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CYLD proteolysis protects macrophages from TNF-mediated auto-necroptosis induced by LPS and licensed by type I IFN

Diana Legarda^{1,10}, Scott J. Justus^{1,2,10}, Rosalind L. Ang¹, Nimisha Rikhi¹, Wenjing Li^{3,4}, Thomas M. Moran^{1,3,4}, Jianke Zhang⁵, Emiko Mizoguchi^{6,7}, Matija Zelic⁸, Michelle A. Kelliher⁸, J. Magarian Blander^{1,2,3,9}, and Adrian T. Ting^{1,2,9}

¹Immunology Institute, Icahn School of Medicine at Mount Sinai, New York, NY 10029, USA

²Graduate School of Biological Sciences, Icahn School of Medicine at Mount Sinai, New York, NY 10029, USA

³Department of Microbiology, Icahn School of Medicine at Mount Sinai, New York, NY 10029, USA

⁴Global Health and Emerging Pathogens Institute, Icahn School of Medicine at Mount Sinai, New York, NY 10029, USA

⁵Department of Microbiology and Immunology, Kimmel Cancer Center, Thomas Jefferson University, Philadelphia, PA 19107, USA

⁶Department of Medicine, Gastrointestinal Unit, Massachusetts General Hospital, Boston, MA 02114, USA

⁷Department of Medicine, Center for the Study of Inflammatory Bowel Disease, Massachusetts General Hospital, Boston, MA 02114, USA

⁸Department of Molecular, Cell and Cancer Biology, University of Massachusetts Medical School, Worcester, MA 01605, USA

⁹Tisch Cancer Institute, Department of Medicine, Icahn School of Medicine at Mount Sinai, New York, NY 10029, USA

SUMMARY

Tumor necrosis factor (TNF) induces necroptosis, a RIPK3/MLKL-dependent form of inflammatory cell death. In response to infection by Gram-negative bacteria, multiple receptors on

CONTACT INFORMATION: Adrian T. Ting, Immunology Institute, Icahn School of Medicine at Mount Sinai, One Gustave L. Levy Place, Box 1630, New York, NY 10029, USA. Tel: +1 212 659 9410; Fax: +1 212 849 2525; adrian.ting@mssm.edu. 10Co-first authors

AUTHOR CONTRIBUTIONS

D.L. and S.J.J. designed and performed the majority of the experiments. D.L. and S.J.J. contributed equally to this work. N.R. performed experiments. R.L.A. performed the real-time quantitative PCR experiments. W. L. and T.M.M. provided *Ifnar1^{-/-}* mice. J.Z. provided the femurs and tibia from $Ripk3^{-/-}$ and $Ripk3^{-/-}$ Fadd^{-/-} mice for BMDM isolation. E.M. provided AIEC strain LF82. M.Z. and M.A.K. generated the $RosaCreER^{T2}Ripk1^{f1/f1}$ BMDMs. J.M.B. provided technical and conceptual guidance during the experimental studies and in drafting of the manuscript. D.L., S.J.J., and A.T.T. wrote the manuscript. A.T.T. directed the studies.

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macrophages including TLR4, TNF and type I IFN receptors are concurrently activated but it is unclear how they crosstalk to regulate necroptosis. We report that TLR4 activates CASPASE-8 to cleave and remove the deubiquitinase CYLD in a TRIF- and RIPK1-dependent manner to disable necroptosis in macrophages. Inhibiting CASPASE-8 leads to CYLD-dependent necroptosis caused by the TNF produced in response to TLR4 ligation. While LPS-induced necroptosis was abrogated in $Tnf^{-/-}$ macrophages, a soluble TNF antagonist was not able to do so in $Tnf^{+/+}$ macrophages, indicating that necroptosis occurs in a cell-autonomous manner. Surprisingly, TNF-mediated autonecroptosis of macrophages requires type I IFN, which primes the expression of key necroptosissignaling molecules including TNFR2 and MLKL. Thus, the TNF necroptosis pathway is regulated by both negative and positive crosstalk.

Graphical Abstract



INTRODUCTION

Multiple forms of programmed cell death occur following microbial infection, serving to eliminate infected cells and to mount an appropriate host response (Campisi et al., 2014; Vanden Berghe et al., 2014). Apoptosis, which is predominantly dependent on effector caspases such as CASPASE-3 and -7, is thought to generate a tolerogenic response if it occurs in the absence of an inflammatory signal. Pyroptosis, which is dependent on CASPASE-1 and -11, occurs following activation of the inflammasome by microbial products. Pyroptosis serves to eradicate infected cells and the release of cellular contents and damage-associated molecular patterns (DAMPs) following plasma membrane permeabilization amplifies the inflammatory response (Bergsbaken et al., 2009; Chen and Nunez, 2010). In contrast to apoptosis and pyroptosis, which are dependent on various caspases, necroptosis or programmed necrosis has recently emerged as a form of cell death that occurs in the absence of caspase activity. Similar to pyroptosis, necroptosis is also characterized by plasma membrane permeabilization with the release of DAMPs and thus also induces a pro-inflammatory response. Necroptosis may allow the host to circumvent the blockade of caspase-dependent death pathways that may be imposed by a pathogen that

encodes caspase inhibitors to block apoptosis or pyroptosis and to retain the ability to mount an inflammatory response to signal danger (Chan et al., 2003; Mocarski et al., 2011; Upton et al., 2010). In this regard, inhibition of host caspases by pathogens and subsequent induction of necroptosis functions effectively as a pathogen-sensing event.

