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A novel role for osteopontin in macrophage-mediated amyloid- β clearance in Alzheimer's models



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ABSTRACT

Osteopontin (OPN), a matricellular immunomodulatory cytokine highly expressed by myelomonocytic cells, is known to regulate immune cell migration, communication, and response to brain injury. Enhanced cerebral recruitment of monocytes achieved through glatiramer acetate (GA) immunization or peripheral blood enrichment with bone marrow (BM)-derived CD115⁺ monocytes (Mo^{BM}) curbs amyloid β -protein (A β) neuropathology and preserves cognitive function in murine models of Alzheimer's disease (ADtg mice). To elucidate the beneficial mechanisms of these immunomodulatory approaches in AD, we focused on the potential role of OPN in macrophage-mediated A β clearance. Here, we found extensive OPN upregulation along with reduction of vascular and parenchymal A β burden in cortices and hippocampi of GA-immunized ADtg mice. Treatment combining GA with blood-grafted Mo^{BM} further increased OPN levels surrounding residual A β plaques. In brains from AD patients and ADtg mice, OPN was also elevated and predominantly expressed by infiltrating GFP⁻ or Iba1⁺-CD45^{high} monocyte-derived macrophages engulfing A β plaques. Following GA immunization, we detected a significant increase in a subpopulation of inflammatory blood monocytes (CD115⁺CD11b⁺Ly6C^{high}) expressing OPN, and subsequently, an elevated population of OPN-expressing CD11b⁺Ly6C⁺CD45^{high} monocyte/macrophages in the brains of these ADtg mice. Corrologram analyses indicate a strong linear correlation between cerebral OPN levels and macrophage infiltration, as well as a tight inverse relation between OPN and A β -plaque burden. In vitro studies corroborate in vivo findings by showing that GA directly upregulates OPN expression in BM-derived macrophages (M Φ ^{BM}). Further, OPN promotes a phenotypic shift that is highly phagocytic (increased uptake of A β fibrils and surface scavenger receptors) and anti-inflammatory (altered cell morphology, reduced iNOS, and elevated IL-10 and A β -degrading enzyme MMP-9). Inhibition of OPN expression in M Φ ^{BM}, either by siRNA, knockout (KO^{OPN}), or minocycline, impairs uptake of A β fibrils and hinders GA's neuroprotective effects on macrophage immunological profile. Addition of human recombinant OPN reverses the impaired A β phagocytosis in KO^{OPN}-M Φ ^{BM}. This study demonstrates that OPN has an essential role in modulating macrophage immunological profile and their ability to resist pathogenic forms of A β .

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1. Introduction

Osteopontin (OPN; encoded by the gene secreted phosphoprotein 1, SPP1) is a multifunctional matricellular glycoprotein secreted by a variety of activated immune cells such as macro-

phages and T cells, among other cell types (Franzen and Heinegard, 1985; Patarca et al., 1989; Scatena et al., 2007). It is highly expressed by bone marrow (BM)-derived myelomonocytic cells and modulates their migration, communication, and immunological responses (Lund et al., 2009; Rittling, 2011). Recent studies indicate that this glycoprotein may curb the pathology of various brain diseases such as stroke, ischemia, and traumatic brain injury via neuroprotective and repair-promoting effects (Brown, 2012;

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Chan et al., 2014; Gliem et al., 2015; Meller et al., 2005; van Velthoven et al., 2011). Examination of OPN is particularly timely, since it is notably upregulated during the inflammation associated with Alzheimer's disease (AD) and other neurodegenerative conditions (Choi et al., 2007; Comi et al., 2010; Kim et al., 2004; Wung et al., 2007).

The past century has seen substantial growth in our understanding of AD and the molecular mechanisms underlying its development. When detected within the brain, extracellular plaques, comprising predominantly of amyloid- β protein ($A\beta$), and intracellular neurofibrillary tangles, composed primarily of hyperphosphorylated tau, are still considered to be the pathological hallmarks of the disease (De Strooper and Karran, 2016; Glenner et al., 1984; Hardy et al., 1998). It has been shown that misfolded $A\beta$ forms, especially aggregates of $A\beta_{40}$ and, moreover, $A\beta_{42}$ alloforms, are highly synaptotoxic and that their accumulation is pathogenic to AD (Shankar et al., 2008). Furthermore, exploration of this disease in central nervous system tissues beyond the brain has revealed $A\beta$ pathology in retinas of AD patients, including those at early stages (Koronyo et al., 2017; Koronyo-Hamaoui et al., 2011; La Morgia et al., 2016; Hart et al., 2016 #59).

Converging data from genetic, physiologic, biochemical, and clinical studies demonstrate a strong association between $A\beta$ accumulation and neuroinflammation, synaptic loss, impaired neuronal function, and ultimately, debilitating cognitive decline (Selkoe, 2001 #73; Wyss-Coray, 2006). The progressive accumulation and aggregation of $A\beta$ peptides in the brain are thought to result from an imbalance between their production and clearance (Mawuenyega et al., 2010; Saido, 1998). Inadequate cerebral $A\beta_{42}$ clearance was detected in sporadic common cases of AD (Mawuenyega et al., 2010). Moreover, the dramatic increase in cerebral $A\beta$ begins as early as 20 years prior to symptom manifestation, far preceding clinical impairment (Bateman et al., 2012; Bilgel et al., 2016; Perrin et al., 2009). Therefore, a strategy capable of reducing $A\beta$ levels in the brain, either by inhibiting production and the resulting aggregation or by promoting clearance, would be advantageous in preventing the development of AD.

Growing evidence suggests that infiltrating innate immune cells may play a key role in the clearance of cerebral $A\beta$ plaques in murine models of AD (Bernstein et al., 2014; Butovsky et al., 2007; El Khoury et al., 2007; Koronyo et al., 2015; Koronyo-Hamaoui et al., 2009; Lai and McLaurin, 2012; Lebson et al., 2010; Malm et al., 2012; Simard et al., 2006). We are among a number of groups to demonstrate that immunomodulation can lead to increased cerebral infiltration by monocytes, cells directly involved in $A\beta$ -plaque clearance. In our studies, we found that beneficial immunomodulation in double transgenic APP_{SWE}/PS1 Δ E9 (ADtg) mice was facilitated either by altered myelin-derived antigens, such as glatiramer acetate (GA; also called Copaxone[®]) and MOG45D, or by adoptive transfer of inflammatory monocytes into the peripheral bloodstream (Bakalash et al., 2011; Bernstein et al., 2014; Butovsky et al., 2006, 2007; Frenkel et al., 2005; Koronyo et al., 2015; Koronyo-Hamaoui et al., 2009, 2011; Lebson et al., 2010). In particular, GA immunization alone and in combination with BM-CD115⁺ monocyte (Mo^{BM}) blood enrichment delayed AD-like progression and profoundly attenuated various abnormalities associated with AD. Following therapy, we observed reduced levels of soluble and insoluble $A\beta_{42}$ in the brain, enhanced proteolytic degradation of $A\beta$ (by MMP-9, ACE, neprilysin), regulation of neuroinflammation, and preservation of synapses and cognitive function in ADtg mice (Koronyo et al., 2015). In BM-derived macrophage (M Φ ^{BM}) cultures, GA activation increased phagocytosis of fibrillar $A\beta$, a change partly attributed to elevation of surface scavenger receptors (i.e. CD36, Scara-1) (Koronyo et al., 2015).

While the exact molecular mechanisms through which GA affects monocyte and macrophage recruitment and function in

AD remain largely unclear, recent reports have suggested a promising link between a subset of these cells and neuroprotective clearance. Among these are the involvement of OPN-producing macrophages in the phagocytosis of fragmented cell debris following stroke (Shin et al., 2011), and an impairment of phagocytic activity in OPN-deficient mice (KO^{OPN}) (Schack et al., 2009; Toyonaga et al., 2015). These findings prompted us to investigate how cerebral OPN expression is impacted following GA immunomodulation in ADtg mice and how it relates to monocyte migration and $A\beta$ -plaque burden. Furthermore, we have examined how OPN inhibition affects the immune phenotype of macrophages and its effect on their ability to phagocytose pathogenic $A\beta_{40}$ and $A\beta_{42}$ aggregates associated with AD.

2. Materials and methods

2.1. Mice

Double transgenic mouse models of Alzheimer's disease (ADtg) from the B6Cg-Tg (APP_{SWE}, PSEN1E9) 85Dbo/J strains and their age-matched non-transgenic (WT) littermates were purchased from Jackson Laboratories (MMRC stock #34832-JAX|APP/PS1), then bred and maintained at Cedars-Sinai Medical Center. ADtg mice carrying the human transgene allow for detection of $A\beta$ forms with anti-human $A\beta$ antibodies. OPN knock-out (KO^{OPN}) mice from the B6.129S6 (Cg)-*Spp1*^{tm1Blh}/J strain (Jackson laboratories stock #004936|OPN KO) were used for in vitro experiments. For our studies, mice with ages spanning from 2 months to 24 months old were used. For all in vitro experiments, bone marrow was collected from young mice, aged 8–16 weeks. In addition, young C57BL/6-tg (UBC-GFP) 30 Scha/J donor mice, aged 8–10 weeks, were used for adoptive transfer of bone marrow-derived GFP⁺ monocytes. These mice express enhanced green fluorescent protein (GFP) under direction of the human ubiquitin C promoter. All mice in this study had a C57BL/6 congenic background, with males and females in equal number and age-matched when applicable. All experiments were conducted and recorded by researchers blinded to mouse genotypes and/or treatment. This research was performed in accordance with Cedars-Sinai Medical Center Institutional Animal Care and Use Committee (IACUC) guidelines under an approved protocol and complied with the current United States applicable laws.

2.2. Human brain tissues

Postmortem brain sections from the frontal cortex of 2 neuropathologically confirmed AD patients and 2 age- and gender-matched (cognitively normal) control individuals were prepared and analyzed by immunohistochemistry. Human tissues were obtained from the Alzheimer's Disease Research Center (ADRC) Neuropathology Core (IRB protocol: HS-042071) at the University of Southern California's (USC; Los Angeles, CA) Department of Pathology.

2.3. Glatiramer acetate (GA) immunization

Symptomatic 10-month-old ADtg male mice received either subcutaneous injections of 100 μ g GA (also known as Copaxone, in 200 μ l PBS; TEVA Neuroscience) or PBS alone (control). Injections were administered twice during the first week, then once a week for 8 weeks. After completing treatment, all mice were anaesthetized and perfused with ice-cold PBS supplemented with 0.5 mM EDTA. Brains were collected and analyzed or fixed in 2.5% paraformaldehyde (Sigma-Aldrich), and cryo-protected in 30% sucrose for further histological analysis.

Flow cytometry analysis was conducted on blood and brain of adult ADtg and WT mice (~3 month old; $n = 4$ mice/per group and genotype) that were immunized with GA or PBS, according to a regimen of twice a week the first 2 weeks then once a week the following 2 weeks (total of 6 injections). Blood (up to 0.5% of mouse body weight) was collected from the saphenous vein before and after the last GA injection. At the end of the experiment, all mice were perfused and brains were collected and immediately processed for flow cytometry analysis ($n = 3$ ADtg mouse brains per treatment group).

2.4. Adoptive transfer

CD115⁺ monocytes from donor mice were isolated as previously reported in [Koronyo et al. \(2015\)](#). In short, after euthanasia, bone marrow cells were harvested from femurs, tibiae, and humeri, and enriched for mononuclear cells on a Ficoll-Paque PLUS (GE Healthcare) density gradient. The CD115⁺ monocyte population was isolated with a MACS enrichment column using the biotinylated anti-CD115 mAb clone AFS98 (eBioscience) and streptavidin-coupled magnetic beads (Miltenyi Biotec), per manufacturer's protocols. After this procedure, the entire CD115⁺ monocyte population ($5\text{--}6 \times 10^6$ cells/mouse) was injected into tail veins of 10-month-old ADtg mice once a month for 2 months.

2.5. Immunohistochemistry (IHC)

Coronal brain sections (30 μm thick) were treated for 30 min in antigen-retrieval solution (Dako) prior to serum-free protein blocking (Dako Cytomation). Sections were then hybridized with various primary antibodies (overnight at 4 °C): goat polyclonal OPN (R&D systems), mouse anti-human A β [residues 17–24, mAb clone 4G8 (1:100; Covance), and residues 1–16, mAb clone 6E10 (1:100; Covance)], rabbit anti-GFAP pAb (1:100; Sigma-Aldrich), rabbit anti-Iba1 pAb (1:250; Wako Chemicals), rabbit anti-GFP pAb (1:500; MBL), rat anti-CD68 mAb (1:100; Abcam), rabbit anti-iNOS pAb (1:100; Cell signaling), rat anti-CD45 mAb clone 30-F11 (1:25; BD Pharmingen), mouse anti-human CD45 mAb (1:20; BD Pharmingen), mouse anti-Tuj1 mAb (1:200; Abcam), rabbit anti-NeuN mAb (1:200; Abcam) and goat anti-MMP9 pAb (1:100; R&D systems). Hybridization with primary antibodies was followed by incubation with biotinylated secondary antibody and conjugation with horseradish peroxidase (HRP) [Vectastain Elite ABC kit from Vector laboratories using DAB-plus substrate chromogen system (DAKO)] or fluorophore-conjugated secondary antibodies (1 h at 37 °C; donkey anti-mouse, anti-rat, anti-goat, and anti-rabbit; 1:200; Jackson Immuno Research Laboratories) conjugated with Cy2, Cy3, and Cy5. Sections were mounted using ProLong Gold with DAPI (Molecular Probes, Life Technologies). Negative controls were processed using the same protocol with the omission of the primary antibody to assess non-specific labeling. A Carl Zeiss Axio Imager Z1 fluorescence microscope equipped with ApoTome (Carl Zeiss MicroImaging, Inc.) was used for microscopic analysis. AxioVision (release 4.6.3) software (Carl Zeiss) was used to process and analyze the images.

