

definitive conclusions on the role of Th2 cells in GvHD from this small study was difficult, Th2 and non-Th2 recipients showed similar rates of acute and chronic GvHD with a similar organ distribution.

Th17

Th17 & Th1/Th2 revisited in autoimmune diseases

Classically, autoimmune diseases have been assumed to be associated with dysregulated Th1 responses on the basis of findings from animal models of autoimmunity such as experimental autoimmune encephalomyelitis (EAE). EAE is a mouse model of multiple sclerosis. However, accumulating evidence argues against the role of Th1 cells in the pathogenesis of EAE. EAE was more severe in IFN- γ -deficient mice than wild-type mice [29], whereas IL-23p19-deficient mice, in which Th1 cells were not reduced, were resistant to EAE [30]. Subsequent studies pointed to the role of IL-23 as an inducer of IL-17 and demonstrated that IL-17-producing Th17 cells are primarily responsible for EAE induction [31,32]. Deficiency of IL-17 or ROR- γ t, a key regulator of Th17 development, attenuates EAE [32,33]. Similarly, blockade of IL-17 ameliorated experimental arthritis [34]. In patients with rheumatoid arthritis, IL-17 levels were elevated in synovial fluids [35]. Infiltration of Th17 cells was observed in the biopsied samples of chronically inflamed tissues in patients with psoriasis, Crohn's disease, rheumatoid arthritis and asthma [36]. These results clearly indicate that Th17 immunity plays a central role in many autoimmune diseases.

Differentiation & function of Th17 cells

Th17 cells are characterized by their expression of proinflammatory cytokines IL-17, IL-21, and IL-22, cytokines involved in neutrophilia, production of antimicrobial peptides and tissue repair [13,31,37,38]. IL-17 belongs to a family of six members: IL-17A, IL-17B, IL-17C, IL-17D, IL-17E (also known as IL-25) and IL-17F. Among them, IL-17A and IL-17F, which form heterodimers, are by far the best characterized cytokines [39]. IL-17 is also produced by innate immune cells in an inflammatory milieu, and may have a central role in the initiation of IL-17-dependent immune responses even before Th17 development [40].

Th17 cells differentiate from naive CD4⁺ T cells by stimulation with antigens in the presence of TGF- β and IL-6 or IL-21 in mice (FIGURE 1)

[13–15,38,41–43]. In humans, different Th17 differentiation factors have been initially reported, but factors inducing Th17 cells in mice and humans seem to be identical [41,44–49]. IL-21 is produced by Th17 cells, and thus, it is a part of a positive feedback loop for amplifying Th17 immunity [38]. Naive T cells do not express IL-23R, except on activation, and IL-23 stabilizes differentiating Th17 cells and promotes further maturation of Th17 cells in part by inducing IL-22 in these cells [13,50]. Specific transcriptional factors such as ROR- γ t, ROR- α and STAT3 define the Th17 transcriptional program because they are indispensable for Th17 differentiation [33,43,51–53]. In addition, Th17 cells can be induced in memory T cells by IL-23 and IL-1 β [13,31,49].

Th17 cells are found in the systemic circulation, secondary lymphoid organs and tissues, particularly in the intestinal mucosa where they protect the host from microorganisms that invade through the epithelium [54]. Th17 differentiation is influenced by innate immunity and by the composition of the intestinal microbiota. Stimulation with binding of bacteria-derived peptidoglycan to Toll-like receptor 2 on dendritic cells (DCs) in the gut induces production of IL-23 and IL-1 β , leading to Th17 differentiation [41]. Luminal ATP and flagellin are critically involved in this process [55,56]. Th17 cells contribute to host defense against certain pathogens by inducing expression of antimicrobial factors and recruiting neutrophils to mucosal surface [57]. They also play a role in maintaining intestinal ecology with commensal bacterial flora [58]. On the flip side, Th17 differentiation is influenced by the composition of the intestinal microbiota; the commensal microbe segmented filamentous bacterium increases Th17 cells in the lamina propria of mice, whereas segmented filamentous bacterium is absent in mice with few Th17 cells [54]. However, such a specific bacterium that can induce Th17 differentiation in the lamina propria of humans remains to be identified. Importantly, broad receptor distribution of IL-17 and IL-22 results in massive tissue responses to Th17 cells; these cells trigger local tissues to release IL-6, TNF- α , IL-1 β and chemokines and recruit neutrophils and macrophages to inflammatory sites [57].

Regulatory T cells

Regulatory T-cell subsets

T-cell-mediated immunoregulation is one of the main mechanisms responsible for maintaining antigen-specific operational tolerance *in vivo*, controlling T-cell homeostasis, and

the homeostatic expansion of T cells in lymphopenic hosts. Although many lymphocyte populations that can suppress antigen-specific immune responses have been described, the best described *bona fide* populations of regulatory cells are CD4⁺25⁺FOXP3⁺ Tregs [16,17]. FOXP3 is an oligomeric component of a large supermolecular complex and critical regulator of the development and function of Tregs [59,60]. Certain *FOXP3* gene mutants are associated with X-linked autoimmunity–allergic dysregulation syndrome and immunodysregulation, polyendocrinopathy and enteropathy X-linked syndrome in humans [61,62]. Individuals with these disorders fail to develop Tregs and experience varied symptoms that include diarrhea, dermatitis, insulin-dependent diabetes, thyroiditis and anemia, with massive T-cell infiltration into the affected tissue. There are other types of CD4⁺FOXP3⁻ Tregs; Tr1 cells are generated *in vitro* by antigenic stimulation in the presence of IL-10 and secrete IL-10 and TGF- β [63]. A subset of CD8⁺ T cells also suppresses CD4⁺ T-cell responses with phenotypic and functional features similar to CD4⁺ Tregs in mice and human [64–67]. However, little is known regarding the role of CD8⁺ suppressor cells in GvHD and this article will focus only on CD4⁺ Tregs.

Development & function of Tregs

Tregs can be divided into two populations: naturally occurring Tregs (nTregs) that develop in the thymus and induced Tregs (iTregs) that are derived from CD4⁺FOXP3⁻ T-cell population in the periphery (FIGURE 1) [68]. Thymic development of nTregs requires high-affinity interactions between their TCR and self-peptide–MHC complexes expressed in thymic stromal cells. Differentiation of conventional CD4⁺ T cells (Tcons) into iTregs in the periphery represents the outcome of cytokine-mediated activation of specific STAT proteins and the induction of lineage-specific transcriptional factors, and depends upon TGF- β .

