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New Cell Sources for T Cell Engineering and Adoptive Immunotherapy

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Abstract

The promising clinical results obtained with engineered T cells, including chimeric antigen receptor (CAR) therapy, call for further advancements to facilitate and broaden their applicability. One potentially beneficial innovation is to exploit new T cell sources that reduce the need for autologous cell manufacturing and enable cell transfer across histocompatibility barriers. Here we review emerging T cell engineering approaches that utilize alternative T cell sources, which include virus-specific or T cell receptor-less allogeneic T cells, expanded lymphoid progenitors, and induced pluripotent stem cell (iPSC)-derived T lymphocytes. The latter offer the prospect for true off-the-shelf, genetically enhanced, histocompatible cell therapy products.

T cells are essential mediators of immune defense against infectious pathogens and cancer. Their insufficiency, which occurs in hereditary or acquired immune deficiencies, results in life threatening infections, increased cancer incidence, and disrupted immunoregulation. T cells can also be harmful and cause normal tissue destruction, as seen in autoimmune disorders, graft rejection, and graft-versus-host disease (GVHD). T cells develop from precursors that rearrange germline antigen receptor VDJ genes in the thymus, thereby generating clonotypic T cell receptors (TCRs) that undergo positive and negative thymic selection (Figure 1). The resulting T cells are self-restricted and tolerant of self tissues. The newly generated T cell clones, known as naive T cells, initially circulate throughout the body at low frequency. Upon encountering antigen, T cells expand and acquire effector and/or memory functions. This T cell priming requires TCR engagement by Human Leucocyte Antigen (HLA)-peptide complexes on the surface of antigen presenting cells (APCs) and concomitant ligation of costimulatory receptors by ligands borne by the APCs (Chen and Flies, 2013; Krogsgaard and Davis, 2005).

Pathogen-specific T cells can be effectively expanded through vaccination, a medical intervention that allows prevention of a number of infectious diseases. In this instance, immunization proceeds in vivo within secondary lymphoid organs where T cells engage their TCRs on professional APCs that initiate productive T cell activation and clonal expansion. Active immunization has, however, proven far less effective when infection or cancer is already established and progressing. In such circumstances, T cells, whether they

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are naturally activated or elicited through immunization, often fail to eradicate disease owing to their inadequate number or suboptimal function.

The infusion of T cells, or adoptive transfer, has proven to overcome the limitations of active immunization in some pathologies. The therapeutic use of isolated T cells began somewhat inadvertently with allogeneic bone marrow transplantation (BMT). The use of whole marrow grafts containing donor T cells revealed the beneficial (graft-versus-tumor responses) and deleterious (GVHD) effects of adoptive T cell transfer (Ferrara and Deeg, 1991). Several forms of T cell therapy subsequently developed, including donor leukocyte infusion (Kolb et al., 2005) and virus-specific T cell therapy (Riddell and Greenberg, 1995). These therapies utilize “donor-derived T cells,” which tap into the alloreactive potential of T cells harvested from a healthy donor but expose the recipient to the risk of normal tissue destruction by graft versus host (GVH) responses. In contrast, autologous T cells, harvested from the intended recipient (Rosenberg et al., 1986), are devoid of such toxic potential. However, autologous T cells with therapeutic potential may be lacking or functionally impaired in patients with refractory infections or progressing cancer. Allogeneic and autologous T cells thus have their respective advantages and disadvantages.

For some cancers, T cells may be isolated from surgically removed tumors, which are enriched in tumor-reactive T cells relative to peripheral blood. Tumor infiltrating lymphocytes (TILs) can be isolated at quite a high frequency from melanoma specimens, but this technique is not feasible or effective in many other tumor types (Rosenberg et al., 2008; Wu et al., 2012). Thus, we and others have sought to generate tumor-targeted T cells through genetic engineering (Ho et al., 2003; Sadelain et al., 2003). The rationale for T cell engineering is to rapidly generate populations of T cells specific for any antigen and, furthermore, to enhance their therapeutic (e.g., anti-tumor) functions. Peripheral blood T cells are easily accessible and are a perfectly suitable cell source for this purpose. Most current therapies utilizing engineered T cells process autologous peripheral blood T cells that are targeted to tumor antigens following retroviral transduction of a TCR or a chimeric antigen receptor (CAR). In recent years, a few clinical trials have resulted in encouraging and sometimes dramatic clinical responses (Couzin-Frankel, 2013). This Perspective article focuses on the sources of T cells for adoptive cell therapy, starting from blood, hematopoietic stem cell-derived lymphoid progenitor cells, embryonic stem cell (ESC), or induced pluripotent stem cell (iPSC)-derived T cells.

