

# Origine de la réactivation virale chez les patients transplantés rénaux avec une néphropathie à BK virus

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INSERM U944 « Cellular biology of viral infections », Institut de recherche Saint Louis, Paris



# JNI

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au mercredi 1<sup>er</sup> septembre 2021



## Déclaration d'intérêts de 2014 à 2020

- Aucun

## Déclaration de liens d'intérêt avec les industries de santé en rapport avec le thème de la présentation (loi du 04/03/2002) :

**Intervenant :** Gras Julien

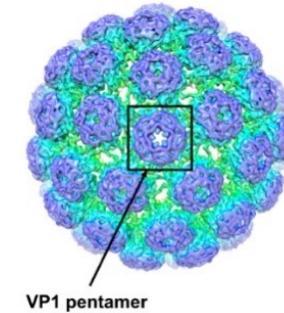
**Titre :** Origine de la réactivation virale chez les patients transplantés rénaux avec une néphropathie à BK virus

L'orateur ne souhaite pas répondre

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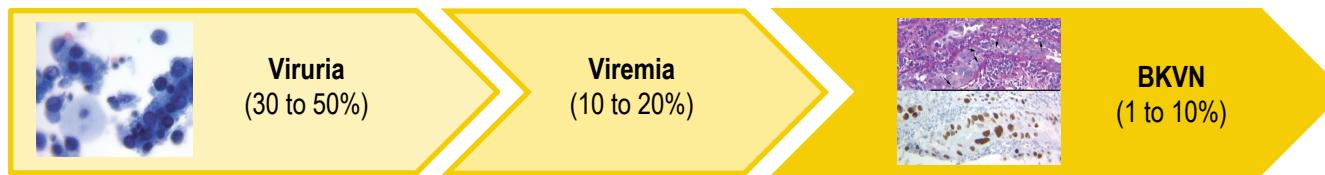
# BK virus: an opportunistic pathogen

- A member of the *Polyomaviridae* family
  - ds-DNA virus (5.2kb genome)
  - Icosahedral capsid (40-45nm)
- Natural acquisition in childhood
  - Persistent infection (renourinary tract)
  - Intermittent reactivation  
(asymptomatic viruria → hemorrhagic cystitis / interstitial nephropathy)
- BKV genetic diversity > Sequence variation of VP1-gene
  - 4 main genotypes (distinct serotypes)
  - BKV-I largely predominant > BKV-IV (>> BKV-II / BKV-III)
  - *In vivo* coexistence of different genotypes



# BK virus associated nephropathy (BKVN)

- A major complication following kidney transplantation (KT)
  - 1 to 10% of KT recipients [KTR] : poor renal prognosis (30 to 65% of graft loss)
  - No specific antiviral drug available
- Natural history of BKV infection in KTR



