9th International Cherry Symposium

Abstract Book

Oral

S2-04

Characterization of sugar content in sweet cherry populations

for genetic analyses

C. Gracia^{1,2}, A. Calle³, E. Arias², A. Wünsch^{1,2}

- ¹Departamento de Ciencia Vegetal, Centro de Investigación y Tecnología Agroalimentaria de Aragón (CITA). Avenida de Montañana 930, 50059 Zaragoza, (Spain)
- ²Grupo de Investigación en Alimentos de Origen Vegetal. Facultad de veterinaria (Universidad de Zaragoza). Instituto Agroalimentario de Aragón-IA250013 Zaragoza, (Spain)
- ³Department of Plant and Environmental Sciences, Clemson University, Clemson, SC (USA)

Keywords: Sweet cherry; Anthocyanin accumulation; BBX transcription factor; Light

Abstract

Sweet cherry is a temperate fruit highly appreciated by consumers. Most genetic studies on fruit quality have focused on physical parameters like size, colour, or firmness. In contrast, chemical parameters associated with organoleptic quality, like sweetness and acidity, have been less studied. In this work, the sugar content was characterized in individuals from five sweet cherry populations for two years. The segregation and heritability of the different sugars identified were investigated. Fruit samples from each individual tree were evaluated for soluble solids content (SSC), and identification and quantification of the main sugars (glucose, fructose, sorbitol, and sucrose) were performed using UPLC. Sugars and SSC segregation were analysed to test whether they conformed to a normal distribution. In addition, the sugars and SSC correlation between years, and the broad-sense heritability (H^2) in each population were studied. These results will be further used for QTLs mapping of these sugars content variability.