Rotavirus infection induces both interferon (IFN) and inflammatory resistance by re-programming intestinal IFN receptor signaling

Adrish Sen[1,2]**, Nima D. Namsa**[1, 2, 3]**, Ningguo Feng**[1,2]**, Harry B. Greenberg**[1,2]

The Department of Microbiology and Immunology, Stanford University School of Medicine, Stanford, California, USA[1]; VA Palo Alto Health Care System, Palo Alto, California, USA[2], Department of Molecular Biology and Biotechnology, Tezpur University (A Central University), Napaam 784 028, Assam, India[3]

<u>Abstract (250 words)</u>

Rotaviruses (RV) cause acute severe diarrhea and are highly infectious in homologous

host species. However, RV infection does not cause appreciable intestinal inflammation

and homologous RV replication resists the antiviral effects of different interferon (IFN)

types. In suckling mice, we found that homologous murine EW RV triggers sustained

induction of IFNs, but not the amplification of interferon-stimulated genes (ISGs) or

inflammatory transcripts. IFNs can trigger both apoptosis and inflammatory necroptotic

cell death pathways. However, during RV infection intestinal epithelial cells (IECs) are

severely impaired in the cleavage of different caspases regulating intrinsic and extrinsic

cell death. Inhibition of caspase cleavage occurs in both RV-infected and -uninfected

bystander IECs. In contrast, RV induces cleavage of receptor-interacting protein 1 (RIP1),

a master regulator of cell death and survival decisions. Comparison of IFN-resistant

(homologous) and -sensitive (heterologous) RV strains in mice demonstrates that

successful RV infection restricts IFN receptor (IFNR)-amplified intestinal inflammatory

cytokine secretion. LPS-mediated intestinal damage, driven by IFNR-induced

inflammatory necroptosis, is also prevented during ongoing homologous RV infection. We

find that the ability of RV to negate IFNR-mediated antiviral and inflammatory functions

may involve a two-pronged attack. First, RV infection induces resistance to IFNAR1-

directed STAT1 activation in bystander cells and inhibits IFNR-feedforward amplification.

Second, ectopic stimulation of different IFNRs during RV infection results in efficient

negative feedback transcription, eliminating several host innate responses to infection.

Our study reveals a novel mechanism underlying the ability of homologous RV to

dismantle the antiviral and inflammatory arms of several intestinal IFN types.

<u>Significance</u>

Interferons mediate distinct functions including amplification of antiviral and inflammatory genes and inflammatory necroptosis of cells. Here we show that rotavirus (RV) infection of small intestinal epithelial cells (IECs) subverts IFNR-mediated antiviral and inflammatory functions. Although RV sensitizes both infected and uninfected bystander IECs to acute necroptosis, it prevents intestinal inflammation and induces cleavage of a key necroptosis regulator, RIP1K. During RV infection, uninfected bystander cells resist IFN-mediated STAT1 phosphorylation and feedforward transcription is blocked. In addition, ectopic stimulation of IFNRs during RV infection triggers negative-feedback signaling and eliminates several host antiviral and inflammatory responses. Thus, perturbation of IFNR feedback, recently linked to IFN resistance in cancer and inflammatory pathologies, is also exploited by an acute viral pathogen, RV.

Introduction

Rotaviruses are non-enveloped segmented dsRNA viruses that cause severe dehydrating diarrhea in many mammals, resulting in an estimated 200,000 human infant deaths annually(1). Acute RV infection in a homologous host (the natural species in which the RV strain replicates to high titer and spreads from individual to individual) is an extremely efficient process; severe diarrhea can be initiated by exposure to a very small infectious inoculum (2). The host innate interferon (IFN) response is a multi-pronged line of defense against viral pathogens as its induction can activate hundreds of different antiviral genes and trigger inflammatory signals – thus eliminating infected cells, limiting the spread of virus infection to uninfected bystander cells, and shaping both the magnitude and quality of subsequent adaptive immune response (3-5). The induction of all three major IFN types (α/β, γ, and λ) is mounted during homologous RV infection of the small bowel(6-10), but the antiviral effects of the IFN response on homologous RV replication are modest at best(10). Although poorly studied, IFN-mediated inflammatory responses are also likely reduced by RV infection, which is associated with only mild intestinal inflammation(11, 12) compared to other acute diarrheal pathogens(13). How RV efficiently subverts the ability of different IFN types to execute their antiviral and inflammatory functions(14) is not well understood.

On the other hand, infection with *heterologous* RVs (RV not typically isolated from the infected host species) generally results in severely restricted intestinal replication and little or no sustained transmission to uninfected hosts(11, 15, 16). In suckling mice, the intestinal replication of certain heterologous RV strains is substantially restricted (~10^4-fold) but can be partially rescued by deleting IFN receptors or the downstream convergent

transcription factor STAT1, thus unmasking the key role of IFN in suppressing RV

infectivity in heterologous host species(7, 10, 11, 16). Depletion of IFNRs in suckling mice

can also result in an acute biliary inflammatory disease (often lethal) following infection

with the heterologous rhesus RV strain (RRV)(11). Thus, identifying the mechanisms by

which homologous RVs suppress host IFN antiviral and inflammatory responses in the

gut is vital for developing more effective strategies to combat RV-associated morbidities

and mortalities, and in understanding the pathogenic potential of naturally circulating RV

strains.

Results

Homologous RV infection restricts intestinal IFN amplification responses. We

compared the intestinal viral replication and antiviral responses between a replication-

restricted, IFN-sensitive simian strain (RRV), and a replication-competent, IFN-resistant

murine RV strain (EW)(10) at 12 and 72 hpi by qRT-PCR (Fig 1A-B). Both RV strains

comparably induced the three major IFN-type transcripts (IFN-α/β, IFN-γ, and IFN-λ, Fig.

1A) at 12 hpi. Similar *induction* of IFN pathways by both RV strains is supported at the

protein level by measurement of intestinal IFN-γ, which is produced in both EW and RRV

RV-infected mouse pups at 24 hpi (Fig. 1C). Several IFN-stimulated genes (ISGs) were

also transcriptionally induced at 12 hpi with similar efficiencies by both EW and RRV

(*IRF7*, *ISG15*, *IFIH1*, *DDX58*, *IRF9*, *IFIT2*, and *MX2*).

Despite inducing IFNs and various ISG transcripts with a similar or lower efficiency,

intestinal EW RV NSP5 transcript levels were markedly higher than RRV at 12 hpi (by

$>10^4$-fold, Fig. 1B). Surprisingly, at 72 hpi the induction of several IFN types and ISG

transcripts continued to be comparable in mice infected with either murine or simian RV

(Fig. 1A and Fig. S1). However, EW RV replication at 72 hpi remained substantially higher

than RRV RV (by ~10^4-fold, Fig. 1B and Fig. S2). The only exception to these findings

was *IL28A* (encoding IFN-λ2), which was significantly higher (~10 fold) in the EW RV-

infected pups at 72 hpi. In terms of kinetics, the induction of type I IFNs (*IFNA4* and

IFNA5) and *IRF7* transcripts by both RV strains significantly increased between 12 to 72

hpi while induction of IFN type II (*IFNG*) and type III (*IL28A*) declined during the infection

time course. Of note, the 12 hour-induction of several interferon-stimulated gene and

inflammatory transcriptional responses in EW RV infected pups was virtually eliminated

by 72 hpi (*IL6*, *CXCL10*, *ISG20*, *IFI203*, *PELI1*, *IFIH1*, *DDX58*, *IRF9*, *IFIT2*, 72 hpi, Fig.

1A) resembling IFN amplification responses to replication-incompetent RRV RV. Thus,

intestinal IFN amplification responses to EW and RRV do not mirror the marked

differences in actual viral replication between these two RV types. A likely explanation for

these results is that despite substantial continued EW RV replication in the gut and the

induction of IFN genes during the first 3 days of infection, homologous murine RV has

suppressed the amplification of intestinal ISGs and inflammatory genes by 72 hpi.

Homologous rotavirus replication suppresses cleavage of several caspases in the

intestine. The results above revealed that several IFNs continue to be transcriptionally

induced in a sustained manner over 72 hrs during homologous RV infection. Exposure of

cells to IFNs (IFN-α/β and -γ) in a sustained manner generally results in a sequential

cleavage of the initiator caspase 8 and the executioner caspase 3, promoting

apoptosis(17-20). How homologous RV contends with the potential for caspase-mediated

cell death while inducing different intestinal IFN types (Fig. 1A, 1C) in a sustained manner

is not well studied. Therefore, we next examined the expression of precursor (full-length,

FL) and cleaved (Cl) forms of the initiator caspase 8, which is critical for extrinsic cell

death(21), in murine EW RV-infected mice. Immunoblotting of small bowel IECs revealed

that RV infection results in a substantially lowered ratio of Cl: FL caspase 8 isoforms (2.2

in mock *vs.* 0.5 in RV infected at 24 hpi, but not at 3 hpi) (Fig. 2A). The defective cleavage

of caspase 8 (Fig.2A) would be expected to impair subsequent cleavage of the apoptotic

executioner caspase 3, so we next examined the status of caspase 3 during RV infection.

