

# 17- $\beta$ -estradiol affects nuclear image properties in MCF-10F human breast epithelial cells with tumorigenesis

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Received June 22, 2007; Accepted August 10, 2007

**Abstract.** Treatment of the human breast epithelial cells MCF-10F with 17- $\beta$ -estradiol (E2) induces transformation and tumorigenesis. E2-transformed MCF-10F cells are known to exhibit progressive loss of ductulogenesis, and invasive and tumorigenic phenotypes. Although DNA amounts and chromatin supraorganization change in E2-transformed MCF cells, no comparative study has yet been undertaken in the resulting cells selected for aggressive invasiveness (C5) and tumor generation in a heterologous host. The aim of this study was thus to determine whether changes in Feulgen-DNA content and chromatin supraorganization are involved during E2-induced transformation and tumorigenesis of the MCF-10F cells. Image analysis was performed for nontransformed and E2-transformed MCF cells, highly invasive cells (C5), and for cell lines (C5-A6-T6 and C5-A8-T8) derived from tumors generated by injection of C5 cells in SCID mice. A decrease in Feulgen-DNA amounts and nuclear sizes induced by E2 treatment was accentuated with selection of the highly invasive tumorigenesis potential. However, in the tumor-derived cells a high variability in cellular phenotypes resulted inclusively in near-polyploidy. Significant changes in textural parameters, including nuclear entropy, indicated chromatin structural remodeling with advancing tumorigenesis. An increased variability in the degree of chromatin packing states in the E2-transformed MCF cells is followed by reduction in chromatin condensation and in contrast between condensed and noncondensed chromatin in the highly invasive C5 cells and tumor-derived cell lines. Studies on epigenetic mechanisms involving DNA methylation and/or the histone code would contribute to a better interpretation of the chromatin supraorganization changes reported herein.

## Introduction

Treatment of MCF-10F estrogen receptor  $\alpha$  (ER- $\alpha$ )-negative immortalized human breast epithelial cells with 17- $\beta$ -estradiol (E2) results in anchorage-independent growth, colony formation in agar-methocel, increase in cell proliferation and in invasive capability in Matrigel (1), loss of heterozygosity in chromosomes 13 and 17 (2,3), and loss of the 9p11-13 locus (4). The 9p11-13 locus is known to contain a putative tumor suppressor gene (5). Treatment of the MCF-10F cells with E2 twice a week for two weeks also results in changes in DNA amounts, nuclear sizes and chromatin higher order packing states (supraorganization) (6). Many of these changes are similar to the effects caused by the carcinogen benzo[a]pyrene (BP) which are indicative of neoplastic transformation *in vitro* (2,3,6,7).

The above-mentioned findings suggest that transformation by E2 may occur by a mechanism independent of the presence of ER- $\alpha$ , since the expression of the cell phenotypes indicative of neoplastic transformation as well as the changes in Feulgen-DNA amounts and nucleus image properties in E2-treated MCF-10F cells are not abrogated when these cells are treated simultaneously with E2 and the estrogen antagonist, ICI182,780 (6,8).

Selection of the most highly invasive E2-transformed MCF-10F cells in Matrigel chambers has generated transformed cells with phenotypic/genotypic variations correlated with their tumorigenic potential in SCID mice (4). Only invasive cells with a 4p15.3-16 deletion were tumorigenic. The generated tumors (adenocarcinomas) exhibited genomic alterations similar to those described in primary breast cancer, as determined by comparative genomic hybridization (4). Tumors and tumor-derived cell lines exhibited loss of chromosome 4, deletions in chromosomes 3p12.3-13, 8p11.1-21, 9p21-qter, and 18q, and gains in 1p, and 5q15-qter (4). The 9p21-qter region includes the p16 and p15 genes, their lack thus possibly contributing to the progression of the invasive to the tumorigenic states in the natural progression of the disease (4). The chromosome arm 18q, a region frequently lost in breast cancer (9-14) may involve one or more tumor-suppressor genes (4). Amplification of 1p has been reported in breast cancer cell lines (15) and in primary breast cancer (13).

