

Salmonella typhimurium engineered to produce CCL21 inhibit tumor growth

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Received: 3 April 2008 / Accepted: 24 June 2008 / Published online: 17 July 2008
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Abstract Intravenously-applied bacteria tend to accumulate in tumors and can sporadically lead to tumor regression. Systemic administration of attenuated *Salmonella typhimurium* is safe and has shown no significant adverse effects in humans. The purpose of this study was to test the hypothesis that engineering *S. typhimurium* to express a chemokine, CCL21, would increase anti-tumor activity. We engineered an attenuated strain of *S. typhimurium* to produce the chemokine CCL21. Attenuated *S. typhimurium* expressing CCL21 significantly inhibited the growth of primary tumors and pulmonary metastases in preclinical models of multi-drug-resistant murine carcinomas, while control bacteria did not. Histological analysis of tumors showed marked inflammatory cell infiltrates in mice treated with CCL21-expressing but not control bacteria. Levels of cytokines and chemokines known to be induced by CCL21 [e.g., interferon- γ (INF γ), CXCL9, and CXCL10] were significantly elevated in tumors of mice treated with CCL21-expressing but not control *S. typhimurium*. The anti-tumor activity was found to be dependent on CD4- and CD8-expressing cells, based on antibody-mediated in vivo immuno-depletion experiments. Anti-tumor activity was achieved without evidence of toxicity. In summary, chemokine-expressing, attenuated bacteria may provide a novel approach to cancer immunotherapy for effective and well-tolerated in vivo delivery of immunomodulatory proteins.

Keywords *Salmonella* · CCL21 · Cancer therapy · Bacterial therapy · Tumor targeting

Introduction

Studies that have used bacteria to treat tumors have been documented as far back as 1893 [1], but the significant toxicity associated with this strategy precluded widespread use [1–3] and has thus far been limited to bacillus Calmette-Guerin for the treatment of superficial bladder carcinomas [4, 5]. Recently, interest in this strategy of employing bacteria to target tumor cells has received heightened interest, particularly with regard to the use of attenuated strains of *Salmonella typhimurium*. *S. typhimurium* are facultative anaerobic bacteria that are able to accumulate within tumors in vivo to concentrations of up to 10^9 colony forming units (CFU) per gram of tumor tissue [6, 7]. This tumor accumulation may be explained by a variety of mechanisms, such as increased access to nutrients provided by rapidly growing tumor and necrotic cells, adaptation of *S. typhimurium* to growth in hypoxic areas, lack of bactericidal activity of macrophages and neutrophils due to reduced oxygen within tumor areas, suppression of the immune system in tumor regions due to secretion of cytokines such as TGF- β , and absence of circulating antibodies and complement factors due to irregular vasculature and positive pressure within tumors.

Recently, a lipid A-negative strain of *S. typhimurium* has been used in phase I clinical trials [8, 9]. Although *S. typhimurium* was found to be safe at doses up to 10^9 CFU/m² body surface area in these studies, and the bacteria did indeed accumulate within tumor tissues, no significant anti-tumor activity was observed. Because attenuated *S. typhimurium* display an excellent safety profile [7, 10]

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and are able to synthesize functional cytokines at high levels [11], we have employed these facultative anaerobic bacteria to express proteins with anti-tumor or immunomodulatory activities, testing them in mouse tumor models [12].

In this study, we engineered an attenuated strain of *S. typhimurium* to stably express the cytokine CCL21. CCL21, also known as 6CKine or secondary lymphoid tissue chemokine, is a member of the chemokine family of small secreted proteins that controls the migration of lymphocytes [13], dendritic cells [14], T [15] and natural killer (NK) cells [16]. The activities of CCL21 suggest that it may induce the colocalization of naïve T cells and antigen-stimulated dendritic cells, which may help to overcome tumor-induced immunosuppression, thereby leading to an effective cell-mediated immune response and subsequent tumor elimination. Several studies demonstrated the feasibility of such an approach to use this chemokine to induce tumor rejection [17–19], but were limited to intratumoral injections. In addition, CCL21 appears to have antitumor functions that do not depend on the immune system, but are mediated through the binding of the chemokine receptor CXCR3 [20], including angiostatic properties [21–24]. In this study, we used an attenuated strain of *S. typhimurium* to deliver the immunostimulatory cytokine CCL21 to tumors and show that CCL21 expression results in inhibition of tumor growth without significant toxicity in murine models.

