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| Title | Image synthesis with deep convolutional generative adversarial networks for material decomposition in dual- energy CT from a kilovoltage CT |
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| Relation | |



| 1 | Title |
|----|--|
| 2 | Image synthesis with deep convolutional generative adversarial networks for |
| 3 | material decomposition in dual-energy CT from a kilovoltage CT |
| 4 | |
| 5 | Abstract |
| 6 | |
| 7 | Generative Adversarial Networks (GANs) have been widely used and |
| 8 | it is expected to use for the clinical examination and image. The objective of |
| 9 | the current study was to synthesize material decomposition images of bone- |
| 10 | water (bone(water)) and fat-water (fat(water)) reconstructed from dual-energy |
| 11 | computed tomography (DECT) using an equivalent kilovoltage-CT (kV-CT) |
| 12 | image and a deep conditional GAN. The effective atomic number images were |
| 13 | reconstructed using DECT. We used 18,084 images of 28 patients divided into |
| 14 | two datasets: the training data for the model included 16,146 images (20 |
| 15 | patients) and the test data for evaluation included 1938 images (8 patients). |

| 16 | Image prediction frameworks of the equivalent single energy CT images at 120 |
|----|--|
| 17 | kVp to the effective atomic number images were created. The image-synthesis |
| 18 | framework was based on a CNN with a generator and discriminator. The mean |
| 19 | absolute error (MAE), relative mean square error (MSE), relative root mean |
| 20 | square error (RMSE), peak signal-to-noise ratio (PSNR), structural similarity |
| 21 | index (SSIM), and mutual information (MI) were evaluated. The Hounsfield |
| 22 | unit (HU) difference between the synthesized and reference material |
| 23 | decomposition images of bone(water) and fat(water) were within 5.3 HU and |
| 24 | 20.3 HU, respectively. The average MAE, MSE, RMSE, SSIM, and MI of the |
| 25 | synthesized and reference material decomposition of the bone(water) images |
| 26 | were 0.8, 1.3, 0.9, 0.9, 55.3, and 0.8, respectively. The average MAE, MSE, |
| 27 | RMSE, SSIM, and MI of the synthesized and reference material decomposition |
| 28 | of the fat(water) images were 0.0, 0.0, 0.1, 0.9, 72.1, and 1.4, respectively. The |

| 29 | proposed model can act as a suitable alternative to the existing methods for the |
|----|--|
| 30 | reconstruction of material decomposition images of bone(water) and fat(water) |
| 31 | reconstructed via DECT from kV-CT. |
| 32 | |
| 33 | Keywords: Deep learning, Medical imaging, Artificial Intelligence, Dual- |
| 34 | energy CT, Material Decomposition |
| 35 | |
| 36 | |
| 37 | |
| 38 | Compliance with Ethical Standards: |
| 39 | Disclosure of potential conflicts of interest |
| 40 | Author Daisuke Kawahara declares that he has no conflict of interest. |
| 41 | |
| 42 | Research involving human participants and/or animals |
| 43 | This article does not contain any studies with human participants or animals |
| 44 | performed by any of the authors. |
| 45 | |

46 **Ethnical approval**

All procedures performed in studies involving human participants were in
accordance with the ethical standards of the institutional and/or national
research committee and with the 1964 Helsinki declaration and its later
amendments or comparable ethical standards.

51

52 Informed consent

53 Informed consent was obtained from all individual participants included in

54 the study.

55

56

58 I. INTRODUCTION

59

Dual-energy computed tomography (DECT) uses two different energy spectra 60 61 (low and high energy) that can be created by combining two datasets acquired 62 over the same region [1]. DECT can obtain different information such as the 63 effective atomic number, monochromatic energy CT (mCT) number, iodine-64 enhanced map, bone-water (bone(water)) density images, and fat-water 65 (fat(water)) density images [2]. A (bone(water)) density image suppresses the 66 water signal and enhances the calcium signal. It is created from the DECT data 67 by estimating the amount of bone mineral, primarily composed of calcium, and subtracting this from the scanned original image [3, 4]. It is used for the 68 69 diagnosis of bone marrow edema, which is a biomarker for arthritis, bone 70 infarction, and hidden fractures. It is difficult to identify with traditional CT 71 owing to the intrinsic low contrast of the involved tissues; it has been detected 72 with magnetic resonance (MR) imaging [5]. MR represents the gold standard 73 for soft tissue imaging and can provide quantitative fat-fraction measurements. 74 A fat(water) image enhances the fat signal and suppresses other signals such as 75 water and bone. Recently, Hyodo et al. reported that the DECT technique can estimate fat quantification in the liver [6]. 76

A GE Revolution CT scanner (GE Healthcare, Milwaukee, WI) can reconstruct
the effective atomic number, 120 kVp equivalent images, monochromatic
energy CT, iodine contrast-enhanced, and calcium-enhanced images using a

gemstone spectral imaging (GSI) technique [7]. The disadvantages of DECTare the increasing radiation dose, scan time, and cost.

