Structural Patterns of Immunoreactivity to Cytokeratin 18 in Normal Prostate and Benign Prostate Hyperplasia: Global and Local Differences

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Abstract

Background: Benign prostate hyperplasia (BPH) is the most common benign disease of the human prostate. The comparison between global versus local changes in spatial patterns of pathological lesions provoked a growing interest in some fields such as neuropathology. To date, there is little data on this subject in prostatic pathology. Given the interest of local parameters to distinguish between normal and pathological structures, the present study will apply first and second order stereological tools to find out if the cytokeratin18 (ck18) immunoexpression shows relevant local changes in BPH compared to normal prostate, independently if global estimates were similar in both groups.

Methods: To verify if the global and local changes in immunoexpression of ck18 are relevant to ascertain differences between normal (CTR) and BPH cases, the following parameters will be applied: Volume fraction of epithelium immunostained for ck18 (V,ck18), both in global and local estimates; dispersion indices of V,ck18; estimates of local variance of V,ck18 (positional and of scale) using wavelet analysis; and lacunarity analysis to measure the tissue heterogeneity. Then, the set of values from the parameters studied that show significant differences between normal (CTR) and BPH cases, the following parameters will be applied: Volume fraction of epithelium immunostained for ck18 (V,ck18), both in global and local estimates; dispersion indices of V,ck18; estimates of local variance of V,ck18 (positional and of scale) using wavelet analysis; and lacunarity analysis to measure the tissue heterogeneity. Then, the set of values from the parameters studied that show significant differences between CTR and BPH will be employed to perform stepwise linear discriminant analyses to determine if locally estimated parameters were able to classify accurately the cases in CTR and BPH groups.

Results and Conclusions: The findings of the present study indicate that changes in the expression of ck18 by the hyperplastic prostatic epithelium are not homogeneous. This limits the use of a single biopsy based markers to predict biological behavior in BPH. On the other hand, the local changes in the expression of ck18 are more evident in terms of V,ck18 and its

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local variability, whereas other parameters that are useful in other pathologies, such as lacunarity, are less relevant in prostatic hyperplasia.

**Keywords**

Prostate, Hyperplasia, Cytokeratin 18, Volume Fraction, Lacunarity, Local Changes

### 1. Introduction

Benign prostate hyperplasia (BPH) is the most common benign disease of human prostate [1] [2], representing approximately 50% of medical consultations made by urological disease [3] [4].

Although the pathogenesis of BPH is unclear, it is known to be multifactorial; being necessary the presence of two factors for prostate growth occurs: the androgen stimulus and age [5]. Different theories have been proposed, based on histological, and hormonal age-related changes, but, currently, not a single explanation is accepted [6]. Androgens possibly act as initiators stromal hyperplasia, which in turn induce epithelial hyperplasia [7] [8]. It has been also suggested that neuroendocrine cells are involved in the genesis of BPH, perhaps enacting some mediation between stroma and epithelium. The cytological findings in BPH are generally of little relevance, and rather unspecific, as alterations including basal cell hyperplasia, increased stromal mass (particularly the amount of smooth muscle cells), enhanced extracellular matrix deposition, reduced elastic tissue, more infiltrating lymphocytes around ducts, acinar hypertrophy and more luminal corpora amylacea and calcifications in the form of prostatic calculi [9].

The cytokeratin class of intermediate filaments has been shown to exist in all epithelia [10]. There are now recognized 19 distinct cytokeratins expressed in human epithelia and each epithelial type has a distinct phenotype with regard to these proteins [11] [12]. Early studies using polyclonal anticytokeratins demonstrated that the basal cells of the prostate have cytokeratins that are immunoreactively different from the luminal or columnar cells [13] [14]. Several studies using monoclonal anti-cytokeratin antibodies have differentiated the columnar and basal cell populations on the basis of their specific cytokeratin content [15]. The columnar cells react with monoclonal antibodies to cytokeratin 18 [16]. Besides, chemical determination of the cytokeratin phenotype of three established human prostatic carcinoma cell lines also suggests that all three cell lines synthesize cytokeratin 18 (ck18) [16]. In the nonmalignant (BPH) prostatic secretory acini, keratin expression patterns have been examined using monoclonal antibodies for several types of cytokeratins [17]. The keratins recognized are all expressed in the glandular cells, for example, cytokeratin 14 specifically stain the basal cell population, while cytokeratins 18 and 19 react with the columnar cell population.

