



Lexicon Genetics Incorporated – Genentech Project Materials

Genentech ID:	UNQ577	Date of Submission:	4/28/05
Lexicon Contract Name:	DNA221A	Mutation Type:	X Standard Knock out
LexVision Name:	MEM688N1		☐ Conditional
Reference accessions:	NM_026609	Is this gene X-linked?	no

Required Materials: X pKOS clone DNA(s) __pKOS-7,67_____
X Target Vector DNA __pKOS-67TV_____
X Targeted ES Cell DNA __2A12_____
X Genomic Map

Southern Blot Analysis: *External/Internal Probe Strategies*

	<u>5' External</u>	<u>3' Internal</u>
Name of Probe:	14/15	19/20
Restriction Enzyme for Genomic Digest:	BstX1	HpaI
Predicted Wild-type Band (kb):	11.3	10.9+n (~11)
Predicted Mutant Band (kb):	5.8	8.6+n (~8.7)
Probe Size (bp):	356	368

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:	5	5' Primer Name:	Neo3a
3' Primer Name:	6	3' Primer Name:	13
Predicted Wild-type Band (bp):	280	Predicted Wild-type Band (bp):	none
Predicted mutant band (bp)	none	Predicted mutant band (bp)	300

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**

14 5' – CTTCTGTTGGCCTTTAGATTG
 15 5' – TTATGACACTGGCAGCTAGGTG
 19 5' – CCAATTGCTATGCCATACATGC
 20 5' – TGAAGATGCCAATGACCGCTAC

PCR Genotyping

5 5' – CTGACTACGCTTCTCACGTGAG
 6 5' – GAATATGGCAAGCTCCTTAGAC
 13 5' – GACAGCAAGAACAGACGGAAG
 neo3a 5' – GCAGCGCATCGCCTTCTATC

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