One of the best-characterized inducers of necroptotic death is the cytokine TNF, which paradoxically, can also induce a cell survival response within the same cell. Which response is generated is dependent on the ubiquitination status of the signaling molecule RIPK1 following ligation of TNF receptor 1 (TNFR1); non-degradative Lys63-linked ubiquitination of RIPK1 leads to cell survival whereas inhibiting ubiquitination of RIPK1 leads to necroptosis (Justus and Ting, 2015). In some cellular models, blocking ubiquitination (often using SMAC mimetics) causes RIPK1 to first initiate a caspase-signaling cascade leading to apoptosis (O'Donnell et al., 2007; Wang et al., 2008), but if caspases are also blocked (often using zVAD-fmk), then RIPK1 initiates necroptosis (He et al., 2009; O'Donnell et al., 2011). In other cellular models, blocking caspases is sufficient to trigger necroptosis in the presence of TNF (O'Donnell et al., 2011). In the latter models, the fact that a caspase inhibitor switches the TNF response from survival to necroptosis indicates that a caspase normally generates a pro-survival signal. When that survival signal is blocked, necroptosis is switched on. The molecular mechanism underlying this survival versus necroptosis switch has been clarified over the last few years. Following TNFR1 ligation, CASPASE-8, in a complex with FADD and c-FLIP, delivers a pro-survival signal (Dillon et al., 2012) by cleaving and removing the tumor suppressor CYLD (O'Donnell et al., 2011). CYLD is a deubiquitinating enzyme that is essential for TNF-induced necroptosis (Hitomi et al., 2008; O'Donnell et al., 2011; Vanlangenakker et al., 2010). It disassembles Lys63-linked ubiquitination from RIPK1, a requisite step for necroptosis. Removal of CYLD by CASPASE-8 sustains the ubiquitination of RIPK1, leading to a survival response. Thus, the CASPASE-8:CYLD interaction is an early switch that determines survival versus necroptotic death in the TNFR1 pathway.

With the discovery of RIPK3 as an essential molecule in TNF-induced necroptosis (Cho et al., 2009; He et al., 2009; Zhang et al., 2009), the physiological and patho-physiological roles of necroptosis are starting to become clearer. Excessive RIPK3-dependent necroptosis, often revealed by the genetic deletion of CASPASE-8, leads to embryonic lethality (Kaiser et al., 2011; Oberst et al., 2011), mucosal inflammation (Gunther et al., 2011; Welz et al., 2011) and an impaired T cell response (Ch'en et al., 2011). Furthermore, RIPK3-dependent necroptosis has been reported to be beneficial as well as detrimental for the host during bacterial and viral infections (Cho et al., 2009; Duprez et al., 2011; Polykratis et al., 2014; Robinson et al., 2012; Upton et al., 2010). These in vivo studies indicated that RIPK3dependent necroptosis is activated subsequent to an interaction between the pathogen and the host's pathogen-sensing innate cells. Following recognition of LPS derived from Gramnegative bacteria by TLR4 on macrophages, a signature response is the production of TNF. This cytokine is a potent inducer of necroptosis and yet surprisingly, little is known about what effect auto-produced TNF may have on necroptosis in macrophages. In addition to TNF, TLR4 stimulation also leads to the production of type I interferon (IFN) α and β . Type I IFN has also been reported to trigger necroptosis (McComb et al., 2014a; Robinson et al.,

2012; Thapa et al., 2013) but whether type I IFN participates in TNF-mediated necroptosis is unclear.

We report here that TLR4, via the TRIF signaling adaptor, rapidly activates CASPASE-8 to cleave and remove CYLD. This CASPASE-8-mediated cleavage event has a self-preservation function because when it is blocked, macrophages undergo auto-necroptosis caused by the TNF that is produced. Furthermore, the ability of TNF to induce necroptosis in macrophages requires IFN α/β priming because in its absence, macrophages have defective expression of key necroptosis-signaling molecules.

RESULTS

TLR4 activates CASPASE-8 to eliminate the pro-necroptotic molecule CYLD

We previously demonstrated that the deubiquitinating enzyme CYLD is the key substrate cleaved by CASPASE-8 to block TNF-mediated necroptosis (O'Donnell et al., 2011). Upon TNFR1 ligation, CYLD is cleaved by CASPASE-8 at aspartate 215, resulting in the generation of a N-terminal 25 kDa fragment and degradation of the C-terminal fragment containing the catalytic domain (O'Donnell et al., 2011). Since TNF is produced in response to LPS during bacterial infection, we surmised that TNF would activate CASPASE-8 in an autocrine manner to cleave CYLD. Loss of CYLD would then prevent necroptosis and sustain cell survival. Therefore, we sought to confirm that CASPASE-8-mediated cleavage of CYLD occurs after TLR4 ligation. When wild type BMDMs were stimulated with LPS, levels of the 107 kD full-length CYLD (p107) decreased in a time- and dose-dependent manner, coinciding with the appearance of the 25 kD CYLD fragment (p25) generated by CASPASE-8-mediated cleavage (Figure 1A and 1B). This cleavage fragment was evident as early as 30 minutes, implying that CASPASE-8 activity is rapidly induced. This is due to TLR4 as it was not observed in $Tlr4^{-/-}$ BMDMs (Supplemental Figure 1A). The ability of other TLR ligands to induce CYLD cleavage was also tested and only TLR3 and TLR4 ligands could do so after a 4-hour stimulation (Supplemental Figure 1B). Infection of BMDMs with two Gram-negative bacteria, Escherichia coli strain LF82 and Citrobacter rodentium, similarly resulted in the proteolytic processing of CYLD (Figure 1C). Cleavage of CYLD was inhibited by the pan-caspase inhibitor z-VAD-fmk (Supplemental Figure 1C), indicating that this is dependent on the proteolytic activity of caspases. Similar to murine BMDMs, stimulation of the human monocytic cell line THP-1 with LPS also resulted in caspase-dependent cleavage of CYLD (Supplemental Figure 1D). To confirm that the FADD/CASPASE-8 complex cleaves CYLD in macrophages, we compared BMDMs from *Ripk3^{-/-}Casp8^{-/-}* or *Ripk3^{-/-}Fadd^{-/-}* mice to *Ripk3^{-/-}* mice since the *Ripk3* deletion rescued the embryonic lethality of the Casp8-/- and Fadd-/- single knockout mice (Dillon et al., 2012; Kaiser et al., 2011; Oberst et al., 2011). The appearance of the CYLD p25 fragment following LPS stimulation and bacterial infection was detected in *Ripk3*^{-/-} BMDMs, but not in *Casp8^{-/-}Ripk3^{-/-}* (Figure 1D) or *Fadd^{-/-}Ripk3^{-/-}* BMDMs (Supplemental Figure 1E). Thus, the FADD/CASPASE-8 complex rapidly cleaves CYLD after TLR4 stimulation.

