

Recruitment of latent pools of high-avidity CD8⁺ T cells to the antitumor immune response

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A major barrier to successful antitumor vaccination is tolerance of high-avidity T cells specific to tumor antigens. In keeping with this notion, HER-2/neu (neu)-targeted vaccines, which raise strong CD8⁺ T cell responses to a dominant peptide (RNEU₄₂₀₋₄₂₉) in WT FVB/N mice and protect them from a neu-expressing tumor challenge, fail to do so in MMTV-neu (neu-N) transgenic mice. However, treatment of neu-N mice with vaccine and cyclophosphamide-containing chemotherapy resulted in tumor protection in a proportion of mice. This effect was specifically abrogated by the transfer of neu-N-derived CD4⁺CD25⁺ T cells. RNEU₄₂₀₋₄₂₉-specific CD8⁺ T cells were identified only in neu-N mice given vaccine and cyclophosphamide chemotherapy which rejected tumor challenge. Tetramer-binding studies demonstrated that cyclophosphamide pretreatment allowed the activation of high-avidity RNEU₄₂₀₋₄₂₉-specific CD8⁺ T cells comparable to those generated from vaccinated FVB/N mice. Cyclophosphamide seemed to inhibit regulatory T (T reg) cells by selectively depleting the cycling population of CD4⁺CD25⁺ T cells in neu-N mice. These findings demonstrate that neu-N mice possess latent pools of high-avidity neu-specific CD8⁺ T cells that can be recruited to produce an effective antitumor response if T reg cells are blocked or removed by using approaches such as administration of cyclophosphamide before vaccination.

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Abbreviations used: BrdU, bromodeoxyuridine; Dox, doxorubicin; ICS, intracellular cytokine stain; k_{off} , dissociation rate constant; neu, HER-2/neu; neu-N, MMTV-HER-2/neu transgenic mice; T reg, regulatory T.

Central and peripheral mechanisms of T cell tolerance must exist to prevent autoimmunity (1–3). However, these same mechanisms are likely contributors to ineffective T cell responses often observed in cancer patients (4). CD8⁺ T cells specific for tumor antigens have been detected in patients. In particular, HER-2/neu (neu)-specific T cells have been isolated from patients with breast and ovarian cancer (5). In addition, p53-specific T cells have been isolated from patients with colorectal carcinomas, human papilloma virus-specific T cells have been isolated from patients with cervical cancer, and T cells specific for more than 10 different antigens have been isolated from patients with melanoma (6). Although these T cell responses are often observed in patients who have been treated with an antigen-specific vaccine, most are weak and ineffective in controlling tumor growth. In some

instances, this ineffectiveness may result from an ineffective vaccine approach. However, because the majority of tumor-associated antigens are either overexpressed or reactivated developmental self-antigens, tolerance to specific tumor antigens is a likely mechanism for blunted T cell responsiveness in many cases (7, 8).

Intrathymic expression of self-antigens often leads to central deletion of T cells that express high-avidity TCR specific for these antigens. However, T cells that express lower-avidity TCR for these antigens can escape into the periphery (9). In addition, both high-avidity and low-avidity T cells specific for peripherally expressed self-antigens will leave the thymus rather than undergo deletion (10). Therefore, peripheral (extrathymic), nondeletional mechanisms exist that render T cells specific for self-tissues ignorant or functionally impaired (11–13). Understanding the mechanisms of peripheral tolerance in the context of tumor antigens is crit-

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ical for the development of interventions that can reverse the tolerant state and allow these T cells to respond more effectively to tumors. In particular, it is critical to determine whether there are latent pools of tumor-specific T cells in tumor-tolerant individuals that are capable of being coaxed into a functionally active state.

Several murine models are available for dissecting mechanisms of peripheral tolerance (14). Extending this approach to a relevant tumor antigen, we recently described the existence of immune tolerance in the HER-2/*neu* transgenic (*neu*-N) mouse model of breast cancer. These mice overexpress the rat protooncogene *neu* and develop spontaneous mammary tumors between 4 and 8 mo of age (15). Similar to observations in patients with breast and ovarian cancers (5), *neu*-N mice exhibit down-regulated immune responses after *neu*-targeted vaccination. Specifically, *neu*-N mice develop poor antibody and T cell responses to *neu* regardless of the *neu*-targeted vaccination approach used when compared with parental FVB/N mice receiving the same vaccine (16). The presence of the transgene does not result in complete central deletion of *neu*-specific T cells in *neu*-N mice, because anti-*neu* activity is detected. However, the possibility remains that subpopulations of high-avidity *neu*-specific T cells are de-

leted centrally, whereas T cells with lower avidity can leave the thymus but are subject to peripheral mechanisms of tolerance. Alternatively, tolerization of all *neu*-specific T cells may take place completely in the periphery. Various models of antigen-specific tolerance support all of these hypotheses.

We previously reported that treatment of *neu*-N mice with immune modulating doses of cyclophosphamide-containing chemotherapy before vaccination resulted in tumor protection in 10–30% of mice (17). Here we show that the vaccine-enhancing effect of cyclophosphamide is mediated through selectively inhibiting the cycling population of CD4⁺CD25⁺ regulatory T (T reg) cells in *neu*-N mice. Furthermore, high-avidity RNEU_{420–429}-specific CD8⁺ T cells are not deleted in the periphery, as previously thought. Instead, these high-avidity T cells can be recruited and activated to provide a potent antitumor immune response if T reg cells are inhibited.

RESULTS

Adoptive transfer of CD4⁺CD25⁺ regulatory T cells inhibits the antitumor immune response induced by cyclophosphamide given with vaccine

We previously demonstrated that 10–30% of tolerized *neu*-N mice can be cured of *neu*-expressing tumors when treated

