GeneMark

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

PROGRAM INFORMATION

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Sequence         :  Streptomyces phage StarPlatinum complete sequence, 133886 bp including 12199 bp terminal re
Analysis Date    :  3/29/24 at 2:47:06
Pages            :  65
Sequence Length  :  133886 bp
GC Content       :   49.47%

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

PS-Version       :  1.2

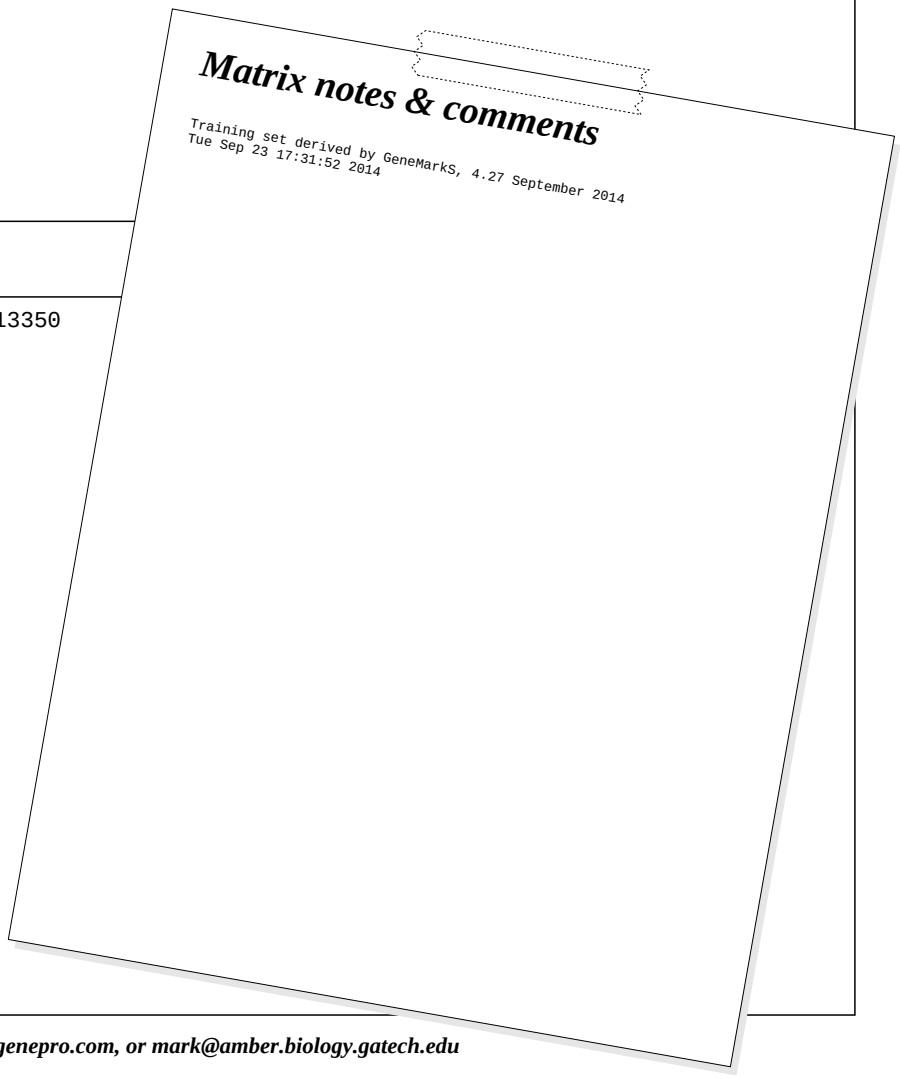
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MATRIX INFORMATION

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Author     :  -

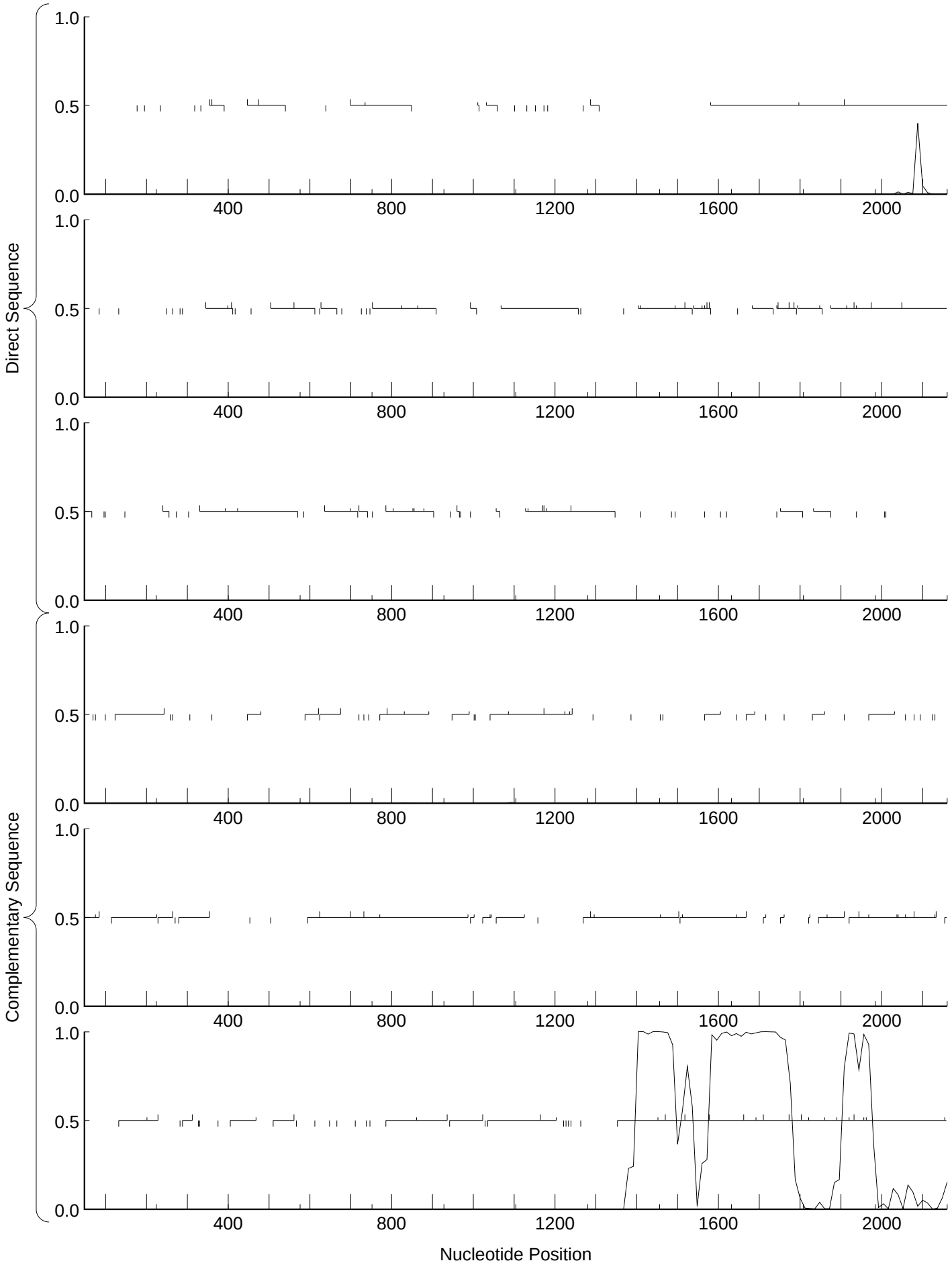
Order      :  4


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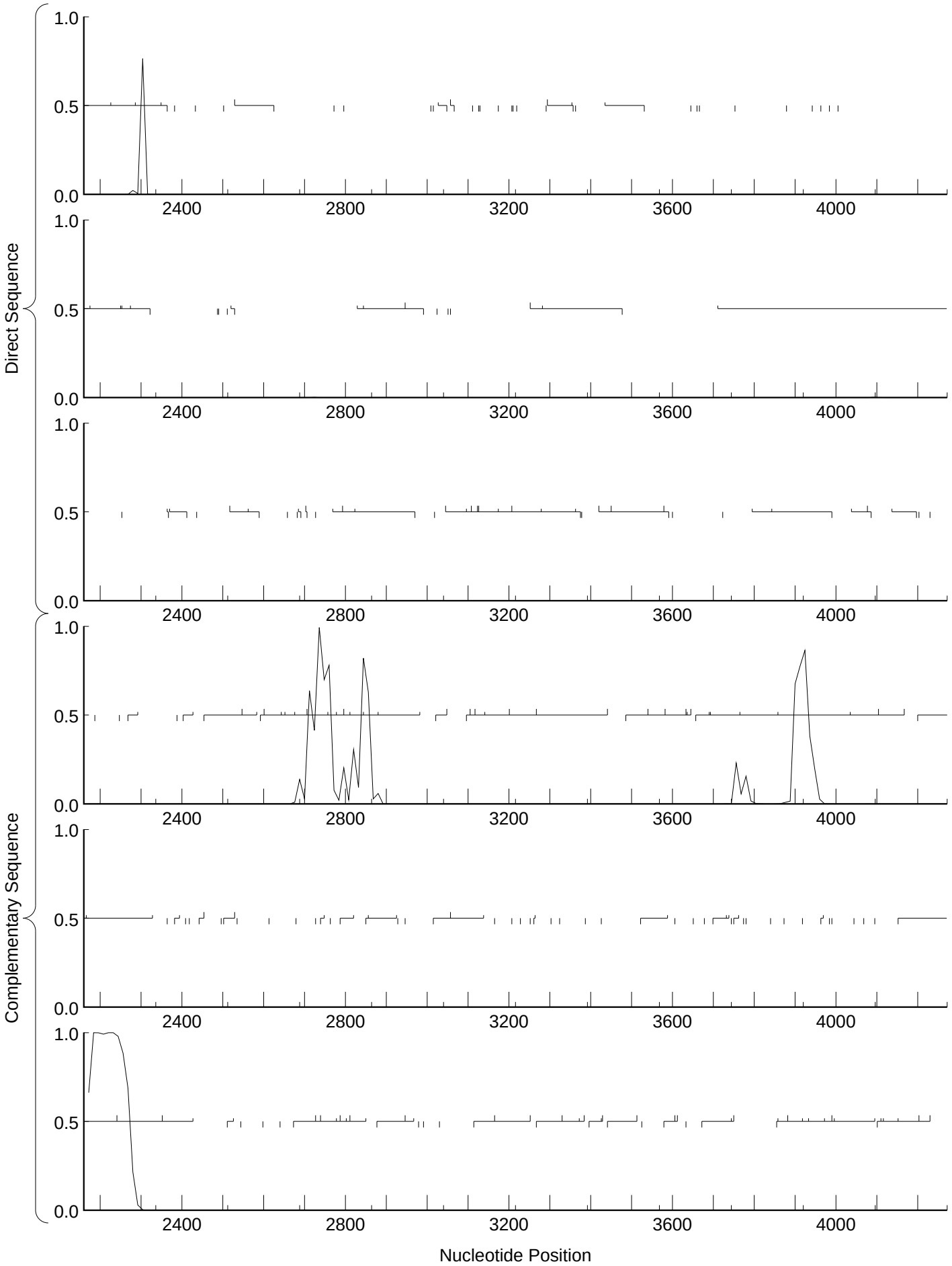


For help or information, send electronic mail to: custserv@genepro.com, or mark@amber.biology.gatech.edu

