



MODEL GENERATION TECHNICAL REPORT

**Generation of a RFP657-LF2A-Cre-F3-ERT2-F3 into Prg2
Knock-In mouse line in a pure BALB/cN genetic background**

Project code: Ros6280 / IR6280

Report updated: 15/02/2023

1 PROJECT DESCRIPTION

2 GENETIC STRATEGY

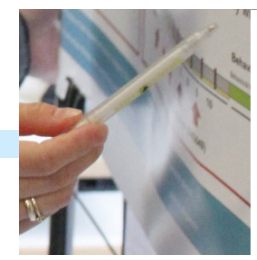
3 HOMOLOGOUS RECOMBINATION
VECTOR CONSTRUCTION

4 ES ELECTROPORATION & SCREENING OF
RECOMBINANT CLONES

5 MICROINJECTION & BREEDING

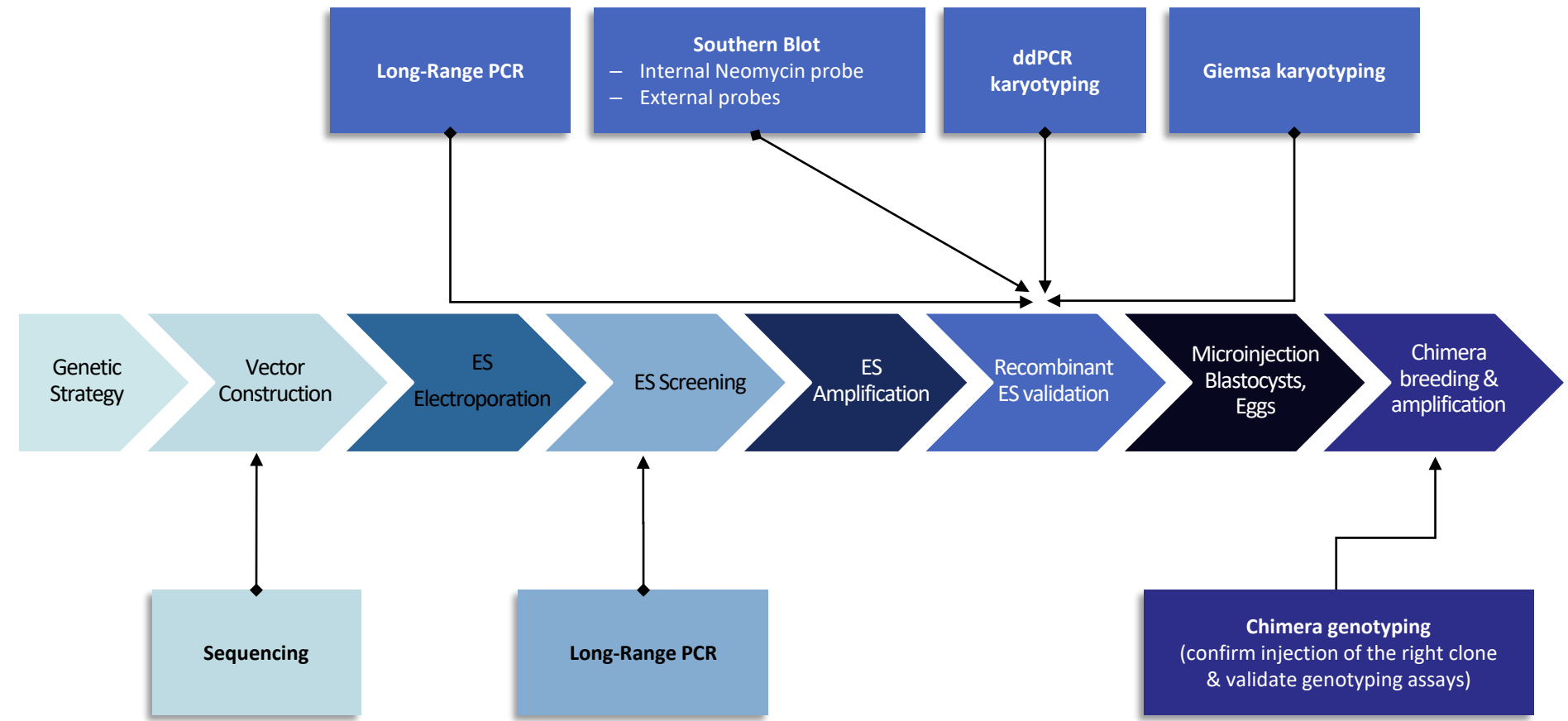
6 SEQUENCE OF THE DELIVERED ALLELE

1 PROJECT DESCRIPTION

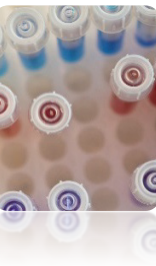


- Aim
- Project process & quality controls

Project process & quality controls



2 GENETIC STRATEGY

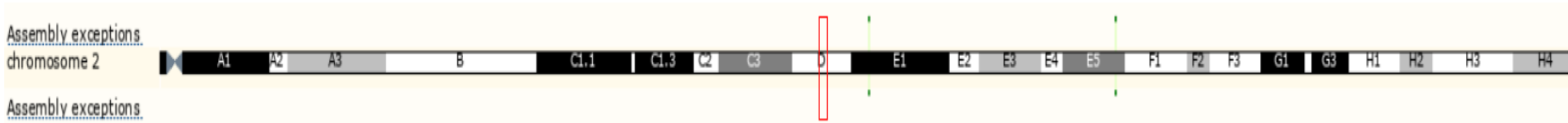


- Target locus structure
- mRNA and protein
- Genetic strategy
- PRO & CONS evaluation of the strategy

