

1 A novel method for comparison of arterial remodeling in hypertension:
2 quantification of arterial trees and recognition of remodeling patterns on
3 histological sections

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15 **Abstract**

16 Remodeling of spatially heterogeneous arterial trees is routinely quantified on tissue sections by
17 averaging linear dimensions, with lack of comparison between different organs and models. The
18 impact of experimental models or hypertension treatment modalities on organ-specific vascular
19 remodeling remains undefined. A wide variety of arterial remodeling types has been
20 demonstrated for hypertensive models, which include differences across organs. The purpose of
21 this study was to reassess methods for measurement of arterial remodeling and to establish a

22 morphometric algorithm for standard and comparable quantification of vascular remodeling in
23 hypertension in different vascular beds. We performed a novel and comprehensive morphometric
24 analysis of terminal arteries in the brain, heart, lung, liver, kidney, spleen, stomach, intestine,
25 skin, skeletal muscle, and adrenal glands of control and Goldblatt hypertensive rats on routinely
26 processed tissue sections. Mean dimensions were highly variable but grouping them into
27 sequential 5 μ m intervals permitted creation of reliable linear regression equations and complex
28 profiles. Averaged arterial dimensions demonstrated seven remodeling patterns that were distinct
29 from conventional inward-outward and hypertrophic-eutrophic definitions. Numerical modeling
30 predicted at least twenty variants of arterial spatial conformations. Recognition of remodeling
31 variants was not possible using averaged dimensions, their ratios, or the remodeling and growth
32 index. To distinguish remodeling patterns, a three-dimensional modeling was established and
33 tested. The proposed algorithm permits quantitative analysis of arterial remodeling in different
34 organs and may be applicable for comparative studies between animal hypertensive models and
35 in human hypertension. Arterial wall tapering is the most important factor to consider in arterial
36 morphometry, while perfusion fixation with vessel relaxation is not necessary. Terminal arteries
37 in organs undergo the same remodeling pattern in Goldblatt rats, except for organs with
38 hemodynamics affected by the arterial clip. The existing remodeling nomenclature should be
39 replaced by a numerical classification applicable to any type of arterial remodeling.
40

41 **Author summary**

42 Arterial hypertension effects modern nations and is characterised by systemic hypertensive
43 angiopathy that affects all organs. Arterial remodeling is a main factor to be analyzed in animal
44 models and human. Despite abundant data, there is a significant lack of comparative analysis on

45 arterial remodeling. The data from the present study have established a novel methodological
46 approach to assess and compare arterial remodeling in hypertension. We have developed an
47 effective algorithm for morphometry of intra-organ arteries to standardize remodeling
48 assessment and allow comparisons between different hypertensive models, organs and species.
49 Our study opens the possibility to assess remodeling using conventional widely used histological
50 tissue sections with no need for special perfusion-fixation. The method will elucidate the
51 improvement and development of animal models of hypertension, and enhance the assessment of
52 experimental therapeutic modalities.

53 **Introduction**

54 The importance of understanding the pathogenesis of hypertension is undisputed, and despite
55 recent decreases in mortality due to heart disease and stroke, the burden of disease remains high.
56 Based on data from 2018, 33% of adults in North America have hypertension, but only 53% of
57 those with documented hypertension have their condition controlled to target levels[1]. An
58 important feature of hypertensive angiopathy is vascular remodeling: a complex structural and
59 spatial modification in small arteries that is of crucial functional consequence since it alters
60 peripheral resistance and impairs contractility[2],[3],[4],[5].

61 The features of arterial remodeling in hypertension have been extensively
62 studied[3],[4],[6],[7]. To date however, there are no data demonstrating how arterial remodeling
63 in spontaneously hypertensive rats (SHR) is distinct from hypertension due to angiotensin II or
64 deoxycorticosterone acetate infusion, inhibition of nitric oxide synthesis, or Goldblatt's model.
65 The utility of animal models must be based on their resemblance to human pathology, yet
66 commonly used measures are unable to quantify how arterial remodeling in experimental
67 animals corresponds to arterial remodeling in humans. Thus, the well-established clinical

68 concept of target organ damage in hypertension[8] has yet to be supported by data indicating that
69 arterial remodeling is more extensive in the kidney or heart, for example, compared to the skin,
70 liver, or other organs.

71 In the majority of studies complex three-dimensional (3D) intra-organ arterial trees have
72 been quantified by simple measures first described more than 80 years ago[9], consisting of the
73 wall-to-lumen ratio (WLR), and/or the mean values for a variety of arterial dimensions, e.g.,
74 external diameter (ED), internal diameter (ID), wall thickness (WTh), external perimeter (EP),
75 internal perimeter (IP), media cross sectional area (MCSA), lumen cross sectional area (LCSA),
76 total cross sectional area (TCSA) and internal radius (IR). These measures are highly variable
77 and inconsistent, and do not permit comparison of remodeling between different organs,
78 hypertension models or species. Presumably, that quantification of tapered 3D intra-organ
79 arterial trees on tissue sections is oversimplified, and conventional averaging of dimensions in
80 vessel morphometry led to incomparable stochastic results. Modern micro-computed tomography
81 achieves 3D images of peripheral vessels with high resolution: $\sim 2.5 \mu\text{m}$ voxel size. However, in
82 this technique, detailed microstructure cannot be described since only the contrast-filled lumen is
83 visualised. While this is useful for angiogenesis studies, analysis of hypertensive remodeling is
84 limited[10].

85 Arterial wall remodeling patterns have been classified as hypertrophic, hypotrophic or
86 eutrophic, associated with inward narrowing or outward widening of the lumen[6],[11]. To date,
87 however, there is significant discrepancy with regards to the type of remodeling that develops in
88 terminal arteries (TAs) of different organs in various hypertensive models. In SHR for example,
89 renal arterioles may demonstrate remodeling characterized as outward
90 hypertrophic[12],[13],[14], inward hypertrophic[15], solely hypertrophic[13] or no

91 change[14],[16]. Some authors conclude that renal arteries $> 60 \mu\text{m}$ do not develop
92 remodeling[17], while others suggest an absence of remodeling for smaller arteries $< 60 \mu\text{m}$ [16].
93 Mesenteric arteries may show inward hypertrophy or no change[12],[18]. Similar apparent
94 contradictions appear in other hypertensive models[19],[20],[21],[22].

95 Such discrepancies could arise since arterial remodeling has previously been classified
96 using empirical drawings, without precise quantitative analysis[11]. Since it was introduced
97 twenty years ago, the classification has been extensively reviewed[5],[23],[24],[25] but has not
98 been challenged with quantitative methods. Furthermore, frequently used parameters, such as
99 WLR, remodeling index (RI), and growth index (GI) have not been rigorously tested as markers
100 of remodeling. We therefore set out to i) study the classification of arterial remodeling patterns in
101 hypertension, using mathematical methods, and ii) elucidate how arterial remodeling differs
102 across a variety of organs. An algorithm for arterial remodeling assessment was developed, and
103 we then determined if it could distinguish variants of remodeling in different organs within one
104 model of hypertension in Goldblatt one-kidney one-clip (1K1C) rats. We hypothesized that, if
105 conventional averaging was avoided, all organs would show the same arterial remodeling
106 pattern. We also hypothesized distinct remodeling patterns for the adrenal gland and kidney,
107 where the 1K1C model creates particular hemodynamic conditions. In this model, arteries within
108 the adrenal glands experience the effects of an activated renin-angiotensin system (RAS), similar
109 to other organs, but also experience enhanced flow due to diversion of blood from the main renal
110 artery as a result of distal stenosis or ligation, similar to hemodynamic models of
111 overflow[26],[27]. The remaining kidney also experiences the vasoconstrictive effects of an
112 activated RAS, but under low blood flow due to the clipped renal artery[28],[29], that
113 corresponds to low blood flow models[26],[30].

114 We also determined if examination of random tissue sections can provide useful
115 information in studying hypertension, compared to use of arterial myography[3],[31]. In this
116 regard, the vast majority of data from *in vitro* myography experiments has been generated from
117 dissected mesenteric arteries, and may not be applicable to other organs. Finally, studies have
118 recommended that arterial morphometry should only be performed on perfusion-fixed organs
119 with pharmacologically relaxed vessels[31],[32]. We therefore examined if routine immersion
120 fixation is appropriate for arterial remodeling analysis to elucidate comparisons between animal
121 and human samples, since for the latter immediate relaxation and perfusion fixation are not
122 practical in general.

123

124 **Results and Discussion**

125 **Part I. Quantification of arterial trees on histological sections**

126

127 **Means of linear sizes and ratios are not applicable for arterial morphometry on
128 tissue sections**

129 While the parenchyma of different organs is well represented on histological sections, arteries
130 are an exception. They appear as circles and irregular strips with a wide range of sizes and
131 shapes, which are difficult to quantify[33]. To date, there are two approaches to arterial
132 morphometry. *In vitro* myography uses similar vascular segments to minimize variability and
133 allow comparison of averaged dimensions or their ratios among different models, or different
134 arteries in the same model[34],[35]. However, myography data are restricted to only a few areas
135 that are suitable for sampling - mainly the mesentery and aorta. Second, quantification of intra-
136 organ arteries is based on casual measurements of arterial segments on tissue sections. However,

137 the majority of studies simply average the sizes and ratios, as for myography data, resulting in a
138 high degree of variability and difficulty in comparing organs. Measured vessels are defined with
139 uncertain ranges. For example, for the kidney: ‘arteries and arterioles’[36]; ‘intrarenal’[37],[38];
140 ‘proximal interlobular’[13],[39]; ‘afferent arterioles’[40],[41]; ‘vessels adjacent to glomeruli in
141 the outer cortex’[42] or ‘at the same level proximal one-third from corticomedullary
142 junction’[43]. Similar ‘interlobular arteries’ could vary significantly in ID (~100 μm [44], ~40
143 μm [13], ~25-50 μm [45], ~30-250 μm [22]).

144 TAs from multiple organs in normal rats were first analyzed using that conventional
145 approach with measurements of mean ED, ID, WTh and WLR (**Table 1**). All values were
146 obtained within an arterial ED range of 10-50 μm , since larger or smaller arteries were difficult
147 to find in sufficient numbers in tissue samples. TAs in each organ were characterized by distinct
148 average linear dimensions and WLRs. For instance, although bronchial and skeletal muscle
149 arteries demonstrated similar WLRs, their EDs, IDs and WThs differed ($P < 0.01$). The WLRs
150 for brain arteries were lower but IDs larger than in liver arteries ($P < 0.001$), although EDs and
151 WThs were similar between these two organs. Importantly, coefficient of variation of arterial
152 dimensions were relatively high, varying between 40-70%, indicating their significant
153 diversity[46]. Accordingly, individual histograms for parameter distribution in each organ were
154 prepared. Histograms for ED, ID, and WTh demonstrated considerable variability, without a
155 normal Gaussian distribution (**Fig 1**). Indeed, as shown in **S1 Table**, the extent of variability for
156 ED, ID and WLR measures did not pass conventional statistical tests for normality.

157

158

159

160 **Table 1. Averaged linear dimensions and wall-to-lumen ratio for terminal arteries with ED**
161 **of 10 – 50 µm in different organs.**

Organ	ED, µm	ID, µm	WTh, µm	WLR, %
Liver (N=63)	21.6±1.4	8.8±0.8**	6.4±0.4	98.6±7.2**
CV	54%	69%	48%	58%
Adrenal (N=61)	22.0±1.1	8.7±0.5	6.6±0.4	81.9±5.1
CV	40%	41%	52%	48%
Kidney (N=148)	32.7±0.6	13.3±0.4	9.7±0.2	78.9±2.2
CV	24%	33%	24%	35%
Skin (N=225)	20.1±0.6	8.5±0.3	5.8±0.2	75.3±1.9
CV	48%	52%	52%	37%
Skeletal muscle (N=98)	19.7±0.7*	8.4±0.3*	5.6±0.2*	70.8±3.1
CV	37%	37%	45%	43%
Bronchial arteries (N=50)	37.1±1.3	18.4±1.2	9.4±0.4	68.7±7.5
CV	24%	48%	32%	77%
Heart (N=115)	26.6±0.9	12.7±0.6	6.9±0.3	63.6±3.0
CV	40%	52%	42%	50%
Spleen (N=203)	17.7±0.5	8.3±0.3	4.7±0.1	61.6±1.3
CV	42%	51%	38%	30%
Small Intestine (N=55)	18.9±1.	9.5±0.8	4.7±0.2	57.2±3.0
CV	42%	65%	38%	39%
Stomach (N=90)	24.9±1.0	12.7±0.6	6.1±0.3	53.5±3.0
CV	41%	46%	54%	54%
Brain (N=144)	23.4±0.6	12.7±0.4	5.4±0.2	45.6±1.7
CV	32%	40%	35%	46%
Pulmonary arteries (N=66)	29.2±1.3	20.9±1.0	4.1±0.2	21.9±1.1
CV	35%	41%	34%	42%

162 Data are mean \pm SEM. CV - coefficient of variation; N - number of measured arteries; *P<0.01
163 vs the bronchial arteries; **P<0.001 vs the brain. Organs are displayed in order of descending
164 WLR values.

165

166 **Fig 1. Detailed statistical analysis of primary data.** Histograms for ED, ID, and WTh in the
167 kidney, heart and pulmonary arteries, used to calculate statistics in Table 1, demonstrated
168 irregular frequencies and profound asymmetry.

169

170 Our detailed analysis (**Table 1, Fig 1, S1 Table**) indicates that mean values have little statistical
171 value with regards to such measurements[46], since even a small range of EDs (10-50 μ m) for
172 TAs in all organs was associated with an abnormal distribution and high variability.
173 Furthermore, increasing numbers of measurements to normalize distribution and reduce
174 variability are not effective for tapering branching objects[33].

175 While WLR is considered independent of diameters and not affected by the size of
176 arteries studied[3],[9],[41], this matter has not been rigorously substantiated in the
177 literature[4],[5],[25],[32],[47]. Furthermore, we found as many as sixteen different formulae to
178 calculate WLR. Some studies employed the diameter or radius: WTh/ID[20],[44],[48]; ED -
179 ID/2ID[49]; ED/ID[50]; WTh/IR[12][17]; a media index WTh/IR with IR specifically defined as
180 the distance from the center of the arterial lumen to the middle point of the media[51];
181 2WTh/ED[52],[53]; 2WTh/ID[54]; WTh/ED[55]; or ID/ED[56]. Others have measured WLR
182 from the perimeters as EP/IP[40],[41],[42], or preferred to use the area:
183 MCSA/LCSA[15],[16],[45]; MCSA/TCSA[37]; LCSA/MCSA[36]; LCSA/TCSA[57]. The
184 WLR has also been interpreted as the ratio of the area to perimeter MCSA/IP[22], or renamed
185 RI[58]. Rarely, point counting has been used to estimate WLR as the average ratio of the volume
186 density of the walls to the volume density of the lumens[16],[59]. However, many studies
187 indicate that WLR is not constant but declines 2-10-fold in arteries of ED 20-500 μ m in the

188 human and rat kidney[51],[60]; human and rat brain[61],[62]; and human and dog liver[48],[63].

189 Our data indicate that WLR is not constant even within a small range of EDs (**Fig 2A**).

190

191 **Fig 2. Detailed statistical analysis of primary data.** (A) WLR scatterplots were irregular with
192 low r^2 (solid lines), and wide 95% prediction bands (dashed lines). Corresponding statistics are
193 shown in **Table 1**. (B) Scatterplots of primary data displayed certain incremental change in
194 arterial dimensions. Coefficients r^2 were moderate, and the pulmonary arteries had the best
195 fitting value.

196

197 In summary, neither averaged WLR nor the mean values for ED, ID or WTh are
198 applicable to even relatively small intervals of vessel caliber. Thus, any comparison of mean
199 values for these measures is not appropriate[46].

200

201 **Application of short intervals provides precise complex profiles and linear
202 regression equations**

203 For vessels with EDs in the range of 10-50 μm , there was a trend towards dependence of ID and
204 WTh on ED, although variability was substantial (**Fig 2B**). We therefore assessed primary data
205 with a complex profile method, which is widely used in cartography, thermodynamics, and
206 engineering[64],[65]. In this technique, a contour line is drawn as a function of two variables,
207 where the gradient of function is perpendicular to the contour isoline, thus representing more
208 than two dimensions on a two-dimensional (2D) graph. To be applied to measuring arteries, each
209 arterial circle on a histological section is first characterized by three values: ED, ID, and WTh.
210 Then all measured vascular circles were arranged in a row in order of ascending ED, and
211 numbered from 1 to N, where N is the total number of measured arterial rings. Subsequently,

212 each number was plotted on the y-axis, while its corresponding ED, ID and WTh were plotted on
213 a bidirectional x-axis (**Fig 3**).

214

215 **Fig 3. Analysis of primary data with complex profiles.** Primary measurement for the kidney,
216 heart, and pulmonary arteries were arranged in the complex profiles. Axis X – the bidirectional
217 common scale for ED (outer contours), ID (inner contours), WTh (shaded regions); axis Y
218 represents the number of measured arteries (variables) in order of ascending ED. Steps in the
219 outer contours (solid arrows) reflected a minimal division on an eyepiece micrometer. Outliers
220 along the internal contours (dashed arrows) corresponded to wider or narrower ID, compared to
221 neighboring ED values.

