

# Abnormal cytokinesis in microsporogenesis of *Brachiaria humidicola* (Poaceae: Paniceae)

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**ABSTRACT.** Microsporogenesis was evaluated in the *Brachiaria humidicola* collection of the Embrapa Beef Cattle Center, represented by 60 accessions. One accession (H121) presented an abnormal pattern of cytokinesis that had never been reported in this genus. Among 900 meiocytes analyzed in the first division, 10.7% underwent precocious and multiple cytokinesis in metaphase I, fractionating the genome and the cytoplasm into two or more parts. The expected cytokinesis after telophase I did not occur. The abnormal meiocytes from the first division entered the second division but the second cytokinesis after telophase II was also abnormal. Among the 857 meiocytes analyzed in the second division, 10.9% presented abnormal, incomplete or total absence of cytokinesis. Dyads and binucleated microspores were recorded among the meiotic products. The use of this accession in the Embrapa breeding program is compromised.

**Key words:** *Brachiaria humidicola*, Breeding program, Cytokinesis, Forage grass, Microsporogenesis

## INTRODUCTION

Microsporogenesis is an ideal process for studying meiotic mutations affecting the steps involved in meiotic reduction of diploid mother cells to form haploid microspores. The pattern of divisions during microsporogenesis is highly controlled and normally yields a predictable end product (Staiger and Cande, 1990).

Numerous mutants that disrupt meiosis at various stages have been reported in different plant species (Gottschalk and Kaul, 1974, 1980a,b; Baker et al., 1976; Golubovskaya, 1979, 1989). In *Brachiaria*, the genus of forage grasses that is most widely cultivated worldwide in the tropics, recent cytological studies have indicated several putative meiotic mutations (Mendes-Bonato et al., 2001, 2003, 2004, 2006a,b; Risso-Pascotto et al., 2002, 2003a,b, 2005; Junqueira Filho et al., 2003; Mendes-Vieira et al., 2005).

After two rounds of chromosome segregation (karyokinesis) and one simultaneous or two successive cytoplasmic divisions (cytokinesis), the final product of male meiosis in flowering plants emerges as a tetrad of haploid microspores in a callose wall. Cytokinesis is accomplished by a typical phragmoplast which is initiated in the spindle midzone during late anaphase and early telophase. The array of parallel phragmoplast microtubules propagates centrifugally and cytokinesis is completed before the next division (Staiger and Cande, 1991).

Several mutants are known to alter normal progression of meiosis and can be correlated with defects in microtubule distribution. One of such mutant, *dy*, has been reported in maize (Staiger and Cande, 1990). In the genus *Brachiaria*, some putative mutations have been reported affecting the pattern of cytokinesis, mainly in *B. humidicola* (Boldrini et al., 2006; Gallo et al., 2007; Calisto V, unpublished data). We report a new pattern of cytokinesis in one accession of this species.

