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Measles control – Can measles virus inhibitors make a difference?

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Abstract

Infection by measles virus (MV) is a major cause of human morbidity and mortality worldwide. In 2001, the WHO, UNICEF and their partners launched the Measles Initiative, the goals of which are to interrupt the transmission of MV in large geographic areas by increasing vaccination coverage and to assess the feasibility of eradicating MV worldwide. An estimated 74% reduction in mortality resulting from measles was achieved between 2000 and 2007, equivalent to a reduction of approximately 200,000 deaths annually. Despite this progress in the control of measles, the highest number of measles cases in more than a decade was observed in 2008 in several European countries and the US, and the virus was again declared endemic in the UK. In the light of this resurgence in the UK and the limitations associated with the current live-attenuated vaccine, this review discusses the means by which safe and effective measles antivirals could augment vaccination and strengthen global efforts to control measles. Important aspects of treatment are the potential to prevent infection effectively after exposure to MV, the improvement of case management, the amelioration of complications that frequently follow MV infection and the influence of antivirals on a potential strategy for global measles eradication.

Keywords

Antiviral; drug; eradication; measles; vaccine; virus

Introduction

Measles virus (MV) is a member of the *Morbillivirus* genus in the *Paramyxovirus* family. MV and other paramyxoviruses, such as mumps virus, respiratory syncytial virus, human parainfluenzaviruses and recently emerged zoonotic hendra and nipah viruses, constitute major pathogens for humans and animals [1]. All of these viruses are highly communicable airborne pathogens that spread via the respiratory route. Furthermore, MV is one of the most infectious viruses identified, with a basic reproduction number (R_0) of 12 to 18 [2-4], meaning that a single infection will cause 12 to 18 secondary cases in a fully susceptible population in the absence of intervention. High infectivity combined with the induction of long-lasting immunity protecting individuals against re-infection means that a population size of approximately 250,000 individuals or greater is required to ensure sufficient births of susceptible individuals to sustain continued MV transmission [5,6]. Because the virus has no non-human reservoir [2,3], MV can have emerged only after human populations reached this size approximately 5000 years ago [3]. As such, measles is a comparatively young human disease.

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MV as an infectious agent

Paramyxovirus particles possess a lipid envelope derived from the host-cell plasma membrane and a nonsegmented, single-strand RNA genome of negative polarity [7]. Paramyxovirus virions are of pleiomorphic shape with an average diameter of 150 to 300 nm. Inserted into the MV envelope are glycoprotein spikes (Figure 1), consisting of the attachment (H) and fusion (F) proteins, which mediate receptor binding, subsequent membrane fusion of the virus, and cellular entry. A helical nucleocapsid core consisting of the RNA genome and the nucleocapsid (N), phospho- (P) and large (L) proteins are tethered to the envelope by the matrix (M) protein (Figure 1B) [1,3].

Signaling lymphocyte activation molecule (SLAM) is a cellular receptor for all characterized MV strains [8,9]. Some laboratory-adapted and vaccine strains of MV can also use the regulator of complement activation (CD46) for efficient cellular entry [10-12]. Receptor binding triggers pH-independent fusion of the viral envelope with the target plasma membrane, followed by the release of the incoming nucleocapsid core into the target cell. Transcription of viral mRNAs and genome replication – the latter via a positive-sense, full-length antigenome – then occurs in the cytosol, mediated by the virus-encoded, RNA-dependent RNA-polymerase (RdRp) complex [13].

Similar to other members of myxovirus families, MV RdRp consists of the viral P and L proteins in addition to the N:RNA template [1]. The non-coding 5' and 3' termini of the viral genome contain cis-acting promoter and encapsidation signals that initiate the synthesis of mRNAs or antigenome, or genome replication [14]. The final assembly of newly synthesized nucleocapsid cores and glycoprotein complexes, enveloping, and budding of progeny viral particles occur at the plasma membrane of infected cells [1].

Multiple steps in the paramyxovirus life cycle are unique to the virus compared with the host cells, and thus constitute attractive targets for pathogen-directed antiviral therapies. Promising pharmacological approaches may include viral entry inhibitors that prevent receptor binding, envelope protein refolding or membrane merger required for infection, and inhibitors of RdRp that suppress viral mRNA synthesis and genome replication. Such viral mechanisms can potentially be targeted safely because human cells lack structural or functional homologs of these viral proteins.

