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メタデータ	言語: eng 出版者: 公開日: 2021-06-28 キーワード (Ja): キーワード (En): 作成者: メールアドレス: 所属:
URL	<a href="https://doi.org/10.24517/00062700">https://doi.org/10.24517/00062700</a>

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# 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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