2.6. Thioflavin-S staining and CAA scores

For amyloid burden assessment in mouse/human brain parenchyma and vasculature, sections were stained with Thioflavin S (Thio-S, 1% w/v in 70% ethanol; Sigma-Aldrich) for 10 min at room temperature following the secondary antibody step, according to a previously described standard protocol ([Koronyo et al., 2015](#)). Various degrees of cerebral amyloid angiopathy (CAA) in animals were defined by analyzing Thio-S-labeled brain sections using a scale of 0–4 (0 indicates no CAA, 4 indicates severe CAA;

5 brain sections per animal), as previously described ([Wyss-Coray et al., 2001](#)).

2.7. Primary cultures of bone marrow-derived macrophages

Bone marrow-derived macrophages (M Φ^{BM}) from WT or KO^{OPN} mice were generated from femurs and tibiae after euthanasia, and cultured for 6–7 days in complete RPMI 1640 medium (Life Technologies) supplemented with 10% FBS, 100 U/ml penicillin, 100 $\mu\text{g}/\text{ml}$ streptomycin, 2 mM L-glutamine and 20 ng/ml MCSF (PeproTech). Primary cultures of M Φ^{BM} were then plated overnight with 1×10^5 cells per well (3–5 wells per condition) in 24-well tissue culture plates on glass coverslips. Next, macrophages were treated with either 30 $\mu\text{g}/\text{mL}$ GA, 50 ng/ml Recombinant Human Osteopontin (rOPN; Peprrotech), Brefeldin A (BFA; Sigma, 1X), or 10 μM Minocycline (Sigma) for a duration of 3 or 24 h (each component diluted in the culture medium). The control group was not treated. Immediately following addition of preformed fibrillar A β_{1-42} or A β_{1-40} (100 nM), plates were briefly centrifuged and incubated at 37 °C for 30 or 60 min. Cells were then rinsed with culture medium to remove non-incorporated A β , washed with PBS, and fixed in cold methanol (99.8%, –20 °C) for immunocytochemistry.

2.8. Immunocytochemistry (ICC)

After serum-free protein blocking (Dako Cytomation), cells were hybridized with various primary antibodies (overnight at 4 °C), including goat polyclonal OPN (R&D systems), mouse anti-human residues 1–16, mAb clone 6E10 (1:100; Covance)], rabbit anti-Iba1 pAb (1:250; Wako Chemicals), rat anti-CD68 mAb (1:100; Abcam), rabbit anti-iNOS pAb (1:100; Cell signaling), goat anti-MMP9 pAb (1:100; R&D systems), Rat anti-CD36 mAb (1:xx; Abcam), rabbit anti-LC3 pAb (1:250; Novus), mouse anti-Golgi mAb (1:50; Sigma-Aldrich), rabbit anti-EEA1 pAb (1:100; Millipore), rabbit anti-Rab7 mAb (1:100; Abcam), rat anti-SCARA1 mAb (1:100; AbD Serotec), and anti IL-10 pAb (1:100; R&D system). Hybridization with primary antibodies was followed by incubation with appropriate secondary polyclonal antibodies (1 h at 37 °C; donkey anti-mouse, anti-rat, anti-goat, and anti-rabbit; 1:200; Jackson Immuno Research Laboratories) conjugated with Cy2, Cy3, and Cy5. Sections were mounted using ProLong Gold with DAPI (Molecular Probes, Life Technologies) and analyzed as previously described.

2.9. Microscopy and quantification

Fluorescence of specific signals was captured with a Carl Zeiss Axio Imager Z1 fluorescence microscope equipped with ApoTome (Carl Zeiss MicroImaging, Inc.) using the same exposure time for each image. Images were analyzed using ImageJ software (NIH), converted to grayscale and using the same post-acquisition threshold for analysis. Manual counting of OPN⁺/CD45^{hi} and Iba1⁺/CD45^{hi} cells was performed with ImageJ software, using the 'analyze' grid, by analyzers blinded to mouse genotype and treatment groups. Quantification of the number and area (μm^2) of OPN⁺ cells, Iba1⁺-CD45^{hi} and OPN⁺CD45^{high} cells, 6E10⁺, Thio-S⁺ and 4G8⁺ A β plaques, as well as OPN, iNOS, CD36, SCARA-1, IL10, and MMP9 immunoreactive area, were determined as follows: 1. For IHC, four to six coronal brain sections at 150- μm intervals per mouse were examined over an area covering both hippocampal and cortical regions (including entorhinal and cingulate cortex); 2. For ICC, five to ten images from each well ($n = 3\text{--}6$ wells per group), covering on average 100 cells per image, were analyzed.

2.10. Enzyme-linked immunosorbent assay (ELISA) on cell samples and brain homogenates

MΦ^{BM} cells in each well were lifted by 2 mM EDTA-PBS, collected in tubes, and centrifuged separating cells from supernatant. Cell pellets were lysed and re-suspended in a cocktail of RIPA buffer supplemented with 1% protease inhibitors (Thermo Scientific), and stored at -80°C until use. Brain tissues, freshly collected after perfusion, were immediately and thoroughly homogenized in PBS buffer with 0.5% Triton X-100 (Sigma) and 1X protease inhibitor cocktail set I (Calbiochem), then processed as previously described (Koronyo et al., 2015).

After determination of protein concentration using the Pierce BCA Protein Assay Kit (Thermo Scientific), concentrations of A β ₄₂, A β ₄₀ and OPN were analyzed with an anti-human A β _{1–42} and A β _{1–40} specific sandwich ELISA kit (Invitrogen) and an anti-mouse OPN quantikine ELISA kit (R&D Systems), per manufacturer's instructions. The optical density of each well was read at 450 nm (with 540 nm correction for OPN) using the same microplate reader (Spectra Max 384 plus, Molecular Devices).

2.11. Western blotting

MΦ^{BM} were lifted from each cell by 2 mM EDTA-PBS, collected in an Eppendorf tube, and centrifuged. Cell pellets were lysed in RIPA buffer (Thermo Scientific) and supplemented with a cocktail of protease inhibitors (Thermo Scientific). Protein concentration was determined using a BCA Protein Assay Kit (Thermo Scientific). Aliquots of protein samples were electrophoretically separated using 4–12% Bis-Tris gels (Invitrogen), then transferred to nitrocellulose membranes, blocked in Tris-buffered saline (TBS) containing 5% (w/v) non-fat dry milk, and hybridized with appropriate primary antibodies, including rabbit anti-OPN pAb (Abcam), goat anti-OPN pAb (R&D systems), and β -Actin (Abcam). Rabbit anti-OPN pAb recognizes the full-length as well as the MMP-cleaved OPN fragments, whereas the goat anti-OPN pAb recognizes and provides a strong signal only for the full-length OPN. Membranes were then incubated with the appropriate HRP-conjugated secondary antibody prior to development with chemiluminescent substrates. Densitometric analysis of blots was conducted using ImageJ software, and each experimental sample was normalized to β -Actin.

2.12. Preparation of A β ₄₂ and A β ₄₀ fibrils

Lyophilized A β _{1–40} and A β _{1–42} peptides (Anaspec) were dissolved in ice cold hexafluoroisopropanol (HFIP) (Sigma) to create a monomeric solution of 100 μM . The HFIP was evaporated in a sterile hood creating a desiccated aggregate-free peptide film that can be stored at -20°C . HFIP was used to ensure preparation of A β fibrils from uniform monomeric A β peptide layer. Leftover traces were removed in a SpeedVac for 2 h. To initiate fibril formation, A β films were resuspended in a sterile mixture containing 45% 20 mM NaH₂PO₄ + Na₂HPO₄ (pH 7.4) and 10% NaOH 60 mM in 45% H₂O, vortexed for 30 s. A β solutions were incubated on a shaker for 2 weeks at 37 $^{\circ}\text{C}$. Pre-formed A β fibrils were then sonicated for 60 s and diluted to 100 nM in culture media prior to the phagocytosis assay.

2.13. OPN knockdown by siRNA in vitro

We used a set of three Stealth OPN siRNAs made up of 25-bp duplex oligoribonucleotides each, with sequences corresponding to the sense and antisense strands of OPN (Invitrogen). Equal amounts (100 nM each) of siRNAs were mixed and diluted in Opti-MEM (Gibco) to a final concentration of 100 nmol/L, then tran-

siently transfected into MΦ^{BM} using Lipofectamine[®] RNAiMAX Transfection Reagent (Invitrogen), per manufacturer's protocol. Control macrophages were transfected with Stealth RNAi[™] siRNA Negative Control. MΦ^{BM} were incubated for 48 h after transfection, and were used for subsequent experiments.

2.14. Cell length measurement

The long and short axes of cells were manually measured in μm from microscopic images (McWhorter et al., 2013) using length tools in the Axiovision Rel. 4.8 software package. At least 100 cells were examined in each experiment where individual cells were assayed by microscopy.

2.15. Flow cytometry analysis of murine peripheral blood

Blood collected into EDTA tubes was processed for flow cytometry analysis as follows.

After centrifugation (1800 RPM, 4 $^{\circ}\text{C}$, 12 min), RBC lysis (1 \times ; #420301, Biolegend) was applied on the cell pellets (2 ml of RBC lysis to 100 μl of whole blood, incubated for 13 min in dark and R.T.). After centrifugation (1800 RPM, 4 $^{\circ}\text{C}$, 5 min) and washes with FACS buffer (2% BSA, 0.1% w/v NaN₃, #438456, Sigma, in PBS), cells were counted in the presence of Trypan blue stain (0.4%; #15250-061, Gibco). Subsequently, cells were blocked with FC receptor blocking (anti mouse CD16/32, #101301), and stained for cell surface markers with the following antibodies: FITC-conjugated anti-mouse-CD11b (clone M1/70; #101206), APC-conjugated anti-mouse CD115 (clone AFS98; #135510) and Pacific blue-conjugated anti-mouse Ly6C (clone H.K1.4; #128013). All required controls were added to the flow cytometry analysis including isotype control antibodies: Rat IgG2b, κ – FITC (clone RTK4530; #400605); Rat IgG2a, κ – APC (clone RTK2758; #400511); Rat IgG2c, κ – Pacific Blue[™] (clone RTK4174; #400717) and Goat IgG – PE (clone Poly24030; #403004). All antibodies were diluted 1:100 and purchased from BioLegend.

After staining (30 min in dark, 4 $^{\circ}\text{C}$), the cells were washed and centrifuged. Next, cells were fixed with 500 μl cold 2.5% PFA in PBS for 15 min in R.T., then washed and centrifuged twice. After fixation, cells were permeabilized with 2 ml Saponin buffer (0.1% w/v Saponin #S4521, Sigma, 0.05% w/v NaN₃, #438456, Sigma, in Hank's Balanced salt solution sterile 1x, Corning) for 5 min. Samples were centrifuged and cells were stained for intracellular markers with the following antibody: PE-conjugated anti-mouse OPN (#IC808P, R&D). After staining (30 min in dark, 4 $^{\circ}\text{C}$), cells were washed and centrifuged twice, and PBS (1x, 300 μl) was finally added to the samples. Samples were analyzed with flow cytometer (BD LSRFortessa[™], BD Biosciences) and data were analyzed with FlowJo software (Tree Star, Inc.).

2.16. Flow cytometry analysis of myelomonocytes in ADtg mouse brain tissue

Whole brains freshly collected after perfusions were mechanically minced in a 70- μm cell strainer (Falcon; Corning Inc.) with ice-cold 2% fetal bovine serum (Atlanta Biological) in PBS. After centrifugation, homogenization and washing, pellets were suspended in 40% sterile Percoll (GE Healthcare) and centrifuged (850g, 4 $^{\circ}\text{C}$, 25 min, without brake). Cell pellets were re-suspended in 70% sterile Percoll and centrifuged (800g, 4 $^{\circ}\text{C}$, 20 min, without brake). Cells located in the top layer after the Percoll gradient were collected and washed. Mononuclear cells were counted in the presence of Trypan blue stain (0.4%; #15250-061, Gibco). Subsequently, cells were blocked with FC receptor blocking, 1 μg per 10⁶ cells (anti mouse CD16/32, #101301), and stained (30 min in the dark, 4 $^{\circ}\text{C}$) for cell surface markers with the follow-

ing antibodies: FITC-conjugated anti-CD11b clone M1/70 (#101206); Pacific blue-conjugated anti mouse -Ly6C (clone H. K1.4; #128013); and PE/Cy7-conjugated anti-CD45.2 clone 104 (#109830). All antibodies were diluted 1:100, and purchased from BioLegend. All required controls were added to the flow cytometry analysis including isotype control antibodies as previously described. After staining (30 min in dark, 4 °C) the cells were washed and centrifuged. Following steps (PFA fixation, saponin permeabilization, OPN intracellular staining and analysis) are similar to the blood processing (see above).

2.17. Statistical analysis

Experimental data was analyzed using GraphPad Prism 6.01 (GraphPad Software). In cases where three or more groups were compared, two-way or one-way ANOVA was performed, followed by the Tukey's, Dunnett's, or Bonferroni's multiple comparison test of paired groups. Two-tailed unpaired Student's *t*-tests were used in two-group comparisons. The statistical association between two or more variables was determined by Pearson's correlation coefficient (*r*) test (GraphPad Prism). Pearson's *r* indicates direction and strength of the linear relationship between two variables. Results are shown as means ± standard errors of the mean (SEMs). Degree of significance between groups is represented as follows: **p* < 0.05, ***p* < 0.01, ****p* < 0.001, and *****p* < 0.0001. A *p*-value lower than 0.05 was considered significant. All data analysis was led by blind examiner; code was revealed when analyses were concluded.

