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A cytogenetic study of hospital workers occupationally exposed to radionuclides in Serbia: premature centromere division as novel biomarker of exposure?

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Abstract

Purpose The health risk of chronic exposure to radionuclides includes changes in the genome (e.g., chromosomal aberrations and micronuclei) that increase chromosomal instability. There are also other phenomena, which seem to appear more frequently in metaphases of exposed persons (such as premature centromere division). The aim of this study was to discover whether or not there is correlation between incidence of named cytogenetic changes in persons occupationally exposed to radionuclides in comparison with unexposed control group, and if significant correlation is determined, can premature centromere division be consider as a biomarker of radiation exposure?

Methods The exposed group comprised 50 individuals occupationally exposed to radionuclides. The reference control group consisted of 40 unexposed individuals. Chromosomal aberrations, micronuclei and premature centromere division were analyzed according to a standard International Atomic Energy Agency protocol. Statistical analyses were performed using SPSS 17.0 statistics.

Results The means for analyzed cytogenetic changes were significantly higher in the exposed group. Positive correlation between them was found in exposed group. Premature centromere division parameter PCD5-10 was

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selected as particularly suitable for separating groups (exposed/unexposed).

Conclusions Identification of other phenomena related to radionuclide exposure, beside well known, may clarify recent problems in radiobiology concerning the biological response to low doses of ionizing radiation and its consequences.

Keywords Chromosomal aberrations · Micronuclei · Premature centromere division · Radionuclides

Introduction

The health risk of chronic occupational exposure to low doses of ionizing radiation (IR) includes changes in the genome that increase chromosomal instability as one of the main forces driving the onset and progression of carcinogenesis (Hoeijmakers 2001; Miller et al. 2008; Morgan 2003; Mothersill and Seymour 2004; Ullrich and Ponnaiya 1998). Genome instability induced by low doses of radiation manifests as changes, such as aneuploidy, micronuclei, chromosomal aberrations, oxidative stress and increased apoptosis incidence (Miller et al. 2008; Engin et al. 2005). Several cytogenetic studies performed on peripheral blood lymphocytes (PBL) from hospital workers exposed to low doses of IR during the course of their employment revealed an enhanced baseline of chromosomal aberrations (CAs) as well as micronuclei (MNs) compared to control populations (Ballardin et al. 2007; Barquinero et al. 1993; Bigatti et al. 1988; Bonassi et al. 1997; Bouraoui et al. 2013; Jovičić et al. 2009; Maffei et al. 2002; Ropolo et al. 2012; Saberi et al. 2013; Sari-Minodier et al. 2007; Thierens et al. 2000; Touil et al. 2002).

Besides them, there are other phenomena visible on cytogenetic level, which seem to appear more frequently in metaphases of persons exposed to IR than in unexposed groups. One of them, certainly, is premature centromere division (PCD) (Jovičić et al. 2010), discovered in peripheral lymphocytes, characterized by chromatids already separated in metaphase of the cell cycle. Centromere dysfunction presented by premature centromere splitting is caused by disorder in spatial and temporal regulation of mitosis, which finally leads to aneuploidy and genomic instability (Litmanovic et al. 1998; Vig and Wodnicki 1974). This phenomenon was predominantly studied as consequence of exposure to genotoxic chemicals (Major et al. 1999).

However, the prevalence of PCD in persons occupationally exposed to ionizing radiation has not been studied thoroughly. Among exposed workers, those who perform nuclear medicine and interventional procedures may receive higher doses and are the subjects of concern (Sari-Minodier et al. 2007).

Regarding all the aforementioned, the aim of this study was to examine the relationship between PCD prevalence in exposed and control group and then to discover whether or not there is correlation between the prevalence of CAs, MN with the prevalence of PCDs in persons occupationally exposed to IR, in nuclear medicine department.

If there is significant correlation between those three phenomena in the exposed subjects, can we then consider PCD as a biomarker of radiation exposure and what are the cutoff values of PCD parameters that are sensitive and specific enough to classify an examined subject to the exposed or unexposed group?