Tregs are indispensable in maintaining self-tolerance and immune homeostasis, establishing tolerance to allografts and promoting fetomaternal tolerance. The disadvantage of Tregs is dampening of antitumor T-cell responses [69]. Tregs suppress effector activities of differentiated CD4⁺ and CD8⁺ T cells and the function of NK cells, NKT cells, B cells, macrophages and DCs. Tregs must be activated through their TCR to be functionally suppressive and IL-2 signaling is required for their expansion and function [70,71]. Tregs suppress immune responses through

contact-dependent mechanisms and the production of soluble factors, including TGF- β , IL-10 and IL-35 [17]. Contact-dependent suppression is mediated through multiple molecules such as cytotoxic T lymphocyte antigen (CTLA)4, lymphocyte activation gene 3, granzyme A and CD95. Naive Tregs do not express these molecules, except on activation. While CTLA-4 expression on Tregs is critical for suppression by modulating APCs, Tregs can also suppress effector T cells through direct contact even in the absence of APCs [72]. The mechanisms of cytokine-mediated suppression include FOXP3 induction in Tcons and attenuation of DC function by TGF- β and IL-10 [68].

Human Foxp3⁺ CD4⁺ T cells are composed of three phenotypically and functionally distinct subpopulations: CD45RO⁺FOXP3^{lo} naive Tregs, CD45RO⁺Foxp3^{hi} effector Tregs, and CD45RO⁺FOXP3^{lo}-activated Tcons that are pathogenic because they produce proinflammatory cytokines [73–75]. Thus, FOXP3 is not always a reliable marker for human Tregs: Foxp3⁺ T cells are heterogeneous, including nonregulatory T cells and naive Tregs that only weakly express Foxp3.

Plasticity of effector & Tregs subsets

Regulatory and effector T-cell subsets are found to be more malleable than originally thought on the basis of the more stringent commitment of Th1 and Th2 cells, and flexibility in T-cell commitment is not an exception [76]. Th17 cells have the potential to convert toward a Th1 profile in mice and humans [77]. Th17 cells isolated from a certain strain of mice convert into Th1 cells after transfer into another strain of mice probably because of the difference in cytokine environment *in vivo* [78]. Th17 cells can shift to a Th1 phenotype in the presence of IL-12 [79]. This conversion occurs through expression of Th1 transcription factors STAT4 and T-bet that also antagonize Th17 development [13].

A common requirement of TGF- β in the differentiation of Th17 cells and iTregs suggests that these subsets may be developmentally linked and Th17 and Treg development programs are reciprocally interconnected. TGF- β is a pleiotropic cytokine that affects multiple cell lineages by promoting or opposing their differentiation, survival and proliferation. TGF- β inhibits Th1 and Th2 differentiation, whereas it induces Th17 and Treg differentiation [80]. Transcriptional factors involved in driving FOXP3 expression may indirectly regulate Th17 development, because FOXP3 inhibits the function of Th17-associated

transcriptional factor ROR- γ t [81]. The local cytokine milieu is of great importance for Treg function and differentiation. Treg generation from antigen-stimulated T cells is promoted by the environment rich in TGF- β and retinoic acid at the expense of Th17 differentiation [82]. Tregs are reciprocally induced by exclusion of IL-6 from the Th17-skewing condition in mice and humans [83], whereas Tregs are converted to IL-17-producing cells when reactivated under Th17-polarizing conditions [84–88] or even under normal conditions [89]. A lymphopenic environment also leads to the loss of FOXP3 expression in Tregs and their differentiation into T follicular helper cells in Peyer's patches [90]. Furthermore, Foxp3⁺ T cells that produce IL-17, IFN- γ or IL-4, and those express ROR- γ t or T-bet have been demonstrated in mice and humans [81,91–97].

For the maintenance of the developmentally established Treg function program, continuous expression of Foxp3 is an absolute requirement. Committed Tregs are resistant to such conversion to effector T cells upon adoptive transfer into lymphopenic hosts, whereas a relatively minor fraction of uncommitted CD25⁺ or CD25^{lo} FOXP3⁺ T cells can lose FOXP3 expression and divert into effector T-cell lineages [98]. By contrast, a recent study using inducible labeling and tracking the fate of Tregs *in vivo* demonstrated that Tregs are remarkably stable under physiological and inflammatory conditions, and that Treg homeostasis is maintained by self-renewal of established mature Tregs [99]. Although that study does not exclude the possibility that recently generated FOXP3⁺ cells can differentiate into effector T cells in a lymphopenic or proinflammatory environment, it argues against plasticity of once committed Tregs. This issue is fundamentally important for those studies directed to manipulate immune responses in a therapeutically useful manner.

Th17 & Tregs in experimental GvHD

■ Th17

Research on Th17 in GvHD is rapidly developing. The role of Th17 responses in GvHD has been evaluated using IL-17-deficient mice; however, it has not yet been completely established. Yi *et al.* showed that infusion of IL-17-deficient donor T cells induced more severe GvHD than that of wild-type T cells [100]. This was in association with enhanced Th1 differentiation of donor T cells by increased production of IL-12 from DCs in the absence of IL-17. By contrast, two other studies failed to show the effects of

IL-17 on GvHD mortality [101,102]. Differences between experimental designs such as strain combinations, intensity of conditioning irradiation and T-cell dose may result in inconsistent results. Furthermore, ROR- γ t-deficient T cells can also induce acute GvHD as severe as that observed with wild-type T cells [103]. IL-17 is not required for transition from acute to chronic GvHD [104]. These results suggest that Th17 differentiation is at least dispensable for GvHD. Yi *et al.* further showed that liver and gut GvHD is largely Th1 dependent, whereas skin GvHD is more Th17 dependent and lung GvHD is Th2 dependent [105]. Thus, each T-helper subset may play a differential role in inducing tissue-specific GvHD; Th17 responses may be involved in skin GvHD.

Recently, Hill *et al.* demonstrated that donor pretreatment with granulocyte colony-stimulating factor (G-CSF) induced Th17 differentiation of donor T cells after PBSCT [106]. PBSCT from IL-17-deficient donors treated with G-CSF significantly reduced skin scleroderma in association with reduced infiltration of macrophages that are important sources of TGF- β , a mediator of fibrosis [107,108]. These findings are highly relevant clinically as PBSCT from G-CSF-mobilized donors is a risk for chronic GvHD compared with bone marrow transplant in humans [109]. However, the role of IL-17 and TGF- β in human chronic GvHD has not yet been identified.

The ability of *ex vivo*-polarized Th17 cells to induce GvHD in each target organ has also been evaluated [103,110]. Th17 cells were generated by repetitive stimulation of naive CD4⁺ T cells with CD3/CD28 in the presence of TGF- β , TNF- α , IL-6, IL-1 β and IL-23, as well as anti-IL-2 monoclonal antibodies (mAbs) and anti-IFN- γ mAbs [110]. On transfer into lethally irradiated mice, these Th17 cells migrated and expanded in GvHD target organs and lymphoid tissue, and subsequently induced more severe pathological lesions of the skin and lung, but equivalent pathological lesions of the liver and intestine in comparison with naive CD4⁺ T cells [110]. Unexpectedly, however, IL-17 blockade ameliorated skin GvHD but had no impact on the mortality and morbidity of systemic GvHD. These Th17 cells produced TNF- α and IFN- γ in addition to IL-17A and IL-17F after transfer. Neutralization of TNF- α significantly reduced the mortality and morbidity of systemic GvHD. Another study also confirmed such a central role of TNF- α in systemic GvHD mediated by polarized Th17 cells [111]. Iclozan *et al.* generated

Th17 cells by stimulating CD4⁺ T cells with anti-CD3 mAbs in the presence of TGF- β , IL-6, anti-IL-4 and anti-IFN- γ mAbs [103]. These Th17 cells had superior expansion and migration capabilities in GvHD target organs, which correlated with their increased pathogenicity, compared with Th1 or naive T cells.

