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Prerequisites for the Antitumor Vaccine-Like Effect of Chemotherapy and Radiotherapy

Dalil Hannani, PhD,^{* †§} Antonella Sistigu, PhD,^{* †§} Il Oliver Kepp, PhD,^{*p§} Lorenzo Galluzzi, PhD,^{*p§} Guido Kroemer, MD, PhD,^{*p¶Lp} and Laurence Zitvogel, MD, PhD^{*p§}**

From the ^{*}Institut Gustave Roussy, Villejuif, France; [†]INSERM, U1015, Villejuif, France; [‡]INSERM, U848, Villejuif, France; [§]Université Paris Sud-XI, Villejuif, France; ^{||}Istituto Superiore di Sanita`, Rome, Italy; [¶]Metabolomics Platform, Institut Gustave Roussy, Villejuif, France; ^LCentre de Recherche des Cordeliers, Paris, France; ^{**}Pôle de Biologie, Hôpital Européen Georges Pompidou, Paris, France; ^{††}Université Paris Descartes, Paris, France.

Correspondance: Laurence Zitvogel, MD, PhD, INSERM, U1015, Institut Gustave Roussy, F-94805 Villejuif, Paris, France. E-mail: zitvogel@igr.fr.

Abstract: For a long time, anticancer therapies were believed to work (and hence convey a therapeutic benefit) either by killing cancer cells or by inducing a permanent arrest in their cell cycle (senescence). In both scenarios, the efficacy of anticancer regimens was thought to depend on cancer cell intrinsic features only. More recently, the importance of the tumor microenvironment (including stromal and immune cells) has been recognized, along with the development of therapies that function by modulating tumor cell extrinsic pathways. In particular, it has been shown that some chemotherapeutic and radiotherapeutic regimens trigger cancer cell death while stimulating an active immune response against the tumor. Such an immunogenic cell death relies on the coordinated emission of specific signals from dying cancer cells and their perception by the host immune system. The resulting tumor-specific immune response is critical for the eradication of tumor cells that may survive therapy. In this review, we discuss the molecular mechanisms that underlie the vaccine-like effects of some chemotherapeutic and radiotherapeutic regimens, with particular attention to the signaling pathways and genetic elements that constitute the prerequisites for immunogenic anticancer therapy.

Key Words: Calreticulin, HMGB1, NLRP3, TLR4, ATP, dendritic cells

IMMUNOGENIC TUMOR CELL DEATH

Although the armamentarium of anticancer therapies is being constantly ameliorated, the number of people succumbing to cancer has been predicted to drastically rise in the future (www.who.int/cancer). This trend basically reflects the facts that efficient therapies for some prominent neoplasms such as lung cancer (which now is the leading cause of cancer-related deaths worldwide) are still missing and that current anticancer regimens are often associated with the insurgence of resistance and therapeutic failure.¹ To counteract this dreadful tendency, further insights into the molecular mechanisms underlying the resistance of cancer cells to conventional therapy and into the signaling pathways that govern the host-tumor crosstalk are urgently awaited. This type of information will allow not only for the refinement of the current therapeutic arsenal, but also for a better stratification of cancer patients and the development of personalized anticancer therapies.

The current clinical approach to cancer most frequently involves surgery (whenever possible) alone or in association with a single-agent or combinatorial treatment based on chemotherapy or radiotherapy. Intriguingly, some cancers can be cured by conventional regimen (such as breast, colon, testicular, prostate, head and neck cancers, Hodgkin and follicular lymphoma, etc.), whereas others still remain a major medical challenge (such as lung and pancreatic cancers), suggesting that the intrinsic properties of the tumor and/or the specificity of the cytotoxic drugs matter. Minimal residual disease or incomplete eradication of tumor (stem) cells associated with the arousal of chemoresistant and/or radioresistant metastases¹ questioned the bases of our current reasoning. Thus, to improve the clinical outcome of anticancer therapies, it is of the utmost importance to understand how therapy-resistant tumor cells can be efficiently targeted and how therapeutic failure can be predicted.

