

GeneMark

Version 2.5p (09.08.06)

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PROGRAM INFORMATION

Sequence : Gordonia phage Nodigi complete sequence, 47205 bp including 10-base 3' overhang (CGGTAGGCAT
Analysis Date : 9/21/23 at 22:44:12
Pages : 24
Sequence Length : 47205 bp
GC Content : 63.34%

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,

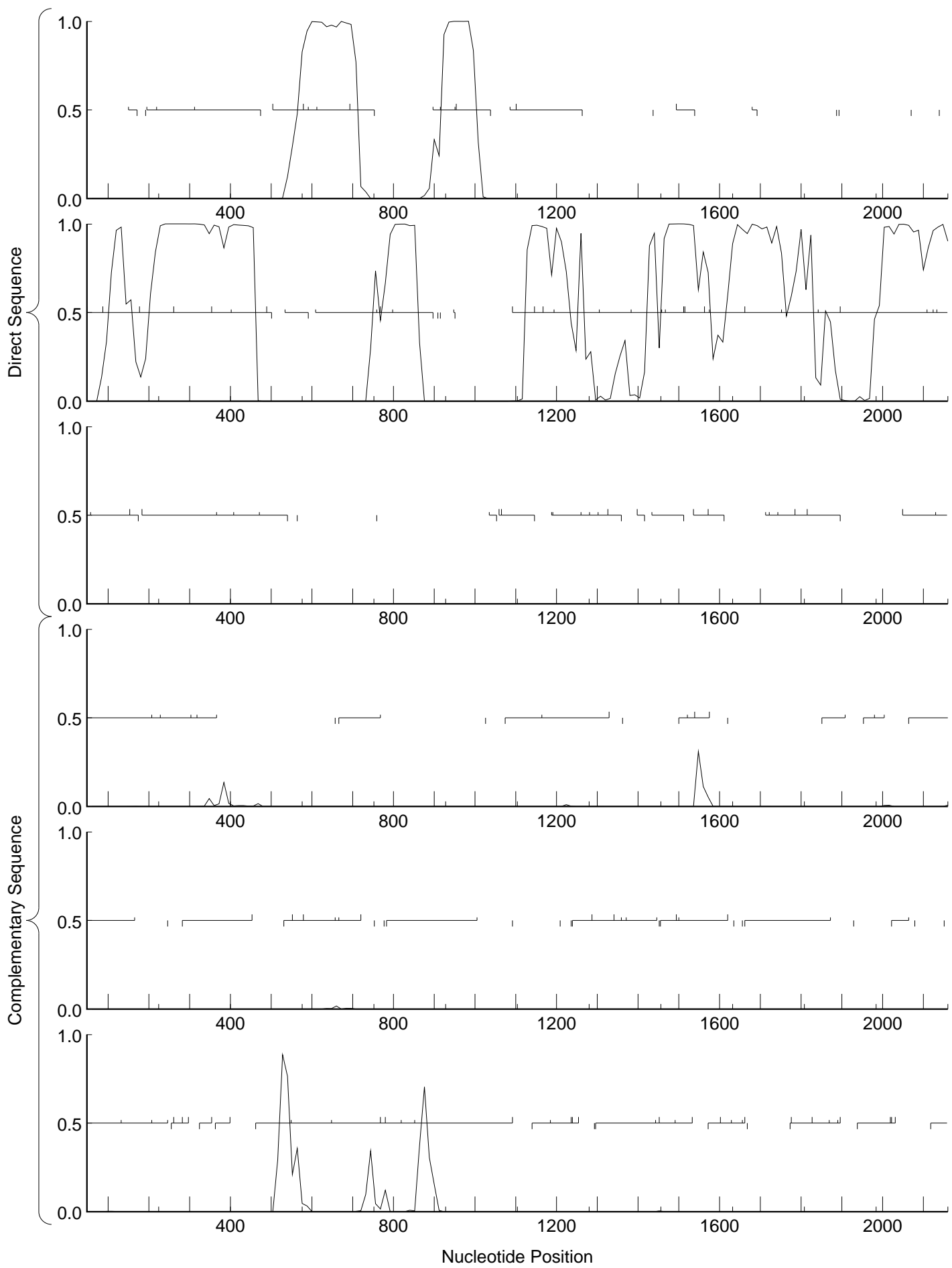
Matrix notes & comments

