



# Cytokine engineering for targeted cancer immunotherapy

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## Abstract

Cytokines are key modulators of the immune responses and represent promising therapeutics for a variety of cancers. However, successful translation of cytokine-based therapy to the clinic is limited by, among others, severe toxicities and lack of efficacy due to cytokine pleiotropy and off-target activation of cells. Engineering cytokines with enhanced therapeutic properties has emerged as a promising strategy to overcome these challenges. Advances in protein engineering and protein-polymer conjugate technologies have fostered the generation of cytokines with enhanced target cell specificity and longer half-life than the native ones. These novel cytokines exhibit reduced systemic toxicities while focusing the activities at the tumor site, thus, enhancing antitumor immunity. The growing toolbox of cytokine engineering strategies will further stimulate the development of smart cytokine-based immunotherapies with enhanced efficacy and safety profiles.

## Addresses

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## Introduction

Cytokines are small proteins (typically < 30 kDa) that are produced by immune cells and play an essential role in maintaining physiological immune homeostasis and regulating pathophysiological processes, such as cancer and autoimmune diseases [1]. Cytokines are divided in different subclasses based on their receptors' conformation, including chemokines, interferons (IFNs), interleukins (ILs), and tumor necrosis factors (TNFs) (reviewed elsewhere [2–5]). They act by binding to specific receptors inducing receptor dimerization or

membrane reorganization. Receptor aggregation activates distinct kinases, which phosphorylate serine and tyrosine residues of the receptor's cytoplasmic domains and activate distinct transcription factors for nuclear translocation and gene expression regulation. For example, many IFN and IL receptors selectively associate with Janus kinases (JAKs), which initiate receptor phosphorylation and subsequent recruitment of one or more Signal Transducer and Activator of Transcription (STAT) proteins. In contrast, TNF binding to its cognate receptor induces the recruitment of specific complexes and kinases, which lead to activation of downstream transcription factors including AP-1 and the nuclear factor  $\kappa$ B (NF- $\kappa$ B) [5–9].

Given their essential role in modulating the immune responses, many cytokines have been identified as therapeutic candidates for the treatment of a variety of cancers [10]. To date, several cytokines have been approved for cancer immunotherapy in the clinic, including recombinant IL-2, IFN $\alpha$ , and TNF [11]. However, the clinical application of cytokine-based immunotherapies is greatly hampered by toxicities and/or modest efficacies. This effect is mainly due to the pleiotropic nature of cytokines causing undesired binding and activation of off-target cells. Moreover, native cytokines typically have a short serum half-life and narrow therapeutic window, which limits their therapeutic efficacy [12].

Engineering of cytokines with enhanced specificity, localized activity, and longer half-life is a promising approach to unleash their full therapeutic potential. Advances in cytokine structural biology and synthetic biology have boosted the development of engineered cytokines either through structure-based protein modifications or through the fusion of cytokines with antibody-derived fragments (Table 1) [12–14]. On the other hand, a variety of polymer–cytokine conjugates and peptide–cytokine fusions have been developed to provide therapeutic cytokines with prolonged half-life, increased specificity, and low immunogenicity (Table 1) [15–18]. Moreover, beyond directly engineering the cytokine molecule, antibodies can be used to target either the cytokine or the cytokine's receptor, thus altering their functional behavior [19,20]. In this review, we highlight recent progress in engineering cytokine molecules for targeted cancer immunotherapy

Table 1

## Summary of recent advances in cytokine engineering strategies.

Strategy	Cytokines	Key results	Representative references
<b>1. Molecular engineering</b>			
Immunocytokines	IL-2, IL-12, IFN $\gamma$ , TNF $\alpha$	↑ Tumor localization, ↑ Half-life	[24]
Site-directed mutagenesis	IL-2	↑ Selective activation of cytotoxic immune cells	[33,34,79]
Combinatorial mutagenesis	IL-2, IL-10, IL-18	↑ Selective activation of cytotoxic immune cells, ↑ Bioactivity	[37,38,40,42,43]
<i>De novo</i> protein design	IL-2, IL-15	↑ Selective activation of cytotoxic immune cells	[41]
<b>2. Chemical engineering</b>			
PEGylation	IFN $\alpha$ , TNF	↑ Half-life, ↑ Aqueous solubility	[15]
Conjugation to synthetic polypeptides	IFN $\alpha$	↑ Half-life, ↑ Stability, ↑ Tissue permeability, ↓ Immunogenicity	[51,53,54]
<sup>a</sup> ECM-targeting cytokines	IL-2, IL12, CCL4	↑ Tumor localization, ↓ Systemic toxicities	[62–65,68]