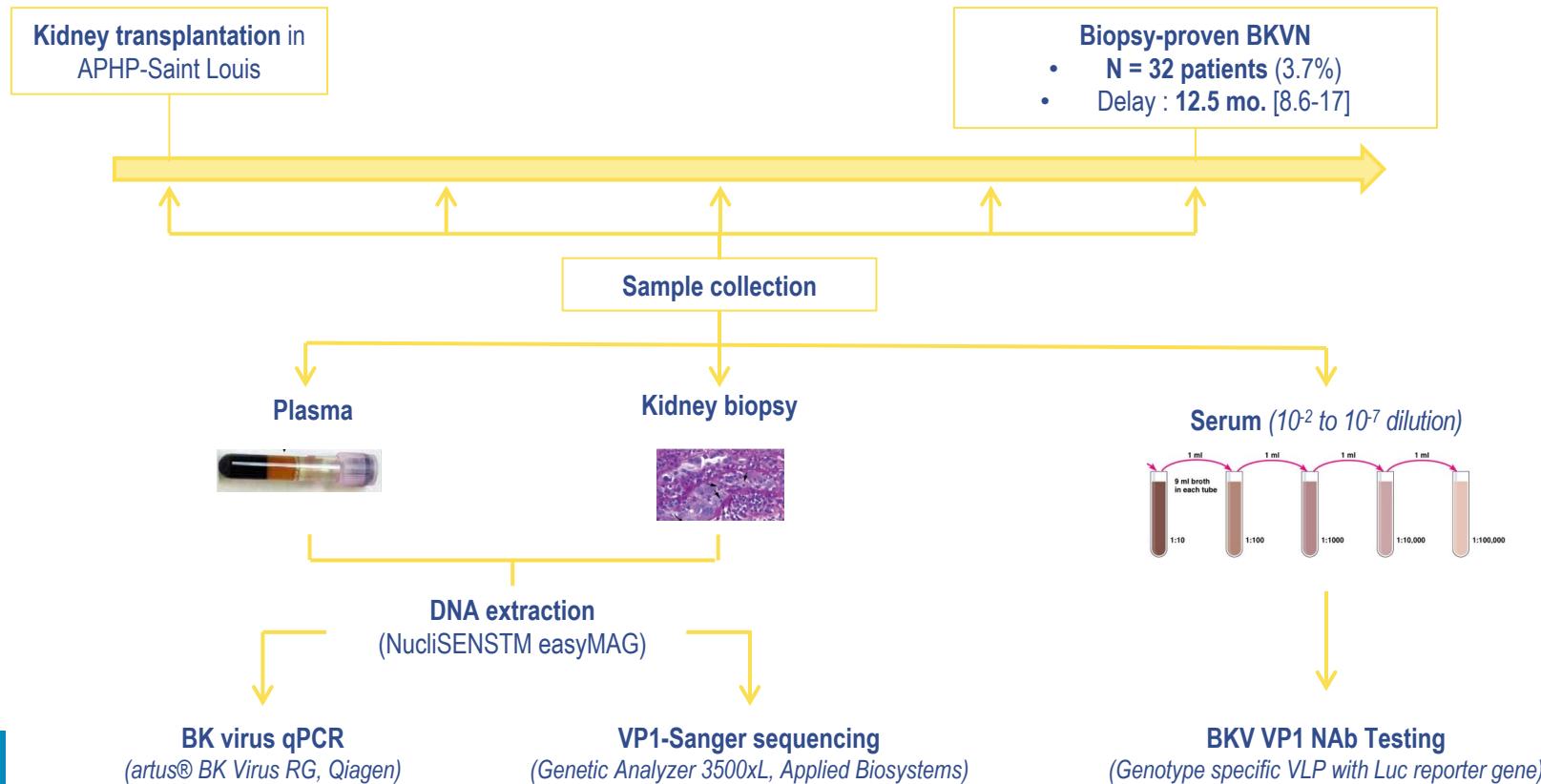
- Physiopathology of BKVN : several unanswered questions
  - Viral origin of the BKV strain responsible for BKVN ?
  - Impact of *in vivo* BKV genetic diversity ?
  - Mechanisms of BKV immune escape ?

# Objectives

**Kidney transplant recipients with biopsy-proven  
BKV associated nephropathy**

1. Analyze the kinetics of BKV replication following kidney transplantation in plasma & kidney
2. Study BKV genetic diversity in different biological compartments using VP1 Sanger sequencing
3. Quantify the anti-VP1 humoral response associated with BKV infection using neutralizing antibody testing

