Flow cytometry evaluation of lead and cadmium effects on mouse spermatogenesis

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Abstract

Flow cytometry (FCM) is a powerful tool to evaluate cell DNA content and ploidy levels. We have assessed the accuracy of two protocols of nuclei isolation from paraffinized samples (P1 and P2) by comparing FCM results with those obtained using fresh material (F1–F3). After isolation, nuclei were stained with propidium iodide and quantitatively analysed by FCM for changes in germ cell ratios. Results obtained with Protocol P2 were similar to those obtained using the protocol that gave best results for fresh tissues (F2). Protocol P2 was then applied to paraffin embedded testicular samples from ICR-CD1 mice exposed to 1, 2 and 3 mg CdCl\textsubscript{2}/kg bw by single subcutaneous injection, and to 74 and 100 mg PbCl\textsubscript{2}/kg bw administered in four repeated doses. The highest doses of CdCl\textsubscript{2} decreased the number of haploid (1C) cells and increased the number of diploid (2C), S phase and tetraploid (4C) cells. Treatment with PbCl\textsubscript{2} did not induce significant changes in testicular cells subpopulations. These results support the usefulness of FCM in evaluating the effect of toxic substances on mouse spermatogenesis, using both fresh and paraffinized material.

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1. Introduction

Flow cytometry (FCM) is a fast and sensitive tool that can provide quantitative analyses of different cell types and an insight into the cell cycle status of those cells. FCM also allows the measurement of DNA content of cell subpopulations in a mixed cell population, using fluorescence intensity distribution histograms, enabling the identification of cells with different ploidy levels within a given tissue [1,2].

FCM has been applied to the study of the toxic effect of substances on spermatogenesis of several species, namely mouse [1,3,4], rat [5,6] and golden hamster [7]. Alterations in testicular germ cell percentages are common responses to chemicals such as vinblastine [3], vindesine sulphate [8], diepoxybutane [1] or methoxyacetic acid [9]. In all cases FCM analyses were performed in nuclei isolated from fresh material. The specific methodologies are quite variable, and include 0.1% pepsin digestion [1], tripsin and DNAse digestion [5] and testis dissection in TNE (0.01 M Tris buffer, 0.15 M NaCl, 0.01 M ethylenediaminetetraacetic acid, pH 7.4) buffer [7]. Other authors prepare nuclei for FCM analysis using citric acid with detergent solutions [10]. Nevertheless, the general requirement for fresh material may condition a broader utilization of FCM in these kinds of studies.

In most laboratories interested in toxicology, histological evaluation of tissues and organs is performed routinely, and usually includes paraffin embedding. In the present work, the possibility of using FCM to analyse mouse testicular cells from embedded samples was assessed. This technique has been developed by Hedley et al. [11] to study DNA content in tumoral samples, and since that time it has been routinely used at the clinical oncology level. Therefore, the development of
a FCM protocol for testicular cell analysis, especially in cases where spermatogenesis is affected, may be a valuable tool in the reproductive toxicology field.

Cadmium is a well-known reproductive toxicant that causes disruption of the blood–testis barrier [12]. Several works describing the histological and ultrastructural effects of cadmium in the testis have been published (e.g. [13,14]). However, the quantification of cadmium-induced alterations in spermatogenesis, in terms of the relative percentages of each testicular cell type, has never been performed. Also, the toxicological effects of lead on male reproductive system have long been studied. The main effects referred in the literature are a decrease in sperm cell number and motility [15,16], and an increase in sperm morphological abnormalities [15]. Although histopathological effects of lead compounds on the testis have been previously described (e.g. [16,17]), those studies have not used more refined techniques such as FCM.

The aim of the present work is thus to demonstrate that, in alternative to fresh material, paraffin embedded samples of testicular material might be used to accurately determine the ploidy level of germ cells using FCM. The validation of paraffinized samples for FCM nuclei analyses was done by comparing the results with data obtained using fresh material. The validated protocol was then used to assess the toxic effect of cadmium and lead chloride on mouse spermatogenesis, at the level of each specific cell-stage.

2. Materials and methods

2.1. Animal housing

Seven weeks old male ICR-CD1 mice were provided by Harlan Interfauna Ibérica SA, Barcelona, Spain. The animals were housed in a constant temperature (22 ± 2 °C) and relative humidity (40–60%) vivarium on a 12-h light/12-h dark cycle. Water and food were provided ad libitum. Mice were allowed to acclimate for one week before experimental use.