## MATERIAL AND METHODS

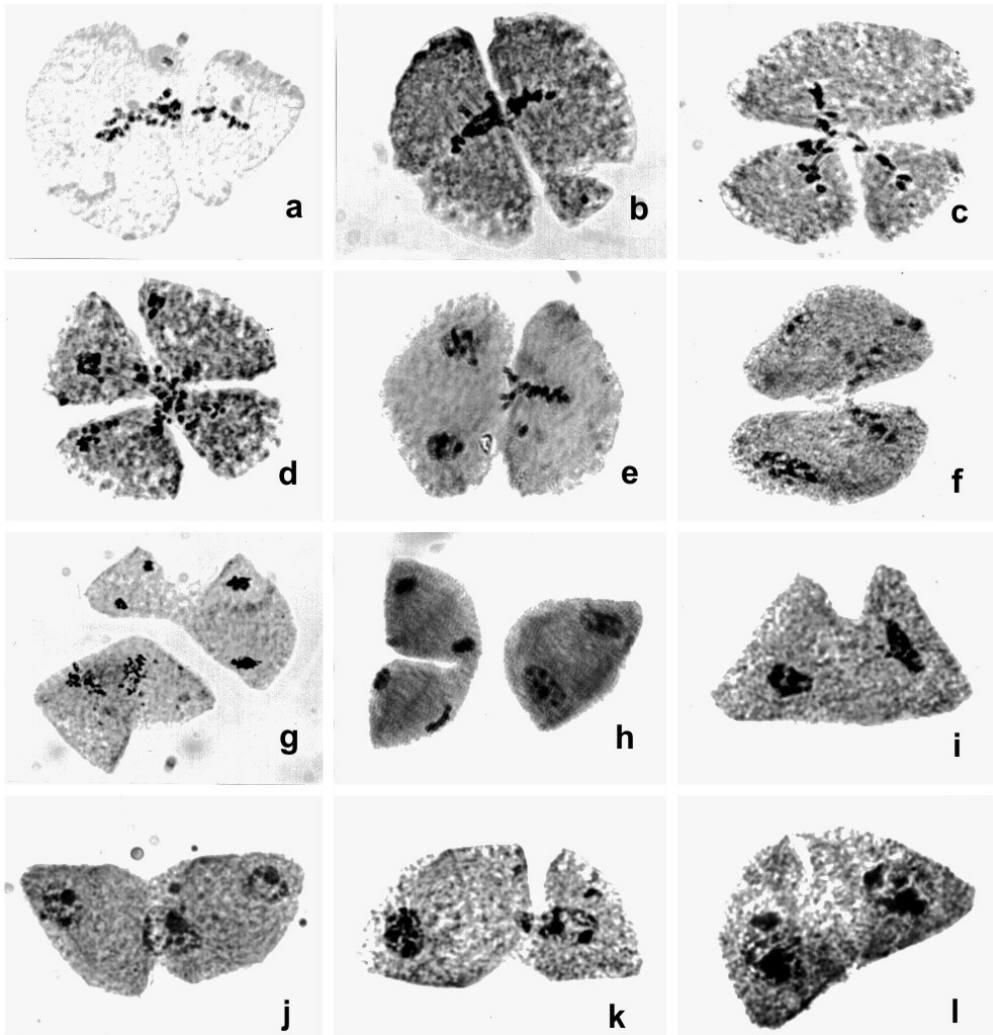
Accessions of *B. humidicola* from the Embrapa Beef Cattle *Brachiaria* germplasm collection (Campo Grande, State of Mato Grosso do Sul, Brazil, originally collected from wild-African savannas in the mid 1980s by CIAT (Colombia) were cytologically analyzed. Site characteristics of the plots in the field at Embrapa Beef Cattle Research Center in Brazil were: climate type Aw: tropical humid savanna; average annual precipitation = 1526 mm; average temperature = 22°C; altitude = 520 m; latitude = 20° 28' S; longitude = 55° 40' W; poor Dark Red Latossol (59% sand; 8% silt; 33% clay; pH = 4.2).

Inflorescences for meiotic study were collected in plots with 16 plants and fixed in a mixture of 95% ethanol, chloroform and propionic acid (6:3:2) for 24 h, transferred to 70% alcohol and stored under refrigeration until use. Microsporocytes were prepared by squashing and staining with 0.5% propionic carmine. Photomicrographs were made with a Wild Leitz microscope using Kodak Imagelink - HQ, ISO 25 black and white film.

## RESULTS AND DISCUSSION

The *B. humidicola* collection at Embrapa Beef Cattle Center is represented by 60 accessions. Among them, 58 accessions have had their meiotic behavior evaluated. One accession (H121) presented an abnormal pattern of cytokinesis that had never been reported

in this genus: 900 meicytes in the first division were analyzed. Among them, 96 meicytes (10.7%) underwent precocious and multiple cytokinesis in metaphase I, fractionating the genome and the cytoplasm into two or more parts (Figure 1a-f). The expected cytokinesis after telophase I did not occur, thus abnormal meicytes from the first division entered the second division (Figure 1g,h). The second cytokinesis after telophase I in these meicytes was also abnormal. Among the 857 meicytes analyzed in meiosis II, 94 (10.9%) presented abnormal incomplete (Figure 1i-l) or total absence of cytokinesis. Among meiotic products, dyads and released binucleated microspores were recorded.



