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Prospect for immunological therapies of the equine malignant melanoma

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Summary: Although equine melanocytic neoplasms occur frequently, at least in grey horses, their existence is neglected frequently, despite their potentially severe clinical consequences. Immune-based therapies have been investigated in human metastatic melanoma and offer encouraging results. Researchers have recently begun to explore the immunological approach in equine malignant melanoma (EMM) with some success. Therefore, this review aims at giving a short overview of the nature of EMM and providing insight into the immunological basis of melanocytic neoplasms in general. Equine malignant melanoma is a true neoplasm with a genetic basis. Initially they show a slow invasive growth, but they have the potential to metastasize. EMM and human malignant melanomas share various common histopathologic features. For the immune system eradication of the tumor is a challenge. Although solid tumors are often infiltrated with immune cells, they usually do not induce a significant tumor remission. The antitumoral immune response depends on factors of the innate immune system interacting in a coordinated fashion with the adaptive immune system. Antigen-presenting cells, neutrophilic granulocytes, and lymphoid cells may elicit antitumoral effects. This innate response precedes a downstream antigen-specific response. In the tumor microenvironment of solid tumors immune suppressive factors can induce immune escape mechanisms of the tumors. Immunological therapies of EMM include blockage of histamine 2 receptors on melanoma cells and lymphocytes, cancer vaccines and gene therapeutic approaches. This article focuses on the immunological therapeutic approach of EMM and gives an outlook on future implications.

Keywords: horse, neoplasia, cancer immunity, immunotherapy, interleukin, DNA vaccine

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Introduction

This review describes the advances made in immune-mediated therapeutic approaches to equine malignant melanoma (EMM). After a brief outline of the nature and clinical signs of EMM, the main immunological principles against melanocytic neoplasms in general are depicted and therapies aiming at immune-mediated effects in the treatment of EMM are described. The review concludes with an assessment of problems encountered when attempting to establish a functional therapy and gives an outlook on future potential prospects.

Equine malignant melanoma is one of the most common neoplastic conditions of the skin in horses, with a considerably high incidence in aging grey horses (*Mcfadyean* 1933, *Fleury* et al. 2000b). It is, nowadays, accepted as a true neoplasm (*Gorham* and *Robl* 1986, *Macgillivray* et al. 2002, *Patterson-Kane* and *Ginn* 2003, Seltenhammer et al. 2004). Several research groups have investigated the genetic basis of melanoma formation and found a link between the autosomal dominant trait of greying with age and melanoma formation in horses (*Rieder* et al. 2000, *Seltenhammer* et al. 2003, *Rosengren Pielberg* et al. 2008, *Sundstrom* et al. 2012).

Typically, grey horses of five or more years are affected with EMM, showing cutaneous tumors at predilection sites, such as the ventral tail, perianal region, prepuce, eye lid, parotid, or guttural pouches (*Rodriguez* et al. 1998, *Patterson-Kane* et al. 2001, *Macgillivray* et al. 2002, *Garvican* et al. 2007). Despite their initially slow invasive growth, they frequently metastasize later (*Baker* and *Leyland* 1975, *Valentine* 1995, *Fleury* et al. 2000a, *Smith* et al. 2002). For further information on clinical signs and conventional therapeutic options, the reader is referred to recent extensive reviews on the matter (*Metcalfe* et al. 2013, *Moore* et al. 2013, *Phillips* and *Lembkke* 2013, *Cavalleri* et al. 2014).

Nature of EMM

Despite some similarities, EMM and human metastatic melanoma differ in their aetiopathology as well as in many known molecular mechanisms. The cellular origin of the neoplasm in EMM is the same as in human malignant melanomas, though. Furthermore, it is a well-accepted fact that the predominant proportion of equine melanomas progress to malignancy (*Sundberg* et al. 1977, *Smith* et al. 2002, *Brown* et al. 2014). Nevertheless, in contrast to human melanomas, EMM tends to show a retardation of metastasis in many cases. Similarly, the pathophysiologic role of ultraviolet (UV) light in many types of human malignant melanomas (*Wood* et al. 2006, *Noonan* et al. 2012) is of no significance in EMM, where typical predilection sites (i.e. prepuce, underside of tail and perineal region) are well protected from sunlight. Hypothetically, a mutation of the melanocortin 1 receptor (MC1R) might be involved in this difference regarding the role of UV light. *Rosengren Pielberg* and colleagues (2008) found an association of an increased risk of developing EMM in horses with an increase in MC1R signalling.

It was stated that EMM could serve as an adequate animal model of naturally occurring melanomas for human dermal melanocytic disease (*Tuthill* et al. 1982, *Heinzerling* et al. 2001). Histopathologic features of EMM and human malignant melanomas were compared in a study conducted by *Seltenhammer* and colleagues (2004). They shared various common features identified by immunohistochemistry. In detail, melanocyte-specific proteins S-100, HMB-45 and T-311, proliferation markers PCNA and Ki-67, and adhesion molecule CD44 stained positively in both equine and human malignant melanoma samples. Immunohistochemically and histomorphologically equine melanomas in this study proved to be comparable to particular types of human malignant melanoma, i.e. the malignant blue nevi and desmoplastic melanomas. These features add to the authors' conclusion that EMM closely resembles certain types of human malignant melanoma and, therefore, might serve as a suitable animal model of the corresponding human malignancy.