Animal experiments were conducted in accordance with institutional guidelines for ethics in animal experimentation (Rule number 86/609/CEE-24/11/92).

2.2. Flow cytometric analysis of fresh and paraffinized samples

Different protocols for the isolation of mouse testicular cells were used from both fresh and paraffin embedded tissues. For each protocol at least four testes from different mice were used.

2.2.1. Fresh samples

Three different protocols for nuclei isolation from fresh samples were tested: Protocol F1: Samples were prepared according to a protocol developed for the golden hamster [7]. Briefly, testes were dissected in TNE buffer (0.01 M Tris buffer, 0.15 M NaCl, 0.01 M ethylenediaminetetraacetic acid, pH 7.4), filtered with a nylon mesh of 55 μm and the cell suspensions that were obtained were supplemented with glycrol 10% (v/v) and stored at −20 °C until analysis. Thereafter, cell suspensions were thawed and centrifuged at 500 × g for 5 min and resuspended in TNE buffer.

Protocol F2: This protocol was based on the protocol described by Spanò et al. [1] for mouse testicular cells. Briefly, testes were dissected in 0.1% pepsin–HCl at pH 1.8 for 10 min. After filtration, the cell suspension was fixed in 96% ethanol and kept at −20 °C. After that, cell suspensions were incubated with pepsin–HCl 0.5% for 10 min, centrifuged at 500 × g for 10 min and resuspended in phosphate buffered saline (PBS).

Protocol F3: In this case, testes were dissected in 0.1 M citric acid and 0.5% (v/v) Tween 20 and incubated for 10 min at RT. After filtration in a 55 μm nylon mesh, the suspension was centrifuged at 300 × g for 10 min, the supernatant was then discarded and the pellet was resuspended in a small volume of PBS. The suspension was fixed in 70% ethanol and was kept for a few days. After that, the cell suspension was centrifuged and cells were resuspended in detergent solution, incubated for 10 min at RT and finally 5 volumes of a 0.4 M Na2HPO4 solution were added.

2.2.2. Paraffin embedded material

Two protocols for isolation of nuclei from paraffin embedded samples were assayed. In both protocols two or more sections (40 μm thick) of five blocks corresponding to five different mice were used. Sections were deparaffinized in xylol and rehydrated.

In Protocol P1, samples were incubated in a water bath at 80 °C with citric acid 2 mg mL−1 at pH 6.0 for 2 h, digested with pepsin, pH 1.5, at 37 °C and washed in PBS [11,18]. Protocol P2 was similar to Protocol P1 except for the incubation with citric acid, which was not performed [19].

2.2.3. Flow cytometry analysis

In all protocols (F1–F3, P1 and P2), samples were treated with 50 μg mL−1 of RNase (Sigma, St. Louis, MO, USA) in order to eliminate RNA, and stained with 50 μg mL−1 of propidium iodide (PI). Samples were analysed within a 15-min period. The relative fluorescence intensity of PI-stained nuclei was measured with a Coulter EPICS XL (Coulter Electronics, Hialeah, FL, USA) flow cytometer. The instrument was equipped with an air-cooled argon-ion laser tuned at 15 mW and operating at 488 nm. Integral fluorescence together with fluorescence pulse height and width emitted from nuclei was collected through a 645 dichroic long-pass filter and a 620 band-pass filter and converted on 1024 ADC channels. Prior to analysis, the instrument was checked for linearity with fluorescent check beads (Coulter Electronics) and the amplification was adjusted so that the peak corresponding to the haploid peak was positioned at channel 200. This setting was kept constant. The results were obtained in the form of three graphics: linear fluorescence light intensity (FL), forward angle light scatter (FS) versus side angle light scatter (SS) and FL pulse integral versus FL pulse height. This last cytogram was used to eliminate partial nuclei and other debris, nuclei with associated cytoplasm and doublets (these events have a higher pulse area but the same pulse height as single nuclei).

2.3. Heavy metal treatment assays

For toxicological studies, ICR-CD1 mice were housed in the conditions described in Section 2.1. For cadmium toxicological assays three groups of five mice each were subcutaneously injected with 1, 2 or 3 mg CdCl2/kg bw. For lead toxicological assays two groups of five mice each were subcutaneously injected with 74 and 100 mg PbCl2/kg bw for four consecutive days. In both experiments control groups of five mice each were injected with the saline vehicle (0.9% NaCl) for equivalent periods.