TLR4-TRIF signaling and not TNF mediates CASPASE-8 dependent cleavage of CYLD

As discussed earlier, our expectation was that LPS-induced cleavage of CYLD would be dependent on auto-production of TNF, which we sought to confirm using Tnf knockout BMDMs. Surprisingly, the cleavage of CYLD was not dependent on TNF since it was observed in both $Tnf^{+/+}$ and $Tnf^{-/-}$ BMDMs following LPS stimulation (Figure 2A). To test the possibility that the CASPASE-8-mediated cleavage of CYLD may be secondary to another cytokine produced, we pre-treated BMDMs with the protein synthesis inhibitor cycloheximide prior to LPS stimulation. Cycloheximide did not block CYLD cleavage and degradation, whereas it efficiently blocked the re-synthesis of $I_KB\alpha$ following LPS stimulation (Supplemental Figure S2A). Therefore, cleavage of CYLD by CASPASE-8 following TLR4 stimulation is independent of *de novo* protein synthesis. This suggested a direct activation of CASPASE-8 by TLR4, prompting us to examine the involvement of MYD88 and TRIF. Cleavage of CYLD induced by TLR4 stimulation was not affected in the *Myd88*^{-/-} BMDMs (Figure 2B), but was abrogated in the *Trif*^{-/-} BMDMs (Figure 2C), indicating that CASPASE-8 activation was dependent on TRIF. Consistent with the requirement for TRIF in the activation of CASPASE-8, the two molecules are able to form a complex when co-expressed in HEK293 cells (Supplementary Figure S2B). Thus, TLR4 activates CASPASE-8 in a TRIF-dependent but MYD88-independent manner to eliminate CYLD, and not via autocrine production of cytokines.

Since the TNF signaling adaptor RIPK1 has been previously implicated in TLR4 signaling (Cusson-Hermance et al., 2005; Meylan et al., 2004) and RIPK1 and CASPASE-8 can associate (Lin et al., 1999), we asked whether RIPK1 has a role in regulating the activation of CASPASE-8 downstream of TLR4. Treatment with the RIPK1 kinase inhibitor Necrostatin-1 prior to LPS stimulation did not inhibit the cleavage of CYLD in BMDMs (Figure 2D), suggesting that the kinase activity of RIPK1 is not required. However, RIPK1 could still regulate the function of CASPASE-8 through a kinase-independent mechanism. We utilized immortalized BMDMs generated from *RosaCreER^{T2}Ripk1^{f1/f1}* mice and treated those cells with tamoxifen (4-OHT) to acutely delete *Ripk1* (Roderick et al., 2014). LPS-induced cleavage of CYLD was abrogated in the RIPK1-deficient macrophages, demonstrating that this is dependent on RIPK1 (Figure 2E). This dependency on RIPK1 for CYLD cleavage may be related to the reduced level of CASPASE-8 observed in RIPK1-deficient BMDMs suggesting that RIPK1 regulates CASPASE-8 expression in macrophages via a mechanism that is unclear at this point.

TLR4 induces auto-necroptosis in macrophages via production of TNF

It was reported that blocking caspase activity in LPS-stimulated murine macrophages resulted in necroptosis (He et al., 2011; Kaiser et al., 2013). Similarly, human THP-1 cells undergo necroptosis when stimulated with LPS in the presence of zVAD-fmk (Supplementary Figure 2C). Although auto-produced TNF is not required for CASPASE-8 to cleave CYLD following TLR4 stimulation (Figure 2A), whether auto-produced TNF is required for necroptosis once CASPASE-8 is blocked remains questionable. Previous studies have suggested that TLR4 induces necroptosis independently of TNF after examining the response of $Tnf^{-/-}$ or $Tnfr1^{-/-}Tnfr2^{-/-}$ double knockout macrophages to high doses of LPS (He et al., 2011; Kaiser et al., 2013). In light of the possibility that the effect of a genetic

deficiency in a Toll-like receptor signaling molecule can be overlooked due to the use of high amount of the ligand (Bonham et al., 2014; Horng et al., 2002), we conducted our experiments using a lower range of LPS. BMDMs from $Tnf^{+/+}$ and $Tnf^{-/-}$ mice were stimulated with varying concentrations of LPS up to 1 ng/ml for 16 hours. Under these conditions, LPS alone did not induce cell death in wild type BMDMs whereas in the presence of zVAD-fmk, 1 ng/ml of LPS was sufficient to achieve near maximal death of BMDMs (Figure 3A), a concentration that is 20–500 fold lower than that used previously (He et al., 2011; Kaiser et al., 2013). Using this lower range of LPS, necroptosis was abrogated in the $Tnf^{-/-}$ BMDMs (Figure 3A). When we increased the concentration of LPS by up to 100-fold more, the requirement for TNF was diminished even though there was still a difference in viability between $Tnf^{+/+}$ and $Tnf^{-/-}$ macrophages (Supplemental Figure 3A). Thus at lower doses of LPS, there is a strong dependency on TNF to induce necroptosis whereas at higher doses, there is diminished dependency. Similar to LPS, E. coli and C. rodentium infection at a low multiplicity of infection (MOI) resulted in necroptosis that is dependent on TNF (Figure 3B). The LPS or bacterial-induced necroptosis in wild type BMDMs was also inhibited by Necrostatin-1 (Nec-1), indicating a RIPK1 kinase dependency (Supplemental Figure 3B and 3C). These results demonstrate that Gramnegative bacteria, as well as stimulation with limiting doses of LPS, induce necroptosis that is dependent on autocrine production of TNF.

To further confirm the role of autocrine TNF, we examined the role of MYD88 since this TLR signaling adaptor protein is required for TNF production in response to LPS (Kawai et al., 1999). $Myd88^{+/+}$ BMDMs succumbed to necroptosis when treated with LPS or infected with bacteria whereas *Myd88^{-/-}* BMDMs survived (Supplemental Figure S3D). This result is consistent with the fact that LPS-induced TNF expression is MYD88-dependent (Kawai et al., 1999) and which remains true under necroptotic conditions (Supplemental Figure S3E). Since the other TLR4 signaling adaptor TRIF has been implicated in TLR3 and TLR4induced necroptosis (He et al., 2011; Kaiser et al., 2013; McComb et al., 2014a) we also examined whether TRIF is required for necroptosis in our low-dose LPS model. Trif-/-BMDMs were more resistant to necroptosis in response to LPS or bacterial infection (Supplemental Figure S3F). The observation that TLR4-induced necroptosis requires TNF (Figure 3A), as well as TRIF (Supplemental Figure S3F), was somewhat puzzling since TNF is not known to signal through TRIF. In addition to its well-known role in the induction of type IFN, TRIF also has a less appreciated role in TNF production as suggested by a previous report (Gais et al., 2010). We therefore considered the possibility that TNF expression may be compromised in the TRIF-deficient BMDMs, which could provide one explanation for why they are more resistant to necroptosis. Consistent with this possibility, TNF expression after LPS stimulation was defective in the TRIF-deficient BMDMs, regardless of the absence or presence of zVAD-fmk (Supplemental Figure S3G).