When comparing the immunoexpression of ck18 in BPH with normal pros-
tate, no remarkable differences are detected [16] [17]. Nevertheless, the percentage of positivity for luminal ck18 was statistically lower for BPH cultures respect to the positivity observed for both prostatic intraepithelial neoplasia (PIN) and prostate cancer (PCa)-derived cultures [18].

All these studies were based on global estimates of immunoexpression of ck18 in malignant of benign prostate cases, but there is lack of information about possible local differences in immunoexpression of ck18, within both normal and pathological prostate samples.

The comparison between global versus local changes in spatial patterns of pathological lesions provoked a growing interest in some fields such as neuropathology [19]. However, there is little data on this subject in the field of prostatic pathology. Recently, several studies have dealt with the estimation of measurements of either acinar or stromal parameters at the local level in normal and pathological human prostate, using second order stereological methods [20]. For example: The study of the distribution of microvessels in normal and pathological prostate [21], the distribution of cell nuclei populations in prostatic adenocarcinoma and PIN [22], the estimation of K function and isotropy of normal prostatic acini compared to cancer [23], and the local changes of glandular pattern in BPH compared to normal prostate [24]. In some of these studies, several discrepancies between global and local results were detected. Thus, the first order (global) parameter of volume fraction of epithelial ck18 does not rapport differential information between normal and cancer prostate acini [23], or in the study of the structural pattern of acinar tree, the global measurements as the average volume of acini were unable to distinguish between normal prostate and BPH; whereas local parameters, such as connectivity density, showed remarkable differences between normal and pathological prostate [24].

Given the interest of local parameters to distinguish between normal and pathological structures, the present study will apply first and second order stereological tools to find out if the ck18 immunoexpression shows relevant local changes in BPH compared to normal prostate, independently if global estimates were similar in both groups.

To check for such local changes in the ck18 immunoexpression in both normal and hyperplastic (BPH) prostate, the following parameters will be applied:

1: Volume fraction of epithelium immunostained for ck18 (V₂ ck18), both in global and local estimates (pixel to pixel of tissue).
2: Dispersion indices of global V₂ ck18, as Morisita index [25] [26].
3: Estimates of the local variance of V₂ ck18 (positional and of scale) using wavelet analysis [27] [28] [29] [30].
4: Lacunarity analysis in order to measure the distribution of gap sizes interspersed among the immunostained acini (i.e.: tissue heterogeneity) [31].

2. Material and Methods

2.1. Material

Twenty prostate specimens were collected over a period of one year (2015-16), at
La Princesa Hospital (Madrid, Spain), 10 were from adults, (CTR group), age (mean ± SD): 45 ± 7; range: 30 - 47 years, all these specimens were of healthy subjects, without endocrine or reproductive pathology, deceased in traffic accidents, and eligible as donors for transplant, the age of the subjects of CTR group was in the range indicated to avoid any histological changes of subclinical BPH, relatively frequent in subjects older than 50 years. The other 10 were surgical specimens (adenomectomies) from patients diagnosed of the adenofibromiomaticous type of benign prostatic hyperplasia, (BPH group), age (mean ± SD): 75 ± 10, range: 65 - 85 years. All the ethical requirements were accomplished in order to obtain the prostatic tissue either at the moment of the multiorganic extraction for transplant (CTR group) or at the surgery (BPH group).

2.2. Processing of the Tissues

Immediately after extraction, the specimens were fixed during a week in 10% paraformaldehyde in PBS, pH 7.4, (Probus, Barcelona, Spain).

After fixation, the specimens from the two groups were thoroughly sectioned into 2-mm-thick slices, performed by isotropic uniform random sampling (IUR sections) in order to preserve the isotropy of the tissue [32].

All the specimens were processed for paraffin embedding. The paraffin blocks were exhaustively sectioned. A total of 20 sections (5-µm-thick) were performed on each block for immunohistochemistry.

2.3. Immunohistochemistry

At least five randomly selected slides per specimen were immunostained for ck18 in CTR and BPH groups. Deparaffinized and rehydrated tissue sections were treated at room temperature for 30 min with hydrogen peroxide 0.3% in phosphate-buffered saline (PBS) pH 7.4, to block endogenous peroxidase. To detect ck18 immunoreactivity, sections were incubated with a monoclonal anticytokeratin 18 antibody (Abcam, Cambridge, UK) diluted at 1:250. Pretreatment of sections by heat in citrate buffer pH 6.0 (using a pressure cooker) [33] was performed to enhance immunostaining.

