

1 **Culture under normoxic conditions and enhanced virulence of phase II *Coxiella***

2 ***burnetii* transformed with a RSF1010-based shuttle vector**

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18 **Running title:** Normoxic growth of *Coxiella burnetii*

19

20 **Abstract**

21 *Coxiella burnetii* is a Gram-negative, facultative intracellular microorganism that can
22 cause acute or chronic Q fever in human. It was recognized as an obligate intracellular
23 organism until the revolutionary design of an axenic cystine culture medium (ACCM).
24 Present axenic culture of *C. burnetii* strictly requires a hypoxic condition (<10%
25 oxygen). Here we investigated the normoxic growth of *C. burnetii* strains in ACCM-2
26 with or without tryptophan supplementation. Three *C. burnetii* strains - Henzerling
27 phase I, Nine Mile phase II and a Nine Mile phase II transformant, were included. The
28 transformant contains a pMMGK plasmid that is composed of a RSF1010 ori, a
29 repABC operon, an eGFP gene and a kanamycin resistance cassette. We found that,
30 under normoxia if starting from an appropriate concentration of fresh age inocula,
31 Nine Mile phase II can grow significantly in ACCM-2 with tryptophan, while the
32 transformant can grow robustly in ACCM-2 with or without tryptophan. In contrast,
33 long-term frozen stocks of phase II and its transformant, and Henzerling phase I of
34 different ages had no growth capability under normoxia under any circumstances.
35 Furthermore, frozen stocks of the transformant consistently caused large
36 splenomegaly in SCID mice, while wild type Nine Mile phase II induced a lesser
37 extent of splenomegaly. Taken together, our data show that normoxic cultivation of
38 phase II *C. burnetii* can be achieved under certain conditions. Our data suggests that
39 tryptophan and an unknown temperature sensitive signal are involved in the
40 expression of genes for normoxic growth regulated by quorum sensing in *C. burnetii*.

41 **Introduction**

42 *Coxiella burnetii* is a Gram-negative, facultative intracellular pathogen and the
43 causative agent of human Q fever. The primary transmission route for human is
44 through inhalation of contaminated aerosols from the secretions and excretions of
45 domestic ruminants (1). *C. burnetii* infections in humans may manifest as an acute
46 disease (mainly as a self-limiting febrile illness, pneumonia, or hepatitis) or as a
47 chronic disease (mainly endocarditis in patients with previous valvulopathy) (2). The
48 majority (~50-60%) of human infections are asymptomatic (2, 3). Chronic infections
49 are rare but can be fatal if untreated. *C. burnetii* is a significant cause of
50 culture-negative endocarditis in the United States (4). Treatment of chronic infections
51 is challenging and currently requires a combined antibiotic therapy with doxycycline
52 and hydroxychloroquine for at least 18 months (5). Worldwide only one vaccine for Q
53 fever called Q-Vax is licensed in Australia to protect high risk populations (6).

54 *C. burnetii* has two phase variants. Virulent *C. burnetii* isolated from natural
55 sources and infections is defined as phase I. It produces full-length LPS that may play
56 an important role in *C. burnetii* persistent infections by masking toll-like receptor
57 ligands from innate immune recognition by human dendritic cells (7). LPS from phase
58 I *C. burnetii* contains two unique biomarkers of methylated sugars (virenose and
59 dihydrohydroxystreptose) at its O-specific chain. When extensively passaged in
60 immunoincompetent hosts, virulent phase I *C. burnetii* mutates to avirulent phase II (8,
61 9). LPS from phase II *C. burnetii* is severely truncated and only contains lipid A and
62 partial core oligosaccharide. Lipid A, the basal component of LPS, is essential for *C.*

63 *burnetii* growth in macrophage-like THP-1 cells but nonessential in non-phagocytic
64 cells (10).

65 The recent description of an axenic culture medium and the subsequent modified
66 axenic culture media provide invaluable tools for *C. burnetii* research (11-15). The
67 first generation of axenic culture of *C. burnetii* requires a complex nutrient-rich
68 medium (Acidified Citrate Cysteine Medium, ACCM) and microaerophilic conditions
69 with 2.5% oxygen and 5% carbon dioxide (11). A second-generation medium
70 (ACCM-2) was generated by replacing fetal bovine serum with
71 methyl-beta-cyclodextrin (12), then was improved by supplementing tryptophan (14,
72 15). A nutritionally defined medium (ACCM-D) further improved *C. burnetii* growth
73 especially with increased bacterial viability (13). The continuous improvement of
74 axenic culture systems significantly facilitates the development of the *C. burnetii*
75 field.

76 Phase I *C. burnetii* is a category B select agent with potential for illegitimate use
77 and requires a biosafety level-3 laboratory for culture (16). Phase II *C. burnetii* is
78 approved for use at biosafety level-2 and is widely used for studying the biology and
79 pathogenesis in tissue culture. The caveat of working with phase II *C. burnetii* is the
80 lack of animal models for assessing *in vivo* pathogenesis. To promote the
81 identification of virulence factors in phase II *C. burnetii*, an SCID mouse model was
82 recently established (17). An alternative infection model using larvae of the greater
83 wax moth *Galleria mellonella* was also used to identify virulence factors in phase II *C.*
84 *burnetii* (18, 19).