Training set derived by GeneMarkS, 4.30 October 2014
Sat Sep 8 00:36:14 2018

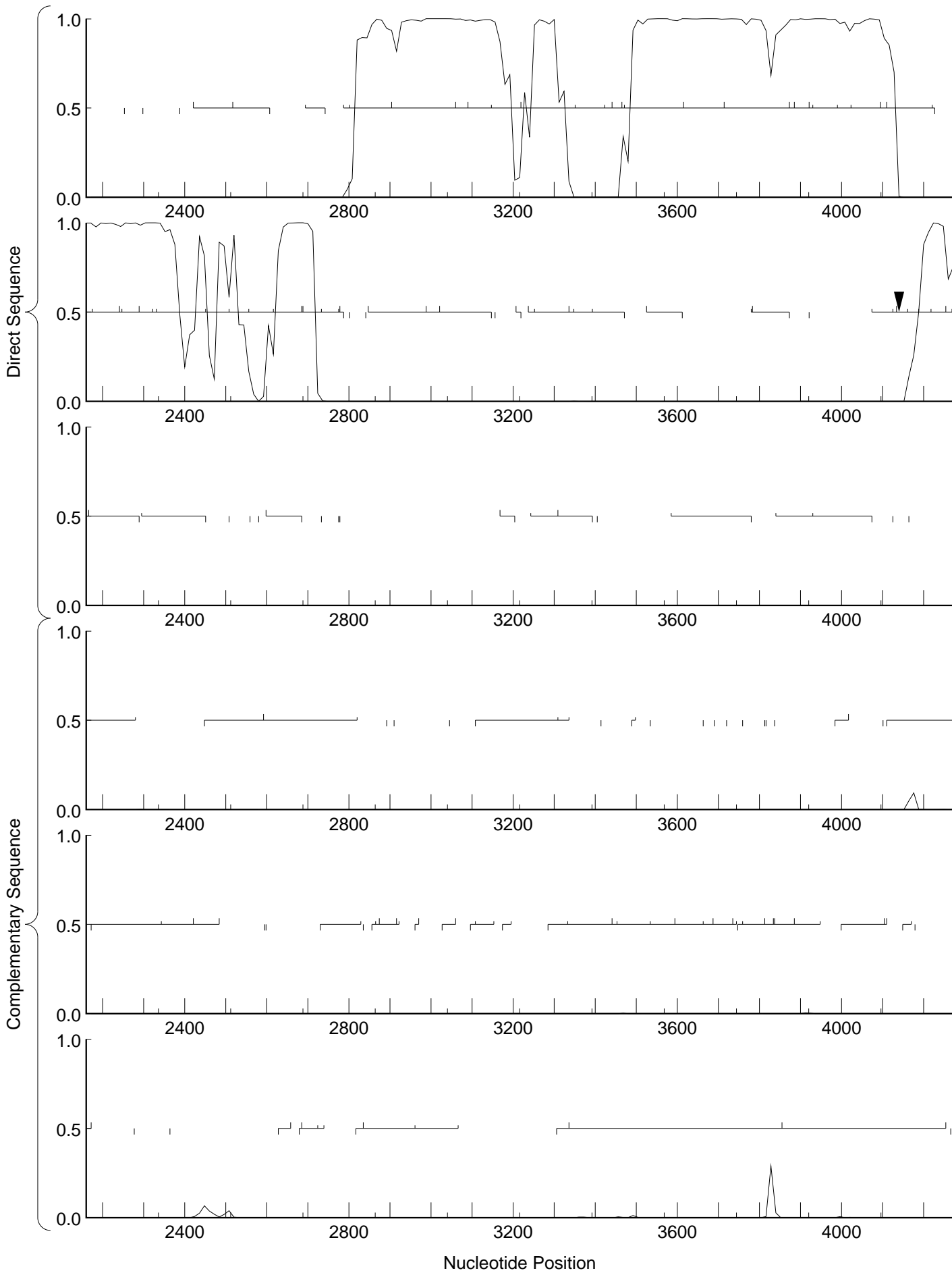
MATRIX INFORMATION

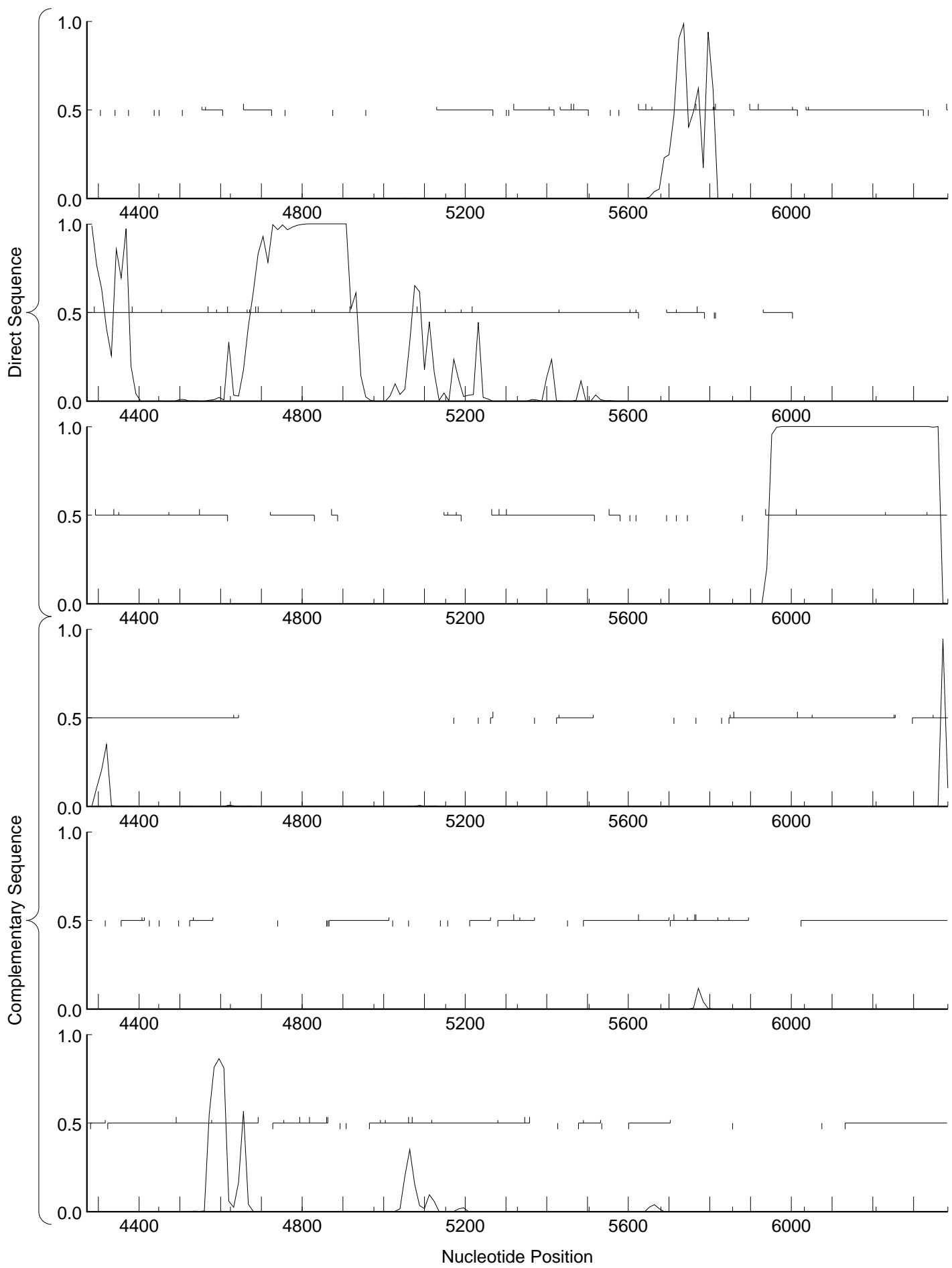
Matrix : -
Author : -
Order : 4

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

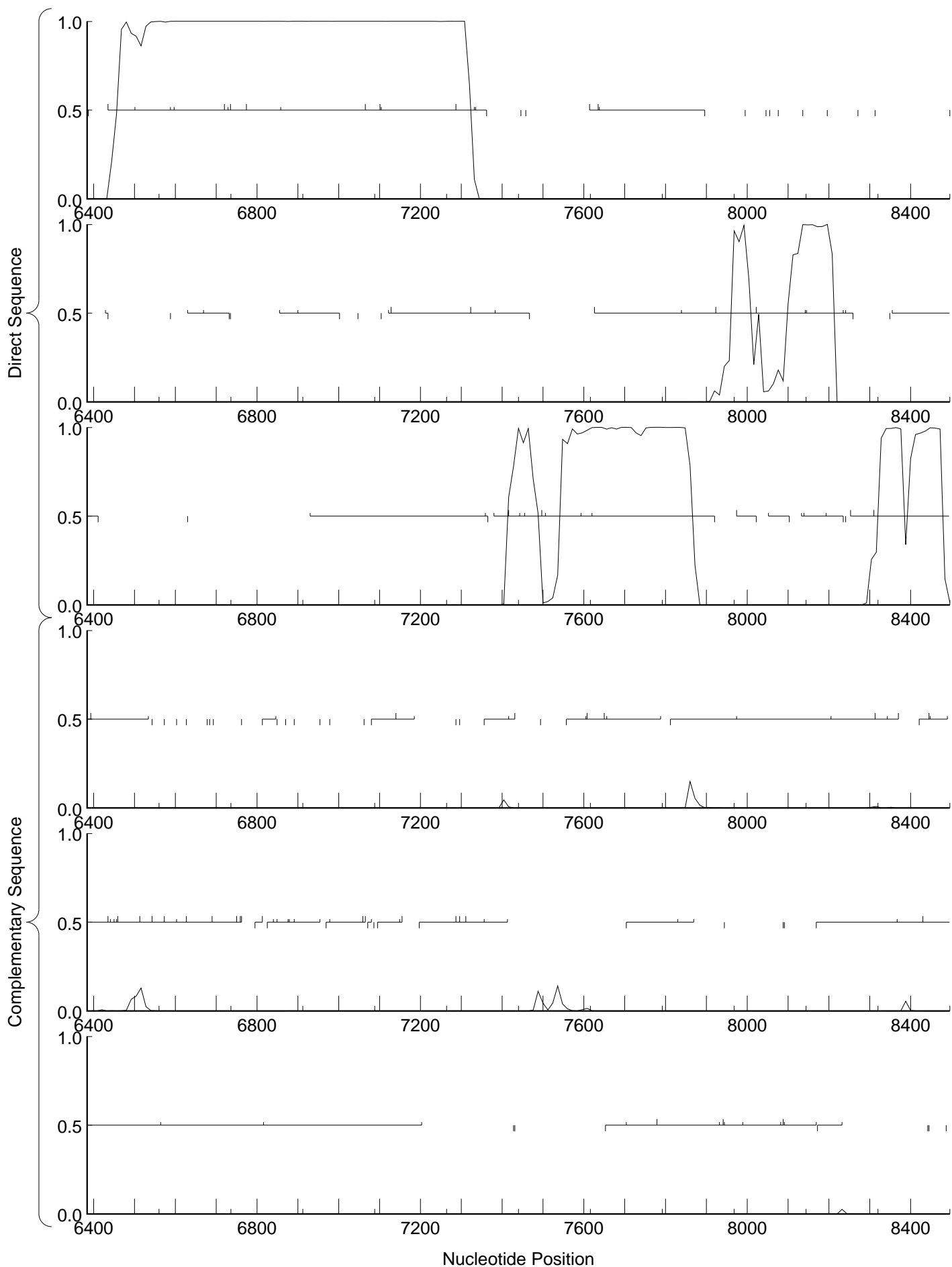


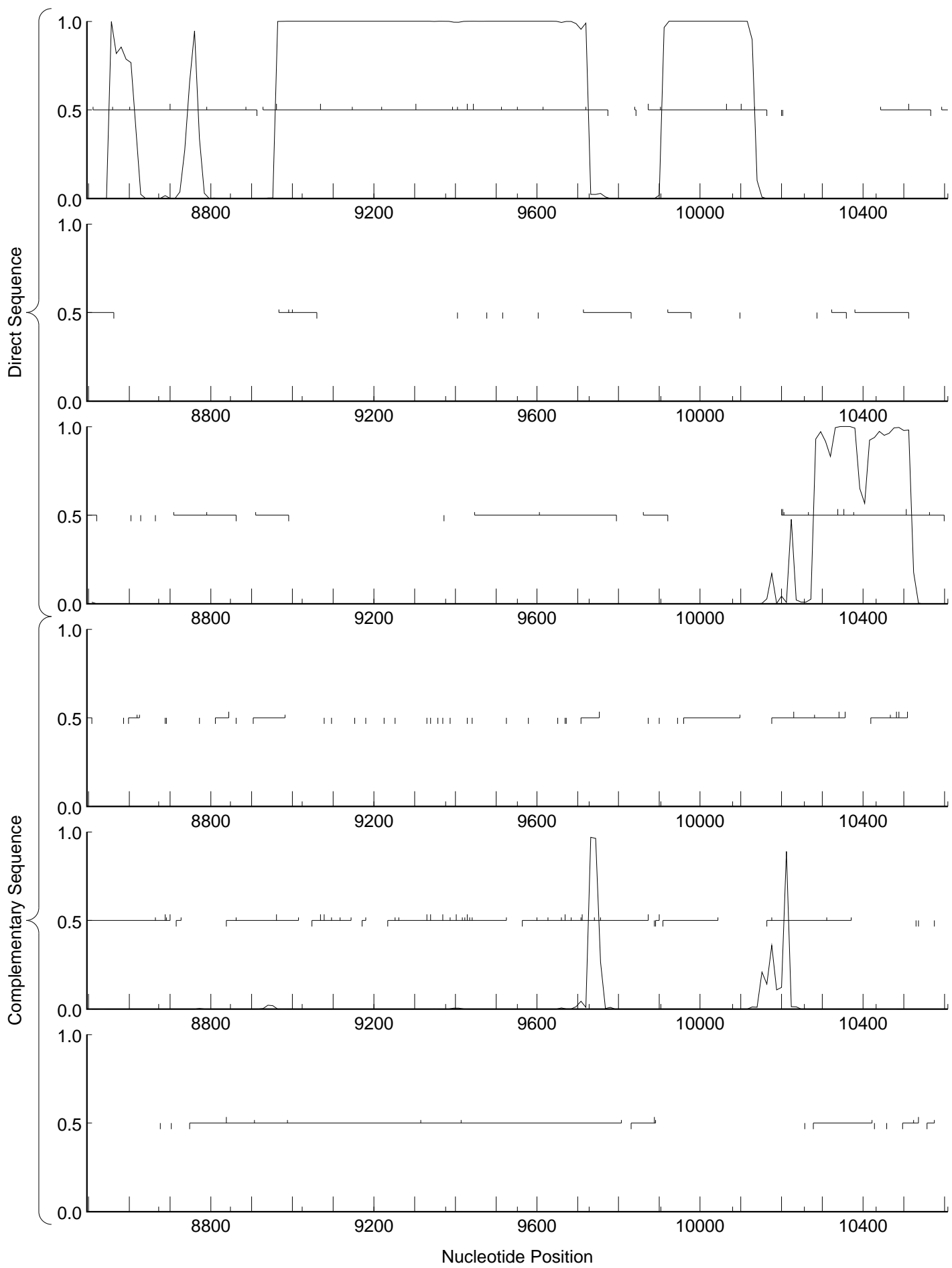
i complete sequence, 47205 bp including 10-base 3' overhang (CGGTAGGCAT), Cluster CT , Order 4, Window 96, Step 12, 3/24