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**Key words:** human breast epithelial cells, 17- $\beta$ -estradiol, tumor-derived cells, chromatin packing states, image analysis

In early E2-transformed MCF-10F cells certain losses of DNA, changes in nuclear sizes, and chromatin remodeling have been assessed by image analysis procedures (6). One of the most heterogeneous distributions of chromatin packing states ever observed, such that an increase in chromatin condensation in certain sectorial nuclear domains coexisted with a decrease in condensation in less packed zones was demonstrated by high nuclear entropy and absorbance variability per nucleus in the Feulgen-stained E2-transformed cells (6). Changes in nuclear image properties, including chromatin supraorganization remodeling, and decreased Feulgen-DNA content, whether similar to those present in the E2-treated MCF-10F cells *in vitro*, have so far not been studied in the highly invasive and resulting tumor-derived cells. This investigation would be required for demonstrating that the previously reported changes in these parameters in early E2 transformed MCF-10F cells could be associated with a complete transformation in a heterologous host.

Nuclear image properties and Feulgen-DNA values were thus evaluated in highly invasive tumorigenic E2-transformed MCF-10F cells and tumor-derived cell lines studied by Russo and co-workers (4) as regards other cell parameters (selection of cells by Matrigel invasion assay, colony formation in agar-methocel, ductulogenic, invasion and tumorigenic assays, verification of cell lineage, screening for E2-induced genetic changes, development of cell lines from tumors, and histopathological, immunocytochemical and DNA fingerprint analyses).

## Materials and methods

**Cells and treatments.** The cells used consisted of the spontaneously immortalized human breast epithelial cell line, MCF-10F, E2-treated MCF-10F cells, the tumorigenic C5 cell line selected from the E2-treated MCF-10F cells, and C5-A6-T6 and C5-A8-T8 cells, obtained from tumors generated in SCID mice injected with C5 cells (4).

MCF-10F cells were cultured in DMEM:F-12 medium containing 1.05 mM calcium, antibiotics, antimycotics, hormones, growth factors and equine serum as described previously (16). E2 cells consisted of MCF-10F cells in their 123rd passage, treated with 70 nM 17- $\beta$ -estradiol as described previously (4). The choice of the E2 concentration was based on previous studies which evaluated colony efficiency in agar methocel, number of solid masses in collagen gel, invasiveness in the Matrigel invasion chamber (2,4,8), and chromatin remodeling (6). E2 cells at passage 10 were used.

Cells expanded from the 10th passage of E2 cells were named C5; these cells developed poorly differentiated adenocarcinomas when injected in 9 out of 10 SCID mice (4). The tumoral cell lines derived from tumors induced in the SCID mice, C5-A6-T6 and C5-A8-T8, were used at passage three, the same passage for which fingerprint and tumorigenic assay analyses have been reported (4).

The cells used were cultured for 96 h (MCF-10F cells) and 48 h (the other cell types) at 70-80% confluence on 9.4 cm<sup>2</sup> plastic chamber slides (17), fixed in an absolute ethanol-acetic acid mixture (3:1, v/v) for 1 min, rinsed in 70% ethanol for 5 min, and air dried. The preparations were then subjected to the Feulgen reaction (4 M HCl at 24°C, 75 min), and

mounted in Cargille oil (Cedar Grove, NJ, USA) ( $n_D=1.54$ ). Three slides were used for each experimental condition.

