CLINICAL INVESTIGATION

PARENCHYMAL CELL PROLIFERATION IN CORONARY ARTERIES AFTER PERCUTANEOUS TRANSLUMINAL CORONARY ANGIOPLASTY: A HUMAN TISSUE BANK STUDY

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Purpose: Restenosis after percutaneous transluminal coronary angioplasty (PTCA) remains a limitation of this technique. Arterial wall cell proliferation is a component of restenosis preventable with intravascular brachytherapy. This study attempts to locate the sites of cellular proliferation after PTCA so as to aid the optimization of this therapy.

Methods and Materials: Autopsy records from January 1, 1985 through December 31, 1995 were reviewed, and 27 patients who received PTCA prior to death were identified who also had evidence of PTCA on histologic examination of the arterial sections. The sections were subjected to immunohistochemical staining for proliferating cell nuclear antigen (PCNA) to detect the proliferating cells in the arterial sections, followed by image analysis to determine the proliferative index (PI) of all regions and layers of the section.

Results: The PI did not differ significantly according to vessel region (plaque, plaque shoulder, or portion of vessel wall with lowest plaque burden), vessel layer (intima, media, adventitia), or evidence of prior PTCA. There was a trend toward a higher PI in young lesions.

Conclusion: Cell proliferation in the vascular wall after PTCA was found throughout the treated arterial section’s axial plane, not only in the periluminal region. © 1999 Elsevier Science Inc.

Restenosis, Angioplasty, Brachytherapy, Immunohistochemistry, Autopsy.

INTRODUCTION

Intravascular brachytherapy as a preventative treatment for restenosis after transcatheter vascular intervention is gaining currency (1, 2). The success of this treatment has prompted its empiric application using several devices. These techniques are in the introductory phase, and further advances are likely to be characterized by better coverage of the treatment target(s). The honing of a brachytherapy procedure to precisely treat the perceived treatment target, as has been done in the treatment of early-stage prostate cancer, leads to improved efficacy and decreased toxicity (3, 4).

It is reasonable to believe that similar results will be achieved with further development of intravascular brachytherapy techniques. The major hindrance to these advances is the definition of the treatment target(s).

The study of the cause(s) of restenosis is an area of evolving consensus. It is clear that cellular proliferation is at least partly responsible (5). This is an important recognition, because ionizing radiation is well known to inhibit cellular proliferation in the clinical setting of keloid prophylaxis (6). This has led to the application of radiation in animal models of restenosis with promising results for restenosis prevention (7, 8). These preclinical studies have prompted clinical trials that also report encouraging results (1, 2, 9). Optimization of this technique may lead to even better results with new devices that are attuned to the biology and anatomy of the vascular lesions treated with PTCA. To this end, it seems that targeting the radiation to the proliferating cells should be a reasonable strategy. This report describes the location and intensity of the proliferation of parenchymal cells within the vascular wall of patients who have had PTCA prior to death, so as to aid such optimization.

Immunohistochemical staining for proliferating cell nuclear antigen (PCNA) is the method chosen for identifying the proliferating parenchymal cells. Traditional methods of
visual evaluation are labor-intensive and prone to investigator bias (10). Computer-based image analysis utilizes consistent criteria to evaluate samples and is therefore reproducible, objective, and reliable, making it an increasingly important option in the evaluation of histological samples (11). Successful image analysis methods for determining proliferation indices in various types of tissue sections have been reported in the literature (10, 12–14). We employ similar methods to determine the endpoints of this investigation using computer-generated proliferation indices.

**METHODS AND MATERIALS**

**Case acquisition**

The autopsy records from January 1, 1985 to December 30, 1995 from The Cleveland Clinic Foundation were reviewed. Sixty-four patients with a history of PTCA were identified, and the microscope slides of the treated coronary artery sections were retrieved from the archives. All slides were reviewed for evidence of the PTCA, for example, vascular trauma (15) and 27 were culled for evaluation because they displayed the stigmata of PTCA. Re-cuts of the 27 arterial sections were subjected to immunohistochemical staining and image analysis.

**Immunohistochemical staining**

PCNA is a 36-kD protein that is highly conserved between species. PCNA functions as a cofactor for DNA polymerase delta in both S phase and in DNA synthesis associated with DNA repair mechanisms (16). The PCNA molecule has a half-life in excess of 20 h and may be detected in non-cycling cells such as those in G₀ phase or those fixed within one day of death. PCNA was therefore chosen to detect proliferating cells in these autopsy specimens because they were fixed within one day of death, and the resting cell division rate in human arteries is longer than five half-lives of PCNA; therefore, the resting proliferative rate will not affect the analysis.