Materials and methods

Subjects

The exposed group comprised 50 individuals (29 females and 21 males; average age 45.24 years, age range 27-59 years) occupationally exposed (daily continual exposure ranged from 5 to 35 years) to radionuclides $(Y^{90}$ and I¹³¹). The reference group consisted of 40 individuals (20 females and 20 males; average age 44.40 years, age range, 34-60 years) who had never been occupationally exposed to IR or chemical mutagens in their workplaces. Among exposed persons, 26 subjects (12 females and 14 males) were regular smokers-smoking between 15 and 20 cigarettes per day, up to 20 years consecutively (16.14 years average)-and 24 were not (17 female and seven male subjects). In the control group, 17 subjects (eight females and nine males) were regular smokers-smoking between 15 and 20 cigarettes per day, up to 20 years consecutively (14.28 years average)—and 23 were not (12 female and 11 male subjects). No subject reported medicinal treatment (including exposure to diagnostic X-rays) over 6 months before blood sampling. The characteristics of the study population regarding gender (male or female), age, working experience (WE), occupational exposure time (DOE duration of occupational exposure in years), smoking habit, use of protective measurements and effective doses (external exposure) for the preceding and last 5 years in mSv (ED₁ and ED₅, respectively) are listed in Table 1.

Table 1	General	characteristics	of the	studied	population
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Groups	Ν	Mean	SD	Min–Max
Gender				
Exposed	50 (29 f; 21 m)			
Control	40 (20 f; 20 m)			
Age (years)				
Exposed	50	45.24	8.37	27-59
Control	40	44.40	6.21	34–60
Smoking (nu	mber of smokers)			
Exposed	26 (12 fs; 14 ms)			
Control	17 (8 fs; 9 ms)			
Smoking dur	ation (years)			
Exposed		16.14	3.28	10-20
Control		14.28	2.56	8-17
Smoking inte	ensity (number of ciga	rettes per d	lay)	
Exposed				15-20
Control				15-20
WE (years)				
Exposed	50	21.94	8.00	5-35
Control	40	19.67	6.17	13-36
DOE (years))			
Exposed	50	17.96	8.12	5–35
Control	40	0.00	0.00	0.00
$ED_5(mSv)$				
Exposed	50	9.87	6.84	0.16-47.38
Control	0			0.00
$ED_{I}(mSv)$				
Exposed	50	2.19	1.41	0.00-10.10
Control	0			0.00
PM gloves				
Exposed	20			
Control	0			
PM-lead app	pron			
Exposed	15			
Control	0			
PM glasses				
Exposed	30			
Control	0			

f female, *m* male, *fs* female smokers, *ms* male smokers, *WE* working experience, *DOE* duration of occupational exposure to radiation, ED_5 effective doses for last 5 years, ED_1 effective doses for last year, *PM* protective measurements

Personal dosimeters were used for estimation of the external exposure. The amounts of IR absorbed by the subjects' bodies were measured with thermo-luminescent dosimeters (TLD) once a month during occupational exposure time and expressed in mSv as the mean of annual effective doses for the period of exposure. Routine personal dosimetry was performed using an automatic TLD reader (Harsaw Model 6600 with LiF:Mg Ti cards; Thermo Scientific, 81 Wyman Street, Waltham, MA 02454). The glow curve was used for sophisticated data processing with Harsaw CGCD (computerized glow curve deconvolution) software (German et al. 2000).

The controls matched the exposed subjects in gender, age and smoking habits.

Detailed information on the occupational and medical history of all examined subjects was obtained by completing a targeted questionnaire. This included demographic data, smoking, alcohol intake and use of medication, use of protective measurements and duration of exposure to radiation or chemicals. Only subjects who had not been exposed to any mutagen except for IR (exposed group) or any mutagen at all (control group) were included in the analysis. The participants signed written informed consent, and study protocol was reviewed and approved by the Serbian Institute of Occupational Health (SIOH) ethical committee. Examinations were performed in accordance with the ethical standards of Helsinki Declaration of 1975 as revised in 2013.