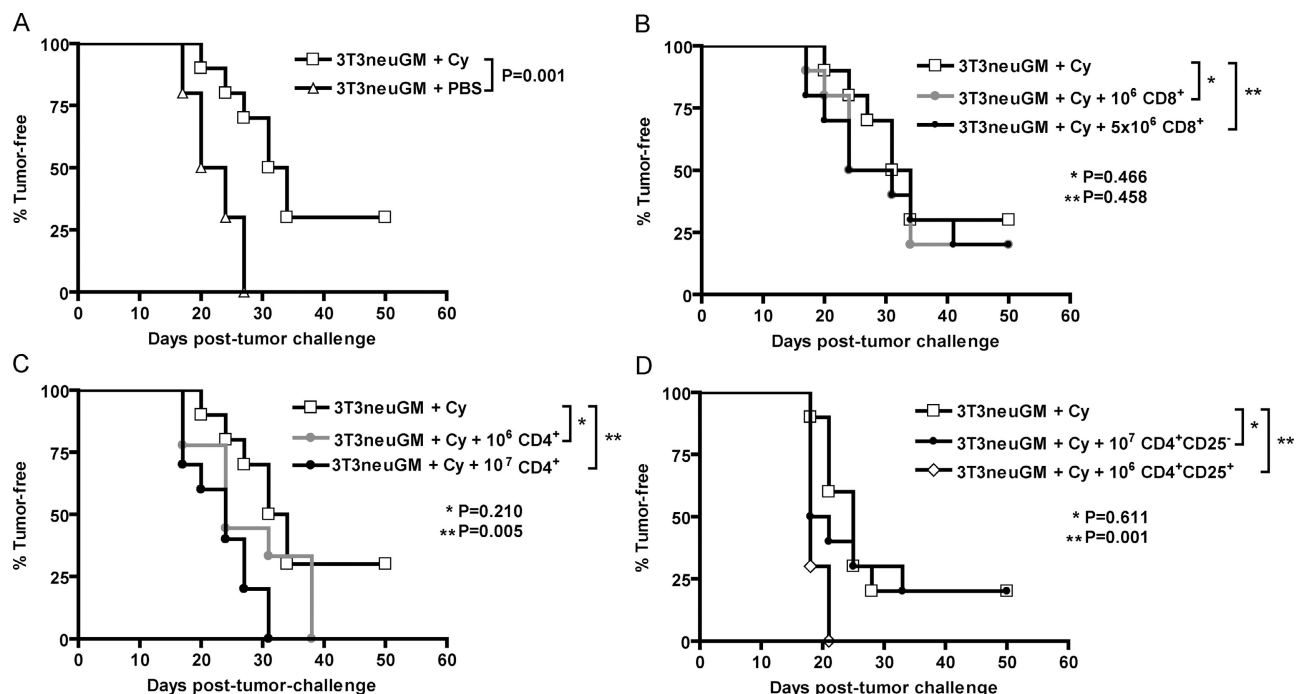


Figure 1. Adoptive transfer of CD4⁺CD25⁺ T reg cells abrogates the immune modulatory effect of cyclophosphamide. 10 *neu*-N mice/group were vaccinated with either 3T3neuGM alone on day 0 or with cyclophosphamide on day -1 and 3T3neuGM on day 0. Adoptively transferred T cells were given on day 13, as described in Materials and methods. On day 14, all mice were challenged with NT2 tumor cells. Mice were monitored for tumor outgrowth two times/wk. (A) cyclophosphamide given before vaccine results in significant protection ($P = 0.001$) from NT tumor challenge. Adoptively transferred, purified CD8⁺ splenocytes (B) were not

capable of suppressing the effect of cyclophosphamide, whereas CD4⁺ splenocytes (C) did significantly inhibit the immune modulatory effects of cyclophosphamide in a dose-dependent fashion ($P = 0.005$ for 10^7 CD4⁺). (D) The experiment was repeated using purified CD4⁺CD25⁻ splenocytes and purified CD4⁺CD25⁺ splenocytes. A significant suppressive effect of the T cells localized to the CD4⁺CD25⁺ splenocytes ($P = 0.001$), and not to the CD4⁺CD25⁻ splenocytes ($P = 0.611$). Each study was repeated at least twice with similar results. Cy, cyclophosphamide.

with immune-modulating doses of cyclophosphamide chemotherapy in sequence with neu-targeted vaccination. In contrast, all *neu-N* mice given vaccine alone develop tumors (17). T reg cells probably play a role in suppressing antigen-specific T cell responses (3). It has been suggested for more than 20 yr, and addressed more recently, that immune-modulating doses of cyclophosphamide may function by inhibiting suppressor T cell activity (18–21). We therefore evaluated T reg cells as a mechanism by which vaccine-induced immunity is suppressed in *neu-N* mice. *Neu-N* mice were given vaccine, alone or with cyclophosphamide, and then tumor challenged 2 wk later. Pretreatment with cyclophosphamide significantly enhanced the vaccine effect ($P = 0.001$) compared with mice given vaccine alone (Fig. 1 A). To determine if cyclophosphamide pretreatment eliminates a suppressive T cell population, *neu-N* mice that were given cyclophosphamide and vaccine received adoptively transferred CD8⁺ T cells (either 10^6 or 5×10^6 total per mouse; Fig. 1 B), or CD4⁺ T cells (either 10^6 or 10^7 total per mouse; Fig. 1 C) 1 d before tumor challenge. Initial experiments used T cells isolated from the spleens of tumor-bearing donor *neu-N* mice vaccinated 14 d before adoptive transfer. Because no enrichment of the transferred cells based on antigen specificity could be performed, we reasoned that any suppressive population would be activated in the vaccinated tumor-bearing donors. Neither dose of CD8⁺ T cells abrogated the antitumor response in the cyclophosphamide- and vaccine-treated mice (Fig. 1 B). However, CD4⁺ T cells from donor *neu-N* mice significantly suppressed the antitumor responses induced by cyclophosphamide and vaccine ($P = 0.005$) when 10^7 cells were transferred (Fig. 1 C). In subsequent experiments, T cells isolated from naive *neu-N* mice were used for the adoptive transfer, yielding the same results (unpublished data). These data suggest that there is a subset of CD4⁺ T cells with T reg cell function that inhibit the activity of tumor-rejecting, neu-specific T cells. In a second set of studies, mice received CD4⁺CD25⁻ T cells (10^7 total per mouse) or CD4⁺CD25⁺ T cells (10^6 total per mouse). The CD4⁺CD25⁻ T cell population failed to inhibit cyclophosphamide-modulated antitumor immunity ($P = 0.611$), whereas the CD4⁺CD25⁺ T cell population completely abrogated the antitumor immune response even at a 10-fold-lower dose ($P = 0.001$; Fig. 1 D). The suppressive activity of the CD4⁺CD25⁺ T cell subset was confirmed by in vitro proliferation assay in coculture with CD4⁺CD25⁻ T cells (22) and detection of *foxp3* expression by RT-PCR (reference 23 and unpublished data).

Vaccine combined with cyclophosphamide chemotherapy uncovers T cells specific for the immunodominant epitope RNEU₄₂₀₋₄₂₉, directly correlating RNEU₄₂₀₋₄₂₉ T cell activity with in vivo tumor rejection

In nontolerized FVB/N mice, neu-targeted vaccine will completely cure mice from a neu-expressing tumor chal-

lenge. In vivo rejection of tumor is associated with a neu-specific CD8⁺ T cell repertoire that is directed at the immunodominant T cell epitope RNEU₄₂₀₋₄₂₉ (24). We have shown that the RNEU₄₂₀₋₄₂₉-specific CD8⁺ T cells are capable of lysing neu-expressing tumors in vitro and eradicating neu-expressing tumors in FVB/N mice in vivo. However, activated RNEU₄₂₀₋₄₂₉-specific T cells are rarely detected in vaccinated *neu-N* mice (24). Because cyclophosphamide chemotherapy combined with vaccine cures 10–30% of *neu-N* mice, we hypothesized that the curative effect was caused by the cyclophosphamide chemotherapy regimen activating RNEU₄₂₀₋₄₂₉-specific T cells. Tumor-challenged *neu-N* mice were given vaccine with or without cyclophosphamide chemotherapy and followed for development of neu-express-

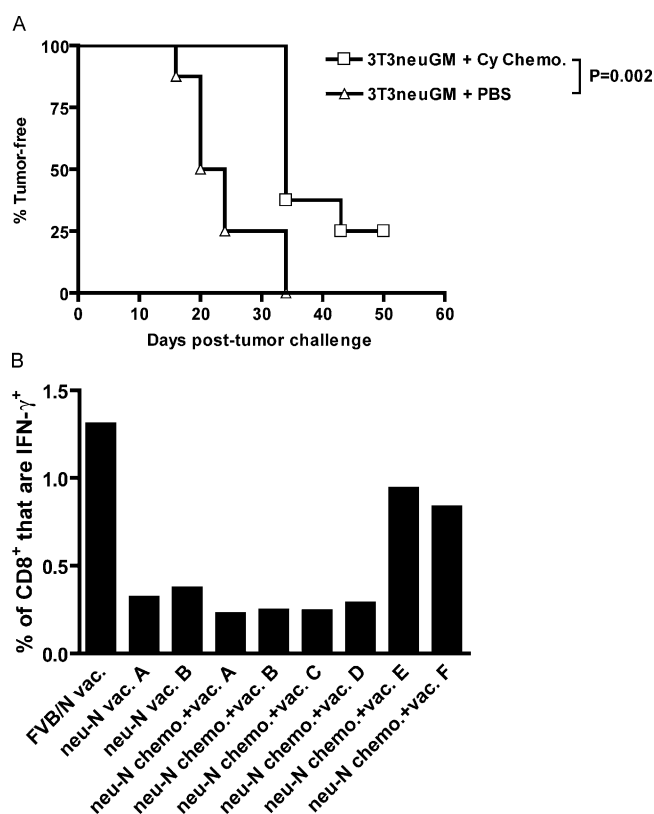


Figure 2. RNEU₄₂₀₋₄₂₉-specific T cells can be identified in polyclonal T cell populations from *neu-N* mice given vaccine and cyclophosphamide chemotherapy that rejected neu-expressing tumors. (A) Eight *neu-N* mice/group were tumor challenged on day -3, followed on day 0 by vaccination with or without cyclophosphamide chemotherapy. Mice were monitored for tumor outgrowth two times/wk. All mice that received mock vaccine with or without cyclophosphamide chemotherapy developed tumor by day 35 (not depicted). (B) At the end of the experiment in (A), splenic T cells were isolated, and reactivity to RNEU₄₂₀₋₄₂₉ was determined by ICS. Plotted is the percentage of CD8⁺ T cells that secreted IFN- γ in response to RNEU₄₂₀₋₄₂₉ minus the percentage responding to NP₁₁₈₋₁₂₆. This experiment was repeated at least three times with similar results. Cy, cyclophosphamide.

