

Quantitative
Imaging
Biomarkers
Alliance



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Profile: DCE MRI Quantification

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Version 1.0

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I. Executive Summary

The RSNA QIBA Dynamic Contrast Enhanced Magnetic Resonance Imaging (DCE-MRI) Technical Committee is composed of scientists representing the imaging device manufacturers, image analysis laboratories, biopharmaceutical industry, academia, government research organizations, and professional societies, among others. All work is classified as pre-competitive. The goal of the DCE-MRI committee is to define basic standards for DCE-MRI measurements and quality control that enable consistent, reliable and fit-for-purpose quantitative transfer constant (K^{trans})¹ and blood normalized initial area under the gadolinium concentration curve (IAUGC_{BN})² results [across imaging platforms (at 1.5Tesla), clinical sites, and time] .

This effort is motivated by the emergence of DCE-MRI as a method with potential to provide predictive, prognostic and/or pharmacodynamic response biomarkers for cancer³⁻¹¹. Remarkably, the results demonstrating this potential have been obtained despite considerable variation in the methods used for acquisition and analysis of the DCE-MRI data. This suggests there are substantial physiological differences (i.e., benign vs. malignant or non-responsive vs. responsive tumors) underlying these observations. Thus, there appears to be a promising future for use of DCE-MRI for both clinical research and in routine clinical practice. However, in order to fulfill this promise it is essential that common quantitative endpoints are used and that results are independent of imaging platforms, clinical sites, and time.

For the application of DCE-MRI in the development of anti-angiogenic and anti-vascular therapies, there is a consensus¹² on which quantitative endpoints should be employed: K^{trans} and IAUGC_{BN} . Hence, the initial focus of the DCE-MRI committee is on these biomarkers. Although there have been general recommendations on how to standardize DCE-MRI methodology^{12, 13}, there are no guidelines sufficient to ensure consistent, reliable and fit-for-purpose quantitative DCE-MRI results across imaging platforms, clinical sites, and time. Hence, in this profile, basic standards for site and scanner qualification, subject preparation, contrast agent administration, imaging procedure, image post-processing, image analysis, image interpretation, data archival and quality control are defined to provide that guidance.

Summary of Clinical Trial Usage

This technique offers a robust, reproducible measure of microvascular parameters associated with human cancers based on kinetic modeling of dynamic MRI data sets. The rigor and details surrounding these data are described throughout the text of this document in various sub-sections.

II. Clinical Context and Claims

One application of DCE-MRI where considerable effort has been focused on quantitative endpoints is its use to provide pharmacodynamic biomarkers for the development of novel therapeutic (in specific anti-angiogenic) agents targeting the tumor blood supply^{4, 9, 14-25}. A growing understanding of the underlying molecular pathways active in cancer has led to the development of novel therapies targeting VEGFR, EGFR-tk, PI3K, mTOR, Akt and other pathways. Unlike the conventional cytotoxic chemotherapeutic agents, many of these molecularly-targeted agents are cytostatic, causing inhibition of tumor growth rather than tumor regression. One example is anti-angiogenesis agents, which are presumed to act through altering tumor vasculature and reducing tumor blood flow and/or permeability. In this context, conventional endpoints, like tumor shrinkage as applied at e.g. Response Evaluation Criteria in Solid Tumors (RECIST), may not be the most effective means to measure therapeutic responses. Other functional MR imaging acquisition and

73 analysis applications (e.g. BOLD, R_2^* perfusion) yield several important candidate imaging biomarkers that
74 can predict and monitor targeted treatment response and can document pharmacodynamic response.
75 However, these are not within the scope of this document. DCE-MRI represents an MRI-based method to
76 assess the tumor microvascular environment by tracking the kinetics of a low-molecular weight contrast
77 agent intravenously administered to patients.

78 The emerging importance of angiogenesis as a cancer therapy target makes assays of vascularity important
79 to clinical research and future clinical practice related to targeted cancer therapy. There are multiple
80 literature reports of the application of DCE-MRI to predict and detect changes associated with angiogenesis
81 targeted therapy^{4, 9, 15, 17, 19, 20, 24, 25}. Further, there is interest in the application of quantitative DCE-MRI to
82 characterize enhancing lesions as malignant in several organ systems, including breast and prostate.

83 In this context, K^{trans} and IAUGC_{BN} can provide evidence of the desired physiologic impact of these agents in
84 Phase 1 clinical trials. For some agents, e.g., VEGFR-targeted agents, evidence of substantially reduced K^{trans}
85 and IAUGC_{BN} is necessary, but not sufficient, for a significant reduction in tumor size^{16, 17}. For other
86 agents, e.g., vascular-targeted agents, evidence of a substantial vascular effect may not be associated with
87 a reduction in tumor size⁹, but is still essential for effective combination with other anti-cancer agents. In
88 either case, lack of a substantial vascular effect indicates a more potent agent is needed, while evidence for
89 a substantial vascular effect indicates further development is appropriate.

90 Utilities and Endpoints for Clinical Trials

91 DCE-MRI is currently not the standard of care in many centers conducting clinical trials in oncology. Since
92 these centers often do not have expertise in DCE-MRI and more than one center is typically involved,
93 therefore effort and precision are required ensure consistent, reliable and fit-for-purpose quantitative DCE-
94 MRI results. Hence, the guidelines provided in this profile will ensure that not only are the relative changes
95 induced by treatment are informative, but that absolute changes can be compared across these studies.

96 Claim:

97 **Quantitative microvascular properties, specifically transfer constant (K^{trans}) and blood normalized initial**
98 **area under the gadolinium concentration curve (IAUGC_{BN}), can be measured from DCE-MRI data obtained**
99 **at 1.5T using low molecular weight extracellular gadolinium-based contrast agents within a 20% test-**
100 **retest coefficient of variation for solid tumors at least 2 cm in diameter.**

101
102 Profile specified for use with: **patients with malignancy**, for the following indicated biology: **primary or**
103 **metastatic**, and to serve the following purpose: **therapeutic response**.

104 III. Profile Details

105 1. Subject Handling

106 1.1 Subject Scheduling

107 *Subject Selection Criteria related to Imaging*

- 108 • Local policies for contraindications for absolute MRI safety should be followed;
109 definition of relative and/or absolute contraindications to MRI are not within the scope of
110 this document.

- 111
- 112 • Lesions that are selected for DCE-MRI analysis should not be within 10 cm of metal
- 113 prostheses, e.g., spinal hardware, hip prostheses, metallic surgical staples, etc.
- 114
- 115 • Patient selection criteria may be guided by the Eastern Cooperative Oncology Group
- 116 (ECOG) status (See Appendix 2) for full description of ECOG performance status). In specific,
- 117 patients meeting ECOG status ≥ 2 will not be eligible for participation in the study because,
- 118 historically, this patient profile has shown poor ability to meet the demands of the
- 119 examination.
- 120
- 121 • The QIBA DCE-MRI committee acknowledges that there are potential and relative
- 122 contraindications to MRI in patients suffering from claustrophobia. Methods for minimizing
- 123 anxiety and/or discomfort are at the discretion of the physician caring for the patient.
- 124
- 125 • The QIBA DCE-MRI committee acknowledges that there are potential risks associated
- 126 with the use of gadolinium-based contrast media. The default recommendations for
- 127 intravenous contrast that follow assume there are no known contraindications in a particular
- 128 patient other than the possibility of an allergic reaction to the gadolinium contrast agent.
- 129 The committee assumes that local standards for good clinical practices (GCP) will be
- 130 substituted for the default in cases where there are known risks.
- 131
- 132 • Recent FDA guidelines
- 133 (<http://www.fda.gov/Drugs/DrugSafety/ucm223966.htm#approved>), outline the safety
- 134 concerns associated with using gadolinium based contrast agents in patients with impaired
- 135 renal function. The DCE-MRI committee echoes these recommendations and advises
- 136 reference to these standards when choosing patients in order to determine eligibility for
- 137 entry into a DCE-MRI clinical trial.
- 138
- 139 • Patients will not be eligible if they have received ANY gadolinium based contrast
- 140 agent within 24 hrs.
- 141

142 **1.1.1. Timing of Imaging Tests within the Treatment Calendar**

143 The DCE-MRI Technical Committee believes that all baseline evaluations should be ideally be within 14 but
144 at least within 30 days prior to treatment start. Otherwise the resulting functional tumor characterization
145 may not reflect the status of the tumor prior to initiation of therapy. The interval between follow up scans
146 within patients may be determined by current standards for GCP or the rationale driving a clinical trial of a
147 new treatment

148 **1.1.2. Timing Relative to confounding Activities (to minimize “impact”)**

149 DCE-MRI examinations should not be performed within 14 days after biopsy.

150 **1.2. Subject Preparation**

151 There are no specific patient preparation procedures for the MRI scans described in this protocol. There
152 are specifications for other procedures that might be acquired contemporaneously, such as requirements
153 for fasting prior to FDG PET scans or the administration of oral contrast for abdominal CT. Those timing
154 procedures may be followed as indicated without adverse impact on these guidelines

155 **1.2.1. Prior to Arrival**

156 The local standard of care for acquiring MRI scans may be followed. For example, patients may be advised
157 to wear comfortable clothing, leave jewelry at home, etc.

158 **1.2.2. Upon Arrival**

159 Staff shall prepare the patient according to the local standard of care, (including e.g. removal of all metal
160 objects and electronic devices). Patients should be comfortably positioned, in appropriate attire to
161 minimize patient motion and stress (which might affect the imaging results) and any unnecessary patient
162 discomfort.

163 **1.2.3 Preparation for Exam**

164 Beyond a clear, simple language description of the image acquisition procedure, no exam preparation is
165 specified beyond the local standard of care for MRI with contrast.

166 **1.3. Imaging-related Substance Preparation and Administration**

167 **1.3.1. Substance Description and Purpose**

168 The literature, which supports the claim, is based on the utilization of an extracellular gadolinium based
169 contrast agent. Although it is known that there is a small degree of protein binding associated with many
170 commercially available extracellular gadolinium contrast agents,²⁶ these are comparable amongst the
171 various vendors. Contrast agents with fundamentally different degrees of protein binding, (e.g.,
172 Gadobenate and Gadofosveset) are not addressed by this profile. The committee therefore recommends
173 using a classical extracellular based gadolinium based contrast agent.