<sup>a</sup> ECM, extracellular matrix; TNF, tumor necrosis factor; IFN, interferon; PEG, polyethylene glycol.

using molecular and structural variations as well as chemical modifications (Figure 1). We also share our vision on the current limitations of cytokine engineering and technological advances needed for the successful translation of these strategies into the clinic.

### Molecular engineering of therapeutic cytokines

Molecular engineering strategies aim at enhancing the therapeutic properties of cytokines by fusing them with other protein domains, such as antibody fragments, or by altering the protein's molecular structure using codon mutagenesis. These techniques have proven successful in overcoming some shortcomings of native cytokines, including dose-limiting toxicities, off-target effects, short half-life, and insufficient activity. In this subsection, we first review recent efforts to increase cytokine tissue specificity by using cytokine–antibody fragment fusions. Next, we illustrate current advances in the modulation of cytokine activities employing mutagenesis-based approaches. For the purpose of this review, modification with peptides, although very similar to proteins in composition, is discussed in the chemical engineering subsection due to their small size and similarity to synthetic polymers.

### Immunocytokines

Targeted delivery of cytokines to the tumor microenvironment (TME) can trigger strong antitumor immunity and turn so-called immunologically “cold” tumors into “hot” tumors with high immune cell infiltration [11,21]. However, systemic administration of cytokine-based therapies is limited by their pleiotropic nature, their short half-life, and substantial side effects [15,16]. Antibody–cytokine fusion proteins, called immunocytokines, consist of the fusion of cytokines to full-sized antibodies or antibody fragments, which provide the molecule with the capability to target tumor-associated antigens. This led to current clinical testing of multiple

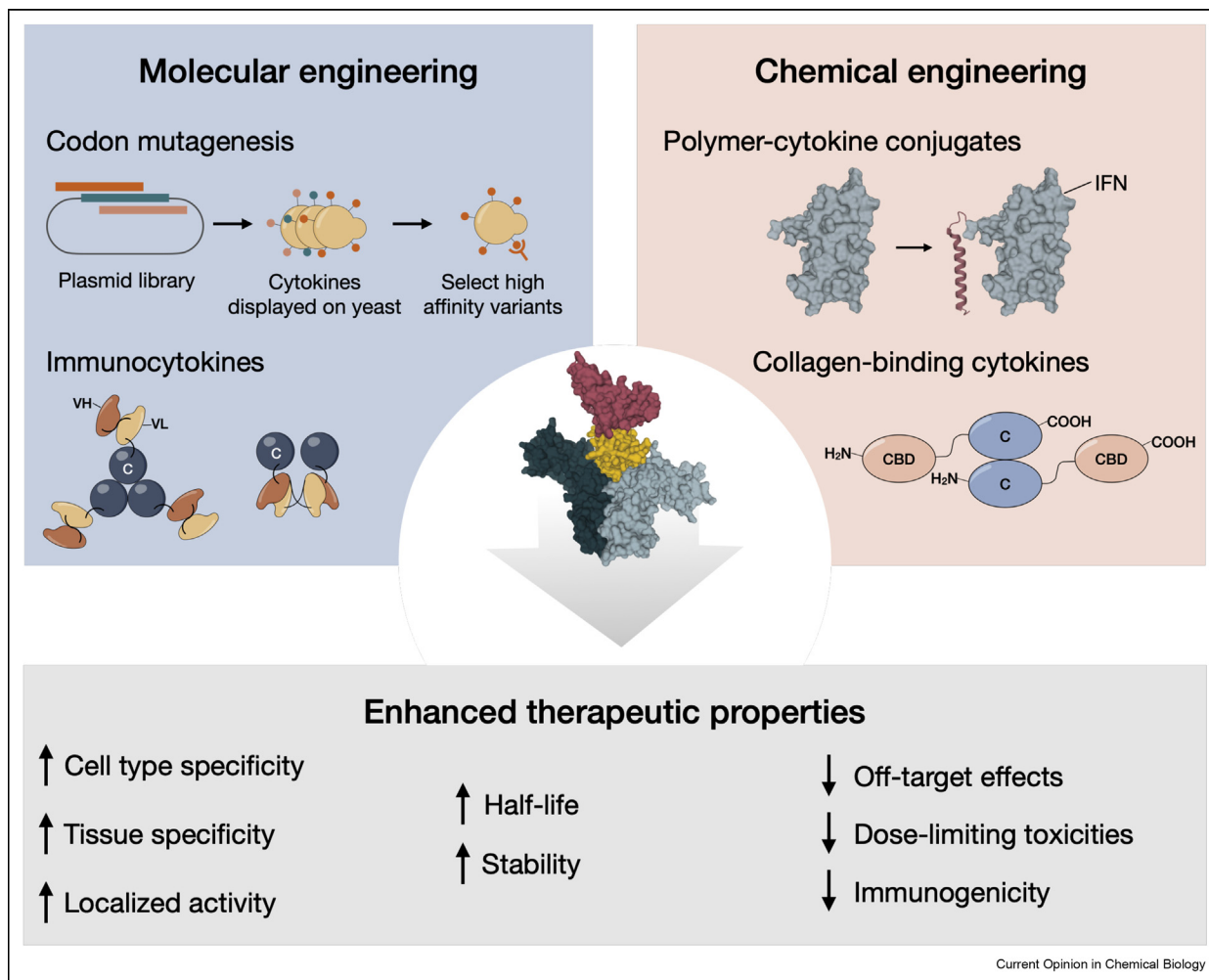
immunocytokines, including IL-2, IL-12, and TNF-based fusions for the treatment of different malignancies (extensively reviewed elsewhere [11,22–26]).

