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Immunotherapy of cancer

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Abstract

Major advances have been made in the field of immunology in the past two decades. A better understanding of the molecular and cellular mechanisms controlling the immune system, has opened the door to many innovative and promising new cancer therapies that manipulate the immune response. For instance, toll-like receptor agonists have been shown to boost immune responses toward tumors. Also, a wide array of cell-based immunotherapies utilizing T cells, NK cells, and dendritic cells have been established. Furthermore, a rapidly expanding repertoire of monoclonal antibodies is being developed to treat tumors, and many of the available antibodies have demonstrated impressive clinical responses. Here, we examine some of these immunotherapeutic approaches currently in use or testing to treat cancer, and we examine available evidence with regards to mechanism and efficacy of these treatments.

Keywords

Immunotherapy; cancer; monoclonal antibody; cell-based therapy; toll-like receptors

1. Introduction

For decades, primary cancer treatments have incorporated both chemotherapeutic agents and ionizing radiation to eliminate the bulk of tumor mass. While many of these therapies have offered substantial benefit and some cures, the incidence of tumor relapse is a significant problem that results from the development of drug resistance mechanisms in a portion of the tumor cells. Therefore, additional therapeutic approaches to eliminate these resistant tumor cells must be established.

One appealing alternative strategy is therapy designed to harness the immune system to induce a potent anti-tumor response. Historically, the first successful immunotherapy to treat cancer involved the use of toxins from *Streptococcus erysipelatis* and *Bacillus prodigious* by William Coley in the 1890's (Coley, 1991). More recently, the development of vaccines to tumorcausing hepatitis B virus and papilloma virus are contributing significantly to preventing cancer

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in a large portion of the human population (Blumberg, 1997; Rogers et al., 2008). Since our understanding of basic immune mechanisms has expanded substantially, a wide array of immune pathways have been identified as attractive targets to promote anti-tumor responses in cancer patients.

Many immune cell types influence tumor growth in humans. The immune system is comprised of both innate cells that mediate immediate, short-lived responses [monocytes, macrophages, dendritic cells, and natural killer (NK) cells] and adaptive cells that develop long-lived responses and memory (T cells and B cells). The functions of these cells in response to cancer are schematized in Figure 1. The innate cells provide the earliest responses by releasing cytokines, directly lysing abnormal cells (NK cells) or capturing debris from dead cells (monocytes, macrophages, and dendritic cells) to present peptide fragments of specific foreign antigens to T cells in the context of major histocompatibility complex (MHC) molecules. The adaptive T and B cells generate antigen-specific responses that are delayed by several days, since minor subpopulations expressing antigen-specific receptors must proliferate and differentiate to generate a multitude of competent and activated effector cells. These proliferation and differentiation events are promoted by inflammatory cytokines [interferon (IFN)-γ, interleukin (IL)-12] produced by the activated innate immune cells or T cells. T cells can be divided into two major subsets consisting of CD8+ cytolytic (CTL) and CD4+ helper (Th) cells that can directly lyse tumor cells or release immunomodulatory cytokines, respectively, while B cells generate a humoral response by secreting antigen-specific immunoglobulin (Ig). Human CTL and Th cells express α/β antigen receptors (TCR) that recognize foreign antigenic peptides presented in the contexts of MHC class I (HLA-A, -B, and –C) or class II (HLA-DR, -DP and –DQ), respectively. Th cells can be subdivided based upon their capacities to secrete various cytokines into: Th1 cells (secrete type-1 cytokines, especially IFN-γ and IL-2, which are considered inflammatory), Th2 cells (secrete type-2 cytokines, including IL-4, IL-5, and IL-13, which are associated with allergic responses), and Th17 cells (secrete IL-17 and IL-22, which can cause inflammation in skin and mucosal surfaces and can also promote autoimmune responses). Additional relevant T cell subpopulations include invariant NK-like T (iNKT) cells, γ/δ T cells, and regulatory T (Treg) cells, and the cytokines produce by these T cell subsets are summarized in Figure 2. The iNKT cells provide a rapid and potent source of IL-4 and IFN-γ, which is triggered through recognition of glycolipids presented by MHC-like CD1d molecules by their invariant α/β TCR. Alternatively, the γ/δ T cells express a γ/δ TCR that is triggered by recognition of a variety of ligands, which appear to be recognized directly in an Ig-like manner (Born et al., 2007). In contrast, Treg cells have been found to infiltrate tumors and draining lymph nodes, where they can repress tumor-specific CTL responses through a variety of mechanisms, including production of the immunosuppressive cytokines IL-10 and transforming growth factor (TGF) β (Allan et al., 2008). Tumor-associated macrophages and myeloid-derived suppressor cells can also release immunosuppressive cytokines within the tumor microenvironment to inhibit dendritic cells functions and reportedly promote angiogenesis and tumor growth (Melief, 2008;Pittet, 2009).

The effectiveness of an anti-tumor immune response can be compromised by the immunosuppressive impacts of either cytotoxic chemotherapeutic drugs or cytokines produced within the tumor microenvironment [especially IL-10, TGF-β, IL-6, and prostaglandin E2]. Recent findings indicate that some tumors establish a protective immune-privileged environment by promoting the production of type-2-related cytokines (IL-4, IL-5, IL-6, IL-10, IL-13, and IL-21). The type-2 cytokines suppress the production of inflammatory type-1 related cytokines (IFN-γ, IL-12, IL-2) to thereby blunt the adaptive anti-tumor immune response (Becker, 2006). In fact, polarization of the immune response toward type-2 cytokine production has been associated with progression of a number of tumor types (Becker, 2006). Additional mechanisms of immune evasion by tumor cells include minimal expression or

shedding of tumor-associated antigens and shedding of ligands for activating receptors, such as the ULBP2, MICA and MICB ligands recognized by the NK cell receptor NKG2D, thereby preventing its function (Salih et al., 2006; Salih et al., 2002; Waldhauer and Steinle, 2006).

The field of cancer immunotherapy covers a vast array of manipulations to provoke immune responses toward tumors in both humans and animals. Therefore, we will only touch upon certain techniques that have promising or proven clinical value in treating cancer in patients. In the following discussion, we will focus on advances in the use of Toll-like receptor agonists, cellular immunotherapies, and antibody-based immunotherapies.

