



Review

Cancer immunotherapy utilizing gene-modified T cells: From the bench to the clinic[☆]

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ABSTRACT

The immune system plays a critical role in the elimination and suppression of pathogens. Although the endogenous immune system is capable of immune surveillance resulting in the elimination of cancer cells, tumor cells have developed a variety of mechanisms to escape immune recognition often resulting in tumor outgrowth. The presence of immune infiltrate in tumors has been correlated with a good prognosis following treatment (Sato et al., 2005; Loi et al., 2013; Clemente et al., 1996; Galon et al., 2006). As such, immune cells such as T cells, have been harnessed in order to target cancer. Tumor reactive lymphocytes, called tumor-infiltrating lymphocytes (TILs) have been isolated and expanded from the tumor and reinfused back into patients for the treatment of melanoma. The promise of adoptive immunotherapy utilizing TILs as a robust treatment for cancer has been highlighted in patients with advanced melanoma with greater than 50% of patients responding to treatment (Dudley et al., 2005). Although TIL therapy has shown promising results in melanoma patients, it has proved difficult to translate this approach to other cancers, given that the numbers of TILs that can be isolated are generally low. To broaden this therapy for other cancers, T cells have been genetically modified to endow them with tumor reactivity using either a T cell receptor (TCR) (Parkhurst et al., 2009, 2011; Chinnasamy et al., 2011) or a chimeric antigen receptor (CAR) (Grupp et al., 2013; Park et al., 2007). This review will outline the origins and development of adoptive immunotherapy utilizing TILs leading to genetic modification strategies to redirect T cells to cancer. Potential hurdles and novel strategies will be discussed for realizing the full potential of adoptive immunotherapy becoming a standard of care treatment for cancer.

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1. Introduction

The maintenance of a healthy host is largely dependent on the ability for the immune system to identify, suppress and eliminate pathogens. Although the endogenous immune system is rarely capable of eliminating established cancer, the presence of immune infiltrate in tumors has been correlated with good prognosis following therapy (Sato et al., 2005; Loi et al., 2013; Clemente et al.,

1996; Galon et al., 2006). Therefore, components of the immune system such as T cells have been utilized in order to target cancer. This has involved isolating and expanding tumor reactive lymphocytes from the tumor, called tumor-infiltrating lymphocytes (TILs). The promise of adoptive immunotherapy utilizing TILs as a robust treatment for cancer has been recently highlighted in patients with advanced melanoma, with greater than 50% of patients responding to treatment (Dudley et al., 2005). However, TILs are challenging to isolate from cancers other than melanoma. To overcome this problem, T cells have been genetically engineered endowing them with tumor reactivity. This has involved modifying T cells with $\alpha\beta$ T cell receptor (TCR) transgenes (Parkhurst et al., 2009, 2011; Chinnasamy et al., 2011). TCR modified T cells have had some promising results in clinical trials in melanoma patients (Morgan et al., 2006). However, tumors are capable of escaping detection by the immune system by downregulating molecules

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such as MHC1. Therefore, recombinant receptors called chimeric antigen receptor (CAR) (Grupp et al., 2013; Park et al., 2007) have been developed, arming the T cell with anti-tumor activity regardless of MHC1 expression. The tremendous promises of the approach have been recently highlighted utilizing CAR T cells specific for CD19, which has resulted in remarkable responses in hematological malignancies (Maude et al., 2014). In this review, we outline the origins and development of adoptive immunotherapy utilizing TILs, TCR modified T cells and CAR modified T cells. Finally we discuss potential hurdles and novel strategies for the potential of adoptive immunotherapy becoming a standard of care treatment for cancer.

2. Adoptive immunotherapy

2.1. Early preclinical studies and clinical trials

It has long been proposed that the presence of a lymphocytic infiltration within tumors is associated with a good prognosis. In 1972, it was reported that a gastric cancer patient with hepatic metastases had a total regression of metastases in his liver in the absence of therapy (Rosenberg et al., 1972). The dense infiltration of lymphocytes in the resected gastric biopsy suggested the importance of these TILs against the growing cancer (Rosenberg et al., 1972). The presence of TILs have since been associated with a favorable prognosis in other cancers including ovarian (Sato et al., 2005), breast (Loi et al., 2013), melanoma (Clemente et al., 1996) and colon cancer (Galon et al., 2006).

Whilst TILs are capable of recognizing tumor associated antigens through their endogenous TCRs, the small number of TILs isolated from patient tumors without expansion are insufficient for clinical intervention. The discovery of interleukin-2 (IL-2) (Smith et al., 1980a, 1980b) allowed for the *ex vivo* expansion of TILs isolated from patients, with early clinical trials initiated by the pioneers of adoptive immunotherapy, Rosenberg et al. (Rosenberg et al., 1988a; Topalian et al., 1988; Lotze and Rosenberg, 1986) at the National Institutes of Health. The first clinical trial of adoptive immunotherapy involved the administration of IL-2 expanded TILs from 12 patients with advanced renal or breast cancer, colon carcinoma or melanoma. A single dose of cyclophosphamide was used to precondition the patients who were infused with 8×10^9 – 2.3×10^{11} cells and varying doses of IL-2. Partial tumor regression was reported in one patient with renal carcinoma, one patient with breast cancer and one patient with melanoma (Topalian et al., 1988). These encouraging results led to further trials to assess the combination of lymphodepleting regimens and the use of IL-2 for enhancing the efficacy of adoptive T cell therapy.

Early preclinical and clinical studies for metastatic melanoma were pioneered by Rosenberg et al. (Rosenberg et al., 1988a; Lotze and Rosenberg, 1986). Tumor reactive TILs were cultured with fresh melanoma samples or melanoma cell lines, and assessed for their ability to secrete cytokines and lyse target cells *in vitro* (Topalian et al., 1987). Following selection for antigen recognition, these TILs were then activated and expanded *ex vivo* in the absence of an immunosuppressive tumor microenvironment. Patients also received high doses of IL-2 to support the activation and proliferation of infused T cells (Forni et al., 1986; Testa et al., 1990). In a preliminary clinical trial with metastatic melanoma, Rosenberg et al. found that 11/20 of patients treated with this regimen had objective regression of their tumors in various sites of the body that lasted from 2 to greater than 13 months (Rosenberg et al., 1988a). The protocol used in this trial became the foundation of future clinical trials.

