GeneMark

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

PROGRAM INFORMATION

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Sequence         :  Streptomyces phage Gilson complete sequence, 128338 bp including 766 bp terminal repeat, Cl
Analysis Date    :  3/28/24 at 12:02:43
Pages            :  62
Sequence Length  :  128338 bp
GC Content       :   47.44%

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

PS-Version       :  1.2

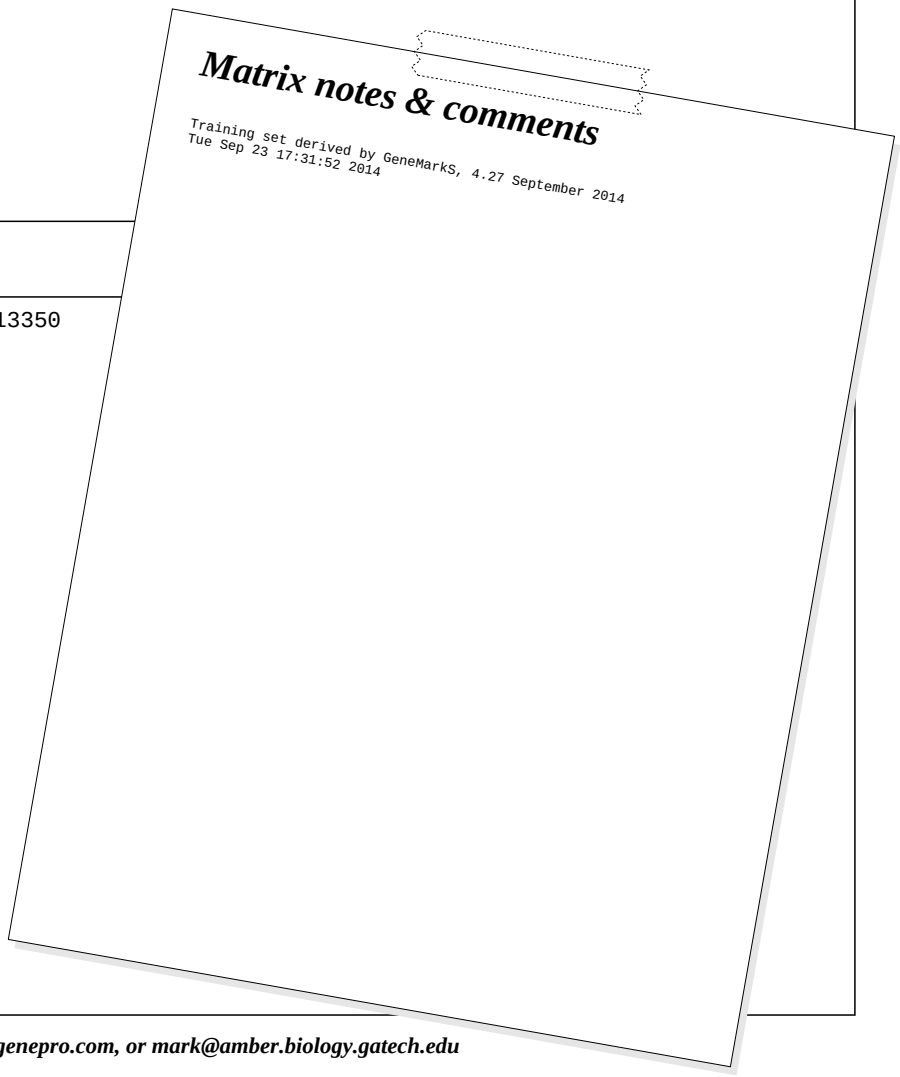
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MATRIX INFORMATION

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Author     :  -

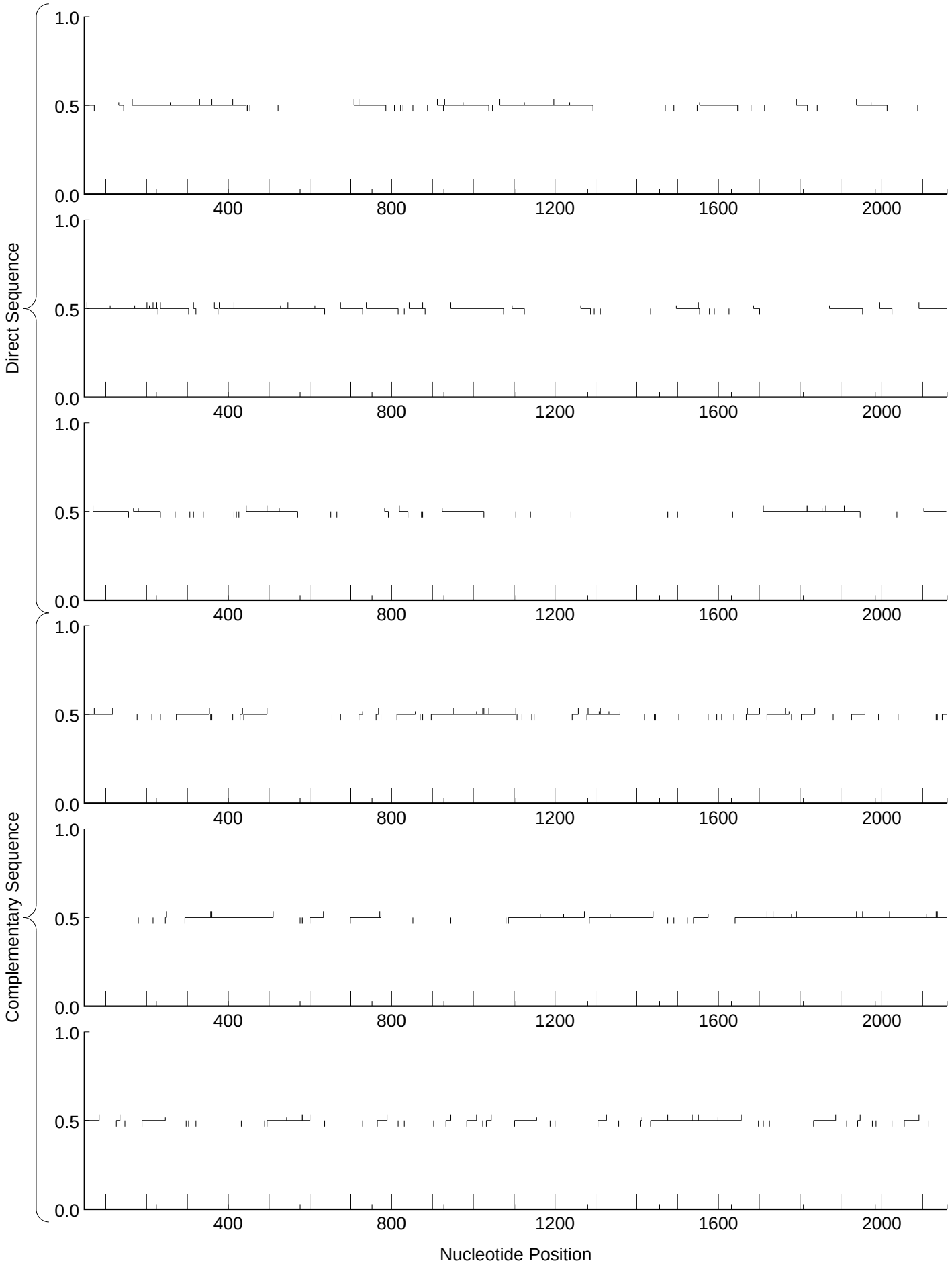
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  Send questions / comments to:
      Dr. M. Borodovsky
      Georgia Institute of Technology
      School of Biology
      Atlanta, GA  30332-0230
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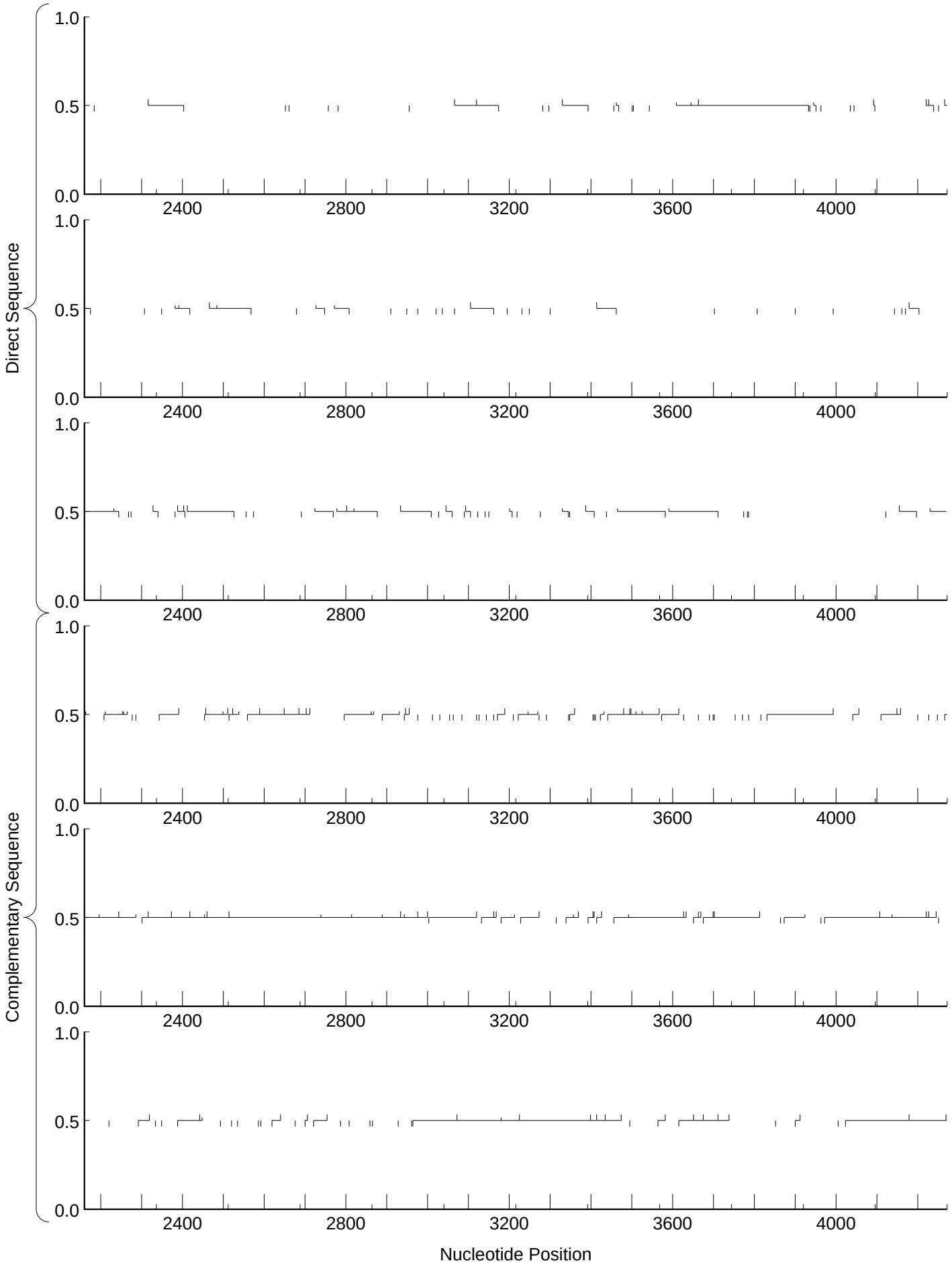


For help or information, send electronic mail to: custserv@genepro.com, or mark@amber.biology.gatech.edu

