# Prevention of Hepatic Apoptosis and Embryonic Lethality in *RelA/TNFR-1* Double Knockout Mice

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**Mice deficient in the nuclear factor** <sup>k</sup>**B (NF-**k**B)-transactivating gene** *RelA* **(***p65***) die at embryonic days 14–15 with massive liver apoptosis. In the adult liver, activation of the NF-**k**B heterodimer RelA/p50 can cause hepatocyte proliferation, apoptosis, or the induction of acute-phase response genes. We examined, during wild-type fetal liver development, the expression of the Rel family member proteins, as well as other proteins known to be important for NF-**k**B activation. We found these proteins and active NF-**k**B complexes in the developing liver from at least 2 days before the onset of lethality observed in** *RelA* **knockouts. This suggests that the timing of NF-**k**B activation is not related to the timing of lethality. We therefore hypothesized that, in the absence of** *RelA***, embryos were sensitized to tumor necrosis factor (TNF) receptor 1 (TNFR-1)-mediated apoptosis. Thus, we generated mice that were deficient in both** *RelA* **and** *TNFR-1* **to determine whether apoptotic signaling through TNFR-1 was responsible for the lethal phenotype.** *RelA/TNFR-1* **double knockout mice survived embryonic development and were born with normal livers without evidence of increased hepatocyte apoptosis. These animals became runted shortly after birth and survived an average of 10 days, dying from acute hepatitis with an extensive hepatic infiltration of immature neutrophils. We conclude that neither** *RelA* **nor** *TNFR-1* **is required for liver development and that RelA protects the embryonic liver from TNFR-1-mediated apoptotic signals. However, the absence of both TNFR-1 signaling and RelA activity in newborn mice makes these animals susceptible to endogenous hepatic infection.** *(Am J Pathol 2000, 156:997–1007)*

The NF- $\kappa$ B transcription factor modulates gene expression in many cellular responses, including inflammation, apoptosis, and liver regeneration.<sup>1–3</sup> The NF- $\kappa$ B transactivating subunit, RelA (p65), plays a critical role in mouse liver development because RelA<sup>-/-</sup> mice die during mid-gestation from massive hepatocyte apoptosis.<sup>4,5</sup> Whereas much work has been done establishing

the roles of NF- $\kappa$ B/Rel proteins in adult tissues, (for recent review, see<sup>6</sup>), little is known about their functions in the developing animal. Except for RelA<sup>-/-</sup> mice, other Rel knockouts have not demonstrated critical roles for Rel proteins during development.<sup>7-9</sup> Schmidt-Ullrich et al generated transgenic mice expressing a NF-kB-driven LacZ reporter gene construct to provide information on the timing and cellular location of  $NF$ - $\kappa$ B activation during murine development. NF-<sub>K</sub>B activity was detected late in development, with no evidence of LacZ-positive cells until E12.5. Furthermore, NF- $\kappa$ B transactivation was predominantly noted within the developing brain, spinal medulla, and thymus of their transgenic lines. Only one of their transgenic lines showed any expression in the fetal liver, starting at E13.<sup>10</sup> Thus, the finding by Beg et al that  *embryos die of massive liver apoptosis is partic*ularly intriguing because of the lack of demonstration of NF- $\kappa$ B activation during mouse liver development.

The tumor necrosis factor (TNF) rapidly and potently activates NF-kB in a variety of cell types, including adult hepatocytes. TNF mediates its cellular responses by signaling through two receptors, TNFR-1 and TNFR-2, but most of its biological functions are signaled through TNFR-1.11,12 TNFR-1 is a death domain-containing receptor that, on ligand binding and trimerization, recruits the adapter protein TRADD. The signals elicited in response to TNF bifurcate at the level of TRADD binding. The overexpression of TRADD stimulates both apoptosis and  $NF-\kappa B$  activation through differential binding domains.<sup>13</sup> The C-terminal portion of TRADD binds the death domain-containing protein FADD, resulting in apoptosis.14 The N-terminal portion of TRADD binds the TNF receptor-associated factor 2 (TRAF-2), and other more recently identified proteins such as RIP, ultimately resulting in the activation of  $NF- $\kappa$ B. <sup>15,16</sup>$ 

NF- $\kappa$ B activation is initiated by its release from the cytoplasmic inhibitory proteins  $I_{\kappa}B_{\alpha}$  or  $I_{\kappa}B_{\beta}$ .<sup>17</sup> Two  $I_{\kappa}B$ kinases, IKK-1 and IKK-2, which are responsible for phosphorylating I<sub>K</sub>B proteins, are components of a large multisubunit complex.<sup>18,19</sup> Phosphorylation of  $I_{\kappa}B$  leads to its ubiquitination and subsequent degradation by the proteasome, resulting in nuclear translocation of NF-

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<sup>k</sup>B.20,21 Recent work has shown that *IKK-2* knockouts die during development of liver degeneration and apoptosis, similar to RelA knockouts.<sup>22,23</sup> This suggests that IKK-2 is directly responsible for activating NF- $\kappa$ B in the developing liver.

Work from this and other laboratories demonstrated that TNF is required for the initiation of liver regeneration in mice, and that its proliferative effect on hepatocytes is strictly dependent on signaling through TNFR-1.<sup>3,24</sup> Mice lacking TNFR-1 have high mortality during the early stages of liver regeneration and the surviving animals have a deficit in DNA replication. *TNFR-1<sup>-/-</sup>* mice fail to activate the transcription factors NF-kB and STAT3 at the start of liver regeneration. However, DNA synthesis can be restored in these animals by interleukin 6 (IL-6) injection, which also restores STAT3 but not NF- $\kappa$ B activation.<sup>3</sup> These and other experiments<sup>25,26</sup> show that, on one hand, TNF, signaling through TNFR-1 and downstream activation of NF-kB, IL-6, and STAT3, is an important proliferative agent for hepatocytes. On the other hand, direct blockage of NF-kB activation, both *in vivo* and in hepatocyte cultures, switches the biological effect of TNF from proliferative to apoptotic. $27-30$  If, as in liver regeneration, *TNFR-1<sup>-/-</sup>* mice fail to activate NF- $\kappa$ B during liver development, it is surprising that these mice develop normally and show no obvious signs of hepatic apoptosis.

