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# Early Neoplastic Progression Is Complement Independent

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## Abstract

Infiltration of leukocytes into premalignant tissue is a common feature of many epithelial neoplasms and is thought to contribute to cancer development. However, the molecular and cellular regulatory mechanisms underlying activation of innate host responses to enhanced neoplastic cell proliferation are largely unknown. Considering the importance of the complement system in regulating inflammation during acute pathologic tissue remodeling, we examined the functional significance of complement component 3 (C3) as a regulator of inflammatory cell infiltration and activation during malignant progression by using a transgenic mouse model of multistage epithelial carcinogenesis, e.g., HPV16 mice. Whereas abundant deposition of C3 is a characteristic feature of premalignant hyperplasias and dysplasias coincident with leukocyte infiltration in neoplastic tissue, genetic elimination of C3 neither affects inflammatory cell recruitment toward neoplastic skin nor impacts responding pathways downstream of inflammatory cell activation, e.g., keratinocyte hyperproliferation or angiogenesis. Taken together, these data suggest that complementindependent pathways are critical for leukocyte recruitment into neoplastic tissue and leukocytemediated potentiation of tumorigenesis.

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#### Introduction

Tumors are composed of genetically altered cells, a variety of responding host cells, including fibroblasts, vascular cells (endothelial and mural cells), and a diverse array of inflammatory leukocytes embedded in a dynamic extracellular matrix (ECM) microenvironment [1,2]. Accumulating clinical and experimental data suggest that inflammatory leukocytes provide essential regulatory roles for developing tumors where their innate activities are conscripted by neoplastic cells to make significant contributions to malignant phenotypes [1–3]. The significance of inflammation associated with cancer development is underscored by the realization that tissues afflicted with chronic inflammatory states, e.g., Crohn's disease, ulcerative colitis, pancreatitis, etc., exhibit increased risk for malignancy [4-7]. Moreover, long-term users of anti-inflammatory drugs demonstrate reduced cancer risk [8-14]. Taken together, it seems reasonable to suggest that therapeutic strategies targeting inflammatory cell recruitment, activation, or response represent tractable anticancer opportunities.

Complement activation is a central event during innate immune defense after pathogenic tissue assault [15,16]. Three different pathways of complement activation have been identified, namely, the classic, the alternative, and the lectin pathways [17–19]. Complement component C3 is a central protein of the complement cascade, expression of which is essential for activation of all three complement pathways [19]. Foreign antigens and immune complexes activate the proteolytic complement activation cascade, resulting in formation of lytic membrane attack complexes (MAC) [17-19] and formation and liberation of anaphylatoxins, e.g., C3a and C5a, potent proinflammatory factors that induce recruitment and activation of various inflammatory cells, in particular mast cells, eosinophils, and neutrophils [20-23]. Deposition of complement component proteins is a common occurrence at sites of inflammation [24-28]. Studies utilizing complement-depleted mice and C3- or C5-deficient mice have identified crucial roles for complement in mast cell recruitment and activation during airway hyperresponsiveness [25,29,30], IgG-antigen complex-mediated inflammation [31], intestinal ischemia/reperfusion [32], delayed-type hypersensitivity [33], and subepidermal blistering disease [34], thus underscoring the central role complement plays in regulating disease pathogenesis.

Using a transgenic mouse model of multistage epithelial carcinogenesis, e.g., HPV16 mice [35], we have investigated the functional role of complement component 3 (C3) as a regulator of inflammatory cell recruitment and premalignant progression. HPV16 mice express the early region genes of human papillomavirus type 16 (HPV16) as transgenes under control of the human keratin 14 promotor/enhancer [36]. By

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Abbreviations: BrdU, bromodeoxyuridine; CAE, chloroacetate esterase; C3, complement component 3; IF, immunofluorescence; HPV16, human papillomavirus 16

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1 month of age, HPV16 mice develop epidermal hyperplasias with 100% penetrance that are characterized by a terminally differentiating hyperproliferative epidermis atop a dermal compartment rich in recruited activated mast cells [37]. Hyperplastic lesions advance focally into angiogenic dysplasias between 3 and 6 months of age and are distinct from hyperplasias based on the prominent hyperproliferative epidermis that fails to undergo terminal differentiation and a dermis where intense mast cell and neutrophil infiltration occurs proximal to dilated and enlarged angiogenic vasculature [37-39]. By 1 year of age, 60% of HPV16 mice develop malignant skin carcinomas, 50% of which are squamous cell carcinomas (SCCs) that metastasize to regional lymph nodes with an  $\sim 30\%$  frequency, and  $\sim 10\%$  of which represent nonmetastatic microcystic adnexal carcinomas (MACs) [38,40]. Mast cell deficiency in HPV16 mice significantly attenuates keratinocyte hyperproliferation and activation of angiogenesis, thus supporting the concept that inflammation, and in particular activation of sentinel mast cells and release of their factors, functionally contribute to early neoplastic development in HPV16 mice [37].

