

1 The chimeric TAC receptor co-opts the T cell receptor yielding robust anti-tumor activity
2 without toxicity

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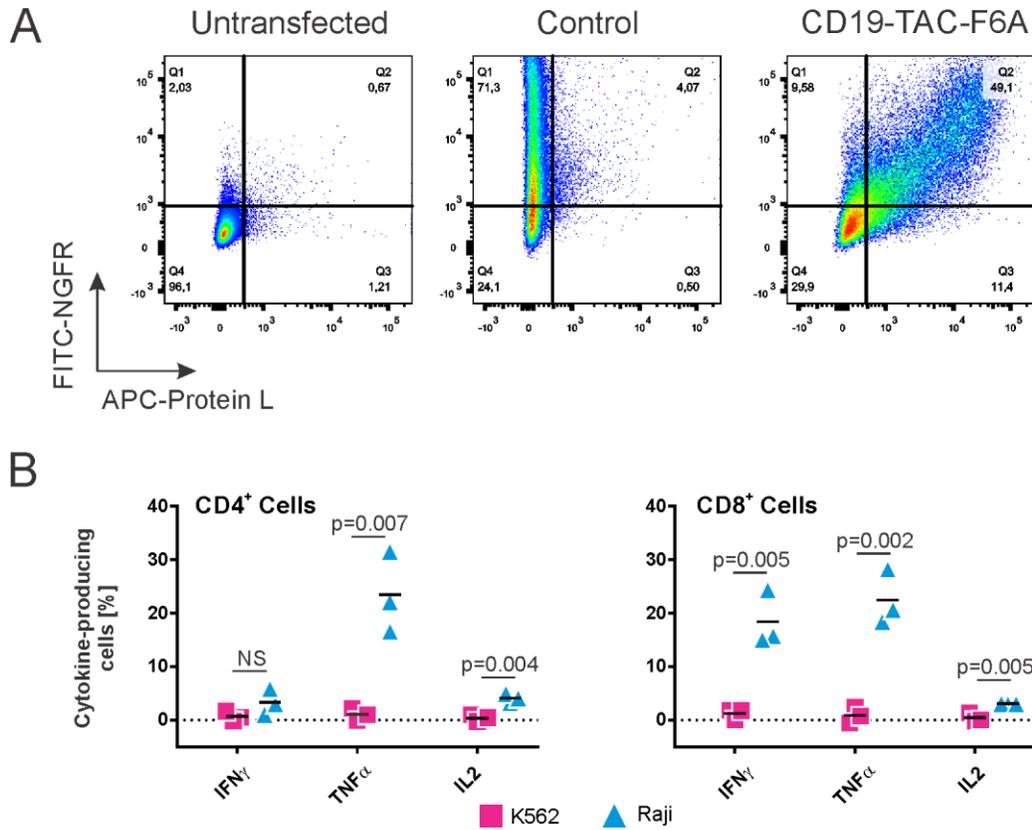
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6 **Supplementary Figures**

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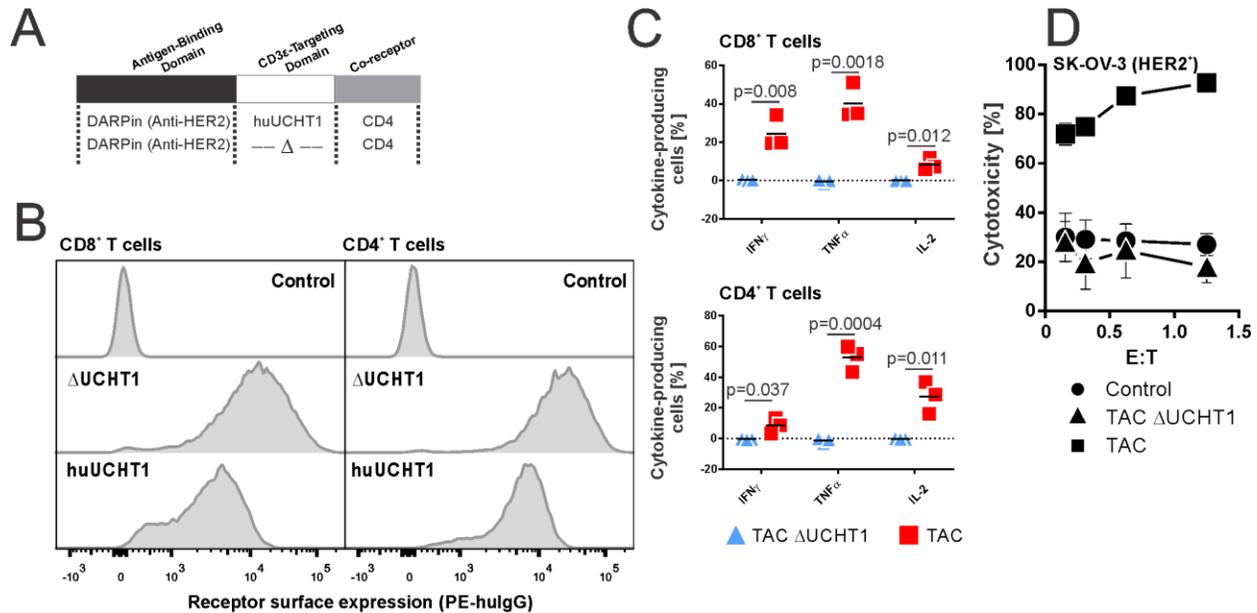
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10 Supplementary Figure 1:

11 **F6A scFv TAC can be detected by Protein L and F6A TAC engineered T cells show antigen**
 12 **specific reactivity.** **A.** Protein L binds the kappa light chain of scFv. HEK 293T cells were
 13 transfected with CD19-TAC-F6A, stained and analyzed for TAC and tNGFR expression by flow
 14 cytometry. For the gating strategy see Supplementary Fig. 16. **B.** CD19-TAC-cells were
 15 stimulated with antigen-positive Raji (triangle) or antigen-negative K562 (square) tumor cells,
 16 respectively. Data are presented as percent of CD4 or CD8 T cells producing cytokine. Lines
 17 represent data medians. Multiple t-test was used to determine significance. For the gating
 18 strategy see Supplementary Fig. 13B.

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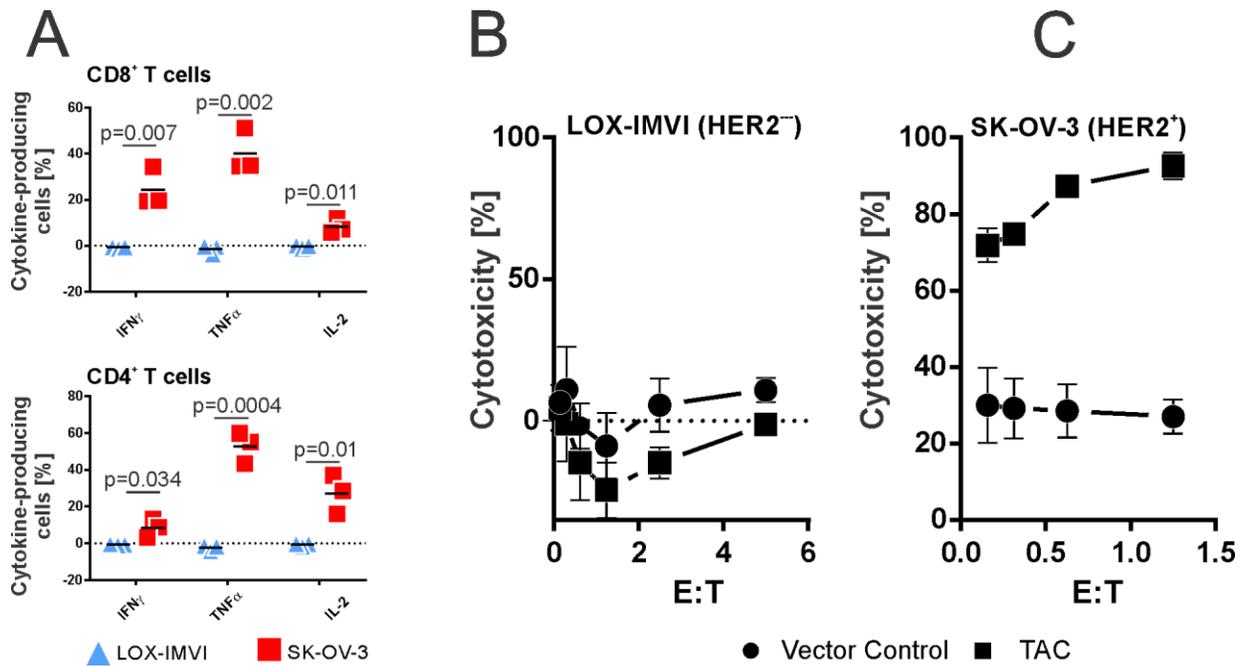
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24 Supplementary Figure 2:

