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Poliovirus tropism and attenuation are determined after internal ribosome entry

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Poliovirus replication is limited to a few organs, including the brain and spinal cord. This restricted tropism may be a consequence of organ-specific differences in translation initiation by the poliovirus internal ribosome entry site (IRES). A C-to-U mutation at base 472 in the IRES of the Sabin type 3 poliovirus vaccine strain, known to attenuate neurovirulence, may further restrict tropism by eliminating viral replication in the CNS. To determine the relationship between IRES-mediated translation and poliovirus tropism, recombinant human adenoviruses were used to express bicistronic mRNAs in murine organs. The IRESs of poliovirus, the cardio-tropic coxsackievirus B3 (CVB3), and the hepatotropic hepatitis C virus (HCV) mediate translation in many organs, including those that do not support viral replication. A translation defect associated with the Sabin type 3 IRES was observed in all organs examined. Poliovirus type 1 and recombinant polioviruses dependent on the IRES of CVB3 or HCV replicate in the CNS of mice and cause paralysis. Although the type 3 Sabin strain is an effective vaccine, polioviruses with a U at base 472 of the IRES cause paralysis in newborn mice. Tropism of wild-type and vaccine strains of poliovirus is therefore determined after internal ribosome entry.

Introduction

The initiation of most eukaryotic protein synthesis is dependent on the recruitment of ribosomes to the mRNA 5' end by the 5' 7-methylguanosine cap structure (1, 2). However, translation of some viral and cellular mRNAs is initiated when a *cis*-acting RNA sequence called an internal ribosome entry site (IRES) binds ribosomes in a 5' end- and cap-independent manner (3–7). The first IRES was identified within the 5' untranslated region of the poliovirus RNA genome (8), which is an mRNA but lacks a 5' cap structure (9, 10). Translation in eukaryotic cells may be regulated at the step of IRES-dependent initiation (4, 7).

It has been suggested that poliovirus tropism, defined as the organs where the virus replicates, is determined by cell type-specific differences in translation initiation by the poliovirus IRES (11–14). Poliovirus replication is limited to the brain and spinal cord, oropharyngeal and intestinal mucosa, tonsils, Peyer's patches, and cervical and mesenteric lymph nodes (15). The poliovirus receptor is not sufficient to determine poliovirus tropism, because this protein is produced in a broad range of animal tissues (16–18). The ability of poliovirus to replicate in different organs may be determined at a postentry step such as translation of the viral RNA. Cell proteins other than the canonical translation-initiation factors have been identified that influence IRES-mediated translation (19). Viral replication could be regulated by organ-specific synthesis, localization, or modification of these cell proteins. Recombinant polioviruses dependent on the IRES of human rhinovirus 2 or hepatitis C virus (HCV) do not accumulate or cause disease in the brain and spinal cord of mice (12, 13, 20). These results have been interpreted as indicating that the IRESs of rhinovirus and HCV do not mediate translation initiation in the

brain and spinal cord. However, there have been no direct measurements in organs of translation initiation dependent on the IRES of rhinovirus, poliovirus, or HCV.

Organ-specific IRES-mediated translation has also been hypothesized to account for the reduced neurovirulence of the Sabin live attenuated poliovirus vaccine strains. These vaccine strains infect the alimentary tract, producing immunity to infection but rarely causing paralysis (21). Genetic analysis has demonstrated that a point mutation within the IRES of all three poliovirus vaccine strains is a determinant of the attenuation phenotype (22–24). A mutation from C to U at nucleotide 472 in the IRES of poliovirus type 3 is sufficient to attenuate neurovirulence in primate and murine models (23, 25, 26). Studies of IRES-mediated translation initiation have demonstrated that a change from C to U at nucleotide 472 causes a translation defect *in vitro* and in cultured cells of neuronal origin (27–29). It has been hypothesized that the translation defect caused by the C472U mutation is specific to the brain and spinal cord and leads to a defect in viral replication in these organs (11, 27, 30). This replication defect might explain the reduced neurovirulence of the oral poliovirus vaccine strains. However, it is not known whether mutations in the poliovirus IRES reduce translation in the brain and spinal cord but not in other organs.

Two experimental approaches were used to determine the relationship between IRES-mediated translation and poliovirus tropism. Using recombinant human adenoviruses to express bicistronic mRNAs in murine organs, the IRESs of poliovirus, coxsackievirus B3 (CVB3), and HCV were found to mediate translation in many organs, including those that do not support viral replication. A translation defect associated with the Sabin type 3 IRES was observed in all organs examined. In a mouse model of poliomyelitis, recombinant polioviruses dependent on the IRES of CVB3 or HCV replicate in the CNS and cause paralysis. Polioviruses with a U at base 472 of the IRES cause paralysis in newborn mice. These findings demonstrate that tropism of wild-type and vaccine strains of poliovirus is determined after internal ribosome entry.

Nonstandard abbreviations used: coxsackievirus B3 (CVB3); hepatitis C virus (HCV); internal ribosome entry site (IRES); passive lysis buffer (PLB); transgenic poliovirus receptor (TgPVR).

Conflict of interest: The authors have declared that no conflict of interest exists.

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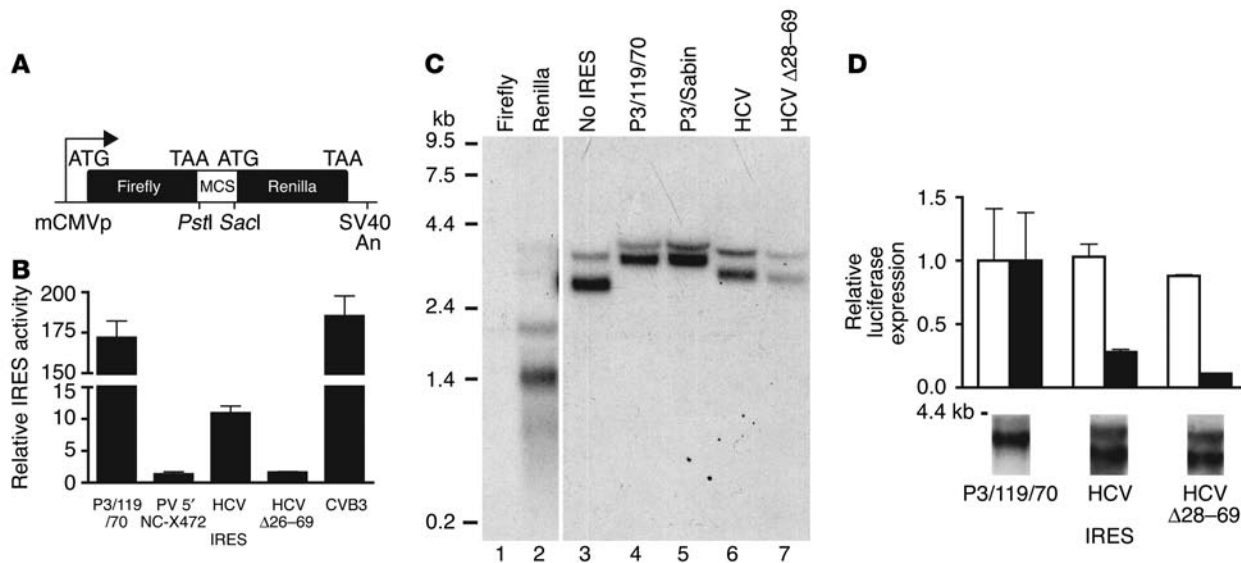


