

PKR Protects Colonic Epithelium Against Colitis Through the Unfolded Protein Response and Prosurvival Signaling

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Background: The dsRNA-activated protein kinase (PKR) phosphorylates the α subunit of eukaryotic translation initiation factor 2 (eIF2 α), a global regulator of protein synthesis in mammals. In addition, PKR activates several signal transduction pathways including STAT3 and AKT. PKR is activated by a number of inflammatory stimuli that are induced in the inflamed intestine. In this study we intended to determine the role of PKR in colonic epithelial cells during experimental colitis in mice.

Methods: Age- and sex-matched $PKR^{+/+,+/-}$ and $PKR^{-/-}$ littermate mice were reconstituted with wildtype bone marrow cells and subjected to dextran sodium sulfate (DSS)-induced colitis.

Results: $PKR^{-/-}$ mice displayed more severe clinical and histological manifestations upon DSS colitis compared with their $PKR^{+/+,+/-}$ littermates. In response to DSS colitis, the colonic epithelial cells of $PKR^{-/-}$ mice exhibited impaired activation of the unfolded protein response (UPR) signaling, including eIF2 α phosphorylation, endoplasmic reticulum (ER) chaperone response, and ER-associated degradation (ERAD) components, as well as antioxidative stress response. In addition, the phosphorylation of STAT3 and AKT, which are protective against epithelial cell death and colonic inflammation, was also impaired in the colonic epithelial cells of $PKR^{-/-}$ mice upon DSS colitis.

Conclusions: These data demonstrate that PKR is a physiologically relevant transducer of inflammatory response signaling in colonic epithelial cells. PKR may promote the homeostasis and survival of intestinal epithelial cells (IECs) through eIF2 α -mediated UPR activation, as well as the activation of STAT3 and AKT pathways. In the absence of PKR, the survival and proliferation of IECs was impaired, thus exacerbating intestinal inflammation.

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Key Words: PKR, DSS colitis, UPR, prosurvival signaling

The endoplasmic reticulum (ER) in eukaryotic cells is the site for the folding, assembly, modification, and maturation of polypeptides destined for secretion, membranes, and intracellular organelles. Protein folding in the ER is exquisitely sensitive to multiple environmental alterations and cellular disturbances including ER Ca²⁺ depletion, oxidative stress, and inflammatory stimuli. Disrupted ER protein folding homeostasis leads to accumulation of unfolded/misfolded proteins in the ER lumen, a condition

termed ER stress. To cope with this cellular stress and restore ER homeostasis, eukaryotic cells have evolved the unfolded protein response (UPR). In mammalian cells, three ER-localized stress sensors signal the UPR: inositol-requiring kinase 1 (IRE1), activating transcription factor 6 (ATF6), and PKR-like ER kinase (PERK). The UPR protein sensors are activated in response to ER stress and signal downstream pathways that control transcriptional, translational, and posttranslational processes. During ER stress, the ER chaperone BiP releases from the luminal domain of PERK, which is activated through dimerization and *trans*-autophosphorylation. Activated PERK then phosphorylates the α subunit of eukaryotic translation initiation factor 2 (eIF2 α) to reduce global protein synthesis. At the same time, phosphorylated eIF2 α induces the translation of mRNA encoding activating transcription factor 4 (ATF4), which subsequently transactivates genes encoding other UPR-associated transcription factors, ER protein-folding machinery, and components of ER-associated protein degradation (ERAD), and promotes cellular homeostasis by inducing the biosynthesis and transport of amino acids, and spurring the antioxidative stress response.¹

In addition to PERK, three cytosolic eIF2 α kinases exist in mammals, including the double-stranded

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RNA-activated protein kinase (PKR).² PKR was originally identified and characterized as a pathogen sensor and mediator of the interferon response to protect the host from viral infection. PKR-mediated eIF2 α phosphorylation is a strategy to shut down viral protein synthesis in host cells.³ However, in contexts other than viral infection induction of downstream UPR signaling and inhibition of global translation through the same eIF2 α phosphorylation by PKR activation may be crucial for ER homeostasis in response to various environmental alterations.² Indeed, a number of reports have demonstrated that PKR does signal protection to ER stress. In addition to dsRNA, PKR can be activated by other inflammatory signals including lipopolysaccharide (LPS) and tumor necrosis factor alpha (TNF- α), as well as oxidative stress. When activated, PKR induces or acts in conjugation with a number of signal transduction pathways, including p38 mitogen-activated protein kinase (MAPK), c-Jun N-terminal kinases (JNK), signal transducer and activator of transcription 3 (STAT3), and Akt/Protein kinase B.⁴ Therefore, PKR functions as a regulatory hub that coordinates inflammatory response signaling, pathogen sensing, and UPR signaling in mammalian cells.