85 In this study, we characterized the growth of various *C. burnetii* strains under
86 different oxygen concentrations in ACCM-2 with or without tryptophan
87 supplementation. We found that, under normoxia if starting from an appropriated
88 concentration of freshly age inocula, *C. burnetii* Nine Mile phase II had significant
89 propagation in ACCM-2 supplemented with tryptophan. Under same conditions, a
90 Nine phase II transformant that contains a RSF1010-based shuttle vector can have
91 robust growth in ACCM-2 with or without tryptophan. Phase I *C. burnetii* had no
92 apparent growth under normoxia in ACCM-2 with or without tryptophan. Long-term
93 frozen stocks of all tested strains unanimously failed to have normoxic growth. In the
94 SCID mouse infection model, compared to wild type Nine Mile phase II, frozen
95 stocks of its transformant caused larger splenomegaly. Our data suggest that *C.*
96 *burnetii* retains functional genes for aerobic growth, and the expression of these genes
97 might be regulated by a temperature sensitive signal of the quorum sensing system.
98 Our data also suggest that the expression of genes relating to aerobic growth is
99 associated with enhanced *in vivo* virulence in *C. burnetii*.

100 **Materials and Methods**

101 ***C. burnetii* strains.** *C. burnetii* Nine Mile phase II and Henzerling phase I are from
102 our laboratory strain collection. *C. burnetii* transformant -NMIIpMMGK was
103 constructed by transforming Nine Mile phase II with a shuttle vector pMMGK by
104 electroporation (20). The pMMGK plasmid is composed of a RSF1010 ori, a repABC
105 operon, an eGFP gene and a kanamycin resistance cassette. Construction of pMMGK
106 was described previously (10).

107 **Cultivation of *C. burnetii*.** Seed stocks of *C. burnetii* Nine Mile phase II, Henzerling
108 phase I and phase II transformant NMIIpMMGK were collected from ACCM-2
109 cultures at 37°C under 2.5% oxygen, and were stored in fresh ACCM-2 at -80°C.
110 Different ages and concentrations of inocula were used to characterize *C. burnetii*
111 growth in two media (ACCM-2 with or without tryptophan supplementation) under
112 hypoxia or normoxia. Inocula were added in 150 µL media per well in 96 well plates
113 (Corning). The plates were placed in a tri-gas CO₂ incubator (2.5% oxygen and 5%
114 CO₂) or a CO₂ incubator (5% CO₂) for seven days at 37°C. Each culture condition has
115 six replicates.

116 **Quantitative PCR.** Genome equivalent (GE) of *C. burnetii* was quantified by using
117 qPCR as previously described with minor modifications (21). Bacterial bodies of *C.*
118 *burnetii* were collected by centrifugation at 15,000 ×g for 30 min. Total DNA was
119 extracted with TIANamp N96 Blood DNA Kit (Tiangen). Genome copy numbers
120 were determined by Taqman probe qPCR specific to *dotA* by using a ViiA™ 7
121 Real-Time PCR System (Applied Biosystems). PCR conditions were as follows:
122 initial denaturation at 94°C for 10 min, followed by 40 cycles of amplification at 94°C
123 for 15 s, 60°C for 1 min.

124 **Mouse infection.** Female SCID (CB17/Icr-Prkdc^{scid}/IcrIcoCrI) mice were purchased
125 from Vital River Laboratory Animal Technology Co., Ltd (Beijing, China). Two
126 separate mouse infection experiments were conducted. Mice were infected with mock
127 or 1×10⁸ GE *C. burnetii* in 200 µL PBS by intra-peritoneal injection. In the first
128 experiment, 18 four weeks old mice were divided into three groups: control (PBS)

129 group and two infection groups. Fresh age (less than one day old) inocula of Nine
130 Mile phase II and NMIIpMMGK were used. In the second experiment, 9 five weeks
131 old mice were averaged into three infection groups: one month old Nine Mile phase II,
132 one-month old NMIIpMMGK and 19 months old NMIIpMMGK. At indicated days
133 post-infection, mice were weighted and sacrificed to harvest spleens to determine
134 splenomegaly (spleen weight/body weight). Each spleen was homogenized in 2 mL
135 PBS. Total DNAs from 20 μ L of each tissue homogenate were purified with DNeasy
136 Blood & Tissue Kit (Qiagen). *C. burnetii* GEs in spleens were determined by
137 quantitative PCR as described above. All animal procedures were carried out in strict
138 accordance with the guidelines for the Care and Use of Laboratory Animals of the
139 National Ministry of Health of China.

140 **Statistical Analysis.** A two-tailed Student *t* test was used for qPCR analysis of *C.*
141 *burnetii* growth yields under various conditions and was used for comparison of
142 splenomegaly and spleen bacterial loads of different infection groups.

143 **Results**

144 **Phase II *C. burnetii* transformants can grow robustly under normoxia in**
145 **ACCM-2 with or without tryptophan supplementation.** Current axenic culture of
146 *C. burnetii* requires a microaerophilic environment (2.5% oxygen) (14). But
147 interestingly, *C. burnetii* encodes terminal oxidases -cytochrome *o* (encoded by
148 *cyoABCDE*) and cytochrome *bd* (encoded by *cydAB*), which are typically associated
149 with aerobic and microaerophilic respiration, respectively (22). The presence of
150 cytochrome *o* suggests the possibility of *C. burnetii* replication under aerobic

151 conditions. In our attempts to culture *C. burnetii* in regular CO₂ incubators (~20%
152 oxygen), we found that a *C. burnetii* Nine Mile phase II transformant -NMIIpMMGK
153 can grow robustly (~3 logs increase) if starting from a reasonably high inoculum
154 concentration in the regular ACCM-2 medium (without tryptophan supplementation)
155 (Figure 1A). The pMMGK plasmid is composed of the pMMB207 backbone (a
156 RSF1010 *ori* and a repABC operon), an eGFP gene and a kanamycin resistance
157 cassette (10). This initial finding of *C. burnetii* normoxic growth was made in T-flask
158 cultures. To thoroughly characterize the influence of multiple culture parameters on *C.*
159 *burnetii* growth, 96 well plates were used for *C. burnetii* cultivation. Tryptophan
160 supplementation can improve the viability of *C. burnetii* in ACCM-2 (14). When
161 cultured in ACCM-2 with tryptophan supplementation, NMIIpMMGK displayed
162 similar growth characteristics, though there was a trend of a bigger yield in ACCM-2
163 with tryptophan supplementation (Figure 1B).

