**Poster Abstract – A.11** 

## **RESPONSES TO DIVERGENT SELECTION FOR COB COLOUR IN A MAIZE POPULATION**

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## Cob colour, P1 gene, marker assisted selection, correlated traits, leaf number

In a previous study, we carried out four cycles of recurrent selection for grain yield in the  $F_2$ population of the maize single cross A632 (red cob) x Mu195 (white cob). Significant gains were achieved for grain yield and other agronomic traits; moreover, an increase of the frequency of the red cob allele was observed throughout the selection process, suggesting an association between grain yield and cob colour (determined by P1 gene in chromosome 1). In a subsequent study, nearisogenic lines with red or white cob were developed by backcross to both recurrent parents A632 (using Mu195 as donor of the white allele) and Mu195 (using A632 as donor of the red allele). These near-isogenic lines were evaluated both per se and in hybrid combination and the lines with red cob (R) yielded more than the corresponding lines with white cob (W) in both A632 and Mu195 backgrounds, thus confirming the association between grain yield and cob colour. These findings prompted us to undertake a divergent selection for cob colour, i.e., for R or W, in the  $F_2$  of A632 x Mu195 to evaluate the effect of this marker assisted selection on grain yield and other traits. About 200 random F<sub>2</sub> plants were selfed and the resulting F<sub>3</sub> seeds were bulked according to the cob colour of the ears. Forty seeds were bulked from each R ear to produce the selected  $F_3$ -R population; similarly, 40 seeds were bulked from each W ear to produce the selected F<sub>3</sub>-W population. Moreover, 20 seeds were bulked from each ear (regardless of the cob colour) to produce the unselected  $F_3$  population. The three populations were tested in two field trials. The  $F_3$ -R mean across the two trials was superior to the F<sub>3</sub>-W mean for number of leaves per plant (19.2 vs. 18.0), days to flowering, whole leaf area, kernel moisture, number of kernels per plant and grain yield (3.44 vs. 3.05 Mg ha<sup>-1</sup>). In contrast, the mean across the two selected populations was rather close to the mean of the unselected  $F_3$  population for most traits, indicating a prevalence of symmetric responses to selection. In comparison with the unselected F<sub>3</sub> population, F<sub>3</sub>-R population showed a superiority for grain yield of 0.22 Mg ha<sup>-1</sup> corresponding to a gain of 7.7%, which is slightly higher than the average gain per cycle (7.3%) achieved in the laborious recurrent selection previously conducted. When differences between F<sub>3</sub>-R and F<sub>3</sub>-W populations were referred to the number of leaves per plant, either directly, as for leaf area (i.e., considering the average area per leaf), or indirectly, as for grain yield (considering the average grain yield per leaf area unit), the differences between F<sub>3</sub>-R and F<sub>3</sub>-W populations were not significant any more. These findings indicate that in the material herein investigated selection for cob colour can lead to an appreciable yield improvement, achieved at least partly by changes concerning the number of leaves per plant and associated traits. These findings could be accounted for by assuming that close to the P1 gene there is a QTL controlling the number of leaves per plant and then date of flowering, whole leaf area, number of kernels per plant, kernel moisture and grain yield, following a sequence of causally related events. To gain a better insight into the genetic effects of this QTL and the frequency of recombination with *P1* gene, we are going to compare the selected  $F_3$ -R and  $F_3$ -W populations with the selected  $F_3$ -R and  $F_3$ -W populations obtained by selfing the plants derived by one cycle of random mating in the same source  $F_2$ .