

Evaluation of radioinduced damage and repair capacity in blood lymphocytes of breast cancer patients

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Abstract

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Genetic damage caused by ionizing radiation and repair capacity of blood lymphocytes from 3 breast cancer patients and 3 healthy donors were investigated using the comet assay. The comets were analyzed by two parameters: comet tail length and visual classification. Blood samples from the donors were irradiated *in vitro* with a ^{60}Co source at a dose rate of 0.722 Gy/min, with a dose range of 0.2 to 4.0 Gy and analyzed immediately after the procedure and 3 and 24 h later. The basal level of damage and the radioinduced damage were higher in lymphocytes from breast cancer patients than in lymphocytes from healthy donors. The radioinduced damage showed that the two groups had a similar response when analyzed immediately after the irradiations. Therefore, while the healthy donors presented a considerable reduction of damage after 3 h, the patients had a higher residual damage even 24 h after exposure. The repair capacity of blood lymphocytes from the patients was slower than that of lymphocytes from healthy donors. The possible influence of age, disease stage and mutations in the BRCA1 and BRCA2 genes are discussed. Both parameters adopted proved to be sensitive and reproducible: the dose-response curves for DNA migration can be used not only for the analysis of cellular response but also for monitoring therapeutic interventions. Lymphocytes from the breast cancer patients presented an initial radiosensitivity similar to that of healthy subjects but a deficient repair mechanism made them more vulnerable to the genotoxic action of ionizing radiation. However, since lymphocytes from only 3 patients and 3 normal subjects were analyzed in the present paper, additional donors will be necessary for a more accurate evaluation.

Key words

- Comet assay
- DNA damage
- DNA repair
- Ionizing radiation
- Breast cancer
- Blood lymphocytes

Introduction

Breast cancer has attracted increasing attention over the last years because of its high incidence among women. In industrialized countries one in ten women develops this

type of cancer some time during her life (1).

The etiology of breast cancer is multifactorial, including various risk factors such as environmental, hormonal and hereditary ones. Most cases (about 80%) are considered sporadic, being caused by somatic events and

about 20% have a family history including a 5% rate of autosomal dominant mutations of high penetrance (2).

Alterations of the tumor suppressor genes BRCA1 and BRCA2 confer a high risk of breast cancer. Mutations in the BRCA1 gene are involved in about 50% of families with a high incidence of breast cancer and in at least 80% of families with a high rate of early breast and ovarian cancers (3). Similarly, mutations of the BRCA2 gene cause early breast cancer in women and also a higher risk in men (4).

Although breast and ovarian cancers are types of neoplastic disease with a strong familial component, many uncertainties exist about the role of hereditary predisposition because of many factors that impair analysis. These include the existence of sporadic cancer and mutations of variable penetrance, in addition to the heterogeneity of the disease as revealed by epidemiological evidence and molecular analysis (1).

The deficiency in repair has been pointed out as a factor of susceptibility to cancer development. The biological importance of the DNA repair mechanism in cancer development is illustrated by recessive autosomal disease such as chromosome fragility syndrome. Affected individuals, i.e., subjects with xeroderma pigmentosum, Fanconi's anemia, or ataxia telangiectasia, are deficient in a type of repair and are at high risk for malignancy when exposed to specific mutagens such as UV light, alkylating agents and ionizing radiation (5). There are many cell repair mechanisms and, depending on the type of DNA lesion, a specific repair mechanism is initiated. Recent molecular studies have shown that the BRCA1 and BRCA2 genes are involved in DNA repair. Both protein products act directly or indirectly with yeast Rad51 protein homologues or *Escherichia coli* RecA protein which participates in DNA double-strand break or recombinant repair (2,6).

In addition to these genes, other tumor

suppressors such as the p53 and AT genes, which are involved in breast cancer, participate in cell repair processes (7,8). On this basis, the analysis of the association between cell repair capacity and genotoxic exposure to environmental agents in breast cancer patients would be of great interest.

There are few studies on this subject and some data reported in the literature are contradictory. Some investigators found a high sensitivity and a reduced repair capacity in peripheral blood lymphocytes from breast cancer patients when exposed to X-rays, gamma and UV light, as evaluated by the determination of chromosome aberrations (9-11) and by the micronucleus test (12). While Hsu et al. (13), using bleomycin in a mutagenic test, did not find a significant difference between blood cells from breast cancer patients and healthy subjects in terms of number of chromatid breaks per cell, Jaloszynski et al. (14) reported a high sensitivity and reduced repair capacity in peripheral blood cells from breast cancer patients using the comet assay. Rothfuss et al. (15) reported a higher sensitivity to gamma radiation and oxygen peroxide in patients with BRCA1 mutation than in healthy subjects using the micronucleus assay but not when using the comet assay.

