

$\gamma\delta$ T-cell function in sepsis is modulated by C5a receptor signalling

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Introduction

$\gamma\delta$ T cells are involved in a wide variety of disease processes and play important roles in both innate and adaptive immunity. However, the underlying mechanisms, particularly the molecules involved and the biochemical bases, are still unclear.¹ Recently, data have been obtained suggesting that $\gamma\delta$ T cells act as early responders that promote inflammation through the production of interleukin-17 (IL-17).^{2,3} $\gamma\delta$ T cells can also act as antigen-presenting cells to initiate immunity, hence acting as a bridge between innate and adaptive immunity.⁴ However, in most cases, the mechanisms used by $\gamma\delta$ T cells and the identity of the molecules involved and the cell surface markers with which they interact have not been fully determined.¹

The complement system is part of the innate immune network and is composed of a sophisticated network of

Summary

We previously showed that $\gamma\delta$ T cells are involved in the pathogenesis of sepsis, but, the underlying mechanisms remained unclear. The present study demonstrates, for the first time, that $\gamma\delta$ T cells express the complement C5a receptor (C5aR, CD88) and that CD88 expression in $\gamma\delta$ T cells was up-regulated in mice following sepsis both at protein and mRNA levels. Complement C5a itself contributed to the regulation of C5aR expression on $\gamma\delta$ T cells, as (i) neutralization of C5a *in vivo* prevented the expression of C5aR on $\gamma\delta$ T cells in septic mice and (ii) incubation of mouse spleen cells or purified $\gamma\delta$ T cells with recombinant C5a *in vitro* increased CD88 expression by $\gamma\delta$ T cells at both protein and mRNA levels. C5a receptor on $\gamma\delta$ T cells also mediates increased interleukin-17 (IL-17) expression as incubation of mouse spleen cells or purified $\gamma\delta$ T cells with recombinant C5a promotes the IL-17 expression by $\gamma\delta$ T cells. Ligation of the C5aR on $\gamma\delta$ T cells activated the phosphoinositide 3-kinase (PI3K)/Akt signalling pathway, which enhances CD88 expression and promotes IL-17 secretion. These results demonstrate that C5a acts directly on the C5aR expressed on $\gamma\delta$ T cells, resulting in cell activation, and subsequently enhances their capacity for IL-17 production. The up-regulation of the PI3K/Akt pathway following C5a stimulation contributes to up-regulation of $\gamma\delta$ T-cell function.

Keywords: C5a; CD88; sepsis; $\gamma\delta$ T cells

soluble and membrane-bound proteins.⁵ Complement C5a is an important soluble pro-inflammatory mediator and can activate innate immune cells after binding to its receptor expressed on neutrophils or dendritic cells.^{6,7} In addition to its role in the innate immune response, C5a has recently been shown to play an important role in the regulation of T-cell immunity.^{8–14} For example, Strainic *et al.*⁸ found that the locally produced complement fragments C5a and C3a provide both co-stimulatory and survival signals to naive CD4⁺ T cells, and Kim *et al.*¹¹ found that the C5a receptor (C5aR) is essential for optimal generation of anti-viral CD8⁺ T-cell responses. Although it is clear that C5a is actively involved in the regulation of both innate and adaptive immunity, its roles in the regulation of $\gamma\delta$ T cells, important mediators of both innate and adaptive immunity, are largely unclear.

We previously found that $\gamma\delta$ T cells play pathogenic roles in the progression of sepsis in mice and that their

Abbreviations: C5aR, complement C5a receptor; CD62L, CD62 ligand; CLP, caecal ligation and puncture; IL-17, interleukin-17; NF- κ B, nuclear factor-k-gene binding; PE, phycoerythrin; PI3K/Akt, phosphoinositide 3-kinase/Akt.

activity can be affected by excessive production of C5a.¹⁵ However, the underlying mechanisms remain unclear. Whether C5a is directly involved in the regulation of $\gamma\delta$ T cells in sepsis and other inflammatory diseases remains to be determined.

Sepsis is a life-threatening medical condition caused by various micro-organisms entering the bloodstream and triggering an uncontrolled inflammatory reaction. It is well known that excessive production of C5a contributes to this inflammatory response and is involved in the pathogenesis of sepsis.^{16,17} As excessive production of C5a is linked to worse survival in sepsis,¹⁸ the expression and regulation of the C5aR in sepsis would be expected to be important, but this has not been demonstrated.¹⁹

Our data show, for the first time, that $\gamma\delta$ T cells constitutively express the C5aR (CD88) and that C5a is directly involved in the regulation of $\gamma\delta$ T-cell activity in sepsis in a phosphoinositide 3-kinase (PI3K) -Akt signal pathway-dependent manner.

Materials and methods

Mice

Male C57BL/6 mice (7–8 weeks old) were obtained from Jackson Laboratories (Bar Harbor, ME) and bred in our facilities under specific pathogen-free conditions. All treatment of mice in this study was in strict compliance with the guidelines for the care and use of laboratory animals set out by the Institute of Basic Medical Sciences.

Reagents and antibodies

Recombinant human C5a was purchased from Sigma-Aldrich (St. Louis, MO). Wortmannin, a specific inhibitor of the PI3K-Akt signalling pathway, was purchased from Cell Signaling Technology, CST (Beverly, MA). Anti-mouse C5aR (CD88) monoclonal antibody was obtained from Cell Sciences (Clone 20/70, Canton, MA). Polyclonal rabbit anti-mouse C5a antibodies were produced as described previously.¹⁵ Briefly, the C-terminal of mouse C5a (amino acids 58–77) was chosen and the peptide was synthesized and coupled to keyhole limpet haemocyanin and used to immunize rabbits. The IgG was purified by protein A chromatography and its reactivity with recombinant mouse C5a (Sigma-Aldrich) was confirmed by ELISA. Rabbit IgG was used as the isotype control.

