## **Supplementary Data**

CRISPR/Cas9 microinjection in oocytes disables pancreas development in sheep

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PDX1	gRNA Target Site L Sheep Locus 5 <sup>,</sup> -gggctggcgctgaagtctggcgcccgggccccgggcccggggcccggggccctgaagtctggggccctggaagtccttgtacagctgtgggccc-	3'
MII oocytes	Mono-allelic mutations Mil#1 5'-GGGCTGGCGCTGAAGTCTGGCGCCGGGCCCCGCTGGAACGCGCAGGGGTCCTTGTACAGCTGTGTGGCCG-3' 5'-GGGCTGGCGCTGAAGTCTGGCGCCGGGCCCCGCTGGAACGCGGGGTCCTTGTACAGCTGTGTGGCCG-3' Bi-allelic mutations	(WT) (-3)
	MII#5 5'-GGGCTGGCGCTGAAGTCTGGCGTCGTTGTACAGCTGTGGGGGCCG-3' 5'-GGGCTGGCGCTGAAGTCTGGCGTCGTTGTACAGCTGTGGGGCCCG-3'	(-27;+1) (-27;+1)
	MII #15 5'-GGGCTGGCGCTGAAGTCTGGCGCCGGGGCCCCGCTGGAACGGGTCCTTGTACAGCTGTGGGCCGCGTA-3' MII #16	(-6) (-6)
	5'-GGGCTGGCGCTGAAGTCTGGCGCCCGGTGCAGCTGGGGGGGG	(7subs.;+18;-27) (7subs.;+18;-27)
	5'-GGGCTGGCGCTGAAGTCTGGCGCCGGGGCCCCGGTGGAACCCAGGGGTCCTTGTACAGTTGGGGGGGCCC-3' 5'-GGGCTGGCGCTGAAGTCTGGCGCCGGGCCCCGGCGGACCCAGGGGTCCTTGTACAGTTGGGGGGGCCC-3'	(5subs.;+1;-3) (5subs.;+1;-3)
	Wiii #19 5'-ACAGGCACGCAGGGGGGCGCGGGGCGCGAAGGGGTCCTTGTGGGCCCTGTGTGGGCCC-3' 5'-ACAGGCACGCAGGGGGGCCTGGAGGCCCTGAAGGGGTCCTTGTGGGCCCTGTGTGGGCCC-3'	(-30;4ins.) (-30;4ins.)
	MII#23 5GGGCTGGCGCTGAAGTCTGGCGCCGGGGCCCCGGGACGGCATGAGTCCTTGTACAGCTGTGTGGCCG-3' 2GGGCTGGCGCTGAAGTCTGGCGCCGGGCCCCGGGACGGCATGAGTCCTTGTACAGCTGTGTGGCCG-3'	(2subs.) (2subs.)
Zygotes	Mono-allelic mutations	
	Z. #3 5'-GGGCTGGCGCTGAAGTGGGGTCCTTGGACCCGTGGGTGGCCGCATGGTACGGGTCCTTGTACATCTGTGTGGCCG-3' 5'-GGGCTGGCGCTGAAGTCTGGCGCCGGGGCCCCGGGGGCCCGCAGGGGTCCTTGTACAGCTGTGGGCCG-3'	(-29) (WT)
	2.93 3'-GGGCTGGCGCTGAAGTCTGGCGCCGGGGCCCCGCTGGAACGCGCAGGGGTCCTTGTACAGCTGTGTGGCCGCCTAGT-3' 5'-GGGCTGGCGCTGAAGTCTGGCGCCGGGGCCCCGCTGGAACGCGCAGGGGTCCTTGTACAGCTGTGTGGCCG-3' 2'20	(1subs.) (WT)
	2: -GGGCTGGTGGCGCATGTAGAGGAAGCCAGGGGGGGGGGG	(-26) (WT)
	Drament inductions   2.#2:   S'-GGGCTGGCGCTGAAGTCTGGCGCCGGGTCCCGGCTGCTGCGGGGGCCGCGTAGTACTGCTCCGCGCAGATCAAAAACTGCTCCTCGC-3'   S'-GGGCTGGCGCTGAAGTCTGGCGCCGGGTCCCGGTCCCGCTGCTGCGGGGCCGCGTAGTACTGCTCCGCGCAGATCAAAAACTGCTCCTCGC-3'	(3subs.;+21;-24) (3subs.;+21;-24)
	L: #12 5'-GGGCTGGCGCTGAAGTCAGGGGTCCTTGTACAGCTGTGTGGCCG-3' 5'-GGGCTGGCGCTGAAGTC	(-26) (-26)
	L.## 5'-GGGCTGGCGCTGAAGTGGGGTCCTTGGACCCGTGGGTGGCCGCATGGTACGGGTCCTTGTAGATCTGTGTGGCCG-3' 5'-GGGCTGGCGCTGAAGTGGGGTCCTTGGACCCGTGGGTGGCCGCATGGTACGGGTCCTTGTAGATCTGTGTGGCCG-3'	(-29;1subs.) (-29;1subs.)

Supplementary Figure S1. Genotypes of the PDX1 edited embryos injected at the MII oocyte or zygote stage. Sanger sequencing results from bi-allelic and a mono-allelic mutant sheep blastocyst are shown. The PAM sequence is underlined and the target region is shown in blue. Red dashes represent deletions and red letters insertions/substitutions.



