1	Novel human polyomaviruses in pregnancy: higher prevalence of BKPyV, but no WUPyV,
2	KIPyV and HPyV9
3	
4	Eszter Csoma ^{a, *} , Tamás Sápy ^b , Beáta Mészáros ^a , Lajos Gergely ^a
5	
6	^a Department of Medical Microbiology, Medical and Health Science Centre, University of
7	Debrecen, Nagyerdei krt. 98., H-4032 Debrecen, Hungary
8	^b Department of Obstetrics and Gynecology, Medical and Health Science Centre, University
9	of Debrecen, Nagyerdei krt. 98., H-4032 Debrecen, Hungary
10	
11	* Corresponding author. Tel.: +36 52 255 425; fax: +36 52 255 424. Email address:
12	csomae@freemail.hu.
13	
14	Abbreviations:
15	WU polyomavirus (WUPyV), KI polyomavirus (KIPyV), human polyomavirus 9 (HPyV9),
16	BK polyomavirus (BKPyV), genome equivalent (GEq), polymerase chain reaction (PCR)
17	
18	
19	
20	
21	
22	Number of words in the Abstract: 248
23	Number of words in the text: 1879
24	
25	

26 Abstract

Background: Immunosuppression due to pregnancy may lead to higher susceptibility to
infections and reactivation of latent infections, such as BK polyomavirus (BKPyV). There is
lack of information about the prevalence of novel human polyomavirus 9 (HPyV9), WU
(WUPyV) and KI (KIPyV) during pregnancy.

Objectives: To study whether pregnancy results in higher prevalence of HPyV9, WUPyV,
KIPyV and their correlation with BKPyV.

Study design: Plasma, urine and throat swab samples from 100 pregnant and 100 non
pregnant women were screened for the presence of WUPyV, KIPyV, HPyV9 and BKPyV by
PCR.

36 Results: No WUPyV DNA was detected in plasma, urine and respiratory samples from pregnant and non pregnant women. KIPyV DNA was found in two plasma samples from non 37 38 pregnant women (2 %) and not detected in other samples from neither pregnant nor non 39 pregnant women. HPyV9 DNA was determined in all sample types of pregnant and non 40 pregnant women, respectively. There were no significant differences between pregnant and 41 non pregnant women in HPyV9 DNA frequencies for plasma (2 % vs. 6 %), urine (3 % vs. 2 42 %) and respiratory samples (2 % vs. 2 %). Prevalence of BKPyV in urine samples was 43 significantly higher (p=0.039) in pregnant women (13 %) then in non pregnant women (4 %); 44 co nfection with KIPyV and/or HPyV9 was not detected.

45 Conclusions: In contrast with BKPyV, infection with WUPyV, KIPyV and HPyV9 was not
46 detected more frequently during pregnancy. To our knowledge HPyV9 was detected first in
47 respiratory samples in our study.

48

49 Key words: human polyomaviruses, pregnancy

51 **1. Background**

Human polyomavirus BK (BKPyV) seroprevalence increases with age reaching high, 52 80-90 % in adult population.¹ Similarly high, 55-90 % adult seropositivities were observed for 53 recently discovered KI² and WU³ polyomaviruses (KIPvV, WUPvV).⁴⁻⁶ Investigation of 54 seropositivity against the newly discovered human polyomavirus 9 (HPyV9)⁷ revealed 47 % 55 56 positivity for healthy adults.⁸ It is well known that after the childhood primary infection with 57 BKPyV, lifelong persistent infection is established mainly in renal and urinary tract cells.¹ 58 Transient immunosuppression due to pregnancy may lead to reactivation of BKPyV resulting in generally asymptomatic viruria with frequency of 3 to 54 %.^{1, 9-11} Beside viruria, BKPyV 59 viraemia was also detected in pregnant women.¹¹ The pathogenic role of the novel WUPyV, 60 KIPyV and HPyV9 is far from clear, only speculative. WU and KI viruses were found in 61 various sample types - respiratory samples, blood, faeces, cerebrospinal fluid, lymphoid 62 63 tissues, urine - and higher prevalence was observed in children and immunocompromised patients.¹²⁻¹⁶ HPyV9 was described from blood and urine samples of kidney transplant 64 patients, then it was found in skin samples, but no in respiratory and fecal samples.^{7, 17} The 65 66 higher frequency of these viruses in immunocompromised patients suggests higher 67 susceptibility or reactivation due to immunosuppression. Up to now only four urine samples from pregnant women were investigated for the presence of KIPyV and WUPyV DNA with 68 negative result.¹⁸ The genetic and possible transmission similarities to BKPvV, and the higher 69 70 PCR prevalence data among immunocopromised patients may suggest that 71 immunosuppression, thus pregnancy may lead to higher susceptibility to infection with WUPyV, KIPyV and HPyV9 or may result in reactivation of possible latent infections. 72

