Triamcinolone Acetonide Selectively Inhibits Angiogenesis in Small Blood Vessels and Decreases Vessel Diameter within the Vascular Tree

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ABSTRACT

BACKGROUND. The steroid triamcinolone acetonide (TA) is a potent anti-angiogenesis drug used to treat retinal vascular diseases that include diabetic retinopathy, vascular occlusions and choroidal neovascularization.

PURPOSE. To quantify the effects of TA on branching morphology within the angiogenic microvascular tree of the chorioallantoic membrane (CAM) of quail embryos.

METHODS. Increasing concentrations of TA (0–16 ng/ml) were applied topically on embryonic day 7 (E7) to the chorioallantoic membrane (CAM) of quail embryos cultured in Petri dishes, and incubated for an additional 24 or 48 hours until fixation. Binary (black/white) microscopic images of arterial end points were quantified by VESGEN software (for Generational Analysis of Vessel Branching) to obtain major vascular parameters that include vessel diameter ($D_v$), fractal dimension ($D_f$), tortuosity ($T_v$) and densities of vessel area, length, number and branch point ($A_v$, $L_v$, $N_v$ and $Br_v$). For assessment of specific changes in vascular morphology induced by TA, the VESGEN software automatically segmented the vascular tree into branching generations ($G_1$–$G_{10}$) according to changes in vessel diameter and branching.

RESULTS. Vessel density decreased significantly up to 34% as the function of increasing concentration of TA according to $A_v$, $L_v$, $Br_v$, $N_v$ and $D_v$. TA selectively inhibited the growth of new, small vessels, because $L_v$ decreased from 13.14 ± 0.61 cm/cm$^2$ for controls to 8.012 ± 0.82 cm/cm$^2$ at 16 ng TA/ml in smaller branching generations ($G_7$–$G_{10}$), and for $N_v$ from 473.83 ± 29.85 cm$^2$ to 302.32 ± 33.09 cm$^2$. In contrast, vessel diameter ($D_v$) decreased throughout the vascular tree ($G_1$–$G_{10}$).

Key Words: triamcinolone acetonide; angiogenesis; retinal neovascularization; fractal; chorioallantoic membrane; CAM; quail; avian.
INTRODUCTION

The steroid triamcinolone acetonide (TA) is a potent anti-inflammatory and anti-angiogenic drug used for treatment of macular edema secondary to retinal vascular diseases including diabetic retinopathy and retinal vascular occlusions, as well as neovascularization including choroidal neovascularization (CNV) and retinal neovascularization in inflammatory diseases. It is believed that corticosteroids decrease levels of retinal thickening and improve macular edema by several mechanisms. TA decreases inflammatory mediators that include tumor necrosis factor, interleukin 5, interleukin 6, interleukin 8, prostaglandins and interferon-gamma.\textsuperscript{1-3} TA modulates cellular calcium levels by interacting with both voltage-dependent calcium channels and calmodulin.\textsuperscript{4, 5} These activities may promote more effective fluid resorption, resulting in decreased macular edema. Corticosteroids also decrease levels of the potent permeability factor, vascular endothelial growth factor (VEGF).\textsuperscript{6, 7} Finally, TA improves the integrity of the blood-retinal barrier.\textsuperscript{8} Regardless of the mechanisms of activity, TA has been documented to significantly improve vision and decrease retinal edema.\textsuperscript{9-12}

Although the clinical effect of TA in these diseases is established, the effect of TA on vascular morphology is not well understood. Improved understanding of how TA affects angiogenesis and vascular morphology would be helpful for therapeutic optimization. For this study, site-specific changes within the vascular tree induced by TA were quantified using an \textit{in vivo} model of angiogenesis.

