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*CORRESPONDENCE Morgane Bomsel Morgane.bomsel@inserm.fr

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Antibody Fc-chimerism and effector functions: When IgG takes advantage of IgA

Andréa Cottignies-Calamarte ^{1,2}, Daniela Tudor ^{1,2} and Morgane Bomsel ^{1,2*}

¹Laboratory of Mucosal Entry of HIV-1 and Mucosal Immunity, Department of Infection, Immunity and Inflammation, Cochin Institute, Paris, France, ²Université Paris Cité, Institut Cochin, Institut National de la Santé et de la Recherche Médicale (INSERM), Centre National de la Recherche Scientifique (CNRS), Paris, France

Recent advances in the development of therapeutic antibodies (Abs) have greatly improved the treatment of otherwise drug-resistant cancers and autoimmune diseases. Antibody activities are mediated by both their Fab and the Fc. However, therapeutic Abs base their protective mechanisms on Fc-mediated effector functions resulting in the activation of innate immune cells by FcRs. Therefore, Fc-bioengineering has been widely used to maximise the efficacy and convenience of therapeutic antibodies. Today, IgG remains the only commercially available therapeutic Abs, at the expense of other isotypes. Indeed, production, sampling, analysis and related in vivo studies are easier to perform with IgG than with IgA due to well-developed tools. However, interest in IgA is growing, despite a shorter serum half-life and a more difficult sampling and purification methods than IgG. Indeed, the paradigm that the effector functions of IgG surpass those of IgA has been experimentally challenged. Firstly, IgA has been shown to bind to its Fc receptor (FcR) on effector cells of innate immunity with greater efficiency than IgG, resulting in more robust IgA-mediated effector functions in vitro and better survival of treated animals. In addition, the two isotypes have been shown to act synergistically. From these results, new therapeutic formats of Abs are currently emerging, in particular chimeric Abs containing two tandemly expressed Fc, one from IgG (Fcy) and one from IgA (Fc α). By binding both Fc γ R and FcaR on effector cells, these new chimeras showed improved effector functions in vitro that were translated in vivo. Furthermore, these chimeras retain an IgG-like half-life in the blood, which could improve Ab-based therapies, including in AIDS. This review provides the rationale, based on the biology of IgA and IgG, for the development of Fc γ and Fc α chimeras as therapeutic Abs, offering promising opportunities for HIV-1 infected patients. We will first describe the main features of the IgA- and IgG-specific Fc-mediated signalling pathways and their respective functional differences. We will then summarise the very promising results on Fc γ and Fc α containing chimeras in cancer treatment. Finally, we will discuss the impact of Fc α -Fc γ chimerism in prevention/treatment strategies against infectious diseases such as HIV-1.

KEYWORDS

antibody-engineering, the rapeutic antibodies, Fc-mediated effector functions, chimeric antibody, Fc γR , Fc $\alpha R/CD89$, IgA

Highlights

- Antibody-based therapies take advantage of the Fc region of antibodies to activate a broad spectrum of Fc-mediated effector functions, thereby linking adaptive and innate immunity.
- IgA conveys stronger FcR signalling and effector functions than IgG, but evaluation of IgA therapeutics is neglected.
- The presence of both $Fc\alpha$ and $Fc\gamma$ gives the resulting chimeric antibodies the ability to bind to a greater number of FcR molecules than the parental antibodies, resulting in increased killing of target cancer cells.
- Chimeric Abs containing Fcα-Fcγ could be effective as antiinfective agents, particularly against HIV-1.

1 Introduction

Antibodies (Abs) used as therapeutic agents have many advantages such as specific targeting provided by their Antigen Binding Fragment (Fab), good bioavailability, and the ability to engage immune effector cells using their crystallizable region (Fc). In humans, IgG and IgA predominate over the other three Ab isotypes. IgG accounts for more than 80% of circulating Abs (IgG1 being the most abundant (1)), and with a proportion of 15%, IgA (mainly IgA1) is the second most abundant circulating isotype (2). In contrast, in the mucosa, more than 90% of mucosal antibodies are IgA, with the ratio of IgA1 to IgA2 subtypes varying according to mucosal site (2).

IgA and IgG share common features, including general shape and functional domains (Figure 1). Both isotypes bind to the antigen using their Fab, which is linked via a hinge region to the Fc. The latter then interacts with the corresponding Fc receptor (FcR) on the surface of the effector cell. However, IgG and IgA differ in their glycosylation patterns, with IgG showing predominantly N-glycosylations while IgA shows Nand O-glycosylations, even in the hinge region (3) (Figures 1A, C). In addition, a tail completes the C-terminus of the IgA heavy chain (HC), which contributes to the dimerisation/multimerisation processes of mucosal IgA through binding to the J-chain (2) (Figure 1B). Despite the compartmentalisation of Ab isotypes that has evolved to counteract tissue-specific pathologies, the current development of therapeutic Abs focuses primarily on IgG, particularly IgG1. This predominance is the result of various technical factors: (i) IgG is the most abundant isotype in blood, easier to produce in large quantities and, therefore, widely studied, unlike the other isotypes; (ii) conversely, IgA, the second most abundant isotype in serum and the most abundant in mucosal sites, is more difficult to sample, produce and purify; (iii) IgG1 and IgG3 are the IgG subtypes that best induce Fc-dependent effector functions; (iv) IgG1 has a longer half-life in blood than IgG3 and IgA; and (v) IgG3, due to a long hinge, like IgA1 (Figures 1A, C), and a complex glycosylation profile, is produced at a higher cost than IgG1 (4-7). Thus, IgG1 has become the preferred isotype for the design of therapeutic antibodies. This choice should be reconsidered because of isotypic differences in Fc-mediated functions that are particularly significant for IgA.

To determine how antibody functions can be improved to meet therapeutic needs, a thorough understanding of antibody biology is required. Therefore, a variety of Fc-mediated functions of IgG and IgA have been extensively characterised over the past 20 years, revealing substantial IgA efficiencies that could be applied to the design of therapeutic mAbs (8). Abs link adaptive and innate immunity by interacting with their corresponding FcR on the surface of effector cells downstream of antigen targeting, resulting in the induction of a wide range of immune responses (8-10). In cancer treatment, one of the main therapeutic Ab mechanisms is initiated by opsonisation of cancer cells, leading to their elimination by innate immune effector cells (11). However, despite the superior ability of IgA-FcoRI/CD89 compared to IgG/FcyR to trigger effector functions such as antibody-dependent cellular cytotoxicity (ADCC) or antibody-dependent cellular phagocytosis (ADCP), the use of IgA as a therapeutic agent remains limited (12-16). To take advantage of the signalling cascades and subsequent effector functions mediated by both isotypes, chimeras containing two Fc domains, namely Fcy and Fc α in tandem, have emerged as new formats of therapeutic Abs (17, 18). This approach is also relevant for mucosal pathogens, such as Human Immunodeficiency Virus type 1 (HIV-1). Thus, prevention of viral entry into mucosa by HIV-1-specific IgA (19, 20) as well as neutralisation of HIV-1 in the circulation by HIV-1-specific IgG correlate with protection against HIV-1 infection in vivo (19-22).

