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# The current understanding of the phenotypic and functional properties of human regulatory B cells (Bregs)

Nawara Faiza Ahsan<sup>1,2</sup>, Stella Lourenço<sup>3</sup>, Dimitra Psyllou<sup>1</sup>, Alexander Long<sup>1</sup>, Sushma Shankar<sup>4</sup>, Rachael Bashford-Rogers (D<sup>1,5,\*</sup>

<sup>1</sup>Department of Biochemistry, University of Oxford, Oxford OX1 3QU, United Kingdom

<sup>2</sup>Centre for Human Genetics, University of Oxford, Oxford OX3 7BN, United Kingdom

<sup>3</sup>Keizo Asami Institute, Federal University of Pernambuco, Recife 50740-520, Brazil

<sup>4</sup>Nuffield Department of Surgical Sciences, University of Oxford, Oxford OX3 7DQ, United Kingdom

<sup>5</sup>Oxford Cancer Centre, University of Oxford, Oxford OX3 7LH, United Kingdom

\*Corresponding author: Department of Biochemistry, University of Oxford, South Parks Road, Oxford OX1 3QU, United Kingdom. E-mail: rachael.bashford-rogers@bioch.ox.ac.uk

#### Abstract

B cells can have a wide range of pro- and anti- inflammatory functions. A subset of B cells called regulatory B cells (Bregs) can potently suppress immune responses. Bregs have been shown to maintain immune homeostasis and modulate inflammatory responses. Bregs are an exciting cellular target across a range of diseases, including Breg induction in autoimmunity, allergy and transplantation, and Breg suppression in cancers and infection. Bregs exhibit a remarkable phenotypic heterogeneity, rendering their unequivocal identification a challenging task. The lack of a universally accepted and exclusive surface marker set for Bregs across various studies contributes to inconsistencies in their categorization. This review paper presents a comprehensive overview of the current understanding of the phenotypic and functional properties of human Bregs while addressing the persisting ambiguities and discrepancies in their characterization. Finally, the paper examines the promising therapeutic opportunities presented by Bregs as their immunomodulatory capacities have gained attention in the context of autoimmune diseases, allergic conditions, and cancer. We explore the exciting potential in harnessing Bregs as potential therapeutic agents and the avenues that remain open for the development of Breg-based treatment strategies.

Keywords: B cells; regulation; Bregs; autoimmune diseaseinfectionIL-10IL-25TGF-B

#### Introduction

B cells are key players in shaping the humoral immune system by antibody production, opsonization, antigen presentation, and activation of T cells. B cells are historically acknowledged as promoters of adaptive immune responses. An imbalance or dysfunction of B cells can result in suboptimal defence against infections, impaired cancer surveillance, and a breakdown of immune tolerance causing autoimmune or autoinflammatory diseases. More recent studies have recognized the immunoregulatory properties of B cells including maintenance of immune tolerance, and suppression of immune responses. In particular, a subset of B cells named regulatory B cells (Bregs) have demonstrated immune suppressive properties in conditions such as allergies, autoimmune diseases, infection, transplantation, and tumours.

The presence of a B cell population with immunosuppressive properties was first suggested in the 1970s, based on findings that B cell-depleted splenocytes correlated with the severity and duration of delayed-type hypersensitivity (DTH) in guinea pig models [1, 2]. This hypothesis of a suppressive B cell population was further supported by studies demonstrating that B cells regulated the differentiation of T cells into suppressor T cells and could act as a suppressive-inducer in feedback control [2, 3]. Janeway *et al.* confirmed the existence of an immunoregulatory subset of B cells which when adoptively transferred in B cell-deficient mice enabled complete recovery from experimental autoimmune encephalomyelitis (EAE) [3]. Soon after, Mizoguchi and colleagues reported a suppressive function of B cells in colitis and intestinal inflammation in mice models and named them 'regulatory B cells' [4–6]. Menna Clatworthy's team were the first to report findings that suggested the existence of human Bregs when they observed that the use of an anti-CD20 monoclonal antibody therapy that depleted B cells in transplant recipients increased the rate of organ rejection, such that the clinical trial was ceased [7]. Human Bregs were then more firmly identified in 2010 by Claudia Mauri's group in the context of Systemic Lupus Erythematosus (SLE) [8]. Since then, our understanding of the characteristics, phenotypes, and functionalities of Bregs in the context of cancer, infection, and other autoimmune diseases has significantly increased.

Briefly, Bregs are capable of modulating immune responses by inhibiting effector functions of CD4<sup>+</sup> and CD8<sup>+</sup> T cells, promoting differentiation of regulatory T cells (Tregs), and producing immunosuppressive cytokines such as Interleukin 10 (IL-10), Transforming growth factor beta (TGF- $\beta$ ), Interleukin 35 (IL-35) [9]. Further, these cells can suppress monocyte proliferation, and interferon-gamma (IFN- $\gamma$ ) production by natural killer (NK) cells, in an IL-10 dependent manner [10]. While Breg deficiencies and/

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or defects have been seen in autoimmune disorders [6], chronic inflammation [11], and graft rejection [12], their abundance has been associated with tumour progression and poorer outcomes in cancer [12], in both animal models and humans. Such diverse functional properties of Bregs makes them potential cellular targets across a range of diseases, including Breg suppression or depletion in cancers and infection, or Breg induction in autoimmunity, allergy, and transplantation. However, harnessing the clinical potential of Bregs is limited by our poor understanding of human Breg biology.

While our insights on the biology of Bregs have expanded in recent decades, most research has been conducted on murine Bregs. As such our understanding of the origin and differentiation of human Bregs remains unclear, which is further hampered by a lack of unique phenotypic markers or transcriptional markers for characterizing human Bregs, as well as contextdependant differences in phenotypes and secretory products and discrepancies amongst papers causing further complications. This review discusses our current knowledge on origin, differentiation, and phenotypic diversity of human Bregs, and elaborates on the functional roles of Bregs in cancer, infection, and autoimmune diseases highlighting opportunities for clinical application and future research.

#### **Origin and differentiation of Bregs**

The progenitors of B cells differentiate from common lymphoid progenitors in the bone marrow, where they undergo a series of progressively committed differentiation steps into early pro-B, pro-B, and pre-B cells. Pre-B cells develop into immature B cells that leave the bone marrow becoming transitional B cells in the spleen, before finally differentiating into naïve mature B cells [13]. These cells can further differentiate into three broad populations namely marginal zone B cells, follicular B cells, and plasmablasts (which ultimately become plasma cells), each with specific functional roles in humoral response [13].

While the position of the developmental pathway of Bregs on the classical B cell differentiation process remains unclear, two potential mechanisms have been suggested. The initial hypothesis deemed Bregs as individual lineages of B cells with specific transcription factors (TFs) conferring expression of immunosuppressive genes. The rational coming from other lymphocytes being derived in this manner (such as Tregs and Foxp3) [9]. However, no study has been able to validate the presence of any lineage-specific markers yet which has led to a second theory that B cells, at different developmental stages, acquire regulatory properties in response to environmental stimuli, such as cytokines, inflammation, and cell signalling. This latter hypothesis is supported by studies documenting that immature B cells, mature B cells, and plasmablasts in both humans and mice can differentiate into Bregs in response to stimuli [9]. This is supported by Glass et al. using mass-cytometry, and live-cell tracing on ex vivo stimulated PBMCs to show that IL-10<sup>+</sup> human B cells arise from multiple subsets, including CD45RB+ CD27+ memory, CD24+ CD38<sup>+</sup> transitional, CD19hiCD11c<sup>+</sup> effector B cells [14]. While immunosuppressive properties of antibody-producing and memory B cells contradict the conventional understanding of their role in inflammation, such regulatory properties of Bregs might be limited to specific subsets of these B cell subpopulations or when these cells are subject to certain stimulatory conditions within the body [15, 16]. This is suggested by the differentiation of regulatory plasmablasts in mice lacking an essential plasma cell transcription factor BCL6 [15]. Hence, it is important to identify

factors inducing differentiation of human Bregs. One transcription factor linked towards immunoregulatory activity in humans is the STAT3 signalling pathway. Shankar *et al.* observed increased STAT3 phosphorylation within their expanded Breg population as well as decreased TIM-1 expression when STAT3 was inhibited [17].

