

**Supplemental Data Table 1A.** Relative mRNA expression levels of the uPA system members in tissue of PCa patients

	No. of patients	Median <sup>1</sup>	Range
<b>uPA</b>			
		<b>P = 0.004</b>	
Malignant tissue	132	0.38	0.08 – 11.5
Non- malignant tissue	125	0.51	0.12 – 2.39
<b>uPAR</b>			
		<b>P = 0.604</b>	
Malignant tissue	132	0.47	0.04 – 8.61
Non- malignant tissue	125	0.54	0.00 - 25.8
<b>PAI-1</b>			
		<b>P = 0.261</b>	
Malignant tissue	132	1.75	0.09 – 121.7
Non- malignant tissue	125	1.31	0.09 – 37.4
<b>Ratio uPA/PAI-1</b>			
		<b>P = 0.036*</b>	
Malignant tissue	132	0.21	0.01 – 3.81
Non- malignant tissue	125	0.33	0.03 – 3.15

mRNA expression levels of the *uPA* system members are normalized to *TATA box binding protein (TBP)* gene expression. The Bonferroni-adjusted threshold for significance is at  $\alpha=0.0125$ . <sup>1</sup> Mann-Whitney test, \* Wilcoxon rank test.

**Supplemental Data Table 1B.** Antigen concentrations of the uPA system members in serum of PCa and BPH patients

	No. of patients	Median <sup>1</sup>	Range
<b>uPA</b>			
		<b>P = 0.303</b>	
PCa serum	81	0.68	0.43 – 1.88
BPH serum	36	0.68	0.51 – 2.21
<b>uPAR</b>			
		<b>P = 0.692</b>	
PCa serum	81	0,76	0,20 – 26,2
BPH serum	36	0.79	0.05 – 1.46
<b>PAI-1</b>			
		<b>P = 0.310</b>	
PCa serum	81	1347,9	557,5 – 2714,0
BPH serum	36	1385.5	940.9 – 5930.2
<b>PSA</b>			
		<b>P &lt; 0.001</b>	
PCa serum	81	9.21	1.29 – 51.9
BPH serum	36	1.78	0.22 – 17.5

Antigen concentration in serum is given in ng/ml; The Bonferroni-adjusted threshold for significance is at  $\alpha=0.0125$ . <sup>1</sup> Mann-Whitney-Test.

**Supplemental Data Table 2.** Overview of literature including the present study: Association of tissue RNA or protein levels of uPA gene family members with clinicopathological parameters in prostate cancer.

Factor	Number of patients	Method	Significant association / Remarks	Reference
<b>Tissue RNA</b>				
<i>uPA</i>	13 locally recurrent hormone refractory PCa PCa cell lines: PC-3, LNCaP, DU145	qPCR; Southern hybridization	<i>uPA</i> mRNA with <i>uPA</i> gene amplification in refractory PCa PC-3 with gene amplification more sensitive to uPA inhibitor	Helenius <i>et al.</i> 2001
<i>uPA, uPAR</i>	25 high grade PCa	<i>in situ</i> hybridization; IHC	<i>uPA</i> and <i>uPAR</i> expression in PCa and BPH; <i>uPA</i> and <i>uPAR</i> expression predominantly in adenocarcinoma cells of high grade PCa; Stromal cells were mostly negative for both <i>uPA</i> & <i>uPAR</i>	Gavrilov <i>et al.</i> 2001
<i>uPA, uPAR, PAI-1</i>	44 PCa, 23 BPH	qPCR	Increased expression of <i>uPA</i> and <i>PAI-1</i> in malignant compared with non-malignant prostate tissue; <i>uPAR</i> and <i>PAI-1</i> with higher GS	Riddick <i>et al.</i> 2005
<i>uPAR, PAI-1</i>	stromal cell cultures		<i>uPAR</i> and <i>PAI-1</i> were primarily expressed by stromal cells	Riddick <i>et al.</i> 2005
<i>uPA, uPAR, PAI-1</i>	16 PCa , 9 BPH	<i>in situ</i> hybridization; IHC	<i>uPA, uPAR</i> and <i>PAI-1</i> expression in both PCa and BPH; <i>uPA</i> and <i>uPAR</i> expression predominantly in macrophages, <i>PAI-1</i> expression mostly in scattered fibroblast-like cells, few in macrophages; No association between <i>uPA, uPAR</i> or <i>PAI-1</i> expression and GS	Usher <i>et al.</i> 2005
<i>uPA, uPAR, PAI-1</i>	132 PCa	qPCR	Strong correlation between expression levels of all three <i>uPA</i> system members; <i>uPA</i> with higher GS	Present study
<b>Tissue Protein</b>				
<i>uPA, uPAR</i>	120 clinically localized PCa	IHC	<i>uPA</i> and <i>uPAR</i> with GS, tumor stage, surgical margin status; <i>uPA</i> with lymph node status	Cozzi <i>et al.</i> 2006
<i>uPA, PAI-1, uPAR</i>	230 clinically localized PCa	IHC	Combined <i>uPA/PAI-1</i> values with extraprostatic extension and seminal vesicle involvement; <i>uPA</i> and <i>PAI-1</i> with biochemical RFS and aggressive disease recurrence; combined <i>uPA/PAI-1</i> values independent predictor	Gupta <i>et al.</i> 2008

			for biochemical RFS and aggressive disease	
uPA, PAI-1, uPAR	153 PCa with organ-confined disease	IHC	uPA and uPAR with tumor stage, GS, lymphatic invasion, surgical margin status, and lymph node metastasis; PAI-1 with tumor stage and surgical margin status; uPA, uPAR and PAI-1 with shorter biochemical RFS; uPA independent predictor for biochemical RFS	Kumano et al. 2009
<b>Blood-based Protein</b>				
uPA, suPAR	72 PCa (65% metastatic), 62 BPH, 54 healthy donors	EIA (Serum)	Increased uPA and uPAR levels in PCa compared to BPH patients; uPA and uPAR with tumor stage and metastasis; Either uPA or uPAR or both with OS	Miyake et al. 1999
uPA, suPAR	429 clinically localized PCa, 19 PCa with lymph node metastasis, 10 PCa with bone metastases, 44 healthy donors	EIA (Plasma)	uPA and uPAR with tumor stage, GS, lymph node metastasis and lymphatic invasion; uPA and uPAR with features of biologically aggressive PCa, disease progression after radical prostatectomy and metastasis; uPA and uPAR in PC patients with bone metastasis>PCa with lymph node metastasis> localized PCa> healthy men	Shariat et al. 2007
uPA, suPAR, PAI-1	423 clinically localized PCa	EIA (Serum)	uPA and uPAR with biochemical RFS	Shariat et al. 2008
suPAR	131 metastatic PCa	TR-FIA (Serum)	Each of the suPAR forms with shorter OS	Almasi et al. 2011
uPA, suPAR, PAI-1	81 PCa, 36 BPH	EIA (Serum)	uPAR with shorter OS	Present study

Table was modified according to Schmitt et al. 2010; if not indicated otherwise, high marker levels were associated with clinical parameters;

Abbreviations: PCa: Prostate Cancer; BPH: benign prostatic hyperplasia; qPCR: quantitative PCR; EIA: Enzyme-linked immunosorbent assay; TR-FIA: Time-resolved fluorescence immunoassay; IHC: immunohistochemistry; GS: Gleason score; OS: Overall survival; RFS: recurrence-free survival