GeneMark

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

PROGRAM INFORMATION

```
Sequence         :  Mycobacterium phage ET08
Analysis Date    :  3/28/24 at 20:11:16
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Sequence Length  :  155445 bp
GC Content       :   64.60%

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

PS-Version       :  1.2

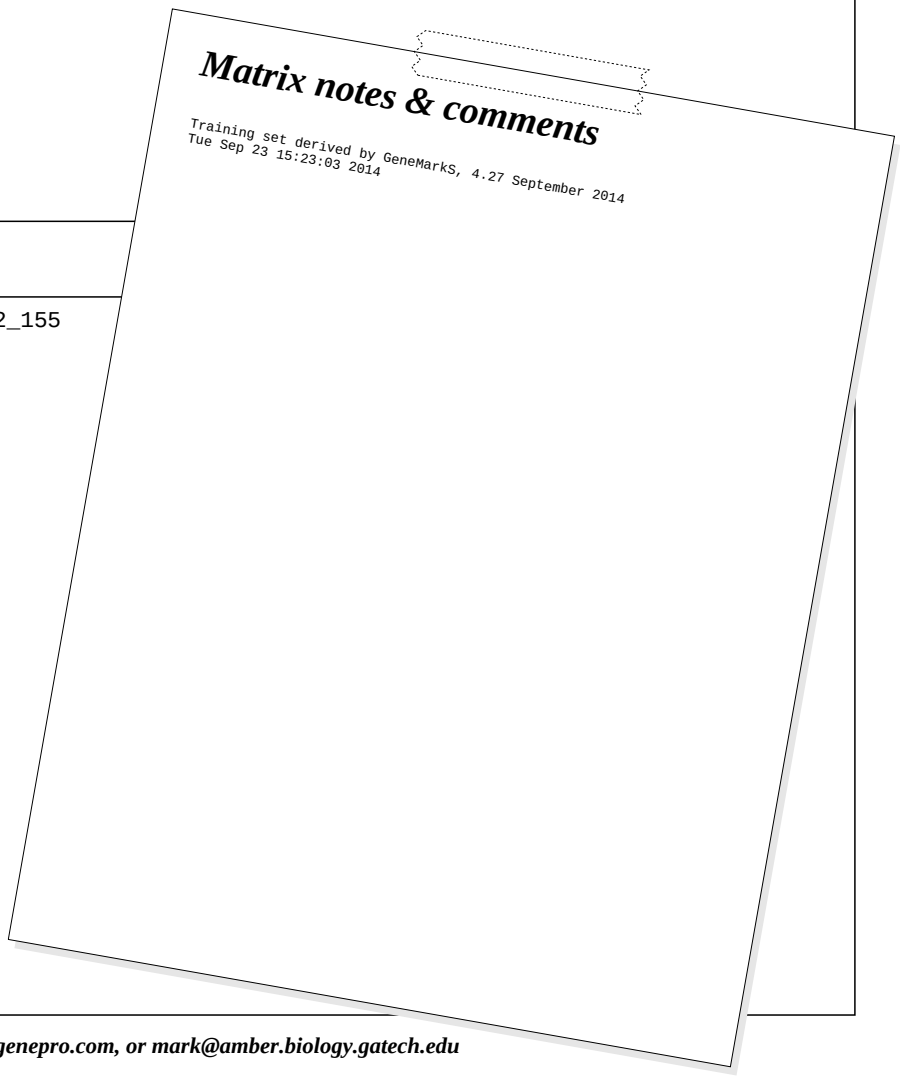
GeneMark Options :  PostScript graph,
                    Mark ORFs / splice sites,
                    List ORFs,
                    List regions and/or splice sites,
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MATRIX INFORMATION

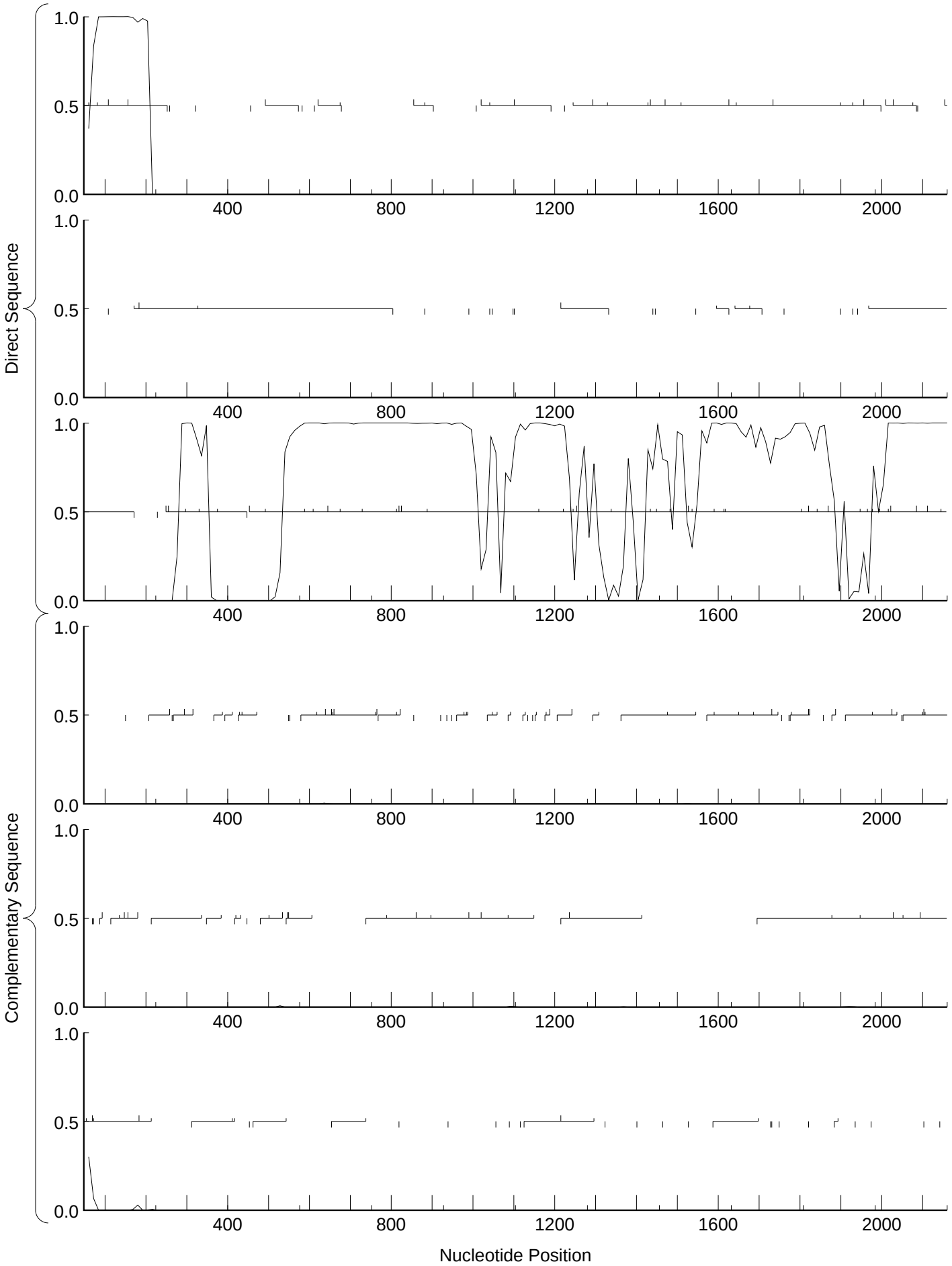
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Author     :  -

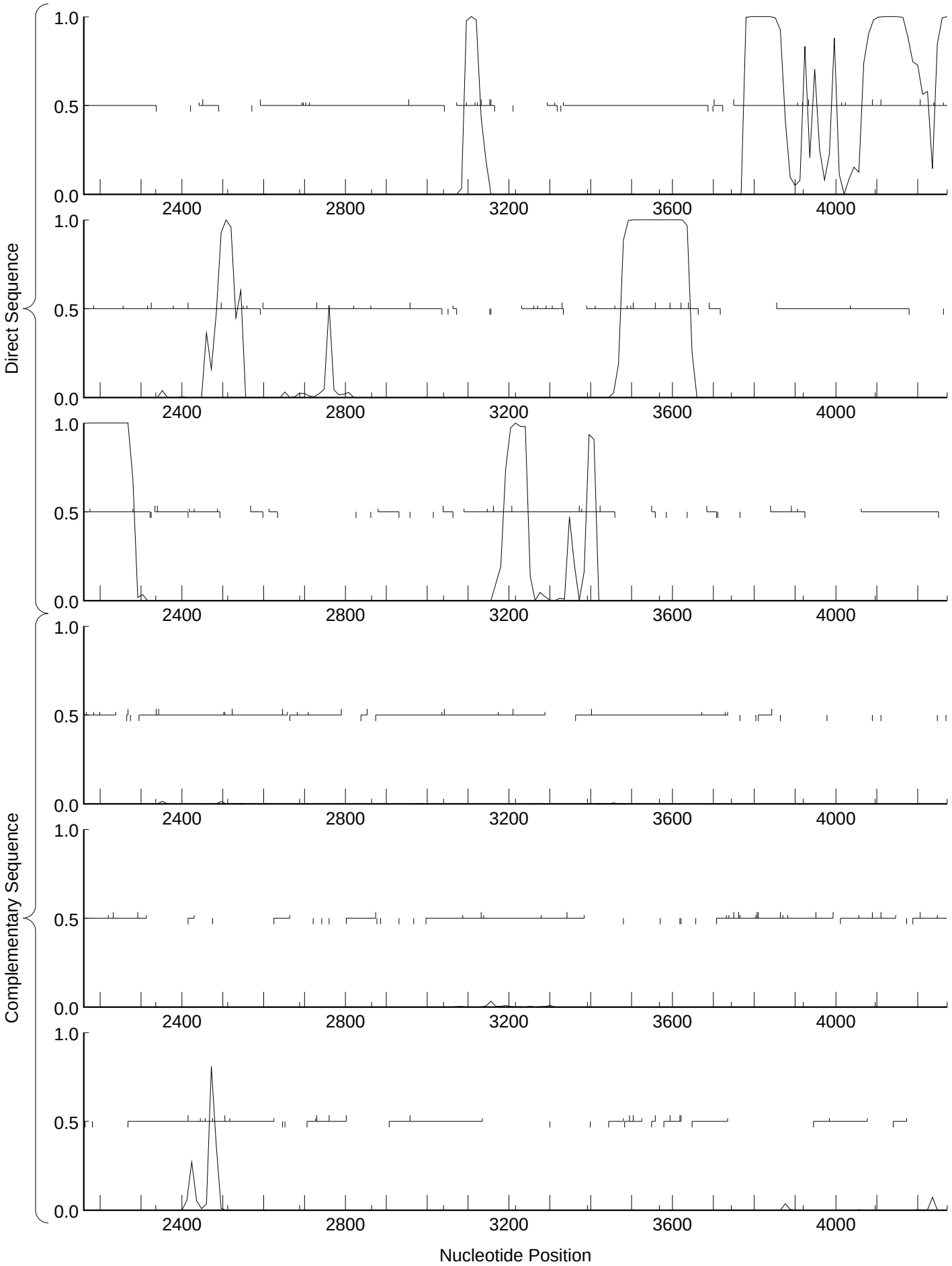
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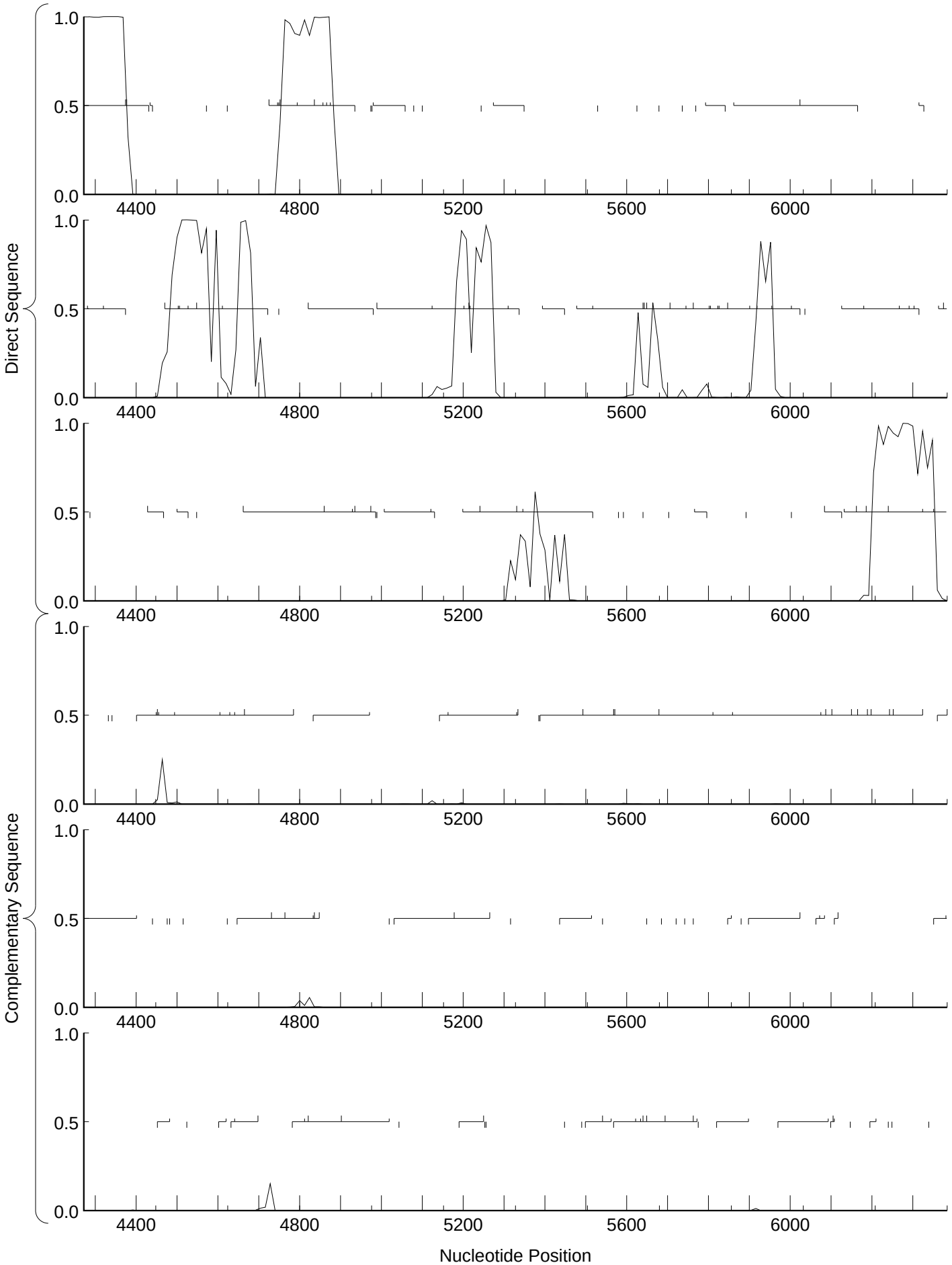

  Send questions / comments to:
      Dr. M. Borodovsky
      Georgia Institute of Technology
      School of Biology
      Atlanta, GA  30332-0230
```

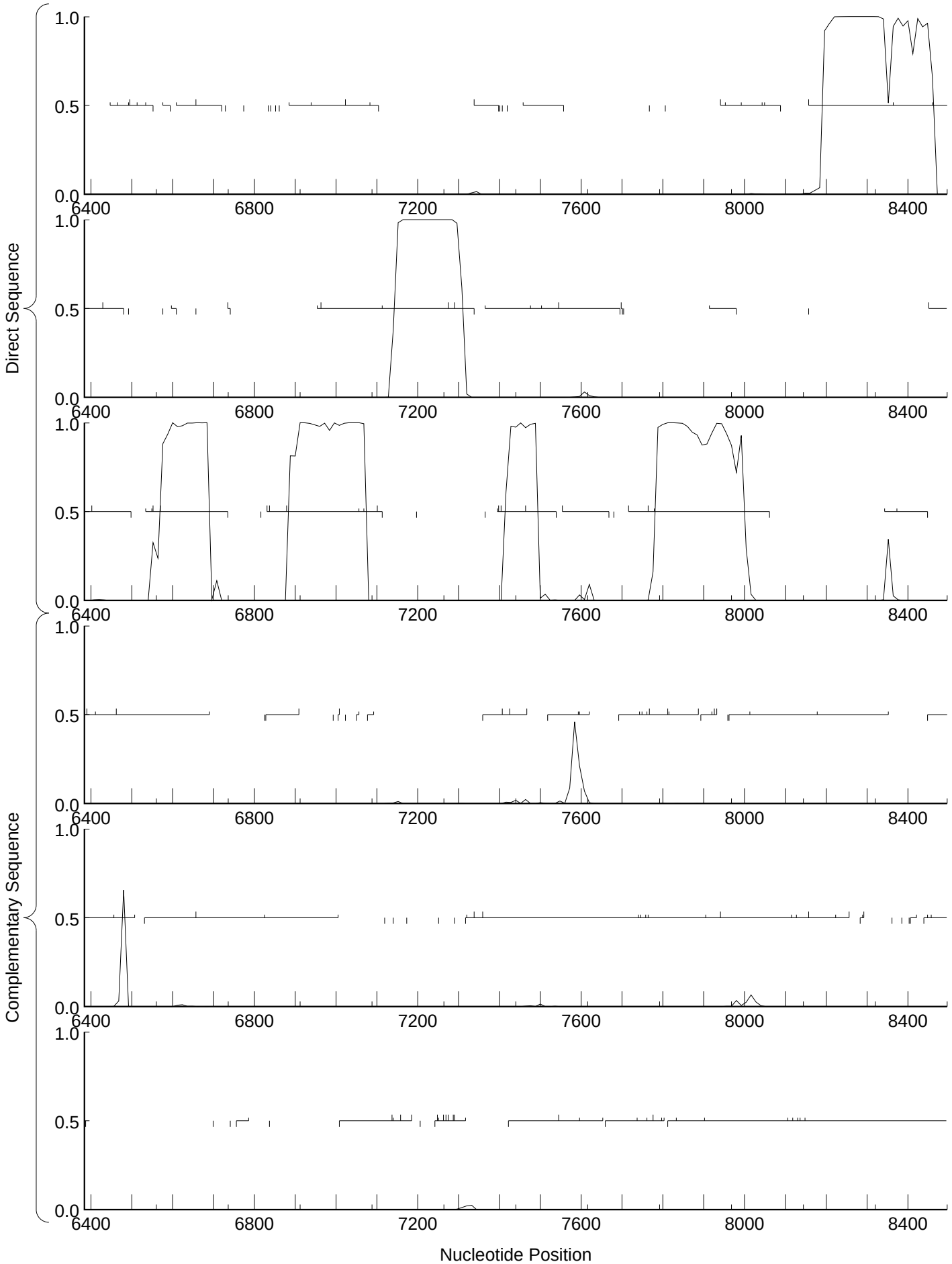


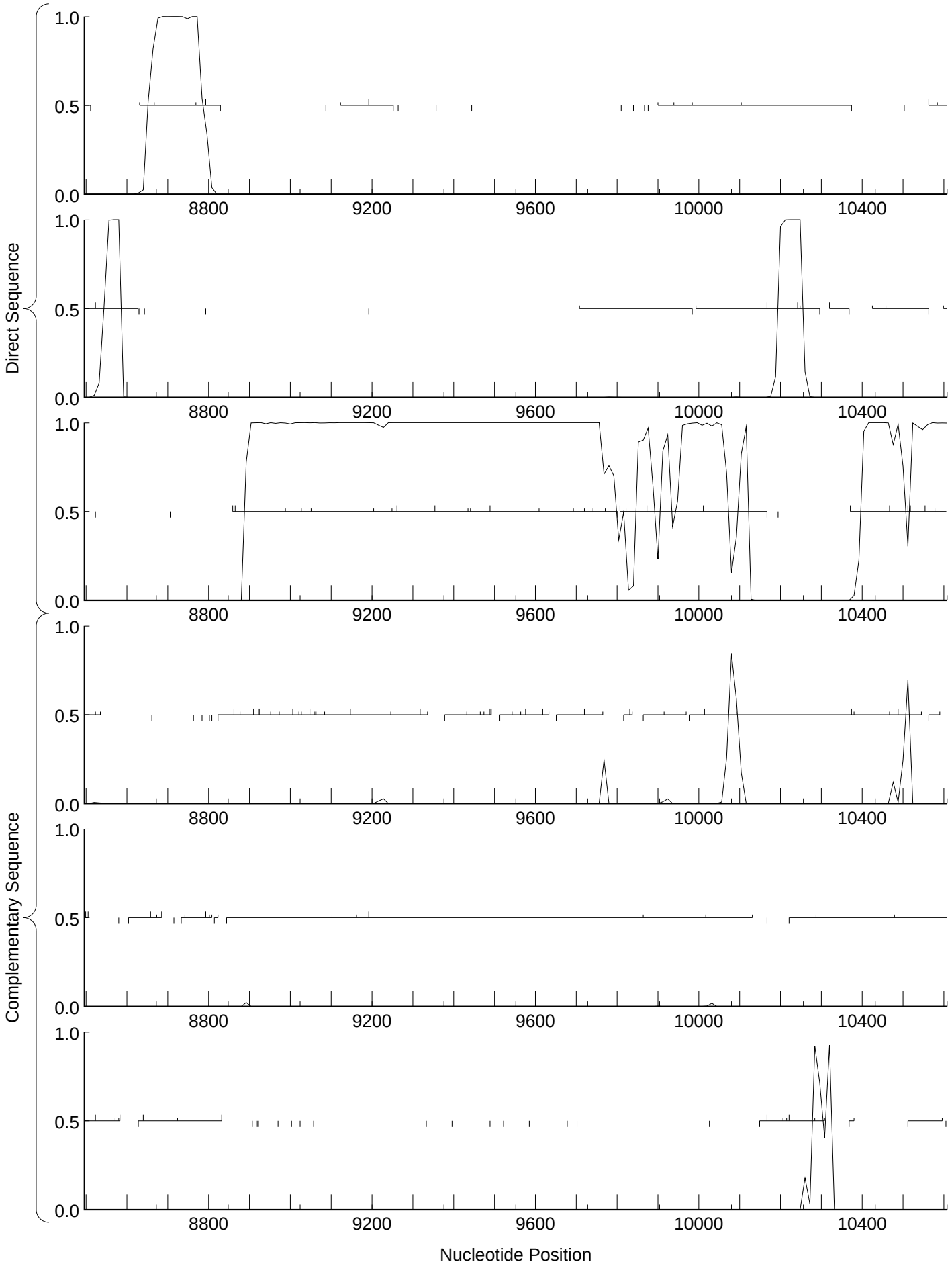
For help or information, send electronic mail to: custserv@genepro.com, or mark@amber.biology.gatech.edu

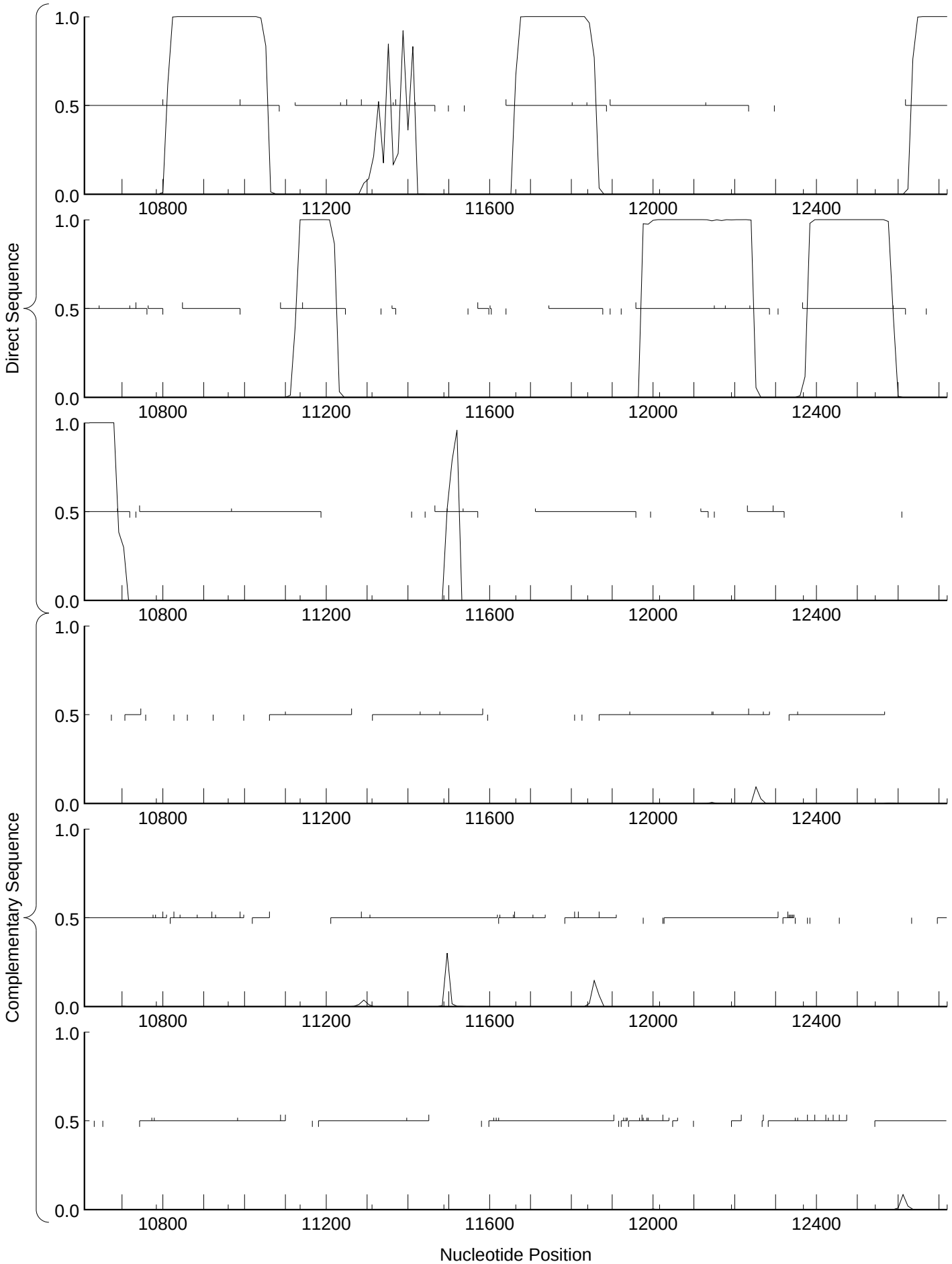

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1600
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Nucleotide Position

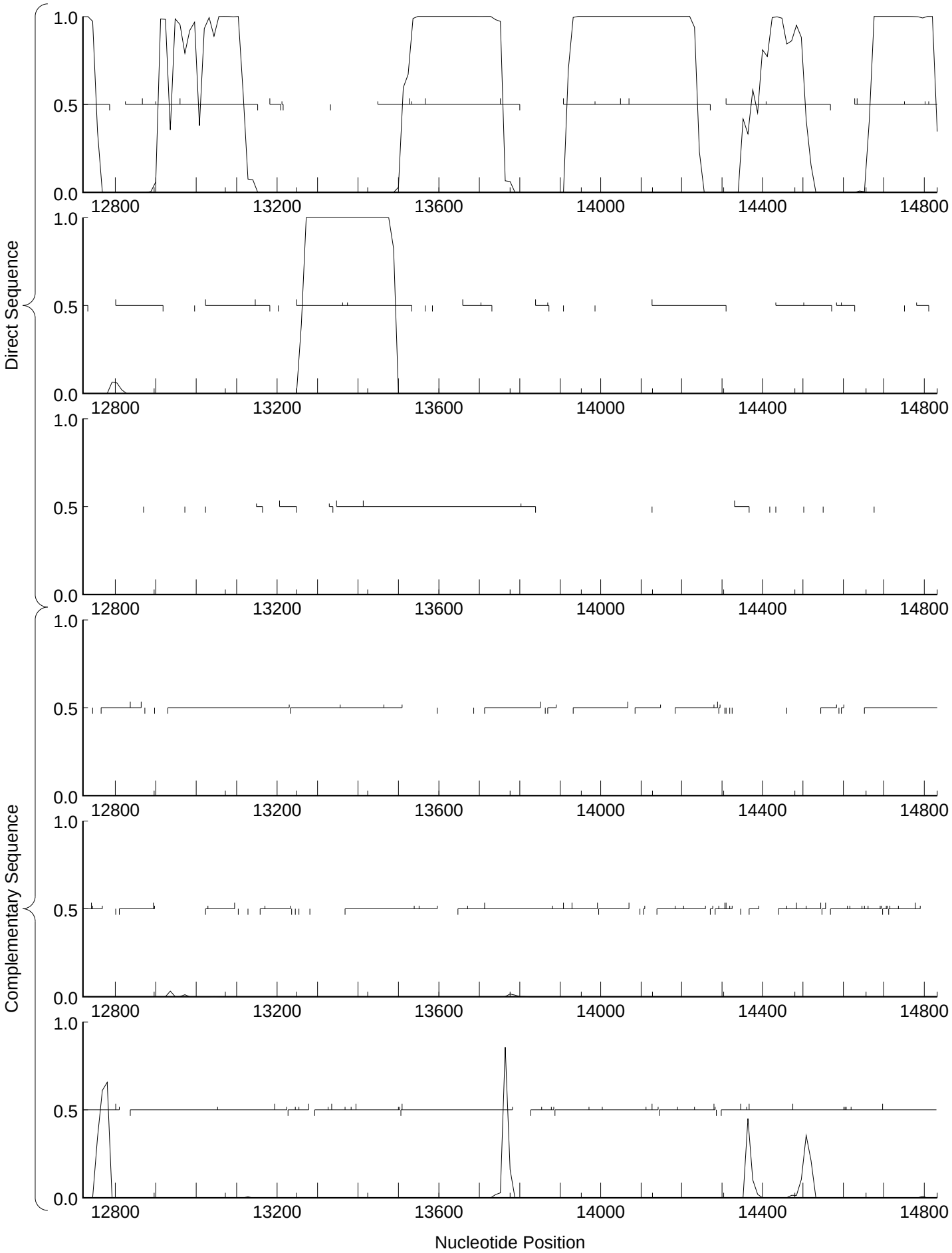

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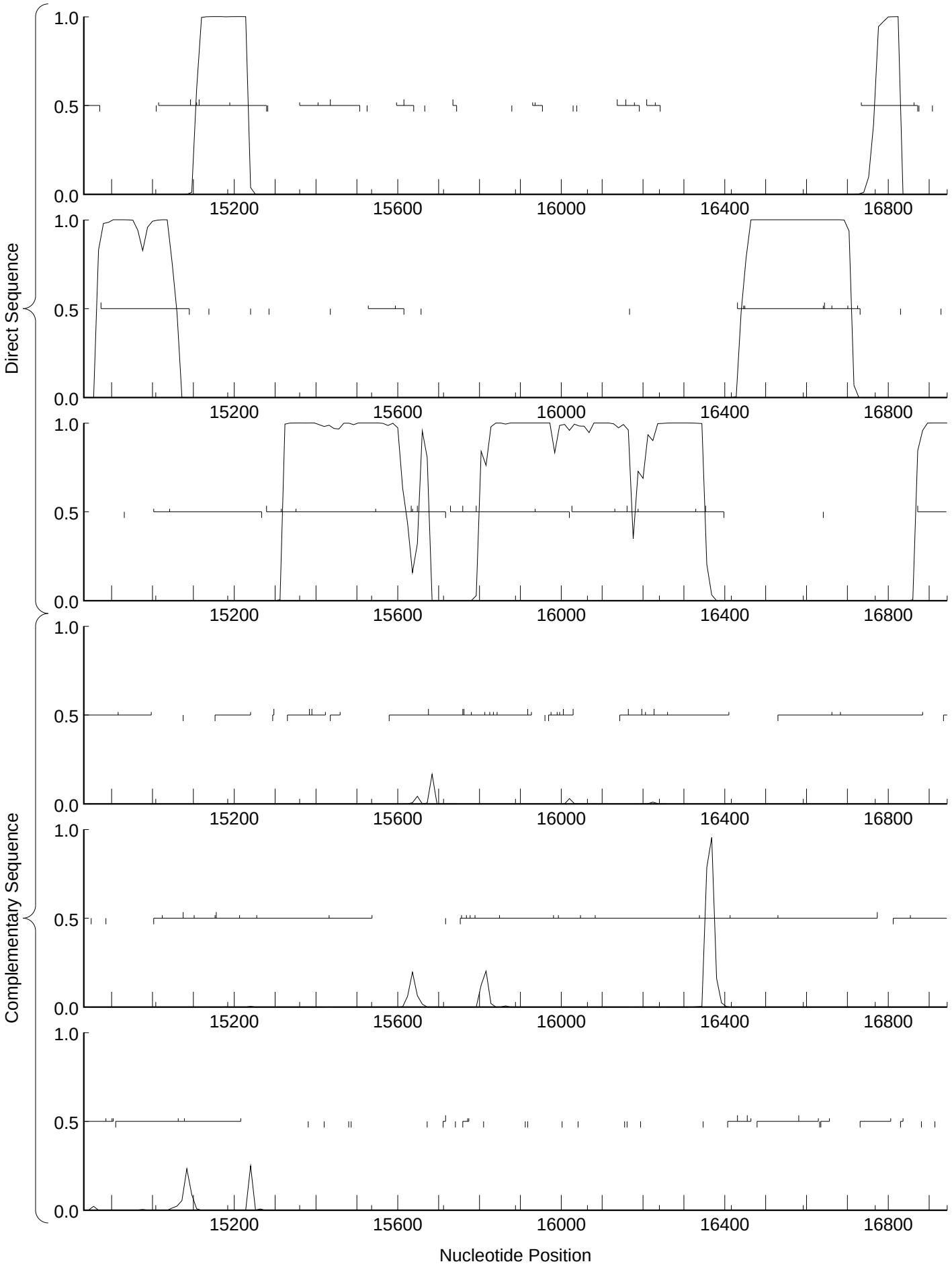

