Low prevalence of Merkel cell polyomavirus in human epithelial thymic tumors

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Abstract

Background: The etiology of thymic epithelial tumors is unknown. Murine polyomavirus strain PTA has been shown to induce thymomas in mice. Recently, using diverse molecular techniques, we reported the presence of human polyomavirus 7 (HPyV7) in thymic epithelial tumors. In the present study, we investigated the prevalence of Merkel cell polyomavirus (MCPyV) in thymic epithelial tumors.

Methods: Thirty-six thymomas were screened for MCPyV by PCR and subsequently tested by DNA and RNA in situ hybridization and immunohistochemistry. Twenty-six thymomas were diagnosed with myasthenia gravis (MG).

Results: MCPyV DNA was detected by PCR in 7 (19.4%) of the 36 thymic epithelial tumors and in six of these, the presence of MCPyV was confirmed by fluorescence in situ hybridization. Of these, 3 (28.6%) revealed weak MCPyV LT-antigen protein expression. In addition, one of the MCPyV positive thymomas tested positive for MCPyV LT RNA with RNAscope. Of interest, two out of the three thymomas that previously tested positive for MCPyV by immunohistochemistry also tested positive for HPyV7. One of the 11 MG-negative and 2 of the 25 MG-positive were positive for MCPyV.

Conclusions: MCPyV DNA and MCPyV protein expression can be detected in human epithelial thymoma; however, to a far lesser extent than HPyV7. Our data strongly indicate that because of its infrequent detection and weak expression, MCPyV is unlikely to play an important role in the etiopathogenesis of human thymomas.

Introduction

Thymomas are rare thymic epithelial tumors that are frequently associated with autoimmune diseases, such as myasthenia gravis (MG, 24–40%). The underlying etiology of thymomas is unknown. Several studies have investigated the role of viruses in thymomagenesis.1–3 We previously assessed the presence of the novel human polyomaviruses 6 and 7 (HPyV6 and 7) in human thymic epithelial tumors, based on the reported induction of thymomas in mouse strains C3H/BiDa and AKR by the polyomavirus strain PTA, and frequently detected HPyV7 in these tumors.1–6 The number of human polyomaviruses detected has continually increased in recent years.7,8 To date, only Merkel cell polyomavirus (MCPyV) has been established as
a novel human tumor virus in the majority of Merkel cell carcinomas (MCC). In MCC, MCPyV is clonally integrated in the tumor DNA and harbors tumor specific truncating oncogenic mutations within the large T (LT) antigen. Truncated LT antigen and small T antigen (sTAg) of MCPyV have been shown to inhibit the tumor suppressor protein retinoblastoma (RB) and p53. (sTAg) of MCPyV have been shown to inhibit the tumor (LT) antigen. In other studies, 86.4% and 56.8–63.6% of adult blood sera (≥ 20 years) were positive for MCPyV- and HPyV7-DNA, respectively. In two independent studies using clinicopathologically not further classified thymomas (n = 10), no MCPyV DNA could be detected by PCR. Because of these reports and our recent finding that 62.2% of thymomas are HPyV7 positive, we aimed to comprehensively assess the prevalence of MCPyV in this clinicopathologically well-classified cohort of human thymic epithelial tumors by diverse molecular techniques.

Testing the prevalence of MCPyV in formalin-fixed and paraffin-embedded (FFPE) tissues of MCC has generated conflicting results in previous studies. According to Moshiri et al., the use of a PCR-based approach in combination with the CM2B4 anti LT-antigen antibody has the highest specificity and sensitivity. Herein, we used a PCR based approach in combination with MCPyV fluorescence in situ hybridization (FISH) and immunohistochemistry (IHC) to elucidate the presence of MCPyV in thymic epithelial tumors.

