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Spatiotemporal Stability of Cu-ATSM and FLT Positron Emission Tomography Distributions During Radiation Therapy

Tyler J. Bradshaw, MS^{*}, Stephen Yip, PhD^{*}, Ngoneh Jallow, MS^{*}, Lisa J. Forrest, VMD[‡], and Robert Jeraj, PhD^{*†}

^{*}Department of Medical Physics and University of Wisconsin, Madison, Wisconsin

[†]Department of Human Oncology, School of Medicine and Public Health, University of Wisconsin, Madison, Wisconsin

[‡]Department of Surgical Sciences, School of Veterinary Medicine, University of Wisconsin, Madison, Wisconsin

Abstract

Purpose—In dose painting, in which functional imaging is used to define biological targets for radiation therapy dose escalation, changes in spatial distributions of biological properties during treatment can compromise the quality of therapy. The goal of this study was to assess the spatiotemporal stability of 2 potential dose painting target—dihypoxia and proliferation—in canine tumors during radiation therapy.

Methods and Materials—Twenty-two canine patients with sinonasal tumors (14 carcinoma and 8 sarcoma) were imaged before hypofractionated radiation therapy with copper(II)-diacetyl-bis(N4-methylthiosemicarbazone) (Cu-ATSM) positron emission tomography/computed tomography (PET/CT) for hypoxia and 3'-deoxy-3'-¹⁸F-fluorothymidine (FLT) PET/CT for proliferation. The FLT scans were repeated after 2 fractions and the Cu-ATSM scans after 3 fractions. Midtreatment PET/CT images were deformably registered to pretreatment PET/CT images. Voxel-based Spearman correlation coefficients quantified the spatial stability of Cu-ATSM and FLT uptake distributions between pretreatment and midtreatment scans. Paired *t* tests determined significant differences between the patients' respective Cu-ATSM and FLT correlations coefficients. Standardized uptake value measures were also compared between pretreatment and midtreatment scans by use of paired *t* tests.

Results—Spatial distributions of Cu-ATSM and FLT uptake were stable through mid-treatment for both sarcomas and carcinomas: the population mean \pm standard deviation in Spearman correlation coefficient was 0.88 ± 0.07 for Cu-ATSM and 0.79 ± 0.13 for FLT. The patients' Cu-ATSM correlation coefficients were significantly higher than their respective FLT correlation coefficients ($P=.001$). Changes in Cu-ATSM SUV measures from pretreatment to midtreatment were histology dependent: carcinomas experienced significant decreases in Cu-ATSM uptake ($P<$

05), whereas sarcomas did not ($P > .20$). Both histologies experienced significant decreases in FLT uptake ($P < .05$).

Conclusions—Spatial distributions of Cu-ATSM were very stable after a few fractions of radiation therapy. FLT spatial distributions were generally stable early in therapy, although they were significantly less stable than Cu-ATSM distributions. Canine tumors had significantly lower proliferative activity at midtreatment than at pretreatment, and they experienced histology-dependent changes in Cu-ATSM uptake.

Introduction

After radiation therapy, most local recurrences occur at the primary tumor site—the region receiving the highest dose—indicating the need for higher tumor doses (1, 2). Increased doses, however, can also lead to increased normal tissue complications. Dose painting has been proposed as a potential solution to these conflicting objectives. In dose painting, a nonuniform dose is prescribed to a tumor volume, tailored to match the spatial heterogeneity of an underlying biological property linked with radioresistance (3), enabling higher doses to be delivered to tumor subregions of suspected recurrence while maintaining, or even lowering, normal tissue doses.

Several biological properties have been considered as potential dose painting targets. Tumor hypoxia is hypothesized to directly decrease the sensitivity of tumor cells to ionizing radiation (4) and is associated with poor clinical outcome (5). With the use of positron emission tomography (PET), hypoxia distributions can be imaged in vivo with radiotracers such as copper(II)-diacetylbis(N4-methylthiosemicarbazone) (Cu-ATSM) or ^{18}F -fluoromisonidazole (FMISO), among others (6). The resulting 3-dimensional (3D) images of hypoxia distributions can then guide prescribed dose distributions, as has been the subject of several feasibility studies (7-9).