Inflammatory bowel diseases (IBDs) represent a set of inflammatory conditions in the gastrointestinal tract. Although IBDs were discovered dozens of years ago, the etiology of IBD is still unclear. Recently, some studies linked ER stress and UPR signaling to the pathogenesis of IBD by showing that patients with active Crohn's disease and ulcerative colitis exhibit signs of ER stress in their ileal and/or colonic epithelium.⁵⁻⁸ Genes that regulate ER homeostasis and the UPR including *XBPI* have been identified to associate with Crohn's disease and/or ulcerative colitis. The murine and human intestinal epithelium contains Paneth and goblet cells, which have massive secretory functions and play essential roles in intestinal homeostasis and host defense. Intestinal epithelial cell (IEC)-specific deletion of *XBPI* caused Paneth cell death and spontaneous inflammation in murine ileum.⁸ A previous study suggested that PKR may regulate the differentiation of a rat intestinal epithelial cell line by promoting the activity of alkaline phosphatase.⁹ However, the role of PKR in intestinal inflammation is unclear. Here we show that the deletion of PKR increases sensitivity to dextran sodium sulfate (DSS)-induced colitis in mice, due to reduced induction of adaptive UPR signaling and prosurvival signaling including the activation of STAT3 in colonic epithelial cells during inflammation.

MATERIALS AND METHODS

Mice

All mice were housed on 12/12 hour light/dark cycles at the Unit for Laboratory Animal Medicine (ULAM) at the Uni-

versity of Michigan Medical Center with free access to water and standard rodent chow. All animal care and procedures were conducted following the protocols and guidelines approved by the University of Michigan Committee on the Use and Care of Animals (UCUCA).

Generation of Bone Marrow Chimeras

Eight-week-old *PKR*^{-/-} and *PKR*^{+/+, +/-} littermate mice¹⁰ were lethally irradiated with 950 rad ionizing irradiation. After 2 hours, 5×10^6 bone marrow cells isolated from the tibias and femurs of wildtype mice were injected into the lethally irradiated recipient mice through the tail vein. After transplantation, mice were treated with a 6-week course of antibiotics in their drinking water and allowed to recover for another 2 weeks prior to treatment with DSS to induce colitis.

To generate chimeric mice with deletion of *PKR* in their hematopoietic compartment, 8-week-old wildtype mice on a C57BL/6 background (Jackson Laboratory, Bar Harbor, ME) were lethally irradiated with 950 rad ionizing irradiation. After 2 hours these mice received a tail vein injection of 5×10^6 bone marrow cells isolated from *PKR*^{-/-} or *PKR*^{+/+} littermate mice. After transplantation mice were treated with a 6-week course of antibiotics in their drinking water and allowed to recover for another 2 weeks prior to experiment.

DSS-induced Colitis

For DSS-induced acute colitis, *PKR*^{-/-} and *PKR*^{+/+, +/-} mice reconstituted with wildtype bone marrow cells as well as wildtype mice reconstituted with *PKR*^{-/-} or *PKR*^{+/+} bone marrow cells received 3% (w/v) DSS (MW 36,000-50,000; MP Biomedicals, Solon, OH) in drinking water for 7 days. Body weight change and rectal bleeding were monitored daily. After DSS administration the mice were euthanized, colon length was measured, and colon sections and colonic epithelial cells were isolated for histological and biochemical analyses.

Isolation of Colonic Epithelial Cells

The entire colons were cut open longitudinally, then feces was removed by washing with ice-cold phosphate-buffered saline (PBS). Colons were then cut into 2-3-mm pieces and incubated in Ca²⁺, Mg²⁺-free PBS buffer containing 10 mM EDTA in a 50 mL conical tube at 4°C for 1 hour with gentle rotation. Then the tubes were rigorously shaken to elute the epithelium from colon sections. The supernatant was removed and sieved through a cell strainer (500 μ m; Fisherbrand, Pittsburgh, PA). The flow-through was centrifuged; the resulting cell pellet was washed twice in ice-cold PBS and snap-frozen in liquid nitrogen for protein and RNA extraction. The purity of isolated colonic epithelial cells was confirmed by FACS and quantitative reverse-transcription polymerase chain reaction (Q-RT-PCR). Trypan blue staining confirmed the presence of >85% viable epithelial cells after the 2-hour isolation procedure.

