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2 Scanner influence on the mechanical response of QCT-based finite
3 element analysis of long bones

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32 **Abstract**

33 **Purpose:** Patient-specific QCT-based finite element (QCTFE) analyses enable highly accurate
34 quantification of bone strength. We evaluated CT scanner influence on QCTFE models of long
35 bones.

36 **Methods:** A femur, humerus, and proximal femur without the head were scanned with K_2HPO_4
37 phantoms by seven CT scanners (four models) using typical clinical protocols. QCTFE models
38 were constructed. The geometrical dimensions, as well as the QCT-values expressed in Hounsfield
39 unit (HU) distribution was compared. Principal strains at representative regions of interest (ROIs),
40 and maximum principal strains (associated with fracture risk) were compared. Intraclass
41 correlation coefficients (ICCs) were calculated to evaluate strain prediction **reliability** for different
42 scanners. Repeatability was examined by scanning the femur twice and comparing resulting
43 QCTFE models.

44 **Results:** Maximum difference in geometry was 2.3%. HU histograms before phantom calibration
45 showed wide variation between QCT scans; however, bone density histogram variability was
46 reduced after calibration and algorithmic manipulation. Relative standard deviation (RSD) in
47 principal strains at ROIs was <10.7%. ICC estimates between scanners were >0.9. Fracture-
48 associated strain had 6.7%, 8.1%, and 13.3% maximum RSD for the femur, humerus, and proximal
49 femur, respectively. The difference in maximum strain location was <2 mm. The average
50 difference with repeat scans was 2.7%.

51 **Conclusions:** Quantification of strain differences showed mean RSD bounded by ~6% in ROIs.
52 Fracture-associated strains in "regular" bones showed a mean RSD bounded by ~8%. Strains were
53 obtained within a $\pm 10\%$ difference relative to the mean; thus, in a longitudinal study only changes

54 larger than 20% in the principal strains may be significant. ICCs indicated high reliability of
55 QCTFE models derived from different scanners.

56 **Key words** QCT; Femur; Finite element analysis; Humerus; Patient-specific; Personalized
57 medicine

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60 **Introduction**

61 Personalized finite element (FE) models of bones based on quantitative computed tomography
62 (QCT) have been used extensively to estimate bone stiffness and strength. Among these, the human
63 femur and humerus attracted significant interest (Dahan et al., 2016; Dall'Ara et al., 2013; Enns-
64 Bray et al., 2016; Hazrati Marangalou et al., 2012; Helgason et al., 2014; Keaveny et al., 2008;
65 Keyak et al., 2005; Keyak et al., 1990; Nishiyama et al., 2013; Pise et al., 2009; Schileo et al.,
66 2008a; Trabelsi et al., 2011; Yosibash et al., 2007b) because of their importance for fracture risk
67 prediction due to osteoporosis, optimal management of bony metastases, and more. QCT-based
68 FE (QCTFE) analysis is considered one of the best current techniques for noninvasive assessment
69 of femoral and humeral strength (Keaveny et al., 2010).

70 Validation of QCTFE predictions has been performed by comparing predicted strain on bone
71 surface and fracture load to these measured in ex vivo experiments (Dahan et al., 2016; Trabelsi
72 et al., 2011; Yosibash et al., 2007b). Most validation studies were based on QCT scans performed
73 on a single scanner; however, to introduce QCTFE into clinical practice for longitudinal studies in
74 diverse clinical centers, scanner influence on QCTFE predictions must be quantified. QCTFE
75 models are directly affected by QCT data: (a) bone geometry is reconstructed from the set of points
76 sampled in the scan; (b) bone density in each voxel, a proxy for material properties, is derived
77 from CT-values expressed in Hounsfield unit (HU). Additionally, indirect effects are present due
78 to algorithmic manipulations of QCT data: (a) boundary corrections and density averaging are
79 performed at the voxel level to overcome the partial volume effect and to reduce noise; (b) material
80 properties are determined using empirical relationships based on HUs and assigned to the FE
81 model at specific locations (integration points).

82 The feasibility of introducing QCTFE into clinical use was recently addressed in a study examining
83 QCTFE model dependence on scan protocols (Dragomir-Daescu et al., 2015). The paper
84 concluded that predicted strength and stiffness of QCTFE models generated from ‘high’ and ‘low’
85 resolution scans could be different if CT settings and reconstruction techniques vary. A second
86 study (Giambini et al., 2015) found that material property estimation may be dependent on scan
87 parameters such as voltage and current as well as post-processing techniques, including the
88 reconstruction kernel, with the differences potentially affecting estimates of strength and stiffness.
89 The basic assumption that QCTFE models are independent of CT scanner was questioned based
90 on a comparison of density and strength evaluated with two different scanners (Carpenter et al.,
91 2014). The study concluded that significant inter-scanner differences remain even after using
92 phantom calibration.

93 This study compares patient-specific QCTFE models using data obtained with multiple CT
94 scanners and hospitals, while considering the steps in the QCTFE model construction pipeline. We
95 analyzed scans of two human femurs and one humerus surrounded by five K_2HPO_4 phantoms.
96 QCT scans were performed using standard clinical protocols. We hypothesized that a carefully
97 designed and implemented pipeline would allow development of reliable QCTFE models of
98 human bones, *i.e.* models that are virtually independent of the CT scanner used to acquire data.

99 **Materials and Methods**

100 A femur from a 70-year-old female (51 kg, 170 cm), a humerus from a 57-year-old male (55 kg,
101 180 cm) (National Disease Research Institute, Philadelphia, PA), and a proximal femur without
102 the head and neck from a 55-year-old male (73 kg, 175 cm) (ScienceCare Anatomical, Phoenix,
103 AZ), were scanned by seven different CT scanners at four medical centers. In total, four CT

104 scanner models from three manufacturers were used: Brilliance (64) and iCT (256) by Philips
105 Healthcare (Eindhoven, The Netherlands), Somatom Definition Edge (128) by Siemens Medical
106 Systems (Erlangen, Germany) and Optima CT660 (128) by GE HealthCare (Milwaukee, WI,
107 USA).

108 Five liquid K_2HPO_4 phantoms (concentrations: 0, 50, 100, 200 and 300 mg/cc), prepared
109 according to (Mindways Software, 2002) were used. The bones and phantoms were immersed in
110 a water bath during all scans (Fig. 1A). Scans were performed according to a standard clinical
111 protocol: exposure 210–500 mAs, X-ray tube voltage 120 kV, and slice thickness and spacing
112 1mm in all protocols except for scan G, where it was 1.25 mm (see Table 2). Images were
113 reconstructed using a “soft/body” filter/convolution kernel (B, B31f, and body filters for Philips,
114 Siemens, and GE scanners, respectively). Details on CT scanners, medical centers and scanning
115 protocols are presented in Tables 1 and 2.

116 *Brief summary of QCTFE pipeline*

117 FE models were generated using the various QCT scans according to a validated procedure
118 (Yosibash et al., 2007a; Yosibash et al., 2007b). Strains, stiffness, and fracture loads computed by
119 QCTFE models were **previously** validated in double-blind in vitro studies, where the modeling
120 pipeline was described in detail (Trabelsi et al., 2011; Yosibash et al., 2014; Yosibash et al., 2010).
121 Briefly, bone boundaries were identified on each slice and surfaces were imported as a point cloud
122 to a computer-aided design software program (SolidWorks by Dassault Systems, Waltham, MA,
123 USA). All QCTFE models were aligned in a uniform coordinate system. This **ensured** that the
124 same boundary conditions were applied and that results were extracted at the same locations in all
125 models. Alignment was performed using an “iterative closest point” algorithm (CloudCompare,

126 <http://www.danielgm.net/cc/>). Geometrical differences between models were due to differences in
127 pixel size and the set of points from which each model was reconstructed. After alignment, these
128 differences were assessed in 9 to 11 cross sections located along the bones at 12–25 mm intervals.
129 Dimensions in x and y directions were extracted and the maximum differences at each section were
130 compared to the average values.

131 HU calibration was performed by K_2HPO_4 phantoms: HU values were averaged along each
132 phantom using a 10×10 mm² area at the center of 10 slices (3000–5800 pixels), see Fig. 1B. Five
133 average HU values (five known densities in phantoms), determined a linear relationship between
134 K_2HPO_4 density and HU: $\rho_{K_2HPO_4} = m \times HU + n$ (in all scans $R^2 > 0.998$). Values for m and n for
135 each scan are given in last two columns of Table 2.

