

ARTICLES

Flice Inhibitory Protein Is Associated With the Survival of Neonatal Neutrophils

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ABSTRACT: Neonatal polymorphonuclear leukocytes (PMN) exhibit delayed apoptosis both constitutively and under inflammatory conditions, and evidence has linked PMN longevity to the presence of antiapoptotic proteins. Activation of the survival-associated transcription factor, nuclear factor kappa B (NF- κ B), promotes the synthesis of several antiapoptotic proteins including Flice inhibitory protein (FLIP). Neonatal and adult PMN were compared in this study to test the hypothesis that FLIP modulates age-related apoptosis. Expression of the short isoform, FLIP-S, was prominent at baseline and persisted during spontaneous apoptosis in neonatal PMN, whereas basal expression was lower and decreased under the same conditions in adult PMN. Stable FLIP-S expression in neonatal PMN was associated with a relative resistance to apoptosis in response to the protein synthesis inhibitor, cycloheximide (CHX), or the NF- κ B inhibitor, gliotoxin. In contrast, similar treatment of adult PMN promoted greater overall apoptosis accompanied by FLIP degradation. Nuclear levels of phosphorylated p65, a critical NF- κ B dimer, were relatively robust in neonatal PMN under basal conditions or after stimulation with TNF- α , a cytokine that induces FLIP. In conclusion, persistent FLIP-S expression is involved in the longevity of neonatal PMN, and our data suggest a contribution of NF- κ B signaling and related survival mechanisms. (*Pediatr Res* 70: 327–331, 2011)

The timely resolution of inflammatory processes is dependent on the efficient clearance of apoptotic neutrophils [polymorphonuclear leukocytes (PMN)] (1), while prolonged PMN survival can contribute to chronic inflammation (2). Neonatal PMN are relatively resistant to both spontaneous and Fas-mediated apoptosis (3,4). In addition, neonatal PMN exhibit marked survival responses to cytokines (3,5) identified with bronchopulmonary dysplasia and other neonatal inflammatory disorders (6,7). Neonatal PMN exhibit diminished functional expression of caspase-3, a critical effector of apoptosis, as well as other key apoptotic proteins (4,8). In addition, neonatal PMN have an impaired ability to alter mem-

brane potential in response to stimulation (9), also observed to contribute to delayed PMN apoptosis during sepsis (10). Neonatal PMN also exhibit a diminished apoptotic response to treatment with the protein synthesis inhibitor, cycloheximide (CHX) (3), consistent with the presence of preformed survival proteins (11). Thus, neonatal PMN are characterized by altered function of death mechanisms that favor survival under basal and inflammatory conditions, although these remain incompletely defined.

Flice inhibitory protein (FLIP) is a prototypical antiapoptotic protein critical to the survival of hematopoietic progenitors, although its role in PMN is less clear (12,13). Synthesis of FLIP and related antiapoptotic proteins is regulated in part by the transcription factor, nuclear factor kappa B (NF- κ B) (14–17). Modulation of the NF- κ B pathway by inflammatory cytokines such as TNF- α (11,18–20) can induce FLIP (14), including the short isoform of FLIP (FLIP-S) in PMN (13). In addition, Vancurova *et al.* (21) observed enhanced TNF- α -mediated activation of NF- κ B in neonatal PMN. However, the contribution of FLIP expression to the prolonged survival of neonatal PMN has not been described. We designed this study to test the hypothesis that increased FLIP expression in neonatal PMN is a mechanism underlying their relative longevity.

METHODS

Donor characteristics. Heparinized blood samples from the peripheral veins of healthy adult donors (aged 18–55 y) and from the umbilical venous cord blood of freshly delivered term placentas were processed in parallel. Cord blood donors met strict criteria including term gestation, planned C-section, normal intrauterine growth, and the documented absence of acute, chronic or gestational illness or infection, medications, tobacco and/or recreational drug use, and fetal genetic/structural abnormalities. Samples were not used in the event of low (<5) neonatal Apgar scores. All blood samples were collected from donors following informed consent according to a protocol approved by the Institutional Review Board for Human Studies at St. Louis University.

