

Oxidative Response and Micronucleus Centromere Assay in HEp-2 Cells Exposed to Fungicide Iprodione

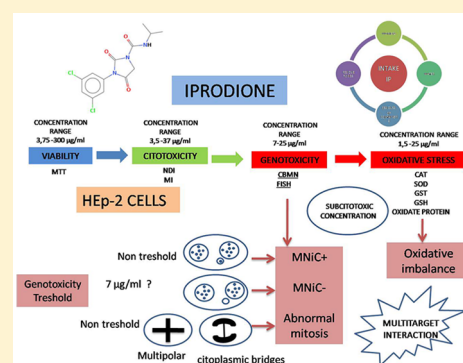
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ABSTRACT: The fungicide agents are a key component in the fruits and vegetables production. The Iprodione residues are one of the pesticide more frequently found in food products. The available data about the cytotoxicity of iprodione and its metabolites are scarce and do not allow characterization of its genotoxic potential and define the risk assessment. The human larynx epidermoid carcinoma cell line (HEp-2) has been shown to be sensitive to the toxic effects of xenobiotics of different origin and have been often used in cytotoxicity and genotoxicity studies. The purpose of this paper is to evaluate the induction of genotoxicity and the role of oxidative stress in HEp-2 cell line by exposure to the IP. The MTT test for viability resulted in CL_{50} 85.86 (77.05–95.68) $\mu\text{g}/\text{mL}$ of Iprodione. On the basis of this result, we proceeded to expose the cells to the sublethal concentrations (below the CL_{50}) during 24 h to analyze the mitotic index and nuclear division index in order to determine the subcytotoxic concentrations of IP which the genotoxicity was evaluated. The subcytotoxic concentrations of 7, 17, and 25 $\mu\text{g}/\text{mL}$ IP induced aneugenic effects as micronuclei centromere positive whereas 17 $\mu\text{g}/\text{mL}$ was a threshold for centromere negative micronuclei induction in HEp-2 cells. The abnormal mitosis was induced for exposition of Hep-2 cells to the three concentrations. According to the result obtained, cytotoxicity and genotoxicity oxidative stress studies were performed in 1.5, 7.0, and 25 $\mu\text{g}/\text{mL}$ of IP. The results showed that the GSH intracellular content, the SOD activity and the levels of oxidative damage of the proteins were affected lead to redox imbalance. The decreased in the SOD activity and protein oxidation were in according to the result obtained to genotoxicity, suggesting that different biological targets could be affected.



INTRODUCTION

The application of fungicide agents constitutes a key component in the fruits and vegetables production during seeds treatment and vegetal growth and the stage of postharvest storage and distribution.¹ As a result of this practice, some potential issues can be identified, for instance, (1) the presence of fungicide residues in fresh or processed food. Most of the fungicide residues can remain stable in food, although the risk of people's consumption is related with the processing factors.^{2,3} (2) The contact of applicators with the fungicide agent. This contact should be reduced by applying safety measures in order to prevent the fungicide from entering the organism.^{4,5} (3) The environmental pollution caused during the fungicide production and application, and the containers disposal.^{6–8}

Studies carried out by the Environmental Working Group have detected fungicide residues in food products for babies, commercialized by Gerber (Fremont, MI), Heinz (Pittsburgh, PA), and Beech-Nut (St. Louis, MO).⁹ Furthermore, these

studies show that one of the pesticide residues more frequently found and with greater content of such substances is iprodione (IP).^{10–12} Studies conducted in animals proved that IP shows low acute toxicity. However, it is classified into the B2 group as “likely” carcinogenic in humans, based on evidence from the induction of liver tumors in mice of both sexes and in male rats.¹³ Studies in mice show that the exposure to IP induces the activity of microsomal enzymes, the proliferation of hepatocytes, and hepatomegaly.^{14,15} In addition, other findings show that it is an endocrine disruptor that causes reproductive abnormalities in Sprague–Dawley rats.¹⁶ Other studies indicate that dicarboximides fungicides, such as IP and vinclozolin, induce lipid peroxidation by means of the action of reactive oxygen species in fungi, but its effect over mammalian cells is not yet clearly studied or defined.^{14,17} The OCDE commission implemented the regulation (EU)

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concerning the nonrenewal of approval of the active substance iprodione in the 2017 year, because available data about the cytotoxicity of this component and its metabolites are scarce and do not allow to characterize its genotoxic potential and residues definition for risk assessment.¹⁸

