

# Legionella pneumophila-induced IκBζ-dependent expression of interleukin-6 in lung epithelium

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ABSTRACT: Severe community- and hospital-acquired pneumonia is caused by *Legionella* pneumophila. Lung airway and alveolar epithelial cells comprise an important sentinel system in airborne infections. Although interleukin (IL)-6 is known as a central regulator of the immune response in pneumonia, its regulation in the lung is widely unknown.

Herein, we demonstrate that different L. pneumophila strains induce delayed expression of IL-6 in comparison with IL-8 by human lung epithelial cells. IL-6 expression depended, at early time points, on flagellin recognition by Toll-like receptor (TLR)5, activity of mitogen-activated protein kinase/extracellular signal-regulated kinase kinase (MEK)1 and p38 mitogen-activated protein (MAP) kinase, and, at later time points, on the type-IV secretion system. In the same manner, but more rapidly, the recently described transcription factor  $I\kappa B\zeta$  was induced by Legionella infection and, binding to the nuclear factor (NF)- $\kappa B$  subunit p50 - recruited to the il6 promoter together with CCAAT-enhancer-binding protein  $\beta$  and phosphorylated activator protein-1 subunit cJun. Similarly, histone modifications and NF- $\kappa B$  subunit p65/ReIA appeared at the  $i\kappa b\zeta$  and subsequently at the il6 gene promoter, thereby initiating gene expression. Gene silencing of  $I\kappa B\zeta$  reduced Legionella-related IL-6 expression by 41%.

Overall, these data indicate a sequence of flagellin/TLR5- and type IV-dependent  $I\kappa B\zeta$  expression, recruitment of  $I\kappa B\zeta/p50$  to the il6 promoter, chromatin remodelling and subsequent IL-6 transcription in L. pneumophila-infected lung epithelial cells.

KEYWORDS: Cytokines, gene regulation, Legionella, pneumonia, signal transduction

n important causative agent of severe community-acquired pneumonia and the second most commonly detected pathogen in pneumonia in those admitted to intensive care units in industrialised countries is Legionella pneumophila [1, 2]. Although >40 Legionella species are known, the majority of human infections are caused by L. pneumophila serogroup 1 [3]. L. pneumophila is a Gram-negative, facultative intracellular pathogen of amoebae in natural and man-made aquatic environments. Infection of humans occurs after inhalation of contaminated water aerosol droplets. Recognition of Legionella by transmembraneous toll-like receptor (TLR)2 and TLR5 and cytosolic Naip5 (and possibly other unknown receptors) seems to activate the eukaryotic immune response [4–7].

With respect to *L. pneumophila* pathogenesis, essential results were obtained by analysing

infection of protozoans or immune cells like macrophages [8, 9]. However, although *Legionella* replicates efficiently within lung epithelial cells and recent studies pointing to the lung epithelium as an important sentinel and effector of innate immunity [7, 10–16], little is known of the consequences of pulmonary epithelial cell infection with *Legionella*.

Interleukin (IL)-6 concentrations in blood and bronchoalveolar lavage fluid of patients suffering from pneumonia are positively associated with disease severity [17–19] and IL-6 promoter polymorphisms are associated with extrapulmonary dissemination in pneumococcal pneumonia [20]. Studies using IL-6-deficient mice further indicated a prominent role of the cytokine IL-6 in pneumonia [21, 22]. IL-6 influenced important innate immune mechanisms by, for example, reducing neutrophil apoptosis and increasing

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European Respiratory Journal Print ISSN 0903-1936 Online ISSN 1399-3003 their cytotoxic capabilities [23, 24]. Its expression is regulated by several transcription factors, including nuclear factor (NF)- $\kappa$ B, activator protein (AP)-1, and CCAAT-enhancer-binding protein (C/EBP) [25, 26]. More recently, a critical role of inducible nuclear protein  $I\kappa$ B $\zeta$  for IL-6 expression was demonstrated [27].

Considering the important role of IL-6 in pneumonia, we here analysed mechanisms of *Legionella*-related IL-6 expression in human lung epithelial cells. *L. pneumophila* induced expression of IL-6 in a flagellin-TLR5- and type-IV secretion system-dependent manner. NF- $\kappa$ B-related expression of I $\kappa$ B $\zeta$  was critical for IL-6 expression in *Legionella*-infected cells. *Legionella*-induced histone modifications and subsequent recruitment of p50 and p65/RelA to the *il6* and  $i\kappa$ b $\zeta$  gene promoter to initiate gene transcription. At the *il6* promotor, I $\kappa$ B $\zeta$  and p50 interact in *Legionella*-infected cells and gene silencing of I $\kappa$ B $\zeta$  reduced *Legionella*-related IL-6 expression. Overall, *L. pneumophila* flagellin activated the expression of I $\kappa$ B $\zeta$  and subsequent transcription of IL-6 in human lung epithelial cells.

#### **MATERIALS AND METHODS**

#### Materials

Fetal calf serum (FCS) and trypsin-EDTA solution were obtained from Life Technologies (Karlsruhe, Germany). Pyrrolidine dithiocarbamate (PDTC), protease inhibitors, NP-40, Triton X-100 and Tween-20 were purchased from Sigma Chemical (Munich, Germany); MG-132 was from Calbiochem-Merck (Darmstadt, Germany) and recombinant flagellin from Alexis (Lörrach, Germany). All other chemicals used were of analytical grade and obtained from commercial sources.

