

In vivo characterization of spontaneous microhemorrhage formation in mice with cerebral amyloid angiopathy

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Abstract

The pathophysiology of microhemorrhages in the context of cerebral amyloid angiopathy (CAA) remains poorly understood. Here we used in vivo two-photon microscopy in aged APP/PS1 mice with mild-to-moderate CAA to assess the formation of microhemorrhages and their spatial relationship with vascular A β depositions in the surrounding microvascular network. Mice with chronic cranial windows were intravenously injected with fluorescent dextran to visualize the vessels and a fluorescently labeled anti-fibrin antibody to visualize microhemorrhages. Focal vessel irradiations resulted in extravascular fibrin-positive clots at individual rupture sites that remained visible for weeks. Spontaneous extravascular fibrin-positive clots were more often observed in 19-month-old transgenic APP/PS1 mice compared to their wild-type littermate controls ($p = 0.039$), after heparin administration. In the transgenic mice, these spontaneous leakage sites frequently occurred at arteriolar segments without CAA at bifurcations or vessel bends. These findings suggest that the presence of vascular A β per se does not directly predispose vessels to leak, but that complex flow dynamics within CAA-affected vascular networks likely play a role. Our in vivo approach for the detection of individual spontaneous leakage sites may be used in longitudinal studies aimed to assess structural and functional alterations at the single-vessel level leading up to microhemorrhage formation.

Keywords

Blood–brain barrier, cerebral amyloid angiopathy, fibrin, in vivo microscopy, microbleeds

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Introduction

Cerebral small vessel disease (SVD) refers to pathological alterations of the small arteries and venules of the brain and is a major contributor to age-related cognitive decline and dementia.¹ Whereas the small vessels escape detection by conventional magnetic resonance imaging (MRI), manifestations of SVD can be readily detected during life and include white matter hyperintensities, infarcts, and microhemorrhages.² Advancements in the development and optimization of blood-sensitive MRI sequences have resulted in increased detection of microhemorrhages,^{3,4} which have been linked to a higher risk of stroke, mortality, and dementia.⁵ Regarding their pathophysiology, microhemorrhages in deep locations of the brain have been associated with hypertensive arteriopathy,

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whereas cortical microhemorrhages are indicative of cerebral amyloid angiopathy (CAA).⁶ CAA is characterized by the accumulation of amyloid β (A β) in the walls of small cortical arterioles and leptomeningeal vessels and increases the risk of large bleeds that are often fatal in patients with the disease.⁷ Although neuropathological examinations have confirmed a positive association between cortical microhemorrhages and severe CAA,^{8,9} the underlying mechanisms and sequence of events that lead to microhemorrhage formation are poorly understood. This is mainly due to a lack of relevant experimental animal models and the fact that microhemorrhages easily escape detection upon neuropathological examination. Moreover, microscopic observations in the brain post-mortem are cross-sectional in nature and do not allow to assess directly causes and consequences of bleeding. Furthermore, histopathology is often limited by the assessment of single thin brain sections, which does not allow studying lesions in relation to the surrounding microvascular network. Therefore, we utilized in vivo two-photon microscopy in a mouse model of A β overproduction with mild-to-moderate CAA and developed a novel approach to assess the formation of microhemorrhages as well as their exact sites of leakage in real-time in 3D and their spatial relationship with vascular A β depositions in the surrounding microvascular network.

Material and methods

Animals

Seven-to-nine months old and 18-to-20 months old male and female transgenic (Tg) APP^{swc}/PSEN1^{dE9} (APP/PS1) mice and their wild-type (WT) littermates were included in this study. APP/PS1 mice express the human mutant amyloid precursor protein (APP) gene containing the Swedish K594N/M595L mutation and the human presenilin 1 gene lacking exon 9, under control of the prion protein promoter.¹⁰ Mice were purchased from The Jackson Laboratory (stock nr. 34829) and bred in our animal facility on a C57BL/6J background. While aging, mice were housed up to four individuals in one cage on a 12-h light/dark cycle with food and water provided ad libitum. After surgery, mice were housed individually. A subset of a separate set of naïve mice (10/12) used for ex vivo histopathology carried a Gad2-Cre transgene, unrelated to the current study.

All animal procedures were performed with the approval of the Massachusetts General Hospital Animal Care and Use Committee and experiments were conducted in compliance with the National Institutes of Health Guide for the Care and Use of

Laboratory Animals. The ARRIVE guidelines were followed for reporting animal experiments.

Ex vivo histopathology

Naïve 20-month-old mice were euthanized by CO₂ asphyxiation and transcardially perfused with 20 mL phosphate-buffered saline (PBS). Brains were extracted and stored at 4°C in a fixation solution containing 4% paraformaldehyde and 15% glycerol, in PBS. Next, the intact brains were embedded in paraffin in coronal orientation and cut in 6 μ m-thick sections, as follows: throughout the brain, four adjacent sections were cut skipping 100 μ m between each set of four sections. The olfactory bulbs and cerebellum were excluded. The first section at each level of the brain was stained with hematoxylin and eosin (H&E) to assess general morphology, and the second section with Perls' Prussian blue to detect Iron-positive deposits, both following standard protocols. A β immunohistochemistry was performed on a single section per brain to confirm the genotype of each mouse.