164 **Tryptophan improved axenic growth of phase II *C. burnetii* under normoxia.** The
165 axenic growth of wild type phase I and phase II *C. burnetii* in early generations of
166 ACCMs without tryptophan supplementation has been extensively investigated and
167 was reported to be microaerophilic (2.5% oxygen) (11, 12). We repeated the
168 cultivation of wild type phase I and phase II *C. burnetii* in ACCM-2 without
169 tryptophan at different oxygen concentrations (Figure 2AC). In ACCM-2 without
170 tryptophan supplementation, these two strains grew normally at 2.5% oxygen but had
171 no significant growth at 20% oxygen, which is consistent with previous reports. In
172 ACCM-2 with tryptophan at 2.5% oxygen, both strains had similar growth yields

173 (Figure 2BD). Interestingly, in ACCM-2 with tryptophan at 20% oxygen, wild type
174 phase II *C. burnetii* grew significantly (~2.5 logs increase) if starting from a high
175 concentration of fresh age inocula (~4×10⁶ GE/mL), while phase I *C. burnetii*
176 consistently had no obvious growth. These results show tryptophan has subtly
177 different effects on the axenic growth of wild type phase I and phase II strains.

178 **Long-term frozen stocks of *C. burnetii* strains failed to have apparent**
179 **propagation under normoxia.** The cultivation of NMIIpMMGK at 20% oxygen was
180 constantly repeated. Its viability at 2.5% oxygen in ACCM-2 with or without
181 tryptophan had no obvious variation after long-term frozen (>three months) in -80°C
182 (Figure 3AB). Interestingly, its growth ability at 20% oxygen in either medium was
183 completely abolished after long-term frozen. Similar with NMIIpMMGK, wild type
184 Nine Mile phase II also failed to replicate at 20% oxygen in ACCM-2 with tryptophan
185 (Figure 3CD). The growth deficiency of wild type Henzerling phase I at 20% oxygen
186 in either medium was consistent (Figure 3EF). Compared to growth of freshly
187 cultivated inocula (Figure 2), lower fold change of growth yield of long-term frozen
188 inocula was observed in either medium (Figure 3C-F), suggesting low viability of
189 both wild type frozen stocks. Taken together, our results suggest that a temperature
190 sensitive factor is associated with the expression of genes involved in *C. burnetii*
191 aerobic growth and viability.

192 **Phase II *C. burnetii* transformants show enhanced virulence in the SCID mouse**
193 **infection model.** The SCID mouse model allows for assessing *in vivo* pathogenesis of
194 phase II *C. burnetii* (17). Our *C. burnetii* phase II strains with or without the pMMGK

195 plasmid showed variable growth capability under normoxia. We next intended to
196 investigate whether pMMGK affects the virulence of phase II *C. burnetii* in the SCID
197 mouse model. Firstly, fresh (one day old) inocula of 1×10^8 genome equivalents (GE)
198 of NMII or NMIIpMMGK were used to infect four weeks old female SCID mice by
199 peritoneal (IP) injections. 16 days post infection similar magnitude of splenomegaly
200 was detected in both infected groups ($p=0.10$) (Figure 4A, C). Notably, two cases of
201 exceptional splenomegaly were observed in the NMIIpMMGK group. To examine the
202 potential influences of inoculum age on *in vivo* virulence, another infection
203 experiment using frozen bacterial stocks and five weeks old female SCID mice were
204 conducted. Compared to the results of using one day old inocula, 18 days post
205 infection one-month-old frozen stock of NMII caused similar size of splenomegaly
206 (Figure 4B, C). Unexpectedly, however, frozen stocks of both one month old and 19
207 months old of NMIIpMMGK induced similar large splenomegaly in all infected mice
208 (Figure 4B, C). The sizes of splenomegaly are in accordance with genome equivalents
209 in all infected mice (Figure 4D). Taken together, frozen stocks of phase II *C. burnetii*
210 transformants showed enhanced virulence in the SCID mouse model.

211 **Discussion**

212 The development of the axenic culture medium significantly advanced *C. burnetii*
213 research (11). The axenic culture medium for *C. burnetii* was continuously improved
214 (11-14). These medium studies found that hypoxic conditions are strictly required for
215 *C. burnetii* axenic replication. In this study, we characterized the axenic growth of
216 different *C. burnetii* strains in ACCM-2 with or without tryptophan supplementation,

217 especially their growth under normoxic conditions. Our goal was to further improve
218 the axenic culture of *C. burnetii* under normoxia.