Figure 1 IRES-mediated translation in A549 cells infected with recombinant adenoviruses. (A) Schematic of bicistronic reporter DNA. The arrow indicates the transcription-initiation site of the murine cytomegalovirus immediate early promoter. Firefly luciferase and *Renilla* luciferase have independent translation initiation and termination codons. SV40 An, simian virus 40 polyadenylation signal. (B) IRES activity in cells infected with adenovirus encoding bicistronic mRNA. The x axis indicates IRES encoded in the recombinant adenovirus. PV 5'NC-X472, poliovirus IRES with *Xho*I linker substitution at nucleotide 472; HCV Δ28–69, HCV IRES with deletion of nucleotides 28–69. (C) Northern blot hybridization of A549 cells. Cells were infected with recombinant adenoviruses indicated at the top. RNA size markers are given at left. Lane 1, monocistronic firefly luciferase; lane 2, monocistronic *Renilla* luciferase; lane 3, bicistronic mRNA, no IRES; lane 4, bicistronic mRNA with P3/119/70 IRES; lane 5, bicistronic mRNA with P3/Sabin IRES; lane 6, bicistronic mRNA with HCV IRES; lane 7, bicistronic mRNA with HCV IRES lacking nucleotides 28–69. First two lanes are from the same gel as lanes 3–7 but were exposed for a longer period of time. (D) Luciferase expression in adenovirus-infected cells. Upper panel: Firefly luciferase (white bars) and *Renilla* luciferase expression (black bars). Luciferase was normalized to the P3/119/70 IRES value for each reporter protein, and also to the quantity of bicistronic mRNA determined by Northern blot hybridization. Lower panels: Representative Northern blot hybridizations of RNA from the same infected cells. IRES encoded by bicistronic mRNA is indicated at the bottom. Data points in B and D, the results of separate experiments, are the mean of three infections, and error bars indicate SD.

Results

Expression of bicistronic mRNAs from recombinant human adenovirus. A system was developed to measure IRES activity in murine organs. Recombinant human adenovirus vectors were produced that express bicistronic mRNAs that encode two reporter proteins separated by an IRES (Figure 1A). These bicistronic mRNAs possess a 5' cap structure, and translation of the first open reading frame occurs by 5' end-dependent initiation. Efficient translation of the second open reading frame requires an IRES to mediate internal binding of ribosomes.

To demonstrate that an IRES-dependent mRNA can be translated when expressed from an adenovirus vector, A549 cells were infected with adenoviruses that encode bicistronic mRNAs with the IRES of poliovirus, HCV, or CVB3. Inclusion of the IRES within a bicistronic mRNA results in significantly higher levels of *Renilla* luciferase activity compared with a short, unstructured sequence (Figure 1B). These results agree with those of previous studies that showed that the inclusion of the IRES of poliovirus, HCV, or CVB3 in an mRNA results in increased expression of the downstream reporter gene (8, 31, 32). The effects of mutations previously shown to abrogate IRES-mediated translation were determined. Substitution of an *Xho*I linker for nucleotides 472–479 of the poliovirus IRES, or deletion of nucleotides 28–69 from the HCV IRES, has been shown to impair translation (33, 34), and these mutations reduce *Renilla* luciferase activity in adenovirus vector-infected cells (Figure 1B). These findings provide genetic

evidence that *Renilla* luciferase activity observed in adenovirus vector-infected cells is a consequence of internal initiation.

To confirm that *Renilla* luciferase is present only in full-length mRNAs, Northern blot hybridization of RNA from adenovirus vector-infected cells was performed (Figure 1C). No species smaller than the full-length bicistronic mRNA was detected in cells infected with these adenoviruses. These findings demonstrate that *Renilla* luciferase is translated by internal ribosome entry on full-length bicistronic mRNAs, and not by 5' end-dependent translation of smaller RNAs produced by RNA degradation, splicing, or aberrant transcription. An additional RNA approximately 500 nucleotides larger than the bicistronic mRNA was also detected (Figure 1C). This RNA likely results from transcription termination at the adenovirus E1b polyadenylation signal 500 nucleotides downstream of the simian virus 40 polyadenylation signal (35).

Because IRES activity is based on standardization of *Renilla* to firefly activity, any effect of the IRES on translation of the upstream reporter protein would prevent a reliable quantitation of translation initiated by the IRES. To address this possibility, firefly and *Renilla* luciferase activities were measured in infected cells and normalized to the amount of bicistronic RNA as determined by Northern blot hybridization. Normalization compensates for differences in luciferase activity that result from variation in infection efficiency or transcription. The findings demonstrate that the IRES has no effect on firefly luciferase translation (Figure 1D).

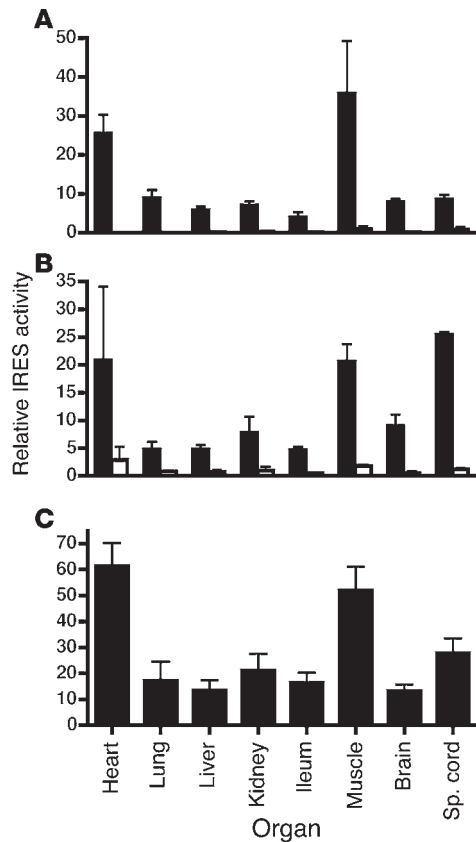


Figure 2

IRES-mediated translation in murine organs infected with recombinant adenoviruses. (A) Poliovirus type 3 119/70 IRES (black bars) and poliovirus type 1 5'NC-X472 IRES (white bars). (B) HCV 1b IRES (black bars) and HCV 1a IRES with nucleotides 28–69 deleted (white bars). (C) CVB3 IRES. Data points are the mean of at least five mice, and error bars indicate SD.

ranging from 10^6 to 10^{10} PFUs, levels of firefly luciferase and *Renilla* luciferase increased nearly 10,000 times (Figure 3A), but activity of the poliovirus IRES remained constant (Figure 3B). Therefore IRES activity in an organ is not affected by the quantity of infecting vector virus.

Does the IRES determine sites of viral replication and disease? Studied in isolation from the viral genome, the IRES of poliovirus, HCV, or CVB3 mediates translation in many organs. In the genome, organ specificity of an IRES could be influenced by viral RNA sequences or proteins produced during viral replication. To address this possibility, the poliovirus IRES was substituted with the cognate sequence from viruses that infect different organs (Figure 4A). The IRES of human poliovirus type 1 was replaced with that of HCV, which is hepatotropic (36), or with the IRES of CVB3, which causes myocarditis (37). These recombinant viruses were named P1/HCV and P1/CVB3. Single-step growth analysis of both recombinant viruses in HeLa cells reveals a defect in replication, comprising an early delay in virus production (Figure 4B). While the final yield of P1/CVB3 approached that of poliovirus type 1, the yield of P1/HCV was significantly lower. Previous results have demonstrated that the poliovirus IRES contains determinants of viral RNA replication (38, 39). It is possible that exchange of the poliovirus IRES has removed *cis*-acting sequences important for RNA replication.