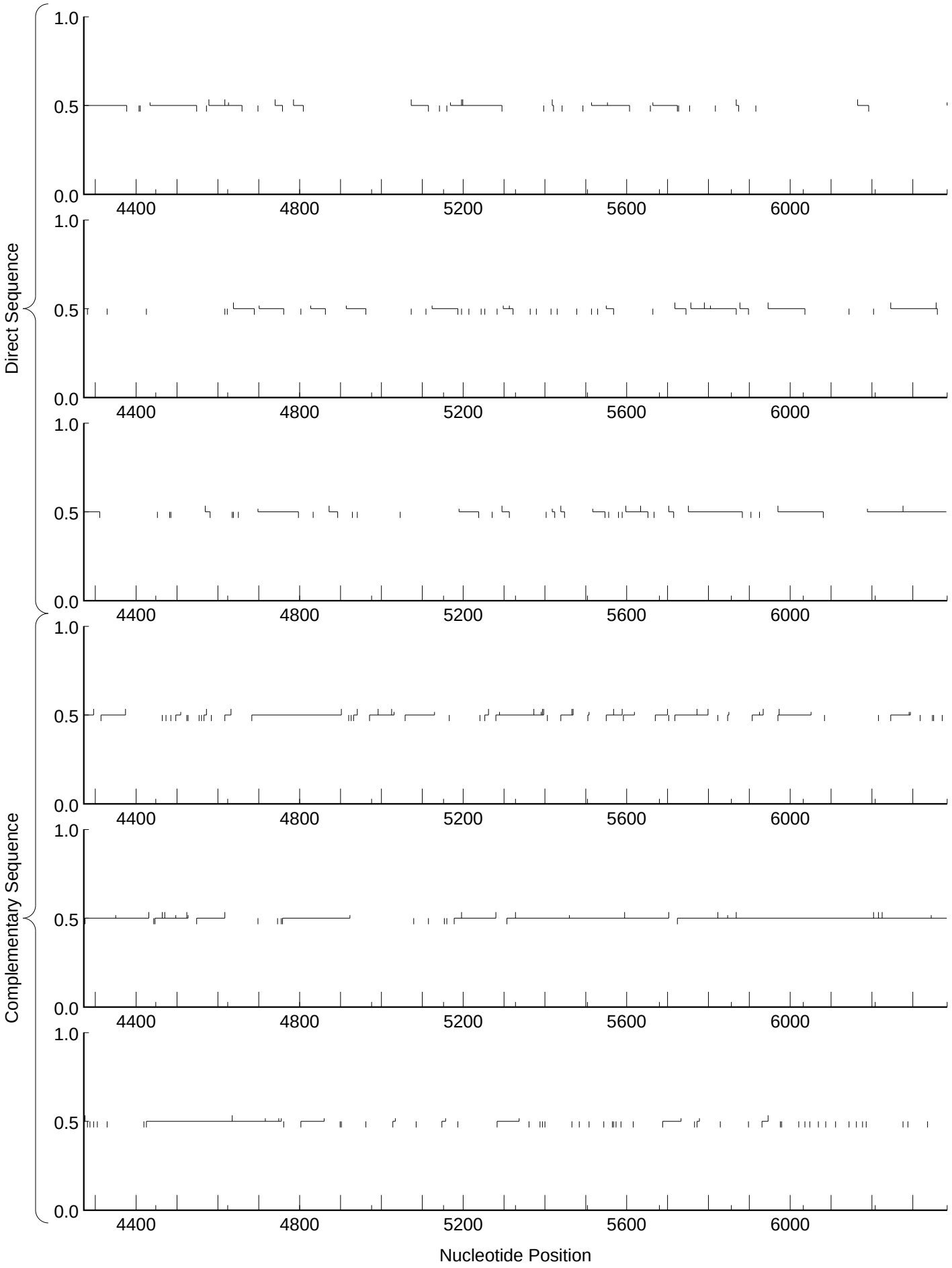
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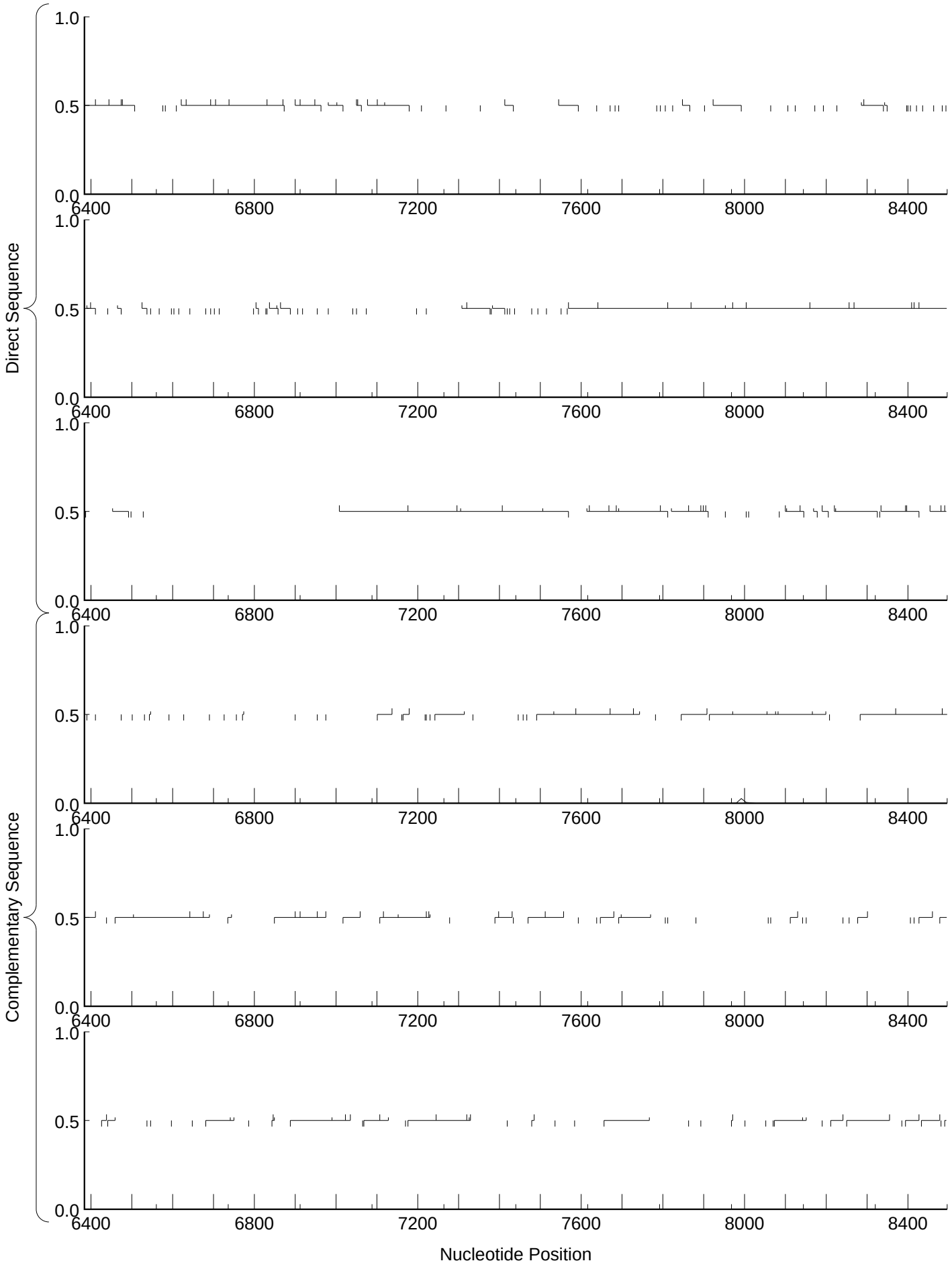
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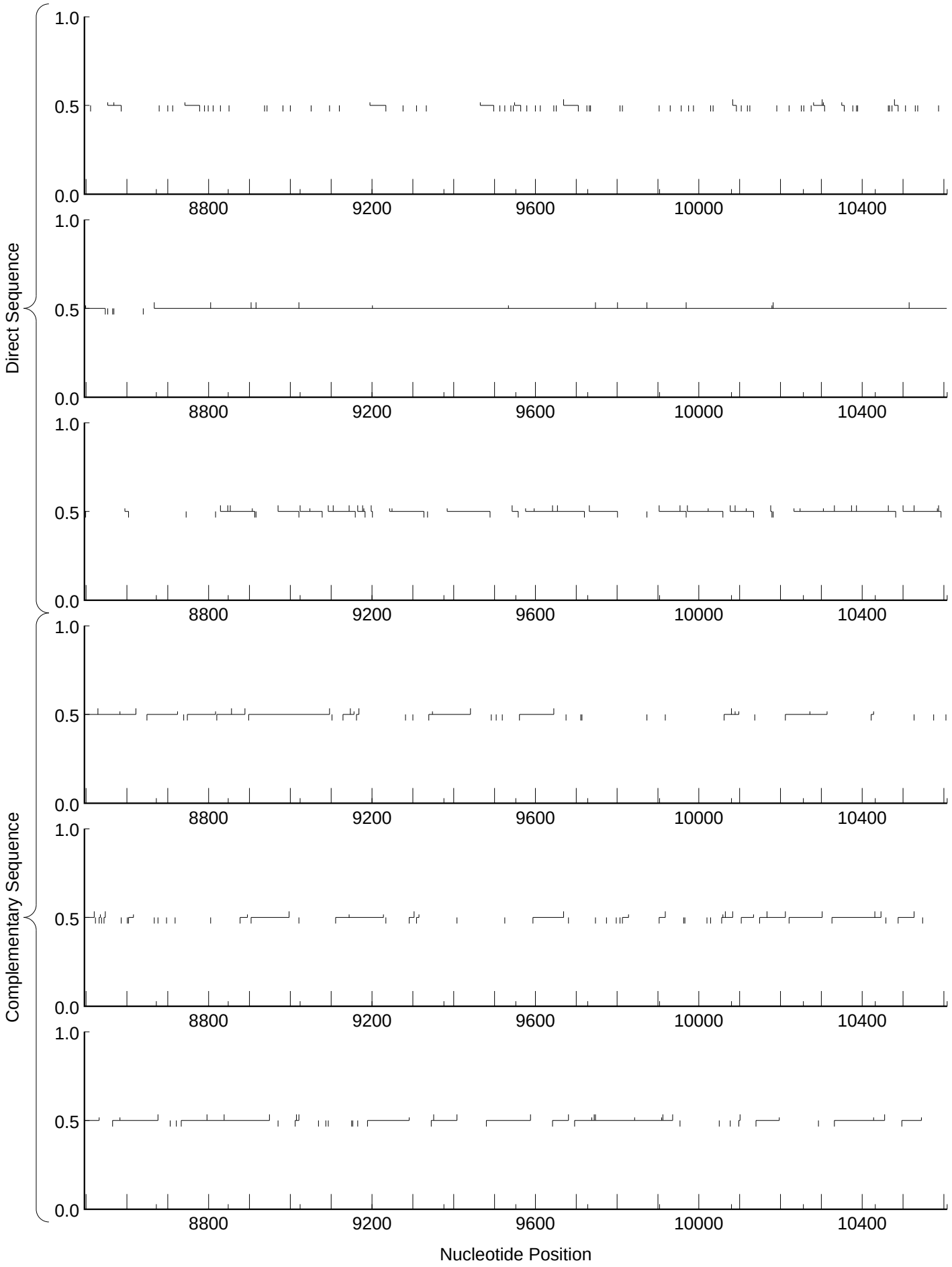
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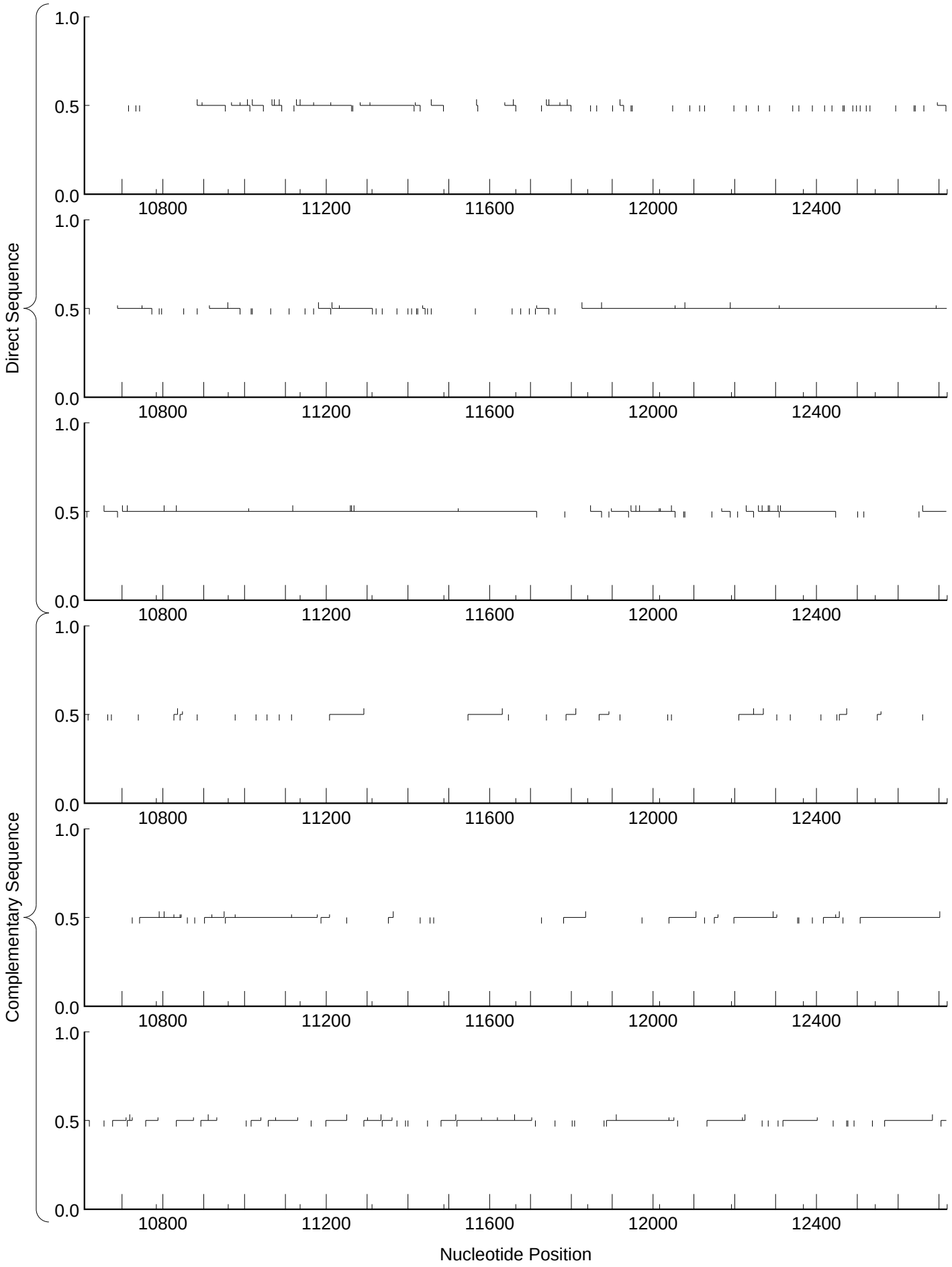
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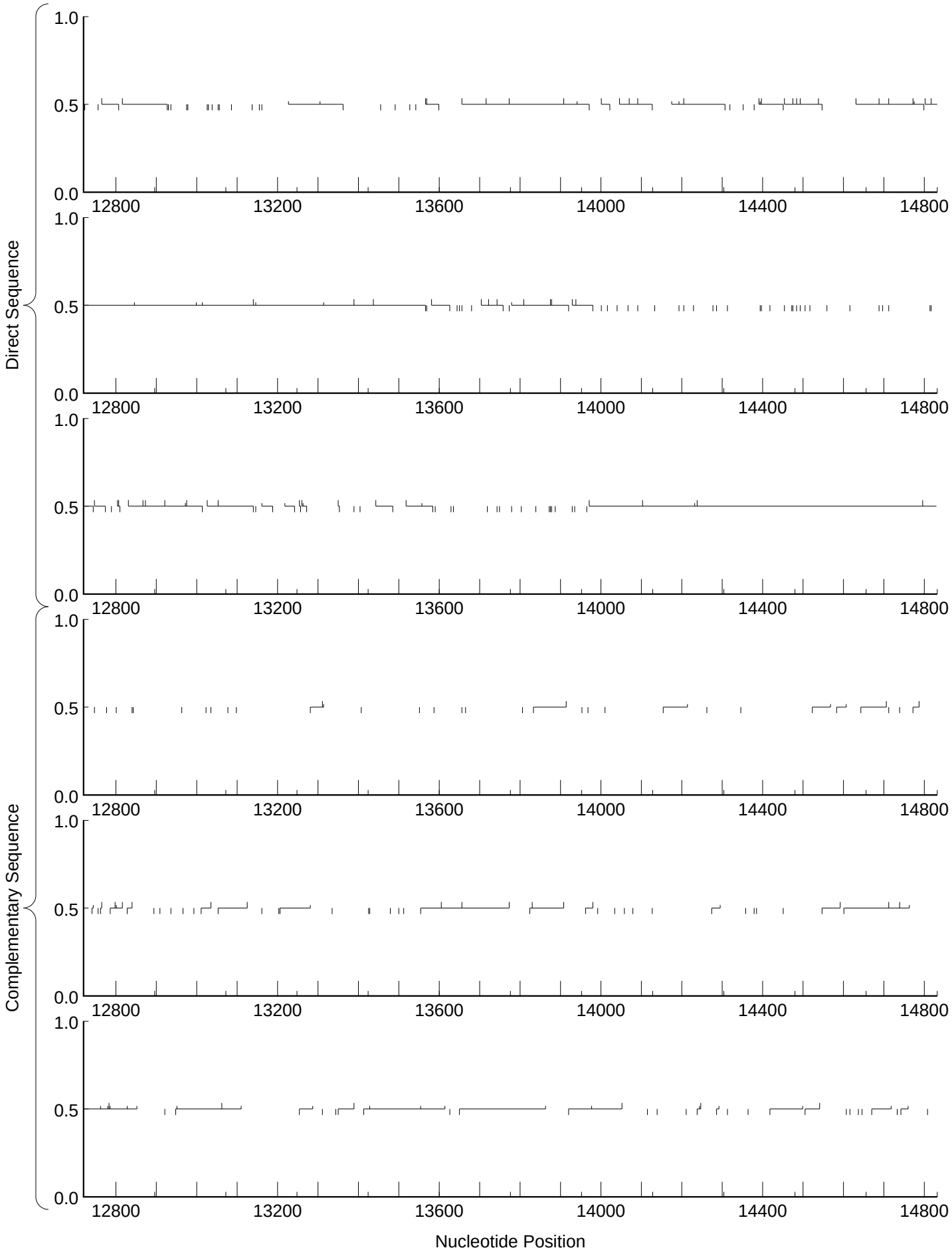
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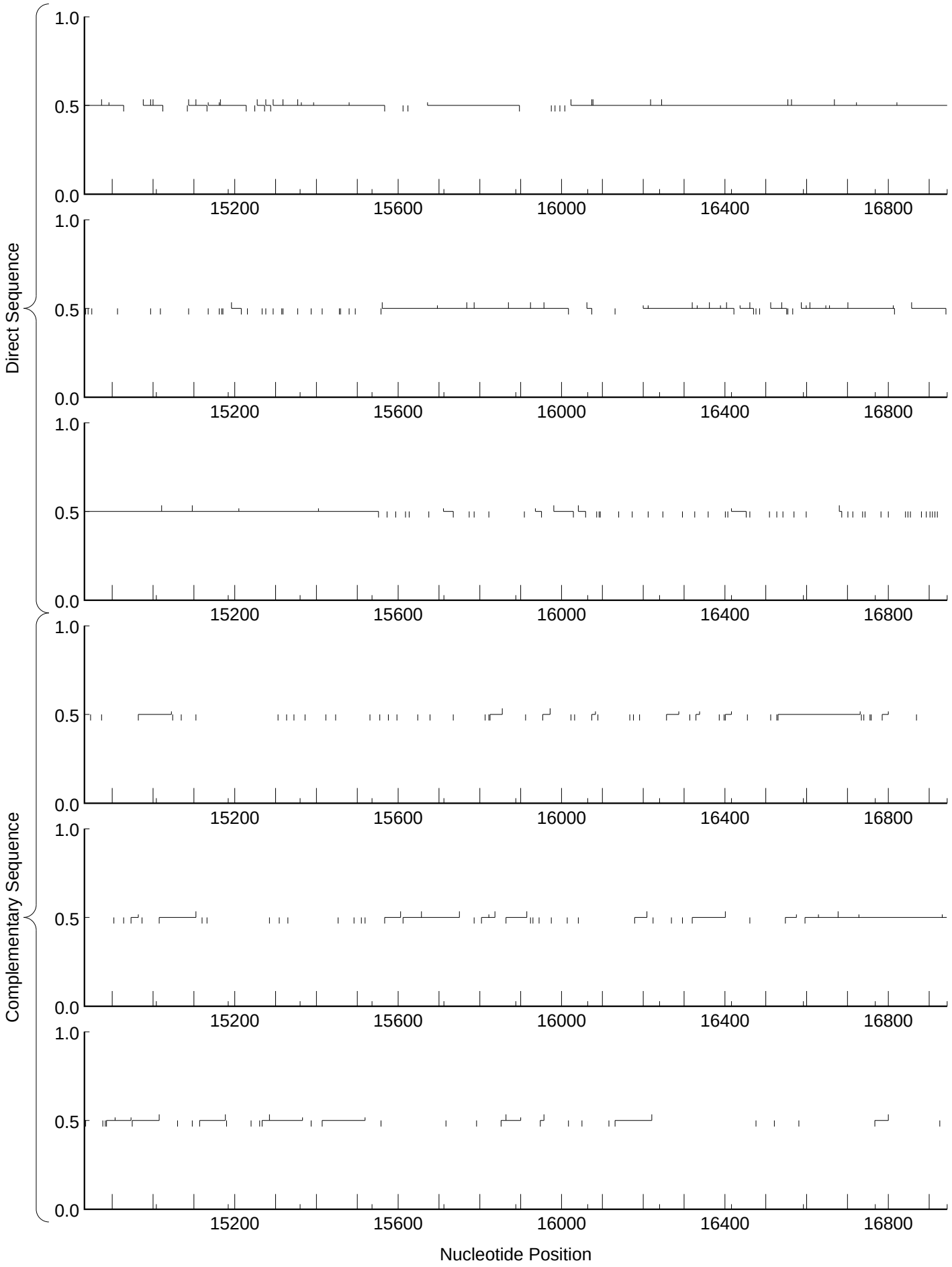
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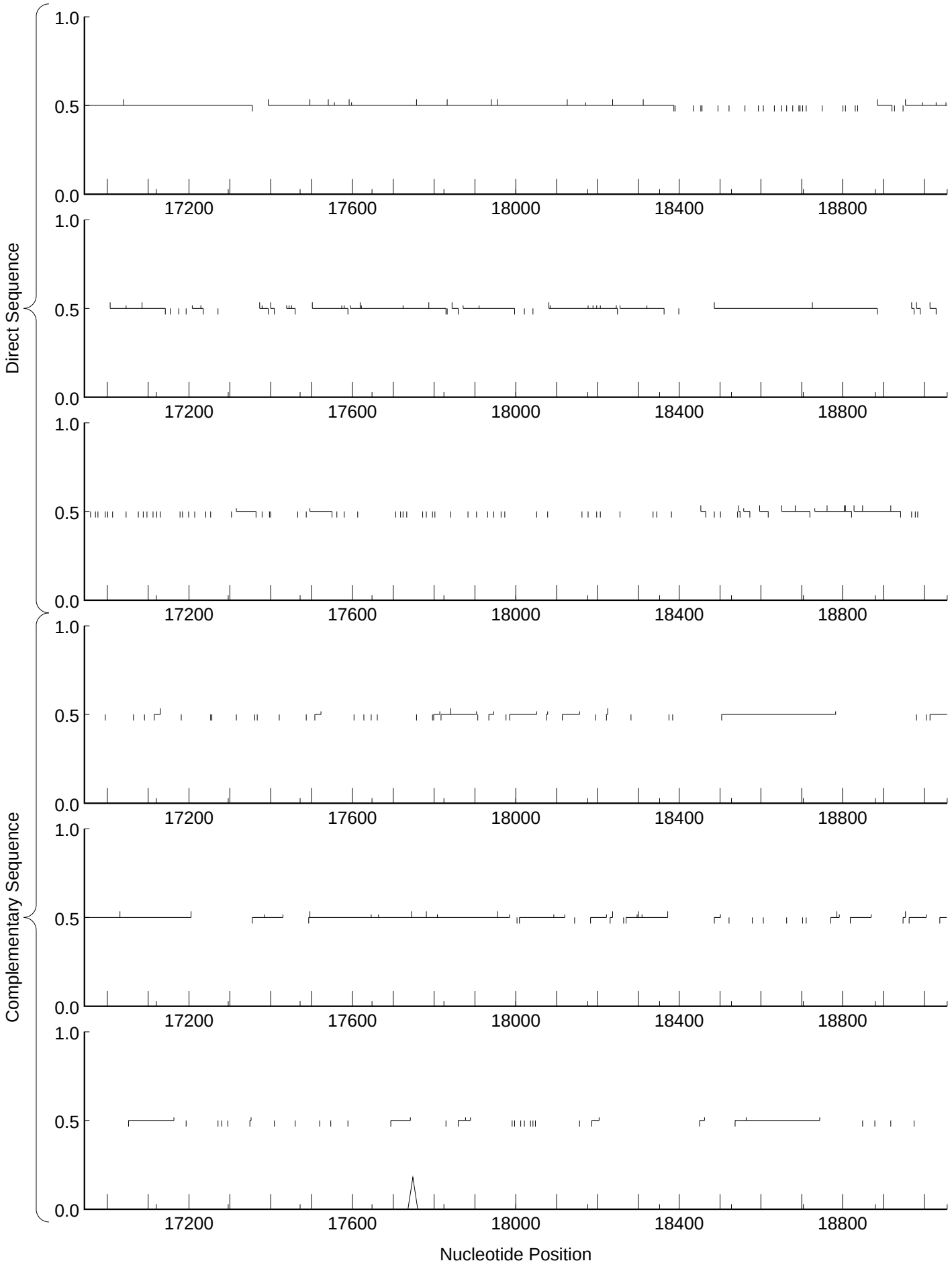
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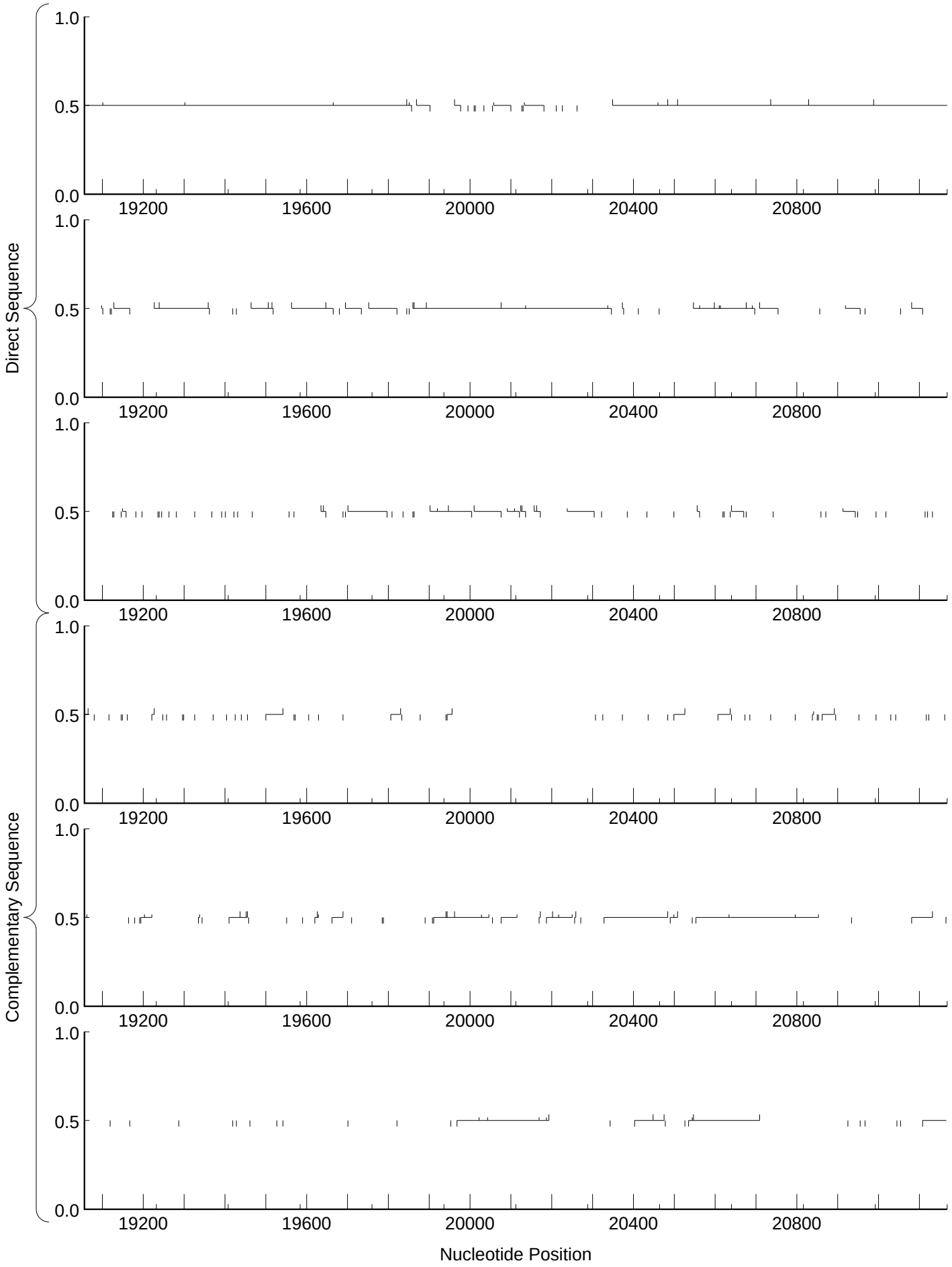
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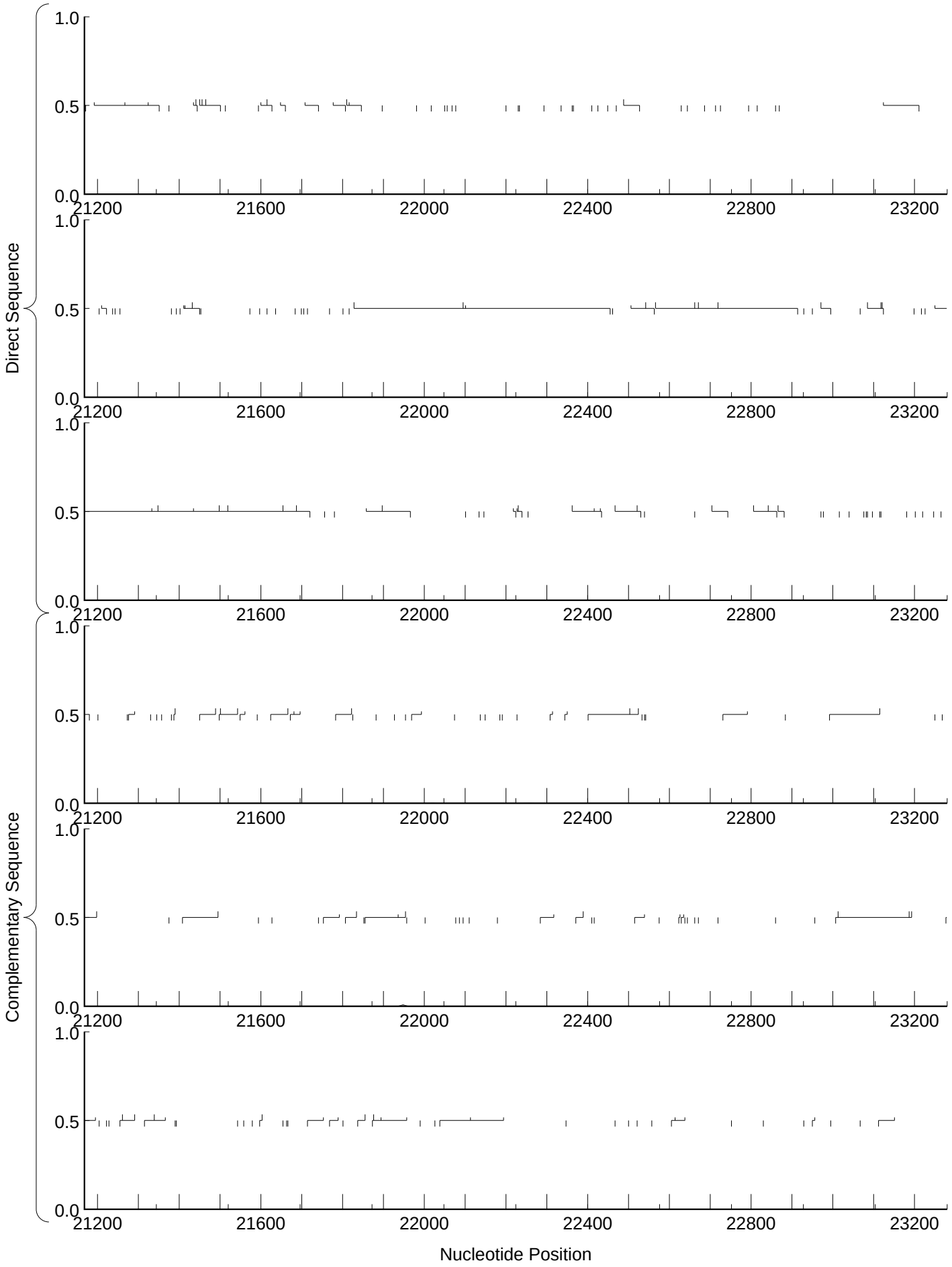
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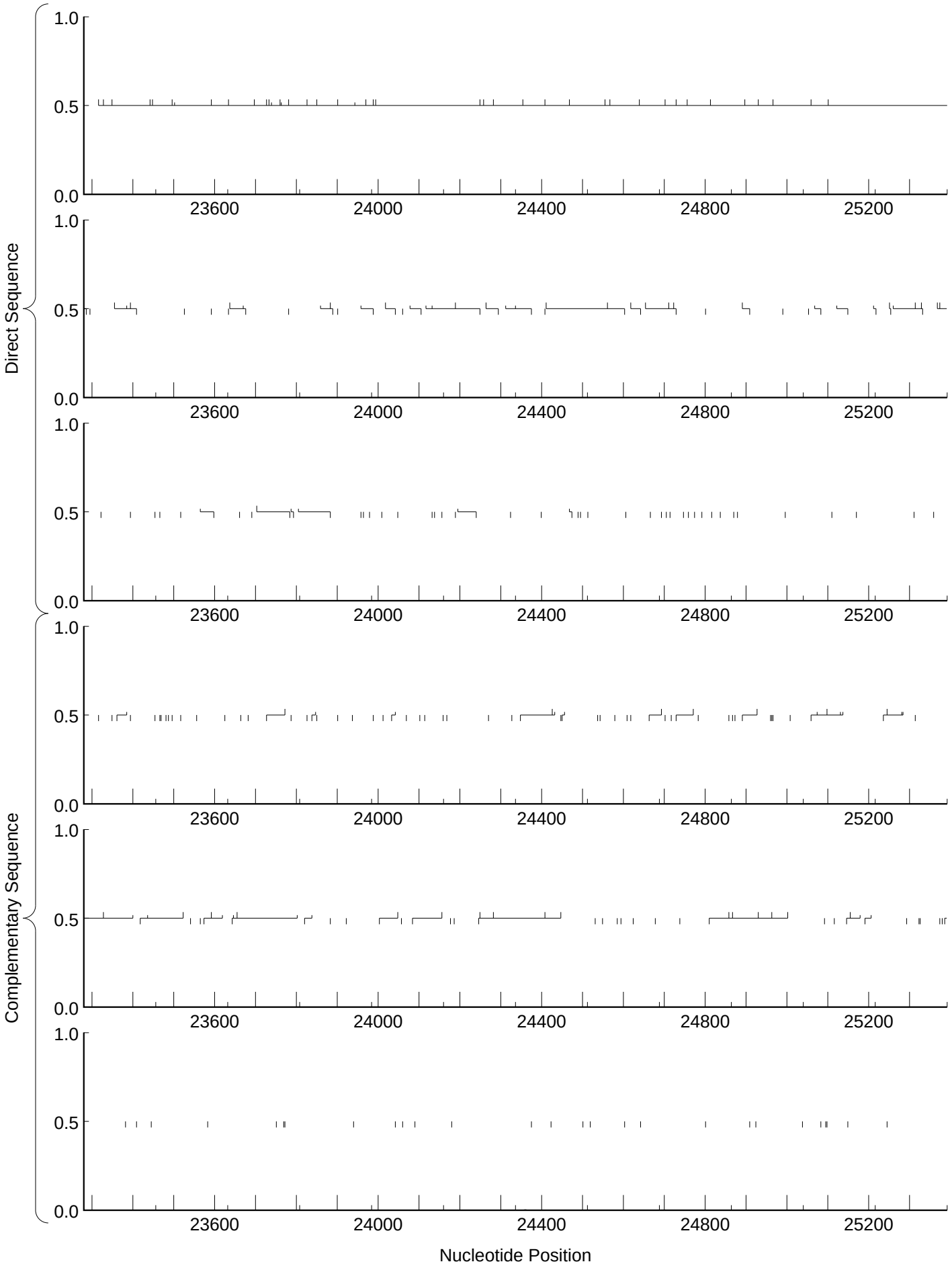
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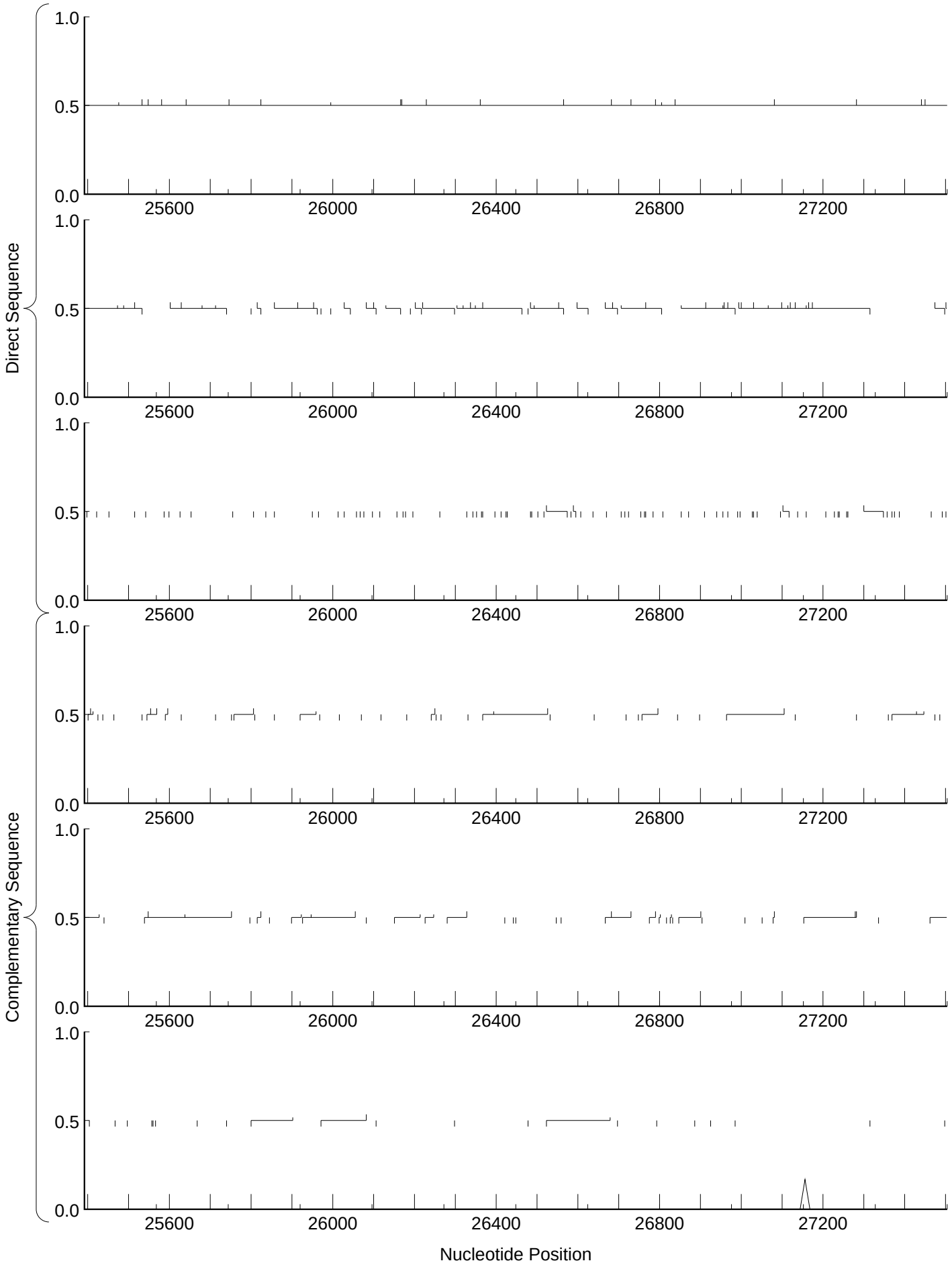
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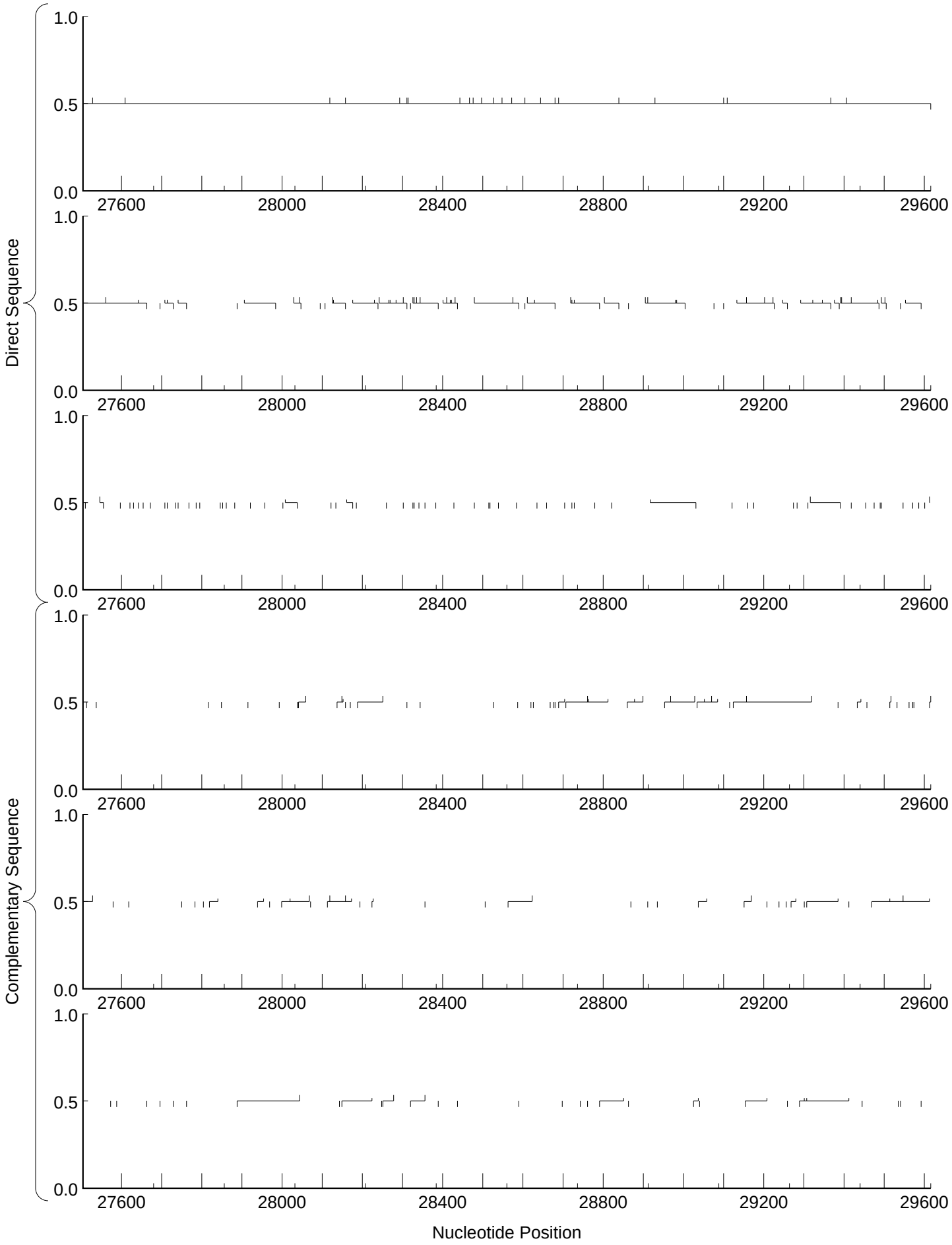
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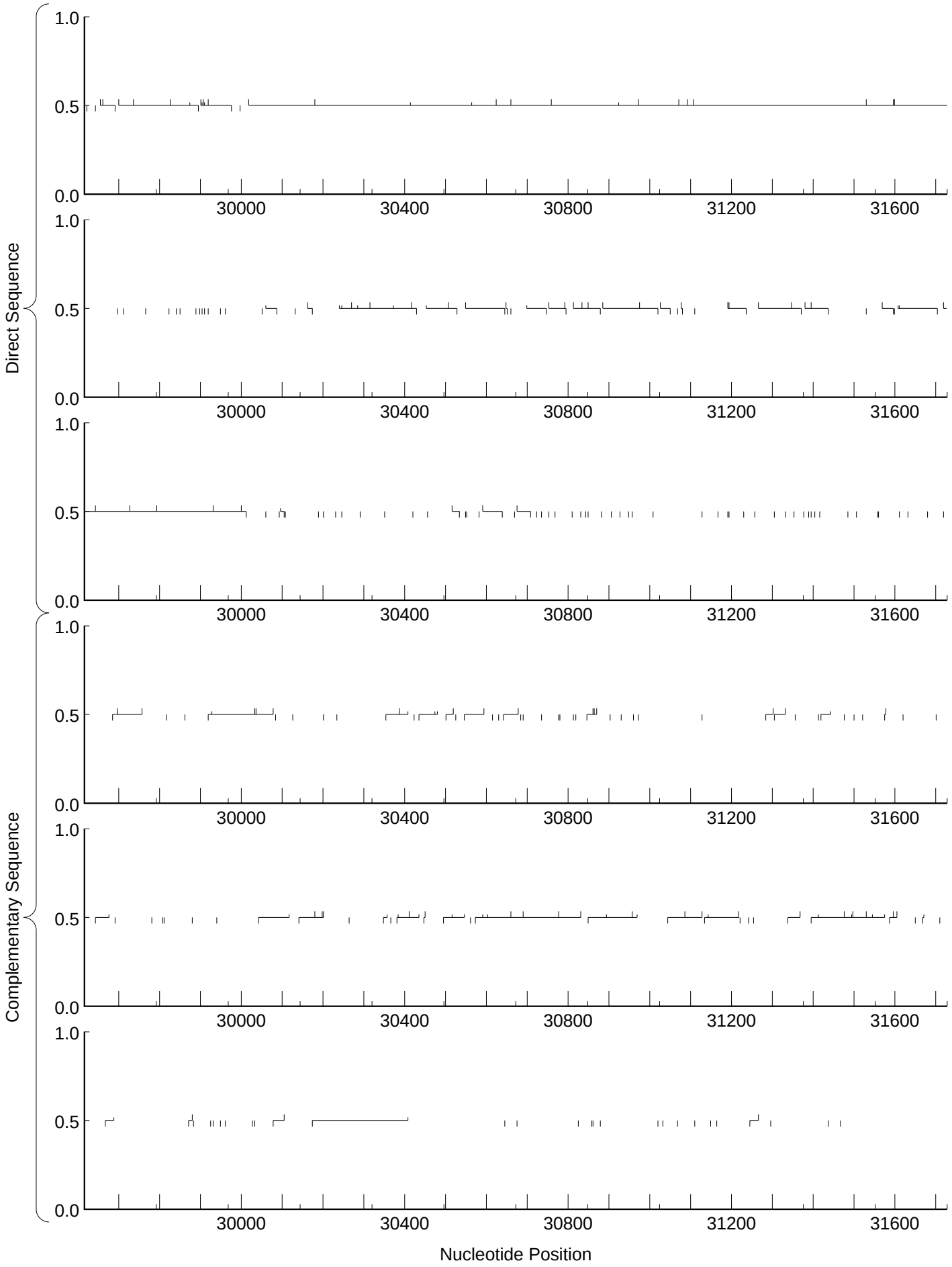
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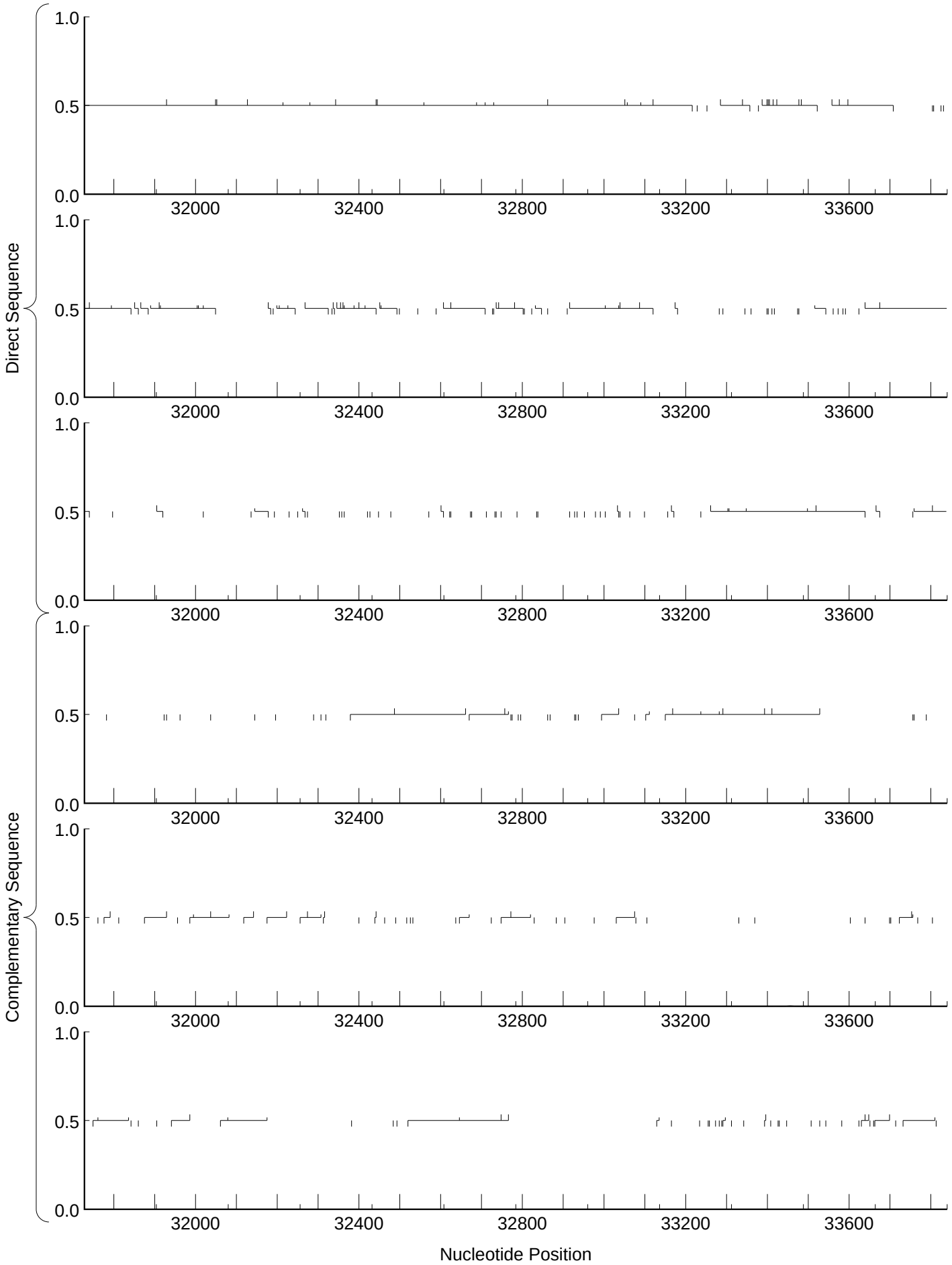
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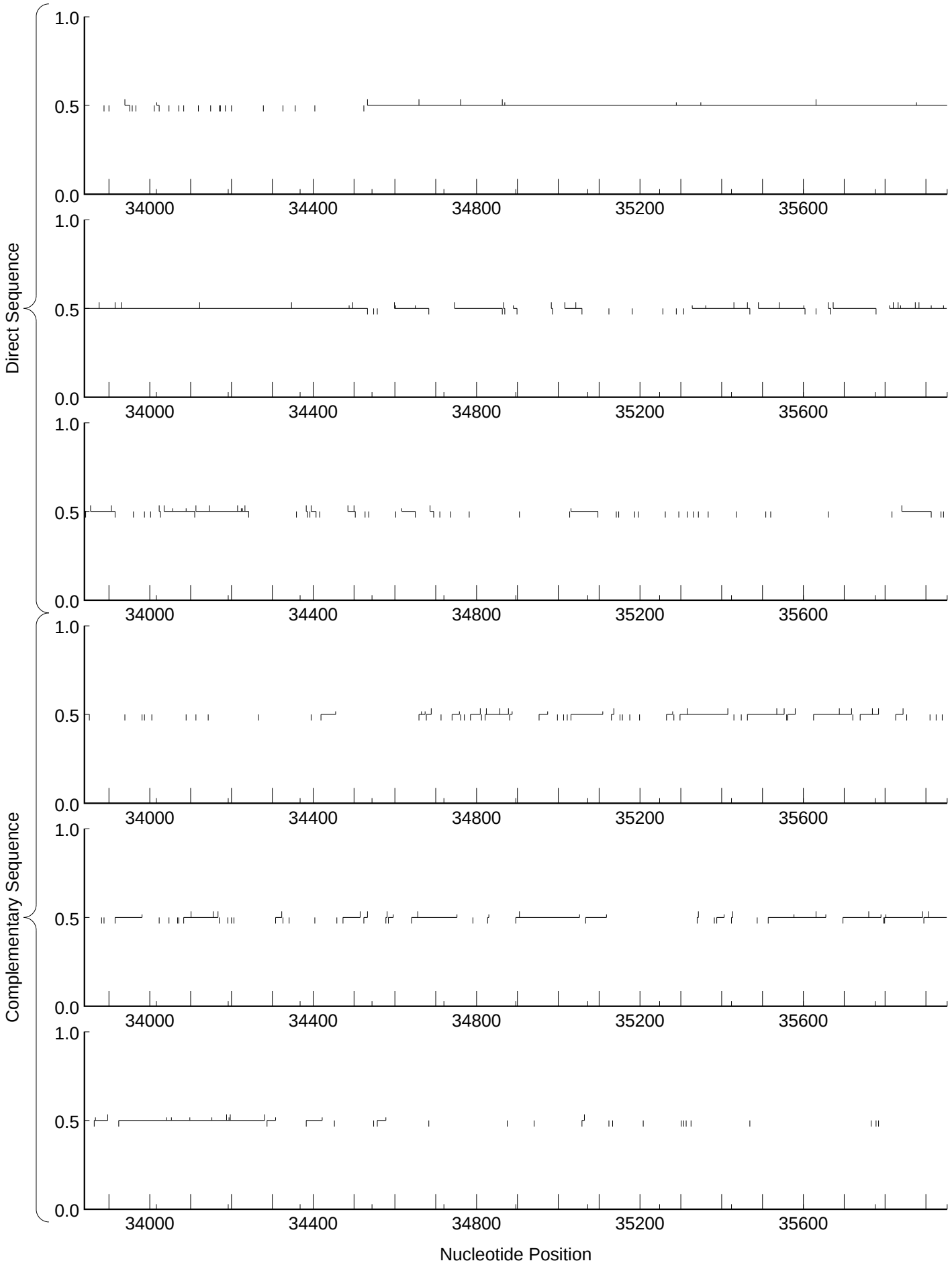
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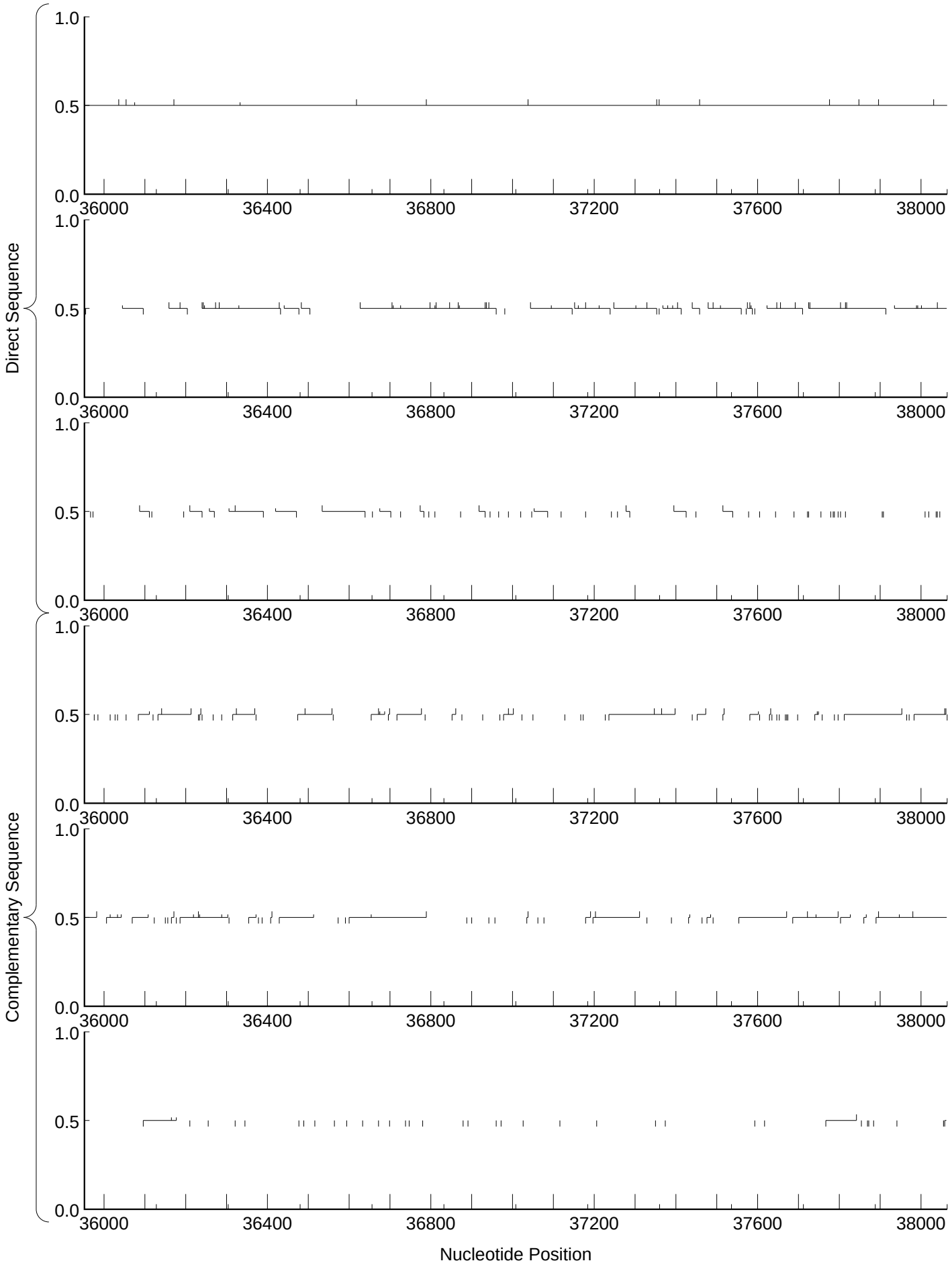
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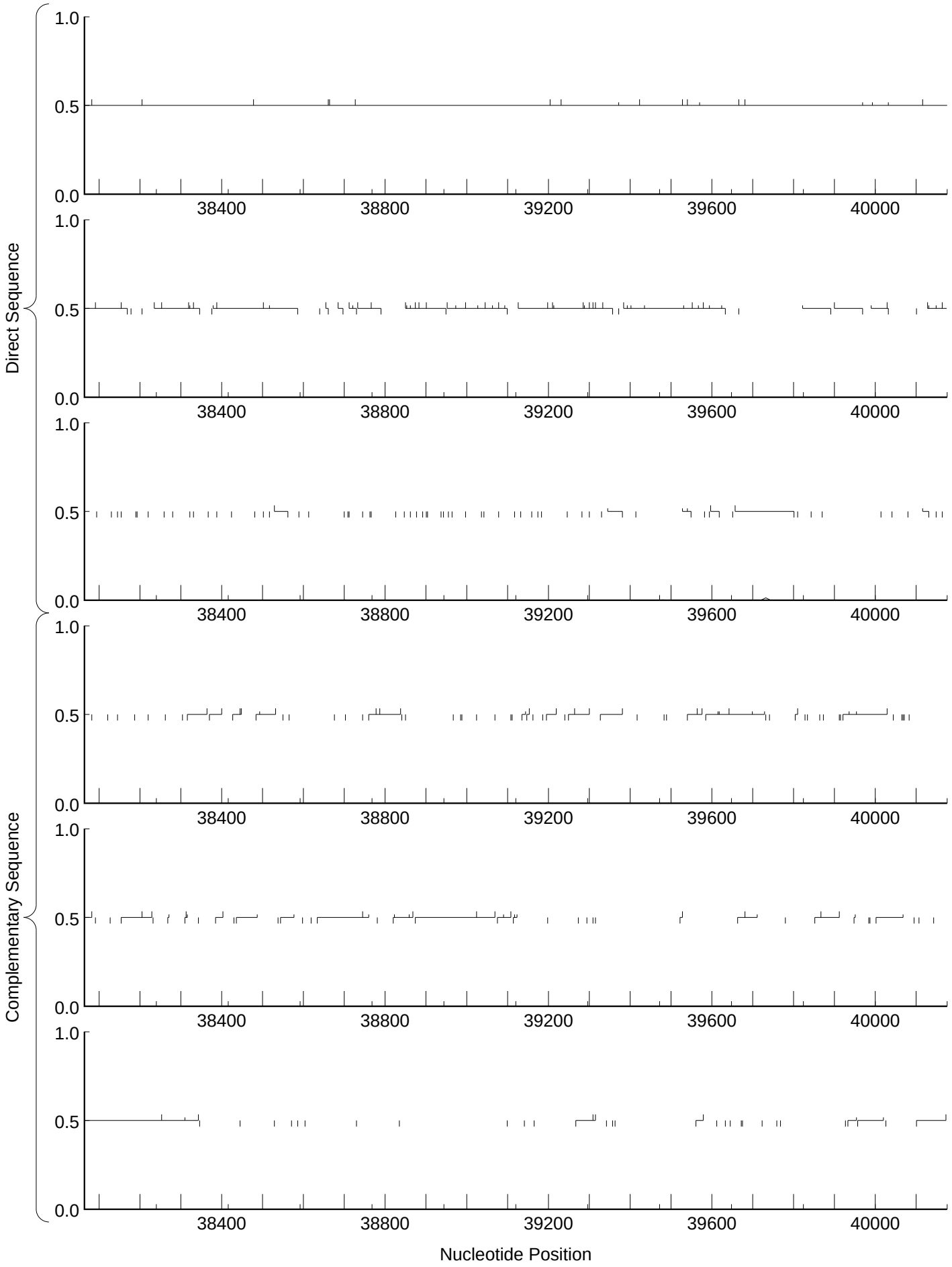
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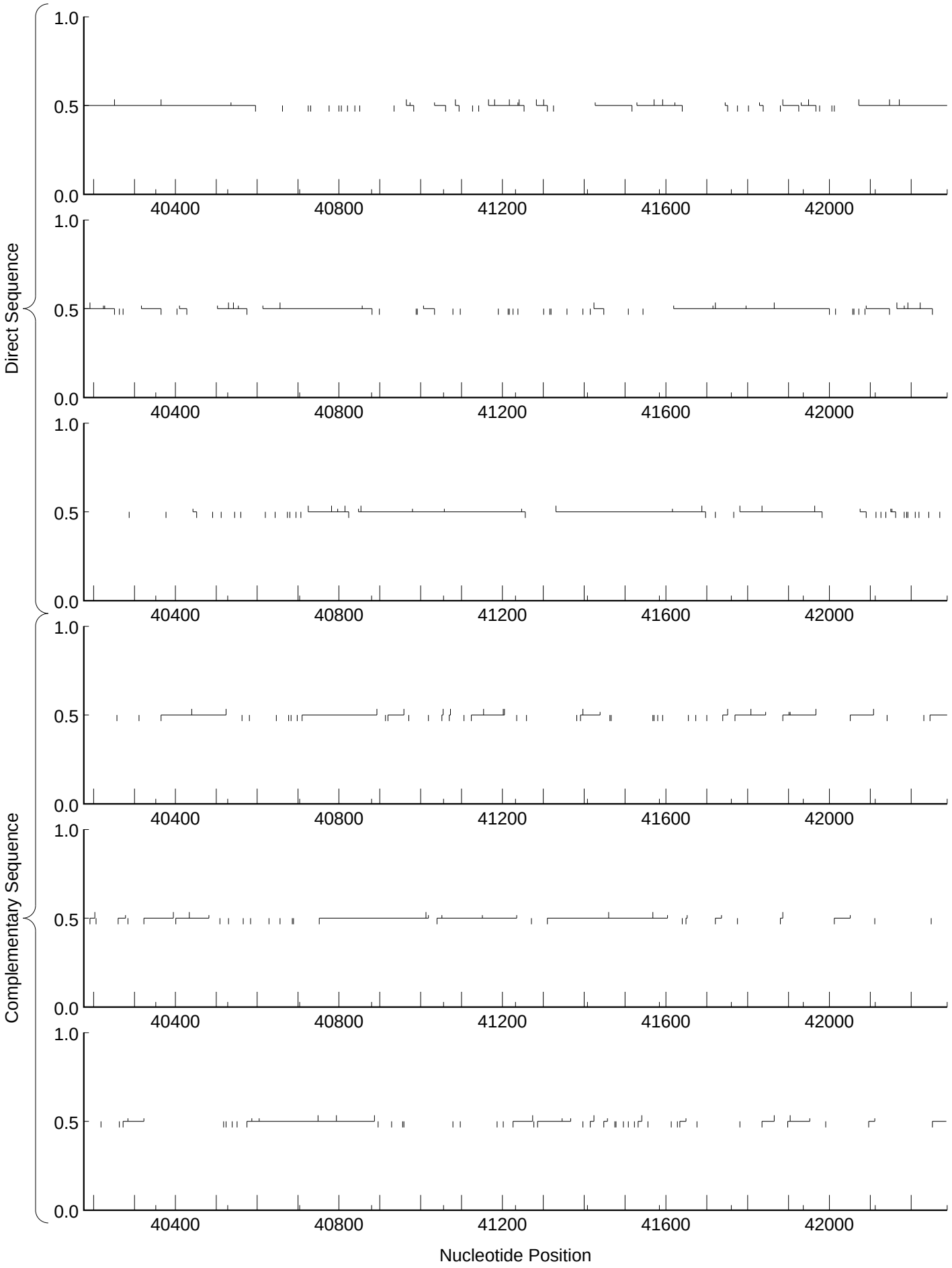
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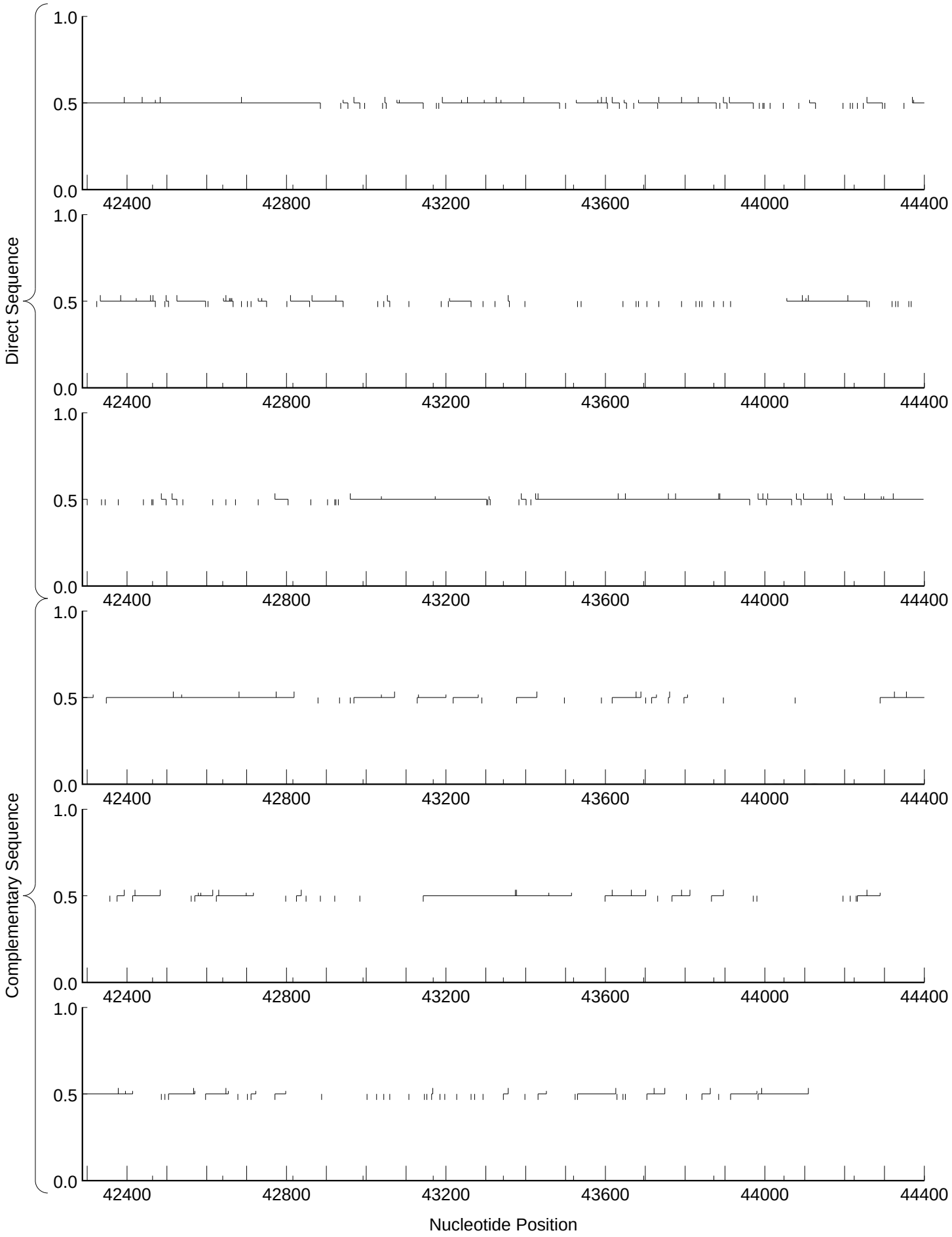
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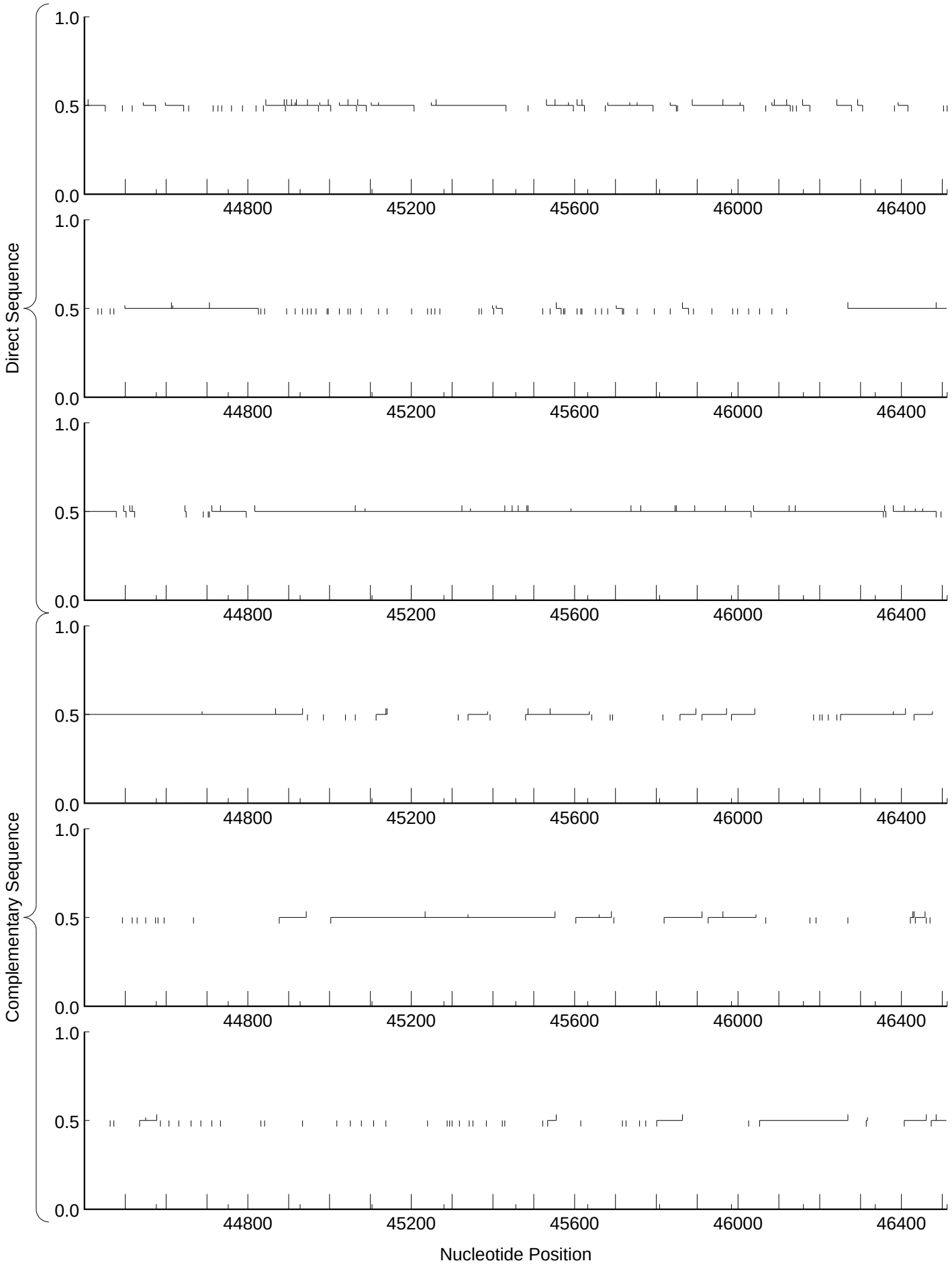
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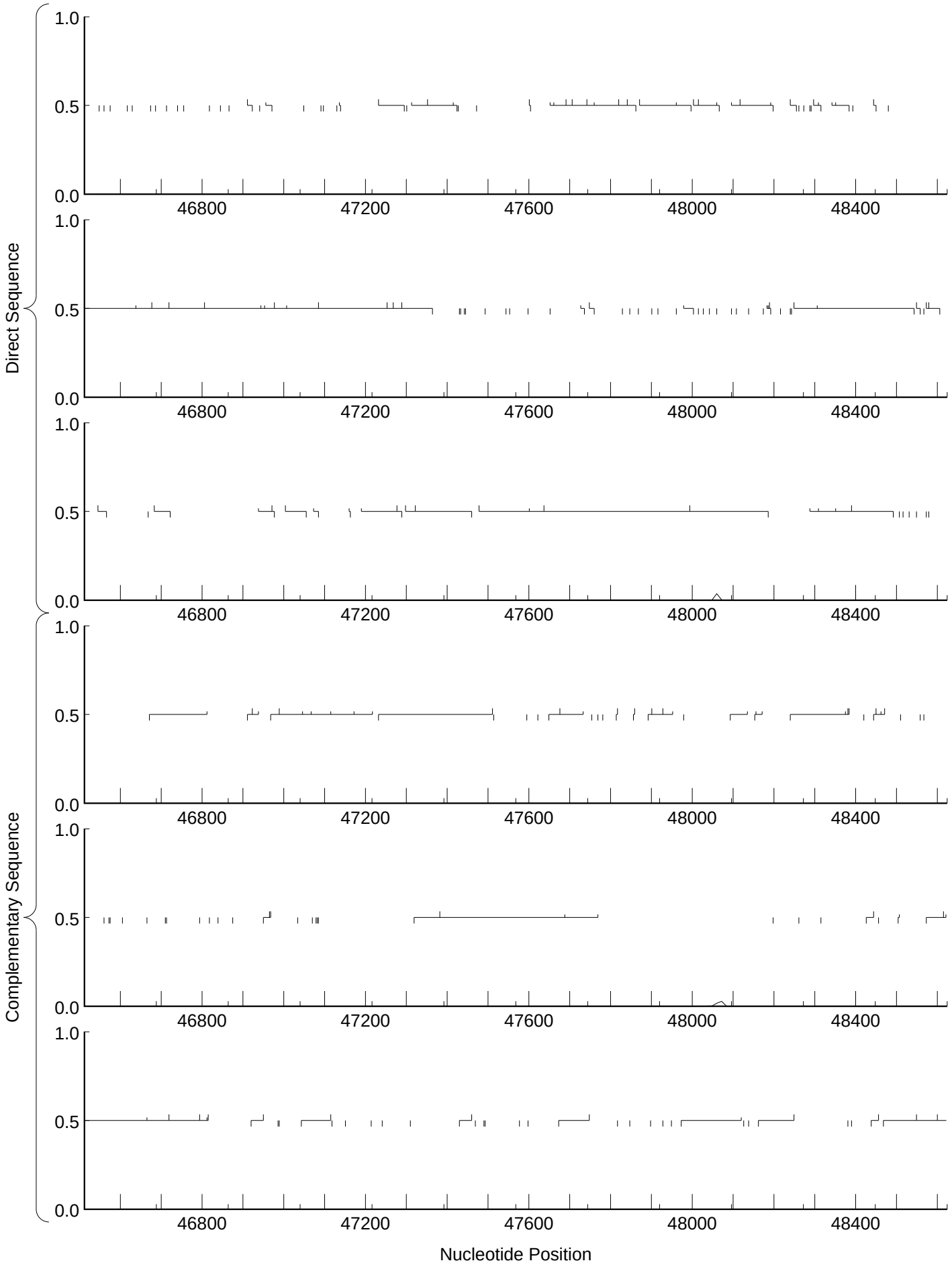
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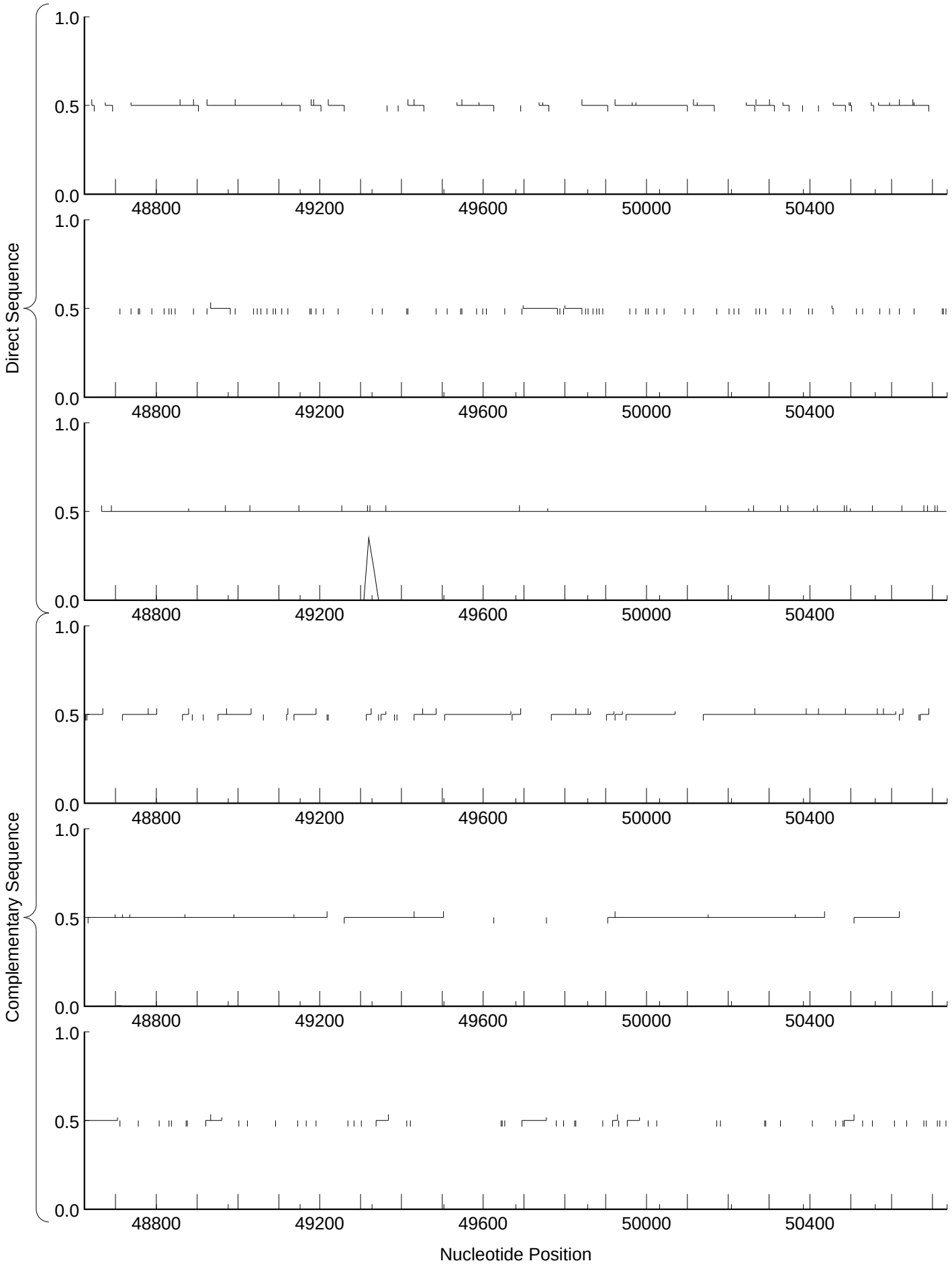
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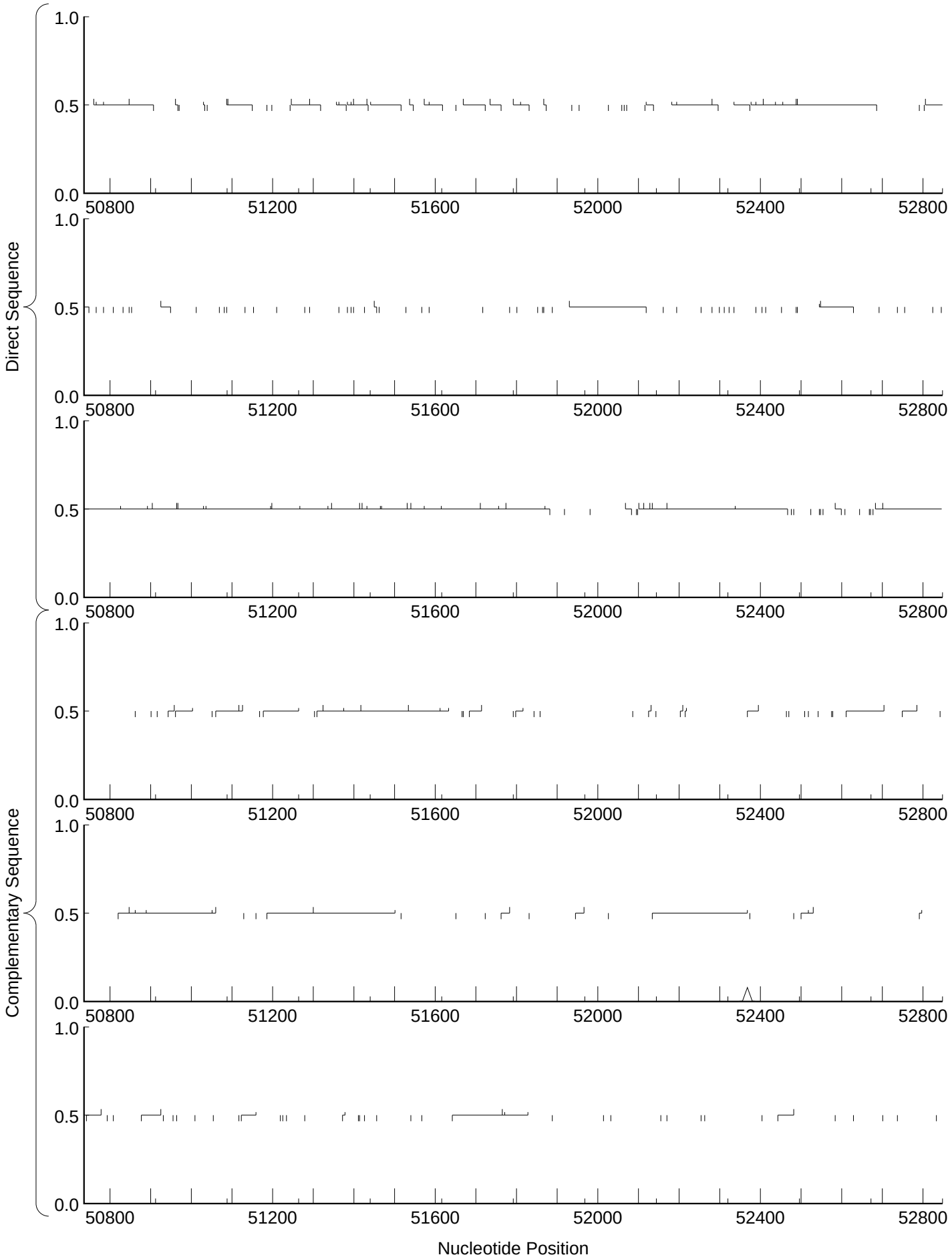
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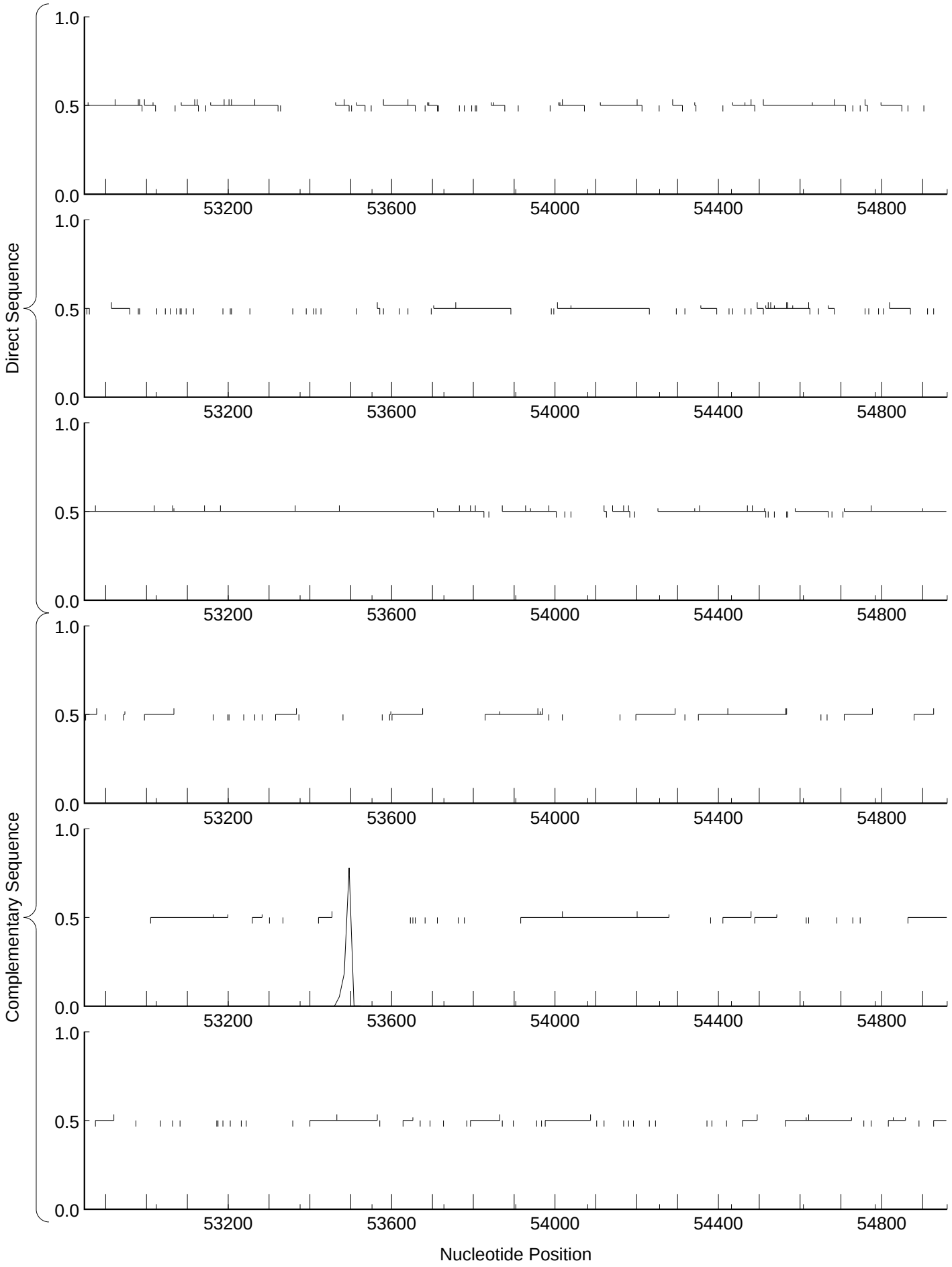
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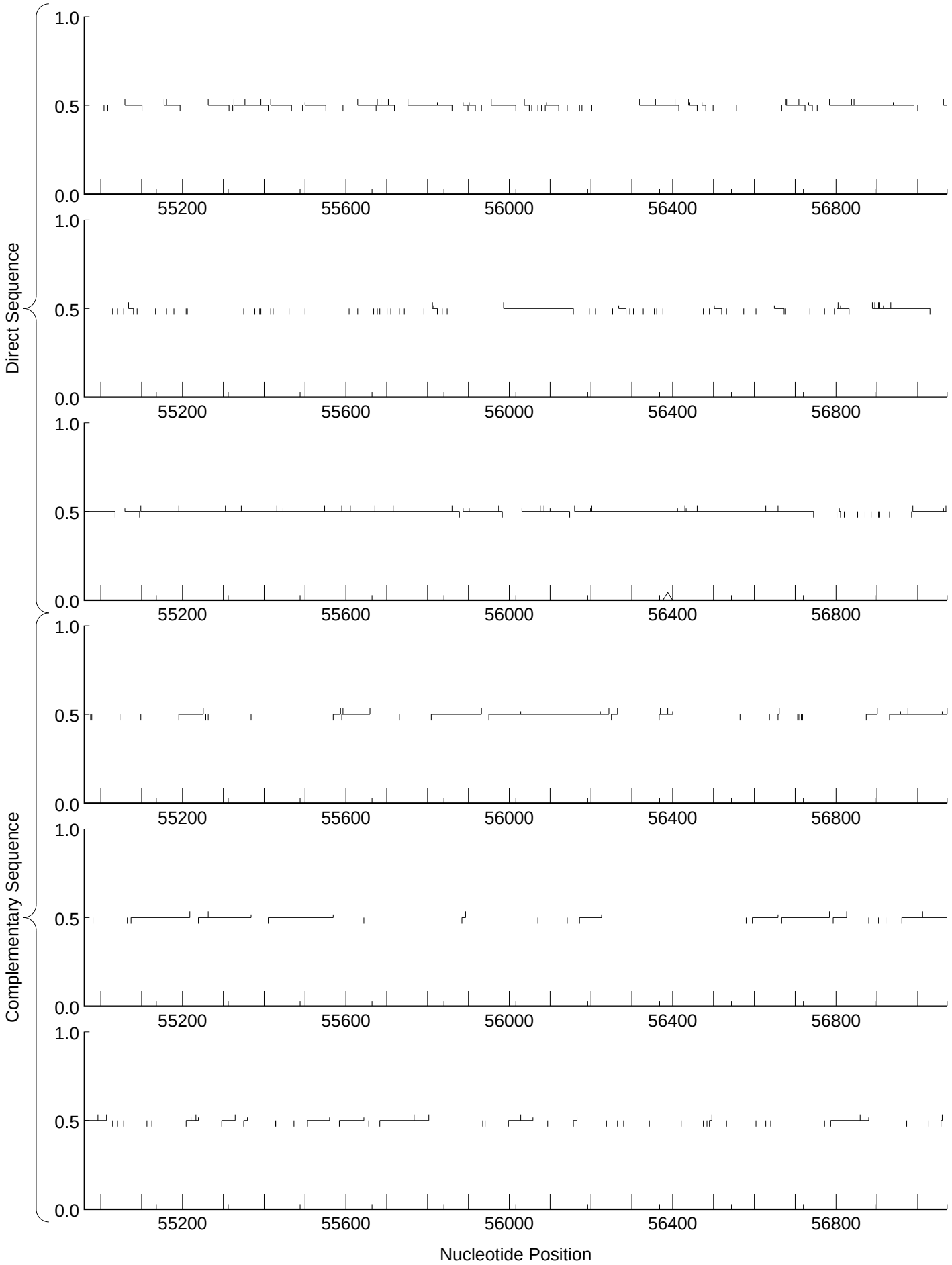
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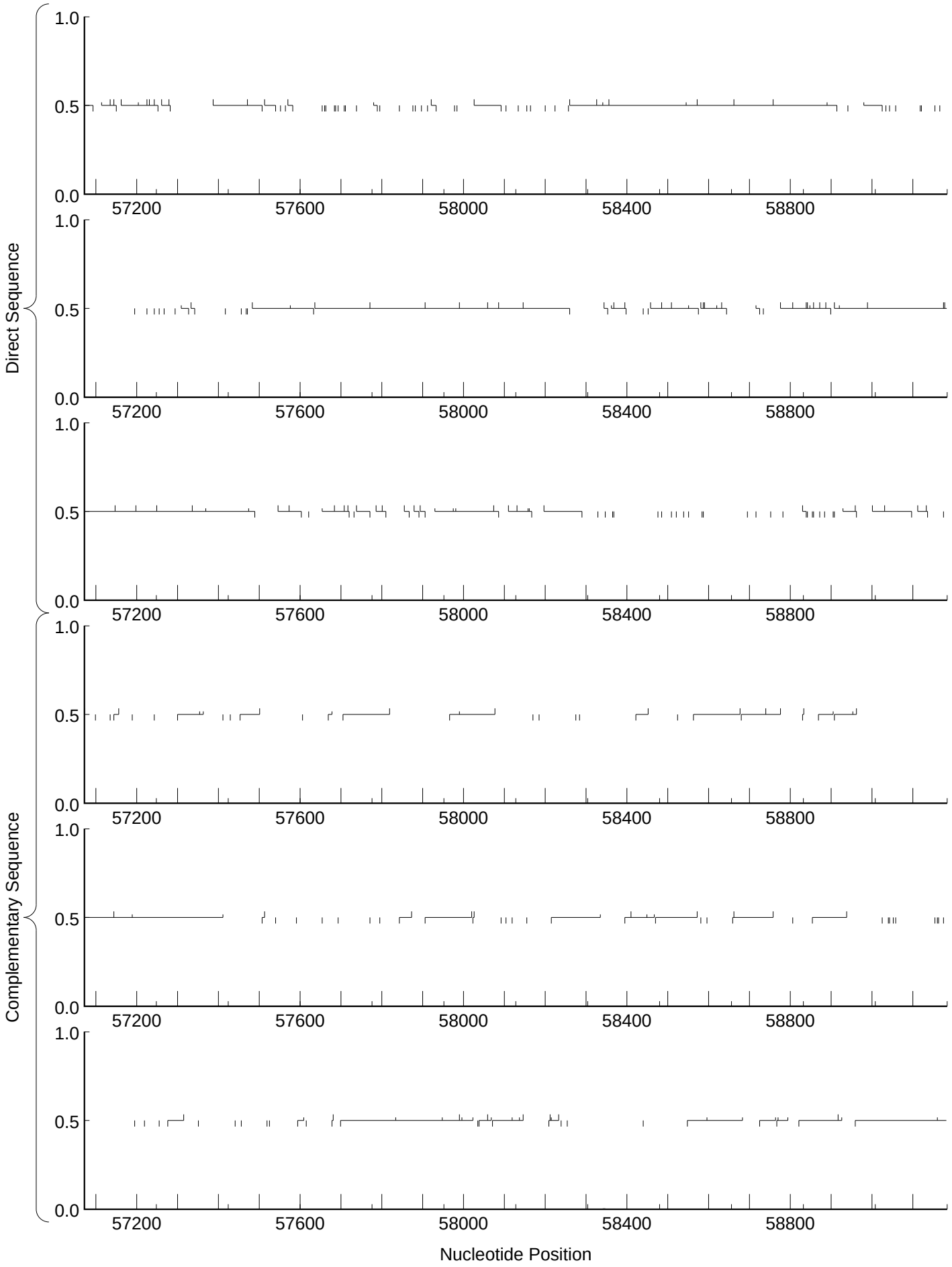
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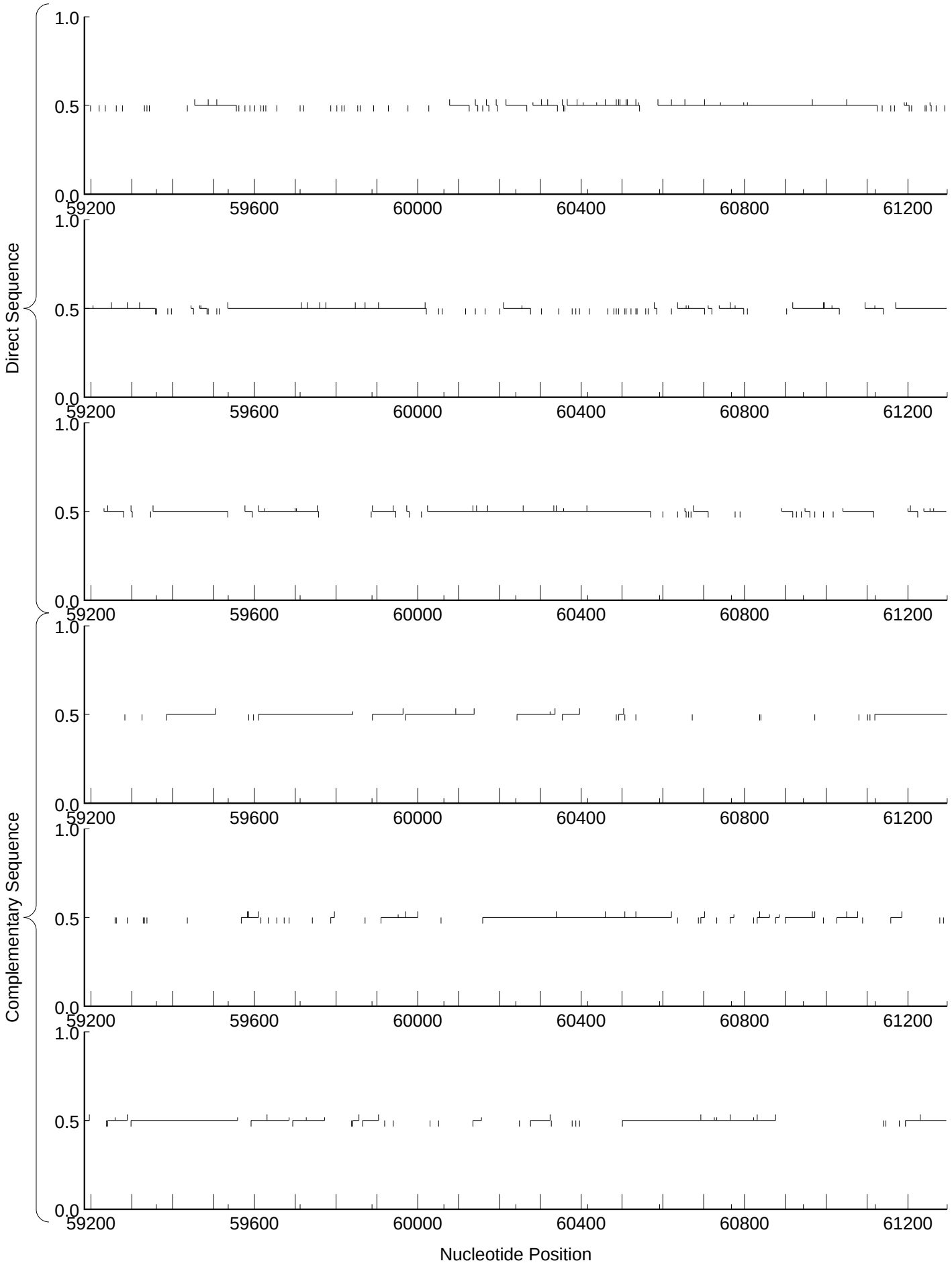
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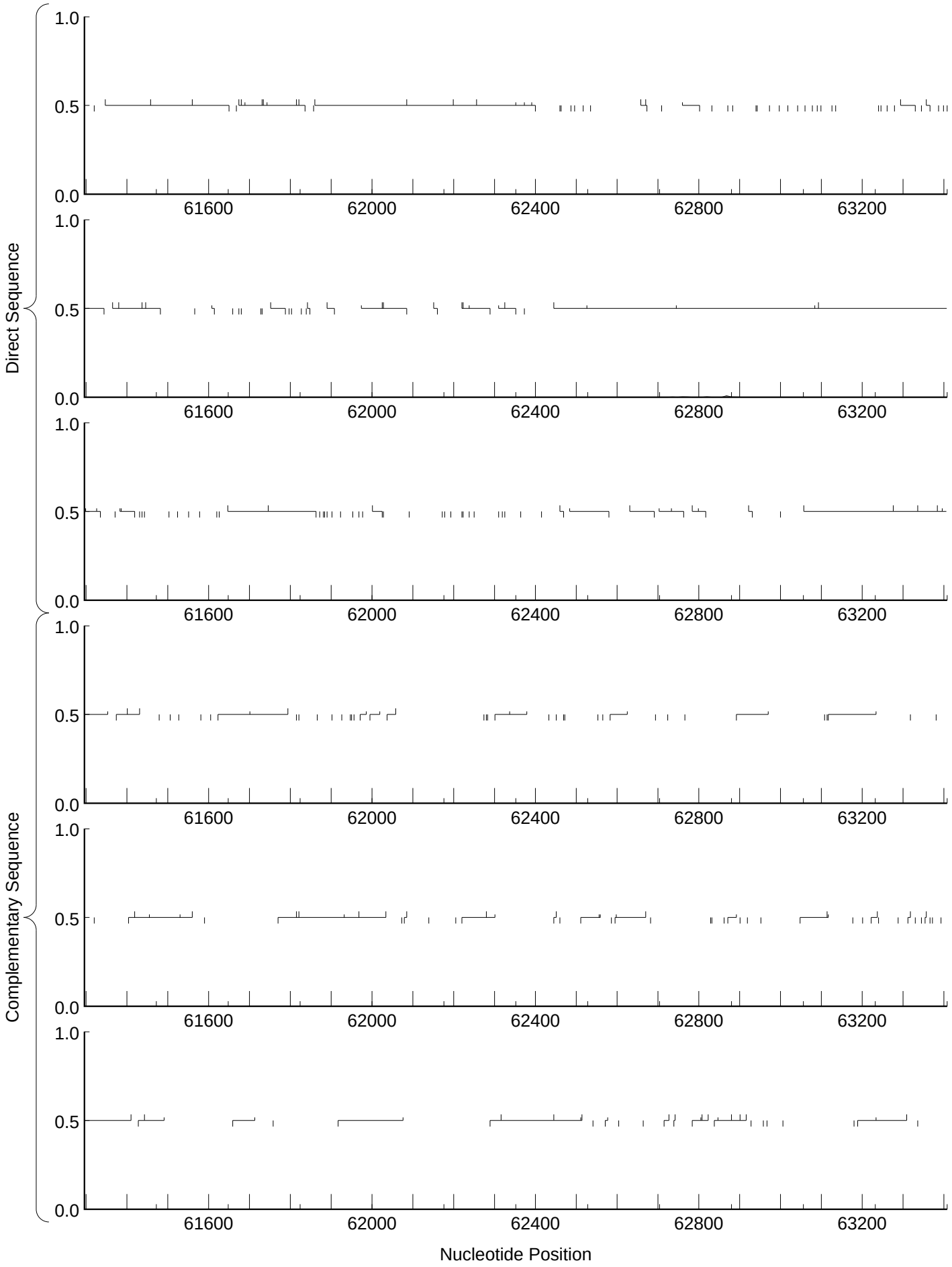
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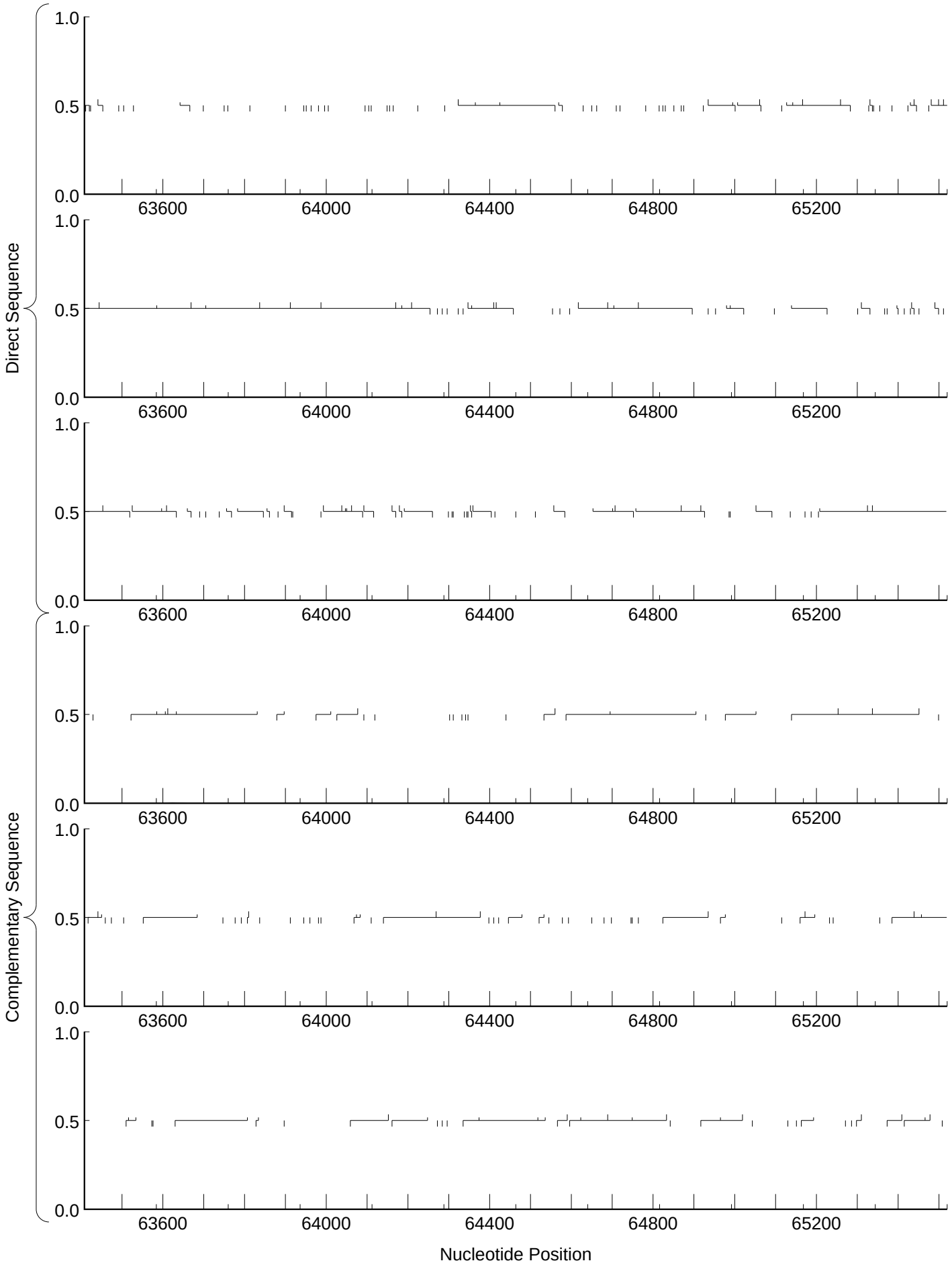
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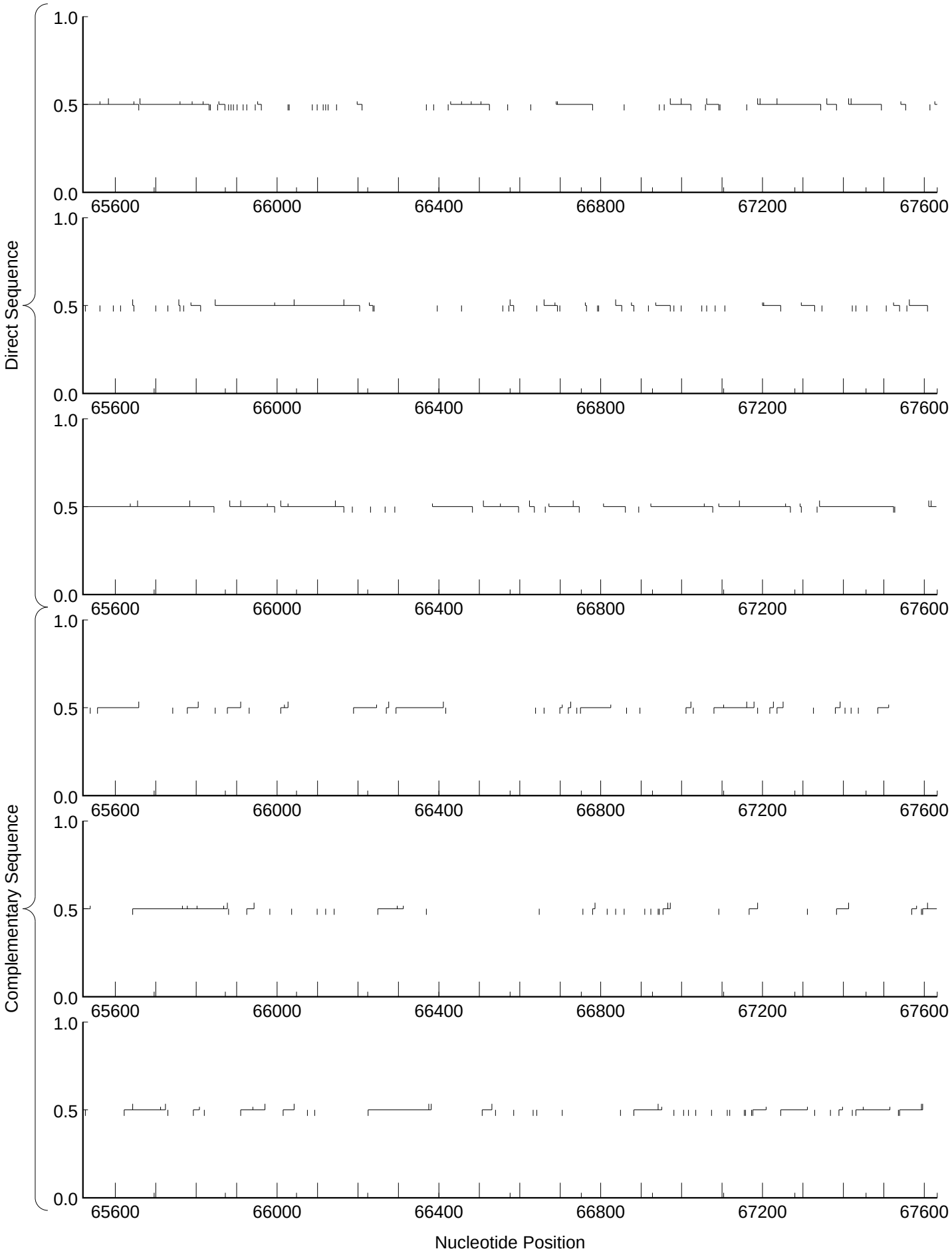
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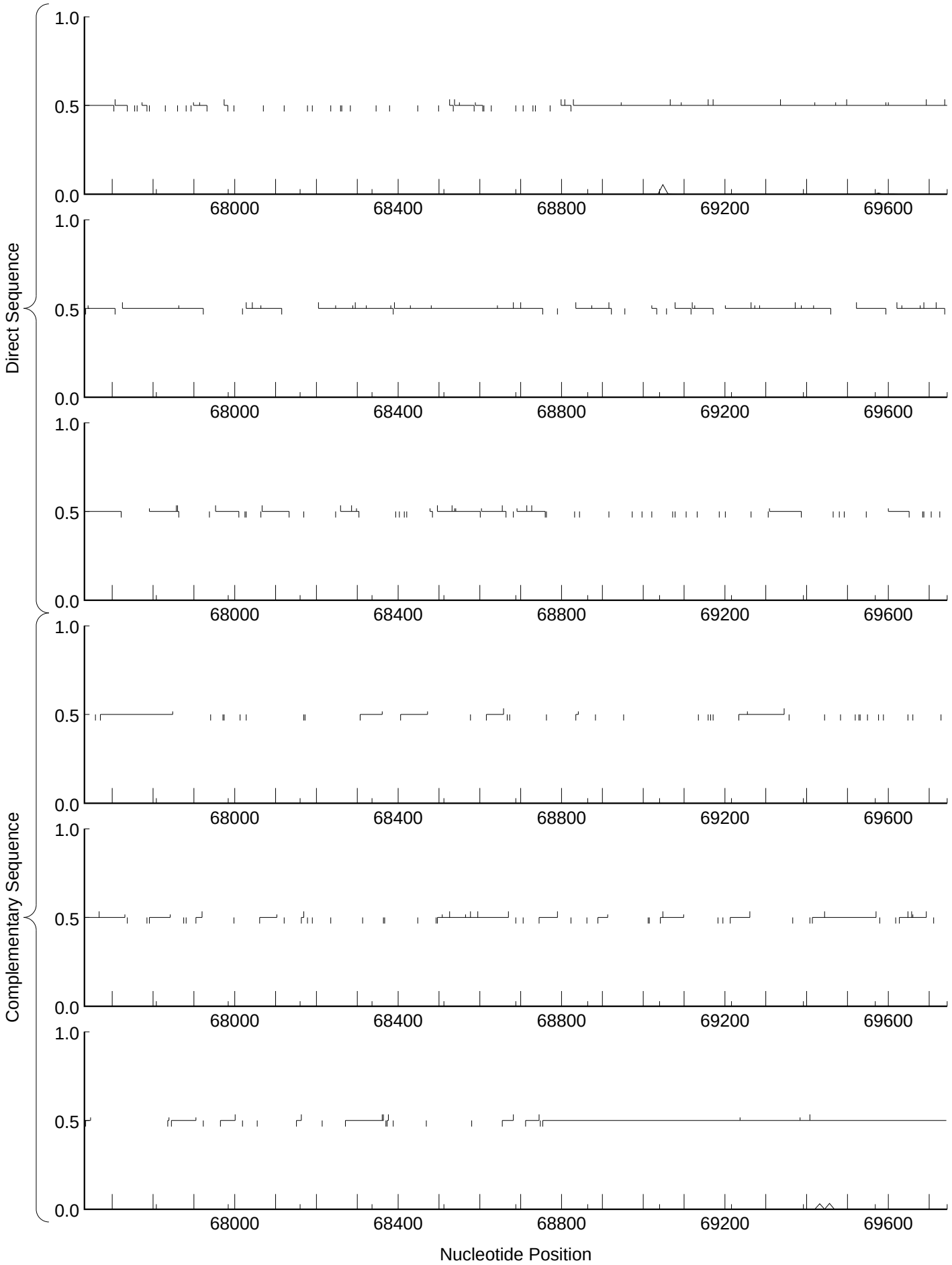
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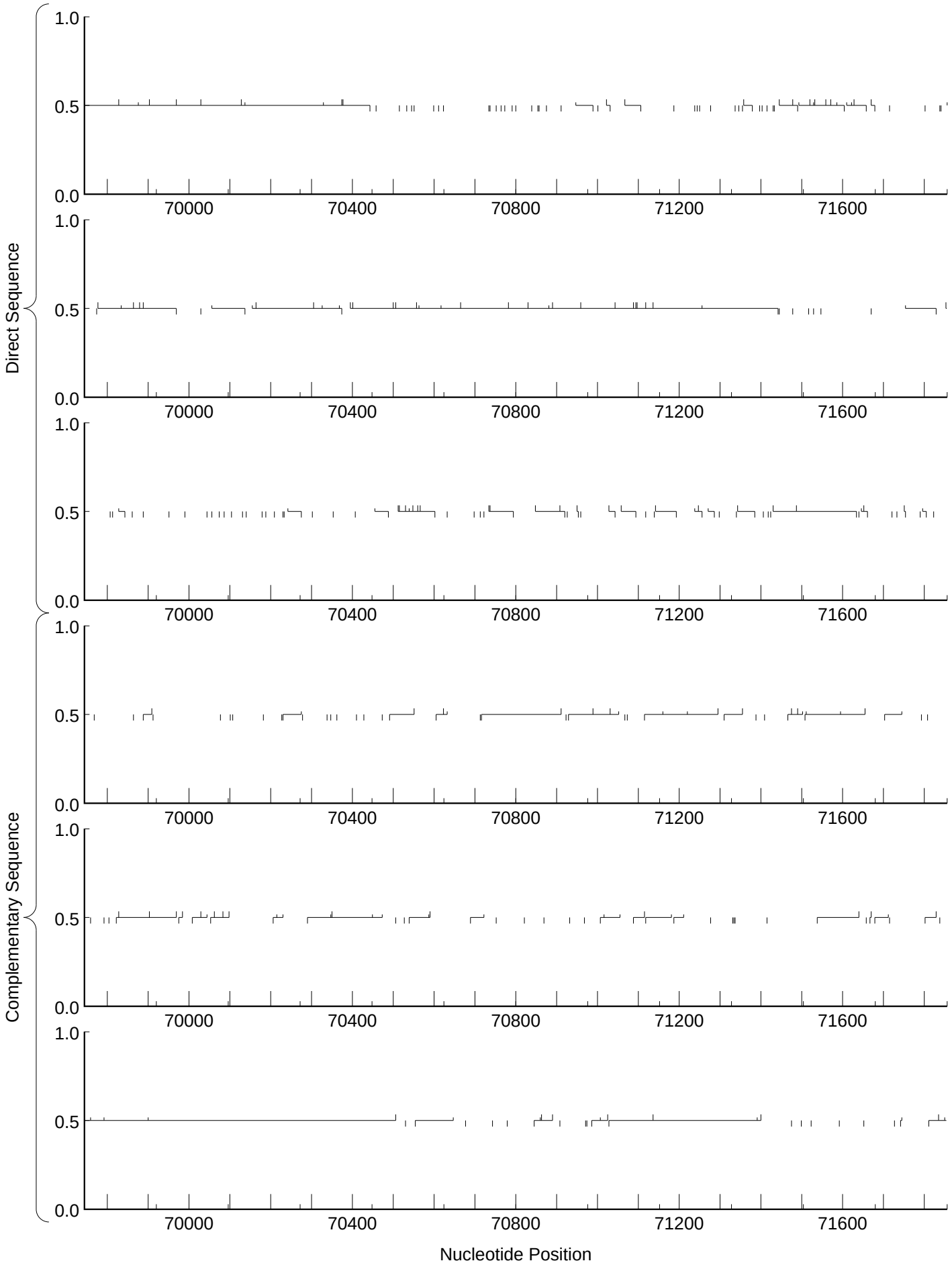
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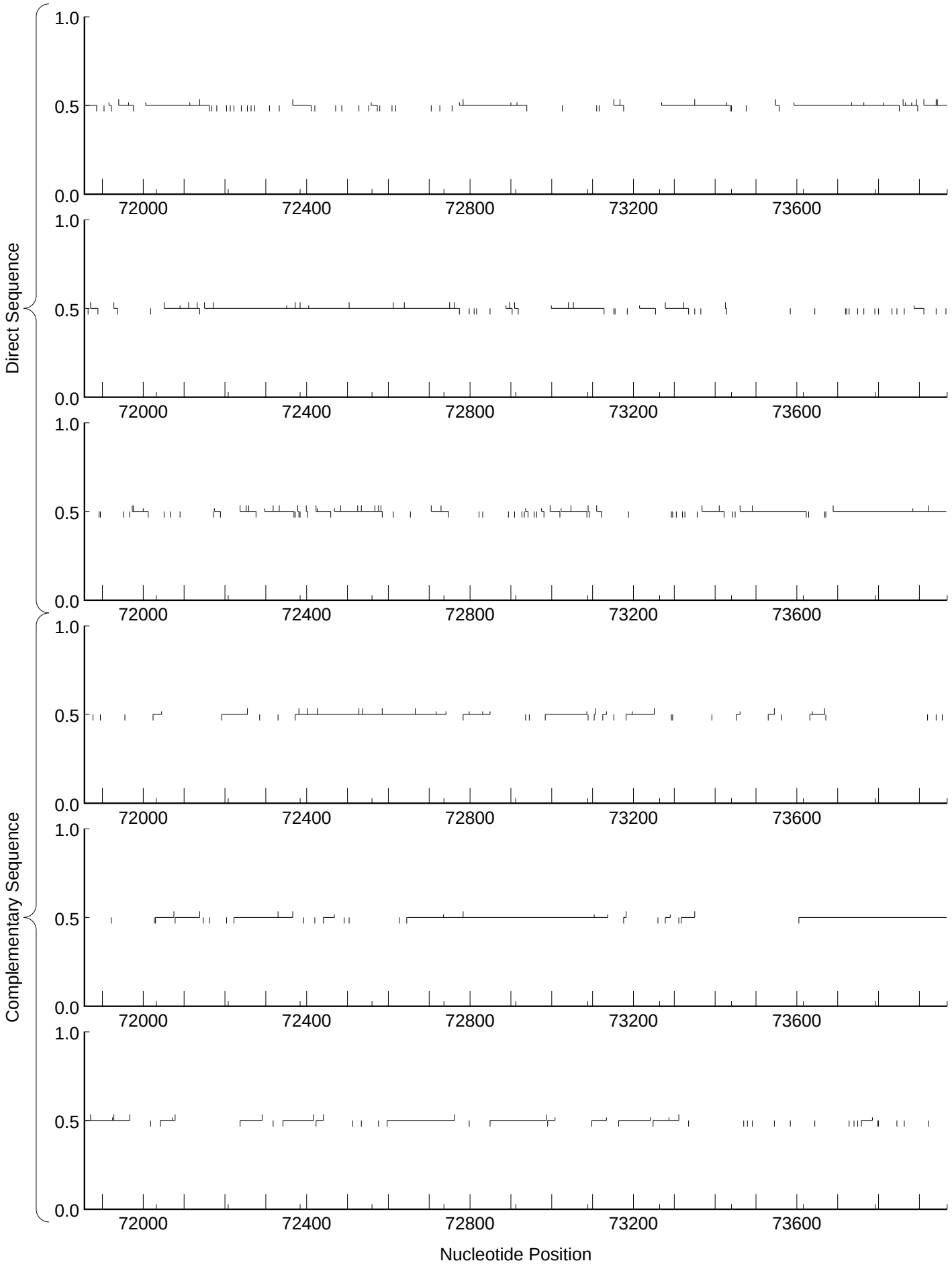
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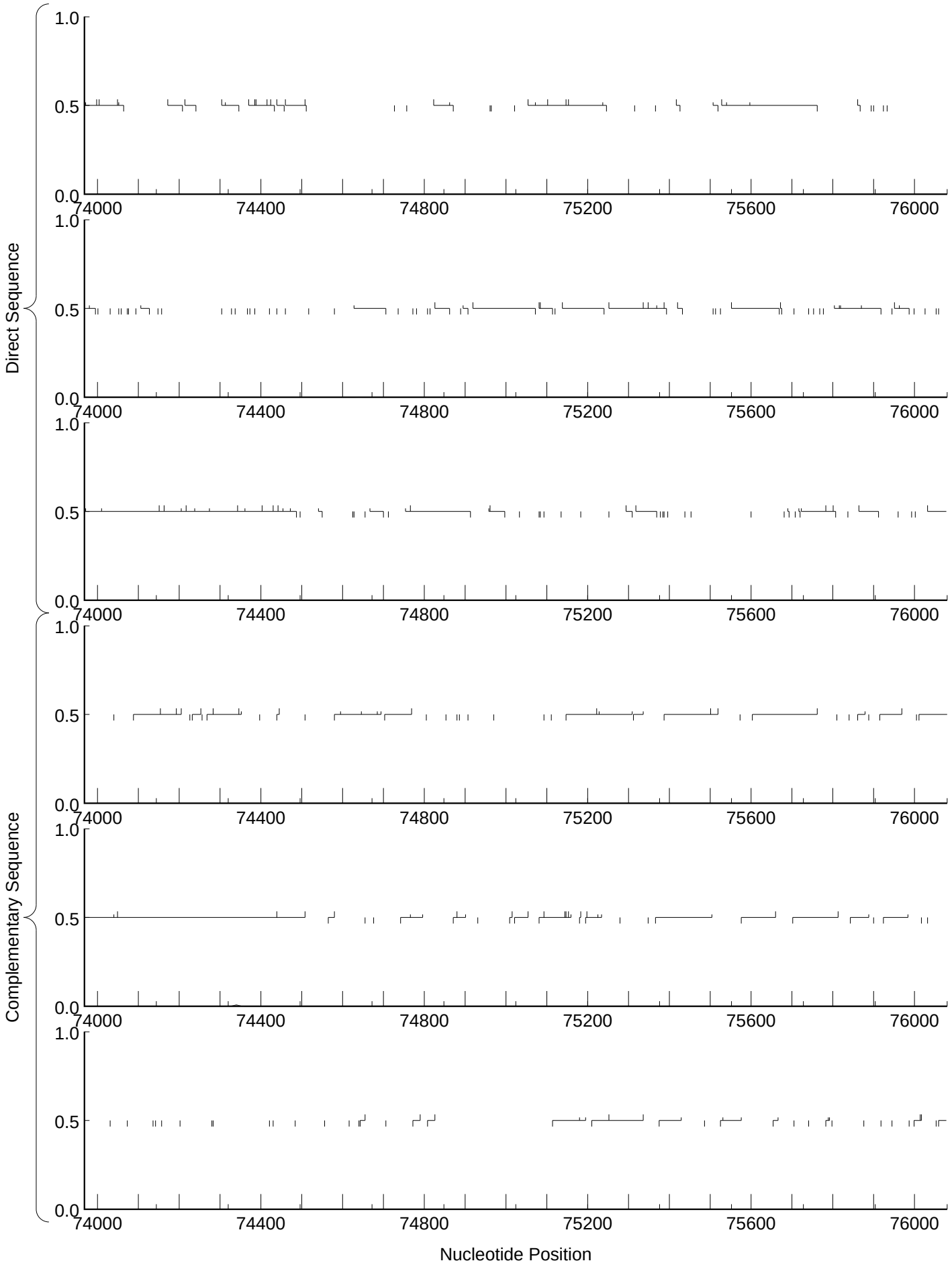
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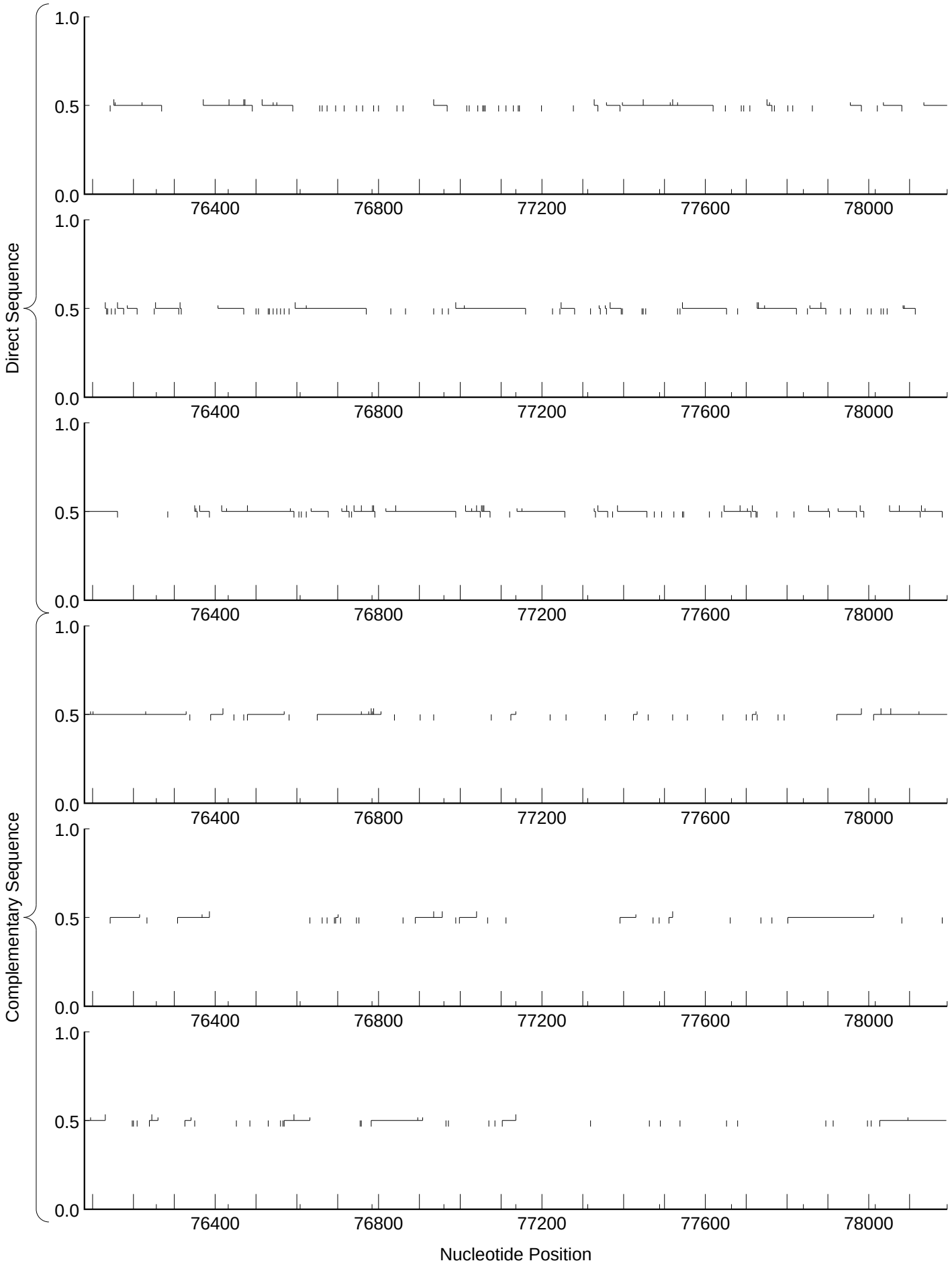
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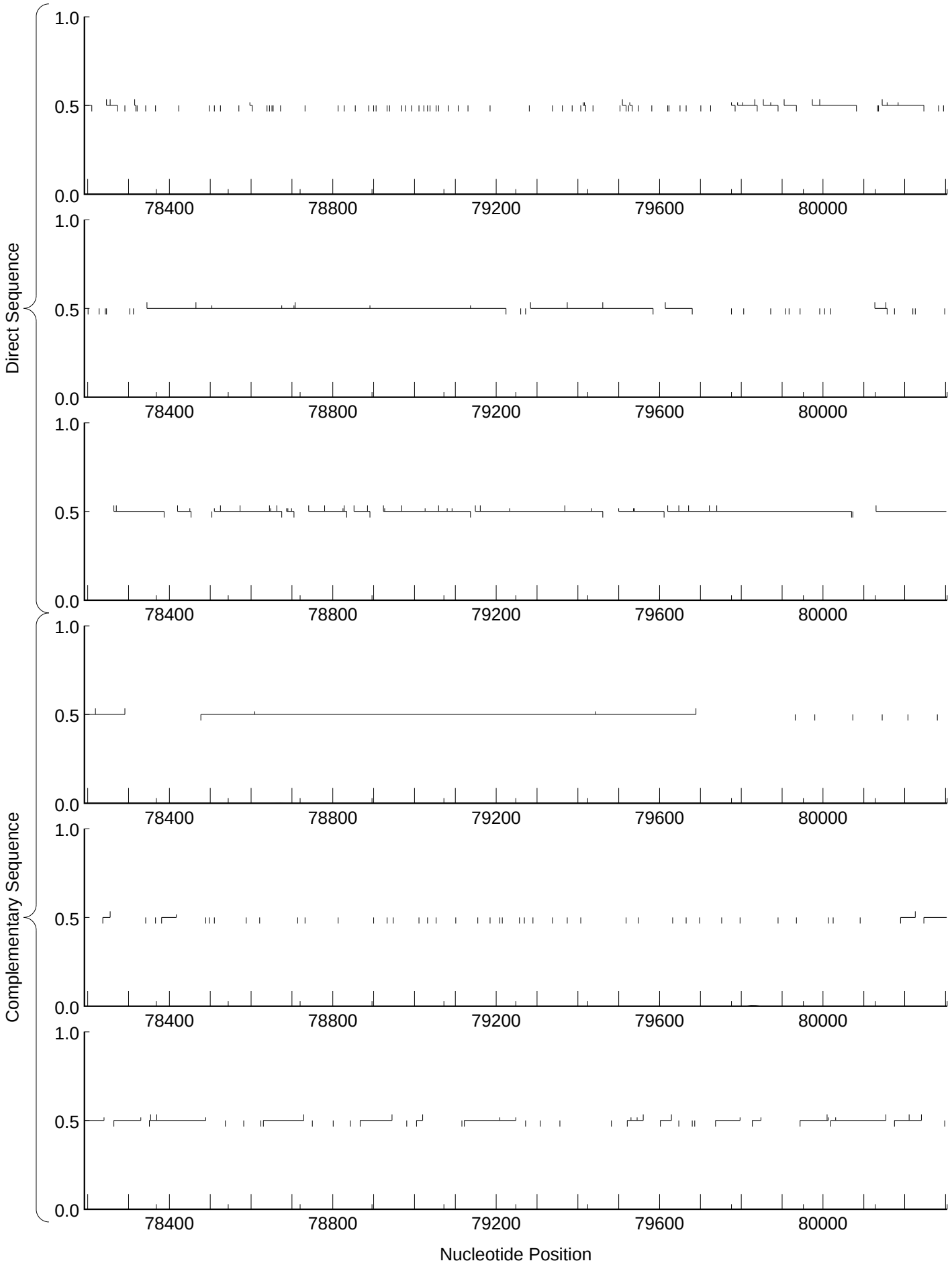
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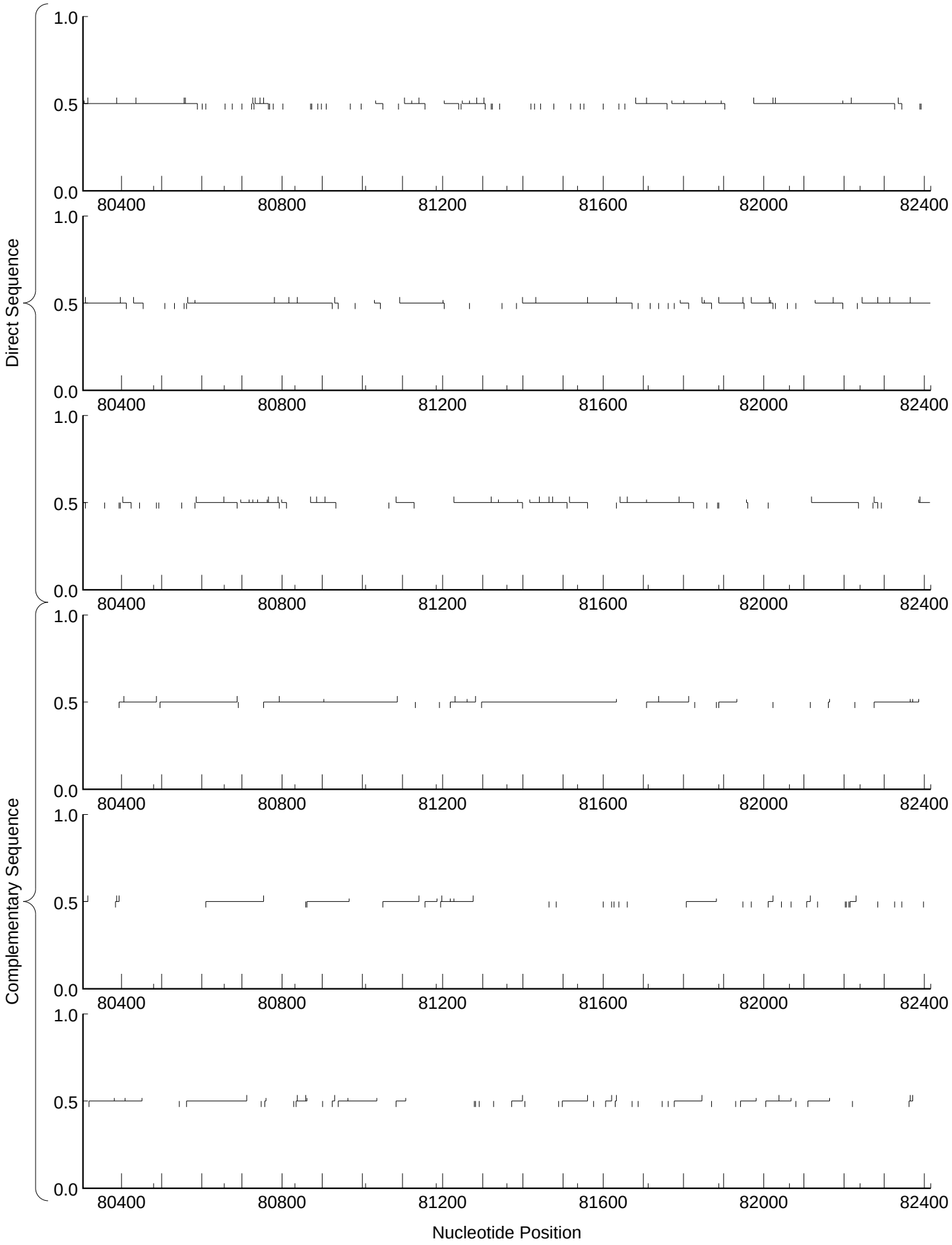
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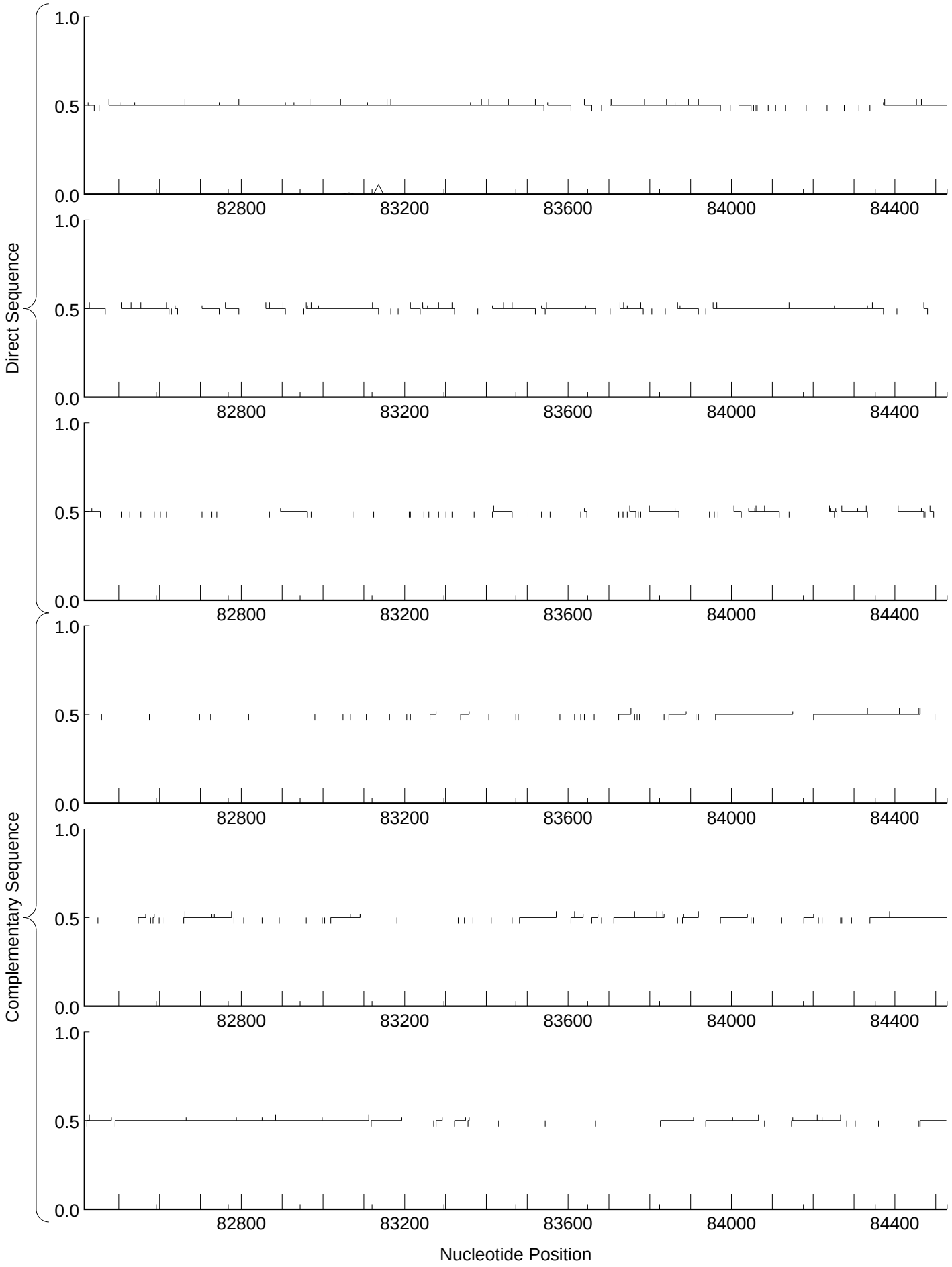
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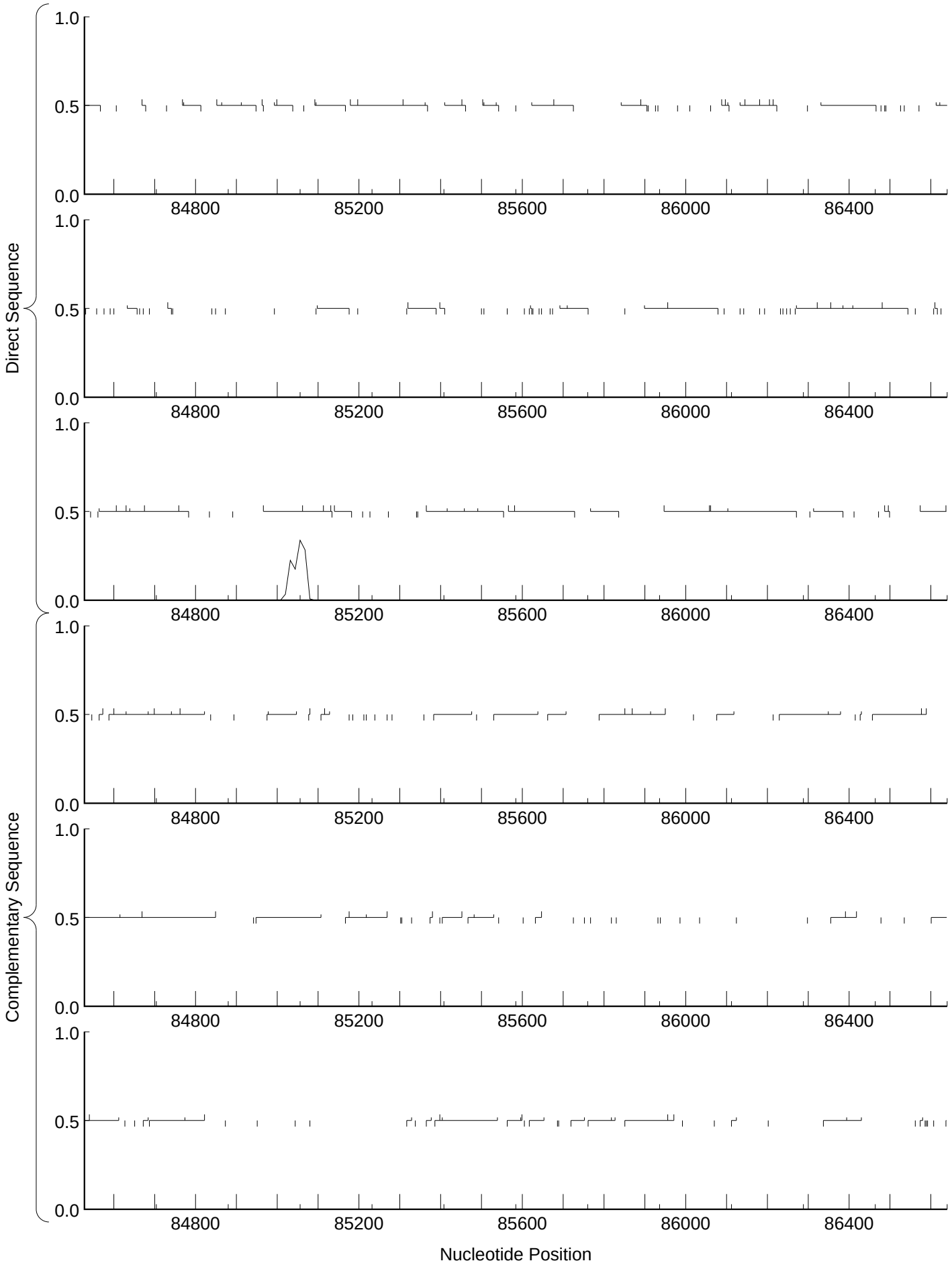
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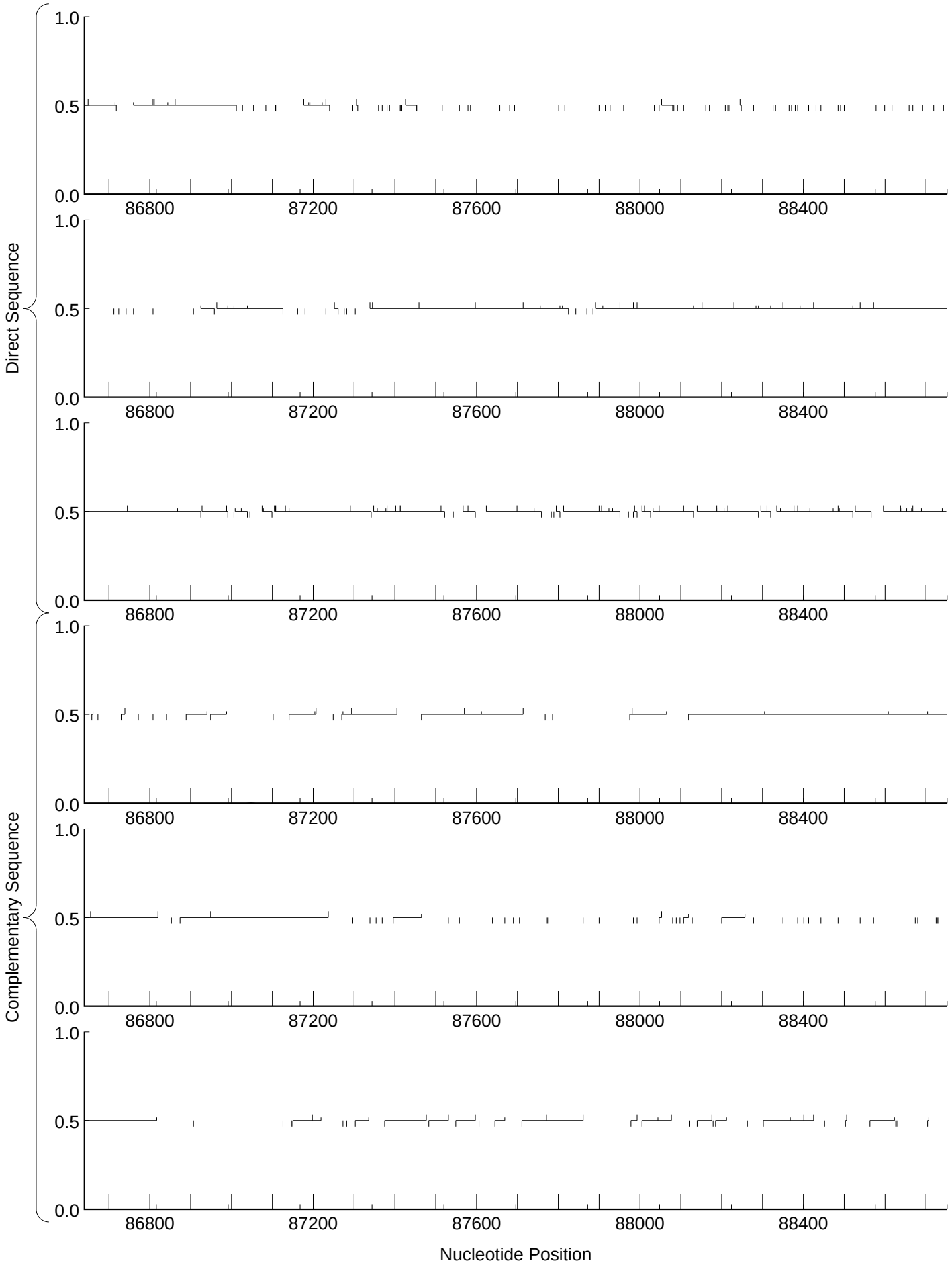
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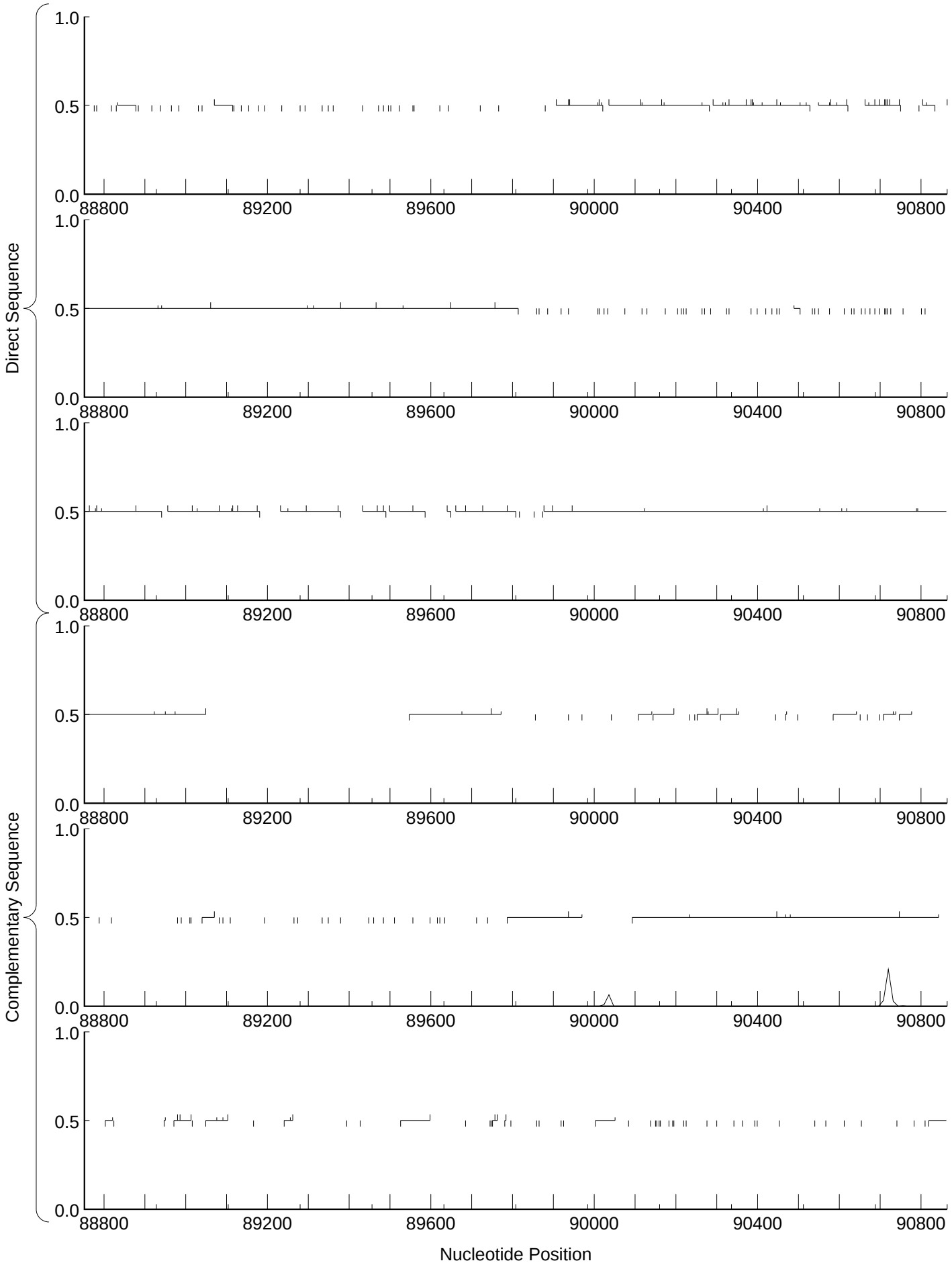
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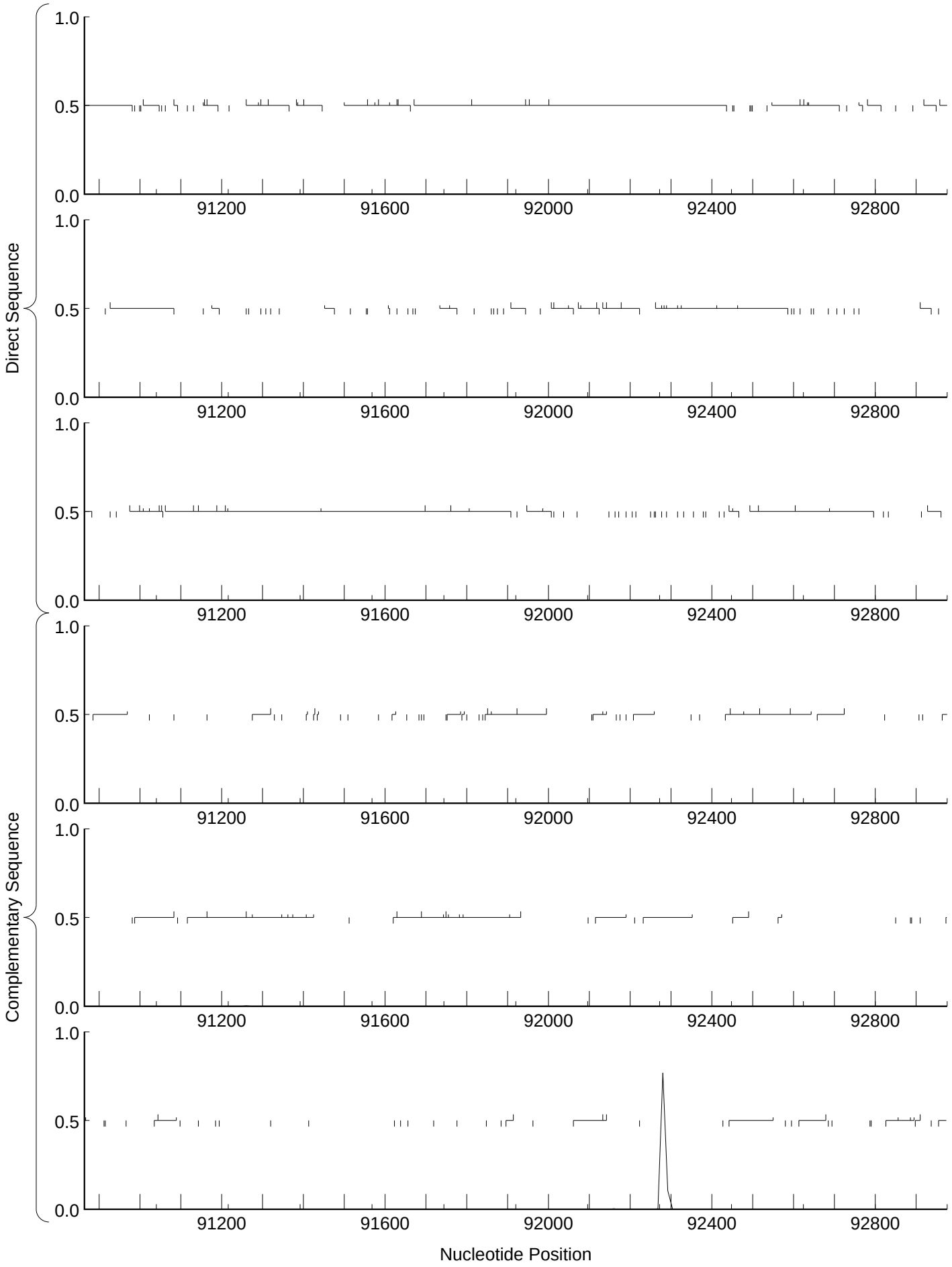
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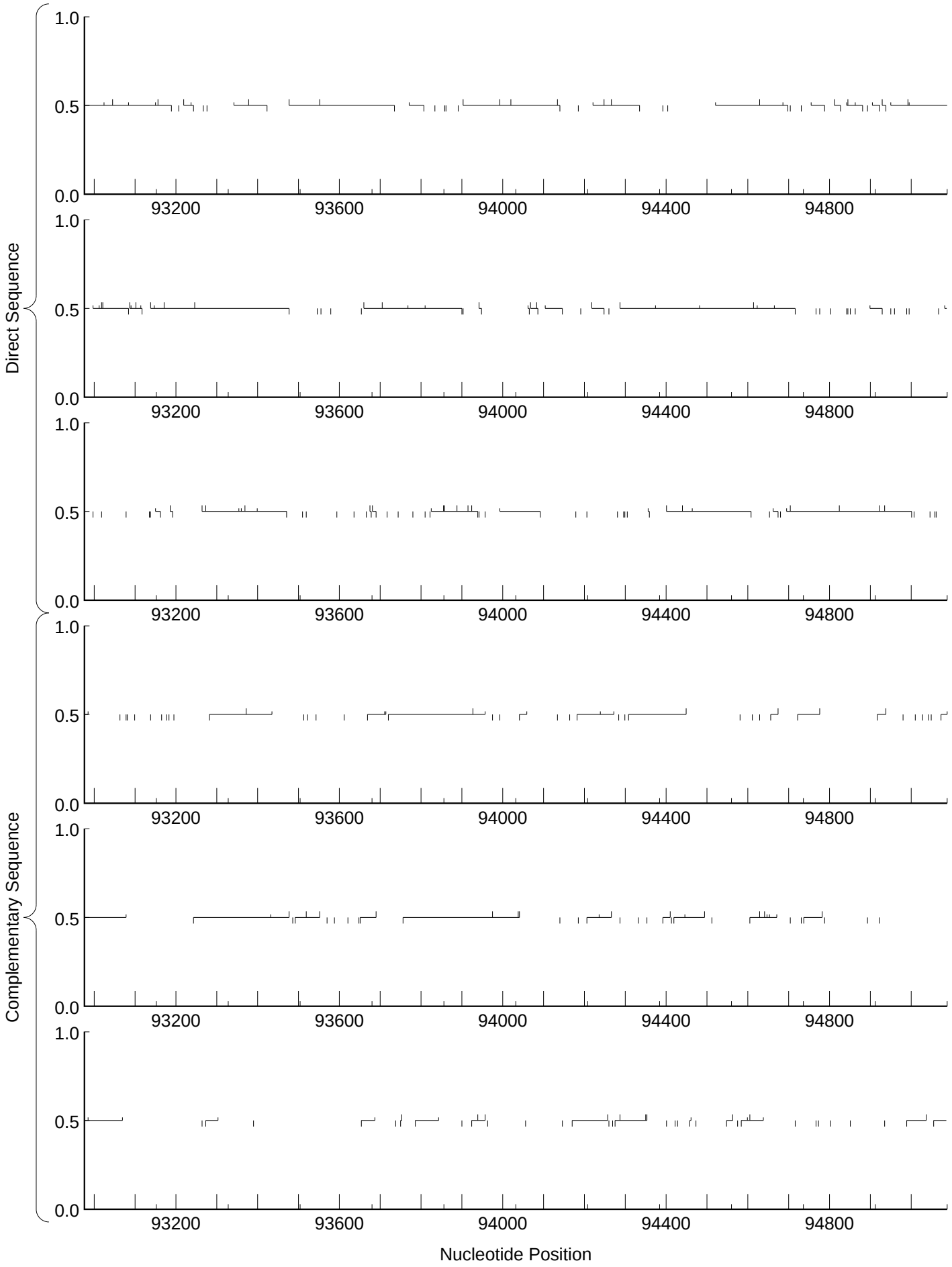
phage Gilson complete sequence, 128338 bp including 766 bp terminal repeat, Cluster BK1, Order 4, Window 96, Step 12, 44/62