Induction of experimental sepsis by caecal ligation and puncture

Specific pathogen-free 7- to 8-week-old male C57BL/6 mice were used to establish the sepsis model. Briefly, the mice were anaesthetized by intraperitoneal injection of ketamine, then approximately two-thirds of the caecum

was ligated through a 2-cm abdominal midline incision and punctured extensively with a 21-gauge needle. After the bowel was repositioned, the abdomen was closed in layers, using a 4/0 surgical suture and metallic clips. Sham-operated mice were handled in the same manner, except that the caecum was not ligated and punctured. Immediately after caecal ligation and puncture (CLP), test animals were injected intravenously with 400 μ g anti-C5a antibodies in 400 μ l PBS, while the control animals received the same amount of normal rabbit IgG (Jing Mei Biotechnology, Beijing, China). This CLP model of sepsis is believed to closely simulate clinical sepsis in humans because of the polymicrobe-driven inflammatory process.

Isolation of $\gamma\delta$ T cells

Briefly, spleens were collected and spleen cells were enriched by Ficoll-Hypaque centrifugation (Pharmacia, Uppsala, Sweden) and washed twice with PBS containing 0.5% BSA and 2 mM EDTA. The cell pellet containing 1×10^7 cells was resuspended in 50 μ l PBS and 5 μ l phycoerythrin (PE) -conjugated anti- $\gamma\delta$ T-cell antibody (clone GL3; eBioscience Inc. (San Diego, CA)) was added and the mixture was incubated at 4° for 10 min. Non-bound antibodies were removed by two washes with PBS and the cells were re-suspended in 50 μ l PBS. Then, 10 μ l anti-PE-Microbeads (Miltenyi Biotec, Bergisch Gladbach, Germany) was added and the suspension was incubated at 4° for 15 min. After two washes, the cells were sorted using magnetic antibody cell sorting Separator columns (Miltenyi Biotec) into bound and non-bound fractions. The bound fraction was collected and its purity was measured by flow cytometry on a FACScaliber flow cytometer using CELL QUEST software (Becton Dickinson, San Jose, CA).

Flow cytometry analysis

To label C5aR (CD88) or C5L2 on $\gamma\delta$ T cells, spleen cells were stained for 30 min at 4° with FITC-conjugated anti- $\gamma\delta$ T-cell receptor antibody (eBioscience), allophycocyanin-conjugated anti-mouse CD3 antibody (eBioscience), and then for surface staining, PE-conjugated anti-mouse CD88 antibody (C5aR; Abcam, Cambridge, MA) were washed and used for flow cytometry analysis; for intracellular (C5L2) staining, the above cells were fixed overnight at 4° with 1 ml fixation buffer (Fix and Perm cell permeabilization kit; eBioscience). After washing, purified rat anti-mouse C5L2 (R&D, Minneapolis, MN) was added, followed by incubation with PE-conjugated anti-rat IgG (Santa Cruz Biotechnology, Santa Cruz, CA). After 30 min of incubation, the cells were washed and collected for flow cytometry analysis. For intracellular cytokine (IL-17) labelling, spleen cells were incubated for 4–6 hr at 4° with 1 μ g/ml brefeldin A, 50 ng/ml PMA and 1 μ g/ml ionomycin (all from Sigma-Aldrich), then they were stained for

30 min at 4° with FITC-conjugated anti- $\gamma\delta$ T-cell receptor antibody (eBioscience) and allophycocyanin-conjugated anti-mouse CD3 antibody and fixed overnight at 4° with 1 ml fixation buffer (Fix and Perm cell permeabilization kit; eBioscience). After washing, PE-conjugated anti-IL-17 antibody (eBioscience) was added, then, after 30 min of incubation at 4° and washed twice with 1 × permeabilization Buffer (eBioscience), the cells were collected for flow cytometry analysis on a FACSCalibur flow cytometer using CELL QUEST software.

Real-time-PCR

Quantitative reverse transcription PCR was performed with a multicolour real-time PCR detection system (iQ5, MJ; Bio-Rad, Hercules, CA), and DyNAmo HS SYBR Green qPCR kit (MJ BioWorks Inc., South San Francisco, CA).

The real-time PCR was performed following the manufacturer's protocol and carried out in triplicate. The PCR primer sequences were: mouse CD88: sense: 5'-atgcccaacacctaggccagcaag-3'; antisense: 5'-tgggcccacttcgaggtcccga-3'; mouse C5L2: sense: 5'-GCTCCTCCGCACTCCTT-3'; anti-sense: 5'-ACCGCACTTTCCTCATCC-3'; and mouse glyceraldehyde 3-phosphate dehydrogenase (GAPDH): sense 5'-tcttgggctacctgaggacc-3', antisense: 5'-cataccaggaatgagcttga-3'. The GAPDH was used as an endogenous control for any sample-to-sample variation in quantity and quality of RNA and difference in efficiency of reverse transcription and PCR.

Measurement of IL-17 by ELISA

The concentration of IL-17 and interferon- γ was measured using ELISA kits (eBioscience) and an automatic plate reader.

Measurement of intracellular cAMP

The purified $\gamma\delta$ T cells were incubated for 30 min with forskolin (10 nM; Sigma Co.). In some experiments, $\gamma\delta$ T cells were further incubated in the presence or absence of recombinant human C5a (20 nM, 100 nM) for 40 min at 37°. The treated cells were then washed twice with cold PBS and lysed in 250 μ l cell lysis buffer by three cycles of freeze-thaw. The cell lysates were used to measure cAMP using a Parameter cAMP assay kit according to the manufacturer's instructions (R&D Systems).

Western blot analysis

The $\gamma\delta$ T cells were stimulated with recombinant human C5a for the indicated time, then lysed. Equal amounts of protein were subjected to SDS-PAGE followed by transfer to a PVDF membrane and probing with antibodies against nuclear factor- κ B (NF- κ B) p65, κ B, Akt, or

phosphor-Akt, phosphor NF- κ B p65 (all from Cell Signaling Techniques). Bound antibody was visualized using an enhanced chemiluminescence kit (Amersham Biosciences, Piscataway, NJ).