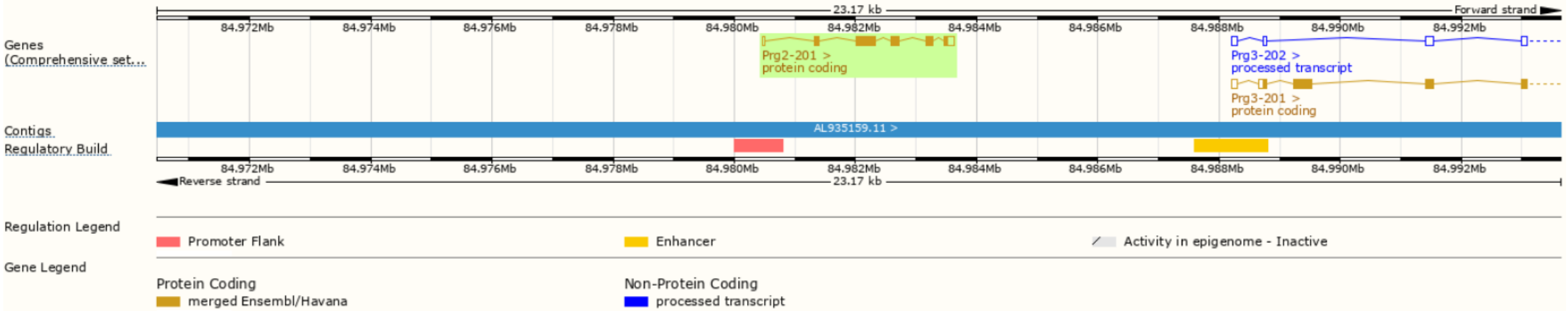
■ mouse genomic locus – structure



Chromosome 2: 84,980,461-84,983,632



Ensembl Gene ID : ENSMUSG00000027073



■ mRNA and protein

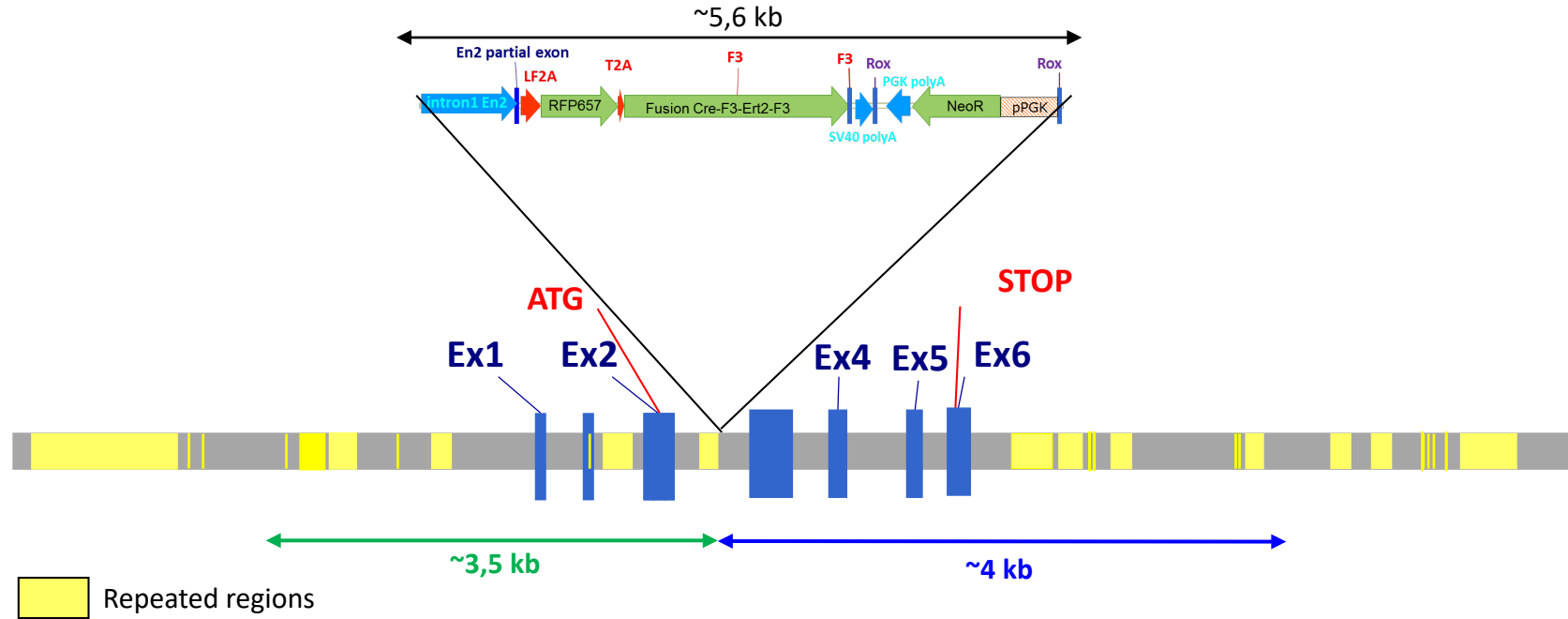


Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	RefSeq	Flags
Prg2-201	ENSMUST00000028467.5	838	223aa	Protein coding	CCDS16198	Q545D8 Q61878	NM_008920 NP_032946	TSL:1GENCOD E basicAPPRIS P1

