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Characterisation of eppin function: expression and activity in the lung

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Take Home Message – Eppin is a low molecular weight protein which is expressed in the human lung during inflammation.
Abstract

Eppin is a serine protease inhibitor expressed in male reproductive tissues. In this study we have demonstrated novel sites of eppin expression in myeloid and epithelial cell lines with further confirmation in primary myeloid cell types. Using immunohistochemistry and Western blotting, eppin was detected in the lungs of patients with Acute Respiratory Distress Syndrome and Cystic Fibrosis lung disease. Expression of eppin in monocytic cells was unaffected by stimulation with TLR agonists, cytokine stimulation and hormone receptor agonist stimulation. However, upregulated expression and secretion of eppin was observed following treatment of monocytes with epidermal growth factor (EGF). Incubation of recombinant eppin with monocytic cells resulted in significant inhibition of lipopolysaccharide (LPS)-induced chemokine production. Furthermore, eppin inhibited LPS-induced NF-κB activation by a mechanism which involved accumulation of phosphorylated IkBα. In an in vivo model of lung inflammation induced by LPS, eppin administration resulted in decreased recruitment of neutrophils to the lung with a concomitant reduction in the levels of the neutrophil chemokine MIP-2. Overall, these results suggest a role for eppin outside of the reproductive tract and that eppin may have a role in the innate immune response in the lung.
**Introduction**

Human epididymal protease inhibitor (eppin, SPINLW1, WFDC7) is a cysteine-rich protein comprised of both a Kunitz-type and a WAP (whey acidic protein) four-disulfide core (WFDC) protease inhibitor consensus motif [1, 2]. In the epididymis, eppin is secreted by epithelial cells and is a key component in the eppin protein complex which mediates the capacitation of spermatozoa [3]. In this role, eppin acts both as a serine protease inhibitor, inhibiting the action of prostate specific antigen [4], while protecting the spermatozoa from potential damage by bacteria [3, 5]. The majority of work on eppin to date has therefore focused on the development of eppin as target for a male contraceptive based on inhibition of the eppin protein complex via specific antibodies [6].

Although eppin is a serine protease inhibitor, the arrangement of cysteine residues differs from that of SLPI and elafin, the two most prominent members of the WFDC family [7]. Similar to SLPI and elafin, recombinant eppin inhibited elastase activity, but unlike SLPI, eppin had no effect on trypsin or chymotrypsin activities [8–11]. Split domain studies demonstrated that the serine antiprotease activity resides solely in the Kunitz domain of eppin, while the WFDC domain retains no antiprotease activity [9]. Like SLPI [12], findings suggest that eppin also possesses antibacterial activity against *Escherichia coli* and that this effect is mediated via membrane permeabilisation [5]. In contrast to its antiprotease activity, the antibacterial activity of eppin against *E. coli* appears to be mediated by both the WFDC and Kunitz domains, with maximal killing achieved by the full-length protein [9].
On the basis of their multiple functions, WFDC proteins are considered to play key roles in innate immunity at mucosal surfaces. In contrast to eppin, elafin and SLPI are constitutively expressed by a variety of cell types, including respiratory and intestinal epithelia as well as key cell types involved in the innate immune response such as mast cells, neutrophils, and macrophages [13–19]. Furthermore, SLPI and elafin expression is altered in response to a number of factors including pro-inflammatory cytokines, bacterial products, proteases and androgens [16, 20–22]. The anti-inflammatory activity of SLPI and elafin in vitro and in vivo is well documented and reviewed elsewhere [7, 23, 24]. Previous work has shown that SLPI is capable of inhibiting lipopolysaccharide (LPS)-induced responses [20, 25–27]. In addition, elafin inhibited LPS activation of monocytic cells by inhibiting NF-κB activation via a process partly involving accumulation of polyubiquitinated IκBα [28]. Little is known about the function of eppin outside of the epididymis. In this study, we expand current knowledge on the localisation and regulation of eppin expression, and explore its potential role as a multifunctional host defence protein.
Results

