

**EMMA ID:** 04608

**Gene:** *Pknox*

**Common name:** *EUCJ0079\_D10*

**Allele:** *Pknox1*<sup>Gt(EUCJ0079d10)Hmgu</sup>

## 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	EUCJ79_D10-F-5'arm	EUCJ79_D10-R-3'arm	427
Mutant	EUCJ79_D10-F-5'arm	Sanger5'	338

### Primer sequences

Primer Name	Sequence 5' --> 3'
EUCJ79_D10-F-5'arm	ACATAGTCACTCCTAGACTGC
EUCJ79_D10-R-3'arm	CTGTCTCCAAGAGTACACACC
Sanger5'	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 <sub>2</sub> (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 <sub>2</sub> O*	13,7	
Final volume	25	

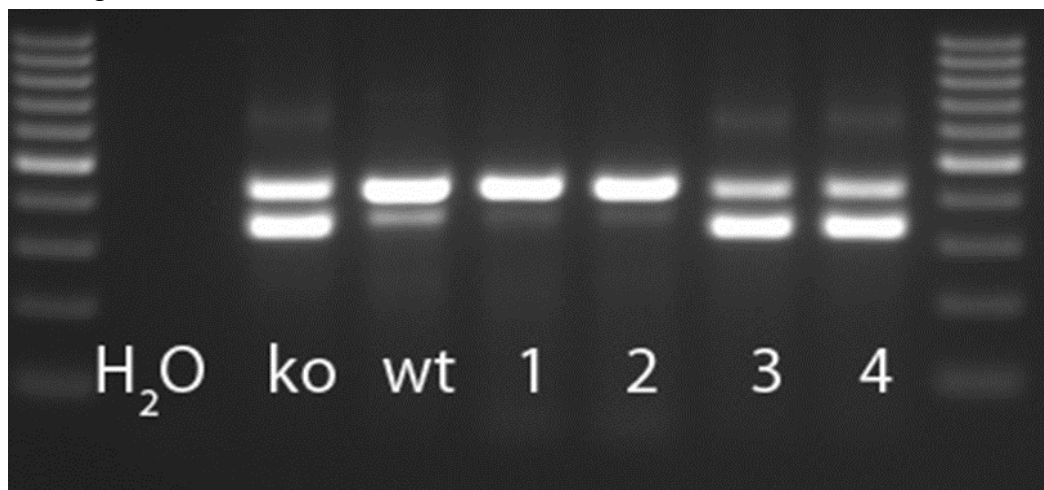
\* The amount of H<sub>2</sub>O is adjusted with the number of primer.

### 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
	55°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

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## Gene Report

**Pknox1** [MGI:1201409](#) [ENSMUSG0000006705](#) [OTTMUSG00000040106](#)

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

[Duplicate Project](#)

Pipeline Stage: Pre-pipeline Designs Vectors ES Cells Mice

[Reset gene to redesign](#)

[ES Cells - Targeting Confirmed](#)