3. Results

3.1. Distinct patterns of cerebral OPN expression associated with Alzheimer's disease

To investigate cerebral OPN distribution in various brain regions and cell types, we initially analyzed brain regions associated with AD (Fig. 1A) in symptomatic 13-month-old male ADtg mice in comparison with age- and gender-matched wild-type (WT) non-Tg littermates. Immunohistochemistry (IHC) revealed pronounced OPN expression patterns in the hippocampus (HC), cingulate cortex (CC), and entorhinal cortex (EC) in ADtg mice within or near Aβ plaques (Fig. 1B, C). Similar patterns of OPN expression were observed in brains of ADtg mice with peroxidase labeling (Fig. 1C). OPN immunolabeling was largely undetectable in control WT mice (Fig. 1B). Next, we screened for the type of cells that express cerebral OPN in both ADtg and WT mice (Fig. 1D, E). In ADtg mice, OPN protein did not colocalize with markers of neurons (Tuj1, NeuN) or reactive astrocytes (GFAP) in CC and EC (Fig. 1D). While OPN did not appear in resting parenchymal Iba1⁺ myeloid cells in association with plaques in both WT and ADtg mice, activated Iba1⁺ microglia/macrophages clustering around amyloid plaques showed strong OPN expression (Fig. 1E). As expected, no such plaque-related cell clusters were found in WT mice. In brain regions not typically associated with AD (i.e. striatum), OPN immunostaining was predominantly confined to neurons in both ADtg and WT mice (Fig. S1). Analysis of cerebral OPN levels by quantitative ELISA showed protein levels increased 1.6–2.2 times in the brains of ADtg versus WT mice, in both 13- and 24-month-old mice, and increased 2–3 times with aging (13- vs. 24-month-old mice) in both genotypes (*p* < 0.01 – 0.0001).

To further explore OPN expression patterns in brains of AD patients, frontal cortices from four AD patients and two age- and gender-matched controls were immunostained and analyzed. In support of findings from murine models of AD (Fig. 1), patients showed a marked increase in cortical OPN expression in comparison with controls (Fig. S2A, B). OPN was located within and around

Iba1⁺ and/or CD45⁺ myelomonocytic cells (i.e. microglia, macrophages), predominantly surrounding Thio-S⁺ amyloid plaques (Fig. S2C–H). Overall, these results indicate that in brain regions affected by AD, OPN is selectively expressed by myeloid cells surrounding Aβ plaques.

3.2. Elevated expression of cerebral OPN following immunomodulation therapy is associated with reduced parenchymal and vascular Aβ burden

Previously, we demonstrated that GA immunization alone or in combination with monocyte blood enrichment attenuates disease progression, and that this effect is mediated by cerebral recruitment and modulation of myelomonocytic populations surrounding Aβ plaques in ADtg mice (Butovsky et al., 2006, 2007; Koronyo et al., 2015; Zuroff et al., 2017). Given that OPN is an immunomodulator highly expressed by myelomonocytic cells, we studied the effects of this approach on OPN cerebral expression (Figs. 2–4). To this end, we analyzed the levels of OPN in AD-associated brain regions of symptomatic (13-month-old male) ADtg mice following s.c. GA immunization (once a week for 2 months) or combined treatment of weekly s.c. GA plus monthly i.v. injections of BM-derived CD115⁺ monocytes to the peripheral blood (for 2 months; GA + Mo^{BM}) in comparison with OPN levels in ADtg mice that underwent i.v. PBS injection (once a month for 2 months; control group). Representative micrographs of coronal brain sections covering HC and CC regions reveal elevated OPN immunoreactivity along with reduced 6E10⁺-Aβ plaque burden in GA- and GA + Mo^{BM}-immunized mice in comparison with PBS controls (Fig. 2A). Similar results were observed in EC (not shown).

Quantitative IHC indicated that while 6E10⁺-Aβ plaque burden diminished in all AD-associated brain regions (HC, CC, EC) of GA-immunized groups (GA and GA + Mo^{BM}) in contrast to that of PBS controls (Fig. 2B), OPN expression increased, especially in cortical regions following combined GA + Mo^{BM} treatment (Figs. 2C and S3A). A quantitative ELISA analysis of OPN protein levels in brain homogenates confirmed that both treatment groups (GA, GA + Mo^{BM}) expressed significantly increased (1.4- and 2.1-fold) OPN compared with PBS control ADtg mice (Fig. 2D; *p* < 0.001). Further, to evaluate changes in OPN immunomodulator levels per inflammatory plaque lesion site, we next analyzed the ratio of OPN expression against area of 4G8⁺ or 6E10⁺ plaque burden (Fig. 2E and S3B, C). Specifically, the ratios of OPN-positive area to Aβ plaque area in EC, CC, HC and total brain regions were calculated. Data indicated greater increases in OPN expression (2.3- and 4.2-fold, respectively) in GA and GA + Mo^{BM} groups versus the PBS-control group (Fig. 2E; *p* < 0.01 – 0.0001). Notably, OPN expression in cortical regions (CC, EC) increased more substantially than in the hippocampus following GA immunization (Fig. S3B). This elevation of OPN levels within Aβ plaques following combined treatment (GA + Mo^{BM}) was consistently larger than that resulting from GA-immunization alone (Figs. 2E and S3C). To evaluate the relationship between OPN and Aβ levels in the brain, we analyzed paired parameters from individual mice using the Pearson's coefficient (*r*) correlation test (Figs. 2F–G and S3D–F). These analyses indicated a strong inverse relationship between OPN and Aβ plaque burden (Figs. 2F and S3D; Pearson *r* = −0.65, *p* = 0.008), especially in cortical regions. Furthermore, total OPN protein levels were inversely related to insoluble Aβ₄₂ (Fig. 2G; *r* = −0.73, *p* = 0.017) but not to soluble Aβ₄₂ (Fig. S3F). Together, these data demonstrate that upregulation of cerebral OPN by GA immunization, with or without monocyte blood enrichment, is concomitant with a reduction of parenchymal insoluble Aβ deposits in ADtg mice.

Cerebral amyloid angiopathy (CAA), a complex pathological feature found in most AD patients, involves vascular changes and cerebrovascular amyloid deposition (Ellis et al., 1996), where

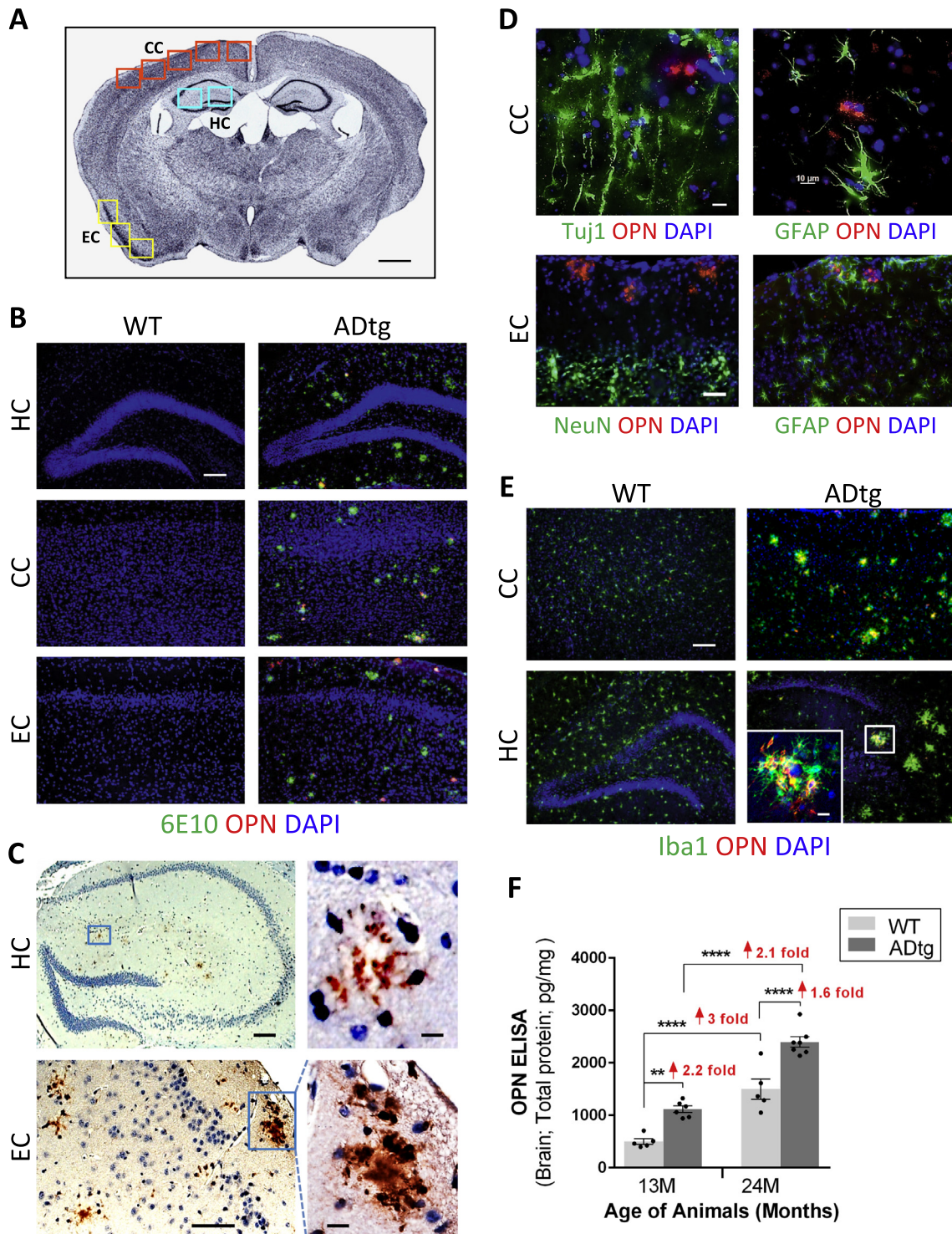


Fig. 1. OPN expression patterns in brains of ADtg and WT mice. (A) Quantification scheme for coronal brain sections. Representative Nissl-stained image of mouse brain at Bregma-2.65 mm. Scale bar: 1 mm. Cingulate cortex (CC), hippocampus (HC) and entorhinal cortex (EC) were included in subsequent quantitative analyses. (B) Representative fluorescent micrographs of brains from ADtg and age-matched WT mice, immunolabeled for anti-OPN (red), anti-human A β (6E10; green), and nuclei (DAPI, blue). OPN immunostaining was detected within and around A β plaques in ADtg mice in all AD-associated brain regions (HC, CC and EC). In WT animals, no A β or OPN immunolabeling was detected. (C) Photomicrographs of brain sections (HC and EC) from ADtg mice show OPN immunohistochemistry by labeling with horseradish peroxidase (HRP)-conjugated secondary antibody. OPN is abundant in layers II/III of EC and often forms plaque-like structures. (D) Representative fluorescent micrographs of ADtg mouse brain sections (CC and EC) display immunostaining for OPN (red) and neuronal markers, Tuj1 or NeuN, or the astrocyte marker, GFAP. (E) OPN (red) was expressed by a subset of Iba1⁺ cells (green) in ADtg mice but not in WT-mice. Scale bars: 100 μ m, inserts: 10 μ m. (F) Quantitative ELISA analysis of OPN levels in brain lysate from ADtg and WT littermate groups at 13 and 24 months of age (equal numbers of females and males). n = 5–6 mice per group. Fold changes indicated in red. Group means, SEMs and individual data points are shown. **p < 0.01, ****p < 0.0001, by one-way ANOVA and Tukey's multiple comparisons post test.

