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Review

Anticancer opportunities at every stage of chemokine function

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The chemokine system, comprising 48 chemokines and 23 receptors, is critically involved in several hallmarks of cancer. Yet, despite extensive efforts from the pharmaceutical sector, only two drugs aimed at this system are currently approved for clinical use against cancer. To date, numerous pharmacological approaches have been developed to successfully intervene at different stages of chemokine function: (i) chemokine availability; (ii) chemokine–glycosaminoglycan binding; and (iii) chemokine receptor binding. Many of these strategies have been tested in preclinical cancer models, and some have advanced to clinical trials as potential anticancer therapies. Here we will review the strategies and growing pharmacological toolbox for manipulating the chemokine system in cancer, and address novel methods poised for future (pre)clinical testing.

Chemokines and their receptors in cancer

Currently, 48 chemokines (*chemotactic cytokines*) and 23 chemokine receptors are documented as members of the chemokine superfamily [1]. Together, they function in a concerted manner to govern the migration and localization of all immune cells in the human body [2]. Accordingly, the chemokine system forms a highly relevant therapeutic target for a plethora of immune and inflammation-related disorders. In particular, the chemokine system is highly involved in cancer, with reported roles in almost all hallmarks of the disease, such as promoting angiogenesis, metastasis, and an immunosuppressive **tumor microenvironment (TME)** (see [Glossary](#)) ([Box 1](#)) [3]. Chemokines and their receptors can be expressed by many types of cells within the TME, including cancer cells, endothelial cells, and immune cells. From here, chemokines can regulate leukocyte recruitment to the TME, as well as leukocyte differentiation or polarization, which can lead to an immunosuppressive TME that favors tumor growth. In addition, chemokine signaling on tumor cells can directly promote cancer cell proliferation, survival, invasiveness and metastasis, cancer stem cell-like phenotypes, and angiogenesis [4–8] ([Box 1](#)). However, antitumor roles have also been described [4,7,8], which adds a layer of complexity when targeting this system. To date, only two drugs targeting the chemokine system are approved for clinical use in cancer: Plerixafor (anti-CXCR4) and Mogamulizumab (anti-CCR4), both against hematological malignancies. The paucity of clinically-approved anticancer drugs aimed at this system provides an impetus to review the current approaches to pharmacologically modulate the chemokine system and assess their respective (dis)advantages to aid in the rational design of future anticancer therapies.

Targeting the chemokine system: from chemokine secretion to downstream receptor signaling

Chemokine function involves different stages, each of which provides opportunities for therapeutic intervention ([Figure 1](#)). First, chemokines are secreted by a variety of cells either in homeostasis or upon induction by inflammatory stimuli. While some of these chemokines remain inside the underlying tissue, a fraction moves to the luminal surface of endothelial cells forming the blood

Highlights

The chemokine system has emerged as a relevant target in immune-oncology, with roles in most hallmarks of cancer.

Therapeutic targeting of this system can be achieved by interfering with chemokines, chemokine receptors, or glycosaminoglycans.

Many therapies aimed at the chemokine system are currently under clinical oncological investigation, including monoclonal antibodies targeting chemokines and their receptors, glycomimetics, and various peptides or small-molecule antagonists for chemokine receptors.

Other strategies showed preclinical anticancer value and appear poised for clinical testing, including nanobodies, small-molecule chemokine inhibitors, engineered chemokines, pepducins, siRNA, and biased ligands.

The emergence of chemokine receptor antagonists with novel mechanisms of action may prove more effective in the clinic.

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vessel wall, where they bind **glycosaminoglycans (GAGs)** with varying degrees of affinity [9,10]. These chemokine–GAG interactions prevent chemokines from rapidly diffusing away in the circulation and are thus key for forming a localized concentration gradient that directs leukocyte migration toward the site of secretion [9,10]. Patrolling leukocytes (or cancer cells) expressing a corresponding receptor then interact with these chemokines, eliciting multiple downstream signaling cascades and distinct biological effects, some of which are cancer-related (Box 1). In the following sections, we discuss the different strategies that interfere with these three key stages of chemokine function in cancer: (i) chemokine availability; (ii) chemokine–GAG interactions; and (iii) chemokine-receptor binding. Given the magnitude of the chemokine system, this work delineates a selection of exemplary cases rather than a comprehensive list of all (pre)clinical therapies targeted at the chemokine system in cancer. An overview of the discussed anticancer strategies is provided in Table 1 (Key table).

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Interfering with chemokine availability

Biologics

One strategy for interfering with tumorigenic chemokine signaling is to preclude the presence of chemokines at the tumor site by neutralization with **monoclonal antibodies (mAbs)**. mAbs offer many desirable features as therapeutics, including a long serum half-life, high selectivity and specificity, and indirect mechanisms of action such as **antibody-dependent cell-mediated/cellular cytotoxicity (ADCC)** or **complement-dependent cytotoxicity (CDC)**

Box 1. The chemokine system in various hallmarks of cancer

Many chemokines and chemokine receptors play key roles in various hallmarks of cancer (Figure 1).

Sustaining cancer cell proliferation

Chemokines can directly activate downstream signaling cascades to enhance tumor cell proliferation, such as PI3K/AKT/NF- κ B by CXCL16, MAPK/ERK by CCL5, and the STAT3 pathway by CCR5 and CCR7 [3,5]. They can also indirectly induce proliferation by transactivation of epidermal growth factor receptor (EGFR), as reported for CXCR4/ACKR3 [89]. Conversely, CXCL14 can suppress proliferative signaling in various cancer types [4].

Altering stress response to favor cell survival

Cell survival can be enhanced by modulation of proapoptotic and antiapoptotic proteins. For example, CCR5 was found to upregulate proapoptotic proteins and down-regulate antiapoptotic proteins via activation of NF- κ B [3]. Similarly, activation of CXCR4, CCR3, CCR7, and CCR8 induces inactivation of proapoptotic proteins via the ERK pathway. Chemokine signaling can also protect cancer cells from autophagic death, such as CXCR4 via the PI3K-mTOR pathway. Finally, chemokine receptors can interact with tumor suppressors, such as TP53, to regulate their anticancer activity [3].

Inducing angiogenesis

Chemokines, such as CXCL8 and CCL2, can promote tumor angiogenesis by directly targeting vascular endothelial cells and promoting their survival, synergizing with vascular endothelial growth factor, or recruiting endothelial progenitor cells. In general, most chemokines have been reported to stimulate angiogenesis; however CCL21, CXCL4, CXCL9, CXCL10, CXCL11, and CXCL14 appear to inhibit it [3,5,6].

Promoting invasion and metastasis

Chemokine signaling can regulate cancer cell invasiveness and metastasis by guiding cell migration and inducing epithelial–mesenchymal transition (EMT). For instance, CXCR4 expression is correlated with metastasis to the bone marrow, lymph nodes, and lungs, which produce high levels of CXCL12 [6,89]. Similarly, the CCL19/CCL21–CCR7 signaling axis is fundamental for lymph node metastasis [6,61]. CCL2, CCL18, and CXCL8 are examples of chemokines that induce EMT [4].