**Methods**

**Patients and tissues**

Thirty-six FFPE thymoma resection specimens were included in this study (from 19 women and 17 men; mean age 58.3 years, range 34–82 years). Thirty-five of these had previously been tested for the presence of HPyV7. Twenty-six thymoma patients were known to have a history of MG, of which 24 were anti-acetylcholine receptor (AChR) antibody positive and two were negative. Twelve of the 26 MG-positive thymoma patients were immunosuppressed. All clinical pathological data are summarized in Table 1. In addition, non-neoplastic thymic hyperplasia (n = 20) and fetal thymic (n = 20) tissues were included in this study (summarized in Table S1). All use of this tissue and patient data was in agreement with the Dutch Code of Conduct for Observational Research with Personal Data (2004) and Tissue (2011, www.federa.org/sites/default/files/digital_version_first_part_code_of_conduct_in_uk_2011_12092012.pdf).

Diagnoses were previously defined by histology in routine diagnostics and were reviewed by two experienced pathologists.

**Merkel cell polyomavirus (MCPyV) detection**

PCR was performed with 250 ng of genomic DNA using the AmpliTaq Gold DNA Polymerase (Thermo Fisher Scientific, Naarden, The Netherlands) in a final volume of 50 μL, as recently described. DNA quality and integrity were assessed by specimen control size ladder (SCS) as previously described and published for the DNA used. All gained PCR fragments were confirmed by sequencing. In addition, Merkel cell carcinoma cell lines MS-1 and MCC26 were used as positive and negative controls, respectively, to detect MCPyV.

**MCPyV detection by fluorescence in situ hybridization (FISH)**

MCPyV FISH was performed as previously described. In brief, deparaffinized 3 μm thick sections were pretreated with 0.2 M hydrochloric acid, incubated with 1 M NaSCN and digested with 1 mg/mL pepsin (2500–3500 U/mg, Sigma Chemical, St. Louis, MO, USA). The biotin labeled “specific” MCPyV DNA probe was added to the samples at a concentration of 5 ng/μL, followed by denaturation of DNA (five minutes, 80°C) and hybridization overnight (37°C, humid chamber; ThermoBrite System, Abbott Molecular, Abbott Park, IL, USA). Unbound MCPyV DNA probe was stringently washed away. Bound probe was detected by sequential incubation in a combination of secondary antibodies: fluorescein isothiocyanate (FITC) avidin secondary antibody (1:500) and biotin conjugated goat anti-avidin (1:100; Vector, Brunschwig Chemie, Amsterdam, The Netherlands). Prior to incubation, aspecific binding sites were blocked with Boehringer Blocking reagent (Roche, Molecular Diagnostics Inc., South Branchburg, NJ, USA). Cell nuclei were counterstained, and cover slipped with 4,6-diamidino-2-phenylindole dihydrochloride (DAPI; 0.2 μg/mL; Vectashield, Vector Laboratories, Burlingame, CA, USA). Samples were visualized using a DM 5000B fluorescence microscope (Leica, Wetzlar, Germany) coupled to an online digital camera (Leica DC 300 Fx) for independent evaluation of FISH signals by two investigators.

**RNA-in situ hybridization**

RNAscope 2.5 HD RED assay (cat. no. 322350; Advanced Cell Diagnostics, Newark, CA, USA) was performed using
Table 1  Clinicopathological data and MCPyV DNA PCR, FISH, IHC, and RISH results with HPyV7 positivity of the tissues

