

Supplementary Material

Non-Targeted Metabolomic Analysis of the Kombucha Production Process

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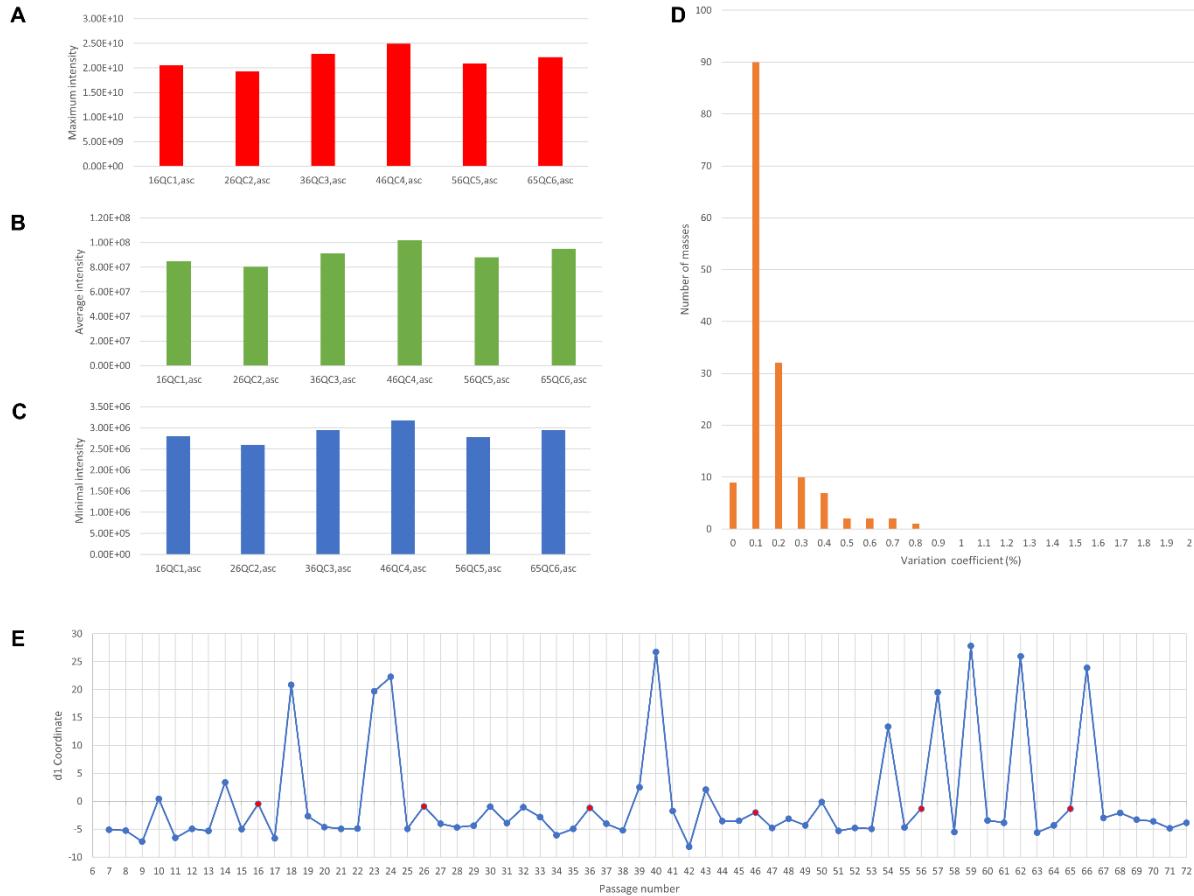


Figure S1: Visualization of Quality control (QC) samples. (A) maximum, (B) average and (C) minimal ion intensities measured in QC samples. (D) Distribution of mass number according to variation coefficient of QC samples. (E) Principal Component Analysis d1 coordinate of samples according to passage number, with QC samples signalized in red.

