



# Effective Splicing Correction of *SMN2* Gene in SMA Cells after Delivery of RNA Interpolyelectrolyte Complexes

## *Mariia Palagina, Marianna Maretina, Anton Kiselev*

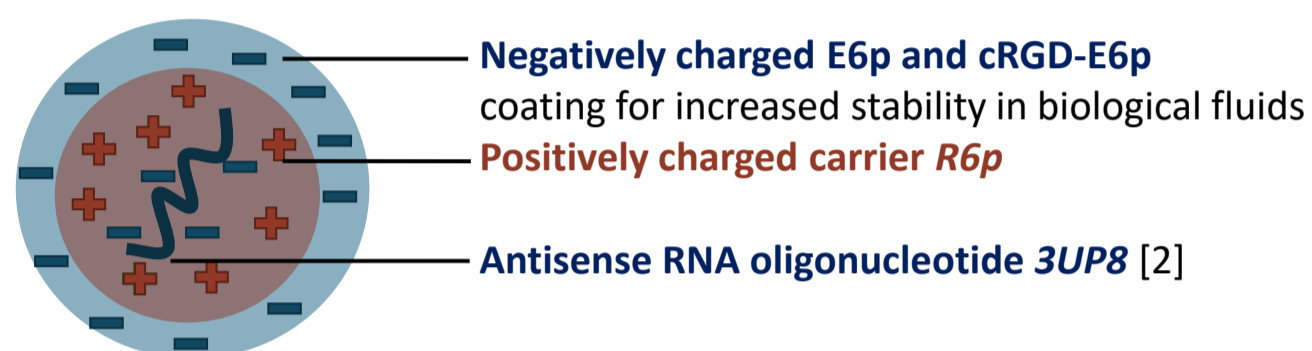
Department of Genomic Medicine Named after V.S. Baranov, D.O. Ott Research Institute of Obstetrics, Gynecology and Reproductology, Mendeleevskaya Line 3, 199034 Saint-Petersburg, Russia  
E-mail: [palagina.marie@gmail.com](mailto:palagina.marie@gmail.com)

## INTRODUCTION & AIM

- Splicing correction of the *SMN2* gene is a key treatment for spinal muscular atrophy.
- Intrathecal administration of antisense RNA oligonucleotides (ASOs) can cause side effects and requires repeated injections.
- The use of non-viral methods for the delivery of antisense oligonucleotides has the potential to facilitate the transition from intrathecal administration to intravenous infusion of these drugs.
- Ternary oligonucleotide–peptide complexes coated by anionic polypeptide are designed to overcome the impossibility of systemic delivery antisense RNA oligonucleotide.

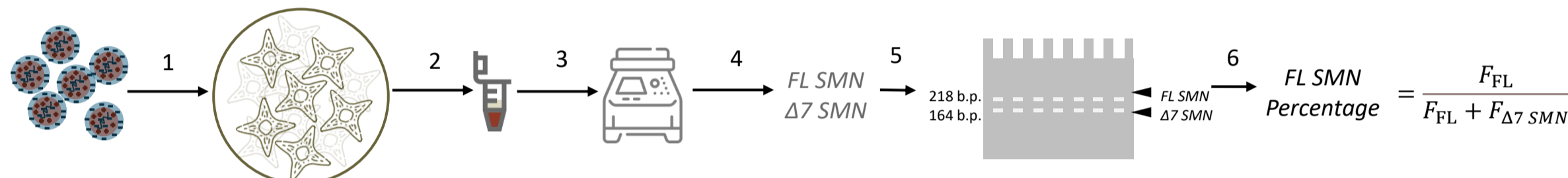
## METHOD

### Ternary oligonucleotide–peptide complexes

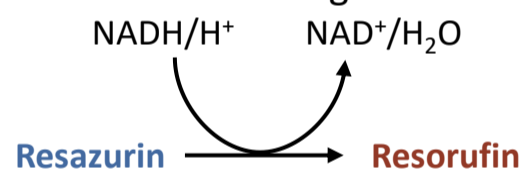


Previously, we developed a **cationic peptide carrier R6p**, and an **anionic peptide coating E6p** and **cRGD-E6p** (ligand-modified to  $\alpha\beta3$  integrins) [1] to deliver ASO 3UP8 [2] to a cellular model of SMA in the presence of serum, which corresponds to physiological conditions. The **RNA/anionic carrier** charge ratios tested in this work: **1/4, 1/8, 1/16, 1/24**.

The study involved transfecting SMA fibroblast cell culture with the serum-stable oligonucleotide–peptide complexes, reverse transcription, semiquantitative PCR, and resazurin assay.



1. Transfection with interpolyelectrolyte complexes of primary fibroblast cell culture obtained from skin biopsy of patients with SMA II [3] in the presence of fetal bovine serum. 2. RNA isolation. 3. Reverse transcription. 4. Semi-quantitative fluorescence PCR with primers for amplification of a cDNA fragment from 6 to 8 exons of *SMN2* [4]. 5. Visualization of PCR by electrophoretic separation of PCR products in 6% polyacrylamide gel. Gel staining with ethidium bromide. Estimation of the luminescence intensity of amplification products using the ImageJ. 6. Calculation of the proportion of full-length *SMN* transcripts (*FL SMN*) to the sum of full-length and truncated *SMN* transcripts ( $\Delta 7$  *SMN*) [5].



