

Supplemental Appendix for

**Deciphering the prognostic and predictive value of urinary CXCL10 in kidney recipients
with BK virus reactivation**

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SUPPLEMENTAL RESULTS

Supplemental Table 1: Oligo sequences and probes used for transcript quantification by qPCR.

Gene	Reference	Sequence	Localization
BKV VP1	J02038	Sense: 5'-TGCTGATATTGTGGCCTGTTACTA-3'	2355-2380
	K00058	Antisense: 5'-CTCAGGC GGATCTTAAAATATCTTG-3'	2438-2414
	V01108	Probe: 5'-FAM-AGCTCTGGAACACAACAGTGGAGAGGCC TAMRA 3'	2383-2410
BAK	Y13231	Sense: 5' CCCACATCTGGAGCAGAGTC 3'	192-212
		Antisense: 5' CAGATGCCATTTTCAGGTCTTG 3'	264-242
		5' FAM CAGGTGACAAGTGACGGTGGCTCCA TAMRA 3'	215-240

Supplemental Table 2: Patient and transplant characteristics in the cross-sectional study.

Variables	N=391 patients
Recipient characteristics	
Men, n (%)	241 (61.6)
Age at transplantation (yr), <i>mean±SD</i>	47.6±15.7
Cause of ESRD	
GN, n (%)	88 (22.5)
Diabetes, n (%)	40 (10.2)
Cystic/hereditary/congenital, n (%)	91 (23.3)
Secondary GN, n (%)	21 (5.4)
Hypertension, n (%)	25 (6.4)
Interstitial nephritis, n (%)	43 (11.0)
Miscellaneous conditions, n (%)	6 (1.5)
Cancer, n (%)	0
Etiology uncertain, n (%)	77 (19.7)
Transplant variables	
Donor age (yr), <i>mean±SD</i> ^a	53.8±17.2
Donor type ^b	
Living donor, n (%)	107 (27.9)
SCD, n (%)	129 (33.6)
ECD, n (%)	148 (38.5)
Retransplantation, n (%)	83 (21.2)
Delayed graft function, n (%)	169 (43.7)
Preformed DSAs, n (%) ^c	178 (46.5)
Immunosuppressive protocol	
Induction therapy, n (%) ^d	369 (98.3)
Basiliximab/Thymoglobulin®, n (%)	186 (49.9)/178 (47.7)
Calcineurin inhibitor, n (%) ^e	379 (98.2)
Cyclosporin/tacrolimus, n (%)	72 (18.7)/307 (79.5)
Purine synthesis inhibitor, n (%) ^e	363 (94.0)
Azathioprine/mycophenolic acid, n (%)	10 (2.6)/353 (91.5)
Mammalian target of rapamycin inhibitor, n (%) ^e	28 (7.3)
Steroids, n (%) ^e	383 (99.2)

Abbreviations: DSAs, donor-specific antibodies; ECD, expanded-criteria donor; ESRD, end-stage renal disease; GN, glomerulonephritis; SCD, standard-criteria donor; SD, standard deviation. ^a Data not available (NA) for 3 patients. ^b NA for 7 patients. ^c NA for 8 patients. ^d NA for 18 patients. ^e NA for 5 patients.

Supplemental Table 3: Patient and transplant characteristics in the nested case-control study including 63 single patients with BKV-DNAemia.