There is concern, however, over the spatiotemporal stability of hypoxia maps, particularly in the context of dose painting (10, 11). Tumor hypoxia levels have been shown to fluctuate over time (12). If hypoxic locations change between pretreatment measurements and the beginning of therapy, or if hypoxic locations change during therapy, the quality of dose painting plans could greatly deteriorate: sensitive regions could be overdosed but resistant regions underdosed. To ensure that hypoxic regions are indeed preferentially targeted, hypoxia imaging would need to be repeated frequently and dose painting plans adjusted accordingly, which may not be technically or economically feasible. The spatial stability of hypoxia distributions has previously been investigated with the use of repeated FMISO PET scans, but with conflicting results: Roels et al (13), Nehmeh et al (14), and Lin et al (15) found poor spatial stability in serial FMISO images, whereas Okamoto et al (16) found very high spatial stability. Therefore, the stability of hypoxic targets before and during radiation therapy remains uncertain.

Cellular proliferation is another biological property that may be useful in guiding dose painting plans. Accelerated repopulation of surviving cancer cells during therapy has been shown to adversely influence treatment outcome (17). Furthermore, drastic changes in uptake of 3'-deoxy-3'- ^{18}F -fluorothymidine (FLT), a PET radiotracer for proliferation, have

been observed after the beginning of radiation therapy (18, 19). This has created interest in using changes in FLT uptake from pretherapy to midtherapy as a possible early response indicator (20, 21). Extending this idea to the voxel level, the change in a voxel's FLT uptake might be indicative of the voxel's response to therapy, so that maps of voxel responses might provide useful information for biological target definition. It is therefore important to characterize how proliferation distributions change in tumors during radiation therapy.

The goal of this study was twofold: to assess the spatial and absolute stability of Cu-ATSM PET during radiation therapy as a surrogate for tumor hypoxia stability, and to measure changes in FLT PET during radiation therapy as a surrogate for changes in cell proliferation. Canine veterinary patients with spontaneously occurring sinonasal tumors were used as subjects so that motion, setup, and image registration could be carefully controlled.

Methods and Materials

Patient population

The study included 22 canine veterinary patients referred “to the University of Wisconsin Veterinary Medical Teaching Hospital”. The research protocol was approved by the Animal Care and Use Committee of “the University of Wisconsin”, and all dog owners signed a written informed consent. The patients received diagnoses made with the use of computed tomography (CT), and they underwent biopsies for histopathologic evaluation. All dogs had nasal or paranasal sinus tumors of stage I to III (mean tumor volume, 55 cm³), with no evidence of distant metastases or intracranial extension. Histopathology revealed 13 adenocarcinoma, 7 chondrosarcoma, 1 squamous cell carcinoma, and 1 osteosarcoma. All imaging and treatment sessions were monitored by a veterinarian.

Imaging and treatment

The patients were treated with intensity modulated radiation therapy (IMRT) delivered by helical tomotherapy with curative intent. They were divided into 2 treatment groups as part of a larger study investigating dose escalation. The first group received the standard-of-care prescription: 4.2 Gy × 10 fractions to the planning target volume for a total of 42 Gy. The second group was prescribed 4.2 Gy × 10 fractions to the planning target volume with an additional boost of 0.8 Gy × 10 fractions to the gross tumor volume (GTV) plus a 2-mm margin.

The imaging and treatment schedules were as follows: Patients received pretreatment [⁶¹Cu]Cu-ATSM and [¹⁸F] FLT PET/CT scans 1 to 3 days before the beginning of therapy, with at least 1 day between scans. Patients received a second FLT scan after 2 fractions of IMRT (8.4 or 10 Gy) and a weekend break. Patients received a second Cu-ATSM scan after the third fraction of IMRT (12.6 or 15 Gy). Midtreatment FLT and Cu-ATSM scans were acquired on consecutive days. One patient did not receive a midtreatment FLT scan because of equipment failure. Patients were injected with 150 to 370 MBq of tracer. After injection, patients were kept in a kennel to limit physical activity. PET/CT scans were performed on a Discovery VCT (GE Healthcare, Waukesha, WI) PET/CT scanner. Patients were anesthetized during scans with an initial propofol bolus injection, and then isoflurane plus

100% oxygen. For Cu-ATSM scans, the time between tracer injection and the start of the PET scan was 3 hours; scans were 20-minute 3D static acquisitions over a single 15-cm bed position. FLT scans were 90-minute 3D dynamic acquisitions over a single 15-cm bed position. Emission data were attenuation corrected and reconstructed by the use of ordered subset expectation maximization (2 iterations, 35 subsets, and 3 mm postfiltering). The image grid was $256 \times 256 \times 47$ with $2.0 \times 2.0 \times 3.3$ mm³ voxel sizes. Voxel activity measurements were converted to standardized uptake values (SUVs) for analysis. For FLT scans, SUVs were calculated by averaging frames from 60 to 90 minutes. To achieve reproducible positioning across PET/CT scans and IMRT treatment sessions, patients' maxillae were positioned into custom dental molds that were affixed to the scanner and treatment couches, and patients' bodies were immobilized with vacuum mattresses (22).