In these early trials, although some patients experienced long lasting regression, the majority of anti-tumor responses were short-lived. To determine the persistence and distribution of the

infused TILs in patients, Rosenberg et al. retrovirally transduced TILs to express the neomycin resistance gene, which allowed the cells to be tracked following infusion (Rosenberg et al., 1990). Using polymerase chain reaction (PCR) analysis, they reported minimal numbers of TILs in the patients' peripheral blood and tumor biopsy and that these numbers decreased significantly after 3 weeks. The relatively short life span and poor infiltration of the infused TILs into the tumor bed warranted further investigation to enhance the tumoricidal capacity of the transferred T cells.

2.2. Lymphodepletion in adoptive immunotherapy

Murine models have demonstrated that lymphodepletion using nonmyeloablative sublethal total body irradiation (TBI) prior to TIL infusion greatly augmented the efficacy of adoptive immunotherapy (Wrzesinski et al., 2010; Gattinoni et al., 2005a). The mechanisms as to why lymphodepletion is beneficial prior to the infusion of T cells are still not completely understood. It has been shown that there is an increase in freely available homeostatic cytokines such as IL-2, IL-7 and IL-15 in the absence of cellular 'sinks', a phenomenon where endogenous lymphocytes such as T cells and natural killer cells (NKs) compete with transferred T cells for cytokine support (Wrzesinski et al., 2010; Gattinoni et al., 2005a; Klebanoff et al., 2005a). Elimination of immunosuppressive cells such as myeloid derived suppressor cells (MDSCs) and T regulatory cells (T_{regs}) and their secreted inhibitory cytokines such as IL-10 may also play a role in this enhanced anti-tumor effect (Gajewski et al., 2013; Gabrilovich et al., 2012). It is also believed that damage to the mucosal surfaces releases toll like receptor (TLR) agonists, inducing dendritic cells (DCs) to mature and stimulate expansion of the activated infused T cells (Kieper et al., 2005; Hill et al., 1997). Therefore, it appears that lymphodepletion using irradiation and chemotherapy are able to skew the tumor microenvironment, depleting pro-tumoral elements and subsequently allowing for a beneficial shift in the ratio of infused T cells to suppressive cells.

These preclinical findings prompted the investigation of the preconditioning regimen including TBI and lymphodepleting prior to adoptive transfer, which results in the temporary elimination of host immune cells from the patient. Several clinical trials treating patients with metastatic melanoma utilized increasing doses of lymphodepletion, achieving objective response rates of 49–72% (Dudley et al., 2005; Rosenberg et al., 2008, 2011) (the response evaluation criteria in solid tumors (RECIST) was used in these studies). In a report consisting of three sequential clinical trials of 93 patients with measurable metastatic melanoma, the combination of chemotherapy, TBI and adoptive immunotherapy led to the regression of melanoma metastasis in the lung, liver, brain, bone, lymph nodes and subcutaneous tissue, which resulted in objective response rates of 72%, with some complete responses reported for more than 3 years at the time of publication (Rosenberg et al., 2011). The results were more remarkable given that some tumors were rendered inoperable due to their size and location.

2.3. Optimal phenotype of infused T cells

While the *ex vivo* culture conditions for T cell expansion favor a more effector phenotype, several studies have focused on investigating which phenotype of infused T cells will thrive and induce the greatest anti-tumor effect following infusion. Early pre-clinical and clinical protocols for the generation of TILs involved the activation and expansion of T cells into large numbers *in vitro* with IL-2. T cells with attributes such as interferon- γ (IFN- γ) secretion (Barth et al., 1991) and cytotoxicity toward tumor cells (Aebbersold et al., 1991; Schwartzentruber et al., 1994) were isolated and infused into patients. These T cells were highly activated and had an effector

phenotype, however more recent investigations indicate that additional parameters such as TIL differentiation status and memory phenotypes are important determinants in anti-tumor activity and persistence following transfer (Klebanoff et al., 2005b; Gattinoni et al., 2005b; Wolfl et al., 2011).

Data emerging from mouse models demonstrated that less differentiated TILs have superior *in vivo* antitumor properties. Gattinoni et al. (2005b) performed studies where B16 melanoma tumor-bearing mice were treated with active immunization, IL-2 and T cells in varying differentiation states. They reported that despite mature effector T cells (T_{EFF}) demonstrating a greater capacity for IFN- γ secretion and cytotoxicity *in vitro*, the differentiation stages of T_{EFF} cells were inversely related to their anti-tumor activity and proliferative capacity *in vivo*. In addition to the maturation status, emerging evidence also demonstrated that T cells with memory properties are superior candidates for adoptive immune therapy. Based on their phenotypic markers, function and migratory properties, memory T cells can be categorized into two populations, central memory (T_{CM}) and effector memory (T_{EM}) cells (Kaech et al., 2002). Gene expression analysis of these tumor-reactive CD8⁺ T cell memory subsets have showed that T_{CM} cells have lower expression of pro-apoptotic signaling genes, Bid, Bnip3 and Bad and higher expression of genes associated with trafficking to secondary lymphoid organs, that include CD62L, CXCR3, CCR7 compared to T_{EM} cells (Klebanoff et al., 2005b). Tumor-reactive T_{CM} cells also mediated greater tumor regression and survival of B16 melanoma-bearing mice compared to T_{EM} cells, indicating that T cells with a T_{CM} phenotype are superior over T_{EM} cells for adoptive immunotherapy.

There has also been increasing interest in a population of CD8⁺ T memory stem (T_{SCM}) cells. T_{SCM} cells were first identified in a model of graft vs host disease (GVHD) (Zhang et al., 2005), as self-renewing T cells capable of generating T_{CM} , T_{EM} and T_{EFF} cells, expressing CD27 and lymphoid homing molecules such as CCR7 and CD62L (Gattinoni et al., 2011). These properties make T_{SCM} cells suited for use in adoptive immunotherapy as they are able to mediate long-term immunity and generate various T cell phenotypes. A comparison of T_{EM} , T_{CM} and T_{SCM} cells demonstrated that T_{SCM} cells were superior to T_{EM} and T_{CM} cells in the ability to mediate tumor regression in a mouse model (Klebanoff et al., 2011).

