

# Generalizable design parameters for soluble T cell receptor-based T cell engagers

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## ABSTRACT

While most biological and cellular immunotherapies recognize extracellular targets, T cell receptor (TCR) therapeutics are unique in their ability to recognize the much larger pool of intracellular antigens found on virus-infected or cancerous cells. Recombinant T cell receptor (rTCR)-based therapeutics are gaining momentum both preclinically and clinically highlighted by recent positive phase III human clinical trial results for a TCR/CD3 bifunctional protein in uveal melanoma. Unlike antibody-based T cell engagers whose molecular formats have been widely and extensively evaluated, little data exist describing the putative activities of varied bifunctional formats using rTCRs. Here we generate rTCR/anti-CD3 bifunctionals directed toward NY-ESO-1 or MAGE-A3 with a variety of molecular formats. We show that inducing strong redirected lysis activity against tumors displaying either NY-ESO-1 or MAGE-A3 is highly restricted to small, tandem binding formats with an rTCR/anti-CD3 Fab demonstrating the highest potency, rTCR/anti-CD3 single chain variable domain fragment showing similar but consistently weaker potency, and IgG-like or IgG-Fc-containing molecules demonstrating poor activity. We believe this is a universal trait of rTCR bifunctionals, given the canonical TCR/human leukocyte antigen structural paradigm.

## BACKGROUND

There are two major categories of cancer immunotherapy. The first involves stimulating disease-targeting immune cells to overcome tumor immunosuppression using activating cytokines or small molecules, immune cell costimulators, or by releasing immune suppression via inhibition of immune checkpoint receptors.<sup>1</sup> The second involves cell-based or soluble recombinant agents that redirect immune cells toward diseased cells expressing tumor-specific antigens (TSAs).<sup>2</sup> Targeting can be mediated by transduction of an antibody on autologous or allogeneic immune cells ex vivo for in vivo patient delivery<sup>3</sup> or via the direct administration of soluble multispecific agents that act in trans to bridge effector immune cells to bind and kill tumor cells.<sup>2</sup> Both categories are transforming cancer therapy, and combinations of these two categories are an emerging next step in cancer treatment.

TSAs commonly targeted by antibody or antibody-like immunotherapies include cell surface markers expressed on diseased hematopoietic cancer cells or cell surface markers overexpressed via gene amplified, epigenetic upregulation or other mechanisms on solid tumor tissues.<sup>2</sup> The number of these extracellular targets is limited and their specificity is rarely exclusive to tumor cells. It is estimated that >70% of tumor-specific targets consist of intracellular oncogenes, mutated tumor-suppressor proteins, or fusion proteins inaccessible to antibody targeting including CAR T cells, antibody-drug conjugates, and bispecific T cell engagers.<sup>4</sup> These intracellular target proteins are accessible to the human immune system through adaptive immune recognition by T cells. In particular,  $\alpha/\beta$  T cells express  $\alpha/\beta$  T cell receptors (TCRs) with the capacity for antibody-like V/J ( $\alpha$ ) and V/D/J ( $\beta$ ) recombination to generate vast diversity to recognize foreign proteins. CD8+ T cells use their  $\alpha/\beta$  TCRs to recognize non-self-peptides complexed and displayed by major histocompatibility complex (MHC) class I proteins that are ubiquitously expressed on somatic cell surfaces. Methods for identifying, sequencing, and functionally verifying  $\alpha/\beta$  TCRs have advanced significantly with the discovery of  $\alpha/\beta$  TCRs against many of the most common intracellular cancer antigens displayed by the most prevalent human leukocyte antigens (HLAs).<sup>5</sup>

Recombinant T-cell receptor (rTCR)-based targeting of tumor cells is gaining momentum as a cancer immunotherapy.<sup>6</sup> This includes (1) the recombinant transduction of an oncogene-specific or cancer mutation-specific  $\alpha/\beta$  TCR into autologous patient T cells and reintroduction of these T cells back into patients with the proper T cell antigen-HLA match or (2) the generation of soluble TCR proteins fused to an anti-CD3 single chain variable domain fragment (scFv) to redirect a patient's T cells to target



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antigen/HLA complexes on tumor cells.<sup>6,7</sup> The first generation of soluble rTCR/anti-CD3 fusion molecules has been dubbed ‘immune mobilizing monoclonal TCRs against cancer’ (ImmTACs). While rTCR-T cell therapies have been more prevalent than ImmTACs, these cell therapies suffer from a variety of issues including competition with endogenous TCRs on T cells for the CD3 signaling components, insufficient rTCR expression, insufficient potency, and heterogeneous mixing of the rTCR α and β chains with their endogenous counterparts, which could lead to uncontrolled specificity issues and safety concerns, and challenges with engraftment.<sup>4,8</sup> Once cell longevity is obtained, dosing and potential long-term toxicity can also be difficult to control, which has spurred the recent development of switches and control methodologies for cell therapies.<sup>9</sup>

Soluble rTCR-based protein therapeutics have not been as prevalent for many reasons but are gaining traction, given advances in the design of antibody-based T cell engagers,<sup>2,10</sup> improved TCR discovery and engineering,<sup>7</sup> and improved methods for their production.<sup>11</sup> Importantly, recent positive clinical results from a phase III study of a gp100/HLA-A2-targeted ImmTAC<sup>12</sup> are the first large-scale clinical trials demonstrating activity for T cell-engaging bifunctionals and highlight the benefits of rTCR-based therapies. Spearheaded by ImmunoCore, the ImmTAC technology uses affinity matured rTCRs with high potency to enable T cell redirection toward tumor cells displaying HLA/peptide complexes with receptor densities as low as 50–100 receptors per cell.<sup>7</sup>

Engineering these molecules for therapy is challenging, given the many parameters required for activity including exquisite specificity and manufacturability. rTCR proteins are generally manufactured in *Escherichia coli* and require solubilization of protein fragments followed by oxidative folding and chain pairing, typically resulting in low yields. These challenges have hindered the ability to characterize rTCR/anti-CD3 molecules with geometries and valency beyond the ImmTAC format.<sup>13</sup> Molecular geometry and epitope play a critical role in the activity of bispecific antibody (BsAb)-based T cell engagers<sup>10</sup>; thus, a greater understanding of the role molecular design plays on the function of rTCR-based T cell engagers is warranted. Unlike BsAbs whose epitopes are highly diverse, TCRs use a canonical epitope/binding mechanism to engage HLA-peptides. Thus, we believe molecular parameters derived for rTCR-based bifunctional proteins (BFPs) that drive their potency and activity will be universal across this class of molecules. Recently, we have described a methodology to express soluble TCRs at antibody-like levels in mammalian cells,<sup>11</sup> opening the door to different rTCR/anti-CD3 BFP architectures. Here, we explore many rTCR/anti-CD3 BFP geometries and contrast them for their ability to redirect T cells to kill tumor cells displaying either MAGE-A3/HLA-A1 or NY-ESO-1/HLA-A2 on their cell surfaces. We uncover strict geometrical requirements for high potency and activity of these molecules with rules common to both sets of antigen/

HLAs. These studies lay a general foundation for the design of rTCR/anti-CD3 BFPs.