Data analysis

We assessed the spatial stability of Cu-ATSM and FLT SUV distributions from pretreatment to midtreatment. GTVs were contoured by veterinarians using pretreatment CT images. For each patient, pretreatment and midtreatment CT images were deformably registered by use of the optical flow algorithm (23), and the resulting transformations were applied to their respective PET data. This allowed for voxel-based comparisons of pretreatment and midtreatment tracer distributions. To quantify tracer spatial stability, voxel-based Spearman rank correlation coefficients (ρ) were calculated for Cu-ATSM ($\rho_{\text{Cu-ATSM}}$) and FLT (ρ_{FLT}) within each patient's GTV, comparing pre-treatment and midtreatment voxel SUV distributions. Spearman correlation coefficients were preferred over Pearson correlation coefficients because SUV distributions were often not normally distributed. Population-averaged ρ were calculated with the Fisher transformation, and 2-sided paired t tests compared patients' $\rho_{\text{Cu-ATSM}}$ and ρ_{FLT} . We also investigated whether or not $\rho_{\text{Cu-ATSM}}$ and ρ_{FLT} were influenced by tumor volume, or by the level of uptake heterogeneity within the tumor, which we quantified using the standard deviation (SD) of the SUV distribution within the GTV (SUV_{SD}).

In addition to assessing voxel correlations, we quantified the degree of spatial overlap between the high-uptake regions at pretreatment and midtreatment. For each tumor, various thresholds were used to segment high-uptake regions. Thresholding was not performed as a percentage of the SUV_{max} , as is typically done in PET segmentation; this type of thresholding can result in extremely small volumes for some tumor distributions. Instead, we performed volume-based thresholding of the sorted SUV distribution, from 10% to 90%, in 10% increments. For instance, a 70% threshold would include 30% of the GTV's highest uptake voxels. For each threshold level, both pretreatment and midtreatment high-uptake volumes were segmented, and Dice coefficients quantified the degree of spatial overlap. The Dice coefficient is defined as the ratio of the intersection of 2 regions to the average volume of the 2 regions. After calculation of each patient's Dice coefficients for both tracers and all threshold levels, 2-sided paired t tests determined whether, across the population, Dice coefficients for Cu-ATSM were significantly different from Dice coefficients for FLT.

We investigated whether or not the magnitudes of Cu-ATSM and FLT uptake were significantly lower at mid-treatment than at pretreatment. The maximum voxel SUV in the

GTV (SUV_{max}), the mean SUV of a 1-cm³ sphere centered at SUV_{max} (SUV_{peak}), and the mean SUV of the GTV (SUV_{mean}) were calculated for pretreatment and midtreatment PET scans (midtreatment SUV measurements were assessed before deformable registration). Two-sided paired *t* tests determined whether changes in the SUV measures from pretreatment to midtreatment were statistically significant across the population, with a significance level of $P < .05$.

Results

A patient's pretreatment and midtreatment PET/CT images are shown in Figure 1. Voxel scatter plots comparing pre-treatment and midtreatment Cu-ATSM and FLT PET distributions are shown for 4 patients in Figure 2. The population-averaged $\rho_{Cu-ATSM}$ ($\pm SD$) was 0.88 ± 0.07 for all patients, 0.88 ± 0.08 for carcinoma patients, and 0.88 ± 0.07 for sarcoma patients. The population-averaged ρ_{FLT} was 0.79 ± 0.13 for all patients, 0.76 ± 0.14 for carcinoma patients, and 0.84 ± 0.12 for sarcoma patients. All patients' $\rho_{Cu-ATSM}$ and ρ_{FLT} are shown in Figure 3. Carcinoma and sarcoma populations were not significantly different in terms of $\rho_{Cu-ATSM}$ or ρ_{FLT} values. Using paired *t* tests, we found that patients' $\rho_{Cu-ATSM}$ were consistently higher than their respective ρ_{FLT} values ($P = .001$). We did not find significant correlations between tumor volume and $\rho_{Cu-ATSM}$ or ρ_{FLT} ($R \approx .2$), or between SUV_{SD} and $\rho_{Cu-ATSM}$ or ρ_{FLT} ($|R| < .2$), indicating that tracer stability was not likely affected by tumor volume or tumor heterogeneity.