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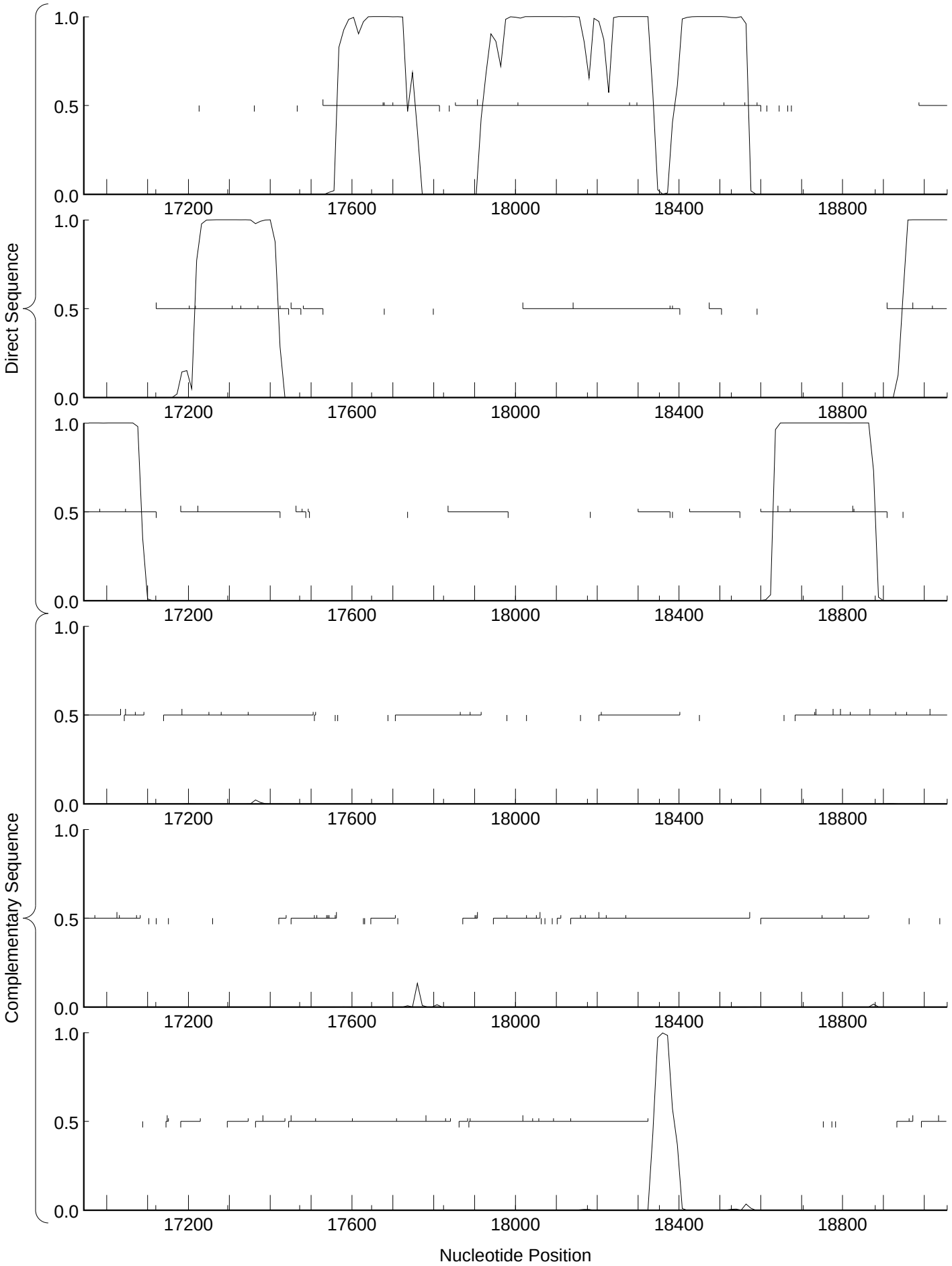

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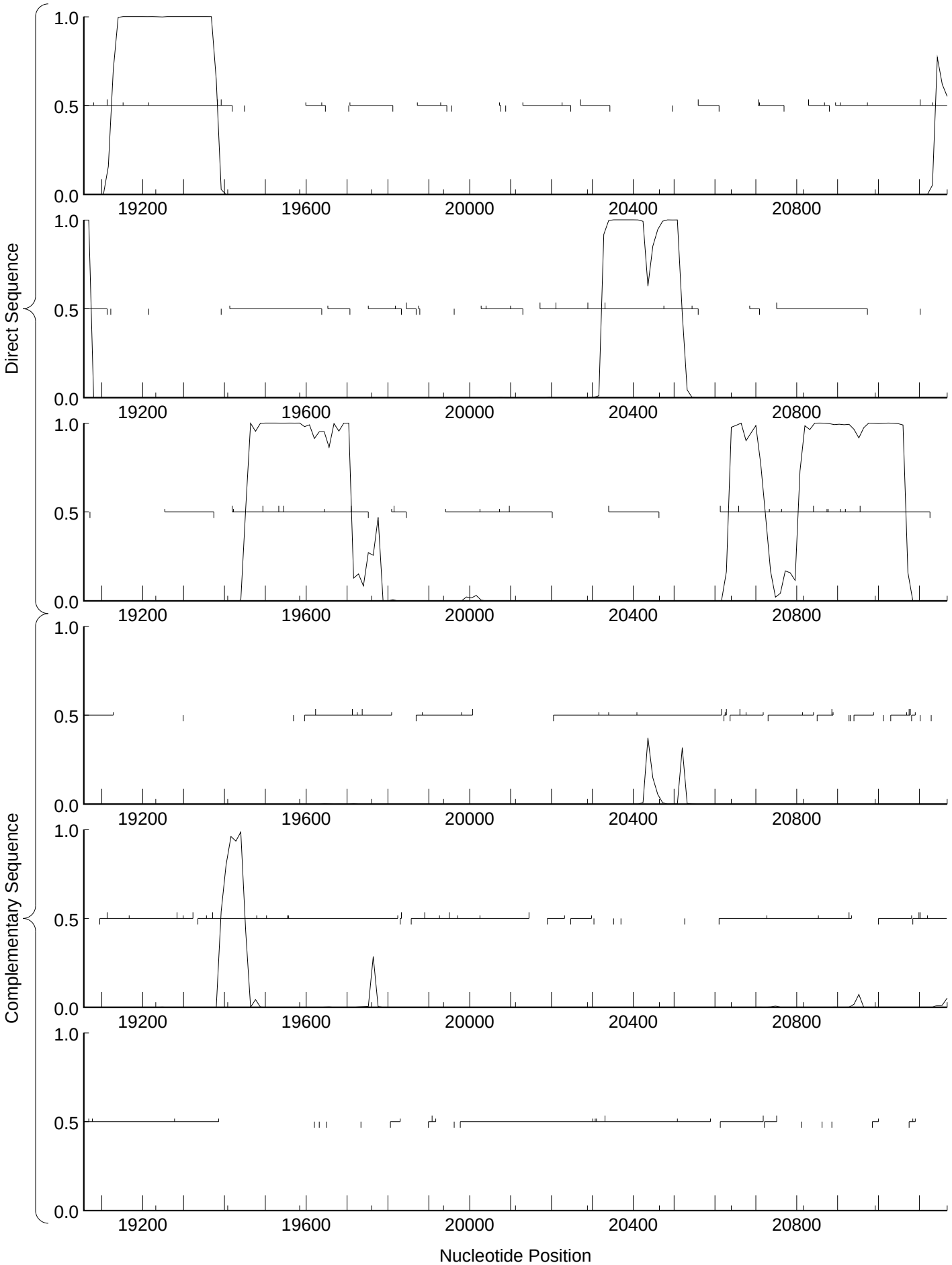

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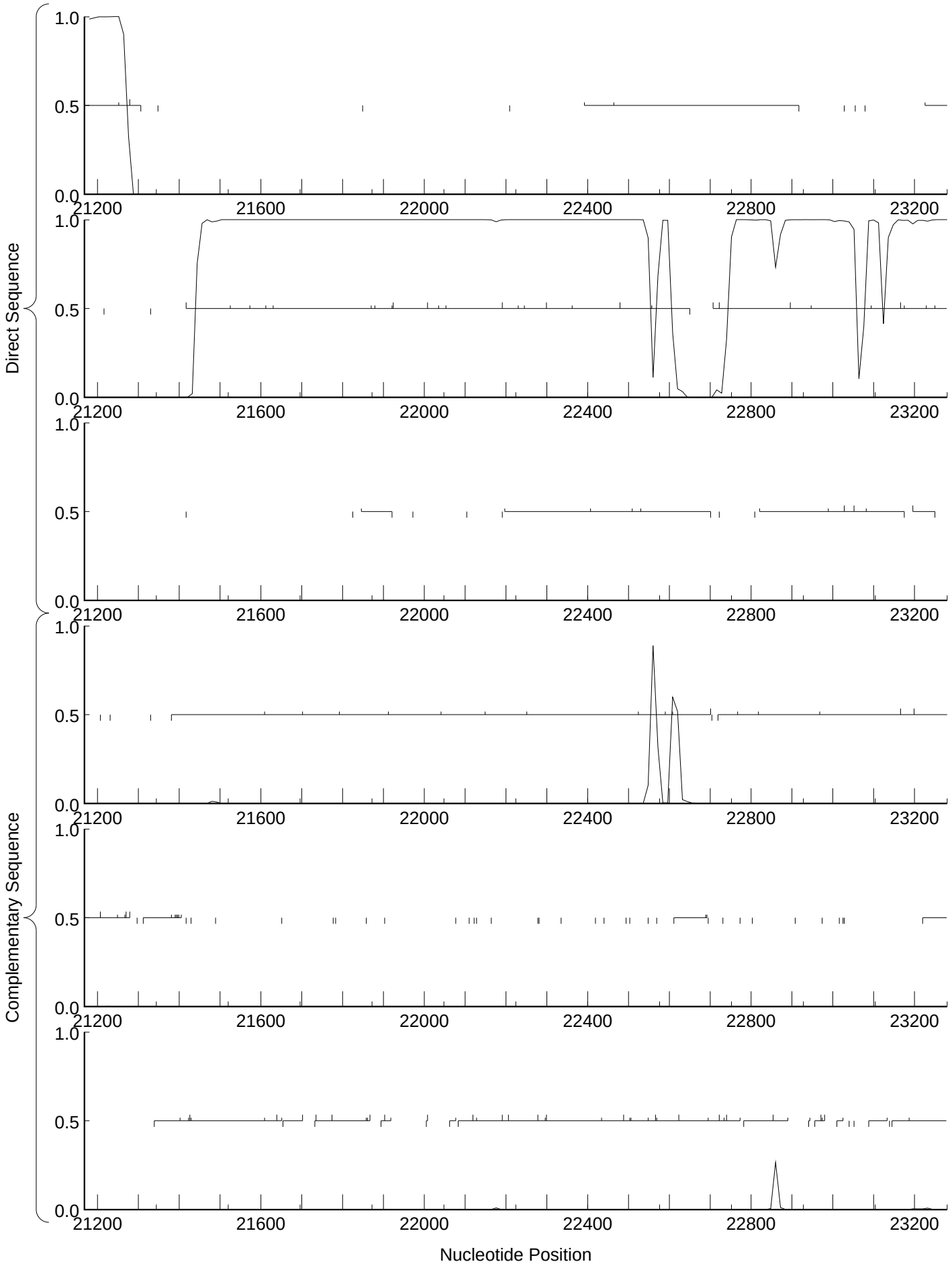

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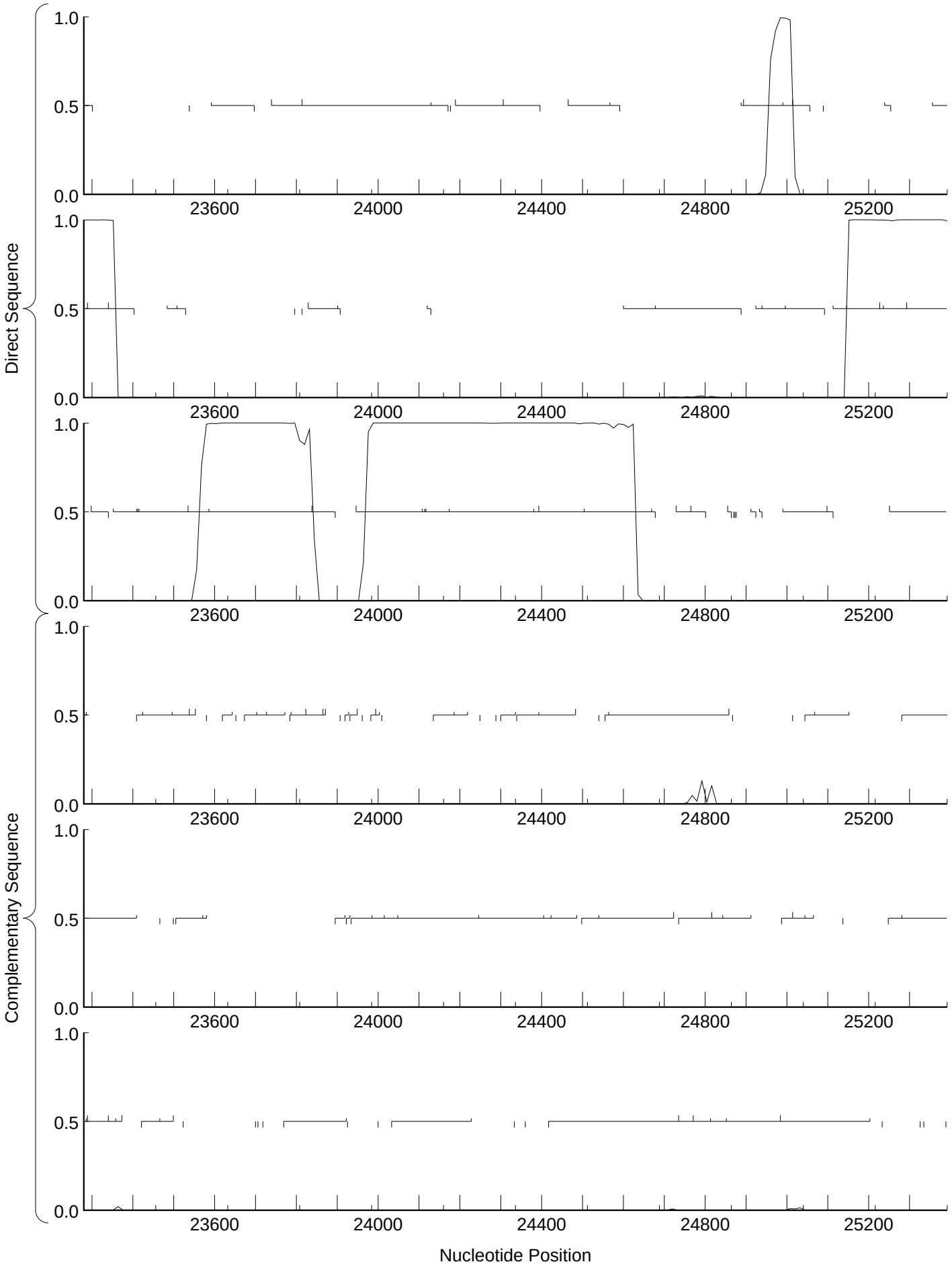

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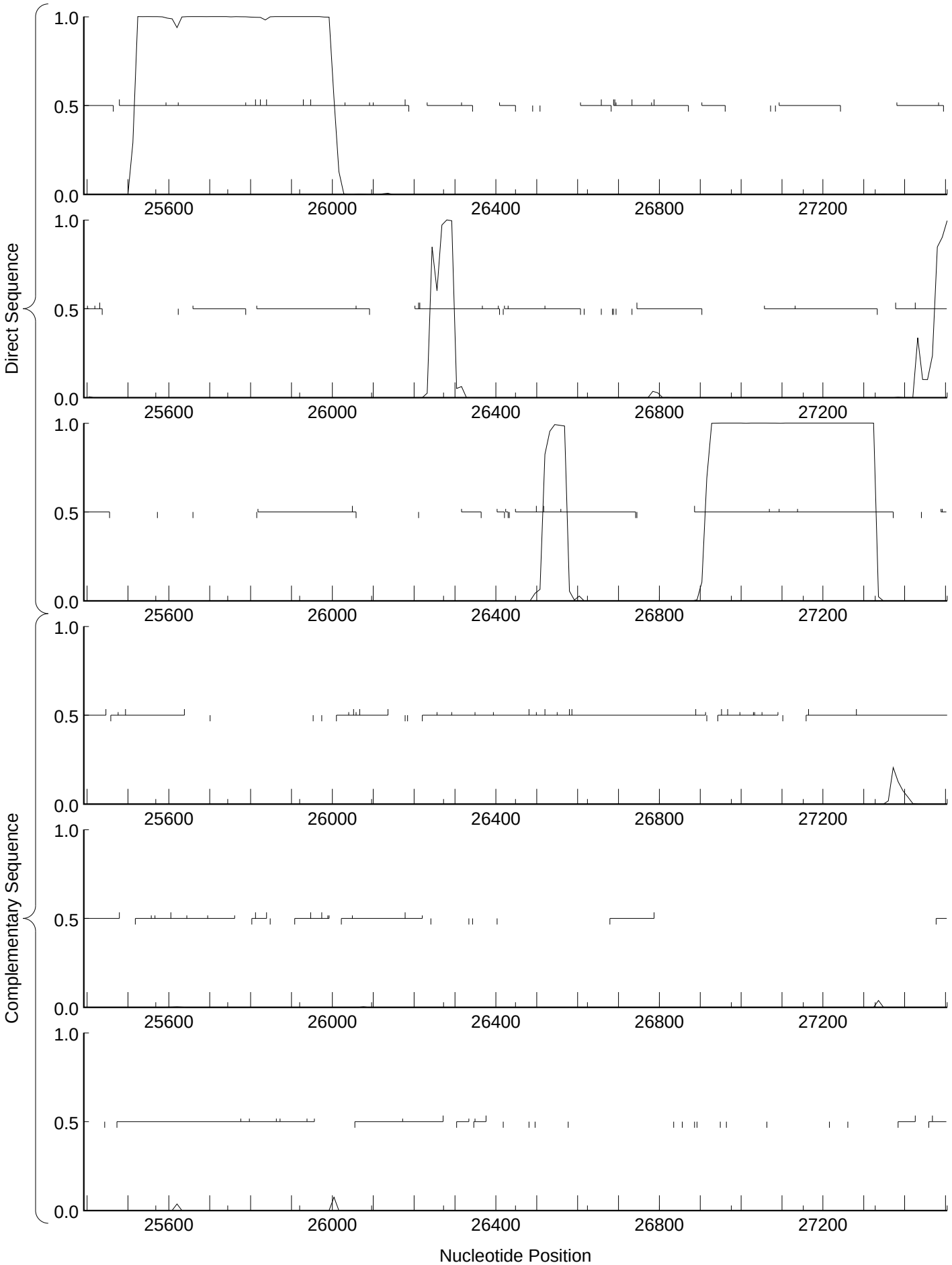

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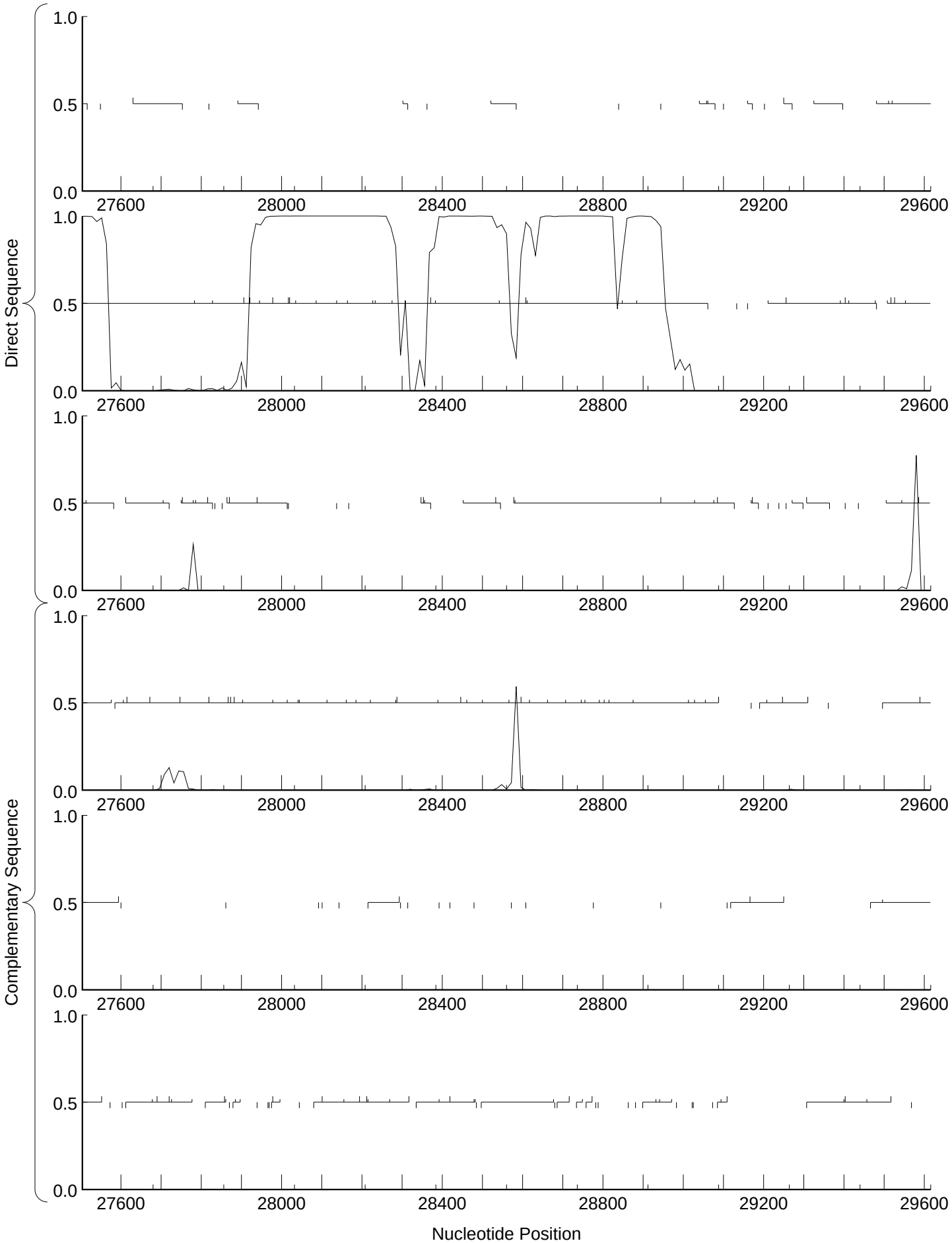

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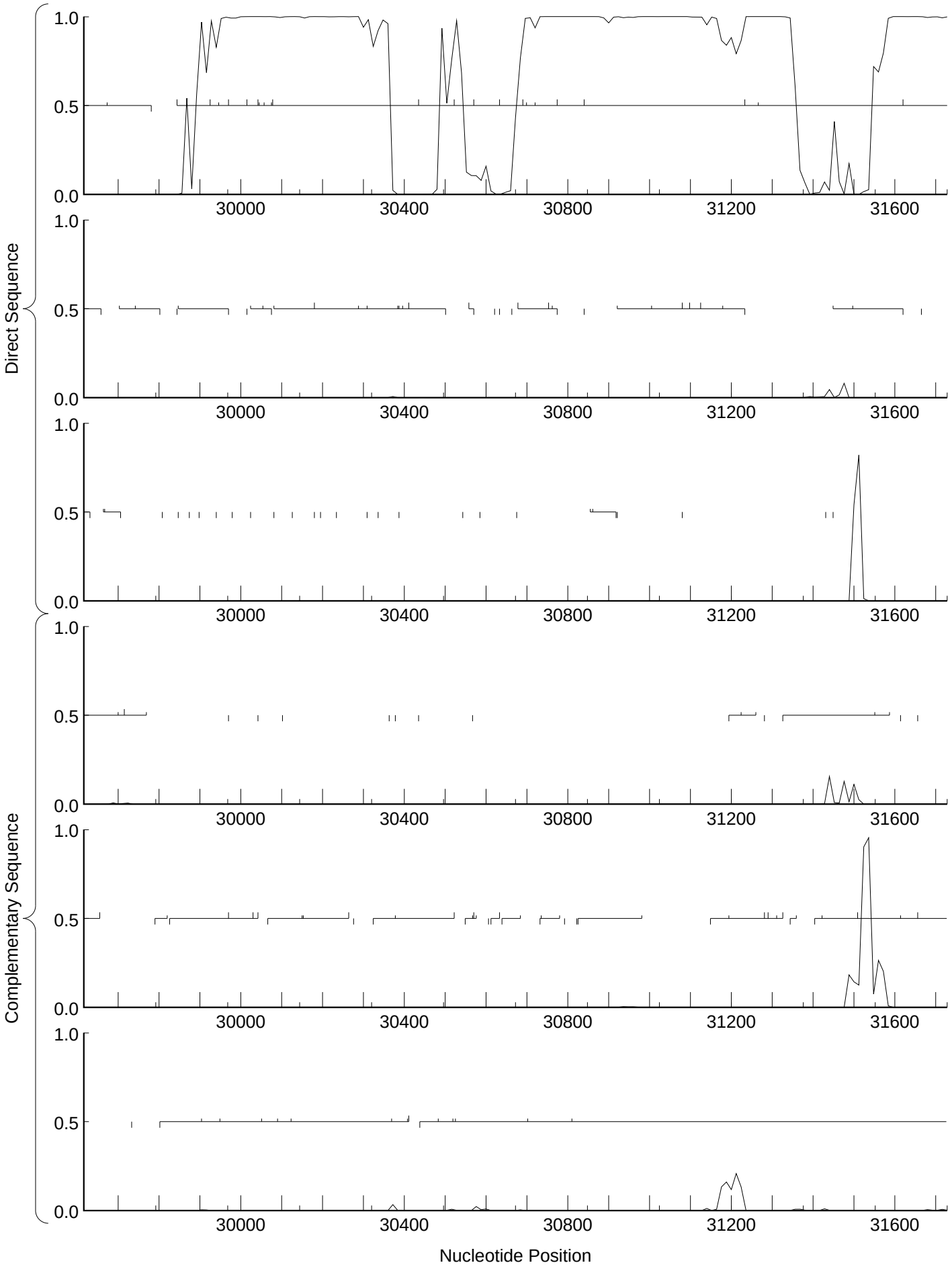

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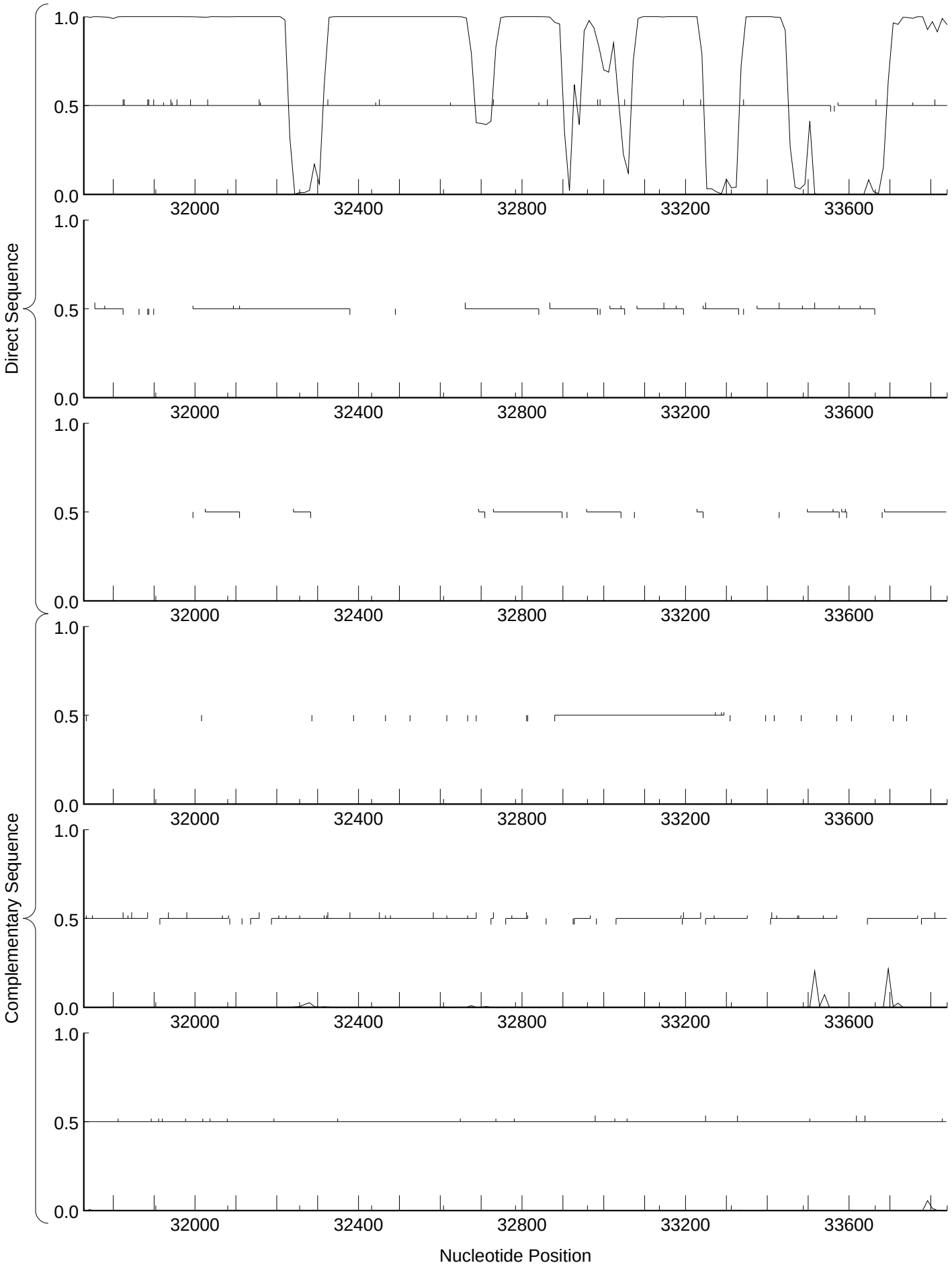

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Complementary Sequence
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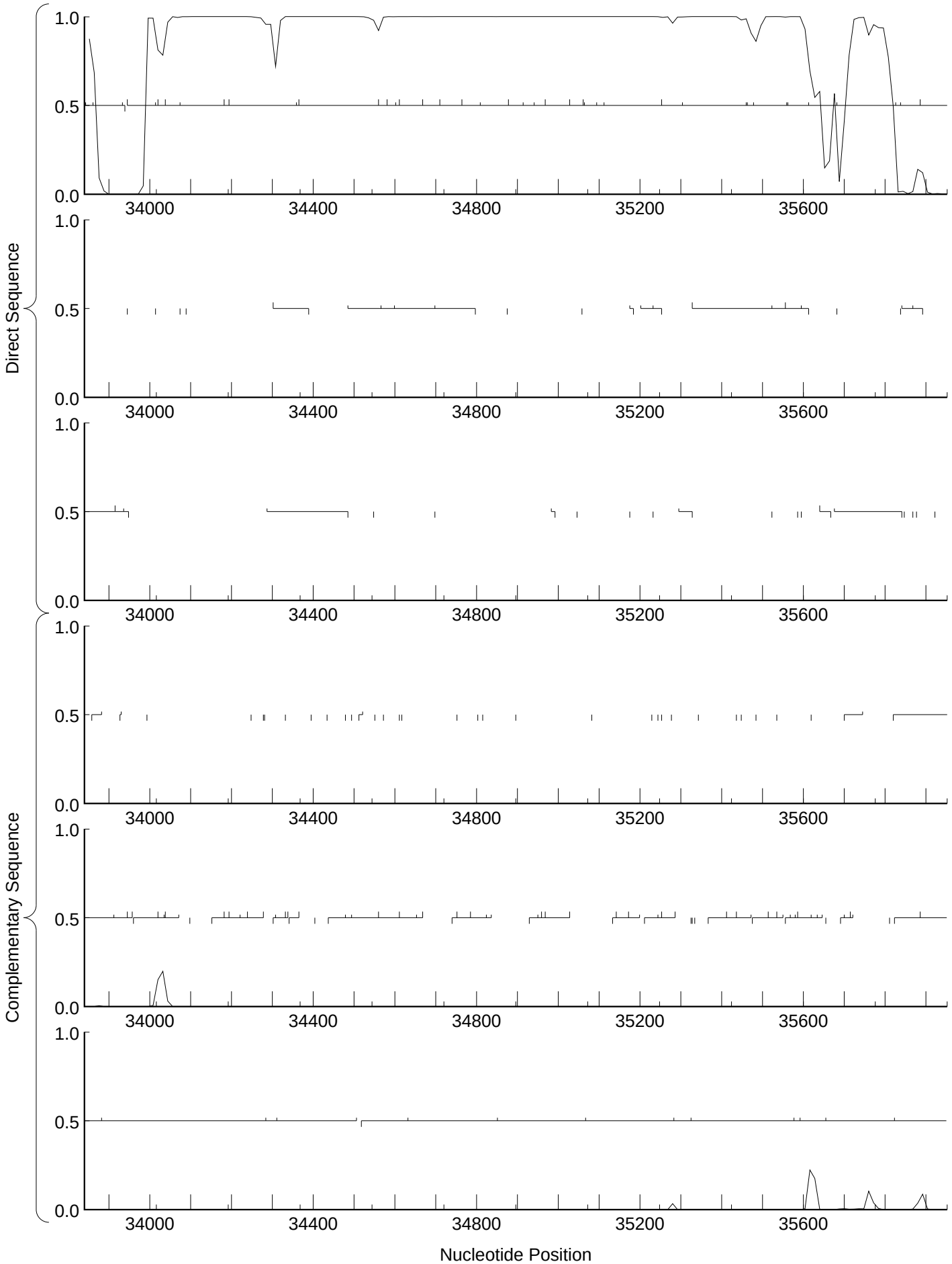

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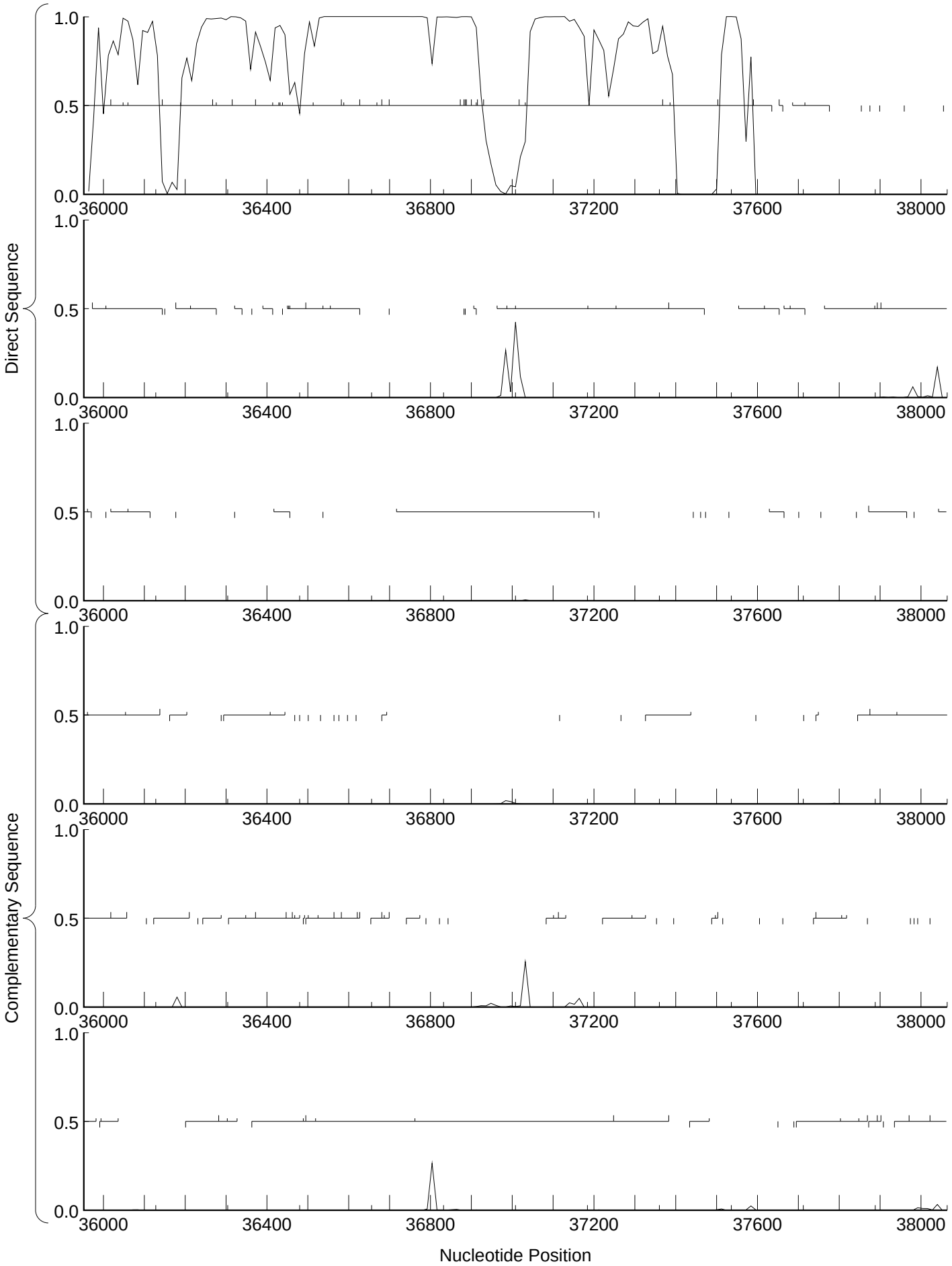

