

The Intestinal Microbiome Primes Host Innate Immunity Against Enteric Virus Systemic Infection Through Type I Interferon

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1 **The intestinal microbiome primes host innate immunity against enteric virus**
2 **systemic infection through type I interferon**

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12

13 **Abstract**

14 Background: Intestinal microbiomes are of vital importance in antagonizing systemic viral
15 infection. However, very little literature has shown whether commensal bacteria play a
16 crucial role in protecting enteric virus systemic infection from the aspect of modulating host
17 innate immunity. Also, only a few specific commensal bacteria species have been revealed
18 to be capable in regulating antiviral innate immune responses mediated by type I interferon
19 (IFN). The underlying mechanisms have not yet been elucidated.

20 Results: We utilized an enteric virus, encephalomyocarditis virus (EMCV) to inoculate PBS-
21 treated or antibiotic cocktail-administrated mice (Abx) orally or intraperitoneally to examine
22 the impact of microbiota depletion on virulence and viral replication in vivo. Microbiota

23 depletion exacerbated the mortality, neuropathogenesis, viremia and viral burden in brain
24 following EMCV infection. Furthermore, Abx-treated mice exhibited severely diminished
25 macrophage activation and impaired type I IFN production and ISG expression in PBMC,
26 spleen or brain. With the help of fecal bacterial 16S rRNA sequencing of PBS and Abx
27 mice, we identified a single commensal bacterium *Blautia coccooides* (*B. coccooides*) that
28 can restore macrophage- and IFNAR-dependent type I IFN responses to restrict systemic
29 enteric virus infection.

30 Conclusion: Our present study demonstrates that intestinal microbiome is fundamental for
31 protecting from enteric virus systemic infection through activating macrophages and type I
32 IFN responses. Reconstitution with *B. coccooides* can inhibit enteric virus infection and
33 mitigate its neuropathogenesis by activating IFN-I and ISG responses in macrophages via
34 IFNAR- and STAT1-mediated signaling pathway.

35 Keywords: microbiota, *B. coccooides*, macrophage, type I interferon, enteric virus

36

37 **Introduction**

38 Mammalian intestines are colonized by trillions of microorganisms composed of bacteria,
39 viruses, archaea and fungi, collectively referred to as intestinal microbiota [1, 2]. The widely
40 diverse intestinal microbial communities have established an immensely complicated
41 ecosystem that is of vital importance in maintaining host homeostasis. Cumulative
42 evidence strongly supports the view that the dynamic crosstalk between the host and its
43 indigenous commensal bacteria is fundamental for the development, induction, education
44 and tuning of the host immune system [3]. As such, perturbation of the microbiota

45 composition, (termed as 'dysbiosis') is linked to a myriad of metabolic and inflammatory
46 diseases both at the intestinal mucosal sites and outside the gastrointestinal tract [2, 4, 5].
47 Over the past decade, rapid and extensive advances have shed light on the nature and
48 utmost importance of intestinal microbes in regulating virus replication, transmission and
49 pathogenesis. Multiple studies utilizing enteric viruses including poliovirus, retrovirus and
50 noroviruses have demonstrated that the intestinal bacteria can promote viral infection [6-
51 8]. The stimulating mechanisms include direct facilitation of viral binding to target cells,
52 virion stabilization and indirect regulatory pathways of suppressing the mucosal immune
53 responses [7, 9, 10]. However, as every coin has two sides, the signals from the indigenous
54 microbiota are shown to be essential to protect the *Drosophila* from enteric virus oral
55 infection by priming the antiviral innate immunity [11].

56 Other than exhibiting regulatory effect at mucosal sites during enteric viral infection, the
57 gut microbiota has been shown to be essential in limiting viral systemic infection by
58 calibrating innate immune responses mediated by mononuclear phagocytes [12-14]. A role
59 for the commensal bacteria to regulate the systemic type I interferon (IFN-I) has been
60 described in these studies, in which impaired production of IFN-I and interferon-stimulated
61 genes (ISGs) were observed in antibiotic (Abx)-treated or germ-free (GF) mice infected
62 with influenza A virus, murine cytomegalovirus, or Sendai virus, showing increased
63 mortality and susceptibility [13, 15, 16]. Recently, two compelling back-to-back research
64 articles reported that plasmacytoid dendritic cells (pDCs) are the main cellular sources of
65 Microbiota-induced IFN-I. Schaupp and colleagues demonstrated that microbiota-driven
66 IFN-I expression by pDCs primes conventional dendritic cells (cDCs) to initiate immune

67 responses following pathogen encounter [17]. Meanwhile, Diamond's group has pointed
68 out that IFN-I produced by microbiota-enabled pDCs prevents alphavirus, Chikungunya
69 virus (CHIKV) in particular, from infection and dissemination in host blood monocytes [18].
70 The latter study has demonstrated that a single commensal bacterium, *Clostridium*
71 *scindens* (*C. scindens*) can modulate prompt IFN-I responses through Toll-like receptor 7
72 (TLR7) and MyD88 signaling in pDC via primary to secondary bile acid (BA) transformation
73 [18]. Although these studies provided great insights into the systemic effect of microbiota
74 to limit viral infection, identification of unknown specific microbiome species that influence
75 type I IFN antiviral responses is still lacking and the molecular links between the gut
76 microbiota and type I IFN-mediated innate immunity are only beginning to be elucidated.
77 Here, we described the impact of the intestinal microbiome on host IFN-I associated
78 antiviral innate immunity using EMCV, a well-studied picornavirus that targets the central
79 nervous system and is transmitted via fecal-oral route. Depletion of the gut microbiota with
80 an Abx cocktail via oral gavage exacerbated the mortality and neurological symptoms to
81 EMCV infection in mice. EMCV infection of Abx-treated mice had increased viral loads in
82 blood and brain tissues. Coincidental with these phenotypes, the Abx-treated mice
83 exhibited diminished innate immune responses reflected by impacted NK cell activity and
84 macrophage activation post EMCV infection, and this was correlated with diminished
85 expression of ISGs in the periphery and brain tissues. We performed the bacterial 16S
86 rRNA sequencing of the fecal samples collected from the Abx cocktails-treated or single
87 Abx-treated mice to identify the possible commensal bacteria that may play important role
88 in priming IFN-I expression. Monocolonization of Abx-treated mice with *B. coecoides*

89 enabled them to regain the capability of restricting EMCV systemic infection by activating
90 NK cells and macrophages through type I interferon expression. Thus, in the present study,
91 we have identified a commensal bacterium that as far as we know, has not been previously
92 documented to be able to regulate type I IFN-mediated innate immunity in the context of
93 enteric virus systemic infection.

94

95 **Results**

96 **Microbiota depletion alters the neuropathogenesis and mortality to EMCV infection**

97 Groups of wild-type C57BL/6J mice (referred to as WT B6) were oral gavaged with a broad-
98 spectrum Abx cocktail, composed of Vancomycin (Van), Neomycin (Neo), ampicillin (Amp)
99 and Metronidazole (Metro) consecutively for 5 days and infected orally with different doses
100 of EMCV while the same Abx were kept in the water until the end of the experiments post
101 inoculation [6, 19, 20]. The clinical symptoms and mortality rate of different groups was
102 observed and documented for 14 days post-infection (dpi). As shown in Figure 1A, 60% of
103 Abx-treated mice inoculated with high dose of 2×10^7 TCID₅₀ units succumbed to infection
104 by 9 dpi, only about 10% of Abx-treated mice succumbed at the dose of 2×10^5 TCID₅₀ units
105 and no mortality occurred at the low dose of 2×10^3 TCID₅₀ units. However, all PBS-treated
106 control survived without clinical symptoms during the entire observation period. Alongside
107 the lethality phenotype, the Abx-treated mice infected with 2×10^7 TCID₅₀ units of EMCV
108 developed various clinical symptoms including hunch and trembling, hind limb paralysis,
109 dyspnea and death at 5 dpi, while the PBS-treated mice exhibited no obvious clinical signs
110 (Fig. 1A). We then harvested the blood and brain tissues of the infected mice at 3 and 5

111 dpi for virus loads detection using real-time RT-PCR. It was shown that EMCV per oral
112 inoculation resulted in significantly higher viral burden in the blood and brain tissues of
113 Abx-treated mice than the PBS-treated mice, indicating that viral replication in the
114 circulatory system and the target tissues was controlled by the presence of microbiota
115 during EMCV acute infection via oral delivery (Fig. 1B). Consistent with the clinical
116 symptoms and viral burden, Immunofluorescent assay revealed that there were more viral
117 signals (minor capsid protein VP2, Magenta) in the brain sections of Abx-treated mice than
118 the PBS-treated mice at 5 dpi (Fig. 1C, right panel). It seems that alteration of microbiota
119 composition does not alter the cellular tropism of EMCV in the brain for the majority of viral
120 signals were still co-stained with the marker of astrocytes (GFAP, red), but not with the
121 marker of microglial (IBA-1, green) in either PBS- or Abx-infected mice. Collectively, these
122 data indicate that the intestinal microbes are essential for limiting EMCV replication in the
123 target cells in the brain, and are required to protect the neurological diseases and cerebral
124 tissue lesions from EMCV oral infection.

125 To further confirm that the intestinal microbiome had an extraintestinal effect on limiting
126 EMCV systemic infection, we next infected both PBS- and Abx-treated mice systemically
127 via intraperitoneal injection to analyze the viral neurovirulence through this infectious route.
128 Indeed, Abx-treated mice were more susceptible to intraperitoneal infection with 200
129 TCID₅₀ units of EMCV than PBS-treated animals and reintroduction of fecal bacteria into
130 Abx-treated animals alleviated EMCV mortality and disease (Fig. 1D). As the onset of viral
131 dissemination and in vivo replication is faster in the context of intraperitoneal inoculation
132 than per oral infection, we measured the viral loads in the blood and brain tissues earlier

133 at 1 and 3 dpi and discovered that the Abx-treated mice supported more significant EMCV
134 replication than PBS-treated mice, whereas viral titers of Abx mice with fecal microbiota
135 transplantation (FMT) were similar to those of PBS-treated mice (Fig. 1E). It was
136 demonstrated that EMCV viremia and replication in target tissues was greatly increased in
137 the absence of microbiota and FMT could restore the capability of systemic viral clearance.
138 Taken together, these data indicate that commensal bacteria are critical in promoting host
139 systemic immunity against viral infection beyond the intestinal mucosal barrier.

140 **Microbiota deficiency severely diminishes innate macrophage antiviral immune**
141 **responses to EMCV systemic infection**

142 The impaired viral clearance in the blood and brain of Abx-treated animals after EMCV
143 infection provoke the hypothesis that the early systemic innate immune response was
144 crippled by microbiota depletion. Thus, we first evaluated the recruitment and activation of
145 early responding innate immune cells post EMCV infection by infecting PBS or Abx mice
146 intraperitoneally with 200 TCID₅₀ units of EMCV and extracting the peritoneal blood
147 mononuclear cells (PBMCs) and splenocytes at 3 dpi. The cell extracts were stained with
148 different antibodies specific for indicated cell types and quantified by flow cytometry.
149 Compared with their perspective mock control mice, infected PBS mice exhibited a
150 significantly increased percentage of inflammatory monocytes, NK cells and macrophages,
151 whereas there was a comparable frequency of pDC and cDC. In contrast with these data,
152 it seems that EMCV infection did not activate any of these cell subsets whatsoever
153 regarding cellular frequencies in infected- versus mock Abx mice (Fig. 2A and Fig. S1).
154 Also, analysis of the adaptive immune cells including T cells and B cells suggests that

155 commensal bacteria depletion did not alter the frequencies of T/B lymphocytes in the
156 context of EMCV infection (Fig. 2B).

