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Regional maps of rib cortical bone thickness and cross-sectional geometry

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Abstract

Here we present detailed regional bone thickness and cross-sectional measurements from full adult ribs using high resolution CT scans processed with a cortical bone mapping technique. Sixth ribs from 33 subjects ranging from 24 to 99 years of age were used to produce average cortical bone thickness maps and to provide average \pm 1SD corridors for expected cross-section properties (cross-sectional areas and inertial moments) as a function of rib length.

Results obtained from CT data were validated at specific rib locations using direct measurements from cut sections. Individual thickness measurements from CT had accuracy (mean error) and precision (SD error) of -0.013 ± 0.167 mm (R^2 coefficient of determination of 0.84). CT-based measurement errors for rib cross-sectional geometry were -0.1 ± 13.1 % (cortical bone cross-sectional area) and 4.7 ± 1.8 % (total cross-sectional area).

Rib cortical bone thickness maps show the expected regional variation across a typical rib's surface. The local mid-rib maxima in cortical thickness

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along the pleural rib aspect ranged from 0.9 mm to 2.6 mm across the study population with an average map maximum of 1.4 mm. Along the cutaneous aspect, rib cortical bone thickness ranged from 0.7 mm to 1.9 mm with average map thickness of 0.9 mm. Average cross-sectional properties show a steady reduction in total cortical bone area from 10% along the rib's length through to the sternal end, whereas overall cross-sectional area remains relatively constant along the majority of the rib's length before climbing steeply towards the sternal end. On average, male ribs contained more cortical bone within a given cross-section than was seen for female ribs. Importantly, however, this difference was driven by male ribs having larger overall cross-sectional areas, rather than by sex differences in the bone thickness observed at specific local cortex sites.

The cortical bone thickness results here can be used directly to improve the accuracy of current human body and rib models. Furthermore, the measurement corridors obtained from adult subjects across a wide age range can be used to validate future measurements from more widely available image sources such as clinical CT where gold standard reference measures (e.g., like direct measurements obtained from cut sections) are otherwise unobtainable.

Keywords: Cortical bone, Rib, Computed Tomography, Cortical thickness, Cross-sectional geometry, Computational models

1. Introduction

2 Ribs provide crucial protection for the thoracic viscera, but are often
fractured in a variety of scenarios (Wuermsler et al., 2011). In motor vehicle
4 crashes (MVCs) specifically, rib fractures continue to be prevalent despite

advances to safety systems and vehicles. Furthermore, the presence of rib
6 fractures increases mortality and morbidity rates in vulnerable populations
(e.g., elderly) (Stawicki et al., 2004; Sirmali et al., 2003). Recent research has
8 highlighted the need for in-depth exploration of rib geometry to better under-
stand whole thoracic response to loading (Murach et al., 2018), and therefore
10 develop injury mitigation techniques. Broadly, this can be accomplished us-
ing computational human body models (HBMs), an important modern tool
12 for injury assessment.

These models rely on accurate input for their prediction of a rib's response
14 to loading. Global and cross-sectional geometry have been identified as im-
portant predictors of this response in ribs (Stein, 1976; Agnew et al., 2013;
16 Murach et al., 2017; Agnew et al., 2018; Holcombe et al., 2016). Rib cor-
tical bone thickness spans approximately 0.1 mm to 2.4 mm (Choi & Kwak,
18 2011; Mohr et al., 2007; Agnew et al., 2018) and is commonly represented in
finite element (FE) computational models using shell elements surrounding
20 a solid trabecular core. Li et al. (2010) found that models which incorpo-
rate variable thickness into their cortical bone definitions can better predict
22 a rib's structural response. However, the precise distribution of thicknesses
along and around rib bones is not well understood. Current sources report
24 thickness values only in aggregation across particular zones around the rib's
circumference or along its length (Agnew et al., 2018; Mohr et al., 2007;
26 Mayeur et al., 2010). In most current models the rib global geometry is
drawn from a single individual, and cortical bone thickness values are drawn
28 from these limited or simplified literature sources (Gayzik et al., 2011; Choi
et al., 2009; Kemper et al., 2007). Furthermore, there is now evidence that

30 cross-sectional bone area and bone distribution may contribute more than
only cortical thickness in predicting rib structural properties (Agnew et al.,
32 2018), highlighting the need to quantify all rib cross-sectional geometry more
thoroughly across the population. Despite the current knowledge that pre-
34 cise rib cross-geometry is crucial for predicting rib fracture properties, the
incorporation of such geometry at the level of detail necessary to reflect true
36 human variation has not been fully realized in current HBMs. When rib
modifications are made to HBMs to simulate population-based differences, a
38 greater emphasis is generally placed on altering material properties and gross
thoracic geometry than cross-sectional rib geometry to achieve the desired
40 structural results (Ito et al., 2009; Schoell et al., 2015).

The Cortical Bone Mapping (CBM) methodology allows for accurate mea-
42 surements of these important geometric factors from CT imaging (Holcombe
et al., 2018; Treece & Gee, 2015; Treece et al., 2010). CBM has been pre-
44 viously applied to ribs for tracking bone thickness reductions after cancer
therapy (Okoukoni et al., 2016), but it has not yet been used to report rib
46 cortical bone thickness distributions from individuals or across populations.

In this study we apply the CBM method to high resolution CT scans of full
48 ribs. We assess the accuracy of this method against cross-sectional histology
images taken at key locations along each rib, and develop full cortical bone
50 thickness maps along and around individual ribs. We spatially register these
maps from multiple individuals to present a detailed average thickness map
52 that is representative of an American adult population. The methodology
presented here can be used to build individualized rib models, while the
54 aggregated maps can be applied to enhance general population models.

2. Materials and methods

56 This study utilizes histology images extracted from, and CT image data
covering, 33 complete sixth-level ribs ethically obtained from anatomical
58 donors in Ohio, USA (16 male, 17 female) with no existing trauma or gross
pathological condition affecting the ribs. Subject ages ranged from 24 to
60 99 years (average \pm SD 65 ± 21) with distributions shown by sex in Fig-
ure 1. Male subjects (70 ± 18 years) were on average older than female
62 subjects (60 ± 22 years), however this difference was not statistically signifi-
cant ($p = 0.15$).

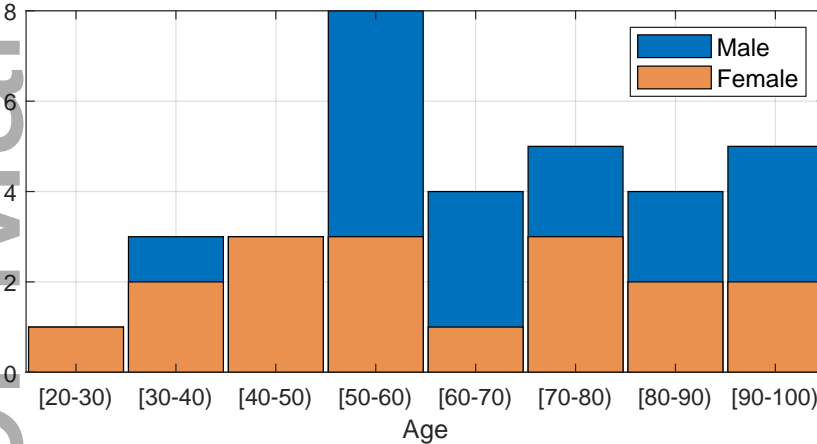


Figure 1: Stacked subject age counts by sex and decade of life.