Materials and methods

Animals, bacterial strains, cell lines, and plasmid vectors

Female mice were purchased at about 6–8 weeks of age from The Jackson Laboratory (Bar Harbor, ME, USA). All animal experiments were performed according to the National Institutes of Health *Guide for Care and Use of Experimental Animals* and approved by the Animal Care Committee of the Burnham Institute for Medical Research (#AUF 04-152). Attenuated lipid A-negative *S. typhimurium* ($\text{pur}^-/\text{msb}^-$) were generated as previously described [6]. The D2F2 cell line was a gift from Dr Wei-Zen Wei (Karmanos Cancer Institute, Wayne State University, Detroit, MI, USA); CT-26 and B16 melanoma cells were purchased from the American Type Culture Collection (ATCC; Manassas, VA, USA). cDNAs encoding luciferase (pGL3; Promega Corporation, Madison, WI, USA), enhanced green fluorescent protein (pEGFP; Clontech, Mountain View, CA, USA), or murine CCL21 (InvivoGen, San Diego, CA, USA), were inserted into pGEN206. This vector is a slightly modified form of pGEN222 [25] in which the *parA* locus has been exchanged with a *parM* and a *parR* locus. In addition, a *repA* locus has been introduced between the origin of replication and the multiple cloning sites.

Immunoblotting

Protein expression of pGEN206-CCL21 was demonstrated by Western blotting of 10^9 CFU *S. typhimurium* (expressing nothing or pGEN206-CCL21) and their respective culture supernatants after resuspending in Laemmli sample buffer and running an SDS-PAGE followed by immunoblotting with a polyclonal goat anti-CCL21 antibody (Abcam, Cambridge, MA, USA).

Mouse experiments

Tumor cell infectivity was performed via a standard gentamycin protection assay. For experiments involving primary tumors, BALB/c mice ($n = 5$) were challenged subcutaneously (SC) with 1.25×10^5 D2F2 breast carcinoma cells and treated as indicated with PBS or 5×10^6 CFU of indicated *S. typhimurium* per mouse after 14 days. Volumes of SC tumors were measured in two dimensions and calculated as $\text{length}/2 \times \text{width}^2$. Mice were challenged with a lethal dose of 2.5×10^5 CT-26 colon carcinoma cells and treated intravenously (IV) with either PBS or indicated *S. typhimurium* (5×10^6 CFU *S. typhimurium* per mouse) after 9, 14 and 19 days. In experiments in which pulmonary metastases were established, BALB/c mice ($n = 8$) were first injected IV with 5×10^4 D2F2 breast carcinoma cells and then treated IV with PBS or indicated *S. typhimurium* (5×10^6 CFU *S. typhimurium* per mouse) after 6, 13, and 20 days. After 50 days, lungs were weighed (mean \pm SE) and examined for metastases using the following scale to determine the percentage of lung surface covered by metastases: 0 = 0%, 1 = <20%, 2 = 20–50%, 3 = >50%. For experiments involving antibody-mediated depletion/blocking, C57/B16 mice ($n = 5$) were challenged SC with 1.25×10^5 B16 melanoma cells and treated IV after 7 days with PBS, *S. typhimurium*, *S. typhimurium* plus empty vector, or *S. typhimurium* plus CCL21-bearing vector, including groups of mice that were treated intraperitoneally (IP) with antibodies (500 $\mu\text{g}/\text{mouse}$; [26]) against CD4+ T cells (anti-CD4 clone GK1.5, ATCC; Manassas, VA, USA), CD8+ T cells (anti-CD8 clone 2.43, ATCC; Manassas, VA, USA), NK cells (anti-asialo-GM; Wako Chemical, Dallas, TX, USA), or with antibodies (100 $\mu\text{g}/\text{mouse}$; [27]) against $\text{IFN}\gamma$ (anti- $\text{IFN}\gamma$ clone R4-6A2, ATCC; Manassas, VA, USA), CXCL9 or CXL10 (gifts from Dr Valbuena; [28]) after 5 and 15 days.