T Cell Engineering with TCRs and CARs

The general premise for engineering T cells for cancer immunotherapy is to rapidly generate tumor-targeted T cells, bypassing the obstacles that preclude the induction and execution of effective immune responses *in vivo*. Two categories of antigen receptors are used to retarget T cell specificity: physiological TCRs and synthetic receptors referred to as CARs (Figure 2). The design of TCRs and CARs has steadily improved over the past 2 decades (Cohen et al., 2006, 2007; Robbins et al., 2008; Sadelain et al., 2003, 2009, 2013; Voss et al., 2008). TCRs are typically cloned from patient tumor-reactive T cell clones (Johnson et al., 2006), from humanized murine models (Cohen et al., 2005; Parkhurst et al., 2009), or through the use of phage display technology (Li et al., 2005; Varela-Rohena et al., 2008). In CARs,

tumor recognition is mediated by a single chain variable fragment (scFv) derived from a monoclonal antibody or an antigen-binding region isolated from an immunoglobulin (Ig) heavy and light chain library. Unlike TCR-mediated antigen recognition, CARs function independently of HLA and can therefore be used in any genetic background. Second-generation CARs (Maher et al., 2002, Figure 2) not only mediate antigen recognition and initiate T cell activation but also harness costimulation to enhance T cell function and prolong T cell persistence (Sadelain et al., 2009). Over a decade ago, we selected the CD19 antigen as a potential CAR target for B cell malignancies (Brentjens et al., 2003) and made it the focus of our CAR therapy program. Impressive results were obtained in patients with relapsed, chemorefractory B cell malignancies. A number of patients with chemorefractory B cell malignancies developed complete responses after a single infusion of CAR T cells, as first reported by the National Cancer Institute for B cell lymphoma (Kochenderfer et al., 2010, 2012), the University of Pennsylvania for chronic lymphocytic leukemia (Kalos et al., 2011; Porter et al., 2011; Brentjens et al., 2011), and Memorial Sloan Kettering Cancer Center for acute lymphoblastic leukemia (ALL) (Brentjens et al., 2013b; Davila et al., 2014a; Grupp et al., 2013; Lee et al., 2015; Maude et al., 2014). These patients were treated with autologous T lymphocytes that were retrovirally transduced with second-generation, CD19-specific CARs (Davila et al., 2012). To date, the most dramatic results have been obtained in adult and pediatric patients with ALL (Brentjens et al., 2013b; Davila et al., 2014a; Grupp et al., 2013; Lee et al., 2015; Maude et al., 2014). Encouraging results have also been obtained in patients with CD19+ lymphomas, reviewed in Ramos et al. (2014) and Kochenderfer and Rosenberg (2013). Third-generation CARs, which contain two costimulatory domains along with an activation domain, may provide superior T cell function (Carpenito et al., 2009; Pule et al., 2005; Till et al., 2012; Zhong et al., 2010), although their effectiveness remains to be evaluated in clinical trials.

Autologous T Cell Manufacture

The genetic modification of autologous peripheral blood T lymphocytes to generate tumor-targeted T cells is now a well-established approach that was developed in a handful of academic centers. The power and promise of TCR and CAR therapies utilizing these manufacturing processes are best illustrated by the exciting clinical results obtained with NY-ESO-1 TCR (Robbins et al., 2011) and CD19 CAR T cells (Brentjens et al., 2013b; Davila et al., 2014a; Grupp et al., 2013; Kochenderfer et al., 2012, 2014).

These cell manufacturing processes combine T cell activation and transduction steps to generate expanded, genetically targeted T cell products. For example, T cells engineered to express specific CARs or TCRs may be initiated from Ficoll-purified PBMCs, which are next activated with anti-CD3 monoclonal antibody (mAb) in the presence of irradiated allogeneic feeder cells and transduced with a vector encoding either the CAR or TCR α and β chains (Till et al., 2012; Morgan et al., 2006). We and others have established cGMP-compliant large-scale transduction and expansion processes, which are applicable to CARs or TCRs, utilizing either γ -retroviral or lentiviral T cell manufacturing (Figure 3). These processes begin with the selection and activation of T cells from patient apheresis products using materials coated with anti-CD3 and anti-CD28 mAbs. In the case of iron beads, CD3⁺CD28⁺ T cells are enriched using a magnetic particle concentrator and subsequently

cultured. Activated T cells are retrovirally transduced in RetroNectin-coated cell bags and inoculated in a WAVE bioreactor where they are expanded with a continuous perfusion regimen, reaching cell densities of 10 million T cells/ml or more (Hollyman et al., 2009). At the end of the production, the beads are removed and the cells are formulated for immediate infusion or frozen for deferred use. The entire process typically takes 10–14 days, depending on the disease and the targeted T cell dose. This semi-closed large-scale manufacturing platform can be easily adapted for various vectors and for the expansion of either patient (autologous) or donor (allogeneic) T cells. It successfully supports several ongoing clinical trials in which therapeutic efficacy has been demonstrated (Brentjens et al., 2011, 2013a; Davila et al., 2014b). This process starts from bulk T cells harvested from each individual subject. Several groups are currently evaluating what T cell phenotype and T cell subset or subsets account for the anti-tumor activity of these cells and what will be optimal tools to activate and expand T cells for different T cell therapies. Various means to enhance the activation and expansion of T cells for adoptive cell therapy have been reviewed elsewhere (Vacchelli et al., 2013).