64 Complete ribs were excised from subjects soon after death, and subse-
quently CT scans of each rib wrapped in saline-soaked gauze were taken
66 using a Phillips Vereos digital PET/CT with 64 slice Ingenuity technology
at an axial resolution of 0.15 mm/pixel with slice spacing of 0.67 mm/pixel
68 (i.e., $0.15 \times 0.15 \times 0.67$ mm voxels). Ribs were oriented with their end-to-end
axis aligned vertically in the scan such that mid-rib regions were approxi-

70 mately co-planar with the scan’s axial plane. After experimental bending
tests, cross-sections perpendicular to the long axis of the rib were taken
72 immediately adjacent to each fracture site (39 total sites at either one or
two fracture sites per rib) while ensuring no disruption to the bone cortex.
74 Approximate fracture site locations—measured manually using string—were
noted as a percentage of rib curvilinear length. Slides were then prepared
76 according to undecalcified hard tissue histology standards (see Agnew et al.
(2018)). High-resolution microscopy (Olympus BX61VS) allowed for direct
78 image capture (*i.e.*, no reconstruction) of the entire rib section at 100x total
magnification and a resolution of 0.69 microns/pixel.

80 *2.1. Histology image processing*

Periosteal and endosteal cortical borders were semi-manually identified on
82 each histology image using ImageJ software (NIH) by an experienced bone
histologist (AA) (Dominguez & Agnew, 2019). These were used as gold stan-
84 dard cortical bone cross-sectional geometries at their specific rib locations,
and each histology image was spatially registered within its corresponding
86 CT image volume as follows. Firstly, rigid registration errors were calcu-
lated between the histology-derived periosteal border and those taken from
88 successive cross-sectional cuts through an initial CT-derived periosteal sur-
face (described below). Local minima in registration error indicated strongly
90 matching regions, and visual overlays of the histology image onto the CT vol-
ume were used for minor adjustment to align cortices and trabeculae between
92 the two modalities. A typical overlay is presented in Figure 2 showing the
local patterns of trabecular bone—clearly visible on histology and slightly
94 blurred on CT—in strong alignment. For all cut histology sections, the loca-

tion in CT image space (along the rib's length) of minimum rigid registration
96 error occurred near to the corresponding physical fracture location that was
noted by hand. In all cases the final chosen position that showed the strongest
98 coherence in trabecular patterns between histology and CT was less than 1 %
in rib length from the position exhibiting minimum rigid registration error.

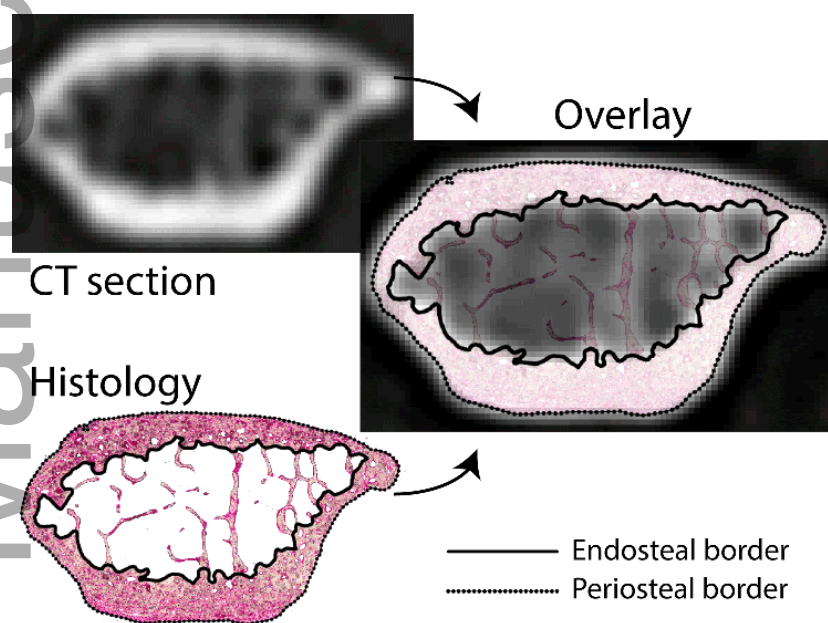


Figure 2: Exemplar histology image overlay showing the spatial correspondence to the underlying CT image volume at each chosen cross-sectional position.

100 *2.2. CT image processing*

An initial and approximate 3D periosteal surface was generated from each
102 rib's CT volume via segmentation performed in MIMICS (v19, Materialise).
All subsequent image and statistical analyses were performed in MATLAB
104 (The Mathworks). A central axis along the rib was formed from the rib head
(at the vertebral or posterior end) to its sternal (or anterior) end by fitting

106 a smoothed spline through the 2D centroids obtained from successive cross-
sectional cuts across this initial periosteal surface. This initial surface was
108 then discretized along the central axis into 301 successive and equally spaced
cross-sections. Each section was further discretized to 80 locations around
110 its circumference to produce an overall surface map as depicted in Figure 3.

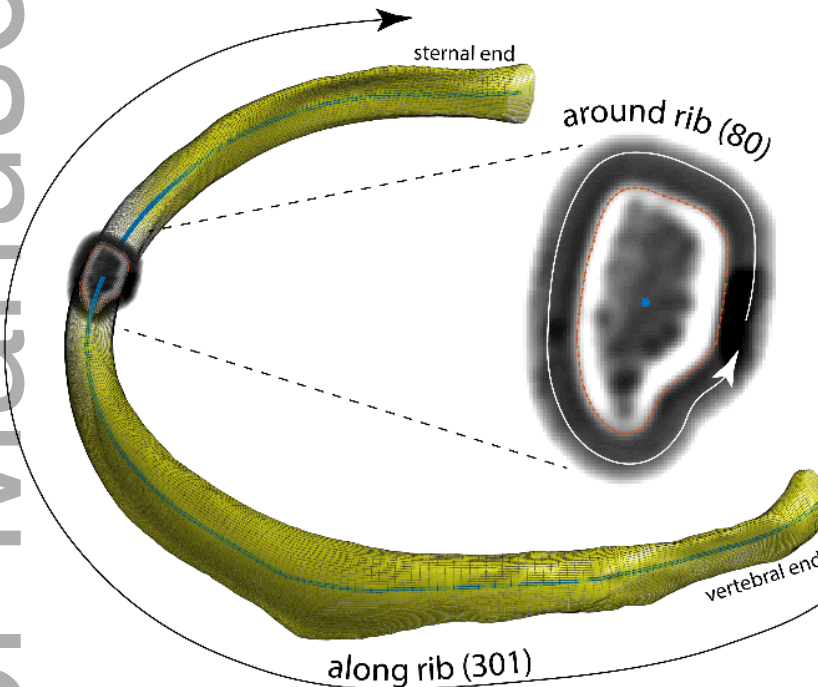


Figure 3: Ribs are discretized to a 301 (along) by 80 (around) grid of rib surface locations.

Local pleural and cutaneous aspects for each cross-section were calculated
112 using the two points on its circumference intersected by that section's minor
inertial axis. A smoothed spline fitted to these points formed overall rib
114 pleural and cutaneous aspects. Co-alignment between rib surface maps was
achieved by aligning the circumferential locations along these aspects.

116 At each surface location (that is, at 301×80 locations per rib), a 1D

cortical signal of the image intensity across the cortex was produced by re-
118 sampling the underlying CT image in a direction normal to the 3D surface at
that location. The Cortical Bone Mapping method (CBM) was then applied
120 to each cortical signal. CBM uses nonlinear optimization to match a model
consisting of three constant density regions (y_0 outside the periosteal border,
122 y_1 within the cortex, and y_2 inside the endosteal border) to the cortical
signal. The distance between the optimally-fitted periosteal and endosteal
124 estimates (x_0 and x_1) provides the local cortical bone thickness, and the
collection of individual estimates across a rib's gridded surface provide overall
126 rib thickness maps. As per previous work (Holcombe et al., 2018; Treece
et al., 2010) the cortical density model parameter (y_1) was fixed within each
128 CBM optimization to a Hounsfield unit value corresponding to the density
seen in the thickest bone region along the rib, and weighting was applied
130 to more aggressively penalize model fitting errors located near the initial
periosteal surface.