In the present study, we report abundant deposition of C3 in neoplastic skin of HPV16 mice. To determine if C3 deposition represented a functionally significant parameter of mast cell recruitment, activation, or response, we generated HPV16 mice harboring a homozygous null mutation in the C3 gene, e.g., HPV16/C3<sup>-/-</sup> mice [15,16]. Despite abundant C3 deposition into the neoplastic microenviroment of HPV16 mice, our data suggest that activation of the complement system does not functionally contribute to recruitment of inflammatory cells, induction of keratinocyte hyperproliferation, or activation of angiogenesis during epithelial neoplastic progression. Because abundant deposition of IgG antibodies is an early event that is sustained throughout neoplastic progression and coincides with inflammatory cell infiltration, we propose an antibody-dependent but complement-independent pathway regulating mast cell recruitment into neoplastic skin.

## Materials and Methods

## Transgenic Mice

Generation and characterization of HPV16 transgenic mice and neoplastic staging based on keratin intermediate filament expression has been described previously [35,36]. The generation of C3–deficient mice has been described [15,16].  $C3^{-/-}$  mice were kindly provided by Dr. M. C. Carroll (Harvard Medical School, Boston, MA). To generate HPV16 mice in the C3-deficient backgrounds,  $C3^{+/-}$  mice were individually backcrossed into the FVB/n strain to N19, where they were then intercrossed with HPV16 mice to generate a breeding colony of HPV16/C3<sup>-/-</sup> and C3<sup>-/-</sup> animals. All mice were maintained within the University of California, San Francisco, Laboratory for Animal Care facility according to IACUC procedures. Genotyping of C3<sup>-/-</sup> mice was based on a C3 ELISA where C3 levels in sera (or tissue) were assayed using a polyclonal goat anti–mouse C3 antiserum

(1:500; ICN/Cappel, Irvine, CA). Because the C3 locus is located near the MHC region, we confirmed H- $2K^q$  expression and absence of H- $2D^b$  expression after backcrossing into the FVB/n strain by FACS analysis using mouse anti-mouse H- $2D^b$  and mouse anti-mouse H- $2K^q$  mAbs (BD Biosciences, San Diego, CA).

## Immunofluorescence Detection of Antibody and C3 Depositions

Age-matched tissue samples from transgenic and control animals were frozen directly in glycerol-based freezing medium (OCT). For immunofluorescence (IF) detection of antibody and C3 deposition, 10- $\mu$ m OCT-embedded tissue sections were cut with a Leica CM1900 cryostat. Sections were air-dried, fixed in acetone for 5 minutes, and subjected to IF staining as follows: sections were blocked for 30 minutes in blocking buffer (5% goat serum/2.5% BSA/PBS). For IF detection of antibodies, a 1:50 dilution of rat anti-mouse CD16/CD32 mAb (BD Biosciences) was added to the blocking buffer to prevent nonspecific antibody binding. Fluorescein isothiocyanate (FITC)-conjugated goat anti-mouse IgG (v-chain specific, 1:100; Sigma, St. Louis, MO), and rat anti-mouse C3 (1:100; Caltag, Burlingame, CA) were diluted in 0.5  $\times$  blocking buffer and incubated with tissue sections for 1 hour at room temperature, protected from light. Sections were then extensively washed with PBS. For IF detection of C3, sections were blocked for 5 minutes in blocking buffer and incubated with FITC-conjugated goat anti-rat (1:100, Jackson ImmunoResearch, West Grove, PA) for 45 minutes at room temperature, protected from light. Slides were then extensively washed with PBS before mounting using Vectashield containing 4',6 diamidino-2phenylindole (DAPI) (Vector, Burlingame, CA). All IF experiments included negative controls for determination of background staining, which was negligible. Data shown are representative of results obtained after examination of tissues removed from a minimum of four mice per group.