2. Toll-like receptor agonists

The family of Toll-like receptors (TLR) recognize specific molecular patterns found on byproducts of various pathogens, especially bacteria. Agonists for TLR2, TLR3, TLR4, and TLR9 are particularly attractive for boosting inflammatory innate immune responses to treat cancer. These stimulants have been historically referred to as "adjuvants" to boost vaccine therapies, and most of their effects are due to boosting the activity of myeloid cells (macrophages, monocytes, and dendritic cells). TLR2 ligands include muramyl peptides, peptidoglycans, and lipoteichoic acids, while polyinosinic acid:cytidylic acid (poly I:C) is a synthetic double stranded RNA that serves as a classic agonist of TLR3. Lipopolysaccharide (LPS) is the well-known agonist for TLR4, which has been identified as the "hemorrhage producing fraction" from the Gram negative bacteria that stimulated antitumor effects in Coley's toxin (Garay et al., 2007). Bacillus Calmette-Guérin (BCG) contains peptidoglycans from the cell wall of mycobacteria, which is an agonist for both TLR2 and TLR4 that has been used for decades to stimulate immune responses in treating bladder cancer (Garay et al., 2007). TLR9 can be stimulated with unmethylated CpG DNA sequences that are significantly more prevalent in bacteria than vertebrates (Jahrsdorfer and Weiner, 2008), and this may account for some non-specific antisense oligonucelotide effects. TLR2 or TLR4 agonists stimulate macrophages and dendritic cells to produce inducible nitric oxide synthetase (iNOS) and numerous proinflammatory cytokines and chemokines, including IL-1, IL-6, IL-8, IL-10, IL-12, IL-23, TNF- α , MIP1 α , and MIP1 β (Garay et al., 2007). On the other hand, plasmacytoid dendritic cells are selectively stimulated by the TLR3 agonist, poly I:C, to secrete IFN-α and IFN-β, which are potent activators of NK cells and CTL. The use of CpG analogs to stimulate TLR9 on dendritic cells can induce production of IL-12, IFN-α and IFN-β (Jahrsdorfer and Weiner, 2008). Although not detailed here, a wide array of compounds have been developed as TLR agonists that are currently being tested to treat cancer. It is important to note, however, that TLR agonists can also stimulate the growth and survival of some tumors (Rakoff-Nahoum and Medzhitov, 2009), so they may only be therapeutically useful under defined conditions.

3. Cellular immunotherapies

3.1 T cells

In view of their importance in establishing adaptive immune responses, many groups have attempted to activate a patient's own (autologous) tumor-reactive T cells by culturing them with IL-2 and other cytokines and reinjecting these cells to treat cancer. Such methods are designed to increase the number of reactive T cells and provide long-term immune protection with minimal autoimmune responsiveness. Importantly, however, the need for specialized cell culture facilities and time intensive cell preparation requirements have limited their use. Success of adoptive transfer has been improved by using chemotherapy and/or total body irradiation to achieve lymphodepletion prior to cell infusion, as well as IL-2 infusions post infusion. Although attempted under a wide variety of conditions, adoptive transfer of ex vivo activated autologous T cells has had limited effectiveness in treating leukemia (Curti et al., 1998; Morgan et al., 2006; Schultze et al., 2001), but some success in treating melanoma (Oble

et al., 2009). In contrast to bacterial or viral pathogens, tumor cells generally lack "foreign" antigens, and therefore, T cell reactivity is directed toward peptide antigens derived from altered or over-expressed gene products. The ex vivo expansion and reintroduction of tumorinfiltrating lymphocytes (TILs) is a particularly attractive approach, since TILs can contain tumor-specific T cell populations (Disis et al., 2009; Oble et al., 2009). $CD4^+$ Th cells have also been found to be more effective than $CDS⁺ CTL$ in adoptive transfer therapies (Perez-Diez et al., 2007). T cell reactivity toward several antigens over-expressed in melanoma (such as MAGE-6, Melan-A/MART-1, NY-ESO-1, and gp100) has been established from the characterization of TILs (Oble et al., 2009). These and other antigens are being studied as potential tumor vaccines. Some success has also been observed using genetically engineered T cells in which an antigen-specific antigen receptor has been introduced by retrovirus (Morgan et al., 2006).

Dendritic cells are potent antigen-presenting cells that serve as critical initiators of T celldependent immune responses. Therefore, they have been widely studied as potential vectors that can be loaded with peptide antigens for presentation to T cells and stimulate adaptive immunity (Disis et al., 2009; Melief, 2008). In particular, many groups have loaded with peptides derived from characterized tumor-associated antigens or material from a patient's tumor to introduce cancer vaccines (Disis et al., 2009; Kuwabara et al., 2007; Melief, 2008; Nagayama et al., 2003; Yamanaka, 2009).

Allogeneic hematopoietic stem cell transplantation has been widely used to exploit the T cellmediated allogeneic immune response in cancer immunotherapy, especially for treating highrisk leukemia patients, where graft-versus-leukemia responses can be beneficial. In allogeneic T cell responses, the donor T cells respond to the MHC molecules of the recipient as foreign antigens. While allogeneic T cell responses mediate the graft-versus-leukemia effect, the development of T cell-mediated graft-versus-host disease can be a serious complication of hematopoietic stem cell transplantation that often requires lifelong immunosuppressive therapy and can be fatal. As mentioned above, ex vivo-activated autologous T cell immunotherapies are designed to avoid autoimmune responses, but these approaches have generated only limited success in treating cancer, and graft-versus-host disease can still occur in the autologous context. The failure of adoptive T cell therapies can be attributed, at least in part, to ineffective homing of the activated T cells to the tumor site, contamination of T cell preparations with Treg cells, short-term survival of ex vivo activated T cells, emergence of antigen-deficient tumor variants, or suppression of T cell responsiveness by immunosuppressive factors within the tumor microenvironment.

3.2 NK cells

The activation of NK cells is regulated by signals derived from a variety of cell surface activating and inhibitory receptors. Activating receptors include the natural cytotoxicity receptors (NCR: NKp30, NKp44 and NKp46), NKG2D, 2B4, NKp80, and DNAM-1. The major control of NK cell activation, however, is mediated by their expression of inhibitory receptors that are specific for class I MHC (MHC-I) molecules, which prevent attack of normal (MHC-I-expressing) cells in the body. Some tumor cells and virus-infected cells avoid recognition by CTL by down-regulating their expression of MHC-I, but these escape mutants consequently become targets for NK cells, since they lack the inhibitory receptor ligands. In this way these inhibitory receptors mediate NK cell tolerance toward normal "self" cells, and their lack of engagement provides the mechanistic basis for recognition and attack of tumors that are lack these "self" ligands, as shown in Figure 3. Killer cell Ig-like receptors (KIR) constitute the major NK cell inhibitory receptors that recognize cognate MHC-I in humans. KIR have evolved in higher mammals as a polymorphic receptor family encoded by distinct genes, each of which can detect a subset of the wide array of polymorphic human MHC-I

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molecules (HLA-A, -B, and –C) (Vilches and Parham, 2002). Expression from individual KIR genes is independently regulated within each NK cell, and during NK cell development, a repertoire of cells is established, in which distinct NK cells stochastically express different members of the KIR family. In order to achieve functional competence, each NK cell must express at least one KIR family member that recognizes at least one of the expressed "self" MHC-I molecules (Anfossi et al., 2006;Cooley et al., 2007). To create further diversity, individuals in the human population inherit different haplotypes of KIR family members (Uhrberg et al., 1997), and most individuals express some KIR that do not recognize their own "self" MHC-I molecules. Due to such diversity, it is possible that a subset of NK cells from one individual will not be tolerized by the MHC-I expressed by another individual, since appropriate ligands are missing for the available inhibitory KIR. This is known as allogeneic NK cell responsiveness, which can occur under conditions of transplantation.