ing mammary tumors. As previously reported, cyclophosphamide chemotherapy resulted in a significant enhancement of the vaccine effect ($P = 0.002$), and 25% of mice receiving vaccine and cyclophosphamide chemotherapy remained tumor-free (Fig. 2 A; reference 17). These mice were killed on day 50, and intracellular cytokine staining (ICS) was performed to assess recognition of RNEU₄₂₀₋₄₂₉ in both tumor-bearing and tumor-free mice. As shown in Fig. 2 B, the two tumor-free *neu-N* mice (mouse E and mouse F) showed the highest percentage of RNEU₄₂₀₋₄₂₉-specific CD8⁺ T cells. The tumor-bearing mice (mice A–D) showed no activation above the background observed in the *neu-N* mice given vaccine alone. Thus, cyclophosphamide chemotherapy combined with vaccine seems to activate RNEU₄₂₀₋₄₂₉-specific T cells in the *neu-N* mice, enabling them to reject the *neu*-expressing tumor challenge. Similar activation of

RNEU₄₂₀₋₄₂₉-specific T cells was observed in tumor-free *neu-N* mice given cyclophosphamide alone with vaccine (unpublished data).

Because the presence of RNEU₄₂₀₋₄₂₉-specific T cells could result simply from a boosting effect of having successfully rejected the *neu*-expressing tumor, we determined if RNEU₄₂₀₋₄₂₉-specific T cells could be detected in *neu-N* mice given cyclophosphamide chemotherapy in the absence of a tumor challenge. RNEU₄₂₀₋₄₂₉-specific CD8⁺ T cell responses were detected in 20–30% of these mice (unpublished data), correlating well with our previous data measuring the antitumor effect of vaccine and cyclophosphamide chemotherapy in tumor-bearing *neu-N* mice (17). Thus, T cells specific for the immunodominant epitope, RNEU₄₂₀₋₄₂₉, can be detected after treatment with cyclophosphamide chemotherapy and vaccine in tumor naive *neu-N* mice.

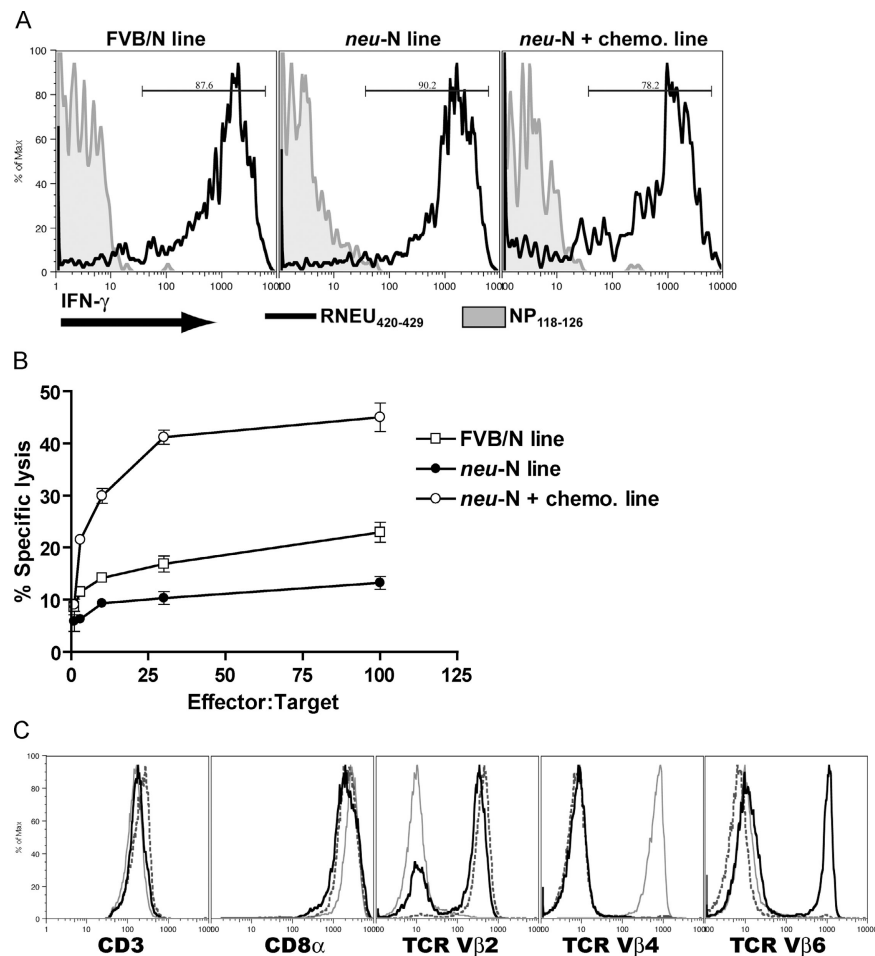


Figure 3. FVB/N and *neu-N*-derived CD8⁺ T cell lines are specific for RNEU₄₂₀₋₄₂₉ but differentially lyse *neu*-expressing mammary tumors despite expressing similar levels of cell surface markers. (A) T cell lines were derived from vaccinated FVB/N mice (FVB/N line), *neu-N* were mice given vaccine alone (*neu-N* line), or *neu-N* mice were given vaccine and cyclophosphamide chemotherapy that had rejected an NT tumor challenge (*neu-N* plus chemotherapy line). IFN- γ ICS was performed after T cells were

stimulated overnight with equal numbers of T2D⁹ cells pulsed with either the irrelevant peptide NP₁₁₈₋₁₂₆ (shaded histogram) or with RNEU₄₂₀₋₄₂₉ (black line). (B) CTL assay using the three T cell lines and *neu*-expressing NT2 tumor targets. These experiments were repeated more than six times with similar results. (C) Cells were stained with antibodies to CD3, CD8 α , TCR V β 2, TCR V β 4, and TCR V β 6 as described in Materials and methods. Black line, FVB/N line; dashed line, *neu-N* line; gray line, *neu-N* plus chemotherapy line.

Comparison of RNEU₄₂₀₋₄₂₉-specific CD8⁺ T cell lines from vaccinated FVB/N and *neu*-N mice reveal differences in T cell avidity that may explain the enhanced therapeutic effect of cyclophosphamide chemotherapy and vaccine in the *neu*-N mice

CD8⁺ T cell lines were generated after *neu*-targeted vaccination in FVB/N and *neu*-N mice. All lines were generated by repeated *in vitro* stimulation with 3T3neuB7-1 target cells that contain the entire *neu* cDNA. Initial analysis of T cell lines derived from the vaccinated FVB/N and *neu*-N mice, as well as *neu*-N mice given vaccine and cyclophosphamide chemotherapy, revealed that all are specific for the RNEU₄₂₀₋₄₂₉ peptide (Fig. 3 A). In contrast, these lines did not recognize the irrelevant H-2D^a-binding peptide NP₁₁₈₋₁₂₆ (25). To characterize functional differences between the FVB/N-derived and the *neu*-N-derived T cell lines, a more rigorous analysis directly comparing the three T cell lines was performed. Each T cell line was evaluated for the ability to lyse NT2 mammary tumor cells, which express naturally processed *neu* peptides. The RNEU₄₂₀₋₄₂₉-specific T cell line derived from vaccinated *neu*-N mice was less effective at lysing tumor cells than the lines derived from vaccinated FVB/N mice and from *neu*-N mice given vaccine and cyclophosphamide chemotherapy (Fig. 3 B). The T cell lines were all >99% CD8⁺ and expressed comparable surface levels of TCR, CD8 α , and CD3 (Fig. 3 C). Because these lines are comparable in the degree of peptide specificity and TCR, CD8 α , and CD3 levels, we hypothesized that the difference in tumor lysis could be caused by a difference in avidity of TCR for the H-2D^a/RNEU₄₂₀₋₄₂₉ MHC/peptide complex.