Statistical analysis

Results were expressed as the mean \pm SEM and subjected to statistical analysis. Student's *t*-test or analysis of variance was used where appropriate to determine significant differences between samples. Values of *P* < 0.05 were considered significant.

Results

$\gamma\delta$ T cells constitutively express the C5a receptor, CD88

C5a is involved in the regulation of $\alpha\beta$ T-cell immunity through the C5aR expressed on CD4⁺ and CD8⁺ T cells.^{8–14} Our previous data¹⁵ also suggested an interaction between C5a and $\gamma\delta$ T cells in sepsis. However, whether C5a directly regulated the function of $\gamma\delta$ T cells, a T-cell subset with important roles in both innate and adaptive immunity, remained unclear. To test this, we first examined whether $\gamma\delta$ T cells express the C5a receptor, CD88. Spleen cells and mesenteric lymph node cells were prepared from naive C57BL/6 mice and C5aR expression on $\gamma\delta$ T cells was examined by flow cytometry. As shown in Fig. 1, C5aR (CD88) was constitutively expressed on the $\gamma\delta$ T-cell population. Besides CD88, C5L2 is another C5a receptor recently identified with biological functions.²⁰ Here we also examined the expression of C5L2 on $\gamma\delta$ T cells. There is no significant difference between the C5L2 staining group and the control group, indicating that C5L2 is not expressed on $\gamma\delta$ T cells (data not shown).

C5aR expression on $\gamma\delta$ T cells is up-regulated in mice following sepsis

Previous data showed that C5aR expression in different tissues is up-regulated following sepsis in mice.¹⁹ To test whether C5aR expression on $\gamma\delta$ T cells was also inducible, we used the CLP sepsis model described previously¹⁵ and measured C5aR expression on $\gamma\delta$ T cells by flow cytometry. As shown in Fig. 2(a,b), C5aR expression on $\gamma\delta$ T cells in a spleen cell suspension was significantly up-regulated in mice following sepsis. We also analysed the total numbers of CD88⁺ $\gamma\delta$ T cells in spleen. We found that after CLP, the total number of CD88⁺ $\gamma\delta$ T cells also increased (Fig. 2c), indicating that there was an enhanced CD88⁺ $\gamma\delta$ T-cell response in mice following sepsis. The $\gamma\delta$ T cells in mesenteric lymph node cells and peritoneal lavage fluid also showed increased C5aR expression in mice following sepsis (data not shown). We examined whether there was

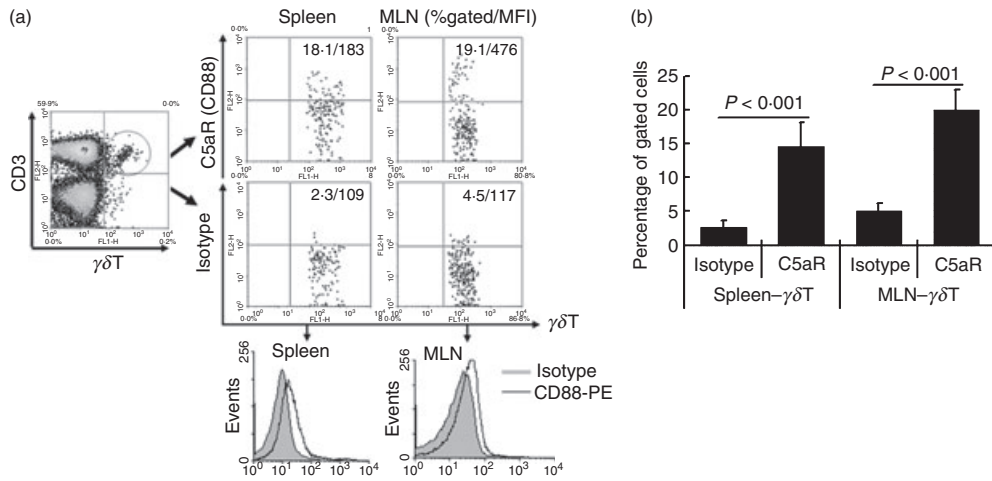


Figure 1. $\gamma\delta$ T cells constitutively express the C5a receptor, CD88. Spleen cells and mesenteric lymph node (MLN) cells were isolated from naive C57BL/6 mice and stained with antibodies against mouse CD3 (allophycocyanin-conjugated), the $\gamma\delta$ T T-cell receptor (FITC-conjugated), and the C5a receptor (CD88) (phycoerythrin-conjugated), then CD3⁺ $\gamma\delta$ T cells were gated to analyse CD88 expression. (a) Representative flow cytometry dot plot and histogram picture analysing the expression of CD88 on gated CD3⁺ $\gamma\delta$ T cells. (b) Statistical analysis of the expression of CD88 on $\gamma\delta$ T cells. The data shown are representative of the results from three independent experiments with at least four mice per group.

