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1 Short communication:

2 **Development of a protocol to quantify local bone**
3 **adaptation over space and time: quantification of**
4 **reproducibility**
5

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1 **Abstract**

2 *In vivo* micro-computed tomography (μ CT) scanning of small rodents is a
3 powerful method for longitudinal monitoring of bone adaptation. However, the
4 life-time bone growth in small rodents makes it a challenge to quantify local
5 bone adaptation. Therefore, the aim of this study was to develop a protocol,
6 which can take into account large bone growth, to quantify local bone
7 adaptations over space and time. The entire right tibiae of eight 14-week-old
8 C57BL/6J female mice were consecutively scanned four times in an *in vivo*
9 μ CT scanner using a nominal isotropic image voxel size of 10.4 μ m. The
10 repeated scan image datasets were aligned to the corresponding baseline (first)
11 scan image dataset using rigid registration. 80% of tibia length (starting from
12 the endpoint of the proximal growth plate) was selected as the volume of
13 interest and partitioned into 40 regions along the tibial long axis (10 divisions)
14 and in the cross-section (4 sectors). The bone mineral content (BMC) was used
15 to quantify bone adaptation and was calculated in each region. All local BMCs
16 have precision errors ($PE_{\%CV}$) of less than 3.5% (24 out of 40 regions have
17 $PE_{\%CV}$ of less than 2%), least significant changes (LSCs) of less than 3.8%, and
18 38 out of 40 regions have intraclass correlation coefficients (ICCs) of over 0.8.
19 The proposed protocol allows to quantify local bone adaptations over an entire
20 tibia in longitudinal studies, with a high reproducibility, an essential
21 requirement to reduce the number of animals to achieve the necessary statistical
22 power.

23

24 **Keywords:** *in vivo* micro-CT; local bone adaptation; mouse tibia; space and
25 time

26

1 **1. Introduction**

2 Bone adaptation is a process in which bone undergoes adaptive changes.
3 While bone keeps its strength through balanced resorption and formation,
4 disorder of bone adaptation can lead to bone diseases, such as osteoporosis,
5 osteomalacia, Paget's disease, etc. [Britton and Walsh, 2012;Shih 2012].
6 Small rodents offer a cost-effective and efficient way for the investigation of
7 bone diseases in preclinical studies. In addition, the development of *in vivo* high
8 resolution micro-computed tomography (μ CT) scanning on the entire bone of
9 small rodents offers a powerful approach to quantify bone adaptations over
10 space and time [Altman et al., 2015; Birkhold et al., 2014; Lambers et al.,
11 2013; Lu et al., 2015]. To quantify bone adaptations, three-dimensional (3D)
12 bone morphometric measurements (trabecular thickness, trabecular separation,
13 cortex thickness, etc.) over a volume of interest (VOI) (proximal mouse tibia,
14 tibial midshaft, etc.) were used [Bouxsein et al., 2010; Campbell et al., 2014;
15 Lambers et al., 2013;Nishiyama et al., 2010]. Although 3D image registration
16 can improve the long-term precision of these measurements [Campbell et al.,
17 2014], these morphometric measurements were averaged values over a region
18 of an entire bone and can hardly be used to quantify local bone adaptations over
19 the entire bone's space. On the contrary, *in vivo* μ CT images obtained at the
20 same anatomical site over different time points were superimposed using the
21 rigid registration, and then bone formation and resorption were quantified from
22 the superimposed images [Birkhold et al., 2014; Schulte et al., 2011].
23 However, in rodents like mouse, bone growth spans across the animal's life
24 time [Glatt et al., 2007], and should be taken into account when interpreting the
25 data [Birkhold et al., 2014]. This is particular true for long bones (e.g. tibia),
26 where changes in length due to growth can be significant. In this case, it may
27 still be valid to quantify bone formation and resorption over a short time
28 interval with rigidly registered images [Birkhold et al., 2014], but this approach
29 would fail in a longer time interval (e.g. 2 weeks) due to the significant shift and

1 changes of bone structure caused by bone growth. Therefore, in this study, a
2 novel protocol that aims to account for large bone growth was proposed to
3 quantify the local bone adaptation over a larger volume of interest (80% of
4 mouse tibia) and over space and time.

5

6 **2. Material and methods**

7 **2.1 Animals**

8 The detailed information on animals can be found in Lu *et al.* [2015]. In
9 summary, eight 14-week-old female C57BL/6J (BL6) mice were used and the
10 mice were well housed before the experiment. All the procedures were
11 approved by the local Research Ethics Committee of the University of Sheffield
12 (Sheffield, UK).

