

EMMA ID: 04087

Gene: *Nfya*

Common name: *EUCJ0004_F10*

Allele: *Nfya*^{Gt(EUCJ0004f10)Hmgu}

Genotyping Information

Genotyping by end-point PCR based on gel is composed of a genespecific short range PCR using primers on wild type allele and a mutant allele-specific short range PCR. The combined results show the genotype of the mice. For example: mutant positive, wild type positive = Heterozygous.

PCR primer pairs and expected size bands

Assay	Forward Primer	Reverse Primer	Expected Size Band (bp)
Wildtype	E4F10-F	E4F10-R	520
Mutant	E4F10-F	Sanger5'seq	450

Primer sequences

Primer Name	Sequence 5' --> 3'
E4F10-F	GAGCATTCTGGGCATTTTCAGCTT
E4F10-R	CTGTATTTTACCGCTTCCACTAGC
Sanger5'seq	GTCCTCCGATTGACTGAGTCGC

PCR setup (Qiagen, Hot Start Plus)

Component	Volume (µl) 1x	Final conc.
DNA (~ 50-100 ng)	2	
Q-Solution (5x)	2,5	0,5
PCR-Buffer (10x)	2,5	1
DNTP mix (10 mM)	0,5	0,2
MgCl ₂ (25 mM)	1,5	1,5
Primer 1 (10 pmol/µl)	1	0,4
Primer 2 (10 pmol/µl)	1	0,4
Taq Polymerase (5 U/µl)	0,3	0,06
H ₂ O*	13,7	
Final volume	25	

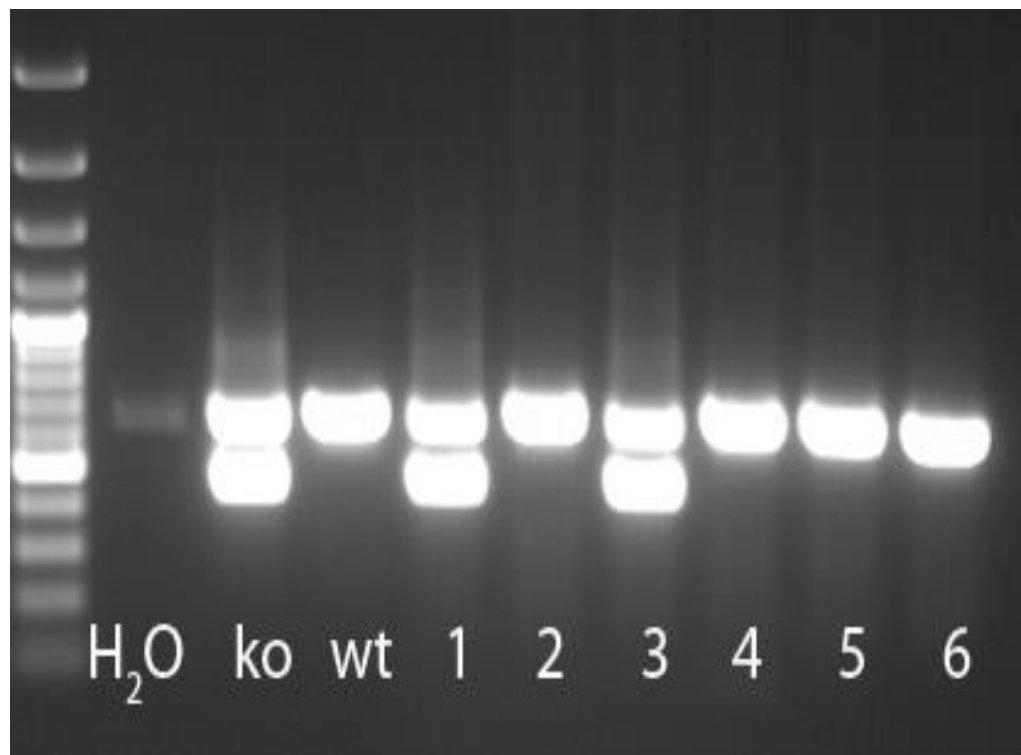
* The amount of H₂O is adjusted with the number of primer.

Amplification conditions

PCR Settings	Temperature (°C)	Time	# of cycles
1 Denaturation (Melting)	95°C	5 min	1
2 Amplification (Melting, Annealing, Polym.)	94°C	30 sec	39
	65°C	45 sec	
	72°C	45 sec	
3 Polymerisation	72°C	10 min	1
4 Cooling	12°C	hold	1

These PCR conditions have been optimized for our methods and preparation kits. Adaptions may be required.

Gel Image



Separated by gel electrophoresis on a 2% agarose gel.