The genetic grounds in EMM development show similarities to human malignant melanoma where a family history of melanoma is one of the strongest risk factors for melanoma (*Miller* and *Mihm* 2006). However, even taking into consideration how many specific molecular mutations are known in human melanoma as opposed to how little is known in EMM, genetic predispositions likely differ substantially. There are many aspects that differ significantly from human melanoma. One unique trait of EMM seems to be the lack of junctional activity in the grey horse EMM (*Fleury* et al. 2000a, *Seltenhammer* et al. 2003). Another dissimilar characteristic is the responsibility of UV exposure for some mutations causing melanoma in humans, and a further difference between EMM and human malignant melanoma clearly is the aggressiveness of the neoplastic growth. The human melanoma is the most aggressive type of skin cancer. The median survival of patients suffering from metastatic melanoma is usually less than one year (*Agarwala* 2009).

Lesion site in EMM frequently corresponds to satellite and in-transit lesions known in human medicine

The term satellite lesion in human malignant melanoma refers to melanoma foci in the near circumference (less than 2 cm) of the primary tumor. In melanoma of the head and neck, they are encountered in approximately 3% of cases (*Hasney* et al. 2008). In-transit lesions, on the other hand, are tumors found at a greater distance to the primary melanoma, but still within the drainage area of the same lymph node draining the primary tumor (*Hasney* et al. 2008). *Moore* et al. (2013) proposed a clinical classification into 4 progressing stages in which horses showing only a single melanoma with a diameter < 2 cm and a slow growth is classified as stage 1. Stage 2 describes horses with multiple small (< 2 cm) tumours without dissemination or metastases and a slow or quiescent growth pattern. Animals showing multiple melanomas < 4 cm in diameter with dissemination or metastases and a slow growth rate are classified as stage 3. Those multiple melanomas with a diameter of > 4 cm and a rapid growth are classified as stage 4. The multiple tumor foci commonly observed at predilection sites accounting for stage 2 or higher normally show lesions corresponding to the human medical definition of satellite and in-transit lesions (Figure 1).

In short, the multifactorial aetiology of EMM has not yet been fully elucidated. Similarities between EMM and human disease facilitate comparisons of disease features or therapies. However, differences between the disease entities clearly exist and have to be considered.

Immune mechanisms against melanocytic neoplasms

Since neoplasms arise from the host organism's own cells, the eradication of the tumor is a challenge to the immune system. Although solid tumors are often infiltrated with a variety of immune cells, those are not usually able to induce a significant tumor remission. Recent work suggests that those T-cell inflamed tumors contain an abundant amount of immune regulatory cells and substances hindering the effective tumor lysis. Amongst these are forkhead box p3 (FoxP3) + regulatory T cells (Treg), programmed death ligand 1 (PD-L1), and indoleamine-2,3-dioxygenase (IDO) inducing immune escape of tumor cells.

Innate immune mechanisms

Innate cells involved in tumor recognition and defense are antigen-presenting cell populations, neutrophils, and lymphoid cells. Macrophages and dendritic cells (DC) act as antigen presenting cells (APC) in antitumoral immune responses. DCs secrete type I interferons and present tumor derived antigens to $CD8^+$ T cells (*Den Haan* et al. 2000, *Lou* et al. 2007). However, depending on the microenvironment of the tumor, DC populations can be modified towards an immunosuppressive phenotype associated with tumor progression (*Woo* et al. 2015). Similarly, polarisation of macrophages towards a functional inflammatory phenotype (M1 macrophages) is associated with a tumoricidal activity, whereas alternatively activated M2 macrophages have a protumor phenotype and are involved in tumor growth (*Sica* and *Mantovani* 2012). Tumor associated macrophages (TAM) usually present an M2-like phenotype. Functional M1 macrophages have a high IL-12, low IL-

Figure 1 Dermal melanomas at a typical predilection site in the perianal region of a grey horse.

10 cytokine profile and are activated by IFNγ and TNFα. M2-like TAMs have been associated with tumor progression and poor prognosis (*Bingle* et al. 2002, *Chen* et al. 2005, *Zabuawala* et al. 2010, *Zhang* et al. 2012). Cytokines and chemokines secreted by tumor cells (like CCL2, colony stimulating factor-1, IL-10 and others) can recruit monocytes and stimulate the conversion into M2-like TAMs (*Hagemann* et al. 2006, *Qian* et al. 2011). They result in blocking T cell proliferation and thereby inhibition of antitumoral T cell responses. Immunosuppressive TAMs were shown to downregulate IL-12p70 but upregulate IL-12p40, IL-23, IL-6 and IL-10 and show a decrease in NFB and C/EBP transcription factors family members (*Rodríguez* et al. 2013). One proposed mechanism is the increased expression of programmed death ligand 1 (PD-L1) (*Kuang* et al. 2009). Blocking of the protumorigenic alternative macrophage activation in an experimental setting of murine melanoma transplants resulted in reduced melanoma outgrowth and enhancement of macrophage proinflammatory responses (*Yaddanapudi* et al. 2013). IFNγ and a combination of CpG and anti-IL10-antibodies activating the TLR9 pathway have been used to switch M2-like TAMs into M1 macrophages (*Guiducci* et al. 2005, *Duluc* et al. 2009). Mirroring the M1/M2 and Th1/Th2 polarisation, neutrophils also display a plasticity resulting in antitumorigenic N1 and protumorigenic N2 neutrophils (*Piccard* et al. 2012). In investigations on infection driven stimulation of neutrophils it was shown that equine neutrophils can exhibit non-phagocytic properties and act as a crucial bridge between innate and adaptive immunity by expression of tumor necrosis factor alpha (TNFα), IL-12p35, IL-12p40, IL-6, IL-8, IL-23p19 and IFNγ (*Nerren* et al. 2009). However, due to their plasticity, tumor associated neutrophils have been shown to be important in tumor development, progression and angiogenesis (*Mantovani* et al. 2011). This has also been proposed for melanoma in a mouse model (*Jablonska* et al. 2010).