Abbreviations: AD, adult; CB, cord blood; CHX, cycloheximide; FLIP, Flice inhibitory protein; FLIP-S, short isoform of FLIP; FLIP-L, long isoform of FLIP; I κ B α , inhibitor kappa B alpha; IKK, I κ B kinase; NF- κ B, nuclear factor kappa B; PCNA, proliferation cellular nuclear antigen; PMN, polymorphonuclear leukocytes; XIAP, X-linked inhibitor of apoptosis

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PMN isolation and culture. PMN were isolated as described (5). Isolated PMN suspended in RPMI 1640/2% FCS were incubated at 37°C, 5% CO₂ for up to 24 h in the presence or absence of specific stimuli or inhibitors.

Reagents. RPMI 1640 and FCS were purchased from GIBCO-BRL (Invitrogen Corporation, Carlsbad, CA) and Hyclone, Inc. (Logan, UT), respectively. Anti-FLIP and anti-XIAP (X-linked inhibitor of apoptosis) antibodies were purchased from Axxora (San Diego, CA), and antibodies against the nuclear marker PCNA (proliferation cellular nuclear antigen) and secondary antibodies were purchased from Santa Cruz Biotechnology, Inc. (Santa Cruz, CA). Antibodies against I κ B α (I kappa B alpha), IKK (I κ B kinase), and the NF- κ B subunit, p65, and their phosphorylated forms were purchased from Cell Signaling Technology (Beverly, MA). Recombinant human TNF- α was purchased from R&D Systems (Minneapolis, MN). Anti- β -actin antibodies, gliotoxin, and CHX were purchased from Sigma Chemical Co.-Aldrich, Inc. (St. Louis, MO).

Apoptosis studies. To correlate *de novo* FLIP synthesis with survival, PMN were cultured with the protein synthesis inhibitor CHX and treated or control cells were analyzed for apoptosis by TUNEL assay (3). To examine a potential survival effect of constitutive NF- κ B activation, PMN were treated with the NF- κ B inhibitor, gliotoxin (11,22), and apoptosis was assessed by TUNEL assay. In parallel studies, the apoptotic effects of each treatment were correlated with FLIP expression in PMN lysates, analyzed by Western blot.

TUNEL assay. A commercial flow cytometric TUNEL assay (*In Situ* Cell Death Detection Kit; Boehringer Mannheim, Inc., Mannheim, Germany) was used to detect and quantify apoptotic cell death by enzymatic labeling of DNA strand breaks with fluorescein (dUTP) and terminal deoxy-nucleotidyl transferase (TdT), as described (3).

Western blots. For whole cell studies, PMN (5×10^6) were lysed in RIPA buffer containing a protease inhibitor cocktail (both, Sigma Chemical Co.-Aldrich). Nuclear fractions were prepared from PMN (10^7) stimulated with TNF- α , using the NE-PER kit (Pierce, Rockford, IL). For phosphoprotein expression, the lysis buffer was supplemented with 2 mM each of Na orthovanadate and Na fluoride. Equivalent protein amounts of lysates in 2 \times Laemmli buffer were separated by SDS-PAGE and analyzed by Western blot using primary antibodies and horseradish peroxidase-conjugated secondary antibodies. Protein sample loading was normalized by reprobing blots with anti- β -actin antibodies or with a nuclear marker, as described (23,24). Protein bands were visualized by chemiluminescence (ECL), and band intensities relative to β -actin or PCNA expression or the ratio of phosphorylated to total cognate protein were quantified by densitometric analysis.

Statistical analysis. Data expressed as mean \pm SD were analyzed by *t* test or by ANOVA, as appropriate, using a statistical software program (Sigma Chemical Co. Stat for Windows, SPSS, Inc.). A two-way ANOVA with a fixed factor effect was performed where appropriate to assess differences between neonatal (CB) and adult (AD) groups across time. All analyses used an α -level 0.05 to determine statistical significance.

RESULTS

FLIP expression during spontaneous apoptosis. Expression levels of FLIP were determined in lysates of PMN at baseline (0 h) and during spontaneous apoptosis (24 h culture) (3, 11). Basal expression of FLIP-S was greater than that of the long form (FLIP-L) and expression of each decreased relative to basal levels during spontaneous apoptosis in both neonatal and adult PMN (Fig. 1A). However, FLIP-S expression was more prominent in neonatal PMN under both basal and apoptotic conditions. Expression of another antiapoptotic protein, XIAP, was similar at 0 or 24 h (Fig. 1B), confirming previous observations (13), and expression did not differ between groups.

