

Induced mitotic homologous recombination by the babesicide imidocarb dipropionate in *Aspergillus nidulans* diploid cells

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ABSTRACT. Imidocarb dipropionate (IMD) is a chemotherapeutic agent prescribed for the treatment and control of babesiosis; it is known to be a nucleic acid synthesis inhibitor. Although it is an effective babesicide, there are reports of persistent IMD residues retained at high levels in edible tissues of cattle, swine and sheep, raising concerns about potential effects on humans. Since the carcinogenic potential of a chemical compound can be assessed through its effect on the homologous recombination, we investigated whether IMD is recombinogenic in *Aspergillus nidulans* diploid cells and whether it is capable of inducing homozygosis in genes that were previously heterozygous. This analysis was done with a homozygotization assay applied to a heterozygous diploid strain of *A. nidulans*. IMD used at non-toxic concentrations (2.5 to 10.0 μ M) was recombinogenic, demonstrated by homozygotization indices higher than 2.0 for diploid markers. A diploid homozygous for genetic markers from chromosomes I and II was also produced. Since DNA replication blockers that induce DNA strand breaks have been classified as potent inducers of homologous recombination,

the recombinogenic potential of IMD may be due to induction of recombinational repair.

Key words: *Aspergillus nidulans*; Homozygotization index; Babesiosis; Genotoxicity; DNA synthesis inhibitors

INTRODUCTION

Imidocarb dipropionate (IMD; N,N'-bis[3-(4,5-dihydro-1H-imidazol-2-yl)phenyl]urea dipropionate) is a chemotherapeutic agent belonging to the family of the carbanilide derivatives commonly prescribed in the treatment and prophylaxis of babesiosis, which is a tick-transmitted disease caused by a protozoa of the genus *Babesia*. In addition to being a well-recognized disease of veterinary importance in cattle, horses, dogs, and other domestic animals, babesiosis has been characterized as a human infectious disease, especially in persons with underlying immunosuppressive conditions, including HIV coinfection, malignancy, immunosuppressive medication, and splenectomy (Froberg et al., 2004; Stowell et al., 2007; Krause et al., 2008; Wang et al., 2009; Haapasalo et al., 2010).

Babesia species are obligate parasites of red blood cells and belong to the phylum Apicomplexa, together with organisms that cause malaria (*Plasmodium* sp) and toxoplasmosis (*Toxoplasma gondii*). Whereas *B. bovis*, *B. caballi* and *B. canis* are species causing babesiosis in cattle, horses and dogs, respectively, human babesiosis is primarily caused by *B. microti*, a rodent species, and *B. divergens*, a cattle species. While symptoms of bovine babesiosis include fever, anemia, anorexia, hemoglobinuria, and death, in humans, the *Babesia* parasitic species may cause a malaria-like syndrome including fever, hemolysis, hemoglobinuria, renal failure, acute respiratory distress, and death (Benach and Habicht, 1981; Uilenberg, 2006; Vial and Gorenflot, 2006; Hasle et al., 2010).

Many drugs have been used over the years as therapeutic or prophylactic agents against infection for hemoprotozoa in domestic animals, of which IMD is currently considered the most efficacious and safest of all available babesicides (Zintl et al., 2003; Vial and Gorenflot, 2006). In addition to providing protection against clinical diseases in domestic animals, IMD also allows a sufficient level of infection for immunity development, which represents an attractive approach of the drug, especially in areas where babesiosis is endemic (Zintl et al., 2003; Vial and Gorenflot, 2006). Despite its efficacy, previous studies have reported that IMD residues are retained in high and persistent amounts in animal edible tissues, such as liver and kidney of cattle, swine and sheep (Lai et al., 2002; Belloli et al., 2006; Wang et al., 2009). The literature indicates two explanations for the retention of IMD in animal tissues after a therapeutic dose: a) the resistance of the drug to biotransformation processes, and b) IMD's higher capacity to bind to nucleic acids (DNA and RNA) than to other macromolecules, such as hemoglobin and serum albumin, providing a cellular mechanism to account for IMD retention in animal tissues such as the liver, which has a higher DNA content (Coldham et al., 1995; Moore et al., 1996; Belloli et al., 2006).