For a long time, the field of clinical oncology was dominated by the notion that efficient anticancer therapies would work exclusively on tumor cells, either by inducing their apoptotic demise (cytotoxicity) and immunologically silent clearance or by permanently arresting their cell cycle progression (cytostasis). Moreover, several anticancer compounds were known to induce different degrees of immunosuppression, reinforcing the belief that the host immune system plays no role in the fight against transformed cells. Even the official guidelines formulated in 1975 by the National Institutes of Health recommended that the efficacy of novel anticancer strategies should be evaluated on human cancers xenotransplanted in immunodeficient mice.² More recently, it has been shown that (i) cancer cells engage in a strict crosstalk with their microenvironment (the tumor stroma, including fibroblasts and endothelial and immune cells such as macrophages) and that this interaction can be specifically targeted to induce tumor regression³; that (ii) highly efficient anticancer regimens can kill tumor cells through nonapoptotic cell death subroutines⁴; and that (iii) apoptosis can also occur in an immunogenic fashion, leading to the elicitation of an anticancer immune response.⁵

Such an immunogenic cell death (ICD) involves the transfer of tumor-derived antigens to immune cells that stimulate a tumor-specific immune response. This is critical for the eradication of residual cancer (stem) cells as it operates irrespective of their resistance to therapy.² Experiments in suitable animal as genetic interventions whereby BAX, BAK, and/or caspase 8 are removed or depleted, blocks CRT exposure and abolishes the tumor-vaccinating effect of cells undergoing ICD.^{6,15}

Third, approximately 5% to 10% of the endogenous CRT pool is exposed together with ERp57 at the surface of dying cells via SNAP and NSF attachment receptor (SNARE)-dependent exocytosis. This occurs well before plasma membrane permeabilization (which occurs as the final step of apoptosis) and also precedes the translocation of phosphatidylserine (PS) from the inner to the outer leaflet of the plasma membrane. Phosphatidylserine is the prototypic eat-me signal of apoptotic cells (although it has been implicated also in nonapoptotic cell death),^{20,21} and the kinetics of its exposure might affect the switch between the silent removal of dying cells by macrophages and the initiation of a cognate immune response by DCs.¹⁶ The receptor that is responsible for antigen uptake by DCs upon CRT binding remains to be determined. Possible candidates include the major CRT receptor CD91 as well as other CRT-interacting proteins such as scavenger receptor A, scavenger receptor expressed on endothelial cell I,²² CD40 ligand, tumor necrosis factor-related apoptosis-inducing ligand (tumor necrosis factor-related apoptosis-inducing ligand), or CD95/FAS ligand.²³ The CRT-driven uptake of tumor antigens by DCs is per se insufficient to elicit an anti-tumor immune response as internalized antigens must be processed and re-exposed for the cross-priming of CD4+ and CD8+ T lymphocytes. This implies that other signaling pathways are involved in ICD.

A systematic study of the response to CDAMPs of distinct Toll-like receptors (TLRs) on naive T cells revealed that TLR4 is both required and sufficient for efficient antigen presentation by DCs.²⁴ Among other proteins, TLR4 binds the non-histone chromatin-binding nuclear protein high-mobility group box 1 (HMGB1), leading to the activation of the downstream effector myeloid differentiation primary response 88 (MYD88).²⁴ This inhibits the fusion between lysosomes and antigen-containing phagosomes, thus facilitating antigen processing and presentation to T cells. High-mobility group box 1 also stimulates the neosynthesis of proIL-1A²⁵ but per se does not serve as a DC maturation signal.²⁴ For a long time, HMGB1 has been thought to exert a proinflammatory function exclusively during necrosis,²⁶ but recent evidence indicates that it also gets released during the late stages of apoptosis.²⁷ The re-lease of HMGB1 from tumor cells succumbing to ICD can be blocked by Z-VAD-fmk and hence depends on the activation of caspases.^{24,27} This process manifests with a dual kinetics whereby HMGB1 first translocates from the nucleus to the cytoplasm and then, following the breakdown of the plasma membrane, gets released into the extracellular space.⁸ Further insights into the molecular mechanisms that underlie HMGB1 release are missing. However, the addition of recombinant CRT or HMGB1 to dying cancer cells does not suffice to stimulate the presentation of tumor antigens by DCs,^{28,29} implying that additional signals are required for the immunogenicity of cell death.

