Lexicon Genetics Incorporated – Genentech Project Materials

Genentech ID:	UNQ1942	**Date of Submission:**	1/17/05
Lexicon Contract Name:	DNA382	**Mutation Type:**	X Standard Knock out
LexVision Name:	SEC565N1		☐ Conditional
Reference accessions:	XM_132070	**Is this gene X-linked?**	No

Required Materials:
X pKOS clone DNA(s) __pKOS 41_____
X Target Vector DNA __pTV 41Bgeo_____
X Targeted ES Cell DNA __1C6_____
X Genomic Map

Southern Blot Analysis:
External/Internal Probe Strategies

	5' External	**3' External**
Name of Probe:	**37 + 36**	**22 + 23**
Restriction Enzyme for Genomic Digest:	**BglII**	**BamHI**
Predicted Wild-type Band (kb):	**11.8 kb**	**14.9 kb**
Predicted Mutant Band (kb):	**16.1 kb**	**10.32 kb**
Probe Size:	**240 bp**	**244 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:	DNA382-13	5' Primer Name:	DNA382-12
3' Primer Name:	DNA382-11	3' Primer Name:	GT-IRES
Predicted Wild-type Band (bp):	202	Predicted Wild-type Band (bp):	none
Predicted mutant band (bp)	none	Predicted mutant band (bp)	376

Primer sequences:

Southern probes
DNA382-22 5' – **CTGAAAGGAGAGTCCGTTCC**
DNA382-23 5' – **GCTGGAGGATTGCAAGCTG**
DNA382-36 5' – **GGAGCTATGACTGTGTGGC**
DNA382-37 5' – **CCTTGACCTCTCTGTAGAAAC**

PCR Genotyping
DNA382-11 5' – **CAGCACACAGTAGTTCTGAG**
DNA382-12 5' – **GTGGGTGCCATAGTATCAAG**
DNA382-13 5' – **CTGGACATCCATGTAAGAGAG**
GT-IRES 5' – **CCCTAGGAATGCTCGTCAAGA**

Genomic Sequence Deleted:

GTGAGGTACTTGCCCCGGCTTTACCTGGACATCCATGTAAGAGAGCTTTGCCTGGTGGGTGGGTGGGGTAGGGTGGATGTGAACTGCCTCTGCTGCTGG
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Genomic Locus: (*The deleted sequence represents nt 10003 – 12093 in the sequence below. KOS41 used to generate the TV represents nt 4086 – 14606 in the sequence below.)*

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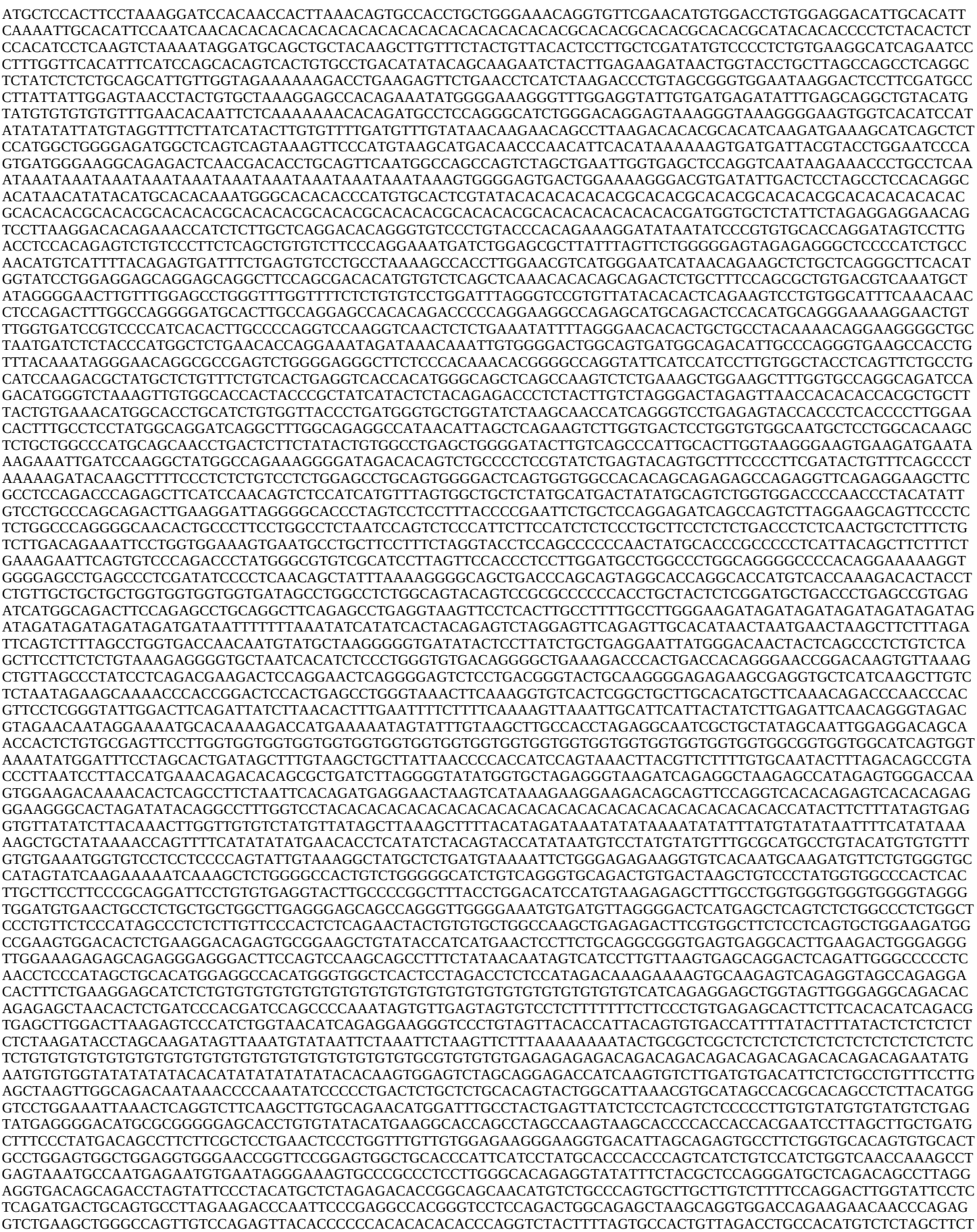

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Selection Cassette:

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Targeted Locus:

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