

# MICROSTRUCTURAL VARIABILITY OF THE LEMMA ABAXIAL EPIDERMIS IN THE GROUP BROMUS SECALINUS, B. COMMUTATUS AND B. RACEMOSUS

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Abstract. A group of closely related species from the section Bromus: B. secalinus, B. commutatus and B. racemosus were described by characteristics of the abaxial epidermis of lemma. B. commutatus appeared as the most extreme unit against other types. B. secalinus exhibited an intermediate nature between commutatus and racemosus morphs. Many interspecific forms of putative hybrid origin were also put in an ordination space as intermediates.

Key words: Bromus secalinus group, lemma epidermis, variation, taxonomy

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## Introduction

Genesis of the genus Bromus L. has been presented in details by STEBBINS (1981). He considered this genus as an intermediate group between the tribes Festuceae Dumort. and Triticeae Dumort. The genus comprises six sections: Neobromus (Shear). Hitchc., *Ceratochloa* (P. Beauv.) Griseb., Pnigma Dumort., Genea Dumort., Nevskiella (Krecz. et Vved.) Tournay and Bromus. All three species, B. secalinus L., B. commutatus Schrad. and B. racemosus L. are members of the section Bromus Sm. (Cvelev 1976; SMITH 1980). At present, B. secalinus is a rare weed in cereal fields. B. commutatus has smaller distribution in Europe than B. secalinus. B. racemosus has a broad geographical distribution in Europe. B. commutatus and B. racemosus are not easily discriminated. Both species were accurately described by Ammann (1981).Many intermediate forms between both species have also been described. Ammann considered both as subspecies in B. racemosus. Plants expressing more cleistogamic behaviour were noted in B. racemosus and recognised as hybrids B. ×litvinovii Roshev. ex Nevski (CVELEV 1976). Such a change of mating system can fix them at a specific rank. Species from the section Bromus express facultative autogamy. For instance, the level of outcrossing in B. mollis L. reaches 4%

(GRANT 1981). A seasonal cleistogamy was also described in B. carinatus Hook, et Arn. (Harlan 1945). Kosina & Szmidzińska (2000) found that *B. arvensis* L. from the section Bromus is more allogamic than species from the section Genea Dumort., while B. secalinus and B. commutatus are intermediate in mating behaviour. Are these mating differences reflected in plant microstructure? Abaxial epidermis characteristics were successfully applied in taxonomy of many grasses. For instance, lemma quantitative characteristics clustered together B. secalinus and B. racemosus and separated both from *B. commutatus* (KOSINA & Zawerbna 2002).

## Material and methods

The following taxa were cultivated on small plots under equal soil and weather conditions in the grass collection of the first author:

1. Bromus secalinus, twelve accessions marked as Bs1 to Bs11 and Bs;

B. racemosus, two accessions Br1, Br2; 2.

3. Intermediate forms B. racemosus -B. commutatus, four accessions Brc1, Brc2, Brc3, Brc4:

4. *B. commutatus*, one accession Bc

5. Intermediate forms B. commutatus - B. racemosus, three accessions Bcr1, Bcr2, Bcr3.



**Fig. 1.** A minimum spanning tree of the group *Bromus secalinus*, *B. commutatus* and *B. racemosus*. OTUs (species) are scattered in a non-metric multidimensional scaling ordination space (x, y and z axes). OTUs were described by micromorphology of the lemma abaxial epidermis.

The random samples of diaspores (n = 30) were treated according to the completely randomised one-way classification. Varnish replicas of the lemma abaxial epidermis were observed under an Amplival microscope. The following types of short cells were counted per a comparable surface of each accession: duplexes (silica and cork cells), papillae, simple short cells and triplexes. Data were elaborated by taxonomic numerical methods including the Kruskal's non-metric multidimensional scaling according to ROHLF (1994). Results are presented in the form of diagram as the minimum spanning tree.

## **Results and discussion**

Characters of lemma epidermis are significantly negatively correlated, as follows (Pearson correlation):

- duplex *versus* papilla,  $r = -0.60^{***}$ ;
- duplex *versus* simple short cell,  $r = -0.46^*$ . This means that papillae and simple short

cells are morphogenetically linked versus duplexes. The one cytokinesis creating single cell is a different morphogenetic event from that giving duplexes (two cytokineses) and both these kinds of meristemoid activity are interchangeable. The papillae are the main morphogenetic event in the lemma. It was proved that epidermal characteristics of lemma or palea independently discriminate species within the genus Bromus (KOSINA 1997). Then, the data of lemma can be applied here for a taxonomic purpose. A matrix of average taxonomic distances was used in the UPGMA clustering method. Numerical results show that maximum epidermal meristemoidal events (cytokineses) were detected in *B. commutatus*, while minimum in the accession Brc4. This picture, maximum versus minimum, is also well presented in Fig. 1, when the nonmetric multidimensional scaling was applied. Many intermediate forms between extreme accessions, including B. secalinus, prove that this group of species is biologically complex.

A pattern of scattering of OTUs in the ordination space corroborates results obtained earlier for another set of bromegrasses (KOSINA & ZAWERBNA 2002). Autogamy plus facultative allogamy, natural hybridization creating a pool of recombinants, and natural mutations noted in flower organs (KOSINA, unpubl.), all these influence the variability of the group. ACEDO & LLAMAS (1997) expressed an opinion that an intensive longitudinal growth of epidermal cells decreases frequency of short cells in Bromus. In the group of *B. secalinus*, interaccessional differences in frequency of epidermal short cells are of quantitative nature and ranges of variation often overlap (Коснмаńsки 2008). This is also evidenced in gross morphology of spikelet (AMMANN 1981). In the group, many cytogenetic defects such as multipolar anaphases, laggards, bridges and elimination of micronuclei were noted (Skowrońska 2005). These cytogenetic events also caused disorders in the development of caryopsis. This proves that many studied types are developmentally unbalanced because of hybrid origin.

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