CORRECTION

Correction: The newborn metabolome: associations with gestational diabetes, sex, gestation, birth mode, and birth weight

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In this article, the legend for Fig. 2 was inadvertently misrepresented; the figure should appear as shown below. The original article has been corrected.

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Fig. 2 Birth weight z-score and cord blood serum NMR metabolomic profiles. The difference in mean metabolite log concentration in SD units per 1 unit increase in birth weight z-score in adjusted models. Panel (a) is amino acids, ketone bodies, glycerides and phospholipids, glycolysis-related metabolites, and fluid balance markers; (b) is fatty acids, apolipoproteins, cholesterol, lipoprotein particle size, and inflammation (GlycA). Error bars represent 95% confidence intervals. Closed points represent associations with significant p value at the 5% FDR level after adjustment using the Benjamini–Hochberg method. The associations depicted here are listed in Supplementary Table S5. C cholesterol, DHA docosahexaenoic acid, HDL high-density lipoprotein, IDL intermediate density lipoprotein, LA linoleic acid, LDL low-density lipoprotein, MUFA monounsaturated fatty acids, PG phosphoglycerides, PUFA polyunsaturated fatty acids, SFA saturated fatty acids, TG triglycerides, VLDL very-low-density lipoprotein.