Currently established signals involved in inducing Breg differentiation are predominantly through Toll-like receptor and/or CD40 activation. Toll-like receptors (TLRs) play a crucial role in the development of innate and inflammatory responses by recognizing pathogen-associated molecular patterns (PAMPs) and damage-associated molecular patterns (DAMPs). Stimulation of TLRs in B cells results in differential production of proinflammatory and anti-inflammatory cytokines [9, 18]. The significance of TLR signalling in regulating suppressive functions of B cells is evidenced by the development of chronic EAE in mice with B cell-restricted deficiency of TLR signalling molecules MyD88, TLR2, and TLR4 [19]. CD40 signalling is also important for Breg activation; CD40 expressed by B cells binds to CD40 Ligand (CD40L or CD154) expressed by activated T cells, to facilitate B cell differentiation into plasma cells, whereas continuous CD40 signalling limits antibody production [20]. Contradicting conventional beliefs, several studies support the role of CD40 signalling in promoting activation of immunosuppressive Bregs. Mice with CD40-deficient Bregs possess fewer IL10-producing cells, increased T helper type 1 (Th1) and T helper type 1 (Th17) responses, and develop a more severe form of EAE [19]. Indeed, Shankar and colleagues demonstrated that CD154-mediated stimulation of human B cells by co-culture with CD154<sup>+</sup> CHO cells, could drive differentiation and expansion of IL10<sup>+</sup> Bregs [17]. Notably, the strength of the CD154 signal determined whether Bregs or activated, non-immunoregulatory B cells were produced. Another important factor in Breg induction is antigen recognition by B cell receptor (BCR), which when impaired inhibits Breg activation and IL-10 production [21], suggesting BCRdependency in Breg generation.

Differentiation and activation of Bregs have also been shown to be affected by inflammation. This is supported by the increase in Bregs during inflammatory phases of autoimmune conditions, while their absence is associated with more severe episodes of arthritis and EAE in mouse models. For example, one study showed an expansion of Breg populations in arthritic mice in response to proinflammatory cytokines such as IL-1b and IL-6 with their absence leading to worsened symptoms [22]. In addition, stimulating B cells with other inflammatory cytokines, such as IFN- $\alpha$ , IFN- $\beta$ , IL-21, and GM-CSF, in combination with IL-15, has also shown to facilitate Breg cell differentiation [18, 23]. However, many of these findings were performed in mice, and it is currently unclear whether these observations will apply also to human Bregs.

#### Mechanisms of action of Bregs

Bregs have been reported to have a variety of mechanisms of action through both cell surface receptor interactions with target cells as well as through secreted molecules, summarized in Fig. 1 and Table 1.

#### Phenotypic diversity of Bregs

Despite the lack of consensus on a marker characteristic to Bregs, several subpopulations with distinct surface markers, effector molecules, and functions, have been reported in both



Figure 1. Key mechanisms of Bregs on suppressing other immune cells. (Top) Cell surface receptors expressed by Bregs with known immunosuppressive effects on iNKT and CD4 T cells, and (bottom) secreted factors with known immunosuppressive effects on CD4 and CD8 T cells as well as Tregs. Created in BioRender. Wu, S. (2024) BioRender.com/b10q919.

human and mice models. Bregs can be also classified into ontogenic subsets, based on the differences in activation modes and resulting cytokine production and cell surface protein production that influence immune cell function within their environment [47]. Thus far in humans, Bregs have been reported to produce three suppressive cytokines namely, IL-10, IL-35 and TGF- $\beta$ . These three categories are used more widely in literature to classify Bregs. However, multiple other Breg subpopulations, characterized by expression of surface markers or protease production, have been noted in both humans and mice.

#### IL-10-producing Bregs (IL-10<sup>+</sup> Bregs)

IL-10 is an immunosuppressant cytokine that inhibits the production of IFN-γ and TNF-α by antigen-specific T cells, the release of inflammatory mediators from dendritic cells (DCs) and mononuclear cells, and differentiation of Th17 cells [48–50]. In the absence of a specific marker, expression of IL-10 is commonly used to identify Bregs in both humans and mice. An expansion of Il- $10^+$  B cell populations have been noted in patients with autoimmune conditions such as arthritis, multiple sclerosis (MS), Systemic lupus erythematosus (SLE) [51, 52], cancers (e.g. breast cancer, gastric cancer (GC)) [10, 53, 54], and infections (e.g. lymphocytic choriomeningitis virus, and vaccinia virus) [55]. However, several groups have demonstrated that whilst a wide spectrum of stimulatory conditions can induce IL-10 expression by B cells, not all IL-10-producing B cells are necessary regulatory [14, 56, 57]. This has made the characterization of definitive cell surface phenotypes of IL-10<sup>+</sup> Bregs somewhat challenging.

Multiple IL-10<sup>+</sup> Breg subsets exist, and we will discuss the main ones here. Tedder's group functionally defined and enumerated a significant proportion of IL-10<sup>+</sup> Bregs by their competence to express IL-10 following 5 hours of ex vivo stimulation with PMA and ionomycin, referring to such Bregs as 'B10' cells. Studies have shown that B10 cells and proB10 cells did not express CD38 and were memory in origin (CD24hiCD27<sup>+</sup> Bregs in human) [58–60]. However, many other IL-10<sup>+</sup> Breg populations can be stimulated to produce IL-10 under different conditions, for example CD24<sup>hi</sup>CD38<sup>hi</sup> Bregs [8] CD24<sup>hi</sup>CD27<sup>+</sup> [61], CD1dhiCD5<sup>+</sup> IL-10<sup>+</sup> B cells in the spleen of mice and humans, CD19<sup>+</sup>CD24<sup>hi</sup>CD27<sup>int</sup> IL-10<sup>+</sup> plasmablasts [62, 63] in autoimmune conditions, and CD25hiCD71hiCD73loIgG4<sup>+</sup> IL-10<sup>+</sup> B cells during allergic reactions. Interestingly regardless of their surface marker combination, these IL-10<sup>+</sup> B cells regulate immune responses in different conditions by producing IL-10.

Initially it was hypothesized that IL-10<sup>+</sup> Bregs differentiate from a pro-IL-10<sup>+</sup> Breg population in response to BCR stimulation by antigen/other cells [62]. However, the presence of B10 cells at different stages of maturation indicates a multi-lineage nature of Bregs, or that IL-10 production occurs in response to microenvironmental stimuli at different stages of B cell development, evidenced by the phenotypic diversity in surface marker expression of B10 cells [62, 64]. For instance, human IL-10<sup>+</sup> Bregs can express