157 Previous studies have shown that NK cell priming is dependent on cytokines expressed by
158 IFN-I-stimulated mononuclear phagocytes including DCs and macrophages [13, 21].

159 Macrophages, but not DCs had decreased expression of surface molecules that are critical
160 during the early responses to lymphocytic choriomeningitis virus (LCMV) or influenza virus
161 infection under the condition of Abx oral treatment [15]. Consistent with these findings, we

162 discovered that surface molecules reflecting macrophage activation, like MHC-I and CD80
163 were greatly augmented in the PBMC macrophages from infected PBS mice, but not in
164 those cells from infected Abx mice at 3 dpi compared to mock animals (Fig. 2C and Fig.

165 S2). To further determine the requirement for macrophages in the systemic antiviral innate
166 immunity and verify the necessity of intestinal microbiome in the macrophage response to
167 viral infection, we treated PBS or Abx mice systemically with clodronate liposomes to
168 deplete macrophages and analyze survival kinetics post EMCV intraperitoneal inoculation.

169 Depletion of macrophages was at least 90% effective as assessed by flow cytometry (Fig.
170 S3). For the mice that received control reagent, Abx treatment still resulted in significantly
171 higher death rate than PBS controls, however for the mice injected with clodronate

172 liposomes, Abx treatment did not make the animals more susceptible to EMCV infection
173 (Fig. 2D). In support of this phenotype, at 1 dpi, greater titers of EMCV were detected in
174 the blood of PBS mice than Abx mice both treated with control reagent, and no difference

175 was observed between PBS or Abx mice treated with clodronate liposomes (Fig. 2E).

176 Colonized mice treated with clodronate liposomes had almost 100-fold higher blood viral

177 loads than control reagent-treated colonized mice, indicating that macrophages are
178 absolutely important for the early control of EMCV systemic infection. Additionally, blood
179 viral titer of commensal bacteria-depleted macrophage knockout mice was also
180 significantly higher than Abx mice without macrophage depletion, suggesting that
181 commensal bacteria are not only required for activating macrophages, but also important
182 for mediating other innate antiviral immune factors (Fig. 2E).

183 Together, these data suggest that gut microbiota can regulate the activation of
184 macrophages, while disrupting the bacterial composition by antibiotic administration results
185 in a significant impact on the macrophage-dependent protective innate immunity to restrain
186 viral systemic infection.

187 **Commensal bacteria depletion greatly impairs the systemic and cerebral IFN-I**
188 **response following EMCV infection**

189 Because IFN-I is the key factor that contributes to the stimulation of macrophages, which
190 subsequently prime NK cells for antigen encounter via cytokine expression, therefore, we
191 hypothesized that the microbiota could regulate the IFN-I response in our model. To test
192 whether the commensal bacteria altered the profile of IFN-I production at different time-
193 points post infection, we inoculated groups of Abx-treated WT mice and PBS control with
194 200 TCID₅₀ units of EMCV intraperitoneally, collected blood and spleen at indicated
195 timepoints post-infection and probed for IFN- β and ISGs by real-time PCR. As early as 12
196 hours post-infection (hpi), there was significantly upregulated expression of IFN- β and
197 ISGs including *Isg15*, *Isg56*, *Oas1a* and *Mx1* in the PBMC extract (Fig. 3A) and spleen
198 (Fig. 3B) of PBS-treated mice compared to Abx-treated mice, suggestive of an impaired

199 innate immune response mediated by IFN-I. Consistent with the affected systemic IFN-I
200 response post-infection, there was reduced expression of IFN- β and associated antiviral
201 defense genes like *Isg15*, *Irf7*, *Irf9* and *Stat1/2* in the brain of Abx mice versus PBS mice
202 at 1 and 3 dpi, indicating that the antiviral innate immunity is impeded not only systemically,
203 but also locally at the target site of dissemination (Fig. 3C). In contrast, no differences in
204 expression of proinflammatory cytokine TNF- α , type II interferon, immune regulatory
205 cytokine IL-17 and IL-10 were observed in the brain tissues of PBS and Abx mice at 3 dpi,
206 implying that the impaired antiviral immunity in microbiome deficient Abx mice was not
207 associated with the production of these cytokines (Fig. 3D).

208 To further confirm that commensal bacteria-driven type I interferon response is critical in
209 antiviral innate immunity, we examined the requirement for the IFN-I induction and
210 amplification pathway in EMCV early control with or without antibiotic suppression. *Irf3*^{-/-} or
211 *Ifnar*^{-/-} mice that lacking key mediators either in type I interferon upstream induction
212 pathway or amplification loop respectively, were utilized to clarify the effects of microbiota
213 on different phases of IFN-I production upon EMCV systemic infection. Despite a narrowing
214 phenotype comparing PBS- and Abx-treated wild-type mice (P=0.0359), the lethality rate
215 discrepancy in *Irf3*^{-/-} mice was still above 30% (P=0.106). On the contrary, all *Ifnar*^{-/-} mice
216 were succumbed to EMCV inoculation by 5 dpi, showing identical susceptibility to EMCV
217 intraperitoneal infection regardless of PBS or antibiotics treatment (Fig. 3E). In consistent
218 with the phenotype on survival kinetics, the dissimilarities between PBS and Abx wild-type
219 mice in viremia (20-fold) and brain viral replication (10-fold) at 1 dpi narrowed to 6.3-fold
220 and 3-fold in *Irf3*^{-/-} mice, whereas the differences were completely lost in *Ifnar*^{-/-} mice (Fig.

221 3G). Collectively, these findings indicate that *Irf3*-mediated IFN-I induction pathway was
222 partially affected by Abx treatment, while intestinal microbiome-correlated IFN-I
223 amplification pathway is absolutely essential for preventing viral systemic infection.

224 ***B. coccoides* monocolonization alleviates EMCV pathogenesis and restricts viral** 225 **infection**

226 We next investigate whether specific commensal bacteria taxa are responsible for the
227 protein against EMCV systemic infection. First, we tested this idea by looking at which
228 antibiotic in the cocktail was required for the effect on IFN-I modulation and whether the
229 phenotype could be narrowed down to a single antibiotic treatment, or to a certain kind of
230 alteration in the intestinal microbiome per se. Groups of WT B6 mice were treated with
231 combination of four antibiotics or single antibiotic of Van, Neo, Amp or Metro, inoculated
232 intraperitoneally with EMCV and the brain tissues were dissected at 3 dpi for EMCV viral
233 burden measurement and IFN- β expression detection. Generally, single Abx treatment
234 significantly increased viral replication in the brain compared to PBS-treated controls
235 except Metro or Neo. Van treatment conferred partial suppressive effect to EMCV early
236 control as brain viral titers of these mice were not as high as Abx cocktail-treated animals.
237 Amp was the most effective single antibiotic for the viral replication was comparable to Abx
238 combination (Fig. 4A). Consistent with these data, Metro-treated mice exhibited very
239 similar level of IFN- β expression to the Abx cocktail-treated controls, suggesting that Metro
240 was not required for IFN- β suppressive effect. Amp, Van or Neo treatment all dampened
241 IFN- β expression significantly compared to PBS mice. Among them, Amp treatment
242 conferred an optimal suppression on IFN- β expression level, Van treatment showed an

243 intermediate effect, while Neo treatment had least impact on IFN- β expression (Fig. 4B).
244 As expected, successful inhibition of IFN- β expression in brain tissues in general
245 correlated with a substantial reduction in detectable 16S rRNA readouts upon antibiotics
246 treatment (Fig. 4C). In comparison, Van, Neo or Amp treatment had an effect is similar to
247 that of the antibiotic cocktail whereas Metro had the least effective bacteria depletion (Fig.
248 4C). Based on these data, one would hypothesize that certain bacteria communities in the
249 PBS control and resistant to Metro treatment, but significantly low in the Neo, Van, Amp or
250 Abx mice, mediated the protective effect to EMCV systemic infection as an IFN-I inducer.
251 Thus, we then carried out 16S rRNA gene sequencing and analysis on fecal samples
252 collected from PBS-, Abx cocktail- and single antibiotic-treated mice at 0 and 3 dpi (Fig.
253 4D). With the hypothesis mentioned above, we chose three bacterial taxa of *Blautia*,
254 *Akkermansia* and *Lactobacillus* because they had similar abundance between PBS- and
255 Metro- treated mice at 0 and 3 dpi in general (Fig. 4E). To explore whether they are the
256 specific bacteria that mediated the protective effect of microbiota against viral systemic
257 infection, we gavaged Abx mice with bacterial strain of *B. coccoides* (herein referred to as
258 B.C), *Akkermansia muciniphila* (*A. muciniphila*, herein referred to as AKK), *Lactobacillus*
259 *reuteri* (*L. reuteri*, herein referred to as L.R), an unrelated gram-positive human symbiont,
260 *Clostridium butyricum* (*C. butyricum*, herein referred to as C.B), or mouse fecal contents
261 (FMT) prior to EMCV intraperitoneal inoculation. Strikingly, colonization of Abx mice with a
262 single bacterial strain B.C fully rescued EMCV mortality to the extent of FMT mice and that
263 of PBS-infected controls (Fig. 4F), whereas gavage with C.B, L.R or AKK did not alter
264 mortality, even though these bacteria strains exhibited efficient colonization (Fig. S4).

265 Similarly, we observed significantly lower viral burden in the brain tissues of B.C colonized
266 Abx mice in comparison to that of Abx, FMT or C.B-colonized Abx mice at 3 dpi (Figure
267 4G). Taken together, these data suggest that specific commensal bacteria *B. coccoides*
268 protects host from EMCV systemic infection by restricting viral replication.

269 ***B. coccoides* monocolonization promotes macrophage activation and IFN-I**
270 **responses to systemic EMCV infection**

271 To further elucidate whether *B. coccoides* restricted EMCV infection by promoting
272 macrophage activation, we then gavaged Abx mice with B.C, C.B or fecal contents to
273 evaluate expression of activation surface markers of MHC-I and CD80 in macrophages of
274 PBMC and spleen. Indeed, B.C colonization in Abx mice remarkably increased the
275 frequency of MHC-I macrophages by 45% in PBMC (7.2% to 52%) and by 13% in spleen
276 (from 13.5% to 26%) macrophages following EMCV infection at 3 dpi, comparable to FMT
277 mice and PBS controls (Fig. 5A). In a similar way, CD80-upregulated macrophages were
278 elevated from 10% to 26.4% in PBMC, and from 7.36% to 53.7% in spleen macrophages
279 post B.C colonization in the context of EMCV infection (Fig. 5B). However, unlike B.C
280 colonized-Abx mice exhibited completely recovered MHC-I upregulation in both PBMC and
281 spleen macrophages to FMT treatment and PBS controls, the CD80 upregulation was only
282 partially rescued by B.C monocolonization especially in PBMC macrophages that the
283 magnitude of increase was significantly smaller than FMT mice (Fig. 4B, 16.4% versus
284 37.5%), indicating that the expression of this surface molecule might not be only regulated
285 by *B. coccoides*, but is dependent on other intestinal bacteria as well. Likewise, C.B
286 colonized-mice remained low level of either MHC-I- or CD80-upregulated PBMC and

287 spleen macrophages compared to Abx mice following viral infection at 3 dpi. To determine
288 whether *B. coccoides* mediates protection from EMCV systemic infection through a
289 macrophage-dependent mechanism, we then colonized macrophage-depleted Abx mice
290 with *B. coccoides* before EMCV intraperitoneal inoculation and observed the survival
291 curves for 14 days. It was shown that clodronate treatment abolished B.C-mediated
292 protection, demonstrating that B.C is required for conferring macrophage-mediated
293 protection against EMCV infection (Fig. 5C).