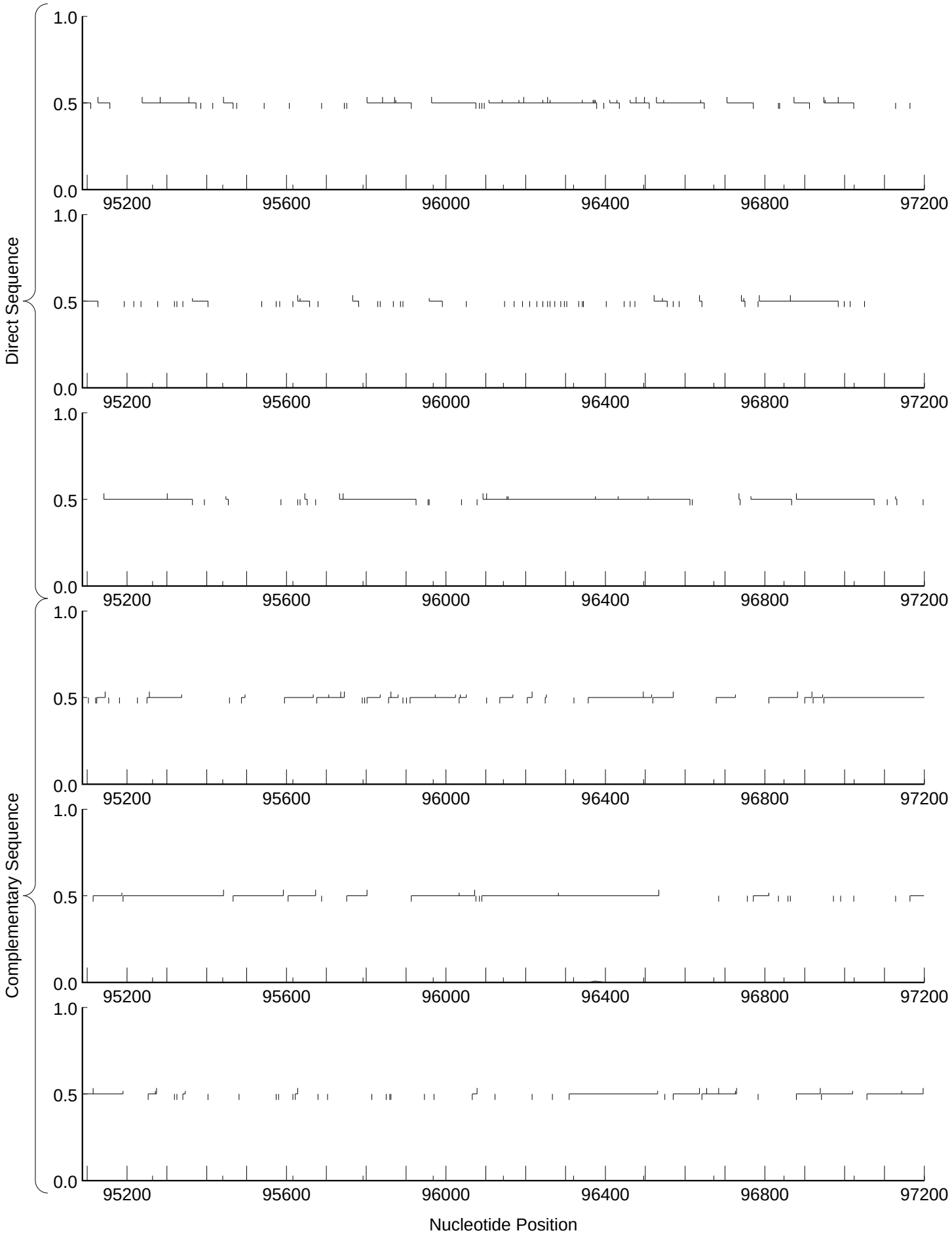
phage Gilson complete sequence, 128338 bp including 766 bp terminal repeat, Cluster BK1, Order 4, Window 96, Step 12, 45/62