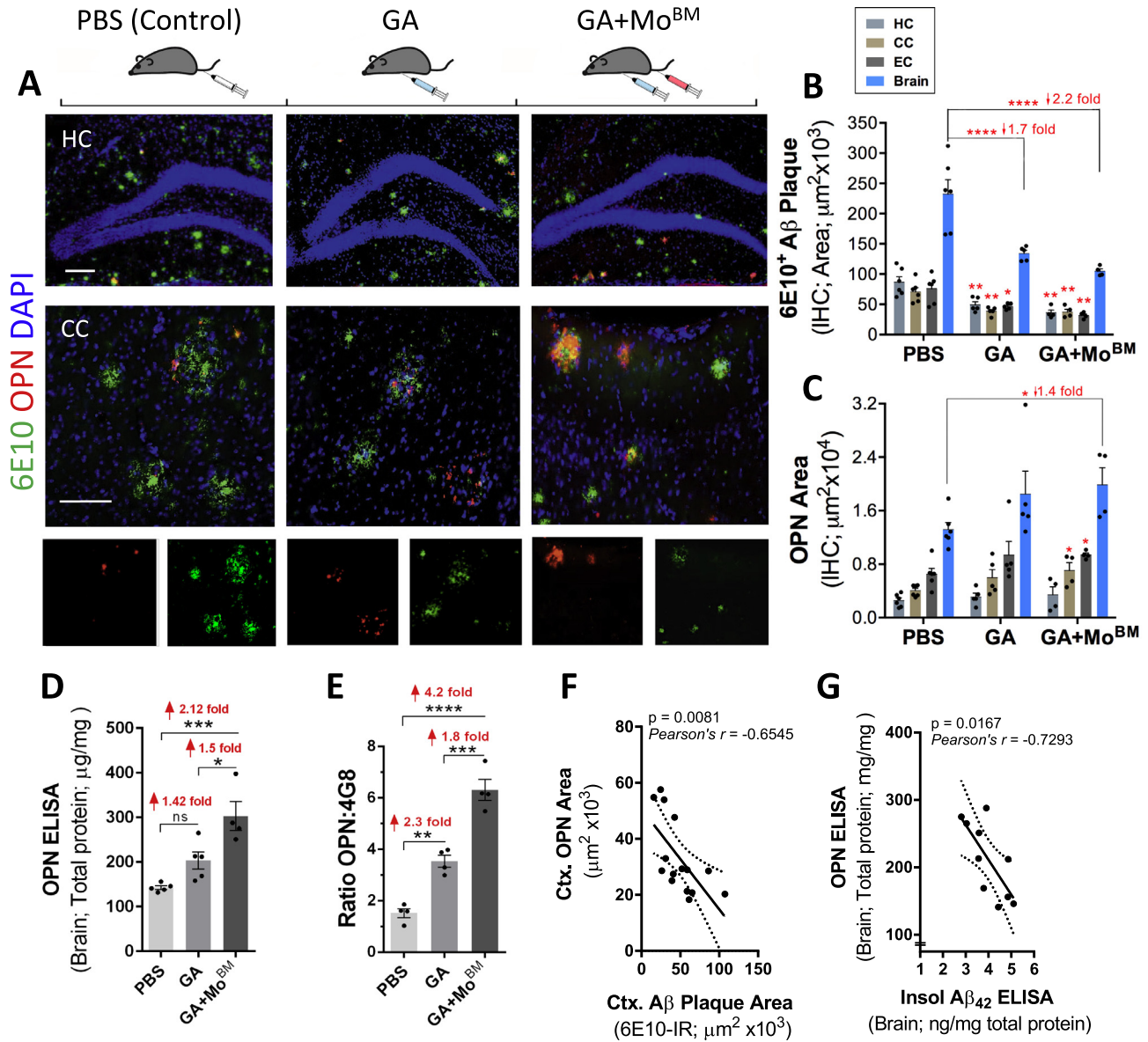


Fig. 2. Increased cerebral OPN expression associated with decreased A β plaque burden. (A) Fluorescent micrographs of coronal brain sections from HC (20 \times magnification, top panel) and CC (63 \times , lower panel) regions display OPN (red) and A β (6E10, green) aggregation patterns for all experimental ADtg mouse groups: s.c. GA immunization, GA + Mo^{BM} [GA s.c. immunization combined with i.v. adoptive transfer of bone marrow-derived CD115⁺ monocytes (Mo^{BM})], and i.v. PBS injection as a control group (n = 4–6 male mice/group). Nuclei stained with DAPI. Lower row displays single channels of OPN (red) and 6E10⁺ (A β , green). Scale bars: 100 μ m. (B, C) Quantitative IHC analyses of 6E10⁺-plaque area and OPN immunoreactive area in brains of ADtg mice (n = 4–6 mice/group). Data by region (HC, CC, and EC) accompanies total brain levels. (D) Quantitative ELISA analysis of OPN levels in brain lysates from all ADtg mouse groups (n = 4–5 mice/group). (E) Ratio between cerebral OPN⁺ area and 4G8⁺-plaque area for all experimental groups (n = 4 mice/group). (F, G) Correlation analyses between cerebral OPN and A β burden: (F) Inverse correlation between cortical (Ctx.) OPN⁺ area and 6E10⁺-A β plaque burden, and (G) inverse correlation between cerebral OPN and insoluble A β ₄₂ levels (ELISA). Correlation analyses performed using Pearson's coefficient (r) test (n = 10–15 ADtg mice). Fold changes indicated in red. Group means, SEMs and individual data points are shown. *p < 0.05, **p < 0.01, ***p < 0.001, ****p < 0.0001, by one-way ANOVA and Tukey's or Dunnett's multiple comparisons post-tests (B–E) for comparison between experimental groups and PBS control or by two-tail unpaired Student's t-test for analysis between two groups (C).

A β ₄₀ is the principal amyloid constituent (Herzig et al., 2006). We examined A β deposition in cerebral vessels by staining cortical and hippocampal brain regions with Thioflavin S (Thio-S) and further determined CAA severity scores (Fig. 3A, B), as previously described (Wyss-Coray et al., 2001). Results revealed that GA immunization significantly decreased vascular amyloid burden by 2.2 times (Fig. 3B; p < 0.001) as compared to PBS-treated ADtg mice. Combined treatment of GA immunization and Mo^{BM} also effectively cleared vascular A β deposits (Fig. 3B; 4.1-fold reduction, p < 0.0001). To determine the relationship between CAA severity and soluble/insoluble A β ₄₀ and A β ₄₂ alloforms, we performed mul-

tipole correlation analyses using the Pearson's coefficient (r) test (Fig. 3C–F). We found a highly significant direct association between CAA and levels of A β ₄₀ peptide, both soluble and insoluble, in the brains of ADtg mice (Fig. 3C, D; r > 0.76, p = 0.0001 – 0.0002). While a positive correlation between CAA and insoluble A β ₄₂ levels was observed (Fig. 3E), no such correlation was found with soluble A β ₄₂ (Fig. 3F). We next analyzed the relationship between OPN and vascular-associated amyloid deposition (Fig. 3G–I), and found statistically significant evidence that cerebral OPN levels are inversely related to both vascular amyloid burden (Fig. 3G; r = 0.72, p = 0.008) and soluble/insoluble A β ₄₀

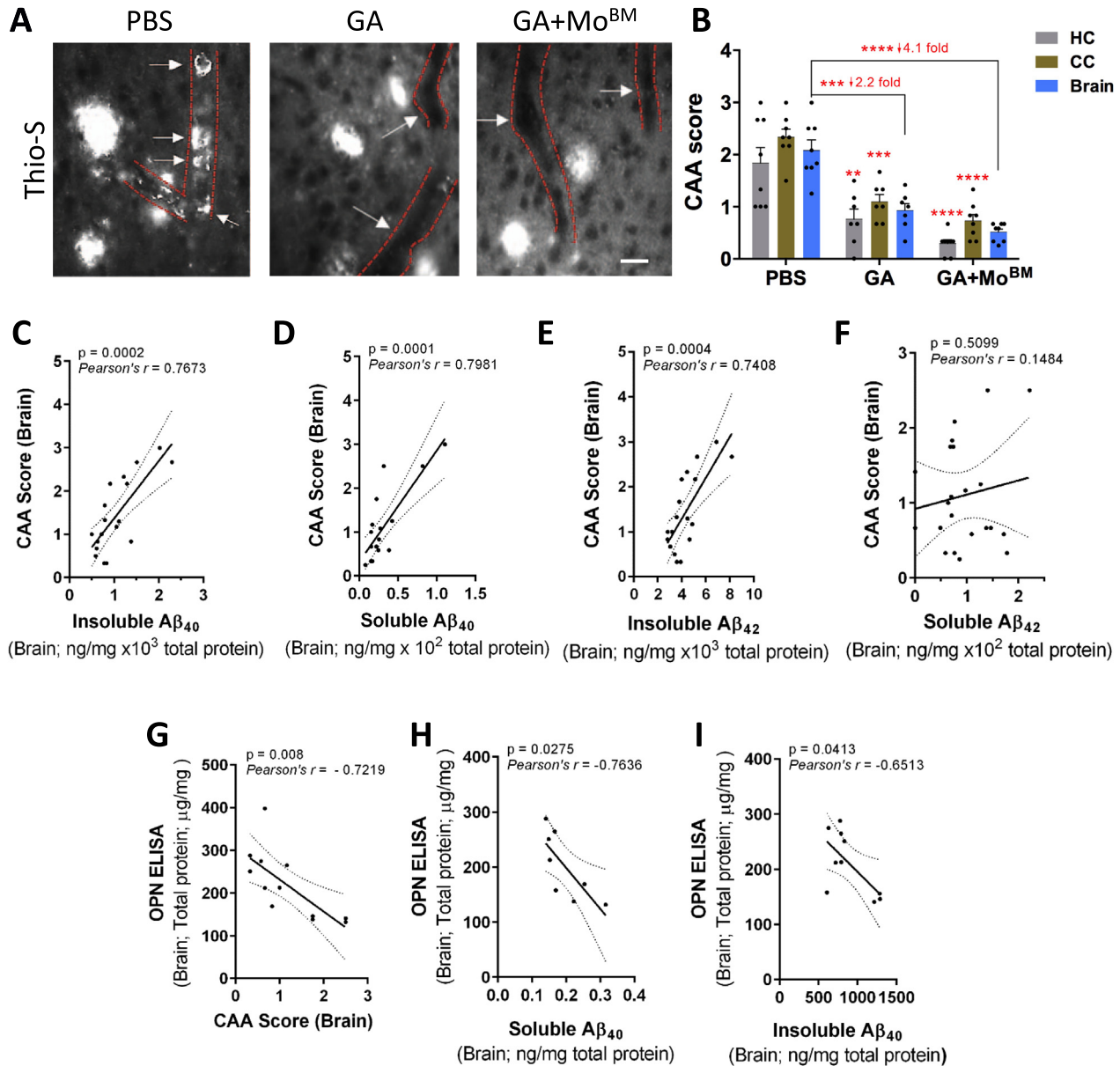


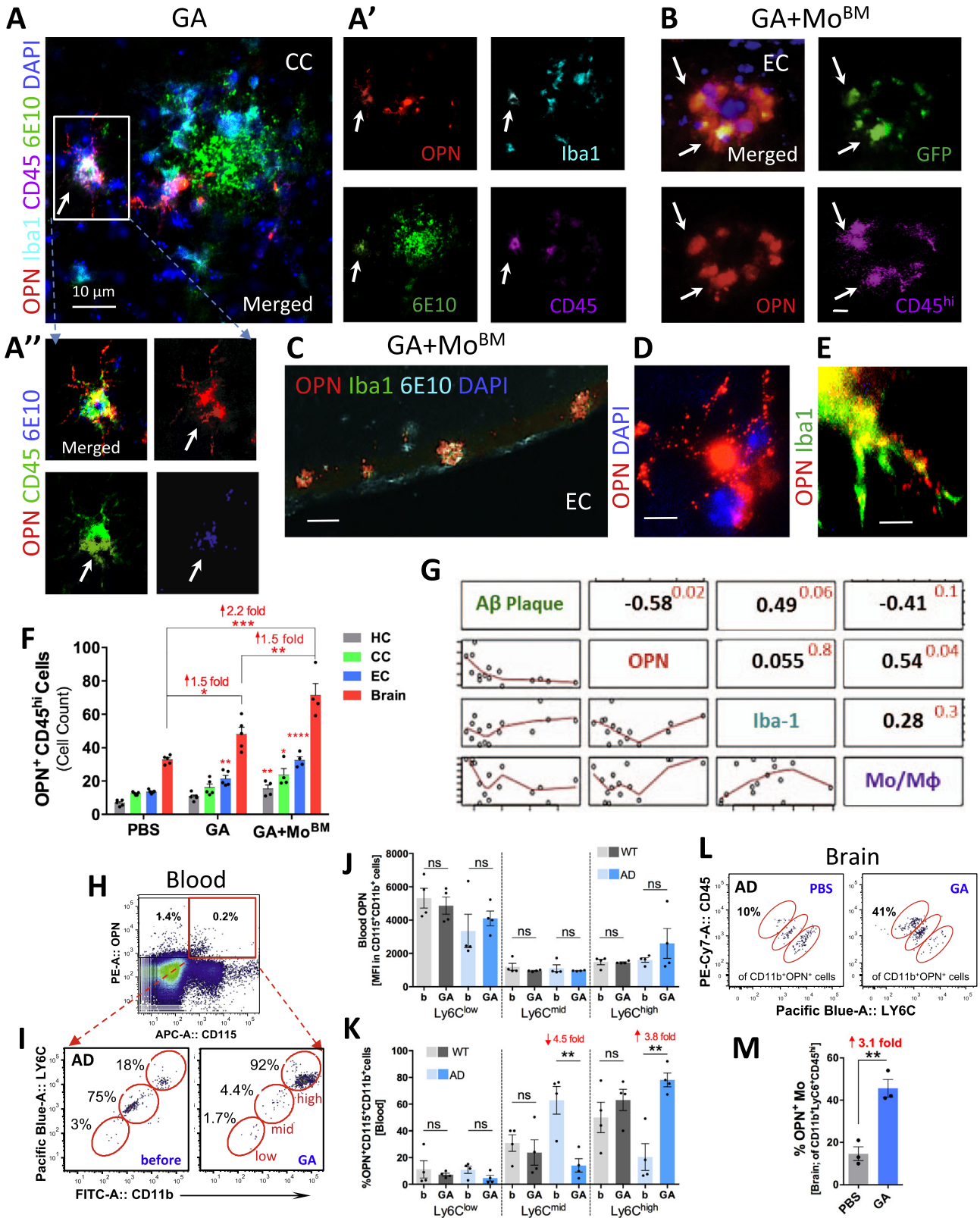
Fig. 3. Reduced cerebral vascular amyloid and A β_{40} burden correlate with increased levels of cerebral OPN. (A) Representative photomicrographs show Thioflavin-S (Thio-S)⁺ amyloid burden in brain parenchyma and vasculature; arrows and red delimitations indicate vascular A β deposits in respective cortical regions of GA-immunized (with and without grafted monocytes) vs. control (PBS-injected) ADtg mice. Thio-S staining revealed not only lesser parenchymal plaque burden in GA immunized groups but also reduction of vascular A β deposits. (B) Quantitative Thio-S⁺ CAA score assessment by brain region (HC and CC) and average in ADtg mouse groups (n = 7–8 mice/group). (C–F) Correlation analyses reveal strong, significant linear correlations between vascular CAA burden and both (C) insoluble and (D) soluble A β_{40} levels in the brain (ELISA) as well as with (E) cerebral insoluble A β_{42} , but no correlation with (F) soluble A β_{42} levels (ELISA). (G–I) Inverse correlations between cerebral OPN levels and (G) CAA scores, (H) soluble A β_{40} , and (I) insoluble A β_{40} in ADtg mouse brains (ELISA). Correlation analyses performed using Pearson's coefficient (r) test (n = 10–15 ADtg mice). Fold changes indicated in red. Group means, SEMs and individual data points are shown. ** $p < 0.01$, *** $p < 0.001$, **** $p < 0.0001$, by one-way ANOVA and Dunnett's multiple comparisons post-tests.