Virulence and tropism of recombinant viruses were determined in mice transgenic for the human poliovirus receptor gene (40). After intraperitoneal inoculation of adult transgenic poliovirus receptor (TgPVR) mice with poliovirus type 1, virus replicates to high titers in the brain and spinal cord (Figure 5A). In contrast, virus titers in the heart steadily decline after infection, and virus

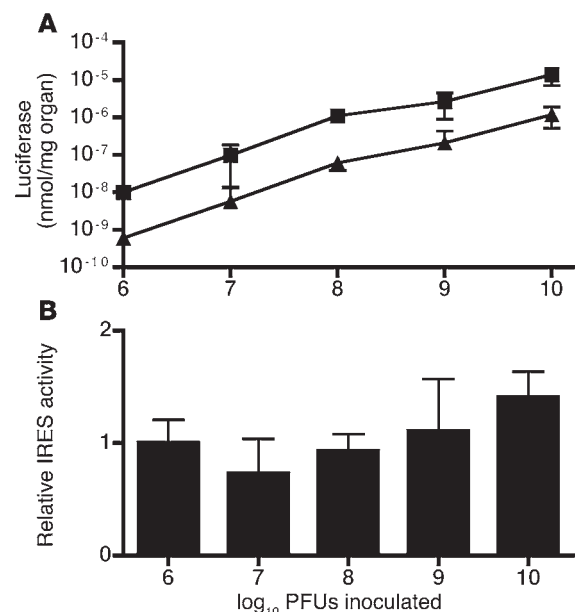
IRES-mediated initiation of translation in murine organs. In animals, IRES-mediated internal initiation of translation could be restricted to specific organs. To address this hypothesis, mice were infected with recombinant adenovirus vectors that encode bicistronic mRNAs. The IRES of poliovirus, HCV, and CVB3 is functional in all organs examined, including brain, spinal cord, skeletal muscle, heart, lung, liver, kidney, and ileum (Figure 2, A–C).

To provide genetic evidence that *Renilla* luciferase translation in mice is mediated by the viral IRES, the effect of mutations in the IRES was assessed. As was observed in cultured cells (Figure 1B), substitution of an *Xho*I linker for nucleotides 472–479 of the poliovirus IRES, or deletion of nucleotides 28–69 from the HCV IRES, nearly eliminates IRES activity in mouse organs (Figure 2, A and B). No firefly or *Renilla* luciferase activity was detected when mice were infected with an adenovirus lacking the promoter necessary for production of bicistronic mRNA with the poliovirus IRES (data not shown). This result shows that translatable *Renilla* luciferase mRNAs are not produced from a promoter within the poliovirus IRES.

IRES activity should be independent of the quantity of infecting virus. When mice were inoculated with amounts of virus

Figure 3

Relationship between dose of recombinant adenovirus and IRES activity in mouse brain. (A) Firefly luciferase (squares) and *Renilla* luciferase (triangles) expression in mouse brain. (B) Activity of the poliovirus type 3 119/70 IRES in the mouse brain, normalized to activity at a dose of 10^6 PFUs. The x axis indicates adenovirus dose in \log_{10} PFUs. Data points are the mean of five infections, and error bars indicate SD.



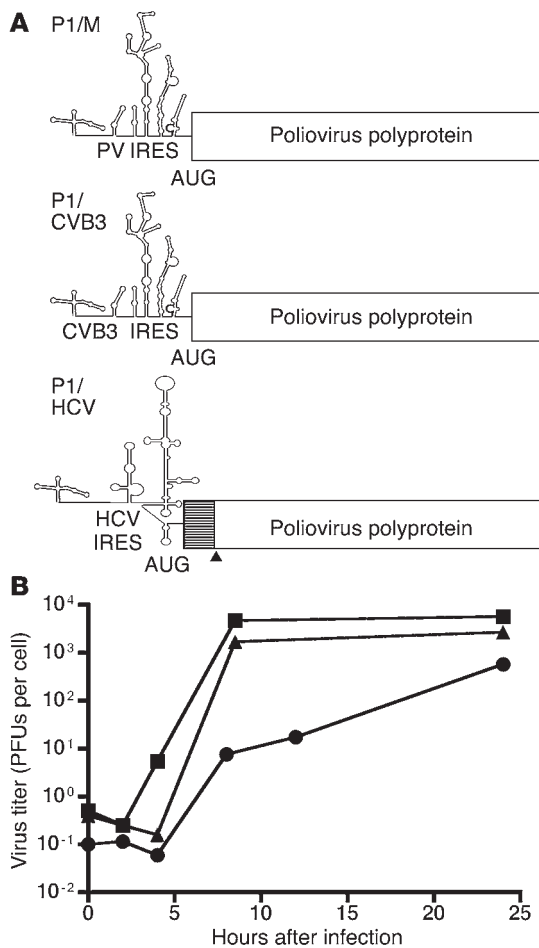


Figure 4

Replication and virulence of recombinant poliovirus strains. **(A)** Genome structure of poliovirus type 1 strain Mahoney, recombinant strain P1/CVB3, and recombinant strain P1/HCV. IRES, predicted AUG initiation codons, and poliovirus polyprotein (open box) are indicated. Translation of P1/HCV is predicted to initiate at the HCV AUG initiation codon, which is followed by 369 nucleotides of HCV polyprotein sequence (lined box). Sequence encoding the recognition site for poliovirus protease 2A^{pro} (triangle) separates HCV sequence and nucleotide 745 of the poliovirus genome. **(B)** Single-step replication analysis in HeLa cells of poliovirus type 1 strain Mahoney (squares), P1/CVB3 (triangles), and P1/HCV (circles). Data points are the mean of two infections.

titers in the pancreas increase by day 3 and subsequently decline (Figure 5A). In newborn TgPVR mice, there is a large increase in virus titer in the brain and spinal cord, and a moderate increase in the liver (Figure 5C). Infection of adult and newborn TgPVR mice leads to flaccid limb paralysis and death (40). Viral replication and pathogenesis are not observed in nontransgenic mice (40).

To determine the effect of IRES replacement on poliovirus pathogenesis, TgPVR mice were infected with recombinant viruses. P1/CVB3 replicates and causes disease in adult TgPVR mice (Figure 5, B and E). Virus titers in the brain and spinal cord increase 10,000-fold during the course of infection (Figure 5B). Some infected mice develop flaccid hind-limb paralysis or die (Figure 5E). CVB3 replication and pathogenesis occur in heart and pancreas of adult mice (41, 42), yet the pattern of P1/CVB3 replication in these organs is unchanged from that of poliovirus type 1 (Figure 5, A and B). When adult TgPVR mice were inoculated with P1/HCV, viral replication and disease did not occur (data not shown). However, P1/HCV replicates and causes disease in newborn TgPVR mice (Figure 5, D and E). Virus titers in the brain and spinal cord increase 500- and 1,000-fold, respectively, during the course of infection (Figure 5D). Some infected mice develop flaccid hind-limb paralysis or die (Figure 5E). HCV replicates and causes disease in primate liver, yet in TgPVR mice P1/HCV is cleared from this organ (Figure 5D). Virulence of virus recovered from paralyzed newborn TgPVR mice was unchanged in newborn and adult mice

(data not shown). Replication of poliovirus type 1 in the brain and spinal cord of mice and development of paralytic disease are dependent on the human poliovirus receptor (40). P1/HCV and P1/CVB3 are cleared from the brain and spinal cord of nontransgenic mice (Figure 5, B and D), and paralysis does not occur (Figure 5E and data not shown). The human poliovirus receptor is required for paralytic disease and increases in viral titers after inoculation, further evidence that P1/HCV and P1/CVB3 replicate in the murine brain and spinal cord.