Quantitative Real-time PCR

RNA from isolated colonic epithelial cells and IEC-6 cells were extracted by using RNeasy kit (Qiagen, Chatsworth, CA); RNA from 5 mm distal colons was isolated using TRIzol reagent (Invitrogen, La Jolla, CA). Q-RT-PCR was performed as previously described.¹¹ Q-RT-PCR results were normalized to the levels of 18S rRNA or the mRNA encoding glyceraldehyde 3-phosphate dehydrogenase. Primer sequences are listed in Table 1.

Western Blotting

The isolated colonic epithelial cells were lysed in RIPA buffer for protein extraction. Protein content was measured by Protein Assay (Bio-Rad, Hercules, CA), and protein samples were analyzed by 10%–20% reducing sodium dodecyl sulfate-polyacrylamide gel electrophoresis (SDS/PAGE) (Tris-HCl precast gels, Bio-Rad) and were detected with the following antibodies: anti-GRP94/BiP (StressGen, Vancouver, BC, Canada), anti-phosphor-PERK (Cell Signaling Technology, Beverly, MA), anti-ATF4 (Santa Cruz Biotechnology, Santa Cruz, CA), anti-CHOP (Santa Cruz Biotechnology), anti-XBP1 (Santa Cruz Biotechnology), anti-LC3 (Novus Biologicals, Littleton, CO), and anti-glyceraldehyde 3-phosphate dehydrogenase (Millipore, Bedford, MA), and anti-phosphor-STAT3 (Cell Signaling Technology). After overnight incubation of primary antibodies at 4°C the membranes were washed and then incubated with secondary horseradish peroxidase (HRP)-coupled antibodies (GE Healthcare, Milwaukee, WI) and developed with a chemiluminescent detection system (GE Healthcare).

Immunohistochemical Staining

Colonic tissues were fixed overnight in 10% formalin, embedded in paraffin, and cut into 5- μ m sections. Paraffin sections were rehydrated and heat-induced antigen retrieval was then performed in 10 mM sodium citrate, 0.05% Tween-20, pH 6.0, for 10 minutes. Primary antibodies: anti-BiP antibody (Santa Cruz Biotechnology) at a dilution of 1:200; anti-ATF4 antibody (LifeSpan BioSciences, Seattle, WA) at a dilution of 1:200; anti-phosphor-STAT3 antibody (LifeSpan BioSciences) at a dilution of 1:200. Hematoxylin was performed for counterstaining.

Histological Scoring

Colonic tissues were fixed overnight in 10% formalin, embedded in paraffin, and cut into 5- μ m sections for hematoxylin and eosin (H&E) staining using a standard protocol. H&E-stained sections were scored in a blinded manner by a licensed animal pathologist for the amount of tissue damage and inflammatory cell infiltration as previously described.¹² The damaged area involved was scored on a scale of 0 to 4: 0 = no involvement, 1 \leq 25%, 2 \leq 50%, 3 \leq 75%, 4 \leq 100%. Inflammatory cell infiltration were scored on a scale of 0 to 3: 0 = occasional inflammatory cells in the lamina propria, 1 = increased numbers of inflammatory cells in the lamina propria,