Variables	N=63 patients
Recipient characteristics	
Men, n (%)	35 (55.6)
Age at transplantation (yr), <i>mean±SD</i>	50±17.0
Cause of ESRD	
GN, n (%)	15 (23.8)
Diabetes, n (%)	7 (11.1)
Cystic/hereditary/congenital, n (%)	19 (30.2)
Secondary GN, n (%)	2 (3.2)
Hypertension, n (%)	5 (7.9)
Interstitial nephritis, n (%)	7 (11.1)
Miscellaneous conditions, n (%)	0
Cancer, n (%)	0
Etiology uncertain, n (%)	8 (12.7)
Transplant variables	
Donor age (yr), <i>mean±SD</i>	58±19.0
Donor type ^a	
Living donor, n (%)	16 (26.7)
SCD, n (%)	14 (23.3)
ECD, n (%)	30 (50.0)
Retransplantation, n (%)	10 (15.9)
Delayed graft function, n (%)	26 (41.3)
Preformed DSAs, n (%)	32 (50.8)
Immunosuppressive protocol	
Induction therapy, n (%) ^b	59 (100)
Basiliximab/Thymoglobulin®, n (%)	24 (40.7)/34 (57.6)
Calcineurin inhibitor, n (%)	62 (98.4)
Cyclosporin/tacrolimus, n (%)	5 (7.9)/57 (90.5)
Purine synthesis inhibitor, n (%)	62 (98.4)
Azathioprine/mycophenolic acid, n (%)	1 (1.6)/61 (96.8)
Mammalian target of rapamycin inhibitor, n (%)	2 (3.2)
Steroids, n (%)	63 (100)

Abbreviations: DSAs, donor-specific antibodies; ECD, expanded-criteria donor; ESRD, end-stage renal disease; GN, glomerulonephritis; SCD, standard-criteria donor; SD, standard deviation. ^a Data not available (NA) for 3 patients. ^b NA for 4 patients

Supplemental Table 4: Sample characteristics in the nested case-control study including 63 single patients with BKV-DNAemia. The low-CXCL10 group was defined as uCXCL10/cr ≤12.86 ng/mmol, and the high-CXCL10 group was defined as uCXCL10/cr >12.86 ng/mmol. MVI is defined by the sum of the glomerulitis and peritubular capillaritis scores.

Variables	All patients (N=63)	Low-CXCL10 group (N=40)	High-CXCL10 group (N=23)	P value
Time from transplantation to biopsy (mo), median (IQR)	10 (19.0)	13 (25.5)	6 (8.0)	0.003
Indication of biopsy				
Screening biopsy, n (%)	4 (6.3)	2 (5.0)	2 (8.7)	0.97
Clinically indicated biopsy, n (%)	59 (93.7)	38 (95.0)	21 (91.3)	
Rise in serum creatinine, n (%)	23 (39.0)	16 (42.1)	7 (33.3)	0.70
Proteinuria, n (%)	0	0	0	
De novo DSAs, n (%)	0	0	0	
Control after rejection, n (%)	1 (1.7)	0	1 (4.8)	0.76
BKV-DNAemia, n (%)	35 (59.3)	22 (57.9)	13 (61.9)	0.98
Other, n (%)	0	0	0	
Banff elementary lesions				
i score, mean±SD ^a	0.21±0.60	0.15±0.54	0.30±0.70	0.14
t score, mean±SD	0.90±1.29	0.78±1.21	1.13±1.42	0.37
ci score, mean±SD ^a	1.16±1.07	1.21±1.08	1.09±1.08	0.67
ct score, mean±SD ^a	1.15±1.05	1.18±1.05	1.09±1.08	0.71
i-IFTA score, mean±SD	0.75±1.09	0.80±1.07	0.65±1.15	0.45
ti score, mean±SD	0.54±0.96	0.53±0.96	0.57±0.99	0.92
g score, mean±SD	0.24±0.64	0.28±0.68	0.17±0.58	0.38
ptc score, mean±SD	0.21±0.60	0.10±0.44	0.39±0.78	0.02
MVI score, mean±SD	0.44±1.03	0.38±0.90	0.57±1.24	0.56
cg score, mean±SD	1.10±0.35	1.10±0.38	1.09±0.29	0.89
ah score, mean±SD	2.40±1.06	2.53±1.01	2.17±1.11	0.21
cv score, mean±SD ^b	2.57±1.02	2.57±1.01	2.35±1.04	0.35
Pathologic primary diagnosis				
Inadequate, n (%)	4 (6.3)	2 (5.0)	2 (8.7)	0.97
BKVN, n (%)	15 (23.8)	11 (27.5)	4 (17.4)	0.55
Acute rejection, n (%)	9 (14.3)	4 (10.0)	5 (21.7)	0.36
Normal, n (%)	3 (4.8)	2 (5.0)	1 (4.4)	0.62
Other lesions, n (%) ^c	33 (52.4)	22 (55.0)	11 (47.8)	0.77
BKV infection characteristics				
Time from 1 st blood BKV-positive PCR to biopsy (d), median (IQR)	43 (113.0)	45.5 (127.8)	43 (70.5)	0.84

