Supplementary Materials

Figure S1. Diagram for screening strategies.

Figure S2. Anti-CD19 CAR-T cells specifically react with CD19⁺ tumor cells *in vitro*.

Figure S3. Tumor growth for mice receiving antibody-mediated eosinophil depletion and without CAR-T infusion

Figure S4. Anti-CCR3 antibody and anti-Siglec-F antibody deplete eosinophils *in vivo* separately.

Figure S5. Flow cytometry gating strategy for eosinophils and anti-CD19 CAR-T cells.

Figure S6. Fewer CAR-T cells were recovered from the tumors treated with eosinophils depletion antibody.

Figure S7. Tumor volume change in CD19-CART treated mice with/without eosinophil transferring.

Figure S8. Expression of intratumoral T-cell attractants

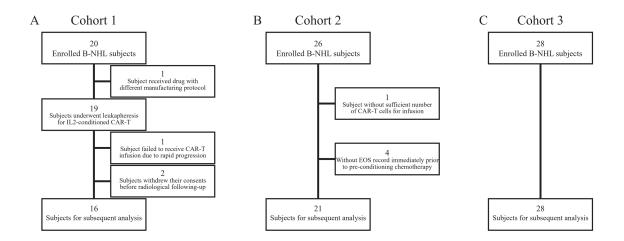


Figure S1. Diagram for screening strategies. Diagrams showed the screening strategies for identifying candidates for evaluating the predictive significance of baseline eosinophils. (A) Cohort 1; (B) Cohort 2; (C) Cohort 3.

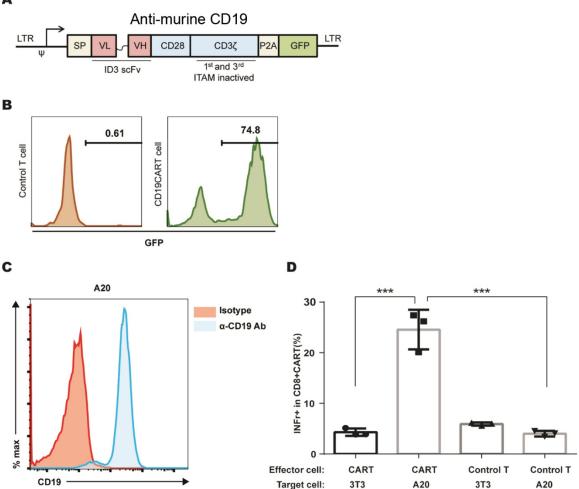


Figure S2. Anti-murine CD19 CAR-T cells specifically react with CD19⁺ tumor cells in vitro. (A) Diagram of the DNA encoding the 1D3-28Z CAR (ψ , retroviral packaging signal). (B) Prior to the mouse infusion procedure outlined in Fig. 3A, anti-murine CD19 CAR-T cells were tested for CAR expression. (C) Expression of CD19 on A20 lymphoma cells. (D) Specific reactivity of anti-CD19 CAR-T cells against CD19⁺ A20 lymphoma cells. Anti-CD19 CAR-T cells (4×10^5 /ml) were co-cultured with A20 or 3T3 cells (CD19⁻) at a ratio of 1:1 for 12 hours, then intracellular IFN- γ staining assays were conducted. IFN- γ expression in CAR-T cells co-cultured with A20 cells was significantly higher compared with 3T3 cell co-culture, demonstrating CAR-T specificity against CD19. n = 3 replicates for each group; *P < 0.05; ***P < 0.001 by unpaired *t*-test)

A

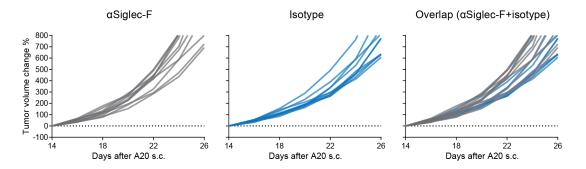


Figure S3. Tumor growth for mice receiving antibody-mediated eosinophil depletion and without CAR-T infusion. Mice were prepared as illustrated in Figure 3A. Infusion of CD19-CART i.v. on day 14 was replaced by saline. Each line represents one mouse in experiment.

