Functional and molecular imaging in cancer drug development

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Abstract: Imaging biomarkers have a potential to identify key metabolic pathways that are up-regulated in cancer cells compared to normal cells. In early drug development, they can provide valuable information on the dissemination of the drug and estimate whether the drug reaches the target and, consequently, to determine the appropriate clinical benefit. The use of imaging as an early surrogate biomarker of response is also appealing, since it allows to tailor treatment regimens in individual patients. The aim of this review is to describe various imaging biomarkers covering most important cancer hallmarks such as cell death, proliferation, metabolism, vascularity, and hypoxia. We highlight the current status of using molecular imaging such as fluorodeoxyglucose (FDG), fluorothymidine (FLT), fluoromisonidazole (FMISO), and fluoroazomycin arabinoside (FAZA) positron emission tomography (PET) as well as advanced magnetic resonance imaging (MRI) techniques such as dynamic contrast enhancing (DCE) and diffusion weighted (DW)-MRI, and their potential roles in cancer drug development.

Keywords: Imaging biomarkers; molecular imaging (FDG-, FLT-, FMISO-, FAZA-PET); advanced magnetic resonance imaging (MRI) techniques (DCE-, DW-MRI); cancer drug development

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Introduction

Biomarkers are becoming an indispensable part of drug development. A biomarker is a representative that serves as an indicator of a patho-physiological process, or as a response to treatment which affects such a process (1). A more ideal use of imaging biomarkers for drug development can serve multi-purposes, such as disease staging, patient stratification, risk assessment, pharmacokinetics/ pharmacodynamics (PK/PD), drug safety and efficacy. The use of non-invasive imaging biomarkers to assess drug therapies has become more common during the last decades. From December 11, 1992, to July 1, 2010, the U.S. Food and Drug Administration (FDA) granted accelerated approval of 47 new indications for 35 anticancer drugs using surrogate endpoints, and most of them were objective response rate and progression-free survival (PFS), typically measured by magnetic resonance imaging (MRI) or computer tomography (CT) (2). However, vigorous debate

has challenged the use of anatomic assessments alone, as it may take two or three months to detect any shrinkage, thus only morphological information can be obtained. But, it may not be a suitable tool to assess response when agents targeting signaling pathways are involved, most notably in patients with gastrointestinal stromal tumor (GIST) treated by cytostatic targeted agents (3). To better understand tumor microenvironment (TME), and thereby select specific agents targeting metabolic key pathways, morphological information is not enough. Therefore, the addition of functional information on TME through imaging biomarkers would aid principle investigators to design personalized treatment planning by using specific targeted drugs (4).

Functional imaging of TME has several advantages: (I) it is a non-invasive procedure; (II) various sites of the tumors can be visualized and quantified simultaneously; (III) functional imaging using biomarkers can generate



Figure 1 Schematic presentation showing how imaging biomarkers identify overexpressed pathophysiological processes in cancer cells. IB, imaging biomarker; IRE's, imaging responsive elements; GLUT-1, glucose transporter-1; HIF-1, hypoxia-inducible factor-1; CC3, cleaved caspase 3; CD 31, platelet-endothelial cell adhesion molecule-1; VEGF, vascular endothelial growth factor; EPO, erythropoietin; FDG-PET, fluorodeoxyglucose-positron emission tomography; FMISO, FAZA-PET, fluoromisonidazole, fluoroazomycin arabinoside-PET; FLT-PET, fluorothymidine-PET; DW-MRI, diffusion-weighted-magnetic resonance imaging; DCE-MRI, dynamic contrast enhancing-MRI; ADC, apparent diffusion coefficient; SUV, standardized uptake value; Ktrans, volume transfer constant; PET, positron emission tomography; MRI, magnetic resonance imaging.

three dimensional images of the tumor which allows better quantification; (IV) moreover, functional imaging is capable of visualizing heterogeneous metabolic processes, such as glucose metabolism or tumor hypoxia, which are important contributors to tumor resistance and progression.