294 To determine whether colonizing Abx mice with *B. coccoides* restores systemic IFN-I
295 responses to viral infection, we harvested PBMCs from Abx mice with B.C, C.B or fecal
296 contents colonization at 3 dpi and subjected to IFN- β expression measurement by qPCR.
297 Apparently, B.C colonization in Abx-treated mice recovered IFN- β expression in PBMCs
298 to the level of that detected in FMT and PBS-treated controls upon EMCV infection at 3
299 dpi, whereas C.B colonization did not (Fig. 5D). Since we have demonstrated that intestinal
300 microbiome-driven IFN-I amplification pathway is absolutely essential for antagonizing viral
301 systemic infection, we next investigated whether *B. coccoides* mediates protection from
302 EMCV infection in an IFN-I dependent manner. We colonized *Ifnar*^{-/-} mice with B.C prior to
303 virus inoculation and discovered that these bacterium colonized-mice did not display less
304 susceptibility to EMCV infection than Abx *Ifnar*^{-/-} mice (Fig. 5E). Furthermore, B.C
305 colonization of Abx-treated *Ifnar*^{-/-} mice did not reduce viremia or brain replication in
306 comparison to Abx *Ifnar*^{-/-} controls at 1 dpi (Fig. 5F). As a whole, these data demonstrate
307 that the protective effect of *B. coccoides* against viral systemic infection required
308 macrophages and IFN-I amplification signaling.

309 ***B. coccoides* colonization promotes type I IFN responses in macrophages to limit**

310 **EMCV infection**

311 To directly clarify whether *B. coccoides* can promote systemic type I IFN responses through
312 macrophages activation, we infected bone-marrow-derived macrophages (BMDMs)
313 isolated from PBS-, Abx-treated, B.C-colonized, or C.B colonized mice with EMCV at MOI
314 of 5 in vitro and detected induction of *Irfn* and associated ISGs at 8 hpi using qRT-PCR.
315 Expression of *Irfn* and ISGs including *Oas1a*, *Isg15* and *Mx1* was drastically reduced in
316 BMDMs isolated from Abx mice compared to PBS mice, suggestive of an intrinsic
317 incapability of responding to viral infection, but this responsiveness was partially regained
318 by B.C, not C.B colonization (Fig. 6A). Furthermore, B.C colonization in Abx mice endowed
319 their BMDMs with the ability to induce ISGs expression in an *Irfn*-dependent manner for
320 ISGs expression was undetectable (or extremely low) in BMDMs isolated from B.C-
321 colonized *Irfn*^{-/-} mice following EMCV infection at 8 hpi (Fig. 6B). As STAT1 conveys
322 signals downstream of IFN receptor engagement and its phosphorylation and nuclear-
323 translocation mediates transcription of ISGs in the IFN-I amplification loop, thus we
324 stimulated macrophages isolated from spleen of PBS, Abx or B.C colonized-Abx mice with
325 IFN- γ in vitro and tested STAT1 phosphorylation using flow. IFN- γ -stimulated
326 macrophages isolated from Abx mice exhibited significantly low level of pSTAT1 compared
327 to those from PBS mice, while B.C colonization in Abx mice resulted in recovered STAT1
328 phosphorylation in splenic macrophages comparable to PBS controls (Fig. 6C and Fig. S5).
329 We also looked at expression of multiple antiviral defense genes expression in
330 macrophages isolated from the spleen of infected PBS, Abx, B.C-colonized animals to

331 confirm whether same phenotype exists in vivo post viral infection. At 1 dpi, splenic
332 macrophages from Abx mice exhibited significantly decreased antiviral gene expression
333 compared to those from PBS mice, but B.C colonization in Abx mice induced significantly
334 higher level of *Irfnb* and ISGs, although not fully restored to the level of PBS controls (Fig.
335 6D). This phenotype is consistent with the data showing B.C colonization partially restored
336 the frequency of CD80-expressing macrophages (Fig. 5C) in PBMC, implying other
337 commensal bacteria species also possess the capability of inducing type I IFN and ISG
338 responses in circulating macrophages. Again, B.C exerts its effects through IFNAR-
339 mediated IFN-I signaling because no detectable or obviously upregulated antiviral gene
340 expression was observed in macrophages sorted from spleen of B.C colonized Abx *Irfnar*^{-/-}
341 mice (Fig. 6E). Overall, these findings suggest that *B. coccoides* is sufficient for maintaining
342 optimal macrophage responsiveness to stimuli (cytokine and viral infection) and it is very
343 likely that *B. coccoides* augments type I IFN signaling in macrophages via IFNAR and
344 STAT1 phosphorylation to protect host from virus systemic infection.

345

346 **Discussion**

347 In the present study, we have described a crucial role for the commensal microbiota in
348 modulating antiviral innate immunity towards enteric virus systemic infection. In particular,
349 the intestinal microbiome protects systemic viral infection by promoting type I IFN
350 responses in peripheral innate immune cells, mainly macrophages, via an *Irfnar*-dependent
351 signaling pathway. Furthermore, we have identified the mechanism by which an under-
352 studied commensal bacteria species, *B. coccoides*, utilizes to restrict enteric virus systemic

353 infection through enhancing type I IFN responses in macrophages.

354 Over the past ten years, it has been gradually acknowledged that the intestinal microbiome
355 can shape host antiviral immunity. Accumulating literature has shown that animals
356 administered with cocktails of broad-spectrum antibiotics or GF mice have impaired innate
357 and adaptive immune responses following infections by various viral pathogens including
358 LCMV, influenza virus, murine cytomegalovirus (MCMV), flaviviruses, VSV or CHIKV [13,
359 15, 16, 18, 22]. However, up till now, all studies on mammalian enteric viruses have
360 reported that intestinal microbiome promotes virus infection [6, 7, 10, 20, 23], with two
361 exceptions. One conducted by Grau et al reported that *C. scindens* primes type III
362 interferon induction to suppress murine norovirus infection in proximal small intestines by
363 bile acids biotransformation[19]. Another example of commensal bacteria-mediated anti-
364 enteric virus protection was provided by Shi et al, demonstrating that segmented
365 filamentous bacteria (SFB) prevent and cure rotavirus infection by accelerating epithelial
366 cell turnover [24]. Unlike norovirus and rotavirus infection only causes self-limited diseases,
367 some enteric viruses, enterovirus 71 and EMCV for instance, can result in systemic
368 infection and lead to a diverse array of neurological disease [25, 26]. To answer the
369 question whether intestinal microbiota could be protective indirectly by regulating host
370 antiviral innate immunity during enteric virus infection, we infected wild-type mice with
371 EMCV per orally to mimic its natural infection, or through intraperitoneal route to
372 immediately establish systemic infection. To our surprise, microbiota depletion with
373 antibiotics exacerbates the neurological disease and enhances viral replication upon
374 EMCV infection under both routes of inoculation (Fig.1). This phenotype resembles the

375 results reported by previous studies on non-enteric viral pathogens mentioned above, but
376 not similar to the milestone study conducted by Kuss et al using poliovirus and reovirus, in
377 which intestinal microbiomes and LPS are reported to promote viral replication and
378 systemic pathogenesis [6]. These discrepancies suggest that microbiota acts very
379 differently in interacting with even two closely related enteric viruses (EMCV and poliovirus
380 are both picornaviruses that share very similar viral structure). It is highly likely that the
381 innate immune responses were also severely diminished by antibiotics treatment in their
382 model (we used the same antibiotic regimens) whereas poliovirus replication in vivo was
383 barely established to reflect the defects on host innate immunity in the absence of
384 microbiota. Conversely, EMCV in vivo replication and dissemination does not require
385 microbiota in our model (Fig. 1B and 1C) so that the neuropathogenesis was deteriorated
386 in the context of antiviral innate immunity being dampened by microbiota depletion.

387 Collectively, multiple studies have demonstrated that commensal bacteria are responsible
388 for shaping host antiviral immunity beyond mucosal sites. GF mice and Abx-treated mice
389 exhibited blunted type I interferon responses that are required for optimizing functions of
390 nonmucosal immune cells including NK cells, macrophages, or CD8⁺ T cells [13, 14, 16,
391 22]. In our study, we showed that macrophages isolated from Abx mice did not display
392 higher frequency or upregulated expression of activating surface markers like MHC-I or
393 CD80 at 3 dpi (Fig. 2A and 2D). Additionally, macrophage depletion equalized the
394 discrepancies in mortality and viral replication between Abx-treated mice and PBS controls,
395 indicating that macrophages are essential for microbiota-mediated protection from EMCV
396 systemic infection (Fig. 2E and 2F). Our data are in consistent with previous studies

397 demonstrating that microbiome can regulate antiviral macrophage responses by inducing
398 the expression of type I IFN and ISG genes after influenza virus infection [15, 16]. Similar
399 to these studies, we also showed that microbiota depletion results in severely diminished
400 type I IFN and ISG responses both in periphery and brain (Fig. 3A-D), which causes
401 unrestricted virus replication and exacerbates neuropathogenesis and mortality. The
402 impaired IFN-I production and ISG expression in the brain tissues of Abx mice suggests
403 major defects in responsiveness of microglia to viral infection because they are the main
404 sources of IFN production in brain. The results are consistent with one manuscript reporting
405 that microglia of GF mice were unable to produce various cytokines and chemokines upon
406 stimulation[27]. Our model figured out that the microbiota-driven IFN and ISG responses
407 are partially dependent on *Irf3*-mediated IFN-I induction and are fully relied on *Ifnar*-
408 mediated signaling pathway (Fig. 3E and 3F).

409 Although accumulated evidences has shown that microbiota can promote ISG responses in
410 multiple cell types, the cellular sources of type I IFN are yet less-well defined until very
411 recently, where two papers demonstrated that microbiota constitutively induces type I IFN
412 production and basal level of ISG expression in pDCs at systemic sites [17, 18]. In our
413 study, we only looked at the frequency of pDCs in PBMC or spleen following EMCV
414 infection at 3 dpi, and observed no differences between PBS and Abx mice (Fig. 2A).
415 Similarly, Abt et al reported that at 3 dpi following influenza virus infection, there was a
416 comparable influx of pDCs into bronchiole alveolar lavage and the pDCs exhibited similar
417 activation profile in PBS and Abx mice [15]. These seemingly discordant observations
418 could be explained by the timing of detection because pDCs respond to virus nucleic acids

419 with massive and rapid secretion of IFN-I (1-3 h post stimulation) independently of the
420 IFNAR-based feedback signaling that is always required for most cell types in IFN-I
421 production [28]. Also, profiling the IFN-I and ISG expression in pDCs could be very subtle
422 especially at steady state (0 dpi) and using different microbiota-deficient mouse model (GF
423 versus Abx-treated) may result in distinct phenotype [17, 18]. Generally, it is plausible that
424 signals of commensal bacteria drive a very swift first wave of IFN-I (especially IFN- β)
425 production in poised pDCs to activate NK cells, macrophages and overall innate immune
426 responses to viral infection.

