Autophagy Negatively Regulates Keratinocyte Inflammatory Responses via Scaffolding Protein p62/SQSTM1

Hye-Mi Lee, Dong-Min Shin, Jae-Min Yuk, Ge Shi, Dae-Kyoung Choi, Sang-Hee Lee, Song Mei Huang, Jin-Man Kim, Chang Deok Kim, Jeung-Hoon Lee and Eun-Kyeong Jo

*J Immunol* 2011; 186:1248-1258; Prepublished online 15 December 2010; doi: 10.4049/jimmunol.1001954

http://www.jimmunol.org/content/186/2/1248

Supplementary Material

http://www.jimmunol.org/content/suppl/2010/12/15/jimmunol.1001954.DC1

References

This article cites 52 articles, 18 of which you can access for free at:

http://www.jimmunol.org/content/186/2/1248.full#ref-list-1

Subscription

Information about subscribing to *The Journal of Immunology* is online at:

http://jimmunol.org/subscription

Permissions

Submit copyright permission requests at:

http://www.aai.org/About/Publications/JI/copyright.html

Email Alerts

Receive free email-alerts when new articles cite this article. Sign up at:

http://jimmunol.org/alerts
Autophagy Negatively Regulates Keratinocyte Inflammatory Responses via Scaffolding Protein p62/SQSTM1

Hye-Mi Lee,*,† Dong-Min Shin,*,‡ Jae-Min Yuk,*,‡ Ge Shi,‡ Dae-Kyoung Choi,‡ Sang-Hee Lee,‡,§ Song Mei Huang,‡,§ Jin-Man Kim,‡,§,‖ Chang Deok Kim,‡,§ Jeung-Hoon Lee,‡,§ and Eun-Kyeong Jo*,†

The scaffolding adaptor protein p62/SQSTM1 (p62) has been shown to be an autophagy receptor that acts as a link between the ubiquitination and autophagy machineries. However, the roles of autophagy and p62 in human keratinocytes are not well understood. In this study, we show that keratinocyte autophagy negatively regulates p62 expression, which is essential for the prevention of excessive inflammation and the induction of cathelicidin in human keratinocytes. Stimulation of TLR2/6 or TLR4 in primary human keratinocytes robustly activated autophagy pathways and up-regulated p62 expression through induction of NADPH oxidases 2 and 4 and the generation of reactive oxygen species. MyD88 and TNFR-associated factor 6, key signaling molecules that mediate TLR activation, played an essential role in the induction of autophagy and p62 expression. Additionally, blockade of autophagy significantly increased the generation of inflammatory cytokines and expression of p62 in primary human keratinocytes. Notably, silencing hp62 through RNA interference resulted in a significant decrease in NF-κB activation, inflammatory cytokine production, cathelicidin expression, and cell proliferation (as well as cyclin D1 expression) in keratinocytes. Epidermal expression of p62 was further found to be significantly higher in psoriatic skin than in skin affected by atopic dermatitis or from healthy controls. Collectively, our data provide new insights into the roles of autophagy and p62 in controlling cutaneous inflammation.

The online version of this article contains supplemental material.

Abbreviations used in this article: AD, atopic dermatitis; Baf-A1, bafilomycin A1; Cdk, cyclin-dependent kinase; DHE, dihydrorhodamine; DPI, diphenyleneiodonium; IKK, IκB kinase; 3-MA, 3-methyladenine; NAC, N-acetylcysteine; NOX, NADPH oxidase; ROS, reactive oxygen species; shAtg5, short hairpin RNA specific to hAtg5; shBeclin-1, short hairpin RNA specific to hBeclin-1; shNS, nonspecific control short hairpin RNA; siRNA, small interfering RNA; TRAF6, TNFR-associated factor.

Received for publication June 15, 2010. Accepted for publication November 7, 2010.

Address correspondence and reprint requests to Dr. Eun-Kyeong Jo, Department of Microbiology, College of Medicine, Chungnam National University, 6 Mahwa-dong, Jungku, Daejeon 301-747, South Korea. E-mail address: hayoungj@cnu.ac.kr

The online version of this article contains supplemental material.

Copyright © 2011 by The American Association of Immunologists, Inc.

www.jimmunol.org/cgi/doi/10.4049/jimmunol.1001954

Autophagy Negatively Regulates Keratinocyte Inflammatory Responses via Scaffolding Protein p62/SQSTM1

The scaffolding adaptor protein p62/SQSTM1 (p62) has been shown to be an autophagy receptor that acts as a link between the ubiquitination and autophagy machineries. However, the roles of autophagy and p62 in human keratinocytes are not well understood. In this study, we show that keratinocyte autophagy negatively regulates p62 expression, which is essential for the prevention of excessive inflammation and the induction of cathelicidin in human keratinocytes. Stimulation of TLR2/6 or TLR4 in primary human keratinocytes robustly activated autophagy pathways and up-regulated p62 expression through induction of NADPH oxidases 2 and 4 and the generation of reactive oxygen species. MyD88 and TNFR-associated factor 6, key signaling molecules that mediate TLR activation, played an essential role in the induction of autophagy and p62 expression. Additionally, blockade of autophagy significantly increased the generation of inflammatory cytokines and expression of p62 in primary human keratinocytes. Notably, silencing hp62 through RNA interference resulted in a significant decrease in NF-κB activation, inflammatory cytokine production, cathelicidin expression, and cell proliferation (as well as cyclin D1 expression) in keratinocytes. Epidermal expression of p62 was further found to be significantly higher in psoriatic skin than in skin affected by atopic dermatitis or from healthy controls. Collectively, our data provide new insights into the roles of autophagy and p62 in controlling cutaneous inflammation.
p62 accumulation may contribute to the pathogenesis of this chronic inflammatory skin disorder.

Materials and Methods

Cell cultures

Human primary keratinocytes and HaCaT cells were cultured and maintained, as described previously (17). Briefly, foreskin epidermal keratinocytes were cultured in keratinocyte serum-free medium supplemented with bovine pituitary extract and recombinant human epidermal growth factor (Life Technologies/Life Technologies, Eggenstein, Germany). HaCaT cells were cultured in DMEM (Life Technologies-BRL, Gaithersburg, MD), supplemented with 10% FCS (Invitrogen, San Diego, CA), 1% l-glutamine, penicillin (50 U/L; Life Technologies-BRL), and streptomycin (50 mg/mL; Life Technologies-BRL). The study protocol was approved by the Institutional Research Board of Chungnam National University College of Medicine. Written informed consent was obtained from each participant.

