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Relation between the rate of tumour cell proliferation and latency time in radiation associated breast cancer

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Abstract

Background: Patients with possible radiation induced cancer could be used to study if the rate of tumour cell proliferation is related to latency time. Such a finding could help researcher to find time periods when other initiating risk factors operate.

Methods: Seventeen women with breast cancer, with a prior history of radiation treatment towards the parts or the whole breast, exclusive of the primary treatment of a breast cancer were identified. Most women had received treatment for benign disorders as hemangiomas, shoulder pain or skin infections. Three patients had been treated with mantle radiation for Hodgkin's disease prior to developing breast cancer. DNA analysis were performed, on remaining tumour tissue after hormone receptor analysis had been done, measuring the fraction of tumour cells in S-phase. Latency time (time between diagnosis and previous radiation treatment) was calculated and related to the S-phase fraction.

Results: A significant inverse relationship between latency time and S-phase was found ($p < 0.0025$), indicating that tumours with a high S-phase had a short latency time and vice versa. Among the possible radiation induced tumours, median S-phase was 14%, comparable with a median latency time of 22 years. Very high S-phase values were associated with short latency times (eg a S-phase of 35% would be compatible with a latency time of 7 years).

Conclusion: Our preliminary results indicate that S-phase is related to latency time and that the median latency time maybe as long as 22 years. Our data may also explain why breast cancer is rare before 30 years of age and if patients are diagnosed at early ages, tumours often show high S-phase values and bad prognostic signs. We postulate that these results from radiation induced breast cancer may be used to extrapolate possible latency times in patients with non radiation induced breast tumours in order to isolate possible time periods for research after other initiating events.

Background

Ionizing irradiation is the only proven initiating factor for

human breast cancer. As such the breast, the thyroid, and the bone marrow appear especially radiosensitive[1]. By

investigating several important cohorts of exposed women, as those surviving atomic bomb irradiation women given radiation therapy for the breast for acute post-partum mastitis or other benign diseases, and women examined repeatedly by fluoroscopy of the chest during treatment for pulmonary tuberculosis with artificial pneumothorax, our knowledge in the field of radiation induced breast cancer has increased [2]. Studies of patients, irradiated for benign conditions or other malignancies than breast cancer may help to describe latency times (time from the first initiating event to diagnosis) in human breast cancer. Studies of breast cancer after exposure to ionizing irradiation points to latency times between 10–30 years [3–5]. In general radiation exposures at very young ages have been associated with longer latency times for breast cancer than exposure at higher ages because radiation induced tumours generally first appear when the normal incidence of tumour is seen [6]. Excess risk has not appeared until 10 years after exposure and rarely before age 30 [2,7–9]. Also the relative risk of diagnosing a tumour seems to further increase in age groups where the natural incidence is high [6]. The risk has foremost and first been shown for women therapeutically irradiated at young ages (childhood, adolescence), while peri- and postmenopausal women irradiated for breast cancer in one breast have not conclusively been found to have a higher risk for contralateral breast cancer because of the radiation treatment [10,11]. This could be due to the fact that the breast epithelium at older ages is less sensitive due to involution and less proliferation.

Cases previously irradiated for benign conditions are becoming less common as the indications for irradiation of non tumour conditions are disappearing. We have previously described a case series in whom women in the past were irradiated for various benign conditions and other malignant diseases (as Hodgkin's disease) [12]. In that previous study, we found that cases with the progesterone receptor positive tumours more often had a history of therapeutic irradiation towards parts of or the whole breast after a pregnancy than before. Women with radiation treatment before the first pregnancy instead more often had progesterone receptor negative tumours.

The aim of the present study was to see if the S-phase fraction (SPF), knowing the age of the woman at diagnosis, could predict the possible latency time and thus give information at what age or age interval the tumour ought to have been initiated. If SPF could be used in such manner epidemiological studies using markers for proliferation in the tumour may assist the epidemiologist in designing studies aimed at identifying initiating factors of breast cancer in a relevant age period of life.

