

1 **The protected physiological status of intracellular *Salmonella enterica***
2 **persisters reduces host cell-imposed stress**

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22 **Abstract**

23 Today, we are faced with increasingly occurring bacterial infections that are hard to treat and
24 often tend to relapse. These recurrent infections can occur possibly due to antibiotic-tolerant
25 persister cells. Antibiotic persistent bacteria represent a small part of a bacterial population that
26 enters a non-replicating (NR) state arising from phenotypic switching. Intracellularly, after
27 uptake by phagocytic cells, *Salmonella enterica* serovar Typhimurium (STM) forms persister
28 cells that are able to subvert immune defenses of the host. However, the clear physiological
29 state and perceptual properties are still poorly understood and many questions remain
30 unanswered. Here we describe further development of fluorescent protein-based reporter
31 plasmids that were used to detect intracellular NR persister cells and monitor the expression of
32 stress response genes via extensive flow cytometric analyses. Moreover, we performed
33 extensive measurements of the metabolic properties of NR STM at the early course of infection.
34 Our studies demonstrate that NR STM persister cells perceive their environment and are
35 capable respond to stress factors. Since persisters showed a lower stress response compared to
36 replicating (R) STM, which was not a consequence of a lower metabolic capacity, the persistent
37 status of STM serves as protective niche. Furthermore, up to 95% of NR STM were
38 metabolically active at the beginning of infection additionally showing no difference in the
39 metabolic capacity compared to R STM. The accessory capability of NR STM persisters to
40 sense and to react to stress with constant metabolic activity may supports the pathogen to create
41 a more permissive environment for recurrent infections.

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43 **Introduction**

44 The global increase of multi-resistant bacteria presents one of the major challenges to human
45 health in the near future. In addition, physicians frequently encounter bacterial infections that
46 are very difficult to treat, and often relapse without the presence of genetic resistance to
47 antibiotics (Blango and Mulvey, 2010; Caygill et al., 1994; Levine et al., 1982; Lin and Flynn,
48 2010; Mulvey et al., 2001). These recurrent infections can only be defeated by several rounds
49 of antibiotic treatment, possibly due to the presence of antibiotic-tolerant persister cells. In the
50 context of recurrent infections, the heterogeneous phenomenon of bacterial antibiotic
51 persistence is becoming increasingly important. Antibiotic persistence describes a phenomenon
52 in which a small part of a bacterial population enters a non-replicating (NR) state that can
53 survive actually lethal concentrations of antibiotics during infection. Persister cells arise from
54 a genetically clonal bacterial population by a transient and reversible phenotype switch, leading
55 to a NR and multidrug-tolerant subpopulation (Bigger, 1944; reviewed in Harms et al., 2016;
56 Keren et al., 2004).

57 Recent *in vitro* investigations of antibiotic persistence observed bacteria entering a dormant
58 state when grown in laboratory media (Conlon et al., 2016; Maisonneuve and Gerdes, 2014;
59 Shan et al., 2017). However, the physiological state and the perceptual properties of
60 intracellular persistent bacteria are still poorly understood. Many questions regarding the
61 interface of antibiotic persisters with their host remain to be answered, and there is demand for
62 sensitive tools to interrogate the interplay of both organisms. Better understanding of the
63 physiology of persisters will help to device new forms of antimicrobial therapy.

64 The facultative intracellular pathogen *Salmonella enterica* causes acute and chronic infections
65 (Gunn et al., 2014), and *S. enterica* serovar Typhimurium (STM) serves as a model organism
66 for a facultative intracellular pathogen forming persister cells during systemic infections. It was
67 shown that intracellular persister cells of STM do not show complete dormancy, but consist of
68 subpopulations of metabolically active bacteria, as well as inactive cells showing decreasing

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69 responsiveness to external stimuli over the course of infection (Helaine et al., 2010). Similar
70 NR but metabolically active bacteria have also been observed in macrophages infected with
71 *Mycobacterium tuberculosis* (Manina et al., 2015). It was shown that many persisters of STM
72 are formed immediately upon phagocytosis by macrophages. Vacuolar acidification as well as
73 nutritional deprivation was mentioned as one of the main factors leading to macrophage-
74 induced persister formation (Helaine et al., 2014). More recently, Stapels et al. (2018) reported
75 that persisters of STM translocate SPI2-T3SS effector proteins to dampen proinflammatory
76 innate immune response, and induce anti-inflammatory macrophage polarization. Such
77 reprogramming of their host cells allows NR STM to survive, and might lead to an advantage
78 during infection relapse after termination of antibiosis (Stapels et al., 2018).
79 Within phagocytic cells, STM encounters harsh environmental conditions and various defense
80 mechanisms including antimicrobial peptides and the respiratory burst (Kagaya et al., 1989;
81 Slauch, 2011; Zhang and Gallo, 2016). For the pathogen it is of crucial importance to sense and
82 react to potentially detrimental factors. Hence, STM has evolved a plethora of defensive
83 virulence mechanisms to withstand antimicrobial effectors and to overcome the clearance by
84 the host (reviewed in Fang et al., 2016). These stress response systems (SRS) are able to sense
85 harmful conditions, as well as perturbations of the bacterial envelope, in periplasm, or in
86 cytoplasm. SRS have to perform efficiently in space and time for successful survival of STM
87 within hazardous host environments.
88 Despite being equipped with an extensive set of defensive and offensive virulence factors, the
89 individual fate of intracellular STM is highly diverse (Helaine and Holden, 2013). It was shown
90 that stress response and formation of persister cells are closely related. Among other, main
91 mediators for persister formation are considered to be SOS response via RecA and LexA,
92 stringent response via (p)ppGpp, oxidative stress response via OxyR and SoxSR, and toxin-
93 antitoxin modules (reviewed in Gollan et al., 2019; Harms et al., 2016; Prax and Bertram, 2014).
94 But what happens after establishment of the persistent status? Is switch from persister state to

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95 normal growth a merely stochastic event, or can persister monitor their environment with the
96 ability to respond to cues? Addressing these questions is challenging due to population
97 heterogeneity of intracellular STM, the low frequency of persisters, and their low metabolic
98 activity.

99 Perception of environmental stimuli and especially stressors are of crucial importance for
100 pathogens to maintain their cell integrity. So far, little is known about the stress response of
101 intracellular NR STM. We investigated whether NR STM after entering persistence are still
102 capable in sensing stress factors and in responding by inducing stress responses. If so, is the
103 level of stress response of NR STM similar to replicating (R) STM, or altered? To address these
104 questions, we developed further the recently introduced reporter system (Schulte et al., 2020)
105 to enable analyses of stress response of intracellular STM persister cells. Here we introduce
106 dual fluorescence reporters which enable extremely sensitive flow cytometric analyses of both,
107 stress response and metabolic properties of intracellular STM persisters. Our analyses reveal
108 that intracellular NR STM persister cells perceive their environment and respond to stressors.
109 NR and R STM can mount responses to stressors, and the low response observed for NR STM
110 indicates that the persistent status of intracellular STM serves as protective niche.

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112 Results

113 *Design of fluorescence protein-based reporters for stress response of non-replicating* 114 *intracellular* *Salmonella enterica* *at single cell level*

115 In this study, we analyzed the stress response of NR, persisting intracellular STM at the single
116 cell level. For this purpose, we changed the basic design of dual fluorescence stress reporters
117 (Schulte et al., 2020) as shown in **Fig. 1**. The constitutive EM7 promoter was replaced by the
118 tet-ON cassette (Schulte et al., 2019) to induce expression of a gene of interest (GOI) by
119 addition of the non-antibiotic inducer anhydrotetracycline (AHT). The resulting dual
120 fluorescence reporters consisted of the tet-ON cassette for controlled expression of DsRed
121 version DsRed T3_S4T (Sorensen et al., 2003), and sfGFP under control of the regulated
122 promoters of SRS genes *msrA*, *trxA*, or *htrA* (**Fig. 1A**).

123 The principle for the detection of persisters was adapted from Helaine et al. (2010). Addition
124 of AHT to subcultures used as infection inoculum resulted in synthesis of DsRed. Before
125 infection, AHT was removed by centrifugation and washing to terminate further DsRed
126 synthesis. The principle of fluorescence protein (FP) dilution is that intracellular R STM
127 continuously dilute DsRed, thus, decrease fluorescence levels, while NR or persistent STM
128 maintain DsRed and consistent fluorescence levels. Furthermore, the sfGFP signal reported
129 exposure to stressors and induction of SRS (**Fig. 1B**).

130 To confirm function of newly generated reporter plasmids, we used *in vitro* cultures of STM
131 WT harboring reporter [$P_{tetA}::dsred$ $P_{msrA}::sfgfp$]. STM was grown in LB broth and samples
132 were collected in hourly intervals for quantification of DsRed fluorescence levels by flow
133 cytometry (FC) (**Fig. S 1AB**). Without further induction, DsRed intensity decreased over time
134 proportional to bacterial replication, validating the principle of fluorescence dilution upon
135 bacterial replication.

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136 For infection of the murine macrophage-like cell line RAW264.7, STM WT harboring the
137 reporter [$P_{tetA}::dsred$ $P_{msrA}::sfgfp$] was grown overnight (o/n) in presence of AHT to induce
138 expression of DsRed. RAW264.7 cells were infected by DsRed-positive STM WT. At 8 h p.i.
139 in culture without AHT, the population was released, subjected to FC and the x-median RFI of
140 DsRed was determined (**Fig. S 1C**). Two different subpopulations were detected when using
141 STM WT [$P_{tetA}::dsred$ $P_{msrA}::sfgfp$] (blue histogram). One subpopulation showed normal, non-
142 diluted DsRed intensities comparable to constitutive DsRed expression via the EM7 promoter
143 (**Fig. S 1C**, black histogram). The second subpopulation showed a lower DsRed intensity due
144 to FP dilution. Without addition of AHT to o/n cultures, no DsRed-positive bacteria were
145 detected (**Fig. S 1C**, red histogram). Furthermore, NR subpopulations were detected 16 h, 24 h
146 and 48 h p.i. using STM WT (**Fig. S 1D**), or various STM mutant strains (data not shown). We
147 also determined the detection accuracy of the cytometer used (**Fig. S 1E-H**). The measurements
148 demonstrated that up to 10,000-fold differences in numbers of STM expressing DsRed or sfGFP
149 were determined very precisely.

150 FC analysis of the distribution of DsRed and sfGFP fluorescence of the intracellular population
151 revealed a very small DsRed-positive NR subpopulation (app. 0.4%), compared to the DsRed-
152 negative R subpopulation (**Fig. S 2A**). Calculating the proportion of NR STM of various mutant
153 strains demonstrated that $\Delta ssaV$ and $\Delta dksA$ showed a higher proportion of NR STM of about
154 7% and 20%, respectively (**Fig. S 2B**), as result of attenuated intracellular replication of R STM.
155 Taken together, analyses of FP dilution allowed discrimination of R and NR STM, and detection
156 of NR STM of mutant strains at various time points p.i.

157 *Stress response of non-replicating intracellular Salmonella enterica*

158 Next, we analyzed the stress response in NR compared to R STM. For the detection of the entire
159 intracellular population, STM WT harboring [$P_{EM7}::dsred$ $P_{xxx}::sfgfp$] for constitutive DsRed
160 expression was used. For detection of intracellular NR STM, STM WT harboring [$P_{tetA}::dsred$
161 $P_{xxx}::sfgfp$] for AHT-induced DsRed expression was used. STM WT harboring the various

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162 reporters were used to infect RAW264.7 macrophages (**Fig. 2**). For STM WT harboring the
163 reporters for detection of NR bacteria, AHT was added to o/n cultures of inoculum. The bacteria
164 were released from host cells 8 h p.i. and subjected to FC (**Fig. 2AB**), or immuno-stained against
165 O-antigen and imaged by fluorescence microscopy (**Fig. 2C-H**). STM WT harboring a plasmid
166 for constitutive expression of DsRed, but lacking expression of sfGFP served as negative
167 control for adjustment of FC gating as described before (Schulte et al., 2020).

168 For all stress reporters investigated, i.e $P_{msrA}::sfgfp$, $P_{trxA}::sfgfp$, $P_{htrA}::sfgfp$, we observed
169 increased sfGFP levels in NR STM (**Fig. 2AB**). In contrast, the overall sfGFP expression of the
170 entire bacterial population was higher (2.72-, 1.79-, and 1.77-fold induction of $P_{msrA}::sfgfp$,
171 $P_{trxA}::sfGFP$, and $P_{htrA}::sfGFP$, respectively, compared to NR STM). Fluorescence microscopy
172 showed comparable results (**Fig. 2C-H**), and R and NR STM were readily distinguished by the
173 absence or presence of DsRed fluorescence, respectively (**Fig. 2C-E**). Correlation to sfGFP
174 fluorescence signals showed lower induction of *msrA*, *trxA*, or *htrA* in NR STM WT harboring
175 the stress reporters with constitutive DsRed expression all displayed DsRed and sfGFP
176 fluorescence, indicating induction of *msrA*, *trxA* or *htrA* (**Fig. 2F-H**). The inoculum without
177 AHT addition lacked DsRed synthesis, and accordingly DsRed fluorescence was not detectable
178 (**Fig. S 3**). NR STM WT resided in LAMP1-positive compartments throughout intracellular
179 presence (**Fig. S 4**), in line with prior findings (Helaine et al., 2010) and proving precision of
180 our dual fluorescence reporter system.

181 We have reported that specific virulence factors of STM contribute to reduce exposure to host
182 defense mechanisms and stressors (Noster et al., 2019; Schulte et al., 2020). Therefore, we
183 compared the stress response of R STM to NR STM strains deficient in SPI2-T3SS or SRS in
184 RAW264.7 macrophages at various time points p.i. (**Fig. 3**). We used P_{msrA} as representative
185 for induction of SRS. For all mutant strains analyzed, we observed a lower *msrA* induction of
186 the NR subpopulation compared to the entire population at 24 h p.i. (**Fig. 3A**). We further
187 calculated the time-resolved stress response of R and NR STM (**Fig. 3B**). The depicted slopes

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188 represent the sfGFP intensity increase over the time of infection (8 – 24 h p.i.). Higher stress
189 induction over a constant period of time resulted in a higher slope as shown for example for the
190 entire population of STM Δ *ssaV* and Δ *dksA* strains (red and yellow line in **Fig. 3B**). We
191 observed that stress induction in NR STM only increases very slightly during the course of
192 infection (dashed lines). In contrast, R STM showed a high increase of the sfGFP signal.

