Enhanced Effector Responses in Activated CD8\(^+\) T Cells Deficient in Diacylglycerol Kinases

Matthew J. Riese\(^1\), Liang-Chuan S. Wang\(^2\), Edmund K. Moon\(^2\), Rohan P. Joshi\(^5\), Anjana Ranganathan\(^1\), Carl H. June\(^4\), Gary A. Koretzky\(^3\), and Steven M. Albelda\(^2\)

Abstract

Recent clinical trials have shown promise in the use of chimeric antigen receptor (CAR)-transduced T cells; however, augmentation of their activity may broaden their clinical use and improve their efficacy. We hypothesized that because CAR action requires proteins essential for T-cell receptor (TCR) signal transduction, deletion of negative regulators of these signaling pathways would enhance CAR signaling and effector T-cell function. We tested CAR activity and function in T cells that lacked one or both isoforms of diacylglycerol kinase (dgk) expressed highly in T cells, dgk\(\alpha\) and dgk\(\zeta\), enzymes that metabolize the second messenger diacylglycerol (DAG) and limit Ras/ERK activation. We found that primary murine T cells transduced with CARs specific for the human tumor antigen mesothelin showed greatly enhanced cytokine production and cytotoxicity when cocultured with a murine mesothelioma line that stably expresses mesothelin. In addition, we found that dgk-deficient CAR-transduced T cells were more effective in limiting the growth of implanted tumors, both concurrent with and after establishment of tumor. Consistent with our studies in mice, pharmacologic inhibition of dgks also augments function of primary human T cells transduced with CARs. These results suggest that deletion of negative regulators of TCR signaling enhances the activity and function of CAR-expressing T cells and identify dgks as potential targets for improving the clinical potential of CARs. Cancer Res; 73(12); 3566–77. ©2013 AACR.

Introduction

Elicitation of T-cell effector responses requires signal transduction through the T-cell antigen receptor (TCR), a protein complex that binds antigenic peptide presented by MHC, as well as through costimulatory receptors such as CD28. The effector responses generated from TCR signal transduction differ across individual T-cell subsets that are classified according to the expression of cell surface molecules (1). Expression of the surface molecule CD8, for instance, identifies a subset of T cells that respond to antigenic peptides presented in the binding groove of MHC class I. CD8\(^+\) T cells are responsible for the recognition and elimination of cells that express antigens derived from intracellular pathogens, such as viruses and intracellular bacteria, and also mutated or embryonic proteins generated by cells that have undergone malignant transformation. Although the extent to which CD8\(^+\) T cells are capable of controlling the development and progression of tumorigenesis remains uncertain, it is clear that deficiency of these cells increases the potential for the development of malignancy and that enhanced function of these CD8\(^+\) T cells can impart robust antitumor responses in both animal model systems and patients (2, 3). It is also clear in a number of models that although there may be an initial, potent CD8\(^+\) T-cell response, this response is often insufficient to fully protect from tumors (4). Mechanisms underlying this failure include (i) the lack of specific antigens with sufficient avidity for the TCR expressed by tumors, (ii) the absence of costimulatory ligands expressed by antigen-presenting cells (APC) within tumor-draining lymph nodes, and (iii) direct suppression of T-cell responses within the tumor microenvironment mediated by inhibitory secreted factors such as TGF\(\beta\), prostaglandin E (PGE)\(-2\), or adenosine, as well as inhibitory cells, such as regulatory T cells (5).

The potential for effective responses by CD8\(^+\) T cells in some instances of incurable malignancies, such as metastatic melanoma, has led to significant interest in defining ways to manipulate these cells to generate more potent responses as well as responses against a more diverse array of tumors. One promising approach has focused on engineering T cells to...
express chimeric antigen receptors (CAR). CARs are transmembrane fusion proteins that consist of an extracellular antibody domain capable of binding to a specific tumor antigen coupled to intracellular signaling domains from TCR and costimulatory components (6). In principle, CARs provide several advantages over the endogenous receptors of T cells. First, the engineered ligand-binding segment of CARs arises from an antibody, obviating the need for MHC presentation. Second, the antibody-binding component of the CAR can be chosen to be both specific and highly sensitive to antigens expressed selectively by tumor cells, increasing avidity of the T cell–tumor interaction and minimizing the potential for destruction of normal “bystander” host cells. Third, engagement of the CAR by ligand stimulates both TCR and costimulatory signaling modules, eliminating a requirement for expression of costimulatory ligands by tumor-draining APCs. CAR-expressing T cells that come into contact with tumor cells expressing the antigen of interest have been shown to develop functional responses that lead to tumor cell lysis and cytokine production.

There has been considerable success in the use of CARs in animal models (6, 7), and recently, CAR-expressing T cells have been shown to be effective in patients to treat refractory chronic lymphocytic leukemias (CLL; refs. 30, 31). Although T cells engineered to express CARs are capable of overcoming some limitations of the endogenous immune system to combat tumors (e.g., CARs are not MHC restricted and hence will lyse tumor cells that have downregulated MHC expression), CAR-expressing T cells still lack intrinsic programming to overcome, perhaps, the most important component that limits CD8+ T-cell antitumor responses: the inhibitory tumor microenvironment. We hypothesized that deletion of proteins that limit the strength of TCR signal might overcome this obstacle of impaired CD8+ T-cell antitumor immunity and impart significantly enhanced antitumor functioning in CAR-expressing CD8+ T cells.

