Protein Kinase PKR Amplification of Interferon β Induction Occurs through Initiation Factor eIF-2 α -mediated Translational Control^{*}

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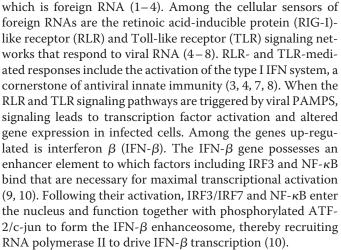
Background: RNA-dependent protein kinase PKR regulates interferon induction.

Results: Knockdown of PKR or mutation of initiation factor eIF- 2α results in increased I κ B- α protein levels and decreased IFN- β induction in RNA-transfected and virus-infected cells.

Conclusion: PKR amplifies IFN- β induction through an eIF-2 α translational control response.

Significance: PKR functions together with additional RNA sensors to modulate signaling leading to interferon gene induction.

The protein kinase PKR is activated by RNA with doublestranded (ds) structure and subsequently impairs translation through phosphorylation of protein synthesis initiation factor eIF- 2α . PKR also mediates activation of signal transduction pathways leading to interferon beta (IFN- β) gene induction following virusinfection or RNA transfection. We previously demonstrated in measles virus-infected cells that PKR is required for the maximal induction of IFN- β gene expression by the interferon promoter stimulator gene 1 (IPS-1) adaptor-dependent cytosolic RNA sensor pathway. While both IPS-1 and PKR are important mediators of IFN-B induction, with PKR contributing to an enhanced NF-kB activation, the mechanism by which PKR enhances NF-kB activity and amplifies IFN- β induction is unresolved. Herein we tested the possibility that PKR could activate signal transduction pathways indirectly through translational control responses. Following transfection with synthetic or natural dsRNAs or infection with measles virus, we observed increased mRNA but decreased protein levels for the inhibitor of NF-κB signaling, IκB-α, that correlated with PKR activation and eIF-2 α phosphorylation. Importantly, knockdown of PKR increased I κ B- α protein levels and impaired IFN- β induction. Additionally, inhibition of translation by cycloheximide treatment rescued IFN-B induction following PKR knockdown but not IPS-1 knockdown. Mutation of eIF-2 α to prevent phosphorylation also impaired IFN- β induction in PKR-sufficient virus-infected cells. These results suggest that an eIF-2 α -dependent translation inhibition mechanism is sufficient to explain the PKR-mediated amplification of IPS-1-dependent IFN-β induction by foreign RNA.



IFN induces the expression of several genes (3), and among them is the double-stranded (ds) RNA-dependent protein kinase (PKR) (11). While inducible by type I IFN signaling, PKR also is constitutively expressed and localizes predominantly to the cytoplasm (12). PKR binds dsRNA as well as structured ssRNAs through tandem dsRNA binding domains present in its N-terminal region. RNA binding can lead to PKR dimerization and activation by autophosphorylation of residues including threonine 446 present in the C-terminal kinase catalytic domain (3, 13, 14). While the PKR protein monomer is able to bind a 15-base pair (bp) dsRNA, ~30-bp are necessary to mediate dimerization of PKR (15), and activation is increased with dsRNA length up to a \sim 85-bp that achieves maximal activation (3, 11, 16). The best characterized substrate of activated PKR remains eukaryotic translation initiation factor eIF-2 α (17), which when phosphorylated on serine 51 subsequently leads to an inhibition of translation initiation (3, 18, 19).

PKR is reported to modulate the functional activity of signal transduction responses, although it is unclear whether these PKR effects are direct or indirect (19). For example, PKR has

eukaryote translation initiation factor- 2α ; IFN, interferon; MV, measles virus; PKR, RNA-dependent protein kinase; PKR^{kd}, PKR knockdown cells; RLR, retinoic acid-inducible protein-like receptor; TLR, Toll-like receptor.



The innate immune system responds to a diverse array of pathogen-associated molecular patterns (PAMPS),³ one of

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³ The abbreviations used are: PAMPS, pathogen-associated molecular patterns; dsRNA, double-stranded RNA; C^{ko} , C knock-out measles virus; eIF-2 α ,

been observed to activate NF- κ B signaling by activating I κ B kinase or NF- κ B-inducing kinase (20, 21) and to interact with TRAF family of protein members (22), however conflicting results were found with two different mouse *pkr* knock-out-derived cell lines regarding NF- κ B activation by PKR (23). In addition to a role for PKR in activation of NF- κ B signaling, PKR is implicated in the activation of proteins including the phosphatase PP2A, p53, stress-activated JNK and p38 kinases, and IRF-1 (19). The mechanism by which PKR activates these diverse pathways largely remains to be elucidated.