The functional, proliferative, and persistence potential of adoptively transferred T lymphocytes is determined by multiple factors. These include the TCR or CAR design, the manufacturing platform, the selected T cell subsets, and the differentiation stage of the harvested T cells. Peripheral blood T cells comprise naive (T_N), stem cell memory (T_{SCM}), central memory (T_{CM}), effector memory (T_{EM}), and terminal effector (T_E) cells (Klebanoff et al., 2012). Several groups have investigated which of these T cell subsets are best suited for use in different adoptive therapy settings (Klebanoff et al., 2012; Riddell et al., 2014). In non-human primates and murine NSG models, T cell transfer studies have shown that virus-specific and CAR-redirection anti-tumor CD8 T_{EM} rapidly mature to terminal effector T cells and do not persist beyond 7–14 days, while a subset of transferred CD8⁺ $T_{E/CM}$ can acquire memory cell features and persist for months and even years (Wang et al., 2012). Polyclonal CD8⁺ T_{CM} isolation from leukopheresis products, followed by CD3/CD28 activation without exogenous feeder cells and cell expansion in IL-2/IL-15, has thus been developed on a clinical scale and is currently in use for the generation of autologous CAR-redirection CD19-specific CD8⁺ $T_{E/CM}$ for adoptive transfer after autologous hematopoietic stem cell transplantation (HSCT) for high-risk CD19⁺ non-Hodgkin lymphomas (Wang et al., 2012). Additional variations on the manufacturing schemas exemplified here have been reported or are under development (DiGiusto and Cooper, 2007; Laport et al., 2003; Savoldo et al., 2011; Somerville et al., 2012; Wang and Rivière, 2015).

It remains to be determined how the cell attributes imparted by in vivo persistence of antigen-specific T cells correlate with those conferring increased anti-tumor efficacy (Biasco et al., 2015; Flynn and Gorry, 2014; Xu et al., 2014). Defining optimal, potent T cell products of specified composition for adoptive cell therapy will require careful phenotypic and biological characterization, taking in account manufacturing and economic practicalities (Heathman et al., 2015).

Allogeneic T Cells as a Substrate for T Cell Engineering

The promising clinical results of engineered T cell therapy could be further amplified and broadened if potent and histocompatible T cells were readily available. Autologous approaches have a proven track record, but personalized manufacture may be challenging in some instances, for example in patients with chemotherapy or HIV-induced immune deficiency or in small infants. While T cells can be easily harvested from donors, their use is compromised by the high alloreactive potential. Owing to their ontogeny, TCRs are naturally prone to react against non-autologous tissues, recognizing either allogeneic HLA molecules or other polymorphic gene products, referred to as minor antigens (Afzali et al., 2007). This propensity underlies the high risk of graft rejection in transplant recipients and of GVHD in recipients of donor-derived T cells. Thus, bulk, unselected donor T cells are prone to cause normal tissue destruction and may be lethal on occasion. To provide an acceptable risk-benefit ratio, allogeneic T cells must be devoid of alloreactive potential. Two strategies designed to overcome the risk of GVH reactions have been proposed, based on the selection of virus-specific TCRs devoid of GVH reactivity or the ablation of TCR expression.

Virus-Specific T Cells

Donor-derived virus-specific T cells can be administered to virus-infected, HLA-matched recipients with a reduced risk of GVHD. For this purpose, donor T cells from a seropositive donor are stimulated in vitro with APCs that present the relevant viral antigens. The APCs may consist of syngeneic Epstein-Barr virus (EBV)-transformed B cells (Heslop et al., 1994), peptide pulsed dendritic cells, or artificial APCs (Latouche and Sadelain, 2000; Papanicolaou et al., 2003). Repeated stimulation with viral antigen gradually increases viral specificity and concomitantly depletes alloreactivity. Although alloreactivity and unanticipated TCR cross-reactivity cannot be prospectively eliminated with full certainty (Cameron et al., 2013; Morgan et al., 2013), virus-specific T cell lines generated in this manner have shown dramatic responses in EBV, cytomegalovirus (CMV) and adenovirus-infected recipients without causing severe GVHD (Heslop et al., 1996; Papadopoulos et al., 1994). Recent studies have suggested that virus-specific T cells can be administered to multiple recipients with limited risk of GVHD (Dobrovina et al., 2012; Haque et al., 2007). Virus-specific T cells may thus serve as cellular vehicles for TCR or CAR therapy. A first trial testing this approach showed that T cells expanded in vivo in response to viral reactivation although anti-tumor activity was modest (Cruz et al., 2013). While the relatively limited expansion potential of virus-specific T cells and the sometimes unpredictable cross-reactivity of TCR-mediated antigen recognition are valid concerns, this approach to treat viral infections represents a first step toward multi-recipient T cell product manufacturing (Wang and Rivière, 2015).

TRC-less Allogeneic Peripheral Blood T Cells

If the endogenous TCR cannot be tamed, one may abrogate its expression, making the engineered TCR or CAR the sole driver of T cell activation and clonal expansion. With the advent of gene disruption technologies, this approach is now within reach. Four technologies based on the use of targeted nucleases, including meganucleases, zinc-finger nucleases (ZFNs), transcription activator-like effector nucleases (TALEN), and CRISPR/Cas9, enable

