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## Conclusions

Since its establishment, the cv. Chardonnay has undergone a – direct or indirect – selection favoring the biotypes with highly acidic must, intended for the production of sparkling wine. This character is very stable within the clone, as shown by the constancy of the figures evaluated on different vintages as well as from the small standard deviation for this character among the progeny seedlings.

Also to be taken into account and further tested in any future breeding program is the finding that in this cultivar a larger bunch but a smaller berry size parallel a higher acidic content.

## References

FANIZZA, G.; RADDI, P.; 1973: The heritability of fruit ripening date in Vitis vinifera L. Vitis 12, 93-96.

FIROOZABADY, E.; OLMO, H. P.; 1987: Heritability and correlation studies of certain quantitative traits in table grapes, *Vitis* spp. Vitis 26, 132-146.

O.I.V.; 1983: Codes des caractères descriptifs des variétés et espèces de *Vitis*. Office International de la Vigne et du Vin, Paris.

VINCOURT, P.; GALLAIS, A.; 1983: Sur la recherche des critères de sélection: La régression géno-phénotypique. Agronomie 3, 827-830.

WAGNER, R.; 1975: Amélioration génétique de la vigne. Ann. Amélior. Plantes 25, 151-175.

## Characterization of *Vitis vinifera* biotypes through biochemical methods

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A b s t r a c t: We have shown that the soluble protein fraction from pollen wall was clone specific and independent of environmental as well as cultural conditions.

Along this line, we have compared several biotypes belonging to the cultivar Nebbiolo; the samples were collected in two distinct, typical areas. When analyzed with two different electrophoretic procedures, the protein pattern varied extensively between the groups while it was constant within each group.

Three supposedly distinct cultivars – Vermentino. Pigata and Favorita – grafted on the same rootstock and grown in the same farm, gave excactly the same electrophoretic pattern for pollen wall proteins as well as storage and enzyme components from the seed. This evidence suggests that the three cultivars share an identical genetic background.