

REVIEW

Bispecific antibodies for viral immunotherapy

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ABSTRACT

Bispecific antibody engineering, in which binding specificities toward 2 distinct epitopes are combined into a single molecule, can greatly enhance immunotherapeutic properties of monoclonal antibodies. While the bispecific antibody approach has been applied widely to targets for indications such as cancer and inflammation, the development of such agents for viral immunotherapy is only now emerging. Here, we review recent advances in the development of bispecific antibodies for viral immunotherapy, highlighting promising *in vitro* and *in vivo* results.

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Introduction

Passive administration of monoclonal antibodies (mAbs) is a promising therapeutic platform for treatment of viral infections.¹ Currently, there is only one FDA-approved antiviral mAb, Synagis[®], for prophylaxis of respiratory syncytial virus (RSV) in premature infants.² However, a number of mAbs are in preclinical or clinical development for chronic viruses (e.g., HIV-1) as well as acute infections (e.g., influenza, Ebola virus, and West Nile virus).^{3–6} In other indications, mAbs are advantageous relative to other therapeutic platforms because they are well tolerated (favorable safety profile), are highly specific for their target with few off target effects, and have long serum half-life due to the Fc region.⁷ These advantages also extend to viral immunotherapy and, in cases where the antibody specificity is directed toward non-host (i.e. viral) epitopes, the safety profile may even be better than other mAbs that target human components.

The majority of antiviral mAb treatments consist of a single mAb that targets a single epitope on the virus surface (“monotherapy”); however, in several cases, a combination therapy consisting of 2 or more mAbs is being advanced. Most immunotherapeutic mAbs are targeted toward the glycoprotein of a particular virus, which is required for cell entry. In many viruses, the glycoproteins across species or strains within the same viral family contain a high degree of sequence variability, and thus a monotherapy may not be effective against all strains. Furthermore, monotherapies that target a single epitope are more susceptible to viral escape mutations and thus development of resistance. Cocktails of antibodies targeting multiple epitopes are capable of increasing potency as well as breadth, and potentially mitigate against escape mutations.^{4,8–11} While there may be clear therapeutic benefits for advancement

of a cocktail of mAbs rather than a monotherapy, the development of mAb cocktails also imposes additional regulatory and manufacturing hurdles.

Recent advancements in recombinant antibody engineering technologies have allowed the generation and clinical development of antibodies with enhanced function.^{12,13} Bispecific antibodies (Fig. 1), which can bind 2 or more separate and distinct epitopes, have been broadly applied across a number of disease indications. They have been engineered to bind 2 epitopes on the same cell, 2 epitopes on different cells, or different epitopes of the same antigen. In viral immunotherapy, bispecific mAb design can be used to lower the complexity of mAb cocktails, allowing simultaneous targeting of 2 or more distinct epitopes in a single entity. This approach potentially combines the therapeutic advantage of targeting multiple epitopes on the glycoprotein with the simplicity of developing and manufacturing a single molecule, as opposed to a mixture of 2 or more conventional mAbs. Furthermore, bispecific antibody engineering provides the opportunity to tailor multifunctional molecules to match the proposed mechanism of action, for example targeting both viral and host components simultaneously.^{13–15}

Here, we provide a brief overview of recent developments toward application of bispecific antibody design for viral immunotherapy. Much of the discussion focuses on formats that contain the Fc-region of immunoglobulin, which provides the potential for long pharmacokinetic half-life *in vivo* and for Fc-mediated antiviral mechanisms (Table 1).⁷ Virus-targeting bispecific antibodies can be grouped into 3 major subclasses (i) molecules with increased species or strain breadth that target several viral epitopes; (ii) antibodies that bind both virus and host epitopes; and (iii) antibodies that recruit the host cells immune system machinery (Fig. 1A). Currently, a variety of

