

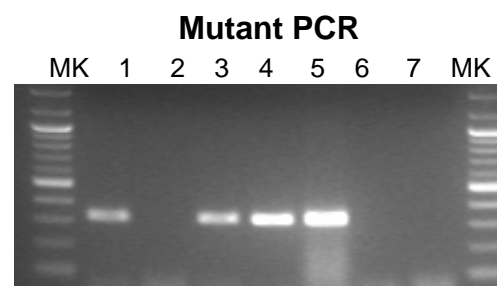
## NIH-0516 Genotyping Strategies

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

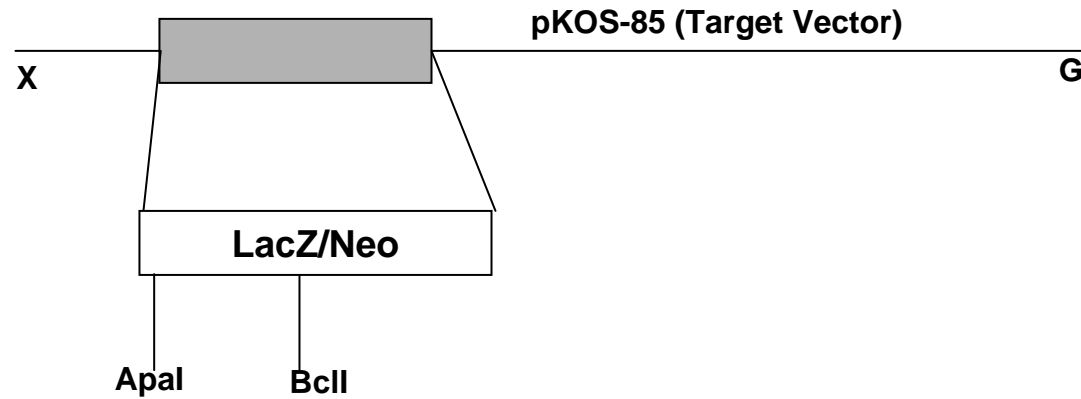
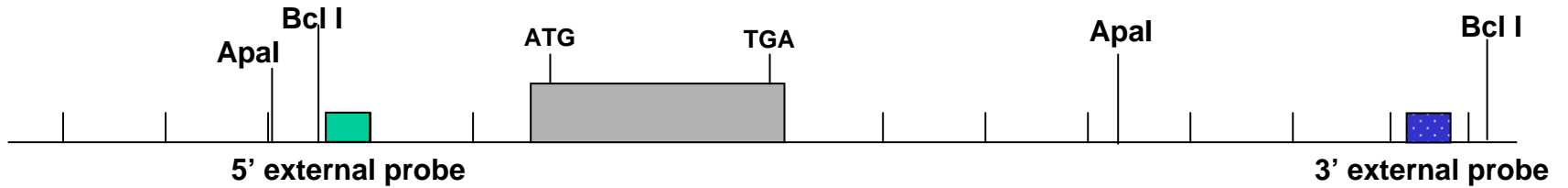
Step	Temp	Time	Note
1	94C	15"	
2	65C	30"	Decrease 1C/cycle
3	72C	40"	Go to 1, 10 cycles
4	94C	15"	
5	55C	30"	
6	72C	40"	Go to 4, 30 cycles

Primer Sequences (5' to 3')	
Mutant PCR: Primer neo3a and Primer 0516-13, 318 bp	
Recommended Wt PCR: Primer 0516-2 and Primer 0516-9, 870 bp	
Primer neo3a	GCAGCGCATCGCCTTCTATC
Primer 0516-13	CAGTCTAGGACTCCTATAATCC
Primer 0516-2	CTACAACCTGACCCGCATGC
Primer 0516-9	AGCCAGGAACCAGGTTAGAGTC



Well	Sample	Genotype
1	54	het
2	55	wt
3	56	het
4	57	het
5	ES DNA	het
6	wt lysate	wt
7	water	no amp