**Video image analysis.** Two hundred interphase nuclei chosen at random for each experimental condition were measured using Carl Zeiss/Kontron equipment and Kontron KS400 software (Oberkochen/Munich, Germany). The microscopic images were obtained with a Zeiss Axiophot 2 microscope equipped with a Neofluar 40/0.75 objective, optovar factor 2, 0.90 condenser, and light of wavelength equal to 546 nm, provided by a Schott interference filter. A 100-W halogen illuminator, a voltage regulator for light intensity maintained constant at point 4, filter wheels 1 and 2 rotated into position 100 (open position), and a luminous-filter diagram (transmitted light) at its maximal opening were used. These illumination conditions were maintained constant for all nuclei investigated. The images to be processed were fed from the microscope into a Pentium computer through a Sony CCD IRIS/RGB Hyper HAD color video camera. Two threshold values determined which gray value range of the image Input is retained or deleted in the image Output. Threshold values (gray values) named Low (L) and High (H) were determined by moving the borders in the gray value histogram such that nuclear images appeared well segmented from each other and from the background (18). Since 'green' was selected as the parameter 'color', the pixels inside the gray value range (L, H) were displayed in this color. In the present study, L and H threshold values were equal to 0 and 116-120, respectively. Under the operating conditions used, 1  $\mu$ m corresponded to 7.23 pixels. The minimum area possible to be measured with this apparatus corresponded to four pixels.

The software provided quantitative information on 1) geometric parameters such as nuclear total area ( $\mu$ m<sup>2</sup>), nuclear absorbing area ( $\mu$ m<sup>2</sup>), nuclear perimeter ( $\mu$ m), and nuclear feret ratio (minimal feret/maximal feret, as an indication of the elliptical shape of the nuclear absorbing image), 2) densitometric parameters such as mean gray value per nucleus [subsequently converted into absorbances or optical density (OD)], and integrated optical density (IOD), and 3) textural parameters, such as standard deviation of the total densitometric values per nucleus or absorbance variability per nucleus (SDtd), and nuclear entropy (E). SDtd reflects the variability in the degree of chromatin packing state per Feulgen-stained image (18,19). Entropy is considered the number of bits necessary to store the densitometric values per nucleus image or amount of gray value variability in a nucleus. Since it measures the complexity of an image (20), a high entropy value for Feulgen-stained nuclei represents a high contrast between condensed and noncondensed chromatin, whereas a low entropy value is an indicator of an image with fairly constant gray levels (homogeneous image) (18,19,21).

For conversion of the mean gray values into absorbances, gray values of 0 and 255 corresponded to light transmittances of 0 and 100%, respectively. IOD, in this case, Feulgen-DNA content, was equal to the absorbances multiplied by the nuclear absorbing area.

**Statistical analysis.** Calculations were done using Minitab12™ software (State College, PA). Applied tests consisted of one-way analysis of variance for unstacked data and Mann-Whitney U test.

Cells	Nuclear area ( $\mu\text{m}^2$ )		Nuclear absorbing area ( $\mu\text{m}^2$ )		Nuclear perimeter ( $\mu\text{m}$ )		Feret ratio	
	X	S	X	S	X	S	X	S
MCF-10F (control)	247.87	79.86	247.08	79.48	107.43	48.41	0.75	0.10
E2	188.03	72.57	187.71	72.01	79.38	32.89	0.75	0.10
C5	147.03	31.39	146.42	31.25	83.15	23.22	0.84	0.06
C5-A6-T6	160.17	56.97	158.54	53.88	100.90	74.22	0.81	0.08
C5-A8-T8	185.24	67.57	183.09	65.51	123.95	72.75	0.81	0.07

n=200; E2, MCF-10F cells treated with 17- $\beta$ -estradiol; C5, cells expanded from E2; C5-A6-T6 and C5-A8-T8, tumor cells; S, standard deviation; X, arithmetic mean.

Table II. Nuclear phenotypes: densitometric and textural parameters.

Cells	Absorbances (OD)		Feulgen-DNA amounts (IOD)		SDtd		Entropy	
	X	S	X	S	X	S	X	S
MCF-10F (control)	0.46	0.03	113.54	36.72	8.32	2.12	4.99	0.35
E2	0.44	0.04	82.58	28.82	12.04	3.02	5.50	0.35
C5	0.40	0.03	58.66	13.11	8.32	2.40	4.96	0.38
C5-A6-T6	0.38	0.02	60.23	19.50	8.20	2.02	4.94	0.36
C5-A8-T8	0.36	0.02	65.63	22.86	7.69	1.85	4.82	0.34

n=200; E2, MCF-10F cells treated with 17- $\beta$ -estradiol; C5, cells expanded from E2; C5-A6-T6 and C5-A8-T8, tumor cells; S, standard deviation; SDtd, standard deviation of total densitometric values per nucleus; X, arithmetic mean.