Sections of artery were fixed in formalin, embedded in paraffin, and cut at 2 µm. Slides were heated in a 60°C oven overnight, deparaffinized through xylene and graded alcohols, and quenched in 0.5% aqueous hydrogen peroxide. Sections were microwaved (850 W) for 10 min in 0.1 M citric acid buffer (pH 6), cooled, and washed in phosphate buffered saline (PBS, pH 7.4). Incubating with 1.5% normal goat serum (Vector Laboratories, Inc., Burlingame, CA) in PBS for 30 min at room temperature blocked nonspecific sites. Monoclonal (PC 10) mouse anti-human Proliferating Cell Nuclear Antigen (Dako, Carpenteria, CA) was applied at a dilution of 1:200 in diluted normal goat serum for 1 h at room temperature. The primary antibody was detected by the VECTASTAIN-ABC kit for mouse IgG (Vector Laboratories), which consists of a biotinylated secondary antibody and an avidin-biotin enzyme complex. Each of these reagents, prepared according to the manufacturer, was applied sequentially to sections for 30 min at room temperature, washing in between with PBS. After a final wash, tissues were incubated with 3,3-diaminobenzidine tetrahydrochloride in hydrogen peroxide (Dako Liquid DAB) for 2 min, producing a reddish-brown precipitate. As controls, commercially prepared slides of human tonsil (Dako) were incubated with primary antibody and separately with PBS, and developed with the Vectastain Kit, under conditions identical to those employed with the sample sections.

**Apparatus and images**

For classification purposes, each arterial cross-section was subdivided into three regions corresponding to basic plaque morphology. As shown schematically in Fig. 1, the thickest region of the arterial wall (indicating the greatest area of neointima and/or plaque deposits) was designated as region 3, the thinnest region of the arterial wall was designated as region 1, and the regions of initial thickening separating regions 1 and 3 were designated as region 2.

Tissue sections were viewed in cross-section on a Nikon TMS light microscope. The images were captured using a DC-330 3CCD color video camera (Dage-MTI, Michigan City, IN) mounted on the microscope and a Flashpoint 128 video capture card (Integral Technologies, Indianapolis, IN). Four images each from the intima, media, and adventitia of each designated region were acquired, resulting in a total of 36 images for each tissue section. Thus, approximately 1.6 mm² of each arterial cross-section was captured for analysis. The color images were stored as uncompressed TIFF (tagged image file format) files in the RGB (red, green, blue) format.
Initial image processing and rationale for using the HSI system

Images were processed and analyzed using the Image-Pro Plus image analysis program (Media Cybernetics, Silver Spring, MD) on a Pentium II personal computer. The original images captured by the video camera (an example is shown in Fig. 2A) were prepared for analysis. Next, the region of interest in each image was isolated, insuring that each image contained only a single designated region and a single arterial area (as shown in Fig. 2B).

The images were then transformed from the RGB color space to the HSI (hue, saturation, intensity) color space for ease of analysis. In RGB color space, the separation of colors is dependent on the intensity; therefore, different colors may cluster closely together, making histogram-based segmentation difficult. In HSI color space, the brown and blue hues of the immunohistochemical staining were clearly separable in a 1-dimensional histogram, ultimately allowing the assessment of consistent proliferation indices of the arterial sections as explained below.

Image analysis

To establish the proliferation index of an artery section, the total number of nuclei and the number of positive-
labeling nuclei must be obtained. Both of these goals can be achieved by histogram-based clustering (thresholding), a widely used technique in image analysis that segments an image into relevant and irrelevant objects based on a pixel-wise classification scheme (17–19). We adapted the thresholding method described by Weaver and Au for human solid tumors to our present analysis of proliferation in artery sections (18). In brief, the method entails utilizing a threshold based on intensity information to establish the total number of nuclei. A second threshold makes use of the hue information to distinguish the positive nuclei from the negative nuclei.

Establishing the total number of nuclei

Interactive thresholding of the intensity spectrum of the HSI provided the required separation of the darker stained nuclei from the cytoplasmic background and cytoplasmic debris (see Fig. 2C). Spurious items such as non-nuclear structures or cellular debris were removed by masking the image with a size range corresponding to the expected nuclei. With both the intensity threshold and the size range applied to the image, a nuclear count was then performed. The result of these steps is demonstrated in Fig. 2D.

Determining the positive nuclei

Positive nuclei were indicated by the brown immunohistochemical staining (the result of the DAB chromagen recognition of the PCNA), while negative nuclei were distinguished by the blue hematoxylin counterstain. This distinction thus allowed us to utilize hue values to separate the positive nuclei from the negative nuclei. Hues representing positive nuclei are located in two clusters at the two ends of the hue spectrum and are separated by the blue hues of the negative nuclei (see Fig. 2E). The result of selecting the brown hue values is shown in Fig. 2F. Because the actual hue of the DAB chromagen was expected to remain relatively constant within the sample group (unlike the intensity of the staining which could be easily affected by variations in the staining environment), the same hue range was applied for all images in the study. The proliferative index (PI) was calculated as the number of positive labeling nuclei/total number of nuclei; this was then multiplied by 100.

Statistical considerations

Two-factor repeated measures analysis was used to determine if the PI differed among layers or regions. The data were initially tested to determine if there was significant interaction between layers and regions. If interaction is not significant, then the PI can be compared among the three regions averaged across layers, or among the three layers averaged across regions. One-way analysis of variance with a repeated measures design was used to determine if the PI differed between lesions that underwent prior intervention and those which did not. Spearman’s rank correlation coefficient was calculated to measure the association between PI and lesion age, separately for each region and layer. The correlation was tested to determine if it differed significantly from zero. All statistical tests were two-sided, and \( p < 0.05 \) was used to indicate statistical significance.