# Retrospective study (01/2016 – 06/2020)



# BKV DNA detection in plasma & kidney biopsy

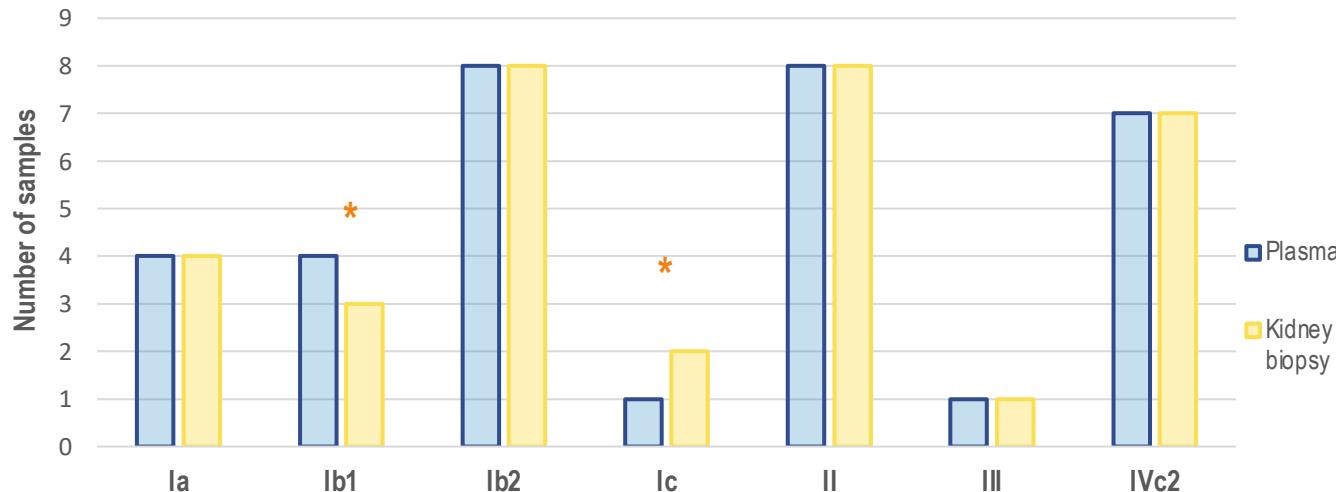
	Day 0 - KT (N=20)	M3 (N=23)	BKVN diagnosis (N=32)
<b>BKV DNA in kidney biopsy</b>			
BKV+ samples (N, %)	8 (40)	12 (52) *	32 (100)
Median VL [IQR] in BKV+ (logIU/ $10^6$ cells)	3.7 [3.5-4.4]	5.7 [3.7-6.4]	8.2 [5.8-8.9]
<b>BKV DNA in concomitant plasma</b>			
BKV+ samples (%)	0 (0)	7 (30)	32 (100)
Median VL [IQR] in BKV+ (logIU/ml)	-	3.5 [3.7-6.4]	5.4 [4.3-6.0]

\* 8/12 of KTR with BKV+ biopsy at month 3 had a negative concomitant plasma

→ BK virus DNA is detectable in kidney from KTR with BKVN before the onset of viremia

