

# Merkel Cell Carcinoma in the Age of Immunotherapy: Facts and Hopes

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## Abstract

Merkel cell carcinoma (MCC) is a rare (~2,000 U.S. cases/year) but aggressive neuroendocrine tumor of the skin. For advanced MCC, cytotoxic chemotherapy only infrequently (<10% of cases) offers durable clinical responses (>1 year), suggesting a great need for improved therapeutic options. In 2008, the Merkel cell polyomavirus (MCPyV) was discovered and is clonally integrated in approximately 80% of MCC tumors. The remaining 20% of MCC tumors have large numbers of UV-associated mutations. Importantly, both the UV-induced neoantigens in virus-negative tumors and the MCPyV T antigen oncogenes that are required for virus-positive tumor growth are immunogenic. Indeed, antigen-specific T cells detected in patients are frequently dysfunctional/"exhausted," and the inhibitory ligand, PD-L1, is often present in MCC tumors. These findings led to recent clinical trials involving PD-1 pathway

blockade in advanced MCC. The combined data from these trials involving three PD-1 pathway blocking agents—avelumab, pembrolizumab, and nivolumab—indicated a high frequency of durable responses in treated patients. Of note, prior treatment with chemotherapy was associated with decreased response rates to PD-1 checkpoint blockade. Over the past year, these striking data led to major changes in advanced MCC therapy, including the first-ever FDA drug approval for this disease. Despite these successes, approximately 50% of patients with MCC do not persistently benefit from PD-1 pathway blockade, underscoring the need for novel strategies to broaden antitumor immune responses in these patients. Here, we highlight recent progress in MCC including the underlying mechanisms of immune evasion and emerging approaches to augment the efficacy of PD-1 pathway blockade. *Clin Cancer Res*; 24(9); 2035–43. ©2017 AACR.

## Introduction

Merkel cell carcinoma (MCC) is a rare (~2,000 U.S. cases/year) but aggressive skin cancer with a high risk of recurrence (27%–31%; refs. 1–3). Although MCC is rare, its incidence is rising steadily (4, 5). Risk factors include advanced age, sun/UV exposure, and chronic immunosuppression (~8% of patients with MCC have hematologic malignancy, solid organ transplant, or HIV/AIDS; ref. 6). Although 92% of patients with MCC are not immunosuppressed, individuals who have chronic T-cell dysfunction have an increased likelihood of developing MCC (10- to 30-fold; refs. 6–8). Only 4% of MCC cases occur in patients under 50 years of age, and MCC risk increases significantly with every additional decade of life (4, 9), likely due in part to increased immune senescence. The disease-associated mortality of MCC is 46% within 5 years (10), highlighting the need for improved therapeutic strategies.

## Presentation/Diagnosis

The presentation of MCC can be challenging for physicians to recognize (Fig. 1), in part, due to its rarity. In two thirds of cases,

physicians suspect a benign lesion based on clinical appearance (6). The following mnemonic summarizes features associated with MCC: Asymptomatic, Expanding rapidly, in an Immune-suppressed patient Older than 50 and on UV-exposed skin (AEIOU; ref. 6). As 89% of MCCs had three or more of these features (6), this mnemonic is sensitive; however, it is not specific for MCC, as such lesions may often represent another nonmelanoma skin cancer or a benign lesion such as an inflamed cyst. MCC diagnosis is confirmed through pathologic review of a biopsied lesion. Pathologic sections of MCC exhibit small cells with little cytoplasm (Fig. 1). The histologic recognition of MCC was greatly facilitated by the determination that perinuclear, coarsely granulated CK20 (KRT20) staining is present in 90% of MCC cases (11, 12).

## Virus-induced MCC

Early studies indicated that MCC can be linked to decreased immune function. One key study found that patients with HIV have a 13-fold increased MCC risk compared with population controls (8). Also, case reports have described the uncommon, spontaneous regression of MCC tumors under a variety of scenarios, further indicating a link to the immune system (13–15). These data collectively suggested that MCC may be linked to a pathogen. In 2008, the Merkel cell polyomavirus (MCPyV) was discovered, and it is now clear that this virus plays a key role in the majority of MCC cases (16).