Since the information on genotoxic effects of IMD is scarce and since the carcinogenic potential of a chemical compound may be assessed through its effect on homologous recombination as well (Arnaudeau et al., 2000; Miyamoto et al., 2007; de Castro-Prado et al., 2009; Franco et al., 2010), this study aimed to investigate whether IMD is recombinogenic in

Aspergillus nidulans diploid cells and capable of inducing the homozygosity of genes previously present in the heterozygous condition.

A. nidulans is a filamentous fungus extensively used for the study of mitotic crossing-over, as its cells pass the greater part of their cell cycle in the G2 phase (Bergen and Morris, 1983). Due to the fact that chromosomes in this phase are in duplicate, they significantly favor mitotic recombination. *A. nidulans* diploid strains have been used to evaluate the recombinogenic potential of chemical compounds such as the trypanosomicidal agent known as benzimidazole (Kaneshima and Castro-Prado, 2005). The recombinogenic potential of IMD was determined using an *A. nidulans* diploid strain that was heterozygous for several genetic markers, as well as the homozygotization assay (Cardoso et al., 2010).

MATERIAL AND METHODS

Strain and culture media

The diploid UT448//A757 strain of *A. nidulans*, carrying markers on chromosomes I, II and IV (Table 1) was prepared by Roper's (1952) technique. Mutant alleles of importance in this study were the following: *w* and *y*, white and yellow conidia, respectively, and *riboA1*, *pabaA124*, *biA1*, and *methA17*, nutritional requirement for riboflavine, p-aminobenzoic acid, biotin, and methionine, respectively. Since the diploid strain is heterozygous for the nutritional markers, it can grow in minimal medium (MM), consisting of Czapek-Dox medium, supplemented with 1% (w/v) glucose. On the other hand, when growing in complete medium (CM), the diploid strain may originate auxotrophic mitotic segregants, which are recognized as normally growing yellow, green or white sectors on UT448//A757 diploid green colonies. CM consisted of MM supplemented with 10.0 g/L glucose, 2.0 g/L peptone, 2.0 g/L yeast extract, 1.0 g/L hydrolyzed casein, 4.0 mg/L inositol, 2.0 mg/L choline chloride, 2.0 mg/L pantothenic acid, 1.0 mg/L nicotinic acid, 1.0 mg/L riboflavin, 0.1 mg/L p-aminobenzoic acid, 0.5 mg/L folic acid, 0.5 mg/L pyridoxine, 0.2 mg/L thiamine, and 0.002 mg/L biotin, whereas supplemented medium (SM) consisted of MM supplemented with all the nutritional requirements of the master strains except one, in each medium type. Solid medium contained 1.5% (w/v) agar.

Table 1. Genotype and origin of *Aspergillus nidulans* strains.

Strain	Genotype	Origin
A757	<i>yA2</i> , <i>methA17</i> , <i>pyroA4</i>	FGSC*
UT448	<i>riboA1</i> , <i>pabaA124</i> , <i>biA1</i> , <i>AcrA1</i> , <i>wA2</i>	Utrecht, Netherlands

riboA1 = riboflavin; *pabaA124* = p-aminobenzoic acid; *biA1* = biotin; *methA17* = methionine; *pyroA4* = pyridoxine. Conidia color: *wA2* = white; *yA2* = yellow. *AcrA1* = resistance to acriflavine. *FGSC = Fungal Genetic Stock Center, University of Kansas Medical Center, Kansas, MO, USA.

