

## Enhanced Effector Responses in Activated CD8<sup>+</sup> T Cells Deficient in Diacylglycerol Kinases

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### 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<sup>+</sup> 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<sup>+</sup> 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<sup>+</sup> 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<sup>+</sup> 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<sup>+</sup> 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

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**Note:** Supplementary data for this article are available at Cancer Research Online (<http://cancerres.aacrjournals.org/>).

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doi: 10.1158/0008-5472.CAN-12-3874

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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<sup>+</sup> 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<sup>+</sup> T-cell antitumor immunity and impart significantly enhanced antitumor functioning in CAR-expressing CD8<sup>+</sup> 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 phosphatidyl (4, 5) inositol bisphosphate by phospholipase C $\gamma$ 1 (PLC $\gamma$ 1) after PLC $\gamma$ 1 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, dgk $\alpha$  or dgk $\zeta$  (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 dgk $\zeta$  results in improved CD8<sup>+</sup> T-cell responses by augmenting signaling via the TCR when mice are challenged with a transplantable subcutaneous tumor (12). However, neither the absence of dgk $\alpha$  or dgk $\zeta$  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 dgk $\alpha$ , dgk $\zeta$ , 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. Dgk $\zeta$ -deficient CD45.2 CD90.2 OT-I mice were created by backcrossing these 2 strains. All experiments were carried out in mice 6 to 12 weeks old. Animal maintenance and experimentation 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<sup>+</sup> T cells were isolated from wild-type or dgk $\zeta$ -deficient CD45.2 CD90.2 OT-I mice by flow cytometry (CD8<sup>+</sup>CD44<sup>lo</sup>) as described (12). Twenty thousand cells were transferred intravenously into CD45.2, CD90.1 recipient mice subsequently infected intravenously with 5,000 cfu *Listeria-ova* 24 hours after T-cell transfer. One week later, CD45.2<sup>+</sup> donor cells were isolated from spleens of recipient mice according to the manufacturer's instructions (Miltenyi Biotec), and  $1.5 \times 10^6$  of isolated cells were transferred intravenously into CD45.1, CD90.2 mice that had been inoculated with  $2.5 \times 10^5$  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 CD8a transmembrane domain, CD3 $\zeta$ , 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 [ $4 \times 10^6$  cells/well in 2 mL T-cell media with 100 U/mL interleukin (IL)-2] coated with  $\alpha$ -CD3 (1  $\mu$ g/mL) and  $\alpha$ -CD28 (2  $\mu$ g/mL). After 48 hours, cells ( $1 \times 10^6$  cells/well) were mixed with retrovirus (1 mL crude viral supernatant) in a 24-well plate coated with Retronectin (50  $\mu$ g/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 tosylactivated 4.5  $\mu\text{m}$  Dynabeads (Invitrogen, #140-13), using the manufacturers' instructions. In brief,  $4 \times 10^7$  beads were incubated 16 to 18 hours at 37°C in the presence of 20  $\mu\text{g}$  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  $\mu\text{L}$ . 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- $\gamma$ .

**WINN assay.** A total of  $1 \times 10^6$  mesothelin-expressing TC1 cells, a murine non-small cell lung cancer line with well-established use in the WINN assay (19), were coinjected into the right flank along with  $2 \times 10^5$  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  $2 \times 10^6$  AE17 meso cells. Seven days later, at which point tumors were approximately 100  $\text{mm}^3$ , mice were injected with  $1 \times 10^7$  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  $0.52a^2b$ , 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  $2 \times 10^6$  mesoCAR-T cells derived from mouse splenocytes replete or deficient in dgks were placed in individual wells of a 24-well plate with or without  $2 \times 10^6$  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,  $1 \times 10^6$  mesoCAR-transduced T cells were incubated either with mesothelin- or

albumin-coated beads in a 1:4 ratio (cells:beads), or with  $\alpha$ -CD3 $\epsilon$  antibody at 2.5  $\mu\text{g}/\text{mL}$  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  $5 \times 10^6$  T cells were subsequently added to 24-well plates that had been seeded with  $5 \times 10^5$  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  $3 \times 10^5$  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  $1 \times 10^5$  live T cells were cocultured with  $5 \times 10^3$  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  $\mu\text{g}/\text{mL}$ . After 18 hours, remaining tumor cells were washed and lysed, and luminescence was evaluated. Cell lysate determinations were corroborated with visual estimate of remaining numbers of tumor cells.

For studies with TGF $\beta$ ,  $1 \times 10^5$  primary human T cells that had been mock infected or transduced with lentivirus-expressing mesoCAR were coincubated with  $5 \times 10^3$  EM-meso-luc cells in the presence of indicated concentrations of TGF $\beta$  for 18 hours, and cell numbers were determined as described above.