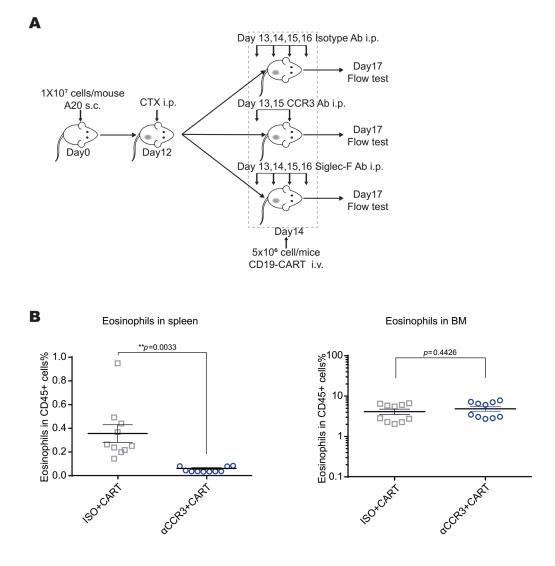


Figure S4. Anti-CCR3 antibody and anti-Siglec-F antibody deplete eosinophils in vivo separately. (A) Experimental schema for analyzing eosinophil and CAR-T cell counts after eosinophil depletion. (B) Eosinophil percentages among CD45⁺ splenocytes (left) and bone marrow (right) for isotype antibody, CCR3 antibody and Siglec-F antibody treated groups. Isotype, gray, n = 10; anti-CCR3, blue, n = 10; Data in left panels in (B) were analyzed by unpaired two-tailed Welch's t-test.

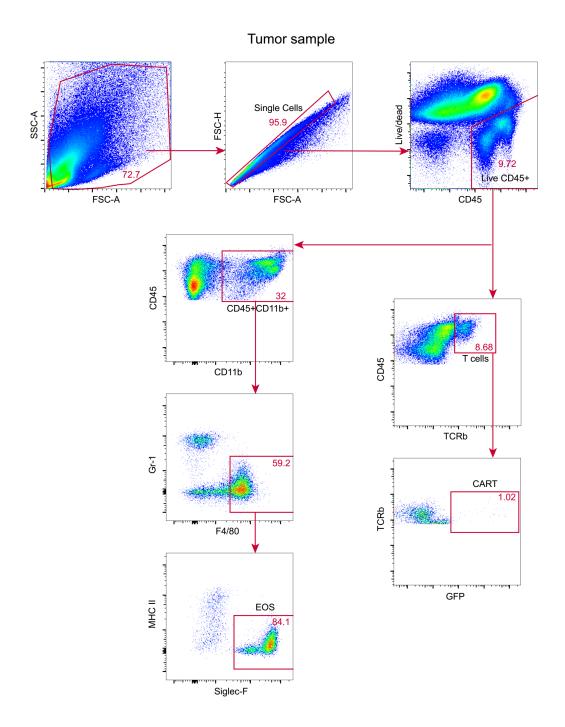


Figure S5. Flow cytometry gating strategy for eosinophils and CD19 CART cells. Tumor samples were gated on FSC-A versus SSC-A followed by a singlet gate excluded any doublets, then a viability CD45⁺ gate was applied to exclude any dead cells. CD19-CART cells were assessed by TCRb and GFP double positive group. Eosinophils were first gated from CD45⁺ live cells, and then CD11b⁺Gr-1^{lo}F4/80⁺Siglec-F⁺MHCII⁻ groups were analyzed as eosinophils.