ConstructQC vector_qc1 esmp_ha
 HTGTDB eucomm_vector esmp_ha

\$Id: header.tt 6045 2011-09-23 14:43:26Z rm7 \$

dp10 htgt1.sanger.ac.uk
 Authenticated username dp10@sanger.ac.uk
 Roles read, edit, eucomm, eucomm_edit

[Home](#) > [Report](#) > Gene Report

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Gene Report

Nfya [MGI:97316](#) [ENSMUSG00000023994](#) [OTTMUSG00000035112](#)

Program: KOMP Pipeline Progress: ES Cells - Targeting Confirmed Publicly Reported: Public

[Duplicate Project](#)

Pipeline Pre-pipeline Stage: Designs Vectors ES Cells Mice

[Reset gene to redesign](#)

[ES Cells - Targeting Confirmed](#)

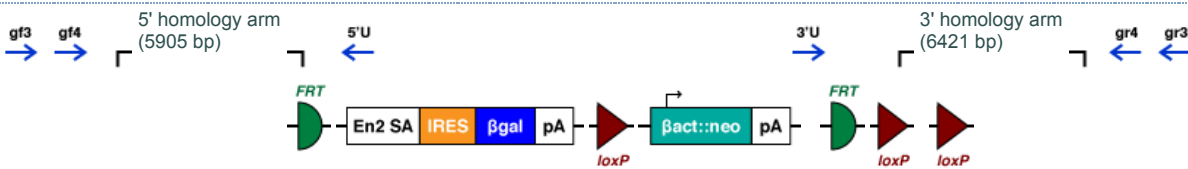
Recovery

Design Tools

ES Cell Clones

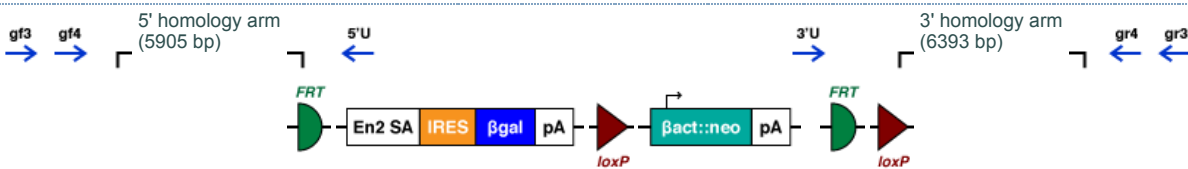
Order	Gene Targeting Vector	# Colonies Screened	# Knockout First Clones	# Targeted Non-conditional Clones	Ship Date
1	Nfya PRPGS00153_B_C04	33	6 GenBank File	13 GenBank File	29-JUN-09 (hzm) 29-JUN-09 (csd)

ES Cell Clones With Conditional Potential



[Promotor-Driven Cassette \(L1L2_Bact_P\)](#)

ES Cell Clones Without Conditional Potential



[Promotor-Driven Cassette \(L1L2_Bact_P\)](#)

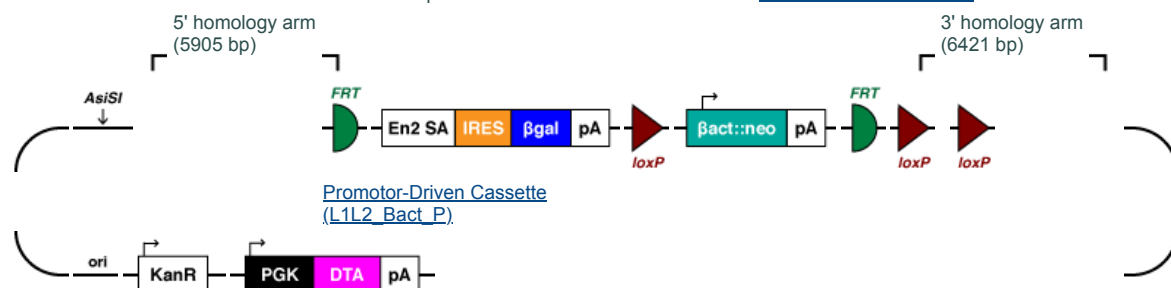
Note: Mutations of type "Targeted Non-conditional" are correctly targeted clones that have lost the 3' loxP site. These mutations cannot be converted into conditional alleles.

LR PCR Genotyping Primers

- 5' Gene Specific (GF3)** GTGTGTGTATGTATAGTAGGTATCTTAAG
- 5' Gene Specific (GF4)** CTGTGGAAGGACTTAGACCTAGTCAGAC
- 5' Universal (LAR3)** CACAACGGTCTTCTGTTAGTCC
- 3' Universal (RAF5)** CACACCTCCCCCTGAACCTGAAAC
- 3' Universal (PNF)** ATCCGGGGGTACCGCGTCGAG
- 3' Universal (R2R)** TCTATAGTCGCAGTAGGCGG
- 3' Gene Specific (GR3)** GCTGAAGCGCTACTGTTAAGTCGGAG
- 3' Gene Specific (GR4)** GCTGGCTGAAGCGCTACTGTTAAGTCGGAG


Targeting Vector

Order	Gene	Design ID	Vector Type	After Flp/Cre	Targeting Vector	Vector Strain	Floxed Exon	Cassette	Backbone	Genbank File
1	Nfya	169999 (153_C04)	Knockout First	Domain Disruption	PRPGS00153_B_C04	C57Bl/6J	ENSMUSE00000137024-	L1L2_Bact_P	L3L4_pZero_DTA_kan	view



[DTA Containing Plasmid Backbone](#)
(L3L4_pZero_DTA_kan)

Intermediate Vector

Order	Gene	Design ID	Vector Type	Intermediate Vector	Vector Strain	Floxed Exon	Genbank File
	Nfya	169999	(153_C04) Knockout First	PCS00153_A_C04	C57Bl/6J	ENSMUSE00000137024-ENSMUSE00000137021	view

Toolbox

[Short Range Primer Tool](#)

Useful Information

- **Links:**
- - [KOMP project](#)
 - [KOMP repository](#)
 - [WTSI Mouse Resources Portal](#)