174 **1.3.2. Dose Calculation and/or Schedule**

175 Total contrast agent dose depending on body weight and renal function:

- 176
- 177 • Before DCE-MRI the patient's renal creatine clearance should be obtained, and estimated
178 glomerular filtration rate (eGFR) determined through well-known and adopted formulas.²⁷
 - 179 • Routine concentration of the Gadolinium contrast agent should be 0.1 mmol/kg.
 - 180 • The decision whether to administer total contrast dosage will be based on GCP and the policies
181 adopted at the institution performing the examination. However, the same body weight adapted contrast
182 agent concentration should be used for repeat studies, and in case of an acute renal insufficiency and/or
183 failure at follow-up a later imaging time point or patient exclusion should be discussed.

184 **1.3.3. Timing, Subject Activity Level, and Factors Relevant to Initiation of Image Data Acquisition**

185 Contrast injection should occur after the following imaging sequences have been acquired (See Section 6):

- 186 • Anatomic imaging for localizing tumors
- 187 • Variable flip angle imaging for native tissue (pre-gadolinium injection) T_1 map calculation

188 Contrast injection should occur after at least 5 baseline acquisitions from the imaging volume have been
189 acquired.

190 **1.3.4. Administration Route**

191 Each subject should have an intravenous catheter (ideally no smaller than 20 gauge), which should be
192 ideally placed in the right antecubital fossa. Injection through a port-a-catheter or permanent indwelling
193 catheter is not recommended. What is critical is that the same injection site and catheter size be used for
194 repeat studies, if at all possible.

195 **1.3.5. Rate, Delay and Related Parameters / Apparatus**

196 Contrast agent and saline flush should be administered in a dynamic fashion with an MR-compatible power
197 injector.

- 198 • At baseline and at each subsequent time-point in any longitudinal study, the same dose of contrast and
199 rate of contrast administration should be performed.
- 200 • The rate of administration should be rapid enough to ensure adequate first-pass bolus arterial
201 concentration of the contrast agent (generally 2-4 ml/sec)
- 202 • The contrast agent should be flushed with between 20 to 30 ml of normal saline injected at the same
203 rate as the contrast agent.

204 **1.3.6. Required Visualization / Monitoring, if any**

205 No particular visualization or monitoring is specified beyond the local standard of care for MRI with
206 contrast.

208 **2. Imaging Procedure**

209 This section describes the imaging protocols and procedure for conducting a DCE-MRI exam. Suitable
210 localizer (scout) images must be collected at the start of exam and used to confirm correct coil placement
211 as well as selection of appropriate region to image. This will be followed by routine non-contrast agent-
212 enhanced sequences to delineate the number, location, and limits of tumor extension. Exact protocols for
213 these imaging sequences may be determined by the local imaging norms, e.g:

- 214 • **Localizer**
- 215 • **Anatomic sequences T_1 , T_2 weighted imaging**
- 216 • **Variable Flip angle (VFA) T_1 weighted imaging (T_1 mapping)**
- 217 • **3D Gradient echo volumetric imaging (dynamic imaging)**
- 218 • **Anatomic, post-contrast T_1 weighted sequences**

219 **2.1. Required Characteristics of Resulting Data**

220 The DCE-MRI portion of the exam will consist of two components, both acquired using the same 3D fast
 221 spoiled gradient recalled echo sequence, or equivalent, and scan locations:

- 222
- 223 (a) A variable flip angle series, for pre-contrast agent native tissue T₁ mapping.
- 224 • Ensure TR and TE values stay constant for all flip angles,
 - 225 • Ensure that the machine gain settings are not reset automatically (using automated pre-scan
 226 features) between each flip angle acquisition so that system gain settings are identical for each flip angle
 227 acquisition.
 - 228 • Flip angles: The range of numbers of flip angles supported in the literature varies from 2-7.
 - 229 • Number of signal averages (NSA or NEX) ≥ 2.
- 230
- 231

232 (b). DCE-MRI Protocol: Pulse Sequence:

- 233 • **Pulse Sequence:** 3D fast spoiled gradient recalled echo or equivalent
- 234 • **Coils:** Transmit: Body coil; Receive: Body coil or phased array receive coil
 235 No parallel imaging options
 236 No magnetization preparation schemes

237 **Imaging plane** - The acquisition plane should include the lesion of interest and a feeding vessel with in-
 238 plane flow.

239 **Frequency encoding direction:** The frequency encoding direction should be adjusted so as to minimize
 240 motion artifact. This decision will be based on the location of the tumor being interrogated and its
 241 relationship to moving structures.

242

Parameter	Compliance Levels	
TE	Acceptable	2.0-2.5ms
	Target	1.5-2.0ms
	Ideal	<1.5ms
TR	Acceptable	5-7ms
	Target	3-5ms
	Ideal	< 3ms

243

244 **Temporal resolution:** The temporal resolution should be less than 10 sec.

245

246 **Flip angles:** Smaller flip angles will lead to potential saturation of the signal intensity vs. gadolinium
 247 concentration, particularly in vessels. Note should be made that SAR limits may affect the maximum
 248 allowable flip angle. Operators should use the maximal allowed flip angle when SAR limitations occur. Flip
 249 angles ranging from 25-35 degrees are recommended in order to minimize saturation effects and to avoid
 250 specific absorption rate (SAR) problems.

251

252 **Receiver Bandwidth:** Greater or equal to ±31.25 kHz (or ~250 Hz/pixel)

253

254 **Field of View (FOV) and Partial Fourier (“fractional echo” and/or reduced phase-encoding FOV) as needed**
255 **to meet temporal resolution requirements**

256
257 **Number of Slices:** Acceptable: ≥ 10 prior to zero fill. Ideal: as many as possible while maintaining ideal
258 temporal resolution.

259
260 **Slice thickness:** *Ideal:* <5 mm, *Target:* 5.1-6 mm, *Acceptable:* 6.1-8 mm

261
262 **Matrix:** 256 x 160 (before applying rectangular FOV) – in order to meet 1-2mm in-plane spatial resolution

263
264 **Number of acquisitions (phases):** Sufficient to allow acquisition of at least 5 minutes of post injection data
265 plus at least 5 phases acquired before contrast agent injection (baseline images).

266
267 **Digitized bit depth:** The maximum dynamic range should be utilized, e.g., “extended dynamic range” or
268 equivalent.

269 **2.1.1. Data Content**

270 All imaging data should be stored in DICOM format.

271 **2.1.2. Data Structure**

272 All VFA data should be clearly labeled as individual series, one per flip angle, or contained in a single series
273 with the data order clearly defined.

274
275 All DCE-MRI data should be contained in a single series.

276 **2.1.3. Data Quality**

277 A quality review, confirming that all imaging parameters were correct, data structure is correct, etc., before
278 the data are submitted for analysis.

279 **2.2. Imaging Data Acquisition**

280 **2.2.1. Subject Positioning**

281 **Patient and coil positioning:**

282
283 • When the general location of the target tumor(s) is known prior to DCE-MRI, for example glioma or
284 local breast cancer evaluation, the patient set up for the MRI should be based on standard operating
285 procedures for patient positioning and coil placement for clinical MRI of that body part taking into account
286 the total scan time (see below).

287
288 • When the subject under investigation may have uncertain tumor location(s), as is common in the
289 setting of patients undergoing therapy for metastatic disease, it will often be necessary for the DCE-MRI

study to be planned with reference to the most recent pre-DCE-MRI imaging (often a CT study). From this study, tumor burden and location should be assessed. Optimally, review of actual imaging by a radiologist involved in the DCE-MRI study planning should be made. At times, if such images are not available for direct review, review of imaging reports (CT, PET) detailing extent of disease is mandatory, both to confirm eligibility (presence of at least one “imageable” target lesion) and to identify the preferred anatomic regions for DCE-MRI (chest, abdomen, pelvis, extremity). Review of prior diagnostic imaging may also be helpful to confirm cystic or necrotic nature of certain lesions, assessments which may be challenging at the time of DCE-MRI planning based solely on T₁- and/or T₂-weighted image sets. When multiple potential target lesions are available, the location of the most suitable lesion(s) should be noted. The most suitable lesion will depend on size, location relative to areas of pulsatile or respiratory artifacts, and presence or absence of necrosis or cystic areas.

- DCE-MRI subject should be placed appropriately in the scanner in order to best image the lesion of interest (e.g. supine for head/neck/thorax/abdomen/pelvis and prone within a breast coil for breast studies).

- When patient condition allows, placement of the arms over the head may avoid undesirable wrap artifact for temporally optimized 3D spoiled gradient echo sequences used for chest and abdomen lesions. However, these positions often cannot be sustained by patients without excessive discomfort. In such cases, arms placed anteriorly over the chest or at the sides may be preferable. For larger patients, side-down arm positioning may require adjustment of the DCE-MRI imaging FOV to avoid undesirable wrap artifact. Appropriate coil placement per area of examination (head, neck, breast, extremity) is then done. For lesions in the chest, abdomen, or pelvis, a torso array coil is then placed in the area of target lesion(s). Ideally, both anterior and posterior coils are centered over the expected target lesion location.

- Tumor size and location on longitudinal studies should be considered in the design of an analysis scheme. Recall, that the claims of this profile are only applicable to lesions greater than or equal to 2cm. If the lesion is large in proportion to the volume imaged by DCE-MRI, precautions should be taken to maximize the possibility that the same portion of the lesion will imaged on longitudinal studies. In general, this requires careful scan location set up on follow-up studies in order to match the same anatomic positions imaged in target organs on earlier studies (e.g. by saving of the planning screen shot). However, because of differences in patient angulation on follow-up studies the same anatomic locations may not be imaged on each study. In this case, an analysis scheme that discards image data from locations that are not included in the imaged volume (after end slice elimination) of all relevant studies is favored. This can be accomplished by registration of images obtained from the dynamic sequences of all studies (for example, images obtained by averaging all dynamic images obtained at the same location) to high-resolution anatomic images obtained (for example) at the initial time point.

