## CLAIMS

1. A plasmid for use with clustered regularly interspaced short palindromic repeat (CRISPR)/ CRISPR-associated 9 (Cas9) integration systems wherein the plasmid comprises in order a left homology arm, a splice acceptor, a transgene, and a right homology arm, and further comprises a polyadenylation signal between the transgene and the right homology arm;

wherein the left and right homology arms specifically hybridize to an adeno-associated virus integrated site q (AAVS1) of chromosome 19 of humans and each has a length of 30-1000bp, and

wherein the right homology arm comprises a nucleic acid sequence selected from SEQ ID NO: 1, SEQ ID NO: 3, SEQ ID NO: 5, or SEQ ID NO: 7, and left homology arm comprises a nucleic acid sequence selected from SEQ ID NO: 2, SEQ ID NO: 4, SEQ ID NO: 6, or SEQ ID NO: 8.

2. The plasmid of claim 1, wherein the left homology arm and right homology arm are the same length.

3. The plasmid of claim 2, wherein the homology arms are 30bp in length, 300bp in length, 500bp in length, or 800bp in length.

4. The plasmid of claim 1, wherein the left homology arm and right homology arm are different lengths.

5. The plasmid of any one of claims 1-4, wherein the transgene comprises a polynucleotide encoding a chimeric antigen receptor for a tumor antigen.

6. An Adeno-associated viral (AAV) vector comprising the plasmid of any one of claims 1-5.

7. The AAV vector of claim 6, wherein the serotype of the AAV comprises AAV6.

8. The AAV vector of claim 6 or 7, wherein the vector further comprises a plasmid encoding a crRNA, tracer RNA (trcrRNA), and a CAS endonuclease.

9. The AAV vector of any of claims 6-8, wherein the vector is a single stranded AAV (ssAAV) and/or a self-complimentary AAV (scAAV).

10. A modified natural killer (NK) cell comprising the plasmid of any one of claims 1-5 or the AAV vector of any one of claims 6-9.

11. The modified NK cell of claim 10, wherein the modified cell is a chimeric antigen receptor (CAR) NK cell that targets a cancer antigen.

12. Use of the modified NK cell of claim 11 for the manufacture of a medicament for the treatment of a subject with a cancer.

13. An ex vivo method of preparing a genetically modified natural killer (NK) cell by homologously directed repair comprising

a) obtaining a ribonucleoprotein (RNP) complex comprising a class 2 CRISPR/Cas endonuclease complexed with a corresponding CRISPR/Cas guide RNA and an adeno-associated virus (AAV) vector comprising a plasmid comprising a transgene that includes a chimeric antigen receptor for a tumor antigen; wherein the transgene is flanked by a left homology arm and a right homology arm, wherein the left homology arm and the right homology arm have a length of 30bp to 800bp, and wherein the right homology arm comprises a nucleic acid sequence selected from SEQ ID NO: 1, SEQ ID NO: 3, SEQ ID NO: 5, or SEQ ID NO: 7 and the left homology arm comprises a nucleic acid sequence selected from SEQ ID NO: 2, SEQ ID NO: 4, SEQ ID NO: 6, or SEQ ID NO: 8; and

b) introducing the transgene and the RNP complex into the NK cell; wherein the transgene is introduced into the NK cell via infection with the AAV into the cell; wherein the RNP complex hybridizes to a target sequence within the genomic DNA of the NK cell and the NK cell's DNA repair enzymes insert the transgene into the host genome at the target sequence within the genomic DNA of the NK cell thereby creating a modified NK cell.

14. The method of claim 13, wherein the NK cell is a primary or expanded cell.

15. The method of claim 14, wherein the NK cell is a primary cell and the primary cell is incubated for 4 days in the presence of IL-2 prior to infection.

16. The method of claim 15, wherein the NK cell is a primary cell that is further expanded for 4 days in the presence of irradiated feeder cells prior to infection.

17. The method of any one of claims 13-15, further comprising expanding the NK cell with irradiated mbIL-21 expressing feeder cells, PM2l particles, EX2l exosomes, or any combination thereof following infection.

18. The method of any one of claims 13-17, wherein the RNP complex is introduced into the NK cell via electroporation, or wherein the RNP complex is introduced into the NK cell via transfection and the RNP complex is encoded on the same or a different AAV.

19. An ex vivo method of preparing a genetically modified natural killer (NK) cell by homology directed repair (HDR) comprising:
(a) introducing into the NK cell (i) a ribonucleoprotein (RNP) complex comprising a class 2 CRISPR/Cas endonuclease complexed with a corresponding CRISPR/Cas guide RNA , and (ii) an AAV vector comprising a plasmid comprising a transgene flanked by first and second homology arms each having a length of no more than 800bp, wherein the first homology arm comprises a nucleic acid sequence selected from SEQ ID NO: 1, SEQ ID NO: 3, SEQ ID NO: 5, or SEQ ID NO: 7 and the second homology arm comprises a nucleic acid sequence selected from SEQ ID NO: 2, SEQ ID NO: 4, SEQ ID NO: 6, or SEQ ID NO: 8, wherein the transgene encodes a chimeric antigen receptor; and
(b) maintaining the NK cell for a time and under conditions sufficient for (i) the RNP complex to hybridize to the target sequence and introduce a double-stranded break (DSB) in the target sequence; and (ii) the NK cell to insert the transgene into its genome at the target sequence and repair the break by HDR, thereby incorporating the transgene into the NK cell genome at the site of the DSB.

20. The method of claim 19, wherein the NK cell is a primary or an expanded cell.

21. The method of claim 19 or 20, wherein the class 2 CRISPR/Cas endonuclease is Cas9.

22. The method of claim 19 or 20, wherein the NK cell is a primary cell and the method

further comprises incubating the NK cell at least 4 days in the presence of membrane bound IL-21 (mbIL-21) prior to step (a).

23. The method of claim 19 or 20, wherein the NK cell is a primary cell and the method further comprises expanding the NK cell for at least 4 days in the presence of mbIL-21 after step (b).

24. The method of claim 23, wherein the mbIL-21 comprises irradiated mbIL-21 expressing feeder cells, PM21 particles, EX21 exosomes, or any combination thereof.

25. A plasmid for use with clustered regularly interspaced short palindromic repeat (CRISPR)/CRISPR-associated 9 (Cas9) integration systems, wherein the plasmid comprises a sequence selected from SEQ ID NO: 13, SEQ ID NO: 14, SEQ ID NO: 15, or SEQ ID NO: 16.