



Alzheimer's Retinopathy: Seeing Disease in the Eyes

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The neurosensory retina emerges as a prominent site of Alzheimer's disease (AD)

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pathology. As a CNS extension of the brain, the neuro retina is easily accessible for noninvasive, high-resolution imaging. Studies have shown that along with cognitive decline, patients with mild cognitive impairment (MCI) and AD often suffer from visual impairments, abnormal electroretinogram patterns, and circadian rhythm disturbances that can, at least in part, be attributed to retinal damage. Over a decade ago, our group identified the main pathological hallmark of AD, amyloid β -protein (A β) plaques, in the retina of patients including early-stage clinical cases. Subsequent histological, biochemical and in vivo retinal imaging studies in animal models and in humans corroborated these findings and further revealed other signs of AD neuropathology in the retina. Among these signs, hyperphosphorylated tau, neuronal degeneration, retinal thinning, vascular abnormalities and gliosis were documented. Further, linear correlations between the severity of retinal and brain AB concentrations and plaque pathology were described. More recently, extensive retinal pericyte loss along with vascular platelet-derived growth factor receptor- β deficiency were discovered in postmortem retinas of MCI and AD patients. This progressive loss was closely associated with increased retinal vascular amyloidosis and predicted cerebral amyloid angiopathy scores. These studies brought excitement to the field of retinal exploration in AD. Indeed, many questions still remain open, such as queries related to the temporal progression of AD-related pathology in the retina compared to the brain, the relations between retinal and cerebral changes and whether retinal signs can predict cognitive decline. The extent to which AD affects the retina, including the susceptibility of certain topographical regions and cell types, is currently under intense investigation. Advances in retinal amyloid imaging, hyperspectral imaging, optical coherence tomography, and OCT-angiography encourage the use of such modalities to achieve more accurate, patient- and user-friendly, noninvasive detection and monitoring of AD. In this review, we summarize the current status in the field while addressing the many unknowns regarding Alzheimer's retinopathy.

Keywords: neurological disease, neurodegenerative disorder, retina, ocular pathology, retinal imaging, optical imaging, amyloid- β plaques, hyperphosphorylated tau

INTRODUCTION

Alzheimer's disease (AD) is the most common age-related neurodegenerative disorder affecting over 50 million people worldwide (Hickman et al., 2016; Patterson, 2018). With no cure and limited options for early unambiguous and noninvasive diagnosis, this devastating and invariably fatal disease remains a major medical, sociological and economical challenge around the globe (Patterson, 2018; Alzheimer's Association, 2019).

Patients with Alzheimer's dementia typically exhibit symptoms of cognitive decline including disorientation, short-term memory loss, confusion, and socio-behavioral impairments. Beyond psycho-cognitive dysfunctions, these patients often experience visual abnormalities such as diminished color and contrast vision and narrowing of the visual field, as well as disruptions of circadian rhythms manifesting as sleep disturbances (Petersen et al., 1999; Perrin et al., 2009; Wang and Holtzman, 2020). Some of these visual dysfunctions and sleep irregularities have been documented early in the prodromal phase of AD in patients with mild cognitive impairment (MCI).

The neuropathological hallmarks of AD – amyloid β -protein (AB) plaques and neurofibrillary tangles (NFTs) comprised of hyperphosphorylated pTau protein - are well established and characterized in the brains of AD patients (Goedert et al., 1989; Hardy and Selkoe, 2002; Selkoe, 2004, 2008; Perrin et al., 2009; Goedert, 2015; Avila et al., 2016). These hallmark pathologies are hypothesized to induce and amplify inflammation and vascular abnormalities, drive synaptic and neuronal loss, and eventually lead to clinical AD-dementia (Choi et al., 2014; Ontiveros-Torres et al., 2016). The preclinical phase of AD-related pathological buildup is an insidious process which can take up to 20 years (Perrin et al., 2009; Bateman et al., 2012; Dubois et al., 2015; Bilgel et al., 2016; De Strooper and Karran, 2016; Dubois, 2018; Douglas and Scharre, 2019). Intervention during this preclinical stage, before definitive clinical symptoms appear and when synaptic and neuronal damage is still limited, should hold promise for increased therapeutic efficacy.

Notably, a recent research framework by the National Institute on Aging and Alzheimer's Association (NIA-AA) classifies AD as compared with other neurodegenerative diseases based on molecular biomarker detection (Jack et al., 2018). The pathobiological phases of AD in living patients, irrespective of cognitive status, are determined on a continuum beginning with Alzheimer's pathologic change and progressing to full AD by molecular biomarkers: $A\beta$ deposition (A), pathologic tau (T), and neurodegeneration [AT(N)]. Accordingly, [N] is not specific to AD and the presence of (A) is necessary to define Alzheimer's continuum (Jack et al., 2018; Gauthier and Rosa-Neto, 2019). Indeed, together with other genetic, longitudinal brain imaging, physiological, and pathological studies, this framework emphasizes that $A\beta$ and tau pathologies are required to define AD. Moreover, as efforts to develop and evaluate other potential AD biomarkers intensify, including those measuring vascular changes, inflammation and synaptic loss (Chen et al., 2018; Baldacci et al., 2020), the AT(N) system remains flexible to incorporate novel biomarkers upon future availability.

In addition to the pathology described above, examination of postmortem AD brains demonstrated the existence and propagation of intra- and extracellular, soluble and synaptotoxic forms of AB oligomers (Selkoe, 2008; Meli et al., 2014; Li et al., 2020) and pTau assemblies (Takeda et al., 2015; Klein et al., 2019). Other key features of AD neuropathology include cerebral amyloid angiopathy (CAA) (Vinters, 1987) and neuroinflammation - the latter usually involving prolonged activation of microglia and astrocytes, release of pro-inflammatory cytokines and chemokines, as well as infiltration of peripheral immune cells [reviewed in Wyss-Coray (2006)]. In recent decades, cumulative evidence supporting the neuroprotective effects of certain immune cell types and inflammatory mediators have caused a historic shift in the view of neuroinflammation away from the common, merely detrimental one (Zuroff et al., 2017; Deczkowska et al., 2018; Schwartz et al., 2020). In this regard, studies showed that subtypes of bone marrow (BM)-isolated peripheral innate immune cells (e.g., BM-derived CD115+Ly6ChiCD45hi monocytes) can be recruited to the diseased brain of AD-model mice. Interestingly, the recruited cells were shown to directly facilitate Aβ clearance, reduce chronic and detrimental inflammation including scar tissue proteins, and induce synaptogenesis and neurogenesis (Simard et al., 2006; Butovsky et al., 2007; Koronyo-Hamaoui et al., 2009; Lebson et al., 2010; Bernstein et al., 2014; Koronyo et al., 2015; Rentsendorj et al., 2018; Koronyo-Hamaoui et al., 2019; Li et al., 2020). These and other promising multi-targeted immunomodulatory strategies of harnessing peripheral immune cells to fight neurodegeneration are currently being developed and tested in various preclinical and clinical trials across the world (Frenkel et al., 2005; Butovsky et al., 2006; Bakalash et al., 2011; Kunis et al., 2013; Villeda et al., 2014; Koronyo et al., 2015; Baruch et al., 2016; Koronyo-Hamaoui et al., 2019; Rosenzweig et al., 2019; Li et al., 2020). Despite these advances, limitations persist due to the lack of readily available in vivo approaches to noninvasively and accurately monitor therapeutic efficacy, motivating scientists to search for new tools that can be widely deployed in the clinical setting.

Indeed, in recent years there has been tremendous progress in the development of AD diagnostic imaging biomarkers, including FDA-approved brain amyloid imaging via positron emission tomography (PET), as well as tau, surrogate markers of inflammation such as TSPO, and synaptic PET imaging (Rabinovici et al., 2007; Schilling et al., 2016; Chen et al., 2018; Edison et al., 2018; Narayanaswami et al., 2018; Chandra et al., 2019; Werry et al., 2019). In addition, structural, functional and metabolic brain imaging are instrumental in evaluating neuronal damage in patients (Johnson K.A. et al., 2012). Yet, most of these methods are not suitable for repeated population screening in the preclinical stages. They are either limited by the use of unsafe ionizing isotopes (radioactivity), high costs, low availability, and/or limited resolution or specificity (Perrin et al., 2009; Ono and Saji, 2011; Johnson J.K. et al., 2012; Johnson K.A. et al., 2012; Koronyo et al., 2012; Mathis et al., 2012; Doraiswamy et al., 2014; James et al., 2015; Kalia and Lang, 2015; Hart et al., 2016; Heurling et al., 2016; Tiepolt et al., 2016; Doustar et al., 2017; Hampel et al., 2018; Baldacci et al., 2020). As it relates to amyloid PET imaging,

available tracers and modalities do not allow for detection of early small or soluble forms of A β accumulation (Lockhart et al., 2007; Ng et al., 2007; Johnson K.A. et al., 2012).

Milestone advances in the development and evaluation of plasma (A β_{1-42} and neurofilament light concentrations) and CSF biomarkers (A β_{1-42} , total-tau, p-tau concentrations) have also revolutionized the prospect of early diagnosis for AD (Hampel et al., 2018; Baldacci et al., 2020). In general, both categories of fluid biomarkers have proven superior to the majority of currently available brain scanning tools with regards to sensitivity, accessibility and cost-effectiveness. Yet, CSF extraction remains an invasive approach and plasma markers are subject to interference from other bodily organs and peripheral metabolic processes, and therefore may not be able to solely represent the underlying neuropathological events occurring in the brain (Johnson K.A. et al., 2012). These challenges provide great incentive to explore other diagnostic methods for deployment in the general population, perhaps through another CNS tissue - the retina.

Serving as the innermost layer of the eyeball, the main role of the retina in visual perception is the conversion of light signals into decodable neuronal impulses for transmission to the brain. It consists of the light-sensitive neurosensory retina and retinal pigment epithelium (RPE). Structurally, the neural retina contains several distinct layers: the innermost inner limiting membrane (ILM), nerve fiber layer (NFL), ganglion cell layer (GCL), inner plexiform layer (IPL), inner nuclear layer (INL), outer plexiform layer (OPL), outer nuclear layer (ONL), outer limiting membrane (OLM), and the outermost photoreceptor layer (PRL) (see left panel in Figure 1). The five types of neurons for visual perception in the retina are photoreceptors, bipolar cells, ganglion cells, horizontal cells, and amacrine cells. The two types of photoreceptor cells, rods and cones, are the primary light-sensing cells. Rods are responsible for sensing dim light, while cones provide color vision. Amacrine cells and bipolar cells are the two intermediate neurons that pass visual information to the ganglion cells, which then transmit signals to the brain. Similar to the brain, the retina also possesses different types of supporting glial cells: Müller cells, astrocytes, and microglia. Müller cells and astrocytes are the two types of macroglia in the retina and provide support to neurons, while microglial cells act like tissue macrophages. Blood is supplied to the retina via the central retinal artery, which enters the optic nerve and supports the inner neural retina, and via choroid blood vessels, which nourish the RPE and outer neural retina.