**Figure 1.** Aspects of abnormal cytokinesis in accession H121. **a-c.** Microsporocytes in metaphase I divided by precocious cytokinesis. Micronuclei are isolated as microcytes. **d.** Metaphase I with the genome fractionated into four cells. **e,f.** Microsporocyte that underwent precocious cytokinesis, showing asynchrony between cells. **g,h.** Microsporocytes in the second division resulting from precocious cytokinesis in the first division. **i-l.** Different aspects of abnormal cytokinesis in binucleated microspores (Magnification 400X).

The accession of *B. humidicola* has  $2n = 54$  chromosomes. Based on the meiotic behavior in normal meiocytes, we suggest that it is a polyploid accession ( $2n = 9x = 54$ ), derived from  $x = 6$ . Chromosome number counts in the *B. humidicola* collection at Embrapa Beef Cattle Center revealed accessions with  $2n = 36, 42,$  and  $54$  chromosomes (Boldrini KR, unpublished data). Among the accessions with  $2n = 54$  chromosomes, another accession (H022) also presented an abnormal pattern of cytokinesis (Calisto V, unpublished data) similar to the abnormality reported in H121. However, in H022 the abnormal cytokinesis occurred only in meiocytes in which chromosome associations were disrupted at the end of diakinesis, separating the chromosomes by desynapsis; the  $54$  univalents were aligned forming a wide metaphase plate. In H121, normal cytokinesis after telophase I was also absent and the second cytokinesis after telophase II occurred normally. Cytological evidence of precocious cytokinesis was not found in the H121.

In another accession of *B. humidicola* (H003) with  $2n = 7x = 42$ , also derived from  $x = 6$ , another abnormal pattern of cytokinesis was recorded (Boldrini et al., 2006). In this accession, the first cytokinesis occurred after telophase II, generating dyads with binucleated microspores, which initiated the second cytokinesis by invagination, giving rise to four normal microspores, after release from the callose wall. In H121 and in H022 (Calisto V, unpublished data), both with  $2n = 54$  chromosomes, cytokinesis was initiated marginally in metaphase I and migrated to the center, similar to an invagination process. Incomplete cytokinesis leading to  $2n$  gamete formation was recorded in another accession of *B. humidicola* (H047),  $2n = 36$  chromosomes (Gallo et al., 2007). In this accession, the first cytokinesis was initiated in the center of the cell after telophase I but did not get to the borders, allowing rejoining of the nucleus after prophase II. The number of microsporocytes affected by abnormal cytokinesis recorded in these accessions of *B. humidicola* was always small. In accession H022, with a similar abnormality, 16.8% of the cells were affected.

A large number of genes, generally dominant, which are stage, site- and time-specific are involved in the control of meiosis (Gottschalk and Kaul, 1974, 1980a,b; Baker et al., 1976; Golubovskaya, 1979, 1989). Among genes acting in the meiotic process, those responsible for the partitioning of the cytoplasm after nuclear division play an important role in the formation of viable gametes. The timing of cytokinesis varies among angiosperms. In most monocotyledons, cytokinesis is successive, i.e., one partitioning of the cytoplasm occurs after telophase I and another after telophase II, so that there is a distinct dyad stage. However, in most dicots it is simultaneous and occurs after telophase II (Peirson et al., 1996). In higher plants, cytokinesis during mitosis is a genetically controlled multistep process. At least three cellular components play important roles in this process: i) the Golgi apparatus produces secretory vesicles and synthesizes the cell wall polysaccharides; ii) Golgi-derived vesicles fuse to form a cell plate, and iii) the cytoskeleton required for phragmoplast formation and expansion controls the cell division planes. Other cellular components, including the endoplasmic reticulum, intermediate filaments, calmodulin, and myosin may also play important roles in cytokinesis (Staehelin and Hepler, 1996). During mitosis in higher plants, a cortical ring of microtubules, called 'the preprophase band', marks the site where the cell plate will be formed, determining the division pattern. Meiosis, on the other hand, lacks the preprophase band, although the division pattern seems to be accurately controlled. Cytokinesis-defective mutants have been characterized in several species of higher plants during mitosis and meiosis (Beadle, 1932; Peirson et al., 1996; Hülkamp et al., 1997; Nickle and Meinke, 1998; Boldrini et al., 2006).

