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Unlicensed NK cells target neuroblastoma following anti-GD2 antibody treatment

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Survival outcomes for patients with high-risk neuroblastoma (NB) have significantly improved with anti-disialoganglioside GD2 mAb therapy, which promotes NK cell activation through antibody-dependent cell-mediated cytotoxicity. NK cell activation requires an interaction between inhibitory killer cell immunoglobulin-like receptors (KIRs) and HLA class I ligands. NK cells lacking KIRs that are specific for self HLA are therefore “unlicensed” and hyporesponsive. mAb-treated NB patients lacking HLA class I ligands for their inhibitory KIRs have significantly higher survival rates, suggesting that NK cells expressing KIRs for non-self HLA are mediating tumor control in these individuals. We found that, in the presence of mAb, both licensed and unlicensed NK cells are highly activated in vitro. However, HLA class I expression on NB cell lines selectively inhibited licensed NK cell activity, permitting primarily unlicensed NK cells to mediate antibody-dependent cell-mediated cytotoxicity. These results indicate that unlicensed NK cells play a key antitumor role in patients undergoing mAb therapy via antibody-dependent cell-mediated cytotoxicity, thus explaining the potent “missing KIR ligand” benefit in patients with NB.

Introduction

Neuroblastoma (NB), an embryonal malignancy of neuroectodermal origin, is the most common extracranial solid tumor of childhood. Nearly two-thirds of patients present at diagnosis with evidence of metastatic disease and have poor long-term survival due to residual disease, despite aggressive approaches, including high-dose multiagent chemotherapy, surgery, radiation therapy, and autologous stem cell transplantation (ASCT) (1, 2). Treatment of patients with high-risk NB with monoclonal antibodies (mAb) targeting the disialoganglioside surface antigen GD2 has resulted in lower recurrence rates and improved overall survival (OS) (3–5). In addition to complement-dependent cytotoxicity, the anti-GD2 mAb 3F8 achieves NB killing through antibody-dependent cell-mediated cytotoxicity (ADCC) mediated by myeloid and NK cells (4).

NK activity is regulated by inhibitory and activating signals following engagement of cell membrane receptors with their cognate ligands on target cells (6). Different mechanisms of NK activation and inhibition have been described upon NK interaction with NB. Untreated NB tumors and cell lines are widely reported to have reduced to no HLA class I expression, rendering them potentially susceptible to NK killing due to lack of engagement of HLA class I–specific inhibitory killer cell immunoglobulin-like receptors (KIRs) (7, 8). In addition to CD16-mediated activation by mAb, direct activation of NK cells by NB through NKp30, NKp44, NKp46, and the DNAM-1 receptor has also been described (9–11). To evade NK surveillance, NB cells exhibit poor cell surface expression of the activating ligands MICA, MICB, and ULBPs; and high serum concentrations of soluble MICA in patients with NB results in depressed NK function (8). Furthermore, the 41g-B7-H3 molecule, widely expressed among solid tumors, including NB (12, 13), is a potent inhibitor of NK function (14).

NK cells spare autologous cells from killing through interaction of inhibitory receptors with self-HLA class I antigens on the autologous cell. KIR2DL2 and KIR2DL3 recognize HLA-C allotypes characterized by Asn80 (grouped as HLA-C1); KIR2DL1, and to a weaker extent some KIR2DL2/3 allotypes, recognize HLA-C allotypes characterized by Lys80 (grouped as HLA-C2); KIR3DL1 recognizes HLA-A and HLA-B allotypes with the Bw4 epitope; and the heterodimeric CD94/NKG2A receptor recognizes complexes of HLA-E bound to peptides from the leader sequences of HLA class I molecules (15, 16). Interaction between self-specific inhibitory KIRs and their cognate HLA ligands is fundamental to a process referred to as licensing (17), in which NK cells expressing inhibitory KIRs for self HLA (S-KIRs) are “licensed” and have higher resting capacity for response (IFN-γ production, cytotoxicity, and ADCC) (18, 19). Lacking S-KIRs, “unlicensed” NK cells are significantly less responsive at rest and include cells expressing inhibitory KIRs for non-self HLA (NS-KIR). Under inflammatory conditions, however, unlicensed NK cells can exhibit higher response (20–22). Independent of KIRs, CD94/NKG2A expression is associated with modest response capacity (19). Because the KIR and HLA genes are located on different chromosomes and segregate independently, approximately 60% of individuals have inhibitory KIRs for which they lack the cognate HLA class I ligands (a condition referred to herein as missing KIR ligand) (23, 24) and therefore potentially possess substantial numbers of unlicensed NK cells expressing NS-KIRs.