Do mutations in the IRES of poliovirus vaccine strains have an organ-specific effect on translation? It has been suggested that neuroattenuation caused by a C472U mutation in the IRES of poliovirus vaccine strains is a consequence of reduced translation of poliovirus RNA in the brain and spinal cord. The effect of the C472U mutation on translation in murine organs was therefore determined. Recombinant adenoviruses were produced that encode bicistronic mRNAs with the poliovirus type 3 IRES and either a C or a U at nucleotide 472. The C472U mutation decreases IRES-dependent translation in continuous cell lines of both neuronal and non-neuronal origin (Figure 6A). No species smaller than the full-length bicistronic mRNA was detected in cells infected with either adenovirus (Figure 6A). The C472U mutation also decreases IRES-dependent translation in murine brain, spinal cord, heart, lung, liver, kidney, ileum, and muscle (Figure 6B).

It has previously been shown that poliovirus strains with the C472U mutation are cleared from the brain and spinal cord of adult mice and fail to cause paralysis (26). To determine whether virus strains with the C472U mutation have lost the ability to replicate within the murine brain and spinal cord, newborn TgPVR mice were infected with polioviruses with either C or U at nucleotide 472. The neurovirulence of these viruses was then determined as a measure of their ability to replicate in the murine brain and spinal cord (Table 1). Poliovirus strain PRV7.3, with U at nucleotide 472, is nearly as neurovirulent in newborn mice as virus strain PRV8.4, which is identical except for a C at nucleotide 472. In contrast, the neurovirulence of virus strain PRV7.3 is attenuated in adult TgPVR mice. The poliovirus type 3 vaccine strain P3/Sabin, which is neuroattenuated in adult TgPVR mice, is virulent in newborn TgPVR mice. As expected, virus strains P3/119/70 and PRV8.4, both with C at nucleotide 472, are neurovirulent in adult TgPVR mice, and virus strain P3/119/70 is highly neurovirulent in newborn mice. These findings demonstrate that the neuroattenuating mutation at nucleotide 472 of the poliovirus genome does not eliminate viral replication in the murine brain and spinal cord.

Viruses with a reversion of the C472U mutation may accumulate during replication of PRV7.3 and P3/Sabin in cell culture

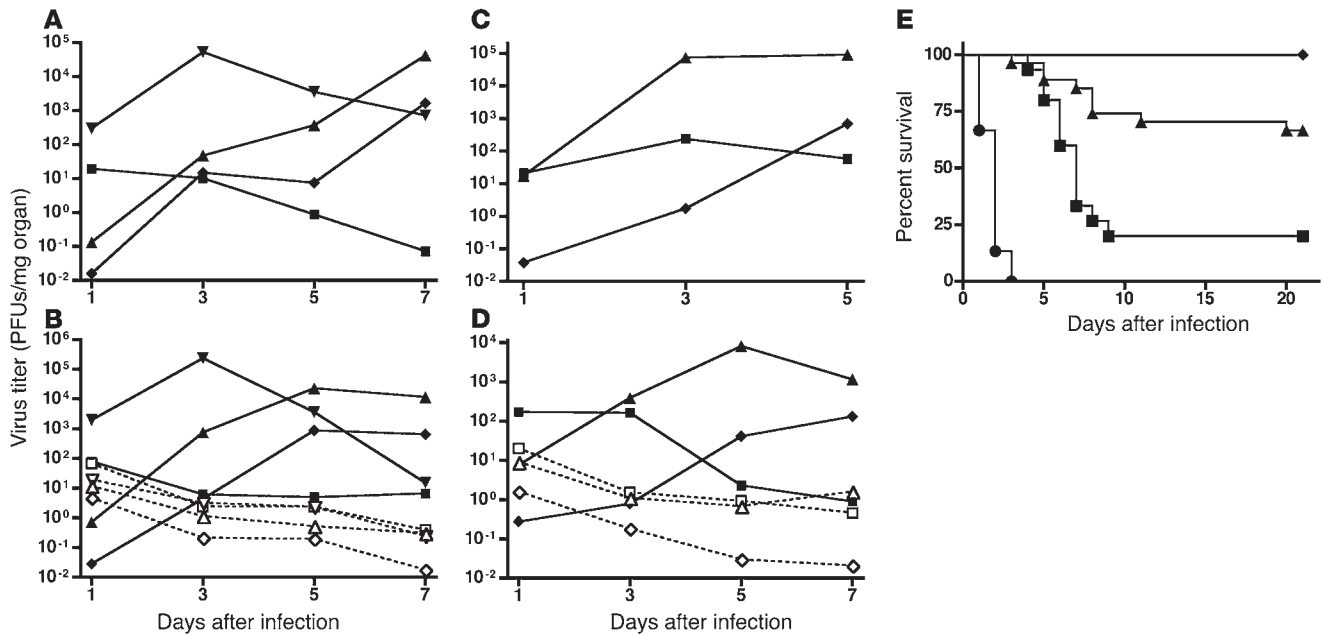


Figure 5 Poliovirus infection of mice. (A–D) Poliovirus replication in murine organs. The y axis indicates virus titer at the indicated days after infection. (A) Virus titers in pancreas (filled inverted triangles), spinal cord (filled triangles), brain (filled diamonds), and heart (filled squares) of adult TgPVR mice infected with poliovirus. (B) Virus titers in spinal cord (filled and open triangles), brain (filled and open diamonds), pancreas (filled inverted and open inverted triangles), and heart (filled and open squares) of adult TgPVR (solid lines) and nontransgenic (dashed lines) mice infected with P1/CVB3. (C) Virus titers in spinal cord (filled triangles), brain (filled diamonds), and liver (filled squares) of newborn TgPVR mice infected with poliovirus. (D) Virus titers in spinal cord (filled and open triangles), brain (filled and open diamonds), and liver (filled and open squares) of newborn TgPVR (solid lines) or nontransgenic (dashed lines) mice infected with P1/HCV. Data points are the geometric mean titer in organs from at least three mice. (E) Virulence of recombinant poliovirus strains. The y axis indicates the percentage of surviving mice at different times after infection. TgPVR mice were infected with 10^7 PFUs P1/CVB3 (squares), 2×10^6 PFUs P1/HCV (triangles), or 10^9 PFUs P1/HCV (circles); nontransgenic mice were infected with 2×10^6 PFUs P1/HCV (diamonds).

or in animals (23, 43). The fraction of viral revertants can be readily determined by restriction enzyme cleavage of a DNA copy of the IRES (44). To confirm that stocks of PRV7.3 and P3/Sabin are free of such revertants, the proportion of U at nucleotide 472 was determined by cleavage of a DNA copy of the IRES with *MboI* (Figure 7). According to a standard curve produced with plasmid DNA (data not shown), more than 99% of the viral RNA in stocks of PRV7.3 and P3/Sabin has a U at nucleotide 472. Similarly, nearly 99% of the PRV7.3 or P3/Sabin viral RNA recovered from the brains of paralyzed mice has a U at nucleotide 472. When plasmid DNA that encodes a C at this position was assayed, cleavage of 100% of *MboI* sites was never observed, most likely because the DNA is damaged during PCR amplification (Figure 7, lane 2). However, the proportion of base C detected at nucleotide 472 in viral RNA from stocks of PRV8.4 and P3/119/70 and in RNA recovered from the brains of mice infected with these viruses was similar to that in plasmid DNA that encodes a C at this position. These findings show that paralysis in newborn mice inoculated with neuroattenuated viruses is not caused by a reversion of the C472U mutation.