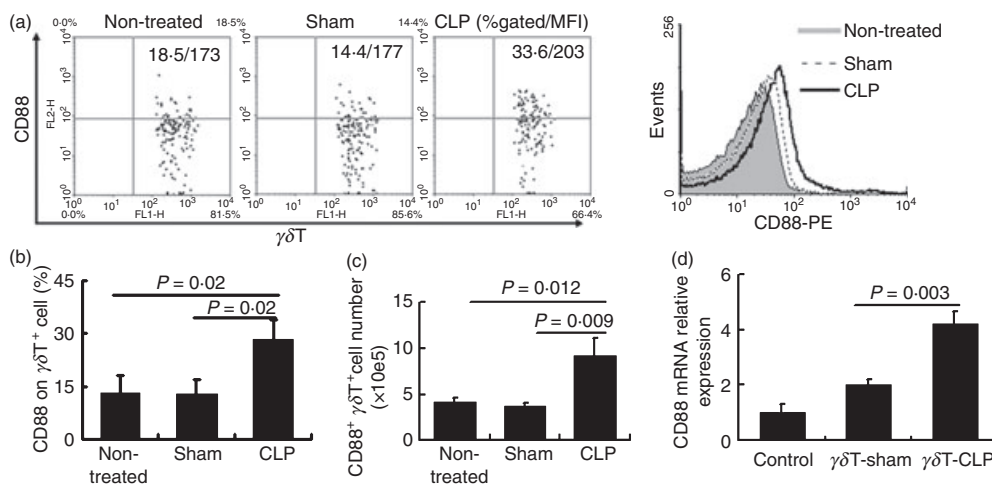


Figure 2. Increased expression of CD88 on $\gamma\delta$ T cells in mice following sepsis. Spleen cells were isolated from untreated, sham-operated, and septic (caecal ligation and puncture; CLP) mice and stained with antibodies against mouse CD3 (allophycocyanin-conjugated), the $\gamma\delta$ T T-cell receptor (FITC-conjugated), and the C5a receptor (CD88) (phycoerythrin-conjugated), then CD3⁺ $\gamma\delta$ T cells were gated to analyse CD88 expression. (a) Representative flow cytometry dot plot and histogram picture analysing the expression of CD88 on $\gamma\delta$ T cells. (b) Statistical analysis of CD88 frequency on $\gamma\delta$ T cells. The data shown are representative of the results for three independent experiments with at least four mice per group. (c) The total number of CD3⁺ CD88⁺ $\gamma\delta$ T in spleen cells from different group were analysed. (d) $\gamma\delta$ T cells were collected from sham control mice and septic mice, respectively. Then CD88 expression at mRNA level was examined by real-time PCR. Un-fractionated total lymphocytes were used as control.

enhanced CD88 expression in $\gamma\delta$ T cells collected from septic mice compared with those from sham controls by real-time-PCR. Our results showed that $\gamma\delta$ T cells from septic mice showed enhanced CD88 expression at mRNA levels compared with those from sham control mice (Fig. 2d). Here we also examined whether there was C5L2 mRNA expression in $\gamma\delta$ T cells from naive or septic mice. Consistent with the above data of flow cytometry, no C5L2 mRNA was detected in murine $\gamma\delta$ T cells (data not shown).

Neutralization of C5a *in vivo* down-regulates CD88 expression on $\gamma\delta$ T cells

We previously found that blocking C5a *in vivo* increases the survival rate and decreases $\gamma\delta$ T-cell activity in septic mice.¹⁵ To test whether excessive production of C5a directly affected the phenotype of the $\gamma\delta$ T cell, CD88 expression on $\gamma\delta$ T cells was examined in septic mice with or without intraperitoneal anti-C5a antibody administra-

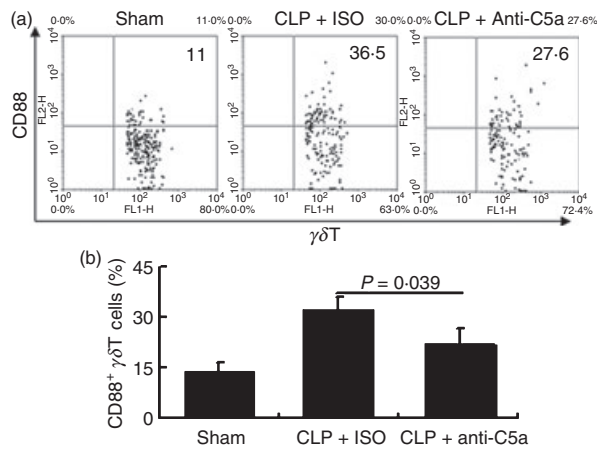


Figure 3. Neutralization of C5a partially prevents the up-regulation of CD88 expression on $\gamma\delta T$ cells in septic mice. Spleen cells from sham-operated mice and mice that have undergone caecal ligation and puncture (CLP) treated with anti-C5a antibody or control IgG were stained with antibodies against mouse CD3 (allophycocyanin-conjugated), the $\gamma\delta$ T-cell receptor (FITC-conjugated), and the C5a receptor (CD88) (phycoerythrin-conjugated), then CD3⁺ $\gamma\delta T$ cells were gated to analyse CD88 expression. (a) Representative flow cytometry dot plot. (b) Statistical analysis of the expression of CD88 on $\gamma\delta T$ cells. The data shown are representative of the results from three independent experiments with at least four mice per group.

tion. As shown in Fig. 3, C5a neutralization significantly down-regulated CD88 expression on $\gamma\delta T$ cells, indicating that excessive production of C5a contributed to the up-regulation of CD88 on $\gamma\delta T$ cells in sepsis. Considering our previous findings that neutralization of C5a *in vivo* significantly increases survival rate in sepsis and down-regulates IL-17 expression by $\gamma\delta T$ cells,¹⁵ these data further indicate that excessive production of C5a is directly involved in the regulation of $\gamma\delta T$ -cell activity in sepsis by up-regulating C5aR expression and enhancing cytokine production by $\gamma\delta T$ cells.

Recombinant C5a promotes up-regulation of CD88 expression by $\gamma\delta T$ cells at both protein and mRNA level

To directly investigate the effects of C5a on CD88 expression by $\gamma\delta T$ cells *in vitro*, we incubated mouse spleen cells with recombinant human C5a for 12 hr, and then measured CD88 expression by $\gamma\delta T$ cells. As shown in Fig. 4(a,b), stimulation with C5a significantly up-regulated CD88 expression by $\gamma\delta T$ cells.

