Combined Alloreactive CTL Cellular Therapy with Prodrug Activator Gene Therapy in a Model of Breast Cancer Metastatic to the Brain

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Abstract

Purpose: Individual or combined strategies of cellular therapy with alloreactive CTLs (alloCTL) and gene therapy using retroviral replicating vectors (RRV) encoding a suicide prodrug activating gene were explored for the treatment of breast tumors metastatic to the brain.

Experimental Design: AlloCTL, sensitized to the HLA of MDA-MB-231 breast cancer cells, were examined in vitro for antitumor functionality toward breast cancer targets. RRV encoding the yeast cytosine deaminase (CD) gene was tested in vivo for virus spread, ability to infect, and kill breast cancer targets when exposed to 5-fluorocytosine (5-FC). Individual and combination treatments were tested in subcutaneous and intracranial xenograft models with 231BR, a brain tropic variant.

Results: AlloCTL preparations were cytotoxic, proliferated, and produced IFN-γ when coincubated with target cells displaying relevant HLA. In vivo, intratumorally placed alloCTL trafficked through one established intracranial 231BR focus to another in contralateral brain and induced tumor cell apoptosis. RRV-CD efficiently spread in vivo, infected 231BR and induced their apoptosis upon 5-FC exposure. Subcutaneous tumor volumes were significantly reduced in alloCTL and/or gene therapy–treated groups compared to control groups. Mice with established intracranial 231BR tumors treated with combined alloCTL and RRV-CD had a median survival of 97.5 days compared with single modalities (50–83 days); all experimental treatment groups survived significantly longer than sham-treated groups (median survivals 31.5 or 40 days) and exhibited good safety/toxicity profiles.

Conclusion: The results indicate combining cellular and suicide gene therapies is a viable strategy for the treatment of established breast tumors in the brain. Clin Cancer Res; 19(15); 4137–48. ©2013 AACR.

Introduction

Breast cancer is the most common malignancy in women in the United States and metastasis is a major cause of morbidity and mortality in these patients. With improvements in the control of visceral and bone metastasis, the incidence of brain metastasis is rising (1–3). The progressive neurological disabilities associated with brain metastasis not only impair the quality of life but also decrease the survival. The biology of the brain and blood–brain barrier, and the fact that brain metastases can present as multiple lesions often characterized by nests of infiltrating tumor cells surrounding the larger brain metastases (4, 5) make the management of brain metastases very challenging. Current therapeutic approaches include whole brain radiation therapy and stereotactic radiosurgery usually administered as palliative care (6–8).

Newer approaches include the use of tumor-targeted CTL for adoptive immunotherapy. A unique form of adoptive T-cell therapy for brain tumors involves the use of alloreactive CTL (alloCTL), potent cytotoxic T-cell effectors that are trained to recognize non- or aberrant-self class I and II HLA. They are generated by sensitization of peripheral blood mononuclear cells (PBMC), isolated from a healthy donor, to the HLA primarily acts as tumor-directed antigen in the brain, because its expression, especially class I, is absent on normal
neuroglia, that is, neurons, oligodendrocytes, and astrocytes (12–14). but is highly expressed on brain tumor cells (13, 15–18).

Another new therapeutic approach is the use of replicating virus vectors for gene therapy or oncolytic virotherapy. Recently, it has been demonstrated that retroviral replicating vectors (RRV), unlike their replication-defective counterparts, exhibit sustained transduction in dividing cancer cells and can efficiently transfer transgenes throughout solid tumors (19–24). Glioma-bearing rats show significantly prolonged survival when given RRV coding for the prodrug activator gene, yeast cytosine deaminase (CD), followed by multiple administrations of 5-fluorocytosine (5-FC) prodrug (20–22). The lack of detectable spread of the RRV to normal tissues additionally provides an element of safety in this approach (21, 23, 25).

Both alloCTL adoptive immunotherapy and RRV-mediated prodrug activator gene therapy have been studied individually in patients with primary brain tumors. Notably, the clinical feasibility and safety of intratumoral alloCTL treatment was initially tested in a small pilot study (9, 11), in which 3 recurrent grade 3 glioma patients receiving this treatment exhibited survival longer than expected. These data led to the initiation of a phase 1 dose-escalation study at University of California Los Angeles (UCLA), now accruing patients (www.clinicaltrials.gov, NCT 01144247). As well, RRV encoding cytosine deaminase for prodrug activator gene therapy has advanced to the clinic for recurrent high-grade gliomas (www.clinicaltrials.gov, NCT 01156584 and NCT 01470794).

Clinically, brain metastases are more common than primary brain tumors (26), and represent a dire clinical situation in need of effective therapies. Here, we evaluated both of these cellular and gene therapy approaches, individually and in combination, for the treatment of intracranial breast tumors. Our results show that alloCTL generated to the breast tumor cell line MDA-MB-231 (231) showed cytotoxicity, proliferation, and IFN-γ production in response to coinoculation with the parental target cell line, as well as with 2 metastatic sublines (27, 28). Intracranially placed alloCTL were capable of migrating from one established intracranial tumor focus to another in contralateral brain over a relatively short time span (6 hours) in immune-incompetent mice, and apoptotic tumor cells were observed in proximity to these CTL. Using subcutaneous and intracranial human breast tumor xenograft models, we found that combining prodrug-activating gene therapy with alloCTL adoptive transfer resulted in improved outcome compared to the benefit seen with each therapy individually. Our findings thus support clinical translation of this multimodal approach.

