

CORRECTION

# Correction: Identifying Human Genome-Wide CNV, LOH and UPD by Targeted Sequencing of Selected Regions

The *PLOS ONE* Staff

## Notice of Republication

This article was republished on June 12, 2015 to correct the omission of Corresponding Author Yu Wang and his affiliation from the Author byline. The publisher apologizes for the error. Please download this article again to view the correct version. The originally published, uncorrected article and the republished, corrected article are provided here for reference.

## Supporting Information

**S1 File. Originally published, uncorrected article.**  
(PDF)

**S2 File. Republished corrected article.**  
(PDF)

## Reference

1. Wang Y, Li W, Xia Y, Wang C, Tang YT, Guo W, et al. (2015) Identifying Human Genome-Wide CNV, LOH and UPD by Targeted Sequencing of Selected Regions. *PLoS ONE* 10(4): e0123081. doi: [10.1371/journal.pone.0123081](https://doi.org/10.1371/journal.pone.0123081) PMID: [25919136](https://pubmed.ncbi.nlm.nih.gov/25919136/)



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**Citation:** The *PLOS ONE* Staff (2015) Correction: Identifying Human Genome-Wide CNV, LOH and UPD by Targeted Sequencing of Selected Regions. *PLoS ONE* 10(7): e0132537. doi:10.1371/journal.pone.0132537

**Published:** July 2, 2015

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