The retina is the only CNS tissue not shielded by bone, allowing for noninvasive imaging and providing a unique perspective into the brain. A developmental outgrowth of the embryonic diencephalon, the retina fittingly shares many structural and functional features with the brain including a blood barrier and populations of neurons and glial cells, which secrete proteins related to the amyloid cascade (e.g. BACE1, γ -secretase, ApoE, clusterin) (Morin et al., 1993; Purves and Fitzpatrick, 2001; Johnson et al., 2002; Byerly and Blackshaw, 2009; Maude et al., 2009; Cai et al., 2012; Li et al., 2016; Trost et al., 2016; Vecino et al., 2016). The retina is physically connected to the brain via axons of the optic nerve (Figure 1A), which facilitate vesicular transport of synthesized A β PP, potentially to and from retinal ganglion cells (RGCs) (Morin et al., 1993). Further, brain and retinal microvasculatures are morphologically and physiologically similar (Patton et al., 2005). Overall, the close relationship between these two CNS tissues and their anatomical sub-structures as well as the feasibility of noninvasive retinal imaging may provide a window into better understanding of processes in the CNS such as healthy aging and neurodegeneration.

Mounting evidence demonstrate AD-related retinal pathology in AD patients and animal models. Recent studies documented parallels between the brain and retinal pathology found both in AD patients and animal models (Koronyo-Hamaoui et al., 2011; Koronyo et al., 2012; Shi et al., 2014; Hart et al., 2016; Doustar et al., 2017; Koronyo et al., 2017; Koronyo-Hamaoui et al., 2020). While early examinations of postmortem eyes isolated from AD patients revealed loss of optic nerve integrity and RGC degeneration (Hinton et al., 1986; Blanks et al., 1989; Sadun and Bassi, 1990; Blanks et al., 1996a,b), it was not until 2010 that Koronyo-Hamaoui and colleagues were able to identify the existence of pathological hallmarks, AB deposits, in retinas isolated from these patients (Koronyo-Hamaoui et al., 2011). These findings were also true for MCI and other early-stage AD cases. Subsequent studies confirmed the original results and further identified pTau, characterized retinal plaque subtypes, as well as demonstrated neuronal degeneration and elevated levels of Aβ alloforms, astrogliosis and microgliosis (Alexandrov et al., 2011; Koronyo-Hamaoui et al., 2011; Koronyo et al., 2012, 2017; Schön et al., 2012; Tsai et al., 2014; Hart et al., 2016; La Morgia et al., 2016; den Haan et al., 2018; Hampel et al., 2018; Grimaldi et al., 2019; Schultz et al., 2020; Shi et al., 2020). In this review, we describe key pathological processes that were found in the AD retina (illustrated in Figure 1), with a focus on characteristic Aβ and tau accumulation and emerging retinal amyloid imaging modalities, which provide promise for advancing noninvasive methods for early disease diagnosis and monitoring.

RETINAL Aβ PATHOLOGY IN ALZHEIMER'S PATIENTS

One decade ago, and more than a century following the identification of $A\beta$ plaques in the postmortem brain of the first person diagnosed with AD, Auguste Deter, AD-specific pathological hallmarks were shown for the first time in the human retina (Koronyo-Hamaoui et al., 2011). In this original study, $A\beta$ plaques were identified in all flatmount retinas isolated from 13 cases with definite and probable AD, as confirmed by both brain pathology and clinical reports (**Figures 2A–C'**). Retinal $A\beta$ -plaque pathology in these patients was in stark contrast to minimal to no pathology found in the retina of age- and gender-matched cognitively normal individuals (**Figures 2B–C'**) (Koronyo-Hamaoui et al., 2011; La Morgia et al., 2016; Koronyo et al., 2017).

This and two subsequent studies on human cohorts of over 50 patients and control donor eyes, examining retinal flatmounts and cross-sections with $A\beta$ -specific monoclonal antibodies



FIGURE 1 | Schematic illustration of retinal pathology in AD patients. A comparison between cognitively normal control and AD retinas along a continuum. AD attributed retinal pathology includes accumulation of Aβ and phosphorylated-tau proteins, degeneration and subsequent thinning, and an inflammatory response. AD retinal vessels display substantial thinning and pericyte loss alongside Aβ protein deposition. Modified illustration from Advances in Retinal Imaging: Retinal Amyloid Imaging (Koronyo-Hamaoui et al., 2020) with permission from Springer Nature via Copyright Clearance Center. Aβ, Amyloid-β protein; CTRL, control; GCL, ganglion cell layer; ILM, inner limiting membrane; INL, inner nuclear layer; IPL, inner plexiform layer; mRGCs, melanopsin-containing retinal ganglion cells; NFL, nerve fiber layer; NFT, neurofibrillary tangles; OPL, outer plexiform layer; ONL, outer nuclear layer; OLM, outer limiting membrane; PRL, photoreceptor layer; RGCs, retinal ganglion cells; pTau, hyperphosphorylated tau.



FIGURE 2 | Increased retinal A β_{42} deposition correlates with cerebral amyloid plaque burden in Alzheimer's patients. (A) Representative micrographs from an AD brain and (B) flat-mount retinas from a cognitively normal (CN) subject and (C) AD patient stained with anti-A β_{42} mAb (12F4). Although smaller in size, retinal A β plaques are similar in morphology to brain plaques. Scale bar: 20 μ m. (C) High-magnification images reveal diffuse, compact, and "classical" mature plaque morphology of retinal A β aggregates. Scale bar: 10 μ m. (D) Quantitation of retinal A β_{42} plaque burden, measured by 12F4 immunoreactive area, in AD patients (*n* = 8) and sex-/age-matched CN control subjects (*n* = 7). Data shown as group mean ± SEM. ***P* < 0.01, unpaired 2-tailed Student's *t*-test. (E,F) Pearson's correlation coefficient tests between retinal A β_{42} plaque load (12F4-immunoreactivity) and mean cerebral neuritic plaque burden (e; *r* = 0.87, *P* = 0.0048, *n* = 8; severity score of Gallyas silver staining) or regional plaque burden either in the entorhinal (F; black symbols; *r* = 0.84, *P* = 0.0092, *n* = 8). Reproduced from Koronyo et al. (2017) with permission of ASCI via Copyright Clearance Center.



cross-sections from definite AD patients showing (**A**) ultrastructure of A β plaque (p), fibrils (fib) and protofibrils (pfb) near a blood vessel (bv). Scale bar: 1 μ m. (**A**') High-magnification image showing A β fibrils, protofibrils and A β deposits (abd). Scale bar: 50 nm. (**B**) A β plaque-like deposits (p); demarcated by red line), near basement membrane (bm) of a blood vessel (bv). Scale bar: 0.5 μ m. (**B**) High magnification image of region marked by red asterisk in (**B**) showing dense A β deposition with structural similarity to annular oligomers (red arrowhead). Scale bar: 40 nm. All sections were prestained with anti-A β_{42} mAb (12F4) and peroxidase-based system and DAB substrate chromogen. Reproduced from Koronyo et al. (2017) with permission of ASCI via Copyright Clearance Center.

(12F4, 11A5-B10, 6E10, 4G8), anti-A β dyes (i.e., Curcumin, Thioflavin-S, Congo-Red) and Gallyas silver stain, showed that all neuropathologically confirmed AD patients exhibited A β deposits in the retina (Koronyo-Hamaoui et al., 2011; La Morgia et al., 2016; Koronyo et al., 2017). Interestingly, through scanning of retinal flatmounts, the team discovered a non-uniform manifestation of A β deposits across the human retina. Plaques were more often detected in peripheral regions, especially in the superior and inferior quadrants (Koronyo et al., 2017).

A quantitative histological analysis of whole-mount retinas in a subset of confirmed AD patients compared to age- and sex-matched cognitively normal controls revealed a significant 4.7-fold increase in A β_{42} -containing retinal plaque burden in patients (**Figure 2D**; Koronyo et al., 2017). Another group, using a sample of retinas isolated from 10 neuropathologically and clinically confirmed AD and 10 control patients, demonstrated a significant 2.7-fold increase in the number of retinal A β_{42} plaques that were also found to be larger in volume relative to deposits detected in normal control tissue (Grimaldi et al., 2019). Importantly, although retinal A β plaques are typically smaller in size compared to brain plaques, their burden in the retina significantly correlated with severity of plaque pathology in the brain (**Figures 2E,F**; Koronyo et al., 2017). In particular, retinal amyloid deposits were more strongly correlated with plaque burden in the primary visual cortex and the entorhinal cortex (**Figure 2F**; Koronyo et al., 2017).

Transmission electron microscopy (TEM) analysis of $(12F4^+)A\beta_{42}$ -positive immunoreactivity in retinal tissues from AD patients revealed the ultrastructure of A β in plaques, fibrils, protofibrils and annular oligomer-like forms (**Figures 3A–B**'; Koronyo et al., 2017). Gallyas silver stain further exposed the existence of retinal neuritic-like plaques. While marked increases in retinal A β pathology were noted in AD patients as compared with age-matched cognitively normal individuals, retinal plaques in patients frequently appeared in clusters and preferentially in the mid- and far-peripheral regions (La Morgia et al., 2016; Koronyo et al., 2017). These findings suggest that regional and geometric differences in plaque density should be considered when examining retinal tissue from patients. Moreover, the use of traditional histological techniques in

retinal cross sections of limited regions could account for the few studies unable to consistently detect $A\beta$ in the AD retina (Schön et al., 2012; Ho et al., 2014). This challenge emphasizes the necessity to standardize approaches for analyzing AD-related pathology across diverse topographical regions of the human retina. Indeed, following untraditional histological protocols developed by Koronyo and colleagues, three additional independent groups were able to detect $A\beta$ deposits in the retina of confirmed AD patients (Tsai et al., 2014; den Haan et al., 2018; Grimaldi et al., 2019).