## Table 1. Key effector molecules associated with Bregs

Breg gene	Protein	Protein type	Role associated with Bregs
CD1D	CD1d	Membrane	CD1d <sup>+</sup> marginal zone B cells are involved in the differentiation
CD274	PD-L1/CD274	Membrane	PD-L1 <sup>+</sup> Bregs can bind to PD-1, inhibiting effector T cell activity [25]
CD80	CD80/B7-2	Membrane	mAb-blockade of CD80 and CD86 in co-cultures of CD24 <sup>high</sup> CD38 <sup>high</sup> Bregs with CD4 <sup>+</sup> T cells reduced Breg
CD86	CD86/B7-1	Membrane	suppressive potency [8]. mAb-blockade of CD80 and CD86 in co-cultures of CD24 <sup>high</sup> CD38 <sup>high</sup> Bregs with CD4 <sup>+</sup> T cells reduced Breg
CR1	CR1/CD35	Membrane	Mucosal thromobospondin-1 producing CD35 <sup>+</sup> B cells promote FOXP3 <sup>+</sup> T cell expansion [26].
ENTPD1	NTPDase1/CD39	Membrane	CD39 expression was increased in CpG and IL-2 stimulated B cells within untreated rheumatoid arthritis patients and healthy controls. This also led to suppressed CD4 <sup>+</sup> and CD8 <sup>+</sup> T cell proliferation [27]. CD35 is an inhibitor of BCR-mediated human B cell activation and differentially regulates TLR7, and TLR9 induced responses [28].
GZMB	Granzyme B (GrB)	Secreted/intracellular	GrB produced from Bregs can cleave the ζ-chain from T cell receptor (TCR) inhibiting T cell activation and growth [29] as well as significantly reduce IFNγ and IL-17 expression in T cells [30].
HAVCR1	T-cell immunoglobulin and mucin domain 1 (TIM-1/HAVCR-1)	Membrane	There was significant co-expression of IL-10 and TIM-1 in transitional B cells after CpG and anti-BCR activation. However, the TIM-1 <sup>+</sup> B cells displayed stronger suppressive capabilities than transitional B cells [31]. Agonistic TIM-1 mAbs augment T cell-mediated immune responses, but antagonistic TIM-1 mAbs inhibits immune responses through regulatory B cells [32]. Ex vivo-expanded human CD19 <sup>+</sup> TIM-1 <sup>+</sup> regulatory B cells suppress immune responses in vivo and are dependent upon the TIM-1/STAT3 axis [17].
IDO1	Indoleamine 2,3- dioxygenase (IDO)	Intracellular	Over a few days, induced Bregs began to express increased IDO levels (through CTLA-4 action) which also led to increase in
IL10	Interleukin-10 (IL10)	Secreted/intracellular	The most common immunosuppressive cytokine secreted by Bregs, IL-10 supresses pro-inflammatory cytokines TNF- $\alpha$ and INE- $\alpha$ and divergence on the secret s
IL12A	Interleukin-12 subunit $\alpha$ (IL-12 $\alpha)$	Secreted/intracellular	Forms a heterodimer with IL-27 $\beta$ as the $\alpha$ subunit to create the regulatory cytokine IL-35 [35].
IL27B/EBI3	Interleukin-27 subunit $\beta$ (IL-27 $\beta$ )	Secreted/intracellular	Forms a heterodimer with IL-12 $\alpha$ as the $\beta$ subunit to create the regulatory cytoking IL-35 [35]
MME	Membrane metalloendopepti- dase (MME/CD10)	Membrane	An increase in CD10 <sup>+</sup> B cells was shown within uncomplicated malaria infection [36].
NT5E	Ecto-5-prime-nucleotidase (NT5E/CD73)	Membrane	Although not an inducible marker, IL-10 <sup>-</sup> B1 cells showed greatly decreased expression of CD73 compared to wild type B1 cells [37]. CD73 is an ecto-5'-nucleotidase that generates adenosine that induces GM-CSF/MDSC-mediated suppression of T cells 35001076 [38].
PDCD1	PD-1/CD279	Membrane	CD4 <sup>+</sup> and CD8 <sup>+</sup> T cells were shown to increase proliferation when an anti-PD-L1 mAb was applied to a coculture of PD-1 <sup>+</sup> Bregs with CD4 <sup>+</sup> and CD8 <sup>+</sup> T cells [39]
PLXNB2	Plexin B2 (PlxnB2)	Membrane	Sema 4C binding to PlxnB2 was found to be the cause of increased IL10 production to WT levels in Sema4C KO variants of murine B cells when co-cultured with Sema4C <sup>+</sup> HEK293 cells [40]
SDC1	Syndecan 1 (SDC1/CD138)	Membrane	CD138 <sup>+</sup> plasma cells showed increased expression of IL10 and IL-27 $\beta$ compared to the less mature variant CD138 <sup>int</sup> B cells suggesting regulatory functionality is gained through this maturation process [41]. CD138 suppresses apoptosis in multiple myeloma by activating IGE1 Recentor [42]
SEMA4C	Semaphorin 4C (Sema4C)	Membrane	Sema4C KO murine CD138 <sup>+</sup> B cells show expression of pro-inflammatory IL-4 cytokine as opposed to Sema4C <sup>+</sup> CD138 <sup>+</sup> B cells which expressed greater levels of IL-10 [40] Sema4C shows significantly greater expression in human tonsillar b cells when stimulated by agonistic anti-CD40 and IL4 [43].
TGFB1	Transforming growth factor β (TGF- β)	Secreted	In melanoma patients, greater numbers of TGF- $\beta^+$ B cells compared to healthy controls. Furthermore, these cells were responsible for increased mediation of T cells to undergo
THBS1	Thrombospondin-1 (TSP-1)	Secreted	Gifferentiation into FOXP3' 1 cells [44] THBS1 can bind to Latency Associated peptide (LAP), disrupting domain interactions and exposing its binding site thus converting LAP into its biologically active form TGF- β [45] Mesenchymal Stem Cells were found to induce IL-10 production in B cells via TSP-1 induced TGF-β production [46].

Table 1. (continued)					
Breg gene	Protein	Protein type	Role associated with Bregs		
TIGIT	TIGIT/VSIG9	Membrane	TIGIT <sup>+</sup> memory B cells were shown to directly target expression of IFN $\gamma$ and IL-17 in CD4 <sup>+</sup> T cells as well as suppress dendritic cell maturation resulting in downstream promotion of IL-10 <sup>+</sup> and suppression of IL-21 <sup>+</sup> and IL-4 <sup>+</sup> CD4 <sup>+</sup> T cells [30].		

memory or plasma B cell markers CD27, CD38, and CD148 [56, 63], or alternatively, they can be enriched in the naïve and transitional CD19<sup>+</sup>CD24<sup>hi</sup>CD38<sup>hi</sup> B cell subset found in peripheral blood [62].

An increase in IL-10<sup>+</sup> Bregs have been documented in the peripheral blood and tumours of patients with malignancy, including breast, gastric, and oesophageal cancers [10, 54, 65]. B10 cells have been shown to promote tumour growth, as tumour infiltrating CD27<sup>+</sup> CD10<sup>-</sup> Bregs inhibited inflammatory cytokine production by effector T cells in GC [53]. Depleting  $\text{IL-}10^+\ \text{B}$  cells can diminish pro-tumour immune responses by enhancing cytolytic abilities of splenic cytotoxic T lymphocyte, and B cells [66]. Elevated levels of IL-10<sup>+</sup> Bregs are also found in chronic Hepatitis B virus (HBV) and Human Immunodeficiency Virus (HIV) infection where they suppressed virus-specific CD8<sup>+</sup> T cells [67, 68]. In contrast, B cell depletion or deficiency in EAE mice exacerbated severity and prolonged duration of inflammatory phases, which was recoverable by adoptive transfer of IL-10<sup>+</sup> Bregs, and is implicative of a protective role in autoimmune conditions [16, 69]. A decrease in IL-10<sup>+</sup> Breg frequency and/or function has also been seen in patients with Rheumatoid Arthritis (RA), Systemic lupus erythematosus (SLE), and asthma [51, 70, 71]. Together, Bregs have been shown to modulate immune responses to cancer, infections, autoimmune diseases via production of IL-10. These different IL-10<sup>+</sup> Breg populations are represented across the spectrum of immunological tolerance in transplantation, autoimmunity, inflammation, cancer and infection. We are yet to identify a common transcription factor or signature of all IL-10<sup>+</sup> Bregs. This may be because there are multiple lineages, or indeed because this immunoregulatory function is multi-faceted and dependent on environmental factors.