It is known that genotoxicity is result of the direct interaction with the DNA or other indirect mechanisms involved interactions with non-DNA targets. Genotoxicity induces abnormalities in cells such as micronuclei (MNi), nuclear buds (NB), and nucleoplasmic bridges (NPB), observed in interphase, as result of loss and break chromosome, dicentric chromosome, and gene amplification, respectively. The cytokinesis-block micronucleus cytome assay (CBMN) allows measure the cytotoxic/cytostatic and genotoxic effects in culture human cells. Abnormalities observed in mitosis include chromosomal fragment, chromosome lagged, anaphasic/telophasic bridges and multipolar divisions. The human larynx epidermoid carcinoma cell line (HEp-2) have been shown to be sensitive to the toxic effects of xenobiotics of different origin and have been often used in citotoxicity and genotoxicity studies.^{19–24}

In view of the foregoing, it can be concluded that there is not enough information on the literature that allows defining the IP genotoxic and oxidative potential in human cells. Therefore, the purpose of this paper is to evaluate the induction of genotoxicity and the role of oxidative stress in HEp-2 cells by exposure to the IP.

MATERIALS AND METHODS

Reagents. Iprodione, 3-(3,5-dichlorophenyl)-N-(1-methylethyl) 2,4-dioxo-1-imidazoline-carboximide (C₁₃H₁₃C₁₂N₃O₃), CAS No. 36734-19-7 was purchased from Sigma-Aldrich (CAS No. 36734-19-7), modified eagle's medium (MEM), MEM vitamin solution, MEM nonessential amino acid solution, and 0.05% trypsin-EDTA, were purchased from Laboratorio Micro Vet SRL (Buenos Aires, Argentina). Fetal bovine serum (FBS) was obtained from BIO-NOS (Buenos Aires, Argentina). The 1-chloro-2,4-dinitrobenzene (CDNB) (CAS No. 97-00-7), 2,4-dinitrophenylhydrazine (DNPH) (CAS No. 119-26-6), 3-(4,5-dimethyl-thiazol-2-yl)-2,5-diphenyl-tetrazolium bromide (MTT) (CAS No. 298-93-1), 5,5-dithiobis(2-nitrobenzoic acid) (DTNB) (CAS No. 69-78-3), cytochalasin B (CAS No. 14930-96-2), glutathione (GSH) (CAS No. 70-18-8) were purchased from Sigma-Aldrich, Giemsa (CAS No. 51811-82-6) was purchased from Biopur S.R.L. (Riccheri 195, Rosario, Argentina), and human pan-centromeric probe was purchased from Lexel S.R.L. (Luis Saenz Peña 1937, CABA, Argentina).

Cell Culture. The human cell line HEp-2 was obtained from the ABAC (*Asociación Banco Argentino de Células* from the Autonomous City of Buenos Aires, Argentina) and it was cultured in minimal essential medium (MEM) supplemented with 10% heat-inactivated fetal bovine serum (FBS), 100 units/ml penicillin, 100 mg/mL streptomycin, and 2.5 µg/mL amphotericin B. Cells were cultured at 37 °C in a humidified atmosphere of 5% CO₂ 95% air. Cell culture medium was renewed twice a week. After 7 days, cells became confluent and ready to use. Prior to use, iprodione was first dissolved in acetone and then diluted in culture medium. The final solvent concentration was less than 1% for all treatments. Negative controls (untreated cells and solvent vehicle-treated cells) were run simultaneously with herbicide-treated cultures. None of the treatments produced significant pH changes in the culture medium. For all experiments, confluent attached cells were removed from cell culture dishes with 0.25% sterile trypsin and diluted with MEM/10% FBS. For MTT assay, cells were reincubated into 96-well plates (0.2 mL; 2 × 10⁴ cells/well), and cytokinesis-block micronucleus (CBMN) assay was performed into 6-well plates (2 mL; 3.8 × 10⁵ cells/well). For the

other experiments, cells were replated into Petri dishes (8 mL; 7.5 × 10⁶ cell/plates).