For evaluation of the effects of TA on the angiogenic vascular tree, the quail chorioallantoic membrane (CAM) is a highly useful model during mid-development, when the rate of CAM angiogenesis is at a maximum.\textsuperscript{13-17} As described previously, the complex spatial patterns of the branching vascular tree and the associated capillary network can be easily visualized by light and fluorescence/confocal microscopy.\textsuperscript{13-17} Using fractal analysis, this
vascular pattern can be precisely analyzed. Fractal analysis is a recent non-Euclidean mathematical innovation\textsuperscript{18} that quantifies the space-filling patterns of complex objects. Fractal geometry is common in nature and includes botanical and vascular trees, snowflakes, coastline topography, and even the spatiotemporal scaling of vascular-based physiological metabolism.\textsuperscript{19} As a fractional, non-integral number that increases according to the increasing density of a space-filling pattern, the fractal dimension (\(D_f\)) is statistically sensitive to small, early-stage changes in the vascular tree.\textsuperscript{13-15, 17, 21} A fractal object typically reaches its greatest space-filling capacity using self-similarity, the geometric property by which a pattern such as vascular bifurcational branching is repeated iteratively at continuously decreasing length scales.

The computer program VESGEN (abbreviated from Generational Analysis of Vessel Branching) was developed by NASA as a fully automated, user-interactive program that quantifies major vascular branching parameters using a single, user-provided image of 2D vascular pattern. The fractal-based VESGEN analysis segments vessels within vascular trees into branching generations (\(G_1, G_2, \ldots G_x\)) according to changes in vessel diameter and branching. Site-specific changes within the vascular tree induced by angiogenic cytokines or other molecular regulators such as TA can then quantified.\textsuperscript{14-16} Thus, the purpose of this study was to evaluate the effects of the TA on branching morphology within the microvascular tree of the quail CAM model using fractal-based VESGEN analysis.
MATERIALS AND METHODS

Embryonic culture, assay, mounting, imaging, and fractal/VESGEN branching analysis used in this study have been described previously, and are summarized below.

Culture, Assay and Mounting

Fertilized eggs of Japanese quail (Coturnix coturnix japonica, Boyd’s Bird Co., Pullman, WA) were incubated at 37.6 ± 0.2 °C under ambient atmosphere, cracked at embryonic day three (E3, following incubation of eggs for 56 hours) and cultured further in 6-well petri dishes (cross-sectional area = 10 cm²). Quail egg culture and experimental protocols were in accord with NIH guidelines and approved by the Chief Veterinarian Officer of NASA (Ames Research Center). At E7 (following incubation for an additional 96 h), prewarmed PBS solution containing TA (0.5 ml at 0-16 ng/ml) was applied dropwise to the surface of each CAM. Sterile-filtered stock solutions of TA (Sigma, T-6501) were prepared at 10 mg/ml in 100% ethanol. Additional concentrations of TA were tested to determine the range of dose effectiveness. The total amount of the molecular regulator, rather than its concentration, is the governing parameter because solutions are quickly absorbed into CAM tissue. Quantities of TA are therefore reported as 0-8 ng/CAM. Following treatment with TA and further incubation for 24 or 48 hours, the embryos (with CAMs) were fixed in 4% paraformaldehyde/2% glutaraldehyde/PBS for several days prior to dissection and mounting for microscopic analysis.

Imaging

Aldehyde fixation of the CAM results in high contrast of the arterial tree due to retention of erythrocytes (or red blood cells, RBC) within arteries, but low contrast of the venous tree resulting from evacuation of RBC from veins during dissection (Fig. 1). Digital images (1392 × 1040 pixels) of (terminal) arterial end-point vessels from the middle region of the CAM were acquired in
grayscale (0-255 intensity) at total 12.5X magnification and resolution of 7.32 µm/pixel (Leica DM4000B microscope attached to Retiga EXi CCD camera, Qimaging, by Image Pro Plus software). Previous studies showed that the CAM arterial end-point regions analyzed by us are representative of changes induced by topical application of angiogenesis regulators throughout the CAM arterial trees, and there is no significant increase in vascular density at a total magnification of 20X in comparison to 12.5X. Grayscale images were converted into binary (black/white) images of vascular morphology (Fig. 1) by semi-automatic computer processing using Adobe Photoshop 7.0 and NIH ImageJ software (http://rsb.info.nih.gov/ij/). The accuracy of vascular image binarization was confirmed by a second independent, experienced operator.