We have shown that PKR is required for maximal IFN- β induction in HeLa cells and that this induction of IFN- β gene expression is dependent upon the RLR interferon promoter signaling 1 (IPS-1) adaptor protein, but not the TLR3 adaptor protein TIR-domain-containing adaptor-inducing IFN-β (TRIF) protein (24). Transcriptional induction of IFN- β is known to require the activation of both IRF and NF-KB transcription factors downstream of the TLRs and RLRs (25, 26). We earlier established that while PKR is not required for IRF3 activation following virus infection or RNA transfection, PKR is required for maximal NF- κ B activity (24, 27). The NF- κ B family members involved in type I IFN signaling include the p65 (RelA) and p50 subunits. These proteins heterodimerize and are kept in an inactive state in the cytoplasm of cells by a family of inhibitors of NF- κ B proteins, among which is I κ B- α (26). NF- κ B signaling is activated following the phosphorylation, ubiquitination and degradation of IkB proteins by the inhibitor of NF-KB kinase (IKK) complex (28, 29). Following degradation of the inhibitor, NF-kB subunits localize to the nucleus and bind sites present within the promoter region of NF-κB regulated genes. Among these is $I\kappa B-\alpha$, whose promoter possesses multiple NF-*k*B binding sites, and following NF-*k*B activation, IκB- α thereby negatively regulates NF- κ B signaling (29, 30).

To probe the mechanism by which PKR amplifies IFN- β induction, we tested the possibility that PKR-mediated inhibition of translation triggered by foreign RNA was sufficient to enhance NF-κB activity and amplify IFN-β transcript induction. We found that the PKR-dependent inhibition of $I\kappa B-\alpha$ protein expression in RNA-transfected and measles virus-infected cells correlated with increased phosphorylation of PKR and eIF-2 α . This response was accompanied by enhanced expression of I κ B- α and IFN- β RNAs, but reduced levels of I κ B- α protein. Knockdown of PKR or expression of a serine-toalanine mutant initiation factor eIF-2 α in PKR-sufficient cells resulted in increased I κ B- α protein expression and decreased IFN- β induction following RNA transfection or virus infection. Cycloheximide treatment, which inhibits translation independently of eIF-2 α phosphorylation, rescued IFN- β RNA induction in PKR knockdown cells, but importantly did not rescue IFN-β induction following IPS-1 knockdown. These results suggest that PKR amplification of IFN-β RNA induction in response to foreign RNA occurs through a translational control mechanism.

EXPERIMENTAL PROCEDURES

Cells and Virus—PKR-sufficient parental (PKR⁺) HeLa cells and PKR-deficient HeLa cells stably knocked down for PKR expression (PKR^{kd}), were as previously described (31, 32).

PKR-mediated Enhancement of IFN- β

Recombinant parental measles virus (MV) MVvac GFP(H) designated wild-type (WT) virus, and isogenic mutant viruses either V-deficient (V^{ko}) or C-deficient (C^{ko}), were generously provided by R. Cattaneo (Mayo Clinic, Rochester, MN) and were as previously described (33). The viruses were constructed based on the Moraten vaccine strain (34, 35), except that a gene encoding green fluorescent protein (GFP) was inserted downstream of the H gene.

For MV infection, PKR⁺ or PKR^{kd} cells were seeded into 12-well plates 24 h prior to infection. Cell monolayers were washed once with Opti-MEM I and then infected with WT, V^{ko}, or C^{ko} MV at a multiplicity of infection (MOI) of five 50% tissue culture infective doses (TCID₅₀)/cell (35, 36).

Transient Knockdown Experiments—Chemically synthesized siRNAs targeting PKR, IPS-1, and TRIF and a control siRNA targeting firefly luciferase (siLUC) were as previously described (24), and were purchased from Dharmacon. For transient knockdown experiments, a double transfection strategy with siRNA was utilized (27). HeLa cells in 60-mm dishes at ~60% confluency were transfected with 50 nM of siRNA with Lipo-fectamine 2000 on day 1 and again on day 3. On day 2, the cells were reseeded in 60-mm dishes for the second siRNA transfection, and on day 4 the cells were seeded in 12-well plates for inducer RNA transfection on day 5.

Double-stranded RNAs and Transfection—Synthetic poly(rI)-poly(rC) was from Sigma. Reovirus genome dsRNA was isolated from purified reovirions as previously described (37). T7 RNA polymerase synthesized 20-bp (Pds20) and 50-bp (Pds50) dsRNAs were made and purified using the Silencer® siRNA construction kit (Ambion) according to the manufacturer's protocol. Further gel purification of the RNAs synthesized in vitro with T7 RNA polymerase was as described by Sambrook et al. (38). The sequence of the Pds20 was the same as the chemically synthesized siLUC; the sense sequence of the T7 transcribed 50-bp Pds50 included this sequence and firefly luciferase sequence 3' of this region, as previously described (24). For RNA transfections, cells were seeded in 24- or 12-well plates and transfected with the indicated amount of RNA complexed with Lipofectamine 2000 in Opti-MEM I.