Ruggeri et al. first reported that NK cell alloreactivity, due to KIR-ligand mismatch, can significantly improve survival advantage when introducing haploidentical hematopoietic stem cell transplants to treat acute myeloid leukemia (AML) (Ruggeri et al., 2002). The benefit was observed when the donor expressed KIRs that lacked MHC-I ligands in the recipient. In addition to a beneficial graft-versus-leukemia effect to eliminate the AML, allogeneic NK cells were shown to substantially diminish the incidence of the deleterious T cell-mediated graftversus-host disease. The reduction in graft-versus-host disease in hematopoietic stem cell transplants under conditions of allogeneic NK cell responsiveness appears to be due to elimination of residual antigen presenting cells by the allogeneic NK cells (Ruggeri et al., 2002). While some groups have shown similar benefits from KIR-ligand mismatch in treating AML (Beelen et al., 2005; Giebel et al., 2003; Hsu et al., 2006; Hsu et al., 2005; Leung et al., 2004; Miller et al., 2007a; Yu et al., 2009), others have failed to observe any benefit under allogeneic NK cell transplantation conditions (Davies et al., 2002; Farag et al., 2006). In comparing the methods used, success of the haploidentical KIR-ligand mismatch approach in treating AML required specific conditions: 1) use of haploidentical donors, 2) myeloablative pre-conditioning of the recipient, 3) extensive T cell depletion of the hematopoietic stem cell graft, and 4) infusion of high numbers of highly purified hematopoietic stem cell.

Umbilical cord blood is also used as a hematopoietic stem cell transplant source for treating hematologic malignancies. Umbilical cord blood contains significantly fewer mature T cells, which is predicted to be advantageous for reducing the incidence of graft-versus-host disease and expected to favor the benefit of KIR-ligand mismatch. A recent study, however, found that the degree of conditioning of the recipient prior to transplant significantly influenced outcome of umbilical cord blood transplantation (Brunstein et al., 2009). In this study, transplantation of KIR-ligand mismatched umbilical cord blood after reduced intensity conditioning resulted in significantly increased incidence of severe graft-versus-host disease and reduced survival, while the introduction of mismatched umbilical cord blood after myeloablative conditioning did not influence graft-versus-host disease or survival. Therefore, although KIR-ligand mismatch may be advantageous under distinct conditions of hematopoietic stem cell transplantation to treat myelogenous leukemia, it is not universally beneficial and may even be deleterious to patients under other conditions.

Recent work has established improved survival of AML patients that have received hematopoietic stem cell transplants from a donor expressing more of a subset of activating KIR (Cooley et al., 2009). In addition to the inhibitory KIR, other members of this receptor family are known to trigger NK cell activation, although their ligands are currently unknown. While many haplotypes of inherited KIR genes (A haplotypes) contain only one activating KIR (KIR2DS4), others contain more than one activating member (B haplotypes). Cooley et al. have found that hematopoietic stem cell transplants to AML patients from B haplotype donors expressing two or more activating KIR result in reduced relapse and improved overall survival,

although incidence of chronic graft-versus-host disease was increased (Cooley et al., 2009). These effects were independent of inhibitory KIR-ligand match/mismatch, indicating that the impacts of KIR and their ligands are complex, but can clearly influence outcome and can benefit patients if optimized accordingly.

Further work is clearly necessary to establish the optimal conditions that favor the benefits of KIR/ligand match/mismatch to promote anti-tumor NK cell responses after hematopoietic stem cell transplantation. As an alternative, the direct adoptive transfer of mature haploidentical NK cells has also been shown to be safe and improve outcome of AML in some patients (Miller et al., 2005a; Passweg et al., 2005). In addition, it may be possible to boost NK cell responsiveness by blocking NK cell inhibitory receptors with antibodies, as has been demonstrated to potentiate ADCC responses in recent *in vitro* studies (Binyamin et al., 2008). Finally, a highly cytolytic NK-like cell line, named NK-92, may have therapeutic potential if injected into cancer patients after irradiation (Klingemann and Martinson, 2004), and genetic manipulation of this or other NK-like cell lines may further enhance this unique therapeutic approach. Therefore, NK cells remain a viable target for cancer immunotherapy.

4. Antibody-based immunotherapies

Antibody-based therapy has emerged as an integral part of effective treatments for a number of malignancies. In the last decade, monoclonal antibodies have become major therapeutic vehicles in the treatment of malignant and nonmalignant diseases. The success of immunebased treatments and an improved understanding of the role of the immune system in cancer development and progression have mapped the road for extensive development of antibodybased therapy.

Paul Ehrlich, a century ago, described the antibody therapy concept to selectively target malignant cells based on the unique expressed determinants profile of the disease (the "magic bullet" hypothesis). The advent of the advanced technology of hybridoma production provided the necessary skills to efficiently produce highly specific monoclonal antibodies (mAb) (Kohler and Milstein, 1975; Kohler and Milstein, 1976). Extensive efforts were taken to apply this technique and to generate mAb for cancer therapy. Many mAb that are currently in use were originally derived from hybridoma-derived murine antibodies (Brekke and Sandlie, 2003). A second generation of engineered antibody was employed, considering the major issue of stimulation of human anti-mouse antibody responses and the inefficient interaction of the murine origin constant region of the antibodies (Fc) with human immune-accessory cells (Mirick et al., 2004). Using an IgG scaffold, humanized and chimeric antibodies were generated by incorporating portions of the murine variable regions into the human IgG backbone (Kim, 2003). Grafting either the entire murine variable regions (chimeric mAb) or the murine complementary-determining regions (humanized mAb), mAb are created that contain human Fc domains and preserve their target antigen specificity (Trail et al., 2003). In addition, using molecular engineering techniques, critical human heavy-chain backbone sequences were grafted onto the xenogeneic murine antibody structure, reducing the immunogenicity and providing the important human Fc domain to the resulting antibodies (Khazaeli et al., 1994). Chimeric alteration from fully murine to human Fc yields prolonged circulation time and more effective cytotoxicity responses. To further increase the power of mAb and to design antibodies that specifically target and subsequently eliminate cancer cells, large combinatorial antibody libraries of murine, human or synthetic origin were constructed. Efficient in-vitro screening systems now allow bypassing immunization and selecting recombinant antibodies of defined specificity without the need for hybridoma production (Benhar et al., 2000; Hoogenboom, 2002). New forms of antibody modules with different size, flexibility and valency suited for in vivo imaging and therapy have been created using genetic manipulation, recombinant

production and antibody conjugation techniques (Irving et al., 2001; Worn and Pluckthun, 2001).

4.1 Mechanisms of tumor cytotoxicity upon antibody binding

Antibodies may mediate anti-tumor effects by employing a variety of effector mechanisms, including antibody-dependent cellular cytotoxicity (ADCC), complement-dependent cytotoxicity, and inducing or blocking intracellular signals, but the exact mechanisms of action of many of these clinically active antibodies remain unknown. Understanding these events will be key to optimizing antibody engineering and to combination therapy to enhance efficacy. In the following sections we will review some of the available data regarding these functions.