In this review, we detail the differences between IgG- and IgAdependent intracellular FcR signalling pathways and Fc-mediated effector functions, and describe how new Ab-based therapeutic strategies can benefit from these properties, especially in the field of Aqcuiered Immunodeficiency Syndrome (AIDS).

2 The basis of FcR-mediated functions

The Fc adopts two dominant conformational states, namely an open and relaxed 'U' shape or a closed and compact 'V' shape, mainly regulated by N-glycosylation. Each conformation of Fc γ engages a distinct class of FcRs, namely type I and type II FcRs, and in turn activates various effector functions. The open 'U' form binds preferentially to type I FcRs while the closed 'V' form binds preferentially to type II FcRs (1, 9, 23). There are also other FcRs for IgG, such as TRIM21 and the neonatal FcR (FcRn), for which no preferential conformation of Fc γ is required for binding (9, 24, 25). In contrast, the role of Fc α conformation in IgA binding of FcRs has been less described (8, 26–29), although Fc α RI/CD89, the major Fc α R, appears to be a type I FcR.

An additional parameter that affects Fc-mediated functions is the oligomerisation state of Ab. Compared to IgG, which is exclusively expressed as a monomeric antibody, IgA is predominantly monomeric in serum and polymeric (mostly dimeric) in mucosal tissues. Dimeric IgA becomes secretory (sIgA) when it is secreted to mucosal surfaces after acquisition by proteolytic cleavage of the secretory component (SC), the extracellular part of the polymeric Ig receptor (pIgR) which transcytoses IgA and allows it access to external fluids (8). Furthermore, the SC protects IgA from degradation (Figures 1A, B).



represent the variable regions of the heavy and light chains (HC and LC respectively), together forming the antigen-specific paratope. The J chain (grey) is present in dimeric and secretory IgA (dIgA and sIgA, respectively). SIgA contains the secretory component (green), which corresponds to the extracellular region of pIgR acquired by proteolytic cleavage, after pIgR had translocated dimeric IgA to mucosal surfaces **(B)**. Inverted triangles and yellow circles represent the N- and O-glycosylation sites, respectively (A, B and C). Hinge length varies between human IgA and IgG subtypes: **(A, B)** IgA1>IgA2>IgG3>IgG3>IgG3>IgG4.

2.1 Understanding the Ab-FcR interaction

IgG can bind to six different type I IgG-specific FcRs, namely Fc γ RI/CD64, Fc γ RII/CD32a, b and c, and Fc γ RIII/CD16a and b (Figure 2A). These type I FcRs are either pro-inflammatory such as Fc γ RI/CD64, Fc γ RIIa/CD32a, Fc γ RIIC/CD32c, Fc γ RIIIa/CD16a and Fc γ RIIIb/CD16b, or anti-inflammatory such as Fc γ RIIb/CD32b (9). The balance between inflammatory and anti-inflammatory properties relies on induced cell signalling, and the subsequent release of specific sets of cytokines, as discussed below. IgG interacts with the type I FcR *via* the upper part of the CH2

domain of Ab located next to the hinge region that activates downstream signals *via* an immunoreceptor tyrosine-based activation or inhibition motif (ITAM and ITIM, respectively) (30). ITAM is contained either in the cytoplasmic domain of the receptor namely the Fc γ RIIa/CD32a and Fc γ RIIc/CD32c, or in the γ -chain, the FcR common adaptor associated with the Fc γ RI/CD64 and Fc γ RIIIa/CD16a, resulting in a cascade of activation signals (Figure 2A). Activation of ITAM engages a pro-inflammatory signalling pathway in effector cells while the ITIM domain activates a downstream anti-inflammatory signalling cascade, described below. Fc γ RIIb/CD32b is the only inhibitory Fc γ R that



FIGURE 2

Human IgG and IgA Fc receptors (FcRs) and their mode of action. (A) Major IgG and IgA FcRs: IgG and IgA FcRs are represented at the cell membrane, associated where indicated with the adaptor protein FcR γ . The respective affinities for IgG from + to +++ are indicated. For FcRn and Fc α RI/CD89, ++/+ ++ and +/+++ refer to the affinity level of antigen-free/antigen-bound antibodies. The activation (ITAM) and inhibition (ITIM) motifs of tyrosine-based immunoreceptors are shown in red and light pink, respectively. The neonatal Fc receptor (FcRn), a non-classical MHC class I only capable of interacting with IgG, is shown associated with its β2-microtubulin chain (red). (B) ITAM-induced activation upon FcyR binding by IgG-IC: IgG binding in the immune complex (IgG-IC) (antigen in light green) to FcyRI/CD64 or FcyRIIa/CD32a results in FcR clustering that triggers the Src kinase Fyn. In turn, Fyn phosphorylates the two tyrosines (white circle) of ITAM contained either in the cytosolic tail of the FcR or in the associated adaptor protein (FcRy chain). These phosphorylations generate docking sites for Syk kinase family that induces a PI3K- and PLCγ-mediated intracellular increase in IP3 and DAG, followed by MAPK activation and an increase in intracellular Ca2+. Subsequently, effector functions such as ADCC and ADCP as well as cytokine secretion and NETosis are triggered. (C) ITAM-induced activation upon FcαRI/CD89 binding by IgA-IC: IgA-IC cross-links FcαRI/CD89 and the associated FcR_γ chain, initiating ITAM phosphorylation by Fyn and the subsequent signalling pathway, similar to that triggered above for IgG-IC. In turn, effector immune cells are activated as well as a pro-inflammatory response, resulting in pathogen elimination by ADCC and ADCP but also cytokine secretion, and NETosis. (D) ITIM/ITAMi signalling: Binding of IgG-IC to FcyRIIb/CD32b carrying ITIM induces monophosphorylation of ITIM. Binding of uncomplexed IgA to FcxRI/CD89 triggers suboptimal phosphorylation of ITAM on the intracellular adaptor FcRy, termed ITAM inhibitor (ITAMi) and schematised in a red/pink hatched rectangle. Then, the ITAMi and ITIM pathways converged: Monophosphorylation of ITIM or ITAMi by Lyn, another receptor-associated Src kinase, promotes the recruitment of SH2-phosphatases (SHP). In turn, SHP opposes Syk activity, thereby aborting the immune response. These two different negative regulatory mechanisms, ITIM and ITAMi, help prevent uncontrolled inflammation and maintain immune homeostasis

carries the ITIM motif in its cytoplasmic domain (Figure 2A), thus controlling many aspects of the inflammatory response. Among IgG isotypes, IgG3 has the highest affinity for $Fc\gamma RI/CD64$, followed by IgG1 and IgG4, while IgG2 is unable to bind to $Fc\gamma RI/CD64$ (10).

