

**UNTARGETED HILIC-MS-BASED METABOLOMICS APPROACH TO
EVALUATE COFFEE ROASTING PROCESS: CONTRIBUTING TO AN
INTEGRATED METABOLOMICS MULTIPLATFORM**

Raquel Pérez-Míguez^{1#}, María Castro-Puyana^{1,2#}, Elena Sánchez-López^{1,2}, Merichel Plaza^{1,2}, María Luisa Marina^{1,2*}.

¹Departamento de Química Analítica, Química Física e Ingeniería Química, Universidad de Alcalá, Ctra. Madrid-Barcelona Km. 33.600, 28871 Alcalá de Henares (Madrid), Spain.

²Instituto de Investigación Química “Andrés M. del Río” (IQAR), Universidad de Alcalá, Ctra. Madrid-Barcelona Km. 33.600, 28871 Alcalá de Henares (Madrid), Spain.

#These authors contributed equally to this work.

*Correspondence: mluisa.marina@uah.es; Fax: +34-918854971; Tel.: +34-918854935
(Departamento de Química Analítica, Química Física e Ingeniería Química, Universidad de Alcalá, Ctra. Madrid-Barcelona Km. 33.600, 28871 Alcalá de Henares (Madrid), Spain).

Figure captions.

Figure S1. Base peak chromatograms (BPC) obtained in positive ionization mode for green coffee (GCB) (A); light coffee (LRC) (B); medium coffee (MRC) (C); and dark coffee (DRC) (D) under optimal separation conditions. HILIC-MS conditions are summarized in section 3.3.

Figure S2. Hierarchical cluster analysis (HCA) in positive (A) and negative (B) ionization modes for the four groups of coffee samples (GCB, LRC, MRC, and DRC) submitted to different roasting process.

Figure S3. Diagrams of the trends observed for all the tentatively and unequivocally compounds both in negative and positive ionization mode along the coffee roasting process.

Figure S1.