Given that CD69, CD44 and CD62 ligand (CD62L) have been implicated as activation markers of $\gamma\delta T$ cells,^{21,22} we also investigated whether ligation of the

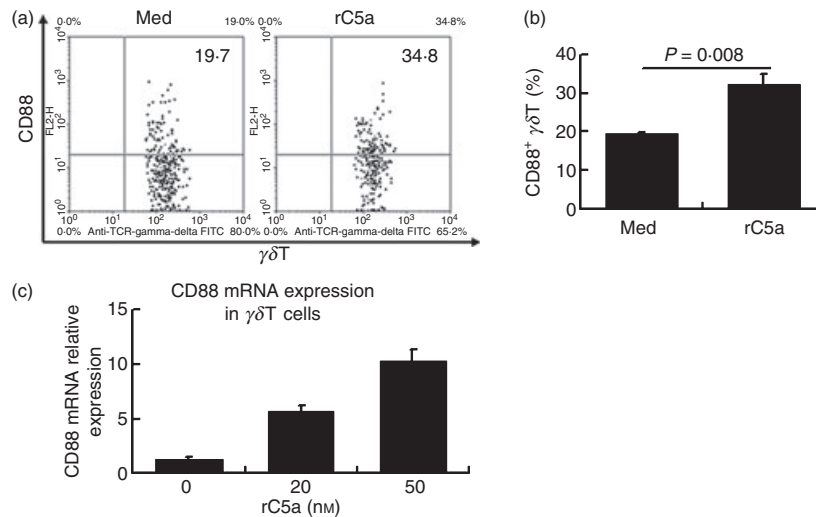


Figure 4. Recombinant C5a promotes up-regulation of CD88 expression by $\gamma\delta T$ cells at both protein and mRNA level. Spleen cells from untreated C57BL/6 mice were incubated with recombinant human C5a (rC5a) (20 nM) or medium (Med) for 12 hr, then were stained with antibodies against mouse CD3 (allophycocyanin-conjugated), the $\gamma\delta T$ T-cell receptor (FITC-conjugated), and the C5a receptor (CD88) (phycoerythrin-conjugated) then CD3⁺ $\gamma\delta T$ cells were gated to analyse CD88 expression. (a, b) Representative flow cytometry dot plots analysing expression of CD88 (a) by $\gamma\delta T$ cells. (b) Statistical analysis of the expression of CD88 by $\gamma\delta T$ cells. (c) $\gamma\delta T$ cells were purified from septic mice as described in the Materials and Methods, and incubated with recombinant human C5a at the indicated dosage. Twelve hours later, cells were collected and used for real-time PCR analysis. The results shown are representative of those from four independent experiments. We also examined the effects of C5a receptor ligation on the activation of $\gamma\delta T$ cells using different markers. Although CD69, CD44 and CD62 ligand expression on $\gamma\delta T$ cells was significantly up-regulated in mice following sepsis, incubation of spleen cells with C5a did not affect the expression of these molecules on $\gamma\delta T$ cells (data not shown).

C5aR also affected their expression. Although CD69, CD44 and CD62L expression on $\gamma\delta$ T cells was significantly up-regulated in mice following sepsis, incubation of spleen cells with C5a did not affect the expression of these molecules on $\gamma\delta$ T cells (data not shown).

To test whether C5a induced CD88 expression because of increased gene transcription, we purified $\gamma\delta$ T cells and incubated them with recombinant complement C5a for 12 hr. Then cells were collected and analysed for CD88 expression. As shown in Fig. 4(c), C5a incubation dose-dependently increased CD88 expression at the mRNA level.

C5a receptor mediates increased IL-17 expression from $\gamma\delta$ T cells

To directly examine the effects of C5aR ligation on the activity of $\gamma\delta$ T cells, we incubated mouse spleen cells or purified $\gamma\delta$ T cells with recombinant human C5a for 12 hr, then measured IL-17 expression by $\gamma\delta$ T cells by flow cytometry. To further confirm that CD88, but not C5L2, was involved in C5a-enhanced activity of $\gamma\delta$ T cells, neutralizing antibodies active against mouse CD88 were added in combination with recombinant human C5a, and then IL-17 expression by $\gamma\delta$ T cells was measured by flow cytometry. As shown in Fig. 5(a,b), incubation of mouse spleen cells with recombinant human C5a significantly increased IL-17 expression within $\gamma\delta$ T cells. Neutralization of CD88 blocked the activity of C5a-enhanced IL-17 expression, again indicating that CD88 but not C5L2 is involved in the activation of $\gamma\delta$ T cells after the addition of C5a.

Subsequently, we purified $\gamma\delta$ T cells from septic mice or sham controls (Fig. 5c) and then co-cultured them with recombinant human C5a with or without recombinant mouse IL-23 for 24 hr, then measured IL-17 levels in the culture supernatants. The ELISA data showed that recombinant C5a alone did not induce IL-17 secretion, whereas IL-23 caused some release and the combination was more effective (Fig. 5c). We previously reported that $\gamma\delta$ T cells from septic mice showed enhanced CD88 expression. As expected, compared with $\gamma\delta$ T cells from sham control mice, $\gamma\delta$ T cells from septic mice showed enhanced IL-17 secretion after C5a stimulation (Fig. 5d). These data directly show that a C5a–C5aR– $\gamma\delta$ T–IL-17 axis exists in the inflammatory response in diseases such as sepsis.

Ligation of C5a activates the PI3K/Akt signalling pathway in $\gamma\delta$ T cells

C5aR signalling, in which the C5aR is coupled to G protein α and $\beta\gamma$ subunits and activates several downstream signalling pathways (e.g. PI3K, NF- κ B, cAMP), has been studied in dendritic cells, macrophages, neutrophils and several other cell types.^{23,24} However, the mechanism of C5aR signalling in $\gamma\delta$ T cells is not known. It has been shown that C5a stimulation activates the Akt pathway in dendritic cells,⁷ raising the possibility that it could have a similar effect on this pathway in $\gamma\delta$ T cells. To identify the signalling pathway(s) responsible for the C5a-mediated activation of $\gamma\delta$ T cells, we first examined whether C5a could affect the activation of the PI3K/Akt and NF- κ B pathways. As shown in Fig. 6(a–c), C5a stimulation enhanced Akt but inhibited NF- κ B-p65 phosphorylation.

