Human derived dimerization tag enhances tumor killing potency of a T-cell engaging bispecific antibody

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Abbreviations: BiTE, bispecific T-cell engager; BsAb, bispecific antibody; HDD, HNF1α dimerization domain; tsc-BsAb, tandem single-chain bispecific antibody.

Bispecific antibodies (BsAbs) have proven highly efficient T cell recruiters for cancer immunotherapy by virtue of one tumor antigen-reactive single chain variable fragment (scFv) and another that binds CD3. In order to enhance the antitumor potency of these tandem scFv BsAbs (tsc-BsAbs), we exploited the dimerization domain of the human transcription factor HNF1α to enhance the avidity of a tsc-BsAb to the tumor antigen disialoganglioside GD2 while maintaining functional monovalency to CD3 to limit potential toxicity. The dimeric tsc-BsAb showed increased avidity to GD2, enhanced T cell mediated killing of neuroblastoma and melanoma cell lines in vitro (32–37 fold), exhibited a near 4-fold improvement in serum half-life, and enhanced tumor ablation in mouse xenograft models. We propose that the use of this HNF1α-derived dimerization tag may be a novel and effective strategy to increase the potency of T-cell engaging antibodies for clinical cancer immunotherapy.

Bispecific antibodies (BsAbs) that engage T cells for antitumor immunotherapy have shown tremendous potential in animal studies and early clinical trials. There exist a number of ways to construct BsAb formats that either utilizes the full immunoglobulin G (IgG) framework, single chain variable domains (scFv), or combinations thereof. One of the most clinically successful BsAb formats is a single polypeptide composed of 2 scFv units in tandem (tsc-BsAb), where one scFv binds to CD3 on the surface of T cells and the other scFv targets a tumor antigen. For CD19-positive leukemias and lymphomas, the tsc-BsAb of anti-CD19xanti-CD3 (termed Bispecific T cell Engager, BiTE) was highly effective in inducing T-cell mediated cytotoxicity and preventing tumor growth in xenograft studies. In a human trial using the anti-CD19xanti-CD3 tsc-BsAb (Blinatumomab), major clinical benefits were seen, including partial and complete remissions. In a subsequent trial, Blinatumomab was highly effective at extremely low doses (0.06 mg/m²/day) against pre-B-cell acute lymphoblastic leukemia (ALL) and non-Hodgkin’s lymphoma with manageable cytokine storm toxicity. Bolus injections, however, can result in substantial central nervous system (CNS) toxicities, the underlying mechanisms of which are not well understood but may be due to leakage of the antibody into the brain. In addition to the CD19xCD3 BiTE, other antibodies using the tsc-BsAb format have been developed to several human tumor targets, including MSCP (CSPG4) for melanoma, EpCAM for pancreatic CA, CEA for epithelial cancers, and EGFR for colorectal cancer. Despite these encouraging results, tsc-BsAbs have limitations. Their relatively small size (~55 KDa) predisposes tsc-BsAbs to succumb to rapid renal clearance, requiring continuous infusions over 4–8 weeks to achieve clinical effect. The modest size of these bivalent constructs may also facilitate their leakage into the CNS. An additional technical hurdle is that the monovalency of tsc-BsAbs toward the tumor antigen requires high affinity of the anti-tumor scFv to achieve optimal tumor epitope binding.

In this study we designed a novel platform to enhance the potency and efficacy of T-cell engaging tsc-BsAbs. We considered the ideal BsAb to have bivalent binding to the tumor antigen for enhanced avidity, and a molecular size of 100–120 kDa for optimal tumor penetration, tumor retention, and serum half-life. Given that excess cytokine release is a known side effect of BsAbs, we also considered the ideal BsAb to have functional monovalency of CD3 engagement in the absence of FcR(n) binding to...
reduce recirculation and prolonged cytokine release. To accomplish these objectives, we engineered a human-derived homodimerization tag to increase the avidity of the bispecific antibody to the tumor target, optimizing serum half-life and potentially reducing leakage into the CNS, while increasing T cell mediated antitumor potency without excessive cytokine release. We chose to use the dimerization domain of human hepatocyte nuclear factor 1α, HNF1α (residues 1–32), which forms a tightly wound anti-parallel 4 helix bundle in the dimer conformation. Using this human dimerization sequence, we engineered bivalent binding of the tsc-BsAbs to their tumor targets without introducing immunogenicity. Given that the anti-parallel orientation of the HNF1α dimerization domain would orient any attached domain in the opposite direction, CD3 engagement was designed to be functionally monovalent, a property crucial for the clinical safety of this dimeric BsAb platform.