Taken together, it seems that Th17 cells are sufficient but not necessary for GvHD induction. However, these studies in which *ex vivo*-polarized T cells were infused have limitations and results from these studies should be carefully interpreted. In these studies, Th17 cells are polyclonally expanded and therefore antigen non-specific. This is not clinically relevant because antigen-specific Th17 cells are supposedly generated *in vivo* after transplantation. Furthermore, incubation of T cells *in vitro* might result in incomplete polarization and partial anergy. Upon transfer, these cells may fail to survive and proliferate or cause secondary conversion to a mixed phenotype or to a different regulatory or effector T-cell subset, as has been stated in previous studies of polarized Th1 and Th2 cells. *In vitro*-generated Th17 cells do not essentially produce IFN- γ , but can produce IFN- γ upon restimulation without Th17-polarizing cytokines. On transfer, such Th17 cells did not maintain their cytokine profiles because they produced large amounts of IFN- γ , although IFN- γ was not required for Th17 cell-mediated GvHD in these studies [103,110]. Whether the production of IFN- γ by Th17-skewed cells is part of the evolution of the T-cell subset *in vivo* or whether IFN- γ is produced by a preformed Th1-like component that subsequently expands is unclear.

■ IL-6, IL-21 & IL-23 in GvHD

IL-6 is one of the central mediators of inflammation. It has a pivotal function in dictating whether T cells differentiate into Tregs or Th17 cells. IL-6-deficient mice do not develop a Th17 response and their peripheral repertoire is dominated by Tregs [42]. Serum IL-6 levels are often elevated during GvHD in mice and humans [112,113]. *IL-6* gene polymorphisms that lead to altered IL-6 expression levels are linked to GvHD severity [114]. Although IL-6 elevation is not specific for GvHD and the association of *IL-6* gene polymorphism and GvHD severity is controversial, these results suggest that IL-6 could be a target molecule to treat GvHD. A recent study showed that blockade of the IL-6 signaling pathway with anti-IL-6R mAbs ameliorates GvHD in association with augmentation

of thymic-dependent and thymic-independent production of Tregs with a concomitant reduction of Th1 and Th17 cells in GvHD target organs [112]. By contrast, Tawara *et al.* reported that anti-IL-6R mAb-mediated attenuation of GvHD was independent of the direct effects on effector T-cell expansion or donor Tregs [113].

IL-21 is produced by Th17 cells and its receptors are broadly expressed on T, B and NK cells. Inhibition of IL-21 signaling on donor T cells attenuates GvHD, particularly in the intestine [102,115,116]. Bucher *et al.* showed an increased number of Foxp3⁺ Tregs in the intestine of anti-IL-21 mAb-treated recipients of Foxp3⁻ T cells, suggesting Treg conversion [115]. However, Oh *et al.* suggested that attenuated GvHD is caused by an impairment of effector T-cell differentiation, rather than by an increase in Tregs [102]. Although multiple mechanisms seem to contribute to GvHD amelioration, no study has showed impaired Th17 differentiation in the absence of IL-21 signaling on donor CD4⁺ T cells.

IL-23 is not involved in the initial Th17 differentiation but is necessary for the generation of a completely functional Th17 response. Expression of *IL-23* and *IL-23R* genes was upregulated during GvHD in mouse colon [117]. IL-23 production by donor-derived APCs is critical for intestinal GvHD, but not for liver or lung GvHD [117]. However, IL-23-mediated intestinal GvHD is dependent on donor-derived IFN- γ , but not on IL-17 [117], although another study showed reduced serum levels of IL-17 in the absence of donor IL-23 [118]. A subsequent study demonstrated that inhibition of IL-23 signaling reduced intestinal GvHD without losing a GVL response [119].

■ Tregs

A role of Tregs in GvHD was initially noted in experiments of costimulatory blockades [120]. Blockade of the CD40/CD40L pathway or B7 pathways ameliorated acute GvHD. This effect was not observed when Tregs were removed from the donor inoculum [120]. The suppressive effects of Tregs on GvHD are not restricted to costimulatory blockades, but rather a general phenomenon. Removal of Tregs from the donor graft dramatically accelerates GvHD [121,122]. Conversely, addition of freshly isolated donor-type Tregs significantly delays GvHD, especially at high regulatory-to-effector T-cell ratios (1:1) [121-125]. In addition, donor or host Tregs were able to ameliorate ongoing chronic GvHD [126,127]. These results revealed the purification of sufficient numbers of Tregs could be

a major limitation for applying this strategy to humans. To date, the only realistic use of Tregs in humans would be to expand them *ex vivo*.

In this context, the ability of *ex vivo*-expanded Tregs to inhibit GvHD has been tested [120,121,124,128]. Donor-derived Tregs were isolated and expanded in culture with host-derived APCs or with CD3/CD28 stimulation and high-dose IL-2 [124]. Exogenous IL-2 is critical because Tregs do not produce sufficient amounts of IL-2 for their own expansion. Injection of *ex vivo*-expanded Tregs significantly ameliorated GvHD induced by Tcons, without interfering with stem cell engraftment and functional immune reconstitution with diverse TCR V β repertoire [124,129]. Improved immune reconstitution is mediated by inhibition of GvHD-induced damage of the thymic and secondary lymphoid microenvironment [129]. Tregs can also be generated *in vitro* by stimulating FOXP3⁺ T cells in the presence of immunosuppressive agents or tolerizing DCs. These iTregs show potent suppressive activity *in vitro* and induce tolerance in transplantation [130–132] and autoimmunity [133,134] *in vivo*, although one study demonstrated that iTregs quickly reverted to FOXP3⁺ T cells and failed to prevent GvHD upon transfer [135]. STATs are critical transcriptional factors involving iTreg generation. Infusion of CD4⁺ T cells constitutively expressing active STAT5b induces attenuated GvHD by enhancing the expansion and function of Tregs [136]. STAT3 ablation enhances Treg reconstitution through thymus-dependent and thymus-independent pathways [137]. These results suggest that pharmacological or genetic modulation of the STAT protein in T cells may be a useful approach to generate efficient iTregs. *In vivo* induction of iTregs may be an alternative approach. Injection of *ex vivo*-generated regulatory DCs also protect against chronic GvHD by enhancing generation of iTregs from FOXP3⁺ donor T cells *in vivo* [138].