IgA binds to two different type I FcRs: $Fc\alpha RI/CD89$ and $Fc\alpha/\mu RI$. IgA binds to $Fc\alpha RI/CD89$ *via* the CH2 and CH3 domains and, like $Fc\gamma RI/CD64$, $Fc\alpha RI/CD89$ associates with the FcR γ adapter chain to induce downstream signalling (8) (Figure 2A). The affinity of $Fc\alpha RI/$ CD89 for IgA varies according to its ability to form an immune complex (IC) with the antigen, whether free or on the surface of opsonised cells or microbes. The affinity of Fc α RI/CD89 for IgA is low when IgA is free of antigen but becomes high when IgA is in IC with its antigen. As will be discussed below, Fc α RI/CD89 can propagate activating but also inhibitory signalling pathways, called inhibitory ITAMs (ITAMi), through the same ITAM domain of the FcR γ chain when activated by antigen-free IgA. Monomeric and dimeric IgA engage Fc α RI/CD89 in an IgA:receptor molecular ratio of 1:2, but the engagement of secretory IgA is limited by the presence of the SC that partially masks the Fc α RI/CD89 interaction domain (27, 31–33). Fc α /µRI is much less well characterized than Fc α RI/CD89. It is expressed by marginal zone B cells, follicular DCs and tonsil cells with expected functional roles (34), but not by circulating lymphocytes.

Type II FcRs are C-type lectins, such as DC-SIGN or Dectin-1, which bind to carbohydrate biomolecules, such as Abs. In contrast to type I FcRs, type II FcRs preferentially interact with the closed 'V' conformation of the Fc domains of IgG but also of IgA. Ab binding triggers anti-inflammatory responses (35, 36), although these results remain debated (37–39).

As summarised in Table 1, FcRs are predominantly expressed on innate immune cells which are therefore considered to be the main effector cells. The expression of FcRs on immune cells throughout the human body is well documented, but comparative expression of Fc γ R and Fc α R between blood cells and mucosal tissues is lacking (40–42). FcR

expression varies qualitatively and quantitatively between blood DC subsets (43), but also between blood and tissue DCs and different mucosal DCs. Genital and rectal mucosal DCs express FcyRII/CD32, low levels of Fc α RI/CD89 and FcyRI/CD64 but lack Fc γ RIII/CD16. Blood DCs have a similar pattern of FcR expression to tissue DCs, although the level of Fc γ RII/CD32 (a, b and c) is higher and that of Fc γ RIII/CD16 is just detectable in blood DCs (40). Blood and mucosal NK cells express only Fc γ RIII/CD16 but not Fc α RI/CD89 in men and at low frequency in the female genital tract correlating with a sexual polymorphism (40).

All IgA and IgG FcRs are expressed by mucosal macrophages in genital and rectal tissues, whereas in the gastrointestinal tract, macrophages are almost devoid of FcRs and express only very low levels of $Fc\gamma$ RI/CD64 and $Fc\gamma$ RII/CD32. In contrast, blood monocytes all express FcRs at a high level (40, 41, 44). No comparison of FcR expression between blood and mucosal neutrophils has been reported, although cervical and colonic neutrophils express Fc γ RIII/CD16 and Fc α RI/CD89 in a similar manner (45). Furthermore, we found that blood neutrophils express only Fc γ RII/CD32 and Fc α RI/CD89 (46).

2.2 Canonical type I FcR signalling

The high-affinity type I receptors, F*c*γRI/CD64 and F*c*αRI/CD89, both use the same intracellular machinery *via* ITAM signalling motifs

TABLE 1	Expression patt	ern of major FcR f	for IgG and IgA in	n human immune an	d non-immune cells.
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found in the cytoplasmic tail of the FcR γ chain, albeit the stoichiometry of IgA and IgG binding to their cognate FcR is different. Indeed, while only one Fc γ of the IgG binds to Fc γ RI/CD64 in a IgG:Fc γ RI/CD64 molar ratio of 1:1, each Fc α of the IgA recruits one Fc α RI/CD89 in a IgA:Fc α RI/CD89 molar ratio of 1:2 (8). Therefore, for an equal amount of Abs, the signalling cascade induced by IgA-IC *via* Fc α RI/CD89 is twice that induced by IgG-IC *via* Fc γ RI/CD64 (13). Worthnoting, monomeric and dimeric IgA have comparable affinities for Fc α RI/CD89, suggesting that Fc α RI/CD89 binds to both forms of IgA in a similar 2:1 stoichiometry (26, 33, 47). Similar to the interaction of Fc γ RI/CD64 with IgG, the medium and low affinity IgG receptors Fc γ RII/CD32 and Fc γ RIII/CD16 bind to the Fc γ region in a 1:1 molar ratio (10).

These Ab : FcR interactions lead to three types of intracellular signalling cascades as follows.

The first two types are triggered only by Abs in IC with free antigen or when they coat the surface of opsonised cells or microbes that can cross-link the receptors and in turn trigger cell activation (Figures 2B, C). During IgG-IC-induced activation, the Src Fyn tyrosine kinase phosphorylates at several sites either the FcR ITAM, i.e. FcγRIIa/CD32a, or the intracellular FcRγ-chain adaptor associated with FcγRI/CD64 (Figure 2B). Upon IgA-IC-induced activation, the Src Fyn tyrosine kinase phosphorylates the intracellular FcRγ-chain adaptor associated with FcαRI/CD89 (Figure 2C). In both cases, this phosphorylation allows the recruitment of SH2-adaptor proteins Syk, which in turn activate PI3K and PKC kinases, ultimately triggering the ERK pathway in association with an increase in cytoplasmic Ca2+ concentration. Together, these actors are responsible for cell activation, phagocytosis, ROS and NET formation, cytokine release and antigen presentation.

In contrast, the last type of intracellular signalling cascades is triggered when IgG forms an IC and engages FcyRIIb/CD32b but also when IgA is devoid of antigen. These two cases of Ab : FcR interactions lead to regulatory inhibition that contributes to immune homeostasis (Figure 2D). In fact, IgG-IC binds to FcyRIIb/CD32b containing ITIMs that become monophosphorylated. Alternatively, antigen-free IgA binds to FcaRI/CD89 triggering suboptimal phosphorylation of the ITAMcontaining intracellular adaptor FcRy, which is called ITAMi. In turn, monophosphorylation of ITIM and ITAMi recruits SH2-phosphatases that counteract Syk activity (48). Therefore, activations of ITIM and ITAMi compensate that of ITAM signalling in the same cells (30, 48). Suboptimal stimulation of FcyRIIa/CD32a by anti-FcyRIIa/CD32a Fab has also been reported to trigger ITAMi signalling in mice (9, 13, 30, 42, 49, 50). However, no suboptimal phosphorylation of ITAM (i.e. ITAMi signalling) has been reported upon engagement of uncomplexed IgG with FcyRI/CD64 and FcyRIIIb/CD16b, the latter associated with the ITAM-bearing adaptor chain FcRy, probably because the affinity of antigen-free IgG for both receptors is too low.

Although IgA can mediate signalling when it is devoid of antigen, IgG acts in the context of immune complexes *in vivo*. In this later case, the size of the immune complex, as well as Ab subclass composition and glycosylation contribute to the strength of FcR engagement and downstream effector functions (51). At the cellular level, the effector cell typically expresses multiple FcRs that can each be engaged by the multiple Fc domains of the Abs comprised in an IC. Therefore, activation (ITAM) and inhibition (ITIM and ITAMi) signals can be triggered simultaneously on effector cells. The resulting net effector function will depend on the balance between the extent of anti-inflammatory and pro-inflammatory FcRs signalling induced.