In the present study, we investigated the damage induced by ^{60}Co gamma radiation and the repair capacity of 3 breast cancer patients (2 with cancer cases in their families) by analyzing peripheral blood lymphocytes immediately and 3 and 24 h after irradiation by the comet assay. The objective was to determine if the irradiated cells of these patients present the same radiosensitivity and repair the damage with the same intensity as observed in healthy individuals.

Material and Methods

Donors

Blood samples from 3 healthy subjects

and from 3 patients with breast cancer were obtained for analysis from Instituto de Radioterapia de São Paulo (São Paulo, SP, Brazil), by written authorization. This project was evaluated and approved by the Ethics Committee on Research of IPEN-CNEN/SP (No. 006/CEP).

Healthy donors. The group of healthy donors consisted of 2 women (donor A, 49 years and donor B, 40 years) and 1 man (donor C, 45 years), all of them nonsmokers who were not consuming alcohol or taking any medication at the time of blood collection. None of them had cases of cancer in the family.

Breast cancer patients. This group consisted of 3 women with ductal breast carcinoma (donor D, 47 years; E, 72 years; F, 65 years) in the primary stage and with no metastasis. All patients had been mastectomized but none had been submitted to chemotherapy or radiotherapy at the time of blood sampling. Donor D had an uncle with intestinal cancer, an aunt with breast cancer and a brother with skin cancer. Donor F had a brother with prostate cancer and a daughter with ovarian cancer. Donor E had no cancer case in her family.

All donors completed a written questionnaire to obtain information related to their life style, such as dietary habits, medical history and exposure to chemical and physical agents.

Blood sample collection and irradiation

About 5 ml of blood was collected from each donor by venipuncture into heparinized tubes. The samples were then fractionated in Eppendorf tubes in equal volumes of 0.8 ml each and irradiated with a panoramic ^{60}Co source at 0.2, 0.6, 1.0, 2.0 and 4.0 Gy (dose rate of 0.722 Gy/min) at room temperature. One tube was maintained as control (0 Gy).

Comet assay

The alkaline version described by Singh

et al. (16) was used, with some modifications. After irradiation, all samples were kept on ice to avoid repair of radioinduced damage (no incubation period) for the evaluation of initial damage.

For repair evaluation, blood samples were incubated at 37°C for 3 and 24 h after irradiation. These periods were chosen based on literature data about the repair kinetics of induced damage evaluated over periods ranging from minutes to some hours after exposure to low linear energy transfer (LET) radiation (16-18). The period of 24 h was adopted because it was considered sufficient to analyze non-repaired damage.

For each fractionated blood sample, irradiated or not, fully frosted microscope slides (in duplicate) were covered with 300 μl of normal melting agarose (Sigma Chemical Co., St. Louis, MO, USA; 0.75% Ca^{2+} and Mg^{2+} free PBS at 65°C) and maintained at 4°C for 5-10 min for gel solidification. On this gel layer, 5 μl of blood was dissolved in 90 μl of low melting agarose (Sigma; 0.5% Ca^{2+} and Mg^{2+} free PBS at 37°C). After solidification at 4°C, another 90- μl layer of low melting agarose was laid.

The slides were then placed vertically in a cuvette containing lysis solution (2.5 M NaCl, 100 mM EDTA, 10 mM Tris, 1% sodium sarcosinate, 1% Triton X-100, and 10% DMSO) for 2 h at 4°C to remove proteins. After lysis, the slides were placed side-by-side in a horizontal electrophoresis tank (10 x 20 cm, Permatron) and immersed in alkaline buffer, pH >12 (1 mM EDTA and 300 mM NaOH) for 30 min to allow DNA damage expression, and electrophoresis was performed at 25 V, 300 mA (Pharmacia, Uppsala, Sweden) for 30 min at 4°C.

The slides were then neutralized three times with 0.4 M Tris buffer, pH 7.5, for 5 min each time and stained with 50 μl ethidium bromide (20 $\mu\text{g}/\text{ml}$; Sigma), covered with coverslips and maintained in a humid chamber protected from light. All steps after lysis were conducted with no light to avoid

additional damage to the DNA.

Microscope slide analysis

The slides were analyzed with a fluorescent microscope (Carl Zeiss) at 200X, with a 515-560-nm exciting filter and a 590-nm barrier filter. Approximately 50-70 comets were analyzed for each radiation dose. All comets were photographed using black and white ASA 400 TMAX Kodak film.