The primary antiserum was diluted in PBS pH 7.4 containing 1% bovine serum albumin (BSA) (Sigma, St Louis, USA) plus 0.1% sodium azide (Merck, Darmstadt, Germany). The incubation with primary antiserum was overnight at 4°C. The second antibody employed was a biotin-caproyl-anti-rabbit immunoglobulin (Biomed, Foster City, CA, USA). The second antibody was diluted at 1/400 in PBS containing 1% BSA without sodium azide, and incubated for 30 min at room temperature. Thereafter, sections were incubated with a streptavidin-biotin-peroxidase complex (Biomed). The immunostaining reaction product was developed using 0.1 g diaminobenzidine (DAB) (Sigma) in 200 mL of PBS, plus 40 µL hydrogen peroxide. After immunoreaction nuclear counterstaining with Harris haematoxylin was performed in some sections immunostained for ck18. No nuclear counter-staining was performed on the remaining sections that were then employed for quantitative purposes. All slides were de-
hydrated in ethanol, and mounted in a synthetic resin (Depex, Serva, Heidelberg, Germany). The specificity of the immunohistochemical procedures was checked by incubation of sections with no immune serum instead of the primary antibody.

2.4. Data Acquisition

Five strips of an average of 20 immediately adjacent quadrats (range 10-40) were explored for each immunostained section from CTR and BPH groups. The origin and sense of the axis for each strip were chosen by systematic random sampling [34] for all the strips. The result was a series of images from the two groups, sized, on average, 512 × 7000 pixels. The final magnification (×100) was such that 1000 pixels represented 1280 μm. At that point, the strips were 9 mm long, on average. Therefore, the total length explored per section (five sections) and per case (10 cases) was 9 × 5 × 10 = 450 mm (for BPH cases, an appreciable percentage of the maximum specimen diameter) [21].

The images were captured using a colour digital camera DP 70 (Olympus Corporation of the Americas, PA, USA) with a resolution of 12.5 mega-pixels, attached to an Olympus microscope fitted with a motorised stage controlled by the stereological software Cast-Grid (Stereology Software Package, Silkeborg, Denmark). This program controls the XY displacement of the microscope stage and allows the selection of fields to be studied by random systematic sampling after the input of an appropriate sampling fraction [22].

The strips were then mounted from the images captured, using the public domain Java image processing program, Image J (version 1.48), developed at the US National Institutes of Health and available on the Internet at https://imagej.nih.gov/ij/index.html [35]. Subsequently, the resultant strips were processed using the same software.

A binary image was produced where the immunostaining to ck18 was shown as black and the pore space (lumina of acini, stroma, etc.) as white (Figure 1).

2.5. Quantitative Measurements

-Volume fractions of immunostained epithelium

The local measurements of the fraction of volume of tissue immunostained to ck18 (\(V_v\) ck18) were obtained as a follows: At each point of the long axis of each image strip, the fraction of pixels belonging to the immunostained epithelium, expressed as a percentage over the space of reference (pore space plus immuno-reactive epithelial component) was automatically recorded by the image analysis system for all the N columns orthogonal to the long axis. The resulting series of N consecutive rational numbers per visual field served as input signals for estimating the \(V_v\) ck18 measurements (per pixel of tissue) [21]. The results were plotted as a space series, being the position (number of pixels transformed on microns) represented in the X-axis and the \(V_v\) ck18 in the Y-axis of the plot.

The global measurement of \(V_v\) ck18 was obtained averaging the local \(V_v\) ck18 over the total number of strips for each case in both CTR and BPH groups.
Figure 1. In (a) the image shown is a strip from a specimen of CTR group immunostained to ck18; In (b) the binarised picture from (a) is depicted, the ck18 immunoreactive cytoplasm is in black and the space of reference (acinar lumina and stroma) in white; In (c) the image shown is a strip from a specimen of BPH group immunostained to ck18; In (d) the binarised picture from (c) is depicted, the ck18 immunoreactive cytoplasm is in black and the space of reference (acinar lumina and stroma) in white. The scale bars represent 326 µm.