## MATERIALS AND METHODS

### Production of rTCR/anti-CD3 BFPs

The NY-ESO-1 and MAGE-A3 α/β TCR as well as anti-CD3 SP34 amino acid sequences have been published.<sup>11,14</sup> DNA GeneBlock up to 900 nucleotides encoding for the rTCR/anti-CD3 BFPs were designed and synthesized by Integrated DNA Technologies. Gene segments with 15 nucleotide 5' and 3' overlaps were designed for In-Fusion (Takara Bio USA) cloning of multiple segments into a cytomegalovirus (CMV) promotor-driven mammalian expression vector using HindIII and EcoRI or BamHI and EcoRI restriction sites (Lonza). Briefly, In-Fusion reactions were transformed into *E. coli* strain Top 10 competent cells (Life Technologies). Colonies were picked and clonal DNA was produced by miniprepping (according to Qiagen MiniPrep cat#27104 protocol procedures) and sequenced by an in-house DNA sequencing core. Medium-scale and large-scale plasmid purifications were performed according to the instructions within the Plasmid Plus Midi Kit (Qiagen cat#12945) and Plasmid Plus Maxi Kit (Qiagen cat#12965), respectively. TCR α/β constant domains contained stabilizing designs.<sup>11</sup> All TCR-scFv, TCR-Fab (antibody antigen binding fragment), and 2TCR-Fab sequences contained an 8×Histag for affinity purification. IgG1-Fc-containing constructs used an Fc-heterodimerization design (7.8.60) to enable proper assembly<sup>11</sup> and L234A, L235A, and N297Q mutations to eliminate CH2-domain N-linked glycosylation and any potential effector function. Mature sequences for every construct described in the manuscript are provided in the online supplemental materials.

Transient transfections and protein expressions in Chinese hamster ovary (CHO) cells (CHO K1S) were performed as described previously.<sup>15</sup> Briefly, plasmids were transfected into CHO cells using 8 mg/L Polyethylenimine Max (PEI Max) from Polysciences (cat#247646-2) per 1.5e6 cells/mL. DNA for transfection was mixed at 1.6 mg/L coding plasmid(s)+1.6 mg/L herring sperm DNA. Cells were maintained in Dulbecco's Modified Eagle Medium (DMEM)-containing media with 10 mM L-glutamine (LM-Growth, SAFC cat#59202C-100) and shaken in flasks at 37°C, 6%–8% CO<sub>2</sub> in a humidified incubator prior to transfection and at 32°C post-transfection. Shaker speed was 250 rpm on a ¾-inch diameter shaker. Protein secretion into the media was achieved by N-terminal fusion of each sequence to a mouse antibody kappa light-chain signal sequence. Supernatants were collected for protein purification by centrifugal pelleting of the cell mass followed by 0.2 μM filtration.

Protein purification was performed using either a one-step or two-step process to achieve >90% pure material based on analytical size exclusion. Histagged proteins were affinity purified using a Ni<sup>2+</sup>-nitrotriacetic acid (NTA) immobilized resin (cOmplete His-Tag, Millipore

Sigma) and an AKTA Pure system (GE Healthcare). The column was conditioned in binding buffer (20 mM Tris-HCl pH 8.0, 0.5 M NaCl) prior to running the supernatants over the column. Protein was eluted using the same buffer but with 250 mM imidazole. Fc-containing proteins were captured on a protein A-containing MabSelect Sure affinity resin (GE Healthcare) conditioned using phosphate buffered saline (PBS). Protein was eluted using 0.1 M glycine, pH 3.0 and neutralized using 1 M Tris pH 8.5. If proteins were >90% pure by analytical size exclusion chromatography (SEC) after affinity capture, they were buffer exchanged into PBS at pH 7.2 using VIVASPIN six concentrators with a 10 kDa MW cut-off; otherwise, they were passed over a preparative SEC column (Zepax SRT-10C SEC3000).

Analytical characterization of the purity of the proteins was performed using an Agilent 1100 HPLC equipped with a Phenomenex Yarra SEC-3000 (300×7.8 cm) column using a 100 mM phosphate, 150 mM NaCl, 0.05% azide, pH 6.8 running buffer. Sodium dodecyl sulfate–polyacrylamide gel electrophoresis (SDS-PAGE) was performed using 4%–12% Bis-Tris gels according to the manufacturer (Life Technologies). Reductions and alkylations were performed with 1 mM dithiothreitol followed by 1 mM N-ethylmaleimide at the heating step.

#### rTCR/anti-CD3 BFP competition flow cytometry assay

The anti-CD3 flow cytometry competition assay was performed essentially as described previously<sup>11</sup> with a few minor modifications throughout the protocol. Jurkat tumor cells (ATCC, cat#TIB-152) grew in suspension in Roswell Park Memorial Institute (RPMI) 1640 media /10% fetal bovine serum (FBS) Corning/gentamicin (Gibco) at 37 °C, 5% CO<sub>2</sub>. Cells were transferred to centrifuge tubes and pelleted by centrifugation at 170 g for 7 min. Cells were washed once with PBS buffer. All subsequent steps were performed on ice. Henceforth, ‘wash buffer’ was PBS/2% FBS/0.05% NaN<sub>3</sub>/10% normal goat serum with extra 10% FBS. Blocking buffer was wash buffer supplemented with human BD Fc Block (BD Biosciences, cat#564220). The cells were resuspended in blocking buffer for 15 min, pelleted, washed 2×, and resuspended in wash buffer before adding 50 µL of the cells (0.5×10<sup>6</sup> cells/well) to 96-well plates (Corning 3799). Next cells were pelleted by centrifugation at 170 g for 7 min and wash buffer was aspirated off each well. In a separate 96-well plate (Corning 3879), the rTCR/anti-CD3 BFPs and control mAbs were added to the wells starting at 1000 nM and titrated using threefold dilutions (working concentrations were 2×). rTCR/anti-CD3 BFP and mAb dilutions were mixed with equal volume of phycoerythrin (PE)-SP34-2 (final concentration of 2.25 µg/mL, BD Biosciences cat#552127). This sample mixture was transferred to the 96-well plate (Corning 3799) with cells at 100 µL/well and incubated 45 min. The cells were pelleted and washed three times. Finally, the cells were resuspended in wash buffer with PI (Molecular Probes cat#P3566) and covered with foil. The cells were

then acquired on a Becton Dickinson LSRII Fortessa flow cytometer with BD FACSDiva software V.8.0.1, and data were analyzed using FlowJo V.10.7.1. The gating strategies for distinguishing live versus dead cells and single cells versus doublets/multiplets that were used for Jurkat cells are provided in online supplemental figures S2 and S3.

