Loss of secreted gelsolin enhances response to anticancer therapies

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ABSTRACT

Type 1 conventional dendritic cells (cDC1) play a critical role in priming anticancer cytotoxic CD8+ T cells. DNGR-1 (a.k.a. CLEC9A) is a cDC1 receptor that binds to F-actin exposed on necrotic cancer and normal cells. DNGR-1 signaling enhances cross-presentation of dead-cell associated antigens, including tumor antigens. We have recently shown that secreted gelsolin (sGSN), a plasma protein, competes with DNGR-1 for binding to dead-cell-exposed F-actin and dampens anticancer immunity. Here, we investigated the effects of loss of sGSN on various anticancer therapies that are thought to induce cell death and provoke an immune response to cancer. We compared WT (wildtype) with Rag1−/−, Batf3−/−, Clec9a−/− or sGSN−/− mice implanted with transplantable tumor cell lines, including MCA-205 fibrosarcoma, 5555 BrafV600E melanoma and B16-F10 LifeAct (LA)-ovalbumin (OVA)-mCherry melanoma. Tumor-bearing mice were treated with (1) doxorubicin (intratumoral) chemotherapy for MCA-205, (2) BRAF-inhibitor PLX4720 (oral gavage) targeted therapy for 5555 BrafV600E, and (3) x-ray radiotherapy for B16-LA-OVA-mCherry. We confirmed that efficient tumor control following each therapy requires an immunocompetent host as efficacy was markedly reduced in Rag1−/− compared with WT mice. Notably, across all the therapeutic modalities, loss of sGSN significantly enhanced tumor control compared with treated WT controls. This was an on-target effect as mice deficient in both sGSN and DNGR-1 behaved no differently from WT mice following therapy. In sum, we find that mice deficient in sGSN display enhanced DNGR-1-dependent responsiveness to chemotherapies, targeted therapy and radiotherapy. Our findings are consistent with the notion some cancer therapies induce immunogenic cell death. What this study adds

Type 1 conventional dendritic cells (cDC1) play a non-redundant role in anticancer immunity.

DNGR-1 is a dedicated cDC1 receptor that facilitates cross-presentation of dead-cell associated antigens.

Secreted gelsolin (sGSN) is a physiological barrier to DNGR-1-dependent cross-presentation and dampens anticancer immunity.

Some cancer therapies may induce an antitumor immune response by triggering immunogenic cell death.

WHAT THIS STUDY ADDS

Loss of sGSN enhances therapeutic response to several immunogenic anticancer therapies in preclinical models including chemotherapy, targeted therapy and radiotherapy.

HOW THIS STUDY MIGHT AFFECT RESEARCH, PRACTICE OR POLICY

Therapeutic inhibition of sGSN function, to unleash DNGR-1-dependent cross-presentation, could be explored as a component of more effective treatment regimens.

BACKGROUND

Type 1 conventional dendritic cells (cDC1) perform non-redundant functions in anticancer immunity.1–3 Mice deficient in cDC1s (eg, Batf3−/−) display lower spontaneous rejection of immunogenic tumors and decreased responses to T cell based immunotherapies.4 In humans, a cDC1 gene signature positively correlates with improved overall survival (OS) in patients with cancer and with responses to checkpoint blockade immunotherapy.1,2 A key role of cDC1 is to prime tumor antigen-specific CD8+ T cells. It is thought that this partly involves internalization by cDC1 of dead cancer cell remnants and subsequent MHC class I (cross-)presentation to CD8+ T cells of antigens derived from the debris.

DNGR-1 (a.k.a. CLEC9A) is a C-type lectin receptor ubiquitously expressed on cDC1 that binds to F-actin exposed on dead cell corpses and signals to enhance cross-presentation of dead-cell associated antigens.3,4 We recently found that secreted gelsolin (sGSN), an extracellular protein found in the plasma of all vertebrates,7,8 forms a natural barrier to DNGR-1-mediated cross-presentation.9 sGSN promotes cancer immune escape by competing with DNGR-1 for binding to...
extracellular F-actin and inhibiting DNGR-1-dependent tumor control.9 In humans, high expression of CLEC9A (encoding DNGR-1) in the tumor microenvironment is associated with greater survival and correlates with gene signatures of immune-mediated cancer control.10 In mice, DNGR-1 can contribute to CD8+ T cell-dependent control of transplantable tumors in animals deficient in sGSN.9