The vaccine-like effect of ICD relies on the elicitation of an IFN- γ -polarized T-cell response, which in turn requires the function of the NLRP3 inflammasome, a multiprotein caspase-1-activating complex.³⁰ Caspase 1 activation is critical for activating an antitumor immune response as it catalyzes the proteolytic maturation of IL-1A.³¹ One of the most abundant factors that activate the NLRP3 inflammasome is ATP,³² and at least in DCs it does so by binding to the purinergic P2RX7 receptor on the cell surface.¹⁰ ATP also constitutes a CDAMP, as it gets released during the final steps of cell death, possibly via voltage-gated hemichannels of the pannexin 1 or connexin type.³³ Accordingly, the depletion of intracellular or extracellular ATP in cells succumbing to ICD abolishes the development of an IFN- γ -polarized response, and P2RX7-deficient mice fail to mount an immune response against syngenic cancer cells succumbing to ICD.¹⁰ Intriguingly, ATP also serves as a “find me” signal for the attraction of immune cells.³⁴ Altogether, these observations highlight the multifaceted and critical role of ATP for the vaccine-like effects of ICD inducers.

SPATIOTEMPORAL CODE

The spatially and temporally regulated emission of immuno-nogenic factors from dying tumor cells accounts for the re-cruitment and activation of immune cells to tumor bed and governs the immune response to cancer cells undergoing ICD (Fig. 1). Thus, the stress conditions that cancer cells confront during chemotherapy and radiotherapy determine whether the subsequent wave of cell death will elicit an antitumor immune response or rather will remain immunologically silent. Normally, cells attempt to cope with stress by arresting normal activities and by activating a series of cytoprotective mechanisms that aim at reestablishing homeostasis. For instance, stressed cells normally arrest protein synthesis, activate DNA repair pathways, and up-regulate factors for the handling of unfolded proteins as well as antioxidant defenses. This is accompanied by alterations of the surface proteome that, in the case of ICD, account for the recognition by immune cells and by the emission of soluble mediators with chemotactic and antichemotactic properties. This is crucial for the “selection” and differentiation/ maturation of engulfing cells, which in turn dictates the immuno-nogenic or tolerogenic outcome of cell death.

In this sense, the exposure of the DC-specific eat-me signal CRT (well before that of PS) paralleled by the disclosure of other, hitherto uncharacterized “don’t eat me” signals (such as CD47^{35Y37}) facilitates the recognition and uptake of dying tumor cells by DCs rather than by macrophages. Other authors have described additional molecular components exposed by dying cells that should be recognized for engulfment by specific DC subsets (such as CLEC9A/DNGR1 and HSP70/90) to elicit adaptive immune responses.^{29,38,39} However, additional stimuli released by dying tumor cells in the proximity of DCs are indispensable for the induction of a tumor-specific immune response. Thus, HMGB1 and ATP facilitate antigen processing/presentation and the release of IL-1A, which is necessary for IFN- γ -polarized T-cell responses. Based on these observations, the spatially restricted and temporally ordered appearance of CRT, HMGB1, and ATP might constitute a “key” that would precisely fit into a series of pattern recognition receptors expressed by DCs (the “lock”) for the conversion of nonimmunogenic into ICD and for the elicitation of an anti-cancer immune response.¹²

THE PERCEPTION OF IMMUNOGENIC CELL DEATH BY THE IMMUNE SYSTEM

Oncogenesis is a multistep mechanism that also involves an escape from immunosurveillance; that is, often cancer cells that sustain tumor growth are poorly immunogenic (immuno-noselection) and/or they actively inhibit immune functions (immunosubversion).⁴⁰ In this context, there are (at least) 2 different pathways whereby the immune system can be re-cruited against tumors: via direct immunomodulatory therapies that relieve immunosuppression or indirectly upon the induction of ICD.

