Correction to iTRAQ-based Quantitative Protein Expression Profiling and MRM Verification of Markers in Type 2 Diabetes

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On page 5531 of the original article, Table 1 was formatted incorrectly. The correct Table 1 is shown here.

Table 1. Characteristics of T2DM (n = 106) and Nondiabetic Control (n = 76) Volunteers[a]

	control	T2DM
Clinical characteristics		
Age	32.2 ± 1.45	53.6 ± 1.43
BMI(kg/m^2)	28.9 ± 0.90	32.37 ± 0.97
HbA_{1C} (%)	5.57 ± 0.05	7.66 ± 0.06
Vitamin D (ng/mL)	17.5 ± 1.40	12.90* ± 1.34
Glucose (mM)	4.88 ± 0.05	8.71** ± 0.10
Triglycyrides (mM)	1.16 ± 0.77	1.72 ± 0.90
LDL (mM)	2.80 ± 0.05	2.54 ± 0.07
HDL (mM)	4.00 ± 1.23	1.23** ± 0.01
Insulin (pM)	72.2 ± 3.69	91.43* ± 4.52

[a]Serum samples obtained from individual subjects were analyzed to determine the levels of glucose, Vitamin D among others (as described in methods). The value spreads within the group for each parameter are indicated as mean ± standard error. The clinical parameters that had a significant change ($P \leq 0.05$) between the control and T2DM groups are indicated with an asterix. As expected, the diabetic cohort showed an elevation in serum glucose, HbA1c, and insulin levels and a decrease in the serum levels of Vitamin D and HDL.