Novel TGF-β inhibitors ready for prime time in onco-immunology

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ABSTRACT
Transforming Growth Factor (TGF-β) inhibitors have been in development for decades with the outmost results of being promising candidates. From the latest clinical results at the 2016 ASCO meeting converging evidences suggest that we have moved from promising to effective drug nomines.

TGF-β pathway stands as one of the most universal and conserved pathways in the animal kingdom and one of the most potent immunomodulators albeit remaining a challenge to tackle for oncolgical applications. Considering the whole TGF-β family, more than 30 family members have been shown involved in various physiological and pathological processes. In oncology, TGF-β has been involved in cell proliferation, angiogenesis, epithelial-to-mesenchymal transition, immune infiltration, metastases dissemination, and drug resistance. Regarding immunomodulation, TGF-β signaling has been shown to switch Th1/Th2 (type 1 and type 2 helper T cells) balance toward Th2 via IL-10 and direct inhibition of the Th1 response, to inhibit M1-type while promoting M2-type macrophages, and to suppress CD8+ T, NK, and dendritic cell functions while increasing CD4+CD25+ T-reg cell functions. Crosstalk between the canonical TGF-β signaling (SMADs) and several non-canonical pathways including MAPK, PI3K, WNT, HH, and NOTCH has been described. Furthermore, the seemingly paradox that most cancer cells display altered or non-functional TGF-β signaling further added complexity for recognizing the value of TGF-β and TGF-β receptors as targets for drug development in oncology. Nonetheless, continuous endeavor to understand the role of TGF-β has started highlighting the crucial importance of the tumor microenvironment, unraveling how TGF-β was every so often behaving as an oncogene at late stage or as a tumor suppressor gene at early stage of tumor development. In advanced tumors, TGF-β inhibition may result in anti-proliferative effects at low concentrations and pro-proliferative effects at higher doses, making uneasy the selection of appropriate concentration in clinical trials. Furthermore, TGF-β may have limited direct effects on tumor cells, and thereby, TGF-β inhibitors may be expected to exert only limited effects on cancer cell proliferation. Interestingly, TGF-β inhibitors primarily exert their antitumor activity by affecting TGF-β responsive cells that are fibroblastic and endothelial cells as well as T-cells in the tumor microenvironment. As a result of modulating tumor microenvironment with limited direct effects on cancer cells, TGF-β inhibitors are expected to yield cytostatic effects and lead to tumor stasis translating in few tumor responses but delayed tumor progression in clinical trials. The immune-modulatory cytokine effects of TGF-β inhibition has also recently gained much interest in the bursting onco-immunology field, which is now rushing to evaluate TGF-β pathway inhibition together with checkpoint inhibitors.

Development of small molecules TGF-β pathway inhibitors has long been impeded by on-target toxicities, especially cardiac injuries. However, significant work on drug pharmacokinetic and pathway inhibition threshold using pharmacodynamic biomarkers can considerably reduce their toxicities. Other strategies using RNAi or monoclonal antibodies are usually displaying manageable toxicities.

Small molecules inhibiting TGF-β
Several TGF-β pathway inhibitors have now reached the clinic (Fig. 1, Table 1), and some evidences, especially using small molecule inhibitors have started to emerge. Small-molecule inhibitors are primarily aimed at inhibiting TGF-β receptors and are based on a dihydropyrrolopyrazole scaffold (LY550410 and LY580276 from Eli Lilly Research Laboratories) or on...
imidazole scaffolds (SB-505124 from GlaxoSmithKline). Other inhibitors of interest are based on pyrazolopyridine, pyrazole, imidazopyridine, triazole, pyridopyrimidine, and isothiazole scaffolds. Among small molecule inhibitors galunisertib (LY2157299 Monohydrate), a drug developed by Eli Lilly, is one of the most advanced and have shown interesting results in two phase II trials. An interesting figure is that galunisertib displays a very safe toxicity profile across various clinical trials.