By shaping T-cell responses, DCs are the first-line decision makers of the innate immune system, and their role in immunogenic chemotherapy has been deeply investigated. Experiments in transgenic mice that express the diphtheria toxin receptor under the control of a DC-specific promoter (allowing for in vivo DC depletion)⁹ revealed the essential role of DCs in the perception and decoding of “come and get me” signals emitted by dying tumor cells during ICD.²⁴ Similarly, the in vivo depletion of CD8⁺ T cells with specific antibodies has been instrumental to highlight the critical role of this lymphocyte subset for the vaccine-like effect of chemotherapy and radiotherapy in a large panel of murine tumor models, including CT26 colon cancer, EL4 thymomas, TS/A mammary carcinomas, MCA205 fibrosarcomas, Glasgow osteosarcoma osteosarcomas,^{11,24} and spontaneous methylcholanthrene-induced sarcomas.^{24,41} In line with these observations, CD8⁺ T cells have been shown to mediate potent anticancer immune effects in clinical settings, for instance, in colorectal tumors, where immune infiltration might serve as a prognostic factor.⁴²

Moreover, it has recently been shown that a precise orchestration of the T-cell response is required for immune effectors to eradicate tumors.⁴³ In this context, the IL-1 γ -dependent activation of IL-17-secreting F/C T cells had to precede the infiltration of tumors by Tc1 lymphocytes for the efficacy of immunogenic chemotherapy *in vivo*.⁴³ Thus, a finely regulated crosstalk between components of the innate (DCs) and cognate (F/C and CD8⁺ T cells) immune system is required for cell death to be perceived as immunogenic, for the elicitation of an anticancer immune response, and for complete tumor eradication leading to therapeutic success (Fig. 2). However, how resident macrophages and/or adverse inflammatory monocytes, which contribute to the proangiogenic and protumoral microenvironment and dominate the scenario before chemotherapy, become “transformed” and/or “overruled” by a subset of antigen presenting cells capable of eliciting a protective anticancer responses in the context of ICD remains to be established.

IMMUNOGENIC ANTICANCER CHEMOTHERAPY

The current definition of immunogenic chemotherapy is based on the ability of a limited array of antineoplastic drugs to elicit ICD rather than the stereotypical, immunologically silent or even tolerogenic apoptotic pathway. A plethora of preclinical^{44,51} and clinical⁵² studies revealed that DCs can take up apoptotic tumor cells and cross-present the internalized anti-gens on major histocompatibility complex class I molecules to CD8⁺ T cells, thus eliciting a productive immune response.

These studies suggest that the immunogenic outcome of cell death is influenced (among other factors) by the nature of the tumor cell as well as by the type of cell death inducer. In this context, a wide arsenal of stimuli including ER stressors (thapsigargin, tunicamycin, brefeldin), lysosome-targeting agents (bafilomycin A1), mitochondrion-targeting compounds (arsenite, betulinic acid, ceramide), proteasome inhibitors (MG132, lactacystin, ALLN), and DNA damaging molecules (Hoechst 33342, camptothecin, etoposide, mitomycin) is *per se* not immunogenic.⁵ Conversely, some cytotoxic chemicals that are currently used for anticancer therapy such as anthracyclines, oxaliplatin (but not cisplatin), and cyclophosphamide induce a type of cell death that is immunogenic, yet is accompanied by all known biochemical and morphologic hallmarks of apoptosis.^{53,54} Tumor cells that have been killed *in vitro* with such chemotherapeutic agents elicit a vaccine-like effect when they are injected subcutaneously into immunocompetent mice. This leads to the long-term protection of mice against subsequent rechallenges with live tumor cells of the same type. Cancer cells respond to DNA damaging agents (which constitute an important class of clinically used chemotherapeutics) with a complex signaling pathway that either allows for DNA repair (if the damage is limited) or engages apoptotic mechanisms (if the damage is excessive).⁵⁵ Prominent players of the DNA damage response include the ataxia telangiectasia mutated (ATM) and the ATM-related kinases, checkpoint kinases 1 and 2 (CHK1 and CHK2),⁵⁶ and the tumor suppressor protein TP53.⁵⁷