#### **Cell lines**

Type-II alveolar cell line A549 was obtained from ATCC (Rockville, MD, USA) and cultured in Ham's F-12 medium with L-glutamine (PAA Laboratories, Pasching, Austria) and 10% FCS without antibiotics [28]. Primary human small airway epithelial cells (SAEC) were obtained from Clonetics/Cambrex (SAEC System; Cambrex, Baltimore, MD, USA) and cultured in SAEC BulletKit® (Clonetics/Cambrex) according to the supplier's instruction [15].

#### Infection with bacterial strains

L. pneumophila sg1 130b wildtype (ATCC BAA-74, kindly provided by N.P. Cianciotto, Northwestern University Medical School, Chicago, IL, USA) [29], JR32 wildtype [30] and JR32  $\Delta dot A$  mutant (LELA 3118, both kindly provided by H. Shuman, Columbia University, New York, NY, USA) [31], Corby wildtype and Corby  $\Delta flaA$  mutant (both kindly provided by K. Heuner, Berlin, Germany) were routinely grown on buffered charcoal-yeast extract (BCYE) agar for 2 days at 37°C [32] and subsequently resuspended in epithelial cell medium at 37°C. Bacterial density was checked by determining the optical density at 660 nm (OD<sub>660</sub>) with a Beckman spectrophotometer DU520 (Beckman Coulter, Unterschleissheim, Germany). A549 or SAEC (106) were infected with 10<sup>5</sup>, 10<sup>6</sup> and 10<sup>7</sup> colony-forming units (CFU) bacteria, i.e. a multiplicity of infection of 0.1:1-10:1, per milliliter and incubated in infection medium (Ham's F-12 with L-glutamine without antibiotics or SAEC medium) for a given time at 37°C and 5% CO<sub>2</sub>. Extracellular bacteria were not routinely killed with antibiotics. *L. pneumophila* strains did not significantly grow in epithelial cell growth medium as controlled by serial dilutions were plated on BCYE agar. For heat-inactivating *L. pneumophila*, bacteria were incubated at 95°C for 30 min [14].

#### IL-6 ELISA

Confluent A549 cells or SAECs were infected as indicated in a humidified atmosphere. After incubation, supernatants were collected and processed for IL-6 quantification by immunoassay according to the manufacturer's instructions (R&D Systems, Minneapolis, MN USA).

#### Chromatin immunoprecipitation (ChIP)

A549 cells were infected with *L. pneumophila* as indicated and then subjected to ChIP assay as previously described [14, 28, 33, 34] using anti-mRNA polymerase II (Pol II) (N-20), anti-p65 (C-20), anti-p50, anti-C/EBPβ (Santa Cruz Biotechnologies, Santa Cruz, CA, USA), anti-P-cJun (Cell Signalling Technology, Danvers, MA, USA), anti-P<sup>Ser10</sup>/Ac<sup>Lys14</sup>-H3, Pan-Ac-H4 (Upstate, Lake Placid, NY, USA) and anti-Iκβζ (Acris Antibodies GmbH, Hiddenhausen, Germany). The il6 and iκbζ enhancer region was amplified by PCR using HotStarTaq polymerase (Qiagen, Hilden, Germany) and specific primers as followed: il6 sense 5'-ACAAATTAACTGGAACGCT-3', antisense 5'-ATTGGGGG TTGAGACTCTAA-3' and iκbζ sense 5'-AGGGGAATGTCC GGGACT-3', antisense 5'-TAATGTCTGACCTCGTGGCAA-3'. PCR amplification of total input DNA in each sample is shown as a control.

For Chromatin-IP-IP, DNA was eluted after first IP, refilled with ChIP-RIPA buffer and incubated with anti-I $\kappa$ B $\zeta$  and anti-p50 antibodies, respectively for the second IP. As a control an IP with anti-sheep immunoglobulin (Ig)G (Li-COR) was performed.

#### Western blot

For determination of IκBζ and COX-2 induction, A549 or SAE cells were infected as indicated, washed twice, and nuclear (IκΒζ, actin), membrane (TLR5) or whole cell extract (COX-2, ERK2) were harvested. Cells were lysed in buffer containing NP-40 or Triton X-100, subjected to SDS-PAGE, and blotted on nitrocellulose membrane (Amersham Biosciences, Freiburg, Germany). Immunodetection of target proteins was carried out with specific antibodies: ΙκΒζ (Acris Antibodies GmbH, Hiddenhausen, Germany), COX-2 (Santa Cruz Biotechnologies, Santa Cruz, USA) and subsequently incubated with secondary antibodies (IRDye 800-labeled anti-mouse or Cy5.5-labeled antirabbit, respectively, and IRDye 800-labeled anti-goat) (Li-COR Inc., Bad Homburg, Germany). Simultaneous detection of actin or ERK2 (Santa Cruz Biotechnologies) by using an Odyssey infrared imaging system (Li-COR Inc.) confirmed equal protein load as described [28, 35].

