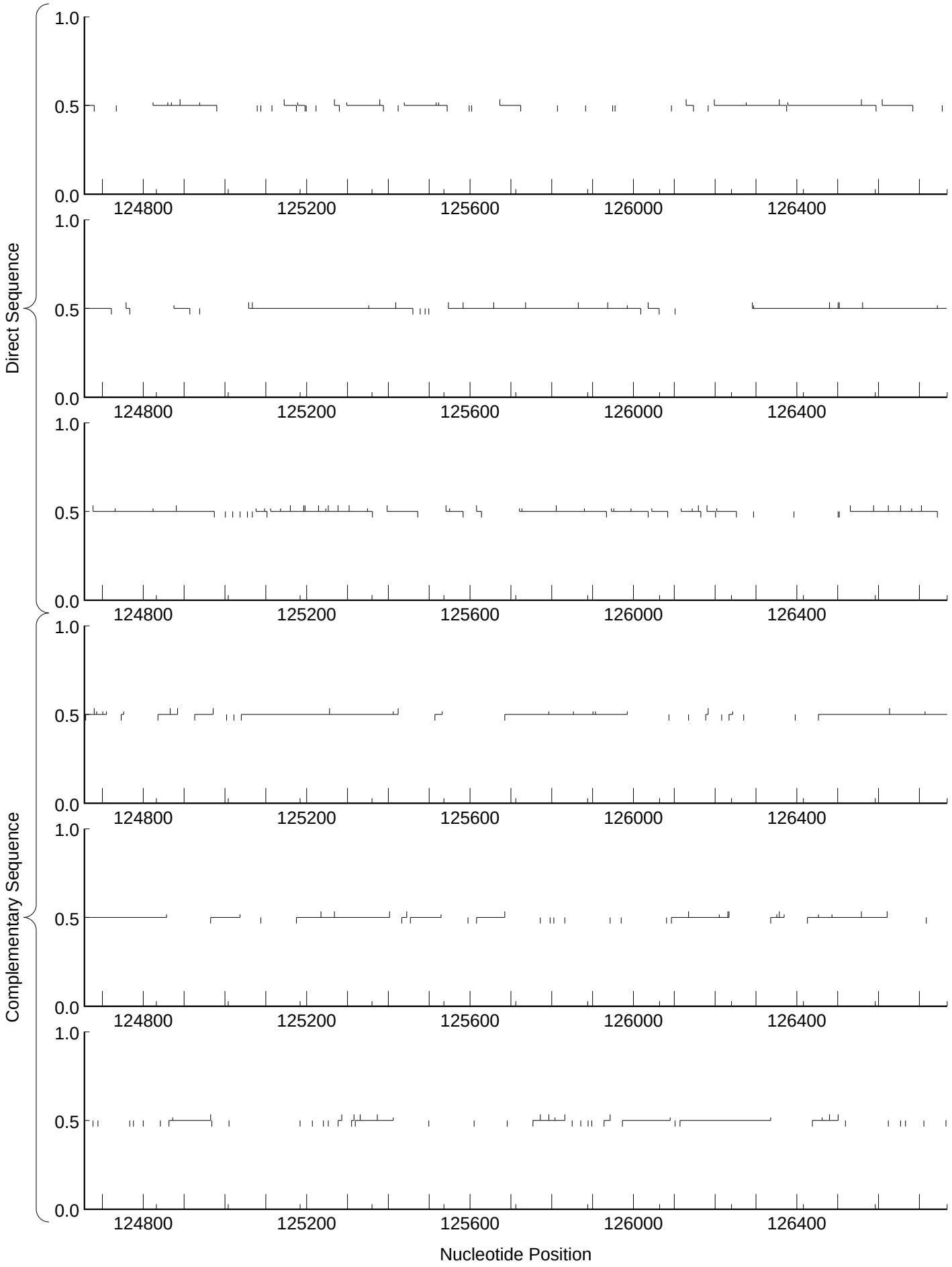
phage Sham complete sequence, 130327 bp including 787 bp terminal repeat, Cluster BK1 , Order 4, Window 96, Step 12, 59/63