The architecture of the antibody fragment greatly impacts the targeting performance of immunocytokines. In general, the size and the spatial arrangements are considered key optimization parameters as they can influence binding affinity, pharmacokinetics, extravasation and tumor uptake, and effective doses [24]. Recently, sophisticated immunocytokine formats were developed for the treatment of so far refractory tumors. As an elegant example, Dougan et al. developed an alpaca-derived heavy-chain only fragment (VHH) targeting programmed death-ligand 1 (PD-L1) fused to IL-2 or IFN $\gamma$  to treat pancreatic ductal adenocarcinoma, which was unresponsive to native immune checkpoint blockade antibodies [27]. They showed that immunocytokines consisting of a small antibody fragment favored tumor penetration and conferred enhanced antitumor activity compared to their full-sized counterparts.

In addition, in a study by Weiss et al., several immunocytokines containing different L19-derived fragments, an antibody that targets a tumor-associated epitope of extracellular fibronectin, were generated for the treatment of glioblastoma [28]. Fusions of L19 fragments in single chain variable fragment (scFv) or diabody (a dimer of scFvs) formats with TNF $\alpha$  and IL-12 could enable effective antitumor activity both in mouse models and patients. Notably, the intravenous route of administration of the immunocytokines represents a significant advantage compared to previously used intratumoral injection route [29] or surgery for local delivery of a cytokine encoding viral vector [21].

Despite the promising results of immunocytokines, work by Tzeng et al. showed that the biodistribution of a

Figure 1



Overview of cytokine engineering strategies in the context of cancer immunotherapy. VH, heavy-chain variable domain; VL, light chain variable domain; C, cytokine; IFN, interferon; CBD, collagen-binding domain.

full size antibody-IL-2 immunocytokine was mainly governed by cytokine–receptor interactions rather than by tumor antigen–antibody fragment interactions [30]. This study pointed out the complexity of *in vivo* fate of immunocytokines, due to multiple potential targeting interactions, and the need for a case-by-case assessment of tumor targeting capacity. In summary, these preclinical and clinical studies highlight the potential of immunocytokines for tumor treatment. Nonetheless, robust optimization is required to find the right cytokine–antibody combination and structure.

### Mutagenesis

Mutagenesis technologies can be used to directly engineer cytokine's binding sites with altered receptor specificity and affinity. To date, different mutagenesis systems have been applied to cytokines including site-

directed mutagenesis and combinatorial library-based platforms, such as phage or yeast display [14].