#### RT-PCR

For analysis of IkB $\zeta$ , IL-6, IL-8, granulocyte macrophage colony-stimulating factor (GM-CSF) and GAPDH gene expression in A549 and SAECs, total RNA was isolated with the RNeasy Mini kit (Qiagen) and reversely transcribed using moloney murine leukaemia virus reverse transcriptase (Invitrogen, Karlsruhe, Germany). Generated cDNA was amplified by PCR using specific primers for IkB $\zeta$  (forward:



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5'-TGAATGCACTTCACATGCTG-3'; reverse: 5'-TTCGTTCT CCAAGTTCCGAGT-3'), IL-6 (forward: 5'-TTCTCCACAAGC GCCTTC-3'; reverse: 5'-TGGACTGCAGGAACTCCTTA-3'), IL-8 (forward: 5'-CTAGGACAAGACCAGGAAGA-3'; reverse: 5'-AACCCTCTCTGCACCCAGTTTTC-3'), GM-CSF (forward: 5'-GTCTCCTGAACCTGAGTAGAGACA-3'; reverse: 5'-AAGG GGATGACAAGCAGAAAGTCC-3'), GAPDH (forward: 5'-C CACCCATGGCAAATTCCATGGCA-3'; reverse: 5'-TCTAGA CGGCAGGCAGGTCAGGTCCACC-3'). All primers were purchased from TIB MOLBIOL (Berlin, Germany). After 20–35 amplification cycles, PCR products were analysed on 1.5% agarose gels, stained with ethidium bromide, and subsequently visualised. To confirm use of equal amounts of RNA in each experiment, all samples were checked for GAPDH mRNA expression.

#### RNA interference

#### Statistical methods

Data are presented as mean  $\pm$  SE of at least three independent experiments. Effects were statistically evaluated employing paired t-tests. A p-value <0.05 was considered to be significant (if not indicated otherwise, test was performed *versus* control).

#### **RESULTS**

## L. pneumophila-induced expression of IL-6 and $I\kappa B\zeta$ in human lung epithelial cells

Infection of human lung epithelial cell line A549 (fig. 1a and fig. S1A in the supplementary material) with *L. pneumophila* strain 130b, strain Corby (fig. 2a and b), and strain JR32 (fig. 3a and b) as well as of primary human lung SAEC with *L. pneumophila* strain 130b (fig. 1a and fig. S1B in the supplementary material) induced the expression of IL-6 as well as of IL-8 mRNA. IL-6 protein liberation was shown in *L. pneumophila* 130b- (fig. 1c), Corby- (fig. 2c and d), and JR32-infected (fig. 3c) A549 and 130b-exposed SAEC (fig. 1d and e). Heat-inactivation of *L. pneumophila* 130b reduced, but did not block, liberation of IL-6 in infected A549 cells (fig. 1c).

As the mRNA expression of IL-6 was observed later than that for IL-8, a possible role for IkB $\zeta$  protein expression in the regulation of IL-6 expression was considered, and IkB $\zeta$  expression was analysed. Early induction of IkB $\zeta$  mRNA (fig. 1a) and protein (fig. 1b) was observed before the start of IL-6 mRNA expression in *L. pneumophila* 130b-infected A549 and SAEC (fig. 1a) as well as in Corby- (fig. 2a and b) and JR32-infected (fig. 3a and b) A549 cells. Expression of IkB $\zeta$ , IL-6

and IL-8 mRNA persisted for 24 h (fig. 1a). Thus, different strains of *L. pneumophila* induced both, IκΒζ and IL-6 expression in human lung epithelial cells in a specific temporal order.

## L. pneumophila-induced expression of IL-6 and $I\kappa B\zeta$ dependent on flagellin-related activation of TLR5 and on the Legionella type-IVB Dot/Icm secretion system

Detection of Legionella flagellin by TLR5 seems to be important for activation of eukaryotic host cells [5, 14]. Indeed, Legionella deficient for flagellin (FlaA) showed to be less potent with respect to the induction of mRNA for IκBζ and IL-6 as well as of IL-8, another important pro-inflammatory cytokine in pneumonia [36], especially at early time points (fig. 2b). In addition, liberation of IL-6 protein was significantly lower in cells infected with flagellin-deficient Legionella compared with wildtype bacteria (fig. 2c and d). As published previously, FlaA-deficient Legionella replicated twice as well as wildtype bacteria over 24 h in lung epithelial cells but did not alter cell death as determined by lactate dehydrogenase release (data not shown and VINZING et al. [37]). Addition of recombinant flagellin to cell cultures demonstrated to be sufficient to induce expression of mRNA of IκBζ, IL-6 as well as of IL-8 in human lung epithelial cells (fig. 4a). Finally, exogenously added recombinant flagellin induced IL-6 protein secretion (fig. 4b). Suppression of endogenous TLR5 expression by specific siRNA (fig. 4c) significantly reduced flagellin-related IL-6 liberation in A549 cells (fig. 4d). Overall, these data indicate a critical role of TLR5-related detection of Legionella flagellin for IκBζ and IL-6 expression.

Type-IVB Dot/Icm secretion system-related activity is known to be important for cytosolic recognition of *Legionella* flagellin and subsequent cell activation [38]. Upon infection of A549 cells with a *L. pneumophila* JR32 mutant deficient for *dotA*, an integral part of the *Legionella* type-IVB system [8], we observed a similar induction of IkB $\zeta$ , IL-6, and IL-8 mRNA at early time points compared with infection of cells with JR32 wildtype, but a clear reduction after 5 h (fig. 3a and b) as well as reduced liberation of IL-6 (fig. 3c). Overall, intracellular, type-IV-dependent recognition of *Legionella* flagellin seems to be important for the persistent induction of IkB $\zeta$  and IL-6 in A549 lung cells.