MV pathogenesis and diseases potentially associated with MV infection

MV is transmitted by respiratory secretions from infected individuals, either person-to-person via larger respiratory droplets or airborne via aerosols. Initially after infection of a susceptible host, the replication of the virus is supported by dendritic cells in the respiratory tract, lymphocytes and regional lymphatic tissues [15]. This phase of localized replication is followed by primary (2 to 3 days after invasion) and secondary (5 to 7 days after invasion) viremia, with viral spread to multiple organs including the kidney, liver, gut and skin [2,3, 16] and predominantly to SLAM-positive lymphocytes and dendritic cells [17]. After an average incubation period of 10 to 12 days, clinical features of acute measles include a prodromal fever, a respiratory infection with rhinitis and severe cough, coryza, conjunctivitis, and pathognomonic enanthem [18]. The maculopapular rash that characterizes measles then appears approximately 2 to 4 days after the prodrome [3,19]. Shedding of the virus occurs from the nasopharynx from the onset of the prodrome until 3 to 4 days after the beginning of the rash [18].

Complications of severe measles infection can include acute demyelinating encephalomyelitis (ADEM) [20] and measles inclusion body encephalitis (MIBE) [19], which manifest soon (weeks to months) after infection. A lethal late complication, subacute sclerosing

panencephalitis (SSPE), can present years after the primary infection [3,19,21] and is largely untreatable with currently available therapeutics (see discussion in the *Current and experimental drugs for the management of measles* section). Worldwide, there were approximately 200,000 measles-related deaths in 2007, rendering the virus a major cause of human morbidity and mortality [22]. A prolonged state of immunosuppression of several months that follows acute cases of measles frequently predisposes patients to bacterial otitis media and bronchopneumonia [19]. Most deaths associated with measles are attributable to secondary viral, bacterial or parasitic infections that occur in this immunocompromised state [3,21].

In addition to direct complications and secondary infections associated with MV infection, a potential MV-related etiology has been discussed for a variety of persistent human diseases including rheumatoid arthritis [23] and multiple sclerosis [24-26]. A possible contribution of MV to lung cancer and Hodgkin's lymphoma has similarly been investigated [27-29]. However, a clear causal relationship between MV infection, and these and several other sequelae cannot be established based on the currently available data. That is, the link between the virus and numerous patient conditions is not supportable by the scientific evidence (for a comprehensive review, see reference [21]).

Measles vaccination and global control efforts

MV meets several prerequisites essential for possible global eradication, including an absence of a non-human reservoir, availability of accurate diagnostic tests and the existence of an effective vaccine [30]. In 2001, the WHO, UNICEF and their partners launched the Measles Initiative, the goals of which are to interrupt the transmission of MV in large geographic areas by increasing vaccination coverage and to assess the feasibility of the worldwide eradication of MV [31,32]. Since this program was established, the number of fatalities caused by worldwide measles was approximately 200,000 in 2007, an estimated 74% decrease compared with 2000 [22], and the virus is no longer considered endemic in the Americas [33].

Despite the successful application of the current live-attenuated measles vaccine [34], several factors contribute to the ongoing morbidity and mortality associated with MV worldwide. First, a trend of increasing measles case numbers has appeared in several industrialized countries, mostly as a consequence of elective exemption from vaccination because of personal or parental philosophical or religious beliefs [35,36]. As a result of its high infectivity, measles is one of the first diseases to reappear when vaccination rates decline [35]. Although in the European region (as defined by the WHO) the incidence of measles was reduced from 8223 cases in 2006 to 3909 in 2007 [36], the highest numbers of cases in more than a decade were observed in several European countries in 2008 [35,37,38]. The UK alone reported 1217 cases from January to November 2008 [39], placing it together with Romania, Germany, Switzerland and Italy as one of the European countries with the most cases of measles [40]. In June 2008, the virus was again declared endemic in the UK [39,41], 14 years after it had been eliminated, and it appears unlikely that the goal of elimination in Europe by 2010 can be achieved [40]. In the UK in particular, parental concerns about vaccination safety were heightened by a report that associated the trivalent measles-mumps-rubella (MMR) vaccine with the onset of autism and intestinal disease [42]; a claim that has since been demonstrated to be unsubstantiated [43-45]. When accompanied by a decline in public awareness of the disease, a scenario of waning immunity based primarily on philosophical beliefs cannot be excluded for the US. The number of cases of measles in the US also reached a 10-year high in 2008, mainly as a result of greater transmission after importation of the virus [35].

