



Article

Analysis of mir-9 expression pattern in rat retina during postnatal development

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Table S1. Multiple comparison of miR-9 expression in different time points analyzed by ordinary one-way ANOVA with Tukey's post-hoc test. Upper part represents data obtained from RT-qPCR, while lower section shows miRNA-sequencing data. (* p<0.05, ** p<0.01, *** p<0.001)

		P1	P3	P5	P7	P10	P15	P21
RT-qPCR	P1	*** <0.0001	ns	ns	*	0.0274	ns	ns
	P3		*** <0.0001	*** 0.0003		ns	*	*** <0.0001
	P5				ns	ns	ns	ns
	P7					ns	ns	ns
	P10						ns	** 0.0072
	P15							ns
		P1	P3	P5	P7	P10	P15	P21
miRNA-Seq	P1	ns	ns	ns		*	0.0143	ns
	P3		ns	ns		** 0.0061	ns	ns
	P5			ns		*** 0.0005	ns	ns
	P7					*** 0.0002	*	ns 0.0219
	P10						0.0194	0.0006



Table S2. Multiple comparison of OneCut2 mRNA expression in different time points analyzed by ordinary one-way ANOVA with Tukey's post-hoc test. (* p<0.05, ** p<0.01, *** p<0.001)

	P1	P3	P5	P7	P10	P15	P21
P1	ns	ns		*	**	***	***
P3		ns	ns	0.0497	0.0041	0.0005	<0.0001
P5			ns	ns	ns	0.0008	*
P7				ns	ns	ns	0.0297
P10					ns	ns	
P15						ns	



Table S3. Multiple comparison of synaptotagmin-17 mRNA expression in different time points analyzed by ordinary one-way ANOVA with Tukey's post-hoc test. (* p<0.05, ** p<0.01, *** p<0.001)

	P1	P3	P5	P7	P10	P15	P21
P1	ns	ns	ns	ns	ns	*	
P3		ns	ns	ns	ns	ns	
P5			ns	ns	ns	ns	
P7				ns	ns	ns	
P10					ns	ns	*
P15						0.031	