In addition to using the *Tnf* and *Myd88* knockouts to demonstrate a requirement for TNF in inducing necroptosis following TLR4 ligation, we also utilized a soluble TNF antagonist to inhibit necroptosis. We confirmed the effectiveness of the murine TNFR1-Fc fusion as an antagonist, as it was able to fully block necroptosis of BMDMs induced by the addition of exogenous TNF (Figure 3C). Surprisingly, it had only a modest effect on necroptosis induced by LPS (Figure 3C). Since the genetic deletion of *Tnf* blocked necroptosis induced

by LPS but the soluble mTNFR1-Fc did not, this strongly suggests that death is not dependent on secreted TNF. This observation suggests that death mediated by TNF occurs in a cell-autonomous manner as a consequence of a ligand-receptor interaction in an intracellular compartment that is not accessible to the soluble mTNFR1-Fc antagonist. We have termed this process auto-necroptosis.

Eliminating CYLD protects macrophages against TLR4-induced auto-necroptosis

We previously showed that removal of CYLD by CASPASE-8 generates a survival response in MEFs but when this cleavage event is inhibited, CYLD-dependent necroptosis ensues (O'Donnell et al., 2011). Removal of CYLD following TLR4 stimulation (Figure 1) may serve a similar pro-survival and anti-necroptotic function in macrophages. Therefore, we examined the functional consequence of the loss of CYLD using BMDMs from control or *Cyld* conditional knockout mice. These conditional knockout mice were generated by crossing *Cyld*^{flox/flox} mice, which contains LoxP sites flanking exons 8 and 9, with the *LysM-Cre* strain to selectively delete CYLD in macrophages (*Cyld* CKO) (Supplemental Figure S4A). Wild type macrophages succumbed to necroptosis in response to LPS and zVAD-fmk whereas CKO cells survived (Figure 4A). A similar result was observed with *E. coli* or *C.* rodentium infection (Figure 4B). To rule out the possibility the CYLD-deficient BMDMs were resistant to necroptosis due to a lack of TNF, we examined TNF expression in the CYLD-deficient BMDMs. Under necroptotic conditions, TNF production was slightly elevated in the CYLD-deficient macrophages, arguing against defective TNF production as a reason for their resistance to necroptosis (Supplemental Figure S4B).

MLKL was recently identified as a key effector molecule of necroptosis. It is phosphorylated by RIPK3 to form an oligomeric structure that translocates to the plasma membrane to disrupt membrane integrity and this can be used as a biochemical signature for necroptosis (Cai et al., 2013; Chen et al., 2013; Dondelinger et al., 2014; Wang et al., 2014). TLR4 engagement in the presence of zVAD-fmk did not lead to MLKL oligomerization in CYLD-deficient cells (Figure 4C), demonstrating that removal of CYLD prevents TLR4 from inducing necroptosis. Since TLR4-induced production of TNF could potentially also activate apoptosis, we tested to see if CYLD was involved in apoptosis. We stimulated BMDMs with LPS after pre-treatment with cycloheximide (CHX) in order to sensitize the cells to apoptosis. Wild type and CYLD-deficient BMDMs were equally sensitive to apoptosis induced by LPS in the presence of cycloheximide (Supplemental Figure S4C) indicating that CYLD is not required for this model of apoptosis. As a control in this experiment, LPS-induced necroptosis was abrogated in CYLD-deficient macrophages (Supplemental Figure S4C).

While removal of CYLD by CASPASE-8 will render macrophages resistant to necroptosis, the possibility remains that CYLD may not be the only substrate cleaved by CASPASE-8 to block necroptosis. To test whether there may be additional CASPASE-8 substrates, we reconstituted $Cyld^{-/-}$ BMDMs with wild type Cyld or the cleavage-resistant $Cyld^{D215A}$ mutant (O'Donnell et al., 2011) and examined whether the D215A mutation negated the need for caspase inhibition. In contrast to our previous observation in MEFs (O'Donnell et al., 2011), macrophages reconstituted with the cleavage-resistant CYLD-D215A did not

undergo necroptosis unless zVAD-fmk was present (Supplementary Figure S4D). This suggests that CYLD may not be the only substrate cleaved by CASPASE-8 following TLR4 ligation. RIPK1 is another necroptosis-signaling molecule that has been implicated as a substrate for CASPASE-8 (Lin et al., 1999; Martinon et al., 2000; McComb et al., 2014b). We therefore used an antibody raised against the N-terminal region of RIPK1 to detect its potential cleavage. LPS stimulation induced the appearance of several smaller fragments of RIPK1 that were abrogated in the presence of zVAD-fmk (Supplementary Figure S4E). This LPS-inducible appearance of these RIPK1 fragments is consistent with the possibility that multiple substrates including CYLD and RIPK1 are cleaved by CASPASE-8 when TLR4 is stimulated. Their removal in a TRIF-dependent manner likely generates a self-preservation signal by preventing the induction of auto-necroptosis in macrophages from the ongoing TNF production during an infection.

TNF-dependent auto-necroptosis in macrophages also requires priming by type I IFN