Beyond their role in the DNA damage response, ATM and CHK1 are known to induce the expression of natural killer (NK) cell group 2D (NKG2D) ligands, thus sensitizing tumor cells to NK-mediated lysis.^{58,61} In addition, TP53 might mediate NKG2D ligand-independent immunogenic effects by inducing cell senescence, a status that has been surmised to be recognized by NK cells and macrophages, leading to tumor eradication.⁶² Recently, multiple chemotherapeutic agents have been shown to up-regulate the expression of mannose-6-phosphate receptors on the surface of tumor cells, thereby promoting a perforin-independent increase in the permeability to granzyme B released by CD8⁺ lymphocytes.⁶³ Altogether, these observations suggest that multiple anticancer agents that are currently used in the clinical setting induce or at least facilitate ICD.

Some chemotherapeutic agents improve anticancer immunity by exerting direct immunomodulatory effects. For instance, cyclophosphamide which is widely used against lymphomas and leukemia selectively reduces the frequency of tumor-induced regulatory T cells,⁶⁴ induces the differentiation of T_H17 cells,⁶⁵ enhances the long-term survival and proliferation of lymphocytes,^{66,67} and resets DC homeostasis.^{54,68,71} Along similar lines, imatinib a tyrosine kinase inhibitor used in bcr/abl and Kit-induced malignancies activates NK-dependent antitumor effects in mouse models⁷² and stimulates the NK-mediated secretion of IFN- γ in patients with gastrointestinal stromal tumor, thus improving long-term survival.⁷³ Thus, several chemotherapeutic agents provide therapeutic benefits not only via tumor cell intrinsic, on-target effects but also by modulating immune responses in an off-target fashion.

RADIOTHERAPY AS A POTENT ANTICANCER VACCINE INDUCER

Focused ionizing radiations induce cancer cell death upon the induction of DNA damage and the overgeneration of reactive oxygen species.⁷⁴ For many years, the direct cytotoxic effect of radiotherapy has been considered as the sole determinant of its therapeutic success. However, multiple lines of evidence have accumulated suggesting that the therapeutic effects of radiotherapy cannot be accounted for by tumor cell death alone and hence might depend at least in part not only on endothelial cells but also on the host immune system.^{75,76} Thus, radiotherapy appears to be more efficient in immunocompetent mice than in their immunodeficient counterparts.⁷⁷ Moreover, the irradiation of primary tumors is known to inhibit the growth of nonirradiated metastases that are localized at distant sites (a phenomenon known as “abscopal effect”).⁷⁵ In line with this notion, the irradiation of primary 4T1 tumors (mouse breast cancer that spontaneously metastasizes) induced a CD8⁺ T cell-mediated immune response that controlled the growth of lung micrometastases when combined with an inhibitor of cytotoxic T lymphocyte-associated antigen 4 (CTLA-4 receptors (to overcome tumor-induced T-cell tolerance)).⁷⁸ Recently, these findings have been corroborated in other murine models of cancer, namely, in TSA-derived breast cancer and MCA-38Y-derived colon carcinoma. In these settings, fractionated radiotherapy (but not single dose) combined to a CTLA-4 blocking antibody led to potent abscopal effects that were paralleled by the production of consistent levels of IFN- γ .⁷⁹ Importantly, the frequency of CD8⁺ T cells elicited by radiotherapy appears to correlate with the intensity of the abscopal effect. Interestingly, radiotherapy influences the chemokine pattern of the tumor microenvironment (promoting CXCL16 secretion by irradiated tumor cells), facilitating the entry of effector CD8⁺CXCR6⁺ T cells into irradiated tumor beds.⁸⁰ Altogether, these studies demonstrate that radiotherapy induces a T cell-dependent antitumor effect, by inducing and/or recruiting tumor-specific T cells into the tumor bed.^{81,82}