Vascular Quantification

Statistical Sampling

Thirty-nine representative CAM specimens were quantified from three independent experiments, including at least three controls from each experiment (a total of eleven controls), and seven specimens for each of the four concentrations of TA. Additional specimens and experiments served as qualitative confirmation of the quantified results. Variation was assessed by calculating the standard error (S.E.). P-values were obtained for the four treatment groups (1-8 ng TA/CAM) compared to control (0 ng TA/CAM) by two-tailed, heteroscedastic Student’s t-test.

Analysis by VESGEN

The NASA Glenn computer code VESGEN (Fig. 2) was used to measure parameters of vascular morphology that include vessel length density (L_v), vessel area density (A_v), vessel branch point density (Br_v), vessel number density (N_v), vessel tortuosity (T_v) and vessel diameter (D_v) for each branching generation G_1 through G_{10}. For example, D_{v1-2} denotes D_v with respect to branching generations G_1-G_2. By VESGEN analysis, only one image was found to contain 11 branching generations (i.e., a few vessels of G_{11}), which were therefore merged into image results for G_{7-10}. 
(G<sub>7</sub>). L<sub>v</sub>, A<sub>v</sub>, N<sub>v</sub> and Br<sub>v</sub> were expressed as density functions by normalization to the area of the image containing the major arterial tree extracted as region of interest (ROI) or (2) the entire image (Fig. 2). Vessel diameter was calculated as D<sub>v</sub> = A<sub>v</sub>/L<sub>v</sub>. A trimmed skeleton was used to obtain accurate measurements for L<sub>v</sub> in specific branching generations such as L<sub>v1</sub>.<sup>16</sup> Tortuosity (T<sub>v</sub>) was estimated by the ratio of the length of a trimmed vessel (L<sub>v</sub>) determined by VESGEN to the shortest distance between the vessel endpoints.

Vessel branching generations (G<sub>1</sub>-G<sub>X</sub>) are determined by VESGEN according to relative decreases in vessel diameter, as first established for branching vascular trees in the dog and pig heart and lung.<sup>22</sup>-<sup>24</sup> Blood flow is conserved at a symmetric vessel bifurcation when the diameter of a symmetric offspring vessel decreases to 71% (1/square root of 2) of the parent vessel diameter, and therefore the decrease of vessel diameter to 71% was used as the primary determinant of a new branching generation. As can be seen in biological branching trees (Fig. 2), however, the branching of relatively symmetric offspring vessels is not perfectly symmetric, and the diameters of very few offspring vessels are of the 71% ideal value. In addition, vessels tend to taper. To accommodate a range of vessel diameters within a branching generation, VESGEN therefore contains a 15% default tolerance factor that is user-adjustable. For the TA study, the tolerance factor was left at ± 15%. For a very small number of vessel segments (only 5 vessel segments out of 39 total images), the automatic segmentation by VESGEN was incorrect according to the established criteria of generational classification based on vessel diameter and branching. The user-interactive features of VESGEN were therefore used to override and correct these few inaccuracies. A further consideration is that the most frequent branching event in a vascular tree is the asymmetric offshoot branching of a much smaller vessel from a larger vessel, presumably due to space-filling requirements of vascular branching (Fig. 2). It is important to note that although the branching pattern of each vascular tree in a CAM or a human retina is unique, the space-filling properties of the vascular trees are remarkably uniform.<sup>13</sup>,<sup>21</sup>
VESGEN now functions as a fully automated, Java-based software operating as a plug-in to NIH ImageJ. A binary vascular image is the single input required by VESGEN to quantify vascular trees, networks or tree-network composites of highly 2D tissues such as the avian CAM, human retina and rodent retina. VESGEN will be publicly available in the near future, and its full capabilities are being described elsewhere.

