

Supplementary File

1. Number of animals in different subgroups of the experiment

Placebo, 5nM Relaxin and 20 nM Relaxin were used for organ flushing and preservation in 8, 5 and 6 donor pigs, respectively. The numbers of kidneys in different subgroups of the experiment are presented in Table S1.

Subgroups of the experiment	Number of animals, n
Placebo 24 h SCS	n=8
Placebo 48 h SCS	n=8
Relaxin 5nM 24 h SCS	n=5
Relaxin 5nM 48 h SCS	n=5
Relaxin 20nM 24 h SCS	n=6
Relaxin 20nM 48 h SCS	n=6

Table S1. The numbers of kidneys in different subgroups. SCS: static cold storage.

2. Subgroup analysis

2.1 24. vs. 48 hours of SCS in Placebo and RLX groups

Oxidative stress-related genes (Figure S1) and apoptosis, necroptosis-related genes (Figure S2) expression, immunohistochemistry (Table S1), graft function representing parameters (Figure S3) and graft survival (Figure S4) was similar between the subgroups of grafts stored for 24 and 48 hours within the RLX and placebo groups.

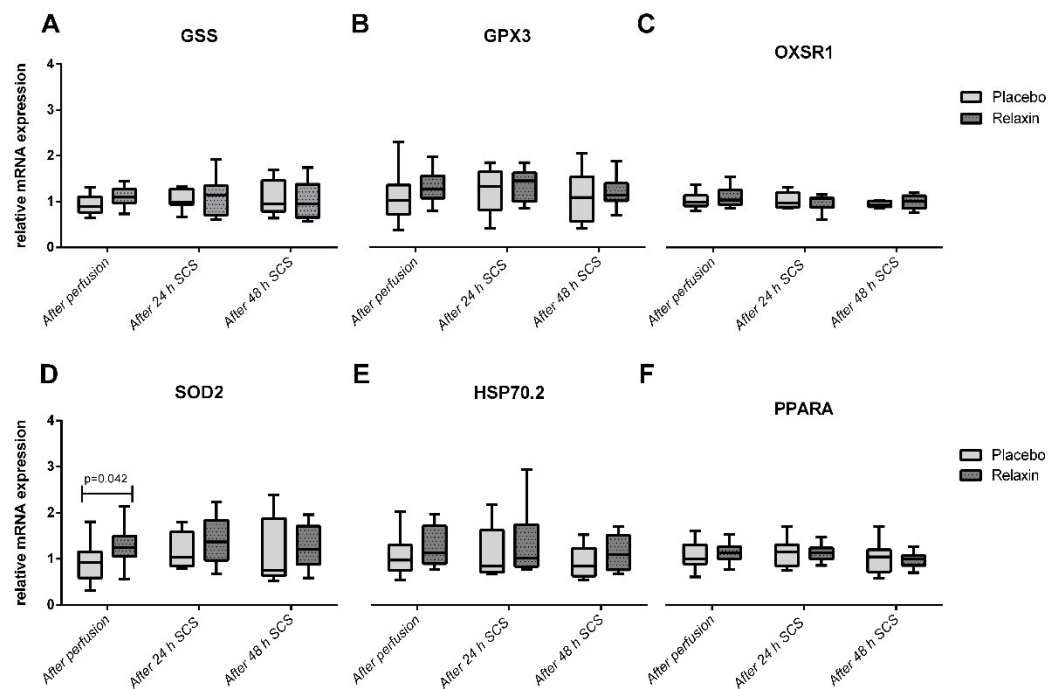


Figure S1. Oxidative stress-related genes expression in kidney tissue after perfusion and static cold storage in Relaxin and Placebo groups for 24 and 48 hours. GSS: Glutathione Synthetase; GPX3: Glutathione Peroxidase 3; OXSR1: Oxidative Stress Responsive Kinase 1; SOD2: Superoxide Dismutase 2; HSP70.2: Heat Shock Protein 70.2; PPARA: Peroxisome Proliferators Activated Receptor Alpha.