## Results

### Dgk $\zeta$ -deficient activated CD8 $^+$ T cells show enhanced response to tumor

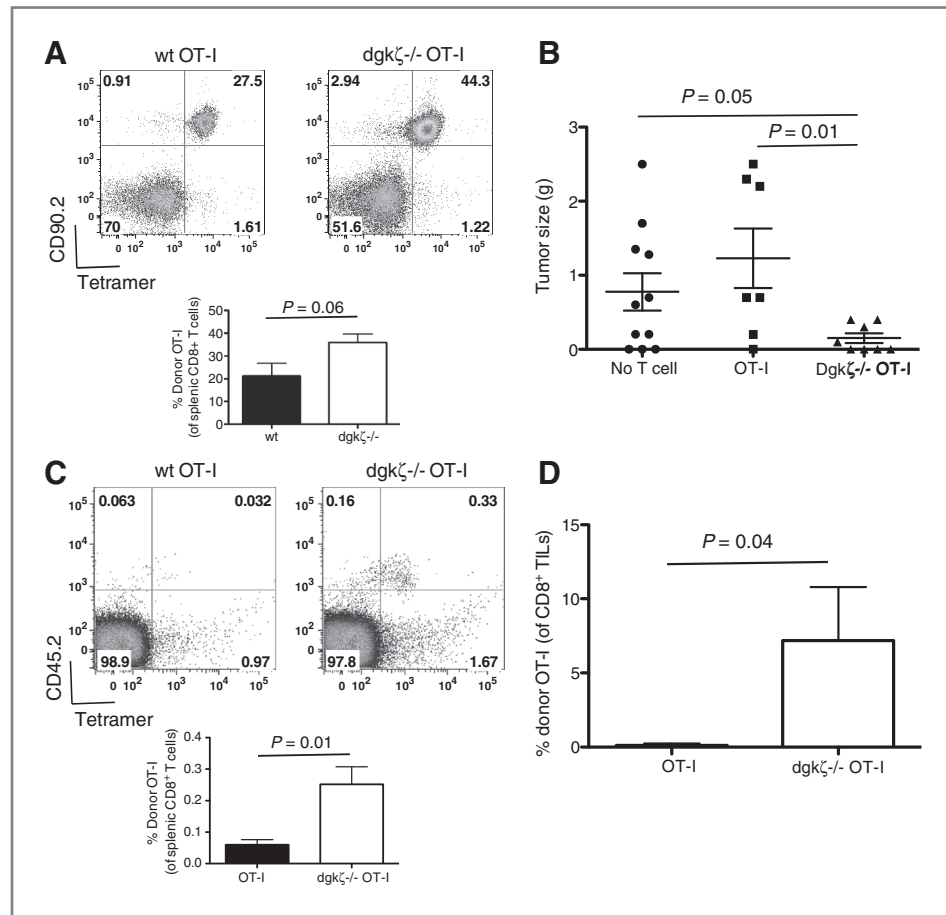
We have previously shown that naïve dgk $\zeta$ -deficient CD8 $^+$  T cells specific for ovalbumin (OT-I cells) are better able to control tumor growth and undergo activation than naïve wild-type OT-I cells after transfer into mice bearing EL4-ova-expressing tumors (12). However, in those studies, implanted tumors were not completely eradicated. We wondered whether the effect of dgk $\zeta$  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 CD8 $^+$  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 dgk $\zeta$  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