Chromosomal aberrations, premature centromere division and micronuclei

Chromosomal aberrations and PCD in PBL were analyzed according to a standard protocol (IAEA 2001). Lymphocyte cultures were prepared by adding whole blood to PB-MAXTM Karyotyping Medium (Gibco). Lymphocytes were incubated in vitro for 48 h at 37 °C. During the last 2 h of incubation, colchicine (0.05 μ g/ml; Sigma-Aldrich, 3050 Spruce St., St. Louis, MO 63103) was added to the medium. The cells were exposed to hypotonic solution (20 min) by stepwise addition of 0.075 M KCl followed by fixation (3 × 20 min) with cold methanol/acetic acid (3:1). Fixed cells were spread on slides and dried over a flame. The slides were coded and scored blind under a light microscope.

For the cytokinesis block micronucleus (CBMN) test, lymphocytes were cultivated the same way as for classical cytogenetic analysis (IAEA 2001). After incubation for 44 h, 0.1 ml of cytochalasin B (Sigma-Aldrich, 3050 Spruce St., St. Louis, MO 63103) solution in DMSO (Sigma-Aldrich, 3050 Spruce St., St. Louis, MO 63103) (final concentration of 3 μ g/ml) was added and cultivation continued for another 24 h. The cultures were then treated with 0.9 % NaCl solution, followed by cold hypotonic solution to lyse red blood cells. The supernatant was removed and replaced with fixative, consisting of methanol/acetic acid (3:1) with 1 % formaldehyde. After incubation for 20 min at room temperature, the cells were washed with four changes in fixative without formaldehyde. The cells were then resuspended gently and the suspension dropped onto clean glass slides and allowed to dry. Cells were stained using 10 % Giemsa in distilled water.

Cytogenetic analysis

Chromosomal aberration parameters: chromatid (CB) and isochromatid (iCB) breaks, acentrics (ACE), dicentrics (DIC) and ring chromosomes, as well as three parameters of PCDs, were evaluated for at least 200 metaphase cells, while five parameters of micronuclei (MN) were evaluated for at least 1000 binucleated cells. PCD was diagnosed when separation between sister chromatids was equal to or more than the thickness of the chromatid (Madan et al. 1987; Ruskovsky et al. 2003). Three PCD parameters were included in the analysis: tPCD (number of metaphases per individual with 11 or more chromosomes with separated chromatids), PCD_{1-5} (number of metaphases per individual with 1-5 chromosomes with PCD) and PCD₅₋₁₀ (number of metaphases per individual with 6-10 chromosomes with PCD). Five MN parameters were included in the analysis: total number micronuclei (ΣMN), frequency of binuclear (BN) cells with one, two, three and four MN (MN_1 , MN_2 , MN₃ and MN₄, respectively).

Statistical analysis

The incidence was estimated for several predictor variables including IR exposure (control group vs. exposed group), gender (men vs. women), smoking habit (smokers vs. non-smokers), age and working experience (WE). The control and exposed groups were coded as "control" and "exposed," respectively, gender qualifications were coded as "m" for male and "f" for female, and smoking habits were coded as binary (0 or 1). Age, WE and effective doses were used as continuous predictor variables.

The data for CAs, PCDs and MNi were tested for normal dispersion. Parameters PCD_{5-10} , ΣMN and MN_1 fitted a normal distribution curve (p > 0.05). The frequencies of total aberrant cells (tAC), DIC, ring chromosomes, ACE, CB and iCB, tPCD, PCD_{1-5} , MN_2 , MN_3 and MN_4 showed significant differences from normal distribution (p < 0.05).