## Immuno- and Enzyme Histochemistry

Age-matched tissue samples from transgenic and control animals were immersion-fixed in 10% neutral-buffered formalin followed by dehydration through graded alcohols and xylene, and embedded in paraffin. Five-micrometer-thick paraffin sections were cut using a Leica 2135 microtome. Sections were deparaffinized and subjected to immunohistochemical detection as previously described [41]. Primary antibodies were diluted in blocking buffer (5% goat serum/ 2.5% BSA/PBS) at 1:50 dilution for rat anti-mouse CD31 (BD Biosciences) and 1:500 for rat anti-mouse neutrophilspecific primary antibody (Cedarlane Labs, Hornby, Ontario, Canada). Sections were incubated with primary antibody for 2 to 4 hours at room temperature, followed by PBS washing, brief (5 minutes) incubation in blocking buffer, and subsequent incubation with biotinylated secondary antibody (rabbit anti-rat IgG 1:200, Vector) for 45 minutes at room temperature. After PBS washing, Vectastain Elite ABC reagent (Vector) was applied for 30 minutes. Sections were then washed in PBS and endogenous peroxidase activity blocked

by incubation in 30% H<sub>2</sub>O<sub>2</sub> in methanol for 20 minutes. Antibodies were visualized by treatment with Fast 3,3'diaminobenzodine (Sigma), dehydrated in graded alcohols (70%, 95%, and 100% ethanol), and mounted in Cytoseal 60 (Richard-Allan Scientific, Kalamazoo, MI).

Chloroacetate esterase (CAE) histochemistry was performed to visualize serine esterase activity in mast cells as previously described [37]. Briefly, 1 mg of naphthol AS-D chloroacetate (Sigma) was dissolved in 20 µl of N,N-dimethyl formamide (DMF) and 1 ml buffer (8% DMF, 20% ethylene glycol monoethylether in 80 mM Tris-maleate, pH 7.5). Subsequently, 1 mg Fast Blue BB salt (Fluka, Buchs, Switzerland) was added, the solution filtered using a 0.45-µm filter and applied to deparaffinized paraffin sections for 5 minutes. Sections were rinsed in PBS, dehydrated, and mounted in glycerol. Quantitative analysis of mast cells and neutrophils was performed by counting cells in five highpower fields (40  $\times$ ) per age/staged tissue section from five mice per neoplastic stage. Data presented reflect the average total cell count per field from the ventral ear leaflet. Statistical analysis was performed using GraphPad InStat version 3.0a for Macintosh (GraphPad Software, San Diego, CA). The specific tests that were used were the Mann-Whitney (unpaired, nonparametric two-tailed) test and the Fishers (unpaired) t test. P < .05 was considered statistically significant. All immunolocalization experiments were repeated on multiple tissue sections and included negative controls for determination of background staining, which was negligible. Data shown are representative of results obtained after examination of tissues removed from a minimum of four different mice per distinctive stage of neoplastic progression.

#### Determination of Keratinocyte Proliferation Index

To detect bromodeoxyuridine (BrdU)-positive keratinocytes in neoplastic and control tissues, animals received intraperitoneal injections of BrdU (Roche, Indianapolis, IN) dissolved in PBS at 50 µg/g total body weight 90 minutes before sacrifice and preparation of tissue samples. Fivemicrometer-thick paraffin sections were deparaffinized in xylenes, rehydrated in graded ethanol, boiled in Citra antigen retrieval solution (BioGenex, San Ramon, CA), washed in PBS, and blocked in blocking buffer and BrdU-positive cells were detected essentially as described by the manufacturer's recommendations using the BrdU Labeling Kit II (Roche), developed by Vector Red Alkaline Phosphatase Kit (Vector), counterstained with eosin, dehydrated through graded ethanol and xylenes, and mounted in Cytoseal 60 (Richard-Allan Scientific). Images used for quantification were captured at high magnification (40  $\times$ ) on a Leica DM-RXA microscope attached to a Leica digital camera operated by OpenLab software (Improvision, Lexington, MA). The proliferative index was quantified from five high-power  $(40 \times)$  images per neoplastically staged tissue and included four mice per category as the percentage of BrdU-positive nuclei over the total number of keratinocytes. Statistical analysis was performed using GraphPad InStat version 3.0a for Macintosh (GraphPad Software). The specific tests that were used were the Mann-Whitney (unpaired, nonparametric two-tailed) test, and the Fishers (unpaired) t test. P < .05 was considered statistically significant.

