

Supplementary Information

Targeted insertion of an anti-CD2 monoclonal antibody transgene into the *GGTA1* locus in pigs using *FokI*-dCas9

Mark B. Nottle, Evelyn Salvaris, Nella Fisicaro, Stephen McIlfatrick, Ivan Vassiliev, Wayne J. Hawthorne, Philip J. O'Connell, Jamie L. Brady, Andrew M. Lew, Peter J. Cowan

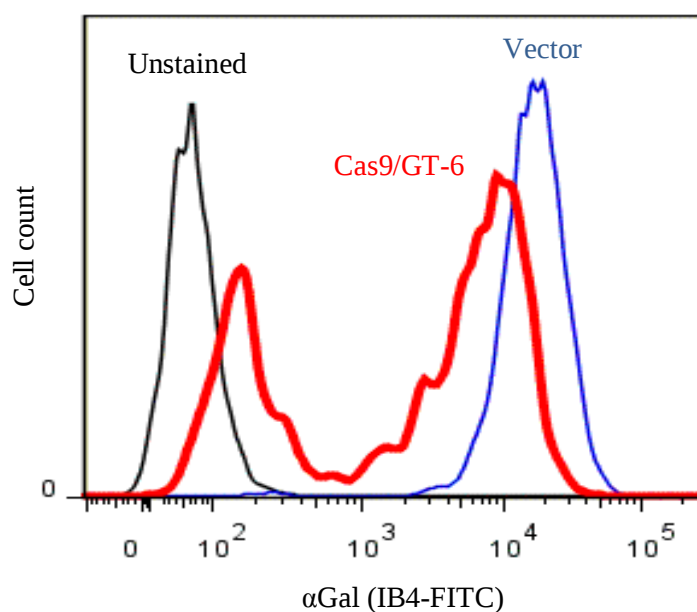
Supplementary Figures 1-6

Supplementary Tables 1 and 2

Supplementary Figure 1

Targeting of *GGTA1* in WT pig fetal fibroblasts using Cas9

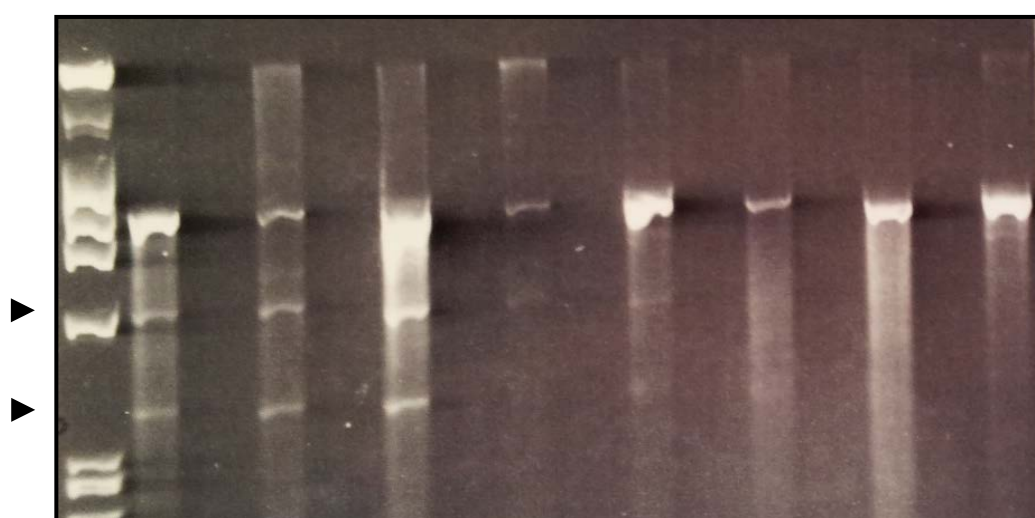
WT fibroblasts were co-transfected with expression vectors for Cas9 (pGS-CMV-hCas9, GenScript) and guide RNA GT-6. Flow cytometric analysis of the cells at day 3 (red line) demonstrated that ~30% were α Gal-negative; blue line, control vector-transfected WT cells; black line, unstained WT cells.