13 **2.2 *In vivo* μ CT scanning**

14 The details of the *in vivo* μ CT scanning were in Lu *et al.* [2015]. In summary,
15 the entire right tibia of every mouse was scanned four times consecutively (the
16 scanning of each tibia took approximately 40 minutes) with an *in vivo* μ CT
17 system (vivaCT 80, Scanco Medical, Bruettisellen, Switzerland) at 14-week-old.
18 For the duration of the scanning, the mice were placed on a heating pad,
19 maintained under anaesthetic gases (isoflurane). Between each scan, the mouse
20 (kept under anaesthesia) was repositioned in the sample holder to simulate a
21 longitudinal study design. The scanner was operated at 55 keV, 145 μ A, an
22 integration time of 200 ms and a nominal isotropic image voxel size of 10.4 μ m.
23 The radiation dose from the μ CT scanning was estimated to be approximately
24 500 mGy for each scan, which has been proved to cause no significant effect on
25 bone adaptations [Laperre *et al.*, 2011].

26 **2.3 Image processing and calculation of bone parameters**

27 In the image processing chain, first, an alignment procedure was defined so
28 that all tibiae, regardless of their positions in the scanner, were aligned to the

1 same anatomical reference system. In the alignment procedure, the tibia from
2 the baseline scan was taken as the reference, referred as baseline scan 1
3 thereafter. The tibia of the baseline scan 1 (**Figure 1a**) was placed back into its
4 anatomical position, i.e. the long axis of the tibia was approximately aligned
5 with the global z axis and the y-z plane passed through the centre line of the
6 articular surfaces of the medial and lateral condyles (**Figure 1b**). The tibiae
7 from the repeated scans and from other mice were rigidly registered and
8 transformed to the transformed tibia of baseline scan 1 (**Figure 1d**) and then
9 resampled using the Lanczos kernel [**Turkowski and Gabriel, 1990**].

10 After the image transformation, the tibial length (L) was measured as the
11 distance from the most proximal tibial bone pixel until the most distal tibial
12 bone pixel. Afterwards, a region of 80% of L (**Figure 1d**), starting from the end
13 of the proximal growth plate [**Klinck et al., 2008**] was cropped out [Amira
14 5.4.3, FEI Visualization Sciences Group, France]. Then the tibial VOI was
15 extracted by removing the proximal part of fibula (**Figures 1e and 1f**) (Matlab
16 v2015a, the Mathworks, Inc. USA).

17 To investigate the spatial adaptation of the Bone Mineral Content (BMC),
18 the tibial VOI was partitioned into 40 sub-volumes. In the tibial longitudinal
19 (proximal-distal) direction, the tibial VOI was divided into 10 regions (**Figure**
20 **1e**). In the tibial transverse (x-y) section, a polar coordinate system was created
21 for each image slice. The system was originated at the centre of mass of each
22 slice and the x-axis was defined from the tibial medial side towards the lateral
23 side (**Figure 1g**). In the tibial transverse section, the tibia was then divided into
24 4 sectors (anterior, medial, posterior and lateral sectors), starting from the
25 position that is 45 degrees away from the x-axis (**Figure 1g**).

26 To calculate BMC in each sub-volume, firstly, the grayscale VOI datasets
27 were smoothed with a Gaussian filter (convolution kernel [3 3 3], standard
28 deviation = 0.65) and then binarised into bone and background using a fixed
29 single level threshold, i.e. 25.5% of maximal grayscale value (around 420 mg

1 HA/ccm) [**Klinck et al., 2008**], close to the values applied in other studies
 2 performed on mouse bone with the same image resolution [**Birkhold et al.,**
 3 **2014; Lambers et al., 2015; Lukas et al., 2013**]. The BMC in each sub-
 4 volume was calculated as the volume of image voxel times the Bone Mineral
 5 Density (BMD) values summed over all bone voxels. In addition, the cortical
 6 and trabecular compartments were separated [**Buie et al., 2007**] and the total
 7 BMC for each bone type [Ct.BMC and Tb.BMC] was calculated.

8 Following the standard procedure, the bone morphometric measurements
 9 were quantified in order to ensure the quality of the images by comparing with
 10 literature data. Trabecular bone volume fraction (Tb.BV/TV), trabecular
 11 thickness (Tb.Th), trabecular separation (Tb.Sp), trabecular number (Tb.N)
 12 were computed in the region (Region 1 in **Figure 1c**) extending 1.00 mm
 13 distally from the growth plate, with an offset of 0.20 mm from the most distal
 14 break in the calcified cartilage bridge of the growth plate observed in the
 15 grayscale CT slice [**Nishiyama et al., 2010; Klinck et al., 2008**]. Cortex
 16 thickness (Ct.th) was calculated in a 1.00mm region centred at the tibial mid-
 17 shaft (Region 2 in **Figure 1c**).