In addition to the differentiation status and memory phenotype, it is also believed that shorter *ex vivo* culture processing is beneficial as prolonged cell culture conditions can result in cells reaching replicative senescence. Hinrichs et al. reported that T_{EFF} cells derived from naïve CD8⁺ T cells exhibited greater anti-tumor immunity compared to the T cells derived from T_{CM} cells. The T_{EFF} cells derived from naïve T cells showed low levels of KLRG-1 expression, a marker for replicative senescence and cell terminal differentiation (Hinrichs et al., 2009). This study implies that naïve rather than T_{CM} cells may be the important source of T_{EFF} cells and prevention of terminal differentiation is beneficial in the generation of T_{EFF} cells for adoptive immunotherapy of cancer. A lack of telomerase activity and shortening of telomere lengths have also been correlated with poor persistence of infused TIL in patients with metastatic melanoma (Shen et al., 2007). In human CD8⁺ T cells, telomeres are longer in naïve T cells than memory T cell subsets (Weng et al., 1995) and this may account for the improved persistence of TILs generated from naïve T cells. In clinical trials performed by Rosenberg et al., the number and percent of CD8⁺ CD27⁺ T cells in cultures as well as the length of telomere repeats in chromosomes corresponded closely with improved patient outcomes (Rosenberg et al., 2011).

It is possible that the suboptimal results of several early clinical trials were due to the extended culturing process (Topalian et al., 1987; Rosenberg et al., 1988b). Investigations into the ideal phenotype of infused T cells by several groups led to clinical trials utilizing

'young' or short-term cultured TILs (Besser et al., 2010; Dudley et al., 2010). The benefits of the protocol to generate younger TILs, included a much shorter culture period which is more cost effective and less labor intensive. Other advantages of the shorter protocol include the generation of T cells with higher levels of CD27 and longer telomeres (Tran et al., 2008) and increased survival of CD4⁺ T cells, which may also contribute to tumor immunity (Bevan, 2004; Ossendorp et al., 1998), although this requires further investigation.

In a Phase II clinical trial using short-term cultured TILs in a cohort of 20 metastatic melanoma patients, Besser et al. (2010) reported that non-selected 'young' TILs that had been expanded for 14 days could mediate tumor regression in 50% of recipients. In another trial, Dudley et al. (2010) compared two different lymphodepletion protocols in combination with 'young' TIL therapy. Objective responses were observed in 58% and 48% of patients treated with young TILs and non-myeloablative conditioning or TBI respectively. As a result of these preclinical and clinical studies, more naïve and less differentiated T cells appear to be better suited to adoptive immunotherapy. Phase III clinical trials comparing 'young' TIL and existing TIL protocols need to be initiated in order to determine which populations are best for the treatment of cancer.

3. Genetic modification of T cells

Due to the success of adoptive immunotherapy for the treatment of metastatic melanoma, numerous attempts have been made to isolate and expand tumor reactive TILs in other malignancies, including renal cell carcinoma (RCC), ovarian, colorectal, and breast cancer (Baxeianis et al., 1994; Bouet-Toussaint et al., 2000; Freedman et al., 1994; Yannelli et al., 1996; Mulder et al., 1995). Figlin et al. (1997) reported that a clinical trial combining nephrectomy, IL-2 and CD8⁺ TILs for the treatment of metastatic RCC resulted in moderate clinical benefit, where 9.1% and 25.5% of patients had a complete response and partial response respectively. However, a subsequent Phase III trial with the same protocol showed no clinical benefit (Figlin et al., 1999). The poor outcome of the adoptive immunotherapy trials in solid cancers other than melanoma may be due to several factors including the reduced immunogenicity of tumor-associated antigens (TAAs) in cancers other than melanoma. In addition, exomic sequencing data indicated that melanomas contain more mutations than other type of solid cancers (Walia et al., 2012). Therefore, it is likely that T cells recognizing these neo-antigens have not been deleted by the thymus.

Nevertheless, the poor clinical response in other solid cancer settings and some inherent limitations of TIL therapy, such as the need to perform invasive procedures to obtain TILs and the inability to grow TILs from some patients, has led to the development of alternative approaches to generate tumor reactive T cells. Development in genetic modification techniques has resulted in the ability to engineer T cells to endow them with anti-tumor activity, using TCRs or CARs specific for a range of TAAs. To date, the use of gene-modified T cells has been investigated for the treatment of several types of cancer including ovarian cancer, leukemia, lymphoma and RCC (Kershaw et al., 1996, 2006; Lamers et al., 2006; Till et al., 2008). The types of T cells used in adoptive immunotherapy are shown in Fig. 1.

3.1. Preclinical studies using TCR modified T cells

TCR gene therapy involves the transfer of tumor-specific TCR $\alpha\beta$ genes into autologous T cells, which are then expanded *ex vivo* and re-infused back into the patient. The choice of TCR is dependent on the target antigen and human leukocyte antigen (HLA) alleles expressed by the patient. TCR gene therapy has been

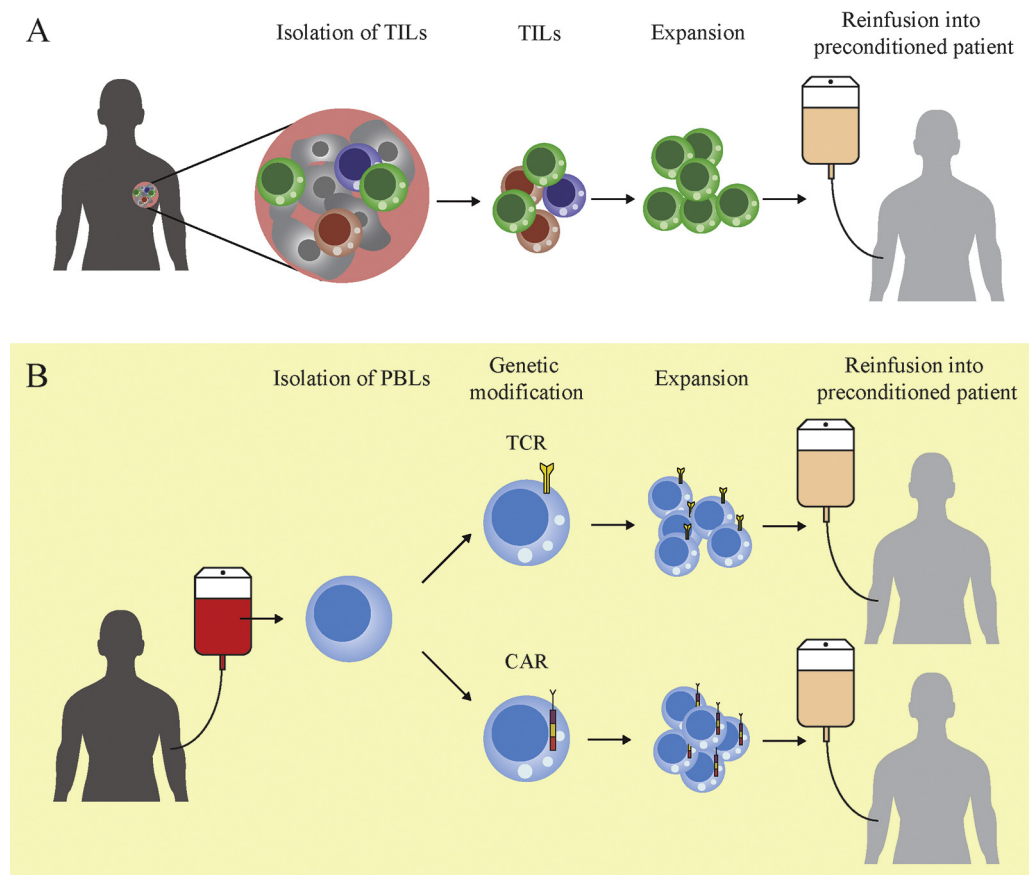


Figure 1. Treating patients with TILs or genetically modified tumor-reactive T cells.

