Clinical Application of Sleeping Beauty and Artificial Antigen Presenting Cells to Genetically Modify T Cells from Peripheral and Umbilical Cord Blood

M. Helen Huls¹, Matthew J. Figliola¹, Margaret J. Dawson¹, Simon Olivares¹, Partow Kebriaei², Elizabeth J. Shpall², Richard E. Champlin², Harjeet Singh¹, Laurence J.N. Cooper¹

¹Division of Pediatrics, U.T. MD Anderson Cancer Center
²Department of Stem Cell Transplantation and Cellular Therapy, U.T. MD Anderson Cancer Center

Correspondence to: Laurence J.N. Cooper at ljncooper@mdanderson.org

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Abstract

The potency of clinical-grade T cells can be improved by combining gene therapy with immunotherapy to engineer a biologic product with the potential for superior (i) recognition of tumor-associated antigens (TAAs), (ii) persistence after infusion, (iii) potential for migration to tumor sites, and (iv) ability to recycle effector functions within the tumor microenvironment. Most approaches to genetic manipulation of T cells engineered for human application have used retrovirus and lentivirus for the stable expression of CAR. Our approach, although compliant with current good manufacturing practice (GMP), can be expensive as it relies on the manufacture and release of clinical-grade recombinant virus from a limited number of production facilities. The electro-transfer of nonviral plasmids is an appealing alternative to transduction since DNA species can be produced to clinical grade at approximately 1/10⁶ the cost of recombinant GMP-grade virus. To improve the efficiency of integration we adapted Sleeping Beauty (SB) transposon and transposase for human application. Our SB system uses two DNA plasmids that consist of a transposon coding for a gene of interest (e.g. 2nd generation CD19-specific CAR transgene, designated CD19RCD28) and a transposase (e.g. SB11) which inserts the transgene into TA dinucleotide repeats. To generate clinically-sufficient numbers of genetically modified T cells we use K562-derived artificial antigen presenting cells (aAPC) (clone #4) modified to express a TAA (e.g. CD19) as well as the T cell costimulatory molecules (stimulation cycle) of γ-irradiated aAPC (clone #4) in the presence of soluble recombinant human IL-2 and IL-21. Typically 4 cycles (28 days of continuous culture) are undertaken to generate clinically-appealing numbers of T cells that stably express the CAR. This methodology to manufacturing clinical-grade CD19-specific T cells can be applied to T cells derived from peripheral blood (PB) or umbilical cord blood (UCB). Furthermore, this approach can be harnessed to generate T cells to diverse tumor types by pairing the specificity of the introduced CAR with expression of the TAA, recognized by the CAR, on the aAPC.

Video Link

The video component of this article can be found at http://www.jove.com/video/50070/

Protocol

Day 0 or Before

1. Isolation of Mononuclear Cells (MNC) from PB and UCB

1. Dilute PB with equal volume, and UCB with four volumes of PBS-EDTA.
2. Slowly layer diluted blood (25 ml) onto Ficoll (12 ml) in a 50 ml centrifuge tube(s) and centrifuge at 400 x g for 30 - 40 min (no brake).
3. Collect and transfer the mononuclear cell fraction (interface) using a transfer pipette to a fresh 50 ml centrifuge tube.
4. Bring up the volume to 50 ml with PBS-EDTA and centrifuge at 450 x g for 10 min.
5. Aspirate the supernatant, gently re-suspend the cell pellet(s) in 50 ml of Complete Culture Media (CCM) and, centrifuge at 400 x g for 10 min.
6. Gently re-suspend and pool the cell pellets in CCM and perform a cell count using Trypan blue exclusion method (Cellometer, PBMC program).
7. MNC can now be used for electroporation (Nucleofection) or cryopreserved for future use.