We previously reported a significant association between KIR/HLA genotypes predictive of missing KIR ligand and survival in 169 patients with NB treated with the anti-GD2 mAb 3F8 following ASCT (25). A smaller study of 38 patients treated with Hu14.18-IL2, an anti-GD2 mAb fused to IL-2, also suggested improved response among patients missing KIR ligand (26). The improved clinical outcomes suggest that unlicensed NK cells expressing NS-KIRs may play...
an important role in patients treated with mAb. In this study, we confirm the impact of missing KIR ligand in patients with NB receiving 3F8 and demonstrate that ASCT is not necessary for this association. We demonstrate that 3F8 in vitro activates NK cells expressing S-KIRs, NS-KIRs, and NKG2A for ADCC, but cytokine-induced upregulation of HLA class I expression on NB targets selectively abrogates activation of S-KIR–positive and NKG2A-positive NK cells. These results not only demonstrate the dominance of the unlicensed NK cell in ADCC and antitumor cytotoxicity but also confirm that missing KIR ligand KIR/HLA genotype combinations represent an important prognostic marker in patients with NB treated with mAb. KIR/HLA genotypes indicative of missing KIR ligand may therefore be a positive prognostic indicator for other patient populations with malignancies treated with mAb.

**Results**

Missing KIR ligand is associated with improved survival in patients with high-risk NB receiving anti-GD2 mAb. We previously reported that patients receiving ASCT with 3F8 had superior OS and progression-free survival (PFS) if they lacked HLA class I ligands for autologous inhibitory KIRs (25). To test whether ASCT is necessary for this benefit, we evaluated updated outcomes of 166 previously reported patients who received 3F8 following ASCT (25) and 76 new patients who received 3F8 following chemotherapy without ASCT. Patient characteristics for both cohorts are listed in Table 1. The median follow-up was 74 months from the institution of 3F8 immunotherapy, and the OS and PFS were similar in both patient groups (data not shown).

Patients were considered missing KIR ligand if they lacked one or more HLA class I ligand for their inhibitory KIRs. In contrast, patients with "all ligands present" possessed all HLA class I ligands for their inhibitory KIRs. 153 patients (63%) were missing KIR ligand, and the proportions of missing KIR ligand were comparable among patients who received chemotherapy without ASCT (62%) and those who received ASCT (64%). In both the non-ASCT and ASCT settings, missing KIR ligand was associated with increased OS and PFS (Figure 1, A and B), indicating that ASCT is not necessary for the "missing ligand" effect and suggesting that the effect may be more related to the mAb 3F8 or to NK-NB interaction in general.

Among patients receiving 3F8, the median OS and PFS were 114 and 50 months, respectively, for patients missing KIR ligand and the proportions of missing KIR ligand were comparable among patients who received chemotherapy without ASCT (62%) and those who received ASCT (64%). In both the non-ASCT and ASCT settings, missing KIR ligand was associated with increased OS and PFS (Figure 1, A and B), indicating that ASCT is not necessary for the "missing ligand" effect and suggesting that the effect may be more related to the mAb 3F8 or to NK-NB interaction in general. Among patients receiving 3F8, the median OS and PFS were 114 and 50 months, respectively, for patients missing KIR ligand and the proportions of missing KIR ligand were comparable among patients who received chemotherapy without ASCT (62%) and those who received ASCT (64%).}

**Table 1**

| Characteristics | Patients treated with chemotherapy plus 3F8 (n = 76) | Patients treated with ASCT plus 3F8 (n = 166) |
|----------------|---------------------------------------------------|---------------------------------------------|
| Stage IV cancer | 75 (99%)                                           | 166 (100%)                                  |
| Age            |                                                   |                                             |
| <1.5 years     | 6 (8%)                                            | 15 (9%)                                     |
| >1.5 years     | 70 (92%)                                          | 151 (91%)                                   |
| MYCN status    |                                                   |                                             |
| Nonamplified   | 45 (59%)                                          | 95 (57%)                                    |
| Amplified      | 21 (28%)                                          | 55 (33%)                                    |
| Unknown        | 10 (13%)                                          | 16 (10%)                                    |
| Lactate dehydrogenase |                                             |                                             |
| <1,500 units/mol | 43 (57%)                                         | 83 (50%)                                    |
| ≥1,500 units/mol| 20 (26%)                                          | 45 (27%)                                    |
| Unknown        | 13 (17%)                                          | 38 (23%)                                    |
| Bone metastases|                                                   |                                             |
| No             | 17 (22%)                                          | 37 (22%)                                    |
| Yes            | 59 (78%)                                          | 129 (78%)                                   |
| Disease status at 3F8 therapy |                                             |                                             |
| First CR/VGPR  | 26 (34%)                                          | 80 (48%)                                    |
| Second CR/VGPR | 2 (3%)                                            | 17 (10%)                                    |
| Primary refractory | 47 (62%)                                      | 41 (25%)                                    |
| Secondary refractory | 0 (0%)                                       | 3 (2%)                                      |
| Others*        | 1 (1%)                                            | 25 (15%)                                    |
| KIR genotype   |                                                   |                                             |
| 3DL1 present   | 73 (96%)                                          | 157 (95%)                                   |
| 2DL2 present   | 44 (58%)                                          | 80 (48%)                                    |
| 2DL3 present   | 65 (86%)                                          | 151 (91%)                                   |
| 2DL1 present   | 75 (99%)                                          | 163 (98%)                                   |
| KIR/HLA ligand |                                                   |                                             |
| All ligands present | 29 (38%)                                     | 60 (36%)                                    |
| Missing ≥1 ligand | 47 (62%)                                     | 106 (64%)                                   |