To address this issue, we stained the three T cell lines with decreasing concentrations of the H-2D^a-RNEU₄₂₀₋₄₂₉ tetramer. This study revealed that the T cells derived from *neu*-N mice given vaccine alone stained with a much lower intensity than the FVB/N-derived T cells. T cells derived from *neu*-N mice given vaccine and cyclophosphamide chemotherapy showed a staining profile similar to the FVB/N-derived line (Fig. 4 A). Even a 10-fold dilution of the tetramer resulted in loss of staining of the *neu*-N T cell line, whereas the other two lines showed staining even at a 100-fold dilution of the tetramer. Further quantitation revealed that at least 16 μ M of tetramer complex was required to stain the T cell line derived from vaccinated *neu*-N mice. In contrast, only 0.16 μ M of tetramer was required to demonstrate similar staining of the FVB/N-derived T cell line and of the T cell line derived from *neu*-N mice given vaccine and cyclophosphamide chemotherapy. Additionally, binding kinetics studies were performed by binding tetramer to T cell lines and competing it off with the H-2D^a-specific antibody 30-5-7S (26). As shown in Table I, the dissociation rate constant (k_{off}) for the *neu*-N line was roughly sixfold higher than that of the FVB/N line and the T cell line derived from *neu*-N mice given vaccine and cyclophosphamide chemotherapy. Other reports have indicated that the expression level of CD8 β influences functional avidity (27). How-

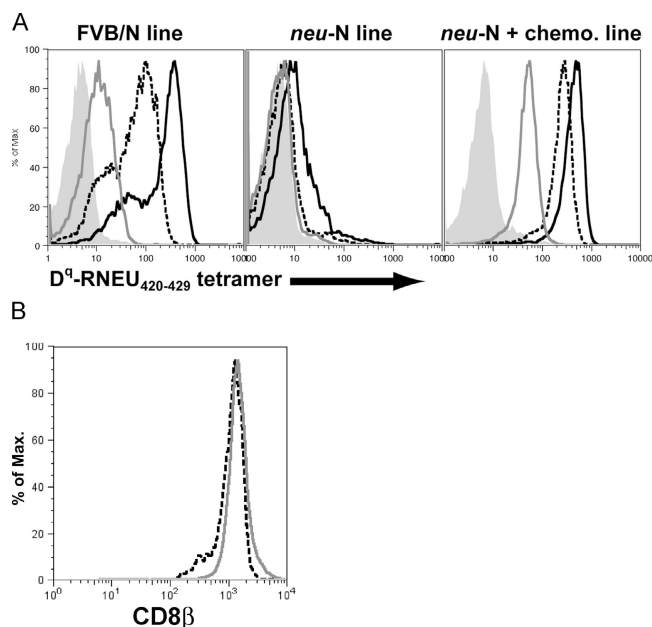


Figure 4. RNEU₄₂₀₋₄₂₉-specific CD8⁺ T cell lines derived from FVB/N and *neu*-N mice demonstrate quantitatively different avidities for the H-2D^a/RNEU₄₂₀₋₄₂₉ MHC/peptide complex but no difference in CD8 β staining. (A) The three T cell lines were stained with decreasing amounts of H-2D^a/RNEU₄₂₀₋₄₂₉ tetramer. Shown is the tetramer staining of gated CD8⁺ T cells. Black solid line, 1:5 tetramer dilution; dashed line, 1:50 tetramer dilution; gray line, 1:500 tetramer dilution; shaded, no tetramer. (B) T cell lines from *neu*-N mice given vaccine alone (dashed line) and *neu*-N mice given cyclophosphamide chemotherapy (gray line) were stained for CD8 β on day 6 after stimulation.

ever, in our studies, no difference in CD8 β staining was observed between the T cell lines from the *neu*-N mice given vaccine alone and the *neu*-N mice given vaccine and cyclophosphamide chemotherapy (Fig. 4 B). These data strongly support our hypothesis that the *neu*-N-derived T cell line has a lower avidity for the MHC class I/RNEU₄₂₀₋₄₂₉ epitope complex than the FVB/N-derived T cell line or the T cell line derived from vaccine and cyclophosphamide chemotherapy treated *neu*-N mice. Importantly, low avidity correlated with reduced lysis of the *neu*-expressing mammary tumor (Fig. 3 B).

High-avidity RNEU₄₂₀₋₄₂₉-specific T cells are detected in a polyclonal population of splenic T cells isolated from *neu*-N mice given vaccine and cyclophosphamide chemotherapy

The data analyzing T cell lines suggest that treatment with immune-modulatory doses of cyclophosphamide chemotherapy before vaccination results in the recruitment of functional high-avidity *neu*-specific T cells in up to 30% of *neu*-N mice. The studies analyzing splenic T cells from *neu*-N mice demonstrate that RNEU₄₂₀₋₄₂₉-specific T cells can be isolated from a polyclonal T cell response after vaccine combined with cyclophosphamide chemotherapy (Fig. 2 B). To demonstrate that these RNEU₄₂₀₋₄₂₉-specific T cells are high avidity, 42

Table I. k_{off} for RNEU₄₂₀₋₄₂₉-specific T cell lines^a

T cell line	k_{off} (min^{-1})	SD
FVB/N	41.5×10^{-3}	$\pm 10.1 \times 10^{-3}$
<i>neu</i> -N	272.7×10^{-3}	$\pm 164.1 \times 10^{-3}$
<i>neu</i> -N + chemotherapy	17.5×10^{-3}	$\pm 3.9 \times 10^{-3}$

^aAssay performed as described in Materials and methods. Values represent an average of three experiments per T cell line.

additional tumor-challenged *neu*-N mice were vaccinated in sequence with cyclophosphamide chemotherapy and monitored for tumor development. As expected, 20% of the mice remained tumor-free (unpublished data). On day 56, splenocytes were isolated from tumor-bearing and tumor-free mice, and reactivity to RNEU₄₂₀₋₄₂₉ was assessed by ICS after one in vitro stimulation (unpublished data). TCR avidity was determined by dilutional tetramer staining of splenocytes from vaccinated FVB/N mice, tumor-free *neu*-N mice given vaccine and cyclophosphamide chemotherapy, and tumor-bearing *neu*-N mice given vaccine alone (included as a negative control to show background tetramer staining because no reactivity to RNEU₄₂₀₋₄₂₉ was observed by ICS). RNEU₄₂₀₋₄₂₉-specific T cells from the vaccinated FVB/N mice and cyclophosphamide chemotherapy-treated *neu*-N mice exhibited equivalent tetramer staining, comparable to the tetramer staining observed in the high-avidity T cell lines (Fig. 5). These staining studies were performed by first gating on CD8⁺ T cells that are CD62L^{lo}, demonstrating that the tetramer-positive RNEU₄₂₀₋₄₂₉-specific T cells are activated. We therefore conclude that the CD8⁺ T cell responses to neu in parental FVB/N mice, but not in tolerized *neu*-N mice, are dominated by high-avidity T cells specific for RNEU₄₂₀₋₄₂₉. In addition, immune-modulating doses of cyclophosphamide chemotherapy given in sequence with neu-targeted vaccination can uncover activated, high-avidity T cells specific for the immunodominant neu epitope in *neu*-N tolerized mice.