Immune modulation and the TME

The chemokine system can regulate the immune system to avoid cancer cell destruction or induce tumor-promoting inflammation. For instance, the CXCL16–CXCR6 axis drives the differentiation of macrophages toward a tumor-promoting phenotype. CCL2, CCL5, and CXCL12 recruit immunosuppressive leukocytes, such as tumor-associated macrophages and myeloid-derived suppressor cells [7]. Conversely, the CXCL9/CXCL10–CXCR3 axis can enhance tumor immune destruction [4].

Metabolic rewiring

The CXCL12–CXCR4 axis regulates expression and secretion of phosphoglycerate kinase 1, an enzyme involved in metabolic reprogramming from oxidative phosphorylation to glycolysis [3].

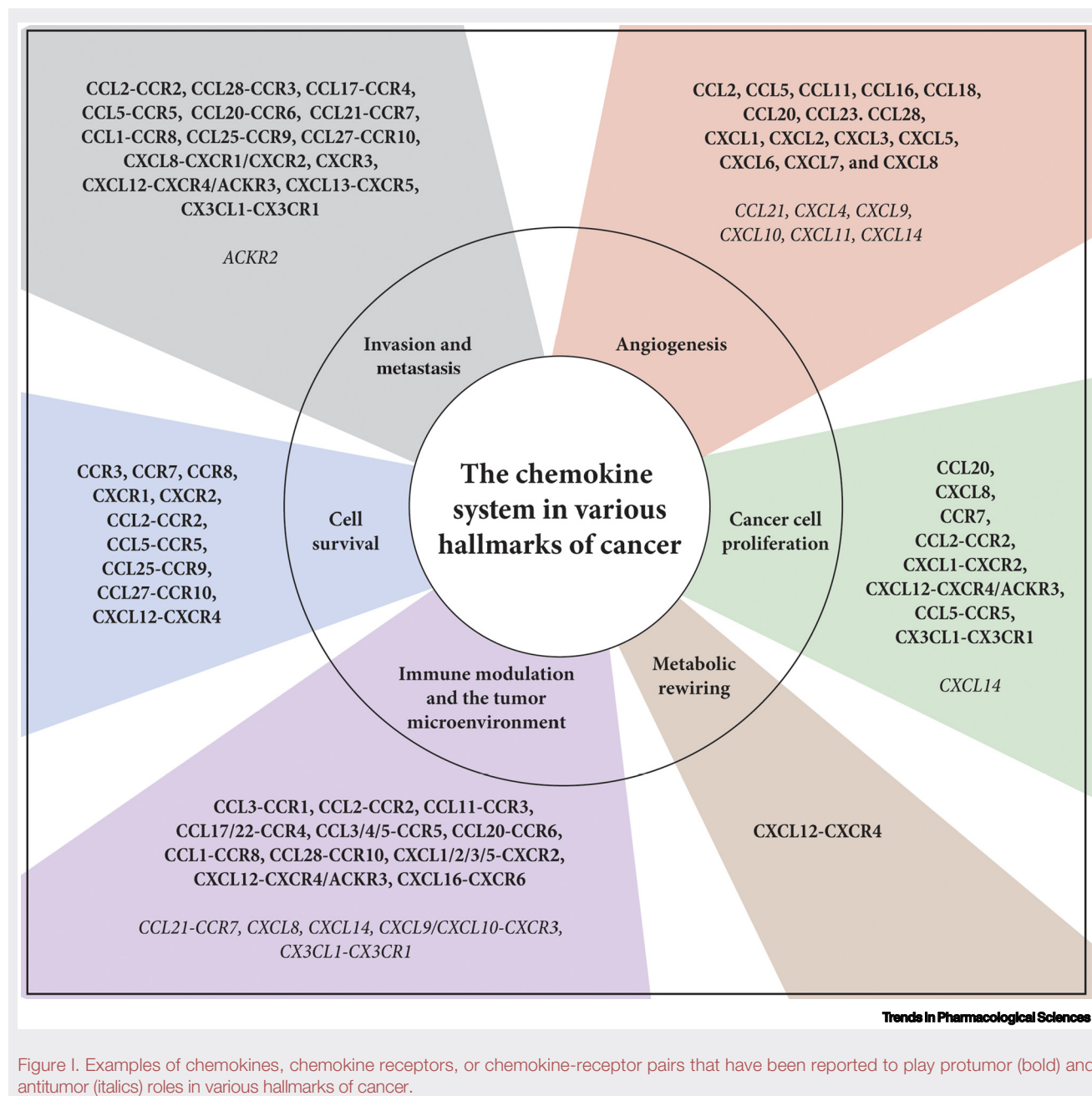
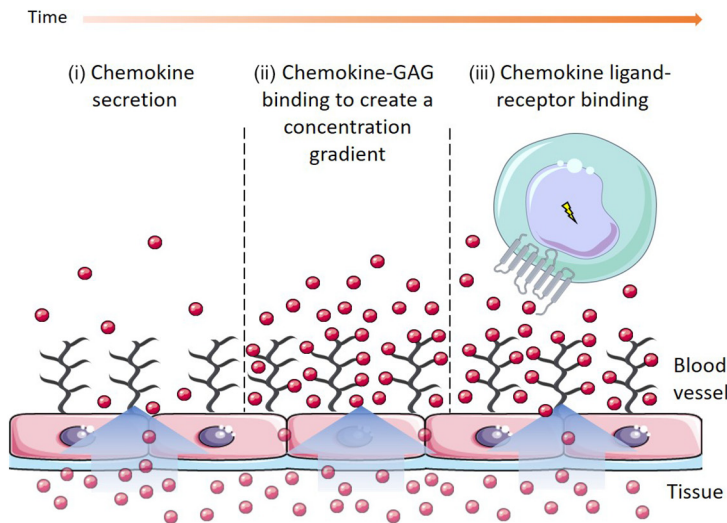


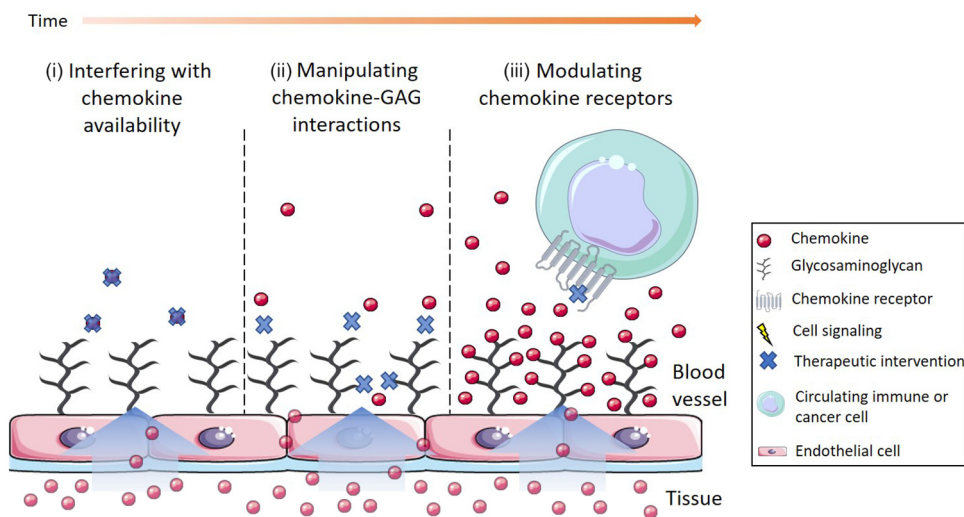
Figure 1. Examples of chemokines, chemokine receptors, or chemokine-receptor pairs that have been reported to play protumor (bold) and antitumor (italics) roles in various hallmarks of cancer.