Molecular imaging using various labelled radioactive tracers such as fluorodeoxyglucose (FDG), fluoromisonidazole (FMISO), fluorothymidine (FLT), and functional imaging using advanced techniques such as dynamic contrast enhancing (DCE)-MRI and diffusion weighted (DW)-MRI, gain an increasing importance in cancer drug development (Figure 1). Quantitative measurements of imaging biomarkers compared to mere visual evaluation allow for more objective evaluation of disease, and more accurate monitoring through time (1). Therefore, the purposes of this review are (I) to summarize the basic principle and qualification of various imaging biomarkers; (II) to investigate key metabolic pathways up-regulated in cancer cells by using imaging biomarkers and to facilitate targeted cancer drug development; and (III) to describe pitfalls and recommendations when imaging biomarkers are

implemented in multicenter trials.

Imaging measurements and qualification

FDG (glucose metabolism)

The most frequently used positron emission tomography (PET) tracer in oncology is FDG for measuring glucose metabolism of the cell (5). However, FDG is not a substrate for metabolism in the glycolytic pathway. Therefore, the degree of trapped FDG uptake in the cells reflects the level of glucose metabolism and could be potentially used as imaging biomarker for early treatment response assessment in cancer patients (6). Maximum standardized uptake value (SUVmax) is a quantitative index to characterize FDG biomarker uptake, hence approximating the glucose metabolism; high SUVmax is associated with aggressive tumor metabolism and poor survival (7,8).

The transport of FDG, a glucose analogue, into cells is mediated by glucose transporters (GLUT-1 and 2) through the plasma membrane (9). Several published studies support significant positive correlation between FDG-PET uptake and the expression of GLUT examined by immunohistochemical staining (10-12). Primarily, the overexpression of GLUT characterizes enhanced tumor glucose metabolism and thereby increased FDG uptake is noticed on PET scan.

Demetri *et al.* (13) showed that, in all GIST patients with a response, the FDG-PET uptake in the tumor had decreased from baseline as early as 24 hours after a single dose of imatinib administration. In addition to that, in all patients, increased FDG-PET uptake from baseline is associated with disease progression. Also, FDG-PET uptake results were correlated with progression on CT or MRI.

Multiple studies have evaluated the role of FDG-PET and showed it promising in assessing response to treatment in solid tumors (14-16). However, the interpretation of SUV is not straightforward, with many factors affecting the values that can be derived. It was shown that a reliable drop in SUV, indicating a tumor response, is only seen in patients with high initial SUV (17). Caution should, therefore, be exercised when we interpret quantitative molecular imaging.

FAZA, FMISO (tumor bypoxia)

Tumor hypoxia is an important adverse prognostic factor and contributes to resistance for both chemotherapy and radiotherapy in several tumor types (18). Under hypoxic cell conditions, tumor hypoxia biomarkers undergo definite reductive metabolic pathways, resulting in reactive tumor metabolite markers which selectively bind to macromolecular cell components that can be detected by the PET signal, but which are washed out from normoxic cells (19,20).

FMISO was the first tracer tested clinically for tumor hypoxia, and it is still widely used (21-23). The novel hypoxia specific tracer, fluoroazomycin arabinoside (FAZA), has generated higher tumor-to-background ratios compared to FMISO in preclinical studies (24,25). FAZA also becomes a more attractive tracer for clinical use due to its more rapid clearance of unbound tracer from non-hypoxic tissues (24).

A clinically relevant exogenous hypoxic biomarker is pimonidazole. With this biomarker, high resolution image of hypoxia distribution at micro-regional level can be obtained using immunohistochemistry. The tumor hypoxia determined by pimonidazole binding assay is consistent with radiobiologically relevant hypoxic volume (26). Dubois *et al.* (27) found significant correlation between the hypoxic area derived from pimonidazole stained tumor section with the FMISO-PET defined hypoxic volume in an experimental rat tumor model (r=0.9066; P<0.0001).

