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Community Challenge towards Consensus on Characterization of Biological Tissue: C⁴Bio's First Findings

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17 **Abstract**

18 This study investigates methodological variability across various expert labo-
19 ratories worldwide, with regards to characterizing the mechanical properties
20 of biological tissues. Two testing rounds were conducted on the specific use
21 case of uniaxial tensile testing of porcine aorta. In the first round, 24 labs
22 were invited to apply their established methods to assess inter-laboratory
23 variability. This revealed significant methodological diversity and associated
24 variability in the stress-stretch results, underscoring the necessity for a stan-
25 dardized approach.

26 In the second round, a consensus protocol was collaboratively developed
27 and adopted by 19 labs in an attempt to minimize variability. This involved
28 standardized sample preparation and uniformity in testing protocol, includ-
29 ing the use of a common cutting and thickness measurement tool. Despite
30 protocol harmonization, significant variability persisted across labs, which
31 could not be solely attributed to inherent biological differences in tissue sam-
32 ples.

These results illustrate the challenges in unifying testing methods across
different research settings, underlining the necessity for further refinement
of testing practices. Enhancing consistency in biomechanical experiments
is pivotal when comparing results across studies, as well as when using the
resulting material properties for in silico simulations in medical research.

33 *Keywords:* Biomechanical characterization, Standardization,
34 Methodological variability, Uncertainty

1. Introduction

Characterization of the mechanical properties of biological tissue serves many purposes. Firstly, it can help to improve our fundamental understanding of physiological function, due to a strong structure-function relationship for many organs and tissue systems. Secondly and as a consequence of the former, ageing as well as various diseases are known to alter the mechanical properties of the involved tissues, such that quantification of these alterations can aid diagnosis and monitoring. Thirdly, in the field of tissue engineering and regenerative medicine, it is imperative that the engineered tissues replicate the mechanical behaviour of native tissues to ensure proper functioning. Likewise mechanical data informs the design of implants, prosthetics, and surgical tools to ensure compatibility with tissue mechanics and minimize adverse reactions. Indirectly, mechanical properties can be used to inform constitutive models which can in turn be used in computational models, for various applications in the field of *in silico* medicine (Motiwale & Sacks, 2025). However, although the scientific literature therefore abounds with articles experimentally characterizing the mechanical properties of biological tissues, there are still no widely recognized testing standards for these experiments. This shortcoming has consequences for the interpretability of the results, especially when comparing across studies. Moreover, when this experimental data is used to derive material properties used as input parameters to *in silico* models, the associated error and uncertainty propagates into these simulations.

Simulation-driven medical device development and *in silico* trials are gaining importance (Pappalardo et al., 2022; Viceconti & Emili, 2024). Clearly, the quality of these simulations is of utmost importance if *in silico* medicine is to take its rightful place in medical research and development. Consequently and analogously to the well-known concepts of ‘good medical practice’ and ‘good laboratory practice’ (Robertson & Williams, 2009; Stevens, 2003), there is a growing emphasis on the development of guidelines for ‘good simulation practice’. In 2018, the American Society of Mechanical Engineers introduced a standard known as ‘ASME V&V 40 - Assessing Credibility of Computational Modeling through Verification and Validation: Application to Medical Devices’ (ASME, 2018), and in 2023, the FDA issued a guidance document entitled ‘Assessing the Credibility of Computational Modeling and Simulation in Medical Device Submissions’ (FDA, 2023). For example, a reporting checklist for verification and validation of finite element analysis was

made more specifically for orthopaedic and trauma biomechanics (Oefner et al., 2021).

Such guidelines span various aspects related to model development, verification, calibration and validation, and focus their attention on the quantification of the uncertainty that accumulates in each of these aspects. Apart from modeling assumptions and discretization errors, it is intuitive that model input data, whether used to calibrate model parameters or validate the model outcomes, are an important source of uncertainty. Indeed, credible numerical simulations require input parameters and underlying acquisition methods that are both traceable and reliable (ASME, 2018). However, where numerical analysts are usually well-equipped to test the validity of their modeling assumptions or to minimize discretization errors, not every numerical analyst has the expertise and/or facilities to assess the quality and uncertainty of their model input data, especially when it involves data related to biological tissue properties.

Ideally, the required material properties and their uncertainty should come from shared, trusted databases. However, various challenges are faced when attempting to create such a database. Firstly, depending on the context of use of a simulation (Viceconti et al., 2021), different types of material properties might be required (mechanical, thermal, electrical, etc.). Even when focusing on mechanical properties alone, a simulation might require properties related to a specific material constitutive law and its characteristics, such as the tensile elasticity, viscoelasticity, anisotropy, compressibility, nonlinearity, ultimate tensile strength, pre-stress state etc. Each of these properties can, in turn, be acquired by various test setups and methods, such as uniaxial or biaxial tensile tests, compression tests, shear tests, indentation tests, dynamic mechanical analysis, and creep tests. The selection of the appropriate testing method depends on the tissue type, its anatomical location, and the desired application/objective to which the simulation responds. Moreover, apart from the actual testing method, the sample collection, preservation and preparation method will also influence the resulting properties, even if such effects are still under debate (Blaker et al., 2024; Oswald et al., 2017; Chow & Zhang, 2011; Stemper et al., 2007).

The second challenge relates to the population- and subject-related variability. Indeed, material properties of biological tissues exhibit significant

110 variability due to physiological and demographic factors. Sex-related vari-
 111 ations and ethnic diversity contribute to changes in tissue composition and
 112 structure, leading to differences in material properties for a given tissue type
 113 (Smoljkić et al., 2023; Åstrand et al., 2011). Age-related changes, such as
 114 decreased elasticity and increased brittleness in tissues like skin and bone, or
 115 other pathological conditions, induce further variability (Kirilova-Doneva &
 116 Pashkouleva, 2022; Mirzaali et al., 2016). Even within a subject, properties
 117 will vary depending on the location within the body (Krueger et al., 2011).
 118 Ideally, a material properties database would encompass this broad repre-
 119 sentation, which would require the collection of an extremely large range
 120 of experimental data. Moreover, in certain research communities, there ap-
 121 pears to be an unsubstantiated trust in the ‘known values’ taken from the
 122 literature. As Hammer & Klima (2019) also noted in their review, the imple-
 123 mentation of further region-specific studies is partially inhibited, as reliance
 124 is placed on a supposedly existing broad database, instead of carrying out
 125 new studies adapted to the research question. Using numerical studies of
 126 the biomechanics of the sacro-iliac joint as an example, they point out the
 127 frequent nested literature references and the often accompanying decrease in
 128 relevance of the basic source for the intended loading scenario.

129
 130 With that in mind, perhaps the most important challenge related to build-
 131 ing up such a database is related to methodological variability. Indeed, even
 132 when comparing literature results of the same tissue type of a similar an-
 133 imal strain or patient group, tested according to - reportedly - the same
 134 method, results vary widely. As mentioned, there are currently no estab-
 135 lished standards for measuring the material properties of biological tissue,
 136 nor is there a unified approach for sample preparation and storage, or for
 137 reporting and assessing the obtained results. Some groups have proposed
 138 guidelines for specific aspects or applications, e.g. Kurz et al. (2023) have re-
 139 cently suggested a standardized approach for characterization of the human
 140 lumbopelvic system, Wale et al. (2021) have investigated sample preparation
 141 and optimal sample shape to induce reproducible failure for tensile testing of
 142 musculoskeletal soft tissues, Scholze et al. (2020) have proposed a clamping
 143 system for simple and reproducible sample clamping. Though valuable, these
 144 studies are limited to a single research group, and it remains to be seen if and
 145 how they will be adopted by the scientific community. Lin et al. (2024) have
 146 recently performed a systematic review of uniaxial tensile testing of human
 147 soft tissues across research groups. They found large variations in sample

148 shape and especially underreporting of various aspects of the protocol, in-
 149 cluding the clamping mechanism. Similarly, Fehervary (2018) investigated
 150 variations in planar biaxial testing of arterial tissue, both in experimental
 151 setup and data processing, and formulated guidelines for testing and report-
 152 ing thereof. Although this lack of standardization and high variability is
 153 commonly known, to the best of our knowledge, a quantitative analysis of
 154 the current degree of methodological variability has not yet been attempted.
 155 It is therefore an essential first step towards standardization, ultimately lead-
 156 ing to reliable material properties, including their uncertainty, which can be
 157 used in computational analyses for *in silico* medicine.

