**PROGRAM INFORMATION**

Sequence: Rhodococcus phage DocB7 complete genome, 75772 bp, circularly permuted  
Analysis Date: 4/7/18 at 14:48:52  
Pages: 37  
Sequence Length: 75772 bp  
GC Content: 56.66%  
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: Mycobacterium_tuberculosis_H37Rv  
Author: -  
Order: 4

Send questions / comments to:  
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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
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Rhodococcus phage DocB7 complete genome, 75772 bp, circularly permuted, Order 4, Window 96, Step 12, 9/37
Rhodococcus phage DocB7 complete genome, 75772 bp, circularly permuted, Order 4, Window 96, Step 12, 11/37
Rhodococcus phage DocB7 complete genome, 75772 bp, circularly permuted, Order 4, Window 96, Step 12, 13/37
## Rhodococcus phage DocB7 complete genome, 75772 bp, circularly permuted, Order 4, Window 96, Step 12, 14/37

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