



GATA4: a key factor for liver fibrosis regression

Salazar Martínez, Miguel (1), Arroyo de Alba, Noelia (1), Díaz Contreras, Irene (1) y Rojas González, Anabel (1,*)

(1) Centro Andaluz de Biología Molecular y Medicina Regenerativa (CABIMER), Avda. Américo Vespucio 24. Parque Científico y Tecnológico Cartuja, 41092, Sevilla, España

Motivation

Several studies have shown that the reversion of active HSCs to a quiescent state is required for liver fibrosis regression. Previous work in our lab have identified GATA4 as key factor for the maintenance of HSCs quiescence. Injection of adenovirus overexpressing GATA4 (Ad-GATA4) via the tail vein in CCl₄treated mice have been used as an approach to revert the fibrosis. Using this method, adenoviruses are sequestered by all hepatic cells. As the first objective of this project we will quantify the % of Adenoviruses that reach the HSCs and other liver cells to test whether the liver regression is due to the overexpression of GATA4 in active HSCs.

Transcriptional repression of HIF2a by GATA4 has been uncover in the group as a molecular mechanism by which GATA4 inactivates fibrotic HSCs during liver fibrosis regression. Two conserved GATA sites in the Hif2a gene have been identified. As a second objective we will test whether the repression of Hif2a by GATA4 requires these two GATA sites. Finally, as a third objective, we will develop a vector containing HSCs-specific enhancer from the Gata4 gene (hG2) fused to a luciferase reporter gene. Using this construct (hG2-luc), we will generate a stable cell line to screen FDA-approved drugs that activate Gata4 expression in active (fibrotic) HSCs to induce the quiescence.



Immunofluorescence in liver sections of mice treated with Ad-GFP for

GFP and Desmin (HSCs marker)

or

GFP and HNF1 (hepatocyte marker)

Quantification of double marked cells will be performed using ImageJ software

- pCI-GATA4 or pCI-Ø expression vector, and
- control vector for Renilla luciferase expression



Luciferase activity was measured using a commercial kit

construct. This construct will allow the FDAapproved drugs screening to identify activators of *Gata4* expression in HSCs

Luciferase activity will be measured using a commercial kit



Results and conclusions

- GFP and Desmin (HSCs marker)





pGL4-hG2-TK-Luc construction has been successfully generated to develop a stable cell line. As shown, digestion of the vector obtained with restriction enzymes Nhel/HindIII and Xhol/HindIII released, respectively, a fragment of ~ 1513 bp (hG2-TK) and ~ 63 bp (TK). pGL4-hG2- pGL4-hG2-TK-Luc TK-Luc

- GFP and HNF1 (hepatocyte marker)



Data from preliminary quantification showed that around 50% of HSCs were positive for GFP and Desmin. So that, adenoviruses injection via tail vein in mice efficiently reach HSCs.

As its seen in the graphic, mutation of GATA sites in the *Hif2a* avoid transcriptional repression by GATA4, letting know that GATA4 to transcriptionally repress Hif2a via two identified conserved GATA sites.

The difference between basal groups can be explained by the activation of endogenous GATA4 that exists in the 293 cell line.

The 1,51% of repression observed in HIF2aMUT must be due to a third GATA site non mutagenized in the *Hif2a*, as well as GATA4 is forced to bind to the endogenous GATA sites of 293 cells.



References

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