In general, a polarisation towards M1, N1 or Th1 allows for a synergistic antitumoral immune response. Upstream of tumor-induced T cell priming type I IFNs seem to be essential for antitumor activity (*Gonzalez-Navajas* et al. 2012, *Fuertes* et al. 2013). They have been shown to activate DCs and natural killer (NK) cells and potentially inhibit infiltration of solid tumors by TAMs (*Swann* et al. 2007, *U'ren* et al. 2010). Innate lymphoid population (NK cells, NKT cell, γδT cells) are able to respond early to inflammation and may in this way be crucial in early recognition of malignant cell alterations (Woo et al. 2015). Natural killer cells have fundamental antitumoral features: They kill cells with absent or reduced MHC class I molecule expression, which is often the case during tumor progression (*Davies* et al. 2014). Infiltration of solid tumors with NK cells may lead to a more favourable prognosis (*Pietra* et al. 2012). DNA damage of tumor cells can result from replication stress and deouble-strand helical breaks in proliferating tumor cells or tumors treated with radiation or chemotherapy. This damage can lead to a DNA damage response resulting in activation of NK cells through increased expression of NK cell-activating receptor ligands (*Bartkova* et al. 2005, *Gorgoulis* et al. 2005, *Lord* and *Ashworth* 2012, *Woo* et al. 2015). Various cytokines (IL-12, IL-18, IL-15, IFNα) stimulate NK cell antitumoral effects by activation and maturation (*Waldhauer* and *Steinle* 2008). Antitumoral effects are mediated by perforin and granzyme release and ligands binding to Fas antigen (FasL) and the TNF-related, apoptosis-inducing ligand (TRAIL) (*Cheng* et al. 2013). IFNγ secretion adds to their tumor destructive properties by enhancing the TRAIL pathway (*Yang* et al. 2014) and stimulates proliferation and activation of macrophages and monocytes as well as differentiation of T cell subsets (*Martin-Fontecha* et al. 2004). Another innate immune cell type playing a role in antitumoral immune surveillance is the Natural Killer T cell (NKT) (*Smyth* et al. 2000, *Crowe* et al. 2002). Two different forms of NKT cells are known, the CD1d dependent iNKT cells and the type II NKT cells recognizing antigens on alternate T cell receptors (*Woo* et al. 2015). Amongst antigen ligands binding on CD1d is the tumor-derived disialoganglioside GD3 (*Wu* et al. 2003) but also CD1d-presented self-antigens on IL-12 producing DCs (*Matsuda* et al. 2008). Upon activation iNKT cells show cytotoxic properties and secretion of IFNγ. In contrast, type II NKT cells are described as immunoregulatory through expression of IL-13 and signal transducer and activator of transcription 6 (STAT6) in tumor models (*Terabe* et al. 2000, *Berzofsky* and *Terabe* 2008). The existence of an NKT/CD1d system in horses has been proven (*Looringh van Beeck* et al. 2009). A further T cell subset, γδT cells, is described in horses as well as in other species (*Schrenzel* and *Ferrick* 1995). Similar to NK cells they contain cytotoxic granules and are able to express/produce IFNγ, TNFα, FasL and TRAIL (*Kondo* et al. 2008). They are even capable of phagocytosis (*Wu* et al. 2009). They are known to act as a bridge between innate and adaptive responses by various features, e.g. indirectly by activation of DCs and expressing the surface molecule CD16 and aid in antibody-dependent cellular cytotoxicity (ADCC) (*Tokuyama* et al. 2008). However, they might also have regulatory functions in certain tumor settings (*Woo* et al. 2015). Interestingly, extracellular ATP release from chemotherapeutically treated tumor cells has been proposed to play an important role in activating and inducing different innate immune populations and indirectly in recruiting $CD8^+$ T cells (*Elliott* et al. 2009, *Ghiringhelli* et al. 2009). Complement, a central section of the innate immunity, has been demonstrated on malignant cells and complement activation has been shown to be increased in tumor patients (*Niculescu* et al. 1992, *Lucas* et al. 1996, *Rikklin* et al. 2010, *Ajona* et al. 2013).

Adaptive immune mechanisms

The immunogenicity of melanoma antigens has been shown in several species (*Speiser* et al. 2003, *Gyorffy* et al. 2005, *Wolchok* et al. 2007, *Yuan* et al. 2009, *Manley* et al. 2011) and can be presumed for equine melanoma as well due to the antitumoral response seen after immune modifying therapies in EMM-bearing horses (*Jeglum* 1997, *Heinzerling* et al. 2001, *Finocchiaro* et al. 2009, *Müller* et al. 2011, *Phillips* and *Lembcke* 2013). Tumor-associated antibodies (TAA) are a central part in triggering this antitumoral response. Several TAAs known in melanoma and proposed as targets for immunotherapy are melanoma antigen recognized by T cells A (MART A), glycoprotein (gp) 75, gp100, tyrosinase, melanoma antigen (MAGE), S100, MELOE-1 and -2, PNL-2 and protein gene product (PGP) 9.5 (*Brichard* et al. 1993, *Bakker* et al. 1994, *Desmet* et al. 1994, *Gaugler* et al. 1994, *Kawakami* et al. 1994, *Brasseur* et al. 1995, *Wang* et al. 1995, *Godet* et al. 2010, *Ramos-Vara* et al. 2014). Tyrosinase, gp100, PNL-2, PGP9.5 and S100 have been identified in EMM (*Roels* et al. 2000, *Seltenhammer* et al. 2004, *Ramos-*

Vara et al. 2014). TAAs are presented to T cells either by antigen presenting cells (APC) or through major histocompatibility complexes (MHC). T cell recognition of TAA can then induce activation of effector cells, release of chemokines and cytokines and the induction of co-stimulatory molecules in order to stimulate a rather cellular Th1 and/or predominantly humoral Th2 response (*Spurrell* and *Lockley* 2014).