193 We can conclude that intracellular NR STM perceive external stress conditions and did not
194 encounter the same stress levels as R STM.

195 *Metabolic activity analyses of intracellular persisters at the early phase of infection*

196 The lower induction of stress reporters in NR STM can be explained by i) lower levels of
197 stressor acting on this population compared to R STM, or ii) reduced biosynthetic activity
198 resulting in lower synthesis of sfGFP. To distinguish, we analyzed the metabolic activity of NR
199 STM WT. For this, dual fluorescence reporter plasmid harboring arabinose-inducible *dsred*
200 expression and AHT-inducible *sfgfp* expression was generated (**Fig. S 5A**). Presence of
201 arabinose during culture of inoculum resulting in synthesis of DsRed. After removal of
202 arabinose and infection of RAW264.7 macrophages, R STM dilute DsRed while NR STM
203 maintain DsRed levels. sfGFP expression was induced by addition of AHT to infected cells,
204 serving as proxy for the biosynthetic capacity (metabolic activity) of STM (**Fig. S 5BC**). For
205 functional control, STM WT harboring the double-inducible dual fluorescence reporter
206 [$P_{BAD}::dsred$ $P_{tetA}::sfgfp$] was grown in LB medium and samples were collected for
207 quantification of DsRed and sfGFP levels by FC (**Fig. S 5D-F**). Arabinose was added at the
208 beginning of subculture (-4 h). After 4 h, the x-median RFI of DsRed increased, however, the
209 sfGFP intensity remained constantly low. After removal of arabinose and start of a further
210 subculture without the presence of arabinose but in the presence of AHT (0 h), the intensity of
211 DsRed decreased within 3 h of subculture due to fluorescence dilution. In contrast, the x-median
212 RFI of sfGFP increased (+3 h), confirming induction by AHT.

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213 Next, we analyzed the metabolic activity of NR STM WT [$P_{BAD}::dsred$ $P_{tetA}::sfGFP$] in
214 RAW264.7 macrophages at 24 h p.i. (Fig. 4) following the experimental design depicted in Fig.
215 S 5C. At 22 h p.i., AHT was added directly to the cell culture medium to induce expression of
216 sfGFP. At 24 h p.i., bacteria were released from host cells and subjected to FC as described
217 above. Plotting the population against their DsRed and sfGFP intensities allowed clear
218 discrimination of the various subpopulations (Fig. 4A). STM positive only for DsRed represent
219 metabolically inactive NR persisters. STM positive both for DsRed and sfGFP represent
220 metabolically active NR persisters. STM only sfGFP-positive represent a metabolically active
221 R population. Particles both negative for DsRed and sfGFP represent the background signal
222 containing host cell debris. Without the addition of arabinose to o/n cultures, and without
223 addition of AHT to infected cells, neither DsRed- nor sfGFP-positive STM could be detected
224 by FC (Fig. 4A). Comparing the amount of metabolically active to inactive NR STM revealed
225 that app. 30-40% of all persisters showed metabolic activity at 24 h p.i. (Fig. 4B). Comparing
226 the metabolic activity of active persisters to active R STM showed that the metabolic capacity
227 did not differ significantly (Fig. 4C), demonstrating that metabolically active NR STM showed
228 the same metabolic activity compared to R STM.

229 However, if about half of the persistent population was inactive at 24 h p.i., why did we only
230 detect a homogeneous stress-induced population instead of two subpopulations (Fig. 2B)? Is
231 there one subpopulation that showed stress responses and another lacking response? To
232 measure the metabolic activity at 24 h p.i., AHT was added 22 h p.i. If a persister is
233 metabolically active at 22 h p.i., this cell starts sfGFP synthesis (Fig. 4D), but inactive persisters
234 do not. However, a persister metabolically inactive at 22 h p.i. may have been active in the prior
235 period of infection (Fig. 4E). If this holds true, the proportion of active persisters should be
236 high at beginning of infection and decrease over time of infection, similar to result by Helaine
237 et al. (2014) for time points 24-72 h p.i. Since stress induction is measured by accumulation of
238 sfGFP, an active persister harboring the dual fluorescence stress reporter could synthesize

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239 sfGFP at the early phase of infection, become inactive e.g. at 10 h p.i., thus appears
240 metabolically inactive at 24 h p.i. However, such cell would still contain sfGFP synthesized in
241 response to stressors and should be detected as stress signal-positive at 24 h p.i. (**Fig. 4E**).

242 To test this hypothesis, we calculated the proportion of metabolically active and inactive
243 persisters in the early phase of infection (**Fig. 4GH**, FC histogram shown in **Fig. 4F**). For that,

244 AHT was added at indicated time points, however, host cells were lysed 24 h p.i. for clear
245 separation of R and NR STM. At 2 h p.i., appr. 95% of all NR STM were metabolically active.

246 Starting 8 h p.i., the amount of active persisters decreased constantly. The metabolic capacity
247 of this active persistent population was always on the same level compared to active R STM

248 (**Fig. 4I**). However, in the early phase of infection from 2-10 h p.i., the metabolic capacity of
249 active NR STM was even higher compared to active R STM. The higher x-median RFI of sfGFP
250 of both, active R and NR STM at the early time points post infection was resulting because all
251 samples were lysed 24 h p.i.

252 We also calculated the metabolic capacity of the entire NR STM population, because when
253 measuring stress induction of NR STM we are not able to discriminate between metabolically

254 active and inactive NR STM (**Fig. 4JK**). In addition, we calculated the ratio of the metabolic
255 activity of all NR compared to active R STM (**Fig. 4L**). Values above a ratio of 1 (red line)

256 indicate higher metabolic activity of the entire NR population. The metabolic activity of the
257 entire NR population up to 10 h p.i. was approximately at the same level as that of R STM. To

258 conclude, because of the same activity of R and NR STM, the lower stress response of persisters
259 8 h p.i. is not due to a lower metabolic capacity of persisters.

260 *Antibiotic exposure of persisters results in decreased stress response*

261 Since intracellular persisters are able to withstand bactericidal antibiotic exposure, we exposed
262 STM in RAW264.7 macrophages to cefotaxime and measured stress induction of NR STM WT

263 at 24 h p.i. As representative indicator for stress response induction P_{msrA} was used. At 10 h p.i.,
264 200 μ g x ml⁻¹ cefotaxime was added, following incubation for further 14 h. After cefotaxime

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265 treatment for 14 h, the subpopulation of R STM was highly reduced (**Fig. S 6A**). The proportion
266 of NR STM WT increased from 0.32% to 13% (**Fig. S 6B**) representing a 41.04-fold increase
267 (**Fig. S 6C**). However, the replicating population has not been fully removed, in contrast to prior
268 observations (Helaine et al., 2014). Fluorescence microscopy of infected RAW264.7
269 macrophages at 24 h p.i. (**Fig. S 6DE**) showed signal patterns supporting the FC data. Without
270 antibiotic treatment, STM WT replicated efficiently in macrophages also containing NR STM
271 showing DsRed fluorescence (**Fig. S 6D**).

272 After cefotaxime treatment, RAW264.7 macrophages contained both, R and NR STM WT (Fig.
273 S 6E, middle panel, indicated by red and green arrows), or NR STM WT only (**Fig. S 6E**, lower
274 panel). Analysis of stress response showed that after antibiotic treatment the x-median RFI of
275 sfGFP decreases, however, cefotaxime-treated NR STM WT still showed an active stress
276 response (**Fig. 5A**). When we checked the entire intracellular population, we observed that after
277 cefotaxime treatment the *msrA* induction also dropped to a level comparable to non-treated NR
278 STM.

279 Since persisters are a subpopulation of NR STM that are able to resume growth after release
280 from host cells, we controlled that we indeed analyzed growth-competent persister cells. For
281 that, we inoculated the released population after lysis of macrophages into fresh LB medium
282 and calculated the relative amount of DsRed-positive persisters 0, 2, 4 and 6 h after
283 reinoculation (**Fig. S 7A**). The number of DsRed-positive events (NR persisters) did not
284 decrease within the first 4 h after reinoculation (**Fig. S 7BC**). After that, the number of DsRed-
285 positive events decreased to about 50%, indicating that NR persisters are still viable and
286 competent to regrow. As determined in **Fig. S 1E-H**, ratios of up to 1:10,000 between STM
287 DsRed and STM sfGFP were precisely determined. In addition, detection of the optical density
288 of the reinoculated cultures showed that cefotaxime-treated STM WT started to replicate after
289 4-6 h (**Fig. S 7D**). Therefore, the persisters analyzed were able to regrow 6 h after reinoculation
290 that was in line with optical density of cultures.

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291 *Stress response of persisters within primary human phagocytes*

292 Finally, we determined the stress response of NR STM within primary human macrophages.

293 Primary cells were isolated from buffy coat and showed M1 polarization after differentiation

294 (*data not shown*). P_{msrA} was used as representative for stress response induction. Infection and

295 FC analysis was performed as described above and NR STM were detected inside human

296 macrophages at 8 h p.i. (**Fig. 6A**). Comparison of *msrA* induction of NR STM in murine and

297 human macrophages revealed an equal x-median RFI of sfGFP (**Fig. 6B**). In addition, the *msrA*

298 induction of the NR subpopulation [$P_{tetA}::dsred$ $P_{msrA}::sfgfp$] and the entire population

299 [$P_{EM7}::dsred$ $P_{msrA}::sfgfp$] showed the same level in human macrophages. Plotting the bacterial

300 population against their AHT-induced DsRed and *msrA*-induced sfGFP intensity showed that

301 in primary human macrophages only NR STM were present, while a subpopulation of R STM

302 was absent (**Fig. 6C**). To conclude, also in primary human macrophages NR STM WT showed

303 an active stress response that was comparable to RAW264.7 macrophages.

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305 **Discussion**

306 *NR intracellular STM are capable to respond to host-imposed stressors*

307 In this study, we further developed and applied FP-based reporters for analysis of stress
308 response of intracellular NR STM at single cell level. This approach enables quantification of
309 the stress response of distinct intracellular subpopulations of STM. Our study demonstrates that
310 NR STM persister cells perceive their intracellular environment and respond to stressors by
311 activating a stress response even under antibiotic pressure (Fig. 7). These cells regrew in rich
312 medium after infection, proving that proper NR STM persister cells were examined.

313 Our findings support previously reported metabolic activity of intracellular STM persisters, and
314 translocation by T3SS-SPI2 (Helaine et al., 2014; Helaine et al., 2010; Stapels et al., 2018).

315 STM persisters express SPI2 genes and for this, bacteria must perceive their environment to
316 activate gene expression (Stapels et al., 2018). Our results corroborate that STM persister cells
317 are not segregated from the intracellular habitat, but rather sense their environment and changes
318 within. Persisters that maintain effector delivery and are able to respond to stress factors but
319 cease to grow, still provide evolutionary benefit. Since the persister status is reversible and a
320 single cell can give rise to a new susceptible population, the intracellular heterogeneity of
321 *Salmonella* provides adaptive advantage and favors survival of the entire population during
322 infection of host cells or in general, under adverse environmental conditions. Since we observed
323 lower stress response of persister cells compared to R STM, independent from the duration of
324 infection or function of virulence factors, we conclude that the persistent status serves as a kind
325 of niche by protecting NR STM from severe stress exposure (Fig. 7). The host cell may no
326 longer be able to exert stress on NR STM persister cells, which in turn results in lower stress
327 response. Beyond this, STM persisters reside individually within segregated SCV during the
328 whole course of infection, further promoting their protective niche. This compartment shields
329 STM from the cytosolic environment and host cell defense mechanisms such as xenophagy

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330 (reviewed in Gomes and Dikic, 2014). In addition, the SCV of NR STM is separated from the
331 SCV/SIF continuum containing R STM (Liss et al., 2017).

332 *At initial infection, a high proportion of NR STM is metabolically active*

333 We observed that at initial infection of macrophages, the proportion of metabolically active NR
334 STM was as high as 95%. Thus, the lower stress response of NR STM is not a consequence of

335 a lower metabolic capacity. Starting about 8 h p.i., the proportion of active persisters constantly
336 decreased. This is in line with previous observations that detected about 70% of active NR STM

337 in RAW264.7 macrophages 12 h p.i., and about 25% active NR STM at 24 h p.i. (Helaine et
338 al., 2010). Our studies extended these analyses to early time points p.i. In addition, we compared

339 the level of metabolic activity, which was comparable to R STM until 10 h p.i. if we average
340 the total activity of the entire NR population. If only metabolically active NR STM persisters

341 are considered, these did not show a significantly different activity compared to active R STM
342 over the entire period of infection. If a small reservoir of persisters always remains active over

343 a longer period of time, this may allow formation of new populations in the host, and thus to
344 trigger relapses. The metabolic activity of NR STM at the beginning of infection may serve to

345 successfully establish their persistent status. For *M. tuberculosis* or *E. coli*, *in vitro*
346 transcriptome studies indicated that persisters showed downregulation of metabolic genes and

347 therefore decreased metabolism (Keren et al., 2011; Shah et al., 2006). Genes with functions in
348 energy production and TCA cycle were downregulated in *Burkholderia cenocepacia* persisters

349 (Van Acker et al., 2013). For intracellular STM, however, there is evidence that distinct
350 subpopulations of metabolically active and inactive persisters exist (Helaine et al., 2010;

351 Staples et al., 2018). The observation of metabolically active NR bacteria is not limited to
352 *Salmonella* species, but also reported for *M. tuberculosis* in macrophages, and could therefore

353 be a widespread phenomenon (Manina et al., 2015).

