Introduction: Gynaecological cancers, including cervical cancer, often require a multidisciplinary approach that includes external beam radiotherapy, chemotherapy, and/or surgical treatment. Biological parameters of the tumour evaluated in ¹⁸F-FDG-PET/CT are used for target volume delineation in radiotherapy planning. The choice of segmentation method may affect the assessment of metabolic tumour volume (MTV) in ¹⁸F-FDG-PET/CT.

Aim of the study: To find the optimal segmentation method for the assessment of primary MTV in ¹⁸F-FDG-PET/CT in cervical cancer patients for radiotherapy planning.

Material and methods: Retrospective analysis was performed on a group of 30 patients with newly diagnosed, histologically confirmed cervical cancer. The primary MTVs were assessed by SUV_{max} and SUV_{mean} values; three segmentation methods were used to assess the primary MTV: constant threshold of SUV_{max} of 2.5, threshold of SUV_{max} 35%, and threshold of SU-V_{max} 45%. The MTVs were compared with the tumour volumes obtained in magnetic resonance imaging (MRI), which was the "gold standard", to select the best optimal segmentation method reflecting the tumour size. Wilcoxon-Mann-Whitney and *t*-test were used for statistical analysis.

Results: Depending on the segmentation method chosen, significant differences in the MTVs were obtained (p < 0.001). The highest volumes were obtained using the method based on constant SUV_{max} of 2.5, while the smallest in case of threshold of SUV_{max} of 45%. Regarding the volume determined by MRI, a 35% SUV_{max} threshold was chosen as the most reliable method.

Conclusions: The choice of appropriate segmentation method has a significant impact on the primary MTV assessment in ¹⁸F-FDG-PET/CT in patients with cervical cancer.

Key words: cervical cancer, segmentation, ¹⁸F-fluorodeoxyglucose, PET/CT, SUV.

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The effect of different segmentation methods on primary tumour metabolic volume assessed in ¹⁸F-FDG-PET/CT in patients with cervical cancer, for radiotherapy planning

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Introduction

In 2014 cervical cancer was the sixth most common gynaecological cancer diagnosed, seventh in terms of mortality among malignant tumours in Poland [1], and ranks fourth for both incidence and mortality worldwide [2]. External beam radiotherapy is one of the treatment options of this cancer and is more commonly used as either an independent method of treatment or supplementary to chemotherapy or surgery. Appropriate determination of the primary tumour boundaries in the treatment planning system plays a pivotal role not only in minimising the dose to the surrounding organs at risks, but also in minimising the risk of recurrence [3].

Metabolic tumour volume (MTV) assessed by PET/CT imaging using ¹⁸F-fluorodeoxyglucose (¹⁸F-FDG), a fluorine-radiolabelled glucose analogue, helps to assess the size of the biologically active tumour [4]. Based on Otto Warburg's observation of increased glucose utilisation in tumour cells [4], the radiopharmaceuticals, which are captured similarly to glucose, e.g. ¹⁸F-FDG, can be used in the imaging of cancerous tumours [5, 6]. The algorithm used to evaluate MTV in PET/CT can be based on: visual assessment, fixed maximum standardized uptake value (SUV_{max}), or the threshold method based on the percentage of SUV_{max} [7].

The aim of the study was to assess the effect of different segmentation methods on primary metabolic tumour volume assessed in a ¹⁸F-FDG-PET/CT study in patients with cervical cancer.

Material and methods

A retrospective analysis was performed on 30 newly diagnosed patients with cervical cancer (mean age: 55 ±11 years, range: 82–28 years), admitted to the Department of Radiotherapy and Gynaecological Oncology at the Greater Poland Cancer Centre between January 2012 and October 2014 for radiotherapy planning. Each patient underwent physical and ultrasound gynaecological examination with an abdominal and vaginal probe. Additionally, magnetic resonance imaging (MRI) was performed to determine the stage of disease and for radiotherapy planning. A ¹⁸F-FDG-PET/CT examination was performed in the Nuclear Medicine Department in Greater Poland Cancer Centre on a Gemini TF 16 PET/CT scanner prior to treatment, within a month

