

CHARACTERISATION AND EVALUATION TOWARDS SELECTION OF MAIZE LANDRACES WITH THE BEST *PER SE* PERFORMANCES

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ABSTRACT

The narrow genetic base of commercial maize varieties emphasise the necessity for conservation, characterisation and utilisation of germplasm stored within gene banks. Broad genetic variability preserved in the Maize Research Institute Zemun Polje (MRIZP) gene bank, which includes accessions which originated from the Western Balkan, as a part of European corn-belt, is an exceptional source of desirable traits for enriching breeders' working collections for maize breeding under temperate conditions. Preliminary screening for abiotic stress tolerance, which marked 321 maize landraces, served as the first step in stratification process of selecting a smaller number of accessions from the entire gene bank local collection. After classification of these landraces into eleven homogenous groups, the objective of this study was to continue the stratification process of selection (as a second step), based on evaluation of agro-morphological traits of interest for breeding. The conducted evaluation highlighted 40 landraces with the best *per se* performances, important for breeding. Out of them, 28 early-maturing flint landraces with stiff stalks, low positioned ears, high yield potential and good general ear assessment could be considered as valuable source for their introgression into elite flint germplasm pool.

Keywords: flints, gene bank, heterotic potential, pre-breeding, *Zea mays* L.

INTRODUCTION

Richness of genetic resources stored in gene banks is very little used in plant breeding. The huge gap between genetic variability maintained in gene banks and commercial breeding is caused by lack of documentation, of appropriate collection description and of information important for breeding, as well as by the fact that many samples have limited adaptability and seed quantities (Technow et al., 2014). Moreover, large number of gene bank accessions loaded with many undesirable properties from the aspects of modern cropping practices and breeding, require expensive and long-term process of their characterisation and improvement, while the final results, in a commercial sense, are uncertain.

The evaluation of five distinct Central European OP varieties, 85 hybrids and their parental components with the total of 55 SSR markers revealed that genetic variation within

and among cultivars decreased significantly during the last five decades (Reif et al., 2005) and is almost certainly limited in comparison to the large genetic diversity available in gene banks (Le Clerc et al., 2005). Numerous unique alleles present in these OPVs, were lost in the elite flint pool, particularly in the last two decades, which impose a necessity to enlarge genetic base in breeding programmes that encompass only 3-5% preserved maize variability (Curry, 2017). Landraces adapted to local growth conditions could play significant role in the process of continuous introgression into commercial breeding material (Mitrović et al., 2016; Vega-Alvarez et al., 2017; Mikić et al., 2017).

The preservation of genetic resources diversity, significance of genetic variability for the survival of any species and its breeding, as well as previously mentioned actual difficulties for the efficient use of these resources, generated the pre-breeding concept (Ortiz et al., 2010).

The implementation of long-term pre-breeding programmes is the most promising way of linking genetic resources and commercial breeding programmes. The concept encompasses programmes and activities for the purpose of identifying desirable properties and/or genes from non-adapted (exotic and semi-exotic) material, including adapted material that will be under certain selection pressure, as well as programmes and activities that improve gene bank materials up to the level acceptable to commercial breeding programmes (Shimelis and Laing, 2012). However, the fact that there are only few such programmes is a limiting factor for more efficient utilisation of landraces from gene bank collections (Carena, 2005). Successful examples of maize pre-breeding programmes are the Latin American Maize Project (LAMP), the Germplasm Enhancement of Maize (GEM) (Carena et al., 2009a) and EU GENRES (Gousnard et al., 2005).

Within Maize Research Institute Zemun Polje (MRIZP) gene bank, that encompasses 2217 local maize accessions, pre-breeding activities are carried out through the long-term stratification process of characterization and evaluation of gene banks accessions and the selection of a smaller number of genotypes with desirable properties, in order to introgress them into elite breeding material (Vančetović et al., 2015).

Previous screening for drought tolerance marked 321 local maize landraces as possible sources of tolerance (Babić et al., 2012a), and was exploited as the first step in stratification process. After classification of these landraces into eleven homogenous groups based on morphological similarity (Babić et al., 2015), the objective of this study was to continue the stratification process of selection

(as a second step) of a smaller number (approximately 10-15% of representative landraces of each homogenous group) for the following pre-breeding activities. The selection of landraces was based on characterisation and evaluation of agronomic traits of interest for breeding, giving the priority to flint type.

MATERIAL AND METHODS

Plant material

Within the long-term pre-breeding activities, we focused on 321 MRIZP gene bank western Balkan landraces (<https://mrizp.rs/emdb/default.htm>), previously designated as possible sources of drought tolerance (Babić et al., 2012a). Due to the large number of genotypes (321), their assignment into homogeneous groups according to CIMMYT/IBPGR descriptors, was undertaken. Based on 27 morphological traits, along with the application of non-hierarchical and hierarchical cluster, correspondence and discriminant analyses, eleven homogenous groups were created (Figure 1), which were fully described in Babić et al. (2015). However, 11 landraces were excluded from further evaluation, due to too long/short vegetative period and high percentage of broken and barren plants.

