

## NIH-0739 Genotyping Strategies

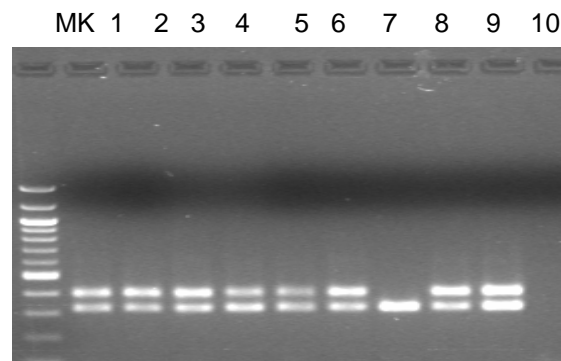
Reaction Components	Vol (ul)
10X Sigma Buffer	5
25mM MgCl <sub>2</sub>	3.5
10mM dNTPs	2
Primer 20 uM	1.5
Primer 20 uM	1.5
5 U/ul Taq polymerase	0.5
Water	31
Total mix volume	45
Tail lysate (1:20 dilution)	5
Total reaction volume	50

Step	Temp	Time	Note
1	94C	15"	
2	65C	30"	Decrease 1C/cycle
3	72C	40"	Go to 1, 10 cycles
4	94C	15"	
5	55C	30"	
6	72C	40"	Go to 4, 30 cycles

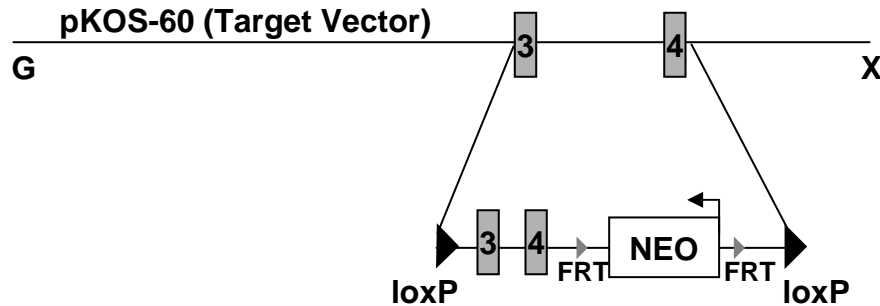
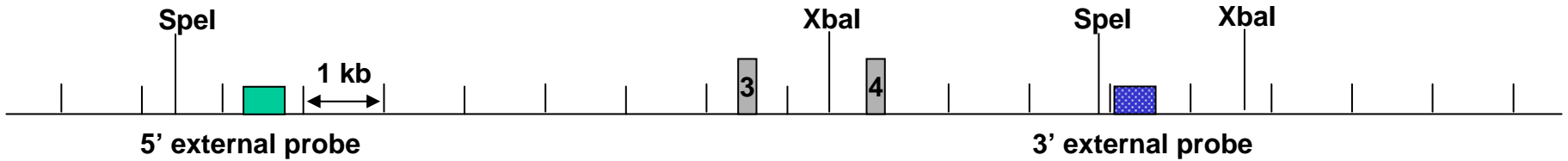
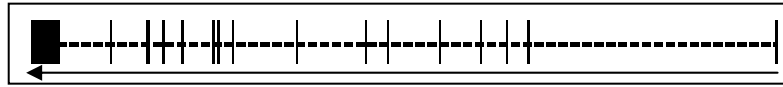
Primer Sequences (5' to 3'):	
Cre excision PCR: Primer 0739-lox5 and Primer 0739-3'cre, 426 bp	
LoxP Mutant PCR: Primer 0739-lox5 and Primer 0739-lox3, mutant 415 bp, wt 325 bp	
Primer 0739-lox5	CCCAAGAGAAGCCACCTAGC
Primer 0739-lox3	AGCCCAGTCCAGGATCAAG
Primer 0739-3'cre	TCAAATTTTACTGTCTACCAGTCCC

Well	Sample	Genotype
1	5	het
2	6	het
3	10	het
4	16	het
5	18	het
6	37	het
7	22	wt
8	23	het
9	es DNA	het
10	water	no amp