FLT (tumor cell proliferation)

FLT was introduced by Shields *et al.* (28) as a PET proliferation imaging biomarker. FLT is monophosphorylated by thymidine kinase 1 (TK1), which leads to intracellular trapping. Since the concentration of TK1 is upregulated during the S phase of the cell cycle, the uptake of FLT reflects proliferation.

Tsuyoshi *et al.* (29) evaluated the effect of gemcitabinebased secondary chemotherapy with FLT- and FDG-PET imaging biomarkers in patients with stage IIIc recurrent ovarian cancer. FLT SUVmax decreased earlier than FDG SUVmax. Interestingly, FLT SUVmax correlated better with a reduction in size as measured by CT. Given the good imaging properties and strong correlation between functional imaging parameter (proliferation) FLT uptake and CT morphological parameters, SUVmax of FLT appears to be a promising biomarker for monitoring response to gemcitabine-based secondary chemotherapy treatment in recurrent ovarian cancer patients.

The rationale behind the FLT-PET uptake in tumors is based on TK1 activity and Ki-67 index dependence on proliferation. Since the concentration of TK1 and Ki-67 is overexpressed during the active proliferation phase of the cell cycle (S phase), the uptake of FLT is supposed to depend on TK1 and Ki-67 concentration. In a preclinical study, Rasey et al. (30) showed strong correlation between FLT and cell growth, TK1 activity and also with the percentage of cells in S phase of cell cycle (28,31). Recently, Yamamoto et al. (31) demonstrated a significant positive correlation between the proliferation index derived from Ki-67 immunohistochemistry with the FLT-PET uptake (r=0.81, P<0.01) in patients with newly diagnosed and recurrent gliomas (n=56). Given the strong correlation between the FLT uptake and TK1 and Ki-67, FLT appears to be a promising tracer for imaging proliferation.

DW-MRI (cell density)

DW-MRI is an advanced MR technique widely used for the detection and characterization of cancer as well as for monitoring the response to therapy. DW-MRI depends on the microscopic mobility of water in tissues, and it provides a unique imaging biomarker of water interaction with cellular, subcellular and macromolecular entities that impede free water movement (32). In oncologic imaging,

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DW-MRI has been used to evaluate tumor microstructure, e.g., cell membrane integrity and cellularity, which reflects lesion aggressiveness and tumor response. The acquisition of DW-MRI is non-invasive, does not require any exogenous contrast agents, does not use ionizing radiation, can be obtained relatively rapidly, and is easily incorporated into routine patient evaluation. The apparent diffusion coefficient (ADC) is the quantitative parameter of DW-MRI, and has been shown to be of high potential value for assessing treatment response (33,34). A low ADC reflects restricted diffusion and can be found in hypercellular tissues such as tumors, lymph nodes or in areas of fibrosis. A high ADC reflects less restriction of extracellular water motion and can be found in tissues with high glandular components or distinct necrosis. Cell kill due to efficient drug treatment leads to a loss of cell membrane integrity and reduction in tumor cell density with increase in the interstitial space, and hence it changes ADC measurement in the tumor tissue.

Foroutan *et al.* (34) evaluated the correlation between ADC and cell death in an osteosarcoma xenotransplant model at pre-treatment and at early time points following treatment. Pixel-by-pixel histograms were produced for each mouse prior to and following the treatment to quantify ADC. Cleaved caspase 3 (CC3) was used as an immunohistochemical marker to quantify cell death. Statistically significant differences in ADC maps were observed between control mice and treated mice, which demonstrates an increase in ADCs towards higher values in treated animals compared to controls. CC3 activity was also significantly higher in the treated animals compared to controls. Overall, a positive correlation was observed between increase in ADC values and cell death depicted by CC3 staining.