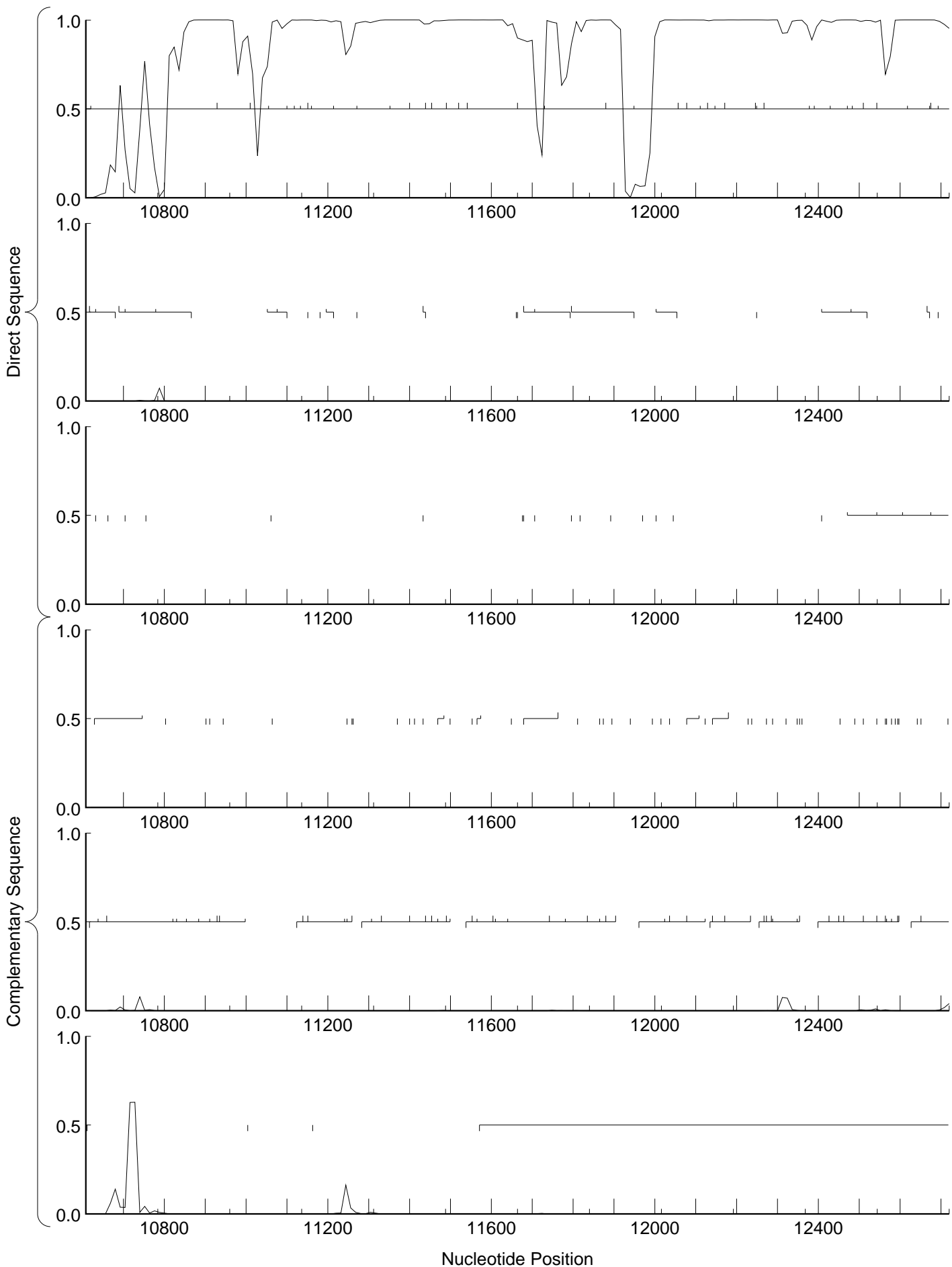


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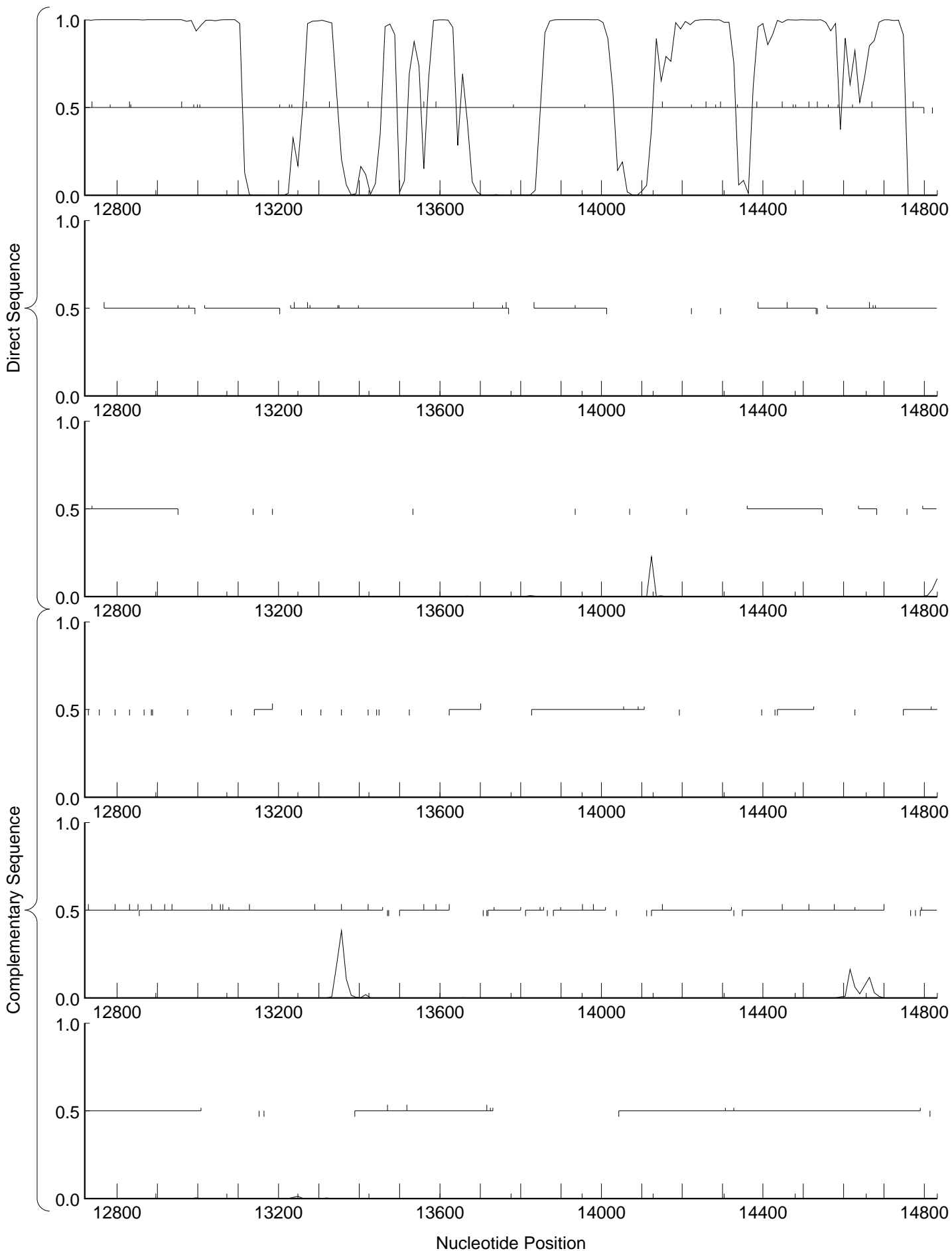