In general, cytotoxic $CD8^+$ T cells (CTL) are known to have effective antitumoral properties in melanocytic neoplasms (*Tuting* 2013). This is presumed to be equivalent in horses. CTL generation is favoured in Th1-biased immune responses. A study on plasmid gene therapy coding for equine cytokines IL-12 and -18 induced an in vitro expression of IFNα stimulated by IL-12 (*Müller* et al. 2011). The apparent positive effect of intratumoral Th1 $CD4^+$ T cells (*Lowes* et al. 1997, *Wagner* et al. 1998, *Conrad* et al. 1999) has yet to be proven in EMM.

Tumor infiltrating T-cells (TILs) in human melanoma may correlate positively with patient survival (*Oble* et al. 2009). In the case of a Th1 response, positive effects on tumor regression were reported for tumor tissue infiltrating CTL, T helper cells or a combination of both T cell populations (*Topalian* et al. 1989, *Mihm* et al. 1996, *Dudley* et al. 2002, *Taylor* et al. 2007, *Hillen* et al. 2008). However, infiltrating Tregs inhibit immune surveillance in the tumor microenvironment and have been associated with an increased recurrence rate in human melanoma patients (*Miracco* et al. 2007).

Synergism of innate and adaptive immune mechanisms in cancer surveillance and destruction

Links between innate and adaptive immunity are necessary to coordinate the immune response with innate response preceding a downstream antigen-specific response (Figure 2). One of these is IFNα, an important cytokine for stimulation of $CD8^+$ T cells and NK cells (*Moreno* et al. 2008, *Terabe* and *Berzofsky* 2008). Furthermore, type I IFNs (mainly IFNα and IFNβ) appear to be critical in bridging innate and adaptive immune responses in the tumor setting (*Woo* et al. 2015). Besides, cytosolic DNA-sensing pathways might contribute to innate immune sensing. Tumor-derived DNA has been detected in tumor-infiltrating DCs (*Woo* et al. 2014).

Escape mechanisms

Tumor cells are able to modify immune cell functions to promote tumor growth (*Lacy* et al. 2012). In the tumor microenvironment of solid tumors immune suppressive factors such as IDO, TGF-β, prostaglandin E2 (PGE2) and NKG2D are found (*Martinet* et al. 2009a, *Martinet* et al. 2009b). Progressing areas of human melanoma exhibited a greater number of IL-10- and TGF-β-expressing T cells than thinner regressive tumor areas (*Conrad* et al. 1999). Thus, the tumor environment tends to induce regulatory T cells and inhibit antitumoral Th1 cells. Metastatic melanoma contain more Treg TILs than

