

# TEXTURE ANALYSIS IN PERFUSION IMAGES OF PROSTATE CANCER—A CASE STUDY

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The analysis of prostate images is one of the most complex tasks in medical images interpretation. It is sometimes very difficult to detect early prostate cancer using currently available diagnostic methods. But the examination based on perfusion computed tomography (p-CT) may avoid such problems even in particularly difficult cases. However, the lack of computational methods useful in the interpretation of perfusion prostate images makes it unreliable because the diagnosis depends mainly on the doctor's individual opinion and experience. In this paper some methods of automatic analysis of prostate perfusion tomographic images are presented and discussed. Some of the presented methods are adopted from papers of other researchers, and some are elaborated by the authors. This presentation of the method and algorithms is important, but it is not the master scope of the paper. The main purpose of this study is computational (deterministic and independent) verification of the usefulness of the p-CT technique in a specific case. It shows that it is possible to find computationally attainable properties of p-CT images which allow pointing out the cancerous lesion and can be used in computer aided medical diagnosis.

Keywords: prostate cancer, perfusion computed tomography, medical image analysis, pattern recognition.

## 1. Introduction

Prostate cancer (PCa) is one of the most common malignancies among men (ACS, 2009; NCR, 2009). In the last years there has still been observed a growth in the number of registered cases. And although it is partially connected with better and better diagnostic methods and increased knowledge among patients (resulting in better detectability of this type of cancer), there is no doubt that PCa is a serious medical and social problem.

Early detection of PCa is a key to survival. Unfortunately, routine medical tests like measuring blood concentration of prostate specific antigen (PSA), digital rectal examination (DRE), transrectal ultrasound (TRUS), and biopsy often fail (Hricak *et al.*, 2007; Roscigno *et al.*, 2004; Selley *et al.*, 1997). For example, on TRUS, cancer lesions can be hypoechoic, hyperechoic or even isoechoic (Daehnert *et al.*, 1986; Norberg *et al.*, 1997; Sudoł-Szopińska and Szopiński, 2005). In view of this, there is an obvious need for other diagnostic methods which could manage this problem in some cases which are too difficult for standard (above mentioned) methods.

There are many studies of new techniques which could address this problem, including, for example, the EPCA test (Bradford *et al.*, 2006; Leman *et al.*, 2007). Perfusion computed tomography (p-CT) is also one of these methods (still under investigation). This method allows evaluating the parameters of perfusion such as blood flow (BF), blood volume (BV), mean transit time (MTT), permeability surface (PS) in specified areas of prostate (ROI—region of interest) (Cenic *et al.*, 2000; Wintermark *et al.*, 2001).

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Nowadays the p-CT examination is used mainly in the diagnosis of brain acute stroke (Miles and Griffiths, 2003; Hartel et al., 2006; Hoeffner et al., 2004; Rosenberg et al., 2004), but the usefulness of this method has also been tested on other organs (Miles and Griffiths, 2003; Blomley et al., 1993; Dugdale and Miles, 1999; Dziubińska et al., 2006; Fukuya et al., 1995; Groell et al., 2001; Sahani et al., 2005; Wolfkiel et al., 1987; Zhang and Kono, 1997), including prostate (Henderson et al., 2003; Ives et al., 2005; Łuczyńska et al., 2008; Prando and Wallace, 2000). Its application to detecting cancerous lesions is based on documented evidence of the creation of new blood vessels in tumor (angiogenesis) (Charlesworth and Harris, 2006; Miles, 1999; 2002). Although prostate is not highly vascularized, it is supposed that p-CT can indicate these suspicious areas also in this gland.

The purpose of this study is computational verification of usefulness of the p-CT technique in a specific case, described in the paper (Łuczyńska *et al.*, 2006). In that case, diagnostics correctly pointed cancerous lesions on the p-CT image, while on TRUS there were no visible suspicious regions. However, that indication was founded only on visual assessment, so it can be considered undeterministic and unreliable.

