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ORIGINAL ARTICLE

Biologic targets identified from dynamic ¹⁸FDG-PET and implications for image-guided therapy

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Abstract

Purpose. The outcome of biologic image-guided radiotherapy depends on the definition of the biologic target. The purpose of the current work was to extract hyperperfused and hypermetabolic regions from dynamic positron emission tomography (D-PET) images, to dose escalate either region and to discuss implications of such image guided strategies. *Methods.* Eleven patients with soft tissue sarcomas were investigated with D-PET. The images were analyzed using a two-compartment model producing parametric maps of perfusion and metabolic rate. The two image series were segmented and exported to a treatment planning system, and biological target volumes BTV_{per} and BTV_{met} (perfusion and metabolism, respectively) were generated. Dice's similarity coefficient was used to compare the two biologic targets. Intensity-modulated radiation therapy (IMRT) plans were generated for a dose painting by contours regime, where planning target volume (PTV) was planned to 60 Gy and BTV to 70 Gy. Thus, two separate plans were created for each patient with dose escalation of either BTV_{per} or BTV_{met} . *Results.* BTV_{per} was somewhat smaller than BTV_{met} ($209 \pm 170 \text{ cm}^3$ against $243 \pm 143 \text{ cm}^3$, respectively; population-based mean and s.d.). Dice's coefficient depended on the applied margin, and was 0.72 ± 0.10 for a margin of 10 mm. Boosting BTV_{met} gave smaller dose differences between the respective non-boost DVHs (such as D_{98}). *Conclusions.* Dose escalation of one of the BTVs results in a partial dose escalation of the other BTV as well. If tumor aggressiveness is equally pronounced in hyperperfused and hypermetabolic regions, this should be taken into account in the treatment planning.

Positron emission tomography (PET), using the glucose analogue 18F-fluorodeoxyglucose (¹⁸F-FDG) as a tracer, can be used for detection of metastasis and prediction of therapy outcome for certain cancers such as in the head and neck using the standard uptake value; SUV [1–3]. PET has been proposed as basis for dose escalation through dose painting by numbers or contours, for both non-small cell lung cancer [4,5] and head and neck cancer [6,7]. Local recurrence for these cancers frequently occurs in PET active regions, but to cover all tumor extensions multi-modality imaging is required [8].

Dynamic contrast enhanced (DCE) imaging utilizes the image contrast formed by an intravenously administered contrast agent, which either is paramagnetic, often gadolinium-based [in the case of magnetic resonance imaging (MRI)] or radiopaque, iodine-based [in the case of computed tomography (CT)]. DCE imaging may be used to assess vascular features such as tumor perfusion [9], which has shown a predictive role, e.g. cervical cancer [10]. However, dynamic ¹⁸F-FDG-PET, where tissue accumulation of ¹⁸F-FDG is followed in space and time, has been shown to provide both estimates of perfusion and metabolism [11,12]. Thus, dynamic ¹⁸F-FDG-PET is a multiparametric imaging modality. As tumors may be a heterogeneous mass with respect to perfusion and metabolism, volumetric information on such tissue properties may have implications for the optimal dose distribution during radiation therapy [13,14].

There are ongoing radiotherapy trials where advanced medical imaging is used as basis for focal dose escalation in order to improved treatment outcome. These trials either employ conventional ¹⁸F-FDG-PET (non-small cell lung cancer) [4] or

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functional MRI (prostate cancer) [15], in particular, DCE-MRI. For the former, the hypermetabolic part of the tumor is boosted, while in the latter the hyperperfused part is boosted. In these single-modality imaging studies, the rationale for defining the boost regions was based on significant correlations between FDG-PET or DCE-MRI and pathology (for lung and prostate cancer, respectively), as pathology may be reflected in both elevated metabolism and perfusion. However, studies correlating multiparametric images with pathology are, to the best of our knowledge, lacking, and vital pathology may thus be missed if only a single imaging parameter is considered. Therefore, if both hypermetabolic and hyperperfused regions can be depicted in the same tumor, the question rises whether these regions overlap. Furthermore, if the regions do not overlap, what are the consequences with respect to dosage of the region that is not considered as a biologic target in the treatment planning? These issues are addressed in the present work.