25 **CD3-binding domain is required for TAC-engineered T cell function.** Full-length and
26 Δ UCHT1 TAC receptors (**A**) were expressed on the surface of primary human T cells (**B**).
27 Relative TAC surface expression is measured by flow cytometry. Cells were stained for CD4,
28 CD8, tNGFR and TAC (via its Myc Tag), and gated on either CD4⁺NGFR⁺ or CD8⁺NGFR⁺;
29 representative TAC expression data are presented as histograms. For the gating strategy see
30 Supplementary Fig. 13A. **C**. HER2-TAC-T cells (bearing huUCHT1 (square) or Δ UCHT1
31 (triangle)) are stimulated with antigen-positive SK-OV-3 tumor cells. Data are presented as
32 percent of CD4⁺ or CD8⁺ T cells producing cytokine. Lines represent data means. Multiple t-test
33 was used to determine significance. For the gating strategy see Supplementary Fig. 13B. **D**.
34 HER2-TAC-T cells (bearing huUCHT1 (square) or Δ UCHT1 (triangle)) and vector control T
35 cells (circles) are co-cultured with SK-OV-3 tumor cells to measure TAC-T cell-mediated
36 cytotoxicity. Data are from 3 independent experiments with 3 different donors; error bars show
37 standard deviation.

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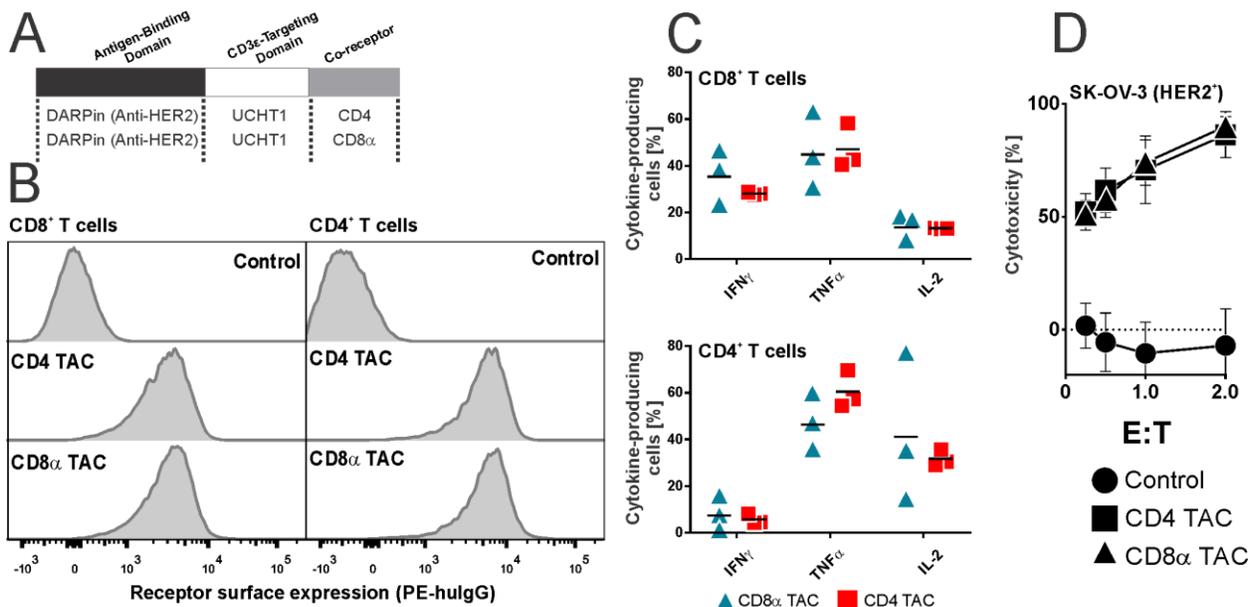
42 Supplementary Figure 3:

43 **TAC-T cells show no evidence of auto-activation in the absence of target antigen.** Data
44 originates from the same experiment as Supplemental Figure 2. **A.** HER2-TAC-T cells are
45 stimulated with antigen-positive SK-OV-3 (square) or antigen-negative LOX-IMVI (triangle)
46 tumor cells. Data are presented as percent of CD4⁺ or CD8⁺ T cells producing cytokine. Lines
47 represent data means. Multiple t-test was used to determine significance. For the gating strategy
48 see Supplementary Fig. 13B. **B-C.** HER2-TAC-T cells (bearing UCHT1; square) and vector
49 control T cells (circle) are co-cultured with LOX-IMVI or SK-OV-3 tumor cells to measure
50 TAC-T cell-mediated cytotoxicity. Data are from 3 independent experiments with 3 different
51 donors; error bars show standard deviation.

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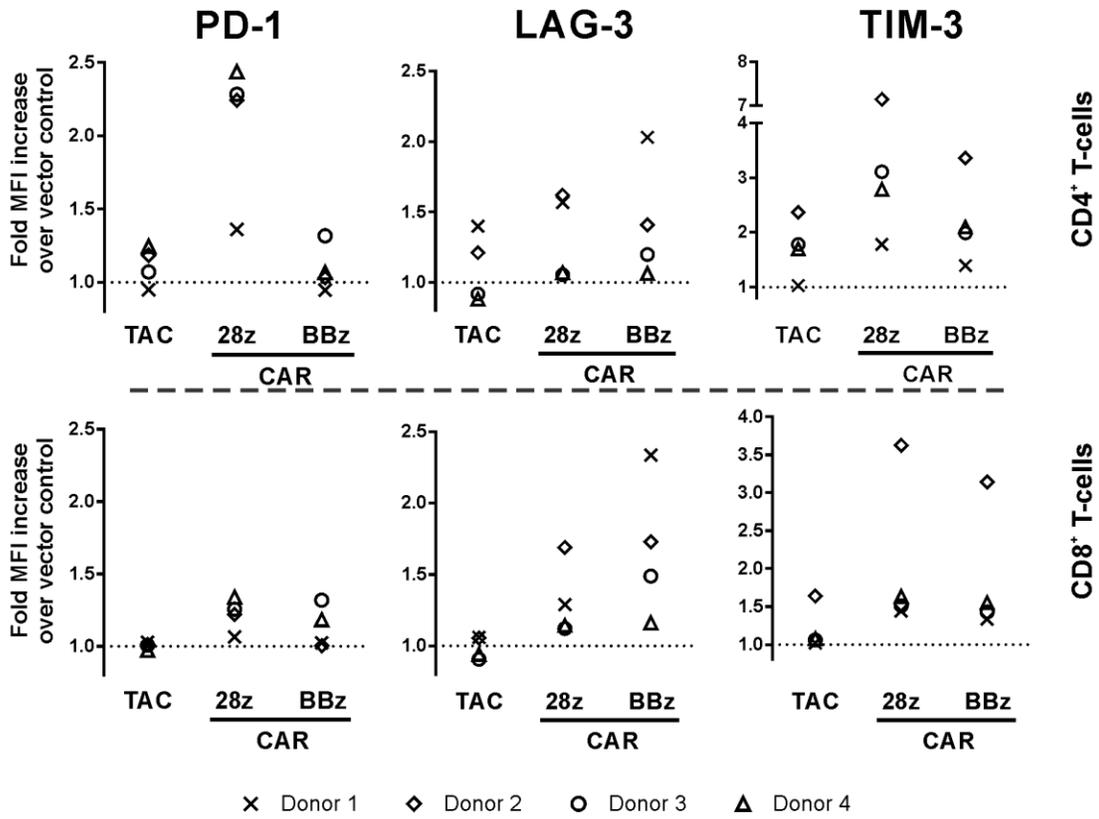
56 Supplementary Figure 4:

57 **Evaluation of cytosolic TAC domains.** **A.** Schematic representation of CD4 and CD8 α TAC
 58 constructs containing the anti-HER2 DARPin and UCHT1 CD3-binding domain. **B.** Cells were
 59 stained for CD4, CD8, tNGFR, and TAC expression, and gated on either CD4⁺NGFR⁺ or
 60 CD8⁺NGFR⁺; representative TAC expression data are presented as histograms. For the gating
 61 strategy see Supplementary Fig. 13A. **C.** Cytokine production by CD4 TAC- (square) and CD8 α
 62 TAC- (triangle) T cells stimulated by HER2⁺ SK-OV-3 tumor cells are shown. Lines represent
 63 data means. Multiple t-test was used to determine significance. For the gating strategy see
 64 Supplementary Fig. 13B. **D.** Cytotoxicity was measured by co-culturing HER2⁺ SK-OV-3 tumor
 65 cells with TAC- (CD4 co-receptor (squares) or CD8 α co-receptor (triangles)) or vector control
 66 (circles) T cells. Data are from 3 independent experiments with 3 different donors; error bars
 67 show standard deviation.