Stratification process for selection of the best performing landraces, as a procedure which allows simultaneously retaining as much variability as possible, is presented through scheme of planned complete pre-breeding activities (Figure 2). Due to the large number of observed genotypes and economic aspects, the selection in the initial steps was performed on the basis of phenotypic characterisation.

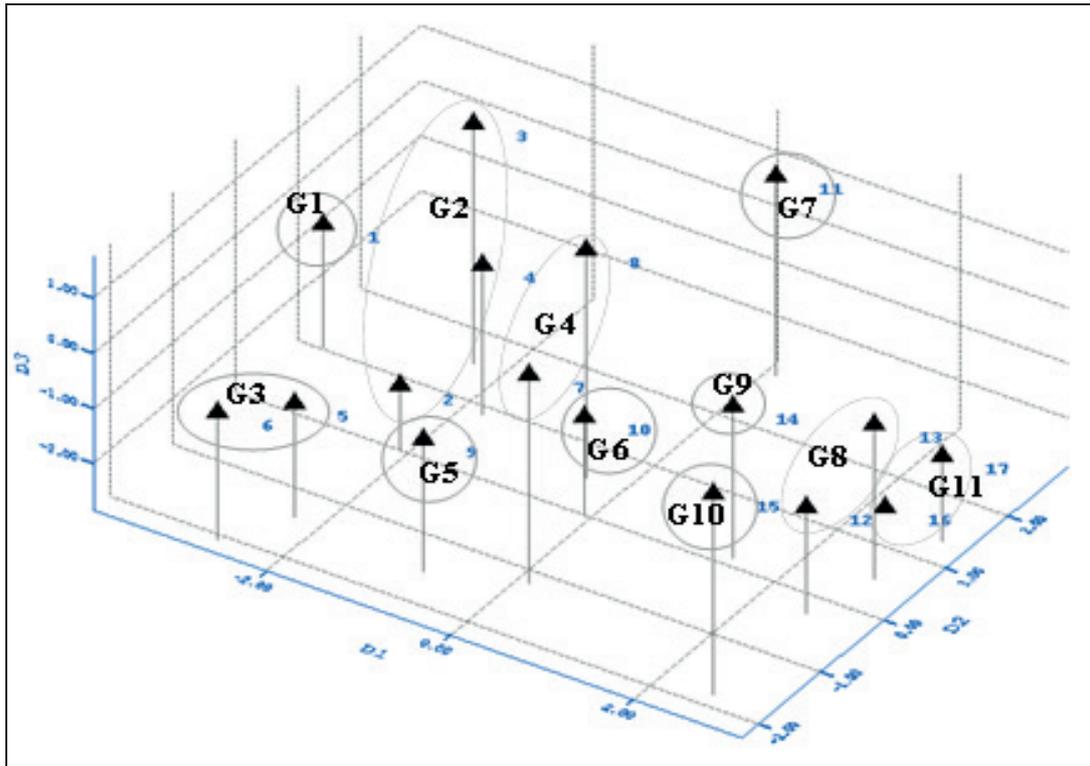


Figure 1. Distribution of cluster's centres, and groups for investigated local landraces in 3D discriminate space