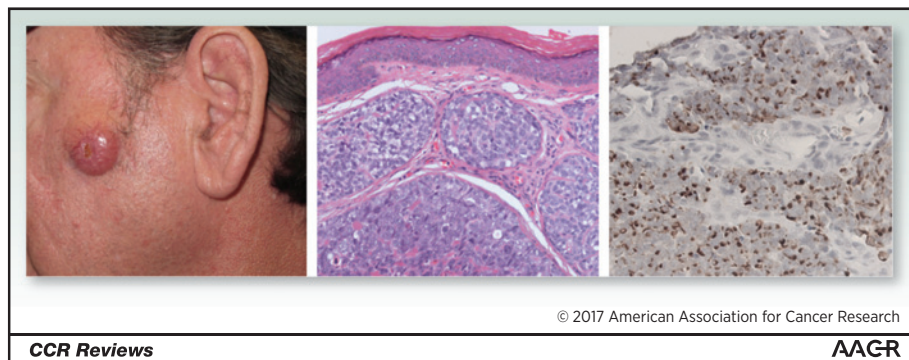
MCPyV is a member of the polyomavirus family comprised of nonenveloped, double-stranded DNA viruses and is the first virus from this family known to cause cancer in humans. MCPyV-specific antibodies have been detected in approximately 45% of children and in 80% of individuals 50 years or older, indicating that it is highly prevalent in the population (17).

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**Figure 1.**

Clinical and histologic appearance of MCC. Left, clinical appearance of an MCC arising on the left cheek of a 55-year-old man. The tumor was red, firm, nontender, and rapidly growing on sun-exposed skin. The differential diagnosis would include other types of nonmelanoma skin cancer. Center, intradermal tumor with pleiomorphic cells with large nuclei and scant cytoplasm. Right, cytokeratin 20 (CK20) IHC staining exhibits the characteristic perinuclear expression of CK20, a highly diagnostic finding for MCC.

Interestingly, despite this high prevalence, MCPyV has not been shown to cause any disease other than when it very rarely leads to MCC. We now understand key aspects of the mystery of how a virus with an extremely high incidence leads to a cancer that is very rare.

MCPyV-related oncogenesis requires two separate events likely accounting for its rarity: (i) The circular double-stranded genome must be linearized and integrated into the host genome, perhaps after a DNA-damaging event (MCPyV-positive tumors frequently occur on sun-exposed skin), and (ii) the virus must be mutated, with loss of expression of the C-terminus of the large T (LT) antigen that is required for viral DNA replication (Fig. 2). Virus-induced MCC is driven, in part, by expression of truncated large T antigen that binds to and inactivates the tumor suppressor Rb (RB1; Fig. 2; ref. 18), promoting cell-cycle progression and uncontrolled proliferation (19, 20). Small T (sT) inhibits the proteasomal degradation of large T (21) as well as the oncoprotein cMyc (MYC) and cyclin E (CCNE1; ref. 21). Both large T and small T have been demonstrated to drive transformation in mammalian cells *in vitro* (18, 20, 22); however, numerous attempts to generate mouse models of MCC at best only partially emulate the disease in adult animals (23–25). These data indicate that additional, as yet undetermined factors are required for induction of MCPyV-associated MCC. Although several groups have successfully generated xenografts using MCC cell lines and postoperative tumor tissue, engraftment can be done only in NOD SCID IL2Rgamma<sup>-/-</sup> (NSG) mice, which have a severely impaired immune system. These xenograft models mimic the gross pathologic features of the corresponding patient's tumor but fail to recapitulate the tumor-immune interactions that are now understood to greatly affect patient outcomes. *In vitro* experiments have demonstrated that ongoing expression of MCPyV oncoproteins is required for survival of virus-positive MCC cells (26–28). These persistently expressed non-self-antigens can potentially elicit host immune recognition, and the limited size of MCPyV T antigens (<400 amino acids) has facilitated immune studies of MCPyV-specific T-cell responses (29–32).

### Antibodies to MCPyV T Antigen Correlate with Tumor Burden

The robust response to MCPyV-positive tumors can include both T-cell and humoral components (33–35). At the time of diagnosis, approximately half of MCC patients make antio-

odies to MCPyV oncoproteins. Knowing a patient's serostatus (MCPyV positive or negative) can be helpful for his or her subsequent care. The prognosis of seronegative patients is less favorable (42% higher risk of recurrence than seropositive patients; refs. 35, 36), and thus, these patients need to be followed closely with scans (36). For seropositive patients, antibody titers correlate with tumor burden (33, 34), and a rising titer is an early indicator of disease recurrence (33). These findings have recently been validated in a large prospective cohort (36), and the test is now included in the 2018 National Comprehensive Cancer Network (NCCN) guidelines for MCC (37). Effective surveillance is relevant to patient care because if disease recurrence is discovered early (when tumor burden is lower), immunotherapy may be more effective (38).