427 Up till now, only a few publications have begun to identify specific commensal bacterial
428 species that play an important role in inducing type I IFN-mediated innate antiviral immunity
429 [16, 18, 29, 30]. Although Abt and colleagues' study revealed an interplay between
430 commensal bacteria and poised low-level tonic antiviral interferon signaling, they have not
431 figured out the specific bacterial species that regulates the 'steady-state' readiness of
432 antiviral pathways in macrophages [15]. Our work builds on the limited existing literature
433 by identifying a less documented-commensal bacterium, *B. coccoides* that plays a crucial
434 role in protecting from enteric virus systemic infection through IFN-I induction. Although the
435 genus of *Blautia* only comprises less than 4% of total colonic microbes (Fig. 4E), this
436 bacterial community seems to be highly capable of activating macrophages as *B.*
437 *coccoides* colonialization almost fully recovered the upregulation of MHC-I and CD80 on
438 the surface of PBMC and spleen macrophages in Abx-treated animals following EMCV
439 infection (Fig. 5B and 5C). Single colonization of *B. coccoides* in Abx mice can also rescue
440 the intrinsic defects in expression of IFN-I and key ISGs in BMDMs in an *Ifnar*-dependent

441 manner following EMCV infection in vitro (Fig. 6A and 6B), or restore the STAT1
442 phosphorylation in splenic macrophages following IFN- γ stimulation in vitro (Fig. 6C). The
443 present study adds a “missing piece” to the previous work by demonstrating a single
444 commensal bacteria strain *B. coccoides* can restore the impotency of macrophages in
445 responsiveness to type II IFN or virus because of microbiota depletion.

446 Since *B. coccoides* monocolonization in Abx-treated mice restored protection of EMCV
447 systemic infection by restricting in vivo viral replication in an IFNAR- and macrophage-
448 dependent manner (Fig. 4F, 4G; 5D and 5F), a next question will be what does *B. coccoides*
449 utilize to induce IFN-I responses in macrophages to confer antiviral systemic protection?

450 Two recent reports suggest that *C.scindens*-derived metabolite deoxycholic acid or
451 glycolipids of *B.fragilis* can impact host antiviral innate immunity by inducing type I IFN
452 production and ISG expression in pDCs or a subset of colonic cDCs [18, 30]. Short-chain
453 fatty acids (SCFA), primarily produced by Gram-positive bacteria Firmicutes and
454 Bacteroidetes through fermentation of undigested polysaccharides, are not only important
455 local energy sources for gut microbiota and intestinal epithelial cells, but also crucial
456 regulators for shaping immune systems at extraintestinal sites [5]. As the most abundant
457 SCFA in the colon, nearly one-third of acetate is produced by acetogenic bacterial
458 communities such as *Blautia* [31]. It was shown that acetate derived from high-fiber diet
459 protects against respiratory syncytial virus infection by promoting type I IFN production and
460 ISG expression in pulmonary epithelial cells through GPR43 [32]. Future studies will be
461 carried out to examine the effects of specific SCFAs on influencing host anti-enteric virus
462 innate immunity using microbiota deficient mouse model.

463

464 **Conclusion**

465 Our present study demonstrates how fundamental innate immune mediators such as
466 macrophages and type I IFN are regulated by commensal bacteria to antagonize enteric
467 virus systemic infection. In particular, we have identified a novel commensal bacterial strain,
468 *B. coccoides* can restrict enteric virus replication and neuropathogenesis by activating IFN-
469 I and ISG responses in macrophages via IFNAR- and STAT1-mediated signaling pathway.
470 This finding may provide insight into the development of novel therapeutics in preventing
471 enteric virus infection or possibly, mitigating its clinical diseases.

472

473 **Methods**

474 **Cell, bacteria isolates and virus**

475 BHK-21 cells were cultured in DMEM supplemented with 10% fetal bovine serum
476 (GIBCO™, Invitrogen Corporation, Carlsbad, CA, USA) at 37°C/5% CO₂.
477 *B. coccoides* was purchased from ATCC (ATCC 29236) and cultured in Modified Chopped
478 Meat Medium (ATCC Medium 1490, ELITE-MEDIA) at 37°C under anaerobic conditions.
479 *C. butyricum* was purchased from ATCC (ATCC 19398) and cultured in Thioglycollate
480 Medium (Hopebio) at 37°C under anaerobic conditions. *A. muciniphila* was kindly provided
481 by Dr. Lan-juan Li (The First Affiliated Hospital, College of Medicine, Zhejiang University,
482 Hangzhou, P. R. China) and cultured in Brain Heart Infusion Medium (BHI, OXOID) at 37°C
483 under anaerobic conditions. *L. reuteri* was purchased from China Center for Type Culture
484 Collection (CCTCC, AB 2014289) and cultured in DeMan-Rogosa-Sharpe medium

485 (OXOID) at 37°C under anaerobic conditions. The concentration of each bacterial species
486 was quantified based on CD600.

487 Recombinant virus derived from full-length clone of EMCV strain BJC3 (GenBank:
488 DQ464062) was used in this study [33] and viral titers of EMCV stocks were determined
489 by standard TCID₅₀ assay described previously [34].

490 **Mice and infections**

491 Mice were maintained in specific pathogen-free (SPF) facility with temperature- and
492 humidity-controlled environment (22 ± 2°C, 50 ± 10% humidity), and all animal experiments
493 were strictly carried out in accordance with protocols approved (#117113) by the
494 Institutional Animal Care and Use Committee of Zhejiang University. Six- to eight-week-old,
495 sex-matched mice were used for all experiments. The C57BL/6J wildtype mice were
496 purchased from the Model Animal Research Center of Nanjing University (Nanjing, China).
497 Type 1 interferon receptor knockout mice (referred to as *Ifnar*^{-/-}) were kindly gifted by Dr.
498 Yu Chen (Wuhan University, Hubei, P. R. China) and interferon regulatory factor 3 deficient
499 mice (referred to as *Irf3*^{-/-}) mice were provided by Dr. Jin Jin (Zhejiang University, P. R.
500 China).

501 For all wild-type B6 mice studies, animals were inoculated orally with 2×10⁷, 2×10⁵ or
502 2×10³ TCID₅₀ units of EMCV in 25 μL inoculum, or infected intraperitoneally with 200
503 TCID₅₀ units of EMCV in 200 μL inoculum. For EMCV virulence assay on *Ifnar*^{-/-}, *Irf3*^{-/-}
504 mice or wild-type mice treated with clodronate liposomes, animals were inoculated
505 intraperitoneally with 50 TCID₅₀ units of EMCV in 200 μL inoculum. Clinical symptoms
506 were observed and documented blindly using following scoring criteria: 0, normal; 1,

507 hunch-back and trembling; 2, hind limb paralysis; 3, dyspnea and unresponsive to touch;
508 4, sudden death.

509 For viral titer determination, tissue samples at indicated time points were harvested,
510 weighed, and homogenized with stainless beads in 1 mL of DMEM media supplemented
511 with 2% FBS and titrated by qPCR. Briefly, tissue samples were homogenized at 45 Hz for
512 1 minutes and the homogenates were clarified by centrifugation at 12,000 rpm for 5 min.
513 Total RNA was extracted with Trizol Reagent (Invitrogen) and subjected to quantitative
514 reverse transcriptase PCR (qRT-PCR) using One-step qPCR Kits (TOYOBO) on an ABI
515 7500 Fast Instrument. Standard cycling conditions and primers/probe (forward: 5'-
516 TGAGTCATTAGCCATTTCAACCCA-3'; reverse: 5'-CGTGAGATACAAACCCGCCCTA-3';
517 probe: 5'-TCCCATCAGGTTGTGCAGCGA-3') were described previously [35]. Viral
518 burden was expressed on a log₁₀ scale as EMCV genomic RNA equivalents per milligram.

519 **Antibiotics treatment, fecal microbiota transplantation (FMT) and bacterial**
520 **colonization**

521 Mice were administrated with antibiotic cocktail comprised of 10 mg each of ampicillin,
522 neomycin, metronidazole and vancomycin (167 mg/μL) daily for 5 days via oral gavage.
523 After the fifth day of oral gavage, antibiotics were added to the drinking water at a
524 concentration of 1 g/L for ampicillin, neomycin and metronidazole and 500 mg/L for
525 vancomycin. Fecal samples collected from microbiota-depleted mice at the 5th day post
526 treatment were homogenized, plated on brain-heart infusion (BHI) agar with 10% sheep
527 blood and cultured under anaerobic conditions at 37°C for 2 d followed by aerobic
528 conditions at 37°C for 1 d to confirm efficient microbial depletion. Animals were maintained

529 with Abx- or PBS-containing water for the duration of the experiment [6, 20].
530 For FMT experiments, 200 mg of pooled feces pellets were homogenized with sterile silica
531 beads in 1.5 ml PBS at 45 Hz for 1 minutes and filtered with 70 μ m strainers. Mice were
532 administrated Abx as described above, Abx administration was discontinued on day 6 and
533 Abx mice were gavaged with 150 μ l filtered stool homogenates individually [16]. For
534 bacterial colonization experiments, Abx mice were gavaged with 10^{10} cfu of *B. coccoides*,
535 *A. muciniphila*, *C.butyrum* or *L. reuteri* in 150 μ l PBS at the 6th day post Abx oral
536 administration. At 48 hours post FMT or bacterial colonization (on day 8 post Abx oral
537 gavage), stool samples were collected to determine the efficiency of colonization and
538 colonized-mice were inoculated intraperitoneally with EMCV at indicated doses.

539 **Immunofluorescent assay (IFA) and viral signal quantification**

540 Brain tissues were collected from either mock or infected mice at 3 or 5 dpi and fixed in 4%
541 buffered paraformaldehyde and embedded in paraffin. Deparaffinized brain sections were
542 incubated with 10% normal goat serum for 30 min to block nonspecific binding, co-stained
543 with anti-EMCV VP2 (1: 500) and anti-GFAP monoclonal antibodies (1: 1200, Abcam), or
544 co-stained with anti-VP2 and anti-IBA1 monoclonal antibodies (1: 800, Invitrogen).
545 Following primary antibodies incubation at 4°C overnight, a Cy3-conjugated anti-mouse
546 secondary antibody at a dilution of 1:100 (Santa Cruz Biotechnology, Santa Cruz) was
547 added to the sections and incubated for 30 min at room temperature. Stained sections
548 were imaged with a NIKON ECLIPSE C1 Upright Fluorescent Microscope. The number of
549 antigen-positive cells from 10 fields of view (200 \times) was combined when a single section
550 was observed and averaged over three coronal brain sections per mouse from six infected

551 mice [33].

552 **Splenocytes, bone marrow derived macrophage and peripheral blood mononuclear**
553 **cells generation and treatment**

554 Splens were dissected and single cell suspensions of splenocytes were generated by
555 grinding through a 70-mm strainer. Erythrocytes were lysed with Red Blood Cell Lysis
556 Buffer (RCLB, HyClone), and remaining cells were resuspended in PBS supplemented
557 with 2% FBS and 1 mM EDTA.

558 Bone marrow derived macrophages (BMDMs) were generated by isolation of bone marrow
559 cells from mouse femurs and tibiae after sacrifice. Cells were cultured at 37°C in DMEM
560 supplemented with 20% FBS and 30% supernatant of filtered-L929 cells for 7 days prior to
561 experimental procedure. BMDMs were infected with EMCV at MOI 5 for 8 hours prior to
562 harvest for indicated groups.

563 Peripheral blood mononuclear cells (PBMCs) isolation was performed by using density
564 gradient centrifugation with a PBMC isolation kit (TBD).

565 **IFN-β protein analysis**

566 IFN-β protein quantification was performed using Mouse IFN-beta ELISA Kit (ABclonal)
567 according to manufacturer's instructions.