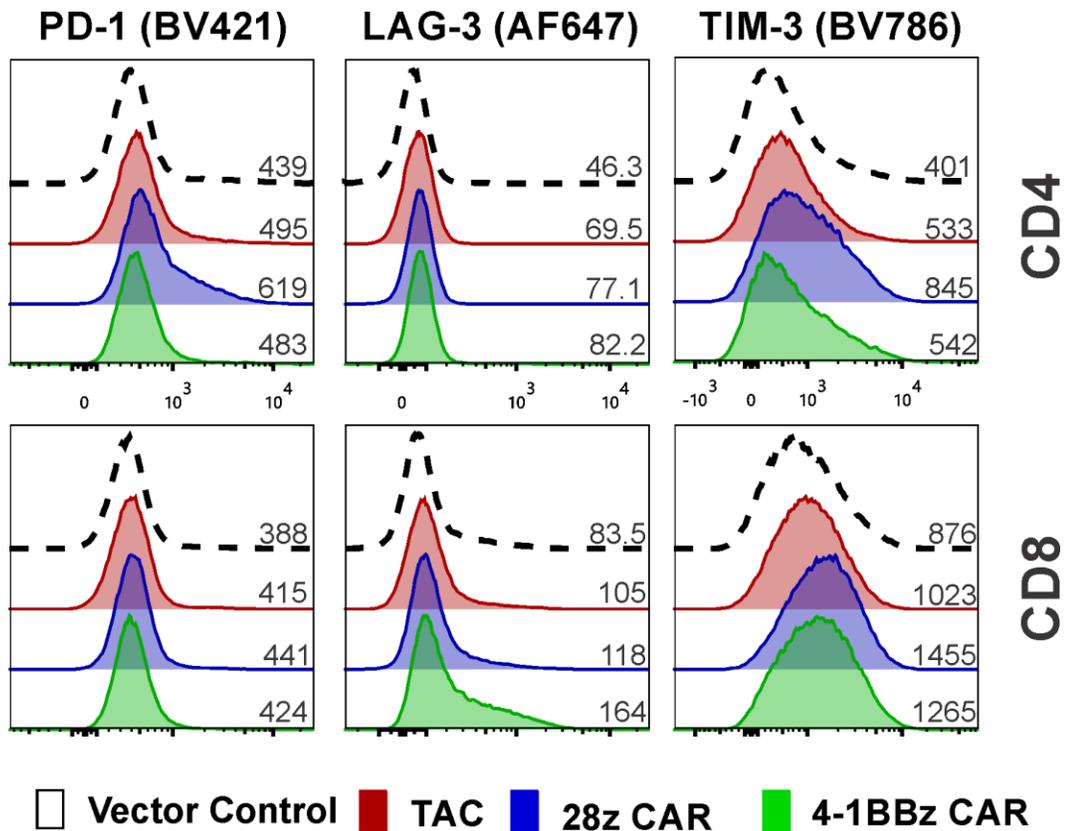
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A



B



71 Supplementary Figure 5:

72 **Relative expression of checkpoint receptors in CAR- and TAC-engineered T cells.** T cells

73 are transduced with HER2-TAC, anti-HER2 28ζ CAR or anti-HER2 BBζ CAR, stained for

74 surface marker expression, and analyzed by flow cytometry. **A.** Expression of checkpoint

75 receptors PD-1, LAG-3 and TIM-3. Populations are gated on either CD4⁺NGFR⁺ or

76 CD8⁺NGFR⁺. All data are normalized to vector control (vector only carrying tNGFR). Data is

77 derived from 4 individual donors across two separate experiments. Each donor is represented by

78 a unique symbol to visualize donor-to-donor variations. **B.** Representative histograms of HER2-

79 TAC (red), anti-HER2 28ζ CAR (blue) or anti-HER2 BBζ CAR (green). Populations are gated on

80 either CD4⁺NGFR⁺ or CD8⁺NGFR⁺. Representative, normalized histograms per marker are

81 shown. Values shown indicate mean fluorescence intensity. For both A and B the gating strategy

82 is shown in Supplementary Fig. 14.

83

85 Supplementary Figure 6:

86 **First-generation CAR-, second-generation CAR-, and TAC-T cells exhibit similar *in vitro***

87 **potency. A.** Schematics of TAC and first-generation CAR constructs. **B.** Comparison of cytokine

88 production from CD4⁺ or CD8⁺ TAC- (square) and first-generation CAR- (triangle) T cells when

89 stimulated with HER2⁺ OVCAR-3 tumor cells (minus cytokine production triggered by HER2⁻

90 LOX-IMVI tumor cells). Lines represent data means. **C.** Cytotoxicity of TAC- (square) and first-

91 generation CAR- (triangle) relative to vector control (circle) T cells against OVCAR-3 and

92 LOX-IMVI tumor cells. Data are from 3-4 independent experiments; error bars show standard

93 error of the mean. **D.** Schematics of the TAC and second-generation CD28-based CAR

94 constructs. **E.** Comparison of cytokine production from CD4⁺ or CD8⁺ TAC- (square) and

95 second-generation CAR- (inverted triangle) T cells when stimulated with OVCAR-3 (minus

96 cytokine production triggered by HER2⁻ LOX-IMVI tumor cells). Lines represent data means. **F.**

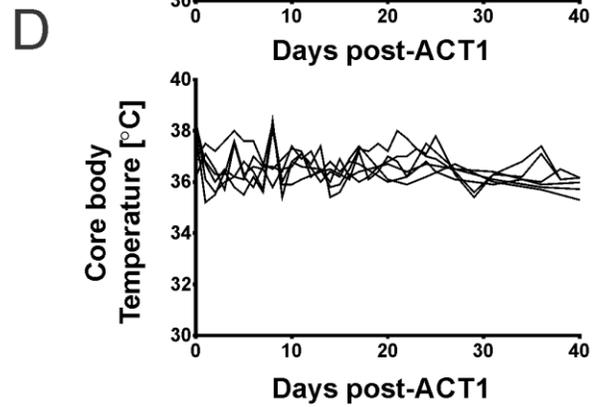
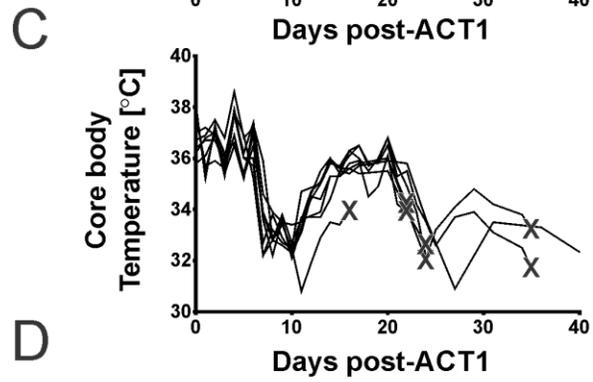
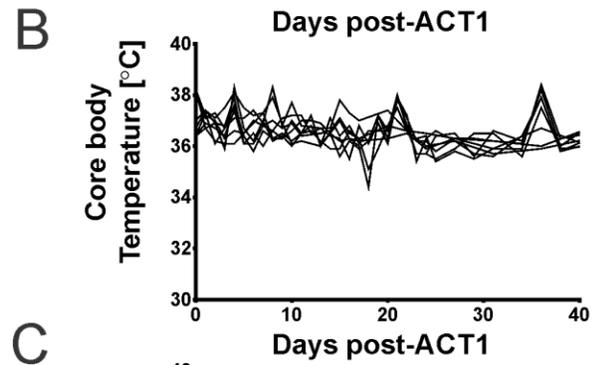
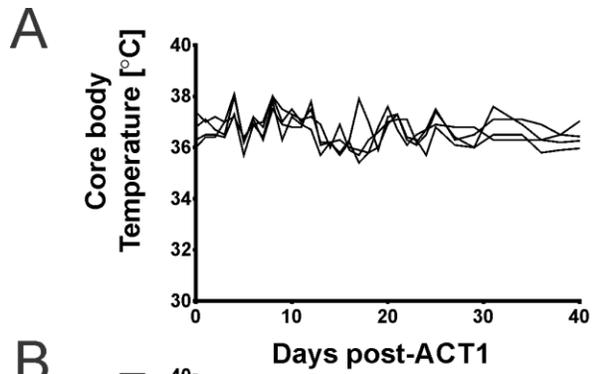
97 Cytotoxicity of TAC- (square) and second-generation CD28-based CAR- (inverted triangle)