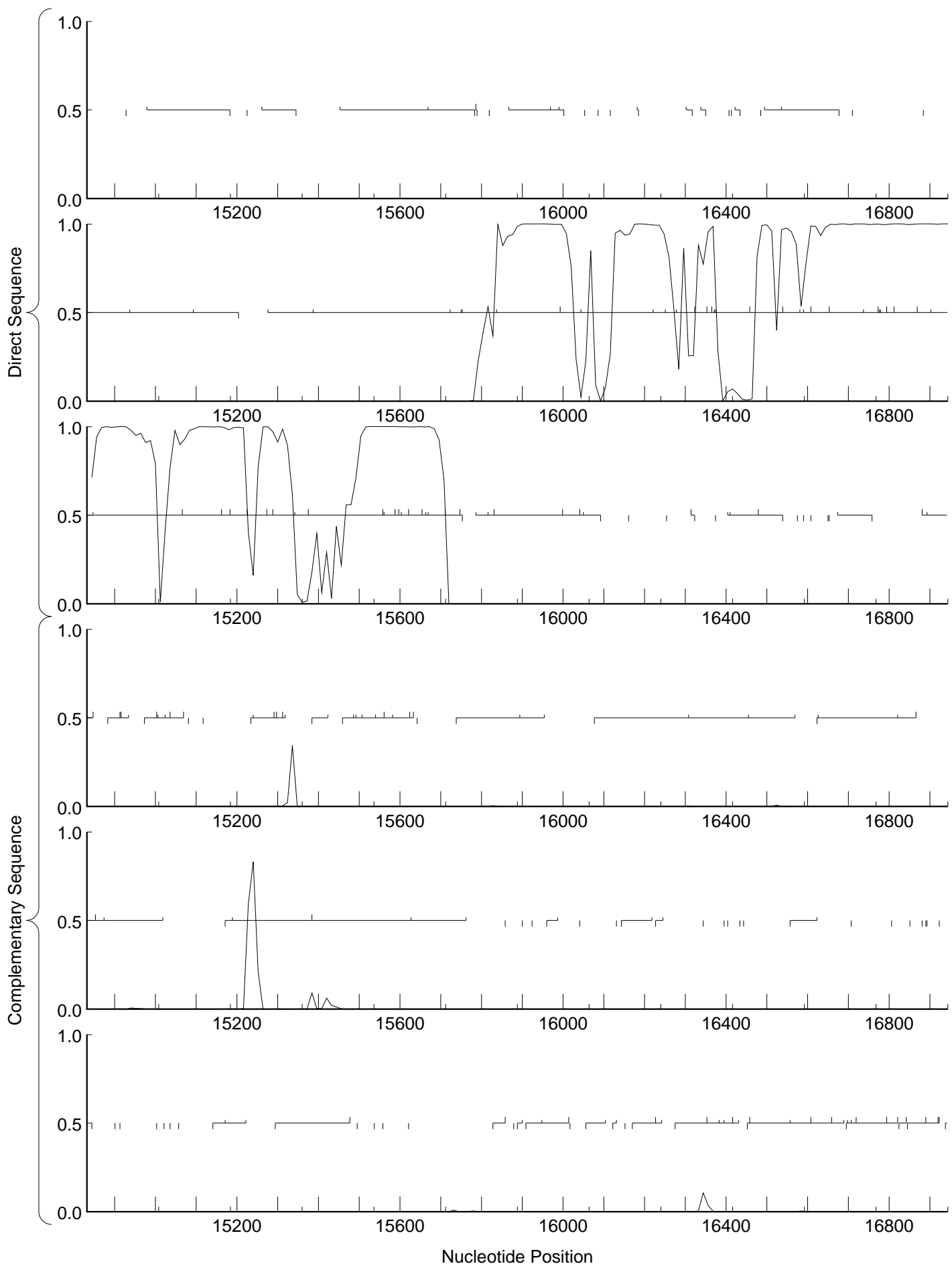


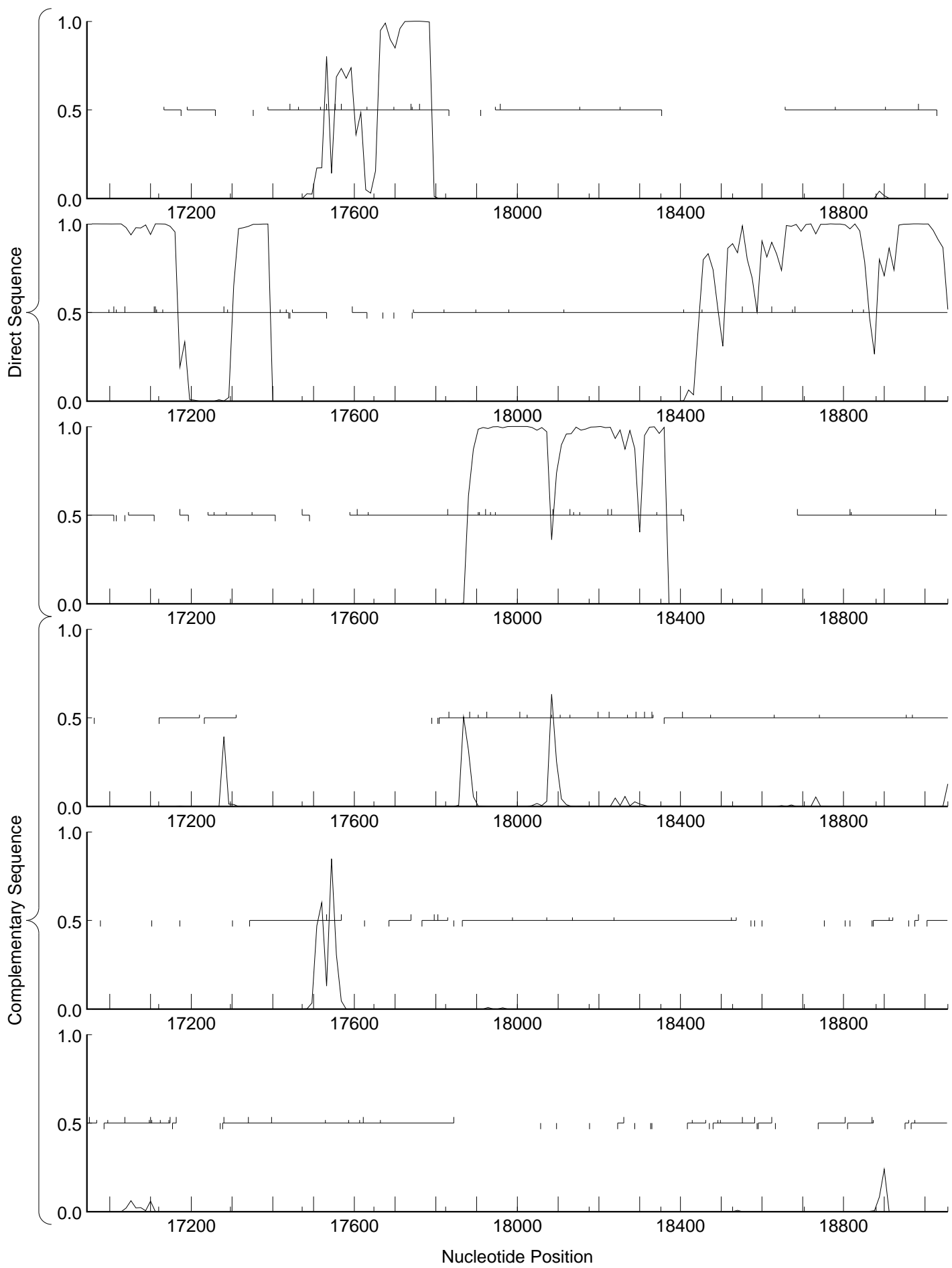
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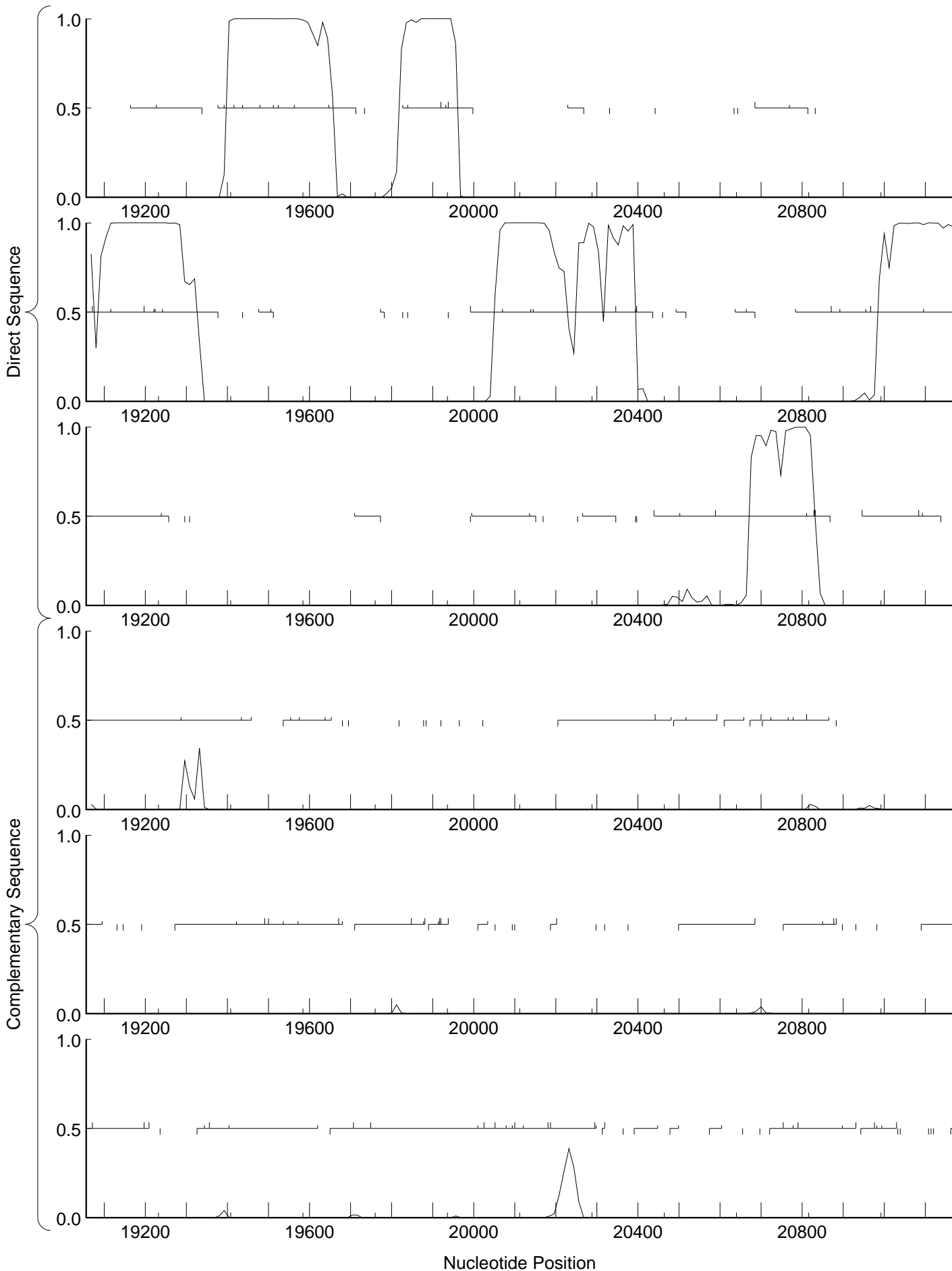