For our studies, we chose to target an inhibitor of diacylglycerol (DAG), an essential second messenger that is created by the cleavage of phosphatidylinositol 4,5 bisphosphate by phospholipase Cy1 (PLCy1) after PLCy1 is phosphorylated and activated by the protein tyrosine kinases that are recruited to the stimulated TCR (1). DAG activates signaling molecules leading to several second messenger cascades, most notably the Ras/ERK pathway that is known to be essential for T-cell activation (7). After its generation, DAG is actively metabolized into phosphatidic acid by one of the 2 isoforms of diacylglycerol kinases (dgk) present within T cells, dgkα or dgkζ (8). Previously, we and others observed that deletion of either dgk isoform potentiates DAG-mediated Ras and extracellular signal-regulated kinase (ERK) activation and augments TCR-induced cytokine production and T-cell proliferation (9–11). We have found further that deletion of dgkζ results in improved CD8+ T-cell responses by augmenting signaling via the TCR when mice are challenged with a transplantable subcutaneous tumor (12). However, neither the absence of dgkα or dgkζ is sufficient to enable a completely successful antitumor response. We speculated therefore that combining CAR therapy with dgk deficiency might boost the ability of T cells to respond to a tumor challenge. Herein, we report the results of our studies showing this augmentation, suggesting that such a combined therapeutic approach may have use in future clinical trials.

Materials and Methods

Mice

Mice deficient in dgkα, dgkζ, or both backcrossed to C57Bl/6 have been described previously (9, 10). C57Bl/6 mice containing a transgene for the OVAp TCR (OT-I mice) were obtained from the Jackson Laboratories. Dgkζ-deficient CD45.2 CD90.2 OT-1 mice were created by backcrossing these 2 strains. All experiments were carried out in mice 6 to 12 weeks old. Animal maintenance and experiments were carried out in accordance with the Institutional Animal Care and Use Committee at the University of Pennsylvania (Philadelphia, PA).

Listeria infection and EL4-ova tumor model experiment

Splenic CD8+ T cells were isolated from wild-type or dgkζ-deficient CD45.2 CD90.2 OT-1 mice by flow cytometry (CD8+ CD44+Flt3−) as described (12). Twenty thousand cells were transferred intravenously into CD45.2, CD90.1 recipient mice subsequently infected intravenously with 5,000 chListeria-ova 24 hours after T-cell transfer. One week later, CD45.2+ donor cells were isolated from spleens of recipient mice according to the manufacturer's instructions (Miltenyi Biotec), and 1.5 × 106 of isolated cells were transferred intravenously into CD45.1, CD90.2 mice that had been inoculated with 2.5 × 105 EL4-ovalbumin (EL4-ova) tumor cells, a murine lymphoma line that stably expresses ovalbumin (13), in the right flank 2 weeks prior. Tumors were barely palpable at time of T-cell transfer. One week later, mice were euthanized, tumor size was measured, and T cells from spleens and tumors were analyzed.

T-cell transduction

Mesocar, a fusion protein that contains the antigen-binding region of an antibody specific for the human tumor antigen mesothelin fused with CD8α transmembrane domain, CD3ζ, and the costimulatory domain of 4-1BB, has been described previously (14). cDNA encoding mesocar was subcloned into the MIGR retrovirus (15), which also expresses GFP using an internal ribosomal entry site. The sequence of antimesothelin Fv was provided by Ira Pastan (National Cancer Institute, Bethesda, MD; ref. 16). Infective particles were generated from the supernatants of 293T cells transfected with retroviral vector plasmid and helper plasmids using Lipofectamine 2000 (Invitrogen), as previously described (17). Primary murine T cells were isolated as suggested by the manufacturer (Miltenyi Biotec) from the spleens of wild-type or dgk-deficient mice and incubated in 24-well plates [4 × 106 cells/well in 2 mL T-cell media with 100 U/mL interleukin (IL)-2] coated with α-CD3 (1 μg/mL) and α-CD28 (2 μg/mL). After 48 hours, cells (1 × 106 cells/well) were mixed with retrovirus (1 mL crude viral supernatant) in a 24-well plate coated with Retronectin (50 μg/mL; Clontech) and centrifuged without braking at room temperature for 30 minutes at 1,200 g. After overnight incubation, cells were expanded with 50 U/mL of IL-2 for 48 hours.
Coating beads with recombinant human mesothelin

Target antigens were chemically crosslinked to tosylated 4.5 μm Dynabeads (Invitrogen, #140-13), using the manufacturers’ instructions. In brief, 4 × 10^5 beads were incubated 16 to 18 hours at 37°C in the presence of 20 μg of recombinant human mesothelin (RayBiotech, #230-00043) in 0.1 mol/L sodium phosphate buffer (pH 7.4) with shaking. After incubation, beads were washed and resuspended in PBS containing 0.5% bovine serum albumin to a final volume of 1 mL.