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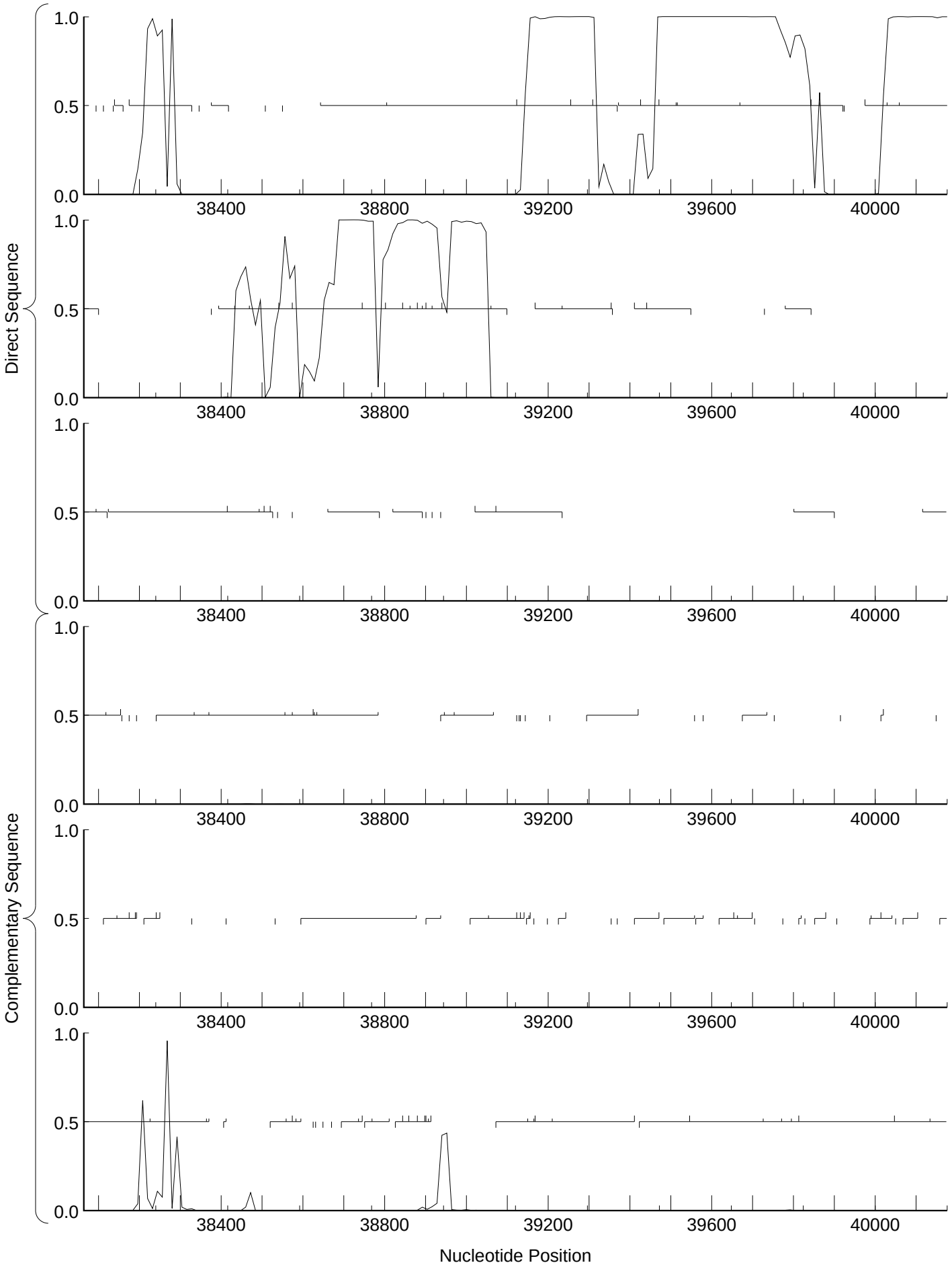

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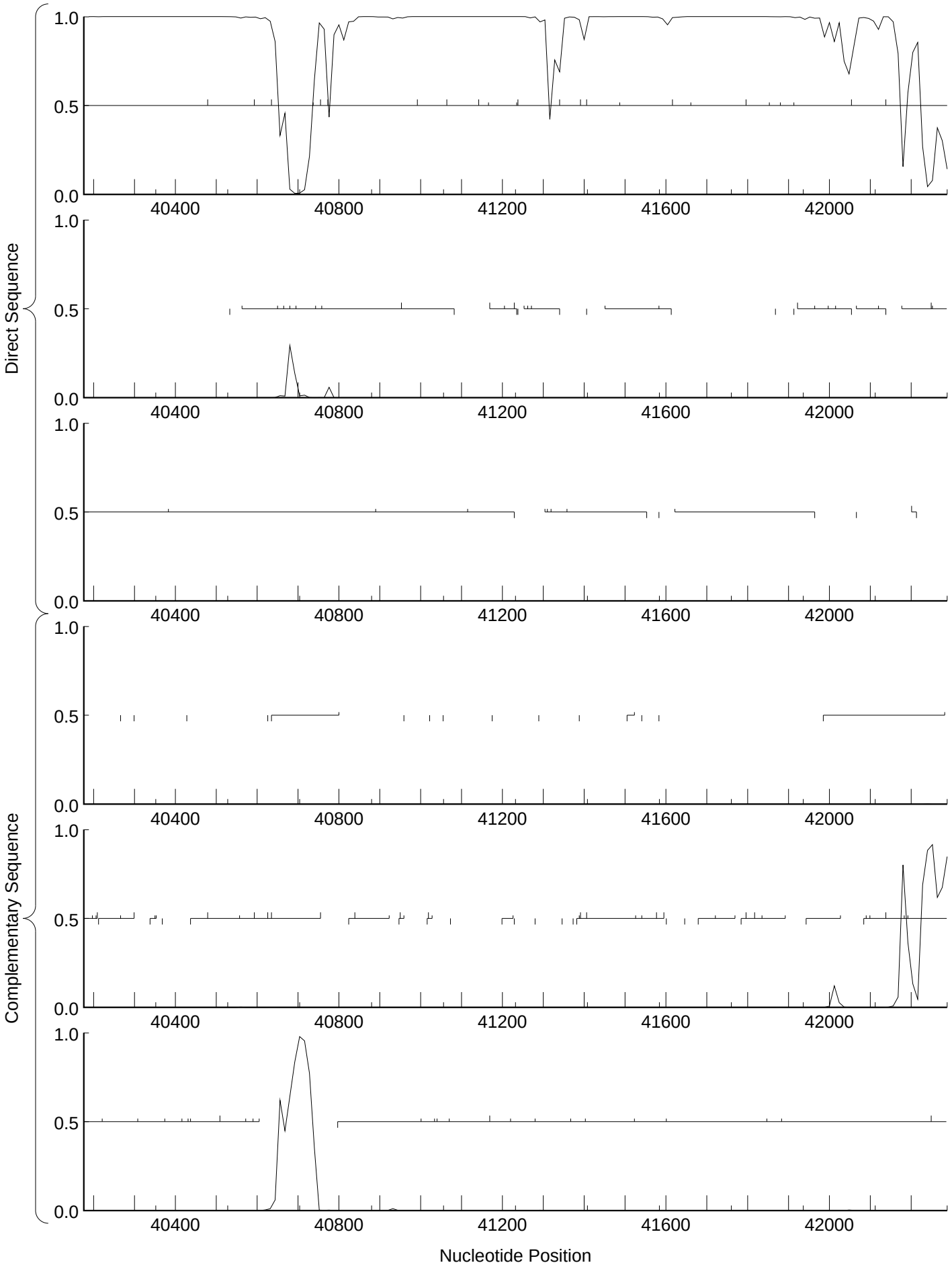

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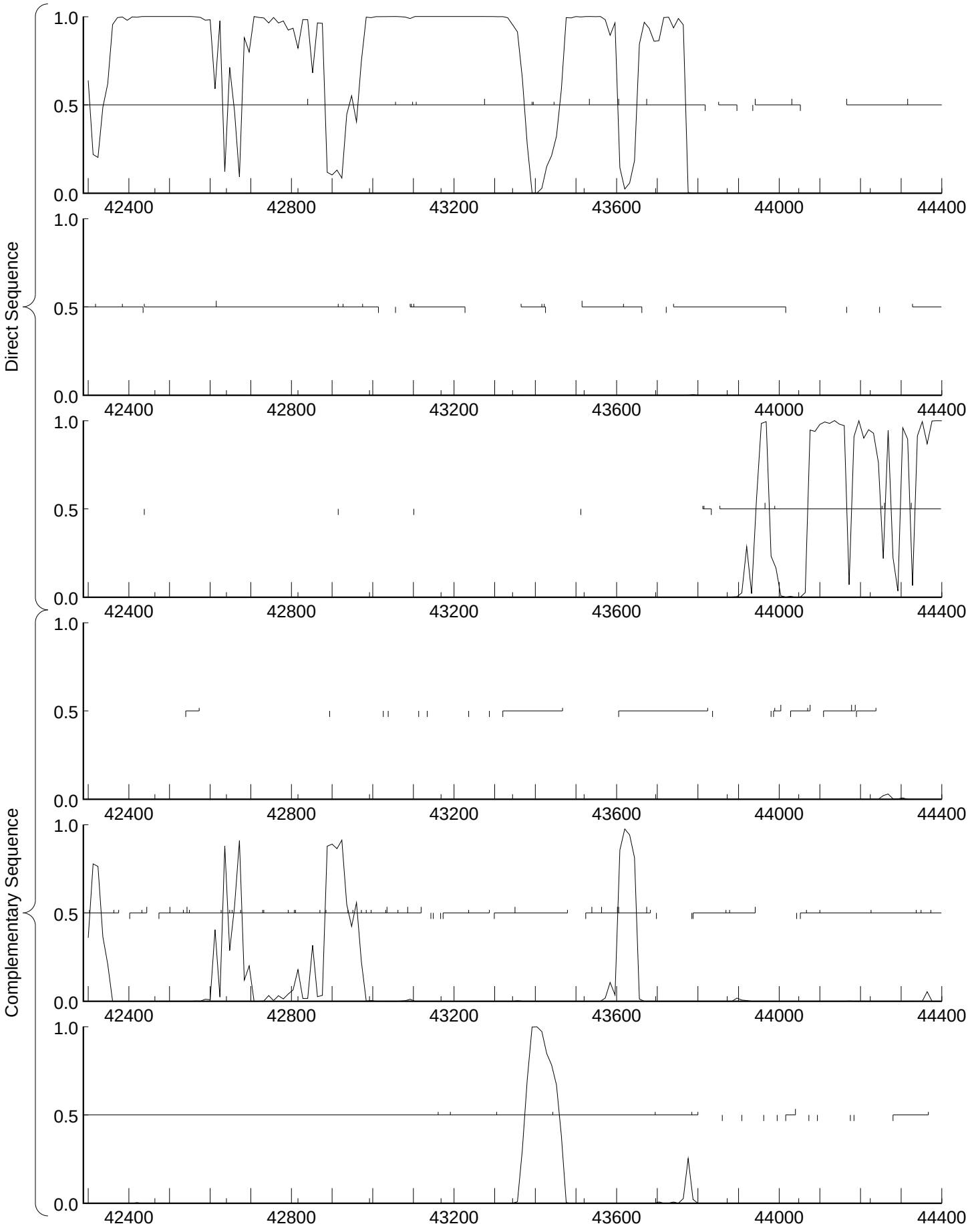

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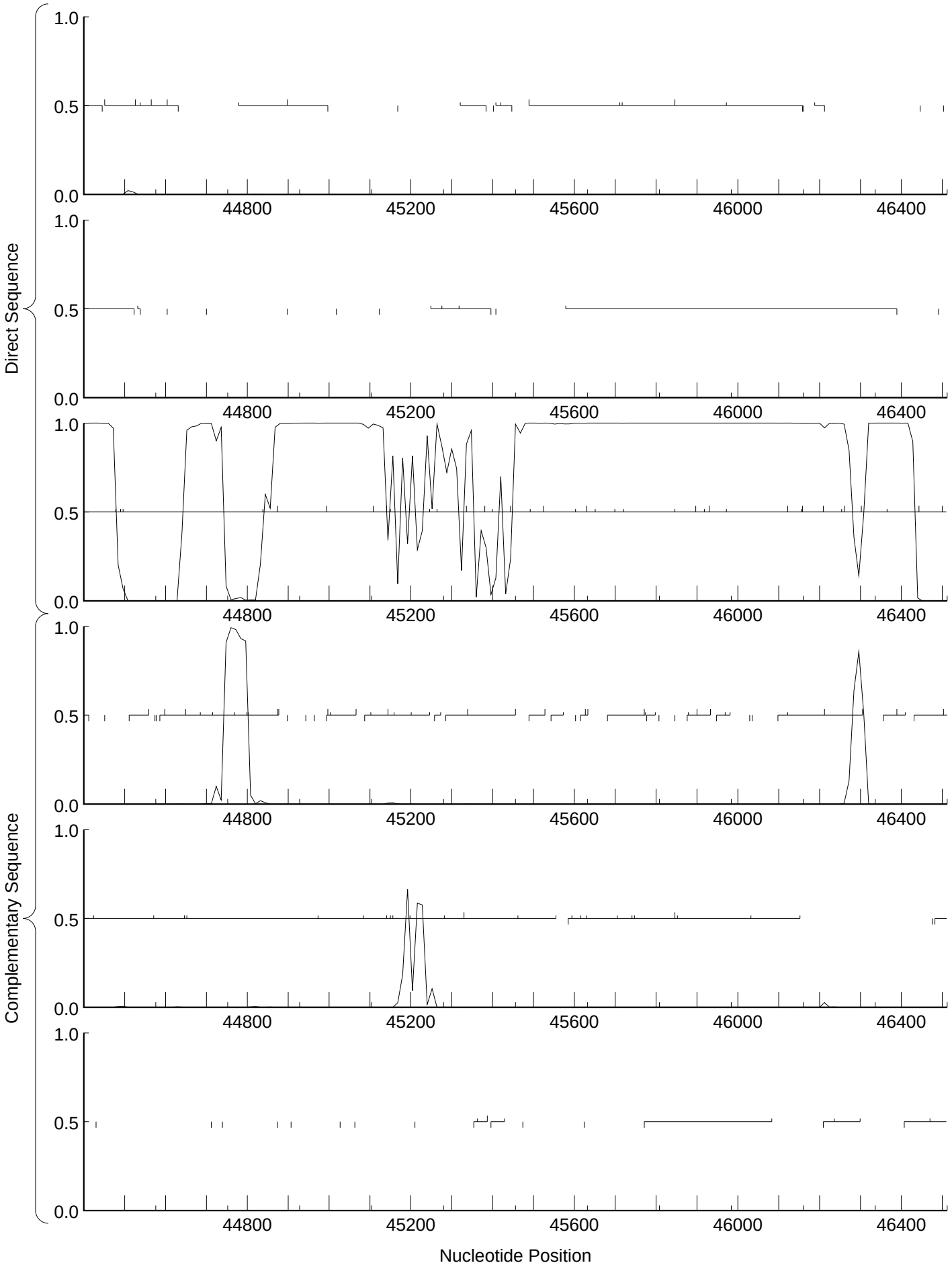

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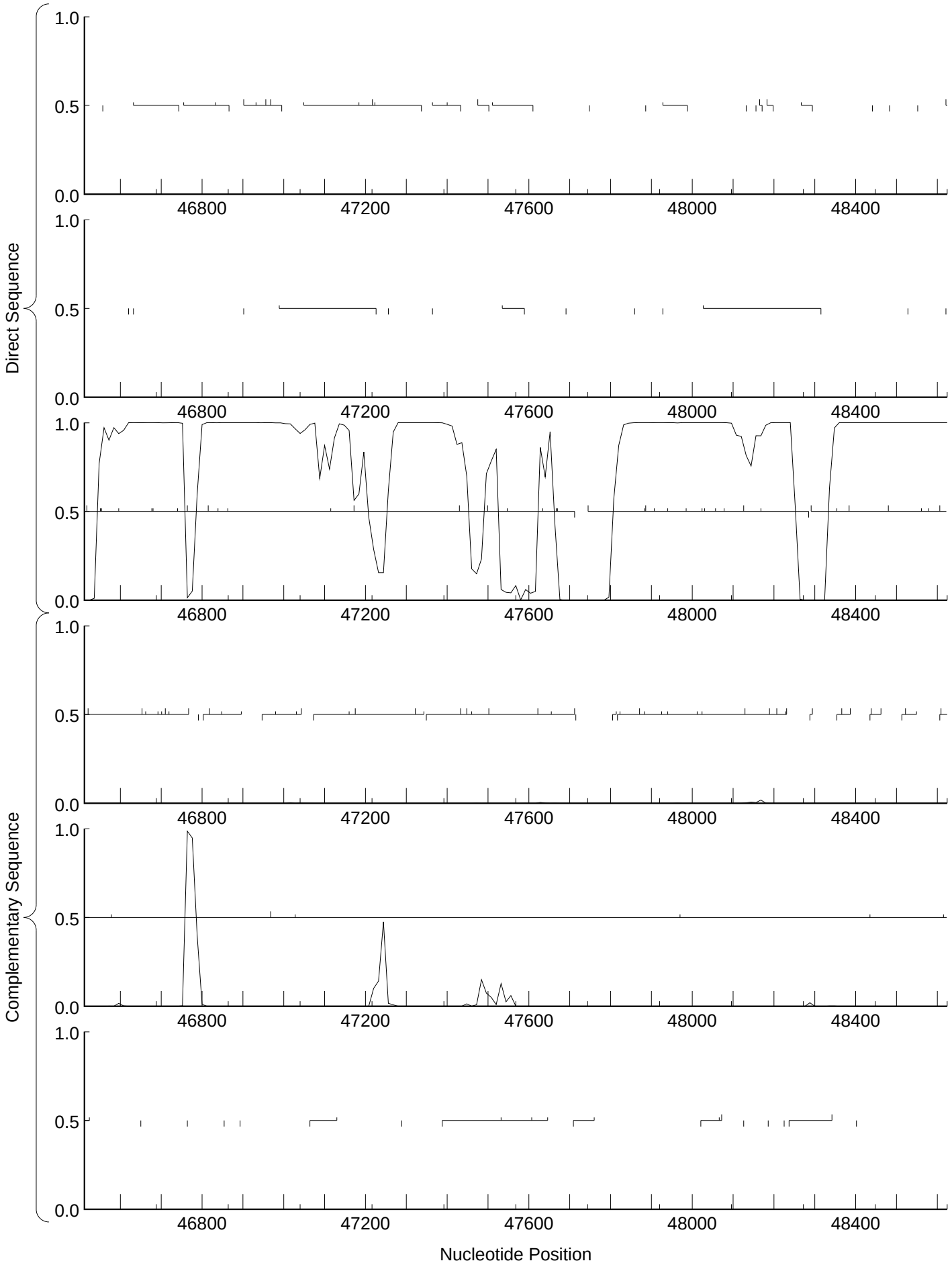

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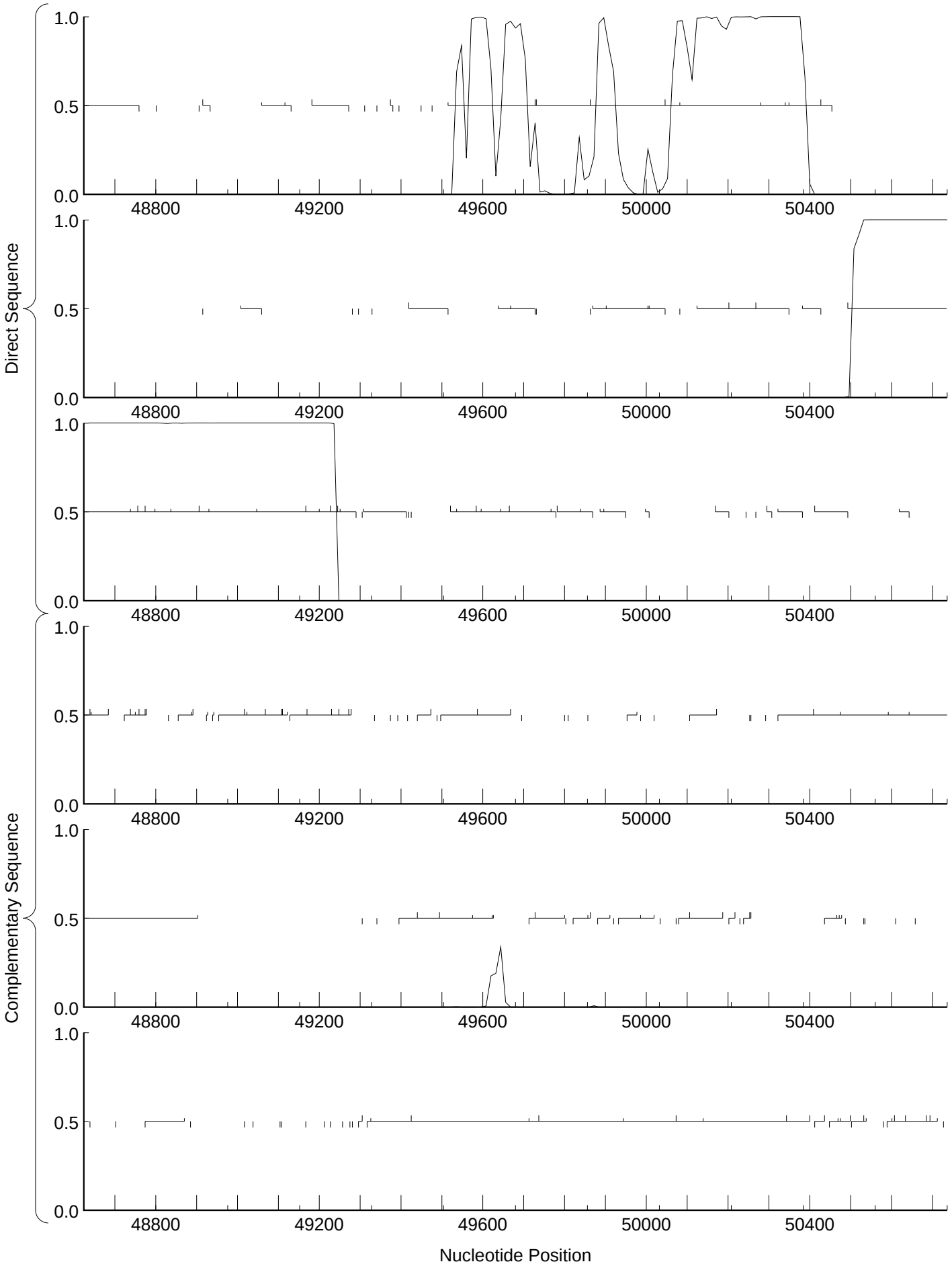

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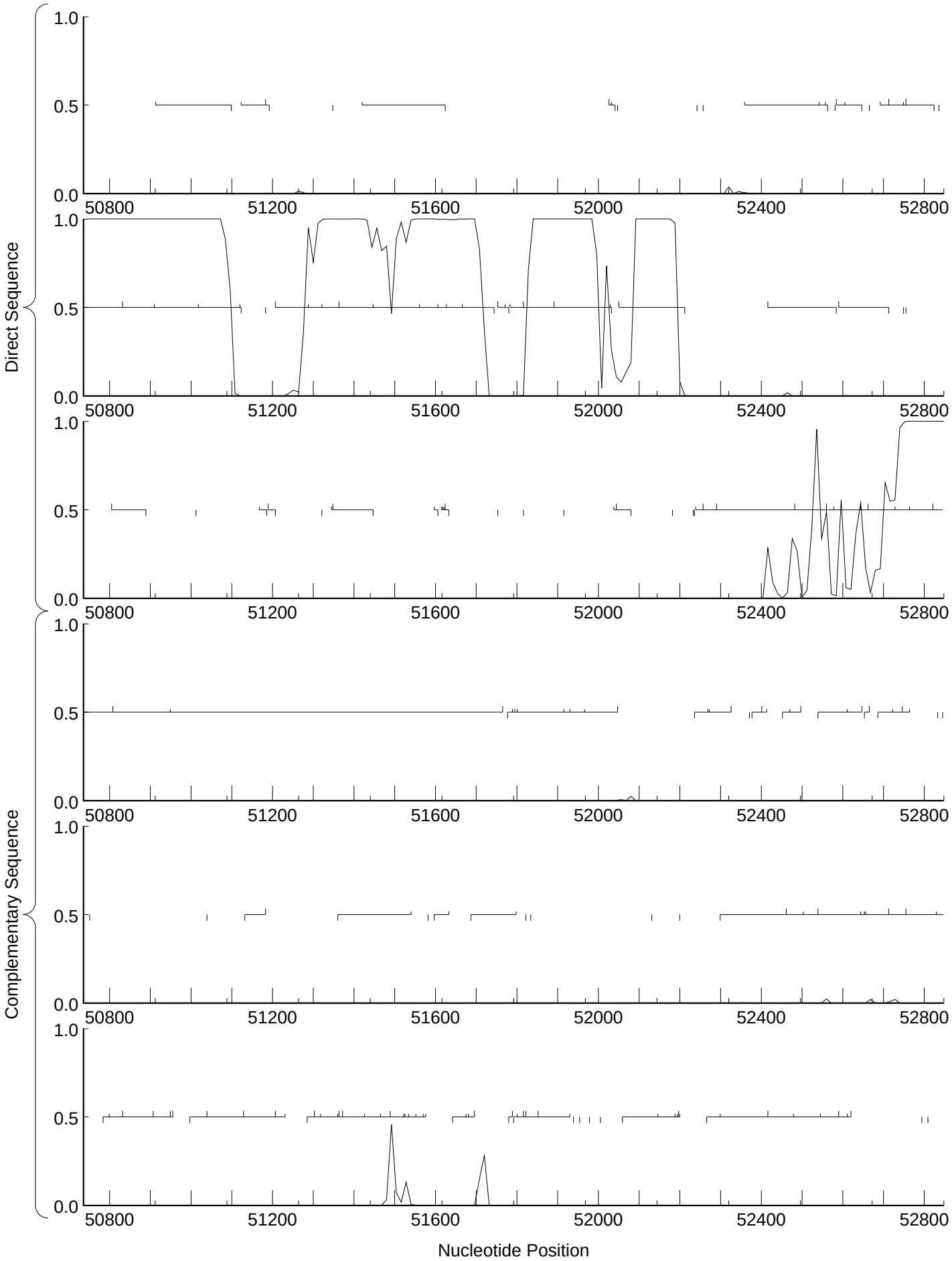

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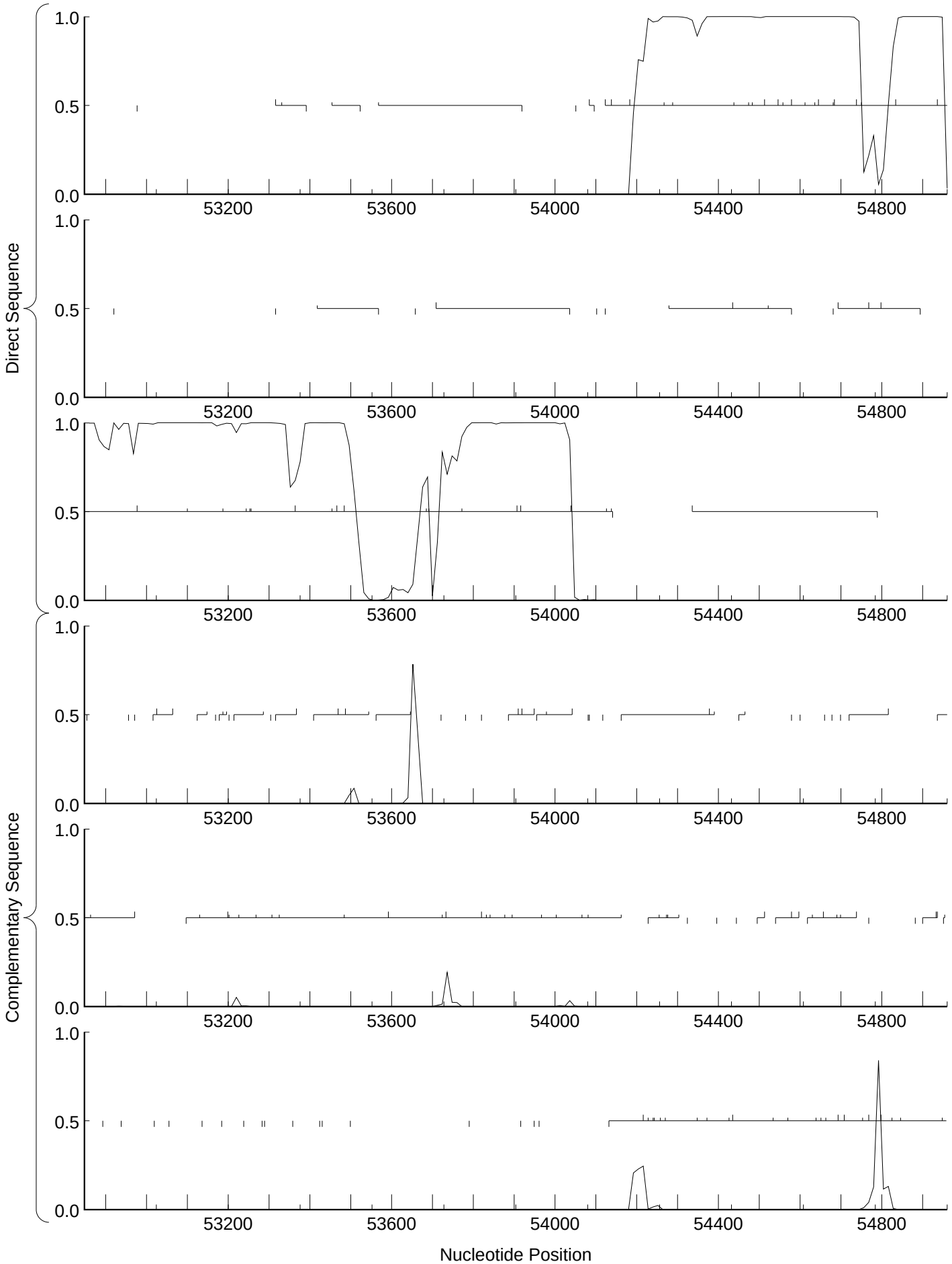

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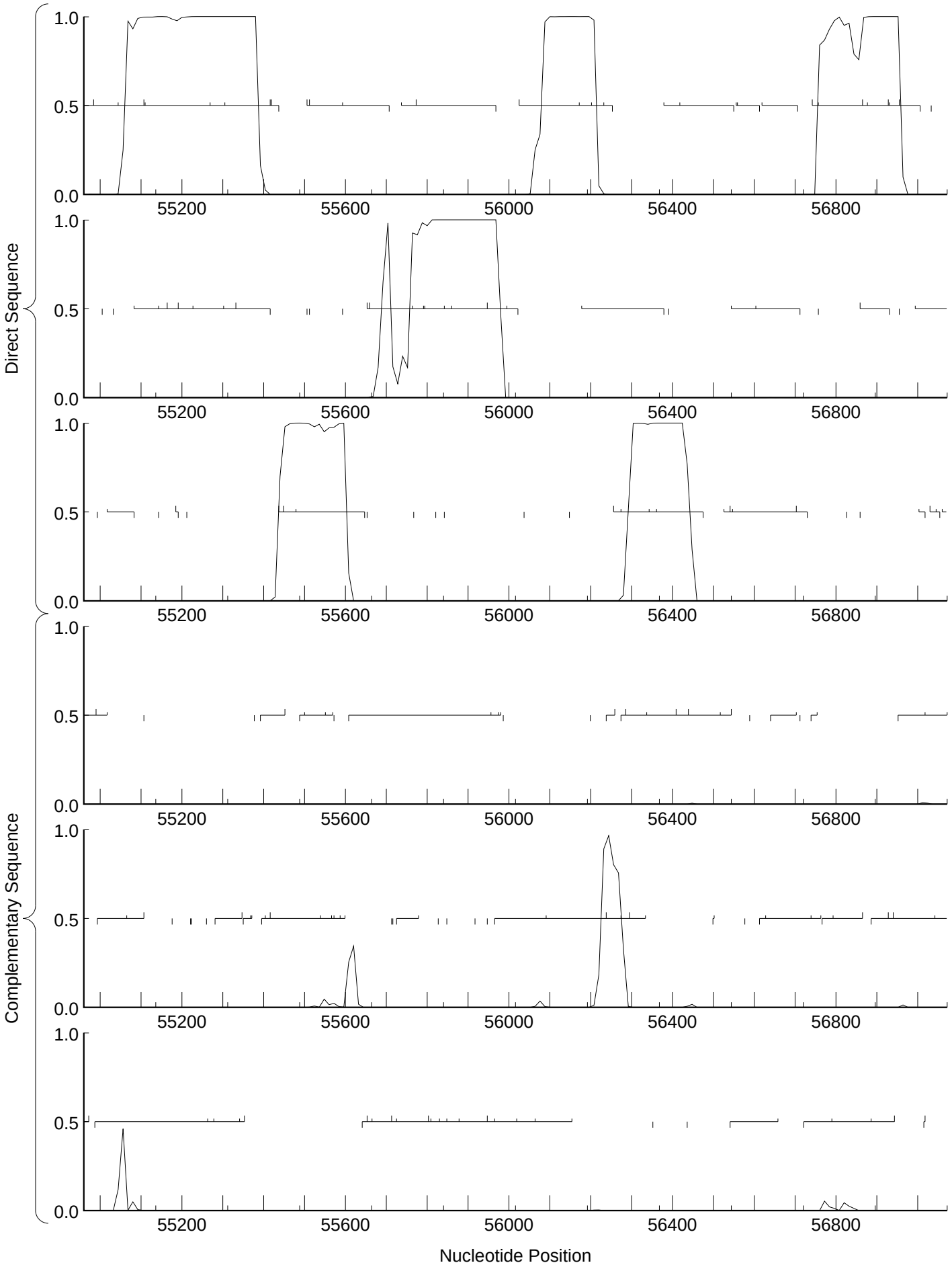