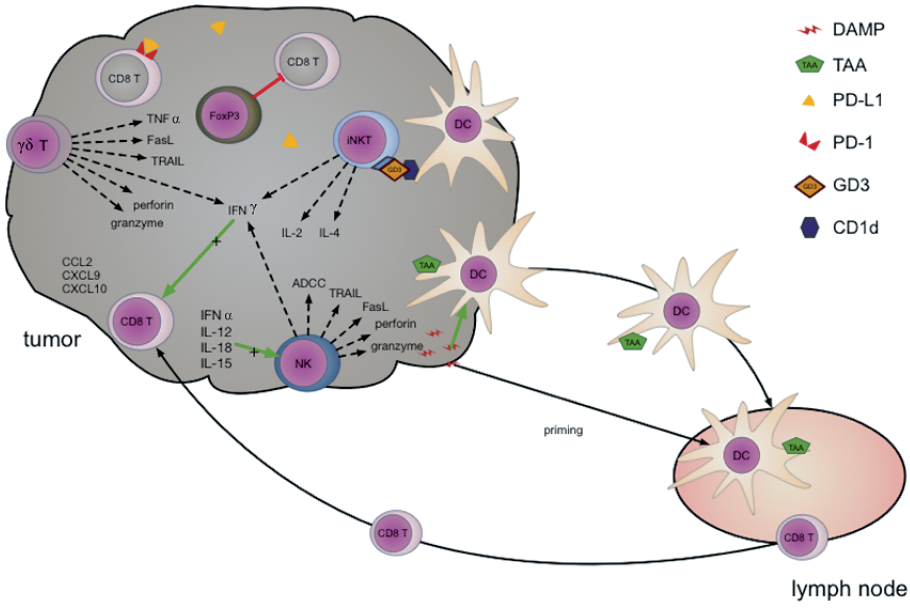

Figure 2 Line drawing of suspected immune mechanisms in solid tumors like EMM.The tumor microenvironment is abundant of antigens. Here, tumor-associated antigens (TAA) are taken up by antigen-presenting dendritic cells (DC), which after activation by damage-associated molecular patterns (DAMPs) migrate to the draining lymph node where they stimulate naive antigen-specific T-cells to become effector T cells. Primed T effector cells (here: CD8+ T-cells) traffic back to the tumor, attracted by chemokines in the tumor microenvironment like CCL2, CXCL9 and CXCL10, where they perform anti tumoral activity. In the tumor microenvironment, regulatory T cells (Tregs, FoxP3-positive) may inhibit other activated effector T cells. Similarly, tumor cells can express PD-L1 (programmed death ligand 1), an inhibitory ligand of PD-1 (receptor expressed on activated CD8+ T-cells). Tumor-infiltrating NK cells (NK) are activated by receptors sensing tumor cell damage. This can occur spontaneously during tumor progression, by proinflammtory cytokines or in response to anti tumoral therapy. Upon activation NK cells may kill tumor cells by releasing perforin and granzyme-containing granules, expression of Fas ligand, TNF-related apoptosis-inducing ligand (TRAIL), antibody-dependent cellular cytotoxicity (ADCC) and secretion of Interferon gamma (IFNγ). Released TAA by lysed tumor cells are then taken up by DC. The release of Th1- and Th2 cytokines by innate NK T-cells (iNKT) can be induced after their recognition of tumor derived disialoganglioside GD3 bound to CD1d on DCs. Tumor-infiltrating gamma delta T cells (γδT) can be activated by stress induced mediators resulting in the release of perforin, granzymes as well as IFNα, TNFγ, FasL and TRAIL.

superficial thin melanoma lesions. Reduced immune reactions, represented by an increased expression of the regulatory T cell marker FoxP3 in the primary tumor were associated with significantly decreased survival (*Gerber* et al. 2014).

Melanoma cells are able to express T-cell inhibitory molecules, such as programmed death-ligand 1 (PD-L1) or cytotoxic T lymphocyte antigen 4 (CTLA-4) (*Jiang* et al. 2013, *Laurent* et al. 2013). PD-L1 binds to the immune inhibitory receptor PD-1, expressed on T cells, where it inhibits the activation of antitumoral T cells and stimulates Treg proliferation (*Pardoll* 2012). No studies regarding these inhibitory immune responses for equine melanoma have been published yet. While PD-1/PD-L1 interactions play an important regulatory role in prevention of autoimmunity in infection/inflammation control, they can induce tumor immune escape and apoptosis of activated T cells in the tumor microenvironment. In human melanoma PD-L1 expression in tumors varies and has been associated with prognosis of response to immunotherapy targeting PD-1 or PD-L1 (*Topalian* et al. 2012).

Immunological melanoma therapies

Research and clinical application of immunological therapies against human malignant melanoma is extensive and, therefore, beyond the scope of this review. Hence, the reader is referred to review articles on immunological melanoma therapy for further reading (*Finn* 2008, *Mouawad* et al. 2010, *Gyorki* et al. 2013).

The immune reaction has to be able to overcome the inhibition induced by tumor cells in order to induce melanoma remission. This can be accomplished with therapeutics directed against tumor-specific or tumor-associated antigens (*Graziano* and *Finn* 2005). Since conventional tumor treatments like radiotherapy, chemotherapy, and photodynamic therapy have been shown to provoke apoptosis or necrosis of tumor cells, released damage-associated molecular patterns (DAMP) may enable antitumor immunity (*Obeid* et al. 2007, *Garg* et al. 2012a, *Garg* et al. 2012b, *Dudek* et al. 2013). In fact, combination therapy using conventional treatment modalities in combination with immunological therapy have been used with varying success in human melanoma patients and might therefore be interesting to try in EMM (*Hoshimoto* et al. 2012, *Davar* et al. 2013).

Blockage of host molecules

One early anti-melanoma therapy in horses targeting the immune system was the use of the histamine 2 (H2) receptor antagonist cimetidine as an immune response stimulating agent. H2 receptors were demonstrated on Tregs and melanoma cells (*Sahasrabudhe* et al. 1987, *Reynolds* et al. 1996). Histamine regulates the Th1 and Th2 responses via H1 and H2 receptors on these cells (Jutel et al. 2001). *Uçar* (1991) demonstrated a direct effect of the H2 receptor antagonists cimetidine and ranitidine on murine and human melanoma cells inhibiting cell proliferation. This is an additional mechanism described to the otherwise suggested immunological effect through inhibition of Tregs (*Rocklin* and *Haberek-Davidson* 1981). Furthermore, the augmentation of natural killer cell activity against melanoma was shown for cimetidine (*Flodgren* et al. 1985). *Goetz* and colleagues described a positive therapeutic effect of cimetidine in the treatment of equine melanomas in 1990 (*Goetz* et al. 1990). However, in a later report, cimetidine treatment in metastatic melanoma was not successful in 5/5 horses (*Macgillivray* et al. 2002). Similarly, *Laus* and colleagues (2010) report no benefit of cimetidine monotherapy in EMM in a controlled clinical trial using two different dosing regimes (3.5 mg/kg bwt BID or 7.5 mg/kg bwt SID for 60 days) and a group size of five horses. Helle drew the same conclusion in a doctoral thesis of the University of Utrecht in 2012 (*Helle* 2012) in a study on 40 grey horses with melanoma, evaluating cimetidine treatment. Orally administered cimetidine has a low and variable bioavailability (*Smyth* et al. 1990), potentially impeding the successful therapy of melanoma in horses.