### UV-induced MCC

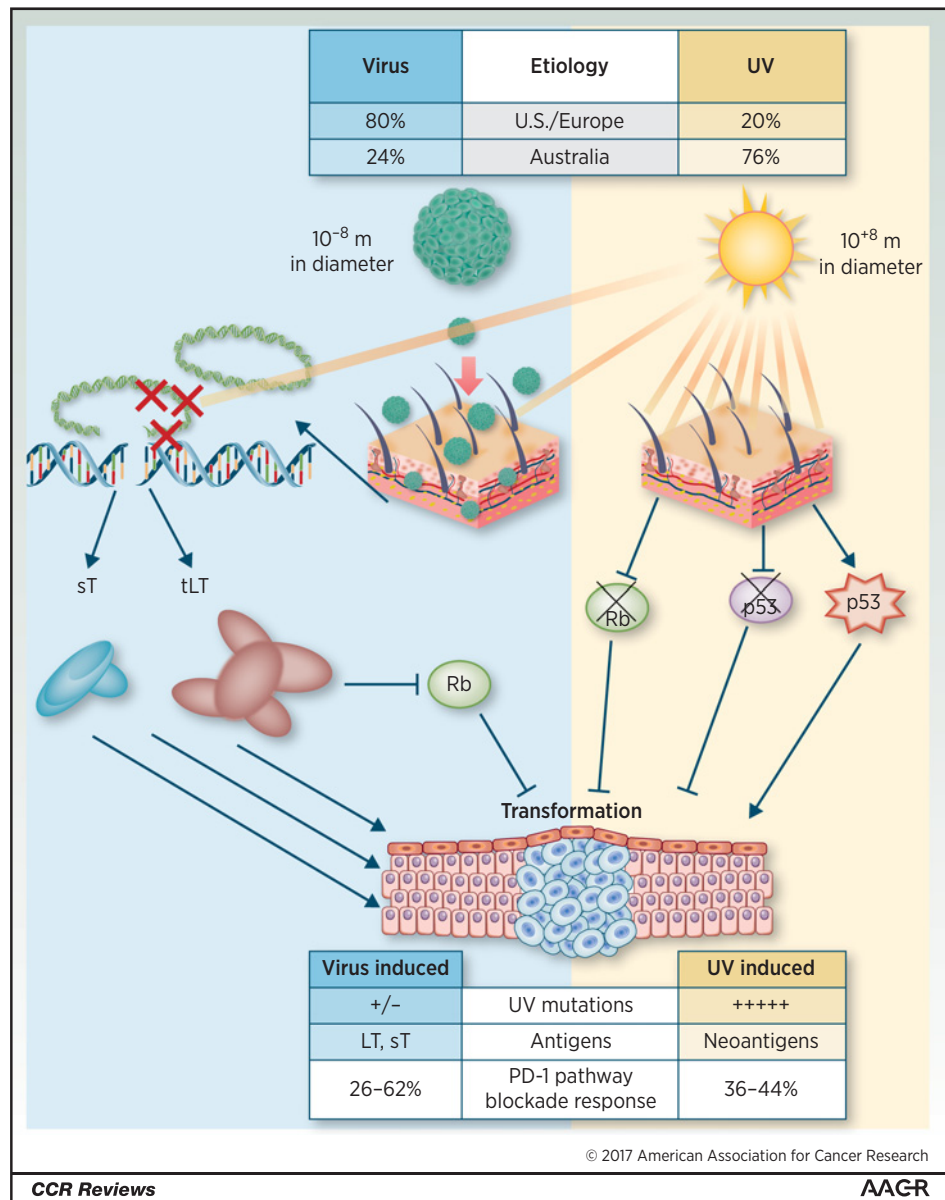
Some MCC tumors have no MCPyV detectable by either DNA-PCR or IHC, which raised the question of whether virus-negative MCC exists or whether viral detection techniques were insufficient (39). Recent studies have demonstrated that MCPyV-negative MCC tumors do indeed exist, with variable incidence around the world (~20% in United States/Europe vs. 76% in Australia; see Fig. 2; refs. 40–43). Strikingly, virus-negative MCC is among the most mutated of all solid tumors, including melanoma and non-small cell lung cancer (40–42). These mutations are mostly UV signature mutations (40–42). The high mutational burden (187–4,707 somatic single-nucleotide variants per exome) in MCC correlates to frequent amino acid changes and large numbers of UV-induced neoantigens (42). The most common mutations found in MCPyV-negative MCC are in *p53* (*TP53*; 75% of cases) and *Rb* (67% of cases), commonly resulting in loss of functional protein expression (42). However, activating mutations also comprise a large proportion of the *p53* mutations detected (45% of *p53* mutations in MCC; ref. 42).