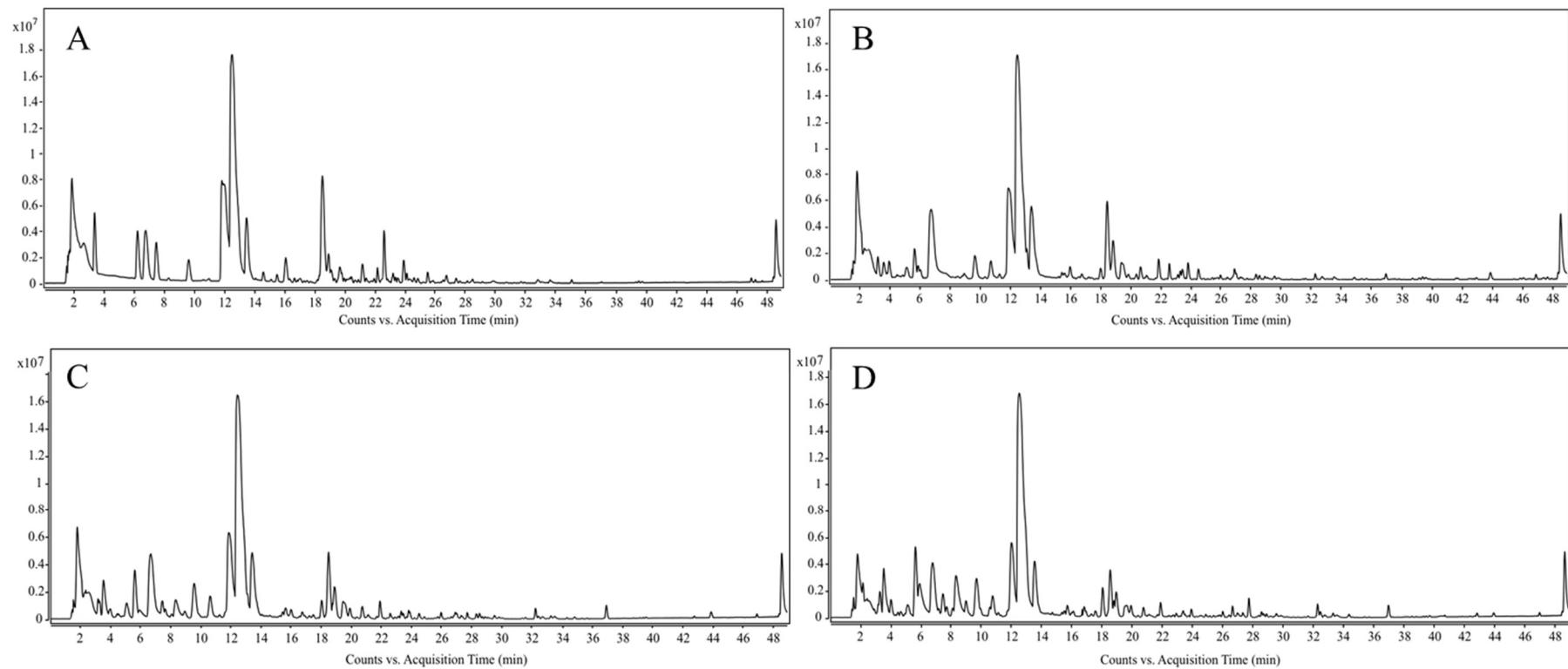


Figure S2-A

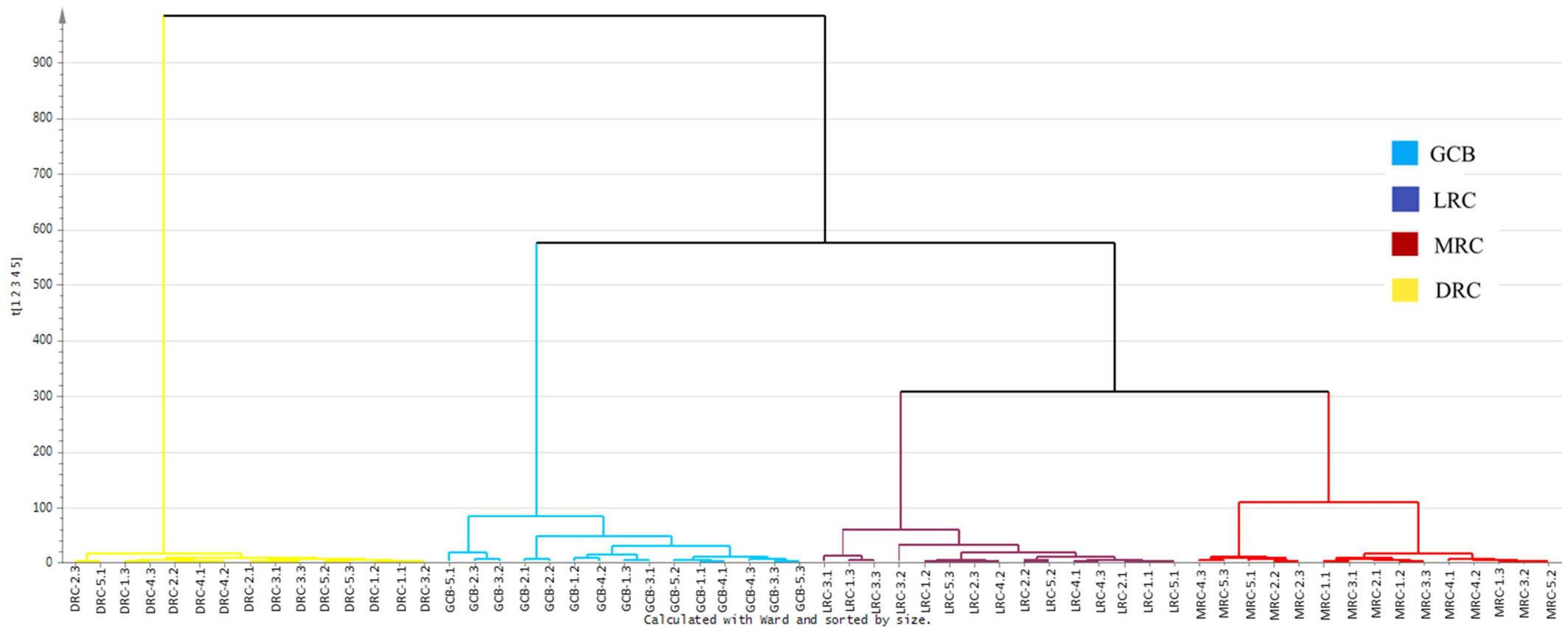


