

The Regulating Function of Heterotrimeric G Proteins in the Immune System

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Abstract Heterotrimeric guanine nucleotide-binding proteins (G proteins), which consist of an α -, a β - and a γ -subunit, have crucial roles as molecular switches in the regulation of the downstream effector molecules of multiple G protein-coupled receptor signalling pathways, such as phospholipase C and adenylyl cyclase. According to the structural and functional similarities of their α -subunits, G proteins can be divided into four subfamilies: *G α s*, *G α i/o*, *G α q/11* and *G α 12/13*. Most of the α - and the $\beta\gamma$ -subunits are abundantly expressed on the surface of immune cells. Recent studies have demonstrated that G proteins are a group of important immunomodulatory factors that regulate the migration, activation, survival, proliferation, differentiation and cytokine secretion of immune cells. In this review, we summarise the recent findings on the functions of G proteins in immune regulation and autoimmunity.

Keywords G proteins · Immune regulation · Lymphocyte development · Autoimmune diseases

Introduction

Mammalian cells do not exist in isolation. Thousands of molecular messages need to be received by the receptors on

the cell surface, and the signals then need to be transmitted from outside to inside the cell through a chain of signalling molecules (Bockaert et al. 2002). Receptors can sometimes activate intracellular effectors more indirectly through a second messenger system, such as the G protein-mediated transmembrane signalling system. In this complex system, heterotrimeric G proteins function as critical molecular switches (Koelle 2006). The G proteins, together with G protein-coupled receptors (GPCRs), effectors and various regulators of G protein signalling (RGS), transduce the extracellular stimuli, such as hormones, chemokines, neurotransmitters, nucleotides, amino acids, sensory stimuli, biogenic amines and ions, into intercellular signals that ultimately convert these stimuli into the appropriate physiological response (Wettschureck and Offermanns 2005).

The heterotrimeric G protein consists of an α -subunit, which is a 37–42-kDa protein that contains a guanine nucleotide-binding pocket and intrinsic GTPase activity that binds and hydrolyses GTP, a 35-kDa β -subunit and an 8–11-kDa γ -subunit. The β - and γ -subunits form an indissociable complex, which is often referred to as a single entity, the $\beta\gamma$ -subunit (Entschladen et al. 2011). When a ligand interacts with a GPCR on the surface of the cell, the *G α* exchanges GTP for GDP and detaches from the *G $\beta\gamma$* dimer. Each of the activated components then interacts with different effectors, such as adenylyl cyclase, phosphodiesterase, phospholipase C (PLC), G protein-regulated kinase and voltage-dependent Ca^{2+} channel, to initiate various cellular responses. The α -subunit of the heterotrimeric G protein defines the basic properties of the G protein. Based on the differences in their α -subunits, G proteins can be divided into four subfamilies: *G α s*, *G α i/o*, *G α q/11* and *G α 12/13* (Wettschureck and Offermanns 2005). To date, only 17 different α -subunits, 5 different

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β -subunits and 12 different γ -subunits have been identified. Thus, a relatively small number of G proteins transmit the signals that are received by the huge number of different GPCRs (Goldsmith and Dhanasekaran 2007).

Due to its high complexity and versatility, the G protein-mediated signalling system has turned out to be the most widely used transmembrane signalling system in higher organisms (Wettschreck and Offermanns 2005). Numerous studies have focused on the function of G protein signalling in multiple tissues and organs. Some of these studies focused on the functions of this signalling pathway in the regulation of smooth and cardiac muscle contraction, platelet activation, synaptic transmission, and carbohydrate and lipid metabolism (Wettschreck and Offermanns 2005). A large number of GPCRs, including chemokine receptors, and most of the G protein subunits are widely expressed in various cell types of the immune system (Druey 2009). Through the use of knockout (KO) mice and chemical inhibitors, the functions of G proteins in the immune system have been recently studied. Increasing amounts of accumulated data have indicated that G protein signalling systems are key determinants in innate and adaptive immunity. These systems are involved in lymphocyte development, immune cell survival and migration, and tolerance induction (Table 1). In this review, we summarise the recent findings that elucidate the role of heterotrimeric G proteins in immune regulation and highlight the emerging roles of these proteins in autoimmune diseases.

G Proteins Are Required for Lymphocyte Development

The major causative factors that induce the onset and development of autoimmune diseases are a disruption in the normal development of the lymphoid tissues and an alteration in the function of lymphoid cells. The development of lymphocytes is regulated at many levels, and some G proteins have emerged as regulating factors in the commitment of precursor cells to the lymphoid lineage.

The initial study in $G\alpha_q$ -KO mice found that proportions of B cells, T cells, granulocytes, and monocytes from the peripheral blood, spleen, thymus, lymph nodes, Peyer's patches, and bone marrow are not significantly different from those observed in wild-type (WT) animals (Davignon et al. 2000). However, a further analysis of this subpopulation of immune cells in $G\alpha_q$ -KO mice revealed that $G\alpha_q$ can regulate the transition between pro- and pre-B cells in the bone marrow and the development of follicular and marginal zone B cells in the spleen. The transitional 1 (T1) B cells and marginal zone B (MZB) cells of $G\alpha_q$ -deficient mice were largely resistant to anti-IgM-induced cell death *in vitro*. The abnormal survival of B cells in the transitional

stage (from T1 to T2 B cells) can reduce the stringency of the negative selection of $G\alpha_q^{-/-}$ B cells and allow the entry of potentially autoreactive cells into the mature MZB compartment (Meyer-Bahlburg et al. 2008; Misra et al. 2010; Thien et al. 2004).

