CORRIGENDUM

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A proposed method for the relative quantification of levels of circulating microRNAs in the plasma of gastric cancer patients

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Following the publication of this article, we noticed a few inaccuracies, which we would like to correct for the record, as follows:

On p. 3112, in the subsection “Data normalization to evaluate microRNA expression levels”, the word “manually” (“...evaluated manually...”) should not have been included, the word “global” (“...specific global expression profile...”) had been inadvertently omitted, and the citation for Fig. 2 was not in the correct place. The two sentences here should therefore have read (additions highlighted in **bold**):

“Following this normalization strategy, no specific **global** expression profile was observed that would allow the distinction of non-atrophic gastritis from GC, or IGC from DGC. Subsequently, each microRNA was evaluated and normalized against the combination of hsa-miR-18a-5p and hsa-miR-29a-3p (Fig. 2).”

Also, on p. 3113, the title of Fig. 2 should have read as “Differential expression levels of 84 microRNAs in plasma samples from patients with GC or non-atrophic gastritis, normalized using hsa-miR-18a-5p and hsa-miR-29a-3p” (i.e., not as “...normalized using snoRNAs/snRNA panel.”). Associated changes to the text were also required for the figure legend, and therefore Fig. 2 is reproduced here, together with the corrected title and legend. We regret any inconvenience these errors have caused.

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Figure 2. Differential expression levels of 84 microRNAs in plasma samples from patients with GC or non-atrophic gastritis, normalized using hsa-miR-18a-5p and hsa-miR-29a-3p. Of the 84 microRNAs that were evaluated, no specific **global** expression profile was identified that would allow the distinction of non-atrophic gastritis (IC and 2C) from GC samples (1-6GC), or of intestinal-type GC samples (1GC, 3GC and 6GC) from diffuse-type GC samples (2GC, 4GC and 5GC). GC, gastric cancer; min, minimum; avg, average; max, maximum.