

GeneMark

Version 2.5p (09.08.06)

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PROGRAM INFORMATION

Sequence : Corynebacterium phage TouchMeNot complete sequence, 40526 bp, circularly permuted, Cluster
Analysis Date : 7/16/18 at 12:29:09
Pages : 21
Sequence Length : 40526 bp
GC Content : 60.03%

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, 4.27 September 2014
Tue Sep 23 15:23:03 2014

MATRIX INFORMATION

Matrix : Mycobacterium_smegmatis_MC2_155
Author : -
Order : 4

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