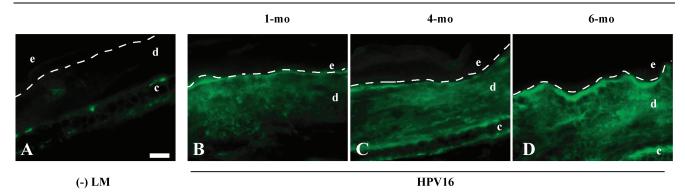
#### Flow Cytometry

The magnitude of inflammatory infiltrate in neoplastic tissues was analyzed by flow cytometry. Ear skin biopsies from (-)LM, HPV16 mice, and HPV16/C3<sup>-/-</sup> mice at specific time points were manually minced using a scalpel, followed by a 13-minute enzymatic digestion with 2.5 mg/ml collagenase Type II (Worthington, Lakewood, NJ), 2.5 mg/ml collagenase Type IV (Gibco, Carlsbad, CA), and 0.5 mg/ml DNase (Sigma) in PBS containing 1% BSA (Sigma) (PBS/ BSA) at 37°C under continuous stirring conditions. The digest was quenched by adding Dulbecco's modified Eagle's medium (DMEM) (Gibco) containing 10% FBS (Gibco) and was filtered through a 70-µm nylon filter (Falcon). Single-cell suspensions were treated with PharM Lyse ammonium chloride lysing reagent (BD Biosciences) for 10 minutes to remove erythrocytes. Cells were washed with DMEM containing 10% FBS, followed by a wash with PBS/BSA. Cells were incubated for 10 minutes at 4°C with rat anti-mouse CD16/CD32 mAb (BD Biosciences) at a 1:50 dilution in PBS/ BSA to prevent nonspecific antibody binding. Subsequently, cells were washed and incubated for 20 minutes with 50  $\mu$ l of 1:100 dilution of APC-conjugated anti-mouse CD45 (eBioscience, San Diego, CA). Cells were washed twice with PBS/BSA and 7-AAD (BD Biosciences) was added at a dilution of 1:10 to discriminate between viable and dead cells. Data acquisition and analysis were performed on a FACSCalibur using CellQuestPro software (BD Biosciences). Statistical analysis was performed using GraphPad InStat version 3.0a for Macintosh (GraphPad Software, San Diego, CA). The specific tests that were used were the Mann-Whitney (unpaired, non parametric two-tailed) test, and the Fishers (unpaired) t test. P < 0.05 was considered statistically significant.

#### Results

## Neoplastic Progression in HPV16 Mice Is Characterized by C3 Deposition

Early neoplastic progression in HPV16 mice is characterized by a dramatic influx of cells of the innate immune system, in particular mast cells and neutrophils [37,38]. To investigate whether the observed accumulation of inflammatory cells in the neoplastic microenvironment was accompanied by activation of the complement system, C3 deposition was examined by IF microscopy on tissue sections and ELISA with tissue lysates (Figure 1). Whereas only minimal detection of C3 was observed in nontransgenic skin, abundant dermal deposits of C3 were prominent in hyperplastic and dysplastic skin of HPV16 mice (Figure 1). Using a C3 ELISA, we found a statistically significant increase in C3 levels in hyperplastic skin lysates from 1-month-old HPV16 mice compared with nontransgenic controls (P<.05, Fishers (unpaired) t test; data not shown). As expected, C3 staining and immunoreactivity (based on ELISA) was not observed



**Figure 1.** Deposition of C3 in the neoplastic microenvironment. Immunofluorescence for C3 (FITC signal) on 10- $\mu$ m OCT-embedded frozen tissue sections in (A) negative littermate (–LM) ear skin, (B) hyperplastic (1 month of age), (C) early dysplastic (4 months of age), and (D) late dysplastic (6 months of age) ear tissue from HPV16 mice. Dashed line indicates epidermal–dermal interface. The epidermis (e), dermis (d), and cartilage (c) are indicated. Scale bar: 50  $\mu$ m (A–D).

in C3<sup>-/-</sup> mice (data not shown). These data suggest that neoplastic progression in HPV16 mice is characterized by early activation of the complement system and C3 accumulation coincident with initial development of hyperplasias. Moreover, C3 accumulation becomes more pronounced as neoplastic progression to dysplasia ensues, suggesting that activation of complement represents an early parameter of premalignant progression in HPV16 mice.

## Infiltration of Inflammatory Cells during Neoplastic Progression Is Not Reduced in the Absence of C3

Previous studies using HPV16 mice have revealed an important role for bone-marrow-derived inflammatory cells during skin carcinogenesis [37–39,42]. Given the critical role of the complement system in recruitment and activation of inflammatory cells during pathologic tissue remodeling [25,29,30,32–34,43,44], in combination with the observation that deposition of C3 is already detectable in hyperplastic skin lesions of HPV16 mice (Figure 1), we hypothesized that complement mediates recruitment of inflammatory cells toward neoplastic skin. To test this hypothesis, we took a genetic approach and generated HPV16/C3<sup>-/-</sup> mice [15,16] and examined several key characteristics of neoplastic progression in the absence and presence of C3.