#### IL-35-producing Bregs (IL-35<sup>+</sup> Bregs)

IL-35 belongs to the IL-12 cytokine family and consists of a  $\beta$ -chain subunit called EBI3 encoded by the IL27B gene, and an  $\alpha$ subunit called IL-12p35, encoded by IL12A gene [72]. IL-35 can regulate the activity of immune cells through the expansion of Tregs, Bregs and suppressing effector T cells, Th1 cells, and Th17 cells, as well as macrophages [41, 73]. The identification of IL-35producing Tregs that suppressed colitis in mice prompted researchers to explore the existence of IL-35-producing Bregs [43]. Immunoregulatory properties of IL-35<sup>+</sup> Bregs were noted when treating experimental autoimmune uveitis (EAU) mice with IL-35 resulting in the expansion of IL-10 and IL35-producing Bregs, and suppressed uveitis [35]. This was complemented by Dambuza et al.'s findings showing that the transfer of IL-35<sup>+</sup> Bregs induced an increase in Tregs and IL-10<sup>+</sup> Bregs, and inhibited proliferation of Th17 and Th1 cells, thereby suppressing EAU [74]. Subsequent studies in humans have shown that the frequency of IL-35<sup>+</sup> B cells is significantly decreased in SLE, which correlates with reduced plasma IL-35 levels. Further, an abundance of IL-35<sup>+</sup> Bregs is negatively associated with SLE severity in these patients [75]. A reduction in IL-35<sup>+</sup> B cell frequency and activity is also seen in ulcerative colitis (UC) [76]. IL-35<sup>+</sup> Bregs can limit inflammatory T-cell responses in patients in an IL-10dependent manner, whilst inhibiting anti-microbial immunity through the production of IL-35 [41, 72, 76] and have been shown to play a role in promoting pancreatic cancer [77, 78]. Although several studies have established the immunoregulatory properties of IL-35<sup>+</sup> Bregs, the mechanisms underlying the suppressive activity of IL-35 remains less understood.

#### TGF- $\beta^+$ producing Bregs (TGF- $\beta^+$ Bregs)

Transforming growth factor- $\beta$  (TGF- $\beta$ ) regulates immune responses in different diseases via mediating T cell differentiation, proliferation, and function. Several studies have established the inhibitory roles of TGF  $-\beta$  <sup>+</sup> Bregs in humans and mice [79]. These Bregs can be stimulated to secrete TGF- $\beta$  via multiple pathways, such as the classical signalling involving the TLR, BCR, and CD40 receptors, or growth factor-induced signalling (e.g. phosphatidylinositol-glycan biosynthesis class F protein (PIGF) induced differentiation of TGF-β secreting Bregs in glioma), subjective to the local environment of cells [18]. In mouse models of EAE, a B cell-specific deletion of TGF-β1, resulted in an earlier onset, greater disease burden, and inflammatory cytokine production, illustrating a role of TGF- $\beta^+$  Bregs in autoimmune diseases [80]. TGF-β secreting Bregs have also been shown to modulate immune tolerance in allergic diseases in humans [81]. mAb blocking experiments demonstrated that CpG-stimulated B cells which were initially isolated from human peripheral blood, were able to suppress CD4<sup>+</sup> T cell proliferation in a TGF-β-dependent mechanism, whilst promoting the differentiation of Tregs independent of IL-10 [33]. Similarly, TGF- $\beta$ 1<sup>+</sup> Bregs were found to suppress immune responses in GC patients by mediating the conversion of effector T cells into CD4<sup>+</sup> Foxp3<sup>+</sup> Tregs [34].

#### CD1d<sup>+</sup> Bregs

B cells present glycolipids to natural killer T (NKT) cells via CD1d, a non-polymorphic molecule, which subsequently secrete IFN- $\gamma$ and IL-4 to induce proliferation, maturation and antibody production by B cells [18]. Following presentation of lipid by CD1d to the invariant T cell receptor (iTCR), invariant NKT (iNKT) cells proliferate, produce cytokines and exert cytotoxic effects to regulate innate and adaptive immune responses [82]. NKT cells are reported to enhance antitumor immunity, protection against infections and regulation of autoimmunity [83]. Within human B cells, CD1d is predominantly expressed in Marginal Zone (MZ) -like B cells, IL-10<sup>+</sup> Bregs and, to a lesser extent in naïve and memory B cells. Early studies noted upregulated CD1d expression in splenic IL-10<sup>+</sup> B cell subsets in murine models, which correlated with colitis. Indeed, the majority of murine B10 Bregs express high levels of CD1d [84]. Further, CD1d<sup>+</sup> Bregs suppressed intestinal inflammation in mice with colitis, in an IL-10dependent manner [5]. Depletion of CD1d<sup>+</sup> Bregs exacerbated arthritis in mouse models and reduced responses to treatment with  $\alpha$ -galactosylceramide, the iNKT cell glycolipid agonist [24]. Oleinka et al. described a role for CD1d<sup>+</sup> Transitional 2-Marginal Zone Precursor (T2-MZP) Bregs in the differentiation of

suppressive iNKT cells that inhibited pro-inflammatory Th1 and Th17 responses in the context of experimental arthritis, partially via the production of interferon (IFN)- $\gamma$ . These mechanisms were IL-10-independent. Conversely, abnormalities in human and mice NKT cells have also been attributed to dysfunctions in CD1d<sup>+</sup> Bregs. Reductions in CD1d levels may potentially mediate the development of SLE through this mechanism [18, 85].

#### PD-1/PD-L1<sup>+</sup> Bregs

PD-1 is a transmembrane receptor that engages with ligands PD-L1, and PD-L2 to down-modulate immune cell responses. The significance of these interactions is highlighted by the frequent expression of these genes in DCs, NKT cells, B cells and T cells as well as the success of therapeutic approaches such as PD-1 inhibitors in treating cancers, and consequent autoimmune side effects in some patients [86]. PD-L1<sup>+</sup> Bregs were found to inhibit proliferation of CD8+ T cells, CD4+ CD25- T cells and CD49b+ NK cells in a murine model, via reduced IFN- $\gamma$  and TNF- $\alpha$  secretion, which ultimately promoted tumour growth in murine breast cancer models [87]. Further, expression of PD-1 and PD-L1 were found to confer immunosuppressive properties to different B cell subsets, in differentiated thyroid cancers (DTC), HCC, and other solid tumours [25, 39, 88]. Expansion of both PD-1<sup>+</sup> Bregs and PD-L1<sup>+</sup> Bregs has been found to coincide with advanced disease stage and progression in malignancies [25, 39]. Conversely, decreased expression of PD-L1 in Bregs have been associated with increased severity of allergic rhinitis and food allergies [89].

#### Granzyme B (GrB<sup>+</sup>) secreting Bregs

Bregs can modulate immune responses via secretion of cytokines and cytotoxic proteases such as GrB. IL-21 stimulation induced GrB<sup>+</sup> human Breg, which expressed the phenotype CD19<sup>+</sup> CD38<sup>+</sup>CD1d<sup>+</sup> IgM<sup>+</sup>CD147<sup>+</sup> together with upregulation of IL-10 expression. Analysis of multiple tumour types revealed tumour infiltrating GrB<sup>+</sup> Bregs adjacent to IL-21-secreting Treg, suggesting a mechanism within the tumour microenvironment [29]. Further, immunoregulatory properties of GrB<sup>+</sup> Bregs have been inferred due to altered frequencies in association with infections, vaccination, and transplantation. For instance, the peripheral blood of untreated HIV<sup>+</sup> patients demonstrated CD4<sup>+</sup> T cells with enhanced IL-21 expression and high in vivo frequencies of suppressive GrB<sup>+</sup> Bregs. When isolated from these patients, coculture of Il-21<sup>+</sup> CD40L<sup>-</sup> Th1 cells with B cells induced GrB<sup>+</sup> Bregs [90]. Interestingly, the supplementation of CD40L multimers to these co-cultures redirected B cell differentiation toward plasma cells, indicating that CD40L determined the fate of IL-21-dependent B cell differentiation to either regulatory or effector functions.