Surgical procedures

Cranial window surgeries were performed as described previously, with minor modifications.^{11,12} Mice were anesthetized with 5% isoflurane (mixed with pure O₂) and placed in a stereotactic frame. Throughout the procedure, isoflurane levels were maintained at ~2% and adjusted if necessary. After shaving and partial removal of the scalp, a craniotomy (measuring 5 mm in diameter) was performed exposing both hemispheres at the level of the somatosensory cortices. A cover glass (measuring 8 mm in diameter) with sterile PBS was placed over the brain and secured in place with a mixture of dental cement and Crazy Glue. Post-surgically, mice received a single subcutaneous injection of (0.05 mg/kg) buprenorphine and (300 mg/mL) Tylenol was added to their drinking water for three days. Mice could recover from the surgery for at least three weeks before they were imaged.

In vivo two-photon microscopy

Imaging was performed using the commercially available FluoView FV1000MPE two-photon laser-scanning system mounted on a BX61WI microscope, equipped with a 25 \times (numerical aperture 1.05) dipping water immersion objective (Olympus). A Mai Tai[®] DeepSee[™] Ti:Sapphire mode-locked laser (Spectra-Physics) generated two-photon excitation at 800 nm. External detectors containing three photomultiplier tubes (PMTs, Hamamatsu) collected emitted light in the range of 420–460 nm (blue), 495–540 nm (green), and 575–630 nm (red). PMT settings were kept constant

for each imaging session, but laser power was adjusted as needed. The day prior to imaging mice were intraperitoneally injected with 300 μ L (5 mg/kg dissolved in 3% Cremophor[®] and PBS) Methoxy-X04 (Glilx lab), a blue fluorescent Congo red derivative that crosses the blood–brain barrier and labels A β depositions (both parenchymal plaques and CAA) for *in vivo* detection.¹³ Prior to imaging, mice were anesthetized with isoflurane (5% inhalation, maintained at 1.5–2% throughout the imaging session) and intravenously injected with 200 μ L (12.5 mg/mL) 70 kDa fluorescein dextran (Invitrogen) to provide an angiogram. To identify extravascular clots at leakage sites, mice also received an intravenous injection of 150 μ L (0.5 mg/mL) anti-fibrin antibody (BioMedica Diagnostics) conjugated with DyLight[®] 594 (Thermo Fisher Scientific).¹⁴

Laser irradiation experiments

To visualize microhemorrhage dynamics in real-time in 19-month-old APP/PS1 mice, arterioles and venules were irradiated to allow circulating fluorescent dextran and anti-fibrin to extravasate. Vessel irradiation was achieved by focusing the laser at maximum power in bidirectional scanning mode on a targeted arteriole or venule at 6 \times magnification for 20 s per attempt. Irradiation was immediately stopped when dextran was observed to extravasate from the irradiated site. If no extravasation was observed after six attempts, the vessel segment was excluded from further analysis. A Z-stack was acquired prior to laser irradiation at a speed of 4 μ s/pixel, resolution of 1 μ m² per pixel, matrix of 512 \times 512, and a step size of 5 μ m. Immediately following a successful irradiation, serial Z-stacks were acquired for a period of 21 min with 3-min intervals at 1 \times magnification. All mice were re-imaged again the next day and after three weeks to evaluate irradiation sites. All Tg mice received an additional intraperitoneal injection of Methoxy-X04 the day prior to the imaging session at three weeks.

Heparin experiments

The formation of spontaneous microhemorrhages was assessed in two separate cohorts of APP/PS1 mice, at 8 months and 19 months of age. First, several baseline Z-stacks were acquired in systematically sampled areas throughout the window, at a speed of 4 μ s/pixel, resolution of 1 μ m² per pixel, matrix of 512 \times 512, a step size of 5 μ m, and a magnification of 1 \times . Moreover, the window was screened visually with epifluorescence to capture any additional sites of leakage. Next, to lower the threshold for spontaneous microhemorrhages to occur, each mouse received an intravenous as well as subcutaneous injection containing (0.16 U/g) heparin

in PBS.¹⁵ The same areas were re-imaged after 15 min and the window was screened visually again to capture any additional sites of leakage. During the imaging sessions, the investigator was not blinded for genotype.

Image analysis

Acquired two-photon images were visually assessed and analyzed in FIJI.¹⁶ Arteriolar CAA burden was measured on maximum intensity projections of the Z-stacks taken prior to laser irradiation, in the (blue) Methoxy-X04 channel. For each irradiation site, the corresponding region of interest (ROI) was first outlined on the (green) dextran channel, and then copied to the (blue) Methoxy-X04 channel. After transforming the maximum intensity projections into a binary image, CAA burden was quantified for each ROI as the number of positive pixels divided by the total number of pixels within each ROI. Vessel diameters were calculated based on maximum intensity projections of the (green) dextran channel. In the heparin experiments, for each fibrin clot-containing image, if visible, the parent artery, first-order, second-order, and third-order arteriolar branches were identified at 1 \times magnification. Pial surface vessels were defined as parent arteries, penetrating arterioles as first-order branches, followed by higher order branches. A segment was considered CAA-positive if more than two distinct Methoxy-X04-positive CAA deposits were present. In addition, the distance between the edge of each fibrin clot to the nearest CAA deposit was measured.