Fractal Analysis

To support fractal analysis by the box-counting algorithm, each binary image was rescaled slightly to 1370 x 1024. A left- and right-most square image of 1024 x 1024 was extracted from each binary image and skeletonized (i.e., linearized, Fig. 1) using the NIH ImageJ skeletonizing algorithm. The fractal dimension ($D_f$) was calculated for binary and skeletonized images by implementation of box-counting at a power of 2 using ImageJ. Values of $D_f$ for the left and right 1024 x 1024 images were averaged to obtain an overall $D_f$ for each original image. The fractal box-counting algorithm has since been incorporated into VESGEN to confirm the TA fractal results and for use in future studies. We consider VESGEN to be a fractal-based analysis of vascular trees due to the complex, non-Euclidean space-filling geometry of the vascular branching structures, and VESGEN assignment of vascular parameters to specific, self-similar generations of vascular branching.
RESULTS

Topical application of TA significantly inhibited ongoing angiogenesis in the quail CAM after 24 hours by two major morphological mechanisms: (1) vascular density was inhibited by a targeted decrease in the number of small blood vessels, and (2) decreased vessel diameter throughout the vascular tree. Vascular density decreased as a function of increasing concentration of TA (Fig. 3) by several confirming measures of the entire vascular field in binary and skeletonized images. Skeletonized images are direct representations of vessel density that illustrate the extensive space-filling properties and overall vessel connectivity of a branching vascular tree. Visual inspection of vascular pattern further confirms these results (Figs. 1, 4). By microscopic observation, decreased vascular density and alterations in vessel diameter induced by TA at 4-8 ng/CAM persisted after 48 hours (results not shown).

Within skeletonized images, \( L_v \) and \( Br_v \) decreased up to 23% and 34%, respectively (Fig. 3). \( D_i, L_v \) and \( Br_v \) decreased from 1.406 ± 0.004, 25.7 ± 0.6 (cm/cm²), and 454 ± 28 (cm⁻²) in controls to 1.355 ± 0.008, 19.5 ± 0.9 (cm/cm²), and 301 ± 27 (cm⁻²) in specimens treated at the maximum concentration of 8 ng TA/CAM (\( p \) values = 3xE-4, 1xE-4 and 0.001, respectively). In binary images, \( D_i \) and \( A_v \) decreased from 1.669 ± 0.006 and 0.14 ± 0.00 (cm²/cm²) in controls to 1.623 ± 0.009 and 0.11 ± 0.005 (cm²/cm²) at 8 ng TA/CAM (\( p = 0.0015 \) and 6xE-4). Absolute differences of \( D_i \) in binary and skeletonized images may appear small. However, as a non-Euclidean, 'fractional' measure of space-filling patterns, \( D_i \) is restricted in 2D, black/white images to fractional values between 1 and 2. \( D_i \) is a sensitive, statistically significant, reproducible measure of space-filling vascular density that ranges from approximately 1.34 to 1.55 in skeletonized images and 1.61 to 1.75 in binary images.¹³⁻¹⁵, ¹⁷, ²¹