222

223 The same primary data, represented in this way, revealed a regular hemodynamic structure.
224 Staircase steps of the external contours were relatively small and regular since a conventional
225 microscope with a 40x objective provided a minimal division value of $\sim 2 \mu\text{m}$ to the eyepiece
226 micrometer, so that each ED was rounded off to $\pm 2 \mu\text{m}$. Spikes of internal contours were larger
227 and quite irregular corresponding to some arterial segments with similar ED but different ID
228 and/or WTh, and it was necessary to understand the source of multiple outliers. Presumably,
229 those outliers represented subsets among arterial segments, branching at different hydrodynamic
230 points. We speculated that arteries with similar ED at proximal regions should have thicker
231 media, i.e. higher wall-to-lumen ratio, in response to larger pressure gradients (**Fig 4A**).

232

233 **Fig 4. The hemodynamic explanation for the appearance of outliers.** (A) According to Fick's
234 law, the capillary pressure in proximal (p1) or distal (p2, p3) regions must be similar, while

235 arterial pressure in a proximal artery P1 is higher than in distal arteries P2 and P3. Consequently,
236 $\Delta(P1 - p1) > \Delta(P3 - p3)$. Arteries at proximal regions should have thicker media, i.e. higher wall-
237 to-lumen ratio, in response to larger pressure gradients. (B) The paired sampling of pulmonary
238 arteries. Forty five arterial pairs with the same ED were identified in the proximal axillary and
239 distal dichotomous branches respectively. For example, A1 (the red bar) and A3 (the green bar),
240 sampled into paraffin blocks, and dimensions were measured on stained sections. Scale bar, 100
241 μm . (C) Five pairs in the table are the examples of the total (N=45) measured pairs of branches
242 with the same ED that could have differences from 50-110% in ID, 100-230% in WTh, and 250-
243 600% in WLR ($P < 0.001$). (D) In the complex profile, those neighboring branches create most
244 outliers, as in **Fig 3**.

245
246 To test this hypothesis, dissected pulmonary arteries were measured by the paired sampling:
247 branches of the same ED were identified in the proximal and distal segments, cut at branching
248 points, and their dimensions were measured (**Fig 4B-D**). To compare only two vessels 5 mm
249 apart would require enormously laborious reconstruction of approximately 2000 serial sections
250 while the method of paired branches morphometry was much more effective. Axillary and
251 regular dichotomous branches with the same ED but smaller ID and thicker WTh would be
252 placed closer in profiles, determined as outliers (**Fig 3**). That explained the presence of numerous
253 outliers as an intrinsic property of arterial branching to respond to variable local hemodynamics,
254 that complies with a conception of constrained random branching for smaller vessels [66].
255 Outliers are not evident in primary data plots (**Fig 2**) to be removed. Moreover, a conventionally
256 recommended increase in number of measurements[46] simply increases the number of outliers.

257 To make outliers more evident, instead of conventional averaging of the data for the entire range
258 of 10-50 μm , the ED was divided into short regular intervals (**Fig 5**).
259

260 **Fig 5. Basic approach to arterial morphometry analysis.** (A) In the conventional approach,
261 statistics of arterial dimensions are calculated for a broad range of calibers. (B) In the proposed
262 algorithm, statistics are calculated for fixed 5 μm intervals for TAs with ED \sim 10-50 μm .
263

264 To avoid averaging, some investigators have measured interval means. However, significant
265 discrepancies in choice of interval steps and caliber ranges render such data as highly variable
266 and difficult for comparative analysis. Some studies considered two intervals as ‘interlobular and
267 arcuate’ with ID \sim 70 and \sim 160 μm [59], or ‘small and intermediate’ as ED \sim 50-100 and 100-500
268 μm [60]. Others found it more appropriate to divide ‘interlobar vessels’ into six intervals of <70 ,
269 70-95, 95-120, and >120 μm [16]. Another variant was to cut the range of 90-220 μm into four
270 equal classes[21]. A choice for cerebral arteries was four intervals of IR <20 , 20-49, 50-79, and
271 >80 μm but for older animals it was only 2 of 80-119 and 120-159 μm [61]. A different
272 subdivision was used for the liver: ≤ 20 , 21-50, 51-124, and > 125 μm [48]. In the lung seven
273 non-regular intervals of 0-50, 50-100, 100-200, 500-1000, and >1000 μm were counted[53].
274 Thus the advantage of using intervals could be compromised by subjective choices of interval
275 size that neither improves standardization nor comparison.

276 We tested different intervals, and found that the interval of 5 μm was optimal for all
277 organs. That interval corresponds to adding one layer of vascular smooth muscle cells, which has
278 a normal thickness of 5-7 μm [63],[67]. Indeed, the relation between WTh and the number of
279 VSMC-layers can be represented by a linear regression line ($r = 0.88$)[63].

280 When the complex profiles had been drawn not from every measured value, but from the
281 mean interval values, they demonstrated organ-specific gradients in tapering (**Fig 6A, S1 Fig**)
282 allowing comparison of arterial remodeling between organs. Furthermore, in contrast to data
283 depicted in **Fig 2B**, if outliers had been removed, the means of short intervals revealed very tight
284 linear regression between both lumen and wall thickness and vessel caliber in kidney, heart and
285 pulmonary vessels, and in other organs as well (**Fig 6B, S2 Fig**). All acquired equations
286 demonstrated goodness of fit coefficients r^2 between 0.8 - 0.9 ($P < 0.0001$), with few exceptions
287 for WTh in the brain and bronchial arteries (**S2 Table**).

288

289 **Fig 6. Application of the short fixed intervals significantly improved primary data analysis.**
290 (A) Complex profiles were built from the accumulated frequencies (axis Y) of the means of ED,
291 ID, and WTh for 5- μ m ED intervals. In contrast to scatterplots in **Fig 2B**, terminal arteries
292 exhibited distinctive tapering patterns in organs. Complementary graphs are in **S1 Fig**. (B) Mean
293 values of ID and WTh for the regular 5- μ m ED intervals revealed robust linear regression
294 equations ($P < 0.001$). Complementary graphs are in **S2 Fig**. Points on the lines are mean \pm SD for
295 5- μ m ED intervals.

296

297 To decide whether linear regression equations for TAs were organ-specific, their slopes and
298 intercepts were compared. Equations for pulmonary, bronchial, adrenal, stomach and skeletal
299 muscle TAs were distinct ($P < 0.0001$). Heart and spleen TAs had similar equations, as did brain
300 and intestine, and kidney, liver, and skin (**S3 Fig**).

301 Only a few attempts to apply linear regression to arterial morphometry have been
302 published[18],[22],[51],[63]. Those studies demonstrated that the range of EDs from 10-100 μ m

303 fits linear regression, while wider intervals of ED 100-1000 μm follow an exponential function.
304 We found the range of EDs from 10-50 μm was the most practical to analyze. That caliber is not
305 only the most frequently observed, but also best addresses the functional consequence of
306 remodeling, being responsible for maximum values in peripheral resistance[3],[4],[5].

307 The same calculation, based on means of short intervals, was applied to compute WLR,
308 although the ratio continued to demonstrate high variability (**Fig 7A**), presumably due to the
309 interaction of standard deviation in the ratio of two dependent values[46]. It is evident,
310 comparing different organs, that irregular graphs of mean interval WLRs are difficult to
311 approximate correctly (**Fig 7B**). Therefore we calculated interval WLR from verified linear
312 regression equations (**S2 Table**). The results demonstrated clearly that WLR is not constant, with
313 evidence for nonlinear increases or decreases (**Fig 7C**).

314

315 **Fig 7. Significant variability of the wall-to-lumen ratio in different organs.** (A) Interval
316 means of WLR still varied significantly, although r^2 improved as compared with **Figure 2A** (P
317 $\approx 0.14-0.02$). The pulmonary arteries had the highest r^2 fitting value (P=0.002). Points on the
318 lines are mean \pm SD for 5- μm ED intervals. (B) Means of WLR for the 5- μm ED intervals were
319 irregular, except in pulmonary arteries. (C) WLRs were counted from the linear regression
320 equations. The ratios were not constant, decreasing or increasing by nonlinear functions.

321

322 **Hemodynamic significance of morphometry data**

323 Reliable linear regressions also validated an assessment of relative resistances (RR) that was
324 calculated using Poiseuille's relationship $\text{RR} = 1/\pi r^4$, assuming constant viscosity and length, and
325 where r is the radius of the vessel lumen, as it was recommended[63],[68],[69]. RRs for 5 μm
326 regular intervals across multiple organs are depicted in **Fig 8 and S2 Fig**, with the highest values

327 in liver, and lowest in pulmonary arteries, rising exponentially in the smaller segments, since
328 blood flow is proportional to the fourth power of the luminal radius.

329

330 **Fig 8. Comparative analysis of the relative resistance in different organs.** (A) RR curves
331 were calculated from the regression equations in **S2 Table**. The liver and spleen had the largest
332 values ($P<0.001$), while the pulmonary and stomach arteries - the lowest ($P<0.001$). (B)
333 Diversions in RRs were better revealed on the logarithmic scale.

334

335 Accordingly, terminal relative resistance (TRR) would estimate the resistance for the smallest
336 vessels of $ED= 10-20 \mu m$ that have the largest accumulated frequency in complex profiles (**Fig**
337 **9A**).

338

339 **Fig 9. Correspondence between morphometric and hemodynamic parameters.** (A) TRR
340 demonstrated similarity in organ-to-organ ratio with vascular resistances acquired with
341 physiological methods: modified from[70] (B); [71](C);[72] (D). Data on (A) are mean $\pm SEM$,
342 on (B-D) – mean values.

343

344 Since the complex profiles reveal the average numbers of arterial branches of defined lumen
345 diameters in each organ, prior to the capillary bed, it would be reasonable to measure terminal
346 capacity (TC) in each organ, as the sum of the internal volume of each segment multiplied by its
347 frequency:

348
$$TC = \sum(\pi * (\text{ID}_i)^2 / 4)i\% \quad (1)$$

349 Where ID_i is the mean ID of each interval, calculated from the linear regression equation; $i\%$ -
350 the frequency of that interval in the complex profile (**Fig 10A**).

351

352 **Fig 10. Correspondence between morphometric and hemodynamic parameters.** (A) TC
353 values were in good agreement with organ blood flow rates, estimated by a physiological
354 method[73] (B). Data on (A) are mean \pm SEM, on (B) – mean values.

355

356 The hemodynamics of each organ are characterized by blood flow rate (BFR), regional vascular
357 resistance (RVR), vascular volume, and percentage of cardiac output received[70],[71],
358 reflecting the uniqueness of a vascular bed. Although each organ possesses a distinctive arterial
359 tree, there are currently no standardized morphometric parameters to quantify those distinctions
360 and correlate them to BFR or RVR[68],[69]. Indeed averaged EDs, IDs, and WLRs are only
361 loosely associated with hemodynamic parameters, although they accompany impaired
362 contraction or relaxation in *in vitro* myography experiments[62]. Our algorithm provides a
363 promising solution to this problem. The use of linear regression equations allows for
364 quantification of TRR, which reveals significant similarity to RVRs obtained by physiological
365 methods (**Fig 9B-D**)[70],[71],[72]. The proposed TC formula also shows good correlation with
366 BFR, determined by microsphere method[73] (**Fig 10**).

367

368 **Larger arteries are the most important source of errors**

369 In TAs with $\text{ED} > 40 \mu\text{m}$, only a small number of measurements per interval could be obtained
370 that has a crucial impact on the precision of regression equations, since even one outlier can alter
371 the result significantly, i.e. the means of such intervals are unreliable[46], and therefore the same
372 data set could be approximated with several equations and RR curves, depending on numbers of

373 counted larger vessels (**Fig 11**). Small deviation of regression lines significantly affect RR
374 according to the formula $R = 1 / \pi r^4$. In our study, minor deviations of 8-12% in the largest IDs
375 caused deviations of 25-90% in the smallest IDs, which in turn, had pronounced effects on the
376 calculated TRR values, since blood flow is proportional to the fourth power of the luminal
377 radius. To avoid such errors, the equations must be verified by comparison of corresponding RR
378 curves (**Fig 11**).

379

380 **Fig 11. The equations must be verified to avoid the effect of fewer data points for larger**
381 **vessels.** (A) Heart. Means of intervals were approximated with two equations to compute RR
382 curves. The first (red formula and lines) accounted for vessels of ED 10-50 μm . The second
383 (blue formula and lines) accounted for all vessels found (ED=10-120 μm), with only 1-3 values
384 per interval for the largest TAs. The equations were highly distinct between each other
385 ($P < 0.001$), and every line and curve had the highest goodness of fit coefficient r^2 and $R^2 \approx 0.98$ -
386 0.99. (B, as boxed area in A) For the smallest arteries, the difference in ID of 1-2 μm (arrows)
387 resulted in a 270% difference in RR curves ($P < 0.0001$). If the RR curve was calculated not from
388 the equation but from means of intervals (green line), it was closer to the red ($P > 0.88$) than to the
389 blue curve ($P < 0.0001$), i.e. the equation for 10-50 μm (red) was correct. (C,D) The same
390 calculation for intestine. The equation for 10-70 μm was correct ($P > 0.90$ vs $P < 0.22$). Points on
391 the lines are mean \pm SD for 5- μm ED intervals, n – number of measurements per interval

392

393 **The method enables comparison between species**

394 Using this approach, we compared TAs between normal rat and mouse. While averaged
395 dimensions demonstrate no substantial differences, the complex profiles, linear regression
396 equations and RR curves revealed specific patterns for each species (**Fig 12**). In rats renal TAs

397 have thicker media and narrower lumens compared to mice, but lower resistance, which is
398 consistent with hemodynamic data[74].

399

400 **Fig 12. The algorithm enables comparison between species.** (A) Scatterplots for ED, ID, and
401 WTh for the rat and mouse. (B) Mean values from scatterplots revealed no difference ($P>0.40$),
402 including WLR. The corresponding complex profiles (C), and regression equations with the
403 relative resistance curves (D) were very distinctive ($P<0.001$) for the rat (dashed line) vs mouse
404 (solid line).

405

406 **Part II: recognition of remodeling patterns in hypertensive rats on histological
407 sections**

408 **Conventional measurements exhibit different remodeling patterns in hypertensive
409 rats**

410 At 60 days, the systolic blood pressure in the sham-operated and hypertensive rats was 115 ± 6
411 mmHg and 217 ± 21 mmHg, respectively ($P<0.001$). Morphometry data from multiple organs
412 were first analyzed using conventional measures of mean ED, ID and WTh for the arterial ED
413 interval of 10-50 μ m. 1K1C hypertension significantly affected TAs in every organ (**Table 2**).

414

415

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417

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420

421 **Table 2. The spectrum of remodeling variants in terminal arteries of ED 10 – 50 μ m.**

Organ	Sham rats			1K1C rats			WLR	MCSA	NC
	ED, μ m	ID, μ m	WTh, μ m	ED, μ m	ID, μ m	WTh, μ m			
Brain	23 \pm 7	13 \pm 4	5 \pm 2	19 \pm 8** \downarrow	7 \pm 4** \downarrow	6 \pm 2 * \uparrow	\uparrow	\downarrow	3-4cI
Kidney	33 \pm 8	13 \pm 4	10 \pm 2	30 \pm 9** \downarrow	10 \pm 4** \downarrow	10 \pm 3 =	\uparrow	\downarrow	3-4b
Heart	27 \pm 9	13 \pm 6	7 \pm 3	27 \pm 10 =	11 \pm 4** \downarrow	8 \pm 3* \uparrow	\uparrow	\uparrow	2-4c
Pulmo- nary	29 \pm 8	21 \pm 8	4 \pm 1	33 \pm 9** \uparrow	22 \pm 6 \leftrightarrow	6 \pm 1** \uparrow	\uparrow	\uparrow	1-5c
Skin	20 \pm 9	8 \pm 4	6 \pm 3	23 \pm 9** \uparrow	8 \pm 3 =	7 \pm 3** \uparrow	\uparrow	\uparrow	1-5c
Skeletal muscle	20 \pm 7	9 \pm 4	5 \pm 3	21 \pm 10 =	8 \pm 3** \downarrow	6 \pm 3 ** \uparrow	\uparrow	\uparrow	2-4c
Bron- chial	37 \pm 9	18 \pm 7	9 \pm 1	32 \pm 10** \downarrow	13 \pm 6** \downarrow	10 \pm 2** \uparrow	\uparrow	\downarrow	3-4cI
Stomach	25 \pm 7	13 \pm 5	6 \pm 3	27 \pm 11* \uparrow	10 \pm 5** \downarrow	8 \pm 3** \uparrow	\uparrow	\uparrow	1-4c
Intestine	19 \pm 8	9 \pm 6	5 \pm 1	24 \pm 10** \uparrow	9 \pm 4 =	7 \pm 3** \uparrow	\uparrow	\uparrow	1-5c
Adrenal	22 \pm 8	9 \pm 3	6 \pm 2	24 \pm 11** \uparrow	14 \pm 5** \uparrow	5 \pm 3* \downarrow	\downarrow	\downarrow	1-6aII
Liver	22 \pm 9	8 \pm 4	6 \pm 2	22 \pm 12 =	7 \pm 4 =	7 \pm 3 =	=	=	2-5b

422 Data are mean \pm SD. * $P<0.05$; ** $P<0.01$ NC, numerical classification. \uparrow \downarrow = indicate increase, decrease or no change respectively vs control values.