- Dispersion index of the global $V_v$ ck18 measurements

This procedure takes estimates of $V_v$ ck18 from quadrats and calculates several indices that can be used to identify the spatial patterning of the volume fraction. In the present study the Morisita index was obtained [25]. It is the scaled probability that two measurements chosen at random from the whole population are in the same quadrat. The higher the value, the more clumped the distribution. This parameter was obtained using the PASSaGE software [26], that is a program suitable for pattern analysis and spatial statistics.

- Estimation of local variability of $V_v$ ck18

The local variance for $V_v$ ck18, related to position and scale, was measured in both CTR and BPH groups. To do this, a wavelet analysis was employed. The quadrat variance methods calculate the variance of differences among blocks of data of different sizes or scales and use the pattern of the variance estimates to
determine the scale of pattern [36]. Wavelet analysis is similar to many of the quadrat variance methods; although in some ways it is much more flexible. Wavelets have been heavily studied in mathematics and engineering for signal analysis and data compression, but have had limited use in biology [27] [28] [29] [30].

A wavelet function is a scalable windowing function whose integral equals zero. One way to think of this is that the wavelet function describes a template that can be scaled to a desired size, and then slid onto the space series of \( V_\gamma \) ck18 values along of long axis of the strip. When the template fits the observed data well, the value of the wavelet transform at that position is high; when it does not, the value is low. Then, the adjusting of wavelet function over the space series, obtain the overall variance at a given position and scale [26]. In the present study, wavelet analysis was performed from one to a maximum scale specified as a percentage of the input data size (50%). The wavelet kernel employed was the Haar wavelet function [37]. Wavelet analysis of variance was performed using the PASSaGE software. The results for both CTR and BPH groups were expressed plotting the variance values in function of position (in microns) and scale.

- Estimation of local heterogeneity (clumping) of \( V_\gamma \) ck18

The heterogeneity of distribution for the acini immunoreactive to ck18 (in CTR and BPH cases) was detected by estimation of the patterning of gap sizes among the immunostained structures; this was performed by means of the lacunarity analysis.

Lacunarity analysis derives from fractal mathematics and is a measure of the distribution of gap sizes [38]. Originally developed for binary (presence/absence) data, lacunarity analysis is easily applicable to continuously distributed variables as well [31]. Lacunarity (\( \Lambda \)) for a specific range (scale) is estimated as the ratio of the first (mean) and second (variance) moments of measurements within all possible boxes of that range width. Lacunarity analysis was performed for all scales from one to a maximum specified as a percentage of the input data size. The maximum allowable scale was 50%.

The results were plotted on a log-log plot of \( \Lambda \) versus scale. Random data produces a curve, which is concave upwards. Clumped data produces greater lacunarity and a curve which is concave downwards. Regularly spaced data produces less lacunarity and a curve which is initially straight [39] [38] [31].

Randomization tests were performed to check whether the behavior of the observed data was at random or followed a particular pattern. Randomization tests for lacunarity analysis work by randomizing the order of the quadrats within the space series and recalculating the lacunarity profile for the randomized data. This generates a null distribution of expected lacunarity indices for data with the specific observed values but with no specific relationships. Significant lacunarity indices can be identified from the observed data when the observed values fall outside the expectation generated from the randomization test.

All these calculations described were performed using the PASSaGE software.
2.6. Statistical Analysis

The global $V_V$ ck18 and Morisita index were expressed as mean ± SEM. Comparisons between these means from CTR and BPH groups were performed by a Student t test. For $V_V$ ck18 values along the space series, positional and scale variance of $V_V$ ck18, and lacunarity measurements, the local estimates of mean ± SEM were performed and also compared between CTR and BPH groups by a Student t test. The level of significance was $p < 0.05$. For $V_V$ ck18 values along the space series, positional and scale variance of $V_V$ ck18, and lacunarity measurements, the set of values showing significant differences between CTR and BPH was pooled and employed to perform stepwise linear discriminant analyses [40] in order to determine if locally estimated parameters were able to classify accurately the cases in CTR and BPH groups.