- Tumors that are predominantly solid without significant necrosis or cystic characteristics would be considered the ideal choice of tumor for analysis. Tumors with extensive hemorrhage, or completely cystic or necrotic lesions are considered non-ideal and should be excluded from consideration.

- Tumor locations should be chosen to minimize the effects of excessive respiratory or pulsatile motion. Ideally, these would include the soft tissues of the extremities, posterior chest wall, retroperitoneum and abdomen. Although areas with some respiratory motion (e.g. kidneys, adrenal glands, retroperitoneum, lateral chest wall, pancreas, lung apices, neck) are considered acceptable, lesions

336 within the hila, pericardium and lateral segment of the left lobe of the liver are not ideal because of their
337 significant compromise secondary to respiratory motion.

338 **2.2.2. Instructions to Subject During Acquisition**

339 The patient will be instructed to perform slow, steady breathing during the examination.

340 **2.2.3. Timing/Triggers**

341 All examinations will be performed in slow free breathing state. Timing parameters for the bolus injection
342 of contrast agent will occur after the acquisition of no less than 5 baseline volume scans.

343 **2.2.4. Model-specific Parameters**

344 Appendix G.1 lists acquisition parameter values for specific models/versions that can be expected to
345 produce data meeting the requirements of Section 7.1.

346 **2.3. Imaging Data Reconstruction**

347 All imaging data reconstruction will be performed per vendor specification and will involve Fourier
348 transformation of Cartesian data. No user-selected smoothing or other post-processing will be performed
349 so as to insure the integrity of the data for image analysis.

350 **2.3.1. Platform-specific Instructions**

351 Appendix G.2 lists reconstruction parameter values for specific models/versions that can be expected to
352 produce data meeting the requirements of Section 7.2.

353 **3. Image Post-processing**

354 There are no specific image post-processing requirements in this profile. No user-selected post-processing
355 filters or image normalization methods should be used prior to data analysis as described in the next steps.
356 If phased-array receiver coils are used, image combination and reconstruction should be according to
357 standard manufacturer algorithms.

358 **4. Parametric image formation**

359 Analysis of DCE-MRI data is carried out in a series of distinct steps:

360

- 361 • Generate a native tissue T_1 map using the VFA data.
- 362 • When required, apply time-series motion correction to the dynamic data.
- 363 • Convert DCE-MRI signal intensity data, $SI(t)$, to gadolinium concentration ($[Gd](t)$).
- 364 • Calculate a vascular input function.
- 365 • Identify the region or regions of interest in the dynamic data.
- 366 • Calculate the DCE-MRI imaging biomarker parameters, K^{trans} and $IAUGC_{BN}$.

367 Each of these steps is addressed in detail below.

368 **4.1. Input Data to Be Used**

369 Processed magnitude images will be utilized for image analysis for input into the steps described in the
370 following sections

371 **4.2. Methods to Be Used**

372 **Generate a T_1 Map**

373 The intent of this step is to provide a complete map of pre-contrast T_1 values for the imaged slab. These
374 values will then be used in the signal formation model based conversion of changes in signal intensity to
375 gadolinium concentration. The slice locations, orientation and resolution of these images should be
376 prescribed identically to the dynamic series, and this series should be acquired immediately prior to the
377 dynamic series. The output of this step is an image of T_1 values which can be co-registered to the dynamic
378 series and used in subsequent calculations. The T_1 values at each voxel location are calculated as follows
379 [1]:

- 380 1. Create a vector x containing the signal intensity at each flip angle divided by the tangent of the
381 flip angle.
- 382 2. Create a vector y containing the signal intensity at each flip angle divided by the sine of the flip
383 angle.
- 384 3. For the n acquired flip angles create a set of points $(x_0, y_0) \dots (x_n, y_n)$.
- 385 4. Fit a line with slope s to the set of points defined in Step 3.
- 386 5. $T_1 = -TR/\ln(s)$.

387
388 The use of non-linear curve fitting methods (for example, simplex or Levenberg-Marquard techniques) to
389 extract T_1 from the signal intensities theoretically may be more robust to noise than the linearized solution
390 presented above. Non-linear techniques may be used if they are validated using test images to perform no
391 worse for than the solution above in the expected range of T_1 , equilibrium magnetization and noise of
392 tumors and vessels to be imaged.

394 **Apply Motion Correction to the Dynamic Data**

395 The intent of this step is to correct for patient motion that occurs between acquired phases of the dynamic
396 data due to respiration, swallowing, and other involuntary movements. This step is not intended to correct
397 ghosting artifacts that can occur along the phase encoding direction within a particular image due to
398 patient motion during acquisition. These artifacts are more or less intractable unless the motion is regular
399 and easily modeled, and are best addressed by adjusting the phase/frequency encoding scheme to
400 minimize their impact on structures of interest. In general, simple rigid shift or affine transform based
401 registration methods will not be adequate for this step, due to the fact that the movement in question is
402 typically limited to specific regions within the image – for example, the liver in a coronal scan of the
403 abdomen may move substantially with respiration while the bulk of the body remains relatively motionless.
404 Fully deformable registration methods based on optical flow may provide good results in some cases^{28, 29}.
405 However, these methods will frequently fail for the phases immediately surrounding the contrast injection.
406 Semi-automated registration in which a user identifies the target tumor and only information drawn from
407 that region is used to generate phase to phase shifts provides an alternative approach. This allows rigid
408 shift methods using mutual information³⁰, which tend to be more robust than optical flow methods, to be
409 employed. Finally, registration may be carried out manually or using simple shift registration techniques²¹.
410 Data corrupted with motion must be either corrected prior to analysis or discarded for subsequent
411 pharmacokinetic analysis.

412

Convert SI(t) in the Dynamic Data to [Gd](t)

The intent of this step is to convert the arbitrary signal intensity units in the dynamic data into units of gadolinium concentration. This step should be applied after the regions of interest for analysis have been defined, but prior to the calculation of vascular parameters. Two methods for accomplishing this are defined below.

Method A: Conversion Using a Signal Formation Model Gadolinium concentration at each image pixel is given by (eq 1):

$$C(t) = \left(\frac{1}{T_1(t)} - \frac{1}{T_{10}} \right) / R_{Gd} \quad \text{Eq. 1}$$

Here T_{10} is the pre-contrast T_1 at that pixel, obtained as described above, and R_{Gd} is the relaxivity of Gd (obtained from contrast agent manufacturer's specifications).

$T_1(t)$ can be derived from the SPGR imaging equation (neglecting T_2^* effects) and is given by the following expressions (eqs 2-4): Let

$$E_{10} = \exp(-TR/T_{10}) \quad \text{Eq. 2}$$

$$B = \frac{1 - E_{10}}{1 - \cos \alpha * E_{10}} \quad \text{Eq. 3}$$

$$A = B * SI(t) / SI(0) \quad \text{Eq. 4}$$

where α is the flip angle, TR is the repetition time, and SI(t) and SI(0) are the signal intensities at time t and pre-contrast baseline respectively in the DCEMRI sequence (eq 5). Then,

$$\frac{1}{T_1(t)} = \frac{-1}{TR} * \ln \left[\frac{1 - A}{1 - \cos \alpha * A} \right] \quad \text{Eq. 5}$$

Method B: Conversion Using a Look-Up Table

This method is motivated by the concern that inaccuracies in T_1 mapping and/or co-registration of initial T_1 values to the dynamic data may introduce excessive variability into the final calculated parameters. If this method is used, it is not necessary to acquire the T_1 mapping data described above. This method assumes a high degree of response uniformity, and so may be limited in cases where phased array coils are used. In general it is recommended to use the inherent body coil for both transmit and receive when using this method. It should also be noted that the use of this method will introduce a uniform bias in the estimation of quantitative parameters which will impact absolute measurements, but will not affect quantification of change, for example from one exam to another. This method has been shown to yield better test-retest reproducibility than T_1 -based quantification method.¹⁴

This method requires that a phantom containing a range of concentrations of gadolinium and a range of baseline T_1 values (generally obtained via different concentrations of copper sulfate or a similar compound) is scanned using the dynamic protocol on each scanner that will be used for the study. Data from these phantoms can then be used to construct a look-up table relating baseline T_1 , signal delta, and gadolinium concentration. In order to create this look-up table, a linear correlation is performed between the difference of signal intensity between that in a phantom concentration sample and a sample with no gadolinium concentration (used as x-axis values) and the nominal R_1 ($1/T_1$) of the concentration sample.

452 The resulting slope m then be used to estimate Gd concentration C using the equation $C = m * [SI(t) - SI(0)]$,
 453 where $SI(t)$ is the signal intensity in the dynamic data for a given time point t , and $SI(0)$ is the signal intensity
 454 in the same location at baseline (before contrast agent injection).

456 Calculate a Vascular input Function

457
 458 The intent of this step is to generate an accurate, patient-specific vascular input function to serve as an
 459 input to the vascular model. One way to accomplish this is to have an analyst draw a manual ROI within an
 460 artery, and use the mean enhancement curve within that ROI as the subject-specific VIF, as described by
 461 Vonken et al.³¹. It has been demonstrated previously that this method has significant variability associated
 462 with it³², due primarily to the spatially- and temporally-varying flow artifacts found in major arteries. A
 463 better option is to make use of an automated search technique to generate a locally optimal VIF. Several
 464 methods of accomplishing this have been described previously^{33,34,35}.

465
 466 Although not as intensely scrutinized, population-averaged vascular input functions have shown promise in
 467 some studies, and are currently under investigation.³⁶

468
 469 The signal for the vascular input function can then be converted into concentration using either Method A
 470 or B as described above.

471
 472 In some cases, data driven vascular input functions may be difficult to measure accurately due to anatomy,
 473 motion, flow effects, and T_2^* effects. In these situations, alternative methods of using population averaged
 474 vascular input functions, or reference tissue based vascular input functions may be used. These methods in
 475 general lead to poorer characterization of subject-specific physiology and lead to poorer reproducibility.