424

425 To date, the majority of studies use the widely accepted classification[11] (**Fig 13**). However, the
426 obtained remodeling varieties did not fit into conventional definitions. The classification
427 considers ‘inward’ or ‘outward’ as only simultaneous ED-ID decrease or increase (**Fig 13**). The
428 brain, bronchial and kidney should be named ‘inward hypotrophic’ due to reduced ED, ID and
429 MCSA, but that term does not mark increased or stable WTh. The classification has no ‘inward-
430 outward hypertrophic’ type that appeared in means for the stomach. The heart and skeletal
431 muscle should be defined as ‘inward’ because mean IDs were reduced, but constant mean EDs
432 are not recognised, there is no ‘inward-only hypertrophic.’ The same for constant IDs in the
433 pulmonary, skin and intestine, that should be classified as ‘outward-only hypertrophic.’ For
434 unknown reason, the remodeling had been considered only as keeping constant MCSA, and not
435 constant WTh. The results were based on averaged linear dimensions, and hypertensive values
436 demonstrated a high degree of variability and abnormal distribution (**S4 Fig**). Accordingly,
437 conventional averaging of dimensions led to the designation of diverse remodeling patterns.

438

439 **Fig 13. The descriptive conventional classification[11].** The ‘inward’ or ‘outward’ means
440 reduction or increase in lumen; the ‘hypertrophic,’ ‘eutrophic,’ and ‘hypotrophic’ indicates
441 increase, no change, or decrease in wall cross-sectional area, respectively.

442

443 There is a general assumption that in humans and animals, regardless of the type of hypertension,
444 small arteries develop inward eutrophic or inward hypertrophic remodeling[7],[24]. Other studies
445 have suggested that inward eutrophic remodeling occurs in essential hypertension, while
446 secondary hypertension is associated with hypertrophic remodeling[4]. The concept of uniform
447 remodeling is logical and reasonable since arteries throughout the body have similar structure

448 and regulatory mechanisms. Nevertheless, studies using either random sections or *in vitro*
449 myography present an entire spectrum of possible remodeling patterns. Differences in animal
450 age, heterogeneous post-mortem arterial contraction, variable perfusion pressure, and different
451 histoprocessing and morphometry techniques may account for data inconsistency[3],[47].
452 However, our attempt to find uniform remodeling in 1K1C rats by using the same tissue
453 preparation and morphometry technique simultaneously in ten organs failed, and instead revealed
454 distinctive, and even opposite patterns (**Table 2**).

455 Many authors have presented data avoiding classification[14],[75],[76],[77]. Reviews on
456 this subject have focused on the method of tissue preparation and the potential for sampling bias,
457 and the general approach has not been revised[3],[24],[25],[31]. Indeed, the conventional
458 classification was based on sketch-drawings, with assumptions regarding geometric
459 parameters[11] (**Fig 13**). The classification presumed, for unclear reasons, the simultaneous
460 movement of ED and ID as inward or outward. Such movement would occur if the artery
461 remodeling is considered as simple inflation or deflation for an elastic, homogenous, gel-like
462 structured thick-wall cylinder[78],[79]. However, the arterial wall is categorized as a
463 multilayered, helically arranged, fiber reinforced composite, with independent ED vs ID
464 displacement[80]. Wall remodeling is therefore a multilayered interaction involving
465 hypertrophy[81],[82], hyperplasia[21],[83], apoptosis[81], hyalinosis and fibrinoid necrosis[84],
466 [85] of vascular smooth muscle cells, as well as deposition of extracellular matrix[86],[87]. The
467 extent of these changes varies from inner to outer layers[88],[83],[76]. We also have found that
468 quite often statistical data from studies could not be even assigned to a certain type because
469 statistical casualties break geometry rules. For example, mean ED and ID remains constant but
470 WTh increased[60], mean ID decreased while MCSA and WTH remain constant[89].

471

472 **The conventional classification is unable to categorize remodeling variants**

473 In order to precisely categorize remodeling variants in consideration of the conventional
474 designations of inward-outward and hypertrophic-hypotrophic (**Fig 13**), possible conformations
475 of a blood vessel were modeled with the approximation of arteries as thick-wall cylinders
476 capable of changing ED and ID independently (**Fig 14**).

477

478 **Fig 14. Numerical modeling approach.** (A) For numerical modeling the ED (2) and ID (5)
479 changes independently. ED could moves outward (vector 1) to increase ED or inward (vector 3)
480 to reduce ED. ID could shift inward (vector 4) to decrease ID or outward (vector 6) to increase
481 ID. (B) An example of numerical classification. The number 1, 2 or 3 indicates increased (\uparrow),
482 stable (=), or reduced (\downarrow) ED respectively. The number 4, 5 or 6 reflects decreased, constant or
483 increased ID. The letter a, b or c identifies diminished, constant or increased WTh. White
484 semicircles – control, black – predicted remodeling variants.

485

486 In a simulation procedure, each variant was considered distinct if a value for only one dimension
487 in the set of five (ED, ID, WTh, MCSA, WLR) was different from the value for another set. The
488 numerical modeling (**Fig 15**) revealed not six, as empirically suggested, but twenty variants of
489 arterial remodeling. For example, increasing ED produces eight (1-4, 1-5, 1-6aI, 1-6aII, 1-6b, 1-6
490 cI-cIII) possible variants by combination of changing ID, WTh and MCSA in different
491 directions. A constant ED results in two variants (2-4, 2-6), and decrease in ED generates nine
492 variants (3-4 aI-aIII, 3-4b, 3-4 cI-cIII, 3-5, 3-6). Variants 1-6 cI-III, 3-4 aI-III and 3-4 cI-III are
493 possible because not only the direction but even gradients of change between ED and ID could

494 elicit unique remodeling patterns, which were impossible to classify with conventional
495 definitions.

496

497 **Fig 15. The formalization of numerical modeling for predicted remodeling variants.** The
498 number **1, 2** or **3** indicates increased (\uparrow), stable (=), or reduced (\downarrow) ED. The number **4, 5** or **6**
499 corresponds to decreased, constant or increased ID. The letter **a, b** or **c** identifies increased,
500 constant or diminished WTh respectively, for the same set of ED and ID. The additional Roman
501 numerals **I-II-III** detail the dynamics in MCSA and WLR for the same set of ED, ID, and WTh
502 due to possible gradients in ED vs ID displacement. The 2-5b variant indicates no remodeling.
503 White semicircles – control, black – predicted remodeling variants.

504

505 Therefore, the conventional term “hypertrophic” would equally apply to nine variants: 1-4, 1-5,
506 1-6cI-III, 2-4, and 3-4cI-III. The term “outward” would cover simultaneously 1-4, 1-5, and all
507 six sub-variants 1-6. The term “eutrophic” would equally apply to 1-6b, 3-4b or 3-4cII, while
508 “hypotrophic” could be true for 1-6aI-II, 2-6, 3-4aI-III, 3-5, and 3-6. According to this numerical
509 classification, in 1K1C rats TAs demonstrated seven remodeling variants across organs (**Table**
510 **2**). Studies using either random sections (**Table 3**) or *in vitro* myography (**S3 Table**) are also
511 very inconsistent.

512

513

514

515

516

517 **Table 3. Numerical classification applied to arteries studied on random histological**
 518 **sections.**

References	Experimental model	Arterial location	NC
Liu et al.[17]	2K1C (PF)	renal arterioles 30-60 μ m > 60 μ m	2-4c 2-5b
Helmchen et al.[44]	2K1C (IF)	interlobular	1-5c
Qin et al.[82]	2K1C(PF)	mesentery	1-4c
Korsgaard et al.[90]	1K1C, unfixed, wire myograph	mesenteric III branch renal arcuate arteries	3-4cIII 2-5b
Deng et al.[77]	wire myograph, 2K1C 1K1C	mesentery III order mesentery III order	3-4cIII 3-4cII
Zhou et al.[87]	2K1C, wire myograph	renal arteries	3-4cIII
Kinuno et al.[13]	SHR (PF) SHR+uninephrectomy (PF)	renal interlobular renal interlobular	1-5c 1-6aII
Owens et al.[18]	SHR (PF)	mesentery, I branch II branch III - IV branch	3-4cII 2-5b 3-4cIII
Smeda et al.[16]	SHR (PF)	renal, ID~20-60 μ m renal, ID~300-60 μ m	2-5b 1-5c
Johansson B.[91]	SHR (PF)	cerebral extracranial	1-6cI
	SHRSP (PF)	cerebral intracranial cerebral extracranial cerebral intracranial	1-5c 1-6cIII 1-6aII
Nordborg et al.[12]	SHR (IF)	IF, mesentery	3-4cIII
	SHRSP (IF)	mesentery	3-4cII
	SHR (IF)	renal	1-5c
	SHRSP (IF)	renal	1-6cI
Leh et al.[14]	SHR (IF)	renal afferent arterioles	1-6cI
Ohara et al.[15]	SHR (PF)	PF, renal interlobular	2-4c
Kost et al.[92]	SHR (PF)	renal afferent interlobular arcuate	2-5b 2-5b 3-5a
	SHR+AngII (PF)	renal afferent interlobular arcuate	2-5b 1-5c 1-5c
Lee et al.[47]	SHR (PF)	mesentery, ID~120-250 μ m	1-5c
Limas et al.[50]	SHR, age dependent (PF) renal, ED~50-100 μ m	10 weeks	2-5b
		20 weeks	1-5c
		28 weeks	3-4b
		48 weeks	2-4c
Casare et al.[19]	AngII (PF)	renal afferent interlobular	1-6cII 1-6aII
Berry et al.[21]	DOCA (PF) intrarenal,	ED ~ 400-500 μ m	1-6cI
		ED~ 200-300 μ m	1-5c
Mazzali et al.[40]	Hyperuricemia (IF)	renal afferent arterioles	1-4c

519 2K1C, two kidney-one clip; Ang, angiotensin; DOCA, deoxycorticosterone acetate; IF,
520 immersion fixed; PF- perfusion fixed; SHRSP – spontaneously hypertensive rats stroke prone;
521 NC – numerical classification.

522 We have found only one study that applied a numerical approach to remodeling[31]. The
523 numerical classification proposed here unambiguously defines any wall conformation, and could
524 be applicable to remodeling not only in hypertension but diabetes, atherosclerosis[93],[94], high
525 or low blood flow[48],[27], restenosis[95], vasculitis[96], or bronchial remodeling[97].

526

527 **Remodeling patterns are not identified with WLR, RI or GI**

528 To decide whether remodeling varieties (**Table 2, 3 and S3 Table**) truly exist or are derived
529 from conventional averaging, we tested methods recognizing that arteries follow one of twenty
530 predicted variants.

531 Increased or decreased WLR is widely used as the main criteria of ‘inward’ vs ‘outward’
532 wall conformation[3],[4],[5]. It is not appropriate because WLR could be similarly increased in
533 nine, decreased in eight, and unchanged in three variants. Therefore the frequently used WLR
534 could unambiguously define neither wall thickening nor lumen narrowing (**Fig 15**).

535 RI and growth index GI are also regarded as the main parameters to estimate
536 remodeling[35],[98],[24]. According to the primary data[6], basilar arteries in SHR developed
537 the variant 3-4cIII as a combination of hypertrophy and rearrangement (remodeling) of vascular
538 smooth muscle cells (**Fig 16**). The proposed combination was explained with two formulae. The
539 formula (1) below was based on the first presumption: what would the ID be if vessels
540 underwent the variant 3-4cII (ID3-4cII) i.e. hypertensive MCSA (MCSA_{hr}) remains equal to
541 normal MCSA (MCSA_n)? The formula (2) below was based on the second presumption: what
542 would the ID be if vessels developed the variant 2-4c (ID2-4c) i.e. hypertensive ED (ED_{hr})

543 remains equal to normal ED (EDn)? To create the formula (1), the larger MCSAhr was ignored
544 in favor of a proposed MCSAn=MCSAhr to calculate percent of encroached lumen if the variant
545 3-4cII would occur:

$$546 \text{Percent of encroached lumen} = \frac{IDn - \sqrt{EDhr^2 - 4MCSAn/\pi}}{IDn - IDhr} = \frac{IDn - ID(3-4cII)}{IDn - IDhr} \quad (1)$$

547 For the formula (2), the smaller EDhr was ignored in favor of a proposed EDn=EDhr to calculate
548 percent of encroached lumen if the variant 2-4c would occur:

$$549 \text{Hypertrophy} = \frac{IDn - \sqrt{EDn^2 - 4MCSAhr/\pi}}{IDn - IDhr} = \frac{IDn - ID(2-4c)}{IDn - IDhr} \quad (2)$$

550 The primary definition “percent of encroachment lumen” in the first formula was renamed the
551 “remodeling index”, and the second formula for “hypertrophy” was modified to “growth index”
552 by[98]. Unlike the second formula, GI counted true MCSAs:

$$553 GI = \frac{MCSAhr - MCSAn}{MCSAn} \quad (3)$$

554 Accordingly, the formulae were intended to quantify only *a subjective interpretation* of 3-4cIII
555 as a combination of 2-4c and 3-4cII but that interpretation could be valid as any combination
556 between eight variants with diminished ID (2-4c, 3-4aI-III, 3-4b, 3-4cI-III). Therefore
557 calculation of RI is also incorrect because the formulas were designated to analyze only the
558 proposed combination (**Fig 16**).

559

560 **Fig 16. The first attempt to quantify remodeling**[6]. The reductions in both mean ED and ID
561 were interpreted by the authors as a possible combination of hypertrophy (variant 2-4c) and
562 rearrangement of vascular smooth muscle cells (variant 3-4cII). White semicircles – control,
563 black – predicted remodeling variants

564

565 GI does not bear any information about growth, as it simply counts the percentage change in
566 MCSA that could be assigned to different variants (**Fig 15**). This may account for the variable RI
567 and GI values found in *in vitro* myography experiments, which are considered gold standard due
568 to sampling of similar mesenteric arteries branches. According to formulae (1) and (2), the sum
569 RI+GI must be 100%, but it appears only in the first authors' publication (**S3 Table**). The
570 question still remains: how to recognize a particular remodeling variant?

571

572 **Remodeling patterns are not identified on 2D graphs**

573 The analysis of the simple annular ring in histological sections represents a conundrum, since
574 arteries have no 'reference points' from which they have been modified to assume a remodeled
575 shape (**Fig 17**). According to the literature and own data, any of the 20 predicted types of
576 remodeling could occur (**Table 2, 3 and S3 Table**).

577

578 **Fig 17. The conundrum in arterial remodeling assessment.** Each artery could be an image of
579 another one before or after remodeling: there is no "reference point" indicating its previous
580 dimensions. Scale 10 μm .

581

582 As in control rats, primary measurements were organized in 5 μm intervals, the mean value for
583 each interval was calculated, and the complex profile and linear regression equations built from
584 the means of intervals (**S4 Table**). Data analysis revealed that comparison of complex profiles
585 also could not be used to identify remodeling variants because it was complicated by the
586 existence of uncertain starting points for superimposition of a hypertensive graph on the control
587 one (**Fig 18A, S5 Fig**). Furthermore, there are no data indicating how vessel remodeling
588 progresses in a length-wise fashion, except for the presence of increased vessel

589 tortuosity[29],[63]. While the common starting points were unclear, varying accumulated
590 frequencies suggested distinct remodeling patterns even in different segments within the same
591 organ (**Fig 18A, S5 Fig**).

592

593 **Fig 18. Considerations for arterial remodeling assessment.** (A) The profile of hypertensive
594 (red) renal TAs was superimposed on the control (green). The starting point (the curved arrow) is
595 uncertain: inward–outward remodeling (the solid double arrow) is well known, while proximal–
596 distal shifting (the dashed arrow) is unacknowledged. Uneven accumulated frequencies displaced
597 interval values irregularly. Complementary graphs are in **S5 Fig**. (B) Linear regression lines of
598 renal TAs demonstrated decreased ID (solid lines) and increased WTh (dashed lines) in
599 hypertension. The slopes were different ($P<0.0001$). Complementary graphs are in **S6 Fig**.

600

601 Next, we tested if linear regression could identify remodeling patterns. Compared to the
602 complex profiles, the linear regressions demonstrated smaller IDs and bigger WThs for the same
603 ED similarly in all organs, except the kidney and adrenal (**Figure 18B, S6 Fig**).

604 To understand how shifts in regression lines could verify remodeling, every predicted
605 variant, for each organ, was simulated from control linear regression equations (**S4 Table**). Some
606 studies considered that not WTh/ID but other ratios such as EP/IP[42],[41], or
607 MCSA/LCSA[13],[38],[45], or MCSA/IP[22] are more reliable and informative. Therefore we
608 tested all possible dimensions and their ratios to demonstrate the lack of reliability in 2D graphs.
609 Simulation was done for: linear sizes including ED, ID, WTh, internal perimeter (IP), external
610 perimeter (EP); areas, such as MCSA, lumen cross sectional area (LCSA), total area
611 (MCSA+LCSA), and ratios, including WLR, EP/IP, and MCSA/LCSA.

