

**Supplemental Figure 1. Human cultured podocytes constitutively express SGLT2.** Representative images of SGLT2 membrane expression (red) in RPTEC used as positive controls (left) and control podocytes (right) under basal conditions. Nuclei were counterstained with DAPI (blue). Original magnification (X630).



**Supplemental Figure 2. Gene silencing of SGLT2 in human cultured podocytes** (A) *SGLT2* mRNA expression evaluated by qRT-PCR analysis in podocytes transfected with SGLT2 siRNA (si-*SGLT2*; 15 nM) or irrelevant siRNA (si-irrelevant; 15 nM) for 24 hours before albumin (10 mg/ml) stimulation (3 hours). *HPRT1* was used as endogenous control. *SGLT2* levels were normalized to *HPRT1* levels and reported as fold change relative to podocytes (untransfected) exposed to albumin. Data are mean ± SEM (n=3 samples) and analyzed by ANOVA with Tukey post hoc test. (B) Representative Western blot of SGLT2 protein in podocytes transfected with si-*SGLT2* (15 nM) or si-irrelevant (15 nM) for 24 hours before albumin (10 mg/ml) stimulation (6 hours). Actin was used as sample loading control. (C) Representative images of SGLT2 membrane expression in podocytes transfected with SGLT2 or irrelevant siRNA before albumin stimulation (3 hours). Nuclei were counterstained with DAPI (blue). Original magnification (X630).

	S	equencing primer	forward			
481	gatctcagtg	gacatgttct	ccggagctgt	attcatccag	caggctctgg	gctggaacat
541	ctatgcctcc	gtcatcgcgc	ttctgggcat	caccatgatt	tacacggtga	caggagggct
601	ggccgcgctg	atgtacacgg	acacggtaca	gaccttcgtc	attctggggg	gcgcctgcat
		Exon 6-7 junction				
	gRT-PC	CR pri <u>me</u> r forward	<u> </u>			
661	cctcatgggt	tacgccttcc	acgaggtggg	cgggtattcg	ggtctcttcg	acaaatacct
			qRT-PCR primer reverse			
721	gggagcagcg	acttcgctga	cggtgtccga	ggatccagcc	gtgggaaaca	tctccagctt
781	ctgctatcga	ccccggcccg	actcctacca	cctgctccgg	caccccgtga	ccggggatct
	Sequencing primer reverse					
841	gccgtggccc	gcgctgctcc	tcggactcac	aatcgtctcg	ggctggtact	ggtgcagcga



**Supplemental Figure 3. (A)** The portion of human solute carrier family 5 member 2 (*SLC5A2; SGLT2*), transcript variant 1, mRNA reference sequence (NM\_003041.3 at National Center of Biotechnology Information) which comprises the qRT-PCR product (highlighted in grey). Exon 6-7 junction is indicated by red arrow. The position and sequence of qRT-PCR *SGLT2* primers and sequencing primers is indicated. **(B)** Starting from podocyte cDNA, a portion of *SGLT2* was amplified and sequenced in the two directions on ABI 3730 DNA analyser. Alignment of the sequences corresponding to the qRT-PCR amplification product with the *SGLT2* mRNA reference sequence. The inset shows the chromatograms relative to the region that includes exon 6-7 junction.

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