To examine whether TNFR-1-mediated signaling is responsible for the lethal phenotype of *RelA* knockouts, we generated mice deficient in both *RelA* and *TNFR-1*. We show that double-knockout mice survive embryonic development and that gestation in these animals is apparently unaffected because newborn mice have normal liver morphology with little detectable apoptosis. Thus, *RelA<sup>-/-</sup>* mice die from apoptotic signals mediated through TNFR-1 during liver development. RelA and p50, the components necessary for  $NF-\kappa B$  activation, are present during fetal liver development in wild-type (WT) mice from at least E12, suggesting that the timing of NF- $\kappa$ B activation is not related to the timing of embryonic lethality in *ReIA<sup>-/-</sup>* mice. Survival of the *ReIA/TNFR-1* double knockouts suggests that neither *RelA* nor *TNFR-1* is critical for hepatic development in mice. However, newborn animals with these deficiencies become sensitive to infection and the neutrophilic invasion that cause damage to the liver and other organs, resulting in early postnatal mortality.

# Materials and Methods

# *Animals*

The C57BL/6J (WT) mice were obtained from Jackson Laboratories (Bar Harbor, ME).The *RelA* heterozygous breeder mice were obtained from Amer Beg (Columbia University, New York, NY), and the *TNFR-1* knockout mice have been previously described.<sup>3</sup> To obtain fetal samples, timed matings were performed, and the presence of a vaginal plug the following morning was considered embryonic day 0 (E0). To generate mice deficient in

both *RelA* and *TNFR-1*, *RelA<sup>+/-</sup>* females were crossed with *TNFR-1<sup>-/-</sup>* males producing double heterozygous *RelA*<sup>+/-</sup>/*TNFR-1*<sup>+/-</sup> offspring that were subsequently back crossed. The breeding cages from *ReIA<sup>+/-</sup>/TNFR-* $1^{+/-}$  matings were examined daily, and the first day that pups were found in the cage was considered day 1 of postnatal life. The number of pups born to each female was recorded, and, within 24 hours, a small piece of tail was removed from each newborn for DNA analysis.

Animals were maintained in a specific pathogen-free facility under 12-hour dark/light cycles and given standard diet and water *ad libitum*. All animal work was in accordance with policies at the University of Washington.

# *Isolation of Murine Liver Samples*

For isolation of fetal livers, pregnant mice were sacrificed between E12 and E19. Fetuses were dissected free of extra embryonic membranes and decidual tissue and placed in phosphate-buffered saline; livers were carefully removed under a dissecting microscope at  $30\times$ . For the isolation of neonatal livers, newborn pups were killed at 2, 4, 7, and 16 days of age, and livers were carefully dissected from the animal.

# *Polymerase Chain Reaction (PCR) Genotyping*

For fetal samples, a portion of the brain was isolated from each sample. For newborns, a small piece of tail was removed. All samples were digested in GNTK buffer, (50 mmol/L KCl, 1.5 mmol/L MgCl<sub>2</sub>, 10 mmol/L Tris-HCl, pH 8.5, 0.01% gelatin, 0.45% Nonidet P-40, 0.45% Tween-20) supplemented with 100  $\mu$ g/ml proteinase K at 55°C. Proteinase K was heat inactivated, and  $1-3$   $\mu$  of the extract was used in PCR. For *RelA* PCR, a three-primer reaction amplified a 120-bp fragment of the WT allele and a 160-bp fragment of the *RelA* mutant allele. For *TNFR-1*, a four-primer PCR reaction amplified a 120-bp fragment of the WT allele and a 155-bp fragment of the mutant allele. The PCR products were run on 2% agarose (SeaKem, FMC BioProducts, Rockland, ME) gels and visualized with a UVP gel documentation system (Upland, CA).

# *Protein Extractions*

Whole-cell extracts from the fetal livers were isolated by resuspension in a tissue lysis buffer (20 mmol/L 4-(2 hydroxyethyl)-1-piperazineethanesulfonic acid (HEPES),  $pH$  7.5, 2 mmol/L ethylene glycol bis( $\beta$ -aminoethyl ether)-*N,N,N',N'*-tetraacetic acid (EGTA), 10% glycerol, 1% Tween-20, 1 mmol/L dithiothreitol, 0.5 mmol/L phenylmethyl sulfonyl fluoride, 0.5 mmol/L benzamidine-HCl, 10  $\mu$ g/ml aprotinin, 10  $\mu$ g/ml leupeptin, 10  $\mu$ g/ml pepstatin A) and incubated on ice for 30 minutes. Extracts were cleared by centrifugation at  $4^{\circ}$ C at 16,000  $\times$  *g* for 15 minutes and stored at  $-80^{\circ}$ C. Whole-cell extracts from the newborn-liver samples were homogenized in a glass homogenizer in 0.5–1 ml of the tissue lysis buffer before incubation on ice and centrifugation.

Nuclear extracts from the fetal liver were isolated in a manner similar to the methods described by Han and Brasier.<sup>31</sup> All buffers were supplemented with the following protease inhibitors before use:  $240 \mu$ g/ml antipain, 2  $\mu$ g/ml aprotinin, 0.01 mol/L benzamidine-HCl, 0.2 mmol/L dithiothreitol, 10  $\mu$ g/ml leupeptin, 10  $\mu$ g/ml pepstatin A, 0.5 mmol/L phenylmethyl sulfonyl fluoride, 0.15  $\mu$ mol/L spermine,  $0.5 \mu$ mol/L spermidine. In a brief description of the procedure, livers were resuspended in Buffer A (50 mmol/L HEPES, pH 7.4, 10 mmol/L KCl, 1 mmol/L ethylenediaminetetraacetic acid (EDTA), 1 mmol/L EGTA, 0.5% Nonidet P-40) and kept on ice for at least 10 minutes. The samples were cleared by centrifugation at 4°C at 4000  $\times$  *g* for 5 minutes. Pellets were resuspended in 200–500  $\mu$ l of buffer B (1.7 mol/L sucrose, 50 mmol/L HEPES, pH 7.4, 10 mmol/L KCl, 1 mmol/L EDTA, 1 mmol/L EGTA) and centrifuged at  $4^{\circ}$ C at 16,000  $\times$  *g* for 30 minutes. Pellets were resuspended in buffer C (10% glycerol, 50 mmol/L HEPES, pH 7.4, 400 mmol/L KCl, 1 mmol/L EDTA, 1 mmol/L EGTA), incubated on ice for 30 minutes with frequent vortexing, and centrifuged at 4°C at 16,000  $\times$  *g* for 5 minutes. Extracts were stored at  $-80^{\circ}$ C.