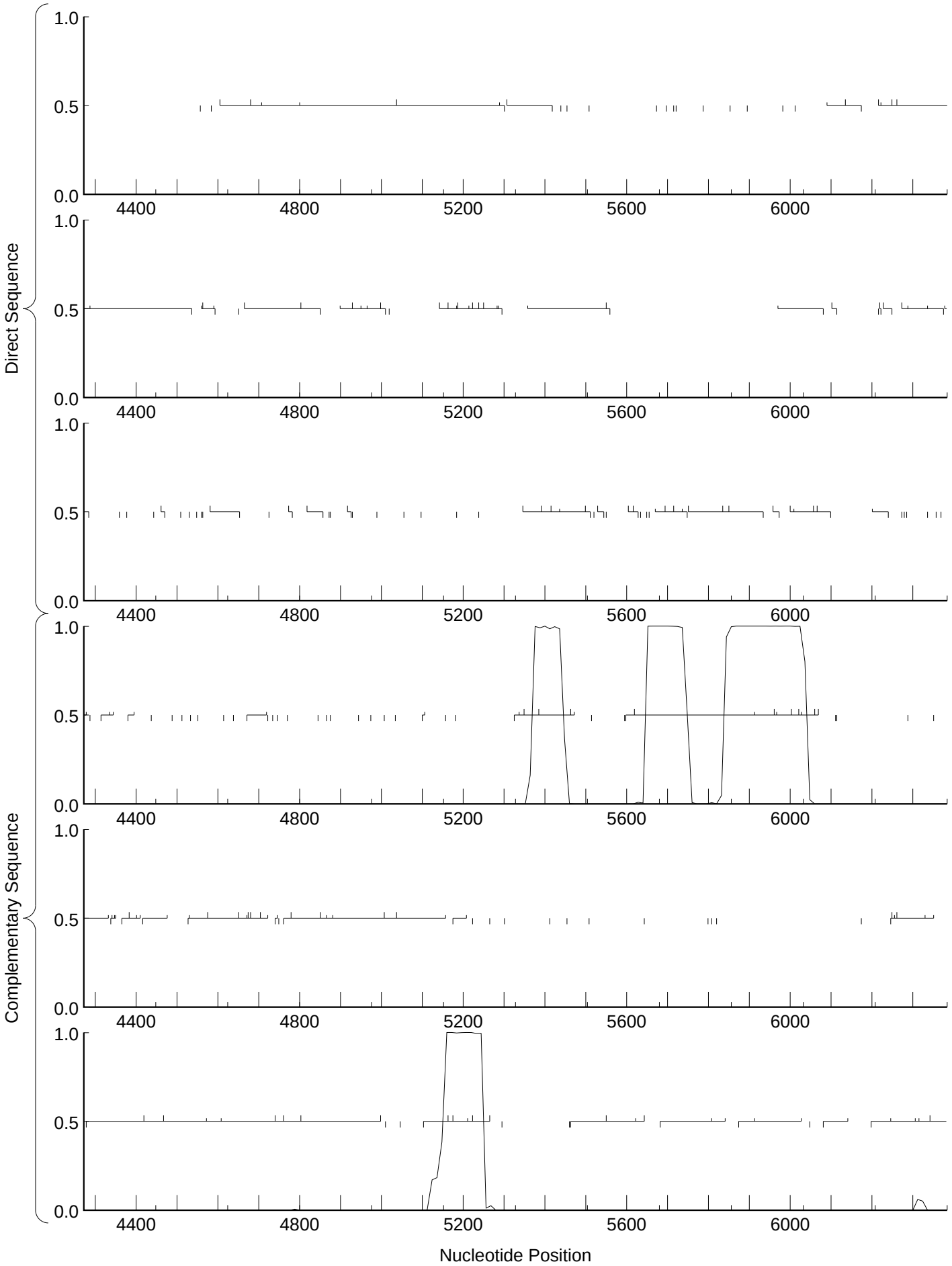
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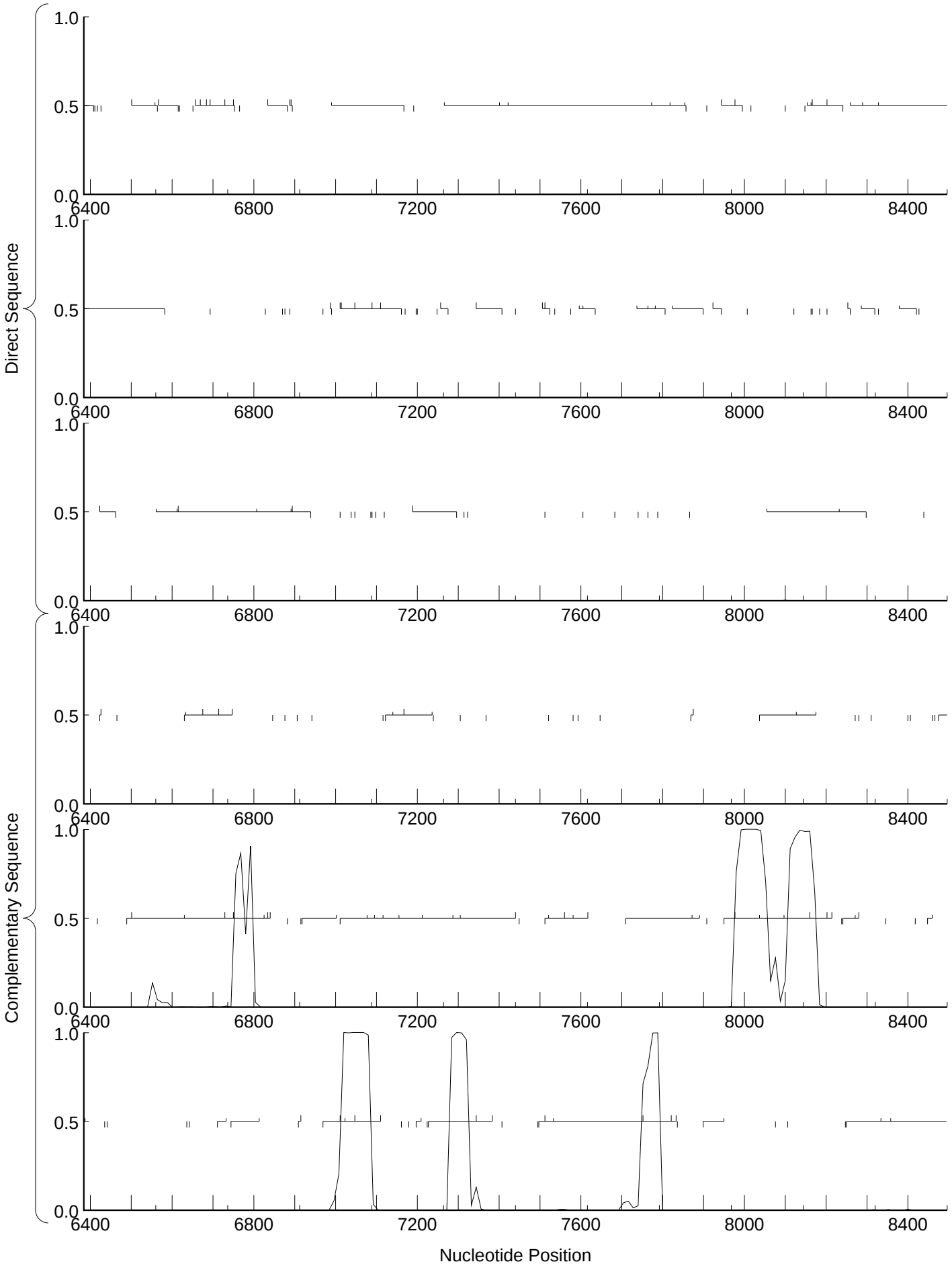
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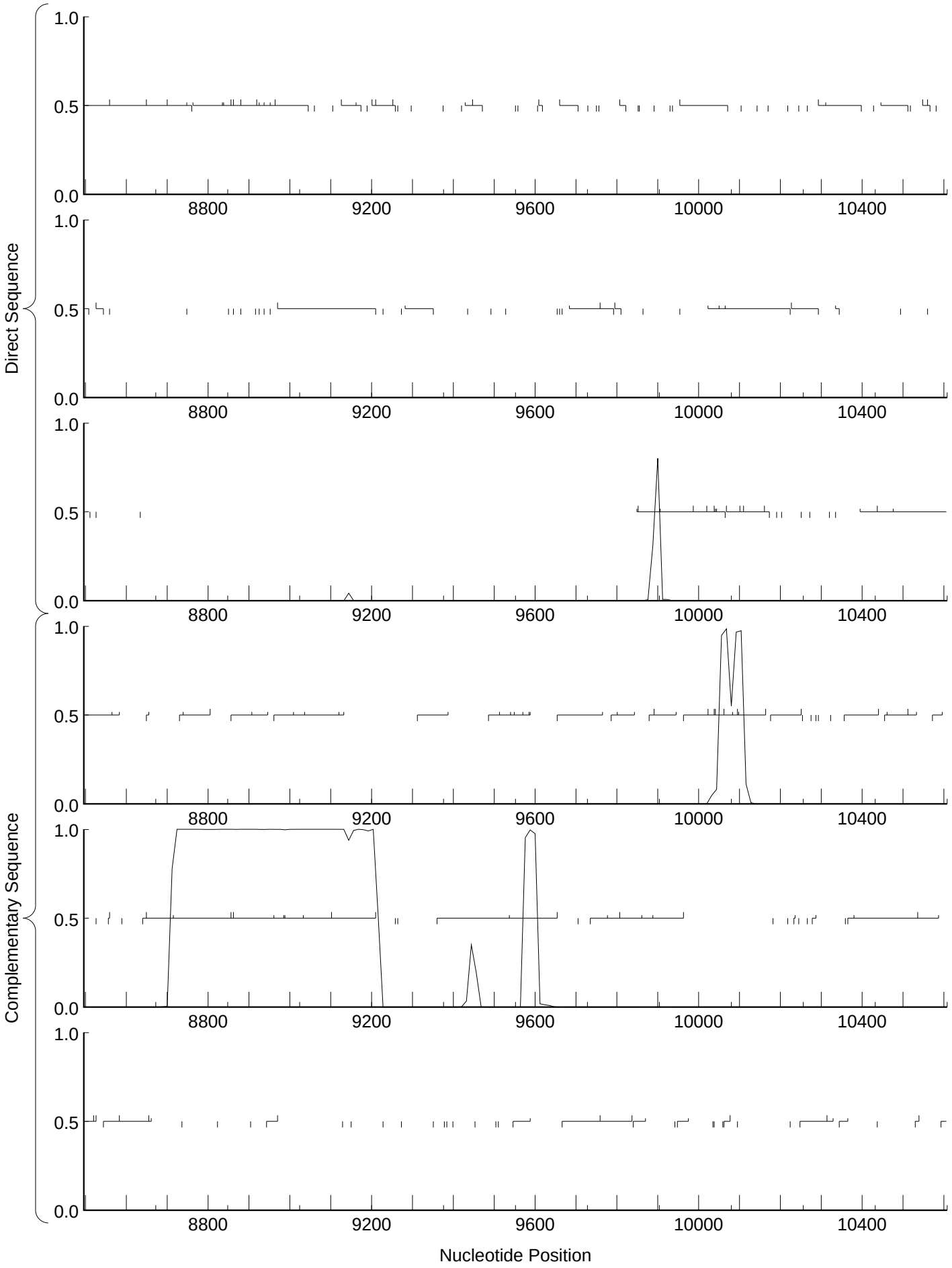
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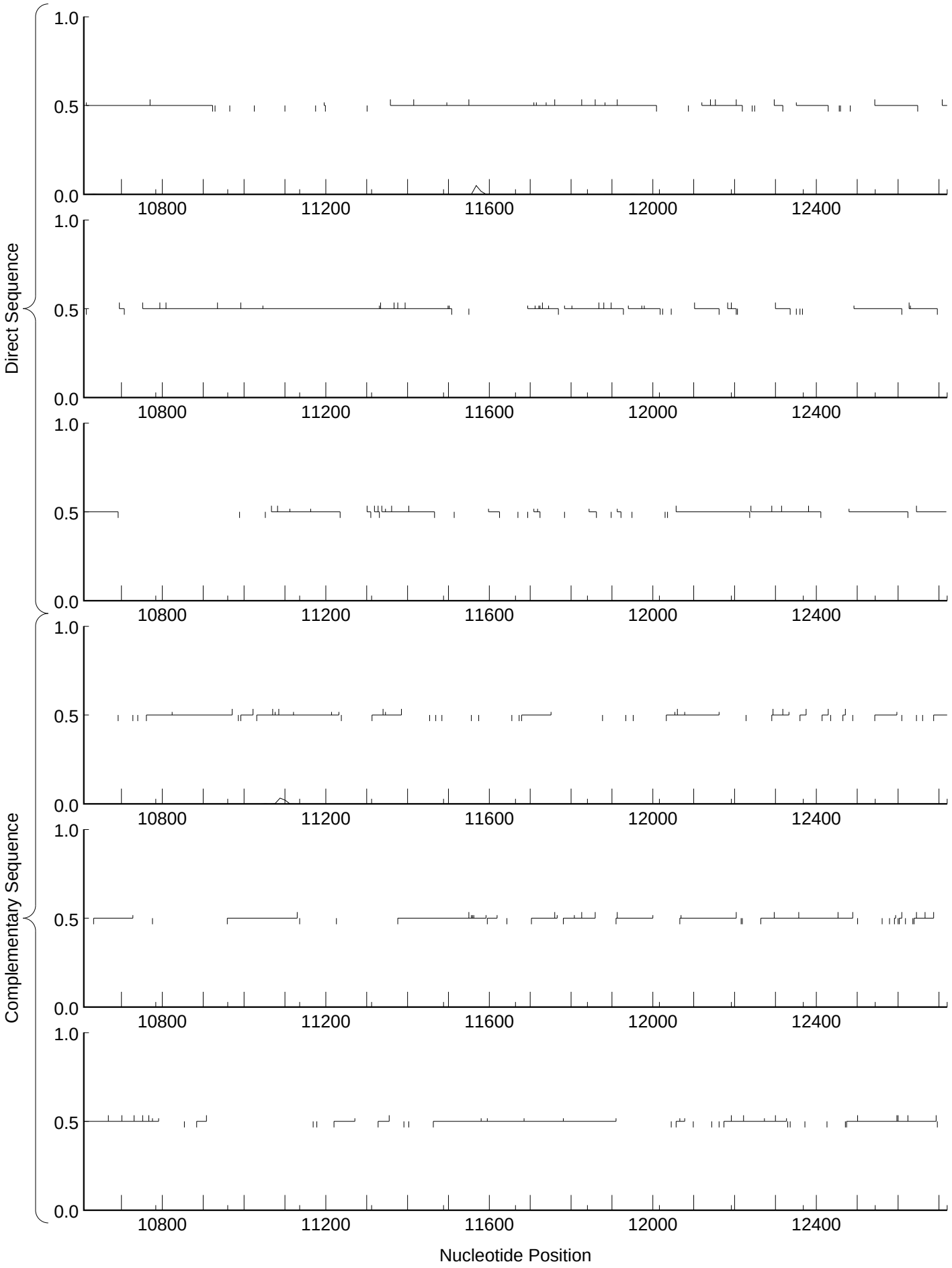
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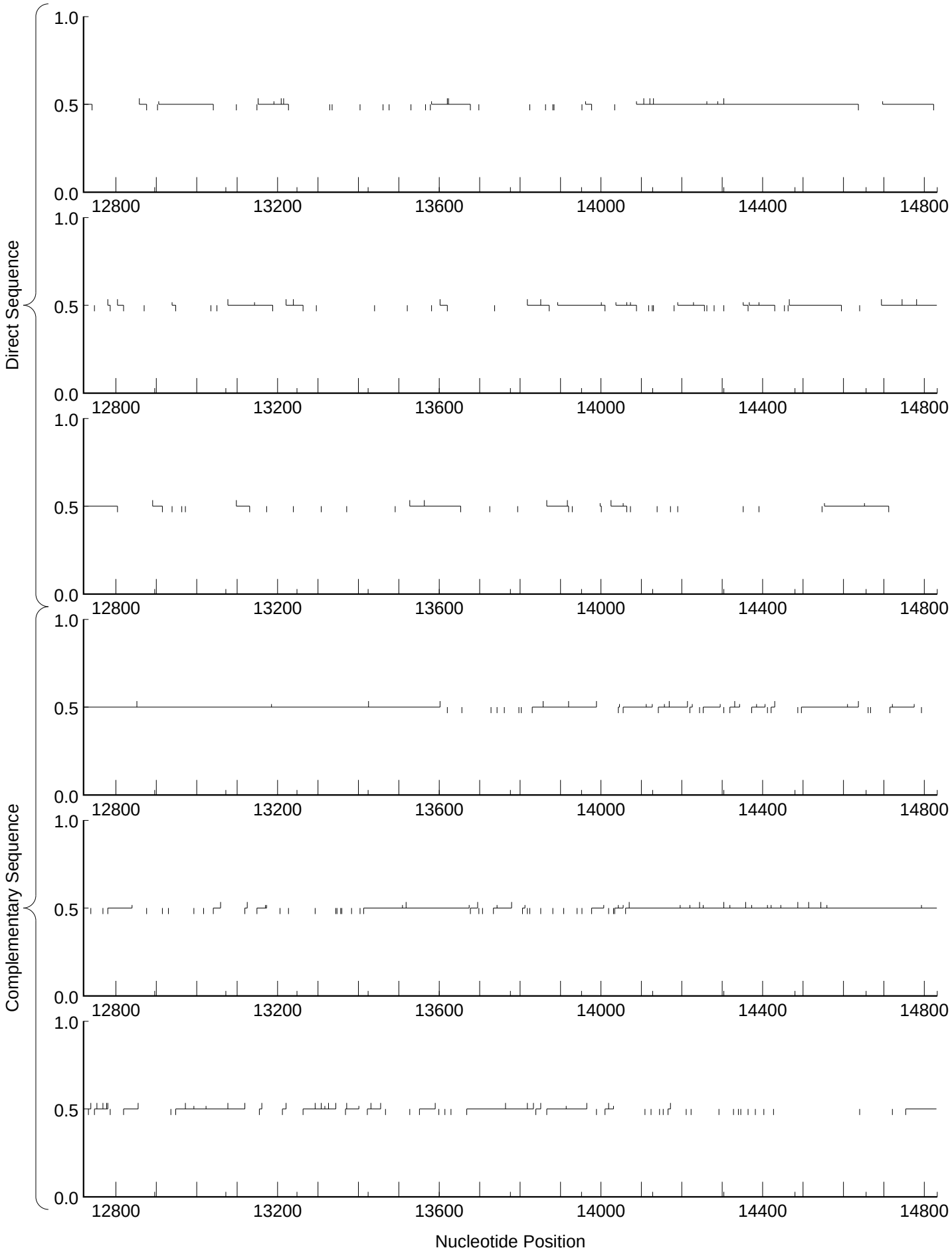
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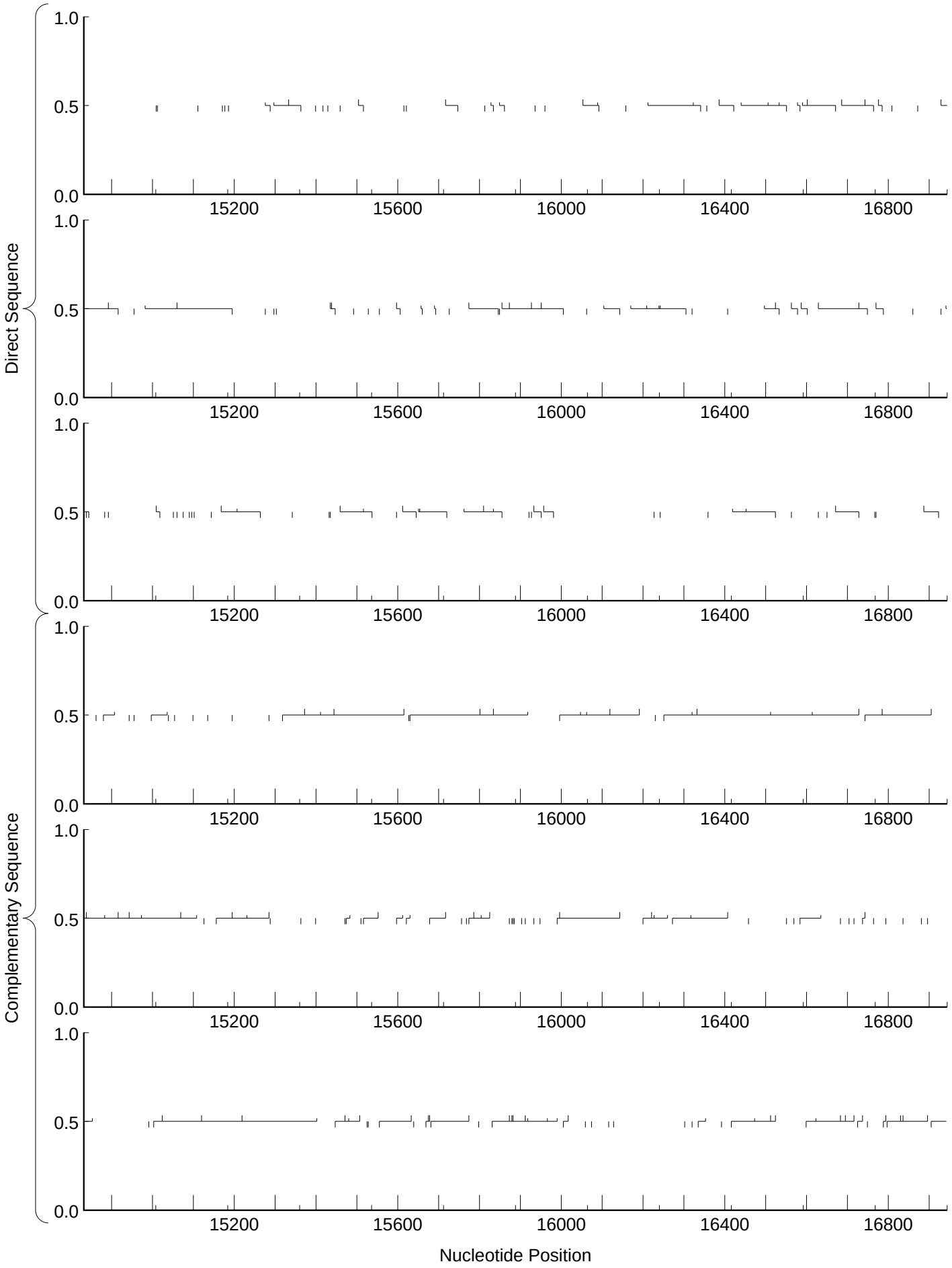
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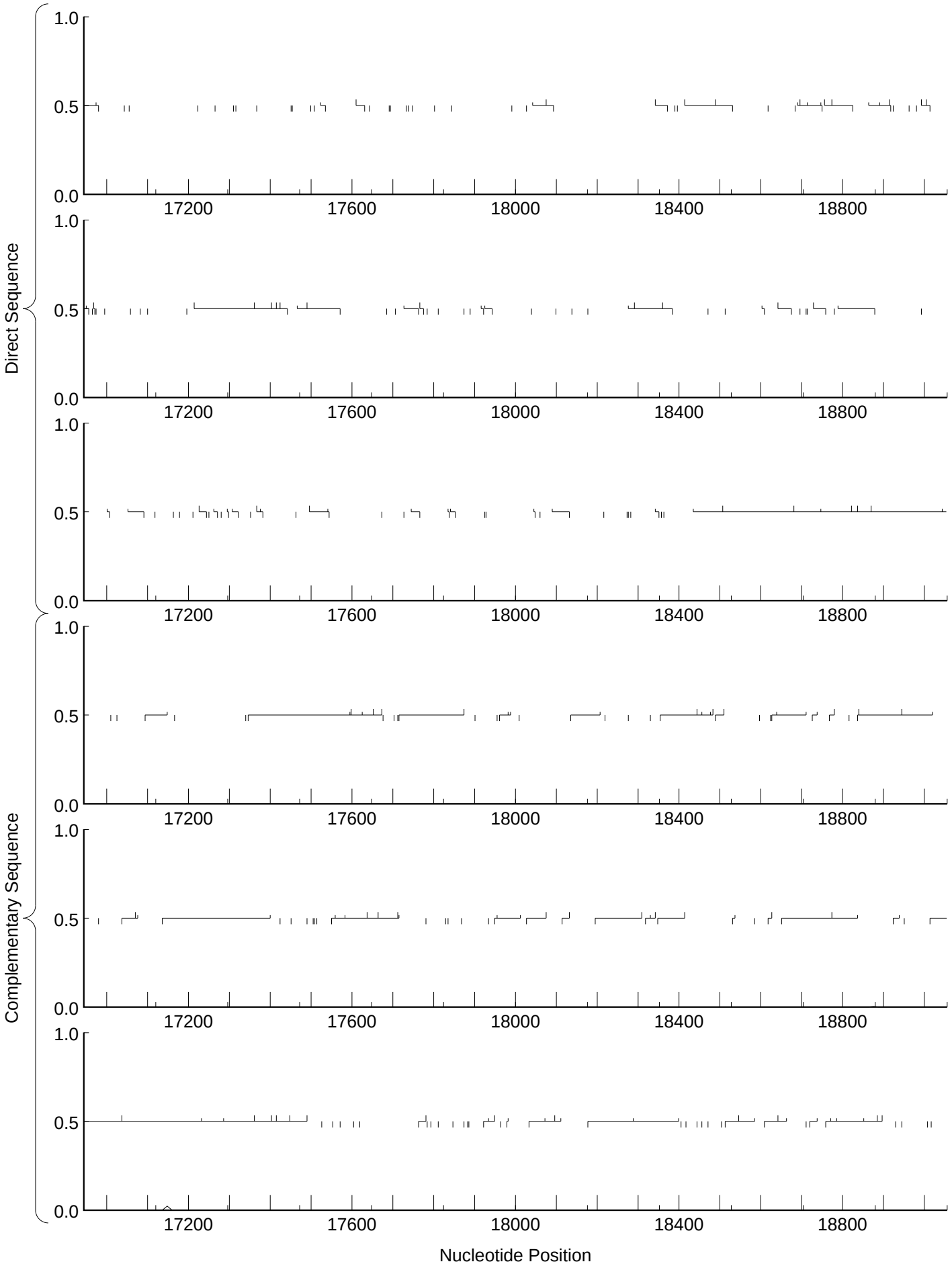
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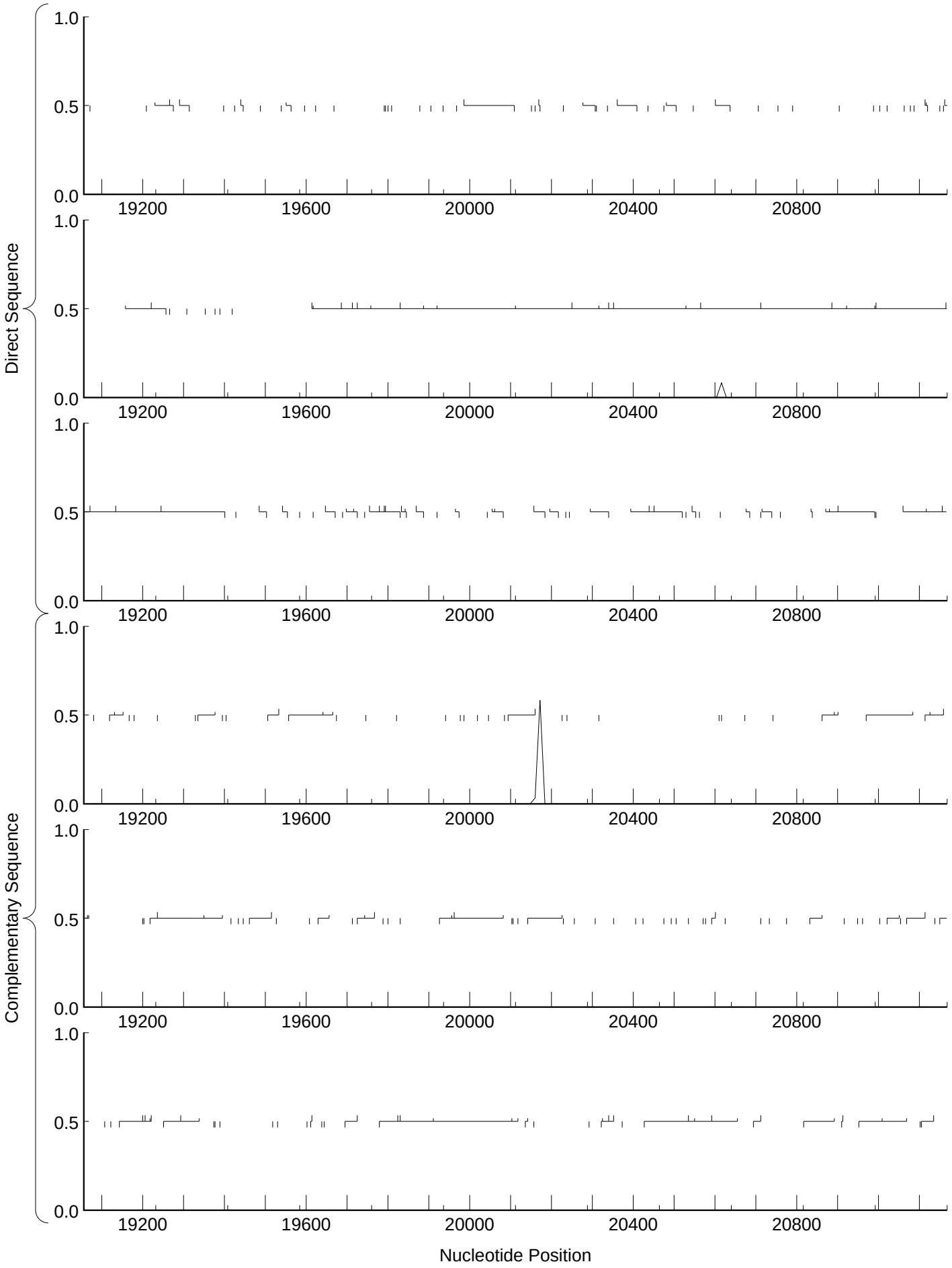
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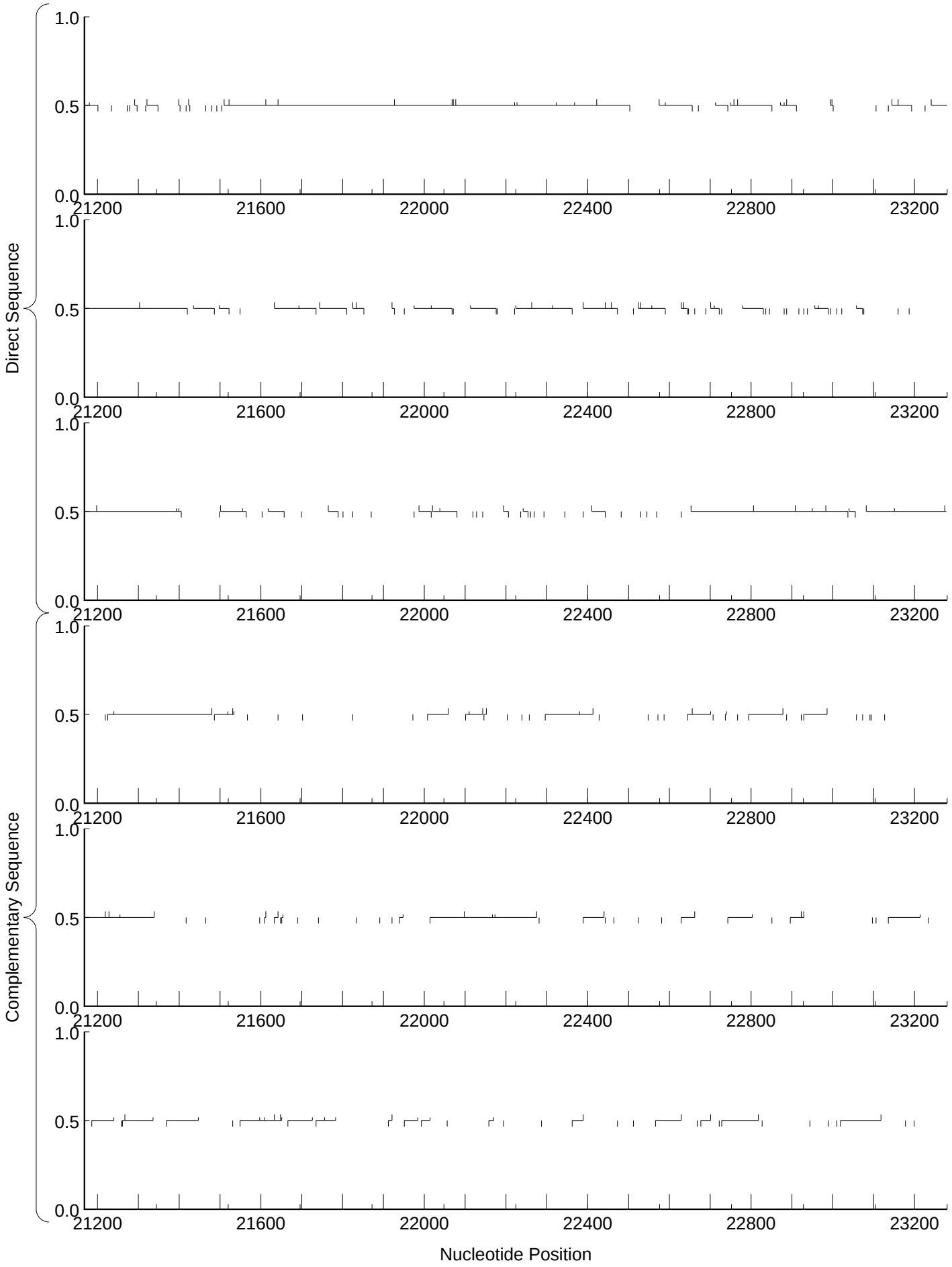
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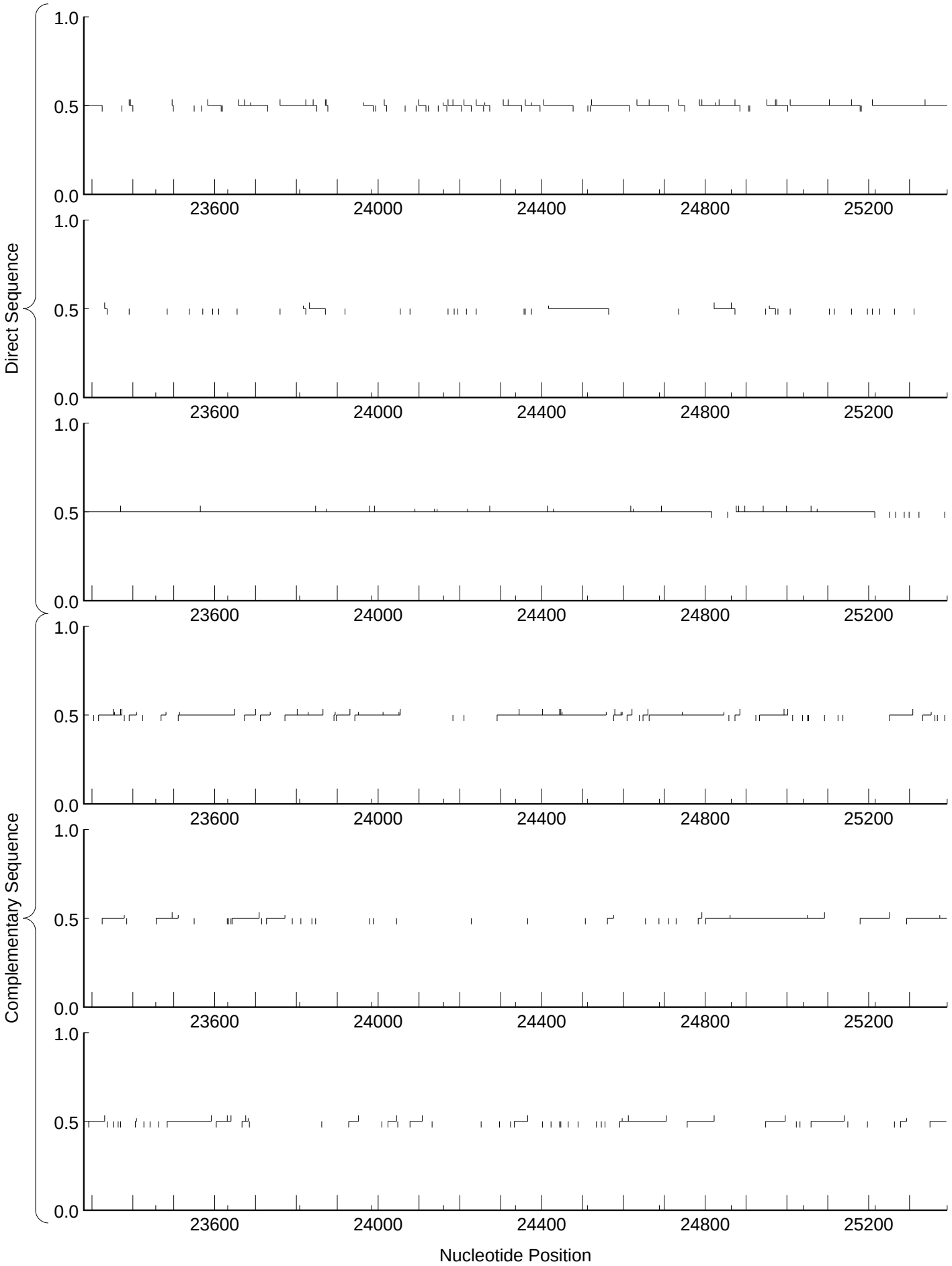
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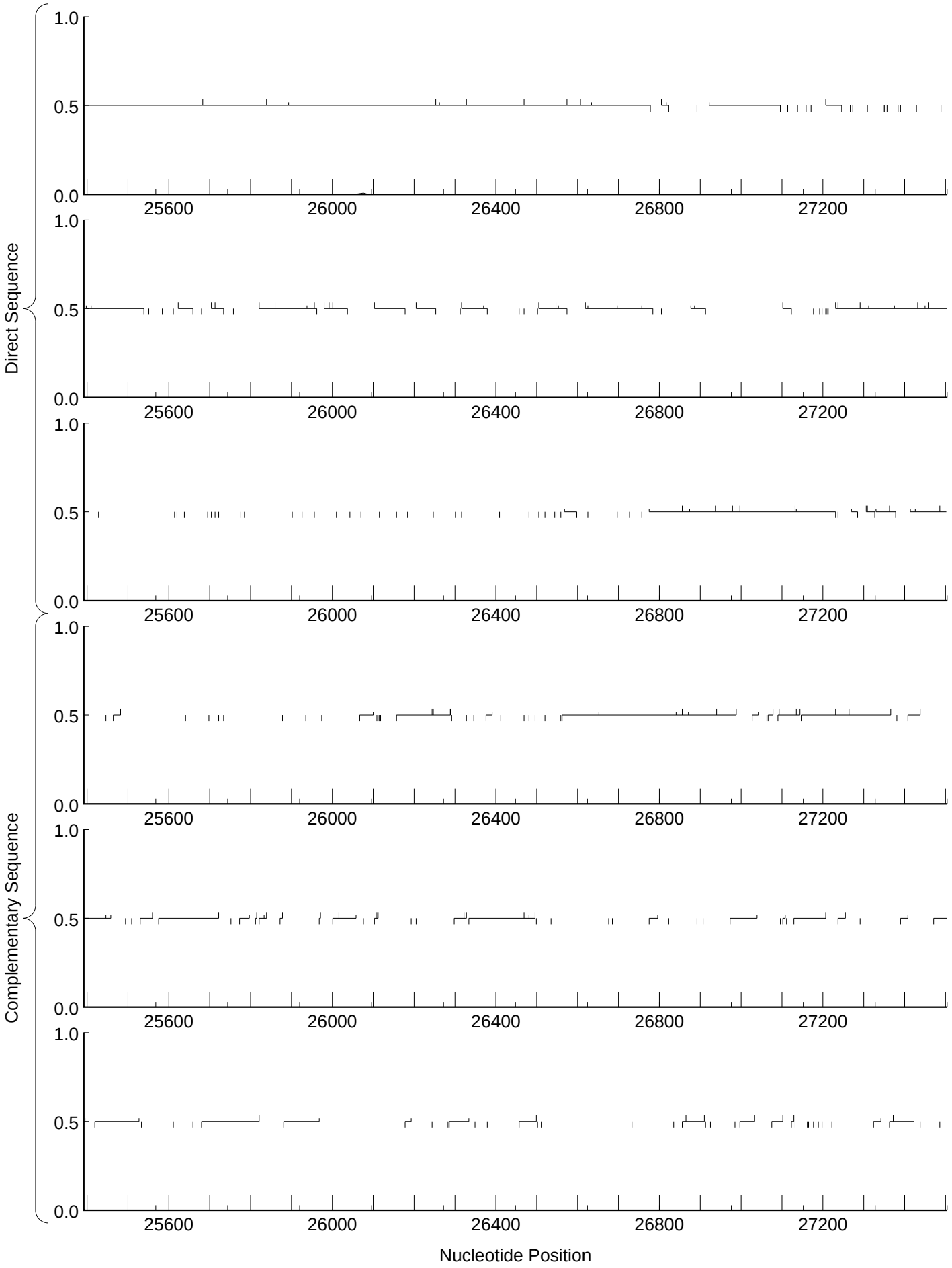
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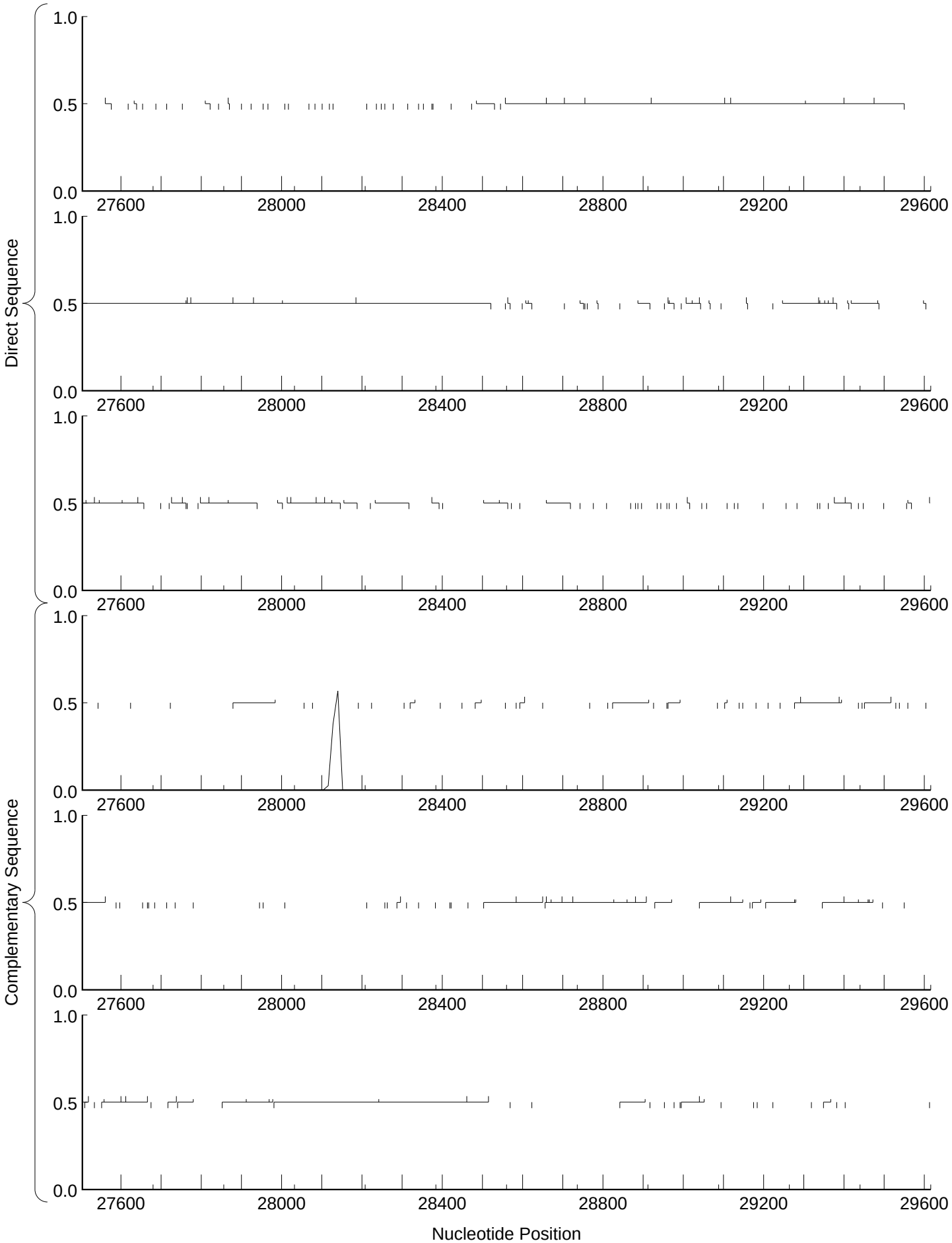
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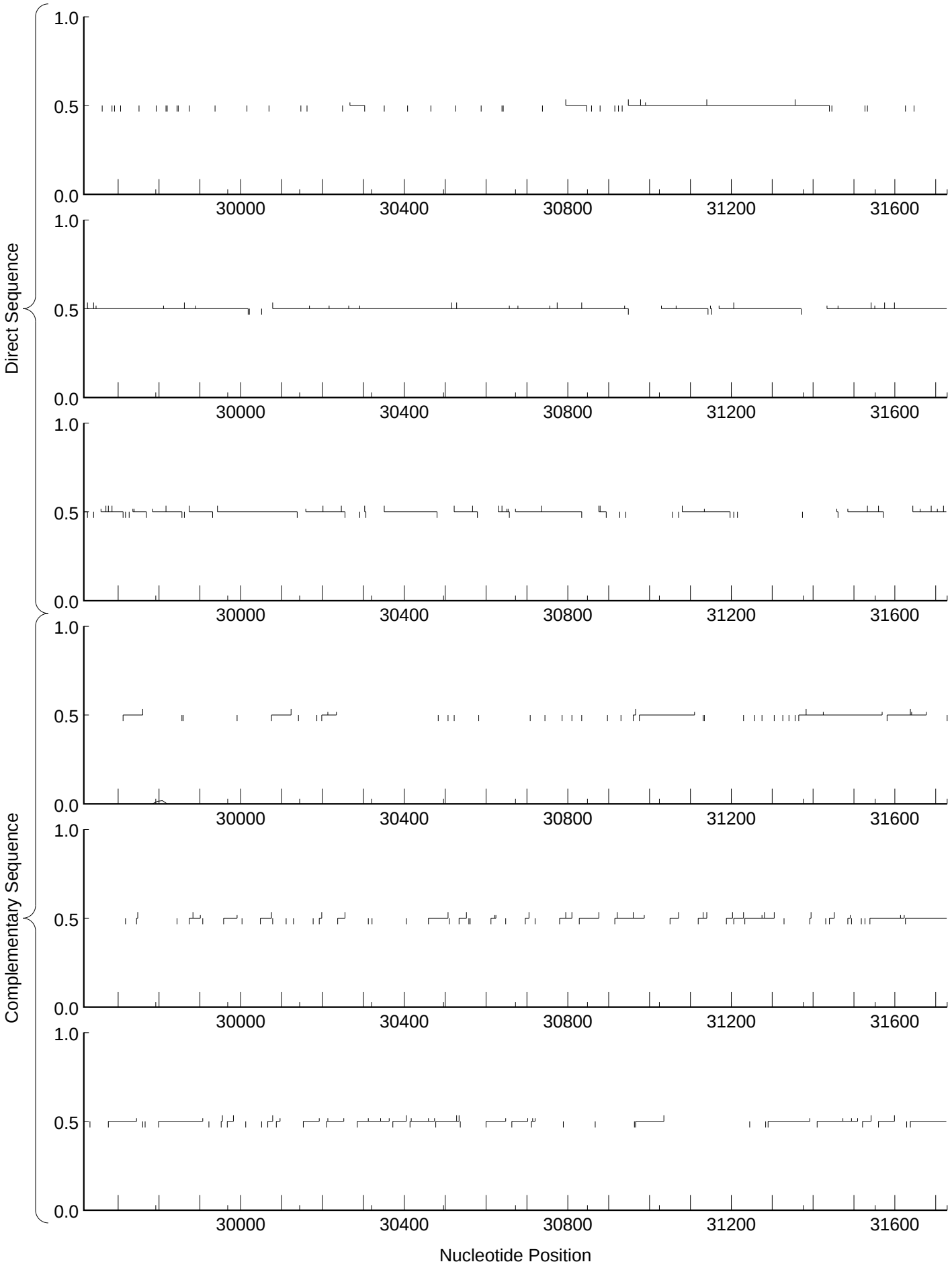
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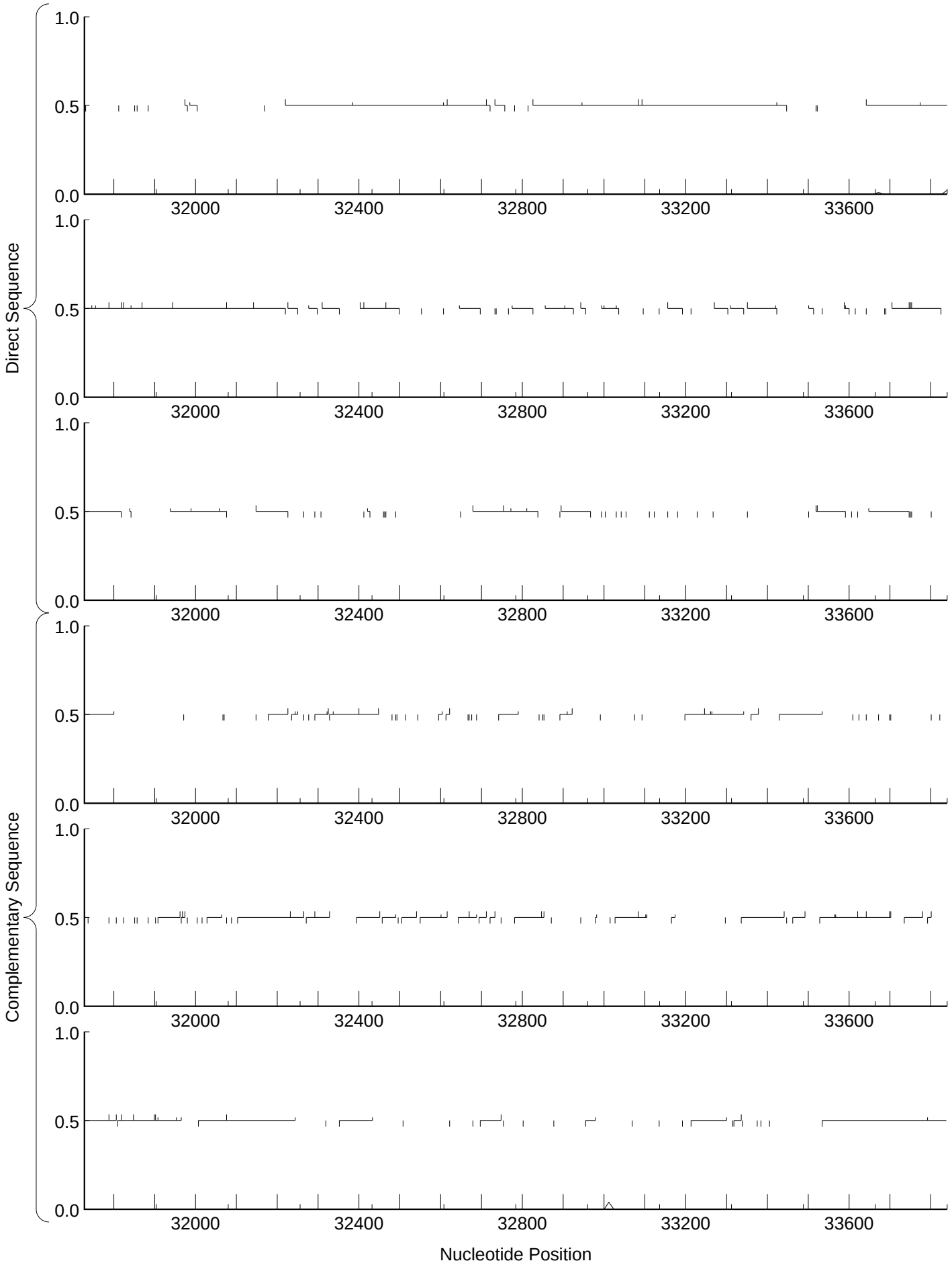
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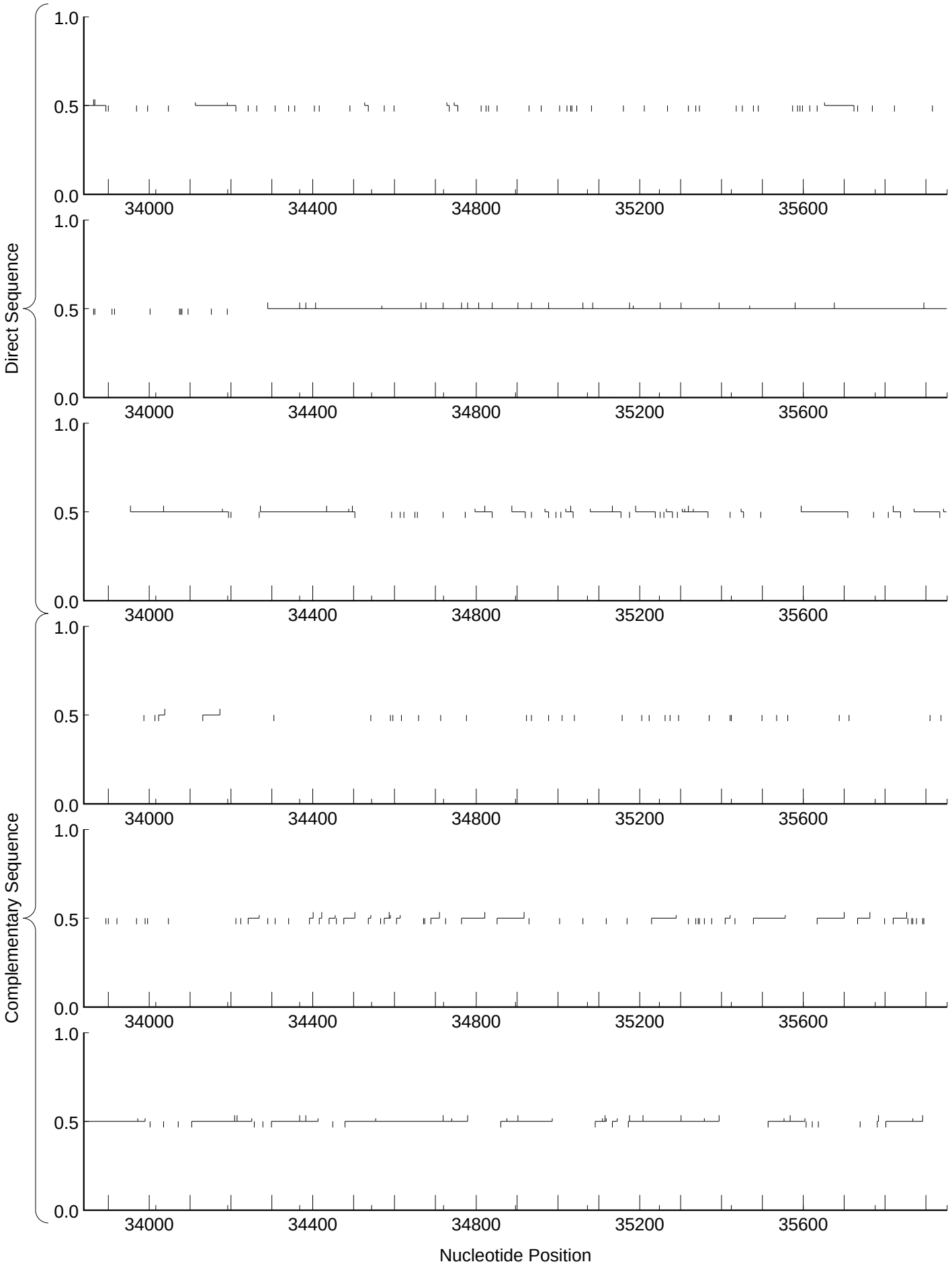
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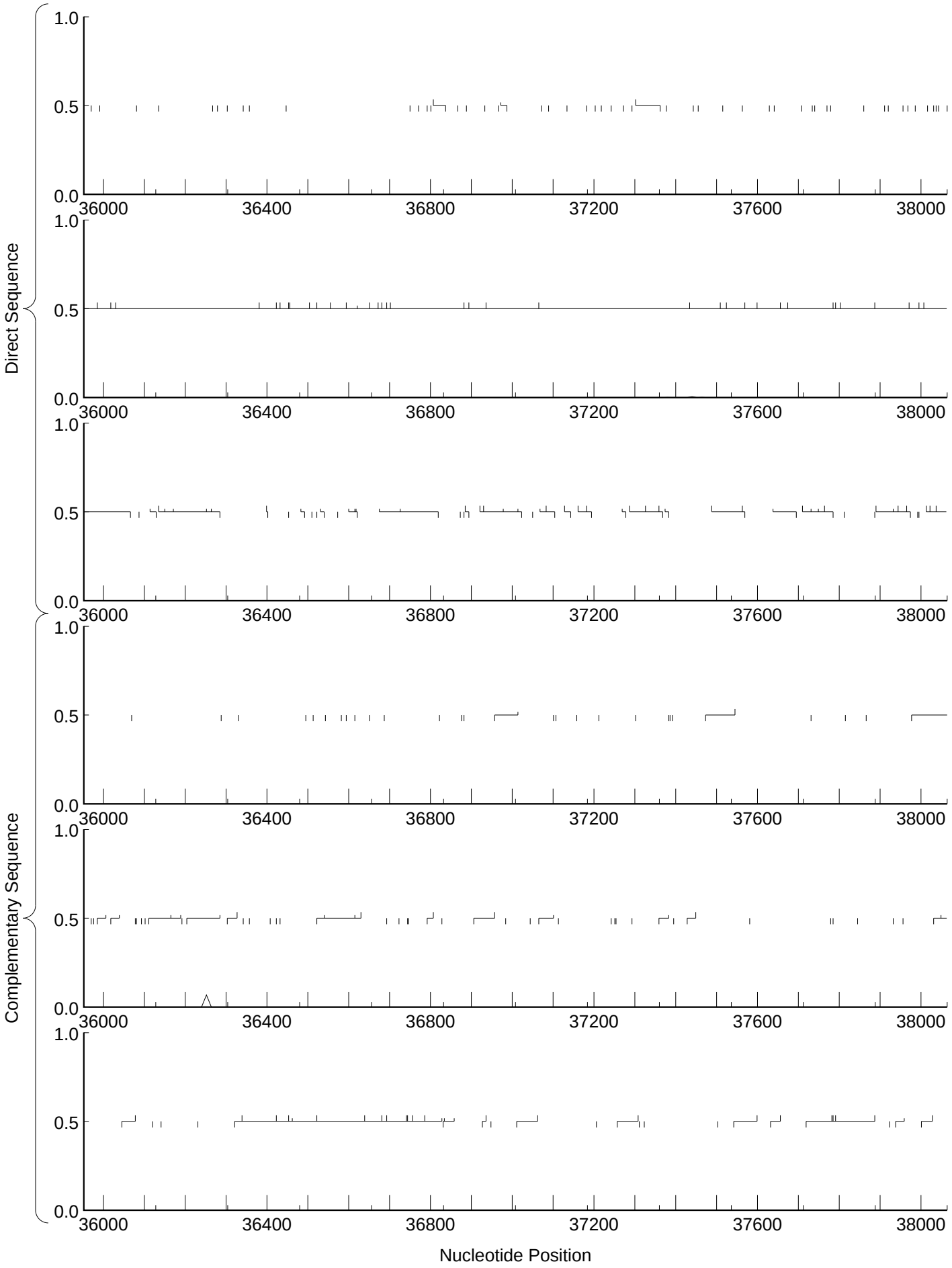
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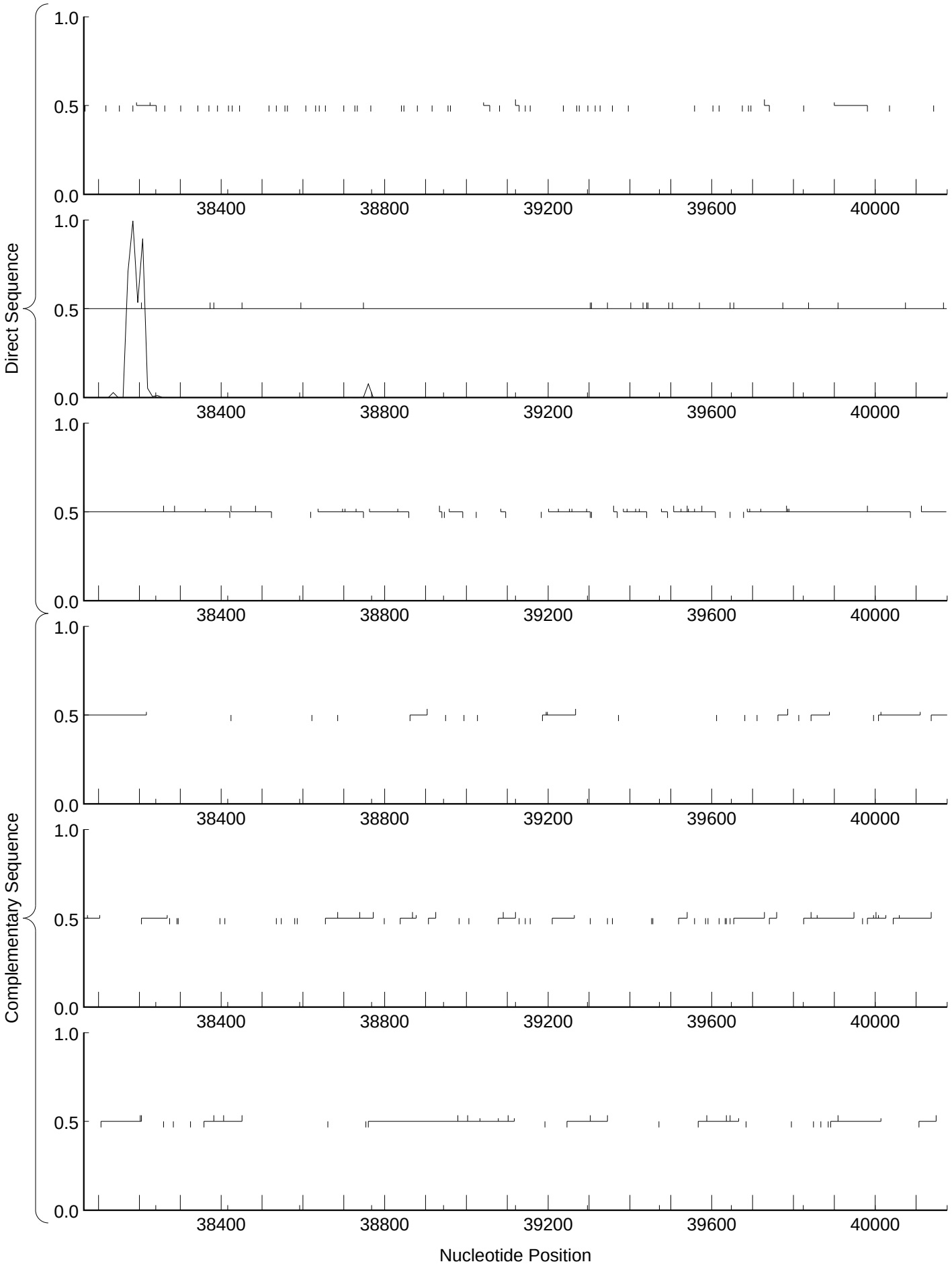
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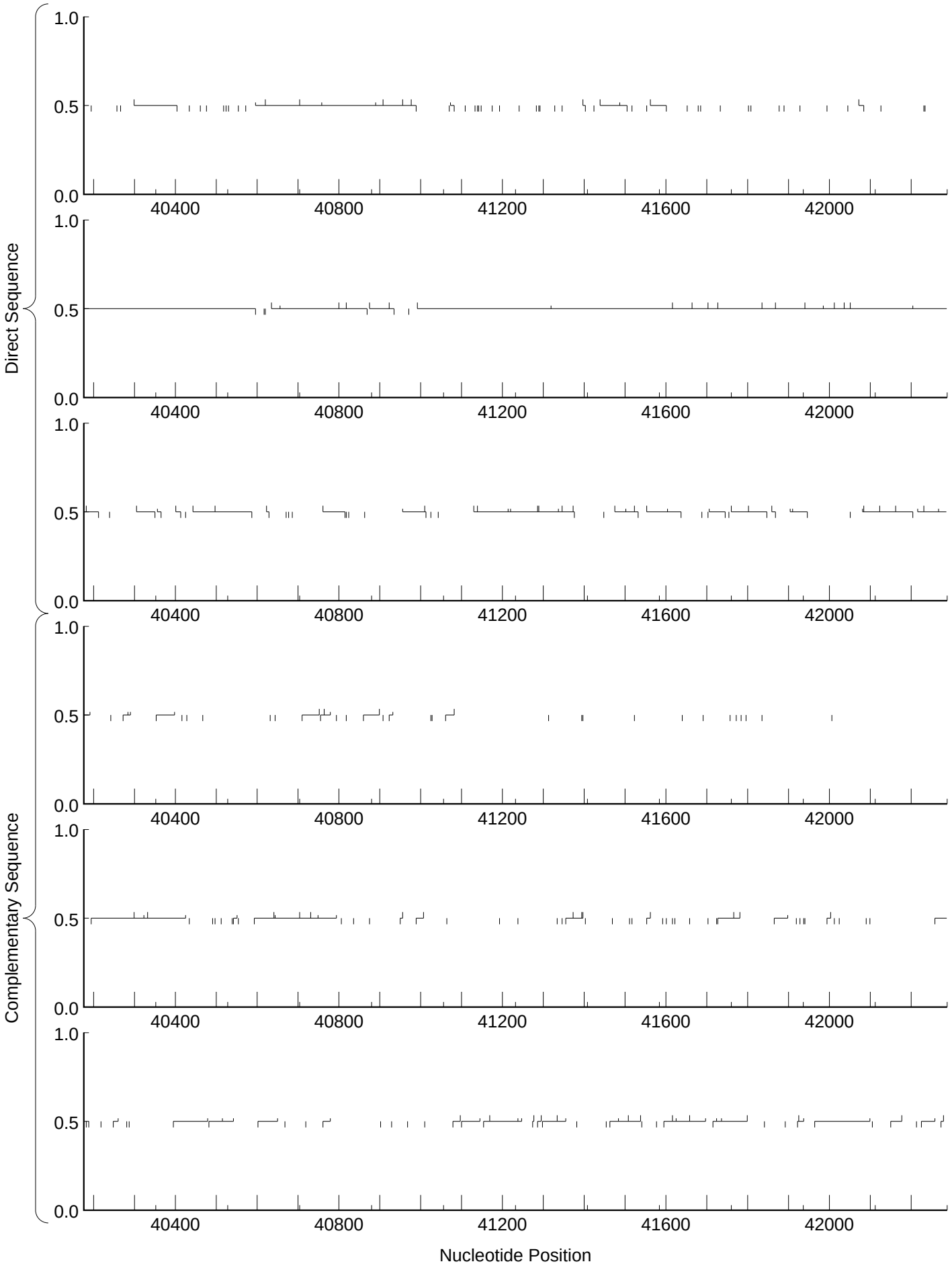
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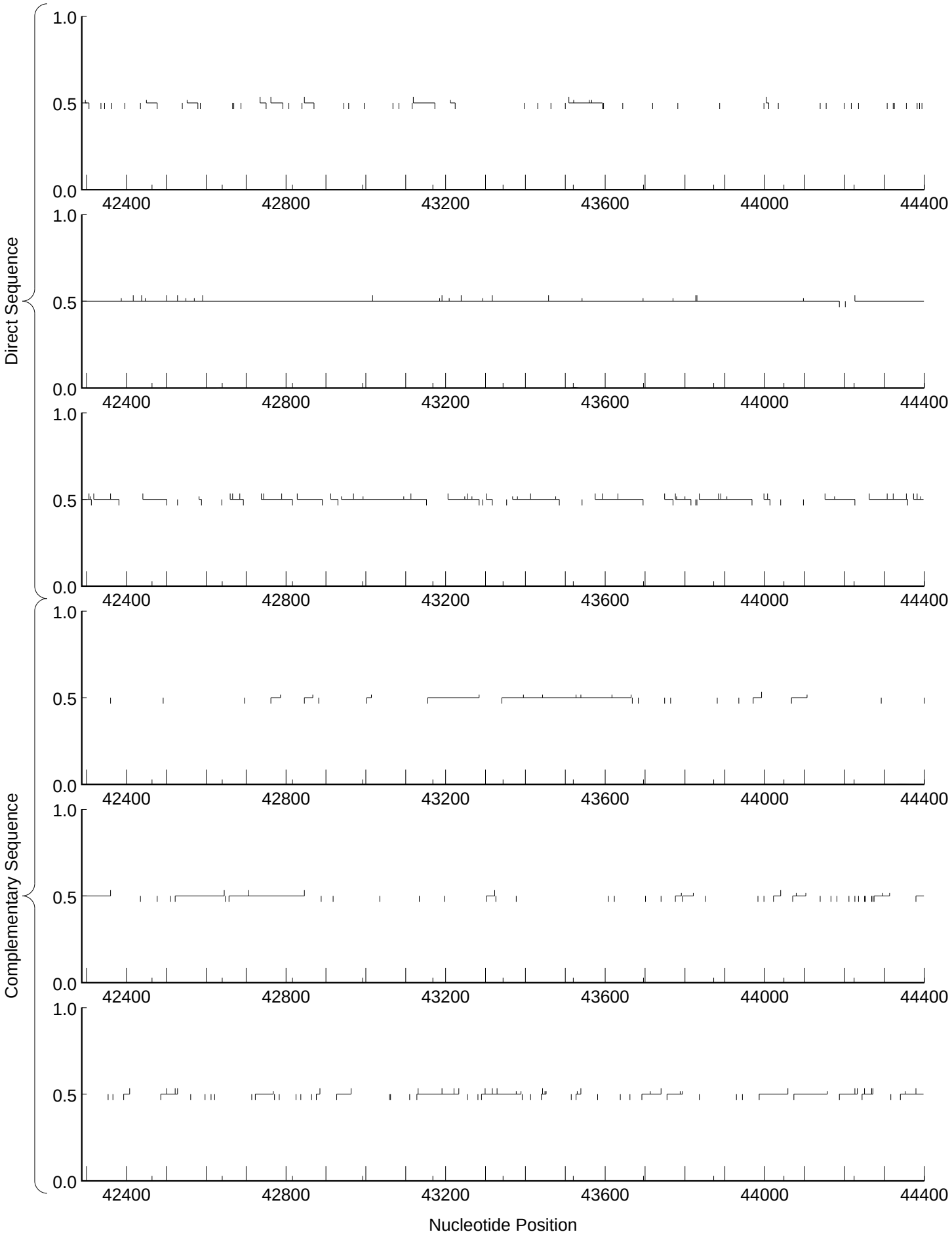
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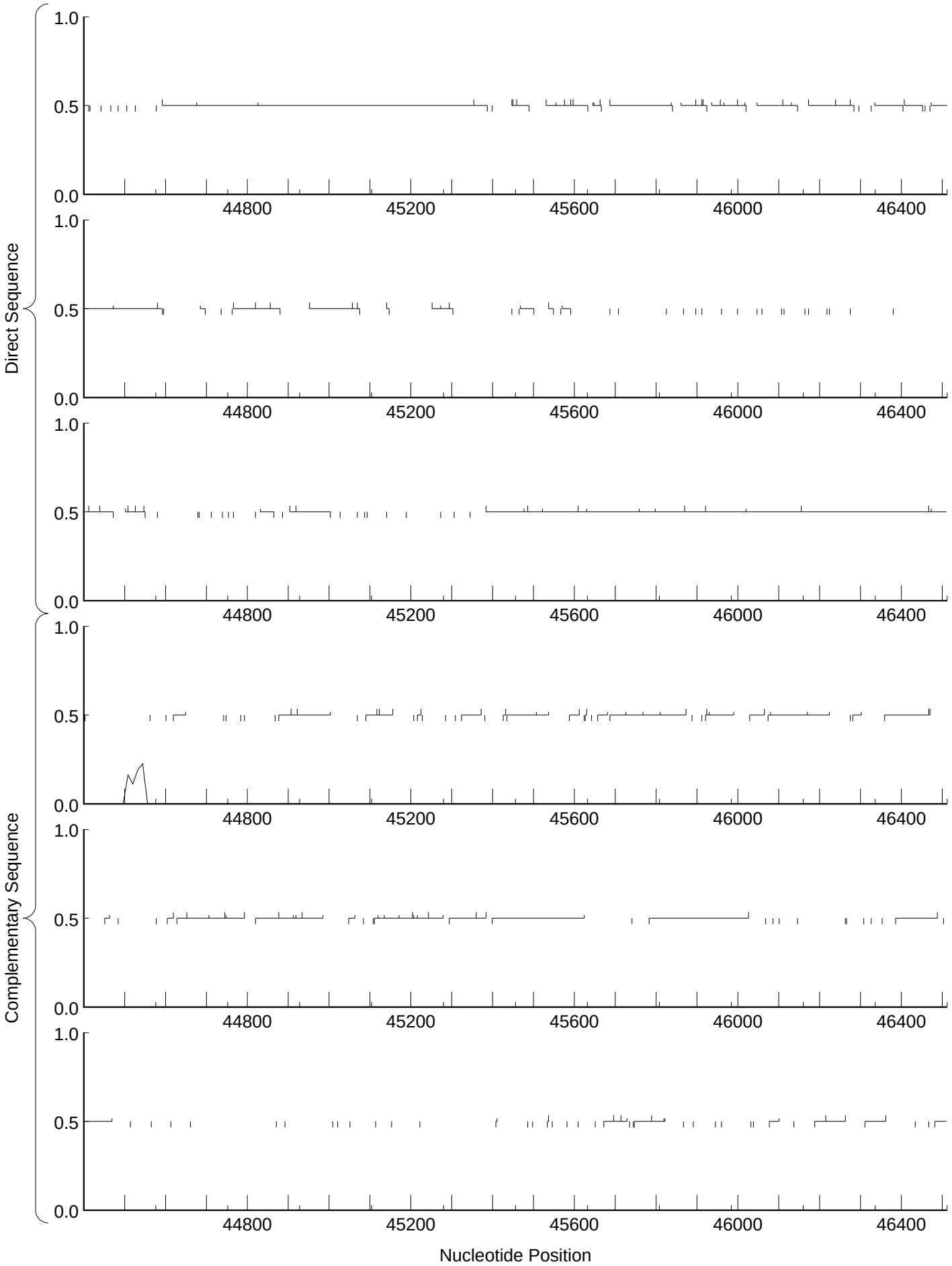
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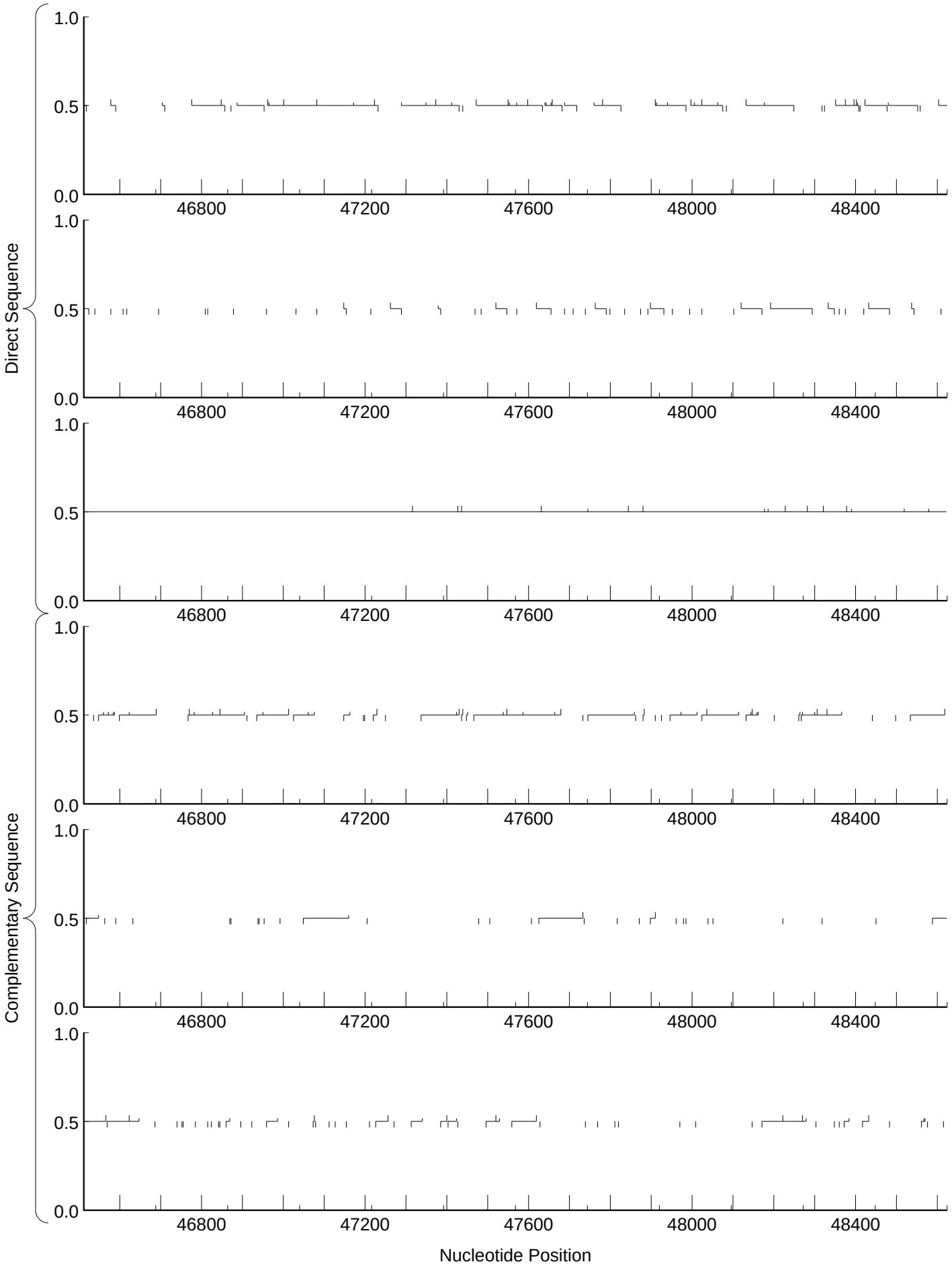
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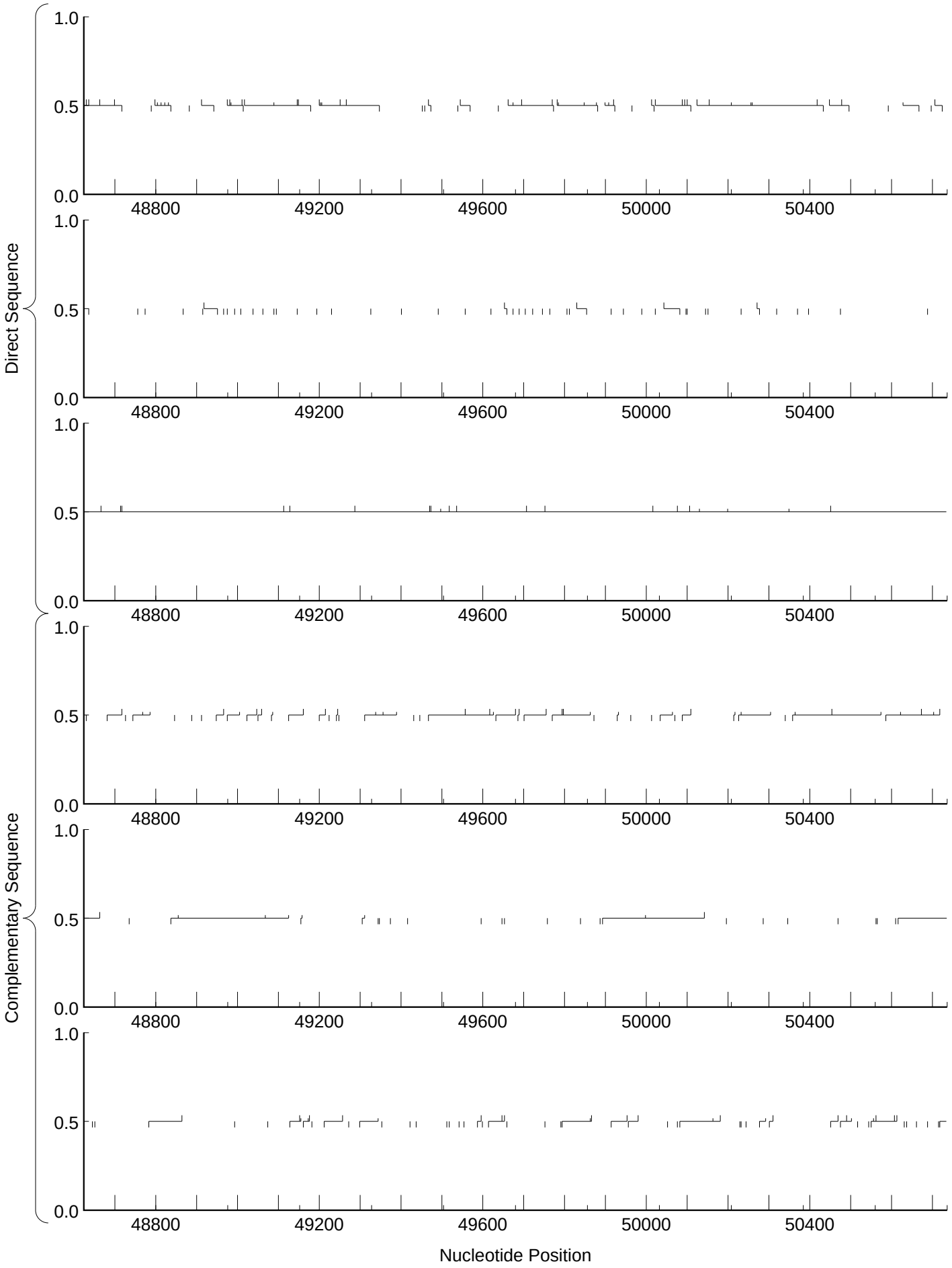
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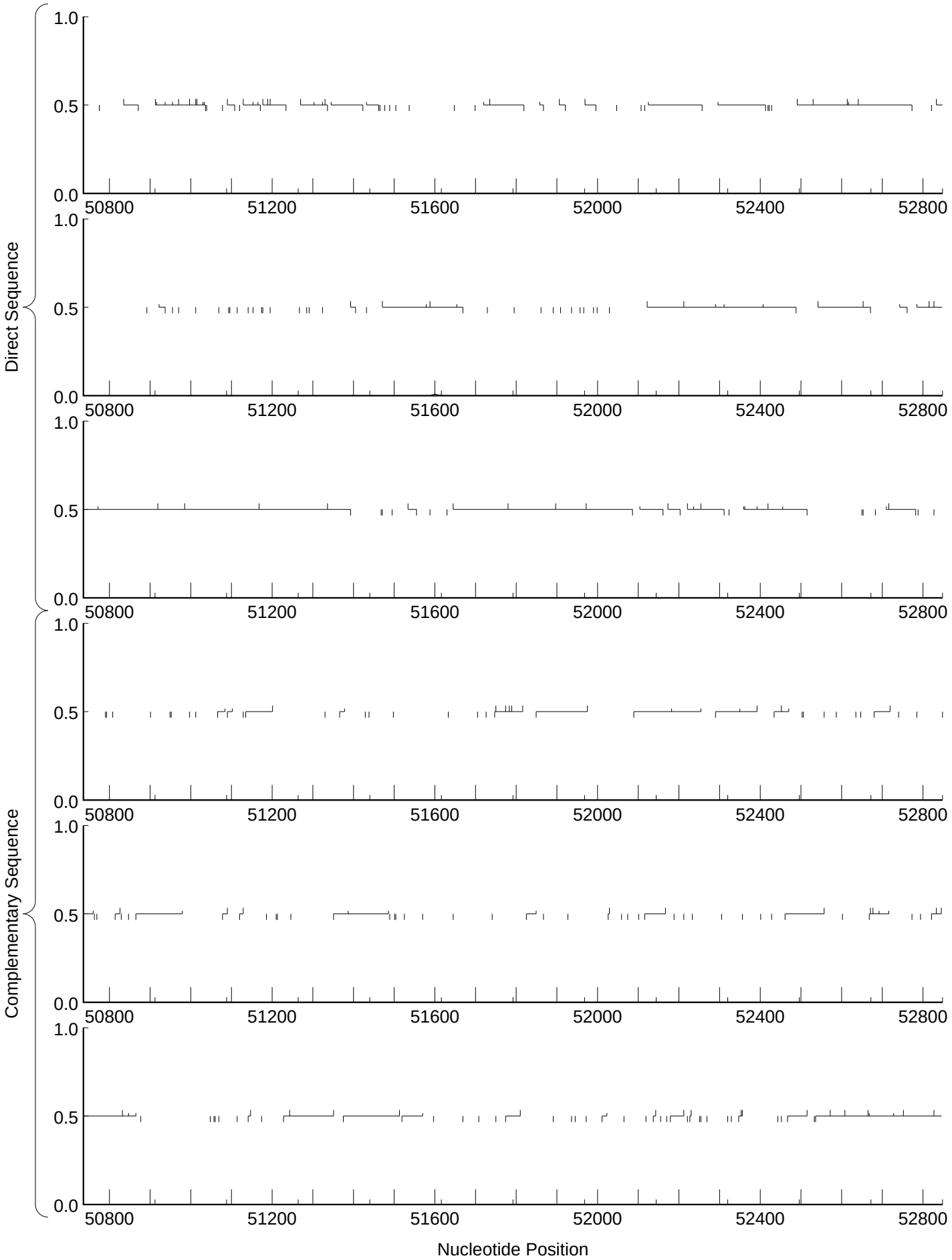
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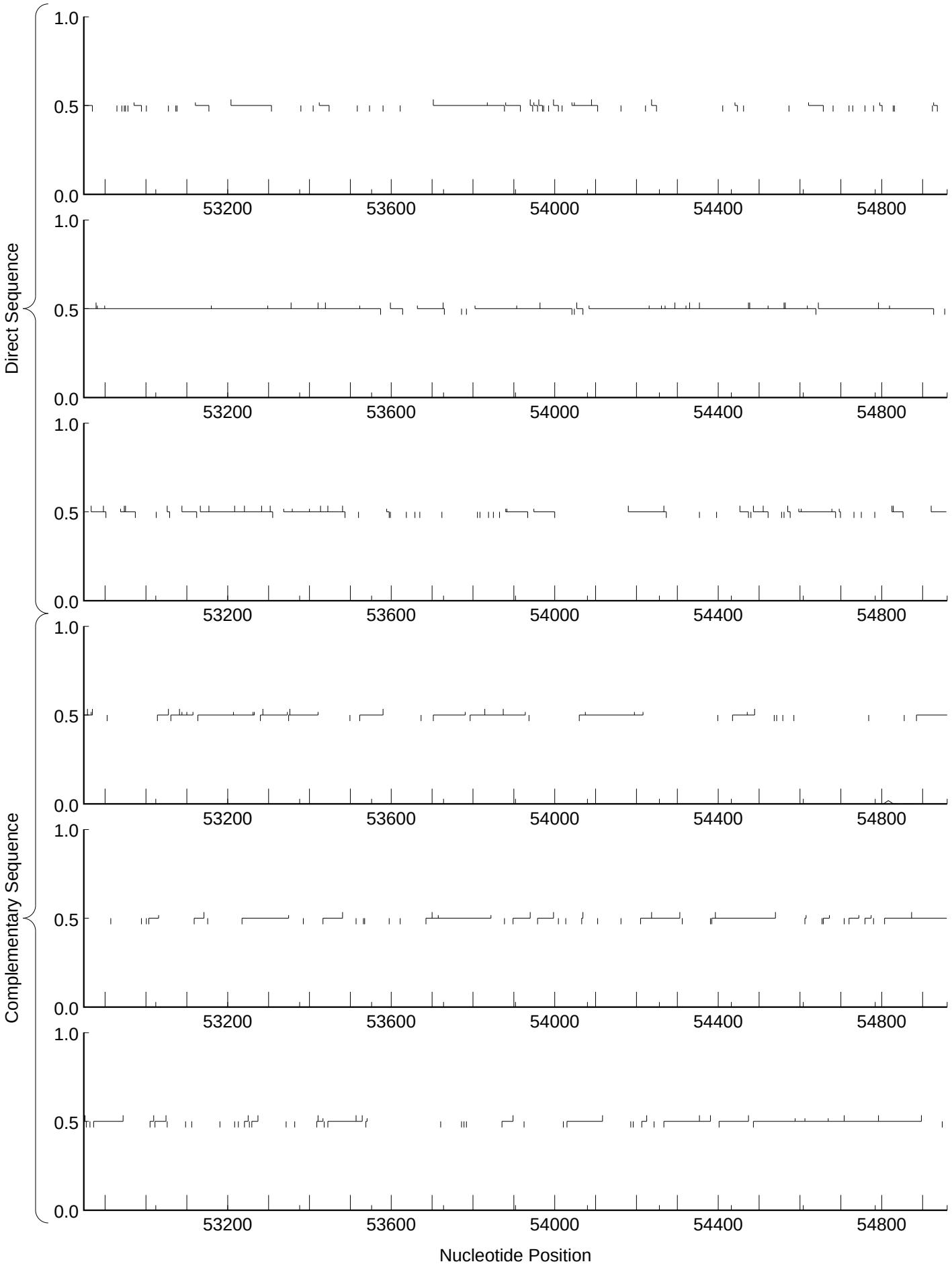
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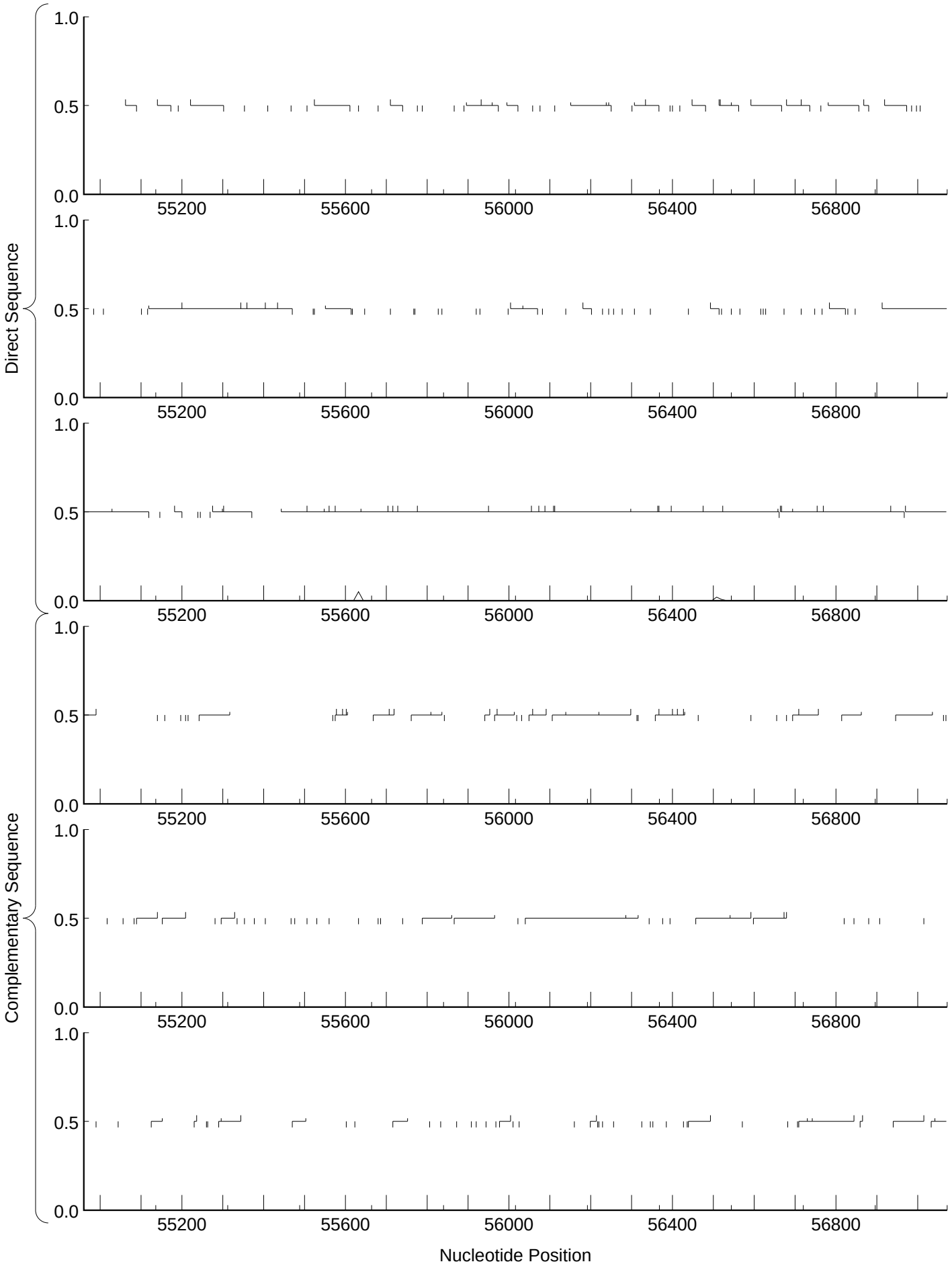
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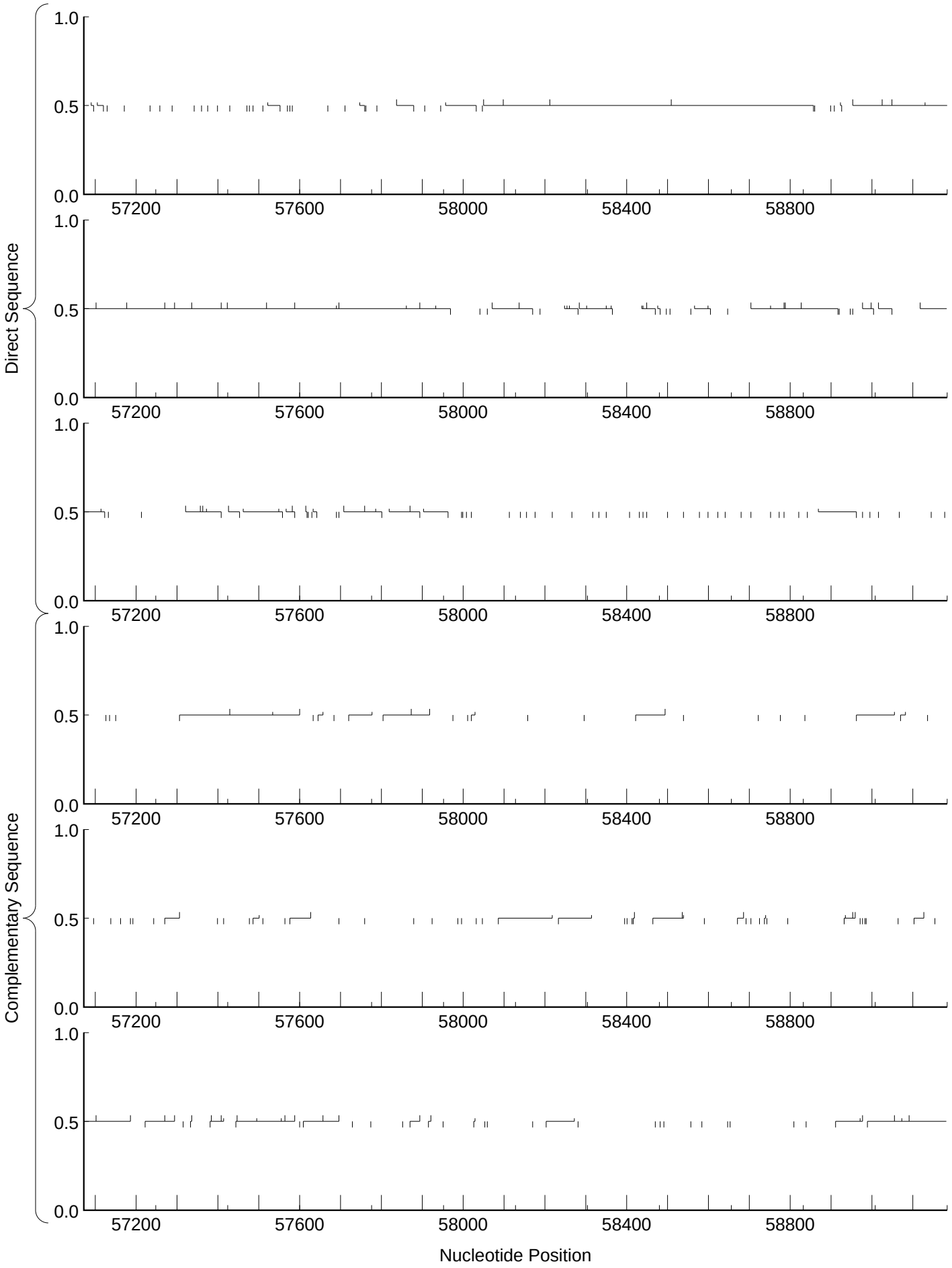
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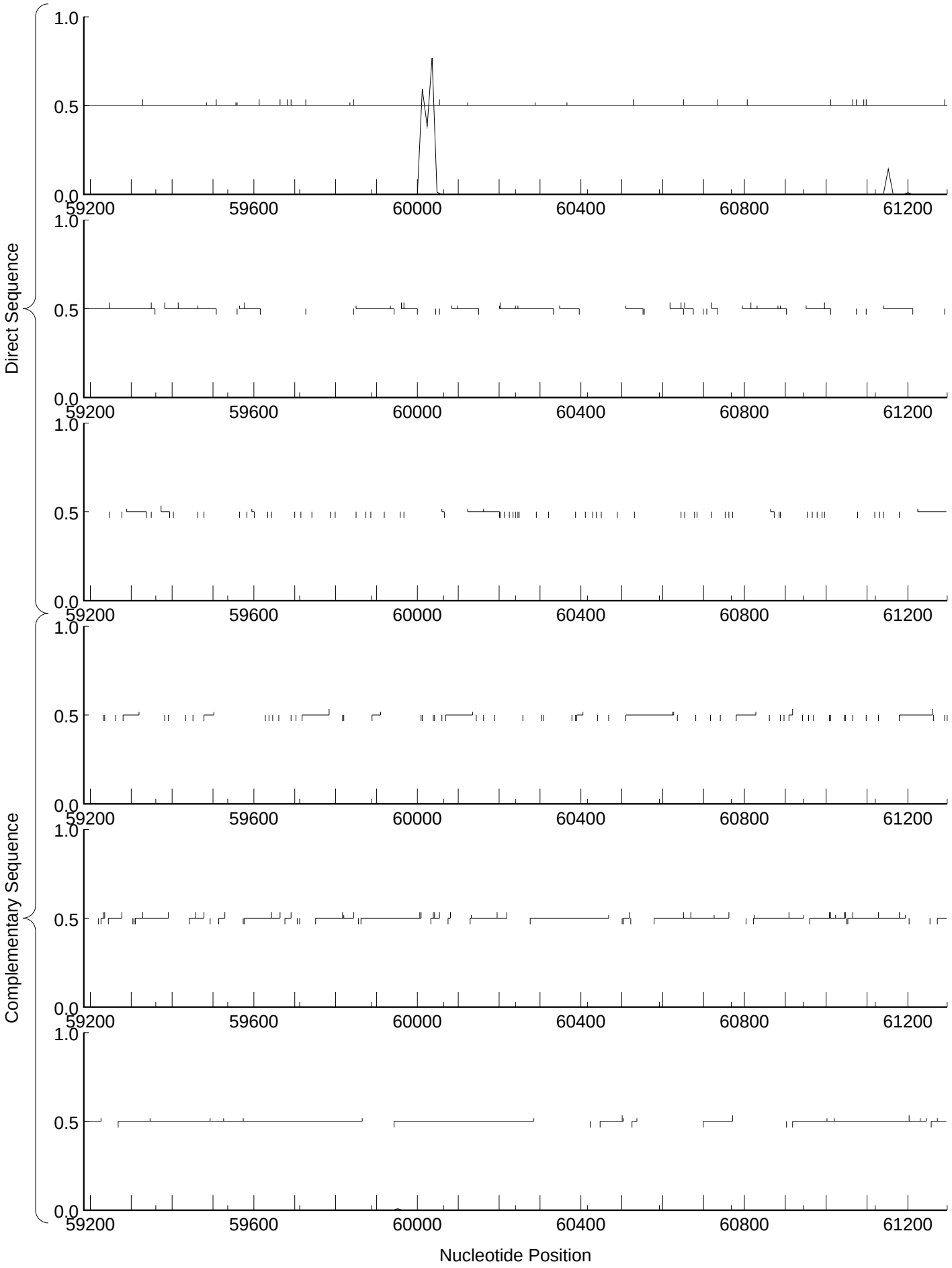
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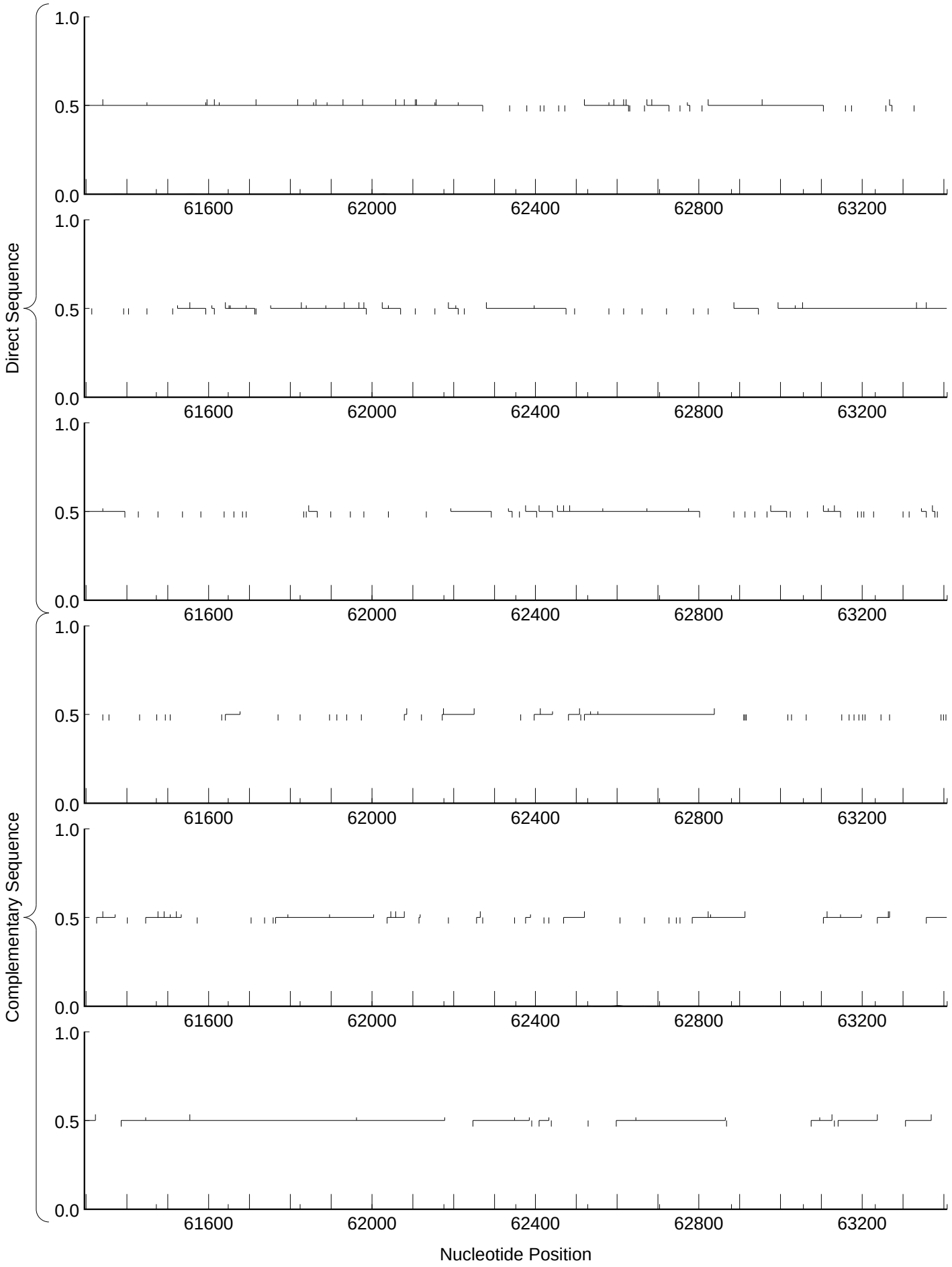
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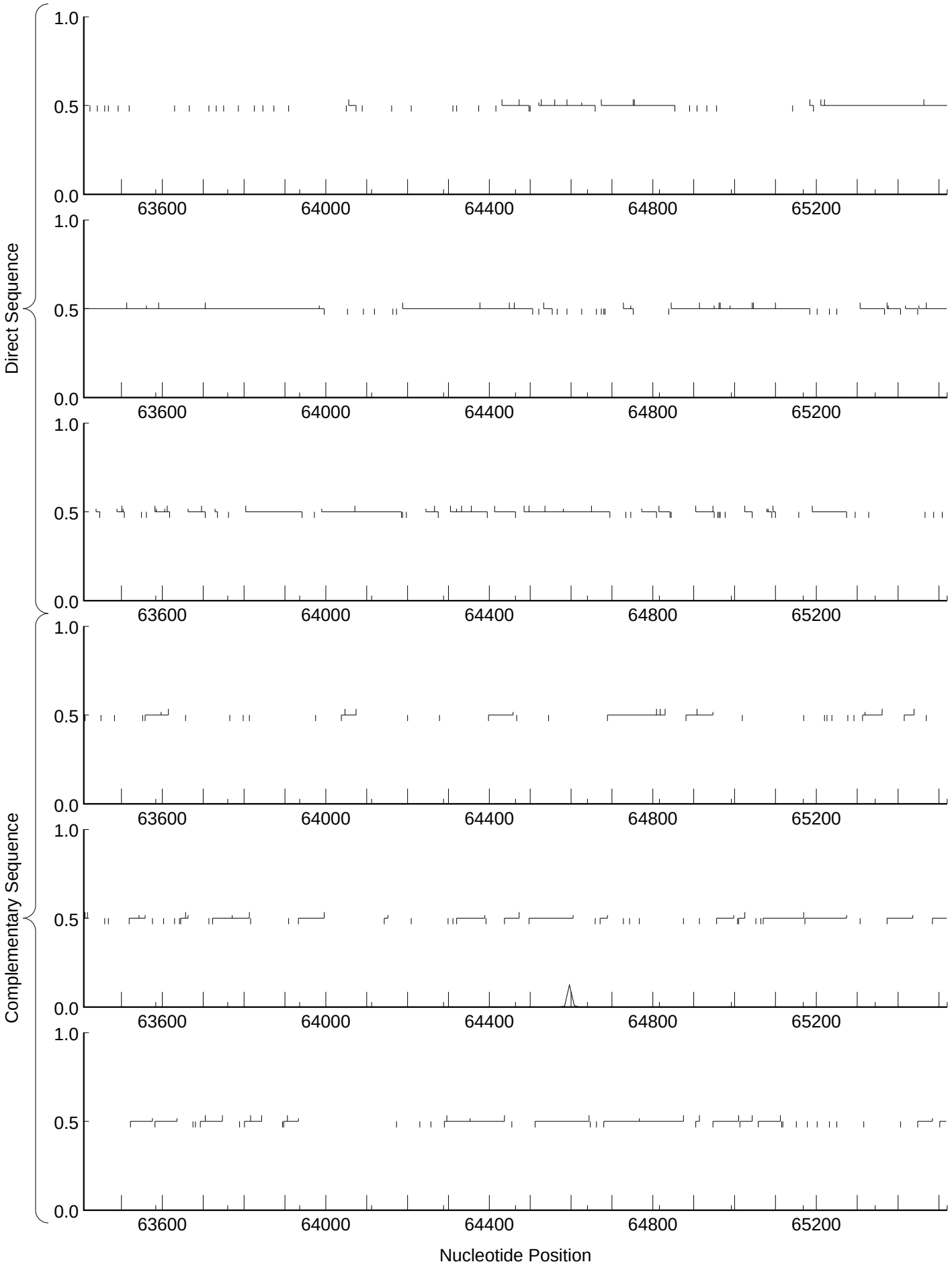
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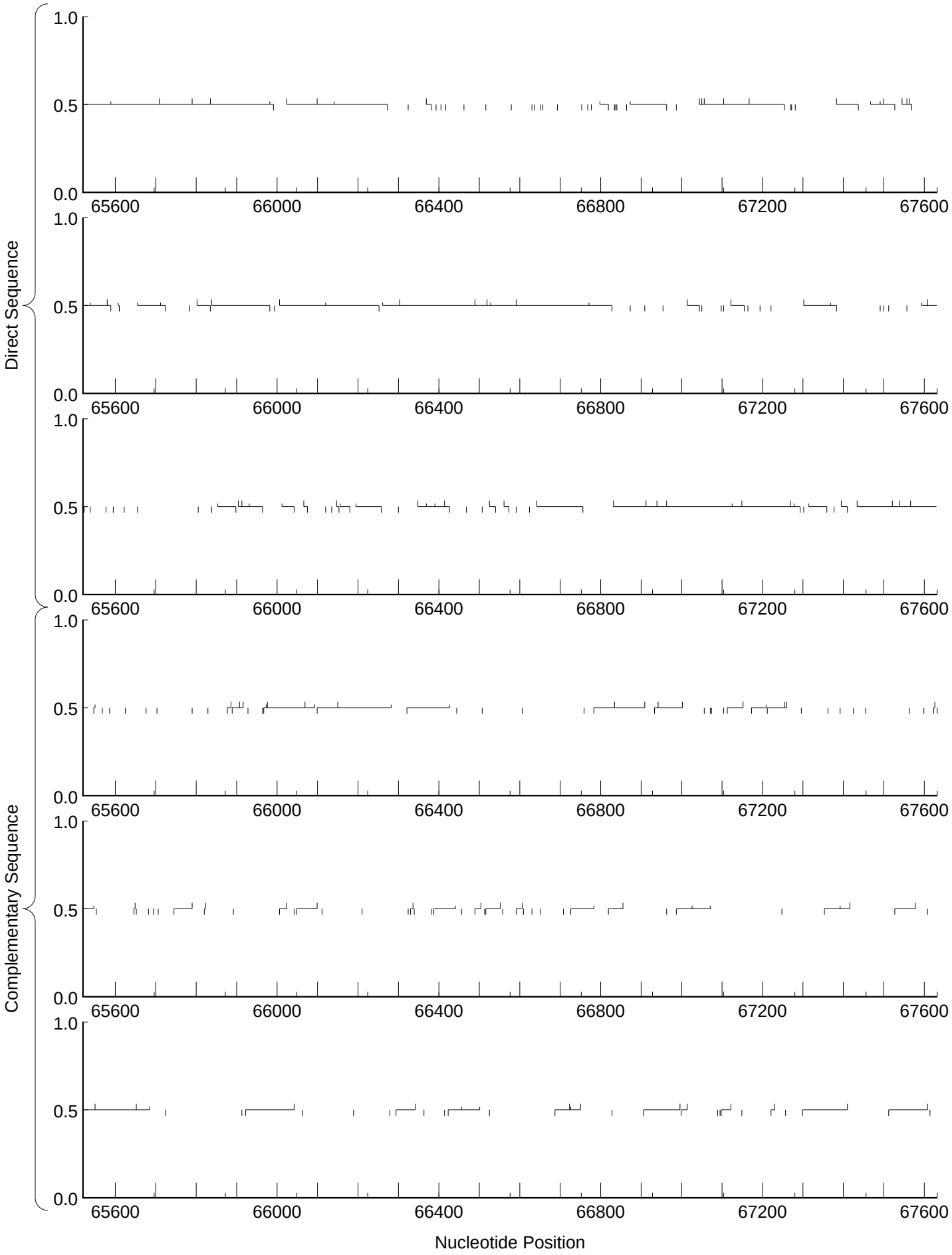
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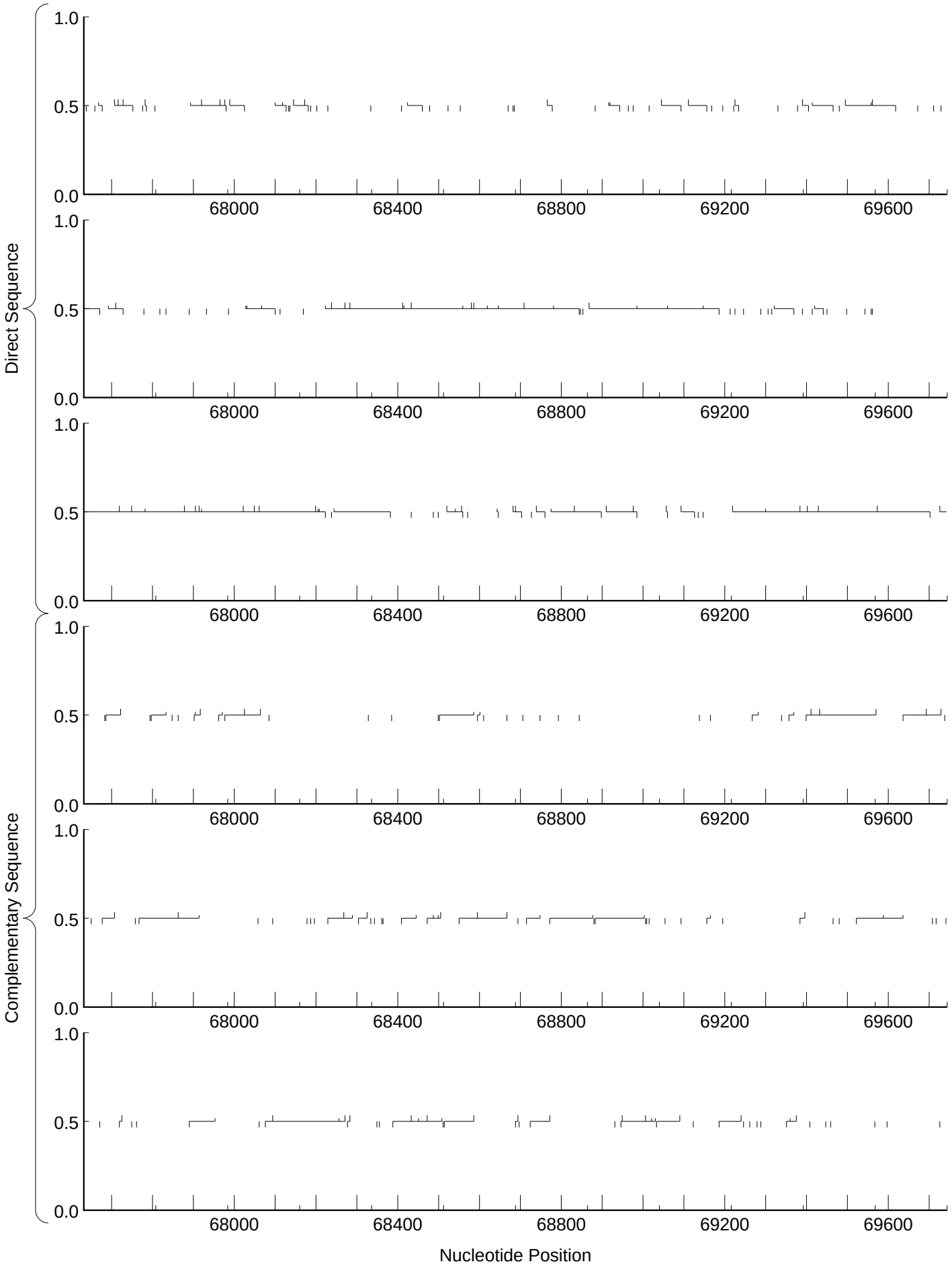
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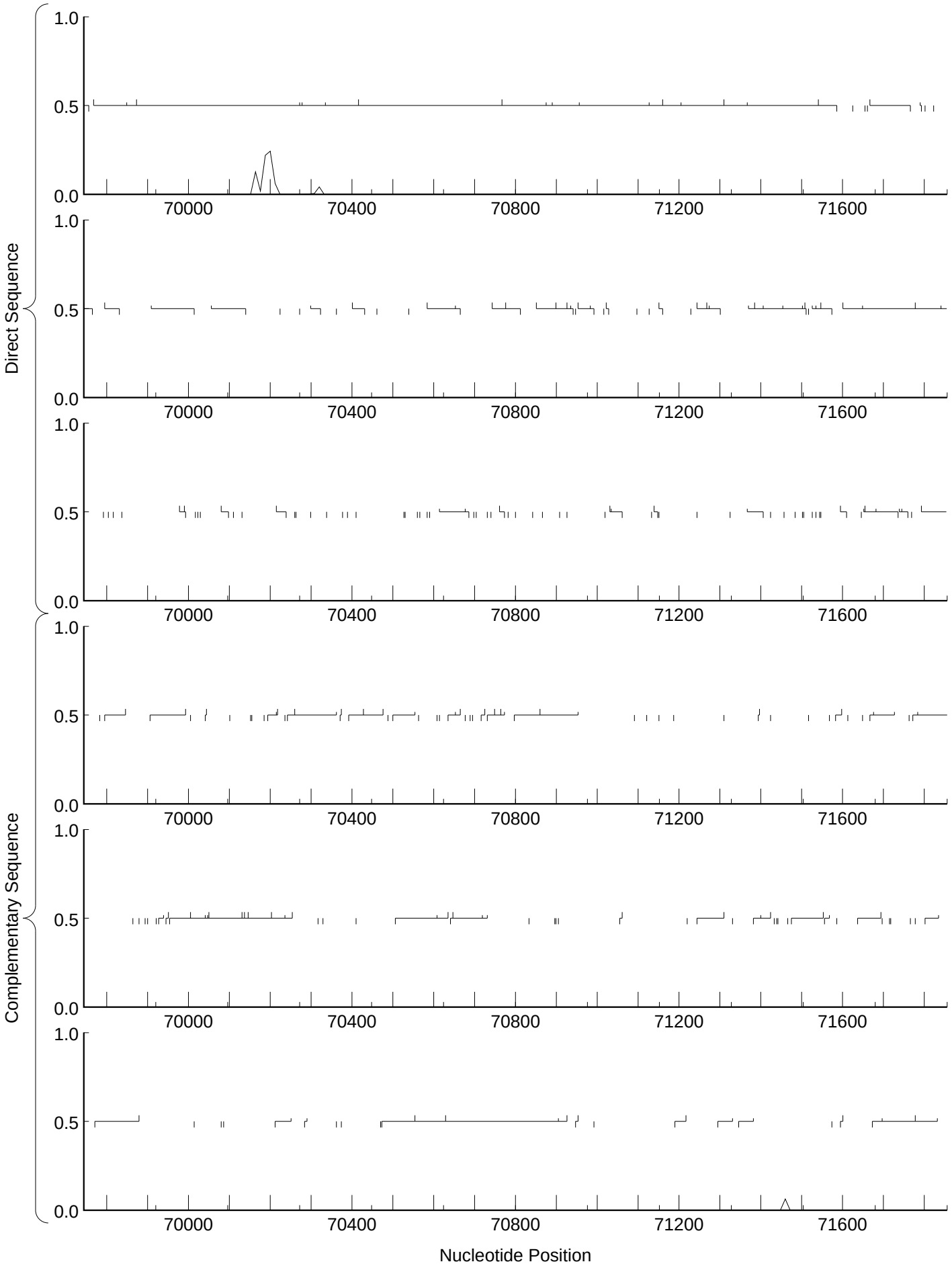
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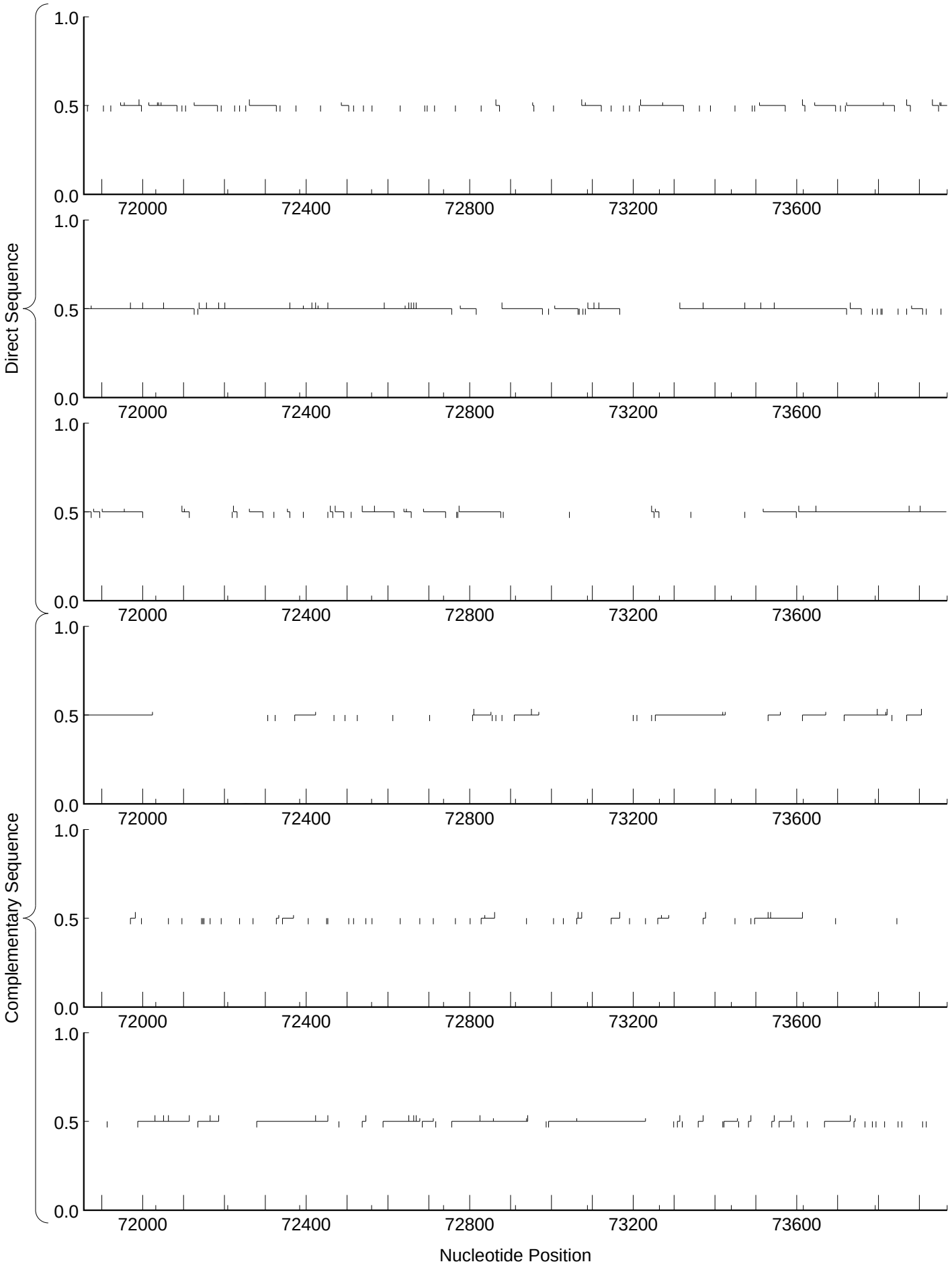
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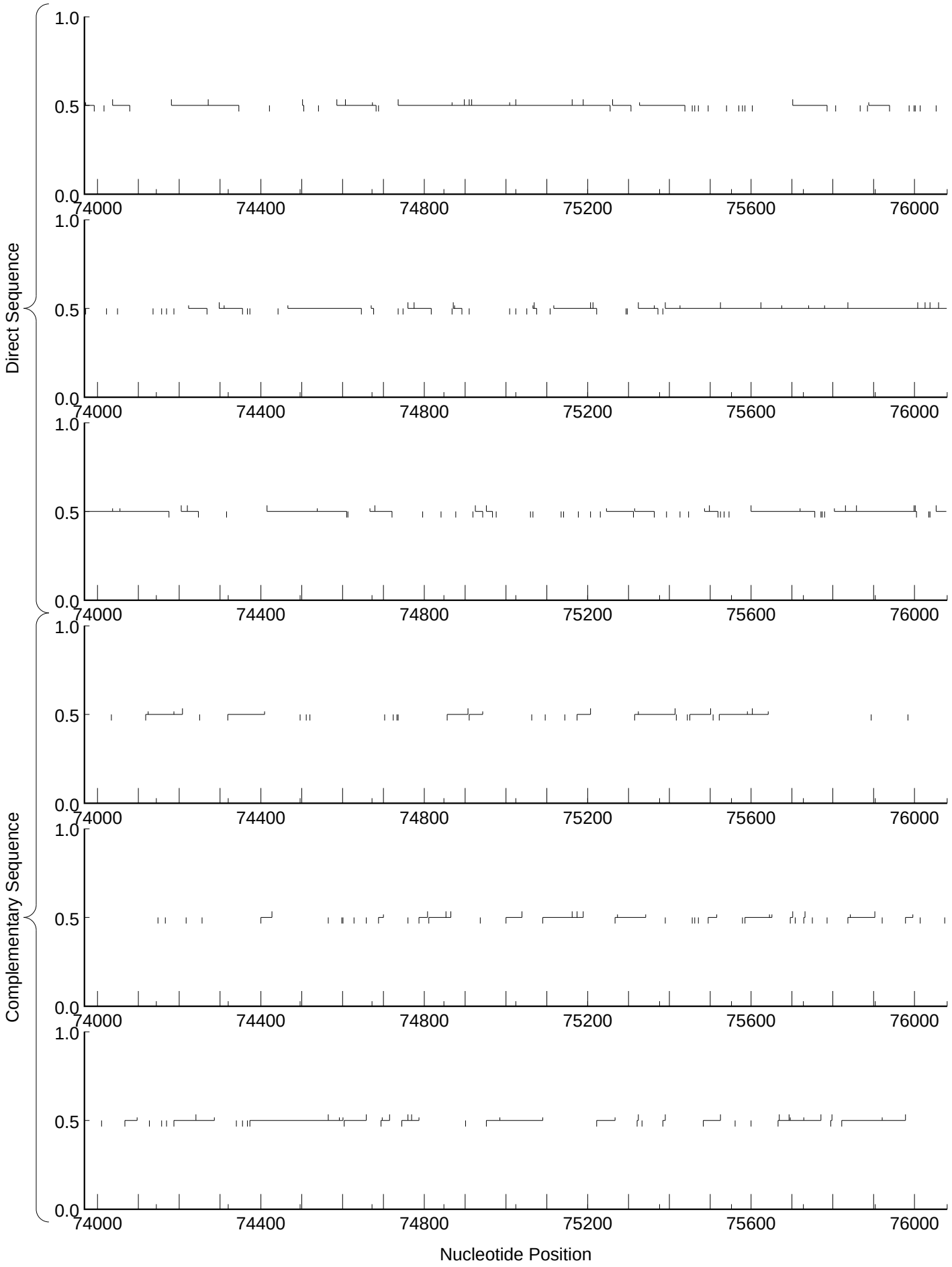
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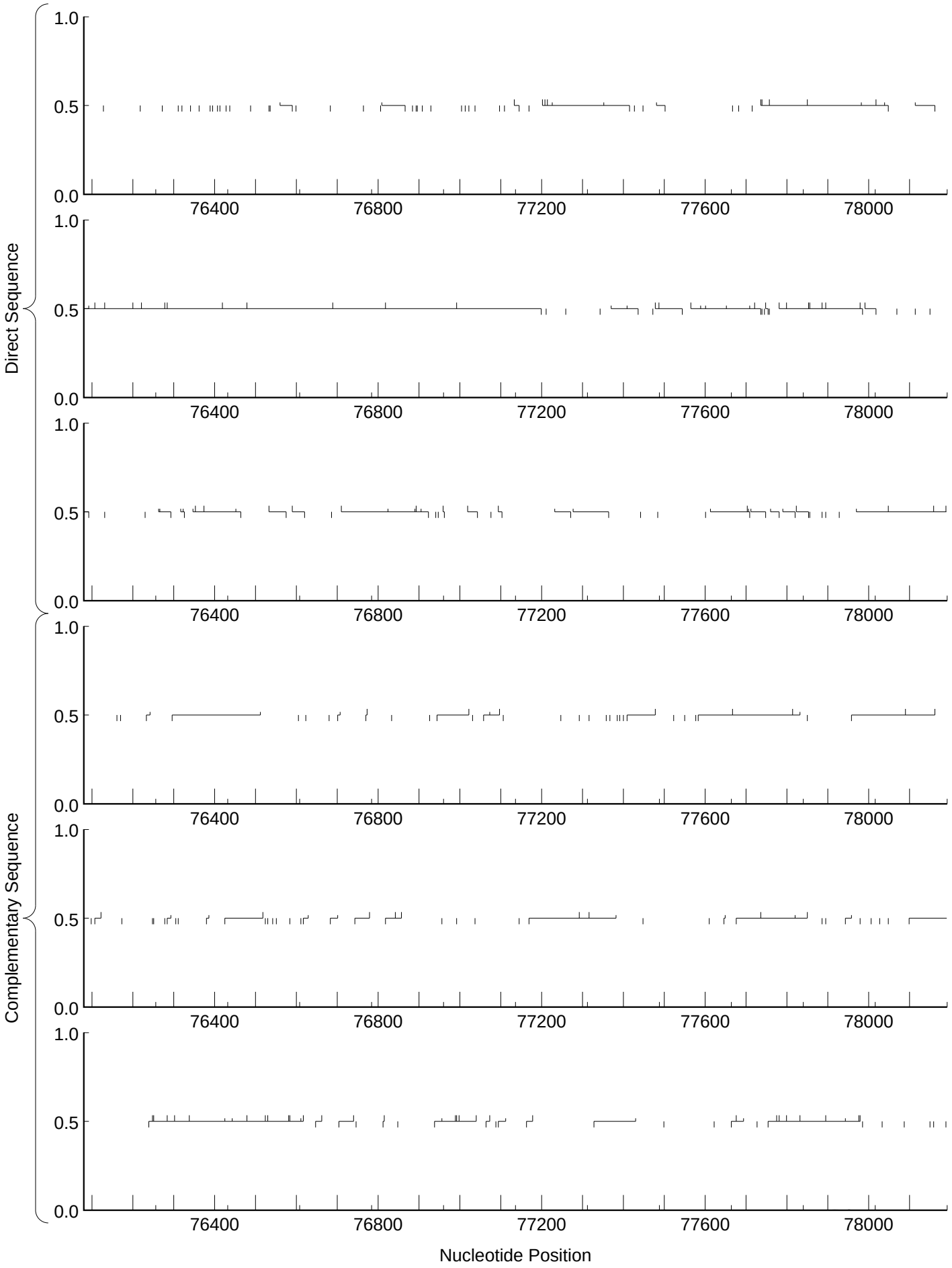
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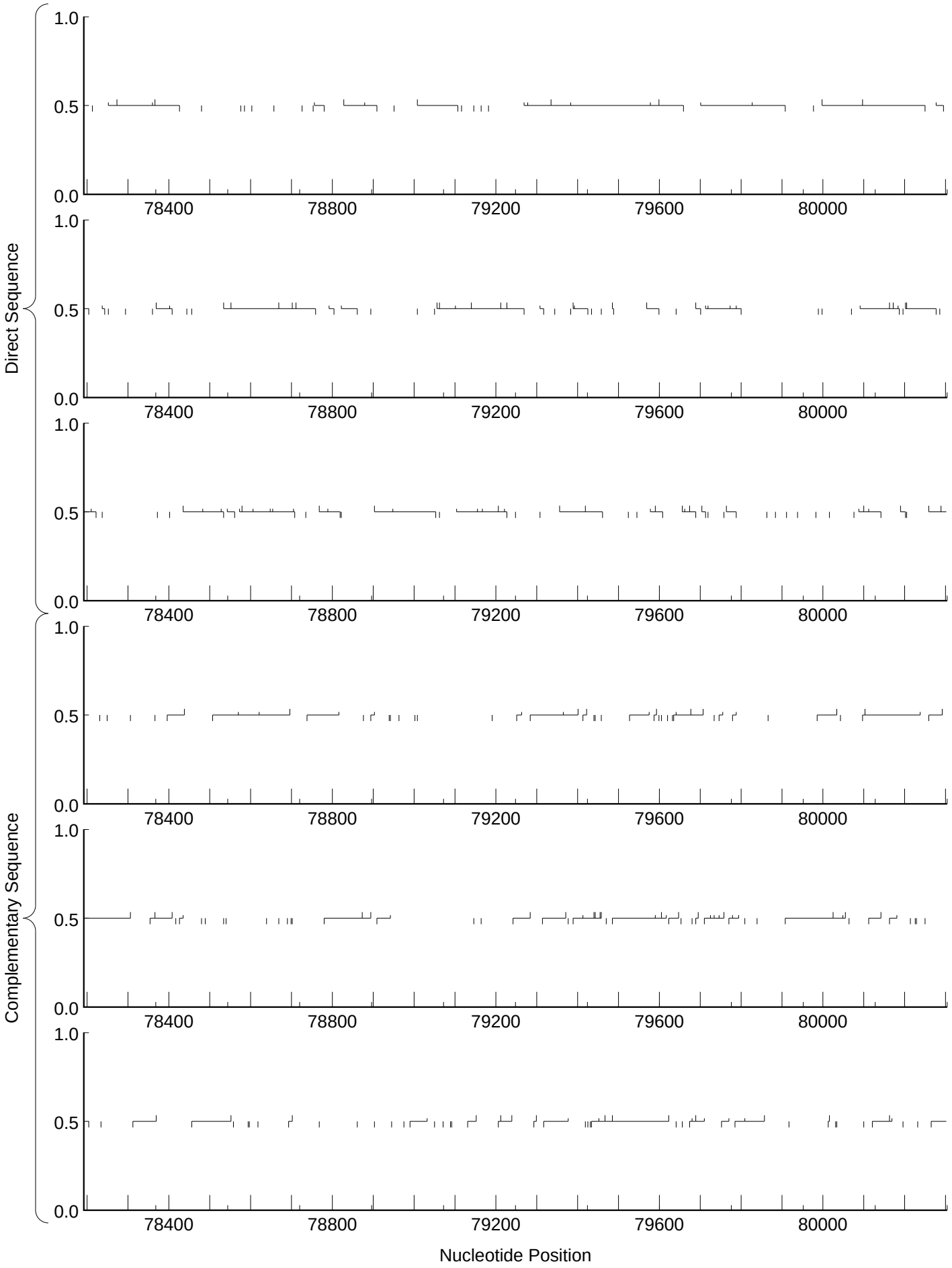
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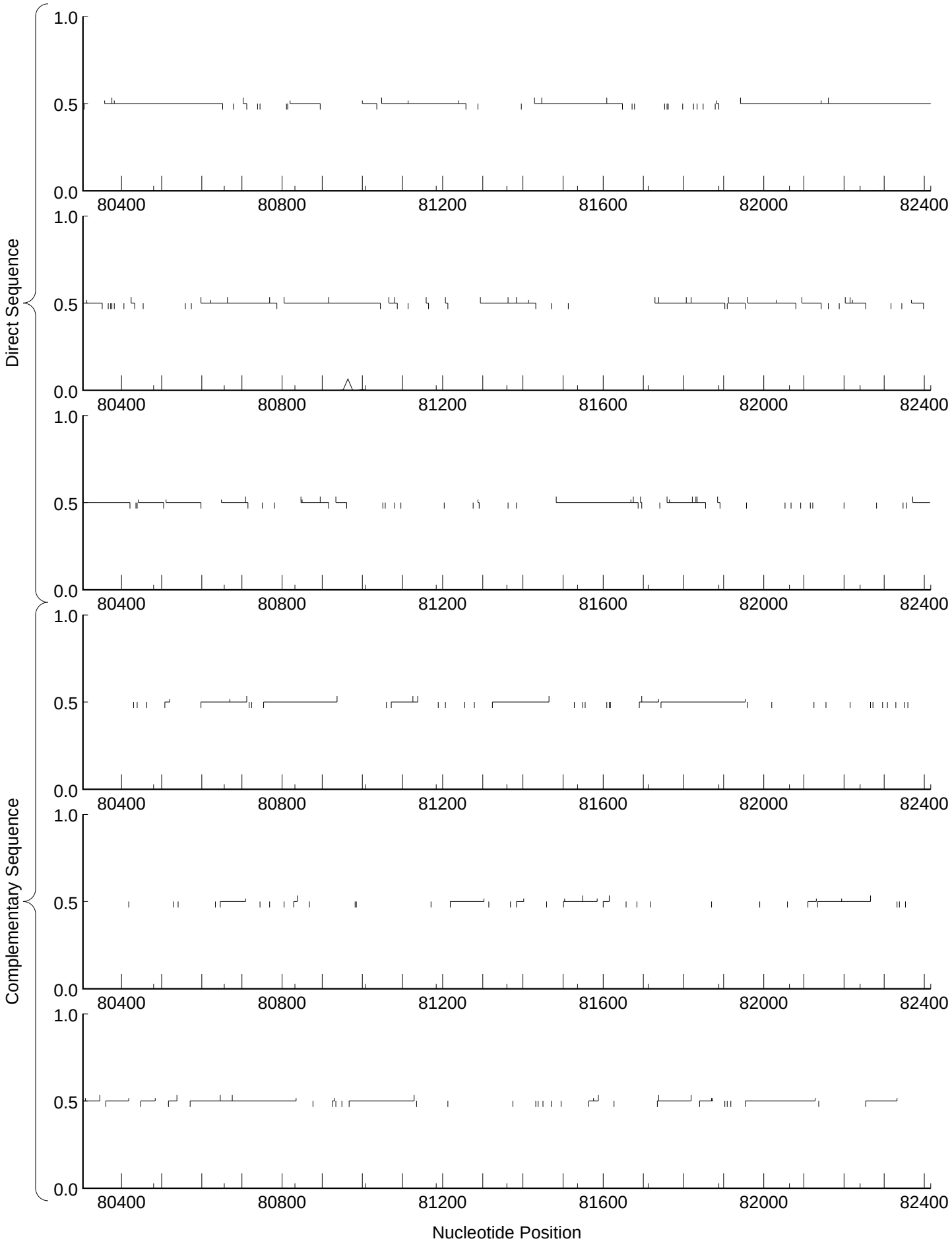
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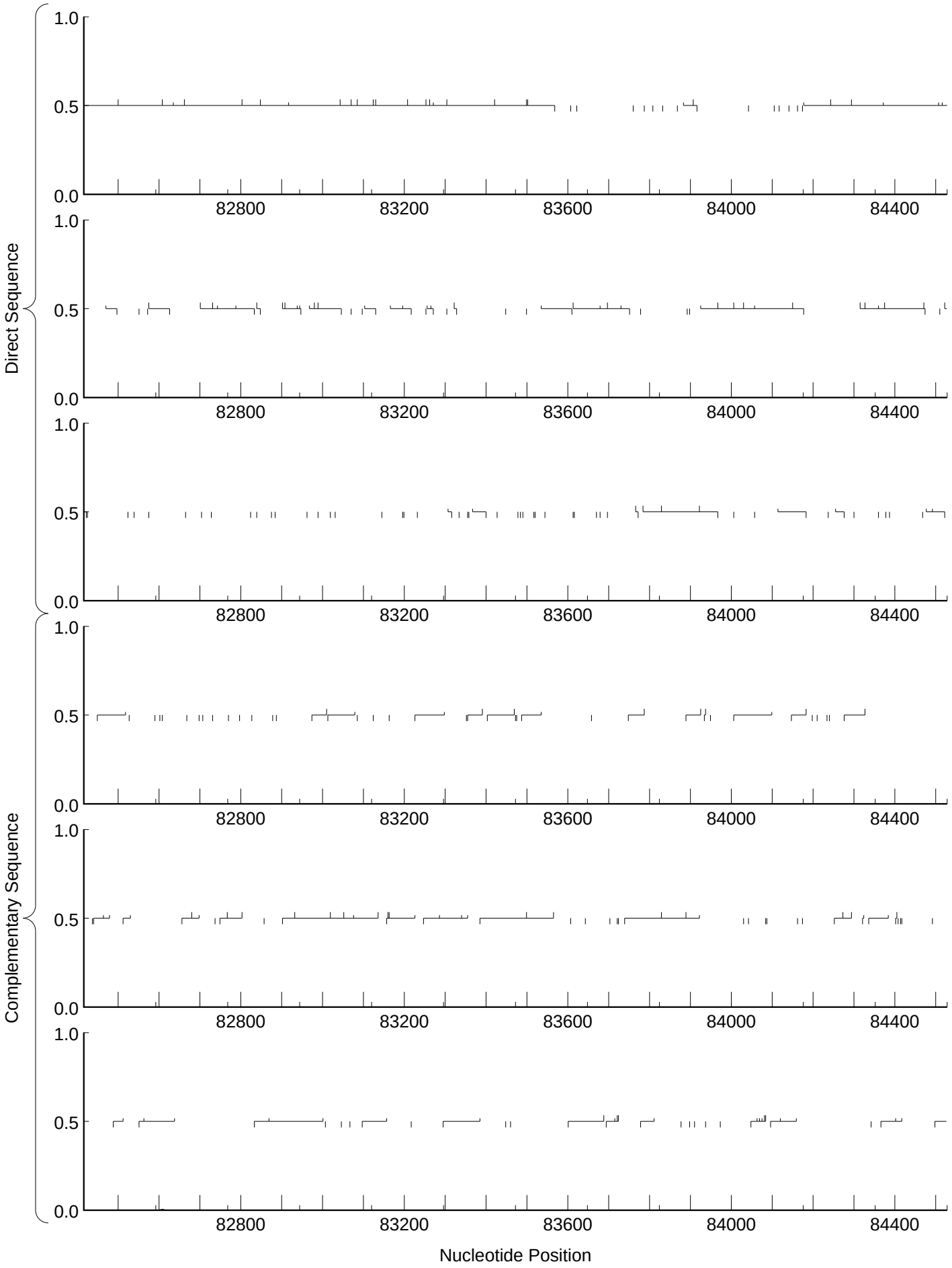
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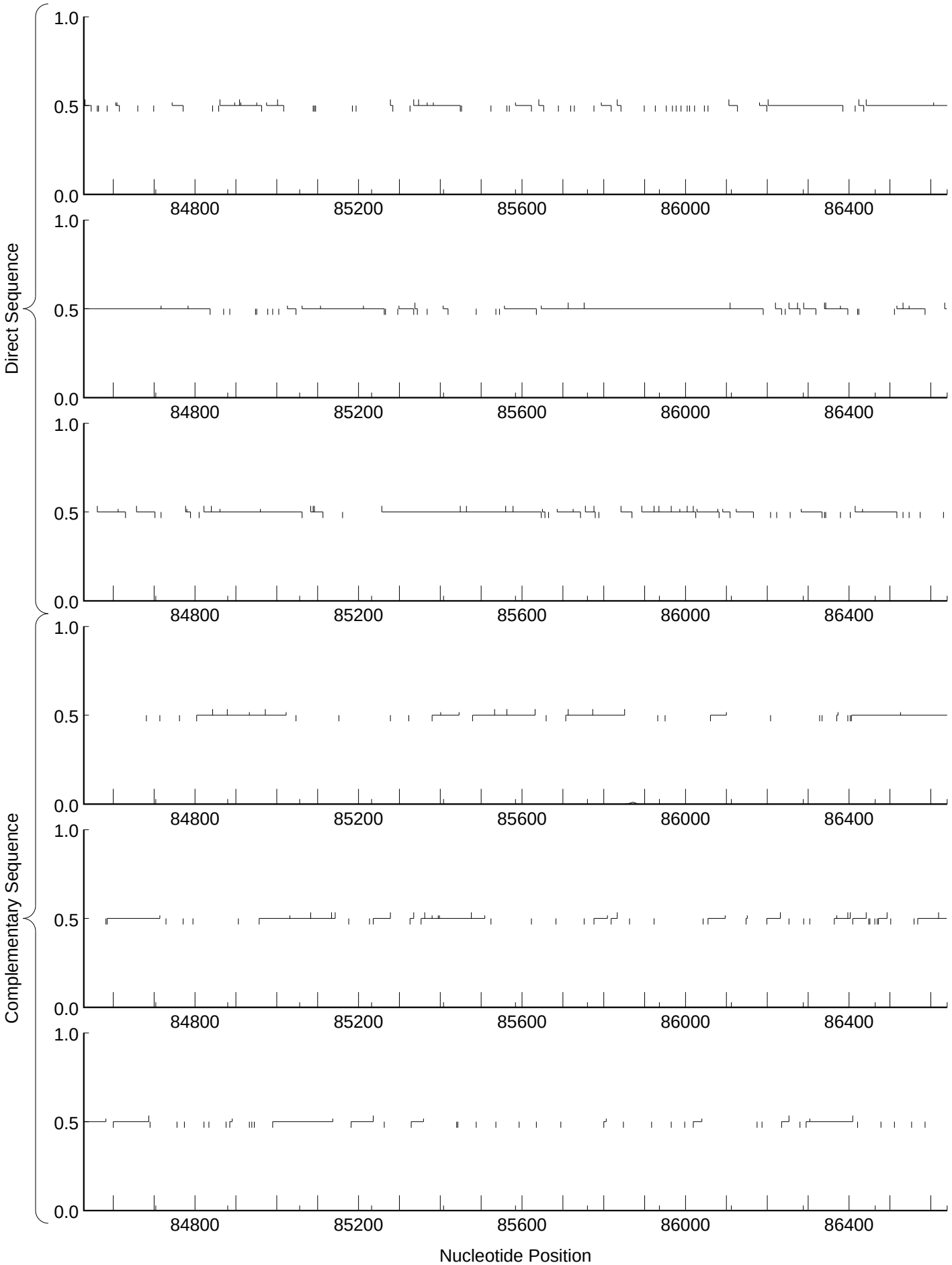