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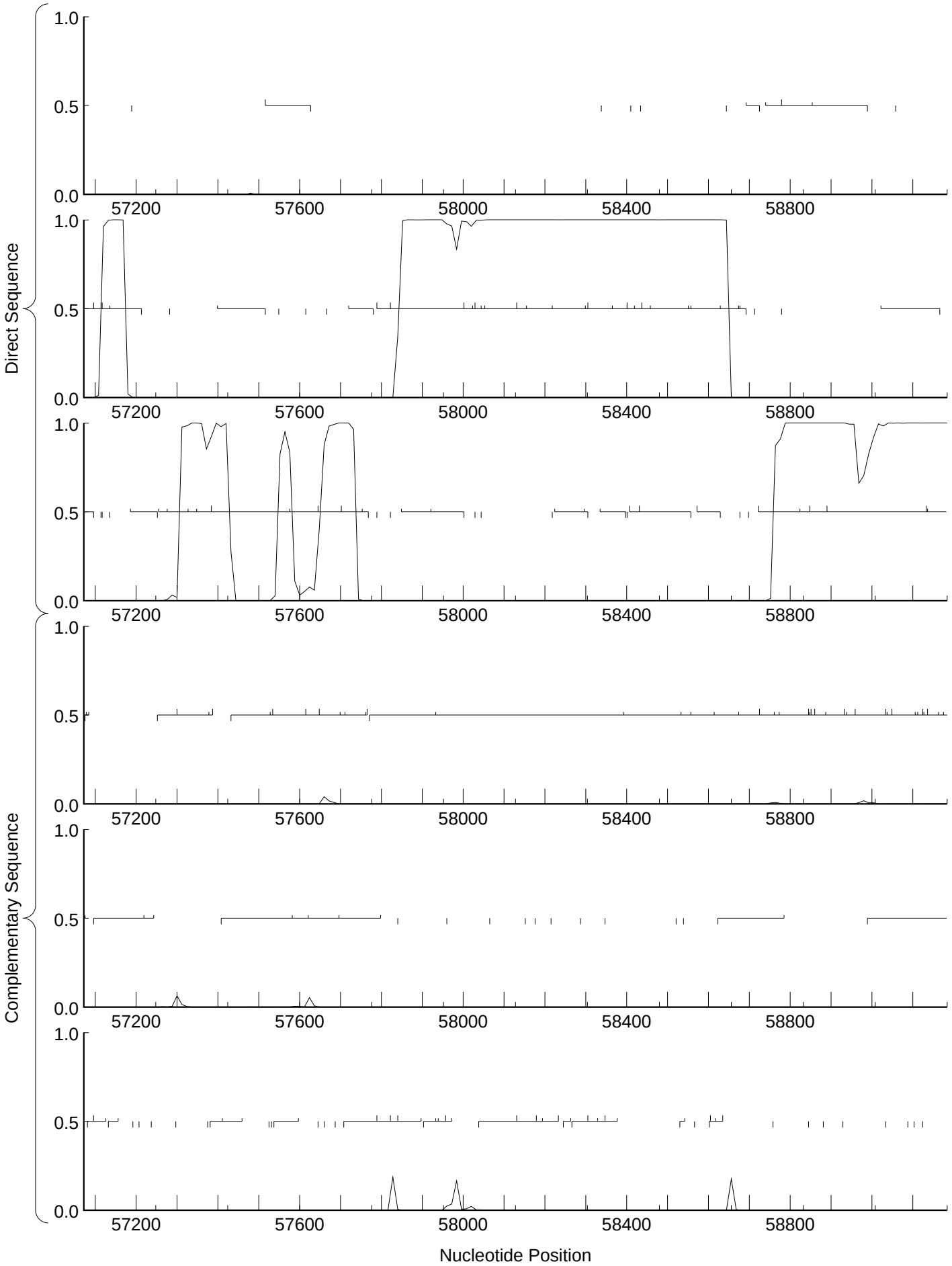

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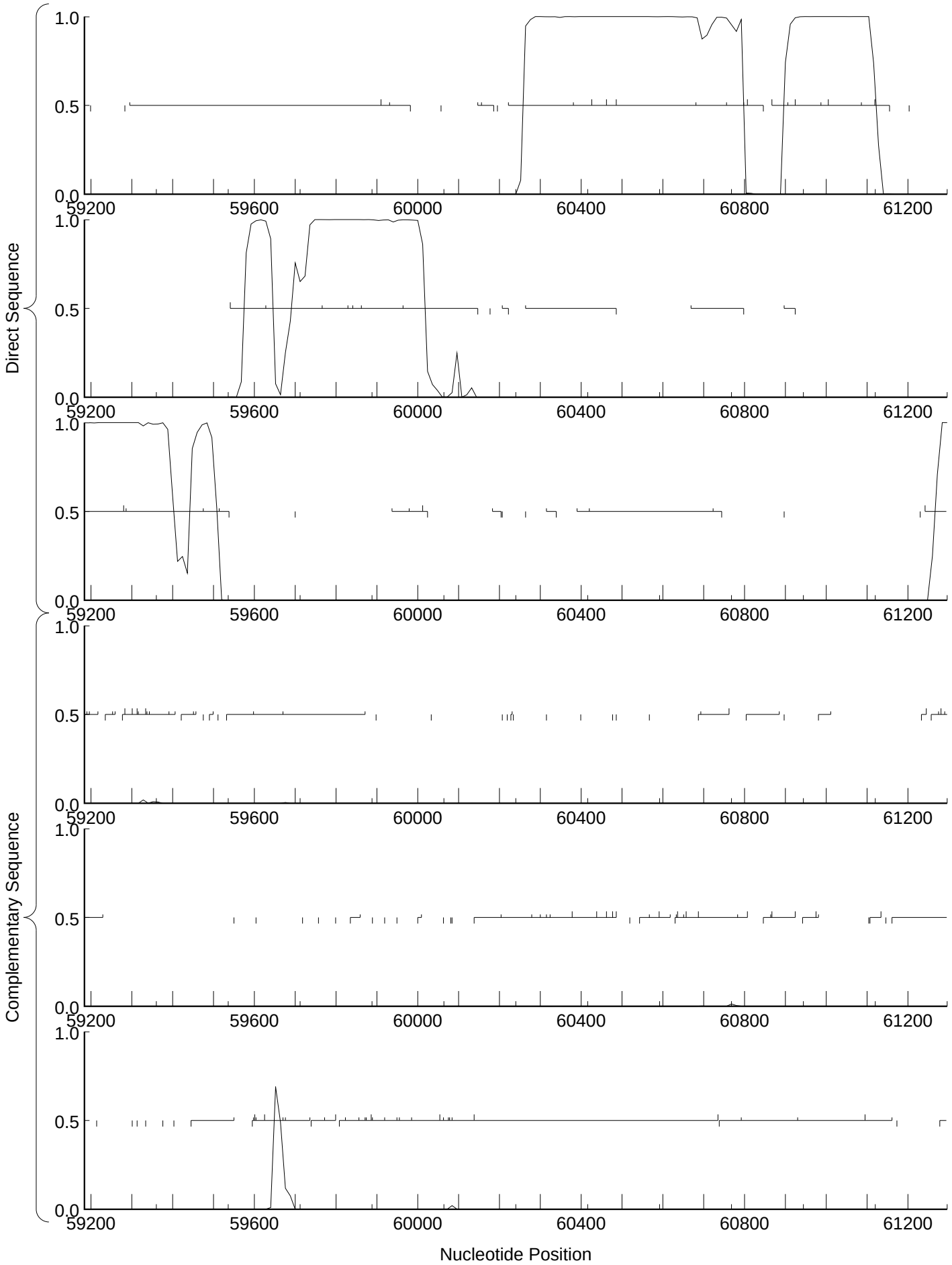

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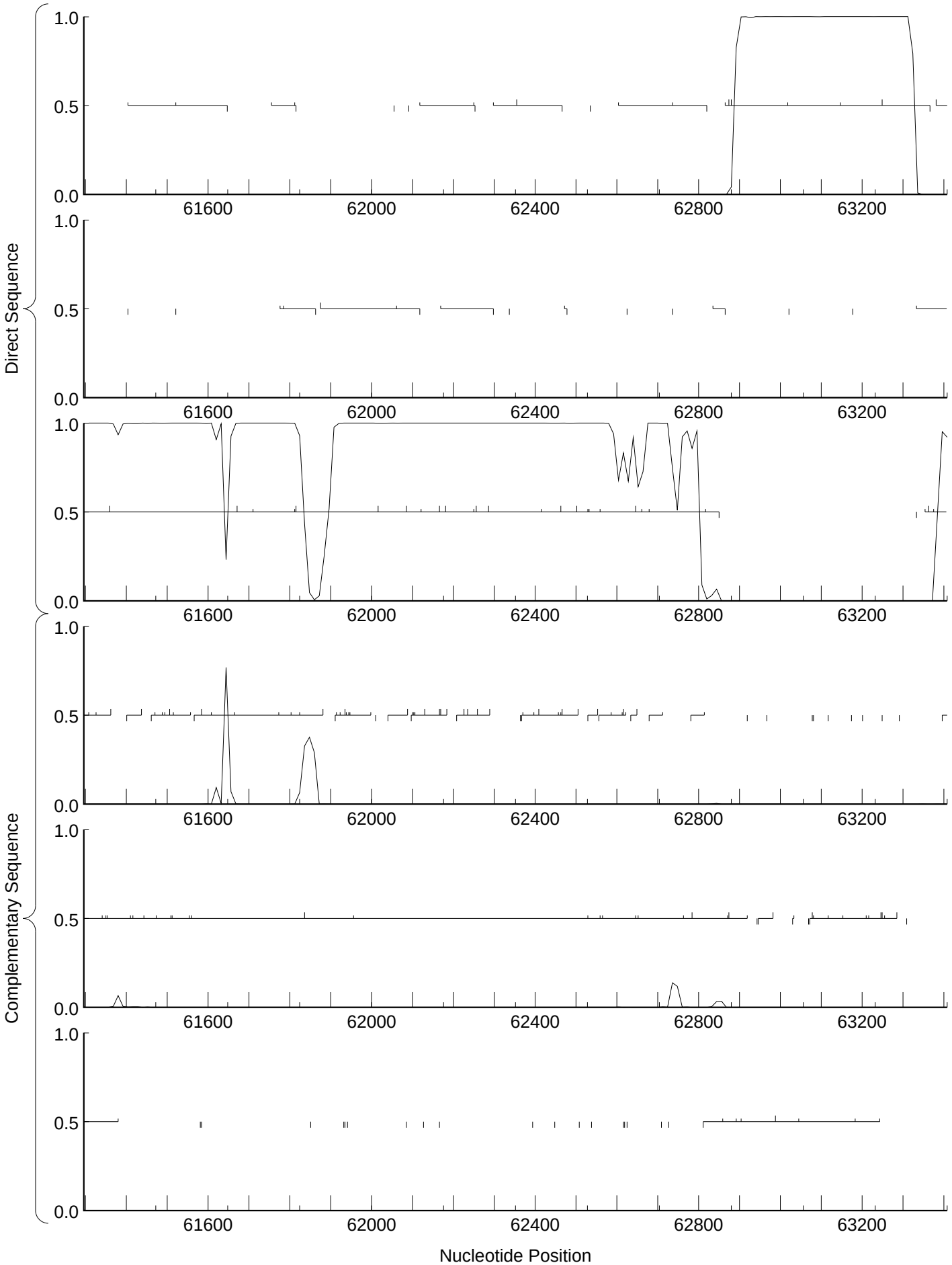

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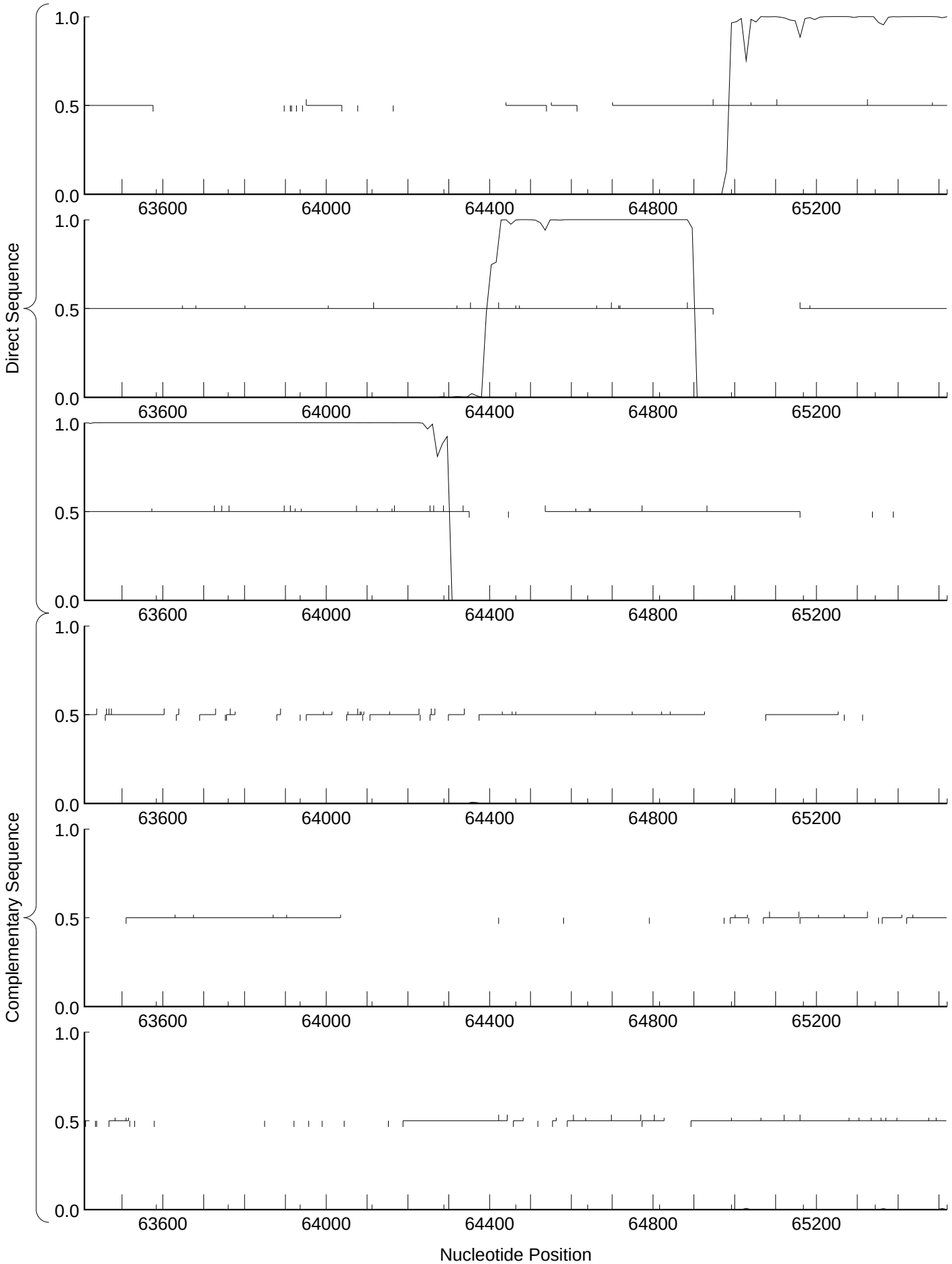

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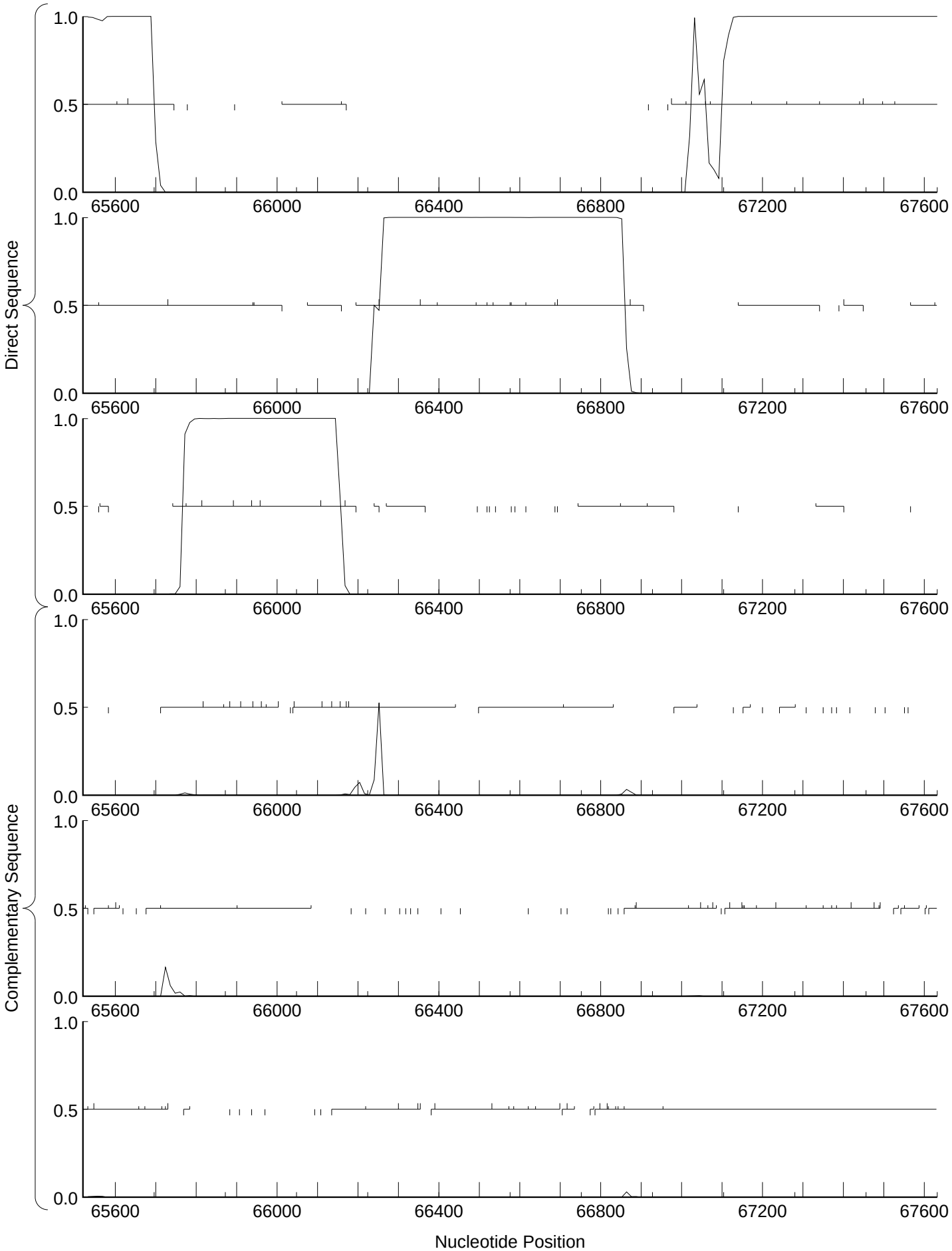

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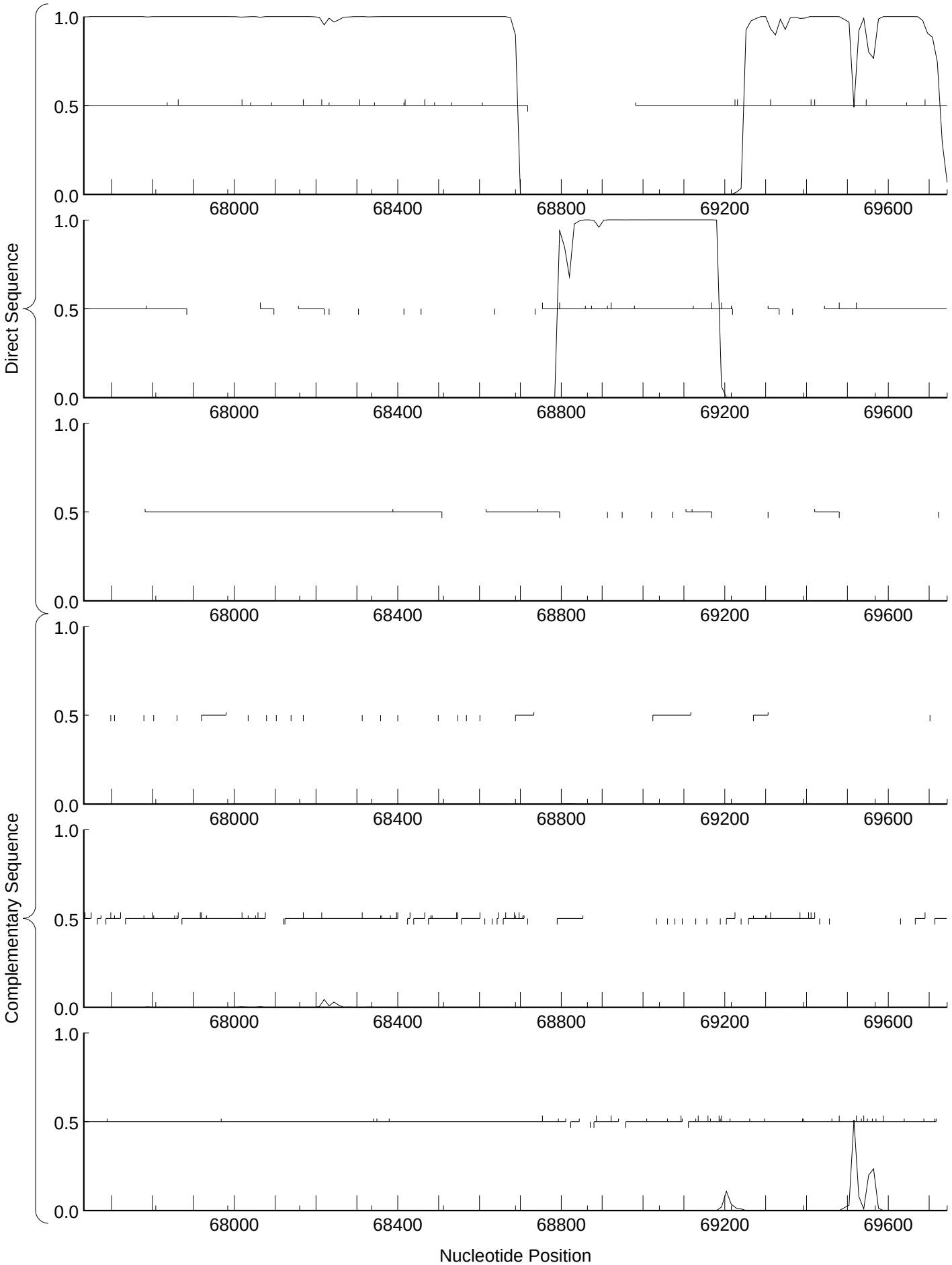

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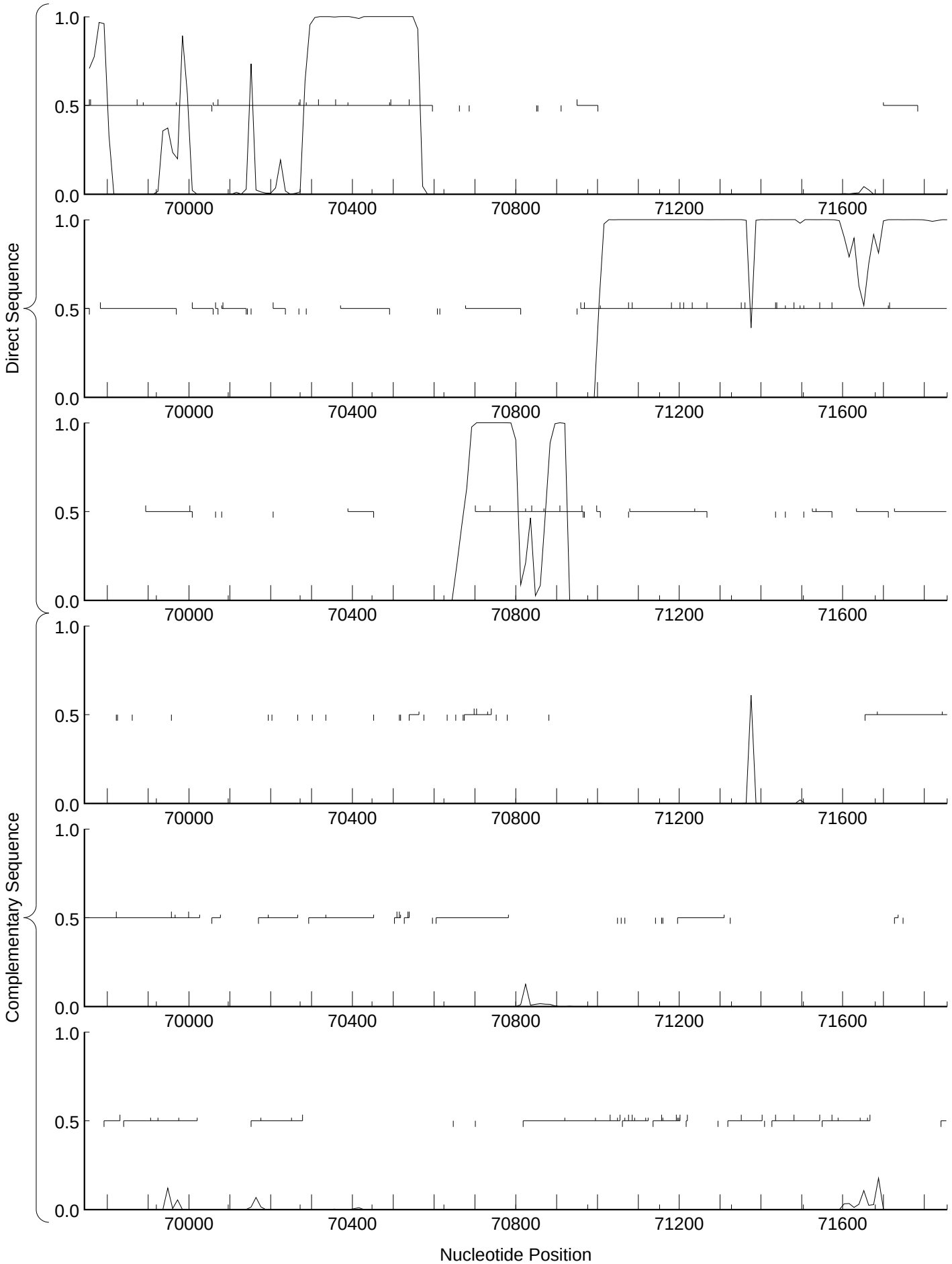

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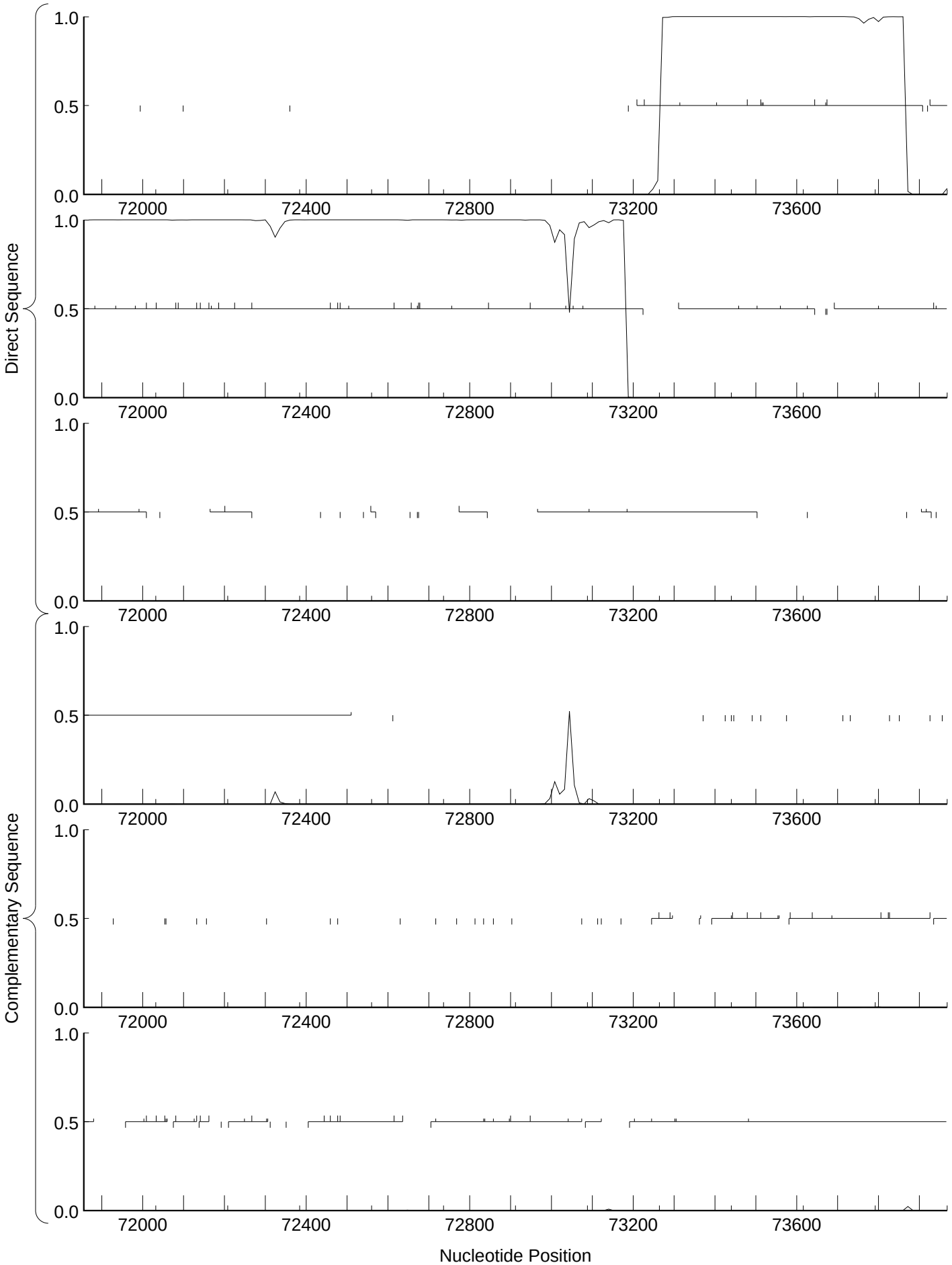

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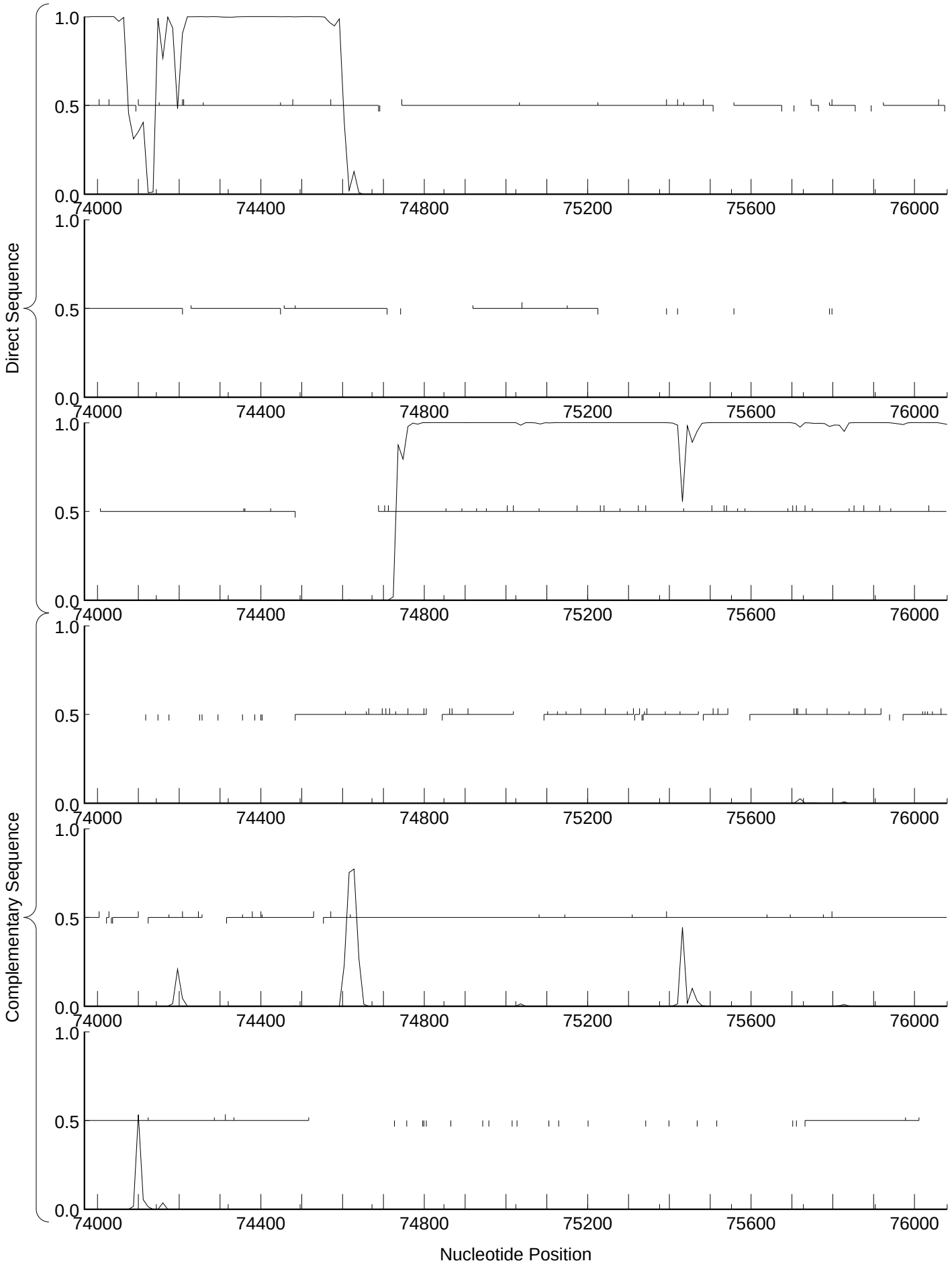

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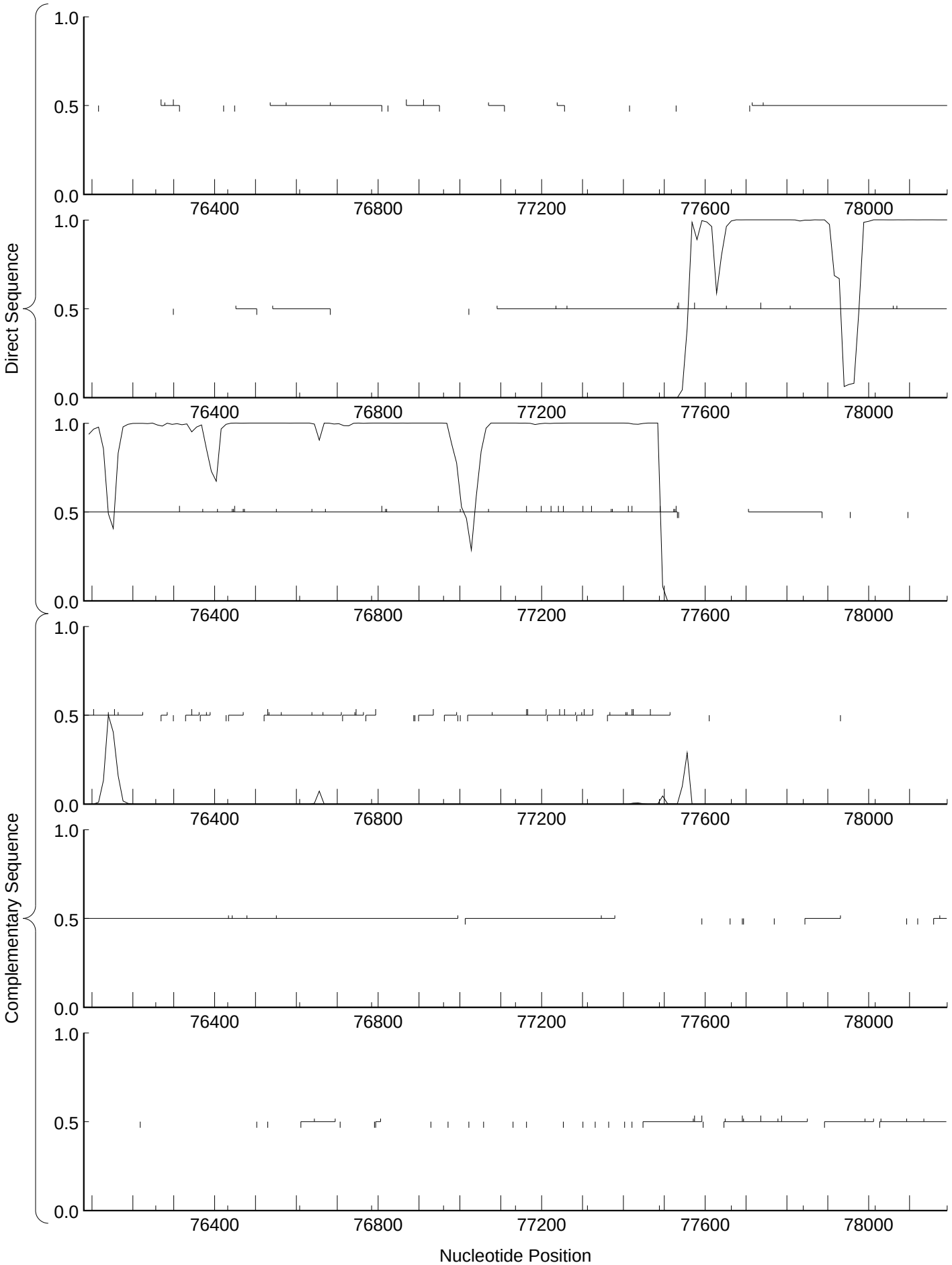