Similarly, increased frequencies of GrB<sup>+</sup> Bregs were identified in the peripheral blood of kidney transplant recipients who were operationally tolerant—that is, recipients who did not require pharmacological immunosuppression to maintain stable graft function. GrB expression by Bregs was again dependent on IL-21 stimulation by Th1 cells [91]. In contrast, decreased frequency and dysfunction of GrB<sup>+</sup> Bregs negatively correlated with disease severity in patients with Rheumatoid Arthritis (RA) [92]. Bregs can secrete GrB to regulate immune responses across a spectrum of immune-mediated disorders.

#### Tim-1<sup>+</sup> Bregs

Tim-1 is an type I membrane protein with an IgV domain followed by a heavily glycosylated mucin domain, a transmembrane domain and an intracellular cytoplasmic tail with one tyrosine phosphorylation motif [93]. TIM-1 is a T-cell costimulatory molecule which regulates CD4<sup>+</sup> T effector cell differentiation and responses in autoimmune and alloimmune settings [32, 94]. Whilst TIM-1 can function as a co-stimulatory molecule for T cell activation [95], it is also important in modulating the function of regulatory T cells (Tregs) [96, 97]. More recently, a role in B cell behaviour has been identified: ligation of TIM-1 induces IL-10 production by TIM-1<sup>+</sup> B cells and, in so doing, may promote immune tolerance [32].

Using mice which harboured a loss of function TIM-1 mutant, Kuchroo's group demonstrated that defects in TIM-1 signalling within B cells resulted in reduced IL-10 production by Bregs and severe multi-organ tissue inflammation. Tim-1-deficient B cells promoted Th1 and Th17 responses and inhibited the generation of Tregs, resulting in an increased severity of experimental autoimmune encephalomyelitis (EAE) [98]. Whilst the mechanisms underpinning human TIM-1<sup>+</sup> Bregs are less well characterized, studies have shown that TIM-1<sup>+</sup> B cells are more regulatory than TIM-1<sup>-</sup> B cells in an IL-10-dependent manner [99], and TIM-1 can serve as a marker for a human IL-10<sup>+</sup> Breg subpopulation which is partially overlapping with transitional B cells [31]. TIM-1+ Bregs have been shown to be reduced in frequency and altered in function in autoimmune diseases including systemic sclerosis [31] and T1D [100], but are enriched in cutaneous squamous cell carcinoma (SCC) [17]. Recent work has shown that ex vivo-expanded human CD19<sup>+</sup> TIM-1<sup>+</sup> Bregs suppress immune responses in vivo, they showed that by blocking TIM-1 activity they could significantly reduce the suppressive capability of the Bregs much more than when blocking IL-10 [17]. Although most mechanistic studies are performed in mice, studies have shown that targeting TIM-1 enhances type I interferon (IFN-I) responses, promotes B cells antigen presentation and activation, and enhances T cell anti-tumour responses, thus inhibiting tumour growth [101]. There is now a growing excitement around using TIM-1 as a potential target in immunotherapy [102].

# Functional roles of Bregs in human health and disease

As previously mentioned, immunoregulatory mechanisms of Bregs have been explored and elucidated in several cancers, infections, and autoimmune conditions. While the expansion of immunosuppressive Bregs promotes immune escape of tumour cells and increases severity of cancers, and infections; they are essential in limiting autoimmune conditions and ensuring transplant tolerance. Table 2 provides an overview of literature in human Bregs in autoimmune diseases, infections, and cancer, summarized in Fig. 2.

#### Bregs in auto-immune diseases

In the context of immune diseases, human Bregs were first discovered in SLE where CD19<sup>(+)</sup>CD24<sup>(hi)</sup>CD38<sup>(hi)</sup> B cells exhibit regulatory capacity in healthy individuals but are functionally impaired in SLE patients [8]. Later studies established the role of IL-10-producing Bregs in immune suppression and maintenance of tolerance, typically impaired in autoimmune diseases. The role of Bregs in autoimmune conditions such as SLE or RA is complicated by the variety of phenotypes, frequencies, and functional roles of the Breg populations described [103]. However, extensive validation within mouse models point to Bregs as an exciting therapeutic option for these autoimmune disorders.

An abundance of CD19<sup>+</sup> CD5<sup>+</sup>CD1d<sup>hi</sup> Bregs, and CD19<sup>+</sup>CD24<sup>hi</sup>CD38<sup>hi</sup> Bregs were observed in active SLE patients

Table 2. Phenotypic diversity of Bregs in autoimmune conditions, cancer and infections

Disease	Surface marker	Effector molecu	le Findings	Reference
Pancreatic Cancer (PC)	CD19 <sup>+</sup> IL-10 <sup>+</sup> Bregs	IL-10, IL-18	Elevated frequencies of CD19 <sup>+</sup> IL-10 <sup>+</sup> Bregs in PC patients versus. healthy controls, which correlated with tumour-node-metastasis (TNM) stage and poor survival. Increased plasma levels of pro-oncogenic IL-18 and	[110]
Ulcerative colitis (UC)	CD1d <sup>+</sup> IL-10 <sup>+</sup>	IL-10	Decreased II-18K expression by ILLO Bregs of PC patients. Decreased frequency in CD19 <sup>+</sup> CD1d+IL-10 <sup>+</sup> B cells in UC	[111]
Human schistosomiasis	CD1dhi CD24hi CD27+	IL-10, TGF-β	Increased II-10 production and a conversion of the production of the provided	[112]
Hepatitis C virus (HCV) related HCC	CD24+/hi CD38+/hi	IL-10, IL-35	Higher frequency of Bregs in HCC patients than controls, and positively correlation with disease stage, serum IL-10 and U-35 lavels	[113]
HCC	CD24 <sup>+</sup> CD38+	IL-10, TGF- <b>β</b> 1	Increased frequency of CD24 <sup>+</sup> CD38 <sup>+</sup> Bregs in tumour margin and blood coincide with cancer progression. Bregs induced proliferation and invasion of HCC cells via CD40-CD154 signalling	[114]
Acute Myeloid Leukaemia (AML)		_	Breg frequency was higher in PBMC and BMMC of AML patients, and coincided with advanced disease stage,	[115]
Invasive Breast Cancer		IL-10, PD-1	IL-10 <sup>+</sup> CD24 <sup>+</sup> CD38 <sup>+</sup> Bregs was elevated in peripheral blood and tumours of IBC patients compared to healthy controls. Peripheral CD19 <sup>+</sup> B cells from IBC patients but not healthy controls could induce CD4 <sup>+</sup> CD25+Foxp3+Tregs during co-culture in a PD-1-dependent manner	[116]
Lupus Nephritis (LN)	CD24hi CD38hi	IL-10	Decreased frequency of IL-10 <sup>+</sup> CD19 <sup>+</sup> CD24hiCD38hi Bregs	[51]
HIV	CD24hiCD27+, CD24hiCD38hi	IL-10	An increase in frequency of IL-10 <sup>+</sup> Bregs correlated with Tregs in untreated HIV patients which reduced	[68]
Non-Small Cell lung cancer (NSCLC)	CD24hiCD38hi	IL-10	Co-culture of Bregs with tumour-infiltrating follicular cytotoxic CD8 <sup>+</sup> T (Tfc) cells increased Tfc expression of IL-10 and reduced Tfc expression of pro-inflammatory	[117]
Gastric Cancer (GC)		IL-10	IFN- $\gamma$ , INF, and IL-2 in NSCLC patients. Increased % of peripheral IL-10 <sup>+</sup> CD19 <sup>+</sup> CD24hiCD38hi Bregs in GC patients compared to healthy cohort, which correlated with tumour stage. TIL-Breg frequency was enriched compared to peripheral blood in GC. Bregs inhibited secretion of IFN- $\gamma$ and TNF- $\alpha$ by CD4 <sup>+</sup> T cells, and induced CD4 <sup>+</sup> FORP3 - Tree differentiation	[34]
HBV		IL-10	In chronic HBV infection, IL-10 <sup>+</sup> Bregs frequency is in- creased in patients, and are temporally associated with peaks in viral load and hepatic inflammation. In vitro depletion of Bregs led to an increase in virus-specific CD <sup>8+</sup> T cells	[67]
HIV		IL-10	HIV <sup>+</sup> individuals have higher frequency of IL-10 <sup>+</sup> Bregs. Frequency of Bregs is positively associated with viral load and T cell exhaustion. Bregs impair proliferation of CD8+T cells in an IL-10-dependent manner, which can be rescued by Brog dopletion	[118]
Systemic Lupus Erythematosus (SLE)		IL-10	CD19 <sup>+</sup> CD24hiCD38hi Breg-mediated inhibition of T helper (Th1) cell differentiation is modulated by IL-10, CD80, and	[8]
Type 1 Diabetes (T1D)		IL-10	IL-10 production by Bregs is significantly reduced in	[119]
Rheumatoid Arthritis (RA)		Il-10	RA patients. RA patients have reduced number of CD19 <sup>+</sup> CD24hiCD38hi B cells compared to a healthy cohort. Bregs from RA patients failed to induce conversion of	[120]
Rheumatoid Arthritis (RA)	CD24hiCD38hi, CD24hi CD27 <sup>+</sup> , CD5 <sup>+</sup> , B10	IL-10	Decreased frequency of IL-10 <sup>+</sup> B cells in RA patients compared to a healthy cohort, with inverse correlation to disease severity. B10 cells from RA patients fail to induce differentiation of Trees.	[121]
Melanoma	CD27 <sup></sup> IgM <sup>+-</sup> IgD <sup>+-</sup>	PD-L1	Frequency of PD-L1 <sup>+</sup> Bregs was greater in metastatic melanoma patients, and positively correlated with disease stage. PD-L1+ naïve B cells inhibited effector T cell functions in tumours.	[25]