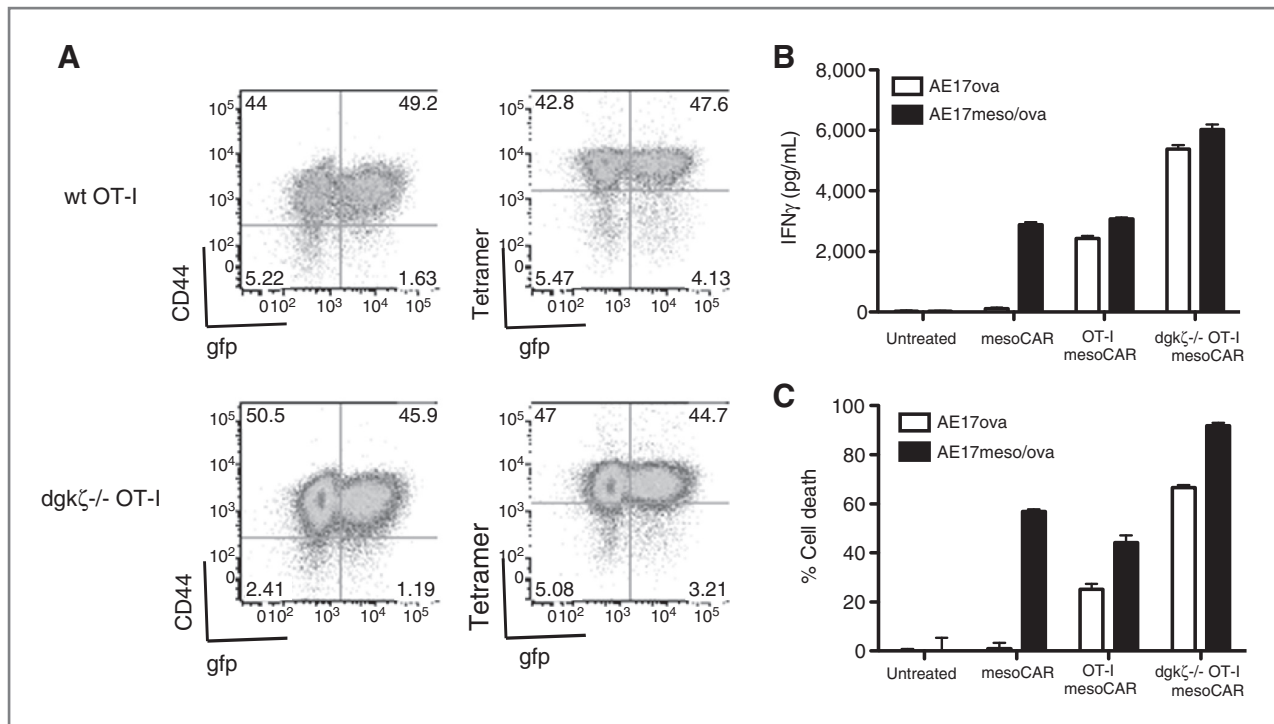
spleens and transferred into EL4-ova tumor-bearing mice. Initially, we noted that expansion of naïve dgk $\zeta$ -deficient OT-I cells was more robust when compared with naïve wild-type OT-I cells in response to the antigenic challenge with *Listeria-ova* (Fig. 1A), as in other dgk $\zeta$ -deficient 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 dgk $\zeta$ -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 dgk $\zeta$ -deficient activated OT-I cells were significantly ( $P = 0.05$ ) reduced in size compared with untreated animals (Fig. 1B). Dgk $\zeta$ -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 dgk $\zeta$  confers enhanced antitumor potential in preactivated T cells. As in our previous study with naïve dgk $\zeta$ -deficient OT-I CD8 $^+$  T cells, however, we found that transfer of activated dgk $\zeta$ -deficient CD8 $^+$  T cells was insufficient to completely eradicate tumors, suggesting that the strategy of targeting dgk $\zeta$  alone is insufficient in curtailing the progression of established tumors.

### Deletion of dgk $\zeta$ enhances functional responses of T cells downstream of CARs

Given that deletion of dgk $\zeta$  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 dgk $\zeta$  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 CD3 $\zeta$  chain and the inducible T-cell costimulatory receptor 4-1BB. Wild-type or dgk $\zeta$ -deficient activated OT-I 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 dgk $\zeta$ -deficient and wild-type OT-I cells were then compared in their ability to produce IFN $\gamma$  and mediate target

**Figure 1.** Dgk $\zeta$ -deficient activated CD8 $^+$  T cells show enhanced tumor responses *in vivo*. **A**, twenty thousand naïve (CD44 $^{\text{lo}}$ ) CD8 $^+$  wild-type (wt) or dgk $\zeta$ -deficient OT-I cells were injected intravenously into congenically marked (CD90.1) mice. Twenty-four hours later, the mice were injected intravenously with 5,000 cfu *Listeria-ova*, and one week later, 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  $1 \times 10^6$  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).





**Figure 2.** Enhanced CAR effector function in dgk $\zeta$ -deficient CD8<sup>+</sup> T cells. CD8<sup>+</sup> wild-type (wt) or dgk $\zeta$ -deficient OT-I T cells were isolated and transduced with mesoCAR retrovirus expressing gfp from an internal ribosomal entry site. A, cells were evaluated for expression of gfp, CD44 as a marker of activation, and ovalbumin tetramer to assess TCR expression. B and C, mesoCAR-transduced wt or dgk $\zeta$ -deficient OT-I cells were incubated with 5,000 AE17 ovalbumin cells or AE17 cells expressing both mesothelin and ovalbumin at a ratio of 40:1 in a 96-well plate for 24 hours, and the presence of IFN $\gamma$  (B) or luciferase (cytotoxicity; C) in cell supernatants was assessed. Calculated estimates of cytotoxicity were confirmed by visual evaluation of cell culture wells. One of the 3 representative experiments is shown; each well was carried out in triplicate. (B and C,  $P < 0.0001$  between dgk/AE17ova and wt/AE17ova or dgk/AE17ovameso and wt/AE17ovameso. B,  $P = 0.0327$  between dgk/AE17ova and dgk/AE17ovameso,  $P = 0.002$  between wt/AE17ova and wt/AE17ovameso. C,  $P < 0.0001$  between dgk/AE17ova and dgk/AE17ovameso,  $P = 0.007$  between wt/AE17ova and wt/AE17ovameso).

