

**Supplementary information**

---

**Enhanced intratumoural activity of CAR T cells engineered to produce immunomodulators under photothermal control**

---

In the format provided by the authors and unedited

**Contents:**

**Supplementary Figure 1:** qPCR screen of HSPs in primary murine T cells.

**Supplementary Figure 2:** Transduction efficiencies of primary human T cells from three donors.

**Supplementary Figure 3:** Thermal switch specificity in Jurkat T cells.

**Supplementary Figure 4:** Gating strategy for viability flow staining.

**Supplementary Figure 5:** Longitudinal heating of primary human T cells.

**Supplementary Figure 6:** Repeated heat treatments do not affect CAR T cell cytotoxicity.

**Supplementary Figure 7:** CD19 expression on K562 and Raji tumour cells.

**Supplementary Figure 8:** Longitudinal control of intratumoural CAR T cells using photothermal pulses.

**Supplementary Figure 9:** TS-Fluc  $\alpha$ CD19 CAR T cell infiltration into K562 and Raji flank tumours.

**Supplementary Figure 10:** Cytokine support improves proliferation of T cells receiving low levels of CD3/28 stimulation.

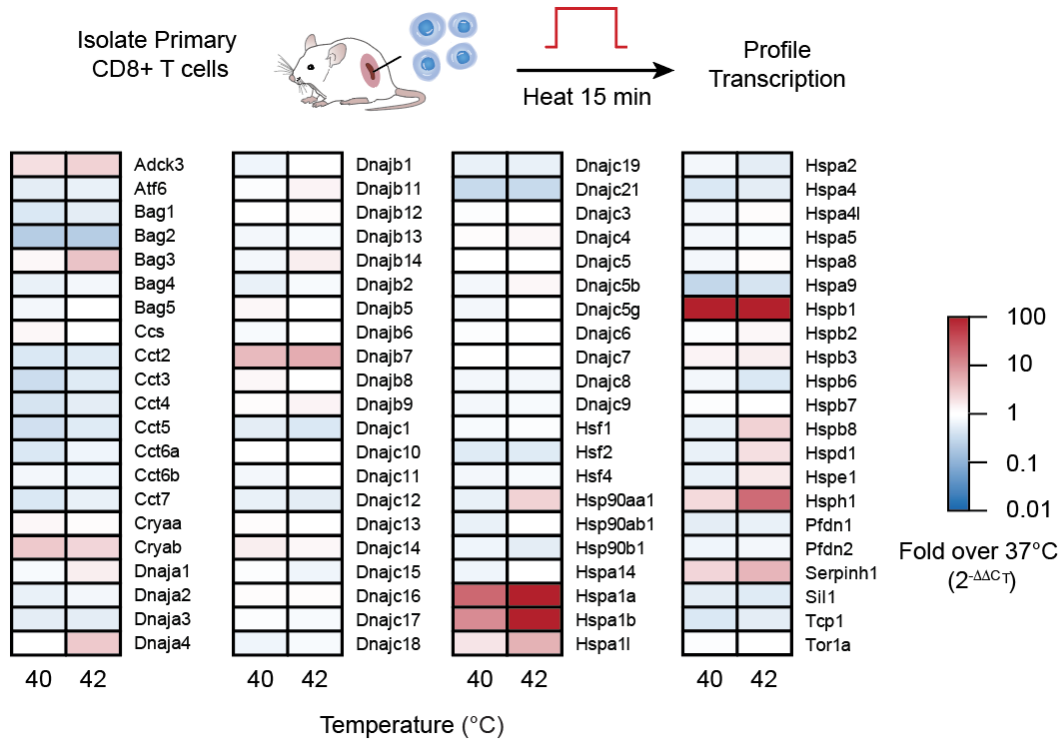
**Supplementary Figure 11:** Gating strategy for mixed proliferation experiment.