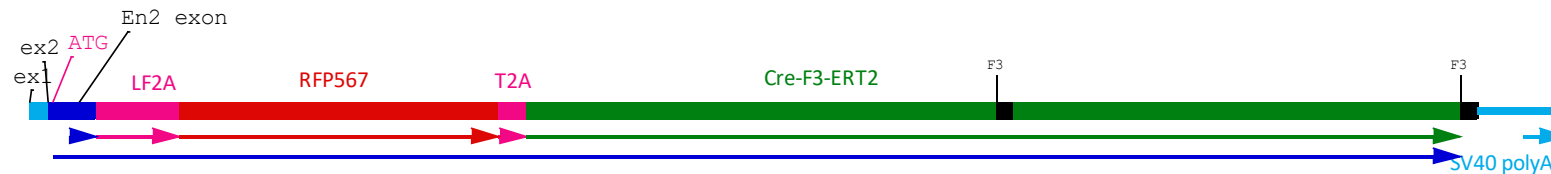
Strategy



Approach: Homologous recombination in BALB/cN ES cells



mRNA and protein expressed after Cre mediated excision (sequence detail see next page)



■ PROs& CONs evaluation of the strategy



■ Pros

- Use of the TagRFP657 and dual cassette (Cre-F3-ER^{T2}-F3)

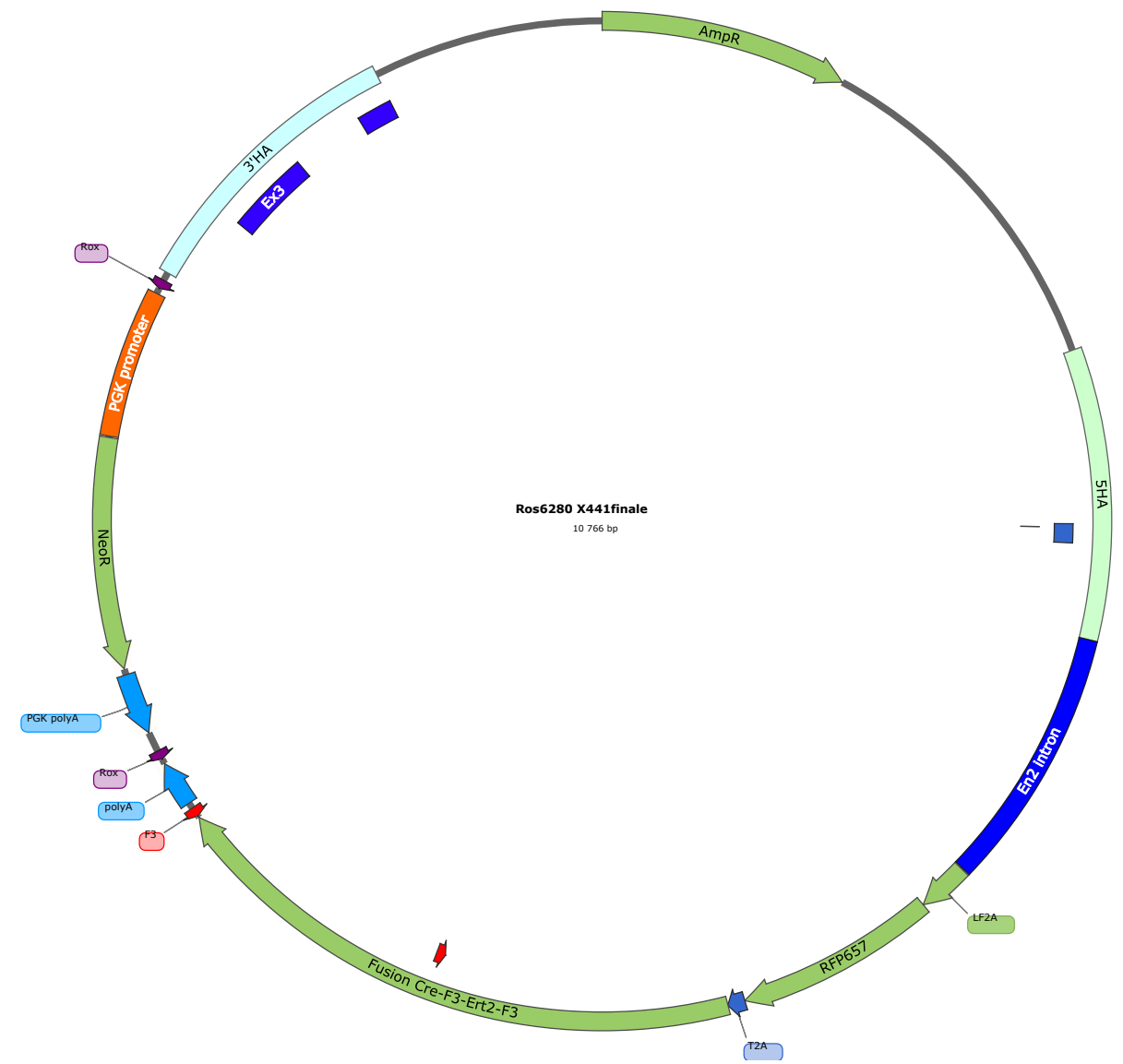
■ Cons

- Presence of repeated sequences in both homology arms might render PCR amplification or LR-screen difficult