In addition to TNF production, macrophages also produce type I interferon (IFN) following sensing of Gram-negative bacteria by TLR4 (Doyle et al., 2002; Kawai et al., 2001). Therefore during this interaction between bacteria and macrophages, multiple signaling pathways are engaged including the TLR4, TNFR and IFNAR pathways. How these different pathways cross-regulate each other to control necroptosis is unclear. There is increasing evidence that IFNs can regulate necroptosis (Doyle et al., 2002; Kawai et al., 2001; Robinson et al., 2012; Thapa et al., 2013) but there are conflicting reports on the role of type I IFN in TLR-induced necroptosis (He et al., 2011; Kaiser et al., 2013; McComb et al., 2014a). For these reasons, we wanted to examine whether type I IFN has any role in our low-dose LPS model where we know that autocrine TNF is inducing the necroptosis response. If $nar1^{+/+}$ BMDMs succumbed to necroptosis whereas If $nar1^{-/-}$ BMDMs survived (Figure 5A). Similar to *Ifnar1^{-/-}* BMDMs, *Stat2^{-/-}* BMDMs were also resistant to low-dose LPS-induced necroptosis confirming a requirement for type I IFN signaling in this model (Supplementary Figure S5A). Infection with E. coli or C. rodentium in the presence of zVAD-fmk similarly induced necroptosis in *Ifnar1*^{+/+} but not in *Ifnar1*^{-/-} BMDMs (Figure 5B). The results from Figures 3A and 5A showed a surprising finding: necroptosis of macrophages following TLR4 ligation requires downstream signaling responses generated by both TNF and type I IFN. This raised the question of the relationship between these two cytokines and suggested that induction of necroptosis by TNF itself maybe dependent on IFN α/β . To test this directly, *Ifnar1*^{+/+} and *Ifnar1*^{-/-} BMDMs were stimulated with murine TNF in the presence of zVAD-fmk. *Ifnar1^{-/-}* BMDMs were resistant to necroptosis induced by TNF (Figure 5C). The requirement for IFN α/β reveals an unappreciated pathwayextrinsic regulation of the TNF necroptosis pathway in macrophages by another cytokine. IFN α/β may be functioning to directly induce necroptosis following its expression in response to TLR4 stimulation as suggested by others (McComb et al., 2014b) or it may have an indirect role by priming the cells to respond to TNF. To discriminate between the two mechanisms, we used an IFNAR1-neutralizing monoclonal antibody previously shown to be effective in blocking type I IFN responses (Sheehan et al., 2006). Surprisingly, anti-IFNAR1 did not inhibit necroptosis induced by LPS (Figure 5D) whereas it inhibited STAT1 phosphorylation (Figure 5E), which is known to be dependent on IFN α/β , confirming its neutralizing function. The inability of the neutralizing anti-IFNAR1 to block necroptosis

argues against acutely produced IFN α/β as a direct trigger of necroptosis. Rather, the combined results from the *Ifnar1*^{-/-} BMDMs and the neutralizing anti-IFNAR1 experiments are more consistent with a licensing role for type I IFN in priming macrophages for necroptosis.

Type I IFN licenses TNF-mediated necroptosis by inducing TNFR2 and MLKL expression

Why TNF requires IFNAR signaling to induce necroptosis was unclear. The defect in the TNF necroptosis pathway in the *Ifnar1*^{-/-} macrophages could be at the level of the TNF receptors or further downstream. Our group and others have shown that in order for TNFR1 to induce necroptosis, ubiquitination of RIPK1 has to be disrupted to switch RIPK1 from a pro-survival to a pro-death signaling molecule (Arslan and Scheidereit, 2011; O'Donnell et al., 2012; O'Donnell and Ting, 2011). Physiologically, this is achieved by the simultaneous ligation of TNFR1 and TNFR2 whereby TNFR2 ligation leads to the rapid degradation of the TRAF2/cIAP1/2 E3 ubiquitin ligase complex (Chan and Lenardo, 2000; Fotin-Mleczek et al., 2002; Li et al., 2002; Ruspi et al., 2013). The loss of TRAF2/cIAP1/2 reduced RIPK1 ubiquitination, facilitating the conversion of RIPK1 to a death-signaling molecule upon TNFR1 ligation. Consistent with this model, TNFR2 was shown to be necessary for necroptosis in T cells (Chan et al., 2003). Therefore, we hypothesized that there may be a defect with TNFR2 expression in the *Ifnar1*^{-/-} BMDMs. We first confirmed that TNFR2 has</sup> a role in TNF-induced necroptosis in macrophages. We treated wild type BMDMs with either murine TNF, which binds both murine TNFR1 and TNFR2, or with human TNF, which binds only to murine TNFR1 (Lewis et al., 1991). Murine TNF induced necroptosis whereas human TNF did not (Figure 6A), suggesting that ligation of both TNFR1 and TNFR2 is needed for necroptosis. Furthermore, murine TNF was able to induce necroptosis in $Tnfr2^{+/+}$ but not $Tnfr2^{-/-}$ BMDMs (Figure 6B), confirming the requirement for TNFR2.

While TNFR1 expression is quite ubiquitous, TNFR2 expression appears to be more restricted and is induced on macrophages and other hematopoietic cells upon cellular activation (Bethea et al., 1997; de Kossodo et al., 1994; Tannenbaum et al., 1993). We thus hypothesized that TNFR2 expression induced by LPS stimulation is dependent on IFN α/β . Using flow cytometry, we found that LPS induced the expression of TNFR2 in *Ifnar1*^{+/+} but not in *Ifnar1*^{-/-} BMDMs (Figure 6C). Similar to the *Ifnar1*^{-/-} BMDMs, TNFR2 expression was also compromised in the *Stat2*^{-/-} BMDMs (Supplementary Figure S5B). However, the level of *Tnfr2* mRNA was not reduced in the *Stat2*^{-/-} BMDMs (Supplementary Figure S5C), suggesting that a STAT2-dependent gene may regulate TNFR2 expression on *Ifnar1*^{-/-} BMDMs could affect necroptosis, we compared the sensitivity of *Tnfr2*^{+/+} and *Tnfr2*^{-/-} BMDMs to necroptosis induced by LPS or bacteria infection. Consistent with our hypothesis, *Tnfr2*^{-/-} BMDMs were more resistant to necroptosis than *Tnfr2*^{+/+} BMDMs (Figure 6D), similar to that seen in *Tnfr*^{-/-} BMDMs (Supplemental Figure S5D).

Since IFN α/β induces a large array of genes collectively known as IFN-inducible genes (ISG), it is possible that in addition to TNFR2, IFNAR1 deficiency may also affect the expression of other necroptosis molecules. Of particular interest is MLKL, the RIPK3 kinase substrate that is essential for TNF-induced necroptosis (Sun et al., 2012; Zhao et al., 2012),

which was recently shown to be an ISG (Dillon et al., 2014; Thapa et al., 2013). We examined the protein expression level of MLKL in *Ifnar1*^{+/+} and *Ifnar1*^{-/-} BMDMs after stimulation with varying doses of LPS. In contrast to other necroptosis signaling molecules including CYLD, RIPK1 and RIPK3, MLKL protein expression was reduced in *Ifnar1*^{-/-} BMDMs (Figure 6E). Since MLKL-deficient macrophages were shown to be resistant to necroptosis (Murphy et al., 2013; Wu et al., 2013), the diminished MLKL expression in the *Ifnar1*^{-/-} BMDMs also likely contributes to their resistance to necroptosis. Accordingly, MLKL oligomerization was also severely impaired in *Ifnar1*^{-/-} BMDMs (Figure 6F). In *Stat2*^{-/-} BMDMs, the basal level of MLKL was also reduced (Supplementary Figure S6A), albeit to a lesser degree than in the *Ifnar1*^{-/-} BMDMs. MLKL oligomerization was also impaired in the *Stat2*^{-/-} BMDMs (Supplementary Figure S6B). Consistent with *Mlk1* being an ISG, its mRNA level was significantly reduced in IFNAR1 and STAT2-deficient BMDMs (Supplementary Figure S6C).