For decades, chemotherapy, radiotherapy and, more recently, targeted therapy, have been the mainstay of treatment for most patients with cancer. There is an increasing evidence that these therapies work not only by directly killing cancer cells but also by inducing or boosting antitumor immune responses.11–13 We, therefore, hypothesized that therapy-induced cell death would increase the antigenic visibility of tumors, augmenting DNGR-1 triggering by F-actin in sGsn−/− mice and resulting in improved anticancer immunity. Here, we show that mice deficient in sGsn display enhanced DNGR-1-dependent responsiveness to chemotherapy, radiotherapy and targeted therapy of tumors, highlighting sGSN as an attractive target for combinatorial cancer treatments that engage immunity.

**METHODS**

**Mice**

Batf3−/−, Clec9aΔβ/Δβ, sGsn−/−, sGsn−/− Clec9aΔβ/Δβ, Rag1−/− and WT (wildtype) mice on a C57BL/6 background were bred and maintained in specific-pathogen free conditions in the Biological Research Facility (BRF) at The Francis Crick Institute. In all experiments, male and female mice were used. Experiments were initiated when mice were between 6 and 14 weeks. In all experiments comparing groups of mice of different genotypes, they were co-housed with sex-matched and age-matched WT controls for at least 3 weeks to eliminate any microbiota-dependent effects. Alternatively, in some experiments where WT littermate controls were used, this is indicated in the figure legends. All animal experiments were performed on prospective approval of a study plan by the BRF at The Francis Crick Institute and adhered to the UK Animals (Scientific Procedures).7

**Transplantable tumors**

Transplantable tumor cell lines used include MCA-205 fibrosarcoma (kind gift from George Kassiotis), 5555 BrafV600E melanoma,14 and B16-F10 LifeAct (LA)-ovalbumin (OVA)-mCherry.9 All cell culture procedures were performed in a sterile condition in a laminar flow hood. Cultured syngeneic tumor cells were dissociated with trypsin (0.25%) (Gibco) and washed for three times in PBS (Gibco). The final cell pellet was resuspended in PBS for the required concentration of cells accordingly and injected subcutaneously (s.c.) in a volume of 100 µl in the shaved right flank of each mouse. Mice were monitored for tumor growth at least two to three times per week. The values of the longest (L) and perpendicular shortest (W) tumor diameters were measured using digital Vernier calipers (Mitutoyo), and tumor volume calculated using the formula: 0.5 x (L x W)^2.15 expressed in the standard metric units mm³. Mice-bearing tumors were monitored until the predetermined humane endpoint (eg, mean tumor diameter reaching ≥15 mm, weight loss ≥15%, or tumor ulceration) or for at least 30 days following tumor transplantation.

**Chemotherapy model**

For in vivo chemotherapy, mice were inoculated s.c. with 0.5×10⁶ cells/100 µl MCA-205 fibrosarcoma. Doxorubicin (Sigma-Aldrich) was administered at a dose of 2.5 mg/kg in a final volume of 50 µl sterile PBS intratumorally. Treatment was administered when tumors became palpable, usually on day 6 or 7, when the mean tumor diameter reached 6–8 mm.

**Radiotherapy model**

For radiotherapy, mice were inoculated s.c. with 0.2×10⁶ cells/100 µl of B16-F10 LA-OVA-mCherry melanoma. On day 7, mice received a single fraction of 10 Gy X-ray irradiation in an X-ray irradiator cabinet (Xstrahl RS320 Research System). The machine was calibrated immediately prior to every use, and radiation was delivered at a consistent dose rate of 1.5–1.6 Gy/min (200kV, 16mA) at a field source distance of 20 cm. Radiation was targeted to the right flank tumor, precisely positioned using laser beams, through a 15 mm aperture at the center of an overlying customized lead shield (2 mm thickness). The shielded portion of the body received a maximum of 0.3 Gy radiation whereas 10 Gy was delivered to target tumor area. Mice received reversible general anesthesia throughout the duration of the procedure using a ‘sleep mix’ of fentanyl 0.05 mg/kg, midazolam 5 mg/kg and medetomidine 0.5 mg/kg, and reversed with a ‘wake mix’ of naloxone 1.2 mg/kg, flumazenil 0.5 mg/kg and atipam 2.5 mg/kg, administered intraperitoneally and dosed according to weight per mouse. Mice were kept in a warming chamber (35°C–37°C) (Datisand Thermacage) during anesthesia, both in preparation for administration of radiation and also for recovery. Mice receiving sham-irradiation were put through reversible anesthesia and recovery in the warming chamber for the same duration.