[11, 12]. Chemokine-neutralizing mAbs have shown some success in preclinical cancer models with reduction of TME tumorigenic signaling, tumor growth, angiogenesis, and metastasis [13, 14]. For example, blocking CCL1-CCR8 signaling with an anti-CCL1 antibody attenuated the immunosuppressive function of regulatory T cells (T_{reg} s) in the TME, without inhibiting effector T cell function in a mouse model of breast cancer [13]. In addition, treatment with an anti-CXCL1 mAb, which prevents signaling mainly via CXCR2, inhibited angiogenesis, reduced tumor growth,

(A) (Patho-)physiological signaling



(B) Therapeutic modulation at each stage of chemokine functioning



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Figure 1. Avenues for therapeutic intervention in chemokine ligand–receptor signaling. (A) Schematic visualization of the steps involved in chemokine ligand–receptor signaling over time. (i) chemokine secretion; (ii) chemokine–glycosaminoglycan (GAG) binding to create a concentration gradient; and (iii) chemokine ligand–receptor binding. (B) Schematic visualization of how therapies can interfere with these steps. Therapies can: (i) intervene with chemokine availability; (ii) prevent chemokines from binding to GAGs and thus, precluding a chemokine gradient; or (iii) block chemokine receptors to prevent chemokine binding and (oncogenic) downstream signaling. This figure was created using elements from Servier medical art.

and induced apoptosis in mice bearing bladder and prostate tumors [14]. The CXCL8–CXCR1/CXCR2 signaling axis has also been targeted with chemokine antibodies. In fact, the anti-CXCL8 antibody BMS-986253 became the first chemokine antibody to undergo clinical trials in patients with advanced solid tumors. A Phase I study showed the antibody to be well-tolerated and effective at reducing serum CXCL8 levels [15]. BMS-986253 is currently under clinical investigation as co-therapy with anti-programmed cell death (PD)-1 antibody Nivolumab in advanced

Glossary

Allosteric binding site: any site(s) in a protein where molecules can bind other than the binding site of the endogenous ligand(s)/substrate(s).

Antibody-dependent cell-mediated/cellular cytotoxicity (ADCC): a mechanism whereby effector immune cells bind to the Fc region of an antibody–tumor cell complex and subsequently secrete factors leading to lysis of the tumor cell.

Biased signaling: preferred activation of one signal transduction pathway over another upon a ligand binding to a receptor

Chemotaxis: directed movement of cells toward a molecular concentration gradient (e.g., of chemokines).

Complement-dependent cytotoxicity (CDC): a mechanism whereby the complement protein C1q binds to the Fc region of an antibody–tumor cell complex to induce tumor cell lysis through the complement pathway.

Glycosaminoglycans (GAGs): long, linear, and highly charged polysaccharides, that can occur as soluble entities in the extracellular matrix or in a bound form as part of endothelial cell-surface proteoglycans that, in turn, form the glycocalyx.

Insurmountable antagonism: the ability of an antagonist to decrease the maximum level of receptor activation despite a high concentration of the endogenous agonist, as well as to possibly decrease the agonist’s potency.

Monoclonal antibodies (mAbs): antibodies produced by clones of a single B-lymphocyte fused with an immortal myeloma cell. These biologics comprise two antigen-binding fragments (Fab) and a fragment crystallizable region (Fc).

Multitarget ligands: molecules that can bind more than one protein target with reasonable selectivity (i.e., other than unintended promiscuous target binding).

Nanobodies: antibodies consisting of only a single antigen-binding fragment, derived from camelids. Due to their single-domain nature, they are much smaller than monoclonal antibodies.

Orthosteric binding site: the site in a protein where its endogenous ligand(s)/substrate(s) bind(s).

Peptiducins: lipid-linked peptides with an amino acid sequence derived from one of the intracellular loops or the C-terminus of the target G protein-coupled receptor (GPCR). The lipid moiety

Key table

Table 1. Overview of discussed (pre)clinically tested strategies to modulate the chemokine system in cancer

Target	Name/type of therapy	Model/latest stage clinical trial	Key result/comment	Refs/identifiers
Chemokine-oriented therapies				
CCL1	α -CCL1 (antibody)	<i>In vivo</i> BALB-neuT mouse TUBO model of breast cancer	Reduced immunosuppressive function and <i>de novo</i> conversion of T _{regs}	[13]
CCL2	Carlumab (antibody)	Phase II clinical trials (castrate-resistant prostate cancer)	Lack of antitumor activity and elevated CCL2 upon therapy cessation	[25]
CCL18	SMC-21598 (small molecule)	<i>In vivo</i> NOD-SCID mouse MDA-MB-231 model of breast cancer	Reduced lung metastasis but not tumor growth	[26]
CXCL1	HL2401 (antibody)	<i>In vitro/in vivo</i> , bladder (T24) and prostate (DU145, PC3) cancer cells, xenograft mouse models	Inhibited angiogenesis and proliferation, and induced apoptosis <i>in vivo</i> . Inhibited cancer cell proliferation and invasion <i>in vitro</i>	[14]
CXCL8	BMS-986253 (antibody, a.k.a. HuMax-IL8)	Phase I/II clinical trials (advanced cancers, hormone-sensitive prostate cancer, non-small cell lung cancer, hepatocellular carcinoma, head and neck squamous cell carcinoma)	Currently ongoing. In combination with anti-PD1 antibody, Nivolumab, luteinizing hormone-releasing hormone antagonist, or anti-CTLA-4 antibody	NCT03400332, NCT03689699, NCT04050462, NCT04848116, NCT04123379
CXCL10	3NB12 (nanobody)	<i>In vitro</i> MDA-MB-231 mouse model of breast cancer	Reduced chemotaxis	[17]
CXCL11	11B1 and 11B7 (nanobodies)	<i>In vitro</i> mouse L1.2 model of precursor B cell lymphoma	Reduced chemotaxis	[18]
CXCL12	1A4 (nanobody)	<i>In vitro</i> mouse L1.2 model of precursor B cell lymphoma	Reduced chemotaxis	[18]
Glycosaminoglycan-oriented therapies				
CCL5 ^a	OTR4120 and OTR4131 (glycomimetic)	<i>In vitro</i> Huh7 and Hep3B hepatocellular carcinoma model	Reduced CCL5-induced tumor cell migration	[30]
CXCL12 ^a	RGTA (several glycomimetics)	<i>In vitro</i> Huh7 model of hepatocellular carcinoma	Reduced chemotaxis and anchorage-independent cell growth	[31]
CXCL12, VEGF-A, FGF2, and P-selectin ^a	M402 (glycomimetic, a.k.a. Necuparanib)	Phase II clinical trial (primary metastatic pancreatic cancer)	In combination with chemotherapy. Insufficiently improved clinical outcome	[35]
CXCR4- and E-selectin specific ^a	GMI-1359 (glycomimetic)	Phase Ib clinical trials (metastatic breast cancer)	Currently ongoing	NCT04197999

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becomes tethered in the cell membrane and anchors the peptide at the intracellular binding site of the target GPCR.

