Diversity in Viral Anti-PKR Mechanisms: A Remarkable Case of Evolutionary Convergence

Elena Domingo-Gil¹⁹, René Toribio¹⁹, José Luis Nájera², Mariano Esteban², Iván Ventoso¹*

1 Departamento de Biología Molecular and Centro de Biología Molecular "Severo Ochoa" (CSIC-UAM), Universidad Autónoma de Madrid, Madrid, Spain, 2 Centro Nacional de Biotecnología (CSIC), Campus Universidad Autónoma de Madrid, Madrid, Spain

Abstract

Most viruses express during infection products that prevent or neutralize the effect of the host dsRNA activated protein kinase (PKR). Translation of Sindbis virus (SINV) mRNA escapes to PKR activation and eIF2 phosphorylation in infected cells by a mechanism that requires a stem loop structure in viral 26S mRNA termed DLP to initiate translation in the absence of functional eIF2. Unlike the rest of viruses tested, we found that Alphavirus infection allowed a strong PKR activation and eIF2 α phosphorylation *in vitro* and in infected animals so that the presence of DLP structure in mRNA was critical for translation and replication of SINV. Interestingly, infection of MEFs with some viruses that express PKR inhibitors prevented eIF2 α phosphorylation after superinfection with SINV, suggesting that viral anti-PKR mechanisms could be exchangeable. Thus, translation of SINV mutant lacking the DLP structure (Δ DLP) in 26S mRNA was partially rescued in cells expressing vaccinia virus (VV) E3 protein, a known inhibitor of PKR. This case of heterotypic complementation among evolutionary distant viruses confirmed experimentally a remarkable case of convergent evolution in viral anti-PKR mechanisms. Our data reinforce the critical role of PKR in regulating virus-host interaction and reveal the versatility of viruses to find different solutions to solve the same conflict.

Citation: Domingo-Gil E, Toribio R, Nájera JL, Esteban M, Ventoso I (2011) Diversity in Viral Anti-PKR Mechanisms: A Remarkable Case of Evolutionary Convergence. PLoS ONE 6(2): e16711. doi:10.1371/journal.pone.0016711

Editor: Jean-Pierre Vartanian, Institut Pasteur, France

Received August 12, 2010; Accepted January 11, 2011; Published February 2, 2011

Copyright: © 2011 Domingo-Gil et al. This is an open-access article distributed under the terms of the Creative Commons Attribution License, which permits unrestricted use, distribution, and reproduction in any medium, provided the original author and source are credited.

Funding: This work was supported in part from the VIRUS-HOST interaction programme (Comunidad de Madrid) and by grants from the Ministerio de Ciencia e Innovación (SAF2006-09810; SAF2008-02036) and the Fundación Mutua Madrileña (FMM 2008). The institutional support from Fundación Ramón Areces is also acknowledged. E.D was a recipient of VIRUS-HOST programme postdoctoral contract. R.T. was a recipient of the SAF2006-09810 contract and I.V. was a researcher of Ramón y Cajal Programme. The elaboration of this work took about 2 years and the estimated cost was 3000 euros excluding salaries. The funders had no role in study design, data collection and analysis, decision to publish, or preparation of the manuscript.

Competing Interests: The authors have declared that no competing interests exist.

* E-mail: iventoso@cbm.uam.es

9 These authors contributed equally to this work.

Introduction

During virus-host coevolution, the acquisition of an antiviral pathway by the host was generally followed by the appearance of a viral countermeasure so that the current host-parasite interactions are keeping on a dynamic equilibrium (the Red Queen principle) [1,2,3]. A paradigmatic example of these complex interactions are the different mechanisms that viruses have evolved to evade or subvert the antiviral effect of interferons (IFNs) and other proinflammatory cytokines that are secreted by vertebrate cells in response to virus and other pathogens [4,5,6,7,8,9]. Thus, viruses express products that impair the detection of viral proteins or nucleic acids by host pattern-recognition receptors (PRRs; Tolllike receptors and RIG-I-like receptors), block the signaling pathways that lead to the synthesis of IFN α/β and other cytokines, or prevent the activation of some IFN-stimulated genes such as dsRNA-activated kinase (PKR) [6,9,10,11,12,13,14]. Moreover, in some cases such as poxvirus or HCV, one or more viral product can interfere with IFN secretion or signalling at multiple points, ensuring a more efficient viral escape to innate immunity of the host [6,9].