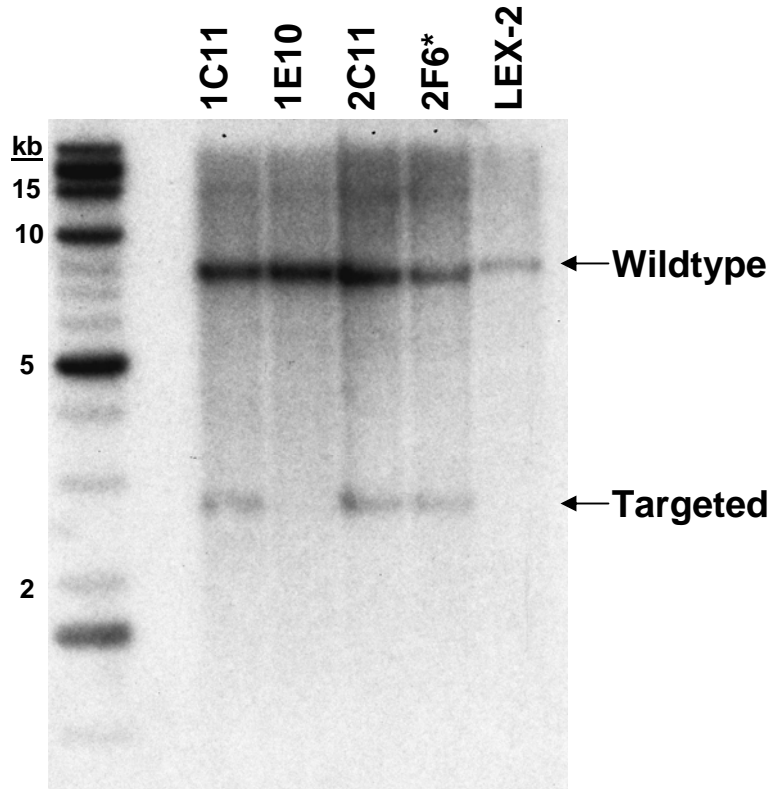
# Targeting Strategy



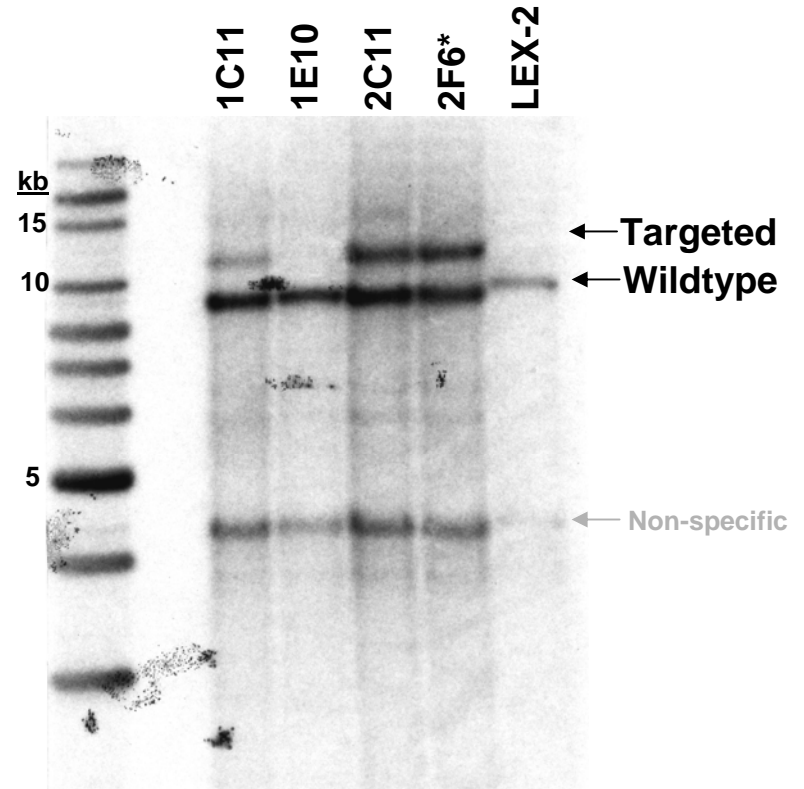
## Southern Strategies

Probe	5' external 	3' external 
Enzyme	Apal	Bcl I
Wildtype	8.2 kb	9.1 kb
Targeted	2.8 kb	11.1 kb

# Southern Data



5' external probe  
 Apal digests  
 Wildtype 8.2 kb  
 Targeted 2.8 kb

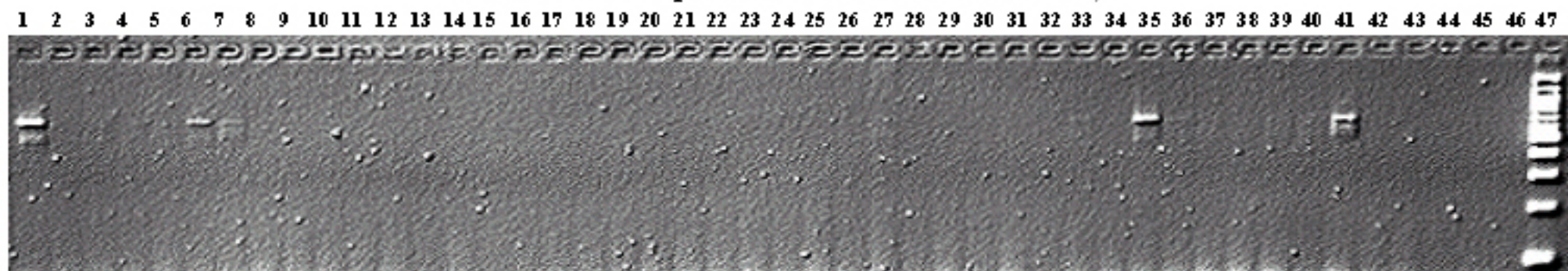


3' external probe  
 Bcl I digests  
 Wildtype 9.1 kb  
 Targeted 11.1 kb

\*Clone achieving germline transmission

## RT-PCR Analysis

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



12/11/2002

**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-0516 (LEXKO-222)

Reference accession(s): \_ AF033585, AK021164

Standard KO or Conditional: \_ Standard

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**Materials Submitted:**  Target Vector \_ pKOS-85TVneo  
 KOS clone(s) \_ pKOS-85

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**Southern Blot Genotyping Strategies:**

	<u>5' External</u>	<u>3' External</u>
Name of Probe:	20/21	18/19
Restriction Enzyme for Genomic Digest:	ApaI	BclI
Predicted Wild-type Band (kb):	8.2	9.1
Predicted Mutant Band (kb):	2.8	11.1
Probe Size:	194 bp	320 bp

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**Primer sequences:**

**Southern probes**

0516-20 5' – GTGCGCGCATGTAACAGGCAAC  
0516-21 5' – CCACCTCCCAAGTGCCTGAGC  
0516-18 5' – CGTGGGCTCAGAGCACAG  
0516-19 5' – GGGTAACTGGAAATGACGAGGC

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**Genomic Sequence Deleted:**

CCGCGACGTGGCGCGCAGATCCCGGGACGCGGGATGGAGCGGTGCGAGTGAGCCCGAGCGGCCGCGCCCGGGGGCGATGGCCGTGCCACCGCTGCTCC  
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**KOS clone sequence:** (note: pKOS-85 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)

**LacZ/Neo**

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