DNA Plasmids and Transfection—The expression constructs for human eIF-2 α , mutated at serine 51 to either alanine (S51A) or aspartic acid (S51D) in the pMSCV-HA3iresGFP vector, were generously provided by M. Hatzoglou (Case Western University, Cleveland, Ohio). The empty vector (Vec) was generated by deletion of the eIF-2 α cDNA insert from the S51D expression plasmid using EcoRI and XhoI. The eIF-2 α double mutant construct S48,51A was generated by mutating serine 48 to alanine in the S51A plasmid background using the Quick-Change site-directed mutagenesis strategy (Stratagene) and the following primers: forward 5'-GAAGGCATGATTCTTCTT-GCTGAATTGGCCAGAAGGCG-3' and reverse 5'-CGC-CTTCTGGCCAATTCAGCAAGAAGAATCATGCCTTC-3' (mutations are designated by underlined font). Constructs were verified by direct sequencing and restriction enzyme analysis. HeLa cells in 6-well plates were transfected with the indicated eIF-2 α expression plasmid or empty vector (1.6 μ g/well) using Fugene HD transfection reagent (Roche) according to the manufacturer's protocol. After 20 h the transfected cells were in-



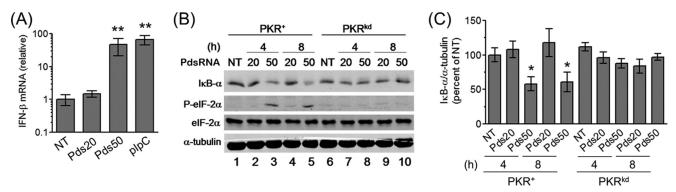


FIGURE 1. dsRNA length-dependent induction of IFN- β correlates with PKR-dependent reduction in I_KB- α protein and eIF-2 α phosphorylation. *A*, relative IFN- β induction measured by quantitative PCR (*qPCR*) using RNA extracted 5 h following incubation with lipid alone (*NT*) or RNA transfection in parental PKR⁺ HeLa cells. IFN- β mRNA levels were normalized to GAPDH using the $\Delta\Delta$ CT method. Standard deviation determined from three independent experiments. *B*, Western blot for I_KB- α , phosphorylated eIF-2 α (P-eIF-2 α) and total eIF-2 α , and α -tubulin loading control, at 4 h and 8 h following transfection of either parental PKR⁺ or knockdown PKR^{kd} HeLa cells with the indicated dsRNA. *C*, quantitation of I_KB- α protein level normalized to α -tubulin and compared with cells not transfected (*NT*). Mean and standard deviation determined from two independent experiments. *, *p* < 0.05 compared with NT and Pds20 at the corresponding time point in PKR⁺ cells, **, *p* < 0.01 compared with NT and Pds20.

fected with C^{ko} virus at a MOI of 1, maintained in DMEM containing 5% (V/V) fetal bovine serum for 24 h, and then RNA was isolated for qPCR analysis or extracts prepared for Western immunoblot analysis.

Cyclohexamide Treatment—For cycloheximide (CHX) (Sigma) treatment prior to protein or RNA isolation, cells were incubated with 10 μ g/ml of CHX as indicated for 4–6 h in Opti-MEM I with or without transfection complexes. In experiments where the time to harvest was greater than 4 h, CHX containing maintenance media was placed on cells following the removal of transfection complexes at 4 h. Cells were then harvested for RNA isolation for qPCR analysis or extract preparation for Western immunoblot analysis.

Real Time PCR Analyses—Total cellular RNA was purified at the indicated time after transfection or infection using an RNeasy mini kit (Qiagen) for 24-well plates or Trizol (Invitrogen) for 12-well plates following the manufacturer's protocol. Random-primed reverse transcription was carried out using 500 ng of RNA and SuperScript II (Invitrogen) according to the manufacturer's protocol. For qPCR reactions, the GAPDH and IFN- β primer pairs were as previously described (27). qPCR reactions were performed in duplicate or triplicate with each RT template, using IQ SYBR Green Supermix (Bio-Rad) and a Bio-Rad MyIQ multicolor real-time qPCR instrument (3-min hot start followed by 30 s at 95 °C, 45 s at 58 °C, 45 s at 72 °C, for 40 to 45 cycles). IFN- β values were normalized to GADPH values.

Western Blot Analyses—Whole cell extracts were prepared with buffer containing protease inhibitors as previously described (39); for measurement of eIF-2 α and PKR phosphorylation, 50 nm NaF and 2 nm Na₂VO₃ (31) or commercial phosphatase inhibitors (Sigma) were also present. Protein fractionation and immunoblot analysis were as described (27, 40) using the following primary antibodies: PKR and I κ B- α (Santa Cruz Biotechnology); phosphorylated-PKR threonine 446 (Epitomics); eIF-2 α (Cell Signaling Technology) and phosphorylatedeIF-2 α serine 51 (Cell Signaling Technology or Epitomics) and α -tubulin (Sigma). Antibody against MV H protein was generously provided by Dr. R. Cattano (Mayo Clinic, Rochester, MN). Blots were visualized and quantified using a Li-Cor Odyssey infrared imager system (Li-Cor Biosciences).