136 Due to the partial volume averaging effect in voxels on bone surface, a boundary correction
137 algorithm was applied. A total width of ~ 1 mm, corresponding to 2 pixels, was usually affected.
138 HU values in all voxels were corrected for noise effects by a moving average algorithm on a cubic
139 volume of 27 bone voxels. Young's modulus (pointwise "stiffness") was thereafter computed
140 based on ash density (ρ_{ash}), using the empirical relationship described in (Schileo et al., 2008a) and
141 the relationship between hydroxyapatite and K_2HPO_4 densities (Goodsitt, 1992):

$$142 \quad \rho_{ash} \left[\frac{gr}{cc} \right] = 1.061 \times \rho_{K_2HPO_4} + 0.08 \quad (1)$$

143 CT scanner output data was compared by inspecting histograms of both raw HU and resulting ρ_{ash}
144 obtained after moving average and boundary correction algorithms. An inhomogeneous Young's
145 modulus (E), based on ρ_{ash} for cortical (Keller, 1994) and trabecular bone (Keyak et al., 1993) was

146 assigned to the QCTFE models, considering only values of $\rho_{ash} > 0 \left[\frac{gr}{cc} \right]$):

$$E_{cort} = 10200 \times \rho_{ash}^{2.01} [MPa], \quad \rho_{ash} \geq 0.486 \left[\frac{gr}{cc} \right] \quad (2)$$

$$E_{trab} = 2398 [MPa], \quad 0.3 < \rho_{ash} < 0.486 \left[\frac{gr}{cc} \right] \quad (3)$$

$$E_{trab} = 33900 \times \rho_{ash}^{2.2} [MPa], \quad \rho_{ash} \leq 0.3 \left[\frac{gr}{cc} \right] \quad (4)$$

147 The Poisson ratio was set to $\nu = 0.3$, as commonly used in bone QCTFE models (Schileo et al.,
148 2014; Varga et al., 2018; Wirtz et al., 2000; Yosibash et al., 2007a).

149 On the femur head, a hip contact force representing a stance position of 1250N was applied parallel
150 to the shaft axis on a 335 mm² area. On the humerus head, a 230N \hat{x} + 650N \hat{z} load was applied on
151 a 320 mm² area with the z axis rotated at 20° to the shaft axis and the x axis pointing laterally. On
152 the proximal femur, a normal displacement $u_n = 0.75$ mm was applied on an area of 810 mm²
153 inclined by 35° to the shaft axis (as illustrated in Fig. 2). All three bones were clamped at the distal
154 surface ($\vec{u} = 0$). A mesh of p -tetrahedral elements with a maximum element volume of ~ 150 mm³
155 was generated. Linear elastic QCTFE analyses were performed by increasing the polynomial
156 degree over elements from $p=1$ to $p=8$, to allow monitoring of numerical errors in energy norm
157 and strains (Szabó and Babüska, 1991). We emphasize the advantages of p -FE methods (Trabelsi
158 et al., 2011; Yosibash et al., 2007b) over conventional FE methods: they accurately represent bone
159 surfaces and polynomial degree (p) of shape functions is increased systematically to achieve
160 convergence. p -FEs also allow a smooth variation of material properties within the model,
161 elements are much larger, may be far more distorted, and produce considerably faster convergence
162 rates.

163 To isolate the effect of the QCT reconstruction algorithm, we also performed FE analyses with a

164 homogeneous Young's modulus, $E=10$ GPa.

165 Comparison between QCTFE models was performed by means of principal strains
166 (compression/tension), as they may serve as a criterion to predict femoral/humeral fracture
167 (Bayraktar et al., 2004; Dahan et al., 2016; Schileo et al., 2014; Schileo et al., 2008b; Yosibash et
168 al., 2010). Regions of interest (ROIs) on the bone surface were arbitrarily selected on the shaft,
169 neck, and intertrochanteric/intertuberosity zones (13, 12, and 8 ROIs on the femur, humerus and
170 proximal femur surfaces respectively, Fig. 2). ROIs were 3 mm lines resembling the commonly
171 used strain gauges (SG) in validation experiments (Dahan et al., 2016; Yosibash et al., 2007b).
172 The average strain value along each line was compared relative to the mean strain value at a given
173 ROI across all QCTFE models from the different scanners. Standard deviation of the strains was
174 calculated for each ROI. Relative standard deviation (RSD [%], also known as coefficient of
175 variation (CV)), defined by the ratio of the standard deviation to the mean strain in the ROI is
176 reported. The mean RSD (averaged across all ROIs) for each bone is also presented. Based on the
177 ROI strains, linear correlations were plotted for each pair of the QCTFE models.

178 To evaluate the reproducibility of results obtained from different scanners, Intraclass Correlation
179 Coefficient (ICC) estimates (two-way random effects, absolute agreement, single rater (Koo and
180 Li, 2016; McGraw and Wong, 1996)) and their 95% confidence intervals (CI) were calculated
181 using SPSS (IBM, Chicago, IL, USA). ICCs were calculated based on strain values in the different
182 ROIs.

183 In addition to ROIs, indicating a global agreement between QCTFE models, the maximum tension
184 and compression principal strains and their locations (surrogate of load to fracture and fracture
185 location) were compared (averaged over a circular surface with a radius of $r=2$ mm). To quantify

186 and compare the obtained locations, a reference point representing the ‘average’ location was
187 computed using least-mean-squares. We report mean, maximum, and RSD of the distance from
188 the reference point.

189 Within-scanner repeatability of the QCTFE was examined by performing two consecutive scans
190 of the femur on each CT scanner. Maximum tension and compression principal strains were
191 computed and compared.

192 **Results**

193 The relative difference in dimension at each cross section is presented in Figure 3. The maximum
194 relative difference in dimensions between all models was less than 2.3%. All QCTFE models had
195 1,170,000–1,390,000 degrees of freedom and converged to <6.5% relative error in energy norm at
196 $p=8$. Strains converged to <1% relative error between consecutive solutions (increased polynomial
197 degree). The influence of the geometrical reconstruction, as examined by the FE models with
198 constant $E=10$ GPa, is presented in Table 3. The maximum RSD in ROI principal strain that can
199 be attributed to the geometry reconstruction algorithm (segmentation process and pixel size in the
200 various QCT scans in the three considered bones) is 5.5%; however, the average RSD is between
201 2.1% and 3.1%. Detailed strain values are provided in the Supplemental Material (Appendix B).

202 Histograms of scans’ output data are shown in Figure 4. Raw HU histograms can be seen in
203 Figure 4A-C; manipulated raw HU resulting in ash density (ρ_{ash}), which is used to determine
204 inhomogeneous material properties is shown in Figure 4D-F. Since one proximal femur was cut
205 and cleaned in a previous study, some of the fat and marrow tissue was lost and replaced by air
206 (HU=-1000), accounting for the many negative values in the proximal femur's histogram.

207 The mean and the RSD% of the absolute maximum principal strains at each ROI for all QCTFE
208 models are presented in Figure 5. These represent all differences in QCTFE models including
209 geometry, material property distribution, and algorithmic influence due to HU manipulation.
210 Maximum RSD in the femur is 7.3%, in the humerus 7.8%, and in the proximal femur 11%;
211 average RSDs are 4.97%, 4.72%, and 6.1%, respectively. Detailed graphs showing the specific
212 values obtained in each FE model are presented in Appendix A. **Correlations between each pair of**
213 **models yielded linear regression equations with slopes varying between 0.93 to 1.05 ($r^2 > 0.96$**
214 **for all pairs), the complete linear correlation matrix is provided in Appendix C.** ICC estimates and
215 their corresponding 95% CIs are 0.959 [0.901, 0.986], 0.988 [0.971, 0.996], and 0.981 [0.952,
216 0.995] for the femur, humerus, and proximal femur, respectively.

217 The location and magnitude of maximum principal strain is the most important information to the
218 medical community because it is associated with risk of fracture. Maximum tensile and
219 compressive principal strains in the femur and proximal femur, and compressive principal strain
220 in the humerus (no tensile significant strains at most loading conditions) are presented in Figure 6.
221 In the femur, a 6.7% RSD was obtained in the maximum tensile principal strain, with the largest
222 maximum difference about 10% from the mean value for scanner A. In the humerus, an 8.1% RSD
223 was obtained in the maximum compressive principal strain, with maximum differences for
224 scanners F and G of about -10% and 10% from the mean, respectively. In the proximal femur, a
225 13.3% RSD was obtained in the maximum compressive principal strain with maximum differences
226 for scanners D and G of about -15% and 15% from the mean value, respectively. The differences
227 in the location of the obtained principal strains compared to the reference point are summarized in
228 Table 4; the maximum difference was <2 mm.

229 Within-scanner differences in the maximum tensile and compressive principal strains was less

230 than 4.1% for six scanners, but was 7.7% for scanner G. The average difference across all seven
231 scanners was 2.7% (standard deviation 2%).

232 **Discussion**

233 This study investigated whether a rigorous QCT protocol combined with a unique pipeline for the
234 generation of a QCTFE model enabled reliable values of principal strain in three human bones,
235 regardless of the CT scanner used for image acquisition. Principal strains are quantities used to
236 estimate risk of fracture and indirectly assess bone stiffness, so are of special interest.

237 Bone dimensions from the various QCT scans showed very small differences (maximum relative
238 difference <2.3%). These differences introduced a maximum RSD of 5.5% in the ROI principal
239 strains for the three bones when homogeneous material properties were used. The average
240 differences were much smaller, usually half of the maximum RSD.