Reagents, DNA, and Abs

MALP-2, a diacylated lipoprotein, was purchased from Immunex (San Diego, CA). N-acetylcyesteine (NAC), diphenylene iodonium (DPI), allopurinol, and wortmannin were obtained from Calbiochem (San Diego, CA), and 3-methyldadenine (3-MA), bafilomycin A1 (Baf-A1), rapamycin, LPS (Escherichia coli 0111:B4), and DMSO were from Sigma-Aldrich (St. Louis, MO). piRNA-h7SKGFp2eo plasmids specific for hTLR2 and hTLR4 were purchased from InvivoGen. Cells were transfected using LipofectAMINE 2000, as suggested by the manufacturer (Invitrogen, Carlsbad, CA). Anti–phospho-Akt (Ser473) and anti–phospho-ERK1/2 (Thr202/Tyr204) were obtained from Cell Signaling Technology, an anti-LC3 Ab from Novus Biologicals (Littleton, CO), and Abs specific for p62/SQSTM1 (H-300), Atg5 (FL-275), MyD88 (N-19), TRAF6 (H-274), and cyclin D1 (H-295), cyclin-dependent kinase (Cdk) 4 (H-22), Beclin-1 (NM_004849), and reverse 3'-TTGCCATCTGAGCCAAAG-3' were obtained from Cell Signaling Technology, an anti-LC3 Ab from Novus Biologicals, an anti-LC3 Ab from Cell Signaling Technology, an anti-LC3 Ab from Novus Biologicals, an anti-LC3 Ab from Cell Signaling Technology, a primary Ab (rabbit, 1:400) overnight, and then incubated with secondary Ab (anti-rabbit IgG-tetramethylrhodamine isothiocyanate (TRITC)) (C-21) were purchased from Cell Signaling Technology (Beverly, MA).

For immunofluorescence analyses, an anti-LC3 Ab was purchased from MBL International (Woburn, MA). Anti–mouse IgG-Cy2, anti–rabbit IgG-Alexa488, and anti–rabbit IgG-tetramethylrhodamine isothiocyanate Abs were obtained from Jackson ImmunoResearch Laboratories (West Grove, PA). The NF-κB luciferase reporter plasmid was a gift from G.M. Hur (Chungnam National University, Daejeon, Korea). A NF-κB-luc adenoviral vector (AdHSV-NF-κB-luc) was purchased from Genetransfer Vector Core (Iowa City, IA).

Lentiviral packaging and short hairpin RNA transduction

Human pLKO.l lentiviral constructs from Open Biosystems, targeted to hbeclin-1 (NM_003766), hatg5 (NM_004849), and hpy62/SQSTM1 (NM_00142298), were supplied by Sigma-Aldrich as glycerol stocks. Virus production was performed, as described previously (17, 18). Briefly, lentiviruses were transduced into human keratinocyte 293T cells with pLKO puro, 1 nonspecific target short hairpin RNA (shRNA) plasmid, and the following packaging plasmids: pMDL-RRE (5 μg), PSV-REV (2.5 μg), and pVSV-g (3 μg). After 48 h, human epidermal keratinocyte 293T cell supernatants containing target viruses were collected, filtered, and concentrated by ultracentrifugation. Collected viruses were aliquoted and stored at −80°C. For viral titration, 2 × 10^6 HaCaT cells were seeded in 6-well plates, and then nontarget pLKO.1 nonspecific control shRNA (shNS) or target lentiviral particles were serially added with 8 μg/mL polybrene (Sigma-Aldrich). After 12 d, relative titers were determined by colony counting under puromycin (1 μg/mL) selection.

Transfection of HaCaT cells

Small interfering RNA (siRNA) transfection was performed, as described previously (17). Briefly, HaCaT cells were transfected with scrambled, nonspecific control siRNA (shNS), or siRNAs specific for hCOX2, hNOD4, hMDR88, hTRAF6, and hpy62/SQSTM1 (Santa Cruz Biotechnology) using the transfection reagent LipofectAMINE 2000 (Invitrogen), according to the manufacturer’s protocol.

Cell proliferation assays

Cell viability and the proliferation of keratinocytes were measured using the Cell Counting Kit-8 assay (Dojindo Molecular Technologies), performed according to the manufacturer’s instructions. Optical absorbances were measured, as described previously (17).

RNA extraction and RT-PCR analysis

The extraction of RNA from cells or paraffin-embedded skin tissue and semiquantitative RT-PCR were performed, as described previously (17). Briefly, total RNA was extracted using the TRIzol reagent (cells; Invitrogen) or High Pure RNA Paraffin Kit (tissue; Roche Applied Science, Mannheim, Germany). PCR reactions involved 35 cycles of annealing, extension, and denaturation. The annealing step consisted of 30 s at one of various annealing temperatures, as follows: 52°C for hpy62/SQSTM1 and β-actin; 54°C for Atyg5; 58°C for hTLR2 and hTLR4; 59°C for hLL-37; 63°C for Beclin-1 and IL-6; and 66°C for TNF-α. The following primers were used: hpy62/SQSTM1 forward 5′-CTTCCGGACACTAGCCTGTTG-3′ and reverse 5′-ACACTTAAAGGCGCCAGGAG-3′; hTLR2 forward 5′-TGCGTATCGTGACACAGGG-3′ and reverse 5′-ttcggacctcatctgggaaagt-3′; hATRG forward 5′-TGGGATTGCAAAATGAC-3′ and reverse 5′-CATGGATGGAGTAAGTGTTG-3′; and hTLR4 forward 5′-CTGAAATGTGAGGCACCCCAACA-3′ and reverse 5′-TCCCCCACCATTCAGAGGTTCC-3′. hIL-6 forward 5′-GGCCGAAAGATTTTCACAAAT-3′ and reverse 5′-AATGGTCTTTGATGTTGCC-3′ and reverse 5′-AATGGTCTTTGATGTTGCC-3′. hβ-actin forward 5′-TGGCAAT-GAATTTTTATGC-3′ and reverse 5′-TCCCCATCTGAGGAAGTTTTG-3′. hNF-κB-luc forward 5′-CTCCACCTCAAGGTTCCCTG-3′ and reverse 5′-AATGGTCTTTGATGTTGCC-3′. hGLU forward 5′-CTTGGGCACCATCATCTGACCC-3′ and reverse 5′-TCCCCCACCATTCAGAGGTTCC-3′. hIL-6 forward 5′-TCCCGACCACTCTTGGACGAAGA-3′ and reverse 5′-TCCCCCACCATTCAGAGGTTCC-3′.