## NF- $\kappa B$ -related transcription of cytokines and $I\kappa B\zeta$ in L. pneumophila-infected epithelial cells

Since NF-κB seems to be central for the expression of IκBζ and important pro-inflammatory cytokines, including IL-6, two unrelated inhibitors of the NF-κB pathway, PDTC and MG-132 [39, 40], were tested with respect to *Legionella*-related cell activation. On mRNA level, both inhibitors reduced the expression of IκBζ, IL-6, GM-CSF, and IL-8 (fig. 5a and c). Moreover, pre-incubation of cells with the inhibitors significantly reduced *L. pneumophila* 130b-induced IL-6 (fig. 5b and d) and IL-8 (data not shown) protein expression as demonstrated by ELISA indicating the central role of NF-κB pathway for *Legionella*-related epithelial cell activation.

### Impact of Legionella-related signalling on the iκbζ and il6 gene promoter

Since our data indicated an important role of  $I\kappa B\zeta$  and NF- $\kappa B$  in IL-6 expression, we made use of ChIP to assess the

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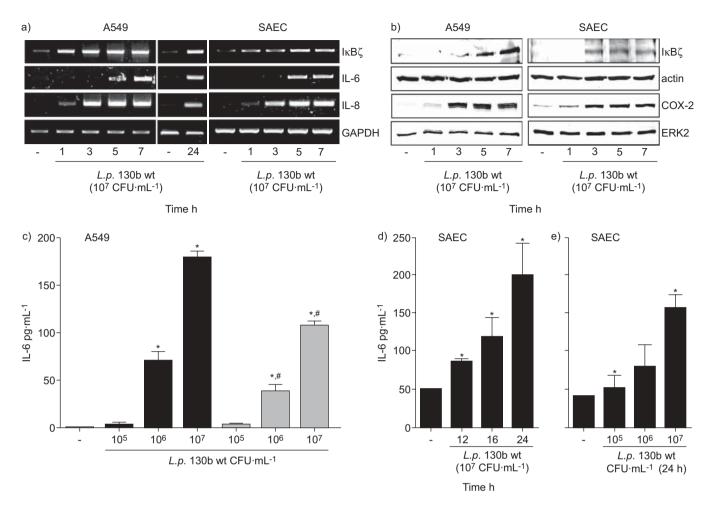


FIGURE 1. Legionella pneumophila-induced expression of interleukin (IL)-6 and I<sub>κ</sub>Bζ in human lung epithelial cells dependent on bacterial viability. Alveolar epithelial cells (A549) or human primary small airway epithelial cells (SAEC) were infected with *L. pneumophila* wildtype strain 130b ( $10^7$  CFU·mL<sup>-1</sup>) for a, b, d) the indicated times or c, e) with the indicated concentrations for 24 h. a) mRNA levels of I<sub>κ</sub>Bζ, IL-6, IL-8, and GAPDH were detected by RT-PCR, and b) protein expression of I<sub>κ</sub>Bζ, COX-2, actin and ERK2 were detected by western blot. Representative gels/blots of three independent experiments are shown. (c) Alveolar epithelial cells (A549) were infected with the indicated concentrations of viable ( $\blacksquare$ ) or heat-inactivated ( $\blacksquare$ ) *L. pneumophila* wildtype strain 130b for 24 h and IL-6 release was determined in the supernatant. Human primary SAEC were infected with *L. pneumophila* wildtype strain 130b ( $10^7$  CFU·mL<sup>-1</sup>) for d) the indicated times or e) with the indicated concentrations for 24 h, and IL-6 release was determined in the supernatant. \*: p<0.05 unstimulated versus stimulated cells; \*\*: p<0.05 viable versus heat-inactivated bacteria.

endogenous  $i\kappa b\zeta$  and il6 gene promoters in more detail in L. pneumophila-infected cells (fig. 6). At the  $i\kappa b\zeta$  gene promotor, Legionella infection resulted in phospho-acetylation of histone H3 ( $P^{Ser10}/Ac^{Lys14}$ ) and pan-acetylation of H4 indicating genetranscription promoting remodeling of the gene (fig. 6a). Recruitment of NF- $\kappa$ B subunits p50 and p65 was observed and accompanied by binding of Pol II indicative for start of gene transcription (fig. 6a). At the il6 gene promoter, we noted the same pattern of histone modification and recruitment of p50- and p65-NF- $\kappa$ B subunits (fig. 6b). Moreover, recruitment of I $\kappa$ B $\zeta$  protein as well as of Pol II to the il6 gene promoter was documented in L. pneumophila-infected epithelium.

Since Yamazaki *et al.* [41] evidenced strong interaction of IkB $\zeta$  with p50, but weak binding to p65, we tested the interaction of both proteins (IkB $\zeta$  and p50) at the *il6* gene promoter in *Legionella*-exposed cells. In ChIP-IP, experiments we found an association of endogenous p50 and IkB $\zeta$  proteins indicating cooperative action of both molecules at the *il6* gene promoter (fig. 6c).