### Chemotherapy: The Previous Standard of Care

Definitive treatment of primary MCC includes surgery and/or radiation. This has been quite well established, and the consensus is summarized in the 2018 NCCN guidelines (37). Historically, chemotherapy was the preferred treatment option for advanced MCC despite a lack of data rigorously assessing its benefit in this setting. Recently, several careful retrospective studies have been carried out in the United States and Europe

**Figure 2.**

Comparison of virus-positive and virus-negative MCC tumors. This schematic depicts the two major causes of MCC, their prevalence, differences in their potential immune targets, and frequencies of response to immune therapy. Top, differences in MCC prevalence—United States (U.S.)/ Europe versus Australia. Left, virus-induced tumorigenesis—the highly prevalent MCPyV is often found on normal skin. Rarely, MCPyV will integrate into the host genome, and through a separate rare event, large T will become truncated (tLT; depicted by red Xs) prior its C-terminal. Expression of the sT and tLT viral oncogenes is tumorigenic through multiple pathways including inhibition of wild-type cellular Rb (see text). Right, UV-induced tumorigenesis—sun exposure results in the generation of many UV signature mutations (C→T mutations). The most common of which are in *Rb* and *p53*. *Rb* is frequently found to be inactivated in UV-induced MCC tumors (67%). Mutation of *p53* includes both activating and inactivating mutations (16, 18–21, 40–43, 71–75).



that document chemotherapy response rates and their durability (summarized in Table 1). A U.S. academic center–based study of 62 patients with distant metastatic MCC showed a first-line chemotherapy objective response rate (ORR) of 55%;

however, the median progression-free survival (PFS) was only 94 days after chemotherapy initiation, and the median overall survival was 9.5 months (44). Second-line chemotherapy was even less favorable, with an ORR of 23% and a median PFS of

**Table 1.** Selected data for chemotherapy and anti-PD1/PD-L1 in MCC

| Line                     | Chemotherapy  |          | Nivolumab<br>≥1st line | Avelumab<br>≥2nd line | Pembrolizumab<br>1st line |
|--------------------------|---|----------|------------------------|-----------------------|---------------------------|
|                          | 1st line  | 2nd line |                        |                       |                           |
| Cohort size              | 62–67   | 20–30    | 22                     | 88                    | 25                        |
| Agent                    | Etoposide and platinum-based agent <sup>b</sup>                   |          | Anti-PD-1              | Anti-PD-L1            | Anti-PD-1                 |
| ORR                      | 31%–55%   | 9%–23%   | 68%                    | 32%                   | 56%                       |
| 9-month PFS <sup>a</sup> | 15%–26%   | 0%–3%    | N/A <sup>c</sup>       | 33%                   | 56%                       |
| Publications             | Becker, 2017 <sup>d</sup> (46); Cowey, 2017 (45); Iyer, 2016 (44) |          | Topalian, 2017 (52)    | Kaufman, 2016 (54)    | Nghiem, 2016 (50)         |

<sup>a</sup>Values estimated from charts.

<sup>b</sup>Most commonly used agents.

<sup>c</sup>9-Month PFS is not yet available; however, 3-month median PFS is 82%.

<sup>d</sup>Data for second-line chemotherapy only.

61 days (44). An independent study of 67 patients with metastatic MCC in the US Oncology Network also assessed responses to first- and second-line chemotherapy (45). This study found a first-line chemotherapy ORR of only 31%, with a median PFS of 4.6 months. Patients on their second or later line of chemotherapy had an ORR of 20% and a median PFS of 2.1 months (45). In a cohort of 34 patients from Europe whose disease had progressed following at least one line of chemotherapy, the patients' next line of chemotherapy had only a 9% ORR and a median duration of response of 1.9 months (46). These studies indicate that although MCC has a relatively high response rate to chemotherapy in the first line, responses are typically short-lived and resistance develops quickly. Multiple mechanisms are likely involved with the disappointing long-term benefit of chemotherapy in MCC. These may include its immunosuppressive effects in the setting of this immunogenic cancer as well as established mechanisms such as resistance to apoptosis (47).

### Immunotherapy: A New Standard of Care

Over the last decade, several lines of evidence have suggested that immune status is linked to clinical outcomes in MCC, indicating that augmenting cell-mediated immunity could be beneficial. An early study focusing on tumor-infiltrating lymphocytes found that patients with robust CD8<sup>+</sup> lymphocyte infiltration into MCC tumors had 100% MCC-specific survival compared with 60% survival in those with little or no CD8<sup>+</sup> infiltration (48). These data indicated that infiltration by CD8<sup>+</sup> T cells had profound prognostic value and that augmenting immune function could benefit patients with MCPyV-driven MCC. The specificity of CD8<sup>+</sup> lymphocytes was then studied, and MCPyV oncoprotein-specific cells were found to be present in MCC patient blood and enriched in patients' tumors (29, 30). Importantly, signs of dysfunction were evident in MCPyV-specific CD8<sup>+</sup> T cells from patients, as they expressed both PD-1 (PDCD1) and Tim3 (HAVCR2), the combination of which suggests functional exhaustion (29). When the tumor microenvironment was investigated, 49% of 49 tumors contained PD-L1 (CD274, typically expressed on antigen-presenting cells) and expression tended to correlate with the presence of intratumoral lymphocytes (49). In aggregate, these findings made a compelling case for testing PD-1 pathway blockade in MCC.