TABLE 1. Sequence of Primers

Primer Name	Oligo Sequence (5' to 3')
mouse IL-1 β (f)	CAACCAACAAGTGATATTCTCCATG
mouse IL-1 β (r)	GATCCACACTCTCCAGCTGCA
mouse TNF α (f)	CCCTCACACTCAGATCATCTTCT
mouse TNF α (r)	GCTACGACGTGGGCTACAG
mouse iNOS (f)	CAGCTGGGCTGTACAAACCTT
mouse iNOS (r)	CATTGGAAGTGAAGCGTTTCG
mouse BiP (f)	TCATCGGACGCACTTGGG
mouse BiP (r)	CAACCACCTTGAATGGCAAGA
mouse GRP94 (f)	AATAGAAAAGAATGCTTCGCC
mouse GRP94 (r)	TCTTCAGGCTCTTCTTCTGG
mouse ERp72 (f)	AGTCAAGGTGGTGGTGGGAAAG
mouse ERp72 (r)	TGGGAGCAAATAGATGGTAGGG
mouse ERdj4 (f)	CCCCAGTGTCAAACCTGTACCAG
mouse ERdj4 (r)	AGCGTTTCCAATTTCCATAAATT
mouse P58IPK (f)	TCCTGGTGGACCTGCAGTACG
mouse P58IPK (r)	CTGCGAGTAATTTCTTCCCC
mouse calreticulum (f)	GAGTGGCTTGGACCAGAAGG
mouse calreticulum (r)	GGACCGCAGATGTCCGG
mouse Ero1 α (f)	GCATTGAAGAAGGTGAGCAA
mouse Ero1 α (r)	ATCATGCTTGGTCCACTGAA
mouse Ero1 β (f)	GGGCCAAGTCATTAAGGAA
mouse Ero1 β (r)	TTTATCGCACCCAACACAGT
mouse PDI (f)	CAAGATCAAGCCCCACCTGAT
mouse PDI (r)	AGTTCGCCCCAACCACTACTT
mouse XBP1 (f)	AAGAACACGCTTGGGAATGG
mouse XBP1 (r)	ACTCCCCCTTGGCCTCCAC
mouse XBP1s (f)	GAGTCCGCAGCAGGTG
mouse XBP1s (r)	GTGTCAGAGTCCATGGGA
mouse ATF4 (f)	ATGGCCGGCTATGGATGAT
mouse ATF4 (r)	CGAAGTCAAACCTTTTCAGATCCATT
mouse Chop (f)	GTCCCTAGCTTGGCTGACAGA
mouse Chop (r)	TGGAGAGCGAGGGCTTTG
mouse Gadd34 (f)	CCCGAGATTCTCTAAAAGC
mouse Gadd34 (r)	CCAGACAGCAAGGAAATGG
mouse ATF6 α (f)	CTTCTCCAGTTGCTCCATC
mouse ATF6 α (r)	CAACTCCTCAGGAACGTGCT
mouse Herpud1 (f)	AGCAGCCGGACAACCTAAT
mouse Herpud1 (r)	CTTGGAAGTCTGCTGGACA
mouse Wfs1 (f)	CCATCAACATGCTCCCGTTC
mouse Wfs1 (r)	GGGTAGGCCTCGCCAT
mouse Nrf2 (f)	ACATCCTTTGGAGGCAAGAC
mouse Nrf2 (r)	GCCTTCTCTGTTCTTCTG
mouse GAPDH (f)	TTCAACGGCACAGTCAAGG
mouse GAPDH (r)	CATGGACTGTGGTCATGAG

2 = confluence inflammatory cells, extending into the submucosa, and 3 = transmural extension of the infiltrate.

Statistical Analysis

All data are presented as means \pm SEM. Statistical significance of the difference between groups was analyzed using Student's *t*-test. $P < 0.05$ was considered significant.

RESULTS

Mice with PKR Deletion in Nonhematopoietic Cells Are More Sensitive to DSS-induced Colitis

In the absence of inflammatory insults, the colon of $PKR^{-/-}$ mice is morphologically indistinguishable from that of their wildtype littermates. To analyze a possible requirement for PKR in IECs to prevent IBD, lethally irradiated age- and sex-matched $PKR^{+/+, +/-}$ and $PKR^{-/-}$ littermate mice were reconstituted with wildtype bone marrow and then fed 3% DSS in the drinking water for 7 days to induce colitis. RT-PCR demonstrated that the level of *Pkr* mRNA in colonic epithelial cells of $PKR^{-/-}$ mice was less than 4% of that in $PKR^{+/+}$ or $PKR^{+/-}$ littermate mice 8 weeks after bone marrow replacement (Fig. 1A), indicating that the intestinal epithelium was not repopulated by wildtype stem cells from the bone marrow transplantation. $PKR^{-/-}$ mice showed more severe clinical manifestations, as demonstrated by body weight loss, rectal bleeding, and colon shortening upon DSS challenge (Fig. 1B–D). In addition,

$PKR^{-/-}$ mice displayed significantly higher levels of damaged area in the epithelium and inflammatory cell infiltration in both the proximal and distal colon (Fig. 2A,B). Consistent with the histological observations, the expression of proinflammatory genes *Il-1 β* , *Tnf- α* , and *Inos* were dramatically induced in the colon of $PKR^{-/-}$ mice upon DSS colitis compared with their wildtype and heterozygous littermates (Fig. 3). In contrast, wildtype mice reconstituted with $PKR^{-/-}$ bone marrow cells showed similar body weight loss upon DSS colitis as compared with wildtype mice reconstituted with $PKR^{+/+}$ bone marrow cells (Fig. 1E), indicating that the deletion of *PKR* in hematopoietic cells does not alter the sensitivity to DSS-induced colitis.

