Neutrophils are essential for induction of vaccine-like effects by antiviral monoclonal antibody immunotherapies

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Using a mouse retroviral model, we have shown that mAb-based immunotherapy can induce life-long endogenous protective immunity (vaccine-like effects). This observation has potentially important consequences for treating life-threatening human viral infections. Here, we investigated the role of neutrophils in this effect. Neutrophils are innate immunity effector cells with wellestablished microbe-killing activities that are rapidly mobilized upon infection. They are also emerging as orchestrators of innate and adaptive immunities. However, their immunomodulatory activity during antiviral mAb immunotherapies has never been studied. Our data reveal that neutrophils have an essential role in immunotherapy-induced immune protection of infected mice. Unexpectedly, neutrophils have a limited effect in controlling viral propagation upon passive immunotherapy administration, which is mostly mediated by NK cells. Instead, neutrophils operate as essential inducers of a potent host humoral antiviral response. Thus, neutrophils play an unexpected key role in protective immunity induction by antiviral mAbs. Our work opens approaches to improve antiviral immunotherapies, as it suggests that preserving neutrophil functions and counts might be required for achieving mAb-induced protective immunity.

Introduction

Neutralizing mAbs are now considered as a potential therapeutic approach for the prevention and treatment of chronic and acute viral infections, including newly emerging viral infections (1). In recent years, there has been a dramatic increase in the development of new mAbs with improved neutralizing activity (1–14). Notably, in addition to antirespiratory syncytial virus (RSV) mAb used to treat infant respiratory disease, several mAbs directed against human cytomegalovirus, HIV, influenza, Ebola, and rabies virus are currently engaged in promising clinical trials (1, 3, 9, 14). While the direct effects of antiviral mAbs on viral propagation have been studied extensively, little attention has been paid to their potential immunomodulatory effects until recently (10, 15, 16).

Using a mouse model of persistent retroviral infection, i.e., the erythroleukemia induced by the murine leukemia virus, FrCasE, we have shown that treatment of infected mice with a highly neutralizing mAb can induce life-long protective immunity (vaccine-like effects) (see ref. 10). This observation may have major therapeutic implications for humans. Indeed, increased antiviral immune responses after mAb treatments were recently observed in preclinical models of HIV, RSV, and henipavirus infections (see ref. 10). Moreover, passive immunotherapy with broadly neutralizing mAbs stimulated antiviral antibody responses in HIV-1–infected patients (14). Similarly, vaccine-like effects have also been documented in preclinical models of cancer immunotherapies (17–19) as well as in clinical trials using anti-CD20, anti-EGFR, or anti-HER2 mAbs (20–23). Yet, neither the mechanisms at play nor the possible long-term protective consequences have been addressed in such clinical trials.

In humans, technical, ethical, and cost limitations strongly limit investigations. Instead, immunocompetent animals offer the possibility of identifying the cellular and molecular actors involved in the induction of vaccine-like effects by antiviral mAbs. The FrCasE-induced erythroleukemia is a model system permitting extensive analysis of the endogenous immune response after passive mAb-based immunotherapy under conditions of both chronic infection and pathological development. We found that a 5-day treatment of FrCasE-infected mice with the neutralizing mAb 667 (an IgG2a recognizing the retroviral envelope

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glycoprotein; Env) shortly after infection provides long-lasting (>1-year) protective antiviral immunity of the Th1 type with enhanced IgG2a humoral and cytotoxic T cell (CTL) responses and reduced regulatory T cell activity (24–29). The immunomodulatory action of mAb 667 depends on its ability to interact with receptors for IgGs (Fc γ R). Notably, immune complexes (ICs) formed between the administered mAb and viral determinants enhance antiviral CTL responses through Fc γ R-mediated binding to DCs (27).

Importantly, IC-Fc γ R interactions concern several Fc γ R-bearing innate effector cells other than DCs, including neutrophils that are rapidly recruited at sites of infections. While they have long been viewed as simple and direct scavengers of extracellular pathogens, there is recent evidence that neutrophils can be key cells in the orchestration of innate and adaptive immunity via the interaction with different myeloid and lymphoid cells and the modulation of their functions (30, 31). However, their role in antiviral immunity is still largely unknown. Indeed, most investigations have studied neutrophil functions in viral pathogenesis or in the control of viral propagation through their effector functions, i.e., phagocytosis, reactive oxygen species production, neutrophil extracellular traps formation, etc. (32, 33). In contrast, the immunomodulatory properties of neutrophils have been poorly studied in viral infections and not addressed in the context of antiviral immunotherapies.

Here, we used the FrCasE model to address the general question of neutrophil participation in the induction of protective immunity by antiviral mAbs. We found that they are essential for the protection of infected, 667 mAb-treated mice. Unexpectedly, and unlike NK cells, neutrophils had a limited effect on viral propagation control. Instead, their protective effect relied on their capacity to induce potent humoral responses due to the acquisition of B cell helper activity upon mAb treatment. Our findings should facilitate improved design of mAb-based antiviral therapies, as they suggest that preserving neutrophil functions and counts might be required for achieving optimal protection by mAb.

Results

Neutrophils are necessary for protection of mice infected by antiviral mAbs without any effect on viral propagation. We first addressed neutrophil mobilization in FrCasE-infected mice with, or without, 667 mAb treatment (infected/treated vs. infected/nontreated). Passive immunotherapy was administered on the same day after establishment of viral infection (27) (Figure 1A) and at days 2 and 5 postinfection (p.i.). Age-matched naive mice were used as controls. Neutrophil recruitment was assessed in the spleen, one of the most viremic organs, at day 8 p.i, when viral replication is maximal in infected/nontreated mice (27). Neutrophil abundance in infected/treated mice was comparable to that observed in naive mice, whereas it was significantly higher in infected/nontreated animals (Figure 1B). Interestingly, this increased neutrophil frequency was associated with a higher percentage of infected spleen cells (Figure 1C), as assessed by flow cytometry using the H34 antibody (recognizing a Gag protein epitope expressed on the surface of FrCasE-infected cells) (34–36).