98 relative to vector control (circle) T cells against LOX-IMVI and OVCAR-3 tumor cells. Data are

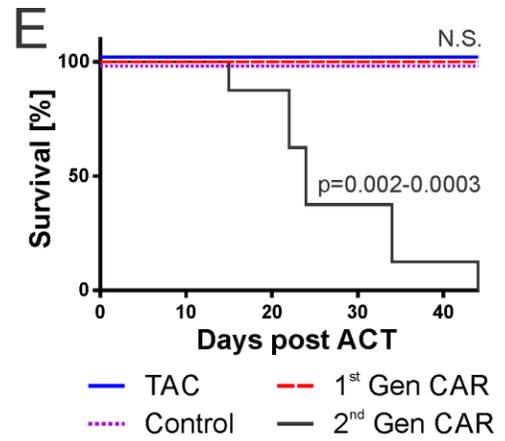
99 from 3 independent experiments; error bars show standard error of the mean. For B and E the

100 gating strategy is shown if Supplementary Fig. 13B.

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TAC
1st Gen CAR
2nd Gen CAR
Vector Control

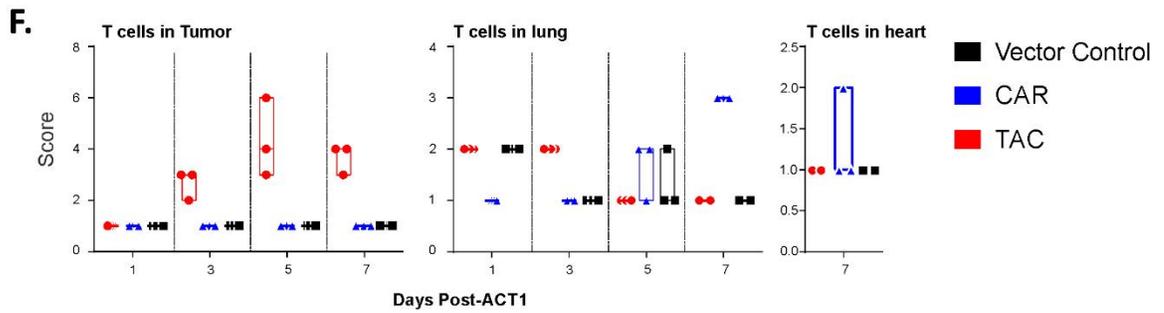
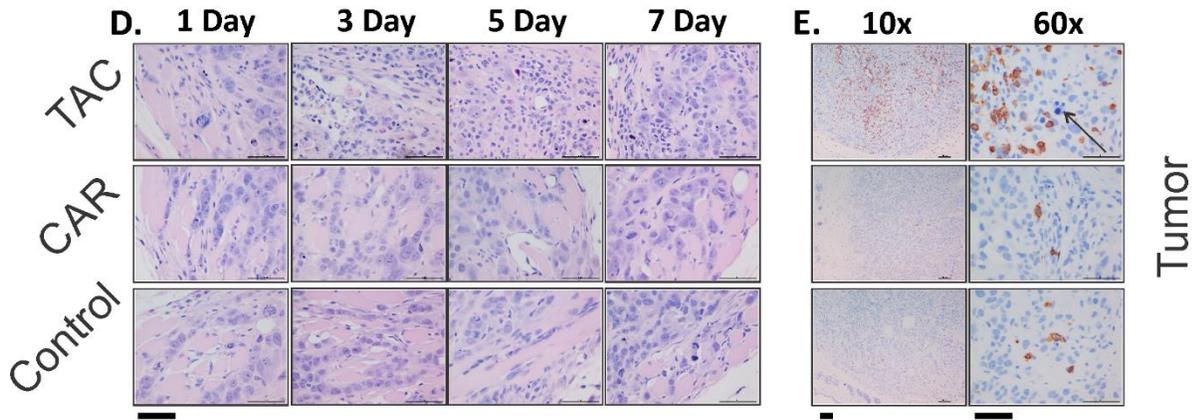
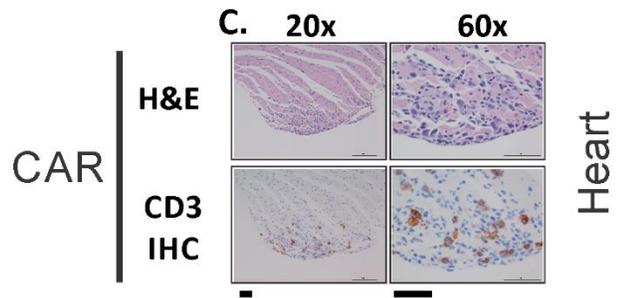
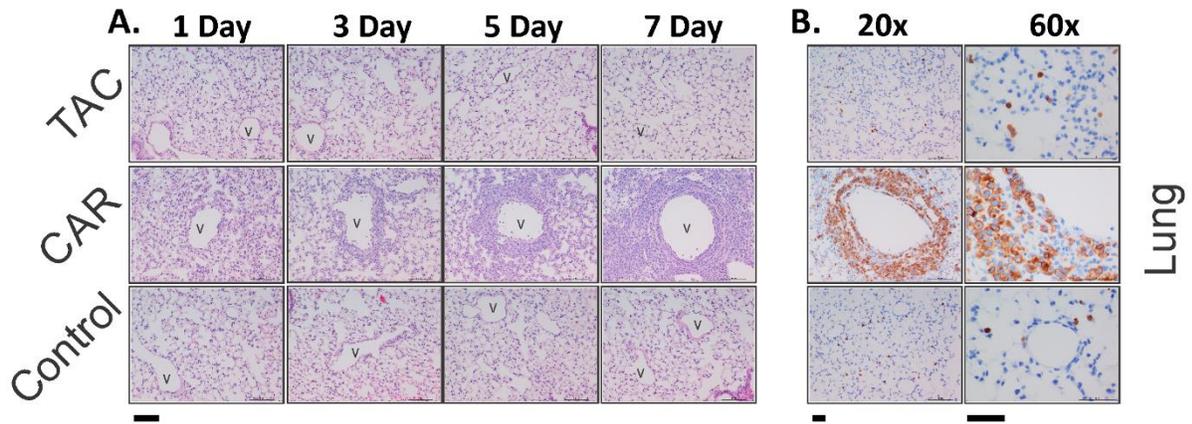


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108 Supplementary Figure 7:
109 **HER2-TAC-T cells demonstrate an enhanced safety profile over first- and second-**
110 **generation HER2-CAR-T cells *in vivo*.** OVCAR-3 tumor-bearing mice are treated with 2.0×10^6
111 HER2-TAC-T cells (**A**), first-generation anti-HER2 CAR-T cells (**B**), second-generation anti-
112 HER2 28 ζ CAR-T cells (**C**), or a matched total number of vector control T cells (**D**). Mice are
113 followed for change in core body temperature; each curve displays data from a single treated
114 mouse. When mice reach a toxicity-induced endpoint, this is indicated via an X. Data has been
115 replicated in an additional independent experiment. The change in core body temperature
116 between vector control, 1st gen CAR, and TAC was not significantly different. Significance was
117 determined via curve fitting and multiple t-test. Survival is shown in (**E**). Vector control (purple,
118 dotted), TAC (blue, solid) or first-generation CAR (red, dashes) treated mice showed no toxicity-
119 induced endpoints and were not significantly different. In contrast, second-generation 28 ζ CAR
120 (black, solid)-treated mice all reached toxicity-induced endpoint within 44 days. Mice were not
121 followed to tumor volume endpoint. Significance was determined via the Log Rank test.