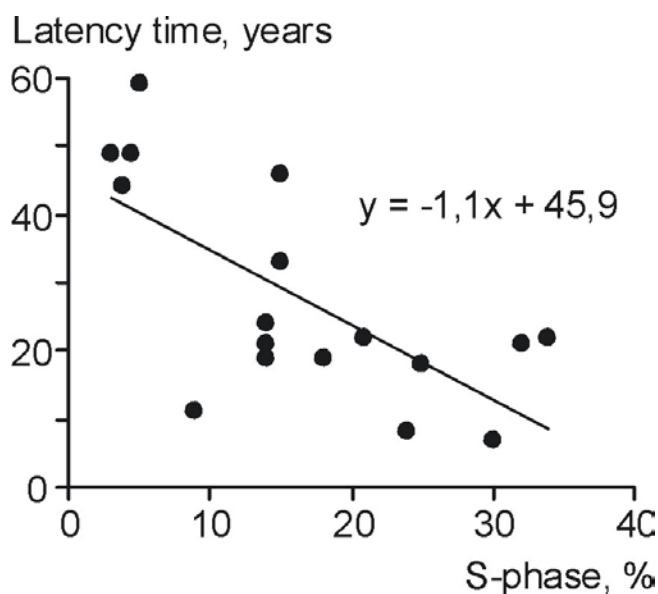


Figure 1

Methods

Seventeen women with breast cancer, with a prior history of radiation treatment towards part of or the whole breast, exclusive of the primary treatment of a breast cancer, were identified. Fourteen women had received treatment for benign disorders as hemangiomas, shoulder pain or skin infections. Three patients had been treated with mantle radiation for Hodgkin's disease prior to developing breast cancer. The radiation dose to the breast tissue varied between 1–40 Gy. All women also had remaining tumour tissue stored at our department after hormone receptor analysis had been done.

Flow cytometric DNA analysis (FCM) was performed according to the following protocol. Tissue was stored at -80°C and a nuclei suspension was prepared with a one-step procedure [13,14] according to current protocol at our lab [15]: 100–200 mg of tissue per tumour sample was thawed in 100–200 μl of citrate buffer (sucrose 250 mM, trisodium citrate 40 mM, dimethyl sulphoxide 5%, pH 7.6) containing trout and chicken red blood cells (106/ml). The tissue was mechanically disintegrated with forceps and thereafter 2 ml of a nuclear isolation medium containing propidium iodide (PI) was added (50 mg PI/ml, SIGMA P-5264; RNase 0.1 mg/ml, SIGMA R-5125; Nonidet P-40 0.6% (v/v; SIGMA N-3516) in isotonic buffered saline; GIBCO). Before incubation in the dark (10 min at room temperature), the sample was filtered (nylon mesh 140 μm) in order to remove tissue remnants from the obtained nuclei suspension. The samples were then

kept at +4 C° until the analysis. An Ortho 50H flow cytometer (Ortho Instruments, Westwood, Massachusetts, USA) was used for the nuclear DNA content analysis. The obtained DNA histogram distributions were judged either as DNA diploid (one cell population) or DNA non-diploid (two or more cell populations). By assuming the cell population to asynchronously synthesizing the DNA, rectangular distribution, the fraction of cells in the S phase (SPF) was estimated planimetrically according to Baisch et al [16]. In case of bimodality in the DNA diploid region reflecting near diploidy (hypo- or hyper diploidy) a mean SPF value was reported for both cell populations together. If DNA index of the non diploid cell population exceeded 1.3 the SPF was reported for this cell population alone. When two or more non diploid cell populations were seen (multiploidy) the SPF for the most prominent cell population was calculated. In those multiploid cases where the cell populations did not appear separated, a mean SPF value was calculated. SPF was not calculated if the histogram showed a debris distribution pattern grossly influencing the S phase region or the fraction of non diploid G0/G1 nuclei was less than 15% of all observed histogram events of the cell cycle. Furthermore, SPF was not reported for those cell populations lacking a visible G2 peak or the CV (coefficient of variation) of the G0/G1 peak exceeded 8 %. Aggregates from nuclei were gated out using the build in DNA analysis software of the Ortho 2140 data handling system.

Latency time (time between diagnosis and previous radiation treatment) was calculated and related to the SPF by regression analysis testing for a linear or a nonlinear relationship.

Results

Median age at radiation exposure was 34 years, range 1–72 yers and median age at breast cancer diagnosis was 64 years range 39–79 years.

The median SPF was 15 (range 3.0–34%). Latency time varied with a median 22 years (range 7–59 years)

A significant inverse relationship between latency time and S-phase was found ($p < 0.0025$), see Figure 1. A linear relation with latency time as the dependent variable and SPF as the independent variable best fitted the values, $Y = 45.89 - 1.09X$, with a $r^2 = 0.47$. An increase of one percent in S-phase decreased latency time on average with one year (95% confidence interval 0.5–1.73 years). A quadratic model gave a slightly better fit, $r^2 = 0.60$, but in view of the small number of observations it was not possible to further explore the functional relationship in more detail and for the same reasons prediction intervals was not presented. Data visualised in Figure 1 suggested that a latency

times less than 8 years was difficult to reach as well as SPF levels below 3%.

No significant relationship was seen between radiation dose and latency time ($p = 0.30$).

Discussion

The time of initiation of an individual breast cancer can only rarely be assessed. The only known exogenous exposure that can be used to possibly date the first initiating event of human breast cancer is ionizing radiation. The practise to treat benign disorders with radiotherapy was prevalent in the past but has become more and more uncommon during the last 30 years. Especially long term side effects as secondary tumors have discouraged the use of radiotherapy for non tumour diagnoses. This also means that the cohort of women, who previously has been exposed to ionising radiation treatment for benign diseases, is an "out dying" cohort and the time period when it is possible to use the cohort for important biological studies is gradually disappearing. In our setting we were fortunate to have a population based breast tumour bank for biological studies that was started 1979 where remaining tumour specimens after hormone receptor analysis have been saved as frozen specimens in -80°C.