73

74 **2. Objective**

The aim of the present study was to evaluate the prevalence of three new human polyomaviruses (WUPyV, KIPyV and HPyV9) during pregnancy, to study whether immunosuppression due to pregnancy may lead to higher prevalence as it was found in case of BKPyV. The possible correlations of these viruses were also investigated.

79

80 **3. Study design**

81 *3.1. Patients and samples*

Urine, plasma (from EDTA blood samples) and throat swab samples were collected on the same day from 100 healthy pregnant women (age 16.5-41.9 years, median 32.1 years; pregnancy 5-39 weeks; median 26 weeks) and 100 non pregnant women (age 18-44.3 years, median 31.6 years) between September 2011 and December 2011. Samples from pregnant women were collected in all three trimesters: first trimester n=28; second trimester n=27; third trimester n=45. The control samples were taken from healthy, non pregnant, fertility exam visitor women.

Immediately after collection, nucleic acid was isolated from samples using High Pure
Viral Nucleic Acid Kit (Roche, Switzerland) according to the manufacturer's instructions.
Briefly, nucleic acids from 200 µl plasma, 200 µl urine specimen and throat swab sample
washed in 200 µl buffer were eluted in 50 µl and stored at -20 °C until use.

93 The study was approved by Regional and Institutional Ethics Committee of94 University of Debrecen. All patients were asked to sign written informed consent.

95

96 3.2. Nested and real-time PCR for WUPyV, KIPyV, HPyV9 and BKPyV

All PCR methods were carried out with 10 µl nucleic acid in a final volume of 25 µl.
For nested PCR AmpliTaq Gold 360 Master Mix, for WUPyV and KIPyV real-time PCR
TaqMan Universal PCR Master Mix (Applied Biosystems, USA) were used. The calibrants

100 for quantitative PCRs were serial dilutions of KIPyV plasmid (in which the genome of KI 101 polyomavirus isolate Stockholm 60 was incorporated) and AP-p003 plasmid (containing the 102 2228 bp half genome of WU polyomavirus) kindly provided by Tobias Allander and David 103 Wang. WUKI nested PCR and real-time PCR for WU and KI virus were performed as described previously.¹⁶ HPyV9 PCR was carried out with diagnostic primers and annealing 104 temperature published by Scuda et al.⁷ For the first round of BKV nested PCR, k1 (5' 105 106 TGAAGCATATGAAGATGGCC 3') and k2 (5' GTTACAGCCTCCCACATC 3') primers 107 were used with 60 °C annealing temperature, while for the second round b1 (5' 108 GATGGCCCCAACCAAAAG 3') and b2 (5' CTAGAACTTCTACTCCTCC 3') primers and 109 56 °C annealing temperature were applied. PCR products were visualized by electrophoresis 110 in 1.5 % agarose gel containing ethidium bromide (0.5 μ g/mL). The amplified PCR products 111 from WUKI and HPyV9 nested PCR were cut, purified with QIAquick Gel Extraction Kit 112 (Qiagen) according to the instructions and sequenced by using ABI PRISM 3100 Genetic 113 Analyzer (Applied Biosystems). To determine BKPyV viral load BKV virus R-gene 114 quantification kit was used (Argene, USA) according to the manufacturer's instructions.

115

116 3.3. Statistical analysis

Difference in frequency for categorical variables was analysed by Fisher's exact test.
For continuous variables Mann-Whitney U test was applied. Difference was considered
significant if p value was less then 0.05.