To date, three antibodies targeting the PD-1 axis have been studied in MCC, with all three showing substantial response rates and impressive durability of responses (summarized in Table 1). Although the numbers of patients studied are small compared with other more prevalent cancer types, these early trials have demonstrated frequent therapeutic durability, whereas there was previously little hope for patients with advanced MCC. A National Cancer Institute-sponsored clinical trial studied pembrolizumab (anti-PD-1) in 25 patients with advanced MCC who had not received prior systemic therapy. The investigators found an ORR to pembrolizumab of 56% including a 16% complete response rate. Of the 14 responsive patients, the response duration ranged from at least 2.2 months to at least 9.7 months. Overall, the trial had an estimated PFS of 67% at 6 months. Pembrolizumab was effective in both virus-negative and virus-positive tumors (ORR of 62% and 44% respectively, not significantly different; ref. 50). The early results of this trial led to pembrolizumab being listed as a

treatment option for advanced disease in the 2017 NCCN guidelines for MCC (51).

An international, single arm, open-label trial of nivolumab (anti-PD-1) included both patients who had and who had not received prior chemotherapy (36% and 64%, respectively). In this study, 15 of 22 patients (68%) had objective responses, and PFS at 3 months was 82% with the trial still ongoing (52).

A large international clinical trial studied avelumab (anti-PD-L1) in 88 patients with distant metastatic disease who had previously received at least one line of chemotherapy. This trial found an ORR of 33%, with a complete response rate of 11%. At 6 months, PFS was 40%, and the estimated PFS at 1 year was 30%. As with pembrolizumab, avelumab was found to be effective in both virus-positive and virus-negative tumors (ORR of 26% and 35%, respectively, not significantly different; refs. 53, 54). In March 2017, these remarkable data in chemotherapy-refractory MCC led to the first-ever FDA approval of a drug for this cancer. Avelumab was granted accelerated approval in advanced MCC in patients at least 12 years of age whether or not they have previously received chemotherapy (55).

Now that avelumab has been approved for treatment of advanced MCC, an important question remains: namely, whether treatment with PD-1 pathway blockade in the adjuvant setting is appropriate and/or beneficial for treatment of this aggressive disease. As with other cancer treatments in general, catching and treating the tumor early correlates with improved prognosis. This possibility, in the context of PD-1 pathway blockade in primary MCC, will be addressed by two (one of which is double blinded and randomized) clinical trials that are now recruiting (Table 2).

Anti-PD-1 checkpoint blockade therapies have proven to be well tolerated in a majority of patients. However, altering the balance of immune homeostasis can induce autoimmunity that results in grade 3 or grade 4 toxicity in 10% to 22% of cases (56, 57). As such, informed consent of patients is critical, particularly because immune-related adverse events (irAE) are typically idiosyncratic, making their early recognition and treatment challenging.

Importantly, the unique therapeutic benefits of these agents raise the question of whether they are indicated in patients who have a known autoimmune condition or previous irAE to ipilimumab. Indeed, patients with MCC exhibit higher numbers of autoimmune conditions than the population at large. Treatment of autoimmune disease is a major known iatrogenic cause of chronic, severe immune suppression that can increase the risk of multiple cancer types, including MCC (58). A recent retrospective analysis of 52 melanoma patients with prior autoimmune disease treated with PD-1 pathway blockade found comparable ORRs (33%) to those observed in clinical trials that have excluded patients with autoimmunity (59). Although 20 (38%) patients had a flare of autoimmune disease and another 15 (29%) developed other irAEs, only eight patients exhibited grade 3 toxicity of a preexisting autoimmune process or irAE, and just two patients permanently discontinued treatment. A separate study of 67 patients who had prior major ipilimumab toxicities exhibited a 40% ORR with PD-1 blockade (59). In this cohort, 25 (37%) patients experienced recurrence of ipilimumab-induced irAEs or developed new/different irAEs. Although 14 (21%) patients exhibited grade 3 to 4 irAEs, only eight (21%) patients discontinued therapy. In both of these cohorts, a majority of the immune toxicities could