612 For all tested dimensions, the regression line displacement from the control line was
613 similar for as many as 11 variants (**Fig 19, S7 Fig**). Any linear size, area or their ratio exhibited
614 similar displacement for many different remodeling variants. While the use of linear regression
615 equations can verify the presence of remodeling, the equations are unable to define a specific
616 remodeling pattern. Thus, it was not possible to distinguish arterial spatial conformations on
617 routine 2D graphs.

618

619 **Fig 19. Remodeling variants are not distinguishable in two-dimensional graphs.** The
620 simulated remodeling of renal arteries for the variants 1-4c, 1-5c and all 1-6. For any parameter a
621 line shift was similar to others, and could distinguish only increase or decrease from control but
622 not verify a particular pattern. MCSA - media cross sectional area, LCSA – lumen cross
623 sectional area. Complementary graphs are in **S7 Fig**.

624

625 **A method of 3D-modeling simulation recognizes remodeling type**

626 A 3D-modeling simulation was applied to address the limitations with 2D graphs. There were
627 four steps to this method. For example, for the brain, IDs and WThs were calculated from the
628 control linear regression equations (**S4 Table**) and placed in 3D-graphs (**Fig 20, S8 Fig**). Then,
629 IDs and WThs for each of twenty possible remodeling variants were computed from the control
630 equations and added to the 3D graphs. Every remodeling pattern was simulated by calculating its
631 possible maximal deviation in dimensions that were arbitrarily limited to 300% for wall
632 thickening or lumen widening, and 99% for wall thinning or lumen narrowing. IDs and WThs
633 were then calculated from the hypertensive regression equations (**S4 Table**) and also added to
634 3D graphs. Finally, line proximity between real and predicted values was verified with methods

635 of analytic geometry in 3D space[99]. From all possible variants the hypertensive remodeling in
636 cerebral TAs corresponded only to the variant 2-4c.

637

638 **Fig 20. 3D modeling simulation for the brain.** In 3D graphs, the control equation (green lines)
639 and each predicted variant (blue lines) are connected by the planes (grey) representing sets of all
640 possible values that ED, ID, and WTh could acquire during transformation. The red lines
641 represent the regression equation for hypertensive rats (**S4 Table**). The red lines were not
642 congruent with any of the predicted planes, except for the variant 2-4c. The method readily
643 distinguishes the gradients Δ ED, Δ ID, and Δ WTh for the variants 1-6 and 3-4 that could only be
644 implied in 2D graphs in **Fig 19**. Axis X – ED, axis Y – ID, axis Z – WTh, μm . Complementary
645 graphs are shown in **S8 Fig., S19 and S20 Video files**. Throughout the article, coordinates in
646 some graphs were rotated at different angles to optimize display.

647

648 The brain, heart, lung, bronchi, liver, stomach, intestine, skin, and skeletal muscle developed the
649 same remodeling variant 2-4c, supporting the concept of uniform remodeling (**Fig 21**).

650

651 **Fig 21. The common remodeling variant in hypertensive rats.** The same variant 2-4c was
652 found in most organs.

653

654 We also found two novel variants of remodeling, as predicted. The adrenal TAs exhibited more
655 complex spatial rearrangements. The distal segments with ED \sim 10-20 μm followed the variant 2-
656 4c, as other organs (**Fig 22A and S9 Fig**). The proximal segments of ED \approx 30-50 μm
657 transformed oppositely – through the variant 1-6aI with a distended lumen and reduced WTh and

658 MCSA (**Fig 22B and S10 Fig**). This was evident even on 2D graphs (**S6 Fig**). The adrenals in
659 the 1K1C model experienced a high blood flow. The blood, shunting from clipped or ligated
660 main renal arteries, induced the variant 1-6aI in the proximal segments, i.e. wall distention and
661 thinning, as described for remodeling in high flow models[26],[27],[30]. However, distal TAs
662 demonstrated the same variant 2-4c as in other organs. It may be that smallest arteries (arterioles)
663 are structurally more resistant to distention from high blood flow, due to the absence of elastic
664 laminae and adventitia[100]. Such segmental arterial remodeling has not been verified previously
665 to our knowledge, and indeed was present in TAs with EDs ranging from 10-50 μ m.

666

667 **Fig 22. The distinct remodeling variants in hypertensive rats.** (A) Adrenal distal segments of
668 ED \sim 10-20 μ m were congruent with the variant 2-4c, as other organs. Complementary graphs are
669 in **S9 Fig**. (B) Adrenal proximal segments of ED \sim 30-50 μ m were congruent with the variant 1-
670 6aI. Complementary graphs are in **S10 Fig**. (C) The variant 3-4cIII appeared in the kidney.
671 Complementary graphs are in **S11 Fig**.

672

673 Renal TAs developed the variant 3-4cIII (**Fig 22C and S11 Fig**), that was responsible for the
674 largest TRR increase. Reduced blood flow in stenotic kidneys[28],[29] should initiate one of the
675 variants (3-4a-b), which has been described for low-flow models[26],[30]. The renal arteries also
676 experienced enhanced RAS activity that likely initiated the variant 2-4c in most organs.
677 Development of the variant 3-4cIII might therefore reflect an interaction between flow-induced
678 and pressure-induced stimuli. We speculated that remodeling by the variant 3-4cIII is a
679 phenomenon of great importance that could be extended beyond the 1K1C model. It is well
680 known that hypertension is combined with atherosclerosis in 70-80% of patients[101]. Therefore

681 TAs in the brain and heart could be exposed to the low flow due to stenotic atherosclerotic
682 plaques in proximal segments of cerebral and coronary arteries, and activated RAS as well.
683 Presumably, TAs would develop the same variant 3-4cIII with the most increased resistance and
684 reduced flow, which might be responsible for lesions in the heart and brain, being recognized as
685 critical target organs.

686 We have provided the first quantitative evidence that pulmonary arteries respond to the
687 hypertensive stimulus in 1K1C rats, and in the same remodeling pattern as other organs in the
688 systemic circulation. Our data also represent the first evidence that substantial TAs remodeling
689 occurs in the liver, intestine, and bronchi, which are not considered typical targets in
690 hypertension[5],[25].

691

692 **The algorithm enables comparison among different studies**

693 Contradictions of two types have been found in conventional remodeling analysis. First, results
694 do not fit the geometric formulae for annular ring dimensions. We applied the term “statistical
695 artifact” (SA) to indicate them. Second, results could be geometrically correct but showed
696 different, opposite, or no remodeling patterns for the same model, organ, or arterial segments
697 (**Table 2, 3 and S3 Table**). In order to demonstrate that our method could resolve this problem,
698 we used 3D modeling to reanalyze data from studies where linear regression had been calculated,
699 although only few articles could be found. Comparative analysis would not be possible with
700 primary averaged data only.

701 Two studies claimed unusual non-pressure-related wall thickening after comparing
702 arterial remodeling between aortic coarctation and Goldblatt-induced hypertension in
703 rats[17],[102]. One study[102] demonstrated many SAs (**S5 Table**). After coarctation, means of
704 WThs and WLRs were elevated despite stable EDs, IDs, and MCSAs. In 1K1C rats, stable IDs

705 with greater EDs, WThs, and WLRs pointed to the variant 1-5c, although stable MCSAs did not
706 support this designation. A separate study[17] was also inconsistent (**S6 Table**). In both models,
707 averaged dimensions indicated no remodeling in larger segments, but the variant 2-4c in smaller
708 arteries. In coarctation, SA occurred in arteries of ED interval 31-60 μm : unchanged MCSAs did
709 not correspond to stable EDs with lumen narrowing. In two kidney one clip (2K1C) rats,
710 constant EDs and IDs did not follow MCSA and WTh expansion.

711 Using primary data[17],[102], we calculated linear regressions and applied 3D-modeling.
712 Aortic coarctation exhibited no remodeling in renal or cremaster arteries (**Fig 23A, S12 and S13**
713 **Fig**), i.e. there is no non-pressure related TAs thickening distally to aortic coarctation, in
714 normotensive renal and cremaster arteries, as mean values indicated[102],[17]. No further data
715 have been published to support such normotensive thickening[5],[24]. In contrast, the variant 1-
716 6aI has been recognized in the liver[48]. Goldblatt hypertension induced the same variant 2-4c in
717 all segments of cremaster and renal arteries (**Fig 23B and 23C, S14 and S15 Fig**). We also used
718 primary data to recalculate morphometry in renal arteries in SHR[39], and found the variant 2-4c
719 (**Fig 23D and S16 Fig**). Accordingly, our data suggest that the variant 2-4c is predominant in
720 hypertensive remodeling. The variant 1-5c (stable lumen with wall thickening) was identified in
721 a clinical case of pulmonary hypertension due to congenital mitral stenosis[55] (**Fig 23E and**
722 **S17 Fig**). That example underlines the precision of our proposed algorithm, since primary data
723 included only fourteen measurements from one pulmonary biopsy. This type of pulmonary
724 hypertension is developed during organogenesis and intensive arterial vascular smooth muscle
725 cells proliferation[103], which could be associated with a stable lumen and wall thickening.
726

727 **Fig 23. The method enables direct comparison between different studies.** (A) Aortic
728 coarctation had no effect on remodeling in renal and cremaster arteries. Complementary graphs
729 are in **S12 and S13 Fig.** (B, C, D) The same variant 2-4c characterized remodeling in cremaster
730 arteries of 1K1C, untouched kidneys of 2K1C rats, and kidneys in SHR respectively.
731 Complementary graphs are in **S14-S16 Fig.** (E) The distinctive variant 1-5c was detected in
732 pulmonary arteries due to congenital mitral valve stenosis Complementary graphs are in **S17.**
733 (F) Data of the subcutaneous arteries was approximated inappropriately with linear regression.
734 Complementary graphs are in **S18 Fig.**

735
736 We also tested the linear regression on subcutaneous arteries, reported in patients with essential
737 hypertension[104]. 3D-modeling detected multiple curved lines (**Fig 23F and S18 Fig**) that were
738 the result of an incorrect approximation of the area / length scatterplot by the linear regression.
739 Plotting areas against lengths follows the exponential function[21],[22] (**Fig 19**). The same error
740 we found in other studies[18].

741
742 **Connecting measurements to hemodynamic values**
743 In addition to proposed TRR and TC, it was also desirable to estimate how media volume had
744 changed in remodeling. The terminal media volume (TMV) could be calculated from complex
745 profiles, similar to TC. However, in the current study we found that variation in accumulated
746 frequencies would significantly affect TC and TMV values. The same negative issue for complex
747 profiles to distinguish remodeling patterns was described above (**Fig 18A**). Accordingly, the
748 linear regression equations, established for each organ, with counting ID and ED for the largest
749 (50 μ m) and smallest (10 μ m) vessel calibers, provided more reliable estimates. TC and TMV
750 were then computed by the formula for the truncated cone volume:

751
$$TC = \pi h^*(IDlc^2 + IDlc* IDsc + IDsc^2) / 3 \quad (4)$$

752
$$TMV = \pi h^*(EDlc^2 + EDlc* EDsc + EDsc^2) / 3 - TC \quad (5)$$

753 where EDlc, IDlc and EDsc, IDsc were the largest and smallest caliber limits, respectively, and
754 h was the range of calibers ($50 \mu\text{m} - 10 \mu\text{m} = 40 \mu\text{m}$).

755 Using this calculation, 1K1C hypertension induced a significant 2 - 5-fold increase in
756 TRR in all organs (**Fig 24A**), correlating significantly with data obtained by physiological
757 measures: in the Goldblatt model increased RVR has been shown in every organ, and is most
758 pronounced in kidneys[105],[106],[107],[108]. It is noteworthy that the kidney has the highest
759 increase in TRR (9-fold), while the pulmonary arteries – the lowest. The increased TRR and TC
760 observed in the adrenal glands were due to different remodeling patterns in the proximal and
761 distal arterial segments.

762

763 **Fig 24. Hemodynamic parameters are enhanced variously in different organs.** (A) There
764 was no clear correlation between the values of TRR, TC, TMV. For example, in the brain
765 enhanced TMV corresponded to a large drop in TC but a small increase in TRR. In the kidney a
766 small elevation in TMV corresponded to a small lowering in TC but augmented TRR. (B) An
767 equal 50% increase in wall expansion in kidney and lung arteries would cause marked
768 differences in TRR and TC increases, while TMV would only increase mildly and remain
769 comparable between the two organs. Different dynamics for TRR, TMV, and TC are the result of
770 different initial organ specific dimensions. x - fold increase.

771

772 In normal rats TC correlated with the regional BFR (**Fig 10**). In 1K1C rats TC demonstrated
773 maximum decrease in the brain, liver and intestine, and minimal lowering in the skin, pulmonary

774 and skeletal muscle arteries (**Fig 24A**). However, it was difficult to compare our results with
775 pathophysiological studies because regional BFR has been shown as decreased, increased, or
776 stable as well[105],[109],[110]. TMV indicated arterial wall thickening, with maximum of 20-
777 25% in the brain, intestine and pulmonary arteries, and minimum of 2-5% in skeletal muscle,
778 skin and kidney, which were not proportional to TRR and TC (**Fig 24A**). Presumably, basic
779 organ-specific arterial dimensions could have significant impact on hemodynamic consequences
780 of remodeling being the main cause to various organ dynamics of TRR, TC, and TMV (**Fig**
781 **24B**). Evidently, hemodynamic correlations of TRR, TC and TMV need further investigation.

782 An important goal of our work was to determine if random tissue sections are a reliable
783 source of data. Numerous studies of remodeling, exploring random tissue sections have been
784 published in the past three decades, with most in the 1980s or 1990s, so the majority of our
785 references are dated 15-20 years back. Since then *in vitro* myography studies dominate
786 significantly, while morphometric studies on random sections have been infrequent.

787 Started from the study[51], the post-mortem contraction has been considered a main
788 cause of high variability in arterial morphometry. Perfusion fixation was intended to eliminate
789 post-mortem arterial contraction, especially with preliminary application of a vasodilator[31] or
790 vessel deactivation[111]. The present study indicates that perfusion fixation is not mandatory for
791 morphometry on random tissue sections. In fact, perfused vs non-perfused organs or myograph
792 experiments demonstrated similar stochastic remodeling patterns (**S3 and S4 Table**). Here, we
793 avoided fixation deliberately, performing only immersion fixation in order to have animal (and
794 potentially human) material in the same condition for comparison, since for the latter perfusion is
795 unlikely to be applied. Our data prove that careful consideration of arterial tapering is the most

796 important factor for elaborated morphometry analysis. A similar approach has been widely
797 accepted in atherosclerotic remodeling[93],[57],[112],[113].

798 In conclusion, we have developed an algorithm to quantify and standardize arterial
799 remodeling analysis. Tissue sampling should be random and representative, according to known
800 recommendations[33],[85]. Following fixation, sections must be stained with periodic acid–
801 Schiff and Masson trichrome. Hematoxylin and eosin staining does not always clearly
802 differentiate muscular and adventitial components, especially when hyalinosis, fibrinoid
803 necrosis, or perivascular inflammatory infiltrates occur[85]. Random measurements of EDs and
804 IDs are performed to obtain about 80-100 measurements of arteries with EDs between ~10-50
805 μm . Then, variables are arranged in order of increasing ED, divided into 5 μm intervals, and
806 statistical analysis performed for each interval. Finally, the regression equations, complex
807 profiles, remodeling variant, hemodynamic parameters are computed from interval statistics and
808 compared among models or organs. The algorithm does not require additional counting or data
809 gathering, compared to conventional morphometry of arteries on histological sections, and
810 represents a more informative, standardized approach to arterial profiling.

811

812 **Material and Methods**

813 Experimental protocols were approved by the Animal Ethics Committee at the University of
814 Ottawa and performed according to the recommendations of the Canadian Council for Animal
815 Care. Analyses in normal rats were performed on 20 male Wistar rats (Charles River, Montreal,
816 Québec, Canada), age 20-25 months, and weighing 600-800 g. Five normal male C57BL6 mice
817 (age 20-30 weeks) were used for comparative morphometry. Goldblatt 1K1C hypertension was
818 induced in 20 male Wistar rats (Charles River Laboratories, Montreal, Québec, Canada), age 20-

819 25 months, weighing 600-800 gm. Under isoflurane anesthesia a silver clip with internal
820 diameter 0.26 mm was placed around the left renal artery and the right kidney was removed. On
821 both sides the manipulation was distal to the adrenal arteries to keep them intact. The control
822 sham-operated group (5 animals) was kept under the same living conditions as the experimental
823 animals. Systolic blood pressure was measured weekly by tail-cuff plethysmography. After 60
824 days rats were euthanized with an intraperitoneal overdose of sodium pentobarbital. The brain,
825 lung, heart, stomach, liver, small intestine, spleen, kidneys, adrenals, hip skeletal muscle, and tail
826 skin were excised and immersion-fixed in 10% buffered formalin for 24 h, dehydrated and
827 embedded in paraffin. Three fresh lungs from control rats were prepared for special analysis: the
828 entire lobular pulmonary arterial tree was cleaned from parenchyma up to branches of ED 20-30
829 μm by dissection under a microscope in phosphate-buffered saline. The lung is the only organ
830 which arteries are relatively easy to separate from parenchyma, compare to other organs. Pairs of
831 branches with the same ED but located in proximal and distal segments, 1-2 mm in length were
832 sampled where a branch began, and also embedded in paraffin. Tissue blocks were specifically
833 oriented according to the known anatomical distribution of arteries in order to obtain
834 predominantly cross-circular arterial sections. Histological sections (5 μm thickness) were
835 stained with hematoxylin-eosin and Masson's trichrome. In sections of three to four blocks from
836 each organ, small arteries and arterioles were traced by visual scanning of the entire section.
837 Only vessels with a long- to short-axis ratio < 1.50 were measured. In this way, the error
838 associated with calculating the diameter by averaging the maximum and minimum diameters was
839 minimized ($< 3\%$)[33]. The ED and ID were measured in arteries if the profile had a visibly non-
840 interrupted circular or ellipsoid shape. From those two values the derivatives were calculated as
841 follows: $\text{WTh} = (\text{ED} - \text{ID})/2$; $\text{WLR} = \text{WTh} / \text{ID}$; media cross sectional area (MCSA) = π^*

842 (ED²/2-ID²/2). Microscopy was performed with a Zeiss AX10microscope (Oberkochen,
843 Germany) and images were analyzed by ImagePro Plus software (Media Cybernetics, Bethesda,
844 MD, USA).