After infusion, Tregs colocalize with alloreactive T cells and expand in secondary lymphoid organs. Alloreactive donor T cells are activated by alloantigens expressed on host APCs in secondary lymphoid organs [2,3]. Stimulation of donor Tregs by host APCs is also necessary for the induction of GvHD protection, suggesting a double-edged role of host APCs in inducing and regulating GvHD [139]. Tregs suppress the initial activation and functional maturation of alloreactive donor T cells in secondary lymphoid organs. A recent study using an *in vitro* human skin explant GvHD model showed that donor-derived Tregs effectively suppressed

CD8-mediated skin damage only when added to the culture during initial effector T-cell priming with alloantigens [140]. Therefore, Treg therapy may be more effective when given as GvHD prophylaxis than as a treatment of established GvHD. Several costimulatory pathways such as CD28/B7, OX40/OX40L, 4-1BB/4-1BBL, CD30/CD153 and PD1/PDL1 are critical for Treg-mediated suppression [141–143]. However, these pathways are also important for Tcon activation and exhaustion of effector T cells in GvHD [11,144].

Upon stimulation, Tregs rapidly switch expression of adhesion molecules and chemokine receptors, acquire a new migratory capacity and migrate to GvHD target tissues [145]. The important T-cell homing receptor CD62L is required by Tregs to suppress Tcons at the priming sites (secondary lymphoid organs) [128,146]. On the other hand, CCR5⁺ Tregs fail to migrate to GvHD target tissues [147]. CD103⁺ Tregs that express high levels of CCR5 but low levels of CD62L directly migrate to GvHD target tissues and strongly suppress GvHD upon transfer [127].

A recent study in a mouse model suggested an association between progressive loss of Tregs during the course of acute GvHD with subsequent development of chronic GvHD by allowing expansion of pathogenic Th1 and Th17 cells [148]. On the other hand, thymic injury during acute GvHD allows the emergence of autoreactive T cells that escape from negative thymic selection and cause chronic GvHD, even in the presence of functional Tregs [149]. Whether such an imbalance between regulatory and effector T cells induced during acute GvHD could be a cause of chronic GvHD remains to be determined in humans.

Whether a graft-versus-leukemia (GVL) effect is maintained when Tregs are envisaged for the control of GvHD is a critical point to be determined. Since Tregs suppress the initial activation of alloreactive donor T cells in secondary lymphoid organs, Treg therapy probably abrogates GVL activity of donor T cells. However, Treg therapy did not result in a complete loss of GVL activity in several animal studies [124,125,150]. This is probably due to an incomplete inhibition of alloreactive T-cell activation that allows induction of at least some GVL effects mediated through the perforin–granzyme pathway. This pathway plays a major role in GVL but may play a lesser role in Treg-mediated suppression of GvHD [151]. Factors related to leukemia, such as leukemia burden, its aggressiveness or the localization of its growth, could also determine whether GVL activity would be lost in Treg therapy [124].

Recent experimental and clinical studies suggest that exposure of the fetus to noninherited maternal antigens (NIMAs) during pregnancy has an impact on allogeneic SCT performed later in life [152–154]. In mice, ‘child-to-mother’ bone marrow transplant from a NIMA-exposed donor reduced morbidity and mortality of GvHD in an antigen-specific manner [155]. Tolerogenic NIMA effects were mediated by exposure to NIMAs both *in utero* and through breastfeeding after birth, and required Tregs [156]. Substantial numbers of maternal cells cross the placenta to reside in fetal lymph nodes, inducing the development of Tregs that suppress fetal antimaternal immunity and persist at least until early adulthood in humans [157].

Tregs could suppress alloresponses through antigen-specific suppression, linked suppression or infectious tolerance. Linked suppression is the phenomenon of suppressing responses to a specific antigen by copresenting it simultaneously with another antigen, against which tolerance has previously been established. iTregs recognizing ovalbumin (OVA) prevent GvHD mediated by alloantigen-specific polyclonal effector T cells only when allogeneic recipients express OVA [132]. Tolerance can be infectious. Coactivation of Tregs with Treg cell-depleted CD4⁺ T cells results in anergized CD4⁺ T cells that in turn inhibit the activation of Tcons [158]. In the presence of Tregs that actively maintain allograft tolerance, naive T cells are newly recruited to the graft and differentiate into graft-specific Tregs [159]. Whether Tregs exert their Treg function through such a linked suppression or infectious tolerance remains to be determined in humans.

In summary of this part, apparently, any Th-cell lineage has the potential to mediate GvHD, whereas Tregs limit the disease. The fine balance between Th1, Th2, Th17 and Treg subsets may be associated with the severity, manifestation and tissue distribution of GvHD.

Th17 & Tregs in human GvHD

■ Th17

The initial phase of GvHD involves cytokine release during preconditioning of SCT recipients and cytokine gene polymorphisms are associated with GvHD development and its severity. Carvalho *et al.* analyzed the associations between variants in *IL-17A*, *IL-17F* and *IL-23R* genes and clinical outcome in T cell-depleted allogeneic SCT [160]. Donor polymorphism of *IL-17A* and *IL-23R* genes was the most important prognostic factor for survival and GvHD. This suggested

the potential usefulness of Th17 genotyping in donor selection, although validation studies are required to confirm this association. Clinical studies assessing the correlation between Th17 cells with GvHD have shown conflicting results. Th17 cells are increased in peripheral blood of patients with acute GvHD in one study [161], but not another study [162]. Ratio of Th17 to Treg was increased in the biopsied samples of GvHD tissues in one study [161], whereas it was decreased in another study [163]. Th17 cells were not increased in the skin in contrast to a significant increase in IFN- γ -producing cells at the onset of acute GvHD [162].

Ustekinumab is a clinical-grade antibody against p40, a common component of IL-12 and IL-23, and their use lead to a blockade of Th1 and Th17 differentiation. The efficacy and safety of ustekinumab has been demonstrated in Phase III trials for treatment of psoriasis [164–166]. It has also been tested in Phase II studies for multiple sclerosis, sarcoidosis and Crohn’s disease [167,168]. On the basis of results from animal studies, blockade of Th1 and Th17 are also considered promising for GvHD prevention [19,105,148]. The anti-IL-17 mAb LY2439821 has been tested for treatment of rheumatoid arthritis [169]. However, results from animal studies indicate that blockade of the IL-17 pathway alone may have limited effects on GvHD.

Modulation of the Th17/Treg balance by cytokine blockade is a realistic and attractive strategy to prevent GvHD. Tocilizumab, a humanized mAb against IL-6R, has been shown to be effective against several autoimmune and inflammatory diseases such as rheumatoid arthritis and Castleman’s disease [170–172]. Although an animal study demonstrated that IL-6 blockade attenuated acute GvHD by increasing Tregs [112], whether IL-6 blockade could shift from Th17 responses toward Tregs in humans remains to be determined. A single case report suggested the effectiveness of tocilizumab in a patient with refractory gastrointestinal GvHD [173]. Tocilizumab may also be effective for treatment of sclerodermatous skin lesion in chronic GvHD because a recent study demonstrated its efficacy for the skin lesion in patients with systemic sclerosis [174].

