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CORRIGENDUM

HMGB1/RAGE Axis Mediates the Apoptosis, Invasion, Autophay, and Angiogenesis of the Renal Cell Carcinoma [Corrigendum]

Wu CZ, Zheng JJ, Bai YH, Xia P, Zhang HC, Guo Y. *Onco Targets Ther*. 2018;11:4501–4510.

Following a review of the data post-publication, the authors found the incorrect images were used for Figure 3A and 3C. The correct images for Figure 3A and 3C are shown in this corrigendum. All the correct images were collected from the second independent experimental group which formed part of the original data. The image backgrounds of this group appear different from the first independent experimental group, shown in the original article, due to the different microscopes used to collect the images (Nikon Ti-s and LEICA DMi1, respectively).

On page 4506, Figure 3A and 3C should be presented as follows:

The authors wish to explain that the analysis of all three data sets was performed prior to the photographs being taken, thus this correction has no impact on the findings of the study.

The authors apologize for this error and any confusion caused.

Received: 2 March 2022
Accepted: 2 March 2022
Published: 9 March 2022

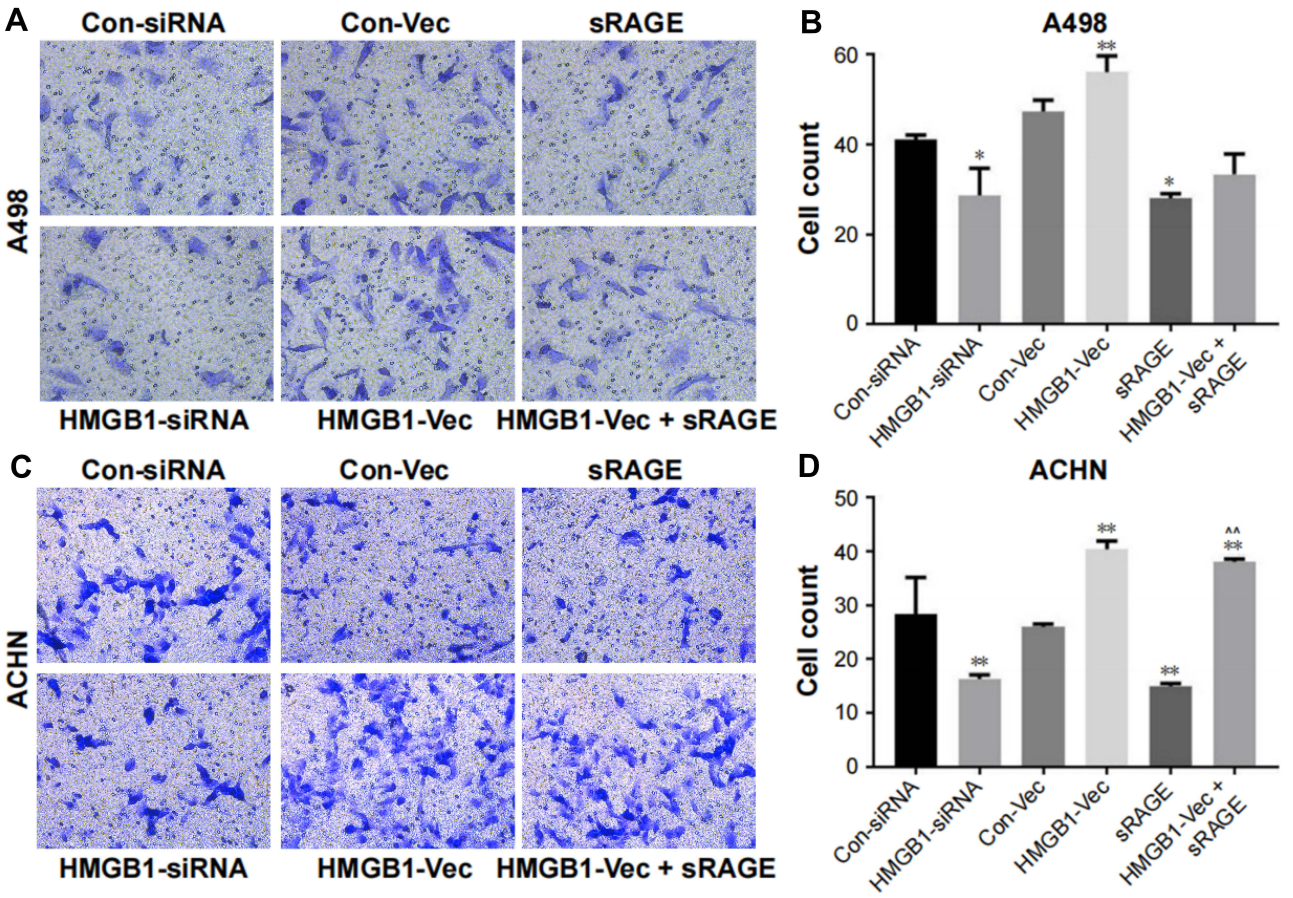

Figure 3 HMGB1 knockdown suppressed the invasion of RCC cells.

Notes: (**A**) The metastatic ability of A498 cells transfected with HMGB1-shRNA, Con-shRNA, and the indicated vectors were analyzed by Matrigel invasion assays. (**B**) The number of A498 cells that invaded through the Matrigel-coated membrane. (**C**) The metastatic ability of ACHN cells transfected with HMGB1-siRNA, Con-siRNA, and the indicated vectors were analyzed by Matrigel invasion assays. (**D**) The number of ACHN cells that invaded through the Matrigel-coated membrane. Data are presented as mean ± SD from 3 independent experiments. **P*<0.05, ***P*<0.01 vs Con-siRNA group; ^^*P*<0.01 vs sRAGE group.

Abbreviations: Con, control; HMGB1, high mobility group box 1 protein; RAGE, receptor for advanced glycation end products; RCC, renal cell carcinoma; siRNA, small interfering RNA.

https://doi.org/10.2147/OTT.S364726