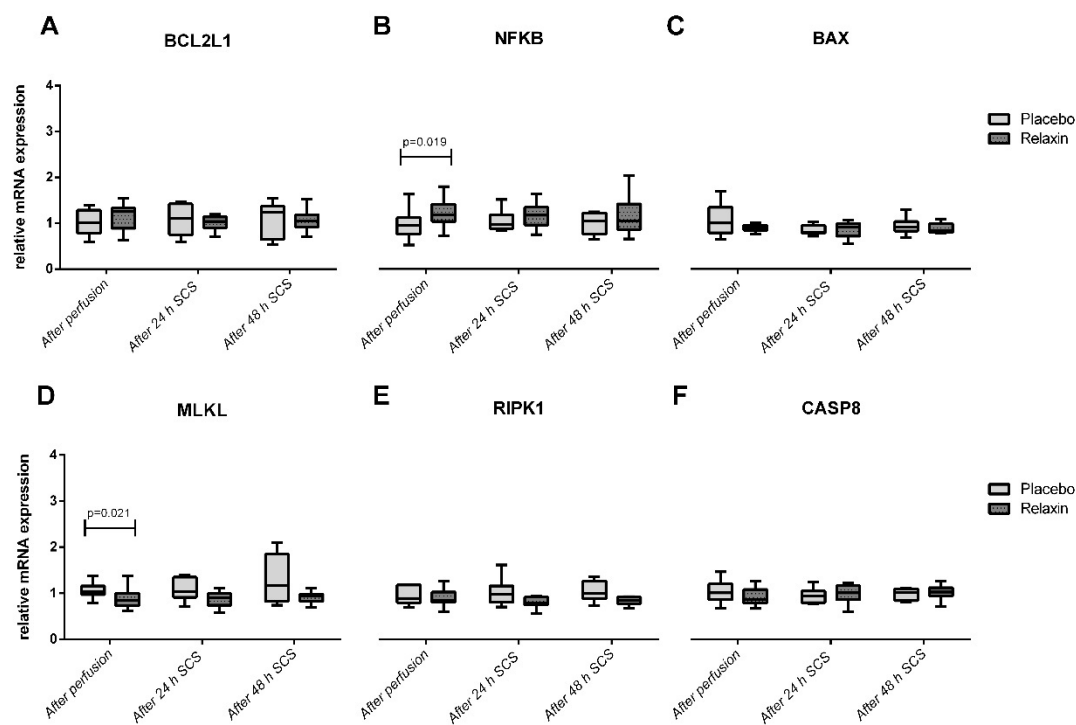


Figure S2. Apoptosis and necroptosis related genes expression in kidney tissue after perfusion and static cold storage in Relaxin and Placebo groups for 24 and 48 hours. BCL2L1: BCL2 Like 1; NFKB: Nuclear Factor of Kappa Light Polypeptide Gene Enhancer In B-cells; BAX: BCL2 Associated X Protein; MLKL: Mixed-lineage kinase domain-like protein; RIPK1: Receptor Interacting Serine/Threonine Kinase 1; CASP8: Caspase 8.

		Caspase 3			Myeloperoxidase		
		RLX	Placebo	p value	RLX	Placebo	p value
<i>Kidney</i>	After perfusion	0 (0; 0)	0 (0; 0)	0.999	0 (0; 0)	0 (0; 0)	0.999
	After 24 hours SCS	0 (0; 1)	1 (0.2; 1.5)	0.068	0 (0; 0)	0.5 (0; 1.25)	0.177
	After 48 hours SCS	0 (0; 0)	1.1 (0; 1.9)	0.051	0 (0; 1)	1 (0.25; 1.5)	0.062
<i>Renal artery</i>	After perfusion	0 (0; 0)	0 (0; 0)	0.999	0 (0; 0)	0 (0; 0)	0.999
	After 24 hours SCS	0 (0; 0)	0 (0; 0.5)	0.768	0 (0; 0)	0 (0; 0)	0.999
	After 48 hours SCS	0 (0; 0)	0 (0; 0.6)	0.768	0 (0; 0)	0 (0; 0.5)	0.594
<i>Renal vein</i>	After perfusion	0 (0; 0)	0 (0; 0)	0.811	0 (0; 0)	0 (0; 0)	0.999
	After 24 hours SCS	0 (0; 0.1)	0 (0; 1)	0.513	0 (0; 0)	0 (0; 1.1)	0.284
	After 48 hours SCS	0 (0; 0)	1 (0.3; 1)	0.019	0 (0; 0)	0.5 (0; 1)	0.199
<i>Ureter</i>	After perfusion	0 (0; 0)	0 (0; 0)	0.999	0 (0; 0)	0 (0; 0)	0.999
	After 24 hours SCS	0 (0; 0)	0 (0; 1)	0.240	0 (0; 0)	0 (0; 0)	0.768