Among innate antiviral mechanisms of vertebrates, PKR activation constitutes one of the first line of antiviral defense acting at the immediate-early phase of virus replication that

precedes the eventual secretion of IFN. PKR is present at basal levels in most of mammalian tissues, but its amount increases after priming cells with IFN α/β [15,16]. PKR binds dsRNA molecules generated during the replication of RNA viruses as well as in some transcripts from DNA viruses, leading to the activation of the kinase by a sequential wave of autophosphorylation events [17,18,19,20]. Activated PKR phosphorylates and inactivates eukaryotic initiation factor 2 (eIF2), the only well-described substrate of the kinase that is also phosphorylated by other members of eIF2-kinase family in vertebrates [17,21]. As result of this, the general translation is rapidly inhibited in an attempt of the infected cells to block viral translation and abort virus spreading [16,22]. However, as in other pathways of innate response, viruses have evolved a variety of strategies to prevent or overcome the activation of PKR in infected cells (reviewed in [12]). Among these mechanisms, the most frequent found are viral products that prevent the activation of PKR by sequestering its activator (dsRNA), by direct binding to the regulatory element of the kinase or by expressing a pseudosubstrate that competes with eIF2 for binding to the kinase (see figure 1). Other viruses such as poliovirus and Rift Valley virus induce the degradation of PKR by a mechanism that has not been well characterized yet [23,24]. In some cases such as Herpes virus-infected cells, eIF2 phosphorylation is rapidly reversed by the action of viral phosphatases that



Figure 1. Current known viral strategies to prevent or counteract PKR activation can be grouped into three categories. *A*) First, viruses can prevent the activation of PKR by expressing proteins or RNAs that bind the kinase in an inhibitory manner, by sequestering dsRNA activator molecules or by inducing the degradation of PKR. Examples of this group are vaccinia E3 and K3, influenza NS1 and Adenovirus VAI RNAs and Rift Valley virus NS1 protein. In the second category are viruses that promote elF2 α dephosphorylation such as Herpes virus γ 34.5 protein which acts as a regulatory subunit of cellular protein phosphatase 1a (PP1). Herpesvirus also express products from the first group (Us11 gene, a PKR inhibitor). In the third category, some viruses do not prevent PKR-mediated elF2 α phosphorylation, but viral mRNA can initiate translation in an elF2-independent manner by means of a RNA structure (DLP) that stalls the scanning 40S ribosome on the initiation codon. To date, this strategy has been found only in Alphavirus. *B*) Secondary structure prediction of the 5' extreme of 26S mRNAs of SINV and SFV. The stability of DLP structure (ΔG°) is expressed in kcal. mol⁻¹. Arrows show the initiation codon. Circled nucleotides in SINV were mutated to adenines in Δ DLP mutant virus as described previously [29]. doi:10.1371/journal.pone.0016711.g001

are expressed along the infection [25,26,27]. For Alphavirus, the strategy is markedly different; PKR is strongly activated upon infection with SINV and Semliki Forest (SFV) viruses, so that eIF2 factor is completely phosphorylated [28,29]. Translation of viral subgenomic mRNAs (26S), however, resists due to the presence of a prominent hairpin loop structure in mRNA located downstream of the initiation codon (DLP) that allows the 40S ribosome to initiate in the absence of eIF2 [29].

The existence of these different mechanism stress the critical role of PKR in regulating host-virus interaction and show a remarkable example of convergent evolution to generate different patterns of molecular mimicry [8]. For example, VV K3L mimics eIF2 α acting as a competitive inhibitor of PKR, whereas Adenovirus VAI RNAs mimics dsRNA activators of PKR by binding the kinase in an inhibitory manner [8,12]. Fingerprints of this antagonist coevolution has been found recently in the

accelerated rates of positive evolution found in PKR gene for defeating viral anti-PKR products such as poxvirus E3 and K3 [30,31]. However, an important issue that has not been addressed to date is whether these different anti-PKR mechanisms might be exchanged between non related viruses to get a functional complementation. In this work we show that a cis-acting RNA structure of Alphaviruses responsible for counteracting PKR activation can be replaced by a viral protein such as vaccinia E3.

Results

Alphavirus are unique in allowing PKR activation in cultured cells and infected animals

We and others have reported before that infection of 3T3 fibroblasts with SINV virus triggered the complete phosphorylation of eIF2 due to a strong PKR activation [28,29] (Figure 2). We wanted to test whether such phosphorylation of host $eIF2\alpha$ was also detected in cells infected with other Alphaviruses and with members of other unrelated families of RNA viruses. For this, we infected cells with SFV (Alphavirus), VSV (Rhabdovirus), Influenza A (Orthomyxovirus) and EMCV (Picornavirus) (Figure 2A). Equivalent or even stronger PKR activation was observed in cells infected with SFV as judged by the mobility shift of PKR band in SDS-PAGE that was indicative of autophosphorylation associated to kinase activation (Figure 2B). As for SINV, activation of PKR in SFV-infected cells induced a strong phosphorylation of eIF2 α that was not observed in the rest of viruses analyzed with the exception of VSV, where a slight increase in eIF2 phosphorylation was observed as reported before [16]. The use of PKR gene knock-out cells $(PKR^{o/o})$ showed that eIF2 phosphorylation in Alphavirus-infected cells was almost completely attributable to PKR activation as described before (Figure 2B and [29]. With the exception of Alphavirus, the rest of RNA viruses tested here did not trigger a substantial activation of eIF2 kinases in infected cells.

