PD-L1 Expression in the Merkel Cell Carcinoma Microenvironment: Association with Inflammation, Merkel Cell Polyomavirus, and Overall Survival


Abstract
Merkel cell carcinoma (MCC) is a lethal, virus-associated cancer that lacks effective therapies for advanced disease. Agents blocking the PD-1/PD-L1 pathway have shown objective, durable tumor regressions in patients with advanced solid malignancies and efficacy has been linked to PD-L1 expression in the tumor microenvironment. To investigate whether MCC might be a target for PD-1/PD-L1 blockade, we examined MCC PD-L1 expression, its association with tumor-infiltrating lymphocytes (TIL), Merkel cell polyomavirus (MCPyV), and overall survival. Sixty-seven MCC specimens from 49 patients were assessed with immunohistochemistry for PD-L1 expression by tumor cells and TILs, and immune infiltrates were characterized phenotypically. Tumor cell and TIL PD-L1 expression were observed in 49% and 55% of patients, respectively. In specimens with PD-L1 (+) tumor cells, 97% (28/29) showed a geographic association with immune infiltrates. Among specimens with moderate-severe TIL intensities, 100% (29/29) showed PD-L1 expression by tumor cells. Significant associations were also observed between the presence of MCPyV DNA, a brisk inflammatory response, and tumor cell PD-L1 expression: MCPyV(-) tumor cells were uniformly PD-L1(-). Taken together, these findings suggest that a local tumor-specific and potentially MCPyV-specific immune response drives tumor PD-L1 expression, similar to previous observations in melanoma and head and neck squamous cell carcinomas. In multivariate analyses, PD-L1(-) MCCs were independently associated with worse overall survival [HR 3.12; 95% confidence interval, 1.28–7.61; P = 0.012]. These findings suggest that an endogenous immune response promotes PD-L1 expression in the MCC microenvironment when MCPyV is present, and provide a rationale for investigating therapies blocking PD-1/PD-L1 for patients with MCC.

Cancer Immunol Res; 1(1); 54–63. ©2013 AACR.

Introduction
Merkel cell carcinoma (MCC) is an uncommon but aggressive cutaneous malignancy, showing both epithelial and neuroendocrine features. The majority (at least ~80%) of cases are associated with the oncogenic Merkel cell polyomavirus (MCPyV; refs. 1–3). MCC is diagnosed predominantly in the sun-exposed skin of older, fair-skinned individuals. An increased risk of developing MCC is observed in immunocompromised patients and those with other primary cancers (4–6).

MCC has historically been difficult to treat (7). It recurs following surgical excision of localized disease in about one-third of patients (8). While lymphadenectomy and adjuvant radiotherapy or chemotherapy may decrease the likelihood of recurrence, a significant percentage of patients develop metastatic disease, for which no therapy has been shown to prolong survival. Commonly used chemotherapy regimens for patients with metastatic or unresectable local disease include cisplatin or carboplatin plus etoposide, topotecan, or cyclophosphamide, doxorubicin (or epirubicin), and vincristine (CAV). Though MCC is often initially chemosensitive, resistance generally develops quickly and tumor regressions are rare. The 2-year survival rate for patients with American Joint Committee on Cancer (AJCC) stage IV disease is estimated to be 26% (9). Accordingly, new therapeutic options are sorely needed for patients with MCC in the adjuvant and advanced metastatic settings.

MCC, similar to other virus-associated malignancies, is highly immunogenic, and patients with MCPyV-positive tumors have improved overall survival compared with those with MCPyV-negative tumors (10). Oncoproteins such as MCPyV large and small T antigens and capsid proteins often stimulate the generation of antigen-specific T cells and...
antibodies in patients with MCC, and high serum antibody titers against viral capsid protein 1 (VP1) have been associated with improved progression-free survival (11, 12). T cells specific for viral antigens can be detected within MCC tumors and in the blood. Moreover, patients with MCC whose tumors harbor high numbers of CD3+, CD8+ tumor-infiltrating lymphocytes (TIL) show improved overall survival (13, 14). Conversely, immunocompromised patients have a higher than expected rate of MCC and display a higher MCC-specific death rate (15). These studies support an important role for the host immune system in containing MCC.