Deep learning has been widely used for denoising applications [8, 9]. The 82 83 denoising technique has been improved using a wavelet residual network, 84 which synergistically combines the expressive power of deep learning and the 85 performance guarantee from framelet-based algorithms [10]. The improvement 86 of the image resolution has led to a reduction in the radiation dose. 87 Convolutional neural networks (CNNs) have been successfully applied to 88 image synthesis and image processing. Dong *et al.* performed super-resolution 89 imaging using a CNN algorithm. Streak artifacts due to beam hardening and photon starvation have been potentially problematic. Zhang et al. suppressed 90 91 the artifacts dramatically using the CNN-based metal artifact reduction 92 framework, which fuses the information from the original and corrected images. 93 For material decomposition, deep learning plays a significant role. Liao and 94 Lyu simulated pseudo-high-energy images from low-energy CT images to 95 improve the quality of the material decomposition with a simple U-Net 96 architecture [11, 12]. Clark et al. used multi-energy CT with DECT and 97 spectral CT for material decomposition with a U-Net-based CNN architecture [13]. Another approach to the crossover architecture that incorporates two 98 99 material generation pathways for the bone(water) density image and water-100 bone (water(bone)) density images was introduced by Zhang et al. [14]. It 101 used both kV-CT images at 80 and 140 kVp that used the DECT scan. These

studies did not directly predict material decomposition images from the singleenergy CT (SECT) images.

104 In a recent study, an image-synthesis technique of cross modality with a 105 generative adversarial network (GAN) was performed. GANs function by 106 training two different networks: a generator network synthesizes an image, and 107 a discriminator network distinguishes between the synthesized and reference 108 images [15]. Florkow et al. proposed an image-synthesis framework of MR 109 images to CT images with a two-dimensional (2D) CNN model [16]. For 110 radiotherapy, the synthesis of PlanCT-like images from Cone beam computed 111 tomography (CBCT) images with planning CT and CBCT datasets with GAN 112 to improve the image quality of the CBCT was introduced in [17]. 113 The current study proposes an image-synthesis approach to material 114 decomposition images of bone(water) and fat(water) reconstructed on DECT

115 from the SECT of an equivalent kilovoltage CT (kV-CT) image at 120 kVp

116 directly using a GAN-based CNN architecture.

117

118

120 **II. MATERIALS AND METHODS**

121

A) Data acquisition 122

123 The DECT image for each patient was acquired with a Revolution DECT 124 scanner (GE Healthcare, Princeton, NJ, USA). The DECT scans were 125 performed at tube voltages of 80 and 140 kVp and exposures of 560 mA. The 126 other scanning parameters were a rotation time (RT) of 1.0 s, slice thickness of 127 0.5 mm, and field of view (FOV) of 360 mm. The material decomposition 128 images of bone(water) and fat(water) and equivalent kV-CT images were 129 reconstructed using the GSI technique. A total of 18,084 images from 28 130 patients were analyzed as part of an institutional review board-approved study. 131

132 **B)** Deep learning model

133 A 16-bit Digital Imaging and Communications in Medicine (DICOM) image 134 was converted to an 8-bit red-green-blue (RGB) portable network graphics 135 (PNG) image, and the output 8-bit RGB PNG image from the 2D CNN model 136 was converted to 16-bit DICOM images. The pixel number in the CT image 137 ranged from -1000 to 3079 Hounsfield units (HU). The unused pixel value was 138 eliminated. Subsequently, the values of the pixels in the CT images were 139 converted to 8-bit (0-255) images by dividing by 16, which is the value 140 obtained by dividing the maximum pixel value, that is, 3079 HU, by 256. The 141 process of radiomics analysis is presented in Fig. 1. The pixel values of the 142 DECT and kV-CT images were rescaled using the RescaleSlope and143 RescaleIntercept tags from the DICOM header as follows:

144 Image Data = (Image Data) × RescaleSlope + RescaleIntercept +
145 1000. (1)

Before calculating the radiomics features, we applied a medium smooth filter 146 147 to the rescaled image data. An overview of the GAN network model is 148 displayed in Fig. 1. This includes a generator (to estimate the material 149 decomposition image) and discriminator (to distinguish the real material 150 decomposition image from the generated image). The generator attempts to 151 produce realistic images that confuse the discriminator. These CNN networks 152 are trained simultaneously by evaluating $\theta_{G,D}$. The generator comprised an 153 encoder and decoder. The encoder mapped the image from the input image of 154 512×512 resolution using a stack of eight convolutional layers, each followed by LeakyReLU activation functions and batch normalization. The number of 155 156 convolutional filters was 64, 128, 256, 512, 1024, 1024, 1024, and 1024, 157 respectively, with a kernel size of 4×4 and stride of 2. The decoder mirrored 158 the encoding architecture, albeit utilizing fractionally strided convolution 159 (deconvolution). The number of the deconvolution filters were 1024, 1024, 160 1024, 1024, 512, 256, 128, and 64, respectively, in the layers that utilized ReLU 161 activation functions. The discriminator used seven convolution layers to extract 162 the features from the image and generate the output image. The number of 163 convolutional filters was 64, 128, 256, 512, 1024, 1024, and 1024, respectively, 164 with a kernel size of 4×4 and stride of 2.