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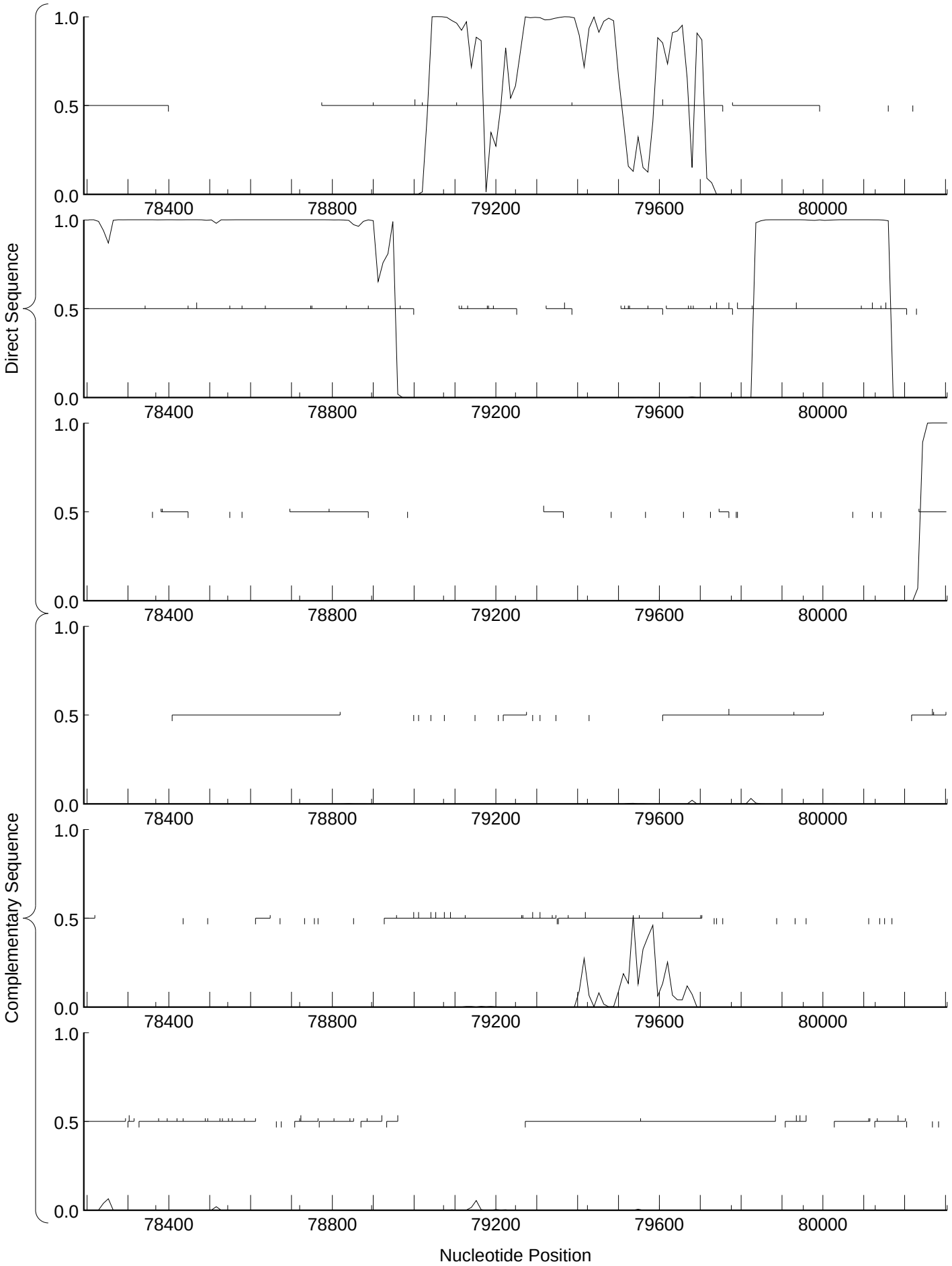

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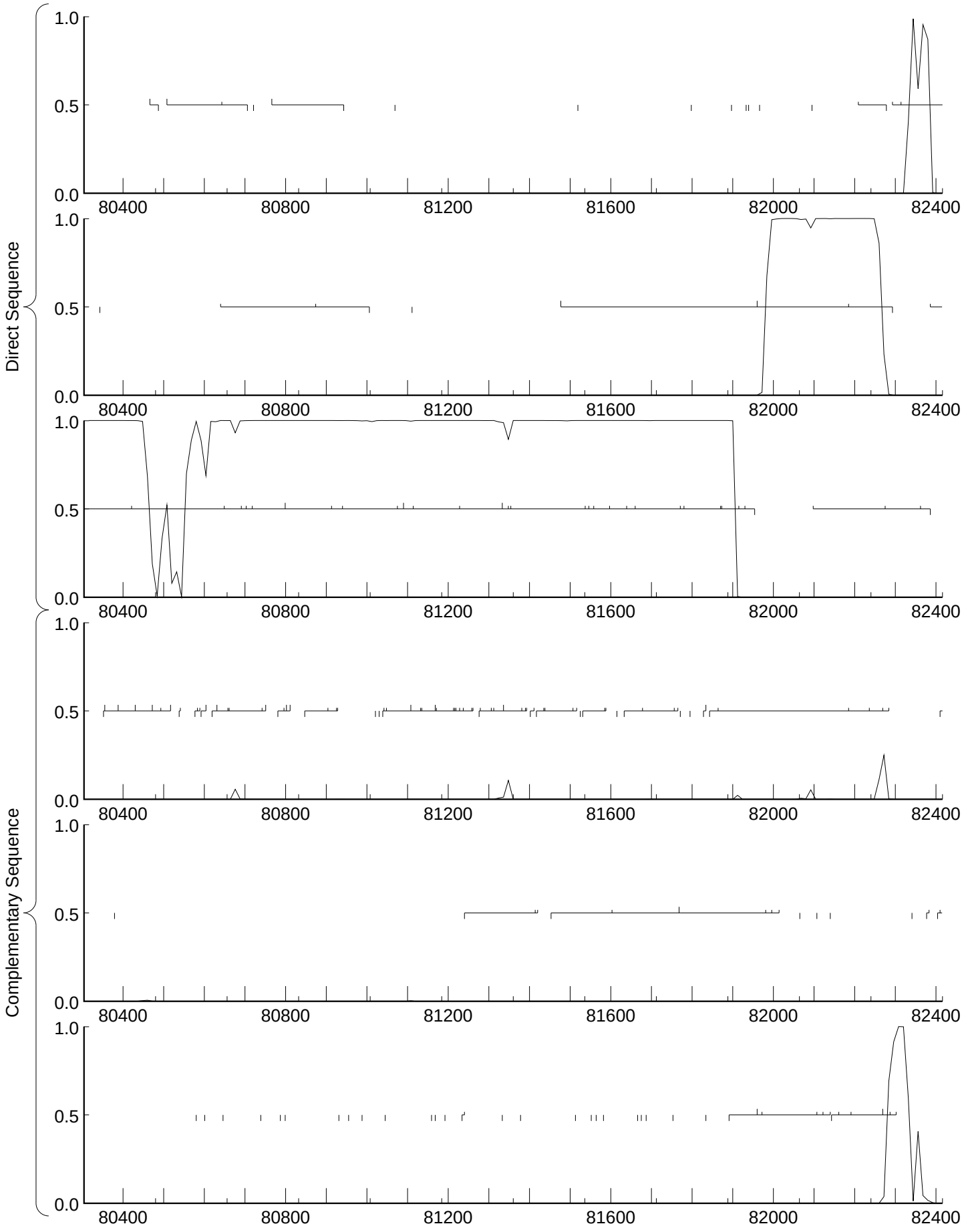

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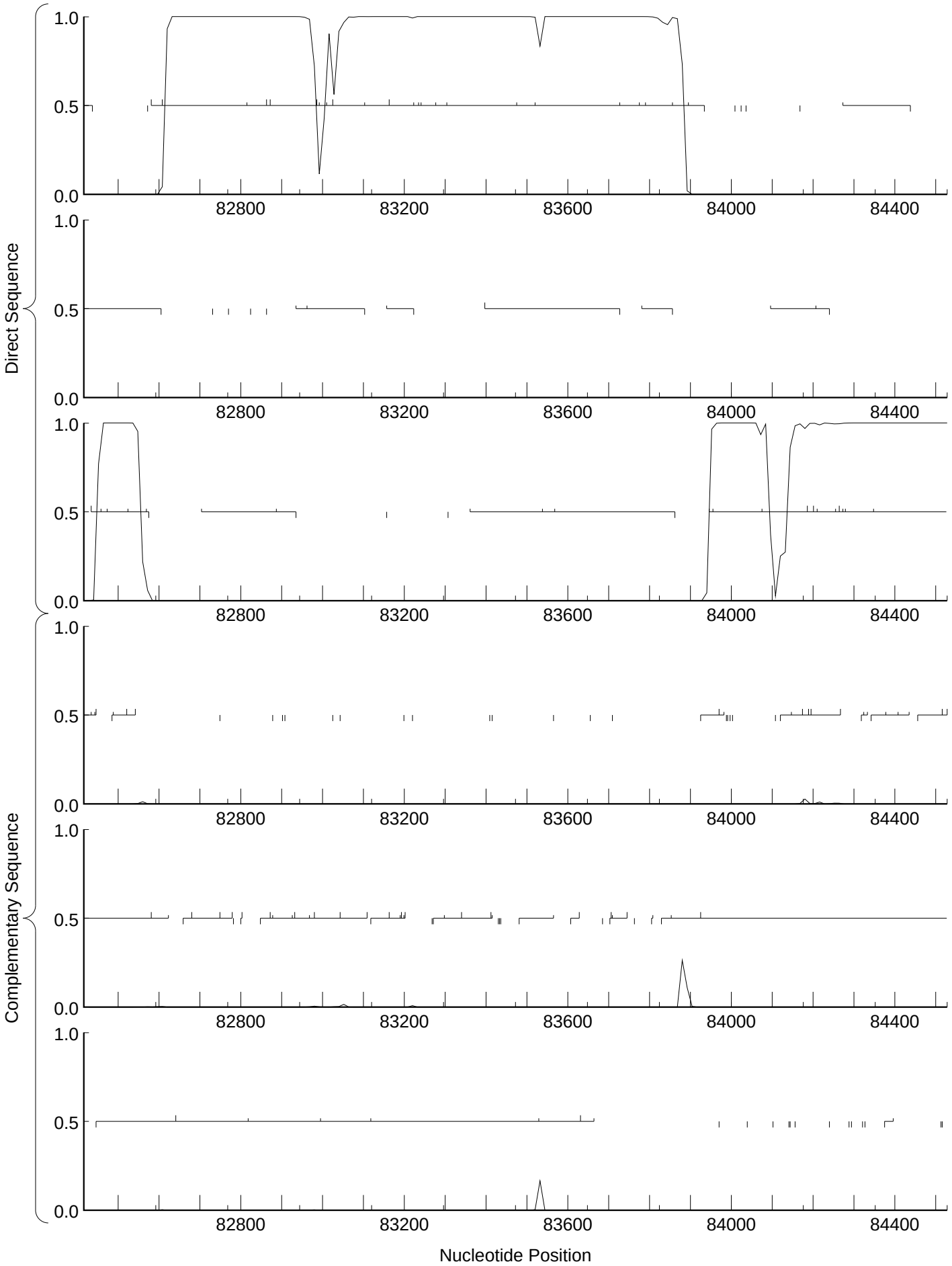

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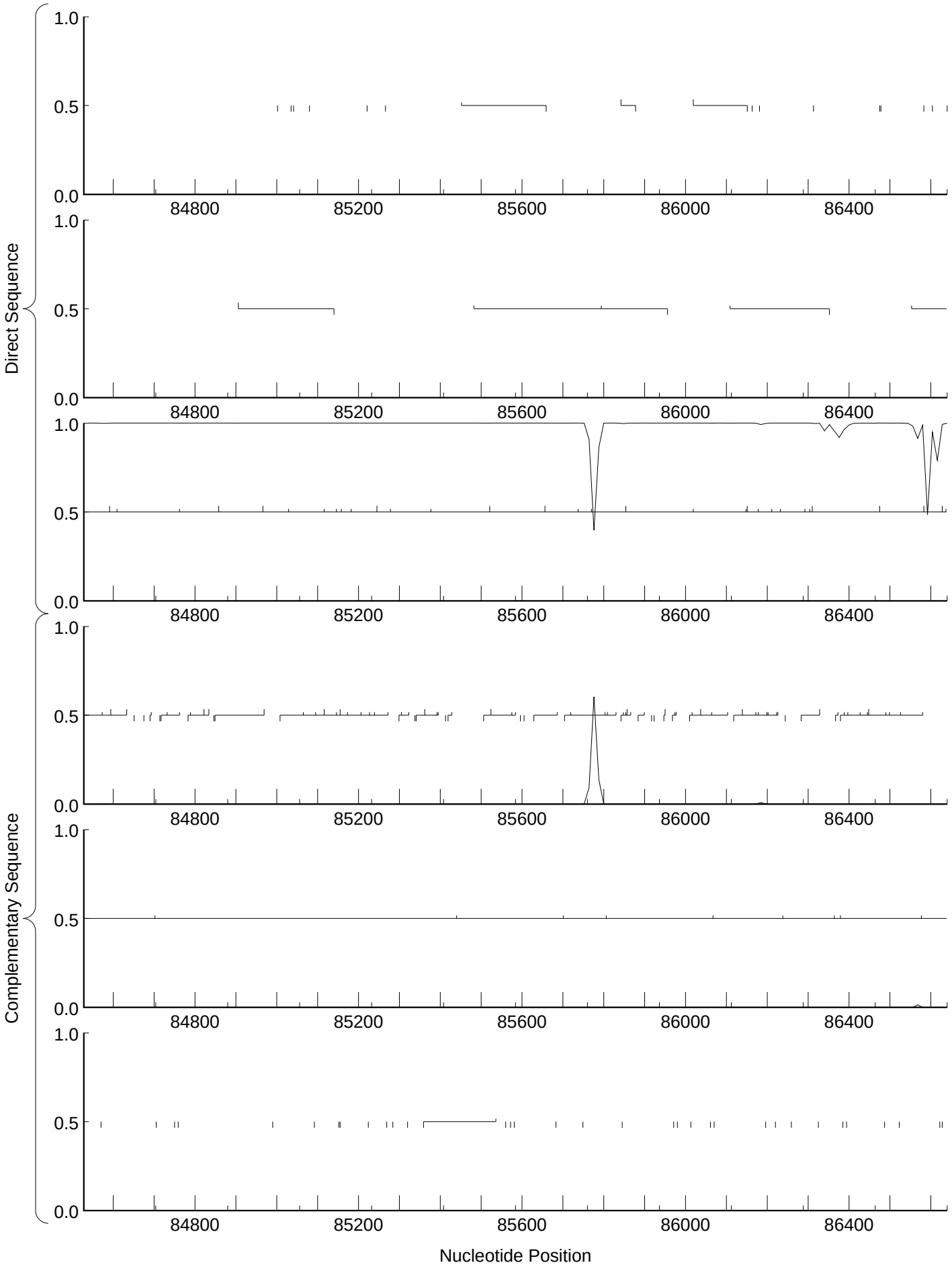

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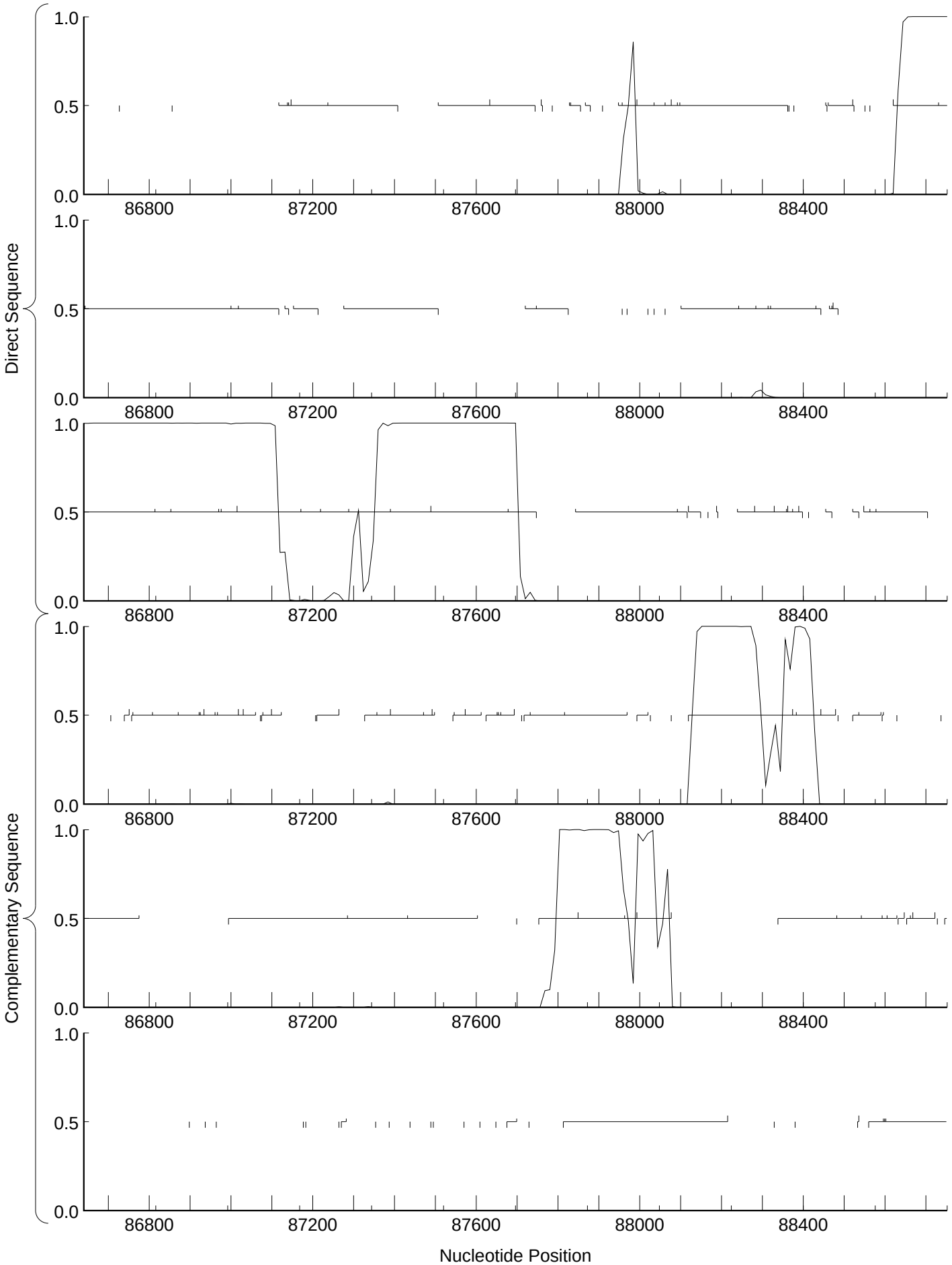

Direct Sequence
Complementary Sequence
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Nucleotide Position

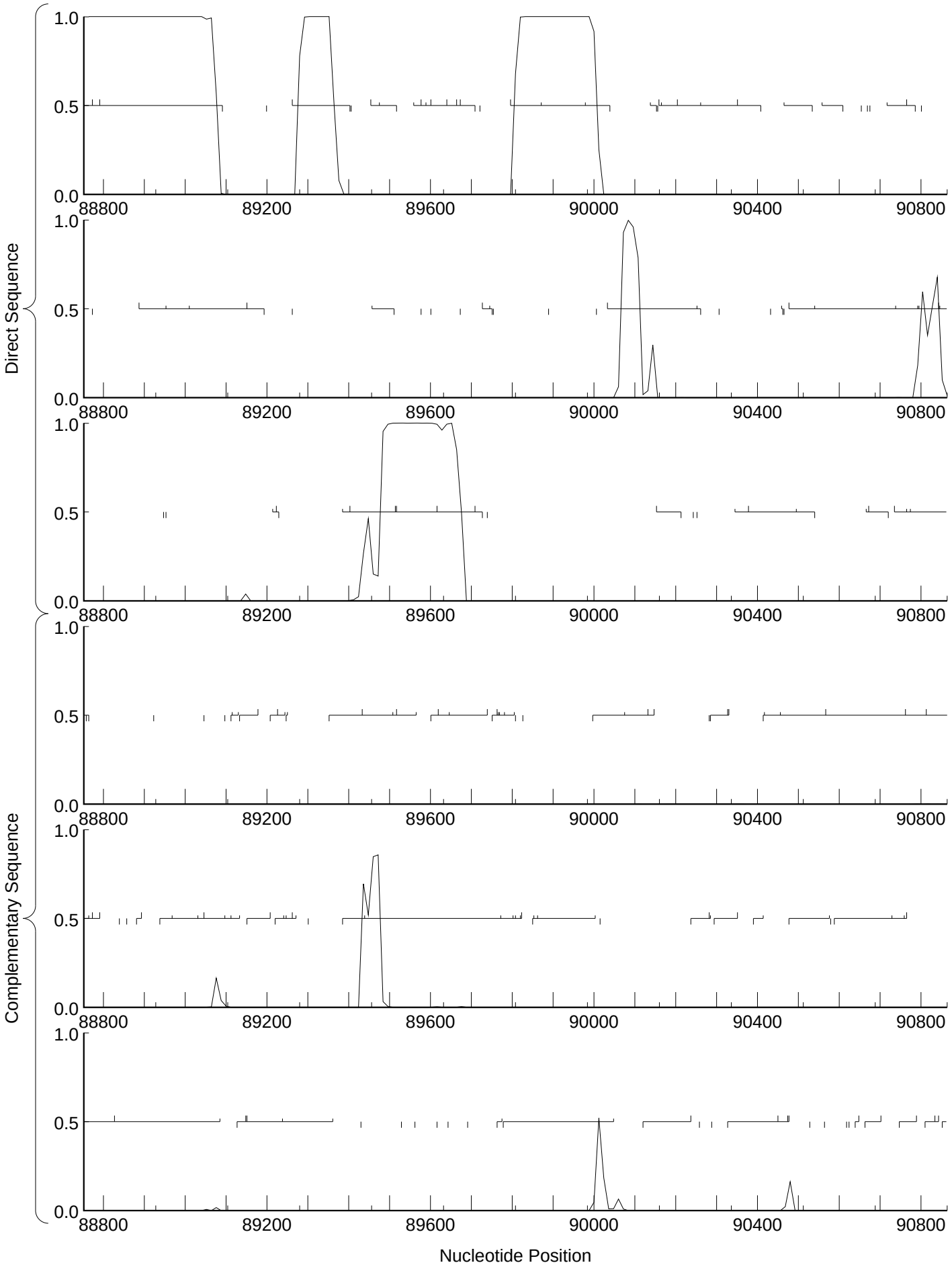

Direct Sequence
Complementary Sequence
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0.5
0.0
78400
78800
79200
79600
80000
Nucleotide Position

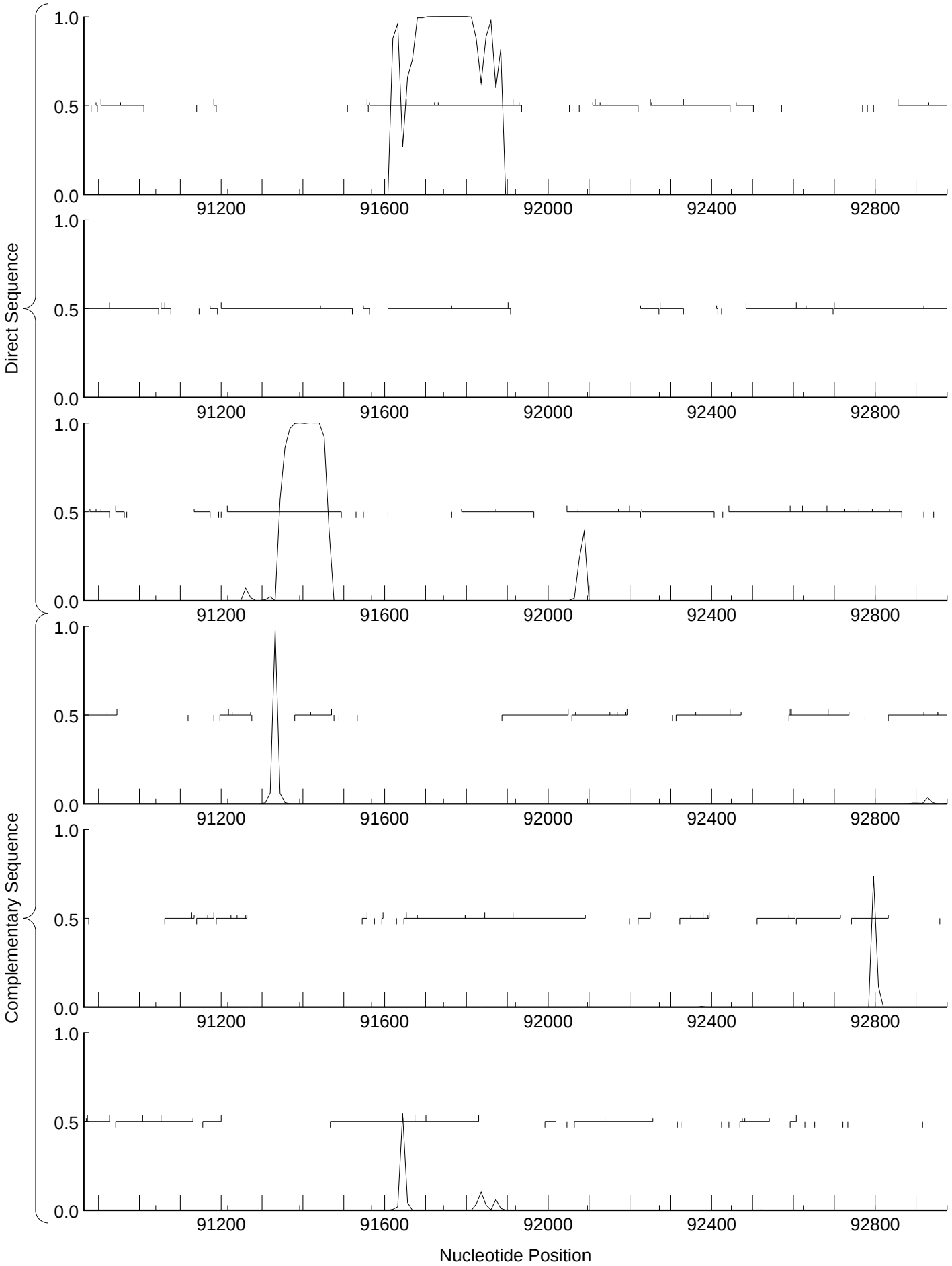

Direct Sequence
Complementary Sequence
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0.0
80400
80800
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82000
82400
Nucleotide Position

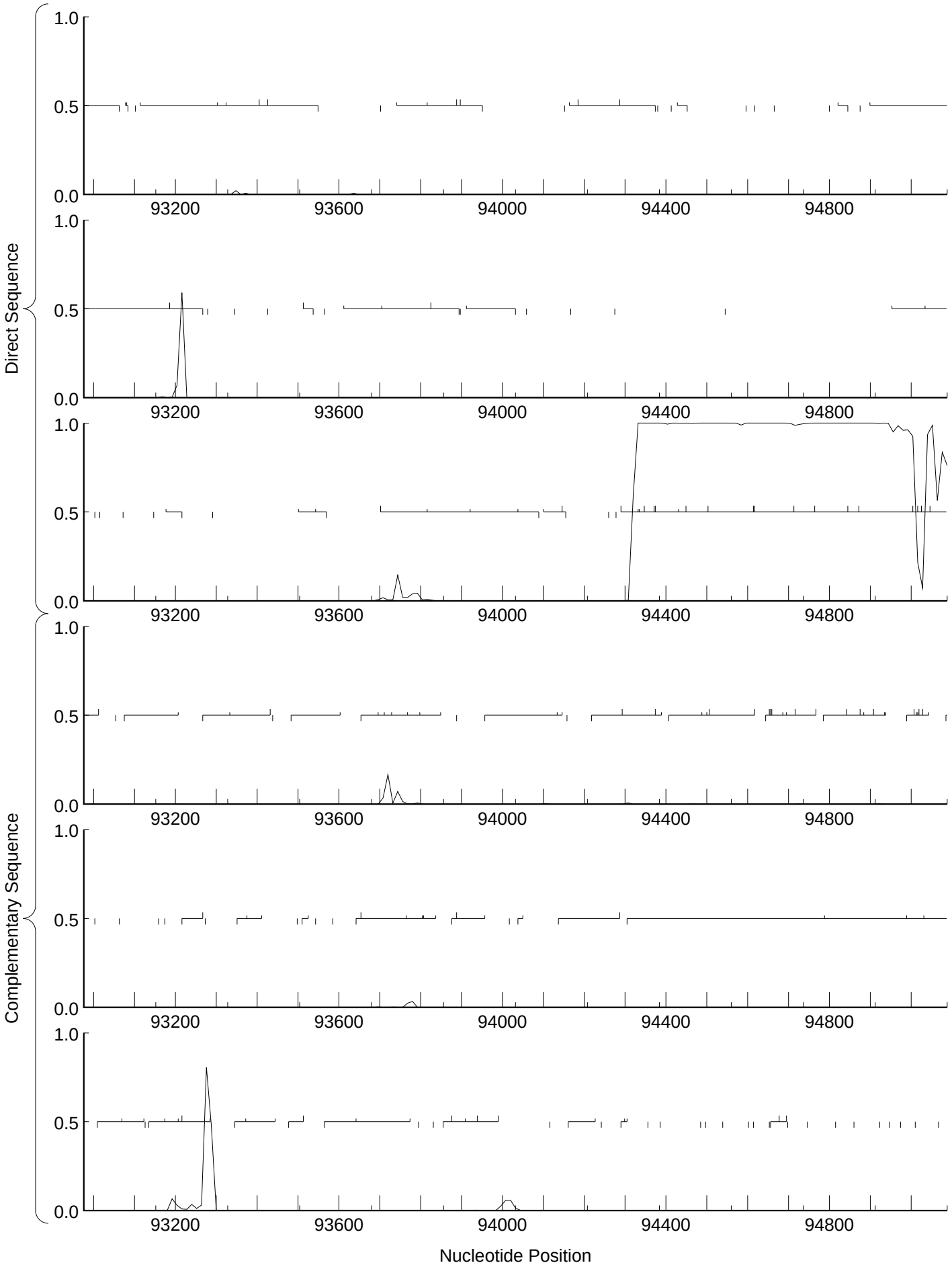

Direct Sequence
Complementary Sequence
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82800
83200
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84000
84400
Nucleotide Position

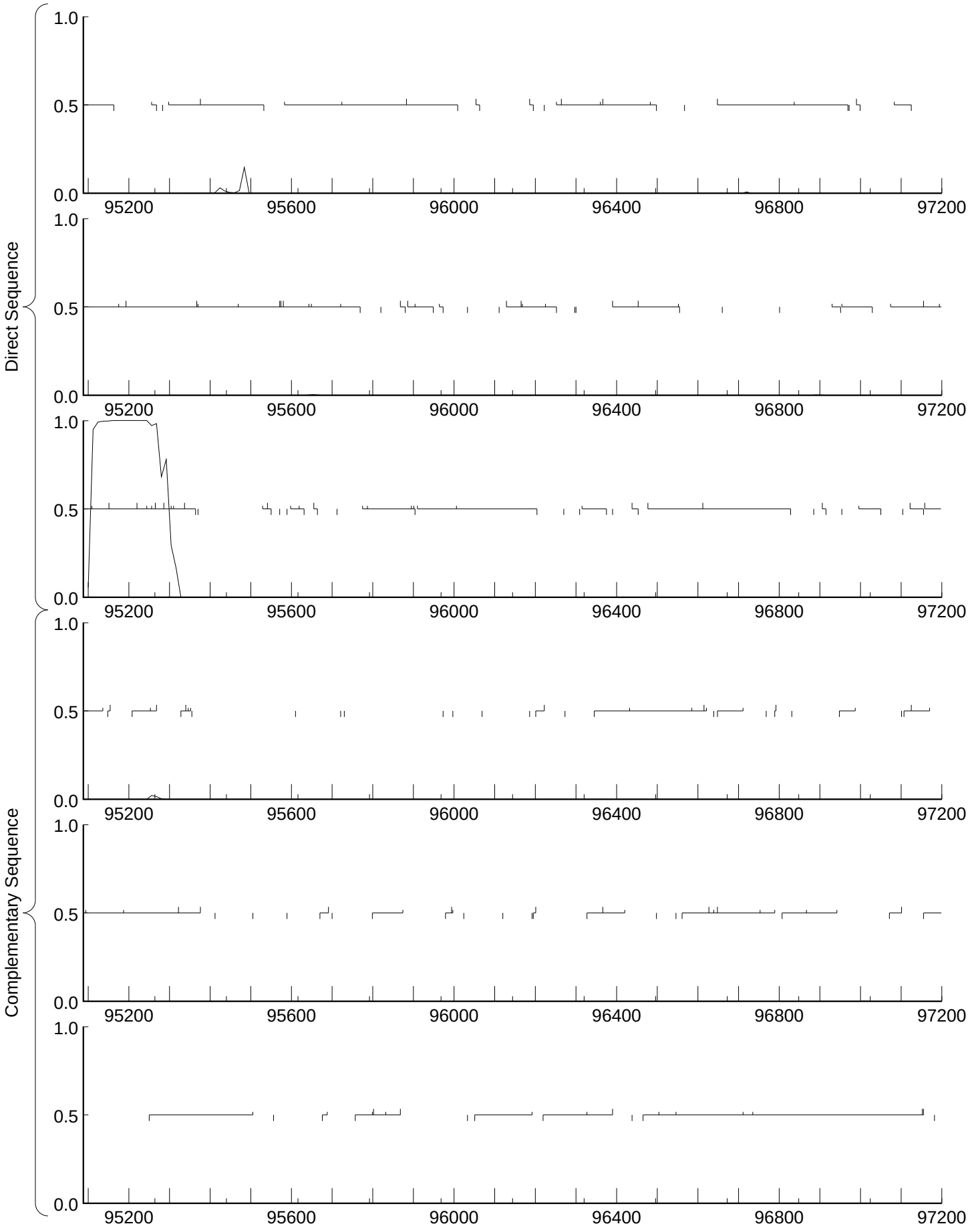

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Complementary Sequence
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85600
86000
86400
Nucleotide Position