To test the utility of this HNF1α dimerization domain (HDD), we constructed tsc-BsAb against the carbohydrate epitope on disialoganglioside antigen GD2, a proven tumor target on the cell surface of a broad spectrum of human cancers. These tumors include neuroblastoma, soft tissue sarcomas, osteosarcoma, small cell lung cancer, melanoma, retinoblastoma, and brain tumors. More recently, several laboratories have reported the presence of GD2 on breast cancer stem cells, neuroectodermal and mesenchymal stem cells. The anti-GD2×anti-CD3 tsc-BsAb (GD2xCD3) were composed of single polypeptide chains containing the scFv of anti-GD2 monoclonal antibody 5F11 and the humanized scFv of the anti-CD3 antibody OKT3 without and with the HNF1α dimerization domain (HDD) (see Fig. 1A and 1B). We previously described the potency of this 5F11 GD2xCD3 monomeric BsAb against GD2-positive tumor cell lines and tumor xenografts. The HDD tag was placed at the carboxyl end of anti-CD3 scFv and distal to the anti-GD2 scFv. This was done to maximize the functional affinity for the tumor antigen (GD2) at the N-terminal end of the molecule, but not to CD3, since the anti-CD3 scFvs were sterically restricted due to the anti-parallel arrangement of the HDD helices, which would orient the anti-CD3 scFvs ~180° apart. Enhancement of GD2 binding would potentially enhance tumor killing, whereas enhancement of CD3 binding could lead to anergy or overactivation of T cells causing cytokine storm. The molecular size of the dimerized tsc-BsAb (MW ~118 kDa) would increase to above the renal clearance threshold (~60 kDa), thereby enhancing serum half-life. We compared the HDD tag with 2 other homo-dimerization domains: the synthetic helix-loop-helix (dHLX) peptide which has previously been shown to form an anti-parallel 4 helical bundle that can dimerize scFv fragments, and the human IgG1-Fc domain which forms covalent homodimers (see Fig. 1C and 1D).

We focused our investigation on enhancing the properties of the tsc-BsAb format because it has demonstrated the highest T cell engaging potency compared to other published formats (see Figure 1.)

**Figure 1.** Engineering bispecific antibodies for cancer immunotherapy. Bispecific antibody formats that were compared in this investigation include: (A) GD2xCD3 tandem single chain variable fragment (scFv) bispecific antibody (BsAb) (tsc-BsAb) monomer, (B) GD2xCD3-HDD dimer generated by the addition of the hepatocyte nuclear factor 1α (HNF1α) dimerization domain (HDD) to C-terminus which forms an anti-parallel 4 helix bundle, (C) GD2xCD3-dHLX dimer using a synthetic helix-loop-helix (dHLX) peptide, (D) GD2xCD3-Fc dimer using a human IgG1 hinge and CH2 and CH3 domains. Other possible bispecific antibody formats considered: (E) diabody (non-covalently associated dimers in which each chain comprises 2 domains consisting of VH and VL domains from 2 different antibodies), (F) tandem diabody (TandAb), (G) quadromas (IgG like BsAbs formed by hybrid hybridomas or by engineering Fc heterodimers), and (H) Dock-and-Lock BsAbs (covalent trimer of 2 antitumor Fab fragments and one anti-CD3 scFv).
Fig. 1E-H) such as the diabody\textsuperscript{24} (non-covalently associated dimers in which each chain comprises 2 domains consisting of VH and VL domains from 2 different antibodies), tandem diabody\textsuperscript{25} (TandAb), quadromas (IgG-like BsAbs formed by hybrid hybridomas\textsuperscript{26} or by engineering Fc heterodimers\textsuperscript{27}), and Dock-and-Lock\textsuperscript{28} BsAbs (covalent trimer of 2 antitumor Fab fragments and one anti-CD3 scFv). Direct comparison showed that the tsc-BsAb BiTE format (CD19xCD3) was 700–8000 fold more potent than the tandem diabody,\textsuperscript{29} which in turn was superior to diabodies or quadromas\textsuperscript{25} at redirecting T cells to kill B cell lymphoma cell lines. The superiority of BiTE format over diabody was independently confirmed using the EGFRxCD3 BsAb,\textsuperscript{30} strongly suggestive of a structural preference of the tsc-BsAb format for efficient T-cell engagement. More recently, the CD19xCD19xCD3 Dock-and-Lock BsAb\textsuperscript{28} was generated, but it too showed significantly weaker tumor cytotoxicity when compared to the published cytotoxicity of the BiTE format described by Molhoj et al.\textsuperscript{29} We therefore chose the tsc-BsAb format to test the dimerization tag HDD. Here, we present biophysical data on the dimerization and \textit{in vitro} binding kinetics of the HDD-tagged GD2xCD3 BsAb,\textsuperscript{30} strongly suggestive of a structural preference of the tsc-BsAb format for efficient T-cell engagement. More recently, the CD19xCD19xCD3 Dock-and-Lock BsAb\textsuperscript{28} was generated, but it too showed significantly weaker tumor cytotoxicity when compared to the published cytotoxicity of the BiTE format described by Molhoj et al.\textsuperscript{29} We therefore chose the tsc-BsAb format to test the dimerization tag HDD. Here, we present biophysical data on the dimerization and \textit{in vitro} binding kinetics of the HDD-tagged GD2xCD3 BsAb, and \textit{in vitro} T cell-mediated tumor cell killing, as well as \textit{in vivo} antitumor effects using mouse xenograft models.