158
 159 In 2019, the Virtual Physiological Human institute together with the Avi-
 160 cenna Alliance launched a task force on tissue characterization, out of which
 161 the C⁴Bio (C⁴Bio, 2025) initiative was born, an acronym for ‘Community
 162 Challenge towards Consensus on Characterization of Biological Tissue’. Con-
 163 sidering the wide range of properties, tissue types and testing methods, the
 164 decision was made to direct the initial focus to ‘simple’ uniaxial tensile testing
 165 (i.e. the method most commonly found in literature) of porcine aorta. This
 166 paper first describes the methods employed to compare the results obtained
 167 by 24 expert laboratories from around the globe who characterized biological
 168 tissues and synthetic samples during two test rounds. In a first testing round,
 169 laboratories were instructed to use their own established method, to allow
 170 an unbiased evaluation of existing methodologies. In a second testing round,
 171 participants worked together to create a ‘consensus methodology’ and were
 172 then instructed to perform their tests accordingly.

173 2. Materials & methods

174 2.1. Participant recruitment & overall methodology

175 The initiative was announced in the fall of 2020 through various com-
 176 munication channels. All research groups with experience in material char-
 177 acterization of biological tissue were invited to join, provided the following
 178 prerequisites:

- 179 • the availability of infrastructure suitable for uniaxial tensile testing of
 180 samples within a length range of 10-60 mm and a linearized Young’s
 181 modulus range of 0-5 MPa,
- 182 • the clearance to test biological tissue in the laboratory,

- 183 • a demonstrated experience in mechanical experiments on biological tis-
184 sue through scientific references,
- 185 • the willingness to publicly share the experimental results.

186

187 In round 1, 24 laboratories from all over the world participated (see Fig-
188 ure S1 in the supplementary material, section Appendix A, for a map), and
189 19 of them further participated in round 2. The participants that were no
190 longer able to participate indicated that this was due to a lack of budget
191 and/or person power.

192

193 For both testing rounds, all samples were procured and prepared cen-
194 trally and shipped as described in Section 2.2. In round 1, 24 laborato-
195 ries performed mechanical testing and analysis according to their preferred
196 methodology, with only a limited number of instructions given (see Section
197 2.3). After testing, raw and processed data were collected and analysed cen-
198 trally according to the methods described in Section 2.4. When providing the
199 raw and the processed data, participants were also asked to fill in a survey,
200 querying on various aspects of the testing methodology they used. Following
201 the central analysis, the observations were discussed with the participants
202 over multiple online meetings, and a consensus protocol was established and
203 used in round 2. The full protocol as shared with the participants can be
204 found in the supplementary material (section Appendix A).

205 *2.2. Sample preparation & shipping*

206 *2.2.1. Biological tissue*

207 In both testing rounds, 2 sets of a proximal and a distal segment of one
208 porcine descending aorta was prepared per participant at FIBEr, the KU
209 Leuven core facility for Biomechanical Experimentation. Each round, aortas
210 were collected from 70 animals from a local slaughterhouse, and brought to
211 FIBEr in physiological medium (saline solution) at 4°C. Care was taken to
212 minimize biological variability, by harvesting material as much as possible
213 from animals of the same strain, average weight and age, and distributing
214 these randomly over the participants. From each aorta a distal and proxi-
215 mal segment of approximately 2 cm in diameter and 5 cm in length was cut,
216 with a small proximal anterior incision, to allow tracking of the orientation of
217 the sample (see Figure S2 in the supplementary material, section Appendix

218 A). Samples were placed in individual containers with physiological medium,
219 while keeping track of the animal and the location (proximal vs distal). All
220 samples were transferred to a -80°C freezer inside the containers within 8
221 hours of excision from the animal, and stored in FIBEr until shipment. The
222 samples were shipped to each of the participants on dry ice in styrofoam
223 boxes. Participants were instructed to keep the aortic tissue in frozen condi-
224 tions (i.e. at -80°C) until they were ready for testing, or to start the thawing
225 and testing process immediately if no freezing capability was available in
226 the laboratory. Upon receipt of the package, the participants verified that a
227 sufficient amount of dry ice was still present in the box to ensure proper sam-
228 ple preservation. For all samples, the duration of frozen storage was recorded.
229

230 2.2.2. *Synthetic samples*

231 In round 1, two sheets of synthetic elastomer material of approximately
232 $4.5 \times 9 \text{ cm}$ were shipped to the participating research group along with the
233 biological tissue, in casu ‘Leartiker-K73’ and ‘Leartiker-K51’, provided by
234 Leartiker s.a. (Spain).

235 In round 2, to use a material with more biologically relevant properties
236 as compared to round 1, samples were 3D printed by Materialise NV (Bel-
237 gium) using the company’s proprietary HeartPrintFlexPlus (HPF+) material
238 (Schickel et al., 2019) in the dogbone shape as agreed upon in the consensus
239 protocol (see section 2.3.2). In both rounds, the synthetic material was kept
240 and shipped in dry conditions at ambient temperature.
241

242 2.3. *Mechanical testing*

243 2.3.1. *Round 1*

244 In the first testing round, only minimal instructions were given to par-
245 ticipants. This was done on purpose to allow for an objective comparison
246 of the participating laboratories preferred methodologies. It was requested
247 that all tests were executed by the same person, to avoid any inter-user vari-
248 ability. Also, the same equipment and methodology had to be used for both
249 the aortic and the synthetic samples. The only exception in the process was
250 that the synthetic material should always be tested in dry conditions.
251

252 The evening before the testing day, the participating groups placed the
253 aortic specimen in a fridge (4°C) to thaw overnight. This allowed a thawing

254 time of around 16 h, whereby the exact time per specimen was recorded.
255 Next, the specimens were divided into test samples. From each tubular seg-
256 ment, at least 3 uniaxial test samples were prepared, with geometry and
257 cutting method varying per participant. Proximal and distal segments were
258 used to prepare circumferentially and longitudinally oriented samples, re-
259 spectively. Participants were instructed to take a picture or make a sketch to
260 indicate the location of each of sample with respect to the original specimen.

261
262 For the synthetic specimens, at least 6 samples were cut out of each pro-
263 vided sheet, in a direction of choice since the material is considered to be
264 isotropic. Figure S3 in the supplementary material (section Appendix A)
265 gives an overview of all samples (biological and synthetic) tested per partic-
266 ipant.

267
268 For each (biological and synthetic) sample, the undeformed sample thick-
269 ness, undeformed sample width in the neck region, and undeformed length
270 of the neck region were measured according to each participant’s method-
271 ology. Finally, participants also selected individually the appropriate loading
272 protocol and data processing methodology to determine the elastic stiffness
273 moduli at various strain levels and the ultimate tensile strength.

274 275 2.3.2. Round 2

276 In round 2, the same thawing instructions were given as in round 1. After
277 thawing, test samples were excised from the aortic segments and at least three
278 circumferentially oriented samples were prepared from the proximal segments
279 and three longitudinally oriented samples from the distal segments. This
280 time, the samples were cut into a dogbone shape using a cutting tool that
281 was shipped to the participating research group along with the tissue. The
282 dogbone shape and cutting tool are shown in Figure 1a and b and more details
283 are available in 2.2. The time required for each phase in the preparation and
284 testing protocol was logged for each sample.

285 Participants were required to measure the undeformed width in the neck
286 region and overall length of each sample using calibrated images taken with a
287 high-resolution camera on millimeter paper. For the thickness measurement,
288 the sample was inserted into a tool that was provided to the participants,
289 shown in Figure 1c. This tool gently squeezes the sample with the help of
290 identical elastic bands between two plates with known dimensions, such that

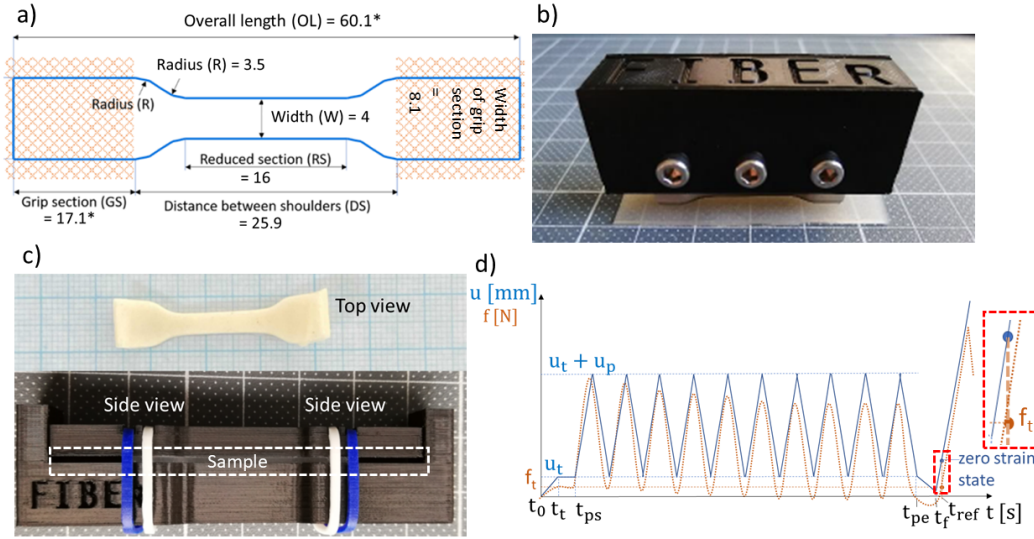


Figure 1: In round 2, a) samples were cut into a dogbone shape as agreed upon by all participants. *These dimensions can be increased if more gripping surface is required. All dimensions are in millimeter. b) A cutting tool was provided to ensure consistency in the shape. c) Calibrated pictures were taken during sample preparation for dimension measurements. Top: aortic sample on millimeter paper. Bottom: synthetic sample in a custom-made thickness measurement tool, tightened with elastic bands. d) Loading protocol for round 2, showing actuator displacement u and expected reaction force f as a function of time t . Time t_t corresponds to the time at which the threshold force of f_t is reached for the first time. Times t_{ps} and t_{pe} correspond to the start and end of the preconditioning cycles, during which a preload amplitude u_p is applied for 10 cycles. Time t_f corresponds to the start of the final upward ramp and t_{ref} corresponds to the assumed zero strain state.