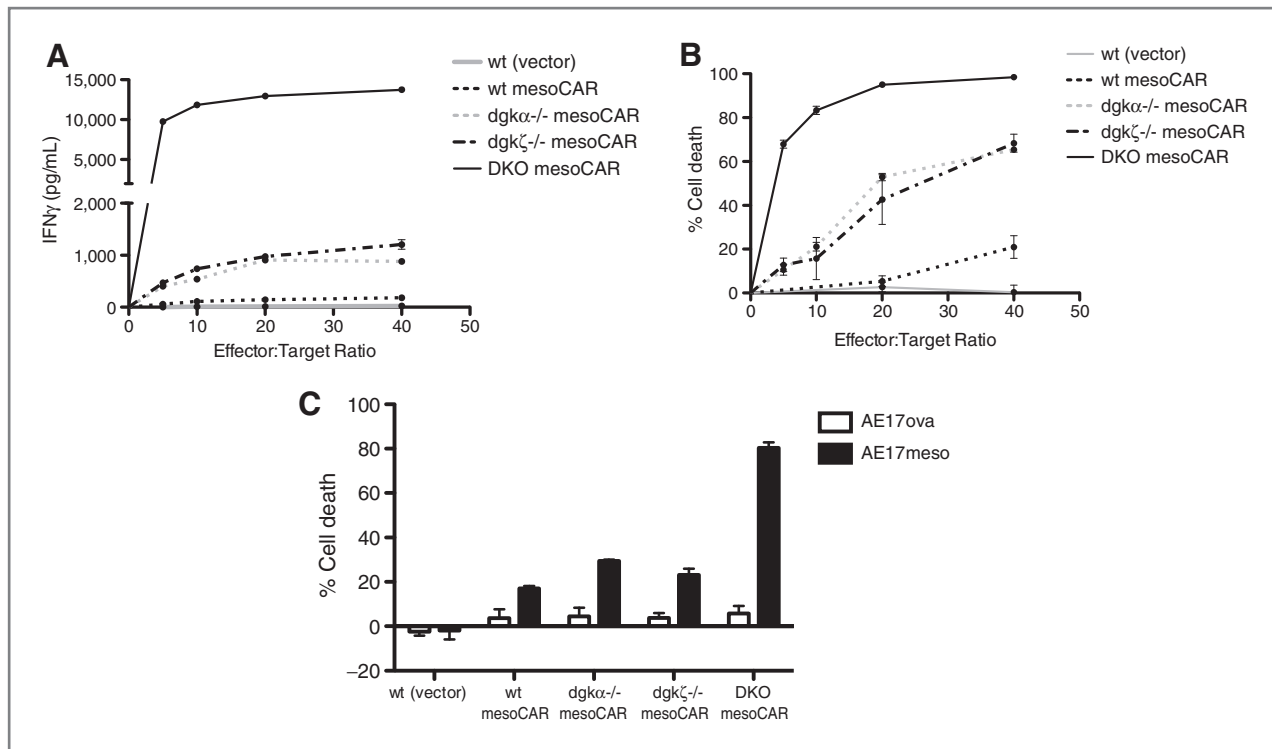
cell lysis after incubation with AE17ova-meso, a murine cell line engineered to express both ovalbumin and human mesothelin. Transduced OT-I cells lacking dgk $\zeta$  displayed enhanced IFN $\gamma$  production and enhanced cytotoxicity after incubation with mesothelioma cell lines (Fig. 2B and C), indicating that deletion of dgk $\zeta$  enhances the function of CAR-transduced CD8<sup>+</sup> T cells against AE17 cells that express both ova and mesothelin.

#### Combined deletion of dgk $\zeta$ and dgk $\alpha$ markedly enhances T-cell responses after stimulation of mesoCAR

The finding that deletion of dgk $\zeta$  enhanced mesoCAR T-cell functional responses suggested that these 2 strategies may be used together for potentiating CD8<sup>+</sup> 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, dgk $\alpha$ , operates in a similar fashion as dgk $\zeta$  in T cells and may blunt the effects of targeting dgk $\zeta$  alone in augmenting T-cell function, we intercrossed dgk $\alpha$  and dgk $\zeta$ -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, dgk $\alpha$ -/-, dgk $\zeta$ -/-, or DKO mice and infected with retrovirus encoding mesoCAR under high IL-2 concentrations that favored CD8<sup>+</sup> T-cell growth (cells were 85% CD8<sup>+</sup> T cells at the end of incubation). As observed with deletion of dgk $\zeta$  in OT-I cells, deletion of either dgk $\alpha$  or dgk $\zeta$  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 tosylactivated beads coated with albumin (as a control) or beads coated with