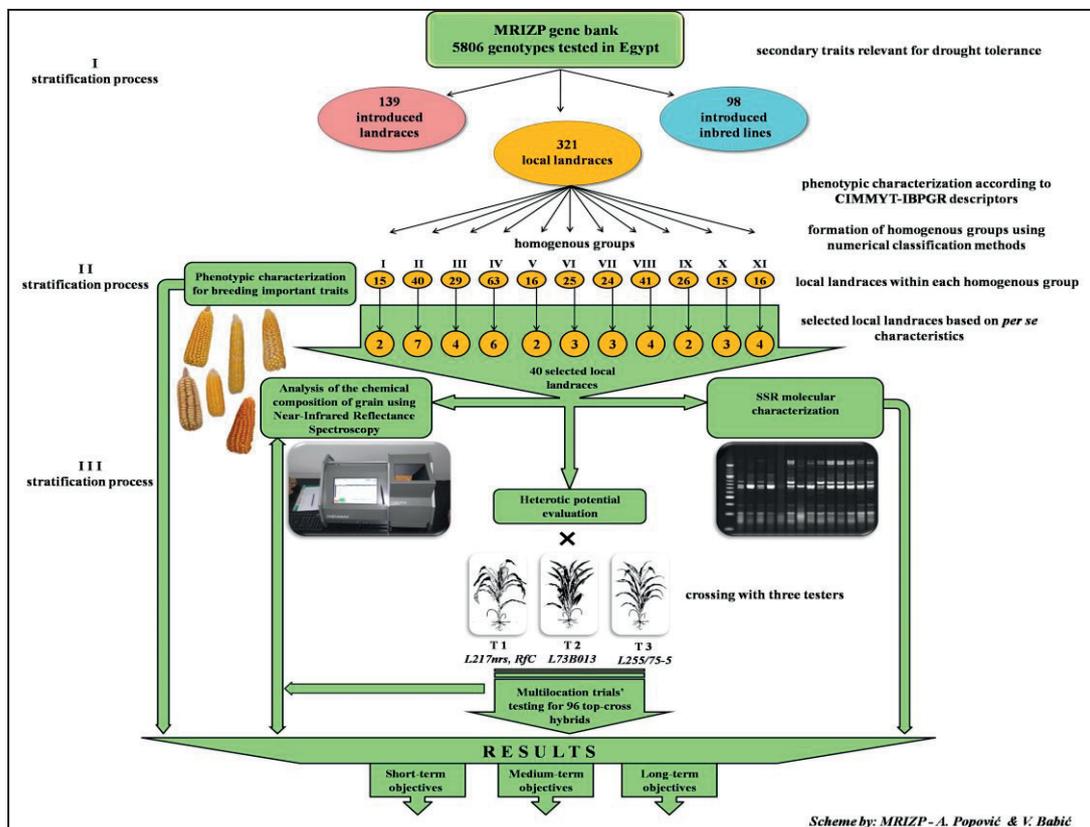


Figure 2. Selection of local maize landraces through stratification process

Field experiment

The experiment was carried out in 2013 and 2014 in Zemun Polje, Serbia (44°52'N, 20°19'E, 81 m asl). Landraces (310) were sown within eleven homogenous groups and further characterisation based on breeding important traits was conducted. The number of landraces per each homogenous group was presented in Figure 2. Each landrace was sown in two rows, 30 plants per row (15 hills were spaced at 40 cm, while the inter-row distance was 75 cm, two plants per hill). The experiment was set up according to the randomised complete block design, in two replications.

Beginning and mid-flowering dates (i.e. 10% and 50% tasseling, anthesis and silking) were recorded. The plant and ear height were measured on 15 plants per landrace, in both replications. The intensity of leaf green colour was estimated on the 1-3 scale: 1 - light green, 2 - medium green and 3 - dark green.

At harvest, the following traits were recorded: the total number of plants per row, the number of lodged and broken plants, as well as the number of ears per landrace. The yield per plot was determined. A sample of five ears was used to determine the grain moisture content. Visual ear evaluation based on shape, length, diameter, number of kernel rows, seed set and general ear assessment (1 - poor, 2 - medium, 3 - good) was estimated. The grain type was also determined in the same sample according to the International Union for the Protection of Cultivated Varieties of Plants (UPOV) Descriptor: 1 - flint, 2 - flint-like, 3 - intermediate, 4 - dent-like, 5 - dent, 6 - sweet, 7 - pop. Plants and ears were photographed.

In parallel, five commercial inbred lines (potential testers) were sown, in order to evaluate the date of flowering under the same environmental conditions, as important information for subsequent crossings.

Statistical analysis

IBM SPSS Statistics 20 programme was used for assignment of maize landraces into 11 homogenous groups, which include application of nonhierarchical, hierarchical

cluster, correspondence and discriminant analyses (Babić et al., 2015). Descriptive statistics were used for all agro-morphological traits measured. Two-way ANOVA was used for determination of differences in grain yield within each homogenous group.

RESULTS AND DISCUSSION

Despite the extensive genetic potential contained in gene bank collections, the valuable variations remain mostly unused by plant breeders (Carena et al., 2009b; Masuka et al., 2017). Lack of information significant for breeding is frequently a limiting factor when using gene bank material. Although there were 27 traits of selected landraces characterised by the CIMMYT/IBPGR Descriptor, there was no information on stalk quality, combining ability and grain yield, the traits important for breeding. Since agro-morphological characterization of germplasm accessions is fundamental in order to provide information for plant breeding programs, the following traits were tested during the preliminary evaluation: early and final growth, tasseling and silking, plant and ear height, lodged and broken plants, ear number per plant, endosperm type, kernel colour, ear quality and yield (Nascimento et al., 2011; Badu-Apraku and Fakorede, 2017). Therefore, in this experiment, the selected landraces (310) were sown according to defined groups and the preliminary evaluation for breeding important traits was done.

The plant and ear heights are presented in Figure 3A. The lowest plant and ear height, as well as the ratio for these two parameters were noticed in the first three groups of populations. Somewhat higher average values of the same parameters were recorded in landraces from the groups IV, V, VI and VII. Study of plant architecture, phenology and yield components on 336 maize lines, revealed that yield components distinguished dents from lower yielding genetic groups (Bouchet et al., 2017). This was in line with our findings that the higher participation of dents is followed by the higher average plant height, and it was observed in the groups VIII

and XI. These groups comprised higher yielding medium to late maturing dent maize landraces. Also, it was noticed that ascendant group number of a homogenous group exhibited the trend of the constant increase of ratio between PH and EH, ranging from 24% (for the group I) to 39% (for the group XI).