#### MHC-peptide tumor cell binding of rTCR/anti-CD3 BFPs

Cell surface MHC–peptide binding of the BFPs was performed essentially as described previously<sup>11</sup> with minor modifications throughout the protocol. Cell culture was performed as described previously. The day before running flow cytometry, T25 flasks were seeded with Saos-2 (ATCC cat#HTB-85) or HCT-116 (ATCC cat#CCL-247) cells in ‘growth buffer’ containing RPMI 1640/10% FBS (Corning), gentamicin (Gibco) at 37 °C, 5% CO<sub>2</sub>. The next morning, Saos-2 and HCT-116 tumor cells were washed once with growth buffer. NY-ESO-1 peptide SLLMWITQC (CPC Scientific, product#824724) was added to Saos-2 cells and MAGE-A3 Peptide: EVDPIGHLY (GenScript, product#SC1848) was added to the HCT-116 tumor cells (or not as a negative control) at 6 µg/mL for 3 hours at 37 °C, 5% CO<sub>2</sub>. Next, excess peptide was aspirated off and the cells were washed 3× with PBS buffer. The tumor cells were lifted from the T25 flasks using Accutase (Innovative Technologies cat#AT104). All subsequent steps were performed on ice. Henceforth, wash buffer was PBS/2% FBS/0.05% sodium azide/10% normal goat serum with extra 10% FBS. ‘Blocking buffer’ was wash buffer supplemented with Human BD Fc Block (BD Biosciences cat#564220). The cells were resuspended in blocking buffer for 15 min, pelleted, washed 3×, and resuspended in wash buffer before adding 50 µL of the cells (0.2×10<sup>6</sup> cells/well for tumor cells) to 96-well plates (Corning 3799). For the tumor cells, the rTCR/anti-CD3 BFPs and recombinantly produced anti-CD3 chimeric SP34 hIgG1\_N297Q<sup>16</sup> were added to the wells at 30.0, 3.0, 0.3, and 0.03 µg/mL and incubated 45 min. The cells were pelleted and washed, and the supernatants were aspirated again before adding 100 µL R-PE-conjugated goat anti-human lambda (1:500 dilution; Southern Biotechnology cat#2070-09) or PE-mouse anti-Histag antibody (1:20 dilution, BioLegend cat#362603) in wash buffer for 45 min. The cells were pelleted and washed again. Finally, the cells were resuspended in wash buffer with 1:1000 propidium iodide (PI, Molecular Probes cat#P3566) and covered with foil. The cells were then acquired on a Becton Dickinson LSRII Fortessa flow cytometer with BD FACSDiva software V.8.0.1, and data were analyzed using FlowJo V.10.7.1. The gating strategies for distinguishing live versus dead cells and single cells versus doublets/multiplets that were used for Saos-2 are shown in online supplemental figures S4 and S5, respectively, and virtually identical to the gating strategies for HCT-116 cells.

## T cell-directed lysis activity of the rTCR/anti-CD3 BFPs using exogenous MAGE-A3 or NY-ESO-1 peptide

Redirected lysis assays using exogenous cancer testis antigen peptides were performed essentially as described previously<sup>11</sup> with minor modifications throughout the protocol. Saos-2 and HCT-116 cells were cultured as described earlier. Primary naïve T cells were from Stem-Express (cat#PB03020C, Donor#D001003581) or AllCells (cat# PB009-1F, Lt# 3009286). For the assay, tumor cells were removed from their culture flasks using Accutase as described previously. The tumor cells were resuspended in complete media and seeded at 5000 cells/well in 96-well black, clear bottom plates (PerkinElmer) overnight at 5% CO<sub>2</sub> and 37°C. The next day, the media were removed and SLLMWITQC (NY-ESO-1) peptide or HCT-116 used EVDPIGHLY (MAGE-A3) peptide was added to Saos-2 or HCT-116 cells, respectively, 100 μL total volume, for 3 hours. Next, 50 μL of the peptide solution was removed from each well and rTCR/anti-CD3 BFPs were added (50 μL/well at 4×) or control antibodies (chimeric SP34 IgG1-N297Q anti-CD3 and irrelevant IgG1) were then titrated onto the cells using 1:10 serial dilutions starting at 200 nM (2×) and ending at 0.002 nM (2×) in triplicate for 30 min. During this period, primary T cells were thawed and washed twice in complete media and gentamicin. T cells (100 μL) were then added at 50 K cells/well (200 μL total volume). After the T cells were added, the rTCR/anti-CD3 BFPs were 1× (starting at 100 nM and ending at 0.001 nM) and the peptide was at 1.5 μg/mL for the duration of the experiment. Non-peptide plates were treated the same as those with peptide except for the absence of peptide in the procedure. Cells were incubated for 48 hours at 5% CO<sub>2</sub> and 37°C. At 48 hours, the plates were washed gently (twice) with serum free RPMI 1640. Finally, to determine the amount of tumor cells alive at the end of the incubations, 100 μL RPMI 1640 and 100 μL Cell Titer Glo reagent (Promega) were added, mixed for 2 min on a shaker (slowly) and incubated for 10 min in the dark. Lastly, the luminescence was read on an EnVision V.2130 multilabel reader.

## T cell-directed lysis activity of the rTCR/anti-CD3 BFPs using endogenous levels of MAGE-A3 or NY-ESO-1 peptide

Redirected lysis assays using endogenous levels of cancer testis antigen were performed essentially as described previously<sup>11</sup> with minor modifications throughout the protocols. Primary T cells (StemCell cat#70024, lot # 200172003C, Donor RG1010) were thawed and resuspended in complete media (RPMI 1640/10% FBS Corning/gentamicin, Gibco) containing anti-CD28 (BD Biosciences 555725, CD28.27, 2.5 μg/mL) and interleukin-2 (R&D systems 202-IL/CF, 2 ng/mL) and expanded by being placed in flasks precoated overnight with anti-CD3 (BD Biosciences cat#555329, UCHT1, 5 μg/mL). T cell expansion was allowed to proceed until day 4. On day 3, the tumor cells were resuspended in complete media and seeded at 5000 cells/well in 96-well black, clear bottom plates (PerkinElmer) overnight at 5% CO<sub>2</sub> and 37°C. On day 4, 50 K expanded T cells/well were washed

2× in complete media and added to each well with HCT-116 and A375 tumor cells. Prior to the addition of the T cells, media was removed and the rTCR/anti-CD3 BFPs were titrated into the plates with tumor cells starting at a final concentration of 200 nM with 1:10 serial dilutions ending at 14:00 (0.002 nM). The T cell/tumor cell/BFP (or MAbs) mixtures were allowed to incubate for 24 hours, then the plates were washed gently (twice) with serum free RPMI 1640. Finally, to determine the amount of tumor cells alive at the end of the incubations, 100 μL RPMI 1640 and 100 μL Cell Titer Glo reagent (Promega) were added, mixed for 2 min on a shaker (slowly) and incubated for 10 min in the dark. Lastly, the luminescence was read on an EnVision V.2130 multilabel reader.