**Results**

HDD-tag induced BsAb dimerization and enhanced avidity to tumor antigen GD2

The ability of the HDD tag to induce dimerization was assayed by dynamic light scattering and size exclusion chromatography (see Table 1 and Fig. S1). GD2xCD3 BsAb (MW 54 kDa) had a diameter of 7.2 ± 1.7 nm and was 95% monomeric, whereas GD2xCD3-HDD BsAb (MW 59 kDa for monomer, 118 kDa for dimer) had a diameter of 11.1 ± 0.5 nm and was 97% dimeric. In addition to higher purity, the GD2xCD3-HDD BsAb had a 2-fold increase in yield when stably expressed in CHO cells. To test whether the HDD tag enhanced the functional affinity to tumor antigen GD2, Surface Plasmon Resonance (SPR) and ELISA experiments were done using purified GD2 (Table 1, Fig. 2 and Table 2). SPR analysis showed that the Kon for GD2xCD3 BsAb and GD2xCD3-HDD BsAb were similar (9.07 × 10\textsuperscript{4} 1/Ms and 8.83 × 10\textsuperscript{4} 1/Ms, respectively), but the Koff were very different (2.27 × 10\textsuperscript{-2} 1/s and 3.45 × 10\textsuperscript{-3} 1/s, respectively). This resulted in a 6-7 fold difference in the overall K\textsubscript{D} (250 nM for GD2xCD3 BsAb and 39 nM for GD2xCD3-HDD BsAb). The kinetic analysis confirmed that the HDD tag enhanced the ability of the BsAb to retain its distal target. ELISA binding assays showed an 8-fold enhancement of GD2 binding for GD2xCD3-HDD BsAb when compared with GD2xCD3 BsAb. We also compared the HDD tag to the synthetic dHLX and the human Fc domain. To this end, GD2xCD3 BsAb was produced with either the HDD, dHLX domain, or the Fc domain at their respective C-termini and assayed for GD2 antigen binding (Table 2 and Fig. 2C). The GD2 binding showed that covalent dimers formed by GD2xCD3-Fc BsAb (0.3 nM EC\textsubscript{50}, 99% dimeric by HPLC-SEC) had a two-fold enhancement in avidity to GD2, when compared with the non-covalent dimers formed by GD2xCD3-HDD BsAb (0.6 nM EC\textsubscript{50}). ELISA results showed that binding of both of these constructs were several fold higher than monomeric GD2xCD3 BsAb (5.0 nM

**Table 1.** Bispecific antibody (BsAb) biophysical characterization by Dynamic Light Scattering (DLS), Size Exclusion Chromatography (SEC), and Surface Plasmon Resonance (SPR).

| BsAb Construct | Yield (mg/L culture) | Diameter (nm ± SD) | Purity (by HPLC-SEC) | Kon (1/Ms)\textsubscript{SPR} | Koff (1/s)\textsubscript{SPR} | K\textsubscript{D}\textsubscript{SPR} |
|----------------|----------------------|--------------------|----------------------|-----------------------------|-----------------------------|-----------------------------|
| GD2xCD3        | 2                    | 7.2 ± 1.7          | 95%                  | 9.07 × 10\textsuperscript{4} | 2.27 × 10\textsuperscript{-2} | 250 nM                     |
| GD2xCD3-HDD    | 4                    | 11.1 ± 0.5         | 97%                  | 8.83 × 10\textsuperscript{4} | 3.45 × 10\textsuperscript{-3} | 39 nM                      |

![Figure 2](https://example.com/figure2.png) Comparison of bispecific antibody GD2 tumor antigen-binding kinetics. Bispecific antibody GD2 binding kinetics by surface plasmon resonance of (A) GD2xCD3 BsAb and (B) GD2xCD3-HDD. Traces are shown at the following BsAb concentrations: 62.5, 125, 250, 500, 1000, and 2000 nM. Analyses of the association and dissociation rates are shown in Table 1. (C) Binding to tumor antigen GD2 by ELISA for 4 BsAb constructs. Data are shown as the mean ± SD. The resultant EC\textsubscript{50} of binding are shown in Table 2.
Table 2. Analysis of binding of the indicated bispecific antibody (BsAb) to GD2 tumor antigen by ELISA (mean ± SE).

| BsAb Construct | GD2 binding | EC50 (nM) | Relative binding | P-value |
|----------------|-------------|-----------|------------------|---------|
| GD2xCD3        | 3.0 ± 4.8   | 193       | 1.0              | <0.0001 |
| GD2xCD3-HDD    | 0.6 ± 0.1   | 8.3       | 0.1              | <0.0001 |
| GD2xCD3-Fc     | 0.3 ± 0.1   | 16.7      | 0.5              | <0.0001 |
| GD2xCD3-dHLX   | 2.9 ± 0.6   | 1.7       | 0.1982           |         |

Table 3. Analysis of binding of the indicated bispecific antibody (BsAb) to CD3 on T cells by cytofluorometric analysis (mean ± SE).

| BsAb Construct | CD3 binding | EC50 (nM) | Maximal binding | P-value |
|----------------|-------------|-----------|-----------------|---------|
| GD2xCD3        | 42 ± 21     | 14.0      | 2.4             |         |
| GD2xCD3-HDD    | 73 ± 23     | 16.7      | 2.2             | 0.6963  |
| GD2xCD3-Fc     | 24 ± 3.6    | 18.2      | 0.8             | 0.0043  |
| GD2xCD3-dHLX   | Poor fit    | Poor fit  | NA              |         |

Figure 3. Bispecific antibody binding to human T cell CD3 and cytokine release. T cells purified from human peripheral blood mononuclear cells were incubated with the indicated bispecific antibody (BsAb) and characterized for CD3 binding (cytofluorometric analysis) and cytokine release (ELISA). (A) Binding of GD2xCD3, GD2xCD3-HDD, GD2xCD3-Fc, and GD2xCD3-dHLX BsAbs to CD3 on the surface of T cells by flow cytometry. (B) Cytokine release from T cells induced by GD2xCD3 and GD2xCD3 BsAb when compared with a huOKT3 IgG. (C) Cytokine release from T cells co-cultured with neuroblastoma IMR-32 cells in the presence of the indicated BsAb. Data points are shown as mean ± SD. Cytokine values and statistical analyses are shown in Table S1 and Table S2.