RESULTS

DsRNA Length-dependent Induction of IFN-β and PKR-dependent Reduction of IκB-α—We previously showed that T7 phage RNA polymerase synthesized RNAs (PRNAs) induce IFN-β in a size-dependent manner (24) as also has been shown by others (41, 42). This size-dependent induction is illustrated in Fig. 1*A*, where the 50-bp dsRNA Pds50 induced IFN-β in HeLa cells as measured by qPCR, while the 20-bp dsRNA Pds20 did not induce IFN-β. The long (>100 bp) synthetic dsRNA pIpC, tested as a positive control, induced IFN-β similar to Pds50.

Because NF- κ B is one of the transcription factors required for IFN-β induction and the activation of NF-κB and induction of IFN- β are enhanced by PKR in measles virus-infected and RNA-transfected cells (24, 27, 43) we hypothesized that differential NF-KB signaling may be responsible in part for the differences in IFN- β expression triggered by the different-sized dsRNAs. We previously observed that the inhibitor of NF- κ B signaling, $I\kappa B-\alpha$, was degraded following Pds50 and pIpC transfection (24). To compare this effect with Pds20, we measured the protein levels of $I\kappa B-\alpha$ following either Pds20 or Pds50 transfection in both parental PKR⁺ HeLa cells and the PKR stable knockdown cells. We observed that a decrease in I κ B- α was only seen following transfection with Pds50 and not Pds20, and that this I κ B- α decrease was PKR-dependent (Fig. 1, B and *C*). Furthermore, the decrease in I κ B- α seen in transfected cells correlated with enhanced eIF- 2α phosphorylation (Fig. 1*B*).

Measles Virus C^{ko} Mutant Infection Reduces $I\kappa B - \alpha$ Protein but Increases $I\kappa B - \alpha$ mRNA in a PKR-dependent Manner—Using wild type (WT) measles virus or isogenic mutants lacking either the V (V^{ko}) or C (C^{ko}) viral accessory proteins that impair innate immune system activation (44), we previously established that the C^{ko} virus induces the highest levels of IFN- β (27). We also observed significant PKR activation measured by Thr-446 phosphorylation, but no change in total PKR level following C^{ko} virus infection, and we showed that maximal induc-



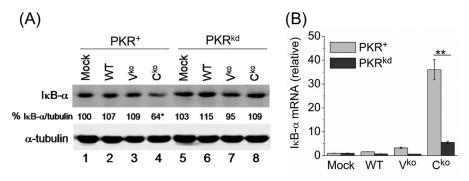


FIGURE 2. **PKR-dependent reduction in** I κ **B**- α **protein level inversely correlates with** I κ **B**- α **mRNA level following MV infection.** *A*, Western blot for I κ B- α protein following mock, WT, V^{ko}, and C^{ko} measles virus infection of PKR⁺ and PKR^{kd} cells. I κ B- α protein levels normalized to α -tubulin in each lane are expressed as a percentage of the mock infected lane. *B*, I κ B- α mRNA transcript levels measured by qPCR at 24 h postinfection in PKR⁺ and PKR^{kd} cells as described in Fig. 1. Mean and standard deviation are shown from two independent experiments analyzed in triplicate. *, *p* < 0.05 compared with mock treatment in PKR⁺ cells, **, *p* < 0.01.

tion of IFN- β required PKR and IPS-1. Using gel shift and NF- κ B luciferase reporter assays, we demonstrated that the measles virus C^{ko} mutant activated NF- κ B signaling, and that maximal NF- κ B activation was dependent upon both PKR and IPS-1 (27).

To test whether the PKR-dependent decrease in $I\kappa B-\alpha$ protein that we observed following dsRNA transfection (Fig. 1) also occurred in the context of viral infection, we examined the $I\kappa B-\alpha$ level in cells infected with WT, V^{ko} or C^{ko} measles virus at the 24 h time point where we previously observed maximal IFN- β induction, PKR phosphorylation, and NF- κB activation (27, 33). Following infection, C^{ko} virus-infected cells displayed a lower $I\kappa B-\alpha$ protein level than uninfected cells or cells infected with either the WT or V^{ko} virus in a PKR-dependent manner (Fig. 2*A*).

We next examined $I\kappa B-\alpha$ RNA levels by gPCR in order to determine the relationship between the levels of I κ B- α protein and I κ B- α mRNA, and to assess whether the reduction in I κ B- α protein was because of a reduction in $I\kappa B-\alpha$ RNA. While untreated and WT virus-infected cells showed no significant change in I κ B- α mRNA transcript levels and the V^{ko} virus only slightly increased I κ B- α mRNA in the parental PKR⁺ cells, the C^{ko} virus significantly increased I κ B- α mRNA levels in a PKRdependent manner (Fig. 2*B*). I κ B- α protein levels were lowest in PKR⁺ cells following C^{ko} virus infection even though the IκB- α mRNA level was ~7-fold higher in the PKR⁺ cells than PKR^{kd} cells. The enhanced induction of I κ B- α mRNA in PKR⁺ cells was not unexpected, because NF-kB is known to induce IκB- α (30) and, as described above, PKR⁺ cells display enhanced NF- κ B signaling. These results suggest that PKR either enhanced I κ B- α protein degradation or the inhibition of IκB- α protein synthesis.