The deep learning model was a conditional GAN that required paired images from the kV-CT and DECT images that were co-registered with voxel-wise correspondence. The label was the kV-CT image before synthesis. The loss was evaluated using the generator and discriminator.

169

170
$$\mathcal{L}_{GAN}(G, D) = \mathbb{E}_{y}[\log D(y)] + \mathbb{E}_{x,z}[\log(1 - D(G(x))] \quad (2)$$

171

where *G* is the generator network, \mathbb{E} is the expectation value dependent on both *x*, the set of kV-CT images, and *y*, the set of target images that are DECT images. Moreover, it included an additional loss based on the absolute difference between the synthesized DECT image and input kV-CT image (L1 norm loss). The L1 norm loss was calculated as

177

178
$$\mathcal{L}_{L1}(G) = \mathbb{E}_{x,y}(|y - G(x)|_1).$$
 (3)

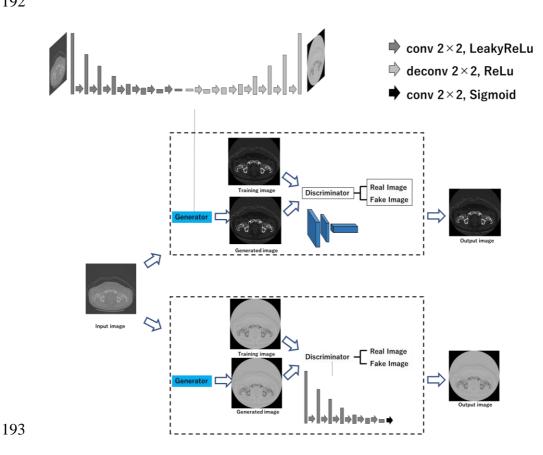
179

180 The adversarial loss in Eq. (1) was calculated using the binary cross-entropy 181 cost function. The final cost function used to optimize the network was a 182 weighted summation of the losses in Eq. (2) and Eq. (3):

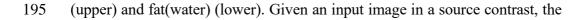
183
$$\theta_{G,D} = \arg \underbrace{\min}_{G} \underbrace{\max}_{D} \left(\mathcal{L}_{L1}(G) + \mathcal{L}_{GAN}(G,D) \right).$$
(4)

184 Here, hyperparameter lambda is the weighting factor for the L1 loss, which

185 was set to 100. The proposed models were implemented using TensorFlow 186 packages (V1.7.0, Python 2.7, CUDA 9.0) on a Ubuntu 16.04 LTS system. Three hundred epochs were used to operate the model on an 11-GB NVIDIA 187 188 GeForce GTX 1080 GPU. All three models were trained with instance 189 normalization and identical hyperparameters, except for the batch size. For 190 each iteration, a mini-batch of 2D images was randomly selected from the 191 training set.



194 Fig. 1 GAN framework of the material decomposition images of bone(water)



196 generator learns to generate an image of similar anatomy in a target contrast 197 and the discriminator learns to discriminate between the synthesized and real 198 pairs of the material decomposition images.

199

200

201

202 C) Evaluation

The prediction accuracy of the model for the synthesized and reference material decomposition images of bone(water) and fat(water) was evaluated using the following five metrics: relative mean absolute error (MAE), relative root mean square (RMSE), structural similarity index (SSIM), signal-to-noise ratio (PSNR), and mutual information (MI). These metrics are defined as follows:

209

210
$$MAE = \frac{1}{n_x n_y} \sum_{i,j}^{n_x n_y} \frac{|r(i,j) - t(i,j)|}{r(i,j)}.$$
 (5)

211

Here, r(i,j) is the value of pixel (i,j) in the planning CT image, t(i,j) is the value of pixel (i,j) in the target image, and $n_x n_y$ is the total number of pixels. RMSE is defined as

215
$$RMSE = \sqrt{\frac{1}{n_x n_y} \sum_{i,j}^{n_x n_y} \left(\frac{r(i,j) - t(i,j)}{r(i,j)}\right)^2}.$$
 (6)

The SSIM is computed based on consideration of the contrast, structure, andluminance to compute a similarity score between two images.

218 The SSIM between two images \vec{x} and \vec{y} can be computed as [18]

219
$$SSIM(\vec{x}, \vec{y}) = \frac{(2\mu_x \mu_y + C_1)(2\sigma_{xy} + C_2)}{(\mu_x^2 + \mu_x^2 + C_1)(\sigma_x^2 + \sigma_y^2 + C_2)},$$
(7)

220
$$C_1 = (k_1 Q)^2, \quad k_1 = 0.01,$$
 (8)

221
$$C_2 = (k_2 Q)^2, \quad k_2 = 0.03,$$
 (9)

where C_1 and C_2 are constants used to prevent a zero denominator and to maintain the stability of the formula. Q is the maximum CT value for the synthesized and reference images. The values of k_1 and k_2 are typically obtained from [19]. σ_x is an estimate in the discrete form