Drug treatment

Filter-sterilized aqueous solutions of IMD, C₂₅H₃₂N₆O₅, FW 496.56, CAS No. 55750-06-6, 99.8% pure; Riedel-de Haën, C.O.O. Germany) were added to molten MM. Non-cytotoxic concentrations of IMD (2.5, 5.0 and 10.0 µM), which showed bovine DNA-binding capacity (Moore et al., 1996), were used for the homozygotization assay. In the case of toxicity

measurements, UT448//A757 diploid colony diameters were determined six days after incubation, at 37°C. The rates in the presence (treatment) and in the absence (control) of babesicide were compared by one-way analysis of variance and by the Bonferroni post-test, for $P < 0.05$ (data not shown). Benznidazole, the antichagasic agent ($C_{12}H_{12}N_4O_3$, FW 260.25, 99.8% pure; Roche), previously characterized as recombinogenic in human blood lymphocytes, human hepatoma cell line (Hep G2), and in *A. nidulans* diploid cells, was used as a positive control (Santos et al., 1994; Kaneshima and Castro-Prado, 2005).

Homozygotization assay

Colonies of the UT448//A757 diploid strain of *A. nidulans* (Figure 1A) were grown on Petri plates containing MM (negative control), MM + benznidazole (100 μ M, positive control) and MM + IMD (2.5, 5.0 and 10.0 μ M, treatment). These plates were incubated for six days at 37°C and then visually inspected for diploid sectors arising on the diploid strain colonies. The treatment (MM + IMD) produced nine morphologically identifiable diploid sectors, which were homozygous (+/+) or heterozygous (+/- or -/+) for nutritional markers, but they were never recessive homozygotes (-/-), since the latter cannot grow in MM. The untreated (negative control) and the treated diploid strains were purified in MM, individually transferred to the CM plates and afterwards processed by spontaneous haploidization. The haploidization process, which is the loss of one member of each chromosome pair through successive mitotic divisions, results in the haploid condition of the nuclei. After haploidization, each diploid produced haploid mitotic segregants (Figure 1B and C), which were purified in CM and then evaluated for mitotic stability in CM + benomyl (0.2 μ g/mL). The mitotically stable haploid segregants at the final stage were the only ones selected for the recombinogenesis test (Figure 1D). Such segregants were individually transferred to different SM for their phenotypic analyses. The mitotic crossing-over causes homozygotization of heterozygous-conditioned genes. If IMD induces mitotic crossing-over in the original diploid strain, only heterozygous (+/- or -/+) or homozygous (+/+) diploids will develop in MM and the nutritional markers will segregate among the haploids in the proportion of 4+ to 2-. However, if the drug fails to induce crossing-over, the proportion will be 4+ to 4-. This is due to the fact that the initial selection process limits the growth of -/- diploids. The ratio of prototrophic to auxotrophic segregants is described by the homozygosity index (HI) (Pires and Zucchi, 1994), or rather, an HI equal to or higher than 2.0 indicates the recombinogenic effect of the babesicide drug. The recombinogenic potential of IMD was assessed by comparing the homozygotization indices of the nutritional markers with the Yates corrected chi-square test, with contingency tables, $P < 0.05$.

RESULTS

The recombinogenic potential of IMD was assessed by determining the HI values of the UT448//A757 diploid strain nutritional markers. HIs for the *ribo*, *paba*, *bi*, and *meth* genes were statistically significant in diploid strains treated with IMD in comparison to the untreated diploid strains (negative control) (Table 2). The UT448//A757 original diploid strain produced eight prototrophic diploids with green ($y+//y$) (Figure 1B and C) and one with yellow conidia ($y//y$, designated D5) when growing in MM in presence of 2.5, 5.0 or 10.0 μ M IMD. The treatment in MM does not allow the development of auxotrophic diploids, specifically those

that are homozygous for the nutritional markers *ribo*, *paba*, *bi*, and *meth*. On the other hand, diploids homozygous for the conidia color markers (*y* or *w*) can be produced and selected in MM, since they are prototrophs.