| ID  | G  | Age | MG   | Thymtype | Masaoka-Koga | Anti AChR IS/ster. | M1/M2 (178 bp) | VP1 (351 bp) | LT3 (308 bp) | MCPyV FISH | MCPyV RISH | MCPyV CM2B4 | sTAG-PCR | LTAg-PCR | FISH | IHC | 2t10 |
|-----|----|-----|------|----------|-------------|-------------------|-----------------|-------------|-------------|------------|------------|------------|----------|----------|------|-----|------|
| 1–1 | F  | 73  | +    | B1/B2    | I           | +                  | +               | –           | –           | NA         | +          | +          | –         | –        | –    |     | –    |
| 1–2 | F  | 75  | +    | B1/B2    | I           | +                  | +               | –           | –           | NA         | +          | +          | –         | –        | –    |     | –    |
| 1–11| F  | 34  | +    | B       | I           | +                  | –               | –           | –           | –          | –          | ++         | –         | –        | –    |     | –    |
| 1–12| F  | 36  | +    | A        | I           | –                  | +               | (+)         | –           | +          | –          | –          | –         | –        | –    |     | –    |
| 1–16| F  | 34  | +    | A        | I           | +                  | +               | (+)         | –           | +          | n.d.       | –          | –         | –        | –    |     | –    |
| 1–17| M  | 69  | +    | A8       | I           | +                  | –               | –           | –           | –          | ++         | –          | –         | –        | –    |     | –    |
| 1–21| F  | 38  | +    | A8       | I           | –                  | –               | –           | –           | –          | +          | ++         | –         | –        | –    |     | –    |
| 1–22| M  | 65  | +    | A/B/B2   | I           | +                  | –               | (+)         | NA         | n.d.       | –          | –          | –         | –        | –    |     | –    |
| 1–28| M  | 38  | –    | B1       | I           | NA                 | NA             | –           | –           | –          | –          | –          | –         | –        | –    |     | –    |
| 1–31| F  | 82  | +    | A        | I           | +                  | +               | –           | –           | –          | +          | +          | +         | –        | –    |     | –    |
| 1–32| M  | 47  | +    | B2       | I           | +                  | –               | –           | –           | n.d.       | –          | n.d.       | –         | –        | –    |     | –    |
| 1–34| M  | 59  | –    | A8       | I           | NA                 | NA             | –           | –           | –          | +          | ++         | –         | –        | –    |     | –    |
| 1–36| F  | 82  | –    | A8       | I           | NA                 | NA             | –           | –           | –          | –          | –          | (+)       | ++       | +    |     | –    |
| 1–39| F  | 78  | –    | B1       | I           | NA                 | NA             | –           | –           | –          | –          | –          | –         | –        | –    |     | –    |
| 1–43| F  | 78  | –    | A8       | I           | NA                 | NA             | +           | –           | +          | n.d.       | –          | –         | –        | ++   | (+) | –    |
| 1–23| M  | 37  | +    | A8       | IIA         | +                  | –               | –           | –           | –          | +          | –          | –         | –        | –    |     | –    |
| 1–24| M  | 68  | +    | B2       | IIA         | +                  | –               | –           | –           | –          | +          | ++         | +         | –        | –    |     | –    |
| 1–33| F  | 43  | +    | B2       | IIA         | +                  | –               | –           | –           | –          | –          | ++         | +         | –        | –    |     | –    |
| 1–35| M  | 45  | +    | A8       | IIA         | +                  | –               | –           | –           | –          | –          | ++         | +         | –        | –    |     | –    |
| 1–37| M  | 77  | +    | A8       | IIA         | +                  | –               | –           | –           | –          | –          | +++        | (+)       | –        | –    |     | –    |
| 1–38| F  | 80  | +    | A8       | IIA         | +                  | –               | –           | –           | +          | –          | ++         | +         | –        | –    |     | –    |
| 1–3 | F  | 57  | +    | B2/B3    | IIB         | +                  | –               | –           | –           | –          | –          | –          | –         | –        | –    |     | –    |
| 1–5 | M  | 37  | +    | B3       | IIB         | +                  | +               | –           | –           | –          | +          | ++         | +         | –        | –    |     | –    |
| 1–7 | F  | 79  | +    | A        | IIB         | –                  | –               | –           | –           | –          | +          | ++         | +         | –        | –    |     | –    |
| 1–8 | M  | 58  | +    | A/B2     | IIB         | (+)                | (+)            | +           | ++         | (+)        | (+)        | +          | ++        | (+)      | –    |     | –    |
| 1–9 | M  | 64  | –    | B2       | IIB         | NA                 | NA             | –           | –           | –          | +          | ++         | (+)       | –        | –    |     | –    |
| 1–15| F  | 53  | +    | B3       | IIB         | +                  | +               | –           | –           | –          | –          | –          | –         | –        | –    |     | –    |
| 1–26| M  | 65  | –    | A8       | IIB         | NA                 | NA             | –           | –           | (+)        | –          | –          | –         | –        | –    |     | –    |
| 1–42| F  | 57  | +    | A8       | IIB         | +                  | –               | –           | –           | –          | –          | –          | –         | –        | –    |     | –    |
| 1–10| F  | 73  | +    | B3       | III         | +                  | +               | –           | –           | –          | –          | –          | –         | –        | –    |     | –    |
| 1–13| F  | 54  | –    | B2       | III         | NA                 | NA             | –           | –           | –          | –          | –          | –         | –        | –    |     | –    |
| 1–20| M  | 64  | –    | B3       | III         | NA                 | NA             | ++          | ++         | ++         | +          | n.d.       | –         | –        | –    |     | –    |
target probes to MCPyV on tissue sections according to the manufacturer’s instructions and methods followed by Wang et al. Human peptidylprolyl isomerase B (Hs-PPIB) and bacterial dihydrodipicolinate reductase (DapB) were used as positive and negative controls, respectively. FFPE HeLa cells (Advanced Cell Diagnostics) were also used as negative and positive controls, as suggested by Advanced Cell Diagnostics. In addition, we used a FFPE MCC tissue block that previously tested MCPyV-positive by DNA-PCR and immunohistochemistry as a positive control. Four investigators independently evaluated the slides.