Decreases in vascular density as measured by \( D_i \) and \( A_v \) in binary images can result from several morphological changes that include decreased vessel length and number of vessels, and/or decreased vessel diameter (see Fig. 1B, E). Quantitative analysis by VESGEN confirmed visual
observations that the morphological mechanisms of decreased vascular density induced by TA include: (1) decreased number densities restricted to smaller vessels, and (2) overall thinning of vessel diameter. Vessel tortuosity as measured by $T_v$ was unaffected, varying typically between 1.04 and 1.17 within a specimen. By $N_v$ and $L_v$ (Fig. 5), vessel density decreased significantly for the smallest vessels of $G_7$-$G_{10}$, but not for large and medium-sized vessels of $G_1$-$G_2$, $G_3$-$G_4$ and $G_5$-$G_6$. $N_{v7-10}$ decreased significantly from $474 \pm 30 \text{ cm}^{-2}$ in controls to $302 \pm 33 \text{ cm}^{-2}$ at 8 ng TA/CAM ($p = 0.0017$), whereas $N_{v1-2}$, $N_{v3-4}$ and $N_{v5-6}$ remained relatively constant (see Fig. 5; $p = 0.69$, 0.53 and 0.61, respectively). For example, $N_{v1-2}$ was $5.04 \pm 0.32 \text{ cm}^{-2}$ in controls compared to $5.15 \pm 0.05 \text{ cm}^{-2}$ at 8 ng TA/CAM. Small but consistent decreases in vessel diameter ($D_v$) were induced by TA throughout the vascular tree (Fig. 6). $D_{v7-10}$ decreased from $28.1 \pm 1.0 \mu\text{m}$ in controls to $25.5 \pm 0.4 \mu\text{m}$ at 8 ng TA/CAM ($p = 0.03$), $D_{v5-6}$ from $61.1 \pm 3.1 \mu\text{m}$ to $53.0 \pm 3.1 \mu\text{m}$ ($p = 0.08$), $D_{v3-4}$ from $118.1 \pm 6.0 \mu\text{m}$ to $101.6 \pm 5.8 \mu\text{m}$ ($p = 0.07$), and $D_{v1-2}$ from $228.4 \pm 12.1 \mu\text{m}$ to $199.0 \pm 11.2 \mu\text{m}$ ($p = 0.09$).

A statistical study of the effects of vessel generational branching on our results confirmed the value and validity of site-specific VESGEN analysis. The binning (i.e., lumping or merging) of $G_1$-$G_{10}$ output parameters demonstrated increasingly strong statistical confidence resulting from increasingly fine binning of the generational data. For example, measurements of $D_v$ using a two-fold binning of generations for controls and 8 ng TA/CAM into larger vessels of $D_{v1-6}$ and smaller vessels of $D_{v7-10}$ (this group identical to results cited above) yielded a $p$-value of relatively insignificant difference for the larger vessels ($p = 0.15$). Similarly, for a three-fold binning of generations into $D_{v1-3}$, $D_{v4-6}$, and $D_{v7-10}$, $p$-values for the two groups of larger vessels were also inconclusive ($p = 0.25$ and $p = 0.15$, respectively). As described above, a finer four-fold binning of $D_{v1-2}$, $D_{v3-4}$, $D_{v5-6}$, and $D_{v7-10}$ showed considerably higher levels of statistical significance for trends ($p \leq 0.10$) and confidence intervals ($p \leq 0.05$). Nonetheless, some binning of generational results as
performed for this study improves and smooths the data due to increased statistical sampling\textsuperscript{14, 17} and provides greater clarity in presentation of results.
DISCUSSION

Topical treatment with the corticosteroid triamcinolone acetonide (TA) resulted in multimodal changes to the vascular tree in the quail CAM, as quantified by VESGEN analysis. The angiogenesis of small blood vessels was selectively inhibited by TA, although vessel diameter decreased throughout the branching tree. Other critical aspects of vessel morphology remained normal following treatment by TA. For example, the density and number of larger vessels were unaffected, the vascular tree appeared to taper smoothly, and vessel tortuosity remained normal. It is not surprising that TA, as a potent angiogenesis inhibitor, selectively decreased the density of only small, new blood vessels within the vascular tree. It is possible that TA decreased overall vessel diameter by inhibiting VEGF activity, because VEGF stimulates increases in vessel diameter (particularly of larger vessels) as well as in vessel density.

In this study, TA was applied topically to the quail chorioallantoic membrane during mid-embryonic development when angiogenesis is occurring at its maximum rate, and angiogenic cytokines and regulators can be applied easily and uniformly in solution. The transparent CAM membrane develops rapidly, is highly vascularized, and is essentially 2D. Complex spatial patterns of the branching vascular tree and associated capillary network are easily visualized by light and fluorescence/confocal microscopy, and quantified by fractal-based VESGEN analysis. As a relatively convenient experimental model, the angiogenic CAM exhibits some useful morphological and functional similarities to angiogenic diseases of the quasi-2D retinal vascular tree. For example, region-based fractal methods developed in the CAM were successfully extended to the quantification of progression in diabetic vascular disease using clinical images of the human retina.