The lowest number of days to 50% tasseling was recorded in landraces from the group I (47.4), while the number of days in landraces from groups X and XI ranged from 58.3 to 59.1, respectively (Figure 3B). The highest anthesis-silking interval (ASI) was in

the group XI (3.8), and the lowest in the groups I and II (0.1 and 0.6, respectively). More pronounced variation in ASI (even up to four days) was observed on individual landraces level. However, the 4-days interval difference could be considered as continuous, undisturbed pollination, if it occurs under optimal conditions (i.e. under conditions without extreme temperatures that lead to pollen abortiveness or without the combination of high temperatures and humidity that disturb physiological role of silk up to 40%) (Cerović et al., 2014).

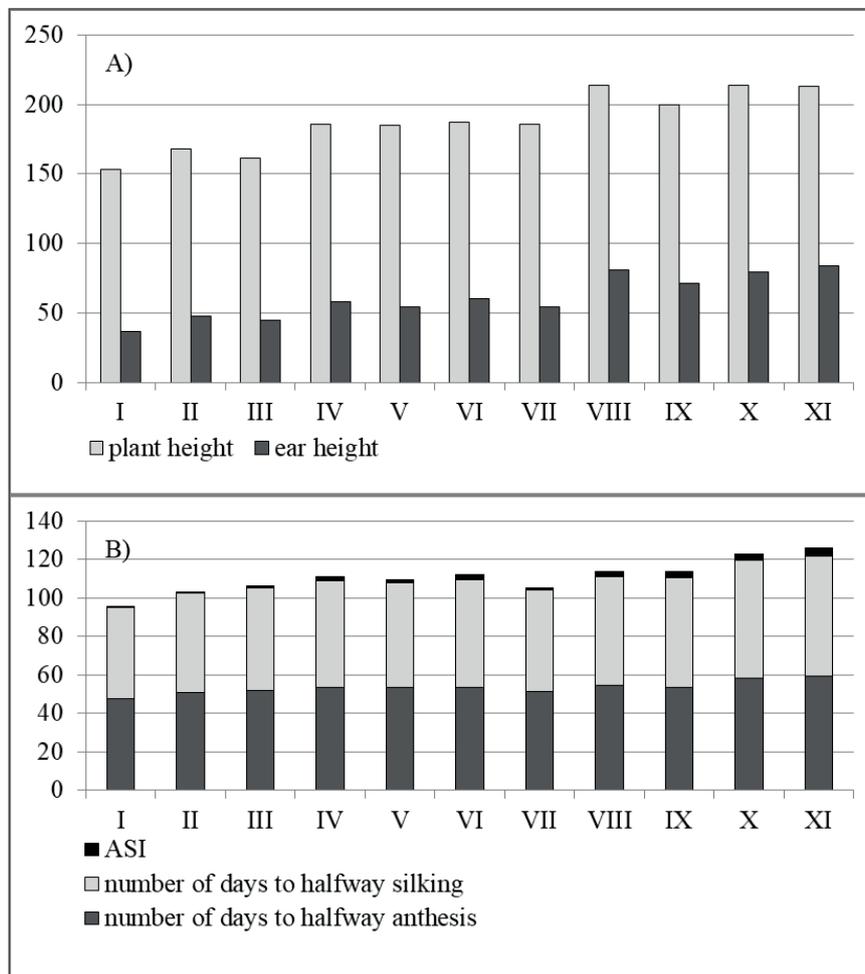


Figure 3. Average values of: A) plant and ear height, B) number of days to flowering and ASI, over homogenous groups

The grain type was evaluated according to the UPOV Descriptor, and then the percentage of participation of different kernel types per homogenous group was estimated (Figure 4A). The group I exclusively comprised extra-early maturing

flint landraces originated from the sites where landraces were grown on rather isolated fields. Inability to cross with other maize landraces might be reflected in preservation of original introduced form.

Flints and flint-like landraces, originating from the hilly regions, prevail in groups II and III (90 and 89%, respectively), are similar to those from the group I, although with a slight percentage of derived and semi-dent kernel types and a certain number of pop maize landraces. Even a slight participation of dent germplasm in landraces from groups II and III contributed to their more successful spreading and this is confirmed by a higher number of collected accessions. Flints and flint-like kernel types, which prevail in the groups IV (63), V (16), VI (25) and especially in the group VII (24), differ from the flints of the first three groups. These are the early-maturing flint landraces, with more robust and higher plants, wider leaves and longer ears. However, certain percentages of dent-like and dent kernel types were recorded in the groups IV and VI.

In the group VIII (41 landraces) predominate dent and dent-like kernel types (80.2%), which is followed by 12.2% of intermediate, and 7.6% of flint and flint-like kernel types. These landraces probably originated from the fourth introduction (at the end of 19th century) of American Corn Belt dents, crossed with already existing flints (Babić et al., 2012b). The group IX (26 landraces) encompasses 50% flint and flint-like types, 11.5% intermediate and 38.4% dent-like and dent kernel types. In the group X prevailed dent-like and dent landraces (53.3%), with 33.3% of intermediate kernel type. Sixteen landraces of the group XI, predominantly collected in Pannonian Plain, consist of dents morphologically the most similar to dents from South America.