**Targeted therapy model**

Mice received 0.2×10⁶ cells/100 µl s.c. inoculation of 5555 BrafV600E melanoma. When tumors were palpable, generally at day 6 after tumor cell inoculation, mice received BrafV600E kinase inhibitor PLX4720 (MedKoo) suspension at a dose of 45 mg/kg in total volume of 200–250 µl by oral gavage (o.g.) daily for a total of 2 weeks, with 5% DMSO as vehicle control.

**Statistics**

Statistical analyses were performed using GraphPad Prism V.9 for Mac OS (V.9.3.0, updated 2021; San Diego, California, USA). Data are shown as mean±SE of the mean (SEM) unless otherwise stated. In general, for all tumor growth profiles, two-way analysis of variance was used to compare the means of two or more groups obtained at
every single timepoint, and this was adjusted for multiple comparisons with post-hoc Bonferroni correction. The statistical differences annotated on the plots represent the comparison at the final timepoints within the respective follow-up period of each experiment. In all instances, p values are two tailed and \( p \leq 0.05 \) is considered the threshold for statistical significance. Where indicated in figures, the following scheme is used to represent the levels of statistical significance: \(* p \leq 0.05, ** p < 0.01, *** p < 0.001, **** p < 0.0001\); ns, not significant.

RESULTS
Loss of sGSN enhances response to immunogenic chemotherapy
First, we confirmed that chemotherapy with doxorubicin in the transplantable MCA-205 primary fibrosarcoma requires an immunocompetent host for full effectiveness. Consistent with that notion, intratumoral doxorubicin administration reduced tumor growth in WT mice but therapeutic benefit was reduced in \( \text{Rag1}^{-/-} \) mice which lack T and B cells (figures 1A,B). The efficacy of doxorubicin was similar in DNGR-1-deficient and WT mice, suggesting that DNGR-1 is dispensable in a sGSN-sufficient setting (online supplemental figure S1a). However, when tested in mice lacking sGSN, we observed that the administration of doxorubicin resulted in significantly greater tumor control compared with treated WT mice (figure 1A,C).

The efficacy of immunogenic radiotherapy is improved in the absence of sGSN
We next tested the effect of sGSN loss in tumor responses to radiotherapy (figure 1D). We used the poorly immunogenic B16-F10 melanoma cell line engineered to express the model antigen OVA fused to mCherry and a 17-amino acid LA F-actin binding peptide (B16-F10 LA-OVA-mCherry).1 We found that effective and sustained control of B16-F10 LA-OVA-mCherry tumors following a single fraction of X-ray irradiation was lost in \( \text{Batf3}^{-/-} \) mice lacking cDC1 or \( \text{Rag}^{1/-} \) mice lacking T and B cells (figure 1E). As in the doxorubicin chemotherapy model, tumor control following radiotherapy was unaffected by deficiency in DNGR-1 (online supplemental figure S1b). However, it was significantly enhanced in sGSN-deficient hosts (figure 1F).

Targeted therapy induces immunogenic cell death and its efficacy is further enhanced in absence of sGSN
We also sought to investigate whether sGSN deficiency might benefit the response to targeted therapy with BRAF inhibitor in a mouse mutant-Braf melanoma model. Tumor-bearing mice implanted with 5555 Braf\(^{V600E}\) primary melanoma tumor cells were treated or not with PLX4720 (figure 1H). We found that the therapeutic effect of the drug was decreased in mice lacking T and B cells (figure 1H). Notably, 5555 Braf\(^{V600E}\) tumors were controlled to a greater extent in \( sGsn^{-/-} \) compared with WT mice even in the absence of treatment with PLX4720 (figure 1I). Importantly, however, tumor control by PLX4720 was further enhanced by sGSN-deficiency (figure 1J and online supplemental figure S1c), confirming the importance of sGSN as a barrier to dead cell-induced immunity.