241 Considering the material properties, the cortex (high HU value) dominates the mechanical
242 behavior of bones. Discrepancy in low HUs has a negligible influence and is deemed unimportant.
243 For example, a tissue having HU <200 ($\rho_{ash} < 0.15$ gr/cc), corresponds to $E < 0.5$ GPa,
244 considerably smaller than typical values of $E = 10\text{--}20$ GPa of the cortex. Focusing on high HU
245 values in the histograms, discrepancies between raw HU values obtained from the different QCT
246 scans are clearly visible, suggesting that the HU without calibration is scanner-dependent (Fig.
247 4A-C). Boundary correction and moving average algorithms only influenced the likelihood of HU
248 values already present in the bone (the y axis in the histograms). In contrast, HU values (the x axis)
249 are affected by the phantom calibration alone; therefore, phantom calibration reduces scanner
250 dependence, although some variation is still present (Fig. 4D-F).

251 The maximum RSD was <8% in principal strains along the bones in a femur and humerus, and
252 <11% in a proximal femur without the head. The mean RSD was about half of the maximum RSD,
253 $\leq 5\%$ in "regular" bones and <6.1% in the proximal femur. However, the total maximum difference
254 observed between two scanners in all three bones and all ROIs was 20%. **Linear correlations**
255 **between each pair of models showed very good agreement.**

256 **ICC values >0.9 indicate excellent reliability (Koo and Li, 2016) among the different scanners.**
257 **However, these findings should be interpreted with caution. The ICC highly depends on the**
258 **variance between the examined subjects, i.e. the ROI strains. These were taken from different**
259 **regions along different bones and thus characterized by high variability. Additionally, the ICC**
260 **provides only a general measure of the agreement level, with no quantitative estimate of the**
261 **variability between the scanners.**

262 The maximum tensile/compressive principal strains, the quantities of interest when risk of fracture
263 is considered, were obtained at similar locations in all models (Fig. 6). The high variability in the
264 maximum principal strains values obtained in the proximal femur (RSD <13.3%) can be related to
265 the highly complex geometry at the unusual location of the maximum strains (Fig. 6-C) and thus
266 may not be representative. Using a clinical "soft/body" filter/convolution kernel at 120 kV and
267 calibration with liquid K₂HPO₄ phantoms, the maximum RSD between QCTFE maximum
268 principal strains in "regular" bones (such as the femur and humerus), is <8%. Nonetheless, there
269 may be differences up to 20% between maximum principal strains estimated using two scanners
270 with the most extreme differences; thus, in a longitudinal study only changes larger than 20% in
271 principal strains may be significant. Such analyses are important, for example, to determine
272 prognosis in a patient with osteoporosis or metastases. It is noteworthy that previous experience
273 suggests that in pathological situations such as benign or metastatic tumors, changes in the strain

274 fields for the pathological femur are of tens of percent compared to the healthy femur, so the 20%
275 difference from different scanners would be of smaller significance in these cases (Sternheim et
276 al., 2018).

277 Repeated scans on same CT systems resulted in an average difference of 2.7% in the maximum
278 principal strains. These findings suggest that one should expect similar or higher variability
279 between CT scanners.

280 Exposure (mAs) has a negligible influence on differences in QCTFE results between CT systems,
281 as demonstrated in previous studies (Giambini et al., 2015; Nazarian et al., 2008). We conducted
282 two identical scans where a change in the exposure level was the only difference in protocol and
283 obtained almost no change in the HU histograms.

284 Several previous studies have emphasized the use of phantoms to quantify HU variability between
285 CT scanners. A study by (Suzuki et al., 1991) investigated the variance between QCT scans of
286 cadaveric vertebrae together with K_2HPO_4 phantoms obtained from 16 CT scanners, including 10
287 different models. They reported “good correlations ($r > 0.97$) ... observed for QCT values obtained
288 by various CT scanners.” An excellent review (Cann, 1988) showed good reproducibility of QCT
289 values obtained from several scanners, although CT systems used in the study are by now outdated.
290 These publications however did not evaluate differences in strains computed by QCTFE. To the
291 best of our knowledge, the only study addressing the latter issue examined the femoral fracture
292 load predicted by QCTFE analyses, as assessed with two different CT scanners (Carpenter et al.,
293 2014).

294 It is important to note that most publications using QCTFE were based on the classical h-version

295 of the FE method, which does not include numerical convergence studies. Numerical errors
296 inherent in h-version techniques may contribute substantially to the differences in predictions. As
297 an example, a difference in strains of $\sim 3\%$ was found at the neck between the FE analysis
298 performed at $p=4$ compared to the converged strains at $p=8$.

299

300 *Limitations and Future Studies*

301 This study did not consider body size effects that may be taken into consideration using
302 anthropomorphic standardization phantoms (ASPs) to assess inter-scanner differences. Carpenter,
303 et al. (Carpenter et al., 2014) found that differences in body size may result in varying estimates
304 of femur strength between scanners. In addition, tube voltage (kV) has an influence on HU
305 distribution histograms. We performed scans using 80–140 kV (data not presented here) and will
306 address the influence of kV on estimation of bone's mechanical response in a future publication.

307 We studied three manufacturers and four CT scanners at four different medical centers. While
308 these are believed to represent the majority of CT scanners in the Western world, future studies
309 may include a larger number of CT systems and manufacturers.

310 **The conclusions of our study are based only on descriptive statistics. Future studies should include**
311 **more subjects, i.e. more bones of the same kind.**

312 Finally, although past investigations suggest the use of phantoms in the range of 0–300 mg/cc
313 (Bessho et al., 2007; Miura et al., 2017; Yosibash et al., 2014), cortical tissue has an equivalent
314 mineral density of about 1200 mg/cc, about four times higher. The slope parameter m in Table 2
315 is sensitive to inaccuracies in the extreme concentration phantoms (0 and 300 mg/cc), thus it

316 strongly influences predicted strains. To improve the calibration, two alternatives will be
317 considered in a future investigation: Using phantom concentrations covering the entire range of
318 HUs, or preferably, developing phantom-less methods as suggested in (Lee et al., 2017; Suzuki
319 et al., 1991).

320

321 ***Conclusions***

322 QCTFE analyses of long bones performed by seven different CT scanners showed a maximum
323 20% differences in principal strains. In a longitudinal study, only changes larger than 20% in
324 principal strains may be significant.

325

326 **Conflicts of Interest**

327 Prof. Yosibash has a financial interest in PerSimiO, a technology company specializing in
328 personalized simulations in orthopedics. Prof. Sosna is a member of the PerSimiO Board of
329 Advisors. Neither Prof. Yosibash or Prof. Sosna derived any direct financial support, research
330 support, or other benefit from the company that could jeopardize their independence in this
331 research. Primary data was jointly controlled by Yekutiel Katz and Gal Dahan. They and the
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Table 1: CT scanners and hospitals in this study.

Scan Abbreviation	Medical Center	CT scanner model (# detector rows)
A	MC 1	Brilliance Philips Healthcare (64)
B	MC 2	
C	MC 4	
D	MC 1	iCT Philips Healthcare (256)
E	MC 2	
F	MC 3	Somatom Definition Edge Siemens Medical Systems (128)
G	MC 1	Optima CT660 GE HealthCare (128)

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456 **Table 2: Detailed information on the different QCT scans. All scans performed at 120 kV.**

Scan	Scanner (slices)	Date of Scanning	Exposure [mAs]*	Slice spacing & thickness (mm)**	Pixel size [mm]	$\rho_{K_2HPO_4} = m \times HU + n$	
						m	n
A	Philips Brilliance (64)	May 17, 2015	374	1	0.496	0.816	1.82
		Jan 18, 2017	83-283	1	0.601	0.814	-3.20
B	Philips Brilliance (64)	Dec 12, 2015	232	1	0.46	0.795	3.08
		Jan 18, 2017	223-370	1	0.523	0.818	-3.82
C	Philips Brilliance (64)	July 19, 2016	301	1	0.492	0.793	3.85
		Jan 18, 2017	164	1	0.518	0.826	-4.50
D	Philips iCT (256)	May 17, 2015	374	1	0.473	0.794	0.60
		Jan 18, 2017	81-106	1	0.557	0.811	-2.28
E	Philips iCT (256)	Dec 12, 2015	406	1	0.574	0.806	-2.96
		Jan 18, 2017	321-1104	1	0.502	0.813	-3.49
F	Siemens Somatom Definition (128)	July 21, 2015	210	1	0.440	0.869	0.00
		Jan 18, 2017	96-215	1	0.621	0.857	0.89
G	GE Optima CT660 (128)	May 17, 2015	500	1.25	0.523	0.823	1.00
		Jan 18, 2017	3-6	1.25	0.594	0.831	-6.83

457 * A range of mAs values indicates scans with adaptive current algorithm.

458 ** The slice thickness and slice spacing are equal for each scan.