Evaluation of CAR T-cell effector functions

Cytotoxicity and IFN ELISA. A stable cell line of the mouse mesothelioma line AE17 expressing human mesothelin subsequently engineered to express luciferase has been described (14, 18). Cytokine release assays were conducted by coculture of T cells with target cells at the described ratios, in triplicate, in 96-well round bottom plates in 200 μL. After 18 hours, cell lysis was determined from the detection of luciferase from the remaining cells using a previously described assay (14). An ELISA Kit (Biolegend) was used to measure IFN-γ.

WINN assay. A total of 1 × 10^6 mesothelin-expressing TC1 cells, a murine non–small cell lung cancer line with well-established use in the WINN assay (19), were co-injected into the right flank along with 2 × 10^5 CAR-transduced T cells (routinely 50% of which were GFP positive, and thus transduced with CAR). Ten days later, mice were euthanized, and tumor volume was assessed.

Intravenous transfer of CAR-T cells in mice with preexisting tumor. C57Bl/6 mice were inoculated subcutaneously with 2 × 10^6 AE17 meso cells. Seven days later, at which point tumors were approximately 100 mm^3, mice were injected with 1 × 10^7 CAR-transduced T cells intravenously by tail vein. Tumor development was monitored by caliper measurement of tumor diameter over an additional 10 days. Each volumetric determination was derived from the formula 0.52πa^2b, with a representing the minor axis and b representing the major axis. Alternately, mice were sacrificed at 3 or 6 days after transfer, and the presence of T cells within spleen or tumor was determined by evaluating for GFP expression within T-cell subsets by flow cytometry.

Expression of cytotoxic markers following CAR activation. A total of 2 × 10^6 mesoCAR-T cells derived from mouse splenocytes replete or deficient in DGKz were placed in individual wells of a 24-well plate with or without 2 × 10^6 mesothelin-coated beads for 18 hours, at 37°C in the presence of 30 U/mL of IL-2. After incubation, T cells were stained for the presence of the surface markers TRAIL (eBioscience, #12-5951-82) or Fasl (eBioscience, #12-5911-81), or the intracellular markers granzyme B (BD, #51-2090KZ) or perforin (eBioscience, #12-9392-82), using protocols described by the manufacturer. Flow cytometry histograms of marker expression were evaluated from cells that were positive for GFP (indicating expression of CAR) and CD8, and negative for CD4.

T-cell immunoblotting and CD69 upregulation

To assess for Erk phosphorylation, 1 × 10^6 mesoCAR-transduced T cells were incubated either with mesothelin- or albumin-coated beads in a 1:4 ratio (cells:beads), or with α-CD3ε antibody at 2.5 μg/mL. final concentration for indicated time points. Lysates were prepared and immunoblotted for phosphorylated Erk, total Erk, or tubulin (antibodies all from Cell Signaling) as previously described (12). Alternately, protein-bead stimulations were allowed to proceed for 5 hours, and then the surface upregulation of CD69 was determined by flow cytometry.

Primary human CAR-T-cell assays

Primary human T cells were obtained from the University of Pennsylvania Clinical Cell Production Facility and mock infected or transduced with lentivirus expressing mesoCAR as previously described (14). A total of 5 × 10^6 primary human T cells were subsequently added to 24-well plates that had been seeded with 5 × 10^5 cells either from the epithelial mesothelioma (EM) human mesothelioma line or a stable derivative cell line, EM-meso, engineered to express high levels of mesothelin, in a 24-well dish. After 96 hours of coincubation (which included the addition of another 3 × 10^7 EM or EM-meso cells at the midpoint of coincubation), cells were resuspended, and T cells were isolated via Lymphoprep density gradient separation (Axis-Shield) as suggested by the manufacturer. Cells were assessed for viability using Trypan blue, and 1 × 10^5 live T cells were cocultured with 5 × 10^5 EM-meso-luc cells expressing luciferase in 96-well plates in the presence or absence of the dgk inhibitors DGK1 (R59022) or DGK2 (R59949; Sigma) at 5 μg/mL. After 18 hours, remaining tumor cells were washed and lysed, and luminescence was evaluated. Cell lysis determinations were corroborated with visual estimate of remaining numbers of tumor cells.

For studies with TGFβ, 1 × 10^6 primary human T cells that had been mock infected or transduced with lentivirus-expressing mesoCAR were coincubated with 5 × 10^5 EM-meso-luc cells in the presence of indicated concentrations of TGFβ for 18 hours, and cell numbers were determined as described above.