354 When examining persistent bacteria, it is highly important which time point is analyzed, and to
355 be aware that overall metabolic activity of the NR population changes over time. This refers to

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356 the decreasing proportion of metabolically active persisters over time, and the associated
357 activity of the entire population on average. In many studies, late time points were selected to
358 investigate persisters, at which a small proportion of metabolically active persisters was
359 detected, which further decreased (Helaine et al., 2010). Our studies are in line with these results
360 and our data also showed that when the entire NR subpopulation is considered, a decreased
361 metabolic activity from 14 h p.i. was observed. This is also in line with a model of reduced
362 metabolism in persisters cells. Importantly, before 14 h p.i. the entire population of NR STM
363 persisters on average does not show a reduced metabolism and the metabolic activity is not
364 lower at any time point analyzed if only metabolically active NR STM are considered.
365 However, since the proportion of metabolically active persisters decreases over time, the
366 metabolic activity of the entire population of NR STM on average decreases accordingly. In
367 turn, a reduced metabolic activity is measured when the entire population of NR STM is
368 observed. Conventional transcriptome analyses or other population-wide approaches average
369 the entire NR population and fail to distinguish between active and inactive persisters. Single
370 cell analyses that are capable of addressing small subpopulations of intracellular STM, such as
371 the approaches applied here, provide a more precise insight into the physiological state of
372 persisters. Novel single cell transcriptomics approaches for bacteria are emerging (Imdahl et
373 al., 2020) and may provide a future option for analyses of persisters.

374 *Stress response of NR STM in non-permissive primary human macrophages*

375 We investigated stress response of NR STM persisters in much less permissive human
376 phagocytic cells. We observed that the level of stress response triggered human macrophages
377 was comparable to that of STM in RAW264.7 macrophages being rather permissive for STM
378 intracellular proliferation. These results suggest that the induced stress response of NR STM is
379 either independent of the killing capacity of the host cell, or because persisters are so well
380 protected that even phagocytes with high antimicrobial activity do not induce higher levels of
381 stress response.

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382 Typhoidal serovars (TS) of *S. enterica* cause systemic infections in humans such as typhoid
383 fever, and persistent or recurrent forms of the disease are frequent (Gotuzzo et al., 1987).
384 Advanced analyses of stress response shall be extended to persister cells of typhoidal serovars
385 Typhi and Paratyphi A to gain insights into their physiological state, and to test the ability to
386 perceive stress factors. As part of the ‘stealth strategy’ (Dougan and Baker, 2014), persister
387 formation in typhoidal serovars may occur at higher frequency. In relation to heterogeneity, it
388 will be of interest if intracellular Typhi and Paratyphi A show distinct subpopulations with
389 different levels of stress response. Knowledge of persister cell frequency, the proportion of
390 metabolically active and inactive persisters, and the level of stress response of TS in comparison
391 to STM may lead to new insight into patho-mechanisms of TS and recurrent infections. This
392 will require further analyses in other cell lines like human phagocytic cell line U937 or primary
393 phagocytic cells.

394 *Stress response of persisters and new therapeutic options?*

395 Since persisters that emerge during infection are extremely well protected and the clinical
396 relevance of relapsing infection due to persister is high, new strategies to eliminate persister
397 cells are of utmost therapeutic importance. Current approaches include prevention of persister
398 formation, identification of antimicrobial compounds that act on persister cells, or resuscitation
399 of persister cells to restore sensitivity to conventional antibiotics (reviewed in Defraine et al.,
400 2018). Promising studies have been performed to identify compounds acting on bacterial
401 persister cells, however further action is required. High-throughput screening methods are often
402 applied, but many screens were performed using planktonic persisters *in vitro*. Recent work
403 showed that the efficacy of drugs against tuberculosis is significantly different when applied *in*
404 *vivo* compared to *in vitro* conditions (Liu et al., 2016). This may also apply to other drugs,
405 making analyses of effects on persister cells during interplay with their host essential. One
406 approach could be host-directed therapies focusing on host-pathogen interactions, as already
407 has been demonstrated for *M. tuberculosis* infections by treating infections with small

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408 molecules to interfere with host responses or persistence (Kim and Yang, 2017; Kiran et al.,
409 2016). Antibody-antibiotic conjugates are another emerging approach. Antibiotics are coupled
410 to antibodies against a specific pathogen to increase the efficiency of antibiotics against
411 intracellular pathogens including persisters (Mariathasan and Tan, 2017). Zhou et al. (2016)
412 showed that *S. aureus* bacteremia in mice was successfully treated, hence, this approach could
413 serve as a novel therapeutic platform in the future. Our finding of the continuing function of
414 SRS in persister cells and the ability of persisters to sense stressors may lead to new options for
415 resuscitation of persisters (examples in Song and Wood, 2020). If deregulation of SRS by decoy
416 compounds is possible, this may lead to re-initiation of normal growth and increased antibiotic
417 susceptibility. Approaches to interfere with quorum sensing for interbacterial communication
418 (reviewed in Defoirdt, 2018) or biofilm matrix production (Dieltjens et al., 2020) enabled new
419 antimicrobial strategies. In contrast, potential interference with SRS in persisters needs to target
420 single cells in complex populations within host organisms.

421 *Conclusions and outlook*

422 In summary, we have developed and applied new dual FP reporters that enable extremely
423 sensitive analyses of stress response and metabolic activity of NR persister cells. Of particular
424 interest is that NR STM perceive their environment and display a lower stress response with
425 constant metabolic activity during the early phase of infection. Our findings provide deeper
426 knowledge that besides the ability to subvert immune defenses of the host, NR STM persister
427 cells maintain capability to sense stressors and react to stress. These persisters exhibit constant
428 metabolic activity at the beginning of the infection, which supports the pathogen to create a
429 more permissive environment for recrudescent infections.

430

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437 **Materials and Methods**

438 *Generation of reporter plasmids*

439 For the generation of a reporter plasmid to detect non-replicating STM, plasmid p5084
440 ($P_{EM7}::dsred P_{msrA}::sfgfp$), p5085 ($P_{EM7}::dsred P_{trxA}::sfgfp$) and p5055 ($P_{EM7}::dsred P_{htrA}::sfgfp$)
441 with a constitutive expression of Dsred and *msrA*, *trxA* or *htrA* regulated expression of sfGFP
442 was used. Via Gibson Assembly (GA) of PCR fragments, the EM7 promoter was replaced by
443 the tet-ON cassette to enable artificial induction of DsRed by anhydrotetracycline (AHT).

444 For the generation of a single FP reporter plasmid to detect non-replicating STM, plasmid
445 p5204 ($P_{EM7}::dsred P_{cypD}::sfgfp$ (*frameshift*)) with a constitutive expression of *dsred* and a
446 frameshift mutation in *sfgfp* was used. Via GA, the EM7 promoter was replaced by the tet-ON
447 cassette to enable artificial induction of *dsred* by AHT.

448 For the generation of a dual fluorescence vitality reporter plasmid to detect the metabolic
449 activity of non-replicating STM, plasmid p4928 ($P_{EM7}::tagrfp-T P_{tetA}::sfgfp$) with a constitutive
450 expression of *tagrfp-T* and an AHT-inducible expression of *sfgfp* was used. Via GA, the EM7
451 promoter was replaced by *araC* P_{BAD} cassette to enable artificial induction of *tag-rfpT* by
452 arabinose. Subsequently, p5419 ($P_{BAD}::tagrfp-T P_{tetA}::sfgfp$) was used to amplify the P_{BAD} and
453 tet-ON cassette. Via GA, the EM7 promoter and the *msrA* promoter of p5084 ($P_{EM7}::dsred$
454 $P_{msrA}::sfgfp$) was replaced by the P_{BAD} and tet-ON cassette to enable artificial induction of *dsred*
455 by arabinose and *sfgfp* by AHT. The resulting single and dual FP reporter plasmids are listed in
456 Table S 1 and oligonucleotides used for construction are listed in Table S 3.

457 *Bacterial strains, growth conditions AHT and arabinose induction*

458 *Salmonella enterica* sv. *Typhimurium* strain NCTC12023 (STM) was used as wild-type strain
459 and isogenic mutant strains used in this study are listed in Table S 2. Bacteria were cultured in
460 lysogeny broth (LB) at 37 °C overnight (o/n) using a roller drum at 60 rpm with aeration. For
461 maintenance of plasmids, carbenicillin was added at 50 µg x ml⁻¹ as a selection marker. For

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462 induction of expression of P_{tetA} -controlled dual FP reporter (p5205, p5300, p5202 or p5418) or
463 P_{BAD} -controlled dual FP reporter (p5426) AHT or arabinose always was directly added to LB
464 to a concentration of 50 ng x ml⁻¹ or 13.3 mM, respectively, in the o/n culture and removed
465 when necessary by centrifugation for 3 min at 5,000 x g and washing with fresh LB.

466 *Isolation of primary human macrophages*

467 Buffy coat was obtained from pooled samples of voluntary, anonymous blood donors via the
468 blood bank of Deutsches Rotes Kreuz (Springe, Germany). Lymphocytes were prepared from
469 buffy coat by Ficoll-Hypaque density gradient centrifugation (see Bonifacino et al., 2004).
470 Buffy coat was diluted in PBS (ratio 1:3) and Ficoll-Hypaque was added, following
471 centrifugation for 20 min at 800 x g. Afterwards, 5 x 10⁸ cells were cultured in Roswell Park
472 Memorial Institute (RPMI) medium containing 5.5 g x l⁻¹ NaCl, 5.0 mg x l⁻¹ phenol red, 2.0 g
473 x l⁻¹ NaHCO₃, 25 mM HEPES, 4 mM stable glutamine without sodium pyruvate (Biochrom),
474 100 units x ml⁻¹ penicillin, 100 µg x ml⁻¹ streptomycin, 2.5 ng x ml⁻¹ GM-CSF (granulocyte-
475 macrophage colony-stimulating factor) and supplemented with 10% human plasma. After o/n
476 incubation at 37 °C in an atmosphere of 5% CO₂ and 90% humidity, non-adherent cells were
477 removed and adherent cells were cryo-preserved in inactivated fetal calf serum (iFCS) and
478 DMSO.

479 *Cell lines and cell culture*

480 For infection experiments murine RAW264.7 macrophages (American Type Culture
481 Collection, ATCC no. TIB-71), RAW264.7 macrophages stably transfected with LAMP1-GFP
482 or primary human macrophages were used. RAW264.7 macrophages were cultured in
483 Dulbecco's modified Eagle's medium (DMEM) containing 3.7 g x l⁻¹ NaHCO₃, 4.5 g x l⁻¹
484 glucose, 4 mM stable glutamine without sodium pyruvate (Biochrom) and supplemented with
485 6% iFCS (Sigma-Aldrich). Primary human macrophages were cultured in RPMI medium
486 containing 5.5 g x l⁻¹ NaCl, 5.0 mg x l⁻¹ phenol red, 2.0 g x l⁻¹ NaHCO₃, 25 mM HEPES, 4 mM

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487 stable glutamine without sodium pyruvate (Biochrom) and supplemented with 10% iFCS
488 (Sigma-Aldrich).

489 *Host cell infection (gentamicin protection assay) for cytometry*

490 Before infection, RAW264.7 or primary human macrophages were seeded in surface-treated 6-
491 well plates (TPP) to reach confluence ($\sim 2 \times 10^6$ cells per well) on the day of infection. For
492 infection, *Salmonella* strains were grown o/n in LB broth (app. 18 h). Infection was performed
493 with a multiplicity of infection (MOI) of 10. Bacteria were centrifuged onto the cells for 5 min
494 at $500 \times g$ and infection proceeded for 25 min at 37 °C in an atmosphere of 5% CO₂ and 90%
495 humidity. Afterwards, infected cells were washed thrice with PBS and incubated for 1 h with
496 cell culture medium containing 100 $\mu\text{g} \times \text{ml}^{-1}$ gentamicin (Applichem) to kill non-phagocytosed
497 bacteria. Afterwards, the cell culture medium was replaced by medium containing 10 $\mu\text{g} \times \text{ml}^{-1}$
498 gentamicin until the end of the experiment. If cefotaxime was used during infection
499 experiments, cells were washed with PBS before addition of cell culture medium containing
500 200 $\mu\text{g} \times \text{ml}^{-1}$ freshly dissolved cefotaxime.

501 *Host cell infection for microscopy*

502 RAW264.7 or LAMP1-GFP RAW264.7 macrophages were seeded in surface-treated 24-well
503 plates on glass cover slips to reach 80% confluence (ca. 3.6×10^5 cells per well) on the day of
504 infection. Cells were infected with STM strains as described above with a MOI of 50 for 8 or
505 24 h. Afterwards, the cells were washed thrice with PBS and fixed with 3% PFA in PBS for 15
506 min at RT. After that, cells were directly prepared for subsequent immunostaining.

507 *Immunostaining and imaging*

508 Immunostaining of intracellular STM was performed as described before (Müller et al., 2012).
509 After fixation of cells with 3% PFA in PBS cells were washed thrice with PBS and directly
510 incubated in blocking solution containing 2% goat serum, 2% bovine serum albumin and 0.1%
511 saponin in PBS for 30 min at RT. STM was stained with anti-*Salmonella* O-antigen group B

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512 factors 1, 4, 5, 12 (BD Difco, diluted 1:500 in blocking solution) for 1 h at RT. Subsequently,
513 cells were washed thrice in drops of PBS following incubation with secondary antibody Cy5-
514 coupled goat anti rabbit IgG (Jackson Immuno Research, 1:1,000 in blocking solution) for 1 h
515 at RT in the dark. Afterwards, cells were washed thrice, mounted with Fluoroprep (bioMérieux)
516 and sealed with Entellan (Merck). Fluorescence imaging was performed using the confocal
517 laser-scanning microscope Leica SP5. Image acquisition was performed using the 100x
518 objective (HCX PL APO CS 100×; numerical aperture: 1.4 to 0.7) and the polychroic mirror
519 TD 488/543/633 for the three channels GFP, DsRed, and Cy5 (Leica, Wetzlar, Germany). For
520 setting adjustment, image acquisition and image processing the software LAS AF (Leica,
521 Wetzlar, Germany) was used.