226
$$\sigma_x = \left(\frac{1}{N-1}\sum_{i=1}^N (x_i - \mu_x)^2\right)^{1/2}.$$
 (10)

227 The correlation coefficient between \vec{x} and \vec{y} is defined as σ_{xy} . It is 228 expressed as

229
$$\sigma_{xy} = \frac{1}{N-1} \sum_{i=1}^{N} (x_i - \mu_x) (y_i - \mu_y), \qquad (11)$$

230 where μ_x is the mean intensity and can be expressed as

231
$$\mu_x = \frac{1}{N} \sum_{i=1}^N x_i.$$
 (12)

232 The PSNR is calculated as

233
$$PSNR_{GL} = 10 \times log_{10} \left(\frac{(MAX)^2}{MSE}\right).$$
(13)

Here, MAE and MSE are the possible maximum signal intensity and mean
square error (or difference) of the image, respectively. MI is used as a crossmodality similarity measure [20]. It is calculated as

238

239
$$I(r:t) = \sum_{m \in I_r} \sum_{n \in I_t} p(m,n) \log\left(\frac{p(m,n)}{p(m)p(n)}\right), \tag{14}$$

where *m* and *n* are the intensities in the targeted monochromatic energy CT image I_r and predicted monochromatic energy CT image I_t , respectively. p(m, n) is the joint probability density of I_r and I_t , whereas p(m) and p(n) are marginal densities. p(m, n) can be calculated as follows:

244
$$p(m,n) = \frac{h(m,n)}{\sum_{m \in I_t} \sum_{n \in I_t} h(m,n)},$$
 (15)

where h(m, n) is the histogram of the pixel values in the reference monoenergetic CT image I_r and synthesized monoenergetic CT image I_t . Furthermore, the difference in the synthesized and reference monoenergetic CT numbers in the region of interest (ROI) was evaluated for several slices, from the feet to chest, in a manually drawn ROI, as depicted in Fig. 2.

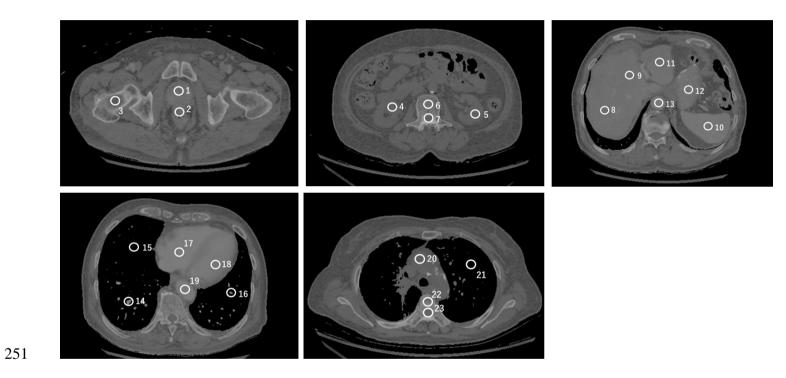
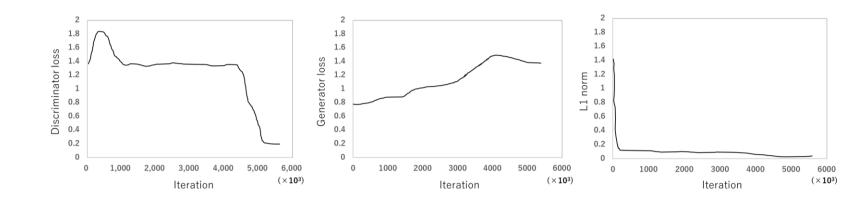


Fig. 2 Method of measurement in the evaluation of the HU in the material decomposition images from feet to chest slice. The

average and SD values of the HU were measured by creating a circular ROI, 2 cm in diameter.

255 III. RESULTS

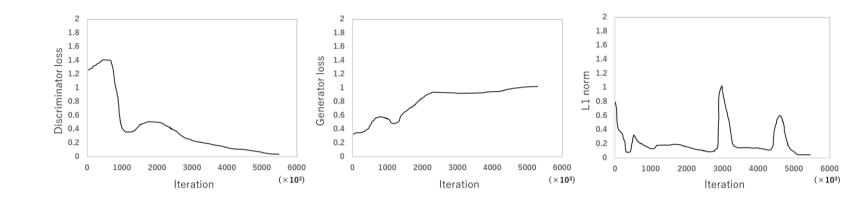
| 257 | The time required to create the image synthesized model was approximately |
|-----|--|
| 258 | 142.2 ± 3.1 h for conversion of the kV-CT to the material decomposition |
| 259 | images of the bone(water) and fat(water). The rate to create the synthesized |
| 260 | monochromatic energy CT images using all the trained models was |
| 261 | approximately 7.2-8.1 images/s. The generator loss, discriminator loss, and L1 |
| 262 | norm loss in each prediction model are displayed in Figs. 3 and 4. |



263

264 Fig. 3 Average training losses in the generator and discriminator in the CT-based prediction model for conversion of kV-CT to

265 the material decomposition images of bone(water).