219 We found that, under normoxic conditions if starting from appropriate
220 concentrations of fresh inoculum, *C. burnetii* transformants of pMMGK can have
221 robust growth in ACCM-2, regardless of tryptophan supplementation. Similarly, wild
222 type *C. burnetii* phase II can grow significantly in ACCM-2 with tryptophan
223 supplementation. Clearly, the *C. burnetii* transformants have a greater capability of
224 growing under normoxia. The mechanism of *C. burnetii* normoxic growth is unknown,
225 but likely its expression of *cyoABCDE* and *cydAB* is coordinately regulated in a
226 reciprocal fashion in response to oxygen concentration as in *E. coli* (23).

227 *C. burnetii* has been recognized as an intracellular parasite in nature. Inside the
228 infected cells, *C. burnetii* replicates in a hypoxic, phagolysosome-like vacuole (24).
229 The cytochrome *bd* oxidase, with its increased affinity for oxygen, is likely used by *C.*
230 *burnetii* under hypoxic conditions. The alternative terminal ubiquinol oxidase in *C.*
231 *burnetii* -cytochrome *o* (typically used for aerobic respiration), correlates with our
232 finding of normoxic growth. Intriguingly, the normoxic growth is affected by multiple
233 factors including the shuttle vector pMMGK, the age and concentration of inoculum,
234 and the phase of *C. burnetii*. The differential normoxic growth of different strains
235 suggests that, in *C. burnetii* phase II, a temperature-sensitive quorum sensing signal
236 molecule like RNA might be involved in the regulated expression of genes for aerobic
237 respiration, especially cytochrome *o*.

238 The capability of normoxic growth in ACCM-2 without tryptophan

239 supplementation is restricted to *C. burnetii* transformants. Their enhanced virulence is
240 likely associated with normoxic growth. How might the shuttle vector pMMGK
241 confers these “gain-of-function” phenotype to *C. burnetii*? The shuttle vector
242 pMMGK brings new proteins (RepA, RepB, RepC, eGFP and Kan^R) and new RNAs
243 to *C. burnetii*. The potential effects of these proteins and RNAs on bacterial
244 physiology are impossible to predict (25, 26). The RSF1010 *ori*-based shuttle vectors
245 are commonly used for *C. burnetii* transformation (27, 28). Whether pMMGK or
246 other similar vectors can restore normoxic growth of other *C. burnetii* phase I and
247 phase II strains will be worthy of further investigation.

248 The normoxic cultures in this study were performed under static conditions. Given
249 that oxygen supply in static cultures is much less than in shaken cultures, it is
250 necessary to test *C. burnetii* propagation in normoxic shaken cultures. This will help
251 determine the influence of inoculum concentration on *C. burnetii* growth under
252 normoxia, as one may argue that high concentrations of inoculum might reduce the
253 physical concentration of oxygen in media. Despite this possibility, static cultures of
254 *C. burnetii* under normoxic and hypoxic conditions have distinct outcomes in infected
255 macrophages (29). *C. burnetii* replication in macrophages is prevented in static
256 hypoxic cultures, but not in static normoxic cultures (29).

257 In conclusion, we characterized the growth of various *C. burnetii* strains in
258 ACCM-2 with or without tryptophan under normoxic and hypoxic conditions.
259 Consistent with previous reports, *C. burnetii* had robust growth under hypoxia. What
260 is surprising is that we found a *C. burnetii* phase II transformant and the wild type *C.*

261 *burnetii* phase II can have significant to robust growth under certain normoxic
262 circumstances. Compared to the wild type strain, the transformant displayed enhanced
263 virulence in the SCID mouse model. Normoxic growth of *C. burnetii* is affected by
264 the shuttle vector pMMGK, the age and concentration of inoculum, and the phase of
265 *C. burnetii*. Our findings suggest that current axenic culture of *C. burnetii* under
266 hypoxia might possibly be further modified to allow routine axenic culture under
267 normoxia.

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357 **Figure legends**

358 **Figure 1.** Growth of *C. burnetii* NMIIpMMGK in ACCM-2 (A) and ACCM-2 with
359 tryptophan supplementation (B) at 2.5% and 20% oxygen. At 2.5% oxygen, regardless
360 of its inoculum concentration, the NMIIpMMGK strain consistently has robust growth
361 in ACCM-2 and ACCM-2 plus tryptophan. At 20% oxygen, in both ACCM-2 and
362 ACCM-2 plus tryptophan, the growth yield of NMIIpMMGK ranges from 0-3.6 logs,
363 depending on its inoculum concentration.

364 **Figure 2.** Growth of wild type *C. burnetii* phase I and phase II strains in ACCM-2 and
365 ACCM-2+Trp at 2.5% and 20% oxygen. In ACCM-2, at 20% oxygen, the wild type
366 NM II and Henzerling I strains have no significant growth. In ACCM-2 plus
367 tryptophan, at 20% oxygen, the growth of NM II ranges from 0-2.5 logs, depending
368 on its inoculum concentration, while the Henzerling I strain consistently fails to grow.