18 **2.4 Statistical analysis**

19 The reproducibility of the global and local bone mineral content variables
 20 was characterized by the precision errors (PEs) [**Glueer et al., 1995**], the least
 21 significant change (LSCs) [**Burghardt et al., 2013; Shepherd and Lu, 2007**]
 22 and the intraclass correlation coefficients (ICCs) [**Schrouf and Fleiss, 1979**].
 23 PEs were expressed as the coefficients of variation (CV) (PE_{%CV}).

$$24 \text{ PE}_{\%CV} = \sqrt{\sum_{j=1}^m \%CV_j^2 / m} \quad (1)$$

25 with

$$26 \%CV_j = \frac{SD_j}{\bar{x}_j} \times 100\% \quad (2)$$

1 where, m is the subject number (m = 8 in the current study) and \bar{x}_j is the mean
2 of all x_{ij} for subject j.

3 The LSC was calculated as follows:

$$4 \text{ LSC} = Z \times PE_{\%CV} \sqrt{\frac{1}{n_1} + \frac{1}{n_2}} \quad (3)$$

5 where, Z-score corresponds a two-tailed 95% confidence level (Z=1.96), while
6 n_1 and n_2 are the number of measures performed at baseline (n_1) and follow-up
7 (n_2), respectively.

8 The ICC is the ratio of the between-subject variance divided by the
9 population variance [Nishiyama et al., 2010].

$$10 \text{ ICC} = \frac{F_0 - 1}{F_0 + (n - 1)} \quad (4)$$

11 where, F_0 is the ratio of between-subject mean squares over the residual within-
12 subject mean squares and n is the number of repetitions (n = 4 in this study).

13

14 **3. Results**

15 The tibial length has a $PE_{\%CV}$ of 0.11%, a LSC of 0.13% and an ICC of 0.99.
16 Bone morphometric parameters have $PE_{\%CV}$ ranging from 0.49% (Ct.Th) to 3.59%
17 (Tb.BV/TV), LSC from 0.56% to 4.14%, and ICCs from 0.93 (Tb.Sp) to 0.99
18 (Ct.Th), the values of which are comparable to the data in literature (**Figure 2**).
19 The Ct.BMC and Tb.BMC have $PE_{\%CV}$ of 1.58% and 3.04%, LSC of 1.82% and
20 3.51%, and ICCs of 0.95 and 0.98, respectively.

21 Regarding the local BMC measurements, 24 out of 40 regions (60%) have
22 $PE_{\%CV}$ less than 2%, 15 regions (37.5%) between 2% and 3%, and one region
23 (2.5%) with 3.2% (**Figure 3**). LSCs for the 40 regions were less than 3.80%,
24 ranging from 1.46% to 3.78%. 29 out of 40 regions (72.5%) have ICCs over
25 0.90, 9 regions (22.5%) between 0.80 and 0.90 (**Figure 3**). With respect to the
26 anatomical location of the tibia, there is no spatial variability pattern for the
27 reproducibility.

1 The mean \pm SD values of tibial morphometric measurements, global and
2 local BMC measurements, and their precision errors and ICCs are reported in
3 **Appendix A, Table A.1.**

4 5 **4. Discussion**

6 In this study, a novel protocol, which can take into account large bone
7 growth, was developed to quantify local bone adaptations over space and time.
8 High precision and reproducibility of the local BMC measurements, calculated
9 through the protocol, were found. Although the reproducibility values cannot be
10 directly migrated to the images obtained from other μ CT systems or other voxel
11 size scans with the same system [Verdelis et al., 2011], this paper proposed a
12 protocol to evaluate local bone adaptations over space and time and this
13 protocol is irrespective of the μ CT systems and μ CT voxel size.

14 The proposed protocol was made efficient by selecting 80% of the tibial
15 length as the VOI to represent the entire tibia. The tibial growth plates were
16 excluded, which not only cause noise and errors in the calculation of BMC, but
17 also impede the automation of the protocol. The BMC was selected as the
18 parameter to quantify local bone adaptation, but other parameters, e.g.
19 periosteal/endocortical perimeters, bone marrow area, etc. [Bouxsein et al.,
20 2010], can be quantified using the protocol developed in this study.

21 In this paper, the tibial VOI was partitioned into 40 sub-volumes. Our
22 preliminary investigations showed there was a conflict between the desire to
23 quantify bone adaptation with the highest possible spatial resolution, and the
24 need for highly reproducible measurements. We found that the partitioning
25 proposed is a reasonable compromise between these two conflicting needs and
26 that smaller compartments would provide less reproducible measurements
27 (Table A.2 in the Appendix), and larger compartments would not further reduce
28 it while losing spatial resolution.