(Fig. 3H, I; $r > 0.65$, $p < 0.05$). Collectively, these results suggest that GA immunization, and to a greater degree its combination with monocyte blood enrichment, leads to an upregulation of cerebral OPN that is tightly associated with reduced vascular and parenchymal A β burden in ADtg mice.

3.3. OPN is predominantly expressed by infiltrating monocyte-derived macrophages involved in A β clearance

To identify OPN-expressing cells in ADtg mouse brains, we used two different approaches (Koronyo et al., 2015). To distinguish infiltrating monocytes/macrophages (Mo/M Φ) from microglia, we

first used the combined Iba1 and CD45 markers (Fig. 4A). Fluorescent IHC analysis revealed that OPN is selectively expressed by infiltrating Iba1⁺CD45^{high} Mo/M Φ in contrast to Iba1⁺CD45^{int-low} resident microglia (Fig. 4A, merged image; separate channels in Fig. 4A'). These infiltrating OPN-expressing Iba1⁺CD45^{high} Mo/M Φ accumulated at and around plaque lesion sites, and the localization of A β within these cells (Fig. 4A, arrow; enlarged framed image in Fig. 4A'') suggests their direct involvement in A β phagocytosis. Brain-infiltrating monocytes expressing OPN were also identified in GA + Mo^{BM}-treated ADtg mice through adoptive transfer of GFP-labeled BM-derived CD115⁺-Mo into the peripheral blood (Fig. 4B). In EC, GFP⁺CD45^{high} Mo/M Φ frequently



expressed high levels of OPN, in contrast to resident microglia (GFP^{neg}CD45^{intermediate-low}). Arrows indicate the localization of OPN in GFP⁺CD45^{high} Mo/MΦ in Fig. 4B. Notably, Mo/MΦ expressing high OPN were often located in EC and clustered at plaque lesion sites (Fig. 4C). We also examined the subcellular localization of OPN within infiltrating Mo/MΦ. OPN showed a punctate staining pattern in vesicular compartments and across the cell body and processes (Fig. 4D, E). OPN expression was particularly strong in the perinuclear subregion (Fig. 4D), apparently in the trans-Golgi network compartments (not shown; further in vitro examination below revealed OPN colocalization with a Golgi biomarker).

Next, we quantified OPN⁺CD45^{high} cells manually in AD-associated brain regions. Substantial 1.5- and 2.2-fold increases in number of infiltrating OPN⁺CD45^{high} cells, particularly in EC, were respectively detected in GA and GA + Mo^{BM} groups in contrast to PBS controls (Fig. 4F; $p < 0.01 - 0.0001$). Multifactor correlogram analyses confirmed that OPN expression inversely relates to Aβ plaque burden and associates directly with brain-infiltrating Mo/MΦ (Figs. 4G and S4A). Similarly, Pearson's correlation analysis demonstrated a significant direct linear relationship between cerebral OPN expression and recruited Iba1⁺CD45^{high} Mo/MΦ in ADtg mice (Fig. S4B; $r = 0.57$, $p = 0.025$). We further found a tighter association between expression of the OPN cytokine and brain-infiltrating CD45^{hi} myeloid cells per plaque (Fig. S4C; $r < 0.61$, $p = 0.0001$).

Previously, we found that our immunomodulation approach altered the immunological environment in brains of ADtg mice, characterized by elevation of the Aβ-degrading enzyme matrix metalloproteinase (MMP)-9 (Bakalash et al., 2011; Butovsky et al., 2006; Koronyo et al., 2015; Koronyo-Hamaoui et al., 2009) as well as the chemokine monocyte chemoattractant protein-1 (MCP-1/CCL2), a key regulator of Mo/MΦ migration into the brain (El Khoury et al., 2007). We evaluated whether changes to these factors relate to cerebral OPN levels through Pearson (r) correlation analyses (Fig. S4D–G), and our data suggest strong positive associations between levels of OPN and MCP-1 (Fig. S4D; $r = 0.73$, $p < 0.03$), as well as between OPN area or protein levels and MMP-9 (Fig. S4E, F; $r = 0.62$, $p < 0.05$). Cortical OPN-expressing CD45^{hi} Mo and MMP-9 protein levels also show a tight linear correlation (Fig. S4G; $r = 0.74$, $p = 0.0016$).

To assess how GA affects OPN expression in peripheral blood monocytes, an additional cohort of adult ADtg mice and age- and gender-matched WT littermates underwent flow cytometry analysis of peripheral blood before and after GA immunization (Figs. 4H–K and S5A–H). Immunolabeling of blood monocytes with

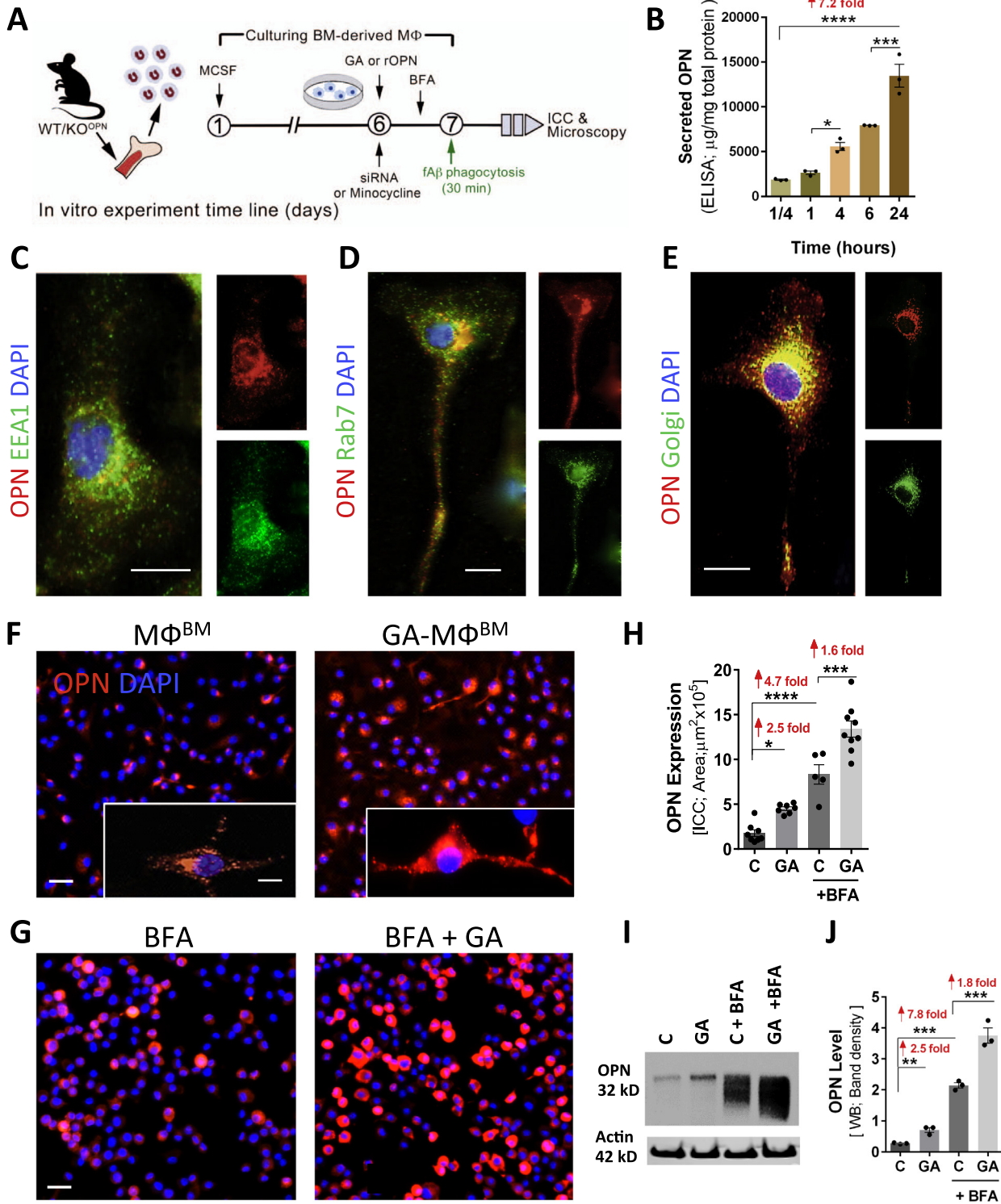
several biomarkers (CD11b, CD115, and Ly6C) allowed detection of OPN-expressing monocyte subpopulations. CD115⁺OPN-expressing monocytes represent a small ~0.2% population of murine blood cells (Fig. 4H, red box). In general, there were no significant differences in percentage of total blood cell population or mean OPN expression (MFI) in blood monocytes following GA immunization in both WT and ADtg mice (Fig. S5B, C). Further gating of OPN⁺CD115⁺ into three monocyte subpopulations, CD11b⁺Ly6C^{low}, CD11b⁺Ly6C^{medium}, and CD11b⁺Ly6C^{high} showed no differences in MFI values before and after GA immunization in either genotype (Fig. 4J). However, significant differences in percentage of CD11b⁺Ly6C^{medium} and Ly6C^{high} monocytes were detected after GA immunization in ADtg mice (Fig. 4K). Following GA immunization, a shift occurred in subpopulations of CD115⁺CD11b⁺Ly6C^{mid} and CD115⁺CD11b⁺Ly6C^{high} monocytes, with the percentage of the former reducing 4.5 times and the latter inflammatory group growing 3.8 times in ADtg mice (Figs. 4K and S5D, E; $p < 0.01$). Examination of CD115^{neg}CD11b⁺ cells that express OPN and exhibit Ly6C^{low}, Ly6C^{medium}, or Ly6C^{high} (Fig. S5F) in WT and ADtg mice indicated that GA immunization did not affect these populations in the blood (Fig. S5G, H). While our data indicate no change to average OPN expression in blood monocytes, they do demonstrate an increase in the OPN-expressing inflammatory CD115⁺CD11b⁺Ly6C^{high} monocyte subpopulation following GA immunization, specific to ADtg mice.

We have previously (Koronyo et al., 2015) and above (Fig. 4A–F) demonstrated that the inflammatory monocyte subpopulation increasingly infiltrates the brain parenchyma and migrates to amyloid plaque sites in GA-immunized ADtg mice. Flow cytometry analysis of infiltrating OPN-expressing monocytic cells (CD11b⁺Ly6C⁺CD45^{high}; Figs. 4L and S5I) in brain tissues of ADtg mice ($n = 6$ mice) indicated a significant 3.1-fold increase in their percentage in the brains of GA-immunized as compared to PBS-injected mice (Fig. 4M; $p < 0.01$). Collectively, we report that GA immunomodulation, with and without monocytes, increases brain-recruitment of OPN-expressing Mo/MΦ tightly associated with Aβ-plaque clearance in ADtg mice.

3.4. OPN expression patterns in primary cultures of BM-derived macrophages

To investigate how OPN impacts macrophage phenotype and its role in macrophage-mediated Aβ phagocytosis and degradation, we further studied OPN loss- vs. gain-of-function in BM-derived MΦ exposed to pathological forms of Aβ. The in vitro experimental

Fig. 4. Selective expression of OPN by blood monocytes and brain-infiltrating monocyte/macrophages involved in Aβ uptake following GA immunization. (A–E) Fluorescent micrographs of cortical regions of GA-immunized ADtg mice, with and without grafted GFP⁺-Mo^{BM} (GA + Mo^{BM}). Scale bars: 10 μm unless otherwise indicated. (A) Infiltrating Mo/MΦ identified by combined Iba1⁺ and CD45^{high} immunolabeling in GA-immunized mouse brains. Cerebral OPN, Aβ (6E10) and nuclei (DAPI) also shown. (A') Images of separate channels; arrow indicates an infiltrating OPN-expressing Iba1⁺CD45^{high} MΦ engulfing Aβ. (A'') Region of interest in box above illustrates the colocalization of Aβ (6E10) with OPN-positive Iba1⁺CD45^{high} MΦ. (B) Brain-infiltrating monocytes expressing OPN were also identified in GA + Mo^{BM}-treated mice through a second approach: adoptive transfer of a GFP-labeled BM-derived CD115⁺-monocyte subset injected into the tail vein of symptomatic ADtg mice. In EC, GFP⁺CD45^{high} Mo/MΦ expressed OPN abundantly, in a manner distinct from resident microglia (GFP^{neg}CD45^{intermediate-low}). Arrows indicate colocalization of OPN and GFP⁺CD45^{high} Mo/MΦ. (C) OPN-expressing Iba1⁺ myelomonocytic cells were associated with plaques, especially at the border of EC. Scale bar = 50 μm. (D, E) Puncta staining patterns of subcellular OPN in vesicular compartments across cell body and along Iba1⁺ processes, with particularly intense signal in the perinuclear subregion. (F) OPN⁺CD45^{high} cell count in HC, CC, EC and combined brain regions ($n = 4-5$ ADtg mice/group). Fold increases relative to PBS controls or GA-treated group indicated in red. (G) Quantitative multifactor correlogram analysis (Pearson's test) demonstrated significant linear association between OPN and monocyte infiltration in ADtg mice, and inverse relations to Aβ plaque burden ($n = 15$ mice). (H–K) Flow cytometry analysis on blood monocytes isolated from additional cohort of adult ADtg and age- and gender-matched WT littermates before and after GA immunization ($n = 4$ mice/genotype). (H) Initial gating of CD115-positive cells expressing OPN in red box. (I) OPN⁺ CD115-positive blood monocyte subpopulation in red box further analyzed by flow cytometry using monocyte biomarkers CD11b and Ly6C. (J, K) The OPN-expressing CD11b⁺CD115⁺Ly6C monocyte subpopulation in the blood was gated into three groups, Ly6C^{low}, Ly6C^{mid} and Ly6C^{high}, showing differences in percentage of total cells before and after GA immunization in ADtg mice. Quantitative analysis of (J) mean fluorescence intensity (MFI) values and (K) proportions of cells in percentage before (b) and after immunization (GA). (L, M) Flow cytometry analysis on myelomonocytes isolated from perfused brains of GA-immunized vs. PBS-injected ADtg mice ($n = 3$ mice/group). (L) Final gating of OPN⁺CD11b⁺ myelomonocytes in the brain into three populations of Ly6C⁺CD45^{hi}, Ly6C⁺CD45^{mid} and Ly6C⁺CD45^{low} (shown in red ellipses). (M) Percentage of OPN⁺CD11b⁺ Ly6C⁺CD45^{hi} infiltrating monocytes in the brain of GA vs. PBS control ADtg mice. Group means, SEMs and individual data points are shown. NS, no statistical significance. * $p < 0.05$, ** $p < 0.01$, *** $p < 0.001$, **** $p < 0.0001$, by one-way (F) or two-way (J–K) ANOVA followed by Dunnett's or Bonferroni's multiple comparisons post tests or two-tail unpaired Student's t -test (M).