2.3 Other non-canonical FcRs

Two non-canonical FcRs have been described. The first, TRIM21, has been proposed to act as an intracellular cytoplasmic FcR for IgG and IgA (monomers or dimers) (52–54). By binding to Abs, TRIM21 triggers intracellular degradation of virus-Ab immune complexes by the Ub-ligase activity of TRIM21. This mechanism is independent of Ab neutralising activity, although it is referred to as Antibody-Dependent Intracellular Neutralization (ADIN). Indeed, ADIN refers to the neutralisation of the virus due to its degradation rather than the ability of Ab to directly block virus entry and infection of target cells (52–54). In addition, TRIM21 facilitates antiviral functions through proteasomal degradation, and exposure of the viral genome to cellular sensors (25, 52, 55).

The second, FcRn, a non-classical MHC-I molecule, is ubiquitously expressed. FcRn binds Ab only at low pH, which occurs after endocytosis of Ab by different FcyRs and its release into an acidic compartment. FcRn is responsible for the recycling of antigen-free IgG to the cell surface and its release at neutral pH. This recycling prevents antigen-free IgG from undergoing proteasomal degradation and results in a prolonged systemic half-life of IgG. In contrast, FcRn routes IgG-IC to degradation, resulting in cross-presentation (24). Therefore, binding of IgG to FcRn increases the cross-presentation of peptides to the corresponding specific CD8+ T cells in vitro (56), providing an additional bridge between adaptive and innate immunity. FcRnmediated cross-presentation may have contributed to a superior HIV-1-specific CD8+ T-cell response in non-human primates (NHPs) infected with recombinant simian virus-HIV-1 (SHIV) pretreated with HIV-1-specific IgG carrying an LS mutation that increases Ab binding to FcRn, compared to untreated animals (57). The role of FcRn in cross-presentation has been further confirmed in animal models of cancer (56, 58-60). This non-classical MHC-I molecule deserves further study. FcRn is also responsible for the bidirectional transport of IgG across the epithelium (24).

Abs can also bind FcR-Like receptors (FCRL), a family of receptors named by homology to the canonical FcγRI/CD64 (61). The FCRL family currently contains 6 members (1–6) and are preferentially expressed on B cells, at least for FCRL1-5. Yet, FCRL3 is also expressed by Th17 and Treg (6a3), while FCRL6 is mainly expressed by certain subsets of T cells and NK cells (62, 63). As recently reviewed (64), the intracellular region of FCRLs contains either a single ITAM domain, as in FCRL1, or an ITAM domain with a variable number of ITIM domains, namely from one to three in FCRL2-6. This heterogeneity in the ITIM/ITAM configuration probably leads to major functional differences between the various FCRL members. Nevertheless, only FCRL1 functions as an activating receptor, the others being considered as negative regulators.

Importantly, only FCRL3, FCRL4 and FCRL5 act as Ab receptors. FCRL3 appears to bind IgA but only in its secretory sIgA form, and interaction of artificially cross-linked sIgA with FCRL3 results in inhibition of Treg functions and promotion of a Th17 phenotype (65). This suggests that FRCL3 may act as a mucosal sensor of ICs promoting a local inflammatory environment, and driving regulatory T cell plasticity to help control pathogen invasion. This hypothesis has yet to be validated using sIgA naturally cross-linked by their specific antigen. In contrast, FCRL4 on memory B cells binds only to serum IgA but not to sIgA (66), and is thought to participate in mucosal tolerance. FCRL4 also binds weakly to heat-aggregated IgG3 and IgG4 (67), although further experiments with native IgG alone or in IC are needed to functionally validate these interactions. FCRL5 binds to all IgG isotypes although with different affinities (67, 68). In FCRL5 transfected cell lines, IgG binding to FCRL5 increases with Ab de-glycosylation, this later increasing with IgG lifetime (68). This result needs to be validated in primary FCRL5+ cells.

The biology and functions of FCRL are poorly understood. It has been proposed that FCRL3, FCRL4 and FCRL5 expressions are involved in B cell development in different B cell subsets (69–71). For example, FCRL5 cross-linking on B cells counteracts BCR signalling while increasing the proliferation index, a major feature for B cell development (69). As recently summarised, FCRL3-5 may exert important roles in mucosal protection, not only as immune-promoting sensors but also as depletion markers in autoimmune disorders and chronic viral infections (72). Overall, cross-linking of FCRLs by IC leading to the elimination of the Ab-targeted cell/microorganism occurs only in rare cases.

2.4 *In vivo* models to study the effector functions of antibodies

Animals such as mice or Non-Human Primates (NHPs) are key models to fully understand how Abs mediate effector functions and contribute to pathophysiology. However, one of the key questions in these studies is the specific engagement of Abs with FcRs, whether in mice or NHPs. The affinity of human antibodies for animal FcRs and, conversely, that of animal antibodies for human FcRs differ from an autologous situation. In particular, human IgG has a slightly lower affinity for mouse FcRs (including FcRn) than for human FcRs, and the level of FcyR expression varies between animal and human effector cells. These characteristics may limit the extrapolation of Ab-FcR interaction results from animals to humans (8, 10, 42, 73–76). Furthermore, as mice express pIgR but not FcaRI/CD89, the use of CD89 Tg mice is mandatory when studying IgA effector functions in mouse models (2, 73, 76). Alternatively, NHPs, a more human-like animal model, express all FcyRs, FcaRI/CD89 and pIgR, and are therefore more suitable for preclinical studies on therapeutic Abs, including IgA (73, 74, 77). Some FcyR expression patterns are peculiar, notably in neutrophils. In human, neutrophils express only FcyRIIa/CD32a and FcyRIIIb/CD16b whereas in NHPs, although lacking FcyRIIIb/CD16b, neutrophils express FcyRI/CD64, FcyRIIa/CD32a and FcyRIIb/CD32b. However, Ab and FcR species matching studies in the animal model would require the development of expensive humanised animal models, especially if the various human FcRs are expressed in each lineage and mediate the corresponding effector functions.

2.5 Fc-dependent and Ab-mediated effector functions

The integration at the effector cell level of activating and inhibitory FcR engagement after IC formation results in a wide range of functions including complement-dependent cytotoxicity (CDC), ADCC, ADCP, cytokine release, ROS production, transcytosis as well as recycling, as illustrated in Figure 3 for FcRmediated effector functions (8, 10). Antibody-mediated transcytosis and recycling, although crucial in Ab biology and engineering, have already been reviewed (24) and will not be discussed.

2.5.1 Complement dependent cytotoxicity

The recognition by Abs of antigens, expressed on the surface of tumour or virus-infected cells, leads to their opsonisation. In turn, complement is recruited by Abs to the cell surface, forming a multi-protein complex that disrupts the lipid bilayer of the plasma membrane and kills the opsonised cell (78). Both IgG and IgA are capable of recruiting the complement system, but using different mechanisms. These differences make a direct comparison between the complement-dependent activities of the two isotypes difficult (79–81). The two pathways converge to form the terminal complement complex C5b-C9, although the underlying initiation mechanisms remain to be described.