Differential responses of neonatal and adult PMN to CHX-induced apoptosis and FLIP degradation. Neutrophils were incubated with CHX to determine a link between increased FLIP expression and *de novo* protein synthesis. The higher dose of CHX tested (100 μ g/mL) induced a greater apoptotic response in adult *versus* neonatal PMN after incubation for 3 or 6 h, whereas a difference between groups was observed for the lower dose (50 μ g/mL) only after the longer incubation (Fig. 2A), confirming our previous observations

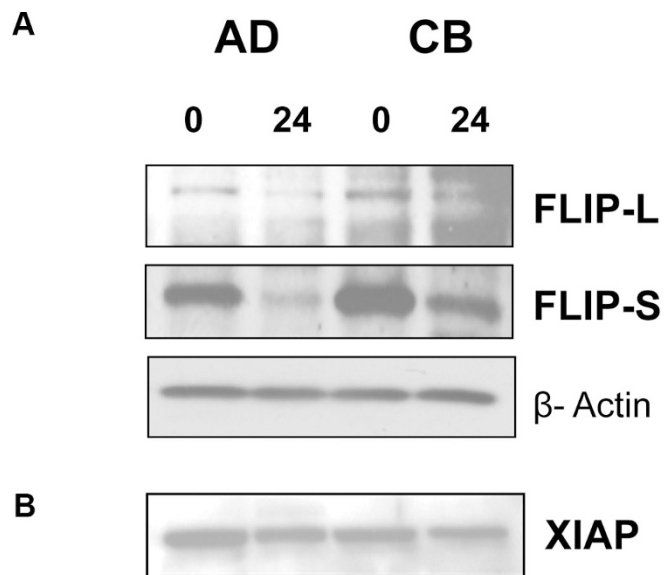


Figure 1. FLIP expression in neonatal and adult PMN. FLIP isoforms and β -actin proteins were detected by Western blot in lysates of adult (AD) and neonatal (CB) PMN (PMN) cultured for 0–24 h. Lysates of PMN harvested at each time point were prepared and blotted with monoclonal anti-FLIP or anti-XIAP antibodies. The blots shown are representative of five (A) or two (B) separate paired (AD, CB) experiments.

(3). Diminished apoptosis of neonatal PMN in response to CHX (3 h) (Fig. 2B) was accompanied by persistent expression of FLIP isoforms, with a more prominent expression of FLIP-S (Fig. 2C). In contrast, CHX treatment of adult PMN was associated with decreased expression of both FLIP isoforms (Fig. 2C) that paralleled the greater apoptotic responses of these cells (Fig. 2B). No differences were observed either in the proportion of apoptotic cells or in the expression of FLIP-S compared with respective baseline values in control PMN from either group.

TNF- α induced activation of the NF- κ B pathway in neonatal and adult PMN. In studies to compare NF- κ B activity in neonatal and adult PMN, nuclear expression levels of the cognate and phosphorylated forms of IKK, I κ B α , and p65 were determined under basal conditions and after TNF- α treatment (Fig. 3A and B). Transient phosphorylation of IKK peaked at 30 min in neonatal PMN and at 60 min in adult PMN. Prominent phosphorylation of nuclear I κ B α in neonatal PMN was observed at baseline and throughout the study interval and was associated with I κ B α degradation at 30 min. In contrast, in adult PMN, phosphorylated I κ B α levels were most evident at 15 and 30 min of treatment, and I κ B α expression levels were lowest at 60 min. The phosphorylation status of nuclear p65/RelA changed significantly from basal values over the treatment period in both neonatal and adult PMN ($p < 0.001$), whereas levels of the unphosphorylated protein remained similar (Fig. 3A and B). In neonatal PMN, phospho-p65 levels were robust under basal conditions and phosphorylation status decreased over the ensuing 30 min, followed again by prominent expression at 60 min. In contrast, the phosphorylation status of p65 was not as pronounced in adult PMN; phospho-p65 levels were minimal at baseline and levels peaked at 15 min of treatment, decreasing thereafter.