Viability MTT Assay. To determine the range of concentrations of IP that keep cells metabolic activity, we exposed cell cultures at iprodione range from 3.75 to 300 µg/mL and negative control. We used the method of Mossman²⁵ with minor modifications. Briefly, cells were exposed for 24 h in serum-free medium to different dilutions of iprodione. Following incubation, treatment cell culture medium was removed; cells were washed with PBS and replaced with 1 mg/mL of sterilized MTT solution. This MTT solution was freshly prepared in MEM containing no FBS, since it has been shown that FBS can dose-dependently inhibit formazan crystals formation with a 50% decrease in these crystals when media with 5–10% FBS is used.²⁶ The plates with added MTT solution were then placed in the 5% CO₂ incubator for 90 min at 37 °C. MTT solution was removed and 200 µL of ethanol was added to each well to dissolve the blue formazan crystals. Optical density was measured at 570 nm with background subtraction at 655 nm, in a BIO-RAD Benchmark microplate reader (BIO-RAD Laboratories, Hercules, CA). Results were expressed as percentage of control (100% cell metabolic activity). Each assay involved 12 wells per condition and the values correspond to the average of 4 independent studies.

Cytotoxicity and Genotoxicity - Cytokinesis-Block Micronucleus (CBMN) Assay. According to the results obtained from MTT test, the sublethal concentrations (lower to CL₅₀/2) were select for the CBMN assay performed following the method of Fenech.²⁷ The cells were subcultured on glass coverslips in 6-well plates. Twenty-four hours after seeding, the medium was removed, and cells were treated with iprodione at 0, 3.5, 7, 17, 25, and 37 µg/mL in serum-free medium for 24 h. Then, cells were washed with PBS and incubated with medium containing cytochalasin B (final concentration 4 µg/mL) for 18 h. After these treatments, cells were washed with PBS and fixed with glacial acetic acid/methanol (1:3 v/v) for 10 min at room temperature. Then cells were washed twice with PBS, stained with Giemsa (%10 p/v), washed again with PBS, and after that the slides were mounted. Three independent experiments were carried out.

Fluorescence in Situ Hybridization (FISH). In order to discriminate between clastogenic and aneugenic effects, fluorescence in situ hybridization technique (FISH) using pan-centromeric probe was performed (Rhodamine, red). The FISH procedure is conducted using total centromeric probes following the method of Steinberg et al.²⁸ with minor modifications. Briefly, 0.5 µL of probe are dissolved in 3 µL of hybridization mixture. This hybridization mixture contains 30% formamide, 30% polyethylene glycol, 10% 20× SSC, 28% NaI, and 2% Tween. The resulting mixture is denatured at 70 °C for 7 min and then kept at –20 °C for a few min (no more than 10) before hybridization. Drops of hybridization mixture containing the probe were applied to glass slides, sealed with synthetic glue, and codenatured at 71 °C for 8 min. Hybridization is then conducted in a wet chamber at 37 °C overnight. Posthybridization washes consist in placing the preparations in a koplín jar with 0.4× SSC/0.3% Tween at 70 °C for 2 min and then place them in another koplín with 2× SSC/0.1% Tween at room temperature for 2 min. Slides are then counterstained with DAPI (Sigma). The DAPI solution was prepared by diluting 5 µL of a DAPI stock solution (20 µg/mL in ultrapure water) in 1 mL of antifade solution (50 mg p-phenylenediamine dihydrochloride, Sigma, in 5 mL of PBS). A 15 µL drop of this mixture is then placed over the preparation and covered with a coverslip. The slides are analyzed under a fluorescence microscope. MNi containing chromosomes with centromere are indicated as MNi cent +, while MNi containing chromosomes fragments without centromere are indicated as MNi cent –.

Scoring of Slides and Data Analysis. The slides were examined under a Leica DMLB light microscope (1000×). One thousand cells per slide from the negative control and treated groups were examined to calculate the binucleate cells (BN) frequency in interphase in order to estimate nuclear division index (NDI), MNi, NB, and NPB. The mitotic index (MI) and abnormal mitosis (AM) were quantified in mitosis.

$NDI = (M_1 + 2M_2 + 3M_3)/N$, where M_1 – M_3 represent the number of cells with 1–4 nuclei and N is the total number of cells scored.

MNiC+ = number of BN cells with MNi cent +/- totals BN cells

MNiC- = number of BN cells with MNi cent -/ total BN cells

NB = number of nuclear buds per 1000 interphase cells.

AM = number of abnormal mitosis/total mitosis

MI = (number of mitosis/1000 cells) \times 100

Oxidative Stress and Antioxidative Response. Antioxidant Enzyme Activities. For enzyme activities determination (Catalase, Superoxide dismutase, and Glutathione S-transferase), cells were grown at confluence with the different treatments for 24 h at 3 concentrations of iprodione (1.5, 7, and 25 $\mu\text{g}/\text{mL}$). Determinations were carried out in 11000g supernatants from cells lysates.