477 Calculate the Vascular Parameters

478 The intent of this step is to generate the parameter set which will be used to characterize the tissues of
 479 interest. Parameters will be calculated based on the standard Tofts model³⁴, which is derived from the Kety
 480 equations³⁷. The vascular bed is modeled as a linear system, such that (eq 6):

$$481 \quad C_t(t) = C_p(t) \otimes h(t) \quad \text{Eq. 6}$$

482 with impulse response $h(t)$ given by (eq 7):

$$483 \quad h(t) = K^{trans} * \exp(-k_{ep}t) \quad \text{Eq. 7}$$

484 where K^{trans} is the volume rate constant between blood plasma and extra-cellular extra-vascular space (EES)
 485 and k_{ep} is the rate constant between the EES and blood plasma. Given the tissue uptake curve $C_t(t)$ and the
 486 VIF $C_p(t)$, K^{trans} and k_{ep} are estimated using a gradient-descent energy minimization scheme, by using
 487 already established Levenberg-Marquardt or Minpack-1 curve fitting algorithms, both of which require
 488 adequate baseline sampling³⁸. Delay correction should be performed to shift the VIF curve to match the
 489 arrival time of the tumor curve prior to curve fitting.

490 A full parameter set will be calculated for each voxel within the defined tumor boundaries. Parameters may
 491 be reported out either as mean and median values per tumor or as histograms.

492 The blood normalized IAUGC_{BN} is measured from the area under the concentration curve up to 90 seconds
 493 post injection, normalized by dividing the area under the vascular input function curve also up to 90
 494 seconds post injection.

495 4.4. Platform-specific Instructions

496 Appendix G.4 lists image analysis parameter values for specific models/versions that can be expected to
497 produce data meeting the requirements of Section 9.

498 **5. Parametric image analysis**

499 Derivation of quantitative parameters characterizing the response associated with a lesion of interest from
500 parameter maps is a multistep process, most, if not all, of which are being studied by on-going research.
501 There are several choices that can be made at any of these steps, and the effect of these choices on the
502 validity of results and variability of parametric maps has not yet been fully characterized.

503

504 When multi-institutional trials are undertaken, a central site for analysis is highly recommended so as to
505 reduce variability in analysis.

506 **5.1. Input Data to Be Used**

507 The input data that will be utilized will be in the form of concentration curves, and parametric maps of K^{trans}
508 and IAUGC_{BN} from which ROI analysis can be performed. One shortcoming of the 3D fast spoiled gradient
509 recalled echo technique used to acquire the dynamic images is that initial and end slice locations give
510 inaccurate results due to wraparound artifact and variability in excitation profile. The extent of this
511 wraparound artifact is dependent on slice-oversampling and other vendor specific techniques. Image
512 analysis can begin by removing areas that are subjectively compromised by wraparound artifact. One
513 method that can be used to determine which slices to discard is to closely examine the T_1 maps obtained at
514 the initial and end slice locations. Marked non-physiologic overestimations of T_1 on initial and end slices are
515 indicative of artifact.

516 **5.2. Methods to Be Used**

517 The following methodology for image interpretation of parametric maps should be performed in order to
518 ensure complete reproducible and interpretable results.

519

520 **Tumor ROI Definition.**

521 • The first step in the extraction of quantitative parameters (K^{trans} or IAUGC_{BN}) associated with a
522 particular lesion is to segment this lesion from adjacent tissues. Which techniques of segmentation are
523 ideal or even acceptable for a given application is the subject of on-going research, but it is clear that the
524 segmentation techniques used must be tailored to the particular organ system being studied with DCE-MRI.
525 The following guidelines are proposed:

526

527 • The committee does not recommend an analysis scheme where an operator defines a lesion
528 by placing regions of interest directly on parameter maps as that will introduce bias into the results

529

530 • Less subjective results can be obtained by using correlative imaging to define the lesion.
531 These correlative images may be obtained at the same imaging session but not directly related to the DCE-
532 MRI images. (For example, a T_2 -weighted image of an organ, which clearly delineates lesions and their
533 boundaries, may be used.) Correlative images should be obtained in the same imaging plane as the DCE-
534 MRI series, with similar FOV and spatial resolution, if feasible. In this scenario, a registration step will likely
535 be required (see 9.2)

536

537 • An alternative approach, which may be helpful if the lesion is well delineated on contrast-
538 enhanced T₁-weighted MRI, is to create summation images (images obtained by adding together images
539 obtained on the dynamic series for each slice location). The average images can be used to segment the
540 lesion on one or more slices, and because these segmentations are (in the absence of patient or organ
541 motion) registered to the dynamic series, the segmentations can be used to directly extract lesion-based
542 parameters from parametric maps.

543
544 • Because of the presence of image noise on source images of the dynamic series, along with
545 time-dependent changes in signal intensity which may blur or even obliterate the border between lesion
546 and background tissue, analysis schemes in which lesions are segmented independently on each image of
547 the dynamic series should be avoided where possible. In the case of moving organs, it may be necessary to
548 segment the lesion of interest on early (preferably, before the arrival of the contrast bolus) or late dynamic
549 images and estimate the position of the segmented lesion in intermediate time points.

550
551 • Although lesions can be segmented using manual techniques, several techniques are
552 available that allow a semi-automated approach to be used. The training of operator or operators in
553 performing segmentations should be documented, preferably with training sets.

554 **Registration of segmentations and parameter maps.**

555
556 Unless the segmentations are derived from relatively motion-free or motion-corrected dynamic images (for
557 example, summary images) image registration techniques may need to be used to place the segmentations
558 and parameter maps into a single anatomic framework (see Section 9.2). The choice of registration
559 technique to be used depends upon the organ system being imaged; the details of this are beyond the
560 scope of this document. In performing registration techniques, either images aligned with the parametric
561 maps or correlative images upon which the segmentation was performed are used as the target image for
562 registration. The registered images are then interpolated from the source images. In interpolating
563 parameter maps to match correlative images, tri-linear techniques are favored to avoid artifacts that may
564 be associated with more advanced interpolation techniques.

565 **Extraction of values for statistical comparison**

566
567 To derive values for statistical comparison from K^{trans} or IAUGC_{BN} parameter maps, median, mean and
568 standard deviation of the pixel values should be calculated, and the median is considered the primary figure
569 of merit. In a patient with multiple lesions due to metastatic disease, each lesion should be reported and
570 tracked separately. In a patient with multiple lesions due to recurrent local tumor (for example, recurrent
571 glioblastoma) per-patient figures of merit should be reported by aggregating the results of the multiple
572 lesions.

573 **Choice of time point for segmentation.**

574
575
576 As a rule, the K^{trans} or IAUGC_{BN} at a given time point should be extracted using tumor ROIs segmented from
577 the same imaging examination. However, in the situation where anti-angiogenic therapies are evaluated
578 and post-therapy imaging is performed within 72 hours of initial treatment with the anti-angiogenic agent,
579 it is acceptable to use a recent (within 1 week) pre-therapy time point to provide the segmentation used to
580 define the lesion on the immediate post-therapy imaging session. In this case, it is presumed that changes
581
582

583 in the appearance of lesions on immediate post-therapy study are due to immediate decreases in
584 permeability or blood flow rather than decrease in lesion volume.

585
586 In settings where analysis is performed retrospectively, all time points should be made available to the
587 reader simultaneously to allow for consistency in choice of tumor(s) for segmentation, and to ensure that
588 similar regions of large tumors have been sampled and segmented. In the case of manual VIF segmentation,
589 such workflow analyses also allow for greater standardization of the region of the aorta or other artery
590 used in the analysis. In such settings, the reader should be blinded to the nature of each time point, so that
591 inherent bias in tumor and/or VIF segmentation does not influence the results.

592 **6. Archival and Distribution of Data**

593 Archival and data distribution procedures are recommended so that all analysis results can be recomputed
594 for verification and validation purposes. In addition to saving of all original images in DICOM formats, the
595 following information must be archived along with the image data:

596 **6.1. Post-Processed Data**

597 **VIF:** Detailed specification of the vascular input function selection. This may include a binary mask of
598 pixels selected for arterial input function, or may consist of a tabulated text file containing RAS coordinates
599 co-ordinates of the VIF pixel locations.

600
601 **Registration:** Recorded parameters and user inputs required for image registration, if used. Time-series
602 image registration may be used to align data spatially over time. Any parameters which control the
603 performance of the registration algorithm (metric used, optimization parameters, user click points/sub
604 regions used for alignment, etc) must be stored in suitable format. It is preferred to save the registration
605 transform parameters so that identical registration can be reproduced in a multi-center environment.

606 **6.6. Analysis Results**

607 All regions of interest where analysis is performed and statistics are computed should be saved. In addition,
608 all computed maps (K^{trans} and IAUGC_{BN}), should be saved in DICOM and DICOM secondary capture modes.
609 $K^{\text{trans}} \text{ min}^{-1} * 10000$.

610 **6.7. Interpretation Results**

611 See 11.6

612 **7. Quality Control**

613 The following section deals with all aspects of quality control in DCE-MRI studies. This includes selecting
614 and qualifying an MRI imaging center, MRI personnel, and specific MRI scanners. In addition, the use of
615 phantom imaging (prior to study initiation and ongoing) is discussed. Finally, post image acquisition quality
616 assessment is detailed. Details of these processes will vary for investigator-initiated single site studies
617 versus sponsor-driven multi site studies.

618
619 Mechanisms for appropriate patient and tumor selection, image acquisition, and post processing are

620 discussed throughout the document.

