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DEVELOPMENT OF A MARKER ASSISTED SELECTION PROGRAM FOR THE IMPROVEMENT OF DURUM WHEAT (*TRITICUM DURUM* DESF.)

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Italy is the main producer of pasta in the world and the genetic improvement of durum wheat represents a strategic activity for the entire agro-industrial sector.

DNA markers have an enormous potential to improve the efficiency and precision of conventional plant breeding via marker-assisted selection (MAS). This paper describes the work carried out at the Cereal Research Centre (CRA-CER) for the development of a MAS breeding program dedicated to the pyramiding of genes for low lipoxygenase (LOX) activity (Lpx-B1.1), high protein content (GPC; Gpc-B1), high yellow pigment content (YPC; Psy-A1) and disease resistances. The following R genes were considered: leaf, stripe and stem rust (Lr14c, Yr36 and Sr13 or Sr26, respectively), powdery mildew (Pm36) and soil borne cereal mosaic virus (SBCMV; QSbm.ubo-2BS).

A set of durum wheat varieties and introgression lines carrying the desirable genes were chosen as donor lines, while the recipient line was the Italian durum cultivar PR22D89, characterized by a high gluten quality and good yield.

The crosses were performed separately for each donor line with PR22D89, than the introgressed genes were first fixed in a homozygous after the screening of the F_2 populations. Then, the F_2/F_3 plants homozygous for the same genes and meeting the required phenotypic standards were selected for further crosses in order to combine up to 4 genes of interest segregating in the same populations.

Presently, several hundreds of genotypes are under evaluation and some $F_{3:5}$ lines are carrying genes at the homozygous state combining four different traits of interest: high GPC, low LOX activity, resistance to stripe rust and powdery mildew; as well as high GPC, low LOX activity, resistance to stripe and leaf rust. These lines exhibited a good increase in GPC with a very limited negative impact on grain kernel weight.