Peptibodies: one or more peptides fused to a fragment crystallizable region derived from an antibody to enhance the circulation half-life of the peptide(s). Fusion of multiple peptides enables multitarget binding.

Small interfering ribonucleic acids (siRNAs): short-length single or double-stranded RNA molecules that can prevent the expression of a target gene.

Tumor-associated macrophages (TAMs): macrophages that are present in the tumor microenvironment. They are derived from monocytes that are recruited to the tumor microenvironment, or from tissue-resident macrophages. These macrophages become polarized to a protumorigenic phenotype by the cues from this environment.

Tumor microenvironment (TME): the immediate surroundings of cancer cells, including blood vessels, infiltrating immune cells, fibroblasts, extracellular matrix, epithelial cells, and other (possibly tumor-produced) extracellular signaling factors. Cancer cells are influenced by the TME, for instance, in their development and response to therapy.

Table 1. (continued)

Target	Name/type of therapy	Model/latest stage clinical trial	Key result/comment	Refs/identifiers
GAGs (syndecan-4)	dnCCL2-HSA (modified chemokine)	<i>In vivo</i> C57BL/6 mouse MC-38, 3LL, and LLC1 models of colon and lung cancer	Significantly reduced lung metastasis	[29]
GAGs (heparin, heparan sulfate)	CXCL12 α (modified chemokine)	<i>In vivo</i> SCID mouse LMD231 model, lung-metastasized breast cancer	Significant reduction of liver metastasis	[28]
Chemokine receptor-oriented therapies				
CCR2	Plozilzumab (antibody, a.k.a. TAK-202 and MLN1202)	Phase II clinical trial (solid tumor with bone metastases)	Reduced bone turnover rates in 14% of 41 patients, serious toxicity in 7%	NCT01015560
CCR2	PF-04136309 (orthosteric small molecule)	Phase Ib clinical trials (pancreatic cancer)	In combination with chemotherapy 16/33 patients achieved objective response. Toxicity concerns	[48,49]
CCR2	CCX872-B (orthosteric small molecule)	Phase Ib clinical trial (pancreatic cancer)	In combination with chemotherapy. Currently ongoing	NCT02345408
CCR2	CNP/siCCR2 (siRNA)	<i>In vivo</i> BALB/c mouse 4T1 model of breast cancer	Reduced CCR2 and TAM levels, tumor growth, and metastatic behavior	[76]
CCR4	Mogamulizumab (antibody, a.k.a. Poteligeo)	Approved clinical use for cutaneous T cell lymphoma	Currently on the market	
CCR4	FLX475 (orthosteric small molecule)	Phase I/II clinical trials (advanced cancer, metastatic gastric cancer, advanced melanoma)	As monotherapy or in combination with anti-PD1 antibody or anti-CTLA-4 antibody. Currently ongoing	NCT03674567, NCT04768686, NCT04894994
CCR5	Maraviroc (orthosteric small molecule)	Phase I clinical trials (metastatic colorectal cancer)	Reduced tumorigenic signaling in the TME in one trial, another trial is ongoing	[51,52], NCT04721301
CCR5	Leronlimab (antibody, a.k.a. PRO140)	Phase Ib/II clinical trials (CCR5 ⁺ metastatic triple-negative breast cancer/solid tumors)	Clinical studies are ongoing	NCT04313075, NCT03838367, NCT04504942
CCR8	Anti-CCR8 antibody (α CCR8)	<i>In vivo</i> BALB/c mouse CT26 and MC38 model of colorectal cancer	Enhanced effector T cell tumor infiltration and long-term survival	[37]
CCR9	91R and 92R (antibodies)	<i>In vitro/in vivo</i> BALB/c Rag2 ^{-/-} mouse MOLT-4 model of leukemia	Reduced <i>in vivo</i> tumor growth (91R/92R), angiogenesis (91R) (leukemia)	[38,39]
CXCR1	Anti-CXCR1 antibody	<i>In vitro</i> H460 and MOR/P non-small cell lung cancer cells	Significantly attenuated cancer cell proliferation	[36]

Table 1. (continued)

Target	Name/type of therapy	Model/latest stage clinical trial	Key result/comment	Refs/identifiers
CXCR1/CXCR2	Reparixin (allosteric small molecule)	Phase II (metastatic triple-negative breast cancer)	Well-tolerated and reduced cancer stem-cell population	[60]
CXCR1/CXCR2,	X1/2pal-i3 (pepducin targeting intracellular loop 3)	<i>In vivo</i> NCR Nu/Nu mouse OVCAR-4 model of ovarian cancer. <i>In vivo</i> SCID mouse PC3, DU145 and C4-2 xenograft models of prostate cancer	Impaired tumor growth and angiogenesis in ovarian cancer. Reduced antiapoptotic protein expression and tumor growth in PTEN-deficient prostate cancer	[72,73]
CXCR2	AZD5069 (orthosteric small molecule)	Phase I/II clinical trials (metastatic head and neck carcinoma, pancreatic ductal carcinoma, and castration-resistant prostate cancer)	Low objective response rate and high toxicity risk in the pancreatic carcinoma study. The other studies are ongoing	NCT02583477, NCT02499328, NCT03177187
CXCR2	Navarixin (allosteric small molecule, a.k.a. SCH-527123 and MK-7123)	Phase II clinical trial (non-small cell lung cancer, castration-resistant prostate cancer, and colorectal cancer)	In combination with anti-PD1 antibody. Currently ongoing	NCT03473925
CXCR4	Mavorixafor (allosteric small molecule, a.k.a. X4P-001, AMD-070, and AMD-11070)	Phase I/II clinical trials (advanced renal cell carcinoma, Waldenström's macroglobulinemia, and advanced melanoma)	In combination with kinase inhibitor, or anti-PD1 antibody. Preliminary results show low toxicity and increased antitumor CD8 ⁺ T cell TME levels	[62], NCT04274738
CXCR4	Ulocuplumab (antibody, a.k.a. BMS-936564 and MDX-1338)	Phase II clinical trials (pancreatic adenocarcinoma, small cell lung cancer, and Waldenström's macroglobulinemia)	Insufficient therapeutic response and high toxicity. Ongoing trial with Waldenström's macroglobulinemia	NCT02472977, NCT03225716
CXCR4	238D2 and 238D4 (nanobodies)	<i>In vitro</i> , Jurkat leukemia cells	Attenuated CXCL12-mediated cancer cell migration	[41]
CXCR4	VUN400-Fc (nanobody-Fc construct)	<i>In vitro</i> , CCRF-CEM leukemia cells	Induced CDC- and ADCC-mediated cancer cell killing	[42]
CXCR4	Plerixafor (orthosteric small molecule, a.k.a. AMD3100 and Mozobil)	Approved for clinical use against lymphoma and myeloma	Currently on the market. Ongoing Phase II trials for use in other cancer types	NCT04177810, NCT03746080
CXCR4	Motixafor tide (peptide, a.k.a. BL8040 and BKT140)	Phase III (multiple myeloma). Also in clinical trials for other cancer types	Currently ongoing. Preliminary results show low toxicity and enhanced levels of circulating hematopoietic stem cells	[64]