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461 **Table 3: Mean RSD% and maximum RSD% for each of the three bones with homogeneous**
462 **constant Young modulus.**

	Femur	Humerus	Femur w/o head
Mean RSD%	2.08%	3.08%	2.51%
Max RSD%	3.45%	5.53%	4.90%
Location of Max RSD	N2	S6	N3

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465 **Table 4: Mean and maximum distances [mm] and standard deviation of the location of the**
 466 **maximum principal strains (QCTFE based on the 7 scanners).**

	Femur		Humerus	Proximal Femur	
	ϵ_1	ϵ_3	ϵ_3	ϵ_1	ϵ_3
Mean distance [mm]	0.48	0.91	0.66	0.73	1.00
SD [mm]	0.32	0.40	0.23	0.23	0.52
Max distance [mm]	1.10	1.57	1.05	1.01	1.97

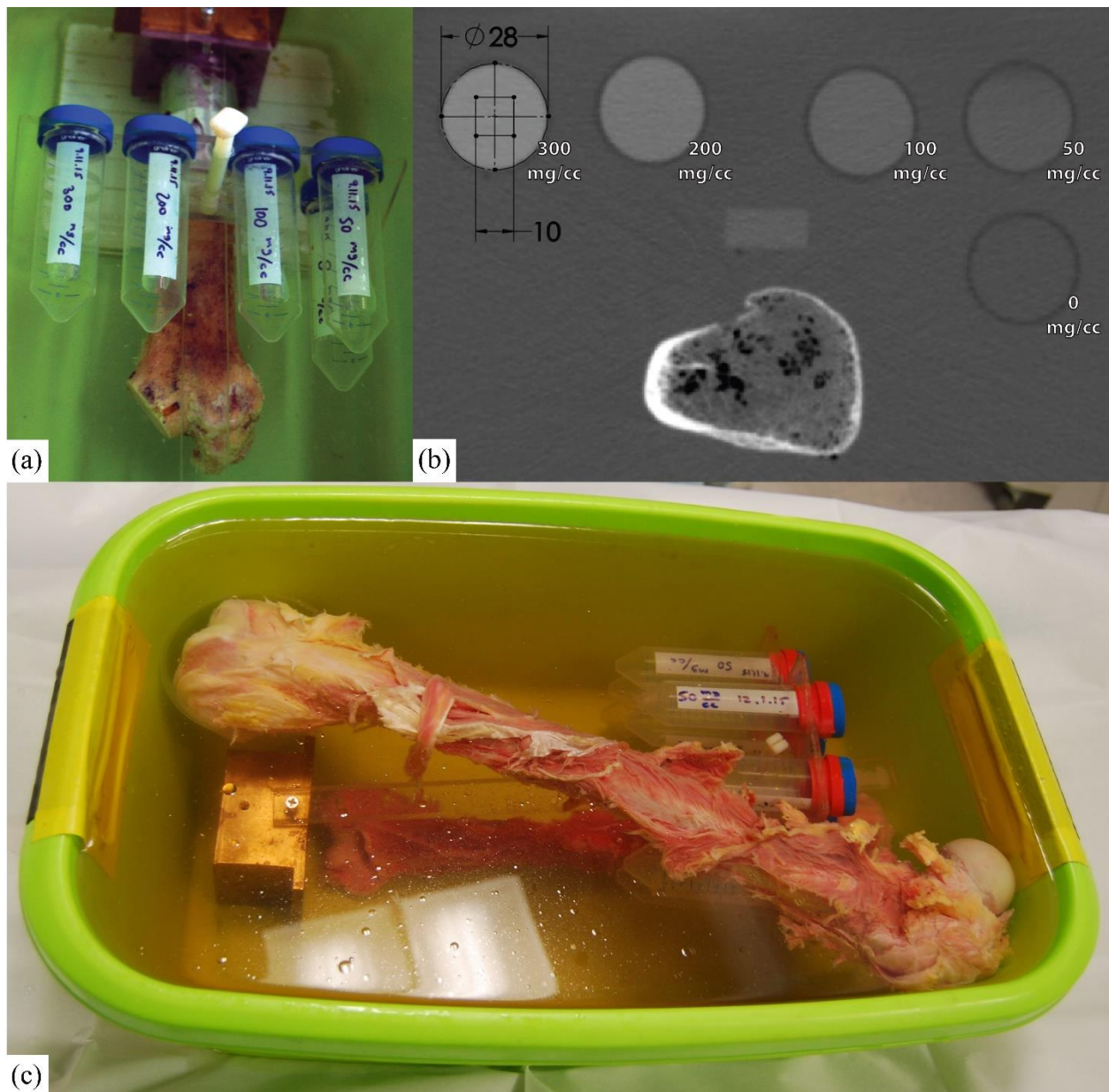
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Figures

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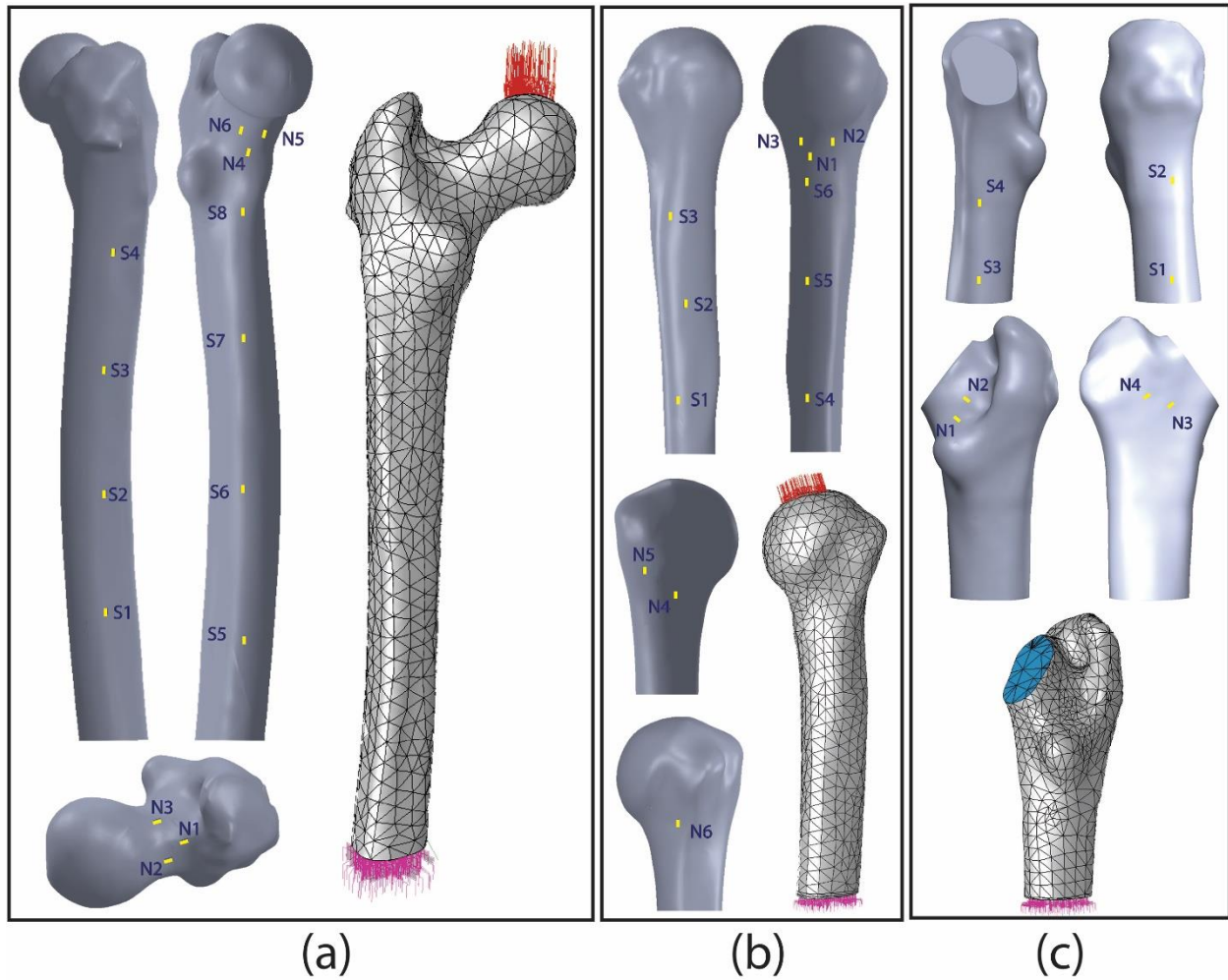
473 **Figure 1** (a) Proximal femur in the water bath surrounded by five liquid potassium phosphate (K_2HPO_4)

474 phantoms. (b) A representative CT slice of the femur and phantoms. (c) Whole femur and humerus in

475 water bath with the liquid phantoms.