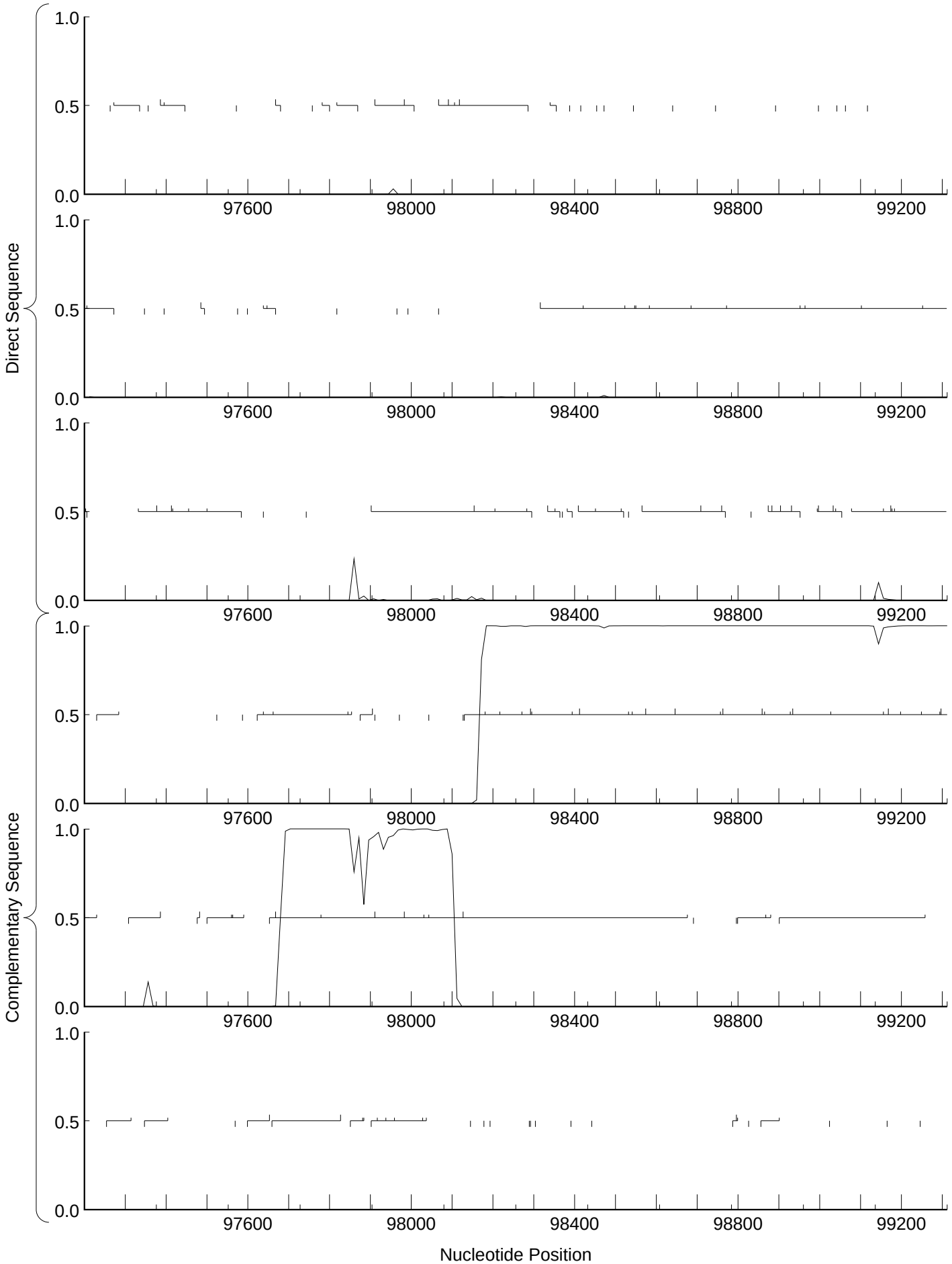

Direct Sequence
Complementary Sequence
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0.0
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87200
87600
88000
88400
Nucleotide Position

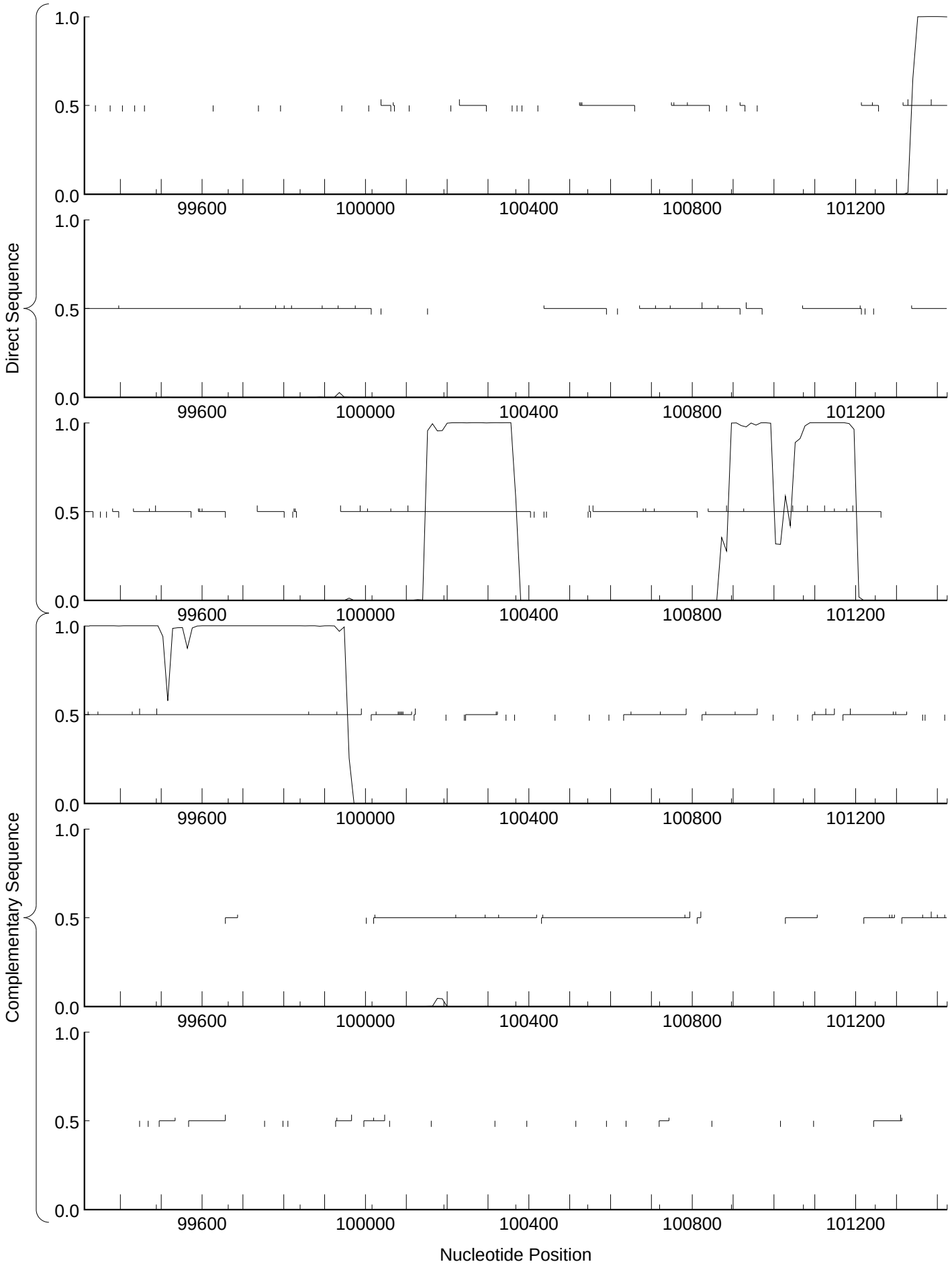

Direct Sequence
Complementary Sequence
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89600
90000
90400
90800
Nucleotide Position

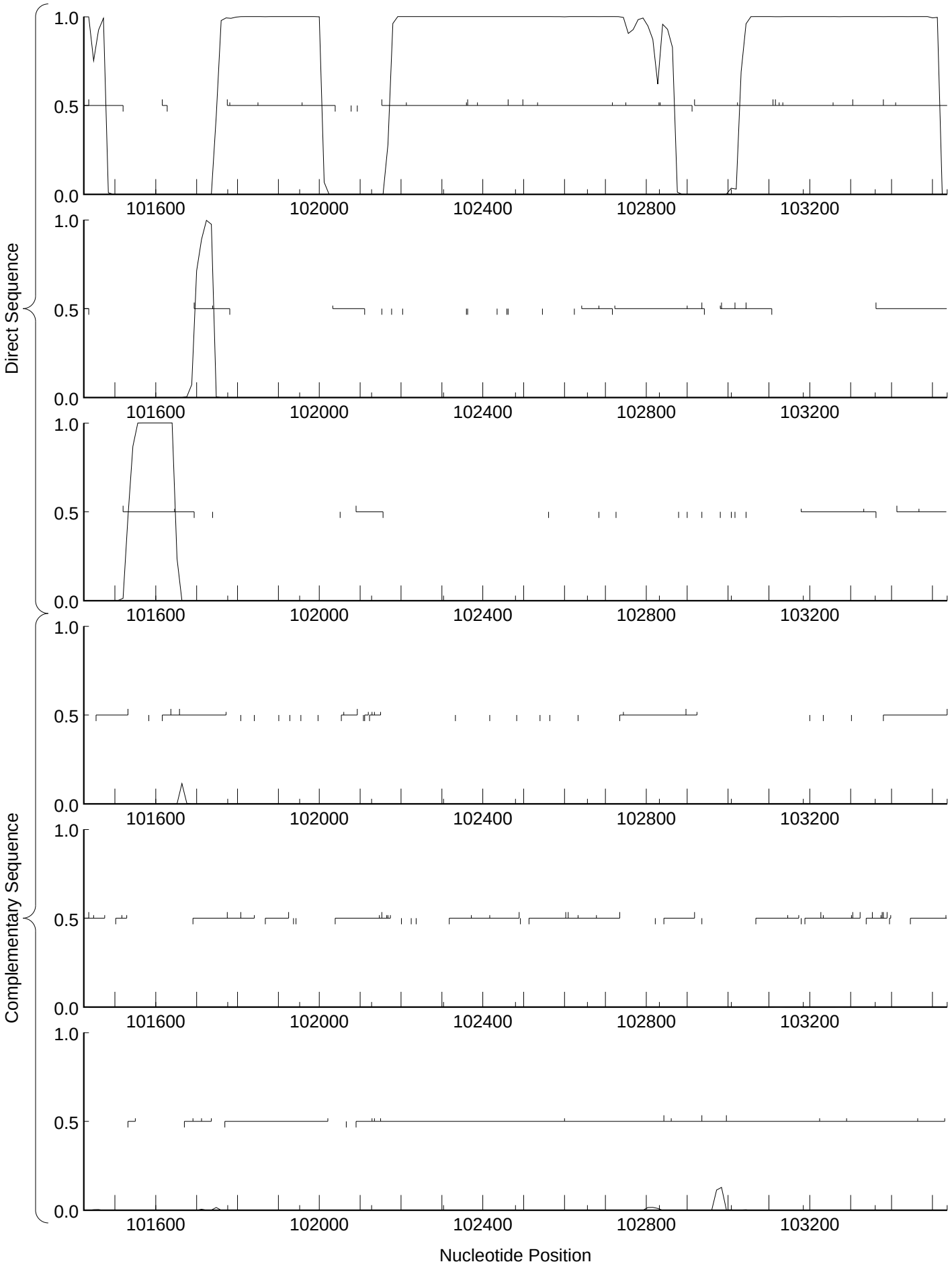

Direct Sequence
Complementary Sequence
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91200
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92000
92400
92800
Nucleotide Position

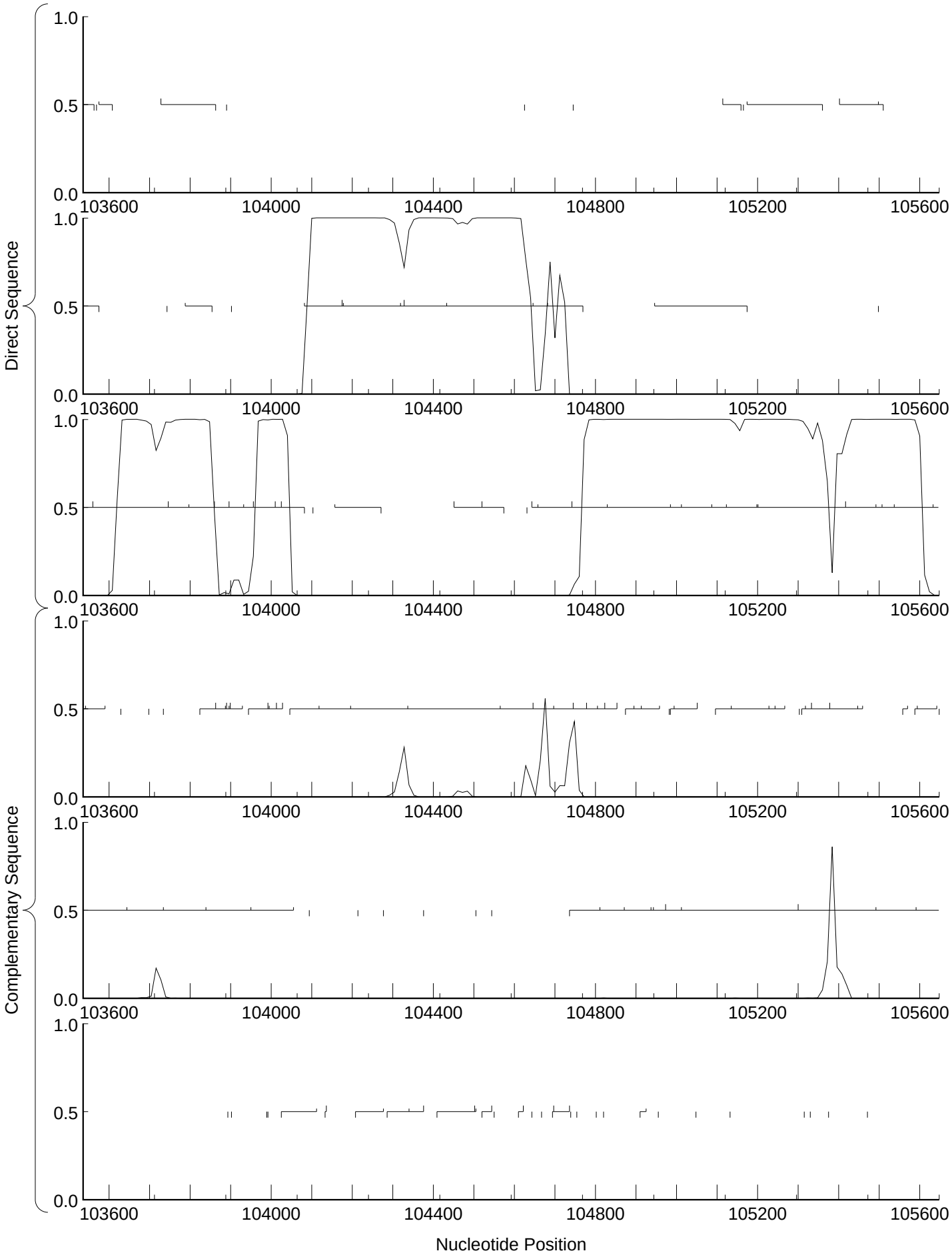

Direct Sequence
Complementary Sequence
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94000
94400
94800
Nucleotide Position

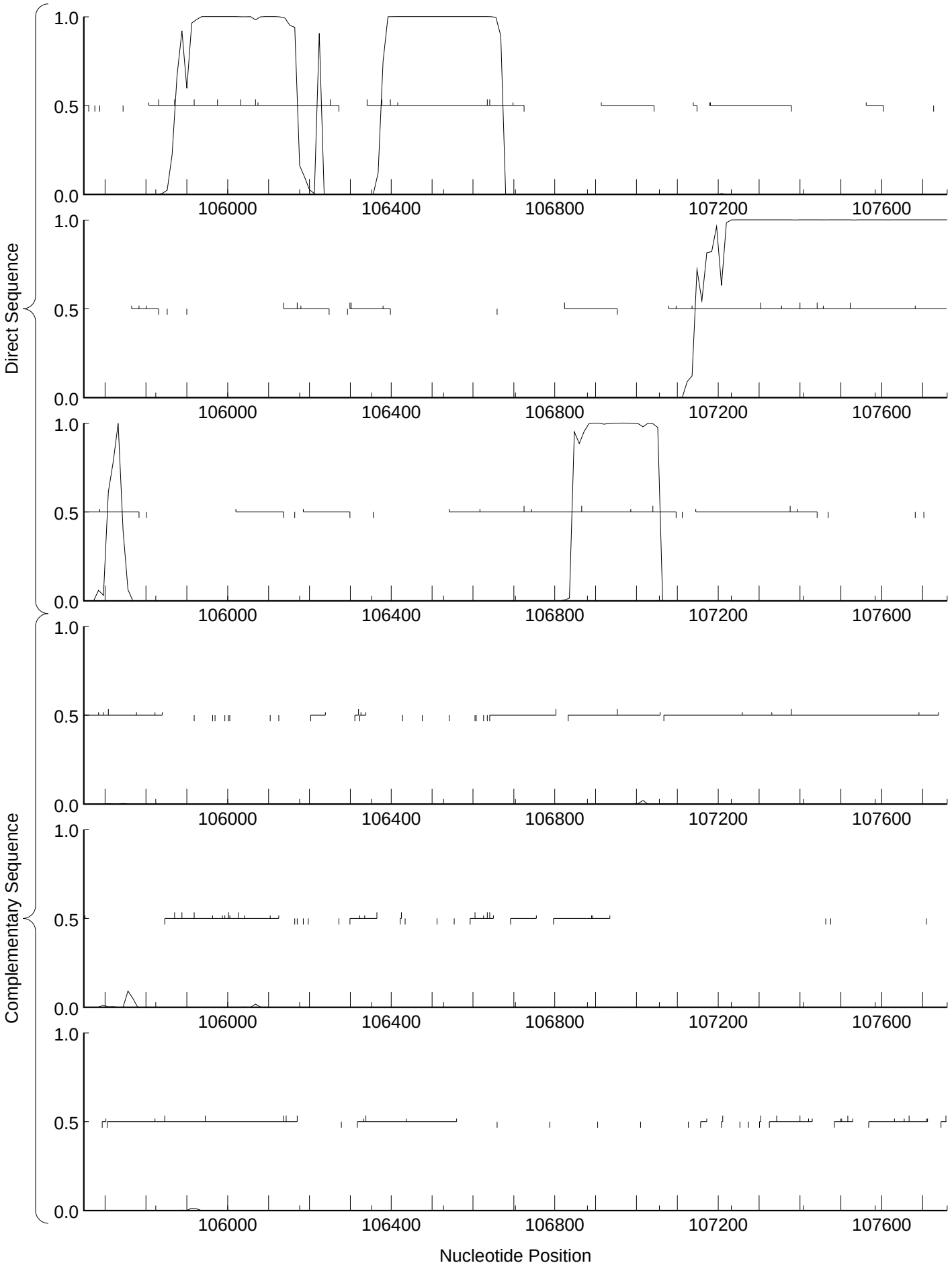

Direct Sequence
Complementary Sequence
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96000
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96800
97200
Nucleotide Position

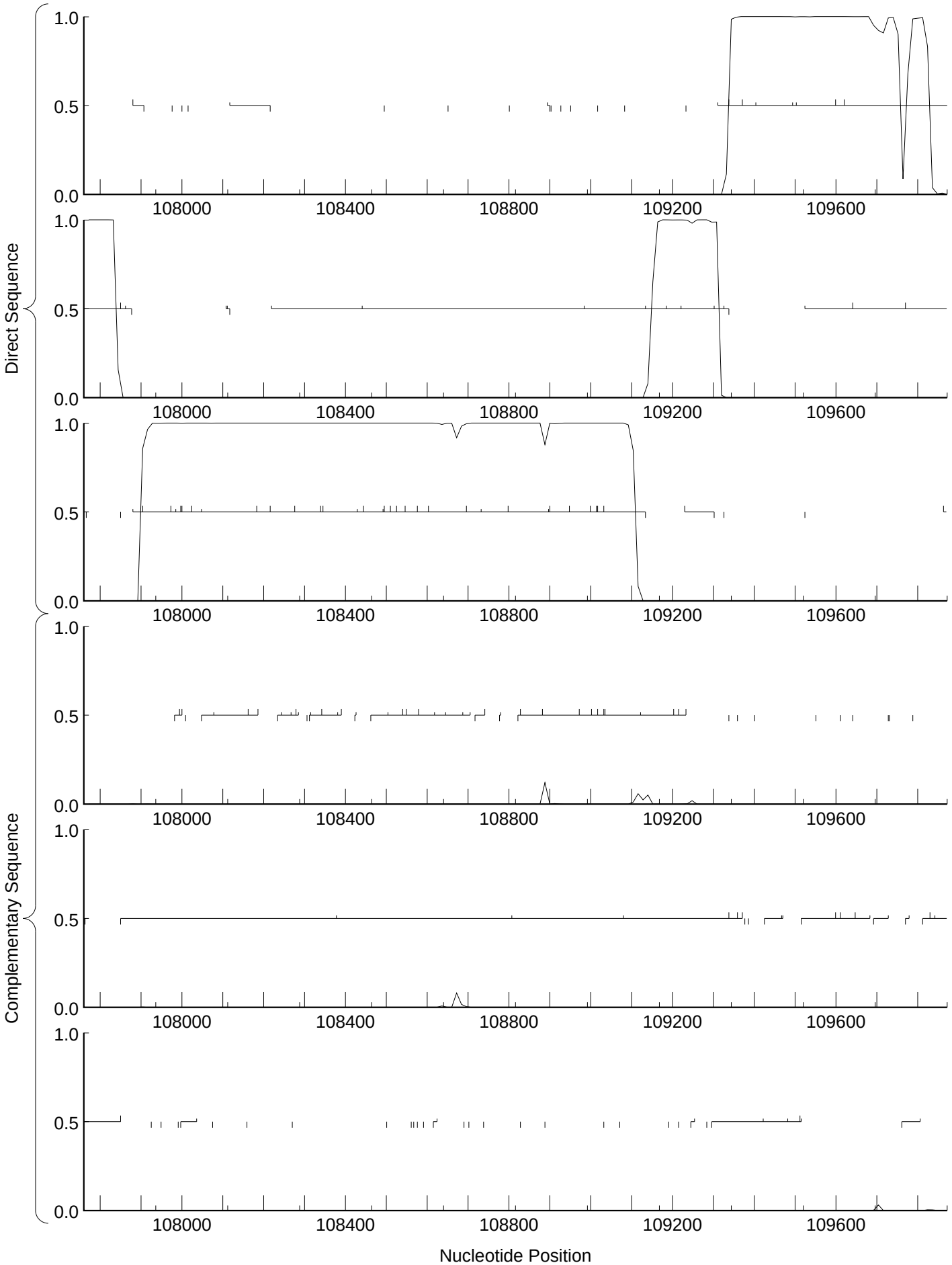

Direct Sequence
Complementary Sequence
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98000
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98800
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Nucleotide Position

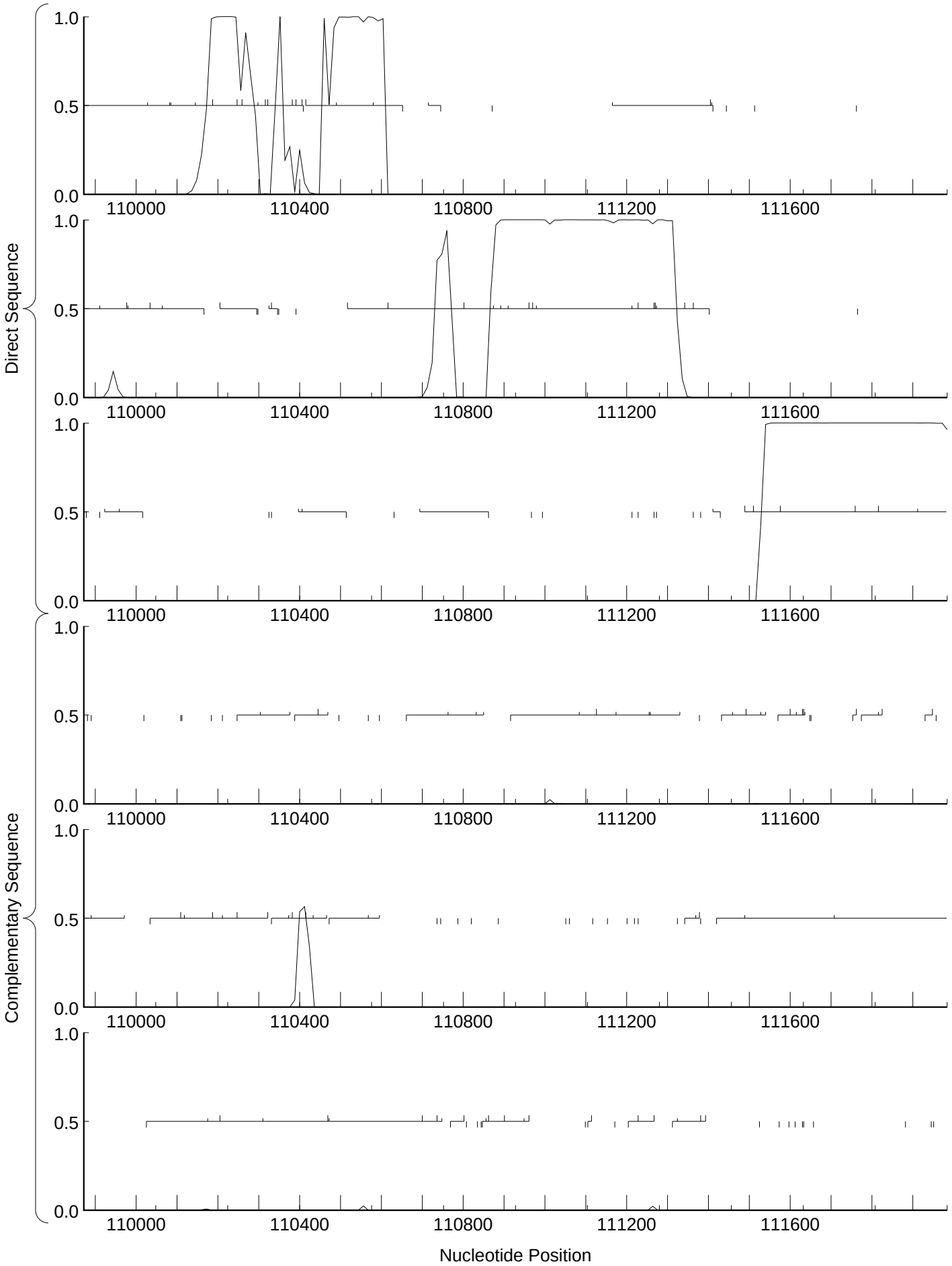

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Complementary Sequence
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100800
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Nucleotide Position

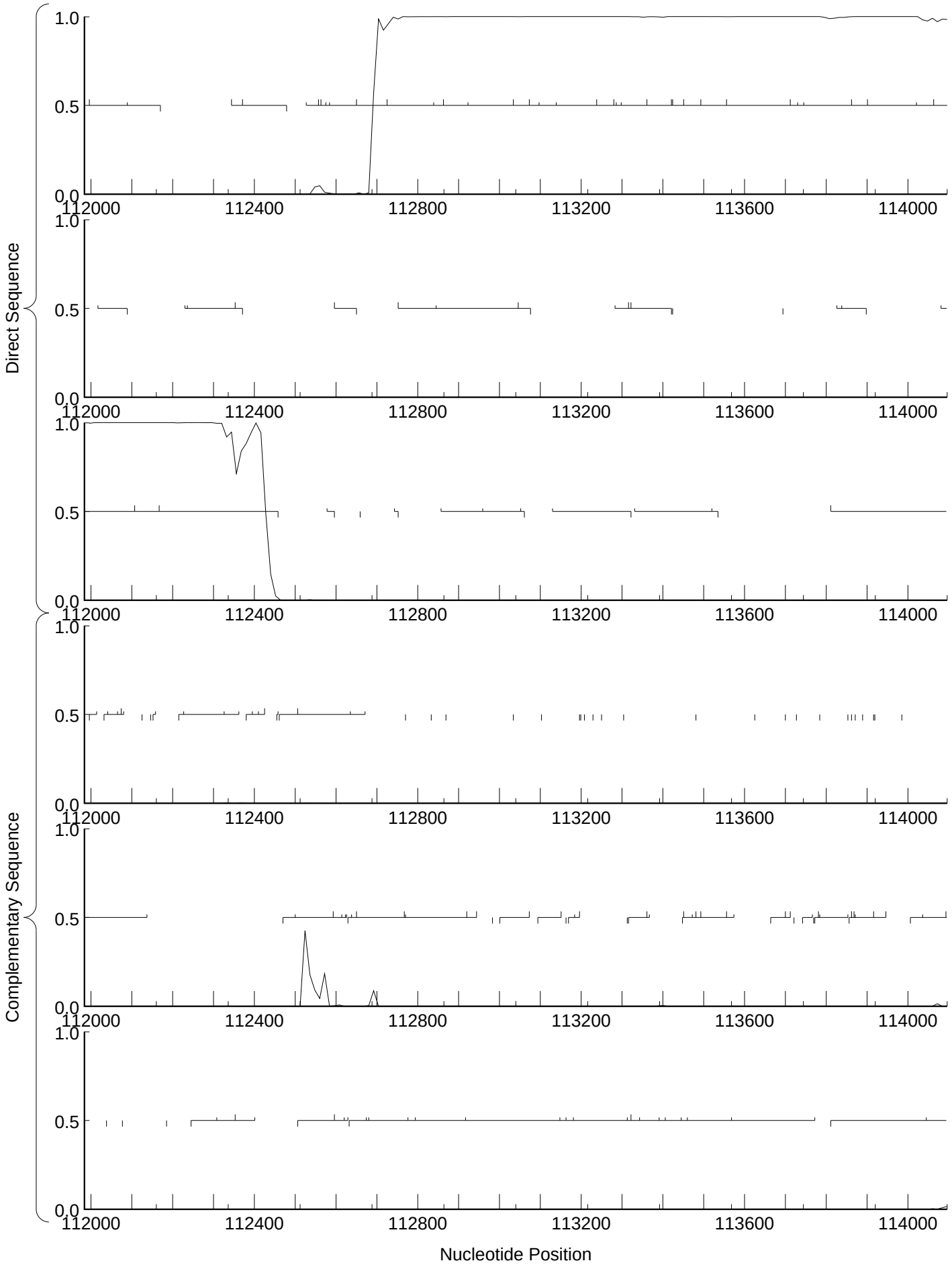

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Complementary Sequence
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102800
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Nucleotide Position

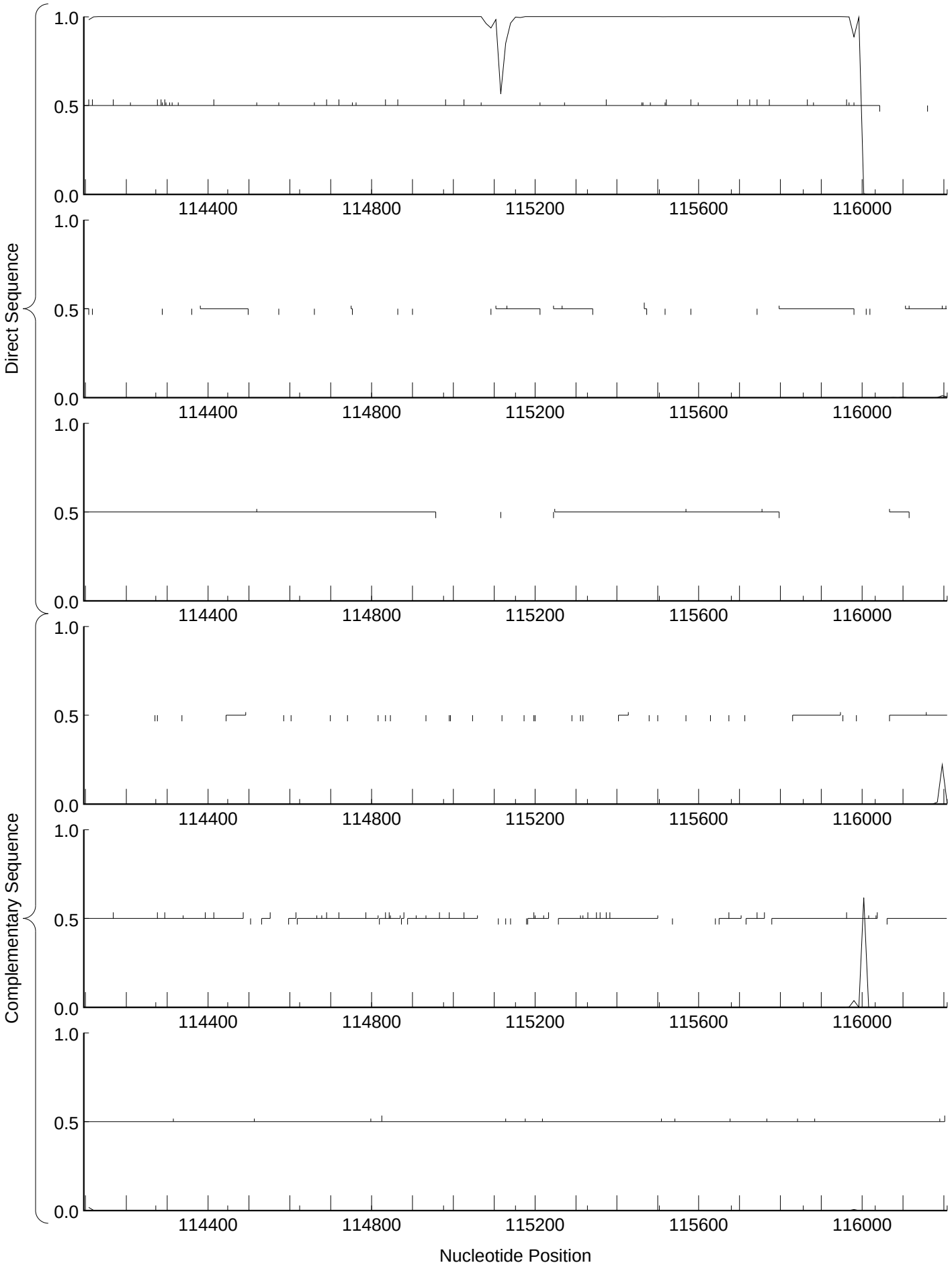

Direct Sequence
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105200
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Nucleotide Position

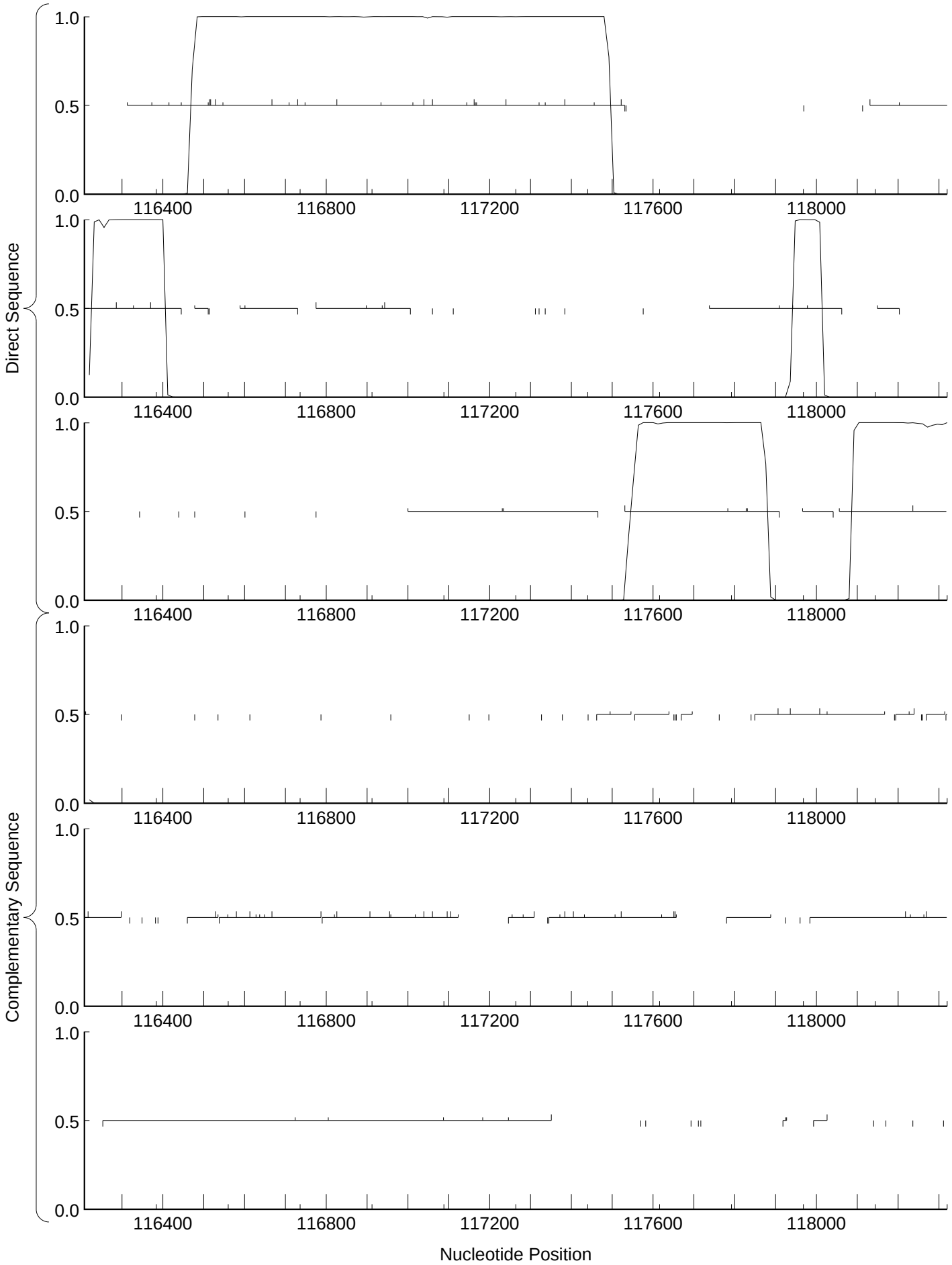

Direct Sequence
Complementary Sequence
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106800
107200
107600
Nucleotide Position