Nuclear extracts from newborn livers were isolated as previously described.3

## *Western Blot*

Protein was quantitated by the Bradford method (Bio-Rad, Hercules, CA) and 25  $\mu$ g of protein per lane was resolved on a 10% sodium dodecyl sulfate-polyacrylamide gel. The gels were transferred to nitrocellulose membranes (Amersham, Piscataway, NJ) and blocked overnight at 4°C in Tris-buffered saline containing 0.1% Tween-20 and 5% nonfat milk. Antibodies were obtained from Santa Cruz Biotechnologies, Inc. (Santa Cruz, CA). RelA (#SC-372-G),  $kB\alpha$  (#SC-371), and  $I_{\kappa}B\beta$  (#SC-945) were used at 1:2000 dilutions and TRAF-2 (#SC-876) was used at 1:1000 dilution. The membranes were incubated for one hour at room temperature, washed with Tris buffered saline-T, and bound with the appropriate horseradish peroxidaseconjugated secondary antibody; blots were washed and, subsequently, detected by ECL (Amersham).

# *Electrophoretic Mobility Shift Assay (EMSA)*

Four  $\mu$ g of nuclear protein were preincubated at room temperature for 10 minutes in a binding buffer (20 mmol/L HEPES, pH 7.5, 60 mmol/L KCI, 5 mmol/L  $MgCl<sub>2</sub>$ , 0.2 mmol/L EDTA, 8% glycerol, 1  $\mu$ g poly dl/dC, 1% Nonidet P-40, 0.1 mmol/L dithiothreitol, 0.1 mmol/L phenylmethyl sulfonyl fluoride). A  $[^{32}P]$ -labeled probe (1  $\times$  10<sup>5</sup> cpm) was added per reaction, and incubation was continued for an additional 30 minutes. Double-stranded probes were obtained from Santa Cruz Biotechnologies, Inc. The  $NF-\kappa B$  binding sequence probe used is 5' AGTT-GAGGGGACTTTCCCAGGC 3'. The SIE binding sequence probe used for Stat3 is 5' GTGCATTTCCCGTA-AATCTTGTCTACA 3'. Reactions were electrophoresed through a 5% polyacrylamide,  $1 \times$  Tris-glycine-EDTA gel.

For binding specificity, an unlabeled probe was added to one reaction 30 minutes before the addition of the labeled probe. For supershifts, after the 30-minute incubation of the labeled probe, specific antibodies obtained from Santa Cruz Biotechnologies, Inc. (RelA #SC-372X, p50 #SC-114X, p52 #SC-298X, c-Rel #SC-70X, and Stat3 #SC-482X) were added to indicated samples for an additional 30 minutes. The gels were dried and exposed to film for the detection of bands.

## *Immunohistochemistry*

Embryos isolated at each gestational stage were fixed overnight in 10% buffered-formalin (Fisher Scientific, Pittsburgh, PA) and processed for paraffin embedding. The sections were cut onto glass slides and processed for immunohistochemistry by using the VECTASTAIN ABC kit and protocol (Vector Laboratories, Burlingame, CA). Antibodies were used at 1:100 dilution, and the control reactions were performed with an antibody prebound to a control peptide (Santa Cruz) before incubation on the slide (data not shown). The ReIA,  $I_{\kappa}B_{\alpha}$  and TRAF-2 antibodies were the same as those used in Western blots. TNF- #SC-1348 and TNFR-1 #SC-1069 were also used. Biotinylated secondary antibodies and VEC-TASTAIN ABC solution were used at 1:100 dilutions. Antigen detection was through diaminobenzidine tetrahydrochloride (Sigma, St. Louis, MO) staining. The slides were counterstained with hematoxylin (Sigma).

# *Histology*

The samples from newborn pups were fixed overnight in 10% buffered formalin (Fischer Scientific) and processed for paraffin embedding. The sections were cut onto glass slides and processed for routine hematoxylin and eosin staining.

# **Results**

# *Expression of RelA Protein in WT Fetal Livers Does Not Correlate with Timing of Embryonic* Lethality in RelA<sup>-/-</sup> Mice

 $Re/A^{-/-}$  mice die within a specific time period during development (E14–E15). Therefore, we first investigated in WT (C57BL/6J) mice whether the appearance of RelA during liver development would coincide with the time of lethality observed in *RelA* knockouts. Whole cell protein samples were obtained from fetal livers of WT mice between E12 and E19. The top panel of Figure 1 demonstrates that RelA protein was present throughout liver development, from at least E12. These livers also contained the p50 subunit of  $NF-\kappa B$  (data not shown), indicating that the components of the classic  $NF-\kappa B$  heterodimer were present from a very early stage of liver formation.

In many adult cell types, including hepatocytes, NF-kB is retained in an inactive complex with an inhibitory  $I_{\kappa}B$ 



**Figure 1.** Protein expression in WT fetal liver development. Western blots using antibodies against RelA,  $I\kappa B\alpha$ ,  $I\kappa B\beta$ , and TRAF-2 illustrate protein expression throughout liver development. **Lanes 12–19:** fetal liver protein from gestational days 12–19, respectively.

protein.<sup>17,32</sup> Examination of  $I_{\kappa}\text{B}\alpha$  and  $I_{\kappa}\text{B}\beta$  proteins in WT fetal livers demonstrated expression from E12 through E19 (Figure 1, middle panels). Because RelA, p50,  $I_{\kappa}B_{\alpha}$ , and  $I_{\kappa}B\beta$  were all expressed in the fetal liver from the earliest time points analyzed, these data demonstrate that embryonic lethality in  $Re/A^{-/-}$  mice (E14–E15) does not correspond to a critical time when expression of these proteins becomes necessary in the fetal liver.

Proteins have been identified in the TNF-signaling pathway that are important for the activation of NF- $\kappa$ B. TRAF-2 is a TNFR-associated factor downstream of TNFR-1 that is important for  $NF-\kappa B$  activation after TNF stimulation.<sup>15</sup> TRAF-2 is recruited to the TNFR-1 signaling pathway by the adapter molecule TRADD.<sup>14</sup> Both TRAF-2 (Figure 1, bottom panel) and TRADD (data not shown) were expressed in the E12–E19 embryonic liver.