Next, we analyzed whether eIF2a phosphorylation was also detected in animals infected with Alphavirus. For this, mice were infected with SINV by the intranasal route, and 4 days later brains were subjected to immunofluorescence (IF) analysis. As described previously, SINV showed a marked tropism for cortical and spinal cord neurons [32,33,34,35]. Groups of neurons expressing viral antigens were easily detected in pyriform, motor and somatosensorial areas of brain cortex as well in hippocampus. Notably, the vast majority of these infected cells showed a strong staining of phospho eIF2 (Figure 3A), with no signal detected outside of areas of viral replication. Clearly, no eIF2 phosphorylation was detected in infected neurons of $PKR^{0/0}$ mice, showing that as occurred in vitro PKR was responsible for eIF2 phosphorylation in response to infection of animals (Figure 3A). We extended this in vivo analysis to other viruses endowed with anti-PKR mechanisms such as VV. Clearly, no eIF2 phosphorylation was detected in spleens of mice infected with VV despite the efficient replication detected in this organ (Figure 3B). The absence of eIF2 phosphorylation in mice infected with VV was not attributable to a low expression of the kinase in spleen, since a comparable expression of PKR was detected in all murine organs analyzed (Figure 3C). Taken together, our results show that PKR-induced eIF2 phosphorylation seems to be a specific feature of animals infected with Alphavirus.

Resistance to PKR activation is critical for replication of Alphavirus in animals

Viral anti-PKR products commonly act as virulence factors necesary for efficient viral replication and pathogenesis in animals [25,36,37,38]. To test whether translational resistance of SINV virus to PKR activation was important for replication *in vivo*, we compared the replication of wild type virus and a SINV mutant lacking the DLP structure (Δ DLP) in 26S mRNA. This structure in RNA has been reported before to be essential for translation and virus replication in 3T3 cells [29]. Clearly, replication of Δ DLP virus was greatly hampered in wild type, but not in *PKR*^{6/o} mice (Figure 3D). About 3–4 log reduction in viral yield of Δ DLP virus were found in *PKR*^{+/+} mice (wild type) when compared to *PKR*^{6/o} counterparts, whereas wild type viruses replicated at similar levels in both animal types. This result shows that as predicted from other viruses, PKR resistance in Alphavirus is essential for efficient replication *in vivo*.

Rescue of SINV Δ DLP translation by poxvirus E3 protein

In a first attempt to test whether anti-PKR mechanisms could be exchangeable among viruses, we analyzed the phosphorylation state of eIF2 in SINV-infected cells that had been first infected with low doses of some RNA (Influenza A, VSV and EMVC) and DNA (VV) viruses (Figure 4A). Time-lag between infections was set to allow the simultaneous expression of products from both viruses. Thus, the previous infection with the indicated viruses did not prevent the replication and accumulation of SINV proteins in doubly infected cells (Figure 4A). Interestingly, pre-infection with VV and EMCV viruses completely prevented eIF2 phosphorylation triggered by SINV superinfection. However, other viruses such as VSV were unable to prevent PKR activation after superinfection with SINV, whereas Influenza A infection exerted only a moderate preventive effect (${\sim}50\%$ reduction). In this experiment we also included a VV mutant lacking the E3L gene, the main PKR inhibitor in VV-infected cells [39,40,41]. Notably, mutant VV- Δ E3L was unable to prevent eIF2 α phosphorylation after SINV superinfection, showing that the inhibitory potential of VV on PKR is largely attributable to the action of E3 protein. This trans-inhibitory activity of E3 opened the possibility to rescue translation of SINV Δ DLP mutant by expressing VV E3L gene in MEF cells. For this, we used a previous established MEF line that expressed the VV E3 protein in an inducible manner after tetracycline withdrawal [42]. As is shown is Figure 4B, induction of E3L gene partially prevented eIF2 phosphorylation after SINV infection. Moreover, mobility shift of PKR in SDS-PAGE was prevented by expression of E3 protein, showing that this viral product targeted PKR to block its activation. Notably, expression of E3 protein partially restored translation of Δ DLP virus mRNA that was strongly inhibited due to eIF2 phosphorylation in non induced cells. This allowed a \sim 50 fold increase of Δ DLP virus yield in induced cells, showing that a deficiency in a cis-acting RNA structure involved in translation can be complemented by an unrelated product but that participates in the same biological trait.

Discussion

Adaptation of non-related parasites to the same host or ecological niche is generally associated to events of convergent evolution (homoplasy), where analogous genetic elements arise during the acquisition of the same biological trait [8,43]. A paradigmatic example of this is the plethora of viral mechanisms to counteract the innate response of vertebrates (reviewed in [6]). Collectively, viruses have acquired mechanisms to subvert or counteract virtually all the biochemical routes of the host that lead into the production of IFN and other antiviral cytokines (reviewed in [6,8]). Among them, the appearance of mechanisms to prevent or counteract PKR activation likely increased the replication capabilities of viruses during their adaptation to vertebrate hosts.