Materials and Methods

Cell lines and HLA expression

The human breast tumor cell line MDA-MB-231 [231; American Type Culture Collection (ATCC), Manassas, VA] was grown in L15 medium supplemented with 10% FBS (Aleken). The brain-seeking 231 subline, 231BR (27), the bone-seeking 231 subline, 231-1833 (28), and 293T human embryonic kidney cells (ATCC) were maintained in Dulbecco’s Modified Eagles Medium supplemented with 10% FBS, sodium pyruvate, and penicillin/streptomycin (Sigma). Class I HLA display was determined on breast cancer cell lines by flow cytometric analysis using RPE-conjugated mouse anti-human HLA-ABC antibody and a mouse RPE-conjugated immunoglobulin G served as isotype control (eBiosciences). The percentage of positive cells and relative antigen density indicated by mean fluorescence intensity (MFI) were determined (13). Specific HLA-ABC types were determined by low-resolution molecular HLA typing (University of California, San Diego, Torrey Pines Clinical Laboratory, San Diego, CA).

Generation of alloCTL-enriched cultures by one-way mixed lymphocyte tumor reaction

PBMC were isolated from leukopaks of healthy donors undergoing leukapheresis (UCLA Institutional Review Board approved protocol) using density gradient centrifugation as previously described (10). Before one-way mixed lymphocyte tumor reaction (MLTR), 231 cells underwent 48-hour incubation in culture medium with 500 IU/mL recombinant human IFN-γ (RD Systems Inc.) to upregulate their HLA expression (13). The stimulator (S) 231-cell monolayers were then washed with PBS and detached with 2 mmol/L EDTA containing 1% bovine serum albumin, washed and inactivated with 7000 rad (X-ray Irradiator, Guilmy Medical, Inc.). Responder (R) PBMC were mixed with inactivated S cells at a R:S ratio of 10:1. The cell mixtures were placed into Aim-V medium (Life Technologies, Inc.) containing 5% heat-inactivated autologous plasma and 60 International Units (IU) of recombinant human interleukin-2 (IL-2)/ml (Proleukin; Novartis). Growth was monitored daily and cultures were fed and maintained as...
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Previously described (10). AlloCTL preparations were used in assays between 12 and 14 days post-MLTR.

Plasmid constructs and viral vector production

The pAC3-yc2 and pAC3-GFP plasmid constructs, encoding amphotropic Moloney murine leukemia virus carrying expression cassettes consisting of an internal ribosome entry site followed by either cytosine deaminase or GFP transgenes, respectively, have been described previously (23, 24).

Viral vector stocks were generated by transfection of 293T cells using the FuGENE6 transfection agent (Roche Molecular Biochemicals) per manufacturer instructions (22). Conditioned medium (viral supernatant) was collected 48 hours after transfection with either the pAC3-GFP or pAC3-yc2 plasmids (23, 24).

Concentrated viral vector supernates were prepared from 293T stable vector producing cells. Conditioned medium was harvested after 72-hour incubation, and clarified by centrifugation before 100-fold volume concentration using a Vivaspin-20 (Sartorius Stedim) per manufacturer protocol. Control supernatant from nontransduced, nonvector producing 293T cells was harvested and concentrated in parallel. Concentrated viral titers were approximately 10^6 transducing units/mL when tested using either 293T or 231BR cells. The titer was determined 48 hours after adding 50 μM of 3′-azido-deoxothymidine (Sigma Aldrich) to the cultured cells to prevent further virus replication as described (20, 29).

For subcutaneous tumors, 10 μL of concentrated vector producing or nonvector producing supernatants were admixed with tumor cells immediately before injection.

Antitumor functionality assessments of alloCTL

**Cytotoxicity assay.** Lytic activity of alloCTL was assessed at various effector-to-target ratios (E:T) by 51Cr release cytotoxicity assays at 12 to 14 days post-MLTR (9). Target cells were labeled with 100 μCi Na251CrO4 (Amersham) and the assay was conducted as previously described (30). Maximal release was obtained from targets incubated with 0.1 M HCl, and spontaneous release was the radioactivity released from targets in assay medium alone. The percentage specific release was calculated by the formula: [(51Cr_{experimental} - 51Cr_{spontaneous})/(51Cr_{maximal} - 51Cr_{spontaneous})] × 100%. Mean specific release ± SEM was determined from triplicate wells.

**Cytokine production.** Human IFN-γ concentrations were measured in clarified supernates collected 48 hours after coincubation of day 12 alloCTL with relevant tumor target cells. Aliquots of the alloCTL from the same preparation either were or were not restimulated at day 12 post-MLTR with 231 or sublines, 231-1833 or 231BR, at a R:S of 10:1. Manufacturer instructions were followed for a Quantikine ELISA kit specific for human IFN-γ (R&D Systems) with a sensitivity of 2 pg/mL (30).

**Immunophenotype and proliferation of restimulated alloCTL.** At 12 days post-MLTR, alloCTL were restimulated with 231 cells or sublines for 72 hours at a R:S ratio of 10:1 in culture medium containing bromodeoxyuridine (BrdU; BD Biosciences). GolgiStop (BD Biosciences) was added to block protein transport during the last 5 hours of culture. Cells were washed and surface stained with APC-conjugated anti-CD4, PeCy5-conjugated anti-CD8 for 15 minutes at 4°C, washed again, fixed, and permeabilized according to instructions for the BD BrdU Flow Kit (BD Biosciences). Cells were then intracellularly stained with phycoerythrin-conjugated anti-IFN-γ and fluorescein isothiocyanate-conjugated anti-BrdU for 30 minutes at 4°C, washed, resuspended, and immunophenotyped using a BD LSRII flow cytometer running BD-DIVA acquisition software. Dot blots were analyzed using FlowJo flow cytometric analytic software.