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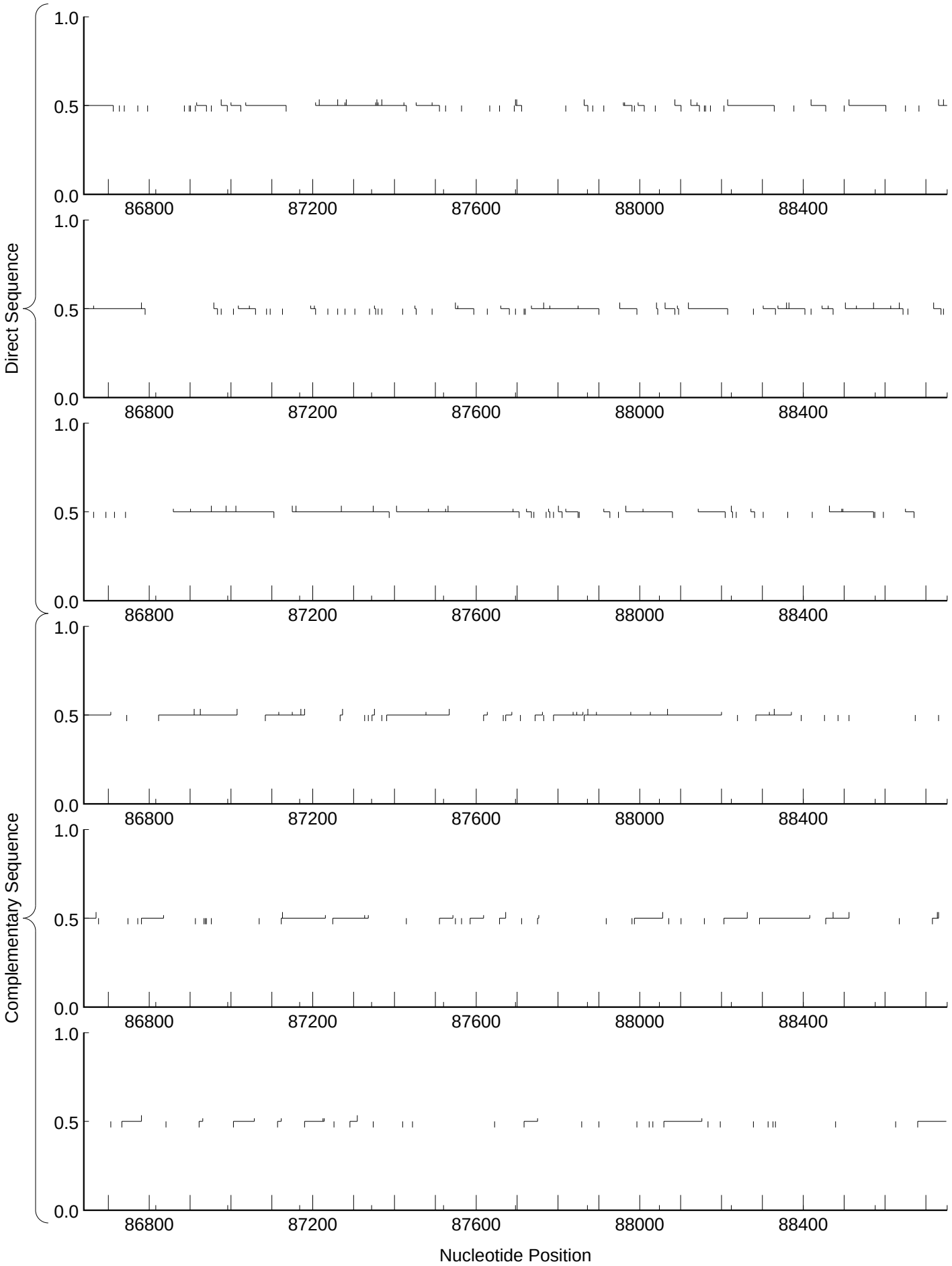