(continued on next page)

Table 1. (continued)

Target	Name/type of therapy	Model/latest stage clinical trial	Key result/comment	Refs/identifiers
CXCR4	LY2510924 (peptide)	Phase II (metastatic clear cell renal cell carcinoma, extensive-stage small cell lung carcinoma)	Low toxicity, but no improved outcome over standard treatment	[68,69]
CXCR4	Balixafortide (peptide, a.k.a. POL6326)	Phase III (HER2 ⁻ locally recurrent or metastatic breast cancer)	Currently ongoing	NCT03786094
CXCR4	e23sFv-9R/CXCR4si (siRNA)	<i>In vivo</i> BALB/c nude mouse HER2 ⁺ BT-474 model, breast cancer	Reduced CXCR4 expression to inhibit tumor growth and metastasis	[75]
CXCR4	PZ-210 and PZ-218 (peptiducins targeting intracellular loop 3 and 1, respectively)	<i>In vitro</i> patient cells, and <i>in vivo</i> NSG mouse Raji lymphoma model, leukemia and lymphoma	Increased survival (PZ-210). Blocked migration of lymphoma cells, and induced apoptosis in patient cells <i>in vitro</i> (both)	[74]
ACKR3 (CXCR7)	NB4 (nanobody)	<i>In vivo</i> nude mouse 22A model, head and neck cancer	Reduced tumor growth and angiogenesis	[43]

^aNote that for glycomimetics, the chemokine for which the compound was tested is shown. It cannot be ruled out that they also affect other soluble GAG-binders (e.g., other chemokines), as their selectivity was not always reported.

solid cancers, prostate cancer, and hepatocellular carcinoma, among others (NCT03400332, NCT03689699, NCT04050462, NCT04848116, and NCT04123379).

In addition to mAbs, several **nanobodies** have been developed to block chemokines. These nanobodies are much smaller (~10 times) than antibodies, which facilitates production, oral administration, and tumor penetration, but results in a short serum half-life [16]. Thus far, nanobodies targeting CCL2, CCL5, CXCL10, CXCL11, and CXCL12 have been reported [17,18]. *In vitro* studies showed **chemotaxis** inhibition of CXCR3-overexpressing breast cancer cells by a CXCL10 nanobody [17]; as well as chemotaxis inhibition of CXCR3 or CXCR4-expressing murine precursor B cell lymphoma cells by nanobodies against CXCL11 and CXCL12 [18]. However, *in vivo* (pre)clinical data is currently lacking to evaluate the potential of nanobodies as cancer therapeutics. Finally, biologics have been developed to alter chemokine availability by delivering specific chemokines to attract antitumoral immune cells to the TME, as reported previously with CCL16-chTNT3 (anti-necrotic DNA antibody) and CCL21-B3 (anti-PDL1 nanobody) conjugates [19,20].

These illustrative cases suggest that chemokine neutralization with biologics may be a viable anticancer strategy. However, previous work also revealed caveats that require consideration. First, Bonvin *et al.* demonstrated that some antibodies could recognize both free and GAG-bound chemokines, while others only recognized the free form [21]. Furthermore, the authors found that the antibody targeting the free form elicited a much better therapeutic response compared to the one that recognizes both free and GAG-bound forms. As the vast majority of chemokines are GAG-bound, antibodies that recognize this form will be drained by chemokine–GAG complexes, leaving the free chemokines available to interact and signal via chemokine receptors.

Thus, this study suggests that anti-chemokine antibodies should be designed to only recognize the free chemokine form to maximize therapeutic effect. Similarly, the surface charge of CCL21 in the aforementioned CCL21-nanobody conjugate required engineering to prevent sequestration to GAGs [20]. Second, studies on the CCL2-CCR2 axis illustrate the risk of cessation-induced complications. Bonapace *et al.*, found that stopping the anti-CCL2 antibody treatment in syngeneic mouse models of metastatic breast cancer resulted in CCL2 expression exceeding pretreatment levels at the metastatic site [22]. This, in turn, translated to a higher incidence of metastasis and higher mortality. A similar increase in metastasis after cessation of anti-CCL2 therapy was reported by Kersten *et al.*, while continued treatment led to metastasis inhibition [23]. Although no exacerbated metastasis was found in Phase I and II clinical trials with the anti-CCL2 antibody Carlumab in prostate cancer patients, CCL2 levels exceeding pretreatment levels were found during administration and after dosing [24,25].

Small molecules

Chemokines can also be neutralized with small molecules. Contrary to biologics, small molecules are easily manufactured, have a smaller size, and high stability that enables oral administration. Yet, thus far only one chemokine, CCL18, has been targeted with a small molecule in a cancer setting [26]. *In vivo* experiments showed that injection of CCL18 and small-molecule CCL18-antagonist SMC-21598 into breast cancer xenografts in mice reduced lung metastasis compared to injection of CCL18 alone. This effect was attributed to the small-molecule binding to CCL18, preventing CCL18 from interacting with its cognate receptors to elicit downstream signaling [26]. In this regard, CCL18 signals via several receptors, including CCR8, membrane-associated phosphatidylinositol transfer protein 3 (PITPNM3 or ACKR6), and G protein-coupled receptor for estrogen (GPR30) [27].

Manipulating chemokine–GAG interactions

Modified chemokines

A second strategy to impair chemokine-induced oncogenic signaling is to interfere with chemokine–GAG interactions. One way to do this is to administer chemokine mutants that are modified to exhibit enhanced GAG-binding and impaired chemokine receptor binding. Indeed, such mutants can disrupt the formation of a chemokine gradient by displacing endogenous chemokines from GAGs and blocking newly-secreted chemokines from binding GAGs. In addition, due to their impaired receptor binding, they prevent the recruitment of chemokine receptor-expressing cells (e.g., cancer cells). This mechanism can yield anticancer effects, as shown by studies with CXCL12 and CCL2, which interact with CXCR4/ACKR3 and CCR2, respectively [28,29]. Administration of lung-metastasized breast cancer cells, along with an engineered CXCL12 variant (CXCL12 α), significantly reduced the amount of liver metastases in mice, compared to control [28]. Similarly, mice that were treated with an engineered CCL2 variant (dnCCL2) before and after injection of colon cancer or Lewis lung carcinoma cells, showed significantly reduced lung metastasis compared to untreated mice [29]. In both cases, the antimetastatic effects were attributed to the modulation of the metastatic microenvironment in the target tissue. Notably, the CCL2 mutant only exerted its antimetastatic effect when fused to human serum albumin (HSA), as it otherwise had insufficient serum half-life, presenting an important consideration for future *in vivo* studies.

