OMTM, Volume 23

Supplemental information

Multiplex CRISPR/Cas9 genome editing in

hematopoietic stem cells for fetal hemoglobin

reinduction generates chromosomal translocations

Clare Samuelson, Stefan Radtke, Haiying Zhu, Mallory Llewellyn, Emily Fields, Savannah Cook, Meei-Li W. Huang, Keith R. Jerome, Hans-Peter Kiem, and Olivier Humbert



Supplementary Figure S1. Indel patterns in individual BFU-Es following single or dual

CRISPR/Cas9 editing. Indel patterns observed at HBG-113 and BCL11A-ee within individual BFU-Es were analyzed, following single, simultaneous dual or sequential dual editing reactions. A reduction in larger indels was observed with simultaneous dual editing, particularly at the HBG-113 locus, consistent with saturation of the MMEJ repair pathway. *Abbreviations*: BCL11A: BCL11A-ee; HBG: HBG-113.



Supplementary Figure S2. Distribution of allele numbers edited in each reaction at each locus, on analysis of single BFU-Es (%)



Supplementary Figure S3. Editing in humanized mouse tissues following transplantation with single or dual-edited CD34⁺ cells.

(A) HBG-113 total editing within human cells in the PB, BM and spleen at necropsy: ratio to editing frequency within infusion product for each arm (mean \pm SD). (B) BCL11A-ee total editing within human cells in the PB, BM and spleen at necropsy: ratio to editing frequency within infusion product for each arm (mean \pm SD). (C) HBG-113 13-nt deletion frequencies within human cells in the PB, BM and spleen at necropsy (mean \pm SD). (D) BCL11A-ee 13- and 15-nt deletion frequencies within human cells in the PB, BM and spleen at necropsy (mean \pm SD). (E) HBG-113 13-nucleotide deletion frequencies within human cells in the PB, BM and spleen at necropsy (mean \pm SD). (E) HBG-113 13-nucleotide deletion frequencies within human cells in the PB, BM and spleen at necropsy: ratio to editing frequency within infusion product for each arm (mean \pm SD). (F) BCL11A-ee 13- and 15-nt deletion frequencies within human cells in the PB, BM and spleen at necropsy: ratio to editing frequency within infusion product for each arm (mean \pm SD). (F) BCL11A-ee 13- and 15-nt deletion frequencies within human cells in the PB, BM and spleen at necropsy: ratio to editing frequency within infusion product for each arm (mean \pm SD). (F) BCL11A-ee 13- and 15-nt deletion frequencies within human cells in the PB, BM and spleen at necropsy: ratio to editing frequency within infusion product for each arm (mean \pm SD). (F) BCL11A-ee 13- and 15-nt deletion frequencies within human cells in the PB, BM and spleen at necropsy: ratio to editing frequency within infusion product for each arm (mean \pm SD). Abbreviations: BCL11A: BCL11A-ee; HBG: HBG-113; nt: nucleotide.