Twenty-four hours after the last injection, animals were sacrificed and both testes were removed, the right testis was fixed in neutral 10% buffered formalin and the left testis was fixed in Bouin’s solution. Both samples were then dehydrated and embedded in paraffin wax. For the histopathology analysis sections were performed in a microtome and stained with haematoxylin and eosin. For FCM analysis samples were treated as described above for Protocol P2.

2.4. Statistical analysis

Statistical analyses were performed using a one-way analysis of variance (ANOVA) (SigmaStat for Windows Version 3.1, SPSS Inc., USA) to compare the values of germ cell percentages. A multiple comparison Tukey test was applied to assay the differences between control and metal treated groups. The correlations between the results obtained with the different treatments were performed by the Pearson Correlation test [20]. In all cases the level of statistical significance was set at p ≤ 0.05.
3. Results

3.1. Flow cytometric analysis of fresh and paraffinized samples

Forward angle light scatter is proportional to cell-surface area or size and side angle light scatter is proportional to cell granularity or internal complexity. The different methods of nuclei isolation gave different results in terms of nuclei light scatter properties. Nuclei isolated from fresh samples (Protocols F2 and F3) were homogeneously distributed in one single population (Fig. 1A and D). Contrarily, the nuclei from embedded origin showed discrete populations in terms of relative size and complexity (Fig. 1G and J). The best separation of nuclei in terms of light scatter was observed in nuclei isolated using Protocol P2 (Fig. 1J).

Linear fluorescence light intensity represents the linear fluorescence of the DNA stained with PI (Fig. 1B, E, H and L). Protocol F1 produced unsatisfactory definition of peaks and for this reason the results are not presented. DNA histograms from the other two fresh and both embedded samples revealed four main peaks corresponding to different ploidy levels. The sub-haploid peak (HC) consists of elongated spermatids and the haploid peak (1C) refers to round spermatids. The discrimination of the haploid nuclei in two peaks is due to different staining of elongated and round spermatids, since the former present a highly condensed chromatin and PI intercalation with DNA is lower. Chromatin condensation occurs as a result of the sperm maturation process where exchange of histones with protamines reduces PI intercalation [21]. Somatic cells, spermatogonias and secondary spermatocytes are recorded in the diploid peak (2C). Cells in the G2/M phase of the cell cycle and primary spermatocytes are included in the tetraploid peak (4C). The region between the diploid and the tetraploid peaks corresponds to cells that are actively synthesising DNA (S phase).

The mean CV obtained for all protocols are shown in Fig. 1B, E, H and L. The higher values were found in the HC subpopulation. This higher CV value can be explained by the chromatin status of the HC nuclei (elongated spermatids), that present different degrees of chromatin condensation, due to different degrees of PI intercalation between the bases.

The cytograms with pulse integral versus pulse height were computed to define a gating that excludes doublets. Nuclei from fresh samples (Fig. 1C and F) showed more doublets than paraffin samples (Fig. 1I and M), with the lower number of doublets being observed in Protocol P2.

Protocols F2 and F3 gave different results ($p < 0.001$) with respect to the subpopulation ratios. Both Protocols F2 and F3 presented similar discrete peaks and CV’s. However, some differences were found in the percentage of each subpopulation obtained by each protocol: with the Protocol F2 (Figs. 1B and 2), the HC subpopulation represented 34% of the nuclei population, whereas it represented only 17% of the nuclei isolated with the Protocol F3 (Figs. 1E and 2). In paraffin embedded tissues, the ploidy percentages were similar to the values obtained using Protocol F2 (Fig. 2). The results obtained with Protocol F2 were positively correlated with those obtained by both Protocol P1 and P2 (Pearson correlation, $r > 0.96$, $p = 0.007$ for Protocol P1 and $p = 0.009$ for Protocol P2). Paraffin protocols were also positively correlated with each other (Pearson correlation, $r = 0.992$; $p = 0.0008$). For toxicological assays we therefore chose Protocol P2 for FCM analyses.

3.2. Heavy metal treatment assays

The highest doses of cadmium chloride (2 and 3 mg/kg bw) induced a significant increase in absolute and relative testis weight after 24 h, but both body weight and absolute and relative weights of the epididymis were not affected by cadmium chloride treatment (Table 1). In the lead chloride assay neither testis nor epididymis weights were significantly affected (Table 2).

Cadmium chloride at the concentration of 1 mg/kg bw did not induce any histological alteration to the seminiferous epithelium (Fig. 3B). The higher concentrations of cadmium chloride (2 and 3 mg/kg bw) provoked severe alterations in the seminiferous tubules, namely haemorrhage, interstitial edema and disruption of the germinal epithelium with desquamation of cells into the lumen (Fig. 3C and D), particularly at the highest concentrations of compound. Lead chloride did not induce any typical changes on testis histology, in terms of percentage of tubules with germ cell degeneration or depletion, and tubular diameter when compared with the respective controls ($p \leq 0.05$) (data not shown).