Immunohistochemistry and ELISA

Mice were sacrificed with an overdose of IP-administered Avertin anesthetic (0.017 ml/g body weight; Aldrich, Milwaukee, WI, USA) followed by cervical dislocation. For histological analyses, tissues were fixed in paraformaldehyde, embedded in paraffin, and stained with hematoxylin

and eosin (H&E). For immunohistochemistry, paraffin sections were stained using an antibody specific for the Ly-6G antigen (BD, Franklin Lakes, NJ, USA). For ELISA, tumors from mice ($n = 3$ per group) were extracted using a solution consisting of PBS containing 1% NP-40 and protease inhibitors, normalized for total protein content, and analyzed by ELISA for cytokines CXCL9, CXCL10 (R&D Systems, Minneapolis, MN, USA) and $\text{IFN}\gamma$ (eBioscience, San Diego, CA, USA). Results represent amounts of cytokines per gram of protein (mean \pm SD).

Statistics

Statistical significance of differential findings between controls and treatment groups was determined by two-tailed Student's t -test (Figs. 2c, 3) or by ANOVA. The significance of metastases scores was determined by Mann–Whitney U -test. Findings were regarded as significant if P -values were less than 0.05.

Results and discussion

CCL21 protein expression by attenuated *S. typhimurium*

To engineer the attenuated *S. typhimurium* strain $\text{purI}^-/\text{msbB}^-$ to express CCL21, we used the plasmid pGEN206, in which a murine CCL21 cDNA was expressed under the control of an *ompC* promoter. The engineered bacteria were grown in culture, recovered by centrifugation, and the resulting cell-containing pellet and cleared culture supernatant were tested for CCL21 protein by immunoblotting. The results demonstrated the presence of CCL21 in both cells and supernatant (Fig. 1). The production of CCL21 in vitro in bacterial cultures is >50 pg/ μg . Control *S. typhimurium* (transformed with empty plasmid) and their culture supernatants did not contain CCL21 protein (Fig. 1).

S. typhimurium-expressing CCL21 inhibit primary and metastatic tumor growth

Recently, we showed that the attenuated strain of *Salmonella* used here accumulates in subcutaneous tumors in vivo in mice, using luciferase-expressing bacteria in conjunction with bioluminescence imaging [12]. We tested the anti-tumor activity of CCL21-expressing bacteria in vivo by performing experiments in which mice were challenged SC with CT-26 colon (Fig. 2a) or D2F2 breast carcinoma cells (Fig. 2b) to form SC tumors, or IV with CT-26 cells to form experimental lung metastases (Fig. 2c). Following tumor challenge, mice were treated IV with PBS, empty *S. typhimurium* or *S. typhimurium* bearing control or CCL21 vectors. In the case of SC tumors, mice received treatment

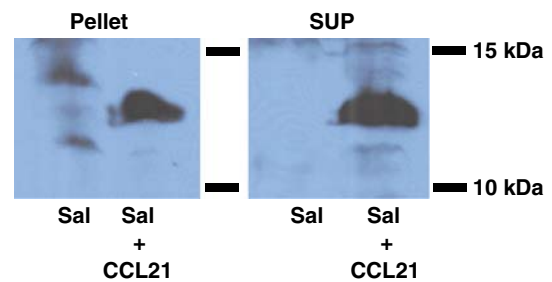


Fig. 1 CCL21 expression by *S. typhimurium*. Attenuated lipid A-negative *S. typhimurium* ($\text{pur}^-/\text{msb}^-$) were generated as previously described [6]. A cDNA encoding murine CCL21 (InvivoGen, San Diego, CA, USA) was inserted into pGEN206. This vector is a slightly modified form of pGEN222 [25] in which the *parA* locus has been exchanged with a *parM* and a *parR* locus. In addition, a *repA* locus has been introduced between the origin of replication and the multiple cloning sites. Expression of the 12 kD CCL21 protein was demonstrated by immunoblot analysis of cell lysates (“pellet”) or culture supernatants (“SUP”) from 10^9 CFU of *S. typhimurium* containing either empty vector (Sal) or CCL21-bearing plasmid-transformed *S. typhimurium* (Sal + CCL21). Samples were normalized for total protein content, resuspended in Laemmli sample buffer and subjected to SDS-PAGE followed by immunoblotting with a polyclonal goat anti-CCL21 antibody (Abcam, Cambridge, MA, USA). Results are representative of three independent experiments