UPR Induction Is Impaired in Colonic Epithelial Cells of $PKR^{-/-}$ Mice

Our previous studies demonstrated that UPR molecules, e.g., BiP and ATF4, are induced in colonic epithelium in response to DSS colitis (unpublished data). Given the role of PKR as an inducer of UPR signaling, we then examined whether UPR activation is impaired in $PKR^{-/-}$ colonic epithelial cells during inflammation. Western blotting demonstrated that colonic epithelial cells of $PKR^{-/-}$ mice exhibited reduced levels of eIF2 α phosphorylation and its downstream target ATF4 compared with their wildtype littermate mice. The induction of the ER chaperones BiP and GRP94 was also impaired at both the protein and

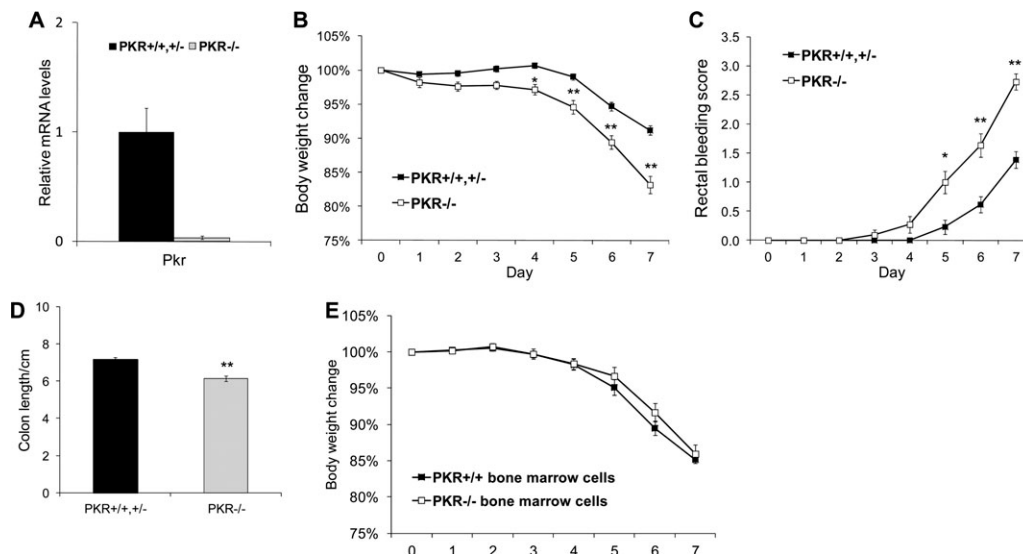


FIGURE 1. $PKR^{-/-}$ mice are more sensitive to DSS-induced colitis. $PKR^{-/-}$ and $PKR^{+/+, +/-}$ littermates were reconstituted with wildtype bone marrow cells. After 8 weeks the mice were fed 3% DSS in drinking water for 7 days; their body weight and rectal bleeding were monitored daily. (A) Colonic epithelial cells isolated from $PKR^{-/-}$ mice after bone marrow transplantation show very low levels of *Pkr* mRNA. (B) $PKR^{-/-}$ mice show more significant body weight loss during DSS colitis. (C) $PKR^{-/-}$ mice show more severe rectal bleeding during DSS colitis. (D) $PKR^{-/-}$ mice show more severe colon shortening after 7-day of DSS administration. $PKR^{+/+, +/-}$: $n = 13$; $PKR^{-/-}$: $n = 11$; * $P < 0.05$, ** $P < 0.01$. (E) Two-month-old C57BL/6J wildtype mice were reconstituted with $PKR^{+/+}$ or $PKR^{-/-}$ bone marrow cells. After 8 weeks the mice were fed 3% DSS in drinking water for 5 days followed by 2 days of free water; their body weight was monitored daily. $PKR^{+/+}$: $n = 8$; $PKR^{-/-}$: $n = 7$.