522 *Flow cytometry analysis*

523 FC of liberated STM from host cells was performed as described before (Noster et al., 2019).
524 Briefly, FC was performed on an Attune NxT instrument (Thermo Fischer Scientific) at a flow
525 rate of 25 μ l x min⁻¹. At least 10,000 bacteria were gated by virtue of the constitutive/induced
526 DsRed fluorescence. Per gated STM cells, the intensity of the sfGFP fluorescence was
527 determined and x-medians for the sfGFP intensities were calculated.

528 For the measurement of liberated bacteria (replicating and non-replicating) from host cells,
529 infected cells were lysed at the indicated time points by 0.5% Triton X-100 in PBS for 10 min
530 at RT with shaking to release the intracellular bacterial population. The lysate was transferred
531 to a test tube and after pelleting of host cell debris by centrifugation for 5 min at 500 x g, bacteria
532 were recovered from supernatant. Bacteria were further centrifuged for 5 min at 20,000 x g and
533 fixed in 3% PFA in PBS for 15 min at RT. After fixation and a further centrifugation step, fixed
534 bacteria were resuspended in 250 μ l 100 mM NH₄Cl in PBS for quenching of residual free
535 aldehydes. After that, samples were directly subjected to FC.

536 For detection of regrowth of cefotaxime-treated intracellular non-replicating STM, the whole
537 liberated bacterial population was reinoculated into 3 ml fresh LB medium after the bacterial

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538 supernatant was transferred to a clean test tube (Fig. S 7) and incubated at 37 °C using a roller
539 drum at 60 rpm with aeration. At time point 0 h, 2 h, 4 h and 6 h post re-inoculation samples
540 were taken, diluted with PBS and fixed with 3% PFA in PBS for 15 min at RT. Then, PFA was
541 removed by centrifugation for 5 min at 20,000 *x* g and the pellet was resuspended in 250 µl of
542 100 mM NH₄Cl in PBS for quenching of residual free aldehydes. After that, samples were
543 directly subjected to FC and amounts of DsRed-positive events (non-replicating STM) was
544 measured and depicted by events per µl.

545 To measure the minimal detectable threshold of bacterial events at the cytometer, various mixed
546 ratios of constitutive DsRed and sfGFP fluorescent STM were prepared. For that, the optical
547 density (OD₆₀₀) of overnight cultures of STM [p5204] and STM [pWRG167] was determined
548 following dilution of the strains to an OD₆₀₀ of 1 (predicted amounts of bacteria: 1.1 x 10⁹
549 bacteria x ml⁻¹). Afterwards, serial dilutions were prepared and mixed ratios of red and green
550 fluorescent STM were prepared in PBS. Either a constant high amount of green fluorescent
551 STM (app. 2,000 events x µl⁻¹) mixed with an equal, 10-, 100-, 1,000-, or 10,000-fold reduced
552 amount of red fluorescent STM or a constant high amount of red fluorescent STM (app. 4,000
553 events x µl⁻¹) mixed with an equal, 10-, 100-, 1,000-, or 10,000-fold reduced amount of green
554 fluorescent STM was measured by FC. The relative amount of green and red fluorescent STM
555 (events x µl⁻¹) measured when the equal amount of green and red fluorescent STM was present
556 in the sample was set to 100%. As controls, only green, red or no fluorescent STM were
557 measured.

558

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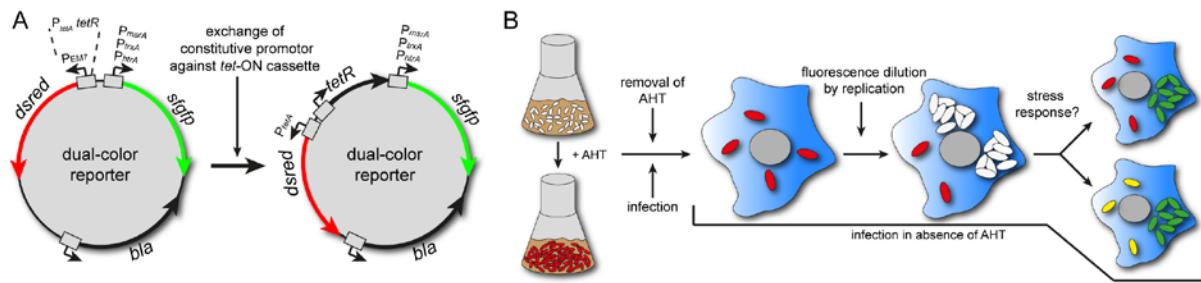
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694 *Staphylococcus aureus* in mice. MAbs 8, 1612-1619. 10.1080/19420862.2016.1229722
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697 **Figure legends**

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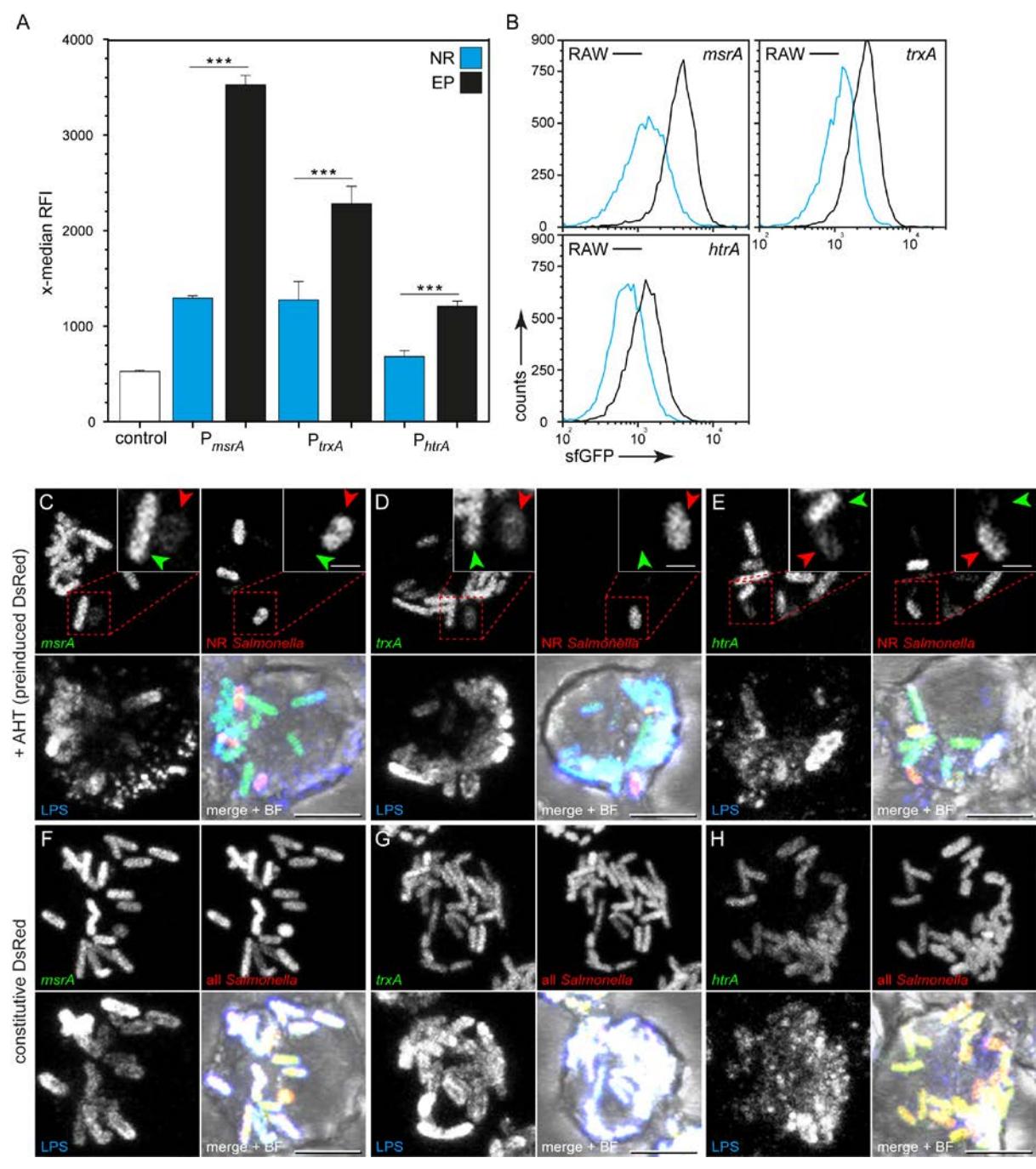


701 **Fig. 1: Design of reporters to measure stress response of non-replicating STM.** A) For
702 detection of NR intracellular STM, the EM7 promoter of dual FP reporters for *msrA*, *trxA*, or
703 *htrA* was replaced by the tet-ON cassette to enable controlled induction of *dsred* expression by
704 AHT. B) Addition of AHT to growing cultures of STM induced DsRed synthesis. Before
705 infection of macrophages, AHT was removed by centrifugation and washing. Infection was
706 performed in absence of AHT. After infection, R intracellular STM dilute cellular DsRed levels.
707 NR intracellular STM maintain cellular DsRed levels and were detected as DsRed-positive
708 events. In addition, the *msrA*-induced sfGFP intensity provides information about response to
709 stress imposed by the host cell.

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713 **Fig. 2: Lower levels of stress response of non-replicating STM WT compared to**
714 **replicating intracellular STM.** STM WT harboring AHT-inducible (NR, blue bars) or
715 constitutive (EP, black bars) dual fluorescence reporter for *msxA*, *trxA* and *htrxA* were grown o/n
716 in LB medium containing 50 ng x ml⁻¹ AHT. Before infection of RAW264.7 macrophages,
717 AHT was removed. At 8 h p.i., infected cells were lysed, or fixed for FC analyses (A, B), or
718 microscopy (C-H), respectively. For FC, released STM were recovered, fixed, and analyzed.
719 As non-induced control, dual FP reporter plasmids were used with a frame shift mutation in

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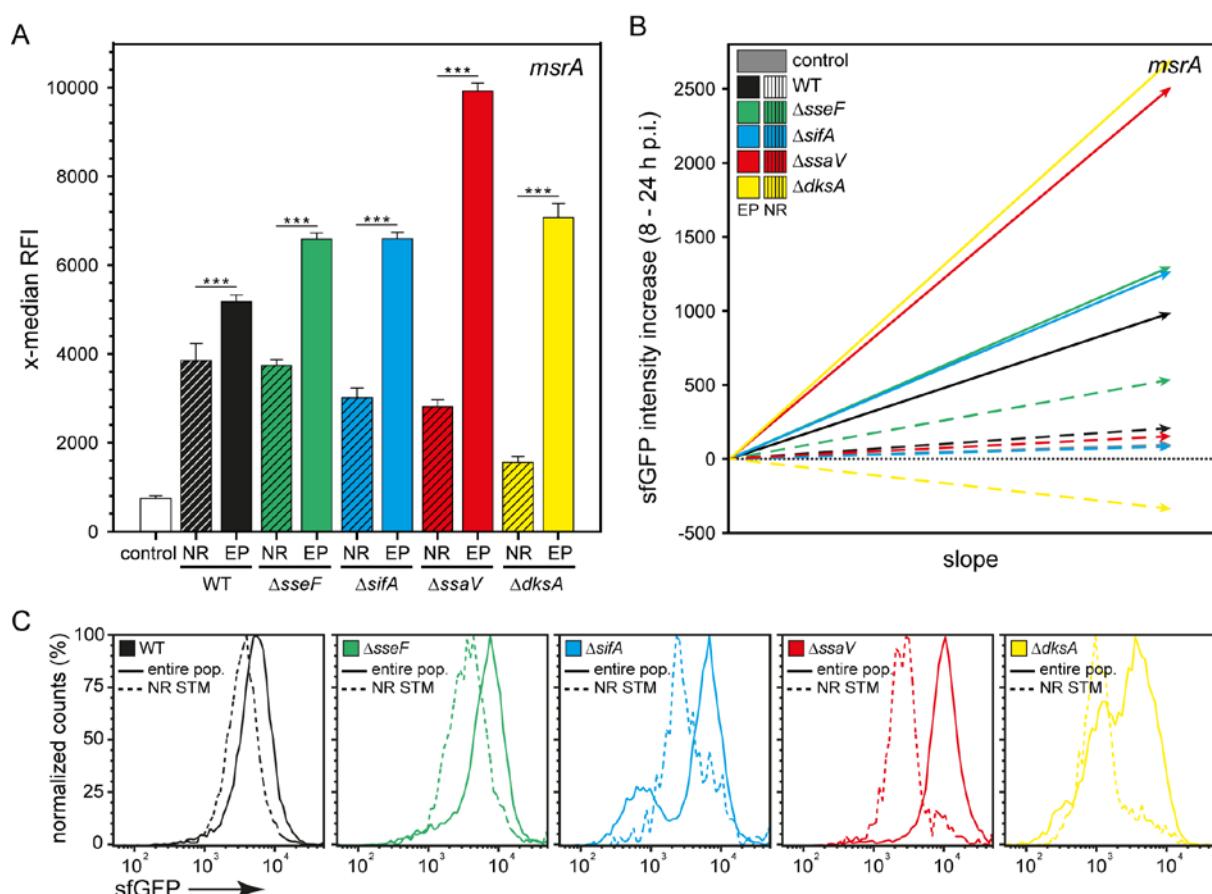
Stress response of *Salmonella* persisters

720 *sfgfp* (described before in (Noster et al., 2019). A) The x-median represents the stress-induced
721 sfGFP signal of the DsRed-positive intracellular bacterial population 8 h p.i. Means and
722 standard deviations of one representative experiment are shown. B) Representative histograms
723 corresponding to A) are shown. Statistical analysis was accomplished by SigmaPlot by One
724 Way ANOVA and significance levels are indicated as follows: *, p < 0.05; **, p < 0.01; ***, p
725 < 0.001; n.s., not significant. C-H) The entire population of intracellular STM was detected by
726 immuno-staining of O antigen (blue). F-H) In addition, constitutively *dsred*-expressing STM
727 show the entire intracellular bacterial population. C-E) NR and R STM were positive or
728 negative for DsRed fluorescence (red), respectively. sfGFP signals (green) indicate induction
729 of *msrA*, *trxA* or *htrA*. Representative NR STM and R STM are indicated by red and green
730 arrowheads, respectively. Scale bars, 5 and 1 μ m in overview and detail, respectively.