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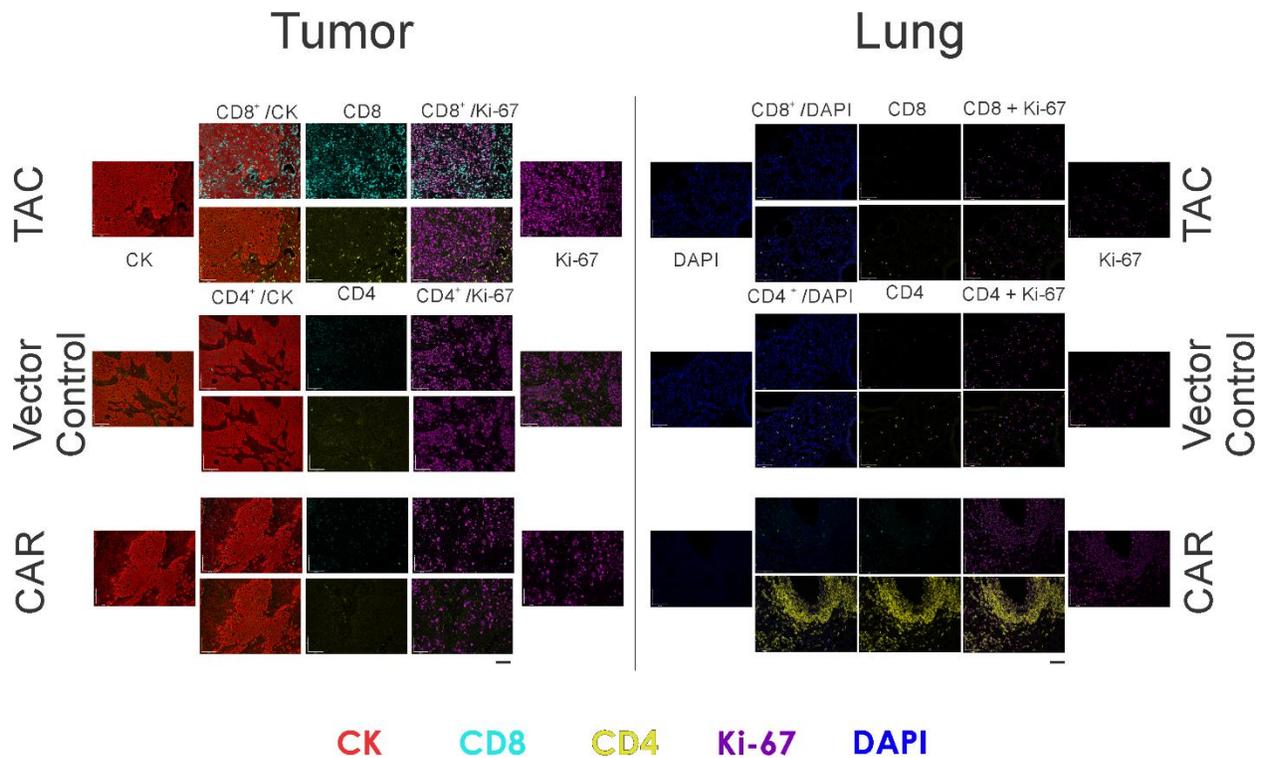
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127 Supplementary Figure 8:

128 **TAC- and CAR-T cells show differential patterns of localization *in vivo*.** OVCAR-3 tumor-
129 bearing mice were treated with 6.0×10^6 second-generation anti-HER2 28 ζ CAR- or HER2-TAC-
130 T cells, or a matched total number of vector control T cells. At 1, 3, 5, and 7 days post-ACT1
131 mice (n = 3 per treatment) were perfused and tissues were formalin fixed and paraffin embedded.
132 **A.** Timecourse of H&E (hematoxylin and eosin) stained lung sections at 20X magnification;
133 vasculature is indicated by a “v”. Scale bar indicates 100 μm . **B.** CD3 IHC staining of lung
134 sections at 7 days post-ACT1. Scale bar indicates 50 μm . **C.** H&E and CD3 IHC of cardiac tissue
135 at 7 days post-ACT1. Scale bar indicates 50 μm . **D.** Timecourse of H&E stained tumor sections
136 at 60X magnification. Scale bar indicates 50 μm . **E.** CD3 IHC staining of tumor sections at 7
137 days post-ACT1; arrow indicates a necrotic tumor cell. Scale bar indicates 50 μm . In all cases,
138 images are representative of observations in all mice (n = 3 each). **F.** CD3 IHC was performed
139 on heart, lung, and tumor tissue; T cell infiltrate was scored as % infiltrate based on tissue area in
140 10% intervals (score 1 = <1%, score 2 = 1-10%, score 3 = 10-20%, score 4 = 20-30%, score 5 =
141 30-40%, and score 6 = 40-50%). Data presented as an average score for n = 3 mice per time
142 point.

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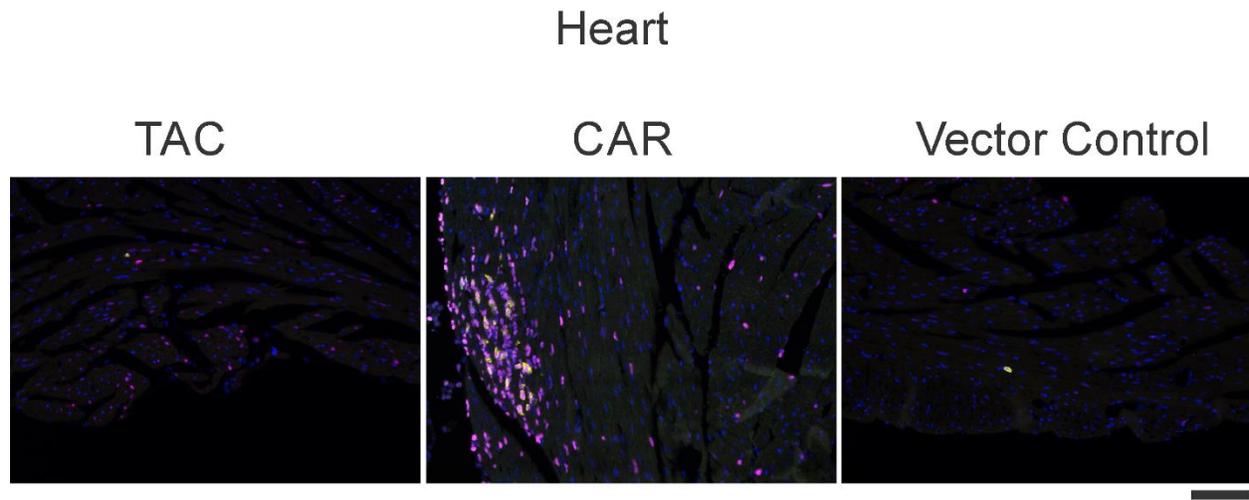
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146 Supplementary Figure 9:

147 **Single color and composite multicolor IHC tissue analysis.** OVCAR-3 tumor-bearing mice
 148 were treated with 6.0×10^6 second-generation anti-HER2 28 ζ CAR- or HER2-TAC-T cells, or a
 149 matched total number of vector control T cells. At 7 days post-ACT1 mice (n = 3 per treatment)
 150 were perfused and tissues were formalin fixed and paraffin embedded for subsequent multicolor
 151 IHC analysis (tumor or lung tissues were stained for human cytokeratin (CK, red), cellular
 152 proliferation marker Ki-67 (purple), CD8 (cyan), CD4 (yellow), and DAPI (blue)).
 153 Representative single-color images are shown alongside 2-color overlay images. Scale bar
 154 indicates 100 μ m.