	After 48 hours SCS	0 (0; 0)	0.8 (0.1; 2)	0.036	0 (0; 0)	0 (0; 0.5)	0.606
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Table S2. Immunohistochemical evaluation for activated Caspase-3 positivity as well as Myeloperoxidase positivity in kidneys after perfusion and static cold storage for 24 and 48 hours in Placebo and Relaxin groups. Values are median (Quartile 1; Quartile 3);

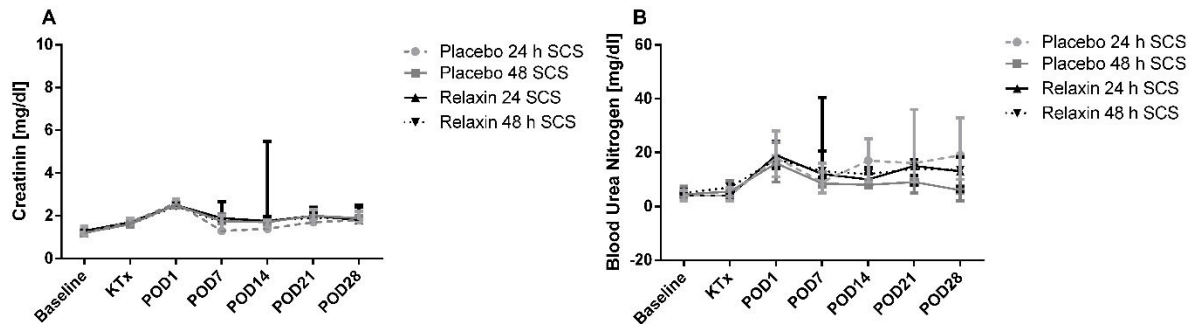


Figure S3. Plasma creatinine (A) and blood urea nitrogen (B) levels after kidney transplantation in Placebo and Relaxin subgroups stored for 24 and 48 hours. POD: postoperative day.

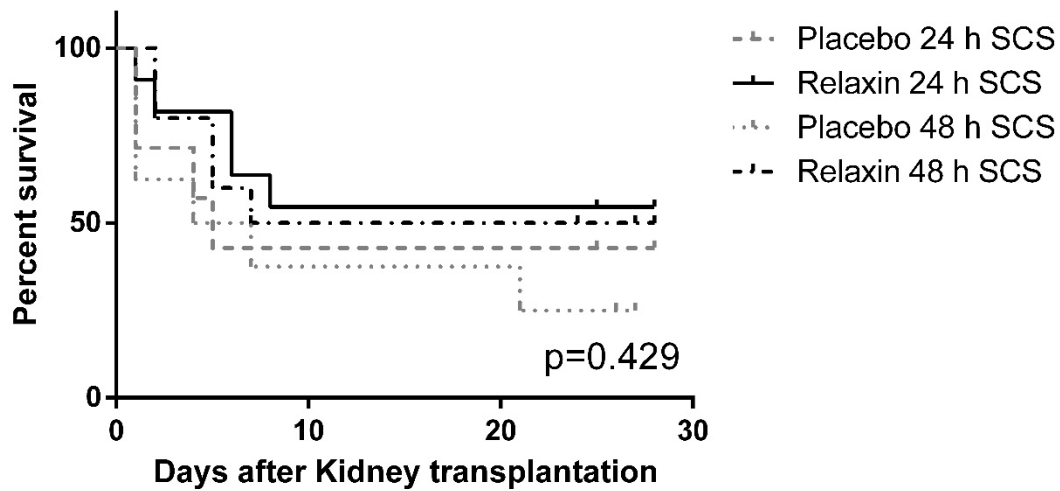


Figure S4. Graft survival after kidney transplantation in Placebo and Relaxin subgroups stored for 24 and 48 hours.