120

121 **4. Results**

4.1. Detection of WUPyV, KIPyV and HPyV9 DNA in plasma, urine and respiratory samples
Table 1 shows the results of PCR detections for the various samples. WUPyV DNA
was not detected in plasma, urine and respiratory samples neither from pregnant nor from non

125 pregnant women. KIPyV was found in two plasma samples of non pregnant women, but was 126 not determined in any other samples. To confirm the positive PCR results and to determine KI 127 or WU virus DNA was detected, PCR products were sequenced. The viral loads were below 128 the limit of detection (< 250 GEq/mL; genome equivalent/mL) by real-time PCR. HPyV9 129 DNA was detected in urine, plasma and respiratory samples from both studied groups. To 130 prove the results from PCR, all PCR products were sequenced. In details, the prevalence of 131 HPvV9 DNA in plasma samples was higher in control, non pregnant group then in pregnant 132 women (6/100; 6 % vs. 2/100; 2%), but the difference was not statistically significant. The 133 two positive samples were taken in the second trimester of pregnancy. Two samples from 134 control, non pregnant women with HPyV9 viraemia were also positive for KIPyV DNA. In 135 respiratory samples the frequency of HPyV9 DNA was the same in both studied groups 136 (2/100; 2% and 2/100; 2%). Both of the positive samples in pregnant women group were collected in the first trimester. Three urine samples from pregnant women were HPyV9 PCR 137 138 positive (3/100; 3%), while in control group 2 samples were positive (2/100; 2%) which is not 139 statistically significant difference.

140 4.2. Prevalence of BKPyV in urine and plasma samples

BKPyV was not detected in plasma samples. Frequency for BKPyV viruria was 13 % (13/100) in pregnant women and 4 % (4/100) in non pregnant, control group (Table 1.). The difference is statistically significant (p=0.039). The BKPyV viral load in samples from pregnant women (range 50-1.86 x 10^8 ; median 11.82 x 10^3 GEq/mL) did not show statistically significant difference from the viral load in control samples (range 2.25 x 10^2 -3.58 x 10^5 ; median 2.98 x 10^2). BKPyV presence in urine samples was found in all trimesters.

147

148 **5. Discussion**

In our study significantly higher prevalence of BK viruria was observed in pregnant women in contrast with non pregnant women. Human polyomavirus 9 was found in plasma, urine and respiratory samples from pregnant women but not more frequently then in samples from non pregnant women. WU and KI viruses were not detected in any of the studied samples from pregnant women.

154 BK polyomavirus is ubiquitous in the human population, the primary infection 155 generally occurs during childhood without significant clinical consequences, respiratory 156 diseases might occur. Transmission of the viruses is not well clarified, but it is suggested that 157 these viruses are acquired mainly through respiratory, faecal-oral and urinary routes, alternatively by blood transfusion and organ transplantation.¹ After the primary infection, 158 lifelong persistence of the virus is established mainly in kidney and urinary tract.^{19, 20} Lytic 159 infection with viruria occurs in 5-10 % of immunocompetent individuals²¹, but more 160 frequently in immunocompromised patients.²² During pregnancy immunologic changes 161 162 together with hormonal effects may result in viral infections, reactivations. Viruria was detected for 3-54 % of pregnant women, while viraemia was found to be less frequent.^{1, 10, 11,} 163 ²³ In accordance with literature, in this study 13 % of pregnant women had active BKPyV 164 165 replication resulting in viruria, but no viraemia. The possible effect of BK virus replication during pregnancy is not clarified. Although viral DNA was demonstrated in fetal tissues ²⁴ the 166 hypothesis of transplacental transmission was not confirmed ^{10, 11}. Recently serological 167 evidence for vertical transmission of BKPyV was published.⁹ 168

169 Hitherto, there are no prevalence data about the novel WU, KI and human 170 polyomavirus 9 during pregnancy. Bofill-Mas et al. investigated 4 urine samples from 171 pregnant women, but WU and KI viruses were not found. ¹⁸ Foetal tissues were also negative 172 for WU and KI viruses. ²⁵ In this study WU and KI viruses were not found in urine, plasma 173 and respiratory samples collected during pregnancy. KIPyV DNA was detected in two plasma

174 samples, but not in urine and respiratory samples from control, non pregnant women. The 175 high, 55-90 % seropositivity in adult population, and the higher PCR prevalence in samples from children suggest childhood primary KI and WU virus infection.^{6, 15} Viruses were found 176 177 with frequency 0.4-14 % in various samples types including respiratory samples, blood, 178 faeces, cerebrospinal fluid, lymphoid tissues and urine samples, with generally higher 179 frequency in immunocompromised patients. The possible way of transmission might be faecal-oral.^{2, 3, 12-16} The higher PCR prevalence data of 180 respiratory and/or 181 immunocompromised patients suggests that immunosuppression might result in reactivation of these viruses, or might establish higher susceptibility to KIPyV and WUPyV infection.¹ It 182 183 was hypothesized that similarly to BKPyV, transient immunosuppression due to pregnancy 184 might result in higher frequency of WU and/or KI viral infections, but no evidence for it was 185 found during this study. However it is important to note, that it was not a follow up study, 186 samples were collected once randomly during pregnancy.