In human medicine, inhibitory proteins like PD-L 1 and CTLA-4 provide targets for antitumoral therapies using monoclonal antibodies. Blockade of CTLA-4 with the monoclonal antibody ipilimumab leads to less suppression of T cells (*Peggs* et al. 2006) and improved survival in humans with metastatic melanoma (*Hodi* et al. 2010). Nivolumab, the monoclonal antibody targeting the T-cell surface molecule programmed death (PD) 1, has also been shown to be effective in the treatment of human melanoma (*Brahmer* et al. 2010, *Brahmer* et al. 2012, *Hamid* et al. 2013). Both antibodies were also used successfully in combination (*Wolchok* et al. 2013). However, clinical potential of this therapeutic strategy remains to be elucidated in EMM.

Cancer vaccines and gene therapeutic approaches

A specific immunotherapeutic approach using a whole-cell vaccine against equine melanoma was suggested as early as 1924 in a case report about the autologous vaccination against melanoma in one grey horse (*Mertens* 1924). Later, a case series was reported using an autologous vaccine as an irradiated whole-cell preparation injected subcutaneously over regional lymph nodes of 12 melanoma-bearing horses. Whole cell vaccination carries the chance to offer a wide range of tumor antigens to the immune system. To prepare the vaccine, surgical excision of a tumor was performed and a single cell suspension was prepared and cryopreserved by the laboratory. Vaccines were administered every other week for six vaccinations and every six weeks thereafter. Eleven out of 12 horses treated showed tumor regression over a follow-up period of up to two years (*Jeglum* 1997). Another study reports on three horses treated with an autologous vaccine prepared by the same laboratory in addition to systemic cimetidine treatment. Those horses received vaccinations for 12 to 36 months. All three horses eventually died from metastatic melanoma disease (*Macgillivray* et al. 2002).

At the University of Florida researchers investigated the immune response to disialoganglioside GD3 vaccination against melanoma in dogs (*Milner* et al. 2006). A similar therapeutic approach is under investigation in horses, too. However, to the authors' knowledge, no information on clinical trials of this vaccine in horses has been published yet.

A case report of a grey stallion with EMM describes the combination therapy of a local suicide gene therapy with ganci-

clovir and a systemic autologous anti-cancer vaccine and adjuvant xenogeneic cell cytokine therapy (*Finocchiaro* et al. 2009). In this report, surgical excision of several, but not all, superficial melanomas was performed with infiltration of the surgical margins with a combination of lipid-complexed plasmid DNA encoding herpes simplex virus thymidine kinase (HSVtk) suicide gene and ganciclovir. The HSVtk suicide gene had previously been shown to sensitize transfected cells to ganciclovir (*Finocchiaro* et al. 2009). An autologous melanoma cell vaccine, produced from excised melanomas, was then subcutaneously injected together with live irradiated Chinese hamster ovary cells modified to synthesize human interleukin 2 and human GM-CSF. The vaccine was applied at weekly intervals for five weeks, followed by injections every other week up to day 105, and thereafter, every fourth week until day 245. At the times of vaccination, all but three remaining superficial melanomas were infiltrated with the suicide gene/ganciclovir combination. This treatment resulted in partial or complete tumor regression, even in the three untreated tumors. Thus, indicating a systemic rather than just local effect of the treatment.The disadvantages of autologous cell therapies are relatively high costs, need for a specialised laboratory to prepare the cell suspension and the patient specificity of the cell suspensions prepared.

Several other gene therapeutic approaches have been published for the treatment of EMM. One of the advantages of gene therapy is that it may reduce systemic toxicity while improving the specificity of effects (*Schmidt-Wolf* and *Schmidt-Wolf* 1996). Different approaches to cancer gene therapy include inserting suicide genes into the tumor (*Finocchiaro* et al. 2009), enhancing the immunogenicity of the tumor or the antitumoral activity of effector cells (*Heinzerling* et al. 2001, *Müller* et al. 2011, *Lembcke* et al. 2012, *Phillips* and *Lembkke* 2013). Other strategies targeting the inhibition of neoangiogenesis, blockade of oncogene activity or insertion of wild-type tumor suppressor genes (*Schmidt-Wolf* and *Schmidt-Wolf* 1996) have not been reported in EMM treatment. Plasmids coding for antitumoral interleukins (*Heinzerling* et al. 2001, *Müller* et al. 2011) or DNA vectors encoding tumor-associated antigens (*Lembcke* et al. 2012, *Mahlmann* et al. 2015) have been evaluated for their use in EMM therapy.

The therapeutic application of genes coding for human IL-12 or equine IL-12 and -18 in grey horses with EMM resulted in partial tumor regression (*Heinzerling* et al. 2001, *Müller* et al. 2011). In a double-blind placebo-controlled investigation of 26 grey horses bearing melanomas, interleukin-18 encoding plasmid DNA, interleukin-12 encoding plasmid DNA or empty plasmid DNA were injected intratumorally. Both treatment groups resulted in significant tumor regression in contrast to the placebo-treated control horses, in which tumor growth was noted over the treatment period (*Müller* et al. 2011).

Cell type-specific differentiation antigens came into the focus of research for a more specific antitumoral vaccination (*Weide* et al. 2008). Due to self-tolerance towards these differentiation antigens, the immunogenicity of vaccines targeting these proteins is weak. A fundamental aspect of therapeutic vaccinations against melanoma is to overcome these self-tolerance mechanisms (*Houghton* 1994, *Naftzger* et al. 1996). Administration of DNA coding for xenogeneic melanocyte differentiation antigens was able to induce an immunological response against melanomas in several species (*Overwijk* et al. 1998, *Weber* et al. 1998, *Zhou* et al. 1999, *Gold* et al. 2003, *Bergman* et al. 2006, *Yuan* et al. 2009). It is suggested that the immune response against the foreign protein confers cross-reaction with the autologous antigen (*Haupt* et al. 2002, *Houghton* and *Guevara-Patino* 2004, *Srinivasan* and *Wolchok* 2004).