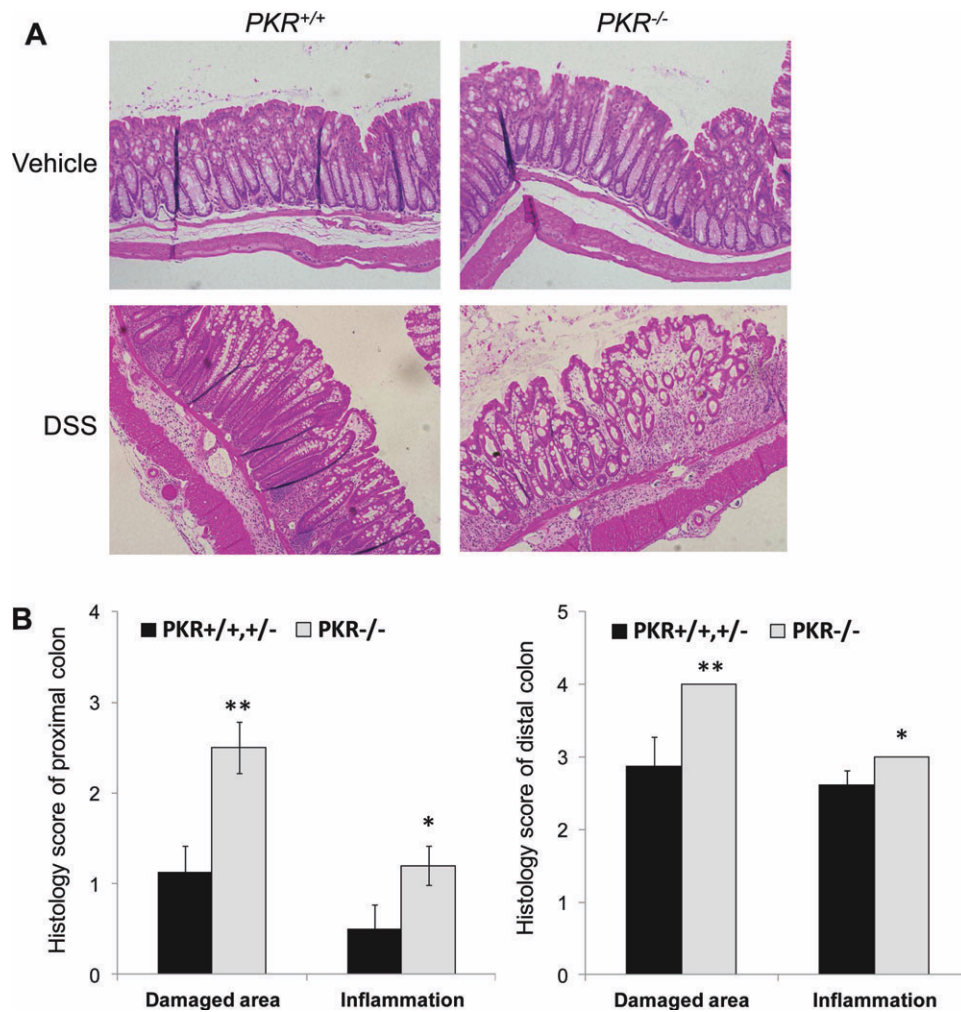


FIGURE 2. $PKR^{-/-}$ mice show more severe histological manifestation after 7 days of DSS administration. $PKR^{-/-}$ and $PKR^{+/+, +/-}$ littermates reconstituted with wildtype bone marrow cells were fed 3% DSS in drinking water for 7 days. (A) After DSS administration the colons were isolated and fixed for H&E staining. Representative images are shown (100 \times). (B) Histological scores including damage area involved and inflammatory cell infiltration taken from $PKR^{-/-}$ and $PKR^{+/+, +/-}$ littermates with DSS colitis. $PKR^{+/+, +/-}$: $n = 8$; $PKR^{-/-}$: $n = 10$; * $P < 0.05$, ** $P < 0.01$.

RNA levels (Fig. 4A,C). In addition to ER chaperones, the induction of a number of UPR pathways were compromised in $PKR^{-/-}$ mice upon DSS colitis, including the UPR transcription factors XBP1s and ATF6, ER protein folding machinery ERp72, Ero1 α , and PDI, ERAD components Herpud1 and Wfs1, and the antioxidative stress response nuclear factor-erythroid 2-related factor 2 (Nrf2).

Previous studies demonstrated that these adaptive UPR signaling pathways are essential during intestinal inflammation. The UPR-induced chaperone response in colonic epithelium has been shown to protect against DSS colitis in mice.¹³ Loss of Nrf2 increases susceptibility to DSS colitis, probably a consequence of enhanced oxidative damage and proinflammatory responses in the colonic mucosa.¹⁴ Therefore, PKR may protect colonic epithelial cells during inflammation through the induction of UPR signaling. In the absence of PKR, UPR signaling was impaired

and may exacerbate ER stress and induce apoptosis. Consistently, the cleavage of caspase-3 was highly activated in colonic epithelial cells of $PKR^{-/-}$ mice after DSS challenge (Fig. 4A).

Prosurvival Signaling Is Compromised in the Absence of PKR During DSS Colitis

Given that PKR is an upstream inducer of STAT3 and Akt phosphorylation,⁴ we next examined whether the activation of these two signaling molecules was impaired in colonic epithelium of $PKR^{-/-}$ mice. Western blotting demonstrated that colonic epithelial cells isolated from $PKR^{-/-}$ mice challenged with DSS exhibited reduced phosphorylation of STAT3 and Akt compared with their wild-type littermate mice (Fig. 4A). Consistently, the nuclear staining of phosphor-STAT3 in the epithelial cells was diminished in the colon of $PKR^{-/-}$ mice (Fig. 4C).