### Mutant PCR



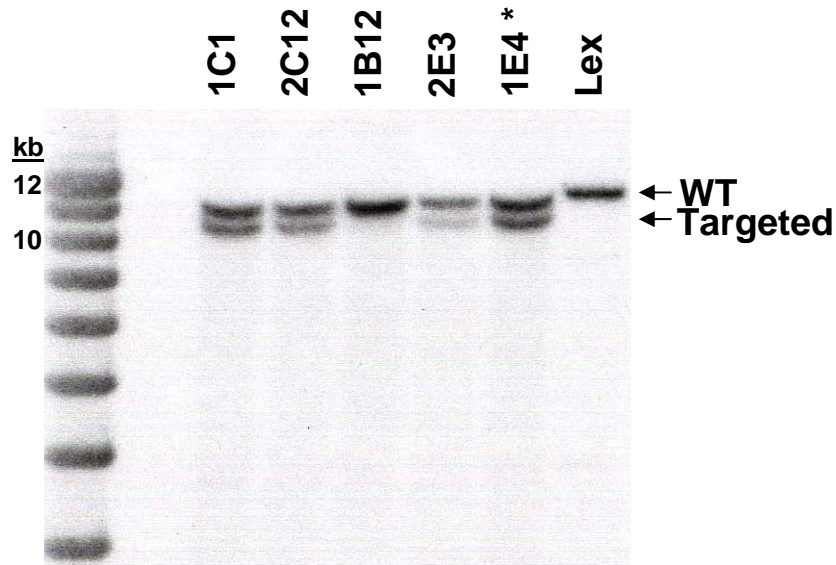
# Targeting Strategy



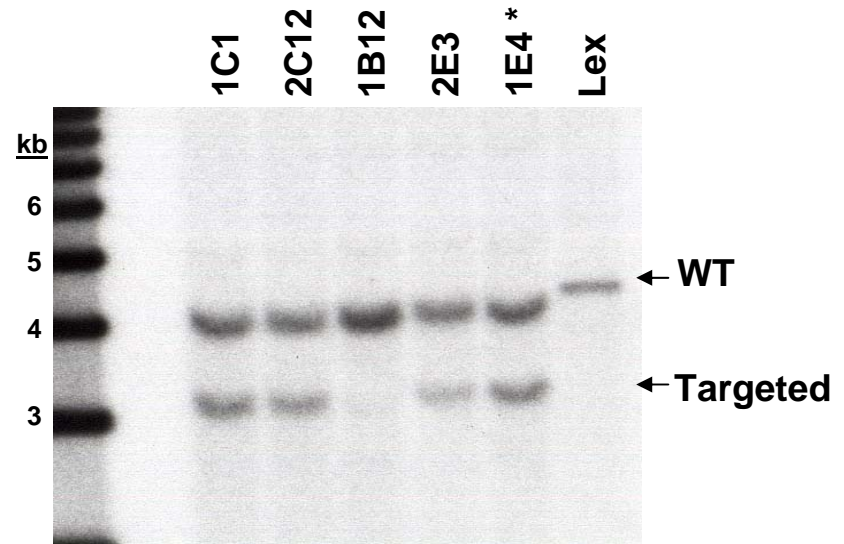
## Southern Strategies:

Probe	5' external 	3' external 
Enzyme	SpeI	XbaI
Wildtype	11.0 kb	4.4 kb
Targeted	10.0 kb	3.3 kb

