



Bioinformatics
doi.10.1093/bioinformatics/xxxxxx
Advance Access Publication Date: Day Month Year
Manuscript Category

OXFORD

Databases

Tellurium Web - Simulation Service and Database

Matthias König ^{1,*}, Janek Grzegorzewski ¹, Kyle Medley ^{2,}, Kiri Choi ^{2,}, Frank Bergmann ^{3,} and Herbert Sauro ^{2,}

Associate Editor: XXXXXXX

Received on XXXXX: revised on XXXXX: accepted on XXXXX

Abstract

Contact: name@bio.com

Supplementary information: Supplementary data are available at Bioinformatics online.

1 Introduction

$$\sum x + y = Z \tag{1}$$

2 Approach

3 Methods

- for bulleted list, use itemize
- for bulleted list, use itemize
- for bulleted list, use itemize

© The Author 2017. Published by Oxford University Press. All rights reserved. For permissions, please e-mail: journals.permissions@oup.com



¹Department, Institution, City, Post Code, Country and

²Department, Institution, City, Post Code, Country.

^{*}To whom correspondence should be addressed.





2



Sample et al.

3.1 This is subheading

Text Text Text Text Text. Figure 2 shows that the above method Text Text Text Text Text Text. Bauer et al., 2007 might want to know about Bauer et al., 2007 might want to know about text text text text. Text Text Text Text Text Text. Figure 2 shows that the above method Text Text Text Figure 2 shows that the above method Text Text. Bauer et al., 2007 might want to know about Bauer et al., 2007 might want to know about text text text Text Text Text Text Text Text. Figure 2 shows that the above method Text Text Text









Tellurium Web 3

3.1.1 This is subsubheading

Text Text Text Text Text Text. Figure 2 shows that the above method 2007 might want to know about text text text Text Text Text Text Text Text Text. Figure 2 shows that the above method Text Text. Bauer et al., 2007 might want Text Text. Bauer et al., 2007 might want to know about text text text Text Text Text Text Text Text Text. Figure 2 shows that the above method Text Text. Figure 2 shows that the above method Text Text. Bauer et al., 2007 might want to know Text Text. Bauer et al., 2007 might want to know about text text text text Text Text Text Text Text. Figure 2 shows that the above method Text Text.

Text Text Text Text Text Text. Figure 2 shows that the above method Text Text. Figure 2 shows that the above method Text Text. Bauer et al., 2007 might want to know Text Text. Bauer et al., 2007 might want to know about text text text text Text Text Text Text Text. Figure 2 shows that the above method Text









Sample et al.

Table 1. This is table caption			
head1	head2	head3	head4
row1 row2 row3 row4	row1 row2 row3 row4	row1 row2 row3 row4	row1 row2 row3 row4

This is a footnote

FPO

Fig. 1. Caption, caption

3.2 Test1

4 Discussion

5 Conclusion

- 1. this is item, use enumerate
- 2. this is item, use enumerate
- 3. this is item, use enumerate











Tellurium Web 5

Acknowledgements

Text Text Text Text Text Text Text. Bauer et al., 2007 might want to know about text text text text

Funding

This work has been supported by the... Text Text Text Text.

References

Bauer, M. *et al.* (2007). Accurate multiple sequence-structure alignment of RNA sequences using combinatorial optimization. *BMC Bioinformatics* **8** 271

Friedrich, J. *et al.* (2005). ProfDist: a tool for the construction of large phylogenetic trees based on profile distances. *Bionformatics*, **21**(9), 2108–2109.

Jukes, T. and Cantor, C. (1969). Evolution of protein molecules. InH. Munro, editor, *Mammalian Protein Metabolism*, pages 21–132.Academic Press, New York, USA.