at a consistent time point after tumors reached a clearly visible size. Marked reduction of subcutaneous tumor growth was observed in mice receiving *S. typhimurium* expressing CCL21 compared with control groups (Fig. 2a, b). Recovering bacteria from excised tumors and various organs of the mice sacrificed at the termination of the study confirmed accumulation of *Salmonella* predominantly in tumor tissue: (938 ± 259 million CFU/g tumor tissue compared to 0.10 ± 0.26 million CFU/g liver tissue, 0.16 ± 0.38 million CFU/g spleen tissue, 0.90 ± 1.34 million CFU/g kidney tissue and 0 ± 0 million CFU/g lung tissue) (Supplemental data). In addition to suppressing growth of subcutaneous tumors, a marked reduction in lung metastases was observed after treatment with CCL21-expressing bacteria compared with control mice. This suppression of metastatic growth was assessed by determining the weight of the lungs of sacrificed animals (Fig. 2c) and determining metastases scores for excised lungs (Fig. 2d).

CCL-21-expressing bacteria induce inflammatory cell infiltration into tumors and elevate intratumoral levels of cytokines and chemokines

We examined the intratumoral levels of $\text{INF}\gamma$, CXCL9, and CXCL10, chemokines and cytokines known to be induced by CCL21 [23, 24]. In mice treated with CCL21-expressing *Salmonella*, intratumoral levels of $\text{INF}\gamma$, CXCL9, and CXCL10 were significantly elevated, compared to either PBS- or control *Salmonella*-treated mice (Fig. 3a). Consistent with elevated levels of cytokines and chemokines,