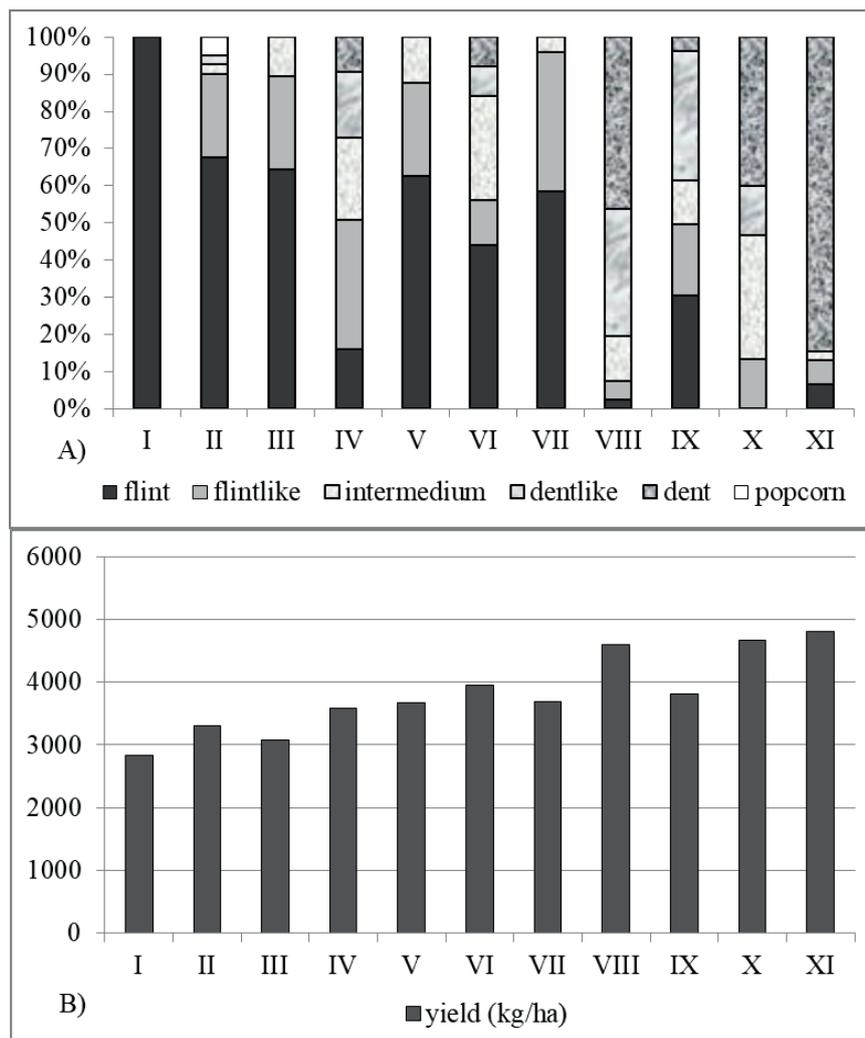


Figure 4. Average values of: A) kernel types B) grain yields over homogenous groups

By comparing average yield of these eleven groups, it can be concluded that the first three homogenous groups represent early-maturing, mainly yellow-orange flints, characterised with lower yield and similar morphological properties (Figure 4B). The yield in the homogenous groups IV, V, VI, VII and IX is approximately equal, ranging from 3.59 t ha⁻¹ to 3.96 t ha⁻¹. The yield of landraces from the groups VIII, X and XI, which are predominantly late-maturing dents and their derived types, is over 4.5 t ha⁻¹.

Besides good *per se* performances, the initial breeding population must have good combining ability with another population, or with a group of populations or inbreds, that will serve as a second parent in the process of hybridisation (Vančetović et al., 2015). The main idea of this work was to select the most representative landraces of each of eleven homogenous groups to be crossed with elite commercial testers in order to define their heterotic pattern.

Flowering synchrony between male and female parents of maize hybrids is an important driver for improved kernel set and grain yield (Worku et al., 2016). In order to evaluate the date of flowering under the same environmental conditions, five commercial testers were sown. Three inbred testers were chosen for subsequent testing: L225/75-5 (Lancaster germplasm), B73B013 (BSSS x Iowa dent) and L217 (Iowa dent). Due to extremely poor agronomic traits (poorly developed tassels, short pollen shading period and small amount of pollen), inbreds F2 and Polj17 were excluded.

When selecting the initial material adapted to the testing conditions, a high level of desirable traits is important, as it provides better chances to obtain superior inbreds through direct selection (self-pollination)

from the landrace and/or requires a lower number of selection cycles to achieve the desired level of traits within the landrace (Gorjanc et al., 2016). In this evaluation, landraces with small ASI values, low positioned ears, high yield *per se*, dark green leaves, a low number of lodged and broken plants, as well as a good general estimation of ears were preferable. In order to be selected, landrace should obtain yield significantly higher than the average value of yield from the homogenous group to which the landrace belongs. This requirement was not achieved by the landraces 589, 1379 and 2144. However, they were selected for the following reasons: landraces 589 and 1379 are early-maturing genotypes with favorable plant architecture, cylindrical shape of ear and pests/disease damage-free. The landrace 2144 was characterized by low positioned, well developed cylindrical and long flinty ear.