Notwithstanding the immunogenicity of MCC, the immune system often ultimately fails to control its growth. One potential explanation for this seeming contradiction is the expression of immune-inhibitory ligands in the tumor microenvironment such as programmed death ligand-1 (PD-L1, B7-H1, CD274). PD-L1 shares homology with B7.1 and B7.2, 2 members of the B7 immunoglobulin superfamily (16). The major receptor for PD-L1, namely programmed death-1 (PD-1), is expressed by activated T lymphocytes and delivers inhibitory signals upon PD-L1 ligation (17, 18). PD-L1 is physiologically expressed by both activated hematopoietic cells and inflamed epithelium as a means to downmodulate antigen-stimulated adaptive immune responses in peripheral tissues, thereby avoiding collateral damage to normal tissues (19). However, PD-L1 may be aberrantly expressed by tumor cells, creating a shield against immune attack (20). PD-L1 expression has been shown in many different human cancers, but whether MCC exploits this mechanism of immune resistance has yet to be established.

As new data emerge about immunotherapies targeting the PD-1/PD-L1 pathway, the potential expression of PD-L1 in the MCC tumor microenvironment is of increasing interest. Recent trials of nivolumab (BMS-936558, MDX-1106; Bristol–Myers Squibb), a blocking antibody against PD-1, and BMS-936559 (anti-PD-L1) have shown the power and durability of an immune-inhibitory system extending away from vessels; or (3), "severe" (either extending around the majority of the perimeter showing immunostaining was scored as 0–4%, 5–9%, and then at increasing 10% intervals by 2 independent pathologists (J.M. Taube and J.G. Vincent) who were blinded to clinical outcomes. Scoring differences greater than 10% were adjudicated. Cases showing at least 5% membranous expression of PD-L1 were considered positive. The pattern of PD-L1 expression in the tumor microenvironment was also assessed. PD-L1 expression was scored as absent (no PD-L1 expression by tumor) and no TILs or no PD-L1 expression by tumor cells, creating a shield against immune attack (20). PD-L1 expression has been shown in many different human cancers, but whether MCC exploits this mechanism of immune resistance has yet to be established.

As new data emerge about immunotherapies targeting the PD-1/PD-L1 pathway, the potential expression of PD-L1 in the MCC tumor microenvironment is of increasing interest. Recent trials of nivolumab (BMS-936558, MDX-1106; Bristol–Myers Squibb), a blocking antibody against PD-1, and BMS-936559 (anti-PD-L1) have shown the power and durability of an immune-inhibitory system extending away from vessels; or (3), "severe" (either extending around the majority of the perimeter showing immunostaining was scored as 0–4%, 5–9%, and then at increasing 10% intervals by 2 independent pathologists (J.M. Taube and J.G. Vincent) who were blinded to clinical outcomes. Scoring differences greater than 10% were adjudicated. Cases showing at least 5% membranous expression of PD-L1 were considered positive. The pattern of PD-L1 expression in the tumor microenvironment was also assessed. PD-L1 expression was scored as absent (no PD-L1 expression by tumor) and no TILs or no PD-L1 expression by TILs, present on both tumor cells and TILs in immediate proximities, or present on TILs but not on tumor cells.

Characterization of immune infiltrates

The recommended surgical pathology scoring system for TILs in MCC specimens applies to primary lesions (28), thus a modified scoring system was adopted. The degree of CD3+ TILs and associated histiocytes was scored as (0), none; (1), "mild" (rare intratumoral cells, mostly perivascular); (2), "moderate" (locally present at periphery of tumor and/or intratumoral extending away from vessels); or (3), "severe" (either extending around the majority of the perimeter of the tumor deposit, or diffusely present throughout tumor; refs. 27, 29). As Merkel cells and lymphocytes can sometimes be difficult to differentiate on hematoxylin/eosin (H&E) staining alone, a CD3 immunostain was used to aid in assigning TIL score. When PD-L1 expression by lymphohistiocytic infiltrates was observed along the periphery of tumor, the proportion of the perimeter showing immunostaining was scored as 0–4%, 5–9%,
10%, and then at 10% intervals up to 100%. In addition, intratumoral and peritumoral CD8+ T cells were scored as "low" or "high" density with a cut point of approximately 60 CD8+ cells per typical high power field (HPF; ref. 13). Finally, the CD4:CD8 ratio in intratumoral and peritumoral TILs was estimated as 4:1, 3:1, 2:1, 1:1, 1:2, 1:3, or 1:4.