4.1.1 Antibody-dependent cellular cytotoxicity (ADCC)—ADCC is a well-recognized immune effector mechanism in which antigen-specific antibodies direct immune effector cells of innate immunity to the killing of the antigen-expressing cancer cells (Ahmad and Menezes, 1996). This property is dependent upon interactions between cellular Fc receptors (FcR) on immune accessory cells and the Fc portion of an antibody (Steplewski et al., 1983). The binding cross-links FcR to activate the effector cells [e.g., natural killer (NK) cells are triggered to kill cancer cells and also release cytokines and chemokines] (Iannello and Ahmad, 2005). While ADCC can be mediated by NK cells, monocyte-macrophages and neutrophils, NK cells are the principal ADCC effectors as they express low affinity type IIIA Fc receptor (FcRγ IIIa; CD16a) on their surface (Ernst et al., 2002; Metes et al., 2001; Morel et al., 1999) and CD16a plays a dominant role in triggering NK cell-mediated ADCC (O'Hanlon, 2004). NK cells kill target cells by directed release of cytotoxic granules containing perforin and granzymes. Antitumor activity of clinically effective mAb against human tumor xenografts growing in either wild-type mice or in murine FcγRII/III knock out mice (Clynes et al., 2000) was reduced in the Fcγ receptor knock-out mice, and was conserved when only the inhibitory FcγRII receptor isoform was removed, stressing the importance of FcγRIIIa (CD16a) in anti-tumor responsiveness (Clynes et al., 2000). Antibodies designed to take advantage of Fc functions, such as rituximab, are IgG1 subclass, as opposed to immunoconjugates, designed to kill by drug or toxin delivery, that are usually IgG4 subclass to limit complement activation and toxicity. The role of CD16a in the anti-tumor effects of rituximab is further supported by the finding that patients with CD16 polymorphisms that generate a high affinity receptor for Fc correlate with improved outcome of follicular lymphoma responses to rituximab (Weng and Levy, 2003b). Thus interactions between the antibody Fc domain and the Fc receptor underlie at least some of the clinical benefit of some mAb, and imply the importance of ADCC. Ongoing research strategies include molecular manipulation of the mAb Fc region to enhance complement activation and/or ADCC. Combined computational and experimental methods have identified mutations within the Fc domain of mAb to selectively tune the affinity for FcγRIII and other Fcγ receptors (Hayes et al., 2002). An alternative strategy to enhance ADCC by mAb is to modify Fc glycosylation by the producing cell line (Umana et al., 1999). Modification of the Fc region to interact with activating or with inhibitory Fc receptors also has potential to enhance antigen presentation by dendritic cells and can be biased to promote or to inhibit the generation of cytotoxic T-cell responses against the targeted antigen. In summary, numerous findings indicate that the induction of ADCC can lead to adaptive immune responses and finally to the elimination of tumor cells (Rafiq et al., 2002; Weiner and Carter, 2005).

4.1.2 Antibody binding inducing intracellular signals—Many antibodies targeting cell surface receptors are designed to alter intracellular signaling pathways. Examples include impacts on epidermal growth factor receptor (EGFR) and HER2/neu blockade. Recent data that mutations in intracellular pathways (i.e. K-ras) determine whether antibody binding is cytotoxic indicate an important role for intracellular pathways in antibody anti-tumor activity.

CD20 is not generally thought of as a signaling molecule, but is known to affect calcium channels (Bubien et al., 1993). Rituximab binding to CD20, particularly when cross-linked, initiates a cascade of intracellular signals, many quite similar to signaling through the B cell receptor (Mathas et al., 2000). When CD20 is bound, it redistributes into cell membrane lipid rafts (Deans et al., 2002; Deans et al., 1998), which appears to be necessary for signal activation. The result of stimulation of these complex signaling pathways (i.e. cell cycle arrest or apoptosis) appears to be cell line specific.

Recent data from patients with chronic lymphocytic leukemia treated with rituximab, supports a role for the induction of apoptosis through the mitochondrial pathway (i.e. activation of caspases 3 and 9, but not caspase 8) and down-regulation of XIAP and mcl-1 (Byrd et al., 2002). Alterations in the set-point for apoptosis, perhaps by changes in levels of bcl-2 or its related family of anti-apoptotic proteins, apoptosome components such as APAF-1, or other regulators of apoptosis sensitivity could lead to antibody resistance and provide mechanisms that could be exploited to enhance activity.

4.1.3 Antibody-induced fixation of complement—Complement activation by the F_c portion of an antibody bound to a tumor cell surface can lead to complement-dependent cytotoxicity. Complement activation does take place in patients treated with rituximab, and it appears that the ability of CD20 to move into lipid rafts is required for complement-dependent cytotoxicity to occur (Cragg et al., 2002). Protein fragments indicative of complement activation are measurable in patients. Complement activation has effects other than lysis, including depositing C3, C3b and C3b breakdown products on the cell surface, which are potential targets of C3-receptors on effector cells, thereby enhancing cell killing by ADCC (see above) (Kennedy et al., 2003). Complement-mediated lysis is regulated by a series of complement inhibitory proteins, including CD35 (complement receptor type, CR1), CD46 (membrane cofactor protein), CD55 (decay accelerating factor) and CD59 (membrane inhibitor of reactive lysis, protectin) (Charles and Foerster). CD55 and CD59 inhibition may enhance antibody-mediated complement-dependent cytotoxicity (Golay et al., 2001; Golay et al., 2000). Fludarabine, clinically used in chronic lymphocytic leukemia and indolent non-Hodgkin lymphoma, down-regulates CD55 and CD59 (Di Gaetano et al., 2001), a potential mechanism of synergy between fludarabine and rituximab.

Subsequent discussion will provide an overview of clinically active antibodies for treating different types of cancer and discuss the available efficacy data for these agents.

4.2 Antibody therapy of hematologic malignancies

4.2.1 Rituximab—The chimeric monoclonal antibody rituximab contains a murine derived Fab targeting CD20, a pan-B cell transmembrane protein, and a human Fc segment (Reff et al., 1994). Features making CD20 an attractive target are having high level expression that is not down-regulated by antibody binding and low plasma levels (Grillo-Lopez, 2000). Rituximab kills normal and malignant CD20+ cells, but fortunately depletion of normal B cells is minimally toxic. The major attributes of Rituximab and other antibodies described in this section are summarized in Table 1.

Single agent rituximab was approved for relapsed/refractory indolent CD20+ non-Hodgkin lymphoma based on a response rate in heavily pretreated patients of 48%, with median duration 13 months (McLaughlin et al., 1998). Given the lack of rituximab bone marrow suppression, full doses of chemotherapy and rituximab can be combined and uniformly give better outcomes than chemotherapy alone. Major randomized trials leading to the currently approved indications for rituximab include documented prolonged remissions as initial therapy for indolent non-Hodgkin lymphoma with rituximab-cyclophosphamide-vincristine-prednisone (R-CVP) (Marcus et al., 2005) or rituximab scheduled re-treatment following CVP (Hochster

et al., 2009),versus CVP alone; and prolonged progression-free and overall survival as initial therapy for aggressive B cell non-Hodgkin lymphoma with rituximab-cyclophosphamidedoxorubicin-vincristine-prednisone (R-CHOP) vs CHOP alone (Coiffier et al., 2002; Habermann et al., 2006; Pfreundschuh et al., 2006; Sehn et al., 2005). Maintenance dosing of rituximab seems to prolong progression-free survival in indolent non-Hodgkin lymphoma, though dose and schedule have been variable and effects on survival require longer follow-up.