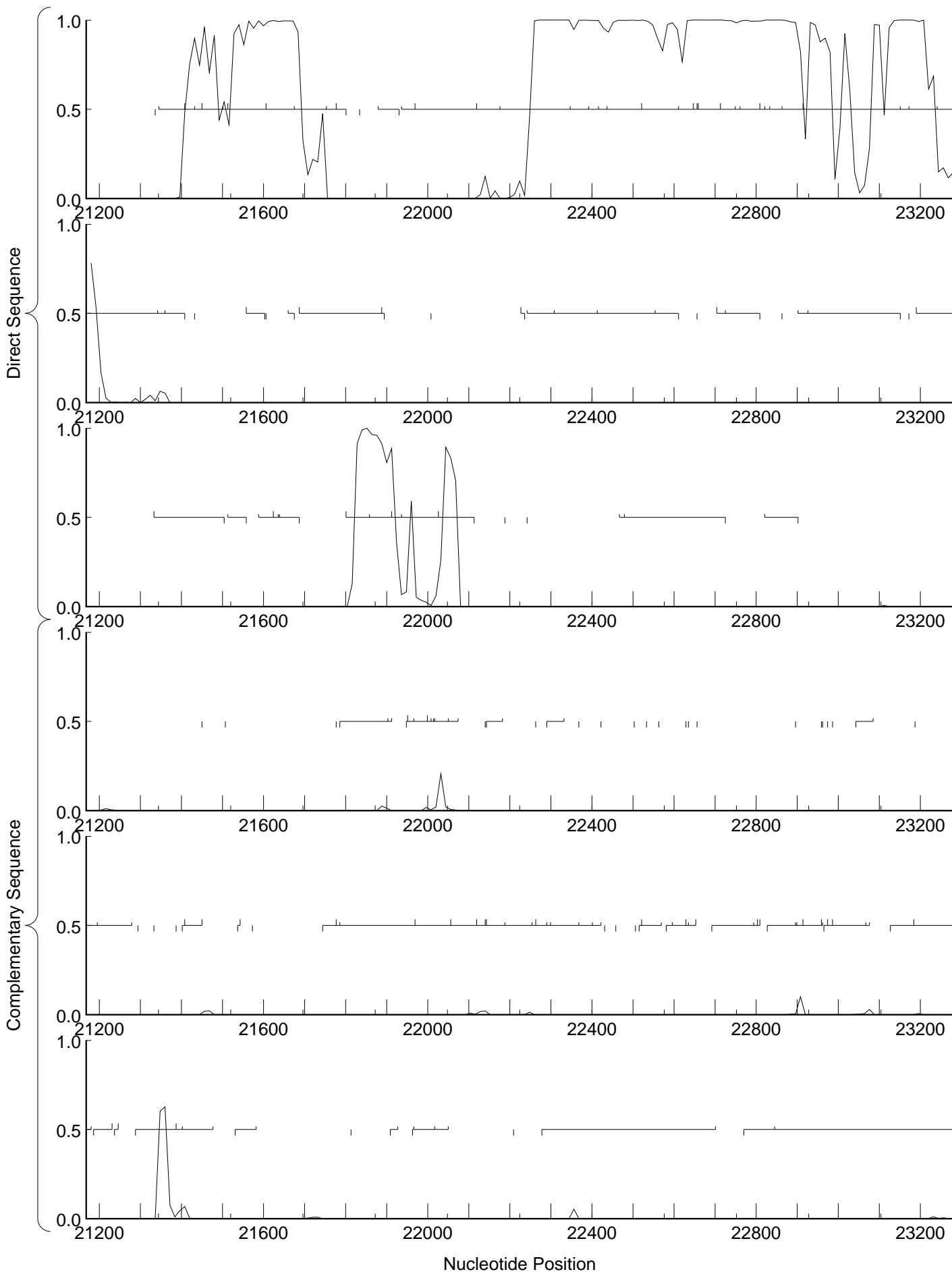
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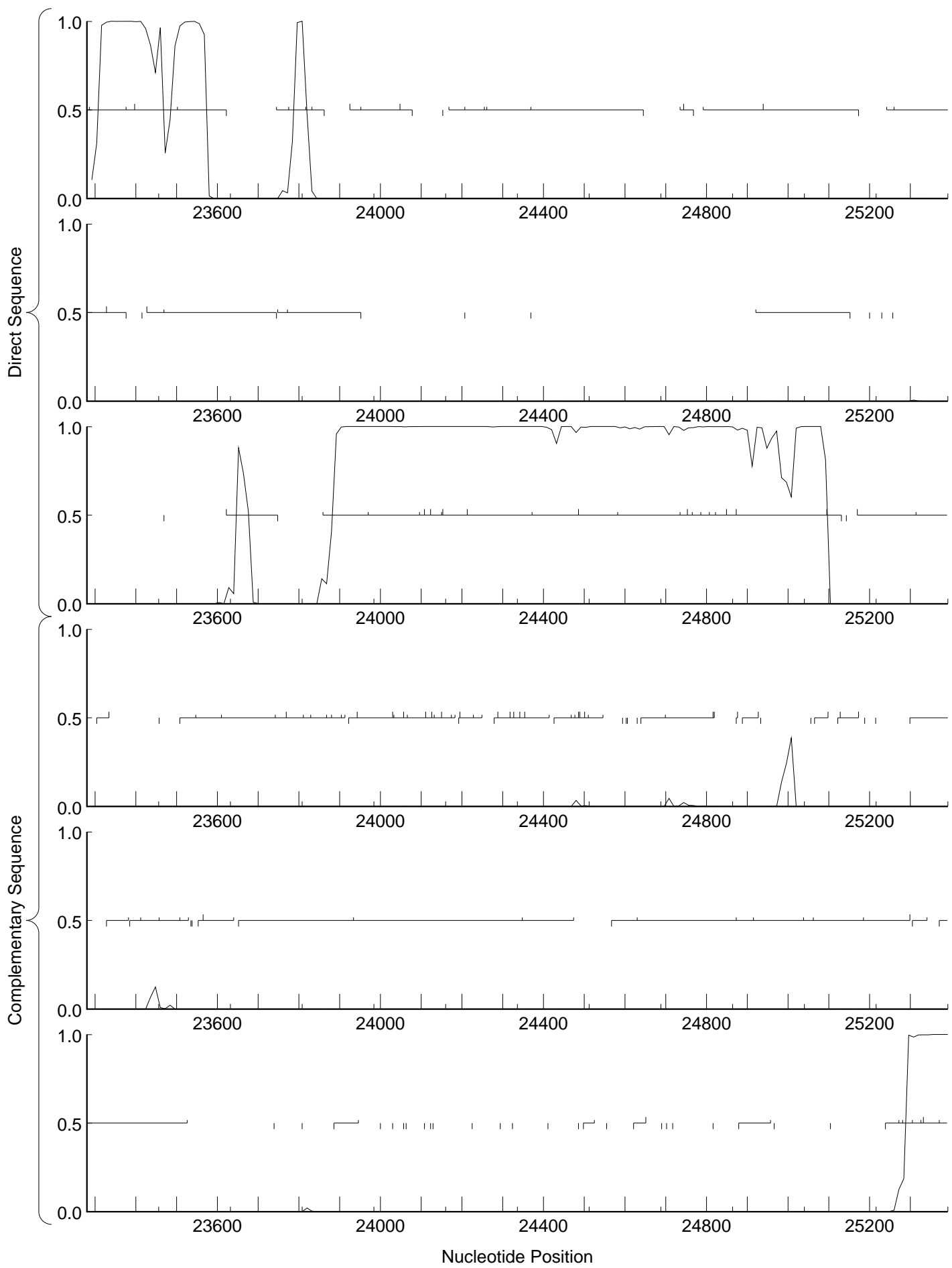




