

## Title

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## Abstract

## Author Summary

## Introduction

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Text. [2,3]

## Results

### Subsection 1

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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**