Statistics

Graphs were created and statistical analyses performed in Graphpad Prism (version 5.03). Differences between genotype and conditions were compared using a χ^2 test for proportions and a Mann–Whitney U test or *t*-test for continuous variables. Linear correlations were calculated using a Spearman's rank correlation coefficient. For visualization of correlations, linear regression analysis was performed to generate fitting lines.

Results

Microhemorrhages on histopathology

To assess evidence of red blood cell extravasation in APP/PS1 mice that exhibit moderate CAA at older age, we first screened for iron-positive deposits with *ex vivo* histopathology. The brains of 12 20.0 \pm 0.6 months old naïve APP/PS1 mice (7 Tg (4 F) and 5 WT (4 F) littermates) were extracted and cut in serial sections to assess presence and number of areas with focal parenchymal iron-positive deposits. On average, 40 sections per brain were visually screened on a brightfield

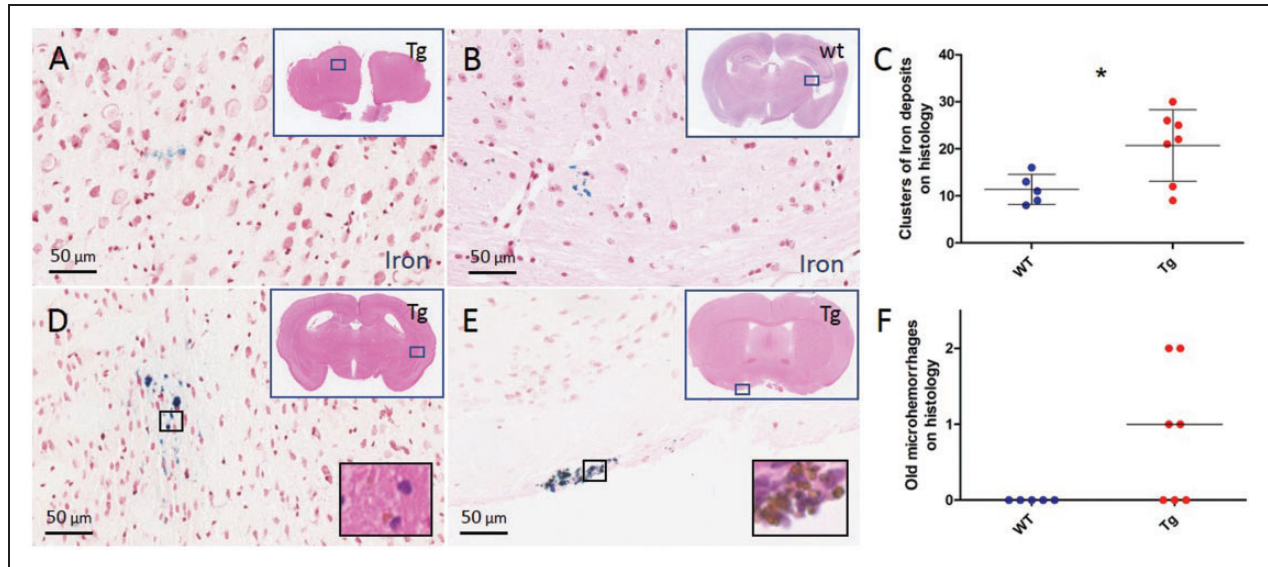


Figure 1. Histopathological evidence of leakage in the brains of 20-month-old APP/PS1 mice. Representative examples of areas with focal iron-positive deposits on a Prussian blue-stained section of the post-mortem brain of a 20-month-old Tg APP/PS1 mouse (a) and an age-matched WT mouse (b). A higher number of areas with clusters of focal iron-positive deposits was observed in Tg compared to WT mice (c, t -test $p=0.029$). Areas with clusters of >20 focal iron-positive deposits (d, e) with evidence of hemosiderin-containing macrophages on the adjacent H&E-stained section (insets) were rare and only observed in Tg mice (f, Mann–Whitney U test $p=0.065$). Lines and error bars in C indicate mean and standard deviations. Lines in F indicate median. Note that each dot in C and F represents a mouse. $*p < 0.05$.

microscope, which revealed that focal areas with clusters of Iron-positive deposits were more frequently present in Tg APP/PS1 mice (mean \pm standard deviation (SD) 21 ± 8) compared to WT mice (mean \pm SD 11 ± 3), $p=0.029$) (Figure 1). The majority of these areas consisted of only a few iron-positive deposits, suggestive of subtle blood–brain barrier leakage rather than frank microhemorrhages. Areas with >20 deposits and evidence of hemosiderin-containing macrophages on the adjacent H&E-stained section – indicative of old microhemorrhages – were rare and only observed in Tg APP/PS1 mice (six old microhemorrhages in $n=4$ Tg mice, $p=0.065$) (Figure 1).

