Transcript levels of aldo-keto reductase family 1, subfamily C (AKR1C) are increased in prostate tissue of patients with type 2 diabetes

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Supplementary Materials

Table S1. Patient characteristics. Statistical significance was calculated among BEN, noT2D vs BEN, T2D as well as PCa, noT2D vs PCa, T2D samples using Mann-Whitney tests. Statistical significance was considered as p<0.05. BMI: body mass index; n: number of patients; BEN: benign prostate tissue; PCa: prostate cancer tissue.

Parameter	BEN, noT2D	BEN, T2D	BEN	PCa, noT2D	PCa, T2D	PCa
	mean ± SD	mean ± SD	p-value	mean ± SD	mean ± SD	p-value
Age [year]	67 ± 5	63 ± 6	0.0765	65 ± 8	64 ± 3	0.7352
BMI [kg/m²]	27.6 ± 3.3	29.5 ± 4.9	0.3940	26.5 ± 3.2	28.9 ± 3.8	0.1879
Gleason 7a	-	-	-	7	7	-
Gleason 7b	-	-	-	4	4	-
n	17	17	-	11	11	-

Table S2. Relative gene expression levels were measured in benign (BEN) prostate and prostate cancer (PCa) tissues of patients with (T2D) or without (noT2D) type 2 diabetes. Statistical significance was calculated among BEN, noT2D vs BEN, T2D as well as PCa, noT2D vs PCa, T2D samples using Mann-Whitney tests. Significant differences are shown in bold, non-significant differences are indicated in italic. Statistical significance was considered as p<0.05.

Statistical significance was considered as p 4.00.											
Pathway	Gene	BEN, noT2D	BEN, T2D	BEN	PCa, noT2D	PCa, T2D	PCa				
		mean ± SD	mean ± SD	p-value	mean ± SD	mean ± SD	p-value				
HIF1α	ANGPTL4	0.0152 ± 0.0164	0.0146 ± 0.0099	0.4332	0.0078 ± 0.0065	0.0100 ± 0.0101	0.8470				
	GLUT1	0.0925 ± 0.0519	0.0932 ± 0.0557	1.0000	0.0842 ± 0.0327	0.0853 ± 0.0308	0.9873				
	HIF1α	0.6066 ± 0.2650	0.7629 ± 0.3382	0.0730	0.8476 ± 0.2779	0.8516 ± 0.5967	0.1713				
NFκB	BIRC5	0.0016 ± 0.0012	0.0013 ± 0.0010	0.6339	0.0036 ± 0.0020	0.0036 ± 0.0030	0.6994				
	GPX2	0.0220 ± 0.0298	0.0139 ± 0.0101	0.7273	0.0091 ± 0.0110	0.0100 ± 0.0094	0.5190				
	NOS2	0.0014 ± 0.0015	0.0032 ± 0.0047	0.1270	0.0212 ± 0.0563	0.0014 ± 0.0014	0.5943				
	RELA	0.2805 ± 0.1127	0.3498 ± 0.1109	0.0232	0.3421 ± 0.1221	0.3573 ± 0.0509	0.9487				
	SOCS2	0.0100 ± 0.0078	0.0065 ± 0.0056	0.2415	0.0069 ± 0.0083	0.0051 ± 0.0049	0.8977				
proli-	MKI67	0.0146 ± 0.0122	0.0099 ± 0.0066	0.1932	0.0171 ± 0.0079	0.0281 ± 0.0228	0.3653				
feration	PCNA	0.1108 ± 0.0574	0.0978 ± 0.0540	0.1932	0.0939 ± 0.0255	0.1155 ± 0.0259	0.0192				

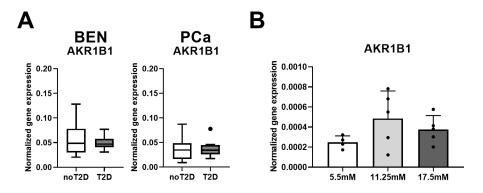


Figure S1. A Gene expression of *AKR1B1* was measured in benign (BEN) prostate (n=17-17) and prostate cancer (PCa) (n=11-11) tissues of patients with (T2D) or without (noT2D) type 2 diabetes. Data are shown as Tukey box plots. Statistical significance was calculated among BEN, noT2D vs BEN, T2D as well as PCa, noT2D vs PCa, T2D samples using Mann-Whitney tests. **B** Gene expression of *AKR1B1* was measured in 5.5-, 11.25- and 17.5 mM glucose treated PC3 cells (n=5 independent experiments). Statistical significance was calculated using paired Friedman test with Dunn's multiple comparison. Statistical significance was considered as p<0.05.