PKR-dependent Decrease of IκB-α Protein and Increase of IκB-α mRNA Occurs following Transfection with Naturally Occurring dsRNA—We next examined the PKR dependence of decreased I*κ*B-*α* protein and the relationship between the level of I*κ*B-*α* mRNA and I*κ*B-*α* protein in PKR⁺ and PKR^{kd} cells transfected with purified reovirus genome RNA. Reovirus genome is composed of ten segments of dsRNA of lengths varying from 1.2 to 3.8 kbp (45), and is not known to include inosine as is present in the synthetic pIpC dsRNA. Reovirus dsRNA activates PKR (37) and both the RIG-I and MDA5 cytosolic

RNA signaling pathways which utilize the IPS-1 adaptor protein (42). Following reovirus RNA transfection, we found a significant reduction in I κ B- α protein only in the PKR⁺ cells and not in the PKR^{kd} cells (Fig. 3*A*). This reduction was observed between 2 and 12 h after transfection and was seen only in PKR⁺ cells where enhanced eIF-2 α phosphorylation was observed (Fig. 3*A*).

Following transfection with either pIpC or reovirus genome dsRNA, the level of I κ B- α mRNA was highest in PKR⁺ cells and was reduced significantly in PKR^{kd} cells (Fig. 3*B*), just the opposite of what we observed for $I\kappa B-\alpha$ protein levels (Fig. 3A). We next confirmed this result obtained with PKR^{kd} cells stably deficient in PKR by using cells transiently knocked down for PKR and observed a similar reduction in I κ B- α mRNA was seen following reovirus genome RNA transfection (Fig. 3C). We also transiently knocked down IPS-1 to test whether $I\kappa B-\alpha$ mRNA induction was reduced, since IPS-1-dependent signaling has been shown to activate the IKK complex through the NF-KB modulator (NEMO, also known as $I\kappa K-\gamma$) protein (46). IPS-1 knockdown but not TRIF knockdown also reduced I κ B- α mRNA induction in response to transfected reovirus genome RNA (Fig. 3C), suggesting that NF-κB signaling is activated, or at least enhanced, by PKR and IPS-1-dependent responses.

Inhibition of Translation by Cycloheximide in PKR^{kd} Cells Reduces IKB- α Protein Levels to That Seen in PKR⁺ Cells following dsRNA Transfection—The mechanism by which PKR enhances NF-KB activation (27) and NF-KB-dependent gene induction (Figs. 2 and 3) could involve the direct activation of NF-*k*B signaling components. Alternatively, an indirect translational control mechanism also may be operative in which the activity of NF-KB signaling would be affected through a PKRdependent eIF-2 α phosphorylation and subsequent inhibition of synthesis of negative regulatory proteins such as $I\kappa B-\alpha$. For the latter explanation, $I\kappa B-\alpha$ protein degradation could be initially similar in PKR⁺ and PKR^{kd} cells (47), but subsequent synthesis of I κ B- α protein would be reduced in PKR⁺ cells due to an inhibition of $I\kappa B-\alpha$ mRNA translation compared with that in PKR^{kd} cells. The observed disparity between $I\kappa B-\alpha$ mRNA and protein levels (Figs. 2 and 3) is consistent with the notion that PKR may enhance activation of NF-κB signaling through a translational control mechanism.



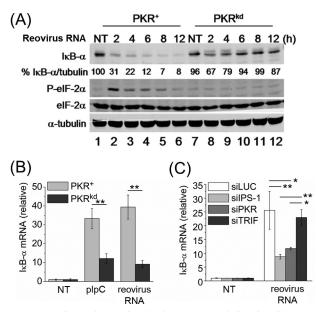


FIGURE 3. PKR-dependent reduction in I κ B- α protein level and increase in I κ B- α mRNA level following transfection with reovirus genome dsRNA. *A*, Western blot for I κ B- α , phosphorylated eIF-2 α (P-eIF-2 α) and total eIF-2 α , and α -tubulin loading control, at the indicated times after transfection of PKR⁺ or PKR^{kd} cells with purified reovirus genome dsRNA. Percentage (%) indicates I κ B- α protein level normalized to α -tubulin and compared with cells not transfected. *B*, I κ B- α mRNA transcript level measured by qPCR with RNA isolated from PKR⁺ and PKR^{kd} cells at 5 h following transfection with reovirus genome dsRNA or plpC. Error bars represent the standard deviation determined from three independent experiments. *C*, qPCR measurement of I κ B- α mRNA transcript level following transfection with reovirus genome dsRNA of HeLa cells transiently knocked down for PKR, IPS-1 or TRIF or transfected with a control siRNA (*siLUC*). Standard deviations determined from two independent experiments analyzed in triplicate. *, p < 0.05; **, p < 0.01.