291 the sample thickness can be derived from a side view image focussed on the
 292 tissue surface.

293

294 Participants were advised to use digital image correlation (DIC) to track
 295 sample deformation during the test, which required application of a speckle
 296 pattern on the sample's intimal surface, by sprinkling graphite powder on a
 297 piece of paper and subsequently placing the intimal side of the sample onto
 298 the paper. Alternatively, if no DIC system was available, participants used
 299 a marker tracking technique, placing four markers at ~ 6 mm intervals in the
 300 central area of the sample.

301 Similar to round 1, the participants were asked that all samples were

302 tested using the same equipment by a single operator. After sample accli-
 303 matisation in saline solution at 37°C for approximately 10 minutes, the load
 304 cell was zeroed. For a horizontal set-up, this was done before mounting the
 305 sample, whereas for a vertical set-up, this was done after the sample was
 306 fixed in the upper clamp. During testing, the biological samples were im-
 307 mersed in a saline bath at 37°C, whereas the synthetic samples were kept in
 308 dry condition.

309 The specific loading protocol as depicted in Figure 1d was applied. After
 310 reaching a threshold force $f_t = 0.1$ N, 10 preconditioning cycles were applied
 311 in the physiological loading regime with an amplitude $u_p = 8$ mm. This
 312 was followed by a displacement-controlled ramp loading until failure at a
 313 displacement rate of 1.3 mm/s. During the test, resulting displacement,
 314 force, applied strain, local samples strains, and where applicable, images,
 315 were acquired with a minimum sampling rate of 10 Hz. The full description
 316 of the loading protocol is provided in the supplementary material (section
 317 Appendix A, see `Instructions_2nd_testround.pdf`).

318 *2.4. Data analysis*

319 *2.4.1. Method variation*

320 In addition to the testing data submission, in round 1 the participants
 321 were also requested to fill in a questionnaire to report their choices regarding
 322 pre-, intra-, and post-testing conditions. These included pre-testing stor-
 323 age conditions, test sample preparation and shape characterization, testing
 324 device characteristics, and testing protocol.

325 While the consensus protocol was used in round 2, the research groups
 326 were asked if they had any intentional or unintentional deviations from the
 327 protocol to identify potential sources of variability.

328 *2.4.2. Data submission*

329 For each test, a table as shown in supplementary material was filled in,
 330 which differed slightly depending on the testing round. Apart from the afore-
 331 mentioned sample geometry information, actuator displacement, load cell
 332 values and local stretch estimation in the testing direction were obtained as
 333 a function of time. The participants were also asked to indicate whether the
 334 sample ruptured in the central area or at the clamp site.

335

2.4.3. Data processing

Dimensions of interest. In round 2, the method and software used to measure dimensions from the calibrated images (see section 2.3.2) were left to the teams' discretion.

Engineering stress & stretch. The raw force data was divided by the cross-section area of the reduced section to compute the raw engineering stress. The stretch data was retrieved by the participants using their own DIC analysis or marker tracking software. Subsequently, the raw stress and stretch data of all groups were processed by two data analysts to avoid any processing errors, before being synthesized by just one of them using Matlab 2019a and 2023a (Mathworks.com): only the test data between the end of preconditioning and the highest stress value was kept, filtered (forward-backward moving average with a window of 5%) and resampled to 1000 points. A qualitative double-check of the cleaned data against the raw data was performed. Per step of 0.1 stretch ratio, a tangent modulus (TM) was computed using a least-squares line-fitting.

Comparison of stress/stretch curves. Various parameters for each team and per type of sample (i.e. a set) were computed to compare the results. The mean and standard deviation of the stress was computed at every 0.1 interval of the stretch, and the mean and standard deviation of the stretch was then computed at every 0.1 interval of the stress. Next, the mean stress/stretch curve a set of curves was computed point per point. I.e, for any given stress/stretch point of the 1000 points discretizing each curve, the mean of all the values at that point within the set was used.

Analytic uncertainty on stress. A theoretical uncertainty on the obtained stress values $U(\sigma)$ was determined for each team and sample type, incorporating measurement uncertainties as follows:

$$U(\sigma) = \sqrt{\sigma^2 \left(\left(\frac{U(N)}{\bar{N}} \right)^2 + \left(\frac{U(w)}{\bar{w}} \right)^2 + \left(\frac{U(t)}{\bar{t}} \right)^2 \right)} \quad (1)$$

In the above equation, σ represents the engineering stress (in MPa), N is the normal force and w and t are the undeformed width and thickness of the sample, respectively. The measurement uncertainty on the force, $U(N)$, is considered the same for all groups and on average assumed to be 0.1 N (based on a ISO 376 class 1 load cell measuring around 20 N), while the uncertainties on the width, $U(w)$, and thickness, $U(t)$, are both set to 0.02

369 mm (based on the maximum permissible error for external jaws on calipers
370 with measurement length 0-5 cm, according to ISO 13385-1). The notation
371 $\bar{\cdot}$ represents the mean value of each parameter for a given team and sample
372 type.

373 **Zero-strain state** In the consensus protocol, the zero-strain state of the
374 final upward ramp was considered to correspond to the time point t_{ref} (see
375 Figure 1d) at which the threshold force $t_f = 0.1$ N was reached. However,
376 during post-processing to increase consistency, this was modified to the strain
377 state corresponding to the time point at which the measured stress was equal
378 to $2*U(\sigma)$ (see equation 1). This was equal to 0.03 MPa for biological samples
379 and 0.06 MPa for synthetic ones.

380 3. Results

381 3.1. Submissions

382 In round 1, 20 participants successfully submitted their results for both
383 the biological and synthetic data. In round 2, we received 17 successful
384 submissions for the synthetic samples and 13 for the biological samples.

385 3.2. Method variation

386 The questionnaire results, regarding the choices of the research groups
387 in round 1 for pre-, intra-, and post-testing conditions, are given in Figure
388 2 together with how many groups (in %) employed a certain technique or
389 device.

390 Although the research groups were asked to use the consensus protocol
391 in round 2, intentional or unintentional deviations from the protocol were
392 observed by some groups and summarized in supplementary material (sec-
393 tion Appendix A). For example, as the surface of the synthetic samples
394 started showing cracks during testing, the stretch measurements could not
395 be obtained with the intended DIC approach. Hence, many research groups
396 reported the clamp-to-clamp distance instead. Another common deviation
397 related to the removal of supposedly surrounding connective tissue in the
398 biological samples. Some groups removed this tissue, possibly removing the
399 adventitia along with it, others did not. Besides these, there were some
400 occasional, unintentional deviations from the loading protocol, such as dif-
401 ferent preconditioning displacement or time point at which load-taring was
402 performed.