Glycomimetics

The large structural overlap of the GAG binding epitopes and the receptor binding epitopes on chemokines can also be exploited in anticancer therapies. For instance, exogenous GAGs or molecules that mimic GAGs (i.e., glycomimetics), can be used to hinder chemokine ligand–receptor binding. Indeed, Sutton *et al.* [30] showed that binding of CCL5 to CCR1-expressing hepatocellular

carcinoma cells could be inhibited by preincubation of CCL5 with the glycomimetics OTR4120 or OTR4131. As a result of this impaired CCL5–CCR1 interaction, CCL5-induced hepatoma cell migration was strongly attenuated *in vitro* [30]. In subsequent studies, glycomimetics effectively inhibited CXCL12–CXCR4-mediated cell growth and chemotaxis of hepatocellular carcinoma cells *in vitro* [31]. These precedents suggested a potential for glycomimetics as anticancer strategy, and paved the way for the glycomimetic GMI-1359 to be investigated in prostate cancer preclinical studies, which showed reduced metastasis as monotherapy or in combination with chemotherapy *in vivo* [32]. After completion of a Phase I trial in healthy volunteers (NCT02931214), GMI-1359 is currently undergoing a Phase Ib clinical trial (NCT04197999) to investigate its safety and tolerability in patients with hormone receptor positive breast cancer.

A potential caveat in the design of glycomimetics is their selectivity profile, as exogenous GAGs may naturally target more soluble GAG-binding elements than chemokines. Although this may raise safety concerns, promiscuous GAG-binding might actually be advantageous in some cases. For example, the GAG-mimetic Necuparanib was found to bind vascular endothelial growth factor (VEGF), fibroblast growth factor 2 (FGF2), P-selectin, and CXCL12 [33]. As a result, Necuparanib attenuated VEGF-A and FGF2-induced angiogenesis, P-selectin-mediated tumor cell seeding, and CXCL12-induced cell migration *in vitro*. This translated to a significant survival benefit in an orthotopic 4T1 murine mammary carcinoma model characterized by spontaneous metastasis [33]. Results from a Phase I clinical trial indicated acceptable safety of Necuparanib in patients with metastatic pancreatic cancer [34]; however, a Phase II clinical trial in metastatic pancreatic ductal adenocarcinoma patients showed no improved outcome upon addition of Necuparanib to standard therapy. Reasons for failure may include an insufficient dose to overcome the TME, the advanced stage of disease, or the presence of intrinsic resistance mechanisms in pancreatic cancer [35].

Modulating chemokine receptors

Biologics

Chemokine receptors can also be targeted by biologics to block chemokine binding. The therapeutic value of anti-chemokine receptor antibodies is best exemplified by the anti-CCR4 antibody Mogamulizumab, which is currently approved for clinical use in Japan, the United States, and Europe against adult or cutaneous T cell lymphoma. Besides Mogamulizumab, many other anti-chemokine receptor antibodies have been tested in a variety of preclinical studies [12]. Antibodies against CXCR1 [36], CCR8 [37], and CCR9 [38,39] were respectively found to reduce cancer cell proliferation *in vitro* [36], attenuate tumor growth and improve long-term survival in colorectal cancer mouse models [37], and impair angiogenesis and tumor growth in a xenograft mouse model of human acute lymphoblastic leukemia [38,39]. Furthermore, mAbs against CCR2, CCR4, CCR5, and CXCR4 have advanced to clinical trials as monotherapy or combination therapy in a variety of cancers (Table 1) [5,12,40]. As this topic has been extensively reviewed elsewhere [12], we will not cover it further.

Jähnichen *et al.* were the first to show that chemokine receptors can also be targeted by nanobodies [41]. They engineered two nanobodies, 238D2 and 238D4, which bind different epitopes at the extracellular site of CXCR4. Both nanobodies effectively inhibited CXCL12 binding to CXCR4 as well as CXCL12-mediated migration of leukemia T cells *in vitro*. Of note, they later engineered a CXCR4-nanobody-fragment crystallizable region (Fc) construct, which is similarly able to inhibit CXCR4 but also benefits from ADCC- and CDC-mediated tumor killing in leukemia cells [42]. Moreover, the same group engineered nanobodies directed against ACKR3, such as NB4, which were found to reduce angiogenesis and tumor growth of head and neck cancer cells both *in vitro* and *in vivo* [43].

Small molecules

The most widely investigated approach to interfere with chemokine ligand–receptor signaling is the use of small molecules that target chemokine receptors. Since these receptors canonically bind proteins rather than small molecules, their **orthosteric binding sites** are comparatively large, flexible, and highly exposed to the polarity of waters [44]. Thus, it is challenging to target them with small molecules, which generally require deep hydrophobic pockets [45]. Nevertheless, several orthosteric antagonists have been successfully developed, and many of them have been or are currently being tested as potential anticancer therapies in (pre)clinical trials [5]. These include antagonists for CCR2, CCR4, CCR5, CXCR2, and CXCR4 [5,46–49]. Of note, the CXCR4 antagonist Plerixafor has been clinically approved for the treatment of lymphoma and myeloma [50] and is currently being tested for use against other cancer types in Phase II clinical trials (NCT04177810 and NCT03746080). Furthermore, the CCR5 antagonist Maraviroc, which is clinically approved against HIV, may become repurposed for clinical use in metastatic colorectal cancer. Indeed, Phase I clinical trials recently investigated the use of Maraviroc as monotherapy [51], in combination with chemotherapy [52] or with immune checkpoint inhibitors (NCT03274804, NCT04721301). Similarly, other CCR5 antagonists are currently under clinical investigation [53].

Small molecules can also bind chemokine receptors at **allosteric binding sites**. Thus far, four chemokine receptors have been co-crystallized with an allosteric ligand: CCR2 [54], CCR7 [55], CCR9 [56], and CXCR2 [57]. Although several allosteric sites exist, these crystal structures all demonstrate the presence of the same intracellular binding pocket just above helix 8. As reviewed elsewhere [58], targeting chemokine receptors at allosteric sites may have several advantages over orthosteric binding. Most notably, by evading competition, it allows target binding and inhibition even in the presence of abundant chemokines (i.e., **insurmountable antagonism**). Examples of intracellular allosteric ligands include Reparixin and Navarixin, targeting both CXCR1 and CXCR2. Based on preliminary efficacy in a Phase I trial [59], Reparixin was investigated in combination with Paclitaxel as a treatment for metastatic triple-negative breast cancer in a Phase II clinical trial (NCT02370238). In addition, a recently terminated Phase II study in HER2⁺ breast cancer suggests some reduction in cancer stem cell numbers after treatment with Reparixin as monotherapy [60]. Navarixin is currently being investigated in Phase II clinical trials as part of a combination therapy with an anti-PD1 antibody in non-small cell lung cancer, castration-resistant prostate cancer, and microsatellite stable colorectal cancer (NCT03473925). Of note, the recent CCR7 crystal structure suggests that Navarixin also binds to this receptor [55]; as such, the anticancer effects of this drug might also be mediated by CCR7 [61]. Another example of an allosteric antagonist is Mavorixafor, which binds to an extracellular region of CXCR4 and is currently being tested in clinical trials for advanced renal cell carcinoma (NCT02667886), Waldenström's macroglobulinemia (NCT04274738), and advanced melanoma (NCT02823405). Preliminary results have been disclosed for the study in advanced melanoma, where Mavorixafor was found to be well tolerated, and to increase antitumor CD8⁺ T cell levels in the tumor tissue. Furthermore, the therapeutic effect of Mavorixafor was further enhanced when administered in combination with the anti-PD1 antibody Pembrolizumab [62]. Although only few allosteric ligands have been clinically tested yet, these examples demonstrate the viability of allosteric modulators as anticancer therapies.