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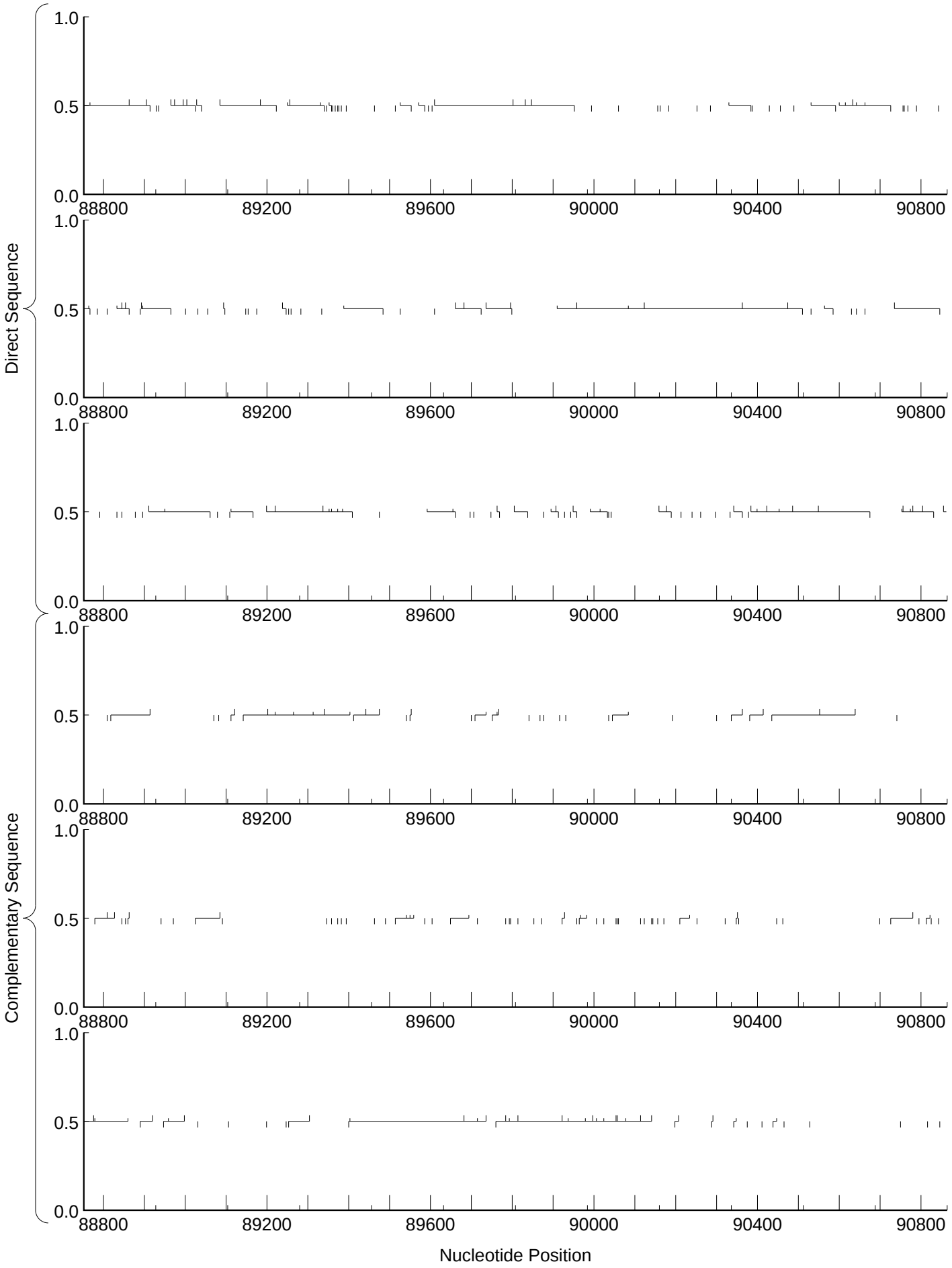