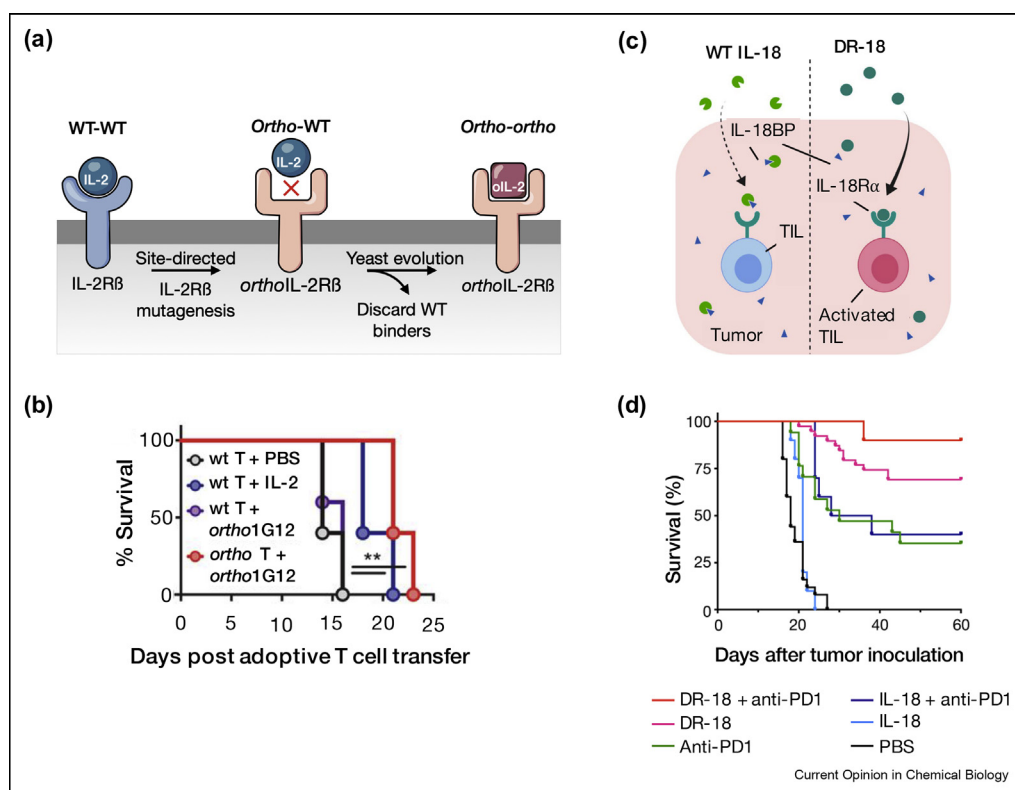
One cytokine that has been extensively explored for mutagenesis is IL-2, which plays a pivotal role in the immune response by promoting immune cell proliferation, differentiation, and cytolytic activity [31]. In the past decades, wild-type IL-2 has been used in the clinic for the treatment of several cancers, but it has been shown to induce severe toxicities and side effects [32]. These effects are mainly due to the pleiotropic nature of IL-2, which acts on effector cytotoxic immune cells as well as immunosuppressive regulatory T cells (Tregs). Specifically, IL-2 signals through activation of the IL-2 receptor (IL-2R), which exists in a dimeric or trimeric format. The intermediate-affinity dimeric receptor consists of an IL-2R $\beta$  and a  $\gamma_c$  subunit, whereas the

high-affinity trimeric receptor includes an IL-2R $\alpha$ , an IL-2R $\beta$ , and a  $\gamma_c$  subunit. Dimeric receptors are mainly expressed on cytotoxic T cells and natural killer (NK) cells, while trimeric receptors are predominantly found on Tregs. Importantly, this effect results in preferential stimulation of Tregs leading to immunosuppression in the TME [31].