Western blot, ELISA analysis, and NF-κB luciferase assays

Western analysis and cytokine ELISAs (for TNF-α and IL-6) were performed, as described previously (17). NF-κB luciferase assays were also performed, as described previously (17). Briefly, cells were transfected with a NF-κB-luc construct and siRNA (for HaCaT cells) or transduced with AdHSV-NF-κB-luc. Following stimulation with MALP-2 (100 ng/ml) for 24 h, cells were lysed and their luciferase activity was analyzed using the Luciferase Assay System (Promega, Madison, WI), according to the manufacturer’s instructions.

Immunofluorescence analysis

Immunofluorescence analysis for LC3 was performed, as described previously (18). Briefly, cells were fixed in 4% paraformaldehyde at 4°C for 10 min. They were then permeabilized with 0.01% Triton X-100, before being blocked with 10% BSA for 1 h. Cells were stained with primary Ab (rabbit anti-human LC3, diluted 1:400) overnight, and then incubated with secondary Ab (anti-rabbit IgG-tetramethylrhodamine isothiocyanate or anti-rabbit IgG-Alexa488) for 1 h. Unbound Ab was removed with PBS, and cells were imaged using a Zeiss LSM510 META confocal microscope (Zeiss).

Transmission electron microscopy

Transmission electron microscopy was performed, as described previously (18). Briefly, after MALP-2–stimulated human primary keratinocytes were fixed with 3% formaldehyde and 2% glutaraldehyde (Electron Microscopy Sciences, Road Hatfield, PA) in 0.1 M sodium cacodylate buffer (pH 7.4; Ted Pella, Redding, CA) for 1 h, they were then washed with 0.1% cacodylate buffer and postfixed with 1% osmium tetroxide (Electron Microscopy Sciences) and 0.5% potassium ferricyanide in cacodylate buffer for 1 h. Cells were next dehydrated with increasing concentrations of ethanol, and infiltrated with Epon-Araldite resin at 80°C for 24 h. Ultrathin sections (70–80 nm) were obtained using an ultramicrotome (RMC MT6000-XL). Sections were counterstained with uranyl acetate and lead citrate and viewed using a Zeiss EM 900 transmission electron microscope.

Immunohistochemistry

The subcellular localization of human p62 in tissue sections was determined through immunohistochemical staining, performed as described previously (17). Biopsies were obtained from patients with childhood-onset dermatitis (AD), as well as five healthy controls, was fixed in formalin, embedded in paraffin, and sectioned to a thickness of 3 μm. After overnight incubation with the primary Ab (rabbit polyclonal p62 Ab, diluted 1:400; Abgent) at 4°C, tissue sections were incubated with the substrate 3,3’-diaminobenzidine (Dako) to generate a colored signal. They were then lightly counterstained with Mayer’s hematoxylin and mounted using Immu-Mount (Thermo Shandon, Pittsburgh, PA). Negative control sections were treated with serum from nonimmunized rabbits (Dako) in place of primary Ab. Cells displaying cytoplasmatic staining were considered to be positive. A single pathologist examined all the immunostained tissue sections.

Downloaded from http://www.jimmunol.org/ by guest on April 20, 2017
**Statistical analysis**

All data are presented as the mean ± SD of independent experiments. Statistical analysis was performed using a paired Student t test with Bonferroni correction or, for multiple comparisons, ANOVA. The p values <0.05 were deemed to be statistically significant.

**Results**

**TLR2/6 stimulation activates autophagy in primary human keratinocytes**

It has previously been shown that TLR ligands stimulate the formation of autophagosomes and autophagolysosomes (7, 9, 10). However, it is not known whether TLR agonists activate autophagy in primary human keratinocytes. To determine whether TLR2/6 stimulation activates autophagy in human keratinocytes, we stimulated primary human keratinocytes with MALP-2 (100 ng/ml) and measured the formation of organelles displaying punctate LC3 staining (autophagosomes) using endogenous LC3 as an autophagic marker. As shown in Fig. 1A, MALP-2 induced autophagy in primary human keratinocytes, with peak activation occurring at 36 h. In addition, posttranslational modification of LC3-I, yielding LC3-II (19), was assessed in MALP-2–stimulated human keratinocytes. Significant increases in the LC3-II:LC3-I ratio were observed in primary human keratinocytes (Fig. 1B) and HaCaT cells (Supplemental Fig. 1A) after MALP-2 stimulation. LC3 autophagosome formation was significantly inhibited in cells pretreated with 3-MA or wortmannin, specific inhibitors of autophagy (20, 21) (Fig. 1C). Additionally, MALP-2–induced punctate LC3 staining was significantly reduced in cells in which Beclin-1 or Atg5 was silenced through transfection with a specific lentiviral shRNA (Fig. 1D).

Colocalization of LC3 and lysosomes (the latter stained with LysoTracker) was significantly increased in primary human keratinocytes following MALP-2 stimulation (Fig. 1E). In primary keratinocytes, the ability of MALP-2 to increase the LC3-II:LC3-I ratio (measured by Western blotting) was attenuated by pretreatment with 3-MA (Fig. 1F). Furthermore, pretreatment with the vacuolar H+–ATPase inhibitor Baf-A1, which prevents lysosomal...
acidification and the fusion of autophagosomes with lysosomes (19, 22), significantly increased the LC3-II:LC3-I ratio in MALP-2–stimulated cells (Fig. 1F). Using a transmission electron microscope, we analyzed numbers of autophagosome-like vacuoles with double-membrane structures (data not shown). As shown in Supplemental Fig. 1B, morphologic analysis of keratinocytes by transmission electron microscopy revealed the cytoplasmic accumulation of autophagic vacuoles. Thus, MALP-2 actively induces both autophagy and autophagosome–lysosome fusion in primary human keratinocytes.