845 Descriptive statistics calculated mean, standard deviation (SD), standard error of mean
846 (SEM), coefficient of variation (CV), and frequency distribution to build histograms and
847 complex profiles. Distribution was assessed by Kolmogorov-Smirnov, D'Agostino & Pearson
848 and Shapiro-Wilk tests. Outliers were identified by the combined robust regression and outlier
849 removal (ROUT) method. Acquired linear regression equations were tested to determine if
850 slopes and intercepts were significantly different (P < 0.05) and the goodness of fit coefficient r^2
851 was counted for each equation. Nonlinear regression was approximated with exponential growth
852 equations and the goodness of fit coefficient R^2 was counted for each equation. Extra sum-of-
853 squares F-test and Akaike's Information Criteria (AICc) were used to compare the best-fit values
854 between nonlinear regression equations (P < 0.05). Statistical analyses were performed with
855 GraphPad Prism software (GraphPad Software, La Jolla, CA, USA). Modeling procedures were
856 made with MathCAD version 15.0 (Parametric Technology Corporation, Needham, MS, USA).
857 Three-dimensional graphs were built in OriginPro version 2016 SR0 b9.3.226 (OriginLab
858 Corporation, Northampton, MA, USA).

859

860 **Author contributions**

861 Conceived and designed the experiments: AG SA. Performed the experiments: AG SA.
862 Analyzed the data: AG KB PB. Supervision: KB SP. Writing ± review & editing: AG PB KB
863 CK.

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1160 **Supporting information captions**

1161 **S1 Table.** Normality tests for averaged arterial dimensions in the kidney, heart and

1162 pulmonary arteries.

1163 **S1 Fig. Complex profiles revealed different tapering patterns in organs.** Complementary

1164 graphs to **Fig 6A**. Axis X – the bidirectional common linear scale for the external diameter (ED),
1165 wall thickness (WTh), and internal diameter (ID); axis Y – accumulated frequency of variables
1166 (%).

1167 **S2 Fig. Linear regression equations and RR in different organs.** Lines and RR (blue curves)
1168 represent the best fit for different organs. Complementary graphs to **Fig 6B**. Points are mean \pm
1169 SD for 5- μ m ED intervals; r^2 - goodness of fit coefficients. Corresponding equations are in **S2**
1170 **Table.**

1171 **S2 Table. Linear regression equations from mean \pm SD for 5 μ m ED intervals.**

1172 **S3 Fig. Comparison of linear regressions for ID and WTh between different organs.**

1173 Equations for lungs, bronchi, adrenal glands, stomach and skeletal muscles were very distinctive
1174 ($P<0.0001$). Brain and intestine (*) shared similar equations ($P>0.41$ for ID and WTh). Heart
1175 and spleen (**) were also close ($P>0.43$ for ID and >0.83 for WTh). Equations for kidney, liver
1176 and skin (***) were similar ($P>0.77$ for ID and >0.87 for WTh).

1177 **S4 Fig. Histograms of terminal arteries in the kidney, heart and brain.** The significant
1178 irregularity and asymmetry for dimensions used to calculate statistics in **Table 2**. Data did not
1179 pass conventional statistical tests for normality (negative, $P<0.001$).

1180 **S3 Table. Numerical classification applied to arteries studied via in vitro myography.**
1181 1K1C, one kidney-one clip; 2K1C, two kidney-one clip; Ang, angiotensin; DM2, diabetes
1182 mellitus type 2; EHT, essential hypertension; eNOS, endothelial nitric oxide synthase; GI,
1183 growth index; L-NAME, Nitro-L-arginine methyl ester; op/+, osteopetrosis heterozygous; PM,
1184 pressure myograph; RI, remodeling index; RVH, renovascular hypertension; SA, statistical
1185 artifact; SHR, spontaneously hypertensive rats; SHRSP, spontaneously hypertensive rats stroke
1186 prone; SOD, superoxide dismutase; WM, wire myograph.

1187 **S4 Table. Linear regression equations of terminal arteries derived from mean \pm SD for**

1188 **5- μ m ED intervals.** All equations for hypertensive rats were significantly different from

1189 equations for control rats ($r^2 = 0.99$; $P < 0.0001$).

1190 **S5 Fig. Control vs hypertensive complex profiles in organs.** Control (green) and hypertensive

1191 (red) complex profiles were superimposed. Remodeling patterns are not recognizable.

1192 Complementary graphs to **Fig 18A.**

1193 **S6 Fig. Linear regression lines displaced similarly in most organs.** The ID slopes decreased

1194 (solid arrows), and the WTh slopes increased (dashed arrows). Adrenal arteries demonstrated

1195 opposite directions: the increased ID slopes and decreased WTh slopes. Complementary graphs

1196 to **Fig 18B.**

1197 **S7 Fig. The remodeling of renal arteries has been simulated for variants 2-4c, 2-6a, all 3-4,**

1198 **3-5a, and 3-6a.** Displacement of linear regression lines up or down for any parameter was

1199 similar for many remodeling variants. Complementary graphs to **Fig 19.**

1200 **S8 Fig. 3D-modeling simulation for the brain.** No variants were congruent. Complementary

1201 graphs to **Figure 20.**

1202 **S9 Fig. 3D-modeling simulation for the distal arterial segments in the adrenals.** No

1203 variants, except 2-4c, were congruent. Complementary graphs to **Fig 22A.**

1204 **S10 Fig. 3D-modeling simulation for the proximal arterial segments in the adrenals.** If the

1205 variant 1-6aI appeared, congruence to the variant 2-6a must also be present because data sets

1206 overlap. Complementary graphs to **Fig 22B.**

1207 **S11 Fig. 3D-modeling simulation for the kidney.** If the variant 3-4cIII occurred, simultaneous

1208 congruence to 2-4c and 3-4cII (but not 3-4cI) must be present because data sets overlap.

1209 Complementary graphs to **Fig 22C.**

1210 **S5 Table. Averaged arterial dimensions in rats with aortic coarctation and 1K1C**

1211 **hypertension** (modified from[102]). NC, numerical classification; SA – statistical artifact;

1212 ↑ indicates increase in dimensions vs sham values, *P<0.05, ** P<0.01.

1213 **S6 Table. Averaged arterial dimensions in rats with aortic coarctation and 2K1C**

1214 **hypertension** (modified from[17]). 2K1C, two-kidney, one clip; NC, numerical classification;

1215 SA – statistical artifact; ↑ ↓ indicates increase or decrease in dimensions vs sham values,

1216 *P<0.05, ** P<0.01, *** P<0.001

1217 **S12 Fig. Cremaster arteries in normotensive rats with aortic coarctation have no**

1218 **remodeling.** The control and experimental equations were not different (the variant 2-5b;

1219 P > 0.6-0.7). Calculated from[102]. Complementary graphs to **Fig 23A.**

1220 **S13 Fig. Renal interlobular arteries in normotensive rats with aortic coarctation have no**

1221 **remodeling.** The control and experimental equations were not different (the variant 2-5b;

1222 P > 0.8). Calculated from[17]. Complementary graphs to **Fig 23A.**

1223 **S14 Fig. Cremaster arteries in hypertensive 1K1C rats developed the variant 2-4c.**

1224 Calculated from[102]. Complementary graphs to **Fig 23B.**

1225 **S15 Fig. Renal arteries in 2K1C rats were congruent to the variant 2-4c.** Calculated

1226 from[17]. Complementary graphs to **Fig 23C.**

1227 **S16 Fig. Renal arteries in SHR showed congruence to the variant 2-4c.** Calculated from[39].

1228 Complementary graphs to **Fig 23D.**

1229 **S17 Fig. Pulmonary arteries in a case of pulmonary hypertension due to congenital mitral**

1230 **stenosis developed the variant 1-5c.** Calculated from[55]. Complementary graphs to **Fig 23E.**

1231 **S18 Fig. Recalculated data from human subcutaneous arteries[104].** The graph length vs

1232 area has been approximated with the linear regression, resulting in curved lines. Complementary

1233 graphs to **Fig 23F**.

1234 **S19 Video. 3-D modeling for the brain.** The video demonstrates the variant 2-4c as congruent.

1235 **S20 Video. 3-D modeling for the brain.** The video demonstrates the variant 3-4aIII as an

1236 example of a non-congruent variant.

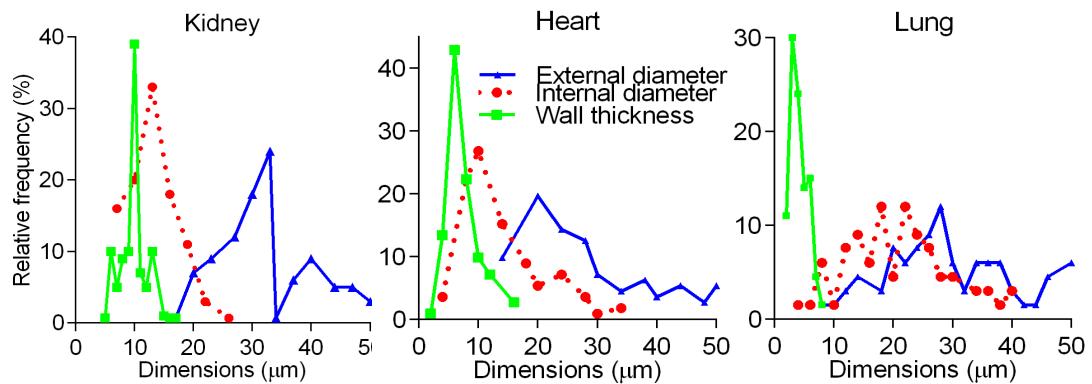


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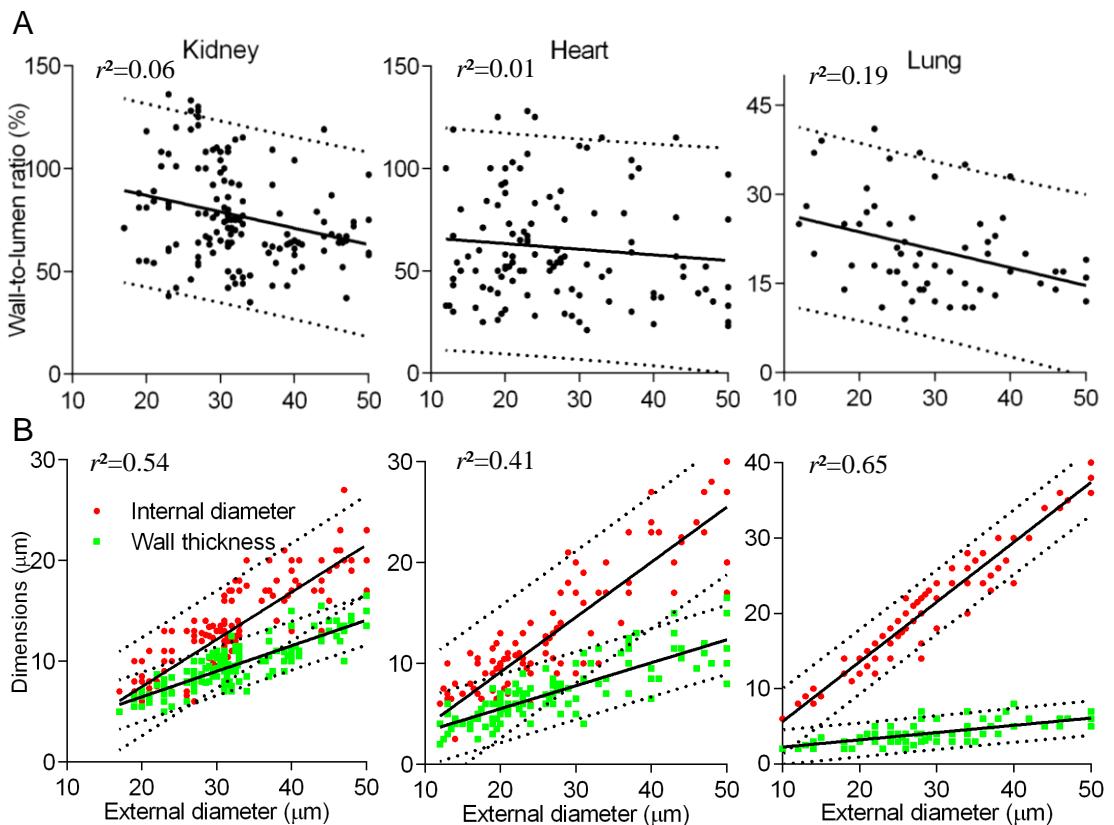


Fig 2.

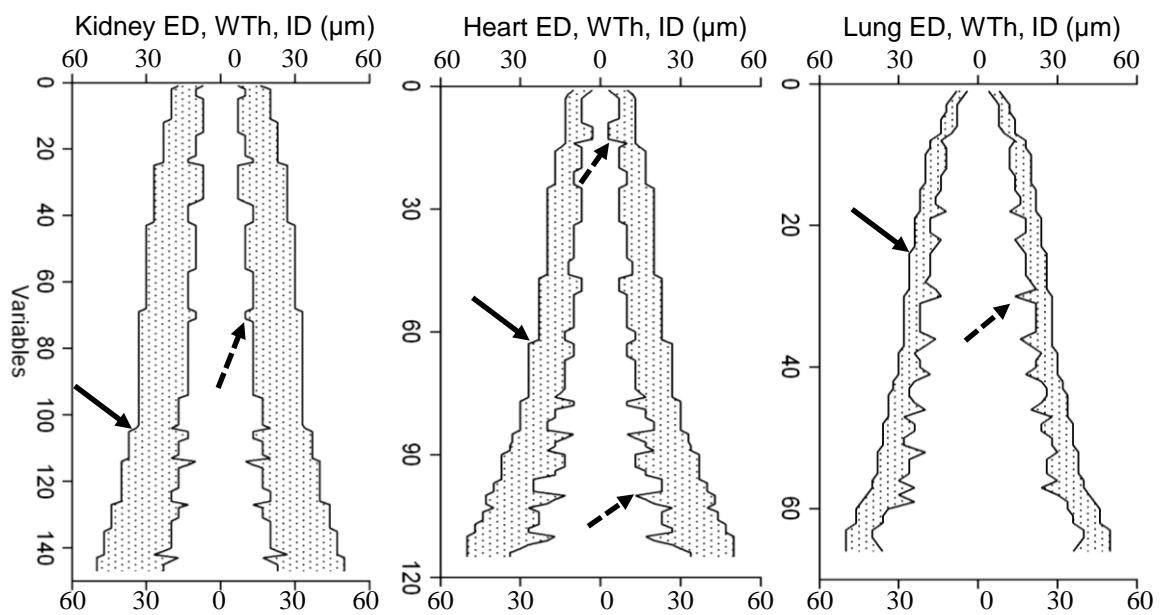
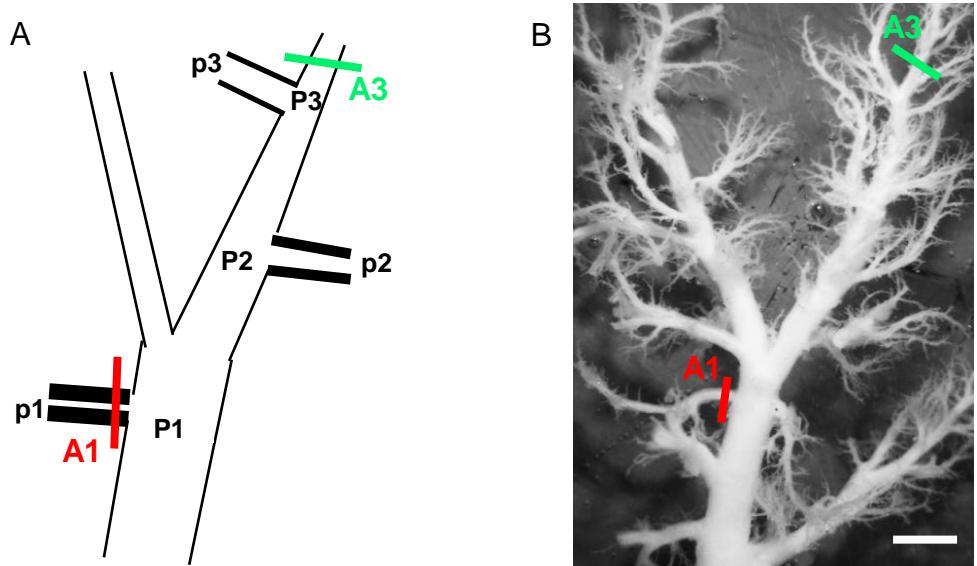


Fig 3.