paradigm included isolation of BM from young WT or OPN-null (OPN^{-/-} knockout; hereafter referred to as KO^{OPN}) donor mice, their differentiation into mature macrophages (MΦ^{BM}), and a series of experiments involving treatments such as GA (GA-MΦ^{BM}), recombinant OPN (rOPN), and/or siRNA inhibition, followed by a 30-min exposure to Aβ₄₀ and Aβ₄₂ fibrils (Fig. 5A). We initially explored secretion patterns of the cytokine OPN in MΦ^{BM} media over the course of 24 h (Fig. 5B). Our analysis suggests that OPN is constitutively expressed and secreted to the extracellular space by MΦ^{BM}, is detectable even at the 15-min time point, and accumulates in substantial amounts over a full-day period (Fig. 5B; 7.2-fold increase, $p < 0.0001$). Next, we examined the patterns of intracellular OPN expression in MΦ^{BM} using various antibodies recognizing subcellular biomarkers (Figs. 5C–E and S6A). In MΦ^{BM} cultures, OPN often displayed intense perinuclear immunolabeling and punctate patterns across the cell body and processes, suggesting that OPN was located in transport vesicles (Fig. 5C–E). OPN did not typically colocalize with early endosomes (early endosomal antigen EEA1/Rab5a; Fig. 5C), late endosome/lysosomes (Ras-related protein Rab7; Fig. 5D), or autophagosomes (light chain 3 protein LC3; Fig. S6A): biomarkers representing internalization/uptake vesicles fated for intracellular degradation. However, OPN was mostly localized within the Golgi apparatus, part of a major secretory pathway, as detected via the anti-Golgi marker 58K protein antibody (Fig. 5E), implying that the cytokine is constantly expressed and sorted through the trans-Golgi network (TGN) for post-translation modifications prior to secretion.

We subsequently assessed the impact of GA pretreatment on OPN expression in MΦ^{BM} cultures in the absence of Aβ fibrils (Figs. 5F–J and S6B). This revealed that GA significantly induced OPN expression in MΦ^{BM}, mirroring our *in vivo* results, and further increased when TGN-derived secretion vesicles were inhibited by Brefeldin A (BFA; Fig. 5F vs. G) (Rosa et al., 1992). While MΦ^{BM} without BFA showed typical elongated morphology and a punctate/vesicular OPN signal (Fig. 5F, inserts), MΦ^{BM} undergoing BFA blockage of TGN-derived protein secretion showed decreased cell adhesion with a rounded morphology, and newly-synthesized OPN trapped within (Fig. 5G). The BFA condition showed substantial accumulation of OPN in MΦ^{BM} in short periods of time (i.e. within 1–3 h). Quantitative immunocytochemistry (qICC) confirmed significant 1.6- and 2.5-fold increases in OPN production/expression (with and without BFA, respectively) in GA-MΦ^{BM} vs. untreated MΦ^{BM} (Fig. 5H). A quantitative ELISA analysis and Western blot assessments of OPN in cell lysates validated that regardless of BFA condition, GA markedly increased OPN expression and production 1.4–2.5 times in MΦ^{BM} (Figs. 5I, J and S6B; $p < 0.01 - 0.001$). Taken together, our data indicate that the immunomodulator OPN is constitutively expressed and secreted by MΦ^{BM} and that GA substantially increases its levels.

3.5. Phagocytosis of fibrillar Aβ is OPN dependent

To study whether OPN gain-of-function induces MΦ^{BM} phagocytosis of pathogenic Aβ aggregates, we first examined whether increased OPN expression via GA is associated with improved phagocytosis of Aβ₄₀ and Aβ₄₂ fibrils. Representative high-magnification Z-stack images demonstrate that intracellular OPN expression in MΦ^{BM} is accompanied by phagocytosis of fibrillar (f)Aβ₄₂ (Fig. 6A). MΦ^{BM} were then pretreated with GA (30 μg/mL, overnight) and exposed to fAβ₄₂ for 30 min. Enhanced fAβ₄₂ uptake was associated with increased OPN expression in GA-MΦ^{BM} when compared with untreated MΦ^{BM} (Fig. 6C vs. B). Representative images show a distinct pattern of OPN immunoreactivity within somas and along processes of CD68⁺ (a phagocytic macrophage marker) phagocytes, with strong perinuclear and evenly distributed vesicular signals, accompanied by separated punctate Aβ labeling. This staining was substantially intensified after GA treatment (Fig. 6A–C). Exposure of MΦ^{BM} to either form of fAβ for 30 min showed that fAβ₄₀ yielded an intracellular signal stronger than that of fAβ₄₂ (Fig. 6D vs. E). Increased OPN expression (2.7-fold) along with augmented fAβ₄₀ and fAβ₄₂ uptake by MΦ^{BM} (2.3- and 2.9-fold, respectively) were detected by qICC analysis (Fig. 6F–H; $p < 0.001 - 0.0001$). Quantitative ELISA analysis of cell lysates validated the qICC results, indicating increased intracellular OPN (2.3-fold; Fig. S6B) with Aβ₄₂ uptake elevated 2.5-fold in GA-treated vs. untreated MΦ^{BM} (Fig. S6C; $p < 0.001$).

To study Aβ phagocytosis in OPN-deficient MΦ, we first knocked down OPN expression using RNA interference (siRNA) (Fig. 6I–K). Previous studies have suggested that intracellular OPN deletion in macrophages impairs general phagocytic activity (Schack et al., 2009; Shin et al., 2011; Toyonaga et al., 2015). Here, to assess OPN involvement in the specific phagocytosis of fAβ, MΦ^{BM} were first transfected with either siRNA^{OPN} or control siRNA^{Neg}, and 48 h later they were exposed to Aβ fibrils. Fluorescent micrographs revealed a substantial reduction of OPN that was associated with decreased fAβ₄₂ uptake by CD68⁺MΦ^{BM} due to siRNA-OPN deficiency (Fig. 6I). The qICC analyses validated our observations and indicated OPN inhibition of 6.8 fold associated with a 2.7 fold reduction in intracellular uptake of fAβ₄₂ (Fig. 6J, K; $p < 0.0001$). Negative control scrambled siRNA did not affect OPN expression or Aβ phagocytosis. The substantial OPN inhibition by siRNA was also confirmed by analysis of protein levels in MΦ^{BM} cell lysates using quantitative ELISA (Fig. S6D). Importantly, GA treatment failed to increase OPN expression and did not impact MΦ^{BM} phagocytosis of Aβ fibrils when OPN was knocked down (Fig. 6J, K). Furthermore, taking advantage of a genetic knockout OPN mouse model (KO^{OPN}) allowed us to study fAβ₄₀ and fAβ₄₂ uptake in OPN-deficient MΦ^{BM} (Fig. 6L, M). Our qICC data suggest that, regardless of GA treatment, phagocytosis of Aβ fibrils is impaired

Fig. 5. Patterns of expression, secretion, and GA-mediated upregulation of OPN in BM-derived macrophage cultures. (A) Schematic illustration of *in vitro* studies: BM was isolated from WT mice (8- to 12-weeks-old) and cultured for 6 or 7 days in MCSF-enriched media to differentiate into macrophages (MΦ^{BM}). On day 6, cells underwent overnight treatment with GA, siRNA or minocycline, except for untreated control cells (labeled 'C'). On day 7, fibrillar Aβ (fAβ₄₀ or fAβ₄₂) was added in a subset of experiments, followed by phagocytosis assays. Brefeldin A (BFA) treatment was performed 3 h prior to phagocytosis. (B) Constitutive secretion of OPN by MΦ^{BM} during a 24 h period. Primary MΦ^{BM} media were collected after 15 min, and then after 1, 4, 6, and 24 h. OPN protein levels were measured at each time point (ELISA; $n = 3$ wells/time point, 1×10^6 cells/well, in triplicates). (C–E) Intracellular OPN expression in MΦ^{BM}. Scale bars = 10 μm. Representative fluorescent micrographs of MΦ^{BM} immunostained for OPN, (C) early endosomal antigen (EEA1) marker, (D) late endosome-lysosomal marker Ras-related protein (Rab7), or (E) Golgi marker 58 K protein, and nuclei (DAPI). The merged images demonstrate subcellular OPN expression within vesicles, predominantly confined to the trans-Golgi network (yellow punctate signal). (F–J) Upregulation of OPN in MΦ^{BM} by GA treatment. (F, G) Representative fluorescent micrographs demonstrate the effect of GA treatment on MΦ^{BM} expression of OPN in the absence of Aβ, either (F) without BFA treatment or (G) with BFA pre-treatment. MΦ^{BM} were immunostained for OPN (red), which was more highly expressed following GA treatment. (F, insert) Subcellular OPN within transport vesicles in MΦ^{BM}. (G) Round-shaped MΦ^{BM} after BFA inhibition of OPN secretion. Scale bars: 20 μm, insert scale 5 μm. (H) Quantitative ICC of OPN-immunoreactive area revealed a significant upregulation of OPN in MΦ^{BM} following GA treatment, with or without BFA inhibition. Means of individual cell fluorescent areas are indicated ($n = 5-9$ images/well, average 100 cells/image, and $n = 3-5$ wells/treatment group). (I) Western blot image of cell lysates from above-mentioned experimental groups. (J) Corresponding densitometry results of OPN levels in Western blots (normalized to β-actin levels; $n = 3$ wells/group, 1×10^6 cells/well; repeated experiments). Group means, SEMs and individual data points are shown. Fold increases in mean values compared with controls indicated in red. * $p < 0.05$, ** $p < 0.01$, *** $p < 0.001$, **** $p < 0.0001$, by one-way ANOVA and Dunnett's multiple comparisons post-tests.