(continued)

Гable 2. (continued)					
Disease	Surface marker	Effector molecul	e Findings	Reference	
Gastric Cancer (GC)	CD27 <sup>+</sup> CD10 <sup>-</sup>	IL-10, CD80/86	IL-10 <sup>+</sup> CD27 <sup>+</sup> CD10 – Bregs were enriched in TIL-B cells, which reduced inflammatory cytokine production by CD4 <sup>+</sup> T cells and CD8 <sup>+</sup> T cells in an IL-10 and CD80/CD86- dependent manner	[53]	
COVID-19	CD27 <sup>+</sup> CD24hi	IL-10	IL-10 <sup>+</sup> Breg population frequency is diminished in patients with critical and severe COVID-19, and corresponds to a hyperinflammatory response marked by increase in clinical inflammatory parameters including neutrophil/ lymphocyte ratio. D-dimer presence etc in patients	[109]	
Differentiated Thyroid Cancer (DTC)	CD27 <sup>+</sup> IL-10+	IL-10	Enrichment of TIL-Bregs correlated with higher frequency of Foxp3+Tregs within tumours of DTC patients. Peripheral CD27 <sup>+</sup> Bregs from DTC patients produced more IL-10 than those from healthy cohort, and inhibited IFN-γ expression by CD4 <sup>+</sup> T cells, perforin and GrB expression by CD8 <sup>+</sup> T cells.	[122]	
Breast, ovarian, cervical, colorectal, prostate carcinomas	CD38 <sup>+</sup> CD1dhi IgM <sup>+</sup> CD147+	GrB	IL-21 induced GrBhi Breg is enriched in tumour tissue and promotes GrB-dependent degradation of TCR, inhibiting T cell proliferation.	[29]	
Head and Neck cancer	CD39 <sup>+</sup> CD73+	Adenosine (ADO)	Frequency of ADO producing Bregs was significantly decreased in tumour microenvironment compared to BBMC, but created an immunocurpressive environment	[123]	
Ulcerative colitis (UC)	CD5 <sup>+</sup> CD24hiCD38hi	IL-10	Lower number of CD5 <sup>+</sup> and CD24hiCD38hi Bregs and increased in CD95 <sup>+</sup> exhausted B cells in UC natients	[124]	
Allergic Asthma	CD5 <sup>+</sup> CD1d+	IL-10	Decreased frequency of CD5+and CD1d <sup>+</sup> CD5+transitional B cells in patients with asthma. Oral corticosteroids inhibited IL-10 production by these cells in patients.	[125]	
Oesophageal squamous cell carcinoma (ESCC)		IL-10	Increased frequency of IL-10 <sup>+</sup> Bregs in ESCC patients versus. healthy cohort. Exosomes isolated from peripheral blood of ESCC patients could induce differentiation of IL-10 <sup>+</sup> B cells and PD1 <sup>+</sup> Bregs.	[126]	
Cervical Cancer (CC)		IL-10	Frequency of CD19 <sup>+</sup> CD5 <sup>+</sup> CD1d <sup>+</sup> Bregs and serum IL-10 level was higher in CC patients. Bregs were strongly associated with disease progression and metastasis, but negatively correlated with CD8 <sup>+</sup> T cell frequency.	[127]	
Systemic Lupus Erythematosus (SLE)	CD5 <sup>+</sup> CD1dhigh	IL-10	Increased frequency of CD19 <sup>+</sup> IL-10 <sup>+</sup> CD5 <sup>+</sup> CD1dhigh Bregs in SLE, particularly during inflammatory phases. Frequency decreased during disease remission	[104]	
HCC	CD5hiCD24–/ +CD27hi/+CD38dim	PD-1	Higher frequency of PD-1hi Bregs exhibiting CD5hiCD24–/ +CD27hi/+CD38dim phenotype in HCC tissue. PD-1hi Bregs positively associated with disease progression and recurrence, and plasma IL-10 levels.	[88]	
Hepatocellular Carcinoma (HCC)	CD5hiCD24 – CD27–/ +CD38+/hi TIM-1+	TIM-1/HAVCR1	Proportion of TIM-1 <sup>+</sup> Bregs was elevated in tumours compared to peri-tumoural regions and blood of cancer patients, and positively correlated with disease progression, early recurrence, and invasion. TIM-1 <sup>+</sup> Breg frequency was negatively correlated to overall survival and limited proliferation and function of CD8 <sup>+</sup> effector T cells. TIM-1 <sup>+</sup> Bregs facilitated immune escape in HCC via novel exosomal HMGB1-TLR2/4-MAPK pathways.	[128]	
Rheumatoid Arthritis (RA)	PD-L1+, CD24hiCD38 - PD-L1+, CD24hiCD38hiPD- L1+	PD-L1	Decreased frequency of PD-L1 <sup>+</sup> Bregs in RA patients. % PD-L1 <sup>+</sup> Bregs increases in response to treatment. PD-L1 <sup>+</sup> Bregs suppress CD8 <sup>+</sup> T cells proliferation and cytokine production in vitro.	[129]	
HIV	TIM-1+	IL-10	IL-10 <sup>+</sup> Bregs increase in early HIV-1 infection in human and mice. Bregs suppress anti-HIV-specific T cell responses in an IL-10-dependent manner.	[130]	

in two separate studies [8, 104]. Contradictions on unchanged or reduced numbers of CD19<sup>+</sup>CD24<sup>hi</sup>CD38<sup>hi</sup> cells in SLE patients have also been noted. Blair *et al.* reported a significantly higher frequency of CD19<sup>+</sup>CD24<sup>hi</sup>CD38<sup>hi</sup> Bregs in PBMCs of patients, but a comparable absolute proportion. They proposed that a functional defect in CD19<sup>+</sup>CD24<sup>hi</sup>CD38<sup>hi</sup> B cells was involved in SLE pathogenesis [8]. This is supported by a refractory response to CD40 stimulation and reduced IL-10 production by CD19<sup>+</sup>CD24<sup>hi</sup>CD38<sup>hi</sup> B cells isolated from SLE patients [8]. The discrepancies in findings highlighted above limit our understanding of the mechanistic role of Bregs in SLE pathogenesis, potentially resulting from differences in cell stimulation approaches, IL-10 production levels of the cell population, and stage of diseases in samples assessed. It does not imply absence of the mechanistic role of Bregs in SLE [103]. As discussed in previous sections, Bregs can also modulate immune responses by IL-10-independent mechanisms. This is exemplified by Menon *et al.* [105] demonstrating the regulatory feedback mechanism between plasmacytoid DCs and Bregs via the release of IFN- $\alpha$  and CD40 engagement, which is altered in SLE leading to impaired Breg differentiation. Functional and/or numerical deficiencies in Bregs have also been found in patients with other autoimmune