Figure 4. Bispecific antibody stimulated T cell-mediated cancer cell cytolysis. T cells purified from human peripheral blood mononuclear cells were incubated with the indicated bispecific antibody (BsAb) in the presence of cancer cells. T cell-mediated killing of (A) M14 melanoma, (B) LAN-1 neuroblastoma, and (C) IMR-32 neuroblastoma cells lines was analyzed by chromium51-release assay. Data are shown as the mean ± SD. The EC50, maximal killing, and statistical analyses are shown in Table 4.
CD3 binding, cytokine release, and T-cell proliferation

We also compared T cell CD3 binding by fluorescence cytomtery for all 4 constructs (GD2xCD3, GD2xCD3-HDD, GD2xCD3-dHLX, GD2xCD3-Fc BsAbs) (Fig. 3A and Table 3) and found no significant difference between the binding GD2xCD3 and GD2xCD3-HDD BsAbs (P = 0.6963), indicating functional monovalency to CD3. GD2xCD3-Fc BsAb showed significantly higher CD3 binding (P = 0.0043), with a nearly 2-fold decrease in EC50. This result indicates that the orientation of the anti-CD3 scFv fragments (HDD tagged versus Fc-tagged) can directly influence CD3 binding avidity. The GD2xCD3-dHLX BsAb showed the lowest CD3 binding, likely due to the observation that the sample did not form homogeneous dimers and contained aggregates.

We also characterized the ability of GD2xCD3 and GD2xCD3-HDD BsAb to induce T-cell mediated cytokine release in the presence or absence of GD2-expressing tumor cells (Fig. 3C and 3D, Table S1 and Table S2). The BsAb stimulated release of the cytokines tumor necrosis α (TNFα), interferon γ (IFNγ) and the interleukins IL10 and IL2 were measured. GD2xCD3-HDD BsAbs stimulated comparable cytokine release as GD2xCD3 (i.e., there was no significant difference), suggesting that although GD2xCD3-HDD dimer has 2 anti-CD3 moieties, it functions as a monovalent binder in regards to CD3-mediated cytokine release. When huOKT3 IgG (which is known to have 2 anti-CD3 moieties) was included in the cytokine release assay, it induced substantially more cytokine release than either of the GD2xCD3 or GD2xCD3-HDD BsAbs in the absence of tumor cells (Fig. 3B). However, in the presence of tumor cells a robust cytokine release was observed in the presence of all the BsAbs tested, with no appreciable difference observed between that induced by the monomer vs. the HDD-dimeric form (Fig. 3C).

We also compared the ability of GD2xCD3 and GD2xCD3-HDD BsAbs to induce CD3-mediated T cell proliferation as compared to huOKT3 IgG (Fig. S2). T-cell proliferation was robust in the presence of huOKT3 IgG, while it was negligible when GD2xCD3 or GD2xCD3-HDD BsAbs were used.

Mouse xenograft studies

A series of mouse xenograft studies were conducted to test the efficacies of GD2xCD3-HDD as compared to GD2xCD3 BsAb in vivo. Immunodeficient double knockout (DKO) mice were implanted with subcutaneous neuroblastoma IMR-32 or melanoma M14 cells mixed with peripheral blood mononuclear cells (PBMC) effector cells (see Methods). At day 5 post-implantation, 5 mice per group were given either no treatment, intravenous injections of GD2xCD3 BsAb (6 times per week for 2 weeks), or intravenous injections of GD2xCD3-HDD BsAb (6 times per week for 2 weeks). Tumor volumes and survival data were collected (Fig. 5). The area under the curve (AUC) was calculated for each mouse, and the averages shown in Table 5.

In the case of mice implanted with neuroblastoma IMR-32 cells, administering GD2xCD3 BsAb modestly reduced tumor growth (19% lower AUC, P = 0.541), whereas GD2xCD3-HDD treatment on the other hand, significantly diminished tumor growth (65% lower AUC, P = 0.018). Survival data at the end of the experiment (155 d post implantation) revealed that in comparison to the 3/5 mice that remained alive in the
GD2xCD3-HDD treatment group, only 1/5 mice survived in the GD2xCD3 group and there were no survivors in the untreated group. In the case of mice implanted with melanoma M14 cells, injection of GD2xCD3 BsAb group markedly reduced tumor growth (57% lower AUC), albeit just beyond statistical significance ($P = 0.065$). The GD2xCD3-HDD treatment again significantly reduced tumor growth (75% lower AUC, $P = 0.024$). Survival data at the end of the experiment (82 d post implantation) found that 2/5 mice remained alive in the GD2xCD3-HDD treated group vs. 1/5 in the GD2xCD3 group and no survivors in the untreated group.