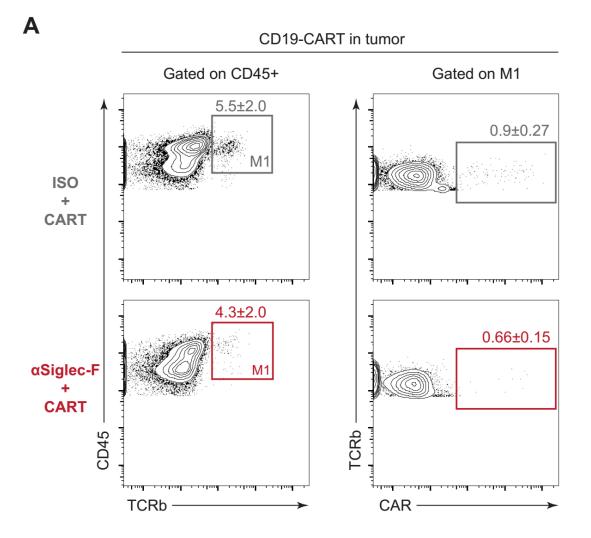


Figure S6. In these eosinophil-depleted animals, markedly fewer CAR-T cells were recovered from the tumor. Representative flow cytometry plots of tumorinfiltrating T cells (left) and intratumoral anti-CD19 CAR-T cells (right). Data are shown as means ±SD.

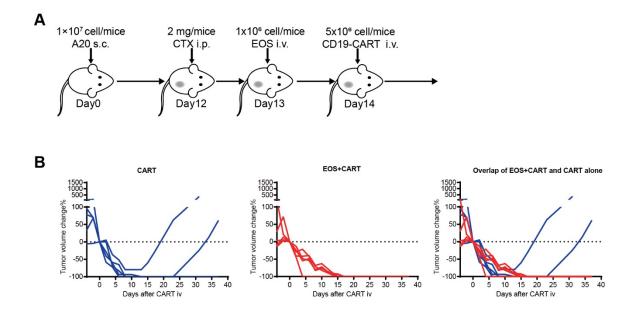


Figure S7. Tumor volume change percentage in CD19-CART treated mice with/without eosinophil transferring. (A) Experimental schema for eosinophils transferring and CAR-T cells treatment in tumor-bearing mice. Balb/c mice were subcutaneously injected with 1×10^7 A20 lymphoma cells. 12 days later, all the mice were treated with 2mg/mouse CTX for preconditioning, following with/without 1×10^6 activated eosinophils intravenously injection on day13. 5×10^6 CD19-CART cells were injected into mice on day14 (Arrow). (B) Tumor volume change percentage in mice transferred with eosinophils or without eosinophils. Blue, mice were transferred with CD19-CART only; red, mice were transferred with eosinophils and CD19-CART cells. (n=5 per group, tumor size at the time point of CART injection was regard as baseline.)

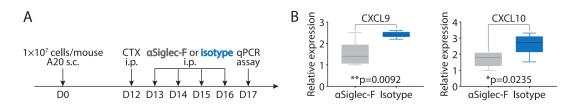


Figure S8. Expression of intratumoral T-cell attractants. (A) Experimental schema. Ten million mouse lymphoma A20 (CD19⁺) cells were subcutaneously injected into syngeneic *BALB/c* mice on day 0. Each mouse was intraperitoneally administered 2 mg cyclophosphamide (CTX) for preconditioning on day 12. Each mouse was intraperitoneally administered 15 µg/d anti-mouse Siglec-F antibody or isotype control for 4 d starting on day 13. Intratumoral CXCL9 and CXCL10 expression levels were measured by qPCR on day 17. (B) Boxplots showing relative CXCL9 and CXCL10 expression in Siglec-F- or isotype-treated tumors (n = 5). Horizontal line: median expression level. Statistical analysis consisted of Mann-Whitney *U*-test. qPCR primers were CXCL9 forward: CTTTTCCTCTTGGGCATCAT; CXCL9 reverse: GCATCGTGCATTCCTTATCA; CXCL10 forward: GACGGTCCGCTGCAACTG; and CXCL10 reverse, CCCTATGGCCCTCATTCTCA.