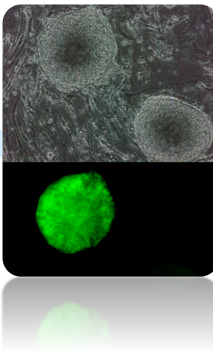
3 HOMOLOGOUS RECOMBINATION - VECTOR CONSTRUCTION



Created with SnapGene®

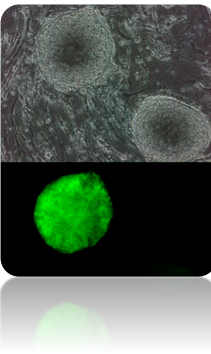


4 ES cell electroporation & Screening of recombinant clones



- Electroporation and screening process
- Long range PCR screening – strategy
- Long-Range 5' PCR screening – results
- Recombinant ES validation by Long Range PCR
- Recombinant ES clones validation by Southern Blot – internal probe
- Recombinant ES clones validation by Southern Blot – External probe
- Aneuploidy screening in ES recombinant clones

■ Electroporation and screening process



The whole process of ES cells validation is described in Erbs *et al.* *.

The targeting vector was co-electroporated with a CRISPR vector (expressing the WT Cas9 and a guide RNA –atagcaccatgaagtac- in the proprietary BALB/cN BA3 ES cell line.

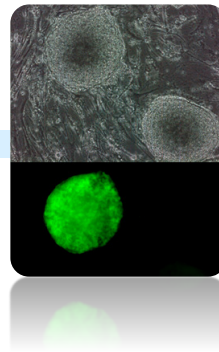
Transfected ES clones were submitted to neomycin selection (G418) and 93 resistant ES clones were isolated. The clones were then submitted to the screening process allowing secured identification of those harbouring the expected recombination events at both ends of targeting vector.

Screening process steps:

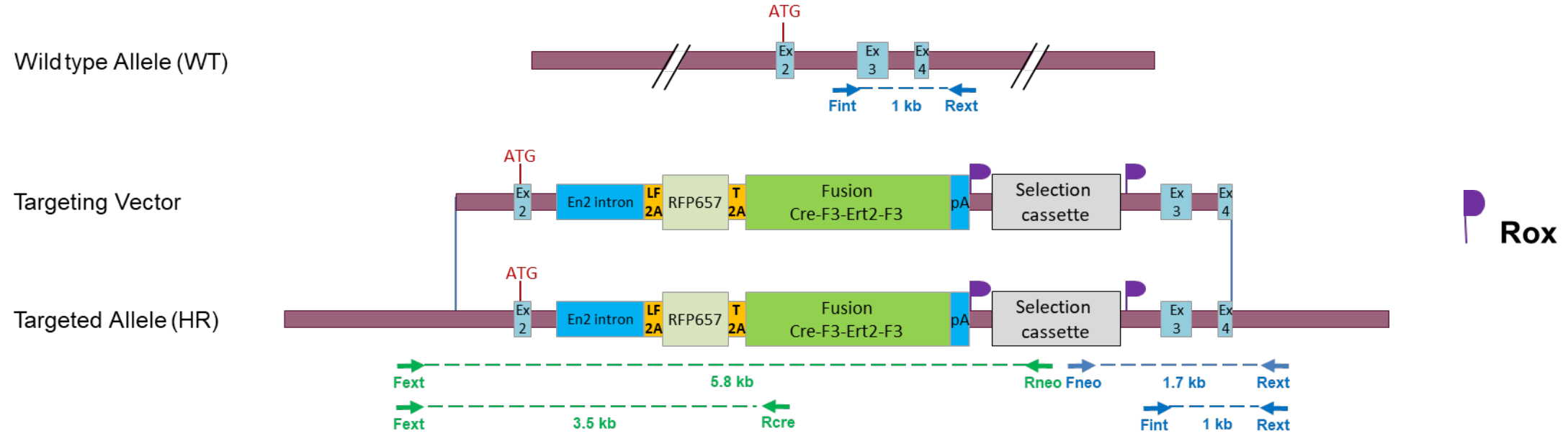
1. Identification of candidate recombinant clones by initial 3' Long-Range PCR
2. Six of 3' PCR positive clones are confirmed for 5' recombination event by Long-Range PCR
3. Positive clones in step2 are further validated by Southern blot analysis using internal and external probes
4. The karyotype of at least 2 validated clones is verified using ddPCR aneuploidy screening and Giemsa staining

*Erbs V, Lorentz R, Eisenman B, Schaeffer L, Luppi L, Lindner L, Hérault Y, Pavlovic G, Wattenhofer-Donzé M, Birling MC. Increased On-Target Rate and Risk of Concatemerization after CRISPR-Enhanced Targeting in ES Cells. *Genes (Basel)*. 2023 Feb 3;14(2):401. doi: [10.3390/genes14020401](https://doi.org/10.3390/genes14020401).