Second, due to the high communicability of the virus, a susceptibility of 5 to 6% of an otherwise highly vaccinated population is sufficient to sustain periodic outbreaks [4,46]. A 'herd

immunity' of greater than 95% [47] is thus required for complete suppression of the virus, which cannot be achieved with a single dose of the vaccine [46,48]. While a second vaccination is routinely administered in developed countries [49,50], there are greater logistical obstacles to repeated vaccination in the developing world. In particular, the current live-attenuated vaccine requires an uninterrupted cold-chain [18], sterile materials and professional healthcare workers for administration [51].

Third, the efficiency of vaccination in infants younger than 12 months of age is compromised by the immaturity of their immune systems and interference from transplacentally acquired maternal antibodies [51-53]. Because maternal antibody titers vary, the immunity of infants to MV is frequently lost at 4 to 9 months of age, creating a window of susceptibility for infection by MV prior to vaccination [51].

Furthermore, concerns have been raised that efforts to eradicate MV might be compromised in the long term by waning protection of the adult population because immunity against the attenuated vaccine strain is less durable than that acquired naturally [54-56]. While currently available data indicate that this might not constitute a major obstacle to the control of MV [30,57,58], the concern of waning immunity could undermine public confidence in vaccination against MV.

Potential role of antiviral therapeutics in the management and control of MV

A novel platform combining prophylactic (vaccination) and therapeutic (antiviral) approaches could overcome both currently encountered and possible future obstacles to the control of MV. However, such an approach would require the identification of antivirals that are safe and effective against MV.

Antiviral drugs can facilitate the rapid control of local viral outbreaks in industrialized and developing countries through post-exposure prophylaxis of the immediate, non-immunized contacts of identified cases in the family and community settings, such as in childcare centers and schools. The long incubation period of MV prior to the onset of viremia offers a large potential window for antiviral treatment. In particular, rapid pre-emptive antiviral treatment could suppress the development of disease entirely in naïve individuals or reduce the severity and longevity of clinical symptoms if naïve individuals are infected before they are vaccinated (either because the vaccination is refused or unavailable). Conversely, long incubation periods may compromise therapeutic potential because clinical disease follows the peak of virus replication. However, the reduction of MV-induced respiratory distress by treatment with ribavirin (see discussion in the Current and experimental drugs for the management of measles section) [59-61], provides some support for the hypothesis that antivirals administered at the stage of clinical disease may improve the management of severe cases. It is possible that such treatment might reduce the disease burden and ameliorate complications and, conceivably, antivirals might also open novel therapeutic options for the treatment of late measles sequelae such as SSPE.

The use of antivirals against MV could, furthermore, contribute to removing the 'window of susceptibility' that is present when maternal antibody titers decrease in infants, by post-exposure prophylaxis prior to vaccination. Preventing this type of transmission would be of special importance in areas of the developing world where the virus is endemic, and in those industrialized countries where declining herd immunity has resulted in the return of endemic MV transmission.

Several logistical advantages render small-molecule antivirals particularly suitable for rapid application in the developing world: large scale production strategies are generally well-established and typically highly cost-effective; small molecules can frequently be optimized

to achieve high shelf-stability at ambient temperatures (unlike the MMR vaccine that requires uninterrupted cold-chains); and compounds can be formulated for aerosolized or oral bioavailability, allowing rapid mass administration. The field strategy for the advanced stages of global MV eradication proposed by the Pan American Health Organization predicts the need for 'mop-up' campaigns to target susceptible children in difficult-to-reach sites of viral outbreaks [62]. Safe and effective antivirals could constitute a desirable additional component of such campaigns because these drugs could provide immediate control of local outbreaks before the trained personnel and sterile materials required for vaccination are available on site.

The HIV/AIDS pandemic is reducing the effectiveness of vaccination for MV; increased rates of failure of both primary and secondary measles vaccination [63] and prolonged shedding of MV have been reported in children infected with HIV [64]. It has been suggested that the high mortality rate among HIV-positive children in the developing world might preclude the formation of a sufficiently large group of MV-susceptible individuals to sustain transmission of MV [2]. However, improved access to antiretroviral therapy may lead to an increased need for effective measles therapy.