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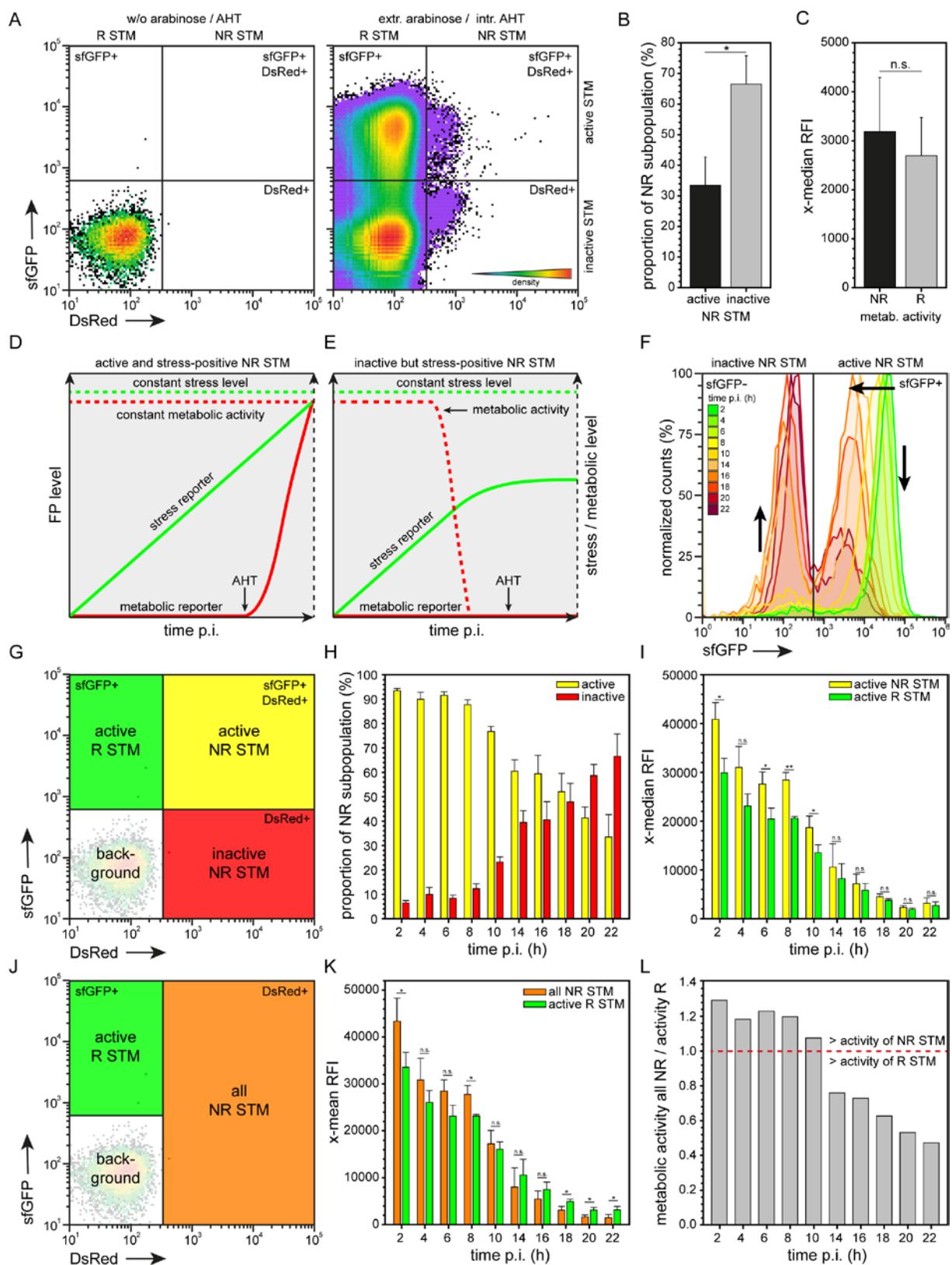


732

733 **Fig. 3: Non-replicating SPI2-T3SS or SR mutant strains also show lower stress response**
734 **compared to respective subpopulation.** STM WT, Δ sseF, Δ sifA, Δ ssaV, and Δ dksA strains
735 harboring AHT-inducible (NR), or constitutive (entire population, EP) dual fluorescence
736 reporter for *msrA* were grown o/n in LB medium. AHT was present for AHT-inducible reporters
737 and removed before infection. RAW264.7 macrophages were infected, lysed 24 h p.i., liberated
738 STM were recovered, fixed, and subjected to FC analyses. As negative control, a dual FP
739 reporter plasmid with *sfgfp* inactivated by a frame-shift was used. A) The x-median represents
740 the *msrA*-induced sfGFP signal of the DsRed-positive intracellular bacterial population at 24 h
741 p.i. Means and standard deviations of one representative experiment are shown. B) The depicted
742 slopes represent the sfGFP intensity increase over the time of infection. C) Representative
743 histograms corresponding to A) are shown. Statistical analysis was performed as described for
744 **Fig. 2.**

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745

746 **Fig. 4: Non-replicating persisters show equal metabolic activity compared to replicating**
 747 **STM WT in the early phase of infection.** STM WT harboring double-inducible dual
 748 fluorescence reporter was grown o/n in LB medium in the presence of arabinose. Before

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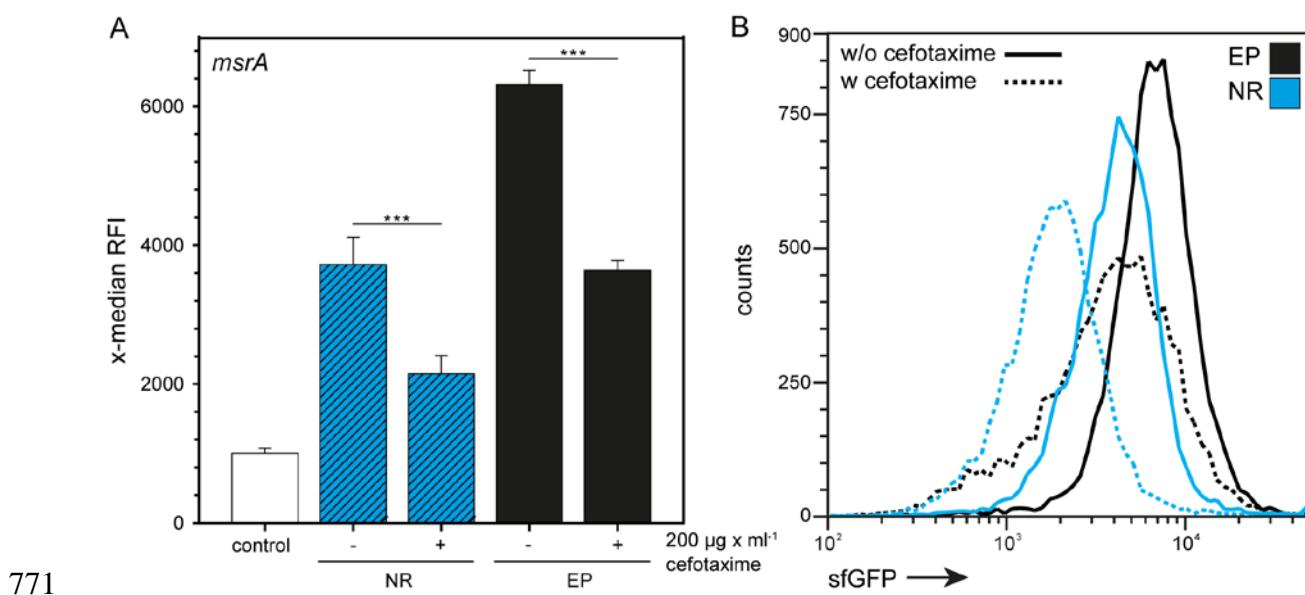
Stress response of *Salmonella* persisters

749 infection, arabinose was removed. RAW264.7 macrophages were infected, AHT was added
750 22 h p.i., and cells were lysed 24 h p.i. Liberated STM were recovered, fixed and subjected to
751 FC. A) Plotting the entire intracellular bacterial population against their arabinose-induced
752 DsRed and AHT-induced sfGFP intensity shows distinct populations. As control, STM without
753 addition of arabinose or AHT was analyzed. B) The proportion of metabolically active and
754 inactive NR STM WT at 22 h p.i. is shown. C) Comparison of the metabolic activity of active
755 NR and R STM WT 22 h p.i. D-H) STM WT was cultured, infected and prepared for FC as
756 described above. AHT was added at various time points p.i. and cells were lysed 24 h p.i. D)
757 FP levels anticipated for metabolically active and stress signal-positive NR STM. E) Detected
758 metabolically inactive NR STM may be stress signal-positive because of metabolic activity
759 before addition of AHT. F) Representative histograms of the metabolic activity of intracellular
760 NR STM WT at various time points p.i. Arrows indicate increase in metabolically inactive NR
761 STM, and decrease in metabolically active NR STM over time of infection. G) Yellow, green,
762 and red gates define subpopulations compared in (H) and (I). H) The proportion of
763 metabolically active and inactive NR STM at the various time points was calculated. I)
764 Comparison of the metabolic activity of active NR and R STM WT at various time points p.i.
765 J) Green and orange gates define subpopulations compared in (K). K) Comparison of the
766 metabolic activity of all NR and R STM WT at various time points p.i. L) The ratio of metabolic
767 activity of all NR STM to metabolic activity of R STM is shown. Means and standard deviations
768 of one representative experiment are shown. Statistical analysis was performed as described for
769 **Fig. 2.**

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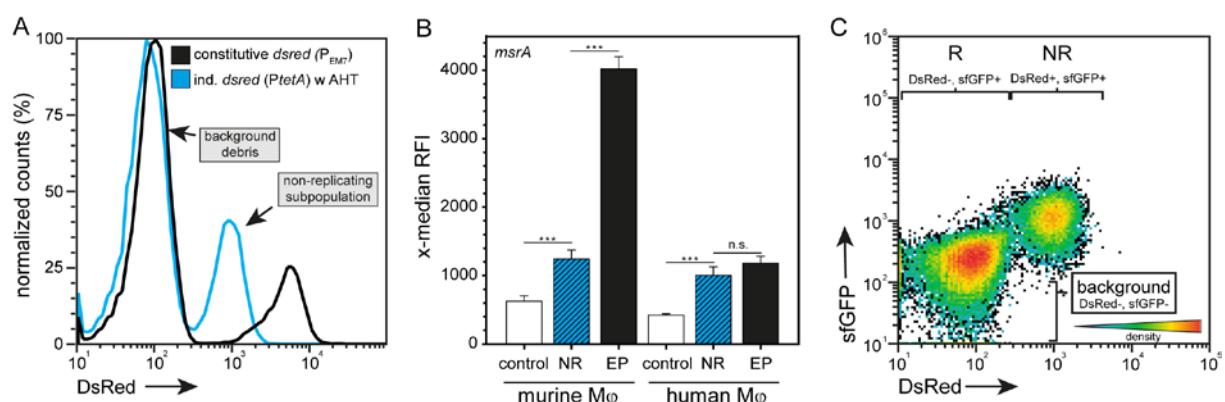
772 **Fig. 5: Cefotaxime-treated non-replicating STM WT show slight induction of stress response.** STM WT harboring AHT-inducible (NR) or constitutive (EP) dual fluorescence reporter for *msrA* was grown o/n in LB medium, in the presence of AHT or the NR reporter. 773 Before infection, AHT was removed. RAW264.7 macrophages were infected, cefotaxime was 774 added to the cells at 10 h p.i. if indicated, and cell were lysed at 24 h p.i. Liberated STM were 775 recovered, fixed and subjected to FC. As negative control, a dual FP reporter plasmid with *sfgfp* 776 inactivated by a frame-shift was used. The x-median (A) represents the P_{*msrA*}-induced sfGFP 777 signal of the DsRed-positive intracellular bacterial population, and (B) shows a representative 778 histogram. Means and standard deviations of one representative experiment are shown. 779 780 Statistical analysis was performed as described for **Fig. 2**.