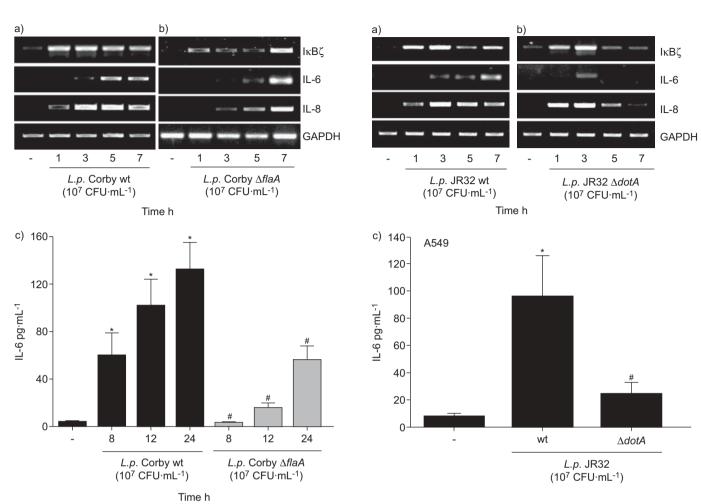
Finally, we wondered if a defect in *Legionella* flagellin expression would result in reduced stimulation of transcription factor and Pol II recruitment to both gene promoters. As it is shown in fig. 6d, flagellin-deficient *Legionella* induced weaker p65, p50 and Pol II recruitment to the  $i\kappa b\zeta$  gene promoter. Along those lines, reduced p65, p50, and IkB $\zeta$  binding and Pol II recruitment was also demonstrated at the il6 gene promoter (fig. 6e). Quantification of the gels in figure 6d and e showed a statistically significant decrease for Pol II as well as for all transcription factors (data not shown).

## Depletion of $I\kappa B\zeta$ reduced IL-6 expression in L. pneumophila-infected epithelial cells

To further analyse the role of IkB $\zeta$  in *Legionella*-related IL-6 expression, we made use of IkB $\zeta$  siRNA. IkB $\zeta$ -specific, but not control siRNA, reduced IkB $\zeta$  as well as IL-6 mRNA expression in *L. pneumophila* infected A549 cells (fig. 7a). Furthermore, we observed significant reduction of IL-6 protein expression (fig. 7b). The relatively low effect on the protein level may be



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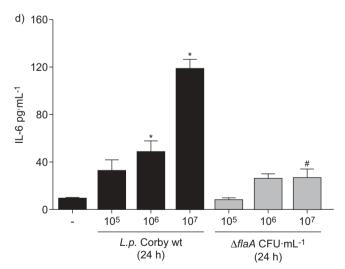


FIGURE 2. Legionella pneumophila induced expression of interleukin (IL)-6 and IκΒζ dependent on flagellin. a) Alveolar epithelial cells (A549) were infected with *L. pneumophila* wildtype strain a) Corby or b) ΔflaA-knock out mutant (10<sup>7</sup> CFU·mL<sup>-1</sup>) for the indicated times and mRNA expression was determined by RT-PCR. Representative gels of three independent experiments were shown. A549 cells were infected with *L. pneumophila* wild type strain Corby (■) or ΔflaA-knock out mutant (■)(10<sup>7</sup> CFU·mL<sup>-1</sup>) for c) the indicated times or d) with the indicated concentrations for 24 h, and IL-6 release was determined in the supernatant. \*: p<0.05 unstimulated *versus* stimulated cells; #: p<0.05 Corby *versus* CorbyΔflaA.

**FIGURE 3.** Legionella pneumophila-induced expression of interleukin (IL)-6 and lκBζ dependent on the Legionella type-IVB Dot/lcm secretion system. a) Alveolar epithelial cells (A549) were infected with L. pneumophila a) wildtype strain JR32 or b)  $\Delta dotA$ -knock-out mutant ( $10^7$  CFU·mL $^{-1}$ ) for the indicated times and mRNA expression was determined by RT-PCR. Representative gels of three independent experiments were shown. c) A549 cells were infected with L. pneumophila wild type strain JR32 or  $\Delta dotA$ -knock out mutant ( $10^7$  CFU·mL $^{-1}$ ) for 24 h and IL-6 release was determined in the supernatant. \*: p<0.05 unstimulated versus stimulated cells; #: p<0.05 L. pneumophila JR32 versus JR32 $\Delta dotA$ .

due to limited transfection efficiency and thus be overcome by siRNA-unaffected cells. To rule out effects of incomplete nucleofection, we created A549 cells stably expressing unspecific or IkB $\zeta$ -specific shRNA. Cells were infected with L. pneumophila wildtype strain 130b ( $10^7$  CFU·mL $^{-1}$ ) for 24 h and IL-6 release was detected by ELISA (fig. 7c). In IkB $\zeta$ -depleted cells, L. pneumophila-induced IL-6 release was reduced by 41% in comparison to cells expressing unspecific shRNA.

## Inhibition of MEK1- and p38 MAP kinase pathway blocked expression of IL-6 but not $I\kappa B\zeta$

Besides IkB $\zeta$  and the canonical NF-kB pathway, several kinase pathways were suspected to be important for bacteria-related activation of eukaryotic cells [10, 14, 42]. In particular, their role in *Legionella*-related IkB $\zeta$  and IL-6 expression is unknown. Therefore, we made use of several chemical inhibitors for important kinases to test their role in IkB $\zeta$ -, and cytokine-expression in *Legionella* infected cells (fig. 8a and fig. S2 in the

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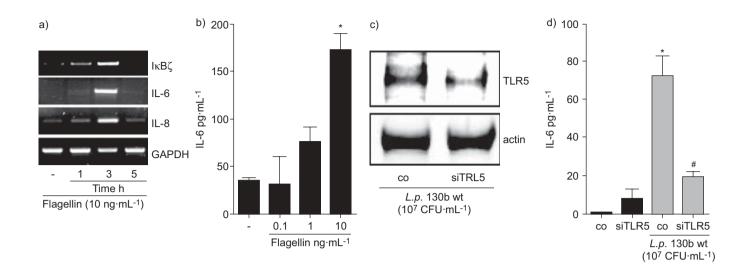