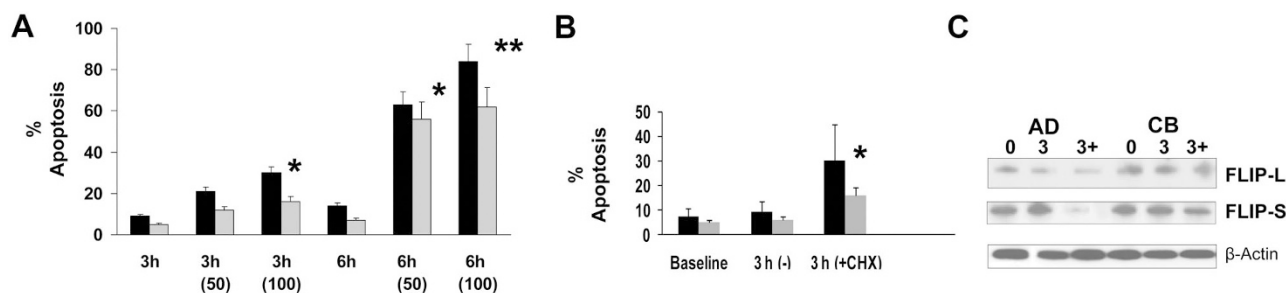


Figure 2. Neonatal and adult PMN apoptosis in response to CHX. (A) PMN isolated from AD (black bars) and CB (gray bars) donors were incubated for up to 6 h in the presence of the protein synthesis inhibitor CHX (50 or 100 $\mu\text{g}/\text{mL}$) or in vehicle alone, and the resultant proportion of apoptotic cells were determined by flow cytometry (TUNEL assay). * $p < 0.05$, ** $p < 0.001$; AD vs CB PMN. Data, representative of seven paired (AD, CB) separate experiments, were analyzed by univariate two-way ANOVA with a fixed factor effect. (B, C) PMN were harvested at baseline (0 h) or after treatment with CHX (100 $\mu\text{g}/\text{mL}$, 3 h) or vehicle control in three separate, paired experiments. (B) PMN were assessed for apoptosis by TUNEL assay. * $p < 0.05$, AD vs CB PMN. (C) FLIP and β -actin were detected by Western blot in AD and CB PMN lysates with monoclonal anti-FLIP antibodies.

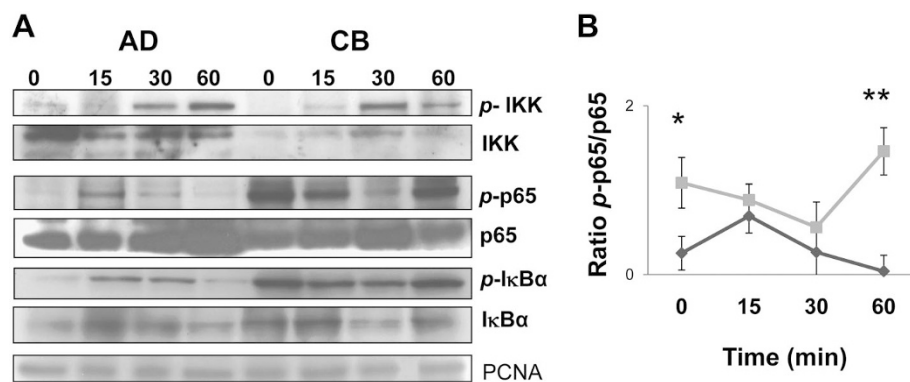


Figure 3. NF- κ B activation in neonatal and adult PMN nuclear fractions. PMN treated with TNF- α (10 ng/mL) were harvested at the indicated times. (A) Nuclear fractions were subjected to Western blotting using antibodies specific to IKK, I κ B α , and p65 (or their phosphorylated forms). Blots were probed with anti-PCNA antibodies as a loading control. (B) Levels of phosphorylated p65 protein were normalized to total cognate protein, and the mean densitometric ratios are shown in the graph. * $p < 0.05$ and ** $p < 0.01$, AD PMN (black line) vs CB PMN (gray line). Blot shown is representative of three paired (AD, CB), separate experiments.