**Figure 2.** Mechanisms of Breg responses in cancer, infection, autoimmune conditions and transplantation. In cancer and infections, an expansion of IL-10<sup>+</sup> Bregs and TGF- $\beta$  secreting Bregs mediate the conversion of effector T cells into Tregs and inhibit the activation of other effector T cell populations. In contrast, a reduction or dysfunction of Bregs have been shown in autoimmune conditions and transplantation responses, where IL-10<sup>+</sup> Bregs inhibit the secretion of proinflammatory cytokines by Th1, Th17, and monocytes. Further, Bregs inhibit expansion of T cells and promote differentiation of Tregs. IL-10<sup>+</sup> Bregs can also induce CD4<sup>+</sup> T cells to become Tregs. TGF- $\beta$ - and GrB-secreting Bregs inhibit proliferation of CD4<sup>+</sup> T and CD8<sup>+</sup> T cells. Bregs surface molecules including PD-1 which bind to T cells and limit their anti-tumoural responses. Created in BioRender. Wu, S. (2024) BioRender.com/s75v103.

diseases, including Type 1 Diabetes (T1D), Lupus nephritis (LN), ulcerative colitis (UC), detailed in Table 1.

#### **Bregs in infections**

Bregs not only limit immune responses in autoimmune and chronic inflammatory diseases but can also dampen immune responses to infection, consequently increasing disease severity. Phenotypically and functionally distinct Bregs have thus been characterized in bacterial, viral, and parasitic infections [106]. The frequency of CD19<sup>+</sup>CD24<sup>hi</sup>CD38<sup>hi</sup> Bregs significantly increased in HIV-infected individuals and correlated to active disease. Further, exposure to HIV-1 facilitates the differentiation of B cells into Bregs with an immunosuppressive repertoire of cytokines including high levels of IL-10, TGF-β1, EBI3 or IL-12 (p35) [107]. These cells were also capable of inhibiting effector functions of CD4<sup>+</sup> T cells and CD8<sup>+</sup> T cells. Similar expansions of IL-10-producing CD19<sup>+</sup>CD24<sup>hi</sup>CD38<sup>hi</sup> regulatory B cells in chronic HBV infections, and IL-10<sup>+</sup> neonatal Bregs in Respiratory Syncytial virus (RSV), demonstrate that IL-10 production by Bregs limits adaptive responses to viral infections [68, 108]. Conversely, a reduction in the frequency of IL-10<sup>+</sup> Bregs has been associated with hyperinflammatory responses in critical and severe COVID-19 patients, supported by an increase in clinical inflammatory parameters, including neutrophil/lymphocyte ratio and the D-dimer presence [109]. An expansion of IL10<sup>+</sup> Bregs were also associated with greater susceptibility to cerebral malaria, and Leishmania [36]. Multiple studies have demonstrated the role of Bregs in murine models of infectious diseases, the implications of which can be extended to gain insights on the mechanistic role of Bregs in human infections.

#### Bregs in cancer

Phenotypically varied Bregs have been shown to suppress immunity and promote tumour growth in several cancers. The expansion of IL-10<sup>+</sup>, TIM-1<sup>+</sup>, PD-1<sup>+</sup> and GrB<sup>+</sup> Bregs has been correlated with progressive disease stage and poorer prognosis in patients, highlighting the intricate functional roles of these cells in cancer pathogenesis, prognosis and outcomes.

An enrichment of IL10<sup>+</sup> CD19<sup>+</sup>CD24<sup>hi</sup>CD38<sup>hi</sup> Bregs promoting the conversion of effector T cells into Foxp3<sup>+</sup> Tregs in vitro was identified in breast and gastric cancer patients [34, 116]. IL-10<sup>+</sup> Bregs dampened the production of pro-inflammatory cytokines IFN-  $\gamma$ , TNF, and IL-17 by CD4<sup>+</sup> T cells and CD8<sup>+</sup> T cells [53, 131]. As described in earlier sections, B cells can also impair immune responses in cancer by secreting cytotoxic proteases such as GrB. Mechanistic exploration demonstrated that IL-21-mediated induction of GrB was dependent on both BCR and TLR signaling pathways, illustrating tight regulation of GrB<sup>+</sup> Breg induction. Elevated IL-21-induced GrB<sup>+</sup> Bregs have been found in the tumour microenvironment (TME) of breast, ovarian, cervical, colorectal, and prostate carcinomas [29]. GrB<sup>+</sup> Bregs limited T-cell proliferation by a GrB-dependent degradation of the T-cell receptor  $\zeta$ -chain, offering novel therapeutic strategies in cancer [29].

The PD-1/PD-L1 Breg axis also offers therapeutic possibilities in cancer treatments. An increased frequency of tumour infiltrating PD-1<sup>+</sup> Bregs was identified in the context of Hepatocellular carcinoma (HCC), which correlated with disease progression, poor survival, and recurrence in patients [88]. Stimulation of tumour-derived PD-1<sup>+</sup> Bregs with PD-1 Ligand (PD-L1) and subsequent co-culture with tumour-derived effector memory T cells, resulted in impaired proliferation and production of granzymes and perforin by CD8<sup>+</sup> T cells in an IL-10-dependent mechanism, thereby contributing to tumour growth. Similarly the frequency of circulating and tumour-derived PD-L1<sup>+</sup> Bregs correlated with disease stage in melanoma patients which were rare in healthy controls [25]. Naive circulating B cells with moderate PD-L1 expression isolated from patients with stage III and stage IV melanoma, could significantly suppress pro-inflammatory T cell responses in a PD-L1-dependent mechanism [25, 116]. Alongside inhibiting effector functions of T cells within the tumour microenvironment, Bregs can also induce differentiation of regulatory T cells (Treg) in a TGF- $\beta$ -dependent manner [79]. IL-10<sup>+</sup> Bregs have also been found to promote the suppressive effect of MDSCs and abrogate NK cell-mediated lysis of tumour cells in cancers such as Multiple Myeloma [25, 132, 133]. Overall, Bregs exert their capacity to exert immunosuppressive effects in cancer via multiple mechanisms.

#### Bregs in transplantation

B cells are thought to mediate acute and chronic rejection in transplant patients by producing *de novo* and donor-specific antibodies, by presenting antigens to alloreactive T cells and costimulating them. Conversely, Bregs are able to modulate the alloimmune response by multiple mechanisms including the secretion of immunosuppressive cytokines, to improve outcomes in experimental models [134]. Adoptive transfer studies have demonstrated that an array of Breg subsets can prolong allograft survival and promote transplant tolerance in an antigen-specific manner, often dependent on multiple mechanisms including IL-10, TIM-1, TGF $\beta$ , GrB and the induction of Tregs [134–137].

