### Quantitative Assessment of the Normal and Abnormal Achilles Tendon in vivo Using a 3D Cones Sequence

Hongda Shao<sup>1</sup>, Michael Carl<sup>2</sup>, Eric Chang<sup>1</sup>, Christine B Chung<sup>1</sup>, Graeme M Bydder<sup>1</sup>, and Jiang Du<sup>1</sup>

<sup>1</sup>Radiology, University of California, San Diego, CA, United States, <sup>2</sup>GE Healthcare, San Diego, CA, United States

#### INTRODUCTION

Tendon is a highly ordered collagen-rich fibril tissue links muscle to bone. Rupture is a common event in both sedentary and athletic population. Typical histopathology features include marked inflammation, angiogenesis, collagen degeneration and disordered arrangement of collagen fibers  $^1.$  Multiple water components with distinct MR relaxation times exist in the tendon. Magnetization transfer (MT) sequences have been employed to probe macromolecules in long T2 tissues  $^2.$  However, conventional clinical MT sequences cannot detect MT effects in tendon which has a very short T2. Ultrashort echo time (UTE) sequences with TEs less than 100  $\mu s$  can potentially detect the multiple water components in tendon, and evaluate the magnetization transfer ratio (MTR) by combining UTE sequences with a MT preparation pulse  $^{3-5}.$  In this study, we aimed to study the multiple water components in the Achilles tendon, and investigate its MTR as a function of MT pulse frequency offset in healthy volunteers and patients with tendon rupture using a clinical 3T scanner.

#### MATERIALS AND METHODS

Two patients (male, age = 41 and 59 years) were recruited in this preliminary study, and both previously had a normal Achilles' tendon and an abnormal Achilles' tendon that had been surgically repaired after a rupture (volunteer A got tendon ruptured 10 years previously, and volunteer B got tendon ruptured two and a half months previously). Both of the Achilles' tendon of each volunteers were scanned with a 3D Cones sequence for T2\* and T1 measurement, and a 3D Cones-MT imaging for MTR measurement using a 3T GE whole-body scanner. The 3D Cones sequence employed a short rectangular pulse for excitation followed by Cones sampling with a nominal TE of 32  $\mu$ s. The 3D Cones-MT sequence employed a Fermi pulse (8 ms, maximal saturation flip angle of 670°) for

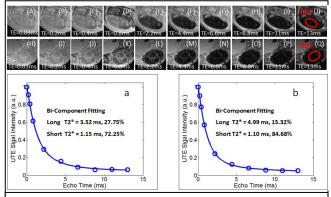
MT preparation. The 3D UTE-MT imaging protocol used the following parameters: TR = 100ms, field of view (FOV) = 10cm, matrix = 256×256, band width = 125kHz, four echoes with TEs of 0.032, 4.3, 8.6, and 12.9 ms, three off-resonance frequencies ( $\Delta f = 1.5$ , 3.0 and 5.0 kHz). Bicomponent T2\* analysis was performed on five interleaved dual-echo 3D Cones data (TEs = 0.032/4.4; 0.2/6.6; 0.4/8.8; 0.8/11; 2.2/13 ms) with a total scan time of 10 min. The effective T1 of both short and long T2 components was measured with a variable TR 3D Cones approach (TR=5.8, 10, 20, 40, 60, 100 ms) ranging from 10 ms to 100 ms). The T1 of the short T2\* components was measured with a 3D IR-Cones approach, using five sets of TR/TI combinations (TR/TI=107/48, 150/64, 200/81, 300/110 and 400ms/131ms) were performed, with each TR/TI combination appropriate to invert and null the long T2 components  $^6$ . A 3-in receive only coil was used for signal reception. The bi-component

analysis algorithm was written in MATLAB and executed offline on Dicom images obtained using the protocols described above. MTR corresponding to different  $\Delta fs$  was calculated using ImageJ software.