In accordance with the set goals, 40 landraces (10-15% from each homogenous group) were selected, such as follows: two, seven, three, seven, two, three, three, four, two, three, four over groups I to XI, respectively (Table 1). The proportional selection of representative landraces from each of eleven groups leads to the equal inclusion of the available gene pool variability. Typical representatives of each homogenous group were presented in Figure 4.

The narrow base of existing breeding collections imposes the necessity of revising flint landraces (Böhm et al., 2014). Considering the high genetic variation found within the Western Balkan flint landraces, it seems promising to use the very best of them for programmes of introgression in order to enlarge the genetic base of the elite flint germplasm pool in Central Europe.

ROMANIAN AGRICULTURAL RESEARCH

Table 1. Basic passport data and observed parameters in selected landraces

| HG | AN | CC | AL | PH (cm) | EH (cm) | PH/EH (%) | LBP | Y (t ha ⁻¹) | GEA | GC | KT |
|----|------|-----|-----|---------|---------|-----------|-----|-------------------------|-----|----|----|
| 1 | 1869 | CRO | 850 | 147 | 30 | 20.47 | 5 | 3.46 | 2 | 3 | 1 |
| | 1890 | CRO | 600 | 147 | 41 | 27.68 | 4 | 4.27 | 3 | 2 | 1 |
| 2 | 594 | MAC | 661 | 175 | 62 | 35.62 | 1 | 3.04 | 2 | 2 | 1 |
| | 773 | SRB | 500 | 188 | 52 | 27.74 | 2 | 4.23 | 3 | 3 | 1 |
| | 1185 | MNE | 215 | 160 | 38 | 23.59 | 3 | 3.45 | 3 | 3 | 1 |
| | 589 | MAC | 661 | 168 | 56 | 33.02 | 2 | 2.39 | 3 | 1 | 1 |
| | 1379 | BIH | 650 | 169 | 44 | 26.42 | 6 | 2.12 | 2 | 1 | 1 |
| | 1381 | BIH | 700 | 190 | 53 | 28.33 | 5 | 4.50 | 3 | 2 | 2 |
| | 1267 | MNE | 500 | 202 | 54 | 27.07 | 0 | 4.53 | 2 | 2 | 3 |
| 3 | 13 | MNE | 900 | 174 | 54 | 31.23 | 3 | 3.14 | 3 | 3 | 1 |
| | 467 | BIH | 500 | 159 | 34 | 21.19 | 2 | 3.68 | 3 | 3 | 1 |
| | 144 | SRB | 900 | 181 | 67 | 36.81 | 1 | 4.39 | 3 | 2 | 1 |
| 4 | 871 | MAC | 105 | 164 | 43 | 26.30 | 3 | 3.55 | 3 | 3 | 2 |
| | 1960 | BIH | 750 | 178 | 46 | 25.83 | 4 | 4.84 | 2 | 3 | 3 |
| | 1276 | MNE | 800 | 195 | 57 | 29.28 | 3 | 4.97 | 2 | 3 | 5 |
| | 846 | BIH | 700 | 182 | 48 | 26.32 | 10 | 3.35 | 3 | 3 | 3 |
| | 1384 | BIH | 650 | 185 | 50 | 27.07 | 7 | 3.99 | 3 | 3 | 3 |
| | 2144 | CRO | 620 | 175 | 45 | 25.77 | 3 | 3.41 | 3 | 2 | 1 |
| | 1534 | SRB | 174 | 197 | 68 | 34.51 | 1 | 5.44 | 3 | 3 | 1 |
| 5 | 642 | SRB | 533 | 184 | 45 | 24.77 | 4 | 4.29 | 2 | 3 | 1 |
| | 1798 | SRB | 550 | 195 | 70 | 35.76 | 0 | 5.36 | 3 | 2 | 3 |
| 6 | 1895 | CRO | 500 | 181 | 56 | 30.53 | 3 | 5.14 | 3 | 2 | 1 |
| | 2033 | CRO | 50 | 198 | 71 | 35.80 | 2 | 5.47 | 3 | 3 | 3 |
| | 2006 | BIH | 800 | 206 | 81 | 39.37 | 3 | 5.48 | 3 | 2 | 1 |
| 7 | 2230 | BIH | 700 | 189 | 53 | 28.14 | 2 | 4.84 | 3 | 3 | 2 |
| | 2236 | BIH | 750 | 205 | 59 | 28.87 | 2 | 4.20 | 3 | 3 | 2 |
| | 2176 | SLO | 620 | 210 | 55 | 26.28 | 4 | 4.19 | 3 | 1 | 1 |
| 8 | 2249 | BIH | 800 | 230 | 90 | 39.08 | 2 | 7.74 | 3 | 2 | 5 |
| | 2036 | BIH | 800 | 205 | 72 | 34.54 | 3 | 5.48 | 3 | 2 | 4 |
| | 1665 | MAC | 700 | 218 | 91 | 41.90 | 0 | 5.79 | 3 | 2 | 4 |
| | 632 | SRB | 400 | 220 | 94 | 43.19 | 4 | 5.34 | 2 | 2 | 5 |
| 9 | 1945 | BIH | 700 | 204 | 72 | 35.02 | 5 | 4.45 | 3 | 3 | 1 |
| | 2047 | CRO | 180 | 209 | 88 | 42.27 | 3 | 4.69 | 2 | 1 | 4 |
| 10 | 1346 | MNE | 200 | 225 | 80 | 35.60 | 6 | 5.88 | 3 | 3 | 5 |
| | 877 | BIH | 41 | 206 | 68 | 33.18 | 1 | 5.36 | 3 | 3 | 3 |
| | 1569 | SLO | 103 | 261 | 121 | 46.23 | 3 | 5.52 | 3 | 3 | 5 |
| 11 | 197 | SLO | 200 | 213 | 77 | 36.03 | 1 | 4.65 | 3 | 3 | 5 |
| | 288 | SRB | 80 | 221 | 85 | 38.32 | 3 | 5.23 | 3 | 1 | 5 |
| | 1509 | SLO | 200 | 216 | 87 | 40.07 | 5 | 6.92 | 3 | 3 | 5 |
| | 1450 | CRO | 300 | 233 | 98 | 41.69 | 8 | 5.73 | 3 | 3 | 5 |