Discriminant variables were selected according to Wilk’s lambda: at each step, the variable that minimises the overall Wilk’s lambda or maximises the associated F statistic is selected (F to enter = 3.84 and F to remove = 2.71). Wilk’s lambda statistic explains the rate of total variability that is not due to differences among groups. A lambda of 1 means that the mean of the discriminant scores is the same in all groups and there is no variability between groups, while a lambda near 0 means that there is a significant difference among groups. Therefore, Wilk’s lambda provides a test of the null hypothesis that the population means are equal. The larger lambda is the less discriminating power is present [41].

For each local variable, the discriminant scores obtained were employed to construct histograms of the relative frequency of CTR and BPH to graphically display their ability to classify cases.

3. Results

3.1. Immunohistochemistry

When comparing CTR with BPH cases, no remarkable differences were observed in relation to ck18 immunoreactivity. In both groups the immunostaining to ck18 was abundant and exclusively detected in epithelial columnar cells. The immunoexpression was mainly located in lateral and apical border of the cells (Figure 2).

3.2. Global Quantitative Findings

No significant differences have been observed in global $V_V$ ck18 when CTR and BPH groups were compared (Figure 3(a)). In the other hand, the Morisita index for $V_V$ ck18 distribution was significantly higher ($p < 0.01$) in BPH than CTR (Figure 3(b)).

3.3. Local Quantitative Findings

The estimate of local $V_V$ ck18 (pixel by pixel) along of the space series shows a similar profile in both CTR and BPH cases. Nevertheless, in several spaced segments of the strips from BPH group, $V_V$ ck18 showed a significant decrease in
Figure 2. Sections immunostained to ck18 and counterstained with haematoxylin: In (a) image from a CTR case; in (b) an image from a BPH case. The cell nuclei were counterstained with haematoxylin. The scale bars represent 50 µm.

comparison with the same segment of strip from CTR group (Figure 4).

The segments of the BPH space series that show the above indicated differences were in the next ranges of distance: 997-1012, 3310-3442, 3503-3557, 3645-3653, and 6151-6336 microns (Figure 5).

The wavelet analysis indicated that positional variance of V \text{ck18} was always higher in BPH than CTR, but these differences were significant only in two patches of the space series, (Figure 6(a)). Respecting to scale variance, no significant differences were observed between CTR and BPH groups (Figure 6(b)).
Figure 3. Bar diagrams expressing mean ± SEM for: (a) global $V_v \text{ck18}$; (b) Morisita Index for dispersion of $V_v \text{ck18}$ global measures, in control (CTR) and hyperplastic (BPH) groups. The means with significant differences between them are connected by zig-zag lines, indicating the correspondent p values.

Figure 4. Diagram for space series of local $V_v \text{ck18}$ values along the X axis (position in µm) of the correspondent strips of CTR (blue line) and BPH (green line) groups. The asterisks indicate the segment of space series were the differences between CTR and BPH are significant (p < 0.05).
Figure 5. Diagram showing the ranges of distance (in µm) where CTR (blue) and BPH (green) show significant differences (p < 0.05) for local VV ck18 values, these values are expressed as mean ± SEM.

Figure 6. Diagrams for positional (a) and scale (b) variances of local VV ck18 values along the X axis (position in µm) of the correspondent strips of CTR (blue line) and BPH (green line) groups. The asterisks indicate the segment of space series were the differences between CTR and BPH are significant (p < 0.05).

The ranges in which the positional variance was significantly different were: 239-244, and 656-714 microns (Figure 7).

The estimate of Λ for VV ck18 in both CTR and BPH groups shows a clear clumping pattern for the ck18 immunoreactiveacini, and differs significantly for lacunarity curves from a random distribution of the data (Figure 8(a) and Figure 8(b)). When comparing Λ between CTR and BPH cases, the CTR group shows higher values of Λ than BPH (Figure 8(c)), but the differences were significant in only a small interval of scale values, ranging between 86 and 118 µm (Figure 9).

