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Rotavirus immune responses and correlates of protection

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Abstract

Selected topics in the field of rotavirus immunity are reviewed focusing on recent developments that may improve efficacy and safety of current and future vaccines. Rotaviruses have developed multiple mechanisms to evade interferon-mediated innate immunity. Compared to more developed regions of the world, protection induced by natural infection and vaccination is reduced in developing countries where, among other factors, high viral challenge loads are common and where infants are infected at an early age. Studies in developing countries indicate that rotavirus-specific serum IgA levels are not an optimal correlate of protection following vaccination, and better correlates need to be identified. Protection against rotavirus following vaccination is substantially heterotypic; nonetheless, a role for homotypic immunity in selection of circulating post vaccination strains needs further study.

INTRODUCTION

We and others have previously reviewed immunity to rotaviruses (RV) [1] and correlates of protection for RV vaccines [2–4]. Here, we will focus on recent and selected topics in the field with emphasis on issues related to improving vaccine efficacy. We will first analyze mechanisms used by RV to evade the innate immune response, mainly based on data from cell culture studies and the mouse model. Second, we will review new data on protection induced by natural infection in humans. Third, we will analyze the correlation of serum IgA levels induced by the two currently licensed safe and effective live RV vaccines with protection [5,6], and finally, we will examine the importance of heterotypic versus homotypic immunity.

Mechanisms used by RV to modulate innate immunity

Recently, a number of studies examining the capacity of RV to evade the innate immune response, and in particular the interferon (IFN) response, have been published. Using non-purified preparations of intestinal epithelial cells from mouse pups, RV infection was found to induce type I (IFN- β) and type III IFN (IFN- λ) mRNA expression [7]. Moreover, treatment of mice with IFN- λ , and to a lesser extent type I IFN, reduces virus replication [7]. In agreement with these findings, mice lacking the receptor for type III IFN, and to a

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lesser degree the receptor for type I IFN, have higher levels of viral replication than wild-type mice, demonstrating the importance of the IFN system in modulating viral replication [7]. Thus, it is not surprising that the RV has developed multiple mechanisms to evade the type I IFN response, many of them through the action of the viral protein NSP1 [8,9].

A simplified model of the mechanism used by RV to evade the IFN response is presented in Figure 1. Viral replication is recognized by RIG-I (see Glossary box) and/or MDA-5, which in turn activate IRF3 and NF- κ B. These transcription factors then induce expression of type I IFN and interferon-stimulated genes (ISGs) [10,11]. RV NSP1 prompts degradation of IRF3 and inhibition of NF- κ B activity [8,9,12]. These effects are complex and depend on viral strain and cell type. For example, NSP1s from some animal RVs degrade IRF3, IRF5 and IRF7; in contrast, NSP1 of human RVs predominantly degrade IRF5 and IRF7 and may, therefore, less efficiently inhibit the IFN response [13]. The degree of extra-intestinal spread and replication of RV appears, at least in part, to be dependent on the capacity of a RV strain to inhibit the IFN system in specific cells: In suckling mice without an intact IFN signaling system, some heterologous strains of RV replicate very efficiently in the biliary tree and pancreas and cause biliary atresia and pancreatic disease, whereas homologous murine strains do not [14]. RV replication in the murine biliary tract is determined by both viral entry, mediated by VP4, and viral evasion of the innate immune response, mediated by NSP1 [15,16]. After RV induction of transcription of IFN, secretion of this cytokine is modulated by PKR [11]. In an autocrine fashion, IFN stimulates type I IFNR expression; IFNR then signals through STAT1/STAT2/IRF9 complexes to further induce IFN and ISG expression and an antiviral state (Figure 1). RV has also been shown to inhibit the function of STAT1 and STAT2 by an unknown mechanism and thus, can potentially inhibit all types of IFN responses [17]. RV has also been shown to stimulate the secretion of TGF- β in polarized Caco-2 cells and this effect has been shown to inhibit the capacity of dendritic cells to activate Th1 T cells [18,19]. It is unknown whether this mechanism occurs *in vivo* but if so, it could potentially explain the poor T cell response to RV seen *in vivo* [20].