In order to achieve favored activation of cytotoxic T cells for cancer therapy, many engineering efforts have been focused on altering the binding affinity of IL-2 towards distinct receptor subunits. Site-directed rational mutagenesis is based on the prediction of free energy changes upon amino acid mutation within a protein–protein interface. However, in the case of cytokine–receptor interactions, the underlying energetics are highly unpredictable. For this reason, site-directed mutagenesis approaches have been mostly used to weaken receptor–cytokine interactions [33–36],

e.g., the binding between IL-2 and the IL-2R $\alpha$  subunit. Conversely, yeast and phage display were successfully used to generate IL-2 variants with increased affinity to one or more receptor subunits. The Wittrup and Garcia groups pioneered the development of several IL-2 variants with redirected specificity towards one distinct IL-2R subunit resulting in preferential expansion of cytotoxic T cells and NK cells [37–39]. While engineered cytokines provide a potent tool to target distinct endogenous immune populations, they are still incapable of precisely targeting adoptively transferred cells. Recent work by Sockolosky et al. has provided convincing evidence that IL-2 specificity can be redirected towards adoptively transferred T cells using an orthogonal (*ortho*) cytokine–receptor system in which both cytokine and receptor are engineered to signal exclusively together (Figure 2a,b) [40]. Notably, engineered *ortho*IL-2 exhibited a high degree of specificity *in vivo*, negligible off-target toxicity, and elicited

Figure 2



**(a)** Schematic overview of orthogonal IL-2/IL-2R pairs consisting of engineered IL-2 and IL-2R that signal exclusively together and do not cross-react with wild-type (WT) cytokine and receptor. **(b)** Survival curves of mice bearing subcutaneous B16F10 tumors treated with adoptive transfer of pmel-1 transgenic CD8<sup>+</sup> T cells expressing WT IL-2R (wt T) or orthoIL-2R $\beta$  (ortho T), in combination with WT or orthogonal IL-2 mutant (ortho1G12). ortho, orthogonal. Adapted with permission from a study by Sockolosky et al. [40]. **(c)** Schematic illustration of the DR-18 technology. Engineered decoy-resistant IL-18 (DR-18) does not bind the decoy receptor IL-18 binding protein (IL-18BP) and maintains the capability to bind tumor-infiltrating lymphocytes. **(d)** Survival curves of mice bearing MC-38 tumors after the treatment with WT IL-18 plus anti-PD-1 antibody, or DR-18 alone or in combination with anti-PD-1 antibody. TIL, tumor-infiltrating lymphocytes; PBS, phosphate-buffered saline; anti-PD1, anti-programmed cell death protein 1 antibody. Adapted with permission from a study by Zhou et al. [42].

enhanced anticancer immune responses against B16F10 mouse melanoma tumors. Although engineering mutant receptor-cytokine pairs as highly potent as the wild-type remains challenging, this approach is promising for the specific expansion of genetically engineered T cells, such as CAR T cells. Alternatively, *de novo* cytokines can be designed and synthesized. Silva et al. used computational methods to elegantly demonstrate this idea [41]. They generated IL-2 and IL-15 mimetics, termed neoleukin-2/15, which specifically bind to the  $\beta$  and  $\gamma$  subunits of IL-2R with no requirement of binding to IL-2R $\alpha$  for signaling initiation. These synthetic cytokines showed a therapeutic activity superior to wild-type IL-2 in several tumor models with greatly reduced toxicity and immunogenicity.

Besides IL-2, other cytokines have been recently employed for mutagenesis. In a recent attempt to enhance the antitumor activity of IL-18, Zhou et al. used directed evolution to engineer a ‘decoy-resistant’ IL-18 variant (DR-18), which does not bind to the decoy receptor IL-18BP, thereby maintaining its activity in the circulation (Figure 2c,d) [42]. Strikingly, administering DR-18 to tumor-bearing mice resulted in a selective expansion of stem-like TCF1<sup>+</sup>CD8<sup>+</sup> T cells in tumor and biased differentiation towards polyfunctional T cells rather than towards an exhausted phenotype. In addition, DR-18-induced proliferation of NK cells in MHC class I-deficient tumors, providing a new tool for the treatment of immune checkpoint blockade therapy-resistant tumors. Another recent example of cytokine mutagenesis involves an engineered IL-10 mutant with increased affinity towards the low-affinity IL-10R $\beta$ , which acts as a strong gene expression regulator in both CD8<sup>+</sup> T cells and monocytes [43].

Collectively, these preclinical studies highlight the immense potential that mutagenesis-based engineering strategies hold for cancer immunotherapy. However, this technology provides solutions to only one drawback of wild-type cytokines, i.e., pleiotropy, thus inevitably passing on the engineered variants several other shortcomings coming from the parental molecule, such as short half-life and low stability. Therefore, the combination of mutagenesis and other cytokine engineering technologies (e.g., chemical engineering) may be necessary to obtain an optimal therapeutic cytokine molecule.