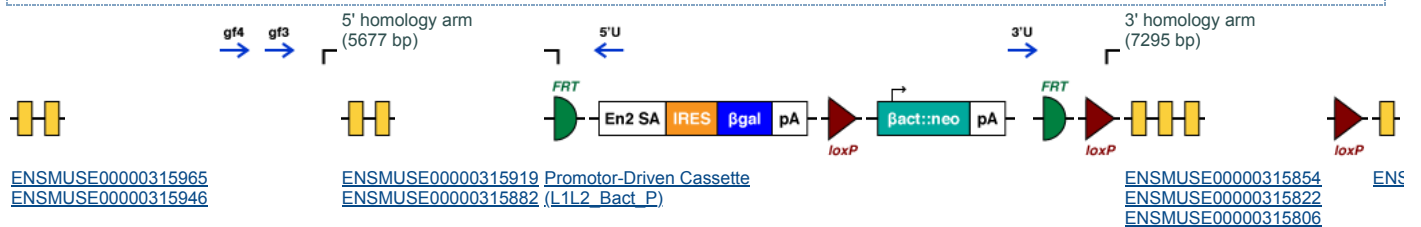
### Recovery

### Design Tools

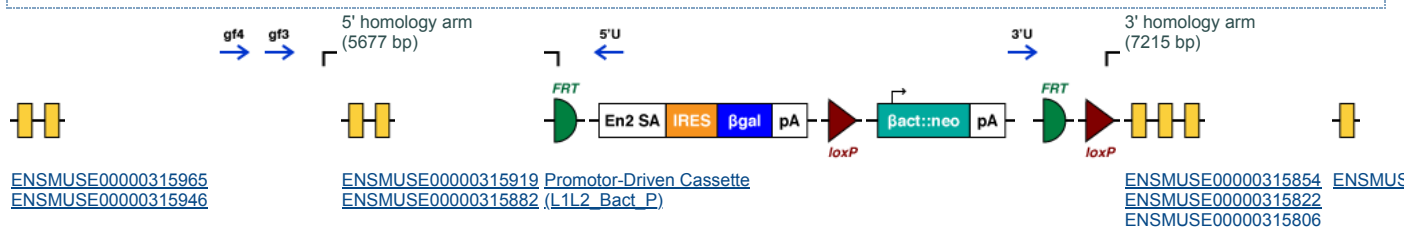
### ES Cell Clones

Order	Gene	Targeting Vector	# Colonies Screened	# Knockout First Clones	# Targeted Non-conditional Clones
1	Pknox1	PRPGS00139_C_B12	33	13 GenBank File	10 GenBank File

#### ES Cell Clones With Conditional Potential



#### ES Cell Clones Without Conditional Potential



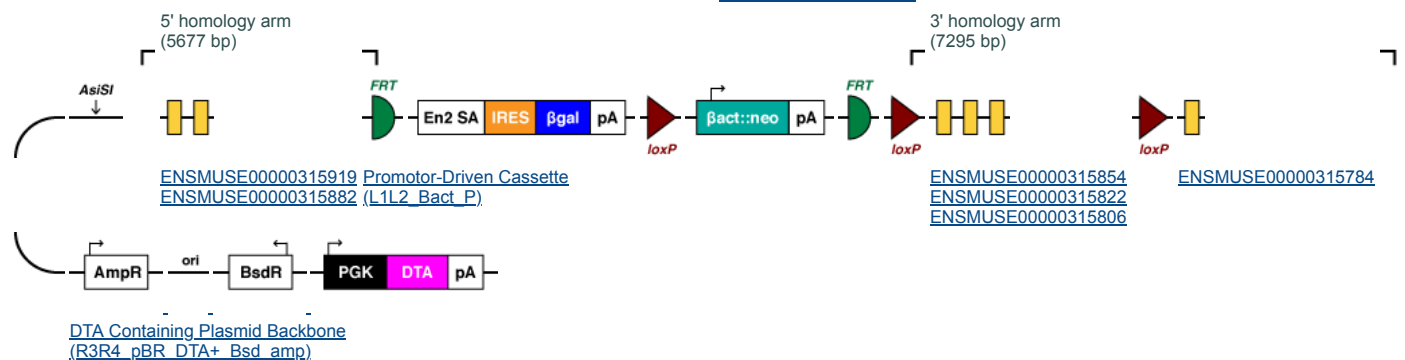
Note: Mutations of type "Targeted Non-conditional" are correctly targeted clones that have lost the 3' loxP site. These mutations cannot be converted into conditional alleles.

### LR PCR Genotyping Primers

5' Gene Specific (GF3) GCACGACGTCTGCCAGCTTCGATGTGGAC  
 5' Gene Specific (GF4) GCACTGTGACACTGTGAGCACAGGTATG  
 5' Universal (LAR3) CACAACGGGTTCTTCTGTTAGTCC  
 3' Universal (RAF5) CACACCTCCCCCTGAACCTGAAAC  
 3' Universal (PNF) ATCCGGGGGTACCGCGTCGAG  
 3' Universal (R2R) TCTATAGTCGAGTAGGCGG  
 3' Gene Specific (GR3) GTGTGGTGATGGTCACAACGGCACCTGCAG  
 3' Gene Specific (GR4) GTGATGGTCACAACGGCACCTGCAGCAGAG

### Targeting Vector

Order	Gene	Design ID	Vector Type	After Flip/Cre	Targeting Vector	Vector Strain	Floxed Exon	Cassette	Backbone	Genbank File
1	Pknox1	210458 (139_B12)	Knockout	Frameshift	PRPGS00139_C_B12	C57Bl/6J	ENSMUSE00000137435- ENSMUSE00000137433	L1L2_Bact_P	R3R4_pBR_DTA+ Bsd_amp	<a href="#">view</a>



## Intermediate Vector

Order Gene Design ID Vector Type Intermediate Vector Vector Strain Floxed Exon Genbank File  
 [Pknex1](#) [210458](#) (139\_B12) Knockout First [PCS00139\\_B\\_B12](#) C57Bl/6J [ENSMUSE00000137435-ENSMUSE00000137433](#) [view](#)

## Toolbox

[Short Range Primer Tool](#)

## Useful Information

- Links:

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