■ Tregs

Although the beneficial effect of Treg appears to be clear in the context of ameliorating GvHD in experimental studies, data from clinical studies show controversies in terms of Treg function in human patients. The association between the

number of Tregs infused or Treg reconstitution post-transplant and clinical outcome is not clear so far [175–177]. In earlier studies, where CD4⁺ Tregs were defined as expressing only CD25, higher numbers of CD4⁺CD25⁺ cells correlated with more acute and chronic GvHD probably due to expression of CD25 on activated effector T cells [178,179]. Daclizumab is a humanized mAb against CD25. In a randomized study, the addition of daclizumab to corticosteroids did not reduce GvHD [180]. This unexpected outcome may be observed because of the elimination or suppression of Tregs by daclizumab. The source of stem cells impacts the Treg content. G-CSF preferentially mobilizes CD62L^{lo} Tregs that have poor suppressive activity [181,182]. T-cell lineage in umbilical cord blood is biased toward immune tolerance but the function of cord blood Tregs appears to be similar to that of adult Tregs [183].

Recent studies thus analyzed FOXP3 expression and demonstrated reduced numbers of FOXP3⁺CD4⁺CD25⁺ T cells in peripheral blood of patients with acute and chronic GvHD [184–186]. Such an association was also found after T-cell-depleted SCT, but not after CD25-depleted SCT [187,188]. By contrast, an increased frequency of a skin-homing Treg population (cutaneous lymphocyte antigen [CTL*]) or a gut-homing Treg population ($\alpha 4\beta 7^+$) in peripheral blood was associated with a reduced risk of skin or gut GvHD, respectively [189]. Immunohistochemical studies of biopsy samples taken from GvHD target organs demonstrated a decreased ratio of FOXP3⁺ cells to CD8⁺ T cells in patients with intestinal GvHD [190,191], although a subsequent study failed to show this association [163]. Increase in FOXP3⁺ cells in the skin was associated with better treatment response [192]. However, in most studies, the patients were sampled at a single time point. Studies repetitively measuring Treg number during the first year after SCT clearly showed the Treg number varied widely depending on the sampling time points [186,193–195]. Moreover, as previously mentioned, the findings that FOXP3⁺ T cells in humans are functionally heterogeneous including non-Treg cells, and that a subset of FOXP3⁺ T cells has a suppressive function may be related to these disparate findings. Thus, these results need to be validated in a larger, multicenter cohort.

Tregs display greater proliferation and metabolic activity than Tcons in the steady state [196,197]. Thus, Tregs are more sensitive to irradiation and cyclophosphamide, both are used as a preparative regimen of SCT [198]. After

allogeneic SCT, the initial phase of T-cell reconstitution is primarily dependent on peripheral expansion of mature T cells present in the stem cell graft. Tregs can also be generated from a FOXP3⁺ T-cell population in lymphopenic host mice [199]. After acute lymphopenia-induced homeostatic proliferation, Treg reconstitution can occur through thymus-dependent generation from donor hematopoietic progenitor cells. Although the relative contribution of thymic and extrathymic pathways to the long-term reconstitution of Tregs remains to be determined, thymic generation of naive Tregs was markedly impaired in adult patients [195].

Treg reconstitution appears to occur at least as fast as Tcons after SCT [187,188,194,195]. During the lymphopenic period after transplantation, Tregs underwent more rapid expansion than Tcons, acquired a predominantly activated/memory phenotype and increased susceptibility to Fas-mediated apoptosis [195]. Thus, a Treg pool was preferentially declined and resulted in a prolonged imbalance between Tregs and Tcons in patients with extensive chronic GvHD. This issue, however, may be still controversial because another study demonstrated an increased ratio of Tregs to Tcons in patients with chronic GvHD following a reduced intensity conditioning and *in vivo* T-cell depletion with alemtuzumab [200]. A recent study demonstrated a positive correlation between Treg recovery and cytomegalovirus-specific CD8⁺ T cells [186]. This may be a reflection of better immune reconstitution in the absence of GvHD.

Stimulation of Treg reconstitution may be a promising approach to induced tolerance. Rapamycin promotes Treg differentiation and as a treatment accumulates Tregs in the skin [201]. Extracorporeal photopheresis is clinically used to treat GvHD. Studies in mice and humans demonstrated that the transfer of white blood cells exposed to UV radiation increased peripheral blood Tregs and serum levels of TGF- β [202–205].

Clinical application of Tregs

Accumulating evidence from experimental animal studies suggest that adoptive transfer of Tregs is a promising strategy in preventing or treating GvHD in humans. In animal studies, a ratio of nearly 1:1 of Tregs:Tcons is required for effective GvHD suppression [121,123–125]. Since alloreactive T cells presumably exist at a high frequency and obtaining a sufficient number of relatively rare Tregs from a single donor is difficult, *ex vivo* expansion of Tregs is largely necessary

to achieve large-scale deletion of alloreactive T cells *in vivo* in order to create transplantation tolerance. Achieving maximum purity of Tregs is also critically important in order to avoid the risk of GvHD exacerbation by contaminating effector T cells.

The FOXP3⁺ T-cell population in humans is heterogeneous and includes non-Treg cells [75]. FOXP3 can be weakly expressed in activating effector cells. Thus, FOXP3 is not always a reliable marker for human Tregs. Moreover, intracellular staining for FOXP3 expression is not adequate for practical use when trying to isolate viable cells. Novel surface markers specific for Tregs have not yet been identified. Currently, antibody-coated magnetic bead separation is most often utilized for Treg isolation to first deplete non-CD4⁺ T cells, followed by a positive selection to enrich for CD25⁺ cells via a saturating concentration of CD25 antibody to capture the CD25^{hi} fraction [206,207]. Alternative isolation strategies include depletion of CD127⁺ cells or selection of CD45RA⁺ or CD49d⁺ cells [208–211]. In culture, repetitive *in vitro* stimulation of Tregs results in a loss of stable FOXP3 expression [212]. CD45RA⁺FOXP3^{hi} memory Tregs are the main source of converting cells, whereas CD45RA⁺FOXP3^{lo} naive Tregs show less conversion. Loss of FOXP3 expression in memory Tregs appears to be the result of a cell intrinsic program initiated in those cells in response to strong and repetitive *in vitro* stimulation [212]. Therefore, CD4⁺CD25^{hi}CD45RA⁺ naive Tregs may be an ideal subset choice for Treg therapy. Depletion of CD127⁺ cells from the CD4⁺CD25^{hi} starting population may not improve the homogeneity of the cell product with respect to FOXP3 expression after *in vitro* expansion [212]. Hoffmann *et al.* provide instructions for isolation, polyclonal expansion, and functional characterization of human Tregs in detail [211].