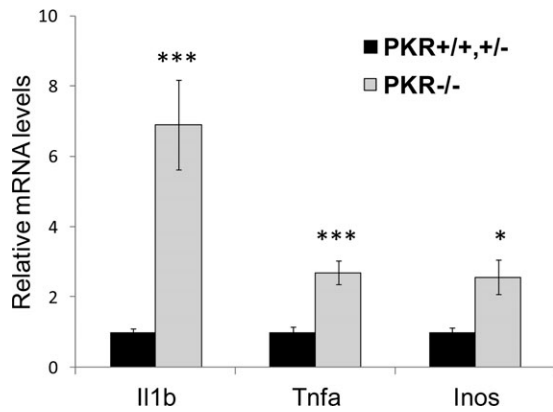


FIGURE 3. *PKR*^{-/-} mice show hyperactivated inflammatory response in the colon upon DSS administration. *PKR*^{-/-} and *PKR*^{+/+, +/-} littermates with wildtype bone marrow cells were fed 3% DSS in drinking water for 7 days. After DSS administration the mice were euthanized and colonic epithelial cells were collected for RNA extraction, cDNA synthesis, and Q-RT-PCR. The mRNA levels were normalized to the expression of *Gapdh*. *PKR*^{+/+, +/-}: n = 6; *PKR*^{-/-}: n = 5; *P < 0.05, **P < 0.01, ***P < 0.001.

DISCUSSION

In this study we demonstrate that PKR is a physiologically relevant activator of UPR signaling in colonic epithelium upon inflammation. In response to DSS colitis, PKR promoted homeostasis and survival of IECs through eIF2 α -mediated UPR signaling. *PKR*^{-/-} IECs showed reduced phosphorylation of eIF2 α upon colitis, consistent with the diminished eIF2 α phosphorylation observed in *PKR*^{-/-} mouse embryo fibroblasts (MEFs) in response to inflammatory stimuli including IFN α/β .¹⁵ Reduced eIF2 α phosphorylation in *PKR*^{-/-} IECs upon colitis impaired the activation of UPR, e.g., transcription factors, chaperone response, ERAD components, and antioxidative stress response. This is consistent with our observations that mice expressing a nonphosphorylatable eIF2 α in IECs are susceptible to DSS colitis and display diminished UPR induction in their colonic epithelial cells upon colitis (unpublished data). The UPR-activated transcription factors XBP1, ATF6 α , and ATF4 are essential transactivators of