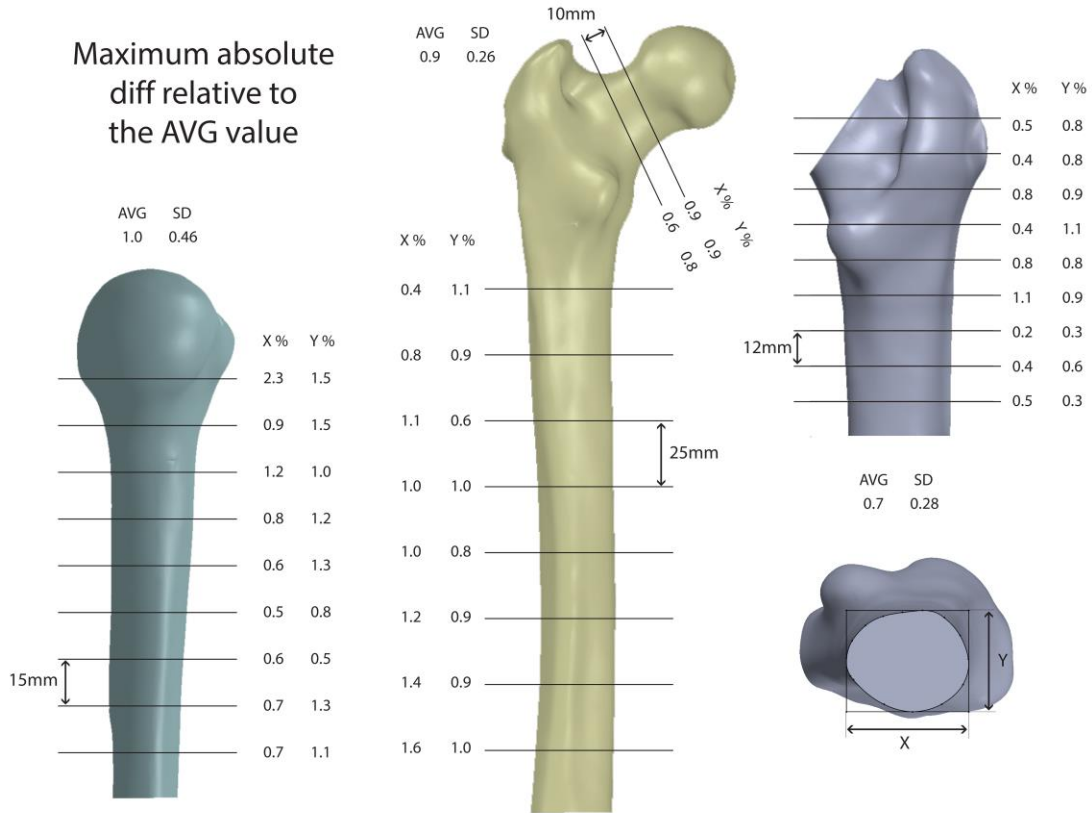
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Figure 2 (a) The femur and regions of interest (ROI) at which strains were extracted with the QCT-based finite element analysis (CTFEA) model and boundary conditions. (b) The humerus and ROI with CTFEA model and boundary conditions. (c) The proximal femur with ROI and CTFEA model with displacement boundary conditions.

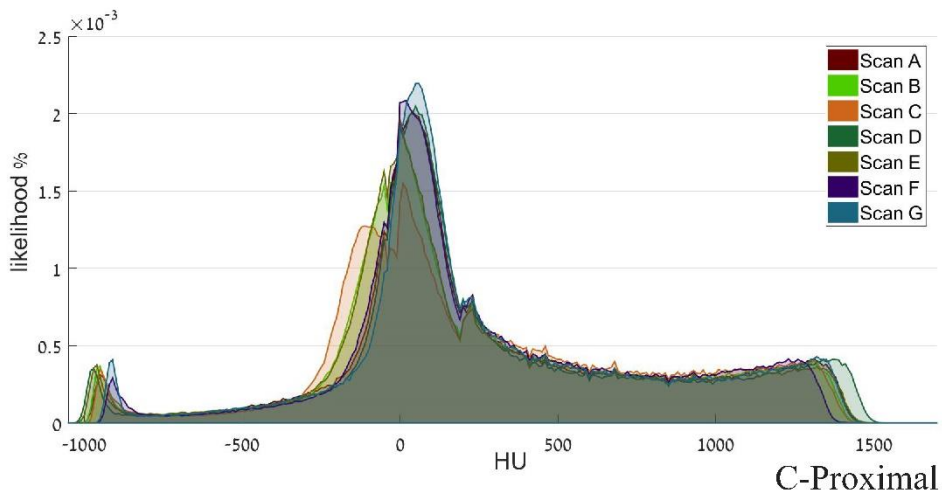
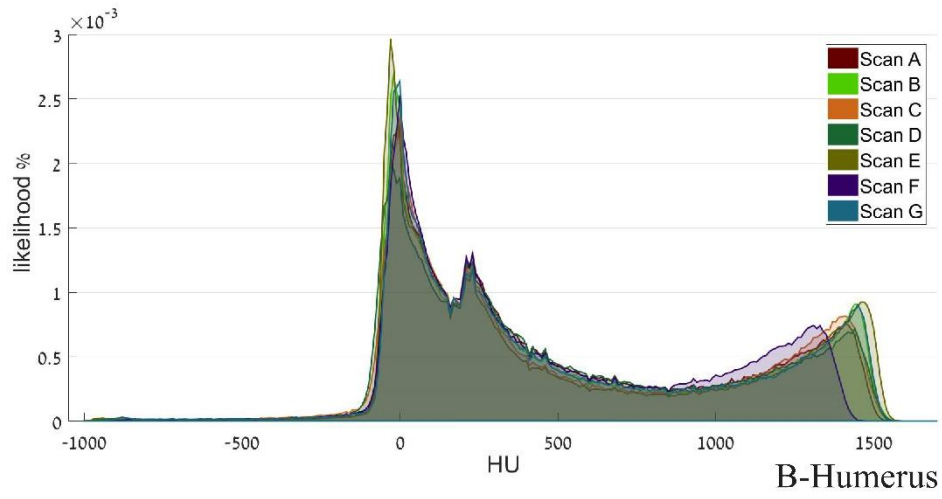
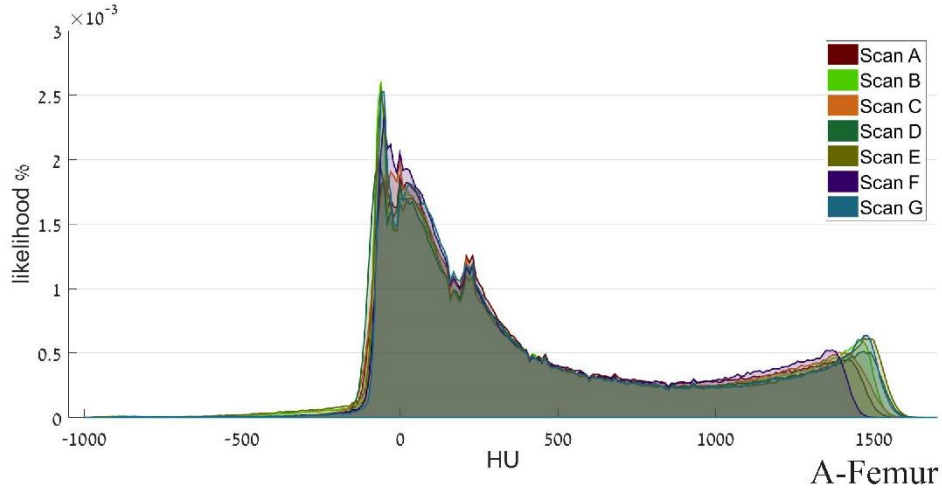