**TLR2/6 stimulation induces p62 expression in human keratinocytes**

It was previously shown that IFN-γ and TLR signaling cooperate to stimulate late expression of p62 in murine macrophages (23). Additionally, TLR ligands induce the formation of autophagosomes and autophagolysosomes (7, 9, 10). However, it is not known whether TLR ligands induce p62 expression and/or activate autophagy in human keratinocytes. To address this, we stimulated human keratinocytes with MALP-2 and measured p62 mRNA and protein expression by RT-PCR and Western blotting, respectively. Endogenous p62 mRNA and protein expression was low in unstimulated cells, but increased time dependently after MALP-2 stimulation (Fig. 2A, 2B; p62 mRNA and protein, respectively). We further assessed the role of TLR2 in MALP-2–induced p62 mRNA expression. Transfection of HaCaT cells with a siRNA specific for hTLR2, but not one targeted to hTLR4, significantly inhibited MALP-2–induced p62 expression (Fig. 2C). Thus, MALP-2 actively induces p62 expression in primary human keratinocytes via TLR2.

**NOX-dependent generation of ROS is required for the induction of p62 expression and activation of autophagy in TLR2/6–stimulated primary human keratinocytes**

ROS derived from superoxide play multiple roles in innate immune responses, including in microbial killing, the regulation of apoptosis, cytokine production, and gene expression, and the activation of antibacterial autophagy (24, 25). We first examined whether MALP-2 increased the intracellular generation of superoxide in human keratinocytes using the oxidative fluorescent dye dihydroethidium (DHE). As shown in Fig. 3A, MALP-2–induced superoxide production was detected within 30 min, peaking 1 h after stimulation. We next examined whether MALP-2–dependent ROS generation was required for p62 expression and the activation of autophagy. Treatment of keratinocytes with either the general ROS scavenger NAC or the NOX inhibitor DPI, but not the xanthine oxidase inhibitor allopurinol, significantly and dose dependently attenuated MALP-2–induced p62 expression (Fig. 3B). Previously, it has been shown that HaCaT cells express significant amounts of the gp91phox homologs NOX2 and NOX4, but only very low levels of NOX1 mRNA (26). Thus, we further investigated the roles of NOX2 and NOX4 in the MALP-2–mediated induction of p62. Silencing NOX2 or NOX4 using specific siRNAs significantly attenuated MALP-2–induced p62 expression in HaCaT cells (Fig. 3C).

The MALP-2–induced formation of autophagosomes displaying punctate LC3 expression (Fig. 3D) and increase in the LC3-II:LC3-I ratio (Fig. 3E) was dose dependently abrogated by pretreatment with NAC or DPI, but not allopurinol. These data suggest that TLR2/6–dependent ROS generation also plays a role in the activation of autophagy in keratinocytes. Transfection of HaCaT cells with siRNAs specific to hNOX2 and hNOX4 significantly inhibited autophagosome formation in response to MALP-2 (Fig. 3F). These data suggest that the NOX–dependent generation of ROS is critically involved in MALP-2–mediated p62 expression and autophagy in keratinocytes.

**TLR4 stimulation also leads to ROS-dependent activation of autophagy and p62 expression in keratinocytes**

Having established the role of TLR2/6 in the induction of autophagy, p62 expression, and ROS generation in human keratinocytes, we examined whether other TLR ligands function similarly in keratinocytes. Although previous studies showed that the TLR4 agonist LPS/endotoxin induced autophagy in macrophages (7, 9), a similar response has not been demonstrated in keratinocytes. As TLR4 was shown to be constitutively and inducibly expressed in HaCaT cells (2), we first examined punctate LC3 expression and LC3-II:LC3-I levels in HaCaT cells after LPS stimulation. Treatment of HaCaT cells with LPS robustly increased punctate LC3 expression and LC3-II band intensities at 18 h, indicating activation of autophagy (Fig. 4A, 4B). The marked increase in punctate LC3 staining was largely abrogated by pretreatment with 3-MA (Fig. 4A), a novel inhibitor of autophagy activation (20, 21). TLR4/LPS stimulation significantly increased the endogenous expression of p62 and LL-37 in HaCaT cells (Fig. 4C, p62 and LL-37 mRNA; Fig. 4D, p62 protein). We further showed that LPS stimulation robustly activated ROS generation in HaCaT cells in a time-dependent manner, and that the peak was observed within 30 min of stimulation (Fig. 4E). Blocking ROS production using the antioxidants NAC and DPI dose-dependently inhibited LC3-II expression (Fig. 4F). Additionally, transfection of HaCaT cells

**FIGURE 2.** TLR2/6 stimulation induces the expression of p62/SQSTM1. A and B, Primary human keratinocytes were stimulated with MALP-2 (2 μg/ml) for the indicated periods of time, and p62/SQSTM1 expression was then analyzed. A, Total RNA was isolated from primary human keratinocytes and subjected to semiquantitative RT-PCR analysis to determine p62 mRNA levels. B, Western analysis for p62 expression. C, Semiquantitative RT-PCR analysis. HaCaT cells were transfected with siNS or siRNAs specific for hTLR2 (siTLR2) or hTLR4 (siTLR4) and then treated with MALP-2 for 6 h. Images are representative of three independent experiments, all yielding similar results (top). Data represent the mean ± SD of three independent experiments (bottom). ***p < 0.001 (versus control).
with siRNAs specific for hNOX2 and hNOX4 significantly inhibited LPS-induced p62 expression (Fig. 4G). These data collectively show that the activation of TLR4, as well as TLR2/6, significantly upregulated NOX-dependent ROS generation, which contributes to autophagy activation and the expression of p62 and LL-37 in human keratinocytes.

**MylD88 and TRAF6 are required for p62 expression and the activation of autophagy in keratinocytes**

It was previously shown that the adaptor protein p62 selectively interacts with the TRAF domain of TRAF6 and plays an important intermediary role in the activation of NF-κB by TNF-α or IL-1 (27). Additionally, MyD88 plays a role in TLR4- or TLR7-induced activation of autophagy (7, 10). We therefore examined whether MALP-2–dependent activation of autophagy was mediated by MyD88 and/or TRAF6, both of which contribute to TLR signaling. When HaCaT cells were transfected with siRNAs specific for hMyD88 and hTRAF6, TRAF6/2-dependent p62 protein (Fig. 5A) and mRNA expression (Fig. 5B) were significantly reduced. Furthermore, the MALP-2–induced increases in LC3-positive autophagosome formation (Fig. 5C) and the LC3-II:LC3-I ratio (Fig. 5D) were significantly reduced in HaCaT cells transfected with siRNAs specific for hMyD88 or hTRAF6. These data suggest that both MyD88 and TRAF6, key signaling intermediates in TLR-dependent signaling, are involved in the activation of autophagy in human keratinocytes.