132 In this study, additional error-based local smoothing was applied to the
collected x_0 and x_1 value maps using a 0.3 mm Gaussian-shaped smoothing
134 kernel that was further scaled by the inverse of the CBM model fitting error
at those same locations. This step served to reduce high-frequency noise
136 in resulting border locations across the rib surfaces and also reduced the
influence of poorly-fitted individual cortical signals.

138 Finally, filters were used to suppress potentially misleading thickness mea-
surements within a given cross-section as described in Holcombe et al. (2018)
140 and summarized below. Firstly, individual signals wherein the parameters
from the CBM method's optimization step did not converge to internal (non-

142 boundary) values were ignored. Secondly, morphological criteria applied to
each circumferential ring of estimated endosteal borders were used to dis-
144 card signals which did not pass through a single isolated cortical wall. This
is most commonly seen near areas of high local curvature such as the costal
146 groove.

2.2.1. Geometric measurements

148 Overall, the steps above served to produce an underlying 301×80 map of
local cortical bone thickness (CT.TH) estimates obtained from CT, with one
150 map for each of the 33 whole ribs. These maps were averaged to produce an
average CT.TH map for the study population.

152 Additionally, the sequence of periosteal and endosteal border positions
around each individual cross-section were joined to produce the geometric
154 shape of that section's predicted cortical shell (for 301 shells per rib). Cross-
sectional geometry measurements of each of these shells were calculated, con-
sisting of the total sub-periosteal area (TT.AR), the cortical area (CT.AR),
156 the endosteal area (ES.AR), and the cortical shell's maximal (or principal)
and minimal (or secondary) area inertial moments (IMAX, and IMIN). The
158 IMAX and IMIN inertial axes intersect the 2D centroid of the cortical shell
and, with rib cross-sections generally elongated, the IMAX inertial axis oc-
curs along an approximately inferior to superior aspect while the IMIN inertial
160 axis lies perpendicular to IMAX along a pleural to cutaneous aspect. As a
descriptor of rib cross-sectional aspect ratio, IRAT was calculated as IMAX
162 divided by IMIN. The six overall cross-sectional measurements were grouped
by position along a rib to report population average values and a $\pm 1SD$ pop-
166 ulation corridor, each calculated as a function of position from the vertebral

rib end to the sternal rib end.

168 Regional sex-based differences in bone thickness and in cross-sectional
geometry measurement distributions were assessed via two-sample t -tests
170 with significance determined at the $p < 0.05$ level.

2.3. Validation against histology

172 Each histological image ($N = 39$) matched a specific cross-sectional po-
sition along the length of one of the 33 whole ribs. A CT validation set was
174 produced by using gold standard measurements obtained directly from the
periosteal and endosteal borders drawn on the histology images, and pairing
176 them with measurements from the spatially equivalent locations within the
full CT image volumes of the same ribs. Therefore, the validation set for all
178 CT measurements consisted of 3120 local cortical bone thickness (CT.TH)
measurement pairs (at 80 locations around each of 39 sections), and 39 pairs
180 (one per histology image section) for the TT.AR, CT.AR, ES.AR, IMAX,
IMIN, and IRAT rib cross-sectional shape measurements. Additionally, the
182 difference in principal inertial axis orientation (IANG) between the cortical
shell shapes obtained using CT and histology was calculated. For IANG, a
184 positive difference indicated rotational misalignment (having the superior rib
aspect rotate towards the pleural side) of the cortical shell obtained using
186 CT compared to the target cortical shell from histology.

Individual thickness values discarded via the morphological filters de-
188 scribed above were excluded from the CT.TH validation set, and their border
positions were linearly interpolated via neighboring successful measurements
190 in order to provide complete cross-sectional geometry. For all measurement

192 pairs in the validation set, the measurement accuracy (mean error) and precision (SD error) of CT-based predictions was calculated.

3. Results

194 3.1. Accuracy assessment

From 3120 histology-matched cortical bone signal locations (at 80 locations sampled around each of the 39 sections with histology), 37 signals did not converge adequately during CBM optimization and 161 were identified by morphological filters as not falling across a clear singular cortex. The remaining 2922 (94%) predictions of CT.TH from CT for comparison to gold standard values from their histology-based pairs, are shown as scattered data in Figure 4.

202 Overall accuracy (mean error) and precision (SD error) of predicted cortical thickness values from CT ($N = 2922$) was -0.013 ± 0.167 mm, and Table 1 also lists the accuracy and precision for CT-based predictions of each of the full cross-sectional property measurements ($N = 39$). In general, cross-sectional property predictions were well correlated with gold standard values ($R^2 > 0.91$ for all properties), but with CT-based predictions on average overestimating area properties by 0.1% (CT.AR), 4.7% (TT.AR), and 6.8% (ES.AR).

210 3.2. Sectional property variation

Average values and 1SD male and female corridors for regional CT.AR, TT.AR, ES.AR, IMAX, IMIN, and IRAT are shown in Figure 5, and all corridor data is included as supplementary information. Specifically, male ribs

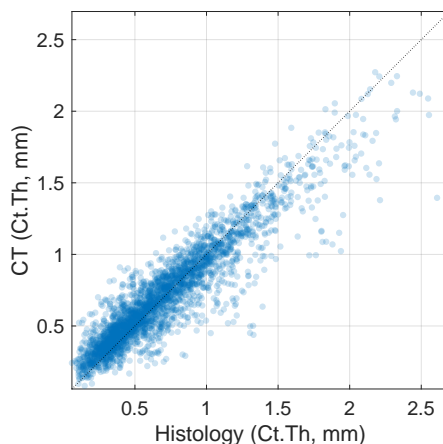


Figure 4: Predicted individual thickness measurements compared to spatially equivalent measures from histology (coefficient of determination $R^2 = 0.84$, $p < 0.0001$).

214 had significantly larger (at the $p < 0.05$ level) CT.AR at 56% of rib loca-
 tions, and significantly larger TT.AR, ES.AR, ES.AR, IMAX, and IMIN at
 216 over 96% of rib locations. Inertial aspect ratio (IRAT) varied along the rib
 with local peaks in aspect ratio occurring near either rib end and within a
 218 region spanning approximately 25% to 50% of the rib's length. This re-
 gion corresponds to the greatest prominence of the rib's costal groove, and
 220 here female ribs were significantly more elongated than male ribs whereas
 rib aspect ratios in other regions were not significantly different. As seen on
 222 Figure 5, the maximal CT.AR along the length of the rib occurs near the
 rib tubercle (approximately 10% rib length) and decreases steadily towards
 224 the sternal end, whereas TT.AR remains relatively constant across most of
 a rib's length before increasing sharply towards its sternal end. These cross-
 226 sectional changes along a rib's length are visualized in Figure 6 which shows
 the CT predictions of cortical bone borders for a number of subjects and rib

Table 1: Values from gold standard histology and their predictions errors from CT (mean \pm SD) for CT.TH (N = 2922) and whole rib section properties (N = 39), with coefficients of determination (R^2) from linear regression ($p < 0.0001$ for all models)

Measure	Histology		CT	
	mean \pm SD	mean \pm SD	Pred. Error	R^2
CT.TH (mm)	0.7 \pm 0.4	0.7 \pm 0.3	-0.013 \pm 0.167	0.84
CT.AR (mm ²)	20.4 \pm 8.0	20.4 \pm 6.3	0.03 \pm 2.67	0.91
TT.AR (mm ²)	64.1 \pm 20.8	67.1 \pm 21.5	3.00 \pm 1.16	1.00
ES.AR (mm ²)	43.7 \pm 17.1	46.7 \pm 18.2	2.97 \pm 2.54	0.98
IMAX (mm ⁴)	247.5 \pm 160.3	262.8 \pm 151.6	15.34 \pm 42.69	0.93
IMIN (mm ⁴)	124.0 \pm 86.5	132.7 \pm 82.8	8.61 \pm 14.44	0.97
IRAT	2.2 \pm 0.9	2.2 \pm 0.9	-0.04 \pm 0.14	0.98
LANG (deg.)			0.63 \pm 2.45	

228 locations, and highlights the overall inter-subject and intra-subject variability in rib cross-sectional geometry.