Detection of merkel cell polyomavirus

The presence of MCPyV in tumor samples was detected by quantitative real-time PCR (qPCR) conducted on genomic DNA extracted from FFPE tissues, as published (30, 31). Ten consecutive 10-μm thick unstained sections were cut from a tumor focus of approximately 2 mm diameter using the Pinpoint Slide DNA Isolation System (Zymo Research) according to the manufacturer’s protocol. After proteinase K digestion, genomic DNA was purified using the QiAamp DNA Mini Kit (Qiagen), according to the manufacturer’s “Blood and Body Fluid Spin” protocol. MCPyV was detected with qPCR for the viral capsid protein 1 (VP1) and viral T antigen 3 (LT3) genes (30, 31). The LT3 primers localize to the first exon and thus detect both large and small T antigens. qPCR was conducted in triplicate for each sample, and the β-actin gene was used as a normalizer for gene abundance. Molecular grade water was used as a nontemplate control. Standard curves were developed by diluting the MCPyV-positive cell line MKL-1. In those instances where only one of the 2 viral genes VP1 or the LT3 was detected, the samples were considered positive for MCPyV. The lowest value we reliably detected was a MCPyV to β-actin ratio of 0.09, which is in keeping with assays reported by others (14, 32).

Statistical analysis

Associations of PD-L1 expression with clinicopathologic features were evaluated with Fisher exact test and Wilcoxon–Mann–Whitney tests. For patients with multiple tumor samples, the samples were treated as independent observations for the clinicopathologic associations. Overall survival was calculated from the date of the original diagnostic biopsy to the date of last follow-up or death with the Kaplan–Meier method and compared with the log-rank test. Multivariate survival analyses were conducted using the Cox proportional hazards model. Survival analyses for patients with multiple specimens were conducted using the sample that had the highest value for the parameter in question. Statistical analyses were conducted using the R statistical package (version 2.15.1). All tests were 2-sided and P values < 0.05 were considered significant.

Results

Patient characteristics and tumor specimens

To assess factors associated with PD-L1 expression in the tumor microenvironment, we studied 49 patients with MCC whose clinicopathologic characteristics are summarized in Table 1. Patients were predominantly male (65%) with a median age of 65 years. Forty-nine percent of patients presented with metastatic disease (AJCC stage III–IV), whereas 51% had localized disease (AJCC stage I–II; ref. 26). Overall,

<table>
<thead>
<tr>
<th>Table 1. Relationship of PD-L1 expression by tumor cells and infiltrating immune cells to patient demographics and clinicopathologic features</th>
</tr>
</thead>
<tbody>
<tr>
<td><strong>All patients</strong></td>
</tr>
<tr>
<td><strong>n = 49</strong></td>
</tr>
<tr>
<td><strong>Gender, n (%)</strong></td>
</tr>
<tr>
<td>Female</td>
</tr>
<tr>
<td>Male</td>
</tr>
<tr>
<td><strong>Age at primary diagnosis</strong></td>
</tr>
<tr>
<td>Mean (SD)</td>
</tr>
<tr>
<td>Median (range)</td>
</tr>
<tr>
<td><strong>Pathologic stage at diagnosis, n (%)</strong></td>
</tr>
<tr>
<td>IA</td>
</tr>
<tr>
<td>IB</td>
</tr>
<tr>
<td>IIA</td>
</tr>
<tr>
<td>IIC</td>
</tr>
<tr>
<td>IIIA</td>
</tr>
<tr>
<td>IIIB</td>
</tr>
<tr>
<td>IV</td>
</tr>
</tbody>
</table>

* Tumor cells or associated infiltrating immune cells were considered PD-L1+ if ≥5% of cells had membranous (cell surface) PD-L1 expression. In this table, patients with multiple specimens are considered PD-L1+ if any specimen was positive.

49% of patients had PD-L1+ tumor cells, and 55% had PD-L1+ infiltrating immune cells, in at least one tumor specimen examined. Expression of PD-L1 in the tumor microenvironment, by either tumor cells or infiltrating immune cells, did not correlate with patient gender, age, or pathologic stage at the time of diagnosis. The median follow-up time for the total cohort of 49 patients was 50 months (range 4–142 months), with a median follow-up of 25 months (range 4–127 months) for those who died and 87 months (range 15–142) for patients still alive. At the end of the follow-up period, 23 patients had died, 13 had disease progression but were still alive, and 13 had no evidence of disease.

Sixty-seven individual tumor specimens were available from 49 patients for assessment of PD-L1 and CD3 expression by IHC (39 patients had a single specimen and 10 patients each had 2–5 specimens). Forty and 39 specimens from 34 patients had both TILs present and tissue available for additional characterization of immune cell subsets by IHC, respectively. Forty-two specimens from 32 patients yielded sufficient material for MCPyV DNA detection by qPCR. For the purpose of the survival analysis on those patients with multiple tumor samples, the patient was considered “positive” for a given parameter if any tumor specimen from that patient was positive.