The autophagy pathway negatively regulates TLR2/6- and/or TLR4-induced inflammatory responses and p62 expression in keratinocytes

Accumulating evidence suggests that autophagy contributes to the regulation of cellular inflammation (28). Recently, it has been reported that autophagy-linked genes are involved in susceptibility to Crohn’s disease (29), and that autophagy defects exacerbate endotoxin-induced activation of the inflammasome (30). We next examined whether the activation of autophagy plays a role in the regulation of inflammatory responses and NF-κB activation in primary keratinocytes. To test this, primary human keratinocytes were pretreated with 3-MA or wortmannin, two inhibitors of the kinase PI3K that is required for the activation of autophagy (31), and transfected with shBeclin-1 or shAtg5 prior to MALP-2 stimulation. Notably, the secretion of TNF-α and IL-6 was dose dependently increased in primary human keratinocytes pretreated with 3-MA or wortmannin (Fig. 6A). Primary human keratinocytes transfected with shBeclin-1 or shAtg5 displayed significant increases in MALP-2–induced production of the proinflammatory cytokines TNF-α and IL-6 (Fig. 6B). Additionally, MALP-2–dependent NF-κB reporter gene activity was significantly increased in HaCaT cells pretreated with 3-MA or wortmannin (Supplemental Fig. 2A) and primary human keratinocytes transfected with shBeclin-1 or shAtg5 (Fig. 6C).

Next, the role of autophagy in regulating p62 expression was determined in human keratinocytes. Blockade of the autophagy
pathway using 3-MA or wortmannin significantly increased MALP-2–induced p62 protein expression in primary human keratinocytes (Fig. 6D). Transfection of primary human keratinocytes with shBeclin-1 or shAtg5 significantly increased the induction of p62 protein expression following MALP-2 stimulation (Fig. 6E). We further characterized the role of autophagy in TLR4-stimulated keratinocytes. Consistent with the results of TLR2/6 stimulation, pretreatment of HaCaT cells with 3-MA or wortmannin (Allo; 0.1 or 1 mM), HaCaT cells were treated with LPS for 18 h and analyzed for LC3 protein expression. Following preincubation for 45 min with medium containing NAC (30 or 50 mM), DPI (20 or 50 μM), or allopurinol (Allo; 0.1 or 1 mM), HaCaT cells were treated with LPS for 18 h and analyzed for LC3 protein expression. G, Western blot analysis of p62 levels in HaCaT cells transfected with siRNAs specific for hnox2 (siNOX2) and hnox4 (siNOX4), or siNS, and then treated with LPS (100 ng/ml). Images are representative of three independent experiments, all yielding similar results (B–D, F, G). L, LPS; U, untreated.

**FIGURE 4.** TLR4-induced expression of p62 and activation of autophagy in human keratinocytes are dependent on ROS. A, Immunofluorescence analysis of LC3 punctate expression. HaCaT cells were stimulated with LPS (100 ng/ml) in the absence or presence of 3-MA (2 h, 10 μM) for 18 h. Cells were stained with DAPI and anti-LC3–Alexa 488 Ab (experimental conditions as described in Fig. 1C). Data represent the mean ± SD of three independent replicates, each comprising at least 250 cells scored in five random fields. B–D, HaCaT cells were stimulated with LPS (100 ng/ml) for the indicated periods of time and analyzed for LC3 and p62 protein expression (B and D, Western blot analysis) and p62 and LL-37 mRNA expression (C, semi-quantitative RT-PCR). E, Measurement of intracellular superoxide generation through fluorescence analysis. HaCaT cells were stimulated with LPS for the indicated periods of time. Cells were then incubated with DHE (10 μM) for 15 min and analyzed for superoxide production by confocal microscopy. Data represent the mean ± SD of three independent experiments. F, Western blot analysis of LC3 expression. Following preincubation for 45 min with medium containing NAC (30 or 50 mM), DPI (20 or 50 μM), or allopurinol (Allo; 0.1 or 1 mM), HaCaT cells were treated with LPS for 18 h and analyzed for LC3 protein expression. G, Western blot analysis of p62 levels in HaCaT cells transfected with siRNAs specific for hnox2 (siNOX2) and hnox4 (siNOX4), or siNS, and then treated with LPS (100 ng/ml). Images are representative of three independent experiments, all yielding similar results (B–D, F, G). L, LPS; U, untreated.

TLR2/6- and/or TLR4-dependent p62 expression is required for the activation of inflammatory signaling and stimulation of keratinocyte proliferation

Recent studies have defined the roles of p62 in NF-κB activation and oncogenesis in autophagy-defective tumor cells (16), suggesting its involvement in inflammatory responses and tumorigenesis. To further explore the role of p62 in inflammatory responses in human keratinocytes, p62 was knocked down in HaCaT keratinocytes through transfection with a specific siRNA. Silencing p62 in this way led to significant decreases in the expression of TNF-α and IL-6 mRNA and protein in response to MALP-2 stimulation (Fig. 7A, 7B). The induction of LL-37 mRNA expression in HaCaT cells by TLR2/6 was suppressed by transfection with siRNA specific to hp62 (si-p62; Fig. 7A).

Following TLR2/6 stimulation, the activation of MyD88-dependent TRAF6 signaling leads to the activation of TGF-activated kinase 1, which phosphorylates the IKK complex (32). The IKK complex, in turn, phosphorylates IκB proteins, resulting in their ubiquitination and degradation (32). This key event triggers the translocation of NF-κB to the nucleus, where it mediates the transcription of target genes (32). To further investigate the underlying mechanisms by which p62 modulates TLR2/6-dependent inflammatory responses, we examined whether p62 played a role in IκKα/β phosphorylation and IκBα degradation in response to MALP-2. As shown in Fig. 7C, MALP-2 stimulation led to the time-dependent phosphorylation of IκKα/β and concomitant degradation of IκBα in human primary keratinocytes transfected with shNS. The transduction of primary keratinocytes with shRNA specific for hp62/SQSTM1 prior to MALP-2 stimulation markedly inhibited the phosphorylation of IκKα/β and degradation of IκBα in primary keratinocytes (Fig. 7C). Moreover, the silencing of p62 in HaCaT cells by p62 siRNA significantly inhibited NF-κB–luciferase reporter gene activity in response to MALP-2 treatment (Fig. 7D). We next examined whether the role of p62 in mediating inflammatory responses was confined to TLR2/6 responses. Silencing of p62 led to the significant inhibition of LPS-induced expression of TNF-α, IL-6, and LL-37, and NF-κB reporter gene activity, in HaCaT cells (Supplemental Fig. 3A–C). Thus, p62 appears to play...
critical roles in TLR2/6- and/or TLR4-induced inflammatory responses through activation of NF-κB signaling.