In melanoma patients treated with TILs (**A**), tumors are surgically resected and the TILs are isolated and expanded before reinfusion back into the patient, who may undergo pre-conditioning with chemotherapy and TBI. In order to generate tumor-specific T cells for targeting other malignancies, (**B**), T cells from patient peripheral blood are isolated and expanded in culture and genetically modified to express either a TCR with specificity for tumor peptide presented on MHC or express a CAR containing a scFv extracellular domain with specificity for TAA linked to transmembrane and cytoplasmic T cell signaling domains. Depicted is a third generation CAR containing the costimulatory domains CD28 and 4-1BB linked to the CD3 ζ chain. These *ex vivo* expanded T cells are reinfused back into pre-conditioned patients. PBL, peripheral blood lymphocyte; TBI, total body irradiation.

investigated for the treatment of cancer and viral-induced malignancies (Gehring et al., 2011; Scholten et al., 2011; Jurgens et al., 2006).

Early preclinical studies utilizing TCR modified T cells focused on the treatment of melanoma, using TCR specific for antigens including melanoma antigen recognized by T cells-1 (MART1), tyrosinase and gp100. Although efforts have been made to isolate TCRs for TAAs in other cancers, tumor specific TILs have proved difficult to isolate and expand, therefore TCRs specific for TAAs in other cancers have been generated using alternative techniques (Parkhurst et al., 2009; Stanislowski et al., 2001). One such technique involved immunizing transgenic mice such as HLA-A2.1 mice with human peptides, resulting in the expansion of murine CD8⁺ T cell clones with peptide specific reactivity. Subsequently, the TCR $\alpha\beta$ chains were cloned into viral vectors and used to transduce TILs. This technique was first used to isolate p53 reactive TCRs (Stanislowski et al., 2001). Another method to enhance the affinity of isolated TCRs is to use phage display techniques. This involves the display of millions of TCR sequences on bacteriophage coat proteins and the identification of tumor reactive TCRs which bind to tumor antigen peptide (Li et al., 2005). These approaches have been successfully used to isolate and transduce T cells with TCRs specific for multiple antigens including Wilms tumor 1 (WT1) (Xue et al., 2005), carcinoembryonic antigen (CEA) (Parkhurst et al., 2009) and murine double-minute 2 (MDM2) (Stanislowski et al., 2001). The use of TCRs isolated from patients or generated using other techniques has been evaluated in

clinical trials for the treatment of cancers including melanoma and colorectal cancer (Robbins et al., 2011; Parkhurst et al., 2011).

3.2. Clinical trials using TCR modified T cells

TCR gene therapy was first investigated for the treatment of patients with metastatic melanoma. In an encouraging pre-clinical study, TCR $\alpha\beta$ genes were isolated from a tumor reactive cytotoxic T lymphocyte (CTL) clone specific for the TAA MART-1 (Cole et al., 1995). Transduction of human peripheral blood lymphocytes (PBL) with the TCR α and β chains was shown to produce CTL with anti-tumor reactivity *in vitro* and could potentially treat patients with metastatic melanoma (Clay et al., 1999). Subsequently, the first clinical trial using genetically engineered MART-1 specific TCRs involved 15 metastatic melanoma patients (Morgan et al., 2006). In this trial, two patients experienced objective responses and modified cells were detected 1 year after treatment. This was the first demonstration of clinical efficacy due to TCR modified T cells in metastatic melanoma. Another clinical trial by the same group (Johnson et al., 2009) utilized higher affinity MART-1 specific T cells and reported objective responses in 30% and 19% of patients respectively. However, investigators also reported the targeting of normal melanocytes in the skin, eye and ear in some patients. This was the first report of “on-target/off-tumor” autoimmunity from TCR modified T cells. Other melanoma target differentiation antigens studied include tyrosinase and gp100, also derived from patients that had

been successfully treated with TIL therapy (Kawakami et al., 1995; Kawakami and Rosenberg, 1997).

Cancer-testis antigens (CTAs) are good candidates for TCR gene therapy as CTAs are expressed by some tumors and are restricted to male germ cells in testis but not in adult somatic tissues. Robbins et al. (2011) utilized TCR modified PBLs targeting the CTANY-ESO-1 that is expressed on a range of epithelial cancers including breast, lung, prostate and ovarian (Barrow et al., 2006; Chen et al., 1997; Goydos et al., 2001; Kienstra et al., 2003). Response rates of 45% and 67% were reported in patients with melanoma and synovial cell sarcoma, respectively. No antigen specific toxicity was observed. This study highlighted the potential of using CTAs as targets in adoptive immunotherapy. Table 1 provides a summary of some of the clinical trials using TCR modified T cells for the treatment of cancer.

3.3. Limitations associated with TCR modified T cells

There are several limitations for using TCR modified T cells for adoptive immunotherapy. One of the main hurdles is that the effect is dependent on HLA/MHC presentation. However, a variety of cancers downregulate their HLA/MHCI (Bubenik, 2004) as a method of immune escape. In addition, TCR mispairing has also proven to be an issue. In preclinical mouse models, the combination of endogenous and transduced TCR has resulted in TCR mispairing, with the formation of potentially auto-reactive TCRs (Bendle et al., 2010). Strategies to reduce the incidence of mispairing have included the introduction of cysteine bonds, leucine zipper motifs and human-murine hybrid TCRs in order to stabilize TCR chains (Chang et al., 1994; Cohen et al., 2006, 2007; Kuball et al., 2007). Furthermore, the safety of retroviral transduction and its potential to insert and enhance dormant oncogenes, on-target/off-tumor toxicity due to infused T cells targeting antigen on normal tissues are also of concern (Parkhurst et al., 2011; Johnson et al., 2009; Linette et al., 2013; Morgan et al., 2013). Strategies to address these concerns are discussed in the latter part of the review, in Section 4.2.