In vivo visualization of focal microhemorrhages in APP/PS1 mice

To assess microhemorrhages in real-time in relation to their surrounding microvasculature, we next utilized *in vivo* two-photon microscopy in 20-month-old Tg APP/PS1 mice and their WT littermates with chronic cranial windows. Angiograms were visualized *in vivo* by injecting fluorescein dextran in the bloodstream prior to imaging. Mice were co-injected with fluorescently labeled anti-fibrin. Focal laser irradiations elicited dextran extravasation and rapid extravascular fibrin clot formation at the site of leakage in both arterioles (21/33 targeted) and venules (8/11 targeted) in $n=6$ mice

(3 Tg (2 F), 3 WT (3 F)) (Figure 2). Notably, over the course of 20 min, fibrin clots appeared to ‘wrap around’ arterioles, whereas this was not observed for venules. The next day, 20/21 clots on arterioles were still visible, whereas only 3/8 clots on venules ($p=0.022$). After three weeks, 10/16 clots on arterioles were still visible, compared to 1/6 clots on venules ($p=0.056$) (Figure 2). Notably, arteriole diameter was not associated with time to rupture (Spearman’s ρ 0.03, $p=0.88$), but more laser irradiation attempts were needed to achieve dextran extravasation from arterioles in Tg APP/PS1 mice compared to their WT littermates ($p=0.0044$). No significant correlation was observed between CAA burden at the irradiation site and time to rupture (Spearman’s ρ 0.36, $p=0.27$) (Figure 2).

Spontaneous and heparin-induced microhemorrhages in old APP/PS1 mice

Next, the prevalence of spontaneous microhemorrhages was assessed in real-time in two separate cohorts of Tg APP/PS1 mice and their WT littermates with chronic cranial windows at 8 months ($n=9$; 4 Tg (2 F), 5 WT (3 F)) and 19 months of age ($n=16$; 8 Tg (1 F), 8 WT (1 F)), after dextran and anti-fibrin were injected intravenously. Only sites of local dextran extravasation that showed evidence of focal extravascular clot formation were counted. Clots on newly