Long range PCR screening – strategy

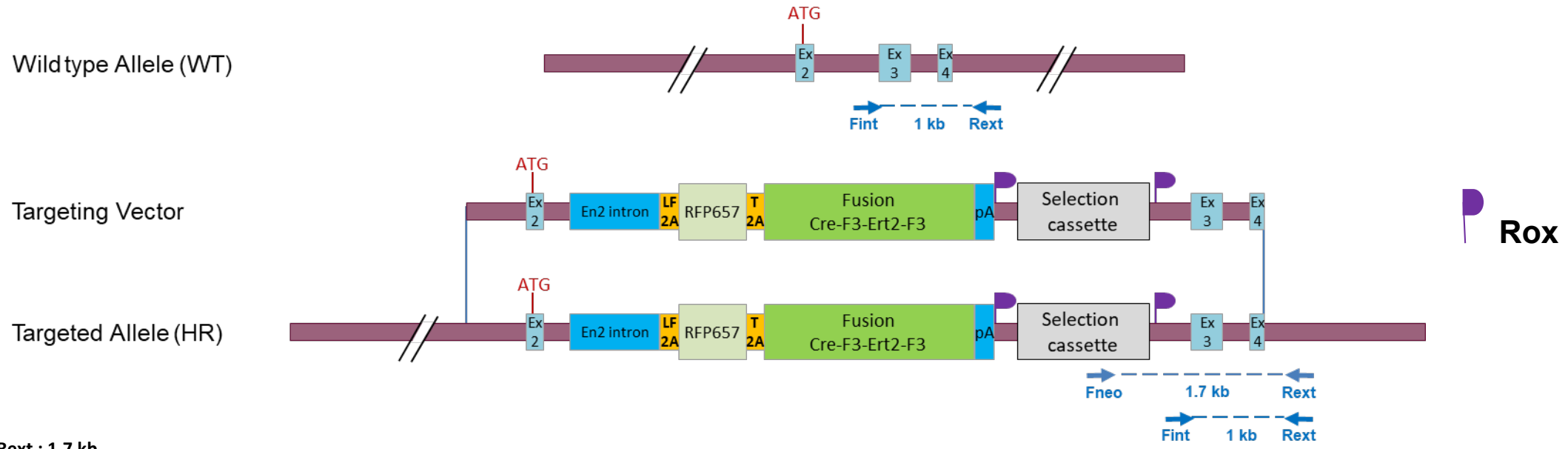
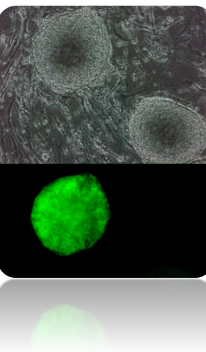


Schematic 5' and 3' PCR screening strategy

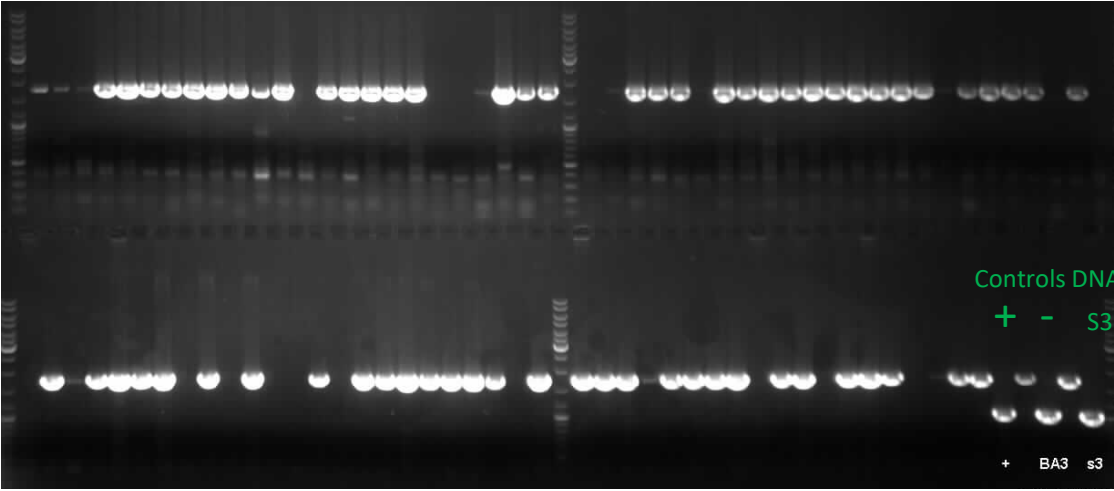


PCR	Primer Name	Primer sequences	PCR product size
5' PCR	Fext	GCAGGTGTCATATGCTGCTTTGAGG	5.8 kb
	Rneo	AGGGGCTCGCGCCAGCCGAAGTGT	
5' PCR	Fext	GCAGGTGTCATATGCTGCTTTGAGG	3.5 kb
	Rcre	CTCTACACCTGCGGTGCTAACCAGC	
3' PCR	Fint	GCTGGTAGTTGCTGGGTGCAGAACA	1 kb
	Rext	GGTAGATACTGGATGGGATCACAAG	
3' PCR	Fneo	GCGGCCGGAGAACCTGCGTGCAATC	1.7 kb
	Rext	GGTAGATACTGGATGGGATCACAAG	