Figure 2. Alphaviruses are unique in allowing PKR activation and elF2 α **phosphorylation in infected cells.** *A*) Susceptibility of MEFs to viruses used in this study. Cells were infected with the indicated viruses to a moi of 25 pfu/cell and 6 h later pulsed with [³⁵S]-Met for 30 min. Labeled proteins were resolved by SDS-PAGE followed of autoradiography. *B*) PKR activation and elF2 α phosphorylation in wild type or *PKR*^{0/o} MEFs infected with the indicated viruses at 4 and 6 hpi. Note the mobility shift of PKR band upon activation in SINV and SFV-infected cells (upper panel). elF2 α phosphorylation in wild type (+/+) and *PKR* knock-out cells (o/o) infected with the indicated viruses. Only Alphavirus-infected cultures showed a strong elF2 α phosphorylation. For VSV-infected cells, a slight increase in elF2 α phosphorylation was also observed. doi:10.1371/journal.pone.0016711.g002



Figure 3. eIF2*a* **phosphorylation and translational resistance of SINV virus also operates in infected animals.** *A*) IF analysis of brains from wild type and *PKR* knock-out mice infected with SINV at 4 dpi. Adjacent sections were incubated with anti-SINV and anti-phosphoeIF2*a* antibodies as described in Materials and Methods. *B*) IF analysis of spleens from mice infected with VV-Luc at 1dpi. Sections were incubated with anti-VVp14 (reactive against the envelope protein A27) and anti-phosphoeIF2*a* antibodies. Note that spleen cells expressing viral antigens did not react with anti-phosphoeIF2*a* antibodies. *C*) Expression of *PKR* in different mouse organs from uninfected animals. Equivalent amounts of protein extracts were analyzed by immunoblot against PKR, total eIF2*a* and β-actin. *D*) Attenuation of ΔDLP mutant virus in wild type, but not in *PKR*^{o/o} mice. Animals were inoculated with 10⁷ of WT and 2×10⁷ of ΔDLP mutant viruses. Viral yields in mouse brains at 4 dpi were titrated by plaque assay. Results are the mean from 10 animals inoculated for each group in three independent experiments. SD from each group is also showed. doi:10.1371/journal.pone.0016711.q003

Thus, the type of anti-PKR mechanism found in a given virus seems to be greatly influenced by the genomic organization and the natural history of the virus during the adaptation to vertebrate hosts. Thus, the singular mechanism of PKR resistance found in Alphavirus probably reflects a peculiar evolutionary adaptation to vertebrates from arthropods (mainly mosquitoes), the vectors that Alphavirus and other Arbovirus use for transmission [35,44,45]. Moreover, the appearance of cis-acting RNA structures such as DLP to counteract PKR-mediated eIF2a phosphorylation was probably limited to simple monocistronic RNA virus such as Alphavirus, whereas in polycistronic RNA viruses (e.g. Influenza A) and in complex DNA viruses (e.g. Poxvirus) the acquisition of trans-acting PKR inhibitors allowed viral mRNAs to be collectively translated in infected cells. Although being functional equivalent, viral anti-PKR mechanisms that prevent or allow PKR activation could have different biological consequences. The strong eIF2 α phosphorylation found in neurons of SINV-infected mice is enough to completely block translation of non-viral mRNAs throughout the infection [46], that probably leads to a rapid and irreversible death of infected cells. This early shut-off of host translation by eIF2 α phosphorylation in SINV-infected cells could also suppress the synthesis of IFN and ISG as described recently for HCV infection [47]. This possibility agrees well with previous findings showing that infection with many Alphaviruses prevented the secretion of IFN α/β in infected cultured cells and animals, despite SINV showed an exquisite sensitivity to IFN α/β when cells were primed with this cytokine before infection [32,48,49]. For other viruses such as poxviruses, however, preventing PKR activation during *in vivo* replication could maintain protein synthesis in infected cells for more time, allowing the virus to express the late genes required for the culmination of the cycle [7,36]. In this case, viruses had to further acquire a more specific mechanism to prevent the synthesis and secretion of interferons by blocking activation of NFkB or IRF7 factors [6,9].

Our results show that, in addition to VV, EMCV expresses a trans-acting inhibitor that prevented the activation of PKR in response to SINV superinfection. However, unlike to that found in poliovirus-infected human cells, anti-PKR activity of EMCV