In the current work, patients with soft tissue sarcoma have been examined with dynamic¹⁸F-FDG-PET prior to therapy. In an earlier work, we found distinct differences between derived images reflecting metabolism and perfusion for these patients [16]. In the current work, we have used pharmacokinetic analysis to produce surrogate maps of metabolism and perfusion, where the maps were used as targets in biological image-guided dose escalation. The resulting maps were compared, and the dose distributions in the respective targets were evaluated.

Materials and methods

Patients and FDG-PET

This study comprised 11 patients with soft tissue sarcoma, and was approved by regional committee for research ethics. Written informed consent was obtained from all patients. The patients were examined at a Biograph 16 PET/CT-scanner (Siemens AG, Erlangen, Germany). The patient cohort and PET examinations have been comprehensively described in a previous study [16]. Briefly, all patients were scanned in dynamic PET mode (duration 45 min) after intravenous injection of ¹⁸F-FDG. The dynamic FDG-PET (D-PET) images were used to derive pharmacokinetic maps, most notably of K_1 (Perfusion), v_p (Vascular fraction), k3 (Metabolic potential), and MR_{FDG} (metabolic rate), using a standard two compartment model and custom made software in IDL (Exelis Visual Information Solutions, Boulder, USA).

Directly after the bolus injection the FDG is concentrated in the blood and the activity will display the vasculature [17–20], largely depicted by K_1 . At late time points, the images will largely reflect metabolism, as depicted by MR_{FDG} though the precise limiting factor is still unclear [21–23].

Treatment planning

The K_1 and MR_{FDG} maps were segmented using Otsu's method, where islands smaller than six voxels were eliminated. The resulting binary masks were exported to the Oncentra treatment planning system (Nucletron – an Elekta Company, the Netherlands) together with corresponding PET/CT images. The gross tumor volume (GTV) was manually delineated in the PET/CT images. A clinical target volume (CTV) was generated from the GTV by adding a 5 mm isotropic margin. Finally, the PTV was generated from the CTV by adding a 1 cm isotropic margin. Hyperperfused and hypermetabolic biological target volumes (BTVs), BTV_{per} and BTV_{met} , were generated based on the thresholded K_1 and MR_{FDG} maps, respectively. An isotropic margin of 1 cm (tentatively accounting for internal motion and setup) was used.

A seven-field IMRT plan was set up for a dose painting by contours regime, where PTV, excluding BTV, was prescribed to a mean dose of 60 Gy and either BTV_{per} or BTV_{met} to a mean dose of 70 Gy. Thus, two separate plans were created for each patient with dose escalation of either BTV_{per} or BTV_{met}. Organs at risk did not impose any restrictions on the resulting dose distributions. Dose volume histograms (DVHs) for PTV, BTV_{per} and BTV_{met} were exported and further analyzed.

Analysis

All analysis was performed using IDL (Exelis Visual Information Solutions, Boulder, USA). Dice's similarity index was used as a measure of the overlap of two segmented maps X and Y: $2 \times X \cap Y/|X| \times |Y|$ Mann-Whitney tests were used to assess differences in dose volume parameters. A significance level of 0.05 was chosen.