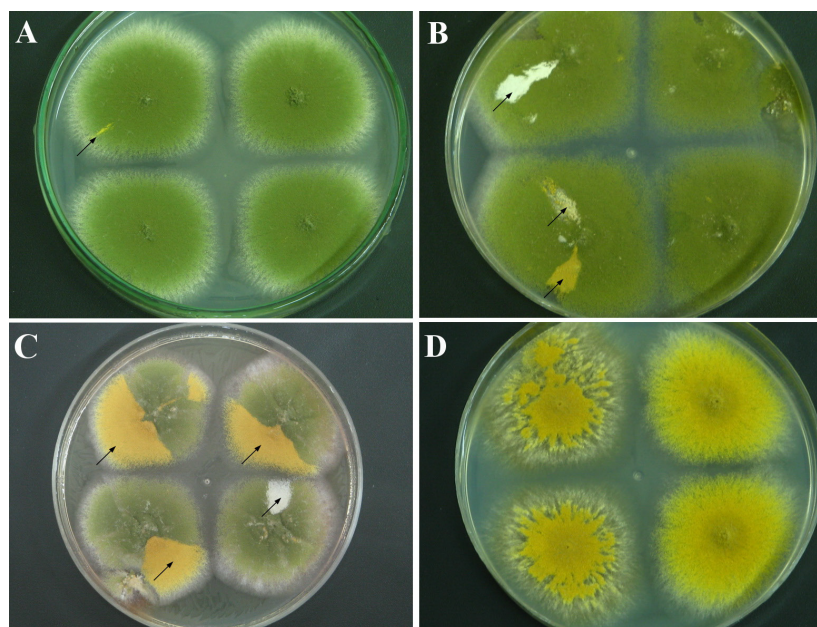


Figure 1. Mitotic instability of imidocarb dipropionate (IMD)-treated diploid strains. **A.** Original UT448//A757 diploid strain growing in the absence of IMD. **B.** and **C.** Diploids obtained respectively with 2.5 and 10.0 μM IMD. **D.** Haploid (right) and aneuploid (left) segregants derived from diploids obtained with 5.0 μM IMD. The arrows indicate the origin of white and yellow mitotic segregants (sectors).

Table 2. Homozygotization index (HI) values for markers from UT448//A757 diploid strain after treatment with 10, 5 and 2.5 μM of imidocarb dipropionate (IMD).

Drug	Concentration (μM)	<i>ribo</i>		<i>paba</i>		<i>bi</i>		<i>meth</i>	
		<i>ribo</i> ⁺ / <i>ribo</i>	HI	<i>paba</i> ⁺ / <i>paba</i>	HI	<i>bi</i> ⁺ / <i>bi</i>	HI	<i>meth</i> ⁺ / <i>meth</i>	HI
Negative control	-	129/101	1.3	123/107	1.1	125/105	1.2	128/102	1.3
Positive control	100	160/55	2.9*	159/56	2.8*	164/51	3.2*	nd	nd
IMD	10	239/103	2.3*	262/80	3.3*	278/64	4.3*	247/95	2.6*
IMD	5	369/168	2.2	369/168	2.2*	369/168	2.2	448/89	5.0*
IMD	2.5	434/166	2.6*	435/165	2.6*	436/164	2.7*	476/124	3.8*

ribo = riboflavin; *paba* = p-aminobenzoic acid; *bi* = biotin; *meth* = methionine. Negative control = diploids did not treat with imidocarb or benzimidazole. Positive control = diploids treated with 100 μM benzimidazole. nd = not determined. *Significantly different from the negative control (contingency table, Yates corrected chi-square test, $P < 0.05$).

Diploid strains obtained with 10.0 μM IMD showed homozygotization indices greater than 2.0 for *ribo*, *paba*, *bi*, and *meth* genes. The phenotypic analysis of haploid segregants derived from diploid D5, with yellow conidia, showed that although D5 was heterozygous for

ribo and *paba* genes, it was homozygous for two other genes, *bi* and *meth*, since auxotrophic *bi* and *meth* segregants were not recovered among their haploid-derived mitotic segregants (data not shown). As mitotic crossing-over is a process that leads to the homozygosity of all genetic markers distal to the position of exchange, we can suppose that diploid D5 was indeed, a mitotic recombinant for chromosomes I and II (Figure 2).

