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Author Correction: Ascorbic acid during the suckling period is required for proper DNA demethylation in the liver

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This Article contains an error in Figure 3 where the baseline of the graph is shifted upward in panel (b).

The correct Figure 3 appears below as Figure 1.

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Figure 1. DNA methylation status and gene expression levels of PPAR α target genes in livers of KO and N offspring. **(a)** Schematic representation of the promoter region of *Fgf21*. Open circles and gray boxes indicate CpG sites and PPAR response elements (PPREs), respectively. The bisulfite-sequencing (BS) analysis region encompassing the transcription start site (TSS) is indicated. **(b)** *Fgf21* DNA methylation status in the livers of N and KO offspring via the BS analysis (n = 4 per group, statistical analysis using a one-way ANOVA with Tukey's multiple comparison test). Representative BS data are depicted below the graph. Data are expressed as the mean $\pm$ SEM. * $p < 0.05$; ** $p < 0.01$; N.S., not significant vs. KO offspring. **(c)** Hepatic *Cpt1a*, *Fgf21*, *Acaa1a*, *Ehhadh*, and *Peci* mRNA expression levels on d16, which were DNA hypermethylated on both d16 and d100. (n = 6 per group). Statistical analysis using an unpaired Student's *t*-test unless otherwise indicated. Data are expressed as the mean $\pm$ SEM. N.S., not significant vs. KO offspring.

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