The melanocyte differentiation antigen tyrosinase gained attention as a potential target for therapeutic vaccination. Tyrosinase catalyses a rate-limiting step in melanin production (*Hearing* and *Tsukamoto* 1991, *Yamaguchi* and *Hearing* 2009), is expressed in all melanin-producing cells (*Hearing* and *Tsukamoto* 1991) and is constitutively expressed in equine melanoma cells (*Seltenhammer* et al. 2004, *Phillips* et al. 2012). Clinical and immunological effects of vaccination with xenogeneic tyrosinase DNA could be shown in dogs (*Bergman* et al. 2006, *Liao* et al. 2006) and humans (*Wolchok* et al. 2007, *Tarhini* et al. 2012). Consequently, the USDA granted a therapeutic human tyrosinase DNA vaccine against canine melanoma (Oncept®, Merial Ltd., Athens, GA) in 2010. This vaccine was able to induce specific antibodies and a cellular immune response in clinically healthy horses (*Lembcke* et al. 2012). The clinical off-label use of this vaccine has resulted in a favourable outcome in several horses making the therapeutical value in the therapy of EMM promising (*Phillips* and *Lembcke* 2013). However, using a different DNA vector, xenogeneic DNA vaccination of melanoma-bearing horses using human tyrosinase DNA or human gp100 DNA did not show a beneficial advantage in comparison to interleukin 12 and 18 DNA therapy alone (*Mählmann* et al. 2015).

Future outlook on immune therapy in EMM

Several approaches have been used to apply functional immunological therapies to the equine melanoma patient. The majority of therapies to date have had limited success in clinical trials. The most successful immunological treatment for EMM seems to be the xenogeneic DNA vaccination with human tyrosinase DNA (*Phillips* and *Lembcke* 2013). However, detailed results on the clinical off-label use of this vaccine have not yet been published. The impairment of inhibitory signals is a treatment mechanism currently receiving a lot of attention in human oncodermatology (*Wu* et al. 2012). Inhibitory T-cell receptors lead to a diminished immune response. Consequently, the effectiveness of the antitumor immune response can be increased by blocking those receptors with specific monoclonal antibodies (*Wu* et al. 2012). Target molecules under investigation in human malignant melanoma include cytotoxic T-lymphocyte antigen 4 (CTLA-4), programmed death 1 (PD1), programmed death-ligand 1 (PD-L1), and B- and T-cell attenuator (BTLA) (reviewed in *Pardoll* 2012). The equine equivalents of those receptors have not been identified and, to this date, no antibodies targeting these inhibitory T-cell receptors are available for the horse. This might be a fruitful field for further investigations in equine medicine.

Combined approaches targeting more than one molecule/effector cell will probably be more effective than monotherapies and should give guidance for future therapeutic strategies in EMM. A combination of local chemotherapy helping to destroy neoplastic tissue and, thereby, exposing

tumor antigens and immune adjuvants activating effector cells might be a target of future research.

List of abbreviations

ADCC	antibody-dependent cellular cytotoxicity
APC	antigen presenting cells
ATP	adenosine triphosphate
BID	bis in die
BTLA	B- and T-cell attenuator
bwt	body weight
C/EBP	CCAAT/enhancer binding protein
CCL	chemokine (C-C motif) ligand
CD	cluster of differentiation
CD1d	cluster of differentiation 1d
CpG	Cytosine-phosphate-Guanosine
CTL	cytotoxic CD8+ T cells
CTLA-4	cytotoxic T lymphocyte antigen 4
CXCL	CXC chemokine ligand
DAMPs	damage-associated molecular patterns
DC	dendritic cell
GD3	ganglioside D3
DNA	desoxyribonucleic acid
EMM	equine malignant melanoma
FasL	Fas antigen ligand
FoxP3	forkhead box p3
GM-CSF	granulocyte monocyte colony stimulating factor
gp	glycoprotein
H	histamine
HMB-45	homatropine methyle bromide-45
HSVtk	herpes simplex virus thymidine kinase
IDO	indoleamine-2,3-dioxygenase
IFN	interferon
IL	interleukin
iNKT	innate NK T-cells
Ki-67	kinase inhibitor 67
MAGE	melanoma antigen encoding
MART A	melanoma antigen recognized by T cells A
MC1R	melanocortin type-1 receptor
MELOE	melanoma-overexpressed antigen
MHC	major histocompatibility complex
NF B	nuclear factor kappa B
NK	Natural Killer
NKG2D	natural killer group 2, member D
PCNA	proliferative cell nuclear antigen
PD	programmed death
PD-L1	programmed death ligand 1
PGE2	prostaglandin E2
PGP	protein gene product
SID	semel in die
STAT6	signal transducer and activator of transcription 6
T-311	tyrosinase-311
TAA	tumor-associated antigen
TAM	Tumor associated macrophages
TGF-ß	transforming growth factor
Th	T helper
TILs	Tumor infiltrating T-cells
TLR9	toll like receptor 9
TNF	tumor necrosis factor
TRAIL	TNF-related apoptosis-inducing ligand
Treg	regulatory T cell
USDA	United States Department of Agriculture
UV	Ultraviolet

Conflict of interest statement

None of the authors of this paper has a financial or personal relationship with other people or organisations that could inappropriately influence or bias the content of the paper.