Supplementary Figure 2

Surveyor nuclease assay to detect indels in *GGTA1* in pig cells

WT fibroblasts were transfected with the combinations of expression vectors shown. pGS-CMV-hCas9 was from GenScript, and pCas9-GFP was from Addgene (plasmid #44719). Three days after transfection, genomic DNA was isolated and a 970 bp region of *GGTA1* encompassing the target site was amplified by PCR using primers SN-F2 and SN-R2. The 970 bp PCR product was analysed using the Surveyor Mutation Detection kit (Transgenomic, Omaha, NE). Bands of approximately 592 bp and 378 bp (arrowheads) indicate cleavage at the expected target site. *FokI*-dCas9 generated mutations with both pairs of guide RNAs, albeit with a lower efficiency than Cas9 with a single guide RNA. Left lane: molecular weight markers (λ HindIII + ϕ X174/HaeIII).



pGS-CMV-hCas9	+	+	-	-	-	-	-	-
pCas9-GFP	-	-	+	+	-	-	+	-
<i>FokI</i> -dCas9	-	-	-	-	+	+	-	-
gRNA	5	6	5	6	1+2	3+4	-	-

Supplementary Figure 3

Sequence of donor DNA template for the generation of anti-CD2 mAb knock-in pig fibroblasts

GCGGCCGCTGCAACTGCAATTAGACCCCTAGTCTGGGAACCTCCATATGCTGCGGATGTGGCCCTAAA
AAGACAAAAAATAAAAAATTAATAAGTGCATCACATCAACACTTTGCACACCTTAACTTACACAA
TGATATATGTCAGTTATATCTGAATAATGCAGGAAAAAAGGAAGAACTAGCATCTGAACTGATTGATA
AGGACAAGGGGATAAAGACTAATTAAGGCAGATGGTAATGAGTTATCACCTTCATTACAGTTAAAGCT
GGTGTTGGAGTATGTGGCTTATAACTGCAGGCCCTGGGCTTCCAAATCTTGAACCACAGAATTAGAAA
GACTTAGGCATCCAGGCACATACAGCACAAGTAACTGCAGGCCTCTGGATGTGGGAGCAGGGCCTGTG
CCACTAAGGTACCACTGCAAGGCCCAGAGCAGCTGAAAACACATGTTCTCTCTGCCTGGTTGGCTTCC
AAGAGTGAGAGAGGAAGGAGCAGGGCTGAGCATGCCCAGCCACCCTGCCAGAATCACCAGTCAGGTAA
GCCACTCCACCTCCCCAAAGCTGAATGACTGAATGGTGGAGAGTAGCTGGGAATGTTACAGCAACAGA
CGTCTCTCATCCAGGATGGGGAAAAATCATTCTTTCTTAACTGCAAAATACAGACTAGATGATAAT
AGCATATTGTCTCCTCTAGAAATCCCAGAGGTTACATTTACCCCATTTCTTTATTTTCAGATACATT
GAGCATTACTTGGAGGAGTTCTTAATATCTGCAAATACATACTTCATGGTTGGCCACAAAGTCATCTT
TTACATCATGGTGGATGATATCTCCAGGATGCCTTTGATAGAGCTGGGTCTCTGCGTTCCTTTAAAG
TGTTTGAGATCAAGTCCGAGAAGAGGTGGCAAGACATCAGCATGATGCGCATGAAGACCATCGGGGAG
CACATCCTGGCCCACATCCAGCACGAGGTGGACTTCCTCTTCTGCATGGACGTGGATCAGGTCTTCCA
AAACAACCTTTGGGGTGGAGACCCTGGGCCAGTCGGTGGCTCAGCTAAGATCTATAACTTCGTATAGCA
TACATTATACGAAGTTATAATTCTACCGGGTAGGGGAGGCGCTTTTCCCAAGGCAGTCTGGAGCATGC
GCTTTAGCAGCCCCGCTGGCACTTGGCGCTACACAAGTGGCCTCTGGCCTCGCACACATTCCACATCC
ACCGGTAGCGCCAACCGGCTCCGTTCTTTGGTGGCCCTTCGCGCCACCTTCTACTCCTCCCCTAGTC
AGGAAGTTCCCCCGCCCCGAGCTCGCGTCGTGCAGGACGTGACAAATGGAAGTAGCACGTCTCAC
TAGTCTCGTGCAGATGGACAGCACCGCTGAGCAATGGAAGCGGGTAGGCCTTTGGGGCAGCGGCCAAT
AGCAGCTTTGCTCCTTCGCTTTCTGGGCTCAGAGGCTGGGAAGGGGTGGGTCCGGGGGCGGGCTCAGG
GGCGGGCTCAGGGGCGGGGCGGGCGGAAGGTCCTCCGGAGCCCGGCATTCTCGCACGCTTCAAAGC
GCACGTCTGCCGCGCTGTTCTCCTCTTCCTCATCTCCGGGCCTTTGGGCTGCAGCCACCATGATTGAA