Time from peak DNAemia to biopsy (d), <i>median</i> (<i>IQR</i>)	0 (40.5)	0 (44.3)	0 (54.0)	0.20
BKV viral load at biopsy (\log_{10} copies/mL), <i>median</i> (<i>IQR</i>)	3.4 (1.8)	3.2 (1.9)	3.4 (1.7)	0.59
Peak BKV viral load (\log_{10} copies/mL), <i>median</i> (<i>IQR</i>)	4.6 (1.8)	4.4 (1.8)	4.8 (1.5)	0.16
Laboratory test results at the time of biopsy				
Serum creatinine ($\mu\text{mol}/\text{L}$), <i>mean</i> \pm <i>SD</i>	163 \pm 51	162 \pm 53	165 \pm 47	0.67
DSAs, <i>n</i> (%) ^d	22 (34.9)	11 (30.6)	11 (47.8)	0.29
Proteinuria/creatininuria ratio (g/g), <i>mean</i> \pm <i>SD</i>	0.38 (0.6)	0.37 (0.6)	0.39 (0.4)	0.19
Urinary CXCL10 (LnCXCL10/cr), <i>median</i> (<i>IQR</i>)	2.21 (1.2)	1.94 (1.3)	3.02 (0.8)	<0.0001

Abbreviations: ah, arteriolar hyalinosis; BKVN, BKV-associated nephropathy; cg, allograft glomerulopathy; ci, interstitial fibrosis; ct, tubular atrophy; cv, chronic vascular changes; DSAs, donor-specific antibodies; g, glomerulitis; i, interstitial infiltrate; IFTA, interstitial fibrosis/tubular atrophy; IQR, interquartile range; Ln, natural logarithm; MVI, microvascular inflammation; PCR, polymerase chain reaction; ptc, peritubular capillaritis; SD, standard deviation; t, tubulitis; ti, total inflammation; v, vasculitis. ^aData not available (NA) for 1 case, ^b NA=8, ^c Including calcineurin inhibitor toxicity, IFTA and recurrent disease, ^d NA=4.

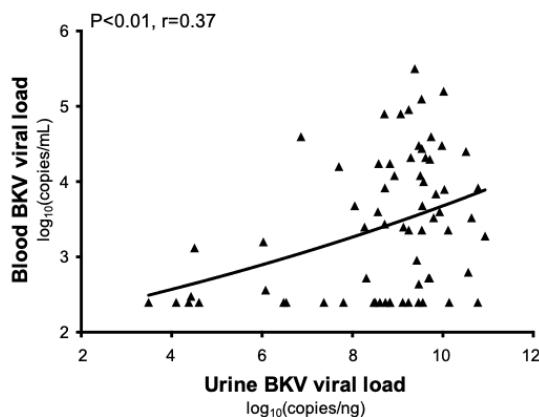
Supplemental Table 5: Patient and transplant characteristics in the longitudinal study including 60 single patients with BKV-DNAemia.

Variables	N=60 patients
Recipient characteristics	
Men, n (%)	36 (60.0)
Age at transplantation (yr), <i>mean±SD</i>	53.6 (15)
Cause of ESRD	
GN, n (%)	15 (25.0)
Diabetes, n (%)	5 (8.3)
Cystic/hereditary/congenital, n (%)	24 (40.0)
Secondary GN, n (%)	4 (6.7)
Hypertension, n (%)	1 (1.7)
Interstitial nephritis, n (%)	2 (3.3)
Miscellaneous conditions, n (%)	2 (3.3)
Etiology uncertain, n (%)	7 (11.7)
Transplant variables	
Donor age (yr), <i>mean±SD</i>	56.3 (18)
Donor type	
Living donor, n (%)	13 (21.7)
SCD, n (%)	16 (26.7)
ECD, n (%)	29 (48.3)
DCD, n (%)	2 (3.3)
Retransplantation, n (%)	17 (28.3)
Delayed graft function, n (%)	8 (13.3)
Preformed DSAs, n (%)	39 (65.0)
Immunosuppressive protocol	
Induction therapy, n (%)	60 (100)
Basiliximab/Thymoglobulin®, n (%)	22 (36.7)/38 (63.3)
Calcineurin inhibitor, n (%)	60 (100)
Cyclosporin/tacrolimus, n (%)	0 (0)/60 (100)
Mycophenolic acid, n (%)	55 (91.7)
Mammalian target of rapamycin inhibitor, n (%)	5 (8.3)
Steroids, n (%)	60 (100)