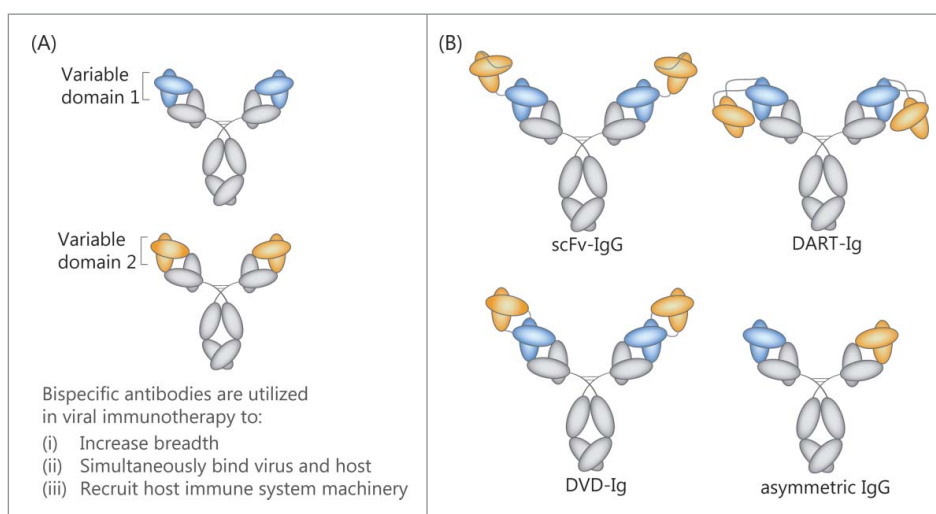


Figure 1. Schematic representation of different antibody formats. The combination of the variable regions of 2 monoclonal antibodies (A) can be achieved via numerous bispecific antibody formats (B). Here exemplified is a single chain variable fragment fusion to an IgG (scFv-IgG), a dual-affinity retargeting immunoglobulin fusion protein (DART-Ig), a dual-variable domain immunoglobulin (DVD-Ig), as well as an asymmetric IgG in which each arm engages a different epitope (e.g., Duobody or CrossMab).

bispecific antibody formats are applied in the development of antiviral multivalent antibodies (Fig. 1B). Methods that have been used for the multimerization of such antibodies include antibody conjugates, asymmetric IgG-like molecules, as well as single chain variable fragment (scFv) fusions to IgG molecules. These different formats and conjugation methods are each associated with particular advantages that may be tailored to the specific application.

Bispecific antibodies targeting 2 distinct viral epitopes

Bispecific antibodies that engage 2 distinct viral epitopes have potential to provide enhanced breadth and potency. As such, Wagner et al. reported a format where 2 full-sized anti-influenza A IgG antibodies with individual subgroup specificity were fused via their C-termini applying sortase transpeptidation and click chemistry.¹⁶ The C-C fusion of this covalently linked IgG antibody heterodimer does not require any mutations and it is therefore believed to not compromise antibody stability or Fc-receptor binding capacity. The antibody dimer retained the activity and stability of the 2 original antibodies against a wide range of influenza subtypes in a suitable mouse model. However, the C-C linkage could be immunogenic, a notion that requires further testing. Zanin et al. developed two neutralizing monoclonal antibodies against H5 influenza viruses of different clades, and combined them into a single bispecific Fc-domain fusion protein (referred to as Fc-dual-affinity retargeting molecule, or FcDART).¹⁷ The generated FcDART consist of 2 Fc-fusion protein chains that dimerize to form antibody derived binding sites. One polypeptide chain connects the light chain variable region (v_L) of the first antibody with the heavy chain variable region (v_H) of the second antibody via a Gly-Ser linker, and the second polypeptide contains the complementary variable regions. Assembly of a DART occurs via heterodimerization of the 2 chains and results in a single protein that is bispecific and tetravalent. The anti-influenza FcDART was shown to possess the broad-

spectrum activity and protective efficacy of both of its parental monoclonal antibodies in mouse infection models. Additionally, the FcDART demonstrated 100% protection in infected ferrets, which represents a more faithful model for influenza infection. Therefore, a therapy consisting of FcDART alone might be as efficacious as a cocktail containing both monoclonal antibodies against a broad spectrum of antigenically diverse influenza viruses.

Similarly, enhanced activity was also demonstrated for bispecific antibodies targeting the antigenically diverse envelope protein (Env) of HIV-1. Asokan et al. constructed asymmetric bispecific IgGs in which each arm binds a different broadly susceptible epitope on Env monovalently, applying the CrossMab format.¹⁸ The generated bispecific IgGs not only retained functional binding and neutralization by both arms, but also neutralized 94–97% of antigenically diverse viruses in a panel of 206 HIV-1 strains. The bispecific molecule with the highest neutralization activity was infused intravenously to rhesus macaques and demonstrated similar pharmacokinetic properties as the combination of its parental antibodies. To test the idea that increased parent affinity might enhance neutralizing activity, Mouquet et al. engineered scFv-Fc IgG-like molecules, also called immunoadhesins, that can bind HIV-1 bivalently by engaging 2 distinct subunits of Env.¹⁹ It was shown that heterotypic binding enhanced neutralization compared with the parental antibodies. By engineering an asymmetric bispecific IgG1 molecule that contains the hinge domain of an IgG3, Bounazos et al. recently reported additional success in the synergistic combination of different anti HIV-Env antibodies.²⁰ The length and flexibility of the IgG3 hinge domain allows for hetero-bivalent binding to 2 essential epitopes on 2 adjacent glycoproteins of the HIV-1 trimer spike, thereby increasing antibody avidity while facilitating spike inactivation. Compared to unmodified asymmetric bispecific molecules, the synergistic activity of molecules with modified hinge domains was shown to be superior in viral neutralization as well as in the protection of HIV-1 infected humanized mice.