Next, we assessed the role of neutrophils in the control of viral propagation as well as in the protection against leukemia in infected mice with or without 667 mAb treatment. To this end, neutrophils were depleted by administering a mAb (1A8) directed to their specific Ly6G cell surface marker (37, 38) or an isotype control mAb (2A3). Depletion started 1 day before infection (Figure 1A), was efficient and specific (Supplemental Figure 1; supplemental material available online with this article; https://doi.org/10.1172/jci. insight.97339DS1), and was maintained for 21 days, i.e., the time necessary to eliminate the therapeutic 667 mAb (27). Neutrophil elimination accelerated disease development in infected/nontreated mice and drastically reduced protection provided by 667 mAb to infected mice (Figure 1D). We next assessed viral propagation in the different groups of mice. In infected/nontreated animals, neutrophil depletion (Figure 2A) was associated with a significant increase in the percentage of infected spleen cells at days 8 (Figure 1E) and 14 p.i. (Supplemental Figure 2A) as well as with a higher viremia (Supplemental Figure 2B). In contrast, in infected/ treated mice, viral propagation was not significantly affected at days 8 (Figure 1E) and 14 p.i. (Figure 2B and Supplemental Figure 2) upon neutrophil depletion and remained lower than in infected/nontreated mice. This suggested that viral control by 667 mAb involved other innate immunity effector cells.

Thus, neutrophils exert different antiviral effects on FrCasE-infected mice depending on immunotherapy. In animals undergoing simple infection, neutrophils participate in the control of viral propagation. Instead, in infected/treated mice, they are crucial during the immunotherapy period for generation of longterm protection against leukemia, despite their limited effect on viral propagation.

NK cells control viral propagation in infected / treated mice. As NK cells can exert antibody-dependent cellular cytotoxicity (ADCC) activity against infected cells exposing determinants such as the retroviral Env protein



Figure 1. Antiviral effects of neu-

trophils. (A) Experimental scheme. Eight-day-old pups were infected and treated with the 667 mAb as indicated. Mice were treated as indicated with the anti-Ly6G 1A8 mAb or the isotype control 2A3 mAb in neutrophil depletion experiments. (B and C) Neutrophil recruitment and infected cell rate in spleen. Splenocytes from naive, infected/nontreated (I/NT), and infected/ treated (I/T) mice were analyzed by flow cytometry on day 8 p.i. for (a) neutrophil recruitment (percentage of Ly6G⁺ cells) and (b) retroviral positivity of splenocytes (percentage of Gag⁺ cells) gated in the CD45.2⁺ population. The data presented correspond to 5 independent experiments, with at least 15 mice per group. (D) Mouse survival. Naive, I/NT, and I/T mice were treated with either the antineutrophil (1A8) or the control (2A3) mAb as indicated in A and followed up for leukemic death. The data represent 2 independent experiments, with 6-9 mice per group. (E) Infected cells rate upon neutrophil depletion. Neutrophils of naive, I/NT, and I/T mice were depleted, or not, as indicated in A and infected splenocytes were assayed as in **C** on day 8 p.i. The data represent 4 independent experiments, with 9-15 mice per group. Data are expressed as mean ± SEM. Statistical significance was established using a parametric 1-way ANOVA test with a Bonferroni correction (*P < 0.05; **P < 0.01; ***P < 0.001).

(27, 39, 40), we asked whether NK cells were involved in the control of viral propagation in infected/treated mice. To this end, NK cells were depleted using an anti-asialo-GM1 antibody (41–44) (Figure 2A). In contrast to neutrophil depletion, the absence of NK cells in infected/treated mice led to a significantly increased viral propagation at day 14 p.i. (Figure 2B). Next, we compared 667 mAb–mediated ADCC activity of NK cells and neutrophils with infected cells using an in vivo antibody-mediated killing assay, relying on the administration of 667 mAb–opsonized FrCasE-infected splenocytes in naive mice (27) (Figure 2D). Depletion of NK cells, but not neutrophils, led to reduced mAb-mediated infected cell lysis (Figure 2D). Finally, as NK cells are crucial for viral propagation control during the immunotherapy period, we assessed their role in the protection against leukemia in infected/treated mice. Importantly, NK cell depletion drastically reduced survival of immunotreated mice (Figure 2E), indicating that antibody-mediated control of viral propagation by NK cells is necessary for long-term antiviral protection.

Thus, NK cells are crucial for protection of infected/treated mice via efficient control of viral propagation by 667 mAb-mediated ADCC.

Neutrophils differentially alter innate lymphoid cell biology in infected/treated and infected/nontreated mice. Innate lymphoid cells (ILCs) are a heterogeneous population of immune cells that includes NK cells and