HG – homogenous group; AN – accession number; CC – country of collection; AL – altitude of collection site; PH – plant height; EH – ear height; PH/EH – plant and ear height ratio; LBP – lodged and broken plants; Y – yield; GEA – general ear assessment; GC – leaf green colour; KT – kernel type; CRO – Croatia; SRB – Serbia; BIH – Bosnia and Herzegovina; MAC – Republic of North Macedonia; SLO – Slovenia.

CONCLUSIONS

Broad genetic variability preserved in the MRIZP gene bank accessions, which originated from the Western Balkan region, generally considered as a part of European corn-belt, is an exceptional source of desirable traits for enriching breeders' working collections for maize breeding under temperate conditions.

Applied stratification process could be recommended as efficient method for selection of small number of landraces from entire gene-bank collections, allowing the preservation of as much gene pool variability as possible. Finally, this method could provide precise information necessary to develop synthetic populations as an initial material in accordance to short-, medium- and long-term goals of the commercial maize breeding programmes under temperate conditions and fulfil so-called missing places in commercial breeding gene pool (particularly with early-maturing flint genotypes).

As a result, forty landraces with the best *per se* performances important for breeding were selected out of 310 landraces. In particular, twenty-eight early-maturing landraces with flint (AN - 1869, 1890, 594, 773, 1185, 589, 1379, 13, 467, 144, 2144, 1534, 642, 1895, 2006, 2176 and 1945), flint-like (AN - 1381, 871, 2230 and 2236) and intermediate kernel types (AN - 1267, 1960, 846, 1384, 1798, 2033 and 877) could be considered as genotypes of special importance for the commercial breeding programme.

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REFERENCES

- Babić, V., Vančetović, J., Prodanović, S., Anđelković, V., Babić, M., Kravić, N., 2012a. *The identification of drought tolerant maize accessions by the two-step cluster analysis*. Rom. Agric. Res., 29: 53-61.
- Babić, V., Ivanović, M., Babić, M., 2012b. *The origin and evolution of maize and its introduction into South-Eastern Europe*. Field and Vegetable Crops Research, 49(1): 92-104. (In Serbian)
- Babić, V., Vančetović, J., Prodanović, S., Kravić, N., Babić, M., Anđelković, V., 2015. *Numerical classification of western balkan maize drought tolerant landraces*. J. Agr. Sci. Tech., 17(2): 455-468.
- Badu-Apraku, B., Fakorede, M.A.B., 2017. *Advances in genetic enhancement of early and extra-early maize for Sub-Saharan Africa*. Springer International Publishing AG (eBook).
- Böhm, J., Schipprack, W., Mirdita, V., Utz, H.F., Melchinger, A.E., 2014. *Breeding potential of European flint maize landraces evaluated by testcross performance*. Crop Sci., 54(4): 1665-1672.
- Bouchet, S., Bertin, P., Presterl, T., Jamin, P., Coubriche, D., Gouesnard, B., Laborde, J., Charcosset, A., 2017. *Association mapping for phenology and plant architecture in maize shows higher power for developmental traits compared with growth influenced traits*. Heredity, 118(3): 249-259.
- Carena, M.J., 2005. *Maize commercial hybrids compared to improved population hybrids for grain yield and agronomic performance*. Euphytica, 141(3): 201-208.
- Carena, M.J., Pollak, L., Salhuana, W., Denuc, M., 2009a. *Development of unique lines for early-maturing hybrids: Moving GEM germplasm northward and westward*. Euphytica, 170(1-2): 87-97.
- Carena, M.J., Yang, J., Caffarel, C., Mergoum, M., Hallauer, A.R., 2009b. *Do different production environments justify separate maize breeding programs?* Euphytica, 169(2): 141-150.
- Cerović, R., Pajić, Z., Filipović, M., Fotrić-Akšić, M., Radičević, S., Nikolić, D., Đorđević, M., 2014. *Pollen germination and pollen tube growth in ZP maize lines*. Genetika, 46(3): 935-948.
- Curry, H.A., 2017. *Breeding uniformity and banking diversity: The genescapes of industrial agriculture, 1935-1970*. Glob. Environ., 10(1): 83-113.
- Gorjanc, G., Jenko, J., Hearne, S.J., Hickey, J.M., 2016. *Initiating maize pre-breeding programs using genomic selection to harness polygenic variation from landrace populations*. BMC Genomics, 17: 30.
- Gouesnard, B., Dallard, J., Bertin, P., Boyar, A., Charcosset, A., 2005. *European Maize landraces: genetic diversity, core collection definition and methodology of use*. Maydica, 50: 225-234.

