Comparison	Figure	Primary Statistical	Post Test
		Test	
Retinal pericyte coverage at P5	2K-L	SRH (Factors:	MWU Tests +
& P10		genotype and region)	Bonferroni Correction
No. of Avascular regions/ mm ²	5E	ANOVA (Factors:	TukeyHSD
		genotype and region)	
No. of Branchpoints/ mm ²	5F	ANOVA (Factors:	TukeyHSD
		genotype and region)	
RGC counts	6M	Kruskall-Wallis test	MWU Tests +
			Bonferroni Correction
Cerebral cortex and retinal	Supp.	SRH (Factors:	MWU Tests
pericyte coverage at P5 by	Fig. S3	genotype and region)	
Desmin staining			

Supplementary Table S3: Statistical Analysis Undertaken

The normality of the data analysed was assessed by Shapiro-Wilk test and by visual inspection and subsequently the appropriate parametric or non-parametric tests were performed.

SRH: two-way Scheirer-Ray-Hare (SRH) extension of the Kruskal-Wallis rank sum test,

MWU Tests: pairwise Mann-Whitney U-tests. ANOVA: two-way ANalysis Of VAriance,

TukeyHSD: pairwise Tukey's Honestly Significant Difference. Prior to analysis the avascular

regions /mm² and branchpoints/ mm² data was root and probit transformed respectively.