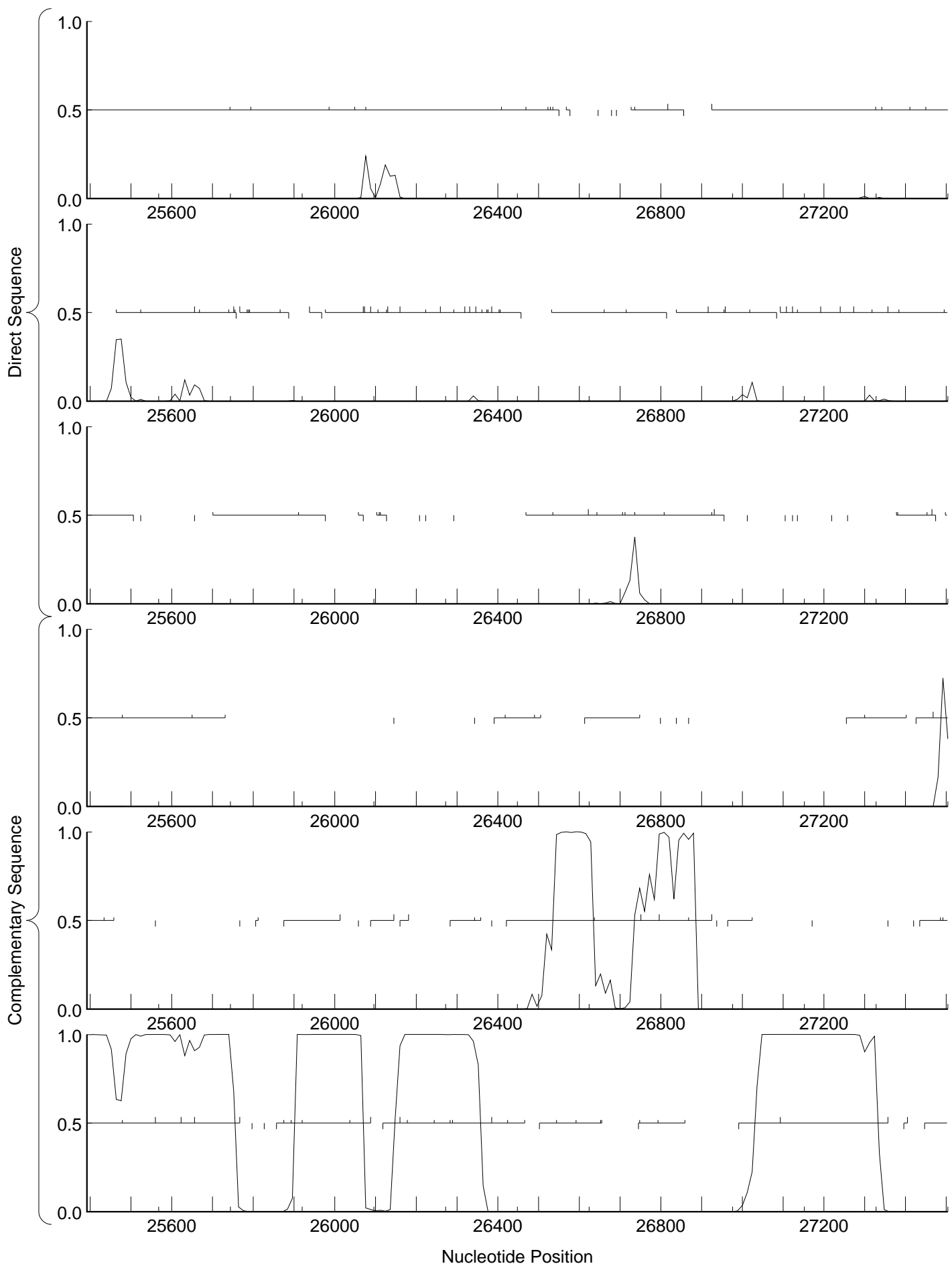




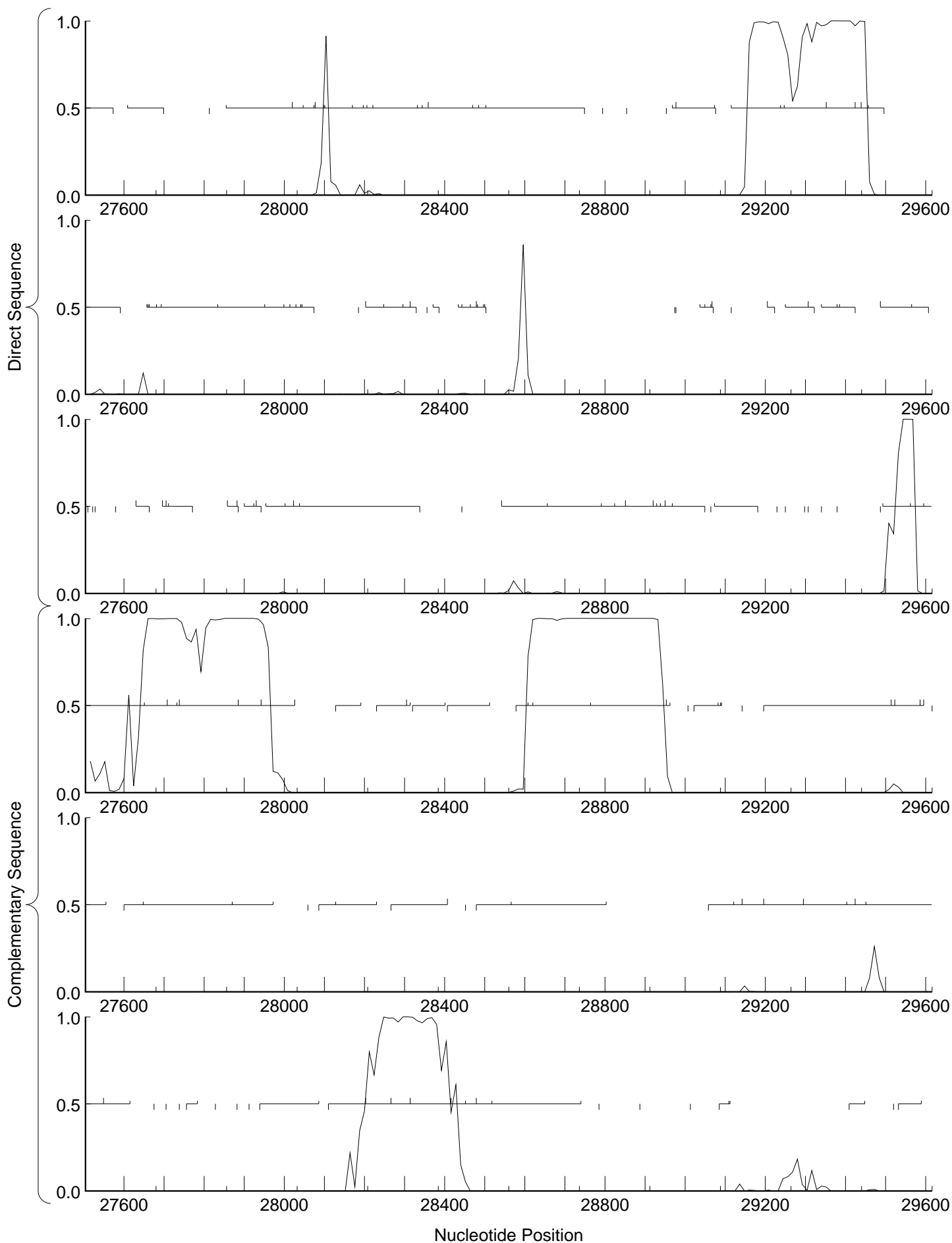
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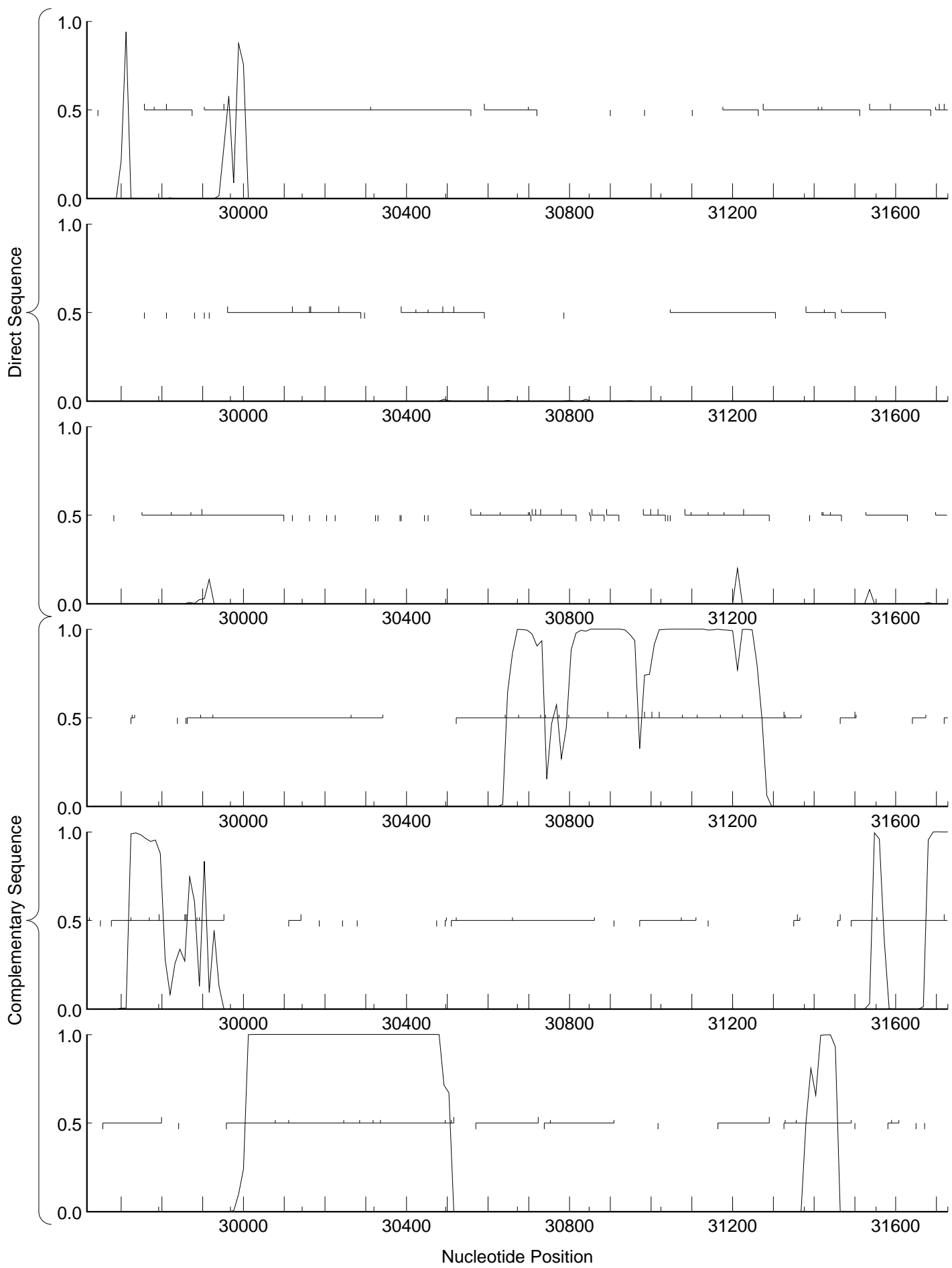
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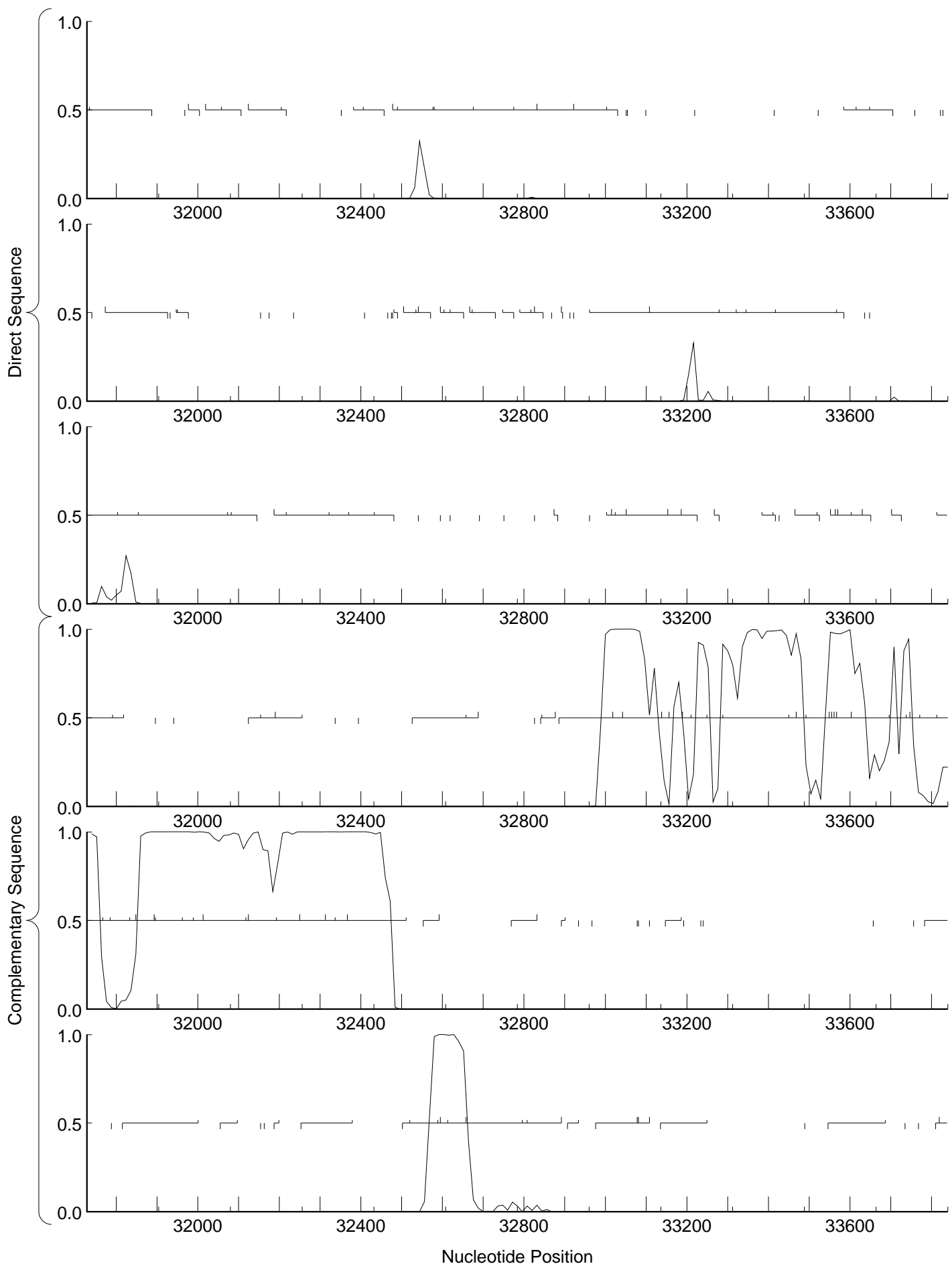
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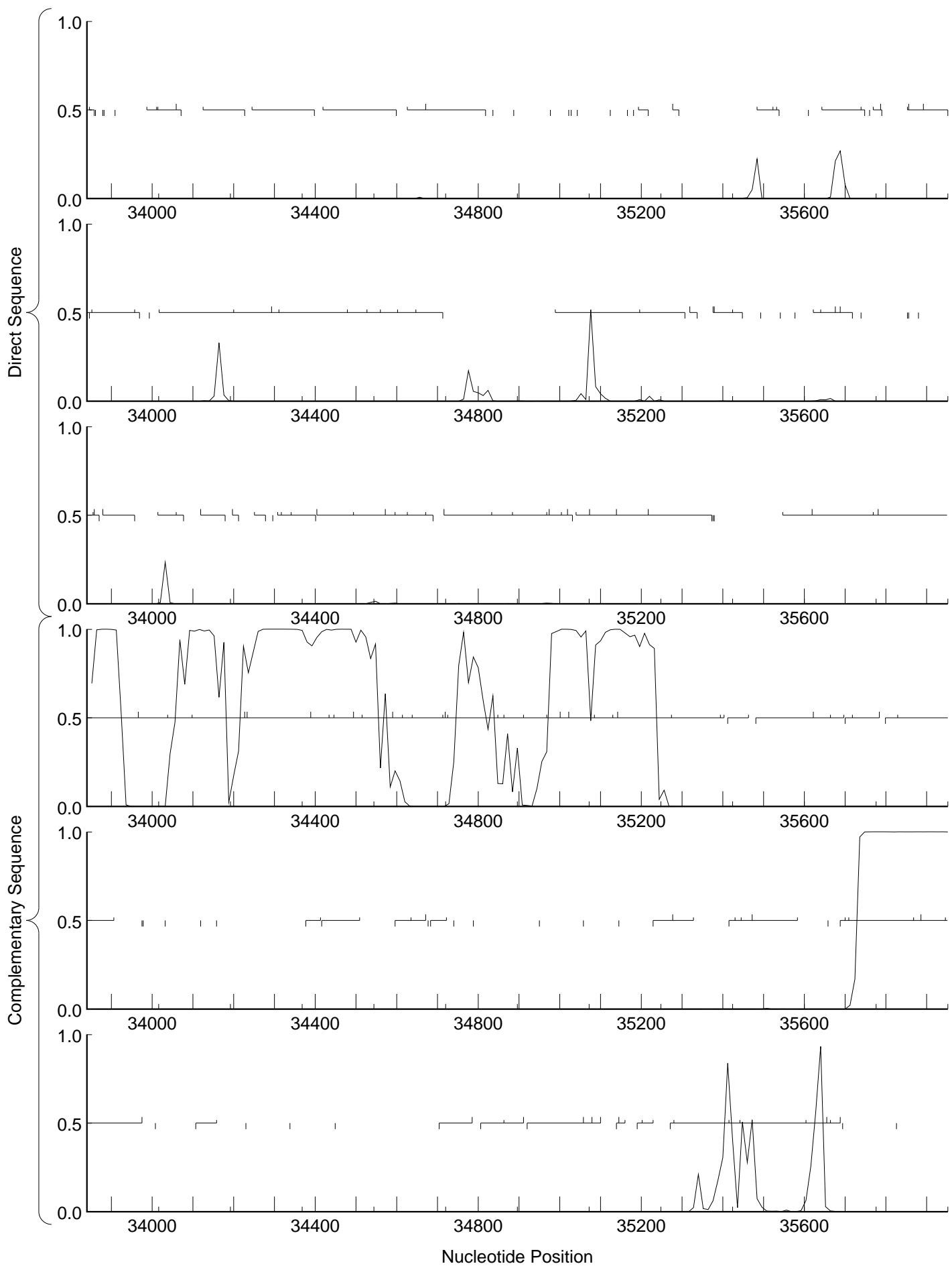


