

Annual Review of Vision Science [Withrawn] Retinal Vasculature in Development and Diseases

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Abstract

The retina is one of the most metabolically active tissues in the body, consuming high levels of oxygen and nutrients. A well-organized ocular vascular system adapts to meet the metabolic requirements of the retina to ensure visual function. Pathological conditions affect growth of the blood vessels in the eye. Understanding the neuronal biological processes that govern retinal vascular development is of interest for translational researchers and clinicians to develop preventive and interventional therapeutics for vascular eye diseases that address early drivers of abnormal vascular growth. This review summarizes the current knowledge of the cellular and molecular processes governing both physiological and pathological retinal vascular development, which is dependent on the interaction among retinal cell populations, including neurons, glia, immune cells, and vascular endothelial cells. We also review animal models currently used for studying retinal vascular development.

1. RETINAL VASCULATURE IN DEVELOPMENT

1.1. Retinal Vessels in the Eye

The retina is supplied by two vascular systems: The central retinal artery supplies the inner retina and the choriocapillaris supplies the retinal pigment epithelium and outer retina (consisting primarily of photoreceptors) (**Figure 1**). The central retinal artery travels along the inferior margin of the optic nerve sheath and then enters the eye through the center of the optic nerve. The artery then branches in the inner retina to form three capillary layers (Netter 2006). The retinal vessels provide blood to the inner retinal neurons. The avascular photoreceptor layer relies on the choriocapillaris lying beneath the retinal pigment epithelium to supply oxygen by diffusion.

The choroid, a densely pigmented vascular sinusoidal plexus, originates from two ophthalmic artery branches: the short posterior ciliary arteries, which supply the posterior portion of the choroid, and the long posterior ciliary arteries, which supply the anterior choroid, ciliary body, and iris. The choroidal vasculature has three interconnected distinct layers: the innermost choricapillaris, the intermediate Sattler's layer, and the outermost Haller's layer (Hartnett 2013). It composes up to 85% of the blood volume in the eye (Anand-Apte & Hollyfield 2011).



Figure 1

Schematic diagram of the ocular vasculature. (*a*) Cross-sectional image of an eye. (*b*) An enlarged cross-sectional view of the retinal and choroidal vasculature. Three interconnected layers of retinal vessels are embedded among inner retinal neurons: The superficial retinal vasculature lies in the NFL; the intermediate and deep retinal vascular networks lie along each side of the INL. The choroidal vessels are located beneath the RPE and Bruch's membrane and supply oxygen and nutrients to the outer portion of the retina, which includes primarily photoreceptors in the outer nuclear layer. (*c*) Retinal vascular development begins with the formation of the superficial vascular plexus from p1 to p10, then the deep vascular plexus from p8 to p12, and finally, the intermediate vascular plexus from p14 to p20. (*d*) Retinal superficial vascular plexus at different developmental stages p0, p3, p7, and p14. Panels *a* and *b* adapted from Liu et al. (2017). Panels *c* and *d* adapted from Joyal et al. (2018). Abbreviations: GCL, ganglion cell layer; INL, inner nuclear layer; IS/OS, inner segment/outer segment of photoreceptor; NFL, nerve fiber layer; ONL, outer nuclear layer; P, postnatal day; RPE, retinal pigment epithelium.

1.2. Development of Retinal Vasculature

During embryonic development, the hyaloid vasculature supplies oxygen and nutrients to the developing inner retina and the lens during its maturation. The hyaloid artery enters the embryonic fissure, extends through the vitreous humor, and surrounds the lens (Hartnett 2013). The hyaloid vessels appear at the fourth week of gestation and reach maximum prominence at the ninth week in the human fetus. When the retinal vasculature develops during midgestation, the hyaloid vessels regress concurrently. The central hyaloid extension from the optic disk to the posterior lens surface regresses last (Hartnett 2013). Retinal vascular development is completed before birth in humans—unlike in mice and rats, where development occurs postnatally to eventually form a vascular network structurally similar to that in primates (Stone et al. 1995).

In humans, retinal vascularization starts in utero at about 16 weeks of gestation and is completed at approximately 40 weeks of gestation, just before birth. The most superficial retinal vascular layer forms first, starting from the optic nerve head and progressing toward the peripheral edge of the retina, reaching the nasal side of the ora serrata at about 36 weeks of gestation. As the superficial layer nears completion, retinal vessels dive into the retina to form the deep retinal vascular layer at the base of the outer plexiform layer. The intermediate layer then forms between the superficial and deep layers, superficial to the inner plexiform layer, forming a well-organized network to complete the three vascular layers (Dorrell et al. 2009). In other mammals, the same pattern of three retinal vascular layers forms, although over varying timescales (Gariano & Gardner 2005). In mice, the superficial plexus starts to form during the first week after birth. During the second week, angiogenic sprouts from the superficial layer grow into the retina to form the deep layer followed by the intermediate layer of the retinal capillary networks, and the entire retinal vasculature matures by three weeks after birth (Stahl et al. 2010a).

Human choriocapillaris is unusual in its development compared with other capillary beds in the body (Chirco et al. 2017), developing via hemovasculogenesis between 6 and 8 weeks of gestation, before the formation of intermediate or large vessels. At 11 to 12 weeks of gestation, the development of intermediate choroidal vessels within Sattler's layer occurs by angiogenesis; this process allows the choriocapillaris to connect with the larger, outer choroidal vessels of Haller's layer. Three distinct layers of choroidal vessels are seen at 21 weeks of gestation, which coincides with the beginning of photoreceptor cell differentiation. Interestingly, the three layers of the choroid (i.e., choriocapillaris, Sattler's layer, and Haller's layer) are found only in the posterior pole. The choroid outside the posterior pole consists of only the choriocapillaris and a layer of larger vessels (Hasegawa et al. 2007). The choriocapillaris appears fully mature, with flat, thinwalled fenestrated vessels, at 22 weeks of gestation (Baba et al. 2009, Hasegawa et al. 2007; for a review, see Lutty et al. 2010).

