Mycobacterium phage Sulley complete sequence, 59873 bp including 11-base 3' overhang (CTCGTAGGCAT), Cluster K1, Order 4, Window 96, Step 12, 2/30
Complementary Sequence Direct Sequence

Nucleotide Position

Mycobacterium phage Sulley complete sequence, 59873 bp including 11-base 3’ overhang (CTCGTAGGCAT), Cluster K1, Order 4, Window 96, Step 12, 3/30
Complementary Sequence Direct Sequence

Nucleotide Position

Mycobacterium phage Sulley complete sequence, 59873 bp including 11-base 3’ overhang (CTCGTAGGCAT), Cluster K1, Order 4, Window 96, Step 12, 4/30
Complementary Sequence  Direct Sequence

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<th>9200</th>
<th>9600</th>
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<td>Complementary Sequence</td>
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Mycobacterium phage Sulley complete sequence, 59873 bp including 11-base 3' overhang (CTCGTAGGCAT), Cluster K1, Order 4, Window 96, Step 12, 6/30
Complementary Sequence Direct Sequence

Nucleotide Position

10800 11200 11600 12000 12400

Mycobacterium phage Sulley complete sequence, 59873 bp including 11-base 3’ overhang (CTCGTAGGCAT), Cluster K1, Order 4, Window 96, Step 12, 7/30
Mycobacterium phage Sulley complete sequence, 59873 bp including 11-base 3' overhang (CTCGTAGGCAT), Cluster K1, Order 4, Window 96, Step 12, 8/30
Mycobacterium phage Sulley complete sequence, 59873 bp including 11-base 3' overhang (CTCGTAGGCAT), Cluster K1, Order 4, Window 96, Step 12, 11/30
Complementary Sequence

Direct Sequence

Mycobacterium phage Sulley complete sequence, 59873 bp including 11-base 3' overhang (CTCGTAGGCAT), Cluster K1, Order 4, Window 96, Step 12, 13/30
Complementary Sequence Direct Sequence

Nucleotide Position

Mycobacterium phage Sulley complete sequence, 59873 bp including 11-base 3' overhang (CTCGTAGGCAT), Cluster K1, Order 4, Window 96, Step 12, 15/30
Complementary Sequence Direct Sequence

Nucleotide Position

Mycobacterium phage Sulley complete sequence, 59873 bp including 11-base 3' overhang (CTCGTAGGCAT), Cluster K1, Order 4, Window 96, Step 12, 16/30
Mycobacterium phage Sulley complete sequence, 59873 bp including 11-base 3' overhang (CTCGTAGGCAT), Cluster K1, Order 4, Window 96, Step 12, 17/30
Mycobacterium phage Sulley complete sequence, 59873 bp including 11-base 3' overhang (CTCGTAGGCAT), Cluster K1, Order 4, Window 96, Step 12, 18/30
Mycobacterium phage Sulley complete sequence, 59873 bp including 11-base 3' overhang (CTCGTAGGCAT), Cluster K1, Order 4, Window 96, Step 12, 19/30
Complementary Sequence Direct Sequence

Nucleotide Position

Mycobacterium phage Sulley complete sequence, 59873 bp including 11-base 3' overhang (CTCGTAGGCAT), Cluster K1, Order 4, Window 96, Step 12, 21/30
Mycobacterium phage Sulley complete sequence, 59873 bp including 11-base 3' overhang (CTCGTAGGCAT), Cluster K1, Order 4, Window 96, Step 12, 22/30
Complementary Sequence Direct Sequence

Nucleotide Position

46800 47200 47600 48000 48400

Mycobacterium phage Sulley complete sequence, 59873 bp including 11-base 3' overhang (CTCGTAGGCAT), Cluster K1, Order 4, Window 96, Step 12, 24/30
Complementary Sequence Direct Sequence

Nucleotide Position

Mycobacterium phage Sulley complete sequence, 59873 bp including 11-base 3' overhang (CTCGTAGGCAT), Cluster K1, Order 4, Window 96, Step 12, 25/30
Mycobacterium phage Sulley complete sequence, 59873 bp including 11-base 3' overhang (CTCGTAGGCAT), Cluster K1, Order 4, Window 96, Step 12, 26/30
Complementary Sequence Direct Sequence

Nucleotide Position

53200 53600 54000 54400 54800

Mycobacterium phage Sulley complete sequence, 59873 bp including 11-base 3' overhang (CTCGTAGGCAT), Cluster K1, Order 4, Window 96, Step 12, 27/30
Mycobacterium phage Sulley complete sequence, 59873 bp including 11-base 3' overhang (CTCGTAGGCAT), Cluster K1, Order 4, Window 96, Step 12, 28/30
Complementary Sequence

Direct Sequence

Nucleotide Position

Mycobacterium phage Sulley complete sequence, 59873 bp including 11-base 3' overhang (CTCGTAGGCAT), Cluster K1, Order 4, Window 96, Step 12, 29/30
Complementary Sequence Direct Sequence

Mycobacterium phage Sulley complete sequence, 59873 bp including 11-base 3' overhang (CTCGTAGGCAT), Cluster K1, Order 4, Window 96, Step 12, 30/30