Table 1. Fc-region containing bispecific antibody formats that have been generated for viral immunotherapy.

Virus	Targets	Format	Results	Reference
1) Targeting 2 distinct viral epitopes				
Influenza	HA protein of different strains	Full-sized IgG antibodies linked via C -termini	Retained activity and stability of original antibodies against various influenza subtypes	[16]
	Head of H5 HA protein	FcDART	Retained activity and efficacy of original antibodies in mice and ferret models	[17]
HIV-1	Various neutralizing epitopes on Env	Asymmetric IgG (CrossMAb)	Retained binding activity of original antibodies; enhanced neutralization of HIV-1 strains; best neutralizer retained pharmacokinetic properties of original antibodies	[18]
	Two subunits of Env	Immunoadhesins (scFv-Fc)	Retained binding activity of original antibodies; enhanced neutralization compared to original antibodies	[19]
	Different epitopes on adjacent GPs	Asymmetric IgG1 molecule with IgG3 hinge domain	Enhanced neutralization and protection in infected humanized mice compared to original antibodies and bispecific antibody without IgG3 hinge domain	[20]
Hepatitis B	Various epitopes on surface antigen	DVD	Enhanced neutralization and inhibition of hepatitis B surface-antigen secretion compared to original antibodies	[21]
Dengue	DII and DIII of surface protein	DVD with mutated Fc-domain	Retained binding activity of original antibodies; enhanced neutralization and protection in mouse model; eliminated ADE of infection <i>in vitro</i>	[22]
	DII and DIII of surface protein	DART-Ig with mutated Fc-domain	Retained neutralizing activity and therapeutic activity in mice as original antibodies; enhanced avidity for isolated recombinant DENV surface protein	[23]
Ebola	SUDV and EBOV GP	scFvs of EBOV antibody fused to SUDV-IgG	Retained neutralizing activity of original antibodies; post-exposure protection of mice from both viruses	[24]
2) Targeting viral and host epitopes				
HIV-1	Env and CD4	scFv of anti-Env antibodies fused to anti-CD4 IgG	Enhanced neutralizing activity and potency compared to original antibodies; neutralization of strains resistant to original antibodies	[14]
	CD4 and CD4 induced epitope on Env	Single-domain antibody (anti-CD4i site) fused to CD4 IgG	Enhanced neutralizing activity and potency compared to original antibodies; neutralization of strains resistant to original antibodies	[25]
	Various neutralizing epitopes on Env and CD4 or CCR5	Asymmetric IgG (CrossMAb)	Enhanced neutralizing activity compared to original antibodies; reduction of viral load in infected humanized mice	[26]
Ebola	Receptor binding site of GP and NPC1	DVD	Neutralization of all known ebolaviruses and post exposure protection against multiple filoviruses in mouse models.	[27]
3) Recruiting host cell machinery				
Dengue	Primate CR1 and DENV GP	Thioester Cross-linked IgGs	Facilitated specific and rapid binding of DENV to monkey and human erythrocytes; clearance of DENV viremia in cynomolgus macaques	[30]
Marburg	Primate CR1 and Marburg GP	Thioester Cross-linked IgGs	Facilitated specific and rapid binding of MARV-GP to monkey and human erythrocytes	[31]
CMV	CD3 and CMV	Thioester Cross-linked IgGs	Redirected specific cytotoxicity to CMV infected cells	[33]
HIV	CD3 and CD4 induced epitope on Env	scFV of anti-CD3 linked to light chain of anti-Env by GS linker	Increased CD8 ⁺ T-cell activation and lysis of infected cells; reduction of latently infected CD4 ⁺ T cells <i>ex vivo</i> ; <i>in vivo</i> safety in ART-treated SHIV-infected macaques	[36]