*Patients who received 3F8 as part of induction therapy in combination with chemotherapy. CR/VGPR, complete response/very good partial response.
ity in vitro, we evaluated unlicensed and licensed NK cell activity from normal individuals in response to NB cell lines in the presence or absence of 3F8. Donors with KIR and HLA genotypes predictive of missing KIR ligand (n = 16) were selected for evaluation, allowing functional comparisons between NK populations expressing S-KIRs or NS-KIRs within each individual. Because donors were of a variety of KIR and HLA ligand backgrounds, licensed and unlicensed NK populations were defined by different KIR/self-HLA combinations (Supplemental Table 2). Four donors with all ligands present were also evaluated.

Multicolor flow cytometry provided single-cell assessment of NK cells exclusively expressing a single inhibitory KIR, permitting evaluation of unlicensed NK cells expressing NS-KIRs distinct from licensed NK cells expressing S-KIRs. Among individuals with missing KIR ligand, unlicensed NK cells exclusively expressing a single NS-KIR represented 3%–35% of the total NK repertoire (Supplemental Table 3). Among these same individuals, licensed NK cells exclusively expressing 1 S-KIR represented 8%–31% of the total NK repertoire. In individuals with all ligands present, licensed NK cells exclusively expressing 1 S-KIR represented 19%–35% of the total NK repertoire. Thus, unlicensed and licensed NK cells expressing a single inhibitory KIR can represent substantial segments of the total NK repertoire.

In each individual, we compared CD107 mobilization as a marker of cytotoxic response among NS-KIR–positive and S-KIR–positive NK cells following stimulation with the HLA class I–negative K562 target cell line or with HLA-genotyped NB cell lines (Supplemental Table 4) in the presence or absence of mAb 3F8. Among all individuals, both NS-KIR and S-KIR NK cell populations had dim CD56 expression, indicating their equiva-

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**Figure 1**

Missing KIR ligand is associated with improved OS and PFS in patients with high-risk NB treated with 3F8. Among patients receiving 3F8, patients lacking class I ligands for autologous inhibitory KIRs (dotted lines) have higher OS and PFS compared with those of patients with all KIR ligands present (solid lines) following (A) chemotherapy alone (OS, HR = 0.53 [95% CI, 0.26–1.08], P = 0.077; PFS, HR = 0.49 [95% CI 0.28–0.88], P = 0.014) or (B) ASCT (OS, HR = 0.57 [95% CI, 0.37–0.88], P = 0.011; PFS, HR = 0.62 [95% CI, 0.41–0.94], P = 0.022), confirming that ASCT is not necessary for the missing ligand effect. (C) Among all patients receiving 3F8, patients missing KIR ligands have higher OS (HR = 0.57 [95% CI, 0.39–0.83], P = 0.003) and PFS (HR = 0.58 [95% CI, 0.42–0.81], P = 0.001) compared with those of patients with all KIR ligands.
lent terminal differentiation and potential for cytotoxic response (27, 28). Furthermore, there was no difference in CD16 expression among the NK populations (data not shown).

As expected, in an HLA-C1/C1, HLA-Bw4/Bw6 individual, licensed NK cells exclusively expressing the self-specific KIR2DL3 or KIR3DL1 receptor were responsive to K562 cells, while unlicensed NK cells expressing the non-self-specific KIR2DL1 receptor were hyporesponsive (Figure 2A). Unexpectedly, the unlicensed NS-KIR spNK cells also NK (spNK) cells for S-KIRs were highly activated when stimulated with the NB cell line LAN-1 (Figure 2A). In the presence of the anti-GD2 mAb 3F8, however, licensed single-positive NK (spNK) cells for S-KIRs were highly activated when stimulated (Figure 2A). Surprisingly, the unlicensed NS-KIR spNK cells also demonstrated higher activation in the presence of 3F8, compared with that of the K562 control. KIR-negative NK cells were also activated but to a lesser degree. These results were confirmed with the K562 control and the 4 other NB cell lines (data not shown), rendering them potentially susceptible to NK cytoxicity, all NK cells, regardless of their licensed status, responded poorly to stimulation with the NB cell line LAN-1 (Figure 2A). In the presence of the anti-GD2 mAb 3F8, however, licensed single-positive NK (spNK) cells for S-KIRs were highly activated when stimulated (Figure 2A). Unexpectedly, the unlicensed NS-KIR spNK cells also demonstrated higher activation in the presence of 3F8, compared with that of the K562 control. KIR-negative NK cells were also activated but to a lesser degree. These results were confirmed with the IMR-32 cell line and 4 other NB cell lines (data not shown), demonstrating that multiple NK subsets are recruited by 3F8 to mediate ADCC against NB target cells. Cytotoxicity assays confirmed that addition of 3F8 was critical for NK-mediated NB lysis (Supplemental Figure 2).