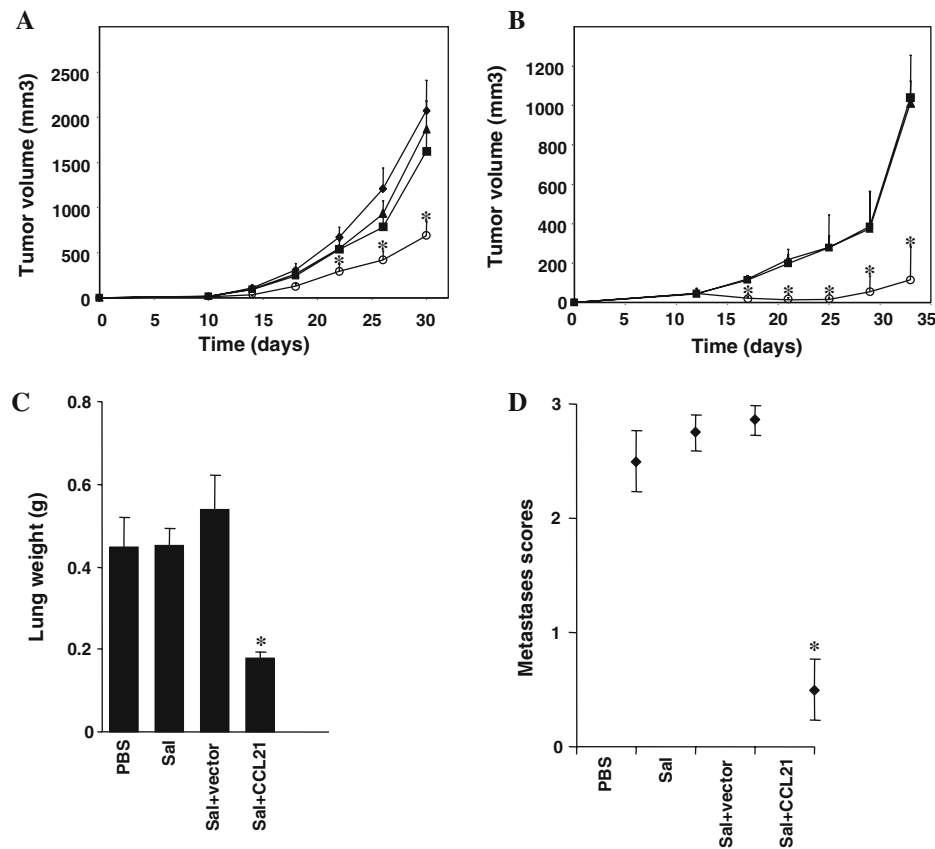


Fig. 2 Effect of CCL21-targeting *S. typhimurium* on tumor growth. Female mice were purchased at about 6–8 weeks of age from The Jackson Laboratory (Bar Harbor, ME, USA). All animal experiments were performed according to the National Institutes of Health *Guide for Care and Use of Experimental Animals* and approved by the Animal Care Committee of the Burnham Institute for Medical Research (#AUF 04-152). **a** BALB/c mice ($n=8$) were challenged SC with 1.25×10^5 CT-26 colon carcinoma cells (from ATCC; Manassas, VA, USA) and treated IV with PBS (dark filled diamond), or 5×10^6 CFU of *S. typhimurium* (dark filled square), *S. typhimurium* with empty vector (dark filled triangle), or *S. typhimurium* with CCL21 vector (open circle) after 14 days. Bar graphs indicate average tumor volumes measured in two dimensions by external calipers and calculated as length/2 \times width² (mean \pm SE). **b** Mice were challenged SC with 2.5×10^5

D2F2 breast carcinoma cells and treated IV with 5×10^6 CFU of *S. typhimurium* (dark filled square), *S. typhimurium* containing empty vector (dark filled triangle), or *S. typhimurium* bearing CCL21 vector (open circle) after 9, 14 and 19 days. Bar graphs indicate average tumor volumes of five mice (mean \pm SD). **c, d** Mice ($n=8$) were injected IV with 5×10^4 CT26 colon carcinoma cells and then treated IV with PBS, or 5×10^6 CFU of *S. typhimurium* (Sal), *S. typhimurium* containing empty vector (Sal + vector), or *S. typhimurium* bearing the CCL21 plasmid (Sal + CCL21) after 6, 13 and 20 days. After 29 days, lungs were weighed (**c** mean \pm SD; normal lung weight \sim 0.15 g) and examined for metastases (**d**), assessing the percentage of lung surface covered by metastases as follows: 0 = 0%, 1 = <20%, 2 = 20–50%, 3 = >50%. Asterisks indicate $P < 0.05$. Results are representative of four independent experiments

histological analysis of tumors excised from treated mice showed abundant infiltrates of immune (mononuclear) and especially inflammatory (polymorphonuclear) cells in mice treated with CCL21-expressing bacteria, with far less cell infiltration observed in mice receiving PBS or control *Salmonella* (Supplementary information). Abundant neutrophils (confirmed by Ly-6G immunostaining) and evidence of tumor necrosis suggested that some of the inflammatory infiltrates may be secondary to tumor lysis.

Anti-tumor activity of CCL21-expressing bacteria requires CD4 and CD8 cells

We challenged C57/B16 mice SC with B16 melanoma cells and various groups of mice were then treated with comple-

ment-fixing antibodies to deplete in vivo CD4⁺ T cells, CD8⁺ T cells, or NK cells. Furthermore, IFN γ , CXCL9, and CXCL10 were neutralized in mice with antibodies (Fig. 3b). As demonstrated previously for intratumoral injected CCL21 [17], CD4⁺ T cells and CD8⁺ T cells (but not NK cells) were indispensable for significant inhibition of tumor growth by CCL21-expressing bacteria (Fig. 3b). Thus, both CD4⁺ and CD8⁺ T-cells appear to contribute to the observed antitumor activity.

Attempts to individually neutralize CCL21-induced cytokine/chemokines with antibodies showed that CXCL9 and CXCL10 significantly contributed to inhibition of tumor growth (Fig. 3b). In contrast, antibody neutralization of IFN γ did not significantly impair anti-tumor activity.