To determine whether absence of C3 in HPV16/C3<sup>-/-</sup> mice altered characteristic inflammatory cell recruitment into early neoplastic lesions, we first profiled infiltration of leukocytes by quantitative detection of CD45 (leukocyte common antigen)-positive cells in age-matched neoplastic tissue from HPV16/C3<sup>-/-</sup> and HPV16 mice by flow cytometry (Figure 2A). This analysis revealed that the magnitude of the inflammatory infiltrate is most striking in early dysplasias (4 months of age), where ~35% of all cells are CD45<sup>+</sup> inflammatory cells (Figure 2A). Surprisingly, the absence of C3 in HPV16/C3<sup>-/-</sup> mice did not result in any quantitative changes in this profile of infiltrating CD45-positive leukocytes (Figure 2A, Mann-Whitney test).

To investigate the possibility that C3 deficiency resulted in a shift in immune cell profile, we quantitatively assessed the two major immune cell populations infiltrating HPV16 neoplastic skin, namely, mast cells and neutrophils [37,38]. We used CAE histochemistry on paraffin-embedded agematched tissue sections to detect serine esterase activity in mast cells *in situ*. As described previously [37], normal nontransgenic skin contains few mast cells distributed broadly throughout the dermis (Figure 2*B*). Mast cell numbers increase beginning in hyperplasias (1 month of age) and further increase with the onset of intense angiogenesis in dysplastic lesions (Figure 2*B*). Quantitative analysis of the number of mast cells *in situ* revealed that the number of infiltrating mast cells in skin from age-matched HPV16/ C3<sup>-/-</sup> mice was indistinguishable from that detected in agematched HPV16 controls (Figure 2*B*, Mann-Whitney test), suggesting that C3 is not required for recruitment of mast cells into premalignant skin lesions.

To assess if C3 instead regulated neutrophil recruitment, we quantitatively compared neutrophils in age-matched HPV16 and HPV16/C3<sup>-/-</sup> mice and found no significant differences (Figure 2*C*, Mann-Whitney test), suggesting that C3 is not involved in the recruitment of either mast cell or neutrophils into premalignant skin. Thus, although deposition of C3 in neoplastic skin coincides with infiltration of inflammatory cells, C3 does not mediate the recruitment of inflammatory cells into the neoplastic microenvironment.

## Activation of Keratinocyte Hyperproliferation and Angiogenesis during Neoplastic Progression Are not Regulated by C3

Although C3 does not appear to be essential for recruitment of inflammatory cells toward neoplastic lesions, we reasoned that C3 could instead be critical for activation of inflammatory cells in the neoplastic microenvironment. Mast cells secrete a variety of mitogenic factors affecting a wide range of cell types, including keratinocytes and endothelial cells [37,42,45-49]. Using mast-cell-deficient/HPV16 mice. we previously reported that mast cells contribute to neoplastic progression by potentiating keratinocyte hyperproliferation and activating angiogenesis [37], in part by release of matrix metalloproteinase-9 (MMP9) [38,42]. Together, these data suggest that activation of mast cells is an important contributing event to tumorigenesis in HPV16 mice. To assess whether C3 was involved in mast cell activation we compared keratinocyte proliferation indices and activation of angiogenic vasculature as readouts for mast cell activation in HPV16 and HPV16/C3<sup>-/-</sup> mice.

Hyperproliferation is an intrinsic property of neoplastic cells. Accordingly, we previously reported that keratinocyte hyperproliferation incrementally increases during neoplastic progression in HPV16 mice and, in part, characterizes progression between premalignant stages [35,36]. Keratinocyte proliferative indices were determined in age-matched skin from HPV16 and HPV16/C3<sup>-/-</sup> mice and were found to be similar at each neoplastic stage examined (Figure 3*A*, Mann-Whitney test), suggesting that C3 is not involved in regulating epithelial hyperproliferation downstream of mast cell activation.