*Eppin is expressed in myeloid and airway epithelial cells*

Given that SLPI and elafin transcripts are found in a variety of cell types, we first investigated whether eppin had a similar pattern of expression. As expected, eppin expression was detected in testes and lung tissue, albeit to a lesser extent in the lung tissue (Figure 1A). Eppin expression was also detected in myeloid cell lines, such as THP-1 and U937 monocytic cells, monocyte-derived macrophages, and neutrophils differentiated from PLB-985 cells (Figure 1A). Furthermore, eppin was expressed in both human peripheral blood mononuclear cells (PBMCs) (Figure 1B) and polymorphonuclear leucocytes (PMNs) (Figure 1A). Expression of eppin was also detected in epithelial cell lines of the respiratory tract including human type-II alveolar cells (A549), as well as tracheal and bronchial epithelial cell lines (Figure 1B). Eppin expression was highest in testes expression and, expression of eppin in the other tissues/cells (relative to eppin at 100%) was: THP-1 monocyte (80%), THP-1 macrophage (75%), U937 monocyte (65%), U937 macrophage (55%), PLB-985 cells (57%), PMNs (80%), A549s (39%), lung (47%), PBMCs (24%), HTEs (17%), CFTEs (17%), HBEs (24%) and CFBEs (22%) as determined by densitometry. Sequencing of PCR products confirmed the identity of human eppin.

**Detection of eppin in human lung tissue, bronchoalveolar lavage fluid (BALF) and sputum**

Human lungs were perfused with LPS as outlined in the Methods section. Tissue samples from non-LPS perfused lobes, LPS-perfused lobes and from cystic fibrosis (CF) patients were used for immunohistochemistry. Some eppin staining was present in non-LPS perfused lung tissue (Figure 2A) and was particularly evident in the LPS-perfused (Figure 2B) and CF
patient (Figure 2C) lung tissues. In agreement with the eppin expression data outlined in Figure 1, positive staining was prominent in epithelial and immune cells using two different eppin antibodies, confirming the presence of eppin in the in vivo lung. In addition, eppin was shown to be present in BALF from patients with Acute Respiratory Distress Syndrome (ARDS) and in sputum from patients with CF but was not present in BALF from healthy volunteers (Figure 3). Eppin was detected as the same high molecular weight complex in ARDS BALF and CF sputum which has been shown previously in semen where eppin is complexed to other proteins including clusterin and lactoferrin. There appears to be low microgram amounts of eppin in CF sputum and ARDS BALF as determined by densitometry of the CF sputum and ARDS BALF blots. Other lower molecular bands were detectable in CF sputum which may represent proteolytic breakdown products of the eppin complex.

**Eppin expression and secretion are upregulated by EGF**

In order to investigate factors that may affect eppin expression, THP-1 cells were stimulated with a range of inflammatory agonists and effects on eppin expression were assessed by RT-PCR. A number of TLR agonists such as LPS (TLR4), Pam3CysSK4 (TLR2) and polyinosinic-polycytidylic acid (TLR3) had no effect on eppin expression (data not shown). Hormone receptor agonists such as estradiol and progesterone are known inducers of SLPI expression [21, 22]. However, estradiol or progesterone did not modulate eppin expression (data not shown). Growth factors such as epidermal growth factor (EGF) and transforming growth factor-β (TGF-β) have also been shown to regulate SLPI expression [29, 30]. To investigate a possible role for growth factors in the modulation of eppin expression, THP-1 monocytic cells were stimulated with EGF, TGF-β and fibroblast growth factor (FGF) at a range of
concentrations. TGF-β and FGF treatment of monocytes had no effect on eppin expression (data not shown). However, eppin expression was increased after treatment with EGF (Figure 4 A and B). In agreement with this result, analysis of cell supernatants by Western blotting revealed the presence of increased levels of eppin in EGF-treated cell supernatants (Figure 4C). The identity of eppin was confirmed by following digestion of protein in SDS-PAG pieces from EGF-treated cell supernatant followed by 1D nLC-MS-MS (reverse phase chromatography) to identify the protein present in the SDS-PAG samples. The two main peptides obtained from this analysis were ANCLNTCK and DRQCQDNKK both of which demonstrate 100% overlap with human eppin.