**PD-L1 expression in the MCC microenvironment**

Previous studies of melanomas and of squamous cell carcinomas of the head and neck (HNSCC) have shown that PD-L1 can be expressed by multiple components of the tumor microenvironment, including tumor cells themselves and infiltrating immune cells (lymphocytes and macrophages/histiocytes; refs. 27, 29). Theoretically, PD-L1 expression by any cell type could exert local immunosuppressive effects, although previous reports suggest that tumor-expressed membrane PD-L1 correlates with response to anti-PD-1 therapy (22). In the current study of MCC, membranous PD-L1 expression by tumor cells was observed in 29 of 67 specimens (43%). Among 29 specimens containing PD-L1+ tumor cells, 100% harbored immune infiltrates, in contrast to only 47% of 38 specimens with PD-L1(-) tumor cells (P < 0.001; Table 2). All but one specimen harboring PD-L1+ tumor cells was characterized by a geographic colocalization of these cells with infiltrating immune cells (Fig. 1); in a single specimen, tumor cell PD-L1 expression was geographically remote from a moderate immune infiltrate (Fig. 2). When observed in association with TILs, focal tumor expression of PD-L1 was seen in ≤10% of the total tumor area represented on the slide (mean = 5%, SD ± 0.94; median 5%, range 5%–10%); expression was located

<table>
<thead>
<tr>
<th>Tumor site</th>
<th>n</th>
<th>TIL(−)</th>
<th>TIL+</th>
<th>TIL(−)</th>
<th>TIL+</th>
</tr>
</thead>
<tbody>
<tr>
<td>Primary</td>
<td>29</td>
<td>8 (50)</td>
<td>8 (50)</td>
<td>0 (0)</td>
<td>13 (100)</td>
</tr>
<tr>
<td>Metastasis</td>
<td>38</td>
<td>12 (55)</td>
<td>10 (45)</td>
<td>0 (0)</td>
<td>16 (100)</td>
</tr>
<tr>
<td>All</td>
<td>67</td>
<td>20 (53)</td>
<td>18 (47)</td>
<td>0 (0)</td>
<td>29 (100)</td>
</tr>
</tbody>
</table>

* Tumor cells were considered PD-L1+ if ≥5% of cells had membranous (cell surface) PD-L1 expression.

* TILs were considered positive if infiltrates were mild, moderate, or severe (scores 1–3; see Materials and Methods).

* Fisher exact test.
Neither tumor nor TIL PD-L1 expression correlated significantly with tumor site (primary lesion versus metastasis). The intensity of the immune infiltrate was characterized by only a mild infiltrate (score 1) to severe (score 3) (not shown; P < 0.001). Similarly, when TIL intensity was studied in relation to TIL PD-L1 expression, PD-L1 expression was significantly associated with increasing intensity of the immune infiltrate (not shown; P < 0.001). Neither tumor nor TIL PD-L1 expression correlated significantly with tumor site (primary lesion versus metastasis).

### Table 3. Correlation of PD-L1 expression by tumor cells with PD-L1 expression by adjacent immune infiltrates in 67 MCC specimens

<table>
<thead>
<tr>
<th>Tumor site</th>
<th>Tumor PD-L1(−), n (%)</th>
<th>TIL PD-L1(−)</th>
<th>TIL PD-L1+</th>
<th>Tumor PD-L1(+), n (%)&lt;sup&gt;a&lt;/sup&gt;</th>
<th>TIL PD-L1(−)</th>
<th>TIL PD-L1+</th>
<th>P value&lt;sup&gt;b&lt;/sup&gt;</th>
</tr>
</thead>
<tbody>
<tr>
<td>Primary</td>
<td>29</td>
<td>12 (75)</td>
<td>4 (25)</td>
<td>0 (0)</td>
<td>13 (100)</td>
<td></td>
<td>&lt;0.001</td>
</tr>
<tr>
<td>Metastasis</td>
<td>38</td>
<td>21 (95)</td>
<td>1 (5)</td>
<td>0 (0)</td>
<td>16 (100)</td>
<td></td>
<td>&lt;0.001</td>
</tr>
<tr>
<td>All</td>
<td>67</td>
<td>33 (87)</td>
<td>5 (13)</td>
<td>0 (0)</td>
<td>29 (100)</td>
<td></td>
<td>&lt;0.001</td>
</tr>
</tbody>
</table>

<sup>a</sup>Tumor cells or associated infiltrating immune cells were considered PD-L1+ if ≥5% of cells had membranous (cell surface) PD-L1 expression.