DCE imaging (blood flow and vascular permeability)

DCE imaging (MRI, CT and ultrasound) allows noninvasive quantification of TME and its vascular structure and function. The degree of DCE signal intensity reflects the pathophysiological factors, which include tissue perfusion and capillary permeability (35). Serial images are acquired dynamically before, during and after administration of a contrast agent: gadolinium for MRI, iodinated contrast for CT and microbubbles for ultrasound. The acquired data are fitted to mathematical models to obtain quantitative parameters through regions of interest. The volume transfer constant (Ktrans) is often used as a marker for the permeability of tumor vasculature. Other measures used are the rate constant Kep and the initial area under the gadolinium concentration curve (IAUGC).

Understanding the dynamics of tissue parameters is crucial for developing anti-angiogenic drugs. Vascular targeting agents such as bevacizumab or vandetanib are developed to reduce vascular permeability and promote tumor necrosis. Kummar *et al.* (35) investigated the effect of the anti-angiogenic drug vandetanib in patients with lymphomas. They observed a positive correlation between DCE-MRI parameters and plasma vascular endothelial growth factor (VEGF) levels. Similar results were reported by Donaldson *et al.* (36) who showed that tumors with poor permeability significantly correlated with the expression of plasma VEGF and the hypoxia marker pimonidazole. High expression of VEGF is associated with tumor angiogenesis and hypoxia, and thereby promotes tumor growth.

Imaging in cancer drug development

Stratifying patients

Molecular and functional imaging provides additional information on tumor characterization, which could help to "pre-select" and "enrich" a patient population. For example, in patients treated with gefitinib, a low baseline SUV of ¹⁸F-FDG has been shown to have prognostic value and to be associated with a higher response rate and a prolonged PFS (37).

Identification of tumor hypoxia could facilitate the use of hypoxia stimulated pro-drugs, which selectively kill hypoxic cells. Tirapazamine (TPZ) is such an example. The relatively limited benefit obtained in a trial reported by the CATAPULT I study group was likely due to poor patient stratification with inclusion of patients with betteroxygenated tumors (38). Recently, Rischin *et al.* (39) compared the cisplatin/5-FU *vs.* cisplatin/TPZ regimen in patients with head and neck squamous-cell cancer, in which FMISO-PET hypoxic imaging was used to stratify the tumors into hypoxic and non-hypoxic ones. The authors have shown that TPZ improved local tumor control in hypoxic but not in non-hypoxic tumors.

Imaging-guided therapy could promote personalizing treatment, for example by adjusting the treatment for nonresponders at an initial phase of treatment. Within the drug development, this sort of response monitoring could be used for selecting a homogeneous patient group for further studies by choosing only those patients who show early

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metabolic response. Several trials are currently investigating the use of FDG-PET/CT for early response-adapted therapy in lymphoma, with therapeutic stratification based on interim FDG-PET/CT results (40-42). The PETresponse-guided treatment has also been investigated in adenocarcinoma of the oesophagogastric junction, and the MUNICON phase II trial showed the feasibility of imaging-guided stratification by using the early metabolic response assessment from FDG-PET to clinical decision making in the treatment of solid tumors (40).

Verifying biological target engagement

The downstream effects of vascular endothelial growth factor receptor (VEGFR) inhibition on DCE-MRI have been documented in more than 30 phase I and II trials with a significant reduction in Ktrans and/or IAUGC being reported with multiple agents (43).

The more direct approach of using PET in cancer drug development is by labelling the drug itself. The anti-human epidermal growth factor receptor 2 (HER2) monoclonal antibody trastuzumab was used to treat breast cancer patients with HER2 expressing tumors and showed improved survival (44). Radionuclide labelled trastuzumab can visualize the affinity of the targeted agent in vivo, which allows us to collect vital information about the pharmacokinetic properties of the drug such as injected dose versus accumulated drug concentration in the organs and its regional bio-distribution. In this case, the use of radionuclide imaging may overcome problems associated with biopsies, including sampling errors and discordance of expression between primary tumors and metastases. Moreover, the drug uptake by the target tissue can be quantified at sequential imaging scans, and it might give us insight into drug's action at the target tissue and its association with the tumor response.