Prior examinations of normal aged eves demonstrated AB immunoreactivity in the sub-retinal pigment epithelium (RPE) (Loffler et al., 1995) as well as toxic AB oligomers in drusen in the macular RPE of aged and AMD patients (Johnson et al., 2002; Anderson et al., 2004; Luibl et al., 2006). As it relates to AD, an early biochemical evaluation of $A\beta_{40}$ and $A\beta_{42}$ alloforms in retinal tissues of patients showed their existence in the human AD retina, albeit without comparing to levels in control retinas nor assessing correlation with respective brain levels (Alexandrov et al., 2011). Importantly, a recent study corroborated these findings of amyloidogenic $A\beta_{40}$ and $A\beta_{42}$ alloforms in the retina of AD patients (Schultz et al., 2020). The study measured $A\beta_{40}$ and $A\beta_{42}$ levels in retinal and hippocampal tissues of human cohorts with neurodegenerative diseases and compared between ApoE £4 carriers and non-carriers (Schultz et al., 2020). Results from this study showed higher levels of retinal and hippocampal $A\beta_{40}$ and $A\beta_{42}$ in individuals with ADrelated pathological changes and ApoE £4 carriers. Further, levels of both alloforms in the retina correlated with their counterparts in the hippocampus, as well as with NFT and A β plaque burden severity (Schultz et al., 2020).

A recent histological study confirmed the presence of retinal plaques in 6 AD patients and 6 healthy controls, including finding similar sized $12F4^+A\beta_{42}$ -containing deposits, and gave additional insight into the spatial distribution and subtypes of AB aggregates in the human retina (den Haan et al., 2018). A β -positive immunoreactivity and deposits were found in various cell layers in postmortem retinas of AD patients, particularly the INL; deposits were found within horizontal, amacrine, and Müller cells (den Haan et al., 2018). Analyses of retinal pathology in AD donors indicated that AB deposits were more abundant in the inner retinal layers, concentrating in the NFL and GCL (Koronyo et al., 2017). A β_{42} was present in both fibrillar and proto-fibrillar forms, confirmed with TEM and Birefringence (apple-green) of Congo redstained retinas under polarized light (Koronyo et al., 2017). More recently, $A\beta_{40}$ was quantified and mapped in a larger cohort of postmortem human AD retinas (n = 47), showing significant increases in both retinal vascular and abluminal $A\beta_{40}$ in AD patients as compared with matched controls (Shi et al., 2020). Retinal $A\beta_{40}$ was especially abundant in the inner retinal layers of the central retina. Increased $A\beta_{40}$ in the retina of AD patients as compared with cognitively normal individuals was further validated by biochemical ELISA analysis (Shi et al., 2020).

Overall, these growing studies confirm the presence of disease-associated $\,A\beta\,$ species in the human retina and highlight the

striking similarities between retinal and cerebral vulnerability to hallmark AD pathologies.

RETINAL TAUOPATHY IN ALZHEIMER'S PATIENTS

Another key characteristic sign of AD neuropathology that strongly reflects neuronal injury and cognitive decline is abnormal tau, specifically hyperphosphorylated tau and its inclusion in NFTs (Iqbal et al., 2009; Buckley et al., 2017; Hanseeuw et al., 2019). The physiological distribution pattern of total non-phosphorylated tau expression in the human retina was first described in 1995 (Loffler et al., 1995). According to this report and subsequent studies, tau is predominantly expressed in the inner retinal layers, most intensely along three distinct bands in the IPL, more diffusely in the OPL and somatodendritically in the INL (Loffler et al., 1995; Leger et al., 2011). Tau is also localized, albeit weakly, in other inner retinal layers such as the GCL and NFL as well as in photoreceptors of the human retina (Loffler et al., 1995; Leger et al., 2011).

Initial post-mortem examinations of late-stage AD retinas did not reveal neurofibrillary inclusions, neuritic plaques or amyloid angiopathy, despite histological observations of GCL degeneration, reduced NFL thickness and optic nerve axonal atrophy (Hinton et al., 1986; Blanks et al., 1989). While a limited number of studies have been unable to histopathologically detect abnormal tau accumulation in retinas of patients (Ho et al., 2014; Williams et al., 2017), the first evidence of diseaseassociated tau hyperphosphorylation in post-mortem retinas of confirmed AD cases was reported by Schön and colleagues in 2012 (Figures 4A,B; Schön et al., 2012). Results from this study were corroborated thereafter by other groups (den Haan et al., 2018; Grimaldi et al., 2019). Different pTau species, recognized by phosphorylation site-specific antibodies such as AT8 (pSer202, pThr205), AT100 (pThr212 and pSer214), and AT270 (pThr181), were primarily found in the inner retinal layers, particularly the plexiform layers, INL, and GCL of AD patients, thus closely mirroring the physiological expression pattern of normal tau (Schön et al., 2012; den Haan et al., 2018). Interestingly, a recent independent report on a quantitative histomorphometric analysis of post-mortem tissue revealed that these particular retinal layers undergo significant pathological atrophy in AD compared to non-demented control cases (Asanad et al., 2019b). Geometric analysis of postmortem retinal tissue from 6 control cases and 6 AD patients showed more intense AT8-immunoreactivity in superior than in medial retinal regions (den Haan et al., 2018). Qualitative observations from the same study also showed a positive gradient away from the optic nerve and toward the periphery. Despite the presence of pTau in all 6 AD retinas, no significant difference was found in retinal pTau area coverage between the two diagnostic groups, likely due to 2 outliers in control cases. Importantly, a novel and significant association was also found between retinal AT8 burden and cerebral amyloid plaque but not NFT severity (den Haan et al., 2018).

Hyperphosphorylation of tau has been shown to drive the formation of fibrillar tau inclusions and neurofibrillary tangles,



leading to disruptions in axonal transport as well as metabolic and oxidative stress, and is closely associated with neuronal death (Iqbal et al., 2009). Although several histological staining approaches have been utilized to confirm the presence of fibrillar inclusions of pTau in human AD retinas, thus far many have been unsuccessful (Schön et al., 2012; den Haan et al., 2018). To date, only one study using Gallyas silver staining detected NFT-like structures in postmortem retinas of definitive AD patients (Figures 4C,D; Koronyo et al., 2017). As mentioned above, there is an apparent gap in results obtained from different groups, possibly driven by the lack of standardized experimental procedures. For instance, the same antibodies against pTau reported to produce negative results by one group (Schön et al., 2012) detected pTau staining patterns similar to AT8immunoreactivity in a later study (den Haan et al., 2018). Still, by utilizing AT100 antibody, two independent groups detected similar patterns of pTau in postmortem retinas from AD patients (Figure 4E; den Haan et al., 2018; Grimaldi et al., 2019). Future investigations would therefore be necessary to confirm the presence or absence of disease-associated tau conformers in the retinas of AD patients.

Moreover, it is still unclear whether abnormal hyperphosphorylation of tau and/or formation of intracellular tangles have similar detrimental consequences in the retinas of patients as those observed in the brain. Analyses of retinal changes in AD transgenic mice support this putative association and will be discussed in the following sections. According to data from clinical studies and meta-analyses, levels of total tau and pTau in the CSF remain among the most reliable and sensitive biomarkers for both AD diagnosis and longitudinal monitoring of disease progression (Ibach et al., 2006; Welge et al., 2009; AlzBiomarker Database, 2018). Several lines of evidence indicate that CSF tau levels are significantly increased in AD patients (AlzBiomarker Database, 2018). Intriguingly, CSF total tau and

pTau-181 concentrations were shown to correlate with retinal changes measured by fluorescent lifetime ophthalmoscopy, a technique suggested to detect the metabolic alterations of tissue represented by fluorescence decay of endogenous fluorophores (Jentsch et al., 2015). Nevertheless, the co-localization of pTau (den Haan et al., 2018; Grimaldi et al., 2019) and sites of neuronal loss in the retina (Koronyo et al., 2017; Asanad et al., 2019b; Grimaldi et al., 2019) are indicative of similar physiological vulnerabilities to pTau accumulation in both the retina and brain.

Retinal disorders such as glaucoma and age-related macular degeneration (AMD) share a number of common features with AD retinal pathology including progressive deposition of protein aggregates, reactive gliosis and pro-inflammatory responses, metabolic dysfunction, oxidative stress, and retinal atrophy (Quigley, 1999; Naskar et al., 2002; Gupta et al., 2006; Guo et al., 2007; Tezel, 2011). In 2008, examination of retinal tauopathy in surgically removed tissue from human glaucoma cases revealed intense localization of AT8-positive pTau in horizontal cells residing in the OPL (Ibach et al., 2006; Gupta et al., 2008). It is hypothesized that the lateral arrangement of horizontal cell processes may predispose them to retinal stretch injury caused by glaucoma-related elevated intraocular pressure (IOP). Although no association was found between high IOP and dementia (Cesareo et al., 2015), it remains to be seen whether similar changes occur due to AD-associated ocular abnormalities. However, increased retinal pTau accumulation and atrophy following injury or due to other neurodegenerative disorders reveal the vulnerability of this neural tissue compartment to structural, functional, and neuropathological abnormalities (Green et al., 2010; Leger et al., 2011; Kim et al., 2017; Satue et al., 2017; Behbehani et al., 2018; Kim et al., 2019).

Aging remains the principal risk factor in AD and is tightly associated with several visual impairments. To date, only two

studies have investigated the relationship between aging and total tau expression in the human retina. An early semi-quantitative analysis found no difference between tau immunoreactivity in post-mortem retinas from young and old healthy subjects, although age ranges were not clearly outlined (Loffler et al., 1995). Another study, examining retinas from enucleated eyes of patients with prior history of ocular disorders, reported a positive correlation between aging and total tau levels in RGCs in a subset of patients, while unable to find evidence of pTau (Leger et al., 2011). Such reports further showcase the disparity in these findings and reiterate the need for replication of these studies. Notably, the effects of aging on the abnormal accumulation of pTau in retinas of healthy individuals and/or patients also remains unexplored.

As it relates to tau imaging in the retina, there are currently no live imaging tools to specifically detect tau aggregates in the human retina. Preliminary results from a study that utilized spectral domain optical coherence tomography (SD-OCT) and fundus autofluorescein (FAF) to visualize pathological tau aggregates in a cohort of PET-confirmed Alzheimer's patients (Kayabasi, 2018) hints at the possibility of noninvasive live imaging and monitoring of neuropathological changes in the retina of MCI and AD patients; however, the specificity of the signal was not clear.