# *I*k*B*<sup>a</sup> *and I*k*B*b *Protein Levels Are Altered during Fetal Liver Development of* RelA *Knockouts but Not* TNFR-1 *Knockouts*

We next examined whether the expression of ReIA,  $I_{\kappa}B_{\alpha}$ ,  $I_{\kappa}B_{\beta}$ , TRAF-2, and TRADD was altered in fetal livers of *RelA* or *TNFR-1* knockouts. We compared the expression of these proteins to WT mice at E12–E14 (Figure 2). As expected, RelA was absent from *RelA* knockouts. In addition, the  $I_{\kappa}B_{\alpha}$  expression in the *ReIA<sup>-/-</sup>* fetal liver was decreased, and  $I_{\kappa}B\beta$  was undetectable. In contrast, levels of RelA,  $I_{\kappa}B_{\alpha}$ , and  $I_{\kappa}B_{\beta}$  in *TNFR-1* knockouts were similar to those observed in WT mice. It should be noted that, in these samples, the level of the  $I_{\kappa}B\beta$  expression is lower at E12 when compared with E13 and E14. These results suggest that, within the developing liver,  $NF-\kappa B$ participates in the regulation of  $I_{\kappa}B_{\alpha}$ . On the other hand, there appears to be a strict dependence of the  $I_{\kappa}B_{\beta}$ protein on NF-kB because this protein was not detectable in *RelA* knockouts. All of these proteins were normally expressed in  $TNFR-1^{-/-}$  fetal livers, indicating that TNFR-1 does not mediate the signal for the activation of



**Figure 2.** Protein expression in the fetal livers of WT mice, *RelA* knockouts, and *TNFR-1* knockouts. Western blots were performed with RelA, IKBa,  $I \kappa B \beta$ , and TRAF-2 antibodies. Lanes indicate fetal liver protein from E12, E13 and E14 for WT fetal livers, RelA<sup>-/-</sup> fetal livers and  $\overline{TNFR-1}^{-/-}$  fetal livers. *RelA*<sup> $-/-$ </sup> fetal livers illustrate absent RelA and I<sub>K</sub>B<sub>β</sub> proteins, diminished I<sub>K</sub>B<sub>α</sub> protein, and unaltered TRAF-2 protein levels. WT and *TNFR-1<sup>-/-</sup>* fetal livers express all proteins.

these genes. The expression of TRAF-2 (Figure 2, bottom panel) and TRADD (data not shown) was similar in WT mice and in *RelA* and *TNFR-1* knockouts during these days of gestation. The evidence of TRADD and TRAF-2 proteins in the  $Re/A^{-/-}$  fetal liver also illustrates that not all proteins are aberrantly expressed in the developing liver of *RelA* knockouts.

# *WT and TNFR-1<sup>-/-</sup> Fetal Livers Have Active NF-*k*B and STAT3 during Development, whereas* ReIA<sup>-/-</sup> Fetal Livers Have Only Active *STAT3*

To determine whether the  $NF-\kappa B$  proteins detected by Western blot were capable of binding DNA, indicative of an active protein complex, we performed EMSA using nuclear protein isolated from WT mice, *TNFR-1* knockout, and *RelA* knockout fetal livers between 12 and 14 days gestation. The  $NF-\kappa B$  complex formation occurred in WT (Figure 3A) and, surprisingly, also in  $TNFR-1^{-/-}$  (Figure 3B) fetal livers. In these samples, as in adult liver, NF- $\kappa$ B was composed of RelA/p50 heterodimers (top band) and p50 homodimers (bottom band). Nuclear protein isolated from E12–E14 ReIA<sup>-/-</sup> fetal livers only demonstrated p50 homodimer formation (Figure 3C). This suggests that the lack of *RelA* is not compensated for by the formation of other NF-kB heterodimers. RelA and p50 specificity was determined by supershift analyses (shown in Figure 3, B and C). These figures seem to indicate that, in  $TNFR-1^{-/-}$ fetal livers, the amount of  $NF$ - $\kappa$ B binding decreases with age, whereas in WT samples the inverse is seen. However, this was not a consistent finding when multiple fetal liver samples were analyzed at these gestational ages. It is interesting that, during E12–E14 of fetal liver development NF- $\kappa$ B complexes are forming, because in the adult liver there is little to no basal NF-kB binding. Antibodies against other Rel proteins, such as c-Rel and p52 (Figure 3B), did not cause band shifts in any samples, indicating that RelA/p50 heterodimers and p50 homodimers were



**Figure 3.** NF-kB complex formation in the fetal liver. **A:** Nuclear extract from WT fetal liver at E12, E13, and E14, demonstrating both RelA/p50 heterodimers (**top band**) and p50 homodimers (**bottom band**). **B:** Nuclear extract from *TNFR-1*2*/*<sup>2</sup> fetal livers at E12, E13, and E14, demonstrating both RelA/p50 heterodimers (**top band**) and p50 homodimers (**bottom band**). Supershift analysis was performed with E13 nuclear protein and anti-RelA antibody, anti-p50 antibody, anti-c-Rel antibody, anti-p52 antibody, or an irrelevant anti-STAT3 antibody, as indicated. Specificity of binding was determined by competition with 30-fold excess of<br>unlabeled oligonucleotide binding sequence probe. C: Nuclear extr Supershift analysis was performed with E13 nuclear extracts and anti-RelA antibody, which did not cause a supershift, and with anti-p50 antibody, which shifted the entire band, as indicated.

the active forms of NF- $\kappa$ B during fetal liver development. In the  $ReIA^{-/-}$  samples, the entire band was shifted with anti-p50 antibody, indicating that other Rel proteins were not being used in the absence of RelA. We conclude from these data that the embryonic lethality in  $ReIA^{-/-}$  mice does not coincide with the timing of  $NF- $\kappa$ B$  activation in liver development. The *RelA/p50* double knockouts<sup>33</sup> have a similar phenotype to that of *RelA* knockouts, but die earlier in development (E12). Thus, the presence of p50 homodimers might provide some protection against early lethality. The finding that *TNFR-1* knockouts have active NF-<sub>K</sub>B during liver development is unexpected, as these mice fail to activate  $NF-\kappa B$  in the regenerating liver.<sup>3</sup>

The  $STAT3^{-/-}$  mice are embryonic lethal early in gestation (E6.5–E7.5), therefore, the role it may have in the liver or other organ development is not yet defined.<sup>34</sup> However, STAT3 activation is an important component in murine hepatocyte proliferation during liver regeneration as demonstrated in *TNFR-1* and *IL-6* knockouts.<sup>3,25</sup> These and other experiments suggest that STAT3 activation in the liver may depend on  $NF$ - $\kappa$ B through induction of IL-6. STAT3 EMSA, and supershift analyses were performed on WT, *RelA* knockout, and *TNFR-1* knockout fetal livers from E12 through E14. Figure 4, A–C, indicates that WT, *TNFR-1<sup>-/-</sup>*, and *ReIA<sup>-/-</sup>* fetal livers were capable of specifically binding STAT3. Band specificity was determined by a complete supershift with an anti-STAT3 antibody; no shift was obtained with antibodies against STAT1 or STAT5 (data not shown). Because STAT3 was active in *ReIA<sup>-/-</sup>* mice during liver development, it appears that a functional NF-kB complex is not required for STAT3 activation in the embryonic liver. The *TNFR-1* knockouts do not show a STAT3 deficit during liver development, in contrast to the findings in liver regeneration.<sup>3</sup>



Figure 4. STAT3 binding in the fetal liver. Nuclear extracts were isolated from WT fetal livers  $(A)$ , *TNFR-1<sup>-/-</sup>* fetal livers  $(B)$ , and  $RelA^{-/-}$  fetal livers (**C**) at E12, E13, and E14, demonstrating STAT3 binding at all time points in all animals examined.