In cell culture, the C472U mutation confers a temperature-sensitive phenotype that can be suppressed by mutations in the 2A^{Pro} coding region (45). Furthermore, mutation to A at IRES nucleotide 537 might restore

neurovirulence by improving base pairing between nucleotides 472 and 537 (46). To ensure that the neurovirulence of PRV7.3 in newborn mice is not a consequence of suppressor mutations, the nucleotide sequence was determined in these regions of viral RNA from the brains of four paralyzed mice. No sequence changes were found in the 2A^{Pro} coding region. In viral RNA from two mice, the parental base, G, was found at nucleotide 537. In viral RNA from two other mice, a mixed population of the parental nucleotide 537G and the mutation 537A was found. These results indicate that mutations within the 2A^{Pro} coding region or at nucleotide 537 are not essential for neurovirulence.

Table 1 Virulence of poliovirus strains in TgPVR mice

Virus	Base 472	LD ₅₀ in mouse, PFUs			
		TgPVR	TgPVR	Non-Tg	Non-Tg
		4 weeks old	1–2 days old	4 weeks old	1–2 days old
PRV7.3 ^A	U	>10 ⁸	1.3 × 10 ⁴	ND	ND
PRV8.4 ^A	C	2.4 × 10 ⁵	3.7 × 10 ³	ND	ND
P3/Sabin	U	>10 ⁸	6.2 × 10 ⁴	>2 × 10 ⁸	>5 × 10 ⁷
P3/119/70	C	2.8 × 10 ⁵	2 × 10 ²	>10 ⁸	>3 × 10 ⁷

^ARecombinant viruses comprising nucleotides 1–784 from P3/Sabin (PRV7.3) or P3/119 (PRV8.4) and the remainder of the genome from P2/Lansing (26). Human poliovirus receptor is not essential for infection of mice with P2/Lansing. ND, not determined.

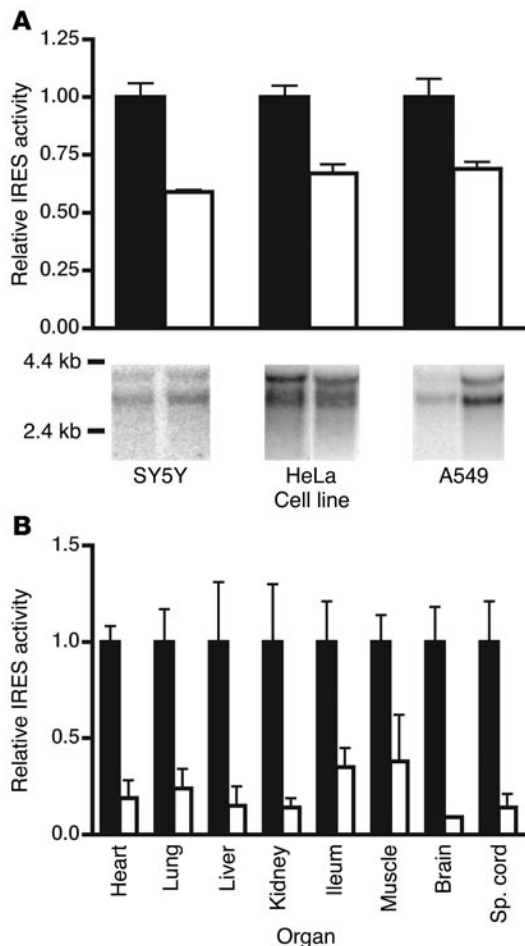


Figure 6

Effect of C472U mutation on translation of bicistronic mRNAs expressed by recombinant adenovirus. (A) Upper panel: Activity of P3/Sabin IRES (white bars) relative to P3/119/70 IRES (black bars) in SY5Y, HeLa, and A549 cells. Lower panel: Representative Northern blot hybridization of RNA from the same infected cells. (B) Activity of P3/Sabin IRES (white bars) relative to P3/119/70 IRES (black bars) in murine organs. Data points are the mean of three infections, and error bars indicate SD.

hypothesis cannot be tested in cell culture, because poliovirus replication occurs in cell lines derived from many organs that do not support viral replication (48). However, direct measurements of viral IRES-mediated translation in organs would address the question of whether IRES activity correlates with the sites of virus replication in the animal. Previous analyses of transgenic mice that express a bicistronic mRNA have demonstrated a lack of organ specificity in translation mediated by the IRES of encephalomyocarditis virus (49) and Theiler murine encephalomyelitis virus (50). In the present study, the use of recombinant adenoviral vectors permitted the analysis of the IRES from multiple viruses in a fraction of the time required for a transgenic approach. Contrary to expectations, the IRESs of HCV, poliovirus, and CVB3 mediate translation in many murine organs, including those that are not sites of virus replication. Therefore, levels of IRES-dependent translation do not determine the organ-specific pattern of poliovirus replication. By expression of bicistronic mRNAs with adenovirus vectors, a pattern of IRES-mediated translation was revealed in organs that was not apparent in studies of cultured cells. It would therefore be of interest to use this approach to study the effect of different physiological states on cellular IRES-mediated translation.

Mouse models of human poliomyelitis may also provide insight into the relationship between poliovirus IRES-mediated translation and viral pathogenesis. Transgenic mice that produce the poliovirus receptor, CD155, are susceptible to poliovirus infection and develop disease that clinically and histopathologically resembles poliomyelitis (40, 51). Although

Discussion

Internal ribosome entry mediated by a cellular or viral IRES is commonly studied by monitoring the synthesis of IRES-dependent reporter proteins. The poliovirus IRES can also be modified in its normal location in an mRNA, the viral genome, and the effects on the viral-replication cycle can then be determined in infected cells or organs. This combination affords a unique approach to the study of translational control in mammalian tissues.

Although the poliovirus receptor is produced in a broad range of organs (17, 18), viral replication and disease are limited to a few sites, including the brain, spinal cord, and alimentary tract. It has been proposed that this restriction is established at the step of IRES-mediated translation of the viral RNA (11–14, 47). This

Figure 7

Base at nucleotide 472 in poliovirus recovered from paralyzed mice. Upper panel: Representative Southern blot hybridization of *Mbo*I-cleaved PCR products. The 93-bp DNA (472U) and 61-bp DNA (472C) are indicated. Lower panel: Proportion of 472U in each sample, determined by quantitative analysis of Southern blots. For mouse brain, data points are the mean of at least three mice except for P3/Sabin, for which one mouse was analyzed. Error bars indicate SD. The x axis indicates template for RT-PCR. Lane 1, plasmid DNA encoding P3/Sabin IRES; lane 2, plasmid DNA encoding P3/119/70 IRES; lanes 3–6, virus stocks used for infection of TgPVR mice; lanes 7–10, RNA from brain of TgPVR mice infected with the indicated virus stock.