RESULTS

The median age (measured in days from time of the most recent intervention to time of death) of the lesions tested is 2 days (mean = 104.9, SD = 413.6, range of 0.1–2162). The majority (19 of 27) is less than 20 days old. Eight of 27 had at least one prior PTCA to the tested site. All sections were suitable for breakdown into the schematic seen in Fig. 1.

The primary endpoint of this analysis is locating the proliferating cells in the arterial wall. The PI did not differ between arterial regions 1, 2, and 3, arterial layers, or according to having a prior intervention (Table 1). Although the \( p \)-values are not significant, a trend was seen indicating the adventitia as a more active site of proliferation.

A statistically significant relationship of the lesion age with PI was not identified. However, it is noted that the sections in which PI was maximal are the younger sections of the series (Fig. 3).
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DISCUSSION

Significance and clinical application

These analyses demonstrate the ubiquity of arterial wall cellular proliferation after de novo or repeat PTCA in human tissue bank coronary artery specimens. Because the most easily obtained images of the coronary arteries are angiograms, much attention has been devoted to that which is imaged by an angiogram: the lumen. This investigation is concentrated on the arterial wall, the tissue that forms, or in atherosclerotic disease states, “permits” the lumen. After a transcatheter intervention to open a stenosed lumen, such as PTCA, the vessel wall again is central to the long-term success of the procedure. The restenosis process will variably affect the lumenal patency through a combination of arterial wall cell proliferation and vessel resizing or remodeling (the interplay between the two is not clearly understood; however, there is reason to believe that negative remodeling may also be a result of cell proliferation).

This study parallels previous work in the porcine model of restenosis (20). In that porcine model, parenchymal cell proliferation (as detected by 5-bromo-2-deoxyuridine labeling) was noted in nearly all sites in the balloon-injured normal coronary artery after balloon overexpansion injury. The major difference between that and the present study is that in the present study, the specimens were obtained from humans with atherosclerotic coronary artery disease that was severe enough to warrant PTCA. Therefore, the patients likely to be eligible for intravascular brachytherapy in the future will be very similar to the present group. In this group, there will be post-PTCA cell proliferation in the periluminal portions of the vessel as well as the adventitia behind the plaque.

The location of the post-PTCA cellular proliferation takes on new importance as intravascular brachytherapy evolves and failure analyses of clinical trials are examined. For example, a single-institution randomized trial of intravascular brachytherapy for in-stent coronary restenosis was completed at Scripps Clinic (1). An important follow-up study to the original publication is the failure analysis in which it was noted that success was heavily dependent on achieving a dose of 8 Gy to the adventitial border (21). The most frequent reason for not delivering at least 8 Gy to the entire adventitial border was eccentricity. With the source placed in the lumen of an eccentric lesion, it was not possible to deliver 8 Gy to the far side of the adventitia without exceeding 30 Gy (the maximum dose the trial would allow) to the near side. From this data, one may assume that there is important target tissue at the far side of the adventitia requiring adequate dosing. The proliferating cells highlighted in the present study have a high likelihood of being that target tissue, as they are present both near the lumen and at the dosimetrically challenging contraluminal locations. The patients in the Scripps Trial with highly eccentric lesions would have received less than 8 Gy to the far side of the adventitia and the proliferating cells it contains.

The work presented here suggests that future intravascular brachytherapy devices and techniques should be designed to correspond to the local vascular anatomy by accounting for the eccentric nature of the diseased artery. In such devices, vascular wall imaging, such as intravascular ultrasound (IVUS), may be valuable because of its ability to image the eccentricity and potentially direct therapy accordingly. In addition, the choice of isotope should be driven by the need to deliver therapeutic doses through tissue that may be 3–4 mm from the luminal border.

Limitations

The difference between the mean and median age of the lesions is vast so it is not surprising that a correlation between lesion age and cellular proliferation was not seen, as it was by others (20, 22). Another possible explanation for the lack of a temporal correlation is that the sections from the different patients were very likely fixed at different times (no record of this exists) after death; therefore, the half-life kinetics of PCNA becomes a confounding variable that cannot be controlled statistically. Conclusions must therefore be based on gross descriptions such as defining the sites of proliferation, rather than attempting to describe temporal phenomena. It is encouraging to see that in Fig. 3 the maximal proliferation was confined to the younger lesions in the series, a finding that parallels earlier data (20, 22). Also, it is heartening to see the trend toward more active cell proliferation in the adventitia (p = 0.11), which corresponds to data from the porcine model of restenosis (20).

CONCLUSION

In summary, arterial parenchymal cell proliferation is found in all arterial layers and regions after PTCA. Intravascular brachytherapy should account for this anatomical finding and deliver radiation according to vascular wall anatomy rather than luminal anatomy.
REFERENCES