Protection induced by natural infection

Development of the RV vaccines currently in use was based on the observation that natural infection can protect against severe RV-induced acute and recurrent gastroenteritis (RVGE). Cohort studies carried out in Mexico [21] and Guinea-Bissau [22] showed that recurrent episodes of RV disease were less severe than the first episode; one episode of RV infection had a protective efficacy of 70% [22] or 77% [21] against RV-induced diarrhea. Two infections, either symptomatic or asymptomatic, protected 100% against moderate to severe disease of any serotype [21]. These observations were challenged, however, by recently published results on a cohort of young children in Vellore (India), in which the severity of diarrhea did not significantly decrease between the first and second infections, but did between the second and third infections; protection after one episode of RV infection was 43% against RVGE, and it took three infections to induce 79% protection against moderate or severe RVGE [23]. In this Indian cohort study there was an earlier occurrence of primary RV infection, with 53% of children infected by 6 months of age, in contrast with 34% and 26% of children infected by that age in Mexico and Guinea-Bissau, respectively. The occurrence of an early primary infection is affected by likelihood of year-round RV exposure and environmental viral loads, with high loads [24] leading to RV infection very soon after the waning of transplacentally acquired maternal antibodies. The early infection of children in the poorest countries of Asia and Africa may also have an effect on vaccine efficacy in countries with high year-round RV infection prevalence rates: Children in the placebo arms of studies in these settings have high rates of RV-specific IgA seroconversion, and these high rates are significantly and inversely correlated with protection (Supplementary Table 1). Given that the capacity to induce neutralizing antibodies to RV is

age dependent [25], early infection of children may not induce an efficient protective immune response and this fact may contribute to the diminished efficacy of natural infection and vaccination to induce high levels of protection in the poorest countries of Asia and Africa.

Serum IgA as a correlate of protection for RV vaccines

Like a number of other oral vaccines, both licensed RV vaccines, the monovalent G1P[8] Rotarix™ vaccine (RV1) [26] and the bovine reassortant pentavalent RotaTeq™ vaccine (RV5) [27,28], are clearly less efficacious in the poorer countries of Africa and Asia than in the Americas and Europe. Protection during the second year following vaccination with RV5 decreases substantially compared to the first year (Table 1). Efficacy of RV1 was not significantly increased with a three-dose immunization schedule compared to the standard two-dose regime [26]; this was contrary to expectations as three natural infections results in significant protection relative to two in some settings [23].

Total serum RV-IgA measured shortly after vaccination has been used as “a measure of vaccine take” and is probably the best (although imperfect) correlate of protection [1]. In previous vaccine studies in Latin America, North America and Europe levels of serum RV-IgA “take” were close to the levels of protection afforded against severe GE for RV5 and against GE of any severity for RV1 (Table 1). In contrast, RV-IgA seroconversion rates generated by RV1 and RV5 were clearly lower in poorer countries of Africa and Asia than in Latin America, North America and Europe, where RV1 was not co-administered with OPV, although there are significant differences among these poorer countries (Table 1). The few studies measuring immunogenicity and efficacy of RV1 and RV5 administered with or without concomitant administration of OPV suggest that concomitant administration of OPV does not significantly interfere with the efficacy of RV vaccines but does somewhat reduce their immunogenicity [29–31]. Analyzing results from Table 1 from settings where OPV was co-administered with RV vaccines (excluding the Mali trial with unreliable data on protection), a significant correlation does not exist between RV-IgA seroconversion and protection against severe RVGE during the first year after vaccination ($p>0.3$).