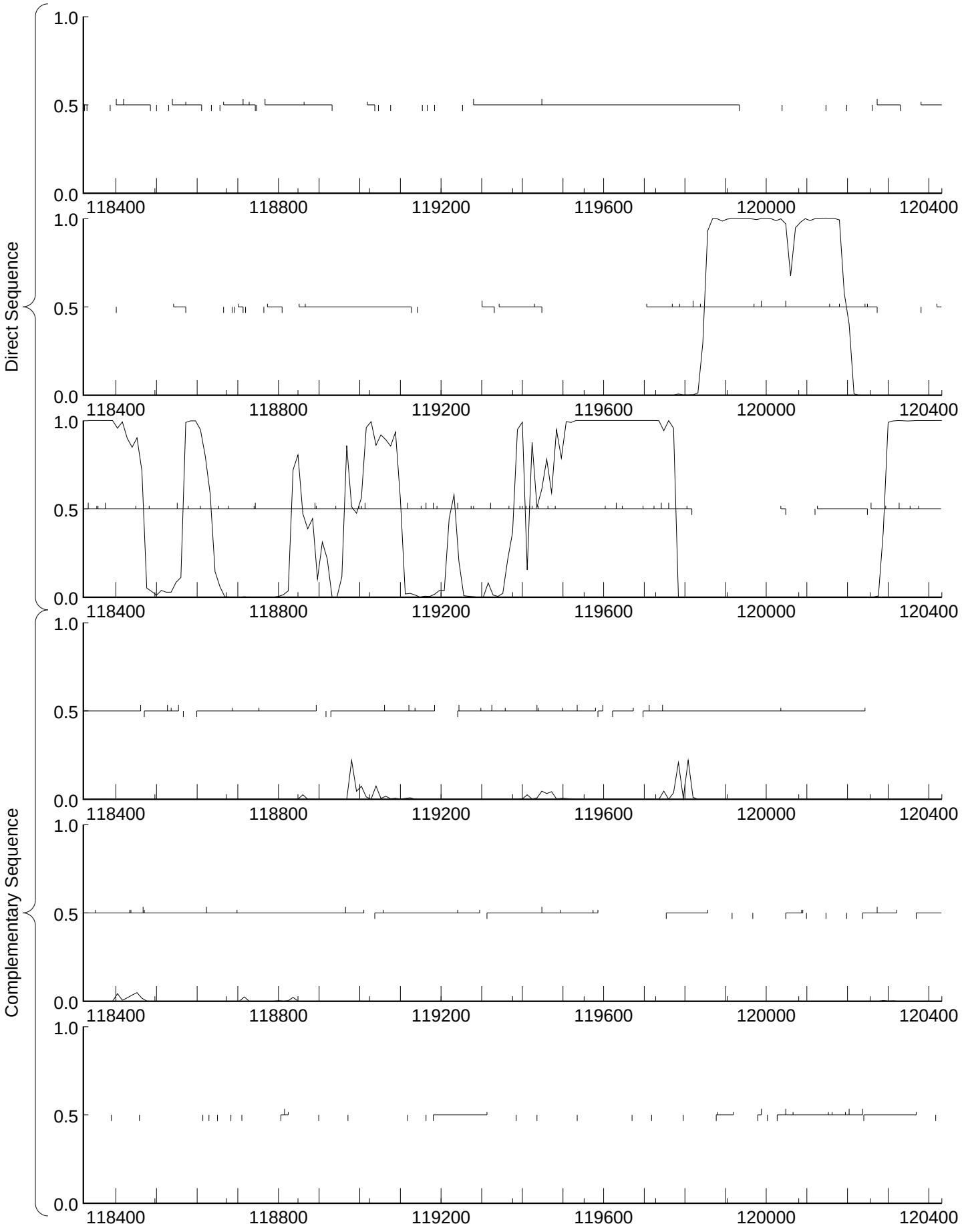

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Complementary Sequence
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109200
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Nucleotide Position

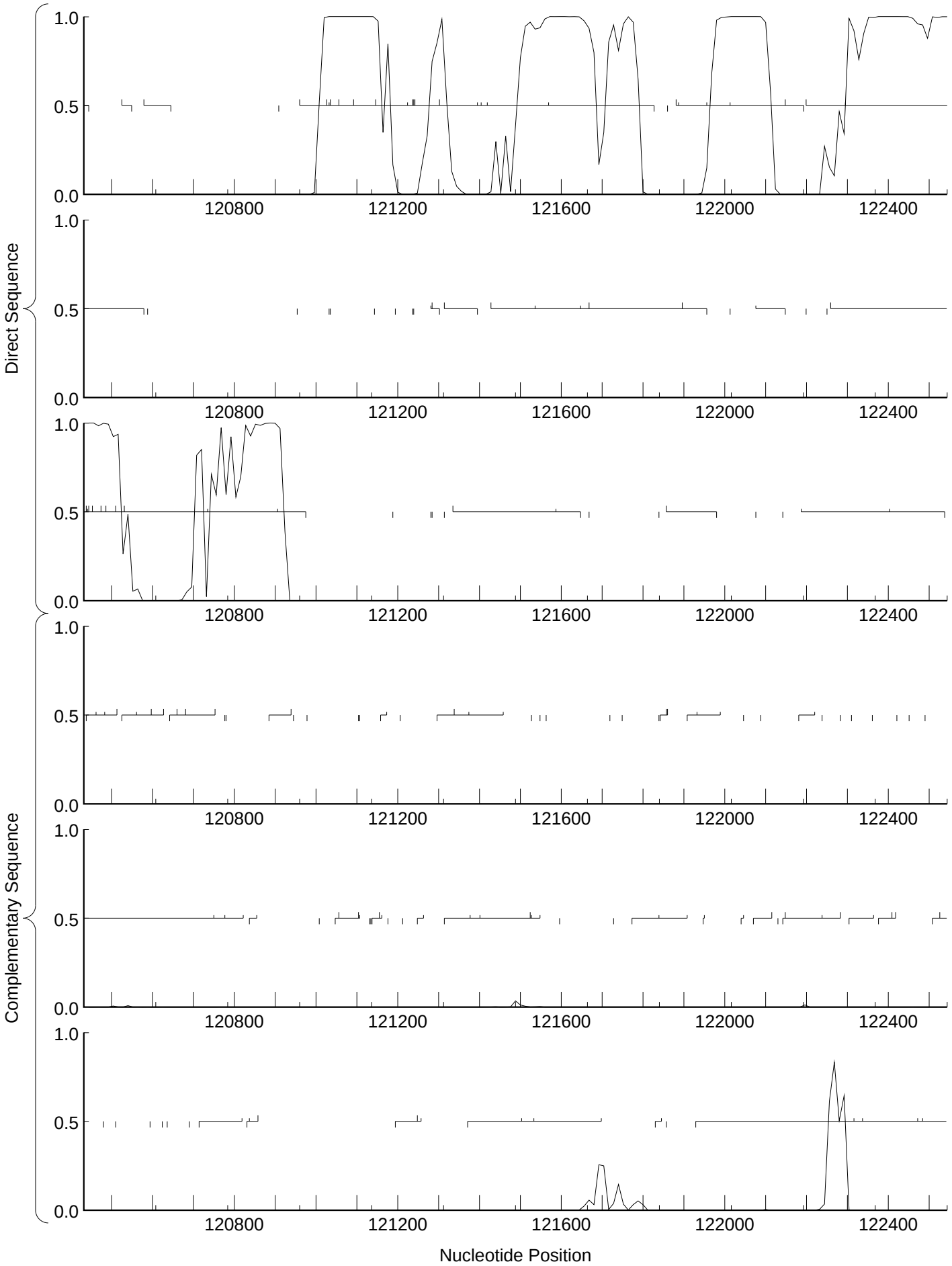

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Complementary Sequence
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110800
111200
111600
Nucleotide Position

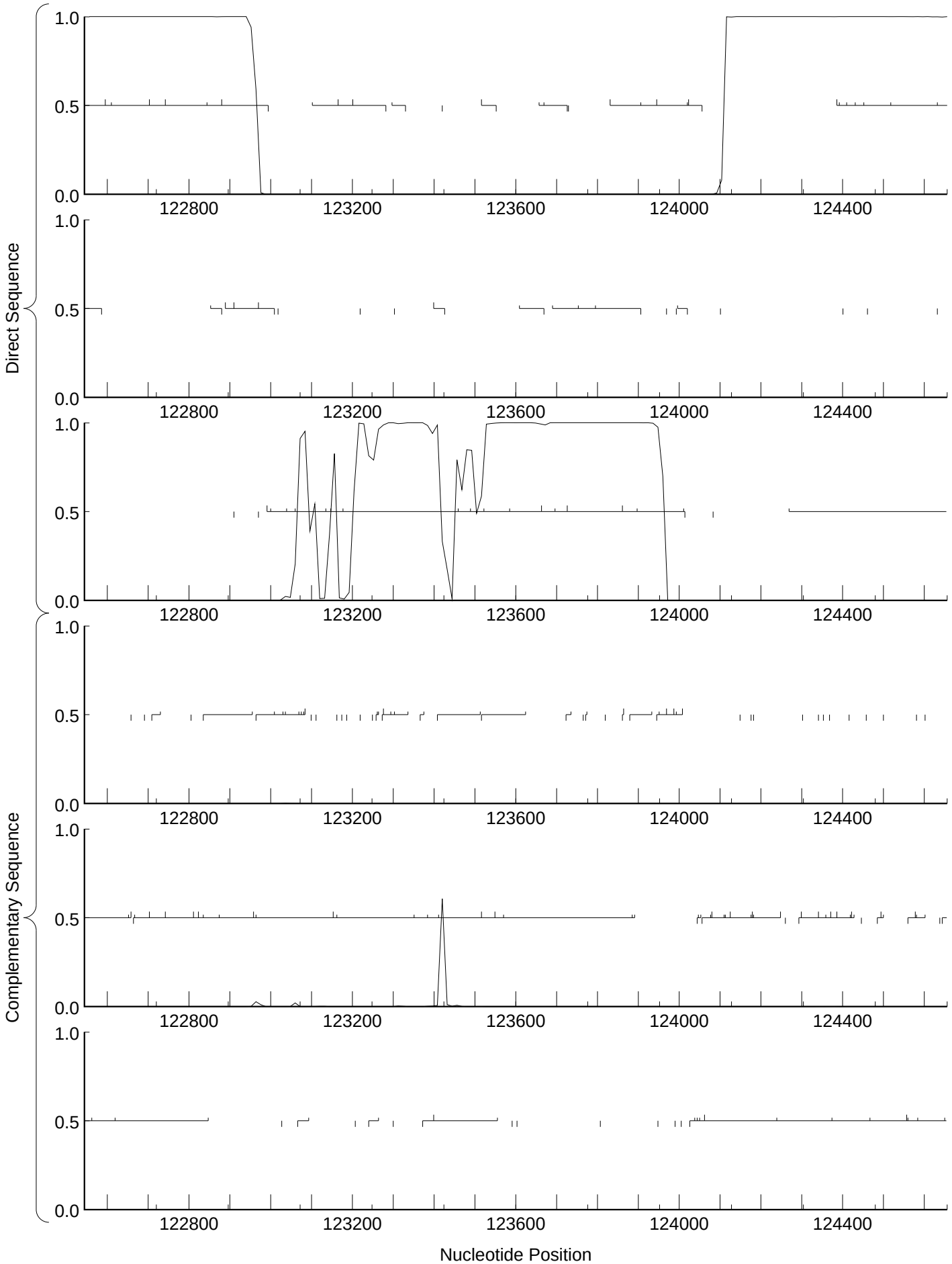

Direct Sequence
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112000
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113200
113600
114000
Nucleotide Position

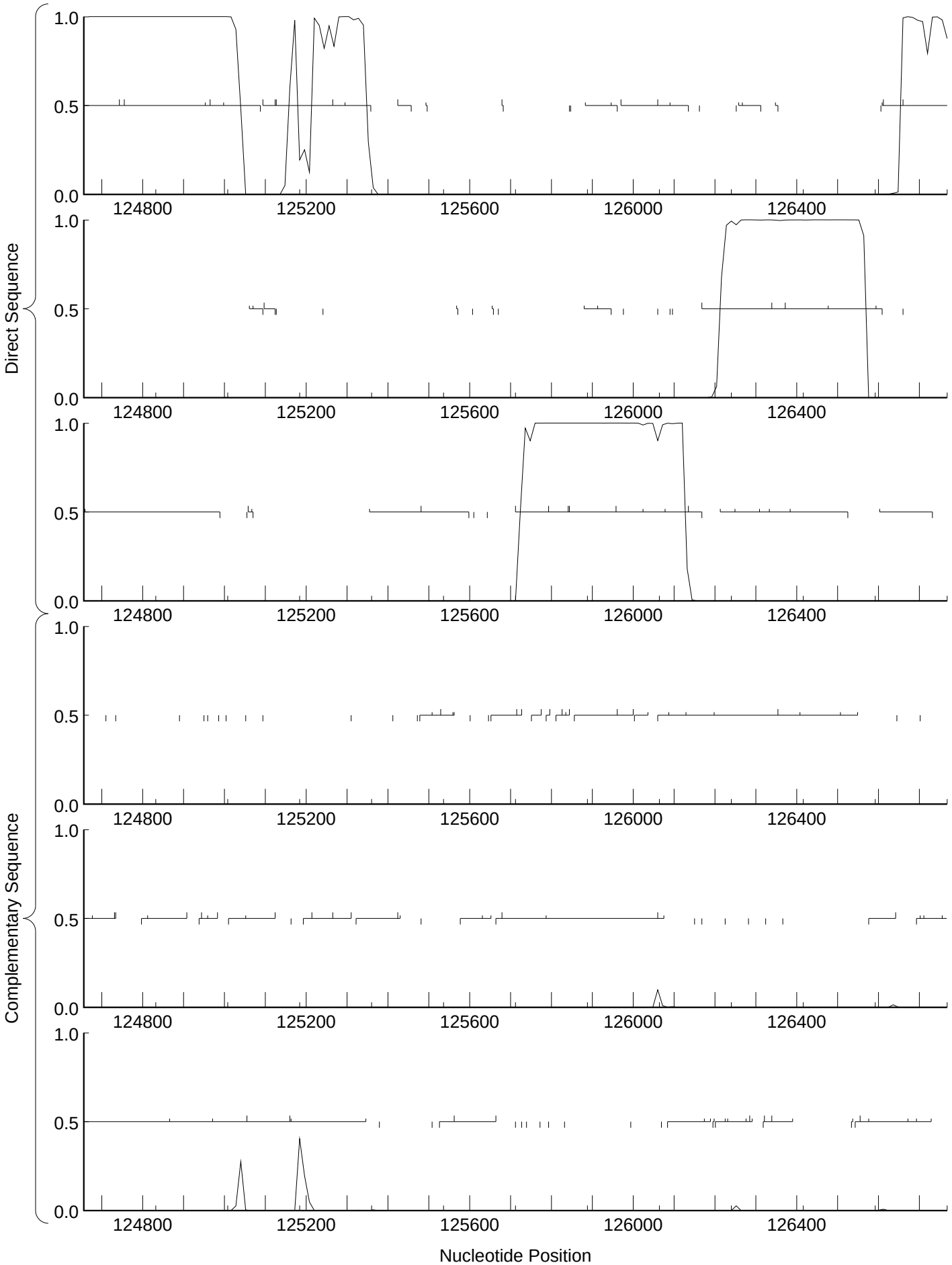

Direct Sequence
Complementary Sequence
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0.0
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114800
115200
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116000
Nucleotide Position

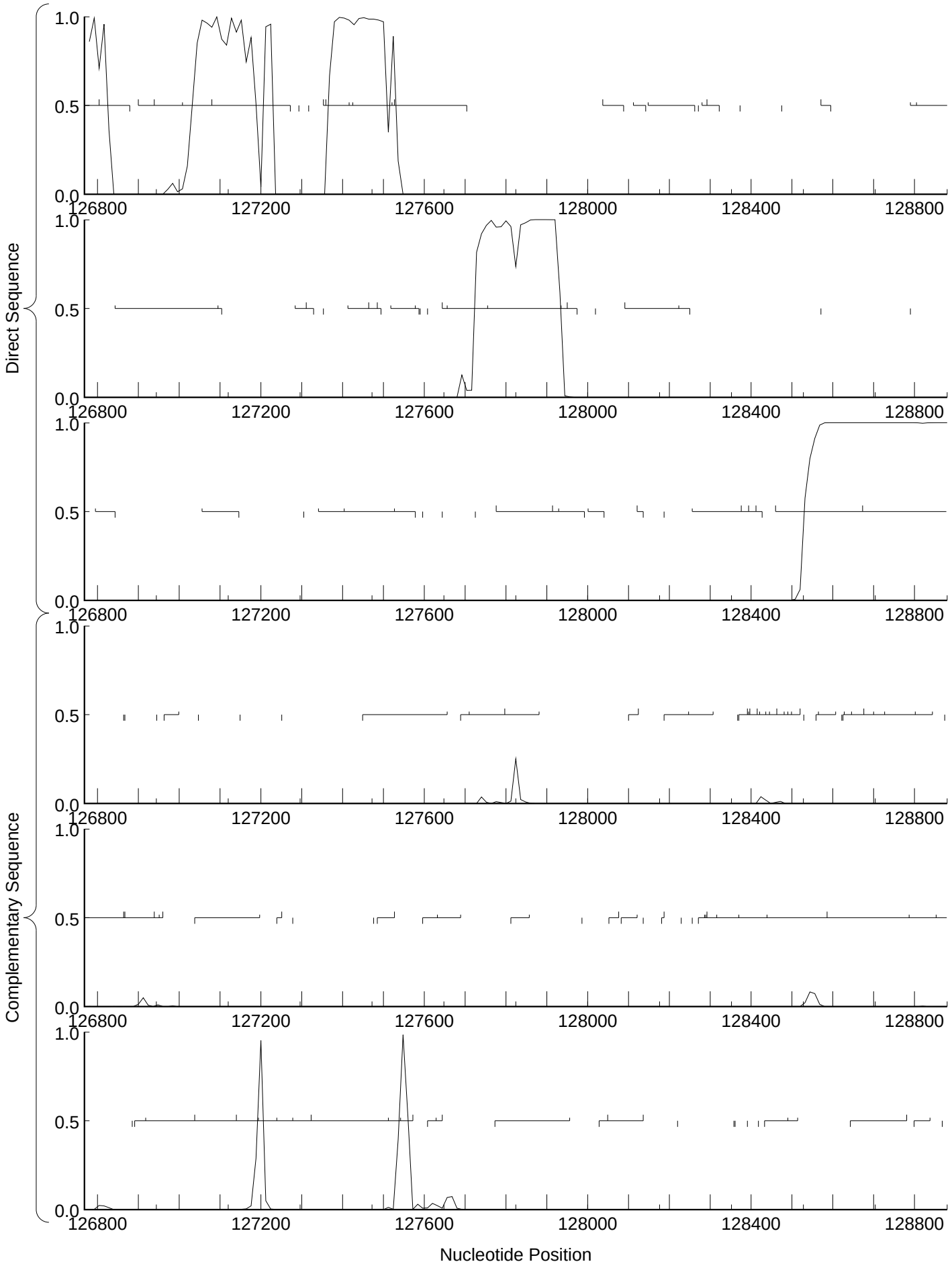

Direct Sequence
Complementary Sequence
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116800
117200
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Nucleotide Position

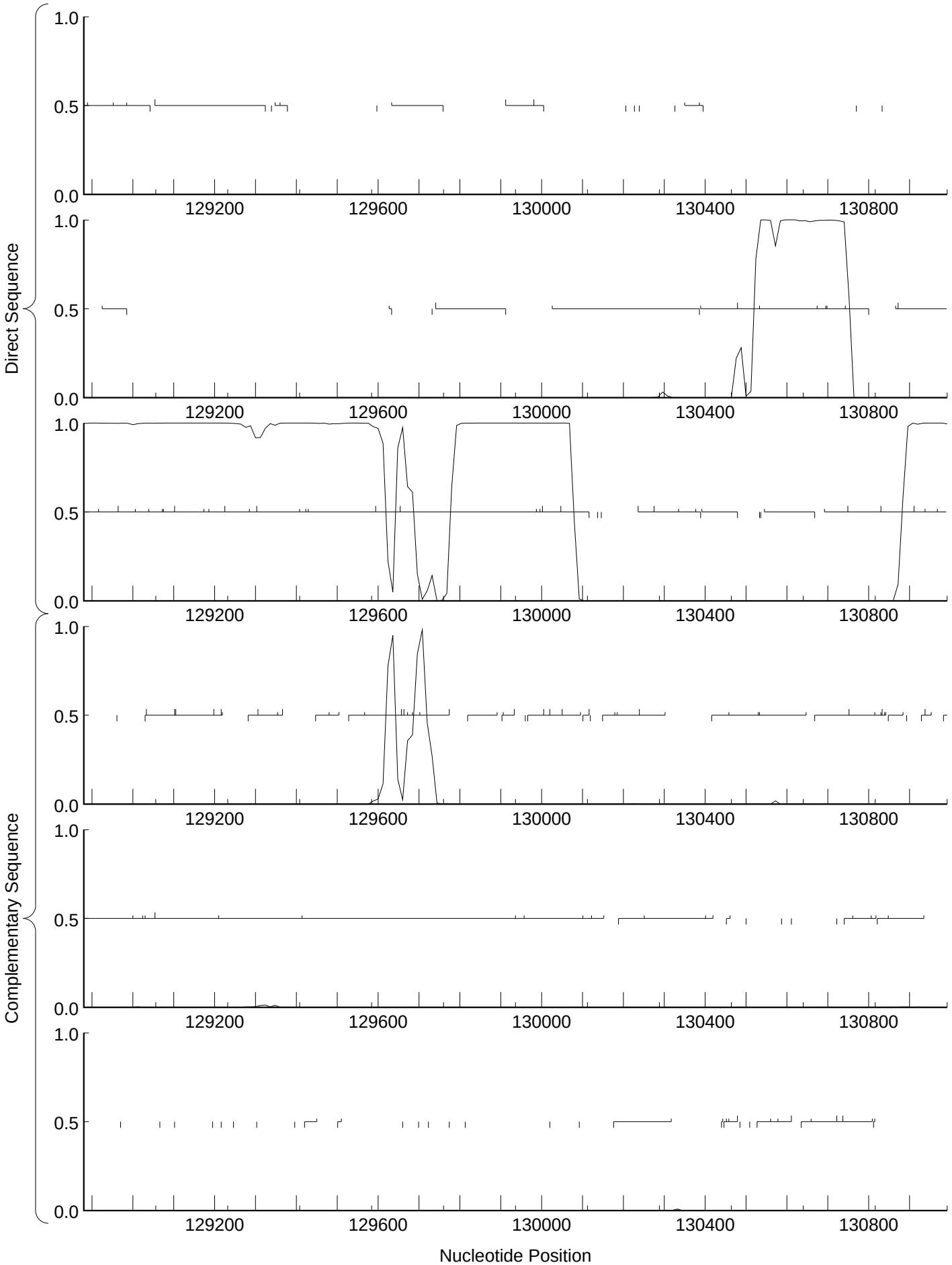

Direct Sequence
Complementary Sequence
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119200
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120000
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Nucleotide Position

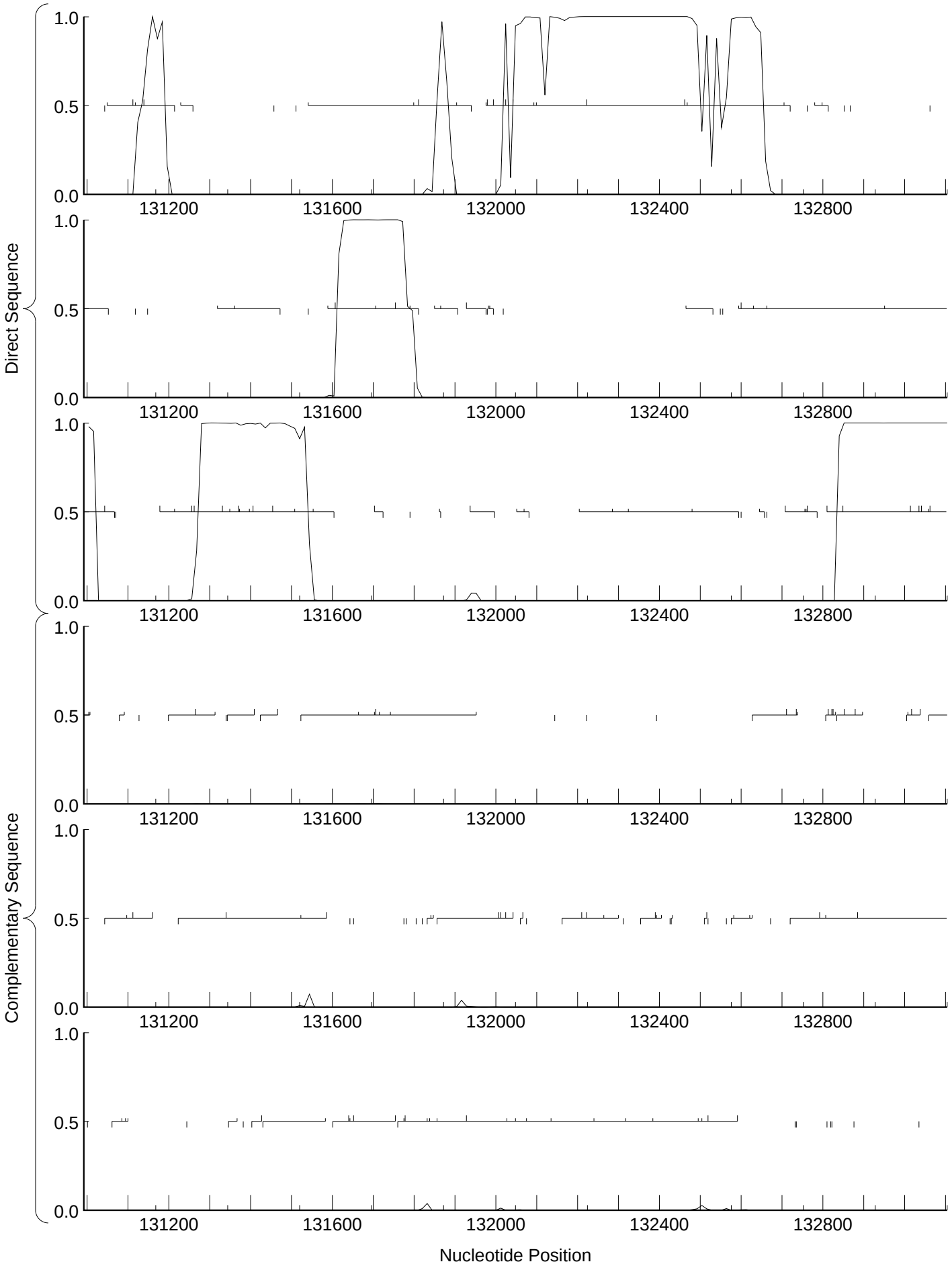

Direct Sequence
Complementary Sequence
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Nucleotide Position

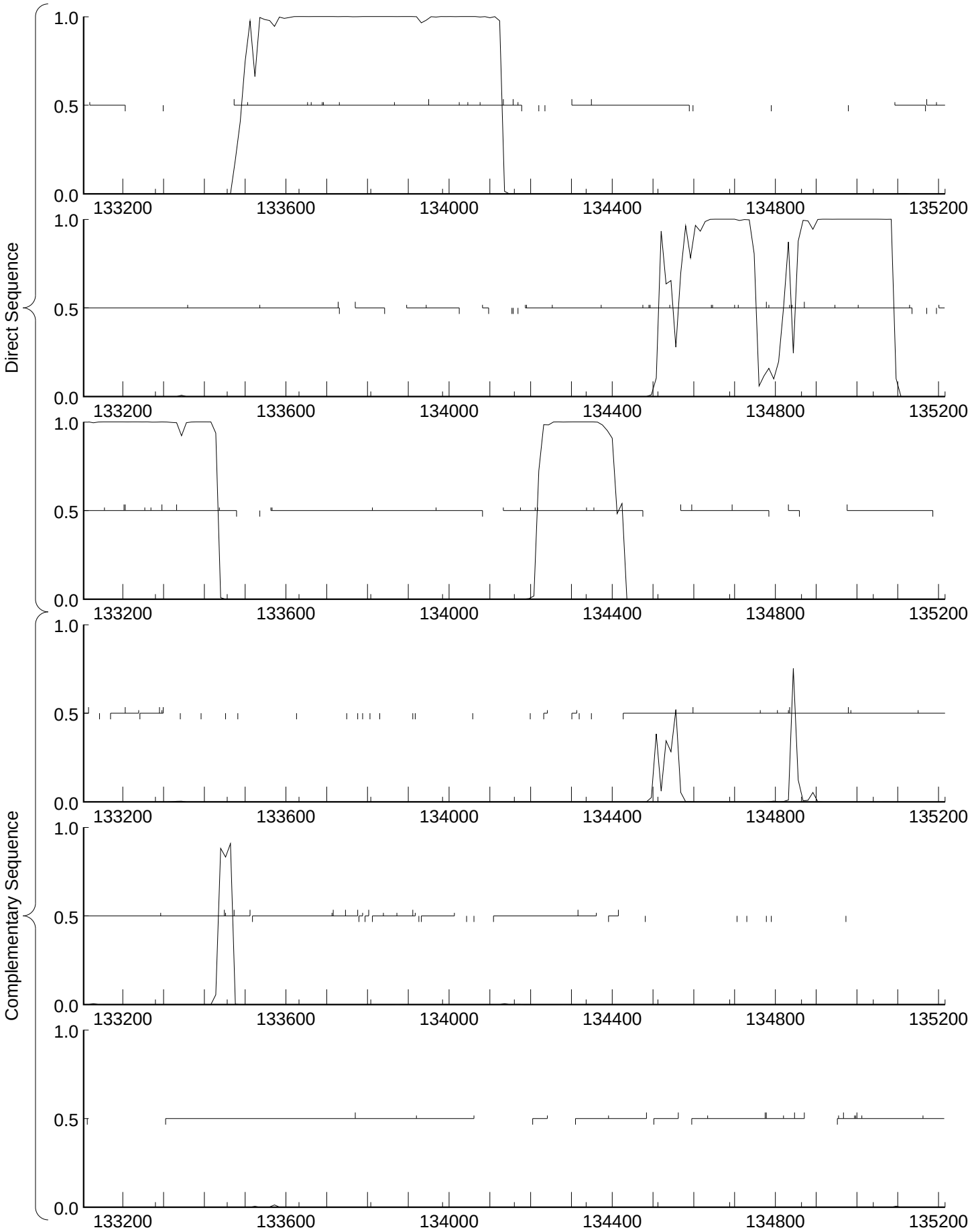

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Nucleotide Position

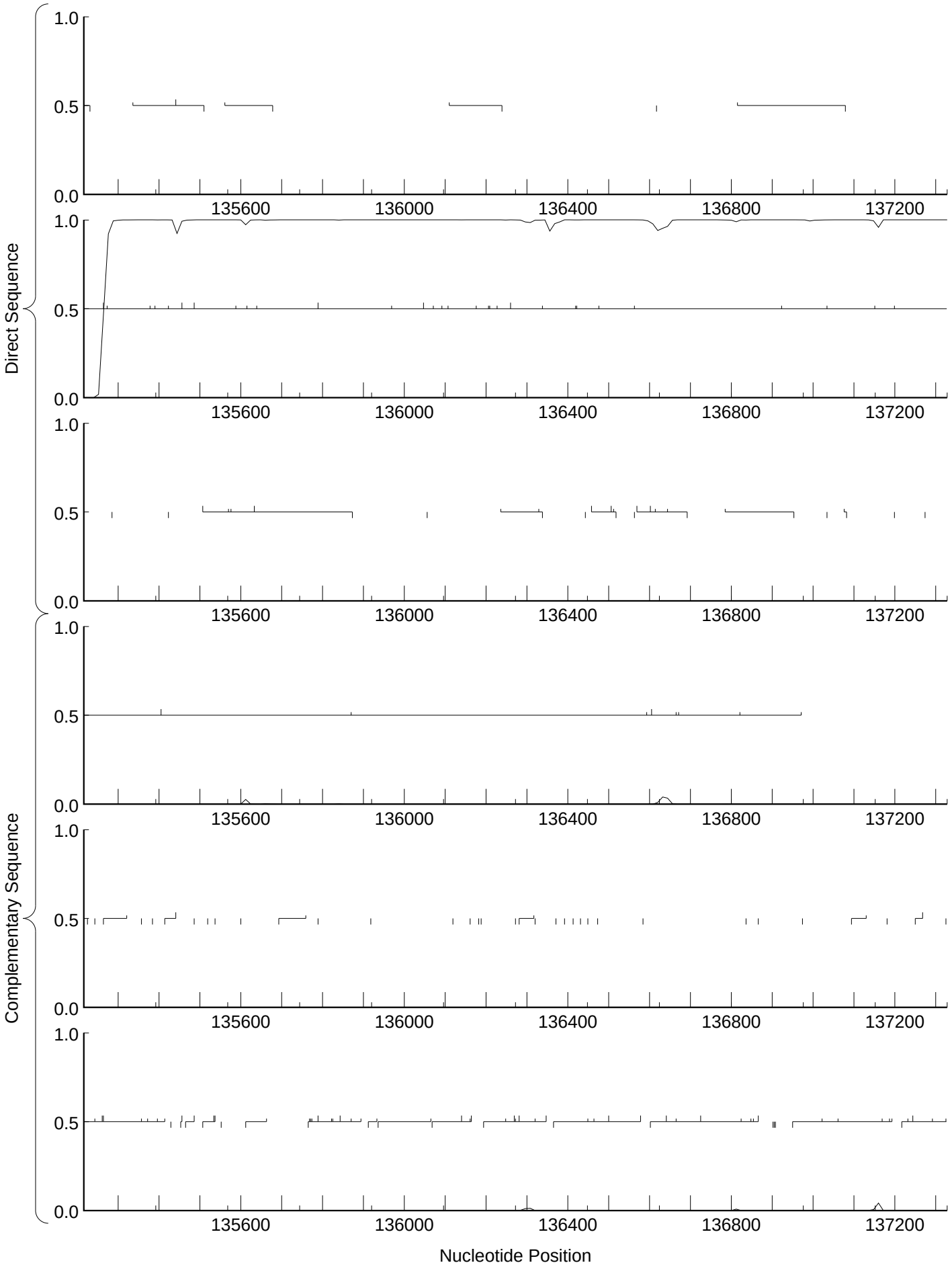

Direct Sequence
Complementary Sequence
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Nucleotide Position