3.4. Discriminant Analysis

The discriminant analyses for the groups of study reveal that:
Figure 7. Diagram showing the ranges of distance (in µm) where CTR (blue) and BPH (green) show significant differences (p < 0.05) for positional variance of local $V_{V, ck18}$ values, these values are expressed as mean ± SEM.

a) From all the local $V_{V, ck18}$ values showing significant differences between CTR and BPH groups, only $V_{V, ck18}$ at 3414 and 1.3 µm respectively have discriminatory power. With these two variables in the model, 82% of the cases were correctly classified into the CTR and BPH groups. Table 1(a) shows the significant reduction of the Wilk’s lambda statistic with these variables included in the model.

b) From all the values for positional variance of $V_{V, ck18}$, showing significant differences between CTR and BPH groups, only those at 1.3 and 241 µm respectively have discriminatory power. With these two variables in the model, 82% of the cases were correctly classified into the CTR and BPH groups. Table 1(b) shows the significant reduction of the Wilk’s lambda statistic with these variables included in the model.

c) From all the Λ values for $V_{V, ck18}$ showing significant differences between CTR and BPH groups, only the Λ value at a scale of 86 µm has discriminatory power. With this variable in the model, 76% of the cases were correctly classified into the CTR and BPH groups. Table 1(c) shows a not significant reduction of the Wilk’s lambda statistic with this variable included in the model.

The histograms showing the distribution of the cases in CTR and BPH groups when the discriminant scores were applied are depicted in Figure 10.

4. Discussion

The immunohistochemical findings observed in the present study are consistent with findings by other authors [15] [16] [17] [42], regarding the absence of changes in immunoreactivity for ck18 in BPH in relation to normal prostate, although some studies show a greater immunoreactivity for ck18 in BPH compared to prostate cancer [18]. Moreover, the global quantification of the fraction of epithelial volume immunostained for ck18 ($V_{V, ck18}$) in the BPH group does not show significant differences with the CTR group, although other studies
Figure 8. Diagrams showing the changes of lacunarity ($\Lambda V_c, ck18$) in relation to scale plotted logarithmically. In (a) and (b), the curves of $\Lambda$ for CTR and BPH are compared with their corresponding distribution obtained by randomizing the data (dashed lines). In (c), the $\Lambda$ curves for CTR and BPH are compared. The asterisk indicates the segment where the differences are significant ($p < 0.05$). For each curve, the envelope of dotted lines represent the SEM of the mean in each point.
Figure 9. Diagram showing the ranges of distance (in log scale) where CTR (blue) and BPH (green) show significant differences (p < 0.05) for $\Lambda V_V \text{ck18}$ values, these values are expressed as mean ± SEM.

Table 1. Discriminant analyses.

<table>
<thead>
<tr>
<th>Entered variable</th>
<th>Wilks' $\lambda$</th>
<th>$p^3$</th>
<th>$p^4$</th>
</tr>
</thead>
<tbody>
<tr>
<td>(a)</td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>$V_V 3414$</td>
<td>0.486</td>
<td>15.88</td>
<td>0.001</td>
</tr>
<tr>
<td>$V_V 1.3$</td>
<td>0.373</td>
<td>11.78</td>
<td>0.001</td>
</tr>
<tr>
<td>(b)</td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Pvar $V_V 1.3$</td>
<td>0.632</td>
<td>8.74</td>
<td>0.01</td>
</tr>
<tr>
<td>Pvar $V_V 241$</td>
<td>0.389</td>
<td>11.99</td>
<td>0.001</td>
</tr>
<tr>
<td>(c)</td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>$\Lambda V_V 86$</td>
<td>0.796</td>
<td>3.843</td>
<td>0.06</td>
</tr>
</tbody>
</table>

Discriminant analyses to classify the CTR and BPH cases after: (a) local $V_V \text{ck18}$, (b) Positional variance (Pvar) for $V_V \text{ck18}$, and (c) lacunarity ($\Lambda$) for $V_V \text{ck18}$. 1Selected variables, a: $V_V \text{ck18}$ at 3414 and 1.3 µm respectively, b: Pvar $V_V$ at 1.3 and 241 µm respectively, and c: $\Lambda V_V$ at 86 µm. 2This column shows the Wilks’ lambda for every variable entered. 3F distribution of Snedecor, the F minimum value for entering the variables was 3.84. 4Level of significance $p < 0.05$.

have described an increase in the fraction of immunoreactive epithelial volume for prostate-specific-antigen (PSA) in BPH, compared to normal prostate [24]. This suggests that, although epithelial compartment in BPH can be expanded, the global expression of ck18 not change in relation to non-hyperplastic prostate. The significant clumping of the ck18 immunoreactive epithelium in BPH respecting to CTR revealed by the increase of Morisita index, indicates a higher heterogeneity in the distribution of hyperplastic acini compared with a more uniform distribution of acini for CTR group. This could be corroborated by the increase of branching pattern of BPH glands as shows the increase in connectivity density described in other study [24].