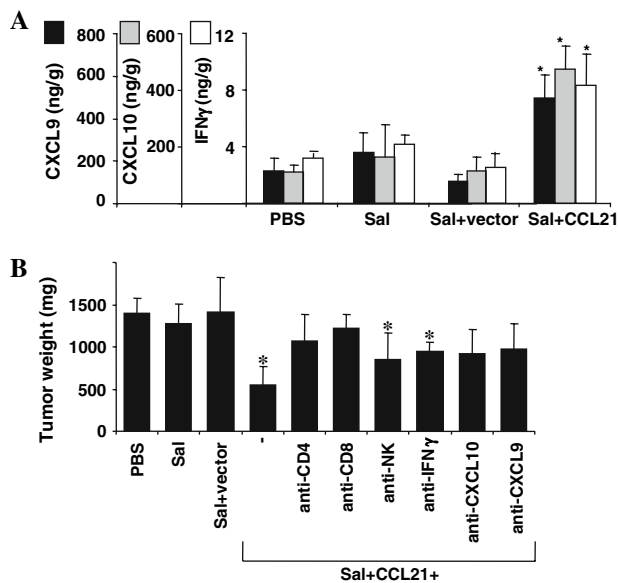


Fig. 3 Analysis of mechanism of anti-tumor activity of CCL21-expressing *S. typhimurium*. **a** Portion of tumors from sacrificed mice ($n = 3$) was extracted using a 1% NP-40 solution in PBS, and analyzed by ELISA (eBioscience, San Diego, CA, USA) for the CXCL9 (dark filled square) and CXCL10 (dark filled square) chemokines as well as IFN γ (open square). Data represent ng cytokine per gram of total protein (mean \pm SD). **b** C57/B16 mice ($n = 5$) were challenged SC with 1.25×10^5 B16 melanoma cells (from ATCC; Manassas, VA, USA). After 7 days, mice were injected IV with PBS, or 5×10^6 CFU of *S. typhimurium* (Sal), *S. typhimurium* containing empty vector (Sal + vector) or *S. typhimurium* bearing CCL21 vector (Sal + CCL21), including groups of mice that were treated IP with (500 μ g/mouse) antibodies against CD4+ T cells (Sal + CCL21 + anti-CD4; anti-CD4 clone GK1.5, ATCC; Manassas, VA, USA), CD8+ T cells (Sal + CCL21 + anti-CD8; anti-CD8 clone 2.43, ATCC; Manassas, VA, USA), NK cells (Sal + CCL21 + anti-NK; anti-asialo-GM, Wako Chemical, Dallas, TX), or (100 μ g/mouse) antibodies against IFN γ (Sal + CCL21 + anti-IFN γ ; ATCC; Manassas, VA, USA), CXCL10 (Sal + CCL21 + anti-CXCL10; gift from Dr Valbuena), or CXCL9 (Sal + CCL21 + anti-CXCL9; gift from Dr Valbuena) after 5 and 15 days. Depicted are the tumor weights (mean \pm SD) after 22 days. Asterisks indicate $P < 0.05$ compared to mice treated with PBS, *Salmonella*, or *Salmonella* + vector. Results are representative of three independent experiments

Though demonstrating potent anti-tumor activity, CCL21-expressing *S. typhimurium* did not cause greater systemic toxicity than the unmodified attenuated *S. typhimurium* strain (data not shown). Grossly, the mice did not show decreased activity or grooming abnormalities until several weeks after *S. typhimurium* injections, after tumor burden increased, suggesting that the tumor (rather than the therapy) contributed to these signs of illness. Histological analysis of multiple organs (data not shown) showed minimal changes that were comparable to the results previously reported for attenuated *S. typhimurium* control bacteria that did not express the CCL21 chemokine [7, 10].

Concluding remarks

We describe a novel bacteria-based strategy using an attenuated strain of *S. typhimurium* engineered to produce a chemokine in vivo. The concept was to exploit the intrinsic capability of these bacteria to accumulate specifically within tumors with the intent to produce functional human chemokine locally within the tumor lesions [11]. We chose CCL21 based on reported activities of this chemokine as a contributor to host responses to tumors [17–19]. Using the same mouse cancer models, we have shown previously [12] and here that the attenuated *S. typhimurium* strain localizes predominantly to tumors in mice, confirming several other studies that have documented tumor targeting by these facultative anaerobes [7, 10].

Our data demonstrate that IV delivery of CCL21-expressing attenuated *S. typhimurium* induces an inflammatory response within tumors and causes a substantial reduction of tumor burden in mice challenged with multidrug-resistant CT26 colon, D2F2 breast, and B16 melanoma cells [26]. In contrast, control *Salmonella* showed little or no anti-tumor activity, thus demonstrating the contribution of CCL21 to the therapeutic effect. The mechanisms responsible for the anti-tumor activity of CCL21-expressing *Salmonella* are probably multifactorial, with the predominant mechanism possibly varying with the tumor involved, anatomical location (e.g., subcutaneous vs. pulmonary), mouse strain, and other variables. In the B16 melanoma model, we showed that the anti-tumor activity of CCL21-expressing bacteria was significantly impaired in mice depleted of CD4⁺ or CD8⁺ T cells, which is in accordance with previous reports involving the intratumoral application of CCL21 [17–19]. In contrast, antibodies targeting NK cells did not significantly impact anti-tumor activity. However, it should be recognized that in vivo antibody immunodepletion is never completely effective for eliminating cell populations, and thus a role for NK cells cannot be entirely excluded. In addition, experiments using neutralizing antibodies targeting cytokines or chemokines known to be induced by CCL21 provided evidence that CXCL9 and CXCL10 contribute significantly to the anti-tumor activity. While antibodies neutralizing INF γ did not significantly impair anti-tumor activity of CCL21-expressing *Salmonella*, we cannot entirely exclude a role for this cytokine because, individually, some cytokines may not be essential but in combination with others, they may make important contributors to the observed anti-tumor activity of CCL21-expressing *Salmonella*.