Guo and colleagues engineered a dual variable domain (DVD) bispecific antibody from 2 distinct monoclonal antibodies against hepatitis B surface antigen (HBsAg) that synergistically neutralized hepatitis B virus (HBV) infection.²¹ In DVDs the v_H and v_L of one antibody are genetically fused to the N-terminus of the heavy and light chain of a second antibody via a short peptide linker, respectively, thereby retaining the Y-shape of an IgG molecule (Fig. 1B). The resulting bispecific molecule was shown to have superior HBV-neutralizing activity in comparison to the combination of both parental monoclonal antibodies, possibly through steric hindrance or induction of HBsAg conformational changes. The same group recently described another DVD bispecific antibody, that neutralizes all 4 closely related but serologically distinct dengue viruses (DENV-1-4).²² The fused variable domains of this molecule bind the domain of the DENV surface protein E, that is responsible for virus attachment (DIII), while the variable domains of the IgG scaffold bind to the E protein

domain DII, which contains the fusion loop that inserts into the host cell during viral entry. The DVD molecule was confirmed to be more effective at protecting against DENV than each of its parental mAbs individually as well as the combination of the 2 in a suckling mouse model. Additionally, the mutation of 9 amino acids within the Fc domain eliminated antibody dependent enhancement (ADE) of infection in myeloid cells expressing Fc- γ receptors (Fc γ R). Earlier, Brien et al. already reported an IgG fusion DART protein (DART-Ig; Fig. 1B) that was also engineered to lack Fc γ R binding capacity and to bind 2 spatially distinct and cross-reactive epitopes on DII and DIII on the surface of the DENV virion.²³ The generated molecule possessed neutralizing activity *in vitro* and therapeutic activity in a mouse ADE disease model. An observed improved avidity of the tetravalent DART-Ig for DENV E protein did not translate into greater inhibitory activity *in vitro* and *in vivo*, possibly due to the quasi-icosahedral structure of the DENV virion, precluding binding

geometries necessary for simultaneous recognition of adjacent epitopes. In 2015, a set of bispecific antibodies was developed in which scFvs of an Ebola virus (Zaire) specific antibody were fused to the N- and C-termini of a Sudan virus-specific antibody (scFv-Ig; Fig. 1B). These two species are the most divergent within the ebolavirus family but have collectively accounted for 95% of Ebola-related deaths from 1976–2014. The bispecific molecules exhibited potent neutralization, and present the first example of antibodies to confer a high degree (in one case 100%) post-exposure protection of mice from 2 distinct Ebola virus species.²⁴

Bispecific antibodies that interact with host and viral epitopes

The approach of constructing bispecific antiviral antibodies that simultaneously bind to viral and host epitopes has been mainly utilized in HIV studies to target CD4 receptor presenting T-cells. By binding to CD4, these bispecifics are able to increase its local concentration at the point of viral entry and prevent HIV from infecting CD4⁺ T cells. Pace et al., developed a bispecific antibody where scFvs of broadly neutralizing anti-Env antibodies were fused to the N-terminus of a humanized anti-CD4 antibody via a Gly-Ser linker.¹⁴ The same research group engineered another bispecific antibody where 2 copies of a single domain antibody that binds to the CD4-induced site on Env is fused to the C-terminus of the heavy chain of the same anti-CD4⁺ antibody by a Gly-Ser linker.²⁵ Both bispecific molecules showed improved antiviral activity and were able to neutralize viruses that exhibit resistance to parental antibodies *in vitro*. Aiming to engineer synergistic bispecific antibodies that preserve the architecture of a normal IgG, Ho and colleagues developed asymmetric IgGs (Fig. 1B) with the Fab from either an anti-CD4 or anti-CCR5 antibody applying the CrossMab technology.²⁶ Two of these molecules showed increased neutralization in a panel of HIV-1 isolates, and one antibody significantly reduced the viral load in HIV-infected humanized mice.

Unlike HIV, which binds to the surface exposed CD4 receptor to facilitate host-cell attachment and entry, ebolaviruses engage the intracellular late endosome residing receptor Niemann-Pick C1 (NPC1) during host cell entry and infection. This engagement is absolutely required for host cell infection and the receptor-binding site (RBS) on the filovirus glycoprotein is shielded prior to physical sequestration in late endosomes. To target this interaction, recently a novel “Trojan Horse” bispecific antibody strategy was described, in which the viable domains of NPC-1 or GP-RBS specific antibodies are fused to an antibody that binds to a conserved surface exposed GP-epitope, applying the DVD technology.²⁷ It was shown that the bispecific molecules, but not their parental monoclonal antibodies, neutralized all known ebolaviruses by utilizing the viral particle themselves for endosomal delivery, and conferred post exposure protection against multiple filoviruses *in vivo*.