# *In vivo* BKV genetic diversity (1)

- BKV genotyping at BKVN diagnosis

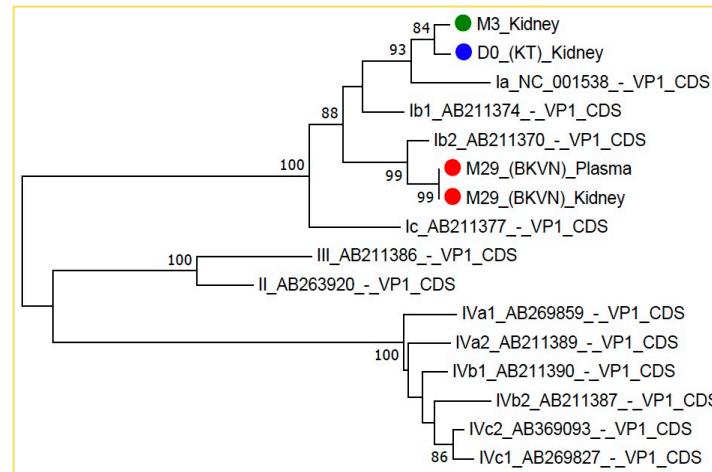
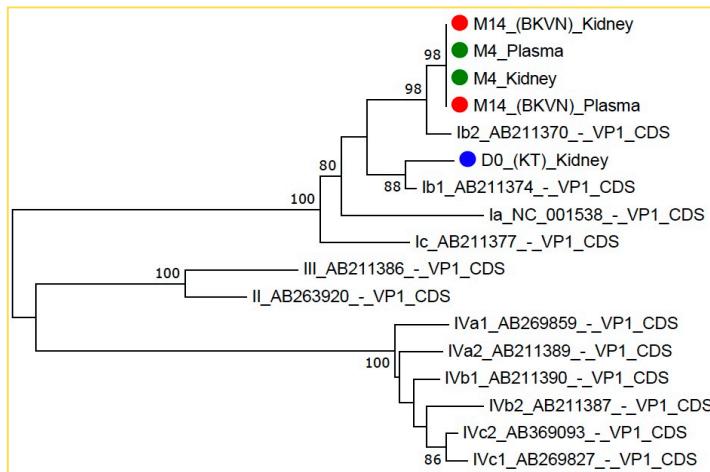


\* In one patient, VP1 Sanger sequencing identified a different BKV-subtype at the time of BKVN diagnosis in kidney biopsy (BKV-Ic) and concomitant plasma (BKV-Ib1)

→ The BKV circulating strain is the one responsible of BKVN at diagnosis

# *In vivo* BKV genetic diversity (2)

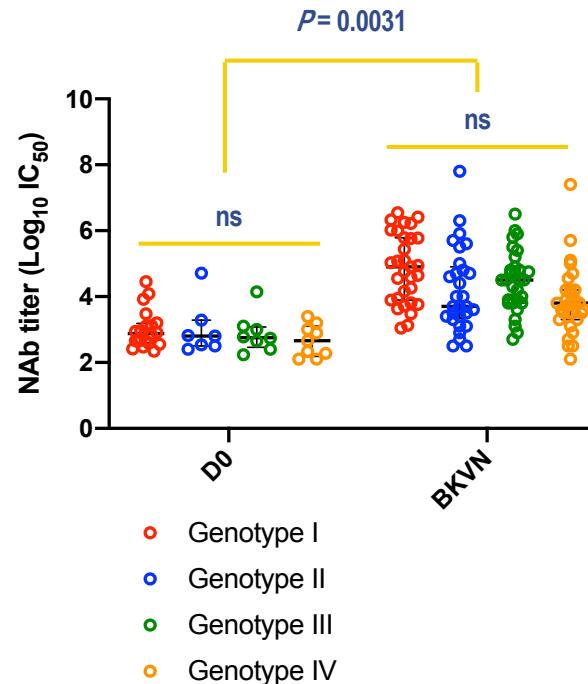
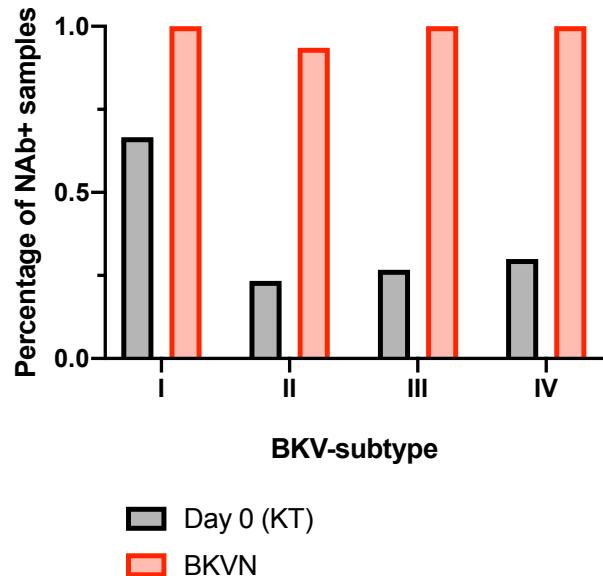
- **Plasma:** Identical BKV-subtype at first viremia & BKVN (N=19/19, 100%)
- **Kidney biopsy:** Different BKV subtype isolated in KB at day 0 ≠ BKVN (N=5/6, 83%)