The deletion of $G\alpha_i2$ greatly augments the response of thymocytes to T-cell receptor (TCR)-mediated stimulation and results in an enhanced proliferation of double-positive (DP) thymocytes upon ligation of the TCRs. This altered response promotes positive selection and may also rescue some thymocytes from death due to their inability to bind MHC molecules (Elgbratt et al. 2007; Zhang et al. 2005). $G\alpha_i2^{-/-}$ mice exhibit a significantly reduced fraction of DP thymocytes and an increased fraction of single-positive (SP) thymocytes. The absence of the $G\alpha_i2$ protein also causes the arrest of thymocyte differentiation at the double-negative (DN) stage (Jin and Wu 2008). The reduced number of DP thymocytes was due to an accelerated transition from DP to SP thymocytes and a reduced transition from DN to DP thymocytes. Thus, the lower number of DP thymocytes resulted in a lower daily production by the thymus; the exact mechanisms behind the altered transition rates provided important information on thymic atrophy during colitis (Elgbratt et al. 2012). In addition, the null mutation of the $G\alpha_i2$ protein resulted in disordered B-cell subpopulations in the spleen and peritoneal cavity. $G\alpha_i2^{-/-}$ mice exhibited a reduced amount of MZB and T2 B cells and significantly increased numbers of follicular B cells. $G\alpha_i2$ was also important for the development and recruitment of B-1a and B-1b cells from the peritoneal cavity (Dalwadi et al. 2003).

The analysis of the cells that are generated from the thymus of these mutant mice revealed that $G\alpha_{13}$ - and not $G\alpha_{12}$ -mediated signalling plays an important role in the proliferation and survival of thymocytes during development and is required for early thymopoiesis (Coffield et al. 2004). Moreover, the $G\alpha_{12}/G\alpha_{13}$ family of G proteins can also regulate MZB cell homeostasis. Although the splenic follicular structure was roughly normal, further studies of the immune cell subpopulation revealed a strong reduction of MZB cell numbers in $G\alpha_{12}/G\alpha_{13}$ -double-knockout ($G\alpha_{12}/G\alpha_{13}$ -DKO) mice (Rieken et al. 2006).

G Proteins Control Lymphocyte Proliferation and Survival

Lymphocyte proliferation and death must be finely regulated to maintain immune homeostasis throughout the lifetime of a mammalian organism (Hildeman et al. 2007; Xu and Shi 2007). The clonal expansion of antigen-specific lymphocytes is required for effective immune responses against invading microorganisms. Shortly after the

Table 1 Signal transduction pathways and phenotype of mice deficient in G protein family members

| Family | Subtype | Effectors | Phenotype of knockout mice | References |
|----------------|----------------|--|--|--|
| <i>Gαs</i> | <i>Gαs</i> | AC↓, Ca ²⁺ influx↑ | <i>Gαs</i> ^{-/-} T cells: IL-17, IL-22 and IFN-γ production↓ | Li et al. (2012) |
| <i>Gαi/o</i> | <i>Gαi2</i> | AC↓, Erk↓?, Smad2↑, Smad3↑, PI3 K-Akt↑ | <i>Gαi2</i> ^{-/-} T cells: activation↑, proliferation↑, survival↓, polarised Th1 immune response, Tregs function↓, DP thymocytes↓, SP thymocytes↑, migration↓, IL-17 production↑ B cells: survival↓, Breg↓? Neutrophils: recruitment↓ Macrophages: recruitment↓, normal phagocytosis, TNF-α, IL-6, IL-1β and MIP-1α production↓ DCs: IL-12, IL-23 production↑, IL-10 production↓ | Elgbratt et al. (2012); Hornquist et al. (1997); Huang et al. (2003); Ngai et al. (2009); Padigel et al. (2007); Thompson et al. (2007); Wiege et al. (2012) |
| | <i>Gαi3</i> | AC↓, Akt↑ | <i>Gαi3</i> ^{-/-} T cells: migration↑ Macrophages: phagocytosis↓, GM-CSF production↑ | Fan et al. (2007); Jin et al. (2008b) |
| <i>Gαq/11</i> | <i>Gαq</i> | PLC-β↑, PLC-γ↓, PI3 K-Akt↓, Erk↓ | <i>Gαq</i> ^{-/-} T cells: proliferation↑, survival↑, activation↑, IL-2, IL-12 and TNF-α production↑ B cells: proliferation↑, survival↑, abnormal differentiation, activation↑ Neutrophils: migration↓ DCs: migration↓ | Misra et al. (2010); Molon et al. (2005); Ngai et al. (2008), (2009); Shi et al. (2007); Wang et al. (2011) |
| | <i>Gα11</i> | PLC-β↑ | <i>Gα11</i> ^{-/-} T cells: alternative TCR signalling pathway, substantial secretion of IL-2 | Bueno et al. (2006) |
| | <i>Gα15/16</i> | PLC-β↑ | <i>Gα16</i> ^{-/-} T cells: activation↓, IL-2 and IL-10 production↓, CD69 expression↓ | Zhou et al. (1998) |
| <i>Gα12/13</i> | <i>Gα12</i> | Rac1↓, Rap1↓, Btk↑ | <i>Gα12</i> ^{-/-} T cells: CD4 ⁺ T-cell activation↑, proliferation↑, CD4 ⁺ T-cell adhesiveness↑ B cells: MZB cells migration↑ | Herroeder et al. (2009); Rieken et al. (2006) |
| | <i>Gα13</i> | RhoGEF↑, RhoA↑ | <i>Gα13</i> ^{-/-} T cells: early thymopoiesis↓, migration↓ B cells: abnormal homing Macrophages: migration↓ | Coffield et al. (2004); Girkontaite et al. (2001); Murakami et al. (2004); Radu et al. (2004) |
| <i>Gβγ</i> | | AC↓, PLC-β↑, PI3 K↑ | <i>Gβ2</i> ^{-/-} Macrophages: migration↓ Neutrophils: impaired migration? | Hwang et al. (2004); Li et al. (2000) |