The enhanced therapeutic response in \( sGsn^{-/-} \) mice is DNGR-1-dependent
The data so far suggest that maximal therapeutic efficacy of some forms of chemotherapy, radiotherapy and targeted therapy in vivo requires an intact adaptive immune system and is further augmented by loss of sGSN. In order to validate that enhanced tumor control in \( sGsn^{-/-} \) mice is DNGR-1-dependent, we tested some of the therapy models in mice deficient in both sGSN and DNGR-1. For both chemotherapy with intratumoral doxorubicin in MCA-205 and radiotherapy in B16-F10 LA-OVA-mCherry, mice deficient in both sGSN and DNGR-1 exhibited similar rate of tumor control to the respective treated WT controls, in contrast to the enhanced therapeutic responses observed in the treated \( sGsn^{-/-} \) subgroups (figure 2A,B). Thus, sGSN impairs DNGR-1-dependent anticancer immunity-induced by therapy.

DISCUSSION
While much of the therapeutic effect of conventional anticancer therapies results from direct cytotoxicity to cancer cells, it is becoming clear that it can also include immune-dependent mechanisms. Specifically, it has been proposed that some cancer therapeutics can cause immunogenic cell death (ICD), allowing for sampling of tumor antigens by DCs that then prime or boost an antitumor T cell response.1,11 DNGR-1 is a receptor that marks the cDC1 subtype of DCs and plays a key role in coupling cDC1 to antitumor CD8\(^+\) T cell immunity.3,6,11 While DNGR-1 activity is often masked by circulating sGSN such that the role of the receptor can be revealed only on loss of the latter protein.9 Here, we show that the immune-dependent efficacy of seemingly disparate therapies in a preclinical mouse model is increased by loss of sGSN. Notably, although DNGR-1 deficiency per se does not impact therapy efficacy, it curtails the therapeutic advantage conferred by loss of sGSN. Thus, these data implicate the sGSN-DNGR-1 axis in decoding ICD induced by cancer treatment.

The contribution of immunity to the effects of conventional chemotherapy and radiotherapy is beginning to be appreciated in patients, but mechanistic insights have been primarily obtained in animal models. In mice, several chemotherapeutic agents, including the anthracycline doxorubicin, result in ICD and lead to antitumor responses that depend on DCs and CD8\(^+\) T cells.11,12 Similarly, both DCs and CD8\(^+\) T cells have been previously demonstrated to contribute to tumor control following radiotherapy.13,15 Our study provides further evidence for the immunogenicity of some anticancer therapies.11,12
showing that their potency is reduced in mice lacking T (and B) cells. We show that this immune benefit can be further enhanced in a DNGr-1 dependent manner on loss of sGSN, concordant with our previous work showing an increase in tumor-specific T cell responses in GSN-deficient animals.9 Thus, DNGr-1-dependent cross-presentation of dead cell-associated antigens can increase anticancer immunity and could be exploited clinically.

In addition to conventional chemotherapy and radiotherapy, a growing repertoire of therapies target driver mutations in cancer. An example is BRAF inhibitors, which have been used to treat BRAF-mutant melanoma.20 However, clinical responses are variable and often curtailed by development of drug resistance. Improving clinical response by co-opting immunity may help tackle resistance to targeted therapy and potentially improve...
Figure 2  Enhanced therapeutic response in the loss of secreted gelsolin settings is dependent on DNGR-1. (A) Tumor growth profile in mice bearing MCA-205 fibrosarcoma receiving chemotherapy. Doxorubicin was administered to cohoused WT (n=10), sGsn−/−(n=10) and sGsn−/− Clec9agfp/gfp (n=10) mice, and compared with cohoused WT (n=10), sGsn−/− (n=3) and sGsn−/− Clec9agfp/gfp (n=3) mice receiving vehicle control. (B) Growth profile of B16-F10 LA-OVA-mCherry melanoma in co-housed WT (X-ray, n=14; sham, n=10), sGsn−/− (X-ray, n=11; sham, n=10) and sGsn−/− Clec9agfp/gfp (X-ray, n=12; sham, n=11) mice. Data are plotted as tumor volume (mm³) ± SEM, and mean tumor volumes were compared using two-way ANOVA with post hoc Bonferroni correction. Where indicated, **p<0.01, ***p<0.001, ****p<0.0001; NS, not significant. Error bars are depicted in all plots; when not visible, errors are small. Data are representative of one experiment respectively. ANOVA, analysis of variance; LA, LifeAct; OVA, ovalbumin; WT, wild type.