Absence of cytokinesis has been recorded for other *Brachiaria* species (Risso-Pascotto et al., 2003a; Utsunomiya et al., 2005). In these cases, the failure of cytokinesis occurred after telophase I or telophase II, generating balanced  $2n$  gametes. In H121 and H022, cytokinesis occurred precociously during metaphase I. We suggest that the genetic control of cytokinesis in these meiocytes is activated very early and is not synchronized with karyokinesis, generating unbalanced gametes. Based on the cytological analysis of the *B. humidicola* collection, we suggest that this species has a greater tendency for abnormal cytokinesis than do other previously analyzed *Brachiaria* species (Mendes-Bonato et al., 2002, 2006a; Utsunomiya et al., 2005). Several putative meiotic mutations have been described in the genus *Brachiaria* (Mendes-Bonato et al., 2001, 2003, 2004, 2006b; Risso-Pascotto et al., 2002, 2003a, 2005; Mendes-Vieira et al., 2005) and also in the post-meiotic process (Junqueira Filho et al., 2003; Mendes-Bonato et al., 2004) which could represent putative mutations, suggesting that these genes were incorporated in the gene pool of the genus during its evolutionary process.

Some promising apomictic accessions of *B. humidicola* are under careful agronomic and grazing evaluation in hopes of selecting new cultivars. This species is well adapted to poorly drained and infertile acid soils (Keller-Grein et al., 1996), for which very few options are available, thus the urgent demand for improved cultivars. The occurrence of precocious cytokinesis detected in this accession affects pollen viability and unbalanced gametes are generated by partitioning of the genome in metaphase I. Other accessions of this species may be better progenitors in intra- and interspecific hybridization as pollen donors. However, the frequency of meiocytes affected by precocious cytokinesis in H121 (about of 10%) may not be enough to require that this accession be discarded from the breeding program. Other abnormalities due to polyploidy ( $2n = 9x = 54$ ) were also detected: precocious chromosome migration to the poles in metaphase I (55.7%) and metaphase II (55.6%), and laggards in anaphase I (82.5%) and anaphase II (58.7%), which generated micronuclei in telophase I (44.9%) and II (11.4%).

Polyploidy in *Brachiaria* is correlated with apomixis, which bypasses meiosis in the megagametogenesis process. The embryo-sac is formed by parthenogenesis of a somatic cell, thus the embryo is maternal. But for seed development, the secondary nuclei of the embryo sac need to be fertilized by a male gamete - pseudogamy. Accessions with a high frequency of meiotic abnormalities due to polyploidy, which severely impair pollen viability, need to be discarded early in the breeding program to avoid passing on defective genes to the progenies. Until now the on-going hybridization program in *Brachiaria* has involved intra- and interspecific crosses only between tetraploid ( $2n = 4x = 36$ ) genotypes derived from  $x = 9$ . The  $2n = 54$  chromosome pool is still impervious to breeding due to a lack of compatible sexual source for crossing. Interploid crosses have been unsuccessful in this genus (Hacker, 1988). Therefore, accession H121 cannot be used in crosses not only due to its high level of ploidy, but also due to its abnormal cytokinesis and differences in its basic chromosome number ( $x = 6$ ).

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