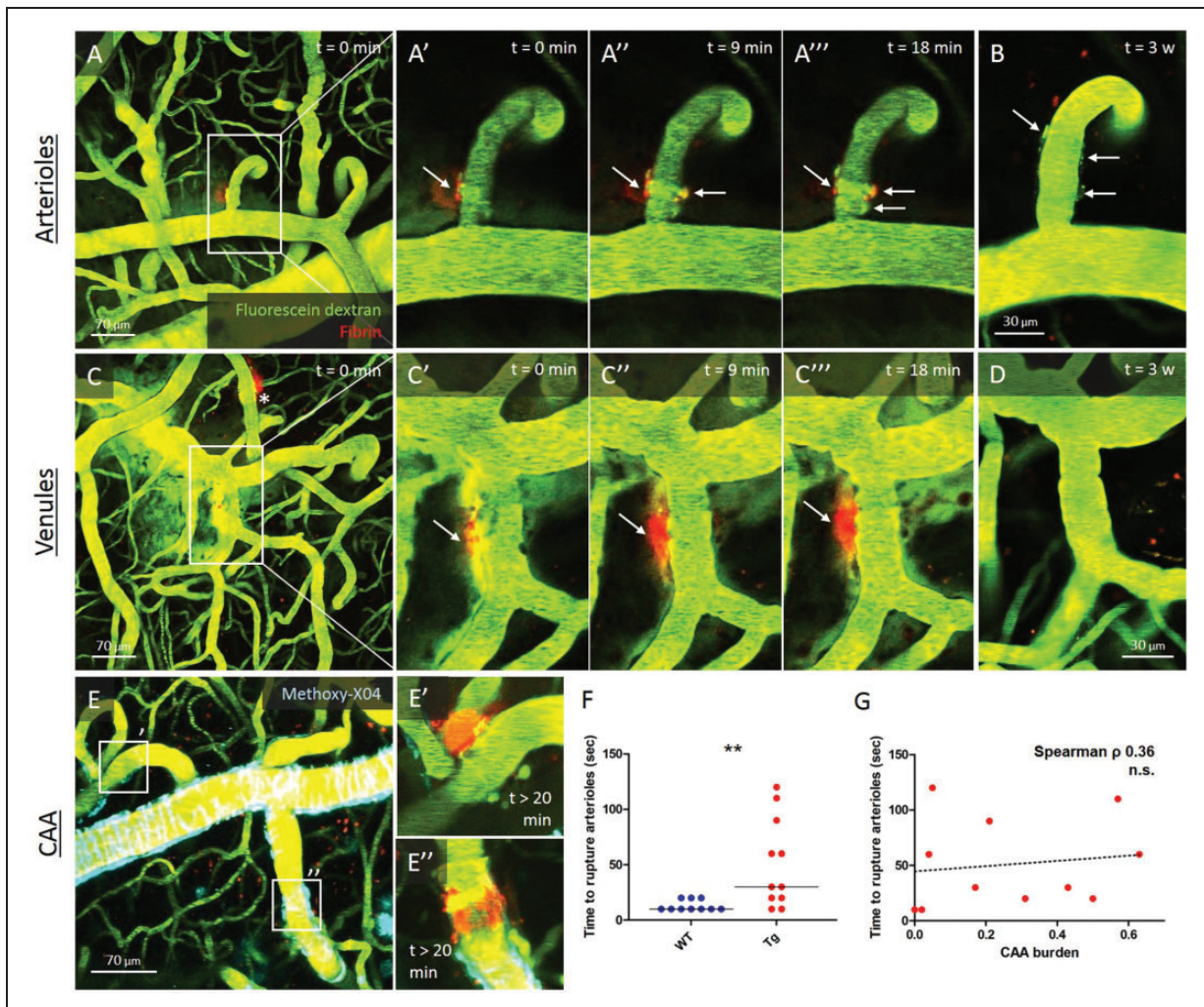


Figure 2. Visualizing microhemorrhage formation in vivo in real-time with two-photon microscopy in 20-month-old APP/PS1 mice. Representative example of the temporal progression of extravascular fibrin clot formation with in vivo two-photon microscopy in a 20-month-old WT mouse, depicted here immediately (a, a', c, c'), 9 min (a'', c''), and 18 min (a''', c''') after focal irradiation of an arteriole and a venule. Note the increasing circumferential appearance of the extravascular fibrin clot over time on the arteriole, but not the venule. Extravascular fibrin clots surrounding arterioles were more often still (partly) visible after one day (not shown) and after three weeks (b), compared to venules (d). In 20-month-old Tg APP/PS1 mice, a similar temporal progression of extravascular fibrin clot formation after focal irradiation was observed, in venules, and in arterioles with low (e') and high (e'') CAA burden. Notably, more laser irradiation attempts were needed to achieve dextran extravasation from arterioles in Tg APP/PS1 mice compared to their WT littermate controls (f, Mann–Whitney U test $p = 0.0044$). However, no significant correlation was observed between CAA burden at the irradiation site and time to rupture (g, Spearman's $\rho = 0.36$, $p = 0.27$). Note the additional extravascular fibrin clot on an arteriole from a previous irradiation in panel C (indicated with *). Lines in F indicate median. $^{**}p < 0.01$.

formed vessels that occasionally grow over the pial surface after chronic window implantation were excluded from the analysis. Mice were administered a high dose of heparin with the aim to lower the threshold for microhemorrhage formation. At eight months of age, no extravasations were observed at baseline in either the Tg APP/PS1 mice or their WT littermates, but heparin administration triggered a few sites to leak (Figure 3). At 19 months of age, baseline extravasations were observed in both Tg APP/PS1 mice (6/8) and their WT littermates (4/8, $p = 0.30$), whereas heparin

administration resulted in a higher occurrence of leakage sites in Tg (7/8) compared to WT mice (3/8, $p = 0.039$) (Figure 3).

Characterization of leakage sites in old APP/PS1 mice

The formation of extravascular fibrin clots at the sites of dextran extravasation allowed us to determine the exact site of leakage and the characteristics of the vessels at the leakage sites in relationship to the