As anticipated, intestines from murine RV infected suckling mice also exhibited

decreased cleavage of caspase 3 (Cl.: FL ratios of 1.4 and 0.4 in mock and EW RV

infected, respectively, at 24 hpi but not 3hpi) (Fig. 2B). These unexpected results reveal

that homologous RV infection inhibits intestinal caspases at the step of their zymogen

activation.

Cleavage of caspases 8 and 3 is inhibited in both rotavirus-infected and uninfected

bystander villous enterocytes. Having observed that RV infection *in vivo* results in

impaired caspase-8 and -3 processing in IECs, we next examined whether this process

occurs specifically in the small intestinal villous epithelium(7). Using flow cytometry, we

gated live villous IECs, using a previously described combination of cell surface markers

(i.e. CD45⁻ESA⁺CD44⁻)(7, 22), and then measured expression of cleaved forms of either

caspase 8 or caspase 3 in the RV infected and mock control pups (Fig. 2C). These

experiments confirmed a substantial decrease in caspase 8 and caspase 3 cleavage in

small intestinal villous enterocytes during RV infection (~75% decreased caspase 8 and

caspase 3 cleavage following infection).

We next examined whether caspase cleavage inhibition occurs solely in RV-infected

mature villous IECs at 24 hpi using a panel of antibodies to define cell type, caspase

activity, and the RV VP6 positivity (as a marker for RV infection status of cells) (Fig. 2D).

Remarkably, efficient suppression of caspase-8 cleavage occurs in both VP6$^+$ (infected),

and in VP6$^-$ (i.e. uninfected 'bystander') mature villous cell population. The relative ratios

of cells staining positive for cleaved caspase 8 in the different populations were: ~3.0

(mock), 0.2 (bystanders), and 0.6 (RV infected) (Fig. 2D, top panels). The efficient

inhibitory effect of RV infection on uninfected bystander (i.e. VP6$^-$) cells was also

observed following analysis of caspase 3 cleavage using a similar approach (Fig. 2D,

relative caspase 3 cleavage ratios were 4.0, 0.3, and 0.9 in mock, VP6$^-$ bystander, and

VP6$^+$ RV-infected cells, respectively; Fig. 2D, bottom panels).

RV inhibition of caspase 8 and 3 cleavage in both RV infected and uninfected

small bowel IECs would be expected to exert a significant impact on intestinal cell viability.

Therefore, we examined cell death in two intestinal cellular compartments. First, viability

analysis of total small intestinal single cell suspensions (i.e. 'bulk' level) revealed a

modest overall decrease in intestinal cell death during EW RV infection (this decline was

not statistically significant) (Fig. S3A-B). Second, a significant reduction in cell death was

observed by TUNEL staining of cells in the intestinal non-hematopoietic (i.e. CD45$^-$)

compartment, which includes both epithelial and stromal cells(7), but not hematopoietic

cells (Fig. S3C, p<0.001). Collectively, these results demonstrate that RV infection *in vivo*

effectively prevents caspase cleavage and suppresses intestinal cell death in both

infected and non-infected bystander cells.

In contrast to caspase 8-mediated extrinsic apoptosis, the initiator caspase 9 is

cleaved during intrinsic apoptosis (23). Interestingly, suckling mice infected with EW RV

also exhibited impaired cleavage of caspase 9 (Fig. 2E), as indicated by an accumulation

of intestinal FL caspase 9 at 16hpi, but not 6hpi, when compared to either RRV or mock-

infected controls (Fig. 2E). To examine the role of IFN in this process, we determined

whether caspase 9 cleavage is blocked in cultured 3T3 fibroblasts infected with the bovine

RV strain UK, which we previously showed triggers MAVS-dependent secretion of IFN-β

in this cell line(24, 25). Infection of 3T3 cells with UK RV for 9 h suppressed caspase 9

cleavage and mediated an accumulation of the FL-caspase 9 zymogen. In contrast,

increased Cl-caspase 9 and decreased FL-caspase 9 expression occurred at 16 hpi (Fig.

2F). These results confirmed that inhibitory effects of RV on intrinsic initiator caspase 9

activity do not require inhibition of IFN secretion and further indicated that caspase

cleavage could be regulated temporally during RV infection.

Kinetic regulation of caspases by RV was further examined in an intestine-derived

cell line by time-course analysis of caspase 8 cleavage in cultured HT-29 cells following

infection with porcine RV strain SB1-A, which we previously showed prevented IFN-β

secretion in this cell line(26). Compared to a mock-infected control, infection with SB1-A

RV inhibited caspase 8 cleavage starting at 4 hpi but with maximal effect occurring at 8

hpi (Fig. 2G). Collectively, these studies indicate that RV broadly inhibits initiatory

caspase cleavage events essential for either extrinsic or intrinsic cell death outcomes and

regulates these processes temporally during the course of infection. The results also

demonstrated that the RV regulatory effects on apoptotic signaling both *in vitro* (Fig. 2F-

G) and *in vivo* (Fig. 2A-E) occur regardless of its ability to suppress IFN induction.

Several RV strains induce cleavage of the cell death regulator RIP1K. In response

to sustained IFN exposure, caspase 8 cleavage results in apoptotic epithelial cell

shedding with typically mild inflammatory consequences(20). In contrast, in cells that are

defective in caspase 8 cleavage (such as IECs during EW RV infection *in vivo*, Fig. 2A-

C), IFN stimulation generally directs a rapid and precipitous form of cell death by

necroptosis that is acutely inflammatory(17, 27). Receptor interacting protein 1 kinase

(RIP1K) is a crucial adaptor dual-function kinase at the crossroads of a cell's decision to

live or die in response to either apoptotic or necroptotic stimuli(28, 29). These opposing

functions of RIP1K are primarily regulated by a proteolytic mechanism(30, 31). While

overexpression of full-length human RIP1K triggers both apoptosis and necroptosis,

caspase-mediated cleavage of the ~75 kDa RIP1K (p75) at multiple sites within the kinase

domain (KD) and interaction domain (ID) generates shorter fragments derived either from

the RIP1K N-terminal (51 kDa, 37 kDa, and 25 kDa) or the C-terminal (39 kDa) regions.

Both p51 and p39 RIP1K potently inhibit p75 RIP1K function and caspase 8 cleavage,

preventing apoptotic and necroptotic cell death in response to sustained inflammatory

stimuli(31).

As RV regulates caspase 8 cleavage in a temporal manner during infection (Fig. 2F-

G), we examined whether RIP1K cleavage is also subjected to RV regulation. HT-29 cells

were infected with representative RV strains differing in their modes of IFN regulation *in*

vitro (IRF3-degrading simian RRV, NF-kB inhibitory porcine SB1-A and human WI-61,

and NSP1-mutant simian SA11-5S RV strain that stimulates IFN-β secretion(32)) for

either 4 or 12 h, and then analyzed by immunoblotting using an anti-(N-terminal) RIP1K

antibody (Fig. 2H). When compared to mock-infected HT-29 cells, infection with the

different RV strains resulted in enhanced p75 RIP1K expression at 4 hpi with varying

efficiencies, and this increased expression was also be detected later during infection (at

12 hpi). In addition, RV infection also generated several shorter p51/p37/p25 RIP1K

cleavage fragments, and strain-specific differences were apparent in the levels of these

cleavage fragments. Notably, all RV strains including SA11-5S, triggered p51-RIP1K

cleavage starting early during infection, with the exception of the human WI-61 RV strain

infection where increased p51 RIP1K expression is primarily detectable by 12hpi.

Moreover, simian (RRV and SA11-5S) and human (WI-61) RV strains induced additional

cleavage of RIP1K (p37/p25 RIP1K - RRV and SA11-5S; p37 RIP1K - WI-61).

Interestingly, SB1-A RV induces increased expression of both the (pro-death) full-length

p75 RIP1K and the shorter (inhibitory) p51 RIP1K isoforms (Fig 2H, lanes 5-6). We thus

examined how functionally antagonistic RIP1K (p75 and p51) fragments that are

expressed during SB1-A infection influence cell death. We found that both cell adherence

and phosphatidylserine expression (a marker for apoptotic cells) at 16 hpi in SB1-A RV-

infected cells resembled mock-infected HT-29 cells, suggesting that increased p75 RIP1K

expression does not trigger cell death. In contrast, HT-29 cells treated with a cocktail of

tumor necrosis factor and cycloheximide exhibited substantial loss of cell adherence and

increased apoptotic staining (Fig. 2I). These results reveal that RIP1K accumulation and

its proteolysis are regulated during RV infection, and a shorter dominant-negative p51

regulatory RIP1K isoform is generated by many RVs early in infection that could

potentially impair apoptosis and necroptosis.