Each of the three main proposed mechanisms for rituximab-mediated cytotoxicity might be manipulated to enhance cytotoxicity. Cell signaling after rituximab is complex and under active investigation (Bonavida, 2007), but could be altered by signal pathway inhibitors. Any clinical effects, however, remain empiric. Complement activity may be enhanced by down-regulating complement inhibitory proteins (Manches et al., 2002) such as CD55 and CD59, which is an effect of fludarabine, raising this as a possible synergistic interaction (Di Gaetano et al., 2001). ADCC effects reflect Fc receptor binding affinities as CD16a polymorphisms alter rituximab single agent effects (Cartron et al., 2002; Weng and Levy, 2003a), however, R-chemo seems to overcome this effect (Carlotti and Pulsoni, 2007). Strategies to enhance ADCC, such as blocking inhibitory signals to killer cells, are under investigation (Binyamin et al., 2008).

Combination of active agents that also are immune modulating, such as lenalidomide and thalidomide, with rituximab may be additive (Kaufmann, 2004), and perhaps synergize by increasing ADCC (Reddy, 2008). The ability to restore the effector cell compartments with cytokines such as interleukin-2, -12 or -15 and myeloid growth factors, or potentially with cellular replacement therapy, may also enhance therapeutic antibody activity, as suggested by pre-clinical data demonstrating that IL-2 can promote NK cell development and enhance rituximab activity (Hooijberg et al., 1995) and clinical trials (Friedberg et al., 2002) (Khan, 2006). Myeloid growth factors in combination with rituximab may also activate ADCC (van der Kolk, 2003). Alternative approaches to induce effector cell activity by combining TLR agonists, such as CpG oligonucleotides, are also being investigated (Warren et al., 2000).

Since most pathways of cytotoxicity ultimately require apoptosis, altering the balance of proand anti-apoptotic signals could generate more cytotoxicity. As anti-apoptotic bcl2 is important in promoting survival of chronic lymphocytic leukemia and non-Hodgkin lymphoma tumors, bcl-2 down-regulation by antisense oligonucleotides was found to enhance rituximab efficacy in pre-clinical testing (Ramanarayanan et al., 2004; Smith et al., 2004). Clinical trials of the bcl-2 antisense, oblimersen, in combination with rituximab have been promising, however, small molecules that bind to the BH-3 domain common to many members of the bcl-2 family of proteins may be better therapeutic agents (Nguyen, 2007; Paoluzzi, 2008; van Delft, 2006).

4.2.2 Additional anti-CD20 antibodies under development—Rituximab can cause infusion related reactions, development of human anti-mouse antibodies and eventually clinical rituximab resistance does develop. The humanized anti-CD20 antibodies veltuzumab (Stein, 2004), ocrelizumab and ofatumumab (Coiffier et al., 2008) may decrease human anti-mouse antibody development and reduce infusion-related reactions. Ofatumumab also binds a different CD20 epitope and is engineered for better complement activation, although it induces less ADCC. Other engineered antibodies enhance ADCC, but activate less complementdependent cytotoxicity (Bowles, 2006).

4.2.3 Additional antibodies to treat non-Hodgkin lymphoma—CD22 is a pan B cell marker internalized upon antibody binding. Epratuzumab is a humanized anti-CD22 that exhibits modest activity alone and combined with rituximab (Leonard JP et al., 2004) (Leonard JP and T, 2005) (Strauss, 2006). Radiolabeled epratuzumab is also being studied (Lindén and Ohlsson, 2005). Building on the success of an anti-CD33-chalicheamycin drug conjugate in treating myeloid leukemia, and since CD22 is internalized upon antibody binding, the anti-

CD22-chalicheamycin drug conjugate inotuzumab ozogomicin, is well tolerated and active in relapsed/refractory B cell non-Hodgkin lymphoma as a single agent (Fayad, 2006) and combined with rituximab. CD22-pseudomonas immunotoxins are also active in treating the indolent B cell malignancy hairy cell leukemia. (Kreitman, 2001) The pan-B cell antigen CD19, for reasons that remain elusive, has not proven to be as effective a target as CD20 (Stone, 1996) (Grossbard, 1999). Further research targeting CD19 is ongoing. CD74, an MHC class II chaperone (Stein, 2007) with survival signaling effects (Starlets, 2006; Stein et al., 2009) is expressed on most B cell malignancies including myeloma, and is targeted by the humanized anti-CD74 antibody milatuzumab.

Humanized alemtuzumab recognizes CD52, which is expressed on B, T and NK cells, monocytes and macrophages, and has been found to be quite immunosuppressive. Approved for chronic lymphocytic leukemia therapy (Demko, 2008), it is also active in T cell diseases such as T-prolymphocytic leukemia (Keating and Maughan, 2002) and cutaneous T cell lymphoma.

4.2.4 Antibodies targeting lymphocyte activation markers—CD23 is expressed on B-chronic lymphocytic leukemia cells, as well as a normal subset of B cells. The chimeric macaque-human anti-CD23 antibody lumiliximab is active against chronic lymphocytic leukemia (Byrd and Wynne, 2007) and has been combined with standard chemotherapy in recent studies (Byrd, 2006). CD25, a component of the high affinity IL-2 receptor expressed on a variety of B and T cell lymphomas, has been targeted by a single chain anti-CD25 Fv fragment-pseudomonas exotoxin conjugate, which is active against CD25+ lymphomas (Kreitman and Waldmann TA, 2000). An IL-2-diptheria toxin conjugate, which also targets CD25, is already approved for use in chronic T cell leukemia and is active against other B (Dang, 2004) and T (Dang, 2007) cell lymphomas. CD80, a B7 ligand receptor family transmembrane glycoprotein, is normally involved in T cell co-stimulatory pathways. It is constitutively expressed on many malignant B cells, Hodgkin lymphoma Reed-Sternberg cells, and transiently on activated B cells and antigen presenting cells. The primatized anti-CD80 antibody, galiximab, may function via typical anti-tumor mechanisms, but also by blocking stimulatory CD28-CD80 interactions, potentially affecting activation of Treg cells and cytokine production to influence the tumor microenvironment (Czuczman, 2005) (Leonard, 2007).

4.2.5 Antibodies targeting TNF-related pathways—CD30 is present on anaplastic large cell lymphoma and Hodgkin lymphoma Reed-Sternberg cells. Antibody binding to CD30 may signal apoptosis, inhibit proliferation or induce complement-dependent cytotoxicity or ADCC. Murine, chimeric and humanized anti-CD30 antibodies (Bartlett and Bernstein, 2008) (Ansell SM and Blanset D, 2007) are well tolerated with limited activity in treating anaplastic large cell lymphoma and Hodgkin lymphoma. Engineered forms of these antibodies having enhanced Fc-receptor binding or as immunoconjugates are in clinical trials. CD40 is important in normal B cell development and expressed on various B cell malignancies, including myeloma and Hodgkin lymphoma Reed-Sternberg cells. The CD40 ligand CD154 is present on activated T cells, and interaction with CD40 on B and dendritic cells activates antigen presentation. Thus, CD40 binding by antibodies may result in a variety of anti-tumor effects, including immune stimulation. CD40 ligand and anti-CD40 antibodies with agonistic or antagonistic effects are also being studied (Vonderheide and Sullivan, 2007) (Advani, 2006). The TNF family of receptors can activate killing, as seen with CD30 and CD40. TNF-related apoptosis inducing ligand (TRAIL) induces apoptosis through the TRAIL receptors DR4 (TRAIL-R1) and DR5 (TRAIL-R2) on tumor cells (Griffith, 1999). Recombinant forms of TRAIL, as well as agonistic anti-receptor antibodies, induce apoptosis of tumors and are currently in clinical trials (Younes, 2005).