FIGURE 4. Legionella pneumophila induced expression of interleukin (IL)-6 and IκΒζ activation of Toll-like receptor (TLR)5. A549 cells were stimulated with flagellin (10 ng·mL<sup>-1</sup>) for a) the indicated times and mRNA expression was determined, or b) with the indicated concentrations for 24 h, and IL-6 release was determined in the supernatant. A549 cells were nucleofected with unspecific (co) or TLR5-specific (siTLR5) siRNA and after 48 h infected with *L. pneumophila* wildtype strain 130b (10<sup>7</sup> CFU·mL<sup>-1</sup>) for 24 h. c) TLR5 expression was determined by western blot and d) IL-6 release by ELISA with (III) and without (IIII) *L. pneumophila* wildtype strain 130b (10<sup>7</sup> CFU·mL<sup>-1</sup>). \*: p<0.05 unstimulated *versus* stimulated cells; #: p<0.05 unspecific *versus* TLR5 siRNA. In c), representative blots of three independent experiments were shown.

supplementary material). Inhibition of MEK1 kinase (U0126, 10  $\mu\text{M})$  and p38 MAP kinase (SB202190, 10  $\mu\text{M})$  both blocked L. pneumophila-related IL-6, GM-CSF and IL-8 expression. Blocking of PI3 kinase (Ly294002, 10  $\mu\text{M})$  and JAK isoforms 1–3 (JAK inhibitor I, 10  $\mu\text{M})$  prevented expression of IL-6 and GM-CSF, but not of IL-8 in Legionella-infected cells. However, inhibitors of PKC (Gö6976, 10  $\mu\text{M})$  and JNK (SP600125, 10  $\mu\text{M})$  only reduced GM-CSF expression. Remarkably, neither inhibitor showed an effect on L. pneumophila-induced IkB $\zeta$  expression

in lung epithelial cells although effective kinase inhibition was indicated by effects of the inhibitors on the addressed cytokines.

Transcription factors C/EBP $\beta$  and AP-1 have been implicated in IL-6 expression and are downstream targets of MEK1 and p38 MAP kinase, respectively [43]. After infection with *L. pneumophila*, an early recruitment of both C/EBP $\beta$  and the phosphorylated form of the AP-1 subunit cJun occurred at the *il6* promoter (fig. 8b and c). Inhibition of MEK1 blocked

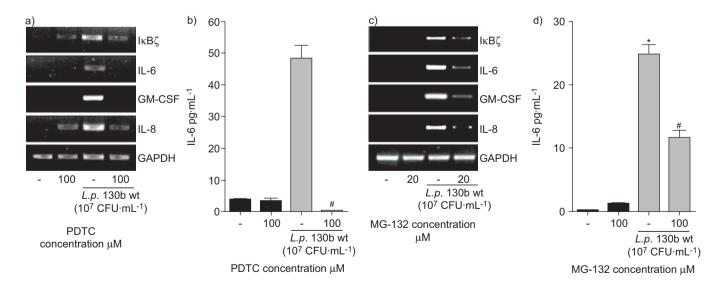
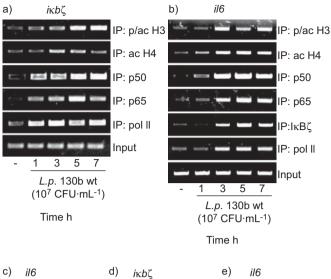
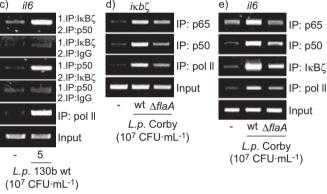


FIGURE 5. Nuclear factor (NF)-κB-related transcription of cytokines and IκB $\zeta$  in Legionella pneumophila-infected epithelial cells. Alveolar epithelial cells (A549) were preincubated for 30 min with inhibitors a, b) PDTC (100 μM) or c, d) MG-132 (indicated concentration) and then infected with *L. pneumophila* wildtype strain 130b (10<sup>7</sup> CFU·mL<sup>-1</sup>)( $\blacksquare$ ). After 7 h, mRNA expression was determined by RT-PCR (a and c). a, c) Representative gels of three independent experiments are shown. After 24 h, interleukin (IL)-6 release was determined in the supernatant (b and d). \*: p<0.05 unstimulated *versus* stimulated cells; #: p<0.05 cells with *versus* without inhibitor.







**FIGURE 6.** Impact of Legionella-related signalling on the  $i\kappa b\zeta$  and il6 gene promoter. A549 cells were infected with  $10^7$  CFU·mL<sup>-1</sup> Legionella pneumophila strains 130b (a–c) or Corby wildtype (wt)/ $\Delta flaA$ -knock out deletion mutant ( $\Delta flaA$ ) (d and e) for the indicated times (a and b), or for 5 h (c–e). Chromatin immunoprecipitation (ChIP) was performed against the indicated targets on the endogenous promoters of il6 (b, c and e) or  $i\kappa b\zeta$  (a and d). Representative gels of three independent experiments are shown.

 $\it L.~pneumophila\mbox{-induced}$  C/EBP $\beta$  recruitment to the  $\it il6$  promoter (fig. 8b).