1 When investigating bone adaptations using rodents, there are essentially two
2 scenarios: the first is that bone undergoes significantly low growth during the
3 experiment (e.g. adult rat or caudal vertebra of adult mouse with the scanning
4 interval of one week) [Altman et al., 2015; Birkhold et al., 2014]. In such case,
5 the voxel difference between the superposed images transformed with the rigid
6 registration [Birkhold et al., 2014; Lambers et al., 2013; Schulte et al., 2011]
7 or the distance vectors between the bone iso-surfaces [Lu et al., 2015] can be
8 interpreted as bone formation and resorption. The second is that bone undergoes
9 continuous relatively large growth (e.g. mouse tibia) during the experiment
10 (**Figure 4**). In such case, the previous methods would produce erroneous results
11 and the bone changes could be measured by using a full elastic registration
12 approach that could be adapted from [Dall'Ara et al., 2014]) or the affine
13 scaling of anatomically referenced partitioning, which was applied in the
14 current study. Our preliminary investigation (Table A.2 in the Appendix)
15 showed that when using smaller size of compartments over which the BMC is
16 averaged would considerably degrade the reproducibility of the measurements.
17 This strongly suggested that when dealing with the bone with large growth, the
18 image voxel-level comparisons need to be replaced with the protocol proposed
19 in this study, which can take into account the relatively large bone growth.

20 In conclusion, a novel protocol, which can take into account large bone
21 growth, was developed to quantify local bone adaptation over space and time
22 and high reproducibility of the local BMC measurements was found. In the
23 future, the protocol can be used in longitudinal image datasets to quantify local
24 bone adaptation over space and time.

25

26 **Conflict of interest statement**

27 The authors have no conflicts to declare.

28

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8

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1 **Figure 1.** Overview of the image processing chain in the reproducibility study.

2

3 **Figure 2.** Reproducibility of mouse cortical and trabecular parameters (tibial
4 length, tibial cortex BMC, trabecular BMC and tibial morphometric parameters)
5 expressed in precision error as coefficients of variation ($PE_{\%CV}$) and the 95%
6 confident intervals ($CI_{95\%}$) shown in terms of error bars, and the intraclass
7 correlation coefficients (ICC) are reported in square brackets (8 mice and 4
8 scans per mouse).

9

10 **Figure 3.** Reproducibility of the mouse tibial local BMC expressed in mean
11 precision error as coefficients of variation ($PE_{\%CV}$) and the 95% confident
12 intervals ($CI_{95\%}$) shown in terms of error bars, the least significant change (LSC)
13 and the intraclass correlation coefficients (ICC) are reported in square brackets
14 (8 mice and 4 scans per mouse) (C01 to C10 corresponds tibial proximal to
15 distal side, see **Figure 1e**).

16

17 **Figure 4.** Superimposition of two mouse tibia sections (a and b) using the rigid
18 registration and visualisation of bone adaptations (d). Over one week, there is
19 little common regions in the trabecular part due to the relatively large growth

1 **Appendix A – the complete statistical data:**

2 To quantify the reproducibility of the variables, in addition to the precisions
3 errors as coefficients of variations ($PE_{\%CV}$) and the intraclass correlation
4 coefficients (ICCs), the precision errors as the standard deviation (PE_{SD}) and
5 confidence intervals (CIs) of $PE_{\%CV}$ were also quantified. The PE_{SD} is defined as
6 below:

$$7 \quad PE_{SD} = \sqrt{\sum_{j=1}^m SD_j^2 / m} \quad (A.1)$$

8 where, m is the subject number ($m = 8$ in the current study) and SD_j is the
9 standard deviation of subject j .

10 CIs were to determine how accurate the PEs were and were determined for
11 each of the $PE_{\%CV}$ values using a chi-squared distribution (χ^2).

$$12 \quad \frac{df}{\chi_{1-\frac{\alpha}{2}, df}^2} PE_{\%CV}^2 < \sigma^2 < \frac{df}{\chi_{\frac{\alpha}{2}, df}^2} PE_{\%CV}^2 \quad (A.2)$$

13 where, df is the total degrees of freedom ($df = 24$ in the current study).

14 The mean \pm SD values of tibial morphometric measurements, global and
15 local BMC measurements, and their precision errors (PE_{SD} , $PE_{\%CV}$ and $CI_{95\%}$)
16 and ICCs are reported in **Table A.1**.

17 The reproducibility data ($PE_{\%CV}$ and ICC) for tibial local BMC at smaller
18 compartments (80 compartments: 20 divisions in tibial long axis and 4 divisions
19 in tibial cross-section) are presented in **Table A.2**. In case of 80 compartments,
20 38 out of 80 regions (47.5%) have $PE_{\%CV}$ less than 2% (compared to 60% for
21 the case of 40 compartments), 38 regions (47.5%) between 2% and 3%
22 (compared to 37.5%), and 4 regions (5.0%) over 3% (compared to 2.5%); 59 out
23 of 80 regions (73.75%) have ICCs over 0.90 (compared to 72.5% for the case of
24 40 compartments), 13 regions (16.25%) between 0.80 and 0.90 (compared to
25 22.5%), and 8 regions (10%) below 0.80 (compared to 0.0%).