High numbers of Tregs can be obtained by polyclonal or antigen-specific expansion. However, even adequately purified Tregs can give rise to a substantial fraction of T cells that produce proinflammatory cytokines in culture. Thus, maintaining Treg purity in expansion is important. Interaction between the sphingosine 1-phosphatase (S1P₁)-mTOR signaling selectively controlled the reciprocal differentiation of Th1 cells and Tregs [213]. The mTOR inhibitor rapamycin and S1P₁ antagonist FTY720 target this pathway to favor Treg differentiation. Addition of rapamycin to Treg expansion cultures blocks Treg differentiation into nonregulatory T cells [214–216]. Accordingly,

Treg proliferation levels were found in patients who received rapamycin for acute GvHD prophylaxis compared with those who received other regimens [195]. In addition, administration of FTY720 has been shown to produce GvHD suppressive activity in mice models by Treg-dependent and -independent mechanisms [217–219]. Histone/protein deacetylases (HDACs) regulate chromatin remodeling and the function of transcription factors. FOXP3 activity is also regulated by HDACs and other transcriptional coregulators [220,221]. HDAC inhibitors promote the generation and function of Tregs [222,223]. Reddy *et al.* demonstrated that HDAC inhibitor vorinostat regulate experimental acute GvHD by multiple mechanisms, although it is not clear whether vorinostat might affect Treg function [224,225]. A Phase II clinical trial is ongoing to determine if vorinostat can reduce the prevalence of acute GvHD following matched related-donor SCT after reduced intensity conditioning.

One of the major concerns of Treg therapy is whether the infused Tregs can survive *in vivo*. Since activated Tregs are prone to die by apoptosis, strategies that clonally expand antigen-specific Tregs while inhibiting excessive activation in culture are required. However, most Treg expansion processes cause Treg maturation. Another concern is whether the infused Tregs can continue to be functionally stable *in vivo*, especially under proinflammatory conditions such as those found early after SCT; the cytokine environment is important for Treg function and differentiation. FOXP3 undergoes post-translational changes as a consequence of extrinsic cellular signals. Stability of FOXP3 expression is associated with demethylation of the Treg-specific demethylated region. Epigenetic imprinting in this region is critical for the establishment of a stable Treg lineage [226]. iTregs generated *de novo* in a TGF- β -dependent process do not display FOXP3 DNA demethylation despite Foxp3 expression, whereas FOXP3 DNA is continuously demethylated in nTregs [227]. Quantitative DNA methylation analysis of Foxp3 is a novel screening method for counting stable Tregs in peripheral blood and tissues and can be used to estimate *in vivo* expansion of functional Tregs after transfer [228].

The suppressive effects of Tregs are not always antigen-specific even after antigen-specific expansion. To achieve efficient therapeutic effects without general immunosuppression, generation of antigen-specific Tregs is required. For this purpose, Tregs are generated in the presence of host

APCs in culture [124]. Such antigen-specific Tregs are much more potent than irrelevant Tregs in controlling GvHD without inhibiting immune reconstitution in mice [124]. However, whether Tregs are antigen-specific in humans where the situation is not as simple as in mice remains unclear. Even once activated by a particular antigen, Tregs can suppress effector T cells irrespective of whether they share antigen-specificity with the Tregs *in vitro* [229]. However, such a bystander suppression by Treg has not been evident *in vivo*. A recent study showed that Tregs can prevent allograft rejection without compromising immunity to viral antigens *in vivo* [230].

Defining the optimal timing of administration in order to achieve the most efficient effects of Treg therapy is also critically important. Treg therapy initiated after GvHD is established appears to have little value for therapeutic application as shown in an *in vitro* skin explant model [140]. Accordingly, Tregs should be expanded before SCT in each case, making Treg therapy cost ineffective and impractical.

An additional potential obstacle to successful use of Tregs is an adverse impact of immunosuppressive agents on Tregs. Calcineurin inhibitors such as cyclosporine and tacrolimus inhibit IL-2 transcription and thereby have an inhibitory effect on Tregs that are dependent on IL-2 [231]. By contrast, other agents such as rapamycin and mycophenolate mofetil, which act through alternative mechanisms, may be permissive for Treg expansion and function [231]. Thus, selection of immunosuppressive drugs that have less suppression on Tregs than on Tcons may be crucial for successful Treg therapy. The net effect of GvHD prophylaxis with a combination of calcineurin inhibitor and rapamycin or mycophenolate mofetil on Treg reconstitution, however, remains unclear. Alternative approaches include use of calcineurin inhibitor-free GvHD prophylaxis as a platform of Treg therapy or infusion of IL-2 to support *in vivo* expansion of Tregs. The latter strategy was tested in a recent clinical trial; administration of low-dose IL-2 after CD4⁺ donor lymphocyte infusion (DLI) enhanced Treg expansion [232].

Several clinical trials are currently ongoing to examine the adoptive transfer of Tregs as prevention against GvHD in humans. The first-in-man clinical results of the treatment of GvHD with *ex vivo*-expanded Tregs was reported from Poland [210]. CD4⁺CD25^{hi}CD127⁺ cells were sorted from the SCT donor buffy coat, polyclonally expanded with anti-CD3/anti-CD28 beads with IL-2, and infused into two patients suffering from GvHD.

The therapy was effective in a patient with chronic GvHD, but not in one with acute GvHD. Umbilical cord blood can be a readily accessible source of Tregs. Results of a prospective and multicenter clinical trial to evaluate the safety of cord blood-derived Tregs were also reported [233]. CD25^{hi} cells were isolated using anti-CD25 magnetic beads and the CliniMACS[®] device from cryopreserved cord blood and expanded similarly. Expansion efficiency was much better in cord blood Tregs than adult Tregs. Upon infusion, no infusional toxicities were observed in 23 patients who underwent double unit cord blood transplantation. Apparently no increase of acute GvHD, infection and disease relapse occurred. The group from Perugia in Italy conducted adoptive immunotherapy with Tregs after HLA-haploidentical PBSCT [234]. Freshly isolated CD4⁺CD25^{hi} cells with CliniMACS were infused 4 days before infusion of Tcons together with purified CD34⁺ PBSCs. Treg therapy made administration of Tcons feasible in a haploidentical setting, with a low incidence of GvHD and cytomegalovirus reactivation. Conversely, the safety and efficacy of Treg-depleted DLI was studied in 17 adult patients with malignancy relapse after SCT [235]. This strategy was safe and induced GvHD and GVL in patients not responding to a classical DLI.