#### **DISCUSSION**

Time h

Here we demonstrate that *L. pneumophila* induced flagellindependent IkB $\zeta$  expression and subsequent IL-6 expression in human lung epithelial cells. *Legionella* induced histone modifications and recruitment of p50 and p65/RelA to the  $i\kappa b\zeta$  and subsequently to the il6 gene promoter to initiate gene transcription. Both, IkB $\zeta$  and p50 interact at the il6 gene promotor in *Legionella*-infected cells and gene silencing of IkB $\zeta$  reduced *Legionella*-related IL-6 expression.

Little is known about the molecular mechanisms of host-pathogen interaction in the lung. This limits the development of innovative intervention strategies in pneumonia despite the emergence of antibiotic-resistant bacteria complicating antibiotic therapy. Therefore, we aimed to elucidate mechanisms of IL-6 expression in *Legionella* infection.

Expression of the pleiotropic cytokine IL-6 during pneumonia is suspected to contribute to the local control of infection and inflammation [20–22] by regulating for example neutrophil recruitment and function [23, 24]. The development of interstitial pneumonia in transgenic mice overexpressing human IL-6 highlights the power of IL-6 to contribute to inflammatory processes in the lung [44]. Infection of lung epithelial cells by *L. pneumophila* strains 130b, JR32 and Corby (this study, [15, 45]) as well as of murine macrophages [46] resulted in strong release of IL-6.

Detection of *Legionella* flagellin by host cell pattern recognition receptors (PRR) seems to be a critical step in legionellosis. At early infection timepoints, flagellin is recognised by transmembraneous TLR5 [47] and a common dominant TLR5 stop codon polymorphism is associated with susceptibility to Legionnaires' disease in humans [5]. Recombinant flagellin induced IL-6 expression, whereas flagellin-deficient L. pneumophila showed reduced IL-6 induction and TLR5-depletion in eukaryotic cells reduced Legionella-related IL-6 expression indicating flagellin-TLR5-related induction of IL-6. In addition, cytosolic recognition of flagellin monomers by Naip5 in macrophages contributes to the restriction of L. pneumophila infection in mice [38]. Noteworthy, the difference of mRNA levels of IκBζ and of IL-6, induced by wildtype or flaA-deletion mutants disappear after 7 h. As the difference in IL-6 protein release persists for 24 h, an additional mechanism, e.g. in translation or release of IL-6 might be involved. In addition, a role of TLR2 can not be ruled out from these data [48].

Flagellin is suspected to reach the host cell cytoplasm by type-IV secretion system-induced perforation of phagosomes [38], and there activating, for example, NAIP5 or IPAF [37]. In agreement, we observed reduced IL-6 mRNA expression at later time points in cells infected with *Legionella* strain JR32 lacking DotA, an integral part of the type-IVB system in these bacteria [8]. Overall, extracellular and possibly intracellular recognition of *L. pneumophila* seems to be important for IL-6 expression in lung epithelium. Interestingly, *Legionella longbeachae* (the common cause of Legionellosis in Western Australia) which contains flagellin but no detectable poreforming activity, does not activate Naip5-related activity, thus implicating possible important strain-specific effects in *Legionella*-related cell activation [49].

Expression of IL-6 is regulated by several transcription factors, including NF-κB, AP-1, and C/EBP [25, 26]. NF-κB-related gene expression is observed in L. pneumophila-infected lung cells [7, 11, 15]. In addition, transcription factors AP-1/cJun and C/EBPB are recruited to the il6 promoter after infection with L. pneumophila. However, much less is known about the role of the inducible nuclear protein  $I\kappa B\zeta$  in lung infection, which seems to be critical for IL-6 expression [27]. We observed induction of IκBζ in L. pneumophila 130b-, Corby-, and JR32-infected A549 and primary human lung epithelial cells. Its expression persisted for 24 hours. In line with the study of YAMAMOTO et al. [27], experiments using recombinant flagellin, and flagellin-deficient L. pneumophila indicated flagellin-related induction of IkBÇ in lung epithelium. Analysis of host cell signalling pathways demonstrated NF- $\kappa B$  dependency of  $I\kappa B\zeta$  expression. However, although inhibition of MEK1, p38, JAK or PI3 kinase suppressed IL-6

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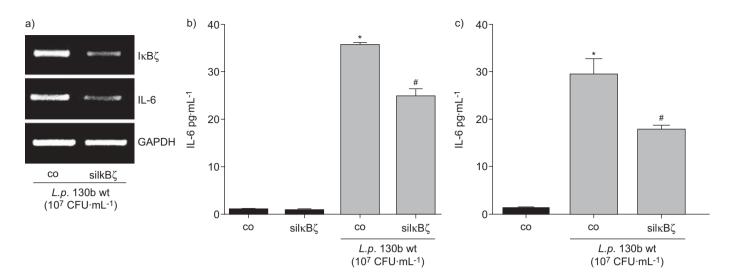
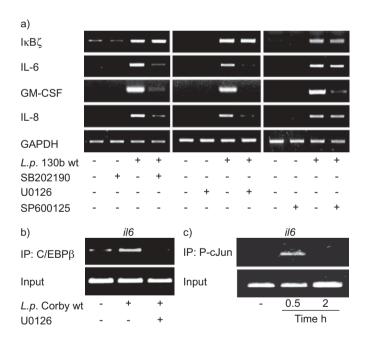


FIGURE 7. Depletion of IκΒζ reduced interleukin (IL)-6 expression in Legionella pneumophila-infected epithelial cells. (a, b) A549 cells were nucleofected with unspecific (co) or IκΒζ-specific (silκΒζ) siRNA and after 36 h infected with *L. pneumophila* wildtype strain 130b (10<sup>7</sup> CFU·mL<sup>-1</sup>) for 24 h. IκΒζ expression was determined by a) RT-PCR and b) IL-6 release by ELISA. c) A549 stably expressing unspecific (co) or IκΒζ-specific shRNA (shIκΒζ) were infected with *L. pneumophila* wt strain 130b (10<sup>7</sup> CFU·mL<sup>-1</sup>) for 24 h and IL-6 release was detected by ELISA. \*: p<0.05 unstimulated versus stimulated cells; #: p<0.05 unspecific versus IκΒζ si/shRNA. In a), representative gels of three independent experiments were shown.