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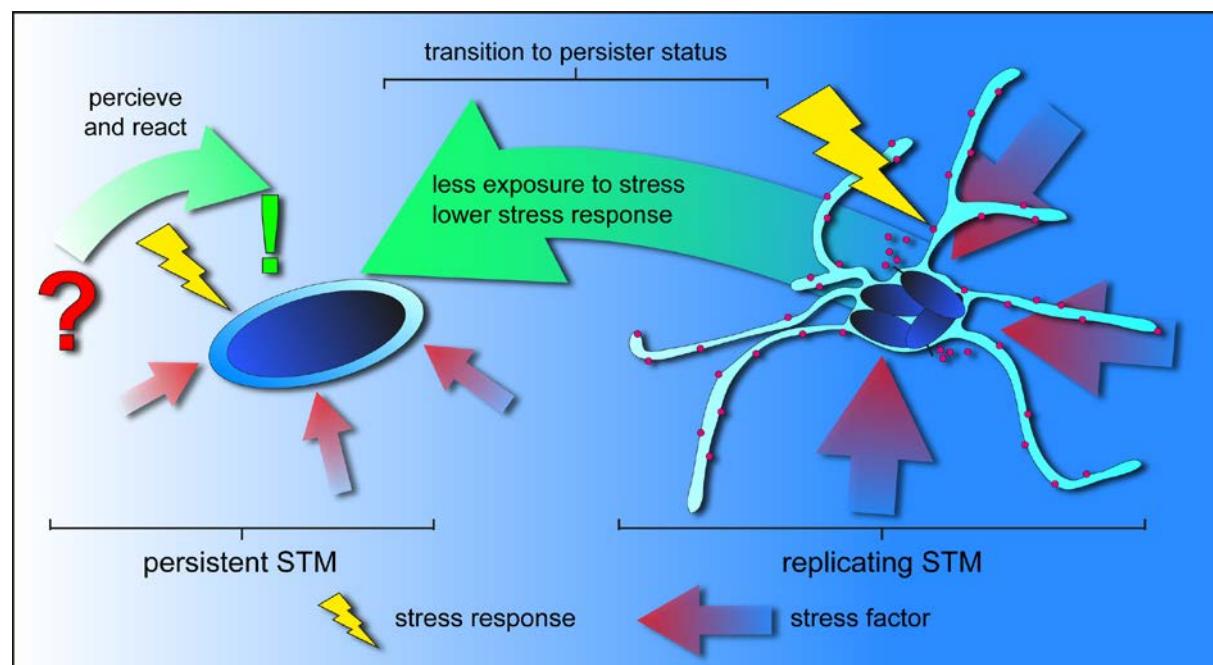


784 **Fig. 6: Stress response of non-replicating STM WT in primary human macrophages**
785 **isolated from peripheral blood.** STM WT harboring AHT-inducible (NR) or constitutive (EP)
786 dual fluorescence reporter for *msrA* was grown o/n in LB medium in the presence of AHT if
787 necessary. Before infection, AHT was removed. Murine RAW264.7 or human primary
788 macrophages from peripheral blood were infected and lysed 24 h p.i. Liberated STM were
789 recovered, fixed, and subjected to FC. As negative control, a dual FP reporter plasmid with
790 *sfgfp* inactivated by a frame-shift was used. A) Detection of intracellular NR STM in human
791 macrophages. B) Comparison of *msrA* induction within murine and human macrophages. The
792 x-median represents the *msrA*-induced sfGFP signal of DsRed-positive intracellular bacteria.
793 Means and standard deviations of one representative experiment are shown. C) STM WT only
794 shows a NR subpopulation when plotting the bacterial population against their AHT-induced
795 DsRed and *msrA*-induced sfGFP intensity. Statistical analysis was performed as described for
796 **Fig. 2.**

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Stress response of *Salmonella* persisters



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799 **Fig. 7: Transition to persister status results in decreases exposure to stressors and enables**
800 **survival of harsh conditions in host cells.** Replicating intracellular STM show a diverse stress
801 response in reaction to harsh environmental conditions. Intracellular persistent STM are able to
802 perceive and actively respond to stressors by turning on SRS. The induced stress response of
803 persister is lower compared to the replicating subpopulation of intracellular STM suggesting
804 that persister encounter lower exposure to stressors. By that, transition to persistent state inside
805 host cells promotes stress tolerance, survival, and dissemination of the pathogen due to re-
806 growth after subsidence of stressful conditions.

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808 **Suppl. Tables**

809 Table S 1. Plasmids used in this study

810	<u>Plasmid</u>	<u>Relevant genotype*</u>	<u>Source/reference</u>
811	p5204	$P_{EM7::dsred} P_{cypD::sfgfp}$ (frameshift)	(Noster et al., 2019)
812	pWRG167	$P_{EM7::sfgfp}$	(Bender et al., 2013)
813	p4928	$P_{EM7::tagrfp-T} P_{tetA::sfgfp}$	(Schulte et al., 2020)
814	p5055	$P_{EM7::dsred} P_{htrA::sfgfp}$	(Schulte et al., 2020)
815	p5084	$P_{EM7::dsred} P_{msrA::sfgfp}$	(Noster et al., 2019)
816	p5085	$P_{EM7::dsred} P_{trxA::sfgfp}$	(Noster et al., 2019)
817	p5205	$P_{tetA::dsred} P_{msrA::sfgfp}$	This study
818	p5300	$P_{tetA::dsred} P_{htrA::sfgfp}$	This study
819	p5302	$P_{tetA::dsred} P_{trxA::sfgfp}$	This study
820	p5426	$P_{BAD::dsred} P_{tetA::sfgfp}$	This study
821	p5418	$P_{tetA::dsred} P_{cypD::sfgfp}$ (frameshift)	This study
822	p5419	$P_{BAD::tag-rfp-T} P_{tetA::sfgfp}$	This study

823 * all plasmids confer resistance to carbenicillin

824

825 Table S 2. *Salmonella enterica* serovar Typhimurium strains used in this study

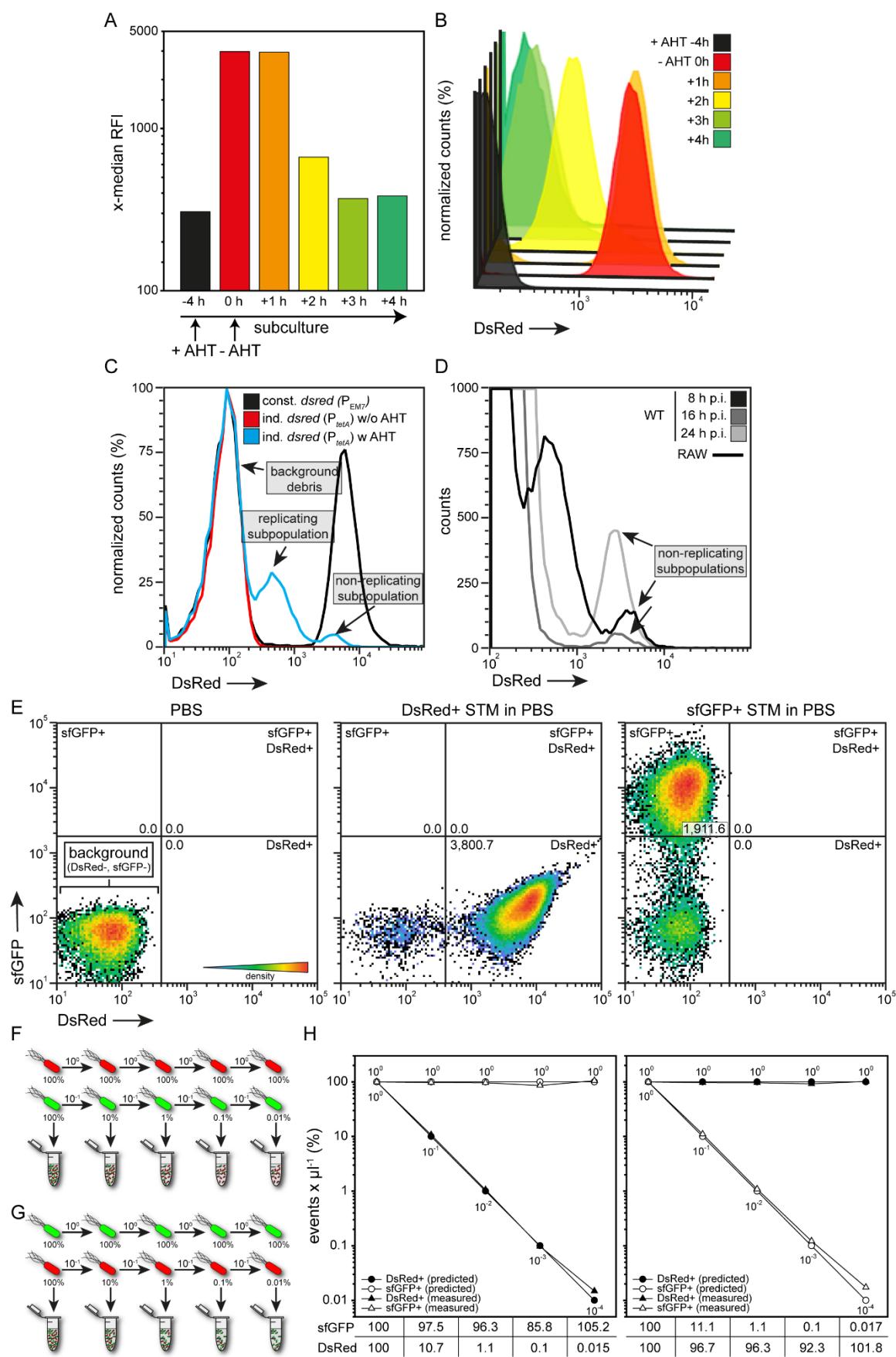
826	<u>Strain</u>	<u>Relevant characteristics</u>	<u>Source/reference</u>
827	NCTC12023	wild type	NCTC, lab stock
828	MvP1890	$\Delta ssAV::FRT$	(Noster et al., 2019)
829	MvP1980	$\Delta sseF::FRT$	(Noster et al., 2019)
830	MvP503	$\Delta sifA::FRT$	(Namakchian et al., 2018)
831	MvP2600	$\Delta dksA::FRT$	(Schulte et al., 2020)

832 Table S 3. Oligonucleotides used in this study

833	<u>Designation</u>	<u>Sequence 5'-3'</u>
834		
835	for Gibson assembly cloning	
836	Vf-p5084 (2)	AAGGAGATGGCGCCCAACAGTC
837	Vr-p5084 (2)	CGGGAAAGAGGGAGAAAAGTATGGC
838	1f-p5084-PtetA (2)	GTTGGGCGCCATCTCCTTGCTTTAAGACCCACTT
839	1r-PtetA-p5084 (2)	CATACTTTCTCCTCTTCCCGTTCACTTTCTCTATCACTGA
840	Vf-p4928	GGCCGCAGAGAATATAAAAAGC
841	Vr-p4928	CGATATCGGTACCGGGAAAGAG
842	1f-p4928-Para	GCTTTTATATTCTCTGCGGCCATCGATGCATAATGTGCCTGTC
843	1r-Para-p4928	CTTTCCCGTACCGATATCGACCATGGTGAATTCCCTCCT
844	Vf-sfGFP	ATGCGCAAAGGCGAAGAACTGT
845	Vf-pMW211-V2	ATGGCATTCCACCGAGGAC
846	1f-p5084-PBAD	GTCCTCGGTGGATGCCATACTTTCTCCTCTTCCC
847	1r-tetA-p5084	GTTCTCGCCTTGCGCATGATATCCTCCTTGCCATCT
848		
849	Check primer	
850	PtetA-p5084-Check-For	TCGCCCTCGATCTCGAACTC
851	PtetA-p5084-Check-Rev	GAATAAGAAGGCTGGCTCTG
852	Para-p4928 Check-For	CCGACCGCTGGGAATGAAAG
853	Para-p4928 Check-Rev	AAGGTGCAGAGCCAGCCTTC
854	dsred-p5419 Check-For	GCACTTGAAGCGCATGAAC
855	dsred-p5419 Check-Rev	GGTCGGCAAACAAATTCTCG

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857 Suppl. Figures and Figure Legends



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859 **Fig. S 1: A dual fluorescence reporter suitable to detect the non-replicating subpopulation**
860 **of intracellular STM.** A) STM WT harboring dual fluorescence reporter with AHT-inducible
861 *dsred* expression and *msrA::sfgfp* was grown in LB medium o/n and diluted in fresh LB medium
862 containing AHT for further subculture. After 4 h of subculture, AHT was removed by
863 centrifugation and washing. Subsequently, bacteria were subcultured in fresh LB without AHT.
864 At indicated time points samples were taken, fixed and subjected to FC. The x-median
865 represents the AHT-induced DsRed intensity of the entire bacterial population, and the
866 respective histograms (B). C) STM WT harboring AHT-inducible or constitutive dual
867 fluorescence reporter for *msrA* was grown o/n in LB medium in the presence of AHT if
868 necessary. Before infection, AHT was removed. RAW264.7 macrophages were infected, and
869 lysed 8 h p.i. Liberated STM were recovered, fixed and subjected to FC. As negative control,
870 AHT was omitted from o/n cultures. The histogram shows that intracellular NR STM can be
871 detected inside RAW264.7 macrophages showing the same DsRed intensity compared to
872 intracellular STM WT harboring the constitutive dual fluorescence reporter for *msrA*. If AHT
873 was omitted, no DsRed-positive STM can be detected. D) NR subpopulations of intracellular
874 STM WT can be detected after various time points post infection. The same experiments were
875 performed for the AHT-inducible dual fluorescence reporter for *trxA* and *htrA*, with the same
876 outcome (data not shown). E-H) Detection accuracy of bacterial particles on an Attune NxT
877 cytometer. STM WT strains constitutively expressing *dsred* or *sfgfp* were grown in LB medium
878 o/n. E) Controls of PBS without STM, only DsRed-positive STM, or only sfGFP-positive STM
879 are shown. Events per μ l are indicated in each gate. Then, mixed ratios of DsRed- and sfGFP-
880 expressing STM were prepared in PBS. Either a constant high amount of sfGFP-positive STM
881 mixed with an equal, 10, 100, 1,000, or 10,000-fold reduced amount of DsRed-positive STM
882 (F), or a vice versa (G) was prepared and directly subjected to FC. H) The relative amount
883 (events $\times \mu$ l $^{-1}$) of sfGFP- and DsRed-positive STM measured in the first sample was set to 100%
884 (10 0 /10 0). Mixtures with gradual 10-fold reduction of red or green fluorescent STM within a

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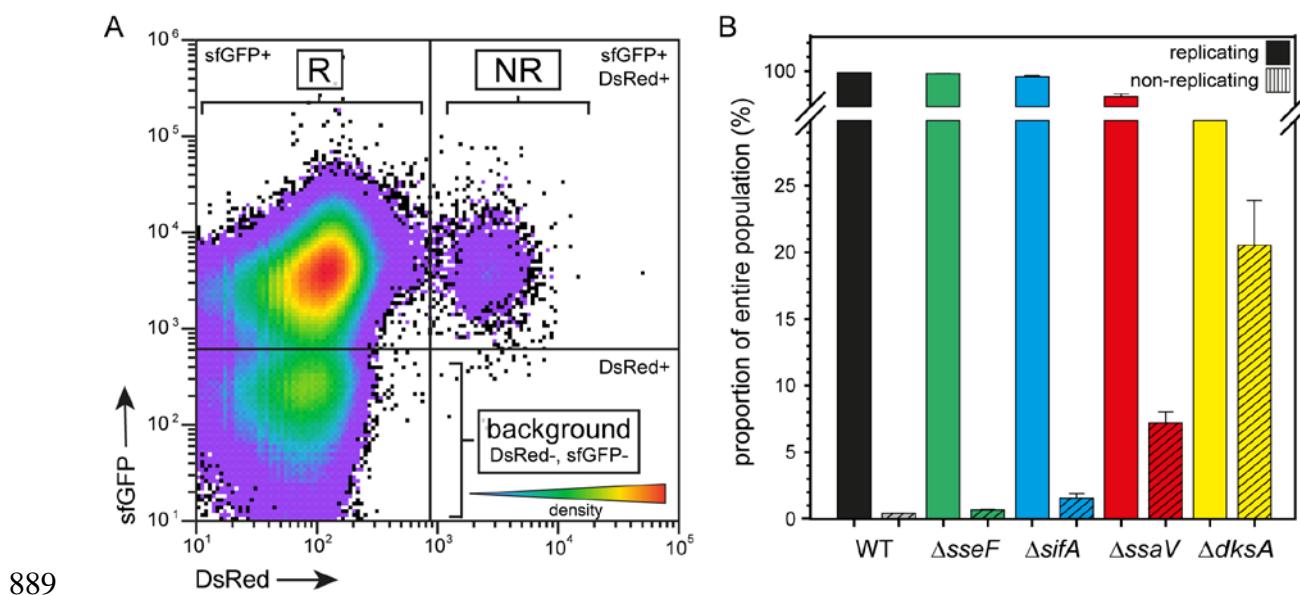
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885 sample containing constant high amount of green or red fluorescent STM, respectively, were
886 quantified. The measured value (events $\times \mu\text{l}^{-1}$ in %) compared to the first sample containing
887 high amounts of both, red and green fluorescent STM, is indicated below.

