

# Bcl10 and Malt1 control lysophosphatidic acid-induced NF- $\kappa$ B activation and cytokine production

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Lysophosphatidic acid (LPA) is a potent bioactive phospholipid that stimulates a variety of cellular responses by acting on cognate G protein-coupled receptors (GPCRs). There is increasing evidence that LPA signaling reprograms gene expression, but the GPCR-induced pathways connecting LPA receptor stimulation to downstream transcription factors are not well characterized. Here, we identify the adapter proteins Bcl10 and Malt1 as essential mediators of LPA-induced NF- $\kappa$ B activation. Both proteins were previously known to activate NF- $\kappa$ B in response to antigen receptor ligation on lymphocytes, but their functions in nonimmune cells are still largely undefined. By using murine embryonic fibroblasts from Bcl10- or Malt1-deficient mice as a genetic model, we report that Bcl10 and Malt1 are critically required for the degradation of I $\kappa$ B- $\alpha$  and the subsequent NF- $\kappa$ B induction in response to LPA stimulation. Bcl10 and Malt1 cooperate with PKCs selectively for LPA-induced NF- $\kappa$ B activation but are dispensable for the activation of the Jnk, p38, Erk MAP kinase, and Akt signaling pathways. In a biological readout, we demonstrate that LPA-induced IL-6 production is abolished in the absence of Bcl10. Thus, our results identify a NF- $\kappa$ B-inducing signaling pathway downstream of GPCRs and reveal previously unrecognized functions for Bcl10/Malt1 signaling in nonimmune cells.

G protein-coupled receptor | signal transduction

NF- $\kappa$ B is a ubiquitously expressed dimeric transcription factor regulating inducible gene expression for cell proliferation, survival, differentiation and inflammation (1). It can be activated by a large variety of stimuli and plays critical roles in normal and disease physiology. Particularly, immune-cell-mediated pathologies and malignancies are causally connected to constitutive NF- $\kappa$ B activity (2–4). The mammalian NF- $\kappa$ B family contains five members: NF- $\kappa$ B1 (p105 and p50), NF- $\kappa$ B2 (p100 and p52), c-Rel, RelB, and RelA (p65). NF- $\kappa$ B dimers are retained in an inactive form in the cytoplasm by interactions with inhibitory I $\kappa$ B proteins. Most physiological and pathological signals for NF- $\kappa$ B activation depend on I $\kappa$ B kinase (IKK)-controlled events (1). Once activated, IKK phosphorylates I $\kappa$ B proteins on conserved serine residues, leading to ubiquitin-mediated I $\kappa$ B degradation and liberation of NF- $\kappa$ B, which then enters the nucleus to regulate the transcription of effector target genes.

A major challenge in the NF- $\kappa$ B field is to understand how distinct upstream stimuli activate IKK in a signal-specific manner (5). Recently, the adapter and scaffold proteins Bcl10 and Malt1 were identified as specific regulators of T cell receptor- and B cell receptor-mediated NF- $\kappa$ B activation (6). Originally, both molecules were isolated from chromosomal translocation breakpoints in mucosa-associated lymphoid tissue (MALT) lymphoma (7). Bcl10 is a caspase recruitment domain (CARD)-containing protein, and Malt1 is a molecule with structural relation to the caspase family of proteases. Bcl10 and Malt1 physically and functionally cooperate to couple antigen receptor-induced PKC signaling (PKC- $\theta$  in T cells or PKC- $\beta$  in B cells) to IKK activation to mediate NF- $\kappa$ B induction (8–12). In addition, a recent report demonstrated essential func-

tions for Bcl10 and Malt1 in Fc $\epsilon$ RI signaling (13). In mast cells, Bcl10 and Malt1 also operate downstream of PKC to selectively control Fc $\epsilon$ RI-induced NF- $\kappa$ B activation for proinflammatory cytokine production. The fact that Bcl10 and Malt1 also are expressed in nonimmune tissues suggests further still uncharacterized functions for Bcl10 and Malt1.

The largest family of cell surface receptors, which are expressed in virtually all mammalian tissues, are the G protein-coupled receptors (GPCRs) (14). In response to growth factors, hormones, neurotransmitters, or other stimuli, these receptors signal through a collection of heterotrimeric G proteins to activate downstream pathways for a broad spectrum of biological responses (15). There is increasing evidence that GPCRs are actively regulating gene transcription, and several GPCRs have been identified, which are able to activate NF- $\kappa$ B through unknown pathways (16). A prototypic GPCR agonist that can activate NF- $\kappa$ B is lysophosphatidic acid (LPA) (17, 18). LPA is a naturally occurring, water-soluble glycerophospholipid possessing growth factor-like activities. LPA can directly or indirectly (through the production of cytokines and chemokines) induce proliferation, migration, and survival of many cell types, normal and malignant (18). LPA activates distinct members of the endothelial differentiation gene (Edg) subfamily of GPCRs, which couple to at least three distinct G proteins (G<sub>q</sub>, G<sub>i</sub>, and G<sub>12/13</sub>) to feed into multiple effector systems, including NF- $\kappa$ B (18, 19).