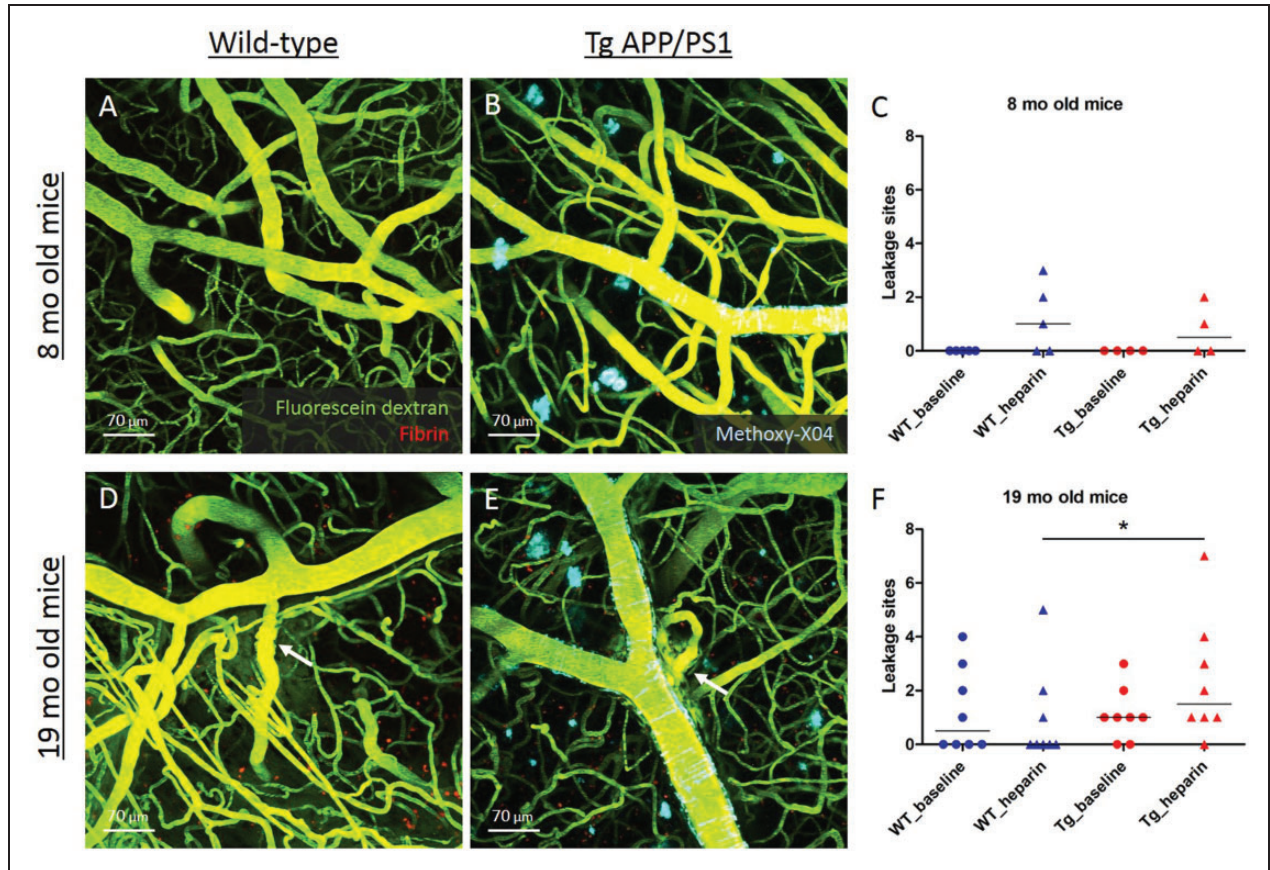


Figure 3. Spontaneous microhemorrhage formation after heparin administration in 19-month-old APP/PS1 mice. Spontaneous extravasations were not observed in 8-month-old Tg APP/PS1 mice (b) or their WT littermate controls (a), and only occasionally after heparin administration (c). At 19 months of age, both Tg APP/PS1 mice (e) and their WT littermate controls (d) had evidence of spontaneous extravasation (f, χ^2 test $p = 0.30$). Subsequent heparin administration resulted more often in extravasations in Tg compared to WT mice (f, χ^2 test $p = 0.039$). Lines in C and F indicate median. Note that each dot or triangle in C and F represents a mouse. * $p < 0.05$.

surrounding microvasculature and local CAA burden. Notably, leakage sites only occurred on arterioles, but not venules. The arteriolar diameters at the site of leakage were on average greater in 19-month-old Tg APP/PS1 mice (mean \pm SD $19 \pm 7 \mu\text{m}$), compared to their age-matched WT littermates (mean \pm SD $13 \pm 5 \mu\text{m}$, $p = 0.0054$) (Figure 4). In Tg APP/PS1 mice, approximately half of all fibrin clots (15/27, 56%) were found on first-order arteriolar branches, as compared to parent arterioles (1/27, 4%), second-order (9/27, 33%), and third-order arteriolar branches (2/27, 7%). One additional clot was found at the capillary level. Notably, CAA-negative first-order arteriolar branches (13/67, 19%) were more likely to bleed compared to CAA-positive first-order branches (2/67, 3%) ($p < 0.0001$) (Table 1). Ten fibrin clots either directly colocalized with or were found close to ($< 20 \mu\text{m}$) CAA deposits, whereas 18 were located remotely from CAA (on average $83 \pm 60 \mu\text{m}$ away, range 21–216 μm). In terms of vessel morphology, most clots

were observed at vessel branchpoints or bends (20/28, 71%) (Figure 4, Supplemental Figure 1).

Discussion

In this study, we demonstrated the feasibility of visualizing individual focal sites of microhemorrhage formation in relation to their surrounding microvasculature in real-time with in vivo two-photon microscopy in Tg mice. Our findings suggest that the vessels in old mice with CAA are more prone to leakage compared to non-Tg age-matched control mice after heparin administration. Leakage does not immediately occur from CAA-laden vessels, but rather from vulnerable vessel segments at bends or bifurcations close to or remote from vascular A β depositions.

Focal laser irradiations coupled with optical imaging have previously been used to assess microhemorrhage dynamics in real-time in vivo¹⁷ to determine effects on the surrounding tissue¹⁸ or the influence of