Future perspective

The balance between regulatory and effector T cells may be an important determinant of GvHD induction or tolerance induction and the fine balance between T helper subsets *in vivo* may impact on clinical manifestation and tissue distribution of GvHD. This assumption suggests that effective strategies to suppress GvHD include selective inhibition of effector T cells while maintaining or enhancing Treg function and reconstitution, as well as manipulation of the balance between T helper subsets to treat organ-specific GvHD. In this context, adoptive Treg therapy has been initiated in SCT patients. These initial studies will provide data regarding safety and feasibility of Treg therapy. We will be paying particular attention to whether Treg therapy could increase the risk for opportunistic infection by their nonspecific immunosuppressive effects, whether it could promote leukemic relapse, and whether it would rather induce GvHD by the contaminated effector cells or by the *in vivo* conversion of Tregs to inflammatory T cells. In addition, development of novel methods to monitor *in vivo* survival and expansion of functional Tregs are required

to estimate impacts of Treg therapy on its outcome. Such a new technology also helps to determine the optimal timing and cell dose of Treg therapy. Selection of drugs for GvHD prophylaxis that do not impair Treg function is also crucial for successful implantation of Treg immunotherapy. Currently performed custom-made Treg preparations for each patient are not realistic for the widespread use of Treg immunotherapy. Improvement of Treg therapy thus requires much better understanding of the checks and balances the immune system to maintain tolerance and induce immunity by overcoming many problems that we now face. Certainly, Treg therapy has a great promise of

avoiding general immunosuppression and many toxicities by currently using potent immunosuppressive agents. In order to modulate the balance of Th subsets *in vivo*, the use of biological products such as cytokine-neutralizing mAbs that has increasingly been used clinically in treating patients with cancer and autoimmune diseases, is a realistic and attractive strategy because cytokine environment is critically important for T helper and Treg differentiation. To establish highly effective Treg therapy or cytokine modulation, consideration of timing of administration is particularly important in the setting of highly inflammatory allogeneic SCT. However, as reduced intensity conditioning has been

Executive summary

The Th1/Th2 paradigm in graft-versus-host disease

- * Differential activation of Th1 or Th2 cells has been suggested to play an important role in GvHD development.
- * The long-held assumption that acute GvHD is a Th1-mediated disease, whereas chronic GvHD is a Th2-mediated disease, needs to be refined.

Th17

- * Th17 cells are characterized by their expression of proinflammatory cytokines IL-17, IL-21 and IL-22.
- * Th17 cells differentiate from naive CD4⁺ T cells by stimulation with antigens in the presence of TGF- β and IL-6 or IL-21.
- * Th17 cells contribute to host defense against certain pathogens by inducing expression of antimicrobial factors and recruiting neutrophils.
- * Th17 cells play a role in maintaining intestinal ecology with commensal bacterial flora.
- * Th17 immunity plays a central role in many autoimmune diseases.

Tregs

- * Tregs are indispensable to maintain self-tolerance and immune homeostasis, establish tolerance to allografts and promote fetomaternal tolerance.
- * Naturally occurring Tregs develop in the thymus while induced Tregs are derived from CD4⁺FOXP3⁺ T-cell population in the periphery.
- * Tregs suppress immune responses through contact-dependent and cytokine-dependent mechanisms.
- * FOXP3⁺ T cells are heterogenous and include nonregulatory T cells.

Plasticity of effector & Treg cells

- * There is a plasticity and flexibility of T-cell commitment to regulatory and effector T-cell subsets.
- * Th17 cells have the potential to convert toward a Th1 profile.
- * Th17 and Treg development programs are reciprocally interconnected.
- * Stable expression of FOXP3 is required for the maintenance of the developmentally established Treg function program.

Th17 & Tregs in experimental GvHD

- * The role of IL-17 in GvHD is still controversial.
- * Th17 cells may be sufficient but not necessary for GvHD induction.
- * Th17 may be involved in skin GvHD.
- * Treg therapy suppresses GvHD only at high regulatory-to-effector T-cell ratios.
- * Therapeutic approach to treat GvHD probably rely on nTregs, not iTregs.
- * Tregs primarily suppress the initial activation of alloreactive donor T cells at priming sites.

Th17 & Tregs in human GvHD

- * Clinical studies assessing the correlation between GvHD with Th17 or Tregs have shown conflicting results.
- * Thymic generation of naive Tregs is markedly impaired in adults.
- * Tregs undergo rapid expansion, acquire an activated/memory phenotype, and increase susceptibility to apoptosis during the lymphopenic period after transplantation.

Clinical application of Tregs

- * *Ex vivo* expansion of Tregs is required to achieve large-scale deletion of alloreactive T cells *in vivo* in order to create transplantation tolerance.
- * Achievement of a maximum Treg purity is required in order to avoid the risk of GvHD exacerbation by contaminating effector T cells.
- * Survival of functionally stable Tregs *in vivo* after transfer remains a major concern of Treg therapy.
- * Suppression of immunity against pathogens and leukemia remains to be estimated.
- * The optimal timing for Treg therapy is not clear.

developed to minimize toxicity of conditioning regimen, advances in transplant medicine will synergistically facilitate development of such novel strategies.

Conversely, results from animal studies suggest that a combined blockade of Th1 and Th17 differentiation pathways of donor T cells may represent a promising strategy for the prevention or treatment of GvHD, while inhibition of either pathway seems to be insufficient to prevent GvHD [19,105,148]. In this context, novel strategies targeting Th1 and Th17 promoting cytokines, miRNAs, transcriptional factors such

as T-bet and ROR- γ t through administration of mAbs, small molecule inhibitors, or siRNA may be promising for the control of GvHD [236,237].

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Postprogression survival for first-line chemotherapy of patients with advanced non-small-cell lung cancer

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Background: Given the growing number of drugs available for non-small-cell lung cancer (NSCLC), an effect of first-line chemotherapy on overall survival (OS) might be confounded by subsequent therapies. We examined the relation between postprogression survival (PPS) and OS in phase III trials of first-line chemotherapy for advanced NSCLC.

Patients and methods: A literature search identified 69 trials that were published during the past decade. We partitioned OS into progression-free survival (PFS) and PPS and evaluated the relation between OS and either PFS or PPS. We also examined whether any association might be affected by the year of completion of trial enrollment.

Results: The average PPS was longer in recent trials than in older trials (6.5 versus 4.4 months, $P < 0.0001$). For all trials, PPS was strongly associated with OS ($r = 0.82$), whereas PFS was moderately associated with OS ($r = 0.43$). The correlation between OS and PPS in recent trials was stronger than that in older trials ($r = 0.89$ and 0.66).

Conclusions: Our findings indicate that, especially for recent trials, PPS is highly associated with OS in first-line chemotherapy for advanced NSCLC, whereas PFS is only moderately associated with OS.

Key words: chemotherapy, non-small-cell lung cancer, overall survival, phase III trial, progression-free survival

Introduction

Lung cancer remains the leading cause of cancer death worldwide [1, 2], with non-small-cell lung cancer (NSCLC) accounting for ~85% of lung cancer cases. Most individuals with NSCLC have metastatic disease at the time of diagnosis and therefore have a poor prognosis. The standard treatment of advanced NSCLC over the past decade has been platinum-based chemotherapy because of the moderate improvement in survival it confers [3–6]. Although many patients initially achieve clinical remission or disease stabilization with first-line chemotherapy, nearly all subsequently experience disease progression and eventually die of advanced NSCLC.

Overall survival (OS) has been traditionally recognized as the most important therapeutic objective for NSCLC patients. However, in view of the growing number of drugs and combinations thereof that are available for the treatment of such patients, any effect of first-line chemotherapy on OS might be confounded by subsequent therapies [7]. Indeed, an improvement in progression-free survival (PFS) has not necessarily resulted in an improved OS in recent randomized trials in patients with NSCLC [8, 9].