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4.2.6 Immunomodulatory antibodies—Along with accumulating evidence that patients with cancer may mount an immune response to their tumors (Hudson et al., 2007; Jager, 2007; Wang et al., 2005), tumors also create an immune suppressive environment locally and systemically that could inhibit such an immune response. Many mechanisms have been proposed for these inhibitory effects. Modulation of co-stimulatory and co-inhibitory molecules is thought to contribute to these immune subversion mechanisms. Examples include molecules and families of proteins such as B7-H1 (programmed death ligand-1) that interacts with the co-inhibitory programmed death receptor-1 (PD-1, CD279) on T cells (Blank et al., 2005) and cytotoxic T lymphocyte antigen-4 (CTLA-4) (Korman et al., 2006). A number of recently introduced candidate antibodies for cancer therapy specifically target these inhibitory receptors or their ligands (Korman et al., 2005; Sanderson et al., 2005).

Initiation of a tumor specific immune response requires the presentation of tumor antigen to T cells by an antigen presenting cell in the context of MHC molecules. T cell activation requires two events: recognition of antigen/MHC complex by T cell receptor and a co-stimulatory signal delivered through CD80 and CD86 on the antigen presenting cell surface interacting with CD28 on the T cell. This process results in T-cell activation and proliferation (Fong and Small, 2008; Korman et al., 2006). Following activation, CTLA-4, is expressed on the surface of T cells, binds CD80 and 86, and provides inhibitory signals to the T cell as part of a negative feedback loop (Fong and Small, 2008). The negative regulation of the immune response maintains peripheral tolerance to self-antigens and prevents damage to normal host tissue (Korman et al., 2006). Theoretically, blocking CTLA-4 with mAbs may sustain the activation and proliferation of tumor specific T cells, promoting the development of an effective tumorspecific immune response (Fong and Small, 2008).

The exact mechanism of action of anti-CTLA-4 antibodies has not been determined. Kavanagh and colleagues recently showed that treatment with the CTLA-4 antibody, ipilimumab, can expand functional Treg cells in humans (Kavanagh et al., 2008). Alternatively, CTLA-4 blockade could act directly on effector $CD4^+$ and/or $CD8^+$ T cells by removing a checkpoint on their proliferation and function (Fong and Small, 2008).

A co-stimulatory molecule, CD137 (4-1BB), has also been studied for its potential role in immune modulation. This molecule belongs to the TNF receptor family and is expressed on multiple immune cell subsets, particularly following their activation. Ligation of CD137 using agonist anti-CD137 monoclonal antibodies (mAbs) leads to co-stimulation of T and NK cells and eradication of established tumors in multiple murine models (Melero et al., 1997; Miller et al., 2002; Nam et al., 2005; Wilcox et al., 2002). Clinical studies of BMS-663513, an IgG4 anti-human CD137 mAb, are ongoing in patients with solid tumors, with objective responses seen, primarily in melanoma (Sznol M., 2008).

PD-1 is a member of the CD28 superfamily and plays an important role as an inhibitor of T cell responses (Greenwald et al., 2005). A higher content of PD-1⁺ cells predicted improved survival in patients with follicular lymphoma (Carreras et al., 2009), while patients with higher grade follicular lymphoma and poor prognostic features had a lower level of PD-1+ cells. Blank and colleagues reported that the interaction between PD-1 and its ligand, PD-L1, negatively regulated T cell effector functions (Blank et al., 2006). These investigators showed that PD-L1 was constitutively expressed on human renal cell carcinoma (RCC) cell lines and upregulated on human melanoma cell lines upon exposure to IFN-γ. PD-L1 expression was found on frozen sections from RCC and melanomas, but not on normal tissues. This supports the notion that PD-L1 is a mediator of tumor escape from tumor-specific T cells (Blank et al., 2006).

Additional trials are ongoing with these and other immunomodulatory antibodies. A better understanding of the mechanism of action of these molecules along with better knowledge regarding their potential side effects is needed before they can achieve widespread use in the clinic, however.

4.3 Antibody therapy of solid tumors

4.3.1 Trastuzumab—Trastuzumab (Herceptin®) HER-2/*neu* (c-erbB-2), a member of the epidermal growth factor receptor (EGFR) family, has been targeted for antibody therapy because it is overexpressed on 25% of breast cancers, as well as other adenocarcinomas of the ovary, prostate, lung, and gastrointestinal tract. Trastuzumab (Baselga et al., 1996; Cobleigh et al., 1999), a humanized antibody derived from the murine mAb 4D5, recognizes an epitope on the extracellular domain of HER-2/*neu.* Over-expression of HER-2/*neu* that is associated with gene amplification correlates with a clinical response to trastuzumab. Earlier studies utilized variable criteria to define HER2 expression, which may account for the differences in the response rates observed in these studies (von Mehren et al., 2003). The major attributes of Trastuzumab and other antibodies described in this section are summarized in Table 2.

Early studies with this agent established its role in patients with metastatic disease. A phase II study of trastuzumab, in women with metastatic breast cancer who were treated with 2 mg/kg weekly, showed an objective response rate of 16% (Cobleigh et al., 1999). Improvement in the median response rate and overall survival were noted.

A large, randomized phase III trial evaluating cytotoxic chemotherapy alone or with weekly trastuzumab has shown the efficacy of combination therapy (Slamon et al., 2001). Patients receiving initial therapy for metastatic breast cancer were treated with cyclophosphamide plus an anthracycline, or with paclitaxel if they had received an anthracycline in the adjuvant setting. Response rates for combination therapy were higher regardless of the chemotherapy used. Furthermore, addition of trastuzumab to chemotherapy improved survival at one year by 16% (Norton et al., 1999) and improved survival at 29 months by 25% (Baselga, 2001a). Myocardial dysfunction was observed more frequently in patients receiving doxorubicin or epirubicin when trastuzumab was added. Therefore, trastuzumab is not recommended in combination with anthracyclines. Based on these clinical trial results, trastuzumab was approved by the FDA to treat women with metastatic breast cancer with HER-2/*neu* overexpression, either alone or in combination with paclitaxel. Trastuzumab also has activity in combination with other chemotherapeutic agents (Burstein et al., 2001; Pegram et al., 1998). In addition two randomized trials have shown that trastuzumab in the adjuvant setting can lead to a 33% reduction in the risk of death (Romond et al., 2005).

Trastuzumab is a success story in breast cancer therapy. However, despite the overexpression of Her2/*neu* in other solid tumors, such as non-small cell lung cancer, to date no significant clinical benefit of trastuzumab has been seen in patients with diseases other than breast cancer.