Direct Sequence
Complementary Sequence
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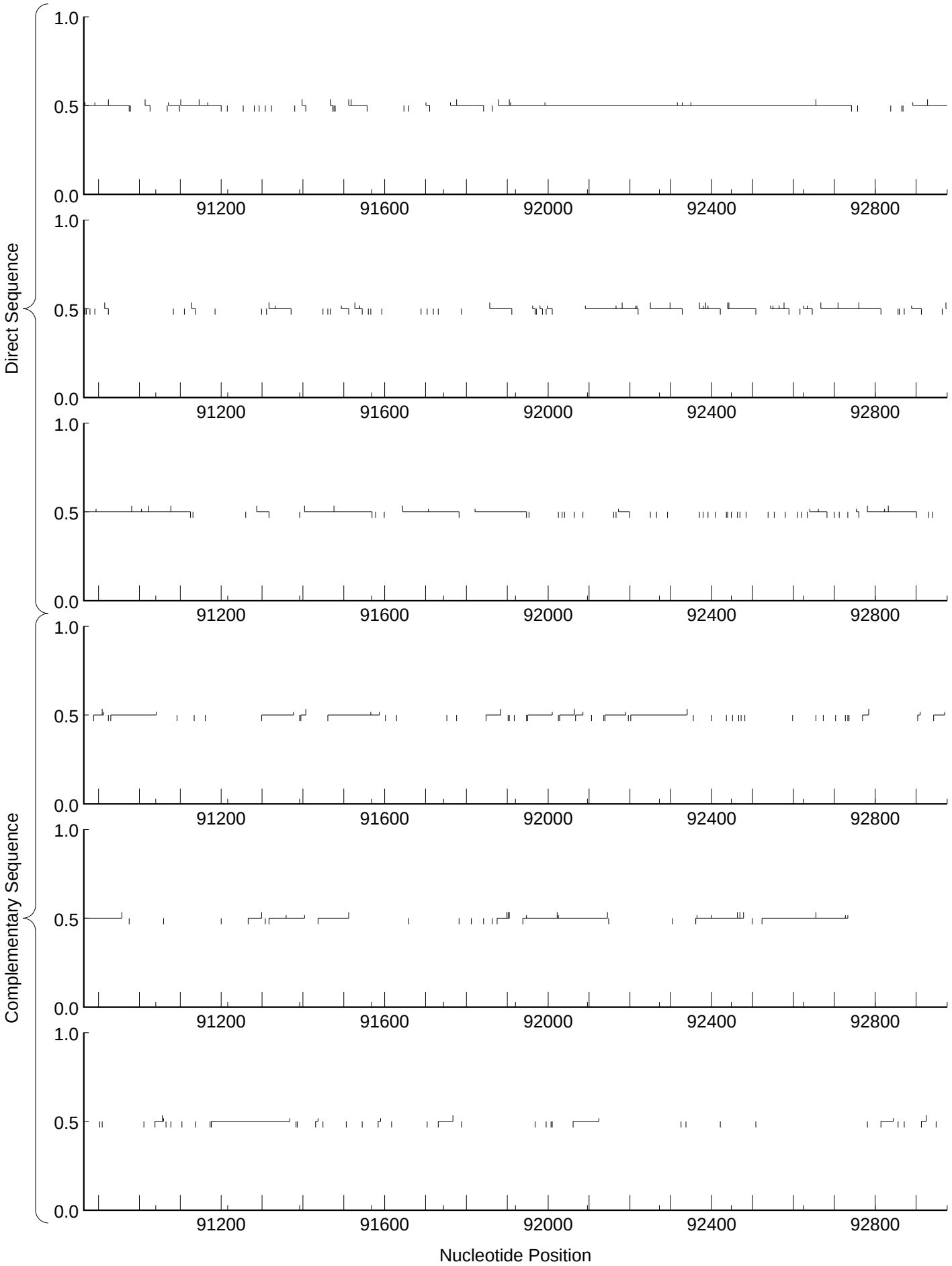
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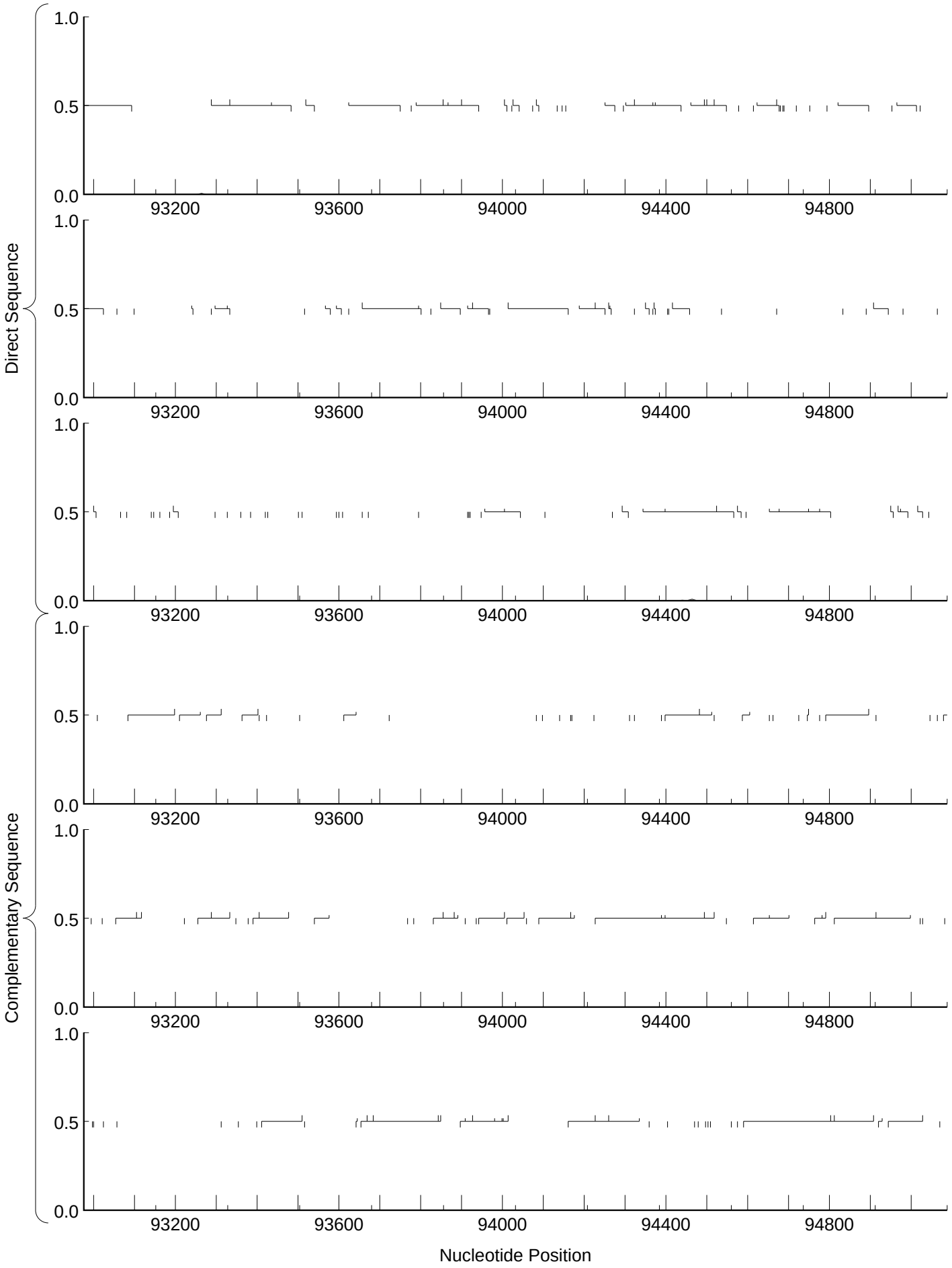
tarPlatinum complete sequence, 133886 bp including 12199 bp terminal repeat, Cluster BE2, Order 4, Window 96, Step 12, 44/65