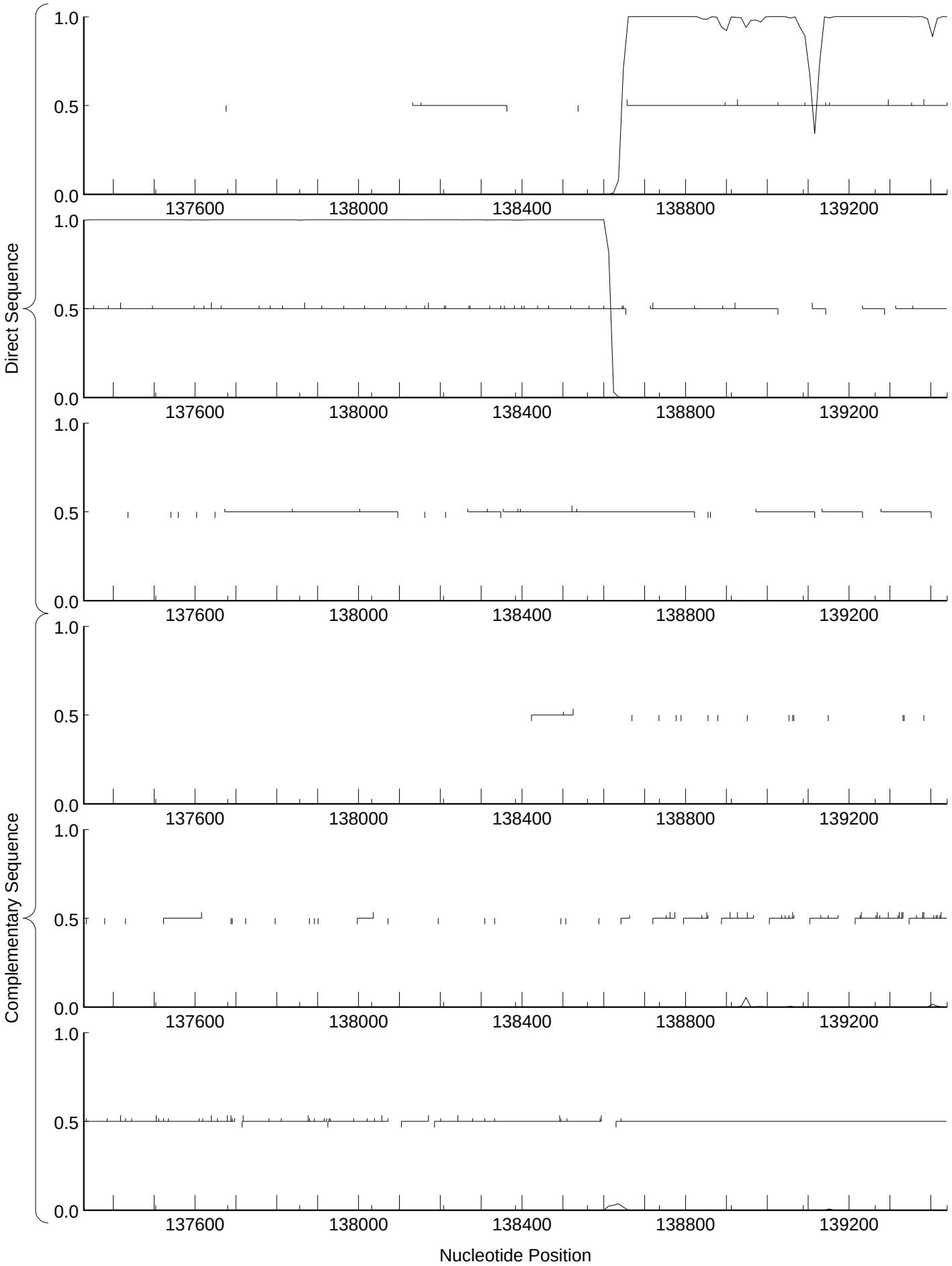

Direct Sequence
Complementary Sequence
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Nucleotide Position

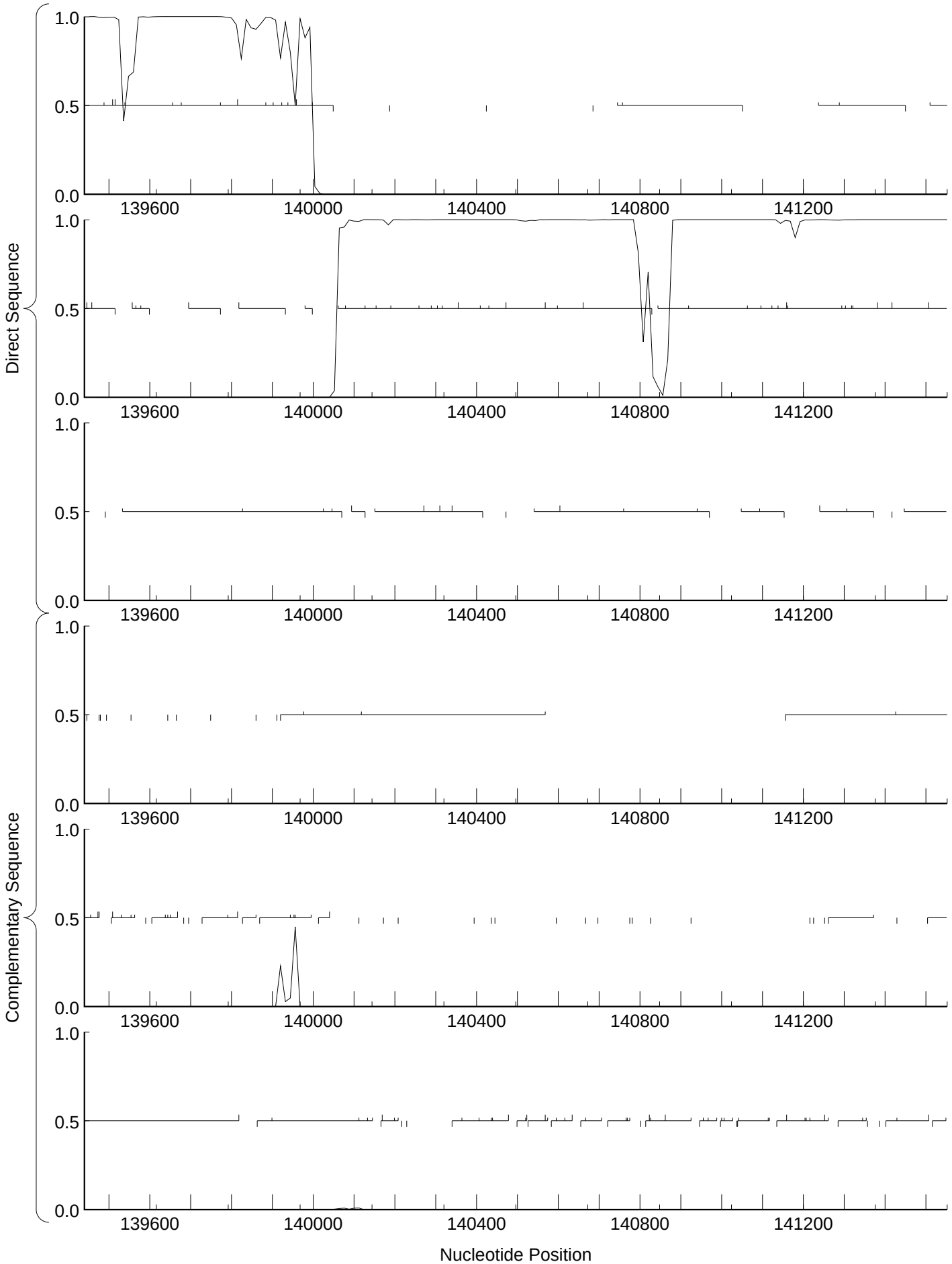

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Nucleotide Position

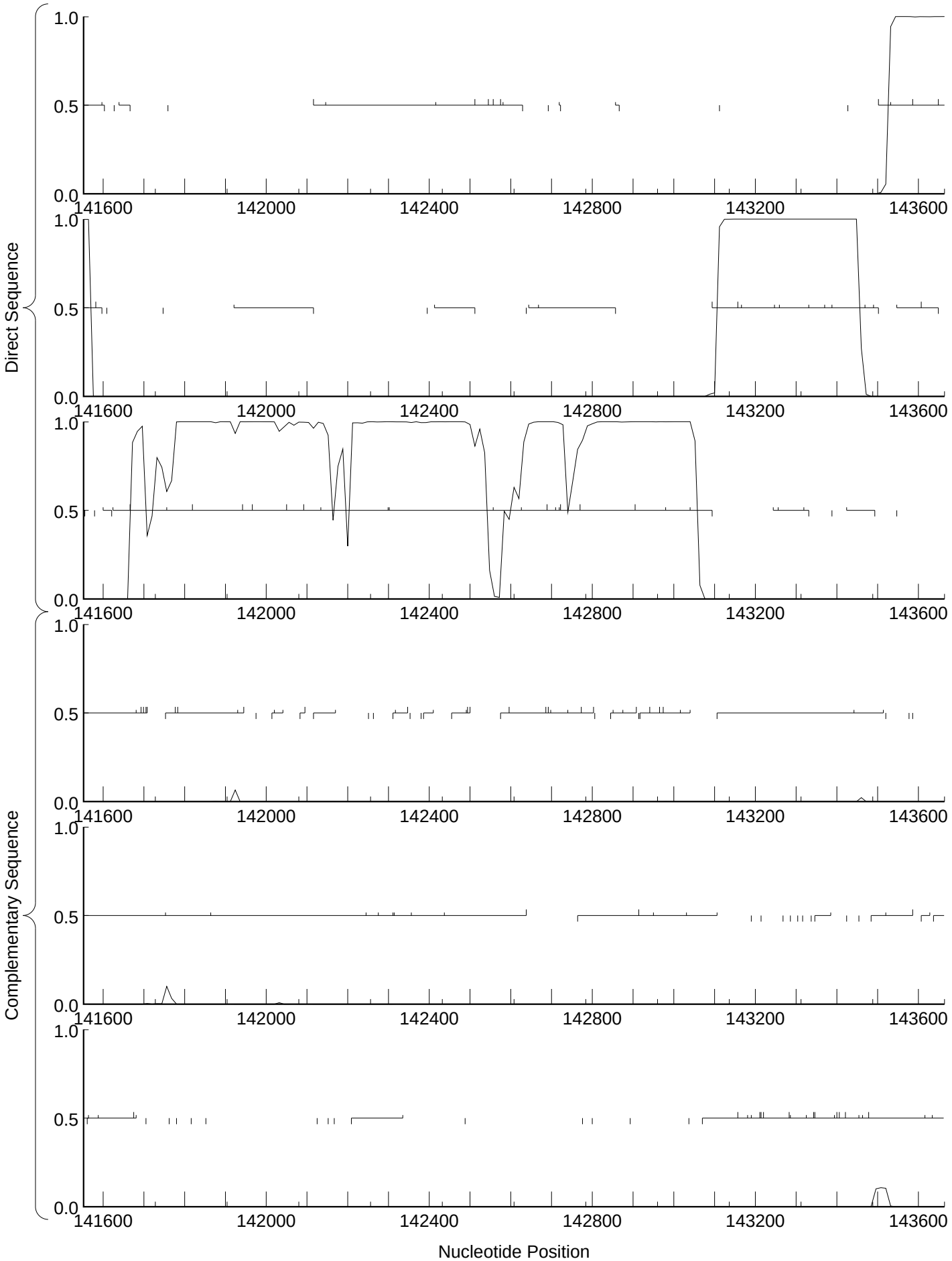

Direct Sequence
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Nucleotide Position

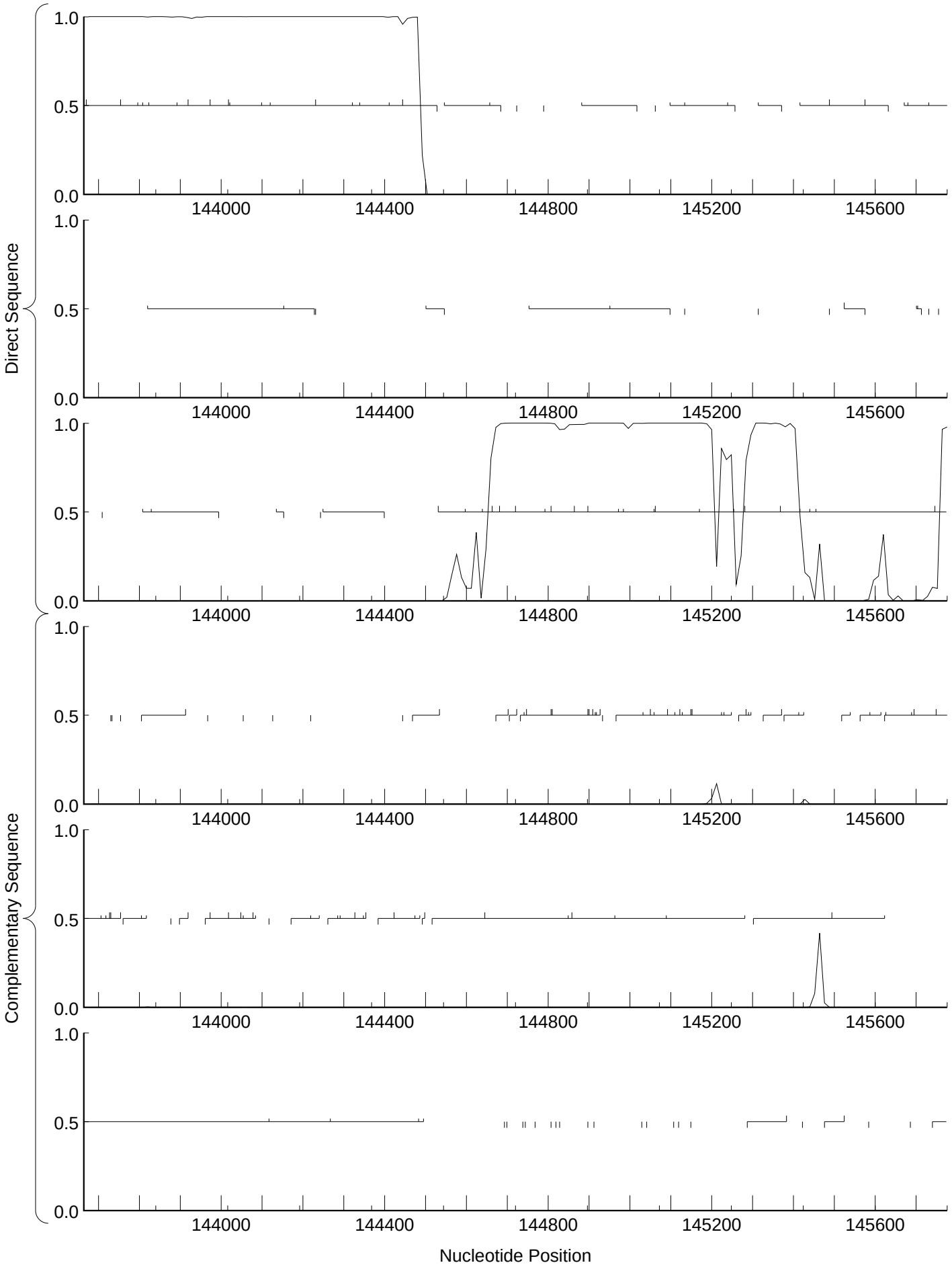

Direct Sequence
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Nucleotide Position

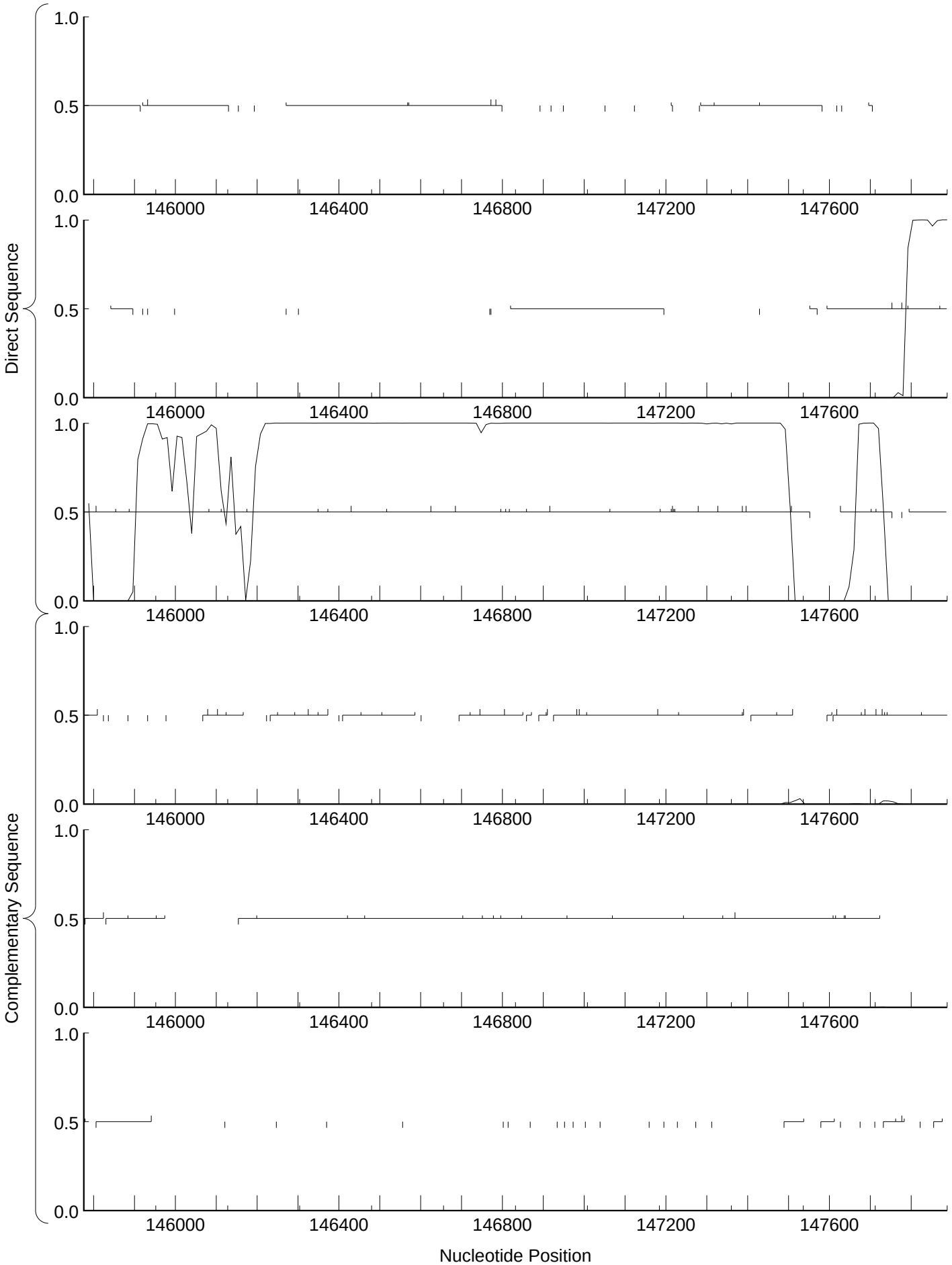

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Nucleotide Position

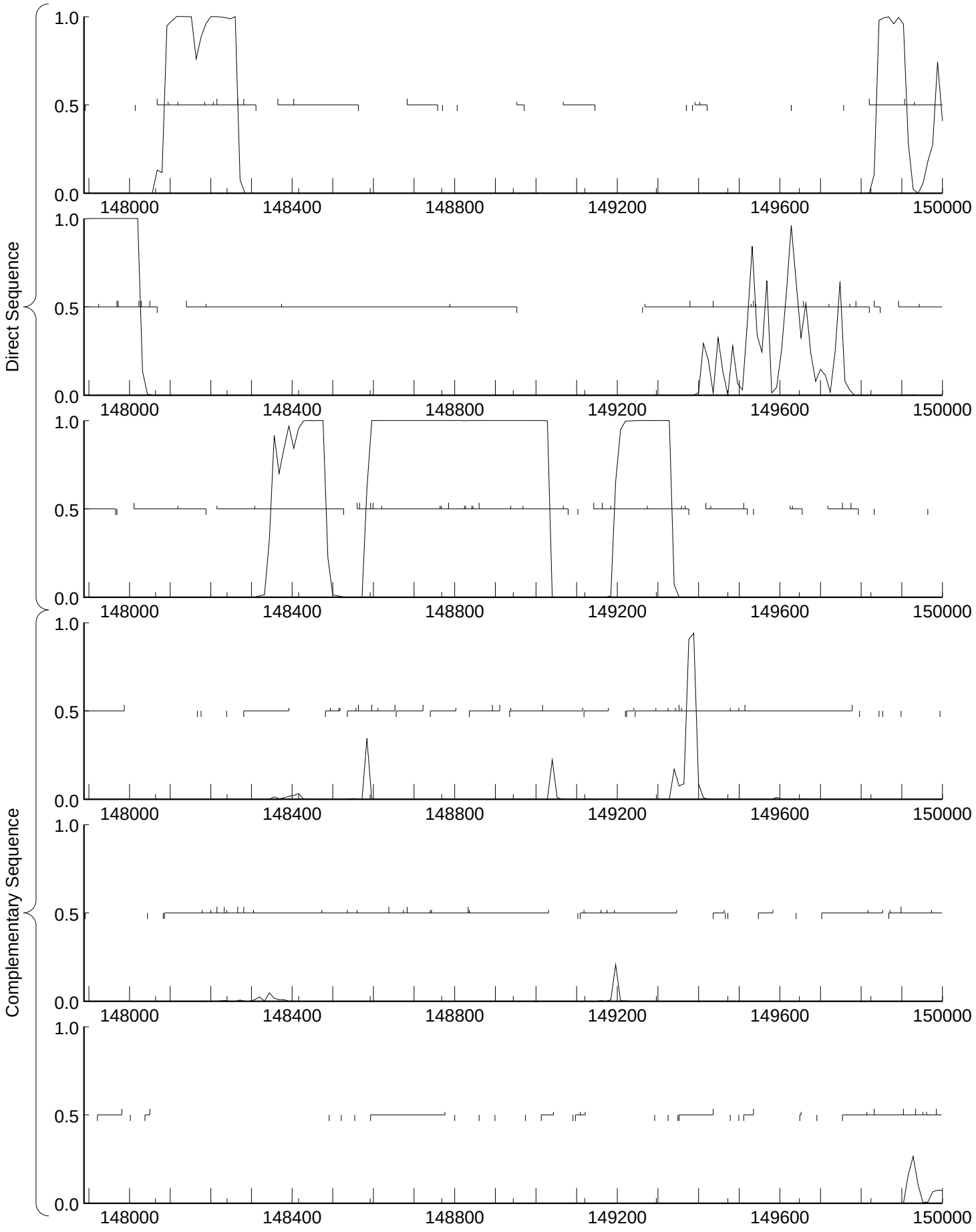

Direct Sequence
Complementary Sequence
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Nucleotide Position

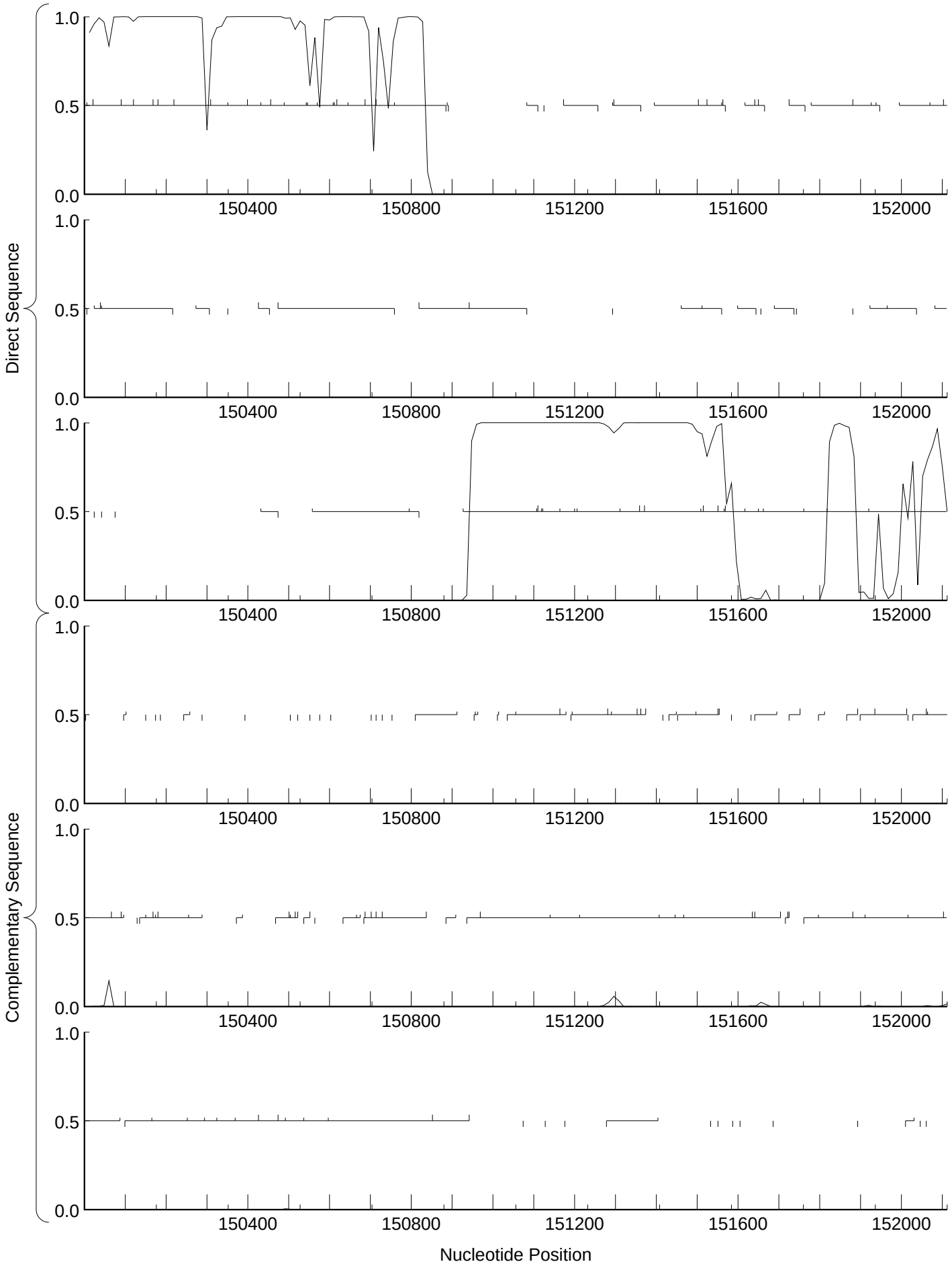

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Complementary Sequence
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Nucleotide Position

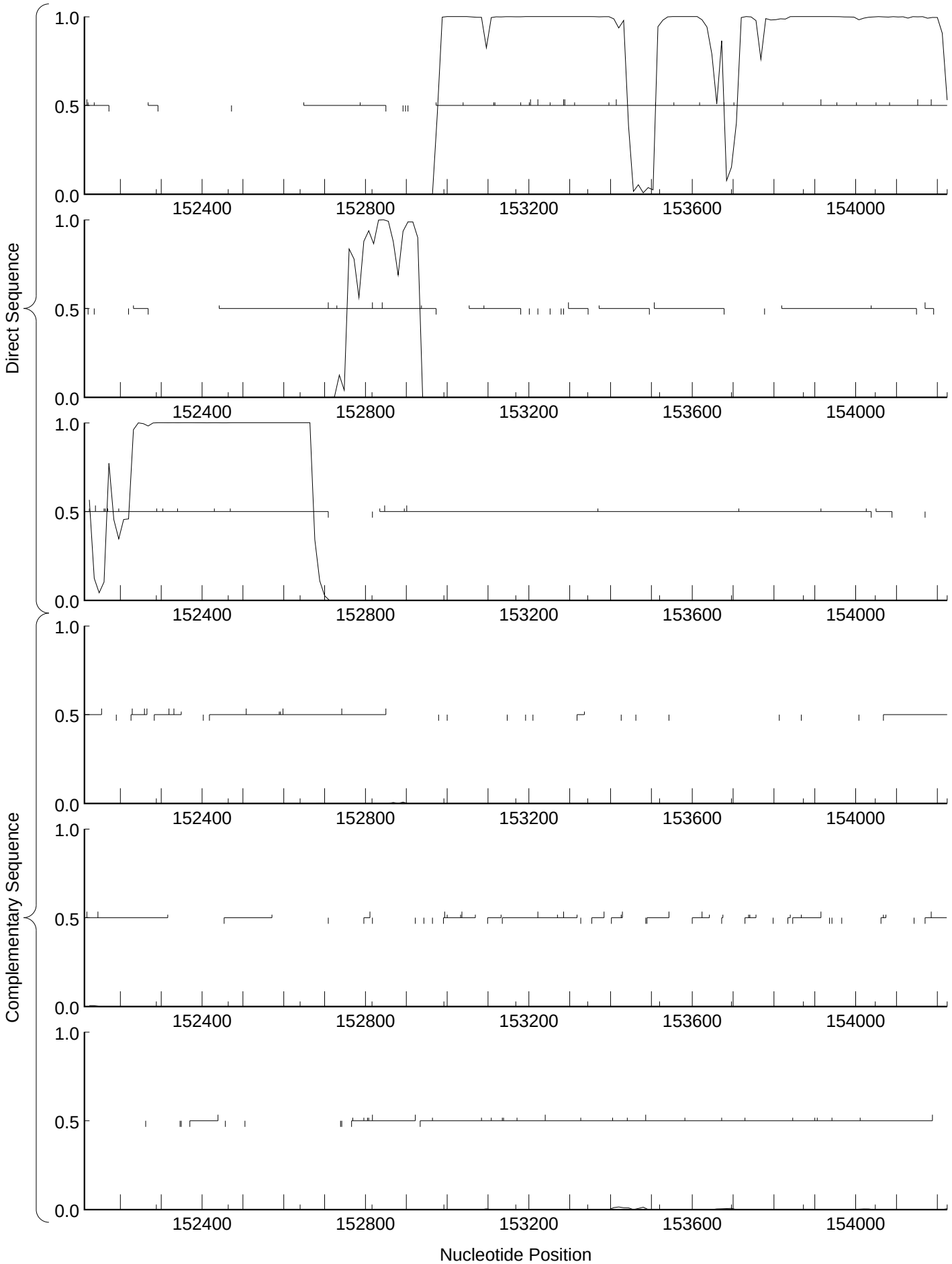

Direct Sequence
Complementary Sequence
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Nucleotide Position

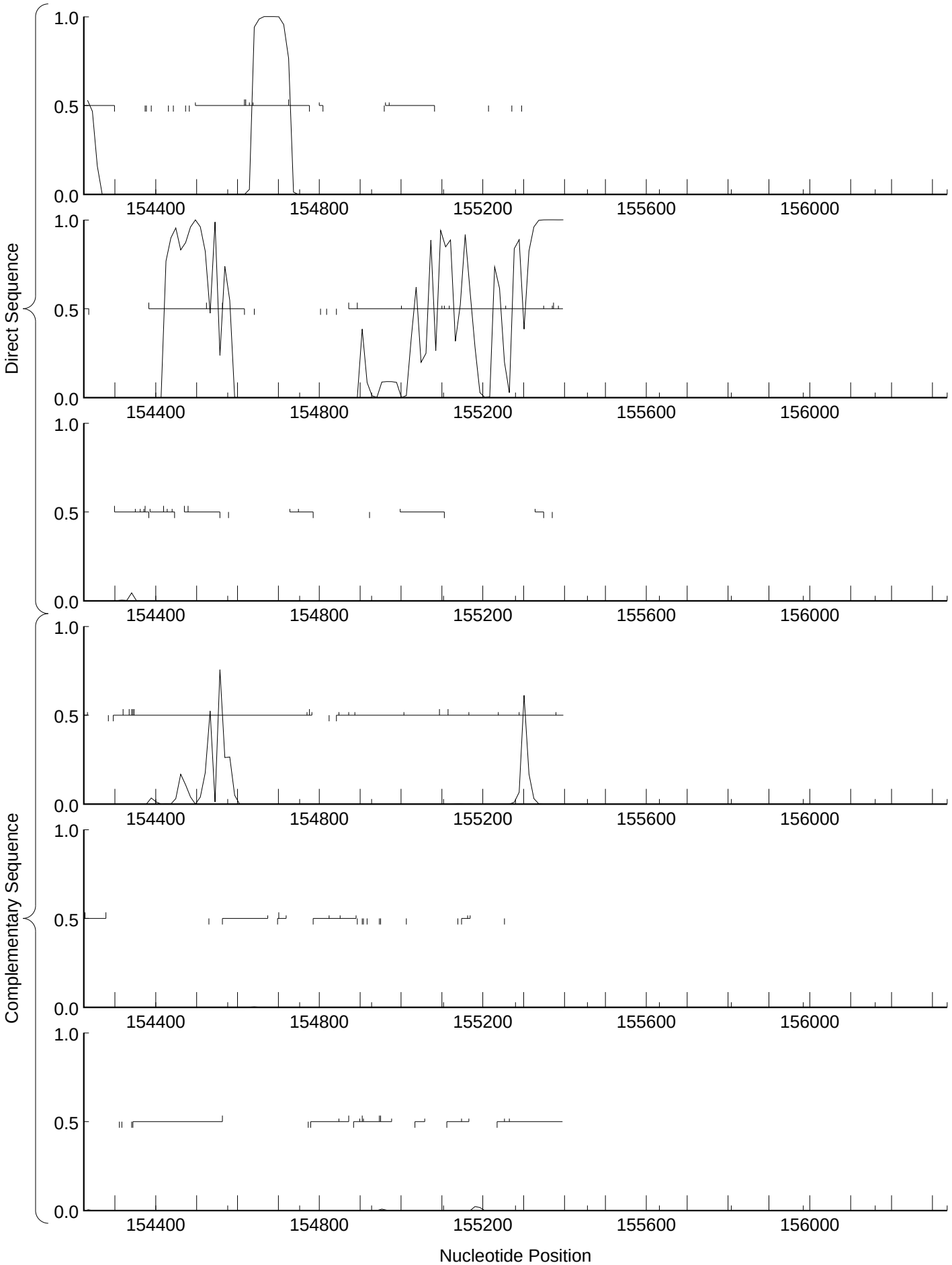

Direct Sequence
Complementary Sequence
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0.5
0.0
144000
144400
144800
145200
145600
Nucleotide Position


Direct Sequence
Complementary Sequence
1.0
0.5
0.0
146000
146400
146800
147200
147600
Nucleotide Position


Direct Sequence
Complementary Sequence
1.0
0.5
0.0
148000
148400
148800
149200
149600
150000
Nucleotide Position


Direct Sequence
Complementary Sequence
1.0
0.5
0.0
150400
150800
151200
151600
152000
Nucleotide Position


Direct Sequence
Complementary Sequence
1.0
0.5
0.0
152400
152800
153200
153600
154000
Nucleotide Position


Direct Sequence
Complementary Sequence
1.0
0.5
0.0
154400
154800
155200
155600
156000
Nucleotide Position