2.2 High vs. low concentration of RLX

Oxidative stress-related genes (Figure S5) and apoptosis, necroptosis-related genes (Figure S6) expression, immunohistochemistry (Table S2), graft function representing parameters (Figure S7) and graft survival (Figure S8) were similar across the animals treated by 5 or 20 nM RLX.

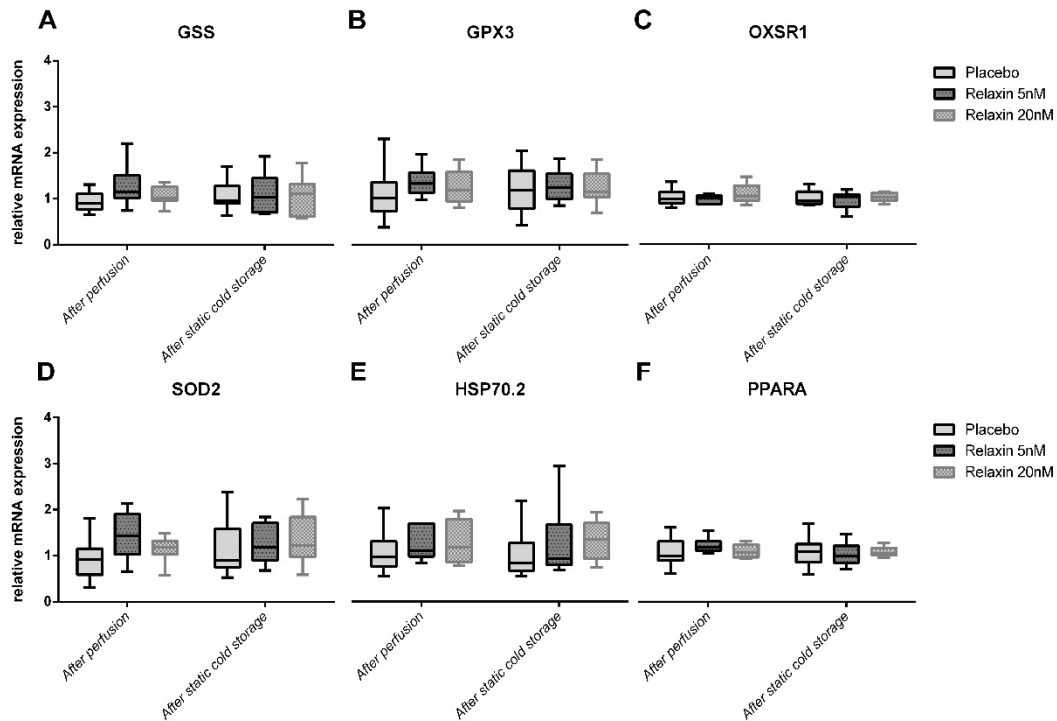


Figure S5. Oxidative stress-related genes expression in kidney tissue after perfusion and static cold storage in Placebo, Relaxin 5nM and Relaxin 20nM subgroups. GSS: Glutathione Synthetase; GPX3: Glutathione Peroxidase 3; OXSR1: Oxidative Stress Responsive Kinase 1; SOD2: Superoxide Dismutase 2; HSP70.2: Heat Shock Protein 70.2; PPARA: Peroxisome Proliferators Activated Receptor Alpha