Figure 2. Antibody-mediated control of viral propagation by NK cells. (A) Experimental scheme. Mice were infected and mAb-treated as in Figure 1A. Infected/treated mice were treated as indicated with the anti-LyGG 1A8 mAb or the isotype control 2A3 mAb to deplete neutrophils and infected/treated mice were treated as indicated with the anti-asialo-GM1 antibody to deplete NK cells. (B) Effect of neutrophils or NK cell depletion in viral spread in infected/treated mice. Percentage of infected cells at day 14 p.i. in the spleens of naive, I/NT, and I/T mice, depleted or not of neutrophils or NK cells assessed as in Figure 1C. The data represent 3 independent experiments, with at least 8 mice per group. (C and D) In vivo cytolysis activity of 667 mAb in naive mice after depletion of neutrophils or NK cells. Splenocytes from noninfected mice (Sp) were labeled using 0.5 μ M of the vital dye CFSE (CFSE^{Io} cells; M1) and mixed at a 1:1 ratio with splenocytes from infected mice (Infected-Sp) labeled using 5 μ M CFSE (CFSE^{Iii} cells; M2) and preincubated, or not, with 667 mAb. Mixed cell populations were administered to naive mice 1 day after depletion of either neutrophils or NK cells with the 1A8 mAb or the anti-asialo-GM1 antibody, respectively. Cytolysis was quantified 5 hours later, as described in Methods section. The data are presented as mean ± SEM of 2 independent experiments, with at least 8 mice per group. Statistical significance was established using a parametric 1-way ANOVA test with a Bonferroni correction (B and D). (E) Effect of NK cell depletion in the survival of infected/ treated mice. I/T, NK cells, depleted or not as indicated in **A**, were followed up for leukemic death. The data represent 2 independent experiments, with 7 mice per group. Statistical significance was established using an unpaired Student's *t* test. Data are expressed as mean ± SEM (**P* < 0.05, ***P* < 0.01,****P* < 0.001).



Figure 3. Effects of neutrophil depletion on innate lymphoid cell recruitment and biology. (A-D) Neutrophils of naive, I/ NT. and I/T mice were depleted, or not. as indicated in Figure 1A, and ILC in the spleen were assayed 14 days p.i. by flow cytometry. (A) Frequency of CD3-NKp46+ cells in the CD45.2⁺ leukocytic population. (B) Frequency of CD117⁺/CD127⁺ cells in the Lin⁻NKp46⁺ population. (C) Maturation (CD11b⁺ cells) and (**D**) expression of IFN-γ in the CD3⁻NKp46⁺ population. (E) In vivo cytolysis activity of 667 mAb in infected/treated mice after depletion of neutrophils. The 1A8, or the 2A3 isotype control mAb, was administered to I/T mice, and 667 ADCC activity was quantified at 30 days p.i., as in Figure 2, C and D. The data represent at least 2 independent experiments. Data are expressed as mean ± SEM. Statistical significance was established using a parametric 1-way ANOVA test with a Bonferroni correction (*P < 0.05; **P < 0.01; ***P < 0.001).

ILC1, ILC2, and ILC3. As NK biology can be affected by neutrophils (45), we first assessed the effect of neutrophil depletion on NK cells. To this end, we measured the frequency of CD3⁻NKp46⁺ cells in the spleens of mice from the different groups. The CD3⁻Nkp46⁺ population contains mostly NK cells but may also include ILC1 and a subpopulation of ILC3 cells. NK cells can be distinguished from such ILCs using the cell surface marker CD49b (46). Notably, over 95% of CD3⁻NKp46⁺ cells were CD49b⁺ (Supplemental Figure 3), indicating that the vast majority of CD3⁻NKp46⁺ cells in the spleen displayed a NK phenotype. As shown in Figure 3A, CD3⁻NKp46⁺ cell recruitment in spleens of both infected/nontreated and infected/ treated mice at day 14 p.i. was similar and stronger than in naive mice. We also observed that neutrophils were involved in CD3⁻NKp46⁺ cell mobilization in infected/nontreated mice but not in infected/treated or in naive mice (Figure 3A). We next assessed the frequency of splenic ILCs other than NK cells by quantifying CD117 and CD127 expression in the NKp46⁺ spleen cell population lacking the common lymphoid and myeloid lineage-associated (Lin-associated) markers (Lin⁻) (47). The recruitment of those ILCs in spleens

of both infected/nontreated and infected/treated mice at day 14 p.i. was not significantly different from that observed in naive mice (Figure 3B). In addition, neutrophil depletion did not significantly alter the frequency of such ILCs (Figure 3B) in any group of mice.

These results show an enhanced recruitment of CD3⁻NKp46⁺ cells, which are mostly NK cells, in both infected/treated and infected/nontreated mice, but these effects are neutrophil dependent only in the latter. They also show that neutrophil depletion does not affect the frequency of splenic Lin⁻CD117⁺CD127⁺ ILCs, in agreement with the lack of effect of anti-Ly6G–mediated neutrophil depletion on splenic ILCs reported in other experimental settings (47).

To better characterize the effect of neutrophils on splenic CD3⁻NKp46⁺ cells, we assessed their maturation by monitoring CD11b and CD27 expression at day 14 p.i. These markers identify different stages of NK maturation in mice (45). Relative to naive mice, both infected/nontreated and infected/treated mice showed similar higher frequencies of CD11b⁺CD3⁻NKp46⁺ cells (including fully mature CD11b⁺CD27⁻ and semimature CD11b⁺CD27⁺ cells) (Figure 3C) at the expense of immature ones (CD11b⁻) (data not shown). Moreover, depletion of neutrophils entailed a strong reduction of the frequency of CD11b⁺CD3⁻ NKp46⁺ cells only in infected/nontreated mice. Further characterization of CD3⁻NKp46⁺ cells indicated higher neutrophil-dependent IFN-γ production in infected/treated mice but low and similar IFN-γ levels in naive and infected/nontreated animals, whether the latter were neutrophil proficient or deficient (Figure 3D). Finally, we assessed whether neutrophils affected the ADCC activity of NK cells in infected/ treated mice. Notably, ablation of neutrophils did not alter 667 mAb–mediated ADCC activity of NK cells (Figure 3E), consistent with the lack of effect on viral propagation (Figure 1E and Figure 2B).

Thus, CD3⁻NKp46⁺ maturation is similarly enhanced in both infected/treated and infected/nontreated mice, but these effects are neutrophil dependent only in the latter. Nevertheless, neutrophils might participate in functional activation of CD3⁻NKp46⁺ cells in infected/treated animals, as our results show that neutrophils modulate their IFN-γ secretion capacity in infected/treated mice.