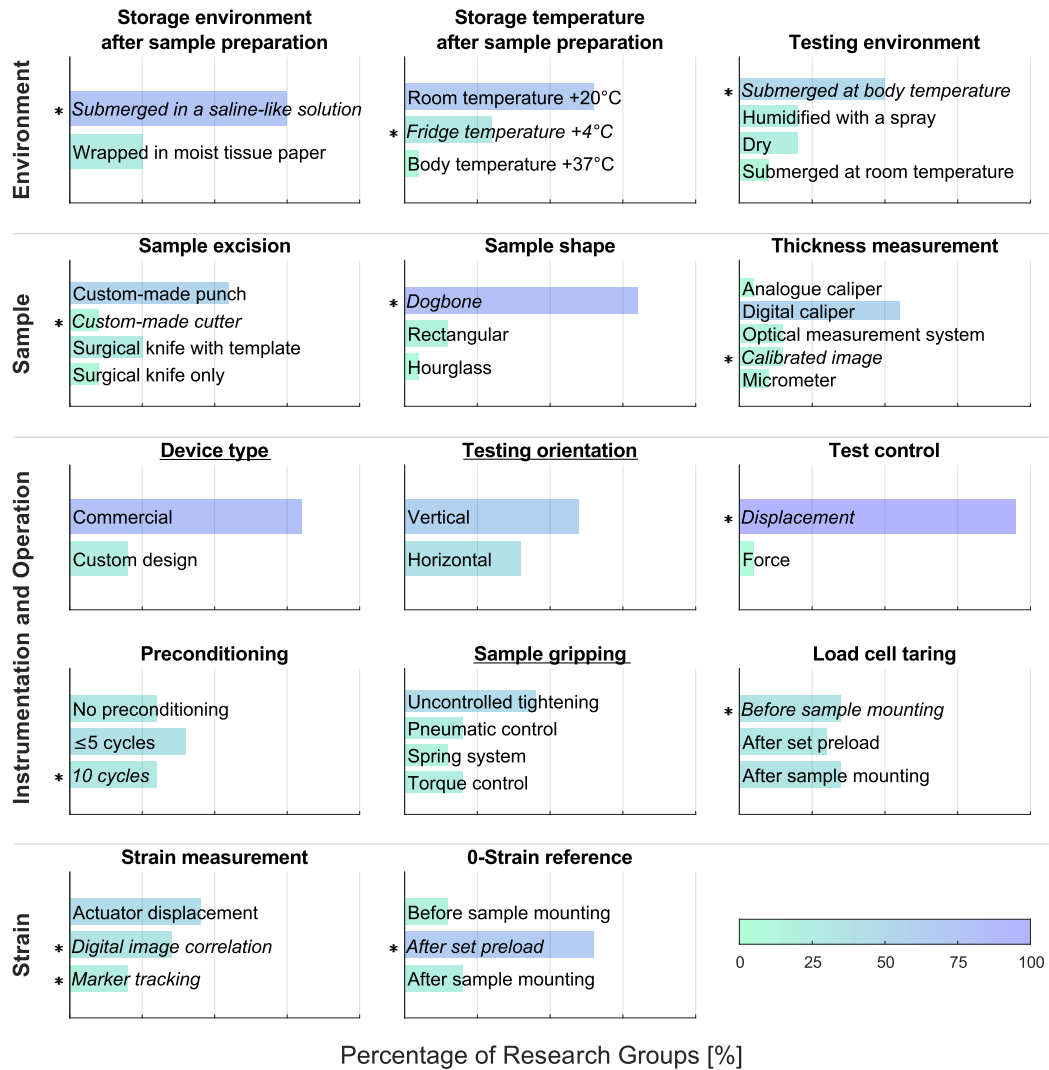


Figure 2: Method variation in round 1 and the percentage of groups using a certain approach. Parameters fixed in the consensus protocol of round 2 are marked with '*'. Test criteria not constrained by the consensus protocol in round 2 are underlined.

3.3. Main results

An overview of all the engineering stress vs. stretch curves obtained by the participants for both testing rounds is given in Figure 3. These curves allow to obtain an impression of the overall variability in stress-stretch response and ultimate strain values of the tested samples.

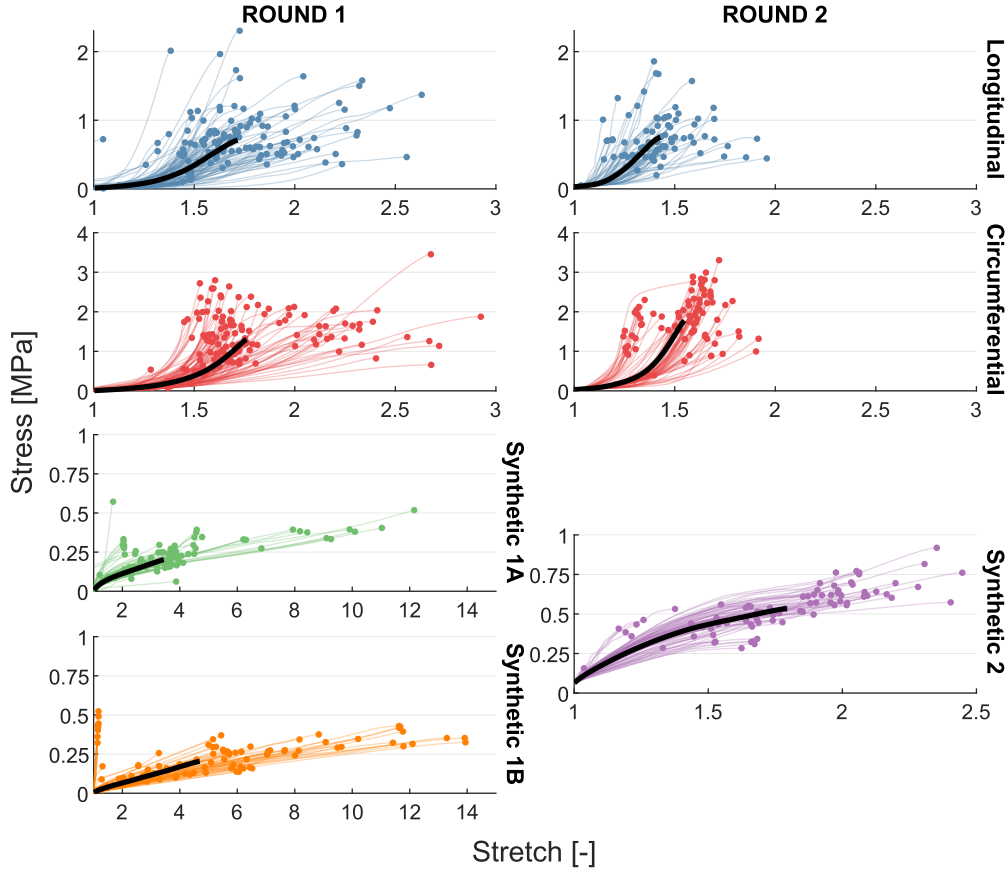


Figure 3: Engineering stress (in MPa) as a function of stretch obtained from uniaxial tension test until rupture, categorised by specimen type and separated per testing round. The average of all curves corresponding to each specimen type and testing round is represented in black.

408 Figures 4a and 4b show the stress vs. stretch curves obtained per research
 409 group on aortic tissue in the longitudinal direction during round 1 and 2 re-
 410 spectively. The corresponding curves for the circumferential direction and for
 411 the synthetic samples can be found in the supplementary material (Figure S4
 412 and S5). The graphs also show the standard deviations of stress and stretch
 413 values at certain intervals. Figure 5 zooms in on this standard deviation and
 414 how it varies between research groups and between rounds. Now once again
 415 grouped for all participants, the graphs show the coefficient of variation of
 416 stress and stretch at two specific stretch levels. The graph also shows the

percentage of research groups that fall under the arbitrary threshold of 0.33 in terms of this coefficient of variation.

The distributions of the tangent moduli measured at 1.2 and 1.4 stretch are illustrated on Figure 6 and 7, respectively. To consider both the stress and strain measurement variabilities in the assessment of the tangent moduli, Figure 8a illustrates the coefficient of variation of stress and stretch at 1.2 and 1.4 stretch, for each group and for the various sample types. This figure discriminates the groups that fall under the arbitrary threshold of coefficient of variation of 0.33 for both stretch and stress.

The distributions of the measurements of the thickness and the width of the samples among the groups and for both rounds are illustrated on Figure 9 and Figure 10, respectively. The coefficient of variation values of these measurements are illustrated on Figure 8b.

4. Discussion

Even for a relatively simple method like uniaxial tensile testing, the results of round 1 demonstrate staggering variability between the participating laboratories. This is true for the obtained stress-stretch curves (Figure 3), the corresponding coefficient of variation values (Figure 5), as well as the resulting tangent moduli (Figure 8a). Looking at Figure 2, one can also observe a large variation in the choices made regarding the test protocol in this first round. However, statistical analysis (not shown here) could not identify any correlation between these protocol variations and the results. This implies that the observed variability cannot be directly attributed to a single queried aspect of the protocol. Indeed, the observed variability is a superposition of biological variability, methodological (or aleatory) uncertainty *and* methodological (or epistemic) discrepancy.

4.1. Variability versus uncertainty in round 1

To minimize biological variability, aorta samples were harvested from animals of the same age, weight and strain, as detailed in section 2.2, and randomly distributed over the participants. However, even in such a population, biological variability is expected. Therefore, in an attempt to isolate the effect of biological variability, we also integrated synthetic samples into our analysis, as they suffer far less from intrinsic variability. Figure 8a shows how the overall coefficient of variation is indeed significantly reduced for the synthetic materials compared to the biological tissue, but still reaches levels

452 close to 50%.