Rotavirus restricts intestinal IFN-mediated inflammatory responses. In addition to

promoting IFNR-mediated and caspase-dependent inflammatory death, different IFN

types can also drive the production of inflammatory cytokines by an IFNR-directed

positive feed-forward gene amplification process(3, 33). Therefore, we next examined

whether, in addition to inhibiting caspase activation *in vivo*, homologous RV infection of

suckling mice also suppresses the synthesis of IFNR-stimulated inflammatory target gene

products. To address this possibility, we used an experimental design similar to Fig. 1A

to measure production levels of different intestinal cytokines at 1 dpi in suckling mice

infected with either homologous murine (EW strain) or heterologous simian (RRV strain)

RVs (Fig. 3A). These studies reveal significant increases in intestinal production of a

variety of cytokines in response to either homologous EW and/or heterologous RRV RV

infection (RANTES, IP-10, IL1-A, EOTAXIN, GRO-A, MCP-3, MIP-1B, IL-4, GM-CSF, IL-

15, LIF; Fig. 3A). Of these, a set of cytokines (MCP-3, IL-15, LIF, IL-10, MCP-1, MIP-2α,

and IL-18) were increased comparably in response to either replication-competent murine

(EW) or replication-restricted simian (RRV) RV. In contrast, a second set of predominantly

pro-inflammatory cytokines (RANTES, IP-10, IL1-α, Eotaxin, GRO-α, MIP-1b, IL-6, IL-4,

GM-CSF, and IL-17A) were primarily induced in the heterologous simian RRV-infected

pups. These cytokines correlate with the poor intestinal replication of heterologous RVs

in mice, although whether they contribute in any manner to RRV growth restriction

remains to be determined. Strikingly, several differentially secreted cytokines in this set

are reported to be IFN-α/β, IFN-γ, and/or IFN-λ mediated effectors (i.e. RANTES(34-36),

IP-10(35, 37), Eotaxin(38), GRO-α(39, 40), and MIP-1b(36, 41, 42)). This suggests that

successful intestinal replication of homologous murine RV is accompanied by a

suppression of IFNR-amplified inflammatory cytokine responses(14).

In order to confirm whether homologous murine RV suppresses IFNR-amplified

inflammation, we used a suckling mouse model of endotoxemia, in which LPS-induced

inflammation and severe intestinal damage are reported to occur primarily through IFNR-

mediated inflammatory cytokine expression(43-45) and necroptotic cell death(17, 27, 46).

In prior work, we found that murine RV infection of the small bowel protected suckling

mice from endotoxin-induced mortality(8). However, whether or not EW RV replication

exerts a local protective effect in the LPS-challenged murine small bowel was not

evaluated. Here, we specifically examined if EW RV infection prevents small intestinal

inflammatory damage during endotoxemia (Fig 3B). Suckling mice orally infected with EW

RV (or mock) for 72h were systemically administered endotoxin and resulting intestinal

damage examined 6h later. In mock-infected mice, parenteral administration of endotoxin

led to severe intestinal necrosis, injury, and tissue destruction after 6h, as expected (Fig.

3C)(47). In contrast, suckling mice infected with murine EW RV (for 3d) prior to endotoxin

challenge exhibit minimal small intestinal damage (Fig. 3C). These results are consistent

with the conclusion that homologous RV effectively prevents IFN receptor-amplified small

intestinal inflammatory damage due to necroptosis or inflammatory gene expression. The

results also indicate, somewhat surprisingly, that efficient caspase 8 inhibition in the small

intestine during EW RV-infection (Fig. 2, C, D) does not sensitize mice to LPS-induced

inflammation.

Different RV strains potentiate IFN resistance in uninfected HT-29 bystander cells.

Cleaved caspase 8 is a key negative regulator of IFN (and LPS)-mediated necroptosis

and acute inflammation(17), and IEC-specific caspase 8 deletion results in spontaneous

intestinal inflammation(48), possibly triggered by constitutive tonic IFN signaling(27, 46).

Inhibition of caspase 8 cleavage is thus considered to be a potent priming event for acute

IFN-directed necroptotic inflammation in the gut(17, 21, 46). Although murine RV infection

reduces the cleavage of caspase 8 isoforms in both infected and bystander IECs (Fig.

2D), and induces various intestinal IFNs *in vivo*(7, 9) (Fig. 1A), these sequelae do not

translate into appreciable ISG amplification, intestinal inflammation or tissue damage

(Figs. 1A, 3A-C), cell death (Fig. S3), or viral growth restriction (Fig. 1B)(7, 10, 11). To

better understand this apparent paradox, we examined whether RV infection might block

IFN receptor (IFNR) signaling in both infected and bystander cells in order to effectively

prevent IFN effector functions. We previously reported that in cell culture, several RV

strains mediate efficient lysosomal degradation of endogenous IFNAR1, IFNGR1, and

IFNLR1 subunits in individual RV+ infected cells, but not in uninfected RV- bystander

cells(8).

We postulated that RV might employ a different strategy to also disable IFNR

functions in uninfected bystander cells. In previous *in vitro* studies, we showed that a

porcine RV strain (SB1-A) (but not several other RV strains) inhibited IFNR-mediated

STAT1-Y701 phosphorylation in RV- bystander cells(26), following a relatively short (i.e.

30-minute) stimulation with high concentrations of purified IFNβ (i.e. up to 2,000 IU/ml).

Here, we more extensively examined whether other RV strains might also suppress IFN

responsiveness in bystander cells, but with a lower efficiency than the SB1-A porcine

strain. This re-examination used an extended administration time for exogenous IFN

stimulation prior to STAT1-pY701 analysis, thus better accounting for possible lowered

efficiencies of IFNR inhibition by other RV strains (Fig. 4A). Human HT-29 cells were

infected with several RV strains (simian RRV and SA11, porcine SB1-A, bovine UK, and

simian SA11 mono-reassortant SOF, encoding a porcine NSP1) for 12h and cells

subsequently stimulated for 6h with purified IFN-β (500 IU/ml) before analysis of STAT1-

pY701 expression in individual RV-VP6$^+$ and VP6$^-$ HT-29 cells. Stimulation of mock-

infected cells with IFN-β activated STAT1-Y701 phosphorylation and this activation was

effectively inhibited by all tested RV strains in RV infected-VP6$^+$ cells as we previously

reported(26). More importantly, STAT1-Y701 activation was also significantly inhibited in

RV-VP6$^-$ bystander cells by the other RV strains (Fig. 4A). These results thus reveal that

all the RV strains tested inhibit IFN receptor signaling in bystander cellular populations *in*

vitro, although their relative efficiencies appear to vary across the strains.

As different IFNR types normally expressed in RV bystander cells(8) are likely to

be functionally impaired (Fig. 4A), we next examined how RV inhibition of IFNR-mediated

STAT1 phosphorylation influences downstream antiviral and inflammatory gene

transcriptional functions. Several ISGs and inflammatory transcripts upregulated by

IFNR-ligand stimulation can also be induced if cells are stimulated with intracellular

dsRNA, primarily through NF-kB and IRF3 signaling pathways(49). Thus, here we

compared the ability of RV to regulate cellular ISG and inflammatory transcriptional

amplification following stimulation with either exogenous IFN-β or intracellular long

dsRNA (Fig. 4B). Control HT-29 cells stimulated with IFN-β or dsRNA triggered both

overlapping (*TNF, TNFAIP3, HERC5, IFI6, IFIT3, IFIT1, IFNB1, RSAD2*) and unique

(*TLR2, IL26, IL21, SERPING* by IFN-β; *IL29, IFNG* by dsRNA only) sets of transcriptional

responses. Regardless of how transcriptional stimulation was carried out, RV infection

effectively prevented further transcriptional upregulation of *HERC5, IFNG, IL22, IL21,*

IFI6, IFIT1, IFIT3, and *RSAD2*. In contrast, we identified a set of pro-inflammatory

transcripts that was specifically blocked by RV in response to exogenous IFN-β, but not

to intracellular dsRNA (*TLR2, IL26, TNF, IRF8, TNFAIP3, IL29, IL1B)*. Among the

transcripts measured, only *SERPING* was efficiently induced by IFNAR1 stimulation

during RV infection, though with a lowered efficiency than in uninfected HT-29 cells. The

stimulus-specific RV inhibition of transcription was not due to RV replication sensitivity,

since it was unaltered under either the IFN-β or dsRNA stimulation conditions used (Fig.

4C). In addition, the induction of certain transcripts was specifically repressed by ectopic

IFNAR1 stimulation during RV infection (*HERC5, IFNG,* and *IL22,* statistically significant

p<0.01*; IRF8, IL29,* and *IL1B,* not statistically significant). These results demonstrate that

RV effectively inhibits IFN-β-receptor mediated transcription of selected ISGs and

inflammatory genes. In addition, they indicate that ectopic IFNAR1 stimulation during the

course of RV infection can elicit negative feedback inhibition of transcription (Fig. 4B, 4D).

Such feedback-based regulation(50) is likely to be particularly relevant to antiviral and

inflammatory functions in uninfected RV- bystander cells that, unlike a RV+ infected cell,

express IFNRs (8) (Fig. 4D).

Murine RV infection re-purposes IFNR-directed intestinal antiviral and

inflammatory signaling. The stimulation of IFNGR1 and IFNAR1 is reported to suppress

STAT1 via negative feedback signaling under certain conditions, fostering IFN-resistance

in the context of cancer and chronic viral infection(33, 50). We examined whether

intestinal IFNR-mediated transcription is altered by homologous RV *in vivo* to suppress

IFN antiviral and inflammatory signaling (Fig. 1A and 3A). Following infection of suckling

mice with murine EW RV for 12h, we parenterally administered purified IFN-γ (or PBS),

and collected small intestines after an additional 12h for transcriptional analysis (Fig. 5A).