Aggregate analysis of NK response to the NB cell line, LAN-1, from 16 individuals demonstrated that NK cells expressing S-KIRs or NS-KIRs showed strong CD107 mobilization mediated by 3F8 (P < 0.0001) (Figure 2B). A higher percentage of S-KIR spNK cells compared with NS-KIR spNK cells exhibited CD107 mobilization (P = 0.004), confirming that licensed NK cells are more activated for ADCC (18, 19). The aggregate analysis also showed response among KIR-negative NK cells, albeit to a lower level than that in the NS-KIR–positive population (P = 0.006).

Based on the results in the KIR-negative NK cells, we surmised that CD94/NKG2A may contribute to the NK cell response to LAN-1 cells in the presence of 3F8. Indeed, NKG2A expression contributed to response among all NK cell populations, including the KIR-negative cells (Figure 2C). Among individuals with missing KIR ligand, NKG2A-positive KIR-negative NK cells represented 13%–51% of the total NK repertoire, and there was no preferential coexpression of NKG2A among NS-KIR or S-KIR populations (data not shown).

**Table 2**

Factors associated with survival

| Variable             | Univariate analysis of OS | Multivariate analysis of OS | Univariate analysis of PFS | Multivariate analysis of PFS |
|----------------------|--------------------------|-----------------------------|---------------------------|-----------------------------|
|                      | HR (95% CI)              | P value                     | HR (95% CI)              | P value                     |
| Missing any ligand   | 0.57 (0.39–0.83)         | 0.003                       | 0.38 (0.32–0.71)         | <0.001                      |
| Age ≥1.5 years       | 2.53 (1.03–6.22)         | 0.043                       | 2.32 (0.92–5.85)         | 0.074                       |
| Bone disease         | 1.20 (0.75–1.91)         | 0.449                       | 1.10 (0.67–1.81)         | 0.709                       |
| MYCN amplification   | 1.04 (0.69–1.56)         | 0.868                       | 1.19 (0.78–1.82)         | 0.407                       |
|                      | 1.06 (0.73–1.53)         | 0.758                       | 1.18 (0.81–1.71)         | 0.397                       |
|                      | 0.52 (0.36–0.74)         | 0.002                       | 0.52 (0.36–0.74)         | 0.001                       |
|                      | 1.04 (0.69–1.56)         | 0.868                       | 1.19 (0.78–1.82)         | 0.407                       |
|                      | 1.06 (0.73–1.53)         | 0.758                       | 1.18 (0.81–1.71)         | 0.397                       |
|                      | 0.52 (0.36–0.74)         | 0.002                       | 0.52 (0.36–0.74)         | 0.001                       |
Figure 3C). Among NKG2A-negative cells, NS-KIR-positive NK cells were more responsive than KIR-negative NK cells (P = 0.001). Similar results were obtained with intracellular IFN-γ response (data not shown). Aggregate analysis of NK activity among 3 individuals with all ligands present was consistent, with inhibition of all S-KIR-bearing NK subsets by NB target cells expressing the cognate HLA class I ligands (P < 0.0001, Figure 3C).

When NKG2A-positive cells were evaluated, NKG2A coexpression enhanced ADCC response of all NK cell subsets, but it did not prevent HLA class I-mediated inhibition of licensed cells (Supplemental Figure 3). Moreover, NKG2A-positive cells were also inhibited by HLA-E expression on the BE(2)N target cell, as evidenced by suppression of activity among the NKG2A-positive KIR-negative subset (Supplemental Figure 3).