Attempts have been made before to calculate latency times in breast cancer using various types of subpopulations as survivors of atomic bombs who later develop breast cancer and populations previously exposed to radiation treatment. The proposed latency time ranges from 10–30 years [1,17–19]. However the increased risk after achieved minimal latency time seems to stay elevated throughout life [20].

By comparing the SPF with the latency time in our study a clear inverse relationship appeared. Our preliminary results indicate that by using the SPF it would be possible in larger materials to predict latency time and that the median latency time maybe as long as 22 years. In men with possible radiation induced breast cancer a possible latency time of 30–40 years have been suggested [4]. Also a minimum latency time of 8 years has been suggested out from radiation exposure studies and breast cancer [5]. In women, irradiated between the time of puberty and the age of 30 for Hodgkin's disease, a median latency time of 15 years have been seen [21,22]. Our results are not contradictory to these previous findings. The data may also explain why breast cancer is rare before 30 years of age as a tumour with a high SPF of 20% still would on average take 25 years to develop. Further data on SPF and latency time also give an explanation for findings of tumour biology in the very young women. For such a tumour to appear with a latency time less than 30 years there is a need to have a SPF above 20% and this tumour necessarily

needs to show bad prognostic signs as a high SPF has been related to poor prognosis [23]. Studies on breast cancer prognosis in different age groups have especially singled out women below age 35 as having a worse prognosis than other women [24–26]. The breast tumours in patients with a prior history of Hodgkin's disease also seem to be more aggressive giving the patients a poorer survival in line with a possibility of higher SPF of their primary tumours because of the rather short latency time [27].

These latency times for radiation induced breast cancer accord very well with time studies of breast cancer development from atypical hyperplasia, through ductal carcinoma in situ and to invasive carcinoma, thought to require 10–20 years or more [28,29].

The median SPF in larger investigations at our laboratory is around 11% [30] compared to the result of the present investigation's 14%. The variation can easily be explained by our small sample size of possibly radiation induced tumours. As the SPF varied between 3–34% a difference in latency time between tumours could be as large as 7–59 years (median 22 years).

A shortcoming in our investigation is that while epidemiological data support that the group of patients may have radiation treatment as a risk factor it is impossible in the individual case to for certain know if radiation caused the tumour. We speculate that if we could accurately know which tumours that were radiation induced, the fit of the model would have been near perfect using the SPF.

We can not fully exclude that the model e.g. is quadratic rather than linear. More cases with biological data would be needed to fully describe the true relationship, but it is reasonable to believe that a minimal latency time of 5–10 years is required for a breast cancer to be able to develop. The reproducibility of analysing the SPF is in our hands very high [31,32] and there is a high correlation with other measures of proliferation [33]. Further SPF calculations give very good prognostic information [23,30]. However there are sometimes potential problems as in patients with low SPF and multiple tumour populations calculations can be contaminated with normal tissue from tumour stroma or inaccuracies can occur in assessing the SPF because of overlapping tumour cell populations [30].

Other potential problems by our approach of relating the SPF to latency time is that cell loss/cell death in individual tumours could not be estimated or accounted for and may vary between tumours. Further it is not fully known from literature how human tumours grow eg if there is a continuous linear growth, exponential growths according to Gompertz or periods of interrupted exponential growths [34]. Further in experimental tumours there is a sugges-

tion that a higher radiation dose may be related to a shorter latency time. However, this has been proposed to be due to a higher statistical probability for transformation at higher doses [35]. In the present study we did not see a statistical significant relationship between radiation dose and latency time, but the small sample size could have prevented us from finding a significant association. In general it has also been stated that radiation induced tumours in man tend to present at common ages for the incidence of a given tumour and thus be related to longer latency times if the radiation exposure took place at younger ages [36].

Most breast cancers are thought to start its development early in life at puberty or even in early childhood or in utero [37]. Previously, it has been proposed from one of the authors that the tumour at diagnosis, at least partially, retains characteristics of the normal breast epithelium at the time of initiation and that varying tumour biology/prognosis at different ages may be explained out from this [38,39]. Experience from atom bomb survivors suggest that reproductive factors and hormone use appear to act independently of radiation exposure on the risk of breast cancer among this population [40].

Conclusion

Results from radiation induced breast cancer may be used to extrapolate possible latency times in patients with non radiation induced breast tumours. Out from a woman's age at diagnosis and SPF, a possible latency time, using our linear formula, possibly can be predicted and initiating epidemiological risk factors can be sought in the responsible time period for initiation. This may enable researcher to more efficiently find new initiating risk factors of breast cancer. It is conceivable that other measures of tumour proliferation rate as mitotic index, labelling index, antibodies as Ki-67, or BrdU labelling may be used in a similar manner and this may allow studies on larger patient materials, where paraffin embedded tumour materials have been saved. These different measures of proliferation (such as S-phase, labelling index, mitotic index, Ki-67) have been found to correlate well with each other in breast cancer [41]. To calibrate curves between tumour proliferation rate and latency time, tumours from possibly radiation induced cases must be identified and used. Therefore radiation induced cases constitute a very important "out-dying" research source.

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