**Effect of recombinant eppin on bacterial growth**

To investigate a potential role for eppin as a host defence protein in the lung, we expressed and purified recombinant human eppin. The ability of eppin to permeabilise clinically relevant Gram negative and Gram positive bacteria as a method of bacterial killing was compared to LL-37, a well-known antimicrobial peptide. Eppin permeabilised *P. aeruginosa* (Figure 5A) and *Staphylococcus aureus* (Figure 5B) bacterial strains and was comparable, if not better than, the permeabilisation observed with LL-37 particularly in the case of *P. aeruginosa*. Eppin also inhibited the growth of *P. aeruginosa* and *S. aureus* when assessed by radial diffusion assay, once again, with significant effect on *P. aeruginosa* growth (Figure 5C).
Effect of recombinant eppin on LPS-induced cytokine production

Previous work has shown that the related WFDC proteins, SLPI and elafin, can modulate LPS signalling in monocyctic cells [26–28]. Pre-incubation of THP-1 monocyctic cells with eppin for 1 h prior to LPS stimulation significantly inhibited both the expression (Figure 6A) and production of IL-8 (Figure 6B). In addition, eppin pre-incubation inhibited LPS-induced production of IL-8 from THP-1 monocyte-derived macrophages (Figure 6C) and MIP-2 from murine bone marrow-derived macrophages (BMDMs) (Figure 6D). We also observed no significant effect of eppin on THP-1 cell viability (Figure 6E).

Effects of eppin on NF-κB activation

To investigate further the effect of eppin on the LPS signalling pathway, we evaluated the impact of eppin on LPS-induced activation of NF-κB. Eppin significantly inhibited LPS-induced p65:DNA binding activity in THP-1 nuclear extracts (Figure 7A), thereby suggesting that the ability of eppin to inhibit LPS-induced cytokine expression may be mediated via its effects on the NF-κB signalling pathway. SLPI has previously been shown to inhibit LPS-induced NF-κB activity by competing with p65 for binding to consensus sites in the promoter regions of target genes such as IL-8 [27]. However, in contrast to SLPI, eppin was unable to bind NF-κB consensus oligonucleotides (Figure 7B). Eppin’s effect on LPS signalling upstream of NF-κB was evaluated by Western blotting. LPS-induced degradation of IκBα was observed in samples over time regardless of whether eppin was present or not (Figure 7C). However, increased phosphorylation of IκBα was observed in the eppin-treated cells compared to cells treated with LPS alone (Figure 7C).
Eppin inhibits LPS-induced lung neutrophil recruitment in vivo

To determine if eppin might alter LPS effects in vivo, the effects of eppin in an in vivo model of LPS-induced acute lung inflammation were investigated. Mice receiving eppin exhibited a significant reduction in pulmonary total cell counts following intratracheal LPS administration (Figure 8A) with a significant decrease in neutrophil counts (Figure 8B). This decrease in neutrophil infiltration was associated with a reduction in levels of the neutrophil chemoattractant MIP-2 in BAL fluid from mice that received eppin and LPS compared to LPS alone (Figure 8C). These data confirm the in vitro findings for MIP-2 shown in Figure 5D and suggest that eppin may play an immunomodulatory role in the airways by reducing LPS-induced MIP-2 production from host cells with a subsequent reduction in neutrophil recruitment.
Discussion

The majority of work on eppin to date has focused on its role within the male reproductive tract. In this study, we have identified novel sites of eppin expression in both myeloid and epithelial cell types. Furthermore, eppin expression was confirmed in primary mononuclear cells and neutrophils. We have also demonstrated the presence of eppin in tissue and airways (BALF and sputum) from patients with ARDS and CF, as well as from human ex vivo lung tissue perfused with LPS. Given the similar sites of expression we have demonstrated between eppin and other WFDC genes, we examined whether the expression of eppin in monocytic cells was modulated by factors known to regulate the expression of SLPI and/or elafin. However, we found that eppin expression was not induced by various TLR agonists, hormone receptor agonists and cytokine stimuli used in this study.

These findings are similar to those reported for HE4 (WFDC2), another WFDC family member. HE4 was originally thought to be limited to the reproductive tract however, the expression profile of HE4 has since expanded to include the trachea, lung and nasal epithelium as well as major and minor salivary glands [13, 31]. HE4 expression was not affected by pro-inflammatory agonists such as IL-1β and TNF-α [14]. Although SLPI expression is induced by hormone receptor agonists such as estradiol and progesterone, elafin expression is unaffected [21, 22]. In addition, growth factors have been shown to regulate SLPI expression in vitro and in vivo [29, 30, 32]. In this study, eppin expression in monocytic cells was unaffected by estradiol and progesterone treatment (data not shown), however, both eppin expression and secretion were up-regulated by EGF treatment. We have also provided evidence of eppin protein in the lungs of patients with Cystic Fibrosis and
ARDS by immunohistochemistry and Western blot indicating that eppin is increased during inflammation with little eppin present in healthy lung. It should be highlighted that further confirmation of eppin presence in diseased lung could be provided by the availability of more specific monoclonal antibodies to eppin as there is a paucity of eppin antibodies available to carry out definitive in vivo studies. However, our immunohistochemistry and Western blot data does correlate to the eppin expression detected in lung epithelial and inflammatory cells as shown in Figure 1. Eppin was not detected as a monomeric species in CF and ARDS airway secretions but was present as a high molecular weight complex or oligomeric species in these biological fluids which has been shown previously in semen where eppin is complexed to other proteins including clusterin and lactoferrin^3.