#### 2. Images

A 60-year-old patient was examined at the Oncology Center in Cracow because of an increasing PSA level. The p-CT examination was performed with a 16-slice CT scanner (*GE Ligh Speed*). The perfusion level was measured during repeated scans of the minor pelvis at 120 kVp and 200 mAs. The scans were started about 10 s after administering of 50 ml of non-ionic contrast medium (370 mgI/ml) at the rate of 5 ml/s and lasted 50 s. The total width of the diagnosed area was 20 mm.

Parametric maps (BF, BV, MTT and PS) were drawn using the *CT Perfusion 3* application on the *Advantage Workstation* at three levels (conventionally base, middle and apex) of the gland.

In order to perform computational analysis, only the area of prostate was selected from the acquired images. The images, originally coded with pseudocolor, where blue symbolizes the area with minimal and red with maximal perfusion, were transformed into a 31-tone grayscale using the LUT table (Tadeusiewicz and Korohoda, 1997), where 0 means maximal visible perfusion (red area in pseudocolor) and 30—no perfusion. (Fig. 1)

Figure 2 presents parametric maps of the prostate (coded with pseudocolor) at the level at which pathological lesions were confirmed (Fig. 3). In this work only the image of Fig. 2(a), which represents the parameter BF, is selected for further analysis.

### 3. Co-occurrence matrices

For automatic description of the texture of particular regions on the analyzed p-CT image, the so-called cooccurrence matrices (Haralick *et al.*, 1973) were selected. There are many other texture analysis methods, but these are most universal and their potential is greatest.

Let  $I : \mathbb{Z}^2 \supset D \rightarrow G = \{1, \dots, N_g\}$  (where  $\mathbb{Z}$  denotes set of integers) be a two-dimensional discrete image with  $N_g$  gray levels. For the given image I, we define the co-occurrence matrix (GLCM):

$$P_0(i, j|d, \theta) = \#\{k, l \in D : I(k) = i, I(l) = j, \\ ||k - l|| = d, \ \angle (k - l) = \theta\}$$
(1)

or, in a normalized version,

$$P(i, j|d, \theta) = \frac{\#\{k, l \in D : I(k) = i, I(l) = j, ||k-l|| = d, \angle (k-l) = \theta\}}{\#\{m, n \in D : ||m-n|| = d, \angle (m-n) = \theta\}}$$
(2)

where  $i, j \in G$  stand for gray levels of points k and l, respectively,  $\angle (k - l)$  is the angle between vector  $\vec{kl}$  and axis  $\vec{0X}$ , d represents the distance between k and l,  $\theta$  is the direction of co-occurrence, #X represents the power (number of elements) of set X.



Fig. 1. Pseudocolor (a), grayscale after transformation (b). The arrow shows a rise in the perfusion values.



Fig. 2. p-CT images of the prostate: blood flow (BF) (a), blood volume (BV) (b), mean transit time (MTT) (c), permeability surface (PS) (d).



Fig. 3. Analyzed image (a) and cancerous area (b)—shown in black.

name

no.