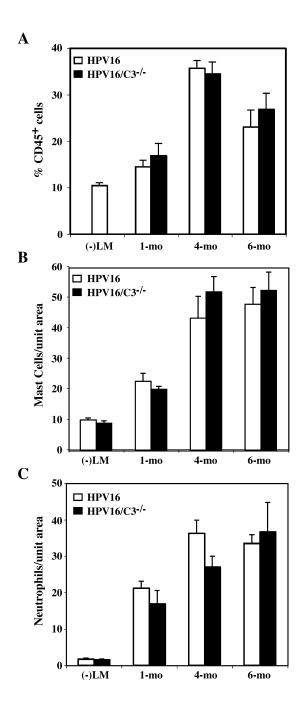
The vascular architecture changes in a characteristic manner during neoplastic progression in HPV16 mice [37,38] and can be monitored by immunodection of the endothelial cell-specific marker CD31. Normal nontransgenic mouse skin contains infrequent capillaries located deep within the dermis (Figure 3B). Neoplastic progression involves an early increase in capillary density first evident subjacent to hyperplastic epithelium, whereas dysplastic lesions contain dilated and enlarged capillaries that are increased in number indicative of an angiogenic switch from vascular guiescence to modest neovascularization in early low-grade lesions (hyperplasias) to a striking upregulation of angiogenesis in high-grade lesions (dysplasias) (Figure 3B). Comparison of vascular architecture, density, and patterning during premalignant progression in HPV16 and HPV16/C3<sup>-/-</sup> mice failed to reveal any differences (Figure 3B), suggesting that C3 does not contribute to inflammatory cell-mediated activation of angiogenic vasculature.

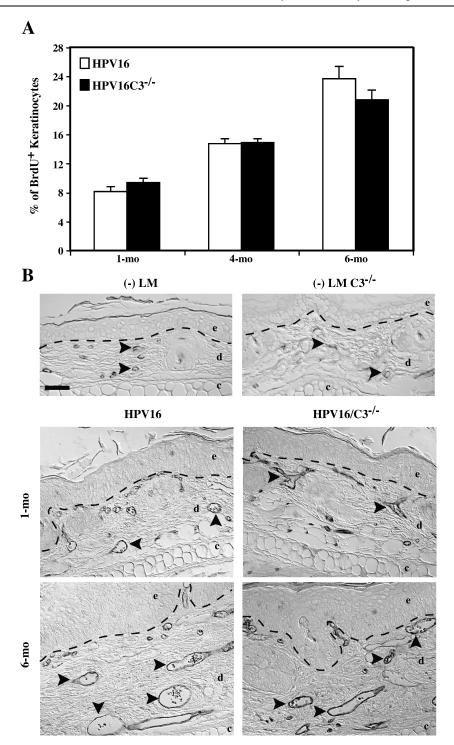
# IgG Deposition Suggests Antibody-Dependent, but Complement-Independent Regulation of Neoplasia-Associated Inflammation

Despite the fact that complement activation plays a prominent role in regulating many immunologic injuries [25,29–34,50] and significant deposition of C3 is observed in dermal compartments of neoplastic skin, the absence of

Figure 2. Infiltration of inflammatory cells during neoplastic progression is not reduced in the absence of C3. (A) Single-cell suspensions derived from negative littermate ear tissue (-LM) and hyperplastic (1 month of age), early dysplastic (4 months of age), and late dysplastic (6 months of age) ear tissue from HPV16 mice and HPV16/C3<sup>-/-</sup> mice were analyzed by flow cytometry to determine the percentage of CD45<sup>+</sup> cells as a percentage of total viable cells (n = 3). Results are shown as mean percentages ± SEM. No statistically significant differences were observed between age-matched neoplastic tissues of HPV16 and HPV16/C3<sup>-/-</sup> mice (Mann-Whitney unpaired t test). (B) Quantitative analysis of mast cells at distinct stages of neoplastic progression in ear tissue from HPV16, HPV16/C3<sup>-/-</sup> mice, and negative littermates (–LM). Mast cells were visualized by CAE histochemistry on 5- $\mu$ m paraffin-embedded tissue sections. Values represent number of mast cells, averaged from five high-power fields per mouse and five mice per category. Error bars represent SEM. No statistically significant differences were observed between age-matched neoplastic tissues between HPV16 and HPV16/C3<sup>-/-</sup> mice (Mann-Whitney unpaired t test). (C) Quantitative analysis of neutrophils at distinct stages of neoplastic progression in ear tissue from negative littermate (-)LM, HPV16, and HPV16/C3<sup>-/-</sup> mice. Neutrophils were visualized by immunohistochemistry on 5-µm paraffin-embedded tissue sections. Values represent number of neutrophils, averaged from five highpower fields per mouse and five mice per category. Error bars represent SEM. No statistically significant differences were observed between agematched neoplastic tissues between HPV16 and HPV16/C3<sup>-/-</sup> mice (Mann-Whitney unpaired t test).