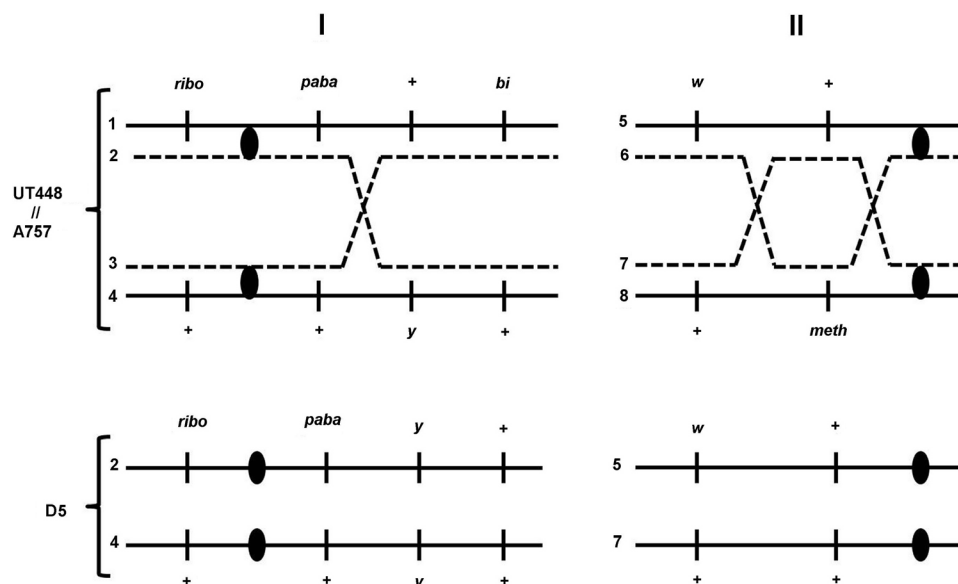


Figure 2. Schematic representation of diploid D5, obtained with 10.0 μM imidocarb dipropionate, generated by a single crossing-over on chromosome I as well as a double crossing-over on chromosome II, followed by the segregation of chromatids 2 (recombinant), 4 (paternal), 5 (paternal), and 7 (recombinant) to the same mitotic pole. For abbreviations, see Table 2.

Diploids obtained with 5.0 and 2.5 μM IMD concentrations also showed HI values greater than 2.0 and significantly different from the HI values for the negative control (Table 2). The recombinogenic action of IMD in *A. nidulans* diploid cells was thus observed for the three drug concentrations tested.

DISCUSSION

The homozygotization assay was employed to assess the ability of IMD to induce mitotic recombination in *A. nidulans* diploid cells. Through the selection and haploidization of heterozygous or homozygous diploid strains obtained after IMD treatment, a greater number of prototrophic rather than auxotrophic segregants were obtained, which increased the HI values. The recombinogenic potential of IMD, at non-toxic concentrations, was demonstrated in the current study by two important results: a) HI values higher than 2.0 and statistically significant for diploid markers obtained with 2.5 to 10.0 μM IMD, and b) the production of a prototrophic diploid (D5) homozygous for *y*, *bi* (from chromosome I) and *meth* genes (from chromosome II) (Tables 1 and 2, Figure 2). *A. nidulans* diploid strains, homozygous for nu-

tritional and conidial color markers, have been previously obtained by treatment with DNA synthesis inhibitors, such as cisplatin and cytosine arabinoside (Kufe et al., 1980; Yarema et al., 1995; Miyamoto et al., 2007). In fact, DNA replication inhibitors, including those that induce the formation of DNA double-strand breaks, have been characterized as potent inducers of homologous recombination (HR) (De Silva et al., 2002; Ludin et al., 2002; Helleday, 2010).