Results

The pharmacokinetic analysis of the D-PET images, resulting K_1 and MR_{FDG} maps in a representative axial plane of a tumor is shown together with the workflow for generating the two BTVs in Figure 1. The derived BTV_{per} and BTV_{met} (including a 10 mm margin) are also indicated. As seen, the tumor is not homogeneous with respect to the two pharmacokinetic parameters. Furthermore, the derived BTV_{per} and BTV_{met} are not spatially co-localized. Figure 2 shows the dependence



Figure 1. Target definition workflow: parametric images (top) produce respective threshold regions (middle), and clinical margins are applied (bottom). Perfusion is in green, metabolic rate in blue, and overlap in red.

of Dice's similarity index (DSI) for BTV_{per} and BTV_{met} on the clinical margin. As seen, patient-averaged DSI increases from 0.51 to 0.72 for an increase in margin from 0 to 10 mm. The latter margin gave mean volumes of 209 ± 170 cm³ and 243 ± 143 cm³ for BTV_{per} and BTV_{met}, respectively (Supplementary Table I, to be found online at http://informahealthcare.com/ doi/abs/10.3109/0284186X.2013.813071). A margin of 10 mm is used in the following.



Figure 2. The Dice's similarity coefficient (DSI) between BTV_{per} and BTV_{met} as a function of the clinical margin for each region.

IMRT planning resulted in appropriate dose distributions in the target volumes for all patients and PTV-BTV combinations. Figure 3 shows two plans based on either boosting of BTV_{per} or BTV_{met} for the patient showing the smallest DSI for these volumes. The boost dose region conforms to the respective BTV, making the dose to the non-boosted BTV rather low. Cohort-based mean DVHs were generated for PTV, $\mathrm{BTV}_{\mathrm{per}}$ and $\mathrm{BTV}_{\mathrm{met}}$ based on the two boost plans (Figure 4). The mean DVHs for the boosted and non-boosted volumes appear rather similar, in particular for the plan where BTV_{met} was boosted. Key dose parameters are reported in Table I. For the plan where BTV_{per} is dose escalated, significant differences were found between DVHs of BTV_{per} and $BTV_{met}.$ For instance, D_{98} was significantly lower in $BTV_{met}.$ The corresponding cohortbased mean doses in $\mathrm{BTV}_{\mathrm{per}}$ and $\mathrm{BTV}_{\mathrm{met}}$ was 69.0 ± 1.0 Gy and 67 ± 3.2 Gy, respectively. For the plan where $\mathrm{BTV}_{\mathrm{met}}$ is dose escalated, there were smaller differences between the DVHs and the median doses were not statistical significant. The cohort-based mean doses in BTV_{met} and BTV_{per} were in this case 69.0 ± 1.0 Gy and 68 ± 1.7 Gy, respectively.

Discussion

Solid tumors are heterogeneous entities, functionally and genetically [24], and both magnitude and spatial heterogeneity of metabolism and vascularity seem to

Table I. Results from the dose planning given as median and range. The p-values results from of a Mann-Whitney test of the dose metrics in BTV_{per} and BTV_{met} .

			*		
Boost	DVH	PTV	$\mathrm{BTV}_{\mathrm{per}}$	$\mathrm{BTV}_{\mathrm{met}}$	p-value
BTV _{per}	D ₉₈	54.9 Gy [52.1–56.0]	62.5 Gy [59.7-64.0]	58.7 Gy [56.0-60.8]	< 0.01
	Mean	60.4 Gy [58.7–61.2]	69.3 Gy [67.1–70.2]	67.2 Gy [60.1–69.2]	< 0.01
BTV _{met}	D_{98}	55.6 Gy [52.4–58.8]	58.7 Gy [57.7-63.6]	62.5 Gy [59.3-63.9]	< 0.01
	Mean	60.3 Gy [59.6-64.1]	68.7 Gy [66.2–71.7]	69.8 Gy [67.7-70.7]	0.05



Figure 3. The dose distribution in patient 3. BTV_{per} in dark green, BTV_{met} in purple, and PTV in blue. The 70 Gy isoshade is in red, and the 60 Gy isoshade is in green. Panel A shows an axial image where BTV_{per} is boosted, while BTV_{met} is boosted in the panel B.

be relevant for describing tumor progression. However, it is not immediately clear which parameters are the most relevant for the various tumor types, though vascular and metabolic information appears to be complementary [25]. During tumor development perfusion and metabolic activity is linked, as continuous tumor growth require an increased vascular density. However, the development of necrosis, vascular occlusion and/or hypoxia can result in a decoupling of the vascular state from the metabolic state, and may lead to a tumor environment with evolutionary selection of aggressive tumor cells through genetic instability and a selective microenvironment [26–28].