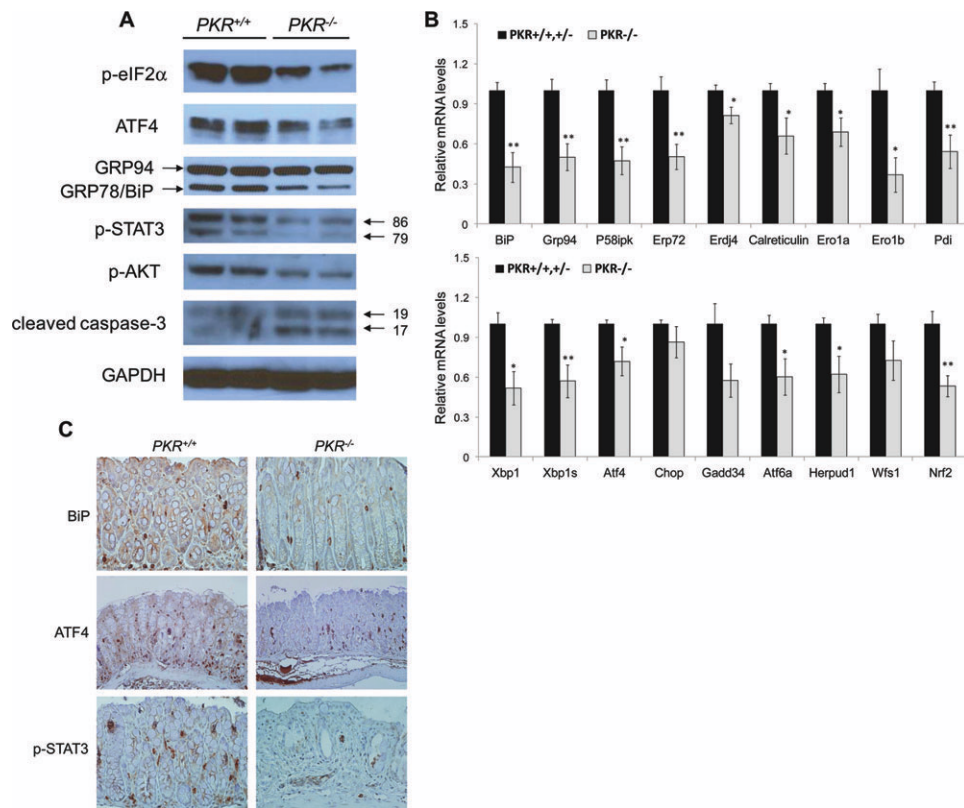


FIGURE 4. Activation of the UPR signaling and prosurvival pathways is impaired in *PKR*^{-/-} colonic epithelial cells upon DSS colitis. *PKR*^{-/-} and *PKR*^{+/+, +/-} littermates with wildtype bone marrow cells were fed 3% DSS in drinking water for 6 days. (A) After DSS administration the mice were euthanized and the colonic epithelial cells were isolated for protein extraction and immunoblotting. (B) After DSS administration the mice were euthanized and the colonic epithelial cells were isolated for RNA extraction, cDNA synthesis, and Q-RT-PCR. The mRNA levels were normalized to the expression of *Gapdh*. (C) After DSS administration the mice were euthanized and the colons were isolated, fixed, and paraffin-embedded for immunohistochemical staining of BiP, ATF4, and phospho-STAT3 (200 \times or 400 \times). *PKR*^{+/+, +/-}: n = 6; *PKR*^{-/-}: n = 6; *P < 0.05, **P < 0.01.

UPR gene induction in response to ER stress. Mice deficient in XBP1 in the intestinal epithelium were sensitive to DSS colitis, although the underlying mechanism is not yet clear.⁸ ATF6 α is activated upon ER stress by sequential cleavage by proteases S1P and S2P, yielding an active ATF6 α p50 that functions as a potent activator of a number of ER chaperones including BiP, GRP94, and P58^{IPK}. The hypomorphic mutation of S1P-encoding gene *Mbtps1* impaired ATF6 α -dependent induction of chaperone response, and increased susceptibility to experiment colitis in mice.¹³ ATF4 is induced at both the transcriptional and translational levels during ER stress and functions as an essential global UPR transactivator.² In addition, ATF4 is activated by oxidative stress and plays an important role in the antioxidative stress response.¹⁶ Although there is little information about the role of ATF4 in IBD, it is possible that ATF4 is protective against ER stress and oxidative stress in IECs during intestinal inflammation. The importance of UPR-induced ER chaperones in DSS colitis is directly supported by our study using a murine model deficient in P58^{IPK}, a heat-shock 40-kDa protein that normally resides in association with the ER chaperone BiP in the ER lumen and promotes proper protein folding.¹⁷ P58^{IPK}^{-/-} mice are highly susceptible to DSS colitis, probably due to unresolved ER stress and selective activation of proapoptotic UPR signaling in colonic IECs during inflammation (unpublished data). There are few studies examining the role of ERAD pathways in IBD. However, given the function of ERAD in eliminating unfolded/misfolded protein and alleviating ER stress, it is possible that ERAD also impacts IEC function during intestinal inflammation. Therefore, the compromised induction of adaptive UPR pathways may exacerbate the dysfunction of colonic IECs in *PKR*^{-/-} mice upon DSS challenge, and consequently worsen the symptoms of colitis.

PKR has been shown to physically interact with STAT3 and induce the phosphorylation at Tyr705 and Ser727, which is required for DNA binding and the transcriptional activation function of STAT3 upon PDGF stimulation.¹⁸ STAT3 is a pleiotropic transcription factor with crucial roles in a variety of cellular processes including proliferation and differentiation. Previous studies demonstrated that STAT3 activity is increased by phosphorylation in IECs during colonic inflammation, and transactivates the genes responsible for the stress response, apoptosis, and wound healing in IECs. IEC-specific deletion of *STAT3* increased the susceptibility to DSS colitis.¹⁹ Similarly, the pleiotropic protein kinase AKT functions to inhibit IEC apoptosis and promote wound healing during intestinal inflammation.²⁰ Therefore, compromised activation of STAT3 and AKT in *PKR*^{-/-} mice may impair function and survival of IECs, thus promoting intestinal inflammation upon DSS challenge.

Several previous studies have linked PKR to apoptotic cell death. In NIH3T3 and COS-1 cells, TNF- α induces apoptosis through PKR-mediated eIF2 α phosphorylation.²¹ *PKR*^{-/-} MEFs were resistant to apoptosis upon challenge with dsRNA, TNF- α , and LPS.²² Another study showed that dsRNA-activated PKR induced the expression of proapoptotic molecules including Fas and Bax in 3T3 L1 cells.²³ In macrophages, cholesterol/7-ketocholesterol induced oxidative stress and activated the PKR-CHOP pathway that was required for the downstream apoptotic response.²⁴ However, all of these studies were performed in vitro and the cells analyzed were very different from colonic epithelial cells at both the cell biological and physiological levels. Therefore, it seems the protective role of PKR in colonic epithelial cells may be both cell-type and/or disease-specific.

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