

Identification and characterization of FGF2-dependent mRNA:miRNA networks during lens fiber cell differentiation

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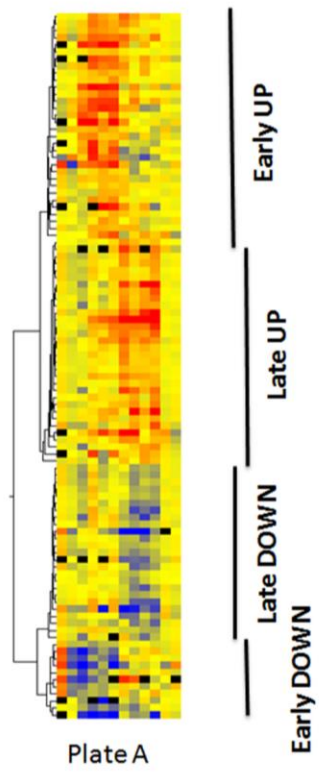
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microRNA profile	Regulated (FC > 1.25; 2 time points)
“Early” UP	50
“Early” DOWN	16
“Late” UP	36
“Late” Down	29

Figure S1 Self organizing maps on duplicate temporal profiles to identify main miRNAs expression profiles.

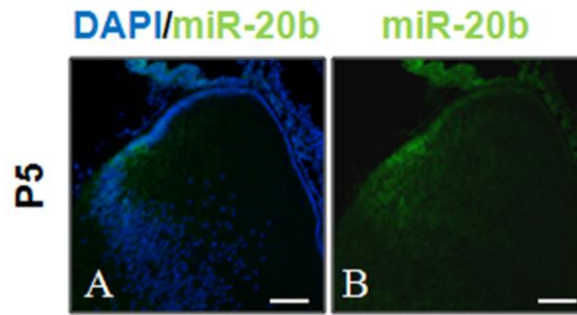


Figure S2 ISH miR-20b. At postnatal day P5, expression of miR-20b (A-B) is detected in the migrating lens cells. Scale bar is 100 μ m.

Files S1-S4 and Table S1

Available for download at <http://www.g3journal.org/lookup/suppl/doi:10.1534/g3.113.008698/-/DC1>

File S1 Two .xls files, inversely correlated genes.

File S2 List of genes identified by GO and KEGG.

File S3 Two .xls files, connectivity and ranking.

File S4 Expression analysis of Bmp2, Bmp4, Bmp7, Id1, Id2, Id3, Fosl1, Fosl2, c-Fos, Ets1, Ets2, Elf1 and Etv1/ER81.

Table S1 Expression data on 131 microRNAs modulated by FGF2 in rat lens explants.