- Le Clerc, V., Bazante, F., Baril, C., Guiard, J., Zhang, D., 2005. *Assessing temporal changes in genetic diversity of maize varieties using microsatellite markers*. Theor. Appl. Genet., 110(2): 294-302.
- Maize Research Institute "Zemun Polje". The ECPGR Maize Database. Available at: <<https://www.mrizp.rs/emdb/default-en.htm>>. Accessed on: Feb. 08. 2019.
- Masuka, B., Magorokosho, C., Olsen, M., Atlin, G.N., Bänziger, M., Pixley, K.V., Vivek, B.S., Labuschagne, M., Matemba-Mutasa, R., Burgueño, J., Macrobert, J., Prasanna, B.M., Das, B., Makumbi, D., Tarekagne, A., Crossa, J., Zaman-Allah, M., Biljon, A. van, Cairns, J.E., 2017. *Gains in Maize Genetic Improvement in Eastern and Southern Africa: II. CIMMYT open-pollinated variety breeding pipeline*. Crop Sci., 57(1): 180-191.
- Mikić, S., Kondić-Šipka, A., Brbaklić, L.J., Stanisavljević, D., Čeran, M., Trkulja, D., Mitrović, B., 2017. *Molecular and phenotypic characterisation of diverse temperate maize inbred lines in Southeast Europe*. Zemdirbyste-Agriculture, 104(1): 31-40.
- Mitrović, B., Stojaković, M., Zorić, M., Stanisavljević, D., Bekavac, G., Nastasić, A., Mladenov, V., 2016. *Genetic gains in grain yield, morphological traits and yield stability of middle-late maize hybrids released in Serbia between 1978 and 2011*. Euphytica, 211(3): 321-330.
- Nascimento, W.F., Silva, E.F., Veasey, E.A., 2011. *Agro-morphological characterization of upland rice accessions*. Sci. Agric., 68(6): 652-660.
- Ortiz, R., Taba, S., Tovar, V.H.C., Mezzalama, M., Xu, Y., Yan, J., Crouch, J.H., 2010. *Conserving and enhancing maize genetic resources as global public goods - A perspective from CIMMYT*. Crop Sci., 50(1): 13-28.
- Reif, J.C., Hamrit, S., Hechenberger, M., Schipprack, W., Maurer, H.P., Bohn, M., Melchinger, A.E., 2005. *Trends in genetic diversity among European maize cultivars and their parental components during the past 50 years*. Theor. Appl. Genet., 111(5): 838-845.
- Shimelis, H., Laing, M., 2012. *Timelines in conventional crop improvement: pre-breeding and breeding procedures*. Aust. J. Crop Sci., 6(11): 1542-1549.
- Technow, F., Schrag, T.A., Schipprack, W., Melchinger, A.E., 2014. *Identification of key ancestors of modern germplasm in a breeding program of maize*. Theor. Appl. Genet., 127(12): 2545-2553.
- Vančetočić, J., Božinović, S., Ignjatović-Micić, D., Delić, N., Kravić, N., Nikolić, A., 2015. *A diallel cross among drought tolerant maize populations*. Euphytica, 205(1): 1-16.
- Vega-Alvarez, I., Santacruz-Varela, A., Rocandio-Rodriguez, M., Córdova-Téllez, L., López-Sánchez, H., Muñoz-Orozco, A., Hernández-Bautista, A., 2017. *Genetic diversity and structure of native maize races from Northwestern Mexico*. Pes. Agropec. Bras., 52(11): 1023-1032.
- Worku, M., Makumbi, D., Beyene, Y., Das, B., Mugo, S., Pixley, K., Bänziger, M., Owino, F., Olsen, M., Asea, G., Prasanna, B.M., 2016. *Grain yield performance and flowering synchrony of CIMMYT's tropical maize (Zea mays L.) parental inbred lines and single crosses*. Euphytica, 211(3): 395-409.