Overall, it is apparent that investigations of retinal $A\beta$ pathology and its relationship with cerebral amyloid plaque burden in AD patients are mounting. However, our understanding of retinal pTau accumulation and associations with brain NFT severity is much more limited. To date, only a few groups have successfully detected tau hyperphosphorylation in post-mortem human retinas from AD patients, and only one study has shown an association between retinal pTau and cerebral amyloid load in a small number of AD patients. Therefore, there is an urgent need for systematic and quantitative analyses of retinal pTau in larger cohorts as well as assessment of both the spatiotemporal and pathomechanistic properties of AD-related tau species in the retina and their relationship with brain disease and cognition.

ALZHEIMER'S DISEASE HALLMARKS IN THE RETINA OF ANIMAL MODELS

In agreement with the above findings in patients, the pathological hallmarks of AD were also described in numerous animal models of AD (see a summary in **Table 1**). Both the soluble and insoluble forms of A β were found in the retina of sporadic models and transgenic mice harboring familial AD (FAD) mutations (Ning et al., 2008; Dutescu et al., 2009; Liu et al., 2009; Perez et al., 2009; Koronyo-Hamaoui et al., 2011; Koronyo et al., 2012; Tsai et al., 2014; Du et al., 2015; Gupta et al., 2016; Hart et al., 2016; Habiba et al., 2020). Intriguingly, early manifestations of retinal A β plaques have also been detected prior to their occurrence in the brain (Koronyo-Hamaoui et al., 2011). Moreover, upon assessment of therapeutic response, researchers found that positive effects of immunotherapy on cerebral A β -plaque reduction were also reflected in the respective retinas

in transgenic animal models of AD (ADtg) (Liu et al., 2009; Koronyo-Hamaoui et al., 2011; Koronyo et al., 2012; Yang et al., 2013; He et al., 2014; Gao et al., 2015; Parthasarathy et al., 2015). These studies illustrate the common retino-cerebral mechanisms of neuroprotection in response to therapy, which encourages the use of retinal imaging to noninvasively assess therapeutic efficacy in real time.

ADtg animals typically express the transmembrane amyloid precursor protein (APP), the source of A β protein, in retinal neurons. Indeed, this protein has been found in the retinas of ADtg drosophila, various ADtg mice (Tg2576, hTgAPP^{tg/tg}, APP_{SWE}/PS1_{ΔE9}, and APP_{SWE}/PS1_{M146L/L286V}), and the naturally occurring sporadic rodent strain *Octodon degus* (*O. degus*) (Ning et al., 2008; Dutescu et al., 2009; Liu et al., 2009; Ardiles et al., 2012; Du et al., 2015). Retinal cytoplasmic APP was found to increase in ADtg models (Ning et al., 2008; Dutescu et al., 2008; Dutescu et al., 2009), but decrease in the sporadic *O. degus* with aging (Du et al., 2015). Ning and colleagues found APP immunoreactivity increased with age in cells of the INL and GCL – but not the ONL – as well as the neuropil of the IPL and OPL and the outer segments (OS) and RPE (Ning et al., 2008).

ADtg rodent models, including Tg2576, APP/PS1, 3xTg, 5xFAD mice, TgF344-AD rat, and O. degus, show cerebral accumulation of soluble and insoluble AB with age, corresponding to AD-like progression (Dutescu et al., 2009; Liu et al., 2009; Perez et al., 2009; Alexandrov et al., 2011; Koronyo-Hamaoui et al., 2011; Williams et al., 2013; Edwards et al., 2014; Park et al., 2014; Tsai et al., 2014; Du et al., 2015; More and Vince, 2015; Parthasarathy et al., 2015; Pogue et al., 2015). Alloforms of A β pathognomonic to AD (A β_{40} and particularly $A\beta_{42}$) were found to be elevated in the retinas of ADtg rodents (Dutescu et al., 2009; Liu et al., 2009; Alexandrov et al., 2011; Williams et al., 2013; Park et al., 2014; Tsai et al., 2014; Du et al., 2015; Parthasarathy et al., 2015) and ADtg drosophila (Greeve et al., 2004; Carmine-Simmen et al., 2009). In addition, plaque and insoluble AB deposits were identified in retinas of Tg2576, APP_{SWE}/PS1 Δ E9, APP_{SWE}/PS1_{M146L/L286V}, 3xTg, and 5xFAD mice, and in O. degus (Ning et al., 2008; Dutescu et al., 2009; Liu et al., 2009; Perez et al., 2009; Alexandrov et al., 2011; Koronyo-Hamaoui et al., 2011; Koronyo et al., 2012; Williams et al., 2013; Yang et al., 2013; Zhao et al., 2013; Edwards et al., 2014; Du et al., 2015; More and Vince, 2015; Hadoux et al., 2019).

Tg2576 mouse retinas were cross-sectioned and analyzed for plaque pathology, which was found in approximately 85% of these transgenic mice but was absent in WT controls (Liu et al., 2009; Williams et al., 2013). Plaques were mostly found in the GCL, INL, and ONL (Liu et al., 2009; Williams et al., 2013). Yet, one study was unable to detect retinal A β pathology in a Tg2576 mouse with cerebral A β (Dutescu et al., 2009).

5xFAD mice present an aggressive model of amyloidosis, with several familial AD mutations that result in the overexpression of A β_{42} . Studies in this model have demonstrated the presence of A β_{42} in ocular tissues including the retina as well as increases in A β_{40} in the RPE (Park et al., 2014; Parthasarathy et al., 2015; Hadoux et al., 2019). In the sporadic *O. degus* model of AD, A β deposition appears to be progressive, accumulating first in the GCL, NFL, INL and photoreceptors (Chang et al., 2020). Whole

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Hallmarks	Animal species	Models and Genotypes	Retinal pathologies	References
APP or Aβ related	Drosophila	AβPP, dBACE-AβPPL, pGMR-Aβ42	Increased APP, A β , A β_{42} toxicity	Finelli et al., 2004; Greeve et al., 2004; Carmine-Simmen et al., 2009; Cutler et al., 2015
	Mouse	Tg2576, hTgAPP ^{tg/tg} , APP _{SWE} /PS1 _{AE9} , APP _{SWE} /PS1 _{M146L/L286} v, 3xTg, 5xFAD, Tg-SwDI	Increased APP, Aβ deposits, Aβ ₄₂ , Aβ ₄₀ , vascular Aβ, Aβ oligomers	Ning et al., 2008; Dutescu et al., 2009; Liu et al., 2009; Perez et al., 2009; Alexandrov et al., 2011; Koronyo-Hamaoui et al., 2011; Koronyo et al., 2012; Williams et al., 2013; Yang et al., 2013; Zhao et al., 2013; Edwards et al., 2014; He et al., 2014; Park et al., 2014; Gao et al., 2015; More and Vince, 2015; Parthasarathy et al., 2015; Pogue et al., 2015; Gupta et al., 2016; Oliveira-Souza et al., 2017; Criscuolo et al., 2018; Hadoux et al., 2019; Habiba et al., 2020; Sidiqi et al., 2020
	O. degus	Spontaneous	Increased APP, Aβ, Aβ oligomers	Inestrosa et al., 2005; Ardiles et al., 2012; Du et al., 2015
	Rat	TgF344-AD	Increased A _β	Tsai et al., 2014
NFT or pTau related	Drosophila	hTau	Accumulation of pTau	Grammenoudi et al., 2006; Chouhan et al., 2016
	Mouse	Tg2576, APP _{SWE} /PS1 _{ΔE9} , APP _{SWE} /PS1 _{M146L/L286V} , 3xTg, P301S, rTg4510	Accumulation of pTau, NFT	Liu et al., 2009; Schön et al., 2012; Yang et al., 2013; Zhao et al., 2013; Chiasseu et al., 2017; Grimaldi et al., 2018; Harrison et al., 2019
	O. degus	Spontaneous	Accumulation of pTau	Du et al., 2015; Chang et al., 2020

APP, amyloid precursor protein; Aβ, amyloid-beta; pTau, phosphorylated Tau; NFT, neurofibrillary tangle; O. degus, Octodon Degus.

retinal histological examination via A β -specific staining revealed the most plaque burden in the central retina (Inestrosa et al., 2005; Du et al., 2015).

Importantly, Koronyo-Hamaoui et al. (2011) developed the first approach to visualize retinal AB deposits in live $APP_{SWE}/PS1_{AE9}$ mice using curcumin as a specific Aβ-labeling fluorophore and optical Micron rodent retinal imager (Phoenix Technology Group, LLC). The identity of AB deposits detected in vivo by curcumin was validated by subsequent ex vivo labeling of the respective whole-mount retinas using anti-Aß monoclonal antibody. Notably, retinal plaques were observed prior to detection in the corresponding brains (Koronyo-Hamaoui et al., 2011). In addition, following intravenous injection of curcumin, flatmount retinas from 2.5-month-old mice were isolated and stained with 4G8 monoclonal antibody, further confirming that the curcumin spots are the same Aβ-specific deposits (Koronyo-Hamaoui et al., 2011). A recent study by Sidiqi and colleagues corroborated the previous reports (Koronyo-Hamaoui et al., 2011; Koronyo et al., 2012, 2017) and demonstrated via retinal curcumin imaging in $APP_{SWE}/PS1_{AE9}$ mice the accumulation of retinal A β plaques with disease progression, which positively correlated with $A\beta$ load in the brain (Sidiqi et al., 2020). Crosssectional analysis of adult APP/PS1 ADtg mice retinas revealed A β deposits in the innermost layers as well as in the choroid and surrounding scleral tissue; WT controls showed only minimal to no plaque deposition (Koronyo-Hamaoui et al., 2011). Studies using the same ADtg model and Tg344F-AD rats (with the same double transgenes) also validated these findings (Ning et al., 2008; Liu et al., 2009; Perez et al., 2009; Tsai et al., 2014).

The initial discovery of tau in the adult mammalian retina dates back to 1988. In this early study, tau was predominantly found in horizontal cells residing in the OPL of adult rat retinas (Tucker and Matus, 1988). On the other hand, diseaseassociated pTau species were detected somatodendritically in RGCs of a commonly studied double transgenic mouse model of AD, known as the APP_{SWE}/PS1_{AE9} mouse (Yang et al., 2013), and in the GCL to ONL of Tg2576 ADtg mice (Liu et al., 2009). In line with these observations, intracellular aggregates of pTau have also been detected in retinas of the APP_{SWE}/PS1_{M146L/L286V} mouse model of AD (Zhao et al., 2013). Further, recent analyses of young, pre-symptomatic 3xTg mouse retinas showed retinal A β plaques and tau tangles in the RGC layer (Grimaldi et al., 2018).