Peptidic antagonists

Chemokine receptors can also be modulated by peptide drugs (different from modified chemokines). In the context of cancer, this avenue has mostly been investigated for CXCR4. Out of the several peptide CXCR4 antagonists developed to date [63], LY2510924 has advanced to Phase II, while Motixafortide and Balixafortide to Phase III clinical trials. The clinical use of Motixafortide as a co-therapy has been evaluated in over ten clinical trials spanning Phase I–III

and across several cancer types, mainly for stem cell mobilization in multiple myeloma [64] and metastatic pancreatic cancer [65]. Balixafortide is currently being tested in a Phase III clinical trial in HER2⁺ metastatic breast cancer (NCT03786094). Finally, the safety of LY2510924 has been demonstrated in several clinical trials, including Phase I and II studies in acute myeloid leukemia [66], advanced solid tumors [67], small-cell lung cancer [68], and advanced renal cell carcinoma [69]; however clinical efficacy has not been demonstrated yet. These cases suggest that peptide antagonists have good toxicity profiles and may provide a viable option for clinical treatment against cancer.

Pepducins are a relatively novel type of peptidic ligands that bind to the intracellular binding site of the target G protein-coupled receptor (GPCR), and therefore likely act as allosteric modulators [70]. In addition, pepducins have desirable pharmacokinetic properties with high bioavailability, and long serum half-lives [71]. As such, pepducins are a promising option in general for novel therapeutics. So far, only three chemokine receptors have been targeted with pepducins: CXCR1, CXCR2, and CXCR4, each in a preclinical setting. The X1/2pal-i3 pepducin, targeting the third intracellular loop of CXCR1 and CXCR2, effectively reduced tumor growth and angiogenesis in a mouse model of ovarian cancer [72]. In another study, X1/2pal-i3 administration was found to reduce CXCL8-induced tumor growth and expression of the antiapoptotic protein Bcl-2 in PTEN-deficient prostate cancer xenografts in mice [73]. In addition, the pepducins PZ-218 and PZ-210, which respectively target the first or third intracellular loop of CXCR4, were found to stimulate apoptosis in cancer cells from leukemia patients. Furthermore, these pepducins enhanced Rituximab-induced apoptosis in a Rituximab-resistant cell line, which could not be achieved with the orthosteric antagonist Plerixafor, and prolonged survival in a mouse lymphoma model [74]. Altogether, these results show promise for pepducins as novel anticancer agents.

Small interfering RNA

Finally, it is also possible to interfere with chemokine-receptor binding by impairing the expression of chemokine receptors with **small interfering ribonucleic acids (siRNAs)**. siRNAs can downregulate chemokine receptor expression directly on tumor cells to prevent oncogenic signaling, or on the TME to impair the recruitment of immunosuppressive cells, such as **tumor-associated macrophages (TAMs)**. The anticancer value of siRNAs targeting chemokine receptors is illustrated by a CXCR4-targeting siRNA, which significantly attenuated tumor growth and metastasis in an *in vivo* model of HER2⁺ breast cancer [75]. In addition, administration of a CCR2-targeting siRNA resulted in a decrease in CCR2 expression on monocytes, reduced TAM levels, suppressed tumor growth, and attenuated metastatic behavior in a 4T1 murine breast cancer model [76]. However, siRNAs are characterized by poor stability, limited intracellular uptake, and poor localization to target cells [75]. These caveats have been addressed by the coupling of the siRNAs to an anti-HER2 nanobody (ensuring localization to HER2⁺ breast cancer cells) [75] or by their encapsulation in a cationic polymeric nanoparticle (ensuring delivery to monocytes) [76].

Concluding remarks

With critical roles in tumor growth, metastasis, and TME formation, the chemokine system embodies a key therapeutic target against cancer [3]. Here, we outlined the potential for anticancer therapies that interfere with chemokine function at three critical stages: (i) chemokine availability; (ii) chemokine-GAG binding; and (iii) chemokine receptor binding. These different stages can be targeted by multiple strategies, including siRNA, biologics, peptides, small molecules, and glycomimetics (Figure 2 and Table 1), each of them exhibiting their own advantages and disadvantages (Table 2). Many of these strategies have progressed to clinical trials, and some are even on the market, emphasizing the clinical potential of targeting this system in cancer. Yet, many challenges remain to be addressed in future research in order to improve their likelihood

Outstanding questions

Which cancer (sub)types and stages will benefit from which specific chemokine intervention strategy?

How can antibodies be designed to only target free, non-GAG-bound chemokines?

What are the underlying principles that define specificity in chemokine-GAG interactions, and how can these be exploited to design therapies that selectively interfere with one chemokine-GAG pair?

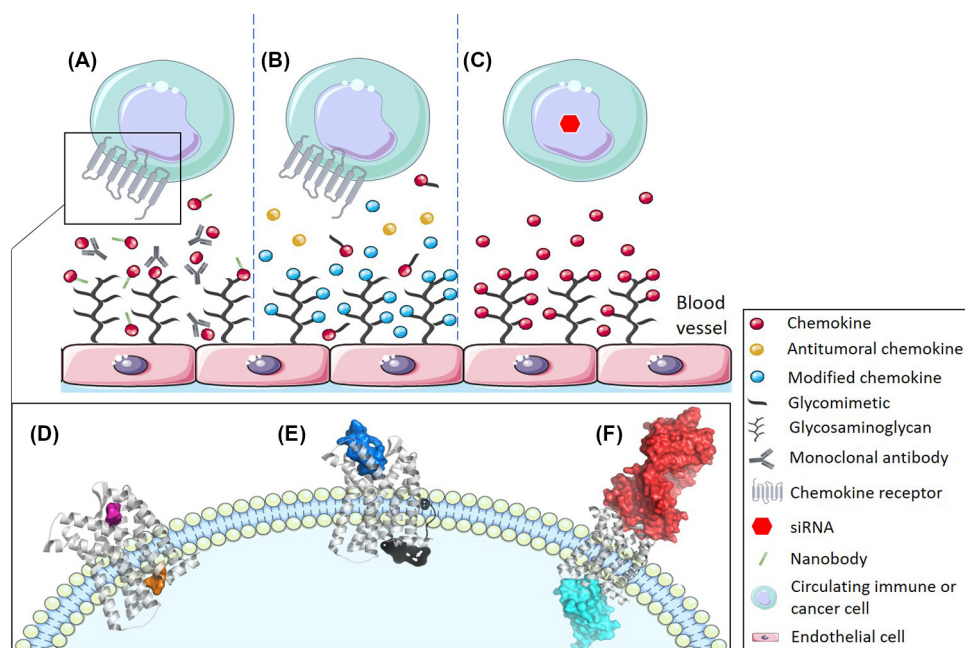
The expression of some chemokines and their receptors has been successfully repressed. Can this be achieved for other chemokines/receptors as well, and how can this be optimized to only occur at the tumor site?