26

1 **Table A.1.** The mean \pm standard deviation (SD) values of the tibial bone
2 mineral content (BMC) (eight mice and four scans for each mouse), of the
3 corresponding reproducibility data (PE_{%CV}: precision error of the coefficient of
4 variation; CI_{95%}: 95% confidence interval of PE_{%CV}; LSC: least significant
5 change for PE_{%CV} ; ICC: intraclass correlation coefficient) (C01L, C01A, C01M
6 and C01P represent the lateral, anterior, medial and posterior regions of
7 compartment 01, as shown in **Figure 1**, respectively)

		Mean \pm SD	PE (CV%)	CI _{95%} (%)	LSC (%)	ICC
Tibial length [mm]		16.83 \pm 0.17	0.11	0.08 – 0.14	0.13	0.989
Ct.BMC [μg HA]		7753.95 \pm 504.26	1.58	1.28 - 2.07	1.82	0.947
Tb.BMC [μg HA]		29.80 \pm 6.00	3.04	2.47 - 4.00	3.51	0.979
Morp homet ric param eters	Tb.BV/TV [%]	4.80 \pm 0.90	3.59	2.92 – 4.73	4.14	0.967
	Tb.N [1/mm]	2.69 \pm 0.322	2.86	2.32 – 3.77	2.30	0.948
	Tb.Th [μm]	45.67 \pm 3.424	1.65	1.34 – 2.18	1.90	0.956
	Tb.Sp [μm]	380.00 \pm 47.41	3.45	2.08 – 4.54	3.98	0.931
	Ct.Th [μm]	172.28 \pm 6.54	0.49	0.40 – 0.65	0.56	0.985
	C01L	263.73 \pm 42.29	2.48	2.01 – 3.26	2.86	0.978
	C02L	207.52 \pm 28.90	2.02	1.64 – 2.65	2.33	0.981
	C03L	165.71 \pm 26.79	2.45	1.99 – 3.23	2.82	0.979
	C04L	142.94 \pm 23.21	3.01	2.45 – 3.97	3.47	0.969
	C05L	156.16 \pm 21.63	2.16	1.75 – 2.84	2.49	0.978
	C06L	152.98 \pm 18.74	1.98	1.61 – 2.61	2.28	0.976
	C07L	154.78 \pm 21.73	3.27	2.65 – 4.30	3.77	0.951
	C08L	133.41 \pm 12.09	2.33	1.89 – 3.07	2.67	0.940
	C09L	160.14 \pm 9.92	1.98	1.60 – 2.60	2.28	0.907
	C10L	242.07 \pm 13.30	2.72	2.21 – 3.59	3.14	0.773
	C01A	277.87 \pm 22.38	2.43	1.97 – 3.19	2.80	0.917
	C02A	286.81 \pm 18.21	1.97	1.60 – 2.59	2.27	0.912
	C03A	306.92 \pm 17.81	1.75	1.42 – 2.31	2.02	0.917
	C04A	285.71 \pm 13.30	1.56	1.27 – 2.06	1.80	0.897

Local BMC [μg HA]	C05A	223.22 \pm 11.58	1.57	1.27 – 2.06	1.81	0.917
	C06A	173.02 \pm 7.52	1.43	1.16 – 1.88	1.65	0.902
	C07A	192.22 \pm 8.63	1.44	1.17 – 1.90	1.66	0.906
	C08A	174.29 \pm 12.37	1.96	1.59 – 2.58	2.26	0.931
	C09A	147.21 \pm 9.13	1.66	1.34 – 2.18	1.91	0.935
	C10A	162.73 \pm 8.55	1.81	1.47 – 2.39	2.09	0.891
	C01M	212.08 \pm 18.17	1.74	1.41 – 2.29	2.01	0.963
	C02M	205.08 \pm 18.01	1.87	1.51 – 2.46	2.16	0.959
	C03M	149.70 \pm 11.94	2.34	1.90 – 3.08	2.70	0.921
	C04M	148.93 \pm 8.62	2.38	1.93 – 3.13	2.74	0.845
	C05M	155.93 \pm 4.01	2.20	1.79 – 2.90	2.54	0.787
	C06M	163.93 \pm 5.71	1.53	1.24 – 2.01	1.76	0.823
	C07M	181.87 \pm 12.53	1.68	1.36 – 2.21	1.94	0.946
	C08M	165.75 \pm 8.87	2.10	1.70 – 2.76	2.42	0.859
	C09M	150.99 \pm 10.25	1.98	1.61 – 2.61	2.28	0.923
	C10M	180.65 \pm 12.02	2.28	1.85 – 3.00	2.63	0.893
	C01P	236.02 \pm 23.97	1.79	1.45 – 2.35	2.06	0.972
	C02P	234.03 \pm 24.08	1.71	1.39 – 2.26	1.97	0.975
	C03P	246.84 \pm 28.31	1.63	1.32 – 2.14	1.88	0.982
	C04P	233.42 \pm 21.61	1.46	1.19 – 1.93	1.68	0.977
C05P	190.46 \pm 14.83	1.27	1.03 – 1.68	1.46	0.976	
C06P	167.41 \pm 7.24	1.28	1.04 – 1.68	1.47	0.921	
C07P	224.66 \pm 11.10	2.16	1.76 – 2.85	2.49	0.824	
C08P	190.27 \pm 10.96	2.08	1.69 – 2.74	2.40	0.880	
C09P	145.36 \pm 6.82	1.96	1.59 – 2.58	2.26	0.840	
C10P	132.12 \pm 10.23	2.28	1.85 – 3.00	2.63	0.921	