Figure 4. Anti-PKR mechanisms can be exchanged between viruses. *A*) Previous infection with some viruses prevented $elF2\alpha$ phosphorylation after SINV virus superinfection. The protocol followed for mixed infections is outlined. Wild type MEFs were infected with the indicated viruses at a moi of 2 pfu/cell. Three hours later, cells were superinfected with SINV at moi of 25 pfu/cell and 5 h later lysed in sample buffer for immunoblot analysis against SINV capsid (SINV C, upper panel) and anti-phosphoelF2 α (bottom panel). *B*) Expression of VV E3L gene rescued translation of ΔDLP SINV mutant. MEF-E3L cells were induced for the expression of VV E3L by tetracycline withdrawal and infected with SINV-WT or SINV- ΔDLP mutant. Five hours later, cells were labeled with [³⁵S]-Met+Cys for 30 min and analyzed by SDS-PAGE followed of autoradiography and by immunoblot against anti-FKR and anti-phospho elF2 α . Parallel cultures were infected at moi of 5 pfu/cell and viral yields were determined by plaque assay 2 days later. Data are the mean \pm SD from three and two independent experiments in PKR^{+/+} and PKR^{o/o} cells, respectively. doi:10.1371/journal.pone.0016711.g004

does not seem to involve degradation of the kinase since no alteration in the steady state of PKR was found in EMCVinfected MEFs (Figure 2). For VSV, the absence of effect on SINV-triggered eIF2a phosphorylation was expected in part given the exquisite sensitivity that VSV showed to PKR activation and eIF2a phosphorylation. Thus, VSV seems to lack of an autonomous anti-PKR activity as suggested from previous results [16], suggesting that this virus could hide the replicative forms of dsRNA in a way that go unnoticed for host PKR [20]. For VV, our results show that E3 protein accounted for almost all the anti-PKR activity in doubly infected cells, stressing the main role of this protein over the K3 product that probably has only a marginal contribution to the anti-PKR activity of the virus as suggested before [39]. The heterotypic complementation obtained for SINV Δ DLP mutant in cells expressing E3 revealed a notable case of functional analogy that reinforced the importance of anti-PKR mechanisms for virus replication in vertebrates. Thus, as showed for other viruses such as VV, Herpes and Influenza A, elimination of anti-PKR mechanism drastically reduced the replication of SINV virus in wild type animals (Figure 4). However, SINV Δ DLP mutant replicated to wild type levels in $PKR^{\alpha/o}$ mice where no eIF2 α phosphorylation in infected neurons was detected. This result showed that cis-acting DLP structure in SINV 26S mRNA is only required to overcome the effect of eIF2 α phosphorylation on translation of viral mRNA, in contrast with other scenarios such as Influenza NS1 or VV E3 proteins whose important functions extend beyond to PKR inhibition [36,38,42,50]. At least for SINV virus, our results demonstrate that resistance to PKR is exerted exclusively at translational level and that probably only involves overcoming eIF2 phosphorylation. In our opinion this is a remarkable point because the existence of other substrates of PKR apart of eIF2 α has been suggested before [51,52] despite the exquisite catalytic specificity that PKR shows for eIF2 [17,21].

Finally, the concept of molecular mimicry where pathogens have taken on proteins or protein domains from their hosts to gain evolutionary advantages could be also applicable to RNA structures such as DLP involved in translational control of some cellular mRNA. Thus, it is possible that Alphavirus had copied a mechanism of translation initiation that already operated in the cell. In our opinion, this is a very exciting possibility that deserves further investigation.

Materials and Methods

Ethics Statement

This study was carried out in strict accordance with the recommendations in the Directive 86/609/EEC of the European Union on the protection of animals used for experimental and other scientific purposes, and that was implemented by the Spanish Government under approval N° 1201/2005. The protocol was approved by the committee on the ethics of animal experiments of Universidad Autónoma de madrid (Permit Number: CEI 20-419). All surgery was performed under isoflurane anesthesia, and all efforts were made to minimize suffering.

Cells, virus and infections

MEFs derived from wild type (C57BL6/129sv) and $PKR^{\rm o/o}$ knock-out mice was described previously [53]. MEFs expressing VV E3L gene under the control of tetracyclin repressor were described previously [42]. SINV, SFV, VSV (indiana strain), EMCV and VV-ΔE3L viruses were grown in BHK21. Influenza A virus (Victoria strain, a gift of Amelia Nieto, CNB, Madrid) and Vaccinia virus (VV and VV-Luc, [54]) were amplified in BSC-10 cells. SINV Δ DLP mutant bearing 7 point mutations that disrupted the secondary structure of downstream loop structure (DLP) in 26S mRNA was described previously [29]. Cells growing in 24 well plates were infected at the indicated multiplicity of infection (moi) in DMEN lacking serum for 1 h. After that, fresh medium containing 10% of calf or foetal serum was added. At the indicated times, cells were washed twice with cold DMEN lacking serum and lysed in sample buffer. To estimate viral yields, infections were done at a moi of 1-2 pfu/cell and two days later both medium and cells were recovered, frozen and thawed three times, centrifuged for 10 min at 8,000×g and virus in the supernatant was titrated in BHK21 cells.