Are allosteric binding pockets present in all chemokine receptors, and, if so, how can allosteric binding be optimized for one particular chemokine receptor over another?

Would it be clinically advantageous to simultaneously target the chemokine system on several levels in the signaling cascade? For example, can the therapeutic effect be enhanced by combining a GAG-mimetic and a chemokine receptor antagonist?

Can multitarget drugs be rationally designed for currently unexplored combinations of chemokine receptors? And do multitarget drugs always enhance the therapeutic response over targeting chemokine receptors with individual modulators?

How can biased signaling be rationally designed?



Trends in Pharmacological Sciences

Figure 2. Overview of the therapeutic avenues to modulate the chemokine system in cancer. (A) By neutralizing chemokines (red) with biologics (monoclonal antibodies or nanobodies), a concentration gradient does not develop and/or chemokines are precluded from interacting with their receptor. (B) Administering modified chemokines (blue) with enhanced GAG-binding and an inability to bind chemokine receptors prevents chemokine ligand-receptor binding. Alternatively, exogenous GAGs (glycomimetics) can bind to chemokines to prevent chemokine ligand-receptor binding, or delivery systems can localize antitumoral chemokines (yellow). (C) Chemokine receptor expression can be repressed with short interfering RNA (siRNA) to prevent chemokine signaling even if a concentration gradient develops. (D) Depicts orthosteric (magenta) and allosteric (orange) antagonism of the chemokine receptor (white) by small molecules. (E) Illustrates how chemokine receptors can be antagonized by peptides (blue) or pepducins (black). (F) Shows neutralization of a chemokine receptor by an antibody (red) or nanobody (cyan). This figure was created using elements from Servier medical art. Abbreviation: GAG, glycosaminoglycan.

of clinical success, including the need to increase our understanding of the chemokine system biology in cancer (see [Outstanding questions](#)).

Clinical success may also be enhanced by uncovering novel strategies. In this context, it is worth noting that, although not tested against cancer yet, intracellular allosteric ligands have also been synthesized for many other chemokine receptors than those discussed in the main text, including CCR1, CCR4, CCR5, and CX3CR1 [77–80]. Long-residence time antagonists might provide another strategy to inhibit chemokine receptors in an insurmountable manner, such as those described for CCR2 [81,82]. Furthermore, the first covalent intracellular antagonist for CCR2 has been recently reported [83], which may improve the therapeutic value via an insurmountable mechanism of action [84]. Another way to improve the clinical efficacy of chemokine receptor inhibitors is to exploit the emerging concept of **biased signaling**, which has been largely overlooked in drug discovery campaigns. In this regard, Hitchinson *et al.* [85] found that inhibition of β -arrestin is responsible for the development of tolerance to CXCR4 antagonist Plerixafor due to an increase of CXCR4 cell surface expression. By contrast, $G\alpha_i$ -biased CXCR4 antagonists SEN071 and X-4-2-6 did not affect CXCR4 cell surface expression and thereby evaded tolerance. Furthermore, recent studies have investigated the development of promiscuous chemokine-blocking antibodies [86] and chemokine-blocking **peptibodies** [87], which aim to

Table 2. Overview of the key (dis)advantages of the different approaches targeting the chemokine system in cancer

Type of treatment	Advantages	Disadvantages
Antibodies targeting chemokines	Long serum half-life Selectively interferes with a single chemokine, whereas targeting its receptor affects the signaling of all chemokines that bind the receptor Can be used as part of pharmacodelivery systems	Difficult and expensive to develop Cannot be administered orally Requires additional engineering to prevent targeting of GAG-bound chemokines Rebound effect after treatment, which may exacerbate the pathology
Antibodies targeting chemokine receptors	Long serum half-life High target specificity Can elicit antitumor immune responses (ADCC and CDC)	Challenging to generate antibodies against chemokine receptors Need to be careful not to target a chemokine receptor that is widely expressed on non-cancer cells
Nanobodies (targeting chemokines or chemokine receptors)	Can reach more epitopes than antibodies Cheaper and higher-yielding than production of antibodies Easier to administer (e.g., through oral administration) than antibodies	Short serum half-life and general instability may require coupling to other molecules
Small molecule chemokine neutralizing agents	Cheaper and quicker to produce than biologics Easier to administer than biologics	Only a few chemokine-targeting small molecules have been developed, resulting in little data for rational design approaches. Requires frequent administration compared to biologics
Modified chemokines	Can preclude the formation of a concentration gradient of tumorigenic chemokines	Requires engineering of the chemokine sequence May need to be fused with other agents (e.g., HSA) due to low serum half-life
Glycomimetics	Can preclude the formation of a concentration gradient of tumorigenic chemokines May attenuate angiogenesis by inhibiting soluble angiogenic factors, such as VEGF	Difficult to design GAG-mimetics that selectively target one chemokine over another May affect unexpected soluble factors other than chemokines
Small molecule chemokine receptor ligands	Easier to produce, purify, and administer than biologics Potential for biased signaling to prevent drug tolerance <u>Orthosteric ligands:</u> Ample data available for rational design of novel compounds <u>Allosteric ligands:</u> Insurmountable activity (i.e., effect even at the presence of abundant endogenous chemokines) Ceiling effect (i.e., maximum physiological effect, reducing toxicity risks) May be more selective than orthosteric ligands	The exact chemical features that determine biased signaling are not entirely understood, hindering the rational design of biased ligands More prone to off-target induced side-effects compared to biologics <u>Orthosteric ligands:</u> Compete with an abundance of endogenous chemokines Target a binding site that is large, flexible, and highly polar <u>Allosteric ligands:</u> Allosteric binding sites have only been confirmed by crystallization for a few chemokine receptors, hindering the rational design of allosteric ligands
Small interfering RNA	Can effectively repress the expression of chemokine receptors on TAMs or tumor cells	Require engineered delivery systems due to poor stability and intracellular uptake Difficult to target only the tumor site
Pepducins	Likely act as allosteric modulators, thus harboring the insurmountability and ceiling effect	Their therapeutic use has not been clinically tested Only three chemokine receptors have been successfully targeted with pepducins so far
Peptides	Have largely shown good toxicity profiles in clinical trials	Have mainly been studied for the CXCR4 receptor

target multiple chemokines simultaneously. Similarly, several **multitarget ligands** have been reported for chemokine receptors [88]. The therapeutic potential of such multitarget ligands can be illustrated by the large number of ongoing clinical trials with the dual CCR2/CCR5 antagonist BMS-813160 in various cancer types (NCT03496662, NCT03767582, NCT03184870, NCT04123379, and NCT02996110). Thus, although there is already a wealth of opportunities to target the chemokine system in cancer, new methods are poised to be added to the pharmacological toolbox. While future efforts are still required to optimize these individual methods, this anticipates great potential for the design of future anticancer drugs aimed at the chemokine system.

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