Abbreviations: DCD, donation after cardiac death; DSAs, donor-specific antibodies; ECD, expanded-criteria donor; ESRD, end-stage renal disease; GN, glomerulonephritis; SCD, standard-criteria donor; SD, standard deviation.

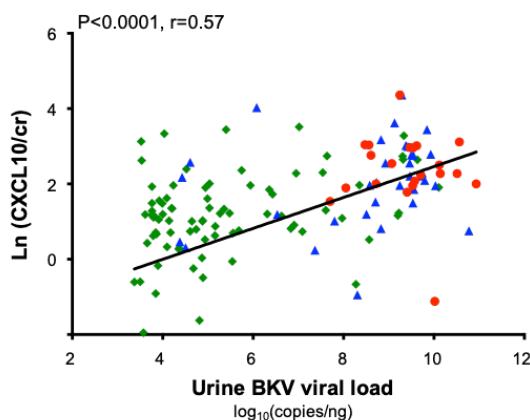
Supplemental Figure 1: Correlation analyses in the cross-sectional study. **(A)** Correlation between blood and urine BKV viral load. **(B)** Correlation between uCXCL10 and urine BKV viral load in a restricted population after exclusion of possible confounders. **(C)** Correlation between uCXCL10 and urine protein output in all samples with BKV DNAemia. Correlations were computed using Pearson's tests. Abbreviations: UTI, urinary tract infection.

A



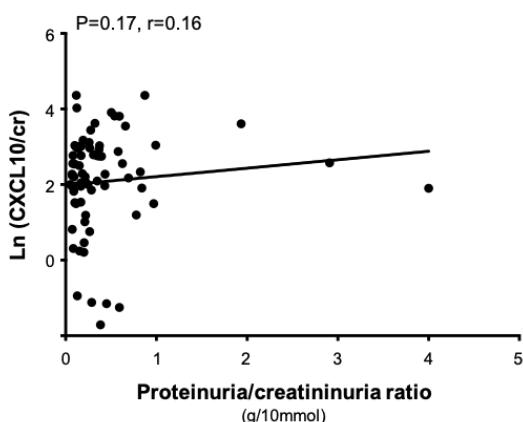
Global population of patients with viremia: N=76 samples of whom 68 have concurrent viruria, see Fig 2A)