CAAGATGGATTGCACGCAGGTTCTCCGGCCGCTTGGGTGGAGAGGCTATTCGGCTATGACTGGGCACA
ACAGACAATCGGCTGCTCTGATGCCGCCGTGTTCCGGCTGTCAGCGCAGGGGCGCCCGGTTCTTTTTG
TCAAGACCGACCTGTCCGGTGCCCTGAATGAACTGCAGGACGAGGCAGCGCGGCTATCGTGGCTGGCC
ACGACGGGCGTTCTTGCGCAGCTGTGCTCGACGTTGTCACTGAAGCGGGAAGGGACTGGCTGCTATT
GGGCGAAGTGCCGGGGCAGGATCTCCTGTCATCTCACCTTGCTCCTGCCGAGAAAGTATCCATCATGG
CTGATGCAATGCGGCGGCTGCATACGCTTGATCCGGCTACCTGCCATTTCGACCACCAAGCGAAACAT
CGCATCGAGCGAGCACGTACTCGGATGGAAGCCGGTCTTGTCGATCAGGATGATCTGGACGAAGAGCA
TCAGGGGCTCGCGCCAGCCGAACTGTTGCCAGGCTCAAGGCGCGCATGCCCGACGGCGAGGATCTCG
TCGTGACCCATGGCGATGCCTGCTTGCCGAATATCATGGTGGAAAATGGCCGCTTTTCTGGATTATC
GACTGTGGCCGGCTGGGTGTGGCGGACCGCTATCAGGACATAGCGTTGGCTACCCGTGATATTGCTGA
AGAGCTTGGCGGCGAATGGGCTGACCGCTTCTCGTGCTTTACGGTATCGCCGCTCCCGATTTCGACGC
GCATCGCCTTCTATCGCCTTCTTGACGAGTTCTTCTGAGCGGGACTCTGGGGTTCGAAATGACCGACC
AAGCGACGCCCAACCTGCCATCACGATGGCCGCAATAAAATATCTTTATTTTCATTACATCTGTGTGT
TGGTTTTTTGTGTATAACTTCGTATAGCATACATTATACGAAGTTATAGATCTGCTCGAGGCCGGCCT
TTCAATATTGGCCATTAGCCATATTATTTCATTGGTTATATAGCATAAATCAATATTGGCTATTGGCCA
TTGCATACGTTGTATCTATATCATAATATGTACATTTATATTGGCTCATGTCCAATATGACCGCCATG
TTGGCATTGATTATTGACTAGTTATTAATAGTAATCAATTACGGGGTCATTAGTTCATAGCCCATATA
TGGAGTTCCGCGTTACATAACTTACGGTAAATGGCCCGCCTGGCTGACCGCCCAACGACCCCCGCCCA
TTGACGTCAATAATGACGTATGTTCCCATAGTAACGCCAATAGGGACTTTCCATTGACGTCAATGGGT
GGAGTATTTACGGTAAACTGCCCACTTGGCAGTACATCAAGTGTATCATATGCCAAGTCCGCCCCCTA
TTGACGTCAATGACGGTAAATGGCCCGCCTGGCATTATGCCAGTACATGACCTTACGGGACTTTACT
ACTTGGCAGTACATCTACGTATTAGTCATCGCTATTACCATGGTGATGCGGTTTTGGCAGTACACCAA
TGGGCGTGGATAGCGGTTTGACTCACGGGGATTTCCAAGTCTCCACCCCATGACGTCAATGGGAGTT
TGTTTTGGCACCAAAATCAACGGGACTTTCCAAAATGTCGTAACAACTGGCCGGCCTTAATTAAGGCG
CGCCTCATAGACTCATGTGTTTGAAAGCTTGGCCACAAGGAGTGGAATGTTAGGAGTGTGACTGTG
TTAGAGGAAGTGTGTCACTTTGAGGTTGGCCTTTGAGGTCTCCTATGCTCATGAGCTGCCAGTGTGG
AATGGAAGCCTCTTCCTGGCTGCATTTGGATCAAGATGTAGAAGTCTCCCCTTCTCCAGCACCATGTC
TACCTGCACACTGGCATGCTTCCTGTCATGATGGACTAAATCCCTGAAACTGTAAGCCAGACCCTATT