**FIGURE 8.** Expression of interleukin (IL)-6, but not IκBζ mRNA, depended on p38 MAP kinase and MEK1. a) Alveolar epithelial cells (A549) were preincubated for 120 min with inhibitors SB202190 (10 μM), U0126 (10 μM) or SP600125 (10 μM) and then infected with *L. pneumophila* (*L. p.*) wildtype strain 130b (10<sup>7</sup> CFU·mL<sup>-1</sup>). After 7 h, mRNA expression was determined by RT-PCR. Representative gels of three independent experiments were shown. A549 cells were infected with  $10^7$  CFU·mL<sup>-1</sup> *L. pneumophila* Corby for 30 min b) after preincubation for 120 min with inhibitor U0126 (10 μM) or c) for the indicated times without preincubation. ChIP was performed against the indicated targets on the endogenous promoter of *il6*. Representative gels of three independent experiments are shown.

expression, we observed no effect on  $I\kappa B\zeta$  expression. In addition, *L. pneumophila*-induced recruitment of C/EBP $\beta$  to the *il6* promoter was blocked by MEK1 inhibitor U0126. These data suggest that kinase-related,  $I\kappa B\zeta$ -independent signals also contributed to IL-6 expression in *L. pneumophila*-infected epithelium. In addition, PKC and JNK pathway neither contributed to IL-6 nor  $I\kappa B\zeta$  expression, but reduced *Legionella*-induced GM-CSF expression. Both kinases have been shown to play important roles in the activation of epithelial cells and macrophages, respectively [11, 50].

Upon infection of cells with *L. pneumophila*, histone H3 gets phosphoacetylated and H4 acetylated at the  $i\kappa b\zeta$  gene promoter, indicative of chromatin remodeling associated with transcriptional activity [14, 33, 34, 51]. This was accompanied by recruitment of p50 and p65 NF- $\kappa$ B subunits, which are known to be important for  $I\kappa B\zeta$  expression [52]. In addition to these modifications and recruitments also documented at the *il6* gene promoter, we observed  $I\kappa B\zeta$  binding at the *il6* gene promoter in *L. pneumophila*-infected epithelial cells.

Recently, Kayama *et al.* [52] demonstrated that LPS-induced histone H3 trimethylation on a subset of promoters depended on IkB $\zeta$  in mouse macrophages. Together with our results, it seems possible that IkB $\zeta$  is involved in TLR- or NLR-induced nucleosome remodeling regulating the expression of secondary genes. This might also explain the species- and cell type-dependent degree of IkB $\zeta$ -dependence [27].

IκBζ seems to preferentially interact with NF-κB subunit p50 [41]. In fact, interaction of p50 with IκBζ was observed at the il6 gene promoter in Legionella-exposed cells by Chromatin-IP-IP. Silencing of IκBζ expression by specific siRNA or shRNA resulted in decreased, but not completely abolished, IL-6 expression in L. pneumophila-infected cells, suggesting that IκBζ plays an important but not indispensable role in IL-6 expression in the system tested.



Finally, flagellin-deficient L. pneumophila induced weaker transcription factor recruitment and Pol II binding at both, the  $i\kappa b\zeta$  and il6 gene promoter, further highlighting the role of flagellin in activation of gene transcription. Overall, these data indicate a sequence of flagellin-dependent chromatin remodeling, p50, p65, and Pol II recruitment,  $I\kappa B\zeta$  expression, and subsequent IL-6 expression in L. pneumophila-infected lung epithelial cells. In addition, other pathways like MEK-1-C/EBP $\beta$  and p38 MAP kinase-AP-1 as well as the canonical NF- $\kappa$ B pathway seem to be involved in IL-6 expression.

From the observation of the complex nature of *Legionella*-related IL-6 expression several questions arise: For example, which cytosolic receptors contribute to IkB $\zeta$  and IL-6 expression? In addition, besides the contribution of NF-kB molecules, the signalling mechanisms leading to IkB $\zeta$  expression are widely unknown and different kinase pathways could be excluded in this study. Besides *L. pneumophila*, other important pathogens which do not express flagella (*e.g.* pneumococci) cause pneumonia and induce IL-6 expression [42]. Furthermore, although *e.g.* the important role of IL-6 for pneumococci dissemination [21, 22] is known, underlying molecular mechanisms of pneumococci-related IL-6 (and potentially IkB $\zeta$ ) expression are unknown. Finally, *in vivo* studies with IkB $\zeta$  deficient mice will help to dissect its role in pneumonia.

In conclusion, our data highlighted a complex pathway of IL-6 induction in human lung epithelium, composed of early flagellin-TLR5-dependent histone modifications,  $I\kappa B\zeta$  expression, and subsequent  $I\kappa B\zeta$ -p50-related IL-6 expression.

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#### STATEMENT OF INTEREST

None declared.

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