267

268 Fig. 4 Average training losses in the generator and discriminator in the CT-based prediction model for conversion of kV-CT to

269 the material decomposition images of fat(water).

271 Figs. 5-10 display samples obtained by cross-modality generation for the 272 synthesized and reference material decomposition images of bone(water) and 273 fat(water). Table 1 presents the difference in HU values between the 274 synthesized and reference material decomposition images of bone(water). The 275 difference between the synthesized and reference material decomposition 276 images is within 5.3 HU. The difference of the monochromatic energy CT 277 number is within the appropriate range of the SD values in all ROIs. Table 2 278 indicates the HU difference between the synthesized and reference material 279 decomposition images of fat(water). The difference of the synthesized and 280 reference material decomposition images is within 20.3 HU for the fat(water) 281 images. The difference of the monochromatic energy CT number is within the 282 appropriate range of the SD values in all ROIs. The RMSE in all ROIs was 0.6 283 for the material decomposition image of bone(water) and 1.2 for the material 284 decomposition image of fat(water).

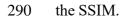
Tables 3 and 4 display the average MAE, MSE, RMSE, PSNR, and MI

286 computed from feet to chest slices for the material decomposition images of

287 bone(water) and fat(water). The MAE, MSE, and RMSE were less for the

288 material decomposition image of fat(water). The PSNR and MI were greater

for the material decomposition image of fat(water). There was no difference in



292

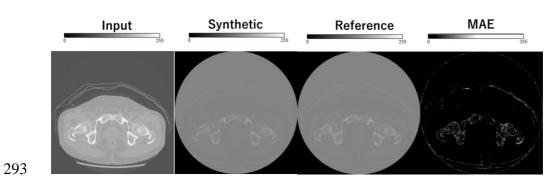


Fig. 5 Samples obtained from a material decomposition image of bone(water) at pelvic level: input image is the equivalent kV-CT image at 120 kVp, synthesized and reference images are the material decomposition images of bone(water), and MAE is the difference between the synthesized and reference bone(water) images.

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300

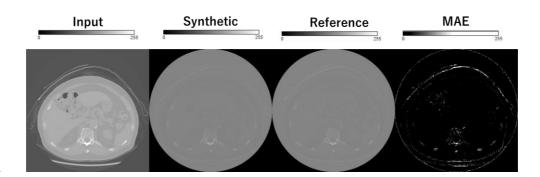


Fig. 6 Samples obtained from a material decomposition image of bone(water)
at abdominal level: input image is the equivalent kV-CT image at 120 kVp,
synthesized and reference images are the material decomposition images of
bone(water), and MAE is the difference between the synthesized and reference
bone(water) images.

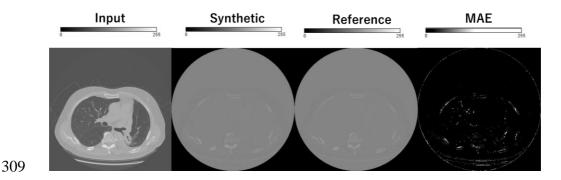


Fig. 7 Samples obtained from a material decomposition image of bone(water)
at chest level: input image is the equivalent kV-CT image at 120 kVp,
synthesized and reference images are the material decomposition images of
bone(water), and MAE is the difference between the synthesized and reference
bone(water) images.

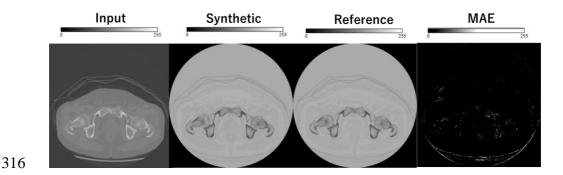


Fig. 8 Samples obtained from a material decomposition image of fat(water) at pelvic level: input image is the equivalent kV-CT image at 120 kVp, synthesized and reference images are the material decomposition images of fat(water), and MAE is the difference between the synthesized and reference fat(water) images.

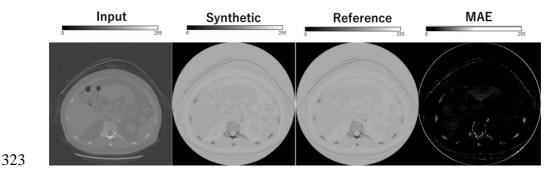
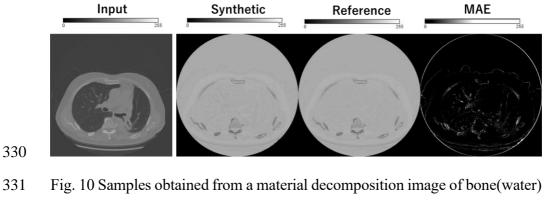


Fig. 9 Samples obtained from a material decomposition image of fat(water) at abdominal level: input image is the equivalent kV-CT image at 120 kVp, synthesized and reference images are the material decomposition images of fat(water), and MAE is the difference between the synthesized and reference fat(water) images.



at pelvic level: input image is the equivalent kV-CT image at 120 kVp,
synthesized and reference images are the material decomposition images of
fat(water), and MAE is the difference between the synthesized and reference
fat(water) images.