In the South American rodent *O. degus*, previously reported to develop several spontaneous AD-like pathologies without genetic manipulation, elevated levels of pTau were primarily detected in the GCL to NFL regions of the retina in both adult and aged animals (Du et al., 2015). In a subsequent study by the same group, early punctate AT8 immunoreactivity in the IPL was reported in young *degus*, compared with the denser expression in IPL to NFL of juvenile and adult animals (Chang et al., 2020). Interestingly, p(tau)-positive aggregates both appeared and propagated to other inner retinal layers earlier than A β deposits in these animals (Chang et al., 2020). Similarly, retinal accumulation of total tau and epitope-specific hyperphosphorylation were also reported to precede onset of behavioral deficits and brain tauopathy as early as 3 months of age in the 3xTg mouse model of AD (Chiasseu et al., 2017).

Several live imaging modalities have been developed and optimized for detection of proteinaceous aggregates and metabolic hyperspectral mapping of retinal tissue in rodent models of AD (Koronyo-Hamaoui et al., 2011; Koronyo et al., 2012, 2017; Schön et al., 2012; More and Vince, 2015; Sidiqi et al., 2020). Using longitudinal scanning laser ophthalmology, Schön et al. visualized and monitored pTau-containing RGCs in the P301S mouse model of tauopathy from 2 to 6.5 months of age and found a steady growth in pTau-positive cell counts (Schön et al., 2012).

In the rTg(tauP301L)4510 tauopathy mouse model of frontotemporal dementia (FTD), accumulation of both tau and

pTau have been observed in RGCs as well as the IPL and INL (Harrison et al., 2019). These areas are associated with reduced neuronal density and optic nerve degeneration in these mice (Harrison et al., 2019). In an experimental rat model of optic nerve crush, injury-induced impaired autophagy was followed by an increase in hyperphosphorylated tau (pSer396), which colocalized with apoptotic markers in dying RGCs (Oku et al., 2019). Silencing the tau gene exerted neuroprotective effects in this model, indicating that tauopathy following retinal injury similar to that observed in the brain plays an essential role in neuronal atrophy (Oku et al., 2019).

AD-related tauopathies have also been investigated in a limited number of non-murine animal models of aging or AD. Selective expression of normal human tau in adult drosophila retina leads to progressive loss of ERG responses without a considerable effect on retinal structure or neuronal density, although TEM analysis later revealed signs of tauinduced retinal synaptotoxicity and abnormal photoreceptor morphology (Chouhan et al., 2016). In the same study, human Aß expression in a separate group of flies caused an agedependent loss of retinal neurons without altering ERG signals (Chouhan et al., 2016). In the UAS-Gal4 drosophila model of AD, species of pTau, phosphorylated at different epitopes to varying degrees and in a cell type-specific manner, have also been detected in both the retina and brain (Grammenoudi et al., 2006). In a small number of young and aged primates, total tau expression in the outer retina was observed in the OPL, ONL and inner segment of photoreceptors, whereas AT8-positive pTau was localized predominantly in the OPL and cones (Aboelnour et al., 2017). In older primates, pTau staining in retinal cones also appeared stronger compared with younger animals (Aboelnour et al., 2017). Altogether, it is apparent that tau expression and pTau accumulation may vary across several species commonly used to model human AD, and future studies should aim to elucidate these differences.

Together with these positive findings, it is important to note that the expression pattern of both APP and tau transgenes in the brain and retina of animal models of AD should not be misinterpreted as a precise reflection of the human disease. The layer and cellular burden of pathological AB and pTau aggregates is most likely governed by the promoter-driven expression of their corresponding transgenes. For instance, expression of the aggregate-prone mutant tau species in the P301S tauopathy mouse model is driven by the murine Thy1 promoter, which leads to the development of pathology in selected CNS cell types and consequently specific regions (Allen et al., 2002). In the retina, Thy1 is uniquely expressed by RGCs (Schmid et al., 1995) and therefore the intracellular aggregation of hyperphosphorylated tau in these cells and their nerve fibers is expected. This may underlie the common disparity in findings from stereological studies attempting to characterize regional and cellular susceptibility to AD pathology in CNS tissue from human versus animal models. Recent developments as well as ongoing efforts to fully characterize complete gene replacement animal models will be invaluable in addressing such limitations in several fields.

RETINAL DEGENERATION IN ALZHEIMER'S DISEASE

In agreement with the observed distribution of retinal AB deposits in AD patients, reports have indicated NFL thinning and RGC degeneration in the GCL chiefly within the superior quadrants of the retina (Blanks et al., 1989, 1996b; Berisha et al., 2007; Lu et al., 2010; Liu et al., 2015; La Morgia et al., 2016; Asanad et al., 2019b; Grimaldi et al., 2019). In vivo OCT imaging of these patient retinas revealed degeneration in multiple retinal layers (La Morgia et al., 2017). Reduced macular thickness and volume, measured by OCT, was also found to correlate with cognitive impairment in AD patients (Iseri et al., 2006). The first study that evaluated melanopsin-containing retinal ganglion cells (mRGCs), photoreceptors known to drive circadian photoentrainment, in the postmortem retina of AD patients described a significant mRGC degeneration reflected in reduction of both dendritic density and cell number (La Morgia et al., 2017). Notably, dendrite loss and cell death were closely linked to A β pathology and were colocalized at sites of A β deposits (La Morgia et al., 2016).

These findings were corroborated in another study, which indicated a substantial loss of retinal cells at sites of Aβ deposition (Figures 5A-D). When compared with age-matched cognitively normal controls, 22-29% retinal neuronal degeneration was detected by Nissl staining (n = 17 subjects) in the GCL, INL, and ONL of AD patients (Figures 5B,C; Koronyo et al., 2017). More recently, Asanad and colleagues analyzed retinal atrophy morphometrically in the superior quadrants of postmortem retinas from 8 AD and 11 age-matched controls (Asanad et al., 2019b). Measurements were acquired along a distance of 4mm from the optic nerve on the supero-temporal (reaching the macular region) and supero-nasal sides. Significant retinal thinning was revealed in the NFL (~40%), GCL (35%), IPL (\sim 20%), and both nuclear layers (25%) in AD patients. In the supero-temporal region, NFL thinning was more pronounced closer to the optic nerve, whereas the other retinal layers showed prominent thinning closer to the macula. However, all analyzed retinal layers showed consistent thinning throughout the superonasal retinal quadrant (Asanad et al., 2019b). Beyond these changes, several studies have demonstrated accumulation of AB in the lens of AD patients (Goldstein et al., 2003; Tian et al., 2014) as well as reductions in the choroid coat - the vascular layer of the eye - and offered insights into the retinal choroid as an oculovascular biomarker for Alzheimer's disease (Tsai et al., 2014; Asanad et al., 2019a).

These findings are further supported by animal models of AD (Frederikse and Ren, 2002; Yang et al., 2013; Park et al., 2014; Tsai et al., 2014). In the hTgAPP^{tg/tg} mouse lens, markers of degeneration included nuclear disorganization, organelle loss, cellular swelling, and shape irregularity (Frederikse and Ren, 2002). In the 5xFAD mouse, A β deposits in the retinal pigment epithelium, hypopigmentation, large vacuoles, and Bruch membrane thickening with Drusen-like deposits were observed (Park et al., 2014). Hypertrophy along with choroid changes were also detected in the TgF344-AD rat (Tsai et al., 2014). Further, in the APP_{SWE}/PS1_{Δ E9} mouse, RGC density was



increased retinal Aβ deposits (red shapes) and neuronal loss (green triangles) in AD patients. Panels a-c are reproduced from Koronyo et al. (2017) with permission of ASCI via Copyright Clearance Center. Panel d is adapted from Hart et al. (2016) under terms of the Creative Commons Attribution 4.0 International License (http://creativecommons.org/licenses/by/4.0/).

lower than in WT controls (Gao et al., 2015; Gupta et al., 2016) and amacrine cell apoptosis was noted (Gao et al., 2015).

A recent finding in 3xTg mice indicated colocalization of both retinal tau tangles and A β plaques with neurodegeneration in the GCL (Grimaldi et al., 2018). In the same mouse model, tau accumulation was associated with retinal neural dysfunction, as measured by deficits in anterograde axonal transport (Chiasseu et al., 2017).

Altogether, observations of retinal atrophy in layers recognized as sites of proteinaceous deposition in both AD patients and animal models of the disease illustrate the vulnerability of retinal cell types to pathological processes traditionally associated with the cerebral disease. In addition, these findings further support the suitability of retinal thinning, in combination with more specific markers, as a diagnostic and monitoring tool for AD.

RETINAL VASCULAR PATHOLOGY IN ALZHEIMER'S DISEASE

Cerebral amyloid angiopathy is defined as a cerebrovascular disease characterized by intense deposition of A β in the walls of cerebral arteries, arterioles, and capillaries, among other vascular damage, and is commonly found in the brains of AD patients

(DeSimone et al., 2017). A β in CAA primarily consists of A β_{40} (Roher et al., 1993; Gravina et al., 1995). In macrovasculature, CAA is composed of AB deposition in tunica media and adventitia of leptomeningeal and cerebral parenchymal arteries (Tian et al., 2004). The AB in CAA will eventually affect all vascular layers and result in degeneration of smooth muscle cells (Keable et al., 2016). CAA is prevalent in the elderly who have developed lobar cerebral hemorrhage (ICH) (Keable et al., 2016). Although CAA remains a distinct clinical entity from dementia, previous studies have demonstrated that over 85% of AD patients have of CAA of varying severity (Arvanitakis et al., 2011; Viswanathan and Greenberg, 2011). The Aβ deposited in the vascular walls triggers several ischemia-induced pathogenic molecular pathways, such as oxidative stress, inflammation and increased blood-brain barrier (BBB) permeability, leading to further hemorrhagic complications (Ghiso et al., 2010). A study based on parametric analysis of neuropathological data from the National Alzheimer's Coordinating Centers' dataset suggested that CAA was facilitating early stage dementia and the transition to moderate dementia (Vidoni et al., 2016). In severe CAA cases, Aβ deposition is usually followed by microaneurysms in cerebral blood vessels (Vonsattel et al., 1991), a vascular pathology shared by retinal vascular diseases such as diabetic retinopathy, which can be easily examined by fundoscopy (Friberg et al., 1987; Hellstedt et al., 1996).