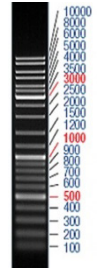
Long-Range 3' PCR screening – results



Pcr Fneo – Rext : 1.7 kb



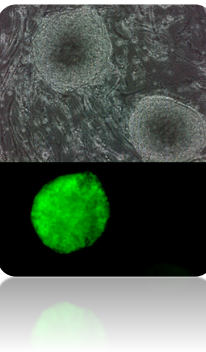
Pcr Fint – Rext : 1 kb



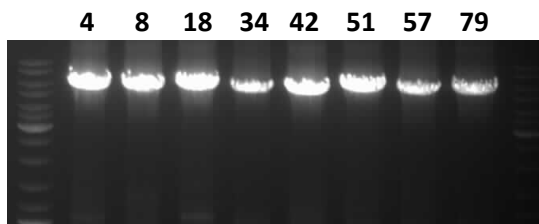
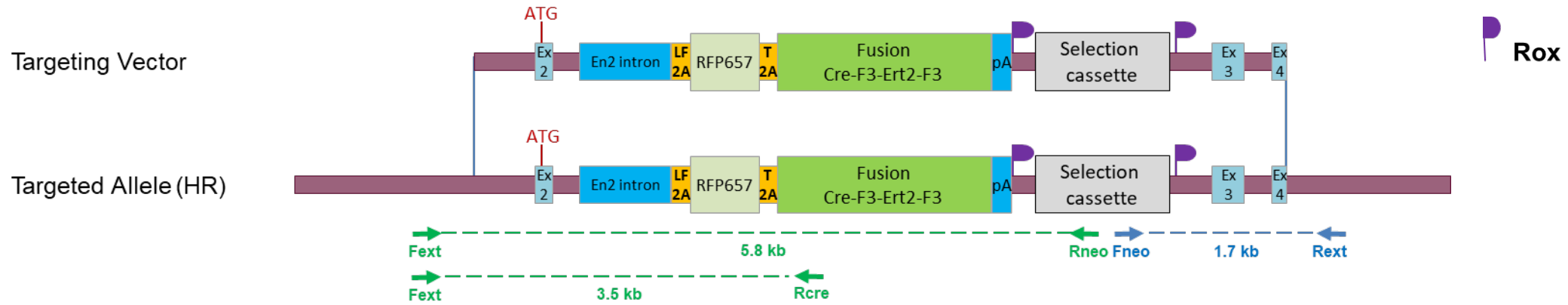
Ladder pattern

Six candidate clones out of the 74 positive clones were selected for 5' Long-Range PCR and Southern blot validation.

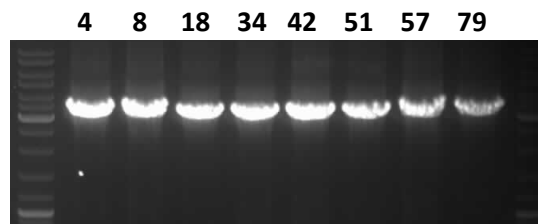
Recombinant ES validation by Long Range PCR



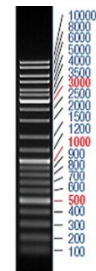
Confirmation and Validation of candidate recombinant ES clones by 5' and 3' PCRs



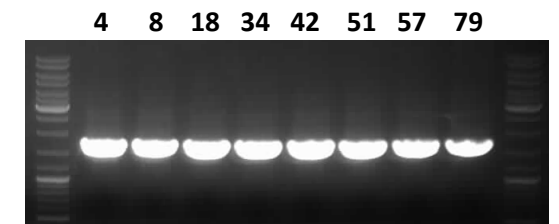
Pcr Fext – Rneo :5.8 kb (+DMSO)



Pcr Fext – Rcre :3.5 kb (+DMSO)



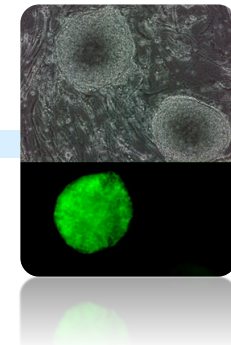
Ladder pattern



Pcr Fneo – Rext : 1.7 kb

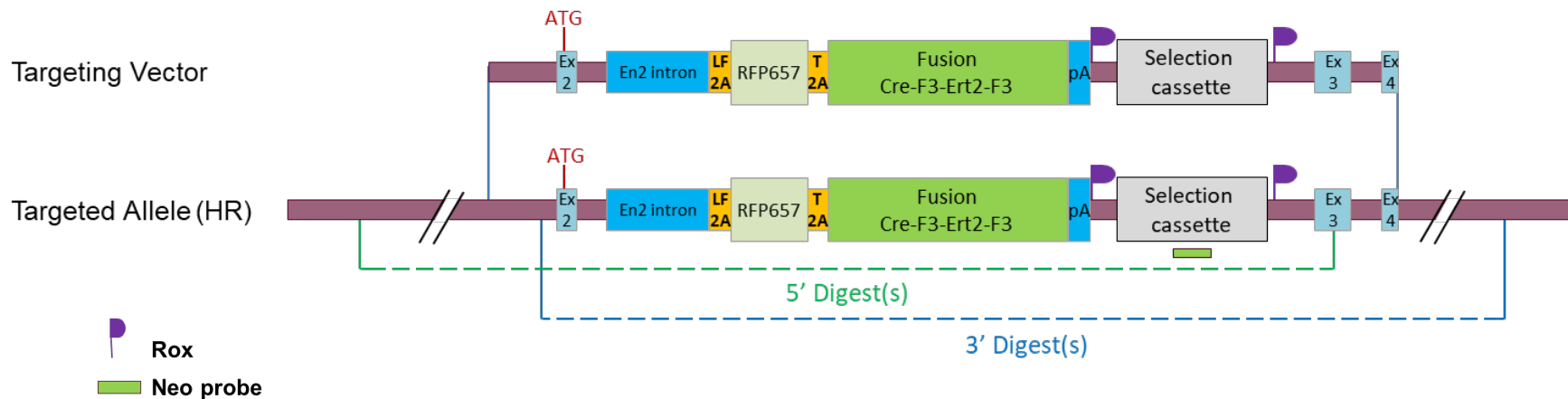
Six candidate clones identified by 3' PCR screening were further analysed by 5' PCR screening. Six clones (clones #4, #8, #18, #34, #42, #51 and #79) were confirmed.