Results

DGKz-deficient activated CD8+ T cells show enhanced response to tumor

We have previously shown that naive DGKz-deficient CD8+ T cells specific for ovalbumin (OT-I cells) are better able to control tumor growth and undergo activation than naïve wild-type OT-I cells after transfer into mice bearing EL4-ova-expressing tumors (12). However, in those studies, implanted tumors were not completely eradicated. We wondered whether the effect of DGKz deletion would be improved if, instead of naïve T cells, we made use of activated T cells that could potentially confer a more robust antitumor response. In addition, this approach would more closely mirror current clinical trials of adoptive CDS T-cell tumor immunotherapy that use cells preactivated before transfer. To generate uniform populations of activated cells, we transferred naïve OT-I cells sufficient or deficient in DGKz into congenically marked mice, and then infected the recipient animals with Listeria engineered to express ovalbumin. One week later, antigen-experienced (CD44-high) donor OT-I cells were recovered from
spleens and transferred into EL4-ova tumor-bearing mice. Initially, we noted that expansion of naïve dgkζ-deficient OT-I cells was more robust when compared with naïve wild-type OT-I cells in response to the antigenic challenge with Listeria-ova (Fig. 1A), as in other dgkζ-deficient CD8+ T cell-models of acute infection (9, 12); however, there was no difference in activation phenotype of the recovered cells as assessed by CD44 expression between the 2 different genotypes (data not shown). After transfer of equal numbers of wild-type or dgkζ-deficient effector cells into EL4-ova tumor-bearing mice, we found that although wild-type OT-I cells conferred no appreciable antitumor effect, tumors in mice treated with dgkζ-deficient activated OT-I cells were significantly ($P = 0.05$) reduced in size compared with untreated animals (Fig. 1B). Dgkζ-deficient effector cells also persisted in increased numbers within the spleen of host animals (Fig. 1C) and were observed in larger quantities within the tumors of host animals (Fig. 1D). These data show that deficiency of dgkζ confers enhanced antitumor potential in preactivated T cells. As in our previous study with naïve dgkζ-deficient OT-I CD8+ T cells, however, we found that transfer of activated dgkζ-deficient CD8+ T cells was insufficient to completely eradicate tumors, suggesting that the strategy of targeting dgkζ alone is insufficient in curtailing the progression of established tumors.

Deletion of dgkζ enhances functional responses of T cells downstream of CARs

Given that deletion of dgkζ conferred enhanced activity of CD8+ T cells against established tumors but did not seem to be curative, we wondered whether inhibition of dgk function might augment other approaches shown to have efficacy in enhancing T-cell responses against tumor. Therefore, we next designed studies to test the impact of dgk deficiency on effector function of CAR-expressing T cells. Our first experiments tested whether dgkζ loss would augment functional responses after ligation of CARs, similar to the augmentation of TCR-induced functions that we have shown previously (9). For this analysis, we made use of mesoCAR, a fusion protein that has high affinity for the human tumor antigen mesothelin, present on human mesothelioma, pancreatic, and ovarian cancer, coupled to the signaling motifs of the TCR CD3ζ chain and the inducible T-cell costimulatory receptor 4-1BB. Wild-type or dgkζ-deficient activated OT-1 cells were transduced with mesoCAR-expressing retrovirus, resulting in approximately 50% transduction efficiency (Fig. 2A). Transduction did not affect the activation state of the T cells, as assessed by expression of CD44, or expression of the endogenous TCR, as assessed with tetramer specific for OT-I (Fig. 2A). MesoCAR-transduced dgkζ-deficient and wild-type OT-1 cells were then compared in their ability to produce IFNγ and mediate target

Figure 1. Dgkζ-deficient activated CD8+ T cells show enhanced tumor responses in vivo. A, twenty thousand naïve (CD44lo) CD8+ wild-type (wt) or dgkζ-deficient OT-I cells were injected intravenously into congenically marked (CD90.1) mice. Twenty-four hours later, the mice were euthanized, and the presence of donor OT-I T cells (CD90.2+; ova tetramer+) was assessed (n = 5, quantitation of 1 of 3 representative experiments is shown). CD90.2- cells from A were isolated magnetically and 1 x 10^6 cells were injected intravenously into CD45.1+ mice bearing 2-week-old subcutaneous EL4-ova tumors. One week later, mice were euthanized and assessed for tumor size (B), persistence of donor (CD45.2+; ova tetramer+) T cells (C), and tumor-infiltrating donor T cells (D). No T cell mice did not receive donor T cells, and CD45.2+ cells were not detected in any organ tissue (data not shown; B, data from 3 pooled experiments. C and D, data from 1 of 3 representative experiments, n = 5 in each group).
cell lysis after incubation with AE17ova-meso, a murine cell line engineered to express both ovalbumin and human mesothelin. Transduced OT-I cells lacking dgkζ displayed enhanced IFNγ production and enhanced cytotoxicity after incubation with mesothelioma cell lines (Fig. 2B and C), indicating that deletion of dgkζ enhances the function of CAR-transduced CD8+ T cells against AE17 cells that express both ova and mesothelin.