C

Branch type	ED	ID	WTh	WLR
	μm	μm	μm	
Proximal, axillary or sympodial	28	14	7	50
Distal, regular dichotomous	28	22	3	
Proximal, axillary or sympodial	32	12	10	83
Distal, regular dichotomous	32	26	3	
Proximal, axillary or sympodial	38	20	9	45
Distal, regular dichotomous	38	30	4	
Proximal, axillary or sympodial	40	22	9	41
Distal, regular dichotomous	40	34	3	
Proximal, axillary or sympodial	44	22	10	45
Distal, regular dichotomous	44	34	5	

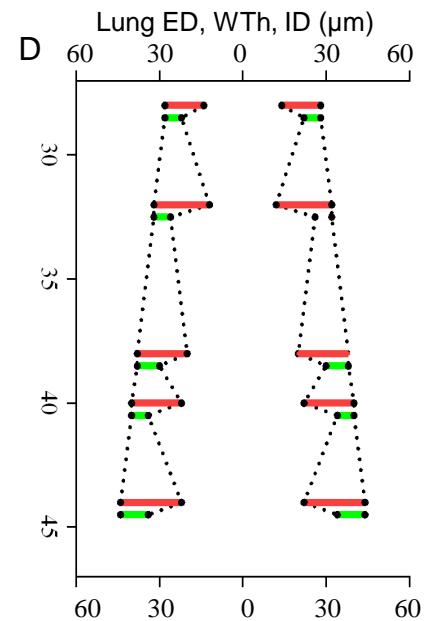


Fig 4.

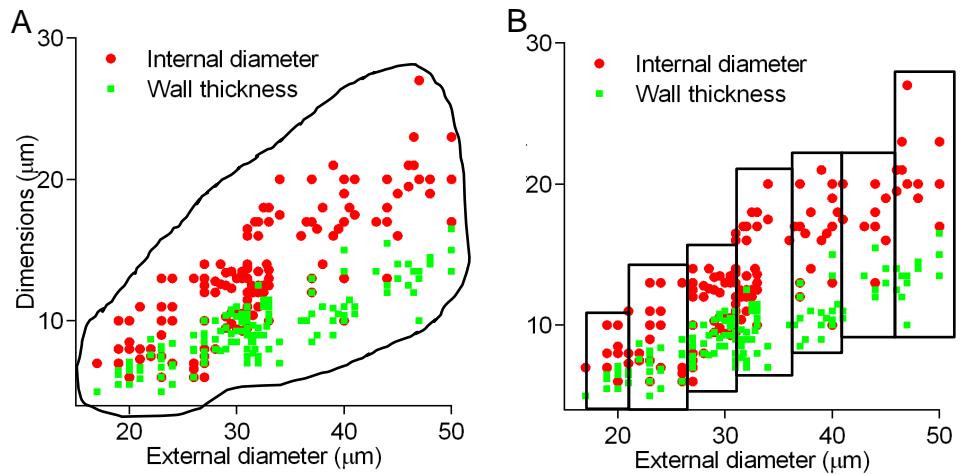


Fig 5.

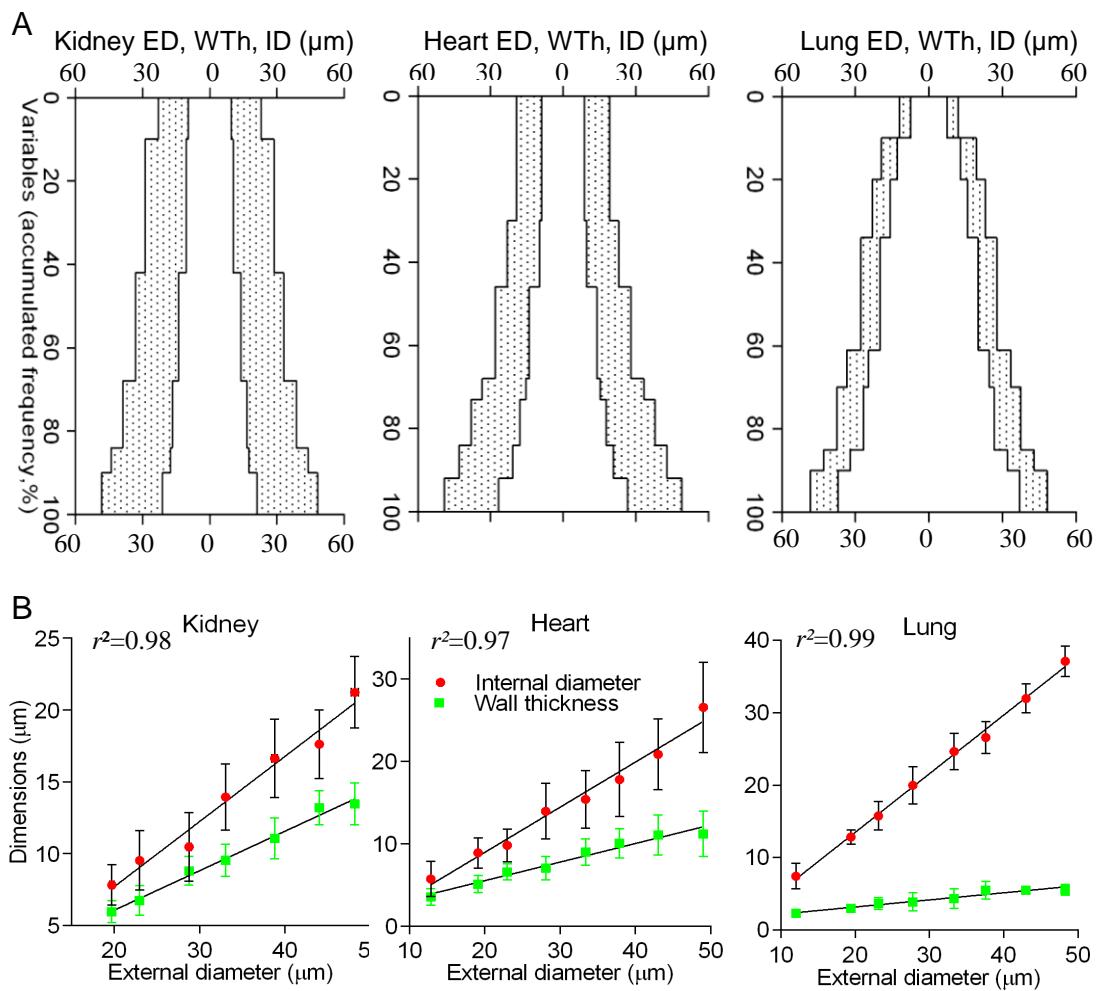


Fig 6.

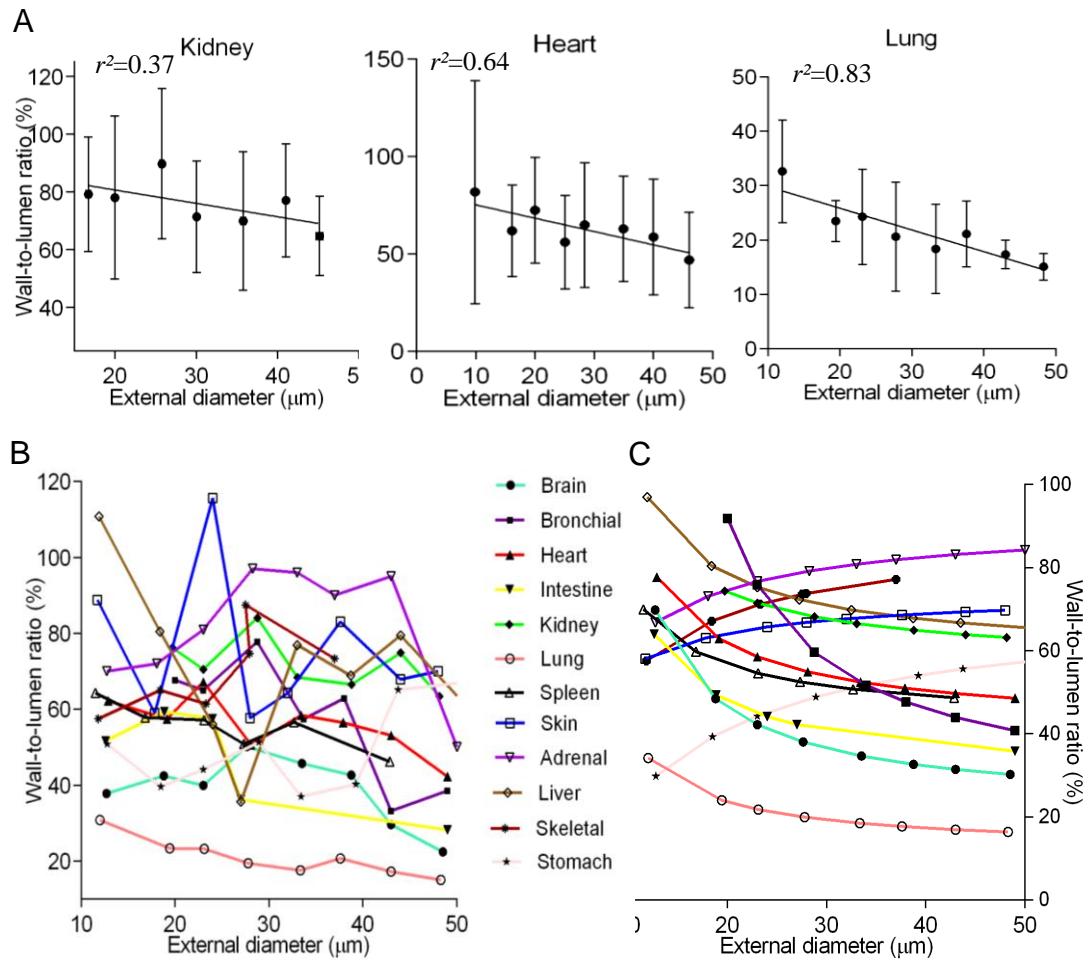


Fig 7.

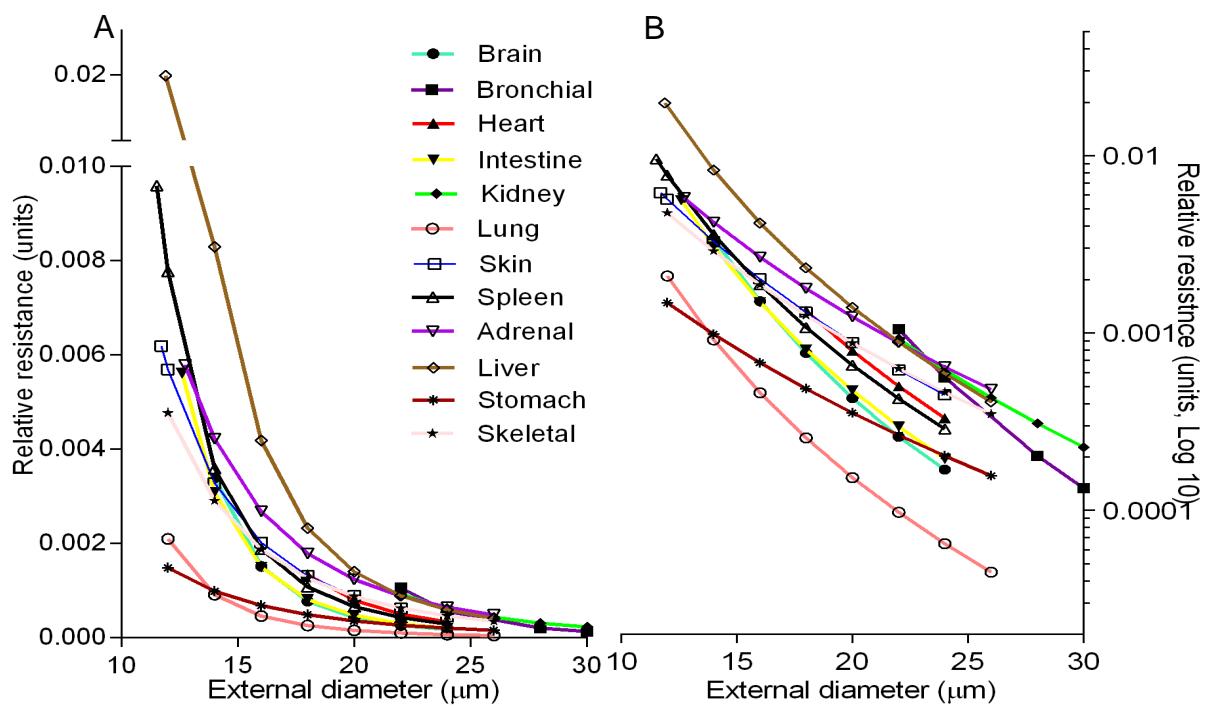


Fig 8.

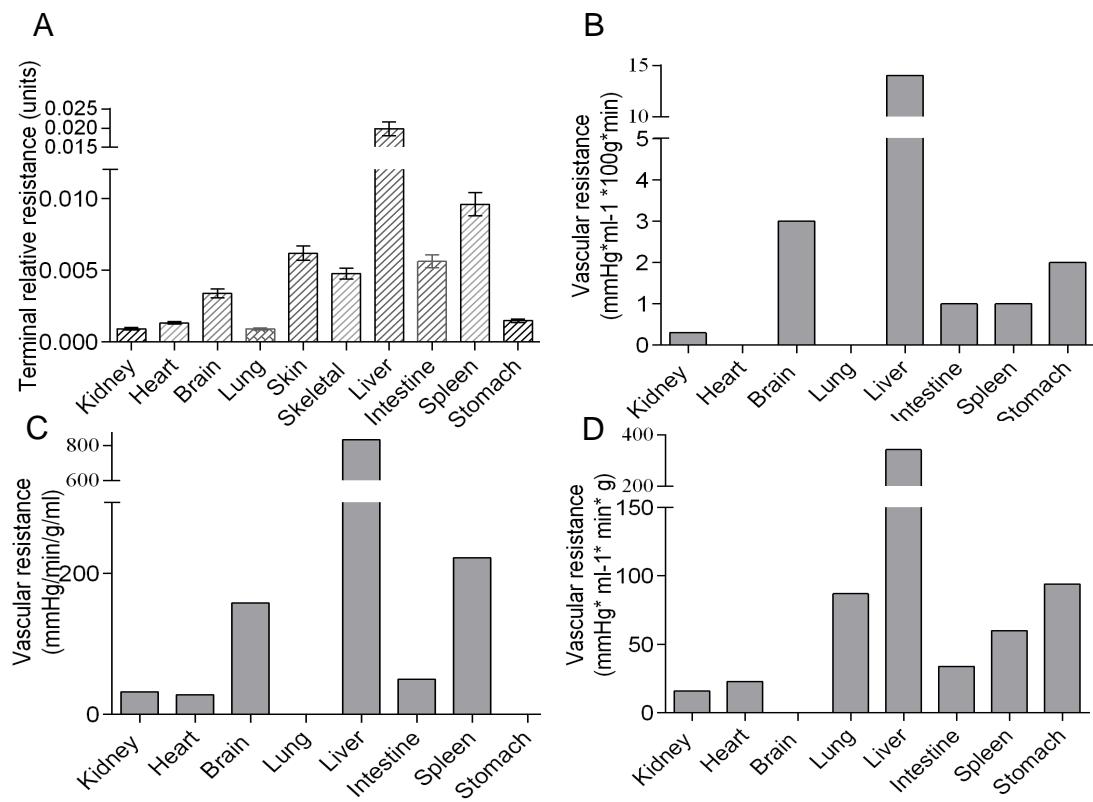


Fig 9.

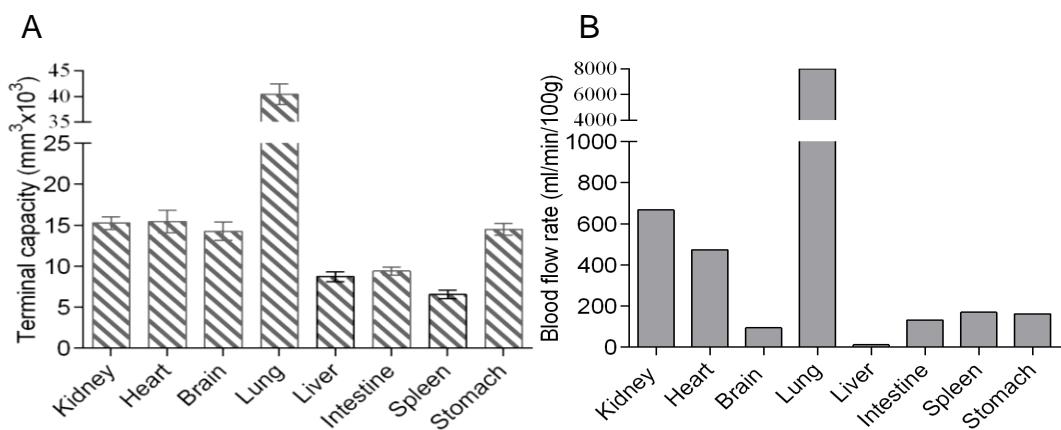


Fig 10.