These results thus far support the model that defective TNFR2 and MLKL expression in the absence of IFN α/β priming contributes to the resistance of the *Ifnar1*^{-/-} BMDMs to necroptosis. To test whether TNFR2 and MLKL expression are the only defects or whether additional defects exist, we complemented *Ifnar1*^{-/-} BMDMs with *Tnfr2* or *Mlk1*. Retroviral transduction of *Ifnar1*^{-/-} BMDMs with *Tnfr2* restored the expression of TNFR2 but was not sufficient to reverse their resistance to necroptosis (Figure 6G). Likewise, retroviral transduction of *Ifnar1*^{-/-} BMDMs with *Mlk1* restored their expression of MLKL but this was also insufficient to reverse their resistance to necroptosis (Figure 6G). Simultaneous reconstitution of *Ifnar1*^{-/-} BMDMs with both TNFR2 and MLKL also failed to reverse the resistance of these cells to necroptosis (Figure 6G). This suggests that in addition to TNFR2 and MLKL, there are additional defects in the TNF necroptosis pathway in the absence of IFN α/β priming. Nonetheless, the reduction in TNFR2 and MLKL observed in the *Ifnar1*^{-/-} cells plays a key role as evidenced by the resistance of the TNFR2 and MLKL-deficient macrophages to necroptosis. This underscores the critical licensing role played by type I IFN in TNF-mediated necroptosis.

DISCUSSION

This current study shows that the determination of survival versus death in macrophages is controlled by multiple pathways (Figure 7). We demonstrated that TLR4 activates CASPASE-8 to proteolyze and remove CYLD. This removal is a mechanism for protecting against necroptosis that can be triggered by the TNF produced during an infection by Gramnegative bacteria. We had previously reported the existence of an early NF-kappaB-independent but ubiquitin-dependent cell death checkpoint in the TNFR1 pathway (Justus and Ting, 2015; Legarda-Addison et al., 2009; O'Donnell et al., 2007; O'Donnell and Ting, 2010). Thus, TLR4 via its activation of CASPASE-8 sustains this early ubiquitin-dependent checkpoint in the TNFR1 pathway by removing the CYLD deubiquitinase. However, when CASPASE-8 is inhibited, this early ubiquitin-dependent checkpoint is disrupted and macrophages undergo auto-necroptosis mediated by the TNF that is produced during the infection. This process occurs in a cell-autonomous manner, likely mediated by a ligand-receptor interaction in cis. This auto-necroptosis response further requires licensing by type I IFN, which is necessary for the expression of TNFR2, MLKL and potentially other

necroptosis-signaling molecules. In the absence of TNFR2 and MLKL, the ability of TNF to induce necroptosis is severely compromised. Our data thus demonstrates an unappreciated pathway-extrinsic regulation of the TNF necroptosis pathway by the TLR4 and type I IFN receptors in a negative and positive manner, respectively.

We previously demonstrated that whether CYLD is cleaved by CASPASE-8 or not is the key determinant of survival versus necroptotic death in MEFs following TNFR1 ligation (O'Donnell et al., 2011). Our current study demonstrates that TLR4, in a TRIF-dependent mechanism, activates CASPASE-8 to remove CYLD and this has an inhibitory effect on the TNF death pathway. We have genetically defined some of requirements for the cleavage of CYLD: it requires TRIF, RIPK1, FADD and CASPASE-8. An interaction between TRIF and CASPASE-8 has been suggested previously (Estornes et al., 2012; Maelfait et al., 2008). Indeed, the two molecules can interact in co-transfection experiments. Alternatively, TRIF can interact with RIPK1 in a RHIM-dependent manner (Kaiser and Offermann, 2005; Meylan et al., 2004) and in turn, RIPK1 can recruit CASPASE-8 (Wang et al., 2008). Therefore it is likely that TRIF, RIPK1, FADD and CASPASE-8 form a proteolytic complex that confers cell survival when macrophages are activated. This remains to be directly confirmed because the analysis of endogenous TRIF-containing complexes in primary macrophages has been hampered by the lack of suitable immunoprecipitating antibodies against TRIF. Our current analysis of how TLR4 regulates CYLD cleavage in macrophages has revealed a couple of aspects that are different from our previous analysis of CYLD cleavage in MEFs. While RIPK1 was not required for CYLD cleavage in response to TNFR1 ligation in MEFs (O'Donnell et al., 2011), it appears to have a role in the TLR4induced response in macrophages, in part by regulating the expression of CASPASE-8 itself, or RIPK1 could also serve to recruit CASPASE-8 to TRIF. Another differing aspect is that we did not detect any RIPK1 cleavage in response to TNFR1 stimulation in MEFs whereas this was observed in response to TLR4. In conjunction, mutating the cleavage site of CYLD was sufficient to sensitize MEFs to undergo TNF-induced necroptosis and negating the need for caspase inhibition, whereas this was insufficient in macrophages. Thus, multiple substrates may be cleaved by CASPASE-8 in macrophages following TLR4 stimulation, indicating receptor and cell type differences in the regulation of CYLD cleavage.

While TNF is not required for the CYLD cleavage observed upon TLR4 stimulation, it is required for the necroptosis induced upon caspase inhibition. Our result differs from previous studies suggesting that TNF is not required (He et al., 2011; Kaiser et al., 2013; McComb et al., 2014a; Robinson et al., 2012) and this is likely due to the low doses of LPS used in the current study compared to the high doses used in the previous studies. We further confirmed that the LPS results were reflective of that seen during bacterial infection by infecting macrophages with a low MOI of viable *C. rodentium* and *E. coli*. The genetic requirement for a molecule in a TLR pathway could be missed when higher doses of ligand are utilized as illustrated by Bonham et al. in their study of the adaptor molecule TIRAP in TLR9 signaling (Bonham et al., 2014). Surprisingly, we were unable to block this TNF-dependent death process using a soluble TNFR1-Fc antagonist indicating that this is not mediated by secreted TNF. Instead, this observation strongly suggests that TNF is engaging its receptors within the cell and activating downstream signaling in a cell-autonomous manner. The biological reason for why macrophages are wired to undergo auto-necroptosis

upon caspase inhibition is unclear. We speculate that this may serve a tripwire function in a sentinel cell such as a macrophage where pathogen or host-mediated alterations in caspase activity following an infection could trigger necroptosis as a danger signal.