568 **Cytokine expression analysis**

569 RNA was isolated from tissues or cells using Trizol Reagent (Invitrogen) as per
570 manufacturer's instructions. Gene expression levels of *Ifnb*, *Isg15*, *Isg56*, *Irf7*, *Irf9*, *Stat1*,
571 *Stat2*, *Oas1a*, *Mx1* and *Cxcl10* (primers used for the assay were listed in Table. S1) were

572 determined via qRT-PCR and normalized to GAPDH expression. Results are presented
573 as fold changes of cytokine expression in infected mice over mock animals ($2^{-\Delta\Delta Ct}$).

574 **Macrophages depletion**

575 Macrophages depletion experiments were performed as previously published literatures.
576 Briefly, animals were injected with 250 μ l per mouse of clodronate liposomes (YEASEN)
577 or control liposomes intraperitoneally 2 days prior to Abx treatment, the day of Abx
578 treatment, and then on days 2 and 5 post Abx treatment.

579 **DNA extraction, 16S rRNA amplicon sequencing and data analyses**

580 Genomic DNA of the fecal samples was extracted using the ALFA-SEQ Advanced Stool
581 DNA Kit (Magen). The quality and quantity of DNA were measured using a NanoDrop One
582 (Thermo Fisher Scientific). Subsequently, Bar-coded PCR primers targeting V3-V4 region
583 of bacterial 16S rRNA genes were used to generate amplicons (Forward:
584 ACTCCTACGGGAGGCAGCA; Reverse: GGACTACHVGGGTWTCTAAT) and multiplex
585 sequencing of amplicons with sample-specific barcodes was performed using an Illumina
586 Novaseq 6000 platform (paired end 2×250 nucleotide reads, Guangdong Magigene
587 Biotechnology Co., Ltd. Guangzhou, China).

588 The raw data of sequencing were filtered using the fastp (version 0.14.1,
589 <https://github.com/OpenGene/fastp>) with parameters -W 4 -M 20 and further processed by
590 the cutadapt (<https://github.com/marcelm/cutadapt/>) to remove the primer sequences to
591 obtain the paired-end clean data. Subsequently, the usearch -fastq_mergepairs tool
592 (version 10, <http://www.drive5.com/usearch/>) was utilized to merge the raw tags, which
593 were later trimmed by the fastp to get the clean tags. The operational taxonomic units

594 (OTUs) were then clustered with a cut-off value of 97% similarity using the UPARSE
595 software. Each representative sequence of these OTUs were assigned using the SILVA
596 database to annotate taxonomic information. The richness of certain commensal bacteria
597 taxa was calculated using the usearch -alpha_div (version 10,
598 <http://www.drive5.com/usearch/>) according to the OTU abundance. Based on the relative
599 abundance of species at each classification level in otu_table, R software was used to
600 draw the histogram, heat map and ternary phase diagram.

601 **Flow cytometry and cell sorting**

602 Splenocytes and PBMCs were harvested to analyze levels of different antigen on the
603 surface of different cell subsets following blockade of Fcγ receptors with anti-CD16/32
604 (eBioscience). Fluorescently conjugated antibodies used include those specific to CD3,
605 CD4, CD8, CD19, CD11b, CD11c (Biolegend), Ly6c, F4/80, MHC II and PDCA-1
606 (eBioscience). Inflammatory monocytes were identified as Ly6c⁺ and CD11b⁺. PBMC and
607 splenic macrophages were identified as F4/80 and CD11b double positive. NK cells were
608 identified as CD3⁻ and NK1.1⁺. cDCs were identified as F4/80⁻, CD11c⁺ and MHC-II^{hi}. pDCs
609 were identified as CD11c^{int} and PDCA-1⁺. CD4 T cells, CD8 T cells and B cells were
610 identified as CD3⁺/CD4⁺, CD3⁺/CD8⁺ or CD19⁺ respectively. For Intracellular staining, cells
611 were permeabilized with Cytotfix/Cytoperm buffer (BD), and stained for INF-γ or granzym
612 B. Splenic macrophages were sorted as F4/80 and CD11b double positive subsets by
613 flow cytometry using the BD FACSVerser.

614 **In vitro phosflow STAT1 staining of macrophages**

615 Splenic macrophages were stimulated with 200 ng/ml recombinant IFN-γ (R&D Systems)

616 for 16 hours, media was subsequently removed and replaced with 0.05% trypsin and
617 incubated at 37°C for 2 min. Cells were then fixed with 4% PFA for 10 min, stained for
618 surface markers, permeabilized with Cytofix/Cytoperm buffer, and stained for pSTAT1 with
619 PE-conjugated anti-STAT1 (pY701) antibody (Biolegend).

620 **Statistical Analysis**

621 Statistical analyses were performed with Prism GraphPad software v 8.0. Error bars
622 represent standard errors of mean in all figures and *P* values were determined by unpaired,
623 two tailed Student's t test. Log-rank test was used for survival curves. (**p* < 0.05; ***p* < 0.01;
624 ****p* < 0.001; **** *p* < 0.0001).

625

626 **List of abbreviations**

627 type I interferon (IFN), encephalomyocarditis virus (EMCV), antibiotic cocktail-
628 administrated mice (Abx), *Blautia coccooides* (*B. C*), interferon-stimulated genes (ISGs),
629 germ-free (GF), plasmacytoid dendritic cells (pDCs), conventional dendritic cells (cDCs),
630 Chikungunya virus (CHIKV), *Clostridium scindens* (*C. scindens*), Toll-like receptor 7
631 (TLR7), secondary bile acid (BA), Vancomycin (Van), Neomycin (Neo), ampicillin (Amp)
632 and Metronidazole (Metro), days post-infection (dpi), ecal microbiota transplantation (FMT),
633 peritoneal blood mononuclear cells (PBMCs), lymphocytic choriomeningitis virus (LCMV),
634 hours post-infection (hpi), *Akkermansia muciniphila* (*A. muciniphila*, herein referred to as
635 AKK), *Lactobacillus reuteri* (*L. reuteri*, herein referred to as L.R), *Clostridium butyricum* (*C.*
636 *butyricum*, herein referred to as C.B), murine cytomegalovirus (MCMV), segmented
637 filamentous bacteria (SFB)

638 **Declarations**

639

640 **Ethics approval and consent to participate**

641 Not applicable.

642

643 **Consent for publication**

644 Not applicable.

645

646 **Availability of data and materials**

647 Not applicable.

648

649 **Competing interests**

650 The authors declare that they have no competing interests.

651

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656

657 **Author's contributions**

658 Xiao-Lian Yang, Gan Wang, Jin-Yan Xie conceptualized the project. Shu J. Zhu, Gan Wang

659 and Jin-Yan Xie wrote the paper. Wei Liu and Shu J. Zhu reviewed and edited the paper.

660 Xiao-Lian Yang performed the virulence studies, viral loads titration, IFN-I and ISGs
661 expression analysis, IFA, flow cytometry of immune cells and activation of macrophages.
662 Gan Wang performed bacteria colonization, pathogenesis studies, macrophage sorting
663 and pSTAT1 staining, IFN-I and ISGs expression analysis. Jin-Yan Xie analyzed and
664 interpreted 16S rRNA sequencing data. Han Li extracted DNA from fecal samples.

665

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671

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779

780 **Figure legends**

781 Fig.1. Microbiota depletion deteriorates EMCV pathogenesis and viral replication in vivo.

782 (A) Groups of 6-8 weeks old C57BL/6J mice were pretreated with PBS or Abx for 5 days,

783 inoculated with different dose of EMCV orally and observed for 14 days. Survival curves

784 (n=12-17) and pathological scores (n=10-15) were documented. (B) Blood and brain

785 samples of PBS or Abx mice inoculated with 2×10^7 TCID₅₀ units of EMCV orally were

786 harvested and tested for viral loads at 3 and 5 dpi by qPCR (n=6). (C) Indirect fluorescent

787 assay (IFA) of brain sections from PBS- or Abx-infected mice at 5 dpi. EMCV capsid protein

788 VP2 and cellular surface markers were double-stained with respective antibodies

789 (representative images from n=6 per group). (D) Groups of PBS, Abx or Abx mice received

790 FMT from conventionally housed PBS naïve mice were intraperitoneally inoculated with

791 200 TCID₅₀ units of EMCV for survival kinetics analysis (n=13-17). (E) Viral titers of brain

792 tissues collected from PBS-, Abx- or FMT-infected mice at 1 and 3 dpi (same dose and

793 route as Fig. 1D, n=5-6). Data represented in the figures were from two independent

794 experiments. Dotted dash lines indicate the limit of detection (LD).

795 Fig. 2. Innate cellular immune responses were severely diminished in Abx-treated mice

796 post EMCV infection. (A) Frequency of a panel of innate immune cells including

797 inflammatory monocytes, NK cells, macrophages, pDC or cDC in PBMC or splenocytes of

798 mock or infected PBS or Abx mice at 3 dpi (n=4-8). (B) Frequency of various adaptive

799 immune cells including CD4⁺ T cells, CD8⁺ T cells or B cells from PBMC or spleen of mock

800 or infected PBS or Abx mice at 3 dpi (n=6). (C) Expression of MHC-I and CD80 on PBMC

801 macrophages from mock- or infected-PBS or Abx mice at 3 dpi (upper two panels,
802 histograms of MHC-I- or CD80-expressing macrophages; lower two panels, frequency of
803 macrophages expressing MHC-I or CD80 in total PBMC macrophages; n=6). (D) Survival
804 analysis of PBS or Abx mice treated with clodronate liposomes or control reagent following
805 EMCV intraperitoneal inoculation (n=10-11). (E) Blood viral titers of PBS or Abx mice
806 treated with clodronate liposomes or control reagent at 1 dpi (n=6). Data presented in the
807 figure were from two independent experiments.

808 Fig.3. Microbiota depletion results in significantly impaired systemic or cerebral type I IFN
809 responses. Fold induction of *Ifnb* and ISG expression in PBMC (A) and spleen (B) of PBS
810 or Abx mice at 12 hpi following EMCV infection relative to respective mock controls (n=5).
811 (C) Expression of *Ifnb* and ISGs in the brain of infected PBS- or Abx-treated mice (fold-
812 change compared to respective mock mice) at 1 or 3 dpi (n=5-6). IFN- β levels in the brain
813 at 3 dpi as detected by ELISA (middle panel). (D) Fold induction of proinflammatory or
814 immune regulatory cytokines expression in the brain at 1 or 3 dpi relative to respective
815 mock mice (n=6). (E) Survival kinetics of PBS- or Abx-treated *Irf3*^{-/-} or *Ifnar*^{-/-} infected with
816 EMCV intraperitoneally, wild-type C57BL/6J mice were served as control (left panel, n=10-
817 12; middle panel, n=9-14; right panel, n=13-15). (F) Viral titers in blood and brain collected
818 from infected PBS or Abx WT, *Irf3*^{-/-} or *Ifnar*^{-/-} mice with at 1 dpi (n=5-6). All data presented
819 in the figure were from two independent experiments.

820 Fig.4. *B. coccoides* protects EMCV systemic infection by restricting viral in vivo replication.
821 Viral titers (A) and induction of *Ifnb* (B) in the brain of PBS-, Abx- or single antibiotic-treated
822 mice were measured by qPCR at 3 dpi (n=5-6). (C) Bacterial richness was defined by the

823 number of the unique taxa. (D) Heat map of relative abundance from different groups of
824 antibiotic treatment. (E) Relative abundance changes of bacteria at genus level. (F)
825 Survival analysis of Abx mice colonized with *B. coccoides* (B.C), *A. muciniphila* (AKK), *L.*
826 *reuteri* (L.R), *C. butyricum* (C.B), or mouse fecal contents (FMT) following EMCV
827 intraperitoneal inoculation (3 experiments, n=16-20). (G) Viral burden in brain tissues
828 harvested from mock- or infected-PBS, Abx or Abx mice colonized with FMT, B.C or C.B at
829 3 dpi (n=6). All data presented in the figure were from at least two independent experiments.