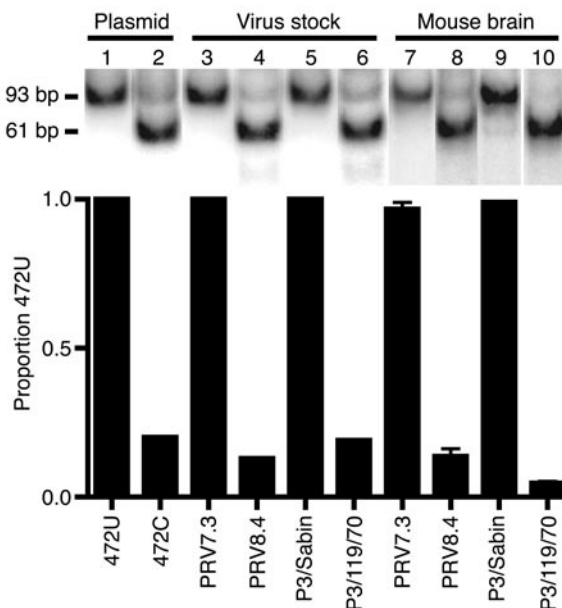




Table 2
Components of PCR reactions

Template	5' Primer	3' Primer	PCR product
pGL3 promoter (Promega Corp.)	5'-CGCGAATTCATGGAAGACGCCAAAAAC-3'	5'-CGCAGATCTGCGGCCGCCTGCAGGATA TCTTACACGGCGATCTTTCC-3'	Firefly luciferase
pRL-tk (Promega Corp.) P3/119/70 subgenomic DNA clone	5'-CGCGAGCTCATGACTTCGAAAGTTTAT-3'	5'-CGCGTCGACTTATTGTTTCATTTTGTAG-3'	<i>Renilla</i> luciferase
P3/Sabin subgenomic DNA clone	5'-CGCCTGCAGTAAAAACAGCTCTGGGGT-3'	5'-CGCGAGCTCTGTGACACTGAAATCCTG-3'	Poliovirus type 3 119/70 IRES
pT7-5'NCR-X472 (ref. 33)	5'-CGCCTGCAGTAAAAACAGCTCTGGGGT-3'	5'-CGCGAGCTCTATGATACAATTGTCTGA-3'	Poliovirus type 1 5'NC-X472 IRES
pC1b (ref. 64)	5'-CGCCTGCAGCCGATTGGGGGCGACACT-3'	5'-CGCGAGCTCAGGATTCGTGCTCATGAT-3'	HCV 1b IRES
pAR100 (A. Rosenfeld and V. Racaniello, unpublished data)	5'-CGCCTGCAGCCGATTGGGGGCGACACT-3'	5'-CGCGAGCTCAGGATTCGTGCTCATGAT-3'	HCV 1a IRES Δ28–69
pP1/CVB3 (see Methods)	5'-CGCGTCGACTTAAAAACAGCTCTGGGGT-3'	5'-CGCGAGCTCTTGCTGATTCAACTTA-3'	CVB3 IRES

an excellent animal model for poliomyelitis, CD155 transgenic mice are not susceptible to poliovirus infection by the oral route, the natural means of infection in humans (52). We found that the tropism of recombinant polioviruses dependent on the IRES of CVB3 or HCV is unchanged in CD155 transgenic mice. These findings support the conclusion that a viral IRES can mediate translation in a wide range of organs and does not determine where poliovirus replicates and causes disease.

Previously, poliovirus dependent on the HCV IRES was reported to be cleared from the brain and spinal cord of adult mice without causing disease (13). These results led to the conclusion that the HCV IRES does not mediate translation initiation in the murine brain and spinal cord. An alternative explanation is that the recombinant virus replicates poorly (53) and is cleared by the immune system. We found that P1/HCV is cleared from adult CD155 transgenic mice but replicates in the brain and spinal cord of newborn CD155 transgenic mice and causes flaccid paralysis. Newborn mice are more susceptible than adults to infection with neurotropic viruses, including poliovirus (54–56), but the basis for the increased susceptibility to poliovirus infection is not known. Taken together with the observation that the HCV IRES mediates translation in multiple organs, the finding that P1/HCV replicates and causes paralysis in newborn mice indicates that the murine brain and spinal cord produce the proteins required for HCV IRES-dependent initiation.

A recombinant poliovirus dependent on the human rhinovirus 2 IRES is neuroattenuated in adult mice (12). A similar recombinant poliovirus is also neuroattenuated in newborn CD155 transgenic mice (data not shown). While the mechanism of attenuation is unclear, the HRV2 IRES may be dependent on cell proteins not present in the brain and spinal cord. This possibility is currently being explored.

Assertions that poliovirus tropism may be determined at the step of translation initiation (12–14) were in part based on previous studies of the mechanism of attenuation of the poliovirus live vaccine strains. The C472U mutation within the IRES of the Sabin type 3 vaccine strain attenuates neurovirulence (23) and was shown to cause a translation defect *in vitro* (28). The Sabin type 3 vaccine replicates poorly in a neuroblastoma cell line but not in HeLa cells, a phenotype attributed to the C472U mutation (30,

57). Poor replication in neuroblastoma cells was believed to be a consequence of reduced IRES-mediated translation (27). These observations led to the hypothesis that attenuating mutations in the Sabin vaccine IRES change poliovirus tropism by causing a translation defect specific to the brain and spinal cord (27, 30). To address this hypothesis, we measured IRES-mediated translation in organs and cells. Contrary to our expectation, the C472U mutation leads to identical translation defects in neuronal and non-neuronal organs and cells. This translation defect should lead to decreased viral replication in neuronal and non-neuronal organs. Our observation that the C472U mutation led to reduced IRES-mediated translation in all organs examined is consistent with the previous finding that this mutation reduces translational efficiency in extracts of non-neuronal cells (28). This observation is not consistent with previous conclusions that the effect of the C472U mutation is specific for cells of neuronal origin and does not decrease viral translation or replication in HeLa cells (27, 30, 57). In our study, variation in mRNA levels that could influence synthesis of the IRES-dependent reporter protein is controlled by the use of bicistronic mRNAs to quantify IRES-mediated translation. Furthermore, IRES-mediated translation might not be the rate-limiting step of viral replication in HeLa cells.

Mutations in the IRES are major determinants of the low pathogenicity of the poliovirus vaccine strains (22–24). The results reported here indicate that the C472U attenuating mutation reduces IRES-dependent translation in all tissues, not only the CNS as previously hypothesized. Furthermore, we find that the C472U mutation does not eliminate viral replication in the CNS, since viruses with this mutation are neurovirulent in newborn mice. How might the C472 mutation reduce poliovirus neurovirulence in humans, yet still allow for sufficient virus replication to ensure protective immunity? Poliovirus infection of humans begins in the alimentary tract, and disease of the CNS is rare (58, 59). Replication defects in the alimentary tract associated with C472U may decrease the probability that a sufficient number of virus particles will reach the brain and spinal cord to initiate a productive infection, without impairing the immunogenicity of the vaccine. Testing this hypothesis will require the development of a transgenic mouse that is susceptible to poliovirus infection by the oral route.



Our results show that polioviruses with the C472U mutation are neurovirulent in newborn but not adult mice. The basis for this difference in pathogenicity is not known. One possibility is that the immature immune system of the newborn mouse cannot effectively clear the virus infection, even though viral replication is compromised by the C472U mutation. Our findings emphasize that attenuation of poliovirus neurovirulence is not simply a matter of whether or not IRES-mediated translation occurs in the brain and spinal cord. The outcome of infection is determined by a complex interplay between virus and host.

Methods

Cells, plasmids, and viruses. S3 HeLa and A549 cells were grown in DMEM (Invitrogen Corp., Carlsbad, California, USA), 10% bovine calf serum (HyClone, Logan, Utah, USA), and 1% penicillin/streptomycin (Invitrogen Corp.). SY5Y and 293 cells were grown in the same medium except that 10% FBS (Invitrogen Corp.) was used. For plaque assays, HeLa cells were grown in DMEM (Specialty Media, Phillipsburg, New Jersey, USA), 0.2% NaHCO₃, 5% bovine calf serum, 1% penicillin/streptomycin, and 0.9% bacto-agar (Difco Laboratories Inc., Franklin Lakes, New Jersey, USA).