	Antibodies				
Cell types	$ReIA*$	$1\kappa B\alpha^{\dagger}$	<b>TNF</b>	TNFR- $1^{\ddagger}$	TRAF-2
Hepatoblasts Hematopoietic cells Megakaryocytes Endothelial cells	M M М М	M H н ⊢	Μ	Μ ⊢ M M	

Table 1. Cell Type Expression of TNF and NF-<sub>KB</sub> Proteins during Fetal Liver Development

E13 WT and *RelA<sup>-/-</sup>* embryos and E14 *TNFR-1<sup>-/-</sup>* embryos were fixed in formalin, processed for parafin embedding, and used for immunostaining as described in Materials and Methods. The levels of expression indicate those seen for all samples, except where indicated. H, high levels of expression (almost all cells express); M, moderate levels of expression (many cells express but many do not); L, low levels of expression (few cells express).

\*No staining in RelA knockouts.

† Decreased staining in RelA knockouts.

‡ Some nuclear staining in TNFR-1 knockouts.

# *Cell Type Expression of TNF and NF-*k*B Proteins Is Not Altered during Fetal Liver Development*

The fetal liver contains diverse cell types, with hematopoietic cells, hepatoblasts, and endothelial cells composing the predominant types in mid-gestation. Whereas the expression of proteins and transcription factor complex assembly can be analyzed by Western blot and EMSA, respectively, they do not provide information regarding the localization of these proteins. To investigate this issue, we performed immunostaining for ReIA,  $I_{\kappa}B_{\alpha}$ , TNFR-1, TRAF-2, and TNF- $\alpha$  in the E13 fetal liver of WT mice and *RelA* knockouts and the E14 *TNFR-1* knockout fetal liver. At E13,  $ReIA^{-/-}$  mice have normal liver morphology similar to that of WT embryos. We found that the staining patterns observed in the fetal livers of WT mice, *RelA* knockouts, and *TNFR-1* knockouts were remarkably similar for all the proteins tested. The results are summarized in Table 1. In the WT mice and *TNFR-1* knockouts, RelA and  $I_{\kappa}B_{\alpha}$  proteins were detected mostly in the cytoplasm of endothelial cells, hepatoblasts, and hematopoietic cells. Cytoplasmic expression of  $I_{\kappa}B_{\alpha}$  was also detected in *RelA* knockout fetal livers. In all instances, we saw occasional cells exhibiting nuclear staining of  $I_{\kappa}B_{\alpha}$ . The  $Re/A^{-/-}$  samples did have a decreased intensity of  $I_{\kappa}B_{\alpha}$  staining, likely due to diminished protein levels observed by Western blot (Figure 2). Small hematopoietic cells were the primary sites of the TNFR-1 expression in the fetal livers of both the WT mice and *RelA* knockouts. Surprisingly, a minimal amount of nuclear staining with the TNFR-1 antibody was observed in the  $TNFR-1^{-/-}$ fetal livers, whereas the WT and  $Re/A^{-/-}$  samples only demonstrated cytoplasmic and membrane-localized staining. However, the  $TNFR-1^{-/-}$  mouse is the result of a functionally mutant TNFR-1 and not a completely absent gene; thus, we may be detecting a small amount of nonfunctional protein sequestered within the nucleus. In addition, the WT,  $ReIA^{-/-}$ , and  $TNFR-1^{-/-}$  mice all had similar TNF and TRAF-2 staining patterns. These data show that the liver development of the *ReIA<sup>-/-</sup>* and *TNFR-* $1^{-/-}$  mice is similar to that of the WT mice, at least through 13 days of gestation. After this time point, the *RelA<sup>-/-</sup>* mice undergo rapid liver degeneration and die, whereas the  $TNFR-1^{-/-}$  mice proceed through gestation in a manner similar to the WT mice. Furthermore, these findings indicate that TNF and TNFR-1 are expressed during gestation in the fetal liver before the time of embryonic lethality seen in the  $Re/A^{-/-}$  mice.

# *Signaling through* TNFR-1 *Is Responsible for Embryonic Lethality in the ReIA<sup>-/-</sup> Mice*

To determine whether signaling through TNFR-1 contributes to the embryonic lethal phenotype of *RelA<sup>-/-</sup>* fetuses, mice deficient in both *RelA* and *TNFR-1* were generated as described in the Materials and Methods section. A total of 115 pups were examined by PCR and monitored for survival. Only seven of nine theoretically possible genetic outcomes were obtained in live-born animals. Table 2 summarizes the results of these crosses. From 115 newborn pups analyzed, 7 (6.1%) had homozygous mutations in *RelA* and *TNFR-1*. Nine newborns (7.8%) had homozygous WT alleles for both genes, and 33 pups (28.7%) were double heterozygous. However, only four of the remaining six genetic possibilities were observed, because newborn mice carrying a null *RelA* genotype were never observed unless *TNFR-1* was also

**Table 2.** Genetic Analysis of Offspring Obtained from RelA/ TNFR-1 Double Heterozygous Matings

			Actual percentage
6.25	9	7.2	7.82
12.5	14	14.4	12.17
12.5	19	14.4	16.52
25.0	33	28.8	28.70
6.25	11	7.2	9.57
12.5	22	14.4	19.13
6.25	O	7.2	Ω
12.5	O	14.4	Ω
6.25	7	7.2	6.09
			Expected Actual Expected number number percentage

Genotyping was performed by PCR analysis of tail DNA using primers diagnostic for both WT and mutant alleles.