gene disruption in human cells (Kim and Kim, 2014; Sander and Joung, 2014). ZFNs and CRISPR/Cas9 are presently the most developed of these tools and have been shown to efficiently target the HIV co-receptor CCR5 (Holt et al., 2010; Mandal et al., 2014; Tebas et al., 2014). The ablation of endogenous TCR expression has been achieved using targeted ZFNs or TALENs that disrupt the constant regions of TCRA and TCRB genes (Berdien et al., 2014; Provasi et al., 2012; Torikai et al., 2012). Unlike unedited, allogeneic T cells, TCR-deleted lymphocytes retargeted by CAR or TCR gene transfer do not mediate GVH reactivity. Their long-term persistence could potentially be compromised, since homeostatic proliferation is partially dependent on TCR-major histocompatibility complex (MHC) interactions. However, T lymphocytes that have acquired a central memory phenotype are less dependent on the TCR for homeostatic proliferation (Surh and Sprent, 2008) and proliferate in response to cytokines (Provasi et al., 2012). Their long-term persistence, relative to that of unedited T lymphocytes, has not yet been fully characterized. Furthermore, gene disruption technologies are still in early stages of development and require optimization to afford high frequency bi-allelic gene targeting without causing off-target mutations, which could potentially alter T cell function or predispose to cell transformation. Prevention of GVHD would require that virtually all T cells bear a disrupted TCR gene, requiring high-efficiency targeting and robust purging of unmodified cells to ensure T cell safety. Similar to the virus-specific T cell paradigm, it is unknown to what extent mature T cells undergoing extensive manipulation, including antigen-specific restimulation, TCR or CAR transduction, gene editing, and cell selection, will yield sufficiently large batches of functional T cells that meet the needs for multiple recipient infusions. Thus, allogeneic T cell approaches are still labor intensive and constrained by the limited replicative potential of mature T cells (Gattinoni et al., 2012).

Interestingly, allogeneic T cells may not cause GVHD in some particular circumstances. Thus, patients infused with T cells collected after allogeneic transplantation have not developed any GVHD-like syndrome (Davila et al., 2014a; Lee et al., 2015; Maude et al., 2014). More strikingly, CD19 CAR-targeted donor T cells show reduced GVHD potential in allogeneic recipient mice (reported at the American Society of Hematology annual meeting in 2012) and in human patients infused with CD19 CAR-modified donor leukocytes (Kochenderfer et al., 2013). A mechanistic explanation for these intriguing observations is still lacking.

Lymphoid Progenitor Therapy

While T cells can cause GVHD, their precursors do not, as they undergo positive and negative selection in the recipient's thymus. Taking advantage of this requires the ability to expand T cell precursors in culture, which is now possible due to advances in understanding T cell development (Awong et al., 2007; Rothenberg, 2011; Shah and Zúñiga-Pflücker, 2014). T cell precursors lack the ability to initiate GVH reactions because they complete their differentiation in the recipient's thymus wherein they become restricted to host MHC and yield T lymphocytes that are host tolerant (Zakrzewski et al., 2006). When transduced with a CAR, allogeneic lymphoid progenitors yield tumor-targeted T cells without causing GVHD (Zakrzewski et al., 2008). The main advantage of using T cell precursors for immunotherapy is that this approach does not require strict histocompatibility between

donors and recipients. In mice, this therapy works with unrelated fully mismatched cells just as well as with autologous cells. T cell precursor immunotherapy may therefore allow for a true “off-the-shelf” therapy, if lymphoid progenitor cell manufacturing can be scaled up.

Pluripotent Stem Cells as a Source of Therapeutic T Lymphocytes

The development of cellular therapeutics relying on functionally validated, banked, broadly histocompatible cell types would have a major impact on the applicability and cost of adoptive T cell therapies. This prospect raises the challenge of artificially generating ideal T cells rather than modifying those naturally formed. Pluripotent stem cells can give rise to a variety of somatic cells (Inoue et al., 2014; Murry and Keller, 2008; Takahashi et al., 2007) and thus have in principle the potential to serve as an endless supply of therapeutic T lymphocytes. A few reports support the feasibility of generating T lymphocytes from human ESCs and iPSCs in vitro (Kennedy et al., 2012; Nishimura et al., 2013; Themeli et al., 2013; Timmermans et al., 2009; Vizcardo et al., 2013).

Antigen Specificity

The first requirement for therapeutic function is specific antigen recognition, which is physiologically mediated by the TCR. ESCs and most iPSCs bear TCR α and β loci in the germline configuration. These undergo random rearrangements during lymphoid differentiation, thus generating polyclonal T cells of undetermined specificity and HLA restriction. This unpredictable repertoire severely limits the usefulness and potential for expansion and functional characterization of T cells derived from ESCs/iPSCs. Two approaches to dictate the specificity of iPSC-derived T cells have been hitherto reported. One utilizes iPSCs that bear rearranged TCR genes, providing a known antigen specificity (Nishimura et al., 2013; Vizcardo et al., 2013; Wakao et al., 2013). Re-differentiation of iPSCs derived from mucosal-associated invariant T (MAIT) cells expressing the invariant T cell receptor $V\alpha 7.2$ or established viral- and tumor-specific T cell clones gives rise to T lymphocytes bearing the same TCR as the parental T cell from which the iPSC clone was established, although re-arrangement of a remaining germline TCR α locus may result in multiple TCRs. This approach to afford antigen-specificity for cancer immunotherapy requires laborious cloning of antigen-specific T cells and the availability of the desired antigen-specific T cells for every prospective recipient. Another approach is to genetically transfer a receptor for antigen of known specificity. We previously demonstrated that T cell-derived iPSCs (TiPSCs) expressing a CAR (CAR-TiPSC) provide an effective means to concomitantly exploit the unlimited proliferative potential of iPSCs and direct the antigen specificity of iPSC-derived T cells (Themeli et al., 2013). In contrast to TCR transfer, CAR engineering yields T cells with unrestricted antigen recognition and enhanced potency owing to the costimulatory signals provided through the CAR.