We are using the computer software VESGEN (for Generational Analysis of Vessel Branching) to quantify major vessel parameters of blood and lymphatic vascular remodeling.
Output parameters of VESGEN include vessel diameter, tortuosity, fractal dimension, and densities of vessel number, branch point, length and area. Most parameters can be obtained by the user for the overall vascular image, the major vascular tree, and for individual or merged branching generations. Vascular trees are decomposed into branching generations so that cytokine- or therapeutic-regulated modifications can be quantified according to site-specific vessel location. For this study on TA, we chose to analyze generational branching within the major vascular tree. Focusing on branching relationships within a single tree can support precise conclusions about regulator-induced changes in vessel branching relationships. This is the first technical report of results generated by the fully mature, newly automated VESGEN software (version 1.0) that now analyzes 2D vascular trees, networks and tree-network composites for a number of experimental and clinical tissue applications in angiogenesis and lymphangiogenesis, including the avian CAM and yolksac, the human retina, rodent retina, and developing coronary vessels in the embryonic heart.

As demonstrated by VESGEN analysis, it now appears that perturbation of rapid, ongoing angiogenesis during the middle stages of CAM development by TA occurs primarily by the selective inhibition or stimulation of new, small blood vessels. As a generalized result for the CAM model, this conclusion appears important although not particularly surprising, because the molecular and cellular characteristics of angiogenic vascular tissues differ from those of more mature, stable vascular tissues.\textsuperscript{25, 26} Angiogenic perturbants quantified to date in this fractal-based CAM model include the inhibitors TA, transforming growth factor β-1 (TGF-β1)\textsuperscript{14} and angiostatin,\textsuperscript{13} and stimulators basic fibroblast growth factor (bFGF)\textsuperscript{15} and vascular endothelial growth factor-165 (VEGF\textsubscript{165}).\textsuperscript{17} Additional regulator-specific effects on vascular morphology such as overall vessel thinning by TA or thickening of larger blood vessels by VEGF\textsubscript{165} have also been quantified. As for TA, morphological response of the vascular tree to VEGF\textsubscript{165} measured by VESGEN was multimodal. Increased vessel density and increased vessel diameter reached
maximal frequencies at lower and higher VEGF concentrations, respectively. The study of TGF-β1 in particular considered tissue growth (rescaling) of the entire CAM vascular tree. Each molecular perturbant of angiogenesis has therefore elicited a unique 'fingerprint' response that is spatiotemporally distinct and quantifiable, despite the apparent generality of inhibition and stimulation of angiogenesis in the CAM at the level of new, small vessels within the growing vascular tree.
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**Fig. 1. Image analysis.**

Representative images of arterial end-point regions of CAM specimens treated with PBS control vehicle (A) or with corticosteroid triamcinolone acetonide (TA, D) were acquired by brightfield microscopy. Aldehyde fixation of the CAM results in retention of RBCs within arterial vessels (arrows) and extraction of large amounts of blood from venous vessels (arrowheads). Arterial trees are thereby conveniently separated from overlapping venous trees to support the semi-
automatic processing of grayscale images into (B, E) binary (black/white) vascular patterns. Binary images and (C, F) corresponding skeletonized images clearly reveal strong inhibition of vessel density by TA. Measurements of decreased vessel density at 8 ng TA/CAM (E, F) relative to control (B, C) include: (E) vessel area density ($A_v$) = 0.122 (cm$^2$/cm$^2$) and fractal dimension ($D_f$) = 1.656, compared to (B) $A_v$ = 0.152 (cm$^2$/cm$^2$) and $D_f$ = 1.683; (F) vessel length density ($L_v$) = 20 (cm/cm$^2$), vessel branch-point density ($Br_v$) = 325 (cm$^{-2}$) and $D_f$ = 1.360 compared to (C) $L_v$ = 26 (cm/cm$^2$), $Br_v$ = 443 (cm$^{-2}$) and $D_f$ = 1.410.
**Fig. 2. Assignment of vessels to branching generations $G_1$-$G_{10}$ by VESGEN.**