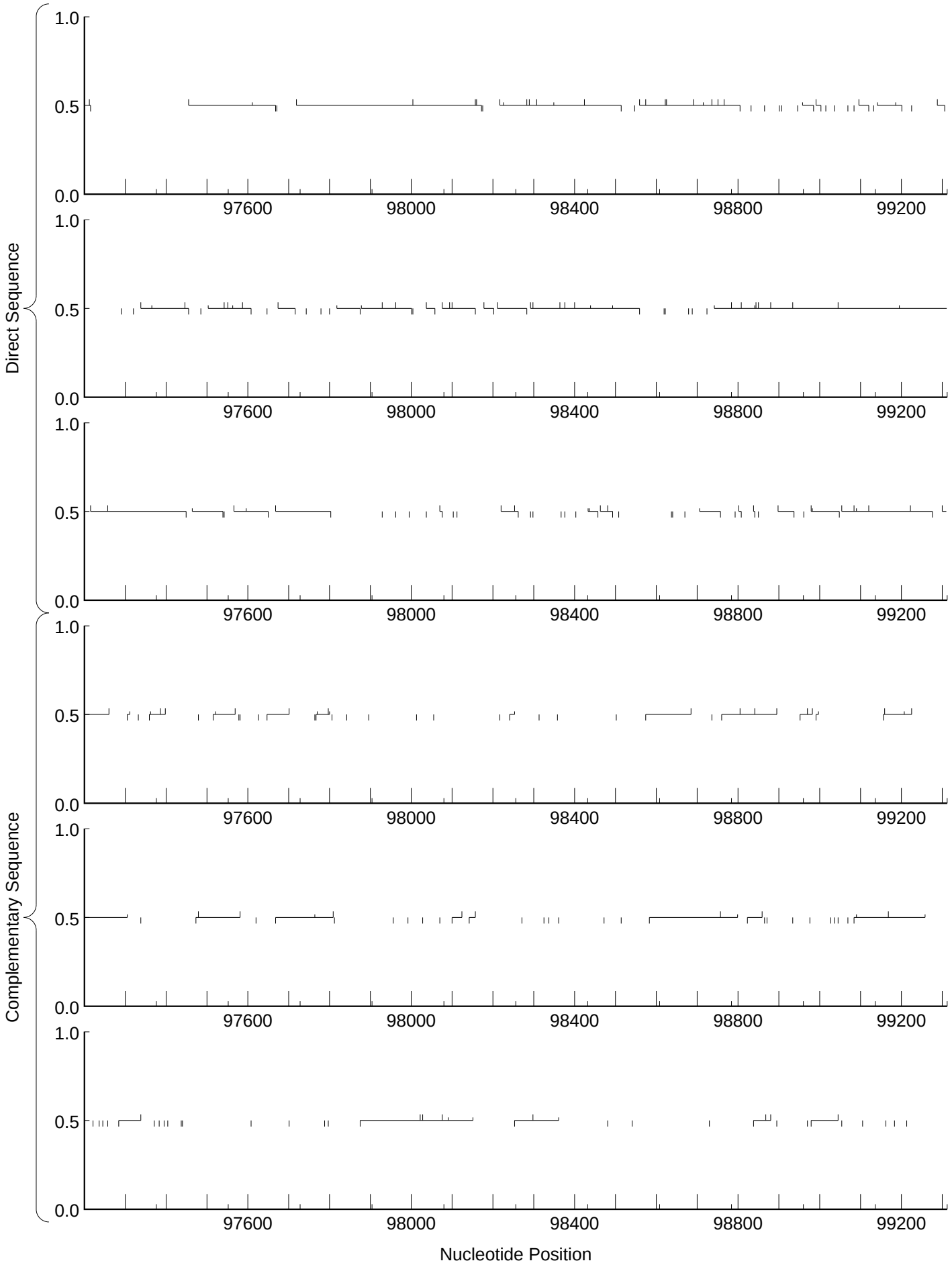
phage Gilson complete sequence, 128338 bp including 766 bp terminal repeat, Cluster BK1, Order 4, Window 96, Step 12, 46/62