In mock-infected control pups, IFN-γ stimulation upregulated several intestinal transcripts

(*STAT1, ISG15, IFI204, IL10RA, IL1RN,* statistically significant*; S100A8, BIRC5, REG3G,*

not significant; Figs. 5B-5D). In contrast, transcription of *CCL5* was significantly repressed

by ectopic IFNGR1 stimulation. Similar transcriptional responses are also upregulated by

EW RV infection, but with magnitudes significantly larger than obtained by IFN-γ

stimulation alone. A set of antiviral (*DHX58, DDX58, MX2*) and inflammatory (*TRAIL,*

NOS2, CXCL10) transcripts was significantly upregulated by RV, but not by IFN-γ

stimulation. Remarkably, ectopic stimulation of the IFNGR1 in EW RV-infected mice

either significantly suppressed (*STAT1, ISG15, TRAIL,* and *CXCL10*) or completely

eliminated (*DHX58, DDX58, MX2, CCL5,* and *NOS2*) these intestinal antiviral and

inflammatory responses (Fig 5B-C). In contrast, IFN-γ stimulated, EW RV-infected pups

upregulated anti-inflammatory and proliferative gene transcription (*S100A8, BIRC5,*

ARG2, REG3G, MKI67, and *IL1RN*) (Fig. 5D). These transcripts were not significantly

induced by either IFN-γ stimulation or by EW RV-infection alone, and synergistically

increased when IFNGR1 stimulation was carried out during EW RV infection. To ascertain

whether the effects of IFN stimulation in EW RV-infected pups could be explained by

altered RV replication levels, we measured intestinal RV RNA by qRT-PCR. The results

demonstrate that EW RV replication was not significantly altered following IFN-γ

stimulation (Fig. 5E). These findings suggest that during homologous RV infection,

intestinal IFNR function can be diverted towards suppressing antiviral and inflammatory

responses and promoting an anti-inflammatory program, revealing a potential

mechanism by which uninfected RV bystander cells resist the antiviral and inflammatory

effects of IFN secreted by IEC and enteric hematopoietic cells(7, 51).

Discussion

Interferons are pleiotropic innate immune effectors that constitute a primary line of

defense against viral pathogens. By signaling through their cognate receptors, different

IFN types potentiate the expression of numerous and multi-functional antiviral and

inflammatory genes(4, 5). In addition, IFNR stimulation can also induce apoptotic and

necroptotic forms of cell death(17-19, 46), possibly to limit viral replication and spread. In

its homologous host species, RV replication is only marginally restricted by the induction

of different IFNs and robust viral replication is not associated with substantial

inflammation(7, 8, 10, 11, 16). However, heterologous RV replication is highly restricted

by each of the three major IFN types(10). How homologous RVs manage to avoid the

antiviral and inflammatory actions of IFNs is still not fully understood.

We found that in suckling mice, mucosal innate immune transcriptional responses

to the virulent and highly infectious murine EW RV strain closely resembled those to a

replication-restricted heterologous simian strain (Fig.1A). The similarity in innate

responses - including IFN type transcription, IFN-γ production, and amplification of ISGs

or inflammatory gene transcripts over time (Fig. 1C) - did not correlate well with the very

different replication levels of the EW vs. RRV (approximately 10,000-fold difference, Fig.

1B). Different IFN type transcripts remained significantly elevated at 72 hpi in EW RV-

infected pups, indicative of a sustained innate immune response to ongoing RV

replication. However, we did not observe amplification of several ISGs (i.e. *ISG15, ISG20,*

IFI203, IFIH1, DDX58, IRF9, and *MX2* (Fig. 1A) during the later stages of EW infection.

These results indicated that although homologous RV replication and IFN induction occur

in a sustained manner during infection, this does not effectively lead to sustained

amplification of multiple antiviral and inflammatory genes.

In response to sustained IFN exposure, apoptosis can occur via proteolytic

activation of the initiator caspase 8(19, 20). IFN-stimulated apoptosis is a relatively slow

process, likely mediated by IFN-stimulated expression of intermediary apoptotic ligands

(e.g., FasL and TRAIL)(18-20), and typically has mild inflammatory consequences. In

addition to triggering apoptosis, proteolytic cleavage of pro-caspase 8 can suppress

necroptosis - a more inflammatory form of cell death(17, 52). Necroptosis is rapidly

activated when cells are deficient for caspase 8, or in the presence of caspase inhibitors,

following exposure to inflammatory ligands like TNF-α or FasL(21). Ligand stimulation of

the IFN-α/β and -γ receptors can also precipitate rapid cellular necroptosis, leading to

acute inflammation(17, 21). This process is mediated by currently unknown but critical

IFNR-dependent effectors (termed constitutively IFN-regulated effectors of necroptosis,

CIRENs)(46). In addition, IFNs can amplify the expression of inflammatory cytokine genes

encoding IFN-sensitive response promoter elements. Both IFN-mediated necroptosis and

inflammatory gene expression require an intact IFNR signaling pathway and are critical

mediators of acute intestinal inflammation in response to endotoxin(17, 21, 43, 46, 53).

Of note, before this set of experiments, the regulated cell death and inflammatory arms

of IFN receptor signaling had not been well studied in the context of RV infection.

Our results suggest that homologous RV infection initiates a two-pronged set of

events to subvert and delay IFN-mediated cell death functions and potentially allow more

time for RV replication within IECs. An estimated 1400 cells/villous/day are normally shed

to maintain intestinal homeostasis in an adult mouse(54). We found that a key step in

epithelial cell turnover, the cleavage of caspases 8 and 3, was significantly inhibited

during homologous RV infection in both virus-infected and uninfected bystander mature

villous IECs (Fig. 2, S3). This blockage of caspase activation may serve to prevent or

diminish IFN-accelerated apoptosis and thus enhance viral replication and spread to

uninfected IECs. The inhibition of caspase 3 activity could itself result from RV-inhibited

cleavage of the upstream initiator caspase 8 (Fig. 2A, 2C-D), which mediates caspase 3

activation by the extrinsic death pathway and is inhibited in both RV-infected and

bystander IECs (Fig. 2D). In addition, homologous murine RV also hampers homeostatic

cleavage of another initiator caspase (caspase 9) that is crucial for executing intrinsic cell

death pathways. The ability of RV to regulate multiple types of caspases *in vivo* is novel

and, to the best of our knowledge, represents an ingenious viral strategy to broadly

regulate and delay several effectors of intestinal cell death. The inhibition of caspase

cleavage also occurs during RV infection *in vitro* and is temporally regulated (Fig. 2F, G).

The mechanism by which cleavage of the different types of caspases is regulated

by RV is presently unknown. Nevertheless, analysis of a key caspase 8-regulated

signaling effector, RIP1K(28, 30, 53), in RV-infected HT-29 cells provides possible insight

into underlying mechanisms. Specifically, we found that *in vitro* different strains of RV

promoted the cleavage of full-length p75 RIP1K into shorter fragments (i.e. p51, p37, and

p25) (Fig. 2H). The cleavage of RIP1K into the p51 fragment is caspase 8-mediated and

generates a negative regulator that can prevent apoptotic and necroptotic cell death in

cells constantly exposed to inflammatory stimuli(31). In contrast to the regulation of RIP1K

by caspase 8, under certain conditions (e.g. ER stress) RIP1K is required for proper

caspase 8 cleavage and apoptosis(55). This suggests the existence of a regulatory

feedback of RIP1K on caspase 8 activity. One possible explanation of our findings could

be that one or more RIP1K cleavage products generated early during RV infection (Fig.

2H) regulate caspase 8 cleavage and cell death during ER stress conditions that occur

later in RV infection(56-59). Time-course analyses of caspase 8 and RIP1K cleavage

during RV infection (Fig 2G-H) indicated that both these processes are impaired

temporally, but do not mediate appreciable levels of cell death or apoptotic markers even

at relatively late times of infection (Fig. 2I). It will be interesting to determine whether these

RIP1K regulatory processes occur in RV bystander cells and to identify the viral and host

effectors involved.

Regardless of the exact underlying mechanisms, impaired caspase 8 cleavage in

both RV infected and bystander IECs (Fig. 2D) creates a potentially problematic situation

in the intestinal epithelium(21). IEC-specific caspase 8 deletion by itself is sufficient to

trigger spontaneous intestinal inflammatory lesions, resulting in terminal ileitis and

colitis(48, 60). As noted above, caspase 8 cleavage inhibition is also a key priming event

that triggers IFN receptor-mediated necroptosis, acute inflammation, and endotoxin-

induced tissue damage(17-19, 27, 46, 53, 60). Paradoxically, we found that murine EW

RV infection significantly protected suckling mice from subsequent lethal endotoxemia

(100% vs. 0% survival, at 24h post-endotoxin exposure)(8). Since endotoxin-mediated

damage is potentiated by IFN receptor-stimulated inflammatory gene expression(44) and

necroptotic cell death(27), it seemed plausible to postulate that homologous RV infection

locally represses IFN-mediated inflammatory functions in the small intestine. Our

histopathologic findings (Fig. 3C) confirm that, at the local small bowel level, EW RV-

infected murine pups are substantially protected from severe inflammatory damage due

to exogenous endotoxin exposure. An interesting question raised by these findings, which

we are currently examining, is whether similar protective effects of RV on inflammation

can occur at extra-intestinal sites. Homologous RV infection also blocked the IFN

receptor-stimulated inflammatory gene expression program, as indicated by the finding

that intestinal secretion of several IFN-stimulated inflammatory cytokines (RANTES(34-

36), IP-10(35, 37), Eotaxin(38), GRO-α(39, 40), MIP-1b(36, 41, 42)) were all significantly

lower in the murine RV-infected pups compared to a replication-restricted heterologous

simian RV, despite substantially higher replication levels of the murine strain (Fig. 3A).