A bicistronic expression plasmid encoding firefly luciferase and *Renilla* luciferase (Figure 1A) was created. A firefly luciferase PCR product was cleaved with *EcoRI/BglII* and ligated to *EcoR/BglII*-cleaved pDC516 (Microbix Biosystems Inc., Toronto, Ontario, Canada). A *Renilla* luciferase PCR product was cleaved with *SacI/SalI* and ligated to a *SacI/SalI* cleavage of the above plasmid. PCR components are described in Table 2. The IRESs of poliovirus strains P3/119/70, P3/Sabin, and 5'NC-X472, HCV strain 1b, HCV strain 1a with nucleotides 28–69 deleted, and CVB3 were PCR-amplified with primers that add *PstI* and *SacI* sites to the 5' and 3' ends (Table 2) and ligated to *PstI/SacI*-cleaved bicistronic expression plasmid.

Plasmid pP1/HCV, an infectious poliovirus DNA clone in which nucleotides 108–745 are replaced with HCV sequence, was created as follows. Plasmid pBSSK was prepared from pBluescript II SK+ (Stratagene, La Jolla, California, USA) by removal of *SacI* and *KpnI* restriction sites and the T7 promoter. The T7 promoter and poliovirus nucleotides 1–107 were amplified by PCR from pT7M, a full-length clone of type 1 poliovirus strain Mahoney, with SK28 (5'-CGCGTCTGACTAATACGACTCACTATAG-3') and SK29 (5'-CGCGAATTCAAGTTACGGGAAGGGAGT-3'), cleaved with *SalI* and *EcoRI*, and ligated to *SalI/EcoRI*-cleaved pBSSK. Nucleotides 746–1,127 of the poliovirus genome were amplified by PCR from pT7M with SK30 (5'-CGCGAATTCGCGAGCTCAAAGGTCTCACAACATATGGTGCTCAGGTTTCATCA-3') and SK31 (5'-CGCGCGGCCGCGATATCAGCGACGTCTGTTCTGT-3'), cleaved with *EcoRI* and *NotI*, and ligated to *EcoRI/NotI*-cleaved pBSSK. Nucleotides 1,119–7,440 of the poliovirus genome were isolated from *AatII/SmaI*-cleaved pT7M and ligated to *AatII/EcoRV*-cleaved pBSSK. The HCV IRES was amplified by PCR from pC1b, cleaved with *EcoRI* and *SacI*, and ligated to *EcoRI/SacI*-cleaved pBSSK.

Plasmid pP1/CVB3, an infectious poliovirus DNA clone in which nucleotides 108–742 are replaced with CVB3 sequence, was created as follows. Nucleotides 743–1,127 of the poliovirus genome were amplified by PCR from pT7M with SK31 and SK90 (5'-CGCGAGCTCATGGGTGCTCAGGTTTCA-3') cleaved with *SacI* and *AatII*, and used to replace the 373-bp *SacI/AatII* fragment from pBSSK at the point prior to insertion of HCV sequence.

The CVB3 IRES was amplified by PCR from pCVB3-28 (60) with SK91 (5'-CGCGAATTCCTTAGAAGTAACACACAC-3') and SK92 (5'-CGCGAGCTCTTTGCTGTATTCAACTTA-3'), cleaved with *EcoRI* and *SacI*, and ligated to *EcoRI/SacI*-digested pBSSK. The sequences of all PCR products were determined and verified to be consistent with published versions.

Recombinant human adenoviruses were produced using the AdMax system (Microbix Biosystems Inc.). Viruses encoding bicistronic mRNAs were created by recombination in 293 cells between calcium phosphate-transfected adenovirus genome plasmid pBHGfrtΔE1,3FLP and pDC516-based plasmids that encode bicistronic mRNAs. Recovered virus was subjected to two rounds of plaque purification as described by the manufacturer (Microbix Biosystems Inc.). Virus stocks were established by infection of 293 cells, and virus was purified from cells by centrifugation onto a cushion of cesium chloride as described by the manufacturer (Microbix Biosystems Inc.).

To produce poliovirus strains P1/M, P1/HCV, and P1/CVB3, viral DNA clones pT7M, pP1/HCV, and pP1/CVB3 were linearized by restriction enzyme cleavage and used as templates for run-off transcription by T7 RNA polymerase (Promega Corp., Madison, Wisconsin, USA). RNA was transfected into HeLa cells using DEAE-dextran, and after 3 days intracellular virus was released by three freeze-thaw cycles and passed once in HeLa cells. Virus was subjected to two rounds of plaque purification, and virus stocks were produced in HeLa cells. Poliovirus strains PRV7.3, PRV8.4, P3/Sabin, and P3/119/70 were produced by transfection of HeLa cells with viral DNA clones using DEAE-dextran, and stocks were produced similarly. PRV7.3 and PRV8.4 are recombinant viruses comprising nucleotides 1–784 from P3/Sabin (PRV7.3) or P3/119 (PRV8.4) and the remainder of the genome from P2/Lansing (26). Human poliovirus receptor is not essential for infection of mice with P2/Lansing.

Assay for IRES-dependent translation in continuous cell lines. Monolayers of adherent cells were infected at a multiplicity of 10 PFUs per cell with adenoviruses encoding bicistronic mRNAs. After 24 hours, medium was aspirated, monolayers were washed with 1 ml PBS, and one-tenth of the cells were lysed in 1 ml passive lysis buffer (PLB; Promega Corp.). For Northern blot hybridization of the same infection (see “Northern hybridization analysis” below), RNA was extracted from the remaining nine-tenths of infected cells. The Dual-Luciferase Assay (Promega Corp.) and the Lumat LB 9507 luminometer (Berthold Technologies GmbH & Co., Oak Ridge, Tennessee, USA) were used to determine firefly luciferase and *Renilla* luciferase activity in lysates. To control for variation in infection or transcription, the ratio of firefly luciferase activity to *Renilla* luciferase activity was determined (IRES activity). Unless otherwise indicated, to control for IRES-independent *Renilla* luciferase translation, the activity of each IRES was normalized to the ratio determined in lysates from cells infected with an adenovirus lacking an IRES. The concentration of luciferase protein was calculated with reference to a standard curve generated using known concentrations of recombinant firefly luciferase (Fisher Scientific Co., Springfield, New Jersey, USA) and *Renilla* luciferase (Chemicon International Inc., Temecula, California, USA).

Northern hybridization analysis. Monolayers of adherent cells were infected at a multiplicity of 10 PFUs per cell with adenoviruses encoding bicistronic mRNAs. After 24 hours, medium was aspirated and total RNA isolated with TRIzol (Invitrogen Corp.). Total



RNA (15 µg) was fractionated in a 1% agarose formaldehyde gel and transferred to GeneScreen membrane (PerkinElmer Analytical and Life Sciences, Boston, Massachusetts, USA) according to standard protocols (61). RNA was UV-crosslinked to the membrane (Stratalinker; Stratagene) and baked 2 hours at 80°C under a vacuum. Blots were hybridized with a PCR DNA product of the *Renilla* luciferase coding region labeled with ³²P-dCTP by random priming (Rediprime; Amersham Biosciences, Piscataway, New Jersey, USA) according to standard protocols (62). For quantitation, a storage phosphor screen (Amersham Biosciences) was exposed to hybridized membranes for 1–6 days, scanned with the PhosphorImager 445SI (Molecular Dynamics), and analyzed with IQMac software (Molecular Dynamics).