The effect of therapies instituted after disease progression on survival in clinical trials is thus of interest. However, little is known about postprogression survival (PPS) in NSCLC. In the

present study, we partitioned OS of phase III trials for chemotherapy-naïve patients with NSCLC into PFS and PPS and assessed the association of each with OS.

Methods

Search strategy and selection of trials

An independent review of PubMed citations from 1 January 2000 to 31 October 2010 was carried out. Key words included in the search were 'non-small cell lung cancer', 'clinical trial', 'advanced', and 'chemotherapy'. The search was limited to randomized controlled phase III trials and articles published in English. We reviewed each publication, and phase III studies that compared two or more first-line systemic chemotherapies (including treatment with molecularly targeted agents) for advanced or metastatic NSCLC were selected. To find any additional trials, we searched the reference lists of included trials as well as of large systematic reviews. We also checked articles that were in press at leading journals and searched websites listing abstracts from conferences (organized by the American Society of Clinical Oncology or the Federation of European Cancer Societies). We included trials that provided data for both OS and either PFS or time to progression (TTP), whether or not these parameters were explicitly defined. Trials were excluded if they investigated only immunotherapy regimens or hormonal therapies. Trials that were designed to assess combined modality treatments, including radiation therapy and surgery, were also excluded. To avoid bias, two observers (HH and IO) independently abstracted the data from the trials.

Data abstraction

We analyzed in detail the primary and secondary efficacy end points, following the definitions of the authors of each trial. When not specifically

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stated by the authors, we considered the primary end point to be that used for calculation of sample size. For the sake of simplicity, two end points (PFS and TTP) based on tumor assessment are collectively referred to as PFS in the present study, similar to the approach adopted in a recent report [10]. Median OS and median PFS were extracted from all trials that provided data for each treatment group. Median PPS was defined as median OS minus median PFS for each trial. We also obtained the following information from each report: year of completion of trial enrollment, number of patients randomized, number of patients in each treatment arm, number of treatment arms in each trial, proportion of patients who were male or had adenocarcinoma, and median age of the patients.

data analysis

We summarized the survival data (median OS, median PFS, median PPS, and median PFS/median OS) as the average and standard error (SE) for trial arms. SE was calculated on the basis of previously described models [11]. We also calculated the percentage of OS accounted for by PPS for each trial arm as: $100 - (100 \times \text{median PFS}/\text{median OS})$. To assess the relation between median OS and either median PFS or median PPS, we used Spearman's rank correlation coefficient. To account for differences in sample size among trial arms, we weighted all analyses by the number of patients in each arm. In addition, all trials were divided into two groups on the basis of the year in which trial enrollment was completed. Given that the median year for completion of enrollment in the 69 analyzed trials was 2002, we dichotomized at year 2002 (older trials, up to and including 2002; recent trials, 2003 and later) in order to evaluate a possible change in PPS, and we assessed whether the evaluated relations might be dependent on the year of completion of trial enrollment. We examined differences in the survival data between older and recent trials by normal approximation of the average survival data (z test). All reported P -values correspond to two-sided tests, and those of P -values <0.05 were considered statistically significant. Analyses were carried out with SAS for Windows release 9.2 (SAS Institute, Cary, NC).

results

characteristics of the trials

Our search yielded a total of 467 potentially relevant publications. Initially, 366 studies were excluded for at least one of the following reasons: they examined other malignancies or combined modality treatments, they were not randomized, they were phase I or II trials, they were review articles, they represented subgroup analyses, or they were duplicates. The selection process for the randomized controlled trials is shown in Figure 1. Review of the remaining 101 publications yielded 69 trials that were considered to be highly relevant for the present study. The main characteristics of the 69 phase III trials included in the analysis are listed in Table 1. A total of 37 986 patients with advanced NSCLC were enrolled, with a median number of patients per study of 433 (range 153–1725). Most of the trials had a high proportion of male patients and of patients with adenocarcinoma. The average median age of the patients was 62.3 years. Ten trials used an end point based on tumor assessment (PFS or TTP) as the primary end point, whereas OS was assessed as the primary end point in 53 trials. The other six trials used response rate or quality of life as the primary end point.

median OS, PFS, and PPS in all trials and in subgroups based on year of completion of trial enrollment

The survival data for trial arms according to the year in which trial enrollment was completed are shown in Table 2. Although the average median PFS in older (up to and including 2002) trials was the same (4.9 months) as that in recent (2003 and later) trials, the average median PPS was ~50% longer in the

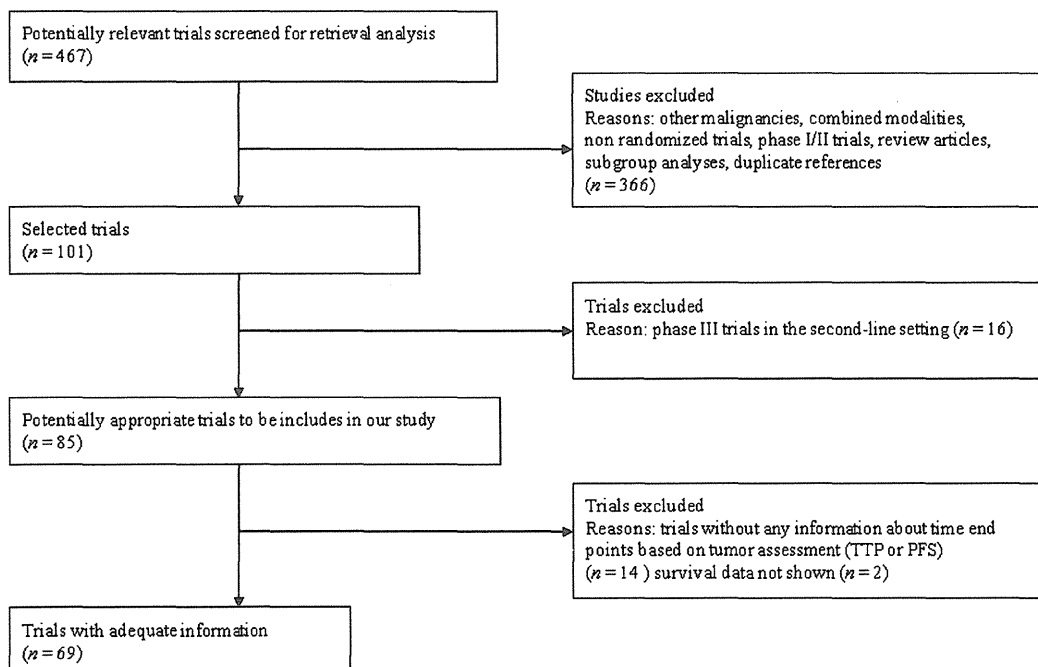


Figure 1. Flow chart showing the progress of trials through the selection process.