**Supplementary Figure 12:** Characterization of engineered Pmel-1 T cells.

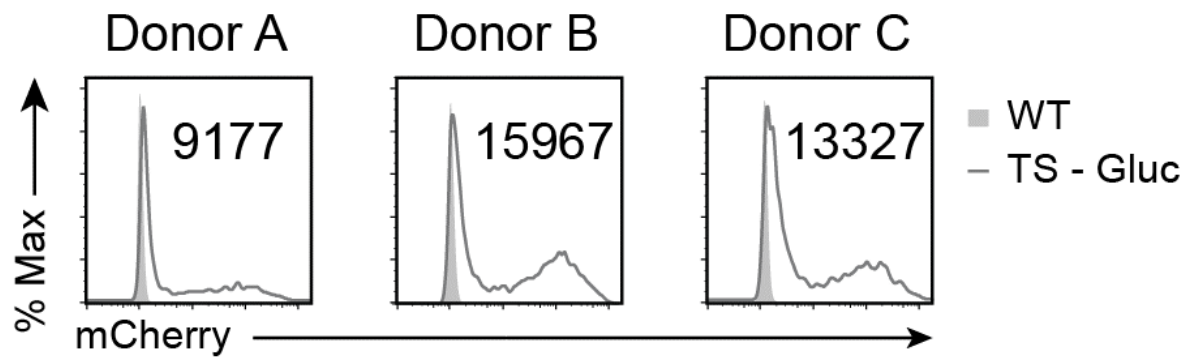
**Supplementary Figure 13:** Validation of MDA-MB-468 transduced with HER2 or Renilla Luciferase.

**Supplementary Figure 14:** TS-BiTE  $\alpha$ HER2 CAR T cells activate when incubated with HER2+ MDA-MB-468 cells.

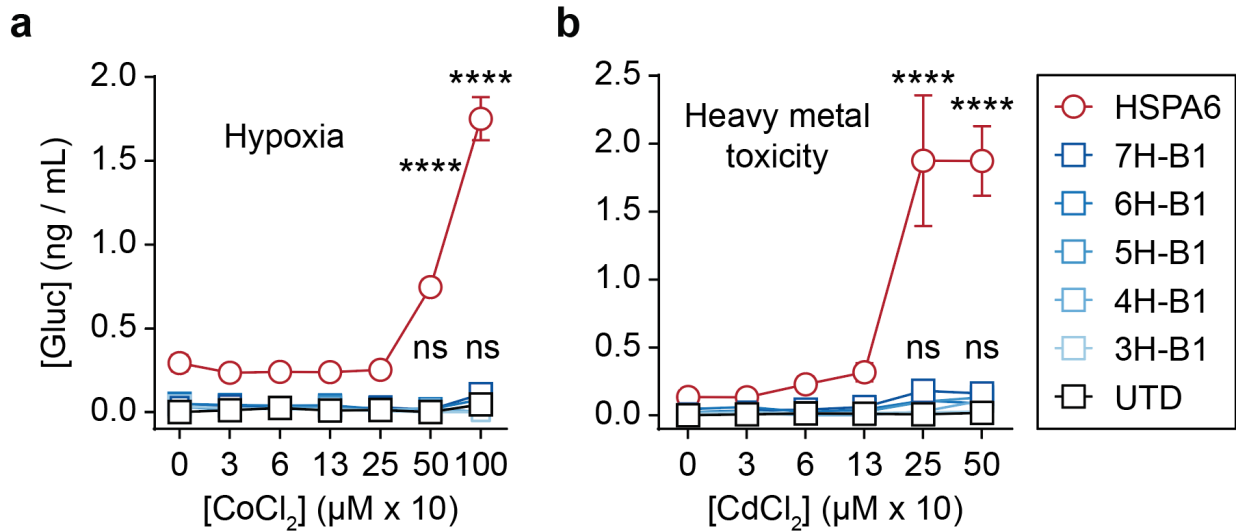
**Supplementary Table 1:** Cell line authentication via STR analysis.