Relative importance of heterotypic and homotypic immunity in protection against RV

The development of homotypic versus heterotypic immunity in people after natural infection or vaccination is a complex and incompletely understood phenomenon [2,4]. In Mexican infants followed prospectively for two years, protection from illness was both homo- and heterotypic in nature with a somewhat stronger, but certainly not exclusive, homotypic response appearing after the first RV infection [21]. In contrast, little or no decrease in the risk of homotypic or heterotypic RV infection or diarrhea was observed in Indian infants after primary infection, although protection did evolve after second or third infections. The basis for these differences is not clear at present [23].

The desire to elicit broad heterotypic protective immunity in children has been a major determinant in the development of the two widely licensed RV vaccines. Controlled clinical trials of RV vaccines in middle income and industrialized countries have not provided conclusive information on the relative importance of each type of immunity, due in part to the limited serotypic diversity of circulating “challenge” strains observed during the actual clinical trials [2]. However, the high levels of protection from severe disease generated by monotypic RV1 indicate that immunization of young children with a single strain of RV provides substantial protective immunity from infection with multiple other serotypes [6,32]. To date, it does not appear as if immunization with a polyvalent vaccine in these populations offers any significant advantage. Moreover, the lack of correlation between the rather low levels of serotype-specific serum neutralization responses and the high levels of

protective efficacy induced by either RV1 or RV5 implies that protective immunity is not likely to be exclusively homotypic and may not be directly linked to homotypic neutralizing antibody, at least in the serum [5].

The results of recent efficacy studies in Africa and Asia, where a high diversity of RV genotypes circulate and infection with multiple serotypes are common, have shed more light on this problem. In the clinical trials performed with RV1 in Malawi and South Africa, 67.7% and 64.1%, respectively, of RV strains that infected children in the placebo group were homotypic G1 or P[8]. The levels of protection induced by RV1 against severe RVGE caused by non-G1 versus G1 strains were similar in Malawi, higher in South Africa and lower in South America (Table 2). Thus, RV1 induces significant protection from severe disease with multiple G and P serotypes not included in the vaccine, supporting the conclusion that immunity to RV has a substantial heterotypic component, at least after enteric immunization with a replicating virus.

The percentage of children with serotype-specific neutralization responses after RV5 co-administered with OPV in Latin American infants was similar to those detected in the United States and Finland, where infants received inactivated polio vaccine (IPV), and lower in infants from Africa and Asia (Table 3). Similar to the results from RV5 studies in middle income and industrialized countries, the levels of protective efficacy induced by this vaccine against each RV genotype were higher than the respective rates of serum neutralizing antibodies (SNA) (Table 3). Of note, most cases (89–100%) of severe RVGE in African and Asian infants vaccinated with RV5 were caused by RV with G or P genotypes, or both, which are covered by the RV5 vaccine; this indicates that the lower efficacy of RV5 in these countries is not due to disease caused by RV strains not included in the vaccine. The similar low efficacy levels in Africa and Asia for both RV vaccines (monovalent and pentavalent) strongly support the conclusion that heterotypic protection is an important component of the protective immune response. The mechanism by which heterotypic immunity is induced following homotypic immunization is still unclear, but possibilities include induction of protective antibody responses to heterotypic epitopes on VP4 or VP7, possible protective effects of antibody to common antigens on VP6 or NSP1 or cross reactive protective T cell responses.

Nonetheless, several new studies suggest a role for homotypic immunity against RV. Modeling of RV genotype variations over time in developed countries are compatible with a role for homotypic immunity [33]. Moreover, detailed analysis of pre-challenge serum samples from selected adult volunteers experimentally challenged with a G1 RV showed that the quantity of both homotypic and heterotypic antibody responses significantly correlated with protection, but that the strongest correlation was with the level of homotypic response [34]. Although RV1 has been clearly shown to protect against the heterotypic G2P[4] strain even after 2 years [35], recent studies in Brazil [36] and Australia [37] suggested that protection against this strain may not be long-lived under some settings. The importance of this finding needs to be confirmed, because opposite findings have been suggested by another study from Brazil [38]. Finally, studies in Brazil [39–41] and developed countries [42–44] suggest that vaccine introduction may result in the selection of serotypically distinct new strains. To date, however, the two vaccines have remained highly effective in preventing severe RV disease in developed countries and large numbers of vaccine selected “drifted” strains have not emerged. Of note, several recent studies from the US have convincingly demonstrated an unexpected elicitation of “herd immunity” following vaccination [45]. Whether this effect might eventually play a role in selecting rotavirus drift variants or in the generation of heterotypic immunity is unknown.

