

Supplementary tables

Mapping of a responsible region for sex reversal upstream of *Sox9* by production of mice with serial deletion in a genomic locus

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Table S2. Genotypes of F₀ mice obtained by injection of sgRNA2/6/7.

Mouse ID	Karyotype	Phenotypic sex	Indel			Deletion		
			F1/R1	F2/R2	F6/R6	F1/R2 (Δ 7-2)	F2/R6 (Δ 2-6)	F1/R6 (Δ 7-6)
#41	XY	Male	indel	indel	indel	○		
#42	XY	Male	indel	indel	indel		○	○
#43*	XX	Female	ND	ND	ND			
#44	XX	Female	ND	ND	indel	○		
#45	XX	Female	indel	indel	indel			○
#46	XY	Female	ND	ND	indel	○		○
#47	XY	Male	indel	indel	indel			
#48	XY	Female	ND	ND	indel	○		○
#49	XX	Female	indel	ND	ND	○		○
#50	XY	Male	indel	indel	indel			
#51	XY	Male	indel	indel	indel			○
#52	XX	Female	ND	indel	indel			
#53	XX	Female	indel	ND	ND		○	
#54	XY	Female	ND	ND	ND			○
#55	XY	Male	indel	indel	indel			
#56	XY	Male	ND	indel	indel			
#57	XY	Male	ND	ND	indel	○		

Columns below 'Indel' and 'Deletion' show pairs of PCR primers used for genotyping.

Parentheses designate deleted regions.

WT: no mutation observed; indel: indel was observed; ND: not detected by PCR amplification; ○: PCR product was identified; Blank: PCR product was not identified

Mice exhibiting XY sex reversal are highlighted in yellow.

*PCR amplicon was not observed by any combinations of genotyping primers.

Table S3. Germline transmission of serial deletion alleles.

sgRNA combination	F ₀ mice	Genotype of F ₀	No. of F ₁ genotyped	No. of F ₁ mouse	
				WT	Deletion*
sgRNA1/2/3/4/5/6	#6	$\Delta 2\text{-}3/\Delta 4\text{-}6$	7	4	1 ($\Delta 2\text{-}3$) , 2 ($\Delta 2\text{-}3$; $\Delta 4\text{-}6$)
	#9	$\Delta 2\text{-}3/\Delta 3\text{-}6/\Delta 5\text{-}6$	8	0	3 ($\Delta 3\text{-}6$) , 2 ($\Delta 2\text{-}3$), 3 ($\Delta 2\text{-}3$; $\Delta 3\text{-}6$)
	#16	$\Delta 2\text{-}6/\Delta 2\text{-}4$	9	1	2 ($\Delta 2\text{-}4$) , 5 ($\Delta 2\text{-}6$)
	#21	$\Delta 2\text{-}5$	7	6	1 ($\Delta 2\text{-}5$)
	#25	$\Delta 4\text{-}6$	5	3	2 ($\Delta 4\text{-}6$)
	#26**	$\Delta 2\text{-}5$	9	0	2 ($\Delta 2\text{-}5$), 7 ($\Delta 5\text{-}6$)
	#33	$\Delta 2\text{-}3/\Delta 4\text{-}6$	3	2	1 ($\Delta 2\text{-}3$)
	#34	$\Delta 2\text{-}4$	14	11	3 ($\Delta 2\text{-}4$)
	#35	$\Delta 5\text{-}6$	7	5	2 ($\Delta 5\text{-}6$)
	#37**	$\Delta 3\text{-}4$	11	5	6 ($\Delta 2\text{-}5$)
sgRNA2/6/7	#45	$\Delta 7\text{-}6$	9	5	4 ($\Delta 7\text{-}6$)
	#57	$\Delta 7\text{-}2$	10	6	4 ($\Delta 7\text{-}2$)

*Established deletion lines are shown in bold.

**Deletion patterns observed at F₁ generation was not identified in F₀ mice. #26 and #37 are considered as mosaic individuals.