



Lexicon Genetics Incorporated – Genentech Project Materials

Genentech ID:	UNQ666	Date of Submission:	May 7, 2004
Lexicon Contract Name:	DNA101	Mutation Type:	<input checked="" type="checkbox"/> Standard Knock out
LexVision Name:	ENZ643N1		<input type="checkbox"/> Conditional
Reference accessions:	ENSMUST00000044148	Is this gene X-linked?	No

Required Materials: X pKOS clone DNA(s) __10_____
 X Target Vector DNA __pKOS.DNA101.10 TV_____
 X Targeted ES Cell DNA __2A1_____
 X Genomic Map

Southern Blot Analysis:
External/Internal Probe Strategies

	<u>5' External</u>	<u>3' External</u>
Name of Probe:	27/29	30/32
Restriction Enzyme for Genomic Digest:	EcoR I	Acc65 I
Predicted Wild-type Band (kb):	9	21
Predicted Mutant Band (kb):	7	15
Probe Size:	456	427

PCR Strategies:*For standard knockouts, give wildtype and mutant-specific strategies**For conditionals, give 5' loxP and cre-excision strategies*

Wild type-specific (absent in targeted allele)		Mutation-specific product (absent in wt)	
5' Primer Name:	DNA101-14	5' Primer Name:	neo3a
3' Primer Name:	DNA101-15	3' Primer Name:	DNA101-25
Predicted Wild-type Band (bp):	573	Predicted Wild-type Band (bp):	none
Predicted mutant band (bp)	none	Predicted mutant band (bp)	273

5' loxP strategy		Distinguish Cre-excised and wt	
5' Primer Name:		5' Primer Name:	
3' Primer Name:		3' Primer Name:	
Predicted Wild-type Band (bp):		Predicted Wild-type Band (bp):	
Predicted mutant band (bp)		Predicted mutant band (bp)	

Primer sequences:**Southern probes**

DNA101-27 5' – CACCCTGCGTACTCTATCCAAACC
DNA101-29 5' – ATAAGGCCTCAGAGAAGAAGTGTG
DNA101-30 5' – CTAACGCTTGCAAGTGGCTCTAA
DNA101-32 5' – CCTTGACTCTTCTGGGCTGGATA

PCR Genotyping

DNA101-14 5' - AGGCAGCTAGTCAACCAGGAAAAG
DNA101-15 5' – ATAACCACCGGCCATGAAGAGAT
DNA101-25 5' – CTCACATTCAAGACTCAAGTCT
neo3a 5' – GCAGCGCATCGCCTTCTATC

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