CORRIGENDUM

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Label-free quantitative proteomic analysis identifies CTNNB1 as a direct target of FOXP3 in gastric cancer cells

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Subsequently to the publication of this article, the authors have realized that they omitted to mention that the proteomic raw data alluded to in the above paper were published as supplementary material in one of their previous articles [Pan D, Gao J, Zeng X, Ma G, Li N, Huang X, Du X, Miao Q, Lian J, Xu L et al: Quantitative proteomic analysis reveals up-regulation of caveolin-1 in FOXP3-overexpressed human gastric cancer cells. Sci Rep 7: 14460, 2017]. The MS proteomics data in question were deposited into the ProteomeXchangeConsortium via the PRIDE partner repository with the dataset identifier PXD007725, and are freely available to access there.

The authors apologize for this oversight on their part, and for any inconvenience caused.