## RESULTS

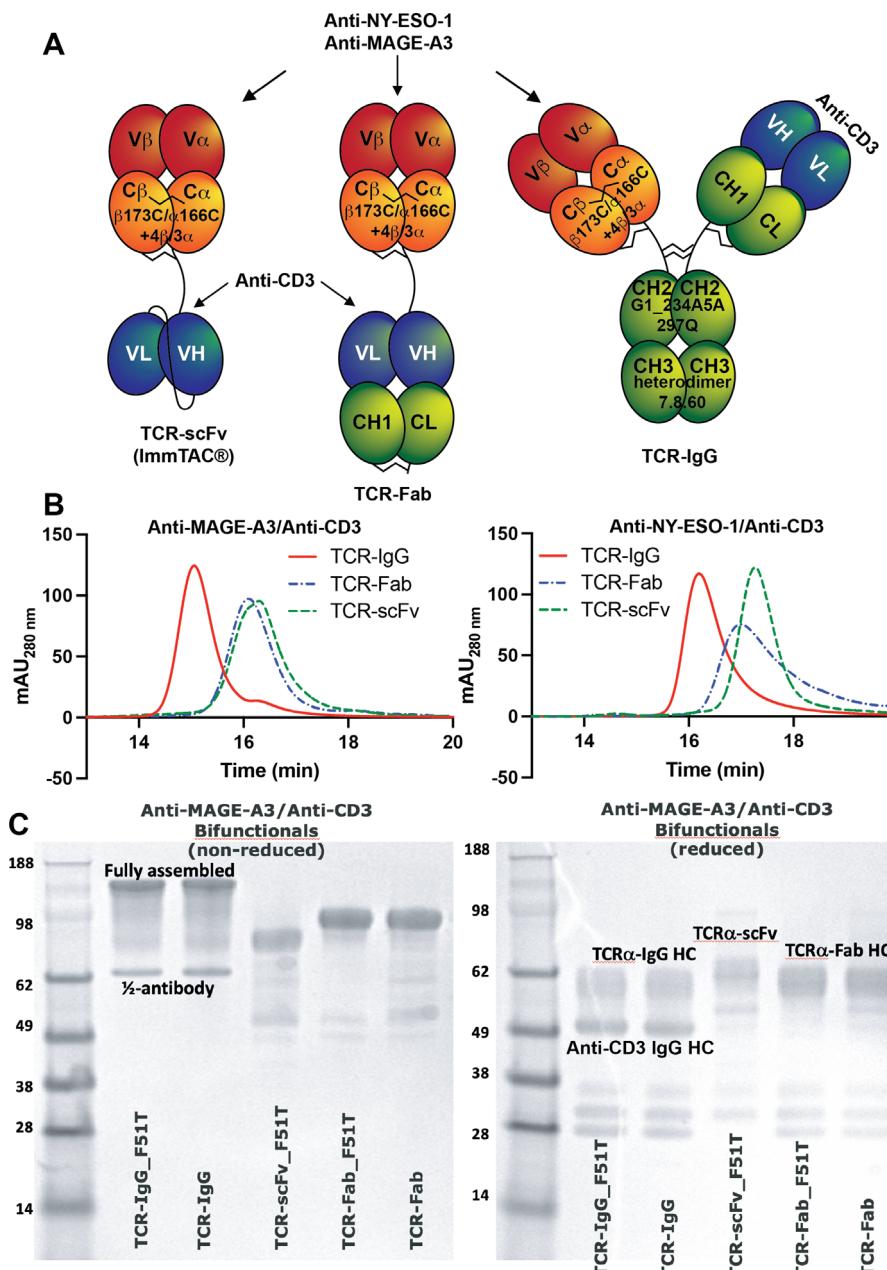
### Expression and characterization of TCR-scFv, TCR-Fab, and TCR-IgG BFPs

Two well-characterized rTCRs were chosen for the study of various rTCR/anti-CD3 BFP geometries. The first was 1G4\_113, an affinity-matured clone of the 1G4 anti-NY-ESO-1/HLA-A2 TCR,<sup>17</sup> and the second was an affinity-matured anti-MAGE-A3/HLA-A1 TCR.<sup>14</sup> Initially, three rTCR/anti-CD3 geometries were assessed: TCR-scFv, TCR-Fab, and TCR-IgG (figure 1A). The anti-CD3 moiety was SP34 constructed as a VH(GGGGS)<sub>4</sub>VL (scFv), a chimeric mouse Fv fused to human CH1/Cλ (Fab), or an effector-less IgG1 constant region (IgG1\_L234A\_L235A\_N297Q).<sup>11</sup> All TCRs contained stabilized Cα/Cβ designs enabling antibody-like transient expression levels, assembly, and secretion via transient expression in CHO cells.<sup>11</sup> TCR-IgG heterodimerization was achieved using a previously described IgG-CH3 domain heterodimerization design denoted 7.8.60.<sup>11</sup>

The rTCR/anti-CD3 BFPs expressed and assembled well. Both the NY-ESO-1 and MAGE-A3 BFPs were mono-disperse by analytical SEC with the TCR-IgGs, TCR-Fabs, and TCR-scFvs eluting in order based on their molecular weight (figure 1B). By SDS-PAGE, the MAGE-A3-directed BFPs displayed the expected molecular weights for full assembly under non-reducing conditions and the expected individual chains under reducing conditions (figure 1C). SDS-PAGE analysis of the 1G4\_113 NY-ESO-1-directed TCR-Fab and TCR-scFv looked similar to their MAGE-A3-directed counterparts (online supplemental figure S1)