2.5.2 Antibody-dependent cellular cytotoxicity

ADCC is an effector function that relies on the interaction of Fc with an FcR on the surface of the effector cell, which triggers the destruction of the opsonised target cell (Figure 3). In cancer therapy, NK-dependent ADCC is one of the main mechanisms used by therapeutic Abs to eliminate cancer cells (82-84). Interestingly, monocyte-, macrophage- and neutrophil-induced ADCC is more effective when triggered by IgA- than IgG-opsonised cells. This IgA-dependent enhancement of ADCC is most likely due to the higher binding stoichiometry of Fca to FcaRI/CD89 (13, 15, 85, 86) than that of Fcy to FcyRI/CD64 mentioned above. In addition, IgG and IgA act synergistically to enhance lysis of HIV-1 infected cells by ADCC (85). This synergy can occur in vivo: indeed, the corresponding FcRs are constitutively expressed by effector cells such as primary monocytes, macrophages (85) and neutrophils (40). Of note, this is not the case in the commonly used myeloid cell line THP-1 (87).

2.5.3 Antibody-dependent cell phagocytosis

Antibody-dependent cell phagocytosis can be performed by monocytes, macrophages and neutrophils stimulated by opsonised cells through activation of Fc γ RIIa/CD32a, Fc γ RI/CD64 or Fc α RI/ CD89, resulting in the uptake of the opsonised target cell into a phagosome (Figure 3). In turn, the phagosome acidifies, forming a phagolysosome in which the opsonised target cell is finally lysed (1, 41, 88). Interestingly, the presence on the effector cell surface of FcRn in addition to Fc γ R increases the ADCP score (59, 89), suggesting cooperation between the two classes of FcRs. Uptake of ADCPopsonised target cells into antigen-presenting cells could result in antigen degradation, efficient cross-presentation (56, 90) and, in turn, in a specific CD8+ T-cell response (91–94). Importantly, this



FIGURE 3

Fc-dependent effector functions triggered by FcR activation. Fc-mediated functions (ADCP, trogocytosis, ADCC, NET formation, cytokine release and ROS formation) initiated by either IgG (light and dark blue) or IgA (red and yellow) engagement with the cognate FcR that activates ITAM. IgA and IgG immune complex antigens are in green. **ADCP**: Antibody-dependent cellular phagocytosis is a potent mechanism for eliminating Ab-coated pathogens or tumour cells. Engagement of FcγR or FcαRI/CD89 expressed on innate effector cells triggers a signalling cascade leading to engulfment of the pathogen by Ab. **ADCC**: Antibody-dependent cellular cytotoxicity is a mechanism by which FcR-expressing effector cells recognise, phagocytose and kill Ab-sponsored target cells. **Trogocytosis**: It is the rapid intercellular transfer of membrane fragments and their associated molecules during intercellular contact. Trogocytosis is typically mediated by the interaction of Fc-FcRs on effector cells with opsonised target cells. Ab-dependent trogocytosis of cancer cells by neutrophils and macrophages can result in target cell death (trogocytosis-mediated cell death). **NETosis**: Neutrophil-mediated target cell **death** after initiation by Abs is characterised by the secretion of large neutrophil extracellular traps (NETs) that capture and destroy target cells. **Cytokine** released depends on the type of FcR stimulated. **ROS** formation: Reactive oxygen species (ROS) induced by FcR cross-linking contribute to a variety of effects such as in ADCP, ADCC and NETosis.

mechanism provides an additional direct link between humoral and cellular immunity, similar to that established by FcRn.

Only a few studies have directly compared IgG-FcyRI/CD64mediated target cell ADCP versus IgA-FcaRI/CD89. On average, IgA induces ADCP more efficiently than IgG. HIV-1 envelopespecific IgA from a single RV144 vaccine recipient achieved robust ADCP (95), although in this study, IgG-mediated ADCP from the same individual was not studied. In addition, the HIV-1-specific broadly neutralising antibody (bNAb) 2F5 produced as an IgA isotype triggers ADCP more efficiently than when produced as an IgG isotype or compared to other anti-gp41 or -gp120 IgG bNAbs. This synergy occurs irrespective of effector cell type (46) and is consistent with other recent publications (12, 15, 16, 96-98). The few studies of IgA- versus IgG-mediated ADCP in bacterial infections have shown that monomeric IgA tends to protect humanised FcaRI/CD89 mice better against multidrug-resistant Mycobacterium tuberculosis infections than IgG, while polymeric IgA and its switched isotype IgG induce Neisseria meningitides phagocytosis in a similar manner (33, 99).

2.5.4 Trogocytosis

Alternative Fc-based mechanisms for mediating Ab-dependent target cell damage have been described. Firstly, the interaction

between opsonised cells and effector cells can result in the capture of a piece of the target cell membrane by the effector cell, which in most cases leads to target cell death. This mechanism is called trogocytosis (Figure 3). All $Fc\gamma R$ engagements, with the exception of $Fc\gamma RIIc/CD32c$ engagement by IgG, result in trogocytosis of opsonised cells by effector monocytes, macrophages and granulocytes (100–103). The relative efficiencies of trogocytosis mediated by different IgG subclasses have not been studied. Furthermore, the ability of mucosal IgA to induce trogocytosis is poorly described and comparative studies between isotypes are lacking (104).

2.5.5 NETosis

Secondly, the formation of nuclear extracellular traps (NETs) results mainly from neutrophil-mediated cell death (Figure 3). NETs are formed after neutrophil activation that triggers the secretion of antimicrobial granular peptides and the release of chromatin from the cell nucleus, which together form extracellular fibres in a process called NETosis (105, 106). NETosis is emerging as an additional tool in the arsenal of antimicrobial and anticancer strategies. NETosis can be induced by both IgG and IgA, although IgA is the strongest inducer (12, 107, 108). The ability of mucosal IgA to induce NETosis remains to be tested.

2.5.6 ROS formation

FcR cross-linking can stimulate the formation of intracellular ROS, which mediate a plethora of functions (Figure 3), including access of opsonised cells to phagosomes, NET formation followed by degranulation, and release of extracellular ROS (50, 106, 109). Thus, intracellular ROS act as second messengers to amplify ADCP and NETosis. Both IgG and IgA have been shown to stimulate ROS formation. However, IgA induces more ROS than IgG promoting NET, ADCC and ADCP (12).

2.5.7 Cytokine production

Finally, cross-linking of the FcR by Abs in the IC also stimulates cytokine release from the innate effector cell (Figure 3). The profile of secreted cytokines depends on the type of FcR stimulated and the resulting signalling. Pro-inflammatory FcRs, namely FcyRI/CD64, FcyRIIa/CD32a, FcyRIIc/CD32c, FcyRIIIa/CD16a, FcyRIIIb/CD16b, and FcaRI/CD89, induce the release of pro-inflammatory cytokines, including IL-6, IL-2, IL-12p40, IFNy and TNFa, while antiinflammatory FcRs, including FcyRIIb/CD32b, will trigger the release of anti-inflammatory cytokines, such as IL-10, or block the secretion of pro-inflammatory cytokines (110-113). Once secreted, cytokines can modulate the engagement of Abs in the FcR through a mechanism called internal and external signalling (114). For example, stimulation of neutrophils by GM-CSF increases the binding of IgG and IgA to the corresponding FcR, but does not alter their expression levels. In turn, neutrophils are more likely to induce ADCC and ADCP in cancer cells (115, 116). This mechanism is particularly effective when GM-CSF synergises with IgA-mediated activation of FcoRI/CD89, triggering PI3K signalling and, downstream, ADCC and ADCP (117). The balance between pro- and anti-inflammatory cytokine secretion depends on the net ITAM and ITAMi signalling induced on FcαRI/CD89 after engagement of IgA-IC or uncomplexed IgA. For IgG, this balance depends on the type of receptor involved, either FcyRIIa/CD32a and c carrying ITAM or FcyRIIb/CD32b carrying ITIM (2, 42, 118, 119).