No global differences in $V_V \text{ck18}$ between CTR and BPH were observed. Nevertheless, at the local level, in limited segments of the space series of cases of the BPH group, a significant decrease of $V_V \text{ck18}$ is detected compared to the nor-
Figure 10. Histograms showing the distribution of percentage of cases of CTR (blue) and BPH (green) groups when the discriminant scores obtained in each discriminant analysis were applied. The best separation between CTR and BPH groups is observed for $V_v$ ck18 (a); and positional variance of $V_v$ ck18 (b); nevertheless, the discrimination for $\Lambda (V_v$ ck18) is poor (c).

Mal prostate. Although the average length of the segments of the space series with significant differences for $V_v$ ck18, represents less than 4% of the average of the total length, in the discriminant analysis carried out was observed that $V_v$ ck18 at two points in the space series, was able to correctly classify 82% of the
In addition, local changes in volume fraction immunostained for ck18, are accompanied by differences in the variability of the parameter, manifested by significant differences in the variance of $V_v$ ck18, detected by Wavelet analysis. This type of analysis has been scarcely used in prostate pathology [43], but in our study it has revealed of interest for detection of local changes in the positional variance of $V_v$ ck18. For all the space series, the positional variance was higher in BPH than CTR, but only in two segments (less of the 1% of the average of the total length) this increase was significant. As in local values for $V_v$ ck18, their positional variance at two locations of the space series, had discriminant power to correctly classify 82% of the cases studied. The local increase of the variability of $V_v$ ck18 could be put in relation to the greater variability of the volume of hyperplastic acini compared to normal acini detected elsewhere [24] and, in general terms, with the heterogeneity of glandular morphometry in BPH, as evidenced by other authors [44].

However, when scale variance was examined, no differences were detected between the CTR and BPH groups, at least in the range of scales studied, whose maximum represents 50% of the total size of the space series. This suggests that the distribution of expression of ck18, both in normal and hyperplastic prostate, not shows a preferred scale [45], although in certain situations, the scale of the processes is of great importance, for example for some models of prostate tumor growth [46].

Lacunarity analysis has proven useful to study the distribution of gaps and structural elements in different pathological situations [47]. Thus, it is a useful tool to describe tissue heterogeneity, and, for instance, the behavior of tumor cells [48]. In the present study, the profile of $\Lambda$ values in relation to the scale of analysis, agrees with the clumped distribution of acini in both CTR and BPH groups. This distribution can not be attributed at random, as suggested by the differences between $\Lambda$ curves for real cases and those constructed after randomization of the data in both CTR and BPH groups.

Although the $\Lambda$-profile for BPH always shows a greater heterogeneity for the distribution of acini ck18 immunoreactive than in controls, this difference is only significant for a very small range of scales. In addition, its discriminatory power to classify cases in the two groups studied is poor. Conversely to what is described in other types of pathologies such as bone dysplasia where the usefulness of $\Lambda$ is clearly shown to discriminate between normal and pathological bone [49].

The findings of the present study can be summarized, indicating that changes in the expression of ck18 by the hyperplastic prostatic epithelium are not homogeneous, so it is not accurate what some authors [50] indicate, that a single prostate biopsy is representative of the entire structure of the adenoma. Rather, it is necessary to consider what others suggest in the sense that benign prostatic hyperplasia is heterogeneous in terms of tissue morphometry and expression of single important genes. This finding limits the use of a single biopsy based
markers to predict biological behavior, and has significant impact on the ability of distinguishing longitudinal changes in tissue composition from sampling artifacts [51].

On the other hand, in the case of BPH, the local changes in the expression of ck18 are more evident in terms of $V_c \text{ ck18}$ and its local variability, whereas other parameters that are useful in other pathologies, such as lacunarity, are less relevant in prostatic hyperplasia.

Two possible limitations of this study are indicated below: The work has focused exclusively on one type of BPH, the adenofibromiomatosus variety, it may be worth exploring with the same quantitative tools the hyperplasias of stromal and fibro-muscular predominance.

Another factor that would be interesting to analyze is how the resolution of the images' influences on the local variance and the lacunarity measurements, for that, it would be necessary to explore those variables on images captured with different magnifications.

References


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