**Supplementary Figure S2.** The two left panels are macroscopic appearance of the vestigial pancreas (arrowheads) of the PDX1-KO fetus and the pancreas of a WT fetus (black dashes) at 4 months old of gestation. St.: stomach; D.: duodenum. The right panel is representative images (100X) of the pancreas and vestigial structure stained with hematoxylin and eosin.



**Supplementary Figure S3.** Confocal microscopy of PDX1 (green) and Insulin (red) double immunostaining, and DAPI staining (blue) of a PDX1-KO 4 month-old fetus compared to a WT fetus of the same age. Scale bars in overviews 500 µm, in details 50 µm.



2 – Wild type 3 – Bi-allelic 5 – Mono-allelic

**Full gel image for Figure 4a.** Gel electrophoresis of PCR product of Sheep embryos injected with PDX1 sgRNA 1 & PDX1 sgRNA2.



**Full gel image for Figure 4d.** Gel electrophoresis of PCR product -using specific primers for *PDX1-* from different tissues (liver, lung, heart, kidney, muscle and spleen) of the mutant fetus.

Replicate	IVP method	Group	Lysis rate (%)	Blastocyst rate (%)
1	IVF	Control		7/41 (17.1%)
		MII	4/82 (4.9%)	14/78 (17.9%)
		Zygote	13/74 (17.6%)	4/61 (6.6%)
2	IVF	Control		21/97 (21.6%)
		MII	3/66 (4.5%)	12/63 (19%)
		Zygote	7/67 (10.4%)	6/60 (10.0%)
3	IVF	Control	· · ·	19/31 (61.3%)
		MII	1/58 (1.7%)	25/57 (43.9%)
		Zygote	14/85 (16.5%)	13/71 (18.3%)
4	PA	Control		20/50 (40%)
		MII	2/63 (3.2%)	20/61 (32.8%)
		Zygote	1/50 (2%)	9/49 (18.4%)
5	PA	Control		20/57 (35.1%)
		MII	0/110 (0%)	29/110 (26.4%)
		Zygote	12/100 (12%)	21/88 (23.9%)

Table S1. Lysis and development rate after microinjection of MII oocytes and Zygotes. Embryos were produced by *in vitro* fertilization (IVF) or parthenogenetic activation (PA).

Table S2. Oligos/Primers used in this study (5' - 3'). Underlined sequences are: gRNAs (Oligos for gRNA synthesis); and 16 bp barcodes (Primers for NGS).

Oligos for gRNA synthesis					
Oligo name	Sequence				
oPDX1-single gRNA	GAAATTAATACGACTCACTATAGGG <u>GGCCCCGCTGGAACGCGCAG</u> GTTTTAGAGCTAGAAATAGC				
oPDX1- dual gRNA1	TAATACGACTCACTATA <u>GCGTACGGGGGGGGGGGGGGGG</u>				
oPDX1- dual gRNA2	TAATACGACTCACTATA <u>GCACGCGTGGAAAGGCCAGT</u> GTTTTAGAGCTAGAAATAGC				
T7-Reverse constant	AAAAGCACCGACTCGGTGCCACTTTTTCAAGTTGATAACGGACTAGCCTTATTTTAACTTGCTATTTCTAGCTCTAAAAC				
Primers for PCR amplification					
Primer name	Sequence				
oPDX1-F	GAACCGCGAGGAGCAGTA				
oPDX1-R-single gRNA	GAGCGGAGGCACCTCGTAT				
oPDX1-R-dual gRNA	CGACGGCACTGAGGAGTC				
Primers for NGS					
Primer name	Sequence				
oPDX1-F-BC1	<u>TCAGACGATGCGTCAT</u> GAACCGCGAGGAGCAGTA				
oPDX1-F-BC17	<u>CATAGCGACTATCGTG</u> GAACCGCGAGGAGCAGTA				
oPDX1-F-BC29	<u>GCTCGACTGTGAGAGA</u> GAACCGCGAGGAGCAGTA				
oPDX1-F-BC34	<u>ACTCTCGCTCTGTAGA</u> GAACCGCGAGGAGCAGTA				
oPDX1-F-BC38	<u>TGCTCGCAGTATCACA</u> GAACCGCGAGGAGCAGTA				
oPDX1-F-BC40	<u>CAGTGAGAGCGCGATA</u> GAACCGCGAGGAGCAGTA				
oPDX1-R-BC48	<u>TCACACTCTAGAGCGA</u> GAGCGGAGGCACCTCGTAT				
oPDX1-R-BC52	<u>GCAGACTCTCACACGC</u> GAGCGGAGGCACCTCGTAT				
oPDX1-R-BC54	<u>GTGTGAGATATATATC</u> GAGCGGAGGCACCTCGTAT				
oPDX1-R-BC62	<u>GACAGCATCTGCGCTC</u> GAGCGGAGGCACCTCGTAT				
oPDX1-R-BC70	<u>CTGCGCAGTACGTGCA</u> GAGCGGAGGCACCTCGTAT				
oPDX1-R-BC09	<u>CTGCGTGCTCTACGAC</u> GAGCGGAGGCACCTCGTAT				