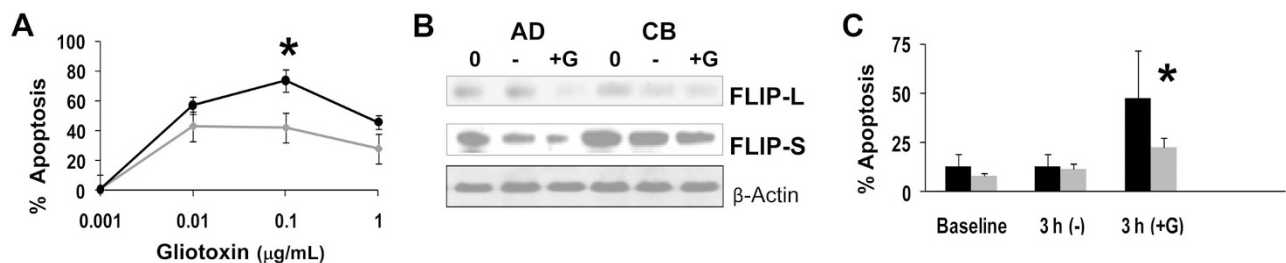


Figure 4. Sensitivity of neonatal and adult PMN to gliotoxin-induced apoptosis. (A) Neonatal (CB) and adult (AD) PMN were incubated for 6 h with increasing concentrations of gliotoxin, a specific NF- κ B inhibitor. The proportions of apoptotic PMN in gliotoxin-treated cultures relative to apoptotic PMN in control cultures were determined by flow cytometry (TUNEL assay). The percentages of apoptotic PMN in CB cultures (gray line) peaked at a lower gliotoxin dose, and peak levels were lower than for AD PMN (black line; * $p < 0.001$). Data represent the mean values of eight separate, paired (AD, CB) experiments. To correlate FLIP expression with apoptosis, PMN were incubated for 3 h with gliotoxin (0.1 $\mu\text{g}/\text{mL}$); PMN lysates or intact PMN were assessed by Western blot (B) or flow cytometry (C), respectively. (B) FLIP and β -actin proteins were detected by Western blot in PMN lysates. Blot shown is representative of three paired (AD, CB), separate experiments. (C) Apoptotic fractions in AD (black bars) and CB (gray bars) PMN were determined by TUNEL assay. * $p < 0.05$, adult vs neonatal PMN. Data represent the means of five separate, paired studies.

Differential gliotoxin effect on apoptosis and FLIP expression in neonatal and adult PMN. We next examined whether augmented constitutive activation of NF- κ B might underlie the relative resistance of neonatal PMN to spontaneous apoptosis (3). Neonatal and adult PMN were incubated with gliotoxin, a specific NF- κ B inhibitor (22), and apoptosis determined by flow cytometry. Gliotoxin treatment induced a peak apoptotic response in neonatal PMN (0.01 $\mu\text{g}/\text{mL}$, 49 \pm 18%) at a 10-fold lower dose than the higher peak response in adult PMN (0.1 $\mu\text{g}/\text{mL}$, 76 \pm 10%; $p < 0.001$; Fig. 4A). As FLIP expression is known to be regulated by NF- κ B (14), we examined the effect of gliotoxin treatment on FLIP expression

in conjunction with apoptosis. FLIP expression remained prominent in treated neonatal PMN, whereas it decreased in adult PMN (Fig. 4B). FLIP-S expression was greater than that of FLIP-L, as previously reported (13). Persistence of FLIP-S expression in gliotoxin-treated neonatal PMN paralleled relatively diminished apoptotic responses (Fig. 4C), whereas diminished levels of FLIP-S corresponded to increased apoptosis in adult PMN.

DISCUSSION

Studies from our laboratory and of others suggest the existence of preformed survival factors (3) and enhanced

NF- κ B activation (21,25) in neonatal PMN. These observations led us to hypothesize that the constitutive presence of FLIP, an antiapoptotic protein regulated by NF- κ B, might be one mechanism underlying the relative resistance of neonatal PMN to spontaneous apoptosis (3,4). FLIP is a critical negative regulator of apoptosis associated with the death receptor complex and caspase activation (26). FLIP exists in two main forms: a prominent longer isoform (FLIP-L) that contains tandem death effector domains (DED) similar to caspase-8, followed by a caspase-like domain, and a shorter isoform, FLIP-S, that possesses only DEDs (27,28). Neutrophils express lower levels of FLIP-L relative to FLIP-S (13), an observation confirmed in this study.