Defining dose setting

In phase I trials, dose-escalation is usually undertaken to define the maximum tolerated dose (MTD), under the assumption that the most pronounced changes are likely to be detected at the highest dose. But, target saturation may already be reached at lower dose levels. Through direct visualization of target inhibition, imaging changes are likely to be apparent at lower doses than the MTD, and imaging may be used in choosing the optimal biological dose. In a study of brivanib, a dual VEGFR and fibroblast growth factor receptor (FGFR) tyrosine kinase inhibitor, Jonker et al. (45) evaluated DCE-MRI responses in several dose schedules in selected patients, known to respond to anti-VEGFR therapies, and then selected the optimal schedule for a phase II trial. Despite this experience, imaging is not commonly used for selecting dose or schedule, and such data are limited, so the use of imaging to determine the optimal schedule of a targeted agent or to monitor drug activity has to be further explored for cancer drug development.

Novel surrogate endpoint for early evaluation of drug activity

A growing understanding of the underlying molecular pathways active in cancer has led to the development of novel therapies targeting VEGFR, EGFR, phosphoinositide 3-kinase (PI3K), mammalian target of rapamycin (mTOR), protein kinase B (Akt) and other pathways. Unlike the cytotoxic chemotherapy, many of these molecular targeted agents are cytostatic, causing inhibition of tumor growth rather than tumor regression. In this context, using tumor shrinkage as a surrogate endpoint may not be the most adequate mean to measure therapeutic response, as the response rates only based on change of tumor size are low, despite a high percentage of patients having prolonged stable disease and sometimes even improvements in survival. Therefore, functional imaging provides a unique potential opportunity to assess antitumoral activity at early stage.

Many have stimulated the FDA to accept novel surrogate endpoints, such as novel imaging endpoints that can be measured earlier than tumor shrinkage and are likely to predict clinical benefit. A qualified biomarker accepted by the FDA as a surrogate endpoint needs to match several important criteria: (I) the endpoint must have an accepted, standardized definition; (II) data from multiple clinical studies must demonstrate a strong correlation of the surrogate endpoint with clinical outcome; (III) wellpowered prospective studies must have been performed to validate the surrogate endpoint (i.e., truly predictive of clinical benefit with meaningful improvement in patient outcome) (46). The strength of evidence will vary, depending on whether the surrogate is intended for use in accelerated approval or definite regulatory approval.

FDG uptake (SUV) has been proposed as an appropriate novel surrogate endpoint for early evaluation of drug activity in clinical trials. There have been many retrospective and some prospective studies in a variety of cancer types that have demonstrated a promising correlation

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between SUV decrease and survival (41,47). To date, these studies have been primarily performed in single institutions with small numbers of patients. To our knowledge, there are two ongoing multicenter trials prospectively designed to validate FDG-PET as a surrogate endpoint in lymphoma (CALGR-53030) and non-small cell lung cancer (RTOG-0235/ACRIN6668). Large prospective multi-center clinical trials are needed to assess the degree of correlation by comparing a pre-defined threshold in SUV change to clinical outcome.

Imaging in multicenter clinical trials

Standardization

Although many imaging biomarkers have been described for cancer research, few of them are widely considered adequate to provide unambiguous assessment of response, and enough for making decisions to stop or continue drug development processes. Implementing molecular and functional imaging to assess response requires that an observed change of the imaging biomarker due to treatments must be greater than the intrinsic and extrinsic variability of the biomarker in the absence of treatment. High reproducibility of molecular and functional imaging techniques relies on good quality data and standardized procedures. Standardization is the first and crucial step when imaging is implemented in multicenter trials. In this context, the EORTC-PET study group issued recommendations for the measurement of [¹⁸F]FDG uptake in monitoring treatment response in 1999 (48). These recommendations included suggestions for patient preparation, pre-therapy and post-therapy imaging delays, and techniques for measuring SUV. Following that, guidelines of the National Cancer Institute (NCI) and the European Association of Nuclear Medicine (EANM) for tumor PET imaging enriched the standardized procedures (49,50), making it more feasible to include PET in large multicenter trials. Regarding advanced MRI techniques, such as DCE, the techniques are relatively simple but require strict protocols, careful acquisition, accurate dosing of contrast agent and suitable selection of injection rate, image timing, and image analysis for quantification. In US, the Quantitative Imaging Biomarker Alliance (QIBA) DCE-MRI technical committee provided guidelines and defined basic standards for DCE-MRI measurement and quality control that enable consistent, reliable and fitfor-purpose quantitative measurements when DCE MRI is implemented in multicenter trials (51). In Europe, the