Recently, we have demonstrated that tumor cells succumbing upon irradiation elicit a cognate tumor-specific response when injected subcutaneously into syngenic mice, thereby exerting a vaccine-like effect and protecting mice against subsequent challenges with living tumor cells of the same type.^{6,7} Irradiation-induced ICD, which might account for, at least part of, the abscopal effect,^{83,84} also turned out to rely on the preapoptotic exposure of CRT and the TLR4/Myd88 pathway (see above). Alternatively, it has been suggested that dormant antitumor immunity might get reactivated by radiotherapy-induced inflammation and cytokine release, which together would trigger the recruitment of T cells into the tumor bed.^{80,85} Irrespective of the fact that the molecular mechanisms underlying the abscopal effect remain poorly understood, radiotherapy appears as a potent trigger of ICD.

GENETIC BACKGROUND AND CLINICAL OUTCOME

During ICD, TLR4 and P2RX7 on DCs are critical for sensing and decoding the immunogenic message conveyed by the release of HMGB1 and ATP, respectively.^{10,12,24} Both TLR4 and P2RX7 present several nonsynonymous single nucleotide polymorphisms,^{86,87} and loss-of-function TLR4 and P2RX7 mutants (Asp299Gly and Gly496Ala, respectively) display low ligand-binding affinity. In line with the fact that TLR4 plays a critical role during ICD (see above), the TLR4 Asp299Gly allele has been shown to negatively affect the progression-free survival of breast cancer patients who received anthracycline-based adjuvant chemotherapy²⁴ as well as of patients bearing colorectal cancers that were treated with oxaliplatin-based regimens.⁸⁸ Furthermore, among breast cancer patients who carried wild-type TLR4, the P2RX7 Gly496Ala allele was associated with shortened progression-free survival.¹⁰ These results strongly suggest that the defect in the molecular mechanisms by which ICD is perceived by the immune system limits the efficacy of anticancer chemotherapy. This also provides further support to the concept that anticancer immune responses and hence all the genetic and environmental factors that affect such responses are crucial for therapeutic success.

In view of these considerations, it is tempting to speculate that detailed information on the patient and tumor genetic background would allow for the design of tailored anticancer regimens with optimal efficacy and limited adverse effects. In particular, such information might (at least partially) predict the proficiency of a tumor to undergo ICD and elicit a cognate immune response and, if required, suggest the development of interventions aimed at restoring the immunogenicity of cell death. For instance, defects in the ER stress module that is required for CRT exposure during CRT might be corrected by the direct absorption of recombinant CRT to the tumor.⁶ Along similar lines, TLR4 loss-of-function mutations (which result in the deficient perception of HMGB1-conveyed signals) might be compensated by combining chemotherapy with alternative TLR agonists or with lysosomal inhibitors such as chloroquine,²⁴ whereas defects in P2RX7 signaling might be reverted by the administration of exogenous IL-1A¹⁰ or apyrase inhibitors.

Alternative approaches for the elicitation of an anticancer immune response focus on the reversal of tumor-induced tolerance by means of immunomodulatory agents such as monoclonal antibodies targeting suppressive pathways (such as CTLA-4, PD-1, Lag3, Tim-3) or engaging activating receptors (such as CD40, CD27, 4-1BB), cytokines, and cell based approaches (T and DC) in combination with conventional therapies.^{89,91} In preclinical models, these strategies have been shown to enhance the vaccine-like effect of both chemotherapeutic and radiotherapeutic regimens, thereby constituting promising approaches.^{91,79} Gulley and colleagues⁹² reported that, in prostate cancer patients, the combination of radiotherapy with an admixture of a recombinant vaccine against prostate-specific antigen (PSA) and B7.1 costimulatory molecules induced a significant increase in PSA-specific and MUC1-specific T cells. The expansion of MUC1-specific T cells indicates that a tumor antigen cross-priming occurs in vivo following radiotherapy. Brody and colleagues reported a phase I/II trial where low-dose radiotherapy combined with in situ TLR9 agonists induced lymphoma remission, not only at the site of the treated lesion but also at distant sites, associated with detectable anticancer T-cell responses.⁹³