C3 did not functionally alter leukocyte recruitment or activation during premalignant progression in HPV16 mice. Inflammatory cells can, however, be recruited and activated directly by antibodies through cross-linking of antibody surface receptors, e.g., Fc $\gamma$  and/or Fc $\varepsilon$  receptors [51,52]. To study whether the humoral immune response is involved in regulating inflammation in HPV16 mice, the spatial and temporal deposition patterns of IgG antibodies in neoplastic skin were analyzed by IF (Figure 4). In negative littermate nontransgenic skin, low IgG immunoreactivity was detectable in the dermis (Figure 4). In HPV16 hyperplastic skin (1 month of age), extensive regions of IgG antibodies were visualized throughout the dermis that was further expanded and intensified in dysplastic lesions (4 to 6 months of age) (Figure 4). HPV16/C3<sup>-/-</sup> mice displayed





**Figure 3.** Activation of keratinocyte hyperproliferation and angiogenesis during neoplastic progression is not affected in the absence of C3. (A) Quantitative analysis of the percentage of BrdU-positive keratinocytes (keratinocyte proliferative index) at distinct stages of neoplastic progression in ear tissue of HPV16 and HPV16/C3<sup>-/-</sup> mice at 1, 4, and 6 months of age. BrdU<sup>+</sup> cells were visualized by immunohistochemical detection on 5- $\mu$ m paraffin-embedded tissue sections. Values represent keratinocyte proliferative indices, averaged from five high-power fields per mouse and four to eight mice per category. Error bars represent SEM. No statistically significant differences were observed between age-matched neoplastic tissues between HPV16 and HPV16/C3<sup>-/-</sup> mice (Mann-Whitney unpaired t test). (B) Immunolocalization of CD31/PECAM-1 (black staining) in 5- $\mu$ m paraffin-embedded sections of age-matched neoplastic car tissue of neoplastic ear tissue of HPV16 and HPV16/C3<sup>-/-</sup> mice at all ages tested. Dashed line indicates epidermal –dermal interface. Arrowheads indicate representative blood vessels. The epidermal (e), dermal (d), and cartlage (c) regions of the tissues are indicated. Scale bar: 50  $\mu$ m.

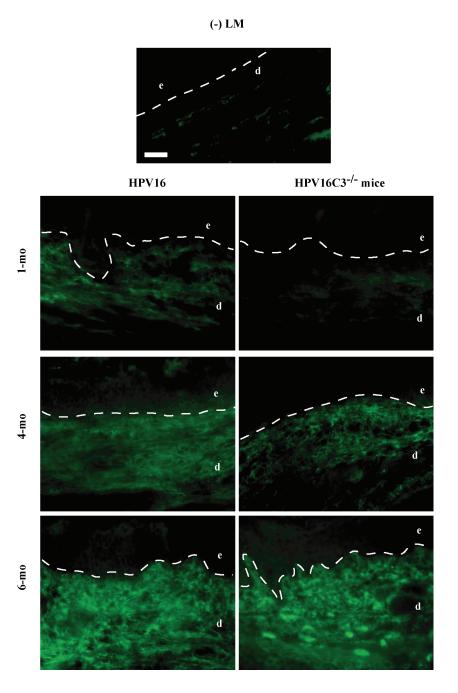
similar IgG deposition patterns during neoplastic progression (Figure 4), suggesting that an antibody-mediated immune response independent of complement activation and C3

deposition might therefore be involved in mediating recruitment of inflammatory cells during malignant progression in HPV16 mice.

#### Discussion

In the present study we examined the functional significance of C3 deposition and complement activation in a mouse model of epithelial carcinogenesis characterized, in part, by prominent inflammatory cell infiltrates in premalignant lesions. Results from this study suggest that C3 is neither required for the dramatic accumulation of inflammatory cells in premalignant skin nor influences the contributory role of inflammatory cells to squamous carcinogenesis, e.g., keratinocyte hyperproliferation or activation of angiogenic vasculature. These data stand in contrast to studies showing a crucial role for complement in recruitment and activation of inflammatory cells in other states of pathologic remodeling, e.g., airway hyperresponsiveness [25,29,30], IgG-antigen complex-mediated inflammation [31], intestinal ischemia/ reperfusion [32,50], delayed-type hypersensitivity [33], and subepidermal blistering disease [34].