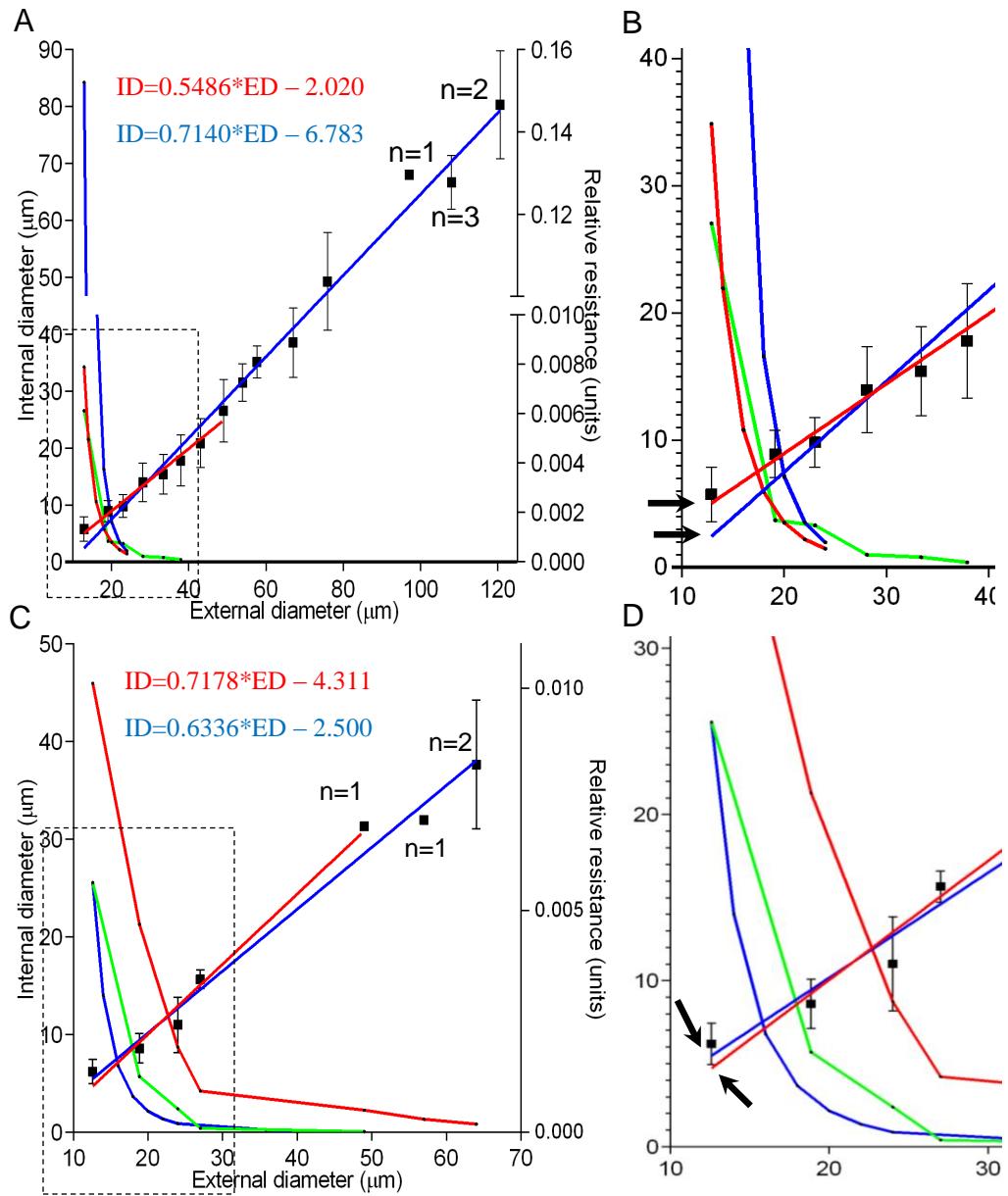
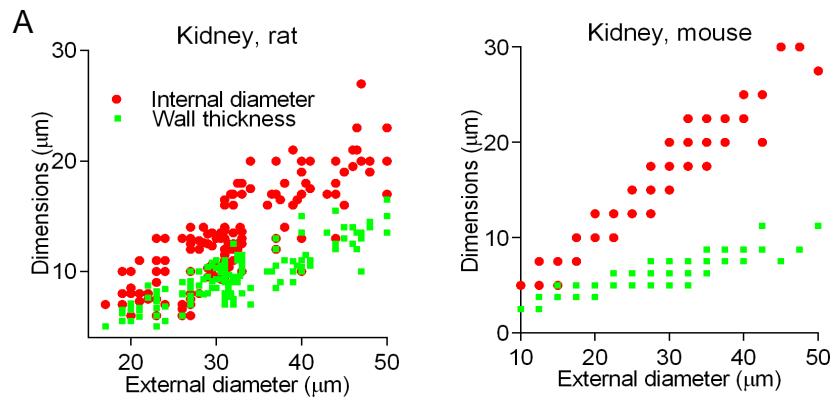


Fig 11.



B

	ED	ID	WTH	WLR
Rat	32.7 ± 7.8	13.3 ± 4.4	9.7 ± 2.3	72.9 ± 27.1
Mouse	30.0 ± 8.7	12.8 ± 5.9	8.6 ± 1.6	67.4 ± 13.5

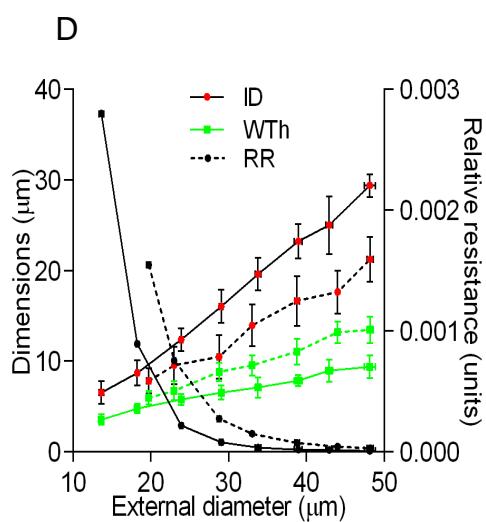
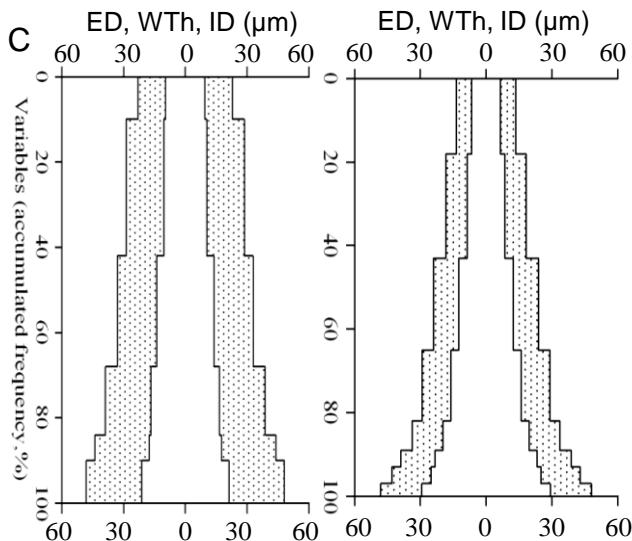


Fig 12.

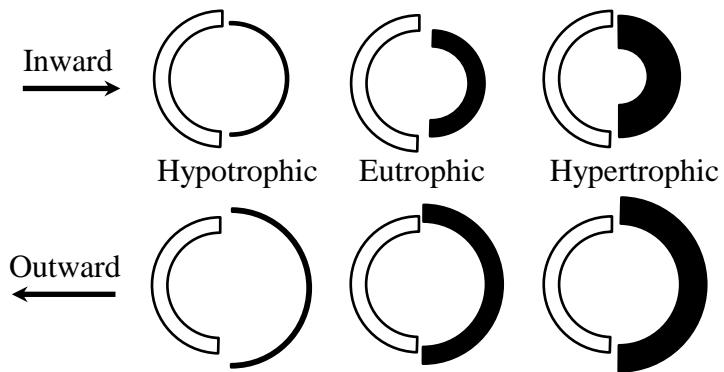


Fig 13.

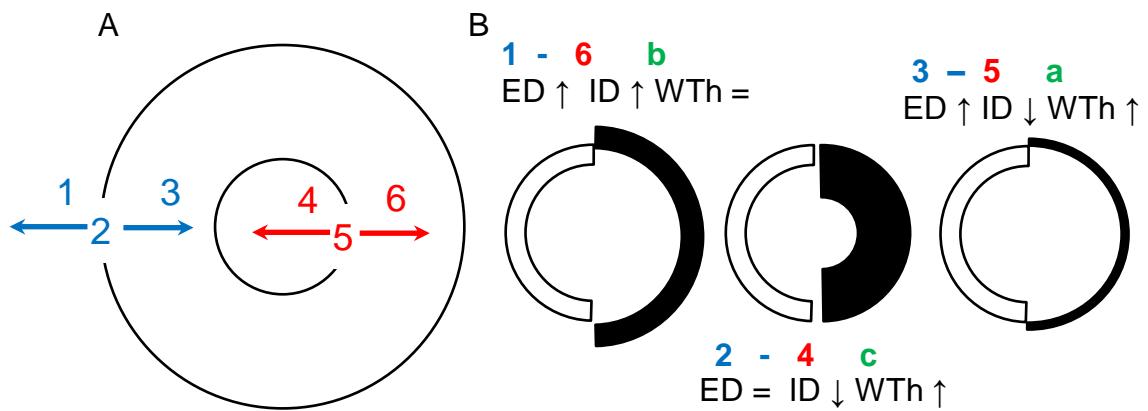


Fig 14.

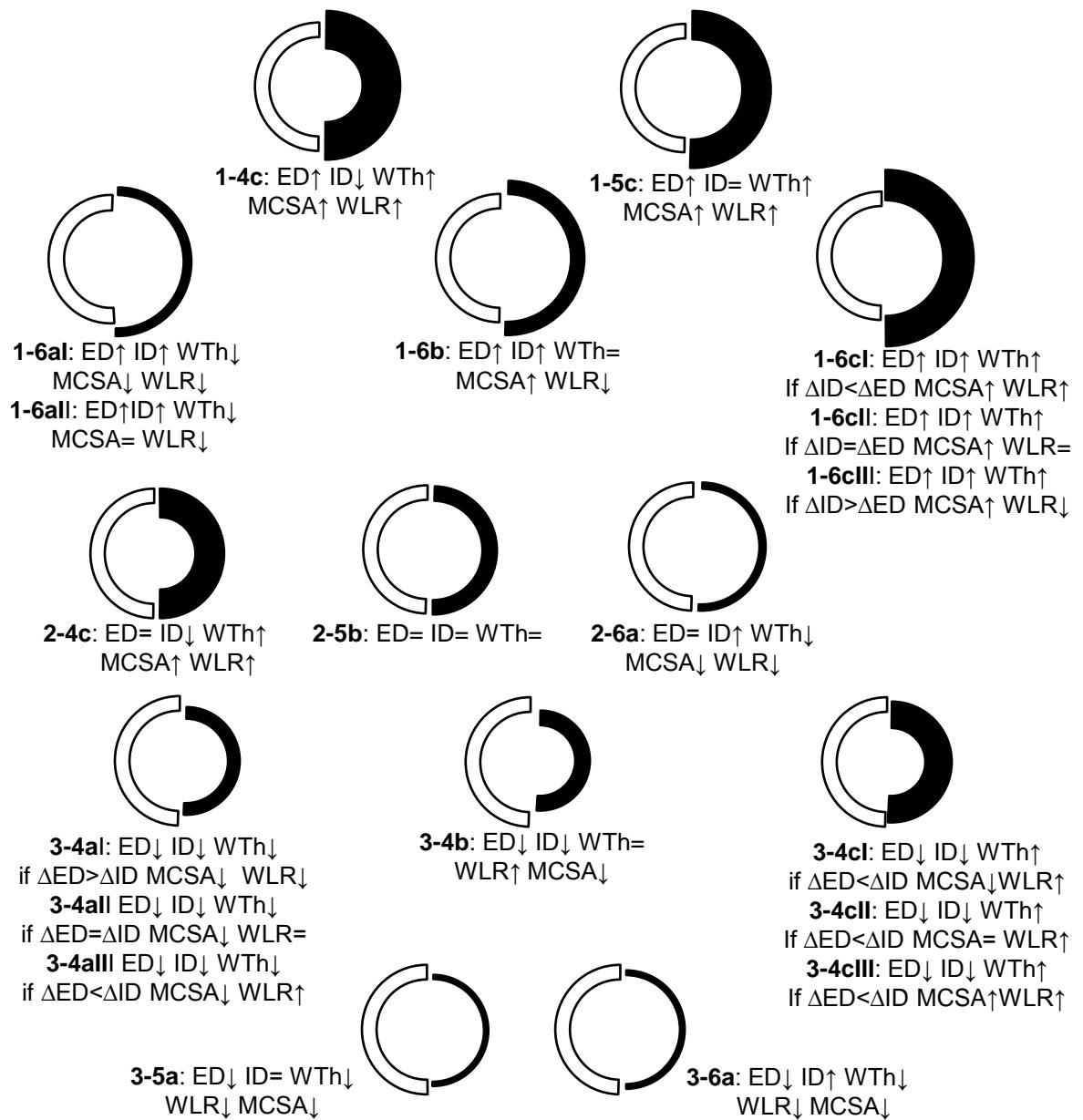


Fig 15.



Fig 16.

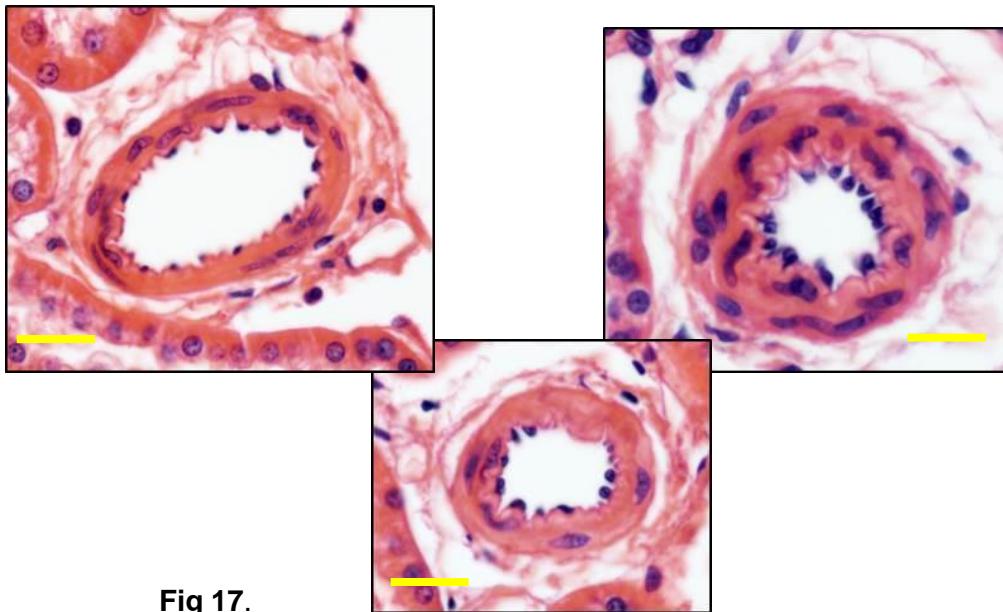


Fig 17.

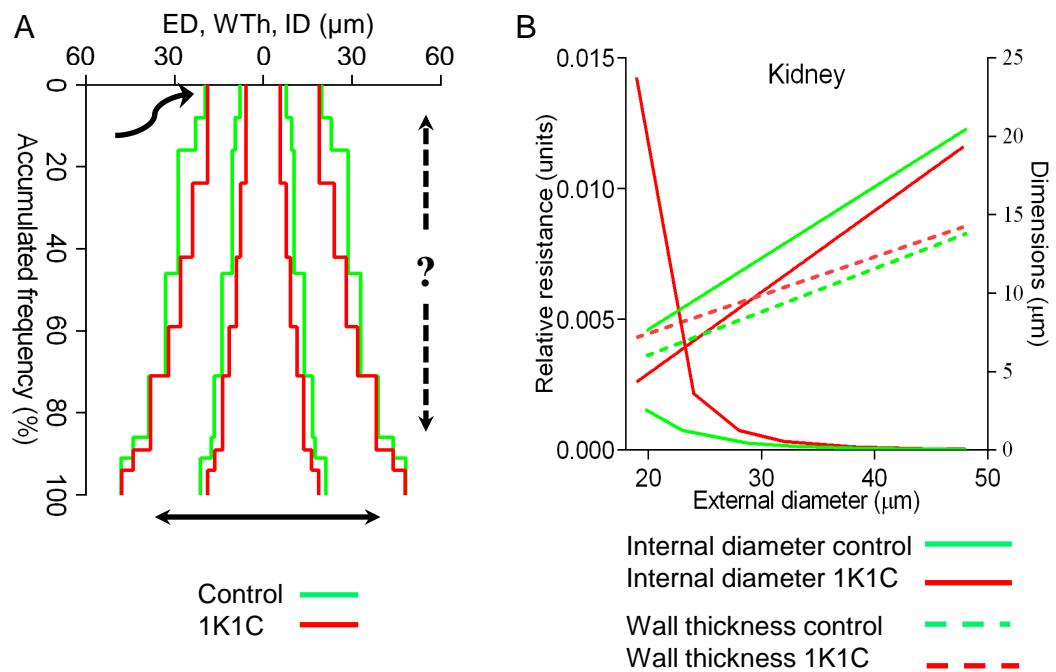


Fig 18.