Retinal blood vessels in primates form a four-lobed pattern along the major vascular arcades and do not appreciably cross the horizontal raphe (Gariano 2010). Nasal–temporal retinal asymmetry is absent primarily because of the contour of perifoveal vessels. Ganglion cell axon bundles and an astrocytic plexus in the nerve fiber layer show a similar pattern. Murine retinal vessels, in contrast, are simply radially organized, without reference to a horizontal midline or macula (Gariano 2010).

2. RETINAL VASCULATURE IN DISEASES

Pathological blood vessels in the eye pose threats to normal vision. Angiogenesis, important in both physiological vascular development and pathological neovascularization, occurs as endothelial cells proliferate and form new vessels following guidance cues and angiogenic stimulators and inhibitors (Potente et al. 2011). Dysregulated angiogenesis disrupts delivery of oxygen and nutrients,

resulting in unbalanced metabolic demand and supply and disturbed neural retinal function. Pathological angiogenesis is associated with many diseases, including cancers, cardiovascular diseases, neurodegeneration, and proliferative retinopathies (Folkman 1995). Pathological retinal neovascularization is characterized by leaky and tuft-like vessels, which are associated with retinal exudates and hemorrhage, leading to retinal damage, retinal detachment, or both (Al-Latayfeh et al. 2012). Abnormal ocular angiogenesis can occur in a broad spectrum of eye disorders, such as retinopathy of prematurity (ROP), diabetic retinopathy (DR), neovascular age-related macular degeneration (AMD), neovascular glaucoma, and corneal neovascularization (Al-Latayfeh et al. 2012).

2.1. Retinopathy of Prematurity

ROP is a major cause of blindness in children, affecting approximately 16,000 infants per year in the United States (Good 2004). Approximately 10% of children worldwide are born preterm (Blencowe et al. 2012), many with lifelong visual impairment. ROP contributes to 6–18% of cases of blindness in the United States (Lad et al. 2009). ROP is a biphasic disease, with an initial phase of vessel loss followed by a second phase of vessel proliferation. Because retinal blood vessels start developing in humans during the fourth month of gestation and reach the retinal periphery just before birth, infants born prematurely have incompletely vascularized retinas with a peripheral avascular zone at birth. The relative hyperoxic environment after birth compared with that in utero suppresses the normal secretion of angiogenic growth factors, leading to regression of existing vessels, the vessel loss phase of ROP. In the absence of an adequate vascular system between birth and gestational age of approximately 30–32 weeks, the retina becomes ischemic, leading to tissue hypoxia and secretion of hypoxia-driven angiogenic growth factors such as vascular endothelial growth factor (VEGF). The vessel-proliferation phase of ROP begins at about 32– 34 weeks of gestational age. In severe ROP, retinal neovessels can cause retinal detachment, leading to blindness.

2.2. Diabetic Retinopathy

DR, the leading cause of blindness in adults 24–70 years of age, eventually occurs in approximately one-third of patients with diabetes (Cheung et al. 2010). The prevalence of DR is higher in patients with type 1 diabetes than in those with type 2 diabetes (Yau et al. 2012) perhaps because of the longer duration of disease. Hyperglycemia and other metabolic dysregulation lead to the development of microangiopathy, including microaneurysms, hemorrhages, and basement membrane thickening (Nentwich & Ulbig 2015). This results in increased vascular permeability of vessels, causing leakage and macular edema (Cheung et al. 2010). Increased capillary occlusion results in retinal ischemia, triggering an increase in the levels of VEGF, which then promotes pathological neovascularization (Cheung et al. 2010, Nentwich & Ulbig 2015).

Classification of DR as either nonproliferative or proliferative is based on the presence of neovascularization. Nonproliferative DR, which can progress to preproliferative DR, exhibits microaneurysms, dot and blot hemorrhages, cotton-wool spots, and capillary nonperfusion owing to microvascular damage and pericyte loss. Microglial changes and diabetic macular edema can also occur in nonproliferative DR. In proliferative DR, neovascularization can result in retinal and vitreous hemorrhages and can lead to retinal detachment (Cheung et al. 2010, Olivares et al. 2017).

2.3. Age-Related Macular Degeneration

Choroidal neovascularization (CNV) and retinal angiomatous proliferation (RAP) are features of neovascular AMD (Campochiaro 2015). CNV lesions refer to the abnormal new vessel growth

from the choriocapillaris, extending through Bruch's membrane into the subretinal pigment epithelium or subretinal spaces. Left untreated, CNV may advance to a stage that forms fibrotic scars underlying the retina, leading to vision loss (Grossniklaus et al. 2010). The progression of CNV is as variable as its response to certain stimuli, depending on the underlying diseases and the stage of CNV development (Grossniklaus et al. 2010). Disruption or degradation of Bruch's membrane is one of the early stages of CNV and is followed by proliferating choroidal neovascular tissue invading the subretinal space. Pathological subretinal vessel formation can originate not only from the choroid but also sometimes from the retinal vasculature. Abnormal intraretinal angiogenesis invading the subretinal space with formation of a retinal–choroidal anastomosis is characteristic of a distinct variant of neovascular AMD known as RAP, which is present in approximately 12–15% of patients with neovascular AMD (Bottoni et al. 2005).

2.4. Other Vascular Eye Diseases

Macular telangiectasia type 2 is a disease of unknown cause with characteristic alterations of the macular capillary network and neurosensory retina. The disease manifests initially temporal to the foveal center but may later encompass an oval area centered on the foveola (Charbel Issa et al. 2013). In several rare human eye diseases, including familial exudative vitreoretinopathy and Norrie disease, incomplete peripheral retinal vasculature and lack of deep retinal vascular layers are observed (Criswick & Schepens 1969). There are many other eye diseases with abnormal angiogenesis, and they have been reviewed elsewhere (Bek 2013, Usui et al. 2015).