Combined deletion of dgkζ and dgkα markedly enhances T-cell responses after stimulation of mesoCAR

The finding that deletion of dgkζ enhanced mesoCAR T-cell functional responses suggested that these 2 strategies may be used together for potentiating CD8+ T-cell tumor responses. However, these initial experiments were complicated by the fact that the target cells expressed antigens for both TCR (ovalbumin) and CAR (mesothelin). Thus, to avoid potentially confounding our results with ovalbumin-specific TCRs, we continued our experiments in non-TCR transgenic animals. Further, as it is well appreciated that an additional isoform of dgk, dgkα, operates in a similar fashion as dgkζ in T cells and may blunt the effects of targeting dgkζ alone in augmenting T-cell function, we intercrossed dgkζ and dgkζ-deficient mice to generate animals deficient in both dgk isoforms to study CAR-T cells generated from double knockout (DKO) mice. Naïve T cells were isolated from wild-type, dgkα−/−, dgkζ−/−, or DKO mice and infected with retrovirus encoding mesoCAR under high IL-2 concentrations that favored CD8+ T-cell growth (cells were 85% CD8+ T cells at the end of incubation). As observed with deletion of dgkζ in OT-1 cells, deletion of either dgkα or dgkζ in this population of cells expressing the mesoCAR receptor conferred enhanced cytokine production and cytotoxicity when the T cells were incubated with tumor cells expressing mesothelin (Fig. 3A and B). Strikingly, DKO cells showed profoundly enhanced effector functions compared with cells with deletion of either dgk isoform alone or wild-type cells. The enhanced cytotoxicity observed in these cell lines was mesothelin specific because mesoCAR-transduced DKO T cells did not lyse cells AE17 cells expressing an unrelated antigen (AE17ova cells; Fig. 3C).

We next evaluated whether the changes in signal transduction that we have previously observed downstream of the TCR in dgk-deficient T cells, for example, enhanced Ras/Erk/AP-1 signaling (9, 10), were also present downstream of CAR. To that end, we developed a means to stimulate mesoCAR T cells that did not require mesothelin-expressing cells because these cells express their own Ras signaling molecules, such as Erk, that could interfere with identifying changes specific to T cells after stimulation. For these studies, we used tosylated activator beads coated with albumin (as a control) or beads coated with
Deletion of both dgks enhances activity of T cells downstream of CARs against tumor in vivo

We next sought to determine whether deletion of dgk isoforms conferred enhanced antitumor responses in vivo making use of the WINN assay. For these experiments, mice were injected with a mixture of TC1meso cells, a murine non–small cell lung cancer line (19), along with mesoCAR-expressing wild-type T cells or mesoCAR T cells lacking dgkξ, dgkζ, or both dgk isoforms. Although mice that received wild-type mesoCAR-transduced T cells or T cells lacking a single isoform of dgk were unable to completely control the growth of mesotheliomas, DKO T cells eradicated the mesotheliomas (Fig. 5A), indicating that, as suggested by in vitro studies, targeting dgk generates meaningful enhancement of CD8+ T cells against tumor. Pronounced differences were also noted when AE17meso cells were used as target cells in the WINN assay (data not shown). Although this experiment offered proof-of-principle that DKO T cells conferred enhanced in vivo activity against mesothelioma, it did not directly assess whether deletion of dgk isoforms would be capable of limiting the growth of established tumors, which is more representative of how CAR-T cells would be used clinically. To that end, AE17meso cells were injected into the flanks of mice, and tumors were allowed to develop to approximately 100 mm³ in size before intravenous administration of CAR-T cells. Under these conditions, although wild-type mesoCAR-transduced T cells were ineffective at limiting tumor growth,
mesoCAR-transduced T cells deficient in either dgk isofrom significantly (P < 0.05) decreased the rate of tumor growth (Fig. 5B), an effect increased by the deletion of both dgk isofroms. In addition to enhanced effector function, this effect could, in part, be related to the increased number of dgk-deficient mesoCAR T cells in tumor-bearing mice because quantitative differences in T cells between mesoCAR wild-type and dgk isoform mice were appreciated 6 days after transfer (Fig. 5C); however, mesoCAR T cells of any genotype were not detected at day 10 or later timepoints under the experimental conditions used.

Inhibition of dgks confers enhanced antitumor responses to human T cells

Although the demonstration of enhanced mesoCAR function in murine T cells lacking dgk isofroms provides a rationale for targeting dgks to augment T-cell responses against tumors, an important next step is to establish a role for dgks in CAR-expressing primary human T cells. Over the course of our studies with primary human T cells transduced with mesoCARs, we have noted that these cells develop reduced functional responsiveness after extended coculture with mesothelin-expressing tumor cells. As shown, 96 hours of coinubation of mesoCAR T cells with EM-meso cells, a mesothelioma line engineered to express high levels of mesothelin, results in significant impairment of mesoCAR T cell lysis of target cells upon reculture (Fig. 6A, left). This effect is reminiscent of various models of antigen-induced anergy, as impaired cytotoxicity is not generated after coculture of mesoCAR T cells with parental EM cells that do not express mesothelin (Fig. 6A). Because we and others have previously shown that deletion of dgks mitigates the induction of anergy (10, 11), we hypothesized that inhibition of dgks might also diminish the impaired cytotoxicity observed in our assay. To test this, we incubated mesoCAR T cells with EM-meso cells for 96 hours, and then assessed their ability to lyse target cells in the presence of dgk inhibitors R59022 (DGK1 inhibitor) or R59949 (DGK2 inhibitor). We observed that the addition of either dgk inhibitor was sufficient to reverse the impaired cytotoxicity present in mesothelin-exposed mesoCAR T cells (Fig. 6A, center and right), indicating that, similar to our findings in mice, inhibition of dgk function seems to augment antitumor activity of primary human T cells expressing CARs. These data also suggest that in addition to augmenting TCR (or CAR)-mediated signaling, blockade of dgk may enhance CAR T-cell antitumor responses by mitigating antigen-induced unresponsiveness of the effector cells.