Neutrophils are crucial for inducing the humoral, but not the $CD8^+$ T cell, antiviral response in immunotherapy-treated mice. Since neutrophils exert no control over viral propagation during the immunotherapy period (Figure 1E and Figure 2B) but are necessary for long-term protection of mice (Figure 1D), we asked whether they could be key for the induction of vaccine-like effects.

We first addressed cellular adaptive immunity by assaying the primary virus-specific CD8⁺ T cell response in infected/nontreated and infected/treated mice at its peak (i.e., 14 days p.i.) (27) with or without neutrophil depletion. Consistent with our previous work, infected/treated mice showed an increased virus-specific CD8⁺ T cell response relative to infected/nontreated mice (27). In neither case was the frequency of virus-specific CD8⁺ T cells altered by neutrophil depletion (Figure 4A). Similarly, neither case showed any difference in the frequency of CD8⁺ T cells expressing IFN- γ (Figure 4B). These data ruled out a major role for neutrophils in the 667 mAb–induced antiviral CD8⁺ T cell response.

Next, we addressed humoral immunity in infected/nontreated and infected/treated mice depleted or not of neutrophils. To achieve this, anti-FrCasE serum Igs from mice of the different groups were assayed by ELISA. Virus-specific IgM titers were assayed at 14 days p.i. (i.e., at the peak of the IgM response upon FrCasE infection and 667 mAb treatment; Supplemental Figure 4) and were not significantly different between infected/nontreated and infected/treated mice (Figure 5A). Interestingly, neutrophil depletion did not alter virus-specific IgM titers in either infected/nontreated or infected/treated mice (Figure 5A). In agreement with published studies in other experimental settings, the absence of neutrophils did not affect the levels of IgM (48). On the contrary, and consistent with our previous work (27), infected/treated mice displayed a much higher level of anti-FrCasE IgGs than infected/nontreated animals (Figure 5B), with a peak (700 μ g/ml) by day 68 p.i. and a still high level (100 μ g/ml) at the end of the experiment. Interestingly, in the absence of neutrophils, serum concentration of antiviral IgGs dramatically decreased in infected/ treated mice (Figure 5B and Supplemental Figure 5). In contrast, neutrophils showed no role in the regulation of the poor nonprotective antibody response elicited in infected/nontreated mice (Figure 5B). Importantly, high anti-FrCasE IgG seric levels correlated with longer survival times, supporting a role for the high humoral antiviral response in protection against disease (Figure 5C). To further characterize the long-term virus-specific humoral response, we tested whether infected/treated mice, depleted or not of neutrophils, could respond to a virus challenge performed 3 months after the first infection (i.e., a time point at which the primary humoral response has strongly declined). Infected/treated mice, depleted or not of neutrophils, were inoculated with FrCasE, and serum samples were collected 1 week later to assay the generation of



Figure 4. Assay of FrCasE-specific CD8⁺ T cells in the presence and absence of neutrophils. (A) Neutrophils of naive, I/NT, and I/T mice were depleted, or not, as indicated in Figure 1A. Frequency of FrCasE-specific CD8⁺ T cells. Spleen cells were isolated at day 14 p.i., and the frequency of virus-specific CD8⁺ T cells in the total CD8⁺ T cells population was assayed by flow cytometry using the H2D^b-GagL MHC tetramer. The data represent 4 independent experiments, with at least 11 mice per group. (B) Expression of IFN- γ by CD8⁺ T cells. Splenic CD8⁺ T cells were analyzed by flow cytometry for the expression of IFN- γ . The data presented represent 3 independent experiments, with at least 7 mice per group. Data are expressed as mean ± SEM. Statistical significance was established using a parametric 1-way ANOVA test with a Bonferroni correction (**P* < 0.05).

endogenous anti-FrCasE antibodies. In the presence of neutrophils, 3 of 5 infected/treated mice showed an increase in the virus-specific humoral response (Figure 5D), while, in the absence of neutrophils, none of the infected/treated mice responded to the viral challenge. Altogether, these results show that neutrophil depletion at early time points after infection and treatment affects humoral immunity with an effect on both primary and memory virus-specific B cell responses.

As marginal zone (MZ) B cells are known to contribute to the mounting of antibody responses (49, 50) and their function might be modulated by neutrophils (51), we next addressed the frequencies of splenic MZ (CD21^{hi}IgM^{hi} CD19⁺ cells) and follicular (FO) (CD23⁺IgM^{lo} CD19⁺ cells) B cells. MZ B cell frequency was enhanced in a neutrophil-dependent manner in infected/treated mice but unchanged in infected/nontreated animals relative to naive mice (Figure 6A). In contrast, the frequency of FO B cells was significantly enhanced in infected/treated mice relative to that in naive mice; however, there was no role for neutrophils (Figure 6B). Interestingly, consistent with a role of MZ



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Figure 5. Enhancement of the humoral antiviral response by neutrophils.

Neutrophils of naive, I/NT, and I/T mice were depleted, or not, as indicated in Figure 1A. (A and B) Serum concentration of FrCasE-specific Igs. (A) Seric FrCasE-specific IgM levels were assayed by ELISA at 14 days p.i. The data represent 2 independent experiments, with 8-11 mice per group (for I/NT and I/T mice) and 3-6 mice per group (for naive mice). (B) Seric FrCasE-specific IgG concentration was assayed by ELISA at the indicated times. The data represent 2 independent experiments, with 7-9 mice per group. Data are expressed as mean ± SEM. Statistical significance was established using a parametric 1-way ANOVA test, with a Bonferroni correction. (C) Correlation between serum anti-FrCasE IgG levels (evaluated as AUC) and survival times, analyzed using the Pearson correlation test. AUC was evaluated until the last time point at which all mice were still alive (day 68 p.i.). All infected/nontreated (n = 8) and infected/treated (n = 9) mice, depleted or not of neutrophils (n = 9 and)n = 7, respectively), showed in Figure 1D were evaluated for such a correlation. (D) FrCasE-specific secondary humoral response. Seric FrCasE-specific IgG levels in I/T mice (depleted or not of neutrophils) were assayed by ELISA before and 1 week after a viral challenge performed at day 93 p.i. The data represent 2 independent experiments, with 5 mice per group. Statistical significance was established using a paired Student's t test (*P < 0.05; **P < 0.01; ***P < 0.001).