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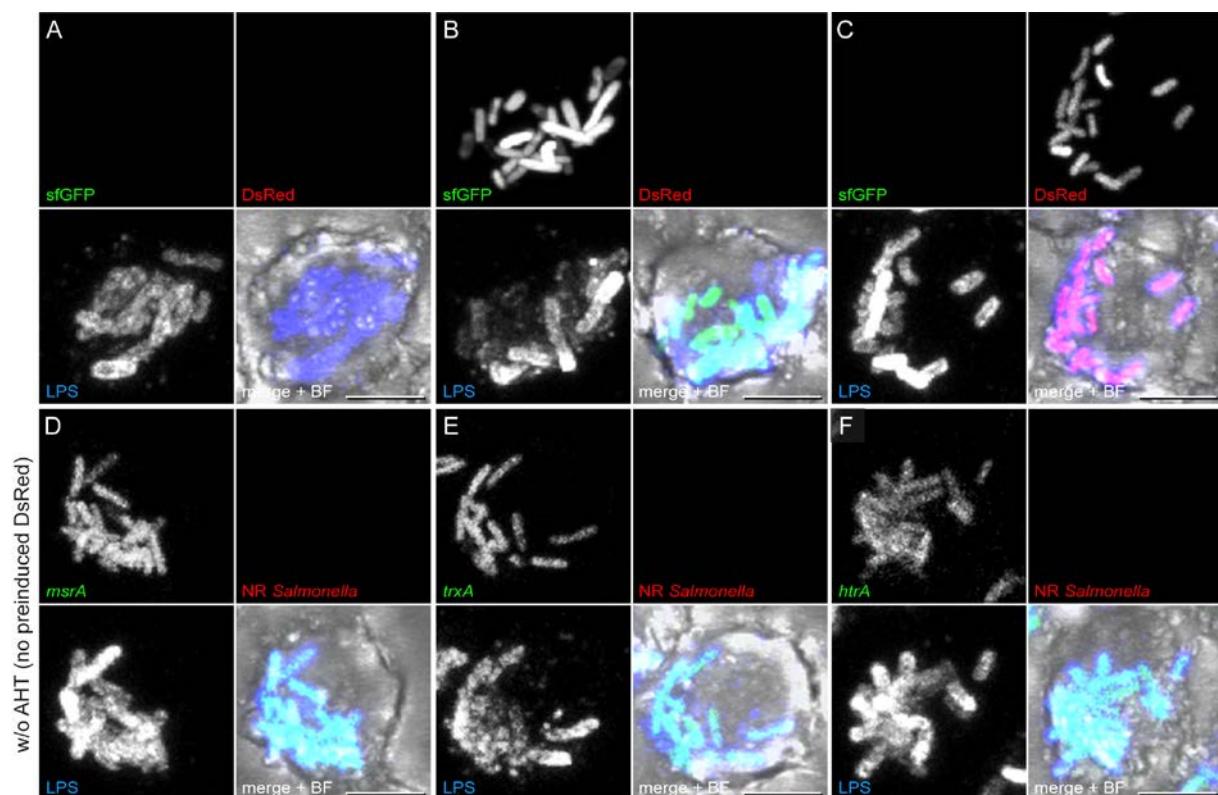
890 **Fig. S 2: A minor fraction of the intracellular population consists of non-replicating STM.**

891 STM WT, $\Delta sseF$, $\Delta sifA$, $\Delta ssaV$ and $\Delta dksA$ harboring AHT-inducible dual fluorescence reporter
892 for *msrA* were grown o/n in LB medium in the presence of AHT. AHT was removed prior to
893 infection. RAW264.7 macrophages were infected and lysed 24 h p.i. STM released from host
894 cells were recovered, fixed and subjected to FC. A) Analyses of the entire intracellular bacterial
895 population for AHT-induced DsRed and *msrA*-induced sfGFP intensity shows three different
896 populations. DsRed-negative (DsRed-) and sfGFP-negative (sfGFP-) events represent the
897 background signal consisting of host cell debris. DsRed- and sfGFP+ events represent the
898 replicating (R) subpopulation of intracellular STM and DsRed+ and sfGFP+ events represent
899 the non-replicating (NR) subpopulation. B) Gating on, and quantification of R and NR
900 subpopulations indicated the small size of the population of NR intracellular STM WT.

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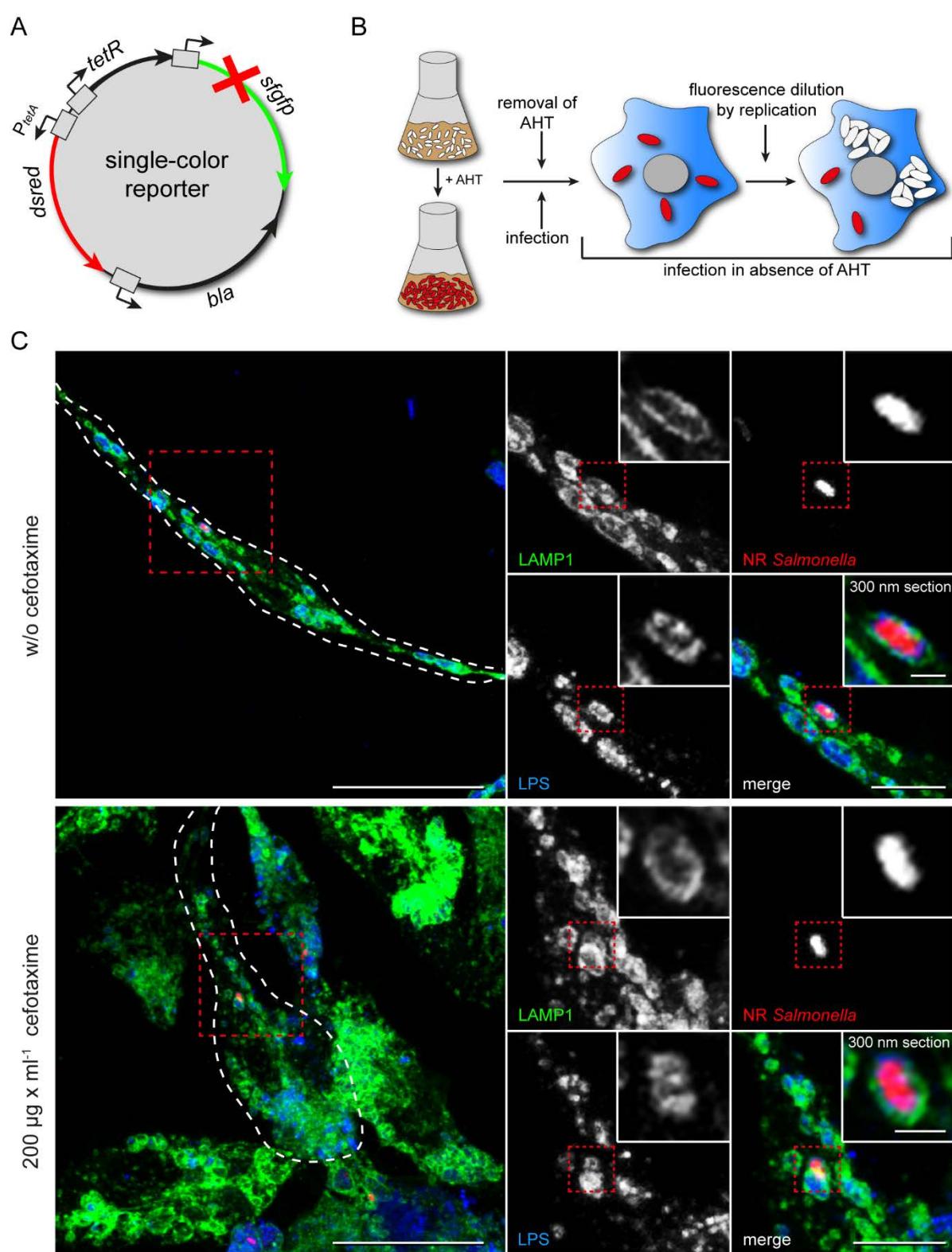


902

903 **Fig. S 3: Functional control of AHT-inducible dual fluorescence reporter without AHT**
904 **induction. A-F)** STM WT harboring AHT-inducible (NR) dual fluorescence reporter for *msrA*,
905 *trxA*, or *htrA* were grown o/n in LB medium without AHT. RAW264.7 macrophages were
906 infected and fixed 8 h p.i. for fluorescence microscopy. STM were immuno-stained against
907 O antigen (blue). As fluorescence controls, STM WT without expression of any FP (A), STM
908 WT constitutively expressing *sfGfp* (B), and STM WT constitutively expressing *dsred* (C) were
909 used. Induction of P_{*msrA*} (D), P_{*trxA*} (E) or P_{*htrA*} (F) is shown in green. NR STM were not detected
910 due to absence of AHT in o/n culture. Representative cells are shown. Scale bar, 5 μ m.

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912 **Fig. S 4: Non-replicating persisters reside inside SCV in infected host cells.** A) For the
913 analysis of NR intracellular STM, the EM7 promoter of the non-induced *sfgfp* frameshift
914 control plasmid was replaced by the tet-ON cassette to be able to experimentally induce the
915 expression of *dsred* by AHT. B) When adding AHT to a culture of STM growing in LB

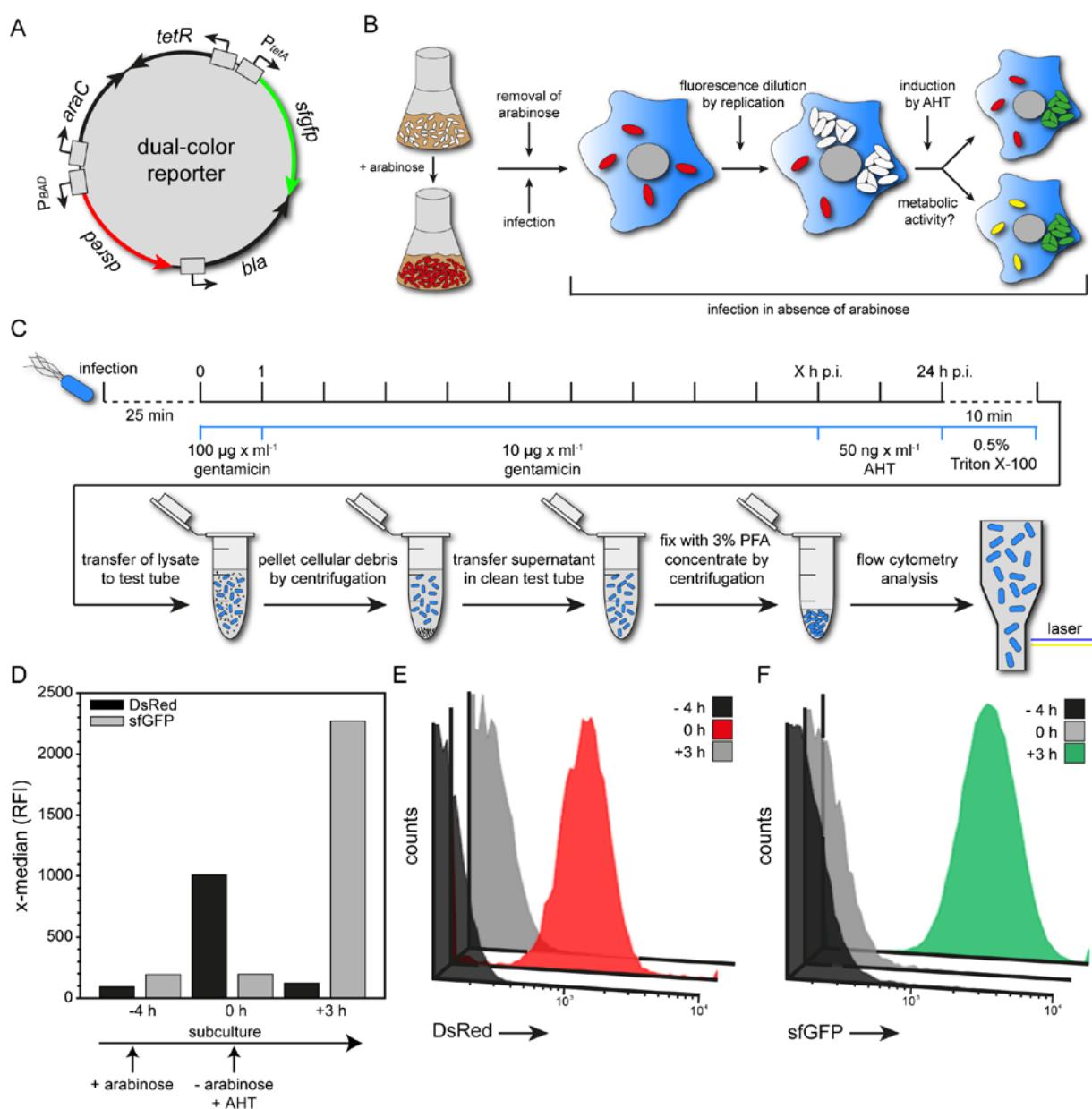
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916 medium, bacteria start to synthesize DsRed. Before infection of macrophages, AHT is removed
917 by centrifugation and washing. Infection is performed without addition of AHT to the cell
918 culture medium. R intracellular STM lose DsRed via fluorescence dilution. NR intracellular
919 STM keep the DsRed inside the cytoplasm and can be detected as DsRed-positive within e.g.
920 RAW264.7 LAMP1-GFP macrophages during microscopy. C) STM WT harboring
921 AHT-inducible single FP reporter was grown o/n in LB medium in the presence of arabinose.
922 Before infection, AHT was removed. RAW264.7 LAMP1-GFP macrophages were infected, 10
923 h p.i. cefotaxime was added to the cells if indicated and fixed 24 h p.i. for fluorescence
924 microscopy. Fixed intracellular STM were immuno-stained against O antigen (blue). LAMP1
925 is shown in green, NR STM show and R STM do not show red fluorescence. Representative
926 cells are shown. Scale bars, 20, 5, 1 μ m in overview, details, and zoom-in, respectively.