### Functional binding of rTCR/anti-CD3 BFPs

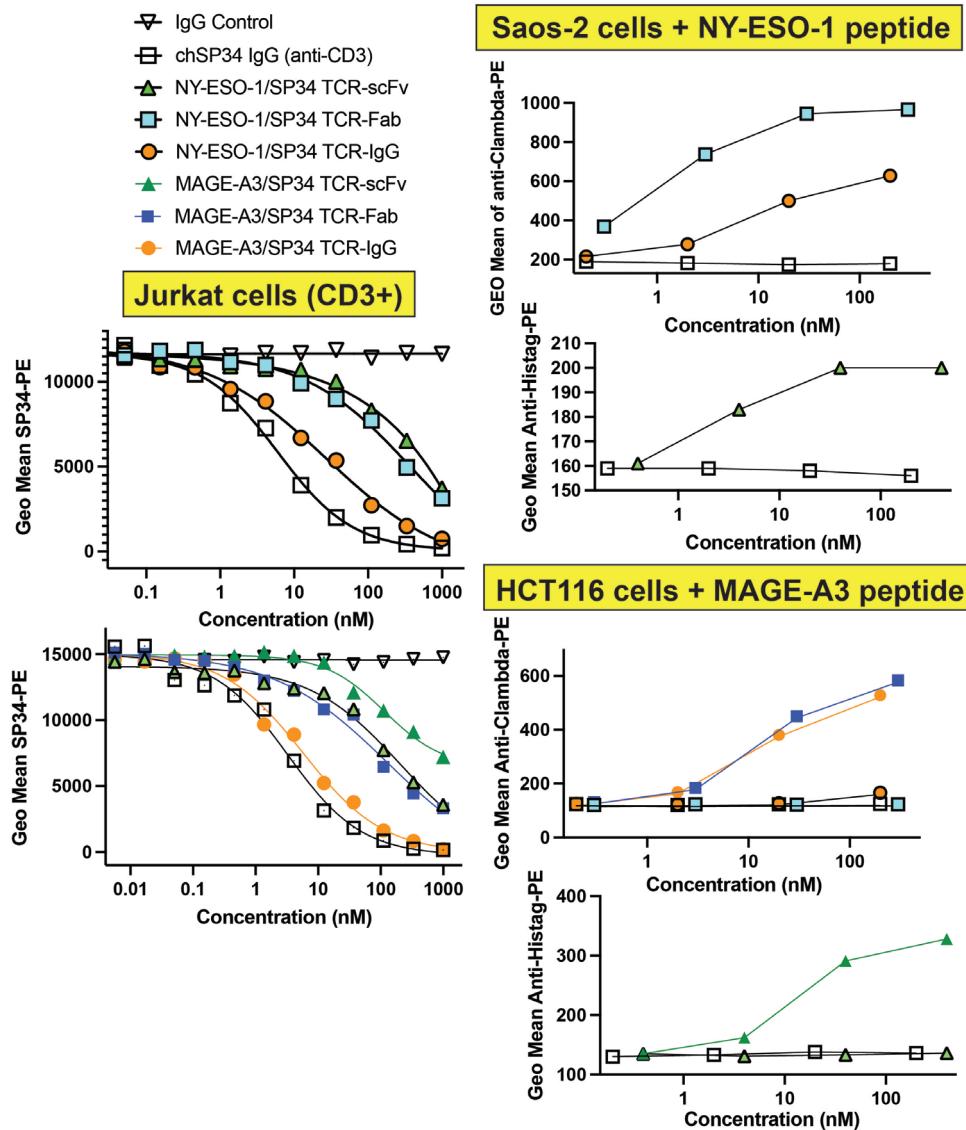
The ability of the rTCR/anti-CD3 BFPs to bind their target antigens was assessed using flow cytometry. First, binding to CD3 antigen was assessed using the Jurkat T cell leukemia line. All the BFPs bound Jurkat cells (online supplemental figures S2 and S3). We found the most quantitative method for comparing the CD3 binding affinity of the BFPs to CD3-expressing cells was to perform a competitive flow cytometry assay that compares the ability of each BFP to compete with a set concentration of PE-conjugated SP34



**Figure 1** Architecture and biophysical properties of rTCR/anti-CD3 BFPs. (A) Schematic diagrams of the TCR-scFv, TCR-Fab, and TCR-IgG BFPs. VH and VL are the anti-CD3 antibody heavy and light chain variable domain fragments, respectively. (B) Analytical size exclusion chromatography of the MAGE-A3-directed (left) and the NY-ESO-1-directed (right) rTCR/anti-CD3 BFPs directly after protein A or Ni<sup>2+</sup>-NTA affinity chromatography. (C) MAGE-A3-directed BFPs analyzed by SDS-PAGE under non-reducing (left) and reducing (right) conditions. BFP, bifunctional protein; rTCR, recombinant T cell receptor-based therapeutic; SDS-PAGE, sodium dodecyl sulfate–polyacrylamide gel electrophoresis; TCR, T cell receptor.

(figure 2 and online supplemental table S1).<sup>18</sup> Unlabeled, bivalent SP34 was titrated as a control and demonstrated a single-digit nanomolar IC<sub>50</sub>. Both of the monovalent TCR-IgG BFPs demonstrated roughly 10-fold weaker potency (figure 2 and online supplemental table S1) similar to what has been observed with fully antibody-based monovalent CD3 BsAbs.<sup>18</sup> Both the TCR-scFv or TCR-Fab BFPs displayed ~10fold weaker potency compared with the TCR-IgG BFPs likely due to the N-terminal fusion of the TCR hindering the kinetics of CD3 engagement (figure 2 and online supplemental table S1).

Next, binding of the BFP rTCR moieties to cell lines displaying specific MHC/peptide complexes was assessed. NY-ESO-1 binding was performed on HLA-A2<sup>High</sup> Saos-2 osteosarcoma cells preloaded with the NY-ESO-1<sub>157-165</sub> SLLMWITQC peptide. All three NY-ESO-1-directed BFPs bound the Saos-2 cell line with roughly similar EC<sub>50</sub>s though the TCR-IgG BFP did not saturate at the same level as the TCR-Fab and its EC<sub>50</sub> was slightly weaker (online supplemental figures S4 and S5, figure 2, online supplemental table S1). All three MAGE-A3-directed BFPs bound HLA-A1<sup>High</sup> HCT116 colon carcinoma cells



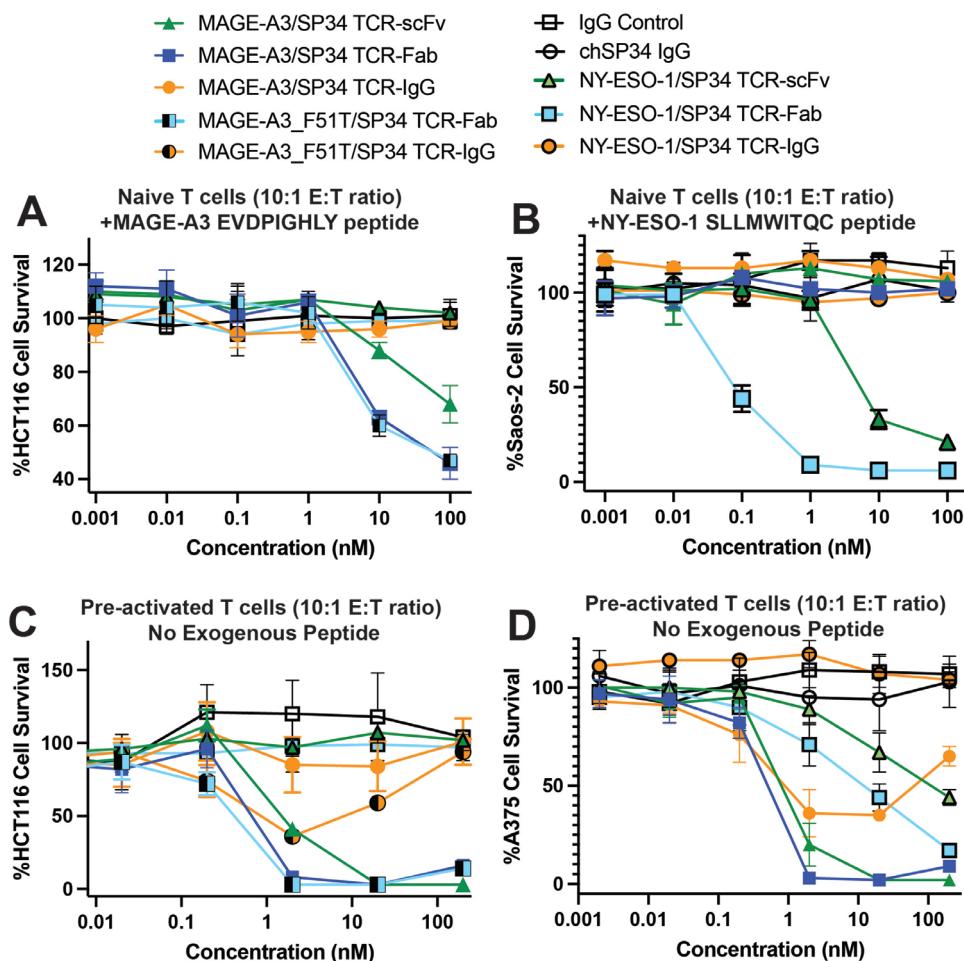
**Figure 2** Cell-binding capability of rTCR/anti-CD3 BFPs to both CD3-expressing Jurkat cells (left), NY-ESO-1<sub>157–165</sub> SLLMWITQC-labeled SaOS-2 (NY-ESO-1) cells, or MAGE-A3<sub>168–176</sub> EVDPIGHLY-labeled HCT116 cells. Jurkat cell potency was assessed by competition with anti-CD3 SP34-PE, while SaOS-2 and HCT116 binding potency was assessed using an anti-human lambda-PE secondary (TCR-IgG and TCR-Fab) or anti-Histag-PE secondary (TCR-scFv). BFP, bifunctional protein; PE, phycoerythrin; rTCR, recombinant T cell receptor-based therapeutic; TCR, T cell receptor.