2. Preparation of T Cells for Electroporation on Day 0

1. If using cryopreserved MNC, quickly thaw sufficient cells for a full scale electroporation (2 x 10^8 adding ~20% to account for cell loss during centrifugation and 2 hr incubation) in 37 °C water bath. If using freshly isolated MNC proceed to step 2.3.
2. Gently re-suspend and transfer cells to an appropriately sized centrifuge tube(s) contained pre-warmed Complete Phenol-Free RPMI culture media (PF-RPMI) and centrifuge at 200 x g for 10 min (no brake), aspirate supernatant.
3. Re-suspend the MNC in PF-RPMI, perform a cell count (Cellometer) and transfer cells to an appropriately sized cell culture vessel(s) at a concentration of 10^6 cells/ml.
4. Incubate in a humidified 37 °C/5% CO₂ incubator for 2 hr ± 30 min.
5. Transfer the MNC to sterile centrifuge tube(s), spin at 200 x g for 5 min (no brakes), aspirate the supernatant and gently re-suspend and combine the cell pellet(s) in PF-RPMI.
6. Perform a cell count (Cellometer) and calculate the volume of the cell suspension required (2 x 10^6 MNC).
7. Transfer the calculated volume to a sterile 50 ml centrifuge tube and spin at 200 x g for 10 min (no brake).
8. Aspirate the supernatant so that no residual media remains and gently re-suspend by tapping side of tube.

3. Electroporation (Nucleofection) of MNC (Full Scale Process Using 10 Cuvettes) on Day 0

1. Pre-incubate a sterile 12-well plate with 10 wells containing 4 ml of warm PF-RPMI in a humidified 37 °C/5% CO₂ incubator.
2. Prepare and pre-warm the Lonza Nucleofector Solution Human T cell kit (reconstituted per manufacturer's instructions, www.lonza.com) to ambient temperature in a Biosafety Cabinet (BSC).
3. Prepare Nucleofector solution/DNA master mix by adding 100 μl of supplemented Nucleofector solution, 15 μg of transposon (supercoiled DNA plasmid designated as CD19RCD28/pSBSO) and 5 μg of transposase (supercoiled DNA plasmid designated as pCMV-SB11) per reaction/cuvette.
4. Disperse the cell pellet (from step 2.8) by gently tapping the side of the centrifuge tube and re-suspend in Nucleofection solution/DNA master mix (Final cell concentration: 2 x 10^7 cells/100 μl).
5. Carefully transfer 100 μl of the cell suspension (from step 3.4) to each of ten (10) Lonza Nucleofection cuvettes, being careful to avoid bubbles.
6. Tap the cuvette once, and electroporate using program U-014 (for unstimulated T cells).
7. Transfer the cuvettes and the 12-well plate (Step 3.1) to the BSC.
8. Harvest the electroporated cells from each cuvette using an Amaxa fine tip transfer pipette, by adding ~500 μl of the pre-warmed culture medium from the corresponding well (12-well plate prepared in Step 3.1) and return the plate to a humidified 37 °C/5% CO₂ incubator for 2 hr ± 30 min.
9. Following the 2-hr incubation, harvest and transfer the cells from all wells to a sterile centrifuge tube.
10. Wash cells by centrifugation at 140 x g for 8 min, ambient temperature, no brake and aspirate and discard the supernatant so that no residual medium covers the cell pellet.
11. Disperse the cell pellet by gently tapping the side of the centrifuge tube and gently re-suspend in Nucleofection solution/DNA master mix (Final cell concentration: 2 x 10^6 cells/100 μl).
12. Perform cell count and adjust cell concentration to 10^6 cells/ml in CCM.
13. Transfer cell suspension to cell culture flask(s) and place in incubator overnight.
14. Process steps 2.1 to 3.8 were repeated for control: EGFP-transfected cells (5 x 10^6 cells/cuvette with 5 μg Amaxa control EGFP supercoiled plasmid, pmaxGFP).

Day 1 of 1st and Subsequent Stimulation Cycles

4. Analysis of CAR Expression by Flow Cytometry on Day 1

1. Harvest the electroporated cells and perform a cell count using Trypan blue exclusion method (Hemocytometer).
2. Stain cells (1 to 2 x 10^6) with antibody specific for CD3, CD4, CD8, and human IgG Fcγ (as a measurement of CAR expression).
3. Acquire cells on FACŚ Calibur and analyze the data using FCS Express software to calculate expression of CAR.
   1. Calculate CAR⁺ cells in culture by the formula: 
     \[(\text{No. of Total viable cells}) \times \text{(% CAR⁺ cells)} = \text{No. of \text{CAR⁺} cells}\]