230 3.3. Cortical bone thickness maps

The full rib cortical bone thickness map (CT.TH) was calculated for each 232 rib, and the resulting average CT.TH map from the full sample is shown in Figure 7. Accumulating all rib locations together, males and females had 234 average \pm SD CT.TH values of 0.71 \pm 0.35 mm and 0.76 \pm 0.38 mm, respectively. Assessing on a regional basis, the majority (90.4%) of rib surface 236 locations did not show a significant difference ($p > 0.05$) in cortex thickness between males and females. Therefore, CT.TH thickness maps have been 238 shown using pooled data from both sexes. This pooled CT.TH map along-

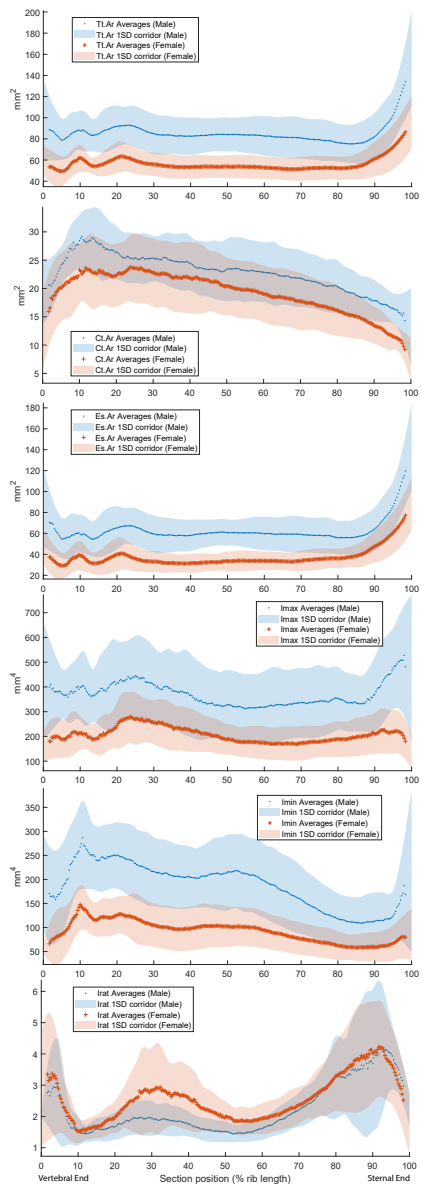


Figure 5: Measurements by sex of rib cross-sectional total area (Tt.AR), cortical bone area (Ct.AR), endosteal area (Es.AR), primary and second inertial moments (IMAX, IMIN) and IMAX/IMIN inertial moment ratio (IRAT) as a function of cross-section location from the vertebral (0%) to sternal (100%) rib ends.

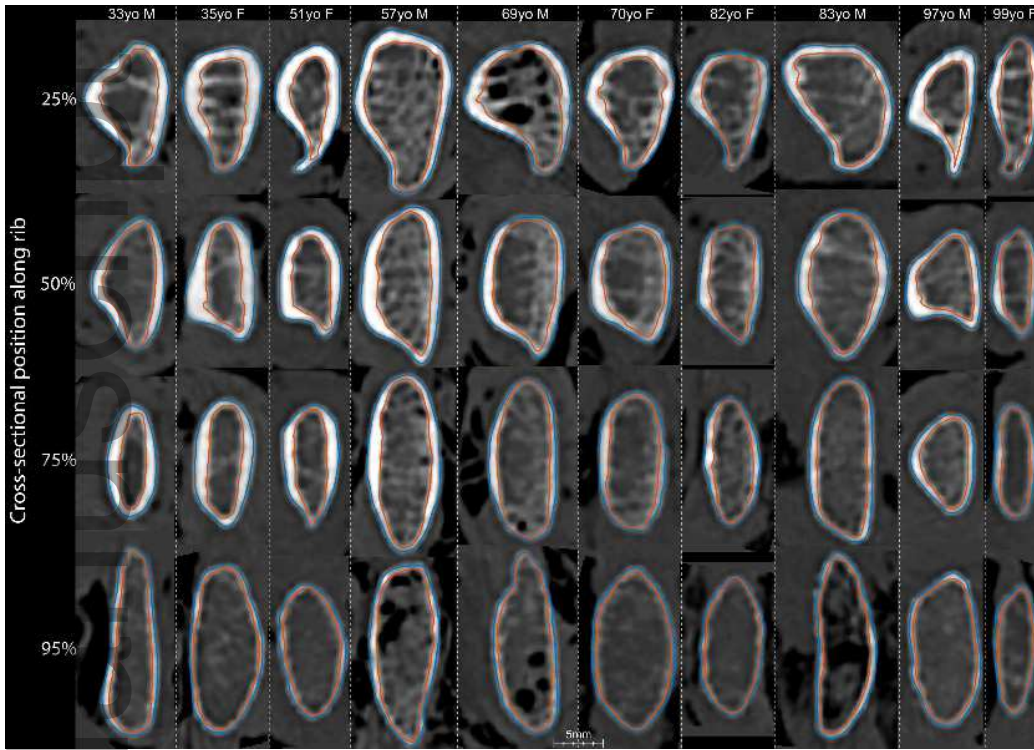


Figure 6: Exemplar cross-sections through CT volumes showing the predicted periosteal (outer) and endosteal (inner) cortical border using a range of male and female subjects (see upper labels) and rib locations (see left labels).

side separated male and female average maps are provided as supplementary
 240 information.

4. Discussion

242 Here we have assessed detailed rib cortical bone thickness and cross-
 sectional geometry from 33 adult sixth ribs. Starting with an initial approx-
 244 imate rib segmentation, the CBM method was applied using error-weighted
 smoothing and morphological filters, and average cortical bone thickness

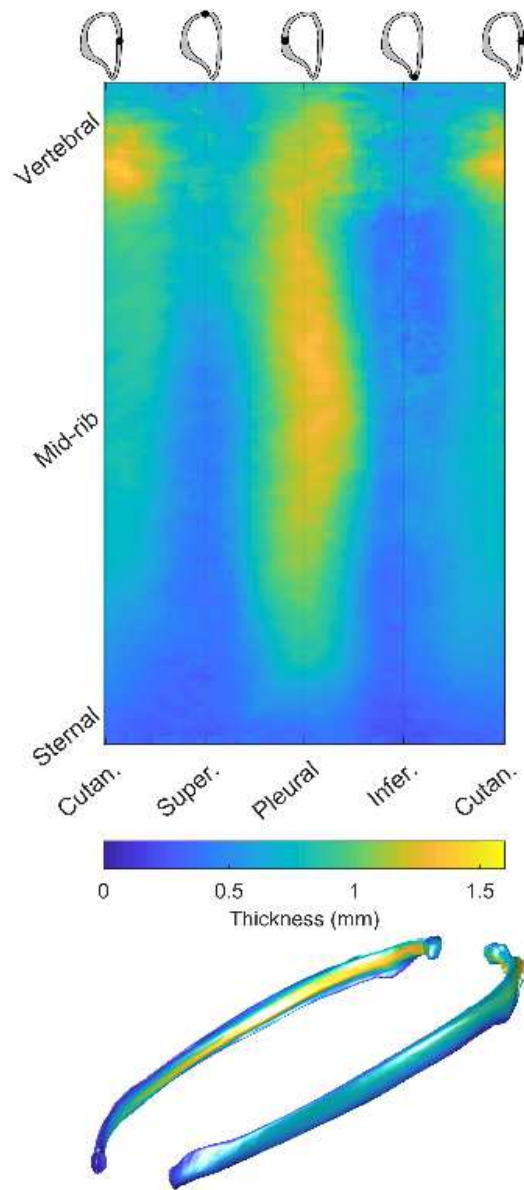


Figure 7: The average rib cortical bone thickness map in gridded form (above) and projected onto exemplar subject geometry (below).