The experimental model of IgG-mediated subepidermal blistering disease [34,43,44] has many characteristics in common with the HPV16 mouse model of *de novo* multistage skin carcinogenesis, namely, deposition of IgG and complement at the dermal–epidermal junction and recruitment of



**Figure 4.** IgG deposition in the neoplastic microenvironment of HPV16 mice. Direct immunofluorescence on 10- $\mu$ m OCT-embedded frozen tissue sections detects IgG antibodies (FITC signal) in negative littermate (–LM) ear skin, hyperplastic (1 month of age), early dysplastic (4 months of age), and late dysplastic (6 months of age) ear tissue from HPV16 and HPV16/C3<sup>-/-</sup> mice. Dashed line indicates epidermal-dermal interface. The epidermis (e), dermis (d), and cartilage (c) are indicated. Scale bar: 50  $\mu$ m.

mast cells and neutrophils [34]. Moreover, subepidermal blister formation is mediated by mast cell activation; mastcell-deficient mice are resistant to experimental subepidermal blistering disease [44]. Likewise, we have previously reported that mast-cell-deficient/HPV16 mice display attenuated characteristics of neoplastic progression [37]. However, whereas subepidermal blistering disease depends on complement activation, as complement component C5–deficient mice failed to develop cutaneous blisters [34], no functional significance of the complement system was observed in our study. To our knowledge, this is the first report in which the role of the complement system in a model of *de novo* neoplastic progression has been investigated.

Why is neoplastic progression and its accompanied inflammation not altered by deficient complement activation? The major difference between inflammation associated with malignant progression and that associated with other states of pathologic remodeling, e.g., subepidermal blistering disease, airway hyperresponsiveness, intestinal ischemia/ reperfustion, and delayed hypersensitivity, is the duration of inflammatory response. Inflammation during neoplastic progression is not abated because the "wound does not heal" [2]; thus, inflammatory cells fail to egress out of neoplastic tissue, therefore representing a chronic disease state. Moreover, antigens that are likely important for activation of the adaptive immune system similarly are present chronically and likely induce chronic B-cell responses, resulting in significant antibody deposition into neoplastic tissue.

If complement is not required for recruitment of inflammatory cells toward premalignant lesions, what other complement-independent recruitment pathways might be involved? Inflammatory cells can be recruited and activated through a complement-independent, antibody-dependent pathway. Antibodies can interact directly with inflammatory cells through antibody binding surface receptors, e.g., Fcy and/or Fcc receptors [51,52]. Fc receptors are multimeric cell-surface receptors that bind the Fc portion of antibodies and are expressed by many hematopoietic cells [53]. Antibody-mediated cross-linking of Fc receptors results in recruitment and activation of inflammatory cells. For example, in IgE-associated hypersensitivity reactions and allergic disorders, mast cell activation is initiated by cross-linking of IgE bound to the high-affinity FccRI receptors, resulting in secretion of products that can have effector, immunoregulatory, or autocrine effects [51]. It has become clear that in addition to IgE-dependent recruitment mechanisms, IgG-dependent mechanisms can regulate recruitment and activation of mast cells through Fcy receptors [52]. In vitro incubation of human mast cells through Fcy RI with IgG1 antibodies resulted in degranulation, increased production of arachidonic acid metabolites, and tumor necrosis factor- $\alpha$  release [52]. Fc $\gamma$ receptor-deficient mice are resistant to development of experimental immune hemolytic anemia and immune thrombocytopenia [54]. Likewise, Fcγ receptor expression on mast cells has been reported to promote the course of experimental allergic encephalomyelitis (EAE) [55]. In addition, antibody-mediated experimental cutaneous Arthus reaction requires expression of Fcy receptors on inflammatory cells

and develops normally in complement-deficient mice [56,57]. Thus, activation of cell-surface Fcy receptors can result in activation of an inflammatory response independent of complement activation. We observed early and abundant deposition of IgG antibodies in premalignant skin lesions (Figure 4) and the spatial and temporal deposition profile of IgG antibodies coincided with the inflammatory infiltrate; thus we are currently investigating whether antibodies localized in the neoplastic microenvironment trigger complementindependent recruitment and activation of mast cells. Because inflammatory cells, and in particular mast cells, are important contributors to the early stages of carcinogenesis in HPV16 mice as well as during human epithelial carcinogenesis, a detailed understanding of the molecules and mechanisms involved in activating the dramatic recruitment will eventually facilitate the rational design of novel therapies against human cancer.

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