5. Preparation of aAPC (clone #4) on Day 1. aAPC (clone #4) were Derived from K562 Cells (Parental Line Obtained from American Type Culture Collection) to Co-express Desired T Cell Co-stimulatory Molecule

1. Thaw an aliquot of frozen 100 Gy irradiated aAPC in a 37 °C water bath.
2. Cells are washed twice by centrifugation at 400 x g, 10 min in CCM and counted using Cellometer (Trypan blue exclusion).
3. Calculate number of viable aAPC required for stimulation: 
   \[(\text{No. of \text{CAR⁺} cells}) \times 2 = \text{No. of irradiated aAPC required}\]
6. aAPC-mediated Stimulation of CAR T Cells on Day 1 Beginning of 1st and Subsequent Stimulation Cycles

1. Mix electroporated cells (expressing CAR) and γ-irradiated aAPC (clone #4) in a sterile container at a ratio of 1:2 (CAR cell: viable aAPC) in CCM.
   1. **Note** the aAPC ratio is adjusted for the expression of CAR based on flow cytometry the day after electroporation.
2. Add IL-21 (30 ng/ml) to the cell suspension.
3. Aliquot in T-75 cm² flask(s) and/or Vue Life Culture bags at a concentration of 10⁶ cells/ml and return to the incubator.

Days 3, 5

7. Continued Culture of CAR T Cells

1. Perform a half-media change, replenish IL-21, and maintain T cells at a concentration of 10⁶ cells/ml.

Day 7

8. End of First aAPC-mediated Stimulation Cycle

1. Harvest cells, count and stain for CD3, CD4, CD8, and Fcγ (CAR) and proceed to step 10.1.

9. Depletion of CD56 Cells (Usually between 7 and 14 days after electroporation)

1. Perform a CD56 depletion using paramagnetic beads if CD56+CD3neg lymphocytes ≥ 10%.

Stimulation Cycles #2, #3, & #4 Corresponding to Days 8 → 14, Days 15 → 21, & Days 22 → 28

10. Recursive Addition of aAPC to Propagate T Cells to Clinically-sufficient Numbers

1. Repeat the stimulation process (4 times) as described in steps 4.3-8.1.
2. Add IL-2 (50U/ml) to the cultures beginning on Days 7, 14 and 21, & then at each media change (three times a week, on a Monday-Wednesday-Friday schedule).
3. Cryopreserve (archive) excess T cells as needed.
   1. T cells frozen using controlled rate freezer.