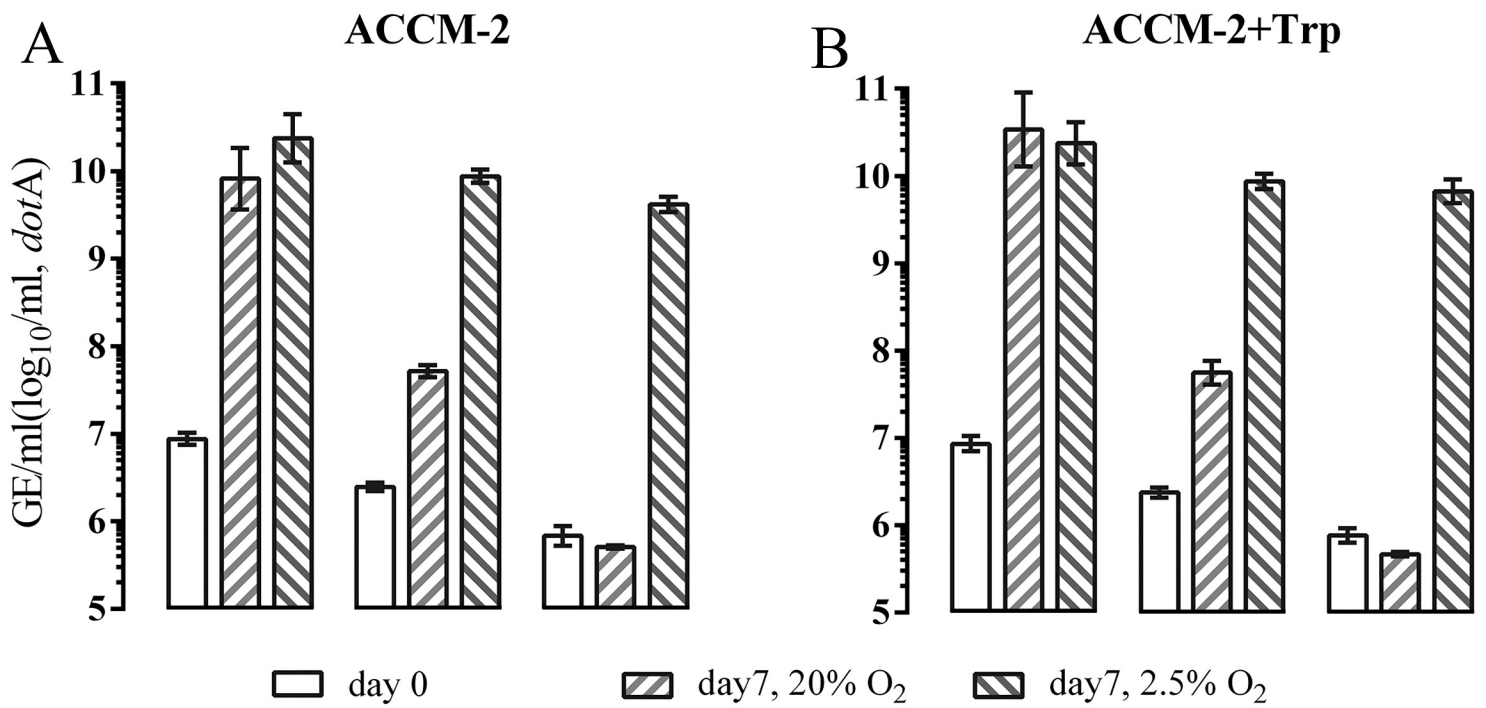
369 **Figure 3.** Growth of frozen stocks of *C. burnetii* strains -Henzerling phase I and Nine
370 Mile phase II and NMIIpMMGK in ACCM-2 with (A, C, E) or without tryptophan (B,

371 **D, F)** at 2.5% and 20% oxygen. In both ACCM-2 and ACCM-2 plus tryptophan,
372 frozen stocks of three *C. burnetii* strains have significant growth at 2.5% oxygen.
373 However, frozen stocks of all strains fail to grow at 20% oxygen, regardless of the
374 media and their inoculum concentrations.

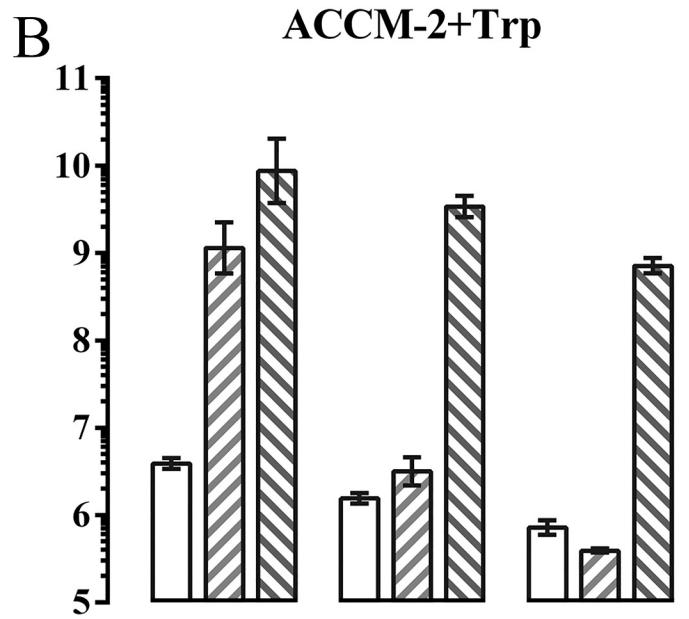
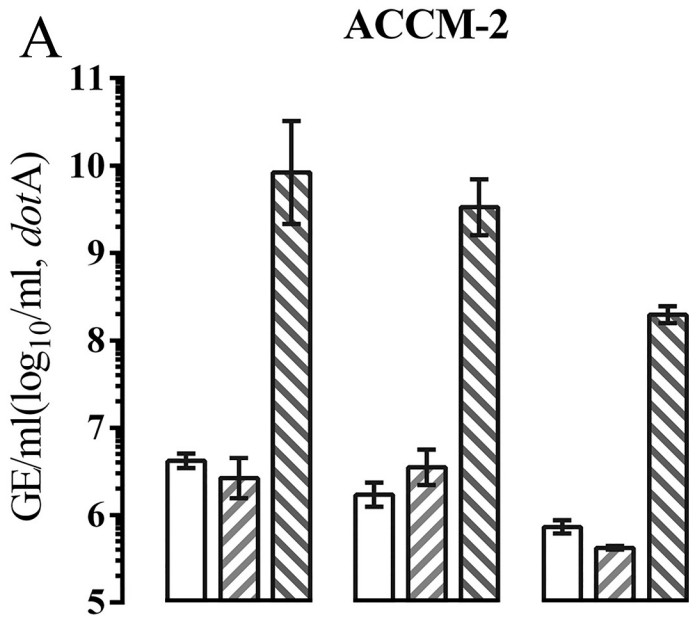
375 **Figure 4.** Intra-peritoneal challenge of SCID mice with different inoculum ages of *C.*
376 *burnetii* strains -Nine Mile phase II and NMIIpMMGK. **(A, B)** Spleens were removed
377 from challenged and control mice. **(C)** Splenomegaly calculated as spleen weight as a
378 percentage of total body weight at the time of necropsy after infection. **(D)** Genome
379 equivalents calculated using Taqman real-time PCR with DNA purified from infected
380 lungs. Error bars represent standard deviation from the mean.