246 maps and cross-sectional property 1SD corridors for males and females have been presented.

248 The measurement techniques used in this study are similar to those pre-
sented in Holcombe et al. (2018) with adjustments to allow for full rib image
250 volumes rather than individual rib section images. The image volumes in the
current study have $0.15 \times 0.15 \times 0.67$ mm voxels, providing resolutions that are
252 both higher (in-plane directions) and lower (out-of-plane direction) than the
highest resolution images of 0.37 mm/pixel used in Holcombe et al. (2018).
254 Correspondingly, the current accuracy and precision of measurement predic-
tions from CT are largely similar to those from this previous study which
256 found prediction errors of -0.03 ± 0.17 mm for CT.TH, -0.6 ± 1.5 mm² for
CT.AR, and 2.1 ± 1.5 mm² for TT.AR.

258 For measuring cross-sectional rib properties using CT, the current CBM-
based methodology is more accurate than traditional CT thresholding meth-
260 ods, which often overestimate the amount of bone in a given cross-section.
Perz et al. (2015) found that simple histogram-based thresholding of CT im-
262 ages resulted in average errors in TT.AR of 8 ± 3 %, while Murach et al.
(2017) used an adaptive histogram-based thresholding technique on 19 CT
264 images of similar resolution to the current study, finding TT.AR errors of
 3 ± 11 %. Those same studies reported that their CT thresholding techniques
266 produced unacceptable CT.AR overestimations of 40 ± 12 % and 71 ± 45 %,
respectively. In the current study we see similar or improved accuracy and
268 precision in TT.AR with errors of 4.7 ± 1.8 %, and greatly reduced errors of
 0.1 ± 13.1 % for CT.AR. It is informative to note that the initial periosteal
270 border in the current study—obtained using a standard 226 HU threshold
for bone segmentation—also overestimated TT.AR on these same images by
272 21 ± 4 %.

4.1. Sex-based sectional differences

274 Results highlighted in Figure 5 show that while there are similar trends
in average rib cross-sectional properties along the lengths of male and female
276 ribs, there were significant sex-based differences in their magnitudes with
males ribs being larger in terms of all area and inertial measurements and at
278 a large majority of positions along those ribs. Comparing results for CT.TH,
on the other hand, sex-based differences were less pronounced with male ribs
280 having significantly thicker cortices at just 2.1% of rib surface locations (at
the $p < 0.05$ level), and female ribs having thicker cortices at 7.5% of rib
282 surface locations. Taken together, these results do indicate that the larger
CT.AR seen in males is primarily due to males having larger overall cross-
284 sectional size to their ribs than females rather than males having rib bones
with thicker cortices than females. Notably also, female subjects in this study
286 were on average older (but not significantly older) than the males, and bone
quantity in general is known to decrease with age. However, (Agnew et al.,
288 2018) found no significant decrease in average pleural or cutaneous CT.TH
with age (or by sex) on a large sample of ribs, suggesting age is likely not a
290 confounding factor here.

As seen in Figure 5, the rib cross-sectional position of highest CT.AR is
292 near the rib tubercle (approximately 10% rib length), and a steady reduction
in CT.AR is seen from this location towards the sternal end. Rib TT.AR,
294 however, remains relatively constant across most of a rib's length before in-
creasing sharply towards its sternal end, despite variability in the qualitative
296 shape seen in Figure 6. Each of these observations match findings by Choi &
Kwak (2011) who measured cross-sectional areas from ribs of seven elderly

298 male cadaveric ribs. While not reporting sixth rib data, their fifth rib results
showed an average TT.AR of 91 mm^2 that was constant from the tubercle
300 to 90% of the rib's length, and a drop in CT.AR across that same region
from 26 mm^2 to 19 mm^2 . Each of these fall within the 1SD male corridors
302 obtained from the current study.

4.2. Cortical bone thickness map

304 Consistent patterns in cortical bone thickness maps were also seen across
individuals, as typified by the population average map (Figure 7). Beyond
306 the tubercle, all ribs showed local CT.TH maxima along the pleural and
cutaneous aspects and local CT.TH minima along the superior and inferior
308 aspects, with these features lessening at the most sternal end of the rib to
form uniformly thin cortices like those seen at the 95th percentile position
310 in Figure 6. The pleural aspect contained the thickest regions of bone, with
CT.TH values peaking to between 0.9 mm to 2.6 mm across the population
312 at approximately mid-rib locations.

4.3. Thickness map registration

314 When performing statistical aggregation it is important that variable sets
(in this case individual thickness maps) are spatially registered to maintain
316 correspondence between regions on the maps from different individuals. The
one-dimensional registration along the length of the rib is straightforward
318 whereby sample locations are equally spaced along the rib's central axis from
the vertebral to sternal rib ends. In this study the further registration of ro-
320 tational positions around the ribs are based only on the surface geometry of
the ribs. We have taken the general approach of positioning the pleural and

322 cutaneous registration aspects at locations where the rib section's secondary
inertial axis intersects with its periosteal border. For continuity, these loca-
324 tions were calculated at each cross-section and a smoothing spline was fitted
to provide their exact location.

326 Having registration depend only on rib surface geometry means that local
features of the thickness maps themselves (which are a product of external
328 and internal surface geometries) are not explicitly aligned. For example, all
ribs in this study were seen to have a regional maxima along or near their
330 pleural margin. Yet, these regional maxima did not align precisely to the
pleural position as determined by only that rib's surface geometry.

332 It would be desirable to have the regional maximum from the average
thickness map correspond to the average regional maxima from each con-
334 stituent rib map. In the current study, the average thickness map registered
by surface features alone actually underestimated the average of the regional
336 maxima along the mid-rib pleural aspect by approximately 0.06 mm (5%).
Similarly, the regional minima at the superior aspect of the mid-to-sternal
338 portion rib was overestimated by approximately 10% compared to the col-
lection of minima from each individual thickness map.

340 4.4. Limitations

342 A primary limitation of the current study is that its results are presented
only for sixth level ribs. With ribs of different levels in the rib cage serving
different mechanical roles, it is expected that both global and local anatomies
344 will differ accordingly. Indeed, ribs do differ by level in terms of global size
and shape (Holcombe et al., 2017; Wang et al., 2016; Weaver et al., 2014),
346 overall mechanical stiffness Kindig et al. (2011), and local cross-section (Choi

& Kwak, 2011; Wang et al., 2016). Nevertheless, the combination of CT and
348 histological image modalities used in this study allows us to establish methods
for validating the typical full-rib properties of this mid-level rib.

350 ■ This reference at the sixth rib level can be used to verify that future
measurements—which may be obtained from sources such as clinical CT
352 scans with less optimal imaging characteristics and without recourse to gold
standards for validation—are free of systematic bias due to their particular
354 imaging conditions.

The current study population covered a wide age range with the inten-
356 tion that average results are seen as typical of an adult (US) population.
In measures where significant sex-based differences were found we have cho-
358 sen to provide male and female results separately. Future work should aim
to broaden the subject population to include children, and to increase the
360 subject count so as to allow statistical analyses that incorporate other de-
mographic factors such as age, stature, body mass, and ancestry.