<sup>b</sup>Fisher exact test.

In 38 specimens that were negative for tumor cell PD-L1 expression, 18 (47%) contained TILs (Table 2) but were characterized by only a mild infiltrate (score 1). In contrast, among 29 specimens with PD-L1+ tumor cells, all were associated with TILs, and TIL intensities were moderate (score 2) to severe (score 3) (not shown; P < 0.001). Similarly, when TIL intensity was studied in relation to TIL PD-L1 expression, PD-L1 expression was significantly associated with increasing intensity of the immune infiltrate (not shown; P < 0.001). Neither tumor nor TIL PD-L1 expression correlated significantly with tumor site (primary lesion versus metastasis).

### Characterization of infiltrating immune cells related to PD-L1 expression

TILs were further characterized according to CD4+ and CD8+ subsets, and the association of the CD4:CD8 ratio and the CD8 density with PD-L1 expression by tumor and by TILs was studied in both the intratumoral and peritumoral compartments (Table 4). CD4+ T cells were present at equal or greater numbers than CD8+ T cells in peritumoral immune infiltrates in 35 of 39 (90%) specimens, while the inverse was seen in intratumoral infiltrates, with CD8+ T cells predominating in 23 of 39 (59%) of specimens. PD-L1 expression by both tumor cells and TILs was strongly associated with an intratumoral CD8-predominant TIL phenotype (P = 0.001 and P = 0.003, respectively); in contrast, there was no significant difference between the composition of peritumoral immune infiltrates in PD-L1+ versus PD-L1− tumors. When CD8+ T-cell density was enumerated in the intratumoral and peritumoral compartments, a high density (>60 per HPF) of CD8+ cells was associated with PD-L1 expression by tumor cells (P = 0.015 and P < 0.001, respectively) and by TILs (P = 0.043 and P < 0.001, respectively). Taken together, these findings are compatible with a role for CD8+ T cells in mediating PD-L1 expression in the MCC microenvironment, and are consistent with our previous findings in melanoma and HNSCC (27, 29).

### Association of viral status with immune infiltrates and PD-L1 expression

Among 42 MCC specimens available for detection of MCPyV, 34 (81%) were positive. There were 6 patients who had multiple specimens available for viral testing, and there was concordance in viral status among specimens from individual patients. Specifically, 2 patients had a primary tumor and 2 metastases for analysis, all of which were positive. Two patients had a primary tumor and one metastasis for analysis, both of which were positive in one patient and both of which were negative in the other patient. Two patients had only metastases available, one with 4 different lesions and one with 2 different lesions, all of which were positive for virus. When analyzed by patient, 78% (25/32) of patients had MCPyV-positive tumors, which is in keeping with the reported prevalence rates of approximately 80%−90% (1, 30, 31, 33).

The association between the presence of MCPyV, inflammatory infiltrates and PD-L1 expression was assessed...
There was a significant association of the presence of virus with tumor PD-L1 expression, and the presence of virus with a brisk (moderate or severe) inflammatory response: 50% of MCPyV+ specimens were PD-L1+ and had a moderate-severe immune infiltrate, compared with 0% of MCPyV(-) specimens. The association of TILs with MCPyV+ tumors is reminiscent of findings from Sihto and colleagues (14). Similarly, HPV+ HNSCC are more likely to be inflamed than their HPV(-) counterparts (29). These findings suggest the hypothesis that an immune response to viral antigens creates a local proinflammatory environment driving tumor PD-L1 expression. PD-L1 expression by the inflammatory cell population alone did not associate significantly with the presence of virus.