Dgk-deficient T cells show reduced sensitivity to TGFβ

Following our observation that inhibition of dgks reverses the antigen-induced inactivation of CAR-expressing T cells, we asked whether deletion of dgk might also reduce sensitivity to other inhibitory influences of T cells. One such inhibitor, TGFβ, is of particular relevance, because secretion of this cytokine by tumor cells has been shown to actively inhibit CD8+ effector T-cell responses against tumors (20). Furthermore, TGFβ is speculated to mediate its effects, in part, by dampening the...
Ras/Erk signal transduction pathway, which is known to be affected by dgks (21, 22). We initially examined whether TGFβ could lead to inhibition of cytotoxicity of human mesoCAR T cells incubated with EM-meso cells. The addition of TGFβ resulted in diminished cytotoxicity by transduced mesoCAR human T cells (Fig. 6B), at levels of TGFβ similar to that present in culture media of EM-meso cells (81.31 pg/mL/24 hours/10^6 cells; data not shown). Next, we tested whether deletion of dgks could diminish the effects of TGFβ. Murine mesoCAR T cells replete or deficient in dgks were incubated with AE17meso cells as described previously, and cytotoxicity and IFNγ secretion were assessed in the presence or absence of TGFβ. As shown before, deletion of either or both dgk isoforms resulted in mesoCAR T cells with enhanced function when compared with wild-type mesoCAR T cells (Fig. 7A and B). These data suggest that deletion of dgks confers relative resistance to TGFβ for mesoCAR T cells. The finding of relative insensitivity to inhibitory stimuli seems not to be solely restricted to TGFβ because greater functional responses were also observed by dgk-deficient mesoCAR-transduced T cells in the presence of the inhibitory stimuli PGE2 and adenosine (Supplementary Fig. S1), although to a lesser extent than that observed with TGFβ. Collectively our data suggest that deletion of dgks augments effector function of CAR-expressing CD8^+ T cells not only by augmenting signaling through the CAR itself but also by reducing sensitivity of the effector cells to physiologically relevant inhibitory signals.

**Increased FasL and TRAIL expression in dgk-deficient T cells**

Because TGFβ expression suppresses mediators of CD8^+ T-cell cytotoxicity (23), such as perforin, granzyme B, FasL, and TRAIL, and because CAR T-cell cytotoxicity is mediated through these effector molecules (reviewed in ref. 24), we hypothesized that dgk-deficient CAR cells would show greater...
upregulation of these cytotoxic mediators when compared with wild-type T cells and that this upregulation may be the basis for the enhanced cytotoxicity observed in dgk-deficient mesoCAR T cells. Wild-type mesoCAR T cells or mesoCAR T cells deficient in one or both dgk isoforms were exposed to mesothelin-coated beads in the presence of IL-2 for 18 hours. Expression of FasL, TRAIL, granzyme B, and perforin were then evaluated by flow cytometry. As predicted, mesoCAR T cells deficient in one or both dgks showed enhanced expression of the cytotoxic cell surface proteins FasL and TRAIL when compared with wild-type mesoCAR T cells (Supplementary Fig. S2). In contrast, no difference was observed in the expression of the intracellular cytotoxic proteins granzyme B- or perforin expressing in cells lacking dgks (Supplementary Fig. S3). These data suggest that FasL and TRAIL may help facilitate the augmented cytotoxicity observed in dgk-deficient mesoCAR T cells.

Discussion

Current strategies aimed at augmenting T-cell immune responses against malignancy have focused either on assisting the initial activation or priming of naïve T cells, or on potentiating the effects of activated or primed T cells. For instance, antibodies that activate CD40 on APCs upregulate costimulatory molecules that help facilitate priming of naïve cells (25–27). In contrast, antibodies that block the T-cell inhibitory cell surface molecule CTLA-4 minimally impact naïve T cells but significantly enhance proliferation and effector function of primed T cells (28, 29). Inhibition of proteins that negatively regulate signal transduction downstream of the TCR has garnered recent attention as a potential strategy for augmenting T-cell responses to malignancy at the time of T cell priming. For instance, deletion of Casitas-B-lineage lymphoma b (cbl-b), an E3 ubiquitin ligase responsible for the degradation of several proteins important in TCR signal transduction, results in T cells with a decreased requirement of costimulation at the time of activation and enhanced antitumor activity of naïve T cells (30–33).