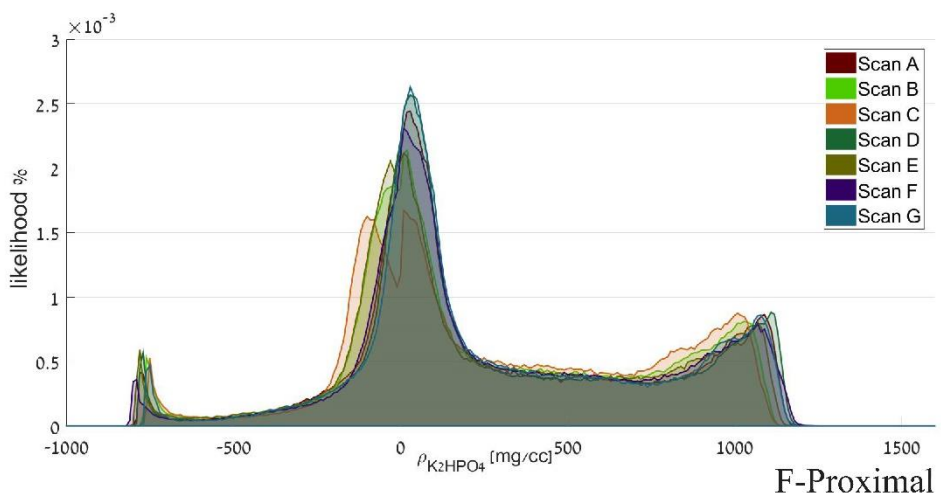
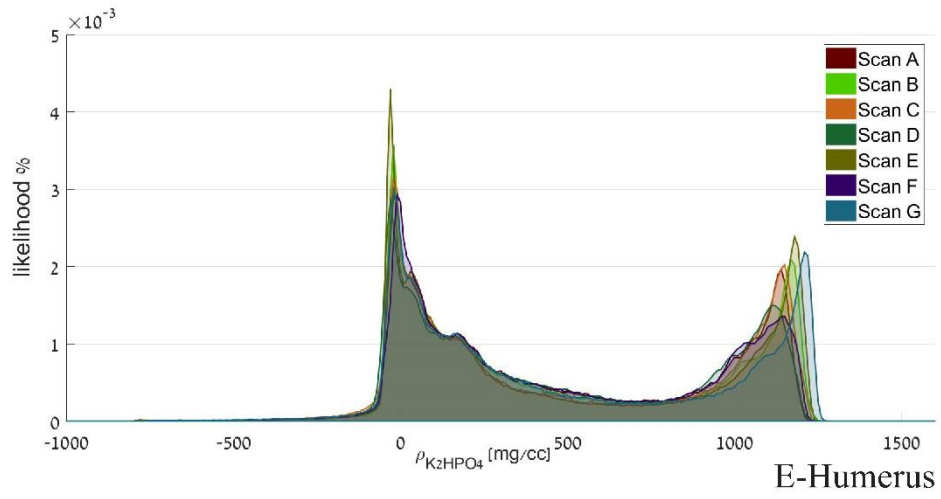
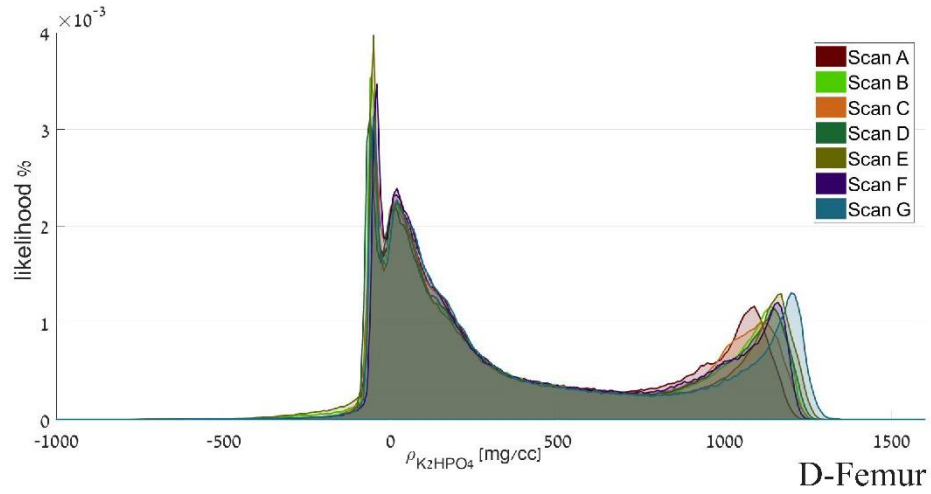
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487 **Figure 3** Maximum difference (relative to scan's average) between all scans at several slice locations
 488 along the segmented femurs and humerus. (AVG, average; SD, standard deviation; X, x-axis; Y, y-axis)

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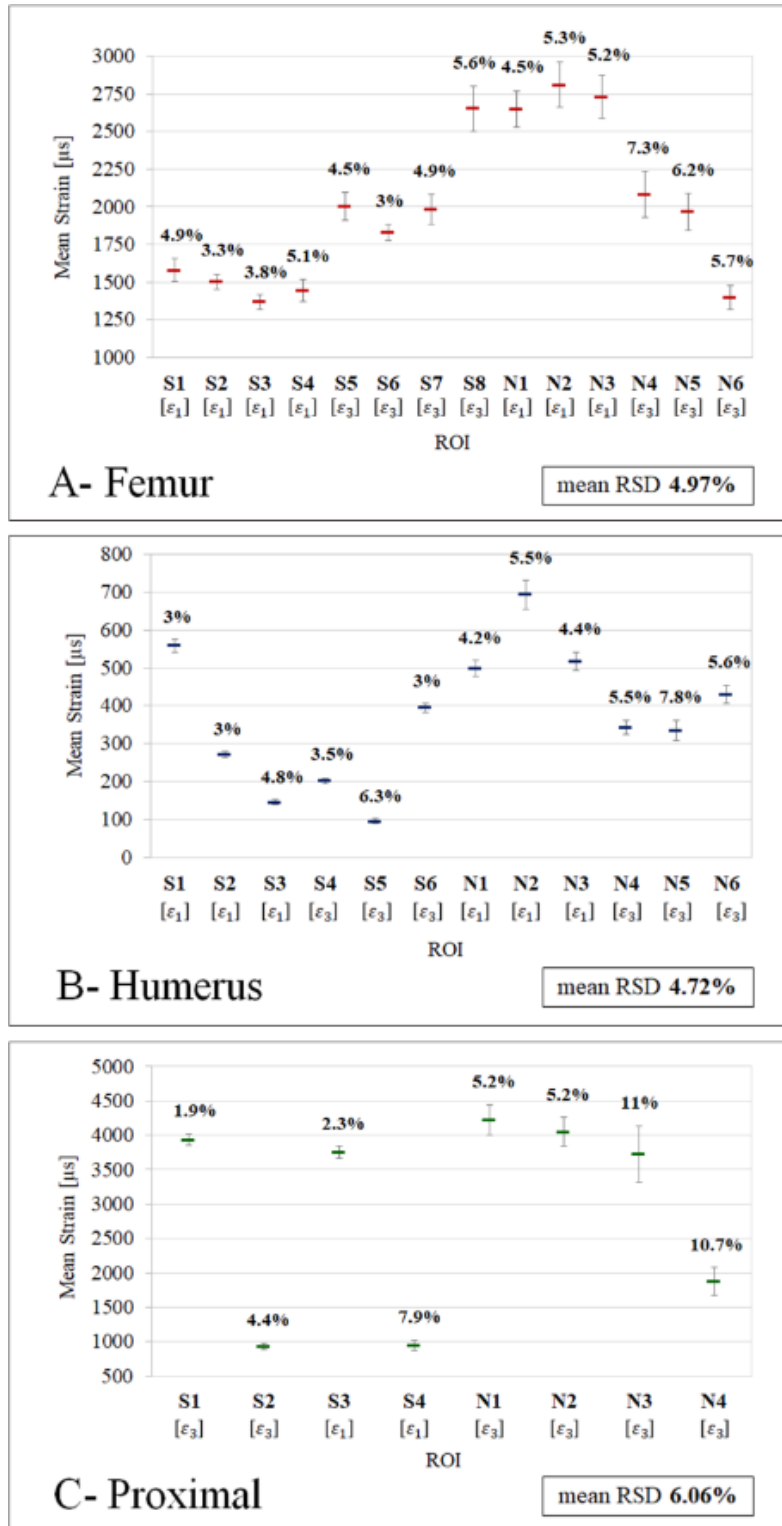
493 **Figure 4** Normalized histograms of Hounsfield unit (HU) values inside the femur, humerus, and proximal
494 femur in scans A through G. The area under the histogram equals 1. (A–C) Histogram of raw HU values.
495 (D–F) Histogram of ρ_{ash} after phantom calibration, moving average, and boundary corrections.