453

454 A rough estimate of the degree of methodological uncertainty on the stress
455 measurement was made using equation 1, based on estimates of the accuracy
456 of the used load cells and geometry measurement methods. Figure 4a shows
457 how this estimated measurement uncertainty relates to the resulting stan-
458 dard deviation in stress. For most groups, this uncertainty represents only a
459 fraction of the total observed variability, even for the synthetic material (see
460 the supplementary material).

461

462 A significant portion of the observed variability must therefore be at-
463 tributed to methodological variability, where we can in turn distinguish be-
464 tween intra-research group variability (also known as repeatability) and inter-
465 research group variability (also known as reproducibility). Although one
466 might expect the former to be minimal, given the instructions to test consis-
467 tently, i.e. on the same day and by the same operator, the results indicate
468 a significant and varying amount of intra-research group variability. Conse-
469 quently, one might expect this repeatability to correlate with the reported
470 protocol variations (see Figure 2). However, as mentioned, further statistical
471 analysis did not reveal any correlation between these protocol variations and
472 the actual results.

473 To disentangle sources of variability, one could attempt to obtain infor-
474 mation on biological variability from other studies reported in literature.
475 Indeed, there are several studies that report mechanical properties of aorta.
476 For example, Shahbad et al. (2025) report on regional variations in stiff-
477 ness of the human aorta along its length, whereas Ryu et al. (2022) report
478 region-dependent stiffness parameters of porcine aorta. Both studies show a
479 response and degree of variability (per region) in the same order of magnitude
480 as our results. Of course, and this is the main message of our publication,
481 it is not possible to distinguish methodological variability from biological
482 variability in any of the studies. Indeed, there is no reason to assume that
483 other laboratories would not suffer from the same degree of methodological
484 variability as our reported intra-research group variabilities.

485 4.2. Consensus protocol and its trade-offs

486 Therefore, in round 2, we aimed to minimize methodological variability
487 of as many aspects of the protocol as possible, leading to a lengthy consen-
488 sus protocol. All participants also used the same cutting tool and thickness

489 measurement tool in this second round. Still, in this quest for a harmonized
490 protocol, we needed to consider two trade-offs. The first was a trade-off
491 between methods that are most likely to yield biologically relevant results
492 and methods with a lower complexity. The latter are more accessible for a
493 broad group of researchers, and might also induce a lower amount of vari-
494 ability. As an example, it is easier to test samples in dried conditions than in
495 physiological, submersed conditions. The latter requires mounting of a fluid
496 bath, which makes it harder to mount the sample, and, depending on the
497 orientation of the set-up, puts restrictions on the type of load cell. Although
498 the dry state of the sample might not alter the intrinsic variability between
499 samples compared to a submerged state, the extra requirements and manip-
500 ulations could induce further methodological variability. Still, we opted for
501 testing in immersed conditions. A counterexample in this respect is the fact
502 that all tissue was frozen until the test day, rather than testing the tissue in
503 fresh conditions. Whereas fresh tissue testing is, of course, more biologically
504 relevant (Chow & Zhang, 2011; Stemper et al., 2007), freezing the tissue was
505 necessary to allow for uniformity in the tissue preservation step. The second
506 trade-off was to be made between methods that are likely to yield the most
507 unbiased results and methods that were available to most participants (in
508 terms of infrastructure and personnel requirements). As an example, the
509 thickness measurement method should ideally not compress the tissue (Kim
510 et al., 2011; O’Leary et al., 2013) or correct for any compression during the
511 measurement (Schwarz et al., 2023), but requires specific hardware and soft-
512 ware that was not available to all participants, which is why we opted for
513 the supplied thickness measurement tool.

514

515 ASTM D412 is a common standard to characterize mechanical properties
516 of rubbers and elastomers using a tensile test. Our resulting consensus pro-
517 tocol deviates from this standard in various ways. Firstly, we used a lower
518 displacement rate (1.3 mm/s vs. 500 mm/min) to approximate ‘physiological’
519 loading conditions. Secondly, where the standard suggests a fixed temper-
520 ature, we allowed variations between laboratories in ambient temperature
521 when testing the synthetic samples, which might contribute to the variabil-
522 ity of these measurements. Due to the limited size of an aorta, our chosen
523 dogbone shape was smaller than the standard, especially regarding the width
524 of the grip section (8.1mm vs. 25mm) and the length of the reduced section
525 (16mm vs. 33mm). This reduced size and aspect ratio might lead to a non
526 uniform strain field within the area of measurement (Lin et al., 2024).

527 4.3. *Variability on geometrical measurements*

528 Despite efforts to reach a consensus on the testing methodology, the sec-
 529 ond round did not display significant improvement in the quantities of inter-
 530 est. One metric that was however drastically improved was the variation on
 531 the sample width measurement (see Figures 8b and 10). This makes sense
 532 given the fact that this dimension was not specified in the first round, whereas
 533 all participants used the same cutting tool in the second round. Thickness
 534 measurement variability was also improved (as can be observed from the re-
 535 duced coefficient of variation of the thickness measurement in Figure 8b),
 536 but not to the same extent as the width measurement. This may be ex-
 537 plained by a greater variation in the thickness compared to the width of the
 538 samples, as the samples were prepared using the cutting tool. Of course,
 539 release of residual stresses may still have led to some width variation in the
 540 samples. Secondly, the samples are approximately twice as wide as they are
 541 thick, reducing the signal to noise ratio for the latter measurement. The
 542 variability on the thickness measurement is indeed significantly lower for the
 543 synthetic samples in both rounds, although one can notice an outlier group
 544 for the measurements on synthetic 1A samples. Apparently, this group had
 545 used an optical laser measurement method, which suffered from laser light
 546 penetration for this sample type, yielding a systematic error in their results
 547 (O’Leary et al., 2013).

548 4.4. *Variability on stress-stretch curves and tangent modulus*

549 Figures 5 and 8a illustrate how the improvements on the dimension mea-
 550 surements did not propagate to the resulting stress-stretch curves or tangent
 551 moduli. For low stretch values, coefficient of variation levels are even slightly
 552 increased, and there is a decrease in research groups reaching the threshold
 553 value of coefficient of variation for the longitudinal stress-strain behaviour of
 554 the aorta. For the higher strain level, we do observe a reduction in coefficient
 555 of variation levels and a higher number of participants reaching the thresh-
 556 old, but the improvement is far lower than initially anticipated.

557
 558 An explanation for this lack of improvement in terms of tangent modulus
 559 variability could lie in the following: due to the standardization of the zero
 560 strain state in the consensus protocol, the origin of the curves is more ho-
 561 mogeneous and the curves have therefore on average shifted towards the left,
 562 leading to higher stresses for a certain stretch level (see Figures 6 and 7). As

563 a consequence, the values of the tangent moduli obtained in round 2 are sig-
564 nificantly higher than in round 1, increasing the amplitude of their variability.

565
566 In all cases, we can observe less variability for synthetic samples than
567 for aortic samples. In round 2, all participants are below the coefficient
568 of variation threshold value for the stress-stretch behaviour and the overall
569 coefficient of variation in tangent modulus is lower for both stretch levels.
570 The resulting tangent modulus is also significantly higher for the synthetic
571 material used in round 2 compared to those of round 1. Indeed, a different
572 material was used, which at least at low stretch levels, matched the average
573 tangent modulus of the aortic material more closely.

574 4.5. *Remaining sources of variability*

575 The estimated methodological uncertainty on the stress measurement
576 (equation 1) is significantly smaller than the observed one, even more so
577 for round 2 (see Figure 4b). This means that much of the variability is still
578 due to other factors, a number of which are discussed below.

- 579 • **Stretch measurement uncertainty** - Either marker tracking or DIC
580 was performed in round 2. The reliability of the former can be affected
581 by marker placement accuracy and tracking resolution, which was not
582 harmonized between the participants. Also for DIC, its reliability de-
583 pends on the quality of the speckle pattern, the resolution of the imag-
584 ing system, as well as on software analysis settings, as was investigated
585 at length in a series of ‘DIC challenges’ (Reu et al., 2018, 2022). Nev-
586 ertheless, these DIC challenges were not specifically aimed at biological
587 tissue applications, which pose particular challenges. For example, im-
588 mersion of the sample in a water bath possibly causes optical distortion,
589 speckle pattern application should ensure proper adherence to the tis-
590 sue, not cause or require dehydration of the tissue, and last through
591 large deformations (Lionello et al., 2014). The sample shape, shorter
592 than the ASTM recommendations for synthetic materials, can induce
593 an inhomogeneous deformation pattern, the degree of which should be
594 investigated (Lin et al., 2024). These inhomogeneities should be dealt
595 with, either through a sample-size specific correction factor, or by ap-
596 plying DIC rather than marker tracking where possible.
- 597 • **Zero-strain state** - The samples were considered to be in their zero
598 strain state at the start of the test, specifically at 0.03 MPa for biolog-

ical samples and 0.06 MPa for synthetic ones. This stress-based definition clearly induces error as well as uncertainty proportional to the force and surface area measurement uncertainty. Moreover, it is generally known that biological tissues exhibit residual strains (Vaishnav & Vossoughi, 1983), further confounding the definition of the zero-strain state.