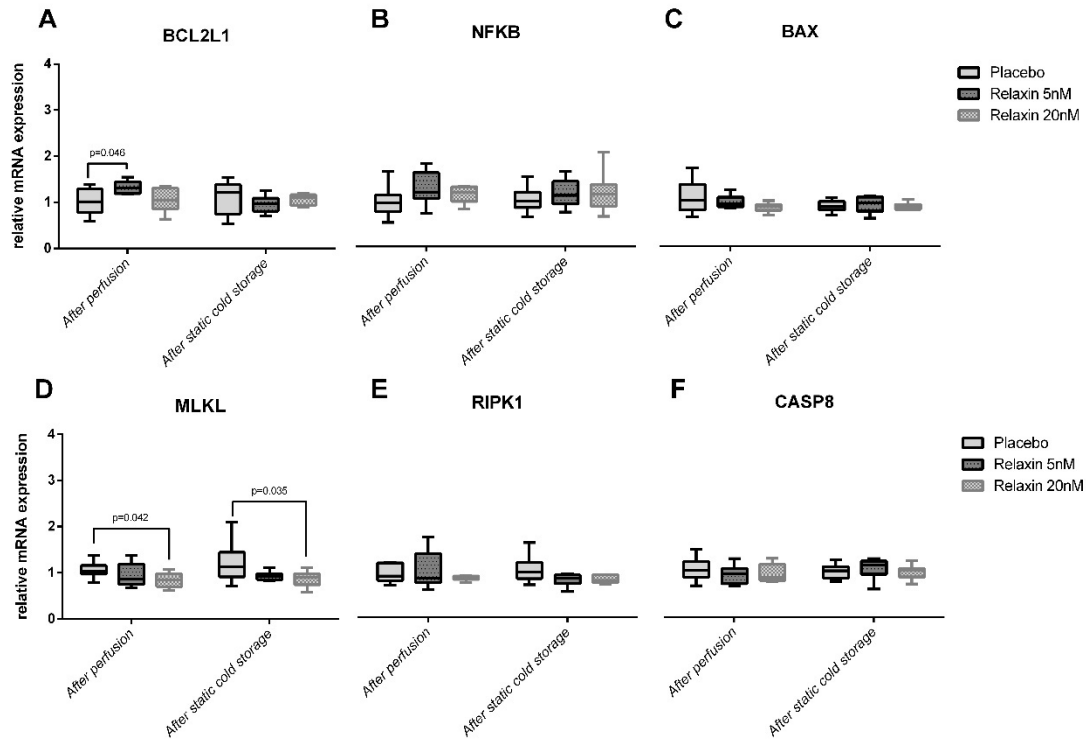


Figure S6. Apoptosis and necroptosis related genes expression in kidney tissue after perfusion and static cold storage in Placebo, Relaxin 5nM and Relaxin 20nM subgroups. BCL2L1: BCL2 Like 1; NFKB: Nuclear Factor of Kappa Light Polypeptide Gene Enhancer In B-cells; BAX: BCL2 Associated X Protein; MLKL: Mixed-lineage kinase domain-like protein; RIPK1: Receptor Interacting Serine/Threonine Kinase 1; CASP8: Caspase 8.

		Caspase 3				Myeloperoxidase			
		RLX 5nM	RLX 20nM	Placebo	p value	RLX 5nM	RLX 20nM	Placebo	p value
Kidney	After perfusion	0 (0; 0)	0 (0; 0)	0 (0; 0)	0.999	0 (0; 0)	0 (0; 0)	0 (0; 0)	0.999
	After SCS	0.5 (0; 1)	0 (0; 0)	1 (0; 1.6)	0.001	0.1 (0; 1)	0 (0; 0)	1 (0; 1.3)	0.009
Renal artery	After perfusion	0 (0; 0)	0 (0; 0)	0 (0; 0)	0.999	0 (0; 0)	0 (0; 0)	0 (0; 0)	0.999
	After SCS	0 (0; 0)	0 (0; 0)	0 (0; 0.2)	0.732	0 (0; 0)	0 (0; 0)	0 (0; 0)	0.329
Renal vein	After perfusion	0 (0; 0.3)	0 (0; 0)	0 (0; 0)	0.449	0 (0; 0)	0 (0; 0)	0 (0; 0)	0.999
	After SCS	0 (0; 0.3)	0 (0; 0)	0.8 (0; 1)	0.013	0 (0; 0)	0 (0; 0)	0 (0; 1)	0.014
Ureter	After perfusion	0 (0; 0)	0 (0; 0)	0 (0; 0)	0.999	0 (0; 0)	0 (0; 0)	0 (0; 0)	0.999
	After SCS	0 (0; 0)	0 (0; 0)	0.6 (0; 1)	0.002	0 (0; 0)	0 (0; 0)	0 (0; 0)	0.626