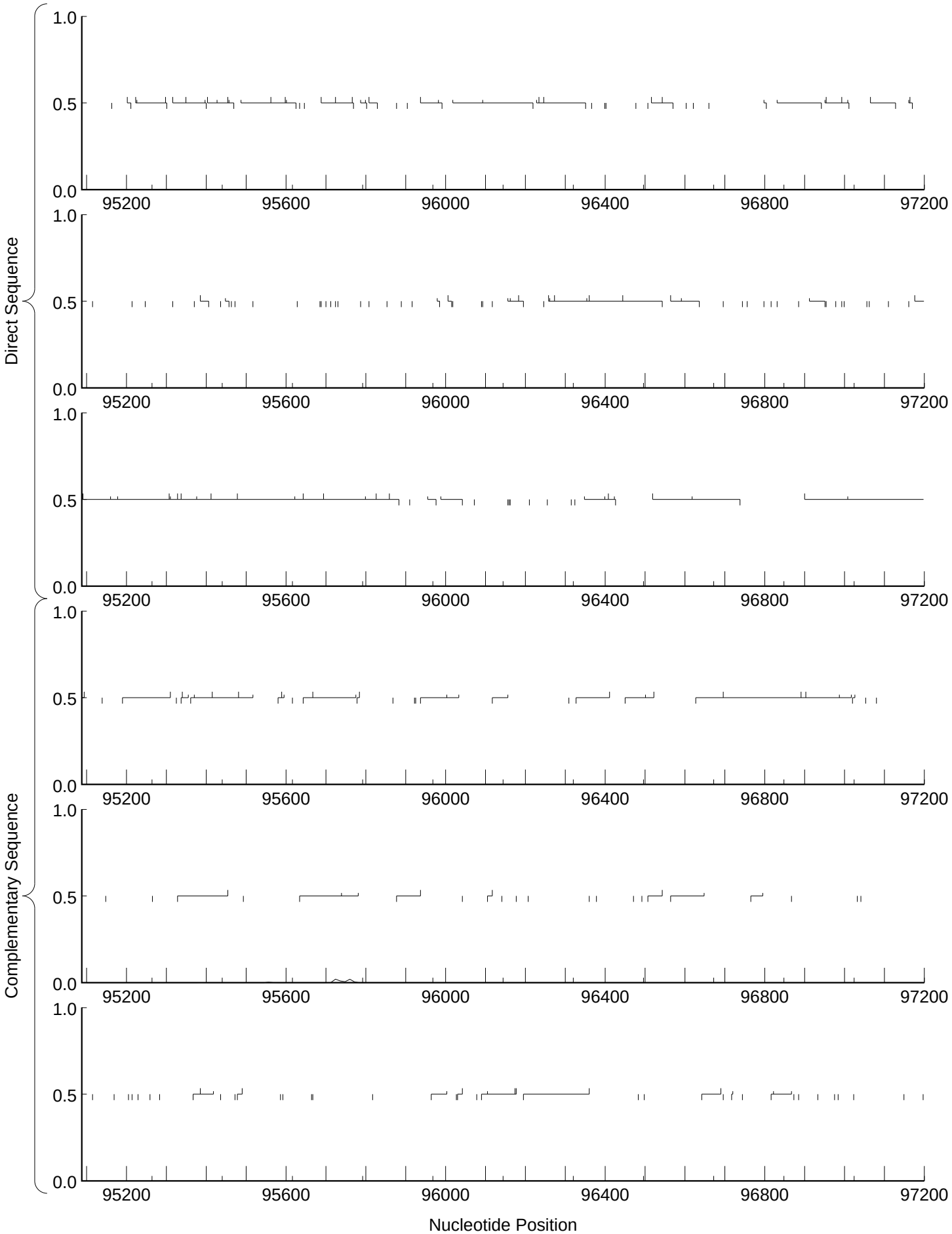
tarPlatinum complete sequence, 133886 bp including 12199 bp terminal repeat, Cluster BE2, Order 4, Window 96, Step 12, 45/65



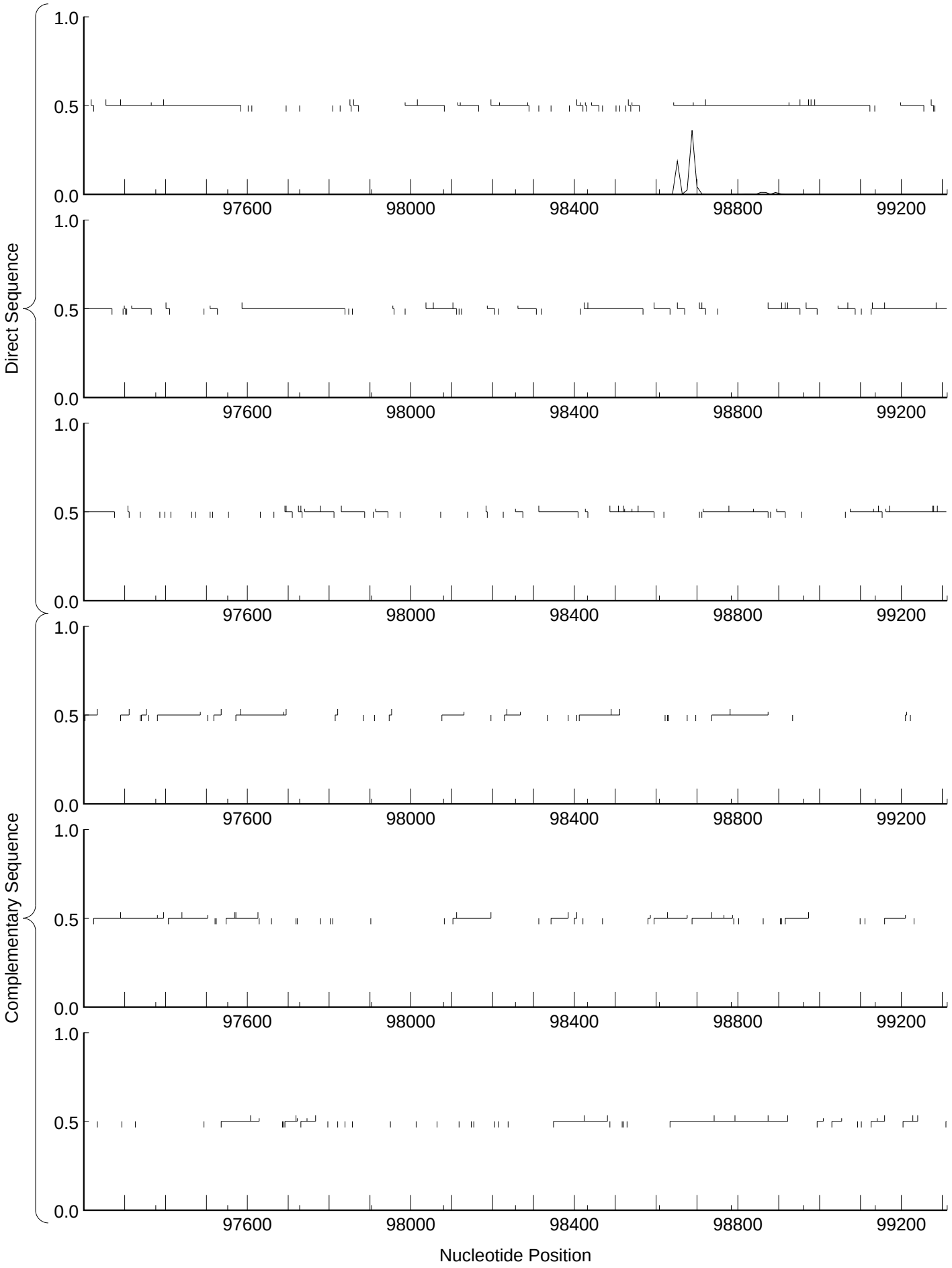
tarPlatinum complete sequence, 133886 bp including 12199 bp terminal repeat, Cluster BE2, Order 4, Window 96, Step 12, 46/65



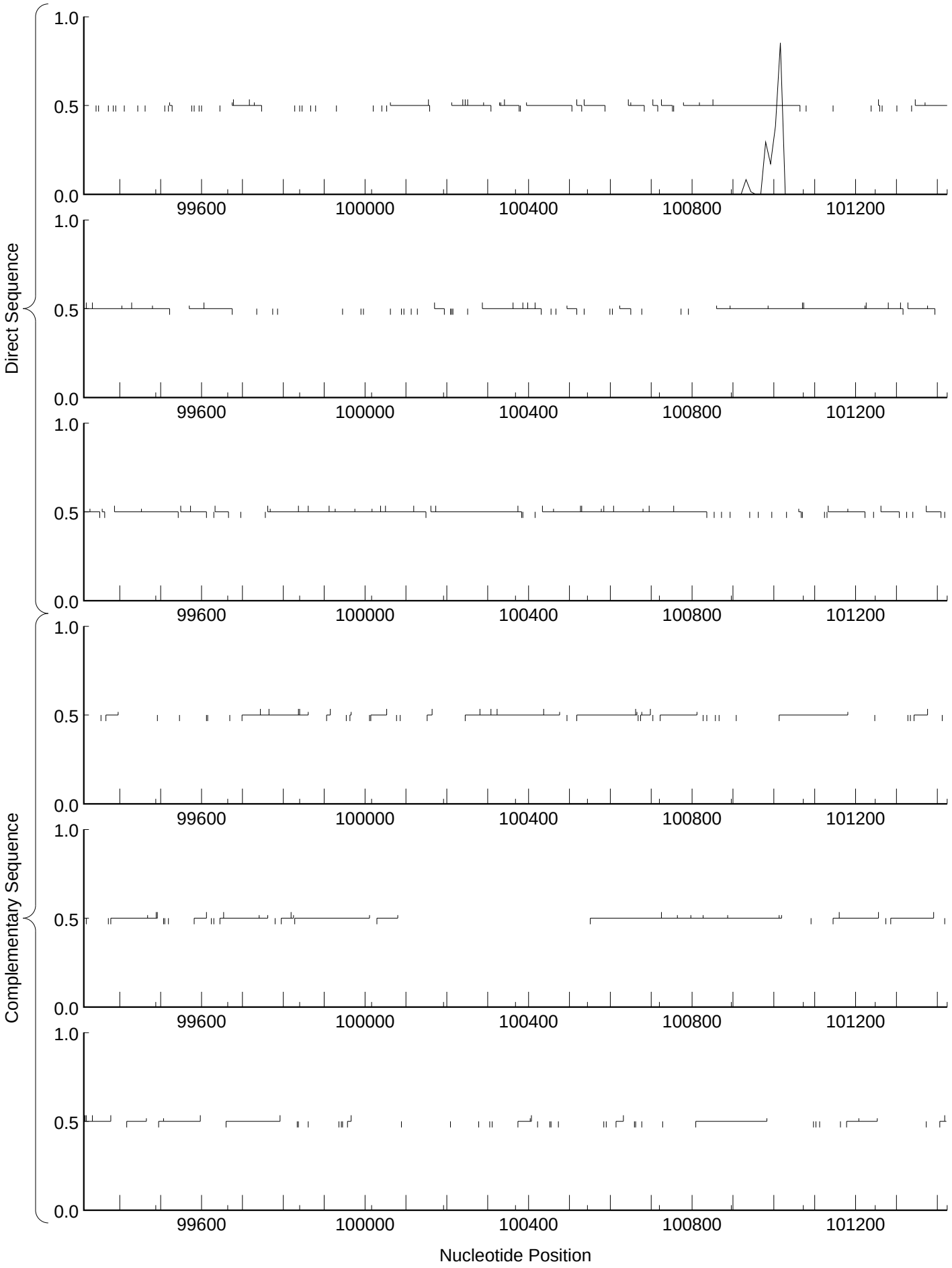
tarPlatinum complete sequence, 133886 bp including 12199 bp terminal repeat, Cluster BE2, Order 4, Window 96, Step 12, 47/65