AAATGTCTCCCTTTAGAAAGGAATGCCTTGGTCATGGTGTTCCTTCACAGCAACAAAACCAAACTAAC
ACACCCAGCACTTAGGATGCAAAGGCAGATGTATCTCTGTGAGTTCAAGGGCAGCCTGATCTACATAA
TGAATTACAGAATAGCCAGGGCTGTGCAGATAGACCTTGTCTAAACAAGCAAACAAACAAAACCCATA
GTAAAATAAGAAGAAAAGTAATGGAGACAGATCCTTTGTGTAGCTGCCTAACCCCAACAGCATCCCTA
GGTTGATCAGGAAGAAACCTAAGAGCATCAAGTCCCCATGGAAAATGCTCACAACATCTAATAAGAGG
GGTTCAGGGAGACAGGGTCTGCTGAATCCTGGTGAGGCTGGGGGGCTGCTGTGGGGATCTGCATGGGG
AGGTGACTGGAGTTCACACCTGTGGTCACACGCATTCAATTGCTCTGTGTGATGAGTGACTCTCTTCAG
GAGTCCCCACCCATCTATGCTCTTCTTTCCACTTCACTTCTGTGACTGATTTGAAAAGATCCACAGAA
TCATGAAATGGGTAAATTTAATCTTTCCCCACACTACTGATCAGGGATGAAACCTCACATCACAGTGT
GTGCTCTCTGGCATGAGAATCATCTTTCTCCAGTGTCACACTGCACAGGCCTGAGGAACTCTGGGG
CCAACTCAATTATCAGAACCCGTGTCCACACAGCTGTGACATTCATGTCCCTCACCTCATCAGCTCA
AAAGTGTGAGTGAAGGTCAGGGAGGGAGGGAGAGGGTCAGAGATTACATAACTTCTACCACACTAGCA
TCCTTATTCTGTGTTATAGTTGGTGACAGCTCCTCTCCTACTGTGATTGAGTCACTAAAGCAACTGCA
CCATGGAAGTGCAGACACAGAGAAGTCCAGTGAGACAGGGATCAGTGCCCTGCTGTCTGTGGTCTGC
ACGGGGTCTCTCTTCAGTGGACAAGGGGGTCTCCTGTGCTGAGACAATGTCCCAGATCCACAGAGACA
AACTCAGGACTCAGAATGAAGATCCTTGTTTCAAATACACACACACACACATACACACACACACAC
ACACACACACACACACACACACACCTGCAGCCACAGACTTTTTCATCTAGGAATTAACACAAGGAATCT
GTGTCTCAGCACAGGGCTGAGAAGACAGATCCTGAGGGGAGAGGCAAAGTCTACACTTGACAGATGAG
AGTCCTGCACTCAGGCTTGCCAGTGTGAGCCGCCATTGCAGGTGAACAGAGCCTGGTCTCTGTGGGG
TCCCTGTGGGGCTTGACAGCCAGCGCCTTGACTTTAAGGAAAAGCCTCTCTCTCCACTGCATCCCTAA
GCGCTTGTGTGCGCCATTGTATTCCCGGAAGAGGCTTTTCTTCTAGAAGACTCCAGGGTCTGACTTCTG
AAGAGAAGAAGAAAGAGGAAGAGTGGAAGAGAGGACACAGAGAGTCTGGCCTGCGGGTCTCTCCTGGT
GTTTTGAGAGTTTCTGGATCAGAACTCGGAGACGACAGCACAGGGTTCAGGCAAAGTCTTAGTCGCCA
GGCAGTGAGGTCAGGGGTGGGGAAGCCCAGGGCTGGGGATTCCCCATCTCCACAGTTTCACTTCTGCA
CCTAACCTGGGTGAGGTCCTTCTGTCCGGACACTGTTGACGCGCAGTCAGCTTTACCCCCATTGGGT
GGCGCGATCACCAAGAACCAATCAGTGTGCGCGGGACGCTGGATATAAAGTCCACGCAGCCCGCAGA
ACTCAGAAGTCGATCAAGCTTATCGATTGGGGTGAGTACTCCCTCTCAAAAGCGGGCATGACTTCTGC
GCTAAGATTGTCAGTTTCCAAAAACGAGGAGGATTTGATATTACCTGGCCCGCGGTGATGCCTTTGA