Table 1. Summary of the main clinical interventions and results.

| Clinical development | Main results |
|----------------------|--------------|
| Small molecule TGF-β inhibitors | |
| Galunisertib (LY2157299) | Trend to OS benefit |
| Phil (NCT01373164): galunisertib + gemcitabine vs. gemcitabine in PAC | OS benefit in patients with >20% decrease in TGF-β1, AFP, and CDH1 levels from baseline |
| Phil (NCT01246986): monotherapy in HCC | Ongoing |
| Phil (NCT01246986): combination with sorafenib in HCC | Ongoing |
| Philb (NCT02734160): combination with durvalumab in PAC | Ongoing |
| Philb (NCT02423343): combination with nivolumab in glioblastoma, NSCLC and HCC | Ongoing |
| Monoclonal antibodies against TGF-β | |
| Fresolimumab (GC1008) | Preliminary evidence of antitumor activity |
| Philb (NCT00356460): monotherapy in melanoma and RCC | Ongoing |
| Phil (NCT01401062): combination with radiotherapy in mBC | Ongoing |
| Philb (NCT02581787): combination with radiotherapy in early stage NSCLC | Ongoing |
| Vaccinations strategies | |
| Lucanix (Belagenpumatucel-L) | OS benefit in patients with prior radiotherapy and/or randomized within 12 weeks of chemotherapy completion |
| Phil (NCT00676507): monotherapy as maintenance therapy in NSCLC | |
| Vigil (Gemogenovatucel-T) | Preliminary evidence of improved relapse-free survival |
| Philb (NCT02346747): monotherapy as maintenance in high-risk ovarian cancer | Ongoing |
| Phil (NCT02511132): monotherapy vs. gemcitabine + docetaxel in Ewing’s sarcoma | Ongoing |
| Phil (NCT02639234): combination with nivolumab in NSCLC after platinum-based therapy | Ongoing |
| Antisense oligonucleotides | |
| Trabedersen | Non significant OS benefit |
| Philb (NCT00761280): monotherapy vs. SOC in refractory GBM or anaplastic astrocytoma | Preliminary evidence of improved overall survival |
| Philb (NCT01016406): monotherapy in melanoma | |

GBM: glioblastoma multiforme; HCC: hepatocellular carcinoma; mBC: metastatic breast cancer; NSCLC: non-small cell lung cancer; PAC: pancreatic adenocarcinoma; Phil: phase II trial; RCC: renal cell carcinoma.
with no dose-limiting event, including no cardiac toxicity in human that was of primary concern with first-generation TGF-\( \beta \) inhibitors tested in the clinic.

From the 2016 American Society of Clinical Oncology (ASCO), data were presented comparing galunisertib in combination with gemcitabine (GG) versus gemcitabine plus placebo (GP) in a 2:1 comparative phase II trial that randomized 156 patients with pancreatic cancer,\(^4\) a disease in which SMAD4 abnormalities are reported in near 60\% of cases. Using advanced statistical analyses, the study was considered positive on its primary endpoint with a median overall survival of 8.9 mo versus 7.1 mo in the galunisertib plus gemcitabine and gemcitabine plus placebo arms, respectively (Hazard ratio: 0.8 (95\% Confidence Interval 0.6–1.09), the probability for the hazard ratio to be <1 being 92\%). Secondary endpoints consistently showed an increase in the disease control rate of 49\% in patients receiving galunisertib versus 20\% receiving placebo. Interestingly, analyses of plasma biomarkers showed that patients with low baseline level of TGF-\( \beta \) showed an increase in the disease control rate of 49\% in patients receiving galunisertib versus 20\% receiving placebo.