Figure 4 shows the population-averaged Cu-ATSM and FLT Dice coefficients, quantifying the degree of overlap between pretreatment and midtreatment high-uptake regions, plotted as a function of segmentation method. For example, with the use of a 70% threshold (ie, the hottest 30% of the tumor volume), 77% of the pretreatment Cu-ATSM-avid regions, on average, overlapped with the mid-treatment Cu-ATSM-avid regions; for FLT and a 70% threshold, the average overlap was 69%. Paired *t* tests found that for a threshold of 30% and for thresholds greater than 50%, patients' Cu-ATSM Dice coefficients were consistently greater than their respective FLT Dice coefficients ($P < .01$).

The resulting pretreatment and midtreatment SUV_{mean} and SUV_{max} measurements are presented in Figure 5. On average, Cu-ATSM SUV_{mean} decreased by 18% at mid-treatment, or 1.2%/Gy when normalized by the prescribed dose. FLT SUV_{mean} decreased by 24% (2.5%/Gy) at mid-treatment. The average decreases in SUV_{max} were 9% (0.6%/Gy) for Cu-ATSM and 36% (3.8%/Gy) for FLT. Table 1 shows the results from paired *t* tests comparing pretreatment and midtreatment SUV measures. On average, patients had significantly lower FLT uptake after 2 fractions of hypofractionated IMRT regardless of histology. A significant reduction in Cu-ATSM uptake was observed for carcinoma patients, whereas sarcoma patients showed no significant changes in Cu-ATSM uptake. Dose level (42 Gy vs 50 Gy) had no significant effect on changes in Cu-ATSM and FLT SUV measures.

Discussion

Spatial stability of tracer distributions

We assessed the spatial stability of Cu-ATSM and FLT distributions during treatment in canine sinonasal tumors. These patients were ideal for this study for several reasons: anesthesia during imaging eliminated motion, given that motion blurring can cause 2 images to appear more similar than they really are (24); the bone surrounding the nasal cavity guided accurate image registration, because registration inaccuracies can significantly degrade voxel-based correlations (25); and precise immobilization techniques enabled repeatable positioning.

Overall, spatial distributions of Cu-ATSM uptake were very stable from pretreatment to midtreatment. Only 3 patients had $\rho_{\text{Cu-ATSM}}$ values less than 0.8. Even the highest uptake regions of the tumors (thresholds of 90%) had high average overlap (67%) between pretreatment and midtreatment. Therefore, in canine sinonasal tumors, Cu-ATSM dose-painting targets are spatially stable early during radiation therapy, so that neither multiple PET scans before therapy nor treatment adaptation based on additional PET imaging early in therapy are necessary. Our results are consistent with the results of Okamoto et al (16), who found average voxel correlations from FMISO test-retest scans to be 0.89 in patients with head and neck cancer. By contrast, Nehmeh et al (14) found the average voxel-based correlations from FMISO test-retest scans in patients with head and neck cancer to be 0.6 over the whole tumor volume, and 0.3 when only the most hypoxic subvolumes were considered.

On average, pretreatment and midtreatment FLT distributions were strongly correlated. This indicates that, for most patients, the most proliferative regions of the tumor generally remained the most proliferative regions early in therapy. This was surprising, considering that FLT uptake at midtreatment was often low (the average midtreatment SUV_{mean} was only 1.2), and one would expect correlations to deteriorate as tumor uptake approached background levels. Considerable interpatient heterogeneity in ρ_{FLT} , however, was observed, with 3 patients' ρ_{FLT} values lower than 0.6. In addition, patients' ρ_{FLT} were significantly lower than their respective $\rho_{\text{Cu-ATSM}}$, and FLT Dice coefficients were significantly lower than Cu-ATSM Dice coefficients for the high-uptake thresholds. Even the highest measured ρ_{FLT} (0.91) was only slightly better than the average $\rho_{\text{Cu-ATSM}}$ (0.88), and the highest FLT uptake regions (threshold of 90%) only had moderate overlap (49%) from pretreatment to midtreatment. This indicates that proliferative response to radiation is not spatially uniform throughout the tumor, and that the magnitude of intratumor heterogeneity in proliferative response appears to vary considerably across patients. Further investigation is warranted on whether intratumor proliferative response heterogeneity could provide useful information for defining dose painting targets.