**In vivo studies with human 231BR xenografts.**

Rag2-/-γc-/- mice were purchased from Taconic Farms (Hudson; ref. 31). All animal experiments were conducted according to institutional guidelines under approved protocols. For subcutaneous tumor model. 231BR cells (2 × 10^6) were coinjected with concentrated RRV-GFP, RRV-CD, or concentrated nonviral control supernate subcutaneously in the right flank of 6-week-old mice. After allowing tumors to establish for 15 days, mice were randomized to treatment groups (n = 4–6). Certain groups were treated with intratumoral alloCTL (1.2 × 10^5). All mice were treated with 5FC (500 mg/kg) intraperitoneally (i.p.) every day from days 21 to 27. Caliper measurements of tumor width and length were conducted every 3 to 4 days. Tumor volumes were calculated for individual animals using the formula: volume = (4/3)π(width/2)^2(length/2). Mean tumor volumes for each treatment group ± SEM were plotted over time.

**Intracranial vector spread.** 100% RRV-GFP transduced 231BR (231BR-GFP) cells were admixed with nontransduced 231BR at 2% or 8% of the total tumor cell inoculum (2 × 10^5/3 μL). Cells were then placed into the right side of the brain through a burr hole, 2 mm lateral and 1 mm anterior to the bregma, at a depth of 3 mm. Groups of mice (n = 3) were sacrificed on days 7, 14, and 20 post-tumor instillation, and brains were harvested. Tumor tissue at the injection site was excised (approximately 5 × 5 × 5 mm^3), minced, and digested in collagenase/dispose (1 mg/mL; Roche) as described (22). Single cell suspensions were stained with fluorescent APC anti-HLA-ABC (eBiosciences) and flow cytometrically analyzed.

**Intracranial alloCTL motility studies.** 231BR cells (2 × 10^6 each) were stereotactically injected into the above coordinates and its enantiomeric position, such that right and left hemisphere tumor foci could develop. Tumors were allowed to establish 18 or 21 days before alloCTL (2 × 10^6) were injected into the left tumor foci. Mice were sacrificed 6 hours later, and brains were harvested, placed into 10% buffered formalin, paraffin-embedded, sectioned, and dianimobenzidine immunostained using rabbit anti-human CD3 (Clone SP7, Genway Biotech Inc.) with hematoxylin counterstain and evaluated by light microscopy.

**Intracranial efficacy studies.** Mice underwent surgery for placement of intracranial cannulas (Plastics One) that were
placed through a burr hole in the right side of the skull at the above coordinates. The cannulas extended 3 mm into the brain and affixed to the skull with resin. Six days later, tumor cells \((2 \times 10^5\) total cells in 3 μL\), consisting of either 100% 231BR, or 98% 231BR cells + 2% 231BR-GFP or 2% 231BR-CD cells, were infused through the cannulas and the animals were placed into various treatment groups. AlloCTL or unstimulated PBMC \((2 \times 10^6/3 \mu L)\) or PBS were infused through the cannulas into the tumor on days 9 and 16 posttumor instillation. All groups of mice \((n = 9-10)\) were treated with 3 cycles of 5 daily i.p. 5-FC \((500 mg/kg)\) injections per cycle beginning on days 12, 26, and 47 posttumor instillation. Mice were monitored for signs of morbidity and weighed every 3 to 4 days for the duration of the experiment.

**Analysis of RRV biodistribution**

To harvest brains, coronal cuts were made at the site of cannula implantation and 4 mm posterior to that cut. The anterior sections were snap-frozen for quantitative real-time PCR (qRT-PCR) to determine RRV biodistribution; the posterior 4 mm sections were fixed in formalin, paraffin-embedded, and the blocks were sectioned \((5 \mu m)\) before staining with hematoxylin and eosin (H&E).

Genomic DNA was extracted using the DNeasy tissue kit (Qiagen) from tissues (liver, lung, spleen, kidney, bone marrow, and brain) of all long-term survivors and representative animals derived from control and experimental groups that succumbed to tumor after intracranial treatment with RRV and/or alloCTL. To detect integrated RRV sequences, qRT-PCR amplification of genomic DNA was carried out in duplicate with TaqMan Universal PCR Master Mix (PE Applied Biosystems) using a My-iQ2 Biorad Thermal Cycler. The primers and probes were designed to target the 4070A amphotropic env gene \((4070A-F, 5'$-GCGGAC-CGGGACTTTTGA-3', 4070A-R, 5'$-ACCCCGACTTTAGTATGC-3'; probe, FAM-CAGGGCACACGTAAAA-NFQ).\)

Human RNase P \((hrNase P)\) and mouse β-actin \((m\beta\text{-actin})\) were also quantified as reference genes using TaqMan RNase P Detection Reagents and Custom TaqMan Gene Expression Assays (Applied Biosystems), which is designed to target mouse β-actin gene \((\beta\text{-actin-forward, } 5'$-GGTCTTACCA-CAGGCCATTGT-3'; \beta\text{-actin-reverse, } 5'$-CCTGTAAGTGCCG-GACAGTG-3'; probe, FAM-CCCGCTCAGGAGTCC-NFQ). A standard curve for copy number was prepared by amplifying serial dilutions of plasmid pAC3-γCD2 in a background of genomic DNA from nontransduced cells.