381

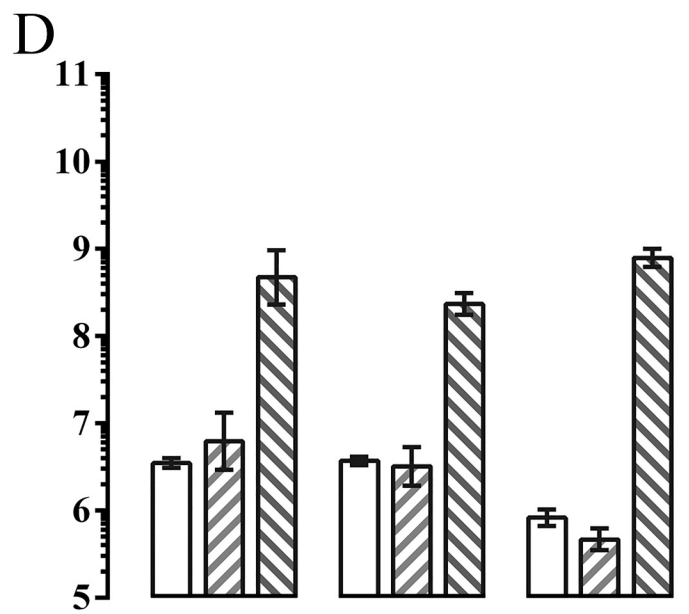
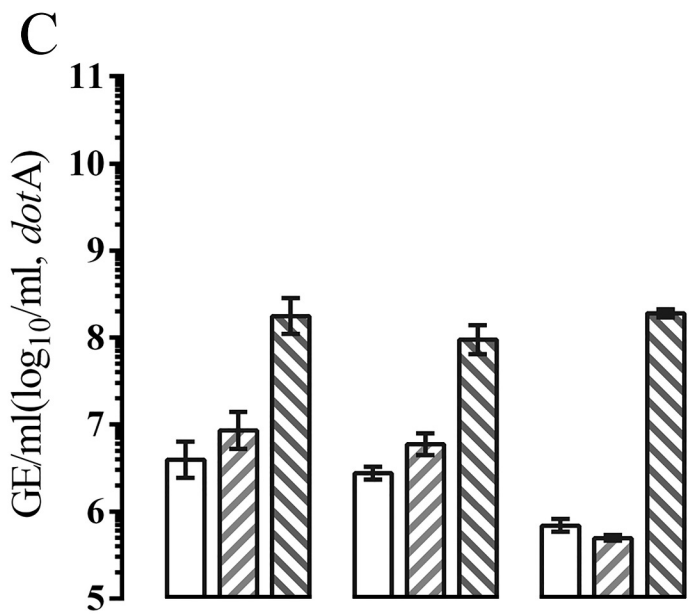
382