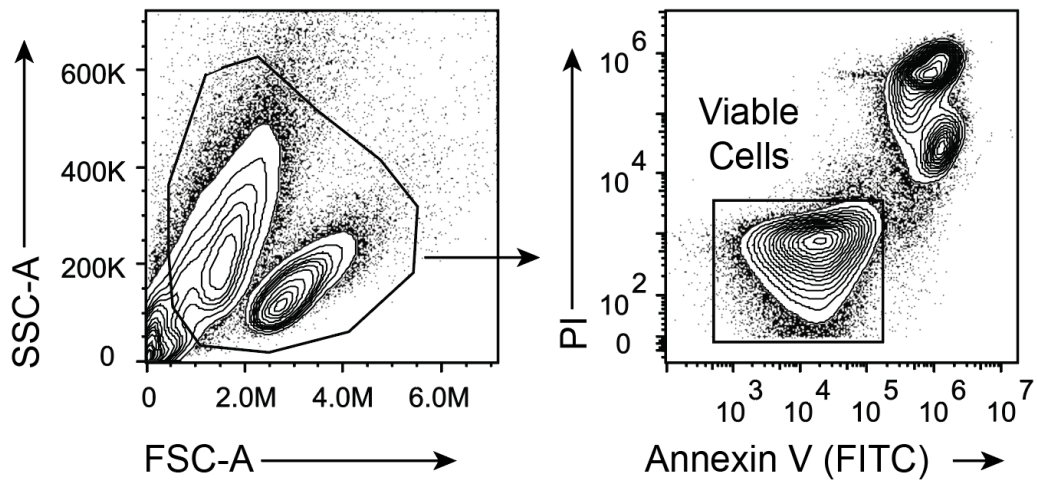
**Supplementary Figure 1: qPCR screen of HSPs in primary murine T cells.** Splenic CD8+ T cells were isolated using the CD8+ T cell isolation kit according to (Miltenyi 130-104-075). Six hours after indicated heat treatments, mRNA was harvested and quantified using the Mouse HSP profiler kit (Qiagen PAMM-076Z) according to manufacturer instructions. Data are displayed relative to unheated controls.



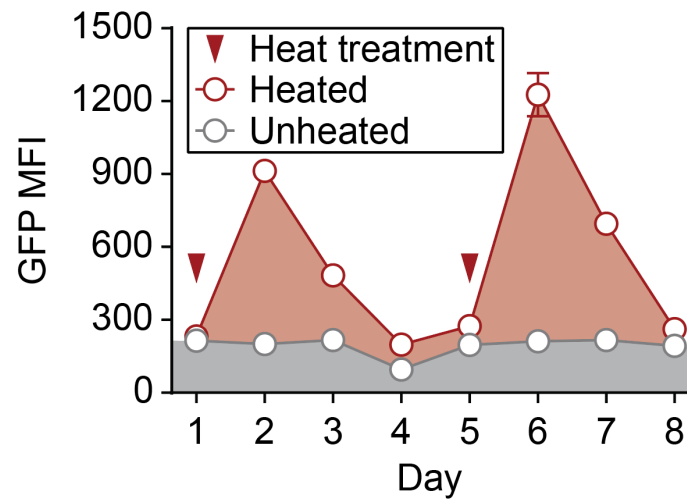
**Supplementary Figure 2: Transduction efficiencies of primary human T cells from three donors.** Flow cytometric plots of primary human T cells derived from 3 donors and transduced with the Gluc expressing 7H-YB Thermal switch containing a constitutively expressed mCherry reporter. Inset shows the mean fluorescent intensity (MFI) of mCherry transduced cells.



**Supplementary Figure 3: Thermal switch specificity in Jurkat T cells.** Gluc activity by Jurkat T cells transduced with synthetic thermal gene switch constructs (blue) or the endogenous HSPA6 promoter (red) following exposure to (a) CoCl<sub>2</sub> to mimic hypoxia or (b) to CdCl<sub>2</sub> to model heavy metal toxicity (ns = not significant, \*\*\*\*P<0.0001, two-way ANOVA and Tukey post-test and correction, error bars show SEM, n = 3).