1

2 **Table A.2.** Reproducibility data (PE_{%CV}: precision error of the coefficient of
3 variation and ICC: intraclass correlation coefficient) for tibial local BMC [μg
4 HA] at smaller compartments (80 compartments: 20 divisions in tibial long axis
5 and 4 divisions in tibial cross-section)

Compartment	Lateral		Anterior		Medial		Posterior	
	PE (CV%)	ICC	PE (CV%)	ICC	PE (CV%)	ICC	PE (CV%)	ICC
C01	2.76	0.973	2.48	0.914	1.92	0.961	1.82	0.971

C02	2.16	0.985	2.41	0.930	1.64	0.966	1.79	0.973
C03	2.05	0.982	2.10	0.929	1.71	0.971	1.78	0.965
C04	1.99	0.979	1.86	0.907	2.24	0.952	1.68	0.981
C05	2.23	0.980	1.80	0.931	2.39	0.878	1.63	0.982
C06	2.77	0.977	1.79	0.908	2.40	0.947	1.64	0.982
C07	3.25	0.967	1.64	0.907	2.40	0.889	1.52	0.978
C08	2.81	0.970	1.58	0.905	2.59	0.794	1.49	0.975
C09	2.25	0.977	1.54	0.925	2.46	0.507	1.35	0.980
C10	2.09	0.979	1.64	0.909	2.04	0.559	1.23	0.968
C11	2.00	0.977	1.56	0.906	1.64	0.757	1.27	0.914
C12	1.97	0.977	1.34	0.900	1.50	0.873	1.30	0.941
C13	2.98	0.955	1.85	0.874	1.88	0.927	3.31	0.916
C14	3.75	0.944	1.47	0.955	1.76	0.955	2.08	0.903
C15	2.61	0.950	1.81	0.948	1.92	0.909	2.34	0.910
C16	2.25	0.915	2.28	0.896	2.58	0.767	2.12	0.754
C17	2.14	0.925	1.98	0.920	2.29	0.905	2.25	0.803
C18	2.33	0.838	1.61	0.939	2.19	0.909	1.98	0.837
C19	2.48	0.859	2.08	0.890	2.49	0.886	2.07	0.935
C20	3.13	0.668	1.88	0.889	2.40	0.879	2.61	0.906

Figure 1

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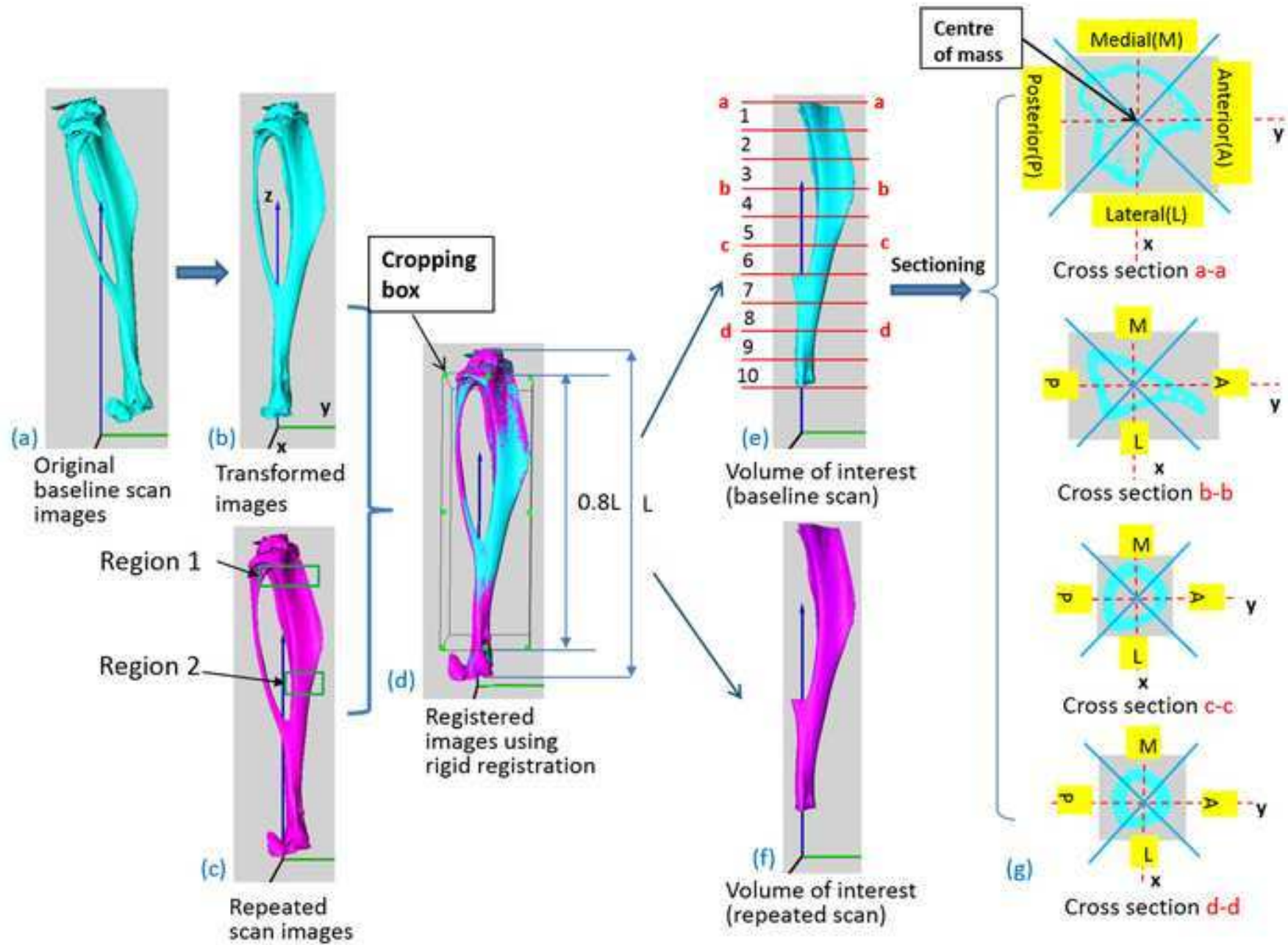