3.4. The development of chimeric antigen receptors

In the past few decades, a novel form of adoptive immunotherapy involving the generation of modified CTL expressing CARs has been developed, which has broadened the application of adoptive immunotherapy to cancers other than melanoma. CARs are generally composed of the single-chain variable fragment (scFv) specific for a TAA (Wang et al., 2010; Kalos et al., 2011). The scFv is linked via hinge and transmembrane domains to intracellular cytoplasmic signaling domains, which are capable of triggering T cell activation. Thus, CARs do not require MHC presentation and provide an alternative to TCR gene therapies. Eshhar et al. (1993) first reported the use of CAR in T cells which had been endowed with the extracellular scFv specific for the hapten 2,4,6-trinitrophenyl (TNP) linked to either the intracellular FcR γ or CD3 ζ chain. T cells modified with this CAR were shown to mediate specific IL-2 secretion and induce target cell lysis. Since then, CAR design has progressed from the initial first generation CARs (with FcR γ or CD3 ζ chains) to second generation CARs (incorporating additional cytoplasmic co-stimulatory domains such as CD28 and CD137 (Kalos et al., 2011; Moeller et al., 2005) and subsequently third generation CARs (comprising three signaling domains) (Zhong et al., 2010; Zhao et al., 2009; Kershaw et al., 2013). However, it remains unclear what number and combination of signaling domains are necessary for eliciting maximal anti-tumor responses.

The use of CARs confer several advantages over TCR transgenes. In addition to the recognition of antigen independent of the expression of MHC, CARs are able to bypass some of the mechanisms that tumors employ to evade immune detection. For example, CARs contain intracellular signaling domains that are designed

to compensate the down-regulation of co-stimulatory molecules by cancer. Furthermore, CARs are not only able to target protein antigens but also carbohydrate (Westwood et al., 2005, 2009; Mezzanzanica et al., 1998; Craddock et al., 2010) and lipid antigens as well as any antigens that can be recognized by an antibody.

Nevertheless, CARs are only able to detect antigens on the cell surface and if tumors lose these antigens, they can escape CAR detection. Furthermore, current CARs incorporate only a fraction of the signaling molecules normally associated with TCR-mediated signaling, and as such CAR-mediated responses still fall short of achieving optimal activity from T cells, as illustrated by the robust TCR-mediated responses against pathogens (van Lier et al., 2003). Therefore studies are currently being performed to determine which combination of signaling molecules will provide optimal activation of a T cell.

3.5. Clinical trials utilizing chimeric antigen receptors

Due to promising preclinical results, clinical trials have been performed to evaluate the efficacy of different CARs against a range of cancers and these trials have generated remarkable responses in patients with hematological malignancies (Maude et al., 2014; Gill et al., 2014; Brentjens et al., 2013). A summary of these trials is shown in Table 2. First generation CARs containing the signaling domain CD3 ζ or FcR γ used for the treatment of various cancers including ovarian, colorectal, renal cell carcinoma and neuroblastoma have generated mixed results (Park et al., 2007; Kershaw et al., 2006; Warren et al., 1998; Louis et al., 2011; Lamers et al., 2013). The first study to demonstrate complete responses in patients treated with CAR modified T cells in solid cancer involved the treatment of 3 neuroblastoma patients with the first generation GD2 specific CAR containing the CD3 ζ signaling domain (Park et al., 2007; Louis et al., 2011). However, other trials utilizing first generation CARs have resulted in only modest anti-tumor activity (Kershaw et al., 2006; Lamers et al., 2011), therefore CARs with additional signaling domains were tested subsequently.

Clinical trials utilizing second generation CARs for the treatment of hematological malignancies have resulted in the most encouraging therapeutic responses (Grupp et al., 2013; Kalos et al., 2011; Brentjens et al., 2011, 2013; Porter et al., 2011; Kochenderfer et al., 2010). Kalos et al. (2011) utilized a CD19 specific CAR comprising the domains CD137 and CD3 ζ to treat patients with chronic lymphocytic leukemia (CLL) and acute lymphoblastic leukemia (ALL). Two out of the three CLL patients experienced complete responses, whilst the third patient experienced a partial response for 8 months. This landmark study was the first to demonstrate the curative potential of CAR T cells. This study also reported that infused T cells were able to expand >1000 fold, infiltrate the bone marrow and remained detectable 6 months post-infusion. Furthermore, each infused T cell was calculated to be responsible for the elimination of more than 1000 CLL cells, indicating the extensive cytotoxic capabilities of CAR modified T cells. In the ALL study, both patients experienced molecular remission, however one patient relapsed with CD19 negative disease (Grupp et al., 2013). Brentjens' et al. also utilized a second generation CD19 specific CAR, with the signaling domains CD28–CD3 ζ for the treatment of ALL. They achieved 88% (14 patients) complete responses out of the 16 patients enrolled in the trial (Davila et al., 2014). However, clinical trials targeting the CD19 antigen have reported on-target/off-tumor toxicity where patients experienced B cell aplasia, due to the targeting of CD19 on B cells (Kochenderfer et al., 2010).

The use of third generation CAR T cells in clinical trials has been limited to trials for lymphoma (Till et al., 2012) and colon cancer (Morgan et al., 2010). Till et al. (2012) treated lymphoma patients with a CD20 CAR composed of CD3 ζ , CD28 and CD137

Table 1

Summary of clinical trials utilizing TCR modified T cells for adoptive immunotherapy.