Recombinant ES clones validation by Southern Blot – Internal probe



Schematic Southern Blot validation strategy

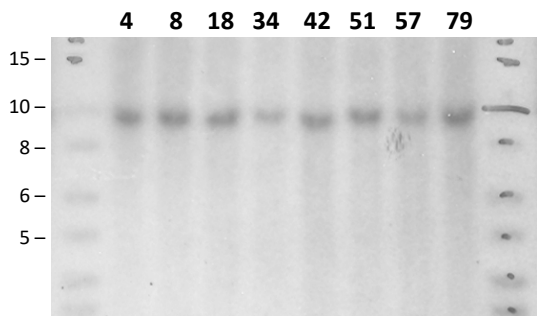
Digests on the scheme illustrate the position of the chosen restriction sites relative to the probe. They don't show the exact position of the restriction sites.



Neo probe sequence

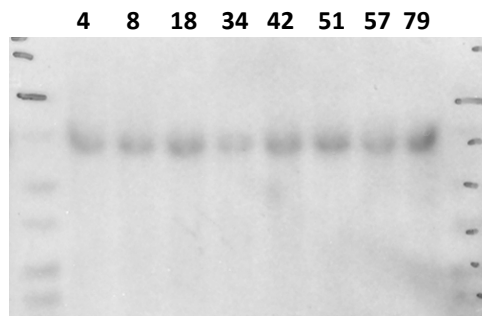
```
AGACTTACAGCGGATCCCCTCAGAAGAAGCTCGTCAAGAAGGCGATAGAA
GGCGATGCGCTGCGAATCGGGAGCGGCGATACCGTAAAGCACGAGGAAG
CGGTCAGCCCATTTCGCCCAAGCTCTTCAGCAATATCACGGGTAGCCA
ACGCTATGTCCTGATAGCGGTCCGCCACACCCAGCCGCCACAGTCGAT
GAATCCAGAAAAGCGGCCATTTTCCACCATGATATTCGGCAAGCAGGCA
TCGCCATGGGTCACGACGAGATCCTCGCCGTGGGCGATGCGCGCCTTGA
GCCTGGCGAACAGTTCGGCTGGCGCGAGCCCTGATGCTCTTCGTCCAG
ATCATCCTGATCGACAAGACCGGCTTCCATCCGAGTACGTGCTCGCTCG
ATGCGATGTTTCGCTTGGTGGTGAATGGGCAGGTAGCCGGATCAAGCG
TATGCAGCCGCCGATTGCATCAGCCATGATGGATACTTTCTCGGCAGG
AGCAAGGTGAGATGACAGGAGATCCTGCCCGGCACCTTCGCCCAATAGC
AGCCAGTCCCTTCCCGCTTCAGTGACAACGTGAGCACAGCTGCGCAAG
GAACGCCCGTCTGGCCAGCCACGATAGCCGCGCTGCCTCGTCTGCAG
```

Southern blot - Neo 5'



SpeI

Southern blot - Neo 3'



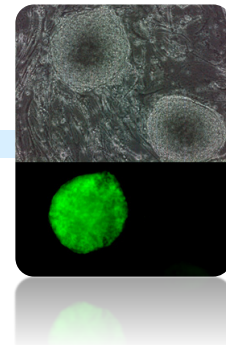
BglII

Digestions used to validate the 5' and 3' insertion

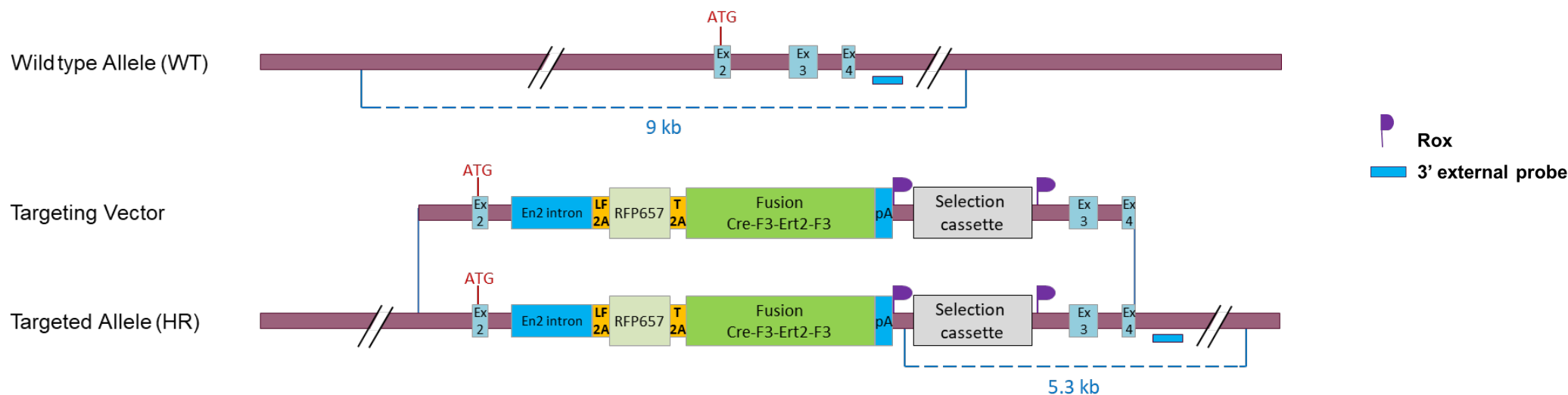
Probe		Genomic DNA digest	Targeted Allele (kb)
Neo	5' digest	SpeI	10.5
	3' digest	BglII	8.6