**Table 2.** Selected immune therapy clinical trials for Merkel cell carcinoma

| NCT identifier                                 | Trial arms   | Recruitment status     | Phase | Targeted enrollment | Comments   | Publications        |
|--|--|------------------------|-------|---------------------|--|---------------------|
| <b>Anti-PD-1/PD-L1 monotherapy</b>             |  |                        |       |                     |  |                     |
| NCT02155647                                    | Avelumab as $\geq$ 2nd line  | Active, not recruiting | II    | 88                  | 28 of 88 chemotherapy-refractory patients achieved a response including eight complete responses (ORR = 32%)   | Kaufman, 2016 (54)  |
| NCT02155647                                    | Avelumab as 1st line   | Recruiting             | II    | 112                 | Preliminary results show an objective response in 11 of 16 patients (ORR = 69%)  | D'Angelo, 2017 (76) |
| NCT02267603                                    | Pembrolizumab as 1st line  | Active, not recruiting | II    | 50                  | Four of 25 patients evaluated had a complete response and 10/25 had a partial response (ORR = 56%)   | Nghiem, 2016 (50)   |
| NCT02488759                                    | Nivolumab as 1st or $\geq$ 2nd line  | Active, not recruiting | I/II  | 25                  | 22 patients initially evaluated on nivolumab alone, 12 had a partial response, and three had a complete response (ORR = 68%)                         | Topalian, 2017 (52) |
| NCT02196961                                    | Avelumab as adjuvant versus observation following resection  | Recruiting             | II    | 113                 | Only in Europe <sup>a</sup>  |                     |
| NCT03271372                                    | Avelumab as adjuvant 1st line  | Recruiting             | III   | 100                 | Stage III/IIIB nodal disease, randomized, double blinded   |                     |
| <b>Checkpoint blockade combination therapy</b> |  |                        |       |                     |  |                     |
| NCT02488759                                    | Nivolumab $\pm$ anti-LAG3 (BMS-9861016) $\pm$ ipilimumab (many arms)   | Recruiting             | I/II  | 500                 | Cohort of patients with virus-associated cancers   |                     |
| NCT03071406                                    | Ipilimumab + nivolumab versus ipilimumab + nivolumab + stereotactic body radiation therapy   | Recruiting             | II    | 50                  |  |                     |
| <b>Innate immunity agents and cytokines</b>    |  |                        |       |                     |  |                     |
| NCT02035657                                    | TLR-4 agonist, GLA-SE  | Completed              | I     | 10                  | Two of three patients with local nodal disease had a complete response, and two of seven patients with distant metastatic disease had stable disease | Bhatia, 2016 (60)   |
| NCT01440816                                    | IL12-EP  | Completed              | II    | 15                  | Four of 15 patients treated with IL12 had an objective response  | Bhatia, 2015 (61)   |
| <b>Cell-based therapies</b>                    |  |                        |       |                     |  |                     |
| NCT02584829                                    | Autologous MCPyV-specific CD8 cells + avelumab + MHC upregulation versus avelumab + MHC upregulation   | Recruiting             | I/II  | 20                  | Four of four patients had responses with 3/4 complete responses  | Paulson, 2017 (66)  |
| NCT02465957                                    | NK cells (activated NK-92) + ALT-803 (modified IL15)   | Closed                 | II    | 24                  | Initial three patients showed no major toxicities, and at least one patient had a response   | Bhatia, 2016 (64)   |
| <b>Oncolytic virus therapies</b>               |  |                        |       |                     |  |                     |
| NCT02819843                                    | T-VEC versus T-VEC + hypofractionated radiotherapy   | Recruiting             | II    | 34                  | Cohort of melanoma and MCC   |                     |
| NCT02978625                                    | T-VEC + nivolumab  | Not yet recruiting     | II    | 68                  | Cohort of refractory lymphomas and refractory nonmelanoma skin cancers   |                     |
| <b>Biomarker-guided combination therapy</b>    |  |                        |       |                     |  |                     |
| NCT03167164                                    | Avelumab, bevacizumab, capecitabine, cisplatin, cyclophosphamide, 5-fluorouracil, leucovorin, nab-paclitaxel, omega-3-acid ethyl esters, stereotactic body radiation therapy, ALT-803, NK-92 (many arms) | Not yet recruiting     | I/II  | 67                  | Treatment customized on the basis of tumor-specific characteristics  |                     |

NOTE: Therapies in the order listed in table: avelumab = anti-PD-L1 (IgG1); pembrolizumab = anti-PD-1 (IgG4); nivolumab = anti-PD-1 (IgG4); ipilimumab = anti-CTLA-4; GLA-SE = glucopyranosyl lipid A in stable emulsion, a TLR-4 agonist; F16-IL = anti-tenascin C mAb-IL2 fusion protein; IL12-EP = IL12 plasmid administered with electroporation; MHC upregulation via radiation or intratumoral IFN $\beta$  administration; NK-92 = activated, irradiated, allogenic natural killer cells; ALT-803 = IL15 superagonist complex; bevacizumab = anti-VEGF; T-VEC = talimogene laherparepvec, an engineered herpes oncolytic virus.