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of diagnosis. The procedure was performed in patients who were fasting for at least five hours before the test (mean glucose level was 93 ±18 mg/dl) in order to avoid problems with the interpretation of the study. The patients after the administration of the isotope stayed in a darkened room at room temperature to rest without any excessive physical activity. Acquisition was performed an average of 60 min after intravenous injection of 374 ±75 MBq of ¹⁸F-FDG. The study protocol extended from the calvaria to the mid-thigh and included 10-12 bed positions, with 1.5 min per bed position and cutting scans 5 mm thick. The study began with the implementation of simultaneous, low-dose computed tomography (CT), beginning in the brain area, through the chest, abdominal cavity, and pelvis to better visualise the areas with increased uptake and to correct attenuation. Afterwards, the PET acquisition was performed without changing the patient's position. Reconstructions were performed at an EBW workstation using the time-of-flight (TOF) technique.

As a result of the PET/CT analysis, areas with increased metabolism considered as active tumour foci were indicated. The primary tumour metabolic activity was assessed by $\mathsf{SUV}_{\mathsf{max}}$ and $\mathsf{SUV}_{\mathsf{mean}}$, and calculated from the formula:

$$SUV = \frac{r}{a/w}$$

where r is the radioactivity concentration (kBq/ml) measured by the PET scanner within a region of interest (ROI), a is the decay-corrected amount of injected ¹⁸F-FDG (MBq), and w is the patient's weight (kg).

Metabolic volumes of the primary tumour were calculated using three segmentation methods based on:

- SUV_{max} value (2.5),
- threshold of SUV_{max} 35% (Th 35%),
 threshold of SUV_{max} 45% (Th 45%).

The volumes taken from ¹⁸F-FDG-PET/CT examination were compared with the volumes obtained in MRI, which was treated as a "gold standard" because it is the most precise imaging method for the assessment of the primary tumour volume in cervical cancer [8], to select the optimal segmentation method in PET. Volume measurements were calculated automatically in dedicated workstations in the Nuclear Medicine and Radiology Department, without user modifications.

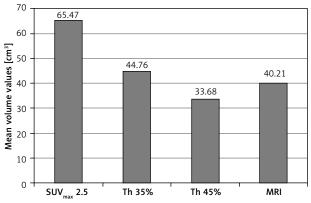


Fig. 1. The effect of different segmentation methods on the volume of primary tumour assessed in the whole group

Normality of the data distribution was assessed using Kolmogorov-Smirnov test or W Shapiro-Wilk test. For statistical analysis the Wilcoxon-Mann-Whitney and t-test were used. Pearson coefficients were used to estimate correlation between parameters. All statistical analysis was performed in the extended STATISTICA application version 13.1 with a medical package, created by StatSoft Polska 2014, as well as the extended Microsoft Excel 2013 application with the Analyse-it package. The mean values of $\mathsf{SUV}_{\mathsf{max}}\!\text{, }\mathsf{SUV}_{\mathsf{mean}}\!\text{, }\mathsf{and}$ MTV parameters were evaluated using three segmentation methods. The value of p < 0.05was considered statistically significant.

Results

The results of the PET/CT study divided the 30 patients into three subgroups: A - disease limited only to the cervix (33%), B – disease limited to the cervix and iliac lymph nodes (37%), C – disseminated disease to the lymph nodes above aorta bifurcation or other organs (30%). In all analyzed groups non-keratinizing squamous cell carcinoma (SCC) was diagnosed.

The analysis of the results performed in the whole group of patients, using different segmentation methods, showed significant differences (p < 0.001) in the mean volumes – the largest tumour volumes were obtained using the method based on $SUV_{max} 2.5$ (MTV = 65.47 cm³), and the smallest with the method based on threshold of SU- V_{max} 45% (MTV = 33.68 cm³), as shown in Figure 1. Different tumour volumes also resulted in a differentiation in metabolic activity parameters measured by SUV_{mean} values. Differences in the $\mathsf{SUV}_{\mathsf{mean}}$ values of the metabolically active tumour, depending on the segmentation method applied, reached 50% (Fig. 2). Depending on the chosen segmentation method, statistically significant differences (p < 0.001) were demonstrated between SUV_{mean} and MTV values for all three groups (Table 1).