 $f_1$ 

 $f_2$ 

 $f_3$ 

 $f_4$ 

 $f_5$ 

 $f_6$ 

 $f_7$ 

 $f_8$ 

 $f_9$ 

 $f_{10}$ 

 $f_{11}$ 

 $f_{12}$ 

 $f_{13}$ 

 $f_{14}$ 

 $f_{15}$ 

 $f_{16}$ 

 $f_{17}$ 

 $\frac{f_{18}}{f_{19}}$ 

 $f_{20}$ 

 $f_{21}$ 

energy	ENE	$f_1 = \sum_{i,j} P(i,j)^2$
entropy	ENT	$f_2 = -\sum_{i,j} P(i,j) \log P(i,j)$
homogeneity	IDM	$f_3 = \sum_{i,j} \frac{1}{1 + (i-j)^2} P(i,j)$
inertia	CON	$f_4 = \sum_{i,j} (i-j)^2 P(i,j)$
correlation	COR	$f_5 = -\sum_{i,j} \frac{(i-\mu_x)(j-\mu_y)}{\sigma_x \sigma_y} P(i,j)$
variance	VAR	$f_6 = \sum_{i,j} (i + j - \mu_x - \mu_y)^2 P(i,j)$
shade	SHA	$f_7 = \sum_{i,j} (i+j - \mu_x - \mu_y)^3 P(i,j)$
prominence	PRO	$\frac{i,j}{f_6 = \sum_{i,j} (i+j-\mu_x-\mu_y)^2 P(i,j)} \frac{f_6 = \sum_{i,j} (i+j-\mu_x-\mu_y)^2 P(i,j)}{f_7 = \sum_{i,j} (i+j-\mu_x-\mu_y)^3 P(i,j)} \frac{f_8 = \sum_{i,j} (i+j-\mu_x-\mu_y)^4 P(i,j)}{2N}$
sum average	SA	$f_{9} = \sum_{i=2}^{2N_{g}} i P_{x+y}\left(i\right)$
sum entropy	SE	$f_{9} = \sum_{i=2}^{2N_{g}} iP_{x+y}(i)$ $f_{10} = -\sum_{i=2}^{2N_{g}} P_{x+y}(i)\log P_{x+y}(i)$
sum variance	SV	$f_{11} = -\sum_{i=2}^{2} (i - f_9)^2 P_{x+y}(i)$
difference average	DA	$f_{12} = \sum_{i=0}^{N_g - 1} i P_{x-y}(i)$
difference entropy	DE	$\frac{f_{12} = \sum_{i=0}^{N_g-1} iP_{x-y}(i)}{f_{13} = -\sum_{i=0}^{N_g-1} P_{x-y}(i) \log P_{x-y}(i)}$
difference variance	DV	$f_{13} = -\sum_{i=0}^{N_{g-1}} P_{x-y}(i) \log P_{x-y}(i)$ $f_{14} = -\sum_{i=0}^{N_{g-1}} (i - f_{12})^2 P_{x-y}(i)$ $f_{15} = \frac{f_2 - \text{HXY}_1}{\max(\text{HX}, \text{HY})}$ $f_{16} = \frac{\sigma(P(i, j))}{\mu(P(i, j))}$ $f_{17} = \max(P(i, j))$
information measure	IMC1	$f_{15} = \frac{f_2 - \text{HXY}_1}{\max(\text{HX}, \text{HY})}$
coefficient of variation	COV	$f_{16} = \frac{\sigma(P(i,j))}{\mu(P(i,j))}$
peak transition probability	MAX	$f_{17} = \max(P(i,j))$
diagonal variance	DIAV	$f_{18} = \sigma^2(P(i,j))$
diagonal moment	DIAM	$f_{19} = \sum_{i,j} \left( \frac{1}{2}  i - j  P(i,j) \right)^{\frac{1}{2}}$
second diagonal moment	DSM	$f_{20} = \sum_{i,j} \frac{1}{2}  i - j  P(i,j)$
triangular symmetry	TRS	$f_{21} =  P(i,j) - P(j,i) $

 Table 1. Coefficients of GLCM.

 abbr.

value

Notation

$$\begin{aligned} \mu_x &= \sum_i i \sum_j P(i,j), \mu_y = \sum_j j \sum_i P(i,j), \\ \sigma_x &= \sum_i (i - \mu_x)^2 \sum_j P(i,j), \sigma_y = \sum_j (j - \mu_y)^2 \sum_i P(i,j), \\ P_x(i) &= \sum_j P(i,j), P_y(j) = \sum_i P(i,j), \\ P_{x+y}(k) &= \sum_{i,j: \ i+j=k} P(i,j), P_{x-y}(k) = \sum_{i,j: \ |i-j|=k} P(i,j), \\ \text{HX-entropy } P_x(i), \text{HY-entropy } P_y(j), \text{HXY}_1 = -\sum_{i,j} P(i,j) \log(P_x(i)P_y(j)) \end{aligned}$$

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The GLCM allows us to evaluate a number of coefficients, which characterize the textures of the analyzed image. Table 1 shows the list of 21 coefficients used in our study.

