Lexicon Genetics Incorporated – Genentech Project Materials

Genentech ID:	**UNQ1903**	**Date of Submission:**	7-13-04
Lexicon Contract Name:	DNA017	**Mutation Type:**	X Standard Knock out
LexVision Name:	SEC440N1		☐ Conditional
Reference accessions:	**BC011475**	**Is this gene X-linked?**	NO

Required Materials:
X pKOS clone DNA(s) __KOS 25_____
X Target Vector DNA __FTV25 NEO_____
X Targeted ES Cell DNA __1H7_____
X Genomic Map

Southern Blot Analysis:
External/Internal Probe Strategies

	5' Internal	**3' External**
Name of Probe:	**15/16**	**13/14**
Restriction Enzyme for Genomic Digest:	**EcoRI**	**BamHI**
Predicted Wild-type Band (kb):	**7.4**	**10.6**
Predicted Mutant Band (kb):	**5.9**	**7.4**
Probe Size:	**443 BP**	**313 BP**

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:	DNA017-5	5' Primer Name:	NEO 3A
3' Primer Name:	DNA017-6	3' Primer Name:	DNA017-11
Predicted Wild-type Band (bp):	473 BP	Predicted Wild-type Band (bp):	none
Predicted mutant band (bp)	none	Predicted mutant band (bp)	507 BP

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
DNA017-15 5' – CCAGTAGATCACGGGTGGTA
DNA017-16 5' – GCCTGGGATTTATTCTCTGACGA
DNA017-13 5' – CTGGCCAGGAAGGGGATAGAATA
DNA017-14 5' – CTGTGTCTTGTAGGGGGTTC

PCR Genotyping
DNA017-5 5' – CTCCCTGAGTGTCCGATAAGATG
DNA017-6 5' – CGCCATTTTTGTGACTCT
DNA017-11 5' –GGCCTTACCTGACGGACATCACC
Neo3a 5' - GCAGCGCATCGCCTTCTATC

Genomic Sequence Deleted:

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Genomic Locus: ***(the deleted sequence represents nt 8011-11199-in the sequence below)***

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Selection Cassette:

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Targeted Locus:

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