Damage and repair evaluation

The radioinduced damage was evaluated in two ways:

Comet measurement. The analysis was done on the negative by projecting the comet image as a photo slide (Zoom Cabin). DNA damage was quantified for each cell by measuring the total length (head and tail) according to the criteria adopted by McKelvey-Martin et al. (19).

Damage category. Damage was assigned to 5 classes (0-4) based on the visual aspect of the comets, considering the extent of DNA migration according to the criteria established by Visvardis et al. (20). Comets with a bright head and no tail were classified as class 0 (cells with no DNA migration) and comets with a small head and a long diffuse tail were classified as class 4 (severely damaged cells). Comets with intermediate characteristics were assigned to classes 1, 2 and 3.

Radioinduced DNA damage (DD) and repair capacity (R) were estimated quantitatively 3 and 24 h after irradiation using the equations described by Jaloszynski et al. (14). DD values ranged from 0 to 400 arbitrary units (au), corresponding to situations ranging from no damaged comets to all comets extremely damaged.

$$DD = (n_1 + 2n_2 + 3n_3 + 4n_4)/(\Sigma/100)$$

$R = (DD \text{ immediately after exposure} - DD \text{ after exposure at time 3 or 24 h}) / (DD \text{ imme-}$

diately after exposure - DD_0) 100%, where, DD: DNA damage (au), n_1 - n_4 : number of class 1 to 4 comets, Σ : total number of scored comets, including class 0, and DD_0 : DNA damage in nonirradiated samples.

Statistical analysis

The radioinduced damage and repair capacity 3 and 24 h after irradiation were compared between groups by nonparametric two-way ANOVA. Dose-response curves were fitted according to nonlinear regression:

$$Y = A.e^{(-kD)} + B$$

where, Y: tail length (μm) or DNA damage (au); A, B, k: constants; D: radiation dose (Gy).

The statistical analyses were performed using the Graph Pad Prism Software.

Results

In this study, radioinduced damage and DNA repair capacity of blood lymphocytes from breast cancer patients and healthy subjects were compared by the comet assay 3 and 24 h after ^{60}Co irradiation. Nonirradiated cells showed a nuclear matrix with a fluorescent halo formed by DNA filaments limited to the original nuclear area (nucleoid or nucleus-like structure) (19). The nucleoid diameter was about 28-29.5 μm . The irradiated cells formed images similar to comets with a head and tail, and tail length increased with radiation dose.

Evaluation of nonirradiated cells

Comets of nonirradiated cells from both groups were analyzed by measuring tail length and DNA damage 3 and 24 h after irradiation (Table 1).

No significant differences in basal values were detected between groups by visual classification ($P > 0.05$) although a slight ten-

dency to an increase was observed in the cancer group. In contrast, significant differences were detected by tail length measurement ($P < 0.05$).

No significant difference was observed in either parameter with respect to time of incubation ($P > 0.05$).

Evaluation of radioinduced damage

The results for blood cells irradiated with ^{60}Co gamma radiation from healthy donors (A, B, C) and from breast cancer patients (D, E, F) analyzed at different times after irradiation are presented in Tables 2 and 3.

In both groups, total length and tail length increased with dose, but the values were smaller at 24 h, and head diameter tended to decrease with dose.

As expected, the percentage of affected cells (comets with tails) increased with dose in both groups. Three hours later the frequency of affected cells was higher in pa-

tients at all radiation doses.

The quantitative estimation of radioinduced DNA damage based on visual classification is presented in Table 4. In both groups the DD values increased as a function of dose, with a tendency to decrease with time.

Figure 1 shows the dose-response curve for DNA migration (tail length and DNA damage) fitted by nonlinear regression. There

Table 1 - Basal values obtained by the comet assay for blood samples from 3 healthy donors and 3 breast cancer patients.

Data are reported as means \pm SEM.

Group	Time after irradiation	Tail length (μm)	DNA damage (au)
Healthy	0 h	1.14 \pm 0.11	13.33 \pm 1.76
	3 h	1.32 \pm 0.52	10.00 \pm 9.26
	24 h	1.21 \pm 0.57	12.89 \pm 2.56
Patient	0 h	2.07 \pm 0.19	15.85 \pm 1.89
	3 h	2.39 \pm 1.38	14.03 \pm 1.46
	24 h	4.00 \pm 0.66	14.93 \pm 4.47

Table 2 - Mean values of comet length (head and tail) obtained for blood samples from 3 healthy donors, processed immediately or 3 and 24 h after in vitro exposure to ^{60}Co .

Tail length = Total length - head length.