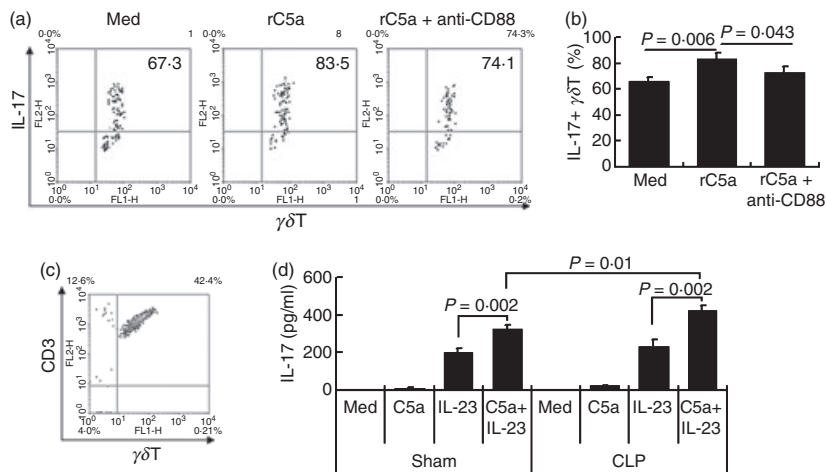


Figure 5. C5a receptor CD88 mediates increased interleukin-17 (IL-17) expression by $\gamma\delta$ T cells. Spleen cells from untreated C57BL/6 mice were incubated with recombinant human C5a (rC5a; 20 nM) or medium (Med) or rC5a plus anti-mouse CD88 monoclonal antibody (2 μ g/ml) for 12 hr, then were stained with antibodies against mouse CD3 (allophycocyanin-conjugated), the $\gamma\delta$ T-cell receptor (FITC-conjugated), and IL-17 (intracellular staining; phycoerythrin-conjugated), then CD3⁺ $\gamma\delta$ T cells were gated to analyse IL-17 expression (a, b). (c) $\gamma\delta$ T cells were purified from septic or sham control mice and (d) were incubated with recombinant human C5a (20 nM) and/or recombinant mouse IL-23 (10 ng/ml) for 24 hr, then the culture supernatants were assayed for IL-17 by ELISA. The results shown are representative of four independent experiments.

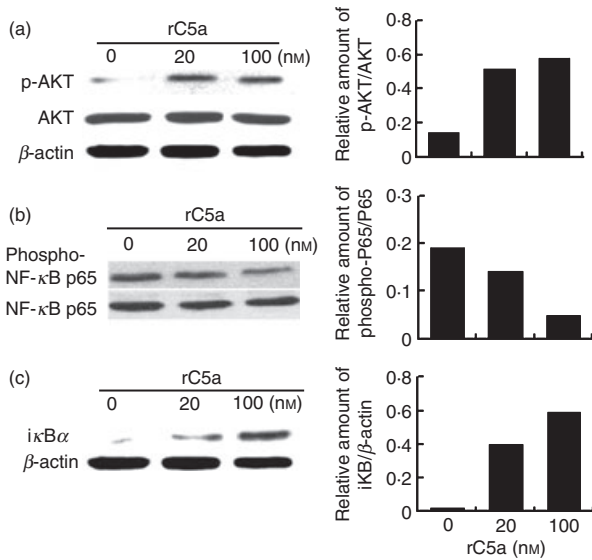


Figure 6. C5a stimulation activates the phosphoinositide 3-kinase (PI3K)/Akt signalling pathway while inhibit nuclear factor-κB (NF-κB) signalling in $\gamma\delta$ T cells. Purified $\gamma\delta$ T cells were stimulated with recombinant human C5a at the indicated dose for 15 min, then cell lysates were tested for phosphorylated Akt (a) and NF-κB p65 (b), as well as $\text{i}\kappa\text{B}$ (c) by Western blotting using antibodies against phospho-Akt, phospho- NF-κB p65, Akt, NF-κB p65, $\text{i}\kappa\text{B}$ or β -actin respectively and the bands were analysed by densitometry.

To test whether and how C5a stimulation affected other signalling pathways in $\gamma\delta$ T cells, we measured cAMP levels in $\gamma\delta$ T cells after C5a stimulation, as the cAMP pathway is a key signalling pathway for G protein-coupled receptors²⁵ and cAMP functions as an intracellular second messenger,²⁶ which regulates a wide range of important cellular processes. In contrast to those found in dendritic cells, in which C5aR, a G protein-coupled receptor, decreases cAMP production, we found that ligation of C5aR increased cAMP levels in $\gamma\delta$ T cells (data not shown). The underlying mechanisms remain to be determined.

Finally, to test the role of the PI3K-Akt pathway in C5a/C5aR-mediated $\gamma\delta$ T-cell activation, we pre-treated spleen cells for 1 hr with the PI3K/Akt inhibitor wortmannin, then incubated them with recombinant human C5a in the continued presence of wortmannin for 12 hr. As shown in Fig. 7(a,b), wortmannin prevented the C5a-induced up-regulation of CD88 on $\gamma\delta$ T cells. We also pre-treated $\gamma\delta$ T cells with wortmannin before stimulating them with C5a and IL-23 and measuring IL-17 secretion and found that wortmannin also inhibited C5a/IL-23-induced IL-17 secretion by $\gamma\delta$ T cells (Fig. 7c). As we use a combination of IL-23 and C5a here, to test whether wortmannin affects pathways downstream of the IL-23R or the C5aR, we examined whether wortmannin affects the signalling of IL-23 in $\gamma\delta$ T cells. Our data showed that