# Southern Data



5' external probe  
SpeI digests  
Wildtype 11.0 kb  
Targeted 10.0 kb

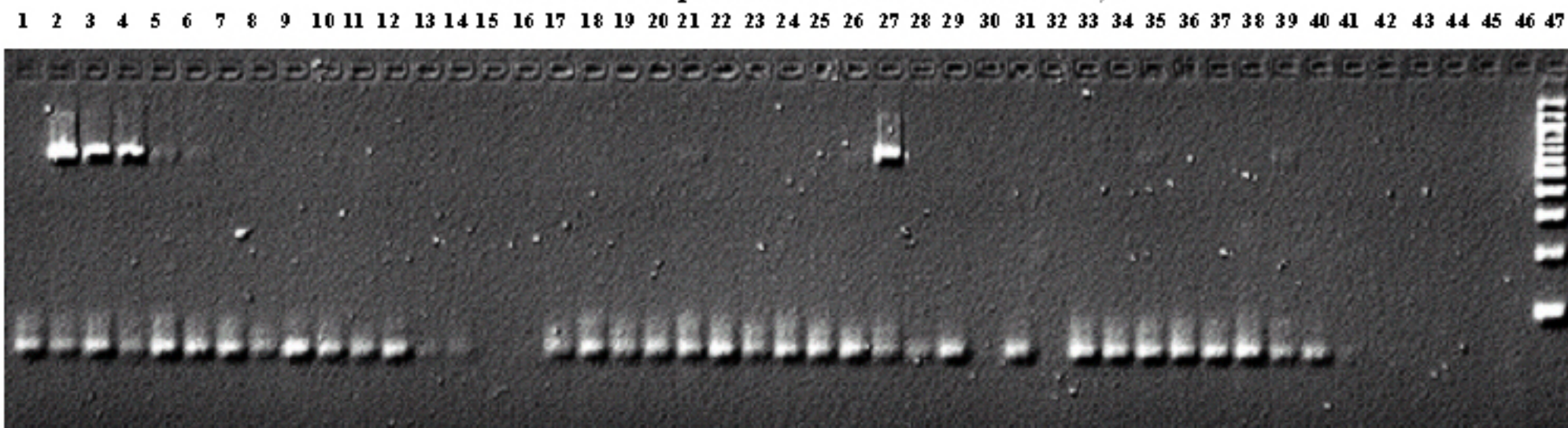


3' external probe  
XbaI digests  
Wildtype 4.4 kb  
Targeted 3.3 kb

\*Clone achieving germline transmission

# RT-PCR Analysis

## mouse random primed cDNA with Primers: 1,2



12/14/2002

**Note:** mouse ES cDNA used as positive (+) control. Expected band size denoted by arrow adjacent to 100bp ladder/marker.

### Mouse cDNA Tissues

1	Brain	19	Ovary	37	Mammary Gland
2	Thymus	20	Gall Bladder	38	Placenta
3	Spleen	21	Lymph node	39	9.5 day Embryo
4	Lung	22	Cerebellum	40	12.5 day Embryo
5	Kidney	23	Esophagus	41	Fetal Brain
6	Liver	24	Prostate	42	Fetal Lung
7	Testis	25	Tongue	43	Fetal Liver
8	Adrenal Gland	26	Thyroid	44	Fetal Kidney
9	Pancreas	27	Bone Marrow	45	(-) Control
10	Salivary Gland	28	Spinal Cord	46	(+) Control
11	Stomach	29	Trachea	47	100 bp ladder/marker
12	Small Intestine	30	Aorta		
13	Colon	31	Whole Blood		
14	Skeletal Muscle	32	Eye		
15	Heart	33	Pituitary gland		
16	Uterus	34	Skin		
17	Adipose	35	Nasal Epithelium		
18	Bladder	36	Whole Bone (femur)		



**Lexicon Genetics Incorporated  
Molecular Genetics Project Materials**

**Catalog Number:** NIH-0739 (LEXKO-331)

**Reference accession(s):** NM 010583

**Standard KO or Conditional:** conditional

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**Materials Submitted:** x Target Vector pKOS-60TVpLFneo  
KOS clone(s) not available

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**Southern Blot Genotyping Strategies:**

	<u>5' External</u>	<u>3' External</u>
Name of Probe:	73 + 74	67 + 68
Restriction Enzyme for Genomic Digest:	SpeI	XbaI
Predicted Wild-type Band (kb):	11.0	4.4
Predicted Mutant Band (kb):	10.0	3.3
Probe Size:	597 bp	262 bp

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**Primer sequences:**

**Southern probes**

0739-73 5' – CATTAGCTAACCTAGCCC  
0739-74 5' – CCTGGCTCGTTCTTCACATC  
0739-67 5' – TCCCAACCCAGAAGCCAG  
0739-68 5' – AGGCATGGGGTATGTGTGCT

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**Genomic Sequence Floxed:**

TGATGATAAAATACTGAGATGGACCTGACATGGGCTCCCCATGTCTGGGTACCCTGTTCATCGTAGACTCTTGTGTACTTA  
CAGGTCGTGCATGACAACTATCTCCTGTATGTGTTTGTCTCCAGACTGTGAGAGTCCGGCAGCGCTGGGTGCTGACCCTTA  
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**KOS clone sequence:** (note: pKOS-60 was used to generate the TV and that is the sequence included here)

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**Selection cassette sequence:** (note: linker sequences may vary and are not provided)

GGCCATAGCGGCCGGCCATAACTTCGTATAGCATA CATTATA CGAAGTTATGGCGCGGAGTCGACGATCAAGCTTTCG  
AAGATCTACGTGGCGGCCCTCGAGCTTTCGGAAGTTCCTATTTCGGAAGTTCCTATTCTCTAGAAAAGTATAGGAACTTCTC  
GAGATCCGATATCGAATTCCTCGCCCCAGCTGGTCTTTCCGCCTCAGAAGCCATAGAGCCCACCGCATCCCCAGCA  
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