Determining T Lineage Commitment and T Cell Fate

The functional properties of T lymphocytes depend not only on their differentiation stage and engineered features, as discussed above, but also on their lineage subtype ($\gamma\delta$ or $\alpha\beta$ T cells, effector or regulatory subsets) (Vantourout and Hayday, 2013). It is therefore essential to generate T cells of the desired functional subset. Natural human T lymphoid development

is outlined in Figure 1. Lymphopoietic progenitors seem to follow these steps overall throughout ESC/iPSC in vitro differentiation. TiPSCs generated from an $\alpha\beta$ -TCR-bearing T cell indeed give rise to $\alpha\beta$ -TCR⁺ cells. However, their phenotype, whether pre or post antigen-expansion, may not be that of a typical $\alpha\beta$ -T cell. Expanded TiPSC-derived T cells are CD3⁺CD7⁺CD5^{lo}TCR $\alpha\beta$ ⁺CD56⁺ and either double negative for CD4 and CD8 or CD8 α ⁺CD8 β ⁻ (Nishimura et al., 2013; Themeli et al., 2013). Antigen-activated and expanded CAR-TiPSC-T cells display an effector memory phenotype (CD45RA⁺CD27⁻CD28⁻CCR7⁻) (Themeli et al., 2013). Microarray gene expression analyses and detailed immunophenotypic profiling provided some clarification for these unexpected findings, establishing that the in vitro generated CAR-TiPSC-T cells possess an innate $\gamma\delta$ T cell-like profile, even though they express their endogenous $\alpha\beta$ -TCR (Themeli et al., 2013). Significantly, their in vivo anti-tumor function was comparable to natural, peripheral blood-derived $\gamma\delta$ T cells collected from the same donor and transduced with the same CAR (Themeli et al., 2013). Furthermore, Nishimura et al. (2013) observed the emergence of a few central memory TiPSC-derived T cells (CCR7⁺CD27⁺CD28⁺), although these cells had very low CCR7 and CD28 expression. In aggregate, these findings indicate that although the TiPSC-T cells express their rearranged endogenous $\alpha\beta$ -TCR on their surface, they acquire a phenotype and functional properties that do not correspond to that of natural naive or memory CD8 $\alpha\beta$ ⁺ T lymphocytes. One study showed efficient generation of CD4⁺CD8 α ⁺TCR⁺ cells, which almost exclusively differentiated into CD8 $\alpha\alpha$ ⁺ cells upon stimulation with antigen (Vizcardo et al., 2013). Other studies showed absent or minor generation of double positive CD4⁺CD8 α ⁺ cells, but no detection of CD8 β has yet been reported. A better understanding of the requirements for inducing CD4⁺CD8 $\alpha\beta$ ⁺ cells will pave the way for the generation of CD8 $\alpha\beta$ + and CD4⁺ T cells, including effector and regulatory T cells.

Interestingly, both CAR-TiPSC-T cells and regenerated MAIT cells (Wakao et al., 2013) express CD56 and CD161, suggesting that the innate nature of the TiPSC-T cells is independent of the identity of the initially reprogrammed T cell subtype. It is noteworthy that lineage diversion has been previously observed in transgenic TCR $\alpha\beta$ mice (Baldwin et al., 2005; Egawa et al., 2008; Terrence et al., 2000), wherein T cells distinct from wild-type natural killer (NK), NK-T, or CD4 or CD8 single-positive T cells displayed $\gamma\delta$ T cell features, including expression of CD8 α and low levels of CD5 (Terrence et al., 2000). Furthermore, in vitro differentiated T cells derived from TCR-engineered human CD34⁺ hematopoietic progenitors display an NK cell-like phenotype (Zhao et al., 2007). These observations suggest that the presence of rearranged TCR genes influences T cell fate, similar to reports in TCR transgenic mice (Baldwin et al., 2005). Accordingly, mature CD4⁺CD8⁺ and single-positive T cells developed from TCR-engineered CD34⁺ hematopoietic progenitors when the TCR cDNAs were introduced in the pre-T cell stage of differentiation (Snauwaert et al., 2014), consistent with time-dependent TCR expression causing lineage diversion. Alternatively, considering that some features of TiPSC-derived T cells, such as their CD8 α ⁺CD8 β ⁻ phenotype, expression of CD161 and low expression of CD5, are also found in innate-like T cells generated in fetal development (Cupedo et al., 2009; Spits and Cupedo, 2012), it may be that their innate character is imparted by a fetal-like hematopoietic stem cell intermediate committed to innate lymphopoiesis (Kennedy et

al., 2012; Mold et al., 2010; Yuan et al., 2012) and intrinsically skewed toward embryonic characteristics (Murry and Keller, 2008). Further investigation of the mechanisms underlying in vitro T lymphoid differentiation of TiPSCs is needed to better direct T cell subset differentiation and further shape the functional attributes of induced T cells.