B cells in the generation of plasma cells (49), the higher frequency of splenic MZ B cells at day 14 p.i. in infected/treated mice was associated with a higher frequency of CD138⁺ plasma cells in bone marrow (BM), as compared with infected/nontreated animals. This effect was neutrophil dependent (Figure 6C). Finally, histological analyses of spleens of infected mice at 14 days p.i. revealed larger germinal centers (GCs; defined by staining of MZ CD169⁺ macrophages and B220⁺ cells) in infected/ treated versus infected/nontreated mice (Figure 6D). In agreement with a role of neutrophils in the enhancement of the humoral response, neutrophil depletion in infected/treated mice led to smaller GCs (Figure 6D).

Thus, neutrophils are essential for the vaccine-like effects induced by the 667 mAb immunotherapy through the stimulation of the humoral, but not the CD8⁺ T cell, antiviral response. This effect is associated with a neutrophil-dependent increase in both splenic MZ B cells and BM plasma cells.

Neutrophils are differently activated in infected/treated and nontreated mice. Finally, we addressed the functional activation of splenic neutrophils in infected mice with or without 667 mAb immunotherapy at day



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Figure 6. Effects of neutrophil depletion on B cell responses. Neu-

trophils of naive, I/NT, and I/T mice were depleted, or not, as indicated in Figure 1A. (A and B) Frequency of MZ and follicular (FO) B cells. Spleen cells were isolated at day 14 p.i. and were analyzed by flow cytometry for the frequency of MZ (CD21^{hi}lgM^{hi}) (A) and FO (CD23⁺IgM¹⁰) (B) CD19⁺ B cells, as depicted in the gating strategy. (C) Frequency of plasma cells. BM cells were isolated at day 14 p.i. and were analyzed by flow cytometry for the frequency of CD138⁺ (CD19⁺) B cells. The data represent 5 independent experiments, with 7-12 mice per group for naive mice and 17-21 per group for I/NT and I/T mice. Data are expressed as mean ± SEM. Statistical significance was established using a parametric 1-way ANOVA test with a Bonferroni correction (*P < 0.05; ***P* < 0.01; ****P* < 0.001). (D) Histological analyses of spleen sections. Immunolabeling of B cells (B220⁺) and macrophages of the MZ (CD169⁺) was performed in sections from spleens of infected/nontreated and infected/treated mice (depleted or not of neutrophils) recovered at 14 days p.i. to visualize germinal centers. The images are representative of 4 separate mice for each experimental condition. Scale bar: 200 μ m.

8 p.i. by monitoring cell surface activation markers and quantifying various cytokine mRNA and protein levels. In infected/treated and infected/nontreated mice, splenic neutrophils were similarly activated, as deduced from CD62L shedding and increased expression of CD11b (Figure 7A). However, neutrophils from infected/treated mice showed a significantly stronger expression of two cytokines involved in B cell activation, B cell–activating factor (BAFF) and lymphotoxin α (LT α), compared with infected/nontreated animals (Figure 7B), suggesting a role for the therapeutic mAb in the functional activation of neutrophils. Splenic neutrophils sorted from infected/treated mice also showed a trend for higher secretion capacity of BAFF and LT α , as compared with infected/nontreated mice neutrophils, even though such a trend did not reach statistical significance. To further assess whether mAb triggering (through Fc γ R cross-linking) could enhance BAFF and LT α release by neutrophils, we isolated BM neutrophils from naive mice and stimulated them for 24 hours in plates coated with the 667 mAb. Experiments were conducted in the presence or the absence of the proinflammatory cytokine IFN- γ to assess the extent to which inflammatory conditions could synergize with 667 mAb triggering. Interestingly, in the presence of IFN- γ , 667 mAb triggering led to a significantly increased release of BAFF and LT α by neutrophils (Figure 7C).



Figure 7. Activation of splenic and BM-isolated neutrophils. (A) Expression of

CD11b and CD62L. Spleen cells from naive, I/ NT. and I/T mice were isolated at day 8 p.i. and were analyzed by flow cytometry for assaying cell surface expression of CD11b and CD62L. The data represent 5 independent experiments, with at least 18 mice per group. Data are expressed as mean ± SEM. (B) Expression and protein release of BAFF and $LT\alpha$ by neutrophils. Neutrophils from naive, I/NT, and I/T mice were sorted from spleens at day 8 p.i. and assessed for cytokine expression or protein release. Cytokine expression (left) was assessed by RT-qPCR normalized to β -actin. The data show fold changes in cytokine expression by neutrophils from I/NT and I/T mice as compared with naive mice and are representative of 3 independent experiments, with 8-10 mice per group. Protein release (right) was assessed by ELISA in supernatants of sorted neutrophils cultured at a density of 2 \times 10⁵ cells/well for 24 hours. The data show BAFF and LT α release by neutrophils from I/ NT and I/T and are representative of 3 independent experiments, with 8-10 mice per group. The dashed line represents the level of BAFF released by neutrophils sorted from naive mice. No $LT\alpha$ release was detected from neutrophils sorted from naive mice. (C) BAFF and LT a release by BM-isolated neutrophils. BAFF and $LT\alpha$ release was assessed by ELISA in supernatants of neutrophils isolated from BM of naive mice (>95% purity) and cultured for 24 hours in 667 mAb-coated 24-well plates at a density of 2 × 10⁶ cells in 500 µl medium. Experiments were done in the presence and in the absence of the proinflammatory cytokine IFN-γ (100 ng/ ml). 667 mAb-noncoated plates were used as control. The data represent 4 independent experiments. Data are expressed as mean ± SEM. Statistical significance was established using a parametric 1-way ANO-VA test with a Bonferroni correction (A and C) or a paired Student's t test (B) (*P < 0.05;</p> ***P* < 0.01; ****P* < 0.001).