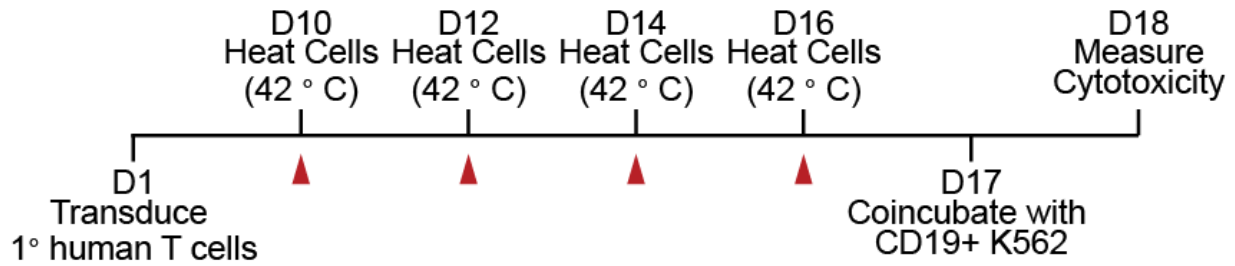


**Supplementary Figure 4: Gating strategy for viability flow staining.** Primary human T cells were heated at 42 °C for 60 minutes as a positive control for thermal damage. Shorter regimens were used for subsequent experiments. Because many of the AnnexinV+ or PI+ events were not within tighter FSC/SSC gates, this conservative gating strategy was used as it better represented the sample's overall viability.

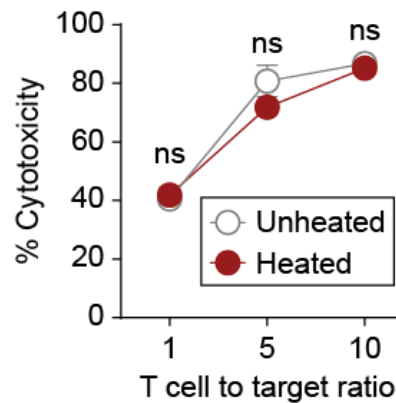


**Supplementary Figure 5: Longitudinal heating of primary human T cells.** Primary human T cells transduced with an HSPA6-GFP switch were repeatedly heated once GFP signal had returned to baseline after previous heat treatment (n = 3 biologically independent wells, error bars show SEM). Two independent experiments were performed with similar results.