830 Fig.5. *B. coccoides* colonization restricts enteric virus systemic infection by activating
831 innate cellular immune responses and type I IFN expression in a macrophage- and *Ifnar*-
832 dependent manner. Frequency of macrophages expressing MHC-I (A) or CD80 (B) in
833 PBMC or spleen isolated from PBS, Abx or Abx mice colonized with FMT, B.C or C.B at 3
834 dpi following EMCV inoculation (representative flow cytometry histograms of showing
835 surface staining of MHC-I or CD80, n=6). (C) Survival kinetics of infected Abx and B.C-
836 colonized Abx mice injected with clodronate liposomes (n=9-10). (D) Expression level of
837 *Ifnb* in PBMC isolated from infected PBS, Abx, or Abx mice gavaged with FMT, B.C or C.B,
838 comparing to their respective mock controls (n=5). (E) Survival kinetics of infected Abx and
839 B.C-colonized Abx *Ifnar*^{-/-} mice (n=6-9). (F) Viral titers in blood and brain collected from
840 infected PBS, Abx or B.C-colonized WT or *Ifnar*^{-/-} mice at 1 dpi (n=5-6). All data presented
841 in the figure were from two independent experiments.

842 Fig.6. *B. coccoides* colonization promotes type I IFN and ISG responses in macrophages
843 to limit EMCV infection. (A) *Ifnb* and antiviral defense gene expression in BMDMs isolated
844 from PBS, Abx, B.C- or C.B-colonized Abx mice at 8 h post EMCV infection (MOI=5) in

845 vitro (n=5-6). (B) Relative mRNA expression of *Oas1a* and *Mx1* in BMDMs isolated from
846 WT or *Ifnar*^{-/-} mice that pretreated with PBS, Abx or Abx plus *B. coccoides* colonization at
847 8 hpi in vitro (n=5-6). (C) Splenic macrophages sorted from spleen of PBS, Abx or B.C-
848 colonized Abx mice were stimulated with 200 ng/ml IFN- γ in vitro. The data are reported
849 as the averaged frequency of pSTAT1-stained CD11b⁺ cell subsets at 16 h subtracted by
850 the frequency of these cells tested at 0 h post IFN- γ stimulation (3 experiments, n=6). (D)
851 *Ifnb* and antiviral defense gene expression in splenic macrophages isolated from EMCV-
852 infected PBS, Abx, or B.C-colonized Abx mice at 1 dpi (n=6). (E) Relative mRNA
853 expression of *Cxcl10* and *Stat2* in splenic macrophages isolated from EMCV-infected WT
854 or *Ifnar*^{-/-} mice that pretreated with PBS, Abx or Abx plus *B. coccoides* colonization at 1 dpi
855 (n=5-6). Data presented in the figure were from at least two independent experiments.

856 Fig. S1. Flow cytometry gating strategy of NK cells in PBMC of mock or infected mice w/wo
857 Abx treatment at 3 dpi (related to Fig. 2). Representative flow cytometry plots showing
858 gating scheme for NK1.1-PE positive cells in PBMC isolated from PBS mock, Abx mock,
859 PBS-infected or Abx-infected mice.

860 Fig. S2. Flow cytometry gating strategy of macrophages in PBMC of mock or infected mice
861 w/wo Abx treatment at 3 dpi (related to Fig. 2). Representative flow cytometry plots showing
862 gating scheme for F4/80-CD11b double-positive cell subsets in PBMC isolated from PBS
863 mock mice and representative histogram image of MHC-I^{hi} (A) or CD80 (B) staining on the
864 surface of these cell subsets (left, lower panel); same gating strategy is utilized for all mice
865 groups and the histogram overlay images are presented (right, lower panel).

866 Fig. S3. Depletion of splenic macrophages following administration of clodronate

867 liposomes (related to Fig. 2). Histograms showing CD11b⁺ cells isolated from spleen of
868 mice treated with control reagents (A) or clodronate liposomes (B).

869 Fig. S4. Relative abundance of bacteria strains in fecal samples collected from single
870 bacterium colonized-mice. Groups of Abx mice were colonized with *B. coccoides* (A) or *C.*
871 *butyricum* (B) and fecal relative bacterial abundance at 48 h post colonization was
872 determined by 16S rRNA sequencing (n=6).

873 Fig. S5. Flow cytometry gating strategy of sorted splenic macrophages from PBS, Abx or
874 B.C-colonized Abx mice that stained for pSTAT1 post IFN- γ stimulation in vitro (related to
875 Fig.6). Representative flow cytometry plots showing sorting scheme for F4/80-CD11b
876 double-positive cell subsets of splenocytes isolated from PBS mice (upper panels) and
877 representative histogram image of intracellular staining of pSTAT1 in sorted macrophages
878 at 0 h post IFN- γ stimulation (left, lower panel); same gating strategy is utilized for pSTAT1
879 staining of sorted splenic macrophages from PBS, Abx or B.C-colonized Abx mice and the
880 histogram overlay images of pSTAT1 staining at 16 h post IFN- γ stimulation are presented
881 (right, lower panel).

Figures

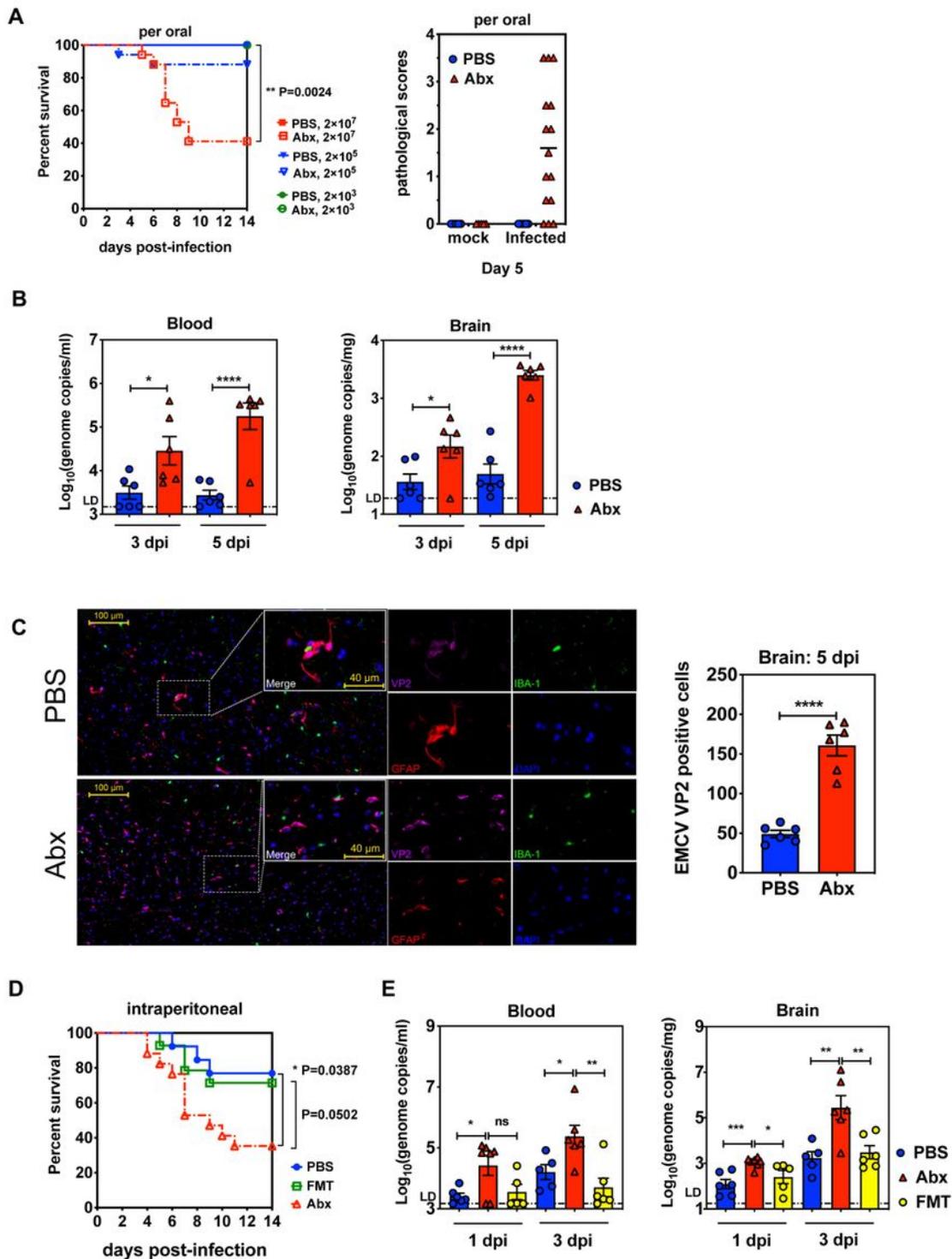


Figure 1

Microbiota depletion deteriorates EMCV pathogenesis and viral replication in vivo. (A) Groups of 6-8 weeks old C57BL/6J mice were pretreated with PBS or Abx for 5 days, inoculated with different dose of EMCV orally and observed for 14 days. Survival curves (n=12-17) and pathological scores (n=10-15) were

documented. (B) Blood and brain samples of PBS or Abx mice inoculated with 2×10^7 TCID₅₀ units of EMCV orally were harvested and tested for viral loads at 3 and 5 dpi by qPCR (n=6). (C) Indirect fluorescent assay (IFA) of brain sections from PBS- or Abx-infected mice at 5 dpi. EMCV capsid protein VP2 and cellular surface markers were double-stained with respective antibodies (representative images from n=6 per group). (D) Groups of PBS, Abx or Abx mice received FMT from conventionally housed PBS naïve mice were intraperitoneally inoculated with 200 TCID₅₀ units of EMCV for survival kinetics analysis (n=13-17). (E) Viral titers of brain tissues collected from PBS-, Abx- or FMT-infected mice at 1 and 3 dpi (same dose and route as Fig. 1D, n=5-6). Data represented in the figures were from two independent experiments. Dotted dash lines indicate the limit of detection (LD).