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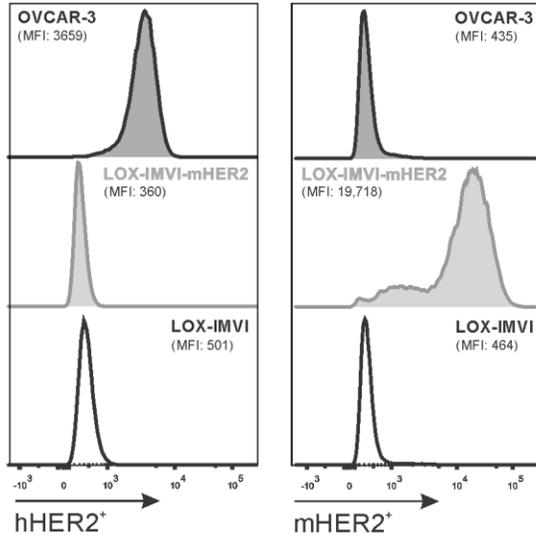
CD8 CD4 Ki-67 DAPI

158 Supplementary Figure 10:

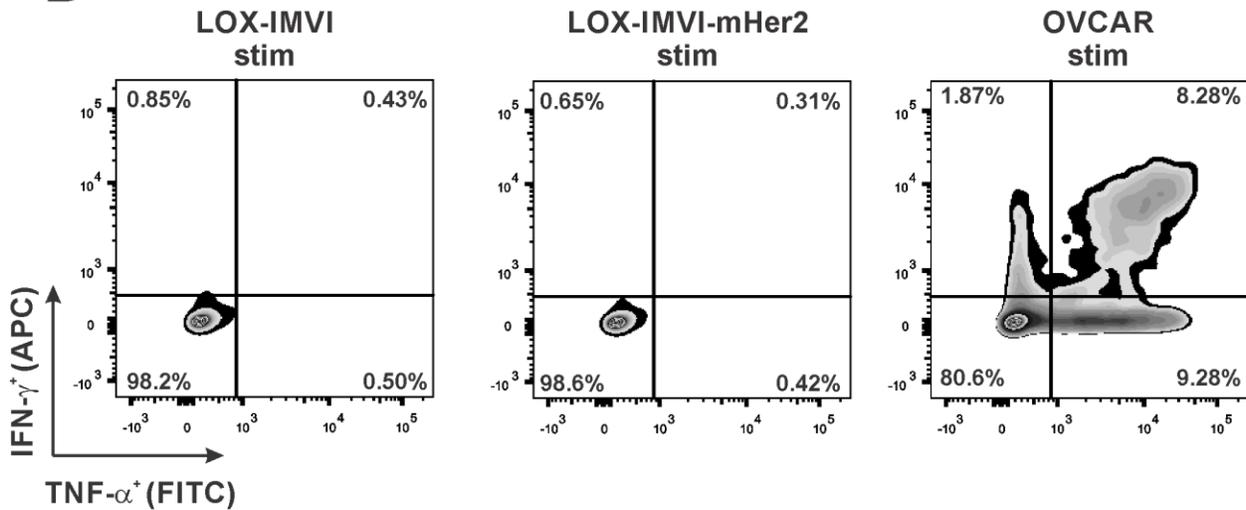
159 **Multicolor IHC of cardiac tissue at 7 days post-ACT1.** OVCAR-3 tumor bearing mice were
160 treated with 6.0×10^6 second-generation anti-HER2 28 ζ CAR- or HER2-TAC-T cells, or a
161 matched total number of vector control T cells. At 7 days post-ACT1 mice (n = 3 per treatment)
162 were perfused and tissues were formalin fixed and paraffin embedded for subsequent multicolor
163 IHC analysis (cardiac tissue was stained for cellular proliferation marker Ki-67 (purple), CD8
164 (cyan), CD4 (yellow), and DAPI (blue)). Scale bar indicates 100 μ m. Representative images are
165 shown.

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A



B



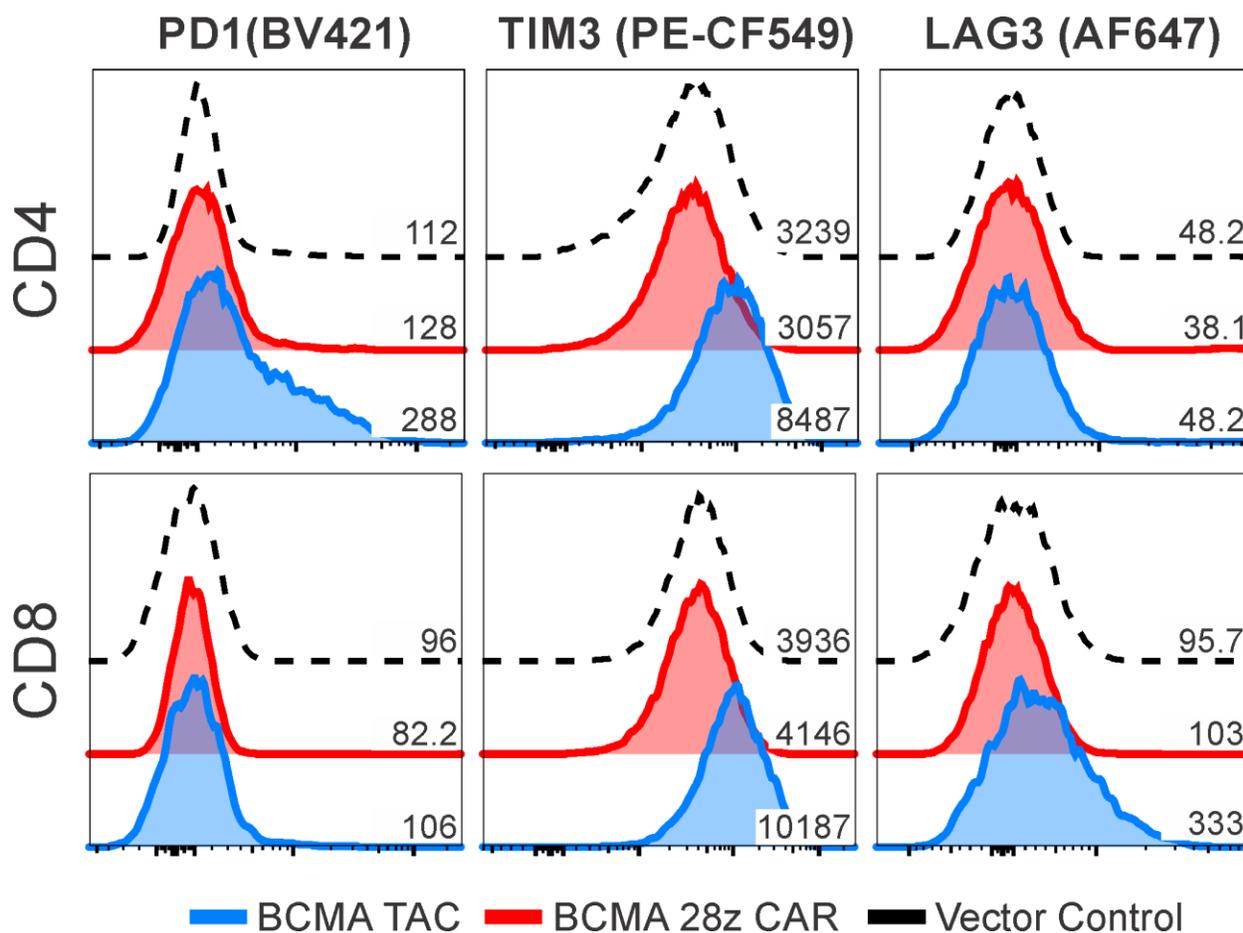
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Supplementary Figure 11.

Second generation anti-HER2 28z CAR-T cells do not respond to murine HER2. A.

Expression of human or murine HER2 (hHER2 or mHER2, respectively) on OVCAR-3, LOX-IMVI, and LOX-IMVI-mHER2 tumor cell lines as determined by flow cytometry. For the gating strategy see Supplementary Fig. 16. **B.** Production of activation cytokines (IFN- γ and TNF- α) by CD8⁺ CD28z-CAR T cells after stimulation with tumor cell lines (as indicated) was determined by flow cytometry. Data is representative of findings from two independent experiments. For the gating strategy see Supplementary Fig. 13B.