Overall, there is consensus on the IgA-mediated increase in effector functions. As the isotype differences probably rely on a higher binding stoichiometry to their corresponding FcR, the more FcRs engage Abs, the more efficiently downstream effector functions will be stimulated. With a greater ability to engage downstream ITAM activation than IgG (13, 18, 85, 120), IgA should be considered in the design of therapeutic Abs, alone or in combination with IgG, to enhance effector functions.

3 Chimeric antibodies containing Fc γ -Fc α enhance Fc-mediated functions: From the laboratory to the bedside

3.1 Increasing the affinity and binding avidity of Ab to FcRs

Knowing that effector functions rely mainly on the Fc domains of Ab's, the insertion of additional Fc domains into the design of Abs could result in chimeric antibodies with enhanced FcR binding and downstream Fc effector functions and therapeutic activities. To this end, from the parental Abs harboring one Fc γ (Figure 4A), chimeric Abs against respiratory syncytial virus (RSV) were designed with two or three Fc γ domains, resulting in a chimeric 2-Fc Ab with a tetrahedral-like geometry (121), as shown in Figure 4B. Surface plasmon resonance experiments showed that each 2-Fc chimera molecule binds to two Fc γ R molecules, thereby improving the Fc avidity of the Ab while maintaining the Fab antigen specificity of the original Ab. Each 2-Fc chimera molecule also binds to four FcRn molecules, thereby prolonging the pharmacokinetics of the chimera. However, this strategy is not optimal because significant amounts of single-chain Fc-Fab are produced along with the 2-Fc chimera. The authors did not evaluate whether increasing the avidity of the 2-Fc chimera for Fc γ Rs improved effector functions (121).

Furthermore, Rituximab, an anti-CD20 IgG, harbouring tandemly duplicated or triplicated Fcys broadens the spectrum of bound FcyRs from FcyRI/CD64, the only FcyR bound by the parental IgG, to FcyRIIa/CD32a (122). Simultaneous binding of FcyRI/CD64 and FcyRIIa/CD32a results in a fivefold increase in ADCC and ADCP activities against CD20 + tumour B cells in mice *in vivo* (122, 123). Such a strategy of Fcy chimeras has not yet been developed against infectious pathogens.

3.2 IgG-IgA bi-isotype chimerism

IgG-IgA bi-isotype Fc chimerism was initially explored with the socalled 'IgGA cross-isotype' format, in which the Fc domain of the anti-HER2 IgG1 therapeutic, Trastuzumab, was genetically engineered to acquire the ability to bind to both the $Fc\gamma R$ and $Fc\alpha R$ (Figure 4C) (124). The authors elegantly determined the residues controlling the engagement of Fc α and γ with their cognate FcRs by first using alanine scanning approaches. As a result, the amino acids CHy₁2 and CHy13, which are not responsible for IgG engagement with the FcyR, were replaced with the corresponding amino acids of $CH\alpha_1 2$ and $CH\alpha_1 3$ responsible for IgA binding to the FcaRI/CD89. The resulting Trastuzumab IgGA chimera contained both Fca and Fcy elements. This cross-isotype IgGA chimera was well produced and had an affinity for FcyRs comparable to that of the parental IgG, although the affinity for FcaRI/CD89 decreased by two compared to the parental IgA. In addition, the affinity for C1q increased compared to parental Trastuzumab IgG and IgA. Due to its ability to bind Fcq- and FcqR, the IgGA-Trastuzumab chimera mediated higher ADCC and ADCP than the parental IgG, but did not reach the level mediated by Trastuzumab-IgA. Furthermore, the cross-isotype format showed a higher CDC compared to IgG and IgA isotypes (124). Unexpectedly, Fc engineering in the IgGA chimera abolished commitment to FcyRIIIa/ CD16a and FcRn, suggesting low bioavailability in vivo (24). In this case, isotypic cross-transformation may therefore decrease therapeutic benefit. This study indicates that mutations focusing exclusively on amino acids directly involved in the interaction with the FcR are not functionally effective: such point mutations would probably be too targeted and unable to preserve the overall structure of each Fc domain essential for FcR engagement by Fc α and Fc γ together. Instead, it can be hypothesised that expression of the full $Fc\alpha$ and $Fc\gamma$ is necessary to endow a chimeric Ab with a true gain of ADCC and ADCP effector functions accompanied by extended pharmacokinetics through enhanced FcRn engagement.

To enable an IgG to target FcoRI/CD89 in addition to FcyR (using its Fcy), bi-specific IgGs with one Fab specific for FcαRI/CD89 have been designed. An example of this strategy is given by TrisoMab, a bispecific IgG (more precisely IgG1) directed against both EGFR and FcoRI/CD89 (14). This chimera offers the advantage of targeting three distinct molecules (Figure 4D). Indeed, TrisoMab recognises EGFR, the target of the parental IgG, via one Fab, as well as FcoRI/ CD89 via the second Fab, while retaining its ability to engage FcyR via the Fcy antibody (14). Consequently, TrisoMab binds to effector cells, namely neutrophils, NKs and macrophages, in turn amplifying effector functions in vitro and in vivo. As a result, neutrophils were recruited more efficiently to tumour colonies, showed improved TrisoMab-mediated ADCC and trogocytosis of cancer cells compared to the parental EGFR-specific IgG, However, the improvement over EGFR-specific IgA, whose Fab targets EGFR and own Fca targets FcaRI/CD89, was limited. The enhancement of trogocytosis by TrisoMab may have resulted in cross-presentation of tumour antigen allowing the generation of therapeutically beneficial cytotoxic CD8+ T cell activity. In addition, macrophages showed an increase in TrisoMab-mediated tumour cell ADCP and ADCC compared to EGFR-specific IgG and IgA. Finally, in mice humanised for CD89, TrisoMab showed a prolonged half-life that was double that of EGFR-specific IgA, although similar to the IgG isoform. Improved survival and smaller tumour size in an hCD89+ mouse tumour model (14) demonstrated a therapeutic gain of the chimera over the parental EGFR-specific IgG and IgA. These results

illustrate the chief role of the IgA-Fc α RI/CD89 axis in Abbased therapy.