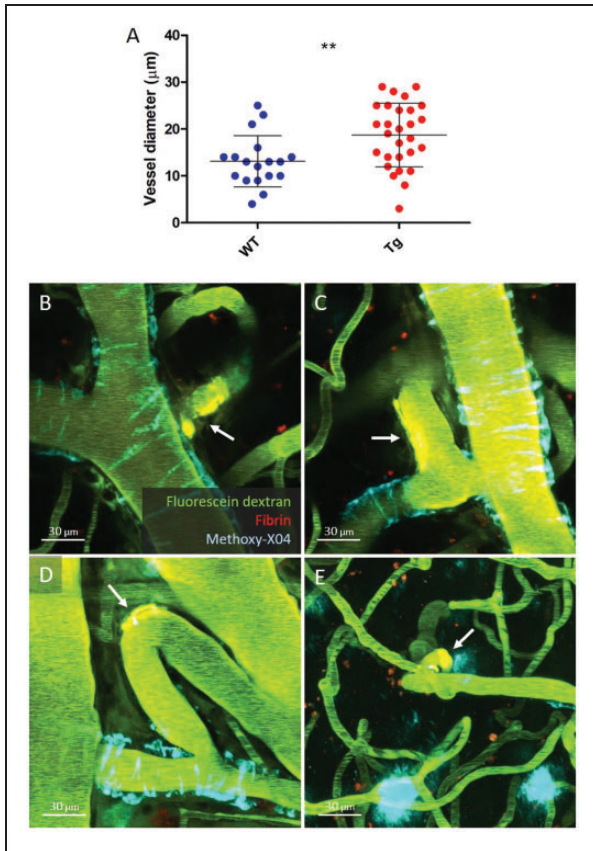


Figure 4. Characterization of spontaneous leakage sites in 19-month-old Tg APP/PS1 mice. Spontaneous extravasations in Tg APP/PS1 mice were on average found on larger arterioles (mean diameter at leakage sites $19 \pm 7 \mu\text{m}$) compared to their WT littermate controls ($13 \pm 5 \mu\text{m}$) (a, *t*-test $p = 0.0054$). In Tg APP/PS1 mice, the majority of extravascular fibrin clots (arrows) were found on CAA-negative first order arteriolar branches (b, c, d), as compared to higher order arteriolar segments (e). Notably, most clots (71%) were observed at vessel branchpoints (b, c) or bends (d, e) and remote from CAA deposits. Lines and error bars in A indicate mean and standard deviations. Note that the vessel in B is the same as depicted in Figure 3(e). Note that the Methoxy-X04 deposit directly adjacent to the clot in panel E is a parenchymal plaque and not CAA. $**p < 0.01$.

drug treatment on microhemorrhage size.^{15,19} Here we utilized this technique in combination with the detection of extravasated fluorescently labeled fibrin to identify the exact sites of microhemorrhages in vivo as a detection method for screening spontaneously occurring microhemorrhages in subsequent experiments. Using the same fluorescently labeled antibody, a previous study observed increased and more rapid fibrin extravasation after treatment with matrix metalloproteinase-9 (MMP9) in Tg2576 mice compared to age-matched WT littermate controls.¹⁴ MMP9 was topically applied on the surface of the mouse brain, resulting in widespread extravasation of

Table 1. Characterization of vessel types affected by fibrin clots in Tg APP/PS1 mice.

	Total number	Without clots	With clots	χ^2 (p-value)
CAA+ parent artery	30	29	1	
CAA- parent artery ^a	0	0	0	n/a
CAA+ 1st order arteriole	42	40	2	
CAA- 1st order arteriole ^a	25	12	13	0.0001
CAA+ 2nd order arteriole	11	10	1	
CAA- 2nd order arteriole ^a	44	36	8	0.47
CAA+ 3rd order arteriole	2	2	0	
CAA- 3rd order arteriole ^a	25	23	2	0.68

^aVessels that were classified as parent arteries had an average diameter of $40 \pm 11 \mu\text{m}$, first-order arterioles $22 \pm 4 \mu\text{m}$, second-order arterioles $16 \pm 5 \mu\text{m}$, and third-order arterioles $13 \pm 4 \mu\text{m}$.

fibrin. Therefore, individual leakage sites could not be determined, or whether leakage originated from vessel segments with or without vascular A β depositions. In our study, we observed focal extravascular clot formation after targeted laser irradiations that were comparable between 20-month-old APP/PS1 mice and their WT littermate controls. No notable differences were observed in terms of clot morphology between Tg and WT mice, whereas clots had a different appearance on arterioles compared to veins. The circumferential appearance of the fibrin clots on arterioles (and not veins) resembles the spiral-like deposition of CAA between vascular smooth muscle cells and could potentially reflect cellular uptake of fibrin in the perivascular space, an interesting topic for future studies.^{20,21} Unexpectedly, in mice with CAA, it took significantly longer for laser irradiation to rupture arterioles compared to arterioles in control mice, with a positive trend towards more attempts needed to rupture vessels with increased CAA burden. This may indicate a relative protective effect of mild-to-moderate CAA locally or the possibility that Methoxy-X04, the fluorescent dye that binds A β depositions, absorbed some of the laser power.