In comparison with MRI, the highest correlation (r =0.85, p = 0.01) in reading the primary tumour metabolic volumes in PET/CT was achieved using the threshold method of segmentation based on 35% of SUV_{max}. In the case of the SUV_{max} 45% threshold and constant value of SUV_{max} 2.5, the correlation was lower (r = 0.75, p = 0.03and r = 0.46, p = 0.07, respectively).

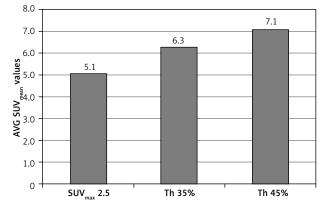


Fig. 2. Average SUV_{mean} values depending on the chosen segmentation method in the whole group

Table 1. The influence of different segmentation methods on the evaluated parameters depending on the group

	SUV _{max} 2.5			Th 35%			Th 45%		
	Α	В	С	Α	В	С	Α	В	С
SUV _{max}	11.7 ±4.9	13.1 ±4.7	11.1 ±4.4	12.1 ±5.1	13.1 ±4.7	11.1 ±4.4	12.5 ±5.6	13.1 ±4.7	11.1 ±4.4
SUV_{mean}	4.8 ± 1.1	5.5 ±1.4	4.8 ±1.3	6.0 ±2.1	6.9 ±2.6	5.7 ±2.6	6.9 ±2.3	7.7 ±2.7	6.4 ±2.8
MTV (cm³)	51.01 ±62.79	68.90 ±62.96	67.28 ±52.85	34.13 ±32.06	44.48 ±35.59	49.60 ±36.22	26.06 ±28.23	33.65 ±28.42	35.47 ±27.85

A – disease limited to the cervix (33%), B – disease limited to the cervix and iliac lymph nodes (37%), C – disseminated disease to the lymph nodes above aorta bifurcation or other organs (30%)

Discussion

Determining the volume of the primary tumour is crucial in radiotherapy planning. Because the $^{18}\text{F-FDG-PET/CT}$ examination is based on Warburg's observation of increased glucose consumption in malignant tumours, it is regarded as an effective method of determining the metabolic tumour volume of cancerous tissue [9, 10]. According to Sridhar et~al., metabolic volumes assessed by $^{18}\text{F-FDG-PET/CT}$ are more reliable for large tumours (\geq 10 ml) than for small tumours (\leq 10 ml) [11]. In this study we did not observe such a dependence. The volumes of primary tumours were comparable to those from MRI imaging when a threshold of SUV $_{\rm max}$ 35% method was chosen, regardless of the size of the primary tumour.

Many methods have been proposed for determining tumour volume in the ¹⁸F-FDG-PET/CT study. One of them is the manual method in which a nuclear medicine specialist defines the border between tumour and normal tissue. This method largely depends on the selection of the right window for CT imaging as well as tumour and surrounding tissue characteristics. It is also dependent on the person evaluating the test, with a variability ranging from 5% to 45% [11, 12]. Hatt et al. analyzed 25 patients with nonsmall cell lung cancer (NSCLC) and compared the volumes obtained with CT and PET. They showed that volumes obtained with CT (55 \pm 74 cm³) were significantly higher (p <0.0001) than those assessed in the PET study (ranging from 18 ±25 to 47 ±76 cm³, depending on the chosen delineation method) [13]. The method based on 50% threshold cut-off point proposed by Wu et al. led to the greatest underestimation of MTV, with errors up to +1.8 cm (+32%) [14]. Some authors have reported underestimation of tumour volumes assessed by SUV-based segmentation methods, especially in the case of small tumours, which is of particular importance when determining the size of lymph nodes or small primary lesions [15]. Our study showed the largest volumes in the segmentation method based on SUV 2.5 values in all three groups of patients, which did not correlate with the volumes assessed by MRI. Currently, the threshold adaptive method is one of the most commonly used segmentation methods, among others in head and neck, lung, and cervical cancers [11].