In summary, FrCasE infection activates neutrophils independently of immunotherapy. However, the activation state substantially differs between the two groups of animals, notably with a significantly stronger expression of $LT\alpha$ and BAFF in immunotherapy-treated mice. In addition, inflammatory conditions synergize with mAb-mediated activation of neutrophils leading to an enhanced release of BAFF and LT α .

Discussion

We have previously shown that antiviral mAb-based therapies can induce life-long protective immunity. This finding potentially has important therapeutic implications, as evidence suggests that it may also apply to diverse severe human viral diseases (10, 14). A paramount task is now to identify the mechanisms at the origin of the mAb-induced vaccine-like effects and exploit them for more efficient mAb-based treatment of patients.

Here, we report that neutrophils are essential during the immunotherapy period for long-term survival of FrCasE-infected mice, not because they control viral propagation but because they are crucial for inducing a

protective humoral response without an effect on the CD8⁺ T cell response. In contrast, we show that, upon 667 mAb treatment, NK cells are crucial for the elimination of infected cells by 667 mAb-mediated ADCC activity and are necessary for long-term survival of infected/treated mice. Thus, our work indicates that both innate effector cells have distinct but complementary roles in the protection of infected mice by mAb; NK cells have an early and predominant role in the control of viral spread, while neutrophils are essential for the emergence of a potent host antiviral humoral response. Our findings contrast with the current view of neutrophils — they are usually considered as simple frontline agents against invading pathogens — and highlight the hitherto unreported role of neutrophils as key cells in the modulation of adaptive antiviral immunity upon mAb treatment.

Our data indicate that survival of infected/treated mice depends on an efficient anti-FrCasE humoral response. Infected/treated mice show neutrophil-dependent (a) increased frequency of MZ B cells, (b) enhanced formation of GCs, (c) increased plasma cell generation, and (d) enhanced production of antiviral IgGs. Interestingly, the absence of neutrophils during the immunotherapy period in infected mice also leads to impaired development of secondary humoral responses upon viral challenge. Overall, our data suggest that neutrophils are essential for antiviral protection due to their B cell helper activity. Similarly, such a helper function has already been documented under homeostatic conditions (51), in autoimmunity disease-prone mice (52), during emergency granulopoiesis (53), and in bacterial infection (54). Importantly, our findings show that the acquisition of B cell helper functions in infected mice is dependent on immunotherapy, as no modulation of the anti-FrCasE humoral response was detected in infected/nontreated mice upon neutrophil depletion.

Though not excluding the role of other factors, our study suggests potential roles for BAFF and LT α in this B cell helper function. As concerns BAFF, it is interesting to note that its secretion by splenic neutrophils can contribute to the activation of splenic MZ B cells and the acceleration of plasma cell generation (51, 55). Moreover, BAFF administration to mice increases both the frequency of MZ B cells and antibody production (56, 57) and constitutes a signal for both MZ B cell survival and differentiation into plasmablasts (47, 58–61). Together with the fact that MZ B cells can favor the generation of plasma cells upon microbial infection (49), it is reasonable to speculate that BAFF induction in neutrophils of infected/treated mice may also favor the MZ B cell response and the subsequent generation of plasma cells. Interestingly, in agreement with our in vitro results showing enhanced BAFF release by neutrophils after a combination of IFN- γ stimulation and Fc γ R cross-linking (Figure 7C), different proinflammatory stimuli, including ICs, have been shown to act as secretagogues and to synergize with IFN- γ to enhance BAFF secretion by human neutrophils (62). Moreover, neutrophil depletion in autoimmunity-prone mice led to a reduction of autoantibodies titers that correlated with decreased serum levels of IFN- γ and BAFF (52). Thus, it can reasonably be hypothesized that inflammatory conditions synergize with ICs formed after 667 mAb treatment of FrCasE-infected mice to enhance BAFF release by neutrophils.

Although not formally shown, our data suggest that LT α release by neutrophils might play a role in the enhancement of the antiviral humoral response in infected/treated mice. LT α is involved in the formation of secondary lymphoid organs, is expressed by lymphocytes, and mediates a large variety of inflammatory and antiviral responses. It has also been reported to play a role in the development of GC formation and to be required for IgG responses (63–65). Here, we show that this cytokine is expressed and released by neutrophils upon mAb treatment of infected mice. Interestingly, although LT α is not known to be expressed by neutrophils, it has been proposed that nonlymphocytic murine splenic cells are able to produce it (66). We also report that Fc γ R cross-linking by immobilized 667 mAb synergizes with IFN- γ to enhance LT α release by neutrophils. This suggests that, similarly to BAFF, inflammatory conditions and mAb triggering might lead to increased LT α secretion by neutrophils. Further studies will be required to address whether BAFF and LT α combine their actions to stimulate the antiviral humoral response upon mAb immunotherapy. Finally, as IFN- γ potentiates BAFF and LT α release, it will also be important to assess whether and how neutrophil-dependent IFN- γ production by NK cells in infected/treated mice contributes to the stimulation of antibody responses, as this cytokine affects class-switching and long-term maintenance of neutralizing antibody titers in retrovirally infected mice (34) and humoral autoimmunity in humans (67).