4.3.2 Epidermal Growth Factor Receptor (EGFR) Antibodies—EGFR is a widely expressed glycoprotein with three main domains: The extracellular ligand binding domain can bind to a number of regulatory factors, such as EGF and transforming growth factor-α (TGFα), leading to EGFR dimerization and autophosphorylation. The hydrophobic transmembrane portion connects the extra- and intracellular portions. The cytoplasmic domain responds to receptor autophosphorylation by increasing the tyrosine kinase catalytic activity (Lockhart and Berlin, 2005). This, in turn, leads to phosphorylation of other tyrosine motifs and activation of a number of intracellular signaling pathways such as phosphatidylinositol-3 kinase (PI3K) and Akt, which result in cellular proliferation, differentiation, transformation and anti-apoptotic functions.

The possible role of EGFR overexpression in tumor growth and development has been best studied in colorectal cancers (Lockhart and Berlin, 2005). It is generally accepted that EGFR overexpression is found in 65% to 70% of colorectal carcinomas (Messa et al., 1998), even though there is heterogeneity in tumor biopsies and lack of a uniform method of evaluating levels of EGFR. The available anti-EGFR monoclonal antibodies bind to the receptor and block the binding of the ligand to prevent downstream events. Cetuximab (C225) is a chimeric monoclonal antibody directed against the extracellular domain of the EGFR. Panitumomab (ABX-EGF) is a fully human IgG2 monoclonal antibody against EGFR generated through the Xenomouse transgenic strain in which human immunoglobulin genes replace mouse immunoglobulin genes (Lockhart and Berlin, 2005). Both of these antibodies are capable of inhibiting the growth of EGFR expressing cell lines in vitro and in vivo (Baselga, 2001b; Yang et al., 2001). Cetuximab is capable of enhancing the anti-tumor effects of chemotherapy and radiation therapy in preclinical and clinical studies. Proposed mechanisms by which anti-EGFR mAbs can elicit an anti-tumor response include receptor internalization and subsequent degradation, interference with EGFR signaling by preventing ligand binding and induction of ADCC. Another proposed mechanism that is less well understood invokes interference with signaling events via antibody disruption of lipid rafts. The EGFR is intimately associated with signaling microplatforms on the cellular membrane (Puri et al., 2005). High concentration of EGFR ligands or, perhaps, cross-linking with an antibody may shuttle EGFR internalization from the clathrin-mediated pathway of recycling to the caveolin-associated route leading to receptor ubiquitination and degradation (Sigismund et al., 2005).

Cetuximab has been used either alone or in combination with radiotherapy or various chemotherapeutic agents primarily in patients with head and neck or colorectal cancers. The most common side effects seen with this agent are an acneiform rash and occasional allergic reactions. A number of earlier trials showed efficacy of cetuximab as a single agent or in combination with chemotherapy. Although single agent cetuximab activity in colorectal cancer is low, a response rate of 22.5% was achieved in combination with irinotecan (Saltz L et al., 2001; Saltz et al., 2004). A subsequent randomized phase III trial involving 329 patients with EGFR+, irinotecan-refractory colorectal cancers showed a significantly higher response rate (22.9 versus 10.8%) and median time to progression in the combination arm (Cunningham et al., 2004). Cetuximab has been approved by the FDA for the treatment of advanced or refractory colorectal cancers, in conjunction with irinotecan.

The first randomized clinical trial to demonstrate a statistically-significant survival benefit by use of any EGFR antagonist is in squamous cell carcinoma of the head and neck. In the pivotal multi-center trial in head and neck squamous cell carcinoma (Bonner et al., 2006), 424 patients underwent randomization to definitive radiotherapy alone or in combination with weekly cetuximab. Bonner et al. reported doubling of median duration of locoregional control from 14.9 to 24.4 months (log-rank P=0.005). The 3-year survival rates were 54% versus 45% in favor of cetuximab and radiotherapy. Locoregional control and survival benefits persist (Robert et al., 2001). Of interest, there was no influence of EGFR expression on patient outcome.

In advanced stage non-small cell lung cancer expressing EGFR, the combination of cetuximab and standard doublet chemotherapy has been studied in a prospective randomized phase III trial (Pirker et al., 2009). Addition of cetuximab was associated with a slight, but statistically significant, benefit in overall survival over chemotherapy alone (median overall survival 10.1 vs. 11.3 mos, P=0.0441).

4.3.3 Vascular endothelial growth factor (VEGF) antibodies—Pioneering work of Folkman and others (O'Reilly et al., 1994) showed that malignant tumor progression requires new blood vessels capable of delivering essential nutrients and oxygen and blockade or damage of tumor vessels deprives malignant cells of essential nutrients and causes tumor destruction.

Since one vessel supplies a vast number of cancer cells, this approach can amplify the consequences of an initial cytotoxic insult. Moreover, this strategy does not require the penetration of other cells or molecules into the tumor parenchyma.

Of the identified angiogenic factors, VEGF is the most potent and specific regulator of both normal and pathologic angiogenesis (Ferrara and Davis-Smyth, 1997). VEGF produces a number of biologic effects, including endothelial cell mitogenesis and migration, induction of proteinases leading to remodeling of the extracellular matrix, increased vascular permeability, and maintenance of survival for newly formed blood vessels (Ferrara and Davis-Smyth, 1997). The biologic effects of VEGF are mediated through binding and stimulation of two receptors on the surface of endothelial cells: Flt-1 (fms-like tyrosine kinase, VEGFR1) and KDR (kinase domain region, VEGFR2) (Davis-Smyth et al., 1996; Ferrara and Davis-Smyth, 1997).

Increased expression of VEGF has been demonstrated in most human tumors examined to date, including tumors of the lung, breast, thyroid, gastrointestinal tract, kidney, bladder, ovary, and cervix, as well as angiosarcomas and glioblastomas (Ferrara and Davis-Smyth, 1997). Inhibition of VEGF by using an anti-VEGF monoclonal antibody blocks the growth of a number of human cancer cell lines in nude mice (Ferrara and Davis-Smyth, 1997). In addition, the combination of anti-VEGF antibody and chemotherapy in nude mice injected with human cancer xenografts results in an increased antitumor effect compared with antibody or chemotherapy treatment alone (Borgstrom et al., 1999).

Bevacizumab is a recombinant humanized monoclonal antibody targeting VEGF (Presta et al., 1997). Clinical trials including patients with a variety of tumor types: colorectal, breast, lung, and renal carcinoma have established the role of this monoclonal antibody in the treatment of metastatic cancer (D'Adamo et al., 2005; Hurwitz et al., 2004b; Hurwitz et al., 2005; Miller et al., 2005b). In a randomized phase III study in patients with metastatic colorectal cancer, Hurwitz and colleagues assigned 815 patients to receive either irinotecan, 5-flourouracil and leucovorin (IFL) with bevacizumab or IFL with placebo (Hurwitz et al., 2004a). The addition of bevacizumab led to a statistically significant increase in the response rate and a 4.7-month prolongation in median overall survival (to 20.3 months, vs. 15.6 months with IFL and placebo). In this study, bevacizumab was associated with reversible hypertension and proteinuria and was relatively well tolerated.