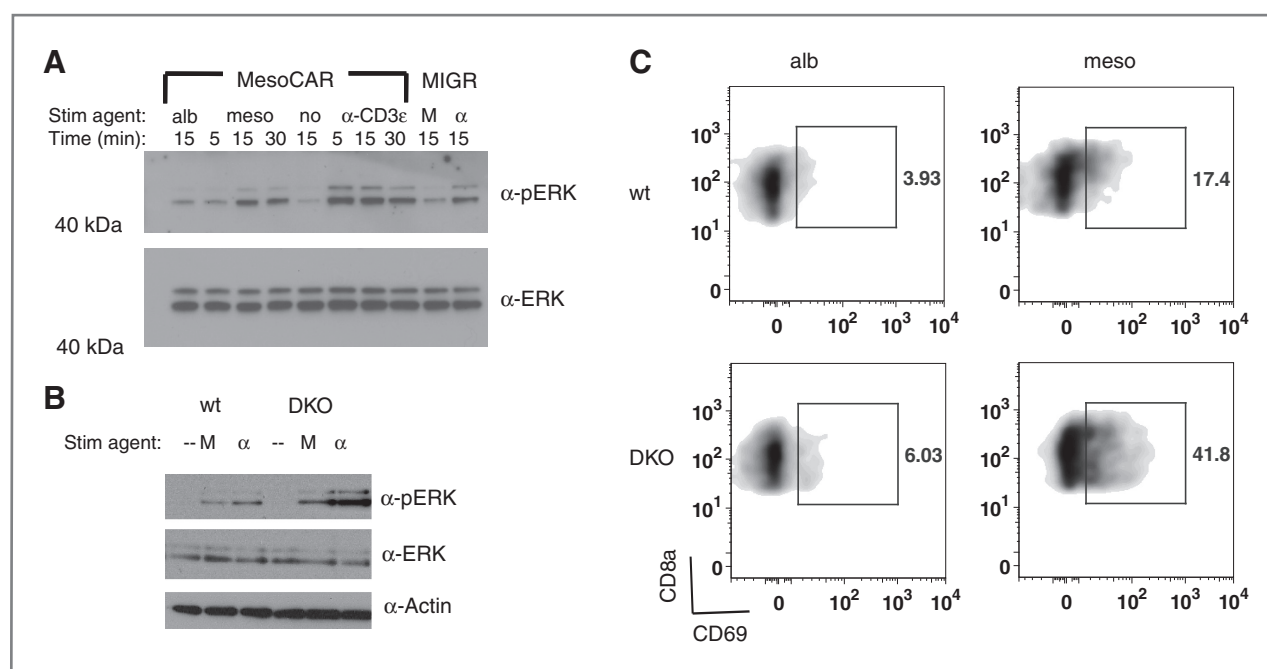


**Figure 3.** Deletion of both T-cell dgk isoforms significantly enhances CAR-T cell effector functions. T cells were isolated from wt, dgk $\alpha^{-/-}$ , dgk $\zeta^{-/-}$ , or DKO mice, transduced with mesoCAR, and assessed for IFN $\gamma$  production and cytotoxicity of target cells at indicated ratios as described in Fig. 2. In C, a ratio of 40:1 was used for experimental (AE17ovameso) or control (AE17ova) cell lines. One of the 3 representative experiments is shown; each data point was conducted in triplicate ( $P$  for all mesoCAR-expressing constructs  $< 0.0001$  for AE17ova and AE17ovameso).

mesothelin to stimulate the mesoCAR-expressing cells. Although phosphorylation of Erk was not observed during incubation with control beads, Erk phosphorylation could be readily detected during incubation with mesothelin-coated beads or, as expected, after stimulation of the TCR complex through CD3 $\epsilon$  (Fig. 4A). Moreover, activation of Erk by mesothelin beads required expression of mesoCAR because activation of T cells transduced with control retrovirus (MIGR) was not observed (Fig. 4A, right lanes). To test whether deletion of dgks enhanced Erk activation downstream of mesoCAR, we then repeated this experiment with mesoCAR-transduced T cells derived from DKO mice. Similar to the enhanced activation of Erk known downstream of the TCR in T cells deficient in dgks, loss of dgks augmented the activation of Erk downstream of mesoCAR (Fig. 4B). We extended our analysis by investigating the upregulation of CD69 in mesothelin-stimulated wild-type or DKO T cells, as CD69 expression is controlled by activation of the transcription factor AP-1 following Ras/Erk signaling (9). Consistent with the biochemical enhancement of Erk activation observed in DKO-transduced T cells, upregulation of CD69 was also increased in mesoCAR DKO T cells compared with wild-type cells (Fig. 4C), confirming a role for dgks in regulation of this pathway downstream of CAR. Together, these data suggest that dgk influences CAR signaling in a manner similar to the TCR and that the combination of CAR expression and dgk deletion could represent an effective strategy for augmenting CD8 $^{+}$  T-cell antitumor responses.