#### Chemical engineering of therapeutic cytokines

In addition to engineering cytokines by modifying their amino acid sequence, another approach to improve their pharmacological properties (e.g., solubility and pharmacokinetics) is through conjugation with synthetic polymers or peptides. Functional groups on the surface of cytokines are typically used to covalently link polymers. The most well-known strategy of this kind is the

conjugation of proteins with polyethylene glycol (PEG), known as PEGylation. PEGylation has gained vast success in prolonging cytokines’ half-life, leading to several clinically approved PEGylated cytokines [15]. Despite these successes, the varying protein bioactivity caused by nonspecific conjugation as well as the nondegradable nature of PEG have recently drawn increasing concerns [44,45]. To address these issues, biodegradable and non-immunogenic alternatives to PEG have emerged, such as amino acid-based synthetic polypeptide.

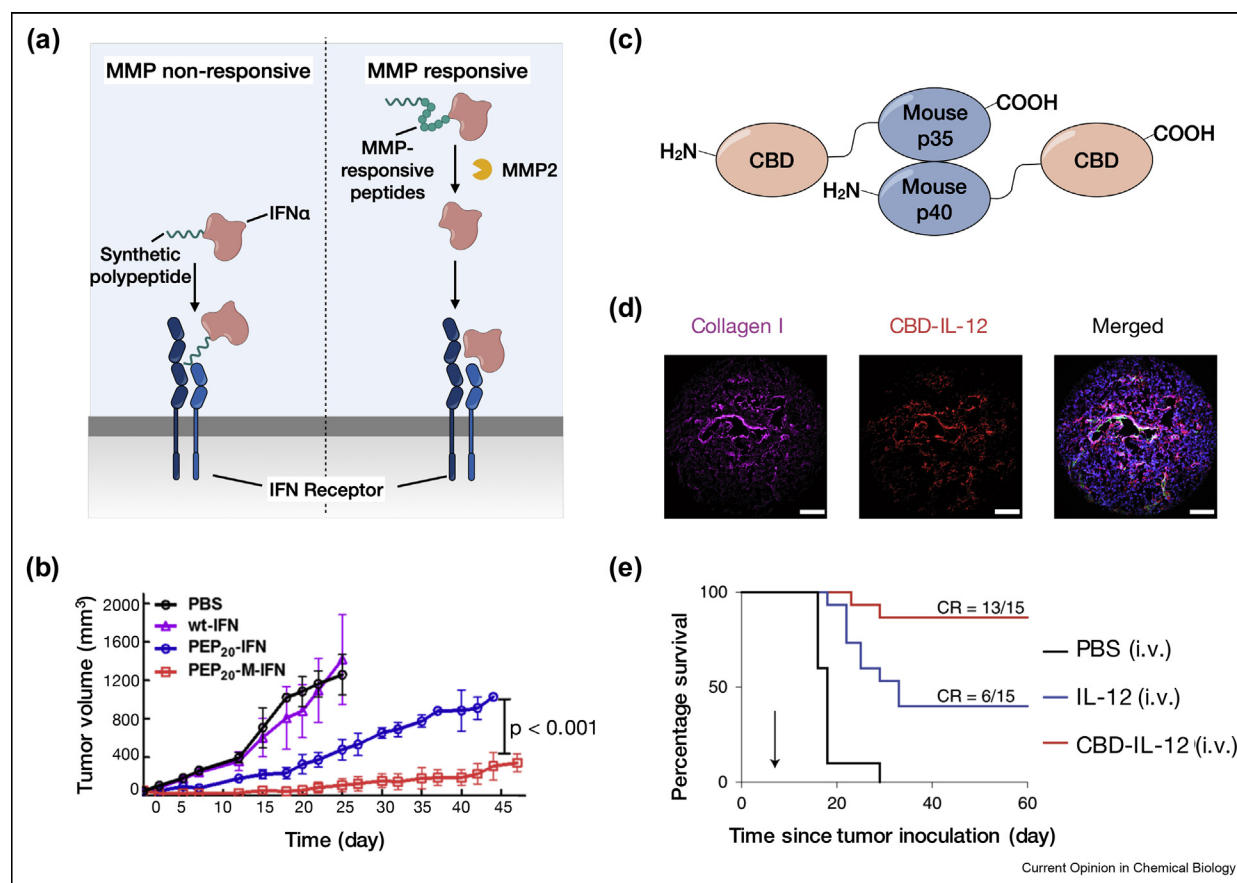
#### Alternatives to PEG

In recent years, a wide range of PEG alternatives has been investigated [46–49]. Despite the potential success of these methods, none of them yet provided satisfactory biodegradability and pharmacological performance. Chemical conjugation of synthetic polypeptides to therapeutic cytokines, known as PEPylation, represents a valid alternative to PEG because of its excellent chemical modularity and increased biodegradability [16]. In an elegant example, site-selective conjugation of polypeptide with protein was achieved upon the introduction of two orthogonal “chemical handles” at the C- and N-termini of the polypeptide [50]. Importantly, the PEPylated IFN $\alpha$  showed significantly enhanced protease resistance and thermostability. Building on this work, additional efforts were focused on employing cyclic polypeptide structures in the modification of antitumor therapeutic biomolecules. In two distinct studies, Lu and colleagues investigated the effects of a PEPylated IFN with a head-to-tail macrocyclic polypeptide [51] and another PEPylated IFN bearing a fixed helical polypeptide [52], respectively, in mouse models. Strikingly, both strategies resulted in improved protease resistance, thermostability, and cytotoxicity *in vitro*, as well as low immunogenicity *in vivo*. In particular, macrocyclization of the IFN–polypeptide conjugate displayed enhanced pharmacokinetics as well as tumor retention and tissue penetration. These examples provide promising insights into the use of PEPylated cytokines for cancer therapy.