Cytokines released by activated NK cells induce HLA class I expression on NB cell lines. Induced HLA expression has been described in NB tumors isolated from patients receiving treatment (29). To determine whether upregulation of HLA class I molecules on NB target cell lines is directly induced by agents commonly used in the treatment of NB, the NB cell lines LAN-1 and BE(2) N were incubated with GM-CSF, isotretinoin, or with different chemotherapy agents. No increase in HLA class I expression on the NB cell lines was detected following treatment with these agents (data not shown). We then investigated whether cytokines released from NK cells activated by 3F8-dependent ADCC could induce HLA class I expression on NB cells. PBMCs and NB targets were coincubated for 24 hours. Supernatant obtained following 24 hours of coculture was collected and used as a culture medium for the NB cell lines LAN-1 and BE(2)N (Figure 4). Supernatant from PBMCs alone did not affect HLA class I expression on NB cell lines, and the addition of GM-CSF to PBMCs resulted in supernatant that minimally increased HLA class I expression on the NB target. Coincubation of 3F8 with PBMCs and NB cells, however, produced a supernatant capable of inducing HLA class I expression on LAN-1 and BE(2)N cells.
Licensed NK cells are selectively inhibited by HLA class I ligands induced on NB targets. (A) Incubation with IFN-\(\gamma\) results in increased HLA class I antigen expression on LAN-1 and BE(2)N cells. HLA-Bw4 and HLA-E antigens are readily induced on BE(2)N but not LAN-1 cells. (B) CD107 degranulation and IFN-\(\gamma\) production in NKG2A-negative, KIR-positive NK cells in response to IFN-\(\gamma\)-treated NB targets in the presence of 3F8. When NK cells from individual no. 17 (HLA-C1/C1/Bw6/Bw6) are incubated with LAN-1 cells, mimicking effectors and NB cells in a patient lacking HLA-C2 and HLA-Bw4 ligands, both NS-KIR–positive (KIR2DL1 single positive [KIR2DL1sp] and KIR3DL1sp) and S-KIR–positive (KIR2DL3sp) NK cells are activated in the presence of 3F8, but only KIR2DL3 spNK cells are inhibited by IFN-\(\gamma\)-induced expression of self HLA on tumor target. When NK cells from individual no. 20 (HLA-C1/C2/Bw4/Bw4) are incubated with BE(2)N cells, mimicking effectors and NB cells in a patient with all ligands present, S-KIR–positive NK subsets (KIR2DL1sp, KIR2DL3sp, and KIR3DL1sp) are inhibited by IFN-\(\gamma\)-induced expression of cognate ligands on the tumor target. In both individuals, blocking antibodies to HLA class I fully restores response among S-KIR–positive NK cells. (C) Aggregate CD107 response among NK populations from 9 HLA-C1/C1 individuals to LAN-1 cells. HLA class I upregulation on the target inhibits S-KIR spNK cells, resulting in a comparatively stronger response from NS-KIR spNK cells (\(P = 0.003\)). Aggregate CD107 response among spNK cells from 3 HLA-C1/C2/Bw4 individuals to BE(2)N demonstrates that HLA class I upregulation on the target inhibits all self-specific NK subsets. Symbols represent individual samples (mean ± SEM). **\(P < 0.01\), ***\(P < 0.001\).
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(Figure 4A) to levels nearly equivalent to the level attained following the direct addition of IFN-γ (Figure 3A). HLA-Bw4 expression was also specifically induced, albeit to a lower level than that induced in the other class I molecules.

We surmised that release of cytokines by tumor-activated PBMCs in the presence of mAb was responsible for the supernatant-mediated upregulation of HLA class I on the tumor target. Using ELISA, we determined the concentrations of IFN-γ released in the different supernatants. PBMCs alone did not produce detectable IFN-γ, and the addition of GM-CSF resulted in the release of a minimal amount of IFN-γ at 24 hours (Figure 4B). However, PBMCs activated by NB target cells in the presence of 3F8 released a substantial amount of IFN-γ, peaking at 12 hours, and the addition of GM-CSF to 3F8 further enhanced IFN-γ production, reaching 60 pg/ml by 24 hours (Figure 4B). Using titration assays, we determined that HLA expression on LAN-1 and BE(2)N cells could be induced with as little as 10 pg/ml IFN-γ (Figure 4C). We therefore conclude that concentrations of IFN-γ released by tumor-activated PBMCs in the presence of 3F8 and GM-CSF are adequate to induce HLA expression on the tumor.

Discussion
When treated with anti-GD2 mAb, patients with NB who lack one or more HLA ligand for their inhibitory KIRs respond better to treatment, have lower rates of relapse, and survive longer compared with patients who possess all HLA KIR ligands. These results are independent of stem cell transplantation and suggest that the missing ligand effect is a result of either the 3F8 mAb therapy and/or the interaction between NK cells and the NB target. The response and survival advantages associated with missing KIR ligand genotypes, denoting individuals who lack HLA ligands for his/her autologous inhibitory KIR, suggested to us that the