Day 28

11. End of Last aAPC-mediated Stimulation Cycle: Harvest T Cells

1. Cryopreserve T cells for release testing and infusion.

**Representative Results**

We report that electro-transfer of DNA plasmids and propagation of T cells on γ-irradiated aAPC can be used to generate clinically-appealing numbers of T cells derived from PB and UCB for human applications. These genetically modified T cells express an introduced CAR that recognizes the TAA CD19, independent of major histocompatibility complex. The SB-derived DNA plasmids to express the (i) transposon, a 2nd generation CAR (CD19RCD28) that signals through CD28 and CD3-ε[314], and (ii) transposase, SB11[15], have been previously described[13,16,17]. The plasmids used in the current study were produced commercially by Waisman Clinical Biomanufacturing Facility (Madison, WI). The aAPC (clone #4), derived from K562 cells (parental line obtained from American Type Culture Collection), co-express desired T cell co-stimulatory molecules (each introduced molecule at ≥90% on cell surface of aAPC), as previously described[12]. Here we show that CD19-specific T cells could be generated from mononuclear cells (MNC) derived from PB or UCB using SB transposition to introduce the CAR followed by addition of aAPC to numerically expand the T cells in a CAR-dependent manner (Figures 1, 4).[13,18] Ten cuvettes (2x10⁷ MNC/cuvette) are electroporated for each recipient using 15 μg of DNA plasmid (CD19RCD28/pSBSO) coding for transposon (CAR) and 5 μg of DNA plasmid (pCMV-SB11) coding for transposase (SB11). The number of cuvettes can be reduced if MNC are limiting or scaled back for laboratory work. The day of electroporation is defined as “Day 0” of Stimulation cycle #1. As controls for flow cytometry and culture conditions, autologous T cells are mock electroporated (without DNA plasmid) and numerically expanded on γ-irradiated aAPC (clone #4) that had been pre-loaded with OKT3 to cross-link CD3 to sustain T cell proliferation. We routinely assess the efficiency of electrotransfer and viability of the T cells the day after electroporation (Figure 2B). The expression of EGFP from control DNA plasmid (designated pmaxGFP) and CAR at this initial time point reflects protein expression from the integrated and episomal plasmid. Typically, the day after electroporation we measure EGFP expression at ~60% and CAR expression at ~40% (Figure 2A) with T cell viability between 40-50%. Recursive additions of γ-irradiated aAPC in the presence of soluble recombinant human IL-2 and IL-21 retrieve T cells stably expressing CAR (CD19RCD28). CD3negCD56+ NK cells are depleted from the culture using CD56-specific paramagnetic beads if the percentage of these NK cells is ≥10% and especially if the percentage of CAR expressed on the T cells is low. This depletion prevents the rapid overgrowth of NK cells which interferes with the ability of aAPC to sustain the proliferation of CAR T cells. On occasion, depletion of NK cells from CAR T cells is undertaken during the last two stimulation cycles, but this introduces a...
loss of desired cells due to co-expression of CD56 on some CAR+ T cells. The T cells were grown in a functionally closed system using Vue Life culture bags past Day 14. A subset of the genetically modified and propagated T cells are typically cryopreserved at Day 14 or Day 21 (end of Stimulation cycles #2 or #3) of co-culture on aAPC to serve as a source of archived material for future analyses and to be thawed if unanticipated problems subsequently occur during the manufacturing process. T cells are typically harvested on or about Day 28 of culture (Figure 3) that routinely express >90% CAR and are >80% viable (Figure 2C, D). We have previously shown that, after four weeks of co-culture on aAPC the average fold-expansion of CD3+ T cells is 19,800±11,313 with CAR+ expression being 90%±7.5 (13). These T cells are cryopreserved and undergo in-process and release testing that informs on the safety and therapeutic potential of the manufactured product. Release testing is undertaken in compliance with clinical laboratory improvement amendments (CLIA) to generate a certificate of analysis prior to infusion into recipients on clinical trials.

Figure 1. Steps outlining the process to electroporate and propagate CAR+ T cells from PB and UCB. Click here to view larger figure.
Figure 2. Characterization of genetically modified T cells from PB. (A) Expression of EGFP at Day 0 of first stimulation cycle to assess the efficiency of gene transfer. Expression of CD19-specific CAR (CD19RCD28) as assessed by flow cytometry on CD3⁺, CD8⁺ and CD4⁺ T cells at (B) approximately 24 hr after electroporation and (C) 28 days after co-culture on aAPC. Similar expression of CAR was observed with UCB-derived T cells. (D) Kinetics of CAR expression. Click here to view larger figure.
Figure 3. Propagation of PB-derived CAR$^+$ T cells. Rate of numeric expansion of CD3$^+$ and CAR$^+$ T cells derived from PB by repeated coculture on γ-irradiated aAPC in presence of recombinant human soluble IL-2 and IL-21. Upward arrows indicate the additions of γ-irradiated aAPC that mark the beginning of each Stimulation cycle. UCB-derived CAR$^+$ T cells exhibit similar rates of numeric expansion.

Figure 4. Schematic of the manufacturing process using SB and aAPC systems to genetically modify and propagate CAR$^+$ T cells derived from PB and UCB. CD19-specific CAR$^+$ T cells were generated by electro-transfer of SB-derived supercoiled DNA plasmids and subsequent co-culture on K562-derived aAPC (clone #4) in the presence of recombinant human soluble IL-2 and IL-21. Click here to view larger figure.