621 **7.1. Selection of appropriate imaging centers for DCE-MRI studies**

622 Typically sites are selected for DCE-MRI due to their competence in clinical oncology and access to a
623 sufficiently large patient population under consideration. Sites must also be highly competent in clinical
624 MRI techniques appropriate to the area(s) of anatomy to be imaged during the DCE-MRI study. In order to
625 ensure high quality DCE-MRI results, it is essential to implement procedures that ensure quality assurance
626 of the scanning equipment and reliable image acquisition methodology. These processes must be set-up at
627 the outset, and followed throughout the duration of the study. A site “imaging capability assessment” prior
628 to site selection is therefore a requirement for any DCE-MRI study. This will include assessment of:

- 629 • appropriate imaging equipment and quality control processes (see section 12.1.1)
- 630 • appropriate injector equipment and contrast media
- 631 • experienced MR technologists or technical MR experts
- 632 • experienced MR radiologists or other anatomic experts
- 633 • procedures to assure imaging protocol compliance during the trial

634 **7.1.1 DCE-MRI Acquisition Scanner**

635 DCE-MRI studies as developed in this profile require a 1.5 T MR scanner. The scanner software version
636 should be identified and tracked across time, with updates and changes in scanner software noted during
637 the course of a trial.

638

639 Proper receiver coil maintenance must be performed to ensure adequate coil performance. Alternate
640 receiver coil systems must be available in the event that coil malfunction is identified prior to or during a
641 DCE-MRI study.

642

643 The MRI scanner and receiver coils must undergo routine quality assurance and quality control processes
644 (including preventive maintenance schedules) appropriate for clinical MRI applications. In addition, in
645 order to assure adequate quantitative MR imaging results, additional quality control measures are
646 required, as discussed below.

647

648 It is beneficial to identify and qualify more than one 1.5T MRI scanner at the site, if such are available for
649 study use. This will ensure that if the primary MRI scanner is temporarily unavailable, the DCE-MRI study
650 may proceed on a secondary scanner.

651

652 **7.1.2 DCE-MRI Power Injector**

653

654 A power injector is required for all DCE-MRI studies. The power injector needs to be properly serviced and
655 calibrated.

656

657 **7.1.3 MR Technologists or other Site Personnel performing DCE-MRI studies**

658 MR technologists or other imaging expert(s) performing DCE-MRI procedures should be MR certified
659 according to local regulations or institutional requirements. These individuals should have prior experience
660 in conducting dynamic contrast enhanced imaging. The personnel should also be experienced in clinical
661 study related imaging and should be familiar with good clinical practices (GCP). Competence in the

662 performance of DCE-MRI should never be limited to a single individual at the imaging center, as scheduled
663 and unplanned personnel absences are to be expected in the course of a DCE-MRI trial.
664

665 **7.1.4 MR Radiologists or other anatomic experts**

666 As tumor identification and selection is a critical component of the DCE-MRI study, sites performing DCE-
667 MRI must have access to highly qualified MRI radiologists or other experts in MRI anatomic assessment.
668 These individuals must be available during each DCE-MRI study to confirm adequate tumor selection and
669 slab placement. In some settings, (e.g. brain tumors), it may be feasible for tumor identification and slab
670 placement to be performed by the MR technologist, with oversight by a neuro-radiologist. In other cases
671 (e.g. wide-spread metastatic disease in the chest, abdomen, or pelvis), it is accepted that a radiologist or
672 other anatomic specialist must be available to identify tumor locations prior to contrast injection. It is
673 expected that more than one anatomic specialist be available at a site performing the examination, should
674 the primary anatomic specialists not be available for a given study.
675

676 **7.1.5 Site compliance with protocol requirements**

677 Imaging centers participating in DCE-MRI trials must adhere to accepted standards of quality control in
678 imaging studies. This includes processes to identify patients, who are participants in research studies,
679 personnel familiar with local IRB and other regulatory practices, familiarity with source documentation, and
680 reporting of protocol deviations and adverse events. Imaging centers must be able to document their
681 compliance with DCE-MRI procedures in order to facilitate central quality control and auditing processes.
682 Centers participating in multi-site trials must be familiar with protocol-directed methods for image transfer
683 of HIPAA-compliant anonymized imaging data, properly annotated, to central analytic laboratories.
684
685

686 **7.2 Site qualification process**

687 **7.2.1 Site readiness**

688 Site readiness for DCE-MRI should be documented prior to the initiation of the DCE-MRI trial. In single-site
689 studies initiated by in-house investigators, imaging procedures should be reviewed with the DCE-MRI team
690 prior to study initiation. In multi-site studies, site readiness assessment can begin with a simple
691 questionnaire completed as a pre-qualification step. A subsequent site visit prior to DCE-MRI study
692 initiation is recommended. During the site visit, study related imaging procedures and protocols are
693 discussed. Ideally, all DCE-MRI scan parameters are reviewed and entered at the MR scanner at the time of
694 the study visit. In some cases, initial phantom scanning can be performed during the site visit to familiarize
695 local MR personnel with proper phantom handling, set-up, and scanning.
696
697

698 **7.2.2 Scanner qualification**

699 MR scanners should be identified based on their vendor, model, and machine name. Hardware
700 specifications (maximum gradient strength, slew rate, etc.) should be documented. Software versions in
701 place at the time of trial initiation, and at all upgrades should be documented as well. Local receive coils to
702 be used should be noted, with quality checks per local institutional methods documented. Power injector
703 models should be noted, including date of most recent calibration.
704
705
706

7.2.3 Phantom imaging

To qualify the MRI scanner, a phantom imaging process is required. The QIBA DCE-MRI phantom, or a similar multi-compartment phantom with range of R_1 relaxation rate values appropriate for DCE-MRI should be utilized. With the exceptions noted below, imaging of the phantom should otherwise be performed using the same R_1 mapping and DCE-MRI acquisitions that are to be used in the clinical research protocol. Coil placement should approximate that which would be used by the site for the typical patient and anatomy of interest.

7.2.4 Phantom imaging data analysis

Phantom data should be analyzed in a uniform method by a centralized DCE-MRI image analysis center. Assurance should be made by the central site that the phantom scan orientation is correct, and appropriate image rotations or inversions were performed (and documented by the image analysis center).

For all phantom image tests, a single central slice is utilized. Uniform 2cm ROI spheres are placed within each phantom compartment, avoiding the edges of the compartments where signal intensity may be altered by Gibbs lines or other artifacts. Mean and standard deviation of the signal intensities within each ROI should be noted. There are three categories of DCE-MRI phantom data analysis: signal stability, signal linearity, and R_1 precision. In all cases, analysis should use a single central slice of the phantom data for analysis.

7.2.4.1. Signal stability

The signal stability test is performed using the DCE-MRI acquisition method to be used for the dynamic gadolinium enhanced imaging. The duration of this scan should be at least 6 minutes to test magnet stability. A single R_1 compartment with adequate SNR (10:1 or higher) is required. The mean SI in the ROI is then plotted over time. The plot should be linear and horizontal with no upward or downward trends. The root mean squared (rms) noise calculation should be similar across all aspects of the scan.

Marked deviations or drift of signal intensity over time indicate magnet instability, and should initiate a thorough evaluation of the magnet by the on-site MR physicist or site engineer prior to use in the DCE-MRI trial. The source of magnet instability should be determined and corrected prior to use in the DCE-MRI trial.

7.2.4.2 Signal linearity

In cases where signal intensity differences are to be used as a marker of tumor gadolinium concentration (see section 9), the linearity of MRI signal intensity with respect to R_1 over a range of R_1 values is required. While published guidelines on the allowed deviation from linearity do not exist, a linear correlation coefficient between SI and R_1 of 0.9 or higher is expected.

If a good linear correlation between SI and R_1 is not achieved, it is recommended that the receive coil array used for phantom imaging be evaluated to ensure that coil failure was not a cause of the abnormal results. The phantom image may be repeated with a different local coil array, or with the body coil as receiver to further evaluate this issue.

753
754 If linearity of SI vs. R_1 is still not achieved, it is recommended that the phantom scan be repeated with a
755 larger flip angle, in order to increase the relative T_1 weighting of the images.

756 757 *7.2.4.3 R_1 precision*

758
759 If T_1 -dependent analysis is intended for the DCE-MRI study, the fidelity of T_1 measurement should be
760 assessed based on the phantom imaging. As uncertainty in the measurement of T_1 is an important
761 contributor to concentration measurement bias³⁹, the measured phantom R_1 values based on the VFA
762 method (see Section 9) should be compared within the known R_1 values calibrated based on non-flip angle
763 dependent methods (such as IR-prepped imaging). Simulation studies suggest that variation in the R_1 value
764 by greater than 15% from actual may severely affect the reliability of the DCE-MRI quantification when T_1 -
765 dependent modeling of tumor gadolinium concentration in DCE-MRI studies is used.

766
767 If accurate R_1 values cannot be reproduced, it is recommended that T_1 -dependent modeling not be
768 performed.

769 770 *7.2.3 Ongoing MRI scanner quality control*

771
772 The phantom scans and analysis should be repeated at regular intervals, such as every 3 months, during the
773 course of the study. Any changes to scanner equipment, including major hardware changes or any software
774 version change, need to be documented and will result in the need for imaging qualification renewal prior
775 to repeat imaging. In particular, it is strongly recommended that patients undergoing longitudinal study be
776 scanned on the same MRI system with the same software version whenever possible. Sites performing
777 DCE-MRI studies should be informed of planned software upgrades, when possible deferring such upgrades
778 until serial imaging of all currently enrolled patients is complete.

779 780 *7.2.4 Use of Human test subjects*

781
782 Given the complexities of local site regulatory environments, it is recognized that the use of human test
783 subjects to qualify DCE-MRI imaging scanners and sites may not be feasible. In such cases, it is strongly
784 encouraged that the initial patients accrued on a DCE-MRI trial at each site be considered a “test subject”
785 and that the data for that patient not be analyzed with the remainder of the cohort. Central analysis of the
786 initial patient data will then serve to finalize the qualification of the DCE-MRI center.

787 788 **7.3. Quality Control of DCE-MRI studies**

789 *7.3.1 Determination of suitable tumor lesions*

790
791 Patients suitable for DCE-MRI analysis must possess at least one tumor ≥ 2 cm, well removed from areas
792 subject to large degrees of cardiac pulsatility artifact, that is not largely cystic or necrotic. Determination of
793 patient eligibility is usually based on pre-enrollment imaging (often CT or clinical MRI) which then serves as
794 a baseline study for subsequent assessments for tumor response or progression. The site radiologist then
795 reviews these images prior to enrollment to ascertain the location of the most suitable tumor lesion(s) for
796 analysis.