preloaded with the MAGE-A3<sub>168–176</sub> EVDPIGHLY peptide with near identical EC<sub>50</sub>s (figure 2, online supplemental table S1). No non-specific binding of NY-ESO-1-directed BFPs to the HCT116 line preloaded with the MAGE-A3 peptide occurred, highlighting the HLA/peptide specificity of the BFPs (figure 2). An affinity matured version of the MAGE-A3-directed TCR-IgG and TCR-Fabs (F51T)<sup>14</sup> demonstrated only a 2-fold EC<sub>50</sub> improvement; therefore, we focused subsequent efforts on the original MAGE-A3 TCR-directed BFPs for functional analyses. Overall, both sets of rTCR/anti-CD3 BFPs demonstrated strong cell surface MHC/peptide and CD3ε binding capabilities (online supplemental table S1).

#### T cell-engaging activity of rTCR/anti-CD3 BFPs

The BFPs were assessed for their ability to redirect T cells to kill tumor cells displaying the correct HLA-peptide

complex. First, the BFPs directed toward MAGE-A3 were mixed with MAGE-A3-peptide-loaded HCT116 cells and naïve (rested/non-activated) T cells. The TCR-Fab and TCR-scFv were able to stimulate and redirect the T cells to kill HCT116 in a dose-dependent manner (figure 3A), while the TCR-IgG was inactive. Interestingly, the potency of the TCR-Fab was significantly higher than that of the TCR-scFv. The NY-ESO-1-directed BFPs had no activity against HCT116 cells preloaded with MAGE-A3 peptide. When applied to HLA-A2<sup>High</sup> Saos-2 cells preloaded with NY-ESO-1 peptide, the NY-ESO-1 TCR-Fab and TCR-scFv BFPs could stimulate and redirect T cells to kill the Saos-2 cells (figure 3B). As with the MAGE-A3-directed TCR-IgG, the NY-ESO-1 TCR-IgG was inactive. Overall, both BFP datasets demonstrate that the TCR-Fab and TCR-scFv moieties can stimulate and redirect T cells towards tumor cells



**Figure 3** Comparative T cell redirected killing of HLA/peptide<sup>+</sup> tumor cell lines via the rTCR/anti-CD3 BFPs. BFP-induced redirection of naïve T cells to kill HLA-A1<sup>+</sup> HCT116 colorectal cancer cells (A) or HLA-A2<sup>+</sup> Saos-2 osteosarcoma cells (B) preincubated with MAGE-A3<sub>168–176</sub> EVDPIGHLY or NY-ESO-1<sub>157–165</sub> SLLMWITQC peptides, respectively. BFP-induced killing of HLA-A1<sup>+</sup>/MAGE-A3<sup>+</sup>/HLA-A2<sup>+</sup>/NY-ESO-1<sup>-</sup> HCT116 cells (C) or HLA-A1<sup>+</sup>/MAGE-A3<sup>+</sup>/HLA-A2<sup>+</sup>/NY-ESO-1<sup>-</sup> A375 malignant melanoma cells (D) using preactivated T cells. BFP, bifunctional protein; E:T, effector cell to tumor cell ratio; rTCR, recombinant T cell receptor-based therapeutic; TCR, T cell receptor.

displaying their target antigens, while the TCR-IgG format is either inactive or much less active. In both cases, the TCR-Fab format was more potent than the TCR-scFv format.

Next, we evaluated the ability of the MAGE-A3 and NY-ESO-1 BFPs to redirect T cells to kill tumor cells expressing endogenous levels of antigen/HLA. The first cancer cell line evaluated was HCT116, which endogenously expresses the MAGE-A3/HLA-A1, but does not express NY-ESO-1.<sup>19 20</sup> The MAGE-A3-directed TCR-Fab and TCR-scFv could induce T cell-mediated killing of HCT116 tumor cells while the NY-ESO-1-directed BFPs had no activity. Again, the TCR-Fab was more potent than the TCR-scFv. The MAGE-A3-directed TCR-IgG was inactive; however, the higher affinity F51T TCR-IgG BFP variant<sup>14</sup> displayed activity that diminished in a bell-shaped profile at higher concentrations (figure 3C). When the MAGE-A3- and NY-ESO-1-directed BFPs were applied to A375 cells, which endogenously express both MAGE-A3/HLA-A1 and NY-ESO-1/HLA-A2, both sets of TCR-Fabs and TCR-scFvs induced activated T cells to kill the A375 cells (figure 3D). Overall, the MAGE-A3 BFPs

more potently induced T-cell killing of the A375 cells compared with the NY-ESO-1 BFPs (figure 3D). Again, the TCR-Fabs were slightly more potent than the TCR-scFvs (figure 3D). The MAGE-A3 TCR-IgG did redirect T cells with high potency to kill A375 cells, but killing only reached 60% and demonstrated a bell-shaped activity curve similar to the data observed for the F51T MAGE-A3 TCR-IgG on HCT116 cells pre-loaded with the MAGE-A3 peptide (figure 3C,D).