**a**

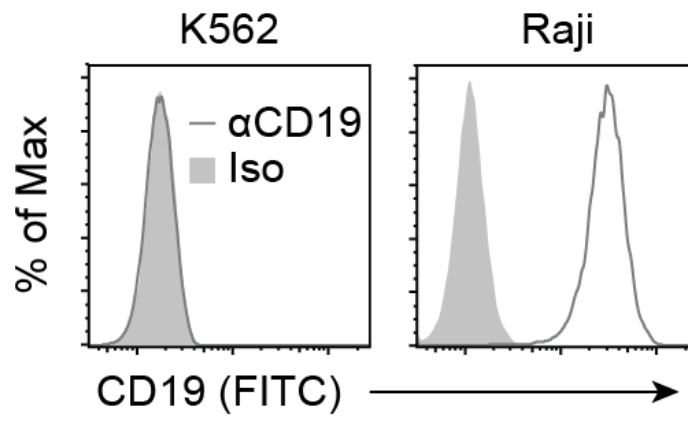


**b**

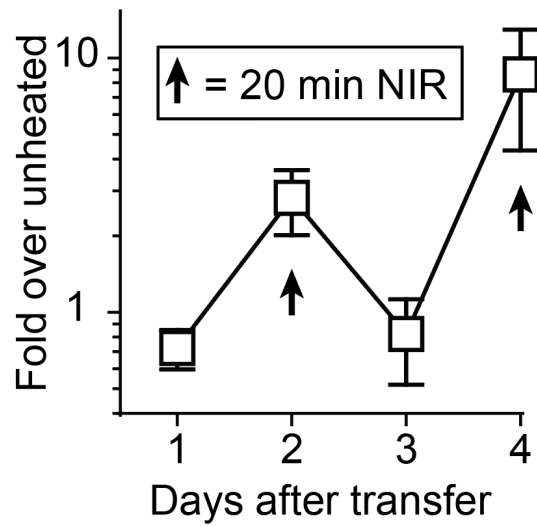


**Supplementary Figure 6: Repeated heat treatments do not affect CAR T cell cytotoxicity.** (a) Primary human T cells were transduced to constitutively express an  $\alpha$ CD19 CAR following CD3/CD28 bead activation. Heat treatments were performed at indicated timepoints prior to coincubation with luciferized, CD19+ K562s according to the timeline. (b) Percent cytotoxicity was quantified by loss of luminescence in wells relative to control wells containing only target cells (b) (ns = not significant, two-way ANOVA and Sidak post-test and correction, mean  $\pm$  SEM is depicted, n = 3 biologically independent wells).

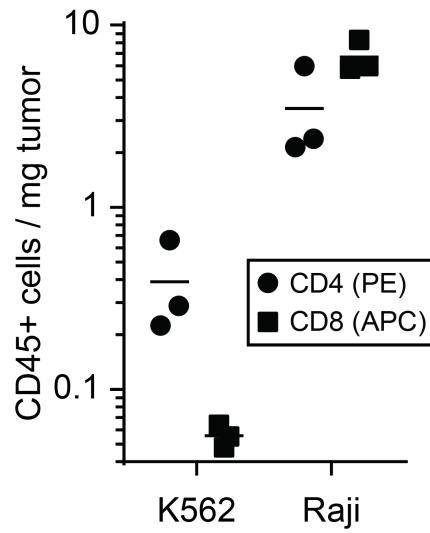




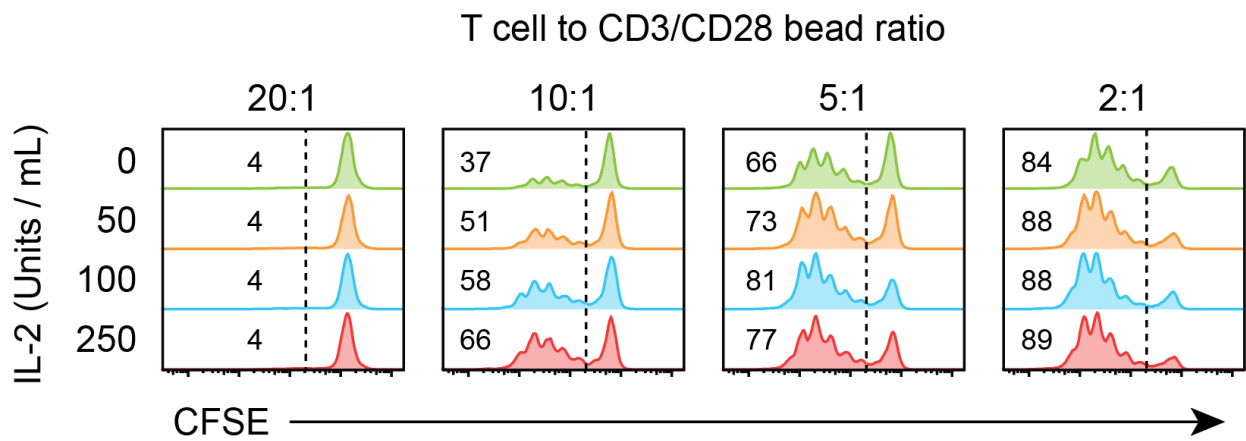
**Supplementary Figure 7: CD19 expression on K562 and Raji tumour cells.** Representative flow cytometric plots of CD19 staining on K562 and Raji cell lines (Iso = isotype control).