Time after irradiation	Dose (Gy)	Number of cells	Total length (μm)	Head length (μm)	Tail length \pm SEM (μm)	Number of cells without a tail (%)	Number of cells with a tail (%)
0 h	0.0	168	30.16	29.02	1.14 \pm 0.11	151 (89.9)	17 (10.1)
	0.2	187	46.14	28.67	17.47 \pm 3.80	53 (28.3)	134 (71.7)
	0.6	163	54.84	28.10	26.74 \pm 8.60	34 (20.9)	129 (79.1)
	1.0	155	65.54	29.16	36.38 \pm 5.83	22 (14.2)	133 (85.8)
	2.0	163	72.10	28.74	43.36 \pm 4.27	4 (2.5)	159 (97.5)
	4.0	177	104.40	27.75	76.65 \pm 11.21	0 (0.0)	177 (100.0)
3 h	0.0	171	31.49	30.17	1.32 \pm 0.52	160 (93.6)	11 (6.4)
	0.2	202	32.34	29.58	2.76 \pm 0.63	171 (84.7)	31 (15.3)
	0.6	193	35.46	30.34	5.12 \pm 0.93	143 (74.1)	50 (25.9)
	1.0	180	35.19	30.04	5.15 \pm 0.63	124 (68.9)	56 (31.1)
	2.0	211	41.23	29.43	11.80 \pm 1.85	113 (53.6)	98 (46.4)
	4.0	223	44.46	28.04	16.42 \pm 2.05	92 (41.3)	131 (58.7)
24 h	0.0	163	30.60	29.39	1.21 \pm 0.57	152 (93.3)	11 (6.7)
	0.2	195	33.33	28.60	4.73 \pm 2.02	149 (76.4)	46 (23.6)
	0.6	170	36.69	27.44	9.25 \pm 1.71	110 (64.7)	60 (35.3)
	1.0	152	35.03	27.18	7.85 \pm 1.74	94 (61.8)	58 (38.2)
	2.0	198	40.97	27.09	13.88 \pm 0.56	106 (53.5)	92 (46.5)
	4.0	208	47.07	26.16	20.91 \pm 1.59	85 (40.9)	123 (59.1)

Table 3 - Mean values of comet length (head and tail) obtained for blood samples from 3 breast cancer patients, processed immediately or 3 and 24 h after in vitro exposure to ^{60}Co .

Tail length = Total length - head length.

Time after irradiation	Dose (Gy)	Number of cells	Total length (μm)	Head length (μm)	Tail length \pm SEM (μm)	Number of cells without a tail (%)	Number of cells with a tail (%)
0 h	0.0	198	29.98	27.91	2.07 \pm 0.19	182 (91.9)	16 (8.1)
	0.2	169	34.72	27.47	7.25 \pm 1.48	111 (65.7)	58 (34.3)
	0.6	191	46.31	26.88	19.43 \pm 2.20	62 (32.5)	129 (67.5)
	1.0	180	58.07	25.77	32.30 \pm 1.62	10 (5.6)	170 (94.4)
	2.0	186	71.60	25.97	45.63 \pm 2.00	1 (0.5)	185 (99.5)
	4.0	167	91.79	25.18	66.61 \pm 3.47	0 (0.0)	167 (100.0)
3 h	0.0	206	30.28	27.89	2.39 \pm 1.38	193 (93.7)	13 (6.3)
	0.2	197	36.32	27.55	8.77 \pm 1.73	129 (65.5)	68 (34.5)
	0.6	161	43.50	26.55	16.95 \pm 2.92	78 (48.4)	83 (51.6)
	1.0	170	45.36	26.45	18.91 \pm 3.13	92 (54.1)	78 (45.9)
	2.0	219	49.71	26.70	23.01 \pm 2.45	86 (39.3)	133 (60.7)
	4.0	171	67.28	26.03	41.25 \pm 8.39	36 (21.1)	135 (78.9)
24 h	0.0	183	32.10	28.10	4.00 \pm 0.66	158 (86.3)	25 (13.7)
	0.2	179	33.37	27.79	5.58 \pm 1.18	141 (78.8)	38 (21.2)
	0.6	163	37.02	27.01	10.01 \pm 1.89	118 (72.4)	45 (27.6)
	1.0	186	42.03	26.94	15.09 \pm 5.67	115 (61.8)	71 (38.2)
	2.0	186	45.46	26.40	19.06 \pm 4.32	91 (48.9)	95 (51.1)
	4.0	185	52.72	26.30	26.42 \pm 5.12	72 (38.9)	113 (61.1)

Table 4 - Mean values of DNA damage obtained for blood samples from 3 healthy donors and 3 breast cancer patients, processed immediately or 3 and 24 h after in vitro exposure to ^{60}Co .