## Results

Treatment with E2 decreased the total and Feulgen absorbing nuclear areas, the nuclear perimeter, the absorbances and the Feulgen-DNA values (IOD) of the MCF-10F cells, but increased their SDtd and entropy values. The nuclear feret ratio was not affected (Tables I-III).

In expanded E2 cells (C5 cells) all the parameters, with the exception of the nuclear perimeter and feret ratio, decreased in comparison to the E2 cells. The nuclear perimeter and feret ratio of the C5 cells were bigger than those of the E2 cells (Tables I-III).

Different values for the parameters studied herein, with the exception of those regarding the feret ratio, were obtained for the C5-A6-T6 and C5-A8-T8 tumor cells compared to each other (Table III). Larger nuclear total and absorbing areas, perimeters and Feulgen-DNA values were demonstrated for the C5-A8-T8 tumor cells, whereas larger absorbances, SDtd and entropy values were detected in the C5-A6-T6 tumor cell values (Tables I-III).

In comparison to the C5 cells, larger total and absorbing nuclear areas, and nuclear perimeters, but smaller feret ratio and absorbances were found in the tumor cells. C5 and C5-A6-T6 cells did not differ from each other in terms of Feulgen-DNA, SDtd and entropy values; C5-A8-T8 tumor cells presented larger Feulgen-DNA values but smaller SDtd and

entropy values in comparison to the C5 cells (Tables II and III).

In comparison to the E2 cells, the two tumor cell lines presented larger nuclear perimeters and feret ratio, but smaller absorbances, Feulgen-DNA, SDtd and entropy values. In terms of total and absorbing nuclear areas, C5-A8-T8 and E2 cells did not differ but C5-A6-T6 tumor cells showed smaller values than E2 cells (Tables I-III).

In comparison to the MCF-10F cells, the tumor cells presented smaller total and absorbing nuclear areas, absorbances, and Feulgen-DNA values and larger feret ratios; in terms of nuclear perimeter, the values for the C5-A6-T6 tumor cells were also smaller, whereas those of the C5-A8-T8 cells were bigger than those of the MCF-10F cells. As regards SDtd and entropy, the values presented by C5-A8-T8 tumor cells were smaller than those of the MCF-10F cells, whereas those presented by the C5-A6-T6 did not differ.

Fig. 1 and 2 show the nuclear phenotypes and the distribution of Feulgen-DNA amounts, respectively, in MCF-10F cells treated with E2 and selected for invasivity and tumorigenesis (C5, C5-A6-T6, and C5-A8-T8). In all the E2-transformed cell lines the decrease in Feulgen-DNA amounts is evident; in comparison to the highly invasive C5 cells, the tumor C5-A8-T8 cell line exhibited a large variability in the distribution of their Feulgen-DNA values.

Table III. Statistical comparison of the image analysis parameters of Feulgen-stained E-2-transformed MCF-10F cells.