In the current work, K_1 and MR_{FDG} were used to assess perfusion and metabolism, respectively. Other surrogates could also have been chosen, such as the SUV at early and late time points, respectively, during dynamic PET acquisition [16]. However, we chose to use the pharmacokinetic parameters as these most likely provide the best separation between hyperperfused and hypermetabolic regions.

In the current work, we used Otsu's method to segment the images, and extract biologic targets of high intensity, by maximizing the intra-class variance while minimizing inter-class variance, resulting in regions of comparable size. As there was no direct measure available of neither metabolism nor vasculature, the actual discrepancy could not be measured, though the data indicate that there is partial overlap between different imaging phenotypes.

Some studies have investigated the overlap/correlation between biologic targets defined from multimodal or multiparametric imaging. In a study of DCEMRI and DWI of prostate cancer, using receiver operating characteristics (ROC) analysis for assessing overlap [29], it was found that the consistency between the derived parameter maps varied greatly between patients. Another study used FDG-PET, F-18-fluorothymidine (FLT; proliferation marker) PET, and Cu-61-diacetyl-bis(N4-methylthiosemicarbazone) (Cu-ATSM; hypoxia marker) PET to study biological heterogeneity in oropharyngeal tumors [30]. The FDG and FLT maps showed voxel-by-voxel correlations of typically 0.76, while Cu-ATSM showed slightly lower correlations with FDG and FLT. In our study, the cohort-based mean



Figure 4. Cohort-based mean dose volume histograms from the two boost regimes, either were BTV_{met} (A) or BTV_{per} (B) is boosted. PTV in black, BTV used for boosting in red, and corresponding non-boost BTV in green. The dotted lines indicate one standard deviation.

voxel-by-voxel correlation between K_1 and MR_{FDG} was 0.42 (data not shown), indicating that the heterogeneity in the currently used PET-derived maps were greater than for the multi-tracer PET study. Therefore, it is expected that smaller differences will be present in dose distributions for biologic targets defined from the multi-tracer PET analysis used in the referenced study.

The observed differences in target volume dose distributions were shown to depend on the degree of overlap between the regions and the clinical margins used. In general, boosting BTV_{met} produced an overall better coverage than boosting BTV_{per} , both in volume and dose differences. However, this was partly due to that BTV_{met} on average was larger than BTV_{per} , thus covering a larger part of the PTV. However, this is not necessarily positive with respect to treatment-related side effects, as a larger boost volume implies larger doses to organs at risk.

In the current work, 6 MV photons where used in the treatment planning. As the dose deposition kernels from such photons (and accompanying secondary charged particles) are rather wide, the resulting dose gradients in the tumor are relaxed. Thus, a region in close vicinity to the boost volume will also be partly boosted, as is the case in our work. The situation is different for proton therapy, where protons may show better conformity to small target volumes than photons [31]. However, if pathology (not visible in the functional images) is found outside the proton boost region, it may be preferable to use photons.

In conclusion, our work on multiparametric imaging and dose painting by contours revealed quite large interpatient variations in biologic targets and resulting dose distributions. For many of the patients, boosting one region will also cover the other region. However, for a subgroup of patients where the biological features are not overlapping, other approaches than single region boosting may be required. However, more knowledge from large patient cohorts is certainly needed before the clinical value is appraised and possible applications of multiparametric imaging may be considered.

Declaration of interest: The authors report no conflicts of interest. The authors alone are responsible for the content and writing of the paper.

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Supplementary material available online

Supplementary Table I.

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