Catalase (CAT, EC 1.11.1.6) activity was determined by following hydrogen peroxide decomposition at 240 nm in a reaction mixture containing 50 mM potassium phosphate buffer (pH 7.0) and 3 M hydrogen peroxide.²⁹ Results were expressed as percentage of control (100% of activity).

Superoxide dismutase (SOD EC 1.15.1.1) activity was measured using a modify procedure in microplate of Beauchamp and Fridovich.³⁰ The standard assay mixture contained enzymatic sample, 0.1 mM EDTA, 13 mM DL-methionine, 75 μM nitro bluetetrazolium (NBT), and 2 μM riboflavin, in 50 mM phosphate buffer (pH 7.9), to a final volume of 0.3 mL. Samples were exposed to intense cool white light for 5 min. One SOD unit was defined as the enzyme amount necessary to inhibit 50% the reaction rate. Results were expressed as percentage of control.

Glutathione S-transferase (GST, EC1.11.1.9) activity was measured by Habig et al.³¹ technique. Briefly, standard assay mixture in 100 mM phosphate buffer (pH 6.5) contained enzymatic sample, 100 mM GSH solution, and 100 mM 1 chloro-2,4 dinitrobenzene (CDNB) in ethanol to a final volume of 0.8 mL. After adding CDNB, change in absorbance at 340 nm was followed for 180 s. One GST unit was defined as the amount of enzyme that catalyzes the formation of 1 μmol of GS-DNB per minute at 25 $^\circ\text{C}$; results were expressed as percentage of control.

GSH Equivalents Content. GSH levels were measured in HEp-2 cells following the Anderson³² procedure with some modifications. Briefly, after being incubated into Petri dishes at early confluence (80–90% confluent) at 3 concentrations of iprodione (1.5, 7, and 25 $\mu\text{g}/\text{mL}$), cells were collected, washed, and resuspended in PBS. Then, cells were lysed as described before. For the GSH determination, we proceeded as Coalova et al.³³ Results were expressed as percentage of control.

Oxidative Damage. Protein oxidation was quantified as carbonyl groups according to Reznick and Packer.³⁴ Carbonyl content was calculated from the peak absorbance (355–390 nm) using an extinction coefficient of 22000 $\text{M}^{-1} \text{cm}^{-1}$. Results were expressed as nmol carbonyl per mg protein. Results were expressed as percentage of control.

Statistical Analyses. The LC_{50} value to MTT was estimated by nonlinear regression sigmoidal dose–response method. For oxidative stress and cytotoxicity parameters, statistical analyses were performed using one-way analysis of variance (ANOVA) followed by Dunnet's test using significant levels of $p < 0.05$. Normality and homogeneity of variances were tested with Lilliefors and Barlett test; respectively, Graph Pad Prism 6 software was used for statistical analyses.

To study the factors that affect the abnormal genotoxicity biomarkers such as MNi and AM, generalized analysis linear mixed models (GLMMs) was performed. The explanatory variables used were treatment, NDI, and IM. For MNi and AM, binomial distribution was assumed. For binomial distribution, also samples were included as random factors to control overdispersion.³⁵ The models were ranked according to their Akaike Information Criteria (AIC), the lower AIC selected as the definite model. The R 3.4.3 software was applied³⁶ using the glmer function of the lme4 R-package³⁷ for binomial distribution. To analyze the difference in the treatment, Tukey test from the multitau package was performed to the genotoxic response comparing all the pair of samples.

RESULTS

The MTT test resulted in CL_{50} 85.86 (77.05–95.68) $\mu\text{g}/\text{mL}$ of iprodione (Figure 1). On the basis of this result, we

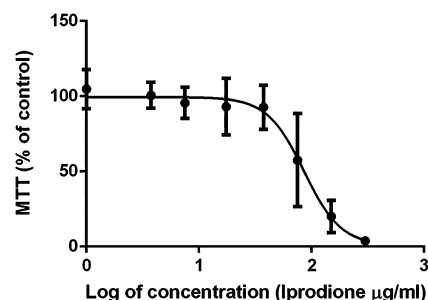


Figure 1. Dose–response curve for iprodione effects fitted by nonlinear regression. These effects were evaluated by the MTT test. Data are expressed as mean + SD ($n = 36$), relative to control cells (100% viability). MTT, 3-(4,5-dimethyl-hiazol-2-yl)-2,5-diphenyltetrazolium bromide; SD, standard deviation.