AC adenylyl cyclase, *GM-CSF* granulocyte–macrophage colony-stimulating factor, *PLC* phospholipase C, *Treg* regulatory T cell, *Breg* regulatory B cell, *RhoGEF* Rho guanine nucleotide exchange factor, *RhoA* Ras homolog gene family member A, *MIP-1α* macrophage inflammatory protein-1α, *Btk* Bruton's tyrosine kinase, *MZB cells* marginal zone B cells

pathogens are controlled, the expanded effector cells must be eliminated to prevent the non-adaptive accumulation of cells and ensure the return to immune homeostasis (Holzman et al. 2000). A fine balance between the survival and the death of lymphocytes ensures an effective immune system and maintains the immunological tolerance to self (Strasser and Bouillet 2003; von Boehmer and Melchers

2010). Although lymphocyte proliferation and death are not directly regulated by G proteins, the G protein-mediated signalling system might have important modulatory roles in these immune functions.

In recent years, studies with KO mice supported the pivotal role of the *Gαq*-mediated signalling pathway in the regulation of lymphocyte proliferation and survival (Misra

et al. 2010; Molon et al. 2005; Ngai et al. 2008, 2009; Wang et al. 2011). The G protein α q-subunit can directly inhibit PI3 K activation and prevent the activation of Akt. The PI3 K-Akt signalling pathway regulates many normal cellular processes, including cell proliferation, survival, growth, and motility processes (Browne et al. 2009; Cantrell 2002; Yanamadala et al. 2009). When chemokine receptors signal through $G\alpha_q$, the activated T cells showed increased proliferation and cytokine production (Molon et al. 2005). Because $G\alpha_q^{-/-}$ B cells are intrinsically defective and more resistant than WT B cells to cell death-inducing signals, such as B-cell-activating factor withdrawal and strong B-cell receptor (BCR) signals, mature $G\alpha_q^{-/-}$ B cells were more fit to survive than WT B cells (Misra et al. 2010). Moreover, the level of $G\alpha_q$ expression can contribute to the determination of the apoptosis and survival of human peripheral blood lymphocytes (PBLs) through the upregulation of Mcl-1 and the downregulation of caspase-3 activity (Wang et al. 2011). Consistent with the findings that have shown that $G\alpha_q$ is involved in human T lymphocyte regulation, the targeted mutation of RGS2 in mice also lead to reduced T-cell proliferation. RGS2 interacts with $G\alpha_q/11$ and accelerates the GTPase activity of the α -subunit, which negatively regulates G protein-coupled receptor signalling (Oliveira-Dos-Santos et al. 2000).

Despite the critical role of $G\alpha_q$ in lymphocyte proliferation and survival, studies with $G\alpha_i2$ -deficient mice have recently demonstrated that the induction of the $G\alpha_i2$ -mediated signalling pathways is sufficient to negatively regulate T-cell proliferation (Gotlind et al. 2011; Hornquist et al. 1997; Jiang et al. 1997; Zhang et al. 2005). $G\alpha_i2^{-/-}$ peripheral T cells display a hyperimmune response, which is characterised by an enhanced proliferation and production of inflammatory cytokines in response to stimulation with various mitogens (Hornquist et al. 1997; Huang et al. 2003). Interestingly, $G\alpha_i2^{-/-}$ T and B lymphocytes exhibited reduced expression of the anti-apoptotic intracellular protein Bcl-2 and significantly increased levels of apoptosis; the size and number of Peyer's patches were also reduced, likely due to the significantly increased T-cell production of interferon (IFN)- γ . Moreover, the upregulated Th1 cytokine production may be in response to the intestinal enteric flora (Ohman et al. 2002, 2005). In addition, an increased frequency of $CD4^+Foxp3^+$ regulatory T cells (Tregs) was observed in $G\alpha_i2^{-/-}$ mice, and this increased number of Tregs had no endogenous functional defect. However, the increased effective numbers of Tregs were unable to regulate the highly potent $G\alpha_i2^{-/-}$ effector T cells in vitro and in vivo. Therefore, these cells cannot prevent the development of autoimmune diseases, such as colitis (Gotlind et al. 2011), because the $G\alpha_i2^{-/-}$ T effector population comprises significantly higher number of cells

with a $CD4^+CD62L^-CD44^+$ effector memory phenotype than WT cells; these memory cells are more pro-inflammatory, more easily activated to proliferate and less susceptible to regulation by Tregs (Huang et al. 2003).

In addition to the $G\alpha_i/o$ and $G\alpha_q/11$ family members, the $G\alpha_{12/13}$ family of G proteins also has important regulatory roles in the T-cell life cycle (Coffield et al. 2004; Herroeder et al. 2009). Using $G\alpha_{13}$ and $G\alpha_{12}$ minigenes, the mutant could bind both $G\alpha_{12}$ and $G\alpha_{13}$ through the RGS domain and thereby prevent these receptors from transducing a signal to downstream effectors. Mice with T cells that exhibited genetically inactivated $G\alpha_{12}$ and $G\alpha_{13}$ have increased cell numbers, particularly $CD4^+$ T cells in the lymph nodes, blood, and the thymus. In addition, these $G\alpha_{12}$ - $G\alpha_{13}$ -double-deficient $CD4^+$ T cells showed an enhanced interaction with dendritic cells (DCs), which suggests that these G proteins have regulatory functions in $CD4^+$ T-cell activation, proliferation and apoptosis (Herroeder et al. 2009).