Identifying neutrophils as key players in the induction of protective immunity by antiviral mAbs has important therapeutic implications for several reasons. First, if combination therapies are considered, it will be of utmost importance that the agent(s) coadministered with the passive immunotherapy do not alter neutrophil functions and counts to avoid inhibiting efficient antiviral humoral immune

responses. Second, in pathological situations leading to neutropenia and/or impaired neutrophils functions, such as certain viral infections (68, 69) and/or drug-induced neutropenia (70), it will be essential to restore them. This could, for example, be achieved through administration of granulocyte colony-stimulating factor (G-CSF), a cytokine already used in the clinic to treat neutropenic patients. Interestingly, this cytokine might also be used in combination with treatments aiming at enhancing the generation of functional neutrophils (70). Furthermore, beyond stimulating neutrophil activation and/or mobilization, G-CSF enhances neutrophil BAFF secretion capacity and, thereby, the ability of neutrophils to stimulate B cells (71). Similarly, reduction of viral load by antiretroviral therapies also permits partial restoration of the impaired functions of neutrophils observed in HIV-infected patients (68). Interestingly, neutrophils have been shown to mediate immunosuppression via the PD-L1/PD-1 pathway in HIV-infected patients (72). Antiviral passive immunotherapies might therefore benefit from combination with the administration of mAbs targeting this immune checkpoint. Finally, alternative approaches, such as engineering the Fc fragment of antiviral mAbs, merit consideration. Increasing their affinity for the FcyRs expressed by neutrophils might, at first, allow superior antibody-mediated phagocytosis, as recently reported in the case the Fc-modified VR01 anti-HIV mAb (73). This could then alter cell signaling and cytokine/chemokines production to ultimately lead to more effective adaptive immune responses. Thus, stimulating neutrophil activity, restoring their impaired functions, and/ or counteracting their immunosuppressive actions should improve the vaccine-like effects of antiviral mAb-based immunotherapies. This might also apply to cancer treatment as enhancement of antitumoral immune responses has also been observed in mAb-based anticancer immunotherapies. This is all the more important to take into consideration, as neutrophils have been shown to play a role in the therapeutic activity of anticancer mAb (74-78).

Methods

Mice. Mice were bred and maintained under conventional, pathogen-free facilities at the Institut de Génétique Moléculaire de Montpellier.

Viral stocks. FrCasE viral stocks were produced, assayed, and stored as described previously (27).

Viral infection, immunotherapy, and mouse follow-up. Eight-day-old 129/Sv/Ev mice were infected i.p. with 50 µl of a virus suspension containing 50,000 focus-forming units (FFU) and treated, or not, with 30 µg 667 mAb 1 hour p.i. and on days 2 and 5 p.i. by i.p. administration. Mice were examined at regular intervals for clinical signs of erythroleukemia (reduction of hematocrits). They were euthanized when their hematocrits reached 35% (experimental endpoint).

Flow cytometry. Spleen single-cell suspensions were obtained by mechanical dissociation of the organs in PBS. BM cell suspensions were obtained by dissection and PBS flushing of tibias and femurs. Cells were stained at 4°C using fluorochrome-conjugated antibodies against CD3e (145-2C11, BD Biosciences), CD4 (RM4-5, eBioscience), CD8 (Ly2, 53-6.7, eBioscience), CD11b (M1/70, eBioscience), CD11c (HL3, BD Biosciences), CD19 (1D3, eBioscience), CD21/35 (eBio8d9, eBioscience), CD23 (B3B4, eBioscience), CD27 (LG3A10, BioLegend), CD45.2 (104, BD Biosciences), CD45R/B220 (RA3-6B2, BD Biosciences), CD49b (DX5, BD Biosciences), CD62L (Ly22, MEL-14, BD Biosciences), CD117 (2B8, BD Biosciences), CD127 (SB/199, BD Biosciences), CD138 (281-2, BioLegend), F4/80 (BM8, BioLegend), Gr1 (RB6-8C5, BD Biosciences), IFN- γ (XMG1.2, BD Biosciences), IgM (eB121-15F9, eBioscience), Ly6G (1A8, BD Biosciences), lineage (Ter119; Gr1; CD45R/B220; CD11c; F4/80; CD3e, BioLegend), and NKp46 (29A1.4, BD Biosciences). FrCasE-infected cells were assayed using an anti-Gag mAb (H34) (35) labeled with Alexa Fluor 647. Forward scatter area and forward scatter time-of-flight, as well as side scatter, were used to remove doublets from flow cytometry analyses. Cells were analyzed on FACSCanto II flow cytometer (BD Bioscience), and the data were analyzed using the FlowJo software (Tree Star).

ELISA of anti-FrCasE antibodies. Plasma anti-FrCasE Igs were assayed by ELISA as previously described (26, 27). Peroxidase-conjugated anti-mouse IgG or IgM rabbit antisera (Serotec) were used as secondary antibodies.

Virus challenge experiments. Infected/treated mice, depleted or not of neutrophils, were injected i.v. with 300 μ l of a 5 × 10⁴ FFU/ml FrCasE suspension mixed to 2 × 10⁶ FrCasE-infected splenocytes 3 months after the first infection. Blood samples were collected 1 week after challenge to assay endogenous anti-Fr-CasE IgG concentrations by ELISA.

In vivo depletion of Ly6G⁺ and NK cells. Neutrophils were depleted by administering a rat anti-Ly6G antibody (1A8; BioXcell), (37, 38) injected i.p. at different time points (150 µg/injection), or isotype control rat IgG (2A3; BioXcell). Neutrophil depletion was monitored by flow cytometry analysis of Gr1^{hi} and CD11b⁺ cells. NK cells were depleted using the anti-asialo GM1 antibody (Wako Pure Chemical Industries Ltd.), injected i.p. at different time points (50 μ l/injection). This antibody has been used to study the in vivo functions of NK cells in mouse strains lacking the NK1.1 allotype, which is a feature of 129 Sv/Ev mice (42).

