

Table S3 RGC type- and injury- specificity of lncRNAs and mRNAs.

Transcript type	# Transcripts detected exclusively in ipRGCs	# Transcripts detected exclusively in normal ipRGCs	# Transcripts detected exclusively in injured ipRGCs	# Total transcripts detected in ipRGCs	# Transcripts detected exclusively in ooDSGCs	# Transcripts detected exclusively in normal ooDSGCs	# Transcripts detected exclusively in injured ooDSGCs	# Total transcripts detected in ooDSGCs
lncRNAs	268 (31% of the total)	294 (35% of the total)	188 (22% of the total)	851	564 (49% of the total)	311 (27% of the total)	293 (26% of the total)	1,147
mRNAs	605 (5% of the total)	895 (7% of the total)	526 (4% of the total)	12,229	1,137 (9% of the total)	505 (4% of the total)	855 (7% of the total)	12,761