## RESULTS AND DISCUSSION

**Figure 1** shows bi-component T2\* analysis of the interleaved dual-echo 3D Cones images. The long T2\* component increased from T2\* $\sim$ 3.52 for the normal tendon to  $\sim$ 4.99ms for the abnormal tendon. T2\* of the short T2\* component remained essentially unchanged (1.15 vs 1.10 ms). The fraction of the long T2 component decreased from 28% for the normal tendon to 15% for the abnormal tendon.

**Figure 2** shows the T1 increased from 362.72 ms for the normal tendon to 464.41 ms for the abnormal tendon.



**Fig 1** UTE-T2\* images of both normal tendon (A-J) and diseased tendon (H-Q) of a 59 year old male volunteer with TEs ranging from 32μs to 13ms, and bi-component fitting of an ROI in the normal tendon (a) and the diseased tendon (b), those shows 28% long T2\* component (T2\*~3.5ms) and 72% short T2\* component (T2\*~1.1ms) for normal tendon, and 15% long T2\* component (T2\*~5ms) and 95% short T2\* component (T2\*~1.10ms) for the diseased tendon.

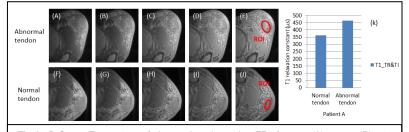


Fig 2 3D Cones T1 mapping of abnormal tendon with a TR of 10 ms (A), 20 ms (B) , 40 ms (C), 60 ms (D), 100 ms (E), and normal tendon with a TR of 10 ms (F), 20 ms (G), 40 ms (H), 60 ms (I), 100 ms (J). Abnormal tendon shows  $\sim$ 30% higher T1 (K).

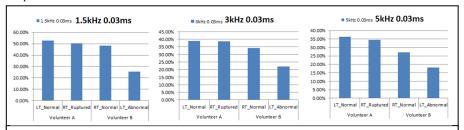


Fig 3 UTE-MT images with different TEs and different off-resonance frequencies of the normal tendon in volunteer A shows excellent detail of its fiber structure (not displayed). The MTR was decreased in the diseased tendon compared with the normal tendon at different  $\Delta f$  (1.5, 3, 5 kHz) with a TE of 32  $\mu s$ .

**Figure 3** shows 3D Cones-MT imaging of the normal tendon of volunteer acquired at different frequency offsets and TEs (images not displayed). The 3D Cones-MT sequence provided high quality morphological images with high signal and resolution, as well as MTR values of the tendon. The MTR chart shows the ruptured tendon has a lower MTR value which especially in the fresh ruptured tendon (volunteer B) at all the different frequency offsets and TEs.

The bi-component T2\* results are a little different from our previous study of volunteer A using a 2D UTE bi-component analysis <sup>3</sup>. This might be caused by multiple factors: tendon changes during the interval of ~2 years, and differences between the 2D UTE and the 3D Cones sequences (e.g., 2D UTE sequence is more sensitive to eddy currents had a longer half-pulse and a shorter rectangular pulse was used in 3D Cones). The histopathologic change in diseased tendon is a complex process<sup>4</sup>. Tendon is known to have multi-components. A bi-component T2\* model is a simplification <sup>5</sup>. Our T1 values are also lower than published values <sup>4</sup>, potentially related to errors in flip angle, which plays a key role in T1 quantification using variable TR approach. Further research is needed to resolve this discrepancy.

# CONCLUSIONS

The study shows differences in long T2\*s and fractions, T1 and MTR between normal and diseased tendons. MTR differences were especially obvious with ultrashort TEs, suggesting that UTE-MTR may be a robust method in evaluating the process of tendon degeneration and recovery.

**REFERENCES** 1. FUKUTA, S., ET AL, MATRIX BIOLOGY 1998; 17(1): 65–73. 2. GROSSMAN RI, RADIOGRAPHICS 1994. 4. Filho GH, et al. AJR 2009; 192:117-124. 5. Grosse U, et al., Magn Reson Med 2013; 70:184-192.

3. ERIC D, ET AL, NMR BIOMED; 2012; 25: 161-168. 6. Du J, et al. PlosOne 2014; e0103296.