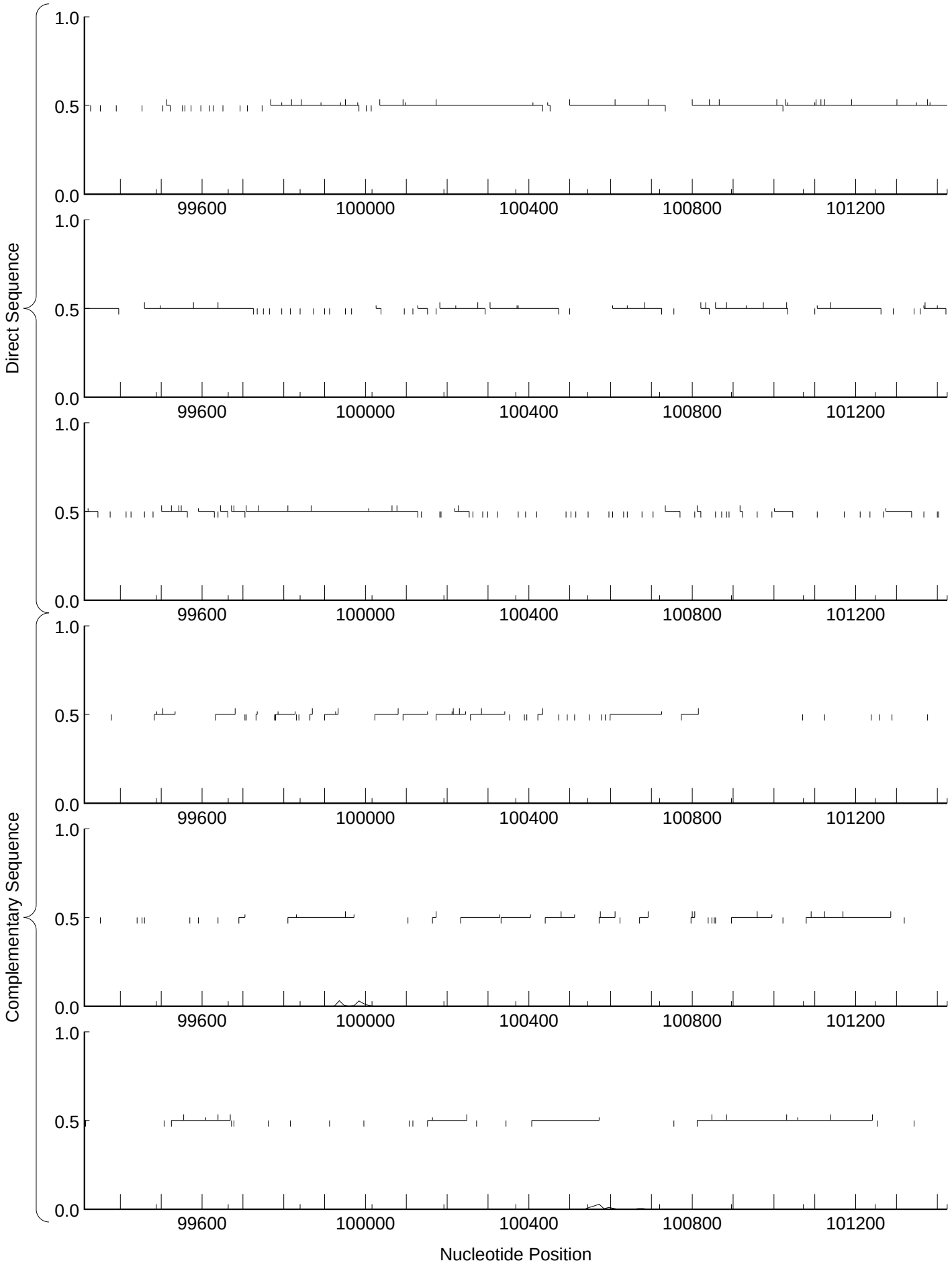
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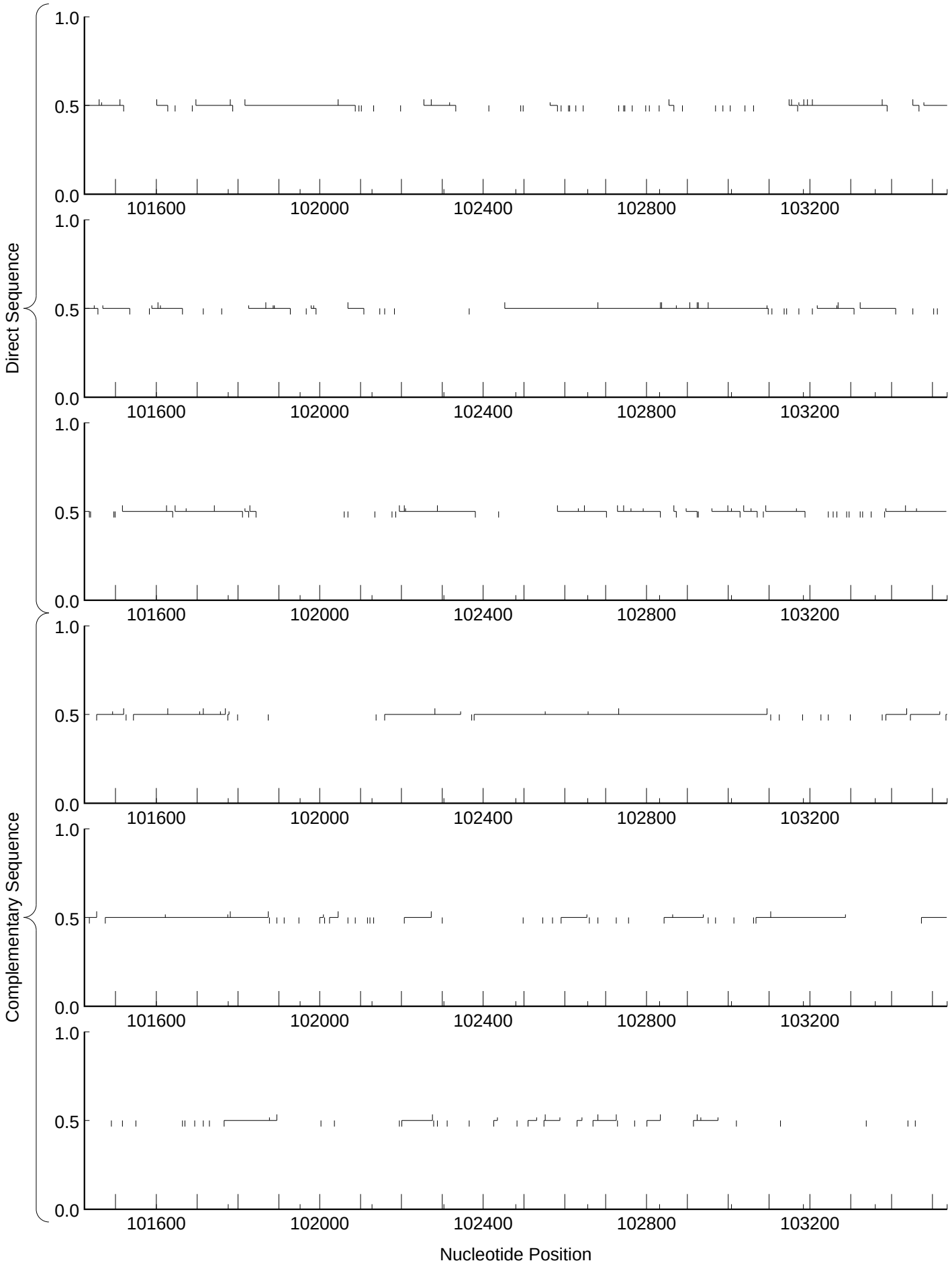
phage Gilson complete sequence, 128338 bp including 766 bp terminal repeat, Cluster BK1, Order 4, Window 96, Step 12, 48/62