Changes in tracer uptake magnitude

The reason for histology-dependent changes in Cu-ATSM SUV measures is uncertain. We have previously shown in this patient population that sarcoma tumors tend to have lower pretreatment uptake of Cu-ATSM and 2-deoxy-2-[^{18}F]fluoro-D-glucose (FDG) than

carcinomas (26), although it is not clear if or how this relates to changes in Cu-ATSM uptake during therapy. Other animal models have also demonstrated cell-line dependence in reoxygenation patterns (27). The reduction in Cu-ATSM uptake, however, may also be a consequence of cell kill. SUV values are affected not only by the biological property of interest (eg, hypoxia) but by the number of tumor cells available to take up the tracer (ie, cellular density). Consequently, decreases in SUV during therapy cannot be fully attributed to either biological changes or anatomic changes alone, but rather to some combination of both. On the basis of our analysis of megavoltage CTs acquired during therapy (data not shown), carcinoma tumors had a greater tendency to shrink during therapy than did sarcomas. This suggests that cell kill may have been a larger factor in carcinomas than in sarcomas, perhaps influencing the histology-specific changes in Cu-ATSM SUV.

We observed significant drops in FLT SUV measures for both histologies during treatment. Other studies have found similar results (18, 19). This reduction in proliferation could be interpreted as radiation-induced DNA damage checkpoint control (28), but it may also represent a drop in cell density. It is interesting to note that FLT uptake decreased significantly in both carcinoma and sarcoma tumors, whereas Cu-ATSM uptake decreased only in carcinoma tumors. If the lower tracer uptake were primarily caused by cell kill, one would have expected that, regardless of histology, if FLT uptake decreased significantly, then Cu-ATSM uptake would have likewise decreased. In addition, we would have expected that tumors with large changes in Cu-ATSM uptake would also experience similarly large changes in FLT uptake (ie, Cu-ATSM response would correlate with FLT response across patients). However, this was not observed: changes in FLT SUV were not correlated with changes in Cu-ATSM SUV ($R \approx .2$, data not shown). This suggests that changes in FLT uptake were driven primarily by functional response rather than by anatomic response.

Limitations

Several of the limitations of this study are addressed in our previous paper involving the same patient cohort (26). Limitations include potential influences from the use of anesthesia and 100% O₂ during imaging (29) (although patients were awake and breathing air during the 3-hour Cu-ATSM uptake period) and uncertainties in extrapolating results from canine tumors to human tumors (30). Perhaps the foremost limitation is Cu-ATSM's inconsistent specificity for hypoxia in different tumor types (31). Studies have found certain tumor types to have poor spatial correlations of Cu-ATSM and FMISO uptake patterns, especially with shorter Cu-ATSM uptake periods (32). We used a 3-hour uptake period for Cu-ATSM based on a preliminary analysis we performed in canine patients; we found that Cu-ATSM uptake at 24 hours after injection had higher spatial correlations to Cu-ATSM uptake at 3 hours after injection than at 1 hour after injection ($R \approx .85$; data not shown). Finally, we emphasize that although the tracer distributions were spatially stable early during treatment, tracer distributions at later midtreatment time points are likely to be less stable.

Conclusion

In conclusion, spatial distributions of Cu-ATSM and FLT PET remained mostly stable after a few fractions of IMRT, despite significant decreases in midtreatment SUV measures. Cu-ATSM spatial distributions were significantly more stable than were FLT distributions.

Acknowledgments

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Summary

There is concern that biological targets for dose painting can change locations over time, compromising the quality of dose painting plans. This study investigated the stability of copper(II)-diacetyl-bis(N4-methylthiosemicarbazone) (Cu-ATSM) and 3'-deoxy-3'-¹⁸F-fluorothymidine (FLT) positron emission tomography uptake distributions in canine sinonasal tumors during radiation therapy. We show that Cu-ATSM spatial distributions are very stable during radiation therapy. FLT distributions are generally stable but not as stable as Cu-ATSM distributions.