Cyclophosphamide exerts its effect predominantly on cycling CD4⁺CD25⁺ T cells

To understand the mechanism of action of cyclophosphamide on CD4⁺CD25⁺ T cells better, the total number of CD4⁺CD25⁺ T cells was monitored in cyclophosphamide-treated naive *neu*-N mice. A significant decrease ($P \leq 0.0001$) in the total number of CD4⁺CD25⁺ T cells was observed in the LN 2 d after cyclophosphamide treatment (Fig. 6 A). This decrease was detected in the LNs at the time when T cell induction would be occurring in vaccinated mice. The percentage of CD4⁺ T cells that are CD25⁺ was also monitored for 2 wk in cyclophosphamide-treated *neu*-N mice. A significant drop ($P = 0.0007$) in the percent of CD4⁺CD25⁺ was also seen within 2 d after cyclophosphamide treatment, which then recovered over the next 2 wk (Fig. 6 B).

Because cyclophosphamide has a relatively modest effect on total CD4⁺CD25⁺ T cell numbers, and the population of

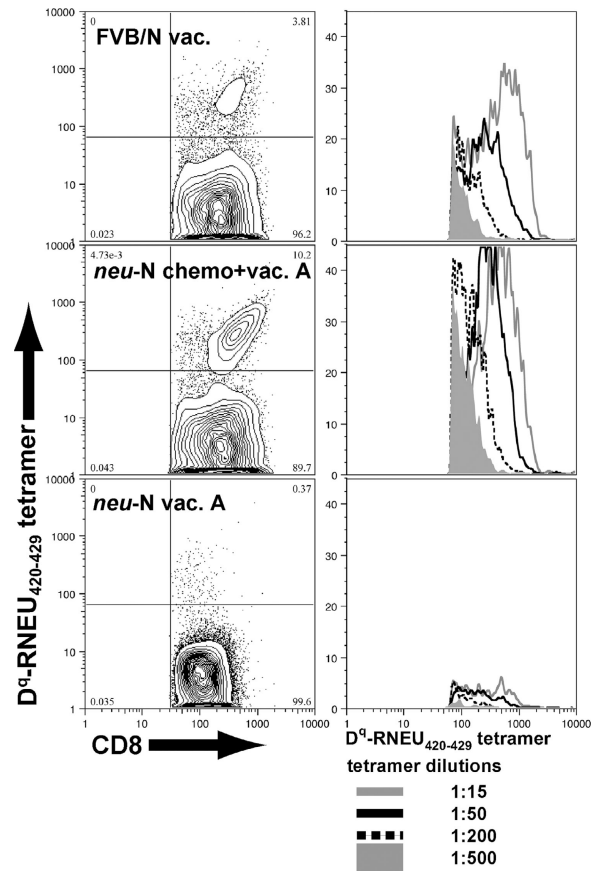


Figure 5. High-avidity, RNEU₄₂₀₋₄₂₉-specific T cells can be identified in polyclonal T cell populations from *neu*-N mice given vaccine and cyclophosphamide chemotherapy that rejected neu-expressing tumors. CD8⁺ enriched splenocytes were stained with decreasing amounts of H-2D^q-RNEU₄₂₀₋₄₂₉ tetramer. Three representative samples are shown. Plots on the left are gated on CD8⁺, CD62L^{lo} lymphocytes and show the 1:50 dilution of tetramer. Corresponding plots on the right show histograms gating on the CD8⁺, tetramer⁺ cells stained with decreasing amounts of tetramer.

CD4⁺CD25⁺ T cells in whole LNs is polyclonal, it is possible that cyclophosphamide affects a specific subset of CD4⁺CD25⁺ T cells. Others have shown that CD4⁺CD25⁺ T cells proliferate in the steady state in response to self-antigen (28). Thus, cyclophosphamide may act on the CD4⁺CD25⁺ T reg cell population that is proliferating to self-antigen in our model. To test this possibility, *neu*-N mice were given cyclophosphamide or PBS on day 0 and then pulsed with a 2-mg dose of bromodeoxyuridine (BrdU) on day 1. On day 2, LNs were harvested, and lymphocytes were stained for CD4, CD8, CD25, and BrdU. In the groups injected with PBS and BrdU, <1% of CD8⁺ T cells and <0.3% of CD4⁺CD25⁺ T cells incorporated BrdU. In contrast, 6.8% of CD4⁺CD25⁺ T cells incorporated BrdU. When treated with cyclophosphamide, the percent of CD4⁺CD25⁺ T cells that incorporated BrdU was not above background (Fig. 6 C). Thus, cyclophosphamide seems to eliminate ac-

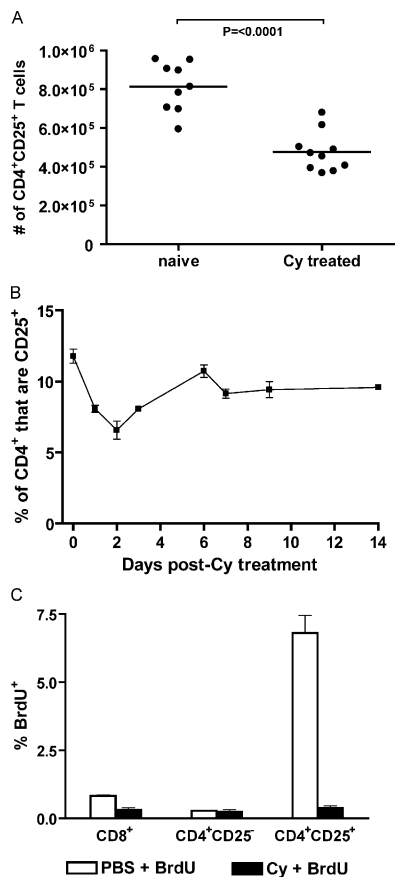


Figure 6. Cyclophosphamide depletes cycling CD4⁺CD25⁺ T cells. (A) Axillary LNs from PBS- or cyclophosphamide-treated *neu-N* mice (10/group) were isolated 48 h after cyclophosphamide administration. Total cell number in each LN was determined, and the number of CD4⁺CD25⁺ T cells was calculated based on the percent of CD4⁺CD25⁺ T cells determined by flow cytometry. (B) LN cells from *neu-N* mice (4/group) given cyclophosphamide were isolated and analyzed for the number of CD4⁺CD25⁺ T cells on the indicated days. The experiment was repeated three times. (C) *Neu-N* mice were given cyclophosphamide or PBS on day 0. On day 1, the mice were given a 2-mg dose of BrdU. On day 2, LNs were harvested, and the lymphocytes were stained for CD4, CD8, CD25, and BrdU. This experiment was repeated three times with similar results. Cy, cyclophosphamide.

tively cycling CD4⁺CD25⁺ T cells. These data provide strong evidence that cyclophosphamide administration depletes cycling T reg cells, allowing the generation and activation of high-avidity RNEU₄₂₀₋₄₂₉-specific T cells.

Direct depletion of CD4⁺CD25⁺ T cells with anti-CD25 antibody allows the detection of high-avidity RNEU₄₂₀₋₄₂₉-specific T cells

Based on these studies, cyclophosphamide seems to act on T reg cells that suppress high-avidity RNEU₄₂₀₋₄₂₉-specific T cell activity in *neu-N* mice. However, it is also possible that cyclophosphamide enhances high-avidity RNEU₄₂₀₋₄₂₉-specific CD8⁺ T cells through a T reg cell-independent mechanism. To address this issue, T reg cells were targeted di-

rectly with the anti-CD25-depleting antibody, PC61, given 4 d before vaccination (29). We first confirmed that a single 1.0-mg dose of PC61 decreased the total number of CD4⁺CD25⁺ T cells from 12% of total CD4⁺ T cells to less than 3% in *neu-N* mice (Fig. 7 A). Importantly, direct depletion of CD4⁺CD25⁺ T cells with PC61 resulted in the ability to isolate RNEU₄₂₀₋₄₂₉-specific T cells from 10–20% of mice (Fig. 7 B). This finding correlated with enhanced vaccine-induced prevention of tumor progression that was similar to observations with cyclophosphamide-modulated vaccine therapy (unpublished data). Furthermore, RNEU₄₂₀₋₄₂₉-specific T cells isolated from mice depleted of CD4⁺CD25⁺ T cells with PC61 before vaccination were high avidity, because they could be detected with multiple dilutions of the H-2D^a-RNEU₄₂₀₋₄₂₉ tetramer (Fig. 7 C).