	Average weight (grams)					
Age (days)	$ReIA^{+/+}/TNFR-1^{+/+}$	$ReIA^{+/+}/TNFR-1^{-/-}$	$ReIA^{-/-}/TNFR-1^{-/-}$			
2	$1.4 \pm 0.04$	$1.4 \pm 0.1$	$1.2 \pm 0.1$			
	$(n = 3)$	$(n = 12)$	$(n = 2)$			
4	$2.3 \pm 0.1$ $(n = 3)$	$2.0 \pm 0.2$ $(n = 14)$	$2.0 \pm 0.1$ $(n = 3)$			
	$4.6 \pm 0.3$	$3.8 \pm 0.6$	$2.5 \pm 0.2^{*+}$			
	$(n = 3)$	$(n = 8)$	$(n = 2)$			
16	$9.6 \pm 0.6$	$8.4 \pm 0.2^*$	$5.3 \pm 1.7^{*+}$			
	$(n = 3)$	$(n = 5)$	$(n = 4)$			

**Table 3.** Average Weight Gain during Early Postnatal Life

Average weight of WT, *TNFR-1<sup>-/-</sup>*, and *RelA<sup>-/-</sup>/TNFR-1<sup>-/-</sup>* pups during the first 16 postnatal days. Mice were weighed at the indicated ages. Tail samples were taken for DNA isolation and PCR genotyping. Numbers represent average  $\pm$  SD.

\*Statistically different from WT mice:  $P \leq 0.001$  to 0.01.

<sup>†</sup>Statistically different from TNFR-1 knockout mice;  $P \le 0.001$  to 0.01.

absent. Taking these two lethal outcomes (ReIA<sup>-/-</sup>/ *TNFR-1<sup>+/+</sup>* and *ReIA<sup>-/-</sup>/TNFR<sup>+/-</sup>)* into consideration, all live-born mice, including *ReIA<sup>-/-</sup>/TNFR-1<sup>-/-</sup>*, were obtained with the expected Mendelian ratios. By monitoring the pups daily, we observed that at birth all animals appeared healthy and nursed well. However, within the first postnatal week, double-knockout mice became runted, failed to thrive despite continuing to nurse, and subsequently died. Our survival analysis on the first seven double-knockout pups revealed that the average survival time was 10 days with a range of 6 to 17 days.

For subsequent studies,  $ReIA^{+/-}/TNFR-1^{-/-}$  mice were used as breeders to obtain litters with a 25% chance of generating *ReIA<sup>-/-</sup>/TNFR-1<sup>-/-</sup>* pups. *ReIA<sup>+/+</sup>/ TNFR-1<sup>+/+</sup>* mice were used to generate age-matched WT controls. At various days after birth, pups were killed and analyzed for body weight, necropsies were performed for histology sections, and pieces of liver were harvested for whole-cell extract and RNA. Consistent with our earlier observations, all newborn pups were similar in weight at 2 and 4 days of age (Table 3). The WT pups and *TNFR-* $1^{-/-}$  pups continued to gain weight, and by 7 days of age, they had doubled their 4-day body weight. In contrast, 7-day-old *ReIA<sup>-/-</sup>/TNFR-1<sup>-/-</sup>* pups had gained only a minimal amount of weight and now appeared significantly smaller, resembling 4-day-old mice. This difference in size is statistically different from both the WT and *TNFR-1* knockout mice at the same age. This size difference becomes even more pronounced if the *ReIA<sup>-/-</sup>/ TNFR-1<sup>-/-</sup>* pups survive to 16 days of age. However, we found that, by 16 days of age, the *TNFR-1* knockout pups were also statistically different in weight from both the WT pups and the double-knockout pups at that age. Therefore, the lack of *TNFR-1* alone may influence body weight to some degree. Thus, the lack of *TNFR-1* rescued *RelA<sup>-/-</sup>* mice from embryonic lethality, demonstrating that signaling through TNFR-1 is responsible for the embryonic death observed in these animals. However, whereas *RelA* may not be necessary for fetal development, its absence, in combination with the lack of TNFR-1 signaling, causes death within a very short time after birth.

All *RelA<sup>-/-</sup>* mice died during E14–E15 with massive hepatocyte apoptosis as previously described.<sup>4,5</sup> The *RelA<sup>-/-</sup>/TNFR-1<sup>-/-</sup>* mice survived gestation and were

born with normal liver morphology. Liver histology was similar between 2-day-old WT mice and double knockouts (Figure 5, A and B). The livers of the *RelA<sup>-/-</sup>/TNFR-* $1^{-/-}$  mice had normal architecture with a persistence of small foci of hematopoietic cells and little, if any, hepatocyte apoptosis. Between 2 and 7 days of age, foci of hematopoietic cells greatly decreased in the WT mice, but at 7 days the livers of the *RelA/TNFR-1* double knockouts had more abundant and larger hematopoietic foci than the WT mice. Furthermore, within the first 2–3 postnatal weeks, the condition of the *RelA/TNFR-1* double knockouts deteriorated, and, by the time of death, acute hepatitis with massive neutrophilic infiltration and necrotic foci was present. The accumulation of immature neutrophils (band cells) started in the periportal areas and progressively infiltrated the liver parenchyma (Figure 5, C–E). Although scattered apoptotic hepatocytes were present, liver damage and, presumably, the lethality of the *RelA<sup>-/-</sup>/TNFR-1<sup>-/-</sup>* mice were consequences of neutrophilic infiltration and hepatocyte necrosis. The livers of the *TNFR-1* knockouts were morphologically normal and were not distinguishable from those of the WT mice. In addition to severe liver abnormalities, neutrophilic infiltration, although not as extensive as that in the liver, was present in the heart (data not shown) and lungs of some *RelA<sup>-/-</sup>/TNFR-1<sup>-/-</sup>* mice. An example of an unusual lung lesion resembling lobar pneumonia that developed in one animal is shown in Figure 5F, in which normal alveoli are seen adjacent to the walled lesion. Spleens of the *RelA<sup>-/-</sup>/*  $TNFR-1^{-/-}$  mice showed a decrease in white pulp areas and less compact germinal centers (compare Figure 5, G and H), whereas in the thymus there were foci of cell degeneration (data not shown). It has been reported that the germinal-center formation is abnormal in *TNFR-1* knockout mice. These changes appear to be accentuated in the  $Re/A^{-/-}/TNFR-1^{-/-}$  mice.

The embryonic survival and subsequent postnatal mechanism of death observed in the *RelA/TNFR-1* knockouts suggest that the absence of *RelA* by itself is not responsible for fetal liver death, but that *RelA<sup>-/-</sup>* cells become sensitive to TNFR-1-mediated apoptotic signals. Removal of these signals in the double-knockout mice prevented embryonic lethality and did not interfere with liver development.

