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How Advances in Imaging Will Affect Precision Radiation Oncology

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Summary
The development of new imaging methods, the advancement of quantitative and validated imaging signals, the adoption of highly automated pipelines for analysis, and the integration with the many other “panomic” cancer features are critical for our success.

Radiation oncology is 1 of the most structured disciplines in medicine. It is of a highly technical nature with reliance on robotic systems to deliver intervention, engagement of diverse expertise, and early adoption of digital approaches to optimize and execute the application of this highly effective cancer treatment. As a localized intervention, the dependence on sensitive, specific, and accurate imaging to define the extent of disease, its heterogeneity, and adjacency to normal tissues directly affects the therapeutic ratio. Image-based in vivo temporal monitoring of the response to treatment enables adaptation and further affects the therapeutic ratio. Thus, more precise intervention will enable fractionation schedules that better interoperate with advances such as immunotherapy. In the data set—rich era that promises precision and personalized medicine, the radiation oncology field will integrate these new data into highly protociled pathways of care that begin with multimodality prediction and enable patient-specific adaptation of therapy based on quantitative measures of the individual’s dose—volume temporal trajectory and midtherapy predictions of response. In addition to advancements in computed tomography imaging, emerging technologies, such as ultra-high-field magnetic resonance and molecular imaging will bring new information to the design of treatments. Next-generation image guided radiation therapy systems will inject high specificity and sensitivity data and stimulate adaptive replanning. In addition, a myriad

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Introduction

The advent of computed tomography (CT), magnetic resonance (MR) imaging, and positron emission tomography (PET) in the past 50 years has transformed the detection and treatment of cancer. These technologies have had a profound impact on our ability to better predict the outcomes of existing interventions and enabled a period of remarkable innovation, bringing new interventions and paradigms of care. The continued pace of advancement in imaging technology, combined with our growing ability to analyze the data from individual patients and aggregate these data across large patient cohorts, is creating a future in which care is becoming increasingly precise and personalized. Thus, the intervention is optimized according to well-defined patient-specific measurement regimens (eg, imaging, genomics, epigenomics) that have been derived from aggregated learning through large cohort data sets. The minimally invasive nature of imaging portrays its unique and critical role as an enabler of precision medicine in radiation oncology—whether that precision is in better defining the patient-specific extent of disease or being used in combination with many other inputs to design the time-dependent pattern of dose to be applied.

Although imaging has always played a central role in radiation oncology—as evidenced by most radiation oncology departments starting as divisions within the radiology department—imaging and image-derived signals will be even more integral to future practice. Image-derived features (eg, tumor size, location, imaging phenotype, texture) are now affecting nearly every facet of patient care and are used at every step (eg, diagnosis, daily treatment delivery, response assessment) in the process. Figure 1 illustrates the many image-related activities in the processes of patient characterization, decision making, care delivery, and follow-up phases of a patient’s journey through modern radiation therapy. Isolating the effect of advances in imaging is challenging, given the wide diversity of roles it plays. Furthermore, advances in 1 domain can enhance or diminish the importance of imaging in another. The development of MR-guided radiation therapy systems is an example in which consolidation of a diagnostic MR scanner with a linear accelerator might enable integrated delineation, adaptation, and response assessment from a single system (ie, daily MR imaging of the patient during treatment) (1). The schematic in Figure 1 also highlights 2 other important concepts—intent and information flow. With the emergence of rich new information during the course of therapy, the clinician must focus on clinical intent instead of “prescription.” Although previously considered “tampering” with evidence-based practice, the idea of adaptation within the course of therapy becomes a potential reality with the emergence of robust, noninvasive imaging signals, including anatomic and functional changes (2, 3). The field of radiation oncology has been a “first mover” in this area, with the development of adaptive radiation therapy enabled by image-guidance systems (4, 5). The effective and efficient flow of information (Fig. 1) is central to enabling this paradigm. Robust data exchange standards, gigabit data transfer speeds, on-demand computing capacity, artificial intelligence-based automation methods, and deformation technologies are enablers, and their scale up in the era of “big data” will open the door for greater consumption of imaging data in the field of radiation oncology (6). The success of the precision medicine paradigm requires that these diverse imaging data be drawn together with genomic data (including epigenetics), phenotypic data (eg, tumor microenvironment), emerging models of radiation biology (eg, radiogenomics, immunotherapy), and, of utmost importance, clinical outcomes.