In contrast to the inhibitory effect TLR4 exerts on the TNF necroptosis pathway, type I IFN receptor signaling exerts a positive effect on this death pathway in a STAT2-dependent manner. Our analysis unequivocally showed that LPS and TNF-induced necroptosis required IFN α/β -mediated responses. This requirement for type I IFN in order for TNF to trigger necroptosis was quite unexpected and prompted us to examine the basis of this crosstalk. We initially directed our attention at TNFR2 and found that its expression after LPS stimulation was impaired in the *Ifnar1*^{-/-} macrophages and it is required for necroptosis to occur. In addition to TNFR2, expression of MLKL was also reduced in the *Ifnar1*^{-/-} BMDMs. There are likely additional defects in the *Ifnar1*^{-/-} cells, as reconstitution with *Tnfr2* and *Mlk1* did not reverse their resistance to death. Since the other molecules essential for TNF necroptosis described to date (CYLD, RIPK1 and RIPK3) are present in the *Ifnar1*^{-/-} BMDMs, it is likely other molecules essential for necroptosis remain to be characterized.

The requirement for IFNAR1 raises the question whether IFN α/β produced following sensing of microbes directly induces necroptosis, as suggested by a recent report (McComb et al., 2014a), or whether basal level of IFN α/β in BMDMs cultures prime these cells to respond to TNF. Our finding that a neutralizing anti-IFNAR1 did not inhibit necroptosis of BMDMs indicates that the IFN α/β acutely produced in response to LPS does not directly trigger necroptosis. Rather, prior exposure of the BMDMs to IFN α/β is necessary to license and prime the necroptosis pathway via the induction of essential signaling molecules. The requirement of IFNAR1 as an essential regulator for necroptosis is likely restricted to certain cell types as suggested by genetic crosses involving the Casp8-/- mice. The embryonic lethality of $Casp8^{-/-}$ mice was rescued by crossing the mice to $Ripk3^{-/-}$ mice (Kaiser et al., 2011; Oberst et al., 2011) but not by crossing to $Stat1^{-/-}$ mice that are defective in type I IFN signaling (Thapa et al., 2013), indicating that the necroptosis pathway in the cell type affected in the Casp8^{-/-} embryos is not dependent on IFN signaling. A report by Duprez et al demonstrated that RIPK3-dependent necroptosis was responsible for the morbidity and mortality in a mouse model of sterile sepsis induced by TNF (Duprez et al., 2011). Prior to the study by Duprez et al., Huys et al., using a similar model of TNF-induced sepsis, showed that IFNAR1 was essential for driving this disease (Huys et al., 2009), further indicating that the interplay of type I IFN and TNF is relevant in vivo.

In summary, our studies showed that necroptosis of macrophages is regulated in a complex manner by multiple crosstalk between the TLR4, TNF and IFN receptors and highlights the insights gained from studying the TNF death pathway in the context of other pathways that are concurrently activated.

EXPERIMENTAL PROCEDURES

Generation of CYLD conditional knockout mice

Mice expressing loxP-flanked (floxed) *Cyld* exon 8 and 9 were generated by Ingenious Targeting Laboratories (Ronkonkoma, New York). These mice were backcrossed to

C57BL/6 for 10 generations, and were then crossed with *LysM-Cre* transgenic mice, which express the Cre recombinase from a myeloid-specific promoter, to selectively delete CYLD from monocytes and macrophages (*LysM-Cre Cyldflox/flox*, or CYLD conditional knockout, or CYLD CKO). For all experiments performed, *Cyldflox/flox* littermates were used as controls (CYLD WT).

Bone Marrow-Derived Macrophages (BMDMs)

BMDMs were generated by culturing bone marrow progenitors in L929 conditioned medium. Immortalization of BMDM from *Ripk1* conditional knockouts was conducted by infection with J2 oncovirus (Roberson and Walker, 1988).

Cell Death Assays

Cell death was analyzed by propidium iodide staining followed by flow cytometry or with the CellTiter-Glo Luminescent Cell Viability Assay kit (Promega).

Western blot analysis

Cellullar lysates obtained using 1% Triton X-100 were resolved by reducing SDS-PAGE. For MLKL oligomerization, lysates generated using either 2% SDS or M2 buffer containing 1% Triton X-100 were resolved by SDS-PAGE under non-reducing conditions (without beta-mercaptoethanol). Blotting was conducted with standard techniques.

Statistical Analysis

Cell death, cell viability, and intracellular staining data are presented as the mean \pm standard deviation. Statistics were performed using T-test. * p<0.05, ** p<0.01, *** p<0.001. Quantitative real-time PCR data is presented as the mean, \pm standard deviation. Statistics were performed using Two-Way ANOVA. Bonferroni posts-tests were performed to compare the triplicate means at different time points. ** p<0.01, *** p<0.001.

Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

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HIGHLIGHTS

- TLR4 activates CASPASE-8 to remove CYLD and disable the TNF necroptosis pathway.
- Autocrine TNF induces necroptosis in a cell-autonomous manner.
- Priming of macrophages by Type I IFN is needed for necroptosis induced by LPS or TNF.
- Type I IFN primes TNFR2 and MLKL expression, which are needed to induce necroptosis.

CYLD p107

CYLD p25

GAPDH

LPS (ng/ml)

50 100 200

10





(A–B) BMDMs from wild type mice were treated with 100 ng/ml of LPS for the indicated times (A) or with the indicated concentrations of LPS for 4 hours (B). (C) BMDMs from wild type BMDMs were infected with viable *E. coli* or *C. rodentium* for 4 hours. (D) BMDMs from wild type, $Ripk3^{-/-}$, and $Ripk3^{-/-}Casp8^{-/-}$ mice were stimulated with either LPS (100 ng/ml), *E. coli* (MOI 1), or *C. rodentium* (MOI 1) for the indicated times. Lysates were subject to western blot analysis using CYLD (D1A10), RIPK3, CASPASE-8, and GAPDH antibodies. All experiments shown are representative of at least three independent experiments with similar results.

