

CORRECTION

# 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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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 JGA0000000022.

## 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. doi:[10.1371/journal.pone.0128066](https://doi.org/10.1371/journal.pone.0128066) PMID: [26043110](https://pubmed.ncbi.nlm.nih.gov/26043110/)



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