As the retina is a developmental outgrowth of the diencephalon (Purves, 2001; Erskine and Herrera, 2014), it shares many vascular morphological and physiological features with cerebral vessels (Patton et al., 2005; Crair and Mason, 2016). For instance, the retina has a highly selective blood retinal barrier (BRB), with many similar structural and functional properties as the BBB, which modulates the influx of ions, proteins and water, as well as limits the infiltration of circulating immune cells (Cunha-Vaz et al., 2011). The recent identification of an ocular glymphatic drainage system in rodents, clearing fluids and metabolites such as $A\beta$ from the retina and vitreous via an aquaporin-4 (AQP4)-dependent pathway, is a sign of yet another structural and physiological similarity shared by the brain and retina (Wang et al., 2020). The same group also demonstrated that this clearance route may be impaired in ocular conditions associated with retinal damage such as glaucoma (Wang et al., 2020). Whether similar glymphatic drainage occurs in the human eye and the extent to which disruptions in this process contribute to pathological changes in neurological diseases such as AD remain to be investigated.

Numerous reports have broadly described vascular dysfunctions in the AD retina, including increased tortuosity, narrowed veins, decreased blood flow, microvascular network damage, and compromised branching complexity (Berisha et al., 2007; Frost et al., 2010, 2013; Cheung et al., 2014; Feke et al., 2015; Williams et al., 2015; Einarsdottir et al., 2016; Abbasi, 2017; Cabrera DeBuc et al., 2018; O'Bryhim et al., 2018), similar to earlier findings in the brain (Smith et al., 1999; Bookheimer et al., 2000; Kimbrough et al., 2015). Such studies have extensively focused on live imaging of retinal blood flow dynamics of AD patients, yet there is a significant lack of understanding regarding the precise molecular and cellular mediators involved in retinal vascular AD pathology, which could lead to the discovery of potential intervention points of treatments and guide the development of next-generation retinal imaging. Thus, more in-depth investigations of the pathogenic mechanisms of retinal vasculature in AD development are needed.

To this end, work by Koronyo-Hamaoui and colleagues demonstrated the existence of different alloforms of AB deposits in retinal vessels of MCI and AD patients in both perivascular and within vessel walls including within the tunica media and outside the basement membrane (Koronyo et al., 2017). These findings were consistent with what is known regarding cerebral vascular Aβ pathology (Vinters and Gilbert, 1983; Vinters, 1987; Bennett, 2001; Bakker et al., 2016; Engelhardt et al., 2016). TEM analysis confirmed the ultrastructures of retinal AB deposits, often in close proximity to or within blood vessels, similar to those found in the brain (Iadanza et al., 2016; Koronyo et al., 2017; Shi et al., 2020). Retinal $A\beta_{42}$ fibrils near a blood basal membrane were approximately 10-15nm in width with typical anti-parallel β -sheets (Koronyo et al., 2017). In the same human study, the existence of retinal A β was verified by Congo red staining under polarized light, by which the investigators revealed positive AB fibrils along blood vessels (Koronyo et al., 2017). According to an animal study with APP-overexpressing mice, vascular amyloidosis resulted in increased rigidity of the blood vessels and decreased blood flow in the brain (Kimbrough et al.,

2015). In addition, a recent clinical study has proposed that cerebral vascular changes may propel other abnormalities in AD pathogenesis (Besser et al., 2016).

Recent reports suggest that vascular pathology in AD brains occurs very early during disease progression (Nikolakopoulou et al., 2017). Yet, it is unclear how early vascular pathology can be detected via retinal imaging and whether it may specifically foretell AD development. Furthermore, the connections between vascular pathology and A β accumulation and clearance in both the brain and retina should be explored in future studies. To date, several studies imply the potential of utilizing retinal vascular biomarkers for early AD or pre-symptomatic stage screening as well as for predicting cognitive decline (Vidal and Mavet, 1989; Gharbiya et al., 2014; Bulut et al., 2016; Cunha et al., 2017; McGrory et al., 2017; Bulut et al., 2018; Jiang et al., 2018; O'Bryhim et al., 2018; Jung et al., 2019).

A recent investigation by Shi and colleagues of retinal vasculature in 62 AD and MCI patients and matched human controls revealed early and progressive PDGFR^β deficiency and pericyte loss along with intense retinal vascular AB accumulation in AD (Shi et al., 2020). Using a modified retinal vascular isolation technique (Figure 6A) in 12 AD patient and control donors, the authors report a significant increase in various types of A β in AD retinal microvasculature, and more importantly, accumulation of A β in retinal pericytes together with pericyte loss (Figure 6B– E'). The existence of $A\beta$ in pericytes was validated by TEM, which showed intense AB42 deposition in retinal pericytes as well as in microvascular lumen and adjacent to microvasculature (Figure 6F). Further, accumulation of Aβ in arterial tunica media in the retina of several AD patients (Figures 6E,E'; Shi et al., 2020) has implications related to lymphatic AB clearance pathways in the human retina: a subject warranting future exploration.

In the same study, histological analysis based on immunofluorescent staining or 3,3'-Diaminobenzidine staining and the use of specific monoclonal antibodies recognizing diverse A β epitopes revealed intense deposition of both A β_{42} and A β_{40} in AD retinal vasculature (n = 28-36 human samples). In addition, early loss of pericyte marker PDGFR β was noted in longitudinal and vertical retinal vessels of MCI and AD patients compared to normal controls (n = 38). The dramatic loss of vascular PDGFR β expression significantly correlated with CAA (n = 14) and Mini-Mental State Examination (MMSE) cognitive scores (n = 10) in a subset of patients, suggesting that retinal pericyte and PDGFR β loss could predict the cerebral vascular disease and cognitive function. Further TUNEL and cleaved caspase-3 staining demonstrated that apoptosis may be the dominant pathway of retinal pericyte death in MCI and AD retina.

In the brain, the BBB plays an indispensable role in mediating clearance of $A\beta$ through its efflux in the cerebral vascular network (Zlokovic et al., 1993; DeMattos et al., 2002; Banks et al., 2003; Do et al., 2015; Zhao et al., 2015; Sweeney et al., 2018). The BBB is established by endothelial cells forming vessel walls, astrocyte end-feet, and pericytes in the basement membrane. In comparison, the BRB is composed of tight junctions of retinal endothelial and epithelial cells together with supporting pericytes. Despite their organ-specific functions, the BBB and BRB display very similar functions in transport and permeation



characteristics (Steuer et al., 2005). A recent study has established a correlation among BBB, ApoE ɛ4 and cognitive decline regardless of AD pathology (Montagne et al., 2020). The recent findings implicated pericytes in AD pathogenesis and future investigations may shed light on the role of the BRB in AD.

Permeability of the BRB may be measured in live laboratory animal models of retinal diseases by the extent of leakage using fluorescent dyes with predefined molecular weight such as fluorescein (Do carmo et al., 1998), Evans blue (Xu et al., 2001) and others (Ivanova et al., 2019). In the clinical setting, fundus fluorescein angiography (FFA) utilizing a fluorescent dye and fundus camera has been an established method for examining retinal vascular circulation and BRB damage (Marmor and Ravin, 2011). Recently, a newly modified OCT method was developed, the OCT-Leakage. By using a proprietary algorithm to identify sites of decreased optical reflectivity, the system quantifies and detects the correlation of retinal extracellular space with degrees of retinal edema (Cunha-Vaz et al., 2016; Cunha-Vaz, 2017). A live imaging study in 28 patients has demonstrated agreement between OCT-Leakage and FFA in identifying sites of impaired BRB in diabetes (Cunha-Vaz et al., 2017), providing a new noninvasive low-cost alternative method to detect and quantify BRB leakage.

Given that retinal vascular amyloidosis has been detected in AD (Koronyo et al., 2017; Shi et al., 2020), future studies should aim to examine if BRB permeability is also altered by disease,

and if this is a cause or effect of retinal vascular amyloidosis. Indeed, a study using OCT angiography in patients revealed increased fovea avascular zone and decreased foveal thickness in eyes of AD patients (O'Bryhim et al., 2018), implying extensive retinal microvascular damage in the AD retina. Accordingly, recent human studies have demonstrated that retinal vascular abnormalities can predict cognitive decline (Baker et al., 2007; Cabrera DeBuc et al., 2018; Deal et al., 2018). Recent progress in retinal amyloid imaging (Koronyo et al., 2017), pericyte imaging by adaptive optics (Schallek et al., 2013) together with FFA and the recently developed OCT-Leakage (Cunha-Vaz et al., 2016) should allow for a comprehensive assessment of retinal A β pathology and BRB damage, potentially revolutionizing AD screening techniques.

RETINAL INFLAMMATION IN ALZHEIMER'S DISEASE

Chronic, low-grade inflammation is a typical sign of Alzheimer's neuropathology (Akiyama et al., 2000). Neuroinflammation in AD is linked with increased astrocyte and microglia reactivity and neurotoxicity, A β and tau seeding and propagation, as well as with microglia-mediated synaptic pruning (Cardona et al., 2006; Wyss-Coray, 2006; Stevens et al., 2007; Fuhrmann et al., 2010; Lee et al., 2010; Wyss-Coray and Rogers, 2012;

Hong et al., 2016; Salter and Stevens, 2017; Ising et al., 2019; Friker et al., 2020). As in the brain, some histological studies have implicated inflammation in the retina of AD patients (Liew et al., 1994; Blanks et al., 1996a; Grimaldi et al., 2018). This is particularly interesting as the eye has historically been considered an immune privileged site (Zhou and Caspi, 2010). In 1996, Blanks and colleagues discovered increased GFAP expression in retinal astrocytes and Müller cells in the GCL of AD retina, suggesting astrogliosis occurs in the retina of these patients (Blanks et al., 1996a). More recently, detrimental astrocyte and microglial activation was observed along with AB plaques, tau tangles, and neurodegeneration in postmortem retinal tissues of 6 AD patients as compared with 6 control subjects (Grimaldi et al., 2019). Notably, molecular mediators of innate immunity including interleukin-1ß (IL-1ß), complement component 3 (C3), osteopontin, and triggering receptor expressed on myeloid cells 2 were found to be upregulated in retinal tissues of AD patients (Grimaldi et al., 2019).