B



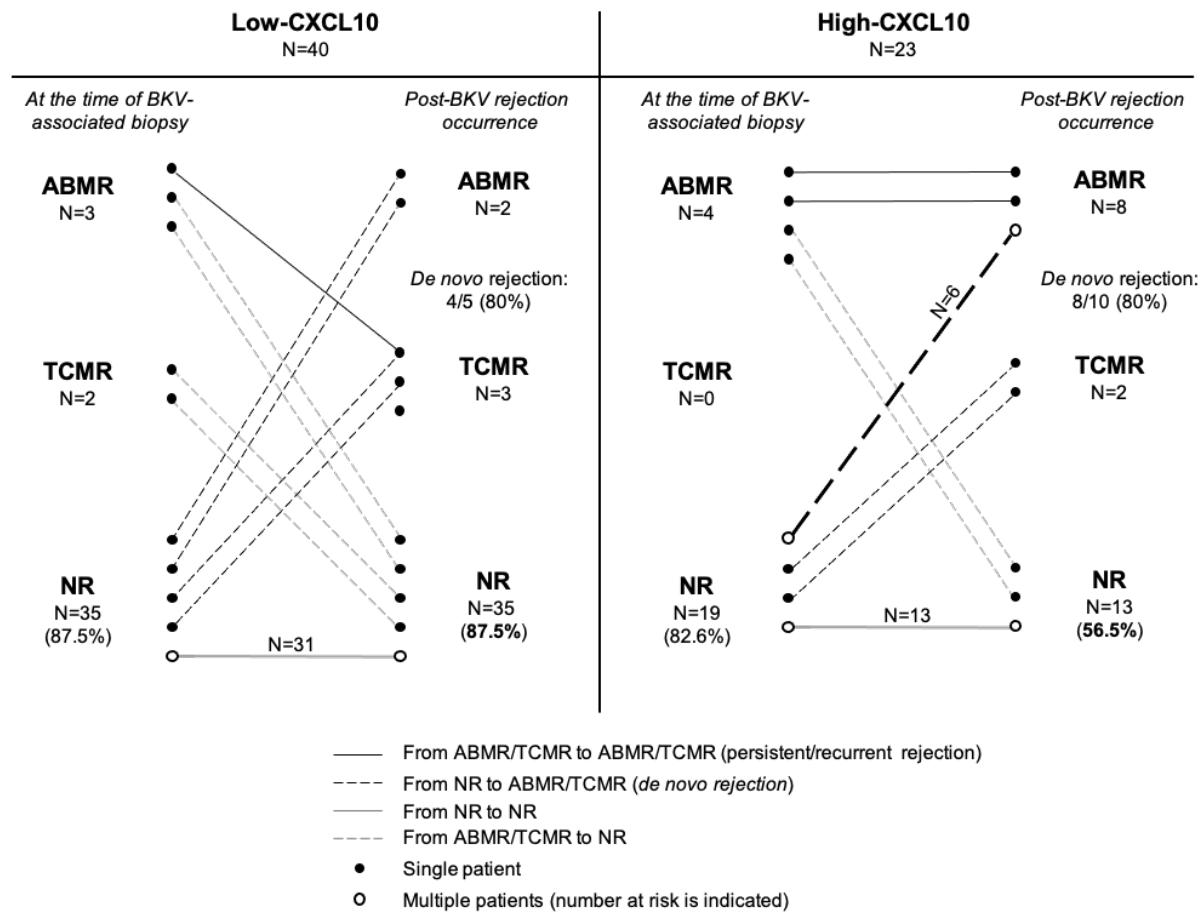
Restricted population of patients with viruria: N=158 samples (after exclusion of N=13 UTI cases and/or N=35 acute rejection cases)

C



Global population of patients with viremia: N=76 samples of whom proteinuria was available for 74 samples

Supplemental Figure 2: Relationship between the rejection phenotypes observed at the time of index biopsy for concurrent BKV-DNAemia and the post-BKV occurrence of rejection in the low-CXCL10 (≤ 12.86 ng/mmol) and high-CXCL10 (> 12.86 ng/mmol) groups from the nested case-control study. Dashed lines indicate a change in category from no rejection to rejection (*i.e.*, *de novo* rejection, in black) or the opposite from rejection to no rejection (in gray). Solid lines indicate patients staying within the same category: with a rejection diagnosis at the time of index biopsy and a post-BKV persistent/recurrent rejection (in black) or no rejection at any time (in gray).



Supplemental Figure 3: Urinary CXCL10/cr (**A**) and blood BKV viral load (**B**) trajectory analyses in the longitudinal cohort split in three groups according to AST Guidelines (Hirsch *et al*, Clin Transplant 2019). Transient BKV-DNAemia is defined by spontaneous BKV clearance without lowering of maintenance immunosuppression (N=18, light gray line). Sustained low-level DNAemia is defined by peak viremia <4 log₁₀c/mL (N=27 KTRs, dark gray line). Sustained high-level DNAemia is defined by peak viremia ≥4 log₁₀c/mL (N=13 KTRs, black line). Trajectories were computed by regression from longitudinal assessments of uCXCL10/cr (samples collected at biopsy and each outpatient clinic visit during the 1st year post-transplantation) and all available blood BKV viral loads over the same period.