T Cell Functionality

Beyond antigen specificity, two critical features that will determine the therapeutic relevance of pluripotent cell-derived T cells are their potential for in vivo persistence and sustained functionality. Few studies have comprehensively assessed the functional profile of ESC or iPSC-derived T cells in vitro or upon adoptive transfer in vivo. As previously mentioned, the random TCR rearrangements occurring in ESC/iPSC-derived T cells limit the feasibility of studying antigen-specific expansion and function. Therefore, ESC/iPSC-derived T cell function has been assessed only in in vitro assays showing IFN γ secretion after unspecific stimulation (Timmermans et al., 2009). Expanded tumor- and viral-specific TiPSC-derived T cells (100- to 1,000-fold) secrete IFN- γ after unspecific stimulation and lyse target cells in an antigen-specific manner in vitro (Nishimura et al., 2013). Re-differentiated MAIT cells were shown to successfully function in vivo against mycobacterium infection, although in a non-antigen-specific manner (Wakao et al., 2013). CAR-TiPSC T cells generated in culture expanded robustly upon CD19 engagement by the CAR (up to 1,000-fold over 3 weeks) and showed anti-tumor efficacy against a CD19⁺ lymphoma in a xenogeneic murine model, comparable to their natural counterparts harvested from peripheral blood (from the same donor) and transduced with the same CAR (Themeli et al., 2013). The latter study provided the proof of principle that human iPSC-derived T lymphocytes generated in vitro possessed anti-tumor function in vivo. Further studies are needed to investigate the in vitro expansion potential and the in vivo capabilities of iPSC-derived T lymphocytes.

Striking a Balance between Preventing Anti-Host Reactivity and Escaping Immune Rejection Is Key

Whereas anti-host reactivity may cause unacceptable toxicity, immune rejection of non-autologous T cells will curtail their efficacy. Therefore, escaping immune rejection, ensuring sufficient persistence and possibly long term engraftment, are further critical requirements to enable off-the-shelf adoptive T cell therapy. One immediate approach to solve this problem is to bank cells with common HLA haplotypes, as proposed for EBV-reactive T cells (Gallot et al., 2014; Leen et al., 2013) or iPSC/ESCs (Gourraud et al., 2012; Nakatsuji et al., 2008; Stacey et al., 2013; Turner et al., 2013). Although this approach is certainly a valuable first step toward broader applicability of adoptive T cell therapy, it is still constrained by HLA matching and by donor availability. Furthermore, the establishment of iPSC/ESC banks requires the generation of multiple iPSC lines from multiple donors, compounded by the eventual need to identify T cells of an appropriate specificity and HLA restriction, followed by extensive safety studies and validation of the individual clones. The alternative is to genetically target HLA genes to generate histocompatible cell products (Riolobos et al., 2013; Torikai et al., 2012). Targeting multiple HLA loci in primary T lymphocytes may be feasible but poses technical challenges owing to the substantial safety validation required for each cell product. HLA engineering and biosafety testing may be easier to perform in

pluripotent stem cells (Riolobos et al., 2013). In contrast to primary T cell manipulations, the genetic engineering of iPSCs results in fully modified clonal lines, which can be extensively evaluated (Papapetrou et al., 2011). However, disruption of HLA loci would expose cells to NK cell-mediated rejection. Further cell engineering including overexpression of HLA-E or HLA-G has been proposed as a solution to confer NK resistance (Riolobos et al., 2013; Torikai et al., 2012).

Perspectives for “Synthetic T Cells”

Stem cell reprogramming not only offers potential access to an unlimited source of therapeutic T lymphocytes, but it also provides an excellent platform for performing additional engineering intended to enhance the therapeutic value of induced T cells. The genetic engineering of TiPSCs with CARs is the first example of an efficient strategy to concomitantly harness the unlimited availability of iPSCs and direct the specificity and functional potential of iPSC-derived T cells (Themeli et al., 2013). The use of iPSCs further opens up new perspectives for the generation of histocompatible, off-the-shelf T cells that could eventually be administered to multiple recipients. The combination of iPSC technology and immune engineering may thus provide an opportunity to generate T cells that uniquely combine favorable attributes including antigen specificity, lack of alloreactivity, enhanced functional properties and histocompatibility (Figure 4). Several challenges remain, including the ability to control T lineage specification (to $\alpha\beta$ - or $\gamma\delta$ -T cells, NK-T, CD8, CD4, or regulatory T cells), differentiation to an optimal maturation stage (e.g., naive or stem central memory T cells) (Gattinoni et al., 2012), and acquisition of an optimal functional and proliferative potential (Sadelain et al., 2003). Natural, autologous T cells represent the best-defined cell source for adoptive cell therapy today, and they are the cornerstone of present cell-based cancer immunotherapy. Induced, engineered T cells derived from allogeneic pluripotent stem cell sources may play an important role in the future.