G Proteins Are Involved in Leukocyte Migration

The coordinated retention and relocation of leukocytes have key roles in the development, maintenance and proper functioning of the immune system, and a loss in the control of leukocyte traffic might contribute to immune suppression and autoimmune diseases (Jin et al. 2008a; Moser et al. 2004). It is now well recognised that chemokines and their receptors are master controllers of leukocyte migration, and studies over the past 30 years also reveal that chemokine receptors represent a new subfamily of GPCRs. These chemokine receptors activate the $G\alpha_s$, $G\alpha_i/o$, $G\alpha_q/11$ and $G\alpha_{12/13}$ families of G proteins depending on the type and activation state of the respective cell (Bennett et al. 2011; Kunkel and Butcher 2002; Wettschureck and Offermanns 2005; Zlotnik et al. 2011).

The role of $G\alpha_i$ in chemotaxis has been extensively investigated, and studies using KO mice have revealed that $G\alpha_i$ has a crucial role in leukocyte migration (Chaffin and Perlmutter 1991; Jin et al. 2008b; Spangrude et al. 1985; Thompson et al. 2007). The irreversible blocking of $G\alpha_i$ -mediated signalling by pertussis toxin (PTX) strongly impairs lymphocyte migration in vitro (Spangrude et al. 1985) and causes the accumulation of mature T cells in the thymus and greatly reduced level of T cells in the peripheral lymphatic organs in vivo (Chaffin and Perlmutter 1991). The PTX-induced blocking of $G\alpha_i$ -mediated signalling also causes defective homing of the peripheral lymphocytes to the spleen, lymph nodes, and Peyer's patches (Cyster and Goodnow 1995; Warnock et al. 1998). The results of a study that used $G\alpha_i2^{-/-}$ - and $G\alpha_i3^{-/-}$ -KO T cells indicate that $G\alpha_i2$ is indispensable for T-cell

migration and GTP γ S incorporation upon CXCR3-stimulation and that the G α i3- KO T cells display a significant increase in both GTP γ S incorporation and migration when stimulated with CXCR3 agonists. More importantly, the increased GTP γ S incorporation could be blocked by the G α i3 protein in a dose-dependent manner (Thompson et al. 2007). Similarly, during the development of graft-versus-host disease, the deletion of G α i2 hampered the trafficking of pathogenic T cells from the secondary lymphoid tissues to the inflammatory sites, and G α i2^{-/-} T cells displayed a defect in their response to CXCL10, CXCL11, and CCL5. In contrast, an aggravated rejection was induced in mice that were adoptively transferred with G α i3-deficient T cells, and the absence of G α i3 augmented the CXCL10- and CXCL11-induced chemotaxis of effector T cells and resulted in the homing preference of these cells to the liver and colon (Jin et al. 2008b). In neutrophils, G α i2 is required for chemokine-induced arrest in response to CXCL1, and G α i2^{-/-} mice show significant defects in neutrophil recruitment in lipopolysaccharide (LPS)-induced inflammatory models (Zarbock et al. 2007). Although a deficiency in the G α i2 protein does not affect neutrophil function, a significantly reduced recruitment of neutrophils into the microenvironment of the parasites in immunised G α i2^{-/-} mice is observed, and these signalling events are necessary for the recruitment of neutrophils that ultimately leads to the host-mediated killing of the larvae (Padigel et al. 2007). Furthermore, G α i2 and not G α i3 is essential for optimal CCL2- and C5a-induced recruitment of macrophages in acute inflammation (Wiege et al. 2012); however, not all effector functions of macrophages are mediated by G α i2-specific signalling, and G α i3 is able to act as a substitute for G α i2 in C5aR-regulated phagocytosis and G α i-dependent cytokine production (Fan et al. 2007). Interestingly, the G α -interacting vesicle-associated protein (GIV), which is a recently discovered non-receptor guanine nucleotide exchange factor (GEF), can also trigger G protein activation (Garcia-Marcos et al. 2009). The G α i3-GIV association is essential for macrophage chemotaxis through the enhancement of Akt activation and the remodelling of the actin cytoskeleton (Ghosh et al. 2008). Further studies are required to clarify the reciprocal function of G α i2 and G α i3 in leukocyte migration.

There is also evidence that the activation of G α q/11-mediated signalling is involved in chemokine-induced leukocyte migration (Al-Aoukaty et al. 1998; Shi et al. 2007; Soede et al. 2001). Macrophage inflammatory protein-3 α receptors can couple to G α q/11 proteins to enhance a robust calcium response flux and induce the motility of interleukin (IL)-2-activated natural killer cells (Al-Aoukaty et al. 1998). In myeloid cells, G α q/11 is required for integrin activation in the bone marrow and induces the integrin LFA-1-dependent aggregation of TAM2D2 T-cell

