Commercial yeast strains do not significantly disseminate in vineyard ecosystems

Dorit Schuller(1), Eva Valero(2), Brigitte Cambon(2), Margarida Casal(1) and Sylvie Dequin(2)

(1) Centro de Ciências do Ambiente, Departamento de Biologia, Universidade do Minho, 4710-057 Braga Codex, Portugal
(2) Institut National de la Recherche Agronomique, UMR Sciences pour l’Oenologie, 34060 Montpellier, France.

Industrial yeasts are traditionally used in winemaking and subsequently released into the environment. Nevertheless, there is very little data showing what becomes of them in the natural environment.

To evaluate the dynamics of industrial yeast strains in the vineyard, a large-scale sampling plan was devised over a period of three years. This includes 36 sampling sites in 6 different vineyards (3 in France, 3 in Portugal) that used industrial starter yeast for at least 5 years. In each vineyard samples were taken before harvest (annual remanence) and at late harvest (immediate release), at three distances from the winery and from opposite directions. Small-scale fermentations were realised. Must samples were plated when 70 g/L of the CO\textsubscript{2} had been released, and 30 colonies were randomly selected. By using selective media non-	extit{Saccharomyces} strains were eliminated, and the remainder were characterised by karyotyping [1] and RFLP of mitochondrial DNA [2].

During the year 2001 the genetic patterns of only 2% of 720 isolates in France (collected at 100 to 1000m from the winery) were identical to those of the industrial yeast. In Portugal, the typing patterns of about 10% of the 570 isolates were identical to those of the commercial yeast, only in post-harvest samples close to the winery (20-40 m). The identification of the 	extit{S. cerevisiae} strains isolated during the year 2002 is now underway.

These first data sets show that dissemination and remanence of commercial yeast in the vineyard is very limited. Next year’s data will complete this study and allow us to confirm these results.