FCM analyses showed that the relative percentage of sub-haploid nuclei (elongated spermatids) was significantly reduced ($p \leq 0.05$) following administration of 2 and 3 mg CdCl$_2$/kg bw (Fig. 4). Administration of 3 mg CdCl$_2$/kg bw decreased the percentage of haploid cells (round spermatids) ($p \leq 0.005$).

### Table 1

<table>
<thead>
<tr>
<th></th>
<th>Control ($n = 9$)</th>
<th>1 mg/kg bw ($n = 10$)</th>
<th>2 mg/kg bw ($n = 10$)</th>
<th>3 mg/kg bw ($n = 10$)</th>
</tr>
</thead>
<tbody>
<tr>
<td>Body weight (g)</td>
<td>33.23 ± 2.536</td>
<td>33.50 ± 2.257</td>
<td>33.31 ± 2.105</td>
<td>34.40 ± 1.808</td>
</tr>
<tr>
<td>Testis weight (g)</td>
<td>0.114 ± 0.017</td>
<td>0.106 ± 0.011</td>
<td>0.144 ± 0.033*</td>
<td>0.156 ± 0.026*</td>
</tr>
<tr>
<td>Epididymis weight (g)</td>
<td>0.046 ± 0.007</td>
<td>0.041 ± 0.006</td>
<td>0.045 ± 0.006</td>
<td>0.046 ± 0.007</td>
</tr>
<tr>
<td>Testis/body weight ratio (%)</td>
<td>0.345 ± 0.040</td>
<td>0.316 ± 0.022</td>
<td>0.435 ± 0.101*</td>
<td>0.452 ± 0.058*</td>
</tr>
<tr>
<td>Epididymis/body weight ratio (%)</td>
<td>0.141 ± 0.018</td>
<td>0.122 ± 0.017</td>
<td>0.135 ± 0.020</td>
<td>0.132 ± 0.017</td>
</tr>
</tbody>
</table>

Values represent mean ± S.D. The symbol * means significant difference between the control and 1, 2 and 3 mg CdCl$_2$ treated samples at $p \leq 0.05$. 
Fig. 1. Flow cytometry graphics of nuclei isolated from fresh tissue by the Protocol F2 (A–C) and Protocol F3 (D–F) and paraffin embedded tissue by Protocol P1 (G–I) and Protocol P2 (J, L and M). The left column cytograms show the forward angle light scatter (FS) vs. side angle light scatter (SS), the middle column histograms show the relative fluorescence light intensity (FL) from propidium iodide and the right column cytograms show the relative FL pulse integral vs. relative FL pulse height.
Fig. 2. Ploidy percentage variation of germ cells isolated by the Protocol F2, Protocol F3 (fresh tissue), Protocol P1 and Protocol P2 (paraffin embedded tissue). Percentages were estimated as number of nuclei in each ploidy level/total number of nuclei × 100 and represent mean ± S.D. for at least four testicular samples. The symbol ‘*’ indicates significant difference between the fresh and paraffin methods at p ≤ 0.05.

Concomitantly, the percentage of diploid cells increased following treatment with 2 and 3 mg CdCl₂/kg bw. The same was observed for cells in S phase by treatment, in this case for all cadmium doses. In what concerns tetraploid cells a significant increase (p ≤ 0.05) was only found after administration of 3 mg CdCl₂/kg bw.

The lead chloride treatment induced some variations in the percentage of germ cells, although the only one that was statistically significant (p ≤ 0.05) was an increase in the percentage of cells in S phase with 100 mg PbCl₂/kg bw (Fig. 5).

### Table 2

<table>
<thead>
<tr>
<th></th>
<th>Control (n = 10)</th>
<th>74 mg/kg bw (n = 10)</th>
<th>100 mg/kg bw (n = 10)</th>
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</thead>
<tbody>
<tr>
<td>Body weight (g)</td>
<td>36.294 ± 2.988</td>
<td>36.444 ± 2.330</td>
<td>36.758 ± 3.518</td>
</tr>
<tr>
<td>Testis weight (g)</td>
<td>0.123 ± 0.018</td>
<td>0.124 ± 0.016</td>
<td>0.128 ± 0.025</td>
</tr>
<tr>
<td>Epididymis weight (g)</td>
<td>0.047 ± 0.005</td>
<td>0.049 ± 0.005</td>
<td>0.048 ± 0.006</td>
</tr>
<tr>
<td>Testis/body weight ratio (%)</td>
<td>0.340 ± 0.040</td>
<td>0.341 ± 0.048</td>
<td>0.353 ± 0.088</td>
</tr>
<tr>
<td>Epididymis/body weight ratio (%)</td>
<td>0.130 ± 0.009</td>
<td>0.135 ± 0.016</td>
<td>0.130 ± 0.019</td>
</tr>
</tbody>
</table>