3. ANIMAL MODELS OF RETINAL VASCULAR DISEASES

The past several decades of angiogenesis research produced several animal models of ocular angiogenesis that mimic various aspects of neovascular eye diseases with retinal and choroidal angiogenesis. Mouse models of oxygen-induced retinopathy (OIR) and laser-induced CNV are two commonly used models for investigating pathological neovascularization in the retina and choroid, respectively. Additionally, transgenic knockout or knockin mice with intraretinal or subretinal neovascularization were developed as experimental tools for studying retinal vascular diseases.

3.1. Animal Models of Retinopathy of Prematurity

The OIR model was established to mimic ROP in humans. It reproduces the two phases of the disease: an initial vaso-obliteration phase with oxygen exposure and a subsequent hypoxiainduced neovascularization phase. The OIR models were established in several vertebrate animals, including cat (Ashton et al. 1954), mouse (Smith et al. 1994), rat (Penn et al. 2001), dog (McLeod et al. 1996), and fish (Cao et al. 2008). These animal models share the basic principle that exposure to high oxygen induces loss of existing retinal vasculature (Stahl et al. 2010a). After being returned to normoxic conditions with inadequate vessels to support retinal neurons, retinas become hypoxic and signal new (abnormal) blood vessel formation.

The cat model of ROP, developed in the 1950s by Ashton et al. (1954), was the first model to use high oxygen concentration to suppress retinal vascular development. Kittens are exposed to 70–80% oxygen for 4 days, causing retinal vascular obliteration, and then returned to normal room air exposure (21% oxygen), resulting in hypoxia-induced vasoproliferation (Ashton et al. 1954). In the less commonly used dog model of OIR, puppies are exposed to 95–100% oxygen for 4 days and returned to normoxic conditions. The pattern and severity of the vascular reaction

to hyperoxia in this model are similar to those observed in the kitten model and in human infants (McLeod et al. 1996).

The advantage of genetic manipulation in mice makes the mouse model of OIR (Smith et al. 1994) one of the most commonly used ROP models. In this model, neonatal mice are exposed to 75% oxygen for 5 days, from postnatal day 7 to 12, and subsequent normal room air exposure from postnatal day 12 to 17. With hyperoxia exposure, the formed retinal vessels regress in the central zone, leading to vaso-obliteration, and vascular growth ceases in the periphery, mimicking the initial phase of ROP. After the pups are returned to normoxic conditions at postnatal day 12, the vaso-obliterated area becomes ischemic and hypoxic, resulting in upregulation of proangiogenic pathways, such as hypoxia-inducible factor (HIF)-dependent pathways (Pierce et al. 1995, Stahl et al. 2010a), including VEGF and erythropoietin (Aiello et al. 1995, Chen et al. 2008, Pierce et al. 1995, Robinson et al. 1996). This leads to both physiological revascularization of the vasoobliterated area and pathological neovascularization (Smith et al. 1994). In the mouse model of OIR, pathological neovessel growth reaches the maximal severity at postnatal day 17. After postnatal day 17, the abnormal neovascularization gradually regresses and completely disappears by about postnatal day 25 (Connor et al. 2009). Vaso-obliteration and neovascularization can be quantified as a percentage of the total retinal area in retinal whole mounts stained with isolectin IB₄ to visualize blood vessels (Connor et al. 2009, Stahl et al. 2009). Vascular leakage can be visualized by fluorescein angiography and quantified by the Miles assay measuring permeability (J. Chen et al. 2011a, Pierce et al. 1995, Stahl et al. 2009, Vähätupa et al. 2016). The mouse model of OIR was used in early work investigating the role of VEGF in ROP (Aiello et al. 1995; Pierce et al. 1995, 1996; Robinson et al. 1996) as well as those factors critical in the third trimester and missing after preterm birth, such as insulin-like growth factor (Hellström et al. 2001; Smith et al. 1997, 1999) and omega-3 polyunsaturated fatty acids (Connor et al. 2007; Fu et al. 2015; Sapieha et al. 2011, 2012; Shao et al. 2014; Smith et al. 1999; Stahl et al. 2010b).

Penn et al. (2001) developed a rat model of OIR utilizing alternating hyperoxia–hypoxia cycles in which the oxygen levels are cycled between 50% and 10% every 24 h from birth to postnatal day 14, and then the pup is exposed to room air through postnatal day 20. The cycling levels of oxygen slow retinal vascular development and lead to a peripheral avascular zone. After returning to room air, pathological neovessels grow at the boundary between vascularized and avascular areas in the midperipheral retina. In the rat model of OIR, fluctuations in oxygen concentrations reflect variable oxygen levels measured in preterm human infants with severe ROP (Hartnett & Penn 2012). Vaso-obliteration in the peripheral retina is geographically similar to the peripheral avascular zone observed in human ROP (Grossniklaus et al. 2010). Overall, the rat model shows many clinically relevant features of ROP. However, it is difficult to genetically manipulate rats, and the levels of neovascularization are relatively low compared with those of the mouse model of OIR, limiting its use in screening antiangiogenic compounds.

3.2. Animal Models of Diabetic Retinopathy

Several animal models have been developed to investigate the etiology and pathogenesis of DR and to create and test therapies to treat the disease. Animal models of DR are generated through surgical injury, laser or chemical damage, drugs, or diet. Genetic models are created with selective gene editing in rodents (Olivares et al. 2017).

Induced models include surgical removal of the pancreas, administration of the drug alloxan, administration of the drug streptozotocin, high-galactose diets, and laser or chemical damage to the eye (Grossniklaus et al. 2010). Of all the methods of induction, the most common is administration of streptozotocin, as it results in the fastest onset of disease (Rakieten et al. 1963).

The most frequently used animal models for inducing DR are mice and rats, but dogs, cats, pigs, rabbits, monkeys, and zebrafish are also used.

There are also several genetic modes of DR in mouse, rat, and zebrafish. These models include spontaneous, strain-specific, and genetically edited mutations. Several inbred mouse strains—for example, nonobese diabetic (NOD) and db/db (Lepr^{db})—exhibit hyperglycemia, one of the main characteristics of diabetes. Genetic mouse models of DR are relatively easy to maintain, have well-characterized genetic backgrounds, and can be manipulated to generate knockout or transgenic models. Genetic models exist for both type 1 and type 2 diabetes, and models for type 2 diabetes can be either obese or nonobese (Weir et al. 2009).