Healthy				Patient			
Time after irradiation	Dose (Gy)	Number of cells	DNA damage (au)	Time after irradiation	Dose (Gy)	Number of cells	DNA damage (au)
0 h	0.0	150	13.33 \pm 3.06	0 h	0.0	177	15.85 \pm 3.27
	0.2	154	103.38 \pm 16.30		0.2	163	62.36 \pm 11.12
	0.6	153	164.09 \pm 29.21		0.6	152	125.31 \pm 25.20
	1.0	154	207.90 \pm 37.54		1.0	152	158.05 \pm 53.84
	2.0	154	232.46 \pm 11.55		2.0	152	232.18 \pm 16.44
	4.0	151	290.59 \pm 18.34		4.0	152	290.82 \pm 20.10
3 h	0.0	150	10.00 \pm 3.46	3 h	0.0	194	14.03 \pm 2.54
	0.2	150	31.33 \pm 16.04		0.2	160	62.15 \pm 11.36
	0.6	152	57.00 \pm 16.64		0.6	139	89.06 \pm 16.08
	1.0	152	55.36 \pm 14.45		1.0	149	84.82 \pm 19.44
	2.0	150	82.67 \pm 11.72		2.0	160	162.17 \pm 55.34
	4.0	154	123.97 \pm 11.12		4.0	152	230.28 \pm 78.26
24 h	0.0	148	12.89 \pm 4.44	24 h	0.0	179	14.93 \pm 7.74
	0.2	150	31.33 \pm 16.29		0.2	164	33.71 \pm 6.83
	0.6	152	48.17 \pm 19.80		0.6	157	53.53 \pm 28.15
	1.0	155	49.44 \pm 21.10		1.0	149	76.57 \pm 9.93
	2.0	154	81.72 \pm 40.10		2.0	181	110.97 \pm 9.33
	4.0	150	125.33 \pm 29.14		4.0	150	197.33 \pm 86.01

was a significant increase in DNA migration as a function of dose at all 3 periods of time analyzed.

Both groups showed a similar response when analyzed immediately after exposure. Analysis of variance showed that there were no differences in comet length between groups ($P>0.05$), but visual classification showed a significant difference ($P<0.05$).

Both tail length and DD values showed a considerable reduction after 3 and 24 h when compared with the values obtained immediately after exposure, with significant differences between groups after 3 h ($P<0.01$) and 24 h ($P<0.05$). The breast cancer group presented a greater quantity of damage than controls.

Evaluation of cell repair

Repair capacity of radioinduced damage in blood cells was estimated by the formula described by Jaloszynski et al. (14). The individual values and means at 3 and 24 h are illustrated in Figure 2.

There were interindividual differences in repair capacity of blood lymphocytes in both groups, but this variability was more marked among patients.

Figure 2 shows that radioinduced damage was less efficiently repaired among patients than among controls after all doses at both times of analysis. Among patients, the blood cells of donor E showed a better effi-