Parameters	Comparisons	Test	P	Decision
Nuclear area	MCF-10F vs. E2, E2 vs. C5, C5-A6-T6 vs. C5-A8-T8,	Mann-Whitney	<0.010	SS
	E2 vs. C5-A6-T6		0.022	S
	C5 vs. C5-A6-T6		1.000	NS
	E2 vs. C5-A8-T8			
Absorbing area	MCF-10F vs. E2, E2 vs. C5, C5-A6-T6 vs. C5-A8-T8,	Mann-Whitney	<0.010	SS
	E2 vs. C5-A6-T6		0.028	S
	C5 vs. C5-A6-T6		0.740	NS
	E2 vs. C5-A8-T8			
Nuclear perimeter	MCF-10F vs. E2, E2 vs. C5, C5-A6-T6 vs. C5-A8-T8,	Mann-Whitney	<0.010	SS
	MCF-10F vs. C5-A6-T6, MCF-10F vs. C5-A8-T8,			
	C5 vs. C5-A6-T6, E2 vs. C5-A6-T6, E2 vs. C5-A8-T8			
Feret ratio	MCF-10F vs. E2	ANOVA	0.385	NS
	E2 vs. C5, C5 vs. C5-A6-T6		<0.010	SS
	C5-A6-T6 vs. C5-A8-T8		0.946	NS
Absorbances (OD)	MCF-10F vs. E2, E2 vs. C5, C5-A6-T6 vs. C5-A8-T8,	ANOVA	0.000	SS
	C5 vs. C5-A6-T6			
IOD	MCF-10F vs. E2, E2 vs. C5, C5-A6-T6 vs. C5-A8-T8,	Mann-Whitney	<0.010	SS
	C5 vs. C5-A8-T8, E2 vs. C5-A6-T6, E2 vs. C5-A8-T8			
	C5 vs. C5-A6-T6		0.846	NS
SDtd	MCF-10F vs. E2, E-2 vs. C5, C5-A6-T6 vs. C5-A8-T8,	ANOVA	0.000	SS
	MCF-10F vs. C5-A8-T8, C5 vs. C5-A8-T8			
	MCF-10F vs. C5		0.653	NS
	MCF-10F vs. C5-A6-T6		0.753	NS
	C5 vs. C5-A6-T6		0.583	NS
Entropy	MCF-10F vs. E2, E2 vs. C5, C5-A6-T6 vs. C5-A8-T8,	ANOVA	0.000	SS
	MCF-10F vs. C5-A8-T8, C5 vs. C5-A8-T8			
	MCF-10 F vs. C5		0.315	NS
	MCF-10F vs. C5-A6-T6		0.102	NS
	C5 vs. C5-A6-T6		0.554	NS

n=200; E2, MCF-10F cells treated with 17- $\beta$ -estradiol; C5, cells expanded from E2; C5-A6-T6 and C5-A8-T8, tumor cells; SDtd, standard deviation of total densitometric values per nucleus. NS, not significant; S, significant; SS, highly significant.

## Discussion

Decrease in the DNA content and nuclear sizes of the MCF-10F cells with the E2 treatment, as detected with quantification of the Feulgen-DNA responses and confirming previous image analysis data (6), is accented with selection of the highly invasive tumorigenesis potential in these cells. Comparative genomic hybridization in E2-transformed MCF-10F cells has revealed loss of the 9p11-13 locus in E2 cells and of the 9p11-13 and 4p loci in C5 cells (4). In addition, it is known that estrogen compromises the DNA repair system and induces aneuploidy through chromosomal instability and centromere amplification in other cell types (22-24). All these facts may

play a part in the diminution of the Feulgen-DNA values and nuclear sizes as found by image analysis in E2 and C5 cells.

In the tumor-derived C5-A6-T6 and C5-A8-T8 cells the Feulgen-DNA values under statistical terms remained constant or were slightly increased in comparison to the selected invasive tumorigenic cells (C5), depending on which tumor had generated them. The tumor-derived cell lines studied herein also differed from each other as regards other image analysis parameters, which gives information on nuclear sizes and contours (nuclear areas and perimeters), and chromatin packing states (absorbances, SDtd and entropy).

According to Russo and co-workers (4), both C5-A6-T6 and C5-A8-T8 cell lines exhibit loss of chromosome 4,