**Statistical analyses**

GraphPad Prism software (version 4; GraphPad Software) was used to analyze the in vitro and in vivo data. The Student \(t\) test determined significant differences between the percentage lysis of target tumor cells with and without the addition of anti-HLA; \(P\)-values \(\leq 0.05\) were significant. Experimental and control groups for the in vitro restimulation experiments measuring proliferation, immunophenotype, and cytokine production, and the in vivo subcutaneous tumor volume comparisons at given times were compared using a two-way ANOVA with Bonferroni correction and \(P\)-values \(\leq 0.05\) divided by the number of comparisons \((3)\) in experiments described in Figs. 2 and 4 were considered significant. Median survival times (MST) from the Kaplan-Meier curves were analyzed by nonparametric log-rank tests.

**Results and Discussion**

**HLA class I on breast tumor cells and upregulation with IFN-γ**

Flow cytometric analysis showed that nearly all \((94–99%)\) of the cells in the parental 231 line, as well as 231-1833 and 231BR sublines, expressed HLA class I and their incubation with IFN-γ resulted in 1.4- to 1.5-fold increases in MFI (Supplementary Table S1). HLA-ABC expression after IFN-γ induction was highest on the parental 231 cell line \((MFI 1402 \pm 49.7)\) compared with slightly lower expression by each of the sublines, thus making the parental 231 cells most desirable as stimulator cells in a one-way MLTR for alloCTL generation. The low-resolution molecular type of HLA-ABC loci in 231 cells was HLA-A (17, 231, C) and in 231BR cells was HLA-A*02, B*41, 40, and C*17, 02 (Supplementary Table S2).

**AlloCTL exhibit antitumor function in vitro**

The standard method for generating alloCTL is by one-way mixed lymphocyte reaction where inactivated stimulator PBMC from the cancer patient are combined with responder PBMC from healthy donors genetically distinct from the patient (10). However, because PBMC are not available from the patient from whom 231 breast tumor cells were derived, to generate alloCTL we used a one-way MLTR where 231 tumor cells were incubated with IFN-γ to upregulate HLA class I expression as stated earlier, then inactivated and used as stimulator cells with responder PBMC isolated from HLA-mismatched allogeneic donors (33).

Four different donors provided PBMC that served as sources of precursor alloCTL for these experiments. All donors were mismatched at 4 to 5 class I HLA-ABC antigens compared to that displayed by the 231 stimulator cells (Supplementary Table S2). The HLA types of alloCTL used for in vitro and in vivo experiments are provided and associated with data in particular figures. Note these preparations will contain CTL directed to minor tumor-associated antigens as well as HLA, albeit the precursor frequency for CTL within PBMC would be higher to major than to minor antigens (34–36).

To determine if the alloCTL made by MLTR had antitumor function in vitro, we conducted cytotoxicity assays on day 14 following one-way MLTR. AlloCTL effector cells were mixed with \(^{51}\text{Cr}\)-loaded 231, 231-1833, or 231BR target cells at a range of E:T ratios for 4 hours. As expected, and in a dose-dependent manner, the ability of alloCTL to kill each of the 3 target cell lines was similar; there were no significant differences in the percentages of lysis obtained at each E:T ratio (Fig. 1). Furthermore, the cytosis induced by alloCTL to 231 and its sublines is in part HLA-restricted, as indicated by significant inhibition of cytotoxicity (33.3–41.3%...
BrdU incorporation between CD4<sup>+</sup> T cells. In contrast, there were no significant differences in proliferation. Lymphocyte proliferation occurred equally well with BrdU upon restimulation and thus proliferated upon stimulation with relevant target cells were significantly higher (P < 0.017) compared to those that were unstimulated (Fig. 2A). The percentages of CD8<sup>+</sup> T cells that incorporated BrdU upon restimulation and thus proliferated upon stimulation with relevant target cells were significantly higher (P < 0.017) compared to those that were unstimulated (Supplementary Table S1). The reduction in percentage lysis of 231 and variants, P < 0.006 upon antibody-mediated blockade of HLA. We replicated the cytotoxicity findings with alloCTL (with and without blocking antibody to HLA) using 2 other breast cancer cell lines, BMC Brain2 and MDA-MB-361 (unpublished data).

**AlloCTL subset proliferation and IFN-γ production following restimulation**

The ability of the CD4<sup>+</sup> and CD8<sup>+</sup> T-cell subsets within the alloCTL preparations to proliferate and produce IFN-γ following restimulation with 231, 231-1833, or 231BR target cells was evaluated. AlloCTL that had been restimulated on day 12 after one-way MLTR for 72 hours at a 10:1 R:S ratio with one of the 3 relevant cell lines, or left unstimulated, were surface stained for CD4 and CD8, and then fixed, permeabilized, and stained for proliferation marker (BrdU) and proinflammatory IFN-γ production. Figure 2 shows data obtained from a representative alloCTL preparation. The percentages of CD8<sup>+</sup> T cells that incorporated BrdU upon restimulation and thus proliferated upon stimulation with relevant target cells were significantly higher (P < 0.017) compared to those that were unstimulated (Fig. 2A). Lymphocyte proliferation occurred equally well with parental 231 cells and the metastatic sublines, 231-1833 or 231BR. In contrast, there were no significant differences in BrdU incorporation between CD4<sup>+</sup> T cells that were restimulated versus those that were unstimulated (Fig. 2A), indicating lack of CD4<sup>+</sup> subset proliferation, regardless of whether restimulation was conducted with the parental 231 cells or the 231-1833 or 231BR subline counterparts.