Five commonly used genetic mouse models of DR are $Ins2^{Akita}$, NOD, db/db (Lepr^{db}), Kimba, and Akimba (Olivares et al. 2017). The $Ins2^{Akita}$ mouse is a model of type 1 diabetes that harbors a missense mutation in the gene *Ins2*, which leads to a conformational change in the insulin protein. Ins2 accumulates in pancreatic β cells, leading to β cell death (Izumi et al. 2003). The NOD mouse model of type 1 diabetes exhibits an autoimmune response in which CD4⁺ and CD8⁺ cells attack pancreatic β cells (Serreze et al. 1997). The db/db (Lepr^{db}) mouse was developed to study type 2 diabetes. db/db mice harbor a mutation in the leptin receptor and develop hyperglycemia and obesity after 4–8 weeks (Chen et al. 1996, Hummel et al. 1966), and disease progression continues for 10 months. A mouse model of DR with hyperglycemia-induced background vascular changes (such as pericyte loss) but with non-hyperglycemia-induced neovascularization can be obtained by breeding two mutant mouse strains. The Akimba mouse was generated by crossing Kimba mice, a nondiabetic model of proliferative retinopathy resulting from overexpression of *Vegf* driven by the rhodopsin promoter (Tee et al. 2008), with the diabetic $Ins2^{Akita}$ mice to create a model of diabetes with retinal neovascularization secondary to non-diabetes-driven expression of *Vegf* in photoreceptors.

Genetic rat models of DR include Zucker diabetic fatty (ZDF), Otsuka Long-Evans Tokushima fatty (OLETF), biobreeding (BB), WBN/Kob, spontaneously diabetic Torii (SDT), and Goto-Kakizaki (GK). ZDF, OLETF, and BB are monogenic models of DR with independent mutations that perturb different nodes of the DR disease pathway. SDT, WBN/Kob, and GK, in contrast, are polygenic models. These models demonstrate the genetic complexity of DR (Olivares et al. 2017).

3.3. Animal Models of Neovascular Age-Related Macular Degeneration

Ideal animal models of CNV should be stable, efficient, and reproducible and exhibit pathological features similar to those observed in CNV lesions in AMD patients. Taking into account the complex conditions that lead to CNV in humans, three main types of animal models of CNV have been developed: laser-induced CNV, surgical-incision-induced CNV, and transgenic animal models. Laser-induced CNV is one of the most widely used models recapitulating CNV seen in neovascular AMD. However, the CNV produced is in response to injury and does not reflect age-driven disease. The laser-induced CNV model is relatively rapid.

Laser-induced CNV models have been used extensively in the rat and mouse to study key pathways involved in CNV formation and to evaluate pharmacologic antiangiogenic intervention (Takahashi et al. 2000). In the mouse model, laser photocoagulation spots are placed through the dilated pupil, avoiding the major retinal vessels, around an area 2-disc diameters from the optic nerve. CNV lesions reach maximal volume and area on day 7 after the laser burn, forming diffuse subretinal fibrovascular tissue, which can be visualized and quantitatively analyzed for leakage and lesion size by means of fluorescein fundus angiography and immunohistochemical staining (Gong et al. 2015). After day 7 post laser, CNV starts to regress and CNV completely disappears

approximately one month post laser. Subretinal fibrosis at the lesion sites can also be observed in this model and this continues to increase through day 35 post laser (Ishikawa et al. 2016). This fibrotic aspect of CNV can be used to study subretinal fibrosis, and the progression of which is often considered a contributing factor to the lack of responsiveness to anti-VEGF therapies.

Although the laser-induced CNV model is frequently used as a neovascular AMD model (Lambert et al. 2013), as noted above, it is an acute injury model and does not recapitulate the aging aspect of AMD. Nor does it mimic macula-specific aspects of AMD, as rodents do not have a macula. In small-rodent eyes, considerable variations in lesion size, even among several lesions within the same eye, are likely due to variations among laser applications. Therefore, sufficient sample size and rigorous application of preestablished inclusion and exclusion criteria are necessary to obtain reliable results (Gong et al. 2015, Poor et al. 2014). Nevertheless, the laser-induced CNV model, together with OIR models (Aiello et al. 1995, Robinson et al. 1996), has been used to evaluate the therapeutic value of anti-VEGF therapies (Krzystolik et al. 2002). In addition to laser-induced CNV, subretinal injection of matrigel is used to induce CNV in rats and mice (Li et al. 2011). This method can be more convenient for researchers without access to laser photocoagulation.

3.4. Genetic Models for Pathological Retinal Vessel Growth

The advantage of genetic manipulation in mice makes them the most commonly used animal model in laboratories for studying retinal vessel growth. Genetic models include transgenic mouse models with intraretinal angiogenesis, subretinal angiogenesis, and deficient intraretinal angiogenesis.

RAP is a distinct form of neovascular AMD. Animal models of RAP exhibit intraretinal, subretinal, and choroidal angiogenic features that mimic the vascular changes seen in human patients. Loss of very-low-density lipoprotein receptor (*Vldh*²) and rhodopsin-driven *Vegf* mutations result in RAP-like lesions (Heckenlively et al. 2003, Hu et al. 2008, Joyal et al. 2016, Okamoto et al. 1997, Sun et al. 2017, Tobe et al. 1998).

