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Integrative Analysis Identifies Four Molecular and Clinical Subsets in Uveal Melanoma

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During the course of revising the manuscript, the authors inadvertently did not include the reference to a paper by Royer-Bertrand et al. in the reference list as well as the corresponding citation in the discussion of copy number variants in uveal melanoma and their consistency in high-risk primary and metastatic tumors.

Royer-Bertrand et al. should have been cited in the second paragraph under the Discussion section as follows: "Prior studies have shown poorer clinical outcomes to be associated with higher chromosome 8q copy number (Royer-Bertrand et al., 2016; Caines et al., 2015; Cassoux et al., 2014; Versluis et al., 2015)."

The reference is now included below and has been added to the online version of the article, along with the citation. The authors apologize for the omission and any confusion it may have caused.

REFERENCES

Royer-Bertrand, B., Torsello, M., Rimoldi, D., El Zaoui, I., Cisarova, K., Pescini-Gobert, R., Raynaud, F., Zografos, L., Schalenbourg, A., Speiser, D., et al. (2016). Comprehensive genetic landscape of uveal melanoma by whole-genome sequencing. Am. J. Hum. Genet. *99*, 1190–1198.

Arid1a Has Context-Dependent Oncogenic and Tumor Suppressor Functions in Liver Cancer

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In the originally published version of this paper, there were two instances in which a paper by Raab et al. was cited, but the full reference was accidentally omitted from the reference list. First, in the subsection titled "Arid1a Haploinsufficiency Alters Global Chromatin Occupancy and Metastasis Genes," paragraph 4, the authors wrote, "To determine if these genes have direct interactions with ARID1A and SWI/SNF components in human HCC, we examined ChIP-seq experiments from HepG2 cells performed by Raab et al." Second, in the legend for Figure 7G, the authors wrote, "ChIP-seq data showing binding of *EMILIN1*, *MAT1A*, *LCN2*, and *IL1R1* loci by ARID1A and SNF5 over input in human HepG2 hepatoma cells (data from Raab et al.)."