Table 1. Difference (Δ) and SD values of synthesized and reference material

- 340 decomposition of bone(water).

| | Bone(water) | |
|-----|---------------|---------|
| | Δ (HU) | SD (HU) |
| (1) | 1.7 | 5.0 |
| 2 | 0.8 | 11.5 |
| 3 | 3.0 | 26.1 |
| 4 | -1.8 | 5.2 |
| 5 | -2.2 | 4.1 |

| $\widehat{0}$ 1.211.7 $\widehat{7}$ -1.15.84 $\widehat{8}$ 1.43.9 $\widehat{9}$ 2.74.5 $\widehat{10}$ 3.04.8 $\widehat{11}$ -0.95.1 $\widehat{12}$ 2.94.4 $\widehat{13}$ 4.64.8 $\widehat{14}$ -0.14.5 $\widehat{15}$ 3.25.7 $\widehat{16}$ 5.36.2 $\widehat{17}$ -0.34.1 $\widehat{18}$ -3.24.3 $\widehat{19}$ -0.14.4 $\widehat{20}$ 1.33.6 $\widehat{21}$ -4.614.3 $\widehat{22}$ 3.415.5 $\widehat{23}$ -1.15.1 | | | |
|---|----------------|------|------|
| (\$) 1.4 3.9 (9) 2.7 4.5 (10) 3.0 4.8 (11) -0.9 5.1 (12) 2.9 4.4 (13) 4.6 4.8 (14) -0.1 4.5 (15) 3.2 5.7 (16) 5.3 6.2 (17) -0.3 4.1 (18) -3.2 4.3 (19) -0.1 4.4 (20) 1.3 3.6 (21) -4.6 14.3 (22) 3.4 15.5 | 6 | 1.2 | 11.7 |
| (9) 2.7 4.5 (10) 3.0 4.8 (11) -0.9 5.1 (12) 2.9 4.4 (13) 4.6 4.8 (14) -0.1 4.5 (15) 3.2 5.7 (16) 5.3 6.2 (17) -0.3 4.1 (18) -3.2 4.3 (19) -0.1 4.4 (20) 1.3 3.6 (21) -4.6 14.3 (22) 3.4 15.5 | \overline{O} | -1.1 | 5.84 |
| | 8 | 1.4 | 3.9 |
| 11-0.9 5.1 12 2.9 4.4 13 4.6 4.8 14 -0.1 4.5 15 3.2 5.7 16 5.3 6.2 17 -0.3 4.1 18 - 3.2 4.3 19 -0.1 4.4 20 1.3 3.6 21 -4.6 14.3 22 3.4 15.5 | 9 | 2.7 | 4.5 |
| 12 2.9 4.4 13 4.6 4.8 14 -0.1 4.5 15 3.2 5.7 16 5.3 6.2 17 -0.3 4.1 18 -3.2 4.3 19 -0.1 4.4 20 1.3 3.6 21 -4.6 14.3 22 3.4 15.5 | 10 | 3.0 | 4.8 |
| 13 4.6 4.8 14 -0.1 4.5 15 3.2 5.7 16 5.3 6.2 17 -0.3 4.1 18 -3.2 4.3 19 -0.1 4.4 20 1.3 3.6 21 -4.6 14.3 22 3.4 15.5 | (11) | -0.9 | 5.1 |
| $ \begin{array}{c ccccccccccccccccccccccccccccccccccc$ | 12 | 2.9 | 4.4 |
| (15) 3.2 5.7 (16) 5.3 6.2 (17) -0.3 4.1 (18) -3.2 4.3 (19) -0.1 4.4 (20) 1.3 3.6 (21) -4.6 14.3 (22) 3.4 15.5 | 13 | 4.6 | 4.8 |
| 165.36.2 17 -0.34.1 18 -3.24.3 19 -0.14.4 20 1.33.6 21 -4.614.3 22 3.415.5 | 14) | -0.1 | 4.5 |
| $ \begin{array}{cccccccccccccccccccccccccccccccccccc$ | 15 | 3.2 | 5.7 |
| 18 -3.2 4.3 19 -0.1 4.4 20 1.3 3.6 21 -4.6 14.3 22 3.4 15.5 | 16 | 5.3 | 6.2 |
| (19) -0.1 4.4 (20) 1.3 3.6 (21) -4.6 14.3 (22) 3.4 15.5 | 17 | -0.3 | 4.1 |
| 20 1.3 3.6 21 -4.6 14.3 22 3.4 15.5 | 18 | -3.2 | 4.3 |
| (2) -4.6 14.3 (2) 3.4 15.5 | 19 | -0.1 | 4.4 |
| 22 3.4 15.5 | 20 | 1.3 | 3.6 |
| | | -4.6 | 14.3 |
| 23 -1.1 5.1 | | 3.4 | 15.5 |
| | 23 | -1.1 | 5.1 |

345 Table 2. Difference (Δ) and SD values between synthesized and reference

346 material decomposition of fat(water).