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928 **Fig. S 5: Dual fluorescence reporters for analyses of metabolic activity of intracellular NR**

929 **STM.** A) For the measurement of metabolic activity of NR intracellular STM the EM7

930 promoter and the *msrA* promoter of the dual fluorescence reporter for *msrA* was replaced by the

931 arabinose-inducible promoter cassette (Guzman et al., 1995), and the tet-ON cassette to be able

932 to artificially induce the expression of *dsred* by arabinose, and *sfgfp* by AHT. B) When adding

933 arabinose to a growing culture of STM in LB medium, bacteria start to synthesize DsRed.

934 Before infection of macrophages, arabinose is removed by centrifugation and washing.

935 Infection is performed without addition of arabinose to the cell culture medium. Replicating

936 intracellular STM lose DsRed via fluorescence dilution. NR intracellular STM keep the DsRed

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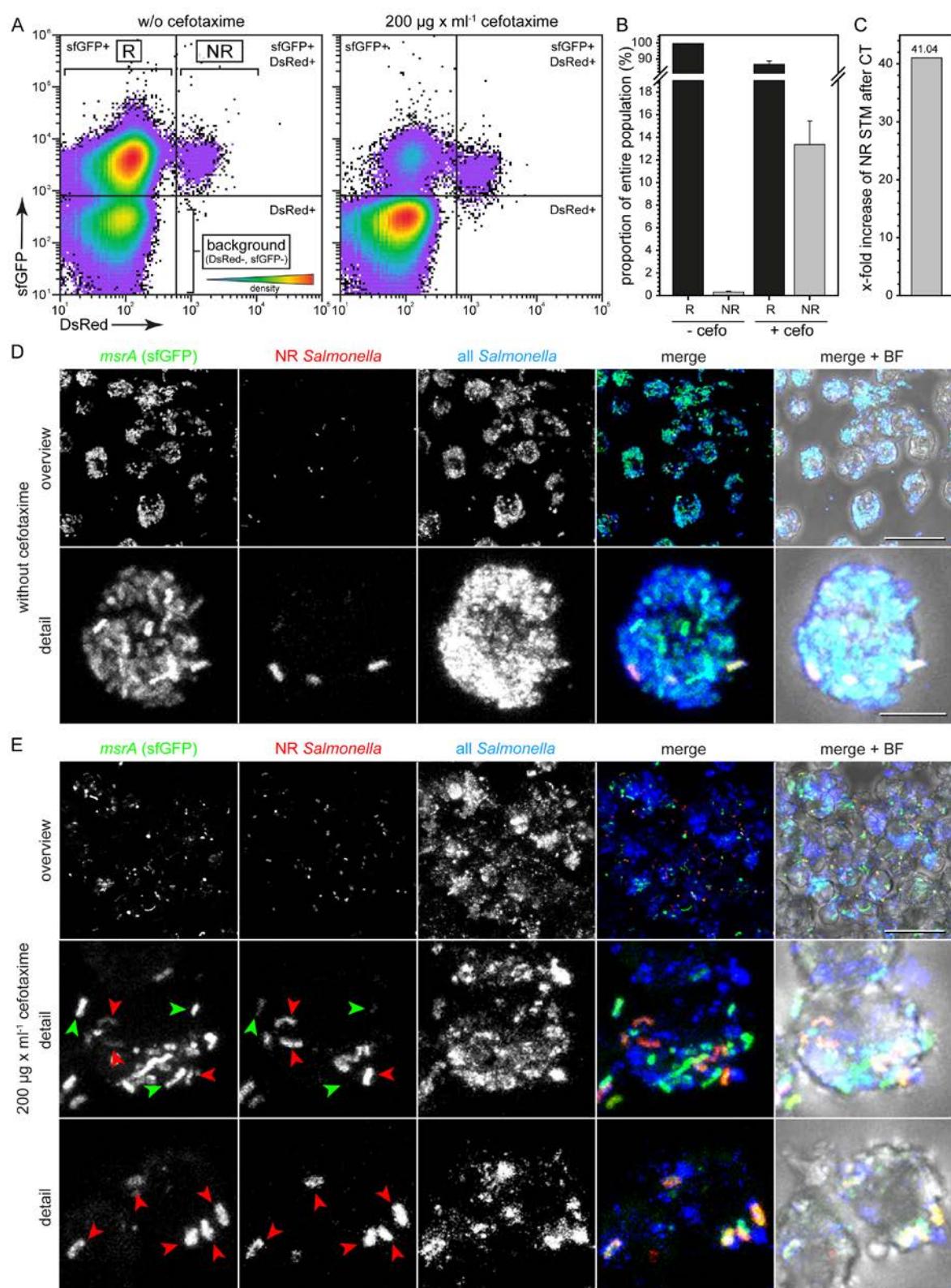
Stress response of *Salmonella* persisters

937 inside the cytoplasm and can be detected as DsRed-positive event after infection. In addition,
938 the metabolic activity can be determined by addition of AHT resulting in synthesis of sfGFP of
939 metabolically active bacteria. C) RAW264.7 macrophages were infected and all
940 non-phagocytosed STM were killed by addition of gentamicin. At indicated time points AHT
941 was added to the cell culture medium. Host cells were lysed 24 h p.i. After removal of cell
942 debris, liberated STM were fixed and subjected to FC. D-F) STM WT harboring double-
943 inducible dual fluorescence reporter was grown in LB medium o/n and diluted in fresh LB
944 medium containing arabinose for further subculture (-4 h). After subculture for 4 h, arabinose
945 was removed by centrifugation and washing (0 h). Subsequently, bacteria were further
946 subcultured for 3 h in fresh LB without arabinose, but containing AHT (+3 h). Samples were
947 taken, fixed and subjected to FC at indicated time points. D) The x-median represents the
948 arabinose-/AHT-induced DsRed/sfGFP intensity of the entire bacterial population. Respective
949 histograms for DsRed intensity (E), and sfGFP intensity (F) are shown.

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953 **Fig. S 6: Cefotaxime treatment increases intracellular population of non-replicating STM**

954 WT. STM WT harboring AHT-inducible (NR) dual fluorescence reporter for msrA was grown

955 o/n in LB medium in the presence of AHT. Before infection, AHT was removed. RAW264.7

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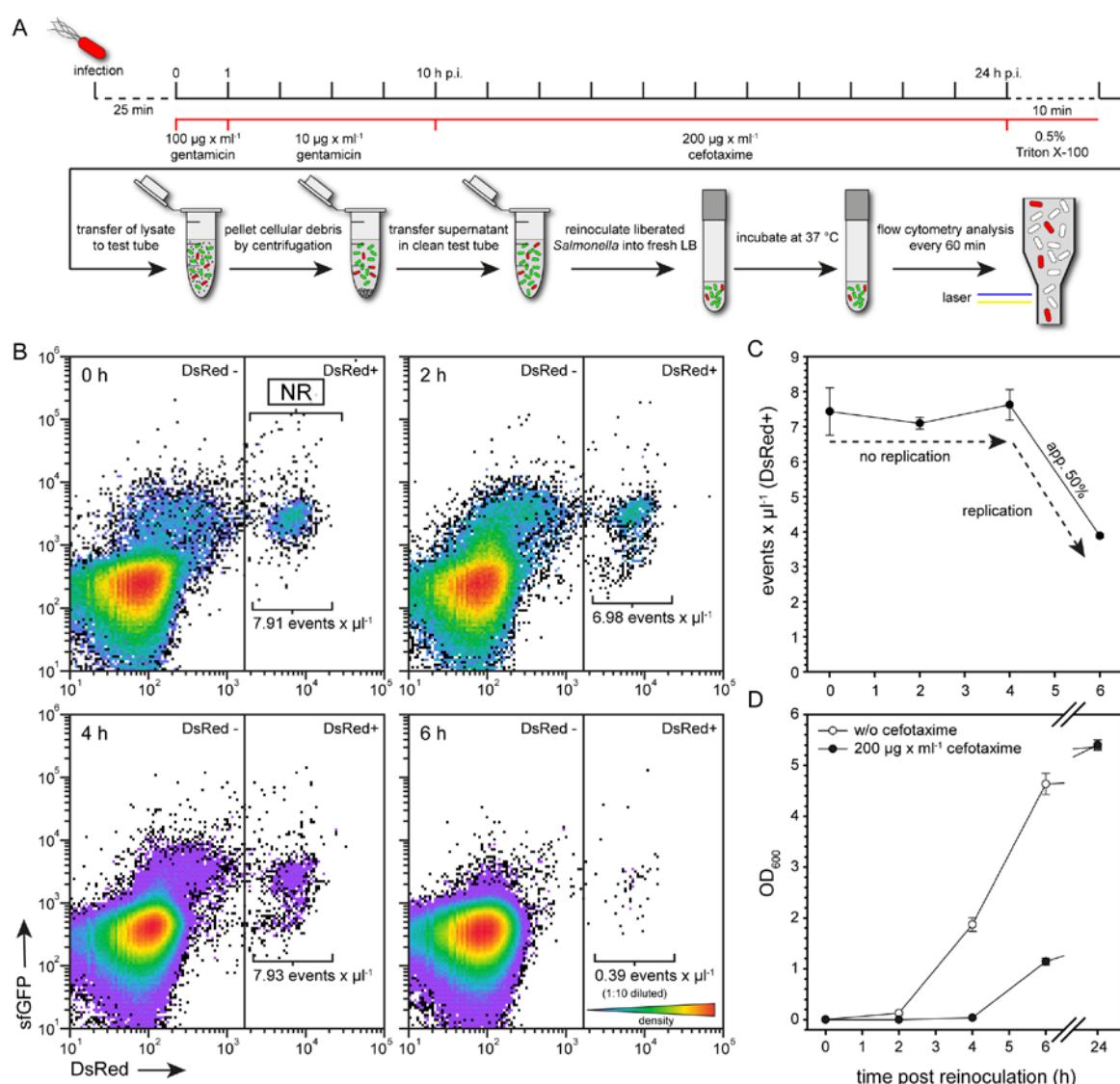
Stress response of *Salmonella* persisters

956 macrophages were infected, cefotaxime was added 10 h p.i. to the cells if indicated, and cells
957 were lysed for 24 h p.i. FC analyses, or fixed for fluorescence microscopy. Liberated STM were
958 recovered, fixed and subjected to FC. A) Plotting the entire intracellular bacterial population
959 against their AHT-induced DsRed and P_{msrA} -induced sfGFP intensity shows a high reduction
960 of R STM after antibiotic treatment. B) Quantification of the proportion of R and NR STM
961 within the population with and without antibiotic treatment. C) Calculated x-fold increase of
962 the proportion of NR compared to R STM after cefotaxime treatment (CT). Means and standard
963 deviations of one representative experiment are shown. D, E) Microscopy of intracellular R and
964 NR STM without (D), and with cefotaxime treatment (E). Fixed intracellular STM were
965 immuno-stained against O antigen (blue). Induction of *msrA* is shown in green, NR STM show
966 and R STM do not show red fluorescence. Merge micrographs with and without overlay of
967 brightfield (BF) channels are shown. NR STM are indicated by red arrowheads, R STM are
968 indicated by green arrowheads. Representative cells are shown. Scale bars, 20 μ m and 5 μ m in
969 overview and detail, respectively.

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Stress response of *Salmonella* persisters



971

972 **Fig. S 7: Schematic overview of infection and lysis of host cells, reinoculation into fresh**
973 **LB, and subsequent cytometric analysis. A)** RAW264.7 macrophages were infected using
974 STM harboring various reporters. After killing of all non-phagocytosed STM by addition of
975 gentamicin, 10 h p.i. cell culture medium was replaced by medium containing 200 $\mu\text{g} \times \text{ml}^{-1}$
976 cefotaxime. After incubation for further 14 h, host cells were lysed 24 h p.i. After removal of
977 cell debris, liberated STM were reinoculated into fresh LB medium and incubated at 37 °C. At
978 indicated time points samples were taken, fixed and subjected to FC. Quantification of DsRed-
979 positive STM was performed using an Attune NxT cytometer. B) STM WT harboring

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Stress response of *Salmonella* persisters

980 AHT-inducible (NR) dual fluorescence reporter for *msrA* was grown o/n in LB medium in the
981 presence of AHT. Before infection, AHT was removed. RAW264.7 macrophages were
982 infected, 10 h p.i. cefotaxime was added to the cells and then lysed 24 h p.i. Liberated STM
983 were recovered, reinoculated into fresh LB and incubated at 37 °C. At indicated time points
984 samples were taken, fixed and subjected to FC. C) The relative amount (events x μ l⁻¹) of DsRed-
985 positive NR STM in the culture was determined. A loss of DsRed-positive events over time
986 indicates that NR STM are growth-competent and start to replicate. D) In addition, the optical
987 density was measured. Means and standard deviations of one representative experiment are
988 shown.