The Global Polio Eradication Initiative demonstrated that in the endgame of viral eradication, outbreaks in countries with remaining endemic transmission – in the case of poliovirus, Nigeria, India, Afghanistan and Pakistan [65] – can result in viral spread across several continents [66]. It is pertinent to consider the different routes of transmission and basic reproduction numbers of MV and poliovirus: MV undergoes airborne transmission and has an R_0 value of 12 to 18 whereas the transmission of poliovirus is fecal-oral and the R_0 value is 5 to 7 [67]. Therefore, sporadic outbreaks of MV comparable to those with poliovirus at a very late stage of eradication could lead to even more rapid viral spread across large geographic areas. The relevance of this scenario will likely be increased once the virus is considered to be nearly eliminated. At this time, the resources in industrialized countries will be sufficient to continue vaccination programs [2,30,51]. This must be offset, however, against the possibility of a rapid erosion of public acceptance and global political will to maintain global vaccination against an 'eradicated' pathogen. Thus, a back-up antiviral prophylaxis strategy to immediately curb local outbreaks through stockpiled, shelf-stable antivirals until mop-up vaccination responses can be implemented in the area would be desirable and likely boost public confidence.

Current and experimental drugs for the management of measles

Currently, no therapeutics for the treatment of measles are available. Ribavirin (which is approved for the treatment of some paramyxovirus infections) and IFN α therapies have been tested clinically against MV, mostly for the treatment of patients presenting with SSPE. Although some studies noted a beneficial effect of IFN α [68,69], the majority of reports documented either a long-term relapse [70-72] or lack of efficacy [73-76]. High-dose ribavirin treatment, either alone or in combination with IFN α , appeared to be more efficacious against MV than IFNα alone [59-61,70,77,78]. Although some studies noted a gradual progression of measles despite therapy [72,74] or a lack of efficacy [73], these reports nevertheless suggest that antivirals might ameliorate the complications of measles even when administered after the onset of clinical symptoms. However, severe side effects, most notably hemolytic anemia, have been attributed to ribavirin when used in combination with pegylated IFN α for the treatment of viral hepatitis C [79,80]. Treatment with high-dose vitamin A has been associated with some reduction in the morbidity and mortality of measles [81], but this effect is most pronounced in children younger than 2 years of age [82,83]. For post-exposure prophylaxis, the administration of high-titer MV-specific immune globulin (Ig) within 6 days of exposure can prevent or modify disease [84]. Such treatment is recommended for temporary protection of the immunocompromised and infants younger than 1 year of age [18]. However, Ig therapy is comparatively expensive, requires sterile materials and an uninterrupted cold-chain, and is

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overall not recommended or feasible across an entire population for the control of large measles outbreaks.

Considering these mixed reports of efficacy and the additional limitations associated with Ig and ribavirin therapies, the development of novel, safe and efficacious inhibitors of MV is required for a combined prophylactic and therapeutic anti-measles platform. Other desired features of a measles antiviral are cost-effective mass production, shelf-stability and the potential for oral or aerosolized delivery.

A variety of different antiviral strategies for MV inhibition have been considered, including antisense molecules, peptidic inhibitors, natural extracts, nucleoside analogs and smallmolecule compounds. Vector-based antisense inhibitors [85,86] and peptide-conjugated morpholino oligomers [87], although effective in *in vitro* models of MV infection, have high production costs, limited storage stability, and delivery and bioavailability issues. Related concerns apply to peptidic inhibitors of MV entry, such as di- and tri-peptides (that are moderately effective in in vitro models of MV infection) [88,89] and highly potent peptides derived from the conserved heptad repeat B domain of the viral fusion protein [90]. The latter peptides act by a mechanism of membrane fusion inhibition considered analogous to that of the efficacious HIV-1 entry inhibitor enfuvirtide (Fuzeon) [91,92] and are potent, with active concentrations in the nanomolar range (EC₅₀ = $0.1 \,\mu$ g/ml). However, because experience with the 36-residue peptide enfuvirtide has also highlighted several obstacles associated with heptad-repeat-derived peptidic antivirals (subcutaneous injection is required, injection-site reactions occur and the cost of therapy is approximately US \$24,000 per year) [93], such inhibitors are unlikely to be a viable strategy for measles therapy and, therefore, have not been pursued for further development.

Multiple natural extracts or synthetic analogs derived from natural products have been reported to possess anti-MV activity [94-103]. However, many of these substances were only moderately active in cell culture, were cytotoxic, were inactive when added to cells post-exposure or the active ingredient remained elusive (reviewed in reference [104]).

