Allogeneic Chimeric Antigen Receptor T Cells Targeting B Cell Maturation Antigen

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Abstract

B cell maturation antigen (BCMA) is a tumor necrosis factor family cell surface receptor that binds B cell activating factor (BAFF) and a proliferation inducing ligand (APRIL) and is involved in the long-term survival of B cells. BCMA has been shown to be expressed broadly on monoclonal gammopathy of undetermined significance (MGUS) and multiple myeloma (MM) cells and for this reason has been pursued as a potential antigen for the treatment of MM. Indeed, clinical studies evaluating autologous chimeric antigen receptor T cells (CAR-T cells) targeting BCMA for the treatment of MM have shown promising efficacy with >80% response rates. However, logistical challenges will potentially limit the number of patients that have access to autologous therapies. Using the CRISPR/Cas9 system, we generated allogeneic CAR-T cells targeting BCMA by disrupting the beta-2-microglobulin ($\beta 2M$) and T cell receptor alpha constant (*TRAC*) genes and inserting an anti-BCMA CAR into the TRAC locus. This results in allogeneic CAR-T cells lacking the major histocompatibility complex (MHC-I) and endogenous T cell receptor (TCR) expression while being potently cytotoxic towards cells expressing BCMA. These data show proof-of-concept for an "off-the-shelf" CAR-T therapy targeting BCMA.

Figure 1: CRISPR Therapeutics Allogeneic CAR-T Pipeline

Program	Editing approach	Research	IND-enabling	Ph I/II	Partner
CTX101: Anti-CD19 allogeneic CAR-T	Disruption & Insertion			IND filing Q4 2018	Wholly-owned
CTX102: Anti-BCMA allogeneic CAR-T	Disruption & Insertion				Wholly-owned
CTX103: Anti-CD70 allogeneic CAR-T	Disruption & Insertion				Wholly-owned
Multiple solid tumor allogeneic CAR-T	Disruption & Insertion				Wholly-owned

Figure 2: CRISPR/Cas9 Gene-Edited Allogeneic Anti-BCMA **CAR-T Cells**



by homology-directed repair after using CRISPR/Cas9 to introduce a site-specific double strand break. To enhance persistence of allogeneic cells, MHC-I expression is eliminated by disrupting the $\beta 2M$ gene.

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(A) Representative FACS plots of TRAC and $\beta 2M$ expression (left panel) and CAR expression following site-specific knock-in to the TRAC locus (right panel) 1 week following gene editing. (B) Editing results in decreased surface expression of TCR and MHC-I, as well as high CAR expression. More than 60% cells possess all 3 desired modifications (TCR⁻/ β 2M⁻/CAR⁺). (**C**) Production of allogeneic anti-BCMA CAR-T cells preserves CD4 and CD8 proportions.

Figure 4: Allogeneic CAR-T Cells Remain Dependent on **Cytokines for Growth Following Editing**



Summary and Conclusion

- due to the editing process have occurred

Figure 3: High Efficiency CRISPR/Cas9 Gene Editing to Produce Allogeneic Anti-BCMA CAR-T Cells

To examine whether editing results in transformations that confer cytokine-independent growth to T cells, cytokines and/or serum were removed from the growth media 2 weeks following editing and CAR knock-in. No further proliferation of T cells was observed in the absence of cytokines, indicating that the T cells remained dependent on cytokines for growth.

Activity In Vitro



Figure 6: Allogeneic Anti-BCMA CAR-T Cells Display Robust Anti-Tumor Activity in a Mouse Model



Using CRISPR/Cas9 gene editing, we have generated an allogeneic anti-BCMA CAR-T product at high efficiency, with over 60% of the cells harboring all 3 desired edits The CAR-T cells maintain a similar CD4/CD8 ratio compared to controls, as well as characteristic cytokine dependency, suggesting neither abnormal tonic signaling from CAR insertion nor transformation

The CAR-T cells selectively kill BCMA⁺ cells and secrete T cell activation cytokines following encounter with BCMA-expressing cells The CAR-T cells eradicate MM cells in a subcutaneous RPMI-8226 tumor xenograft model, confirming potent activity in vivo



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Figure 5: Allogeneic Anti-BCMA CAR-T Cells Show Potent Yet Specific