Although much work in larger retinal samples is needed to investigate the nature and potential mechanisms of retinal inflammation in human AD, various studies in animal models of AD have also provided evidence and insights into retinal inflammation in this disorder. Ning and colleagues first established the correlation of neurodegeneration and inflammation in the retina of ADtg mice (Ning et al., 2008). Other studies have shown increased microgliosis and astrocytosis, infiltration of lymphocytes and monocytes, and upregulation of monocyte chemoattractant protein-1 (MCP-1) in multiple layers of the retina and choroid (Ning et al., 2008; Liu et al., 2009; Perez et al., 2009; Yang et al., 2013; Edwards et al., 2014; Tsai et al., 2014; Antes et al., 2015; Gao et al., 2015; Pogue et al., 2015). A recent study in 3xTg AD mice has further described morphological changes in retinal microglia, including increased microglial cell number, soma size, retraction and reorientation of microglial processes, and change in cell locations (Salobrar-Garcia et al., 2020).

In general, retinal inflammation is implicated in multiple traditional retinal vascular and neurodegenerative disorders, including diabetic retinopathy (Tang and Kern, 2011; Semeraro et al., 2015; Rubsam et al., 2018), AMD (Knickelbein et al., 2015; Kauppinen et al., 2016), and glaucoma (Vohra et al., 2013). During the onset of disease pathogenesis, retinal inflammation is usually triggered by an imbalance of pro- versus antiinflammatory molecules. This can be evoked by a wide spectrum of pathogenic pathways, including overproduction of reactive oxygen species, activation of NF-κB or protein kinase C pathways, inflammasome or microglial activation, advanced glycation end products, or shear pressure and leukocyte invasion due to retinal microvascular damage. In this context, the recent discovery of early retinal pericyte loss in MCI patients (Shi et al., 2020) suggests an early BRB disturbance and perhaps retinal microvascular leakage in AD pathogenesis that may be implicated in retinal inflammation. Future studies should evaluate BRB leakage and the potential relationship with imbalanced retinal inflammatory pathway activation and brain inflammation in AD. Findings from such studies could lead to the discovery of novel retinal biomarkers to facilitate AD detection and monitoring.

FUNCTIONAL AND VISUAL CHANGES IN ALZHEIMER'S DISEASE

As discussed above, the retina undergoes neuropathological changes similar to the brain in AD patients. This is not surprising given the developmental, physiological, and anatomical similarities between the two tissue types, rendering them vulnerable to AD-related neuronal and functional abnormalities (Byerly and Blackshaw, 2009; London et al., 2013; Jindal, 2015). A host of visual and ocular manifestations were reported in both MCI and AD patients, particularly loss of contrast sensitivity, color discrimination deficits, difficulties with depth and motion perception, circadian rhythm irregularities, and sleep disturbances (Sadun et al., 1987; Katz and Rimmer, 1989; Trick et al., 1989; Cronin-Golomb et al., 1991; Cronin-Golomb, 1995; Lakshminarayanan et al., 1996; Cormack et al., 2000; Rizzo et al., 2000; Pache et al., 2003; Ng et al., 2007; Armstrong, 2009; Salamone et al., 2009; Chang et al., 2014; Nolan et al., 2014; Armstrong and Kergoat, 2015; Salobrar-Garcia et al., 2015; Hart et al., 2016; Javaid et al., 2016; La Morgia et al., 2016; Polo et al., 2017; Cerquera-Jaramillo et al., 2018; Colligris et al., 2018; Risacher et al., 2020). Importantly, retinal changes in AD patients such as RGC degeneration, reduced NFL thickness, and decreased blood circulation were linked to specific visual and ocular disturbances (Hinton et al., 1986; Blanks et al., 1989, 1996a,b; Parisi et al., 2001; Berisha et al., 2007; Paquet et al., 2007; Kesler et al., 2011).

Alzheimer's hallmark pathologies Aß deposits and tauopathy are found in retinal cell types and topographical regions shown to undergo degeneration and physiological abnormalities in both patients and transgenic animal models (Koronyo-Hamaoui et al., 2011; Schön et al., 2012; La Morgia et al., 2016; Koronyo et al., 2017; den Haan et al., 2018; Asanad et al., 2019b; Grimaldi et al., 2019; Habiba et al., 2020; Shi et al., 2020). These findings provide a link between AD-related retinal neuropathology and the visual deficits reported in living patients. For instance, histological examination of retinal tissue shows colocalization of pTau and Aβ pathology in regions associated with RGC loss (Schön et al., 2012; La Morgia et al., 2016; Koronyo et al., 2017; den Haan et al., 2018; Asanad et al., 2019b; Grimaldi et al., 2019; Shi et al., 2020). Specifically, degeneration of RGCs would undoubtedly compromise their essential role in receiving sensory input from photoreceptors via bipolar cells, and in early stages of visual information processing and conveyance to the brain (Demb and Singer, 2015). In support of this, a recent study utilizing OCT and pattern electroretinography (ERG) in AD patients showed a linear relationship between NFL thickness and the bioelectrical integrity of RGCs and their nerve fibers, which are essential for afferent signal transduction (Ngoo et al., 2019).

In addition to the above reported changes, irregularities in efferent ocular pathways in conjunction with retinal abnormalities were also associated with altered pupillary light response (PLR) in patients. These alterations include increased latency of pupillary constriction to light, decreased constriction amplitude, faster redilation after light offset, decreased maximum constriction velocity and acceleration (Chougule et al., 2019), and altered pupil dilation response during cognitive tasks (Granholm et al., 2017; Kremen et al., 2019). Therefore, there is growing interest in exploring various aspects of pupillary responses, using specialized pupillometry tools, in pre-clinical and clinically diagnosed AD patients. Further to being governed by both the sympathetic and parasympathetic systems, PLR is also regulated by retinal rods, cones, and mRGCs (Hattar et al., 2003). In line with this, altered PLR in AD patients was associated with retinal mRGC loss (La Morgia et al., 2016), highlighting the consequences of retinal AD pathology for various ocular and visual dysfunctions. Moreover, these intrinsically photosensitive RGCs are also reported to regulate light photoentrainment of circadian rhythms, supported by their tightly regulated communication with photoreceptors and their projections to the central circadian clock in the hypothalamic suprachiasmatic nucleus (SCN) (Lu et al., 1999; Berson et al., 2002; Gooley et al., 2003; Markwell et al., 2010; Chen et al., 2011). Therefore, retinal mRGC loss is suggested to play a key role in the sleep-wake cycle dysfunctions observed in AD patients (La Morgia et al., 2016, 2017). Sleep abnormalities reported in MCI and AD patients include prolonged sleep latency, reduced total sleep duration and rapid eye movement (REM) sleep, sleep fragmentation, frequent awakenings and lower melatonin levels at night, and daytime somnolence (Petit et al., 2004; Ju et al., 2013; Peter-Derex et al., 2015; Feng et al., 2016; La Morgia et al., 2016, 2017; Weissová et al., 2016; Asanad et al., 2019b). Intriguingly, there are reports of sleep disturbances exacerbating or even preceding cognitive impairments (Ju et al., 2014; Bubu et al., 2017; Brzecka et al., 2018). The essential discovery that A β accumulates within or in close proximity to mRGCs, which undergo degeneration and dendrite diameter loss, at least in part, can explain impaired sleep patterns and altered pupil dilation in AD patients (La Morgia et al., 2016). These findings also suggest that certain retinal cells are more susceptible to Aβ-induced neurotoxicity and proposes the first retinal damage-based mechanism for functional disturbances commonly seen in AD patients.

The spatial distribution of retinal AD pathology may indicate the specific functional deficit. Regionally, both AB and pTau pathologies appear to preferentially affect the peripheral superior and inferior retinal quadrants in AD patients (Koronyo-Hamaoui et al., 2011; La Morgia et al., 2016; Koronyo et al., 2017; den Haan et al., 2018). This, along with consistent neuronal loss, degeneration of mRGCs and RGCs, and NFL thinning in these particular peripheral regions, could underlie visual abnormalities such as in contrast sensitivity, motion perception and circadian rhythms described in AD (Trick et al., 1989, 1995; Blanks et al., 1996a; Berisha et al., 2007; Lu et al., 2010; Kesler et al., 2011; Moschos et al., 2012; Ju et al., 2013; Kirbas et al., 2013; Risacher et al., 2013; Kromer et al., 2014; La Morgia et al., 2016). Beyond AD-related manifestations predominantly in the inner layers and peripheral retinal regions, deposits of AB were found in close proximity to blood vessels and within vessel walls (Koronyo-Hamaoui et al., 2011; La Morgia et al., 2016; Koronyo et al., 2017; Shi et al., 2020). Furthermore, vascular amyloidosis positively correlated with the degree of vascular PDGFR- β deficiency as well as A β_{40} and AB42 alloforms accumulated within degenerating pericytes in MCI and AD retinas (Koronyo et al., 2017; Shi et al., 2020). These

findings, together with altered retinal neurovascular coupling and structural vascular abnormalities in clinically diagnosed MCI and AD patients (Berisha et al., 2007; Querques et al., 2019), implicate retinal vascular-associated A β in causing or amplifying retinal metabolic dysfunctions and impaired clearance pathways. These changes may lead to functional-visual abnormalities.

A study in monkeys has recently provided evidence for the involvement of mRGCs in blue color sensitivity (Dacey et al., 2005). AD patients are reported to experience altered color vision with a selective and severity-independent deficiency in blue hue discrimination (Cronin-Golomb et al., 1993; Wijk et al., 1999; Rizzo et al., 2000; Chang et al., 2014; Polo et al., 2017). Interestingly, impaired blue color discrimination also manifests in patients with other neurodegenerative disorders (Paulus et al., 1993; Haug et al., 1995; Birch et al., 1998; Rodnitzky, 1998; Melun et al., 2001; Anssari et al., 2020).

The second most common cause of irreversible vision loss is glaucoma, a condition characterized by optic neuropathy, apoptotic RGC degeneration and NFL damage, driven by increased IOP and accumulation of proteinaceous deposits of AB and pTau (Cordeiro et al., 2004). Indeed, a higher occurrence of glaucoma was reported in AD patients compared to controls (Bayer et al., 2002; Tamura et al., 2006). Diminished contrast sensitivity, altered motion and depth perception, and oculomotor disturbances in both AD and glaucoma patients are partially attributed to a preferential loss of the functionally relevant magnocellular RGCs (M-cells) and their nerve fibers (Sadun and Bassi, 1990; Fiorentini et al., 1996; La Morgia et al., 2017). Moreover, impaired contrast sensitivity in MCI patients was shown to significantly correlate with the cerebral burden of amyloid and tau pathology (Risacher et al., 2020). Given the higher concentration of rods in the peripheral retina and their importance in motion and depth perception, low-contrast sensitivity, and peripheral vision, it is not surprising that these visual functions may be adversely influenced by the increased levels of A β deposits and pTau in the periphery (Gilmore et al., 1994; Risacher et al., 2013). Cumulative evidence of vision loss and ocular impairments accompanied by progressive retinal changes in AD and other neurodegenerative disorders, including glaucoma, multiple sclerosis (MS), PD, and Huntington's disease (HD), robustly support the involvement of specific retinal pathologies in functional and visual manifestations of these conditions (Fisher et al., 2006; Gupta et al., 2008; Archibald et al., 2009; Green et al., 2010; London et al., 2013; Kersten et al., 2015; Andrade et al., 2016; Weil et al., 2016; La Morgia et al., 2017).