GGGTGGCCGCGTCCATCTGGTCAGAAAAGACAATCTTTTTGTTGTCAAGCTTGAGGTGTGGCAGGCTT
GAGATCTGACCATACTTGAATGACAATGACATCCACTTTGCCTTTCTCTCCACAGGCGCGCCGGAT
CCGAATTC-coding region for anti-CD2 monoclonal antibody (2,367 bp)-
GAATTCCGACTGTGCCTTCTAGTTGCCAGCCATCTGTTGTTTGCCCTCCCCGTGCCTTCCTTGACC
CTGGAAGGTGCCACTCCCCTGTCCTTTCCTAATAAAATGAGGAAATTGCATCGCATTGTCTGAGTAG
GTGTCATTCTATTCTGGGGGGTGGGGTGGGGCAGGACAGCAAGGGGGAGGATTGGGAAGACAATAGCA
GGCATGCTGGGGATGCGGTGGGCTCTATGGGTGGTACAAGGCACATCCTGACGAGTTCACCTACGAGA
GGCGGAAGGAGTCCGCAGCCTACATTCCGTTTGGCCAGGGGGATTTTTATTACCACGCAGCCATTTTT
GGGGGAACACCCACTCAGGTTCTAAACATCACTCAGGAGTGCTTCAAGGGAATCCTCCAGGACAAGGA
AAATGACATAGAAGCCGAGTGGCATGATGAAAGCCATCTAAACAAGTATTTCTTCTCAACAAACCCA
CTAAAATCTTATCCCCAGAATACTGCTGGGATTATCATATAGGCATGTCTGTGGATATTAGGATTGTC
AAGATAGCTTGGCAGAGAAAAGAGTATAATTTGGTTAGAAATAACATCTGACTTTAAATTGTGCCAGC
AGTTTTCTGAATTTGAAAGAGTATTACTCTGGCTACTTCCTCAGAGAAGTAGCACTTAATTTTAACTT
TTAAAAAATACTAACAAAATACCAACACAGTAAGTACATATTATTCTTCCTTGCAACTTTGAGCCTT
GTCAAATGGGAGAATGACTCTGTGGTAATCAGATGTAAATCCCAATGATTTCTTATCTGTTCTGGGT
TGAGGGGGTATATACTATTAAGTGAACCAAAAAAAAAAATTGTCATAGGCAAAGAAAAAGTCAGAGACA
CTCTACATGTCATACTGGAGAAAAGTATGCAAAGGGAAGTGTGTTGGCAACAAAATAAGATTGGGAGGG
GTCGTCCTCTTGATTTTAGCGGCCGC