The total percentages of CD8<sup>+</sup> T cells were significantly higher (P < 0.017) within alloCTL populations that were restimulated with relevant target cells (51.8–56%) compared to unstimulated alloCTL (30.9%; Fig. 2B). Because the CD8<sup>+</sup> T cells proliferated when restimulated with 231, 231-1833, and 231BR cells and the CD4<sup>+</sup> cells did not, the total percentages of CD4<sup>+</sup> T cells were significantly reduced (P ≤ 0.017) compared to the unstimulated counterpart (Fig. 2C). The data show that restimulation of alloCTL with 231-1833, or 231BR breast cancer cells results in a significant shift in T-cell subsets that make up the alloCTL culture.

Expression of IFN-γ following 72-hour restimulation with 231, 231-1833, and 231BR, compared to unstimulated cells, was also evaluated. Protein transport was blocked during the last 5 hours of culture to allow intracellular accumulation of IFN-γ. Expression of IFN-γ was significantly higher (P ≤ 0.017) within CD8<sup>+</sup> T cells that were restimulated with 231, 231-1833, or 231BR, compared to cells that were unstimulated (64.6, 45.1, 43.8 vs. 27.5, respectively; Fig. 2D). Production was more robust when restimulation was conducted with the parental cells versus the sublines, correlating with the relative antigen densities (MFIs) of cell-surface class I HLA (Supplementary Table S1), suggesting that this parameter may have some influence over response. In contrast, production of IFN-γ in CD4<sup>+</sup> T cells was not affected following restimulation with any of the cell lines (data not shown).

To confirm the earlier observations, IFN-γ secretion by alloCTL at day 12 post-MLTR was analyzed from supernates collected 48 hours following restimulation with relevant target cells, and compared with IFN-γ secreted by unstimulated alloCTL. The average secretion of IFN-γ by restimulated alloCTL coincubated with 231, 231-1833, and 231BR (17.2 ± 1.2, 16.9 ± 0.50, and 13.2 ± 0.97 ng/mL, respectively) was significantly elevated (P < 0.017) compared to that of unstimulated alloCTL (7.4 ± 0.21 ng/mL).

**AlloCTL migrate through and to distant tumor foci in vivo and induce apoptosis**

We examined the migratory capacity of alloCTL in the context of multifocal intracranial tumors. As shown in Fig. 3A, bilateral 231BR tumors were established intracranially in the right and left hemispheres of Rag2<sup>−/−</sup>γc<sup>−/−</sup> mice, and allowed to grow for 21 days. AlloCTL specific for 231 HLA were then stereotactically injected into the tumor bed on the right side (Fig. 3A), and coronal sections of brain were prepared 6 hours later and immunostained with a human CD3-specific antibody (Fig. 3B–G). Figure 3B shows a representative low power photomicrograph of a coronal brain section with bilateral tumor foci. Intermediate power magnification of the tumor (Fig. 3C) in the right side of the brain shows numerous DAB-stained (rust-color) human CD3<sup>+</sup> T lymphocytes (black arrows) penetrating the tumor mass at the instillation site. A higher power magnification photomicrograph (Fig. 3D) from the same area shows...
CD3$^+$ cells (black arrows) in proximity to a necrotic tumor cell with a fragmented nucleus indicating cell death (white arrows). In addition, CD3$^+$ alloCTL have trafficked and localized to established tumor in contralateral brain (Fig. 3E–G). Rust-colored CD3$^+$ cells are shown having permeated the tumor mass (t) on the left side and also appear in perivascular spaces (asterisks), which is a path for tumor invasion (Fig. 3E), whereas no CD3$^+$ cells are visible in normal brain. Intermediate and higher power magnification photomicrographs (Fig. 3F and G) similarly show larger, activated CD3$^+$ T-cell phenotypes (black arrows) in close juxtaposition to apoptotic tumor cells (white arrows), indicating cytotoxic functionality may be retained by the trafficking alloCTL.

Cellular and gene therapies show therapeutic benefit in subcutaneous 231BR xenograft models