Despite many recent advancements, the field of AD retinopathy is still in its infancy and more studies focusing on mechanistic and causative factors are necessary to determine the connection between specific retinal pathologies, neuronal-type impairments and visual-functional deficits.

RETINAL IMAGING

The mounting case for AD pathology in the retina has motivated investigations to develop various retinal imaging modalities, beginning with blue-light high-resolution photography by Tsai



used to quantify spot number and fluorescent area (μ m²). (**B**,**C**) color-coded overlay images from CN (**B**) and AD (**C**) retinas with curcumin-positive amyloid deposits above threshold shown in red, spots above 1:1 reference but below threshold shown in green and spots below reference in blue. (**D**) Representative graph showing number of color-coded spots (described in **B**,**C**) in AD retina used to determine retinal amyloid index (RAI) score. (**E**) Comparison of RAI scores in AD patients (*n* = 6) and age-matched CN subjects (*n* = 5). Data shown as group mean ± SEM. ***P* < 0.005, unpaired 2-tailed Student's *t*-test. Reproduced from Koronyo et al. (2017) with permission of ASCI via Copyright Clearance Center.

et al., 1991, which indicated a correlation between Alzheimer's Disease Assessment Scale (ADAS) scores and optic nerve head changes in AD patients (Tsai et al., 1991).

Further studies employed cross-sectional imaging by optical coherence tomography (OCT) or optical coherence tomography angiography (OCTA) (Parisi et al., 2001; Kromer et al., 2014; Cunha et al., 2016; Ferrari et al., 2017; Polans et al., 2017; Polo et al., 2017; Bulut et al., 2018; Janez-Escalada et al., 2019; Salobrar-Garcia et al., 2019), which revealed structural abnormalities and cell neurodegeneration in the retina.

As mentioned above, the first *in vivo* imaging of AD hallmark pathology was initially developed and tested in transgenic murine models of AD by Koronyo-Hamaoui and colleagues (Koronyo-Hamaoui et al., 2011; Koronyo et al., 2012). The researchers paired curcumin – a natural and safe fluorochrome that labels $A\beta$ fibrils and oligomers with high affinity and specificity - with a modified rodent retinal optical imaging microscope. Later, this approach was translated to humans and the feasibility to noninvasively detect and quantify individual retinal amyloid plaques was demonstrated in living patients with a modified confocal scanning laser ophthalmoscope following oral administration of highly bioavailable Longvida curcumin (Figures 7A-D; Garcia-Alloza et al., 2007; Gota et al., 2010; Koronyo-Hamaoui et al., 2011; Koronyo et al., 2012, 2017). Studies in murine ADtg models demonstrated the feasibility to longitudinally monitor individual AB deposits including their appearance and clearance during disease progression and in response to immune-based therapy (Koronyo et al., 2012).

Importantly, in the proof-of-principal clinical trial, this noninvasive amyloid imaging technique revealed a significant 2.1-fold greater retinal amyloid burden, termed as retinal amyloid index (RAI), in a group of 10 AD patients as compared to 6 age-matched healthy controls (Figure 7E; Koronyo et al., 2017). Supported by histological findings, retinal AB deposits were often found the mid- and far-periphery of the superior and inferior regions, where previously NFL thinning was reported as more pronounced. Further, amyloid deposits were found above the retinal pigment epithelium in the neurosensory retina, unlike the typical location of drusen (Koronvo et al., 2017). Anecdotal data from 2 AMD patients suggested that retinal curcumin fluorescence signals were diffuse and concentrated at the posterior pole, apparently distinct from findings in the retinas of AD patients (Koronyo et al., 2017). Moreover, results presented at the Alzheimer's Association International Conference on July 15, 2014, from a large cohort of over 150 MCI, AD, and cognitively normal participants of the Australian Imaging, Biomarker and Lifestyle (AIBL) Study (www.aibl.csiro.au) found that retinal amyloid fluorescence imaging predicted cerebral amyloid burden and was significantly higher in AD patients (Frost et al., 2014).

Expanding on this work, Kayabasi and colleagues detected abnormal A β deposits in 30 MCI patients using fundus autofluorescence imaging (FAF) and OCT (Kayabasi et al., 2014). An additional study utilizing OCT revealed a correlation between retinal inclusion bodies and cortical amyloid burden in pre-clinical patients via florbetapir PET and multiple retinal sd-OCT aggregation markers of possible disease burden (Snyder et al., 2016).

Noninvasive *in vivo* retinal hyperspectral imaging (rHSI) has also recently been employed in patients and controls. One study found significant differences in the retinal reflectance spectra of MCI patients with high A β burden, confirmed by brain PET imaging, as compared to age-matched controls, validated in an independent cohort with a second hyperspectral camera (Hadoux et al., 2019). Retinal imaging scores correlated with A β burden in the brain (Hadoux et al., 2019). Intriguingly, a similar study from More and colleagues, the team that initially developed the rHSI technique in murine models (More and Vince, 2015; More et al., 2016), found the largest deviation in rHSI signatures between MCI patients and controls, irrespective of other ocular conditions such as cataracts or glaucoma (More et al., 2019). The same technique was able to distinguish cerebral A β + subjects from A β - subjects with 85% accuracy based on retinal vascular measures including vessel tortuosity and diameter (Sharafi et al., 2019).

Finally, a recent multimodal approach including visual performance tests, advanced retinal imaging, and full-field electroretinogram (ERG) in 69 cognitively impaired subjects revealed sensitivity of the combined approach to cognitive decline (Cabrera DeBuc et al., 2018). Although more robust investigations are needed to validate these findings, this study supports the association of retinal geometric vascular and functional parameters with physiological changes in the retina in cognitively impaired individuals (Cabrera DeBuc et al., 2018). Future tools should assess whether a combination of these multimodal methods with specific retinal amyloid imaging will allow for earlier and/or more accurate assessment of AD.

CONCLUSION

With the number of AD patients estimated to triple by 2050 (Alzheimer's Association, 2019), a reliable, noninvasive and affordable diagnostic technique suitable for widespread clinical use is urgently needed. In recent years, a radical idea has emerged that AD pathology in the brain can also manifest and perhaps be mirrored in the retina. Indeed, the identification of hallmark $A\beta$ deposits in the retina of patients and numerous animal models and their correlation with pathology in the brain have fueled this field of neuro-ophthalmology in AD. Moreover, mounting evidence now supports the existence of diffuse, classical and neuritic-like plaques, pTau aggregates, $A\beta_{42}$ fibrils, protofibrils, and oligomer-like structures, pericyte loss, vascular A β_{40} and A β_{42} accumulation, inflammation and neurodegeneration in the retina of AD patients (Alexandrov et al., 2011; Koronyo-Hamaoui et al., 2011; Schön et al., 2012; Tsai et al., 2014; La Morgia et al., 2016; Koronyo et al., 2017; den Haan et al., 2018; Grimaldi et al., 2018; Hadoux et al., 2019; Schultz et al., 2020; Shi et al., 2020). Despite many recent successes, several limitations including the scarcity of retinal tissue from people with neuropathological data and the application of non-conventional histological and biochemical techniques to examine AD effects in various topographical regions, has historically restricted the knowledge in this field. Future investigations, involving brain biobanks with extended collection of ocular tissues, will allow researchers to determine how early AD pathological processes occur in the retina, their extent, distribution, and relationship to brain pathology.

Histological and biochemical examinations of postmortem retina from AD patients and animal models have proven invaluable in deciphering the pathological burden of disease hallmarks, susceptible retinal layers and cell types, and putative molecular pathways linked to retinal atrophy and functional deficits. Nevertheless, there is still lack of understanding of the impact of $A\beta$ and tau pathologies on retinal cell types; some effects, including nerve fiber thinning, neuronal loss and vascular changes, may not manifest in early stages of disease and may not be specific to AD. For instance, reduced NFL, macular, and foveal thickness have also been observed via OCT examinations of Parkinson's disease (PD) patient retinas as compared to normal controls (Inzelberg et al., 2004; Altintas et al., 2008; Hajee et al., 2009; Cubo et al., 2010; Moschos et al., 2011, 2012; Shrier et al., 2012; Adam et al., 2013; Moreno-Ramos et al., 2013; Satue et al., 2013; Moschos and Chatziralli, 2017).

Notably, according to the 2018 NIA-AA research framework, the presence of A β deposits is a requirement for Alzheimer's pathological change diagnosis and should be considered in early screening efforts and recruitment of individuals for clinical trials (Jack et al., 2018). The ability to detect and quantify A β deposits and related pathologies via noninvasive retinal imaging along with sensitive, routine, minimally invasive plasma AD biomarkers show great promise for diagnostic screening, monitoring of progression, and assessment of therapeutic efficacy (Hampel et al., 2018; Baldacci et al., 2020). The work described here combined with future advances may lead to clinical translation, discrimination of pathophysiological phenotypes during the AD continuum, and eventually a highly anticipated cure for this destructive disease.

Future investigations should focus on standardization of histological techniques, identification of early AD biomarkers in the retina, their spatiotemporal profiles, and enhancing the sensitivity of retinal imaging modalities for detection of these pathological indicators in this new and expanding field of Alzheimer's retinopathy.

AUTHOR CONTRIBUTIONS

NM, HS, MO, and MK-H drafted manuscript. NM, HS, JD, YK, and MK-H did the figure and image preparation. NM, HS, MO, JD, AR, D-TF, JS, KB, YK, and MK-H edited the manuscript. MK-H did the study supervision. All authors read and approved the final manuscript.

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Conflict of Interest: YK, MK-H, and KB were co-founders and stockholders of NeuroVision Imaging, Inc., 1395 Garden Highway, Suite 250, Sacramento, CA, United States.

The remaining authors declare that the research was conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest.

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