Mice lacking *Vldlr* were discovered as a unique spontaneous model of RAP and CNV, both of which are features of neovascular AMD (Heckenlively et al. 2003). Genetic variations such as single-nucleotide polymorphisms in VLDLR are associated with AMD in human patients (Haines et al. 2006). In 1995, Herz and colleagues (Frykman et al. 1995) generated *Vldlr* germ-line knockout (*Vldhr^{-/-}*) mice by targeted disruption of exon 5 of the *Vldlr* gene on chromosome 19. These mice were first characterized as a model of subretinal neovascularization and choroidal anastomosis (Heckenlively et al. 2003). The abnormal neovessels in *Vldlr^{-/-}* retinas sprout from the deep vascular layer in the outer plexiform layer toward normally avascular photoreceptors and invade the subretinal space as early as postnatal day 15 and continue growth through postnatal day 20 (Heckenlively et al. 2003). Subretinal hemorrhages and choroidal anastomosis are evident in *Vldlr^{-/-}* eyes by about 2 months of age (Heckenlively et al. 2003). Moreover, *Vldlr^{-/-}* mice not only recapitulate the pathological vascular features of human RAP, but also exhibit secondary photoreceptor degeneration and subretinal fibrosis (Hu et al. 2008).

VEGF-overexpressing transgenic mice were developed in the late 1990s by overexpressing VEGF in photoreceptor cells by incorporating a full-length complementary DNA of human *VEGF* under the bovine *rhodopsin* promoter (Okamoto et al. 1997, Tobe et al. 1998). This *rhodopsin/VEGF* transgenic mouse model (*rho/VEGF* mouse) shows focal areas of intraretinal and subretinal neovascularization that correspond to areas of vascular leakage on fluorescein angiograms (Okamoto et al. 1997). In adult *rho/VEGF* retinas, blood vessels extend from the inner nuclear layer toward the retinal pigment epithelium to form a large plexus of blood vessels in the subretinal space. This model presents features similar to those of RAP and induces intraretinal

and subretinal neovascularization rather than CNV. This model is useful for investigating early VEGF-induced changes that occur in the retina that ultimately lead to neovascularization and for testing the effects of potential angiogenesis inhibitors (Liu et al. 2017).

The neoretinal vascularization 2 (NRV2, or JR5558) mouse harbors currently unknown genetic mutations (Hasegawa et al. 2014) and has spontaneous intraretinal and subretinal neovascularization (Won et al. 2011). The mouse strain NRV2 was discovered through the Jackson Laboratory Eye Mutant Screening Program (Won et al. 2011). Retinal neovascularization in JR5558 mice originates from the retinal vascular plexus and grows toward the subretinal space, forming neovascular structures at the retinal pigment epithelium and Bruch's membrane interface that mimic the early clinical presentation of RAP in humans (Liu et al. 2017). This genetic model of intraretinal neovascularization provides a useful tool that will help researchers understand the molecular causes of RAP and AMD and evaluate preclinical therapeutics for both diseases (Doyle et al. 2015, Hasegawa et al. 2014, Nagai et al. 2015, Paneghetti & Ng 2016).

There are also genetic mouse models with deficient intraretinal angiogenesis, and they are usually based on mutations in the WNT pathway. Familial exudative vitreoretinopathy and Norrie disease are modeled in *Lrp5* knockout and *Norrin* knockout mice. Mice with genetic deficiency in *Lrp5* (*Lrp5^{-/-}*) and *Norrin* (*Ndp^{y/-}*) show similar ocular vascular abnormalities that resemble human pathologies in familial exudative vitreoretinopathy and Norrie disease, respectively (Wang et al. 2016). *Lrp5^{-/-}* mice have delayed primary retinal angiogenesis, with less dense vascular coverage and fewer branching points in the superficial vascular plexus (J. Chen et al. 2011b, Wang et al. 2016). Similar severe retinal vascular defects are observed in *Ndp^{y/-}* mice, including delayed development of superficial retinal blood vessels, absence of the intermediate and deep layers of retinal vessels flanking the inner nuclear layer, delayed regression of the hyaloid vasculature, and, in some cases, intraocular hemorrhage (Luhmann et al. 2005).

4. KEY FACTORS THAT INFLUENCE RETINAL VASCULAR DEVELOPMENT

4.1. Oxygen in Retinal Vascular Development

Oxygen plays a major role in retinal vascular development. In ROP and DR, an initial lack of vessels or loss of retinal vessels resulting in retinal ischemia precedes pathological vessel growth, supporting the idea of hypoxia as a critical stimulator of new blood vessel growth (Gariano & Gardner 2005). During development, as retinal neurons and glial cells differentiate and mature and their metabolic demands increase, physiological hypoxia leads to the development of new vessels from the center toward the periphery of the retina (Chan-Ling et al. 1995). After new vessels have formed, bringing oxygen and alleviating hypoxia, vascular growth continues radially toward the peripheral retina (Liu et al. 2017).

4.2. Angiogenic Regulators in Retinal Vascular Development

The effect of oxygen on vascular growth is mediated in large part by VEGF, a hypoxia-induced growth factor, which is expressed in the developing retina in a pattern that coincides with retinal blood vessel development (Pierce et al. 1995, 1996; Stone et al. 1995). VEGF and other hypoxia-regulated growth factors are controlled by HIF (Wang & Semenza 1993). A gradient of VEGF precedes the radial growth of the superficial vascular plexus in the retina, and it is postulated that the relative hypoxia of the deeper retinal layers during development also results in a VEGF gradient that favors sprouting from the superficial layer downward, resulting in the formation of

first the deep layer and then the intermediate layer of capillary networks (Pierce et al. 1995, Stone et al. 1995). In addition to VEGF, erythropoietin is a hypoxia-induced growth factor that plays an important role during retinal blood vessel homeostasis (Chen et al. 2008). Erythropoietin also has potent neuroprotective and proangiogenic effects (Caprara & Grimm 2012) and was suggested to be an important factor in the formation of the intermediate plexus of retinal vessels in an HIF-dependent manner (Caprara et al. 2011). These oxygen-regulated hypoxia-responsive factors are therefore considered some of the master regulators of both normal retinal vessel development and proliferative retinopathies (Aiello et al. 1995, Smith et al. 1997, Stone et al. 1995). Besides these hypoxia-regulated growth factors, other non-oxygen-regulated growth factors important for retinal vascular development, partly through modulation of the VEGF response, include the Tie1–Tie2 receptors (Sato et al. 1995), the Tie2 ligand angiopoietin 2 (Hackett et al. 2002), and insulin-like growth factor-1 (IGF-1). IGF-1 is required for maximum VEGF activation of vascular endothelial cell proliferation and survival pathways (Smith et al. 1997, 1999). In premature infants, low IGF-1 levels are associated with an increased risk of ROP (Hellström et al. 2001, 2002, 2003; Löfqvist et al. 2006; Smith 2005), characterized by slowed, deficient development of retinal vessels.