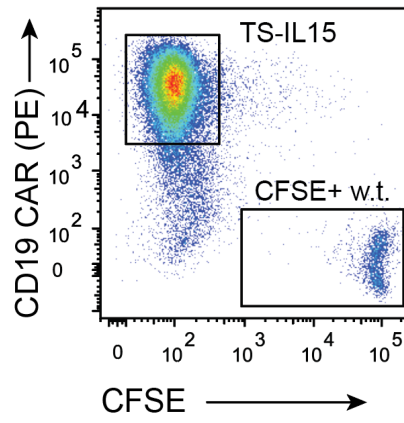
**Supplementary Figure 8: Longitudinal control of intratumoural CAR T cells using photothermal pulses.** Mice bearing Raji tumours (CD19+) were injected i.v. with TS-Fluc T cells. Tumour sites were irradiated on days 2 and 4 using NIR laser light as shown in **Figure 5d**. Luminescence was quantified daily via i.v. injections of D-luciferin (n = 3, biologically independent wells, error bars show SEM).



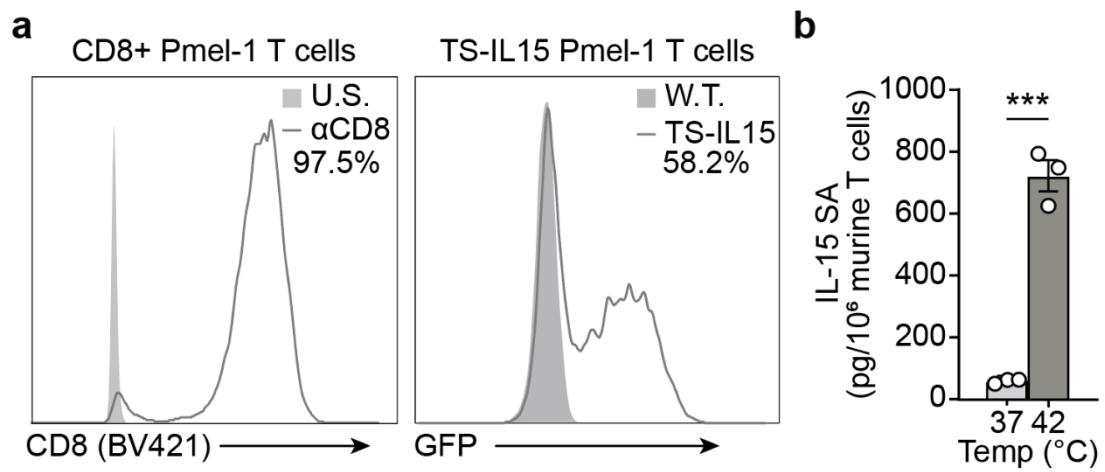
**Supplementary Figure 9: TS-Fluc  $\alpha$ CD19 CAR T cell infiltration into K562 and Raji flank tumours.** TS-Fluc  $\alpha$ CD19 CAR T cells were injected i.v. into tumour bearing mice once tumours had reached  $\sim 250$  mm<sup>3</sup>. After 7 days, tumours were resected, dissociated, and stained to quantify cellular infiltration using flow cytometry counting beads (n = 3, biologically independent wells, error bars show SEM).



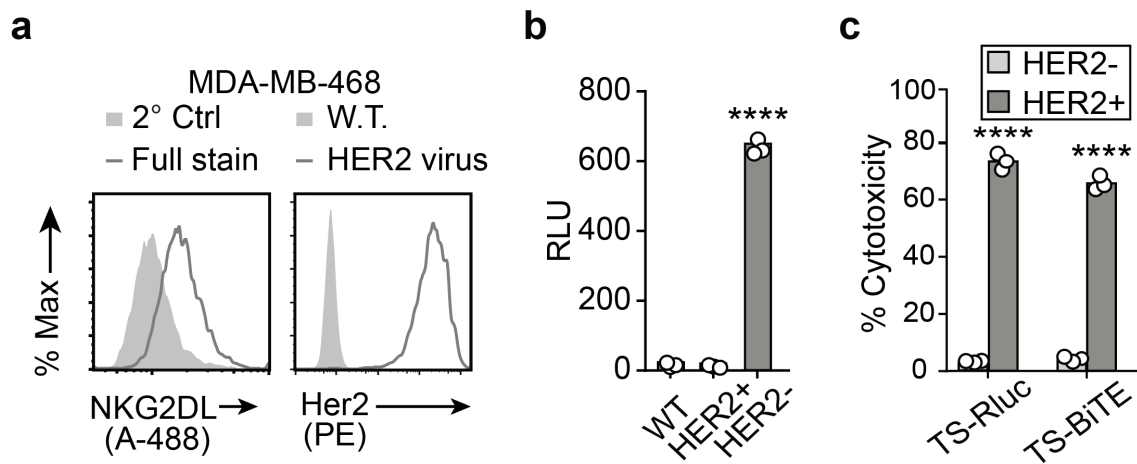
**Supplementary Figure 10: Cytokine support improves proliferation of T cells receiving low levels of CD3/28 stimulation.** T cells were labeled with CFSE and incubated with low levels of activating beads. For reference, routine expansion and culture of T cells uses 3 beads for every T cell. Increasing amounts of IL-2 were added to each bead ratio. All samples were assayed after 4 day incubations at indicated conditions. Two independent experiments were performed with similar results.