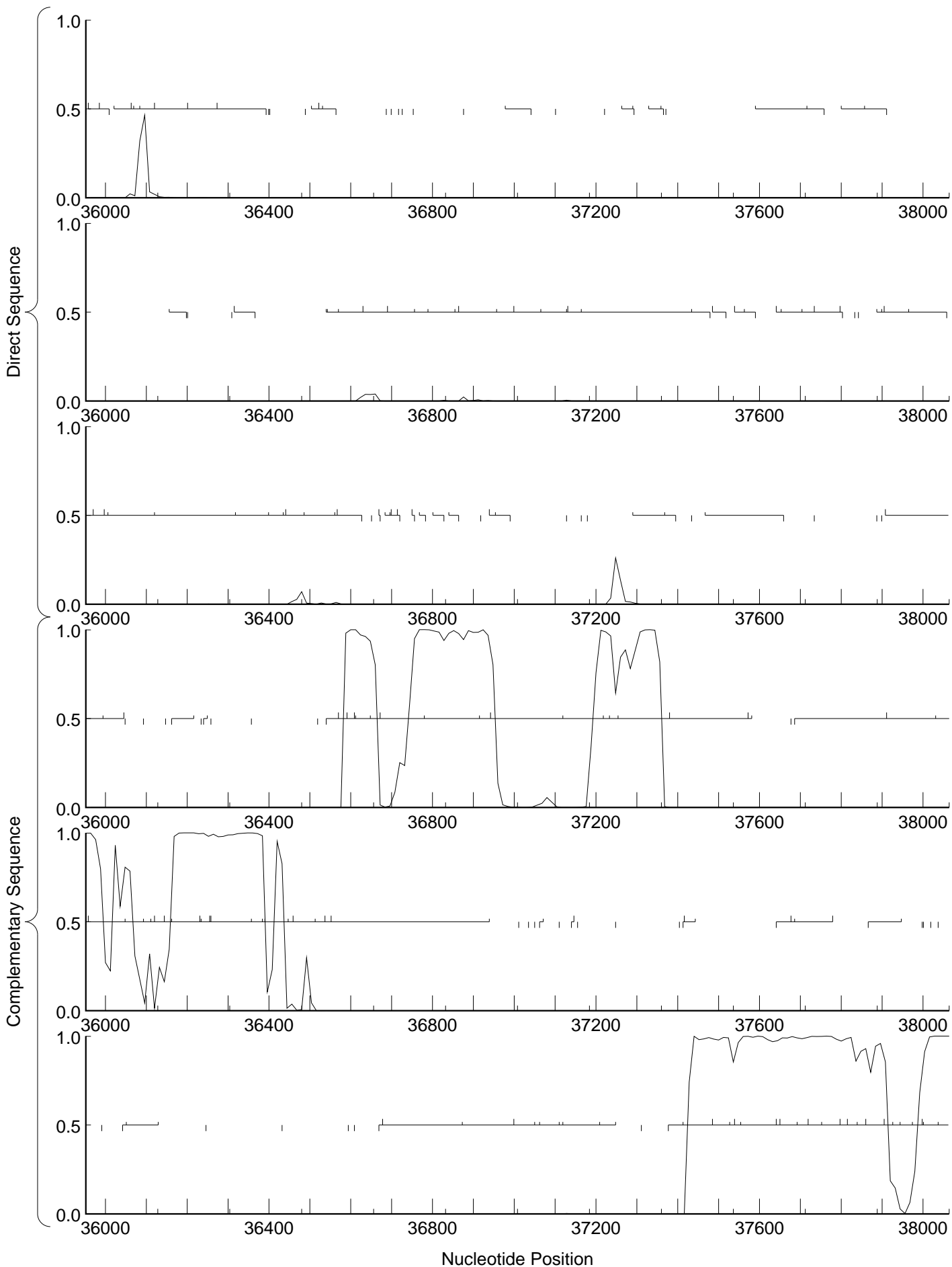
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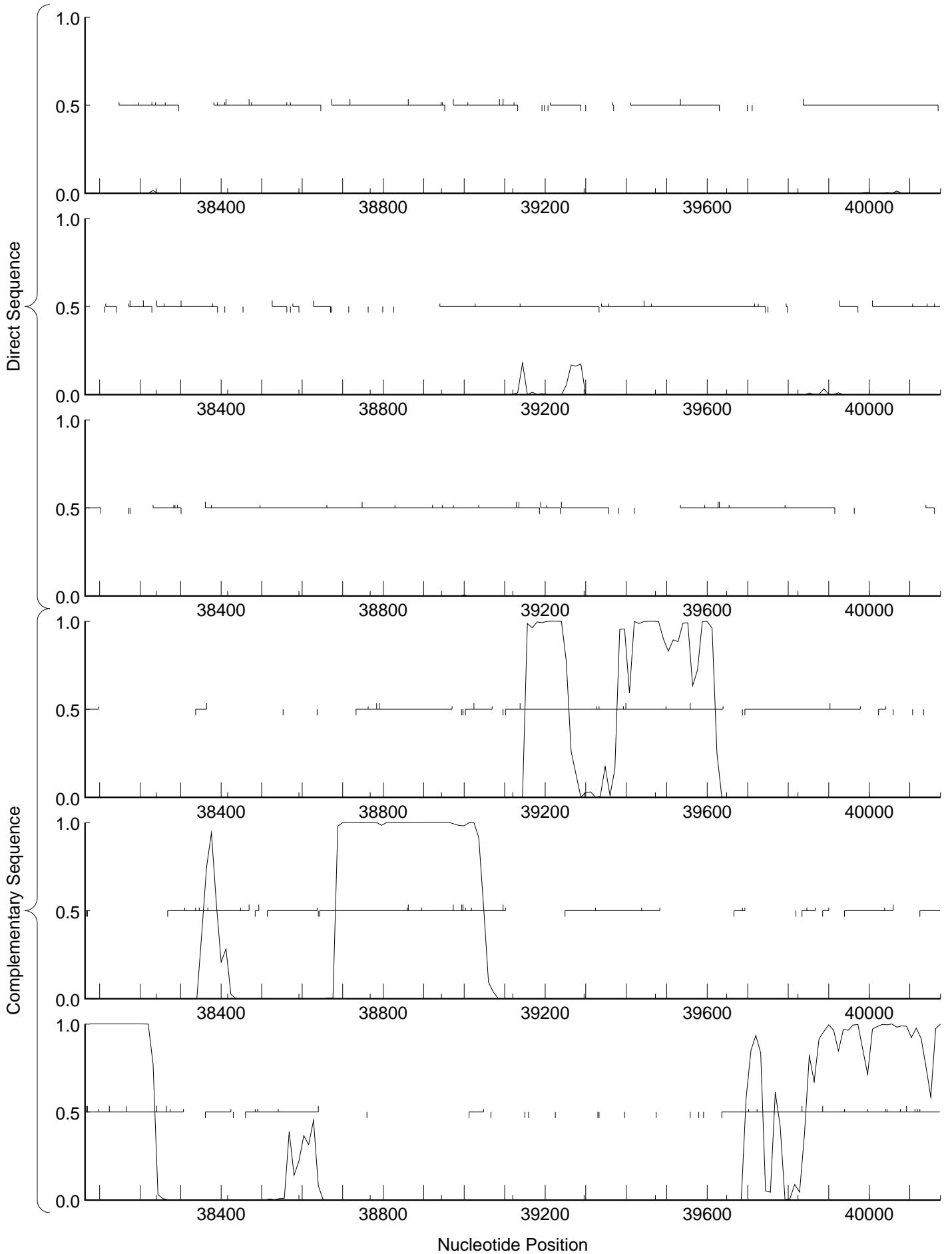