We observed a greater constitutive expression of FLIP-S in neonatal PMN relative to those of adults. Diminished FLIP expression was paralleled by a greater fraction of apoptotic adult PMN, as previously reported (13). In contrast, the persistence of FLIP-S in neonatal PMN during spontaneous apoptosis is consistent with their protracted lifespan (3,4). Persistent FLIP expression has been shown to modulate the resistance of inflammatory macrophages to Fas-mediated apoptosis in patients with rheumatoid arthritis (29,30), and neonatal PMN are also relatively resistant to apoptosis mediated by Fas (3,4). These observations suggest that augmented FLIP expression may directly promote survival in neonatal PMN, although the exact mechanism remains to be discerned. FLIP-S has been shown to inhibit caspase-8/-3 and consequently the intrinsic pathway of apoptosis, particularly in PMN (27,28). Neonatal PMN have blunted activity of these caspases (4,8), which suggests that FLIP may modulate their survival by blocking downstream apoptotic signaling (27) and is a premise that warrants further investigation.

Neonatal PMN persistently expressed FLIP-S protein despite inhibition of *de novo* synthesis (Fig. 2C), consistent with the presence of preformed protein (11). In contrast, FLIP expression was diminished in similarly treated adult PMN, as previously reported in several cell lines, including HL-60 cells (31,32). Neutrophils exhibit constitutive NF- κ B signaling (33,34), which can also induce FLIP synthesis (14,35). This evidence led us to examine a potential contribution of NF- κ B to the prominent basal expression of FLIP-S in neonatal PMN. NF- κ B signaling in PMN involves nuclear IKK activation, phosphorylation/degradation of I κ B α and p65/RelA phosphorylation (23,35,36). In this study, neonatal PMN exhibited earlier I κ B α degradation and prominent basal phosphorylation levels of p65/RelA (Fig. 3), consistent with NF- κ B activation status (23,35–38). NF- κ B signaling is important for constitutive IL-8 promoter activity and IL-8 synthesis (34,39), as well as Akt phosphorylation (40). Pertinently, both IL-8 release and Akt activation status are enhanced under basal conditions in neonatal PMN (5,25), providing further evidence of augmented constitutive NF- κ B activation in neonatal PMN. Akt activation may also be involved in the persistence of FLIP expression by inducing a stabilizing protein that blocks FLIP ubiquitination (41,42), a process that results in proteasomal degradation (43). Although ubiquitination of FLIP has not been reported in PMN, our results could be consistent with the existence of such a protein in neonatal PMN, a mechanism

with implications for their prolonged survival and inflammatory potential (44).

Studies conducted to correlate basal NF- κ B activity with survival function showed a greater sensitivity of neonatal PMN to the apoptotic effects of low-dose gliotoxin under basal conditions, also consistent with enhanced constitutive NF- κ B activity. Unexpectedly, however, higher doses of gliotoxin induced a greater overall apoptotic response in adult PMN. One possible explanation for these observations could be that inhibition of NF- κ B activity in adult PMN unmasked proapoptotic signals, such as caspases and/or cJun N-terminal kinase (13,45,46). This possibility is strengthened by the relative prominence of caspases and other proapoptotic proteins in adult PMN (4,8). Conversely, the muted apoptotic effect of gliotoxin on neonatal PMN could reflect a predominance of survival signaling mechanisms. Inhibition of NF- κ B was shown to promote PMN survival mediated by p38-MAPK (47), although whether p38-MAPK signaling is altered in neonatal PMN is unclear (4,48). Alternatively, robust basal Akt activation status in neonatal PMN (5) suggests that prominence of this survival pathway may be a plausible explanation for our observations. This premise is reinforced by findings that PMN survival may be more dependent on Akt than on NF- κ B activation mechanisms (49). Furthermore, Akt potentially induces FLIP expression and promotes its stability (41,50), which also suggests its contribution to the persistent FLIP expression in neonatal PMN observed in this study. However, further investigation, beyond the scope of this article, will be required to dissect these potential mechanisms.

In summary, this report establishes a novel association between enhanced FLIP-S expression and the preferential survival of neonatal PMN. In addition, our data suggest a link between FLIP expression and a prominence of NF- κ B and related survival signaling pathways. Although the contributory mechanisms remain unclear, age-related differences in FLIP expression are likely important to augmented PMN inflammatory potential. Thus, therapeutic targeting of FLIP or related upstream pathways may be particularly relevant to inflammation in the developing human.

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