797 798 *7.3.2 Selection of target lesion*

799

800 Once the MRI scan commences, the radiologist or anatomic expert will review the pre-gadolinium imaging
801 to identify putative target lesions. The DCE-MRI study then proceeds with slab placement and T₁
802 mapping/dynamic enhanced imaging once the target lesion is identified. Sites should strive inspect these
803 images to ensure absence of substantial artifacts (e.g. phase wrap, pulsatility) overlying the target lesion,
804 with protocol specified adjustments to patient positioning and slab placement prior to continuing the DCE-
805 MRI study. Once the final slab placement is confirmed, grid line overlays of the DCE-MRI slab on routine
806 anatomic imaging (usually axial plane) is recommended to facilitate DCE-MRI slab placement on subsequent
807 visits (e.g. by saving of a screen shot).

808

809 7.3.3 Determination of subjects unsuitable for DCE-MRI analysis

810

811 Despite best efforts and protocol adherence, on occasion, a patient enrolled and imaged in DCE-MRI study
812 will be found to be ineligible for subsequent analysis. Reasons for eliminating patients for analysis include:

813

- 814 • Lack of a tumor of suitable size in the usable DCE-MRI imaging volume
- 815 • Unacceptable pulsatility, wrap, or metallic artifact involving all tumors in the usable DCE-MRI
816 imaging volume
- 817 • All target lesions in the DCE-MRI imaging volume determined to be largely cystic or necrotic

818

819 Determination of patient eligibility should be made by an independent reviewer who is blinded to other
820 attributes of patient data, including (when applicable) randomization arm/drug treatment, toxicity, and
821 clinical outcomes. Decisions on eligibility should be made on the basis of visual image assessment prior to
822 analysis of DCE-MRI data. Quantitative criteria for defining tumors that are largely cystic or necrotic (such
823 as percentage of pixels with enhancement above a certain threshold) should be defined in the protocol to
824 avoid bias in decisions to eliminate patients from further DCE-MRI assessment.

825

826 7.3.4 Determination of DCE-MRI exams unsuitable for DCE-MRI analysis

827

828 In addition, individual DCE-MRI examinations may be deemed nonanalyzable based on a variety of technical
829 deviations. These may include:

830

- 831 • Failure of gadolinium injection
- 832 • Gross patient motion not correctable with motion correcting algorithms
- 833 • Failure of the imaging site to replicate the imaging parameters within acceptable standards of
834 deviation from protocol specifications
- 835 • Failure of the imaging site to replicate anatomic DCE-MRI slab placement

836

837 Whenever possible, all anticipated instances where individual DCE-MRI data will be removed from analysis
838 should be prespecified in the DCE-MRI protocol.

839

840 7.3.5 Editing of DCE-MRI exams prior to DCE-MRI analysis

841

842 It is recognized that DCE-MRI analysis requires post-processing of the DCE-MRI image sets. Most
843 frequently, data sets will be subject to automated or semi-automated motion compensation schemes to
844 eliminate or minimize the effects of image motion of subsequent DCE-MRI analysis. The methodology used

845 for such post processing should be documented, ideally in the DCE-MRI protocol or the standard operating
846 procedures of the central analysis laboratory. Motion correction matrices keyed to each temporal phase
847 may be documented as part of the analysis routine, in order to facilitate replication of the data analysis
848 when required.

849
850 In the course of post processing, individual phases of the DCE-MRI exam may be found to be severely
851 compromised by image blur or degraded by other artifacts (such as random noise spikes). Judicious
852 selection of phases to be eliminated for analysis may be made by the central analysis team, provided that
853 the decision to eliminate such phases is determined prior to data analysis. Elimination of baseline or early
854 post gadolinium phases is discouraged as such post processing may substantially alter the subsequent
855 analysis. Data documenting these forms of post-processing should be maintained by the imaging analysis
856 laboratory.

859 **8. Imaging-associated Risks and Risk Management**

860 MR safety considerations are to be established individually at each institution according to each
861 institutions' radiology departmental guidelines and institutional review board (IRB) considerations to
862 include policy guidelines on the following:

- 863 (1) laboratory screening for renal dysfunction prior to gadolinium based contrast administration
- 864 (2) contrast administration in pregnant patients and in patients who are lactating
- 865 (3) policy on patients receiving gadolinium based agents who have a positive history of a previous adverse
866 event or events to iodinated or gadolinium based contrast agents to include serious and non-serious
867 adverse events. The American College of Radiology Manual on Contrast Media Version 7 2010 can serve as
868 a referenced guideline for each institutional policy development. This manual reflects policy statements
869 previously released by the Food and Drug Administration (FDA) in the United States and its counterpart in
870 the European Union, The Committee for Medicinal Products for Human Use (CHMP).

871 **IV. Compliance**

872 Typically clinical sites are selected due to their competence in oncology and access to a sufficiently large
873 patient population under consideration. For DCE-MRI use as quantitative imaging biomarker it is essential
874 to put some effort into an imaging capability assessment prior to final site selection for a specific trial. For
875 imaging it is important to consider the availability of:

- 876 • appropriate imaging equipment and quality control processes,
- 877 • appropriate injector equipment and contrast media,
- 878 • experienced MR technologists for the imaging procedure, and
- 879 • processes that assure imaging protocol compliant image generation at the correct point in time.

880 **Acquisition Scanner**

881 1.5 T MR machines with 55-70 cm bores need to be available. The scanner needs to be under quality
882 assurance and quality control processes (including preventive maintenance schedules) appropriate for
883 quantitative MR imaging applications, which may exceed the standard requirements for routine clinical
884 imaging or for MR facility accreditation purposes. The scanner software version should be identified and
885 tracked across time. It might be beneficial to identify and qualify a second scanner at the site, if available. If

886 this is done prior to the study start there will be no difficulties later on in case the first scanner is
887 temporarily unavailable. Practically speaking sites are encouraged to perform longitudinal treatment trials
888 on one instrument.

889 **Contrast Inject Device**

890 A power injector is required for DCE-MRI studies. It needs to be properly serviced and calibrated.

891 **Software Analysis**

892 When a site is performing parametric image analysis and interpretation, a DCE-MRI tool that complies with
893 the Toft's model should be utilized. In addition, for multi-institutional trials a central reading site is
894 assumed.

895 **Performing Site**

896 MR technologists running DCE-MRI procedures should be MR certified according to local regulations. The
897 technologists should have prior experience in conducting dynamic contrast enhanced imaging. The person
898 should be experienced in clinical study related imaging and should be familiar with good clinical practices
899 (GCP). A qualified backup person is needed that should fulfill the same requirements. Contact details for
900 both technologists should be available in case of any questions.

902 **Imaging qualification process:**

904 The above-mentioned details can be obtained using a simple questionnaire as a pre-qualification step. If
905 appropriate equipment and personnel are available, a site visit is recommended. During the site visit, study
906 related imaging protocols are discussed and, ideally, all scan parameters are entered at the MR scanner.

908 To qualify the scanner, a phantom imaging process is strongly recommended. The QIBA DCE-MRI phantom,
909 or a similar multi-compartment phantom with range of relaxation rate (T_1) values appropriate for the DCE-
910 MRI study to be performed, should be used if the Profile Claim given above is to be assured. Data should be
911 acquired from the multi-compartment phantom using the same T_1 mapping and DCE-MRI acquisitions that
912 will be used in the proposed clinical application or clinical research protocol (see Section 6).

914 The phantom scans should be repeated on a regular interval (e.g 3 months) during the course of the study.
915 Ongoing image quality inspection on a per scan basis is essential. Any changes to scanner equipment,
916 including major hardware changes or any software version change, need to be documented and will result
917 in the need for imaging qualification renewal.

919 **Site Analysis qualification:**

921 The data analysis procedures to be used in the DCE-MRI application should be used to analyze the T_1
922 mapping data and results compared to the known T_1 values of the various compartments. As uncertainty in
923 the measurement of T_1 is an important contributor to concentration measurement bias³⁹, the measured
924 values should compare within 15 % of the known values over a T_1 range of approximately 50-1000 ms. The
925 DCE-MRI data obtained from the phantom should be analyzed to confirm the correct temporal resolution
926 and to provide SNR measurements and signal intensity vs. T_1 characteristics for the specific DCE-MRI
927 acquisition protocol.

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Significant variations in any of these parameters during the course of an ongoing longitudinal study can affect the resulting imaging biomarker determinations, in the case of this specific claim K^{trans} and $IAUGC_{BN}$, and such changes can readily occur if there are major changes in the scanner hardware or software, e.g., an update to the pulse sequence used for the DCE-MRI and/or T_1 measurements or to the gradient subsystem. All results shall be documented and, if they pass the established acceptance values, will constitute the site qualification documentation for the DCE-MRI procedure. This process ensures study specific training of the site personnel and needs to be documented and signed.