- **Adherence to the testing protocol in round 2** - Despite the provided instructions, we observed variations in how different groups adhered to the testing protocol. For instance, some groups removed the connective tissue of the adventitial layer while others did not. The adventitial layer is known to be collagen-rich and therefore contribute significantly to the nonlinearity of the stress-stretch curve. Additionally, there were discrepancies in the applied preconditioning cycles and in the definition of the state to be reached before starting the final ramp loading. This might be attributed to misunderstanding of the instructions, but also due to the incompatibility of certain testing machine controllers w.r.t. the desired protocol. These deviations are documented in detail in the supplementary material (see ProtocolDeviationsRound2.pdf). Note also that the hypothesis was made that each participant used calibrated sensors and testing machines, for example according to ASTM E4 standards, but this was not explicitly verified.
- **Intra-group variability** - Some teams exhibited greater intra-group variability in round 2 compared to round 1. This could be attributed to the fact that they were following a testing protocol different from their usual practices. For example, managing samples in an immersed test environment can be more challenging and may introduce additional variability, especially when the user is still going through a learning curve.
- **Sample misalignment** - For highly anisotropic samples, ensuring that samples are cut precisely along the circumferential or longitudinal direction is crucial for consistent results. Misalignment can lead to variations in the mechanical properties measured, as the orientation of the fibers in the tissue can significantly affect its behavior. Careful attention to cutting and aligning the samples is necessary to minimize this source of variability, but can never be fully excluded.

- 634 • **Variability of the synthetic materials** - The synthetic samples were
635 considered as isotropic and homogeneous, although it is known that
636 the manufacturing process may introduce a flow-related anisotropy and
637 other imperfections. Hence, even for these materials, it is important
638 to differentiate methodological variability from production-related vari-
639 ability.
- 640 • **Participant preferences** - Certain aspects of the testing protocol
641 were left to the discretion of the participants, such as the orientation
642 of the testing device (vertical vs horizontal), the type of clamps used
643 (pneumatic or manual). Narrowing this down would have led to a much
644 smaller number of participants, which is why this trade-off was made.
645 Scholze et al. (2020) have proposed an easy-to-fabricate clamping sys-
646 tem for uniaxial tensile testing, that could be investigated for use in
647 future testing rounds.
- 648 • **Number of test samples** - 6 samples were tested per condition per
649 research group, each time coming from two different animals. This
650 number is based on previous experiments combined with the practical
651 limitations on the number of aortas that could be collected from the
652 slaughterhouse. Ideally, a dedicated power analysis of the current re-
653 sults should be performed to reveal the required number of tests for
654 future testing rounds for this specific test and sample type.

655 4.6. *Future work*

656 This pilot campaign serves as an exploration of the current landscape,
657 highlighting the existing variability in the mechanical testing to characterize
658 biological tissues. As such, it marks only the start of a much needed effort
659 to further quantify and minimize this variability, and the following areas of
660 future work have been identified.

- 661 • Further statistical analysis of the data should be performed to help
662 identify specific sources of variability. Also, even more detailed analysis
663 of each of the participants' submissions might uncover further devia-
664 tions from the protocol.
- 665 • Each of the above listed remaining sources of variability accounts for
666 just a portion of the overall variability. For example, when observing
667 the degree of anisotropy of the biological tissues tested here, one could

668 argue that the difference between the circumferential and longitudinal
669 behavior is much smaller than the overall variability, such that sample
670 misalignment isn't such a big issue. Indeed, the challenge is not only to
671 identify sources of uncertainty, but also to rank each of these sources in
672 terms of their relative importance. Once a clear overview and ranking
673 of these sources of variability is obtained, further fine-tuning of the
674 consensus protocol can be performed after which a new test round can
675 be launched.

- 676 • To assess the impact of operator variability, it would be beneficial to
677 have the same operator perform the test on different set-ups. In other
678 words, rather than sending samples around, one could exchange re-
679 searchers. Organising training sessions or workshops for all participants
680 could help improve consistency in following the testing protocols.
- 681 • The shape and amplitude of the stress-stretch curves of all the syn-
682 thetic materials deviate strongly from those of the curves for biological
683 samples. This severely limits the relevance of these synthetic samples
684 in terms of protocol verification capacity. Indeed, these synthetic sam-
685 ples do not exhibit the characteristic nonlinear stiffening behaviour,
686 and this absence of a so-called 'toe-region' significantly reduces the
687 sensitivity to slight differences in strain. Therefore, for future testing
688 rounds, we need synthetic materials that do exhibit this nonlinear stiff-
689 ening behaviour. Possible candidates are PVA hydrogels (Millon et al.,
690 2006), fabric-reinforced polymers (Zhalmuratova et al., 2019) or melt
691 electrowritten scaffolds (Mirani et al., 2024). A thorough examination
692 of these materials is required to ensure maximal resemblance to aortic
693 tissue with minimal variability due to the production process, storage,
694 etc.
- 695 • To generalize our findings it is necessary to expand the study to include
696 other types of tests and tissues. This can help to identify whether the
697 observed variability is specific to certain tests or tissues, or if it is a
698 more general issue.

699 This first C⁴Bio campaign serves as a wake-up call that current method-
700 ological variability is too large to enable reliable comparison of results be-
701 tween research groups. This undermines the credibility of resulting material

702 properties reported in literature, and their consequent use in *in silico* simu-
703 lations. This study has made it clear that proper uncertainty quantification
704 is essential, and that through community effort we should aim to further
705 increase the quality of our mechanical testing methods and reduce the un-
706 certainty to a workable level.

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731 Appendix A. Supplementary material

732 All the supplementary material can be found in a repository on Zenodo
733 through the following link.

734 <https://zenodo.org/uploads/14179432?token=eyJhbGciOiJIUzUxMiJ9.eyJpZCI6IjRiMzM5MGlyLTNlYTItNGY4NS04ZTliLWZhMjA2ZTMzN2QyNiIsImRhdGEiOnt9LCJyYW5kb20iOiJjMGZmZDUyZWVlZDIzYzc3MTYwNjAyMmRlNjZhYzAwNyJ9.PG4xkyAfg2JTQuDuZfFh341z2F2nWUedUd9D33ey0aP1AD4s4XTtyyJpmtbftX14oKjP4kSFhu8GZC4ZiJhRfA>
735
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738

739 *Note to reviewers: This link will be shortened in the final submission,*
740 *once the repository is published.*

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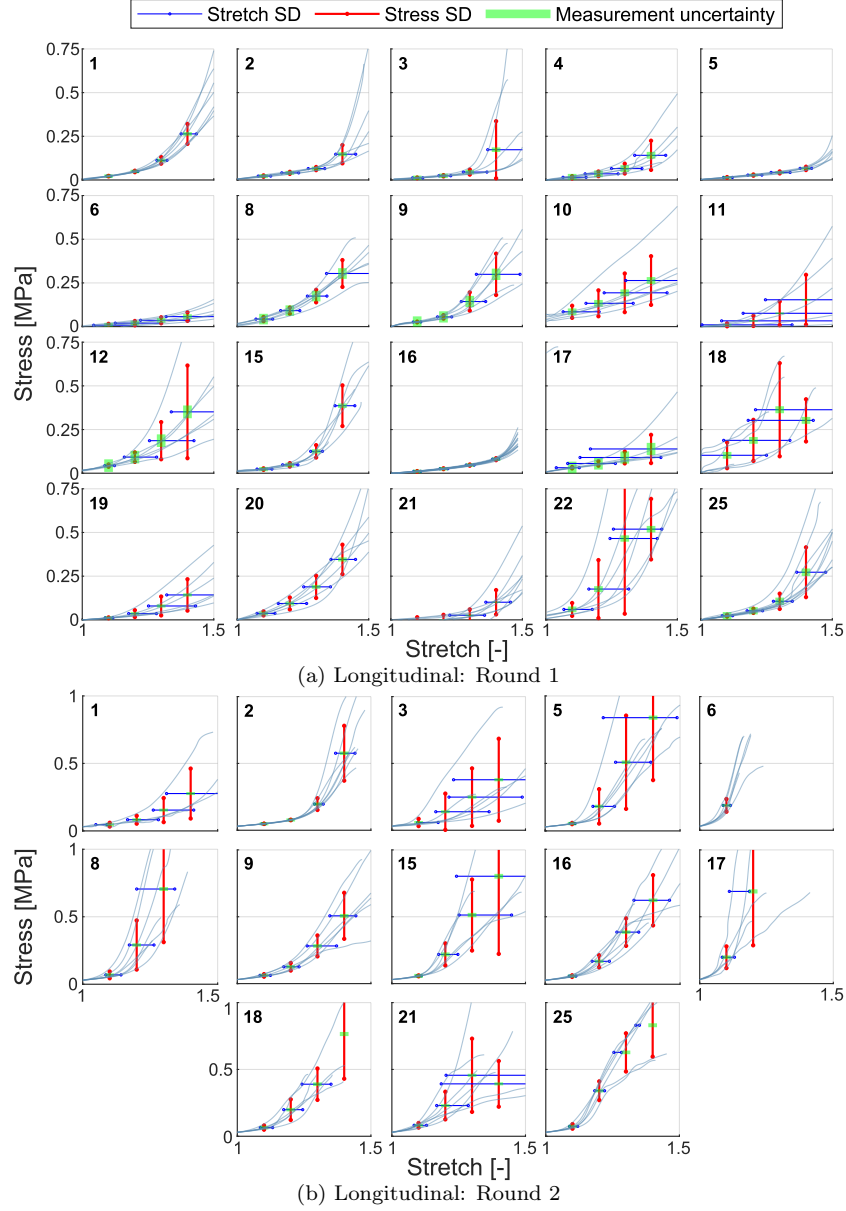