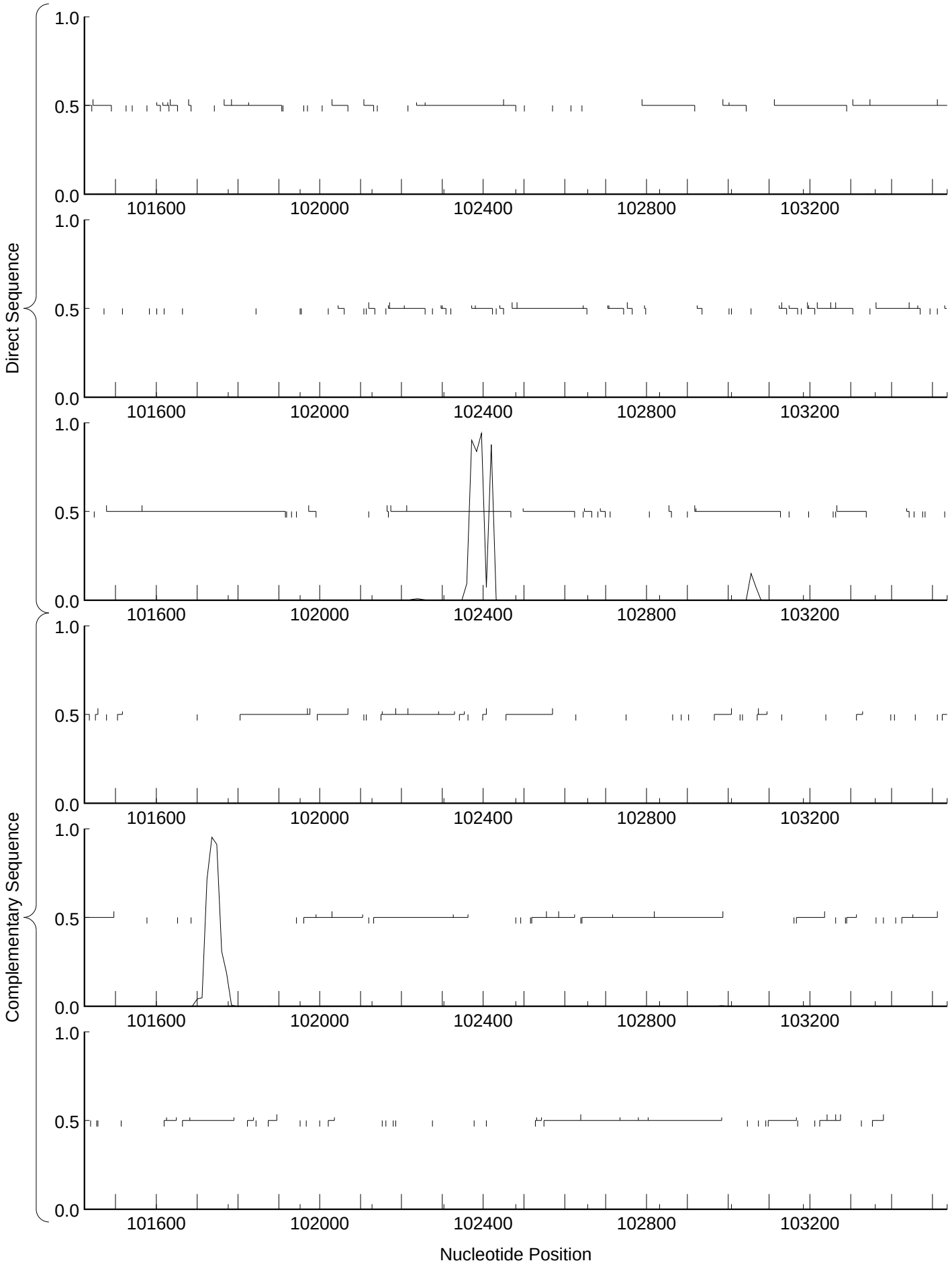
tarPlatinum complete sequence, 133886 bp including 12199 bp terminal repeat, Cluster BE2, Order 4, Window 96, Step 12, 48/65



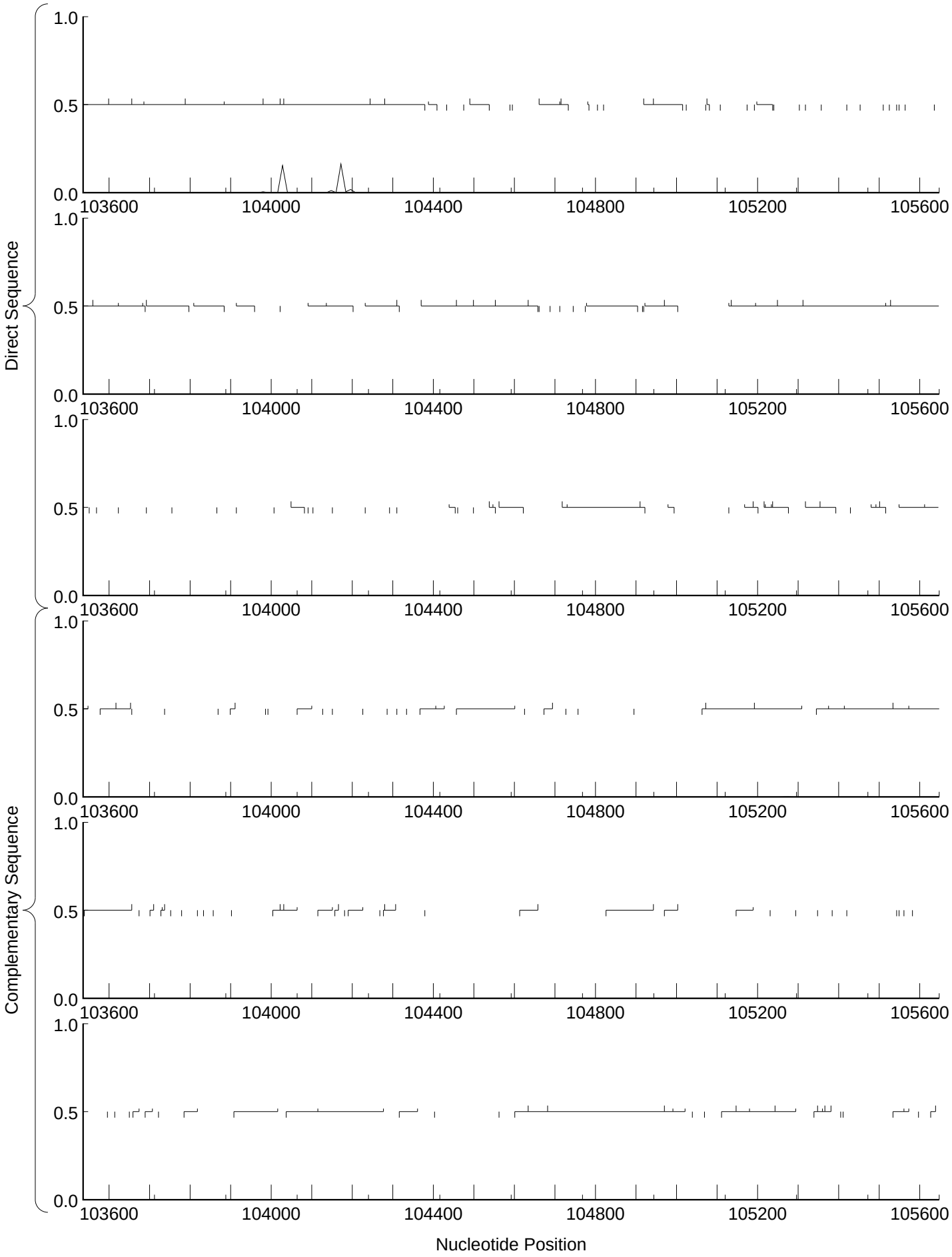
tarPlatinum complete sequence, 133886 bp including 12199 bp terminal repeat, Cluster BE2, Order 4, Window 96, Step 12, 49/65



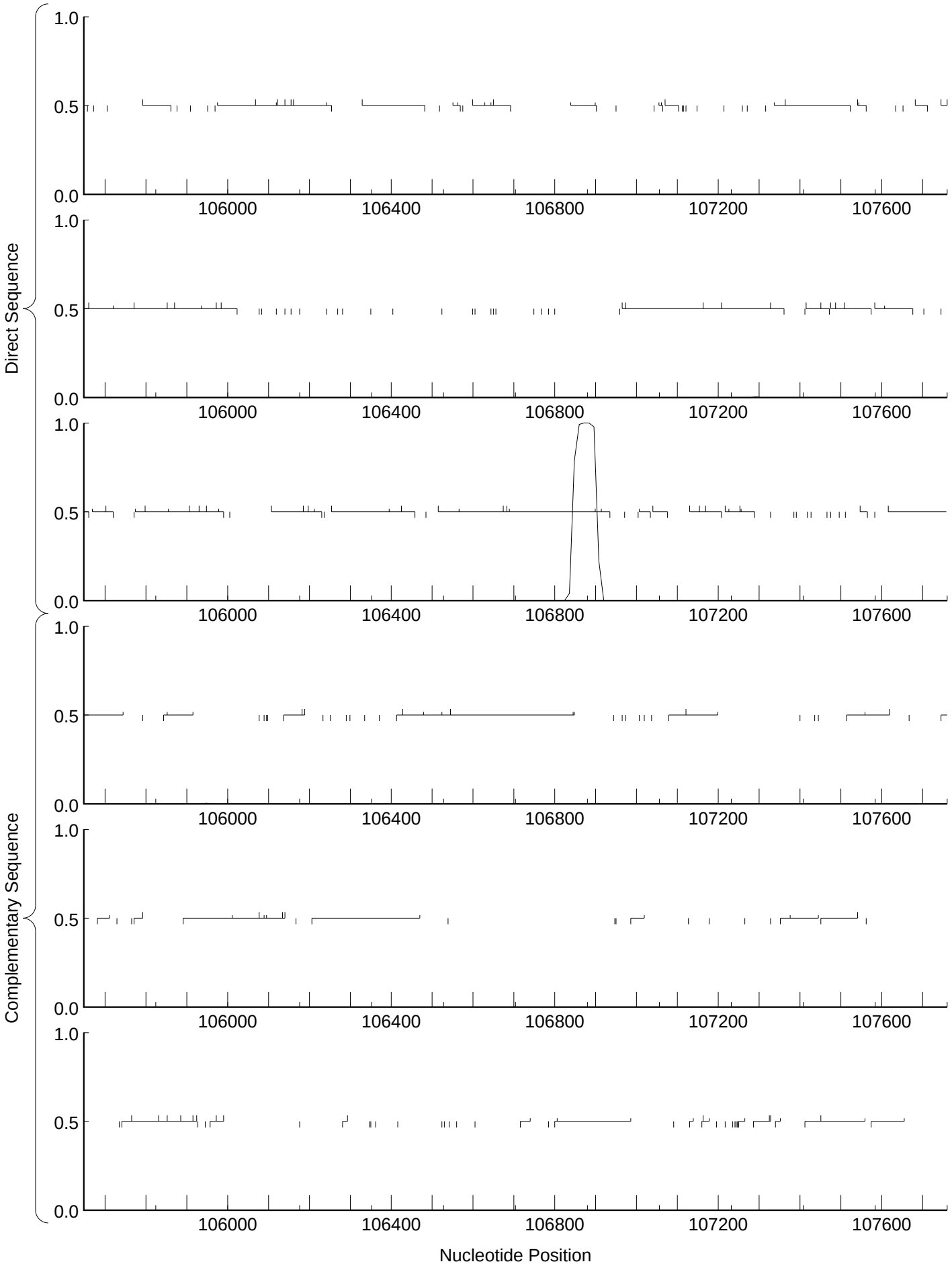
tarPlatinum complete sequence, 133886 bp including 12199 bp terminal repeat, Cluster BE2, Order 4, Window 96, Step 12, 50/65



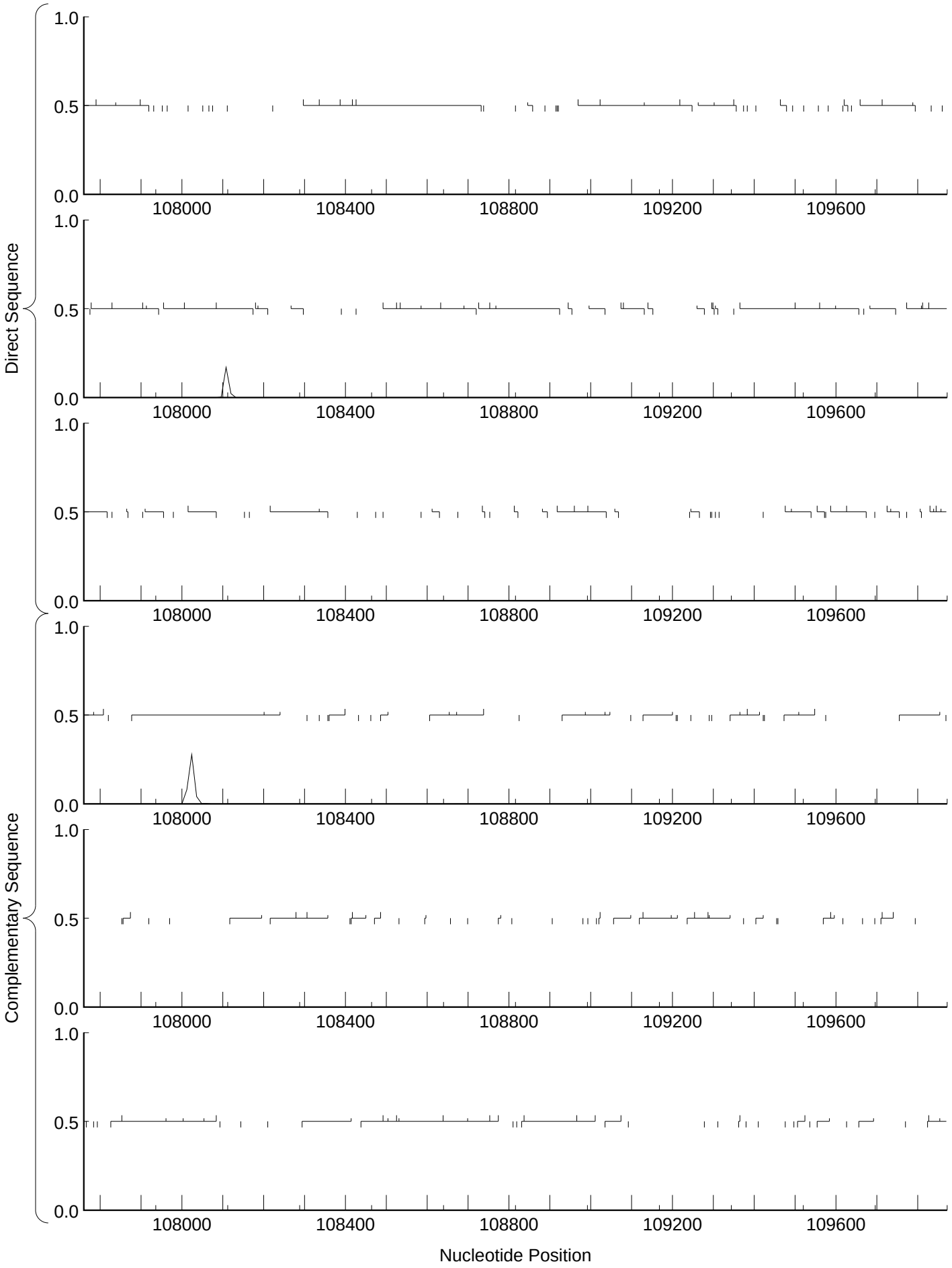
tarPlatinum complete sequence, 133886 bp including 12199 bp terminal repeat, Cluster BE2, Order 4, Window 96, Step 12, 51/65



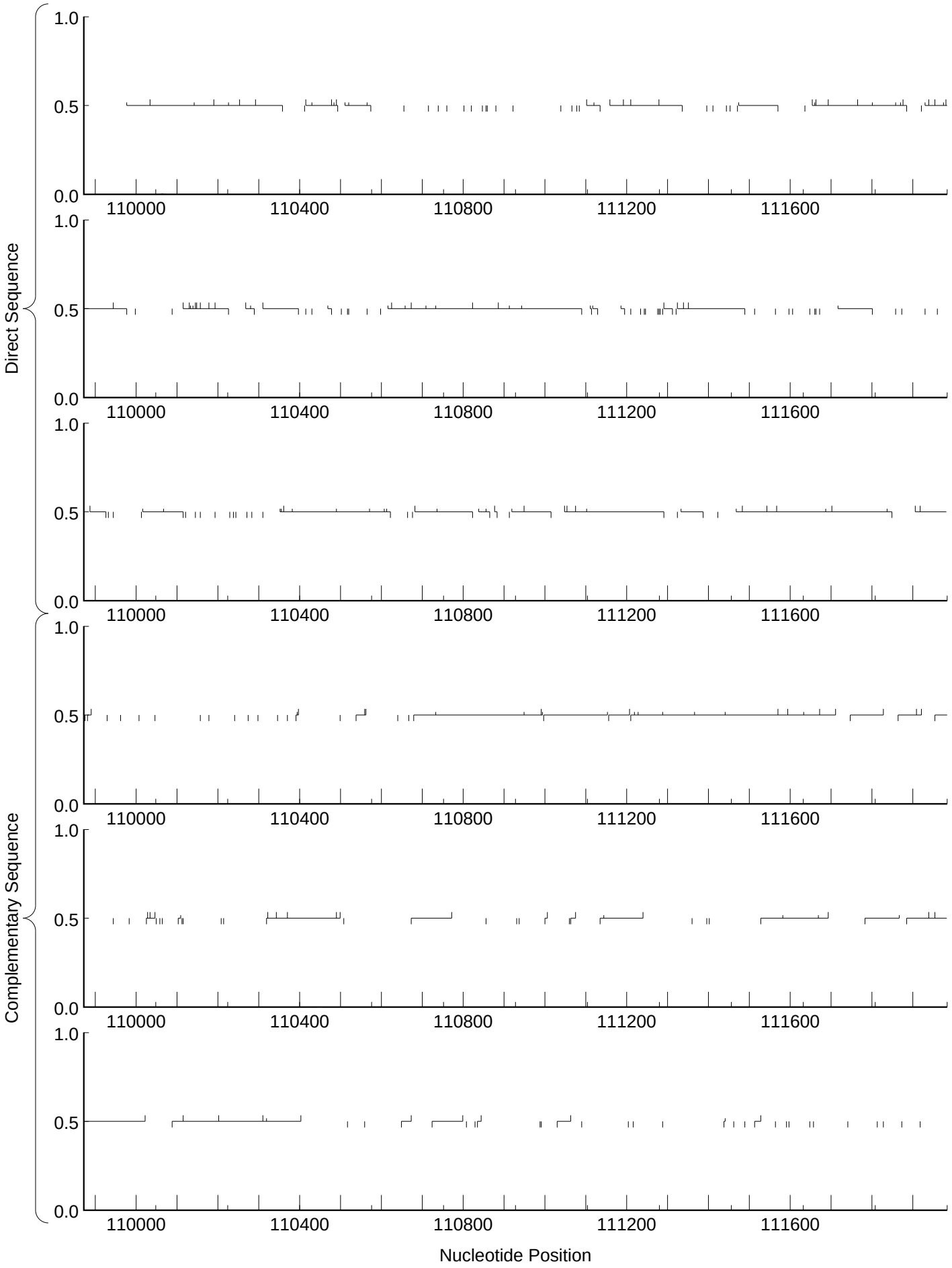
tarPlatinum complete sequence, 133886 bp including 12199 bp terminal repeat, Cluster BE2, Order 4, Window 96, Step 12, 52/65



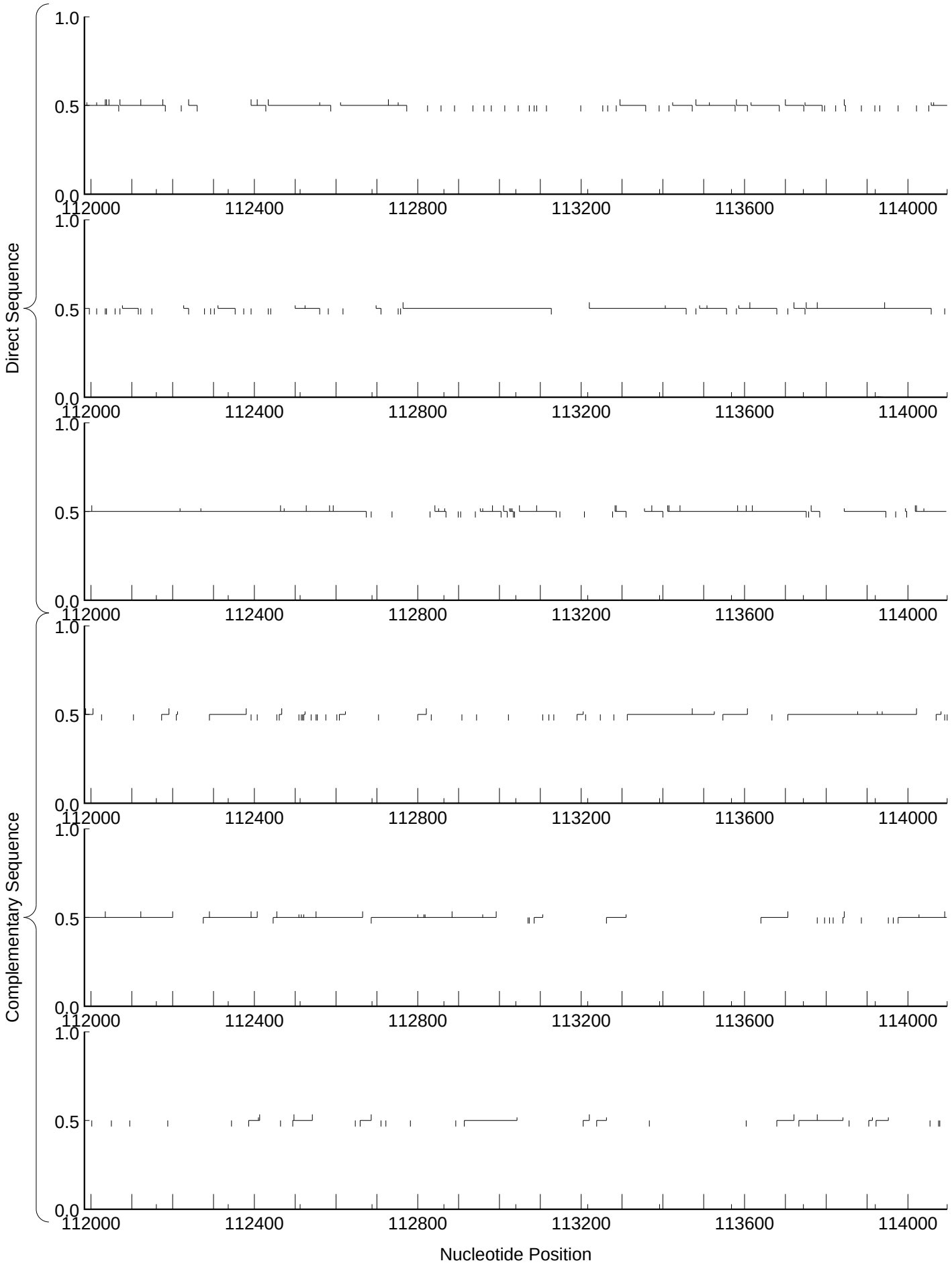
tarPlatinum complete sequence, 133886 bp including 12199 bp terminal repeat, Cluster BE2, Order 4, Window 96, Step 12, 53/65



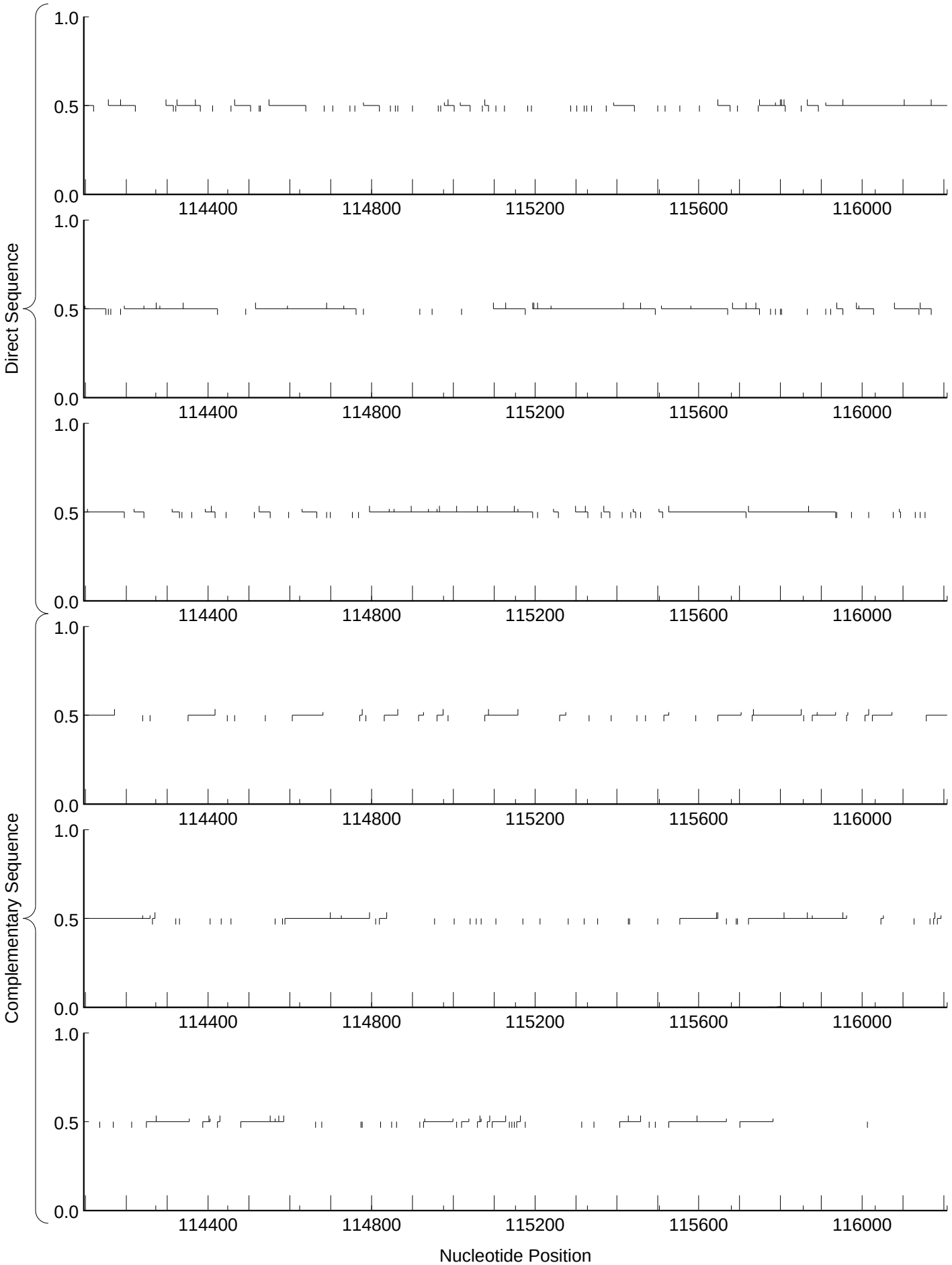
tarPlatinum complete sequence, 133886 bp including 12199 bp terminal repeat, Cluster BE2, Order 4, Window 96, Step 12, 54/65