Rotavirus efficiently degrades multiple IFNR types in infected cells(8), which is

predicted to directly limit both IFN-mediated inflammatory cell death and gene

amplification by depleting surface IFNR expression. The RV protein(s) that mediate IFNR

degradation have not been identified, although the RV-encoded non-structural protein

NSP1 would seem to be the most likely candidate due to its ability to inhibit STAT1

signaling and cause the degradation of both IRF3 and bTrCP(26, 32, 61). In this study we

examined whether IFNR function in bystander cells was also inhibited by RV. We

previously reported that a porcine RV strain (SB1-A) prevents STAT1-Y701

phosphorylation in uninfected RV- bystander HT-29 cells following *a relatively short (i.e.*

30 min) ectopic IFN exposure(26). Whether non-porcine RVs also prevent IFN-mediated

signaling in bystander cells was not explored further using altered IFN stimulation

conditions. Here, using a more prolonged IFN stimulation (i.e. 6 h) combined with FACS

analysis, we demonstrate that several non-porcine RV strains, shown previously to

mediate IFNR degradation in infected cells, also induce effective resistance to IFN-

directed STAT1-pY701 in bystander cells (Fig. 4A).

The ability of various RV strains to block IFN signaling in bystander cells is a

second prong of the viral strategy to subvert antiviral and inflammatory functions mediated

through the IFNRs. The effector functions of different IFNs are precisely titrated by a

sensitive receptor-centered negative-feedback modulating mechanism(62-64), ensuring

that excessive inflammatory damage to the host does not occur by uncontrolled feed-

forward IFNR signaling. Dysregulation of this equilibrium can lead to either IFN resistance

exemplified by cancer cells(65), or to acute inflammatory pathologies(66),such as sepsis.

Previously, we demonstrated that parenteral IFNγ administration results in efficient

STAT1-pY701 activation in small intestinal epithelial cells, a process that is significantly

suppressed during homologous EW RV infection in suckling mice(8). A similar

perturbation of IFNR-mediated STAT1 activation by murine EW RV occurred if purified

IFN-β or IFN-λ was used to stimulate their cognate receptors(8). Earlier studies also

identified a STAT1-independent pathway that regulates IFNγ-stimulated gene

expression(67-70). More recently, the production of inflammatory cytokines in response

to toll-like receptor (TLR) stimulation was significantly suppressed by ectopic IFNγ and

IFNβ stimulation of STAT1-null cells, by re-purposing IFNGR1 and IFNAR1 signaling

toward feedback inhibition(50). In this study, we examined whether such alternate IFNR

signaling could also be usurped by RV to broadly suppress innate antiviral and

inflammatory responses.

Several transcripts upregulated during the IFNR feed-forward amplification phase

can also be efficiently induced by intracellular dsRNA through signal transduction of

different pattern recognition receptors, including the RIG-I-like-receptors (RLRs), toll-like-

receptors (TLRs), and the dsRNA-dependent protein kinase (PKR)(3). We therefore

compared the ability of dsRNA- and IFNβ-stimulation to upregulate antiviral and

inflammatory gene expression during RV infection of HT-29 cells (Fig. 4B). As RV$^+$

infected HT-29 cells are efficiently depleted of different IFNRs, an assumption in our

experiment was that IFNβ-stimulated transcription occurs via the IFNRs expressed on

uninfected RV- bystanders, although we did not directly demonstrate this. The results

clearly indicated an efficient inhibition of feed-forward transcriptional responses to IFNR

stimulation by RV, when compared to intracellular dsRNA stimulation (Fig. 4B).

Interestingly, IFNβ stimulation during the course of RV infection repressed *HERC5*, *IFNG*,

IL22, IRF8, IL29, and *IL1B* transcription, suggesting that IFNAR1 stimulation (presumably

of uninfected RV- bystander cells) might also trigger an inhibitory feed-back transcription

mode (Fig. 4C).

In subsequent experiments, we quantified intestinal IFNR-stimulated transcription

during homologous EW RV (or mock) infection by ectopically stimulating the IFNGR1 in

suckling mice (Fig. 5A). These results demonstrated that, in contrast to a feed-forward

'priming' transcriptional mode in mock controls, IFNγ stimulation during EW RV infection

potentiates decreased ISG and inflammatory transcripts (*STAT1, ISG15, IFI204, DHX58,*

DDX58, MX2, CCL5, TRAIL, NOS2, and *CXCL10*) (Fig. 5B-C). In addition, we found that

ectopic IFNGR1 stimulation during EW RV infection (but not IFNγ stimulation or EW RV

alone) synergistically increases the transcription of anti-inflammatory transcripts (Fig.

5D). Interestingly, several of these upregulated genes (i.e. *S100A8, BIRC5, ARG2,*

REG3G, and *MKI67*) are targets of the STAT3 signaling pathway, whose activation is

reported to suppress STAT1-mediated inflammatory and antiviral programs due to IFNR

stimulation(33, 68-70). We are currently examining whether STAT3 activation occurs

during RV infection and mediates IFNR feedback signaling.

Collectively, our findings demonstrate that RV effectively reprograms IFNR-relayed

transcription towards suppressing antiviral and inflammatory amplification (Fig. 3, 4, 5), a

process whose efficiency is expected to be enhanced by IFNs generated in the local

intestinal milieu during enteric RV infection (Fig. 1). The ability of RV infection to

reprogram IFNR transcriptional function identifies an underlying mechanism by which

other IFNR functions, such as caspase cleavage-mediated cell death could also be

controlled by RV.

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FIGURE LEGENDS

Figure 1. Effects of replication-competent (homologous, murine) and replication-

restricted (heterologous, simian) RV strains on intestinal interferon-stimulated

gene (ISG) and inflammatory transcriptional responses. (A-B) Transcriptional

analysis of intestinal transcriptional responses to murine EW and simian RRV RV strains.

Three- to five-day old suckling mice (N=4 per group) were infected with murine EW or

simian RRV RV strains (or mock), and at 12 h or 72 h small intestines were collected for

transcriptional analysis of host (A) and RV (B) genes. (C) Intestinal secretion of IFN-γ was

measured by Luminex following RV infection at 24 hpi as in (A) (N=3 samples per group,

each sample pooled from 3 pups; ND, not detected).

Figure 2. Rotavirus suppresses the cleavage of multiple caspases and promotes

the cleavage of receptor-interacting protein 1 kinase (RIP1K). (A-B) Analysis of

caspase 8 and caspase 3 expression. Mouse pups were orally inoculated with EW (R) or mock-inoculated (M). At 3 and 24 hpi, bulk intestinal cells were harvested from whole small intestinal tissues and processed for SDS-PAGE. Gels were analyzed by western blot for precursor (FL) and cleaved (Cl.) isoforms of either caspase 8 (A) or caspase 3 (B), and other proteins. The normalized ratios of Cl. to FL caspases determined by densitometry are indicated below individual panels. (C) FACS analysis of caspase 8 and caspase 3 cleavage. Summary of flow cytometry analysis of CD45$^-$CD44$^-$ESA$^+$ IECs following intracellular staining for KI67, and either cleaved caspase 8 or cleaved caspase 3, from EW RV infected or control pups at 24 hpi. (D) The percentage of IECs (CD45$^-$CD44$^-$ESA$^+$) displaying cleaved caspase 8 or cleaved caspase 3 in mock and EW-infected mice at 24 hpi averaged from three experiments; error bars represent the standard errors of the means. (E) Intracellular staining for RV structural protein (VP6) and either cleaved caspase 8 (top panel) or cleaved caspase 3 (bottom panel) in CD45$^-$CD44$^-$ESA$^+$ IECs following EW RV or mock infection at 24 hpi. (F) Immunoblotting analysis of small intestinal tissue from mouse pups infected with EW, RRV, or mock, and harvested at the indicated times. (G) Analysis of caspase 9 cleavage in cultured NIH 3T3 fibroblasts infected with an IFN-stimulatory RV strain, UK. 3T3 cells were infected with bovine UK RV (or mock) for the times indicated and then analyzed by immunoblotting. Numbers below blots indicate normalized ratios of Cl. to FL caspase 9. (H) Human HT-29 IECs were infected with porcine SB1-A RV (or mock) for the times indicated and analyzed by immunoblotting. Numbers below blots indicate normalized ratios of Cl. to FL caspase 8. (I) HT-29 cells infected with the RV strains indicated were analyzed for RIP1K, RIP3K, and VP6 at either 4 or 12 hpi by immunoblotting. (J) HT-29 cells were infected with SB1-

A RV (or mock), or treated with TNF-alpha (10ng/ml) and cycloheximide (10μg/ml) for 4

h, and then analyzed for cell adherence or phosphatidyl serine marker expression. Data

shown is representative of two or more independent experiments.