### 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 dgk $\alpha$ , dgk $\zeta$ , 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 mm $^3$  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,



**Figure 4.** Enhanced CAR signaling in dgk-deficient CD8<sup>+</sup> T cells. **A**, a total of  $1 \times 10^6$  CAR-transduced (MesoCAR) or vector control (MIGR) CD8<sup>+</sup> T cells were incubated with  $4 \times 10^6$  albumin-coated beads (alb) or mesothelin-coated beads (meso or M) or 2.5  $\mu$ g/mL  $\alpha$ -CD3 ( $\alpha$ -CD3 $\epsilon$  or  $\alpha$ ) for the indicated times and assessed for phosphorylated ( $\alpha$ -pERK) or total Erk ( $\alpha$ -ERK) by immunoblotting. **B**, levels of pERK, total ERK, and actin were assessed after stimulation of  $1 \times 10^6$  wild-type (wt) or dgk $\alpha$ <sup>-/-</sup>dgk $\zeta$ <sup>-/-</sup> (double knockout; DKO) CD8<sup>+</sup> CAR-T cells with  $4 \times 10^6$  mesothelin-coated beads (M) or 2.5  $\mu$ g/mL  $\alpha$ -CD3 ( $\alpha$ ) for 15 minutes. Each immunoblot analysis is a representative experiment from at least 3 independent iterations. **C**, a total of  $1 \times 10^6$  wt or DKO-deficient CD8<sup>+</sup> CAR T cells were incubated with  $4 \times 10^6$  beads coated with albumin (alb) or mesothelin (meso) for 5 hours and surface expression of CD69 was assessed by flow cytometry.

mesoCAR-transduced T cells deficient in either dgk isoform significantly ( $P < 0.05$ ) decreased the rate of tumor growth (Fig. 5B), an effect increased by the deletion of both dgk isoforms. 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 $\zeta$  T cells 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.

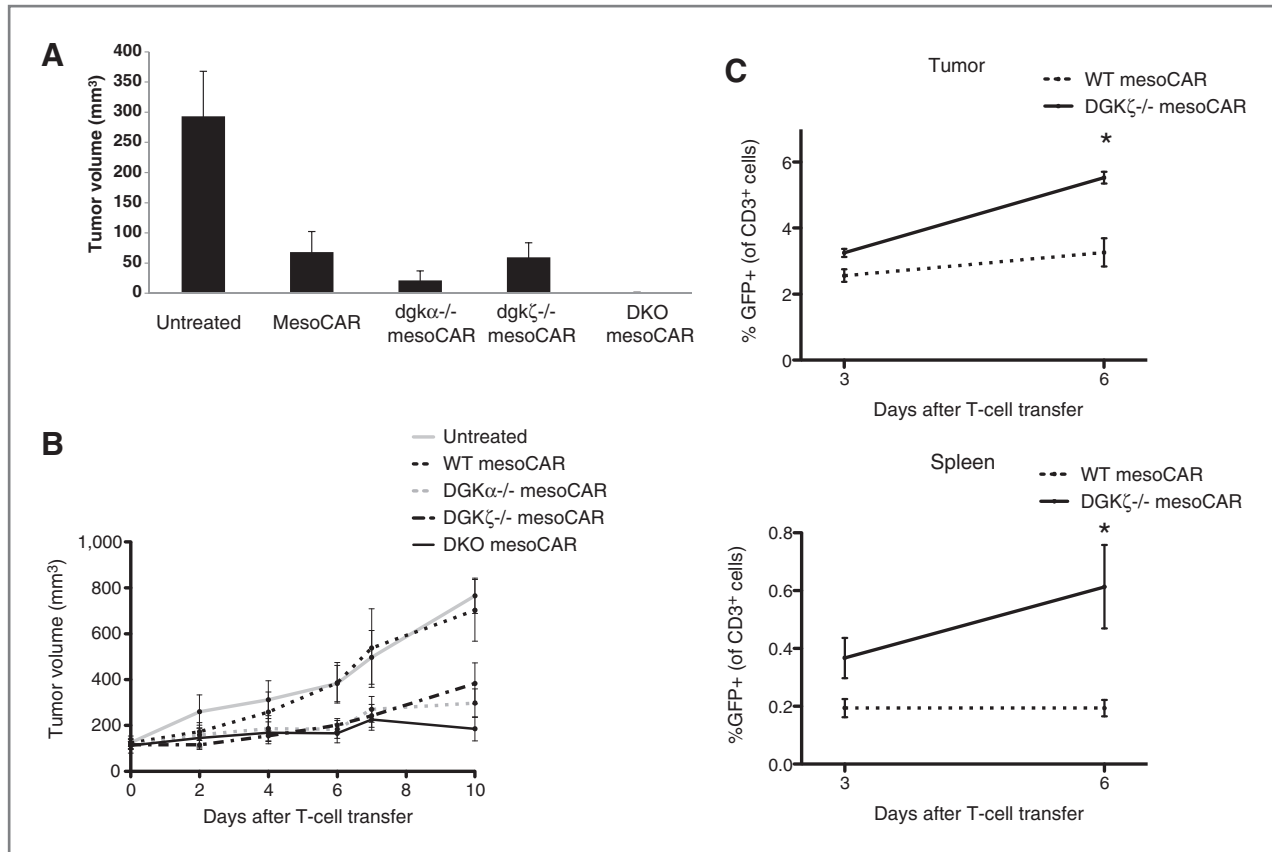
#### Inhibition of dgks confers enhanced antitumor responses to human T cells

Although the demonstration of enhanced mesoCAR function in murine T cells lacking dgk isoforms 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 coculture 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 T-cell antitumor responses by mitigating antigen-induced unresponsiveness of the effector cells.