proceeded to expose the cells to the sublethal concentrations (below the CL_{50}) during 24 h to analyze the cytotoxicity as MI and NDI in order to determine the subcytotoxic concentrations of IP to evaluate genotoxicity. The concentrations chosen were 7, 17, and 25 $\mu\text{g}/\text{mL}$ of IP, because the MI and NDI values do not have significant differences regarding the negative control (Figure 2A,B). The results obtained in CBMN test showed MN induction which included both MNC+ as MNC-, with remarkable differences between them (Figure 3A–D). The GLMMs analysis performed to evaluate genotoxicity indicated that the frequency of MNi are associated with the treatments, the NDI, and MI factors. When the models were performed separately, the MNiC- was associated significantly with the treatments and MI factors, whereas MNiC+ was associated with the treatments, NDI, and MI factors. The most suitable model was based in the ranking according to their Akaike Information Criteria (AIC) where the best-reduced model was the model having the lower AIC.

The multiple comparisons of means to MNiC- throw results significant for 17 $\mu\text{g}/\text{mL}$ and 25 $\mu\text{g}/\text{mL}$ IP showed significant differences with respect to the negative control, whereas the values of MNiC+ were statistically different to the negative control in all the treatments (Table 1).

In mitosis, the abnormalities found were the bridges in anaphase and multipolar division (Figure 4A,B) in the more suitable models, only the treatment factor was associated with the AM. The Tukey contrasts showed significant differences in all concentrations of IP respect to the negative control and for 7–25 $\mu\text{g}/\text{mL}$ IP range (Table 2).

According to the result obtained to cytotoxicity and genotoxicity oxidative stress, studies were performed in 1.5, 7.0, and 25 $\mu\text{g}/\text{mL}$ of IP. The results showed that at 1.5 and 25 $\mu\text{g}/\text{mL}$ of IP the GSH intracellular content was lower with respect to control (Figure 5A). Furthermore, SOD activity is significantly lower in the cells exposed to 25 $\mu\text{g}/\text{mL}$ of IP than those in controls, while at that same concentration the cells have a significant increase in the levels of oxidative damage of the proteins, measured as the content of carbonyl groups. (Figure 5B,E). Neither CAT nor GST showed significant differences in the concentrations tested with respect to the control (Figure 5C,D).

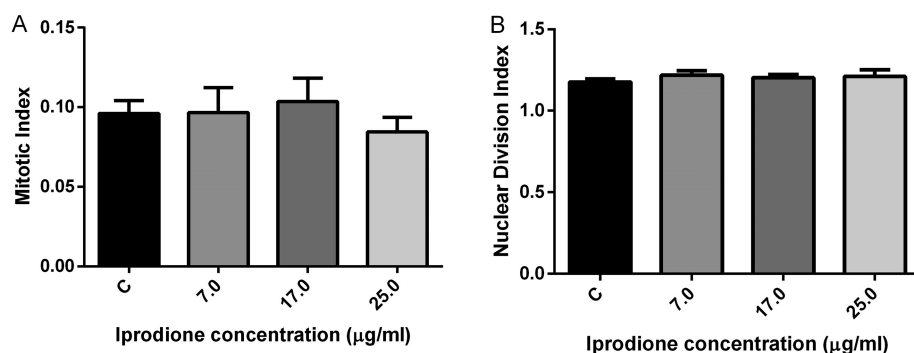


Figure 2. Cytotoxic/cytostatic effects of iprodione based on (A) nuclear division index, (B) nuclear replication index for HEp-2 cells. Results were expressed as mean \pm SD ($n = 9$).