We further investigated whether p62 plays roles in MALP-2–dependent cell proliferation, cyclin D1/Cdk4 expression, and the activation of Akt and ERK1/2 in human keratinocytes. As shown in Fig. 7E, MALP-2 stimulation resulted in a substantial increase in keratinocyte proliferation, as measured in a Cell Counting Kit-8 assay. Silencing p62 expression led to the time-dependent inhibition of keratinocyte proliferation (Fig. 7E). Additionally, transduction of human primary keratinocytes with shRNA specific for p62 efficiently reduced the expression of cyclin D1/Cdk4 and phosphorylation of Akt (Fig. 7F), suggesting that p62 plays key roles in keratinocyte proliferation and cell cycle progression. When primary keratinocytes were transduced with shRNA specific for p62, but not shNS, the phosphorylation of ERK1/2 was significantly reduced (Fig. 7G). These data collectively show that p62 plays important regulatory roles in keratinocyte inflammatory responses through the activation of IKK complex–NF-κB signaling, and in keratinocyte proliferation via cyclin D1-/Cdk4-, Akt-, and ERK1/2-dependent signaling pathways.

Epidermal expression of p62 and LL-37 is up-regulated in patients with psoriasis, but not AD

Based on the roles of p62 in the regulation of inflammatory responses and proliferation, we next extended our analyses to determine whether p62 levels are elevated in inflammatory skin diseases.

**FIGURE 5.** TLR2/6-induced activation of autophagy and upregulation of p62 expression are mediated by MyD88 and TRAF6. HaCaT cells were transfected with siNS or siRNAs specific for hMyD88 (siMyD88) or hTRAF6 (siTRAF6). Following transfection, cells were stimulated with MALP-2 (10 or 100 ng/ml) for 24 h (A, p62/SQSTM1 protein expression; C, LC3 staining; D, Western blot analysis of LC3-II levels) or the indicated periods of time (B, p62 mRNA expression). B, Total RNA was isolated and subjected to quantitative real-time RT-PCR to determine p62 mRNA levels. Data are representative of three independent experiments, all yielding similar results (A, C [top], and D). Data represent the mean ± SD of three independent replicates (B, C [bottom]), each comprising at least 250 cells scored in five random fields (C). Scale bar, 10 μm. ***p < 0.001 (versus control). M, MALP-2; U, untreated.

**FIGURE 6.** Autophagy negatively regulates TLR2/6-mediated cytokine production, NF-κB activation, and p62 expression in keratinocytes. A and D, HaCaT cells were stimulated with MALP-2 (100 ng/ml) in the absence or presence of 3-MA (3MA; 1, 5, or 10 μM) or wortmannin (WM; 100, 200, or 300 nM). A, Supernatants were harvested 18 h after the commencement of MALP-2 stimulation, and levels of TNF-α and IL-6 were measured by ELISA. B, C, and E. Primary human keratinocytes were transduced with shNS, shBeclin-1, or Atg5 shAtg5. The cells were then stimulated with MALP-2 (100 ng/ml) for 18 h (B, ELISA; E, Western blot analysis of p62 protein expression) or 6 h (C, NF-κB-luc reporter gene analysis). Data represent the mean ± SD of three independent experiments (A–C [bottom], and D [bottom]). Images are representative of three independent experiments, all yielding similar results (D [bottom] and E). Transfection efficiency was assessed by RT-PCR (C, inset). ***p < 0.001 (versus control). M, MALP-2; Bec1, Beclin-1; U, untreated.
orders. To test this hypothesis, we performed comparative analyses of epidermal p62 expression in plaque psoriasis and eczema. Psoriasis is characterized by increased inflammation and the release of antimicrobial peptides, as well as hyperproliferation caused by disturbed differentiation of keratinocytes (3, 33). Eczemas represent a group of etiological and regional inflammatory skin disease variants characterized by pruritic, papulovesicular, and weeping dermatitis (34). As eczemas are a diverse group of disorders, we focused on the p62 expression profile in AD, an inflammatory skin disease that is typically familial, and one that is associated with IgE reactivity, Th2 predominance, and mast cell hyperactivity (35, 36).

To examine the role of p62 and cathelicidin LL-37 in inflammatory skin conditions, we measured the expression of p62 and LL-37 in the epidermis of biopsy samples obtained from psoriasis and AD patients and healthy subjects. Immunohistochemical staining revealed high numbers of p62-expressing keratinocytes in the skin of all psoriatic patients studied (35 of 35), but in no AD patients (0 of 22) and only a minority of healthy controls (5 of 16). The p62 expression was primarily detected in the cytoplasm in samples of psoriatic skin (Fig. 8A). As shown in Fig. 8B, RT-PCR analysis of mRNA extracted from patient skin samples revealed significantly higher expression of p62 and LL-37 in psoriasis patients than in AD patients or healthy controls. These results indicate that lesional psoriatic skin displays clear autophagic features, including punctate LC3 staining and colocalization of LC3 and lysosome markers.

Discussion

In human keratinocytes, TLR signaling robustly activates autophagy and p62 expression. In cells in which autophagic function was lost, inflammatory responses were profoundly exacerbated. Expression of p62 increased in parallel with these changes. In MALP-2–treated cells, knockdown of p62 not only reduced proinflammatory cytokine production, but also inhibited proliferation. In psoriatic skin, but not in skin from AD patients, p62 expression was significantly increased. These findings suggest the involvement of autophagy in the regulation of excessive cutaneous inflammation and implicate the upregulation of p62 in the pathogenesis of psoriasis.