CD4⁺CD25⁺ T cells from *neu-N* mice suppress vaccine-induced RNEU₄₂₀₋₄₂₉-specific CD8⁺ T cells in vaccinated, nontolerized FVB/N mice

Additional studies were performed to confirm further that CD4⁺CD25⁺ T reg cells directly suppress RNEU₄₂₀₋₄₂₉-specific CD8⁺ T cell responses. Specifically, groups of parental nontolerized FVB/N mice were vaccinated either alone or 1 d after receiving 5×10^5 adoptively transferred CD4⁺CD25⁺ T cells. Because the vaccine is extremely immunogenic in the nontolerant FVB/N mice, the CD4⁺CD25⁺ T cells were isolated from tumor-bearing, vaccinated *neu-N* mice with the goal of transferring an activated T reg cell population. After 2 wk, mice receiving vaccine alone demonstrated RNEU₄₂₀₋₄₂₉-specific CD8⁺ T cells ranging from 1–3%. However, mice receiving adoptively transferred CD4⁺CD25⁺ T cells demonstrated a significant ($P = 0.016$) decrease in RNEU₄₂₀₋₄₂₉-specific CD8⁺ T cell responses ranging from 0.25–1% (Fig. 8). These data provide additional evidence that CD4⁺CD25⁺ T reg cells have a direct suppressing effect on RNEU₄₂₀₋₄₂₉-specific CD8⁺ T cell responses. Despite the decrease in the percent of RNEU₄₂₀₋₄₂₉-specific T cells, the T cells that were activated seemed to be functional, because no suppression of the antitumor response was observed when the FVB/N mice that received the T reg cell transfer were given a *neu*-expressing tumor challenge (unpublished data).

DISCUSSION

We previously identified RNEU₄₂₀₋₄₂₉ as the immunodominant epitope encoded by the rat *neu* gene and recognized by the majority of T cell lines and clones derived from vaccinated FVB/N mice. We now report three new findings that give insight into the mechanisms of CD8⁺ T cell tolerance in *neu-N* transgenic mice. First, treatment of *neu-N* mice with immunomodulating doses of chemotherapy in sequence with *neu*-targeted vaccination uncovers high-avidity RNEU₄₂₀₋₄₂₉-specific CD8⁺ T cell activity that is associated with more effective eradication of *neu*-expressing tumors in vivo. Second, one mechanism by which cyclophosphamide

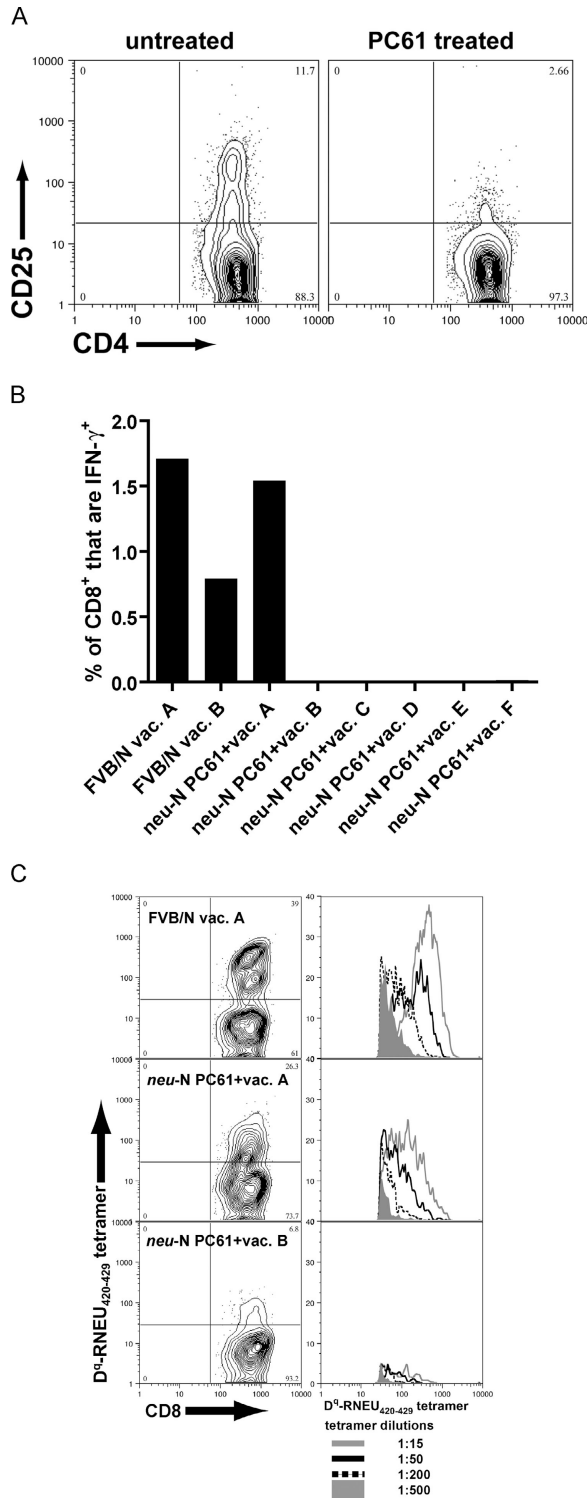


Figure 7. High-avidity RNEU₄₂₀₋₄₂₉-specific CD8⁺ T cells are detected after depletion of CD4⁺CD25⁺ T reg cells with the CD25⁺ T cell-depleting antibody, PC61. (A) Depletion was obtained using a single 1-mg dose of PC61 given 4 d before analysis. Splenocytes were isolated and stained with the noncompeting anti-CD25 antibody 7D4. **(B)** Mice were given 1 mg of PC61 4 d before vaccination. 2 wk after vaccination, splenocytes were isolated, and ICS was performed. Shown are the percent of CD8⁺ T cells that were RNEU₄₂₀₋₄₂₉-specific. Similar results have

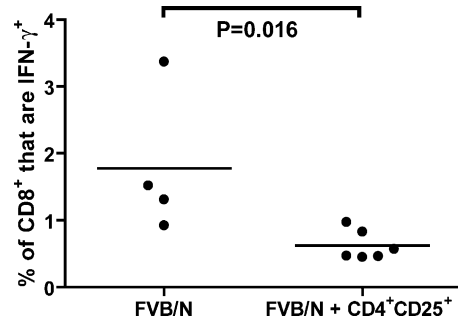


Figure 8. Adoptive transfer of CD4⁺CD25⁺ T cells from vaccinated, tumor-bearing *neu-N* mice suppresses the activation of RNEU₄₂₀₋₄₂₉-specific CD8⁺ T cells in FVB/N mice. CD4⁺CD25⁺ T cells from vaccinated, tumor-bearing *neu-N* mice were isolated by cell sorting. 5 × 10⁵ CD4⁺CD25⁺ T reg cells were transferred into naive FVB/N mice that were vaccinated the following day. On day 14 after vaccination, splenic CD8⁺ T cells were isolated, and ICS was performed. Shown are the percent of total CD8⁺ T cells that produced IFN- γ in response to RNEU₄₂₀₋₄₂₉. This study was repeated twice with similar results. ●, individual mice; solid line, average.

chemotherapy enhances vaccine-induced RNEU₄₂₀₋₄₂₉-specific T cells is through deletion of cycling CD4⁺CD25⁺ T reg cells. Third, unlike FVB/N mice, *neu*-targeted vaccine given alone to *neu-N* mice induces lower-avidity RNEU₄₂₀₋₄₂₉-specific T cell responses.

To our knowledge, this is the first report demonstrating the unmasking of high-avidity CD8⁺ T cell responses against a naturally expressed tissue-specific tumor antigen in a murine model of tolerance. Others have reported the induction of low-avidity antigen-specific T cells upon vaccination of transgenic mice for other model tumor antigens as compared with nontransgenic mice (8, 30, 31). In some cases, T cell avidity can be improved by repeated in vitro stimulation of T cells (11) or repeated antigen exposure in vivo (32). However, the initial generation of high-avidity T cells after in vivo vaccination regimens has not been previously demonstrated.