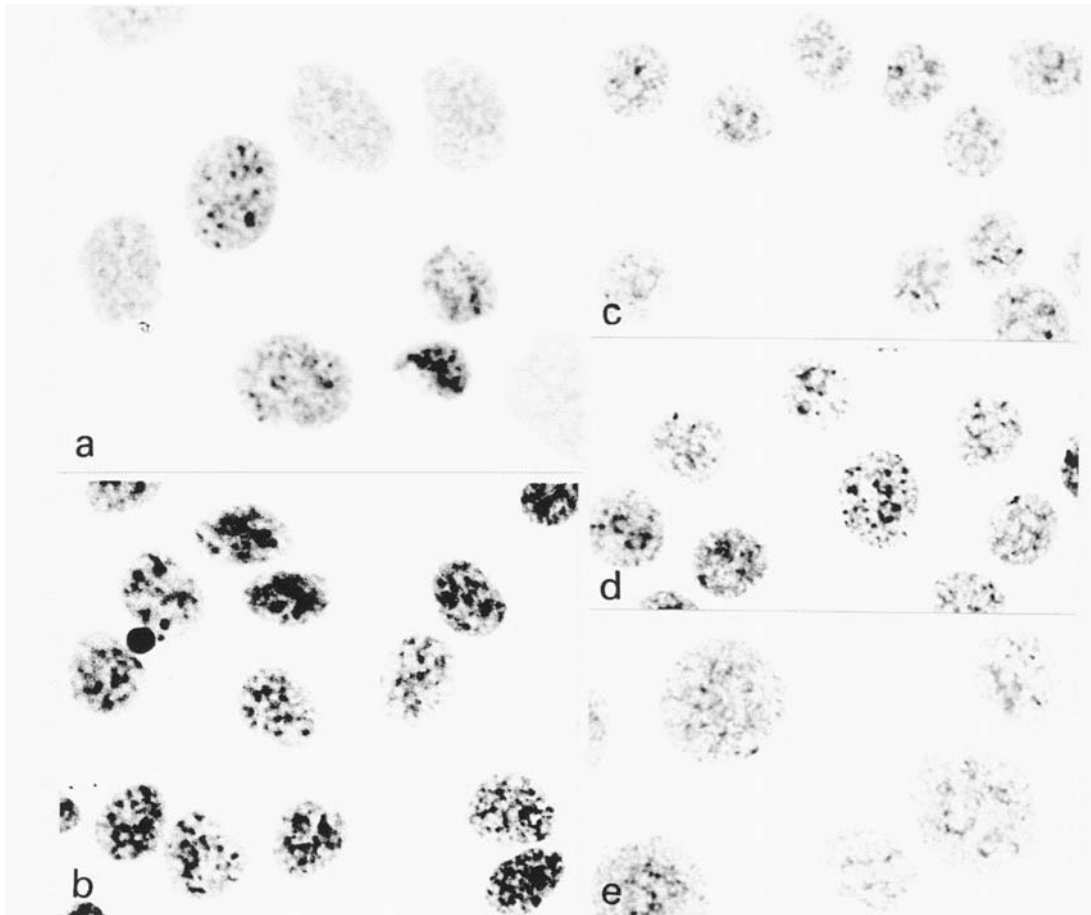


Figure 1. Feulgen-stained nuclei of nontransformed MCF-10F cells (a) and of E2-transformed (b), highly invasive C5 (c) and tumor C5-A6-T6 (d) and C5-A8-T8 cell lines.

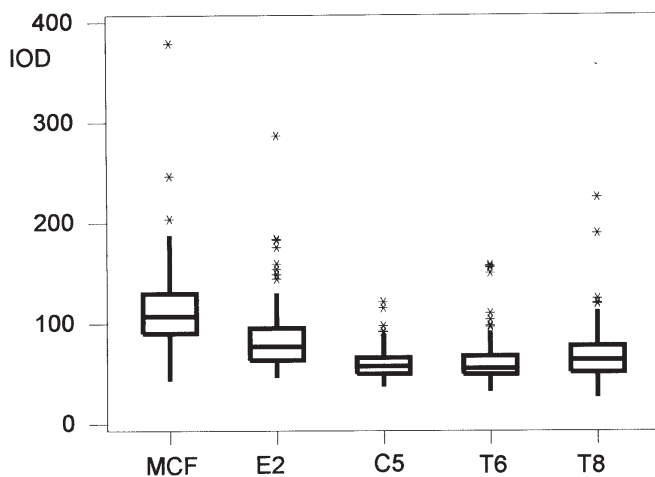


Figure 2. Tukey's box plots showing the distribution of the Feulgen-DNA content of nontransformed MCF-10F cells (MCF) and of E2-transformed (E2), highly invasive C5 and tumor C5-A6-T6 (T6) and C5-A8-T8 (T8) cell lines, n=200.