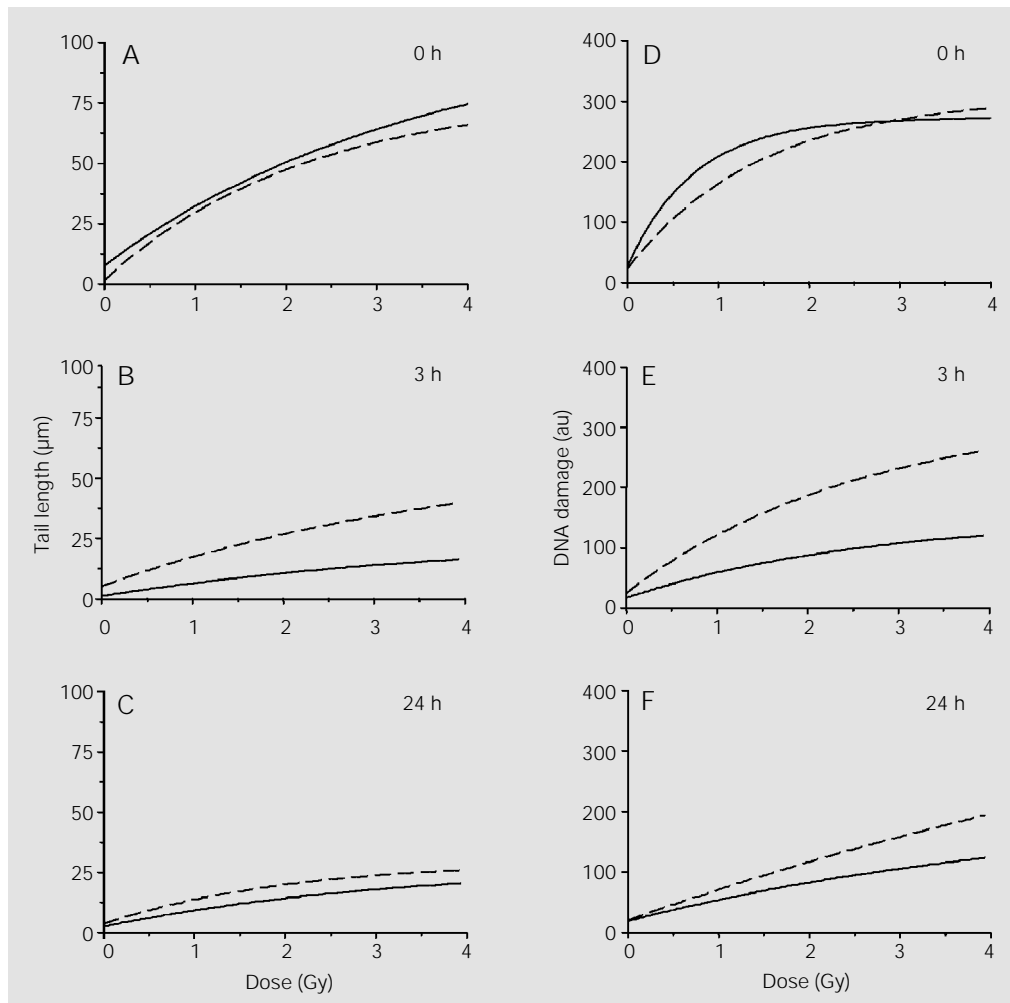
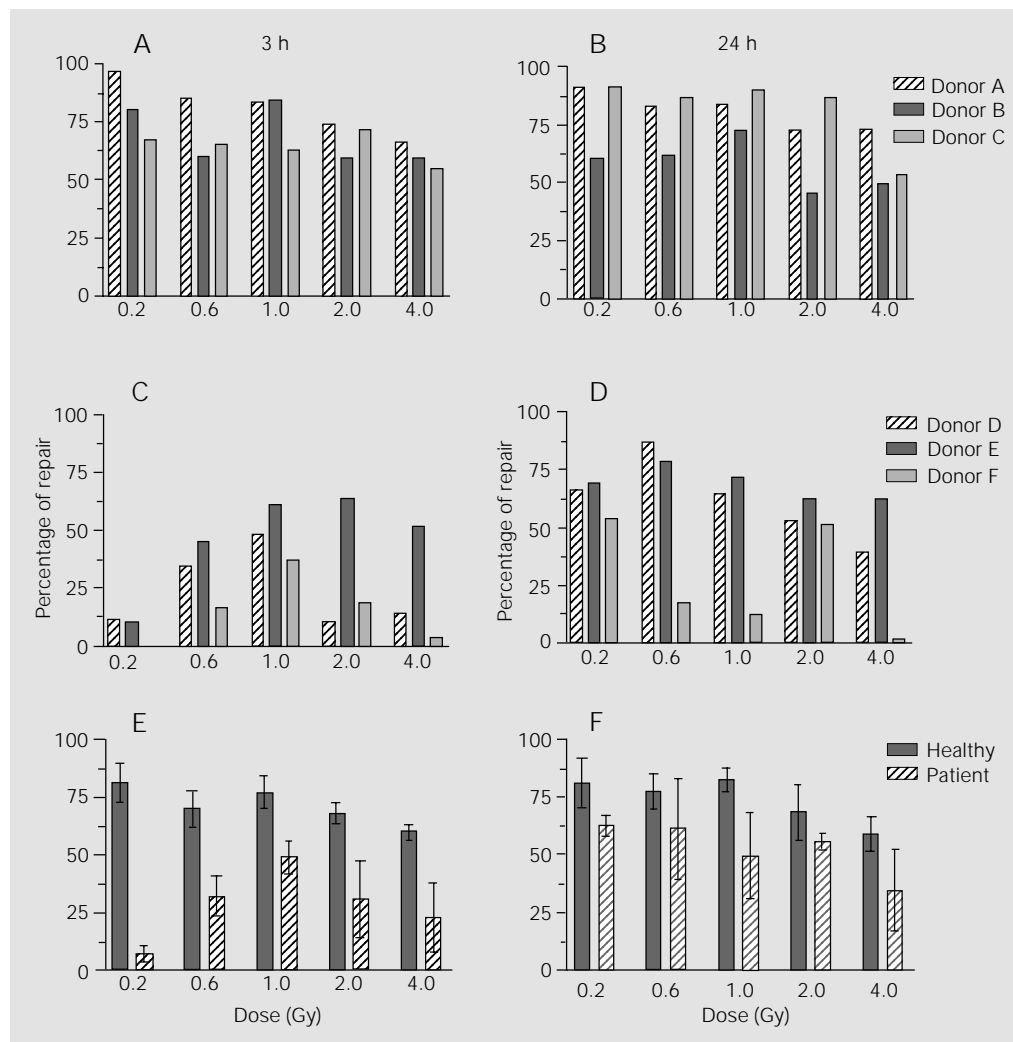


