Revally

Pin-point[™] Base Editing System: A Versatile Editing Platform Driving Cell Therapies

Bronwyn Joubert, Robert Blassberg, Jennifer Harbottle, Olga Mielczarek, Paul Russell, Jesse Stombaugh, Anastasia Lomova Kaufman, John Lambourne, Kevin Hemphill, Matthew Perkett, Žaklina Strezoska, Anja van Brabant Smith, Immacolata Porreca

Horizon Discovery, 2650 Crescent Drive, Suite 100, Lafayette, CO 80026, United States | Email address: baseediting@horizondiscovery.com





A) Schematic of one possible configurations of the Pin-point base editing system. B) Recruitment of APOBEC1 via an aptamer encoded in the sgRNA leads to efficient C to T conversion within a window centred on nucleotides 5-7 of the target sequence. C) Negligible base editing occurs in the absence of an aptamer. Data are average base editing activity in a HEK293 cell line across a series of genomic loci targeted by the Pin-point system.



Figure 4: Multi-gene edited CAR-T cells using the Pin-point system retain therapeutic function A) Percentage of CAR positive T cells in the transduced population. B) Protein knockout of 4 targets in CAR transduced T Cells analyzed by flow cytometry. C) In vitro tumor cell killing assay performed with mock electroporated or T cells edited with the Pin-point system and transduced with either empty or CAR expressing lentiviral vectors. n = 2 T cell donors for A-C.

Figure 7: iPSCs edited at multiple targets with the Pin-point base editor show improved viability A) Base editing at targeted cytosines analyzed by amplicon sequencing 7 days post electroporation. B) Viability of iPSCs edited with single or multiple sgRNAs using Pin-point or Cas9 analyzed by flow cytometry 48hrs post electroporation. Viability was normalized to a no sgRNA control. C) Expression of the pluripotency marker Nanog analyzed by flow cytometry 4 days post electroporation. Data shown in B,C are from two independent experiments with 2 iPSC lines.

One-step simultaneous knock-in and multiplex knock-out in iPSC



Figure 8: Pin-point system enables simultaneous site specific transgene integration and multiple gene KO in

A) Schematic of plasmid-mediated insertion of a transgene (promoter-less GFP) at the B2M locus. B) B2M knockout analyzed by



flow cytometry. C) Indel formation at target loci following Pin-point base editing and Cas9 gene editing analyzed by amplicon sequencing. n = 3 T cell donors.

T cells were activated before delivery of Pin-Point mRNA and synthetic sgRNAs targeting B2M, CD52, PDCD1 and TRAC by electroporation. Electroporated cells were either cultured unstimulated, or stimulated with PMA and lonomycin 5 days post-electroporation for assessment of PD1 knockout. Analysis was performed 7 days post-electroporation.

T cells were activated, transduced with lentiviral particles containing either no insert "Empty", or anti-CD19 CAR, and selected with puromycin. Pin-Point mRNA and synthetic sgRNAs targeting B2M, CD52, PDCD1 and TRAC were delivered by electroporation. Electroporated cells were either cultured unstimulated, or stimulated with PMA and Ionomycin 5 days post-electroporation for assessment of PD1 knockout. Analysis was performed 7 days post-electroporation.

One-step simultaneous knock-in and multiplex knock-out in T cells

ຶ່ 100-

80-

60

20-

Control

CAR



Figure 5: Pin-point system enables simultaneous site-specific CAR integration and quadruple gene KO in T cells

A) Schematic of AAV-mediated insertion of anti-CD19 CAR at the TRAC locus. B) Protein knockout of 4 targets in CAR transduced T cells analyzed by flow cytometry. C) Percentage of CAR positive T cells in the transduced population. D) In vitro tumor cell killing assay performed with T cells edited with the Pin-point system and transduced with AAV to express the CD19 CAR versus untransduced cells edited with the Pin-point system (1:1 and 5:1 are CAR-T : target cells ratios). n = 2 T cell donors.

T cells were activated before delivery of Pin-Point mRNA and synthetic sgRNAs targeting B2M, CD52, PD1 and TRAC by electroporation. This was followed by delivery of AAV particles loaded with the HDR donor DNA for the CD19-CAR by transduction. Flow cytometry analysis was performed 5 days postelectroporation. The killing assay was performed by co-culturing the CAR-T cells with antigen positive cancer cells and measuring the percentage of killed target cells.

Conclusions

- We applied multiplex base editing with the Pin-point system for the development of engineered CAR-T cells and hypoimmunogenic iPSCs.
- Base editing with the Pin-point system achieved greater than 70% knockout efficiency and high purity at therapeutically relevant target sites (B2M, CD52, TRAC and PDCD1) in T cells and iPSCs. Multiplex gene editing with the Pin-point platform substantially reduces guide-dependent off target
- editing and chromosomal translocations compared to Cas9-mediated knockout.
- CAR-T cells retain their cytotoxic activity in vitro following multiplex gene editing with the Pin-point system
- The Pin-point platform enables simultaneous site-specific transgene knock-in and multi-gene knockout in T cells and iPSCs.
- iPSCs edited with the Pin-point platform retain pluripotency.
- We observed greatly improved viability of iPSCs following multi-gene editing with the Pin-point system compared to Cas9.

References

Collantes et al. The CRISPR Journal. Feb 2021.58–68



©2023 Horizon Discovery Group Ltd and its affiliated companies. All rights reserved. Revvity is a trademark of Revvity, Inc., registered in the United States and other countries. Horizon Discovery is a trademark mark of Horizon Discovery Ltd. Dharmacon is a trademark of Dharmacon, Inc. All Revvity, Inc., Horizon Discovery Ltd and Dharmacon, Inc. trademarks are used with permission. Other product names and brand names may be trademarks or registered trademarks of their respective owners.