→ BKVN results at least in part from the **reactivation of the recipient BKV subtype**

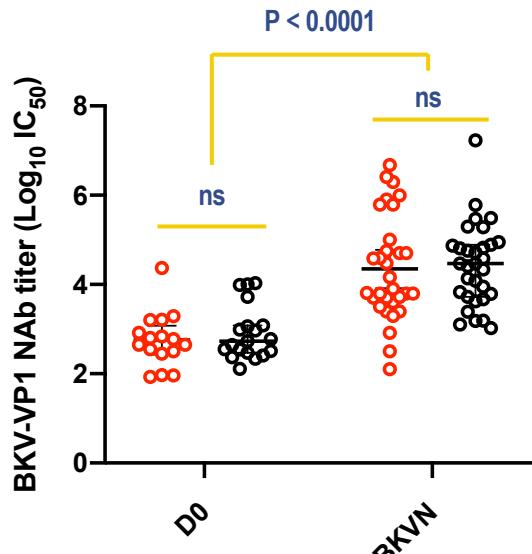
# BKV-VP1 Nab response in KTR with BKVN (1)

- BKV VP1 NAb titers are significantly increased at BKVN diagnosis compared to day 0

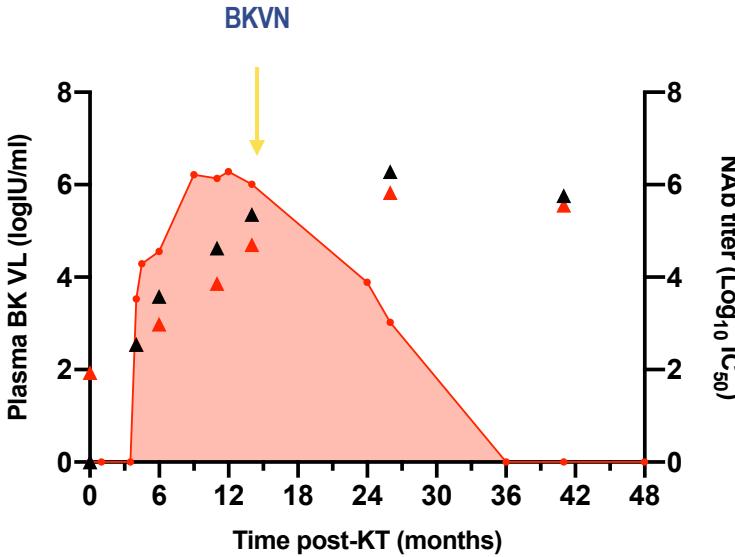


# BKV-VP1 NAb response in KTR with BKVN (2)

- BKV-VP1 NAb response against BKVN strain does not differ from other subtypes



- BKVN-subtype
- Non-BKVN subtype



- Plasma BK VL ( $\log_{10} \text{IU/ml}$ )
- ▲ BKVN subtype NAb titer (IgG)
- ▲ D0 subtype NAb titer (IgG)

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# Conclusion and perspectives

- 1. BK virus is detectable early in kidney of KTR with BKVN**
  - Early marker to better predict the risk of BKVN in KTR ?
  - Comparative analysis with KTR without BKV infection
- 2. BKVN results at least in part from the reactivation of the recipient's strain**
  - Concomitant infection with the donor's virus ?
  - Whole genome sequencing ongoing
- 3. BKV VP1 NAb response is non specific in KTR with BKVN diagnosis**
  - Humoral escape from specific VP1 variants ?
  - Role of impaired cellular immunity

# Acknowledgements



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