For the purposes of investigating a potential host defence role for eppin, we expressed and purified recombinant eppin [9]. Our subsequent analyses focused on exploring the antibacterial and immunomodulatory potential of eppin. Recombinant eppin has previously been shown to inhibit growth of E. coli [5, 9]. In this study, we expand these observations by demonstrating an antibacterial effect of eppin against P. aeruginosa and S.aureus, two clinically relevant pulmonary pathogens. Although there are limitations associated with the radial diffusion assay, the bacterial membrane permeabilisation assay provides more definitive anti-bacterial activity data and the combined results of both assays serve to confirm that eppin has a greater effect on P. aeruginosa than S.aureus. In order to investigate the immunomodulatory effects of eppin, our studies focussed on the effects of eppin on LPS signalling and downstream responses in human monocytic cells and macrophages. Pre-treatment of THP-1 monocytic cells, THP-1 monocyte-derived
macrophages and murine bone marrow derived macrophages with eppin prior to LPS stimulation, significantly decreased production of chemokines such as IL-8 and MIP-2. The anti-inflammatory effect of eppin was also demonstrated in an in vivo model of LPS-induced acute lung inflammation. Treatment of mice with recombinant human eppin resulted in a significant reduction in pulmonary neutrophilia and a concomitant reduction in BAL levels of the neutrophil chemoattractant MIP-2.

Investigation of a potential mechanism to explain these findings demonstrated that eppin reduced NF-κB activity in THP-1 monocytic cells in response to LPS stimulation. In contrast to SLPI [27], the anti-inflammatory effects of eppin were not dependent on binding of eppin to NF-κB DNA binding sites. However, similar to SLPI and elafin, we observed increased levels of phosphorylated IκBα in cells pre-treated with eppin. It is not clear why incubation of cells with eppin results in increased phosphorylated IκBα. Evaluation of proteasome activity has previously shown that the chymotrypsin, trypsin nor caspase-like activities of the proteasome are affected by SLPI or elafin [26, 28]. Therefore, it is possible that eppin affects the ubiquitin-proteasome pathway at other site(s) leading to increased IκBα phosphorylation, decreased NF-κB activity and decreased expression of NF-κB target genes. Moreover, we have demonstrated the presence of eppin in human tissue and airway secretions and have shown an immunomodulatory role for eppin in vivo. As previously discussed, SLPI and elafin play several important roles in the innate immune response. The expression of eppin in myeloid and epithelial cell types, coupled with its documented biological activity, suggest that eppin may play a protective role in the innate immune response similar to SLPI and elafin. The importance of this potential role remains to be fully
explored; however, eppin may represent another important multifaceted member of the WFDC family.

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Methods

Full details are available in the online supplement.

Cell culture

Unless otherwise indicated, cell culture reagents were obtained from Life Technologies Ltd. (Paisley, UK). Human acute monocytic leukemia cells (THP-1), myelomonocytic cells (U937) and type-II alveolar epithelial cells (A549) were purchased from the American Type Culture Collection (ATCC, Manassas, USA). Human diploid myeloid leukaemia cells (PLB-985) were obtained from the German Collection of Microorganisms and Cell Cultures (DSMZ, Braunschweig, Germany). The human bronchial epithelial cell line 16HBE14o- (HBE), the CF bronchial epithelial cell line CFBE41o- homozygous for the F508del mutation (CFBE), the human tracheal epithelial cell line 9HTEo- (HTE), and the CF tracheal epithelial cell line CFTE29o- homozygous for the F508del mutation (CFTE) were obtained as a gift from Prof. Dieter Gruenert (California Pacific Medical Centre Research Institute, San Francisco, CA) [33, 34]

Recombinant protein production

Recombinant eppin was produced as described previously [9].