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 1041

1042 **Appendices**

1043 **Appendix A: Acknowledgements and Attributions**

1044	I. Executive Summary	Jeffrey Evelhoch
1045	II. Clinical Context and Claims	Mitchell Schnell
1046	III. Profile Details	
1047	1. Subject Handling	Alex Guimaraes
1048	2. Imaging Procedure	Ed Jackson/Sandeep Gupta
1049	3. Image Post-processing	Sandeep Gupta
1050	4. Parametric image formation	Ed Ashton
1051	5. Parametric image analysis	Dan Barboriak
1052	6. Archival and Distribution of Data	Sandeep Gupta
1053	7. Quality Control	Mark Rosen
1054	8. Imaging associated Risks and Risk Management	Orest Boyko

1055 **Appendix B: Conventions and Definitions**

- 1056 List of Abbreviations
 1057 - VIF: Vascular input function

- 1058 - DCE-MRI: Dynamic contrast enhanced magnetic resonance imaging
- 1059 - ECOG: Eastern Cooperative Oncology Group
- 1060 - eGFR: estimated Glomerular Filtration Rate
- 1061 - Gd-DTPA: Gadolinium – diethylene triamine pentaacetic acid
- 1062 - IAUGCBN: Initial area under the Gadolinium concentration blood normalized
- 1063 - Ktrans: permeability transfer constant
- 1064 - QIBA: Quantitative Imaging Biomarkers Alliance
- 1065 - ROI: Region of Interest
- 1066 - VEGF: Vascular Endothelial Growth Factor
- 1067 - VFA: Variable Flip angle
- 1068 - VIF: Vascular input function

1069
 1070 **ECOG Performance Status Descriptions, by grade:** ⁴⁰

- 1071 0: Fully active, able to carry on all pre-disease performance without restriction
- 1072 1: Restricted in physically strenuous activity but ambulatory and able to carry out work of a light or
- 1073 sedentary nature, e.g., light-house work, office work
- 1074 2: Ambulatory and capable of all self-care but unable to carry out any work activities. Up and about more
- 1075 than 50% of waking hours
- 1076 3: Capable of only limited self-care, confined to bed or chair more than 50% of waking hours
- 1077 4: Completely disabled. Cannot carry on any self-care. Totally confined to bed or chair
- 1078 5: Dead

1079 **Appendix C: Spreadsheet on reproducibility data**

1080

Table 1. Summary of the data for the following table.

Reference	Year	Modality	System	Manufacturer	Model	Version	Software	Hardware	Configuration	Performance	Compliance	Notes
1	2005	CT	Siemens	Siemens	Hi-CT	1.0	Siemens Hi-CT 1.0	Siemens Hi-CT 1.0	Siemens Hi-CT 1.0	Siemens Hi-CT 1.0	Siemens Hi-CT 1.0	Siemens Hi-CT 1.0
2	2005	CT	Siemens	Siemens	Hi-CT	1.0	Siemens Hi-CT 1.0	Siemens Hi-CT 1.0	Siemens Hi-CT 1.0	Siemens Hi-CT 1.0	Siemens Hi-CT 1.0	Siemens Hi-CT 1.0
3	2005	CT	Siemens	Siemens	Hi-CT	1.0	Siemens Hi-CT 1.0	Siemens Hi-CT 1.0	Siemens Hi-CT 1.0	Siemens Hi-CT 1.0	Siemens Hi-CT 1.0	Siemens Hi-CT 1.0
4	2005	CT	Siemens	Siemens	Hi-CT	1.0	Siemens Hi-CT 1.0	Siemens Hi-CT 1.0	Siemens Hi-CT 1.0	Siemens Hi-CT 1.0	Siemens Hi-CT 1.0	Siemens Hi-CT 1.0
5	2005	CT	Siemens	Siemens	Hi-CT	1.0	Siemens Hi-CT 1.0	Siemens Hi-CT 1.0	Siemens Hi-CT 1.0	Siemens Hi-CT 1.0	Siemens Hi-CT 1.0	Siemens Hi-CT 1.0
6	2005	CT	Siemens	Siemens	Hi-CT	1.0	Siemens Hi-CT 1.0	Siemens Hi-CT 1.0	Siemens Hi-CT 1.0	Siemens Hi-CT 1.0	Siemens Hi-CT 1.0	Siemens Hi-CT 1.0
7	2005	CT	Siemens	Siemens	Hi-CT	1.0	Siemens Hi-CT 1.0	Siemens Hi-CT 1.0	Siemens Hi-CT 1.0	Siemens Hi-CT 1.0	Siemens Hi-CT 1.0	Siemens Hi-CT 1.0
8	2005	CT	Siemens	Siemens	Hi-CT	1.0	Siemens Hi-CT 1.0	Siemens Hi-CT 1.0	Siemens Hi-CT 1.0	Siemens Hi-CT 1.0	Siemens Hi-CT 1.0	Siemens Hi-CT 1.0
9	2005	CT	Siemens	Siemens	Hi-CT	1.0	Siemens Hi-CT 1.0	Siemens Hi-CT 1.0	Siemens Hi-CT 1.0	Siemens Hi-CT 1.0	Siemens Hi-CT 1.0	Siemens Hi-CT 1.0
10	2005	CT	Siemens	Siemens	Hi-CT	1.0	Siemens Hi-CT 1.0	Siemens Hi-CT 1.0	Siemens Hi-CT 1.0	Siemens Hi-CT 1.0	Siemens Hi-CT 1.0	Siemens Hi-CT 1.0

The following table provides a summary of the data for the following table. The table is organized into columns representing the reference number, year, modality, system, manufacturer, model, version, software, hardware, configuration, performance, compliance, and notes. The data is presented in a structured format for easy reference.

Appendix D: Model-specific Instructions and Parameters

1083 The presence of specific product models/versions in the following tables should not be taken to imply that
1084 those products are fully compliant with the QIBA Profile. Compliance with a profile involves meeting a
1085 variety of requirements of which operating by these parameters is just one. To determine if a product (and
1086 a specific model/version of that product) is compliant, please refer to the QIBA Conformance Document for
1087 that product. G.1. Image Acquisition Parameters The following technique tables' list acquisition parameter
1088 values for specific models/versions that can be expected to produce data meeting the requirements of
1089 Section 7.1.

1090 These technique tables may have been prepared by the submitter of this imaging protocol document, the
1091 clinical trial organizer, the vendor of the equipment, and/or some other source. (Consequently, a given
1092 model/version may appear in more than one table.) The source is listed at the top of each table.

1093 Sites using models listed here are encouraged to consider using these parameters for both simplicity and
1094 consistency. Sites using models not listed here may be able to devise their own acquisition parameters that
1095 result in data meeting the requirements of Section 7.1 and conform to the considerations in Section 13.

1096 In some cases, parameter sets may be available as an electronic file for direct implementation on the
1097 imaging platform.
1098

1099 **Siemens**
 1100 QIBA DCE-MRI Abdominal Protocol for VA30 Software
 1101
 1102

parameter	value	notes
Routine tab		
slabs	1	
distance factor	irrevelant	
position	as needed	
orientation	coronal	
phase enc. dir.	R >> L	
rotation	0.0 deg	
phase oversampling	0%	
slice oversampling	0%	
slices per slab	26	Reconstructed images, interpolated by zero-filling. The slab thickness is $4.25 \times 26 = 110.5$ mm
FoV read	400	
FoV phase	81.3%	325 mm
slice thickness	4.25 mm	For 3-D, this is the slice <i>spacing</i> . The true slice thickness is this number divided by the slice resolution, in this case $4.25 / 0.62 = 6.85$ mm.
TR	5.03 ms	
TE	1.9 ms	
averages	1	NEX
concatenations	1	
filter	none	
coil elements	as needed	
Contrast tab		
flip angle	30 deg	
fat suppression	none	
water supp.	none	
Dixon	no	
save original images	on	
averaging mode	short term	
reconstruction	magnitude	
measurements	40	
measurement series	each measurement	
pause after measurement	0 sec	
Resolution tab		
base resolution	256	readout pixel size 1.56 mm
phase resolution	62%	phase pixel size 2.52 mm
slice resolution	62%	Controls zero-filling in slice. If no partial Fourier processing is used, 16 partitions are acquired. The raw matrix is padded with 10 zeros to reconstruct 26 slices: $16 / 0.62 = 26$. Divide the slice spacing by the slice resolution to get the slice thickness: $4.25 / 0.62 = 6.85$ mm
phase partial Fourier	choose 7/8ths here or below (slice)	If 7/8ths is chosen, partial Fourier processing is used to reduce the number of acquired lines to: $256 \times 0.62 \times 0.813 \times 7/8 = 113$
slice partial Fourier	choose 7/8ths here or above (phase)	If 7/8ths is chosen, 14 partitions are acquired to provide the data for 16. Ten additional zeros are added to reconstruct 26 slices.
interpolation	on	In-plane zero-filling to 512×512 .

PAT mode	none	No SENSE or GRAPPA
matrix coil mode	as needed	
image filter	off	
distortion correction	off	also called "large FoV filter"
prescan normalize	off	
normalize	off	Acts on individual slices, so must be turned off.
raw filter	off	
elliptical filter	off	
Geometry card		
multi-slice mode	irrelevant	
series	irrelevant	
special sat.	none	
(remainder)		May be ignored.
System Card		
shim mode	standard	
save uncombined	off	
adjust with body coil	off	
Physio card		
1 st signal/mode	none	
rsp. control	off	
Inline card		
3D centric reordering	off	
(remainder)	off	
Sequence card		
introduction	off	
dimension	3D	
elliptical scanning	off	
asymmetric echo	allowed, weak	
contrasts	1	
bandwidth	250 Hz/pixel	Corresponds to ± 32 KHz.
optimization	min TE	
RF pulse type	normal	
gradient mode	fast	
excitation	slab-sel.	
RF spoiling	on	For the FLASH sequence.
Tool Tips		
readout echo position	38%	Roll over "echo asymmetry."
matrix size	129 x 256	Roll over "phase resolution." This size includes the effects of reduced pixel resolution and rectangular FoV.
slab thickness	110 mm	
pulse sequence	fl3d_vibe	Roll over the pulse sequence abbreviation.

1103

1104

SNR protocol: change measurements to 8 and flip angle to 15°.

1105

Variable flip angle protocol for T₁: one measurement, 4 averages, and flip angles of 2°, 5°, 10°, 15°, 20°,

1106

25°, and 30°.