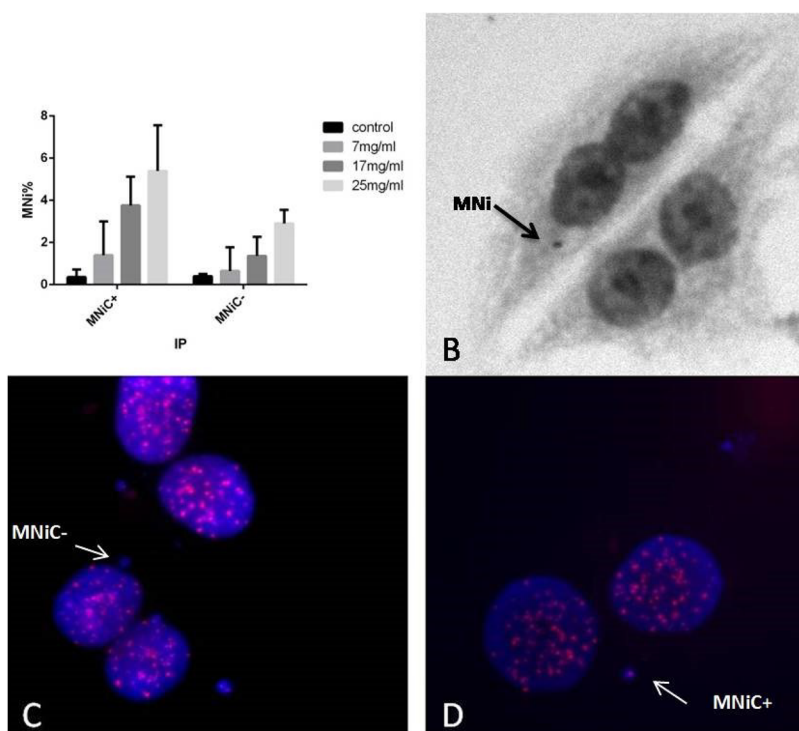


Figure 3. CBMN assay performed in HEp-2 cell line showing (A) MNi containing whole chromosomes are centromere positive (MNiC+) and detected by pan-centromeric probes, while MNi containing chromosome fragments are centromere negative (MNiC-). (B) Nuclear morphology of genotoxic effect of iprodione in HEp-2 cell line. Nuclear DNA was stained with giemsa dye and visualized by optic microscopy. Genotoxic effects were observed in interphase of cells as (C) binucleated cells with MNi cent-stained with DAPI and labeled with pan-centromeric probe respectively, and (D) binucleated cells with MNi cent+stained with DAPI and labeled with pan-centromeric probe, respectively. Nuclear DNA was stained with DAPI dye and visualized by fluorescence microscopy using filters for DAPI (blue) and for pan-centromeric probe (red).

Table 1. Multiple Comparisons of Means to MNiC- and MNiC+ (Tukey Contrasts)^a

treatment $\mu\text{g/mL}$	p values MNiC-	p values MniC+
7-0	0.12690	0.00598**
17-0	0.02695*	<0.001***
25-0	<0.001***	<0.001***

^aSignif. codes: ***, 0.001; **, 0.01; *, 0.05.

DISCUSSION

The biological systems respond to the exposure to xenobiotics by modifying their cellular and molecular status, and this can be detected by means of biomarkers. Biomarkers, such as MTT, MI, and NDI are indicators of the cell activity, thus the

values that are significantly lower than control indicates cytotoxic effects. The oxidative stress and genotoxicity tests are implemented to lower concentrations than the cytotoxic ones in which the metabolic or mitotic activity allows to reveal the effect we intend to characterize for the induction the respective biomarkers.³⁸ Therefore, the subcitotoxic concentrations chosen to performed genotoxicity analysis were in a range containing 7, 17, and 25 $\mu\text{g/mL}$ IP. The more suitable GLMMs used to explain the genotoxicity biomarkers induction showed the association of MNiC+ and MNC- values with all the IP concentrations and MI. The MNiC+ was also associated with the NDI. Considering that MNi are formed by chromosomes or chromatin fragments produced in previous mitosis and they are quantified in the binucleated cells, the MI and NDI variations can influence the MNi induction since they

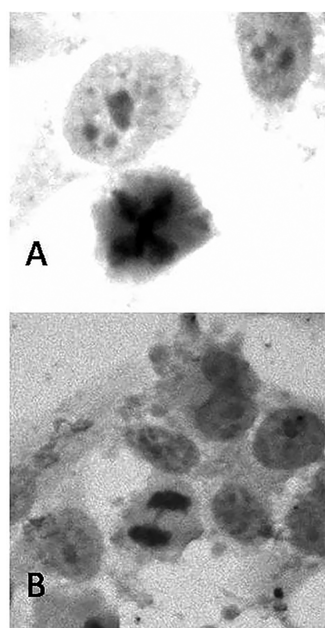


Figure 4. Abnormal mitosis. (A) Multipolarmitosis and (B) nucleoplasmic bridges.