Synthetic small-molecule compounds are by their nature likely to constitute the most suitable class of MV inhibitor because these compounds have a greater potential than other approaches to be mass produced cost effectively, to be stable at ambient temperature and to have desirable bioavailability. Table 1 provides an overview of the classes of MV inhibitor that were investigated previously or are currently under investigation. To minimize potential side effects, a pathogen-directed rather than host-directed antiviral strategy appears preferable for the treatment of measles. For instance, the entry and transcription/replication phase of the virus life cycle, driven by unique viral protein complexes that lack cellular homologs, constitute attractive targets. However, a pathogen-directed approach is at a greater risk for emerging viral resistance. While resistance to treatment is in general a particular challenge for the treatment of persistent viral infections, it must be investigated whether this applies equally to therapy for a pathogen such as MV that causes predominantly acute disease and induces strong immunity. Treatment-resistant variants may be of little clinical relevance if they coincide with reduced efficiency of viral transmission.

A newly developed class of non-nucleoside, target-specific inhibitors of MV polymerase complex activity demonstrated potent antiviral activity, in the nanomolar range, when tested against a panel of wild-type MV isolates representing currently endemic genotypes [105]. Combined with overall desirable pharmacological features of the scaffold (Figure 2), high chemical stability and low cytotoxicity (selectivity index $CC_{50}/EC_{50} \sim 16,500$ [106]), this class is an example of a novel developmental lead that is tailored to the specific requirements for a measles therapeutic.

Conclusions

The Measles Initiative launched in 2001 has made impressive progress towards reducing the global morbidity and mortality caused by measles. This was achieved through a substantial increase in coverage with the live-attenuated measles vaccine, in particular in Africa, and the Eastern Mediterranean and Western Pacific regions [32]. Despite these achievements, the highest numbers of measles case in more than a decade were reported in 2008 for several industrialized countries, and the virus was again declared endemic in the UK. This resurgence, which results mostly from individual or parental reservations against vaccination based on philosophical or religious beliefs, highlights the challenges associated with a decline in public acceptance of vaccination [107]. Further complications arise from the limitations associated with the currently available vaccines as described in detail in the previous section.

In addition to improvements in the control of MV by new vaccines [51,53], the development of novel, safe and cost-effective small-molecule measles antivirals can contribute to overcoming these limitations. Rather than creating an alternative approach, MV inhibitors can be a useful addition to the prophylactic options available against MV, thereby providing a combined prophylactic and therapeutic anti-measles platform. Conceivable areas of immediate use of antivirals include acute and persistent disease (to improve case management), post-exposure prophylaxis, rapid control of local outbreaks before vaccinations become available or in cases of declined vaccination and protection of immunocompromised individuals and infants prior to vaccination. In the long term, antivirals could assist in a prolonged endgame of global eradication, as experienced with poliovirus.

Suitability for these applications sets clear parameters for ideal measles antivirals. The desired drug is safe and effective, mass producible at low cost, characterized by high shelf-stability at ambient temperature and is orally available. None of the experimental measles therapies tested clinically thus far matches this diverse array of requirements, necessitating *de novo* development. Small-molecule antiviral compounds are considered to be best suited to meeting these criteria. To date, promising chemical scaffolds have been identified experimentally that have demonstrated potent antiviral activity. Broadening the current drug discovery efforts and moving promising new hits and current leads through preclinical development is warranted to enable effective antivirals to become clinically available.

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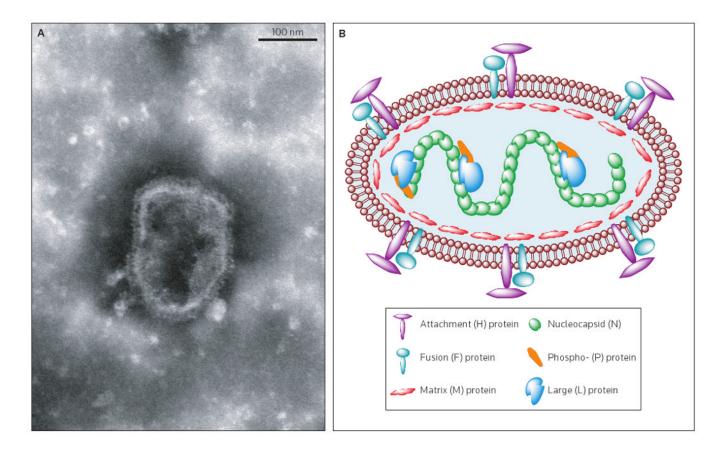