**Figure 4**

IFN-γ released by activated NK cells induces upregulation of HLA class I on NB cells in vitro. (A) HLA-Bw4 and HLA-A, HLA-B, and HLA-C expression on the LAN-1 and BE(2)N NB cell lines is shown following 72 hours in different culture conditions. Supernatants collected from PBMCs co-cultured with LAN-1 and BE(2)N cells and 3F8, with or without GM-CSF, induced HLA class I expression on LAN-1 and BE(2)N cells, respectively; in comparison, supernatant collected from PBMCs incubated with or without GM-CSF did not induce HLA expression. (B) By ELISA, PBMCs alone or PBMCs with GM-CSF produced no or minimal IFN-γ. PBMCs activated by LAN-1 cells in the presence of 3F8 and GM-CSF released a substantial amount of IFN-γ compared with baseline and reached 60 pg/ml at 24 hours. (C) Titration assays demonstrate that HLA class I expression can be induced on LAN-1 and BE(2)N cells with 10 pg/ml IFN-γ and that expression increased in a dose-dependent manner.
unlicensed NK cells expressing KIRs for non-self-HLA ligands in these individuals may be more effective in tumor eradication than the licensed NK cells expressing KIRs for self HLA. We considered 2 possible mechanisms underlying these clinical findings: activation among the NS-KIR-specific NK population by 3F8 and/or selective inhibition of the S-KIR-specific population. For NB treated with 3F8, both mechanisms appear to be important.

In vitro, licensed NK cells expressing S-KIRs and unlicensed NK cells expressing NS-KIRs are both activated by NB targets in the presence of 3F8, but only the licensed cells are inhibited by cytokine-induced expression of self-HLA molecules on tumor targets. While NKG2A expression contributes to the ADCC response, NKG2A-bearing NK cells are also silenced by HLA-E expression on tumor cells (27, 35). Taken together, the data suggest that upon encountering target cells with HLA class I expression, unlicensed NK cells bearing receptors for non-self HLA are the primary mediators of ADCC. Several studies examining tumor samples from patients with NB and mouse models indicate that upregulation of HLA class I occurs in vivo. All patients in this study received chemotherapy, GM-CSF, and anti-GD2 mAb therapy, and most patients received isotretinoin. It has been reported that tumor specimens from patients with NB treated with chemotherapy upregulate class I expression as a result of partial differentiation (29); retinoid derivatives induce cellular differentiation and HLA class I expression on NB cells (30); and, in NB-bearing mice treated with anti-GD2 mAb, recurrent tumors exhibit greater than 5-fold higher MHC class I expression (31). We could not demonstrate in vitro that agents used to treat NB could directly upregulate HLA class I expression on certain NB cell lines, suggesting that in vivo conditions may have contributed to previous findings or that the cell lines we used may not respond with HLA class I upregulation to these agents (30). We show clearly, however, that PBMCs responding to NB in the presence of mAb produce cytokines, in particular IFN-γ, at concentrations sufficient to induce HLA class I expression on NB cells. Other cytokines, such as tumor necrosis factor, can further enhance HLA class I expression, including HLA-Bw4 (data not shown), and it is possible that these cytokines are also released in ADCC-dependent lymphocyte activation. These in vitro findings offer insight to in vivo conditions, in which recruitment of NK cells to the tumor microenvironment may result in effector/target interactions, leading to significant local cytokine concentrations. Taken together, the data indicate that it is highly likely that NK activation in vivo by 3F8-dependent ADCC leads to cytokine release, inducing HLA class I expression on the NB target, inhibiting licensed NK cells, and facilitating the striking missing KIR ligand effect (Figure 5).

The in vitro studies presented in this report represented activity among NK cells expressing a single inhibitory KIR. Examination of the NK repertoire among 20 individuals revealed that the single-positive unlicensed and licensed NK populations, while varying in size from individual to individual, can each represent a significant proportion of the total NK repertoire. The repertoire also includes smaller populations of NK cells that express two or more inhibitory receptors, often resulting in higher levels of responsiveness (19, 36). However, interaction between just one receptor and its HLA class I ligand is sufficient to inhibit these highly responsive NK cells (19). Thus, licensed NK cells expressing S-KIRs, whether expressing one or more than one inhibitory KIR, will be inactivated by the presence of self-HLA class I on the tumor target.

These findings are highly relevant to patients with other malignancies treated with mAbs, such as rituximab, whose mechanisms of action involve NK-mediated ADCC (37). Higher HLA class I expression on CD20-positive lymphoma cells is associated with attenuation of rituximab-induced NK-mediated ADCC (38). However, correlation of KIR/HLA genotyping among patients receiving rituximab and other mAb therapies has not been systematically explored for the missing ligand effect.

Our data support the incorporation of KIR and HLA genotypes as a prognostic marker in patients with high-risk NB receiving antibody immunotherapy. Indeed, KIR/HLA genotypes are as
strong a predictor of response and survival, if not better than several conventional NB biomarkers (25). An analogous study including only patients with NB, particularly those who do not receive antibody treatment, should be performed to confirm whether mAb is necessary for this effect.