<sup>a</sup>Unless otherwise noted, trials include sites within the United States.

be controlled by symptom management, oral steroids, and/or steroid sparing agents (>80% of all irAEs observed). Taken as a whole, this study indicates that, after appropriate informed

consent discussions with the patient, PD-1 pathway blockade may be considered despite the presence of prior autoimmune disease or ipilimumab-induced irAEs (59).



## New Immunotherapy Trials: A Diverse Pipeline

Despite the greatly improved durable responses observed through PD-1 checkpoint blockade therapy compared with chemotherapy, major challenges remain in systemic therapy for MCC in that nearly half of patients do not derive durable benefit from these drugs. To address this issue, numerous clinical trials are underway for MCC, at least nine of which involve immune therapy (Table 2). These trials involve four general strategies that will be summarized below: (i) "removing an additional brake" (i.e., CTLA-4) on the immune system, (ii) "stepping on the gas" by using innate or other immune agonists, (iii) "adding more troops" by infusing more of the relevant cells into the patient, and (iv) "weaponizing viruses" that can specifically target and kill cancer cells while preserving normal tissues.

Activated T cells express CTLA-4 (CTLA4) that suppresses their function after CTLA-4 binds its cognate receptor (CD80/CD86) on an antigen-presenting cell. In this way, CTLA-4 acts as a central type of immunologic "brake." Anti-CTLA-4 antibody (ipilimumab) blocks this binding and allows the T cell to remain in a more active state. Ipilimumab efficacy in MCC is now being determined in clinical trials (Table 2). One trial enrolling patients in Germany will assess the safety and efficacy of ipilimumab or avelumab in the adjuvant setting following surgical resection of local MCC in comparison with resection alone. The ipilimumab arm of this trial has recently closed, whereas the arm investigating avelumab in the adjuvant setting is currently enrolling. In patients where PD-1 pathway blockade is ineffective, one hypothesis is that further augmentation of the immune response is required, possibly via CTLA-4. In a U.S.-based trial, 50 patients with metastatic MCC are being enrolled to test the safety and efficacy of the combination of nivolumab (anti-PD-1) and ipilimumab with and without stereotactic body radiation that can debulk the tumor and may induce immunogenic cell death. The combination of ipilimumab with PD-1 pathway blockade is also being performed in melanoma, and safety data from these trials would be expected to be similar.

Intratumoral immune infiltration and immune recognition/activation is regulated by pro- and anti-inflammatory molecules within the tumor-immune microenvironment. To increase the activity of antitumoral immune responses, several strategies seek to "step on the gas" by adding immune agonists that can reinvigorate antitumor T-cell responses. In a proof-of-concept trial, a Toll-like receptor-4 (TLR4) agonist, glucopyranosyl lipid-A stable emulsion (GLA-SE), was intratumorally injected into superficial MCC tumors (Table 2; ref. 60). In this trial, two of three patients with stage IIIB MCC were recurrence-free at 23+ and 19+ months with one patient having a pathologic complete response after two injections of this TLR-4 agonist (60). In a second cohort, two of seven patients with stage IV MCC had partial responses and were progression free after 13 months at the time of publication. Encouragingly, responses correlated with increased T-cell infiltration and activation of proinflammatory genes (60), providing proof of concept of this therapeutic approach.

Another trial of patients with superficial/accessible MCC tumors explored the utility of intratumoral electroporation of DNA encoding the potent proinflammatory cytokine IL12 (IL12A; Table 2; ref. 61). In this study, three of three patients

with local disease who received definitive surgery and/or radiotherapy at 4 weeks after one cycle of three IL12 treatments had recurrence-free survival of 2+, 9, and 32+ months, with one patient having a pathologic complete response (61). In a second arm of this trial involving 12 patients with metastatic disease, partial responses were seen in three patients and stable disease was seen in one patient (61). Treatment corresponded with induction of IL12 and TNF $\alpha$  (TNF) expression in the tumor microenvironment as well as enhanced T-cell infiltration over baseline. Encouragingly, 40% of the injected lesions exhibited regression (30% complete and 10% partial), and another 40% were stable (61). Regression of noninjected lesions was also observed, and no grade 3 or higher adverse events were reported. Although very preliminary, these results highlight the potential of local IL12 administration.