As this chimera lacked an $Fc\alpha$ domain, the true stoichiometry of an Fca activating two FcaRI/CD89 molecules was not met and rather limited to the activation of a single FcxRI/CD89 molecule (targeted by the TrisoMab anti-CD89 Fab. This reduced engagement of FcoRI/CD89 by TrisoMab could not take advantage of the regular 1:2 IgA : FcxRI/ CD89 stoichiometry. This may therefore have limited the role of FcoRI/ CD89-mediated effector functions. Nevertheless, although not bearing a bona fidae bi-isotype Fc, a TrisoMab molecule activated both FcyRI/ CD64 via its Fcy and FcoRI/CD89, using an anti-CD89 Fab instead of an Fco, with a clear and significant therapeutic advantage. The lower ratio of one chimeric TrisoMab to one $\mbox{Fc}\alpha\mbox{RI/CD89}$ compared to that of two Fca, present in a single IgA molecule, to one FcaRI/CD89 may have limited the potential benefit of IgG-IgA bi-isotype chimerism. It may also explain the lack of complete remission in treated mice. It would therefore be interesting to study a tandem Fcα-Fcγ chimerism combining both Fcγ and $Fc\alpha$ in the same molecule. Such a chimeric Ab would increase the chimeric:FcR binding ratio to 1:3 by simultaneous binding of one FcyRI/ CD64 by Fcy and two Fcc/RI/CD89 by Fcc, compared to a ratio of one parental IgG to one FcyR and one parental IgA to two FcoRI/CD89. Such a chimera would be expected to increase net ITAM signalling due to the engagement of two FcoRI/CD89s in addition to the pre-existing FcyRI/ CD64 (13), thereby enhancing effector functions. Again, no studies using such a bi-isotype Fc chimera enhancing FcoRI/CD89 signalling have been evaluated against infectious diseases.



Fc chimera and parental antibodies. (A) Parental IgA (HC: red, LC: yellow) and IgG (HC: dark blue LC: light blue). (B) Fc γ duplication (dark blue) in tandem or tetrahedral form, (C) Cross-isotype IgGA harbouring a hybrid of Fc γ (dark blue) and Fc α (red), (D) Anti-CD20 and Fc α RI/CD89 bi-specific IgG isotype. The chimera can bind both to FcR *via* its authentic and to Fc α RI/CD89 *via* the anti-CD89 that replaces Fc α used in other constructs. (E) Chimera with Fc α -Fc γ expressed in tandem. (F) Chimera with Fc γ -Fc α expressed in tandem. Both Fc γ -Fc α and Fc α -Fc γ chimeras can harbour either an IgA2 hinge, a linker (G4S)4 or an IgG1 hinge between the two Fc chains.

To overcome these limitations, Borrok et al. designed a novel chimera, the anti-HER2 IgG1 Trastuzumab, in which the Fc α (specifically Fc α 2) and its corresponding hinge were fused into the C-terminus of the Fc γ of the original IgG (17), as shown in Figure 4E. The chimera was produced at a level comparable to IgG and three to four times higher than Trastuzumab-IgA. The affinity of the chimera for each Fc γ R and Fc α RI was comparable with only a slight decrease in C1q affinity compared to parental IgG1. After engagement of both Fc γ RI/CD64 and Fc α RI/CD89 on effector cells, the chimera enhanced ADCC and ADCP of human polymorphonuclear cell-mediated tumour cells and macrophages compared to parental IgG and IgA. In addition, the chimera maintained a similar half-life to that of the parental IgG when injected intravenously into mice. In contrast, that of IgA injected into parallel animals rapidly decreased.

This important study revealed that the presence of human fulllength $Fc\alpha$ and $Fc\gamma$ in tandem in the same chimeric Ab improved Fc α - and Fc γ -mediated functions (17). The limited pharmacokinetics observed in this study could be due to the specie differences between human Ab and mouse FcR in this mouse model, probably in the glycosylation pattern of Fc. Indeed, Fc glycosylation is known to affect both effector functions and half-life of Abs (16, 125, 126). With this in mind, Li et al. developed a Rituximab chimera containing both human Fc α and Fc γ . Then, they evaluated Rituximab efficacy in vivo using Tg mice expressing hFcaRI/CD89 under the control of a CD14 promoter to target myeloid expression (18). In this study, the construct had the Fc γ connected to the Fc α with a polyglycine linker, with both Fc's expressed in tandem. Functionally, in a mouse tumour model, the chimera mediated ADCC in tumour cells at levels similar to parental IgA, but greatly increased compared to parental IgG. Furthermore, when treated with the chimera, Tg $hFc\alpha RI/CD89$ mice had a smaller tumour volume compared to treatment with parental IgG or IgA. All these functional consequences were entirely dependent on hFcaRI/CD89 expression, as no effect was observed in wild-type mice (18).

Taken together, these results support the development of biisotypic Fc-chimeric therapeutic Abs with enhanced effector functions. Such enhancement relies on the presence of full-length Fc α and Fc γ . It should be noted that the role of the IgA hinge region may not affect functional efficacy (17, 18). However, Fc-chimeric Abs in infectious diseases have not been evaluated.

3.3 Fc chimeras: Remaining questions and perspectives

The design of bi-isotype Fc α and Fc γ chimeras is very promising and could offer a powerful tool to target tumour cells using their enhanced Fc effector functions, especially in patients with very aggressive or resistant tumours. This anticipated enhancement would rely on increased affinity/avidity of the chimeras for their cognate FcRs and the subsequent increase in downstream signalling. This increased signalling would occur (via the ITAM domain of the FcR γ chain (paired with Fc γ RI/CD64 and Fc α RI/CD89) or the FcR itself (for Fc γ RIIa/ CD32a or Fc γ RIIc/CD32c). The total phosphorylation of ITAM induced by the chimera compared to parental IgG and IgA is expected to increase Syk recruitment to ITAM, subsequent cellular activation and effector functions, as illustrated in Figure 5.

Production remains an important limiting criterion in the development of such chimeric constructs. However, according to the literature, the chimeras are produced at a good level, at least comparable to that of the parental IgG. The role of the position of Fc α in the IgG/IgA chimeric Ab on effector functions remains to be explored. It could affect the respective binding of FcRs and, consequently, their ADCC and ADCP levels. Furthermore, the role of the Fab isotype, including that of the CH1 domain, also remains a parameter to be studied. In this respect, we revealed that the CH1 domain plays a crucial role in Fab antigen recognition (both in affinity and specificity) with a superiority of CH1 α over CH1 γ (127). Surprisingly, the higher molecular weight of $Fc\alpha$ -Fcy or Fcy-Fcy tandem chimeras compared to parental Abs was not taken into account in these studies, probably introducing a bias. Indeed, comparing the impact of chimerism on effector functions between the chimeras and the parental IgG and IgA tested at the same molar concentration would be more rigorous than testing them at the same weight/volume concentration.

Glycoengineering is another popular method currently used to improve the functions of anti-tumour Abs, although it is mainly limited to the IgG isotype (84, 128). The impact of the glycosylation pattern of Fca and Fcy in chimeric Abs on therapeutic benefit remains to be investigated (86, 125) as IgA has both N- and O-glycosylations (3) and IgG has only N-glycosylations. However, the O-glycosylations of the IgA1 hinge may be too far from the CH2-CH3 domain to affect IgA interaction with Fc α RI/CD89 and downstream effector functions. The impact of IgA glycosylation on Fc-mediated functions has been poorly studied. Therefore, a potential role for IgA glycosylation in the functions of $Fc\alpha$ -Fc γ chimeric Ab is difficult to predict. Nevertheless, desialylation of IgA1, which has been shown to enhance its proinflammatory properties (107), could improve the therapeutic benefits of Fca-Fcy chimeric Abs. However, this study did not address the impact of defucosylation on IgA-mediated effector functions. In contrast for IgG, such glycoengineering is a very promising approach (84).

