

Genetic Interaction Between *Wnt7a* and *Lrp6* During Patterning of Dorsal and Posterior Structures of the Mouse Limb

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In the published version of this article, Table 1 was printed incorrectly. Please find the correct version of Table 1 printed herein.

TABLE 1. Genotypes of Offspring From the Cross *Wnt7a*^{+/-} × *Wnt7a*^{+/-} *Lrp6*^{+/-}*

Parental		Progeny					
		Nonrecombinant		Nonrecombinant + recombinant		Recombinant	
<i>Wnt7a</i> ⁺ <i>Lrp6</i> ⁺	<i>Wnt7a</i> ⁺ <i>Lrp6</i> ⁻	<i>Wnt7a</i> ⁻ <i>Lrp6</i> ⁺	<i>Wnt7a</i> ⁺ <i>Lrp6</i> ⁻	<i>Wnt7a</i> ⁺ <i>Lrp6</i> ⁺	<i>Wnt7a</i> ⁺ <i>Lrp6</i> ⁻	<i>Wnt7a</i> ⁺ <i>Lrp6</i> ⁺	<i>Wnt7a</i> ⁻ <i>Lrp6</i> ⁻
<i>Wnt7a</i> ⁻ <i>Lrp6</i> ⁺	<i>Wnt7a</i> ⁻ <i>Lrp6</i> ⁺	<i>Wnt7a</i> ⁻ <i>Lrp6</i> ⁺	<i>Wnt7a</i> ⁺ <i>Lrp6</i> ⁺	<i>Wnt7a</i> ⁻ <i>Lrp6</i> ⁺	<i>Wnt7a</i> ⁻ <i>Lrp6</i> ⁺	<i>Wnt7a</i> ⁺ <i>Lrp6</i> ⁺	<i>Wnt7a</i> ⁻ <i>Lrp6</i> ⁺
		43	35	56	45	11	7

*Observed genotype frequencies are consistent with linkage of *Wnt7a* and *Lrp6* with a distance of 20 cM.

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