Table 2. Multiple Comparisons of Means to AM (Tukey Contrasts)^a

treatment $\mu\text{g/mL}$	<i>p</i> values AM
7–0	<0.001***
17–0	<0.001***
25–0	<0.001***

^aSignif. codes: ***, 0.001; **, 0.01; *, 0.05.

are a measure of proliferation cells. The MNiC+ were more induced than the MNiC– for all the concentrations (Table 1).

The use of a centromeric probe can fail if the chromosome attaches to the spindle but does not fail if these chromosomes are whole or broken. In order to know if the chromosome is whole, a combination of telomeric and centromeric probes could be used. In this work, we assume this limitation is due to both effects, clastogenic and aneugenic, allowing the general characterization of the damage, whereas the analysis is of the mode of action (MOA) or prediction studies in which the threshold values are required and this is not the goal.

According to more suitable GLMMs, the AM induction is associated with the treatments only and the values of IM and NDI did not show association with AM as it is expected for noncytotoxic concentrations. The genotoxicity evidenced as AM is attributed to the three concentrations of IP in the range studied, according to the Tukey contrasts analysis. The IP exposition induces disturbances on mitotic machinery evidenced as MNiC+ and multipolar division, whereas the clastogenic effect is evidenced as MNiC– and anaphasic bridges. The formation of multipolar spindle is compatible with the microtubule interaction or microtubule associated proteins,³⁹ while the bridge's induction arises from the rupture of the telomeric region which generates sticky ends.⁴⁰ Both effects could be related due to genomic instability produced for interaction with several targets, involving mechanism of indirect genotoxicity.⁴¹ Indirect genotoxicity includes mechanism of oxidative damage, affecting proteins and other biomolecules including DNA. According to this hypothesis,

the parameters related to oxidative stress and damage indicate that the exposure of HEp-2 cells to IP produces an effect in the redox balance and damages the proteins as consequence of exposition to IP. This imbalance is evidenced by the changes in the GSH intracellular content, the low SOD activity and the increase in the levels of oxidative damage of proteins, at the maximum tested concentration of IP. Such effects could be related to the identified genotoxic alterations for the same range of concentrations. In addition, 1.5 $\mu\text{g/mL}$ IP concentration was included in the oxidative stress studies to analyze the behavior of this mechanism at low doses, where the response (GSH consumption together with the GST and SOD activity) is compatible with the oxidative protection. Exposure to higher concentrations would lead to the fall of SOD activity and protein oxidation according to the result obtained. The quantities and quality of the biological target and the interactions could explain the different threshold observed to clastogenic and aneugenic genotoxic effects.⁴²

The few research studies that analyze the IP genotoxicity in the literature present controversial results. Some studies carried out using *Crepis capillaries* show that the IP induces lagging chromosomes and this effect is compatible with the abnormal formation of mitotic spindles.⁴³ By contrast, the MNi test in *Tradescantia* pollen tetrads showed negative results.⁴⁴ The literature about IP genotoxicity in human cells is limited to the study of the biomarkers induced by occupational or environmental exposure to pesticide mixtures which include the IP.^{45–48}

Among the study models conducted to know about the different toxic effects of xenobiotics, the guidelines of the Organization for Economic Coordination and Development, OECD 473,⁴⁹ recommend the human cell lines for the chromosome aberrations' analysis as a resource that allows one to study such responses at a cellular level and identify mechanistic aspects related to the molecular interaction.^{50,51} Thus, the results obtained using human cells *in vitro* are complemented with studies performed *in vivo* in mammals, to obtain conclusions about the risk of human exposure. The HEp-2 is a model used in several studies to evaluate the cytotoxicity and genotoxicity of xenobiotics, in which it is proved that such test system is feasible and sensitive for that purpose.^{52–56,56} The final acceptance of the MNi *in vitro* test for the (OECD) guideline No. 487⁵⁷ places the test for the determination of the genotoxicity potential in the exploratory research, as well as in the regulatory environment. In this context, our results provide valuable and relevant information with data about IP that can be considered for the toxicological risk evaluation due to human exposure and search of genotoxicity threshold values.⁵⁸ The present work carried out in a human *in vitro* model is the beginning of a series of studies necessary for the characterization of genotoxic agents prior to *in vivo* tests in experimental animals which should be avoided as long as possible in according to regulatory genotoxicology. More studies should be focused in finding the threshold to genotoxicity *in vivo* of IP in order to determine the concentration of residues that is compatible with food safety.