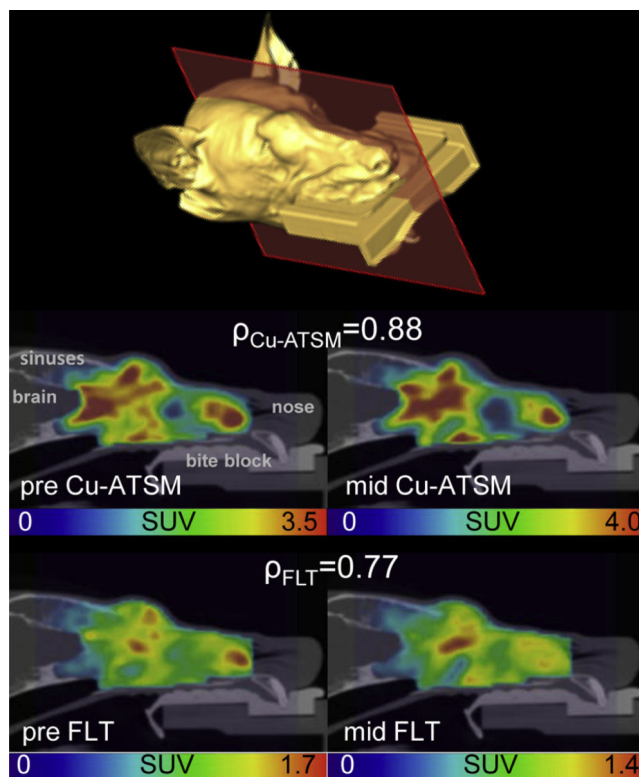


Fig. 1. Example sagittal positron emission tomography/ computed tomography slices are shown from a carcinoma patient's pretreatment (pre) and midtreatment (mid) copper(II)-diacetyl-bis(N4-methylthiosemicarbazone) (Cu-ATSM) (middle) and 3'-deoxy-3'-¹⁸F-fluorothymidine (FLT) (bottom) scans. An isosurface is shown for anatomic reference. This patient is the same as Patient 1 in Figure 2 and had $\rho_{\text{Cu-ATSM}}$ and ρ_{FLT} values similar to the population averages. SUV = standardized uptake value.

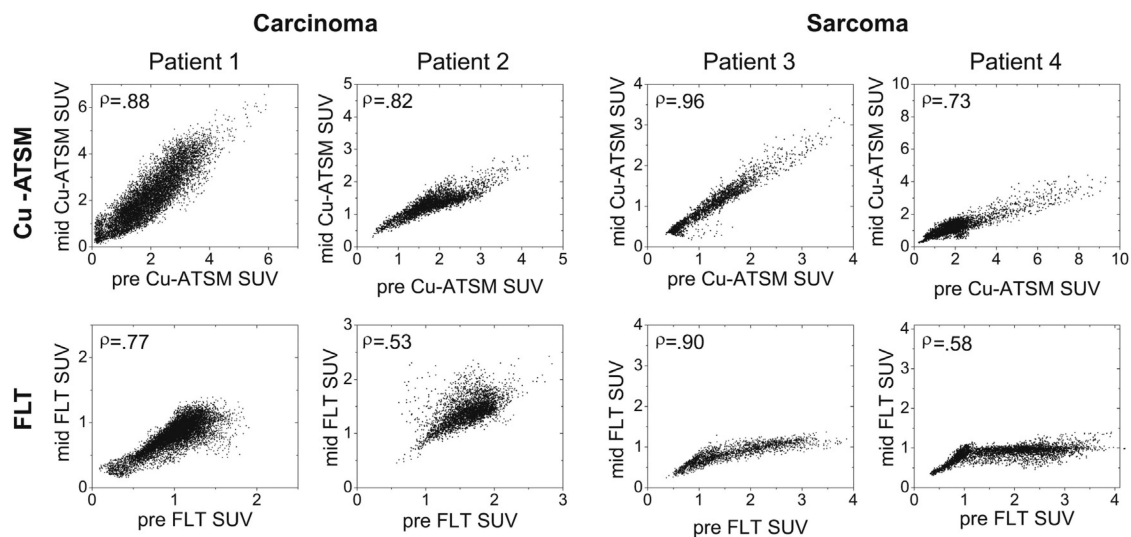


Fig. 2. Voxel-based scatter plots comparing pretreatment (pre) and midtreatment (mid) copper(II)-diacetyl-bis(N4-methylthiosemicarbazone) (Cu-ATSM) standardized uptake value (SUV) distributions (*top row*) and 3'-deoxy-3'-¹⁸F-fluorothymidine (FLT) SUV distributions (*bottom row*) for 4 different patients — 2 carcinomas and 2 sarcomas. Spearman rank correlation coefficients are shown in upper left corners.