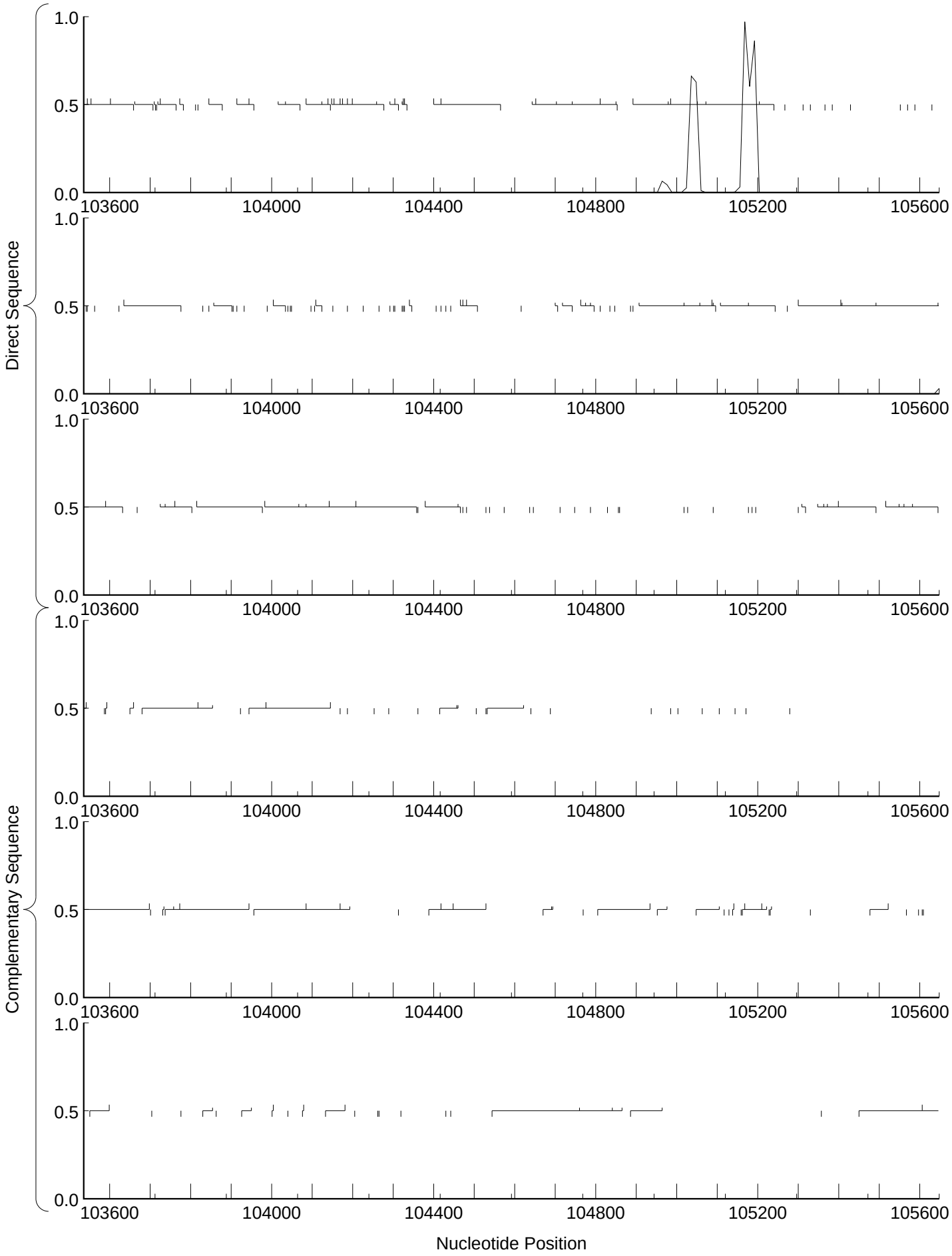
phage Gilson complete sequence, 128338 bp including 766 bp terminal repeat, Cluster BK1, Order 4, Window 96, Step 12, 49/62