A non-targeting animal mouse xenograft study was also performed using the GD2-negative tumor BV-173 (Fig. S3 and Table S3). The results showed that tumor growth in the mice treated with GD2xCD3-HDD and GD2xCD3 BsAbs were comparable to the non-treated control. Another control mouse experiment was also carried out to examine T-cell retention at the tumor site. Mice were implanted with SH-SY5Y cells expressing a low level of GD2 and were subsequently treated with GD2xCD3-HDD or GD2xCD3 BsAb. This line was chosen to test if the HDD tag may enhance the retention of T cells (resulting in possible excess cytokine release) at the tumor site, without being confounded by enhanced avidity of the BsAb for GD2. Tumors were removed and stained for CD3 (Fig. S4 and Table S4). The data demonstrated no significant difference in T-cell retention at the tumor site, providing further evidence that the HDD-dimeric BsAb did not differ from the monomeric BsAb in inducing T-cell proliferation or T-cell survival in the absence of a tumor expressing abundant GD2.

**Pharmacokinetic analysis**

Serum pharmacokinetic analysis was carried out in DKO mice after intravenous injections of BsAbs, in which serum BsAb levels were measured by using an anti-OKT3 and anti-5F11 double sandwich ELISA assay (see Methods). Results are shown in Fig. 6 and Table 6. The Cmax for GD2xCD3 and GD2xCD3-HDD BsAbs were comparable, at 22.95 and 23.72 µg/mL, respectively. However, GD2xCD3-HDD BsAb displayed a significantly longer serum half-life ($t_{1/2}$) of 54.68 ± 17.62 min than GD2xCD3 BsAb (14.04 ± 10.42 min, $P = 0.002$). There was also a 2-fold increase in sustaining GD2xCD3-HDD BsAb concentrations overtime, as shown by the ~2X increased AUC (h × µg/mL) relative to the GD2xCD3 BsAb ($P = 0.018$).
Table 6. Analysis of pharmacokinetics of the indicated bispecific antibody (BsAb) based on non-compartmental analysis (mean ± SD).

|                        | Half life (min) | Cmax (µg/mL) | AUC (hr.µg/mL) | Volume of distribution(mL) | Clearance (mL/hr) |
|------------------------|----------------|--------------|----------------|---------------------------|------------------|
| GD2xCD3 BsAb           | 14.04 ± 10.42  | 22.95 ± 13.07 | 10.64 ± 6.42   | 1.92 ± 1.01               | 7.43 ± 5.90      |
| GD2xCD3-HDD BsAb       | 54.68 ± 17.62  | 23.72 ± 5.38  | 22.09 ± 5.76   | 2.62 ± 0.59               | 2.08 ± 0.51      |
| fold difference        | 3.9            | 1.0          | 2.1            | 1.4                       | 0.3              |
| P-value                | 0.002          | 0.907        | 0.018          | 0.219                     | 0.078            |

Discussion

Tumors evade T cells by downregulating tumor HLA and by upregulating regulatory T cells, as well as interfering with the homing of cytolytic T cells (CTL). Lymphocytes typically comprising low clonal frequency. By engaging CD3 on the surface of T cells, BsAbs activate and redirect polyclonal T cells to tumors, potentially overcoming both issues of insufficient HLA and low clonal frequency in T-cell mediated immunity. In this report, we hypothesized that anti-CD3 scFv units were connected to the HDD tag using a flexible (GGGGS) linker, thereby facilitating bivalent antibody for detection and sandwich ELISA. Parameters from non-compartmental pharmacokinetic analysis are shown in Table 6.

HDD and GD2xCD3-Fc BsAb is the orientation of the anti-CD3 scFv accounting for the distinct CD3 avidities.

ScFv fragments together in a stable way, without limiting the flexibility of the anti-CD3 domain to retarget T cells for cancer cell killing. In fact, we have tested a smaller dimerization domain derived from the peptide hormone Endothelin-1, and found that it was not strong enough to form stable dimers when fused to the GD2xCD3 BsAb (data not shown). The HDD tag had the added benefit of functionally monovalent CD3 binding as compared to GD2xCD3-Fc BsAb, the latter showing enhanced binding to CD3. One of the major differences between the GD2xCD3-HDD and GD2xCD3-Fc BsAb is the orientation of the anti-CD3 scFv accounting for the distinct CD3 avidities.

Since antibodies that bind to carbohydrate antigens such as GD2 are generally of low affinity, the use of stable non-immunogenic peptide sequences to induce homodimerization presents a novel approach to enhance T-cell targeting for cancer immunotherapy. Additionally, dimerization of tsc-BsAbs, which are ~55 kDa in their monomeric form, substantially increase the serum half-life and potential therapeutic efficacy regardless of tumor affinity. The short observed half-life of the monomeric GD2xCD3 BsAb is comparable to that which has been previously reported for tsc-BsAb in mice and Fab fragments that are similar in size. The serum half-life is expected to be substantially longer in humans, based on interspecies scaling.