We had previously shown, similar to cbl-b, that deletion of dgkζ enhanced the effector functions of naïve CD8⁺ T cells (12). Although deletion of cbl-b and dgkζ both result in changes downstream of the TCR, dgkζ and dgkζ, likely act directly to regulate the threshold for activation of T cells downstream of the TCR. Under a currently posited model of TCR signaling, the interplay of 2 Ras-activating proteins, the guanine nucleotide-exchange factors SOS and RasGRP1, determine whether the threshold for Ras activation is met within a T cell after TCR engagement, an event required for T-cell activation (34). In this model, TCR ligation results in the production of DAG, which binds and activates RasGRP1, and generates small amounts of active Ras. If enough active Ras is generated, it is able to bind an allosteric Ras-binding site on SOS, activating SOS and facilitating generation of most of the active Ras within activated T cells. In a manner largely consistent with this model, we had previously found that deletion of dgkζ resulted in a decreased threshold of T-cell activation, a finding that correlated with enhanced responses in naïve CD8⁺ T cells.

In the studies presented herein, we found that deletion of dgkζ also has profound effects on activated T cells. After uniform activation of naïve wild-type or dgkζ-deficient ovalbumin-specific T cells with Listeria-ova and then transfer into mice with subcutaneous ovalbumin-expressing EL4 lymphoma, we found that dgkζ-deficient T cells showed enhanced activity against tumor and increased persistence, both within the spleen and the tumor itself. This finding suggests that alteration of T-cell threshold plays an important role at multiple stages with the T-cell life cycle and identifies dgkζ as a means to simultaneously target both naïve and activated populations of effector T cells.

Figure 6. Dgk inhibitors enhance the cytotoxic capacity of impaired human mesoCAR-transduced T cells. A, mesoCAR-transduced primary human cells were left unexposed or exposed to a human tumor line that does not express mesothelin (EM) or expresses high levels of mesothelin (EM-meso) for 96 hours. A total of 10⁵ T cells were then isolated and recultured with 5 x 10⁴ luciferase-expressing EM-meso cells for 18 hours in the absence or presence of dgk inhibitors R59022 (DGK1 Inhibitor) or R59949 (DGK2 inhibitor) and cell death of target cells was assessed by luciferase release (data from triplicate wells of one of the 3 representative experiments are shown. P of EM-meso exposed T cells to EM-meso exposed T cells = 0.004 in no inhibitor group. P of EM-meso-exposed T cells in the absence of inhibitor to DGK1 inhibitor = 0.006 or DGK2 inhibitor = 0.003). B, lysis of 5 x 10⁵ EM-meso cells was assessed during incubation with no T cells or 10⁵ untransduced or mesoCAR-transduced primary human T cell for 18 hours in the presence or absence of the indicated concentration of TGFβ (data from triplicate wells of one of the 3 representative experiments. P of mesoCAR T cells between 0 and 10 pg/mL = 0.05, 0 and 100 pg/mL = 0.025, and 0 and 1000 pg/mL = 0.017).
The role of dgks in limiting the effector function of activated CD8^+ T cells makes dgks a potential target for CAR-expressing T cells, a strategy gaining traction in the clinical treatment of human malignancies. In current clinical trial protocols, human T cells are transduced with lentivirus-expressing CARs that contain CD3 and CD28 or CD3 and 41BB (CD137), a process that induces T-cell division and activation upon tumor antigen binding (35). However, it is now clear that additional strategies will be necessary to harness the full potential of CAR-T cells, especially in the treatment of solid malignancies. Although clinical trials in CLL seem promising (35, 36), earlier works with CAR-transduced T cells in solid malignancies, such as ovarian cancer (37) and renal cell carcinoma (38), were less encouraging, with an absence of objective tumor response and the lack of T-cell persistence. In the studies described here, we evaluated whether dgks represent a possible strategy for augmenting CAR-expressing T cells.

After establishing a retroviral system to efficiently transduce murine T cells, we found, as with TCR signaling, that deletion of dgk\(\alpha\) augmented Erk activation, a phosphorylation event that occurs downstream of DAG formation, after CAR ligation. Deletion of dgk\(\alpha\) was also found to augment CAR-dependent effector functions because these cells exhibited enhanced cytokine production and target cell killing relative to their wild-type counterparts. Deletion of both T-cell isoforms of dgk resulted in even greater enhancement of effector functions of mesoCAR-transduced cells and resulted in control of tumor \textit{in vivo} in tumor-bearing mice. These results are encouraging for ongoing clinical trials because murine studies of CAR-transduced T cells have accurately predicted clinical outcomes in past trials (6, 35, 39).