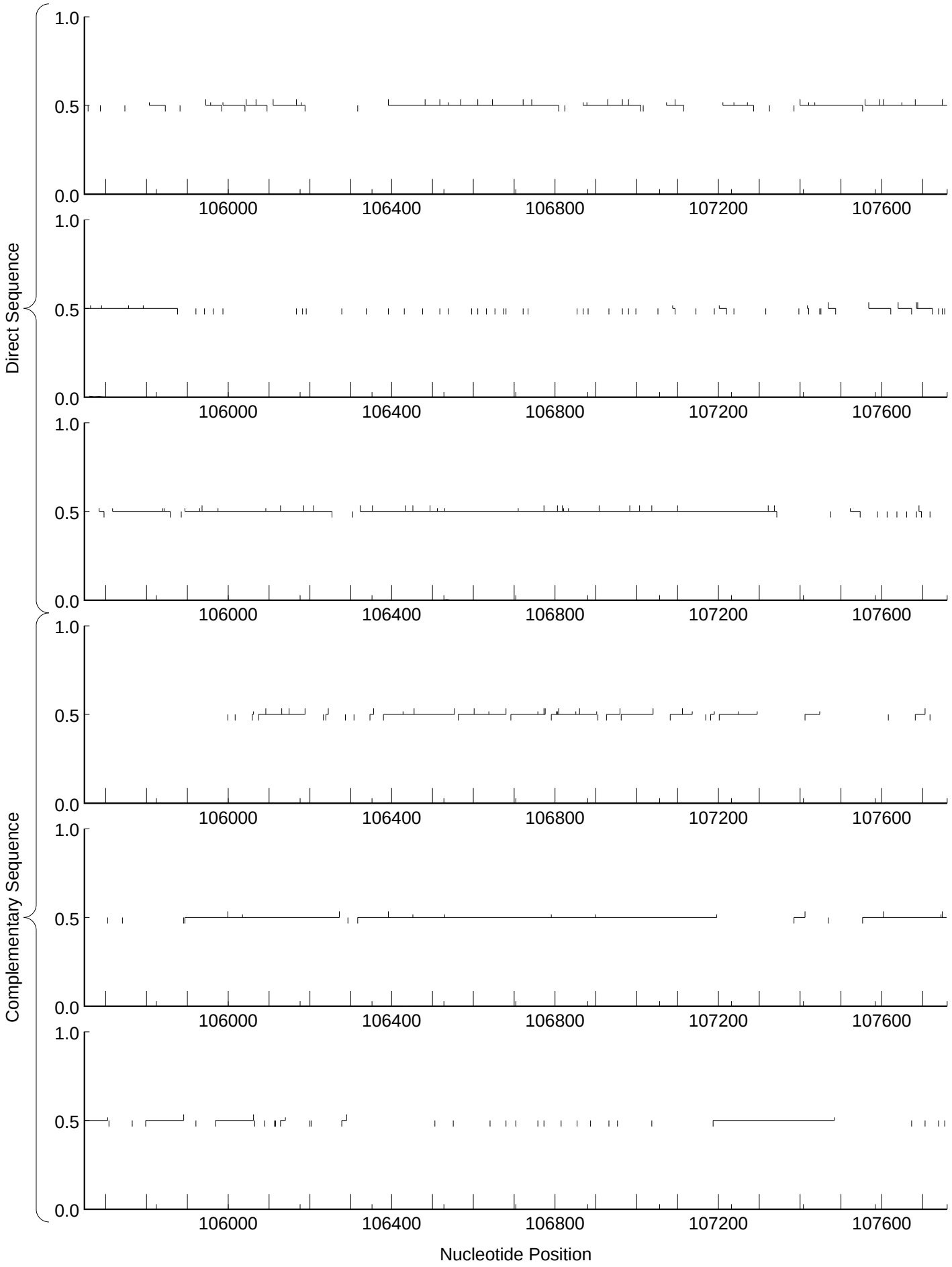
phage Gilson complete sequence, 128338 bp including 766 bp terminal repeat, Cluster BK1, Order 4, Window 96, Step 12, 50/62



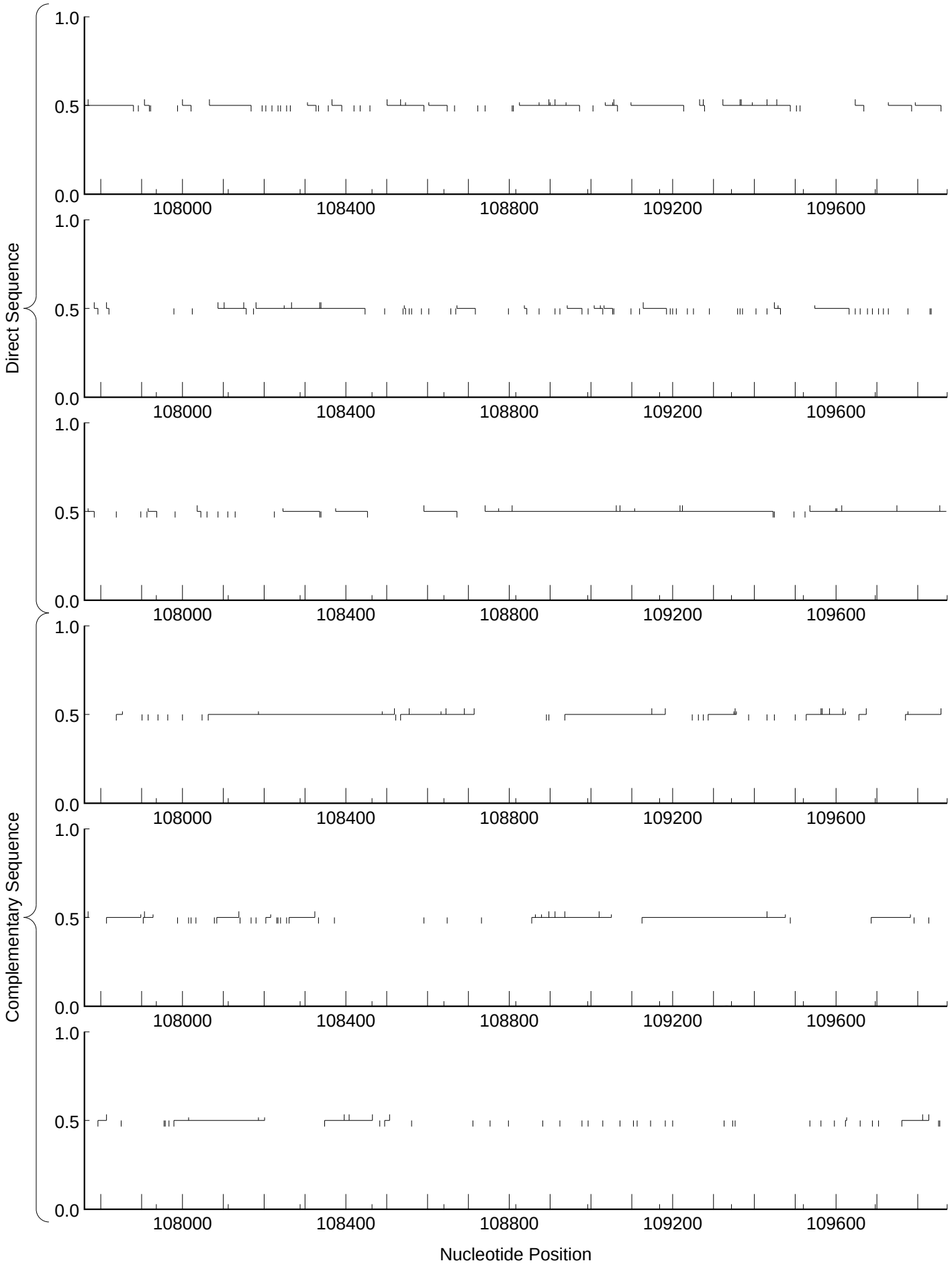
phage Gilson complete sequence, 128338 bp including 766 bp terminal repeat, Cluster BK1, Order 4, Window 96, Step 12, 51/62



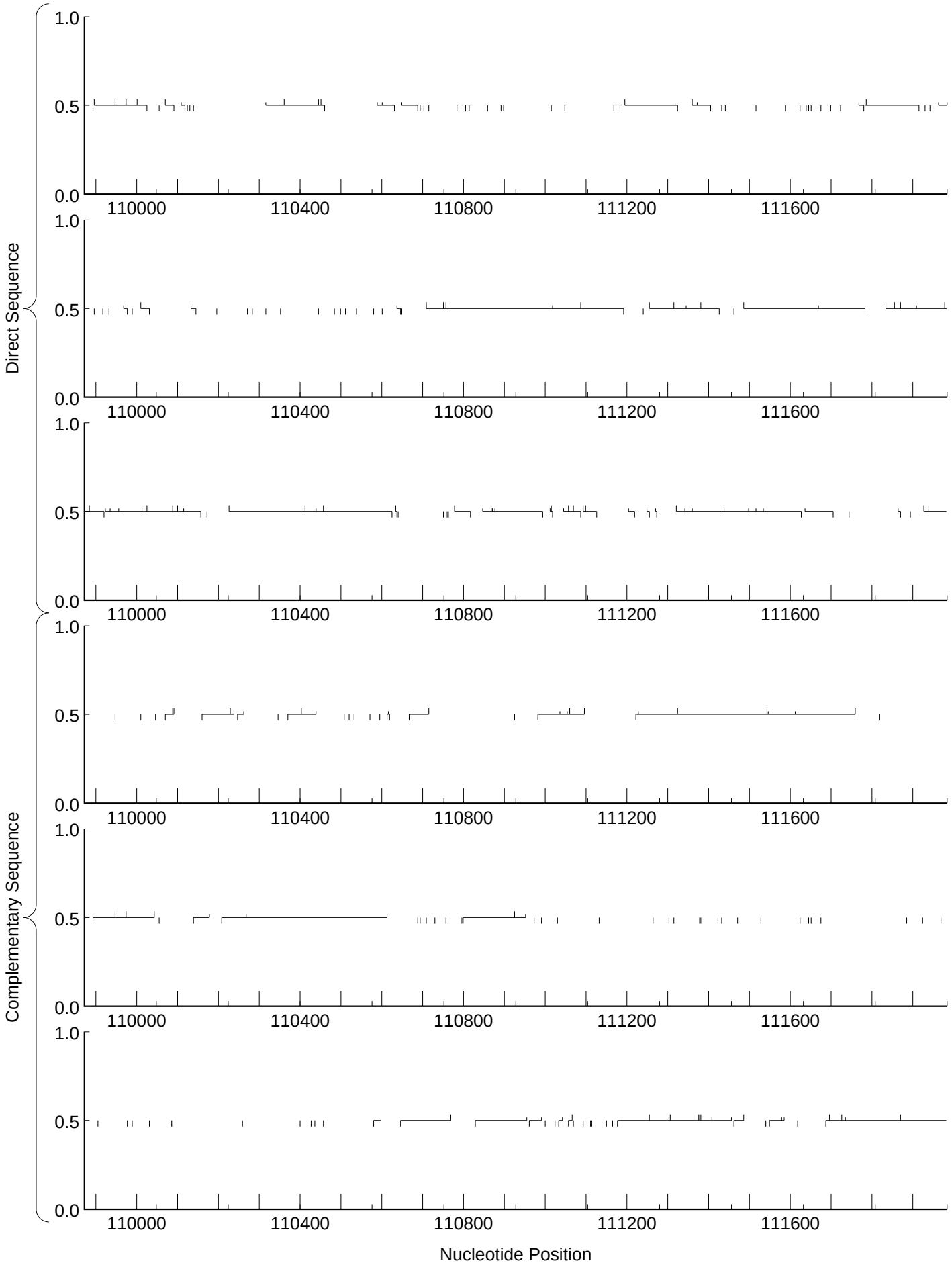
phage Gilson complete sequence, 128338 bp including 766 bp terminal repeat, Cluster BK1, Order 4, Window 96, Step 12, 52/62