Because LPA-induced NF- $\kappa$ B activation involves PKCs (20), we tested the hypothesis that Bcl10 and Malt1 might be required for LPA signal transduction. By using murine embryonic fibroblasts (MEFs) as a model system, we show that Bcl10 and Malt1 are specifically required to mediate activation of NF- $\kappa$ B and subsequent cytokine production in response to LPA treatment. These results identify a novel NF- $\kappa$ B-inducing signaling pathway and indicate previously unrecognized functions for the Bcl10/Malt1 complex in nonimmune cells.

## Results

**Bcl10 Is Required for LPA-Induced NF- $\kappa$ B Signaling.** Because LPA can activate NF- $\kappa$ B in fibroblasts (17), we used MEFs as a model system to study a potential involvement of Bcl10 in LPA signaling (Fig. 1). We first stimulated MEFs from Bcl10-heterozygous (Bcl10<sup>+/-</sup>) or Bcl10-deficient (Bcl10<sup>-/-</sup>) mice with LPA and examined signal-induced degradation of I $\kappa$ B- $\alpha$  by Western blotting (Fig. 1A). Although we detected time-dependent degradation of I $\kappa$ B- $\alpha$  in Bcl10<sup>+/-</sup> MEFs, LPA stimulation did not induce

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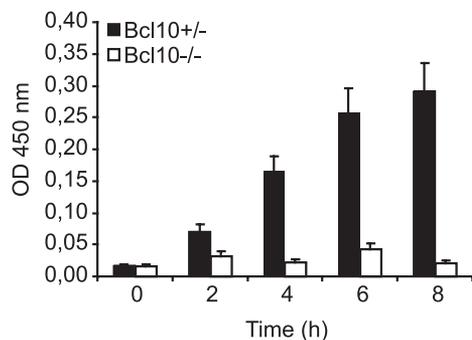
Abbreviations: CARD, caspase recruitment domain; GPCR, G protein-coupled receptor; IKK, I $\kappa$ B kinase; LPA, lysophosphatidic acid; MALT, mucosa-associated lymphoid tissue; MEF, murine embryonic fibroblasts.

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**Fig. 6.** Impaired IL-6 production in Bcl10-deficient MEFs. *Bcl10<sup>+/-</sup>* or *Bcl10<sup>-/-</sup>* MEFs were stimulated with 10  $\mu$ M LPA for the indicated times. Supernatants were collected, and IL-6 concentrations were determined by ELISA. Data are given as means  $\pm$  SEM and are representative of three independent experiments.

Consistent with these results, we detected regular TNF- $\alpha$ -induced NF- $\kappa$ B DNA binding activity in *Malt1<sup>-/-</sup>* MEFs but defective LPA-induced NF- $\kappa$ B induction (Fig. 5 C and D). As shown before in Bcl10-deficient MEFs, activation of Akt and the MAP kinases p38, Erk1/2, and Jnk also was not affected by Malt1 deficiency, indicating that both Bcl10 and Malt1 are dispensable for MAPK and Akt activation (Fig. 5E).

#### Bcl10 Signaling Controls LPA-Induced Production of the Cytokine IL-6.

We next tested whether Bcl10-dependent NF- $\kappa$ B activation might control physiological outcomes of LPA stimulation. NF- $\kappa$ B is known to be required for LPA-induced expression of the proinflammatory cytokine IL-6 (19). Therefore, we analyzed the production of IL-6 in *Bcl10<sup>+/-</sup>* and *Bcl10<sup>-/-</sup>* MEFs in response to LPA treatment by ELISA (Fig. 6). Whereas WT cells produced IL-6 in a time-dependent fashion, LPA-induced IL-6 production was blocked in the absence of Bcl10. Thus, the Bcl10-controlled LPA signaling pathway is essential for physiological effects of LPA stimulation.

#### Discussion

Here, we identify a GPCR signal transduction pathway and discover functions for the Bcl10/Malt1 complex in nonimmune cells. By using MEFs as a model, we demonstrate on a genetic basis that Bcl10 and Malt1 are both essential to transduce signals upon LPA stimulation selectively to the canonical NF- $\kappa$ B pathway, which proceeds via degradation of I $\kappa$ B- $\alpha$ . In a biological readout for this cascade, we show that Bcl10-deficient MEFs have severe defects in the LPA-induced production of the NF- $\kappa$ B target cytokine IL-6.