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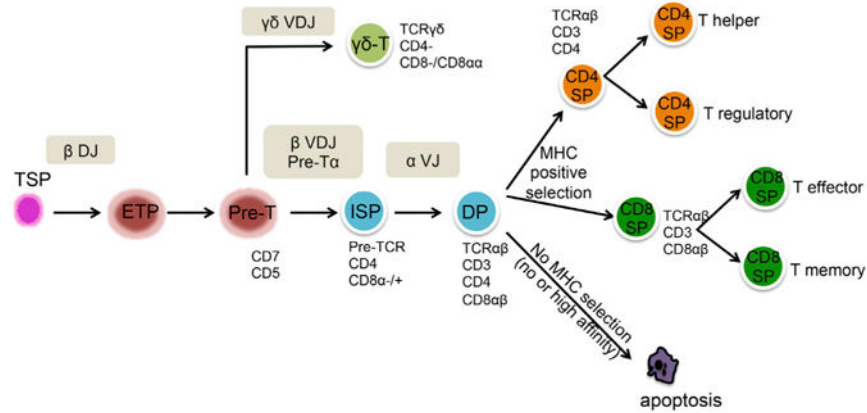


Figure 1. Human T Lymphocyte Development

Hematopoietic stem cell-derived thymus-seeding progenitors (TSPs) migrate into the thymus and differentiate into an Early Thymic Progenitor (ETP) upon rearrangement of the diversity (D) and joining (J) regions of the TCR β locus. ETPs progress to a pre-T cell state expressing CD1a and CD5. At this stage, recombination of the variable (V) region of the TCR β locus to form a complete rearranged VDJ TCR β locus occurs almost simultaneously with the rearrangement of the gene segments encoding the $\gamma\delta$ TCR. Depending on the outcome of the TCR segment rearrangements, the cells can then follow an $\alpha\beta$ or a $\gamma\delta$ differentiation path. A successful TCR β rearrangement leads to the process of β -selection and emergence of a CD4⁺ immature single positive (ISP) T cell. The CD4⁺ ISP cell then develops into a double-positive (DP) cell that expresses both CD4 and CD8 and has begun to rearrange the V and J regions of the TCR α locus. The life span of DP thymocytes is limited as they quickly proceed to apoptosis if they do not receive a TCR-mediated survival signal provided by the self-HLA molecules of the thymic epithelium before maturing into CD4⁺CD8⁻ and CD4⁻CD8⁺ single-positive (SP) T cells.

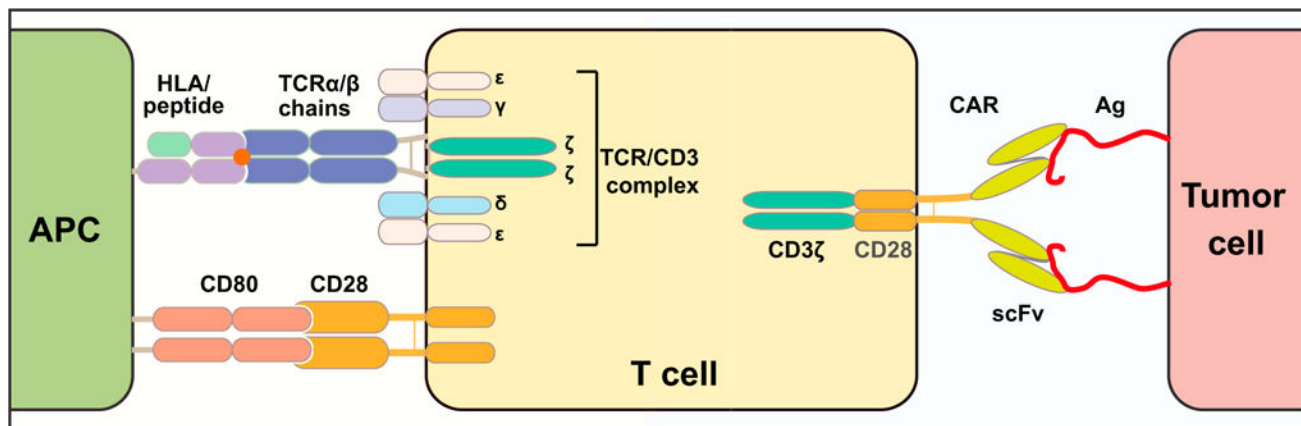


Figure 2. Antigen Receptors Used for T Cell Engineering

Left: Structure of the $\alpha\beta$ heterodimeric T cell receptor (TCR) associated with the multi-chain CD3 complex (γ , δ , ϵ , and ζ) and flanked by CD28, a costimulatory receptor constitutively expressed in most $\alpha\beta$ -T cells. Right: Structure of a prototypical second-generation chimeric antigen receptor (CAR), comprising an scFv for antigen recognition, the CD3 ζ cytoplasmic domain for T cell activation, and the CD28 cytoplasmic domain to enhance T cell function and persistence (Maher et al., 2002).