The nearly 4-fold enhancement in serum half-life resulting from the HDD tag is also consistent with enhancement in serum pharmacokinetics observed between F(ab) and F(ab')2, which are approximately the same sizes as a monomeric and dimeric tsc-BsAb. This limited increase in serum half-life may be more beneficial than the prolonged half-life of a bispecific antibody with an intact Fc domain, which can last in the blood for several weeks due to FcR-mediated recycling and potentiate prolonged...
cytokine release. Other methods exist to prolong serum half-life to a narrow range primarily by pegylation (addition of polyethylene glycol polymers), but controversy exists as to whether pegylation itself is immunogenic. Addition of the HDD tag presents an effective way to enhance the pharmacokinetic properties of the BsAb with a human-derived tag, which is less likely to be immunogenic. In addition, an HDD tagged dimeric tsc-BsAb may be less likely to cross the blood brain barrier, although addressing this possibility requires further investigation.

In this study, we sought to improve the potency of the tsc-BsAb format to stimulate tumor cell killing, while overcoming some technical limitations by using a simple dimerization strategy. We thus utilized the HDD tag with a BsAb that specifically targets the tumor antigen GD2, an immunogen highly expressed in several metastatic cancer types. Anti-GD2 antibodies have shown long-term safety and efficacy in randomized trials, but one of the major side effects is complement-mediated acute pain occurring during antibody infusion. Because GD2xCD3 BsAb lack the Fc portion necessary for complement activation, this strategy of using monomeric or dimeric BsAbs may not be restricted to enhanced tumor cell killing in vitro and in vivo, but may also be associated with reduced pain as a therapeutic side effect. This dimerization strategy can also be applied to other tsc-BsAbs under clinical development, since it can be easily adopted to enhance tumor binding and to increase serum half-life, with the potential to achieve greater therapeutic efficacy without excess cytokine release.

**Methods**

**Molecular cloning**

The sequences of anti-GD2 antibody 5F11 and humanized anti-CD3 have been previously reported. 5F11 scFv (MH-orientation) and anti-CD3 hOKT3 scFv (VH-VL) with an orientation were genetically assembled by a 15 amino acid linker ((GGGGS)5) and synthesized separately (GenScript, Piscataway, NJ). 5F11scFvs were digested with NheI and ApaI, hOKT3 scFv was digested with NheI and ApaI and BamHI, and sequentially ligated into Gluthione synthesis vector (GS) (Invitrogen) to make GD2xCD3 BsAb. Subsequently, 2 cysteine mutations, residue S44C on the heavy chain and residue A100C on the light chain, were introduced by site-directed mutagenesis (Stratagene, CA) to stabilize the 5F11 scFv. Additionally, one affinity maturation mutation P104Y was also incorporated into the 5F11 scFv. The final sequence of the GD2xCD3 BsAb was as follows:

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QVQLQQLPHELVPGPSKISCKTSGYKTFETYTMHWV
KQSHGKCLEWIGGInPNNGNTYNQFKFKGATLTV
DKSSSTAYMEILRSTSEDAYSVCARDTTTVYAWGQ
TTVTVSSGGSAGGSGGGSSGSDIELTQSPAIMASPG
EKTMTCSASSSISHMHWYQQP1GTPSGBKIWYDTKLS
SVPARFSGSGTGYSSTLSSMEADATYYCHQSSYPLT
FCCGTCLEIRASTKGPGGSSGSSGGGSGGQQLV
QSGGGVYVQGPRSRLSCKASYGFTFRYTMHWVRQAPG
KGLEWIGYINPSRGTNYNQFKDRFTISRDNSKNTAFL
QMDLSRPEDTGVYFCARYDYDDHYCLYWQGPTVVS
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The sequence of the GD2xCD3-dHLX was identical to GD2xCD3-HDD, except the HNF1 domain and Fc sequence were added to the C-terminus of the GD2xCD3 BsAb.

For the GD2xCD3-Fc BsAb construct, a human IgG1 hinge domain and Fc sequence were added to the sequence of GD2xCD3 BsAb.

**Production and purification**

Expression constructs were made with the DNA of GD2xCD3, GD2xCD3-HDD, or GD2xCD3-dHLX BsAb followed by a 6×His tag. An expression construct with GD2xCD3-Fc was also made, with no additional purification tags. DNA was transfected into DG44 CHO-S cells (Invitrogen) by electroporation using a Nucelofector II electroporation machine (Amazza) and nuc ellefication solution V. Transfected cells were subject to drug selection with 500 μg/mL G418. Two weeks later, single cells were plated in 96-wells by serial dilution. Irradiated CHO-S cells (5000 per well) were used as feeder cells. Supernatants from each clone were harvested in 3 weeks, and subject to GD2 binding assay. Clones with the highest binding to GD2 were picked, scaled up for large culture via orbital shanking at 125 rpm in 37°C (8% CO2) and cultured until the cells reach 2 million cells/mL and are in log-phase growth. Cultures were harvested when the desired antibody yield was achieved or when the viability dropped to <40%. BsAb proteins with 6×His tags secreted into the culture supernatant were subsequently affinity purified by Ni2⁺ sepharose (GE Healthcare Bio-Sciences, Sweden), eluted with 300 mM imidazole, dialyzed in PBS pH 7.4, and purified by size exclusion chromatography (Superdex 200 GL column, GE Healthcare Bio-Sciences). BsAbs with an Fc domain were purified by affinity chromatography using MabSelect resin (GE Healthcare Bio-Sciences). The hydrodynamic radii of the sample solution were measured by dynamic light scattering using a Malvern Zetasizer Nano (Malvern Instruments, Worcestershire, UK). Sample purity was also evaluated by size-exclusion high-performance liquid chromatography (SE-HPLC). Approximately 20 μg of protein was injected into a TSK-GE-3000SWXL 7.8 mm × 30 cm, 5 μm column (TOSOH Bioscience) with 0.4 M NaClO4, 0.05 M NaH2PO4, pH 6.0 buffer at flow rate of 0.5 mL/min, and UV detection at 280 nm. Ten microliters of gel-filtration standard (Bio-Rad) was analyzed in parallel for MW markers.
ELISA