The first association between homologous recombination and tumorigenesis was provided by the molecular analysis of familial retinoblastoma, in which somatic recombination contributes to as many as 75% of the loss of heterozygosity (LOH) events at the retinoblastoma locus. Since then, LOH by mitotic recombination has been observed in other human malignancies and classified as an important contributor to cellular carcinogenesis (Lasko et al., 1991; Rousseau-Merck et al., 1999; Moynahan and Jasin, 2010). A recent study including 113 neurofibromatosis type 1 (NF1) patients revealed that LOH was detected in 25% of the 518 dermal neurofibromas analyzed. The mitotic recombination, generated by a single crossing-over located between the centromere and the *NF1* gene, was one of the major mechanisms causing LOH in such neurofibromatosis type 1 patients (Garcia-Linares et al., 2011).

Cancer is a multistep process in which a series of genetic or epigenetic events leads to the transformation of a normal cell into a cancerous one. Alterations in tumor suppressor genes, which have the normal physiological role of delaying the cell division process, have been described as relevant in the development of neoplasia (Velasco et al., 2008; Szewczuk et al., 2009). According to Knudson's hypothesis (1993), the heterozygous state of a mutation-bearing cell ("first hit") could be lost when it is exposed to cancer-promoting events ("second hit"). Chemical compounds may act as tumor promoter agents in heterozygous cells of a defective tumor suppression gene through the induction of somatic HR, a process involved in the repair of naturally occurring DNA injuries arising during the S phase of the cell cycle (Knudson, 1993; Arnaudeau et al., 2000; Helleday, 2003).

HR repair requires a homologous sequence to act as a template for the repair event. In the late S or G₂ phases of the cell cycle, such template may be provided by the presence of an identical sister chromatid. This process is error-free and restores the DNA sequence as it was prior to the damage. On the other hand, when the homologous donor is the homologous chromosome, the HR repair has the potential to lead to LOH of parental markers. In fact, somatic recombination between homologous chromosomes leads to LOH of the loci distal to the HR event if the recombinant sister chromatids segregate to the opposite mitotic poles (Johnson and Jasin, 2001; Helleday, 2003; Moynahan and Jasin, 2010).

In addition to displaying structural similarities to compounds with known ability to bind to DNA, such as the cellular polyamines spermine and spermidine (Lambros et al., 1977; Bacchi et al., 1981; Moore et al., 1996), IMD has been described as a potent inhibitor of nucleic acid synthesis. At low concentrations, IMD has been shown to inhibit over 50% of [³H] hypoxanthine incorporation into nucleic acids of *B. bovis in vitro* (Nott et al., 1990). Taken into account the recombinogenic potential of the DNA synthesis blockers, the recombinogenic effect of IMD, shown in the current study, may be associated with the induction of the recombinational repair of DNA strand breaks induced by this babesicide.

The literature describes that a high and persistent concentration of IMD remains in the animal body, especially in the liver and kidney of ruminant species. The "reservoir effect" of the drug in animal bodies has been associated with the presence of detectable IMD concentrations in the milk of ewes and does up to 10 days after the drug injection (Moore et al., 1996;

Lai et al., 2002). Additionally, Lai et al. (2002) found high and long-lasting concentrations of IMD in the brain of sheep and goats, showing that the drug may diffuse across the blood-brain barrier, providing evidence for possible IMD neurotoxic effects.

IMD is characterized as a potent inducer of mitotic recombination (present study). Chemical compounds, such as cisplatin and cytosine arabinoside, previously described as recombinogenic in mammalian cells, were also classified as recombinogenic in the *A. nidulans* diploid strain (Beaula Helen and Subramanyan, 1991; Arnaudeau et al., 2000; Miyamoto et al., 2007). Since mitotic HR may trigger neoplasms, this study highlights the need for further investigation of the genotoxic potential of IMD in mammalian cells, especially due to the long elimination time of IMD from milk and animal tissues, which may pose a food risk to humans.

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