In several models of tumor tolerance, low-avidity T cells induced to recognize self-antigens were sufficient to suppress growth of tumors expressing the antigen (8). However, most other studies have shown that high-avidity T cells are superior antigen-recognition and lytic agents than their low-avidity counterparts (33–36). In one study, low-avidity CTL generated against the melanoma antigen gp100 were capable of lysing only peptide-pulsed cells, whereas high-avidity CTL efficiently lysed endogenously expressed levels of peptide presented by gp100-expressing tumor cells (37). Similarly, high-avidity CTL isolated from B16–GM-CSF-vaccinated mice showed superior in vitro and in vivo antitumor

been seen in more than six independent experiments. **(C)** Splenocytes from **(B)** were stimulated for 1 wk with RNEU₄₂₀₋₄₂₉ and then stained with decreasing amounts of H-2D^q/RNEU₄₂₀₋₄₂₉ tetramer. Three representative samples are shown. Plots on the left are gated on CD8⁺, CD62L^{lo} lymphocytes and show the 1:50 dilution of tetramer. Corresponding plots on the right show histograms gating on the CD8⁺, tetramer⁺ cells stained with decreasing amounts of tetramer.

activity when compared with lower-avidity CTL (38). In our mammary tumor model, the development of high-avidity T cells is associated with protection against tumor outgrowth and an apparent abatement of the tolerance exhibited in *neu*-N mice. We have shown that high-avidity *neu*-specific T cells can be recovered in 10–30% of *neu*-N mice treated with immune-modulating doses of chemotherapy. In addition to the T cell line shown in this paper, we have isolated other high-avidity RNEU_{420–429}-specific T cell lines from *neu*-N mice that have eradicated established transplanted mammary tumors after vaccination and cyclophosphamide chemotherapy. It is not possible, however, to isolate high-avidity RNEU_{420–429}-specific T cells from tumor-bearing *neu*-N mice given vaccine alone. These data provide additional evidence that optimal antitumor immunization will depend in part on the ability to induce high-avidity T cells specific for immunodominant epitopes contained in tumor antigens.

Our data also illustrate the importance of using the optimal vaccine strategy when targeting tumor antigens. We show that unless vaccine is combined with deletion or inhibition of T reg cells, the most potent CD8⁺ T cells will not be recruited to the antitumor immune response. Similar findings have been reported recently by Antony et al. (39) showing in an adoptive transfer model that optimal vaccine against melanoma antigens could be achieved only when T reg cells were removed. Other recent reports suggest that the vaccine itself can overcome the influence of T reg cells if signaling through Toll-like receptors is involved (e.g., a vaccinia-based vaccine) (40). However, our experience in the *neu*-N mice suggests otherwise. Our early reports studying the *neu*-specific immune responses in the *neu*-N mice showed no difference between our GM-CSF-secreting vaccine and a *neu*-expressing vaccinia vaccine (16). Furthermore, the high potency of our GM-CSF vaccine approach is supported by our earlier report that this vaccine given as a single agent results in the regression of 40-mm² tumors in the FVB/N nontolerized mice (17).

The studies presented here focused on the mechanism of cyclophosphamide in potentiating the immune response in vaccinated *neu*-N mice. We are currently investigating the mechanism of doxorubicin (Dox) in the combined chemotherapy regimen. However, our previous data do not support a role for Dox in eliminating T reg cell activity, because Dox inhibits vaccine-induced immune responses when given at the time of T cell priming (17). Dox enhances the effect of vaccine only when given at the time of T cell expansion, probably by enhancing the T cell's cytolytic activity (41). Thus, the ability to uncover high-avidity RNEU_{420–429}-specific CD8⁺ T cells in the *neu*-N mice is not dependent on Dox, because similar results were seen in mice given cyclophosphamide and Dox, cyclophosphamide alone, or PC61 in combination with vaccine.

In this study avidity is defined in part as the degree to which a T cell binds MHC/peptide tetramer. Although nu-

merous studies have shown a correlation between tetramer binding and T cell function (36, 42, 43), others have not (34, 43, 44). Here, the T cell lines that showed high-intensity tetramer staining were able to lyse *neu*-expressing tumor cells to a greater degree than the T cell line that displayed poor tetramer binding. Many investigations have also measured avidity of T cells by their ability to lyse peptide-pulsed cells (7, 8, 30, 37, 45). We confirmed these differences in T cell avidity for the H-2D^a/RNEU_{420–429} MHC/peptide complex by determining TCR k_{off} rates for each T cell line. The T cell line that showed the lowest amount of surface tetramer binding had the fastest k_{off} rate of the three lines analyzed. To the best of our knowledge, this is the first time the TCR k_{off} rate has been quantitated for tolerized versus nontolerized T cells in a murine tumor model.

It is still unexplained why adding cyclophosphamide chemotherapy to the vaccine regimen consistently overcomes tolerance in, at most, 30% of treated mice. Also, each mouse seems to respond in an all-or-none fashion. It is possible that improving the success of cyclophosphamide modulation depends on the long-term depletion or inhibition of T reg cell function. We have found that the T reg cell population in *neu*-N mice recovers within 2 wk of cyclophosphamide treatment. The kinetics is similar to T reg cell recovery after depletion with the CD25-targeted antibody, PC61 (29). It is more difficult to study recurrent treatment with cyclophosphamide, because lymphopenia can abrogate the vaccine-induced antigen-specific T cell response. CD25-targeted depleting antibodies are also difficult to give after an initial dose, because they also target vaccine-induced, activated T cells that up-regulate CD25. Furthermore, other systemic and local mechanisms within the tumor's microenvironment are probably active and require modulation to allow the long-term tumor trafficking and survival of high-avidity antigen-specific T cells. Studies are underway to elucidate additional mechanisms of peripheral tolerance in the *neu*-N mice.

Based on the data described in this report, we propose the following as a model to explain one mechanism of CD8⁺ T cell tolerance in *neu*-N mice. CTL from vaccinated *neu*-N mice that are specific for the immunodominant epitope are rare and are of lower avidity than T cells derived from parental FVB/N mice. These low-avidity T cells are also either weak lytic agents or anergized T cells. However, higher-avidity T cells (similar in avidity to those derived from nontolerant FVB/N mice) specific for the immunodominant epitope can escape thymic deletion and are actively suppressed in the periphery of these mice. Our data strongly support CD4⁺ CD25⁺ T reg cell involvement as one mechanism for the induction of peripheral tolerance. Central tolerance is unlikely to be a major mechanism, because high-avidity CD8⁺ T cells specific for RNEU_{420–429} can be recovered from vaccinated *neu*-N mice treated with T reg cell-depleting agents such as cyclophosphamide or PC61. It is possible that central tolerance does play a role because of uptake of *neu* antigen in the

periphery by immature dendritic cells when mature mice begin to overexpress *neu* in the periphery. Without the proper maturation signals, the outcome is likely to be cross-tolerance of CTL rather than cross-priming (46). However, our data demonstrate that *neu*-N mice develop CD4⁺CD25⁺ T reg cells that actively suppress RNEU₄₂₀₋₄₂₉-specific T cells (47). This is probably one mechanism of peripheral tolerance at work in our model, because treatment with cyclophosphamide or PC61, which have been reported to delete or inhibit T reg cells, reverses tolerance when combined with vaccine (18–21). This possibility is further supported by our data showing that cyclophosphamide selectively inhibits cycling CD4⁺CD25⁺ T reg cells in *neu*-N mice.

In summary, we have described one mechanism of CD8⁺ T cell tolerance to the protooncogene *neu* expressed in *neu*-N transgenic mice. Further investigation is needed to determine the additional mechanisms involved in the induction of *neu*-directed immune tolerance. However, our data strongly suggest that the most effective antitumor vaccine regimens must include a T reg cell-targeting agent to allow the most potent tumor-reactive T cells to be activated.