Accumulating evidence suggests that the recognition of multiple pathogen-derived molecules can activate autophagy, principally in mononuclear phagocytes (7, 9, 10). TLR7 and TLR4 ligands have been reported to be potent inducers of autophagy in macrophages (7, 9). Additionally, stimulation of TLR2 through zymosan treatment can induce autophagy, specifically the formation of phagosomes and their fusion with lysosomes (37). However, it has not been detailed previously whether TLRs activate autophagy in human keratinocytes. Our data provide evidence that TLR signaling robustly activates autophagy in human keratinocytes. Electron microscopic imaging showed the rapid formation of autophagosomes and autolysosomes in human keratinocytes following TLR2/6 stimulation. Moreover, cells stimulated with MALP-2 displayed clear autophagic features, including punctate LC3 staining and colocalization of LC3 and lysosome markers. In addition,
MALP-2 stimulation significantly increased levels of lipid-bound LC3-II. During autophagy, the autophagic marker LC3 is cleaved and covalently conjugated to phosphatidyl-ethanolamine, yielding LC3-II, which is specifically targeted to elongated, Atg12-/Atg5-associated autophagosome precursors, thereafter remaining associated with autophagosomes (38, 39). Moreover, our data show that TLR4/LPS stimulation further induces autophagy in keratinocytes (see Fig. 4). Thus, our data suggest that the stimulation of innate immunity-associated TLRs in primary human keratinocytes actively induces autophagosome formation and regulates the maturation of autophagolysosomes.

We found p62 expression and the activation of autophagy to be dependent on NOX-mediated ROS generation and MyD88- and TRAF6-mediated signaling. Indeed, p62 mediates the TNF-α-and IL-1–dependent activation of NF-κB by interacting with the specific adapters receptor-interacting protein and TRAF6 (27, 40). Recent studies demonstrated that ROS generation contributes to the activation of autophagy, TLR activation, and phagocytosis in PMA-stimulated neutrophils (41). Similar to a previous study (25), we found that NOX activity and ROS generation played important roles in the activation of autophagy in primary human keratinocytes. Moreover, MyD88 has been previously implicated in TLR4- or TLR7-induced autophagy (7, 9, 10). Both key adaptor molecules in TLR signaling, MyD88 and TIR-domain–containing adapter-inducing IFN-β, modulate the interaction between Bcl-2 and Beclin-1, thereby actively triggering and regulating autophagy (10). Together, these data implicate innate immune signaling and key adaptor molecules are involved in the activation of autophagy pathways, not only in leukocytes, but also in nonleukocytic cells.

In this study, we have demonstrated that the inhibition of autophagy results in significantly increased inflammatory cytokine production and p62 expression in primary human keratinocytes. Enhanced inflammatory responses and increased cell proliferation were observed in keratinocytes treated with compounds that interfere with early (3-MA) and late (Baf-A1) autophagic processes or with siRNA targeted to genes essential for autophagy (hBeclin-1, hAtg5). Recently, it was shown that a defect in autophagy potentiated RIG-like receptor signaling and increased IFN secretion caused by ROS generated by dysfunctional mitochondria (42). Evidence suggests that autophagy acts to regulate cellular homeostasis and signaling by altering the availability of key adaptor proteins, such as p62 (43) and Nbr1 (12). Notably, p62 is itself an important intracellular target for autophagy: its cytosolic accumulation represses inclusion body formation (43). Previous studies showed that the inhibition of autophagy leads to the accumulation of p62 in transformed cells, thus compromising the clearance of ubiquitinated proteasomal substrates (44). Together, these data suggest that the pathways controlling the activation of autophagy tightly control p62 levels to prevent homeostasis from being disrupted during inflammatory responses.

Our data further show that knockdown of p62 using a specific siRNA reduced the expression of inflammatory cytokine genes and cathelicidin, cell proliferation (as assessed by measuring cyclin D expression), and NF-κB promoter activity in keratinocytes. Earlier studies showed that dominant-negative mutants and the downregulation of p62 using an antisense plasmid markedly inhibited the activation of NF-κB by TNF-α, suggesting a critical role for p62 (40). It has been reported that cytosolic p62/SQSTM1 expression is induced by Ras, is required for IskB kinase activation through polyubiquitination of TRAF6, and contributes to the development of Ras-induced lung adenocarcinomas in murine models (45). Additionally, sustained p62 expression accompanying defective

**FIGURE 8.** Expression of p62 and LL-37 is increased in the epidermis of psoriatic lesions. A, Immunohistochemical analysis of p62 expression in epidermal cells of normal skin (left), lesional psoriatic skin (middle), and skin affected by AD (right). p62 (brown) and nuclei (blue) are stained in epidermal cells. Scale bar, 400 μm. B, Semiquantitative (left, seven samples from each group are shown) and quantitative RT-PCR analyses (right) of p62 and LL-37 mRNA expression in the epidermis of normal (n = 16), psoriatic (n = 31), and AD-affected (n = 22) skin. ***p < 0.001 (versus normal skin).
autophagy results in altered NF-κB regulation and gene expression, thus favoring tumorigenesis (16). As an atypical protein kinase C scaffold protein, p62 plays an important role in the activation of NF-κB in response to IL-1, TNF-α, and receptor activator for NF-κB ligand through the assembly of an aPKC/p62/TRAF6 multi-protein complex (46). Notably, our data demonstrate that p62 expression is crucial for the TLR-dependent activation of IKKα/β and degradation of IκBα in human keratinocytes (see Fig. 7). Thus, the cytosolic accumulation of p62/SQSTM1 may trigger TLR-induced inflammatory responses in keratinocytes through modulation of IKK–NF-κB signaling.

Both psoriasis and AD are common chronic and relapsing inflammatory skin diseases (33–35). Psoriasis is an inflammatory skin disease characterized by increased levels of TRLS 1, 2, 4, 5, and 9 (reviewed in Ref. 3). It has been suggested that upregulated TLR expression in psoriatic skin contributes to disease pathophysiology by increasing the local production of antimicrobial peptides, and stimulating cutaneous immune responses and epidermal hyperproliferation (3, 33). In contrast, monocytes from AD patients displayed significantly impaired TLR2-mediated production of the proinflammatory cytokines IL-1β and TNF-α (47). Additionally, the immunological factors involved in the pathogenesis of AD are associated with upregulated IgE reactivity, and increased numbers of mast cells, eosinophils, and CD4-activated Th2 cells (34–36). Importantly, we found that the expression of p62 and cathelicidin was higher in psoriatic skin lesions than in lesions from AD patients or skin from healthy controls (see Fig. 8). The current data partly corroborate the findings of previous studies, which showed that psoriatic skin contained high amounts of the antimicrobial peptide cathelicidin/LL-37 (48). LL-37 potently activates phagocytic yeast cells to produce type 1 IFNs by binding nonstimulatory self-DNA and causing it to adopt an aggregated and condensed structure (49). Several previous studies further showed that levels of various antimicrobial peptides, including LL-37 and human β-defensin 2, are significantly decreased in acute and chronic lesions from AD patients (50). The depressed expression of innate immune effectors and antimicrobial peptides may contribute to the increased susceptibility of AD patients to bacterial and viral infections (3, 51, 52). Together, p62 and cathelicidin may play important roles in the homeostatic regulation of cutaneous inflammatory responses, especially in psoriasis. Future studies should clarify the detailed molecular mechanism(s) by which enhanced p62 expression exacerbates psoriatic inflammatory responses by breaking innate tolerance and driving autoimmunity.