Figure 4: Engineering stress (in MPa) - stretch curves of longitudinal specimen types obtained in (a) round 1 and (b) round 2, grouped per participating research group indicated in the top left corner. Vertical lines represent the standard deviation (SD) of stress values at stretch levels of 1.1, 1.2, 1.3 and 1.4. Horizontal lines indicate the SD of stretch values at the mean stress value corresponding to these stretch levels. Measurement uncertainty on the mean stress value is shown at each stretch level, accounting for a 0.1 N load cell uncertainty and 0.02 mm uncertainty in specimen thickness and width measurements.

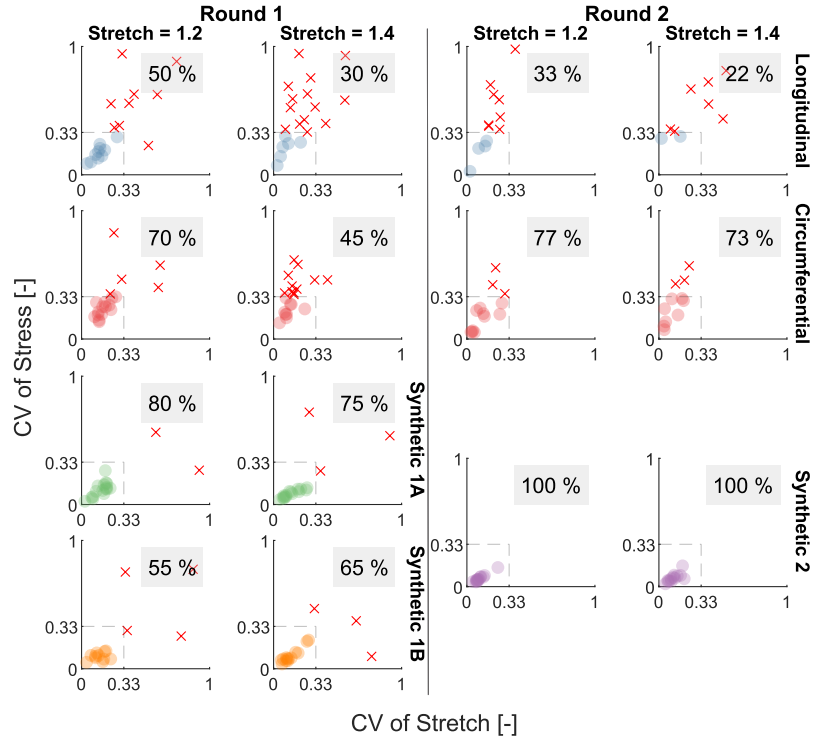


Figure 5: Coefficient of variation (CV) of engineering stress versus CV of stretch at two specific stretch levels, 1.2 and 1.4. CV values are calculated per research group. Research groups for which both CV values fall within the threshold of 0.33 are represented by dots, while those with either CV value exceeding this threshold are marked with ‘x’ markers. The percentage of research groups whose CV values remain within the threshold is indicated on each plot. The data are organised by specimen type and separated by testing round.

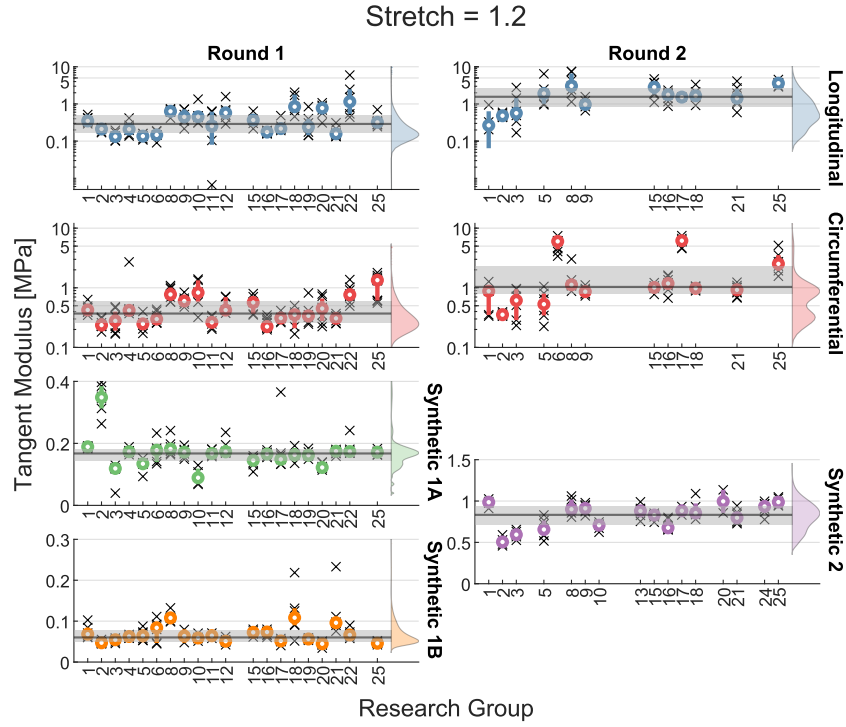


Figure 6: Tangent modulus of the stress-stretch curve at 1.2 stretch per research group, organised by testing round and specimen type. Individual moduli are represented by ‘x’ markers, while the mean for each research group is indicated by a dot and first- to third quantile (Q1-Q3) values are shown as bars. The overall mean and Q1-Q3 values, aggregated across all research groups, are depicted by a line and shaded band, respectively. A density plot is included to represent the distribution of all moduli for each specimen type and testing round, providing insights into the data distribution and variability.

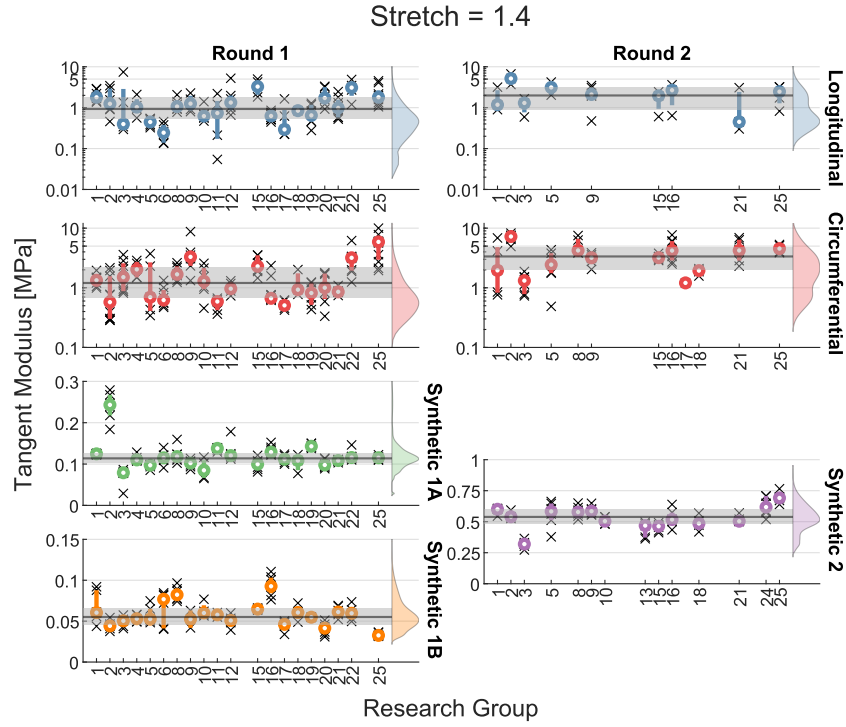


Figure 7: Tangent modulus of the stress-stretch curve at 1.4 stretch per research group, organised by testing round and specimen-type. Individual moduli are represented by ‘x’ markers, while the mean for each research group is indicated by a dot and first- to third quantile (Q1-Q3) values are shown as bars. The overall mean and Q1-Q3 values, aggregated across all research groups, are depicted by a line and shaded band, respectively. A density plot is included to represent the distribution of all moduli for each specimen type and testing round, providing insights into the data distribution and variability.