CONCLUDING REMARKS

New insights into the mechanism used by RV to evade the immune response suggest that a detailed functional analysis of the VP4 and NSP1 proteins of RV1 and RV5 might enhance our understanding of why vaccine efficacy is not optimal and how the safety profile can be improved to allow administration to neonates as has been proposed by some investigators [46].

In children from developing countries, high environmental challenge loads, vaccine interference by maternal antibodies [47,48] and concomitant administration of OPV may in part explain the lower immunogenicity and protective efficacy of the current vaccines compared to developed countries. Withholding breastfeeding prior to vaccination and use of IPV might increase the levels of RV vaccines immunogenicity and protection, and studies to test these interventions should be undertaken.

RV-IgA levels are a poor correlate of protection in developing countries (Table 1). In most RV5 studies, serum IgA conversion rates exceeded protection rates against severe RVGE; this suggests that, as in the case of other Jennerian vaccines based on heterologous animal RV strains, IgA levels may not reflect intestinal immunity [2]. Identifying new correlates of protection that better reflect intestinal immunity may help in designing more efficacious third-generation RV vaccines [49].

RV1 and RV5 have similar efficacy against severe RVGE in countries where a high diversity of strains co-circulate, suggesting an important role for heterotypic protective immunity. However, indirect evidence suggests that homotypic immunity also plays a role in protection against subsequent RV infection. Characterization of RV strains present in the environment post-vaccination is needed to rule out population-based selection of “escape” strains due to long-term pressure of homotypic immunity.

Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

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Glossary box and abbreviations

IFN	Interferon
IFNR	Interferon receptor
IRF	Interferon regulatory factor. A family of transcription factors that regulate the interferon response
ISG	interferon-stimulated genes
MDA-5	Melanoma differentiation-associated gene-5. Member of the RIG-I-like receptor family of cytoplasmic RNA helicases that sense viral RNA and trigger innate immunity
NF-κB	Nuclear transcription factor (NF)- κ B. NF- κ B plays a central role in the cellular stress and inflammatory responses that modulate innate immunity to pathogens

PKR	Double-stranded RNA-dependent protein kinase. Its gene is inducible by IFN and is a key player in the antiviral actions of these cytokines
RIG-I	Retinoic acid inducible gene I (RIG-I). Member of the RIG-I-like receptor family of cytoplasmic RNA helicases that sense viral RNA and trigger innate immunity
SNA	Serum neutralizing antibodies
STAT	Signal transducers and activators of transcription (STAT). A family of cytosolic transcription factors whose activation depends on several growth factors and cytokines including interferon
TGF-β	Transforming growth factor beta. A potent regulatory cytokine key in maintaining immune tolerance and in prevention of immunopathology
TH1 cells	T cells that produce interferon gamma and in general play a role in immunity against intracellular pathogens including viruses

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Highlights

Rotaviruses have developed multiple mechanisms to evade IFN innate immunity

Protection induced by natural infection and vaccination is lower in developing countries

Studies in developing countries indicate that serum IgA is not an optimal correlate of protection for RV vaccines

Protection against RV is highly heterotypic

A role for homotypic immunity in selection of post-vaccination “escape” strains needs further study

