



# Role of Wnt inhibitor *Apcdd1* in retinal angiogenesis and barrier formation

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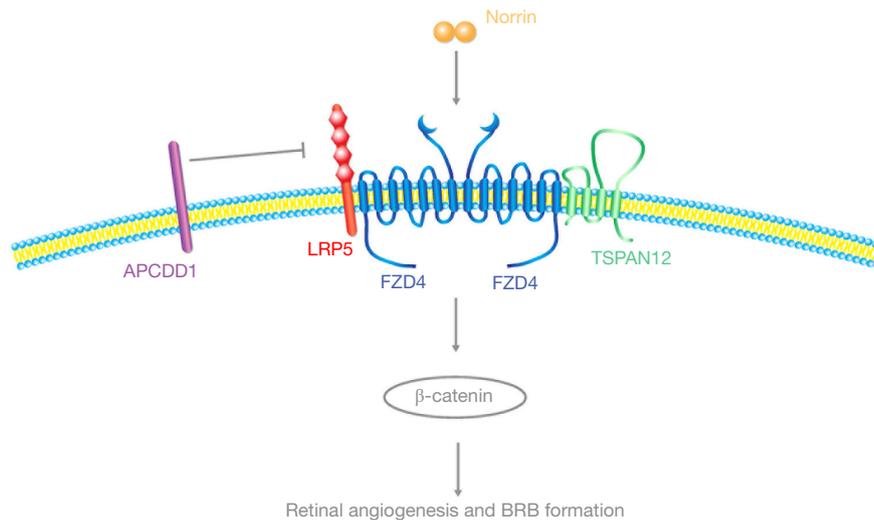
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Angiogenesis and blood-retinal barrier (BRB) formation/maintenance are required for normal function of the eye. These processes are controlled by canonical Wnt ( $\beta$ -catenin-dependent) signaling in retinal endothelial cells (ECs) (1). In the retinal vasculature, canonical Wnt signaling is mediated by the ligand NDP (Norrie disease protein) and a receptor complex composed of FZD4 (frizzled class receptor 4), LRP5 (low-density lipoprotein receptor-related protein 5), and the tetraspanin family member TSPAN12 (2-6). NDP, FZD4, LRP5 and TSPAN12 mutant mice had been reported by several groups to exhibit similar vascular defects characterized by a lack of intraretinal blood vessels and BRB defects (2,7). Additionally, in mutant mice where NDP/FZD4 endo-lysosomal trafficking is inhibited specifically in ECs, similar morphological defects are observed in the retinal vasculature (8). These defects in mice resemble those observed in human patients with FEVR (familial exudative vitreoretinopathy) disease. In these mutant mice discussed, canonical Wnt signaling is downregulated, yet, canonical Wnt signaling upregulation has not been well studied.

Recently, a paper published in *Neuron* by Mazzoni *et al.* presented a mouse model lacking the canonical Wnt inhibitor *Apcdd1* (adenomatosis polyposis coli down-regulated 1) and characterized its role in retinal angiogenesis as well as barrier formation at different stages during early development (Figure 1) (9). In this paper, Mazzoni *et al.* first examined the expression pattern of *Apcdd1* in the retina by RNA *in situ* hybridization from P6-P17 in mice. *Apcdd1* is mainly expressed in retinal ECs but not in pericytes or other neuronal markers, indicating a potential role of *Apcdd1* in retinal angiogenesis. Mazzoni *et al.* comprehensively

examined retinal blood vessels from *Apcdd1* mutant mice (12). Although there are no differences detected between control and *Apcdd1* mutants at P5, knockout mice have significantly increased retinal blood vessel density from P10–P12. This vascular overgrowth phenotype contrasts canonical Wnt signaling loss-of-function phenotype. Additionally, *Apcdd1* has been reported to be downregulated in retinas with mutant *Norrin* or *Lrp5* (13). The increased vessel density observed in *Apcdd1* mutants may be due to upregulated canonical Wnt signaling. Interestingly, Mazzoni *et al.* identified upregulated *Sox17* (Wnt/ $\beta$ -catenin target gene), increased pLRP6 and more active  $\beta$ -catenin at P10 in *Apcdd1* mutant mice (12). These results support *Apcdd1* acts as a negative regulator for canonical Wnt signaling in retinal angiogenesis. However, this overgrowth phenotype is only restricted in superficial vascular layer and eventually resolves after P14, suggesting that *Apcdd1* mutants have a milder defect than *Norrin*, FZD4, LRP5, and TSPAN12 mutants. Their findings provide evidence that canonical Wnt signaling needs to be precisely controlled to ensure normal vascular development in the retina.