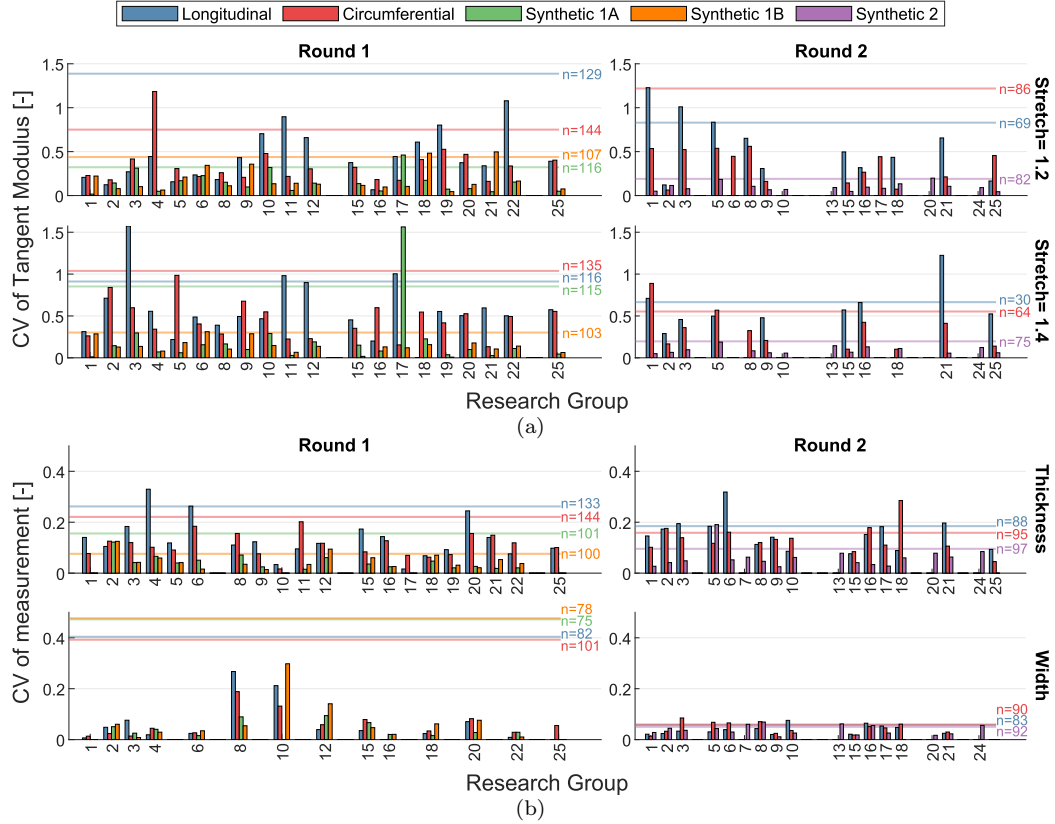


Figure 8: Bar plots showing the coefficient of variation (CV) of (a) the tangent modulus of the stress-stretch curve at 1.2 and 1.4 stretch values, and (b) thickness and width measurements. CV values are represented for each research group across different specimen types, separated by testing round. The overall CV value, calculated by aggregating measurements across all research groups for each specimen type, is represented by a line. The corresponding sample size is annotated adjacent to the line.

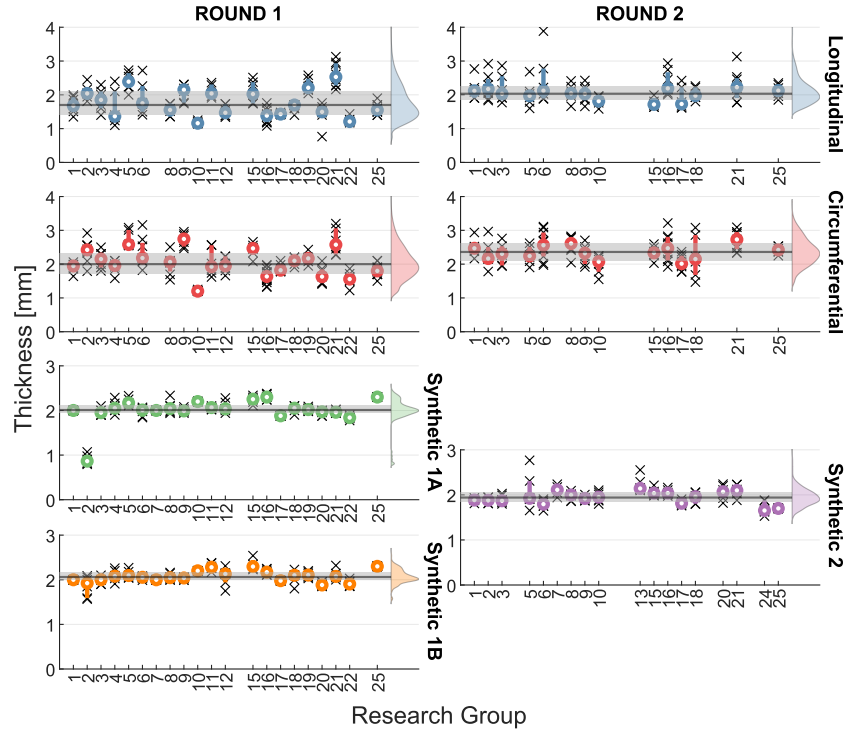


Figure 9: Specimen thickness measurements per research group, organised by testing round and specimen type. Individual measurements are represented by ‘x’ markers, while the mean for each research group is indicated by a dot and first- to third-quantile (Q1-Q3) values are shown as bars. The overall mean and Q1-Q3 values, aggregated across all research groups, are depicted by a line and shaded band, respectively. A density plot is included to represent the distribution of all measurements for each specimen type and testing round, providing insights into the data distribution and variability.

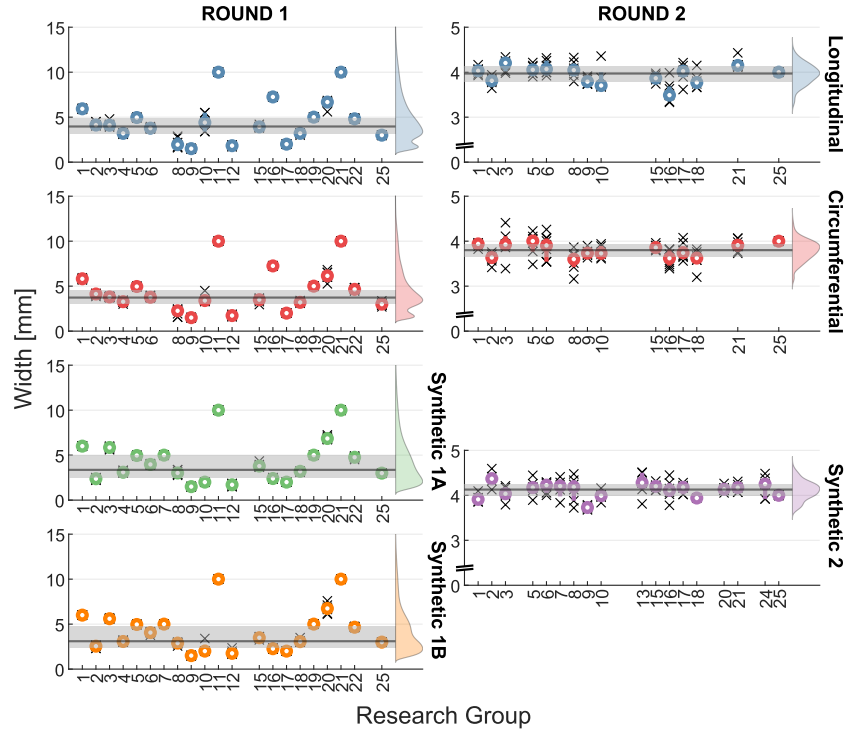


Figure 10: Specimen width measurements per research group, organised by testing round and specimen type. Individual measurements are represented by 'x' markers, while the mean for each research group is indicated by a dot and first- to third-quantile (Q1-Q3) values are shown as bars. The overall mean and Q1-Q3 values, aggregated across all research groups, are depicted by a line and shaded band, respectively. A density plot is included to represent the distribution of all measurements for each specimen type and testing round, providing insights into the data distribution and variability.