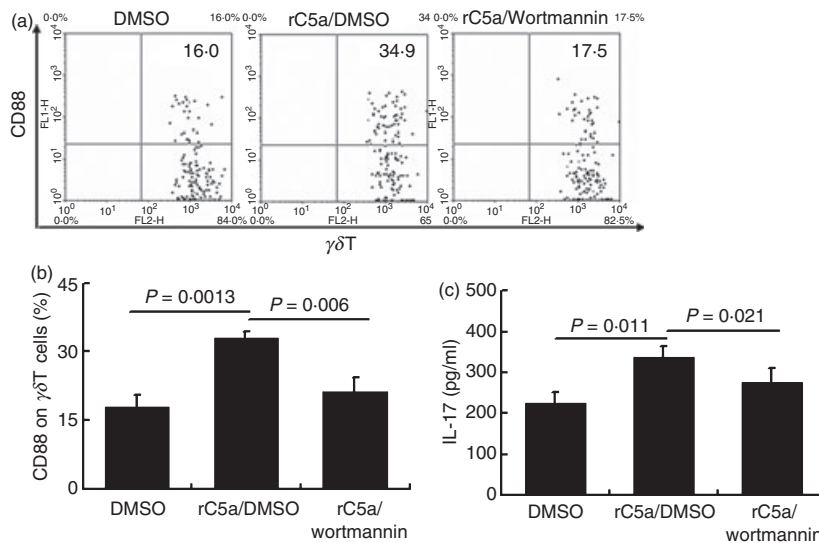


Figure 7. Inhibition of the phosphoinositide 3-kinase (PI3K)/Akt signalling pathway by wortmannin prevents the up-regulation of CD88 and interleukin-17 (IL-17) expression by C5a. (a) Spleen cells were pre-treated with vehicle (DMSO) or the PI3K/Akt inhibitor wortmannin (1 μM) for 1 hr and incubated for 12 hr with recombinant C5a (20 nM), then were analysed by flow cytometry for expression of CD88 on $\text{CD}3^+ \gamma\delta\text{T}^+$ cells. (b) Statistical analysis of CD88 and IL-17 expression by the above $\gamma\delta$ T cells. The data are representative of those for three independent experiments with at least four mice per group. (c) Purified $\gamma\delta$ T cells were pre-treated with wortmannin (1 μM) for 1 hr, then were stimulated with C5a and IL-23 (10 ng/ml) for 24 hr and the culture supernatant was collected and analysed for IL-17 by ELISA. The results shown are representative of those from three independent experiments.

wortmannin did not significantly affect IL-23-induced IL-17 production by $\gamma\delta$ T cells (data not shown), indicating that in the above combination, wortmannin mainly affects the signalling of C5aR.

Discussion

$\gamma\delta$ T cells are actively involved in both innate and adaptive immunity. However, the underlying mechanisms remain to be determined. Here, we showed, for the first time, that $\gamma\delta$ T cells constitutively expressed the C5aR and that its expression was up-regulated in mice following sepsis. Ligation of the C5aR by C5a directly promoted the activation of $\gamma\delta$ T cells in a PI3K-Akt signalling pathway-dependent manner. Our data suggest an important C5a- $\gamma\delta$ T-cytokine (IL-17) axis, which partially explains the pathogenic roles of C5a in inflammatory diseases. The observation that IL-17 secretion by $\gamma\delta$ T cells could also be modified by C5a provides further evidence supporting the existence of a cross-link between the complement system and T-cell immunity.

Excessive production of C5a and up-regulation of the C5aR has been found in different tissues in mice following injury or sepsis.¹⁹ These data suggest that excessive production of C5a might cause its pathogenic effects by binding to the over-expressed C5aR on different cells. Although recent studies have shown regulation of T-cell immunity by the C5a-C5aR axis,⁸⁻¹⁴ whether T cells express the C5aR and its implications in sepsis remained to be determined. In this study, we evaluated, for the first time, the cellular expression and kinetics of the C5aR on $\gamma\delta$ T cells. The observations that CD88 expression on $\gamma\delta$ T cells was up-regulated in mice following sepsis and that C5a neutralization *in vivo* down-regulated C5aR expression (Fig. 3) suggest that C5a is involved in the regulation of $\gamma\delta$ T cells in sepsis. $\gamma\delta$ T cells show increased expression of CD62L, CD44 and CD69 after activation.^{21,22} However, although CD62L, CD44 and CD69 expression on $\gamma\delta$ T cells was significantly up-regulated in mice following sepsis, C5a treatment alone did not affect the expression of these markers on $\gamma\delta$ T cells (data not shown). Here we also examined the absolute number of CD88⁺ $\gamma\delta$ T cells in the spleen after anti-C5a administration. Consistent with our previous findings,¹⁵ anti-C5a antibody did not significantly down-regulate the absolute number of CD88⁺ $\gamma\delta$ T cells *in vivo*, indicating that some other factors are involved in enhancing the activity of $\gamma\delta$ T cells in sepsis (data not shown).

Although $\gamma\delta$ T cells constitute only a small proportion (1-5%) of the lymphocytes in the blood and peripheral organs, they play important roles in both innate and adaptive immunity.¹ Recently, several groups, including our own, suggested that $\gamma\delta$ T cells, which are a major source of IL-17, are involved in the pathogenesis of sepsis.^{15,27,28} In addition, it is clear that, in septic mice,

there is excessive production of C5a and IL-17 and that blockade of either is beneficial for the outcome in sepsis.^{17,29} However, whether there are interactions between these two important pro-inflammatory cytokines remained unclear. Previously, we found that C5a is involved in the regulation of IL-17 production in septic mice and that C5a neutralization *in vivo* increases the survival rate and decreases IL-17 production by $\gamma\delta$ T cells.¹⁵ These data indicate a link between IL-17-producing $\gamma\delta$ T cells and C5a. However, whether C5a could directly regulate the activity of IL-17-producing $\gamma\delta$ T cells was unclear. In this study, we provided evidence that, by binding to C5aRs expressed on $\gamma\delta$ T cells; C5a directly takes part in the regulation of $\gamma\delta$ T-cell immune responses. Hence, we have identified a new axis, in which C5a acts directly on $\gamma\delta$ T cells and increases C5aR expression on $\gamma\delta$ T cells, enhances $\gamma\delta$ T-cell activity (such as promoting IL-17 production), and promotes the inflammatory response in sepsis.