NM II



Henzerling I

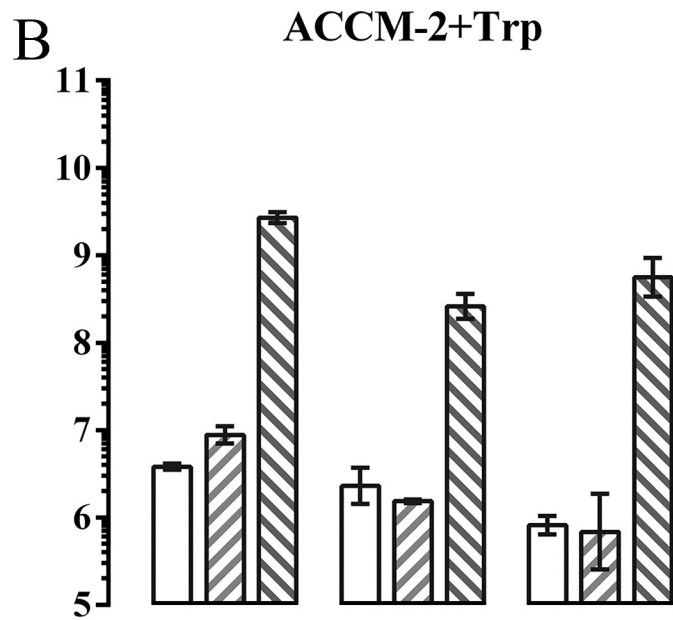
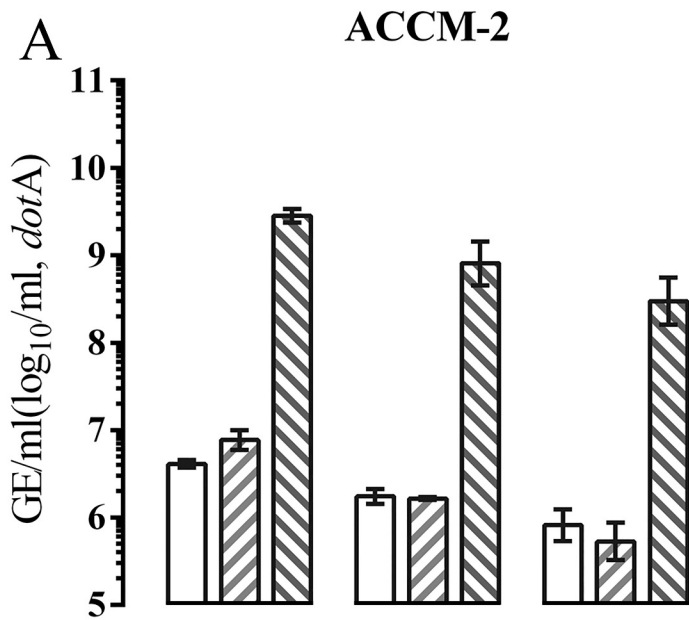


□ day 0

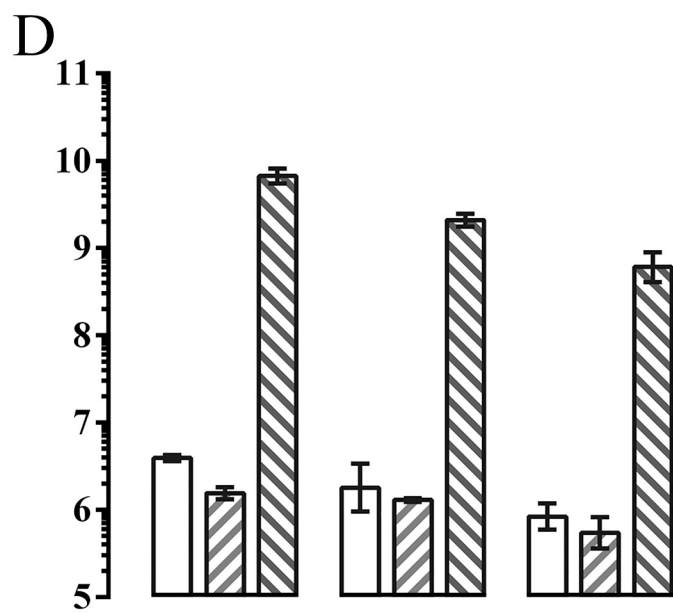
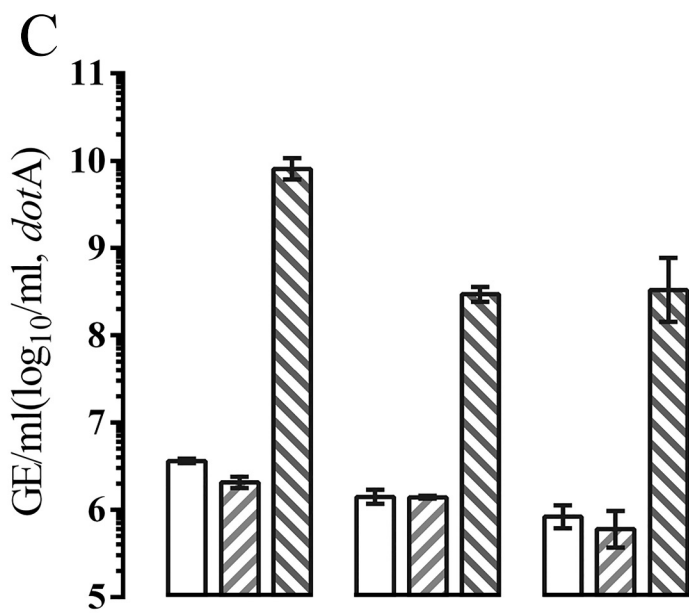
▨ day7, 20% O₂