Figure S2-B

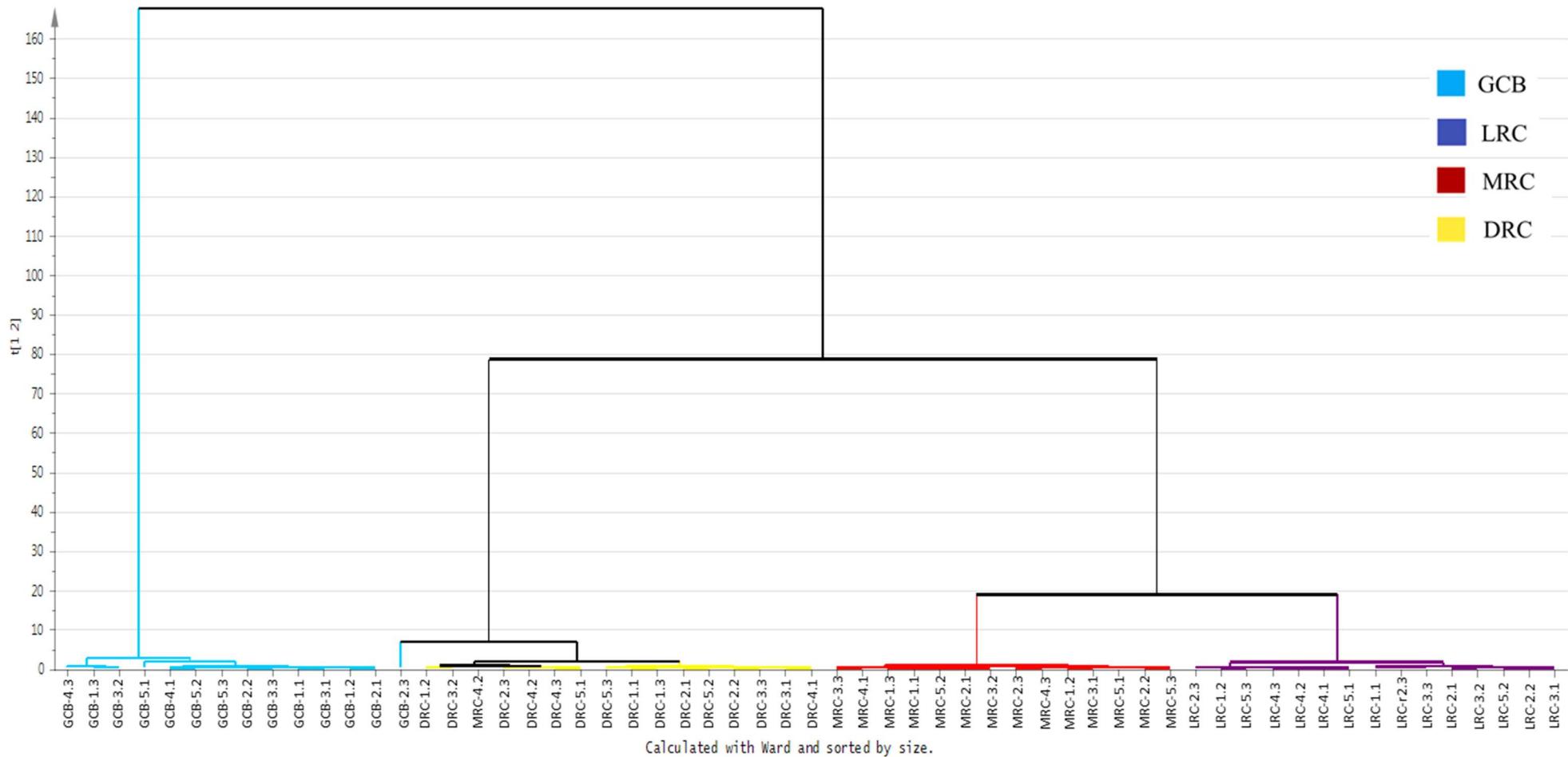


Figure S3.