Bispecific antibodies recruiting the host cell machinery

Bispecific molecules can also be utilized to recruit host cell immune system machinery. Taylor et al., developed heteropolymers (HPs) to clear various prototype pathogens by covalently

crosslinking a monoclonal antibody specific for primate E complement receptor 1 with another monoclonal antibody specific to a target antigen with a nonreducible thioether bond.^{28,29} Potential for viral immunotherapy was shown in HPs that bound to the Dengue glycoprotein and inactivated Marburg virus.^{30,31} The HPs demonstrated rapid and specific binding of their respective pathogens to monkey and human erythrocytes *in vitro*. Furthermore, the dengue-specific HP demonstrated clearance of dengue passive viremia when administered to previously challenged cynomolgus macaques.

There has been much recent interest in engineering bispecific molecules that are able to activate and direct cytotoxic T-cells. For example, bispecific T-cell engagers (BiTEs) contain both an antigen-specific arm and an anti-CD3 arm, which activates and redirects cytotoxic CD8⁺ T-cells to antigen-specific cells.³² Given the success of T-cell engaging molecules in cancer immunotherapy, there is much potential for adoption of similar strategies to eliminate the viral particles themselves, or infected cells. For example, an anti-cytomegalovirus (CMV) bispecific molecule possessing an anti-CD3 arm has been reported.³³ To achieve bispecificity, an anti-CD3 monoclonal antibody was chemically heteroconjugated with a CMV specific human IgG via a nonreducible thioether bond. The bispecific antibody was able to redirect specific cytotoxicity to CMV-infected cells and induce cell lysis of target cells at an effector to target ratio as low as 1:1. Similar approaches have been applied in HIV-1, where the CD3-binding moiety serves a 2-fold purpose in directing cytotoxic CD8⁺ T-cells and activating latently infected cells. In what has been called a “kick and kill” strategy, the anti-CD3 arm promotes viral production by binding to latently infected CD4⁺ T-cells and stimulating Env gene expression. The antibody then crosslinks CD3 on CD8⁺ T-cells to bring the cytotoxic cell in close proximity to recently activated CD4⁺ T-cells with Env on the surface to induce cell lysis.³⁴ Interest in the “kick and kill” strategy for HIV-1 stems from the persistence of latently infected cells in HIV-1 infected individuals treated with antiretroviral therapy (ART). While ART can significantly reduce the viral load in an infected individual, it is unable to fully eradicate HIV-1 from the system due to the virus establishing a stable latent infection in resting CD4⁺ T-cells with minimal expression of HIV genes.³⁵ Fortunately, 2 bispecific anti-HIV antibodies have shown promise in the lysis of these reservoirs and potential use in ART-treated patients. Additionally, an immunomodulatory protein consisting of the scFv of an anti-human CD3 monoclonal antibody that is linked to the light chain of a HIV-1 broadly neutralizing antibody by a Gly-Ser linker showed similar results in CD8⁺ T-cell activation and *in vivo* safety in ART-treated SHIV-infected macaques.³⁶

Multispecific antibody derived formats lacking an Fc-region

Success has also been found in multifunctional anti-viral antibody-mimetics. Genetic fusions of recombinant single-domain antibody fragments derived from camelid heavy chain antibodies (VHHs) have gained attention due to their high stability and easy production through prokaryotic expression. For example, 2 VHHs that bind to different epitopes on the

rotavirus capsid were displayed on the surface of *Lactobacillus paracasei* as linear bivalent proteins that are fused without linkers.³⁷ These were shown to be superior at reducing the rate of diarrhea when used for prophylactic and therapeutic intervention in a mouse model of rotavirus infection. Additionally, Hultberg et al. reported multimeric constructs developed by fusing different llama heavy chain antibody fragments with Gly-Ser linkers of varying length.³⁸ Multivalent molecules were constructed with VHHs against the trimeric envelope proteins of RSV, rabies virus and H5N1 influenza. Despite functional differences in the protein among the viruses, multimerization of the VHHs exhibited increased potency and cross-protection capacity against different viral strains for all 3 viruses. Recently, a DART engineered with the arm of a CD3-specific monoclonal antibody and the arm of one of 2 anti-Env HIV antibodies showed increased CD8⁺ T-cell activation and lysis with lymphocytes derived from patients on ART.³⁹ Additionally, a bispecific antibody-based molecule has been reported in which 2 Fabs of broadly neutralizing anti-HIV antibodies were conjugated to various lengths of double-stranded DNA. The architecture of the Env spike trimer prevents bivalent binding by the Fabs of IgG and reduces neutralization potency. The flexibility and length in the DNA linkers overcome this barrier and allow for bivalent binding of the Fabs to the Env spike trimer via intra-spike crosslinking. When the molecules were engineered with the Fabs of 2 different broadly neutralizing anti-HIV antibodies, the neutralization potency increased 100-fold.