Assay for IRES-dependent translation in murine organs. Four-week-old C57BL/6J mice (The Jackson Laboratory, Bar Harbor, Maine, USA) were injected intraperitoneally with 10⁹ PFUs recombinant human adenovirus for assays of heart, lung, liver, kidney, and ileum, intramuscularly with 5 × 10⁸ PFUs for assays of muscle, and intracerebrally with 5 × 10⁸ PFUs for assays of brain and spinal cord. The volume of all inoculations was 50 µl. Sixteen to 24 hours after infection, mice were sacrificed, and organs were removed and homogenized with a PowerGen 125 homogenizer (Fisher Scientific Co.) in 0.5 ml PLB. Crude protein extracts were prepared, and the Dual-Luciferase Assay and the Lumat LB 9507 luminometer were used to determine firefly luciferase and *Renilla* luciferase activity in the extracts. To control for variation in infection or transcription, the ratio of firefly luciferase activity to *Renilla* luciferase activity was determined (IRES activity). To control for IRES-independent *Renilla* luciferase translation, the activity of each IRES was normalized to the ratio determined in organs from mice infected with an adenovirus lacking an IRES. All experimental mouse protocols adhered to Institutional Animal Care and Use Committee (IACUC) guidelines and were approved by the IACUC of Columbia University Medical Center (New York, New York, USA).

Poliovirus replication in HeLa cells. Monolayers of adherent HeLa cells were infected at a multiplicity of 10 PFUs per cell. At indicated time points, cells were scraped into tubes and subjected to three freeze-thaw cycles to release intracellular virus. The virus titer in each sample was determined by infection of HeLa cell monolayers and plaque assay.

Infection of TgPVR mice with poliovirus. TgPVR mice transgenic for the human poliovirus receptor (40) were genotyped to ensure that they carried the human poliovirus receptor gene. Mice lacking the *pvr* gene were used as negative controls. Tail fragments were incubated overnight at 55°C in 0.2 ml of 50 mM KCl, 10 mM Tris (pH 8.3), 2.5 mM MgCl₂, 0.1 mg/ml gelatin, 0.45% NP40, 0.45% Tween-20, and 60 µg proteinase K (The Jackson Laboratory). Two microliters of the digestion product was used as template for PCR under standard conditions using 20A1C (5'-CTCAC-CACTGTACTCTAGTCTG-3') and 20A1W (5'-AGAAGGACT-CACTAGACTCAGG-3'). The human poliovirus receptor gene was indicated by a 350-bp PCR product.

To assay viral replication, the following inoculations were performed. TgPVR mice (either 4 weeks old or 1–2 days old) were inoculated intraperitoneally with 10⁵ PFUs or 10³ PFUs of poliovirus type 1 strain Mahoney in a volume of 50 µl or 15 µl. TgPVR mice (4 weeks old) were inoculated with 10⁷ PFUs P1/CVB3 in a volume of 50 µl. TgPVR mice (1–2 days old) were inoculated intraperitoneally with 10⁷ PFUs P1/HCV in a volume

of 15 µl. At the indicated time points, mice were sacrificed and organs were removed and homogenized with a PowerGen 125 homogenizer. Intracellular virus was released by three freeze-thaw cycles, and cellular debris was removed by centrifugation at 16,100 g for 15 minutes at 4°C. The titer of infectious virus in the supernatant of each sample was determined by infection of HeLa cell monolayers and plaque assay. To determine survival of infected mice, 15 TgPVR mice were inoculated with P1/CVB3 and 27 TgPVR mice were inoculated with P1/HCV as described above. Nine TgPVR mice were also inoculated with 10⁹ PFUs P1/HCV. Mice were observed daily for paralysis or death, and paralyzed mice were sacrificed immediately.

Neurovirulence assays. TgPVR mice were genotyped as described above, and mice were inoculated intracerebrally with tenfold serial dilutions of virus. Mice 1–2 days old were inoculated with 15 µl, and mice 4 weeks old were inoculated with 50 µl. Between 7 and 11 mice were inoculated with each dilution of virus. Mice were observed daily for paralysis or death, and paralyzed mice were sacrificed immediately. The 50% lethal dose (LD₅₀) value for each virus was calculated according to the number of dead and paralyzed mice at each dilution, following the method of Reed and Muench (63).

Nucleotide sequence of poliovirus RNA recovered from paralyzed TgPVR mice. Viral RNA was extracted from the brains of paralyzed mice as follows. TgPVR mice 1–2 days old were inoculated intracerebrally with 10⁵ PFUs PRV7.3, 5 × 10⁶ PFUs P3/Sabin, 10⁴ PFUs PRV8.4, or 3 × 10³ PFUs P3/119/70. Upon onset of paralysis 10, 7, 6, or 6 days after infection, respectively, mice were sacrificed and brains removed. Brains were homogenized in 1 ml PBS with 0.2% bovine calf serum and subjected to three freeze-thaw cycles to release intracellular virus. Cellular debris was removed by centrifugation at 16,100 g for 15 minutes at 4°C. Total RNA was extracted from 0.2 ml of supernatant with 1 ml TRIzol. RNA was also extracted directly from virus stocks to confirm the sequence at nucleotide 472.

The base at nucleotide 472 in viral RNA was determined by RT-PCR and restriction enzyme cleavage as previously described (44). First-strand cDNA was produced with 5 µg of total RNA using the ImProm-II reverse transcription system (Promega Corp.) and primer SK77 (5'-ACGACTTGCGCGTTACGACAGGCTG-GCTGC-3'). Five microliters of the reaction mixture was used in a PCR amplification with SK76 (5'-TGAGAGTCTCCGGCCCCCT-GAATGCGGCTGAT-3') and SK77. Approximately 0.4 µg plasmid pSK48 or pSK49 was used as PCR template in control reactions. PCR produces a 93-bp product, and the base C at nucleotide 472 creates an *Mbo*I restriction endonuclease site. This is cleaved to produce 61-bp and 32-bp products. Cleavage of 100% of *Mbo*I sites was never observed, most likely because of DNA damage during PCR amplification (Figure 7, lane 2). After incubation with *Mbo*I and electrophoresis in 10% acrylamide, DNA was transferred (Trans-Blot; Bio-Rad Laboratories Inc., Hercules, California, USA) to GeneScreen membrane. Blots were hybridized according to standard protocols (62) with the product of PCR amplification of pSK49 with SK76 and SK77 labeled with ³²P-dCTP by random priming. For quantitation, a storage phosphor screen was exposed to hybridized membranes for approximately 12 hours, scanned with the PhosphorImager 445SI, and analyzed with IQMac software. The proportion of 472U was calculated as the ratio of 93-bp DNA density to the sum of the 93-bp, 61-bp, and 32-bp DNA densities, according to a standard curve generated with plasmid DNA encoding the poliovirus IRES with 472U or 472C. The standard curve was linear from a proportion of 1–100% C at 472.



To determine the nucleotide sequence of the IRES in viral RNA recovered from paralyzed mice infected with PRV7.3, first-strand cDNA was produced with primer SK88 (5'-CGCGAGCTCTGTGACTGAAATCCCTG-3') and used in a PCR amplification with SK88 and SK103 (5'-ATAGCGGCCGCTTAAAACAGCTCTGGGGTTG-3') as described above. To determine the nucleotide sequence of the 2A^{Pro} coding region of viral RNA recovered from paralyzed mice infected with PRV7.3, first-strand cDNA was produced with primer SK107 (5'-GCGGCGGCCGCTTTGTGGGTGAATCCAC-3') and used in a PCR amplification with SK107 and SK106 (5'-CGCCTGCAGTCCCATACTACGGACCAG-3') as described above. Nucleotide sequence of PCR products was determined by the Herbert Irving Comprehensive Cancer Center DNA Facility, Columbia University Medical Center.

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