



(P15) Synthetic RNAs for sorting cells in an *in vitro* model of pancreatic reprogrammed cells

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In order to improve the quality of life of Type 1 Diabetes patients, basic science is focusing on developing Cell Therapies to reconstitute the critical pancreatic beta cell mass to restore glycaemia control. Some strategies involve reprogramming or transdifferentiation from stem cells or differentiated cells to pancreatic beta-like cells, but until now they have to sort many technical impediments before reaching the clinic.

One of the main impediments is the heterogeneous cell population resulting following a reprogramming process, and it represents a high risk if these cells will be transplanted into patients. Trying to avoid this problem, Miki et al., developed a new technique using synthetic RNAs to sort cells, reaching a 99.7% purification efficiency, overcoming the efficiency of some classic methods. This technique depends on the activity of endogenous microRNAs that function as lineage markers. The objective of our work was to apply this technique in an *in vitro* model of pancreatic reprogramming cells to sort microARN-Let7a positive cells. The microRNA-Let7a has been previously associated with insulin-producing cells.

For this purpose, we tested two cell lines, HEK293T as non-reprogrammed cells and MIN6 acting like pancreatic beta cells post-reprogramming. We validated miR-Let7a in order to complement the miR-375 activity as pancreatic beta cell markers; so we synthesized gBlocks for miR-Let7a-BIM-switch (containing the miR-Let7a complementary sequence and encoding the apoptosis inducer protein BIM), miR-Let7a-BFP-switch, miR-375-BFP-switch and we included mRNA-GFP as lipofection control. Constructions were synthesized by MegaScript® SP6 *in vitro* transcription kit. Both cell lines were transfected using Lipofectamine 2000® with different concentrations of miR-Let7a-Bim synthetic switch in order to probe their tolerance to the apoptotic protein, Bim. Apoptotic cells were visualized after 18 hrs by TUNEL assay under a fluorescent microscope. After this time, we designed an *in vitro* model simulating the post-reprogramming environment with a co-culture of HEK293T and MIN6 cells. The synthetic miR-Let7a-Bim was transfected with the miR-375-BFP. After 18 hrs, apoptotic cells were counted under a fluorescent microscope and 85% of HEK293T were dead, so we enriched insulin-positive MIN6 cells to >80% from a heterogeneous population in an *in vitro* model without using any equipment like flow cytometry with a cell sorter.

We confirmed that synthetic RNA switches are a novel tool for purification of living cell types applied in a post-reprogramming environment based on their differential microRNA activity.