Finally, the addition of an Fc α to the rapeutic Abs could enhance the delivery of chimeric Ab to mucosal surfaces via Dectin-1, an alternative IgA receptor (129), and/or FcRn (58) capable of reversing Ab transcytosis in mucosal tissues.

3.4 Therapeutic Fc α -Fc γ antibodies as future anti-HIV-1 bNAbs

Given the role of the IgA isotype against viruses on the mucosal surface where $Fc\alpha RI/CD89$ is specifically expressed (40), we hypothesise that such tandem $Fc\alpha$ - $Fc\gamma$ chimeric Ab (IgA with a fused $Fc\gamma$, Figure 4F) would be effective in the prevention of infectious pathogens such as sexually transmitted HIV-1 (130). Indeed, over the past decade, anti-HIV-1 bNAbs have been developed to either protect from infection or cure HIV-1 infected patients using passive Ab therapy (alone or in combination with antiretroviral therapy) (131). As with anti-cancer drugs, anti-HIV-1 antibodies have been engineered mainly by Fab domain engineering to improve affinity for the viral antigen and to diversify antibody specificity by using bior tri-specific antibodies (57, 126, 132–135). Again, Fc engineering has rarely been studied.



one $Fc\gamma RI/CD64$ (left) and the IgA immune complex binds to two $Fc\alpha RI/CD89$ (middle), the $Fc\alpha$ - $Fc\gamma$ chimera immune complex binds simultaneously to one $Fc\gamma RI/CD64$ and two $Fc\alpha RI/CD89$, This increases downstream ITAM (red) phosphorylation (white circle) and subsequent Syk recruitment (dark green), thereby enhancing the functions of immune effector cells such as ADCC and ADCP.

Despite recent advances in the beneficial role of anti-HIV-1 IgA at the mucosal level (20, 120, 129, 136-139), IgA is routinely overlooked in anti-HIV-1 immunotherapy. As an additional argument for the use of IgA, others and we have shown that IgG and IgA cooperate in effector functions in vitro (85, 97) and in vivo in the protection of vaccinated animals against repeated mucosal challenges (20, 120, 129, 136-139) as well as in the protection of vaccinated individuals as in the RV144 phase III HIV-1 clinical trial (21, 22). From these studies, it can be suggested that $Fc\alpha\text{-}Fc\gamma$ tandem chimeras engineered from bNAbs would be beneficial in protecting against mucosal acquisition of HIV-1 by triggering FcoRI/CD89 and FcyRI/CD64 signalling, thereby enhancing effector functions (Figure 5). This approach would complement the inhibition of mucosal entry through transcytosis and virus neutralisation. These antiviral activities are essential in the protection against sexual acquisition of HIV-1. Hence, eradication of the virus is not yet possible once infection and the resulting viral reservoir are established (20, 120, 129, 136-140). Such chimeric Abs could enhance ADCC involved in protection but also ADCP, NETosis and cytokine secretion that contribute to the establishment/maintenance of a protective immune response (20, 45, 120, 129, 136-139, 141). In addition, bi-isotype Fc chimeras could contribute, in combination with other drugs/approaches, to shock and kill strategies aimed at eliminating the HIV-1 reservoir in HIV-infected patients on suppressive antiretroviral therapy. The combination of Fab engineering and tandem $Fc\alpha$ - $Fc\gamma$ in a single chimera could be superior to specific bi/tri constructs and tandem $Fc\alpha$ - $Fc\gamma$ chimera alone.

Fca-Fcy chimeric Abs can also interact with FCRL4 and 5 on B cells to promote a memory phenotype (69, 71, 72). We have previously shown that human penile mucosa B cells express both FCRL4 and FCRL5 (142). We can suggest that during passive infusion in HIV-infected patients, $Fc\alpha\text{-}Fc\gamma$ tandem chimeras may stimulate mucosal memory B cells, and in turn plasma cells secreting protective anti-HIV-1 IgA. The latter are associated with the prevention of mucosal acquisition of HIV-1 in actively and passively vaccinated macaques and in Highly Exposed Sero-Negatif Individuals that resist to HIV infection despite unprotected sexual intercourses (19, 20, 120). Furthermore, by binding to FCRL3, Fcα-Fcγ chimeras formulated as secretory-type IgA can induce a shift from a Treg to a Th17 profile (65). The latter could contribute to the eradication of HIV reservoirs, as well as to the promotion of tumour invasion by immune effector cells, resulting in tumour clearance (65). However, sIgA cannot bind to FcaRI/CD89 and the chimerism relevant for direct effector functions could be lost in such constructs (31). Nevertheless, FCRL3-5 engagement could also lead to inhibition of B-cell functions, as described elsewhere (143-145). However, the clinical

relevance of binding of therapeutic Abs to FCRL3-5 on B and T cells has not been studied. Therefore, the prediction of the biological effects of these chimeras remains uncertain. Furthermore, as FCRL6, an FcRL expressed on NK cells, cannot bind Abs, it will not interact with chimeric Abs to mediate ADCC by NKs (146).

Finally, the production of chimeric Abs could help to improve the half-life. In particular, the production of anti-HIV-1 IgG using adenoassociated vectors in mouse models prolongs its half-life compared to *in vitro* production in CHO cells (126). As the same coding sequence was used in both systems, this effect may depend on the glycosylation profile of Ab. This strategy could be beneficial for improving the half-life and effector functions of therapeutic HIV-targeted bNAbs, although the packaged coding sequence must be very compact (<4.4kb).

4 Conclusions and perspectives

The development of chimeric Fca-Fcy Abs as drugs still needs improvement, and several unexplored questions remain. The most difficult one concerns the best route of administration for the most relevant bio-distribution and stability to improve clinical efficacy. Secondly, the role of the Fca isotype, i.e. IgA1- or IgA2-derived Fca, in enhancing effector functions needs to be deciphered (147). Similarly, the role of IgG1- versus IgG3-derived Fcy in these chimeras is questionable, with IgG3 being more pro-inflammatory than IgG1, although this relies on the IgG3 hinge (148). The pharmacokinetics should also be evaluated in NHPs to validate the results obtained in mouse models, as well as their half-life at different anatomical sites after intravenous or mucosal applications. Furthermore, we propose that these Fc-based chimeric Abs could offer a more effective treatment against mucosal infectious pathogens, and the generation of a higher CD8+ T cell memory response, providing vaccine-like properties to the chimera. In the case of HIV-1, these Fc-based chimeric Abs would overcome the viral resistance resulting from antiretroviral drugs and offer less toxic alternatives to HIV/AIDS patients on antiretroviral therapy. Advances in Ab engineering, probably assisted by the development of algorithms, undoubtedly rely on the constant updating of structure-function relationships of Abs. Finally, there is

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an increasing need to develop high-throughput screening strategies for the evaluation of Ab efficacy and the improvement of associated production pathways. All these features should encourage and stimulate the development of Ab drugs, the next challenge in immunotherapy.

Author contributions

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Conflict of interest

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