To determine if cellular and gene therapy approaches individually or in combination would be efficacious in vivo, Rag2$^{-/-}$ γc$^{-/-}$ mice were injected subcutaneously with 231BR (initial inoculum 2 × 10^6 cells) admixed with RRV-GFP, RRV-CD, or control nonviral supernatant. Tumors were allowed to establish for 15 days before some experimental groups were treated with intratumoral alloCTL (1.2 × 10^6 cells). All groups (n = 4–6) received the suicide gene therapy prodrug, 5-FC (500 mg/kg i.p.) on days 21 to 27. Tumor volumes monitored over time showed significant differences between the control and experimental groups starting from day 25 ($P < 0.017$; Fig. 4). The
required an natant and/or alloCTL. This experimental paradigm to establish for 1-week before infusing with RRV-CD super-
relevant scenario was tested where tumors were allowed in preparation). In those experiments, a more clinically
subcutaneous U-87MG glioma xenografts (K. Haga et al., similarly efficacious findings in a second tumor model using
2 control groups that received 231BR with nontherapeutic
RRV-GFP or control nonviral naive 293 cell supernatant at
the start of the experiment both exhibited progressive tumor
growth in vivo, with average group tumor volumes at day 28
of 759 ± 243 mm³ and 659 ± 114 mm³, respectively.
Therapeutic benefit was noted for all experimental treat-
ment groups (groups that received RRV-CD, alloCTL, or a
combination of the 2) compared to mice that received
tumor with concentrated nonviral supernatant. Tumors
treated with RRV-CD alone, without alloCTL injection,
showed significant (P ≤ 0.017) tumor growth inhibition
upon application of 5-FC prodrug treatment between days
25 and 27 compared to control groups. Tumors treated with
nonviral supernatant but receiving intratumoral alloCTL
also showed significant (P ≤ 0.017) tumor growth inhibi-
tion compared to control groups, following cellular thera-
policy, but as expected, did not show response to prodrug
administration. Notably, the greatest reduction in mean
tumor volumes was obtained when intratumoral cellular
therapy with alloCTL was combined with prodrug activator
gene therapy with RRV-CD/5-FC (P ≤ 0.017). We obtained
similarly efficacious findings in a second tumor model using
subcutaneous U-87MG glioma xenografts (K. Haga et al.,
in preparation). In those experiments, a more clinically
relevant scenario was tested where tumors were allowed
to establish for 1-week before infecting with RRV-CD super-
natant and/or alloCTL. This experimental paradigm required an in situ transduction of preestablished tumor to
obtain a beneficial effect from administered prodrug. Again,
efficacy was observed for the individual as well as the
combination therapies.

RRV effectively spreads through intracranial breast
tumor xenografts

RRV can achieve significantly enhanced transduction
compared to their replication defective counterparts (19).
RRV exhibit an amplification process that is inherent to the
wild-type virus life cycle, so although viral replication
kinetics and lag times for exponential growth phase repli-
cation can vary, even a slower replication will allow pro-
gressive vector spread within a tumor until prodrug is given.
We have previously demonstrated RRV replication and
gene expression in subcutaneous mammary tumors (29).
To evaluate RRV spread in intracranial breast tumors,
Rag2−/−γc−/− mice were intracranially injected with 2 ×
107 naïve 231BR cells mixed supplemented with either 2% or
8% RRV-GFP transduced 231BR cells (231BR-GFP). Transduction of the tumor by RRV was evaluated by flow
cytometry on days 7, 14, and 20 post-tumor instillation (Fig. 5A and B). To distinguish the human 231BR tumor
cells from mouse brain cells, enzyme-digested tumor
tissues were stained with fluorescently labeled human-
specific anti-HLA-ABC and the percentages of GFP+ cells
within this population were quantified. When the initial
tumor inoculum contained either 2% or 8% 231BR-GFP
cells, by day 7 post-tumor instillation, approximately

![Figure 3. AlloCTL trafficking in vivo through and to tumor foci.Brains were collected 6 hours after alloCTL injection into a 3-week established 231BR tumor focus as seen by human anti-CD3 positivity in ipsilateral and contralateral tumor foci within immune incompetent mouse brain. A, coronal brain section schematic of tumor placement and alloCTL at (ipsilateral injection site. B, gross coronal brain section that was DAB immunostained with anti-CD3 that shows 2 tumor foci (ipsilateral-red and contralateral-blue boxes, respectively). C, low power photomicrograph of rust-colored CD3+ cells (black arrows) in tumor (l) focus #1 and alloCTL instillation site. D, higher power view showing 3 CD3+ cells in juxtaposition to an apoptotic tumor cell (white arrow). E–G, tumor focus #2 in brain contralateral to alloCTL injection. E, low power photomicrograph shows abundant CD3+ cells (black arrows) primarily within the tumor mass (l) but not in normal brain (n) located in the hemisphere opposite to alloCTL instillation. Yellow asterisks show pockets of tumor cells within Virchow–Robin perivascular spaces. (F) Intermediate and (G) high-power magnification photomicrographs at that site again show apoptotic tumor cells (white arrows) in proximity to CD3+ cells (black arrows). The brain sections are counterstained with hematoxylin. Representative photomicrographs are shown from 1 of 10 mice; 2 experiments were analyzed after H&E staining and 2 experiments after DAB immunostaining. Bars = 0.8 mm in B, 30 μm in C, 8 μm in D, 35 μm in E, 15 μm in F, and 10 μm in G.](https://www.aacrjournals.org/doi/10.1158/1078-0432.CCR-12-3735)
concentrated supernate (black diamonds) and the following experimental
with Bonferroni posttests between mice who received MDA-231BR
tumor injection (upward arrows). Data are represented as the average
treated with 500 mg/kg 5-FC i.p. from days 21 to 27 post subcutaneous
(downward arrow; white circles and white diamonds). All mice were
establish intracerebral tumors in 7 groups (n = 6 mice/group).

57% of the human class I HLA-ABC+ tumor cells had been transduced with RRV-GFP. Subsequently, tumor transduction levels rose to ~75% to 83% at 20 days following tumor instillation. Figure 5B shows representative flow cytometric dot-plots from day 14, indicating 71.2 ± 3.7% and 78.9 ± 1.1% GFP transduction of HLA-ABC+ human 231BR cells after initial tumor inoculation with 2% and 8% 231BR-GFP cells, respectively.