### Table 4. Relationship of TIL subsets to tumor and TIL PD-L1 expression

<table>
<thead>
<tr>
<th></th>
<th>All Specimens</th>
<th>Tumor</th>
<th>TILs</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>PD-L1(-)</td>
<td>PD-L1+</td>
<td>n=15</td>
</tr>
<tr>
<td>Intratumoral CD4:CD8, n (%)</td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>CD4 ≥ CD8</td>
<td>16 (41)</td>
<td>12 (80)</td>
<td>4 (17)</td>
</tr>
<tr>
<td>CD8 &gt; CD4</td>
<td>23 (59)</td>
<td>3 (20)</td>
<td>20 (83)</td>
</tr>
<tr>
<td>Peritumoral CD4:CD8, n (%)</td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>CD4 ≥ CD8</td>
<td>35 (90)</td>
<td>15 (100)</td>
<td>20 (83)</td>
</tr>
<tr>
<td>CD8 &gt; CD4</td>
<td>4 (10)</td>
<td>0 (0)</td>
<td>4 (17)</td>
</tr>
</tbody>
</table>

**Table 5. Viral status related to TIL intensity and PD-L1 expression by tumor cells and TILs**

<table>
<thead>
<tr>
<th></th>
<th>All specimens, n=42</th>
<th>MCPyV(-), n=8</th>
<th>MCPyV(+), n=34</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>n (%)</td>
<td>n (%)</td>
<td>n (%)</td>
</tr>
<tr>
<td>Tumor PD-L1(-)</td>
<td>25 (60)</td>
<td>8 (100)</td>
<td>17 (50)</td>
</tr>
<tr>
<td>Tumor PD-L1+</td>
<td>17 (40)</td>
<td>0 (0)</td>
<td>17 (50)</td>
</tr>
<tr>
<td>TIL score 0,1,a</td>
<td>25 (60)</td>
<td>8 (100)</td>
<td>17 (50)</td>
</tr>
<tr>
<td>TIL score 2,3</td>
<td>17 (40)</td>
<td>0 (0)</td>
<td>17 (50)</td>
</tr>
<tr>
<td>TIL PD-L1(-)</td>
<td>21 (50)</td>
<td>6 (75)</td>
<td>15 (44)</td>
</tr>
<tr>
<td>TIL PD-L1+</td>
<td>21 (50)</td>
<td>2 (25)</td>
<td>19 (56)</td>
</tr>
</tbody>
</table>

*aTILs (lymphocytes and associated histiocytes) were scored as (0), none; (1), "mild"; (2), "moderate"; or (3), "severe" as described in Materials and Methods.

**Association of PD-L1 expression in the MCC microenvironment with overall survival**

PD-L1 expression in the tumor microenvironment seems to reflect antitumor immunity and adaptive immune resistance by the tumor, creating a negative feedback loop to mediate T-cell inhibition (27). Negative associations of tumor PD-L1 expression with prognosis have been reported for several cancers (34–36). However, we have reported a positive correlation of PD-L1 expression with improved overall survival in patients with metastatic melanoma (27). Thus, the association of both tumor cell and TIL PD-L1 expression with survival in 49 patients with MCC was assessed in the current study. By univariate analysis, there was a significant association of tumor
cell PD-L1 expression ($P = 0.046$), but not TIL PD-L1 expression, with improved overall survival (Fig. 3). To further analyze this association, we used a Cox multivariate proportional hazards model and controlled for age and whether the specimen studied was a primary versus metastatic lesion. The failure of MCC to express PD-L1 was independently associated with worse overall patient survival by multivariable analysis (HR = 3.12; 95% confidence interval: 1.28–7.61, $P = 0.012$).

Discussion

In 2008, Feng and colleagues reported the association of MCC with a previously undescribed virus, now called MCPyV (1). Further studies have shown that MCPyV is directly carcinogenic; that is, viral DNA is integrated into the tumor genome and the cells that undergo malignant transformation are the same cells infected by the virus (2). Consequently, neoantigens in the form of oncoproteins are expressed by tumor cells, creating an opportunity for immune recognition and tumor elimination. Previous studies indicate that the natural history of MCC is impacted by the resultant host immune response. However, approximately 10%–20% of MCCs are generally not reported to be associated with MCPyV infection, dividing MCC into 2 groups that may differ in the tumor–host immune system interaction. Our results support the notion that the immune microenvironment differs between virus positive and virus negative MCC, as the latter does not express PD-L1, a major component of the PD-1 pathway that downmodulates the amplitude of T-cell activation within the tumor. In contrast, we find that roughly half of virus-positive MCCs express the immunosuppressive ligand PD-L1 on multiple cell types (tumor cells, lymphocytes, macrophages/histiocytes) in the tumor microenvironment. Furthermore, we also show features of geographic colocalization of tumor cell PD-L1 expression with TILs, similar to our previous studies of melanoma and HNSCC, suggesting that cytokine secretion by tumor-reactive immune cells may promote PD-L1 expression to create a negative feedback loop protecting tumor cells from immune destruction. These findings provide a rationale for future studies evaluating the potential role of PD-1/PD-L1 immune checkpoint blockade therapy in treating MCC.