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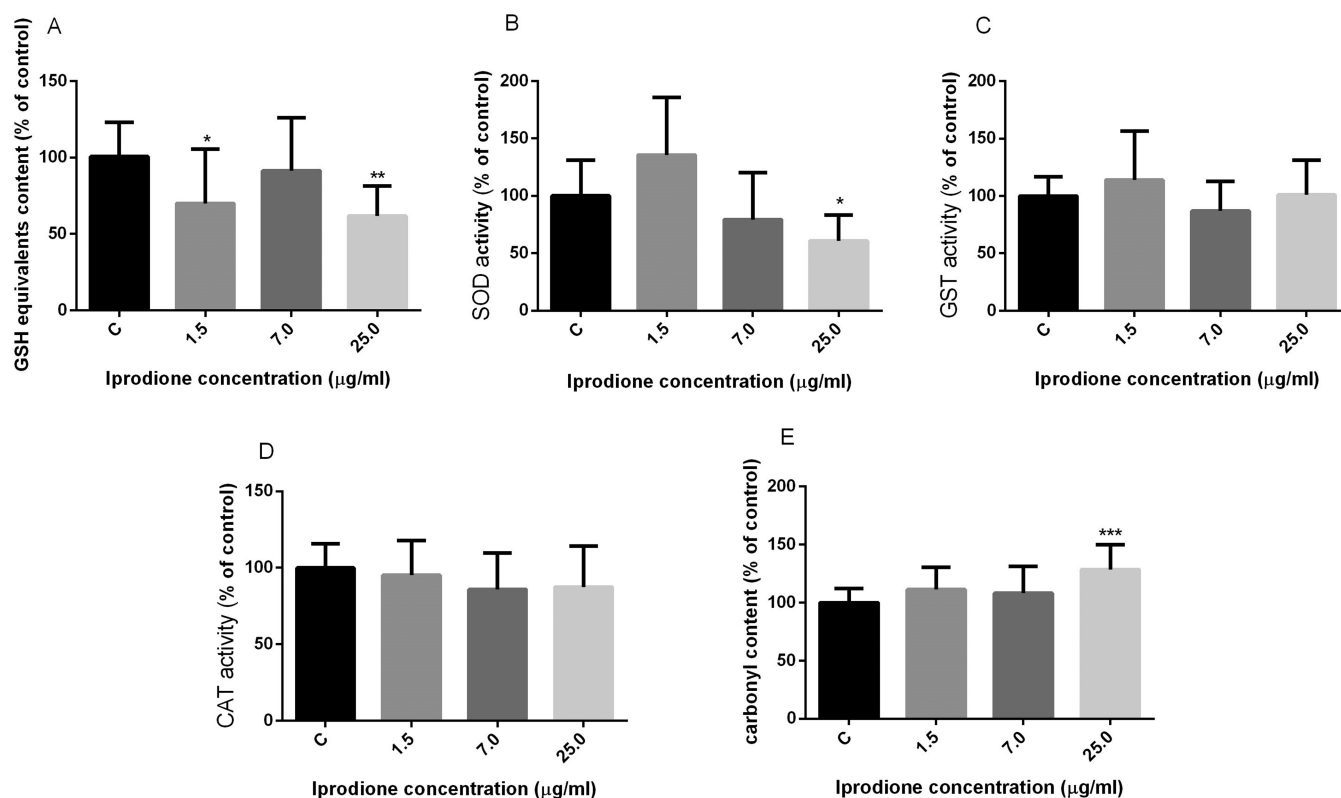


Figure 5. Enzymatic and non enzymatic antioxidant, detoxification capacity, and levels of oxidative damage of iprodione in HEp-2 cells. (A) GSH intracellular levels, (B) superoxide dismutase activity (U SOD mg^{-1} proteins), (C) glutathione S-transferase activity (GST mg^{-1} proteins), (D) catalase activity (CAT mg^{-1} proteins), and (E) carbonyl content (nmol mg^{-1} proteins) in HEp-2 cells at 1.5, 7.0, and 25.0 $\mu\text{g}/\text{mL}$ of iprodione. Results are expressed as mean \pm SD of four independent experiments ($n = 16$). Significant differences between control and treatments are indicated by asterisks: * indicate $p < 0.05$.

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Notes

The authors declare no competing financial interest.

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