Several previous reports have shown that cell stimulation with LPA can activate NF- $\kappa$ B (17, 19, 20, 23–26). LPA-induced NF- $\kappa$ B signaling controls the production of proinflammatory cytokines and chemokines in tumor cells, and can indirectly and directly contribute to the survival of malignant cells, indicating important pathophysiological functions for this pathway (18, 19, 23, 25, 27). We now provide mechanistic insights into this pathway. In line with recent work (20), we find that a pharmacological inhibition of PKCs with the pan-PKC inhibitor rottlerin, but not the inhibition of PI3K/Akt, blocked NF- $\kappa$ B induction by LPA. Several PKC isoforms can be activated by LPA stimulation in various cell types, including PKC- $\alpha$ , PKC- $\beta$ , PKC- $\delta$ , PKC- $\epsilon$ , and PKC- $\zeta$  (20, 28). PKC- $\delta$  has been shown to control LPA-induced IL-8 production via NF- $\kappa$ B in bronchial epithelial cells (20). Yet the specific PKC isoform that regulates LPA signaling for NF- $\kappa$ B in fibroblast remains to be determined, for example, by using specific siRNAs. Because several PKCs are expressed in MEFs, it could be possible that there is redundancy

at the PKC level. Nevertheless, the finding that a pan-PKC inhibition blocks NF- $\kappa$ B to a similar extent as the genetic deletion of Bcl10 or Malt1 indicates a nonredundant role for the Bcl10/Malt1 complex in the PKC-dependent LPA signaling. This model is consistent with the known essential roles of Bcl10 and Malt1 downstream of PKCs in immune cells.

Distinct PKC isoforms control NF- $\kappa$ B activation in response to T cell receptor, B cell receptor, or Fc $\epsilon$ RI signaling in a Bcl10-dependent manner (11–13, 29). Upon antigen receptor ligation, PKC- $\beta$  (in B cells) or PKC- $\theta$  (in T cells) phosphorylate the Bcl10-binding molecule Carma1 (also called Card11/Bimp3). Carma1 is a member of a larger protein family additionally including Carma2 (Card14/Bimp2) and Carma3 (Card10/Bimp1). The Carma proteins possess a C-terminal CARD that can interact with the Bcl10 CARD, and they possess a coiled-coil domain and N-terminal MAGUK region (6). Carma1 is phosphorylated by PKCs in a region between the coiled coil and the MAGUK domain now referred to as the linker region (11, 12). The phosphorylation of Carma1 relieves inhibitory intramolecular interactions, allowing recruitment of Bcl10 and Malt1 to Carma1. These events initiate the assembly of higher-order signalosomes that contain additional proteins, including TRAF6, Tak1, Tak-associated proteins (Tab1, -2, or -3) and the ubiquitination enzymes Ubc13/Uev1A (30, 31). These proteins subsequently induce downstream IKK activation via processes that involve protein oligomerization and ubiquitylation events (30, 31). We consider it likely that the LPA-induced NF- $\kappa$ B pathway also involves the assembly of Bcl10/Malt1 signalosomes, which, similar to the signalosomes in activated lymphocytes, could require a Carma molecule and include tumor necrosis factor receptor-associated factors, Tak1, and Tab proteins. Carma1 is specifically expressed in immune cells, and Carma2 is selectively expressed in the placenta (32, 33). Based on the broad expression of Carma3 in multiple tissues and the responsiveness of multiple cell lineages to LPA stimulation, we speculate that the Carma protein for PKC-dependent LPA signaling could be Carma3. The idea that Carma3 could physiologically regulate PKC-induced NF- $\kappa$ B signaling would be consistent with the findings that the Carma3 linker region can be phosphorylated by PKCs and that Carma3 can functionally replace Carma1 for PKC-dependent cell activation (11). However, Bcl10/Malt1 function for LPA signaling and Bcl10/Malt1 function for antigen receptor signaling differ in several aspects. Previous reports have indicated that p38 and Jnk are regulated by Bcl10 and Malt1 in response to antigen receptor signals (6). In contrast, we report here that LPA signaling to p38 and Jnk does not require Bcl10 or Malt1. Moreover, a recent study has indicated a function for PI3K/Akt signaling in Carma1-mediated NF- $\kappa$ B induction in T cells (21), whereas we do not detect effects of a PI3K/Akt inhibition on LPA-induced signaling to NF- $\kappa$ B. Together, these results indicate that the utilization of upstream and Bcl10/Malt1 effector pathways is cell-lineage and/or stimulus-specific and reinforces the necessity to investigate the function of this signaling complex in a context-dependent manner. Future experimental work is necessary to define the precise components and the regulation of Bcl10/Malt1-mediated NF- $\kappa$ B signaling in response to LPA stimulation.

LPA levels are significantly increased in various malignant effusions (18, 19), and LPA receptors are aberrantly expressed in several human cancers (18). Stimulation of tumor cells with LPA induces cell proliferation (34), cell survival, and drug resistance (35, 36) and promotes cell motility and invasion (37, 38) through still-undefined signaling pathways. Aberrant NF- $\kappa$ B signaling can mediate all these cellular outcomes (2). We show in conclusion that Bcl10/Malt1 signaling controls LPA-induced NF- $\kappa$ B activation. Based on these findings and the known roles of LPA in cancer, we suggest that the activity and function of