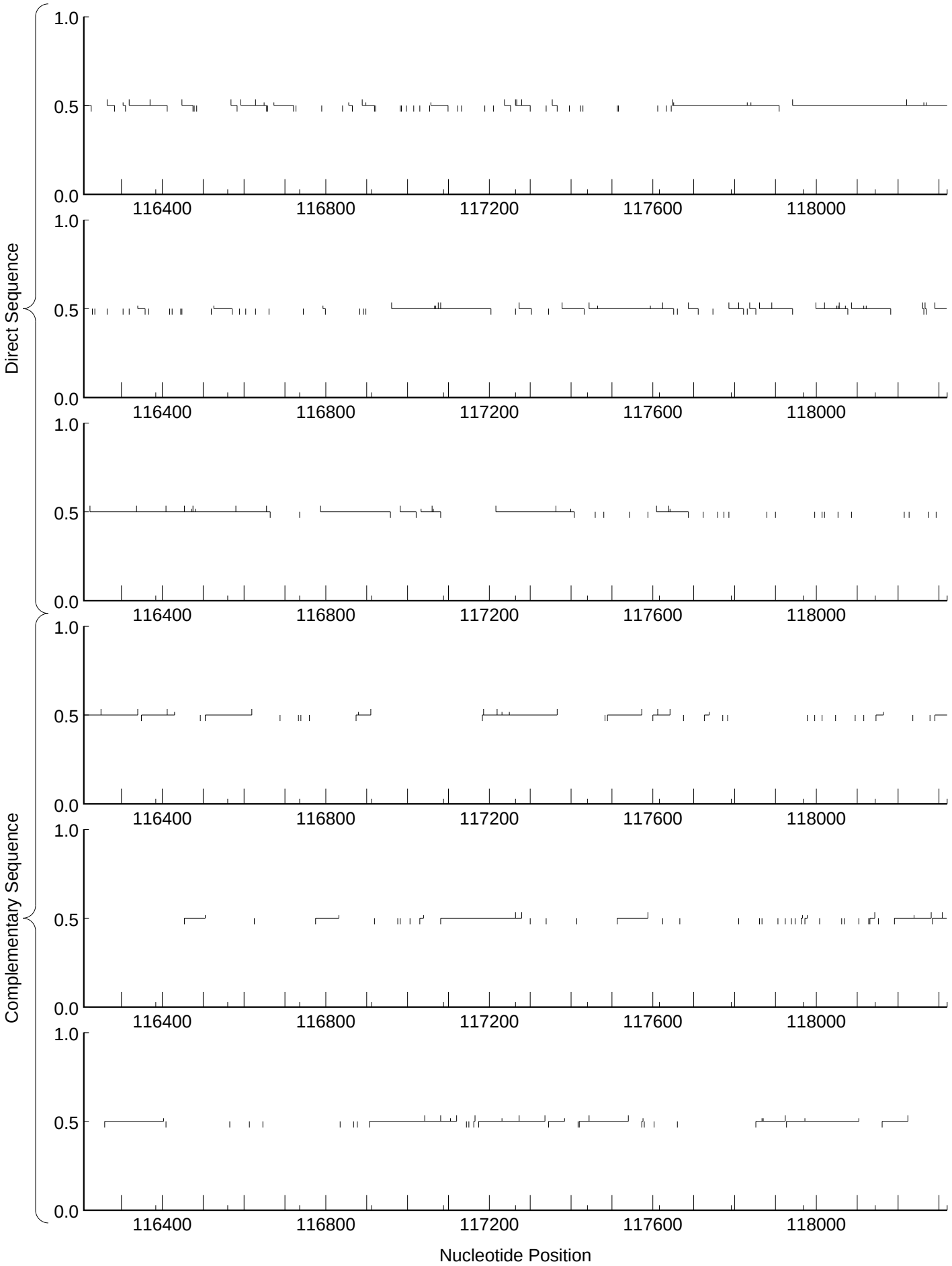
tarPlatinum complete sequence, 133886 bp including 12199 bp terminal repeat, Cluster BE2, Order 4, Window 96, Step 12, 55/65



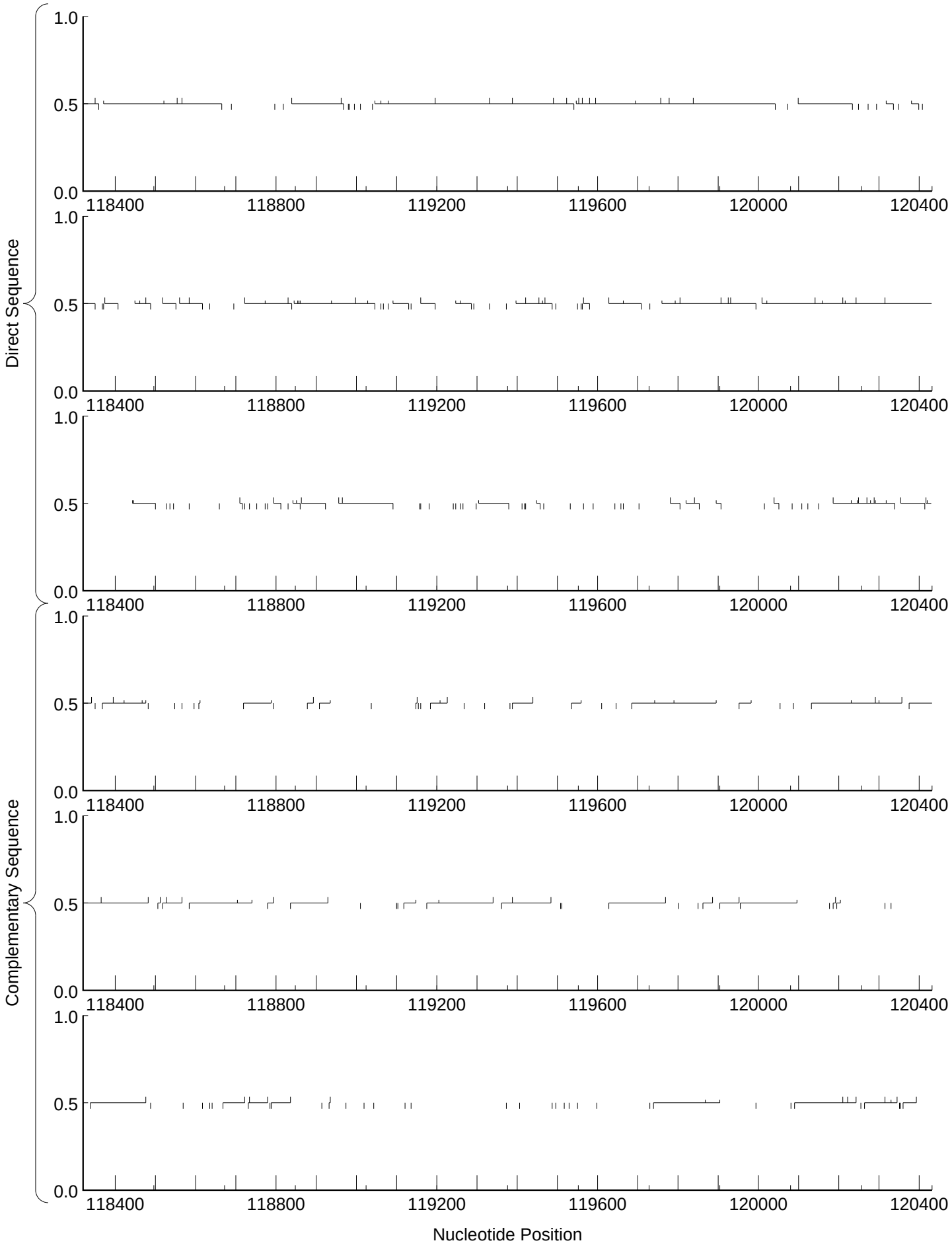
tarPlatinum complete sequence, 133886 bp including 12199 bp terminal repeat, Cluster BE2, Order 4, Window 96, Step 12, 56/65



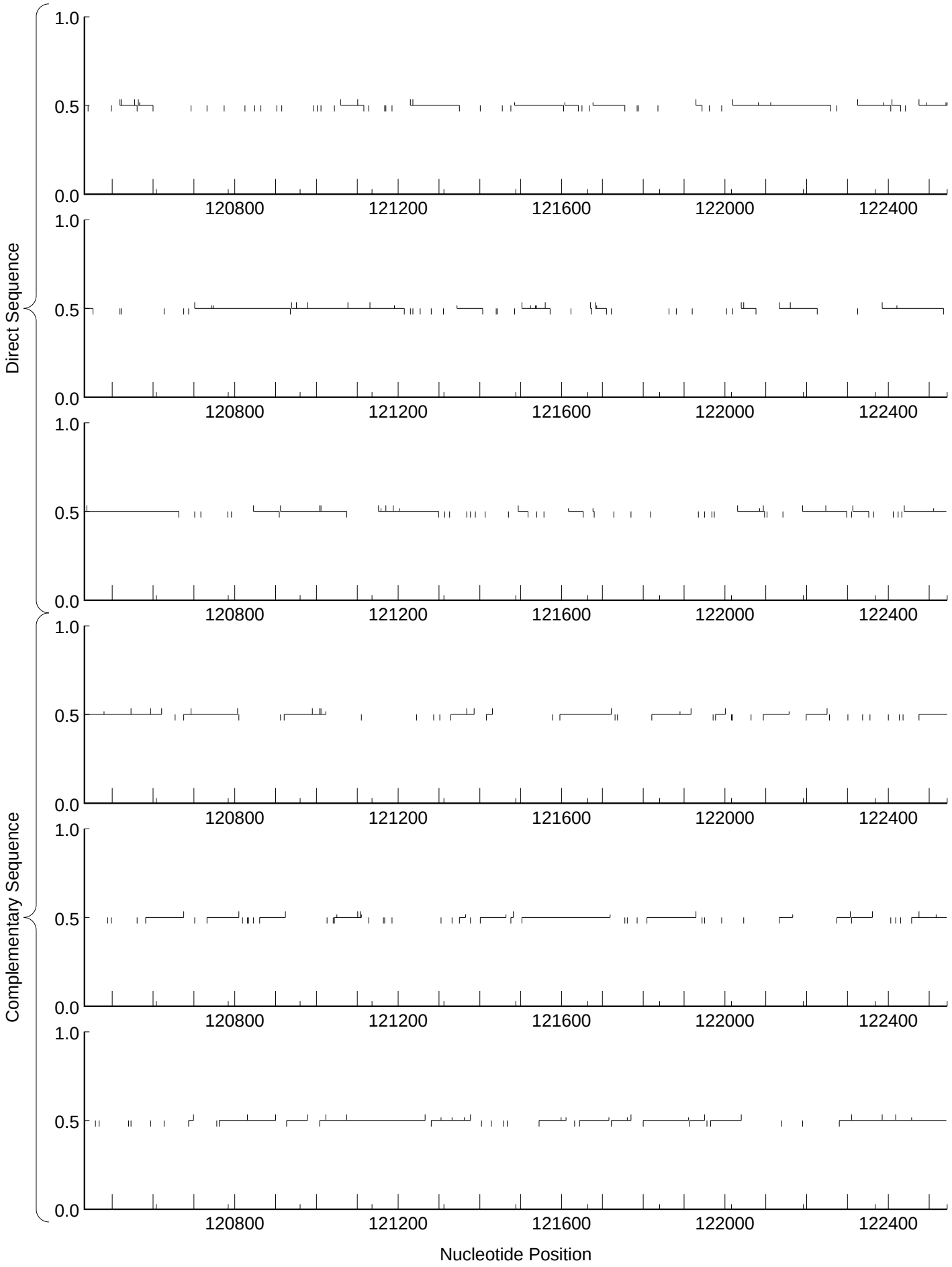
tarPlatinum complete sequence, 133886 bp including 12199 bp terminal repeat, Cluster BE2, Order 4, Window 96, Step 12, 57/65



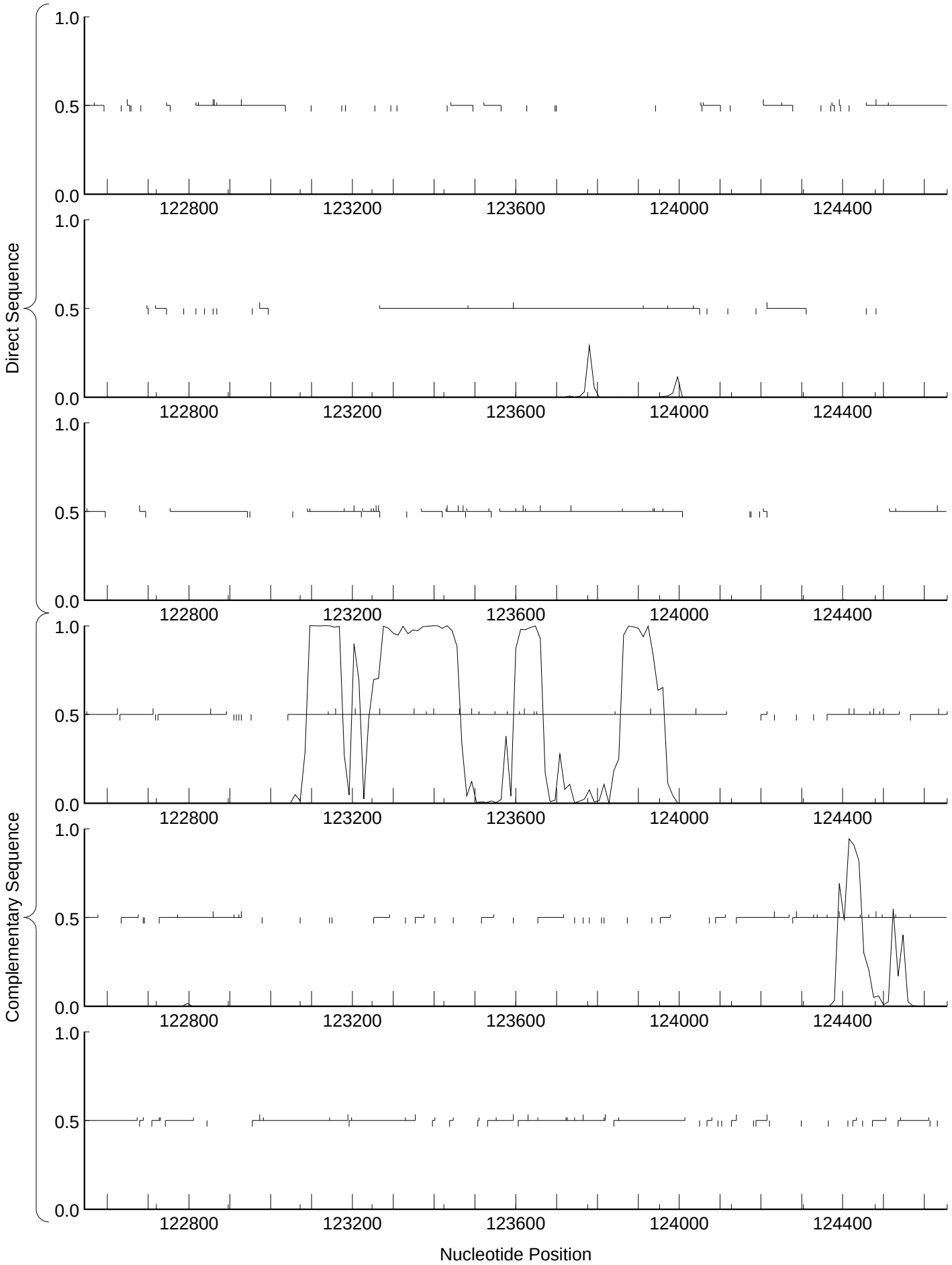
tarPlatinum complete sequence, 133886 bp including 12199 bp terminal repeat, Cluster BE2, Order 4, Window 96, Step 12, 58/65



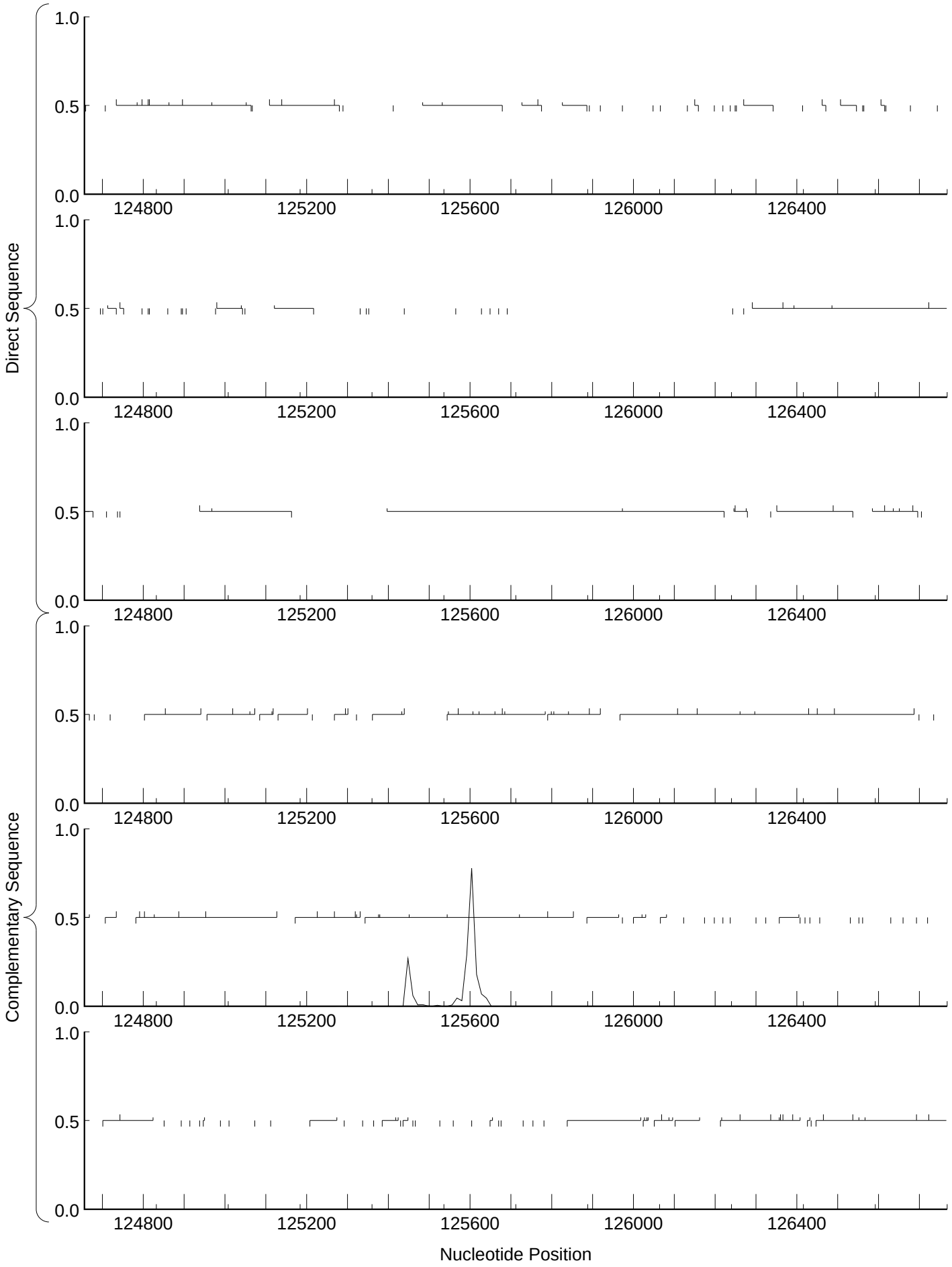
tarPlatinum complete sequence, 133886 bp including 12199 bp terminal repeat, Cluster BE2, Order 4, Window 96, Step 12, 59/65



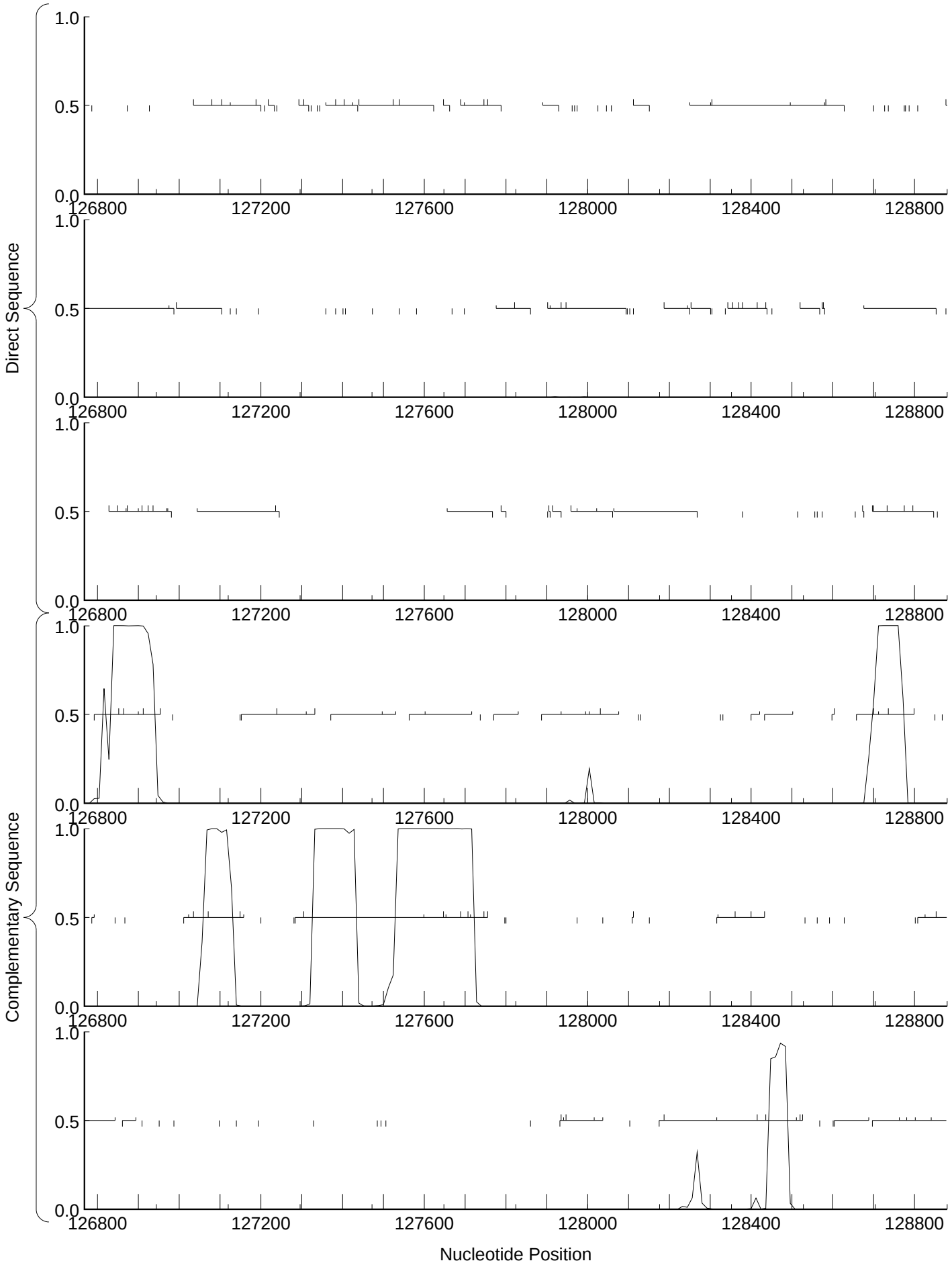
tarPlatinum complete sequence, 133886 bp including 12199 bp terminal repeat, Cluster BE2, Order 4, Window 96, Step 12, 60/65



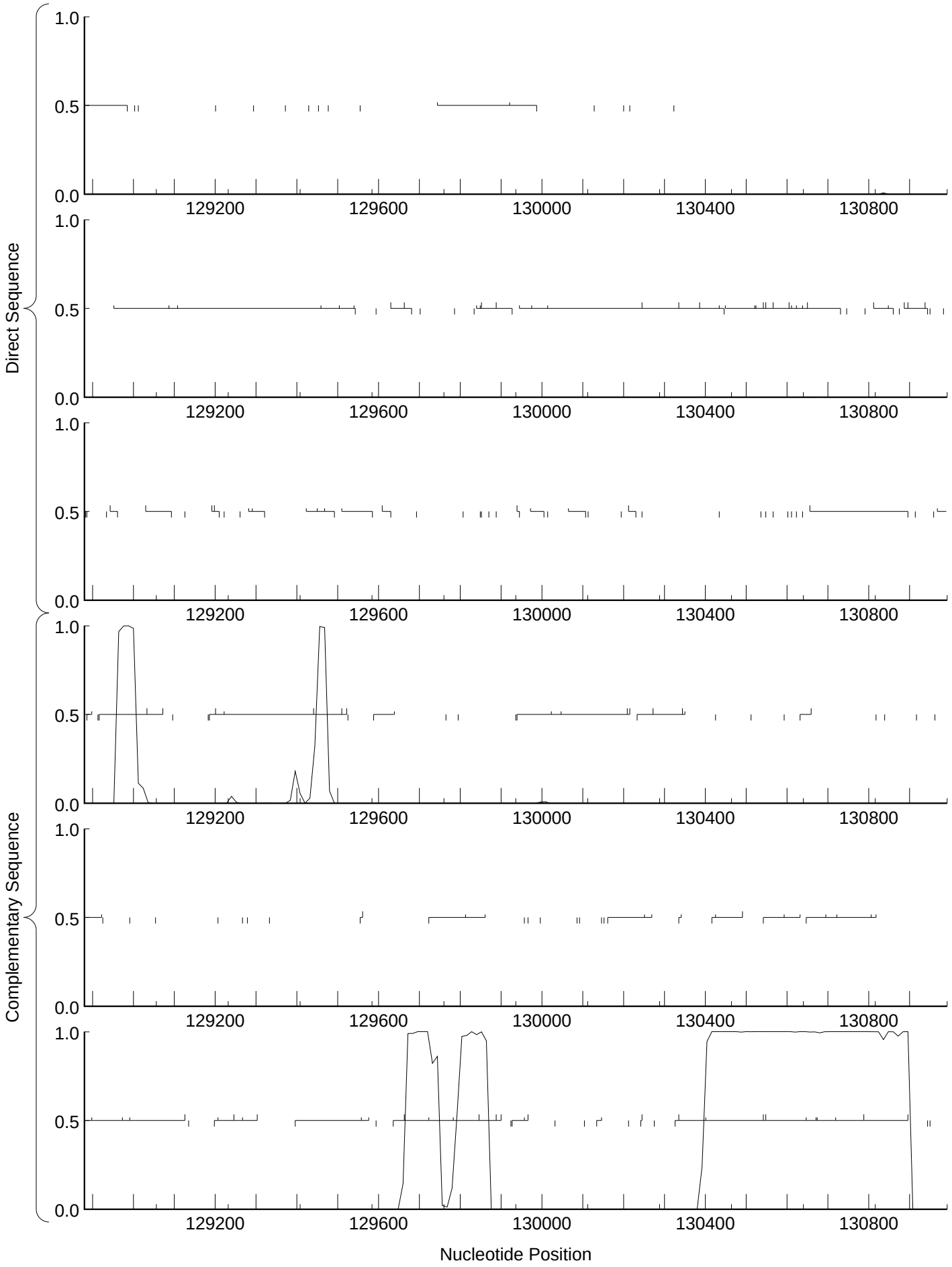
tarPlatinum complete sequence, 133886 bp including 12199 bp terminal repeat, Cluster BE2, Order 4, Window 96, Step 12, 61/65



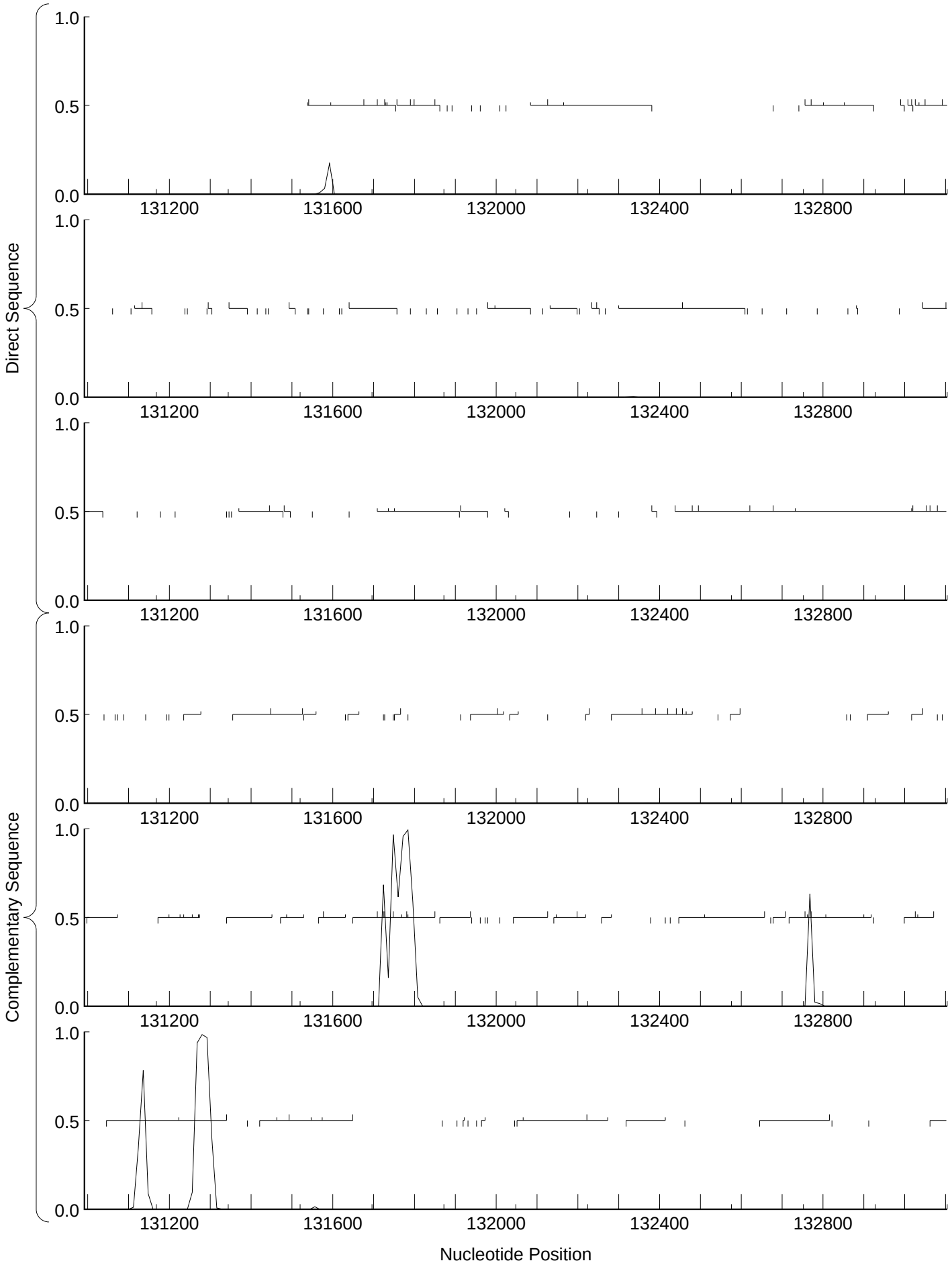
tarPlatinum complete sequence, 133886 bp including 12199 bp terminal repeat, Cluster BE2, Order 4, Window 96, Step 12, 62/65



tarPlatinum complete sequence, 133886 bp including 12199 bp terminal repeat, Cluster BE2, Order 4, Window 96, Step 12, 63/65



tarPlatinum complete sequence, 133886 bp including 12199 bp terminal repeat, Cluster BE2, Order 4, Window 96, Step 12, 64/65



tarPlatinum complete sequence, 133886 bp including 12199 bp terminal repeat, Cluster BE2, Order 4, Window 96, Step 12, 65/65