Figure 2

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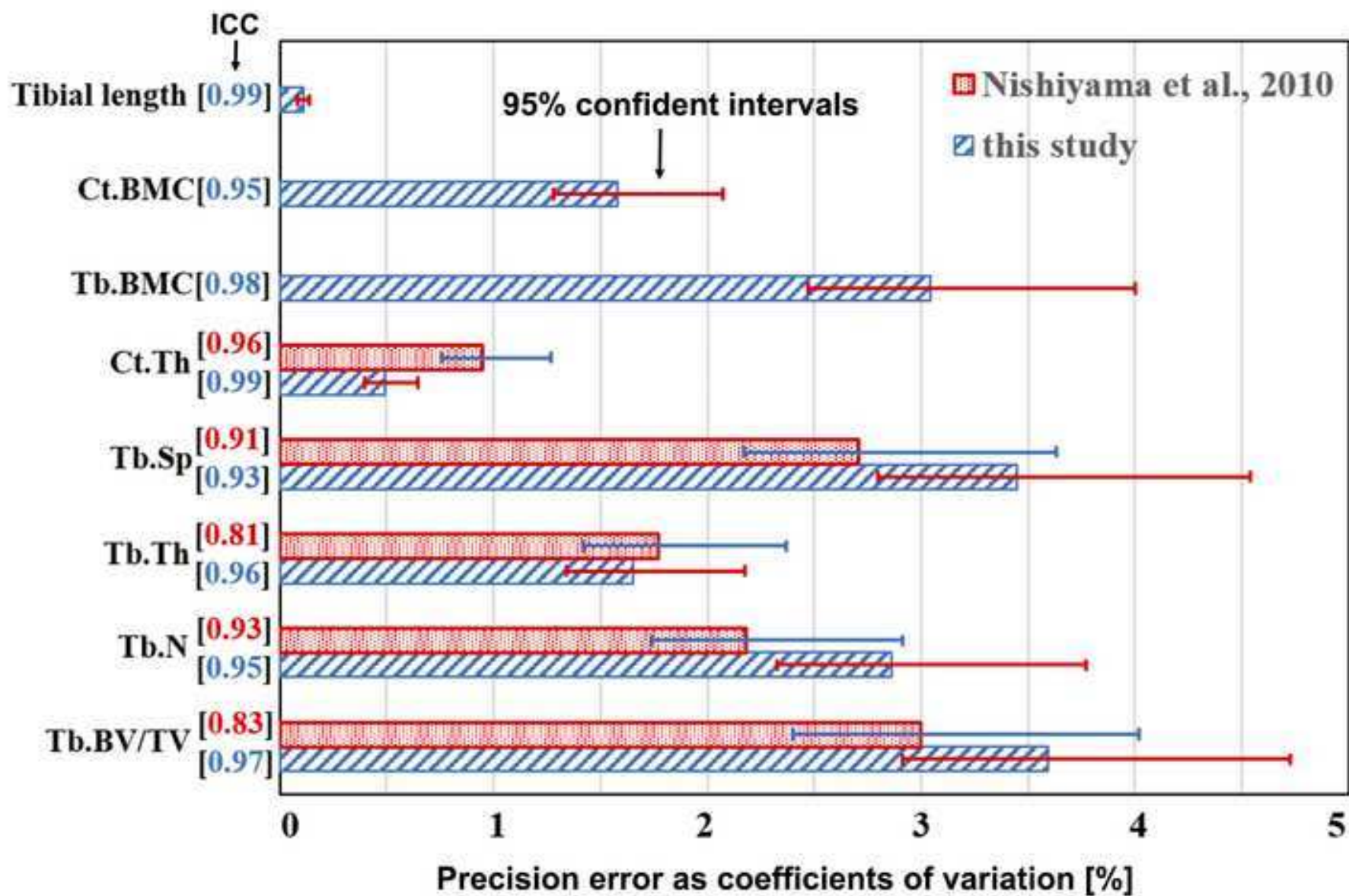