4.3. Neurovascular Crosstalk in Retinal Vascular Development

Neurovascular interactions are important in the maintenance of the nervous system, and defects in this relationship can lead to disease. The retina, which is part of the central nervous system, requires a steady supply of nutrients and oxygen to ensure appropriate neuronal function and sensory transmission. Consequently, neural and vascular systems must be adequately paired so that oxygen and nutrient needs are matched with vascular supply. It has become clear that neurons play an important role in instigating, promoting, and steering angiogenesis within nervous tissue and specifically in the retina (Edwards et al. 2012, Fukushima et al. 2011, Sapieha et al. 2008, Sun et al. 2015a).

4.3.1. Neuronal guidance cues. Endothelial tip cells are enriched in both receptors for angiogenic factors (such as VEGFR) and other receptors that were initially described as neuronal guidance cues but are now understood to function as vascular guidance cues as well as coordinate neural demands with vascular growth (Huber et al. 2003). Tip cells probe the tissue environment through their projecting filopodia and guide the nascent vessel toward gradients of growth factors or guidance cues to its appropriate destination. These neuronal guidance receptors include neuropilins and plexins (for semaphorins), Unc5b (uncoordinated-5 homolog b), neogenin and DCC (deleted in colorectal carcinoma) (for netrins), Eph receptors (for ephrins), and roundabouts (for slits) (Wilson et al. 2006). Tip cells respond to a given cue by advancing, stalling, turning, or retracting depending on the overall intracellular environment of the tip cells (Larrivée et al. 2009). The role of neuronal guidance cues in vascular growth has been comprehensively reviewed elsewhere (De Smet et al. 2009, Gelfand et al. 2009, Larrivée et al. 2009).

4.3.2. Influence of retinal ganglion cells on retinal vascular growth. Among retinal neuron cell types, retinal ganglion cells (RGCs) are the most anatomically coupled with the superficial retinal vascular plexus. A role for RGCs in vascular development was established in mouse models of genetic ablation of RGCs (Edwards et al. 2012, Sapieha et al. 2008). For example, in transgenic mice that express a toxin in newly formed RGCs to eliminate RGCs as they form (Mu et al. 2005), astrocytic networks remain largely intact, yet these mice are completely devoid of a retinal vascular plexus (Sapieha et al. 2008). Similarly, *Math* $5^{-/-}$ mice, which lack 95% of RGCs, do not form a functional retinal vascular layer (Edwards et al. 2012). However, whether astrocytes or retinal neurons such as RGCs are primary drivers of developmental retinal vascular growth is under

debate, as evidence has been provided for both. One possible explanation for the contribution of each cell population during retinal vascular plexus formation is that astrocytes contribute to trophic support of vessels by providing a template for growth (Fruttiger et al. 1996, Uemura et al. 2006) and retinal neurons drive vascular growth to ensure their own metabolic support (Sapieha 2012). In addition, cell-specific depletion of VEGF, HIF-1 α , or HIF-2 α in astrocytes has no effect on developmental retinal vascularization (Weidemann et al. 2010), whereas neuron-specific knockout of HIF-1 α disturbs retinal vascular development (Caprara et al. 2011), suggesting that a neuronal cell population such as RGCs, rather than astrocytes, provides angiogenic factors during retinal development.

In addition, neurovascular crosstalk governs the progression of retinopathy via SOCS3 (suppressor of cytokine signaling 3) (Sun et al. 2015a). Specifically, SOCS3 deficiency in neuronal and glial cells promotes pathological retinal angiogenesis in retinopathy via titrating VEGF signaling. A novel way to suppress retinal angiogenesis may be by inhibiting through SOC3 only the excess VEGF, which is the major cause of proliferative retinopathy. This method may avoid the problems in current treatments that inhibit all VEGF, which is also important for normal vessel growth and maintenance (**Figure 2**).

4.3.3. Influence of photoreceptors on retinal vascular growth: inflammatory signals in photoreceptors. In addition to RGCs, photoreceptors may play a significant role in determining vascular growth. Photoreceptors are the most abundant neuronal cell population in the retina and have the highest oxygen demand (Wangsa-Wirawan & Linsenmeier 2003). Photoreceptors (rods and cones) have an important role in the pathogenesis of ROP (Akula et al. 2007, Hansen et al. 2017). Previous studies of the role of photoreceptors in ROP focused mainly on photoreceptor



Figure 2

Neuronal and glial cell regulation of pathological neovascularization in the eye via STAT3 and SOCS3 signaling pathways. SOCS3 deficiency in neuronal or glial cells promotes pathological retinal angiogenesis in retinopathy via titrating VEGF signaling controlled by activated transcription factor STAT3. A red arrow indicates upregulation, and a green arrow indicates downregulation. Abbreviations: IL-6, interleukin-6; JAK, Janus kinase; SOCS3, suppressor of cytokine signaling 3; STAT3, Signal transducer and activator of transcription 3; VEGF, vascular endothelial growth factor. Figure adapted from Sun & Smith (2015), with permission from *Atlas of Science*.

function during ROP development. Cone photoreceptors develop early before birth with approximately the same birthdate as the other early-born neurons such as RGCs and horizontal cells. Rod photoreceptors develop later than cones (Cepko 2015). The onset of neovascularization in ROP is coincident with the development of the rod outer segment (Fulton et al. 1999). Clinical studies found that ROP has less effect on cone photoreceptors than on rod photoreceptors (Hansen & Fulton 2005, Fulton et al. 2008). Photoresponses suggest that cone photoreceptors are more resistant to the ROP disease process. The similar shape of the b-wave stimulus-response functions in preterm infants and controls implies that ROP does not alter the balance of ON and OFF signals in the cone photoreceptor pathway, suggesting that ROP pathology likely originates from rods instead of cones.