Supplementary Figure 4

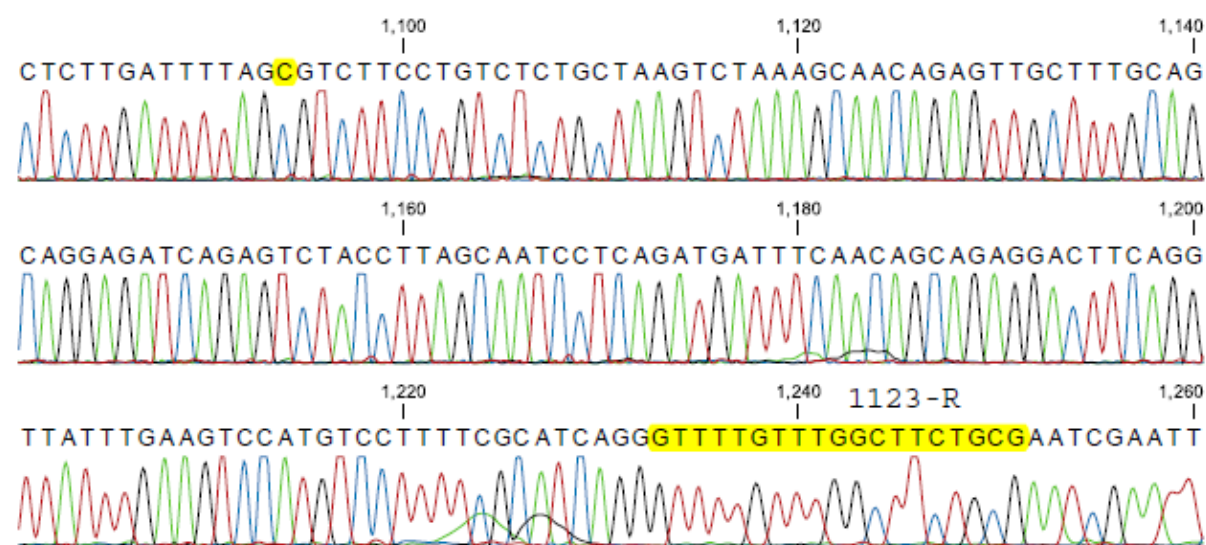
Sequencing chromatograms of the junction regions of the targeted *GGTA1* allele in anti-CD2 mAb knock-in clone #3

(A) Sequence of 5' junction. Highlighted in yellow are (i) the recognition site for primer UKI-F3, which lies upstream of the targeting construct, and (ii) the first base of the 5' homology arm in the targeting construct. (B) Sequence of 3' junction. Highlighted in yellow are (i) the last base of the 3' homology arm in the targeting construct, and (ii) the recognition site for primer 1123-R, which lies downstream of the targeting construct.

A



B



Supplementary Figure 5

Deletions detected within *GGTA1* in WT pig fibroblasts edited using *FokI*-dCas9 with guide RNAs GT-3 and GT-4

