

Title

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Abstract

Author Summary

Introduction

Text. Text. Text. Text. Text. Text. Text. Text. [1] Text. Text. Text. Text. Text. Text. Text.
Text. [2,3]

Results

Subsection 1

Text. Text. Text. Text. Text. Text. Text. Text. [4] might want to know about text text text text Text.
Text. Text. Text. Text. Text. Text. Text. Text. [2,3]

Subsection 2

Discussion

Materials and Methods

Acknowledgments

References

1. Suzuki H, Yano H, Brown CJ, Top EM (2010) Predicting plasmid promiscuity based on genomic signature. J Bacteriol 192: 6045-55.
2. van Passel MW, Kuramae EE, Luyf AC, Bart A, Boekhout T (2006) The reach of the genome signature in prokaryotes. BMC Evol Biol 6: 84.
3. Moreno-Hagelsieb G, Latimer K (2008) Choosing BLAST options for better detection of orthologs as reciprocal best hits. Bioinformatics 24: 319-24.
4. Camacho C, Coulouris G, Avagyan V, Ma N, Papadopoulos J, et al. (2009) BLAST+: architecture and applications. BMC Bioinformatics 10: 421.

Figure Legends

Tables

Table 1. Table title

table information

Table caption