Target antigen	Cancer	Results (number of patients treated in trial)	Reference
gp100	Melanoma	One complete response and two partial responses (16)	Johnson et al. (2009)
MART-1/Melan-A	Melanoma	One partial response (15) Four objective responses (31)	Duval et al. (2006) Morgan et al. (2006) , Burns et al. (2009)
p53	Melanoma	Six partial responses (20)	Johnson et al. (2009)
NY-ESO-1	Melanoma and synovial sarcoma	One partial response (14)	Davis et al. (2010)
	Multiple myeloma	Two complete responses and seven partial responses (17)	Robbins et al. (2011)
		Three complete responses and seven partial responses (11)	Rapoport et al. (2010)
CEA	Colorectal	One partial response (3)	Parkhurst et al. (2011)
MAGE A-3	Melanoma, esophageal, synovial sarcoma	One complete response and four partial responses (9)	Morgan et al. (2013)
	Melanoma and multiple myeloma	Lethal cardiac toxicity (2)	Linette et al. (2013)
HBsAg	HBV-related hepatocellularcarcinoma (HCC)	No clinical response (1)	Qasim et al. (2014)

Abbreviations: CEA, carcinoembryonic antigen; HBsAg, Hepatitis B surface antigen; MART-1, melanoma antigen recognized by T cells; Melan-A, melanocyte antigen; MAGE A-3, melanoma associate antigen 3; NY-ESO-1, New York esophageal squamous cell carcinoma 1.

and reported two of the four patients treated experienced no disease progression for 24 months, with the third patient showing objective partial regression. These early results are promising although a larger cohort of patients will need to be assessed in order to determine whether this approach is better than other FDA approved therapies targeting CD20, such as Rituximab ([McLaughlin et al., 1998](#); [Coiffier et al., 2002](#)). Morgan et al. utilized a similar CAR

targeted to human epidermal growth factor receptor 2 (Her2/erbB2) in colon cancer, however the first patient treated experienced respiratory failure. This was likely due to the targeting of infused T cells to erbB2⁺ lung epithelial cells and subsequently resulted in the cessation of the trial ([Morgan et al., 2010](#)). The results of these two trials highlight the need for further trials using third generation CARs.

Table 2

Summary of clinical trials utilizing CAR modified T cells for adoptive immunotherapy.

Target antigen	Cancer	Receptor	Results (number of patients treated in trial)	Reference/Clinical Trial ID
CD19	CLL	scFv-CD137-CD3 ζ	Two complete responses and one partial response (3)	Kalos et al. (2011) , Porter et al. (2011)
	ALL	scFv-CD137-CD3 ζ	Molecular remission in both patients, one relapsed with antigen negative disease (2)	Grupp et al. (2013) Lee et al. (2014)
	ALL	scFv-CD28-CD3 ζ	Overall complete response rate of 88% (16)	Brentjens et al. (2011)
	ALL and non-Hodgkin lymphoma	scFv-CD28-CD3 ζ	Overall complete response rate of 66.7% (21)	
	CLL and B-ALL	scFv-CD28-CD3 ζ	Stabilized disease (4)	
erbB2	Colorectal	scFv-CD28-CD137-CD3 ζ	Trial ceased due to patient death (1)	Morgan et al. (2010)
anti-FR	Ovarian	scFv- Fc ϵ R1 γ	No response (14)	Kershaw et al. (2006)
GD2	Neuroblastoma	scFv- CD3 ζ	Three complete responses (19)	Louis et al. (2011)
Lewis Y	AML	scFv-CD28-CD3 ζ	One patient experienced cytogenetic remission (4)	Ritchie et al. (2013)
CD20	NHL and mantle cell lymphoma	scFv-CD3 ζ	Two complete response, one partial response, four with stable disease (7)	Till et al. (2008)
	NHL	scFv-CD137-CD28-CD3 ζ	One partial response, two patients without evaluable disease remained progression free (3)	Till et al. (2012)
CAIX	Metastatic RCC	scFv-CD3 ζ	No response (3)	Lamers et al. (2006)
	Metastatic RCC		No response (9) (Two out of nine experienced autoimmune liver toxicity)	Lamers et al. (2013)
CD171	Neuroblastoma	scFv- CD3 ζ	One partial response (6)	Park et al. (2007)
GD2	Neuroblastoma	scFv-CD28-OX40-CD3 ζ	Currently in progress	NCT01822652 NCT02107963
	Non-neuroblastoma GD2 ⁺ solid tumors			
FR	Ovarian	scFv-CD137- CD3 ζ	Proposed	Kandalaft et al. (2012)
F19 protein (FAP)	FAP ⁺ malignant pleural mesothelioma (FAPME-1)	scFv-CD28- CD3 ζ	Proposed	Petrausch et al. (2012)
erbB2	Head and neck squamous cell carcinoma	scFv-CD28-CD3 ζ	Proposed	van Schalkwyk et al. (2013)

Abbreviations: ALL, acute lymphoblastic leukemia; AML, acute myeloid leukemia; CAIX, carbonic anhydrase IX; CLL, chronic lymphocytic leukemia; erbB2, erythroblastosis oncogene B2; FAP, fibroblast activation protein (associated with asbestos); FR, folate receptor; GD2, ganglioside; NHL, non-Hodgkin's lymphoma; RCC, renal cell carcinoma.

Table 3
Adverse events associated with adoptive immunotherapy clinical trials.

Adoptive immunotherapy	Target antigen	Tumor	Adverse events	References
TIL with lymphodepletion and IL-2	Various	Melanoma	Vitiligo (4/13), uveitis (1/1)	Dudley et al. (2002)
Young TIL with lymphodepletion and IL-2	Various	Metastatic melanoma	Vitiligo (1/20)	Besser et al. (2010)
CTL	MART-1	Melanoma	Melanocyte destruction (1/1)	Yee et al. (2000)
TIL and IL-2	Various	Melanoma	Autoimmune thyroiditis (1/25)	Ridolfi et al. (2003)
TIL and IL-2	Various	Melanoma	Systemic and ocular autoimmunity (1/1)	Yeh et al. (2009)
Gene-modified T cells expressing TCR	MART-1, gp100	Melanoma	Melanocyte destruction (29/36)	Johnson et al. (2009)
Gene-modified T cells expressing TCR	MAGE-A3	Melanoma	Neurological toxicity (3/9)	Morgan et al. (2013)
Gene-modified T cells expressing TCR	MAGE-A3	Melanoma and myeloma	Cardiac toxicity (2/2)	Linette et al. (2013)
Gene-modified T cells expressing TCR	CEA	Colorectal cancer	Colitis (3/3)	Parkhurst et al. (2011)
CTL with CAR for CD19	CD19	CLL and ALL	B cell aplasia (3/3)	Kalos et al. (2011)
CTL with CAR for CAIX	CAIX	RCC	Liver toxicity (grade III–IV) (3/3)	Lamers et al. (2006)
CTL with CAR for erbB2	erbB2	Colon cancer	Respiratory failure (1/1)	Morgan et al. (2010)

Abbreviations: ALL, acute lymphoblastic leukemia; CAIX, carbonic anhydrase IX; CEA, carcinoembryonic antigen; CLL, chronic lymphocytic leukemia; CTL, cytotoxic T lymphocyte; TCR, erythroblastosis oncogene B2; MART-1, melanoma antigen recognized by T cells; T cell receptor; TIL, tumor infiltrating lymphocyte.