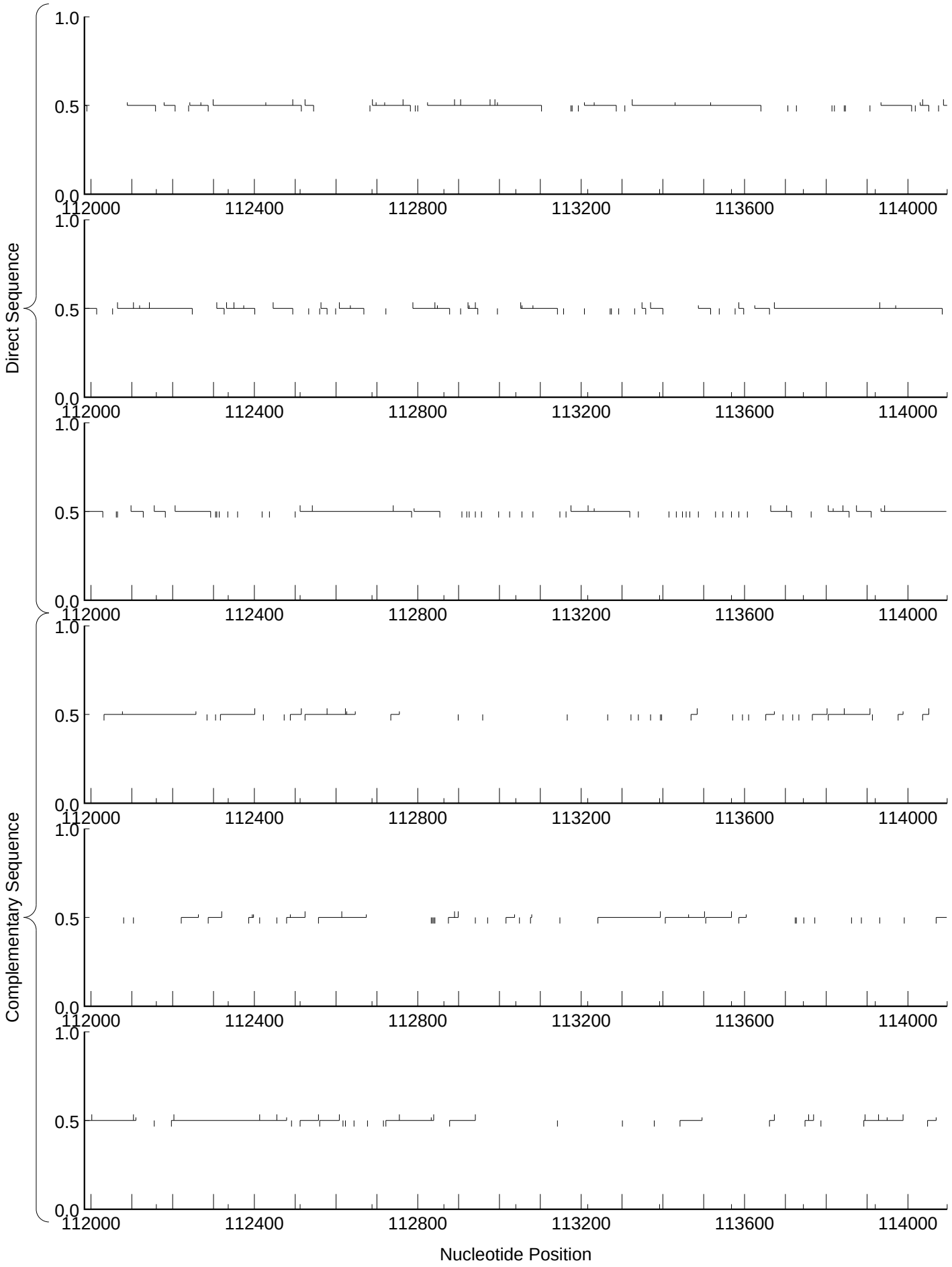
phage Gilson complete sequence, 128338 bp including 766 bp terminal repeat, Cluster BK1, Order 4, Window 96, Step 12, 53/62



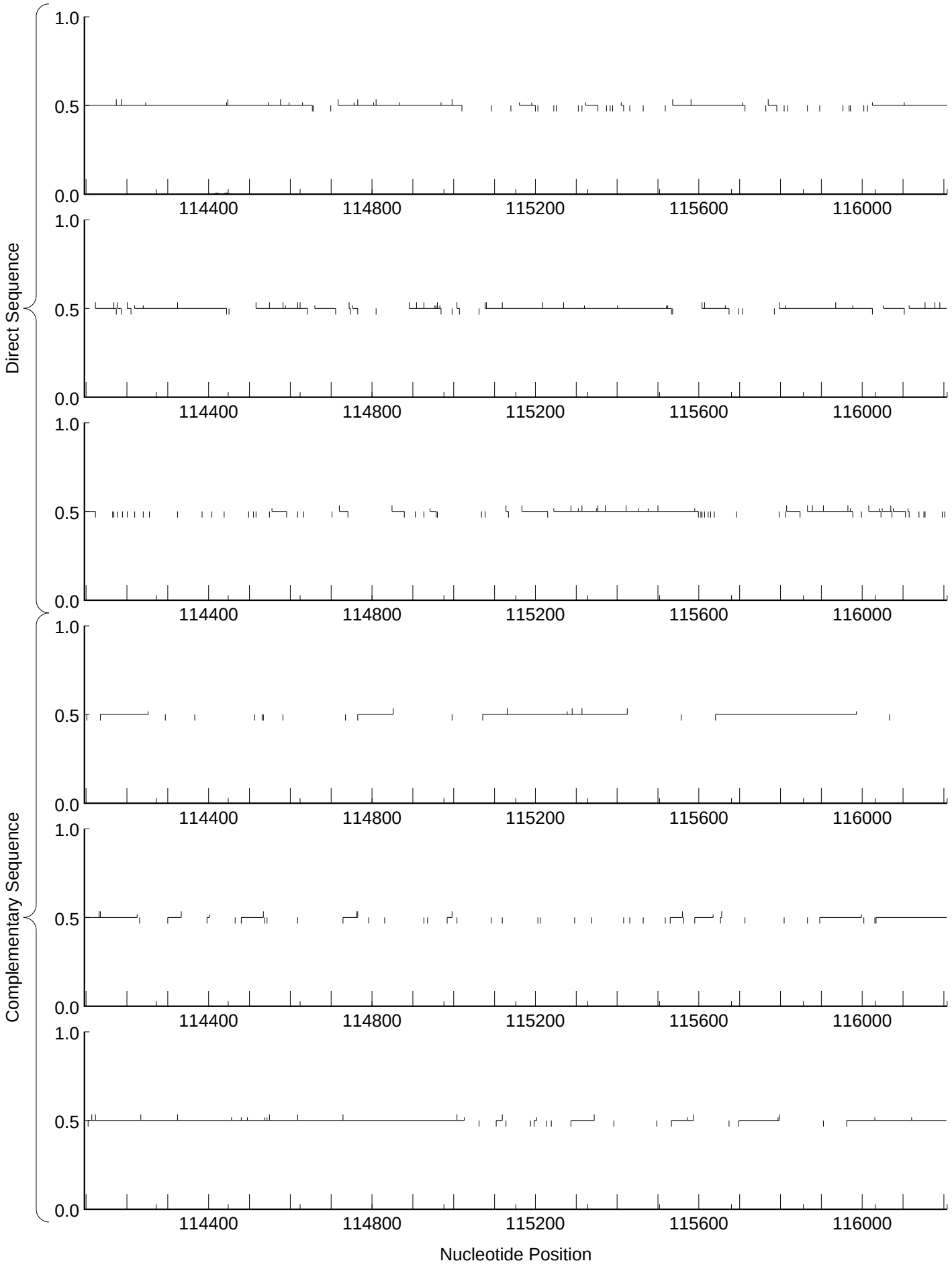
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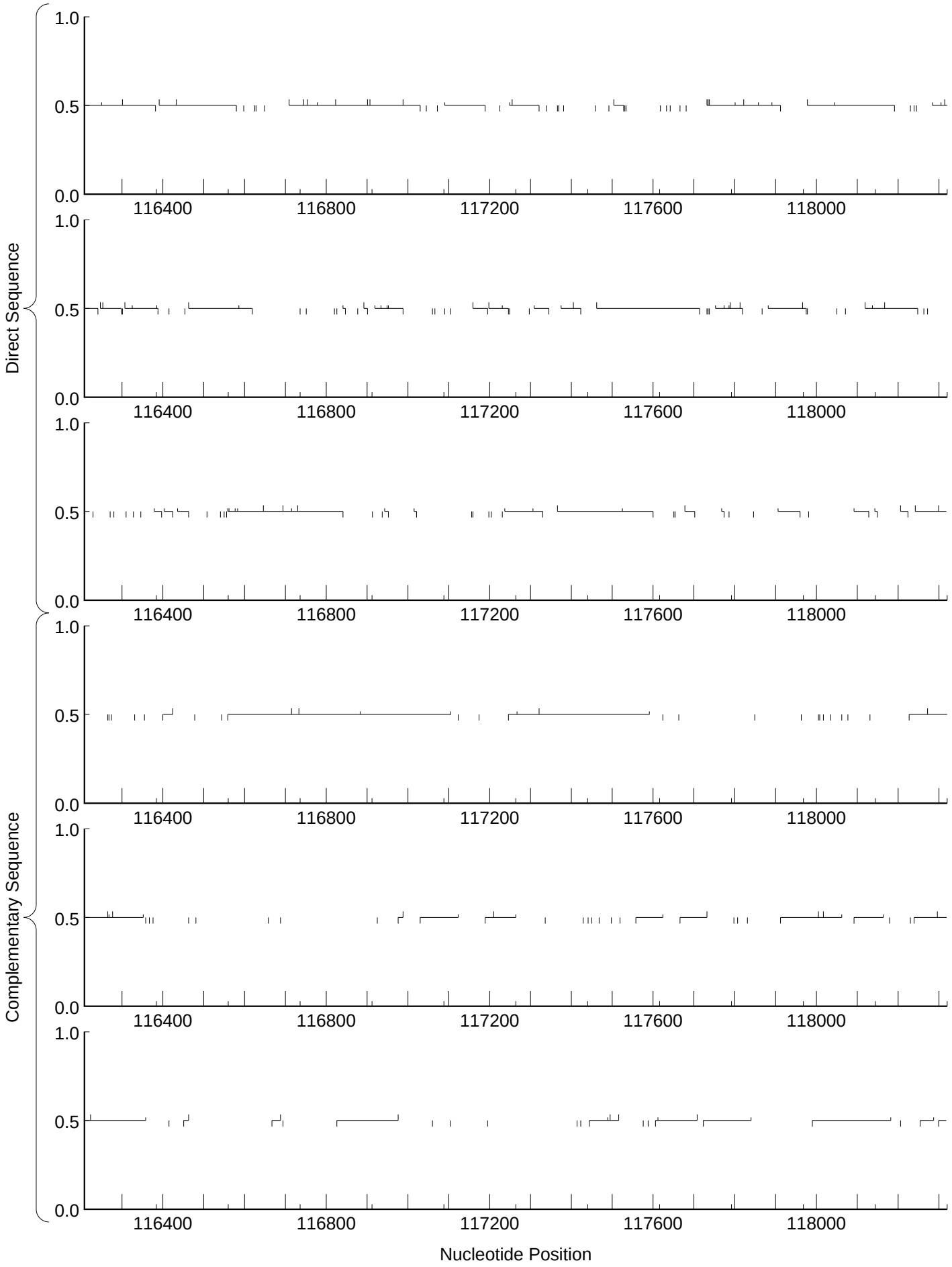
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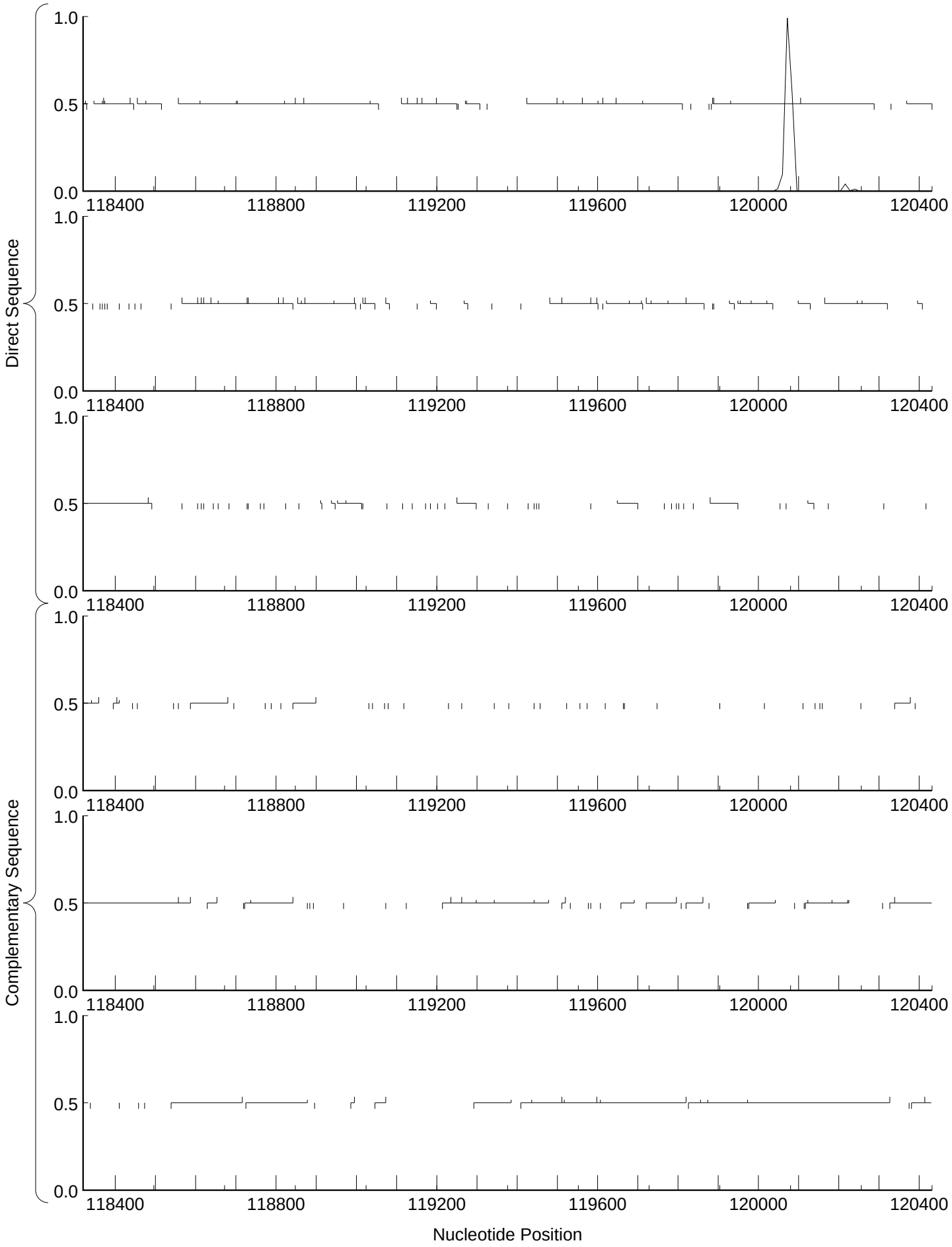
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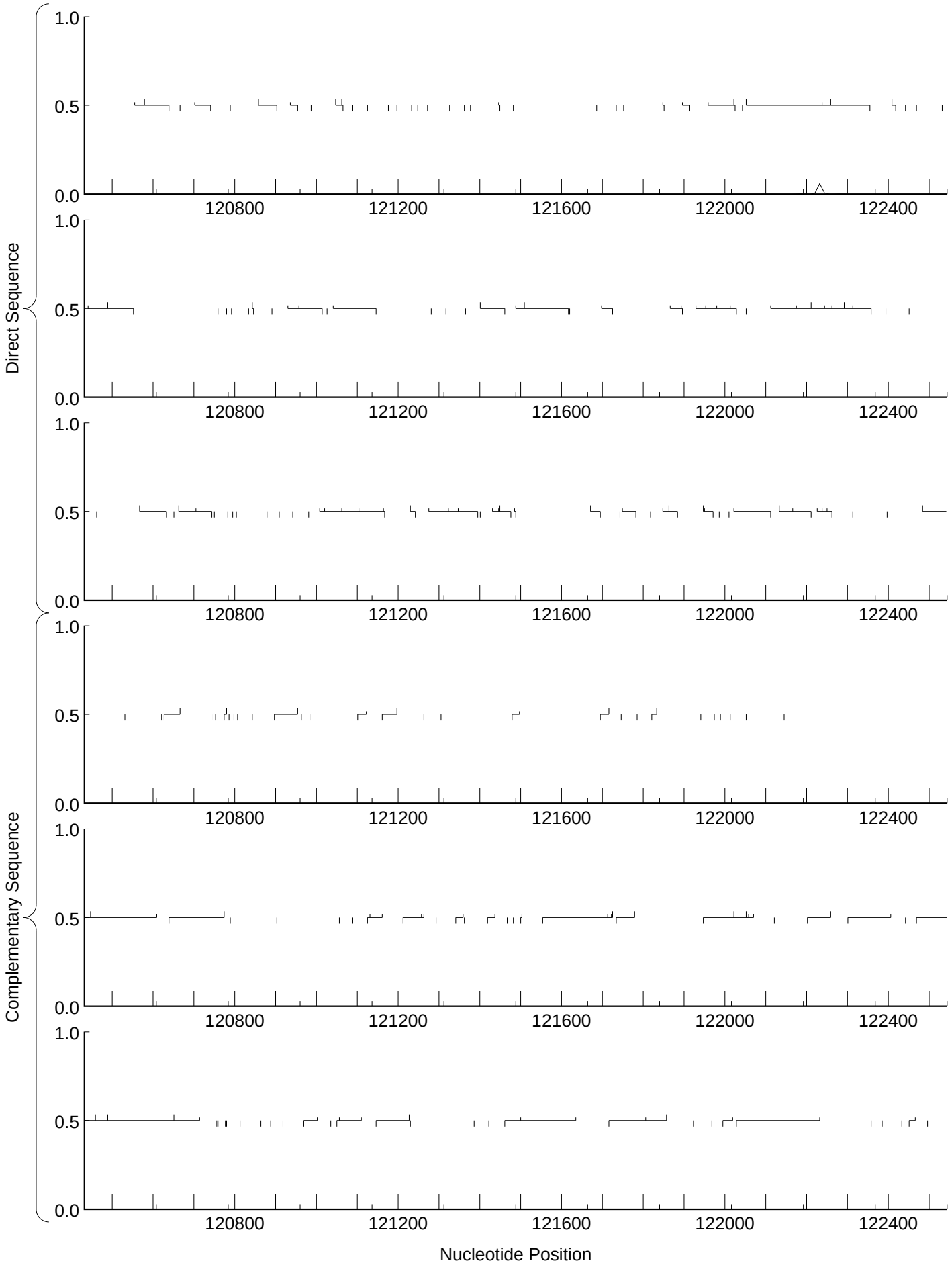
phage Gilson complete sequence, 128338 bp including 766 bp terminal repeat, Cluster BK1, Order 4, Window 96, Step 12, 57/62



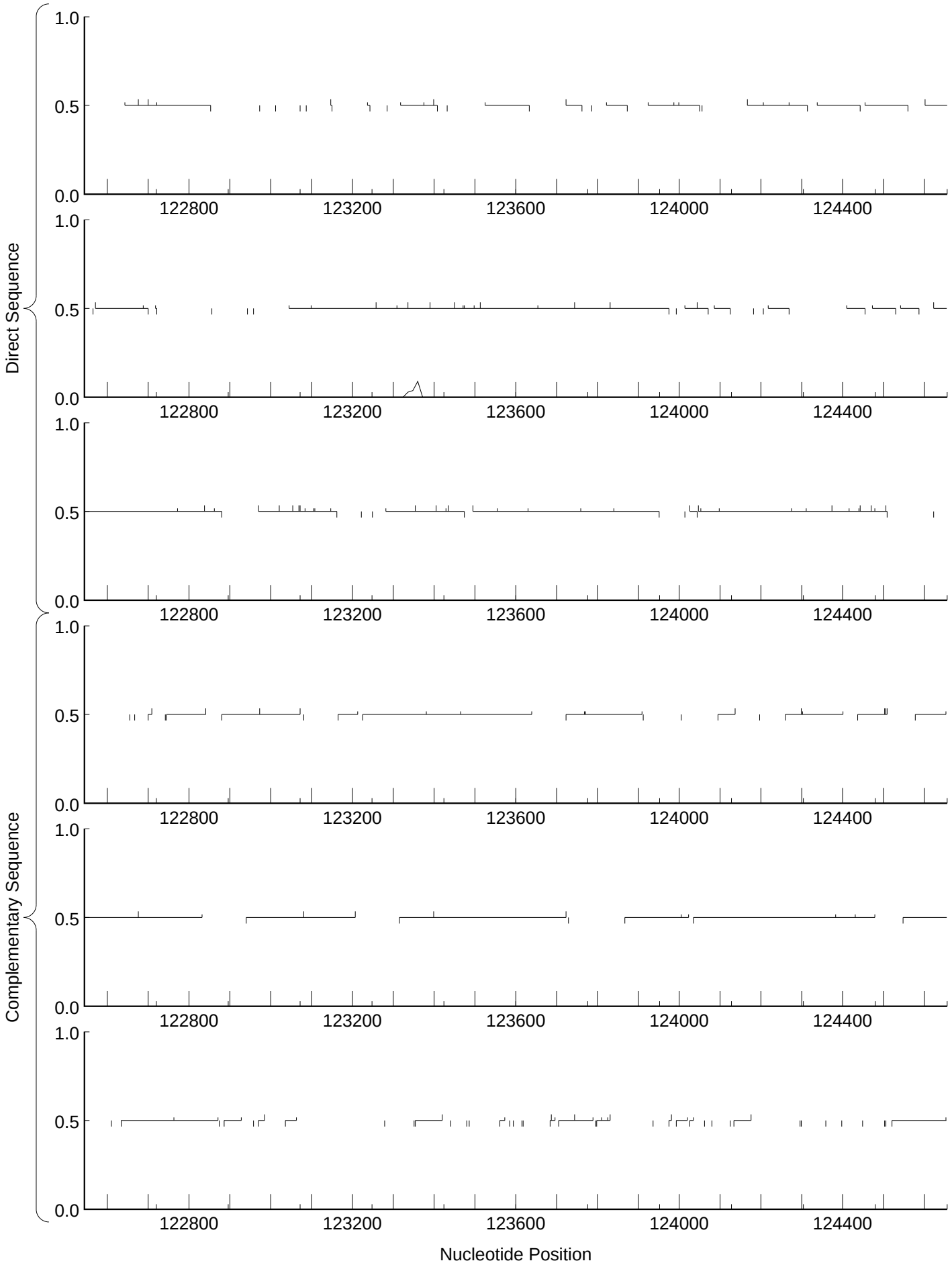
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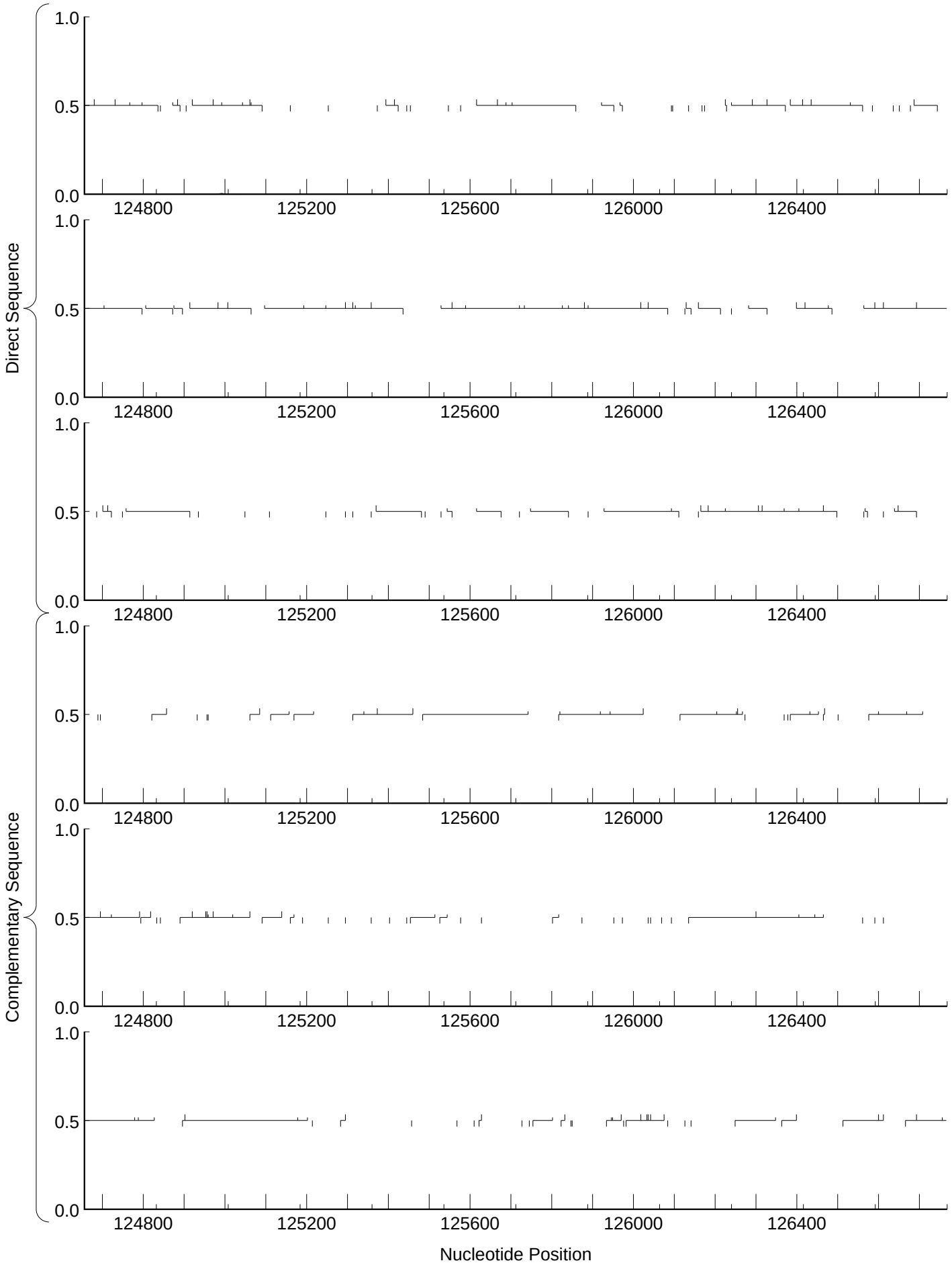
phage Gilson complete sequence, 128338 bp including 766 bp terminal repeat, Cluster BK1, Order 4, Window 96, Step 12, 59/62



phage Gilson complete sequence, 128338 bp including 766 bp terminal repeat, Cluster BK1, Order 4, Window 96, Step 12, 60/62



phage Gilson complete sequence, 128338 bp including 766 bp terminal repeat, Cluster BK1, Order 4, Window 96, Step 12, 61/62



phage Gilson complete sequence, 128338 bp including 766 bp terminal repeat, Cluster BK1, Order 4, Window 96, Step 12, 62/62