The natural history of malignancies linked with viral infections, which are frequently associated with chronic inflammation, is often influenced by the immunologic state of the host. The results of the current study contribute to the emerging theme that the immune regulatory PD-L1/PD-1 axis may play a significant role in the development and progression of virus-associated cancers. Intratumoral expression of PD-L1 and/or PD-1 has been shown in human papillomavirus (HPV)-associated cancers, hepatocellular carcinoma, Epstein–Barr virus-associated nasopharyngeal cancer, and adult T-cell leukemia/lymphoma. The immune regulatory PD-L1/PD-1 axis may play a significant role in the development and progression of virus-associated cancers. Intratumoral expression of PD-L1 and/or PD-1 has been shown in human papillomavirus (HPV)-associated cancers, hepatocellular carcinoma, Epstein–Barr virus-associated nasopharyngeal cancer, and adult T-cell leukemia/lymphoma linked to human T-cell leukemia virus-1. In each of these cancers, the PD-L1/PD-1 axis is thought to be upregulated as a part of host’s innate and adaptive immune response to the presence of virus, yet expression of these molecules enables tumor evasion from cytotoxic T-cell attack. However, the balance between immune activation and tolerance may depend on tumor type. For example, in this study, we showed an association between MCPyV DNA, a brisk immune infiltrate, local expression of PD-L1, and improved patient survival, indicating a degree of MCC containment by the host immune system. We postulate that high lymphocyte infiltration and consequent PD-L1 upregulation in a subset of virus-positive tumors reflects both enhanced antigenicity associated with viral infection and virus-independent factors, such as E-selectin expression on vascular endothelium that promotes lymphocyte egress (41). Of note, Rodig and colleagues, in a recently published study using ultrasensitive IHC and quantitative molecular techniques, showed that MCPyV DNA was detectable in almost all MCC specimens analyzed (3). It may be that integration of MCPyV DNA into host DNA, and subsequent expression of oncogenic viral proteins, beyond a certain threshold is required for immune activation of the tumor microenvironment.

Increased levels of PD-1+ TILs cells have recently been correlated with a favorable clinical outcome in HNSCC (37). In contrast, in patients with HBV+ hepatocellular carcinoma, high levels of circulating PD-1+ and PD-1+ cells were associated with a worse prognosis (38). In virus-associated malignancies, PD-L1 expression likely reflects both a host reaction to the tumor and the chronic inflammatory environment triggered by the presence of virus. Because PD-L1/PD-1 checkpoint blockade has been shown to reverse virus-specific T-cell exhaustion and facilitate virus clearance (42, 43), it is an attractive therapeutic option not only for virus-associated malignancies, but also for treating the underlying infection.
As more knowledge is gained about the expression patterns of PD-L1 in various human cancers, it seems that different cancers may show distinct characteristics reflecting the cell type or anatomic site of origin, the specific inflammatory trigger (e.g., virus or tumor neoantigens), the resultant cytokine milieu, cellular subpopulations dominating the immune infiltrate, and/or activation of aberrant molecular signaling pathways. For instance, in the current study, PD-L1 expression was predominantly seen in the associated lymphohistiocytic infiltrates and was focal when observed in the tumor cells. We have also observed cases of colorectal carcinoma with prominent PD-L1 expression by tumor-associated macrophages and minimal tumor cell surface expression (23). In contrast, as previously reported in melanoma, PD-L1 expression was seen in both the tumor and infiltrating immune cells, in more equal proportions (27). We have reported PD-L1+ tumor cells to be geographically associated with TILs in the majority of MCCs, melanomas, and HPV+ HNSCCs; however, in singular cases, we observed PD-L1 expression out of proportion to infiltrating immune cells, suggesting oncogene-driven expression as described in association with PTEN loss in glioblastoma multiforme (44) and in HPV(-) HNSCC (29). Separate models of innate (constitutive) and adaptive tumor immune resistance via PD-L1 expression have been proposed (45). These are not mutually exclusive; it is possible that both of these mechanisms can be operative independently or concordantly, and their relative contributions may vary by tumor type. Studies investigating the interplay between oncogenic-driver pathways in different tumor types, PD-L1 expression, and different populations of infiltrating immune cells are currently underway in our laboratories.