Another large phase II trial performed in hepatocellular carcinoma (HCC), involving several cohorts, was also presented at ASCO. The initial cohort A included 109 patients with elevated AFP \( \geq \) 1.5 UNL treated with galunisertib as single agent in second-line after sorafenib failure, yielding a median overall survival of 8.3 mo.\(^5\) Interestingly, patients under exposure to galunisertib who decreased the expression levels of any or all the prespecified blood biomarkers, AFP, TGF-\( \beta \)-1, and CDH1 from baseline, had improved outcomes. For instance, patients with high baseline AFP > 200 ng/mL and experiencing >20\% reduction in AFP level at anytime during treatment with galunisertib presented an outstanding median OS of 21.4 mo, suggesting that a subgroup of this poor prognostic population derived a pronounced effect of galunisertib. Cohort B, including 40 patients with normal AFP and treated with galunisertib monotherapy in second line, reported a median OS of 16.8 mo; patients showing >20\% decrease in blood TGF-\( \beta \) level (~30\%) reaching a median OS of 21.8 mo.\(^5\) Expectedly, galunisertib has now moved to first line setting in a randomized phase II trial evaluating the benefit of adding galunisertib to sorafenib (NCT02178358). As well expected, and justified by recent clinical data,\(^6\) a phase I/II trial has been initiated, evaluating addition of the anti-PD-1 nivolumab to galunisertib in advanced recurrent or refractory HCC, NSCLC, pancreatic adenocarcinoma, and glioblastoma patients (NCT02423343).

### Monoclonal antibodies inhibiting the TGF-\( \beta \) pathway

Other major companies may also be interested for clinical development of monoclonal antibodies in oncology. For instance, fresolimumab (GC1008), a human monoclonal antibody initially co-developed by Cambridge Antibody Technology and Genzyme then acquired by Sanofi-Aventis for the treatment of idiopathic pulmonary fibrosis and focal segmental glomerulosclerosis, may be further investigated in renal cell carcinoma and melanoma. Novartis and Xoma also joined for co-development of XOMA089, a fully human, high-affinity, monoclonal antibody designed to neutralize TGF-\( \beta \)-1 and TGF-\( \beta \)-2 while sparing TGF-\( \beta \)-3. Preclinical data have shown that XOMA089 displays activity against tumor growth in preclinical models of squamous cell head and neck carcinoma and breast cancer (BC). Animal data also suggested that XOMA089 might be synergistic with PD-1 inhibition.

### Vaccinations strategies

Belagenpumatucel-L is the most advanced vaccine that uses four TGF-\( \beta \)-2-antisense gene-modified, irradiated, allogeneic non-small cell lung cancer (NSCLC) cell lines to stimulate immune reactions. After promising phase II data, the double blind, randomized phase III trial in patients with stage III/IV NSCLC after frontline platinum-based induction chemotherapy with or without irradiation, did not meet its pre-specified primary efficacy endpoint and was terminated at the second interim analysis for futility.\(^7\) Median OS in the intention-to-treat (ITT) population was 20.3 mo for belagenpumatucel-L compared to 17.8 mo for placebo (hazard ratio (HR) = 0.94, \( p = 0.594 \)). However, in pre-specified analyses, patients who were randomized within 12 weeks after induction chemotherapy, who received prior radiation therapy, or both, seemed to benefit most of the treatment; in these three patient groups, median OS in the vaccine arm were 20.7, 24.8, and 24.8 months, respectively, compared to 13.4, 16.0, and 10.3 mo for the placebo arm (HR = 0.77, \( p = 0.092 \); HR = 0.61, \( p = 0.032 \); HR = 0.47, \( p = 0.013 \), respectively). Owing to the fact that both chemotherapy and radiotherapy can have beneficial effects on antitumor immunity (decreased Treg cell population, improvement of the Treg/CD4\(^+\) and CD8\(^+\) T cells ratios, upregulation of MHC class I), the authors suggested that delaying vaccine treatment allow time for the immune system to recover and for the tumor to rebuild an immunosuppressive microenviroment. Following this hypothesis, a randomized phase IIb has been initiated to evaluate the combination of gemogenovacucel-T (see below) with nivolumab in advanced or metastatic NSCLC after one prior platinum-based systemic therapy (NCT02639234).