There is evidence that photoreceptors control vascular development. The onset of neovascularization in ROP is found at approximately 32 weeks of gestational age (Hansen et al. 2017) no matter the gestational age at birth, which is coincident with rod maturation and outer segment development (Fulton et al. 1999). In addition, patients who have proliferative DR and then develop retinitis pigmentosa with progressive photoreceptor loss show considerably less pathological retinal angiogenesis than diabetic patients with healthy photoreceptors (Sternberg et al. 1984). Similarly, mice with genetically ablated photoreceptors ($Pdeb^{rd1}$ mutant mice) fail to develop retinal neovascularization in the mouse model of OIR (Lahdenranta et al. 2001).

Inflammatory mediators are involved in retinopathy (M. Chen et al. 2011, Hong et al. 2014, Lee & Dammann 2012). Photoreceptors produce soluble inflammatory factors, which in turn can induce inflammatory changes in nearby cells in diabetes (Tonade et al. 2016), suggesting that photoreceptors might play a role in diabetes-induced degeneration of retinal capillaries. Although inflammation is often thought to come from infiltrating inflammatory cells, including macrophages and neutrophils, photoreceptors also signal for blood vessel growth through inflammatory proteins (Huang et al. 2011, Sun et al. 2017, Tonade et al. 2016). In photoreceptors, transcription factor c-Fos, an immediate early gene and pro-oncogene, acting as a master regulator of many inflammatory signals in a retinal angiogenesis model, *Vldlr*^{-/-} mice (Sun et al. 2017). Inflammatory signaling pathways such as c-Fos can be used by stressed photoreceptors to convey a need for blood vessels and are potential upstream targets to control the development of neovascularization (**Figure 3**).



Figure 3

Photoreceptor regulation of pathological neovascularization in the eye via control of inflammatory signals in photoreceptors. Targeting inflammatory regulator c-Fos in photoreceptors to control inflammatory proteins may provide a novel approach to suppress pathological ocular neovascularization, which leads to blindness. Abbreviation: RPE, retinal pigment epithelium.

4.3.4. Influence of lipid and glucose metabolism on retinal angiogenesis. Energy metabolism needs of retinal neurons may modulate vascular network development (Joyal et al. 2018). In retinal neurons, the cellular signaling events triggered by energy metabolites such as lactate (Ahmed et al. 2010), α -ketoglutarate, and succinate (He et al. 2004) have been proposed as contributors to retinal vascularization (Joyal et al. 2016, Sapieha et al. 2008). We find that glucose metabolism of photoreceptors controls normal retinal vascular development through adiponectin (Fu et al. 2018).

Importantly we have shown that lipids (as well as glucose) are used as fuel in photoreceptors (Joyal et al. 2016). Dyslipidemia is associated with AMD (Haines et al. 2006, Lim et al. 2012). Poor lipid uptake in *Vldlr^{-/-}* mice (Goudriaan et al. 2004) with AMD-like RAP and CNV results in high levels of circulating triglycerides and fatty acids (Goudriaan et al. 2004). *Vldlr* is expressed in tissues with a high metabolic rate, including photoreceptors (Dorrell et al. 2009, Joyal et al. 2016), and facilitates the uptake of triglyceride-derived fatty acids (Lopaschuk et al. 2010). High levels of circulating lipids in *Vldlr^{-/-}* mouse retinas signal through GPR40 to suppress expression of the glucose transporter Glut1. Because loss of *Vldlr* also results in poor lipid uptake in photoreceptors, GPR40-mediated suppression of Glut1 and glucose entry into photoreceptors in *Vldlr^{-/-}* mice result in a dual (lipid and glucose) fuel shortage and a reduction in the levels of the tricarboxylic acid cycle intermediate α -ketoglutarate. Low levels of α -ketoglutarate promote the stabilization of HIF1 α and the secretion of *Vegfa* by starved *Vldlr^{-/-}* photoreceptors, leading to neovascularization (Joyal et al. 2016). Dysregulated lipid and glucose photoreceptor energy metabolism might be a driving force in neovascular AMD and other retinal diseases.

The retina is rich in lipids derived from essential omega-3 and omega-6 long-chain polyunsaturated fatty acids, which are critical for many retinal functions. However, they are unlikely to be used as fuel because they are essential fatty acids that must be obtained through diet, as humans lack key enzymes to synthesize α -linolenic acid (omega-3) and linoleic acid (omega-6) (SanGiovanni & Chew 2005). These essential dietary lipids and their metabolites are structural constituents of membranes and can regulate retinal neovascularization and CNV (Connor et al. 2007, Fu et al. 2015, Gong et al. 2016, Gong et al. 2017, Sapieha et al. 2011, Stahl et al. 2010b).

Development of pathological neovascularization in vascular eye diseases is linked to dysregulation of both lipid metabolism (Chew et al. 2010, Keech et al. 2007) and altered inflammation and macrophage function or polarization (Ferguson & Apte 2008). Lipid-induced alteration in inflammatory response is mediated in part by macrophages, which are phenotypically plastic with different polarization states (M1- and M2-like), playing critical roles in regulating ocular angiogenesis during development and in pathologies (Combadière et al. 2007). ROR α (retinoic-acidreceptor-related orphan receptor alpha), a lipid-sensing nuclear receptor and transcription factor (Solt & Burris 2012) that regulates lipid homeostasis and inflammatory cell differentiation and cytokine production (Jetten 2009), was reported to control pathological angiogenesis through direct transcriptional control of Socs3, suggesting the role of lipid-metabolism-driven inflammation in ocular angiogenesis (Sun et al. 2015b) (**Figure 4**).