362 4.5. Applications and Future Work

The technique outlined in this study can be used to create accurate rib
364 endosteal and periosteal surfaces along the entire length of the rib, which
can improve simulation efforts greatly. The rib cortical bone thickness ref-
366 erence data used in current human body models is limited in terms of the
population from which it was drawn, and in terms of the geometric detail
368 that it provides. Li et al. (2010) demonstrated the positive effects of in-
cluding more specific cortical thickness variation into simulation studies, and
370 Agnew et al. (2018) quantified the significant effect that cross-sectional ge-
ometry has on ribs resistance to loading from physical tests. These studies

372 have demonstrated the need for more advanced approaches to understanding
human variation in rib properties and differential rib fracture risk between
374 individuals.

- Future work can target the improvement of HBMs by incorporating the
376 results obtained here for cortical bone thickness maps into HBM rib defini-
tions. Future efforts can also assess the geometric accuracy of such models,
378 testing their ability to represent their target population by comparing their
modeled rib geometries to the typical adult corridors published here.

380 **5. Conclusion**

Numerical models are an important tool for understanding and prevent-
382 ing traumatic injuries to the chest, and the ribs form a key structural model
component. Models are typically developed using CT image data, but tradi-
384 tional CT segmentation methods have been inadequate for obtaining accurate
cortical bone geometry. As such, only simplified cortex data has been applied
386 based on limited available literature from higher resolution sources.

Here we have applied a Cortical Bone Mapping (CBM) methodology to
388 whole human ribs, assessed the accuracy of these techniques against gold
standard measurements from histology, and presented detailed population-
390 based data for rib cortical bone thickness and for rib cross-sectional proper-
ties. The population data presented here for rib cortical bone thickness and
392 cross-sectional area can be used directly to assess and improve the veracity
of current FE models of human ribs. Finally, results here can validate future
394 steps towards personalized and population-based geometry of human ribs
from more broadly sampled yet less detailed image data such as live-subject

396 clinical CT scans.

Disclosures

398 ■ The authors report that there are no conflicts of interest which might affect this work.

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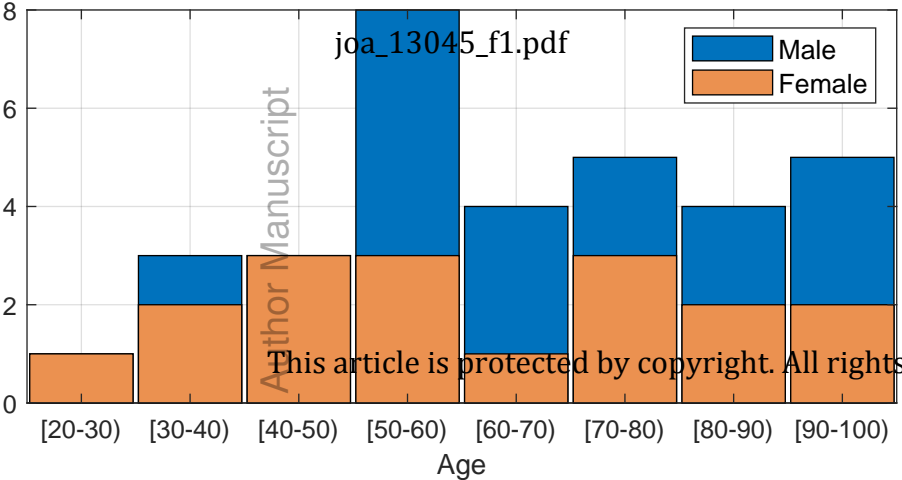
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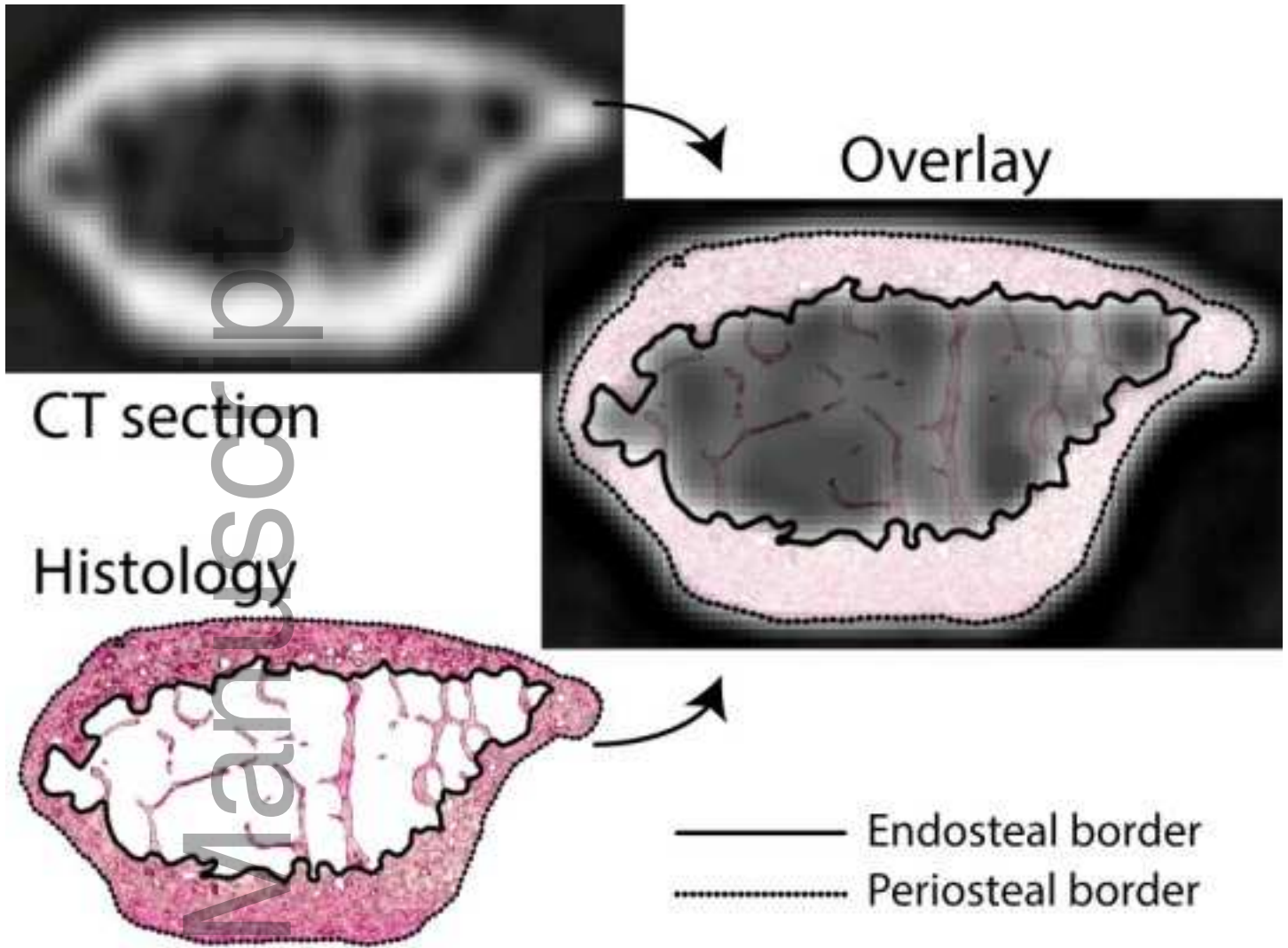
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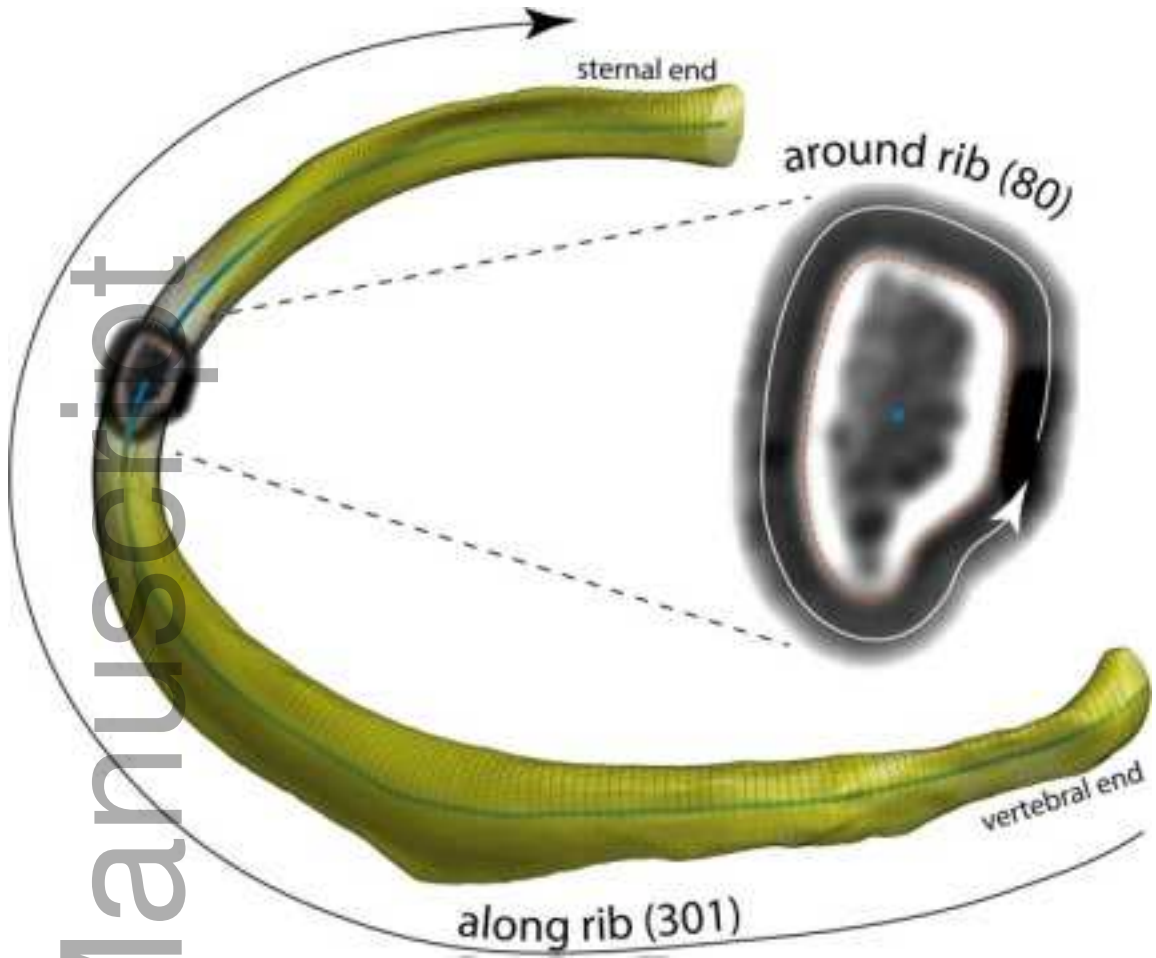
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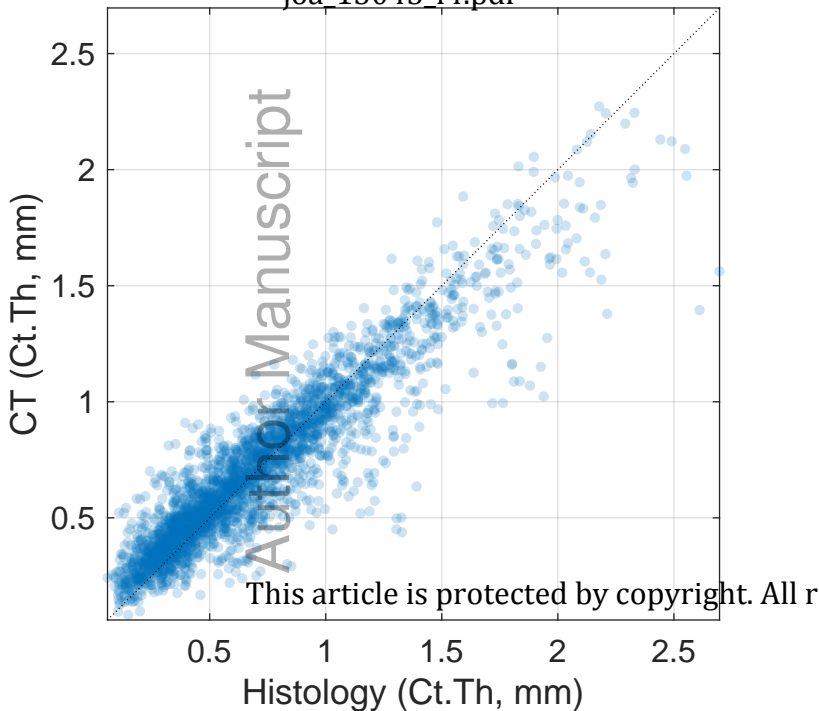