In vivo cytolysis activity. Experiments were conducted as described previously (27, 79). Briefly, red blood cell–free splenocytes were recovered from 10-day-old FrCasE-infected, or noninfected, pups. Splenocytes from noninfected mice were labeled with the vital dye carboxyfluorescein succinimidyl ester (CFSE; Molecular Probes) at a concentration of 0.5 μM (CFSE¹⁰ cells). Splenocytes from infected mice were labeled with 5 μM CFSE (CFSE^{hi} cells) and preincubated, or not, with the 667 mAb (the absence of 667 mAb allowed quantification of spontaneous cell death). Both cell populations were mixed at a 1:1 ratio before retro-orbit-al administration to recipient mice. Cytolysis activity against infected splenocytes was calculated from the ratio of CFSE¹⁰/CFSE^{hi} cells in spleen assayed by flow cytometry 5 hours later. To assess the contribution of NK cells and neutrophils to antibody-mediated cytolysis, 50 μl of the anti-asialo GM1 or 200 μg of the anti-Ly6G 1A8 mAb were administered 1 day prior to the assay.

Flow cytometry assay of CD8⁺ *T cells specific for FrCasE-infected cells*. Splenocytes were labeled with both an APC-conjugated anti-CD8⁺ T cell antibody and a PE-conjugated MHC class I H-2D^b tetramer (Beckman Coulter) displaying the immunodominant Friend virus GagL epitope (27, 80) (D^b-GagL tetramers) as previously described (27).

Assay of IFN- γ production. 10⁶ splenocytes were incubated at 37°C for 5 hours in 12-well plates in 500 µl RPMI culture medium containing phorbol 12-myristate 13-acetate (50 ng/ml) and ionomycin (500 ng/ml) in the presence of brefeldin A (10 µg/ml; MilliporeSigma). IFN- γ production was assayed by flow cytometry using the intracellular Cytofix/Cytoperm Fixation/Permeabilization staining kit (Becton Dickinson).

Histological analyses. Spleens from infected/nontreated and infected/treated mice (depleted or not of neutrophils) were recovered at day 14 p.i. and prepared as previously described (81). Briefly, spleens were initially fixed at 4°C overnight with a paraformaldehyde-lysine-periodate (PLP) solution and then embedded in 4% low-temperature-gelling agarose (type VII-1; MilliporeSigma) prepared in PBS. 300-µm slices were cut with a vibratome (VT 1000S; Leica) in a bath of ice-cold PBS. Sections of tissues were submerged in PBS and transferred to 0.4-µm organotypic culture inserts (Millicell; Millipore) for staining with an anti-B220 (RA3-6B2, Becton Dickinson) and an anti-CD169 antibody (MOMA-1, Bio-Rad) at 37°C for 20 minutes. The images were captured using a Leica SP8-UV confocal scanning microscope.

RT-qPCR quantification of gene expression. Single-cell suspensions of splenocytes were prepared from naive, infected/nontreated, and infected/treated mice 8 days p.i. and immunotherapy. Neutrophils (CD11b⁺Ly6G^{hi} expression) were sorted (>98% pure) using a BD Biosciences FACSAria device. RNA was extracted from 1 \times 10⁶ to 2 \times 10⁶ sorted neutrophils using the RNeasy micro kit (Qiagen). RNA quality and integrity were verified using the Agilent 2100 bioanalyzer. cDNAs were synthesized using the RT2 First-Strand Kit (Qiagen). All quantitative PCRs were performed following protocols optimized for the RT2 quantitative Profiler PCR array using SYBR Green mix (Qiagen) and LightCycler 480 II machine (Roche). All data were normalized to β -actin. Results were expressed as fold increases with respect to naive cells using the $\Delta\Delta$ Ct method.

BAFF and LTα protein release quantification. Soluble BAFF and LTα from cell-free supernatants of cultured neutrophils were assayed using BAFF (R&D Systems) and LTα ELISA (NeoBiotech), respectively. Supernatants were collected from sorted splenic neutrophils (from naive, infected/nontreated, and infected/treated mice at 8 days p.i. and immunotherapy) cultured in 96-well plates at a density of 2×10^5 cells/ well for 24 hours. Alternatively, neutrophils were isolated from naive mice BM using a magnetic-based cell sorting (MACS) neutrophil isolation kit (>95% purity; Miltenyi Biotec) and cultured for 24 hours in 667 mAb–coated 24-well plates at a density of 2×10^6 cells in 500 µl medium, in the presence or in the absence of IFN-γ (100 ng/ml). 667 mAb–noncoated plates were used as controls. G-CSF (R&D Systems) was added at a concentration of 10 ng/ml to neutrophil cultures to maintain cell viability.

Statistics. Statistical analyses were performed using GraphPad Prism 5 (GraphPad Software). Data were expressed as mean \pm SEM, and statistical significance was established using a parametric 1-way ANOVA test with a Bonferroni correction for multiple comparisons or unpaired Student's 2-tailed *t* tests when two groups were compared. *P* values lower than 0.05 were considered as statistically significant.

Study approval. All experimental procedures were performed in accordance with the French national animal care guidelines (CEEA-LR-12146 approval, Ethics committee of the Languedoc-Roussillon Region, Montpellier).

Author contributions

M. Pelegrin, MNG, and M. Piechaczyk defined the research program. MNG, JL, and M. Pelegrin performed the experiments and carried out the data analyses, with a contribution by M. Piechaczyk. M. Pelegrin, MNG, and M. Piechaczyk wrote the manuscript.

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