In June 2016, the American Society of Radiation Oncology, National Cancer Institute, and American Association of Physicists in Medicine co-sponsored a 2-day workshop for radiation oncology physicians and medical physicists focused on precision medicine in radiation oncology. The workshop provided leaders the opportunity to interact with their peers and identify a new foundation of conducting research to bring the precision medicine paradigm to radiation oncology. Specific insights were sought regarding the future scientific direction of radiation oncology, the digital infrastructure needed to participate, and the practical and logistical barriers to enabling precision medicine in radiation oncology. Presentations and dialogue concentrated on genomics, imaging, big data, and real-world challenges to advance precision medicine in radiation oncology research, quality assurance, and safety. In addition to scientific posters and oral presentations from experts in the field, a dedicated workshop session for each
Multiple Roles of Imaging in the Multiple Timescales of Precision Cancer Medicine

Fig. 1. Multiple roles of imaging in the multiple timescales of personalizing radiation therapy. The practice of radiation oncology uses imaging data at every point in the process, including characterization of the patient and his or her disease, prediction of the outcome, design of the dose distribution, targeting and adaptation, real-time tracking, and response assessment. Image-based information informs the intent and enables patient-specific personalization of the treatment. In the absence of precision, this process will not build evidence nor increase our understanding of the disease or the effectiveness of our interventions. Pursuing the precision medicine paradigm requires us to improve our models of response and automate the flow of information.

Rapid Pace of Advances in Imaging—Measurement Is King

Molecular imaging approaches have long held the promise of augmenting the anatomic approach currently used for the design of highly conformal radiation dose distributions (7). The science of radiobiology, augmented by the ability of molecular imaging to assess the spatial heterogeneity of tumors, provided the basis for this approach, with decades of evidence highlighting the variation in the radiosensitivity of cells depending on the type, location in the cell cycle, and microenvironment (8). Without the ability to localize phenotypic heterogeneity, the entire tumor must be dosed to some minimum dose to ensure the most resistant disease is controlled. The effect of this approach on the adjacent normal tissues and the subsequent determination of patient-specific tolerance to a course of radiation therapy severely limited the development of more personalized radiation therapy regimens.

For the purposes of our report, focusing on the topic of hypoxia will be illustrative. The effect of very low oxygen levels on cell radiosensitivity has been established in basic science and is a well-documented negative predictive factor in the outcomes of a variety of cancers, including cancer of the cervix, head and neck, and prostate (9-12). Although most of these studies were performed using an invasive oxygen sensing Eppendorf probe, others have used PET imaging approaches to determine the magnitude and distribution of hypoxia in the tumor (13). In addition to giving valuable input into patient cohort classification, these results also highlight the potential to design hypoxia-escalating and normoxia de-escalating dose distributions (14, 15). Implicit in this approach is the assumption that these molecular patterns are accurate representations of the underlying biology and are stable throughout the course of radiation therapy (16, 17) (Fig. 2).