Figure 3. Effects of replication-competent (homologous, murine) and replication-

restricted (heterologous, simian) RV strains on intestinal cytokine production. (A)

Secretion of intestinal cytokines during RV infection of suckling mice. Suckling mice were

infected as in Figure 1, and small intestinal homogenates were analyzed at 24 hpi for the

indicated cytokine protein levels (N=3 samples per group, each sample pooled from 3

pups and measured in duplicate). (B) Schematic of reported endotoxin-mediated

inflammation that is IFNR-stimulated and negatively regulated by cleaved caspase 8. (C)

Effect of homologous RV infection in suckling mice on endotoxin-mediated intestinal

damage. Suckling mice were mock- or EW RV-infected and 3 d later were parenterally

administered purified endotoxin (15μg/pup) for 6 h before harvesting small intestines for

histological examination by hematoxylin and eosin staining. Data shown is representative

of 2 pups.

Figure 4. Rotavirus potentiates resistance to IFN receptor-mediated STAT1

activation and transcriptional function. (A) Different RV strains trigger resistance to

IFN-mediated STAT1-Y701 phosphorylation in bystander cells. HT-29 cells were infected

with the RV strains indicated and 12 h later, stimulated with 500 IU/ml of purified IFNβ for

6 h before FACS analysis. Numbers in red indicate fold increases in STAT1-pY701

following IFN stimulation in RV VP6+ (infected) and VP6- (bystander) populations. Data

is representative of two independent experiments. (B-C) Effect of RV infection on dsRNA-

and IFN receptor-mediated transcriptional activity. Schematic at the top depicts the

experimental approach used. HT-29 cells were infected with RV (RRV strain, at an MOI

of ~0.5) or mock-infected and at 6 hpi, cells were either transfected with 2μg of purified

dsRNA or were stimulated with 500 IU/ml of purified IFNβ. At 12 hpi cells were analyzed

by qRT-PCR for expression of host (B) and RV (C) transcripts. Data represents two

independent experiments. (D) Schematic summarizing differential inhibition of dsRNA-

and IFN receptor-stimulated transcription by RV. Dashed red line indicates possible

negative feedback signaling upon ectopic IFNR stimulation during RV infection, solid red

lines indicate RV inhibition of IFNRs by receptor degradation (in RV-infected cells(8)) and

by blockage of IFNR-mediated STAT1 activation and transcription (in RV bystanders).

Figure 5. Rotavirus infection re-purposes IFN receptor-directed intestinal and

inflammatory signaling. (A) Schematic of the experimental approach used. (B-E)

Transcriptional analysis of different assigned categories of intestinal transcripts following

ectopic stimulation of the type II IFN receptor. Mice were infected with EW RV (or mock)

(N=3-5 pups per group) and at 12 hpi were administered purified murine IFNγ (1μg/pup,

or PBS, i.p.). Twelve hours later mice were sacrificed, and small intestinal RNA was

analyzed by qRT-PCR for different host (B-D) or RV NSP5 gene (E) transcripts.

MATERIALS AND METHODS

Viruses and reagents. The tissue culture-adapted RV strains (simian RRV, SA11-4F, SA11-5S, human WI-61, porcine SB1-A, bovine UK) were propagated in African green monkey kidney cells (MA104) as previously described (71). Virus titers were determined by plaque assays in MA104 cells. HT-29 cells were plated in 6-well or 24-well cluster plates and completely confluent monolayers were infected 2-3d later as described. The non-tissue culture-adapted murine EW RV strain was prepared as an intestinal homogenate from RV-infected BALB/C suckling mice. The diarrheal dose (DD_{50}) was determined as previously described (16). Human intestinal epithelial HT29 and embryonic kidney HEK293 cells were purchased from the American Type Culture Collection (ATCC) and maintained in Advanced D-MEM Ham's F12 (Cellgro) or D-MEM medium (D-MEM, Cellgro) containing 10% fetal calf serum (Invitrogen) supplemented with non-essential amino acids, glutamine, penicillin, and streptomycin, respectively. Purified human IFN-β (PBL), human IFN-λ (PBL), and human IFN-γ (Millipore) were used for stimulation of cells. Purified carrier-free human IFN-β (PBL), murine IFN-γ, human TNF-α (R&D System, Minneapolis, MN), LPS (Sigma) were used. Cycloheximide (Sigma) and high-molecular weight LyoVec poly(I:C) (Invivogen) were used at 10 µg/mL and 2.5 µg/mL, respectively.

Mouse infection and tissue harvest. All mice were housed in the VA Palo Alto Health Care System (VAPAHCS) Veterinary Medicine Unit and all experimentation followed VAPAHCS Institutional Animal Care and Use Committee approved protocols. 3-5 day old Sv129 pups were infected via oral gavage with 10^4 DD_{50} of murine EW RV, 10^7 PFU of the simian RRV, or mock infected with 1X M199 media (Gibco) as described (16). (16). At the indicated time-points, the small intestines were removed. Sections were stored at

-80 °C or fixed in 4% paraformaldehyde for processing and subsequent analyses. The

protocol for IEC isolation, as previously described (72), was slightly modified. Briefly, the

intestines were removed and opened longitudinally in RPMI media supplemented with

penicillin, streptomycin, and L-glutamine (Gibco). The intestines were washed, cut into

smaller pieces, and placed in 1X PBS containing 5 mM EDTA and 5 mM DTT (Sigma-

Aldrich). Tissues from 3-4 pups (per infection group) were pooled at this point. The tissues

were then placed on a shaker for 10 minutes at room temperature. Released cells were

subsequently filtered through a 100-µm mesh filter into fresh RPMI FBS media (RPMI

supplemented with 10% fetal bovine serum, penicillin, streptomycin, and L-glutamine) and

immediately placed on ice. The remaining tissues were then incubated with fresh RPMI

FBS media and rocked for an additional 10 minutes, followed by a final wash in RPMI

FBS media for an hour. Cells were filtered, washed, and resuspended in PBS for further

analyses.

Mouse stimulation and Luminex assays: To ectopically stimulate IFNGR1, suckling

mice infected with 10^4 DD$_{50}$ of murine EW RV (or mock-infected) were intraperitoneally

administered purified murine IFN-γ (1µg IFN-γ per pup) or IFN- λ2 at 12 hpi and sacrificed

12 h later for analysis of transcripts by qRT-PCR. For endotoxin stimulation, 3-5 day old

129sv mice were orally inoculated by gavage with 10^4 DD$_{50}$ of murine RV-EW. At 3 dpi,

mice were intraperitoneally injected with PBS or purified LPS (Sigma, 10 mg/kg body

weight) and sacrificed 6 h later for analysis of paraformaldehyde-fixed and paraffin

embedded tissue sections by hematoxylin and eosin staining. For measurement of

intestinal cytokines, mouse intestines were harvested and weighed pooled tissues were

homogenized in ice-cold PBS containing a cocktail of protease and phosphatase

inhibitors (Sigma) using a handheld micropestle (Thermo Fisher). Lysates were clarified

by two sequential rounds of centrifugation at 10,000X G for 10 min, and analyzed for

cytokine levels at the Human Immune Monitoring Center at Stanford University. Mouse

38-plex kits were purchased from eBiosciences/Affymetrix and used according to the

manufacturer's recommendations. Plates were read using a Luminex 200 instrument with

a lower bound of 50 beads per sample per cytokine. Custom assay Control beads by

Radix Biosolutions are added to all wells.

Western blot analysis. Preparation of cell lysates and immunoblotting in vitro was

performed as described. For mouse samples, EDTA/DTT-treated IECs were lysed in

radioimmunoprecipitation assay buffer (RIPA) supplemented with protease and

phosphatase inhibitors (Roche). Equal volumes of 2X Laemmli buffer (Sigma-Aldrich)

were added to each sample and analyzed by immunoblotting. Blots were probed with the

following primary antibodies: anti-caspase 3, anti-caspase 8, anti-caspase 9, anti-cleaved

caspase 3, anti-cleaved caspase 8, anti-cleaved caspase 7, anti-PARP, anti-GAPDH,

anti-survivin, anti-STAT-3, anti-p-STAT-3 (Y705) (all from Cell Signaling Technologies)

and an anti-RV capsid antibody (from Santa Cruz Biotechnologies). The blots were

subsequently incubated with either anti-rabbit or anti-mouse secondary antibodies

conjugated to horseradish peroxidase (HRP), incubated with ECL Detection reagent, and

exposed to film (GE Healthcare). All densitometry quantifications of protein blots were

analyzed using ImageJ software analysis and normalized to GAPDH protein bands.

Transcriptional analysis. For *in vivo* analysis of transcripts, mice were sacrificed and

sections of the small intestine were collected on ice in Trizol (Life Technologies). For *in*

vitro studies, cells were first washed in PBS and then lysed in Trizol. Total RNA was

extracted following the manufacturer's instructions and subjected to DNAse digestion

before use in qRT-PCR. Synthesis of cDNA and subsequent microfluidics PCR on the

Fluidigm platform was done as described earlier (73). Serial 10-fold dilutions of mouse or

human reference RNA (Agilent) were run in duplicate for each PCR run.

