

GeneMark

Version 2.5p (09.08.06)

Copyright 1993 - M. Borodovsky, J. McIninch

PROGRAM INFORMATION

Sequence : Arthrobacter phage Kaylissa complete sequence, 44124 bp including 11-base 3' overhang (CGAA
Analysis Date : 11/25/20 at 17:07:52
Pages : 22
Sequence Length : 44124 bp
GC Content : 67.62%

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

Matrix notes & comments

Training set derived by GeneMarkS, 4.30 October 2014
Sat Sep 8 00:22:00 2018









































