

Version 2.5p (09.08.06)

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## PROGRAM INFORMATION

Sequence : Gordonia phage LilyPad complete sequence, 64158 bp, circularly permuted, Cluster DG

Analysis Date : 9/7/23 at 7:47:34

Pages : 32

Sequence Length : 64158 bp GC Content : 59.20% 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, 4.33 September 2016

## MATRIX INFORMATION

Matrix : Author : Order : 4

Send questions / comments to: Dr. M. Borodovsky

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