There are two reasons for retinal vascular overgrowth: retinal EC over proliferation or delayed vessel pruning. Canonical Wnt signaling has been reported to be involved in the both processes (14,15). For example, less EC proliferation and more vessel regression have been reported in *Norrin* mutant mice (15). However, Mazzoni *et al.* performed EdU assay on *Apcdd1* mutant mice from P8–P12 and did not find any evidence of over proliferation (9). On the other hand, some of newly formed ECs did not go



**Figure 1** Norrin/ $\beta$ -catenin signaling in retinal angiogenesis and BRB formation with *Apccdd1* involved in as negative regulator. Mazzoni *et al.* showed *Apccdd1* inhibits  $\beta$ -catenin signaling (9). *Apccdd1* was only reported previously to interact with LRP5 (10) but the question of how exactly *Apccdd1* regulates  $\beta$ -catenin remains complicated. Modified from Lai *et al.* (11).

through the pruning process in *Apccdd1* mutant retinas, which result hyper-vasculature. These results suggest *Apccdd1* negatively controls canonical Wnt signaling but only in the process of pruning.

Canonical Wnt signaling also controls development and maintenance of the BRB (3,5,7). It is possible that BRB formation is affected in *Apccdd1* mutant mice. Mazzoni *et al.* used a low-molecular weight biocytin tracer to test paracellular BRB permeability in *Apccdd1* mutant mice (9). These results demonstrated that *Apccdd1* mutant mice from P10–P14 formed stronger paracellular BRB. Within EC junctions, the occludin mRNA level was not altered but its protein level increased in *Apccdd1* mutants, indicating *Apccdd1* affects BRB formation by regulating occludin protein stability. Consistent with the vessel overgrowth phenotype, stronger paracellular BRB permeability also disappeared at later stages. The BRB is also characterized by a low transcellular trafficking rate. Mazzoni *et al.* tested this trafficking rate in *Apccdd1* mutant mice by intravenous injection of albumin, a transcellular tracer. However, there was no significant change for transcellular trafficking in the retinal ECs of *Apccdd1* mutant mice. Interestingly, isolated brain ECs showed similar decreased paracellular permeability as in the BRB but increased transcellular permeability, which was inhibited by adding Wnt3a.

Mazzoni *et al.* also generated a mouse model for *Apccdd1* gain-of-function specifically in ECs (9). These endothelial

*Apccdd1* overexpression mice display opposite phenotypes to *Apccdd1* knockout mice, including decreased retinal blood vessel density and higher paracellular BRB permeability. All these data support *Apccdd1* playing an important role in regulating retinal angiogenesis and BRB formation.

There are still several questions remaining for the role of *Apccdd1*. First, although *Apccdd1* mutant mice show phenotypes at early stage in retinas, they all revert back to normal at later time points. Is there a parallel pathway compensating for *Apccdd1*? Are there any phenotypes in ECs from other organs? Second, *Apccdd1* has been previously reported to interact with Wnt3a and Lrp5 (10). Does it also interact with Norrin or other molecules in canonical Wnt signaling? Third, it is reported that *Apccdd1* mRNA level is decreased in Norrin and Lrp5 mutant retinas (13). Is *Apccdd1* a direct target for Norrin/Lrp5 or due to a complicated feedback loop? Forth, does inhibiting *Apccdd1* rescue defects from Norrin/FZD4/LRP5/Tspan12 mutant mice? Addressing these questions will not only help further our understanding of how canonical Wnt signaling is regulated but also provide new therapeutic avenues for treating diseases that result from loss of canonical Wnt signaling, such as FEVR disease.

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