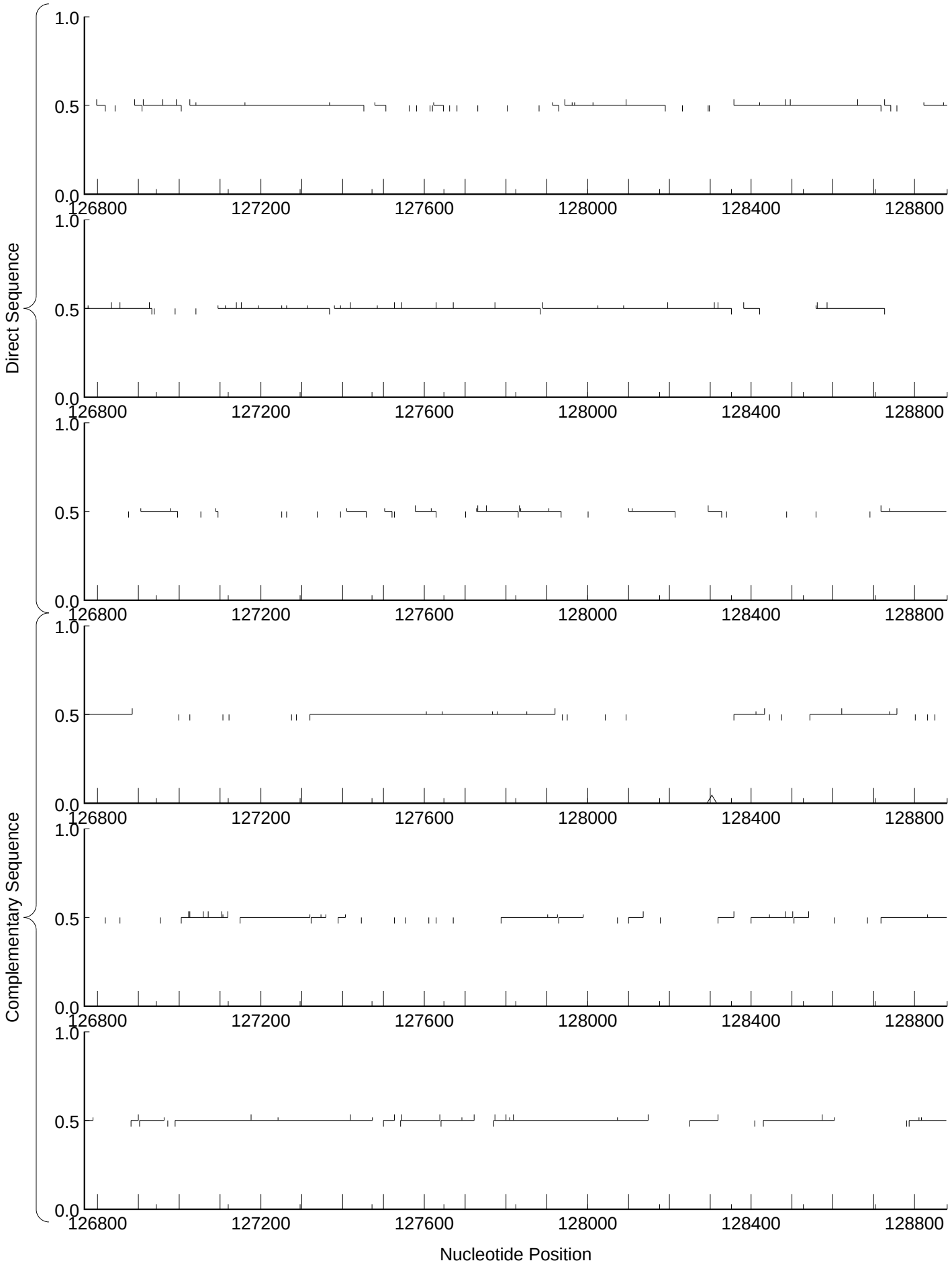
phage Sham complete sequence, 130327 bp including 787 bp terminal repeat, Cluster BK1 , Order 4, Window 96, Step 12, 60/63



phage Sham complete sequence, 130327 bp including 787 bp terminal repeat, Cluster BK1 , Order 4, Window 96, Step 12, 61/63



phage Sham complete sequence, 130327 bp including 787 bp terminal repeat, Cluster BK1 , Order 4, Window 96, Step 12, 62/63



phage Sham complete sequence, 130327 bp including 787 bp terminal repeat, Cluster BK1 , Order 4, Window 96, Step 12, 63/63