Using fluorescent fibrin as a marker to detect individual microhemorrhages in vivo, we observed an age-related increase of heparin-induced leakage sites in APP/PS1 mice. Previous studies have observed an increase in size of separately induced microhemorrhages in heparin-treated mice compared to vehicle-treated mice,^{19,22} indicating prolonged bleeding times of existing bleeds after heparin administration. Using a more sensitive detection method to assess small individual leakage sites, we observed heparin-induced microhemorrhages more often in 20-month-old APP/PS1 mice compared to their WT littermate controls. Although we took care to exclude leakage from newly formed vessels at the surface of the brain or the edge of

the cranial window, it cannot be excluded that some microhemorrhages indeed originated from these permeable vessels and therefore may have resulted in false positive findings. That said, the increased number of microhemorrhages observed in Tg mice was driven by microhemorrhages on larger diameter lower order arteriolar branches (the segments of interest), and not on the smaller newly formed vessels. Interestingly, microhemorrhages were only observed on arterioles and in particular on arteriolar segments without CAA. Although laser-induced fibrin clots were visible longer on arterioles compared to venules, the increased observation of spontaneous microhemorrhages on arterioles is unlikely due to this difference in detection. This is because the formation of the heparin-induced microhemorrhages was observed in real-time (<1 h after administration) and therefore clots on veins would have been detectable within that timeframe. A more likely explanation for the observed arteriolar predilection for microhemorrhages – which is in line with neuropathological observations in humans²³ – is that A β affects arterioles more often than veins in the context of CAA, as arterioles are believed to be the main routes for perivascular A β clearance.²⁴ Also, A β deposition is associated with smooth muscle cell degeneration, which reduces vessel wall integrity and increases the risk for microhemorrhages.²⁵ Even in the absence of vascular A β depositions, the presence of increased soluble A β may induce smooth muscle cell degeneration in vessel segments remote from the visible A β depositions.^{26,27} The frequent observation of microhemorrhages on bends and bifurcations in 20-month-old mice suggests a role for flow network dynamics in combination with focal damage. It may indicate that these anatomically vulnerable segments are progressively weakened due to sustained turbulence in those areas and thus likely to bleed in the context of vessel wall degeneration and overall increased A β levels.²⁵ Notably, microhemorrhages in brain tissue of CAA cases also seem to originate from penetrating arterioles at the sites where leptomeningeal vessels branch into the cortex, similarly to our observations of clots on first-order arteriolar branches in mice.²³ Future studies in mice are warranted to unravel potential functional alterations, such as blood flow or pulsatility changes, within affected vascular networks at those vulnerable locations.

Although 20 months represents an old age for the APP/PS1 mice, the observed CAA appears substantially less advanced compared to humans with severe CAA, suggesting that the spontaneous microhemorrhages in the mice may not be directly comparable to those in post-mortem human brain tissue of CAA patients. First, the mouse microhemorrhages are much smaller in size and perhaps better resemble subtle

blood–brain barrier leakage than frank hemorrhages, as in most cases no extravasation of red blood cells was observed in vivo. Our ex vivo histopathological examination also confirmed that old microhemorrhages are only rarely observed in the brains of Tg APP/PS1 mice post-mortem. Second, microhemorrhages in post-mortem human brains are only observed in the context of moderate-to-severe CAA.^{8,9,28} In these brains, A β depositions are not only found on leptomeningeal vessels, which is the case in APP/PS1 mice, but cover penetrating cortical arterioles and sometimes capillaries as well, resulting in vasculopathies such as fibrinoid necrosis and microaneurysms in the most severe cases. We have previously reported absence of vascular A β at the bleeding site in vessels involved in microhemorrhages in autopsy cases with severe CAA,²⁹ but serial sectioning reveals extensive vessel remodeling prior to bleeding, including thickening of the wall and degeneration of vascular smooth muscle cells,²³ indicating severe vessel wall degeneration not observed in the mice. Therefore, the mechanisms underlying the microhemorrhages (or leakage sites) observed in mice with CAA may not necessarily resemble what happens in patients with severe CAA. Alternatively, they could represent blood–brain barrier leakage that may happen in an earlier stage of the pathophysiology of CAA.

Observations from both post-mortem human brain tissue and our in vivo findings in mice suggest that just the presence of vascular A β per se is not sufficient to make vessels bleed directly but that more complex vascular network dynamics likely play a role. Future studies of spontaneous microhemorrhage formation in real-time in mice with more severe CAA and vascular remodeling, such as APP23 mice,^{30,31} would be informative. Furthermore, our in vivo approach allows the longitudinal assessment of the role of vascular dysfunction (e.g. impaired evoked vascular reactivity) on the formation of spontaneous lesions at the single-vessel level, which is currently not feasible in humans.

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Declaration of conflicting interests

The author(s) declared no potential conflicts of interest with respect to the research, authorship, and/or publication of this article.

Authors' contributions

Susanne J. van Veluw was involved in the concept and design of the study, analysis and interpretation of the data, drafting and revising of the manuscript, and obtained funding.

Matthew P. Frosch was involved in the concept and design of the study and revising of the manuscript.

Ashley A. Scherlek was involved in the analysis and interpretation of the data and revising of the manuscript.

Daniel Lee was involved in the analysis and interpretation of the data and revising of the manuscript.

Steven M. Greenberg was involved in the concept and design of the study and revising of the manuscript, and obtained funding.

Brian J. Bacskai was involved in the concept and design of the study and revising of the manuscript, and obtained funding.

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Supplemental material

Supplemental material for this article is available online.

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