outcomes. Indeed, NK cells are essential for effective melanoma control in mice treated with the BRAF inhibitor PLX4720 and, in human paired-biopsy studies, treatment with BRAF-inhibitor increases tumor infiltration by cytotoxic CD8+ T cells. Here, we show that melanoma control by targeted therapy with PLX4720 requires an immunocompetent host and reveal that efficacy can be enhanced in the absence of sGSN. Thus, sGSN blocking could be a potentially useful adjunct to targeted therapy.

Collectively, these data complement and extend our previous observations that sGSN-deficient mice display enhanced responses to immune checkpoint blockade cancer therapy, and that DNGR-1 expression within tumors is associated with greater OS in several human cancers. Further highlighting the sGSN-DNGR-1 axis as a novel checkpoint in anticancer immunity, we previously reported that in a small subset of patients with stomach adenocarcinoma, high expression of CLEC9A and low expression of sGSN in tumors conferred the best survival outcomes compared with the other stratified subgroups in retrospective analysis. These findings will need to be prospectively validated to identify patients who may benefit most from inhibition of sGSN function to unleash DNGR-1-dependent cross-presentation as a component of more effective treatment regimens.

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Contributors KHJ conducted all the experiments and analyzed data with assistance from EG and OS, AW and ES helped with the set up of the radiotherapy experiments. NCR managed all mouse colonies. KHJ and CReS designed the study, interpreted the data, and wrote the manuscript. JS and CReS supervised the project. All authors reviewed and edited the manuscript.

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Competing interests KHJ, EG and CReS are named as contributors/inventors on a patent application on the use of sGSN for immunotherapies. CReS holds stock options and/or is a paid consultant for Adendra Therapeutics, Bicara Therapeutics, Montis Biosciences, Bicycle Therapeutics and Sosei Heptares. CReS holds a visiting professorship at Imperial College London and honorary professorships at University College London and King’s College London.

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REFERENCES
Supplementary Figure S1

a

MCA-205

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B16 LA-OVA-mCherry melanoma

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C

5555 Braf<sup>V600E</sup>

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SUPPLEMENTARY FIGURE CAPTION

Supplementary Figure S1 | DNGR-1 is dispensable in sGSN-sufficient mice for tumour control following immunogenic chemotherapy, radiotherapy and targeted therapy. (a) Tumour growth profile in mice bearing MCA-205 fibrosarcoma receiving chemotherapy. Doxorubicin was administered to cohoused WT (n = 8) and Clec9a^gfp/gfp (n = 7) mice, and compared to cohoused WT (n = 6) and Clec9a^gfp/gfp (n = 7) mice receiving vehicle control. (b) Growth profile of B16-F10 LA-OVA-mCherry melanoma in co-housed WT (X-ray, n = 12; sham, n = 10) and Clec9a^gfp/gfp (X-ray, n = 11; sham, n = 9) mice. (c) Growth profile of 5555 Braf^V600E melanoma in Clec9a^gfp/gfp mice (PLX4720, n = 8; control, n = 6), compared to respective cohorts of WT mice (PLX4720, n = 7; control, n = 6) previously shown in Figure 1h (experiment performed at the same time). Data are plotted as tumour volume (mm^3) ± s.e.m., and mean tumour volumes were compared using two-way ANOVA with post-hoc Bonferroni correction. Where indicated, *P ≤ 0.05, **P < 0.01, ***P < 0.001, ****P < 0.0001; ns, not significant. Error bars are depicted in all plots, even when not visible. Data are from one experiment.