

Supplementary Materials for

**Widely targeted metabolomics analysis identified characteristic
flavor compounds in the milk from goats, sheep, cows, and buffaloes**

Fuhong Zhang *et al.*

*Corresponding author. Email: luojun@nwafu.edu.cn

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Supplementary Text
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Supplementary Figures

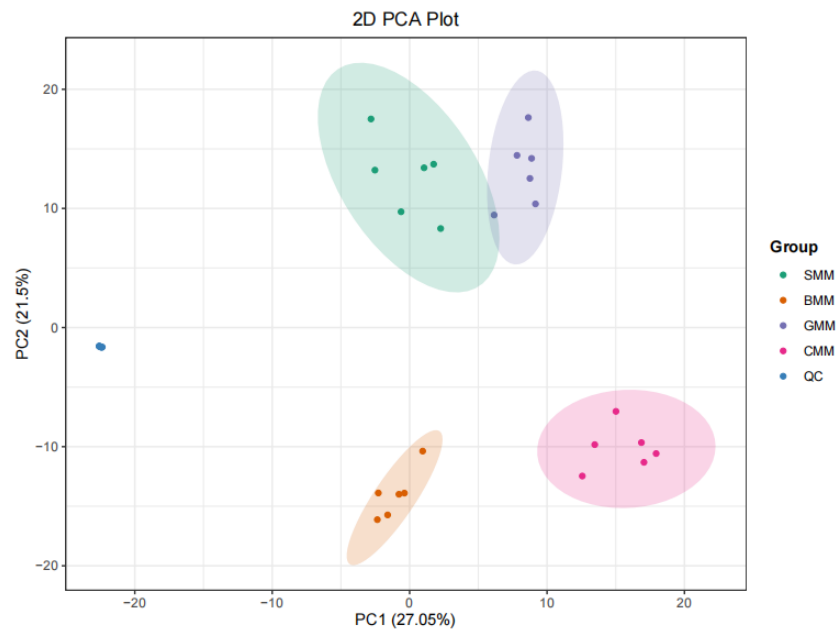


Figure S1. Principal component analysis of the metabolites of four type milk samples. Equal milk samples were mixed and used as quality control (QC).

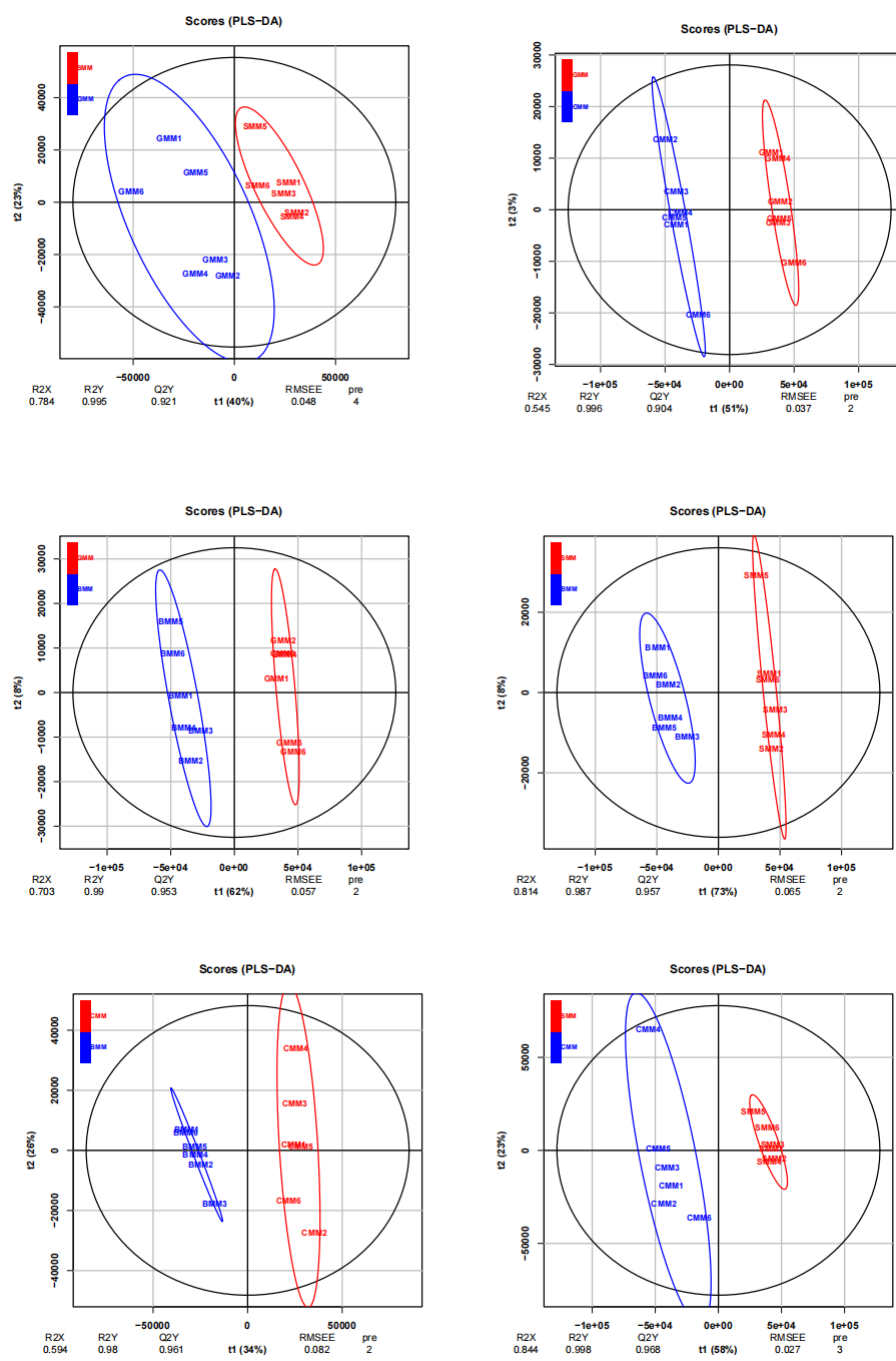


Figure S2. PLS-DA plots exhibiting discernible metabolic disparities among milk samples.

Supplementary Tables

Table S1. A total of 631 metabolites were identified and annotated.

(Excel table)

Table S2. Correlation coefficient between samples.

(Excel table)

Table S3. The results of K-Means cluster analysis.

(Excel table)

Table S4. The results of differential metabolite screening between GMMs and SMMs.

(Excel table)

Table S5. The results of differential metabolite screening between GMMs and CMMs.

(Excel table)

Table S6. The results of differential metabolite screening between GMMs and BMMs.

(Excel table)

Table S7. The results of differential metabolite screening between CMMs and BMMs.

(Excel table)

Table S8. The results of differential metabolite screening between CMMs and SMMs.

(Excel table)

Table S9. The results of differential metabolite screening between SMMs and BMMs.