The current findings demonstrate a major role for autophagy in the regulation of keratinocyte inflammation and provide new insights into the mechanisms by which TLR signaling activates autophagy in human keratinocytes. TLR innate signaling activation in keratinocytes triggers a simultaneous induction of innate inflammatory responses and autophagy pathways. Importantly, the autophagy pathways were found to play a key role in fine tuning of inflammatory responses in keratinocytes. The ability of the inhibition of keratinocyte autophagy to increase inflammation may have important implications for the development of novel therapeutic strategies for improving inflammatory homeostasis in psoriasis patients.

Acknowledgments

We thank K.S. Song, G.M. Hur, and K.K. Kim for discussion and reagents and J.S. Kim for critical reading of the manuscript.

Disclosures

The authors have no financial conflicts of interest.

References


12. Kirkin, V., D. G. McEwan, I. Novak, and I. Dikic. 2009. A role for ubiquitin in the regulation of keratinocyte inflammation and provide new insights into the mechanisms by which TLR signaling activates autophagy in human keratinocytes. TLR innate signaling activation in keratinocytes triggers a simultaneous induction of innate inflammatory responses and autophagy pathways. Importantly, the autophagy pathways were found to play a key role in fine tuning of inflammatory responses in keratinocytes. The ability of the inhibition of keratinocyte autophagy to increase inflammation may have important implications for the development of novel therapeutic strategies for improving inflammatory homeostasis in psoriasis patients.

Acknowledgments

We thank K.S. Song, G.M. Hur, and K.K. Kim for discussion and reagents and J.S. Kim for critical reading of the manuscript.

Disclosures

The authors have no financial conflicts of interest.

Downloaded from http://www.jimmunol.org/ by guest on April 20, 2017


SUPPLEMENTARY FIGURE LEGENDS

FIGURE S1. TLR2/6 stimulation induces autophagy in human keratinocytes. A, Western blot analysis of LC3-I/-II levels in HaCaT cells stimulated with MALP-2 (100 ng/mL). Data are representative of three independent experiments, all yielding similar results. B, Primary human keratinocytes were stimulated with MALP-2 (100 ng/mL) for 24 h and visualized by electron microscopy. Electron micrographs taken under low (left upper) and high (other panels) magnification showing the accumulation of autophagic vesicles (arrow).

FIGURE S2. Autophagy negatively regulates TLR4-mediated cytokine production, NF-κB activation, and p62 expression in keratinocytes. A-C, HaCaT cells were stimulated with MALP-2 (100 ng/mL) or LPS (100 ng/mL) in the absence or presence of 3-MA (1, 5, 10 mM) or wortmannin (WM; 100, 200, 300 nM). A, NF-κB-luc reporter gene analysis after stimulation with MALP-2 or LPS for 6 h. B, Supernatants were harvested 18 h after the start of LPS stimulation, and levels of TNF-α and IL-6 measured by ELISA. C, Western blot analysis of p62 protein expression after stimulation with LPS for 18 h. D, HaCaT cells were transfected with a non-specific siRNA (siNS) or siRNAs specific for Beclin-1 (siBec) or Atg5 (siAtg5). After siRNA transfection, cells were stimulated with LPS (100 ng/mL) for 18 h. p62 protein expression was then analyzed by Western blotting. Data represent the mean ± SD of three independent experiments (A and B; C and D, bottom). Images are representative of three independent experiments, all yielding similar results (C and D, top). U, untreated; SC, solvent control (0.1% DMSO); L, LPS.
FIGURE S3. **p62/SQSTM1 is required for LPS-stimulated inflammatory responses in human keratinocytes.** A-C, HaCaT cells were transfected with a non-specific siRNA (siNS) or an siRNA specific for *p62* (sip62). Following transfection, cells were stimulated with LPS (100 ng/mL) for 6 h (A, RT-PCR analysis) or 18 h (B, ELISA analysis). A, Total RNA was isolated from cells and subjected and analyzed for TNF-α, IL-6, LL-37, and p62/SQSTM1 mRNA levels by semi-quantitative RT-PCR. B, TNF-α and IL-6 levels were assessed by ELISA. C, NF-κB-luc reporter gene analysis. HaCaT cells were transiently transfected with a NF-κB reporter construct (experimental conditions as described in Fig. 6C) prior to stimulation with LPS (100 ng/mL) for 6 h, and then assayed for luciferase activity. Data represent the mean ± SD of five independent experiments (B and C). Images are representative of three independent experiments, all yielding similar results (A). Transfection efficiency was assessed by RT-PCR (C, inset). U, untreated; L, LPS.
Figure S1. Lee et al.

A

MALP-2

<table>
<thead>
<tr>
<th></th>
<th>0</th>
<th>6</th>
<th>18</th>
<th>24</th>
<th>48</th>
</tr>
</thead>
<tbody>
<tr>
<td>LC3</td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>β-actin</td>
<td>0</td>
<td>0.1</td>
<td>0.4</td>
<td>1.0</td>
<td>0.2</td>
</tr>
</tbody>
</table>

B

LC3-II/LC3-I

2 μm

1 μm

2 μm

1 μm
Figure S2. Lee et al.

A

**NF-κB luciferase assay**

- **Relative LUC. activity**

- **MALP-2**
  - U SC 3MA WM SC 3MA WM
- **LPS**
  - U SC 3MA WM

B

**Secretion of cytokine (pg/ml)**

- **TNF-α**
  - LPS
  - U SC 3MA WM
- **IL-6**
  - LPS
  - U SC 3MA WM

C

**LPS**

- P62
- β-actin
- 3MA: 10 20 50
- WM: 0.1 0.2 0.3

D

**siNS**

- **siBec1**
- **siAtg5**

- **p62**
- **Bec1**
- **Atg5**
- **β -actin**

**Arbitrary units**

- **p62/β-actin**
  - U SC 3MA WM LPS
- **p62/β-actin**
  - U L siNS siBec1 siAtg5