

SUV_{max-V} for assessing treatment response in FDG-PET Imaging of Patient-Derived Tumor Xenografts involving Triple-Negative Breast Cancer

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TO THE EDITOR:

In the preclinical arm of a co-clinical trial, Dr Savaikar et al. recently optimized ¹⁸F-FDG-PET imaging biomarkers of response to a combined docetaxel/carboplatin therapy in patient-derived tumor xenografts (PDX) involving triple-negative breast cancer (TNBC)(1). Twenty one necrotic-core phenotype tumors as well as 13 solid tumors were examined. Besides a preclinical μ PERCIST paradigm, 43 imaging metrics were evaluated, both in the whole tumor and in a single highest-intensity tumor slice. These metrics included (i) mean standard uptake values (SUV) obtained from various fixed percentages of the maximal-SUV thresholds (SUV_{Th}), and mean SUVs obtained from the voxels involved in a sphere centered at the maximal-SUV voxel (SUV_{Peak}). The following spherical volumes of 4 – 14 – 33mm³ (radius of 1 – 2 – 3voxels) were considered, leading to SUV_{P4} – SUV_{P14} – SUV_{P33}, respectively. In particular, Bland-Altman plots of test-retest data allowed us to estimate SUV₂₅ reproducibility (also called repeatability) percentage (R; 95%-confidence) of about 20 / 25% for solid / necrotic tumors (from Figure 3C / G, respectively). Finally, a coined quantitative response assessment score favored SUV₂₅ followed by SUV_{P14} as optimal metrics of response to therapy in PDX models.

We would like to stress the central role of R in assessing treatment response for any investigated SUV metrics, that is, the minimal relative change between two SUVs assessed from two successive examinations that is required to consider a significant difference (2). In this

connection, we suggest that a further SUV metrics, i.e., the $SUV_{\max-V}$, might be particularly suitable in the current context involving 21 tumors with a necrotic-core phenotype (and with varying tumor dimensions), thus exhibiting a low ^{18}F -FDG uptake at the core and well-separated ^{18}F -FDG-positive areas (Figure 2 by Savaikar et al.). Indeed, it has been previously shown, in lung cancer patients, that R of $SUV_{\max-N}$, which is an average SUV computed from the N hottest voxels regardless of their location within a ^{18}F -FDG-positive lesion, was significantly lower for $N = 30$ than that of SUV_{Peak} obtained from maximal SUV and its 26 neighboring voxels (3). In a subsequent study, $SUV_{\max-40}$ was found to more likely represent the most metabolically-active portions of tumors than SUV_{Peak} that was obtained from the voxels involved in a 1-mL sphere centered at the maximal-SUV voxel, with close R performance (4). Finally, the $SUV_{\max-N}$ procedure for treatment-response assessment has been described in a Takayasu-arteritis patient, emphasizing that the greater the N value, the lower the $SUV_{\max-N}$ R and, hence, the more efficient the metrics (Table 1 in (5)). Noteworthy, since the voxel volume may depend on the PET system, instead of the $SUV_{\max-N}$, one could alternatively use the $SUV_{\max-V}$ defined as an average SUV computed from an arbitrary total hottest volume (V), regardless of the location within the ^{18}F -FDG-positive lesion of the hottest voxels included in it. When comparing baseline / after-treatment scan, V should be set in the scan showing the lowest total ^{18}F -FDG-positive volume, but at the greatest possible value since the greater the V value, the lower the $SUV_{\max-V}$ R.

To conclude, Savaikar et al. addressed the important issue of reaching a consensus on reproducibility of imaging metrics for assessing response to therapy in oncology animal models (1). We suggest that the $SUV_{\max-V}$ metrics may have a place in this toolbox, with V set at the greatest possible value in the scan showing the lowest tumor uptake (that is expected to be the post-treatment one). Finally, in the current series, whether R of $SUV_{\max-14\text{mm}^3}$ or $SUV_{\max-33\text{mm}^3}$ might be lower than R of SUV_{25} , $SUV_{P14\text{mm}^3}$ and $SUV_{P33\text{mm}^3}$ remains to be assessed.

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