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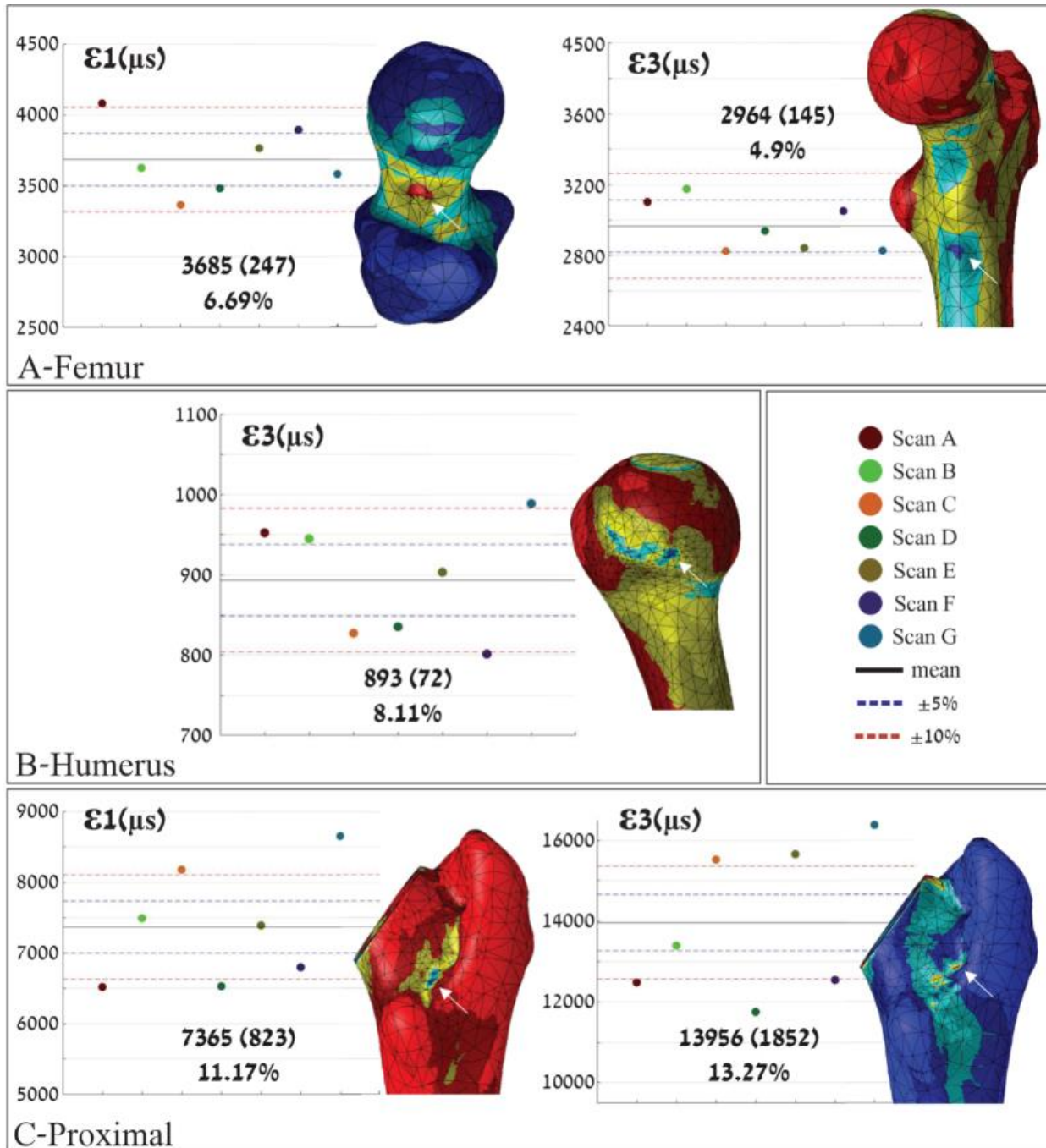
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541 **Figure 5** Mean and RSD% values of the maximum absolute principal strains at each ROI across the
 542 different scans for the femur (A), humerus (B) and proximal femur (C)



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544 **Figure 6** Maximum tensile/compressive (ϵ_1/ ϵ_3) principal strains, in the femur (A), humerus (B), and
 545 proximal femur (C). In each panel the mean (SD) and RSD% of the maximal strain is presented. The
 546 humerus is under compressive state; thus, no tensile graph is presented.

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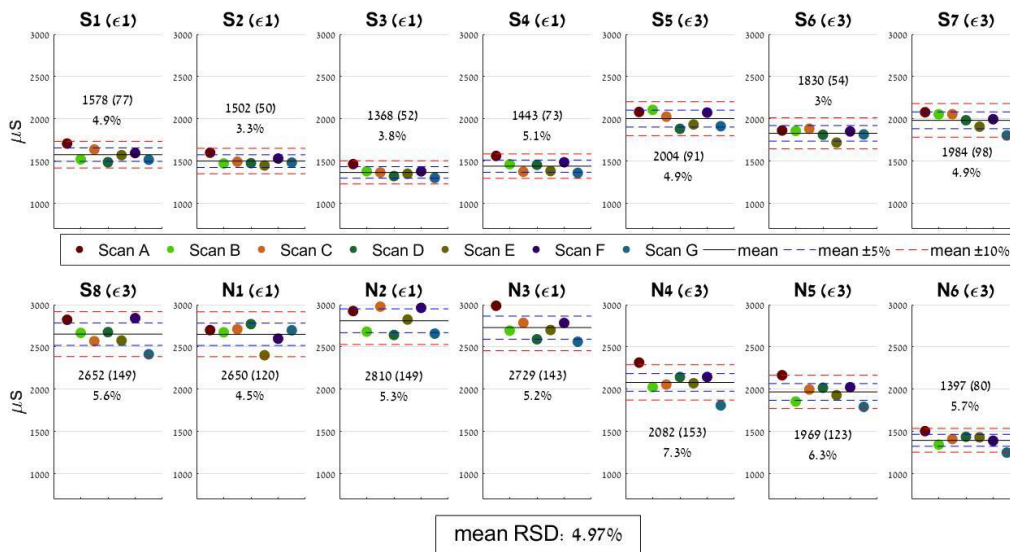
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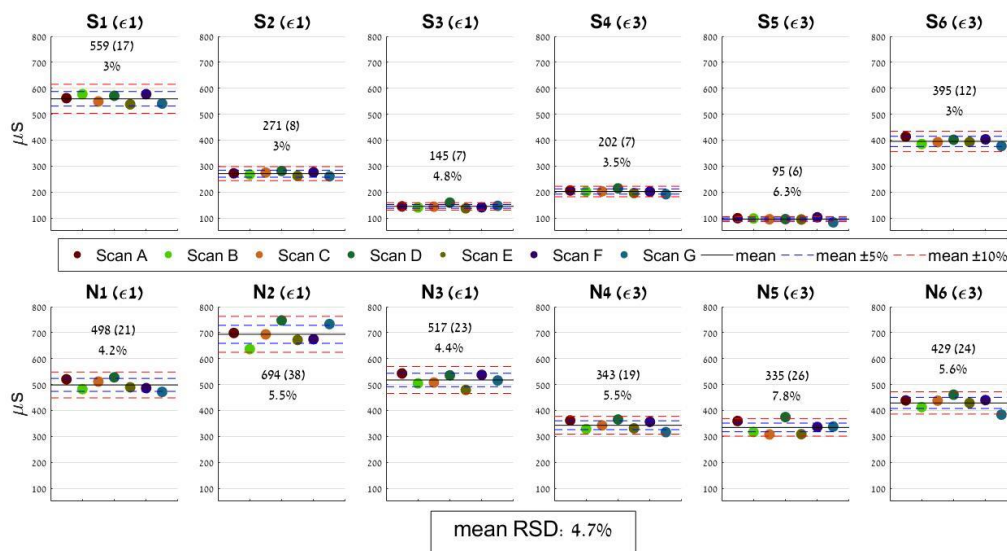
Supplementary Material

Appendix A – Full results of the maximum absolute principal strains at ROIs in the different FE models generated by the data in the different scans for the femur (A.1), humerus (A.2) and proximal femur (A.3). In each panel, the mean (SD) and RSD% for the ROI strain is presented, including the mean RSD% for the entire bone.



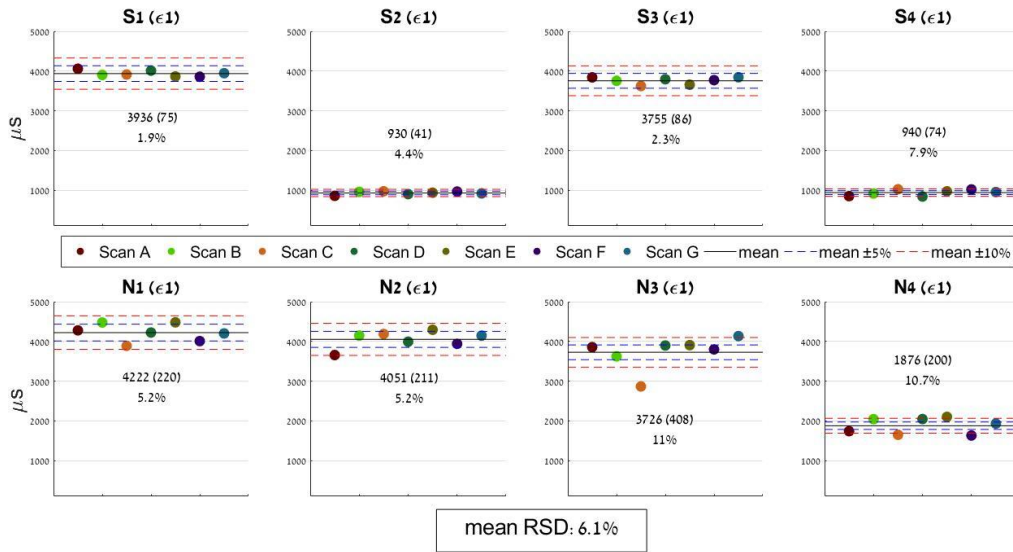
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Figure A.1 - Maximum absolute principal strains at ROIs in the different scans for the femur.



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Figure A.2 - Maximum absolute principal strains at ROIs in the different scans for the humerus.