Table S1: Database annotation of markers

Mass (average)	Formula	Database annotation
169.01427	C ₇ H ₆ O ₅	Gallic acid
173.00917	C ₆ H ₆ O ₆	aconitic acid
179.05612	C ₆ H ₁₂ O ₆	Glucose or Fructose
191.01973	C ₆ H ₈ O ₇	Citric acid
195.05103	C ₆ H ₁₂ O ₇	Gluconic acid
253.21727	C ₁₆ H ₃₀ O ₂	Palmitoleic acid
255.23292	C ₁₆ H ₃₂ O ₂	Palmitic acid
273.07682	C ₁₅ H ₁₄ O ₅	Epiafzelechin
281.24859	C ₁₈ H ₃₄ O ₂	Oleic acid
283.26424	C ₁₈ H ₃₆ O ₂	Stearic acid
289.07175	C ₁₅ H ₁₄ O ₆	Epicatechin
305.06669	C ₁₅ H ₁₄ O ₇	Epigallocatechin
341.10891	C ₁₂ H ₂₂ O ₁₁	Sucrose
425.08788	C ₂₂ H ₁₈ O ₉	Epiafzelechin gallate
441.08278	C ₂₂ H ₁₈ O ₁₀	Epicatechin gallate
503.16189	C ₁₈ H ₃₂ O ₁₆	Dextrin
535.15180	C ₁₈ H ₃₂ O ₁₈	1,4-bêta-D-Glucan

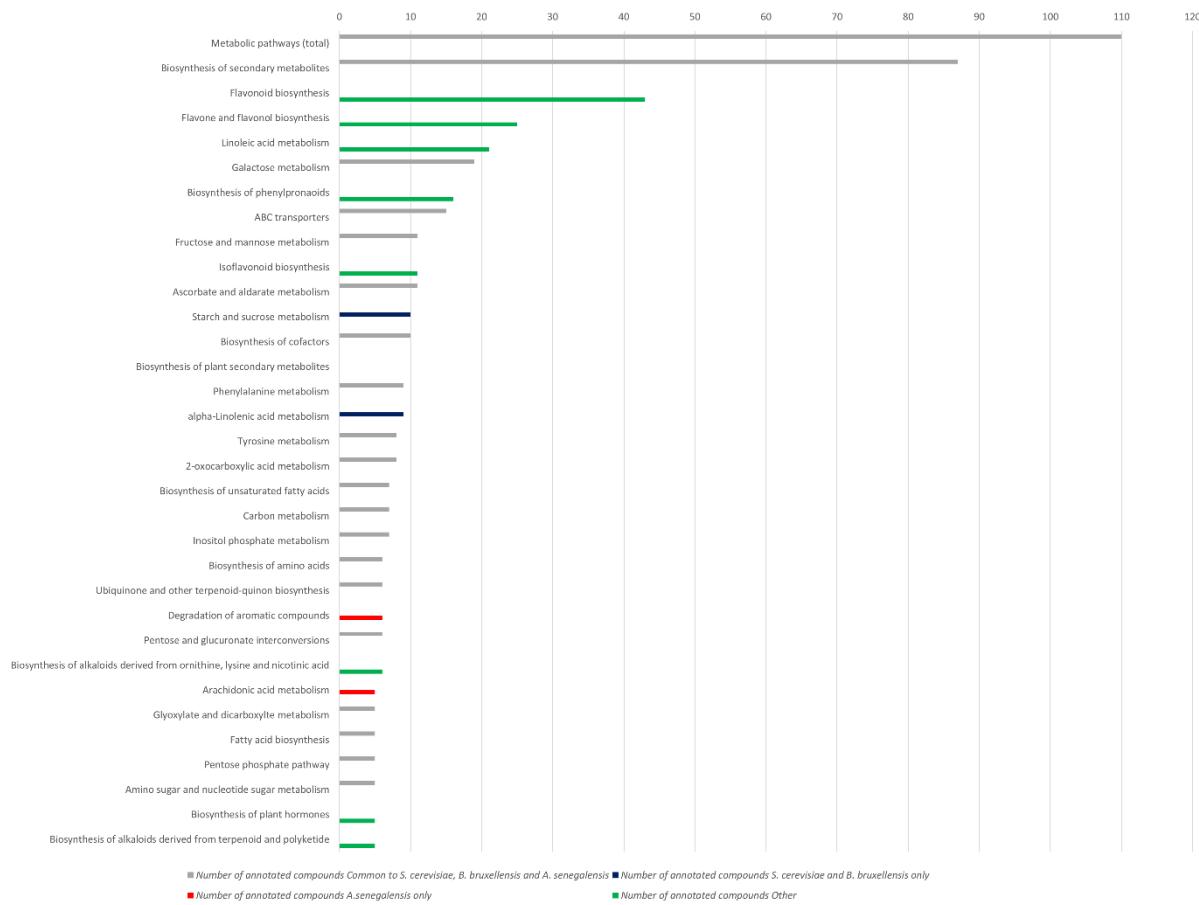


Figure S2: Distribution of annotated compounds using MASSTRIX database according to metabolic pathways according to KEGG Mapper Color.