Authors' contributions

JC designed and drafted the manuscript. KM helped to draft the manuscript. HS contributed to the conception and helped to draft the manuscript and figure. KF helped in the conception and critically revised the manuscript. All authors read and approved the final manuscript.

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Erweiterte Zusammenfassung

Perspektiven für immunologische Therapien des Malignen Melanoms beim Pferd

Melanome sind bei Pferden, in den meisten Fällen bei Schimmeln, häufig auftretende Neoplasien der Haut. Typischerweise sind ältere vergrauende Pferde betroffen, die kutane Tumoren der pigmentbildenden Zellen an Prädilektionsstellen wie der Unterseite der Schweifrübe, der Perianalregion, dem Präputium, Augenlidern, Lippen oder der Parotis zeigen. Nach einem initial meist langsamen invasiven Wachstum kommt es häufig in einem späteren Stadium zur Tumormetastasierung. Eine genetische Komponente der Melanomentstehung wurde durch den Zusammenhang des autosomal dominanten Merkmals mit dem Alter zu Vergrauen und der Entstehung von Melanomen gefunden. Aufgrund der immunhistochemischen und histomorphologischen Vergleichbarkeit des equinen Melanoms mit bestimmten Typen des humanen Melanoms scheinen an Melanomen erkrankte Schimmel geeignet als Modelltiere für bestimmte humane dermale melanozytäre Erkrankungen wie maligne blaue Nevi und desmoplastische Melanome. Aufgrund der Entstehung aus körpereigenen Zellen, ist die Bekämpfung von Neoplasien eine Herausforderung für das Immunsystem. Trotz Infiltration solider Tumoren mit verschiedensten Immuneffektorzellen sind diese häufig nicht in der Lage, eine bedeutsame Tumorregression zu bewirken. Die T-Zellinfiltration in soliden Tumoren beinhaltet häufig regulatorische Zellen und Substanzen, die eine effektive Tumorlyse verhindern. Zur effektiven Tumorbekämpfung ist das konzertierte Zusammenspiel des angeborenen (unspezifischen) mit dem adaptiven (spezifischen) Immunsystem unabdingbar. An der Erkennung und Abwehr von Tumoren sind Antigen-präsentierende Zellen, neutrophile Granulozyten und lymphoide Zellen der unspezifischen Immunabwehr beteiligt. Eine Polarisierung dieser Zellen zugunsten einer antitumoralen Wirksamkeit (Makrophagen vom Typ I, neutrophile Granulozyten vom Typ I, T-Helfer-Zellen vom Typ I) führt zu Synergien in der Tumorabwehr. Interferone spielen hierbei eine Rolle in der Aktivierung und Stimulation der Effektorzellen. Unspezifische lymphoide Zellen sind durch ihre Fähigkeit zur frühen Antwort auf Entzündungsprozesse wichtige Zelltypen in der frühzeitigen Erkennung entarteter Zellen. In der spezifischen Tumorimmunität spielen tumorassoziierte Antikörper (TAA) eine zentrale Rolle in der Antitumorantwort. Verschiedene TAA des Melanoms sind bekannt und können als Ziel einer Immuntherapie fungieren. Sie werden den T-Zellen entweder durch antigen-präsentierende Zellen oder durch MHC-Komplexe dargeboten. Nach Erkennung der TAA durch T-Zellen können dann weitere Effektorzellen aktiviert, Chemokine und Zytokine freigesetzt und co-stimulatorische Moleküle induziert werden. Dadurch wird dann eine eher zelluläre Th1-Antwort oder eine vornehmlich humorale Th2-Antwort stimuliert. Damit die Immunantwort koordiniert abläuft und der unspezifischen Reaktion eine antigenspezifische Antwort folgt, müssen die beiden Anteile des Immunsystems gut vernetzt sein. Interferone übernehmen wichtige Aufgaben in dieser Vernetzung der antitumoren Immunantwort. Auch Tumor-DNA kann im Zytosol dendritischer Zellen erkannt werden und damit zur Aktivierung der unspezifischen Abwehrmechanismen beitragen. Daher und aufgrund des guten Ansprechens Melanome anderer Spezies auf eine immunmediierende Therapie ist die Immuntherapie von equinen Melanomen eine vielversprechende Forschungsrichtung. Es gibt Untersuchungen zur Immuntherapie des equinen malignen Melanoms mit antitumoral wirkenden Zytokinen aber auch mit Tumorvakzinen. Die Mehrzahl der untersuchten therapeutischen Strategien zeigte in klinischen Studien auf partielle Tumorregression begrenzte Erfolge. Eine der erfolgversprechendsten Therapien scheint die Vakzinierung von Melanompatienten mit xenogener (humaner) Tyrosinase-DNA zu sein. Bisher fehlen allerdings noch wissenschafltich publizierte Daten zum klinischen Therapieerfolg der Vakzine. Zukünftig könnten auch beim Pferd Therapieansätze mit mehr als einem Zielmolekül/einer Zielzelle in den Fokus der Melanomtherapieforschung rücken. Solche kombinierten Therapien sind vermutlich effektiver als Monotherapien und könnten beispielsweise auch die Kombination lokaler Chemotherapie zur Tumordestruktion und Freisetzung von Tumorantigenen mit immunstimulatorischen Substanzen zur effektiven Erkennung der freigesetzten TAA einschließen. Der Fokus dieses Übersichtsartikels liegt auf den Möglichkeiten der Immuntherapie equine maligner Melanome und ihrer potenziellen zukünftigen Anwendung am Patienten.