

Supporting Information

Nelson et al. 10.1073/pnas.1207169109

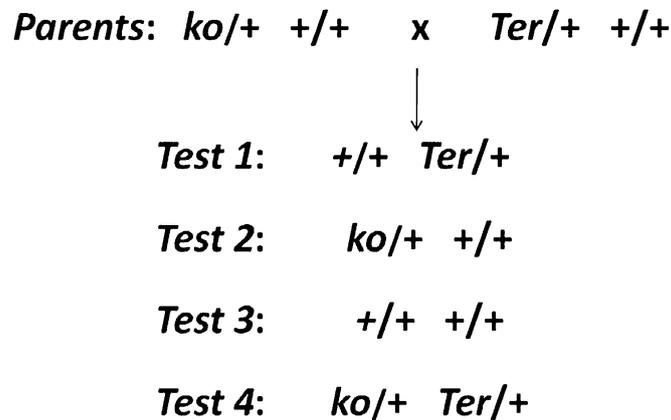


Fig. S1. Design of test for transgenerational gene interactions. Each parent was heterozygous for either *Apobec1^{ko}* or *Dnd1^{Ter}*, and the resulting offspring consisted of four genotypic classes: a test group of double-heterozygotes, and three control groups, each of the single heterozygotes and the wild-type homozygotes (see ref. 1 for details.)

1. Rassoulzadegan M, et al. (2006) RNA-mediated non-Mendelian inheritance of an epigenetic change in the mouse. *Nature*441:469–474.

Table S1. Occurrence of males with a testicular germ cell tumor in various *Apobec1^{ko}* and *Dnd1^{Ter}* crosses

Male offspring genotype	Parental cross	N	No. affected	No. expected	Percent affected	Percent expected	P
<i>ko/+, Ter/+</i>	Total	189					
	<i>Ter/+</i> x <i>ko/+</i>	25	5	1.8	20.0	7.3	0.01
	<i>ko/+</i> x <i>Ter/+</i>	11	5	1.6	45.5	14.1	0.003
	<i>Ter/+</i> , <i>ko/+</i> x 129	10	6	3.4	60.0	34.1	Ns
	129 x <i>Ter/+</i> , <i>ko/+</i>	38	12	12.6	31.6	33.2	Ns
	<i>Ter/+</i> x <i>ko/ko</i>	17	4	6.1	23.5	35.0	Ns
	<i>ko/ko</i> x <i>Ter/+</i>	88	20	31.2	22.7	35.5	Ns
<i>+/+, Ter/+</i>	Total	124					
	<i>Ter/+</i> x <i>ko/+</i>	28	3	8.8	10.7	31.5	0.04
	<i>ko/+</i> x <i>Ter/+</i>	19	2	6.0	10.5	31.5	Ns
	<i>Ter/+</i> , <i>ko/+</i> x 129	26	7	8.2	26.9	31.5	Ns
	129 x <i>Ter/+</i> , <i>ko/+</i>	51	14	16.0	27.5	31.5	Ns
<i>ko/+, +/+</i>	Total	322					
	<i>Ter/+</i> x <i>ko/+</i>	24	2	2.4	8.3	9.2	Ns
	<i>ko/+</i> x <i>Ter/+</i>	17	2	0	11.8	0	Ns
	<i>Ter/+</i> <i>ko/+</i> x 129	30	4	0	13.3	0	Ns
	129 x <i>Ter/+</i> , <i>ko/+</i>	69	6	6.4	8.7	9.2	Ns
	<i>Ter/+</i> x <i>ko/ko</i>	66	9	5.3	13.6	8.1	Ns
	<i>ko/ko</i> x <i>Ter/+</i>	116	8	0	6.9	0	Ns
<i>+/+, +/+</i>	Total	213					
	<i>Ter/+</i> x <i>ko/+</i>	34	4	2.5	12.0	7.2	Ns
	<i>ko/+</i> x <i>Ter/+</i>	61	5	4.4	8.2	7.2	Ns
	<i>Ter/+</i> , <i>ko/+</i> x 129	49	3	3.5	6.1	7.2	Ns
	129 x <i>Ter/+</i> <i>ko/+</i>	69	2	5.0	2.9	7.2	Ns

Table S2. Segregation ratios from heterozygous *Apobec1^{ko}* and *Dnd1^{Ter}* parents at two embryonic stages (embryonic days E3.5 and E12.5) and at weaning

	Time point	<i>Apobec1</i> +/+		<i>Apobec1</i> ko/+		Litter size	N
		+/+	<i>Ter</i> /+	+/+	<i>Ter</i> /+		
<i>Apobec1^{ko/+}</i> ♀ x <i>Dnd1^{Ter/+}</i> ♂	E3.5	13	3	3	1	5.0	20
	E12.5	16	7	10	2	7.2	35
	Weaning	89	37	23	17	6.1	166
<i>Dnd1^{Ter/+}</i> ♀ x <i>Apobec1^{ko/+}</i> ♂	E3.5	6	7	6	3	7.3	22
	Weaning	41	43	54	58	5.4	196

Table S3. Litter size and sex ratio for intercrosses between *Apobec1^{ko}* and *Dnd1^{Ter}* heterozygotes

	<i>Apobec1</i> control cross		<i>Dnd1</i> control cross		<i>Apobec1</i> – <i>Dnd1</i> interaction	
	<i>ko</i> /+	+/+	<i>Ter</i> /+	+/+	<i>ko</i> /+	<i>Ter</i> /+
Maternal genotype	<i>ko</i> /+	+/+	<i>Ter</i> /+	+/+	<i>ko</i> /+	<i>Ter</i> /+
Paternal genotype	+/+	<i>ko</i> /+	+/+	<i>Ter</i> /+	<i>Ter</i> /+	<i>ko</i> /+
Average litter size	7.15	7.09	7.23	6.27	6.05	5.40
No. females	187	128	70	127	64	123
No. males	210	120	90	137	69	109
% females	47	52	44	48	48	53