A pretest—independent sample t test—was performed to determine the possible influence of gender and smoking habit on numeric variables. Then, the effects of those nonparametric predictor variables (IR exposure, gender qualification and smoking habit) on differences in incidence of CAs, PCDs and MNi were tested by an independent sample t test for parameters with normal distribution and the Mann–Whitney U test for parameters with nonparametric distribution.

Correlations between continuous predictor variables (age, WE, ED_1 and ED_5) and CAs, PCDs and MNi parameters were calculated as Pearson's correlations for variables with normal distribution and Spearman rank correlations for nonparametric variables.

Multinomial logistic regression was performed in order to predict affiliation to the control or exposed group for all numerical variables (tAC, ACE, DIC, CB, iCB, total PCD, PCD₁₋₅, PCD₅₋₁₀, Σ MN, MN₁, MN₂, MN₃ and MN₄) as predictor variables. The criterion for classification to the control or exposed group was the Wald coefficient value and its significance. A probability *p* < 0.05 was considered statistically significant in each analysis.

Diagnostic qualities of applied cytogenetic tests were analyzed by determining contingency test parameters (sensitivity, specificity and cutoff value), while graphical analysis is presented by a receiver operating characteristic (ROC) curve, used in medicine to determine cutoff values for clinical/diagnostic tests, and ROC space, known as AUC (area under curve), as an important measure of the accuracy of the clinical test. Uncertainties for all measures of accuracy (including AUC) were expressed as 95 % confidence limits. The analyses were performed using SPSS 17.0 statistics (SPSS Inc.).

Results

A total of 18,000 metaphase cells were scored for CAs and PCDs: 8000 in the control group and 10,000 in the exposed group (200 per individual). A total of 90,000 binucleated cells were analyzed for MN by the CBMN test: 40,000 in the control group and 50,000 in the exposed group (1000 per individual). The results are presented in Table 2. In comparison with the control group, incidence of cells with aberrations/micronuclei was increased by twofold for PCD_{1–5}, Σ MN and CBs, by threefold for iCBs and PCD_{5–10}, by fivefold for DICs, by sevenfold for ACE and by 8.5-fold for tPCD. No cells with ring chromosomes were found in the control group, but in the exposed group, the ratio was 1 cell with a ring per 2500 cells.

The means for CAs, PCDs and MNi were significantly higher in the exposed group, as illustrated in Fig. 1, which gives the values of all CAs and PCDs per 200 analyzed metaphases of each examined individual.

Gender, age and smoking habit were without influence on numerical variables, in both control and exposed groups, except for parameters of CBMN assay, which mean values were higher in females, smokers and subjects older than 45 years.

 Table 2
 Ratios of CAs, PCDs and MN per number of analyzed cells for the control and exposed groups

CAs, PCDs, MN	Control	Exposed	Ratio 5x	
DIC	1/1600	1/303.03		
ACE	1/533	1/75.75	7x	
RING	/	1/2500		
CB	1/615	1/294	2x	
iCB	1/1143	1/400	2.85x	
tPCD	1/800	1/94.33	8.5x	
PCD ₁₋₅	1/49	1/23.53	2x	
PCD ₅₋₁₀	1/116	1/43.1	2.7x	
ΣΜΝ	1/117,30	1/48.26	2.43x	

DIC dicentric chromosome, *ACE* acentric fragment, *RING* ring chromosome, *CB* chromatid break, *iCB* isochromatid break, *tPCD* number of metaphases per individual with 11 or more chromosomes with separated chromatids, PCD_{1-5} number of metaphases per individual with 1–5 chromosomes with PCD, PCD_{5-10} number of metaphases per individual with 6–10 chromosomes with PCD, ΣMN total number of micronuclei

In order to test PCDs as a new genotoxic endpoint, correlations between CAs, MNi and PCDs were examined with the Spearman rank test on the whole sample. There were positive linear correlations between CAs and PCDs for the DIC–tPCD, DIC–PCD_{5–10}, ACE–tPCD, ACE–PCD_{1–5}, CB–tPCD and CB–PCD_{5–10} variable pairs, which indicated that an increase in incidence of one variable was coupled with an increase in incidence of the other. For PCDs and MNi, the only correlation was between MN₄ and tPCD.