The VESGEN output image of a CAM control specimen illustrates the classification of vessels into successively smaller branching generations. Ten branching generations ($G_1$-$G_{10}$) of vascular branching were measured by VESGEN for the arterial end-point region of this control CAM specimen. Vessel branching generations are determined by (1) *decrease in vessel diameter* and (2) *vessel bifurcations that are approximately symmetric* (i.e., when diameters of offspring vessels branching from a parent vessel are approximately equal). The major arterial tree and its corresponding region of interest (ROI, in black) are also identified by VESGEN. The edge of the ROI lies midway between the end points of the major arterial tree and neighboring arteries. In
regions where vessels of the arterial tree extend beyond the edge of the image, the ROI is defined simply by the edge of the image. As seen above, the most frequent branching event within the vascular tree is asymmetric branching, or branching of a small offshoot vessel from a much larger vessel. In this ROI, 35 symmetric branch points and 209 asymmetric branch points were measured by VESGEN.

**Fig. 3. Vessel density decreases as the function of increasing concentration of TA.**

As a measure of the space-filling capacity of a vascular pattern, the fractal dimension ($D_f$) decreased significantly with increasing TA concentration in (A) binary and (B) skeletonized images throughout the entire vascular field, as illustrated in Fig. 1. Vessel density also decreased according to other measures of vessel density that include (C) vessel length density
(L_v) and (D) vessel branch point density (Br_v). Data are plotted as mean ± S.E. P-values ≤ 0.05 and ≤ 0.01 are indicated by * and **, respectively.

**Fig. 4. TA selectively decreases the number of smaller vessels and decreases vessel diameter throughout the vascular tree.**

(A-C) Each representative image displays the major arterial tree with its ROI (in black) as determined by VESGEN, for which vessel diameter (D_v) and the density of key vascular parameters that include vessel number density (N_v), vessel branch point density (Br_v), vessel length density (L_v), and vessel area density (A_v), were measured. Parameters were specified for branching generations G_1-G_{10} (see colorized legend) and denoted D_{v1}, A_{v1}, L_{v1}, N_{v1}, etc. Relative to controls (A), VESGEN results for A_v, L_v, N_v and Br_v in TA-treated specimens (B-C) show that vessel density decreased only in the smallest branching generations G_7-G_{10} (Fig. 5). However, D_v decreased throughout vascular tree (Fig. 6). B above is the VESGEN output image of the input binary image Fig. 1E.
**Fig. 5.** TA selectively inhibits the angiogenesis of small blood vessels.

Relative to controls, vessel density estimated by vessel number density ($N_v$) and vessel length density ($L_v$) decreased only in the smallest branching generations $G_7$-$G_{10}$ for higher concentrations of TA. For clarity, results for various classes of vessels lumped together as $G_{1-2}$, $G_{3-4}$, $G_{5-6}$ and $G_{7-10}$. See Figs. 1, 2 and 4 for qualitative confirmation of the quantitative results. Data are plotted as mean ± S.E. $P$-values ≤ 0.05 and ≤ 0.01 are indicated by * and **.
Fig. 6. TA decreases vessel diameter throughout the vascular tree.

Vessel diameter ($D_v$) decreased in all vessel branching generations ($G_1$–$G_{10}$) with increasing concentration of TA, relative to controls. As for Fig. 5, results were lumped together as $G_{1-2}$, $G_{3-4}$,
G_5-6 and G_7 (G_7-10). For qualitative confirmation of quantitative results, see Figs. 1 and 4. Data are plotted as mean ± S.E. P-values ≤ 0.05 and ≤ 0.01 are indicated by * and **.
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