Values represent mean ± S.D. No statistical differences were found at p ≤ 0.05.
The development of protocols for isolation of nuclei from non-fresh samples may potentiate the use of FCM in reproductive toxicology studies. As Bouin's fixation is routinely used for the study of spermatogenesis [22] its efficacy was evaluated in nuclei fixation for FCM studies. However, this fixative did not produce satisfactory results in terms of DNA content analysis, just as described previously by Hedley [18]. Therefore, Bouin's solution was used for histology analysis and buffered formalin for DNA studies by FCM.

The different methods used to isolate mice testicular nuclei from fresh samples gave clearly different results. Although it has been used successfully in the golden hamster [7], Protocol F1 did not provide a clear definition of peaks, possibly due to species-specific differences in the isolation of nuclei. Protocol F3 has been used for DNA analysis of animal cells [10] and provided a good peak definition (the CV values obtained were among the best), as well as low level of debris. However, this method induced the formation of a great number of doublets and the elongated spermatid population was underestimated, probably due to the high ionic strength of the buffers used. Finally, Protocol F2 provided a good peak definition, and the percentages of cells in the different ploidy levels are consistent with the results previously obtained by other authors [1,3]. On the other hand, although Protocol F3 gave discrete peaks, the CV's obtained showed no great improvement when compared with Protocol F2. As the former is more time consuming and aggressive than the latter, we propose that Protocol F2 is more advisable for this type of studies.

The two protocols of nuclei isolation from paraffin samples gave similar results. However, Protocol P2 was more reproducible in terms of the number of nuclei per minute (data not shown), probably due to the fewer steps involved. Protocol P2 was therefore selected to evaluate the effect of cadmium chloride and lead chloride on mouse testis.

Cadmium chloride induced an increase in absolute and relative testes weight 24 h after administration. This effect was also observed by Shen and Sangiah [23], and it was hypothesized that it could be due to damage of the capillary endothelium and subsequent increase in permeability that lead to testicular edema. This effect on testicular weight was reversed as the time after exposure increased (data not shown). Histological evaluation showed that cadmium chloride seriously injured the testis. These short term effects of cadmium were also observed by FCM, in terms of significant alterations in germ cell percentages. These changes may be due to the effects of CdCl2 on both the blood–testis barrier and vascular endothelium. Cadmium disrupts tight junctions between Sertoli cells and alters Sertoli–germ cell adhesion [12] with consequent exfoliation of spermatids within the seminiferous tubules. This effect is supported by the presence of immature cells in the epididymis lumen (data not shown) and could therefore explain the decrease in the percentage of spermatids observed by FCM. Alternatively, non-specific germ cell degeneration and death can occur due to anoxia caused by changes in interstitial vasculature. No significant changes in mice testis/body weight or epididymis/body weight ratios were observed following treatment with lead chloride, similarly to what was described by Wadi and Ahmad [15]. Lead chloride did not induce striking changes in testis histology. This result was reflected in the absence of alterations in the percentages of testicular germ cells detected by FCM, with the exception of an increase in the percentage of cells in S phase. We hypothesize that this increase may be related with the mitogenic activity of lead previously described in lymphocytes [24] and liver cells [25]. Pinon-Lataillade et al. [26] also did not find prominent alterations in the reproductive system of male rats after exposure to lead oxide or lead acetate. These results suggest that the blood–testis barrier protects the seminiferous epithelium from the toxic effects of lead. The absence of damage to the blood-testis barrier by lead has also been pointed out by other authors [15,27].

In this work we demonstrated that buffered formalin fixed and paraffin embedded material allows good DNA preservation and enables cell extraction and quantification by FCM analysis in toxicological assays using mouse testicular tissue. The great advantage of this technique is that it allows long-term storage of samples and the possibility of making several replicas, thus increasing reliability. Paraffin embedded DNA analysis of testicular cells by FCM should be considered as an useful approach for the screening of substances in terms of testicular toxicity, as well as an important complement to histopathology.

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