After demonstrating expression of C5aR on $\gamma\delta$ T cells *in vivo*, we investigated the effects of C5aR ligation on the surface markers and function of $\gamma\delta$ T cells. As shown in Figs 4 and 5, incubation of recombinant complement C5a increased CD88 and IL-17 expression by $\gamma\delta$ T cells. Blockade of CD88 with antibody abolished the effects of recombinant C5a-induced CD88 and IL-17 up-regulation, indicating that the effects of C5a on $\gamma\delta$ T cells were mediated by CD88 but not another receptor of C5a-C5L2. $\gamma\delta$ T cells from septic mice show up-regulated CD88 expression, which raised the possibility that these $\gamma\delta$ T cells might show an enhanced response to C5a stimulation compared with cells from control mice. As expected, C5a stimulation did induce a significant increase in IL-17 production by $\gamma\delta$ T cells from septic mice compared with those from sham control mice (Fig. 5d). Although recombinant C5a alone did not induce significant IL-17 secretion, it facilitated the activity of IL-23. We also examined the effects of C5a ligation on interferon- γ production by $\gamma\delta$ T cells. We found that C5a alone could not induce the production of interferon- γ . However, C5a enhanced IL-23-mediated interferon- γ production by $\gamma\delta$ T cells (data not shown). These data again indicate that complement components are involved in the regulation of $\gamma\delta$ T cells through the up-regulation of CD88 expression on $\gamma\delta$ T cells. In this scenario, excessive production of C5a itself is directly or indirectly involved in the up-regulation of C5aR expression, then binds to the up-regulated C5aR, and promotes the pathogenic roles of $\gamma\delta$ T cells in septic mice.

Another important finding in this study was the identification of the intracellular signalling pathways by which C5a modulates $\gamma\delta$ T-cell function. The PI3Ks are a conserved family of signal transduction enzymes that are involved in regulating cellular proliferation and survival.³⁰ Akt, the downstream serine/threonine kinase of PI3Ks

(also known as protein kinase B), which was first described as an oncogene, regulates cellular activation, inflammatory responses, chemotaxis and apoptosis.^{31,32} Our data clearly demonstrated that the PI3K-Akt signalling pathway was involved in the activation of $\gamma\delta$ T cells by C5a, as (i) C5a ligation increased Akt phosphorylation and (ii) a PI3K inhibitor, which inhibits the PI3K/Akt pathway, prevented the C5a-induced increase in CD88/IL-17 expression by $\gamma\delta$ T cells. Bommhardt *et al.*³³ reported that over-expression of Akt by lymphocytes results in decreased lymphocyte apoptosis and a better survival in mice in response to CLP-induced sepsis. These data indicate that the PI3K-Akt pathway is involved in maintaining the homeostasis of T cells. Our data showing that ligation of C5aR activates the Akt pathway in $\gamma\delta$ T cells suggest that C5a might also be involved in maintaining the homeostasis of $\gamma\delta$ T cells in inflammatory diseases, such as sepsis.

Nuclear factor- κ B is a ubiquitous inducible transcription factor that stimulates gene expression, in particular of those genes that promote immune and inflammatory responses.³⁴ A variety of extracellular signals can activate NF- κ B.³⁵ To examine the effect of C5a on NF- κ B activation in $\gamma\delta$ T cells, we measured levels of phosphor-NF- κ B-p65 in response to C5a stimulation. Unexpectedly, we found that C5a engagement resulted in NF- κ B suppression (Fig. 6b,c). Although there are data indicating that the PI3K-Akt signalling pathway is involved in cell survival by inhibiting the activation of pro-apoptotic proteins and transcription factors,^{36–38} the underlying mechanisms by which C5a signalling inhibits NF- κ B activation remain to be determined.

Cyclic AMP is a cyclic nucleotide that functions as an intracellular second messenger²⁶ and regulates a wide range of important cellular processes. As the cAMP pathway is a key signalling pathway for G protein-coupled receptors, such as the C5aR, we examined whether the C5aR/cAMP signalling pathway was also involved in the regulation of $\gamma\delta$ T cells. Different from those found in dendritic cells, in which C5aR, a G protein-coupled receptor, decreases cAMP production,⁷ we found that ligation of C5aR increased cAMP levels in $\gamma\delta$ T cells (data not shown). As increased cAMP levels may contribute to the inhibition of NF- κ B,³⁵ our data suggest a dual effect of C5a signalling on the activity of $\gamma\delta$ T cells, on the one hand activating the PI3K signalling pathway and enhancing the activity of $\gamma\delta$ T cells and, on the other, being involved in maintaining $\gamma\delta$ T-cell homeostasis by inhibiting the activation of NF- κ B (Fig. 7). The underlying mechanisms require further investigation.

In summary, our data demonstrate, for the first time, that $\gamma\delta$ T cells constitutively express the C5aR and that, following sepsis, C5aR expression is significantly up-regulated. Excessively produced C5a is involved in the regulation of CD88 expression on $\gamma\delta$ T cells. Ligation of the C5aR by C5a

enhances CD88 and IL-17 expression by $\gamma\delta$ T cells in a PI3-K/Akt signalling pathway-dependent manner. C5a engagement increases cAMP levels and inhibits NF- κ B activation in $\gamma\delta$ T cells. The C5a/C5aR- $\gamma\delta$ T-IL-17 axis provides further evidence for the involvement of C5a in the regulation of T cells and a new mechanism by which C5a influences inflammatory responses. Further studies on cross-talk between the complement system and $\gamma\delta$ T cells and its biochemical basis will provide insight into the pathogenesis of, and therapeutic approaches to, inflammatory diseases.

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Disclosures

The authors have no financial conflict of interest.

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