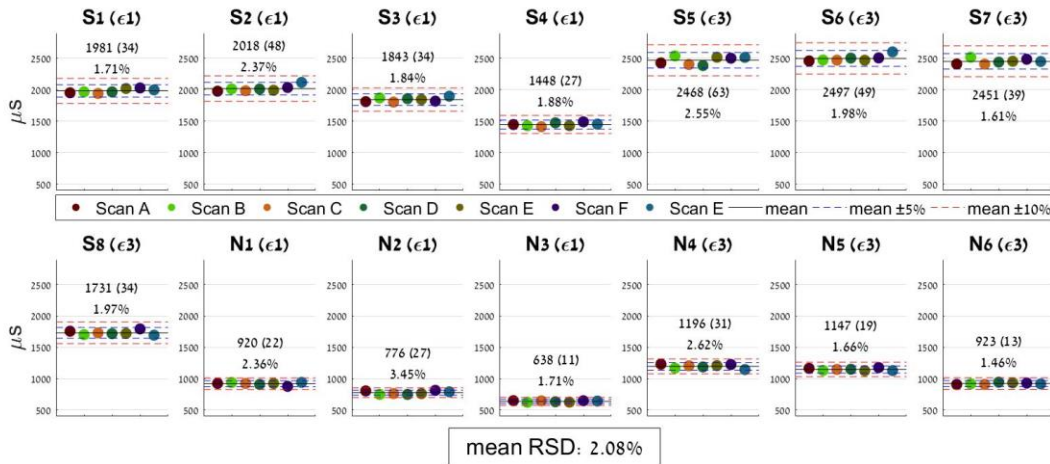


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Figure A.3 - Maximum absolute principal strains at ROIs in the different scans for the proximal femur.

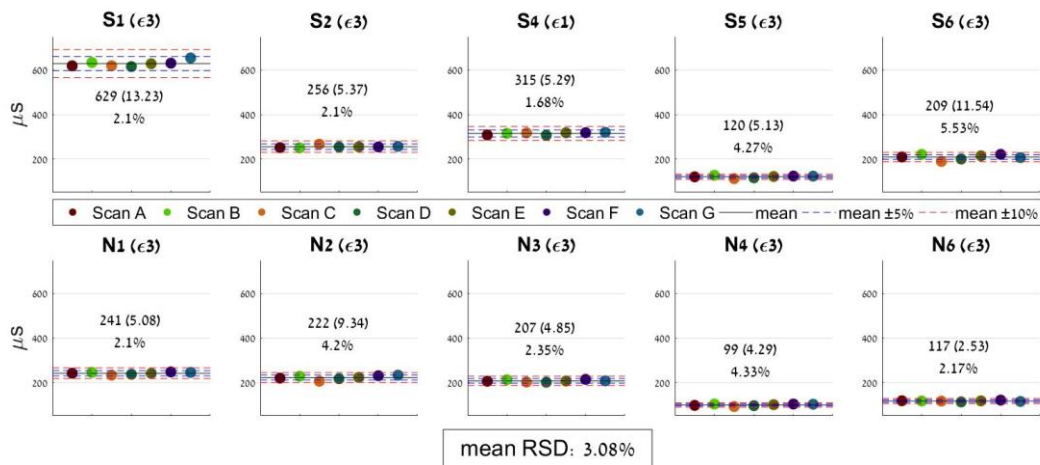
566 **Appendix B - Maximum absolute principal strains at ROIs in the different FE models generated by**
 567 **the data in the different scans for the constant material properties $E = 10$ GPa. This is to investigate**
 568 **the influence of the geometry difference on the principal strains. In each panel, the mean (SD) and**
 569 **RSD% for the ROI strain is presented, including the mean RSD% for the entire bone.**

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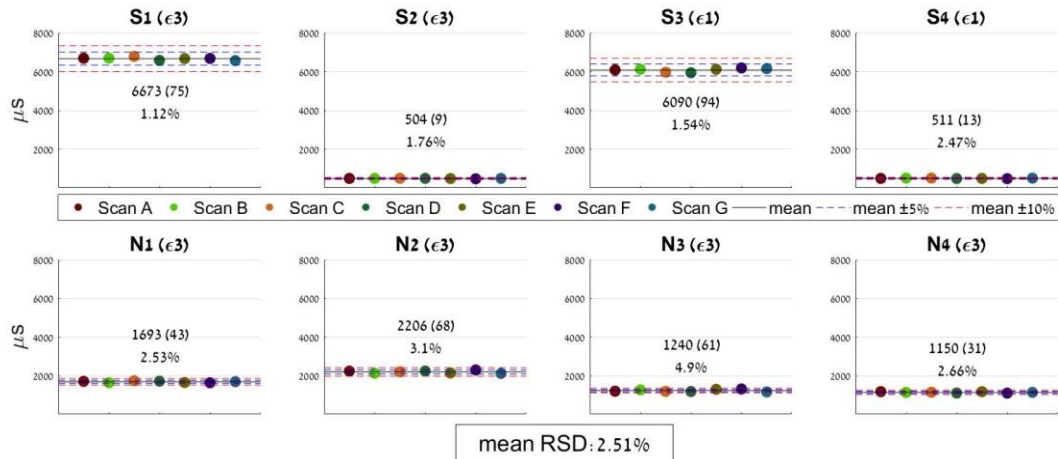
Figure B.1 Maximum absolute principal strains at ROIs in the different scans for the homogeneous femur with constant material properties $E = 10$ GPa.



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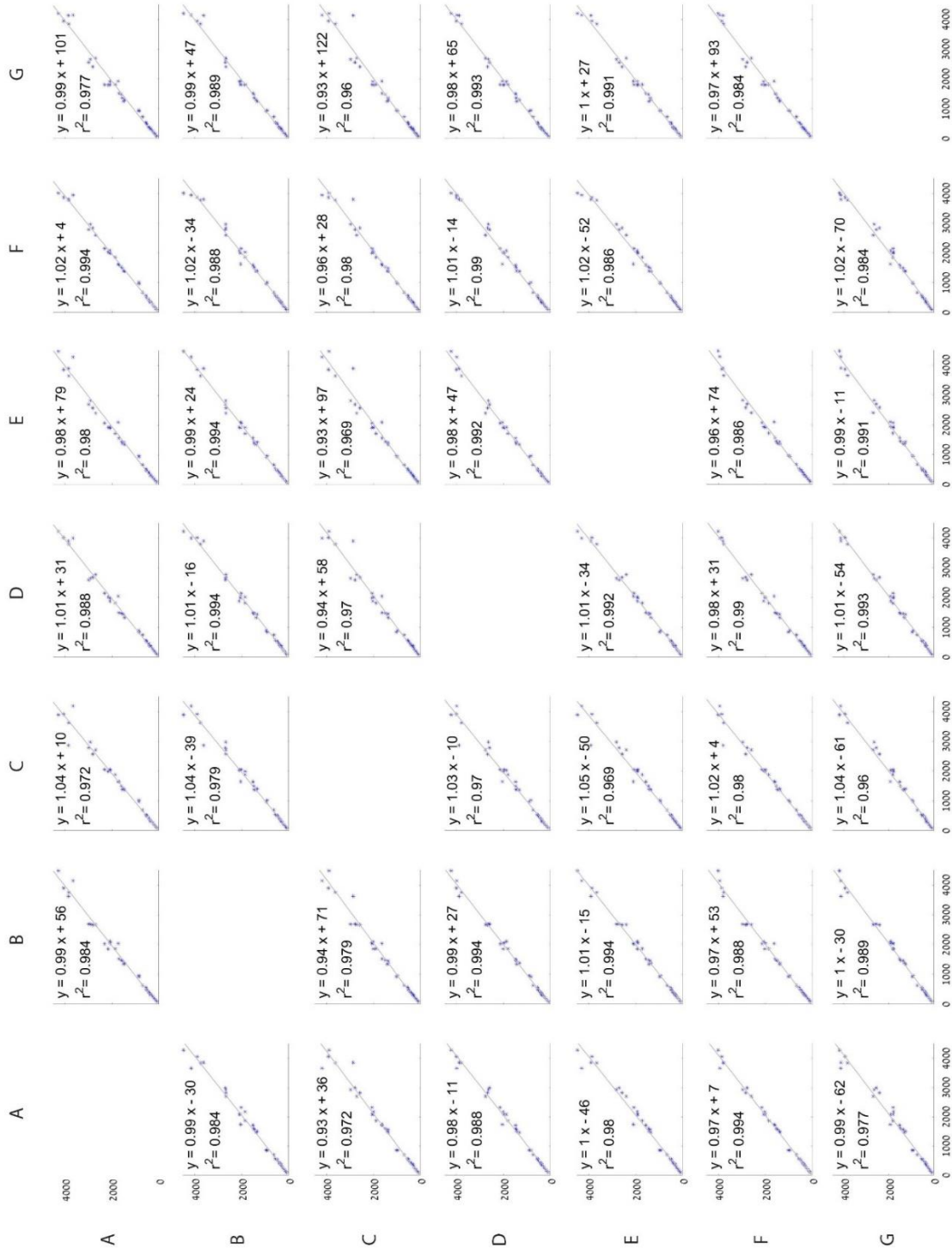
580 **Figure B.2** Maximum absolute principal strains at ROIs in the scans for the homogeneous humerus with
 581 constant material properties $E = 10$ GPa. S3 and N5 are not shown since the strains there are close to zero.
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 586 **Figure B.3** Maximum absolute principal strains at ROIs in the scans for the homogeneous proximal femur
 587 with constant material properties $E = 10$ GPa.
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 592 **Figure C** Linear correlation matrix showing agreement in strains between all pairs of QCTFE models
 593 obtained from the different CT scanners.
 594