The rarity of human Bregs together with challenges in defining specific human Breg subsets and phenotypic discrepancies across species have made the clinical translation of these findings challenging. Depletion of B cells in kidney transplant recipients (KTRs) by Rituximab therapy accelerated rejection in transplant patients, first implicating regulatory B cell involvement in transplant tolerance [7]. Later studies showed that operationally tolerant KTRs (long-term stable graft function without the need for immunosuppressants) had increased frequencies of circulating Bregs compared to KTRs with stable graft function with immunosuppressants or those undergoing T cell mediated rejection (TCMR) [91]. The Immune Tolerance Network have reported similar findings in two USA and European-wide consortia studies, whilst also identifying B cell-based gene signatures specific to transplant tolerant patients [134, 138]. Cherukuri et al. has gone on to demonstrate that within circulating  $\text{CD24}^{\text{hi}}\text{CD38}^{\text{hi}}$ transitional B cells of KTRs, that the ratio of IL-10/TNF- $\alpha$  expression serves as an accurate predictor of 1-year rates of TCMR as well as graft function and graft survival at 5 years posttransplantation [139]. These observations not only suggest markers which could be used in immunosuppression minimization trials in transplantation, but also illustrate that perhaps the balance of Bregs to B effector subsets, rather than simply absolute frequencies, may be an important element in determining clinical outcomes [140]. Recent advances in human Breg expansion protocols have also demonstrated the therapeutic

possibilities of adoptive Breg cell therapy in a clinically relevant humanized mouse model of human skin transplantation [17]. Insights from such work may pave the way to translation for adoptive cell therapy in the clinic.

# Opportunities and outstanding challenges in the clinical applications of Bregs

The immunosuppressive properties of Bregs have been shown to play significant roles across autoimmune diseases, cancers, and infections, which highlights their potential in clinical settings. Here, we identify several such opportunities (Fig. 3).

#### Breg depletion in cancer

Emerging research suggests that Bregs might be manipulated to enhance cancer immunotherapy approaches by modulating the immune response to tumours. As discussed in earlier sections, increased Breg frequency is associated with diminished anti-tumoral immune responses in cancer patients. These observations offer fresh therapeutic possibilities in the form of novel targets for the depletion or inhibition of Bregs which may reactivate immune responses to cancer. Such strategies are already being explored. The inhibition of Mitogen/extracellular signal regulated kinase (MEK) using Cobimetinib in a pre-clinical in vivo colorectal cancer model, resulted in decreased Breg frequency within the draining lymph node, improved T cell infiltration and an enhanced response to anti-PD1 immunotherapy [141]. While Breg depletion by anti-CD20 antibodies has been explored to treat cancers, this approach inevitably has off-target effects when targeting a pan-B cell marker, thus resulting in concurrent depletion of anti-tumour B cell subsets [142]. Further studies investigating the depletion of IL-35-producing Bregs to treat infections and cancers are underway [72]. In addition, there is the potential for the use of Breg phenotypes as a biomarker of treatment response. For example, increase of IL-10<sup>+</sup> Bregs isolated from melanoma patients pre-treatment, positively correlated with a lack of response to anti-CTLA-4 treatment, indicating the potential to utilize IL-10<sup>+</sup> Bregs as a biomarker for treatment response [143]. Definitive clarity of Breg phenotypes will be required to develop precision-based targeting for effective, oncological therapies.

# Breg therapy in autoimmune diseases, inflammatory disorders, and transplantation

Debilitating autoimmune conditions such as EAE, chronic colitis, and CIA, can be experimentally ameliorated by adoptive transfer of Breg subsets in animal models, thus indicating a potential therapeutic opportunity [62]. Harnessing the regulatory functions of Bregs could lead to new therapeutic strategies for managing these inflammatory conditions and in the setting of transplantation, and potentially could be harnessed to control excessive inflammation in conditions like sepsis.

#### In vitro expansion of Bregs

Inducing Breg populations *in vitro* and subsequently transferring them into patients is an alternative strategy to suppress hyperinflammatory reactions in autoimmune diseases. However, this approach has several challenges including determining signals to induce Breg expansion, developing methods for their effective administration in patients, and ensuring their stability in host systems. Recently, protocols for *ex vivo* expansion and maintenance of human Bregs with *in vivo* immunoregulatory properties in humanized mouse models, have been developed [17]. This opens avenues to overcome the challenges of rare human Breg subsets and will facilitate the development of cell therapies in



#### **Outstanding challenges**

Figure 3. Opportunities and challenges of Bregs. Bregs could be targeted for depletion to augment anti-tumour immune responses. In vitro expansion of patient-derived Bregs and adoptive transfer into patients could suppress autoimmune or alloimmune responses in patients with autoimmune disorders or transplants. Boosting endogenous Breg populations via therapeutic agents could promote Breg expansion and induce tolerance in these patients. Created in BioRender. Wu, S. (2024) BioRender.com/a43i376.

transplantation and autoimmune conditions. The recent evolution of Chimeric Antigen Receptor (CAR) technology has revolutionized the T cell therapy landscape [144] and may also do the same for Breg cell therapies to ensure potency and to mitigate off-target effects. Future pre-clinical investigations are needed to determine concentration and specificity of stimuli to induce Bregs, the functional stability of a Breg cell therapy product and the appropriate therapeutic dosages required for clinical purposes [145].

#### Bregs as a prognostic predictor

Given the regulatory and immunosuppressive functions of Bregs suggesting a role in shaping disease outcomes, Bregs have emerged as potential prognostic predictors in various diseases. As described above, many studies have shown that Breg frequencies and functional properties correlate with disease severity and prognosis across multiple disease types. For example, the ratio of IL-10 to TNF $\alpha$  produced by transitional-1 B cells 3 months after transplantation was shown to be a predictive biomarker for clinical and subclinical renal allograft rejection and subsequent clinical course [140]. Investigating the presence and activity of Bregs as prognostic indicators holds promise in tailoring treatment strategies and predicting disease trajectories, contributing to a more personalized approach to patient care.

Challenges persist in standardizing Breg phenotyping and characterizing their specific roles in different disease contexts, necessitating further research and validation. There is no unique surface marker to distinguish them from other B cell subtypes. Bregs are heterogeneous, and their functions can vary. Understanding this diversity and developing approaches to harness specific Breg subtypes will be fundamental to harnessing their translational potential. The precise mechanisms by which Bregs regulate immune responses are not fully understood, making it difficult to develop targeted therapies. The enduring effects of manipulating Bregs and their long-term stability in the context of the wider immune milieu remain to be seen. Finally, developing standardized protocols for Breg isolation, expansion, and administration will be crucial to facilitate translation.

## Future directions and conclusion

B cells are well-characterized contributors to the adaptive and humoral immune responses. The discovery of B cells with immunosuppressive roles across a spectrum of immune-mediated diseases has unveiled novel insights into B cell behaviour in human health and disease. Studies conducted over the last few decades have significantly expanded our understanding of the immunobiology of Bregs and their role in the pathophysiology of cancers, autoimmune diseases, transplantation and infectious diseases. The therapeutic potential presented by these advances in Breg biology are currently limited by a lack of unique phenotypic signatures to identify Bregs. An incomplete understanding of the origin and differentiation of Bregs from B cells in response to pathogenic conditions further constrains efforts to design effective treatments. Given the growing significance of Bregs in health and diseases, the use of single cell multi-omics data (paired RNA-sequencing, B cell receptor sequencing and proteomics), human-centric functional analyses and standardized experimental conditions may present exciting opportunities by which to advance this field from bench to the bedside.

## **Author contributions**

Nawara Faiza Ahsan (Data curation [equal], Writing—original draft [lead], Writing—review & editing [equal]), Stella Lourenço (Data curation [equal], Writing—original draft [supporting], Writing—review & editing [supporting]), Dimitra Psyllou (Visualization [equal], Writing—original draft [supporting], Writing—review & editing [supporting]), Alex Long (Visualization [equal], Writing—original draft [supporting], Writing—review & editing [supporting]), Sushma Shankar (Writing—review & editing [equal]), and Rachael Bashford-Rogers (Conceptualization [lead], Project administration [lead], Visualization [supporting], Writing—original draft [lead], Writing—review & editing [lead])

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# Data availability

There are no new data associated with this article.

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Review