Human polyomavirus 9 was described in 2011.⁷ Up to now, viral DNA was found in 187 188 blood and urine samples from immunocompromised patients and skin samples, but neither in 189 respiratory samples from patients with respiratory failure nor in faeces from children with gastroenteritis.^{7, 17} Based on these data and the recently published 47 % adulthood 190 seropositivity⁸, Van Ghelue et al. hypothesized that HPyV9 is less frequent in the human 191 192 population⁶. We found HPyV9 DNA is all studied samples from pregnant and non pregnant 193 women with frequency of 2-6 %. There was no or not statistically significant difference 194 between the PCR prevalence in the respiratory (2 vs. 2 %), urine (3 vs. 2%) and plasma 195 samples (2 vs. 6 %) between pregnant and non pregnant women. To our knowledge we 196 published first HPyV9 presence in respiratory samples which may suggest respiratory 197 transmission of this virus. In this study higher prevalence of HPyV9 was not found during 198 pregnancy, but the viral loads were not examined which might have been different. Since

mother to foetus transmission of polyomaviruses BK and JC are suggested, this way of transmission cannot excluded in case of the novel WUPyV, KIPyV and HPyV9. Even if this study could not support evidence for higher susceptibility of infection by these viruses, further, follow up study of pregnant women during the whole period of pregnancy might answer this question.

In conclusion, KI and WU viruses were not found in urine, respiratory and blood samples from pregnant women, while HPyV9 was detected in all sample types but with no significantly higher frequency then it was observed for non pregnant women.

207

208 **Conflict of interest**

209 The authors have no conflict of interest.

210

211 Acknowledgements

We thank Tobias Allander from Karolinska Institute (Sweden), David Wang from Washington University (USA) and Bernhard Ehlers from Robert Koch-Institut (Germany) for providing KIPyV, WUPyV and HPyV9 plasmids, respectively. This work was supported by grants from the Hungarian Scientific Research Found (OTKA 73145) and by the TÁMOP-4.2.2/B-10/1-2010-0024 project which is co-financed by the European Union and the European Social Fund.

218

219 **References**

- Jiang M, Abend JR, Johnson SF,Imperiale MJ. The role of polyomaviruses in human
 disease. *Virology* 2009; **384**(2):266-73.
- Allander T, Andreasson K, Gupta S, Bjerkner A, Bogdanovic G, Persson MA, et al.
 Identification of a third human polyomavirus. *J Virol* 2007; **81**(8):4130-6.

- 3. Gaynor AM, Nissen MD, Whiley DM, Mackay IM, Lambert SB, Wu G, et al.
 Identification of a novel polyomavirus from patients with acute respiratory tract
 infections. *PLoS Pathog* 2007; **3**(5):e64.
- 4. Neske F, Prifert C, Scheiner B, Ewald M, Schubert J, Opitz A, et al. High prevalence
 of antibodies against polyomavirus WU, polyomavirus KI, and human bocavirus in
 German blood donors. *BMC Infect Dis* 2010; **10**(215.
- S. Nguyen NL, Le BM, Wang D. Serologic evidence of frequent human infection with
 WU and KI polyomaviruses. *Emerg Infect Dis* 2009; **15**(8):1199-205.
- 232 6. Van Ghelue M, Khan MT, Ehlers B, Moens U. Genome analysis of the new human
 233 polyomaviruses. *Rev Med Virol* 2012;
- Scuda N, Hofmann J, Calvignac-Spencer S, Ruprecht K, Liman P, Kuhn J, et al. A
 novel human polyomavirus closely related to the african green monkey-derived
 lymphotropic polyomavirus. *J Virol* 2011; **85**(9):4586-90.
- 8. Trusch F, Klein M, Finsterbusch T, Kuhn J, Hofmann J,Ehlers B. Seroprevalence of
 the human polyomavirus 9 (HPyV9) and cross-reactivity to the African green
 monkey-derived lymphotropic polyomavirus (LPV). *J Gen Virol* 2012;
- 9. Boldorini R, Allegrini S, Miglio U, Paganotti A, Cocca N, Zaffaroni M, et al.
 Serological evidence of vertical transmission of JC and BK polyomaviruses in
 humans. *J Gen Virol* 2011; **92**(Pt 5):1044-50.
- 10. Kalvatchev Z, Slavov S, Shtereva M,Savova S. Reactivation of Polyomavirus hominis
 1 (BKV) during pregnancy and the risk of mother-to-child transmission. *J Clin Virol*2008; 43(3):328-9.
- Boldorini R, Veggiani C, Amoruso E, Allegrini S, Miglio U, Paganotti A, et al. Latent
 human polyomavirus infection in pregnancy: investigation of possible transplacental
 transmission. *Pathology* 2008; **40**(1):72-7.