Potential drawbacks of bispecific antibodies

From a developmental and regulatory perspective, combining several antibodies into a single therapeutic makes development less complex and more cost-effective, since production and quality control as well as preclinical and clinical testing will be reduced to a single molecule. However, bispecific antibodies are also associated with some disadvantages relative to canonical mAbs, and these limitations should be considered depending on the desired outcome. Asymmetric IgGs (e.g. DuoBodies, CrossMab) bind monovalently to their targets, which might have consequences for activity, especially in the context of viruses that have high glycoprotein spike density per virion and thus may be more potently neutralized by mAbs that engage epitopes bivalently to cross-link subunits or spikes.^{40,41} The production of asymmetric IgGs also requires either a reduction step to convert homodimers into heterodimers, or coexpression of 2 antibody fragments. This additional requirement might hamper high scale production of these molecules. Some bispecific antibody formats might possess reduced physicochemical stability or be prone to aggregation, for example, when lacking a Fc domain, if a constant domain mutation is required to enable IgG heterodimerization, or if long flexible polypeptide linker regions are included in the design. This can comprise stability and may affect binding to Fc receptors.^{42,43} So far, little attention has been given to the immunogenicity and pharmacokinetics of multispecific antibodies within the specific context of viral immunotherapy. However, recent studies on bispecific antibodies generated for different purposes, indicate that the orientation as well as the nature of the fusion partner can affect

the pharmacokinetics of immunoconjugates of IgGs, as it may result in conformational change or misfolding in certain combinations.^{44,45} However, the *in vivo* half-life for any particular antibody (engineered or natural) is unpredictable and requires empirical case-by case assessments. In general, enhanced pharmacokinetics of molecules containing an IgG scaffold can be attributed to 2 major factors; (i) their larger size (>150 kDa) precludes renal clearance, which is responsible for the rapid elimination of smaller constructs (60 kDa), such as scFvs; and (ii) their dynamic binding to the neonatal Fc receptor extends serum half-lives.⁴⁵ An intravenously administered bispecific antibody targeting different epitopes on the HIV-1 envelope was shown to have a similar serum half-life (ca. 10 days) to the parental IgGs in rhesus macaques and persisted in circulation for 4 weeks while retaining neutralization activity in the serum.¹⁸ Like the parental IgGs, this antibody contained a mutated CH3 domain to improve antibody persistence. Development for clinical application might require further engineering of existing bispecific formats. As such, hinge stabilized or aglycosylated IgG4 scaffolds can be applied to decrease heterogeneity.⁴⁶ Furthermore, anti- and pro-inflammatory properties can be regulated through the alteration of glycosylation, and the serum half-life can be increased by Fc engineering.⁴⁶ Other aspects that remain to be carefully addressed when preparing multispecific antibodies for clinical development are formulation and dosing, especially if different amounts of individual components are required in the respective canonical antibody cocktails.

Conclusion

To date, a vast repertoire of bispecific antibody formats is available, which has been primarily applied in the therapy of malignancies and inflammatory conditions. The studies we outlined here highlight that bispecific antibodies offer numerous functionalities and targeting mechanisms that can be exploited for viral immunotherapy. Despite potential limitations, bispecific antibody engineering can be utilized to enhance the effectiveness of antiviral immunotherapies. As individual epitopes may not provide sufficient therapeutic benefit when targeted alone, significant therapeutic benefits of these targets can be gained from additive and/or synergy effects of bispecific molecules. Moreover, bispecific antibody engineering may open up new therapeutic venues when utilizing one of the 2 specificities for the delivery of the second specificity to its target site. Thus, there is much opportunity for the application of such methods to other viruses and to develop next-generation therapeutics.

Disclosure of potential conflicts of interest

JRL and EKN are named coinventors on patents encompassing bispecific antibodies described in references 24 and 27.

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