Although CARs are uniquely positioned to deal with the limited presence of antigen and costimulation found within the tumor environment, they do not address a third issue relating to T-cell response to tumor: inhibitory stimuli. In these studies, we uncovered deletion of dgks as a novel strategy for enhancing T-cell activity in the presence of inhibitory stimuli. Specifically, we found that prolonged exposure to antigen or the tumor microenvironment inhibitors PGE\(_2\), adenosine, and TGF\(\beta\) were less able to suppress CD8^+ effector functions in T cells that lacked one or both T-cell isoforms of dgk. TGF\(\beta\) is thought to be a key mediator of tumor-mediated inhibition because it is secreted by a variety of tumors, and inhibition of TGF\(\beta\) signaling, through the expression of a dominant negative receptor, results in enhanced tumor-specific activity of cytotoxic lymphocytes (40, 41). In fact, the amount of TGF\(\beta\) produced by human cancers, such as prostate cancer, inversely correlates to a patient’s overall prognosis (42, 43).
enhanced Ras activation imparted by the loss of dgks might explain how dgk-deficient lymphocytes develop insensitivity to TGFβ. Because TGFβ is known to result in the reduced phosphorylation of Itk, a Tec kinase important in PLCγ1 activation (+), and because PLCγ1 is the protein directly responsible for DAG generation in T cells, one could envision that deletion of dgks might directly subvert this TGFβ-induced signaling alteration. Future studies will address the potential means by which loss of dgks confers insensitivity to TGFβ, including direct effects, such as enhanced Ras activation downstream of TCR, and indirect effects, such as increased expression of cytotoxic proteins including FasL and TRAIL.

We believe that these findings could be translated clinically. We have shown that pharmacologic inhibition of dgks augment the efficacy of human CAR T cells under inhibitory in vitro conditions. This preliminary finding suggests that dgks play an important role in human T cells and that dgks may represent attractive clinical targets in augmenting CAR T-cell–based therapies. We are currently extending our preliminary data in human T cells by developing model systems in which dgk activity is suppressed through decreased dgk expression (e.g., through expression of shRNA specific for dgks) or inhibited dgk function (e.g., through expression of dominant negative forms of dgks). Of course, as one develops more potent CAR T cells, issues of toxicity may become relevant. We could not assess toxicity in our model system because the CAR T cells are specific for human mesothelin and do not react with an endogenous mouse protein. However, we have developed a second model in our laboratory using CARs specific for the murine antigen mouse fibroblast activation protein (FAP), an antigen overexpressed on cancer-associated fibroblasts. In preliminary studies, we have observed enhanced antitumor efficacy using dgkζ-deficient FAP-CAR T cells in tumor-bearing mice, without evidence of enhanced toxicity (data not shown). Despite this initial indication that dgks can be targeted safely, careful attention to toxicity will be required if dgk knockdown of CAR T cells is moved into clinical trials. One approach that our group has used when introducing CAR T cells with augmented function is to begin the trial using T cells transduced with short-lived CAR mRNA (45), thus mitigating the potential for chronic CAR-induced autoimmunity.

Our data support the notion that combining CAR expression, which improves targeting of T cells to tumors and drives an initial stimulatory response, with inhibition of proteins known to blunt the effectiveness of the TCR signal may synergize to drive an effector response. The additional value of creating effector T cells resistant to the inhibitory environment generated by the tumor is also likely to contribute to the enhanced efficacy observed in this combined approach. Collectively, our data suggest that targeting dgks, as one means to blunt an endogenous inhibitory response, could be a useful mechanism to improve CAR-based strategies in the treatment of human malignancy.

Disclosure of Potential Conflicts of Interest
No potential conflicts of interest were disclosed.

Authors' Contributions
Conception and design: M.J. Riese, L.-C.S. Wang, S.M. Albelda
Acquisition of data (provided animals, acquired and managed patients, provided facilities, etc.): M.J. Riese, L.-C.S. Wang, E.K. Moon, R.P. Joshi, A. Ranganathan, G.A. Koretzky
Analysis and interpretation of data (e.g., statistical analysis, biostatistics, computational analysis): M.J. Riese, L.-C.S. Wang, E.K. Moon, G.A. Koretzky, S.M. Albelda
Writing, review, and/or revision of the manuscript: M.J. Riese, L.-C.S. Wang, E.K. Moon, C.H. June, S.M. Albelda
Administrative, technical, or material support (i.e., reporting or organizing data, constructing databases): G.A. Koretzky
Study supervision: G.A. Koretzky, S.M. Albelda

The costs of publication of this article were defrayed in part by the payment of page charges. This article must therefore be hereby marked advertisement in accordance with 18 U.S.C. Section 1734 solely to indicate this fact.

Received October 10, 2012; revised March 1, 2013; accepted March 27, 2013; published OnlineFirst April 10, 2013.

References
Enhanced Function of dgk-deficient CAR T Cells


Enhanced Effector Responses in Activated CD8+ T Cells Deficient in Diacylglycerol Kinases


Updated version
Access the most recent version of this article at:
doi:10.1158/0008-5472.CAN-12-3874

Supplementary Material
Access the most recent supplemental material at:
http://cancerres.aacrjournals.org/content/suppl/2013/10/02/0008-5472.CAN-12-3874.DC1

Cited articles
This article cites 45 articles, 17 of which you can access for free at:
http://cancerres.aacrjournals.org/content/73/12/3566.full.html#ref-list-1

Citing articles
This article has been cited by 14 HighWire-hosted articles. Access the articles at:
/content/73/12/3566.full.html#related-urls

E-mail alerts
Sign up to receive free email-alerts related to this article or journal.

Reprints and Subscriptions
To order reprints of this article or to subscribe to the journal, contact the AACR Publications Department at pubs@aacr.org.

Permissions
To request permission to re-use all or part of this article, contact the AACR Publications Department at permissions@aacr.org.