

Correction

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## Ringo – an R/Bioconductor package for analyzing ChIP-chip readouts

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### Correction

Following the publication of the article "Ringo – an R/Bioconductor package for analyzing ChIP-chip readouts – *BMC Bioinformatics* 2007, **8**:221" [1], the submitting author became aware that co-authors had been omitted from the publication.

Therefore, this article has been submitted as a correction to the original text, and highlights the contributions that these authors made to the original article.

The author would like to apologise for any inconvenience this may have caused.

### Corrected author list

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### Corrected authors' contributions

JT, OS and TK developed the software. JT wrote the manuscript. JfF and SS provided example data. All authors contributed to design of analytical algorithms. All authors read and approved the final manuscript.

### References

1. Toedling J, Skylar O, Huber W: **Ringo – an R/Bioconductor package for analyzing ChIP-chip readouts.** *BMC Bioinformatics* 2007, **8**:221.