In both melanoma and HNSCC, IFN-γ has been highlighted as a major cytokine driving PD-L1 expression (27, 29). A number of other factors, including LAG-3, II-17, II-10, II-6, IL-21 and other common gamma chain cytokines have also been identified in association with PD-L1 expression (46–49). Although we did not directly investigate soluble factors associated with PD-L1 expression in MCC, we anticipate that IFN-γ plays a strong role. MCC-specific TILs have previously been shown to secrete IFN-γ (11), and Nghiem and colleagues also showed that in patients with MCC with better prognosis, tumors contained CD8+ lymphocytic infiltrates and overexpressed IFN-γ mRNA (13). Consistent with these findings, when tumor cell PD-L1 expression was observed in our cohort, it was in geographic association with CD8+ TILs, and it was shown to be a positive prognostic factor.

In summary, a growing body of evidence suggests that the natural history of MCC is significantly impacted by the immunologic state of the host. Here, we show that PD-L1 expression by MCC is an independent prognostic factor predicting improved overall survival in a cohort of patients from a single institution. The results of our study indicate that MCC tumor cells and the surrounding microenvironment display elements of immune activation that may be augmented with immune checkpoint blocking agents such as anti-PD-1 and anti-PD-L1, and provide a rationale for exploring these therapies in a tumor type that currently lacks effective treatment options for advanced disease.

Disclosure of Potential Conflicts of Interest
H. Xu has a commercial research grant from Bristol-Myers Squibb. R.A. Anders has a commercial research grant from Bristol-Myers Squibb. S.J. Topalian has a commercial research grant from Bristol-Myers Squibb and is a consultant/advisory board member of Bristol-Myers Squibb, Jounce Therapeutics, and Amplimmune Inc. J.M. Taube has a commercial research grant and is a consultant/advisory board member of Bristol-Myers Squibb. No potential conflicts of interest were disclosed by the other authors.

Authors’ Contributions
Acquisition of data (provided animals, acquired and managed patients, provided facilities, etc.): E.J. Lipson, J.G. Vincent, M. Loyo, H. Xu, S.K. Nayar, R.A. Anders, J.M. Taube
Administrative, technical, or material support (i.e., reporting or organizing data, constructing databases): E.J. Lipson, J.G. Vincent, H. Xu, J.M. Taube
Study supervision: E.J. Lipson, S.L. Topalian, J.M. Taube

Acknowledgments
The authors thank Dr. Drew Pardoll (Johns Hopkins University School of Medicine) for critical review of the manuscript, insights, and support. The authors also thank Dr. Leping Chen (Yale University School of Medicine) for helpful discussions and for providing the anti-PD-L1 monoclonal antibody 5H1.

Grant Support
This work was supported by the NIH (R01 CA142779, to J.M. Taube and S.L. Topalian; R01DK081417 and R01DK080736, to R.A. Anders; and P30 CA069637, to R.S. Lubet, H. Wang, D. Sidransky, R.A. Anders, and S.L. Topalian), the Barney Family Foundation (to E.J. Lipson, J.M. Taube, S.L. Topalian), the Laverna Hahn Charitable Trust (to E.J. Lipson, J.M. Taube, S.L. Topalian), and the Commonwealth Foundation (to J.M. Taube, H. Xu).

The costs of publication of this article were defrayed in part by the payment of page charges. This article must therefore be hereby marked advertisement in accordance with 18 U.S.C. Section 1734 solely to indicate this fact.

Received February 20, 2013; revised April 2, 2013; accepted April 11, 2013; published OnlineFirst May 21, 2013.

References


Cancer Immunology Research


Cancer Immunology Research


PD-L1 Expression in the Merkel Cell Carcinoma Microenvironment: Association with Inflammation, Merkel Cell Polyomavirus, and Overall Survival

Evan J. Lipson, Jeremy G. Vincent, Myriam Loyo, et al.


Updated version
Access the most recent version of this article at:
doi:10.1158/2326-6066.CIR-13-0034

Cited articles
This article cites 46 articles, 16 of which you can access for free at:
http://cancerimmunolres.aacrjournals.org/content/1/1/54.full#ref-list-1

Citing articles
This article has been cited by 16 HighWire-hosted articles. Access the articles at:
http://cancerimmunolres.aacrjournals.org/content/1/1/54.full#related-urls

E-mail alerts
Sign up to receive free email-alerts related to this article or journal.

Reprints and Subscriptions
To order reprints of this article or to subscribe to the journal, contact the AACR Publications Department at pubs@aacr.org.

Permissions
To request permission to re-use all or part of this article, contact the AACR Publications Department at permissions@aacr.org.