Taken together, these studies suggest that efficient anti-cancer regimens should combine immunogenic chemotherapy or radiotherapy with immunomodulatory agents that overcome tumor-induced immunosuppression. Moreover, whenever required, the defects in the molecular machinery for the execution and perception of ICD should be compensated to obtain the complete eradication of tumors and long-term tumor-free survival (Fig. 3).

PERSPECTIVES

| As we have discussed above, some (but not all) chemo-therapeutic and radiotherapeutic regimens induce the immu-nogenic death of tumor cells that, in specific circumstances, lead to the elicitation of a potent anticancer immune response. This vaccine-like effect is critical for both therapeutic success and long-term tumor-free survival. The ability of anticancer drugs to induce an ER-stress response that precedes cell death, the intrinsic capacity of tumor cells to emit immunogenic CDAMPs in a defined spatiotemporal order, and the ability of the host to perceive these signals and to overcome tumor-induced immunosuppression appear as fundamental prerequisites for the vaccine-like effect of radiotherapy or chemotherapy. We believe that future anticancer regimens should be tailored to each pa-tient and tumor's genetic background to take into account all these elements, as this will result in combination therapies with optimal cytotoxic and immunogenic profiles and limited adverse effects.

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FIGURE 1. Immunogenic signals emitted by dying cells form a spatiotemporal code unlocking DCs to mount a potent immune response toward tumor cells. (i) Early exposure of ecto-CRT by dying tumor cells, which facilitates engulfment by DCs. (ii) HMGB1 released from dying cells binds to TLR4 on DCs, thus favoring antigen cross-presentation and up-regulating proYIL-A (proYIL-1A). (iii) ATP liberated from dying cells binds to the purinergic receptor P2RX7 on DCs, activates the NLRP3 inflammasome, and leads to the secretion of active IL-1A, which polarizes CD8+ T cells toward IFN-F production. (iv) An additive DC maturation factor remains to be characterized.

FIGURE 2. Precise orchestration of antitumor T-cell responses elicited following ICD: After (immunogenic) chemotherapy, tumor material is phagocytosed by DCs. These later are also activated by ICD signals emitted by dying tumor cells. Within 2 days, IL-17 γ -producing $\gamma\delta$ T cells are recruited to the tumor bed in an IL-1 β -dependent manner. Their arrival precedes and correlates with the IFN- γ -producing CD8 $^{+}$ T cells infiltration, which is critical for tumor eradication.

FIGURE 3. Schematic view of tailored anticancer regimens: To achieve therapeutic success, (i) the anticancer regimen should be able to induce ICD, that is, to induce an ER stress before cell death and ATP release. Nonimmunogenic cytotoxic drug can be combined with ER stressors to restore immunogenicity. (ii) The tumor of the patient should have conserved the intrinsic capacities to emit all the immunogenic signals. If not, the defective signals could be identified and then compensate by recombinant CRT or rIL-1A or ATP superagonists. (iii) The loss-of-function mutation of key receptors involved in the perception of ICD signals might also be compensated by triggering alternative TLR pathway or supplementation with the appropriate cytokine. (iv) The combination of immunotherapy and immunogenic chemotherapy enhances the vaccine-like effect of chemotherapy and radiotherapy by overcoming the tumor-induced immunosuppression. Thus, tailored anticancer regimens should be designed by taking into account the genetic background of both the tumor and the host, with an aim to correctly unlock the immune system to obtain the complete eradication of the tumor and long-term tumor-free survival.