**Immunohistochemistry (IHC) and double staining procedure**

We used an anti LT-antigen MCPyV antibody (clone: CM2B4, dilution 1:50; Santa Cruz Biotechnology Inc., Santa Cruz, CA, USA). Immunohistochemical staining was conducted on a Dako Autostainer Link 48 using the EnVision FLEX Visualization Kit K8008 (Dako, Carpinteria, CA, USA) according to standard diagnostic routine protocols and manufacturer’s instructions.

**Results**

**MCPyV detection by DNA PCR and FISH**

A total of 13 of 36 (36.1%) thymomas tested positive using M1/2, 8 (22.2%) tested positive using VP1, and 4 tested positive using LT3 primers (Table 1). Sequence analyses of the PCR amplicons identified all PCR products as MCPyV DNA sequences. In total, 7 (19.4%) of 36 thymomas tested positive for two out of three different MCPyV DNA PCRs (Table 1, Fig 1) and thus were considered MCPyV-positive. Of these seven MCPyV-positive thymomas, five had a history of MG. In these five, the presence of MCPyV was confirmed by MCPyV FISH (Fig 2a). FISH analysis revealed relative weak punctuate hybridization signals, which were mostly heterogeneous patterns in all cases tested. None of the follicular hyperplasia or fetal thymus tissues were positive for MCPyV-DNA.

**MCPyV LT IHC and MCPyV LT RNA-ISH**

All seven positive MCPyV thymomas were assessed for MCPyV LT-antigen expression by IHC. Three out of seven (42.9%) revealed the presence of LT-antigen in thymoma tissues (Fig 2e, Table 1). The expression of MCPyV LT in these tissues was weak. In addition, three thymoma tissues, two MCPyV-negative (ID 1–26 and ID 1–32) and one MCPyV-positive (ID 1–8); and one MCPyV-positive MCC tissue were analyzed for MCPyV LT-antigen RNA (Fig 2bd,
The MCC cells revealed strong MCPyV RNA detection in the cytoplasm (Fig 2c). ID 1–8 showed weak detection of MCPyV RNA. ID 1–26 and ID 1–32 showed no RNA amplification (Fig 2c, Table 1).

Of interest, thymoma tissue ID 1–8 was positive for both polyomaviruses: HPyV7 and MCPyV (Fig 2f, Table 1), whereas the MCPyV expression was weaker compared to HPyV7.

