PROGRAM INFORMATION

Sequence : Microbacterium phage Clayda5 complete sequence, 39894 including 10-base 3’ overhang (ACTCCCGGCA), Cluster EB
Analysis Date : 1/19/22 at 12:28:51
Pages : 20
Sequence Length : 39894 bp
GC Content : 67.16%
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 INFORMATION

Matrix : -
Author : -
Order : 4

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
Complementary Sequence Direct Sequence

Nucleotide Position

Microbacterium phage Clayda5 complete sequence, 39894 including 10-base 3' overhang (ACTCCCGGCA), Cluster EB, Order 4, Window 96, Step 12, 3/20
Microbacterium phage Clayda5 complete sequence, 39894 including 10-base 3’ overhang (ACTCCCGGCA), Cluster EB, Order 4, Window 96, Step 12, 4/20
Microbacterium phage Clayda5 complete sequence, 39894 including 10-base 3' overhang (ACTCCCGGCA), Cluster EB, Order 4, Window 96, Step 12, 5/20
Complementary Sequence

Direct Sequence

Microbacterium phage Clayda5 complete sequence, 39894 including 10-base 3' overhang (ACTCCGGC), Cluster EB, Order 4, Window 96, Step 12, 6/20
Microbacterium phage Clayda5 complete sequence, 39894 including 10-base 3’ overhang (ACTCCCGGCA), Cluster EB, Order 4, Window 96, Step 12, 7/20
Complementary Sequence Direct Sequence

Nucleotide Position

Microbacterium phage Clayda5 complete sequence, 39894 including 10-base 3' overhang (ACTCCGGCA), Cluster EB, Order 4, Window 96, Step 12, 8/20
Microbacterium phage Clayda5 complete sequence, 39894 including 10-base 3' overhang (ACTCCGGCA), Cluster EB, Order 4, Window 96, Step 12, 9/20
Microbacterium phage Clayda5 complete sequence, 39894 including 10-base 3’ overhang (ACTCCGGCA), Cluster EB, Order 4, Window 96, Step 12, 10/20
Microbacterium phage Clayda5 complete sequence, 39894 including 10-base 3’ overhang (ACTCCGGCA), Cluster EB, Order 4, Window 96, Step 12, 11/20
Microbacterium phage Clayda5 complete sequence, 39894 including 10-base 3’ overhang (ACTCCGGCA), Cluster EB, Order 4, Window 96, Step 12, 12/20
Microbacterium phage Clayda5 complete sequence, 39894 including 10-base 3’ overhang (ACTCCGGCA), Cluster EB, Order 4, Window 96, Step 12, 13/20
Microbacterium phage Clayda5 complete sequence, 39894 including 10-base 3’ overhang (ACTCCGGCA), Cluster EB, Order 4, Window 96, Step 12, 14/20.
Microbacterium phage Clayda5 complete sequence, 39894 including 10-base 3’ overhang (ACTCCCGGCA), Cluster EB, Order 4, Window 96, Step 12, 16/20
Microbacterium phage Clayda5 complete sequence, 39894 including 10-base 3' overhang (ACTCCGGCA), Cluster EB, Order 4, Window 96, Step 12, 17/20
Complementary Sequence Direct Sequence

Nucleotide Position

Microbacterium phage Clayda5 complete sequence, 39894 including 10-base 3' overhang (ACTCCGGCA), Cluster EB, Order 4, Window 96, Step 12, 18/20
Microbacterium phage Clayda5 complete sequence, 39894 including 10-base 3' overhang (ACTCCGGCA), Cluster EB, Order 4, Window 96, Step 12, 19/20
Microbacterium phage Clayda5 complete sequence, 39894 including 10-base 3’ overhang (ACTCCGGCA), Cluster EB, Order 4, Window 96, Step 12, 20/20