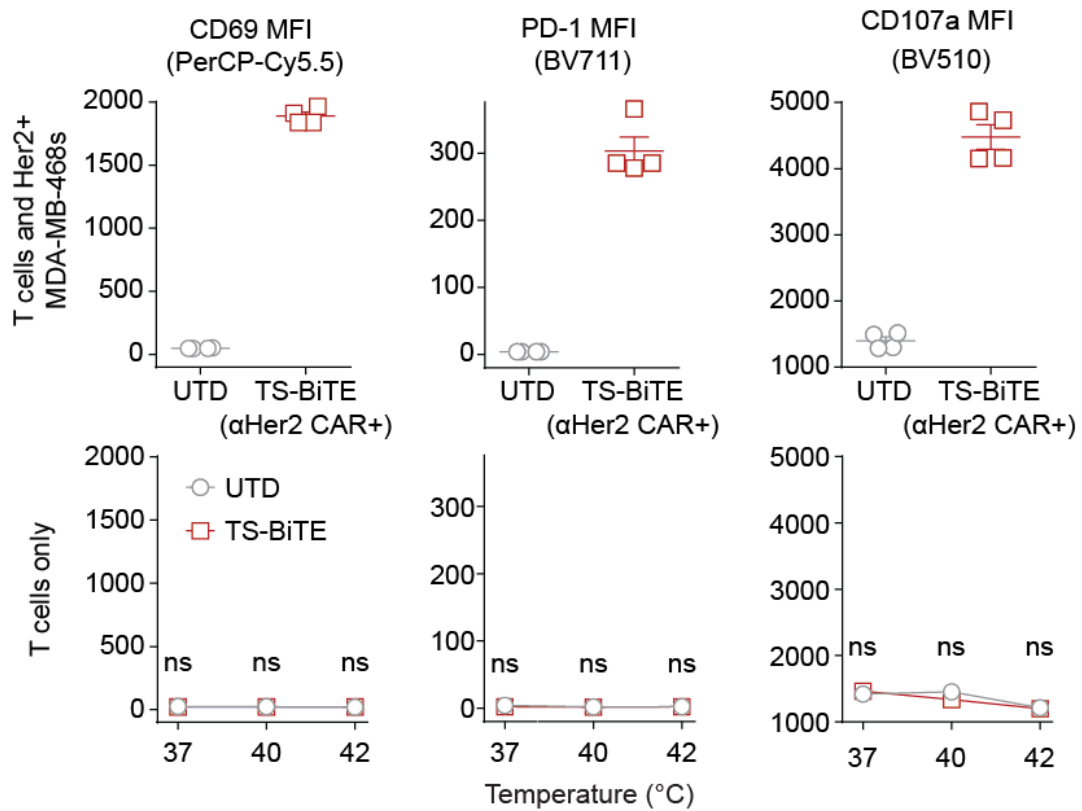
**Supplementary Figure 11: Gating strategy for mixed proliferation experiment.** Transduced TS-15  $\alpha$ CD19 CAR T cells were identified by CAR expression. Proliferation of CFSE+ wild-type cells was assessed by dye dilution and FlowJo proliferation tool.