Clinical trials in neuroblastoma and hematological malignancies highlight the potential of using CAR modified T cells for the treatment of cancer. However, in order to fully harness the capabilities of CAR T cells for the treatment of solid cancers, further studies to characterize and optimize CARs are required.

4. The future of adoptive immunotherapy

Whilst CAR modified T cells have shown great potential in the clinic, future studies are still required to potentially enhance the activity of these chimeric receptors whilst ensuring toxicity does not occur. This includes testing the combination of these CARs with therapies such as checkpoint inhibitors, addressing autoimmunity and the widespread implementation of adoptive immunotherapy as a standard of care therapy.

4.1. Combination of adoptive immunotherapy with existing therapies

Although adoptive immunotherapy using CAR modified T cells for some cancers such as hematological malignancies and neuroblastoma have shown early promise, clinical trials against other solid cancers have only shown moderate results to date. To potentially enhance the anti-tumor effect, CAR T cell therapy could be combined with immune checkpoint blockade antibodies, such as antibodies against cytotoxic T lymphocyte protein-4 (CTLA-4) (Hodi et al., 2010) and programmed death receptor-1 (PD-1) (Topalian et al., 2012) which have shown tremendous anti-tumor activity in patients as single agents or together in combination, particularly against metastatic melanoma with response rates of 40% (Wolchok et al., 2013). Studies have demonstrated that combining adoptive immunotherapy with an anti-PD-1 antibody (Peng et al., 2012; John et al., 2013) resulted in enhanced efficacy of infused T cells. This was found to be due to increased level of IFN- γ from CAR T cells at the tumor site (Peng et al., 2012). Other checkpoint pathways currently under investigation include lymphocyte-activation gene 3 (LAG-3) (Huang et al., 2004), which is being tested in a phase III breast cancer trial and T-cell immunoglobulin domain and mucin domain 2 (TIM-3) (Zhu et al., 2005), which is still in preclinical development. Therapeutic agents have also been developed to target the inhibitory ligands B7-H3 (Loo et al., 2012) (currently in clinical trials) and B7-H4 (He et al., 2011). Several studies have also shown that targeting multiple checkpoint pathways such as PD-1 and TIM-3 can result in the reversal of T cell exhaustion (Fourcade

et al., 2010; Sakuishi et al., 2010), and therefore the combination of multiple checkpoint inhibitors with CAR T cells warrants further investigation. Another target currently under investigation is the ectonucleotidase, CD73, which is upregulated in some cancers and converts immunostimulatory ATP to immunosuppressive adenosine. The accumulation of adenosine, which binds to the A2A receptor impairs anti-tumor T cell responses and NK cell function resulting in the promotion of tumor growth (Spychala, 2000; Deaglio et al., 2007; Beavis et al., 2013). The use of A2A receptor antagonists to target the CD73 pathway has been used in combination with CTLA-4 blockade, which resulted in enhanced anti-tumor immune responses (Iannone et al., 2014). Furthermore, it has been proposed that through other rational combinations, that include epigenetic modulators, such as histone deacetylase inhibitors (Sigalotti et al., 2005), it may be possible to change the immune profile of tumor cells, and therefore increase their recognition and susceptibility to the immune system (Skov et al., 2005; Magner et al., 2000). In future studies, it will be interesting to determine which immune therapies when combined with CAR T cells mediate the most effective therapeutic responses.

4.2. Adverse events

The targeting of normal healthy tissue by genetically modified infused T cells has been reported in several clinical trials (Lamers et al., 2006; Morgan et al., 2010, 2013; Kochenderfer et al., 2010; Cameron et al., 2013) in some cases resulting in the death of patients (summarized in Table 3). Several strategies have been developed (summarized in Table 4) in order to increase the safety of adoptive immunotherapy, including the incorporation of suicide genes (Bonini et al., 1997; Budde et al., 2013; Marin et al., 2012; Sato et al., 2007; Straathof et al., 2005; Tiberghien et al., 1994) and the administration of antibodies which are capable of depleting auto-reactive infused cells (Kieback et al., 2008; Vogler et al., 2010; Wang et al., 2011). The incorporation of suicide genes such as herpes simplex virus-thymidine kinase (Bonini et al., 1997; Berger et al., 2006) or inducible caspase 9 (Budde et al., 2013; Straathof et al., 2005; Di Stasi et al., 2011; Quintarelli et al., 2007; Hoyos et al., 2010) allows cells to be eliminated with the administration of ganciclovir or a chemical inducer of dimerization respectively. Both are currently being evaluated in clinical trials. The use of FDA approved antibodies Rituximab (Vogler et al., 2010; Serafini et al., 2004; Intronas et al., 2000) and Cetuximab (Wang et al., 2011) are currently being investigated in preclinical models to deplete infused genetically

Table 4

Approaches to increase the safety of CAR T cells in adoptive immunotherapy.

Approach	Advantage	Disadvantage	Status	References
HSV-TK suicide gene	Eliminate T cells with the anti-viral drug ganciclovir	Ganciclovir treatment for viral infections will eliminate infused T cells HSV-TK specific T cells detected Loss of therapy	Clinical trial in GVHD	Park et al. (2007), Bonini et al. (1997), Berger et al. (2006)
Inducible caspase 9 (iCasp9)	Eliminate T cells with chemical inducer of dimerization such as AP1903	Loss of therapy	Clinical trial in GVHD Clinical trial in CAR T cells commenced	Budde et al. (2013), Straathof et al. (2005), Di Stasi et al. (2011), Quintarelli et al. (2007), Hoyos et al. (2010)
CD20	Eliminate T cells with Rituximab, a FDA-approved antibody used in non-Hodgkin's lymphoma	Elimination of endogenous B cells Loss of therapy	Preclinical studies in GVHD mouse models	Vogler et al. (2010), Serafini et al. (2004), Introna et al. (2000)
mTMPK	Eliminate T cells with AZT, a FDA-approved drug used in HIV	Loss of therapy	<i>In vitro</i> data using T cell line No data using CARs	Sato et al. (2007)
Truncated EGFR	Eliminate T cells with Cetuximab, a FDA-approved antibody used in colon cancer	Hypomagnesaemia in patients treated with Cetuximab Loss of therapy	Preclinical studies in mouse model using CARs	Wang et al. (2011)
Myc TAG in CAR	Eliminate T cells with Myc-specific antibody	Loss of therapy	Preclinical studies in mouse models using TCR	Kieback et al. (2008)
Targeting multiple TAAs	Complete activation of T cells occurs when two CARs for different TAAs bind to the same tumor cell	Suboptimal activation of T cells in tumors that lose expression of an antigen	Preclinical studies in mouse models using CARs	Duong et al. (2011), Wilkie et al. (2012), Anurathapan et al. (2014)
Combination of activating signals	Activation is dependent on presence of multiple antigens	Suboptimal activation of T cells in tumors that lose expression of an antigen	Preclinical studies in mouse models using CARs	Kloss et al. (2013)
Targeting TAA and normal tissue antigen	Preservation of normal tissue as activation occurs only in presence of TAA	If inhibition is not complete, damage to normal tissue can still occur	Preclinical studies in mouse models using CARs	Fedorov et al. (2013)