## 4. Results

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For the given image of Fig. 3(a) transformed to grayscale we evaluated the first-order statistics calculated directly from the image histogram (Table 2). The mean in a healthy area is smaller than in a cancerous one but the variance is very high in both cases. Therefore, the analysis based only on the first-order statistics of the ROI considered (see below) may not be sufficient (Table 3, Fig. 6).

The ROIs covering the analyzed image were rectangular in shape, 10 pixels wide and 20 pixels high. Each consecutive ROI was selected 10 pixels apart the previous



Fig. 4. Nearest neighborhood of the point (x, y) and directions of co-occurrence (a), co-occurrence can be considered also for a greater distance between pairs of points (b).



- Fig. 5. Example of GLCM: source image with four gray levels (a), illustration of counting co-occurrences for  $d = 1, \theta = 0^{\circ}$  (b), GLCM,  $d = 1, \theta = 0^{\circ}$  (counted co-occurrences are divided by the number of all pairs of points considered (here 9) (c). In this example the values were rounded to two decimal places.
- Table 3. Statistics of the ROI. For each ROI considered, the mean and median were evaluated. In the table we show the minimum and the maximum of those values—separately for ROIs covering the healthy region and separetely for ROIs in the cancerous area.

ROI	m	ean	median		
KOI	min	max	min	max	
healthy region	2.33	9.92	1	10	
cancerous region	8.17	16.42	4	16	

one. Those where less than half of the pixels covered the area of prostate were missed. Each ROI was classified according to the pattern shown in Fig. 3(b). There were 88 ROIs at all: 82 healthy and six cancerous (Fig. 7). For each ROI, normalized GLCM matrices (see Eqn. (2)) and coefficients were evaluated.

There were calculated 21 coefficients (Table 1) for each GLCM characterized by distance d in the range from 1 to 9, and angle  $\theta$  with values 0°, 45°, 90°, 135°, and also d in the range from 10 to 19 and  $\theta = 90°$ . So it was the 966-dimensional feature space. The resulting values for each feature were analyzed in order to eliminate outliers and normalized. The distribution of each feature was equalized using the ladder of powers method (Tukey, 1977; Velleman and Hoaglin, 1981) (see Eqn. 3) with  $\gamma \in (0, 2]$ .

$$\operatorname{error}(\gamma) = \sum_{c=1,2} \left( \int_{x} [\operatorname{cdf}\{x_{c}^{\gamma}\} - \Phi\{\overline{x_{c}^{\gamma}}, \operatorname{var}(x_{c}^{\gamma})\}]^{2} \right), \quad (3)$$

where  $c = \{1, 2\}$  represents classification,  $\operatorname{cdf}(x_c^{\gamma}), \overline{x_c^{\gamma}},$ var $(x_c^{\gamma})$  stand for the distribution function, mean and variance of empirical distribution for class c, respectively,  $\Phi(\mu, \sigma^2)$  is a normal distribution function with mean  $\mu$ and variance  $\sigma^2$ .

We were looking for  $\gamma_{opt}$  which minimizes the function error( $\gamma$ ):

$$\gamma_{\text{opt}} = \min_{\gamma} \operatorname{error}(\gamma).$$
 (4)



Fig. 6. Mean and median of the analyzed ROIs. Light circles healthy regions, dark squares—cancerous regions.



Fig. 7. Analyzed image (a) and pattern (b). The cancerous area is shown in black. Also ROIs classified as cancerous (dark gray) and not analyzed (light gray) are shown on the pattern. Other ROIs, which are not shown, were classified as a healthy area.



Table 2. First-order statistics of the analysed image.