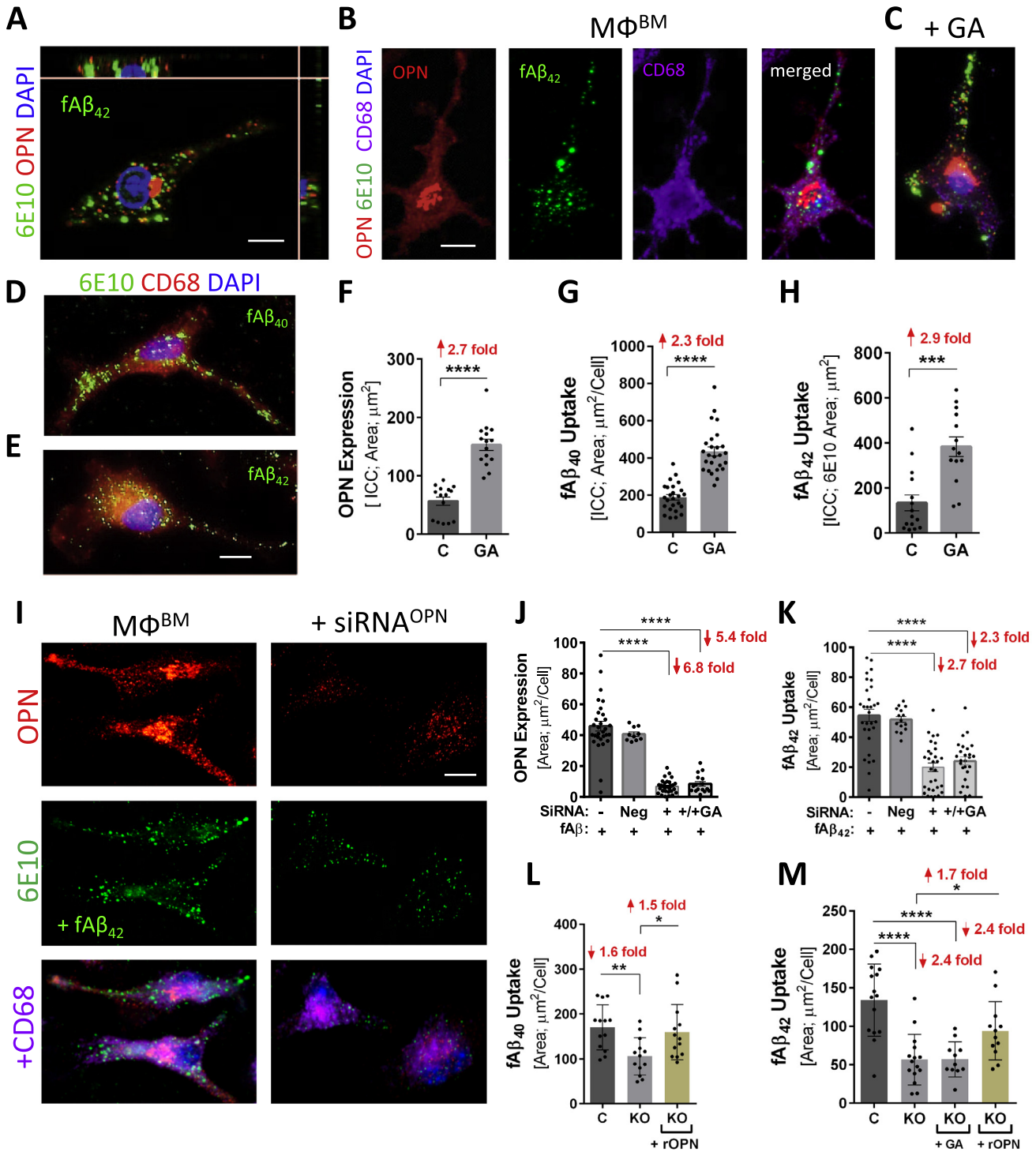
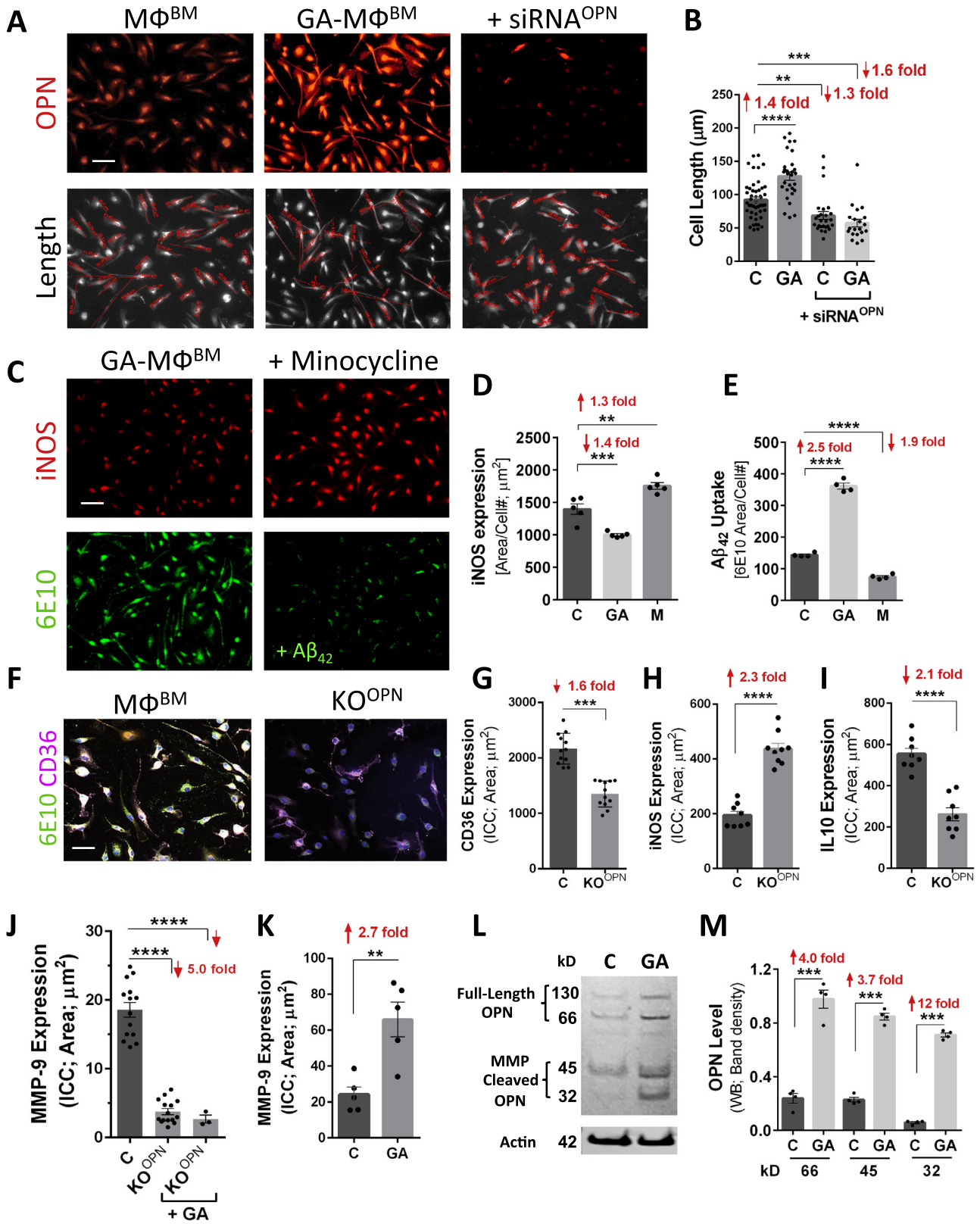


Fig. 6. Phagocytosis of Aβ₄₀ and Aβ₄₂ fibrils by BM-derived macrophages is dependent on OPN expression. (A–H) Representative fluorescent micrographs and quantitative analyses of OPN expression and Aβ uptake in CD68⁺ MΦ^{BM}, in primary cultures pre-treated with GA for 24 h and stimulated with fibrillar (f)Aβ₄₀ or fAβ₄₂ for 30 min. Scale bars: 10 μm. (A) Higher magnification Z-stack image shows intracellular uptake of 6E10⁺-Aβ along with subcellular OPN expression patterns in untreated MΦ^{BM}. (B, C) Increased OPN expression and enhanced cellular uptake of fAβ₄₂ was detected following GA treatment. (D, E) Representative images display phagocytosis of fAβ₄₀ vs. fAβ₄₂ by untreated MΦ^{BM}. (F–H) Quantitative ICC analysis of average OPN expression (F), intracellular fAβ₄₀ (G), and fAβ₄₂ (H) areas per MΦ^{BM} in GA-treated vs. untreated MΦ^{BM}. Along with increased OPN expression, MΦ^{BM} pre-treated with GA exhibit enhanced uptake of Aβ fibrils. (I) Representative micrographs of OPN-silenced MΦ^{BM} (via siRNA^{OPN} knockdown) vs. untreated cells reveal reduced fAβ₄₂ phagocytosis. Scale bar: 10 μm. (J, K) Quantitative ICC revealed that silenced expression of OPN via siRNA^{OPN} leads to impaired fAβ₄₂ phagocytosis in MΦ^{BM}, regardless of GA treatment. Negative control scrambled siRNA^{Neg} affected neither OPN expression nor fAβ phagocytosis. (L, M) Additional experiment utilizing MΦ^{BM} isolated from OPN knockout (KO) mice vs. WT mice (controls, C). Quantitative ICC analysis of intracellular fAβ₄₀ and fAβ₄₂ uptake in control MΦ^{BM}, OPN-deficient MΦ^{BM} (KO), and GA-treated OPN-deficient MΦ^{BM} (KO + GA). OPN-deficient MΦ^{BM} exhibit impaired fAβ phagocytosis, partially restored by supplementation of human recombinant (r) OPN. Means of individual cell fluorescent areas indicated (average of n = 5–10 images per well, with ~100 cells/image, n = 4–6 wells/treatment group). Group means, SEMs and individual data points are shown. Fold increases in mean values compared with controls indicated in red. *p < 0.05, **p < 0.01, ***p < 0.001, ****p < 0.0001, by one-way ANOVA and Tukey's multiple comparisons post-tests or two-tail unpaired Student's *t*-test.



in the absence of OPN (Fig. 6L, M). Notably, addition of full-length rOPN (50 ng/mL, overnight) to KO^{OPN}-M Φ ^{BM} reversed their impairment in cellular uptake of fA β ₄₀ and fA β ₄₂ (Fig. 6L, M; $p < 0.05$). Overall, these results reveal that phagocytosis of pathogenic A β alloforms by M Φ ^{BM} is OPN-dependent, and that upon exposure to GA, it undergoes a gain-of-function mediated by elevated OPN expression.

3.6. OPN plays a key role in macrophage polarization towards an anti-inflammatory phenotype

Macrophages, as well as their circulating counterparts, monocytes, make up a highly heterogeneous population that can acquire various phenotypes and biological functions in response to environmental signals [reviewed in Gliem et al. (2016)]. To further elucidate OPN's effects on M Φ ^{BM}-mediated A β phagocytosis, we investigated whether these effects relate to larger morphological and immunological profile alterations. Past studies have reported that OPN induces M Φ ^{BM} to differentiate into the anti-inflammatory M2 phenotype (Tambuyzer et al., 2012), a polarization associated with elongated cell morphology (McWhorter et al., 2013). Analysis of M Φ ^{BM} cell length under various treatment conditions, including OPN-induction by GA and -silencing by siRNA, revealed significant changes in CD68⁺M Φ ^{BM} cell length as displayed in the representative fluorescent micrographs (Fig. 7A). While GA induced OPN expression, it also increased M Φ ^{BM} cell elongation by 1.4 times (Fig. 7B; $p < 0.0001$). In contrast, siRNA^{OPN} knockdown reversed GA's effects and further decreased cell length by 1.3 times (Fig. 7B; $p < 0.01$). In the absence of OPN, GA did not affect M Φ ^{BM} cell elongation (Fig. 7B; $p < 0.001$), in agreement with our previous A β phagocytosis results (Fig. 6K).

We next assessed the expression of inducible nitric oxide synthase (iNOS), a hallmark of pro-inflammatory M1 macrophages, in different treatment groups; we used the tricyclic antibiotic minocycline to suppress OPN expression (Fig. 7C–E). Minocycline has been shown to modulate inflammatory response, and to reduce MMP-9 activity and generation of proteolytic OPN fragments (Chan et al., 2014). M Φ ^{BM} underwent either OPN-activation by GA or -inhibition by minocycline, and were then exposed to fibrillar A β ₄₂ for 30 min. Fluorescent micrographs demonstrate that GA reduced iNOS expression and increased fibrillar A β ₄₂ phagocytosis, while minocycline induced opposite effects (Fig. 7C). Quantitative ICC assays confirmed a significant 1.3-fold increase in iNOS expression by minocycline-treated macrophages, and a substantial 1.9-fold reduction in A β ₄₂ phagocytosis (Fig. 7D, E; $p < 0.01 - 0.0001$). GA

treatment reduced iNOS 1.4-fold but boosted A β phagocytosis 2.5-fold (Fig. 7D, E; $p < 0.001 - 0.0001$).

Furthermore, we analyzed M Φ ^{BM} phenotype in the genetic OPN-null (KO^{OPN}) models. Downregulation of OPN in the KO^{OPN}-M Φ ^{BM} was accompanied by impaired ability to uptake fA β ₄₂ (Figs. 6L, M and 7F). KO^{OPN}-M Φ ^{BM} showed a significant 1.6-fold reduction in surface expression of the scavenger receptor CD36 (implicated in A β phagocytosis), a 2.3-fold increase in proinflammatory iNOS expression, and 2.1-fold decrease in anti-inflammatory cytokine interleukine-10 expression (Fig. 7G–I; $p < 0.001 - 0.0001$) in comparison with control-M Φ ^{BM}. These studies demonstrate that OPN silencing, either by siRNA, minocycline, or genetic knock-out (KO^{OPN}), all of which impaired macrophages' ability to uptake A β fibrils, was associated with a proinflammatory profile. Importantly, induction of OPN expression by GA induced polarization towards the anti-inflammatory, highly phagocytic macrophage phenotype with elongated pro-healing cell morphology.

In addition to phagocytosis, enzymatic degradation is another major mechanism through which M Φ ^{BM} can clear A β . It has been shown that MMP-9 degrades A β , associates with an alternatively-activated macrophage phenotype, and is regulated by and directly cleaves OPN to generate biologically-active products (Agnihotri et al., 2001; Tan et al., 2013). Quantitative ICC analysis in OPN-deficient (KO^{OPN})-M Φ ^{BM} showed a substantial 5-fold decrease in MMP-9 expression (Fig. 7J; $p < 0.0001$); under OPN loss-of-function conditions, GA treatment failed to reverse MMP-9 levels. In agreement with our previous results (Koronyo et al., 2015), GA treatment significantly increased MMP-9 expression 2.7-fold in WT-M Φ ^{BM} (Fig. 7K; $p < 0.01$); representative Z-stack images show a remarkable increase in MMP-9 expression in GA-treated M Φ ^{BM} (Fig. S6E). Abundant MMP-9 expression, increased fA β phagocytosis, and a 3.1-fold elevation of surface scavenger receptor SCARA-1 are shown in GA-treated M Φ ^{BM} (Fig. S6F, G; $p < 0.0001$). These results mirror our in vivo correlation analysis of ADtg mouse brains, showing a strong linear relationship between OPN and MMP-9 (Fig. S4E, F), which was even tighter between cortex-infiltrating OPN-expressing CD45^{hi} cells and local MMP-9 levels (Fig. S4G).