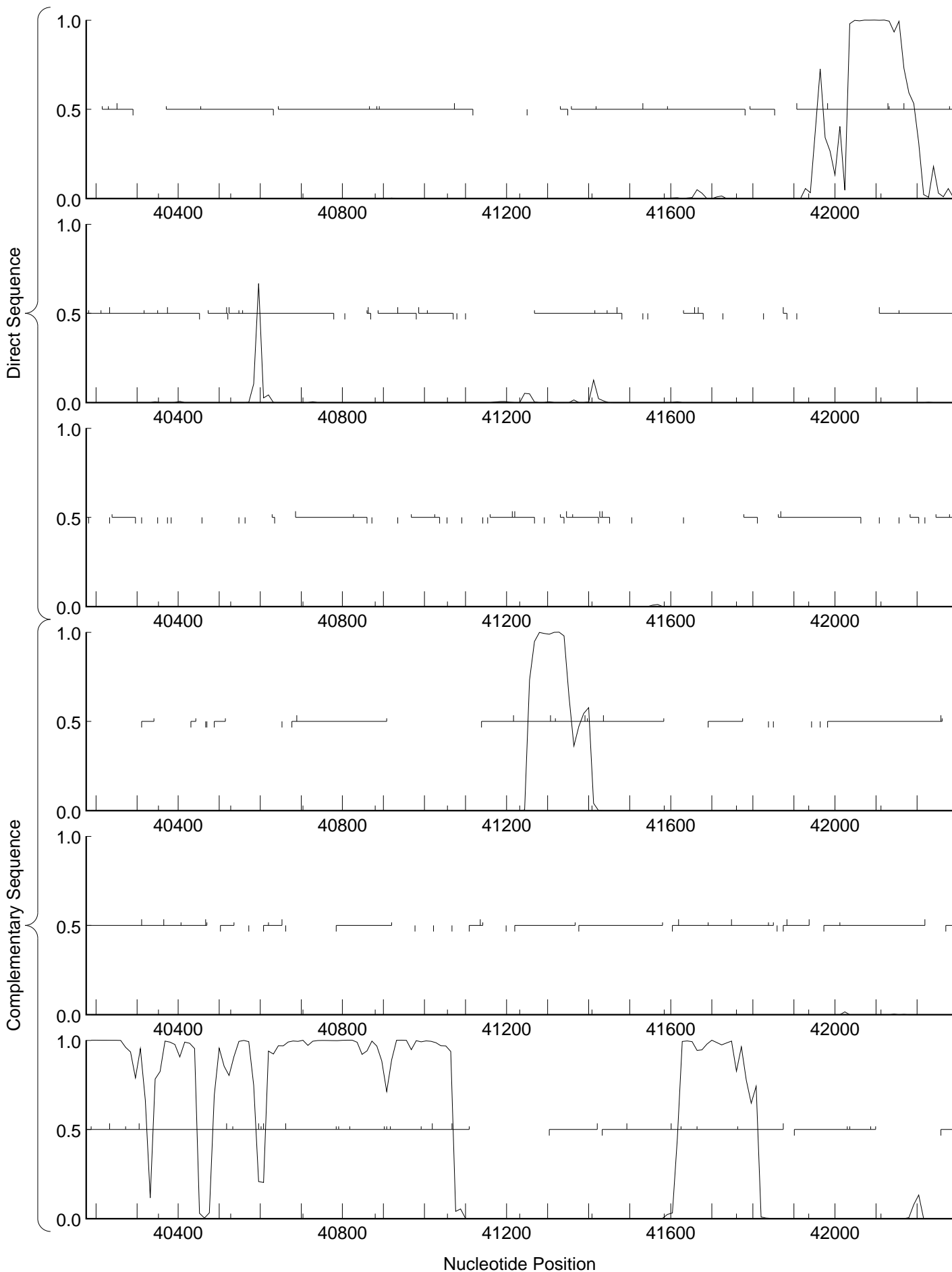
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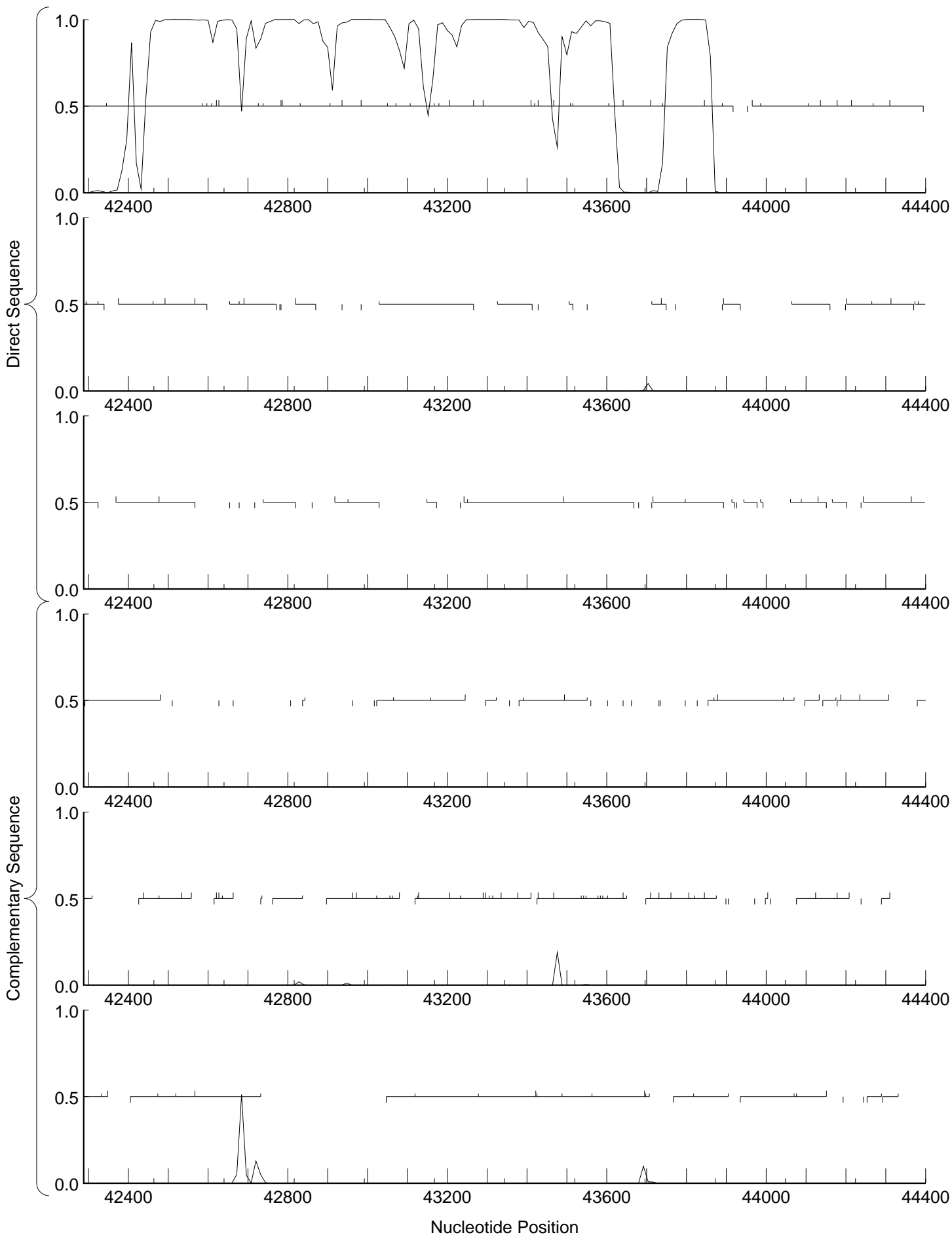








complete sequence, 47205 bp including 10-base 3' overhang (CGGTAGGCAT), Cluster CT , Order 4, Window 96, Step 12, 22/24



complete sequence, 47205 bp including 10-base 3' overhang (CGGTAGGCAT), Cluster CT , Order 4, Window 96, Step 12, 23/24