RNA extraction and RT-PCR

RNA from human lung, testes tissue, primary peripheral blood mononuclear (PBMC) and polymorphonuclear leucocytes (PMN) cells were purchased from BioChain (Hayward, California, USA).
**Analysis of secreted eppin**

THP-1 monocytic cells were incubated in serum free media for 8 h in the absence or presence of EGF (5 ng/ml; Peprotech, London, UK). Supernatants were collected and protein precipitated using a method adapted from Wessel and Fugge [35].

**Human lung tissue**

Lungs harvested from potential donors that were found to be unsuitable for transplantation were obtained from IIAM (International Institute for Advancement of Medicine). Ethical approval for the use of these samples was obtained from the Queen’s University Belfast School of Medicine, Dentistry and Biomedical Science Research Ethics Committee. The lungs were perfused *ex vivo* based on previously described methods [36, 37]. Experimentation ceased 4 hr after LPS instillation, upon which tissue samples for histology were taken from LPS-perfused lobes [36]. Lung tissue sections from CF patients were obtained from the University of Newcastle. Ethical approval for the use of these tissue samples was obtained from Newcastle and North Tyneside Ethics Committee (reference number 11/NE/0291).

**Immunostaining of human tissue for eppin**

Immunostaining for eppin detection was performed similar to previous methods [36].

**Effect of recombinant eppin in a murine model of LPS-induced lung inflammation**

CS7Bl/6 mice (10 – 12 weeks of age) were used in all experiments and were purchased from Charles Rivers Laboratories and also bred in-house.
**Statistical analysis**

All data were analysed using GraphPad Prism 5.0 (GraphPad Software Inc., San Diego, CA). Means were compared by unpaired \( t \)-test, Mann Whitney test, one-way analysis of variance (ANOVA), Kruskal-Wallis test, or one-sample \( t \)-test compared with 100 for densitometry data as appropriate. \( P < 0.05 \) was accepted to indicate statistical significance. Data represent mean ± SEM of \( n = 3 \) unless otherwise stated.
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Figure Legends

Figure 1. Basal expression profile of eppin in cell lines and primary cells.
Eppin and GAPDH expression were detected in (A) THP-1 and U937 monocytc (mono) and monocyte-derived macrophage (mac), PLB-985 differentiated neutrophils (PLB), primary human neutrophils (PMN), human type II alveolar (A549) cells, testes and lung tissue, (B) human peripheral blood mononuclear cells (PBMC), human tracheal (HTE) and cystic fibrosis tracheal (CFTE), human bronchial (HBE) and cystic fibrosis bronchial epithelial (CFBE) cell lines by RT-PCR (35 cycles) as outlined in the Methods. Levels of eppin expression relative to GAPDH expression are shown between panels.

Figure 2. Detection of eppin in human lung tissue.
Tissue samples from (A) ex vivo non-LPS (B) LPS-perfused lungs and (C) cystic fibrosis lungs were formalin-fixed and paraffin-embedded for immunohistochemistry. Tissue sections were stained with 2 anti-human eppin antibodies from (i) Santa Cruz (ii) Abbexa and representative images ($n = 2$) were taken at X 20 objective lens magnification, scale bar = 50 µm. Further zoomed in images (to the right of the main panels) were also taken to highlight cellular staining of eppin. Areas of eppin staining are indicated by white arrows.

Figure 3. Detection of eppin in airway secretions
ARDS patient BALF samples, CF sputum and healthy control BALF samples were electrophoresed and blotted for eppin (Abbexa). Recombinant eppin (rEppin) was run in each blot and migrated as monomeric, dimeric and trimeric bands. Molecular weight
markers are indicated to the left of each blot and the position of eppin complex indicated with arrows.

**Figure 4. Eppin is up-regulated by EGF stimulation.**

(A) THP-1 monocytic cells were incubated with media alone or with EGF (5 ng/ml) for 8 h and eppin expression detected by RT-PCR (35 cycles) as outlined in the Methods. (B) Densitometry of eppin relative to GAPDH. * P < 0.05. (C) Supernatants from control and EGF-treated cells were concentrated by methanol chloroform precipitation and eppin was detected by Western blot. The identity of eppin was confirmed by 1D nLC-MS-MS (Reverse Phase Chromatography). The two main peptides obtained from this analysis were ANCLNTCK and DRQCQDNKK both of which demonstrate 100% overlap with human eppin.