| T-cell source                      | Transposon* | Transposase | T-cell infusion                                             | IRB #      | NIH-OBA #   | IND #  |
|-----------------------------------|-------------|-------------|-------------------------------------------------------------|------------|-------------|--------|
| Autologous (patient-derived)**    | CD19RCD28   | SB11        | After autologous hematopoietic stem-cell transplantation    | 2007-0635  | 0804-922    | 14193  |
| Allogeneic (donor-derived)        | CD19RCD28   | SB11        | After allogeneic hematopoietic stem-cell transplantation    | 2009-0525  | 0910-1003   | 14577  |
| Allogeneic (donor-derived)        | CD19RCD28   | SB11        | After allogeneic umbilical cord blood transplantation       | 2010-0835  | 1001-1022   | 14739  |
Table 1. Clinical trials under the auspices of the FDA at MDACC to infuse CD19-specific CAR+ T-cells propagated on aAPC. *T-cells are rendered specific for CD19 through enforced expression of SB transposon coding for a 2<sup>nd</sup> generation CAR, designated CD19RCD28, that signals through CD28 and CD3-z. **Trial described in reference # 7

Discussion

Transposon and transposase systems, such as from piggyBac<sup>12,19</sup> and SB<sup>18,20,22</sup>, are non-viral approaches to gene therapy that are an alternative to viral-mediated transduction of clinical grade CAR<sup>+</sup> T cells. The SB was chosen as the gene transfer system based on its potential for human gene therapy<sup>19,23</sup>. We developed the dual technologies of SB transposition (to introduce a CAR) and recursive addition of γ-irradiated aAPC (to retrieve genetically modified T cells stably expressing a CAR) to serve as platform technologies in the manufacture of TAA-specific T cells in compliance with cGMP for Phase III/II trials (Figure 4). After 28 days (four 7-day stimulation cycles) of co-culture on γ-irradiated aAPC, we are typically able to generate at least ~10<sup>11</sup> genetically modified T cells suitable for human applications. As needed, additional stimulation cycles can be undertaken to generate larger numbers of genetically modified T cells. Furthermore, if less CAR<sup>+</sup> T cells are needed, the approach to electroporation and propagation can be scaled back employing fewer cuvettes and carrying forward just a sub-set of the numerically expanded T cells for subsequent rounds of proliferation on aAPC at the beginning of each Stimulation cycle. The majority of the electroporated and propagated T cells harvested for infusion stably express the CAR. The outgrowth of CD4<sup>+</sup> and CD8<sup>+</sup> T cells expressing our 2<sup>nd</sup> generation CAR include cells with a memory/naive phenotype and exhibit three hallmarks of re-directed specificity. Firstly, the genetically modified T cells specifically lyse CD19<sup>+</sup> targets, secondly, produce IFN-γ in response to CD19<sup>+</sup> stimulator cells and thirdly, proliferate in response to CD19<sup>+</sup> feeder cells, all in a CAR-dependent manner<sup>13,18</sup>. Our approach to integrate a CAR transgene by electro-transfer of non-viral DNA plasmids from the SB system can be undertaken in quiescent primary T cells derived from PB and UCB. We and others have genetically modified K562 cells to serve as aAPC to efficiently propagate clinically-sufficient numbers of T cells for infusion<sup>24,25</sup>. The aAPC and tissue culture environment (e.g. the addition of IL-21) have been have been modified to generate patient- and donor-derived CD19-specific T cells for infusion after hematopoietic stem-cell transplantation (Table 1)<sup>13,19</sup>. We can produce CAR<sup>+</sup> T cells from PB simply obtained by venipuncture which avoids the cost, discomfort, and inconvenience of obtaining MNC from PB by apheresis. The ability to derive large numbers of CAR<sup>+</sup> T cells from small numbers of MNC is particularly appealing for infusing T cells after allogeneic UCB transplantation. The small size and anonymity of the neonatal donor precludes re-accessing this individual at a later time point and only limited numbers of harvested MNC are available as starting material for T cell manufacture to avoid interfering with hematopoiesis. Further advances to the manufacturing process are currently underway to include a high throughput electroporation device coupled with a fully closed WAVE bioreactor to minimize handling. In aggregate, the SB and aAPC are appealing platforms to generate CD19-specific CAR<sup>+</sup> T cells that can be adapted to generate large numbers of genetically modified T cells that can recognize alternative cell-surface TAAs in compliance with cGMP.

Disclosures

No conflicts of interest declared.

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