



head1	head2	head3	head4
row1	row1	row1	row1
row2	row2	row2	row2
row3	row3	row3	row3
row4	row4	row4	row4

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## ACKNOWLEDGEMENT

*Funding:* Text Text Text Text Text Text Text Text.

Bauer, M., Klau, G. W., and Reinert, K. (2007). Accurate multiple sequence-structure alignment of RNA sequences using combinatorial optimization. *BMC Bioinformatics*, **8**, 271.

Friedrich, J., Dandekar, T., Wolf, M., and Mller, T. (2005). ProfDist: a tool for the construction of large phylogenetic trees based on profile distances. *Bionformatics*, **21**(9), 2108–2109.