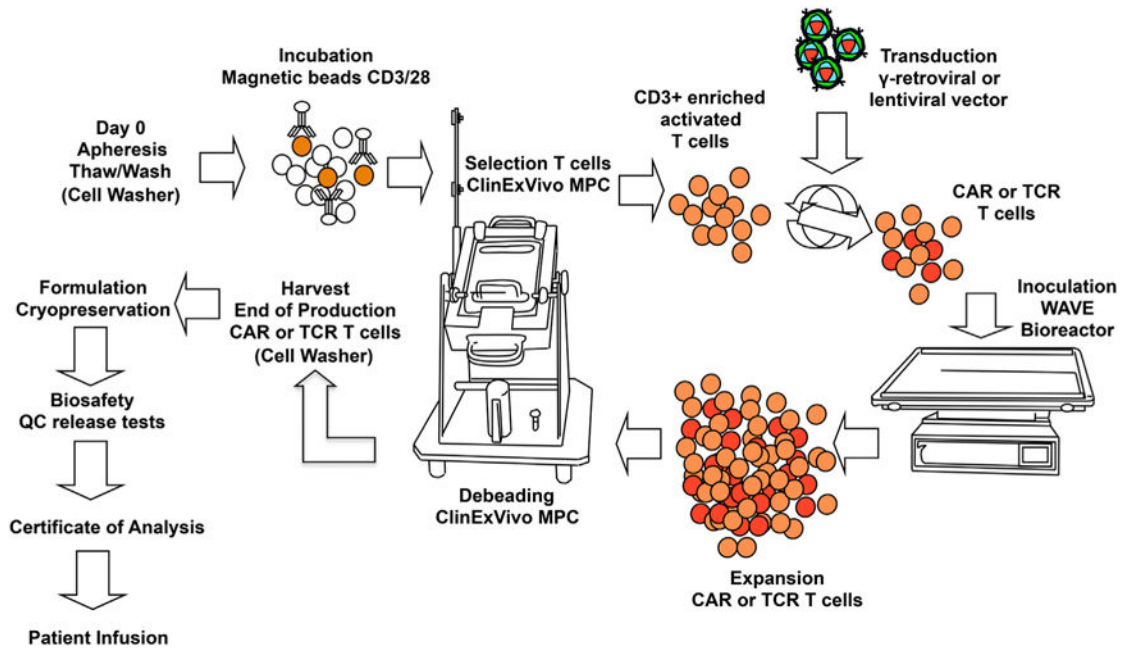


Figure 3. General Schema of Autologous T Cell Manufacturing

The semi-closed system relies on the use of a cell washer to wash the apheresis product before freezing and after thawing, the capture of CD3+CD28+ T cells with magnetic beads or microbeads. For magnetic beads subsequent selection is performed on the ClinExVivo magnetic particle concentrator (MPC). Thereafter, T cells are transduced with a viral vector expressing a TCR or a CAR, expanded in a bioreactor (e.g., Wave Bioreactor), debeaded on the ClinExVivo MPC, and formulated. Adapted from Hollyman et al. (2009).

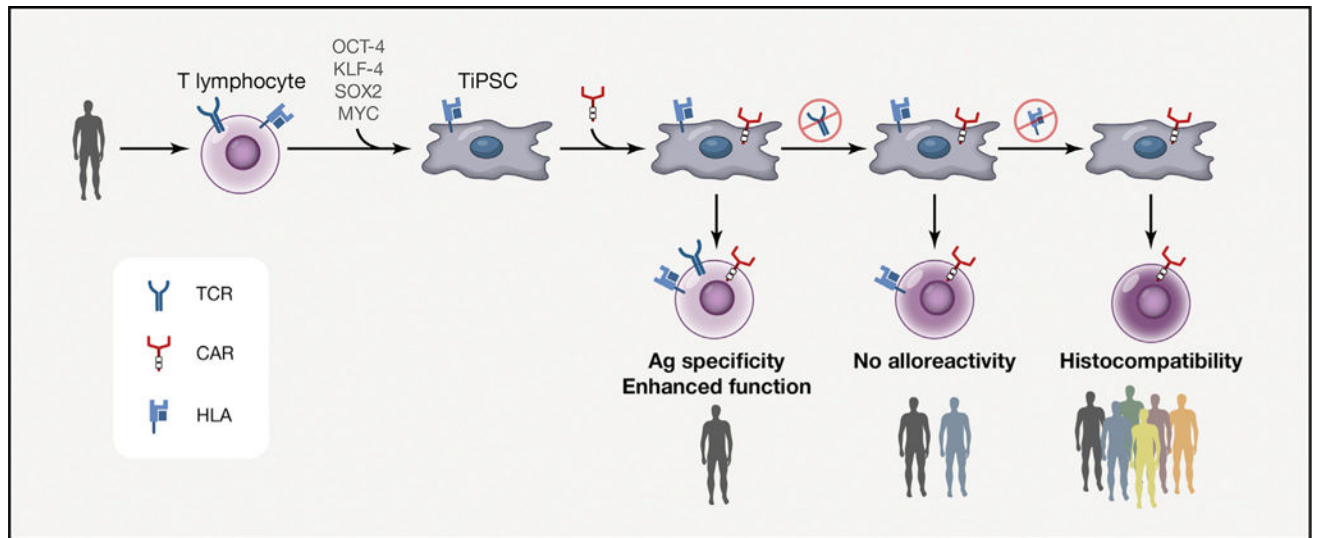


Figure 4. Perspectives for Synthetic TiPSC-Derived T Cell Generation

TiPSCs can be generated from one donor and utilized for the production of synthetic T lymphocytes engineered to possess optimized properties such as (1) antigen specificity through a CAR or TCR, (2) enhanced function (e.g., introducing costimulatory molecules), (3) elimination of alloreactivity and partial broadening of applicability (e.g., strategies to knock out the TCR expression without perturbing the T lymphoid differentiation process), and (4) broad histocompatibility by selection or genetic modification of the HLA loci.