Nevertheless, the engineering of tissue-specific and inducible cytokines remains challenging. Recently, Wang et al. took advantage of the overexpressed matrix metalloproteinases (MMPs) in the TME and designed a peptide–cytokine conjugate linked through a MMP-responsive peptide sequence between the IFN $\alpha$  structure and the synthetic polypeptide (Figure 3a, b) [53]. In the presence of high concentrations of MMP, the peptide substrate is cleaved, and the protein activity can be instantly restored to its full capacity. Moreover, systems that are responsive to multiple stimuli have been developed. For example, a temperature-responsive and MMP-cleavable IFN $\alpha$  have been recently engineered [54]. By fusing a body temperature-responsive elastin-like polypeptide, an MMP substrate, and IFN $\alpha$ , the



Figure 3



authors were able to achieve synergy of the body temperature responsiveness and MMP-cleavability, thus dramatically improving antitumor efficacy compared with the native IFN $\alpha$ . These strategies combine advantages afforded by PEPylation with benefits from unconjugated protein, including high tumor retention, deep tumor penetration, and intact antitumor potency.

Taken together, these studies suggest that the conjugation of synthetic polypeptides to cytokines represents a viable alternative to PEGylation, opening up new opportunities for the development of responsive release systems for cytokine therapeutics. However, the receptor-binding domains of PEPylated cytokines typically did not undergo any modification; thus, the risk of

off-target toxicities deriving from the pleiotropic nature of cytokines persists.

#### Extracellular matrix-targeting cytokines

The TME comprises many different types of cells including stromal cells as well as an evolving extracellular matrix (ECM) [55,56]. Collagen is a major component of the ECM contributing to the molecular architecture and mechanical properties of tissues [57]. Importantly, it has been observed that many tumors contain increased amounts of collagen compared to healthy tissues [58,59]. Early evidence that collagen could act as a promising target was shown by fusing a collagen-binding domain (CBD) to epidermal growth factor receptor (EGFR)-specific antibody fragments

[60,61]. Recently, a variety of ECM-binding cancer therapeutics were developed for enhanced tumor targeting [62,63]. For example, scientists from the Hubbell group elegantly demonstrated the potential of CBD-conjugated cytokines by fusing IL-2 to a peptide, which has high affinity to collagen I and III [64]. Strikingly, CBD-modified cytokines enhanced the tumor infiltration of CD8<sup>+</sup> T cells and achieved increased antitumor efficacy compared to their unmodified counterparts. This strategy took advantage of the increased vascular permeability of tumor blood vessels, which allow the CBD-conjugated cytokines to accumulate in the TME. Building on this work, CBD-conjugation was further used for the tumor-targeted delivery of several cytokines and chemokines including CCL4 and IL-12 leading to more effective and safer cancer immunotherapies [65,66].

