

What is claim is:

1. A mammalian cell comprising a site for integration of a gene of interest, wherein the cell comprises a first Stable Integration Site located in a Genomic Safe Harbor and a second Stable Integration Site that is not located in the Genomic Safe Harbor, wherein the first Stable Integration Site comprises a DNA polynucleotide comprising in 5' to 3' order a first recombinase recognition site (RRS), a first reporter gene encoding a first reporter protein, a Cas9 gene and a second RRS; and wherein the second Stable Integration Site comprises a DNA polynucleotide comprising in 5' to 3' order a first Genomic Safe Harbor homology arm containing a CRISPR single guide RNA (sgRNA) target site, a third RRS, a second reporter gene encoding a second reporter protein, a fourth RRS and a second Genomic Safe Harbor homology arm containing an CRISPR sgRNA target site, wherein the first RRS, the second RRS, the third RRS and the fourth RRS are different, and wherein the first reporter protein and the second reporter protein are different.
2. The mammalian cell according to claim 1, wherein the mammalian cell is a human cell.
3. The mammalian cell according to claim 2, wherein the human cell is a Human Amniotic Epithelial Cell.
4. The mammalian cell according to claim 2, wherein the human cell is a HEK293 Cell.
5. The mammalian cell according to claim 1, wherein the mammalian cell is a CHO cell.

6. The mammalian cell according to claim 1, wherein the second Stable Integration Site is located in a second Genomic Safe Harbor that is different from the first Genomic Safe Harbor.

7. The mammalian cell according to claim 1, wherein the second Stable Integration Site is located in a region that is not a Genomic Safe Harbor.

8. A mammalian cell comprising a site for integration of a gene of interest, wherein the cell comprises a first Stable Integration Site located in a Genomic Safe Harbor and a second Stable Integration Site that is not located in the Genomic Safe Harbor, wherein the first Stable Integration Site comprises a first DNA polynucleotide encoding a first reporter protein, wherein the first DNA polynucleotide comprises in 5' to 3' order a first recombinase recognition site (RRS), a first reporter gene encoding the first reporter protein, a Cas9 gene and a second RRS; and wherein the second Stable Integration Site comprises a second DNA polynucleotide encoding a second reporter protein, wherein the second DNA polynucleotide comprises in 5' to 3' order a first Genomic Safe Harbor homology arm containing a CRISPR single guide RNA (sgRNA) target site, a third RRS, a second reporter gene encoding the second reporter protein, a fourth RRS and a second Genomic Safe Harbor homology arm containing an CRISPR sgRNA target site, and wherein the first RRS, the second RRS, the third RRS and the fourth RRS are different.

9. The mammalian cell according to claim 8, wherein the second Stable Integration Site is located in a second Genomic Safe Harbor that is different from the first Genomic Safe Harbor.

10. The mammalian cell according to claim 8, wherein the second Stable Integration Site is located in a region that is not a Genomic Safe Harbor.

11. The mammalian cell according to claim 8, wherein the mammalian cell is a human cell.

12. The mammalian cell according to claim 8, wherein the mammalian cell is a CHO cell.

13. A method for making at least one protein of interest, comprising:

(a) providing a mammalian cell comprising a site for integration of a gene of interest, wherein the cell comprises a first Stable Integration Site located in a Genomic Safe Harbor and a second Stable Integration Site that is not located in the Genomic Safe Harbor, wherein the first Stable Integration Site comprises a DNA polynucleotide comprising in 5' to 3' order a first recombinase recognition site (RRS), a first reporter gene encoding a first reporter protein, a Cas9 gene and a second RRS; and wherein the second Stable Integration Site comprises a DNA polynucleotide comprising in 5' to 3' order a first Genomic Safe Harbor homology arm containing a CRISPR single guide RNA (sgRNA) target site, a third RRS, a second reporter gene encoding a second reporter protein, a fourth RRS and a second Genomic Safe Harbor homology arm containing an CRISPR sgRNA target site, wherein the first RRS, the second RRS, the third RRS and the fourth RRS are different, and wherein the first reporter protein and the second reporter protein are different;

(b) introducing a polynucleotide encoding the protein of interest into a Stable Integration Site by recombinase mediated cassette exchange, and

(c) culturing the mammalian cell under conditions that allow expression of the polynucleotide encoding the polynucleotide of interest.

14. The method according to claim 13, wherein the mammalian cell is a Human Amniotic Epithelial Cell.

15. The method according to claim 13, wherein the mammalian cell is a HEK293 cell.

16. The method according to claim 13, wherein the mammalian cell is a CHO cell.

17. The method according to claim 13, wherein the second Stable Integration Site is located in a second Genomic Safe Harbor that is different from the first Genomic Safe Harbor.