All 8 clones look correct (only one band of the expected size observed with the 2 restriction digests)

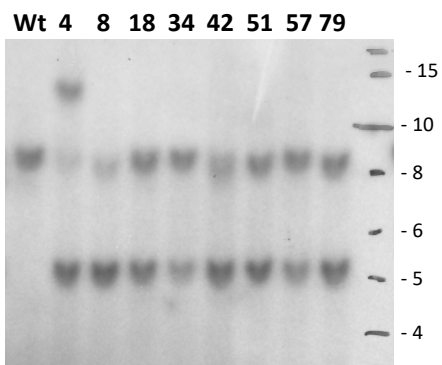
Recombinant ES clones validation by Southern Blot – External probe



Schematic Southern Blot validation strategy



Southern blot – 3' probe



BspHI 9 / 5.3

3' PROBE SEQUENCE

```
CCATCCAGTATCTACCTTGCTGCAGCTTTGTCCGTATTTCCC
CTTTACTATCCCCTAAAGAGCTGGGGACAGAAGGCCACCTAG
GCAAGACATCGAGATATCACAAAGGGTTTAGAATCAGGTTTG
AGTCCAAGTTCTCCAGTTTTTCTATCTTTGTGGCCTGGGG
ACTGTGTCCTTACCTGTAGGTTGCACATGGTTTTCCCTTGCCC
CATTACTGGTTTGTGATATAAGGATGATCATGTCAGAGAATG
GGTTGGTGATGACCACTGTGGACCATTAGTTCATCTGCCAG
TATCCATCATGGTCTGTGGTCAAGGATGTCAGGAGGAGTTA
ACAGGGACAACGCTGGCTCTGCCTCTTCCCTTCCCCCATGT
ATGCTTCTGCTTCTACCTTCTCTAGGGCCGATGCAAACGCT
TTCGATGGGTTGATGGAAGTTCTTGG
```

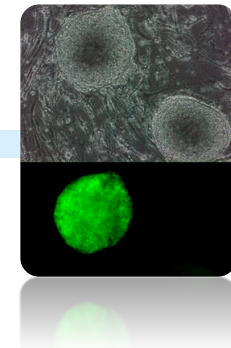
Digestions used to validate the 5' and 3' insertion

Probe	Name	Genomic DNA digest	WT allele (kb)	Targeted Allele (kb)
3' external probe	3' first digest	BspHI	9	5.3

=> All clones show the correct size for the targeted allele.

=> The other allele can be edited (no more WT allele but indels, clearly observed on clone #4)

■ Aneuploidy screening in ES recombinant clones

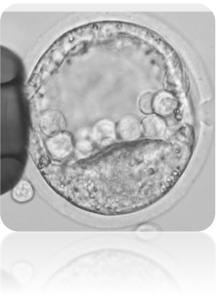


Selected recombinant ES cells clones were karyotyped by ddPCR as described in Codner *et al.*¹ and by Giemsa metaphase staining. Results of aneuploidy analysis are presented in the table below.

Clone ID	ddPCR	Giemsa
#18	Not done	Pass
#51	Not done	Pass

¹ Codner, G.F., Lindner, L., Calder, A., Wattenhofer-Donzé, M., Radage, A., Mertz, A., Eisenmann, B., Mianné, J., Evans, E.P., Beechey, C.V., Fray, M.D., Birling, M.-C., Hérault, Y., Pavlovic, G., Teboul, L
Aneuploidy screening of embryonic stem cell clones by metaphase karyotyping and droplet digital polymerase chain reaction.
BMC Cell Biology 2016 doi:10.1186/s12860-016-0108-6

5 MICROINJECTION & BREEDING



- Microinjection
- Breeding to F1 generation

■ Microinjection



- The ES cells used in the injection experiment were originally derived from a BALB/cN mouse strain (which have white coat colour). These cells were injected into blastocysts derived from an C57BL/6 strain, which have a black coat colour. The resulting offspring are thus chimeras of two different cell types (ES cell-derived cells and host blastocyst-derived cells) and the degree of chimerism was monitored by the percentage of light and dark patches on these animals.
- Recipient blastocysts were isolated from mated C57BL/6 females (Health status SPF Specific Pathogens Free).
- Recombinant ES clones #51 validated in previous project phase was injected into blastocysts to generate chimeric males. The results are presented in the table below.

Clone ID	Number of chimeric males identified according to chimerism rate (Number of chimeric males bred to F1 generation)			
	5 - 40%	45% - 55%	60-100%	Total
#51	1	0	1	2

■ Breeding to F1 generation



- Two highly chimeric males generated in the previous phase by blastocyst injection of the ES clones were mated with wild-type BALB/cN females (health status SPF – Specific Pathogen Free) to investigate whether the recombined ES cells have contributed to the germ layer.
- Germ line transmission was obtained the 24/09/2018
- Allele nomenclature (following MGI guidelines) : **Prg2^{tm2lcs}**



REPORT REDACTION & VALIDATION

Protocol finalized on 2023/05/2023

Prepared by Romain LORENTZ, IE

Verified by Marie-Christine BIRLING, PhD

CONTACT US

By email at mutagenesis@igbmc.fr

By phone at +33 (0)3 88 65 56 57

www.phenomin.fr