Cell-based therapy is an emerging immunotherapeutic approach, particularly in the setting of certain types of immune evasion. MCC evades immune detection through a variety of mechanisms, including downregulation of HLA class I molecules required for antigen presentation, which occurs in 74% to 84% of MCC tumors (62, 63). Natural killer (NK) cells typically target cells that downregulate HLA class I expression. An NK cell-based trial that accepts patients whose tumors were refractory to prior checkpoint therapy involves biweekly infusions of activated, irradiated, allogeneic NK-92 cells (Table 2). Thus far in this study, three patients treated with NK cells showed no major toxicities and although very preliminary, one patient, who had not responded to PD-1 pathway blockade, had a complete response (64).

MCPyV-positive MCC tumors require expression of viral T antigen oncoproteins (26–28). In patients with certain HLA types, MCPyV oncoprotein-specific T cells can be isolated and expanded *ex vivo* prior to therapeutic infusion. In one trial utilizing this strategy (Table 2), three of four patients given T cells plus HLA upregulation (tumor-targeted radiation or interferon) progressed, whereas one that had an initial complete response subsequently progressed after 14 months. It was found that the infused T cells frequently became dysfunctional/"exhausted" upon transfer (65, 66). As such, avelumab (anti-PD-L1) has been added in combination with these autologous virus-specific T cells. In this combined-therapy cohort, all four patients treated with a regimen including T cells, HLA upregulation, and avelumab experienced objective responses, with three complete responses at last follow-up (65, 66). These early results, in a limited set of patients, highlight the potential for the rational design and implementation of transgenic T-cell receptors against virus-positive MCC tumors.

A mechanistically relevant therapy, recently approved for melanoma, is the oncolytic virotherapeutic tamlogene laherparepvec (T-VEC; ref. 67). The viral genes have been mutated so that the construct is replication-defective in normal cells, but constitutively active proliferative pathways in tumor cells allow the virus to replicate and kill those cells. The T-VEC design also includes a granulocyte macrophage colony-stimulating factor (GM-CSF, CSF2) expression cassette to induce a proinflammatory immune response. T-VEC is currently being investigated in two trials that include MCC (Table 2). In the first trial, T-VEC is used alone or in combination with hypofractionated radiotherapy. Another trial combines T-VEC with nivolumab (anti-PD-1) to augment the immune response in conjunction with T-VEC-mediated killing.

## On the Horizon

The diversity of drugs in development and currently being tested in clinical trials greatly outstrips our understanding of the cellular and molecular mechanisms at play (68). Indeed, nearly half of patients do not derive persistent benefit from PD-1 pathway blockade and neither tumor viral status nor biomarker studies accurately identify patients who will not respond (50, 66). In addition, mutation, adaptation, and selection for therapeutically resistant cells are remarkably powerful processes that continue to blunt therapeutic efficacy for all classes of drugs, including immunotherapy (68). To begin to address questions of response and nonresponse, a comprehensive, unbiased examination of host and tumor immune interactions in the tumor microenvironment is required.

MCC offers a particularly fertile hunting ground for studying the immune responses to cancers more broadly due to: (i) the unique relevance of MCC as a model for studying immunogenic cancers (e.g., viral oncoprotein vs. high UV-mutational load); (ii) the robust immune evasion, likely through multiple mechanisms, required for a tumor to persist despite such a heavy viral/neoantigen burden; (iii) the small size of the MCPyVT antigen oncogenes that greatly facilitates immunologic studies; and (iv) the generation of tumor-specific reagents that facilitate both studies of the antitumor immune response and improved therapy. As such, investigations of MCC are poised to contribute to the understanding of the biology of cancer immunogenicity.

Now more than ever, we are able to delve into the cellular and molecular complexities within any given tumor. The cost of next-generation sequencing technologies is rapidly decreasing. Single-cell sequencing is capable of analyzing hundreds to thousands of cells from small core biopsies making serial analysis of tumor tissues following therapy both more feasible and less invasive.

Also, an ever-increasing number of targets can be stained using multiplexed IHC in combination with more sophisticated nucleic acid *in situ* hybridization techniques. In an attempt to combine this arsenal of molecular tools with clinical medicine, one trial will determine the genetic, transcriptomic, and proteomic details of a patient's tumor to customize therapy with immune and more traditional approaches (Table 2).

Detailed molecular analyses of the interactions within the tumor microenvironment in response to various immunotherapies will generate insights into therapeutically relevant targets (69, 70). Importantly, proper assessment of therapeutic efficacy or failure requires that serial tumor biopsies be obtained both before and after immune therapy despite their high costs and logistical challenges. With the recent striking progress in immune therapies for MCC, the diverse pipeline of agents, and forthcoming improvements in our ability to assess the tumor microenvironment, the future for MCC immunotherapy is very encouraging.

## Disclosure of Potential Conflicts of Interest

P. Nghiem reports receiving commercial research grants from Bristol-Myers Squibb and is a consultant/advisory board member for EMD Serono and Merck. No potential conflicts of interest were disclosed by the other authors.

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