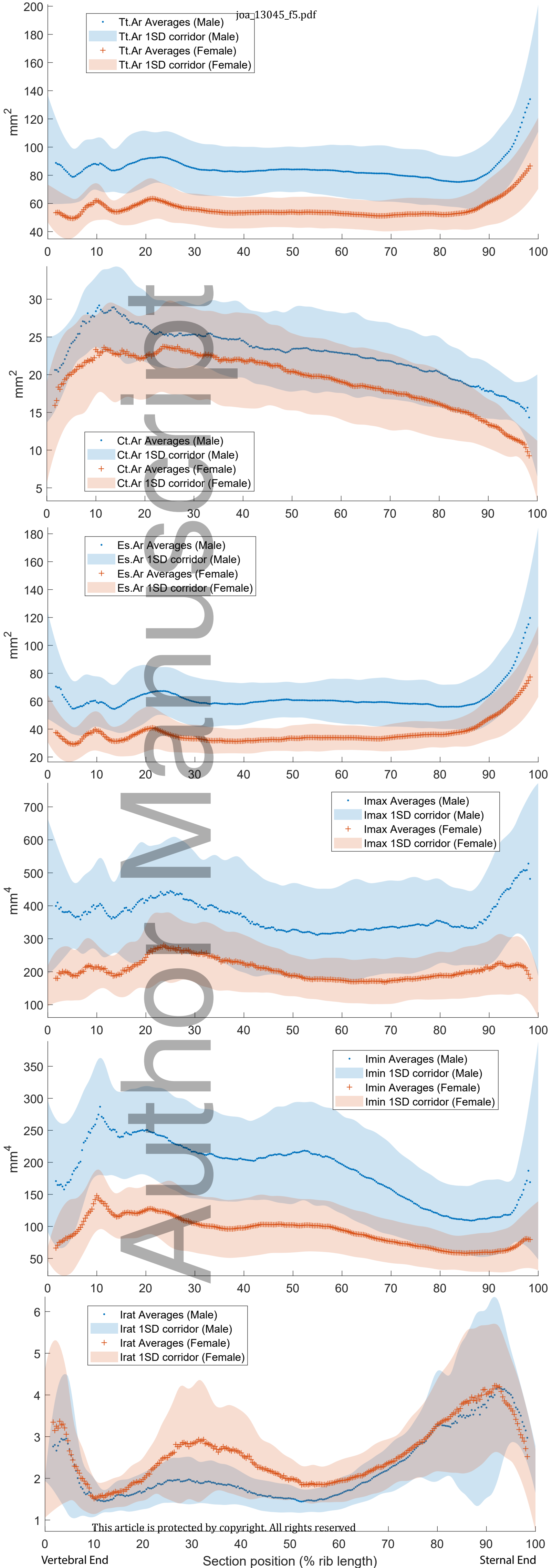


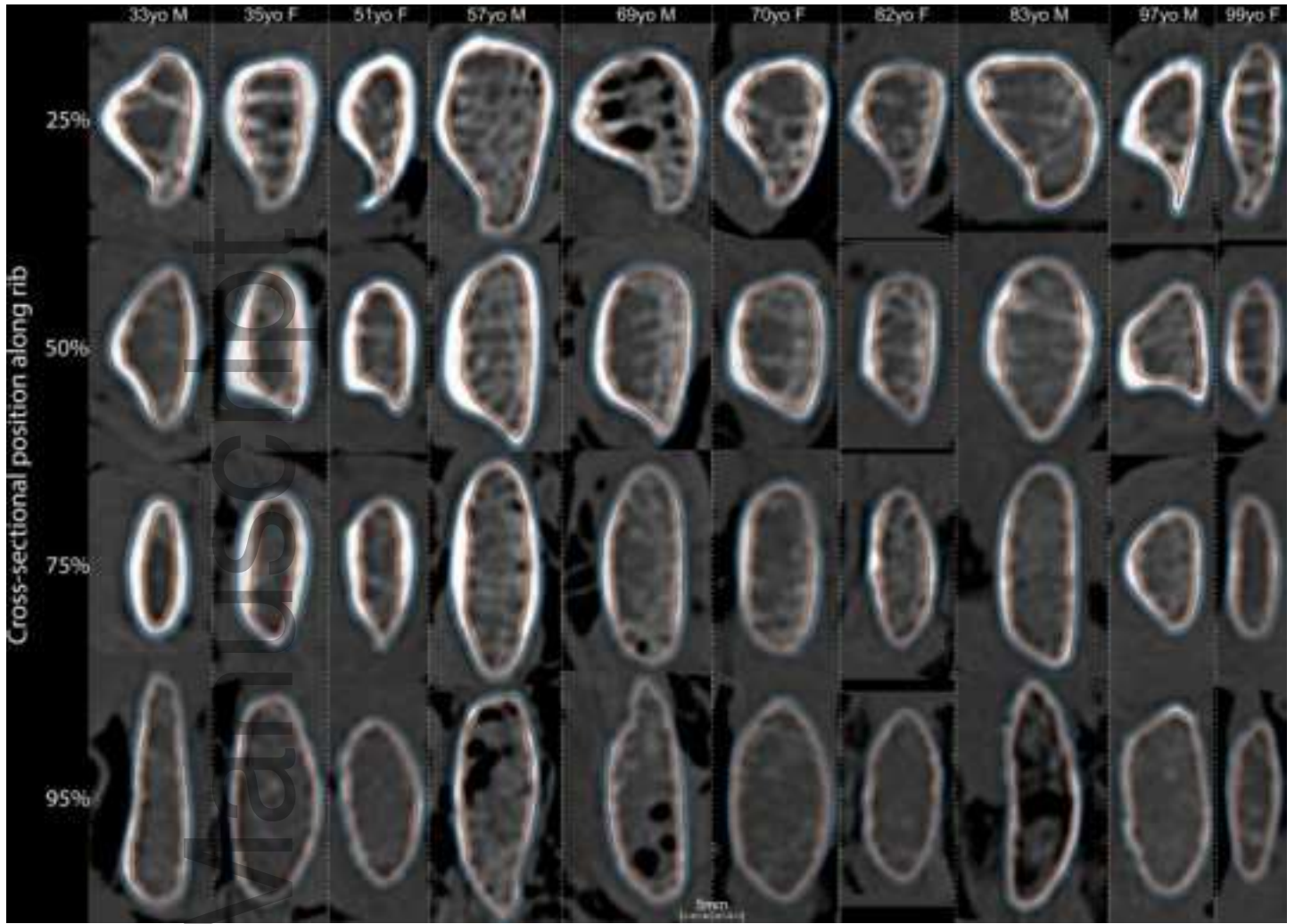
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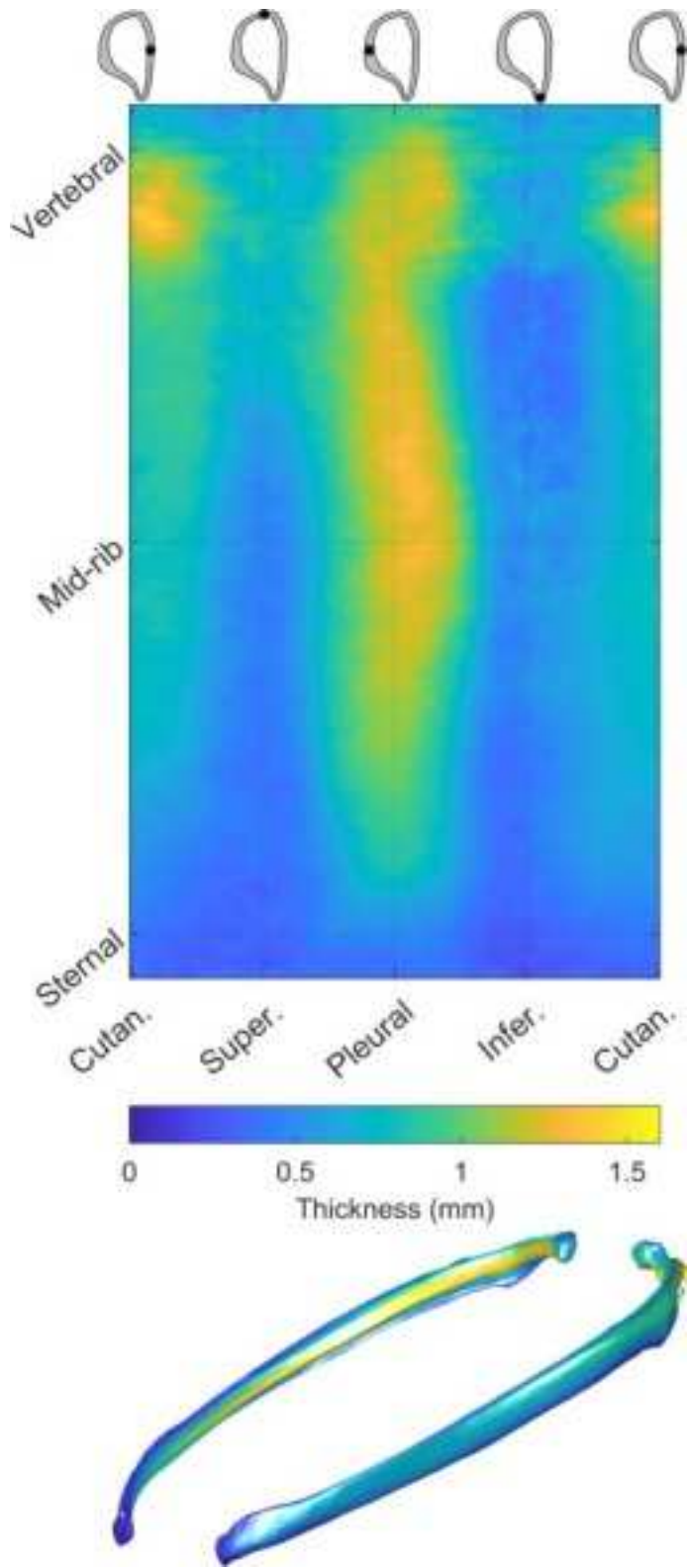






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