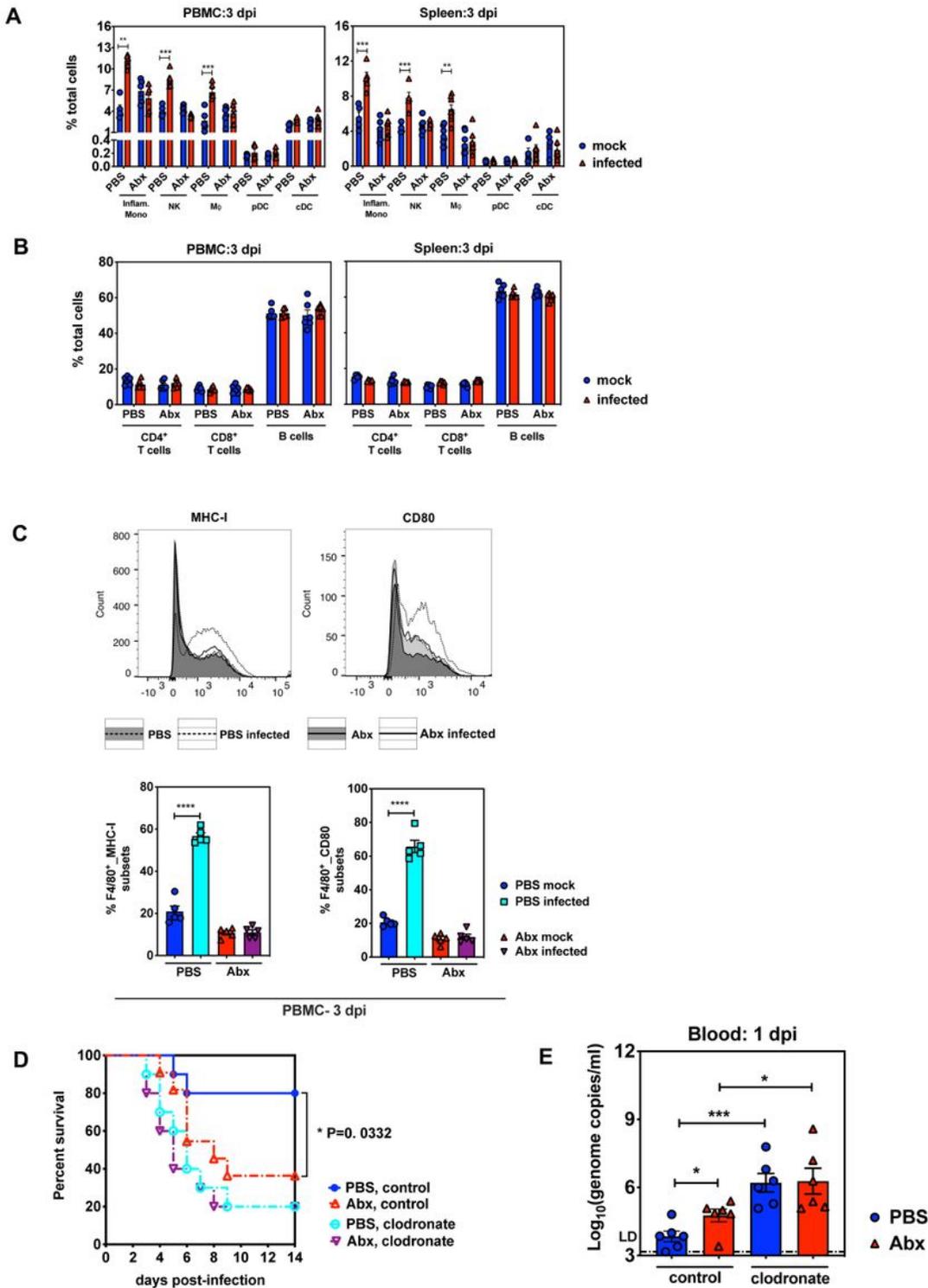


Figure 2

Innate cellular immune responses were severely diminished in Abx-treated mice post ECMV infection. (A) Frequency of a panel of innate immune cells including inflammatory monocytes, NK cells, macrophages, pDC or cDC in PBMC or splenocytes of mock or infected PBS or Abx mice at 3 dpi (n=4-8). (B) Frequency of various adaptive immune cells including CD4+ T cells, CD8+ T cells or B cells from PBMC or spleen of mock or infected PBS or Abx mice at 3 dpi (n=6). (C) Expression of MHC-I and CD80 on PBMC

macrophages from mock- or infected-PBS or Abx mice at 3 dpi (upper two panels, histograms of MHC-I or CD80-expressing macrophages; lower two panels, frequency of macrophages expressing MHC-I or CD80 in total PBMC macrophages; n=6). (D) Survival analysis of PBS or Abx mice treated with clodronate liposomes or control reagent following EMCV intraperitoneal inoculation (n=10-11). (E) Blood viral titers of PBS or Abx mice treated with clodronate liposomes or control reagent at 1 dpi (n=6). Data presented in the figure were from two independent experiments.

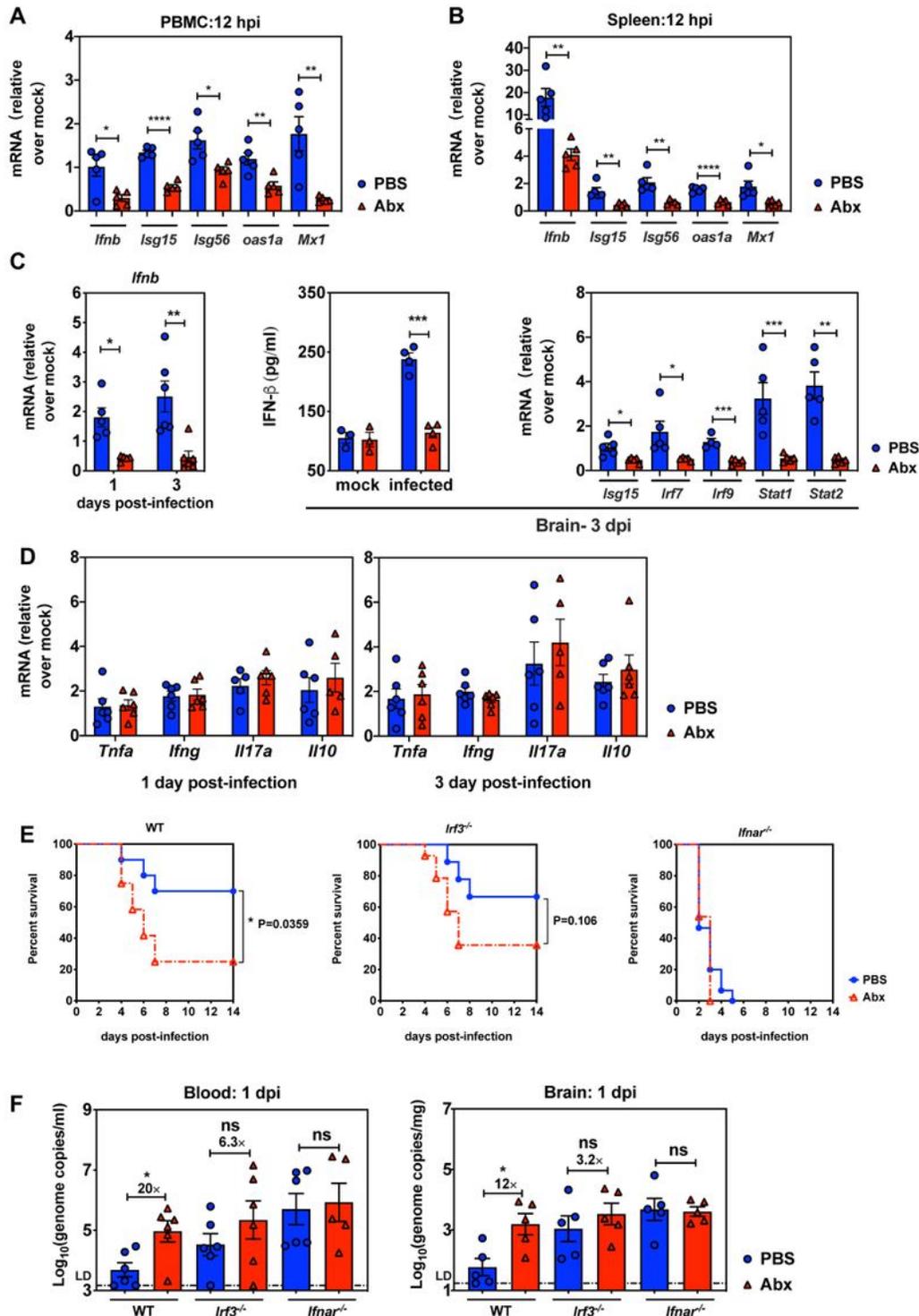


Figure 3

Microbiota depletion results in significantly impaired systemic or cerebral type I IFN responses. Fold induction of *Ifnb* and ISG expression in PBMC (A) and spleen (B) of PBS or Abx mice at 12 hpi following EMCV infection relative to respective mock controls (n=5). (C) Expression of *Ifnb* and ISGs in the brain of infected PBS- or Abx-treated mice (fold-change compared to respective mock mice) at 1 or 3 dpi (n=5-6). IFN- β levels in the brain at 3 dpi as detected by ELISA (middle panel). (D) Fold induction of proinflammatory or immune regulatory cytokines expression in the brain at 1 or 3 dpi relative to respective mock mice (n=6). (E) Survival kinetics of PBS- or Abx-treated *Irf3*^{-/-} or *Ifnar*^{-/-} infected with EMCV intraperitoneally, wild-type C57BL/6J mice were served as control (left panel, n=10-12; middle panel, n=9-14; right panel, n=13-15). (F) Viral titers in blood and brain collected from infected PBS or Abx WT, *Irf3*^{-/-} or *Ifnar*^{-/-} mice with at 1 dpi (n=5-6). All data presented in the figure were from two independent experiments.