IL-12 is considered an attractive antitumor cytokine as it can act on both the innate and adaptive immune system [65]. However, administration of IL-12 in clinical trials resulted in high accumulation in healthy tissues and limited penetration in tumor tissues [67]. In a recent report by the Wittrup lab, IL-12 was fused to lumican, a protein that binds collagen I and IV. Intratumoral injection of lumican-conjugated IL-2 and IL-12 induced increased cytokine retention in the tumor matrix [68]. In a following study, Mansurov et al. [65] developed tumor-targeted CBD-IL-12 therapy that could be administered intravenously (Figure 3c-e). In line with previous reports, CBD-IL-12 administration led to cytokine accumulation in both B16F10 and EMT6 tumors, resulting in enhanced intratumoral IFN $\gamma$  production and CD8<sup>+</sup> T cell recruitment. To summarize, conjugation of cytokine to ECM-targeting peptides/proteins represents a promising engineering approach for tumor-targeted delivery of immunostimulatory molecules for the treatment of solid tumors.

### Perspective

Cytokine-based immunotherapies have been used to treat various tumors. However, considerable toxicities and modest efficacies of therapeutic cytokines pose major challenges that require further cytokine engineering. In this review, we highlighted some of the most recent reports leveraging molecular and chemical engineering strategies to improve safety and efficacy of cytokine-based cancer therapy. However, significant challenges remain to be addressed.

Immunocytokines as well as chemically modified cytokines maintain their pleiotropic capability to target any cell type that expresses their receptor. This effect remains the leading cause of cytokine-derived toxicities and represents a considerable hurdle for clinical

translation. Finding the optimal balance between the activation of desired immune cells versus off-target cells (e.g., CD8<sup>+</sup> T cell versus Tregs) remains a key challenge. Future efforts in cytokine engineering will focus on cytokines with improved pharmacokinetics combined with more specific receptor-targeting capabilities. This may be achieved through masking the receptor-binding domains of cytokines using antibodies or synthetic polymer molecules, thus preventing binding to certain receptor subunits expressed only on undesired cells but not the target cells [69,70].

To achieve targeted immunotherapy, it is crucial to provide cytokines with the capability to discriminate between targeted and non-targeted tissues. We envision that engineering stimuli-responsive cytokine-based immunotherapies (e.g., a cytokine responsive to low pH in TME) will provide a promising solution to on-target-off-tumor toxicities. Moreover, regulating the timing of cytokine delivery with disease progression is crucial for optimal therapeutic performance. In recent years, advancements in cell and genetic engineering technologies have enabled researchers to leverage the capability of immune cells to migrate to tumors and recognize cancer cells for targeted delivery of cytokines. T cells have been engineered to secrete cytokines including IL-2, IL-7, IL-12, IL-15, IL-21, CCL19 [71–75], and the p40 subunit of IL-23 [76], or to carry cytokine-loaded nanoparticles [77,78]. Cell-based cytokine delivery provides inducible cytokine production allowing for precise tuning of the amplitude, duration, and localization of cytokines. Such strategies combining cell engineering with cytokine engineering have shown great promise. However, accompanied toxicities, such as autonomous growth of the cytokine-secreting T cells and cytokine storm, have to be assessed for clinical translation.

Combination therapy with multiple cytokines is an attractive option as it offers the possibility to achieve synergistic therapeutic effects. However, the molecular differences between cytokines lead to dramatically different pharmacological properties, making clinical translation challenging. A solution may reside in the generation of bi-/multispecific cytokines that bind two or multiple targets with one molecule. This strategy could lead to the development of one single molecular product, thus accelerating the approval procedures and reducing development costs. Nevertheless, whether different cytokines combined in one molecule still retain their activities needs to be evaluated.

In conclusion, the growing toolbox of cytokine engineering strategies that can be combined and tuned for maximum safety and efficacy will continue to enable the

development of novel therapeutic solutions for the treatment of refractory diseases.

## Author contributions

L.B. and L.T. planned and wrote the review.

## Declaration of competing interest

The authors declare that they have no known competing financial interests or personal relationships that could have appeared to influence the work reported in this paper.

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