**Supplementary Figure 12: Characterization of engineered Pmel-1 T cells.** 48 hours post isolation and peptide activation, Pmel-1 derived splenocytes were transduced with the TS-IL15 vector containing a constitutive GFP reporter. **(a)** Pmel-1 T cells were characterized via flow cytometry before adoptive transfer to assess CD8+ cell purity expansion (left) and transduction efficiency (right) (U.S. = Unstained Pmel-1 T cells, W.T = Wild type Pmel-1 T cells). **(b)** IL-15 production measured via ELISA from transduced murine T cells after 20 minute heating at indicated temperatures (\*\*\*) P=0.0002, unpaired T test, mean± SEM is depicted, n = 3 biologically independent wells).



**Supplementary Figure 13: Validation of MDA-MB-468 transduced with HER2 or Renilla Luciferase.** (a) Representative flow plots of NKG2DL staining and HER2 Staining of HER2+ MDA-MB-468s. MDA-MB-468 were transduced with lentivirus to stably surface express HER2. (b) Luminescence of Fluc transduced HER2- MDA-MB-468 tumour cells (\*\*\*\*  $P < 0.0001$ , one-way ANOVA and Sidak post-test and correction, mean  $\pm$  SEM is depicted,  $n = 3$  biologically independent wells). (c) Percent cytotoxicity observed via LDH assay in HER2- or HER2+ MDA-MB-468 cells after incubation with T cells constitutively expressing CARs (\*\*\*\* $P < 0.0001$  between HER2- and HER2+ groups, two-way ANOVA and Sidak post-test correction, mean  $\pm$  SEM is depicted,  $n = 3$  biologically independent wells). Two independent experiments were performed with similar results.



**Supplementary Figure 14: TS-BiTE  $\alpha$ HER2 CAR T cells activate when incubated with HER2+ MDA-MB-468 cells.** MFIs of activation and degranulation markers CD69, PD-1, and CD107a on TS-BiTE  $\alpha$ HER2 CAR T cells that were heated at indicated temperatures prior to co-incubation with HER2+ MDA-MB-468 target cells. (ns = not significant, two-way ANOVA and Tukey post-test and correction, error bars show SEM, n = 4 biologically independent wells).



<b>Genetic Locus</b>	<b>Jurkat E6.1</b>	<b>K-562</b>	<b>MDA-MB-468</b>	<b>Raji</b>
D3S1358	15, 17	16	15	15, 16
TH01	6, 9.3	9.3	7	6, 7
D21S11	31.2, 32.2, 33.2, 34.2	29, 30, 31	27, 28	28, 31
D18S51	13, 20, 21	15, 16	17	17
Penta E	10, 12	5, 14	5	5, 13
D5S818	9	11, 12	12	10, 13
D13S317	8, 12	8	12	13
D7S820	8, 12, 13	9, 11	8	10
D16S539	11	11, 12	9	8, 11
CSF1PO	10, 11, 12, 13	9, 10	12	10, 12
Penta D	11, 13	9, 13	8, 10	3.2, 9
vWA	18, 19	16	18	16, 19
D8S1179	13, 14, 15	12	13	14, 15
TPOX	8, 10	8, 9	8, 9	8, 13
FGA	20, 21	21	23	19, 27
AMEL	X, Y	X	X	X, Y
Mouse Contamination	Not detected	Not Detected	Not Detected	Not Detected
<b>% Match</b>	<b>91.89%</b>	<b>100%</b>	<b>100%</b>	<b>100%</b>

**Supplementary Table 1: Cell line authentication via STR analysis.** Jurkat E6.1, K-562, MDA-MB-468, and Raji human cell lines were authenticated via PowerPlex16HS STR profiling (Labcorp). Results were compared to known samples in the Cellosaurus STR Similarity Search Tool (CLASTR 1.4.4) database.