Quantitative Imaging in Oncology: Connecting Cellular Processes to Therapy (QuIC-ConCePT) consortium was created and resourced by the Innovative Medicines Initiative (IMI), Europe's largest public-private initiative (4). It aims to qualify three specific imaging biomarkers of tumor cell proliferation, apoptosis, and necrosis, to allow drug developers to demonstrate reliably the modulation of these pathologic processes in tumors of patients in future trials (4). The precompetitive research and public-private partnerships may reduce the duplication, and develop imaging biomarkers in a most robust, consistent and costeffective way, so as to accelerate drug development.

Recommendation

Providing a benchmark, based on a set of common principles of implementing functional and molecular imaging in multicenter trials, is important to facilitate exchange of data, promote quality, accelerate research and reduce attrition rate for drug developers. In addition to the summary on the utility of imaging biomarkers based on literature review, we provide general recommendations for principal investigators designing and conducting multicenter clinical trials that include functional and moleculare imaging biomarkers (*Table 1*).

Conclusions

In the past decade, advances in biology and genomics have led to the development of targeted agents against cancer. This paradigm shift emphasizes the need for specific imaging biomarkers to identify key metabolic changes within the TME and thereby selecting a specific drug of choice. Non-invasive in vivo imaging offers unique, sensitive and clinically transformable information for cancer drug development, notably via efficient patient selection, imaging-guided therapeutic stratification, verification of biological target modulation and dose adaptation. In addition, functional and molecular imaging may potentially allow us to depict accurate changes in tumors, particularly before anatomic changes are evident, and to predict long-term clinical benefit. However, large prospective multicenter studies are needed to further qualify and validate the potential functional imaging biomarkers by demonstrating a strong correlation with clinical outcome. When imaging is implemented in multicenter clinical trials, we highly recommend designing studies with sound methodology and conducting studies

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Protocol design		Before site activation		During accrual		After accrual
• Early engagement with experts	•	Development of imaging	•	Ensuring of imaging data	•	Closure of database;
from relevant disciplines;		guidelines;		protection;	•	Data analysis
Rational discussion on why the	•	Requirement of Dummy run, i.e.,	•	Compliance of all electronic		according protocol
selected imaging biomarker		test before accruing patient;		processes;		and guidelines;
is appropriate, including	•	Evidence of scanner calibration	•	Quality assurance and	•	Exploration of
feasibility cost-effectiveness;		(e.g., scanner accreditation);		quality control of imaging		potential routes to
Preliminary imaging biomarker	•	Elaboration of standard		data;		integrate imaging
quantification data (e.g.,		operating procedures, quality	•	Appropriate data		biomarker in
reproducibility, accuracy);		assurance & quality control		management and tracking;		future clincial
Understanding of biological		program;	•	Documentation of all		trials, or usage in
mechanism;	•	Organization of imaging central		processes;		clinical routine,
Selection of appropriate		review (e.g., review panel,	•	Interim analysis to reassess		and of necessary
criteria;		procedures, and turn-around		feasibility of approach and		methodological
Early definition of statistical		time);		whether statistical power will		improvements
power calculation with	•	Evidence of proper site training		still be reached (optional)		
simulation and adaption						

Table 1 Recommendations when imaging is integrated in multicenter clinical trials

with adequate standardization of data acquisition and analysis techniques.

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