To assay GD2 binding, GD2 antigen was first coated at 1 μg/mL per well in 90% ethanol on vinyl 96-well plates at room temperature (RT) overnight. On the subsequent day, after blocking the plates with 150 μL/well 0.5% bovine serum albumin (BSA), a dilution series of the BsAb antibodies were added to the plates and incubated at RT for 2 h. Plates were washed with PBS and 100 μL/well of mouse-anti-His-tag antibody (AbD Serotec; dilute 1:1000 dilution in 0.5% BSA) were added and incubated for another hour. Lastly, 100 μL/well goat-anti-mouse-HRP antibody (Jackson ImmunoResearch; 1:3000 dilutions in 0.5% BSA) were added and incubated for 1 h. Plates were developed using 150 μL/well OPD buffer (Sigma) and read at 490 nm on a spectrophotometer.

T-cell binding by fluorescence cytometry

T cells were expanded using CD3/CD28 beads. BsAbs were incubated with 1 × 10^6 T cells per sample for 30 min on ice. After washing, the T cells were reacted with a anti-huOKT3-scFv-mouseFc antibody (Laboratory of Dr. Nai-Kong Cheung, Memorial Sloan Kettering, New York, NY) at 1 μg per 1 × 10^6 cells on ice for 30 min, washed, and then reacted with a goat anti-mouse R-Phycoerythrin (PE)-conjugated antibody (Jackson ImmunoResearch) for 20 min. T cells were washed and analyzed by fluorescence cytometry on a FACS Calibur (BD Biosciences, San Jose, CA).

In vitro binding kinetics by Biacore T-100 Biosensor (GE Healthcare)

A Biacore T-100 Biosensor, a CM5 sensor chip, and related reagents were all purchased from GE Healthcare. The ganglioside GM1 was obtained from ALEXIS Biochemicals (AXXORA L.L. C.) and GD2 was from Advanced ImmunoChemical. Gangliosides were directly immobilized onto the CM5 sensor chip via hydrophobic interaction, as previously described. The active surface contained only GM1. BsAb samples were diluted in HBS-EP buffer (0.01 M HEPES pH 7.4, 0.15 M NaCl, 3 mM EDTA, 0.05% v/v Surfactant P20) at varying concentrations and injected over the sensor surface at a flow rate of 30 μL/min over 1–2 min. Following completion of the association phase, dissociation was monitored in HBS-EP buffer for 3 min at the same flow rate. At the end of each cycle, the surface was regenerated using 10–20 mM NaOH at a flow rate of 50 μL/min over 1 min. The data were analyzed using the Biacore T-100 evaluation software, and the apparent association on rate constant (kon), dissociation off rate constant (koff) and equilibrium dissociation constant (KD = koff/kon) were calculated.

Cell lines

The neuroblastoma cell line LAN1 and the melanoma line M14 were obtained from University of California, Los Angeles. Neuroblastoma IMR-32 cells (from which SKNLD is derived) and the SH-SY5Y cell line were obtained from American Tissue Type Culture Collection (ATCC). Leukemia cell line BV-173 was obtained from the laboratory of Dr. Richard J. O‘Reilly (Memorial Sloan Kettering, New York NY). Authentication of cell lines was accomplished by short tandem repeat (STR) DNA sequencing. Cells were cultured in RPMI1640 (Cellgro) supplemented with 10% FBS (Life Technologies) at 37°C in a 5% CO₂ humidified incubator.

Cytokine release assay

Peripheral blood mononuclear cells (PBMC) were isolated from healthy donor blood by lymphocyte separation medium (Mediatech) centrifugation. Human T cells were purified by Pan T Cell Isolation Kit (Miltenyi Biotec). T cells (50000/per well) was co-cultured with neuroblastoma cell IMR-32 cell (10,000/ per well) in the presence of BsAb in 37°C in a 96-well plate. Supernatants were harvested at 24 h and the concentrations of 4 different cytokines (IL-2, IL-10, IFNγ and TNFα) were assessed using an ELISA-based cytokine assay kit (OptEIA™ human cytokine set, BD Biosciences). The assay was run according to the manufacturer’s protocol and cytokine levels were quantitated using the standards supplied from the kit. Positive control samples were run using T cells activated with CD3/CD28 immuno-beads to verify that cytokines could be adequately detected.

T cell proliferation assay

T cells were purified from human PBMC using the Pan T Cell Isolation Kit (Miltenyi Biotec). Neuroblastoma cells IMR-32 were irradiated at 80 Gy and resuspended in RPMI (GIBCO) at 0.1 million/mL. The IMR-32 cells were mixed at 1 × 10^4 cells/well (100 μL) with 2 × 10^5 purified T cells (100 μL) in the presence of different antibodies (50 μL) added to 96-well cell culture plate to a final volume of 250 μL/well. T cells were cultured and maintained in RPMI supplemented with FBS in 37°C for 6 d and T-cell proliferation was quantitated using Cell Counting Kit-8 (CCK-8) (Dojindo Mol) assay according to manufacturer’s protocol.