Figure 3

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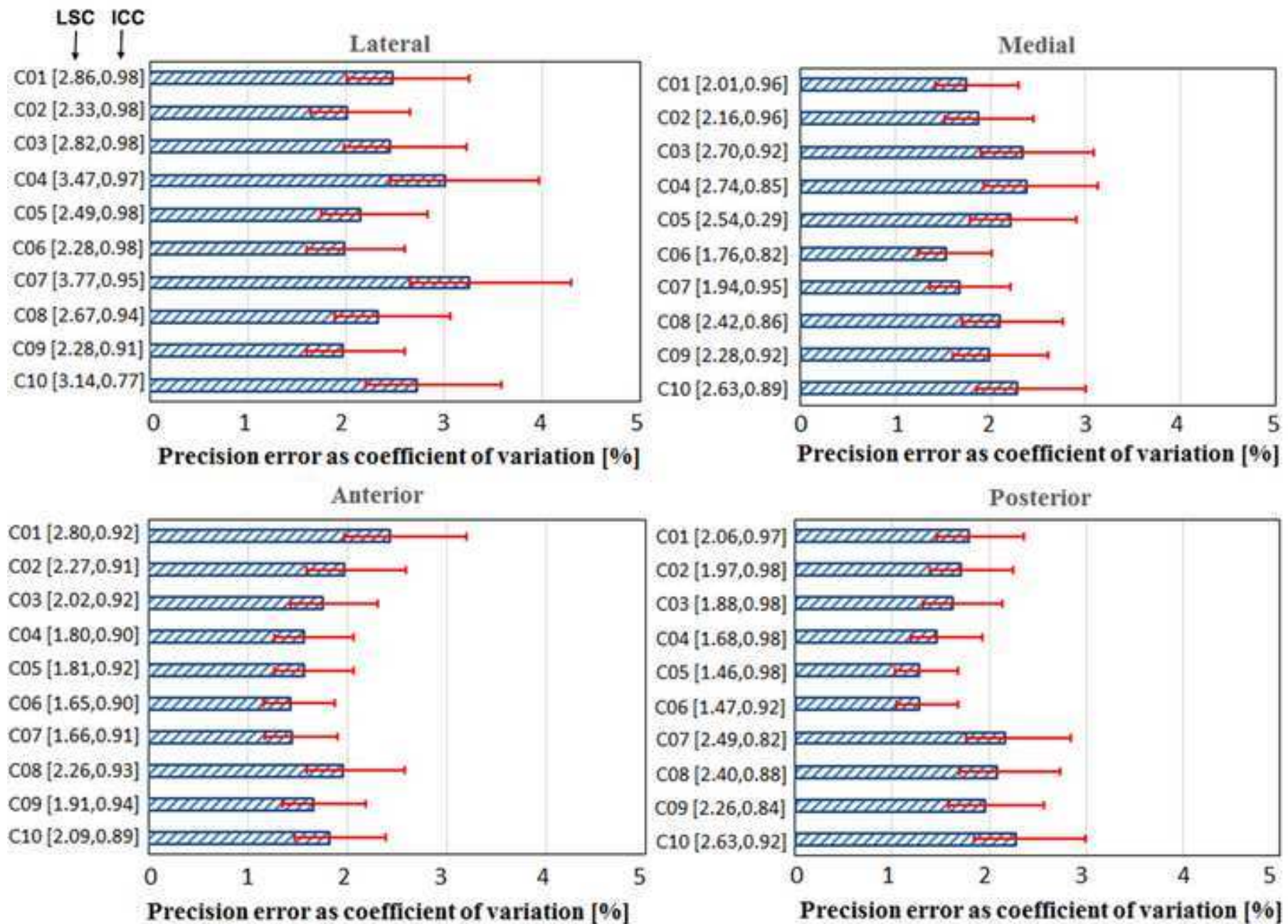


Figure 4

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