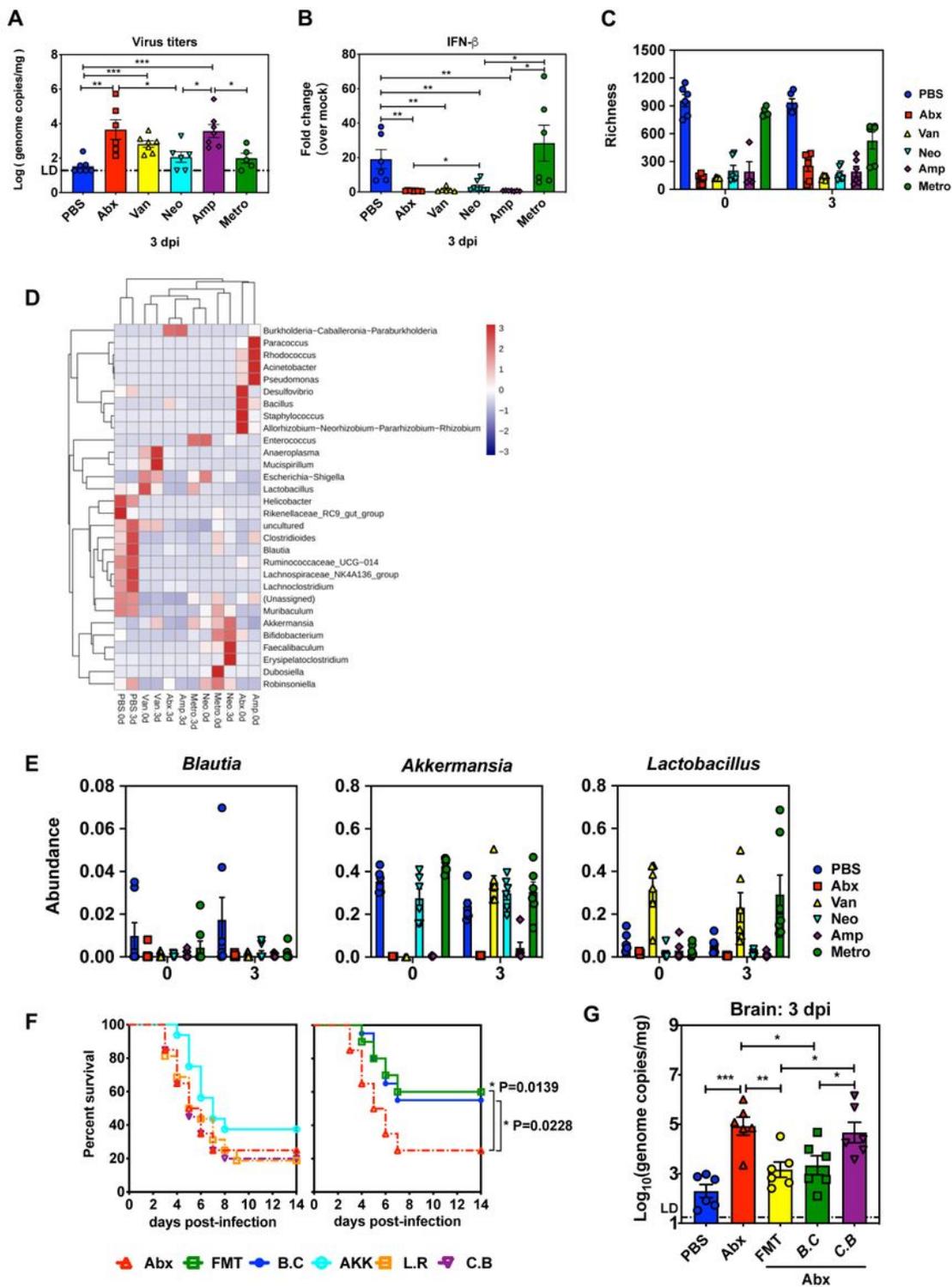


Figure 4

B. coccoides protects EMCV systemic infection by restricting viral in vivo replication. Viral titers (A) and induction of Ifnb (B) in the brain of PBS-, Abx- or single antibiotic-treated mice were measured by qPCR at 3 dpi (n=5-6). (C) Bacterial richness was defined by the number of the unique taxa. (D) Heat map of relative abundance from different groups of antibiotic treatment. (E) Relative abundance changes of bacteria at genus level. (F) Survival analysis of Abx mice colonized with *B. coccoides* (B.C), A. (AKK), L.R, and C.B.

muciniphila (AKK), *L. reuteri* (L.R), *C. butyricum* (C.B), or mouse fecal contents (FMT) following EMCV intraperitoneal inoculation (3 experiments, n=16-20). (G) Viral burden in brain tissues harvested from mock- or infected-PBS, Abx or Abx mice colonized with FMT, B.C or C.B at 3 dpi (n=6). All data presented in the figure were from at least two independent experiments.

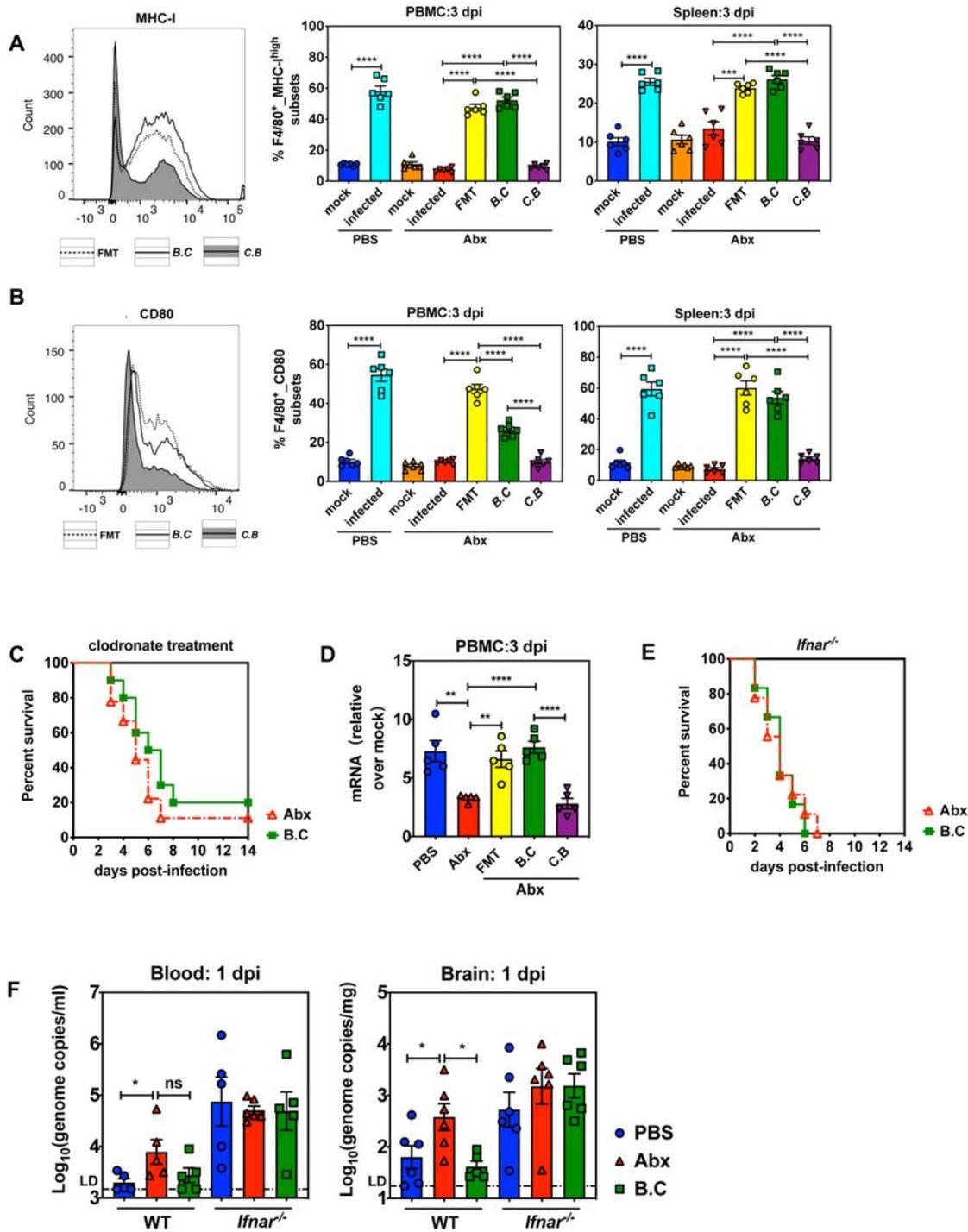


Figure 5

B. coccoides colonization restricts enteric virus systemic infection by activating innate cellular immune responses and type I IFN expression in a macrophage- and *Ifnar*-dependent manner. Frequency of macrophages expressing MHC-I (A) or CD80 (B) in PBMC or spleen isolated from PBS, Abx or Abx mice colonized with FMT, B.C or C.B at 3 dpi following EMCV inoculation (representative flow cytometry histograms of showing surface staining of MHC-I or CD80, n=6). (C) Survival kinetics of infected Abx and B.C-colonized Abx mice injected with clodronate liposomes (n=9-10). (D) Expression level of *Ifnb* in PBMC isolated from infected PBS, Abx, or Abx mice gavaged with FMT, B.C or C.B, comparing to their respective mock controls (n=5). (E) Survival kinetics of infected Abx and B.C-colonized Abx *Ifnar*^{-/-} mice (n=6-9). (F) Viral titers in blood and brain collected from infected PBS, Abx or B.C-colonized WT or *Ifnar*^{-/-} mice at 1 dpi (n=5-6). All data presented in the figure were from two independent experiments.

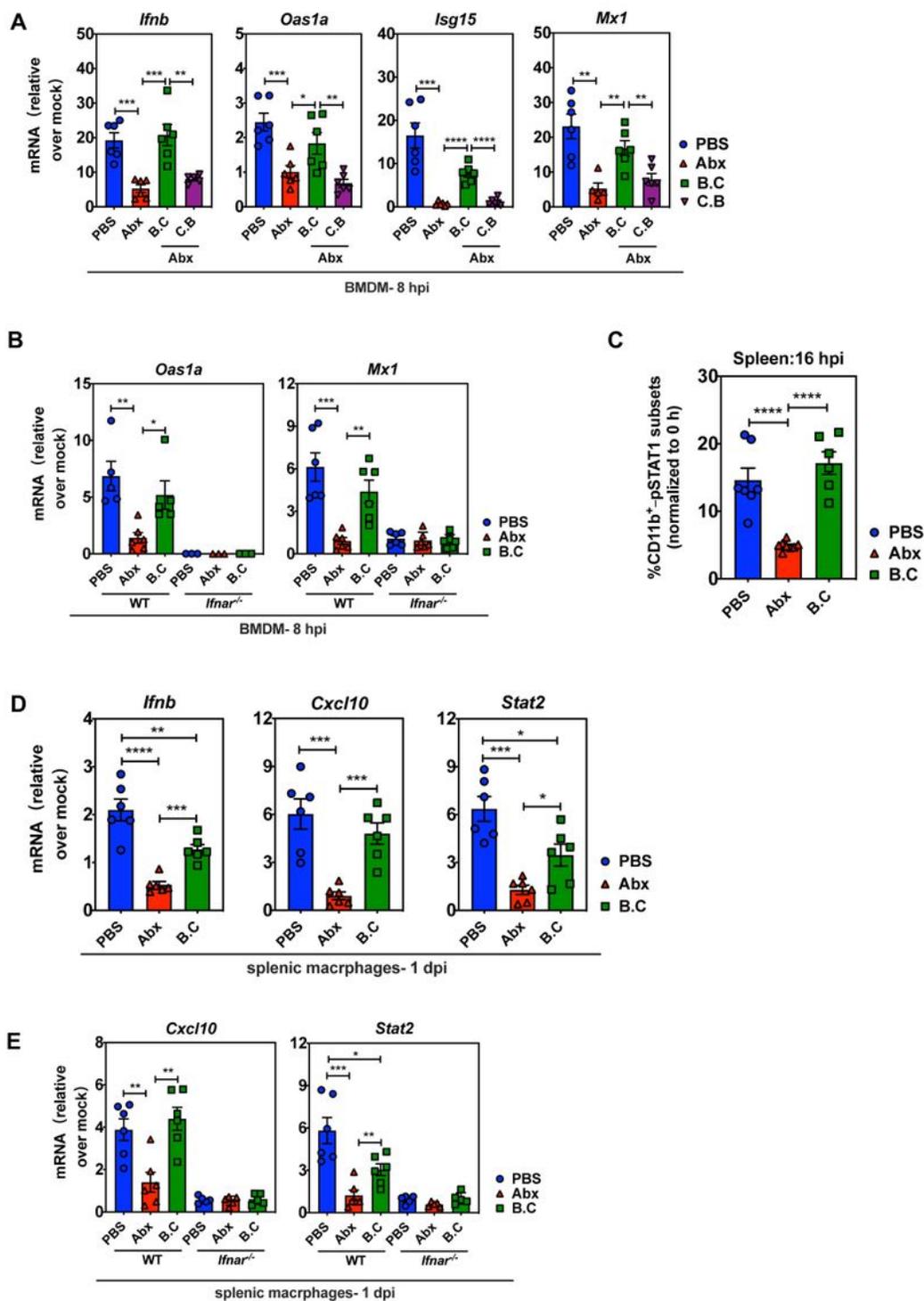


Figure 6

B. coccoides colonization promotes type I IFN and ISG responses in macrophages to limit EMCV infection. (A) *Ifnb* and antiviral defense gene expression in BMDMs isolated from PBS, Abx, B.C- or C.B- colonized Abx mice at 8 h post EMCV infection (MOI=5) in vitro (n=5-6). (B) Relative mRNA expression of *Oas1a* and *Mx1* in BMDMs isolated from WT or *Ifnar*^{-/-} mice that pretreated with PBS, Abx or Abx plus B. coccoides colonization at 8 hpi in vitro (n=5-6). (C) Splenic macrophages sorted from spleen of PBS, Abx

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