Many authors suggest that MTV evaluated by ¹⁸F-FDG-PET/CT study is a predictor of the patient's outcome in solid tumours. Romesser *et al.* compared SUV_{max}, gross tumour volume (GTV) and MTV for disease control and survival in patients with head and neck squamous cell carcinoma (HNSCC) [16]. Using the Kaplan-Meier method they con-

cluded that MTV and GTV are better prognostics factors than SUV_{max} for the prediction of local control and overall survival in HNSCC [16]. In addition, metabolic tumour volume has been shown to be a better prognostic parameter than SUV_{max} itself in predicting patients' outcome, and it may become useful in clinical evaluation [17–19]. Miller and Grigsby noticed that the primary tumour volume determined using ¹⁸F-FDG-PET affects both the overall survival (p = 0.003) and the time to detect the disease spread (p = 0.005) in patients with advanced cervical cancer [20].

Xu et al., in their analysis of 55 patients with planoepithelial cervical cancer, assessed the effect of different segmentation methods on the metabolic volumes obtained in the ¹⁸F-FDG-PET/CT study in relation to the size of the tumour obtained from the pathologist [7]. Depending on the SUV_{max} value, patients were divided into four subgroups, in which statistically significant differences were observed depending on the choice of the segmentation method (p < p0.005). Metabolic volumes assessed based on 20% and 30% thresholds overvalued, while the threshold 40%, 50%, and 60% undervalued the results, in comparison with GTV assessed in pathological examination performed after radical hysterectomy. The most reliable results were obtained using the iterative adaptive segmentation algorithm [7]. Kim et al. compared MTV at threshold 40% in 45 patients with invasive cervical cancer with pathological and prognostic results [21]. The results showed that patients with MTV greater than 20 cm³ had a reduced disease-free survival time compared to those with MTV below 20 cm³ (p = 0.029). Similar studies conducted by Ciernik et al. evaluated the use of PET/CT in the planning of radiotherapy, and 50% SUV_{max} turned out to be a reliable reflection with the tumour volume assessed in the CT technique [22]. Showalter et al. applied a 40% threshold of SUV_{max} to define the primary tumour diameter in the early stage of cervical cancer and found a significant positive correlation with the tumour diameter assessed by the pathologist (r = 0.757, p < 0.0001) [23].

This study has a few limitations. The main one is the small number of patients. Secondly, ¹⁸F-FDG-PET/CT cannot precisely assess the severity of the primary tumour local infiltration, due to high physiological activity of the bladder and rectum. Also, physiological ¹⁸F-FDG uptake is shown, e.g. in urinary tract, ovaries, and uterus during various phases of menstruation, which can mask the malignant lesions. False negative results might be gained in the case of small lesions (< 7 mm) or different tumour subtypes (e.g. neuroendocrine tumour). False positive results could potentially be obtained in the case of inflammatory or infectious processes [24, 25].

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MRI has limited value in the differentiation of metastatic lymph nodes from hyperplasia, in detecting microinvasion in nonenlarged lymph nodes, and for the evaluation of cervical stroma [26, 27]. The limitation in the evaluation of primary tumour size in MRI also includes the impossibility of margin definition, confusing with peritumoral reactions or scars and small tumours (< 1 cm), which might be only visualised on early dynamic contrast-enhanced sequence (DCE) MRI as enhanced foci [28, 29].

Conclusions

In patients with cervical cancer the choice of appropriate segmentation method of primary tumour in $^{18}\mbox{F-FDG-PET/CT}$ has a significant impact on radiotherapy planning. The metabolic tumour volumes obtained in $^{18}\mbox{F-FDG-PET/CT}$ correlated best with MRI volumes when a threshold of 35% of the SUV $_{\mbox{\scriptsize max}}$ value was chosen as a segmentation method.

The authors declare no conflict of interest.

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