Flow cytometry analysis. For flow cytometry analysis of caspase expression and cell

viability *in vivo*, mouse intestines were harvested as described above and equivalent

numbers of EDTA/DTT-treated IECs were stained for cell viability using Invitrogen's

LIVE/DEAD Fixable Aqua Dead Cell Stain Kit following the manufacturer's protocol. Cells

were blocked with purified rat anti-mouse CD16/CD32 Fc Block (BD Biosciences) and

stained for the following murine surface markers: anti- CD45 (clone 30-F11) v450 (BD

Biosciences), anti- CD326 (Ep-CAM/ESA, clone G8.8) PE (Biolegend), and anti- CD44

(clone IM7) PerCP/Cy5.5 (Biolegend) as described earlier (7). Cells were then fixed and

permeabilized with BD Cytofix/Cytoperm Fixation/ Permeabilization buffer according to

the manufacture's protocol and subsequently used for intracellular protein staining. To

perform intracellular protein staining, cells were blocked with BD Fc Block and incubated

with unconjugated antibodies to cleaved caspase 8 (Cell Signaling Technologies) or

cleaved caspase 3 (Cell Signaling Technologies) along with conjugated antibodies to RV

VP6 (1E11-Texas Red) and Ki67 (Alexa Fluor 647, BD Biosciences). Subsequently, cells

were incubated with a secondary goat anti-rabbit-Alexa Fluor 488 antibody (Molecular

Probes) for detection of cleaved caspases. All data was acquired using a LSRII (BD

Biosciences) and analyzed with FlowJo software (Treestar Inc). Staining and flow

cytometry analysis of HT-29 cells was performed as described.

TUNEL and apoptosis assays. For TUNEL, following surface staining of intestinal

epithelial cells (as described above), cells were fixed and permeabilized for subsequent

determination of DNA fragmentation using Roche's In Situ Cell Death Detection Kit-

Fluorescein (following the manufacturer's protocol). Briefly, cells were fixed in 2% PFA

and permeabilized in 0.1% Triton X-100 in 0.1% sodium citrate buffer and incubated with

the terminal deoxynucleotidyl transferase (TdT), along with fluorescein-dUTP labeling

solution for 1 hour at 37 °C. As a positive control, permeabilized IECs were treated with

DNase I (Ambion) for 30 min at 37 °C (to induce DNA fragmentation) prior to the TUNEL

reaction. An additional negative control (labeling reactions carried out without TdT

enzyme) was also included. Incorporation of labeled dUTP was analyzed on a LSRII flow

cytometer. To assay apoptosis marker expression in HT-29 cells, cells were washed with

PBS, stained with Apopxin Green (1:100 dilution, Abcam) at room temperature for 10

minutes, and washed before examination by fluorescence microscopy.

Statistical analysis. Analysis of variance (ANOVA) and 2-tailed student's t-tests were

performed using GraphPad InSTAT software. Figure legends indicate the specific tests

used and P values <0.05 were considered significant. All error bars represent the

standard error of the mean.

REFERENCES

1. Kirkwood CD, Ma LF, Carey ME, & Steele AD (2017) The rotavirus vaccine development pipeline. *Vaccine*.

2. Broome RL, Vo PT, Ward RL, Clark HF, & Greenberg HB (1993) Murine rotavirus genes encoding outer capsid proteins VP4 and VP7 are not major determinants of host range restriction and virulence. *J Virol* 67(5):2448-2455.

3. Talemi SR & Hofer T (2018) Antiviral interferon response at single-cell resolution. *Immunol Rev* 285(1):72-80.

4. Cicchese JM*, et al.* (2018) Dynamic balance of pro- and anti-inflammatory signals controls disease and limits pathology. *Immunol Rev* 285(1):147-167.

5. Yan N & Chen ZJ (2012) Intrinsic antiviral immunity. *Nat Immunol* 13(3):214-222.

6. De Boissieu D, Lebon P, Badoual J, Bompard Y, & Dupont C (1993) Rotavirus induces alpha-interferon release in children with gastroenteritis. *J Pediatr Gastroenterol Nutr* 16(1):29-32.

7. Sen A*, et al.* (2012) Innate immune response to homologous rotavirus infection in the small intestinal villous epithelium at single-cell resolution. *Proc Natl Acad Sci U S A* 109(50):20667-20672.

8. Sen A, Sharma A, & Greenberg HB (2017) Rotavirus degrades multiple type interferon receptors to inhibit IFN signaling and protects against mortality from endotoxin in suckling mice. *J Virol*.

9. Broquet AH, Hirata Y, McAllister CS, & Kagnoff MF (2011) RIG-I/MDA5/MAVS are required to signal a protective IFN response in rotavirus-infected intestinal epithelium. *J Immunol* 186(3):1618-1626.

10. Lin JD*, et al.* (2016) Distinct Roles of Type I and Type III Interferons in Intestinal Immunity to Homologous and Heterologous Rotavirus Infections. *PLoS Pathog* 12(4):e1005600.

11. Feng N*, et al.* (2008) Role of interferon in homologous and heterologous rotavirus infection in the intestines and extraintestinal organs of suckling mice. *J Virol* 82(15):7578-7590.

12. Lundgren O & Svensson L (2001) Pathogenesis of rotavirus diarrhea. *Microbes Infect* 3(13):1145-1156.

13. Morris AP & Estes MK (2001) Microbes and microbial toxins: paradigms for microbial-mucosal interactions. VIII. Pathological consequences of rotavirus infection and its enterotoxin. *Am J Physiol Gastrointest Liver Physiol* 281(2):G303-310.

14. Garrett WS, Gordon JI, & Glimcher LH (2010) Homeostasis and inflammation in the intestine. *Cell* 140(6):859-870.

15. Feng N*, et al.* (2011) Roles of VP4 and NSP1 in determining the distinctive replication capacities of simian rotavirus RRV and bovine rotavirus UK in the mouse biliary tract. *J Virol* 85(6):2686-2694.

16. Feng N, Yasukawa LL, Sen A, & Greenberg HB (2013) Permissive replication of homologous murine rotavirus in the mouse intestine is primarily regulated by VP4 and NSP1. *J Virol* 87(15):8307-8316.

17. Thapa RJ*, et al.* (2013) Interferon-induced RIP1/RIP3-mediated necrosis requires PKR and is licensed by FADD and caspases. *Proc Natl Acad Sci U S A* 110(33):E3109-3118.

18. Apelbaum A, Yarden G, Warszawski S, Harari D, & Schreiber G (2013) Type I interferons
induce apoptosis by balancing cFLIP and caspase-8 independent of death ligands. *Mol Cell*
Biol 33(4):800-814.
19. Balachandran S*, et al.* (2000) Alpha/beta interferons potentiate virus-induced apoptosis
through activation of the FADD/Caspase-8 death signaling pathway. *J Virol* 74(3):1513-
1523.
20. Chawla-Sarkar M, Leaman DW, & Borden EC (2001) Preferential induction of apoptosis by
interferon (IFN)-beta compared with IFN-alpha2: correlation with TRAIL/Apo2L induction
in melanoma cell lines. *Clin Cancer Res* 7(6):1821-1831.
21. Feltham R, Vince JE, & Lawlor KE (2017) Caspase-8: not so silently deadly. *Clin Transl*
Immunology 6(1):e124.
22. Wang F*, et al.* (2013) Isolation and characterization of intestinal stem cells based on
surface marker combinations and colony-formation assay. *Gastroenterology* 145(2):383-
395 e381-321.
23. Wurstle ML, Laussmann MA, & Rehm M (2012) The central role of initiator caspase-9 in
apoptosis signal transduction and the regulation of its activation and activity on the
apoptosome. *Exp Cell Res* 318(11):1213-1220.
24. Sen A, Feng N, Ettayebi K, Hardy ME, & Greenberg HB (2009) IRF3 inhibition by rotavirus
NSP1 is host cell and virus strain dependent but independent of NSP1 proteasomal
degradation. *J Virol* 83(20):10322-10335.
25. Sen A, Pruijssers AJ, Dermody TS, Garcia-Sastre A, & Greenberg HB (2011) The early
interferon response to rotavirus is regulated by PKR and depends on MAVS/IPS-1, RIG-I,
MDA-5, and IRF3. *J Virol* 85(8):3717-3732.
26. Sen A, Rott L, Phan N, Mukherjee G, & Greenberg HB (2014) Rotavirus NSP1 protein
inhibits interferon-mediated STAT1 activation. *J Virol* 88(1):41-53.
27. McComb S*, et al.* (2014) Type-I interferon signaling through ISGF3 complex is required for
sustained Rip3 activation and necroptosis in macrophages. *Proc Natl Acad Sci U S A*
111(31):E3206-3213.
28. Festjens N, Vanden Berghe T, Cornelis S, & Vandenabeele P (2007) RIP1, a kinase on the
crossroads of a cell's decision to live or die. *Cell Death Differ* 14(3):400-410.
29. Matsuoka Y & Tsujimoto Y (2015) Role of RIP1 in physiological enterocyte turnover in
mouse small intestine via nonapoptotic death. *Genes Cells* 20(1):11-28.
30. McComb S*, et al.* (2014) Cathepsins limit macrophage necroptosis through cleavage of
Rip1 kinase. *J Immunol* 192(12):5671-5678.
31. Zhang L*, et al.* (2015) RIP1 Cleavage in the Kinase Domain Regulates TRAIL-Induced NF-
kappaB Activation and Lymphoma Survival. *Mol Cell Biol* 35(19):3324-3338.
32. Barro M & Patton JT (2005) Rotavirus nonstructural protein 1 subverts innate immune
response by inducing degradation of IFN regulatory factor 3. *Proc Natl Acad Sci U S A*
102(11):4114-4119.
33. Ivashkiv LB & Donlin LT (2014) Regulation of type I interferon responses. *Nat Rev Immunol*
14(1):36-49.
34. Kim MO, Suh HS, Brosnan CF, & Lee SC (2004) Regulation of RANTES/CCL5 expression in
human astrocytes by interleukin-1 and interferon-beta. *J Neurochem* 90(2):297-308.