# *I*k*B*<sup>a</sup> *and I*k*B*b *Proteins Are Altered in the Livers of ReIA<sup>-/-</sup>/TNFR-1<sup>-/-</sup> Mice*

To determine whether rescuing the  $Re/A^{-/-}$  lethal phenotype in the *RelA/TNFR-1* knockouts would correct the defects in the  $I_{\kappa}B_{\alpha}$  and  $I_{\kappa}B_{\beta}$  proteins, Western blot analysis was performed. Whole-cell proteins were isolated from the livers of the WT, *TNFR-1<sup>-/-</sup>*, and *ReIA<sup>-/-</sup>/TNFR-* $1^{-/-}$  mice at 2, 4, and 7 days of age, respectively (Figure 6). Similar to the *RelA* knockouts at E12–E14, *RelA/ TNFR-1* double knockouts had decreased  $I_{\kappa}B_{\alpha}$  and no detectable  $I_{\kappa}B\beta$  in comparison with the WT mice and the *TNFR-1* knockouts. This suggests that a deficiency of these proteins is not causing the embryonic lethal phenotype of the *RelA* knockouts. Nevertheless, it could still be contributing to postnatal mortality. As expected, RelA expression was absent in the *RelA/TNFR-1* knockouts, whereas TRAF-2 was unaffected in all of the mice.

# RelA<sup>-/-</sup>/TNFR-1<sup>-/-</sup> Mice Do Not Have an *Increase in TNFR-2 Messenger RNA or Incidence of Apoptosis during the First Postnatal Week*

Up-regulation of TNFR-2 has been shown to result in an inflammatory response.<sup>35</sup> Therefore, we examined whether a TNFR-2 message was being up-regulated in the liver of the *TNFR-1<sup>-/-</sup>* or *ReIA<sup>-/-</sup>/TNFR-1<sup>-/-</sup> pups* during the first week of life. We found that neither animal had up-regulated TNFR-2 messenger RNA compared with the WT mice (data not shown). We also examined whether the double knockouts had increased liver apoptosis that was contributing to their demise. Multiorgan slides of 2-, 4-, 7-, and 16-day-old WT,  $TNFR-1^{-/-}$ , and *RelA<sup>-/-</sup>/TNFR-1<sup>-/-</sup>* pups were examined by terminal deoxynucleotidyltransferase-mediated uridine 5'-triphosphate end labeling assay (data not shown). We did not find an increase in incidence of apoptosis at any age, in any organ, including the liver. Therefore, TNF signaling through TNFR-2 does not result in TNFR-2 messenger RNA up-regulation or in an apoptotic phenotype in the newborn *ReIA<sup>-/-</sup>/TNFR-1<sup>-/-</sup>* pup.

# **Discussion**

The absence of *RelA* results in massive hepatocyte apoptosis and embryonic death between E14 and E15. The timing of this lethal effect may reflect the time at which NF- $\kappa$ B proteins are synthesized or assembled during liver

development. To examine this possibility, we analyzed the expression of ReIA, p50,  $I_{\kappa}B_{\alpha}$ , and  $I_{\kappa}B_{\beta}$  in WT mice from E12–E19. We also examined the formation of active NF-kB complexes in the livers of WT mice from E12–E14. These analyses revealed that all of the proteins and NF-kB complexes, consisting of RelA/p50 heterodimers and p50 homodimers, were present in the liver from at least E12. Because NF-kB complexes are formed in the developing liver of the WT mice at least 2 days before the onset of embryonic death in the  $Re/A^{-/-}$  mice, it is unlikely that lethality corresponds to the timing of  $NF-\kappa B$ activation during hepatic development. An investigation of fetal livers from *RelA* knockouts revealed that they do not form  $NF$ - $\kappa$ B heterodimers, but contain only p50 homodimers. Thus, it appears that, during liver development, a RelA deficiency is not compensated for by other Rel family members. However, *RelA<sup>-/-</sup>/p50<sup>-/-</sup>* embryos die with a similar phenotype to the RelA<sup>-/-</sup> mice, but 2 days earlier in development.<sup>33</sup> It is, therefore, possible that, in the *RelA* knockouts, *p50* partially compensates for the lack of *RelA* before E14.

In comparison with the WT mice,  $Re/A^{-/-}$  fetal livers from E12–E14 had lower levels of  $I_{\kappa}B_{\alpha}$  protein and no detectable  $I_{\kappa}B\beta$  protein. This finding is similar to data reported in  $Re/A^{-/-}$  embryonic fibroblasts.<sup>4</sup> Here we measured  $I_{\kappa}B_{\alpha}$  and  $I_{\kappa}B_{\beta}$  directly in the liver and illustrated that  $I_{\kappa}B\beta$  regulation is dependent on ReIA, whereas  $I_{\kappa}B_{\alpha}$  is partially, but not completely, dependent on RelA. Similar to the WT mice, the *TNFR-1<sup>-/-</sup>* mice had no alterations in ReIA, p50,  $kB\alpha$ , or  $kB\beta$  throughout liver development. TRADD and TRAF-2 proteins were not altered in any of the fetal livers examined.

TNF is a potent NF- $\kappa$ B activator in the adult liver, and signaling through TNFR-1 is required for the initiation of liver regeneration. Although TNF by itself is not generally apoptotic for many cell types,<sup>36</sup> it can cause apoptosis when given in conjunction with drugs that block transcription and translation.<sup>37</sup> Moreover, TNF can cause hepatocyte apoptosis *in vivo* and *in vitro* when NF-kB activation is blocked by an  $I_{\kappa}B_{\alpha}$  super-repressor.<sup>38,39</sup> Because the *TNFR-1* knockouts develop normally, we hypothesized that the elimination of TNFR-1-mediated apoptotic signals could rescue the *RelA* knockouts from embryonic lethality. Therefore, we generated mice that were deficient in both *RelA* and *TNFR-1* and showed that these animals survive embryonic development. At birth *RelA<sup>-/-</sup>/TNFR-* $1^{-/-}$  pups have normal liver morphology, but, after  $4-5$ days of age, most animals become runted and weigh significantly less than the WT mice of the same age. The double knockouts develop acute hepatitis and die with an