**Figure 5. Eppin permeabilises Gram negative and Gram positive bacteria.**

(A) *P. aeruginosa* and (B) *S. aureus* were incubated for 2 h with 5 μM SYTOX® Green nucleic acid stain ± eppin or LL-37 as indicated. The proportion of permeabilised bacteria was quantified by measuring the relative fluorescence at 480 nm (excitation) and emission between 510 and 700 nm using a SPECTRAmax Gemini XS fluorescence microplate reader and results are expressed as a % of the pre-permeabilised positive control. (C) In radial diffusion assays, agarose gels were prepared containing *P. aeruginosa* or *S. aureus* and incubated with increasing concentrations of recombinant eppin (12.5-100μg/ml) or cecropin (100μg/ml) as a positive control.
Figure 6. Eppin inhibits LPS-induced cytokine production in mononuclear cells.

(A) THP-1 monocytic cells were pre-treated with recombinant human eppin (5 μg/ml) for 1 h prior to 6 hr LPS stimulation. Total RNA was extracted and RT-PCR performed for IL-8 and GAPDH as described in the Methods section. Expression levels of IL-8 relative to GAPDH are shown between the panels. (B) THP-1 monocytic cells were pre-treated with recombinant human eppin (5 μg/ml) for 1 h prior to LPS stimulation. Cell-free supernatants were collected after 24 h LPS stimulation and levels of IL-8, were quantified by ELISA. (C) THP-1 monocyte-derived macrophages and (D) murine bone marrow derived macrophages were pre-treated with recombinant human eppin (5 μg/ml) for 1 h prior to 24 h LPS stimulation. Levels of (C) human IL-8 and (D) murine MIP-2 were quantified by ELISA. *** P < 0.001 vs LPS. (E) THP-1 monocytic cells were incubated in cell medium or with recombinant human eppin (0.1, 1 and 10μg/ml) 24 h. Cell viability was assessed using Vision Blue Quick Cell Viability Assay reagent.

Figure 7. Eppin decreases LPS-induced NF-κB activation in THP-1 monocytic cells and increases levels of phosphorylated IκBα.

(A) THP-1 monocytic cells were pre-treated with recombinant human eppin (5 μg/ml) for 1 h followed by stimulation with LPS for 60 min. Nuclear lysates were prepared and p65 activity determined using a TransAm activity ELISA. * P < 0.05, *** P < 0.001 vs LPS. (B) The binding of eppin to the consensus NF-κB binding site was determined by EMSA. (C) THP-1 monocytic cells were pre-treated with recombinant human eppin (5 μg/ml) for 1 h followed by stimulation with LPS for 0-120 min. Cytoplasmic lysates were immunoblotted for IκBα, phospho-IκBα (Ser32/36) and GAPDH.
Figure 8. Recombinant eppin decreases LPS-induced neutrophil recruitment in a mouse model of acute lung inflammation.

Mice received intratracheal saline (sal) or LPS ± recombinant eppin and after 4 h mice were sacrificed and bronchoalveolar lavage (BAL) performed. BAL (A) Total cell counts and (B) neutrophil counts were quantified. (C) Levels of MIP-2 in BAL fluid were quantified by ELISA. $n = 3–7$/group, ** $P < 0.01$, *** $P < 0.001$ vs LPS.
FIGURE 1

A

Eppin

0.84 0.80 0.69 0.58 0.61 0.84
Eppin:GAPDH

0.42 1.06 0.49
GAPDH

Mono  Mac  Mono  Mac  PLB  PMN  A549  Testes  Lung  (-) RT  (-) cDNA
THP-1  U937

B

Eppin

0.25 0.18 0.18 0.26 0.23
Eppin:GAPDH

GAPDH

PBMC  HTE  CFTE  HBE  CFBE
FIGURE 4

A

Eppin

GAPDH

Con    EGF

B

Eppin:GAPDH (% of control)

0    50    100    150    200

Con    EGF

C

kDa

75

50

37.5

25

20

15

Con    EGF

Eppin Dimer
FIGURE 5

A

P. aeruginosa

Normalised Fluorescence

Pre-permeabilised bacteria  | Untreated  | Eppin 25 µg/ml  | LL-37 60 µg/ml

B

S. aureus

Normalised Fluorescence

Pre-permeabilised bacteria  | Untreated  | Eppin 50 µg/ml  | LL-37 125 µg/ml

C

eppin (µg/ml)

100  | 50  | 25  | 12.5

P. aeruginosa
ATCC 27853

S. aureus
ATCC 25923

Control
(100 µg/ml)