Table S3. Immunohistochemical evaluation for activated Caspase-3 positivity as well as Myeloperoxidase positivity in kidneys after perfusion and static cold storage in Placebo, Relaxin 5nM and Relaxin 20 nM subgroups. Values are median (Quartile 1; Quartile 3);

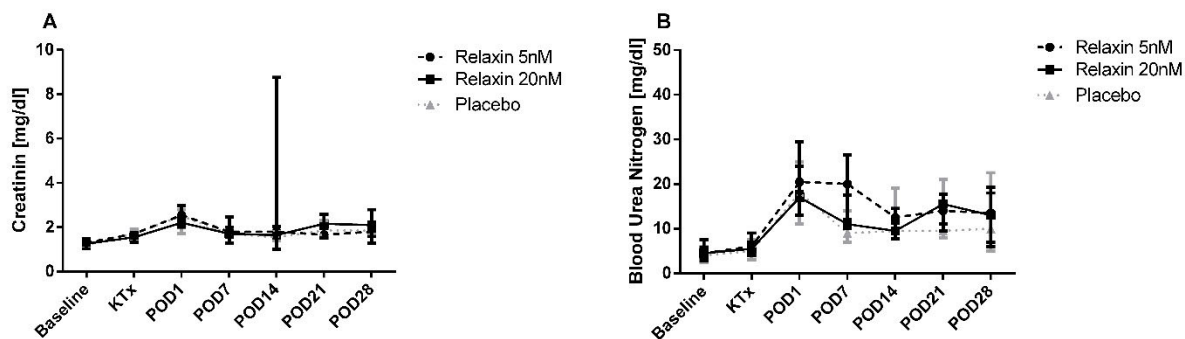


Figure S7. Plasma creatinine (A) and blood urea nitrogen (B) levels after kidney transplantation in Placebo, Relaxin 5nM and Relaxin 20nM subgroups. POD: postoperative day.

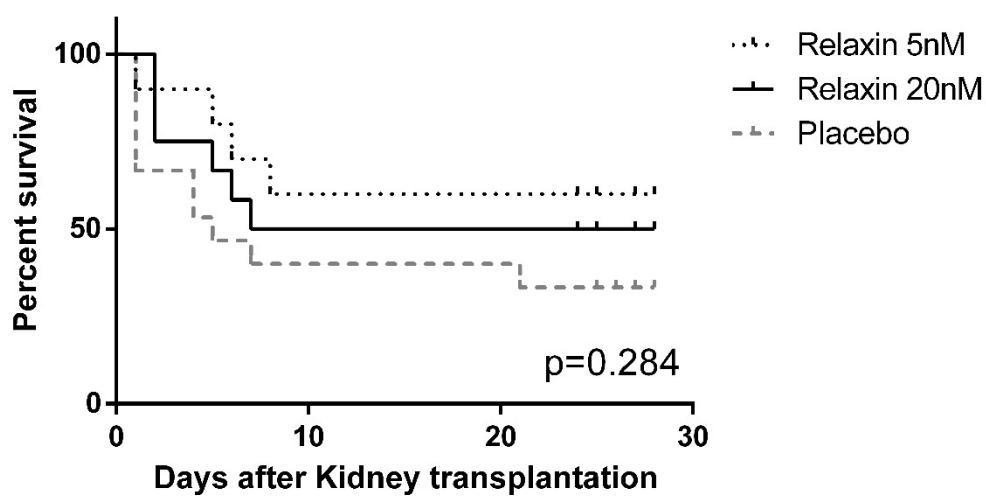


Figure S8. Graft survival after kidney transplantation in Placebo, Relaxin 5nM and Relaxin 20nM subgroups.