These exciting advances in molecular imaging of hypoxia raise many questions and define key areas of research for the inclusion of molecular imaging into the field of radiation oncology. Do we have the spatial resolution in our application of dose that is consistent with the imaging findings, and does the imaging modality have the ability to spatially resolve the underlying biology? (18). Do we consider these signals as absolute quantities, or do...
we consider time-dependent changes as more relevant in the fractionation paradigm of radiation therapy? (19). We are also becoming increasingly aware of the heterogeneity of the biology within tumors. Should we integrate other molecular or physiological signals—metabolism as measured through fludeoxyglucose PET or proliferation as measured through fluorothymidine PET—to pursue an even more accurate description of the underlying biology? (20). If we have an accurate description of the underlying biology, how do we derive the corresponding dose regimen and distribution? (21). Even the simple linkage between an image of hypoxia and the resulting dose distribution requires maturation. It is also important that the field consider the intrinsic uncertainties in the process of radiation therapy delivery when defining its objectives and design integrated solutions that are robust for clinical application.

The intrinsically localized nature of radiation therapy requires the field to be extremely attentive to sensitivity- and specificity-related advances in cancer imaging. More specifically, technologies that are directly “adjacent” to our current practice in terms of technology or care delivery can have immediate effects. Two exciting technologies that illustrate this point are highlighted in Figure 3 (22). First, the development of a highly specific form of prostate-specific membrane antigen that has the potential to affect the care of recurrent prostate cancer; and, second, the commercial development of 7T MR imaging systems (23, 24). In the case of novel prostate-specific membrane antigen agents, the potential to monitor and target recurrent disease is as important as the potential to use these agents to better delineate focal disease in the prostate gland for localized treatment. The potential effect of 7T imaging of the brain for detection, decision to treat, and localization of primary and metastatic lesions in the context of stereotactic radiosurgery is likely but will require investment to characterize the geometric accuracy at these field strengths.

**Image Validation—No More Pretty Pictures**

The ingestion of new imaging signals into the radiation therapy process is a significant challenge. Although the field was aggressive in adopting CT imaging for definition, delineation, and targeting, we have experienced significantly slower uptake of other modalities, including MR and molecular imaging. This is arguably in part owing to the challenges the field has had in ensuring the geometric
and signal accuracy of these new signals and the expense and availability of these modalities. In many ways, CT was easy to adopt owing to its intrinsic geometric and radiometric precision (ie, Hounsfield units). MR imaging provides exquisite sensitivity but has lacked in specificity and geometric integrity, thereby delaying its broad uptake in the planning process. PET imaging, in contrast, provides exquisite sensitivity and specificity but lacks broad availability of radiopharmaceuticals beyond fludeoxyglucose, thereby generally limiting it to academic centers. The field needs to invest in a more formal image ingestion process if these new imaging signals are going to have the effect on patient care that is possible (25). Such a formalization would bring bright minds and industry partners to bear on the current clinically validated problems. It would also provide a “systems thinking” framework to balance the tradeoffs between greater accuracy in delineation against greater precision in dose delivery.

Specific challenges that have been identified include (1) the need for greater transparency in image-signal processing to ensure quantitative performance of the imaging system and quantitative consistency between imaging systems of the same modality used in a trial; (2) addressing the problems of image distortion and registration; and (3) pathology-based validation of the image-derived signals. The development of the NIH’s Quantitative Imaging Network is an excellent example of the broader community’s awareness of the limiting nature these challenges have on the field of cancer (26).

In addition to quantification, validation of the imaging signals must also be aggressively pursued. A founder of modern medical education and practice, Sir William Osler stated it
clearly “as is our pathology, so goes our practice”—clearly, we must be cautious not to blindly “treat to the image.” Although the number of reports on pathology-based validation is small compared with those promoting a novel imaging technique, an increase has occurred in the report of pathology-imaging correlation studies enabled through 3-dimensional mapping technologies (26, 27). The recent report by Menard et al (28) on the correlation between multiparametric MR and biopsies of the prostate gland highlights the scale of the problem, because fused multiparametric MR imaging was inadequate to delineate the tumor boundaries for focal salvage in 10 of 19 patients. Furthermore, the use of a 5-mm expansion margin only reduced this cohort to 7 of 19 patients. This is exciting work because it highlights the field’s growing capabilities to quantify and validate image-based measurement of disease—a critical step toward image-enabled precision radiation oncology.