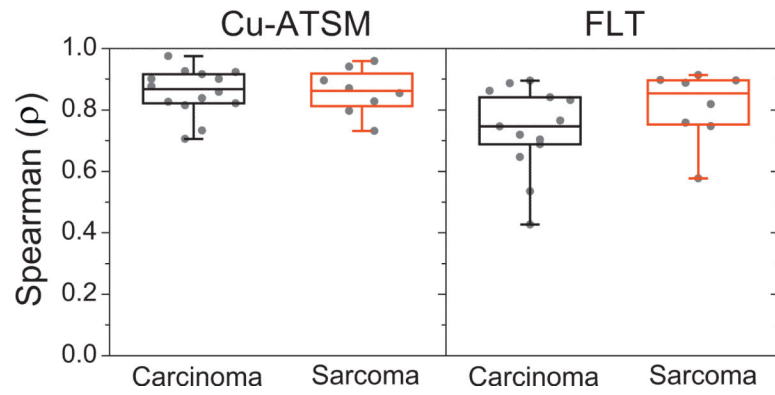


Fig. 3.

Quartile box plots of all patients' $\rho_{\text{Cu-ATSM}}$ (left) and ρ_{FLT} (right), representing the degree of tracer spatial stability through midtreatment, separated by histologic tumor type. Cu-ATSM = copper(II)-diacetyl-bis (N4-methylthiosemicarbazone); FLT = 3'-deoxy-3'- ^{18}F -fluorothymidine.

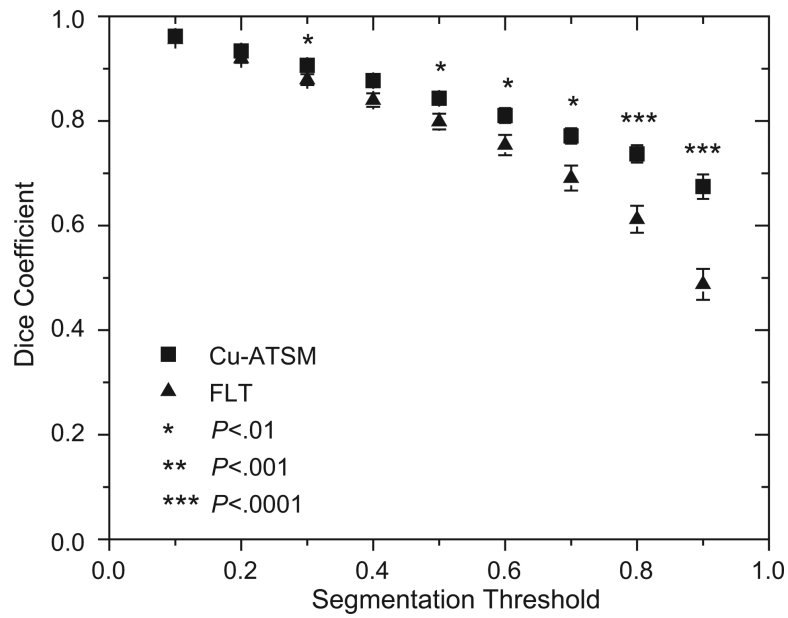


Fig. 4. Population-averaged Dice coefficients quantifying the overlap between high-uptake regions at pretreatment and midtreatment plotted as a function of segmentation method. Also shown are standard errors and results from paired *t* tests comparing patients' respective copper(II)-diacetyl-bis(N4-methylthiosemicarbazone) (Cu-ATSM) and 3'-deoxy-3'-¹⁸F-fluorothymidine (FLT) Dice coefficients.