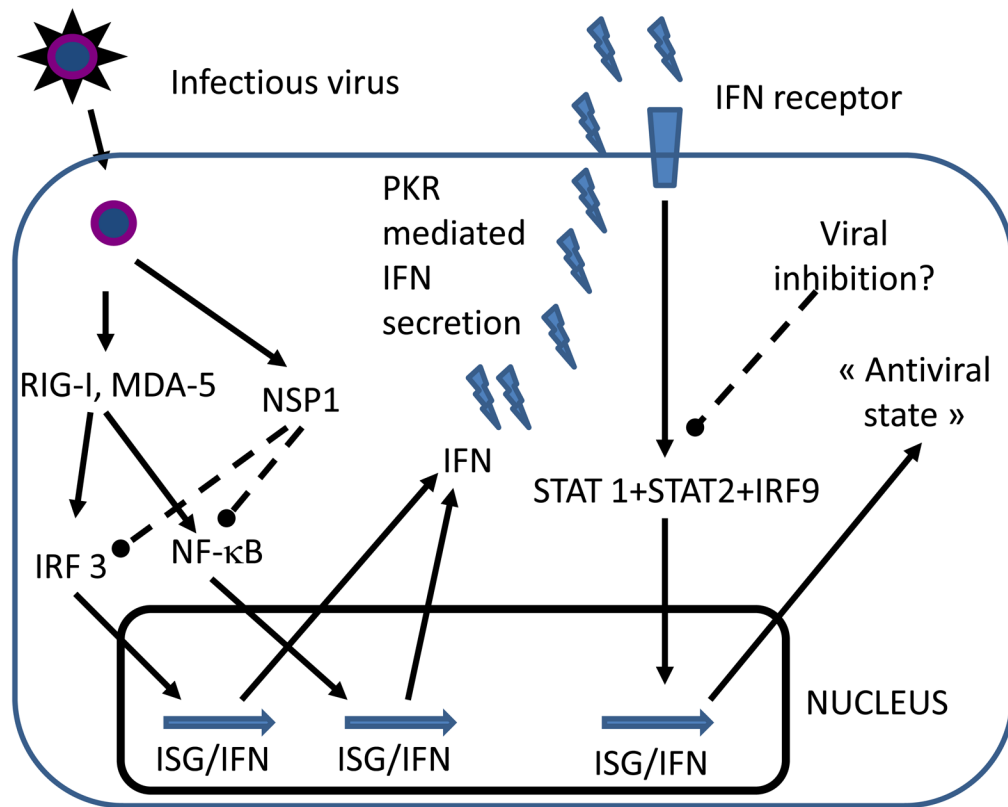


Figure 1. Modulation of the host innate immune system by RV. After viral entry, viral replication activates the pathogen recognition receptors RIG-I and MDA-5, which in turn activate the transcription factor IRF3 and NF- κ B. These molecules translocate to the nucleus, where they induce the transcription of ISGs and IFN. Viral replication produces NSP1 that can induce proteasomal degradation of IRF3. Through NSP1 dependent and independent mechanisms RV may also block NF- κ B translocation. After transcription of IFNs, the dsRNA-dependent protein kinase PKR modulates IFN secretion by an unknown mechanism. Autocrine secreted IFN signals through the IFN receptor to activate transcription factors STAT1, STAT2 and IRF9. These factors translocate to the nucleus and further increase the levels of ISG and IFN transcripts, establishing an “antiviral state”. RV has also been shown to block STAT1 and STAT2 through an unknown mechanism.