hybridoma cells (Soede et al. 2001). More importantly, G α q-deficient mice show defective calcium and chemotactic responses upon stimulation of neutrophils with N-formylmethionine leucyl-phenylalanine (fMLP) and CCL3 and upon stimulation of DCs with CCL2, CCL19, CCL21, and CXCL12. In contrast, the G α q-null T-cell responses to CXCL12 and CCL19 are not affected (Shi et al. 2007). Thus, an alternative G α q-dependent chemokine receptor pathway may control the migration of only a subset of leukocytes (Shi et al. 2007). The novel chemokine receptor signalling pathway appears to be critically important for the initiation of inflammatory responses because G α q is required for the migration of DCs from the skin to the draining lymph nodes after fluorescein isothiocyanate sensitisation and for the emigration of monocytes from the bone marrow to the inflamed skin after contact sensitisation (Shi et al. 2007). In addition, the macrophages of G α 15- but not G α 11- and G α q-deficient mice exhibit only a minor signalling defect in response to complement C5a (Davignon et al. 2000). G α 16, which is the human counterpart of G α 15, can also efficiently couple chemoattractant receptors to NF- κ B activation, which suggests a potential function of G α 16 beyond the activation of PLC β (Yang et al. 2001). However, contradictory results have been obtained for the roles of G α q and G α i2 in the regulation of T-cell migration. A study indicated that the knockdown of G α q resulted in an approximately 80 % increase in CXCL12-induced T-cell migration, whereas the knockdown of G α i2 inhibited CXCL12-induced T-cell migration (Ngai et al. 2008, 2009). Therefore, further studies are required to clarify the roles of G α q and G α i2 in the regulation of T-cell migration.

Due to the embryonic lethality of G α 12/G α 13-DKO, direct evidence for the involvement of the G α 12/13 subfamily in lymphocyte adhesion and migration is not yet available (Gu et al. 2002). However, the G α 12/13 effector RhoA has been repeatedly shown to be involved in lymphocyte traffic. The murine Rho-specific GEF Lsc can couple G α 13 to activate RhoA. The B cells of Lsc-deficient mice exhibit impaired actin polymerisation and motility and abnormal homing (Girkontaite et al. 2001). In addition, defective migration is also observed in mice that lack the orphan G protein-coupled receptor G2A, which may be directly coupled to G α 13. G2A-deficient macrophages and T cells exhibit reduced migration towards lysophosphatidylcholine (LPC), whereas G2A overexpression in a macrophage cell line enhances the migration of these cells towards LPC (Murakami et al. 2004; Radu et al. 2004). T- or B-cell-specific G α 12/G α 13-DKO mice were generated using the Cre-loxP system. The resultant G α 12/G α 13-DKO MZB cells exhibit significantly increased migration towards sphingosine 1-phosphate, calf serum, and mouse serum, whereas the serum-induced migration was not

altered in the follicular B cells (Rieken et al. 2006). Similarly, $G\alpha 12/G\alpha 13$ -DKO $CD4^+$ T cells also exhibit increased adhesiveness and enhanced lymph node entry (Herroeder et al. 2009).

Neutrophil polarisation and directed migration can also be mediated by the G protein $\beta\gamma$ -subunits (Neptune et al. 1999). The complement C5a-induced migration of J774A.1 mouse macrophages critically depends on $G\beta 2$, but not on $G\beta 1$, $G\alpha i 2$, or $G\alpha i 3$ (Hwang et al. 2004). PI3 $K\gamma$, PLC $\beta 2$ and PLC $\beta 3$ are intracellular effectors of the dissociated $G\beta\gamma$ -subunits in neutrophils. Although neutrophils from mice lacking PLC $\beta 2$ or PLC $\beta 3$ exhibit normal and even enhanced chemotactic responses, the migration of PI3 $K\gamma$ -deficient neutrophils is severely impaired, and these cells fail to accumulate at the sites of inflammation in a septic peritonitis model (Li et al. 2000).

Cytokine Production Is Regulated by G Proteins

In general, the physiologic function of all immune responses is to eliminate microbes and other foreign antigens. It has recently become obvious that GPCRs and heterotrimeric G proteins can trigger different aspects of innate and adaptive immunity and exert important modulatory effects on the immune cell effector functions, including cytokine production, lymphocyte differentiation, phagocytosis, and mediator release (Wettschreck and Offermanns 2005).

The treatment of normal mice with PTX inhibited $G\alpha i$ protein signalling and thus mimics the conditions that are observed in $G\alpha i 2$ -deficient mice (Ryan et al. 1998). Various effects of PTX on the immune system have been reported, including a polarised Th1-type immune response, an enhanced capacity of splenocytes to produce IL-12, tumour necrosis factor (TNF)- α , IFN- γ and IL-2 in response to both microbial and non-microbial stimuli, and an augmented expression of co-stimulatory molecules, such as B7-1 and B7-2 on macrophages and their counter-receptor CD28 on T cells (Shive et al. 2000). These findings are further confirmed by the fact that untreated $G\alpha i 2^{-/-}$ mice also exhibit enhanced production of IL-12 and TNF- α by splenocytes and of IL-12 p40 by purified splenic $CD8\alpha^+$ lymphoid DCs (He et al. 2000). Moreover, T cells from $G\alpha i 2^{-/-}$ but not $G\alpha i 3^{-/-}$ mice exhibit hyperresponsiveness in their Th1-type cytokine production after their activation through the TCR, and $G\alpha i 2^{-/-}$ T cells have a relaxed co-stimulatory requirement for IL-2 secretion and proliferation compared to WT cells (Huang et al. 2003). In addition to the induction of Th1 responses, recent studies found that the loss of $G\alpha i 2$ can also promote a Th17 phenotype with significantly greater levels of IL-17, whereas $CD11c^+$ bone marrow-derived DCs produce

higher levels of the inflammatory cytokine IL-23 and a minimal IL-10 response to CpG (Pena et al. 2009). More importantly, the B-cell population of $G\alpha i 2^{-/-}$ mice is functionally deficient in LPS-induced proliferation and IL-10 production, which indicates that $G\alpha i 2$ is required for the development of IL-10-producing B cells (the potential regulatory B cells) (Dalwadi et al. 2003). These immune function abnormalities in $G\alpha i 2^{-/-}$ mice might contribute to lymphocyte-mediated autoimmunity in these mice.