Figure 1 - Dose-response curves of DNA migration obtained for blood samples from 3 healthy donors (full lines) and 3 breast cancer patients (broken lines), processed immediately (A, D) or 3 (B, E) and 24 h (C, F) after in vitro exposure to ^{60}Co .

Figure 2 - Repair capacity observed in blood samples from 3 healthy donors (A, B), 3 breast cancer patients (C, D) and group means (E, F), 3 and 24 h after *in vitro* exposure to ^{60}Co .



ciency in repair capacity. After 24 h, only donor E had a performance equal to controls.

Among controls, donor C had a less efficient repair after 3 h, but presented the best efficiency after 24 h. We observed a tendency to a decrease in repair capacity of blood lymphocytes with increasing doses in both groups at 3 and 24 h. The repair observed in peripheral blood cells from healthy donors after 24 h did not differ from that observed after 3 h, whereas this difference was evident among patients. Statistical analysis showed that the repair capacity of the groups differed both 3 h ($P < 0.001$) and 24 h ($P < 0.05$) after irradiation.

Discussion

Ionizing radiation is an etiologic agent known to act on the induction of breast cancer (21), but on the other hand it is a therapeutic modality used in cancer treatment. Therefore, a better knowledge of the action of ionizing radiation on the cellular response is of importance both clinically and therapeutically.

The head diameters obtained here for cells without DNA migration agreed with data reported by other laboratories (20,22).

A significant difference in relation to basal values was observed between groups

by tail measurement, but not by visual classification. This incongruent result may be explained by the fact that cells with little damage barely differ from cells with no damage so that visual analysis may not permit a distinction between them.

Statistical analysis showed no differences in results after 3 and 24 h of incubation. We conclude that the maintenance of blood samples for 24 h does not cause additional DNA lesions.

About the radioinduced damages, the dose-response curves for DNA migration (tail length) obtained just after irradiation showed an increase in tail length as a function of radiation dose. The increasing dose caused an increase in DNA lesions and so damaged filaments migrated to the anode during the electrophoretic run, giving origin to the comet tails. Statistical analysis showed no difference between groups, whereas the difference determined by visual classification (DD), although small, was considered significant. This result may be explained as a consequence of the slight difference between class 0 and class 1.

In this context, we conclude that both groups had an analogous response when analyzed immediately after exposure. The initial radioinduced damage may have been quantitatively similar.

The quantitative estimation of repair capacity in blood lymphocytes showed that most of the radioinduced damage in the healthy group was repaired within 3 h while patients had more lesions even after 24 h. These data show that the breast cancer donors analyzed in this study had deficient radioinduced damage repair. These data do not agree with those reported by Hsu et al. (13) who used bleomycin as a mutagenic test in peripheral blood lymphocytes from patients with various cancer types. The authors found a higher frequency of chromatid breaks per cell in patients with lung, colon and upper digestive tract cancer but not in breast cancer patients. They suggested that sensi-

tivity to the mutagenic agents may play an important role in the carcinogenesis of tissues and organs in contact with the external environment, a fact that does not occur with mammary tissue.

In contrast, Jalszynski et al. (14) reported that breast cancer patients were more sensitive to bleomycin by expressing higher level of DNA damage than controls, as observed in the comet assay.

The less efficient repair observed in blood lymphocytes in the present patients may explain the high basal level of damage.

Since carcinogenesis may originate from defects in DNA repair (14,23), we may raise various hypotheses. The reduced repair capacity observed in these patients may be a consequence of mutations in genes involved in cell cycle control and DNA repair regulation. The tumor suppressor genes BRCA1, BRCA2, p53 or ATM are fundamental factors for the maintenance of genome integrity (2,24).

Rothfuss et al. (15) observed that familial breast cancer patients with a mutated BRCA1 gene show deficiency in radioinduced damage repair as detected by the micronucleus test. These data show a direct relation between BRCA1 mutation and radioinduced damage repair. Therefore, molecular analysis of the genes revealed that somatic mutations of either BRCA1 or BRCA2 are rare events in the majority of sporadic cases of breast and ovarian cancer (2,6,23,25). Then, some authors such as Futreal et al. (26) raised the hypothesis that sporadic and familial tumors have different etiologies, suggesting that BRCA1 and BRCA2 may not play a critical role in the genesis of sporadic cancer cases.