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QIBA DCE-MRI Abdominal Protocol for VB15, VB17, and VD11 Software
These are the 400 Hz/pixel protocols.

parameter	value	notes
Routine tab		
slabs	1	
distance factor	irrelevant	
position	as needed	
orientation	coronal	
phase enc. dir.	R >> L	
rotation	0.0 deg	
phase oversampling	0%	
slice oversampling	0%	
slices per slab	26	Reconstructed images, interpolated by zero-filling. The slab thickness is $4.25 \times 26 = 110.5$ mm
FoV read	400	
FoV phase	81.3%	325 mm
slice thickness	4.25 mm	For 3-D, this is the slice <i>spacing</i> . The true slice thickness is this number divided by the slice resolution, in this case, $4.25 / 0.62 = 6.85$ mm.
TR	3.61 ms 3.91 ms 4.76 ms	VD11, Aera VB17, Espree VB15B, Verio
TE	1.49 ms 1.48 ms 1.43 ms	VD11, Aera VB17, Espree VB15B, Verio
averages	1	NEX
concatenations	1	
filter	none	
coil elements	as needed	
Contrast tab		
flip angle	30 deg	
fat suppression	none	
water suppression	none	
Dixon	no	
save original images	on	
averaging mode	short term	
reconstruction	magnitude	
measurements	50	as needed
measurement series	each measurement	
pause after measurement	0 sec	for all measurements
Resolution tab		
base resolution	256	readout pixel size 1.56 mm
phase resolution	62%	phase pixel size 2.52 mm
slice resolution	62%	Controls zero-filling in slice. Sixteen partitions are acquired. The raw matrix is padded with 10 zeros to reconstruct 26 slices: $16 / 0.62 = 26$ Divide the slice spacing by the slice resolution to get the slice thickness: $4.25 / 0.62 = 6.85$ mm
phase partial Fourier	off	No further reduction in the number of acquired lines: $256 \times 0.62 \times 0.813 = 129$

slice partial Fourier	off	No further reduction in the number of acquired partitions (16).
interpolation	on	In-plane zero-filling to 512 x 512.
PAT mode	none	No SENSE or GRAPPA
matrix coil mode	as needed	
image filter	off	
distortion correction	off	
prescan normalize	off	
normalize	off	Acts on individual slices, so must be turned off.
B₁ filter	off	
raw filter	off	
elliptical filter	off	
POCS	off	
Geometry card		
multi-slice mode	irrelevant	
series	irrelevant	
special sat.	none	
Set-n-Go Protocol	off	
inline composing	off	
System Card		
shim mode	tune up	
save uncombined	off	
adjust with body coil	off	
confirm freq. adjustment	off	
Physio card		
1 st signal/mode	none	
resp. control	off	
Inline card		
3D centric reordering	off	
(remainder)	off	
Sequence card		
introduction	off	
dimension	3D	
elliptical scanning	off	
asymmetric echo	allowed, weak	
contrasts	1	
bandwidth	400 Hz/pixel	Corresponds to ± 51.2 KHz.
optimization	min TE	
RF pulse type	normal	
gradient mode	fast normal fast	VD11, Aera VB17, Espree VB15B, Verio
excitation	slab-sel.	
RF spoiling	on	For the FLASH sequence.
Tool Tips		
readout echo position	38%	Roll over "echo asymmetry."
matrix size	129 x 256	Roll over "phase resolution." This size includes the effects of reduced pixel resolution and rectangular FoV.
slab thickness	110 mm	
pulse sequence	fl3d_vibe	Roll over the pulse sequence abbreviation.

1111

1112

1113

1114

SNR protocol: change measurements to 8 and flip angle to 15°.

Variable flip angle protocol for T₁: one measurement, 4 averages, and flip angles of 2°, 5°, 10°, 15°, 20°.

1115 25°, and 30°.
1116

QIBA Body Protocol

System: MR450w

Field Strength: 1.5T

SW: 22.0

Notes: Should work fine on other 1.5T systems as well including HDx

3T will likely degrade due to Whole Mode use on HDx platform.

Axial plane with Zoom coil might provide the desired TE/TRs



GE imagination at work

PSD: 3D Vascular TOF SPGR

- TOF SPGR is same as 3D FSPGR but TE/TR optimized for Contrast
- 16 slices Rxed -> 4 kissoffs. 12 slices be reconned pre ZIP2
- All parameters meet either ideal level or target goal

Cor ceMRA GRx Cx 0:05 Details Vascular Advanced

Scan Plane: Coronal ▼ Freq. Dist: 5/1 ▼ # of TECooper Scans: 1.0 ▼ Frequency: 256 ▼

Freq. FOV: 42.0 ▼ TR: 5.4 ▼ TE: Minimum ▼ Phase: 160 ▼

Phase FOV: 0.80 ▼ # Slabs: 1 ▼ Flip Angle: 30 ▼ NEX: 1.00 ▼

Slice Thickness: 5.0 ▼ Locks per Slab: 16 ▼ Intensity Correction: NONE ▼ Bandwidth: 100.00 ▼

Intensity Filter: None ▼ Shim: Auto ▼

A/P S/I R/L Max # Slices: 2016 3D Geometry Correction: Phase Correct: Off ▼

Start P27.5 S21.1 R21.1 # of Acqs: 1 Table Delta: 0.00

End A27.5 S21.1 R21.1 Rel. SNR00: 508 ±

Act End A37.5

Chem SAT: None ▼

Contrast

SAR Est: 2.99 Peak: 5.98 Mode: First dB/dt: First Minimum TE: 1.2 Maximum TE: 11

1119

Advanced Page

- Sequential view order is utilized: Centric and EC adds 1 sec extra per phase
- Turbo Mode=2 for shortest RF -> shortest TE/TR
- Slice resolution 70% for scan time = 5sec/phase

The screenshot displays the 'Advanced' configuration page for a Cor ceMRA scan. The top navigation bar includes 'GRx', '0:05', 'Details', 'Vascular', and 'Advanced'. The main area is divided into several sections:

- Scan Parameters:** Scan Plane: Coronal, Freq. FOV: 42.0, Phase FOV: 0.80, Slice Thickness: 5.0, Freq. Dir: S/I, TR: 3.1, # Slabs: 1, Locs per Slab: 16, Max # Slices: 1024, # of Acqs: 1, ReL SNRQ: 508.
- Start/End Coordinates:** Start (A/P: P27.5, S/I: 521.1, R/L: R21.1), End (A/P: A27.5, S/I: 521.1, R/L: R21.1).
- Chem SAT:** None.
- Contrast:** Contrast.
- User Control Variables Table:**

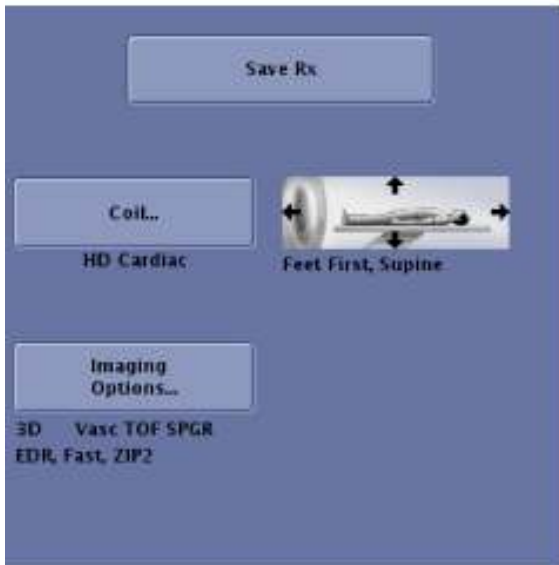
Variable ID	Parameter Name	Value	Min	Max
CV4	Image acq. delay (sec)	0.00	0.0	100.0
CV5	Whole Volume Excitation (0=off, 1=on)	0.00	0.0	1.0
CV6	Turbo Mode (0=off, 1=Fastest, 2=Fastest)	2.00	0.0	2.0
CV11	Reverse Elliptical Centric (0=off, 1=on)	0.00	0.0	1.0
CV12	Elliptical Centric (0=off, 1=std, 2=delay)	0.00	0.0	2.0
CV13	Centric (0=off, 1=on)	0.00	0.0	1.0
CV14	Reverse Centric (0=off, 1=on)	0.00	0.0	1.0
CV23	Slice Resolution (70%-100%)	70.00	70.0	100.0

At the bottom, SAR Est: 2.99 Peak: 5.98 Mode: First and dR/dt: First are displayed.

1120

Imaging Options & Vascular Page

- Turn off Projections and Collapse on Vascular Page
- Turn EDR ON: better dynamic range
- Use ZIP2 and ZIP512 if needed
 - Could help with ringing artifacts



1121
1122
1123

1124 **Phillips**
1125 Philips Achieva 1.5T (edited on release 2.6):
1126
1127 #####
1128 #####
1129 Pulse Sequence: 3D T₁ FFE
1130
1131 NEX = NSA: 2 (change accordingly as needed for ratio map or variable flip angle series)
1132
1133 flip angles: 30, 25, 20, 15, 10, 2 (watch that shortest TR/TE remain constant, or switch to user defined if
1134 needed)
1135
1136 coils: SENSE-body or SENSE-Torso-XL
1137
1138 slice orientation: coronal (for abdomen. for head: axial, adjust FOV as needed)
1139
1140 Foldover direction: RL
1141 Foldover suppression: yes
1142
1143 slice oversampling: user defined: 1
1144
1145 TE/TR: set to shortest, actual values will be: 5.0/2.4 ms (verify it stays constant with changing flip angle)
1146
1147 temporal resolution = dynamic scan time: 8.4 sec (for NSA 2)
1148
1149 receiver bandwidth – corresponding parameter: water fat shift: maximum (313 Hz/pixel for current
1150 parameters)
1151
1152 FOV: FH 420 mm, RL 340 mm, AP 48 mm (for head: 250 AP, 220 RL)
1153 voxel size: FH 1.64 mm, RL 2.1 mm, AP 2 mm (FOV/voxel size ratio yielding matrix: 256x162 for abdominal)
1154 (note, FOV and voxel size are adjustable parameters, corresponding matrix is displayed in info page)
1155
1156 over contiguous slices: yes (acquired slice thickness 4 mm, interpolated into 2)
1157 number of slices: 24 (interpolated – 12 acquired)
1158
1159 SENSE: no, CLEAR: no
1160 Half scan: yes, factor Y 0.65, factor Z = 0.8
1161
1162 Dynamic study: individual
1163 dynamic scans: 42 (giving total scan duration of 05:50)
1164 dynamic scan times: user defined > set 6th dynamic to manual (for injection after 5th dynamic,
1165 leave all other on shortest)
1166 #####
1167 #####
1168
1169