Since licensed NK cells expressing S-KIRs have higher ADCC capacity in general, rescuing licensed NK activity from class I inhibition is desirable and could increase response in all patients, including those with all KIR ligands present. Use of newly developed clonal antibodies directed to the inhibitory KIRs would prevent their engagement with HLA cognate ligands and potentially restore licensed NK function (39). In addition, the use of exogenous NK cells in the treatment of patients with NB may be potentially useful if the patient lacks class I ligands for the donor inhibitory KIRs. Augmentation of innate immunity through adoptive transfer of allogeneic NK cells or the use of agents that increase endogenous NK cell number and activity, such as IL-2, lenalidomide, and anti-CD137 antibody, may also improve NB control, particularly in the presence of 3F8 (40–42).

Unlicensed NK cells expressing NS-KIRs have a clinically significant impact in the control of acute myelogenous leukemia in allogeneic hematopoietic stem cell transplantation (23, 43, 44), and it is clear from this study that their impact extends to other tumor targets. While transplantation and mAb therapy are hardly normal physiologic conditions, they both take advantage of the important pool of unlicensed NK cells, previously thought to be hyporesponsive and therefore potentially less clinically relevant. In mice, unlicensed NK cells play a dominant role in clearing murine cytomegalovirus (45, 46), in which the proinflammatory cytokine environment of viral infection leads to higher function among unlicensed cells. These cells can be recalled upon viral rechallenge to eradicate infected cells more efficiently than licensed NK cells, whose cytotoxic and proliferative function is diminished in the presence of MHC class I expression (45, 46). It is possible that activity among unlicensed NK cells in patients with NB may also be augmented by the proinflammatory state associated with tumor burden. The existence of unlicensed NK cells that have been presumed to be hyporesponsive in an otherwise parsimoniously constructed human immune system has been a paradox. These findings contribute to our understanding of the critical role of the unlicensed NK population in tumor control.

Methods

Patients. This is a retrospective analysis of 242 patients with high-risk NB treated at Memorial Sloan-Kettering Cancer Center between 1994 and 2007 with anti-GD2 mAb 3F8. All but 1 patient received GM-CSF, 76% of patients received oral isotretinoin, and 1 patient received oral β-glucan (http://clinicaltrial.gov/; study nos. NCT00072358, NCT00002560, and NCT00037011). KIR and HLA genotyping. Genomic DNA, extracted from PBMCs or marrow mononuclear cells after remission using the QIAamp DNA Blood Mini Kit (Qiagen), was KIR genotyped using previously published methods (47). Patient HLA alleles were identified by a combination of HLA serology, sequence-based amplification (PCR sequence-specific primer), and oligonucleotide probing of genomic DNA (PCR-specific oligonucleotide probe). HLADQ genotyping for healthy donors was performed at Histogenetics.

Target cells, effector cells, and culture conditions. The human erythroleukemia cell line K562 (ATCC) and human NB cell lines IMR-32 (ATCC) and LAN-1 (provided by Robert Seeger, Children’s Hospital Los Angeles, Los Angeles, California, USA) as well as the human NB cell line NB-1691 (13) (provided by Peter Houghton, Nationwide Children’s Research Institute Center for Childhood Cancer, Columbus, Ohio, USA) and 3 cell lines, BE(1)N, BE(2)N, and SKNLP, derived from patients with NB treated at Memorial Sloan-Kettering Cancer Center were cultured in RPMI-1640 medium supplemented with 10% heat-inactivated fetal bovine serum, 100 U/ml penicillin, 100 μg/ml streptomycin, 1% sodium pyruvate, and 1% 2-mercaptoethanol and incubated at 37°C with 5% CO2. To upregulate HLA class I expression, recombinant human IFN-γ (PeproTech) was added at 1,000 units/ml for 72 hours (33). For supernatant experiments, PBMCs (1 × 10⁴ cells per well) were placed in culture alone or with NB cell lines at a ratio of 1:1, with or without 3F8 and GM-CSF at 2 ng/ml (34) for 24 hours, in 96-well U-bottom plates with 200 μl of medium, described above, per well. The supernatant was centrifuged, filtered through a 22-μm filter (Millipore), and then used as culture media for the same NB cell lines for 72 hours prior to assaying HLA expression. To test induction of HLA class I expression by therapeutic agents, NB cell lines were cultured in complete medium in the presence or absence of vincristine (50 ng/ml and 200 ng/ml), thiopeta (20 μg/ml and 200 ng/ml), and doxorubicin (1 μg/ml) for 48 hours; isotretinoin (5 μM) for 72 hours; or GM-CSF (2 ng/ml) for 72 hours prior to assaying HLA expression. Both PBMCs and purified NK cells from healthy individuals were used for analysis of NK function. PBMCs were isolated by Ficoll-density gradient centrifugation. NK cells were isolated following CD3-depletion and CD56-enrichment using the CultiMacs system (Miltenyi). Effector cells were cultured in medium, described above, supplemented with human IL-2 (Proleukin, Prometheus) at 1,000 U/ml and incubated at 37°C with 5% CO2 for 12 to 16 hours prior to assaying function.