#### Dgk-deficient T cells show reduced sensitivity to TGF $\beta$

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, TGF $\beta$ , is of particular relevance, because secretion of this cytokine by tumor cells has been shown to actively inhibit CD8<sup>+</sup> effector T-cell responses against tumors (20). Furthermore, TGF $\beta$  is speculated to mediate its effects, in part, by dampening the



**Figure 5.** Dgk-deficient mesoCAR-transduced T cells control mesothelioma *in vivo*. **A**, a total of  $1 \times 10^6$  TC1meso cells were coinjected subcutaneously with  $2 \times 10^5$  wild-type (wt) mesoCAR-transduced T cells or mesoCAR-transduced T cells lacking one or both (DKO) of the indicated dgk. Ten days later, mice were euthanized and tumors were measured. One of the 2 representative experiments ( $n = 4-5$ ) is shown.  $P$  of wt versus DKO mesoCAR T cells is 0.05. **B**, a total of  $2 \times 10^6$  AE17meso cells were injected into flanks of C57Bl/6 mice. One week later,  $1 \times 10^7$  CAR-transduced T cells of indicated genotype were injected intravenously into mice and tumors were measured at indicated time points ( $n = 5$  in each genotype,  $P$  for DKO CAR-T cells versus wt CAR-T cells = 0.0141 at day 10 posttransfer). **C**, alternately, mice were sacrificed 3 and 6 days after T-cell transfer of indicated genotypes as in **B** and the presence of mesoCAR T cells in tumor (top) or spleen (bottom) was determined [ $n = 3$  in each group, 1 of 2 representative experiments.  $P = 0.0082$  (tumor) and 0.0461 (spleen) at day 6; day 3 results did not differ significantly].

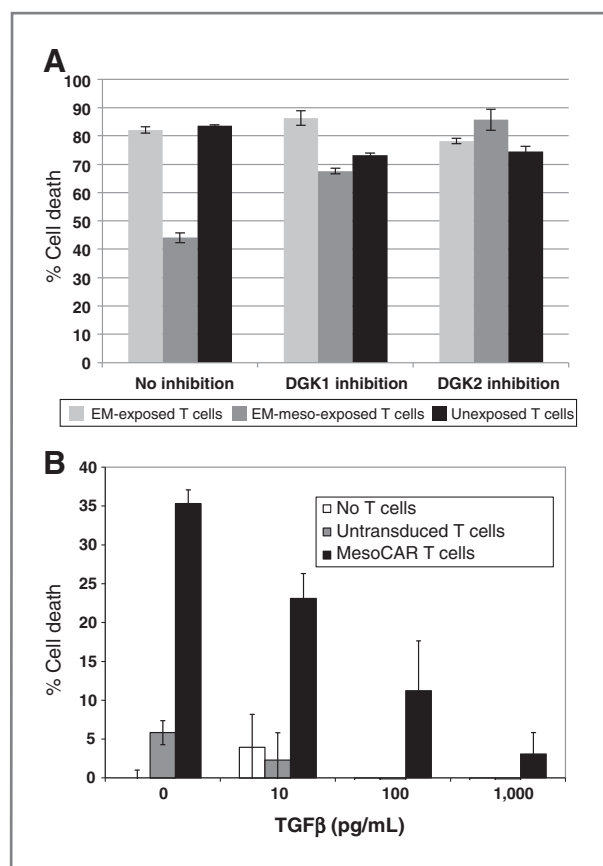
Ras/Erk signal transduction pathway, which is known to be affected by dgks (21, 22). We initially examined whether TGF $\beta$  could lead to inhibition of cytotoxicity of human mesoCAR T cells incubated with EM-meso cells. The addition of TGF $\beta$  resulted in diminished cytotoxicity by transduced mesoCAR human T cells (Fig. 6B), at levels of TGF $\beta$  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 TGF $\beta$ . Murine mesoCAR T cells replete or deficient in dgks were incubated with AE17meso cells as described previously, and cytotoxicity and IFN $\gamma$  production were assessed in the presence or absence of TGF $\beta$ . 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, white bars). Addition of TGF $\beta$  at lower (1 ng/mL) or higher (10 ng/mL) concentrations of TGF $\beta$  were found to reduce cytotoxicity and IFN $\gamma$  secretion in wild-type mesoCAR T cells; however, these functions were not affected in mesoCAR T cells deficient in either or both dgks (Fig. 7A and B). These data suggest that

deletion of dgks confers relative resistance to TGF $\beta$  for mesoCAR T cells. The finding of relative insensitivity to inhibitory stimuli seems not to be solely restricted to TGF $\beta$  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 TGF $\beta$ . 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 TGF $\beta$  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





**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  $10^5$  T cells were then isolated and recultured with  $5 \times 10^3$  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 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  $5 \times 10^3$  EM-meso cells was assessed during incubation with no T cells or  $10^3$  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.

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 expres-

sion 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.