| | Fat(water) | |
|----|---------------|---------|
| | Δ (HU) | SD (HU) |
| 1) | -5.3 | 9.3 |
| 2 | -5.0 | 8.2 |
| 3 | -7.4 | 7.8 |
| 4 | -3.6 | 31.5 |

| 5 | -6.7 | 12.9 |
|------------|------|-------|
| 6 | -8.9 | 14.8 |
| \bigcirc | -2.9 | 9.8 |
| 8 | -5.9 | 9.7 |
| 9 | 4.0 | 17.6 |
| 10 | -0.7 | 18.1 |
| (1) | 0.0 | 58.5 |
| 12 | -3.6 | 9.1 |
| 13 | -6.9 | 8.9 |
| 14 | -8.1 | 9.2 |
| 15 | -5.9 | 20.8 |
| 16 | -3.1 | 8.6 |
| 17 | -6.4 | 10.6 |
| 18 | -1.7 | 12.3 |
| 19 | 5.3 | 12.5 |
| 20 | -3.0 | 42.9 |
| (21) | 7.5 | 182.0 |
| (22) | -7.0 | 51.9 |
| 23 | -8.8 | 93.1 |

Table 3. Average MAE, MSE, RMSE, PSNR, and SSIM computed from feet to chest slices for the material decomposition images of bone(water).

354

| MAE | | MSE | | RMSE | | PSNR | | SSIM | | MI | |
|---------|-----|---------|-----|---------|-----|---------|------|---------|-----|---------|-----|
| Average | SD | Average | SD | Average | SD | Average | SD | Average | SD | Average | SE |
| 0.8 | 0.7 | 1.3 | 1.2 | 0.9 | 0.7 | 55.3 | 15.0 | 0.9 | 0.1 | 0.8 | 0.1 |

Table 4. Average MAE, MSE, RMSE, PSNR, and SSIM computed from feet to chest slices for the material decomposition
images of fat(water).

| MAI | Ξ | MSE | 3 | RMS | E | PSNR | | NR SSIM | | MI | |
|---------|-----|---------|-----|---------|-----|---------|-----|---------|-----|---------|-----|
| Average | SD |
| 0.0 | 0.0 | 0.0 | 0.0 | 0.1 | 0.0 | 72.1 | 3.4 | 0.9 | 0.0 | 1.4 | 0.1 |

361 IV. DISCUSSION

362

DECT enables the separation of several additional materials including calcium, 363 364 fat, and uric acid from a single kV-CT. It provides anatomic knowledge with 365 functional information [21]. DECT requires post-processing to obtain material 366 decomposition images. It requires 5 to 10 min of additional interpretation time 367 after the scan [22]. Conversely, the proposed image-synthesized system can 368 reconstruct a DECT image within 1 min automatically. GE scanners use dual 369 X-ray sources; Siemens Healthiness scanners use dual X-ray sources and two 370 data acquisition systems [23, 24]. Thus, the dual-source scanner cost is 371 considerably greater than the standard SECT scanner. The Philips Healthcare 372 scanner acquires DECT projection data using a layered detector. The high-373 energy data and low-energy data are collected by the posterior and anterior 374 detector layers, respectively [25]. All DECT data acquisition techniques pose 375 a significant burden on the CT system hardware. Zhao et al. reported that, due 376 to this fact, DECT scanners are not widely used in less-developed regions [26]. 377 In addition to the increased cost and complexity of the imaging system, DECT 378 can also increase the radiation dose owing to the additional CT scan. The 379 image-synthesized approach with deep learning is expected to reduce the 380 scanning radiation dose and imaging cost by synthesizing the DECT from the 381 single kV-CT image.

| - | T decomposition that |
|--|---|
| 383 combines conventional penalized weighted-least squa | ares estimation with |
| 384 regularization based on a mixed union of learned transfo | orms model [27]. The |
| 385 RMSE of the density image was 0.04–0.05. To compare th | ne RMSE on the same |
| 386 scale, the material decomposition image was converted to | a density image with |
| 387 a theoretical value. Here, the theoretical density values | were 1 g/cm ^{3} for the |
| 388 water and 1.92 g/cm^3 for the bone used by Li. The RMSE | t in all ROIs was 0.55 |
| 389 for the material decomposition image of bone(water) and | d 1.2 for the material |
| 390 decomposition image of fat(water). The RMSE of the den | sity value in all ROIs |
| 391 was 0.01 for the material decomposition image of bone(w | vater) and 0.02 for the |
| 392 material decomposition image of fat(water), which was s | significantly less than |
| 393 that of Li <i>et al</i> . Although other studies have proposed the | image synthesis of a |
| 394 DECT, the accuracy of the image synthesis could not b | be directly compared |
| 395 because there were no studies evaluating the HU value. | . Moreover, previous |
| 396 studies used other reconstruction images such as low- and | d high-energy kV-CT, |
| 397 multi-energy images, and virtual non-contrast images [1 | 1–13]. These models |
| 398 required multiple images or additional reconstruction | . The current study |
| 399 proposed a prediction model of the material decom | nposition images of |
| 400 bone(water) and fat(water) from a single kV-CT image with | th GAN architectures. |
| 401 The HU difference between the reference and s | synthesized material |
| 402 decomposition images of bone(water) and fat(water) were | e less than 5.3 HU and |
| 403 20.3 HU, respectively. The material decomposition image | es of the bone(water) |