Infection of animals

4–6 week old mice of 129sv strain were infected with 10^7 pfu of SINV by intranasal inoculation. *PKR* knock-out mice in the same genetic background were kindly provided by J.C. Bell (University of Ottawa, Canada) [55]. At 4–5 days postinfection, animals were sacrificed by cervical dislocation, decapitated and brain extracted. For VV inoculation, 5 weeks old mice of BALB/c strain were inoculated by the intraperitoneal route with 2×10^7 pfu of a recombinant virus that expresses luciferase (VV-Luc) [54] and organs were extracted at 1dpi. Brains and spleens were fixed O/N with 4% of PFA, cryoprotected with 20% sucrose in PBS and frozen for cryosectioning. Brains and spleens of some animals were taken to titrate viral yields at 5 dpi. In these cases, brains were homogenated in 1 mL of PBS, centrifuged 10 min at 8,000×g and

References

- 1. Van Valen L (1974) Two modes of evolution. Nature 252: 298–300.
- Webber C, Ponting CP (2004) Genes and homology. Curr Biol 14: R332–333.
 Paterson S, Vogwill T, Buckling A, Benmayor R, Spiers AJ, et al. (2010)
- Antagonistic coevolution accelerates molecular evolution. Nature 464: 275–278.Akira S, Uematsu S, Takeuchi O (2006) Pathogen recognition and innate
- immunity. Cell 124: 783–801.
 Pichlmair A, Reis e Sousa C (2007) Innate recognition of viruses. Immunity 27:
- 370–383.
 Bowie AG, Unterholzner L (2008) Viral evasion and subversion of pattern-
- recognition receptor signalling. Nat Rev Immunol 8: 911–922.
- Sadler AJ, Williams BR (2008) Interferon-inducible antiviral effectors. Nat Rev Immunol 8: 559–568.
- Elde NC, Malik HS (2009) The evolutionary conundrum of pathogen mimicry. Nat Rev Microbiol 7: 787–797.
- Suthar MS, Gale M, Jr., Owen DM (2009) Evasion and disruption of innate immune signalling by hepatitis C and West Nile viruses. Cell Microbiol 11: 880–888.

viral titres in the supernatants were quantified by plaque assay in BHK21 cells.

Immunofluorescence of brain sections

Brains were cryosectioned at ~15 μm and the resulting slides were post-fixed with 4% PFA for 20' at RT, treated with 50 mM of NH₄Cl, permeabilized with 0.2% Triton X-100, blocked in 5% of BSA and incubated O/N at 4°C with the indicated antibodies in a wet chamber. Primary antibodies used were: anti-phospho eIF2 α (Cell Signalling, 1:250 dilution), anti-SINV capsid (1:500) and anti-VVp14 (1:500). Slides were then incubated with secondary antibodies coupled to Alexa 488 or 595 for 2 h at RT, washed with PBS-0.1% Triton X-100, stained with DAPI and mounted. Slides were examined and photographed in a Leika confocal microscope.

Metabolic labeling and Immunoblot analysis

Cells growing in 24 well plates were infected with the indicated virus at a moi of 25 pfu/cell. At the indicated times, cultures were labeled with 25 µCi/ml of Met+Cys (NEB) for 30', washed with DMEN lacking serum and lysed in sample buffer as described previously. Samples were analyzed in a 12% SDS-PAGE, soaked in 1 M sodium salicylate, dried and exposed to X-rays films (AGFA). For immunoblot, proteins were transferred to nitrocellulose membrane and probed with the indicated antibodies: anti-PKR (1:250, Santa Cruz Biotech.), anti-phosphoeIF2a (1:500, Invitrogen), anti-eIF2 (1:500 Santa Cruz) and anti-β-actin (1:2000, Sigma). For PKR detection, protein transfer was carried out O/N at 4°C in a wet transfer apparatus (BIORAD), whereas a semi-dry transfer (BIORAD) was used for detection of eIF2 and b-actin proteins. To detect phosphoeIF2a, blots were first probed with anti-phosphoeIF2 α antibody, incubated with mild stripping buffer (0.1 M NaOH, 15') and then probed with anti-total $eIF2\alpha$ antibody. Blots were revealed by ECL as described previously [29].

Acknowledgments

We are in debt to J. M^a Almendral for having allowed us to work at his laboratory since 2005 and J.J. Berlanga (Centro de Biología Molecular Severo Ochoa) for providing us with mice. Thanks go to J. C. Bell (University of Ottawa, Canada) and B.R. Williams (Cleveland Clinic, USA) for PKR knock-out mice and MEFs, respectively.

Author Contributions

Conceived and designed the experiments: IV. Performed the experiments: EDG RT IV JLN. Analyzed the data: IV EDG. Contributed reagents/ materials/analysis tools: JLN ME. Wrote the paper: IV.