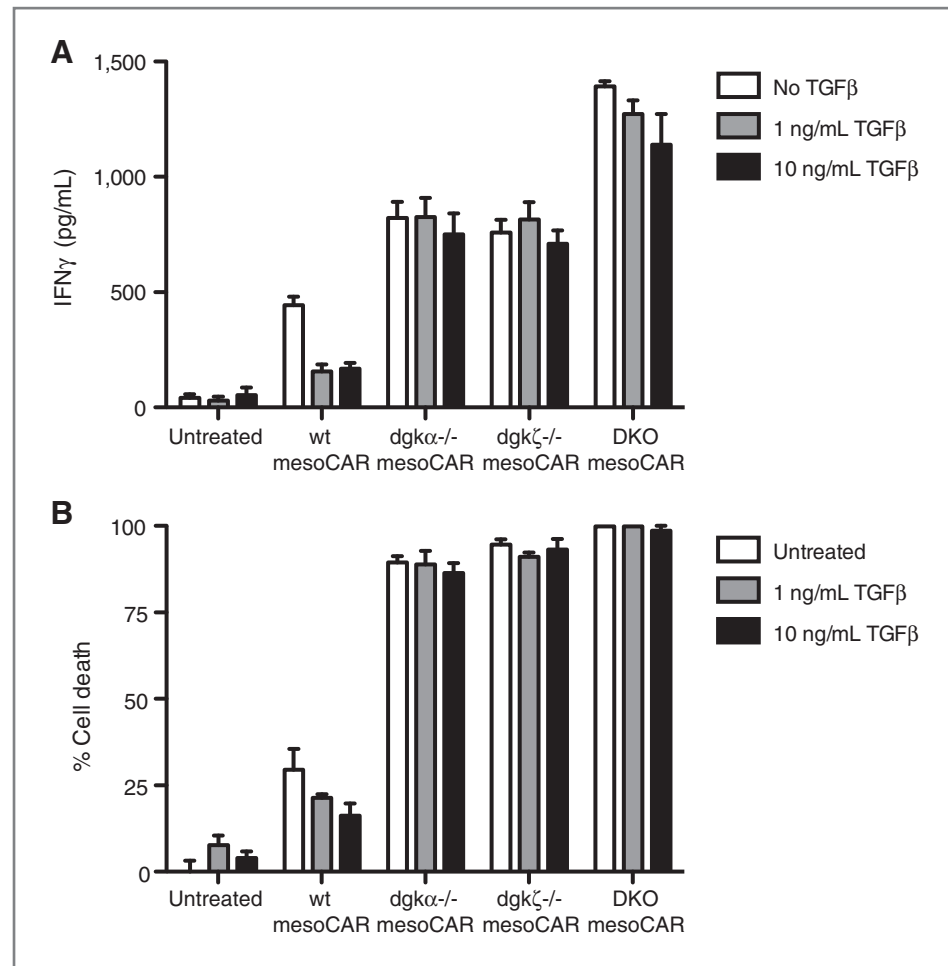
## Discussion

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<sup>+</sup> T cells (12). Although deletion of cbl-b and dgkζ both results in changes downstream of the TCR, dgkζ, and dgkα, 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<sup>+</sup> T cells.

In the studies presented herein, we found that deletion of dgks 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 7.** Dgk-deficient T cells are less inhibited by TGF $\beta$ . A total of  $1 \times 10^5$  naive (CD44<sup>lo</sup>) mesoCAR-transduced T cells of the indicated genotype were incubated with 5,000 AE17meso target cells at a ratio of 40:1 in the absence or presence of TGF $\beta$  at the indicated concentration. A and B, IFN $\gamma$  (A) and cytotoxicity (B) of target cells at 24 hours was assessed as described in Fig. 2. One of the 3 representative experiments is shown. Each data point was conducted in triplicate [ $P < 0.0001$  between wild-type (wt) and all dgk-deficient T cells treated with TGF $\beta$ ].



The role of dgks in limiting the effector function of activated CD8<sup>+</sup> 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 $\zeta$  augmented Erk activation, a phosphorylation event that occurs downstream of DAG formation, after CAR ligation. Deletion of dgk $\zeta$  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 *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<sub>2</sub>, adenosine, and TGF $\beta$  were less able to suppress CD8<sup>+</sup> 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). The

enhanced Ras activation imparted by the loss of dgks might explain how dgk-deficient lymphocytes develop insensitivity to TGF $\beta$ . Because TGF $\beta$  is known to result in the reduced phosphorylation of Itk, a Tec kinase important in PLC $\gamma$ 1 activation (44), and because PLC $\gamma$ 1 is the protein directly responsible for DAG generation in T cells, one could envision that deletion of dgks might directly subvert this TGF $\beta$ -induced signaling alteration. Future studies will address the potential means by which loss of dgks confers insensitivity to TGF $\beta$ , 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 dgk $\zeta$ -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

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**Study supervision:** G.A. Koretzky, S.M. Albelda

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

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