The requirement of $G\alpha q/11$ -mediated signalling for T-cell activation was first demonstrated in RGS2-deficient mice. In addition to a proliferative defect, the RGS2 $^{-/-}$ T cells produced significantly lower levels of the T-cell growth factor IL-2 and induced an impaired antiviral immunity in vivo (Oliveira-Dos-Santos et al. 2000). The activation of the murine thromboxane A2 receptor, which is typically coupled to the $G\alpha q/11$ family of G proteins, impairs the DC-T cell adhesion and inhibits the DC-dependent proliferation of T cells (Kabashima et al. 2003). The abovementioned indirect evidence indicates that $G\alpha q/11$ might be a negative regulator of acquired immunity. Recent studies with KO mice further support the involvement of $G\alpha q/11$ in immune cell effector functions (Bueno et al. 2006; Misra et al. 2010; Ngai et al. 2008). Bacterial superantigens can bypass Lck-dependent TCR signalling through the activation of a $G\alpha 11$ -dependent PLC β -mediated pathway. This alternative signalling pathway leads to the activation of Erk and protein kinase C and to the influx of Ca^{2+} and induces a substantial secretion of IL-2 (Bueno et al. 2006). Similarly, the siRNA targeting of $G\alpha q$ demonstrates a specific role of $G\alpha q$ in TCR signalling. $G\alpha q$ -deficient Jurkat TAg T cells display a reduced activation of Lck but paradoxically show sustained Erk1/2 phosphorylation. Consistent with this finding, $G\alpha q^{-/-}$ primary T cells also show reduced proximal LAT phosphorylation, sustained Erk1/2 phosphorylation and augmented immune responses, including increased secretion of IL-2, IL-5, IL-12 and TNF- α (Ngai et al. 2008). Furthermore, upon anti-IgM stimulation, the levels of phospho-Akt, phospho-PLC $\gamma 2$ and phospho-Erk are significantly increased in $G\alpha q^{-/-}$ B cells, which suggests that there is an increased activation of BCR-mediated signalling in $G\alpha q^{-/-}$ B cells (Misra et al. 2010). In contrast, mutant $G\alpha 16$ transfectants display an inhibition of TCR/CD3-mediated signalling and deficiencies in IL-2 and IL-10 production and CD69 expression (Zhou et al. 1998).

In contrast, the inactivation of $G\alpha s$ -mediated signalling exhibits an immunosuppressive effect. $CD4^+$ T cells in $G\alpha s$ -conditional-KO mice exhibit decreased production of cAMP, reduced Ca^{2+} influx, lower secretion of IL-17, IL-22 and IFN- γ , and normal IL-4 production and cannot mount an antigen-specific Th17 response upon oral CT/OVA immunisation. Due to the selective modulation of

Th17 and Th1 differentiation, the adoptive transfer of naive $G\alpha s^{-/-}$ $CD4^{+}$ T cells into $RAG1^{-/-}$ recipients provokes minimal colonic inflammation, and the mice ultimately fail to develop colitis (Li et al. 2012). The two members of the $G\alpha 12$ family, $G\alpha 12$ and $G\alpha 13$, were ubiquitously expressed and have been reported to mediate actin polymerisation during T-cell activation (Offermanns et al. 1997). The activation of RhoA by $G\alpha 12$ and $G\alpha 13$ is mediated by a subgroup of GEFs for Rho, and both RhoGEF and RhoA have been shown to affect T-cell function (Galandrini et al. 1997; Girkontaite et al. 2001). The indirect evidence suggests that $G\alpha 12$ -mediated signalling might play an important regulatory role in T-cell activation.

G Proteins and Autoimmune Diseases

$G\alpha i 2$ -deficient mice exhibit a pro-inflammatory phenotype and can develop severe immunological problems, such as fatal inflammatory bowel disease (IBD) (Rudolph et al. 1995). IBD is a collective term that refers to chronic, autoimmune, and inflammatory diseases of the bowel, mainly ulcerative colitis and Crohn's disease (Baumgart and Carding 2007). The $G\alpha i 2^{-/-}$ mouse is a well-established model for colitis and is characterised by a Th1 $CD4^{+}$ T-cell response with an increased production of $IFN-\gamma$ and increased levels of IL-1, IL-6 and TNF- α in the inflamed tissue (Hornquist et al. 1997; Ohman et al. 2002). The absence of $G\alpha i 2$ also results in defective arachidonate release and prostaglandin E_2 production in intestinal and