Moreover, we analyzed OPN protein levels in M Φ ^{BM} using a polyclonal antibody recognizing the full-length OPN and its MMP-cleaved fragments, including the anti-inflammatory 32 kD C-terminus fragment (Chan et al., 2014; Scatena et al., 2007). Western blot analysis and subsequent quantification of band densities revealed that all OPN forms, including fragments derived from MMPs' proteolysis, were elevated by GA treatment (Fig. 7L, M). Interestingly, the MMP-generated 32kD C-terminus-containing

Fig. 7. OPN promotes an anti-inflammatory, pro-healing phenotype in BM-derived macrophages. (A) Fluorescent micrographs of M Φ ^{BM} in primary cultures, either untreated, pretreated overnight with GA, or silenced with siRNA^{OPN}, were immunostained for OPN (upper panel). Lower panel displays grayscale images of CD68⁺M Φ ^{BM} with cell length measurements (red) by Zeiss Axiovision software. (B) Quantitative analysis of cell length measurements. Mean and individual cell lengths shown ($n = 20-46$ counts, $n = 3-4$ wells/group). Elongated cell processes in M Φ ^{BM} are linked with polarization towards a highly phagocytic, pro-healing phenotype, resulting from amplified OPN expression following GA treatment. OPN inhibition by siRNA produced shorter cells, and under the resulting OPN deficiency, not even GA treatment could restore their elongated morphology. (C) Fluorescent images of M Φ ^{BM} treated with GA or minocycline (M) for 24 h and then exposed to fA β ₄₂ for 30 min. Cells immunostained for inducible nitric oxide synthase (iNOS) and A β (6E10). (D, E) Quantitative ICC revealed that minocycline treatment increased iNOS expression, which in turn reduced fA β ₄₂ phagocytosis. (F–J) Altered phenotype of OPN-knockout M Φ ^{BM} (KO^{OPN}). (F) Representative fluorescent micrographs of control vs. KO^{OPN}-M Φ ^{BM} labeled for A β (6E10), scavenger receptor CD36, and nuclei (DAPI). Reduced fA β ₄₂ uptake in KO^{OPN}-M Φ ^{BM} is associated with significant decrease in surface CD36 expression (G). Quantitative ICC analysis of (H) iNOS, (I) interleukin 10 (IL10), and (J) matrix metalloproteinase 9 (MMP9) in control vs. KO^{OPN}-M Φ ^{BM}. Reduced OPN expression in KO-M Φ ^{BM} altered their immunological profile; addition of GA did not affect MMP9 expression in KO^{OPN}-M Φ ^{BM}. (K) Quantitative ICC reveals increase in MMP-9 following GA treatment in WT M Φ ^{BM}. Means of individual cell fluorescent areas are indicated ($n = 5-10$ images/well, average 100 cells/image, $n = 3-5$ wells/group). (L) OPN levels as determined by Western blot with rabbit anti-OPN antibody. (M) Respective quantitative band density analysis shows elevation of OPN fragments derived from metalloproteinase (MMP) proteolysis following GA treatment, consistent with elevated MMP-9 production (normalized to β -actin input; $n = 4$ wells/group, 1×10^6 cells/well; repeated experiments). Secretion and proteolysis of OPN by M Φ ^{BM} polarized towards an anti-inflammatory phenotype (mediated by MMP) amplified fA β ₄₂ phagocytosis. Group means, SEMs and individual data points are shown. Fold increases in mean values compared with controls indicated in red. Scale bars: 50 μ m. * $p < 0.01$, ** $p < 0.001$, *** $p < 0.0001$, by one-way ANOVA and Tukey's multiple comparisons post tests or two-tail unpaired Student's *t*-test.

OPN form was elevated 12-fold after GA treatment (Fig. 7M; $p < 0.001$). This fragment's basal levels were very low in untreated $M\Phi^{BM}$, but rose in prominence after GA activation (Fig. 7M). The combined data suggest that OPN affects various aspects of macrophages' immunological profile, including polarization towards a pro-healing, anti-inflammatory, highly phagocytic phenotype.

4. Discussion

In this study, we identified OPN's novel role as a major contributor to $A\beta$ clearance through elevated monocyte-macrophage recruitment into ADtg mouse brains and promotion of macrophage polarization towards an anti-inflammatory, highly phagocytic phenotype. Using both in vivo and in vitro AD models, we demonstrated that treatment with the FDA-approved drug GA effectively upregulates OPN expression in macrophages and boosts phagocytosis of $A\beta$. OPN showed an expression pattern specific to brain regions associated with AD (hippocampus and cortex), was predominantly detected in infiltrating monocytes and macrophages, and increased during aging – especially in diseased ADtg mice. Moreover, in brain tissues from AD patients, we also identified increased OPN within myelomonocytic cells surrounding $A\beta$ lesion sites, mirroring our results in ADtg mice. Our flow cytometry analysis of blood and brain myelomonocytic cells in ADtg mice showed a GA-induced enrichment of OPN⁺ monocyte subpopulations in the blood (CD115⁺CD11b⁺Ly6C^{high}) and, most importantly, in the brain (CD11b⁺Ly6C⁺CD45^{high}). Indeed, histological analysis confirmed that OPN was primarily expressed in brain-infiltrating Iba1⁺CD45^{high} or GFP-labeled monocytes and macrophages, but also to a lesser extent in Iba1⁺CD45^{low-mid} microglia. Our in vitro studies substantiated the in vivo findings by demonstrating that macrophage-mediated $A\beta$ clearance is OPN-dependent: upon gain-of-function, it promotes a phenotype that is highly phagocytic of $A\beta$ and anti-inflammatory, while its inhibition impairs uptake of $A\beta$ fibrils and hinders the neuroprotective effects of GA on macrophage immunological profile. Our multiple correlation analyses strongly suggest that OPN is an immunomodulatory cytokine relevant to resisting AD pathology. It shows tight links to increases in brain-infiltrating monocytes and levels of monocyte-chemoattractant MCP-1, to decreases in soluble/insoluble vascular and parenchymal $A\beta$ burden, and to a macrophage phenotype notable for overexpression of MMP-9. This study urges further investigation of OPN as a tool for therapeutic intervention targeting myelomonocytic cells in AD.

Although increased OPN expression has been detected in reactive microglia following traumatic brain injury (Chan et al., 2014) and in ADtg mouse brains (Keren-Shaul et al., 2017; Yin et al., 2017), here we observed that cerebral OPN expression in AD patients and murine models increases predominantly in monocytes and macrophages compared to the activated microglia surrounding $A\beta$ plaques. This observation is consistent with studies demonstrating that blood-borne infiltrating macrophages phagocytose $A\beta$ plaques more efficiently than do resident microglia (Bernstein et al., 2014; Butovsky et al., 2007; Koronyo et al., 2015; Koronyo-Hamaoui et al., 2009; Majumdar et al., 2008). Two recent reports were unable to demonstrate the impact of brain-infiltrating monocyte/macrophages on reducing cerebral $A\beta$ burden; however, these studies involved invasive techniques and/or compounds that depleted brain microglia (Prokop et al., 2015; Varvel et al., 2015), possibly eliminating the local signals to attract monocytes to $A\beta$ lesion sites (Shechter et al., 2009). In contrast, growing evidence from genetic, physiological, and histological studies overwhelmingly support the key role of peripheral monocytes in cerebral $A\beta$ clearance (Bernstein et al., 2014; Butovsky et al., 2007; El Khoury et al., 2007; Koronyo et al.,

2015; Koronyo-Hamaoui et al., 2009; Lai and McLaurin, 2012; Lebson et al., 2010; Malm et al., 2012; Simard et al., 2006). OPN's known interactions with various surface integrins and scavenger receptors [reviewed in (Rittling, 2011)] further support its direct effects on the phagocytic activity of these inflammatory cells. Notably, the marked upregulation of cerebral OPN following GA immunization was even greater when combined with intravenous CD115⁺ monocyte enrichment. These results parallel our previous report of increased monocyte recruitment to the brain (defined by CD115, Ly6C, and CD45 biomarkers) brought on synergistically through the combination of these immunomodulatory approaches (Koronyo et al., 2015). Given that OPN is an integrin-binding protein expressed by macrophages, our findings could explain its action in opsonization and recruitment of immune cells to inflammatory sites (Kahles et al., 2014; Lund et al., 2009; Rittling, 2011). Therefore, we conclude that when GA induces recruitment of peripheral phagocytic cells to amyloid plaques, this immunomodulatory phenomenon is mediated by OPN.

This study indicates that GA substantially induces OPN expression in BM-derived macrophages in a manner similar to that observed in brains of ADtg mice. We also found consistent evidence for increased OPN and subsequent intracellular uptake of pre-formed $A\beta_{40}$ and $A\beta_{42}$ fibrils by these GA-stimulated primary cultures. As expected, the cytokine OPN was mostly localized to macrophages within the major secretory pathway, the trans-Golgi network vesicles (Verhulst et al., 2002), indicating that it is constantly expressed and, perhaps, sorted for post-translation modifications prior to secretion. This pattern of intracellular OPN expression was different from that of the endosomal-lysosomal localization of $A\beta_{40}$ and $A\beta_{42}$ fibrils following phagocytosis. By utilizing systems that negatively regulate OPN expression (siRNA, genetic OPN^{-/-}, or minocycline) or positively induce it with GA or supplementation of rOPN (Suzuki et al., 2010), we have demonstrated OPN's crucial role in macrophage-mediated $A\beta$ phagocytosis. These results, along with literature describing OPN's involvement in other phagocytic activities, reveal that fibrillar $A\beta$ phagocytosis may directly depend on OPN expression (Shin et al., 2011; Toyonaga et al., 2015).

Different functions of OPN may be achieved through its complex proteolytic processing, such as cleavage by MMPs that permits diverse cellular interactions. Specifically, MMP-9 has been implicated in generating functional, more neuroprotective fragments of OPN than the full length OPN (Chan et al., 2014; Scatena et al., 2007). Our data indicate that the dramatic reduction of MMP-9 expression following OPN inhibition cannot be recovered by GA, implying that GA's mechanistic effects are mediated by OPN. In WT macrophages, GA not only induced MMP-9 expression, but also significantly increased the abundance of full-length OPN and its MMP-cleaved fragments. In agreement with previous studies (Ge et al., 2017; Sun et al., 2013), our results support the notion that OPN and MMP-9 have an intricate relationship, regulating each other's expression levels or functions. Our examination of infiltrating Mo/M Φ in ADtg mouse brains further supports the connection between MMP-9 and OPN, evidenced by their tight correlation and increased expression following GA immunization. These findings align with reports from various groups, all of which indicate MMP-9's role in generating functional forms of OPN (Chan et al., 2014; Scatena et al., 2007; Tambuyzer et al., 2012). It is possible that macrophages may control neuroinflammation through release of MMP-9, which in turn cleaves full-length OPN to generate potent immunoregulatory C-terminal OPN fragments. While future studies should ascertain how each OPN fragment may impact macrophage phenotype in the context of AD, we postulate that GA's induction of a therapeutic macrophage phenotype is mediated by elevating these MMP-cleaved neuroprotective fragments of OPN.

Although generally regarded as a pro-inflammatory cytokine, recent studies have shown that OPN may contribute to repair-promoting processes in the brain (Albertsson et al., 2014; Chan et al., 2014; Meller et al., 2005; Shin et al., 2011; van Velthoven et al., 2011). This study unexpectedly demonstrated that GA polarizes macrophages towards a phenotype that is highly phagocytic (increased uptake of fibrillar A β and associated surface scavenger receptors), anti-inflammatory (reduced iNOS and increased IL-10 and MMP-9), and pro-healing (elongated), a process dependent on OPN expression. Upon inhibition of OPN, this phenotype shift was reversed. Since OPN is known to suppress production of NO and iNOS [reviewed in Rittling (2011)], it was no surprise that OPN inhibition resulted in increased iNOS while OPN elevation (via GA) decreased iNOS. Beyond anti-inflammatory IL-10 induction, elevated OPN expression and consequent down-regulation of NO/iNOS may present a mechanism by which macrophages minimize neuroinflammation in AD. Our previous reports of the marked therapeutic effects of GA on cognitive function in ADtg mice (Bakalash et al., 2011; Butovsky et al., 2006, 2007; Koronyo et al., 2015; Zuroff et al., 2017) warrant future studies to elucidate OPN's potential impact on synaptic and cognitive preservation in vivo. Collectively, we have demonstrated a novel role for OPN as a key immune-regulator of macrophage phenotype and A β clearance in AD models. The specific OPN-dependence of macrophage polarization suggests that this cytokine is a potential key neuroprotectant involved in resisting AD-related pathology and promoting tissue repair.

Compliance with ethical standards

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Ethical approval

All applicable international, national, and/or institutional guidelines for the care and use of animals were followed. All procedures performed in studies involving animals were in accordance with the ethical standards of the institution at which the studies were conducted. This article does not contain any studies with human participants performed by any of the authors.

Author contributions

M.K.-H.: Study conception, design and supervision. A.R., M.K.-H.: Major experimental contribution, planning, data acquisition and analysis. Y.K., J.S., D.-T.F., B.C.S.: Other important experimental contribution, image acquisition, flow cytometry, and/or data analysis; S.L.: assistance with siRNA experiments; E.Y.H., D.B.T.: help with fibrillar A β preparations and rOPN experiments; M.K.-H., A.R., Y.K., J.S., D.-T.F., H.E.S., N.J.H., D.D.: Data analysis, interpretation, and presentation; M.K.-H.: Discussion on intellectual content, supervision, and manuscript editing. M.K.H., Y.K., J.S., D.-T.F., N.J.H. and A.R.: Manuscript editing, figure preparation, and revision. M.K.-H., A.R.: Figure preparation, manuscript writing, revision, and final approval.

Dedication

This paper is dedicated to the memory of Dr. Salomon Moni Hamaoui and Lillian Jones Black, both of whom died from Alzheimer's disease.

Conflict of interest

Authors declare no conflict of interest. International patent application PCT/US2017/033875 filed in the United States 5/22/2017. Legal Ref: 065472.000611WO00. *Inventors*: Koronyo M, Rentsendorj A, Black KL, and Koronyo, Y.

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Appendix A. Supplementary data

Supplementary data associated with this article can be found, in the online version, at <http://dx.doi.org/10.1016/j.bbi.2017.08.019>.

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