Duration of occupational exposure was positively correlated with total number of aberrant cells (tAC) and acentric fragments (ACE)—regarding parameters of CAs and total

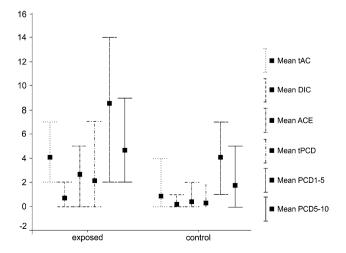


Fig. 1 Means and standard deviations of CAs and PCDs in exposed and control groups

 Table 3
 Spearman correlations

(exposed group)

	tAC	DIC	RING	ACE	CB	iCB	tPCD	PCD_{1-5}	ΣMN	MN_1	MN ₃	DOE
tAC												0.768
ACE												0.753
iCB	0.448											
tPCD		0.515	0.293	0.556	0.318							0.647
PCD ₁₋₅				0.348			0.524					0.659
PCD_{5-10}		0.330			0.295		0.437	0.388				0.670
ΣMN	0.312					0.539						0.818
MN_1						0.408			0.749			0.761
MN_2	0.345					0.355			0.533			0.610
MN_3			0.298			0.296			0.592			
MN_4							0.364		0.438		0.449	

DOE duration of occupational exposure, *tAC* total number of cells with aberrations, *DIC* dicentric chromosome, *RING* ring chromosome, *ACE* acentric fragment, *CB* chromatid break, *iCB* isochromatid break, *tPCD* number of metaphases per individual with 11 or more chromosomes with separated chromatids, PCD_{1-5} number of metaphases per individual with 1–5 chromosomes with PCD, PCD_{5-10} number of metaphases per individual with 1–5 chromosomes with PCD, PCD_{5-10} number of metaphases per individual with 6–10 chromosomes with PCD, ΣMN total number of micronuclei, MN_1 frequency of BN cells with one micronucleus, MN_2 frequency of BN cells with three micronuclei, MN_4 frequency of BN cells with four micronuclei

Only significant correlations were presented

number of micronuclei Σ MN and parameters of micronuclei distribution MN₁ and MN₂. Furthermore, positive correlation was found between duration of occupational exposure (DOE) and all PCD parameters. However, this was not the case for physical dosimetry data (ED₁ and ED₅) and PCDs, or other analyzed cytogenetic parameters (CAs and MNi). Table 3 presents significant correlations for all listed parameters and for the exposed group only.

A logistic regression test was performed in order to predict affiliation of subjects to the control or exposed group for all numerical variables (tACs, ACE, DIC, CB, iCB, total PCD, PCD₁₋₅, PCD₅₋₁₀, Σ MN, MN₁, MN₂, MN₃ and MN₄) as predictor variables. The criterion for classification to the control or exposed group was a Wald coefficient value higher than 1.

ROC analysis was applied in order to investigate ability of all analyzed (particularly PCD) parameters to separate groups, since it is used in medicine as a fundamental tool for diagnostic test evaluation, by determination a cutoff value for a clinical/diagnostic test. Among the listed variables, PCD₅₋₁₀ was found to be the best. From ROC analysis optimum value of 3, 50 cells (per 200 cells examined), with 6–10 chromosomes with PCD, as the cutoff value was selected. The sensitivity of this value is 75.0 %, and the specificity (the ability to separate exposed from unexposed individuals) is 87.5 %. The area under the curve (AUC) is 90.8 % for this value. Figure 2 shows the ROC analysis for PCD₅₋₁₀, while in Fig. 3, ROC analysis for three parameters with the highest Wald coefficient values (PCD₅₋₁₀, tAC and Σ MN) can be seen.