Features where  $\operatorname{error}(\gamma_{opt}) \geq 1$  were excluded from further analysis. For each of the remaining features, the Bhattacharyya measure (Bhattacharyya, 1943) was used for the normal distribution:

$$J = \frac{1}{4} \frac{(\mu_1 - \mu_2)^2}{\sigma_1^2 + \sigma_2^2} + \frac{1}{2} \ln\left(\frac{\sigma_1^2 + \sigma_2^2}{2\sigma_1 \sigma_2}\right), \quad (5)$$

where  $\mu_1, \mu_2$  are means,  $\sigma_1, \sigma_2$  are standard deviations for Classes 1 and 2, respectively. Below, in Table 4, we present a list of the best discriminating properties. As is shown, the best results were produced for the diagonal moment  $(f_{19})$  and various d and  $\theta$ . It should be noted that diagonal directions  $\theta = 45^{\circ}$  and  $\theta = 135^{\circ}$  did not occur in any of the best ten features.

As can be remarked, the above-mentioned consideration is limited to the indication of a single individually best discriminating feature (Fig. 8). It should be observed that, in spite of these limitations, it is possible to indicate such features which individually have the ability to distinguish a healthy and a cancerous area (Fig. 9). However, it is not a universal rule-even for features with a large distance between classes, sometimes these areas cannot be separated (Fig. 10). In such cases it can be helpful to increase the dimension of the feature space (Fig. 11).

0.0	0.1	0.2	0.3	0.4		— <b></b> 0.6	0.7	0.8	0.9 =90°, f	1.0
••••• 0.0	0.1	0.2	0.3	0.4	0.5	0.6	•-•••- 0.7	0.8	0.9	1.0
healthy accercous							c	l=1, θ=	=0°, f <sub>21</sub>	

Fig. 9. Classification of two example parameters from Table 4. It is possible to point out the border value, where lower values suggest a healthy area and higher values mean susceptibility of a cancerous lesion.



Fig. 10. In spite of high discriminant power in these examples we cannot point out the border value.

## 5. Conclusion

In this paper it was shown that it is possible to select such parameters of an image which are deterministic and independent of a personal assessment. Our results confirm the usefulness of the p-CT method applied to PCa diagnosis in the analyzed case. Of course, it is obvious that only one case cannot be generalized, but in this study the potential of this method can be seen.

At the Oncology Center in Cracow the p-CT method is used to examine other patients. Thanks to that it will be possible to verify the usefulness of the proposed algorithm. In further work the authors will also expand research to other perfusion parameters to determine the effectiveness of each one.

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Fig. 11. Improvement of discriminant power after increasing the feature space dimension. Features shown in Fig. 10, which were useless individually, are very good to separate the classes in a two-dimensional space.

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displacement d	angle $\theta$	coefficient	Bhattacharyya measure
8	0	diagonal moment $(f_{19})$	2.494145
10	90	diagonal moment $(f_{19})$	2.441980
1	0	triangular symmetry $(f_{21})$	2.181827
3	90	prominence $(f_8)$	2.074908
3	0	variance $(f_6)$	2.051128
6	90	sum entropy $(f_{10})$	1.962097
11	90	diagonal moment $(f_{19})$	1.860219
4	90	prominence $(f_8)$	1.843247
11	90	second diagonal moment $(f_{20})$	1.839980
4	0	sum entropy $(f_{10})$	1.836315

Table 4. List of ten features with the best discriminant power.

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Fig. 8. Analyzed image (a), illustration of the best discriminating features (b–k), and of the feature with no discriminant power (l).  $d = 8, \theta = 0^{\circ}, f_{19}$  (b),  $d = 10, \theta = 90^{\circ}, f_{19}$  (c),  $d = 1, \theta = 0^{\circ}, f_{21}$  (d),  $d = 3, \theta = 90^{\circ}, f_8$  (e),  $d = 3, \theta = 0^{\circ}, f_6$  (f),  $d = 6, \theta = 90^{\circ}, f_{10}$  (g),  $d = 11, \theta = 90^{\circ}, f_{19}$  (h),  $d = 4, \theta = 90^{\circ}, f_8$  (i),  $d = 11, \theta = 90^{\circ}, f_{20}$  (j),  $d = 4, \theta = 0^{\circ}, f_{10}$  (k),  $d = 8, \theta = 45^{\circ}, f_{15}$  (l).

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