Table 1

Vaccine efficacy and RV-IgA seroconversion rates in selected trials

Vaccine/Ref	Location	Protection against severe RVGE year 1 % (95% IC)	Protection against severe RVGE year 2 % (95% IC)	Protection against RVGE of any severity % (95% IC)	IgA Seroconversion % (95% IC)	OPV
RV1 [26]	Malawi	49.2 (11.1 to 71.7)	NR	34.7 (7.7 to 53.5) ^{&}	47.2 (30.4 to 64.5)	Yes
RV1 [26]	South Africa	72.2 (40.4 to 88.3)	NR	64.2 (52 to 73.4) ^{&}	57.1 (44.7 to 68.9)	Yes
RV1 [31]	Latin America*	81.6 (54.4 to 93.5)	NR	NR	61.4 (53.7 to 68.6)	Yes
RV1 [35]	Europe**	95.8 (89.6 to 98.7)	85.6 (75.8 to 91.9)	87.1 (79.6 to 92.1)	86.5 (83.9 to 88.8)[‡]	No
RV1 [6,50]	Latin America	84.7 (71.7 to 92.4)	79 (66.4 to 87.4)	NR	84.7 (71.7 to 92.4)	No
RV5 [28]	Ghana	65 (35.5 to 81.9)	29.4 (-64.6 to 70.7)	30.5 (16.7 to 42.2) #	78.9 (67.6 to 87.7)	Yes
RV5 [28]	Kenya	83.4 (25.5 to 98.2)	-54.7 (-1752.7 to 82.3)	30.5 (16.7 to 42.2) #	73.8 (60.9 to 84.2)	Yes
RV5 [28]	Mali	≈	≈	≈	82.5 (70.1 to 91.3)	Yes
RV5 [27]	Bangladesh	45.7 (-1.2 to 71.8)	39.3 (-18.3 to 69.7)	42.5 (21.1 to 58.4) +	78.1 (66 to 87.5)	Yes
RV5 [27]	Vietnam	72.3 (-45.2 to 97.2)	64.6 (-47.7 to 93.9)	42.5 (21.1 to 58.4) +	97 (89.6 to 99.6)	Yes
RV5 [5]	Finland and United States	98 (88.3 to 100)	88 (49.4 to 98.7)	74 (66.8 to 79.9)	95.2 (91.2 to 97.8)	No

* Argentina, Brazil, Colombia, Dominican Republic, Honduras and Panama.

** Czech Republic, Finland, France, Germany, Italy, and Spain.

≈ Efficacy studies were unreliable due to problems with follow up of infants.

NR= not reported

[&] Results from groups that received 2 or 3 doses of RV1 were pooled.

[#] Results for Ghana, Kenya and Mali were pooled.

⁺ Results from Bangladesh and Vietnam were pooled.

[‡] GlaxoSmithKline, Clinical study registers Study ID. #102247-036

Available at: <http://www.gsk-clinicalstudyregister.com/> Accessed 9 January 2012.

Direct comparisons between studies of the same vaccine in different geographic and demographic settings are difficult to make, due to differences in study designs, as well as differences in the serotype and inoculum load of circulating rotavirus strains and other environmental conditions.

Table 2

Efficacy of RV1 against severe RVGE caused by G1 and non-G1 strains in selected trials

Trial location	% G1 protection (95% IC)	% Non-G1 protection (95% IC)	Reference
South Africa *	69.8 (32.5 to 87.1)	85.9 (55.1 to 96.6)	[26]
Malawi *	43.7 (<0 to 85.7)	50.3 (17.4 to 70.0)	[26]
South America	100 (<0 to 100.0)	80.6 (51.4 to 93.2)	[31]

* Results are pooled from children receiving two and three doses of RV1.

Table 3

Serum neutralizing antibody (SNA) seroresponse rates and vaccine efficacy (Prot) against severe RVGE in selected RV5 studies.

Strain Serotype	Bangladesh and Vietnam OPV [27]		Ghana, Kenya and Mali OPV [28]		Latin America OPV [30]		US and Finland No OPV [5]	
	% SNA	% Prot	% SNA	% Prot	% SNA	% Prot	% SNA	% Prot
G1	32.1	46.2	18.5	32.3	59.9	NR	76	74.9*
G2	9.9	29.2	9	27.1	36.3	NR	34.3	63.4*
G3	28.2	67	6.3	62.3	29.3	NR	22.1	82.7*
G4	18.3	NR	26.5	NR	50.9	NR	54.1	48.1*
PIA [8]	27.5	49.7	14.4	36.1	43.9	NR	53.7	NR

* Clinical efficacy calculated for RVGE of any severity.

NR = not reported.