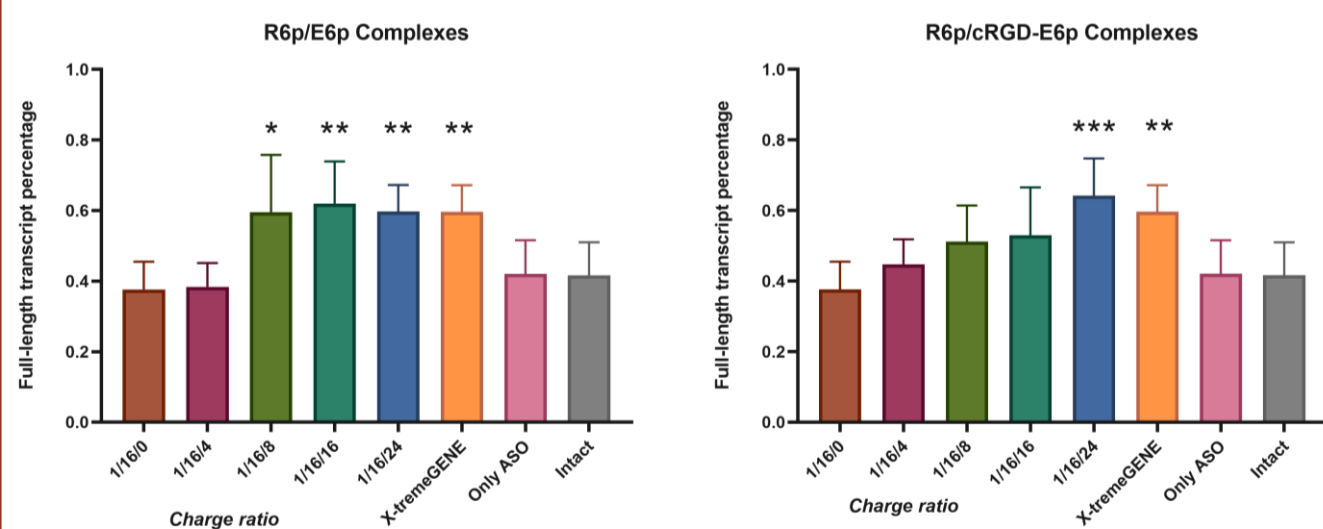
### Resazurin Assay

$$\text{Relative amount of living cells} = \frac{F - F_0}{F_c - F_0}$$

$F_c$  - dye fluorescence intensity in the absence of RNA/carrier complexes,  
 $F_0$  - dye fluorescence intensity in the absence of cells.

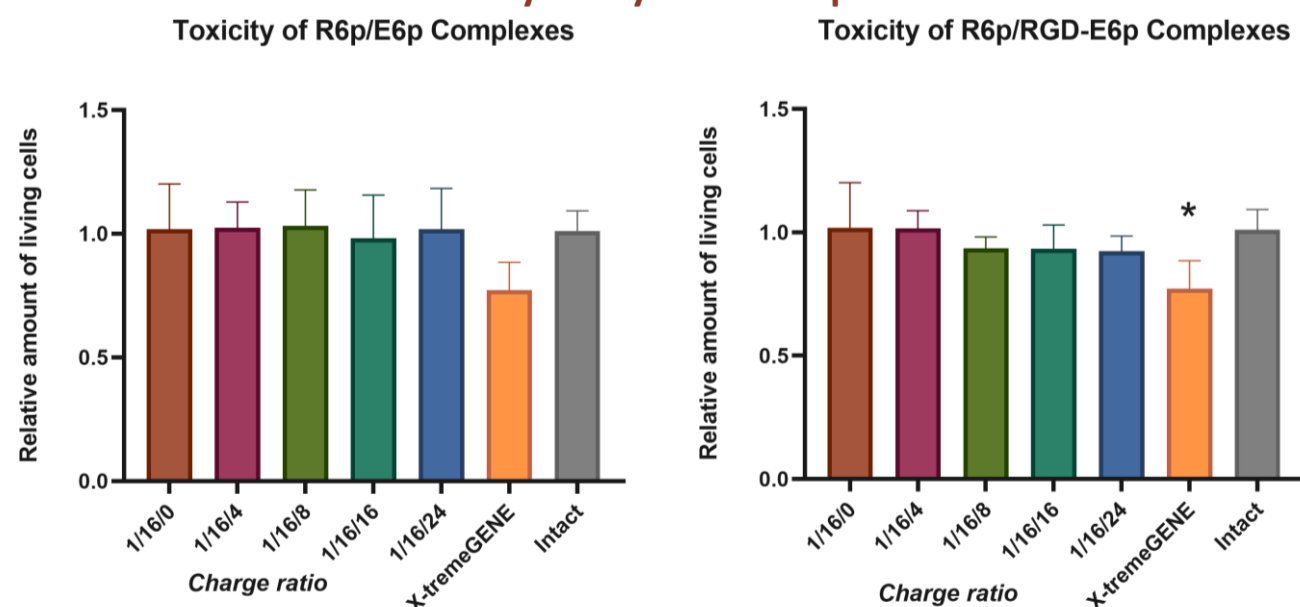
## RESULTS & DISCUSSION

### Evaluation of the Effectiveness of *SMN2* Gene Splicing Correction



The results are presented as a mean  $\pm$  SD  $n = 12$  individual samples from the three independent experiments. \* -  $p < 0.05$ , \*\* -  $p < 0.01$ , \*\*\* -  $p < 0.0001$  compared to intact cells. The level of significance is determined using the Kruskal-Wallis test.

### Toxicity Analysis of Complexes



The results are presented as a mean  $\pm$  SD  $n = 6$  individual samples from three independent experiments. \* -  $p < 0.05$  compared to intact cells. The level of significance is determined using the Kruskal-Wallis test.

Developed system demonstrates **low toxicity** and **effectively facilitates transfection** in cell culture, closely resembling **physiological conditions**.

## REFERENCES

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