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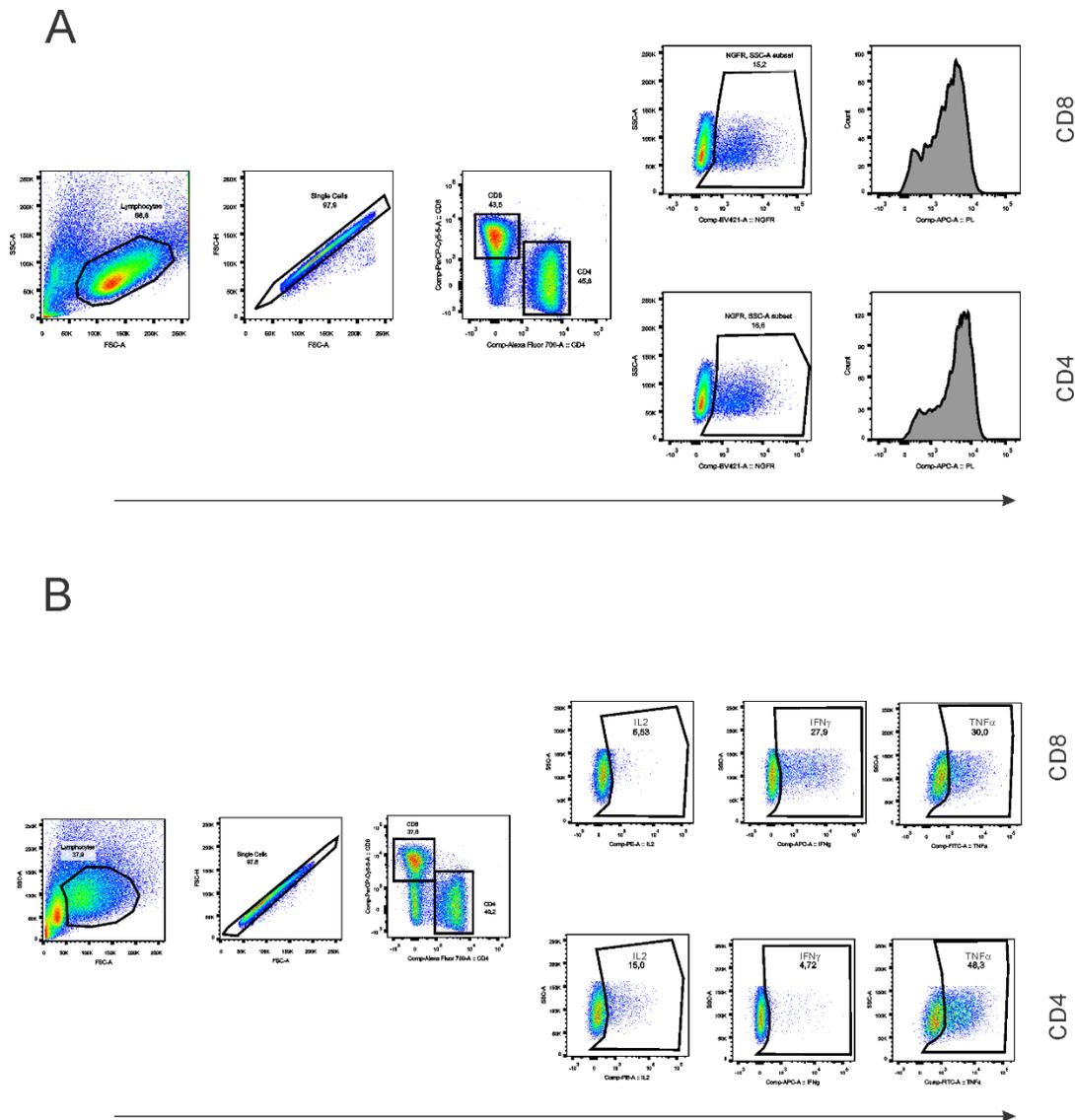


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Supplementary Figure 12:

Relative expression of checkpoint receptors in BCMA-CAR- and TAC-engineered T cells.

T cells were transduced with BCMA-TAC, second-generation BCMA-CD28 ζ -CAR, or vector control (tNGFR). Engineered T cells are stained for surface marker expression and analyzed by flow cytometry. Populations are gated on CD4⁺NGFR⁺ or CD8⁺NGFR⁺ analysis. Values indicate MFI of each marker. Histograms are from one donor, representative of three donors from three independent experiments. For the gating strategy see Supplementary Fig. 14

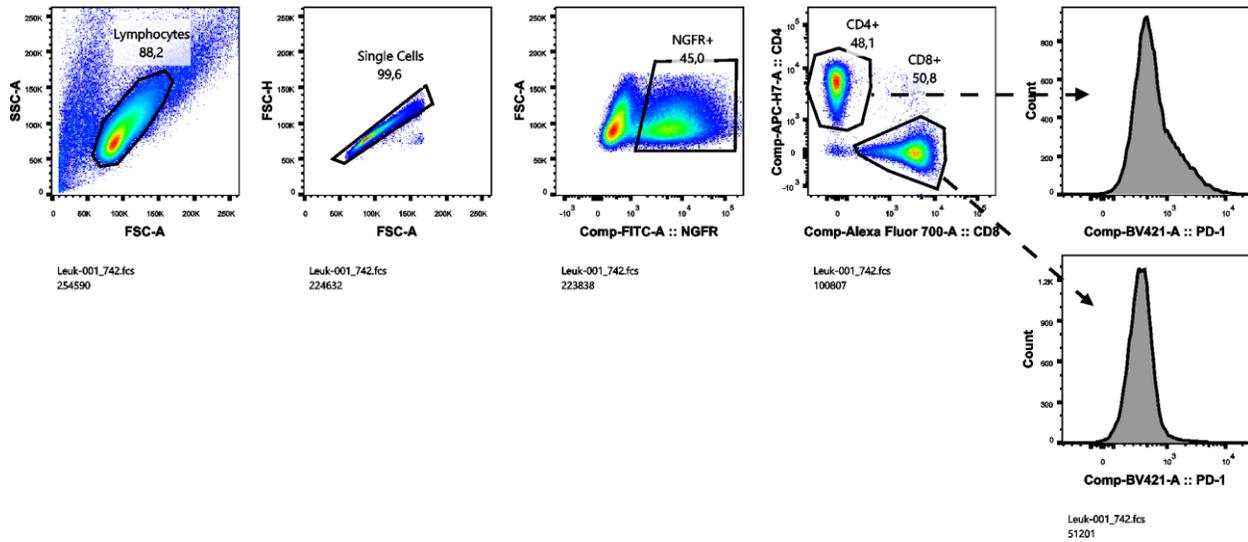


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196 Supplementary Figure 13:

197 **Examples of gating strategies used for the analysis of flow cytometry data. A.** Gating
 198 strategy for phenotypic analysis of engineered T cells. Lymphocytes (SSC-A v. FSC-A) →
 199 singlets (FSC-H v. FSC-A) → CD8⁺ or CD4⁺ cells (CD8-PerCPCy5.5 v. CD4-AF700) →
 200 NGFR⁺ cells (SSC-A v. NGFR-BV421) → TAC receptor expression as a histogram (Protein L⁺
 201 with indirect detection via APC-conjugated streptavidin is shown as an example). NGFR⁺ gates
 202 were set based on fully stained, non-transduced T cell controls. **B.** Gating strategy for functional
 203 cytokine production by tumor cell line-stimulated engineered T cells. Lymphocytes (SSC-A v.
 204 FSC-A) → singlets (FSC-H v. FSC-A) → CD8⁺ or CD4⁺ cells (CD8-PerCPCy5.5 v. CD4-
 205 AF700) → cytokine positive (SSC-A v. TNF-α-FITC or IFN-γ-APC or IL-2-PE). Cytokine⁺
 206 gates were set based on fully stained, PBS- or antigen-negative tumor cell line-stimulated
 207 controls.