identical deletions and genomic gains in other chromosomes, as well as the same histopathological and immunocytochemical profile for cytokeratins, E-cadherin, vimentin, and absence of

estrogen receptor  $\alpha$  and of progesterone receptors. Even so, C5-A8-T8 is the most rapidly growing tumor amongst several other tumors induced in SCID mice by the C5 cells. Similar results have been reported for highly aggressive benzo[a]pyrene-transformed MCF-10F cells obtained after transfection with the *c-Ha-ras* oncogene (25). In this case, the Feulgen-DNA content of the MCF-10F cells, which significantly decreases with cell transformation by benzo[a]pyrene (7), increases in the benzo[a]pyrene-transformed cells additionally transfected with the *c-Ha-ras* oncogene, a situation assumed to be induced by a high variability in DNA content inclusively resulting from polyploidy or near-polyploidy (25). The apparent increase in Feulgen-DNA values in the C5-A8-T8 cells compared to C5 cells may thus result from a similar heterogeneity in the cellular population, which is supported by the larger variability in the distribution of the IOD values in the box plot of the former (Fig. 2).

The morphometric changes regarding the total nuclear areas, the Feulgen-DNA absorbing areas, and the nuclear perimeters are generally in agreement with the above-mentioned changes in Feulgen-DNA amounts. Increased perimeters in C5, C5-A6-T6 and C5-A8-T8 cell nuclei compared to values for E2 cells, reflect irregularity in nuclear contours with the tumorigenesis progress. The irregularity in

nuclear contours is especially variable in the tumor cell lines as revealed by the high standard deviations of their nuclear perimeters. The changes in nuclear feret ratio, although not discriminating the tumor cell lines from each other, indicate a trend to spheroidal rather than to ellipsoidal shapes for the nuclear images with the tumorigenesis progress, since the ratio of minimal feret/maximal feret increased.

The significant changes in SDtd, entropy and absorbances in the Feulgen-stained E2-transformed MCF-10F cell preparations are indicative of chromatin structural remodeling with the tumorigenesis process. In E2 cells, the extremely high SDtd and entropy values and an increased standard deviation for the Feulgen-DNA absorbances reflect an increase in variability for the degree of chromatin packing states (1,19,22). The increase in chromatin condensation in certain sectorial nuclear domains concomitant with a decrease in condensation in less packed zones (contrast enhancement) may be involved with the control of expression efficiency in chromatin domains under relaxation (6,7). With selection of the invasive potential of these cells and progression of their invasive to the tumorigenic states, the contrast between condensed and noncondensed chromatin packing states was reduced, becoming even smaller than that in control non-transformed MCF-10F cells. Considering these findings in association with a progressive decrease in Feulgen-DNA absorbances and diminution in the standard deviation for this parameter in the tumor cell lines, it is assumed that a more homogeneous packing state distribution is occurring for the chromatin of these cells, with a certain unravelling of its condensed chromatin. Epigenetic mechanisms involving DNA methylation and/or the histone code, which are known to change the chromatin packing states (26-31) are possibly playing a part in these results. Global genome hypomethylation, regional hypo- and hypermethylation and altered histone modifications have been described as epigenetic dysregulation in cancer cells (31). The chromatin texture evaluated by image analysis has been found to differ in H69 lung carcinoma drug-sensitive and drug-resistant cells under treatment with the histone deacetylase inhibitor trichostatin A (27). Prominent epigenetic alterations have also been associated with increased malignant properties in MDA-MB-231 and MCF-7 human breast cancer cells (31). Additional studies on DNA methylation, histone methylation/acetylation and non-histone protein associations would contribute to a better interpretation of the chromatin textural properties and presumed functions involved in these cells.

### Acknowledgements

This investigation was supported by grants from FAPESP/Brazil (2006/00066-8), CNPq/Brazil, and the U.S. Army Medical and Research Material Command (grants: DAMD17-00-1-0247 and DAMD17-031-0229).

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