A randomized phase III trial (ECOG 4599) of paclitaxel and carboplatin with or without bevacizumab in patients with advanced non-squamous non-small cell lung cancer led to a significant improvement in median survival (12.5 months vs. 10.2 months; $P = .0075$) for patients in the bevacizumab arm. In addition, patients treated with bevacizumab had a significantly higher response rate $(27\% \text{ vs. } 10\%; P < .0001)$ and a significantly longer progression-free survival time $(6.4$ months vs. 4.5 months; $P < .0001$). A higher incidence of bleeding was associated with bevacizumab (4.5% vs. 0.7%). Five of ten treatment-related deaths occurred as a result of hemoptysis, all in the bevacizumab arm.

A phase III trial randomized 722 patients with metastatic breast cancer with no prior chemotherapy for advanced disease to either paclitaxel or paclitaxel and bevacizumab (Miller et al., 2007b). Progression free survival was significantly better in the paclitaxel plus bevacizumab arm (median, 11.8 vs. 5.9 months; hazard ratio for progression, 0.60; P<0.001) with an increased response rate (36.9% vs. 21.2%, P<0.001). Overall survival, however, was similar in the two groups (median, 26.7 vs. 25.2 months; hazard ratio, 0.88; P=0.16).

Miller and colleagues (Miller et al., 2005c) reported the results of a randomized phase III trial of capecitabine/bevacizumab versus capecitabine alone in 462 anthracycline and taxane pretreated metastatic breast cancer patients. The primary end point of this study, progression-

free survival, did not show a statistically significant difference (4.86 vs. 4.17 months; hazard ratio = 0.98). Combination therapy significantly increased the response rates (19.8% vs. 9.1%; $P = .001$). Overall survival (15.1 vs. 14.5 months) and time to deterioration in quality of life were comparable in both treatment groups.

Five major bevacizumab-associated toxicities have been identified: hypertension, proteinuria, thromboembolic events, hemorrhage and congestive heart failure. The rates of clinically meaningful events have been low, however, and because of these potential adverse events, a number of patients with advanced disease are not eligible for treatment with this agent.

4.4 Radioimmunotherapy

Unlabeled antibody may not have access to all tumor cells nor kill all targeted cells. Many factors affect cell killing, including surface antigen density, antibody penetration into lymph nodes or tumors, and local accessibility to complement and cytotoxic cells. Using antibodydrug conjugates may overcome low antigen density and lack of necessary local immune cells. Several of these are discussed above. Alternatively, radioactively-tagged antibodies can target adjacent cells and may not require immune functions. The two approved radioimmunotherapy agents, Y^{90} -ibritumomab tiuxetan and 131 I-tositumomab appear to have similar efficacy (Davies, 2007; Jacene, 2007). Response rates are high with toxicity mainly delayed by myelosuppression, (Fisher, 2005).(Horning, 2005) (Kaminski, 2005; Witzig et al., 2002). Radioimmunotherapy has been used to consolidate remissions after chemotherapy with promising results (Press, 2007) (Leonard, 2005) (Smith, 2007). Another research area utilizes radioimmunotherapy prior to stem cell transplant (Gopal, 2007; Krishnan, 2008). Radioimmunotherapy utility may be improved by re-treatment, radiosensitizing agents and pre-targeting (Goldenberg, 2008). Radioimmunotherapy responses in rituximab-refractory patients suggests a different mechanism of action, i.e. targeted radiation therapy, rather than antibody-dependent killing. Radioimmunotherapy resistance may therefore indicate an intrinsic altered apoptotic threshold of the lymphoma cell, raising the possibility of overcoming this as another potential therapeutic strategy.

5. Concluding remarks

While targeted immune therapies offer new possibilities to harness the immune response to treat cancer patients, effective manipulation of the immune system may require overcoming barriers, while avoiding potential hazardous complications. Barriers include elimination of the tumor-specific antigen or overcoming immune tolerance within the tumor microenvironment, which can be due to the production of immunosuppressive cytokines (such as IL-10, TGF-β, and type 2 cytokines) or the presence of immuosuppressive cells (such as tumor-associated macrophages, myeloid-derived suppressor cells, and Treg cells). Although not detailed here, many immune-potentiating therapies designed to target these immunosuppressive molecules and cells are also under development. On the other hand, the removal of normal tolerance mechanisms can also trigger autoimmune responses or graft-versus-host disease, which can cause significant and even life-threatening complications. Immune tolerance mechanisms play important roles in maintaining homeostasis and preventing autoimmunity, and altering these mechanisms can have unexpected negative repercussions in patients. Furthermore, it must be recognized that therapies designed to establish adaptive immune responses toward a tumor mass may not effectively target the minor subpopulation of cancer stem cells that support tumor development and recurrence, since those progenitor cells may express distinct antigens. Therefore, future targeted immunotherapies may be necessary that specifically direct immune reactivity toward these cancer stem cells.

One must also consider that many of the radiation and chemotherapeutic agents used to treat cancer can significantly suppress normal immune function. Therefore, immunotherapies must

precede treatment with immunosuppressive anti-cancer agents, since it may be impossible to potentiate a compromised immune system. On the other hand, tumor-targeted radiation treatments do not result in generalized immunosuppression, and may actually enhance antitumor immune responses by releasing antigens from dying tumor cells for uptake and presentation by antigen presenting cells (Ferrara et al., 2009). Immunotherapy will likely not be able to eliminate tumors alone, but combination therapies that incorporate immunotherapeutic agents have great potential for providing clinical success in treating cancer in the coming years.

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Figure 1. Major cells of the innate and adaptive immune systems and their functions in response to a tumor cell

Grey boxes delineate the major cell types constituting the innate and adaptive immune systems. Arrows describe impacts of immune cells on each other or the tumor. Note that this is a very basic schematic designed to define interactions of relevance to the review, and numerous complex molecular interactions and minor immune cell subsets further influence these functions.

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Figure 2. Minor subsets of T cells and their cytokine products Distinct T cell subsets are shown with the major cytokines that they can produce to influence other immune cells or tumors.

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Figure 3. Surface receptor interactions establishing the basis for tolerance and cytolytic responses by NK cells

NK cell activation is regulated by a balance of signals derived from activating and inhibitory surface receptors. Activating receptors recognize an array of cell surface molecules, some of which are expressed ubiquitously on the surfaces of other cells of the body. Inhibitory receptors, such as KIR, recognize MHC class I (MHC-I) molecules, which are expressed on the surface of virtually every normal cell of the body, while many tumor cells lack expression of this "self" molecule. KIR engagement with MHC-I triggers inhibitory signals (predominantly protein tyrosine phosphatase-based) within the NK cell that dominantly suppress activating receptor signals (predominantly protein tyrosine kinase-based) and result in tolerance. Upon encounter with a rare tumor cell lacking MHC-I (so-called "missing self"), the lack of inhibitory signaling at the cell contact interface triggers rapid and directed release of cytolytic granules toward the tumor target cell, which results in specific cytolysis of the target cell.

Table 1

Antibodies for Hematologic Malignancies

*** Indicates approved agents. Remainder are investigational.

Table 2

Antibodies for Solid Tumors