To test the possibility that PKR decreases $I\kappa B-\alpha$ protein through translational control thereby leading to the increase seen in $I\kappa B-\alpha$ mRNA (Fig. 3), we treated PKR^{kd} cells with cycloheximide (CHX), a drug that prevents the elongation step of translation (48), and then examined $I\kappa B-\alpha$ protein levels. In PKR^{kd} cells, CHX treatment significantly decreased $I\kappa B-\alpha$ protein levels (Fig. 4, *lanes 11* and *12*) to a level comparable to that of PKR⁺ cells, suggesting that PKR indeed might contribute to NF- κ B induction solely by preventing the translation of $I\kappa B-\alpha$ mRNA (47). No changes in eIF- 2α phosphorylation were observed in PKR^{kd} cells following CHX treatment.

Cycloheximide Treatment Rescues IFN- β RNA Induction Following PKR-knockdown, but Does Not Rescue Following IPS-1 Knockdown—Most importantly, CHX treatment rescued IFN- β RNA induction in PKR^{kd} cells to a significant extent (p < 0.01) in response to transfected synthetic (pIpC) or natural (reovirus genome) dsRNA (Fig. 5A), indicating that inhibition of synthesis of inhibitors of NF- κ B signaling was a likely explanation for the PKR dependence of IFN- β induction. CHX treatment also restored I κ B- α RNA induction in PKR^{kd} cells to levels found in PKR⁺ cells (Fig. 5B).

While our results are consistent with an amplification of IFN- β RNA induction by PKR-mediated translation inhibition of I κ B- α protein synthesis, they do not clearly differentiate between the IPS-1 and PKR contributions to IFN- β signaling. To test whether CHX treatment specifically rescues PKR-dependent but not IPS-1-dependent induction of IFN- β transcripts, we transiently knocked down either PKR or IPS-1 in

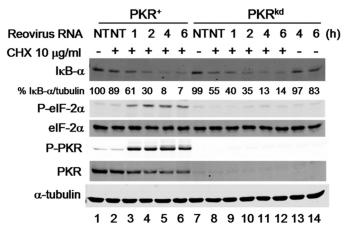


FIGURE 4. **Translation inhibition by cycloheximide prevents IkB**- α **protein expression in PKR**^{kd} **cells.** PKR⁺ and PKR^{kd} cells were transfected with reovirus genome dsRNA for the indicated period of time (*h*) or not transfected (*NT*), and were treated (+) with cycloheximide (*CHX*) or left untreated (-) as indicated. Western immunoblot analyses were performed for IkB- α , phosphorylated elF- 2α (P-elF- 2α), total elF- 2α , phosphorylated PKR (P-PKR), and total PKR, and α -tubulin as a loading control. Percentage (%) is calculated for IkB- α protein level normalized to α -tubulin, and compared with PKR⁺ cells, transfected with reavirus genome dsRNA but not treated with CHX, are shown in *lanes 13* and *14* for comparison.

parental HeLa cells. As we earlier reported (24), transient knockdown of either IPS-1 or PKR significantly reduced IFN- β induction (Fig. 5*C*). Following CHX treatment, however, IFN- β RNA induction increased to near control levels in the PKR transient knockdown cells, but remained significantly reduced in the IPS-1 knockdown cells (Fig. 5*C*). These results suggest that the translation inhibition-dependent contribution of PKR to IFN- β induction is independent of the activation of signaling by IPS-1.

Non-phosphorylatable Mutant eIF-2 α Reduces IFN- β RNA Induction and Increases I κ B- α Protein Levels Following Infection of PKR-sufficient Cells—To further test whether the PKRdependent enhancement of IFN- β transcript induction was translationally mediated through eIF-2 α , the effect of S48,51A, an unphosphorylatable mutant form of eIF-2 α in which the serine phosphorylation sites were replaced with alanine, was examined. As shown in Fig. 6, the phosphorylation defective S48,51A mutant resulted in lower levels of both IFN- β transcript (Fig. 6A) and I κ B- α transcript (Fig. 6B) following C^{ko} measles virus infection compared with either vector-transfected cells or cells expressing the phospho-mimetic S51D mutant. IFN- β transcript levels in the absence of the C^{ko} infection were comparably low in the vector, S51D and S48,51Atransfected cells (Fig. 6).

Expression of the S48,51A phosphorylation defective mutant form of eIF-2 α in PKR⁺ parental cells also led to enhanced I κ B- α protein production in C^{ko} virus-infected cells (Fig. 7*A*, *lane 6*), reaching a level of I κ B- α comparable to that seen in C^{ko}-infected PKR^{kd} cells (Fig. 7*A*, *lane 8*). By contrast, the level of I κ B- α in C^{ko}-infected vector- or S51D-transfected cells was 2–3-fold lower (Fig. 7*B*). Interestingly, PKR⁺ cells expressing the phosphorylation defective S48,51A mutant also gave rise to increased viral H protein synthesis, as was observed in PKR^{kd} cells, compared with the vector control or S51D-expressing cells (Fig. 7*A*).



