

Correction: Integrated copy number and expression analysis identifies profiles of whole-arm chromosomal alterations and subgroups with favorable outcome in ovarian clear cell carcinomas

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著者別表示	前田 大地
journal or publication title	PLoS ONE
volume	10
number	7
page range	e0132751
year	2015-07-06
URL	http://doi.org/10.24517/00062700

doi: 10.1371/journal.pone.0132751



Correction: Integrated Copy Number and Expression Analysis Identifies Profiles of Whole-Arm Chromosomal Alterations and Subgroups with Favorable Outcome in Ovarian Clear Cell Carcinomas

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Published: July 6, 2015 • <https://doi.org/10.1371/journal.pone.0132751>

There are errors in the Data Availability statement. The correct statement is: The expression array data are freely available in the GEO website (The accession number is GSE65986 in GEO website (<http://www.ncbi.nlm.nih.gov/geo/query/acc.cgi?acc=GSE65986>)). The SNP array data are available from NBDC (National Bioscience Database Center) website under controlled access (<http://biosciencedbc.jp/en/>). The NBDC number is hum0030, and the JGA (Japanese Genotype-phenotype Archive) accession number is JGA00000000022.

Reference

1. Uehara Y, Oda K, Ikeda Y, Koso T, Tsuji S, Yamamoto S, et al. (2015) Integrated Copy Number and Expression Analysis Identifies Profiles of Whole-Arm Chromosomal Alterations and Subgroups with Favorable Outcome in Ovarian Clear Cell Carcinomas. *PLoS ONE* 10(6): e0128066. pmid:26043110
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Citation: Uehara Y, Oda K, Ikeda Y, Koso T, Tsuji S, Yamamoto S, et al. (2015) Correction: Integrated Copy Number and Expression Analysis Identifies Profiles of Whole-Arm Chromosomal Alterations and Subgroups with Favorable Outcome in Ovarian Clear Cell Carcinomas. *PLoS ONE* 10(7): e0132751. <https://doi.org/10.1371/journal.pone.0132751>

Published: July 6, 2015

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