**Automation—Managing the 4 ‘V’s of Imaging Data**

The volume, variety, velocity, and veracity of imaging data speak to the scale of the current and future challenges to integrating imaging data into precision radiation oncology. Managing the volume and variety requires alternative approaches to traditional expert observer-based analysis. Radiation oncology has been forced to be a leader in the field of automated delineation to respond to the power of inverse planning for intensity modulated radiation therapy. The burden of delineating normal tissues and target volumes has created a variety of algorithms and software tools that are now in routine use in the clinical setting (29). Similarly, automated methods are being successfully applied to extract and validate features that are beyond the quantification skills of the expert. This is highlighted in the birth of the field of “radiomics.” Although the extraction of phenotype data from images is not novel, the extraction of features that are too subtle in their intensity or their frequency of occurrence for human experts to detect is now emerging as a predictive biomarker of response (30, 31). Although the clinical effect of this work is intriguing, the technical effect of this type of work needs to be recognized for its foreshadowing of cancer medicine and science. The process involved in radiomic analysis requires a massive scale-up in automation to manage the multitude of potentially relevant features, the need to integrate segmentation data, the number of image sets required to reach statistical significance, and the processes for regression analysis and machine learning. Further investments in automation will be required to allow these tools to contribute to clinical decision making, especially as time-dependent signatures become part of the equation. It is likely that robust clinical decision-making tools will need to integrate semantic, algorithm-derived, and machine learning—generated features with more traditional clinical information for maximum patient benefit. As illustrated in Figure 1, the volume, variety, and velocity of data flowing back from the intervention to the decision-making process will require automation. The veracity of the data refers to its quality and affects the certainty with which the data can be acted on—this is of critical importance given the life and death decisions being made in the design of a cancer patient’s treatment.

The adoption of automated “data mining” approaches within the real-world setting of data-rich clinical care will challenge the conventional paradigm of hypothesis-based scientific research. Collecting and analyzing data for patterns and correlations without a predefined mechanistic or conceptual hypothesis has been enabled, and the field of clinical science and the processes of regulatory approval will need to wrestle with the incongruence with their traditional approaches. This was a specific topic of discussion at the Workshop, and it was recognized that a large amount of work must be done on both sides of the debate.

**Open Science—Breaking Down the Silos**

The American Society of Clinical Oncology proposed the term “panomics” to describe the seemingly endless list of measurable quantities of potential importance in the study of cancer and the delivery of modern cancer care. The broad scope of panomics speaks to not only the common interests but also the common methods that the scientific and clinical communities should be using to analyze the data sets found in clinical and basic science of oncology. The creation of a community of researchers that see their data as “features,” regardless of their source—deep sequencing, 3-dimensional CT imaging data, dose-volume histograms, or clinical outcome data—is timely and powerful. Efforts by the National Institutes of Health to create powerful archives are ideal mechanisms to advance these efforts. The Cancer Imaging Archive and The Cancer Genomics Archive are linked with numerous site-specific research groups in bladder, breast, glioma, head and neck, ovarian, and renal cancers collaborating and reporting impactful research findings (32). Paradoxically, although it is work led by researchers in radiation oncology that are demonstrating the value of these “panomic” approaches (33, 34), a need exists for more radiation oncology data sets to be added to the archive. The field of radiation oncology needs to engage fully in the open science oncology community. Progress has been made in this regard through the medical physics archive on arXiv (available at: https://arxiv.org/list/physics.med-ph/recent), where a variety of data sets are available for further analysis. Although exciting, the amount of data is a tiny fraction of the volume of well-curated imaging and dosimetry data sets that sit within individual academic radiation oncology programs. These need to be annotated and drawn together with the corresponding pools of genomic sequencing data in national archives to allow this research to be more fruitful and more broadly effective.
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