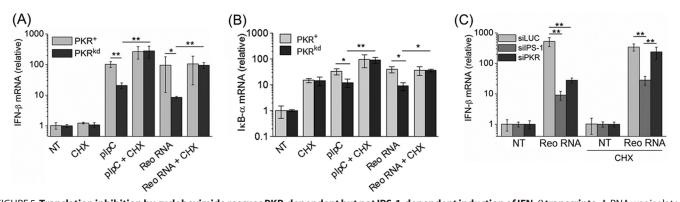


FIGURE 5. **Translation inhibition by cycloheximide rescues PKR-dependent but not IPS-1-dependent induction of IFN-** β **transcripts.** *A*, RNA was isolated from PKR⁺ or PKR^{kd} cells at 5 h after transfection with plpC or reovirus genome dsRNA in the presence or absence of CHX as indicated and the relative IFN- β mRNA transcript level was determined by qPCR as described in Fig. 1. The standard deviation was determined from two or three independent experiments. *B*, qPCR measurement of relative IKB- α mRNA transcript levels following treatment conditions described in *A*. C, relative IFN- β mRNA transcript level in parental PKR⁺ HeLa cells transiently knocked down for either PKR or IPS-1, or transfected with a control siLUC siRNA prior to transfection of reovirus genome dsRNA in the absence or presence of CHX. Standard deviation is shown for two independent experiments. ***, *p* < 0.05; **, *p* < 0.01.

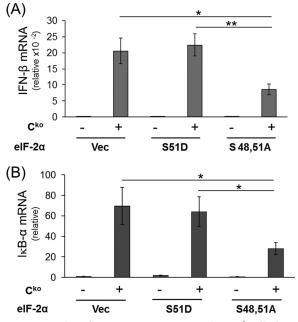


FIGURE 6. Expression of mutant eIF-2 α (S48,51A) in PKR⁺ cells decreases IFN- β induction by measles virus. Cells (PKR⁺) were transfected with either an empty vector (*Vec*) or an expression construct encoding human eIF-2 α mutated to prevent phosphorylation (S48,51A) or a phospho-mimetic (S51D) mutant. At 20 h after transfection, cells were either left uninfected (–) or infected (+) with MV C^{ko} virus, and at 24 h after infection total RNA was prepared and transcript levels determined by qPCR. The results shown are normalized to GAPDH transcript level and are relative to uninfected, vector-transfected cells. The mean induction and standard deviation were determined from three independent experiments. *, p < 0.05; **, p < 0.01. A, IFN- β transcript levels, B

DISCUSSION

PKR has emerged as an important modulator of IFN- β gene induction following infection with RNA viruses, including negative-stranded (27, 43), positive-stranded (49–53) and doublestranded (54) RNA viruses. But, the molecular mechanism by which PKR affects IFN- β induction largely remains unresolved. The objective of our study was to determine the mechanistic basis by which the PKR kinase enhances IFN- β transcript RNA levels following virus infection and RNA transfection. The results reported herein suggest that a translational control mechanism through eIF- 2α is likely sufficient to account for the

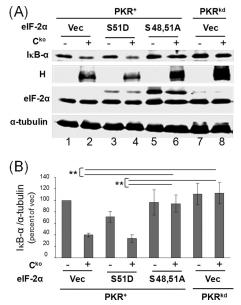


FIGURE 7. Expression of mutant eIF-2 α (S48,51A) increases I κ B- α protein levels in C^{ko} measles virus infected PKR⁺ cells to levels seen in PKR^{kd}infected cells. *A*, PKR⁺ or PKR^{kd} cells were transfected with either an empty vector (Vec) or an expression construct encoding either S48,51A or S51D mutant eIF-2 α as indicated. At 20 h after transfection cells were either left uninfected (–) or infected (+) with MV C^{ko} virus, and at 24 h after infection extracts were prepared and western immunoblot analyses performed using antibodies against I κ B- α , MV H protein and eIF-2 α , and α -tubulin as a loading control. *B*, quantitation of I κ B- α protein levels normalized to α -tubulin; average and standard deviation determined from three independent analyses; **, *p* < 0.01.

PKR-dependent reduction in I κ B- α levels and enhancement of IFN- β induction. In fact, translational control may be broadly important and operative in multiple pathways of IFN signaling as the translational repressors 4E-BP1 and 4E-BP2 also have been shown to negatively regulate the type-I IFN response by inhibiting IRF7 mRNA translation during viral infection (55).