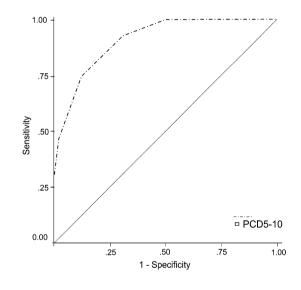


Fig. 2 ROC analysis for the PCD₅₋₁₀ parameter

Discussion

Occupational exposure is an especially delicate area of research, where effects of chronic exposure to low doses play an important role in consideration of the biological response to IR (Hoeijmakers 2001; Jovičić et al. 2009; Miller et al. 2008; Mothersill and Seymour 2004). It is therefore, always useful to investigate new biomarkers visible on the first, cytogenetic level, which can indicate effect of genotoxic exposure. Information on radiation effects can be considered in the context of cancer risk (Preston 2005),

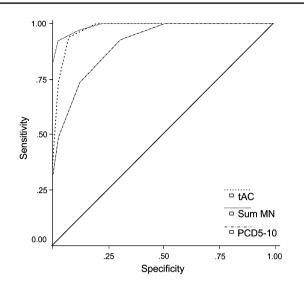


Fig. 3 ROC analysis for PCD_{5-10} , tAC and ΣMN

since there is recent evidence indicating that low doses of radiation exert both suppression and induction of neoplastic transformation (Ko et al. 2006).

The results presented in this paper demonstrate the significantly higher incidence of all cytogenetically visible parameters: CAs, PCDs and MNi in individuals exposed to low radiation doses of radionuclides (ranged from 0 to approximately 10 mSv per year) than in unexposed persons. Different authors have also reported increments of CA and MN incidence in health workers occupationally exposed to IR (Ballardin et al. 2007; Barquinero et al. 1993; Bigatti et al. 1988; Bonassi et al. 1997; Bouraoui et al. 2013; Jovičić et al. 2009; Maffei et al. 2002; Ropolo et al. 2012; Saberi et al. 2013; Sari-Minodier et al. 2007; Thierens et al. 2000; Touil et al. 2002). Similar observations, confirmed by application of FISH procedure, revealed a significant difference (p < 0.001) in incidence between the two groups (exposed and controls) for PCDs as well as for CAs (Jovičić et al. 2010).

Our results also revealed no difference between genders, age and smoking status for the CAs and PCDs. This finding could be expected for at least some of the analyzed CAs (dicentrics and rings) since their formation is caused almost exclusively by ionizing radiation (Ainsbury et al. 2011; Rozgaj et al. 2002). Cytogenetic study of healthy unexposed persons also reported that age and gender were not significant predictors of chromosomal aberrations (Stephan and Pressl 1999). Regarding PCD, authors (Vig et al. 1989) related PCD of acrocentric chromosomes and aneuploidy of some chromosomes and showed that a genderand age-dependent increase in PCD rates of spontaneous aneuploidy of the X chromosome was correlated with PCD (Spremo-Potparevic et al. 2004). All CBMN parameters were significantly higher in females, smokers and subjects older than 45 years. It is well known that the micronucleus prevalence is affected by age, gender, diet, smoking and other lifestyle factors (Antunes et al. 2014; Fenech and Bonassi 2011; Nefic and Handzic 2013).