NK activation and flow cytometric analysis. CD107 mobilization and intracellular IFN-γ production were used as indicators of effector cell activation (48–50). PBMCs (5 × 10⁴ cells per well) or NK cells (2 × 10⁴ cells per well) were incubated with NB target cells at ratios of 1:3 and 1:1, respectively, with or without 0.8 μg/ml 3F8 for 4 hours in 96-well U-bottom plates with 200 μl medium, described above, per well. The 3F8 concentration was selected after titration assays demonstrated no difference in vitro NK activation at higher concentrations (Supplemental Figure 4) and is less than the serum concentration achieved in patients treated with 3F8 (1–4 μg/ml) (51). For HLA class I blocking, NB cells were preincubated for 30 minutes at 37°C with an antibody cocktail composed of DX17 (BD Biosciences), and 6A4 and A6.136 (both provided by Alessandro Moretta, Università di Genova, Genoa, Italy) (52) prior to the addition of PBMCs or purified NK cells and 3F8. Isotype controls were used for each sample. PerCP-Cy5.5-conjugated anti-CD107a or Pacific Blue–conjugated anti-CD107a (H4A3 clone, Biolegend) was added to each well prior to incubation. Cells were stained with anti-CD3-APC-Cy7 (clone SK7, BD Biosciences), -anti-CD56-PE-Cy7 (clone B159, BD Biosciences), or -anti-CD56-EDC (clone N901, Beckman Coulter; anti-KIR2DL1/2DS1-APC (clone HP-MA4, ebioscience), anti-KIR2DL2/2DL3/2DS2-FTTC (clone CH1-L, BD Biosciences), -anti-KIR3DL1-PE (clone DX9, BD Biosciences), or -anti-KIR3DL1–Alexa Fluor 700 (clone DX9, Biolegend; anti-NKG2A-PE (clone Z199, Beckman Coulter); or Pacific Blue–conjugated anti-NKG2A (clone Z199, Beckman Coulter). For IFN-γ evaluation, brefeldin-A (10 μg/ml, Sigma-Aldrich) and GolgiStop (6 μg/ml, BD Biosciences) were added to the mixture 1 hour following incubation, and FIX & PERM (Invitrogen) was used for staining. LIVE/DEAD Aqua Fixable Dead Cell Stains (Invitrogen) were used to exclude dead cells from analysis. Cells were analyzed using multicolor flow cytometry on a LSRII instrument with FACS Diva software (BD Biosciences). Results were interpreted using FlowJo software (Tree Star). For multicolor compensation and gating, unstained, single-color, and FMO controls were used. NK cells exclusively expressing a single inhibitory KIR were evaluated as previously described (19, 22). HLA class 1 and GD2 expression on NB cells was evaluated with...
anti-HLA-ABC-FITC (clone G4-2-6, BD Biosciences), anti-HLA-Bw4-FITC (clone FH007, One Lambda), anti-HLA-E-PE (clone 3D12, Biolegend), and anti-GD2–Alexa Fluor 488 (clone 2Q549, Abcam). NK cytotoxicity. Cell-mediated cytotoxicity was analyzed by flow cytometry with the PKH26 membrane dye and TO-PRO-3 DNA label as previously described (19), with and without 3F8.

Cytokine assay. IFN-γ production was measured by ELISA (R&D Systems) according to the manufacturer’s instructions and analyzed using an MRX plate reader (Dynex Technology) and Biolinx 2.22 software (Dynex Technology).

Statistics. OS was defined as the time from 3F8 therapy to the date of death or last follow-up. PFS was defined as the time from 3F8 therapy to the date of relapse, death, or last follow up. Survival probabilities were estimated using the Kaplan-Meier method. The log-rank test was used to compare survival probabilities among patients missing any or possessing all KIR ligands. HRs of missing KIR ligand indicators were estimated by multivariable Cox regression, controlling for age, MYCN amplification, and bone disease. Fisher’s exact test was used to determine whether the presence or absence of 3F8. Paired 2-tailed Student’s t test was used to compute the comparisons within the licensed or unlicensed populations, and the unpaired 2-tailed Student’s t test was used to compute comparisons between the unlicensed and licensed populations. For relative fold increase calculations, the unpaired Student’s t test was used to compare the relative activity of licensed and unlicensed NK cells. A test with a P-value of less than 0.05 was considered statistically significant.

Study approval. Informed consent for specimen collection was obtained from patients or legal guardians in accordance with institutional review board guidelines at Memorial Sloan-Kettering Cancer Center. These studies were approved by the institutional review board of Memorial Sloan-Kettering Cancer Center.

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