**Figure 5.** Histological analysis of WT and *RelA<sup>-/-</sup>/TNFR-1<sup>-/-</sup>* mice. All sections were stained with H&E. **A:** Liver section from a 2-day-old WT mouse showing normal hepatocytes and presence of focal areas of hematopoiesis. Original magnification, ×200. **B:** Liver section from a 2-day-old *RelA<sup>-/-</sup>/TNFR-1*<sup>-/-</sup> mouse showing normal hepatocytes and presence of focal areas of hematopoiesis. Original magnification, ×200. **C:** Liver section from an 8-day-old *RelA<sup>-/-</sup>/TNFR-1<sup>-/</sup>* mouse illustrating the presence of massive neutrophilic infiltration. Original magnification, ×100. **D:** Higher magnification of the 8-day *RelA<sup>-/-</sup>/TNFR-1<sup>-/-</sup>* liver section, illustrating the immature neutrophils (band cells) responsible for the inflammation seen. Original magnification,  $\times$  400. **E:** Liver section from a 17-day-old *ReLA<sup>-/-</sup>/INFR-1<sup>-/-</sup>* mouse, demonstrating focal <sup>-</sup> mouse, demonstrating focal areas of necrosis. Original magnification,  $\times$ 400. **F:** Lung lesion in a *RelA<sup>-/-</sup>/TNFR-1<sup>-/-</sup>* mouse killed at 16 days of age. The lesion was a solid mass that histologically resembled lobular pneumonia. The area was separated from the rest of the lung parenchyma (**left side**, showing alveoli) by connective tissue surrounded by inflammatory cells. The lesion itself (**right side**) contained closed and thickened alveoli, inflammatory cells, and cell necrosis. Original magnification, 3200. **G:** Spleen section from a 16-day-old WT mouse, showing normal germinal center formation. Original magnification, ×100. **H:** Spleen section from a 16-day-old *RelA<sup>-/-</sup>/TNFR-1<sup>-/-</sup>* mouse, demonstrating less defined white pulp and germinal follicle areas as well an increased numbers of megakaryocytes. Original magnification,  $\times$ 200.





**Figure 6.** Protein expression in livers of WT, *TNFR-1<sup>-/-</sup>*, and *RelA<sup>-/-</sup>/TNFR*newborn pups. Western blots for RelA,  $I \kappa B \alpha$ ,  $I \kappa B \beta$ , and TRAF-2 were performed on liver extracts from 2-, 4-, and 7-day-old WT mice, *TNFR-1<sup>-</sup>* mice, and  $RelA^{-/-}/TNFR-1^{-/-}$  mice as indicated. WT and  $TNFR-1^{-/-}$  mice express all proteins.  $RelA^{-/-}/TNFR-1^{-/-}$  mice have absent RelA and  $I\kappa B\beta$ proteins, diminished IkB<sup>a</sup> protein expression, and unaltered TRAF-2 protein levels.

average survival of 10 days. These data suggest that *RelA* is not critical for liver development but may serve a vital protective function against TNFR-1 apoptotic signaling. Additionally, the *RelA<sup>-/-</sup>/TNFR-1<sup>-/-</sup>* mice may have limited survival because *RelA* provides a critical postnatal function. However, the *ReIA<sup>-/-</sup>/TNFR-1<sup>-/-</sup>* pups do not die with massive liver apoptosis as seen in the *RelA* knockouts. Instead, these animals develop acute hepatitis and neutrophilic infiltration leading to focal necrosis of the liver. Given the intense inflammatory infiltrate found in the livers of the dying mice, it is likely that the lack of TNF signaling and RelA makes these animals extremely susceptible to endogenous infectious agents. However, we cannot rule out the possibility that this phenotype is entirely caused by the absence of *RelA* in the newborn. A conditional *RelA* knockout will be necessary to examine which affects are due to the *RelA* deficiency and which are due to the combined lack of *TNFR-1* and *RelA*. All animals, including the double knockouts, were maintained in a specific pathogen-free facility, reducing the possibility of contamination by external organisms, and were free of murine hepatitis virus. The phenotype of *RelA/TNFR-1* knockouts has similarities to phenotypic abnormalities described in the *RelB* knockouts.<sup>8</sup> These animals also had neutrophilic inflammatory infiltrates and extramedullary hemopoiesis in the liver. It is puzzling that the *RelA* knockouts die only with liver abnormalities because RelA is ubiquitously expressed during normal development. Perhaps RelB or other Rel proteins are compensating for the loss of RelA in hematopoietic organs, although we found no evidence of this within the liver.

While this manuscript was being prepared, Doi et al reported that *RelA* knockouts can be rescued from embryonic lethality by crossing them with  $TNF^{-/-}$  mice.<sup>40</sup> Similar to our findings, the *RelA/TNF* knockouts have normal liver morphology at birth; however, death does not occur until approximately 40 days of age. In *RelA/TNFR-1* knockouts, we have not observed survival beyond 3 weeks of age. Histological analysis of 40-day-old *RelA<sup>-/-</sup>/*  $TNF^{-/-}$  mice demonstrated lung and liver inflammation (similar to our findings in *RelA/TNFR-1* knockouts), as well as kidney abnormalities that we did not observe. A lack of TNFR-1 signaling caused a more profound defect than TNF deficiency, and this might be due to TNF signaling via TNFR-2. However, signaling through TNFR-2 did not cause increased apoptosis in the double-knockout newborns, but it could be contributing to the intense inflammatory reaction. Preliminary evidence suggests that the *TNFR-2<sup>-/-</sup>* mice cannot rescue the  $Re/A^{-/-}$  lethal phenotype, indicating that, whereas a functional TNFR-2 may be contributing to the demise of *ReIA<sup>-/-</sup>/TNFR-1<sup>-/-</sup>* mice, it is not playing a role in  $Re/A^{-/-}$  lethality. It has also been recently reported that *TNFR-1<sup>-/-</sup>* mice can rescue the embryonic lethal effects seen in  $IKK-2^{-/-}$  mice that die from a similar phenotype to that of *RelA<sup>-/-</sup>* mice.<sup>22</sup> However, details of these *TNFR-1<sup>-/-</sup>/IKK-2<sup>-/-</sup>* mice were not described.

Although the main goal of this work was to examine embryonic lethality of RelA<sup>-/-</sup> mice, we also obtained interesting information about *TNFR-1* knockouts. These animals develop normally, but show diminished resistance to infection by intracellular bacteria.41 The *TNFR-1* knockouts fail to activate NF- $\kappa$ B and STAT3 during liver regeneration, leading to high mortality and deficient DNA replication.<sup>3</sup> A major puzzle is to explain how these animals survive embryonic development, if we assume that, as in liver regeneration, the *TNFR-1* knockouts have deficient  $NF$ - $\kappa$ B and STAT3 during liver development. Surprisingly, we found that both  $NF-\kappa B$  and STAT3 were active in the liver of the *TNFR-1* knockouts. This suggests that either TNF does not activate  $NF - \kappa B$  during normal liver development or that, in the *TNFR-1* knockouts, an alternate mechanism of  $NF- $\kappa$ B$  activation (perhaps through IL-1) is used. In any event, these results point out that hepatocyte proliferation in liver development and liver regeneration has different regulatory pathways. Thus, the absence of *RelA* is not directly responsible for fetal liver death, but its absence allows cells to become sensitive to TNFR-1-mediated apoptotic signals. The removal of these signals in *ReIA<sup>-/-</sup>/TNFR-1<sup>-/-</sup>* mice prevents embryonic lethality and does not interfere with liver development.

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