▩ day7, 2.5% O₂

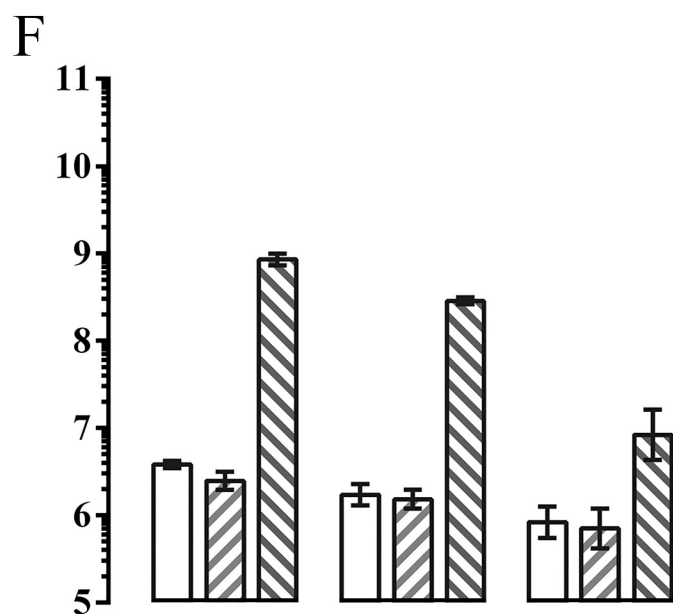
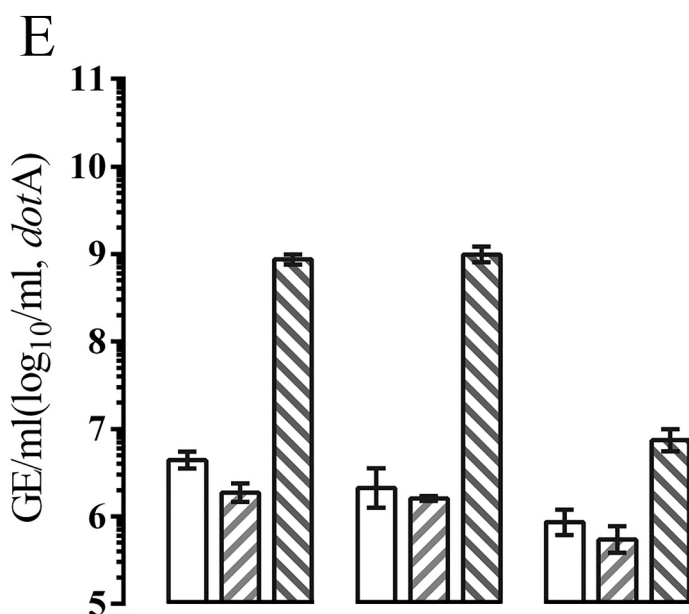
NMIIpMMGK



NM II



Henzerling I

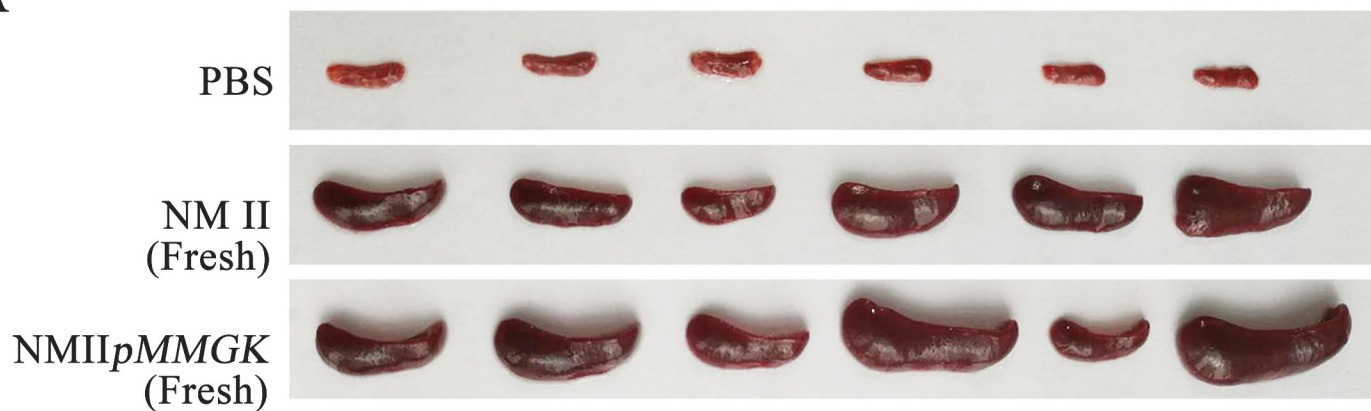


□ day 0

▨ day 7, 20% O₂

▩ day 7, 2.5% O₂

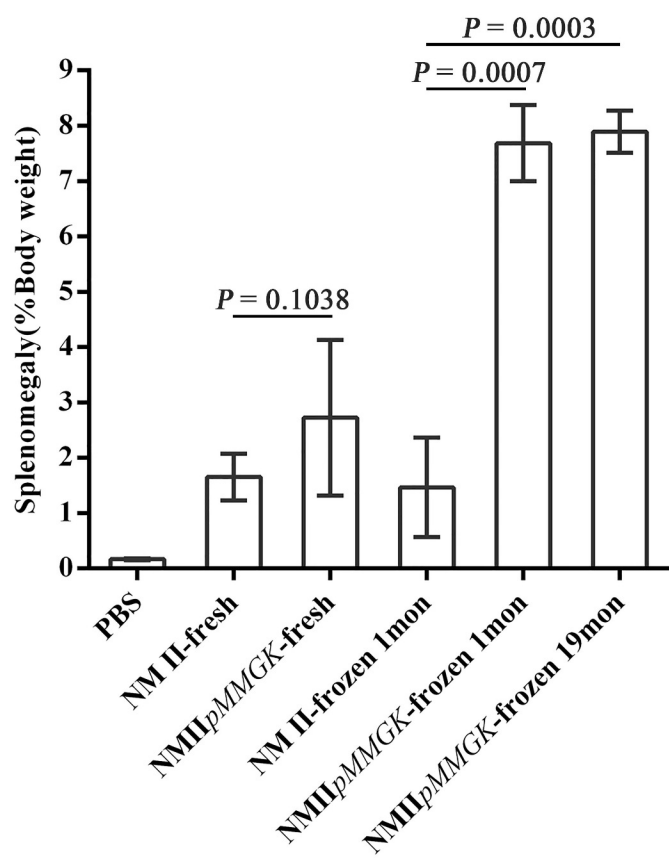
A



B



C



D

