

# CORRECTING A MUTATION IN THE GERM LINE BY CRISPR/CAS9 TO OVERCOME TRANSMISSION OF HEREDITARY INFERTILITY IN MICE

**Bekaert B.** (1), Boel A. (1), Thys V. (1), Stamatiadis P. (1), Chuva de Sousa Lopes S. M. C. (1, 2), De Sutter P. (1), Parrington J. (3), Menten B. (4), Stoop D. (1), Coucke P. (4) and Heindryckx B. (1)

(1) Ghent-Fertility And Stem cell Team (G-FaST), Department for Reproductive Medicine, Ghent University Hospital, Corneel Heymanslaan 10, 9000 Ghent, Belgium; (2) Department of Anatomy and Embryology, Leiden University Medical Center, Leiden, the Netherlands; (3) Department of Pharmacology, University of Oxford, Mansfield Road, Oxford OX1 3QT, UK; (4) Center for Medical Genetics Ghent, Ghent University, Department of Biomolecular Medicine, Corneel Heymanslaan 10, 9000 Ghent, Belgium

**Aims:** Failed Fertilization after ICSI can be caused by mutations in the sperm-related oocyte factor phospholipase C (PLC $\zeta$ ) gene. This can be overcome by artificial oocyte activation (AOA). We aimed to overcome mutation transmission by gene correction through CRISPR/Cas9 delivery during ICSI in a knock-out mouse model. **Methods:** We employed a previously described PLC $\zeta$  knock-out mouse model, harboring a homozygous 22 base-pair mutation (Plcz1 $\Delta$ 22bp), causing failed fertilization after ICSI. Two different gRNA molecules were designed to specifically target the mutant allele. Together with the knock-out sperm, the gRNA/Cas9 complex and a repair template, harboring the wild-type Plcz1 sequence and an additional synonymous mutation to track the use of the template, were injected. AOA was performed following ICSI to obtain normal fertilization rates. **Results:** The different alleles obtained after editing were categorized as follows: unedited Plcz1 $\Delta$ 22bp allele, Plcz1 $\Delta$ 22bp allele harboring additional indel mutations, Plcz1 $\Delta$ 22bp allele corrected with a repair template, and Plcz1 $\Delta$ 22bp allele altered/corrected by alternative mechanisms. An efficient correction rate of 45% of the embryos was observed. Our sequencing data suggest that 25% did use the template during repair attempts and 35% did not. In 40% the template was occasionally used. Mosaicism (=multiple alleles per embryo) was observed in 45% of the embryos. For all parameters, no significant ( $p > 0.05$ ) difference was seen between the two gRNA designs and CRISPR/Cas9 editing did not significantly ( $p > 0.05$ ) affect the activation and embryonic developmental rates. Embryos that were capable to develop into blastocysts had a normal karyotype. **Conclusion:** CRISPR/Cas9 can be efficiently used for the correction of a 22 base-pair deletion.