**Discussion**

We recently reported the presence of HPyV7 in 46.0% (protein level) to 62.2% (DNA level) of thymomas. Herein, we assessed the presence of MCPyV on DNA level by PCR and FISH, on RNA level by RNA-ISH (RISH), and on protein level by IHC. In order to reliably detect MCPyV in FFPE tissues, Moshiriet al. recently reported that the use of a PCR-based approach in combination with the CM2B4 anti LT-antigen antibody yielded the highest specificity and sensitivity. In the present study, we used a PCR based approach in combination with MCPyV FISH, RISH, and IHC in order to comprehensively elucidate the presence of MCPyV in the same clinicopathologically well-classified thymoma cohort in which we recently assessed the presence of HPyV7. None of thymic hyperplasia (n = 20) or fetal thymic tissues were positive for MCPyV assessed by PCR. Overall 19.4% (n = 7) thymomas tested positive for the presence of MCPyV DNA (VP1, M1/2, and LT3). The presence of MCPyV DNA was confirmed in six of seven PCR positive thymomas by FISH analysis. The bands and signals obtained by PCR and FISH were weak. Correspondingly, weak MCPyV LT-antigen expression on protein level was found in three of seven thymomas. The prevalence of MCPyV was far lower than the prevalence of HPyV7 in these thymomas.

Of all MG-positive thymomas tested for MCPyV, five were positive on DNA level and three on protein level. Based on this small number, no conclusion can be drawn concerning an association between MCPyV and MG in thymoma. In addition, the weak DNA and protein detection correlates with the weak prevalence of MCPyV RNA compared to the RNA detection of the MCPyV-positive MCC tissue assessed by RISH in ID 1–8. In contrast to Torrachio et al. and Hashidait al. we used a larger number of MCPyV detection techniques in a larger thymoma cohort, which very likely explains why we were able to identify some MCPyV positive thymoma cases.18,19

**Figure 1** Representative PCR of 11 thymic tissues. Representative results of Merkel cell polyomavirus (MCPyV) M1/2 (178 base pairs [bp]), VP1 (351 bp), and LT3 (308 bp) PCR primer set of 11 human epithelial thymic DNA and two Merkel cell carcinoma cell lines. MS-1 was used as a positive control and MCC26 as a negative control. Amplification could be observed for VP1 and M1/2 primer sets for ID 1-8, ID 1-9 and ID 1-20. For the LT3 primer set, ID 1-8, ID 1-20, and ID-26, a PCR product could be detected. free, free space; H2O, water control; MW, molecular weight marker.
However, given the weak DNA amplification and the low protein expression in these few, it can be concluded that MCPyV is weakly present in three thymomas, thus MCPyV is very unlikely to play a role in human thymomagenesis. In the context of our previously reported frequent finding of HPyV7 within the same thymoma cohort, the low prevalence of MCPyV indirectly supports a role of HPyV7 in human thymomagenesis.

Remarkably, most thymoma cases that tested positive for HPyV7 or MCPyV in this cohort were of stage II or III according to the Masaoka classification (Table 1). Eleven of 14 stage II thymomas were positive for one of these human polyomaviruses. In addition, one of the three MCPyV positive tissues was classified as stage III B (Table 1). The higher thymoma stages are considered to be more invasive. The sTAG of MCPyV was recently observed to provide cells the ability to migrate; however, in as much this applies to HPyV7 and thymic epithelial cells currently remains speculation.

Of interest, to the best of our knowledge, we show for the first time the concomitant detection of HPyV7 and MCPyV in the same tumor tissue using diverse molecular techniques. Interestingly, MCPyV and HPyV7 are also present with a high seroprevalence in the same blood. Detection of the MCPyV genome by PCR and FISH reveals that MCPyV can infect thymic epithelial cells. In the present study we show that thymic epithelial tumors can harbor both HPyV7 and MCPyV, although the prevalence of MCPyV in human epithelial tumors is far lower than reported for HPyV7. Our data strongly indicate that MCPyV is very unlikely to play an important role in the etiopathogenesis of human thymomas.

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**Disclosure**

No authors report any conflict of interest.

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**Supporting Information**

Additional Supporting Information may be found in the online version of this article at the publisher’s website:

**Table S1.** Clinicopathological Data of follicular hyperplasia tissue results with human polyomavirus 7 (HPyV7) positivity, as published by Rennspiess et al., 2017. Anti-AChR, anti-acetylcholine receptor antibodies; F, female; G, gender; ID, laboratory identification; IS/Ster., immunosuppression/steroids; M, male; MG, myasthenia gravis; NA, not applicable; Thym. type, thymoma type.