Based on the same vaccination principle, the personalized approach of the gemogenovacucel-T (Vigil, formerly FANG) vaccine consists in \textit{ex vivo} transfection of autologous tumor cells with the GM-CSF transgene together with a bi-functional short hairpin RNAi (bi-shRNAi) targeting the furin convertase and for the tumor to rebuild an immunosuppressive microenviroment. Following this hypothesis, a randomized phase I/II trial of 12 patients with recurrent advanced/metastatic Ewing’s sarcoma, expression of TGF-\( \beta \)-1 and TGF-\( \beta \)-2 proteins maturation and function. In a phase I trial of 12 patients with recurrent advanced/metastatic Ewing’s sarcoma, expression of TGF-\( \beta \)-1 and TGF-\( \beta \)-2 was dramatically downregulated and treatment was associated with 100\% conversion of immune response to unmodified autologous tumor by IFNy ELISPOT assay.\(^7\) Given a 1-y survival rate of 75\% in this small patient population, a randomized phase Ib comparing gemogenovacucel-T against a combination of gemcitabine and docetaxel was initiated in advanced refractory Ewing’s sarcoma (NCT02511132). Gemogenovacucel-T is also being pursued in a phase II/III trial as maintenance treatment in high-risk stage III/IV ovarian cancer following clinical complete response after primary surgery and adjuvant chemotherapy.
Antisense oligonucleotides to suppress TGF-β expression

Trabedersen, a synthetic TGF-β2 antisens oligodeoxynucleotide, was compared to standard chemotherapy (temozolomide or procarbazine/lomustine/vincristine) in a phase IIb trial that recruited 145 patients with recurrent or refractory glioblastoma multiforme or anaplastic astrocytoma. The trial did not meet its primary efficacy endpoint (tumor growth control at 6 mo) but delayed responses were observed even after treatment discontinuation. Median OS for 10 μM trabedersen was 39.1 mo compared to 21.7 mo for chemotherapy although the difference was not statistically significant. In a phase I/II trial in stage III/IV melanoma patients no longer amenable to established therapy, trabedersen (OT-101) was tested as decloaking agent facilitating immune response. Authors claimed outstanding OS not related to PFS associated with therapeutic benefit primarily in patients who were subsequently treated by chemotherapies.

Biomarkers

Owing to the complex nature of the TGF-β pathway, its role in cell fate and differential activity in tumor cells and their microenvironment, predictive biomarkers may be challenging to identify. Moreover, biomarkers that would tailor TGF-β inhibitors to specific patient population as single agent may not be relevant in combination therapies. Nonetheless, TGF-β signatures have been highlighted in several tumor types such as colon, breast, liver, or pancreatic cancer and may be worth investigating in clinical trials. Given its role in immunomodulation, predictive expression, or mutation in TGF-β receptors (TGFBR1/2) may also be investigated when TGF-β inhibitors are combined with immunotherapy.

In contrast to the still prospective nature of predictive biomarkers, utility of few pharmacodynamic biomarkers to monitor drug effectiveness have been highlighted. Modulation of SMAD phosphorylation is the most often used biomarker of TGF-β pathway activity in preclinical investigations. In the phase I of Galunisertib, inhibition of pSMAD2 in PBMC has been observed in 64% of the patients. However, the inherent difficulty of evaluating phosphorylated biomarkers will likely limit their use in the clinic. In contrast, circulating blood biomarkers are easier to assess and decrease in >20% of TGF-β1, AFP, or CDH1 levels from baseline has been shown to be associated with strong improvement of disease control in the phase II trial of galunisertib in HCC. However, the delay between treatment initiation and decrease in biomarker level seemed highly variable and remain unpredictable. Several other biomarkers have been tested in clinical trials, but they either lacked of significant association with disease control or were tested on a very limited number of patients to be reliably robust, warranting further investigations.

Disclosure of potential conflicts of interest

SF and ER have received consultancy compensation from Eli Lilly.

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