#### Expanded rTCR/CD3 BFP geometry/valency evaluation

The scope of rTCR/anti-CD3 BFP analyses was expanded to include the impact of rTCR affinity and additional BFP geometries. The geometries chosen for investigation were inspired by the most prevalent T cell-engaging BsAb geometries,<sup>10</sup> and our observation that T cell-engaging BsAbs with tandem formats comprising an N-terminal TSA binding domain followed by an anti-CD3 domain showed superior potency over IgG-format bifunctionals in many of our assays.<sup>16 18</sup> Also, many T cell-engaging BsAbs are moving toward 2:1 formats (2 TSA moieties:

1CD3 moiety) that use modest affinity and avidity to achieve improved specific binding to tumor cells over-expressing TSA and to avoid normal tissues expressing lower TSA levels.<sup>10</sup> Lastly, formats with an antibody-Fc to impart antibody-like pharmacokinetics were evaluated. Given the generally similar results obtained comparing the TCR-Fab, TCR-scFv, and TCR-IgG formats directed to MAGE-A3/HLA-A1 and NY-ESO-1/HLA-A2, we decided to focus on the NY-ESO-1 BFPs allowing the construction and testing of more formats.

We first investigated the requirement for high affinity. Previously, we found that rTCR/anti-CD3 BFPs with TCR affinities at or below 1 nM did not show significant potency differences *in vitro*.<sup>11</sup> Therefore, we investigated the impact of reducing the affinity by one or two orders of magnitude. Li and coworkers published a study that included anti-NY-ESO-1 rTCR sequences ranging from low pM to  $\mu$ M  $K_d$ s for HLA-A2/peptide.<sup>17</sup> TCR-Fab BFPs with HLA-A2/NY-ESO-1 peptide binding affinities ranging from 1 nM to 200 nM were generated (1G4\_122~1 nM, 1G4\_006~10 nM, and 1G4\_033~200 nM<sup>17</sup>) and tested for redirected lysis activity on Saos-2 cells loaded with NY-ESO-1 peptide. While the 1 nM TCR-Fab demonstrated redirected lysis potency similar to the 25 pM TCR-Fab, both the 10 nM and 200 nM  $K_d$  TCR-Fab BFPs had highly attenuated activity and potency (>2 orders of magnitude, figure 4A), suggesting high HLA/peptide affinity is crucial for the activity and potency of these molecules in agreement with what has been described previously.<sup>7</sup>

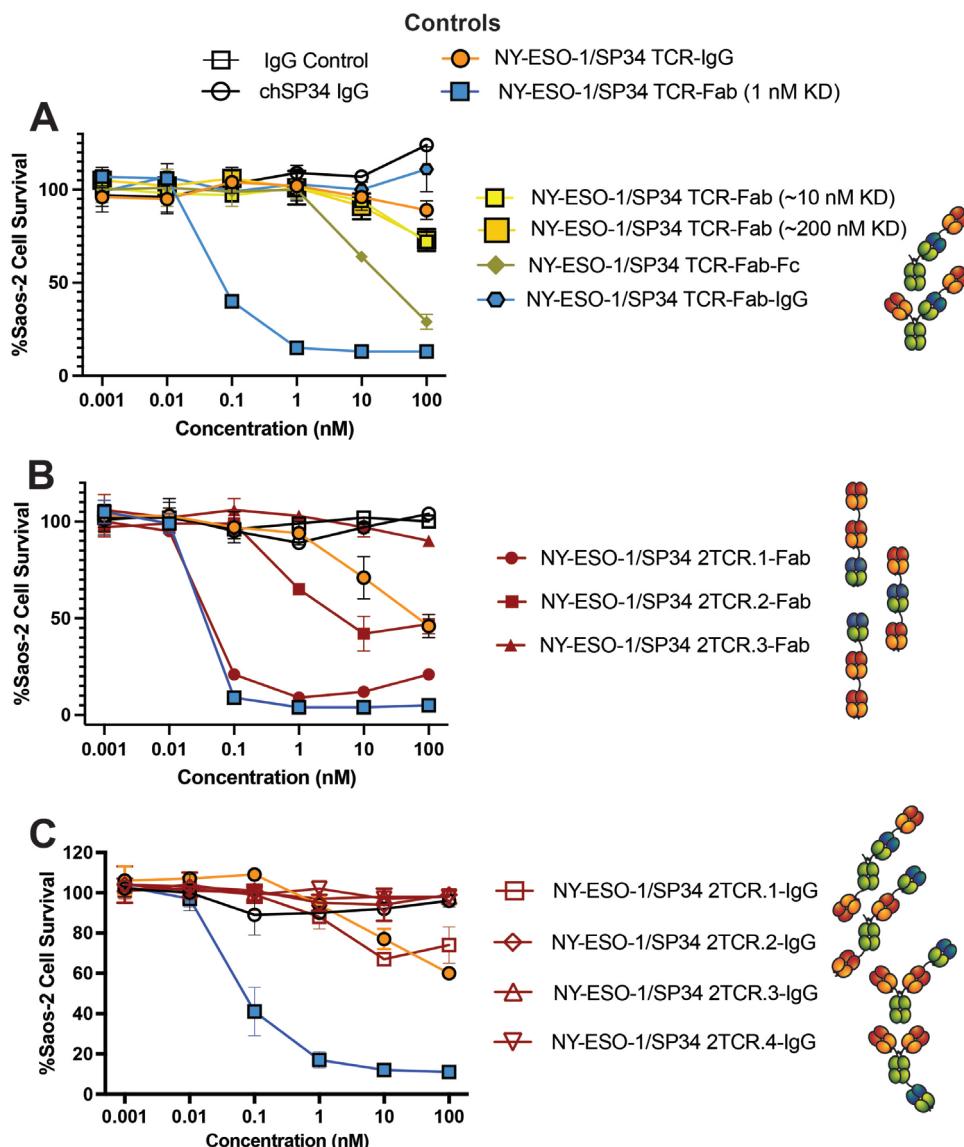
Next, the geometric aspects of the rTCR/anti-CD3 moieties were evaluated in more depth. First, we evaluated modifications of the original TCR-IgG and TCR-Fab BFPs (using the 1G4\_122 anti-NY-ESO-1 TCR with a 1 nM  $K_d$ ) to assess whether moving to the tandem format (TCR-Fab-Fc) or a 2:1 tandem format (TCR-Fab-IgG) would improve the potency of an Fc-containing BFP. The tandem format, TCR-Fab-Fc, did lead to a modest increase in potency over the TCR-IgG format, though still two orders of magnitude weaker than the tandem TCR-Fab format lacking an IgG-Fc (figure 4A). Diversifying the linker length from a (GGGGS)<sub>3</sub> to (GGGGS)<sub>4</sub> or (GGGGS)<sub>5</sub> linker did not impact these results. Adding an additional rTCR arm within the TCR-Fab-IgG format eradicated all activity.