4.4. Influence of Immune Response and Inflammation on Retinal Vessel Development

Blood vessels develop accompanied by microglia, the primary resident immune cells of the retina. Microglia are monocyte-derived tissue macrophages of the central nervous system. Retinal microglia populate the mammalian retina before it becomes vascularized (Dejda et al. 2014) and are rapidly activated after an inflammatory insult (Santos et al. 2008). There is evidence that microglia associate with nascent vessels to modulate normal angiogenesis (Kubota et al. 2009, Stefater et al. 2011).



Figure 4

Regulation of lipid-metabolism-driven inflammation of pathological neovascularization in the eye via the ROR α -SOCS3 pathway. ROR α senses lipid metabolites and regulates target gene expression through binding to RORE elements in their promoter region. ROR α regulates pathological neovascularization (NV) in the eye via control of tissue inflammation and macrophage polarization. Additional abbreviations: ROR α , retinoic-acid-receptor-related orphan receptor alpha; RORE, ROR α responsive element; SOCS3, suppressor of cytokine signaling 3.

The immune response and inflammation are also associated with pathological angiogenesis. There may be an increased risk for ROP in the premature neonate because of an immature immune system (M. Chen et al. 2011). There are higher rates of ROP in infants born to mothers having evidence of chorioamnionitis than in those born to mothers without signs of inflammation (Moscuzza et al. 2011). On the basis of current knowledge about ROP etiology and pathogenesis, it seems likely that O₂/VEGF-related ROP risk is modified by systemic inflammation. The role of anti-inflammatory drugs in preventing severe ROP in the presence of systemic inflammation, and possibly even without systemic inflammation, should be investigated (Lee & Dammann 2012). Recent work suggests a strong link between the SOCS3 signaling pathway and ROP (Stahl et al. 2012; Sun et al. 2015a,b). SOCS proteins are key physiological regulators of both innate and adaptive immunity and inflammatory responses. SOCS3 positively and negatively regulates macrophage and dendritic cell activation, respectively, and is essential for T cell development and differentiation. Evidence is also emerging of the involvement of SOCS proteins in diseases of the immune system (Yoshimura et al. 2007) and the involvement of the immune system in ocular neovascularization.

In addition, inflammatory pathways, including Toll-like receptors (Imai et al. 2008), and the transcriptional activator nuclear factor kappa B (de Oliveira-Marques et al. 2007) may also be modulated by oxidative stress (Vento et al. 2009). Increases in oxidized lipids and hydrogen peroxide in preterm infants are likely because of supplemental oxygen at birth. Activation of inflammatory pathways by oxygen by-products can potentiate inflammatory responses and promote further endothelial injury, resulting in generalized systemic inflammation (Volpe 2008). Prolonged exposure to supplemental oxygen and systemic inflammation [plasma interleukin-6 (IL-6) and tumor necrosis factor-alpha (TNF α)] are strongly associated with severe ROP (Silveira et al. 2011). Inflammatory processes might interfere with normal retinal vascularization in preterm retinas (Dammann 2010). Loss of TNF α in the OIR mouse model appears to be protective (Gardiner et al. 2005). Omega-3-polyunsaturated fatty acids decrease the size of the avascular area, and this protective effect may be mediated in part via suppression of TNF α (Connor et al. 2007). Markers of inflammation also appear to be associated with ROP in human studies (Lavoie et al. 2010). Over the first three postnatal weeks, the systemic levels of eight cytokines and chemokines were different

in infants with ROP compared with controls, and co-occurrence of immaturity, proinflammatory cytokines, and neonatal systemic inflammation is associated with increased risk for ROP. Besides the OIR model, the expression levels of inflammatory cytokine (IL-6 and TNF α) modulated by key inflammatory factor c-Fos in photoreceptors are increased in the retinas of the *Vldlr*^{-/-} mouse model (Sun et al. 2017).

5. CONCLUDING REMARKS

Visual function in the eye is supported by retinal and choroidal vessels. Pathologies in either vascular bed can cause significant visual impairment: Retinal neovascularization occurs in proliferative retinopathy (including DR and ROP), and CNV is seen in AMD. DR, ROP, and neovascular AMD are associated with rapid deterioration of visual function (Lim et al. 2012, Zhang et al. 2012). Although most patients with DR or ROP do not advance to the proliferative stage, once this progression occurs, it can cause profound vision loss. Similarly, although CNV occurs in only 10% of AMD patients, it accounts for up to 90% of vision loss associated with AMD (Apte et al. 2006). Pathological neovascularization continues to be the leading cause of vision loss, and advances in understanding the underlying mechanisms of angiogenesis have led to the development of therapeutics effective in decreasing visual morbidity. This article summarizes the current knowledge of the cellular and molecular processes that govern both physiological and pathological retinal vascular development, which is dependent on the interactions among retinal cell populations, including neurons, glia, immune cells, and vascular endothelial cells. We also review the animal models currently used for studying retinal vascular development and the key factors that influence vascular growth. We anticipate that the identification of key regulators that globally influence pathological neovascularization will lead to the development of new targeted therapies for preventing and treating these debilitating neurovascular eye diseases.

SUMMARY POINTS

- 1. The retina is supplied by two blood sources: the central retinal artery for the inner retina and the choriocapillaris for the retinal pigment epithelium and outer retina. Pathological neovascularization, which threatens vision, can arise from either vascular bed.
- 2. Both oxygen and fuel deficits in neurons play a major role in pathological neovascularization. Neuronal and vascular interactions are important in modulating angiogenesis in the retina.
- 3. Immune response and inflammation influence retinal vessel development.
- 4. Genetic and injury mouse models of intraretinal and subretinal angiogenesis, deficient intraretinal angiogenesis, and CNV are available.

FUTURE ISSUES

- 1. Photoreceptor energy demands may play a significant role in determining vascular growth.
- 2. Lipid metabolism in photoreceptors is an important aspect of energy metabolism modulating the vascular network.

3. Pathological neovascularization driven by inflammatory signals may originate from photoreceptors and may be linked to photoreceptor metabolism.

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