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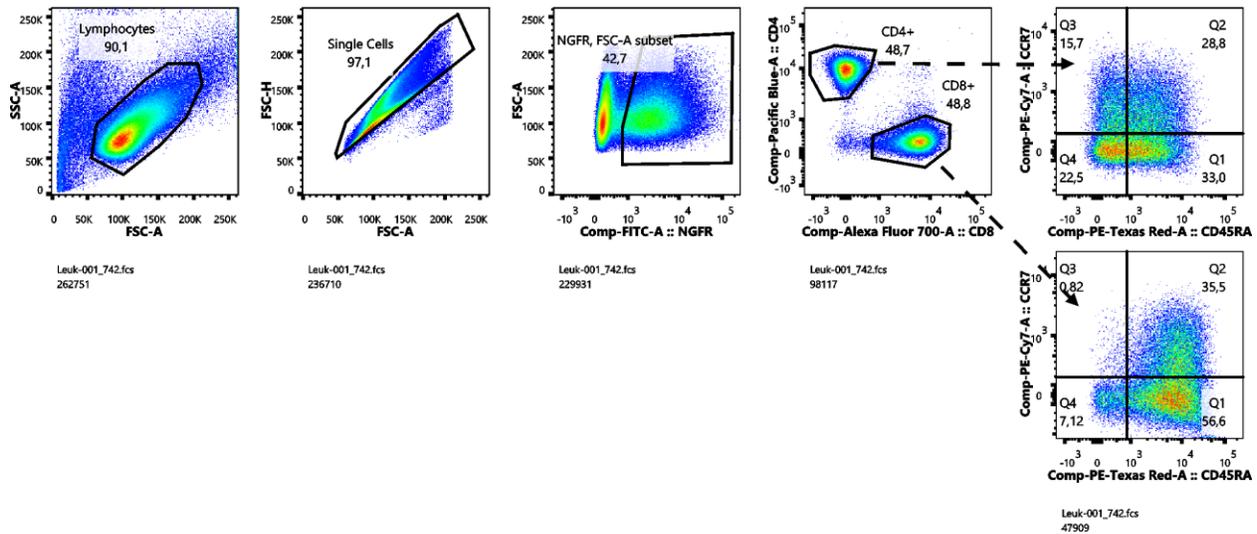
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210 Supplementary Figure 14:

211 **Examples of gating strategies used for the analysis of flow cytometry data.** Gating strategy
 212 for checkpoint analysis of engineered T cells. Lymphocytes (SSC-A v. FSC-A) → singlets (FSC-
 213 H v. FSC-A) → NGFR⁺ cells (SSC-A v. NGFR- VioBright FITC) → CD8⁺ or CD4⁺ cells (CD8-
 214 AF700 v. CD4-APC-H7-A) → Checkpoint receptor expression as a histogram (TIM-3-BV785/
 215 LAG-3-AF647/ PD-1-BV421). NGFR⁺ gates were set based on fully stained, vector control
 216 transduced T cells.

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222 Supplementary Figure 15:

223 **Examples of gating strategies used for the analysis of flow cytometry data.** Gating strategy
 224 for memory population analysis of engineered T cells. Lymphocytes (SSC-A v. FSC-A) →
 225 singlets (FSC-H v. FSC-A) → NGFR⁺ cells (SSC-A v. NGFR- VioBright FITC) → CD8⁺ or
 226 CD4⁺ cells (CD8-AF700 v. CD4- Pacific Blue) → Checkpoint receptor expression was analysed
 227 via quarter analysis (CD27-APC-H7/ CD28-PE/ CD45RA-ECD/ CCR7-PE-Cy7). NGFR⁺ gates
 228 were set based on fully stained, vector control transduced T cells.

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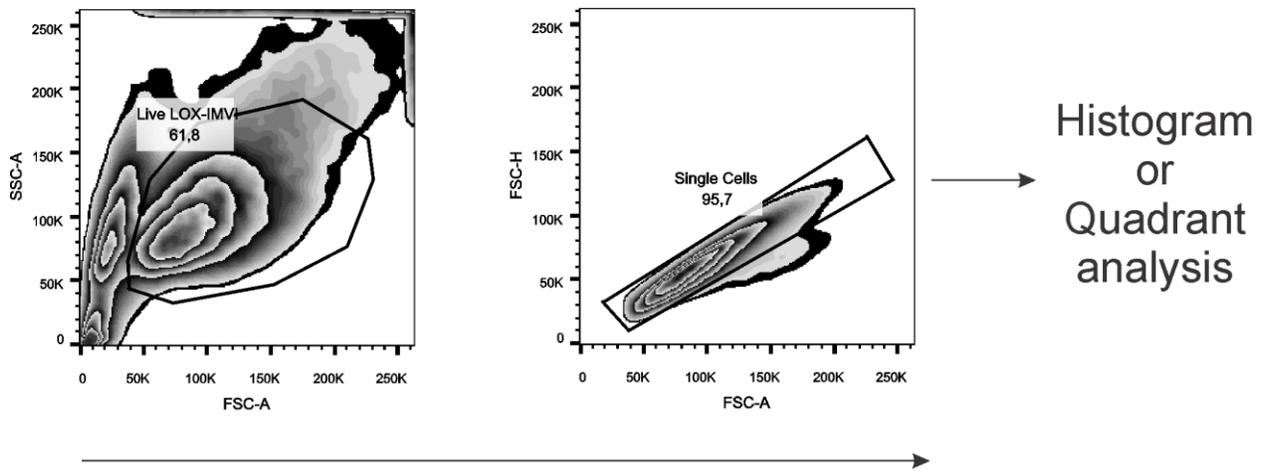
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236 Supplementary Figure 16:

237 **Examples of gating strategies used for the analysis of flow cytometry data.** Gating strategy
238 for various adherent cell lines engineered with various surface expressing proteins. Cell gate
239 (SSC-A v. FSC-A) → singlets (FSC-H v. FSC-A) → Quadrant or Histogram analysis with
240 relevant markers

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Test for significance in cytokine levels

Day 3	Condi- tions	GM-CSF	IFN γ	IL-1 β	IL-2	IL-4	IL-6	IL-8	IL-10	IL-12 (p70)	MCP-1	TNF α	IL-13	IL-5
	TAC- Vector	NS	NS	NS	NS	NS	NS	NS	NS	NS	NS	N/A	NS	NS
TAC- CAR	0.00068	0.00051	NS	< 0.0001	< 0.0001	NS	NS	NS	0.00101	NS	N/A	0.0036	0.00239	NS
CAR- Vector	0.00013	0.00021	NS	< 0.0001	< 0.0001	NS	NS	NS	0.0009	NS	N/A	0.00254	0.00052	NS

Day 7	Condi- tions	GM-CSF	IFN γ	IL-1 β	IL-2	IL-4	IL-6	IL-8	IL-10	IL-12 (p70)	MCP-1	TNF α	IL-13	IL-5
	TAC- Vector	NS	NS	N/A	NS	NS	NS	NS	NS	NS	NS	NS	0.00093	NS
TAC- CAR	< 0.0001	0.00035	N/A	0.00038	0.00522	NS	NS	0.00296	0.00061	0.00652	NS	NS	0.00025	< 0.0001
CAR- Vector	< 0.0001	0.00035	N/A	0.00035	0.00523	NS	NS	0.00325	0.00063	0.00867	NS	NS	0.00025	< 0.0001

242

243 Supplementary Table 1:

244 **Statistical analysis of serum cytokine data.** An unpaired t-test was used to evaluate the
 245 statistical significance of the serum cytokine data (n = 3 for each of HER2-TAC-, HER2-CAR-,
 246 and vector control T cells) presented in Figure 6. Pairwise comparisons between the three
 247 treatment groups are presented for each of the thirteen cytokines tested. P-values are shown. NS
 248 = not significant using a confidence interval of 95%. N/A denotes cases where no analysis was
 249 mathematically possible.

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FWD-1	GGCGCGCCATGGATTTCCAGGTCCAGATTTTC
REV-1	CCCGGGGTTTCAGGTCTTCTTCGCTAATC
FWD-2	ATATGGCGCGCCATGGATTTCC
REV-2	GCTGAACTTCACTCTGCAGTAAAGGGTGATAACCAG
FWD-3	ATCACCTTTACTGCAGAGTGAAGTTCAGCAGGAG
REV-3	ATATGCTAGCTTAGCGAGGGGG
FWD-4	ATATGGATCCGTGGGGTCACCGTCTCCTCAACC
REV-4	ATATGCTAGCTTAGCGAGGGGGCAGGGC
FWD-5	CATGTGACTCCACGGAGTACCGG
REV-5	GTCCGAACGTCCACGGCAGAG
FWD-6	GGTCTTCTTCGCTAATCAGTTTCTGTTCCGGCCTGGCGGCCTGG
REV-6	CCACGCCGCCAGGCCGGAACAGAACTGATTAGCGAAGAAGACC

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254 Supplementary Table 2:
255 Sequence of oligos used in construction of various plasmid constructs described in materials and
256 methods.