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FIGURE 2. TLR4-TRIF signaling and not TNF mediates CASPASE-8 dependent cleavage of CYLD

(A) Wild type and *Tnf*^{-/-} BMDMs were treated with 100 ng/ml of LPS for 4 hours. (B–C) Wild type, *Myd88*^{-/-}, and *Trif*^{-/-} BMDMs were treated with 10 ng/ml of LPS for the indicated times. (D) Wild type BMDMs were pre-treated with 30 μM Necrostatin-1 prior to stimulation with LPS for 4 hours. (E) Immortalized BMDMs generated from *RosaCreER*^{T2}*Ripk1*^{fl/fl} mice were treated with ethanol or with tamoxifen (4-OHT) to delete RIPK1. Cells were then treated with 100 ng/ml of LPS for the indicated times. Lysates were blotted as indicated. All experiments shown are representative of at least three independent experiments with similar results.

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FIGURE 3. TLR4 induces auto-necroptosis in macrophages via production of TNF

(A–B) BMDMs from $Tnf^{+/+}$ or $Tnf^{-/-}$ mice were pre-treated with 25 µM zVAD-fmk or DMSO prior to stimulation with the indicated concentrations of LPS (A) or infected with *E. coli* or *C. rodentium* at the indicated multiplicity of infection (MOI) (B) for 16 hours. Cell death was measured by propidium iodide staining. Data is presented as the mean of three independent experiments performed in triplicate, ± standard deviation. Statistics were performed using T-test. * p<0.05, ** p<0.01, *** p<0.001. (C) Wild type BMDMs were treated with control-FC or TNFR1-FC, in addition to either LPS or TNF, for 24 hours. Viability was determined by CellTiter-Glo assays. Data is presented as the mean of three independent experiments performed in triplicate, ± standard deviation.



FIGURE 4. Elimination of CYLD protects macrophages from TLR4-induced auto-necroptosis (A) BMDMs from *Cyld*^{flox/flox} (CYLD WT) and *LysM-Cre x Cyld*^{flox/flox} mice (CYLD conditional knockout, or CKO) were pre-treated with DMSO or 25 μ M zVAD-fmk and stimulated with LPS for 16 hours. (B) BMDMs from CYLD WT or CYLD CKO were pre-treated with 25 μ M zVAD-fmk prior to infection with *E. coli* or *C. rodentium* for 16 hours at the indicated MOI. (A–B) Cell death was analyzed as in Figure 3A. * p<0.05, ** p<0.01, *** p<0.001. (C) *Cyld*^{+/+} or *Cyld*^{-/-} BMDMs pre-treated with 25 μ M zVAD-fmk were stimulated with 1 ng/ml of LPS for the indicated time. Triton-soluble lysates were subject to non-reducing SDS-PAGE, and blotted with anti-MLKL and β -actin. The experiment shown is representative of three independent experiments.

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FIGURE 5. TLR4-induced auto-necroptosis in macrophages also requires type I IFN

Ifnar1^{+/+} and *Ifnar1*^{-/-} BMDMs were pre-treated with 25 μ M zVAD-fmk prior to stimulation with LPS (A), viable *E. coli* or *C. rodentium* (B), or murine TNF (C) for 24 hours. (D) Wild type BMDMs were treated with isotype control or 1 ug/ml of IFNAR1 neutralizing antibody prior to stimulation with the indicated concentrations of LPS for 24 hours. Cell death in (A–D) was measured by propidium iodide staining or by CellTiter-Glo as in Figure 3. * p<0.05, ** p<0.01, *** p<0.001. (E) Wild type BMDMs were treated as in (D) prior to stimulation with LPS for 24 hours. Lysates were blotted with antibodies to

phosphorylated STAT1 and total STAT1. Experiment shown is representative of at least three experiments performed independently.



FIGURE 6. Type I IFN licenses TNF-mediated necroptosis by inducing TNFR2 and MLKL expression

(A) Wild type BMDMs were stimulated with indicated amounts of murine or human TNF for 24 hours in the presence of zVAD-fmk. (B) $Tnfr2^{+/+}$ and $Tnfr2^{-/-}$ BMDMs were stimulated with murine TNF for 24 hours in the presence of zVAD-fmk. (A, B) Viability was determined by CellTiter-Glo assays as in Figure 3C. (C) $Ifnar1^{+/+}$ and $Ifnar1^{-/-}$ BMDMs stimulated with 1 ng/ml of LPS for 24 hours were stained with a PE-conjugated anti-TNFR2. Data is representative of three independent experiments. (D) Necroptosis in $Tnfr2^{+/+}$ and $Tnfr2^{-/-}$ BMDMs were analyzed as in Figure 3A & B. (E) $Ifnar1^{+/+}$ or $Ifnar1^{-/-}$ BMDMs were stimulated with the indicated concentrations of LPS for 4 hours.

Triton-soluble lysates were blotted with antibodies for CYLD (D1A10), RIPK1, RIPK3, MLKL, and GAPDH. (F) *Ifnar1*^{+/+} or *Ifnar1*^{-/-} BMDMs were pre-treated with 25 μ M zVAD-fmk and stimulated with 10 ng/ml of LPS for the indicated times. MLKL oligomerization was analyzed as in Figure 4C. Experiments were repeated twice with similar results. (G) *Ifnar1*^{-/-} BMDMs were transduced with retroviruses encoding a control protein, TNFR2, or MLKL, as indicated. Lysates from these cells were blotted to confirm MLKL expression. *Ifnar1*^{-/-} BMDMs transduced with TNFR2 or MLKL were stained with PE-conjugated anti-TNFR2 to confirm TNFR2 expression that is equivalent to that of *Ifnar1*^{+/+} BMDMs stimulated with 10 ng/ml of LPS for 24 hours. These cells were stimulated with LPS in the presence of 25uM of zVAD-fmk. Viability was determined by CellTiter-Glo as in Figure 3C.



FIGURE 7. Regulation of survival and necroptosis in macrophages

During an infection, the determination of TNF-induced necroptosis is subject to crossregulation by other receptor pathways. I: TLR4 activates CASPASE-8 through a TRIFdependent manner to proteolyze and remove CYLD. CYLD is essential for TNF-induced necroptosis, and so its removal prevents macrophages from undergoing necroptosis that can be induced by the TNF produced during bacterial infection. II: Inhibition of CASPASE-8 stabilizes CYLD and this results in necroptosis mediated by auto-produced TNF signaling via CYLD. Type I IFN plays a licensing role because it is required for the induction of TNFR2, MLKL and other uncharacterized molecules that are essential for TNF-mediated necroptosis.