Figure 1. Measles virus, a member of the Paramyxovirus family

(A) Electron micrograph of purified MV particles at a magnification of 100,000×. Virions are pleiomorphic with an approximate diameter of 150 to 300 nm. Glycoproteins embedded in the viral envelope are detectable (Image was taken by Brindley MA, Wang JJ and Plemper RK). (B) Schematic representation of an MV particle showing the six structural viral proteins. Not shown are non-structural C and V proteins that are encoded in the P gene and accessed through alternative ribosome initiation and pseudotemplated RNA editing, respectively. Only a small number of envelope glycoprotein complexes are shown for clarity.

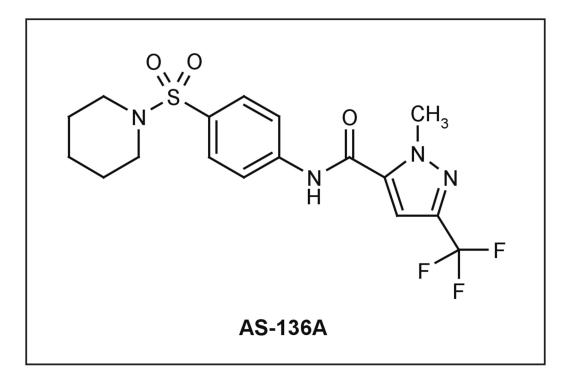


Figure 2. The structure of AS-136A

AS-136A is a specific inhibitor of measles virus RNA-dependent RNA polymerase activity with desirable pharmacological properties.

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Compound or class	EC ₅₀ concentration	Specificity	Proposed target, if known	Minimal efficacious dose	Reference
Neplanocins	0.1 to 1 μg/ml	Broad-range	S-adenosylhomocysteine hydrolase	ND	[108, 109]
Noraristeromycins	0.4 µg/ml	Broad-range	S-adenosylhomocysteine hydrolase	ND	[110,111]
5'-Nor carbocyclic adenosine analogs	< 0.4 µg/ml	MV-specific	ND	ND	[112]
5'-Fluoro-5'-deoxyaristeromycin	13 µM	MV-specific	S-adenosylhomocysteine hydrolase	ND	[113]
5-Ethynyl-1- β -D-ribofuranosylimidazole-4-carboxamide (EICAR)	0.1 to 1 μg/ml	Broad-range	IMP dehydrogenase	120 mg/kg bid	[109,114]
Ribavirin	17.6 to 47 μg/ml	Broad-range	Lethal mutagenesis	360 mg/kg bid	[114-116]
Ring-expanded nucleoside analogs	2 to 10 µM	MV-specific	ND	ND	[117]
Mycophenolic acid	0.2 µg/ml	Broad-range	IMP dehydrogenase	ND	[116]
Coumarin analogs	0.2 to 50 µg/ml	MV specific	Possibly RdRp	ND	[118]
N-phosphonacetyl-L-aspartate (PALA)	8 μg/ml	Broad-range	L-aspartic acid transcarbamoylase	ND	[119]
Pathogen-associated molecular patterns (PAMPs)	0.5 μg/ml	Broad-range	Possibly virus attachment	No therapeutic effect	[114]
Isoquinolin analogs	10 µM	MV-specific	ND	ND	[120]
AS-48	0.6 to 2 µM (wild-type isolates)	MV-specific	MV fusion protein	Inactive	[121-123]
Isothiazole analogs	2.9 μg/ml	Broad-range	ND	ND	[124,125]
AS-136A	10 to 50 nM (wild-type isolates) MV-specific MV RdRp complex	MV-specific	MV RdRp complex	ND	[105,106,126]
When available, postulated targets of the different compounds	, measles virus (MV)-specificity ve	ersus broad-ran	compounds, measles virus (MV)-specificity versus broad-range antiviral activity (viruses of the paramyxovirus and/or other virus families), and the	ramyxovirus and/or other vir	us families), and

minimal efficacious dose in the cotton rat model of MV infection are provided. IMP inosine monophosphate, ND not determined, RdRp RNA-dependent RNA polymerase