- Kang DC, Gopalkrishnan RV, Wu Q, Jankowsky E, Pyle AM, et al. (2002) mda-5: An interferon-inducible putative RNA helicase with double-stranded RNAdependent ATPase activity and melanoma growth-suppressive properties. Proc Natl Acad Sci U S A 99: 637–642.
- Yoneyama M, Kikuchi M, Natsukawa T, Shinobu N, Imaizumi T, et al. (2004) The RNA helicase RIG-I has an essential function in double-stranded RNAinduced innate antiviral responses. Nat Immunol 5: 730–737.
- Garcia MA, Gil J, Ventoso I, Guerra S, Domingo E, et al. (2006) Impact of protein kinase PKR in cell biology: from antiviral to antiproliferative action. Microbiol Mol Biol Rev 70: 1032–1060.
- Kato H, Takeuchi O, Sato S, Yoneyama M, Yamamoto M, et al. (2006) Differential roles of MDA5 and RIG-I helicases in the recognition of RNA viruses. Nature 441: 101–105.
- Wang Z, Choi MK, Ban T, Yanai H, Negishi H, et al. (2008) Regulation of innate immune responses by DAI (DLM-1/ZBP1) and other DNA-sensing molecules. Proc Natl Acad Sci U S A 105: 5477–5482.

- Meurs E, Chong K, Galabru J, Thomas NS, Kerr IM, et al. (1990) Molecular cloning and characterization of the human double-stranded RNA-activated protein kinase induced by interferon. Cell 62: 379–390.
- Balachandran S, Roberts PC, Brown LE, Truong H, Pattnaik AK, et al. (2000) Essential role for the dsRNA-dependent protein kinase PKR in innate immunity to viral infection. Immunity 13: 129–141.
- Dar AC, Dever TE, Sicheri F (2005) Higher-order substrate recognition of eIF2alpha by the RNA-dependent protein kinase PKR. Cell 122: 887–900.
- Romano PR, Garcia-Barrio MT, Zhang X, Wang Q, Taylor DR, et al. (1998) Autophosphorylation in the activation loop is required for full kinase activity in vivo of human and yeast eukaryotic initiation factor 2alpha kinases PKR and GCN2. Mol Cell Biol 18: 2282–2297.
- Green SR, Mathews MB (1992) Two RNA-binding motifs in the doublestranded RNA-activated protein kinase, DAI. Genes Dev 6: 2478–2490.
- Weber F, Wagner V, Rasmussen SB, Hartmann R, Paludan SR (2006) Doublestranded RNA is produced by positive-strand RNA viruses and DNA viruses but not in detectable amounts by negative-strand RNA viruses. J Virol 80: 5059–5064.
- Dever TE (2002) Gene-specific regulation by general translation factors. Cell 108: 545–556.
- Stojdl DF, Abraham N, Knowles S, Marius R, Brasey A, et al. (2000) The murine double-stranded RNA-dependent protein kinase PKR is required for resistance to vesicular stomatitis virus. J Virol 74: 9580–9585.
- Black TL, Barber GN, Katze MG (1993) Degradation of the interferon-induced 68,000-M(r) protein kinase by poliovirus requires RNA. J Virol 67: 791–800.
- Ikegami T, Narayanan K, Won S, Kamitani W, Peters CJ, et al. (2009) Rift Valley fever virus NSs protein promotes post-transcriptional downregulation of protein kinase PKR and inhibits eIF2alpha phosphorylation. PLoS Pathog 5: e1000287.
- Chou J, Kern ER, Whitley RJ, Roizman B (1990) Mapping of herpes simplex virus-1 neurovirulence to gamma 134.5, a gene nonessential for growth in culture. Science 250: 1262–1266.
- 26. He B, Gross M, Roizman B (1997) The gamma(1)34.5 protein of herpes simplex virus 1 complexes with protein phosphatase 1alpha to dephosphorylate the alpha subunit of the eukaryotic translation initiation factor 2 and preclude the shutoff of protein synthesis by double-stranded RNA-activated protein kinase. Proc Natl Acad Sci U S A 94: 843–848.
- Mohr I (2004) Neutralizing innate host defenses to control viral translation in HSV-1 infected cells. Int Rev Immunol 23: 199–220.
- Gorchakov R, Frolova E, Williams BR, Rice CM, Frolov I (2004) PKRdependent and -independent mechanisms are involved in translational shutoff during Sindbis virus infection. J Virol 78: 8455–8467.
- Ventoso I, Sanz MA, Molina S, Berlanga JJ, Carrasco L, et al. (2006) Translational resistance of late alphavirus mRNA to eIF2alpha phosphorylation: a strategy to overcome the antiviral effect of protein kinase PKR. Genes Dev 20: 87–100.
- Elde NC, Child SJ, Geballe AP, Malik HS (2009) Protein kinase R reveals an evolutionary model for defeating viral mimicry. Nature 457: 485–489.
- Rothenburg S, Seo EJ, Gibbs JS, Dever TE, Dittmar K (2009) Rapid evolution of protein kinase PKR alters sensitivity to viral inhibitors. Nat Struct Mol Biol 16: 63–70.
- Binder GK, Griffin DE (2001) Interferon-gamma-mediated site-specific clearance of alphavirus from CNS neurons. Science 293: 303–306.
- Cook SH, Griffin DE (2003) Luciferase imaging of a neurotropic viral infection in intact animals. J Virol 77: 5333–5338.
- Jackson AC, Moench TR, Trapp BD, Griffin DE (1988) Basis of neurovirulence in Sindbis virus encephalomyelitis of mice. Lab Invest 58: 503–509.
- Strauss JH, Strauss EG (1994) The alphaviruses: gene expression, replication, and evolution. Microbiol Rev 58: 491–562.