colonic subepithelial myofibroblasts and dysregulation of the epithelial cell barrier (Edwards and Smock 2006; Edwards et al. 2008; Saha et al. 1998). In addition, the loss of $G\alpha i 2$ results in elevated IL-12 levels and insufficient IL-10 secretion by macrophages and splenic $CD8\alpha^{+}$ DCs (He et al. 2000; Pena et al. 2009). Furthermore, the null mutation of $G\alpha i 2$ can cause a reduction in the MZB and peritoneal B-1b cell subpopulations (Dalwadi et al. 2003) and Treg cell non-function (Gotlind et al. 2011). These facts indicate the multiple requirements for $G\alpha i 2$ in mucosal immune responses. $G\alpha i 2$ -deficient mice can spontaneously develop colitis by the age of 16–21 weeks, and 30–40 % of the animals develop nonpolypoid adenocarcinoma, which is a typical character of human ulcerative colitis (Rudolph et al. 1995). Genetic differences between mouse strains might affect the susceptibility of $G\alpha i 2^{-/-}$ mice to colitis: $G\alpha i 2^{-/-}$ C57BL/6 mice were relatively resistant to colitis, whereas $G\alpha i 2^{-/-}$ 129/Sv mice developed IBD earlier and with greater frequency and severity (Bjursten et al. 2004). Genetic linkage studies in humans have mapped the $G\alpha i 2$ gene within an IBD-susceptible locus at chromosome 3p21, which raises the possibility that $G\alpha i 2$ is one of the candidate genes that contribute to IBD development in humans (Fig. 1) (Hampe et al. 2001). Remarkably, genetically engineered mice raised in germ-free conditions on a 129/Sv strain background appear to be less susceptible to IBD-like diseases compared with mice housed in SPF conditions, and germ-free studies have also revealed the key role of the microbiota in the development of colitis (Arthur and Jobin 2011; Taurog et al. 1994).

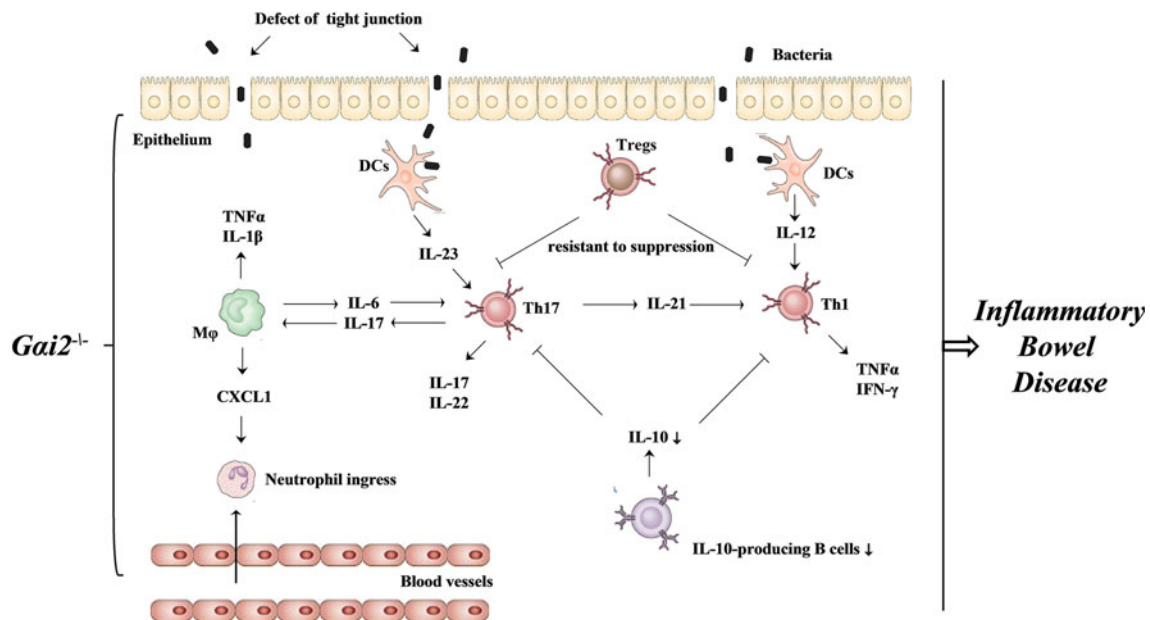


Fig. 1 Possible immune pathogenesis of $G\alpha i 2^{-/-}$ colitis. The loss of $G\alpha i 2$ can lead to an amplification of Th17/Th1 responses, which are likely mediated by a defect in the epithelium tight junctions. IL-17 is

fed back to the macrophages, which then upregulate CXCL1 secretion, and induces neutrophil ingress. The lack of appropriate regulation by Tregs further exacerbates the $G\alpha i 2^{-/-}$ colitis

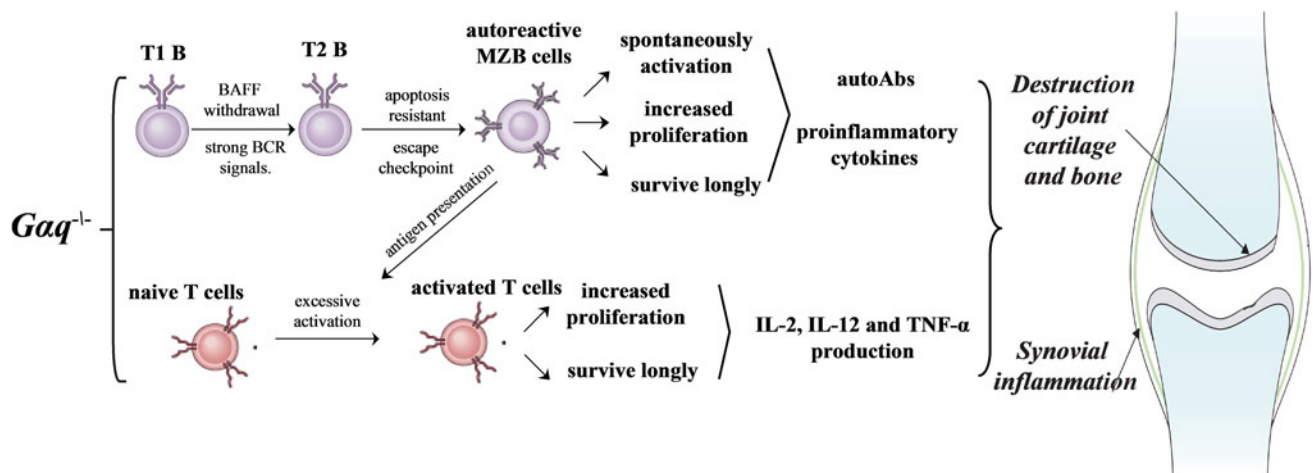


Fig. 2 Possible immune pathogenesis of $G\alpha q^{-/-}$ arthritis. $G\alpha q$ has a critical intrinsic role in the maintenance of the immunological tolerance of peripheral lymphocytes. $G\alpha q$ modulates the peripheral B-cell development and appears to control the numbers of T1-derived

