

Gene Editing Strategies to Create Human Derived Erythroid Progenitor Cells with SC Mutation

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INTRODUCTION

- Sickle cell disease (SCD) is caused by homozygous Beta-sickle mutation on both alleles or compound heterozygous with Hemoglobin (Hb)S on one allele and HbC or a Beta-thalassemia mutation on the second allele.
- Hemoglobin sickle-Hemoglobin C disease (HbSC) accounts for 30% of SCD genotypes. HbS results from A→T mutation in the sixth codon of the HbB gene while HbC results from G→A mutation in the same codon of the HbB gene.
- HbSC patients also have substantial morbidity and mortality. Hence, creating HbSC cell models is crucial to understand disease pathophysiology and developing novel therapies.

OBJECTIVE

- The aim of the project is to optimize a CRISPR/Cas9 gene editing strategy in Human Umbilical Cord Blood-Derived Erythroid Progenitor (HUDEP)-S cell line from a sickle patient to create HUDEP-C and HUDEP-SC cell line.

METHODS

- Two different single stranded oligonucleotide templates (ssODN donor A and B) were trialed with at least 2 different concentrations.
- Delivery of ribonucleoprotein (RNP) complex of CRISPR guide RNA (gRNA) and Cas9 with ssODN templates was done through electroporation.
- Transfected cells were cultured and incubated for recovery before harvested for examination.
- Two methods were used for confirming gene editing:
 - 1) PCR with a digestion reaction using a restriction enzyme (HhaI): If editing is successful and homology directed repair (HDR) occurs, two bands (250 and 207 Bp) should be detected during gel electrophoresis.
 - 2) TIDE (Tracking of indels by Decomposition) assay coupled with sanger sequencing to examine if non-homologous end joining (NHEJ) occurs

RESULTS

ssDON Templates- Edits needed to result in HbC highlighted in red

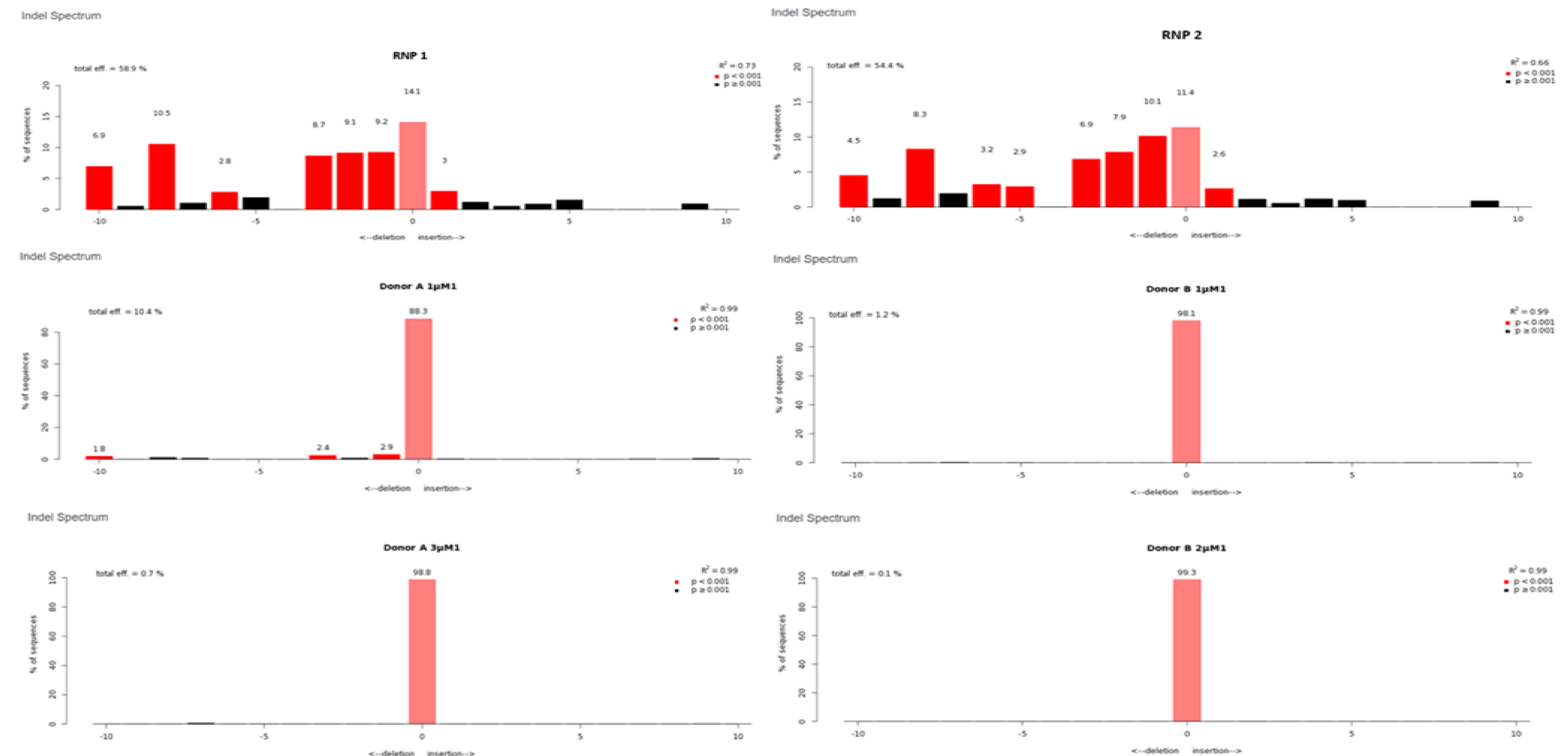
Donor A
TCTGACACA ACTGTGTTCACTAGCAACCTCAAACAGACACCA
TGGTGCATCTGACTCCTaagGAGAAGCTGCGGTTACTGCGC
TGTGGGGCAAGGTGAACGTGGATGAAGTTGGTGGTGA

Donor B
TCAGGGCAGAGCCATCTATTGCTTACATTTGCTTCTGACACA
ACTGTGTTCACTAGCAACCTCAAACAGACACCATGGTGCATCT
GACTCCTaagGAGAAGCTGCGGTTACTGCGCTGTGGGGCAA
GGTGAACGTGGATGAAGTTGGTGGTGAAGCCCTGGGCAGGT

Gel electrophoresis after digestion reaction with HhaI- 100 bp ladder used



TIDE Assay Results: RNP1 and RNP2 samples with indels. Donor A at 1μM with indels, at 3 μM no indels. Donor B at 1 μM and 2 μM with no indels



CONCLUSION

- Our experiment did not result in HDR in HUDEP-S cells while resulted in indels in RNP control groups. This could be attributable to ssODN toxicity, instability or electroporation conditions. Further experiments are needed to optimize gene editing in HUDEP-S cells.