If we consider the hypothesis that donors D and F are carriers of mutated genes, we could partially explain their significantly lower repair capacity than donor E who has no history of cancer in her family, although no molecular gene analysis was done. Donors D and F have first- and second-degree

relatives with ovarian and breast cancer, besides other types of cancer in the families. There seems to be a predisposition to cancer development in their families, although we cannot rule out the hypothesis of exposure of the entire family to the same carcinogenic agent.

Other genes may be involved but further studies are necessary to test this possibility. One of the possible genes involved is p53, which has been described in almost all human cancers, including breast cancer (27), and plays a role in the repair mechanism of radioinduced damage (9). There is evidence that p53 and rad51 are involved *in vivo* (28).

We cannot rule out the possibility of an ATM gene action since this gene acts on BRCA1 phosphorylation after exposure to ionizing radiation as an initial signal for the recombinational repair process of DNA double-strand breaks (24). The sensitivity to ionizing radiation is also observed in ataxia telangiectasia patients who are defective in DNA strand break repair. The disease is related to ATM gene mutation and heterozygotes are 0.5-1% of the population more predisposed to breast cancer (2).

These observations suggest the occurrence of other genes that can act in the same way as suppressor genes. On this basis, various other molecular targets for sporadic tumors may exist (26).

In relation to controls, the blood cells from donor C showed the most efficient repair after 24 h, suggesting a slower repair mechanism than in the other donors.

Patient age was not an important factor because the oldest one, donor E, showed the best repair. The control group ranged in age from 40-50 years. A possible influence of age may have affected the comparison between groups.

The relation between age and damage susceptibility in somatic cells had been studied but data are contradictory. Duffaud et al. (29) and Peace and Succop (30) observed an increase in the frequency of affected cells

with aging. This could be attributed to the accumulation of genetic damage in cells (31) or to the aging process, such as altered cellular metabolism and/or decrease in the efficiency of DNA repair (32). Other authors, however, did not find any association between age and genetic damage in cells (33,34).

The influence of gender is not significant for the biological response to environmental genotoxic agents (29,30), although some evidence shows that women are more susceptible to some carcinogenic agents (35).

In addition to these considerations, there is the possibility that the difference between groups was a consequence of the pathological stage (malignancy) of the tumors. Other factors such as immunological, metabolic and dietary ones, also considering that life style is influenced by various environmental agents that can act synergistically, have been suggested by several authors (29,33). Therefore, nothing that could explain the difference between patients and healthy donors was observed in the replies to the questionnaire.

Ionizing radiation is known to induce multiple DNA lesions (17). Although the comet assay does not permit the direct identification of the damage or repair mechanism involved, it is known that the pH of the electrophoresis buffer is a fundamental factor: under neutral conditions, double-strand breaks are the type of damage most frequently detected, while under alkaline conditions (pH >12), as used in the present study, single- and double-strand breaks as well as alkali-labile lesions make a significant contribution to the DNA quantity found in comet tails (16,18).

Studies of repair kinetics in blood cells from healthy donors showed that many low LET radioinduced lesions such as base damage are repaired within the first 15 min, while a slower repair mechanism that acts on alkali-labile lesions, single-strand breaks and principally double-strand breaks is completed within

2 h (16-18). Olive (36) described a slower repair system for more complex lesions, which was completed only after 30-40 h.

Based on this information, we suggest that the discordant results obtained by Rothfuss et al. (15) may have been due to the period of time used to check the breast cancer cell repair. The authors did not find a significant difference in repair induced by 2 Gy ⁶⁰Co analyzed one hour later by the comet assay, but did find a difference by the micronucleus test used to evaluate the non-repaired damage.

Double-strand breaks are among the most difficult radioinduced lesions to repair. Non-repaired damage can result in cell death, aging and cancer.

Both parameters adopted in the present study, i.e., comet measurement and visual classification, proved to be safe and sensitive and therefore could be used as an alternative method for the evaluation of cellular response.

The results showed that cellular radio-

sensitivity depends not only on the amount of imposed initial damage, but also on the cell repair capacity, among other factors.

Since in the present study only a small number of patients and controls was analyzed, it is necessary to investigate a larger number of donors including other family members, with an equal distribution of patients and healthy donors with respect to age, gender and other risk factors of increased DNA fragility, and to perform molecular analysis of possible genes involved in breast cancer susceptibility for a more accurate evaluation.

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