Individual and combined cellular (alloCTL) and gene (CD) therapies are safe and exhibit therapeutic benefit in intracranial 231BR xenograft models

The efficacy of combined cellular alloCTL immunotherapy with RRV prodrug activator gene therapy was tested in Rag2-/-γc-/- mice bearing established intracranial 231BR tumors. Indwelling intracranial cannulas were used to establish intracerebral tumors in 7 groups (n = 9–10) of mice, consisting of either 100% 231BR or 98% 231BR mixed with either 2% 231BR-GFP or 2% 231BR-CD cells (Fig. 6). Control groups of mice were injected with non-therapeutic 231BR-GFP tumor cells instead of 231BR-CD, or were infused with PBS or unstimulated PBMC in place of alloCTL. On days 9 and 16 post-tumor instillation, effector alloCTL, or control PBMC, or PBS was infused into the established tumor bed through the cannula. All mice were treated with 5-day cycles of 5-FC (500 mg/kg, i.p.); up to 3 cycles were possible, spaced 2 or 3 weeks apart beginning on day 12 post-tumor instillation (Fig. 6A). Kaplan–Meier survival plots and MST for the control and experimental groups are shown in Fig. 6B. The MST of the untreated control group (no RRV, no alloCTL effector cells, group 1) was 31.5 days, whereas the MST of the sham-treated control group given nontherapeutic 231BR-GFP vector producing cells with the tumor inoculum and unstimulated PBMC was 40 days (group 3). The MSTs for mice treated with individual experimental modalities ranged from 50 to 83 days. The MSTs of the 2 groups receiving alloCTL therapy alone (with RRV-GFP infection or no RRV infection) were 65 and 83 days respectively (groups 4 and 2), whereas the MSTs of the 2 groups receiving prodrug activator gene therapy alone (with PBMC or PBS, instead of alloCTL) and up to 3 prodrug cycles were 50 and 57 days, respectively (groups 6 and 5). Statistical significance was reached (P < 0.05) for all of the individual immune or gene therapy–treated groups compared to each of the control groups (Fig. 6C). The MST of the group receiving the combined intratumoral effector alloCTL and RRV prodrug activator gene therapies (group 7) was 97.5 days, thus exhibiting a highly statistically significant survival benefit compared to each of the control groups (Fig. 6C). Furthermore, the group receiving combined immune and gene therapies showed a statistically significant additional survival advantage compared to those groups receiving gene therapy alone (P < 0.0001 and 0.0123, groups 7 vs. 6 and 5, respectively). The combination treatment also showed a trend toward statistical significance compared to the groups receiving alloCTL cellular therapy alone (P < 0.0511 and 0.6380, groups 7 vs. 4 and 2, respectively).

The individual animal weights of Fig. 6 animals were monitored every 3 to 4 days and plotted over the duration of the experiment as one indication of treatment toxicity and/or morbidity due to tumor progression. The plots for the control groups, and the single or combined modality groups are shown starting on day 12 to 106 with prodrug cycle administration indicated on the abscissa (Supplementary Fig. S1). The plots not only show the extension of survival of animals in the experimental groups, but as well, indicate the health of the 11 long-term survivors present in groups 2, 4, 5, and 7. Except in sham-treated control animals (groups 1 and 3), generally little to no weight loss occurred during or shortly after the first prodrug cycle and second alloCTL infusion. Progressive weight loss at later time points was likely associated with higher tumor burden, and notably, stabilization or reversal of weight loss was observed following prodrug administration cycles in individual animals in groups 2, 5, 6, and 7; this was most apparent after the last prodrug cycle in groups 2 and 7, and most of those animals went on to thrive as indicated by stable or increasing weight gains. Thus, treatment with intracranial infusions of alloCTL as well as prodrug administration in RRV-CD–treated animals both contributed positively to overall clinical response to therapy as measured by average body weight, and mice treated with combined immunogene therapy had the highest overall average weight throughout the course of treatment.

Histopathologic findings of the brains from the long-term survivors (n = 11, from Fig. 6 experimental groups 2, 4,
5, and 7) showed no evidence of tumor in H&E-stained sections, whereas there was consistent presence of tumor in animals that succumbed before the end of the survival experiment (Supplementary Fig. S2).

To monitor the spread of RRV-GFP or RRV-CD within the brain and to extratumoral sites, genomic DNA was extracted from brain/brain tumor, lung, liver, spleen, kidney, and bone marrow tissues, then analyzed by qRT-PCR using primers and probe sequences specific for the 4070A amphotropic envelope. Data are shown as RRV copy number/5 × 10^4 cells and calculated as detailed in Supplementary Table S3. In the long-term survivors from the experimental treatment groups (Fig. 6), there were low or no detectable levels of RRV signals, which correlated with the apparent absence of tumor by histopathology. In tumor-bearing animals, there were also higher levels of virus; representative data are shown for one animal in group 7 that succumbed to tumor on day 90. As expected in immunodeficient animal models, RRV signals indicating extratumoral vector spread were largely restricted to hematopoietic tissues, and in other tissues, the lymphoid cell content within them (37).