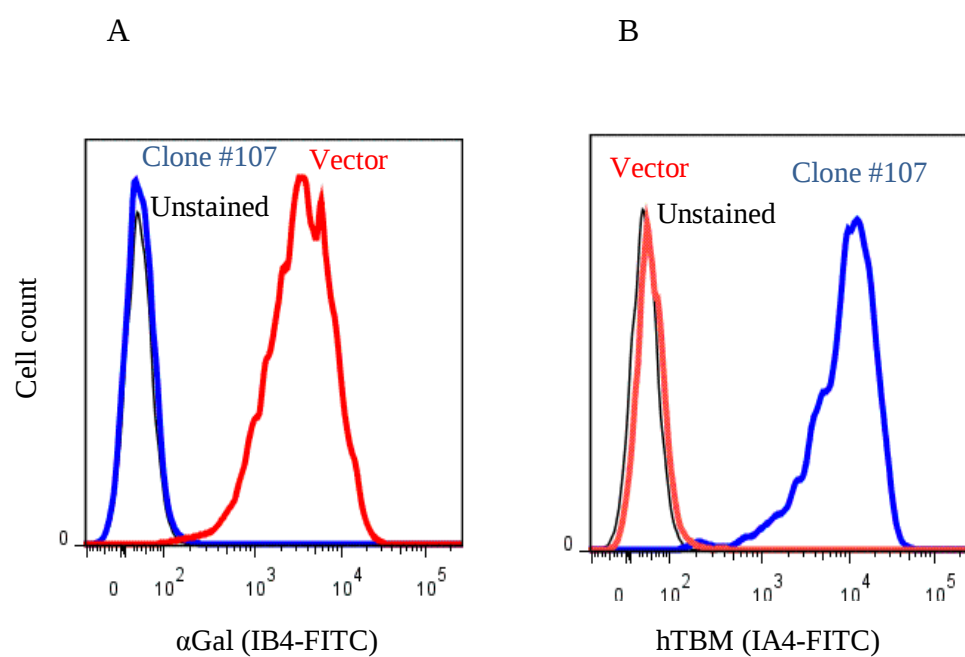
The WT sequence (top) shows the recognition sites for GT-3 and GT-4 (underlined). **(A)** Clone isolated during the initial testing of GT-3 and GT-4, showing a 25 bp deletion. **(B)** Anti-CD2 knock-in clone #3, showing a 43 bp deletion. **(C)** hTBM knock-in clone #107, showing a 14 bp deletion.

```
WT  GTCGGTGGCTCAGCTACAGGCCTGGTGGTACAAGGCACATCCTGACGAGTTCACCTACGAGAGG
A   GTCGGTGGCTCAGCTACAGGCCTGGTGGT-----CTACGAGAGG
B   GTCGGTGGCT-----CCTACGAGAGG
C   GTCGGTGGCTCAGCTACAGGCC-----ACATCCTGACGAGTTCACCTACGAGAGG
```

Supplementary Figure 6

Absence of α Gal and expression of hTBM in hTBM knock-in clone #107

(A) α Gal and (B) hTBM staining. Blue line, clone #107; red line, vector-transfected WT cells; black line, unstained WT cells.



Supplementary Table 1

Single guide RNAs targeting *GGTA1*

Name	Sequence 5'-3'	PAM
GT-1	CTGAGCCACCGACTGGCCC	AGG
GT-2	TGACGAGTTCACCTACGAG	AGG
GT-3	CTGTAGCTGAGCCACCGAC	TGG
GT-4	CGAGTTCACCTACGAGAGG	CGG
GT-5	GGTGGCTCAGCTACAGGCC	TGG
GT-6	CAGTCGGTGGCTCAGCTAC	AGG

PAM: protospacer adjacent motif

Supplementary Table 2

PCR primers

Target	Forward 5'-3' (primer name)	Reverse 5'-3' (primer name)	Size (bp)
<i>GGTA1</i>	GCTGCAACTGCAATTAGACC (GTFS-F1)	GCTAAAATCAAGAGGACGACC (GTFS-R1)	1806
<i>GGTA1</i>	CCTCTAGAAATCCCAGAGGTTAC (SN-F2)	CCCTCAACCCAGAACAGATAAG (SN-R2)	970
Knock-in 5' junction	GGTGTAGGTCATAGACAAAGC (UKI-F3)	AGAAAGCGAAGGAGCAAAG (UKI-R2)	1506
Knock-in 3' junction	GAGGATTGGGAAGACAATAGC (117-F)	CGCAGAAGCCAAACAAAAC (1123-R)	947
<i>FokI</i> -dCas9 vector	GAAGATGGATGGGACGGA (fCas9-F1)	CTTTTGTATGGCTGGCGA (fCas9-R1)	1063
gRNA vector (pGS- U6-gRNA)	ATCCTCATCCTGTCTCTT (gRNA-F2)	GTATGTTGTGTGGAATTGTG (gRNA-R1)	1518