- Diversity in Viral Anti-PKR Mechanisms
- Brandt TA, Jacobs BL (2001) Both carboxy- and amino-terminal domains of the vaccinia virus interferon resistance gene, E3L, are required for pathogenesis in a mouse model. J Virol 75: 850–856.
- Bergmann M, Garcia-Sastre A, Carnero E, Pehamberger H, Wolff K, et al. (2000) Influenza virus NS1 protein counteracts PKR-mediated inhibition of replication. J Virol 74: 6203–6206.
- Donelan NR, Basler CF, Garcia-Sastre A (2003) A recombinant influenza A virus expressing an RNA-binding-defective NS1 protein induces high levels of beta interferon and is attenuated in mice. J Virol 77: 13257–13266.
- Langland JO, Jacobs BL (2002) The role of the PKR-inhibitory genes, E3L and K3L, in determining vaccinia virus host range. Virology 299: 133–141.
- Chang HW, Watson JC, Jacobs BL (1992) The E3L gene of vaccinia virus encodes an inhibitor of the interferon-induced, double-stranded RNA-dependent protein kinase. Proc Natl Acad Sci U S A 89: 4825–4829.
- Romano PR, Zhang F, Tan SL, Garcia-Barrio MT, Katze MG, et al. (1998) Inhibition of double-stranded RNA-dependent protein kinase PKR by vaccinia virus E3: role of complex formation and the E3 N-terminal domain. Mol Cell Biol 18: 7304–7316.
- Domingo-Gil E, Perez-Jimenez E, Ventoso I, Najera JL, Esteban M (2008) Expression of the E3L gene of vaccinia virus in transgenic mice decreases host resistance to vaccinia virus and Leishmania major infections. J Virol 82: 254–267.
- Woolhouse ME, Webster JP, Domingo E, Charlesworth B, Levin BR (2002) Biological and biomedical implications of the co-evolution of pathogens and their hosts. Nat Genet 32: 569–577.
- Weaver SC, Barrett AD (2004) Transmission cycles, host range, evolution and emergence of arboviral disease. Nat Rev Microbiol 2: 789–801.
- Powers AM, Brault AC, Shirako Y, Strauss EG, Kang W, et al. (2001) Evolutionary relationships and systematics of the alphaviruses. J Virol 75: 10118–10131.
- Toribio R, Ventoso I (2010) Inhibition of host translation by virus infection in vivo. Proc Natl Acad Sci U S A.
- Garaigorta U, Chisari FV (2009) Hepatitis C virus blocks interferon effector function by inducing protein kinase R phosphorylation. Cell Host Microbe 6: 513–522.
- Burke CW, Gardner CL, Steffan JJ, Ryman KD, Klimstra WB (2009) Characteristics of alpha/beta interferon induction after infection of murine fibroblasts with wild-type and mutant alphaviruses. Virology 395: 121–132.
- Frolova EI, Fayzulin RZ, Cook SH, Griffin DE, Rice CM, et al. (2002) Roles of nonstructural protein nsP2 and Alpha/Beta interferons in determining the outcome of Sindbis virus infection. J Virol 76: 11254–11264.
- Hale BG, Randall RE, Ortin J, Jackson D (2008) The multifunctional NS1 protein of influenza A viruses. J Gen Virol 89: 2359–2376.
- Cuddihy AR, Wong AH, Tam NW, Li S, Koromilas AE (1999) The doublestranded RNA activated protein kinase PKR physically associates with the tumor suppressor p53 protein and phosphorylates human p53 on serine 392 in vitro. Oncogene 18: 2690–2702.
- Nakamura T, Furuhashi M, Li P, Cao H, Tuncman G, et al. (2010) Doublestranded RNA-dependent protein kinase links pathogen sensing with stress and metabolic homeostasis. Cell 140: 338–348.
- Yang YL, Reis LF, Pavlovic J, Aguzzi A, Schafer R, et al. (1995) Deficient signaling in mice devoid of double-stranded RNA-dependent protein kinase. Embo J 14: 6095–6106.
- Rodriguez JF, Rodriguez D, Rodriguez JR, McGowan EB, Esteban M (1988) Expression of the firefly luciferase gene in vaccinia virus: a highly sensitive gene marker to follow virus dissemination in tissues of infected animals. Proc Natl Acad Sci U S A 85: 1667–1671.
- Abraham N, Stojdl DF, Duncan PI, Methot N, Ishii T, et al. (1999) Characterization of transgenic mice with targeted disruption of the catalytic domain of the double-stranded RNA-dependent protein kinase, PKR. J Biol Chem 274: 5953–5962.