Correlation analysis revealed positive correlations between CA and PCD parameters as well as some of the parameters characterizing MNi and PCDs in the exposed group. Similarly to the results presented here, some other researchers (Jovičić et al. 2010; Major et al. 1999) reported PCD in individual chromosomes in subjects that already had CAs in their metaphases, and they suggested that the two phenomena (CA and PCD) are not mutually exclusive or dependent on each other. Duration of exposure (DOE) and PCD occurrence were also positively correlated. Positive correlation was also found between DOE and total number of aberrant cells (tAC) and acentric fragments (ACE)-regarding parameters of CAs and total number of micronuclei **SMN** and parameters of micronuclei distribution MN₁ and MN₂. This finding suggests that several genotoxic actions are probably more likely to induce both well-established cytogenetic changes and PCD than short-term exposures. However, no correlation was established between physical dosimetry data (expressed by ED_1 and ED_5 in the exposed group) and PCD, or other analyzed cytogenetic parameters (CAs and MNi). These data refer only to external exposure, while the examined subjects were exposed to radionuclides, which could have entered the body, and therefore, the biological endpoints, found at a significantly higher level in PBL of exposed subjects in comparison with unexposed ones, were better markers of exposure. So, our finding of a lack of correlation between physical and biological dosimetry data may be explained by possible exposure to amounts of radionuclides small enough not to be recorded by personal TLDs, but genotoxic enough to cause changes in genetic material that can be seen at the cytogenetic level. The present data also suggest that personal protection is not at a satisfactory level and adequate personal protection could prevent PCD induction (Major et al. 1999). Since little attention has been devoted to studying PCD phenomenon, the question could be raised whether PCD may be considered as an artifact, or could be regarded as biological expression of the cellular response to the action of genotoxic agents, such as IR. Bühler et al. (1987) suggested that the PCD phenomenon was not an accidental finding. If this is the case, the incidence of PCDs should be different in the control and the exposed group. The results of our study confirmed the previous statement. PCD incidence was significantly lower in the control group than in the exposed group. Furthermore, the applied ROC analysis clearly marked one of the three PCD parameters, PCD_{5-10} , as the best for separation of exposed from unexposed persons.

Our results as well as results of other authors (Major et al. 1999) showed significant influence of exposure to various genotoxic agents (IR and genotoxic chemicals) on premature dividing of centromeres. These findings could be explained by the DNA structure of human centromeric regions and its influence on the timing of centromeric separation. According to research of Shueller et al. (2001) as well as some earlier findings (Way and Willard 1987; Willard 1985), human centromeric regions are rich in repetitive sequences. It has been proposed that centromere separation is a function of the timing of replication of the centromere, which splits into two units, presumably after undergoing some sort of maturation. This suggestion is supported by the fact that the mammalian centromeres replicate during late S phase (Broccoli et al. 1989) but do not separate into two units until late metaphase. Mutations in this satellite region due to action of genotoxic agent, such as IR, can cause gain or loss of entire repeat units and slippage during replication (Sankaranarayanan et al. 2013). This could alter replication time of centromeres and cause early replication and consequently early separation of centromeric regions seen in already metaphase of the cell cycle (Vig 1984; Vig and Zinkowski 1986). Higher prevalence of metaphases with split centromeres can also arise as the result of molecular checkpoint defects in the cell cycle influenced by highly potent mutagenic agents, such as IR (Gollin 2005; Matsuura et al. 2000; Michor et al. 2005; Morgan et al. 1995; Resta et al. 2006).

Conclusion

Most human radiation exposure nowadays involves chronic or highly fractionated doses below 1 Gy and occurs most often in occupational settings. The presence of an unstable CA is the best criterion of health consequences and radiation damage (Jovičić et al. 2009), and standardized techniques, such as CA analysis and the CBMN test, are often used in dose assessment. However, detection and identification of other phenomena related to IR exposure may clarify recent problems in radiobiology concerning the biological response to low doses of IR and its consequences. One of these phenomena is certainly premature centromere division.

Since PCD may be observed as a phenomenon representing the manifestation of chromosomal instability in the exposed persons, our study led us to consider PCD as a possible parameter of genotoxic risk for individuals occupationally exposed to low doses of IR.

Studies of PCD could allow better insight into molecular events underlying chromosomal instability considering that cell cycle progression requires control mechanisms which could be associated with the origin of PCD. The results presented here were obtained from preliminary studies on a relatively small sample by applying only conventional cytogenetic techniques. However, further studies that will include a larger number of subjects, application of different molecular techniques and in vitro investigations should be performed in order to understand better the described phenomenon and its association with established biomarkers of radiation exposure.

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Compliance with ethical standards

Conflict of interest The authors declare that they have no conflict of interest.

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