Abbreviations: AZT, 3'-azido-3'-deoxythymidine; CAR, chimeric antigen receptor; EGFR, epidermal growth factor receptor; FDA, Food and Drug Administration; GVHD, graft versus host disease; HIV, human immunodeficiency virus; HSV-TK, herpes simplex virus thymidine kinase; TAA, tumor associated antigen; TCR, T cell receptor.

modified cells. However, as the administration of antibodies or other agents to eradicate infused cells can also ablate the cancer therapy, efforts to address autoimmunity by using a novel approach of multiple chimeric receptors have been investigated (Fedorov et al., 2013; Kloss et al., 2013; Duong et al., 2011). One approach is to genetically modify a T cell so that it expresses two different CARs specific for two TAAs, called dual-specific T cells (Duong et al., 2011). Dual-specific T cells were generated to express CARs specific for the TAAs, erythroblastosis oncogene B2 (erbB2) and folate binding protein (FBP). The dual specific T cells secreted more cytokine and lysed tumor target cells expressing both FBP and Her2, compared to cells expressing FBP or Her2 alone. Although this approach resulted in greater tumor specific responses, cells expressing either antigen alone were not spared completely, therefore autoimmunity could still occur, although possibly to a lesser extent. A recent study utilized an approach where activation of a T cell was dependent on combinatorial recognition of two prostate cancer antigens (Kloss et al., 2013). This approach utilized a combination of anti-prostate specific membrane antigen (PSMA) CAR, which provided suboptimal activation as well as a chimeric costimulatory receptor weakly specific for prostate stem cell antigen (PSCA) in the same T cell. This resulted in the elimination of PSCA⁺PSMA⁺ but not PSCA⁺PSMA⁻ tumor cells in mice. The same group also published a study where they used PD-1 or CTLA-4 incorporated into CARs in an effort to alter the activity of CD19 specific CARs in the same T cell (Fedorov et al., 2013). They generated a PSMA specific chimeric inhibitory receptor (CIR) containing a PD-1 intracellular signaling domain and expressed it in T cells, which had been modified to express a CD19 specific CAR. They demonstrated that T cells, which expressed both CARs exhibited a decrease in effector function when exposed to target cells expressing both CD19 and PSMA, compared to CD19

alone. In the clinical setting, it will be crucial to find the optimal balance between tumor targeting and normal tissue preservation. This could be achieved by altering the ratio of CIRs to CARs, the level of receptor expression (based on the strength of the promoter) and careful selection of antigen based on its expression level and distribution in addition to chimeric receptor affinity. Although most studies have focused on identifying TAAs, some investigations have sought to characterize antigens on normal tissue (Uhlen et al., 2010). In order to translate this approach to the clinic, further investigations are required to determine which normal tissue antigens are expressed on the majority of normal tissue but down-regulated or absent on tumor tissue. One possible target that CIR could be specific for is MHC I (Connor and Stern, 1990) as it is often down-regulated by tumors in order to escape immune detection. Cell surface tumor suppression antigens such as OPMCL, which is frequently inactivated by methylation in cancers including lung, colon and breast (Cui et al., 2008), could also be targeted.

4.3. Transition of adoptive immunotherapy in clinical trials to a standard of care treatment option

In order for adoptive immunotherapy to become standard of care, several issues need to be resolved. There is a need to develop consistent clinical protocols, which can be used as a baseline in multiple institutions. This includes the choice of gene modification technique and which chimeric receptor is used for a particular antigen target. In the past 5 years, several papers have been published pertaining to the use of CD19 CAR modified T cells for hematological malignancies (Grupp et al., 2013; Brentjens et al., 2011, 2013; Kochenderfer et al., 2010, 2012; Savoldo et al., 2011). Although some patients have experienced remarkable remissions, other

patients have only experienced partial responses. This is likely due to several factors, including differences in signaling domains present in the CAR, preconditioning regimens and tumor burden prior to treatment. Furthermore, once the most effective signaling domains in CARs for a particular cancer have been established, a cost effective and consistent genetic modification approach needs to be investigated. Some clinical protocols expand and activate T cells using beads (Grupp et al., 2013), however other groups use high dose IL-2 and the anti-CD3 antibody, OKT3 to stimulate cells prior to infusion (Ritchie et al., 2013). Currently, Carl June's group utilizes a lentiviral vector to modify T cells, however the majority of clinical trials have used retroviral transduction (Ritchie et al., 2013). Laurence Cooper's team is investigating a less complex and cheaper technique involving the electroporation of T cells and using the Sleeping Beauty transposon system (Singh et al., 2013). At present, there have been few studies comparing the efficacy of CAR T cells using these different genetic modification approaches, however in order for adoptive immunotherapy to progress, the optimal genetic modification and culturing technique needs to be determined. Although, adoptive immunotherapy currently costs approximately \$50,000 per patient, it is anticipated that due to the development of closed systems (Klapper et al., 2009) and following the initial investment into the construction of good manufacturing practice grade facilities, the overall cost of adoptive immunotherapy may significantly decrease. Furthermore, it is hoped that once CAR T cells are an established treatment for cancer, the use of cell banks, similar to those currently being used for Epstein–Barr virus specific T cells (Leen et al., 2013) can be used to store cryopreserved allogeneic CAR T cells. Resolution of these various issues relating to adoptive immunotherapy will enable better comparison of clinical trial results, which would allow for the true value of immunotherapy to be evaluated and possibly its widespread application for the treatment of various types of cancer.

5. Concluding remarks

Adoptive immunotherapy has come of age with success in recent clinical trials, in particular using TCR modified cells in melanoma and CAR T cells in hematological malignancies. However devising new strategies for improving the activity but limiting toxicity of these gene modified T cells in patients may broaden the approach to other cancers and increase the feasibility of adoptive immunotherapy as a standard of care treatment for cancer.

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