Subsequently, we evaluated multiple 2:1 rTCR:anti-CD3 tandem formats. Adding a second rTCR to the N-terminus of the TCR-Fab (denoted 2TCR.1-Fab) did not improve potency over the original TCR-Fab format and appeared to induce an activity loss at higher BFP concentrations similar to what was observed for the MAGE-A3 TCR-IgG BFP (figure 4B). Moving the anti-CD3 Fab moiety to the central or N-terminal positions (denoted 2TCR.2-Fab and 2TCR.3-Fab, respectively) each led to weaker potency and activity with the N-terminal anti-CD3 Fab format having no activity (figure 4B). Lastly, we evaluated additional IgG-Fc containing 2:1 formats (denoted 2TCR-IgGs). These formats included a variety of contexts investigating the positioning of the second rTCR or the anti-CD3 Fab moieties.

In short, none of these formats were more active than the TCR-IgG format, and most were completely inactive in the assay (figure 4C). Thus, adding a second rTCR to the original constructs did not result in improvements in potency or activity over the original TCR-Fab or TCR-IgG formats and in most cases reduced activity.

## DISCUSSION

Unlike T cell-engaging BsAbs, the parameters required for potent rTCR/anti-CD3 T cell redirection have been poorly characterized. Still, the anti-gp100 rTCR/anti-CD3 ImmTAC is the first T-cell engager to show strong clinical activity in a solid tumor indication, uveal melanoma, in a large phase III trial,<sup>12</sup> demonstrating the utility of the rTCR-based BFPs and warranting further exploration of rTCR-based T cell engagers. Given the conserved nature of TCR-HLA-peptide interactions, we hypothesized that the parameters dictating activity are likely generalizable, unlike BsAb-based T-cell engagers whose activity depends on both BsAb geometry and the membrane proximity of the targeted epitope.<sup>10</sup> In agreement with this hypothesis, the impact of BFP geometry on the activity and potency was uniform across both the MAGE-A3/HLA-A1- and NY-ESO-1/HLA-A2-directed BFPs. While there is significant diversity in TCR complementarity determining loops and how they engage specific HLA-bound antigenic peptides, the overall angle of approach of the rTCR to its HLA-epitope is a general feature required for the production of a functional immune synapse.<sup>21</sup> To our surprise, both sets of TCR-Fab BFPs had greater potency than the TCR-scFv format, while the activity of the TCR-IgG (antibody-like) format was substantially weaker. We hypothesize that the larger and less flexible TCR-IgG has difficulty forming the tight immune synapse necessary to induce TCR conformational changes, and TCR cross-linking, and possibly the exclusion of large phosphatase receptors as described in the kinetic-segregation model (online supplemental figure S6).<sup>22</sup> Also, in some circumstances, the TCR-IgG and 2TCR.1-Fab (ie, the fusion of 2 rTCRs and one anti-CD3 Fab) formats demonstrated bell-shaped activity profiles with a loss of activity at higher concentrations. This may result from molecular crowding and expansion of the synapse between T cell and tumor cell with these larger molecules. All formats containing an IgG-Fc demonstrated poor potency or a total lack of activity. It is possible smaller modifications, for example, acylation or nanobody fusion, could direct these molecules to human serum albumin for improved pharmacokinetics with binding kinetics that allow for controlled release once the molecules enter the immune synapse. The use of diverse anti-CD3 binders with varied epitopes hypothetically could impact the general observations discovered here; however, the epitope space described for anti-CD3 binders is limited and structures determined of different anti-CD3 mAbs bound to CD3 $\epsilon$  have nearly identical overall epitopes.<sup>23 24</sup> Thus, it seems unlikely that varying the CD3 epitope would modify the geometrical



**Figure 4** Assessment of affinity and diverse architecture on T cell-redirection lysis activity of NY-ESO-1-directed rTCR/anti-CD3 BFPs. Impact of affinity or addition of an IgG-Fc with or without a second rTCR (A); addition of an additional TCR-binding arm to the tandem format in three separate orientations (B); or complex avid structures containing both an IgG-Fc and a second rTCR in various orientations (C) on the ability of the rTCR/anti-CD3 BFPs to redirect naïve T cells to kill HLA-A2<sup>+</sup> SaOS-2 cells labeled with NY-ESO-1<sub>157–165</sub> SLLMWITQC peptide. As controls, NY-ESO-1 TCR-Fab and TCR-IgG BFPs were used in every experiment. The differences in the redirected lysis potency of the control NY-ESO-1 TCR-Fab and TCR-IgG as well as other rTCR/anti-CD3 BFPs across different experiments can be attributed to the use of different T cell donors. Other than the exceptions described (A), all TCRs had a 1 nM  $K_D$  for the NY-ESO-1/HLA-A2 complex. BFP, bifunctional protein; rTCR, recombinant T cell receptor-based therapeutic; TCR, T cell receptor.

requirements of rTCR-based BFPs for T-cell redirection uncovered here.

Like BsAb T cell engagers, a complex relationship between HLA/peptide-targeting affinity and potency exists. There appeared to be a threshold near 1 nM for the NY-ESO-1-directed TCR-Fab BFP that enabled potent T cell redirection that was nearly identical regardless of an affinity difference between 1G4\_113 ( $K_D$  ~25 pM) and 1G4\_122 ( $K_D$  ~1 nM)<sup>11</sup> but which was rapidly degraded by reducing affinity further. Such a result is likely related to the avidity of molecular engagement at the immune synapse. Recently, a diabody-based TCR-mimic BsAb

(~50 kDa) was shown to have antitumor activity against a common mutant of the tumor suppressor p53 protein, mutants of oncogenic Ras enzyme, and other oncogenes.<sup>25</sup> The diabody directed to mutant TP53 had an affinity of 86 nM and was still capable of potent T cell redirection.<sup>25</sup> One possibility for this potency discrepancy is the known high potency of the UCHT1 anti-CD3 with stable binding to the T cell surface, allowing high avidity display of the anti-TP53 portion of their molecule.<sup>26</sup> High-affinity binding to CD3 $\epsilon$ , however, is likely to result in general T cell activation, and numerous studies have shown that low-affinity CD3 $\epsilon$  binding may facilitate less

systemic cytokine release and improved targeting of these compounds to the tissue sites harboring the epitopes of interest.<sup>18 27 28</sup>

We show that the TCR-Fab format had the best potency. This format has not been widely adopted likely because it is less amenable to production in bacterial systems, which has been the standard for rTCR molecules. An intrinsically unfolded CH1 domain<sup>29</sup> and likely issues with three chain oxidative folding would likely preclude bacterial production of a TCR-Fab. Soluble TCR production in mammalian cells has also been fraught with secretion and assembly issues and strong glycosylation, which can result in binding to lectin-containing receptors such as the asialoglycan receptor in the liver or DCSIGN on dendritic cells impacting their pharmacokinetics, trafficking, and immunogenicity.<sup>30</sup> However, design of intrinsically stable rTCRs, as described recently,<sup>11</sup> resulted in significant decreases in glycosylation with good assembly and expression from mammalian cells. Thus, expression of a stabilized TCR-Fab format using mammalian expression systems in the future may significantly improve both the function and production of rTCR/anti-CD3 BFPs for therapeutic use.

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