MZB cell precursors and mature MZB cells. $G\alpha q^{-/-}$ T and B cells are more likely to survive than WT lymphocytes and exhibit significant proliferation. These abnormal $G\alpha q^{-/-}$ lymphocytes might contribute to the development of arthritis

These findings have led to an intense interest in the interactions between the intrinsic (genetic) host defects and extrinsic (microbiota) factors. One study has shown that an ecological disorder of the bacterial community, which is also named “dysbiotic microbiota”, can cause IBD and that the symptoms are more severe in genetic animals models (Neish 2009). Consistently, an antibiotic treatment can also induce remission in IBD patients (Khan et al. 2011). Furthermore, the autoantibodies detected in IBD models are very likely the result of cross-reactivity with bacterial antigens (Targan and Karp 2005).

A previous study with $G\alpha q^{-/-}$ chimeric mice, which were generated through the reconstitution of lethally irradiated C57BL/6 J recipient mice with $G\alpha q^{-/-}$ bone marrow, demonstrated that $G\alpha q^{-/-}$ chimeras could spontaneously develop systemic autoimmunity with multi-organ involvement, including the production of autoreactive antibodies by $G\alpha q^{-/-}$ B cells, the deposition of IgG2a- and IgG2c-containing immune complexes in the kidney, thrombotic microangiopathy, the reduction in the numbers of red blood cells, synovitis, bone resorption, exostotic bone development, and osteolytic activity (Misra et al. 2010). These pathological features may be manifestations of diseases in Lupus or arthritis patients (Misra et al. 2010). Because there is over 99 % identity between the human and mouse $Gnaq$ genes, $G\alpha q$ might also play a role in the pathogenesis of human rheumatoid arthritis (RA). We recently showed that the expression of $G\alpha q$ in the PBLs from RA patients is significantly decreased and that the expression level of $G\alpha q$ mRNA in PBLs from RA patients strongly correlates with RA disease activity (DAS28), anti-cyclic citrullinated protein antibodies, C-reactive protein and rheumatoid factor (Elgbratt et al. 2012). Our results

support the hypothesis that $G\alpha q$ may be involved in the pathogenesis and progression of RA through the regulation of the apoptosis and survival of PBLs (Fig. 2).

The data obtained using $G2A^{-/-}$ mice present indirect evidence that supports the hypothesis that the inactivation of $G\alpha 13$ may be involved in the development of autoimmune-like syndrome. Older $G2A^{-/-}$ mice present multiple signs of autoimmunity with heavy lymphocytic infiltration in various organs, increased immunoglobulin levels, the deposition of immunoglobulin complexes in the glomeruli and the presence of anti-nuclear autoantibodies. Together with the emergence of lymphadenopathy in ageing $G2A^{-/-}$ mice, these observations suggest the development of a late-onset non-organ-specific autoimmune syndrome (Hampe et al. 2001). Similarly, T-cell-specific $G\alpha 12/G\alpha 13$ -DKO mice also exhibit increased $CD4^{+}$ T-cell proliferation and infiltration, enhanced proinflammatory cytokine production and the deposition of IgG-containing immune complexes in the kidney (Herroeder et al. 2009). These facts indicate that $G\alpha 13^{-/-}$ mice may be associated with increased susceptibility and severity of T-cell-mediated abnormal immune responses and autoimmune disease (Hampe et al. 2001; Herroeder et al. 2009).

Concluding Remarks

The past several years have seen an explosion of data that uncover the roles of heterotrimeric G proteins signalling in the regulation of immune cell function and in the development of diverse autoimmune-like syndromes. Many of these insights were derived from studies that used gene-targeted mice. An important point has to be taken into

consideration: chemokines are a superfamily that contains 48 ligands that bind to 19 different GPCRs. In addition to their best recognised function, which is the induction of immune cell migration, chemokines have many functions, including the regulation of immune cell survival and effector responses. With the availability of KO mice, the significance of many aspects of G protein-mediated signalling in the innate and adaptive immune response has been studied, including the roles of G proteins in autoimmunity. It is worth paying more attention to the function of various G proteins in the development and progression of human autoimmune diseases by studying the relationship between G proteins and human autoimmune diseases and verifying the internal links epidemiologically. Due to the functional diversity of G proteins in the immune system, the study of the crosstalk between multiple subforms of G proteins-mediated signalling, such as $G\alpha_q$ and $G\alpha_{11}$, will likely lead to new insights that will help elucidate their function in the maintenance of immune homeostasis. In conclusion, the expression of G proteins in immune cells, such as granulocytes and lymphocytes, represents an important regulatory component of the intracellular signalling pathways that are induced by GPCRs. Thus, the molecular elucidation of the functions of G proteins in immunity might pave the way for the development of novel therapeutics for the treatment of autoimmune diseases, and the modulation of G protein signalling downstream of the receptor might lead to the development of novel drugs with better efficacy and/or fewer side effects.

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