Histological examination demonstrated the induction of an inflammatory response within tumors following infection with CCL21-producing bacteria, with far less immune and inflammatory cell infiltration in the tumors of mice treated with control bacteria lacking CCL21. In contrast,

inflammation of non-cancerous organs was limited principally to spleen, suggesting preferential targeting of the host response to tumors, though we cannot exclude a broader systemic effect of these chemokine-producing bacteria. The infiltrates in tumors of CCL21-treated mice included mononuclear and polynuclear cells. CCL21 is known to be chemotactic for lymphocytes and dendritic cells (mononuclear cells) but not polymorphonuclear cells. However, chemokines induced by CCL21 (e.g., CXCL9, CXCL10) are chemotactic for neutrophils [29]. Also, it bears noting that evidence of tumor necrosis was found histologically in mice treated with CCL21-expressing *Salmonella*, suggesting that some of the neutrophil infiltrates could be secondary to tumor lysis. In this regard, CCL21 has been reported to have a direct anti-angiogenic effect that could contribute to tumor hypoxia and cause subsequent necrosis. The anti-tumor activity of CCL21-expressing *Salmonella* therefore may be multifactorial, providing a distinct advantage over more target-focused therapies for which tumor resistance often emerges, thus making the pleiotropic actions of CCL21-expressing *Salmonella* a distinct advantage. Conversely, it should be recognized that the pleiotropic actions may include adverse mechanisms, such as induction of regulatory T cells and myeloid-suppressor cells, which could reduce efficacy of this immunotherapy strategy [30].

Taken together, our preclinical findings demonstrate the feasibility of a novel microbial-based approach for cancer immunotherapy involving engineering *S. typhimurium* to express the chemokine CCL21. These findings extend our recent analysis of bacteria armed to express the cytokine LIGHT, a TNF-family member [12], and thus demonstrate the potential of using chemokines or cytokines for this application. Future studies involving other types of immunomodulatory proteins should help to reveal the full repertoire of molecules that might be appropriate to consider for cancer immunotherapies using engineered non-virulent *S. typhimurium*. Ultimately, however, human clinical trials will be needed to test the utility of such approaches and the therapeutic index of specific types of molecules, given the important genomic differences between rodent species and humans with respect to their repertoires of genes encoding cytokines and their receptors [31].

Regardless of the relative merits of various chemokines or cytokines for engineering non-virulent *S. typhimurium*, a bacteria-based strategy for de novo production of immunomodulatory proteins may have several advantages over other approaches. For example, virus-based cytokine gene delivery is difficult to terminate, given the general lack of antiviral drugs capable of halting viral infection should unacceptable toxicity be encountered. Most of the viruses under clinical investigation also lack biological features that may preferentially target them to tumors. Furthermore, recombinant cytokines are typically expensive to produce,

lack tumor-targeting features, and often cause severe systemic inflammatory reactions that would limit clinical use. In contrast, genetically engineered bacteria are inexpensively produced, can selectively target tumors, and are readily eradicated by antibiotic therapy in the event of unacceptable toxicity. Altogether, these advantages, along with the results from this and similar studies, indicate that further preclinical evaluations are warranted of chemokine-producing bacteria as a novel microbial approach to cancer immunotherapy.

Acknowledgments We thank M. Hanai and T. Siegfried for manuscript preparation, M. Cuddy for manuscript review, Dr W. Z. Wei for D2F2 cells, and Dr G. Valbuena for CXCL9 and CXCL10 antibodies. We also acknowledge the generous support of the Austrian Academy of Sciences, APART Fellowship Program (M. Loeffler) and the NIH (CA-69381) (J. C. Reed).

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