Overall, our results show proof-of-concept that a unique combination regimen, consisting of cellular therapy with alloCTL and gene therapy using RRV encoding a prodrug activator gene, represents a promising strategy to treat breast tumors metastatic to the brain. We postulate that the mechanisms at work benefiting the combined treatment might involve alloCTL favorably improving the dissemination of RRV to metastatic foci. This could either be by (i) adsorption of virus particles to the surface of trafficking T cells that then transduce tumor cells, that is, a “hitchhiking” mechanism without involving actual infection of T cells (38), or (ii) a small percentage of alloCTL that become transduced and then act as motile vector producing cells to carry the vector to other tumor foci. Suicide gene therapy may also augment the immediate effects of adoptive cellular therapy, because tumor cell injury (combined apoptosis + lysis) induced by effector T cells or suicide gene therapy could improve endogenous immune function by presentation of fragmented tumor cells and antigens to T cells. It is possible that further therapeutic gains can be realized, particularly in immunodeficient xenograft models, by administering additional cycles of prodrug, as previously reported (20), and by adjusting the timing and sequencing of alloCTL infusion versus prodrug administration. Although the xenograft model used in these studies did not allow a determination of the degree of inflammation induced by the viral vector interacting with endogenous immune cells in the brain, prior studies in a syngeneic animal model indicate that this is not a problem (21). In fact, recent studies in syngeneic intracranial glioma models show that an intact endogenous immune system can contribute to complete tumor eradication after RRV-mediated prodrug activator gene therapy, as well as to restriction
Figure 6. Immunogene treatment schema, control, and experimental treatment groups, Kaplan–Meier survival plots with median survival times. A, the treatment schema along with the 7 control or experimental groups are detailed. Mice were first surgically implanted with cannulas. Six days later they were intracranially infused with either nontransduced 231BR cells or 2%/98% mixtures of RRV-transduced/nontransduced 231BR cells. The transduced 231BR cells had RRV coding for either nontherapeutic GFP marker gene (control) or therapeutic CD gene. AlloCTL (therapeutic cells) or PBMC (unstimulated control cells) were infused into the tumor site on days 9 and 16 post-tumor infusion. All animals received the nontoxic prodrug, 5-FC, which was intraperitoneally injected daily for 5 days (one cycle; up to 3 cycles were possible spaced 2–3 weeks apart). Groups 1 and 3 were controls for the suicide gene therapy and cellular therapy treated groups. The experimental groups testing cellular therapy with alloCTL were groups 2 and 4. The experimental groups testing suicide gene therapy were groups 5 and 6. The combination immunogene therapy was group 7. B, the Kaplan–Meier survival plots along with the MST are shown for all 7 control or experimental groups (n = 9–10/group) as observed to day 112 posttumor injection. Long-term survivors were obtained for all individual or combined cellular and gene therapeutic groups. The P values of significance (P < 0.05) calculated by nonparametric log-rank (Mantel–Cox) tests were as follows from this singular experiment:

<table>
<thead>
<tr>
<th>Groups</th>
<th>Treatments</th>
<th>P value</th>
</tr>
</thead>
<tbody>
<tr>
<td>Groups 1,2</td>
<td>PBS vs. alloCTL</td>
<td>0.0011</td>
</tr>
<tr>
<td>Groups 1,4</td>
<td>PBS vs. GFP+alloCTL</td>
<td>0.0365</td>
</tr>
<tr>
<td>Groups 3,2</td>
<td>GFP+PBMC vs. alloCTL</td>
<td>0.0038</td>
</tr>
<tr>
<td>Groups 3,4</td>
<td>GFP+PBMC vs. CD+alloCTL</td>
<td>0.0101</td>
</tr>
<tr>
<td>Groups 1,5</td>
<td>PBS vs. CD</td>
<td>0.0078</td>
</tr>
<tr>
<td>Groups 1,6</td>
<td>PBS vs. PBMC+CD</td>
<td>0.0051</td>
</tr>
<tr>
<td>Groups 3,5</td>
<td>GFP+PBMC vs. CD</td>
<td>0.0093</td>
</tr>
<tr>
<td>Groups 3,6</td>
<td>GFP+PBMC vs. CD+PBMC</td>
<td>0.0004</td>
</tr>
<tr>
<td>Groups 1,7</td>
<td>PBS vs. alloCTL+CD</td>
<td>0.0001</td>
</tr>
<tr>
<td>Groups 3,7</td>
<td>GFP+PBMC vs. CD+alloCTL</td>
<td>0.0001</td>
</tr>
<tr>
<td>Groups 5,7</td>
<td>CD vs. CD+alloCTL</td>
<td>0.0123</td>
</tr>
<tr>
<td>Groups 6,7</td>
<td>CD+PBMC vs. CD+alloCTL</td>
<td>0.0001</td>
</tr>
</tbody>
</table>

Three separate survival experiments were conducted; 2 pilot studies with n = 4 to 6/group gave similar findings.
of systemic viral spread to normal tissues (23). Activation of endogenous antitumoral immune responses due to destruction of the immunosuppressive tumor environment and release of immunostimulatory cytokines from adoptively transferred alloCTL may also contribute to further therapeutic benefit. As alloCTL and RRV therapies have now individually reached the clinical testing stage, we can envision the clinical design for combination immunogene therapy of breast cancer metastatic to brain to be feasible, as shown in Supplementary Fig. S3. In summary, continued preclinical and clinical investigation of combined, local cellular, and gene therapy regimens are warranted for brain metastases.

**Disclosure of Potential Conflicts of Interest**

N. Kasahara has ownership interest (including patents) and is a consultant/advisory board member of Tocagen Inc. No potential conflicts of interest were disclosed by the other authors.

**Authors’ Contributions**


Acquisition of data (provided animals, acquired and managed patients, provided facilities, etc.): M.J. Hickey, K.L. Erickson, H. Soto, ET. Ha, S. Kamijima, M. Takahashi, Y. Kato, B.M. Mueller.


**References**


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