- Dalianis T, Ramqvist T, Andreasson K, Kean JM,Garcea RL. KI, WU and Merkel cell
 polyomaviruses: a new era for human polyomavirus research. *Semin Cancer Biol*2009; 19(4):270-5.
- 13. Bialasiewicz S, Whiley DM, Lambert SB, Nissen MD, Sloots TP. Detection of BK, JC,
- WU, or KI polyomaviruses in faecal, urine, blood, cerebrospinal fluid and respiratory
 samples. *J Clin Virol* 2009; **45**(3):249-54.
- Mourez T, Bergeron A, Ribaud P, Scieux C, de Latour RP, Tazi A, et al.
 Polyomaviruses KI and WU in immunocompromised patients with respiratory disease. *Emerg Infect Dis* 2009; **15**(1):107-9.
- Babakir-Mina M, Ciccozzi M, Perno CF, Ciotti M. The novel KI, WU, MC
 polyomaviruses: possible human pathogens? *New Microbiol* 2011; **34**(1):1-8.
- 260 16. Csoma E, Meszaros B, Asztalos L, Konya J,Gergely L. Prevalence of WU and KI
 261 polyomaviruses in plasma, urine, and respiratory samples from renal transplant
 262 patients. *J Med Virol* 2011; 83(7):1275-8.
- 263 17. Sauvage V, Foulongne V, Cheval J, Ar Gouilh M, Pariente K, Dereure O, et al.
 264 Human polyomavirus related to African green monkey lymphotropic polyomavirus.
 265 *Emerg Infect Dis* 2011; **17**(8):1364-70.
- Bofill-Mas S, Rodriguez-Manzano J, Calgua B, Carratala A,Girones R. Newly
 described human polyomaviruses Merkel cell, KI and WU are present in urban sewage
 and may represent potential environmental contaminants. *Virol J* 2010; 7(141.
- Heritage J, Chesters PM,McCance DJ. The persistence of papovavirus BK DNA
 sequences in normal human renal tissue. *J Med Virol* 1981; 8(2):143-50.
- 271 20. Chesters PM, Heritage J,McCance DJ. Persistence of DNA sequences of BK virus and
 272 JC virus in normal human tissues and in diseased tissues. *J Infect Dis* 1983;
 273 147(4):676-84.

- 274 21. Egli A, Infanti L, Dumoulin A, Buser A, Samaridis J, Stebler C, et al. Prevalence of
 275 polyomavirus BK and JC infection and replication in 400 healthy blood donors. J
 276 Infect Dis 2009; 199(6):837-46.
- 277 22. Dharnidharka VR, Abdulnour HA,Araya CE. The BK virus in renal transplant
 278 recipients-review of pathogenesis, diagnosis, and treatment. *Pediatr Nephrol*279 26(10):1763-74.
- 280 23. Boothpur R,Brennan DC. Human polyoma viruses and disease with emphasis on
 281 clinical BK and JC. *J Clin Virol* 2010; 47(4):306-12.
- 282 24. Pietropaolo V, Di Taranto C, Degener AM, Jin L, Sinibaldi L, Baiocchini A, et al.
- 283 Transplacental transmission of human polyomavirus BK. J Med Virol 1998;
 284 56(4):372-6.
- 285 25. Sadeghi M, Riipinen A, Vaisanen E, Chen T, Kantola K, Surcel HM, et al. Newly
 286 discovered KI, WU, and Merkel cell polyomaviruses: no evidence of mother-to-fetus
 287 transmission. *Virol J* 2010; 7(251.
- 288
- 289