404 had a smaller MAE, MSE, and RMSE, and a greater PSNR and MI than the 405 fat(water). The bone was highlighted and the other organ with a similar density to water in the material decomposition images of the bone(water), were less 406 407 prominent. For the U-Net-based CNN employed by the previous study, the 408 label was image-wise. Conversely, the model proposed in the current study 409 used the label that was pixel-wise. Although the previous study could extract 410 local imaging features, it was required to register the input and output images 411 for model training. The current study used the kV-CT and material 412 decomposition images reconstructed from the DECT image. No differences 413 were observed in the alignment of the images.

There is a possibility that certain patients could be affected by the beamhardening artifact. This could cause errors in the model training in the correlation of the training of the kV-CT and material decomposition images of the bone(water). However, these differences in each ROI were within the SD range. The proposed model could produce highly accurate DECT images within the noise estimations from the kV-CT images.

There are limitations to the current study. First, the current study used 120 kV-CT images reconstructed from DECT. The difference in the image quality of 120 kV-CT images scanned via SECT and the equivalent 120 kV-CT images reconstructed from DECT for clinical patients were evaluated by Tawfil *et al.* The subjective image quality scores between the DECT and SECT groups did not indicate a significant difference. Thus, 120 kV-CT images from the DECT 426 images can be used as equivalent to the SECT images [28]. Moreover, the 427 patients used for the current study were randomly selected, regardless of the presence or absence of disease. Therefore, evaluation of the lesion detectability 428 429 could not be performed. In addition to evaluating the image similarities for complete images, we confirmed that the HU difference was within the noise in 430 431 the local region from the pelvic to the chest level. Further studies will be 432 performed to examine the quality of the synthesized images compared to the 433 original images in terms of diagnostic performance.

434

435

438 V. CONCLUSION

| 440 | The current study proposed an image-synthesis framework using a GAN-based |
|-----|--|
| 441 | CNN architecture for kV-CT to material decomposition images of bone(water) |
| 442 | and fat(water) scanned by DECT. The proposed image synthesis model showed |
| 443 | a highly image quality and the difference of the monochromatic energy CT |
| 444 | number is within the appropriate range of the SD values in the local region. |
| 445 | Synthesized medical image generation can be a cost-effective approach for |
| 446 | developing automated diagnostic technologies. |
| 447 | |

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538 Figure captions

| 539 | Fig. 1. GAN framework of the material decomposition images of bone(water) |
|-----|--|
| 540 | (upper) and fat(water) (lower). Given an input image in a source contrast, |
| 541 | Generator learns to generate an image of similar anatomy in a target contrast |
| 542 | and Discriminator learns to discriminate between synthesized and real pairs of |
| 543 | the material decomposition images. |

544

Fig. 2 Method of measurement in the evaluation of the HU in the material
decomposition images from feet to chest slice. The average and SD values of
the HU were measured by creating a circular ROI, 2 cm in diameter.

548

Fig. 3 Average training losses in the generator and discriminator in the CTbased prediction model for conversion of kV-CT to the material decomposition
images of bone(water).

552

Fig. 4 Average training losses in the generator and discriminator in the CTbased prediction model for conversion of kV-CT to the material decomposition
images of fat(water).

556

557 Fig. 5 Samples obtained from a material decomposition image of bone(water)558 at pelvic level: input image is the equivalent kV-CT image at 120 kVp,

synthesized and reference images are the material decomposition images of
bone(water), and MAE is the difference between the synthesized and reference
bone(water) images.

562

Fig. 6 Samples obtained from a material decomposition image of bone(water)
at abdominal level: input image is the equivalent kV-CT image at 120 kVp,
synthesized and reference images are the material decomposition images of
bone(water), and MAE is the difference between the synthesized and reference
bone(water) images.

568

Fig. 7 Samples obtained from a material decomposition image of bone(water)
at chest level: input image is the equivalent kV-CT image at 120 kVp,
synthesized and reference images are the material decomposition images of
bone(water), and MAE is the difference between the synthesized and reference
bone(water) images.

574

Fig. 8 Samples obtained from a material decomposition image of bone(water)
at pelvic level: input image is the equivalent kV-CT image at 120 kVp,
synthesized and reference images are the material decomposition images of
bone(water), and MAE is the difference between the synthesized and reference
bone(water) images.

Fig. 9 Samples obtained from a material decomposition image of bone(water) at abdominal level: input image is the equivalent kV-CT image at 120 kVp, synthesized and reference images are the material decomposition images of bone(water), and MAE is the difference between the synthesized and reference bone(water) images.

586

Fig. 10 Samples obtained from a material decomposition image of bone(water)
at pelvic level: input image is the equivalent kV-CT image at 120 kVp,
synthesized and reference images are the material decomposition images of
bone(water), and MAE is the difference between the synthesized and reference
bone(water) images.