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Publisher Correction: CNspector: a web-based tool for visualisation and clinical diagnosis of copy number variation from next generation sequencing

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Correction to: *Scientific Reports* <https://doi.org/10.1038/s41598-019-42858-8>, published online 23 April 2019

The original version of this Article contained errors in the Abstract.

“CNspector is written in R and the source code is available for download under the GPL3 Licence from <https://github.com/PapenfussLab/CNspector>. A server running CNspector loaded with the figures from this paper can be accessed at <https://shiny.wehi.edu.au/jmarkham/CNspector/index.html>.”

now reads:

“CNspector is written in R and the source code is available for download under the GPL3 Licence from <https://github.com/PapenfussLab/CNspector>.”

Additionally, the data availability statement contained errors.

“CNspector is written entirely in R and the source code is available for download under the GPL3 Licence from <https://github.com/PapenfussLab/CNspector>. A server running CNspector loaded with the figures from this paper can be accessed at <http://115.146.86.255/CNspector/index.html>. The patient data displayed in the figures is not publicly available due to the presence of potentially genotyping information, but de-identified versions are available from the corresponding author on reasonable request.”

now reads:

“CNspector is written entirely in R and the source code is available for download under the GPL3 Licence from <https://github.com/PapenfussLab/CNspector>. A server running CNspector loaded with the figures from this paper can be accessed at <https://shiny.wehi.edu.au/jmarkham/CNspector/index.html>. The patient data displayed in the figures is not publicly available due to the presence of potentially genotyping information, but de-identified versions are available from the corresponding author on reasonable request.”

These errors have now been corrected in the PDF and HTML versions of the Article.

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