Impaired IFN- β transcript induction in HeLa cells stably knocked down for PKR is known following dsRNA transfection (24) or measles virus infection (27, 43). Using a transient knockdown strategy we earlier had established that, in addition to PKR, the IPS-1 adaptor protein for the RLR cytosolic signaling pathway was required for maximal IFN- β induction by measles



virus deficient in C protein, but that the TRIF mediator of TLR3 signaling was not involved. Because both PKR and IPS-1 were necessary for maximal IFN- β transcript induction in response to either dsRNA transfection or MV infection (24, 27), we attempted to further delineate and distinguish the requirements for these proteins. Short structured RNAs without a 5'-triphosphate are known to be weak inducers of IFN- β (42), but they still activate PKR (24, 37), suggesting that PKR activation is not sufficient for IFN- β induction. Although the activation of IRF3 is PKR-independent in both RNA transfected and measles virus infected HeLa cells, the activation of NF- κ B is enhanced by PKR (27, 43).

Negative regulators of NF-kB signaling, exemplified by IκB- α , are known to be degraded in response to an NF- κ Bactivating stimuli, and then resynthesized to subsequently inhibit NF- κ B activity (29, 47). Hence, we considered that translation inhibition mediated by PKR might lead to enhanced NF- κ B signaling and IFN- β induction. As shown herein, an inverse correlation was found between I κ B- α mRNA and protein levels following dsRNA transfection and virus infection. Furthermore, consistent with previous results (24), we observed an inverse correlation between eIF-2 α phosphorylation and I κ B- α protein levels following these treatments. The steady-state protein expression level of $I\kappa B-\alpha$ was increased and largely rescued when the S49,51A mutant of eIF-2 α was expressed, which suggests that the resynthesis rather than the stability of I κ B- α is the key parameter affected by PKR activation.

As a further test of the translational basis of the PKR-dependent effect, we used a pharmacologic inhibitor, CHX, to inhibit protein synthesis. Importantly, CHX treatment during RNA transfection rescued PKR-dependent, but not IPS-1-dependent, IFN- β transcriptional induction. This finding, together with our observation that the transient knockdown of either PKR or IPS-1 impaired IFN- β transcript induction, suggests that PKR and IPS-1 differentially function in the signaling response leading to activation of IFN- β transcription.

Our results furthermore demonstrate that the effect of PKR on signaling pathway responses following RNA transfection or virus infection is dependent on PKR enzymatic activity through its eIF- 2α kinase activity (40, 56) and hence cannot represent simply a protein adaptor or scaffold role for PKR (22, 57). The observed PKR-dependent decrease in I κ B- α following infection or transfection is consistent with results obtained for the endoplasmic reticulum stress-responsive eIF- 2α kinase, PERK, where maximal NF- κ B signaling following phosphorylation of eIF- 2α by PERK likewise correlated with decreased protein levels of I κ B- α (56).

Our results taken together support the role of translation control of negative regulators, including $I\kappa B-\alpha$, in the modulation of NF- κ B signaling. Similar to the effects of PKR on $I\kappa B-\alpha$ expression established herein, it is conceivable that other negative regulators of the NF- κ B pathway in addition to $I\kappa B-\alpha$ also are modulated in a PKR-dependent manner. One intriguing possibility is the A20 negative regulatory protein, which prevents NF- κ B essential modulator (NEMO) activation through its ubiquitin-editing activity (58). Because A20 is induced by NF- κ B signaling in a manner similar to $I\kappa B-\alpha$ resulting in inhibition of NF- κ B signaling upstream of I κ B- α (59), it is reasonable to speculate that A20 would represent an additional potential target for PKR-dependent translational control of NF- κ B signaling. Since A20 regulates NEMO activity, and NEMO activity is required for the activation of IRF3 through TBK1 and I κ K (60), then PKR-dependent inhibition of A20 translation presumably could also affect IRF activation in some instances, perhaps explaining the PKR-dependent activation of IRF3 observed following mutant vaccinia virus infection (32). In fact, the absence of negative regulators of NF- κ B signaling may also explain the PKR-dependence of I κ B phosphorylation and activation of I κ K shown by others (20, 21). It is also possible that cell type differences in NF- κ B signaling, especially in cancer cells (61), may account for the seemingly different effects of PKR on this pathway.

Our results are consistent with a translational inhibition response through eIF-2 α phosphorylation for the mechanistic basis of the PKR-dependent amplification of IFN-β expression following transfection or infection (40, 56). First, in PKR-sufficient PKR⁺ cells as shown herein, a mutant form of eIF-2 α that cannot be phosphorylated impaired measles virus-induced activation of IFN- β gene expression and enhanced I κ B- α protein production following infection. Secondly, in PKR-deficient cells, catalytically active PKR is necessary in order to complement measles virus-induced PKR-dependent IFN-β transcript induction (40). Our findings, taken together, reveal that PKR functions at least in part through an eIF-2-mediated translational control mechanism to indirectly enhance IFN-β transcript levels, and that IPS-1 functions through direct activation of signaling pathways including initial NF-KB and IRF3 activation as established by others (5).

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