Cell cytotoxicity (chromium51 release assay)

T cells were purified from human PBMC using the Pan T Cell Isolation Kit (Miltenyi Biotec). CD3/CD28 dynabeads (Invitrogen) were then used to stimulate and expand T cells according to manufacturer’s protocol. Expanded T cells were cultured and maintained in RPMI supplemented with FBS and 30 U/mL IL-2. The resultant population of T cells was stained with anti-CD3-perCP Cy5.5, anti-CD4-fluorescein isothiocyanate (FITC), anti-CD8-allophycocyanin (APC), and anti-CD56-PE antibodies (BD Biosciences) and analyzed using a FACS Calibur.

Target tumor cells were harvested with trypsin/EDTA and labeled with sodium51Cr chromate (Amersham, Arlington Height, IL) at 100 μCi/10⁶ cells at 37°C for 1 h. After the cells were washed twice, 5000 target cells/well were admixed with 50,000 effector cells and BsAb antibodies in 96-well polystyrene round-bottom plates (BD Biosciences) to a final volume of 250 μL/well. Samples were prepared in triplicate. The plates were incubated at 37°C for 4 h and then centrifuged at 800 g for 10 min after which 51Cr release into the supernatant was counted.
using a γ-counter (Packed Instrument, Downers Grove, IL). The percentage of specific release was calculated using the formula: 100% (experimental cpm - background cpm) / (10% sodium dodecyl sulfate (SDS) cpm-background CPM), in which cpm are counts per minute of $^{51}$Cr released, total release was assessed by lysis with 10% sodium dodecyl sulfate (SDS; Sigma, St Louis, Mo), and background release was measured in the absence of effector cells.

**Xenograft mouse model**

The immune deficient mouse strain BALB-Rag2$^{-/-}$ -IL-2R-γc-KO (DKO) was kindly provided by Dr. Mamoru Ito, Central Institute for Experimental Animals, Kawasaki, Japan and maintained at Memorial Sloan Kettering Cancer Center. Animals were provided with Sulfatrim chow. All procedures were performed in accordance with the protocols approved by our Institutional Animal Care and Use Committee and institutional guidelines for the proper and humane use of animals in research. In vivo experiments were performed in 6–10 week old mice bred at the Memorial Sloan Kettering Cancer Center. PBMC of healthy donors were isolated from whole human blood from healthy donor (New York blood center). Erythrocytes were depleted by incubation for 15 min with erythrocyte lysis buffer (Lonz) and thrombocytes removed by supernatant aspiration following centrifugation at 100 × g for 10 minutes. All PBMC samples used had similar percentages of T-cell subpopulations (30–50% CD3 positive). Purified PBMC were mixed with tumor cells (1:1 ratio) and implanted in DKO mice subcutaneously. Treatment with BsAb was initiated at day 5. Mice were treated with 10 μg/day of antibody for 2 consecutive weeks (6 d per week). Tumor size was measured by caliper twice per week; both tumor length and width was recorded. Mice were sacrificed if tumor volumes reached 2 cm$^3$.

**Tumor immunohistochemistry**

Tumor sectioning and immunohistochemistry were done at the Molecular Cytology Core Facility of Memorial Sloan Kettering Cancer Center. Paraffin embedded tumor sections were stained for CD3 using Discovery XT processor (Ventana Medical Systems). The primary antibody was done for 3 h, followed by 1 h incubation with biotinylated goat anti-rabbit IgG (Vector labs,) in 10% normal goat serum, 2% BSA in PBS. Incubation with the primary antibody was done for 3 h, followed by 1 h incubation with biotinylated goat anti-rabbit IgG (Vector labs,) at 1:200 dilution. Secondary Antibody Blocker, Blocker D, streptavidin-HRP and DAB detection kit (Ventana Medical Systems) were used according to the manufacturer instructions. Cell staining quantitation was done using MetaMorph (Molecular Devices).

**Pharmacokinetics**

Blood was obtained from tail vein of DKO mice over 6 h after a bolus injection of 50 μg of BsAb. Serum BsAb levels were measured by double sandwich ELISA where BsAb was captured using solid phase human anti-OKT3 antibody (20 μg/mL), and bound BsAb was detected using 5 μg/mL rat anti-5F11 idiotype antibody, followed by mouse anti-rat IgG-HRP antibody at 1:1000 (Jackson ImmunoResearch, West Grove, PA). Pharmacokinetic analysis was carried out by non-compartmental analysis of the serum concentration-time data using WinNonlin software program (Pharsight Corp., Mountain View, CA).

**Statistics**

Binding data and dose response curves were fitted by non-linear regression using GraphPad Prism software, to allow determination of EC50. For comparison of curves, best-fit values for EC50 were analyzed for significance using F tests. Significance of tabulated data were determined by Student’s t-test, and $P < 0.05$ was considered statistically significant. For comparison of different BsAb constructs, molar equivalents were considered be each unit that contained one anti-GD2 domain and one anti-CD3 domain.

**Disclosure of Potential Conflicts of Interest**

MA and NKC were named as inventors in a patent application for multimerization technologies filed by Memorial Sloan Kettering Cancer Center.

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