MATERIALS AND METHODS

Mice. All mice used were between 8 and 12 wk of age. FVB/N mice were purchased from the National Cancer Institute and Taconic Laboratories. *neu*-N mice (15), provided by William Muller, were bred to homozygosity as verified by Southern blot analysis (16) and bred and housed at Johns Hopkins University. All experiments were performed in accordance with protocols approved by the Animal Care and Use Committee of the Johns Hopkins University School of Medicine.

Cell lines and media. The GM-CSF-secreting vaccine cell lines 3T3GM and 3T3neuGM (expressing *neu*) were generated and grown as previously described (24). The IT22, IT22neu, and T2D^q cell lines used for targets in T cell assays were also previously described (24). The NT2 and NT5 *neu*-expressing tumor lines were derived from spontaneously arising mammary tumors excised from *neu*-N mice. NT2 cells used in tumor-challenge experiments express stable *neu* and MHC class I. NT5 was retrovirally infected with the human *B7-1* gene to generate NT5B7-1 cells as previously described (16, 17).

T cell lines were generated from FVB/N or *neu*-N mice given vaccine (3T3neuGM) or from *neu*-N mice that rejected an NT2 tumor challenge after being given vaccine and cyclophosphamide chemotherapy (see Immunization protocols). Splenocytes were initially stimulated every 5 d with irradiated, IFN- γ treated NT5B7-1 cells and then every 9 d by addition of irradiated 3T3neuB7-1 cells and FVB/N-derived splenocytes. T cells were maintained at 37°C and 5% CO₂ in CTL media (RPMI 1640 supplemented with 10% FBS, 1% L-glutamine, 0.1 mM 2-mercaptoethanol (Sigma-Aldrich), and 0.5% penicillin/streptomycin supplemented with 10 μ g/ml murine IL-2, a supernatant from B16 IL-2 line (17).

Peptides. RNEU₄₂₀₋₄₂₉ (PDSLRDLSVF) and NP₁₁₈₋₁₂₆ (RPQASGVYM) peptides were synthesized at >95% purity and purchased from either the Johns Hopkins Biosynthesis and Sequence Facility or from Macromolecular Resources of Colorado State University. The NP₁₁₈₋₁₂₆ peptide is from the lymphocytic choriomeningitis virus nucleoprotein (25) and is used as an irrelevant H2-D^a-binding peptide.

Antibodies and tetramer and flow cytometric analysis. mAb hybridoma supernatants (all acquired from the American Type Culture Collection) were used for staining T cell lines: B20.6 (TCR V β 2), KT4 (TCR V β 4),

and RR4-7 (TCR V β 6). Directly conjugated antibodies anti-CD3 FITC, anti-CD4 CyChrome, anti-CD8 α CyChrome, anti-CD8 β FITC, anti-CD25 PE (PC61 and 7D4), anti-CD62L allophycocyanin, and anti-IFN- γ PE were purchased from BD Biosciences. The CD25-depleting antibody PC61 was purified from the supernatant of PC61.5.3 hybridoma (American Type Culture Collection) grown in Protein Free Hybridoma Media II (Invitrogen).

The H-2D^a/RNEU₄₂₀₋₄₂₉ tetramer was constructed using previously described methods (48). Tetramer staining was performed by first staining only for CD8 (T cell lines) or CD8 and CD62L (ex vivo experiments). Cells were washed, and tetramer was added at varying dilutions and incubated for 30 min at 8–12°C. Cells were washed twice and immediately fixed with fresh 1% paraformaldehyde in PBS. Tetramer staining was analyzed by gating only on CD8⁺ cells (T cell lines) or CD8⁺, CD62L^{lo} cells (ex vivo experiments).

ICS was performed using the mouse ICS kit from BD Biosciences for murine IFN- γ . Purified T cells were incubated for 6–12 h with an equal ratio of indicated targets (IT22neu, IT22, or T2D^q cells pulsed with either RNEU₄₂₀₋₄₂₉ or NP₁₁₈₋₁₂₆) in the presence of GolgiStop (BD Biosciences). Cells were then stained for CD8, fixed and permeabilized, and stained with anti-IFN- γ PE. Analysis was performed by gating on CD8⁺ cells and calculating the percent of total CD8⁺ cells that were also IFN- γ ⁺. The percent of antigen-specific cells was calculated by subtracting the percent of IFN- γ ⁺ cells in the irrelevant antigen sample from the percent of IFN- γ ⁺ cells in the relevant antigen sample.

Flow cytometric data were collected using BD FACScan and BD FACSCalibur cytometers (BD Biosciences). Data were analyzed using CELLQuest (BD Biosciences) and FlowJo software (Tree Star, Inc.).

Immunization protocols. Mice were given vaccine (3T3neuGM), or mock vaccine (3T3GM cells) alone or in combination with cyclophosphamide, or cyclophosphamide plus Dox as described previously (17), or PC61 depletion. Mice given vaccine alone were injected s.c. with 3×10^6 total cells divided equally among one forelimb and two hind limbs. Mice given vaccine and cyclophosphamide received a single i.p. injection of cyclophosphamide, 100 mg/kg in 0.5 ml (Mead Johnson), on day -1 and vaccine (as described previously) on day 0. Vaccine given with cyclophosphamide and Dox included cyclophosphamide on day -1, vaccine on day 0, and Dox, 5 mg/kg given i.v. in 0.5 ml (Gensia Sico Pharmaceuticals Inc.) on day 7 (this regimen is referred to as cyclophosphamide chemotherapy in the text). We and others previously found that Dox given at the time of T cell expansion can enhance the cytotoxic function of CD8⁺ T cells (17, 41). PC61-depletion studies were performed by giving one dose of PC61 (1 mg i.p.) on day -4 followed by vaccine on day 0. In tumor-challenge experiments, NT2 tumor cells (5×10^4 cells injected s.c. in the mammary fat pad) were injected either on day -3 or on day 14. Mice were monitored for tumor outgrowth twice per week.

Adoptive transfer experiments. *neu*-N donor mice were either naive or given an NT2 tumor challenge on day -3 and 3T3neuGM vaccine on day 0. 2 wk after vaccination, donor CD4⁺ and CD8⁺ splenic T cells were isolated using MACS negative isolation kits (Miltenyi Biotec). Donor CD4⁺CD25⁺ and CD4⁺CD25⁻ splenic subsets were isolated from purified CD4⁺ T cells by staining for CD25 and sorting the positive and negative fractions using a FACSVantage cell sorter (Becton Dickinson). All T cell subsets were analyzed by flow cytometry to confirm purity. T cells were washed with PBS and injected i.v. into recipient mice.

Bromodeoxyuridine incorporation assay. *neu*-N mice were given cyclophosphamide or PBS i.p. on day 0. On day 1, mice were given a 2-mg dose of BrdU injected i.p. On day 2, axillary LN cells were isolated, washed, and stained using the FITC BrdU Flow Kit (BD Biosciences).

Chromium-release assays. Lysis assays were performed in triplicate in 96-well V-bottom plates as previously described (16). After 4-h incubation,

supernatant was assayed for ^{51}Cr release and the percentage of specific lysis was determined by the formula: (^{51}Cr release sample - spontaneous ^{51}Cr release)/(maximum ^{51}Cr release - spontaneous ^{51}Cr release) \times 100.

Tetramer off-rate experiments. T cells were washed and resuspended at 10^6 in 100- μl buffer. Cells were incubated with 8×10^{-3} M H-2D^a/RNEU₄₂₀₋₄₂₉ tetramer for 1 h at 4°C and then 20-fold molar excess of the H-2D^a antibody 30-5-7S (American Type Culture Collection) was added to prevent tetramer from rebinding T cells. Mean fluorescence intensity of tetramer staining was determined by flow cytometry at various time points from 0 to 240 min. Resulting data were fit to a first-order exponential decay using the Origin program (Microcal Software, Inc.). Off-rates were calculated as the reciprocal of t_1 .

Statistical analysis. Statistical significance of the tumor-free survival plots was determined using the log rank test. A Student's *t* test was applied to compare statistical significance between treatment groups. All analysis was performed using Prism 4 software (GraphPad Software).

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