35. Wen X, *et al.* (2010) Predominant interferon-gamma-mediated expression of CXCL9,
CXCL10, and CCL5 proteins in the brain during chronic infection with Toxoplasma gondii
in BALB/c mice resistant to development of toxoplasmic encephalitis. *J Interferon*
Cytokine Res 30(9):653-660.
36. Sung PS, *et al.* (2017) CXCL10 is produced in hepatitis A virus-infected cells in an IRF3-
dependent but IFN-independent manner. *Sci Rep* 7(1):6387.
37. Majumder S, *et al.* (1998) p48/STAT-1alpha-containing complexes play a predominant
role in induction of IFN-gamma-inducible protein, 10 kDa (IP-10) by IFN-gamma alone or
in synergy with TNF-alpha. *J Immunol* 161(9):4736-4744.
38. Vargas A, *et al.* (2010) Association between plasma levels of eotaxin (CCL-11) and
treatment response to interferon-alpha and ribavirin in HIV/HCV co-infected patients. *J*
Antimicrob Chemother 65(2):303-306.
39. Gasperini S, Calzetti F, Russo MP, De Gironcoli M, & Cassatella MA (1995) Regulation of
GRO alpha production in human granulocytes. *J Inflamm* 45(3):143-151.
40. Meyer-Hoffert U, *et al.* (2003) Th2- and to a lesser extent Th1-type cytokines upregulate
the production of both CXC (IL-8 and gro-alpha) and CC (RANTES, eotaxin, eotaxin-2, MCP-
3 and MCP-4) chemokines in human airway epithelial cells. *Int Arch Allergy Immunol*
131(4):264-271.
41. Chandrasekar B, Deobagkar-Lele M, Victor ES, & Nandi D (2013) Regulation of
chemokines, CCL3 and CCL4, by interferon gamma and nitric oxide synthase 2 in mouse
macrophages and during Salmonella enterica serovar typhimurium infection. *J Infect Dis*
207(10):1556-1568.
42. Chiu BC, *et al.* (2003) Cytokine-chemokine networks in experimental mycobacterial and
schistosomal pulmonary granuloma formation. *Am J Respir Cell Mol Biol* 29(1):106-116.
43. Siegfried A, *et al.* (2013) IFIT2 is an effector protein of type I IFN-mediated amplification
of lipopolysaccharide (LPS)-induced TNF-alpha secretion and LPS-induced endotoxin
shock. *J Immunol* 191(7):3913-3921.
44. Karaghiosoff M, *et al.* (2003) Central role for type I interferons and Tyk2 in
lipopolysaccharide-induced endotoxin shock. *Nat Immunol* 4(5):471-477.
45. Mahieu T, *et al.* (2006) The wild-derived inbred mouse strain SPRET/Ei is resistant to LPS
and defective in IFN-beta production. *Proc Natl Acad Sci U S A* 103(7):2292-2297.
46. Sarhan J, *et al.* (2019) Constitutive interferon signaling maintains critical threshold of
MLKL expression to license necroptosis. *Cell Death Differ* 26(2):332-347.
47. Guo S, *et al.* (2015) Lipopolysaccharide Regulation of Intestinal Tight Junction
Permeability Is Mediated by TLR4 Signal Transduction Pathway Activation of FAK and
MyD88. *J Immunol* 195(10):4999-5010.
48. Gunther C, *et al.* (2011) Caspase-8 regulates TNF-alpha-induced epithelial necroptosis and
terminal ileitis. *Nature* 477(7364):335-339.
49. Dalet A, Gatti E, & Pierre P (2015) Integration of PKR-dependent translation inhibition
with innate immunity is required for a coordinated anti-viral response. *FEBS Lett*
589(14):1539-1545.
50. Kim HS, *et al.* (2015) STAT1 deficiency redirects IFN signalling toward suppression of TLR
response through a feedback activation of STAT3. *Sci Rep* 5:13414.

51. Pott J*, et al.* (2011) IFN-lambda determines the intestinal epithelial antiviral host defense.
Proc Natl Acad Sci U S A 108(19):7944-7949.

52. Wang H*, et al.* (2014) Mixed lineage kinase domain-like protein MLKL causes necrotic
membrane disruption upon phosphorylation by RIP3. *Mol Cell* 54(1):133-146.

53. Duprez L*, et al.* (2011) RIP kinase-dependent necrosis drives lethal systemic inflammatory
response syndrome. *Immunity* 35(6):908-918.

54. Potten CS & Loeffler M (1990) Stem cells: attributes, cycles, spirals, pitfalls and
uncertainties. Lessons for and from the crypt. *Development* 110(4):1001-1020.

55. Estornes Y*, et al.* (2014) RIPK1 promotes death receptor-independent caspase-8-
mediated apoptosis under unresolved ER stress conditions. *Cell Death Dis* 5:e1555.

56. Guerrero CA & Acosta O (2016) Inflammatory and oxidative stress in rotavirus infection.
World J Virol 5(2):38-62.

57. Crawford SE & Estes MK (2013) Viroporin-mediated calcium-activated autophagy.
Autophagy 9(5):797-798.

58. Trujillo-Alonso V, Maruri-Avidal L, Arias CF, & Lopez S (2011) Rotavirus infection induces
the unfolded protein response of the cell and controls it through the nonstructural
protein NSP3. *J Virol* 85(23):12594-12604.

59. Zambrano JL*, et al.* (2011) Rotavirus infection activates the UPR but modulates its activity.
Virol J 8:359.

60. Welz PS*, et al.* (2011) FADD prevents RIP3-mediated epithelial cell necrosis and chronic
intestinal inflammation. *Nature* 477(7364):330-334.

61. Arnold MM & Patton JT (2011) Diversity of interferon antagonist activities mediated by
NSP1 proteins of different rotavirus strains. *J Virol* 85(5):1970-1979.

62. Rateitschak K*, et al.* (2010) Mathematical modelling of interferon-gamma signalling in
pancreatic stellate cells reflects and predicts the dynamics of STAT1 pathway activity. *Cell
Signal* 22(1):97-105.

63. Swameye I, Muller TG, Timmer J, Sandra O, & Klingmuller U (2003) Identification of
nucleocytoplasmic cycling as a remote sensor in cellular signaling by databased modeling.
Proc Natl Acad Sci U S A 100(3):1028-1033.

64. Wiesauer I, Gaumannmuller C, Steinparzer I, Strobl B, & Kovarik P (2015) Promoter
occupancy of STAT1 in interferon responses is regulated by processive transcription. *Mol
Cell Biol* 35(4):716-727.

65. Benci JL*, et al.* (2016) Tumor Interferon Signaling Regulates a Multigenic Resistance
Program to Immune Checkpoint Blockade. *Cell* 167(6):1540-1554 e1512.

66. Bhattacharya S*, et al.* (2014) Triggering ubiquitination of IFNAR1 protects tissues from
inflammatory injury. *EMBO Mol Med* 6(3):384-397.

67. Ramana CV*, et al.* (2001) Stat1-independent regulation of gene expression in response to
IFN-gamma. *Proc Natl Acad Sci U S A* 98(12):6674-6679.

68. Lu L*, et al.* (2018) Gene regulation and suppression of type I interferon signaling by STAT3
in diffuse large B cell lymphoma. *Proc Natl Acad Sci U S A* 115(3):E498-E505.

69. Qing Y & Stark GR (2004) Alternative activation of STAT1 and STAT3 in response to
interferon-gamma. *J Biol Chem* 279(40):41679-41685.

70. Wang WB, Levy DE, & Lee CK (2011) STAT3 negatively regulates type I IFN-mediated
antiviral response. *J Immunol* 187(5):2578-2585.

71.   Deal EM, Lahl K, Narvaez CF, Butcher EC, & Greenberg HB (2013) Plasmacytoid dendritic
905   cells promote rotavirus-induced human and murine B cell responses. *J Clin Invest*
906   123(6):2464-2474.
907   72.   Guerrero CA, Santana AY, & Acosta O (2010) Mouse intestinal villi as a model
908   system for studies of rotavirus infection. *J Virol Methods* 168(1-2):22-30.
909   73.   Sen A*, et al.* (2012) Innate immune response to homologous rotavirus infection in the
910   small intestinal villous epithelium at single-cell resolution. *Proc Natl Acad Sci U S A*
911   109(50):20667-20672.
912

Figure 1.

Figure 2.

Figure 3.

3A

3B

3C

H & E stain

Figure 4.

Figure 5.