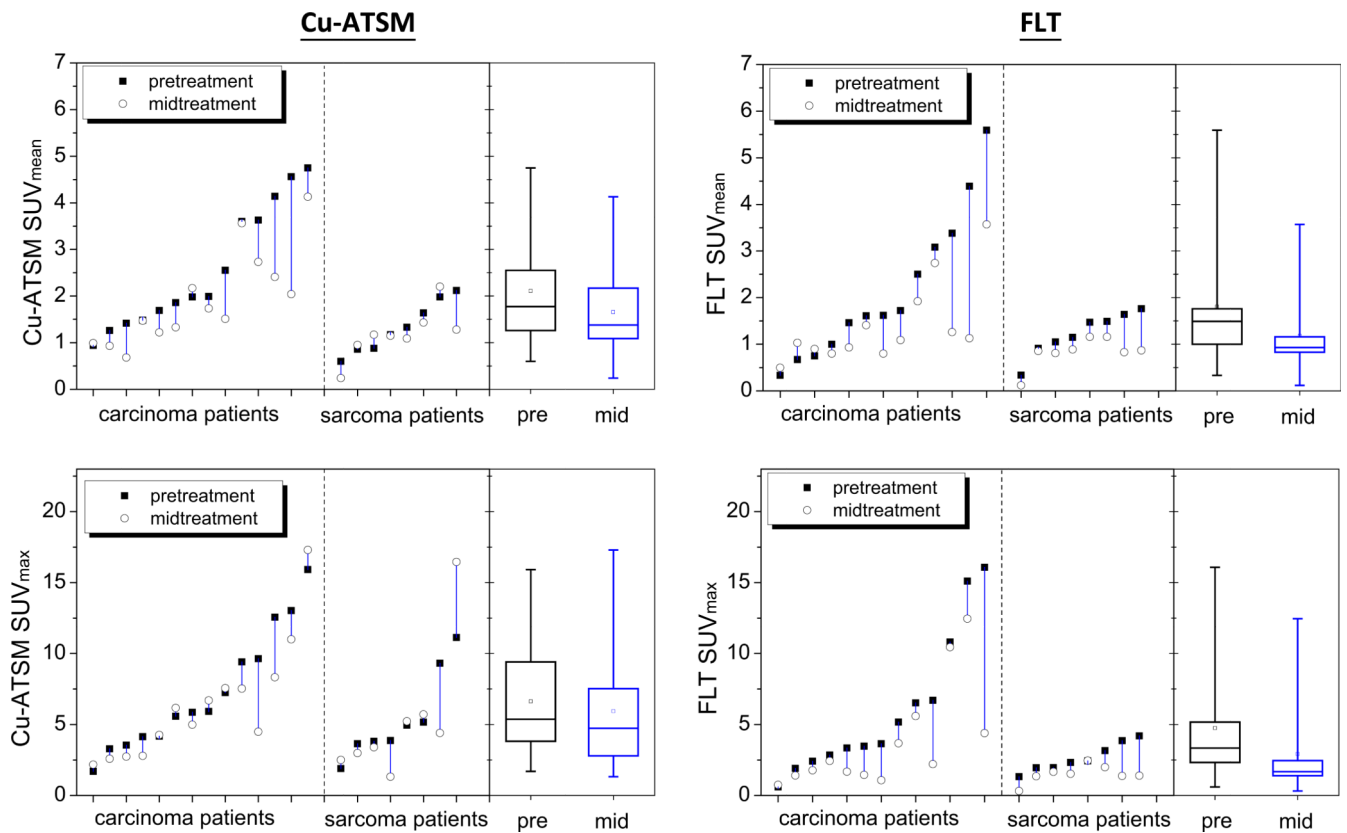


Fig. 5. Parallel line plots comparing each patient's pretreatment (solid square) and midtreatment (empty circle) standardized uptake value (SUV) measures for copper(II)-diacetyl-bis(N4-methylthiosemicarbazone) (Cu-ATSM) SUV_{mean} and SUV_{max} (left) and 3'-deoxy-3'- ^{18}F -fluorothymidine (FLT) SUV_{mean} and SUV_{max} (right), separated by histology. Also included are quartile box plots representing population distributions of respective SUV measures.

Table 1Results from paired *t* tests comparing patients' pretreatment and midtreatment SUV measures

Tracer	SUV measure	Carcinoma (n = 14)	Sarcoma (n = 8)	All patients (N = 22)
Cu-ATSM	SUV _{mean}	<i>P</i> =.006	<i>P</i> =.34	<i>P</i> =.004
	SUV _{max}	<i>P</i> =.08	<i>P</i> =.83	<i>P</i> =.17
	SUV _{peak}	<i>P</i> =.02	<i>P</i> =.81	<i>P</i> =.05
FLT	SUV _{mean}	<i>P</i> =.02	<i>P</i> =.008	<i>P</i> =.003
	SUV _{max}	<i>P</i> =.02	<i>P</i> =.01	<i>P</i> =.004
	SUV _{peak}	<i>P</i> =.04	<i>P</i> =.02	<i>P</i> =.01

Abbreviations: Cu-ATSM = copper(II)-diacetyl-bis(N4-methylthiosemicarbazone); FLT = 3'-deoxy-3'-¹⁸F-fluorothymidine; SUV = standardized uptake value.

Significant values (*P* ≤ .05; bolded) indicate significant decreases in SUV measures at midtreatment.