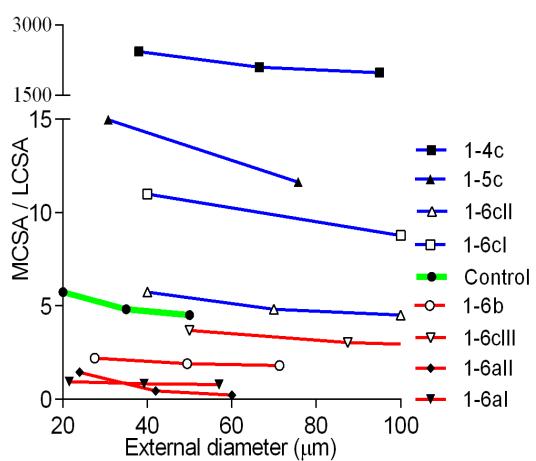
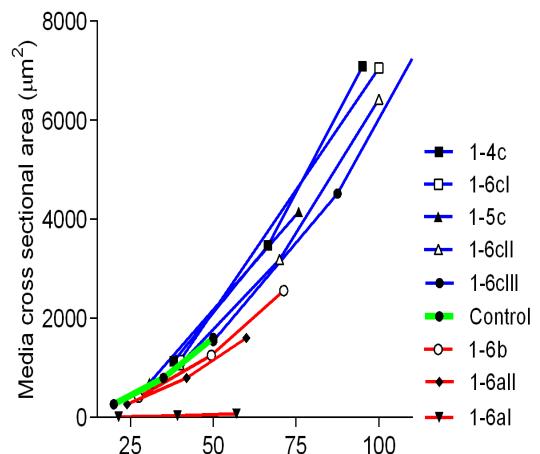
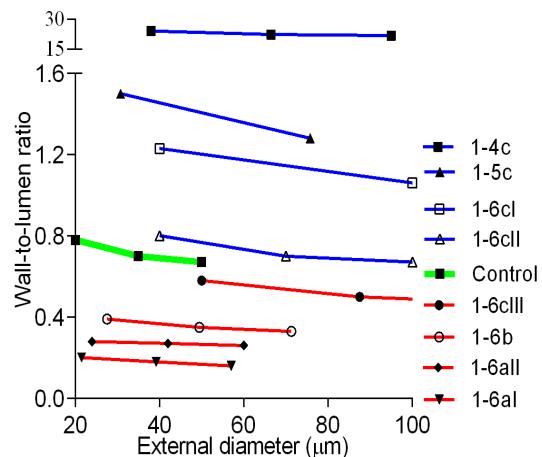
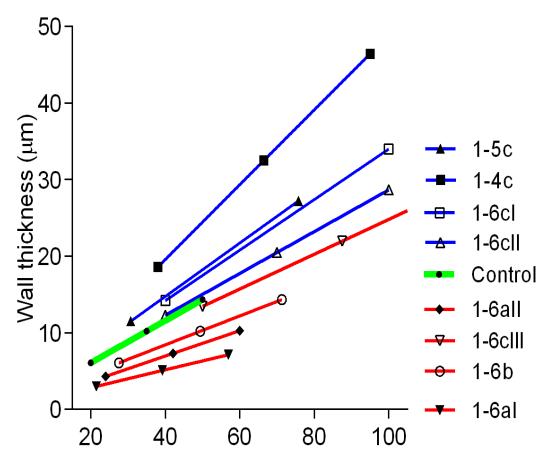
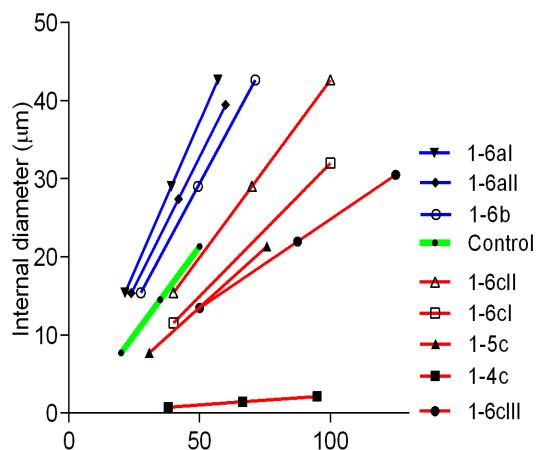


Fig 19.

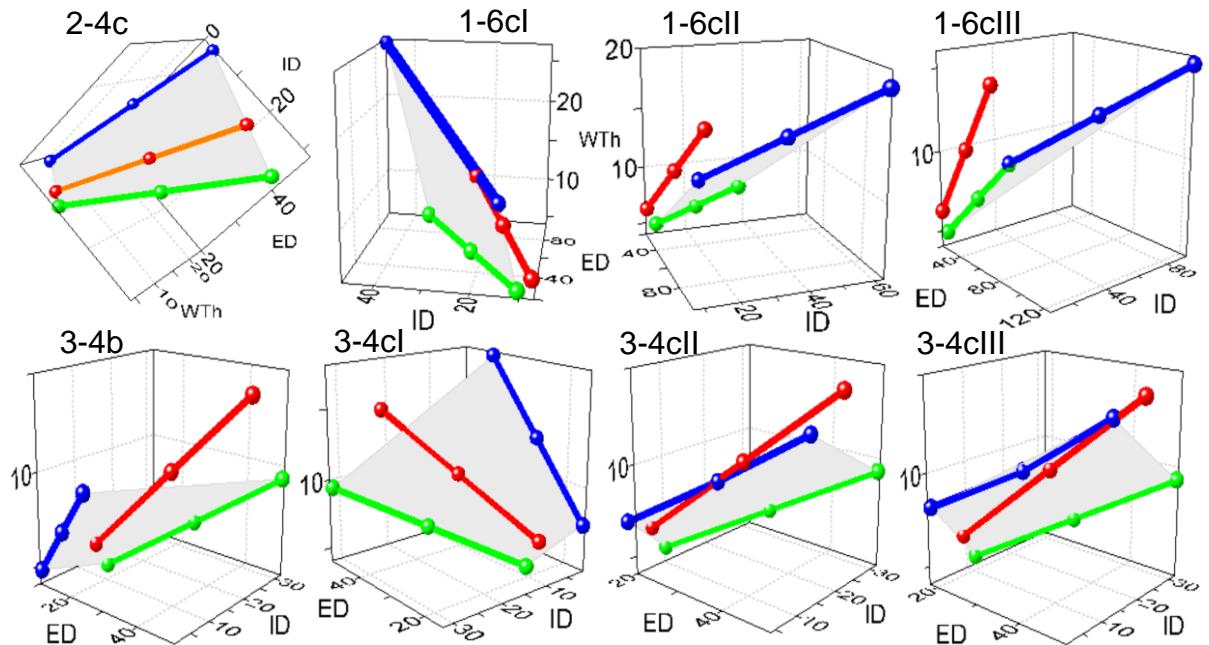


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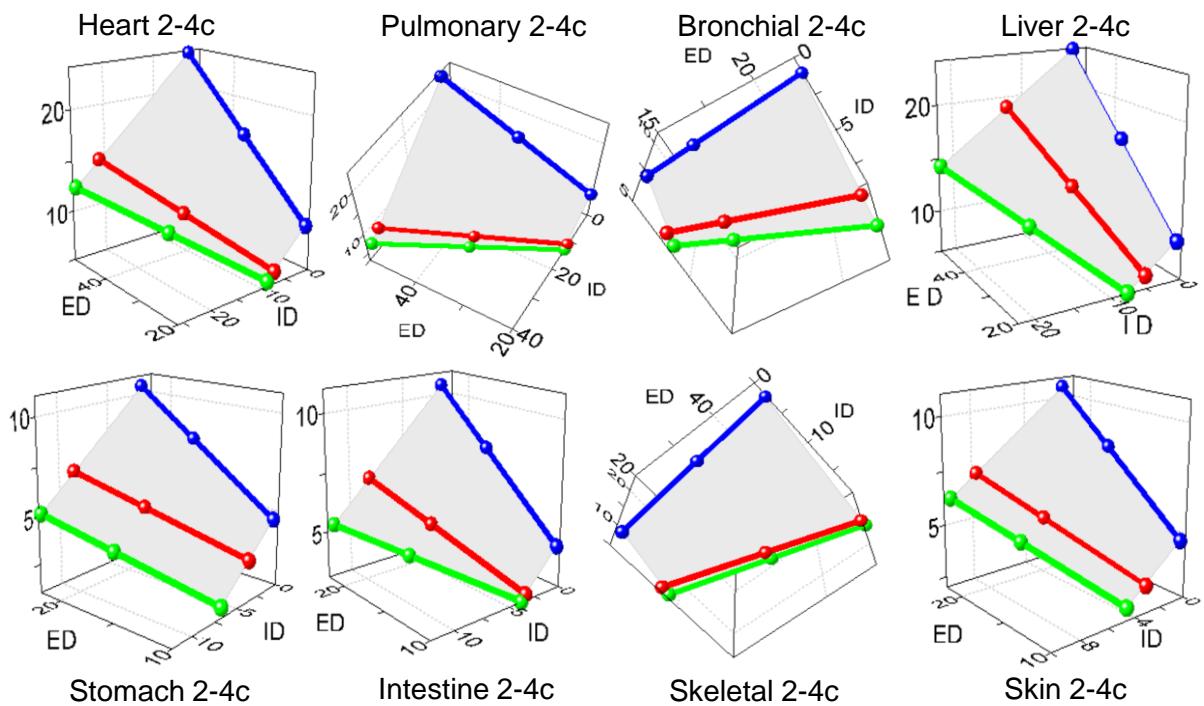


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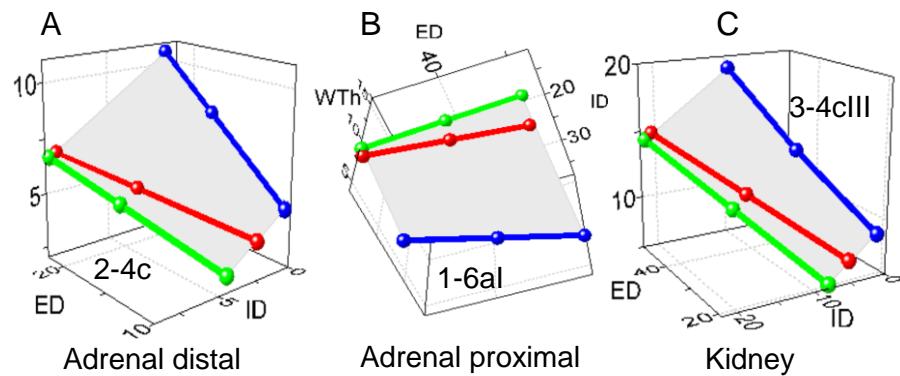


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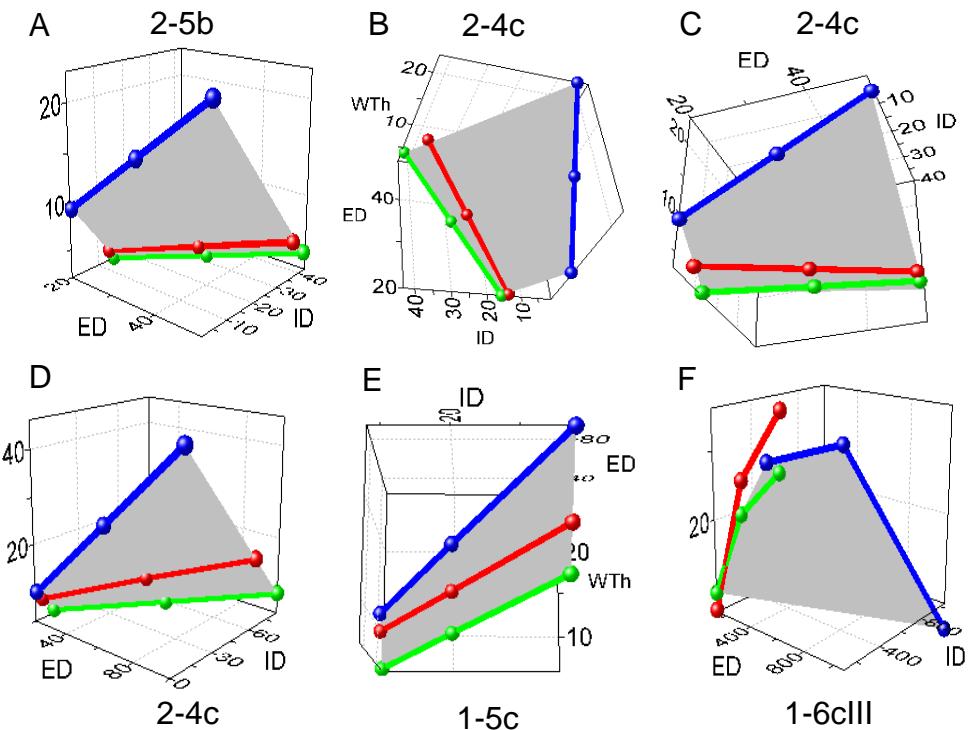


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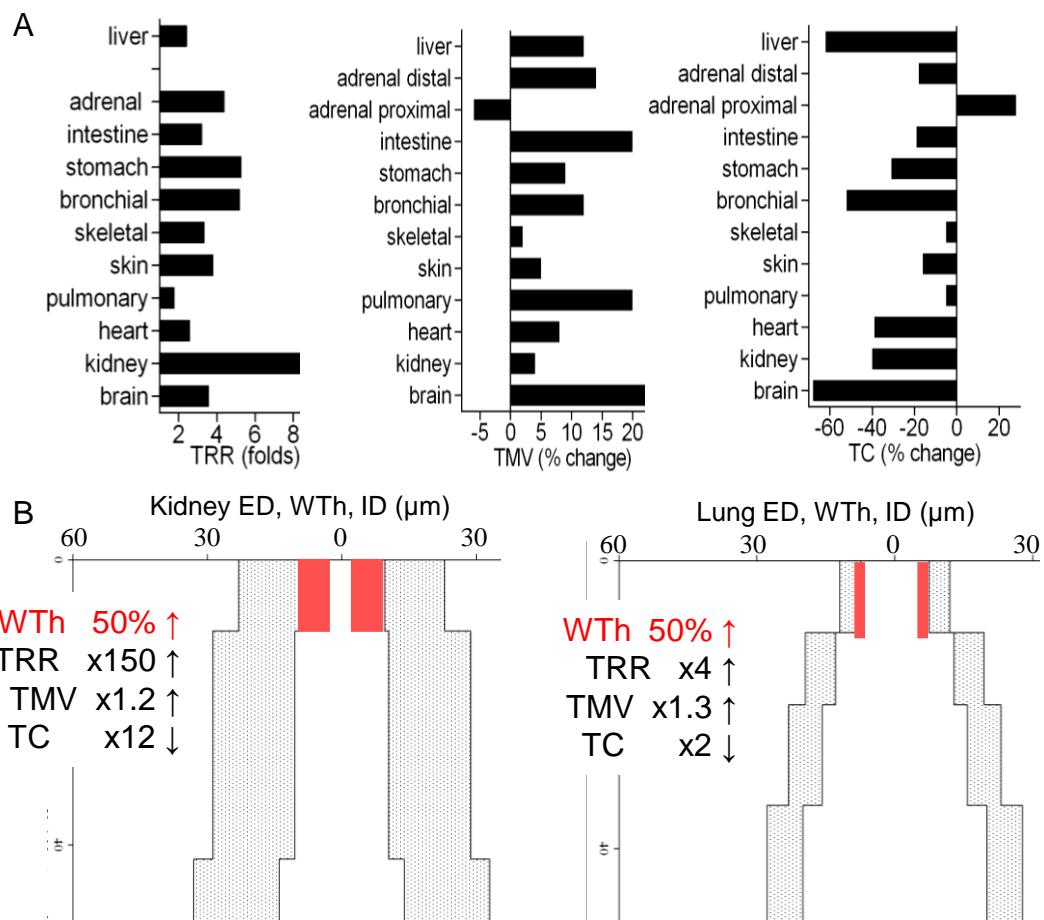


Fig 24.