

1 **Bacterial host interaction of GFP-labelled *Listonella anguillarum* HI-610 with**  
2 **gnotobiotic sea bass (*Dicentrarchus labrax* L.) larvae**

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24 Running title: bacterial interaction in gnotobiotic larval gut

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**Abstract**

The location and cell damage caused by *Vibrio anguillarum*, the causative agent of classical vibriosis, within the developing gut of the newly hatched sea bass (*Dicentrarchus labrax* L.) is unknown. A gnotobiotic sea bass model was used to investigate the early interactions of *V. anguillarum* with sea bass larvae. In the present study, germ-free sea bass larvae were orally exposed to a *V. anguillarum* HI-610 pathogen labelled with the Green Fluorescent Protein (GFP-HI-610) and sampled at regular intervals. Pathogenic colonisation of gut enterocytes was observed 2 hrs post exposure (p.e.) and onwards, whereas bacteria within the swim bladder were visualized 48 hrs p.e and onwards. Ultrastructural findings demonstrated direct bacterial contact with the host cell in the oesophageal mucosa and putative attachment to microvilli of mid- and hindgut enterocytes. The present findings form a starting point for studies assessing the impact of potential candidates (probiotics, prebiotics, antimicrobial peptides) to mitigate bacterial virulence.

*Keywords:* larviculture, gnotobiotic model system, GFP-labelled pathogen, microscopy, ultrastructure

## 45 **Introduction**

46 Knowledge of the location of adhesion and/or translocation of pathogenic bacteria to and  
47 within the enterocytes of various gut segments in fish larvae is of utmost importance in the  
48 design of mitigation strategies. However, the diversity of micro-organisms colonising the gut  
49 in the early larval stages is hindering the study of the cause-effect relationship in microbial  
50 larval interactions. Using a gnotobiotic model (Dierckens, Rekecki, Laureau, Sorgeloos,  
51 Boon, Van den Broeck & Bossier 2009) starting with germ-free larvae, and hence eliminating  
52 indigenous microbiota, may provide an excellent tool to unravel the modes-of-action of host  
53 microbe interactions (Gunasekara, Rekecki, Baruah, Bossier & Van den Broeck 2010).  
54 Recently we described the zootechnical aspects of a standardised challenge test with the  
55 pathogen *Vibrio anguillarum* HI-610 (Dierckens *et al.* 2009). However, the invasion strategy  
56 of this pathogen is unknown in the developing gut of sea bass larvae as the location of and  
57 cell damage caused by *V. anguillarum* has not been investigated yet. In the larviculture  
58 industry, exogenous feeding of sea bass larvae with *Artemia* start at day 8-9 after hatching  
59 (AH) (Chatain 1997). This initial period without food is considered as a stress inducing factor.  
60 Furthermore, bacteria are taken up in the gut before fish larvae switch from endogenous to  
61 exogenous feeding regime (Ringø & Birckbeck 1999). By using the gnotobiotic model, sea  
62 bass larvae are feed-deprived until day 7 AH. During the first two weeks AH, germ-free sea  
63 bass larvae do not possess mucus producing goblet cells (Rekecki, Dierckens, Laureau, Boon,  
64 Bossier & Van den Broeck 2009). As a consequence, the mucus layer, normally acting as a  
65 physical barrier in the gut is lacking, except for in the pharyngeal and oesophageal regions  
66 (Rekecki *et al.* 2009) whereby the mid- and hindgut are exposed and may become more  
67 vulnerable to bacteria at this stage. It is suggested that the absence of indigenous bacteria by  
68 shedding of epithelial cells in stressed fish and the lack of protective mucus might have a  
69 significant role in pathogenesis as the pathogenic bacteria can then attach to cell surfaces,

70 translocate through enterocytes and cause disease (Ringø, Myklebust, Mayhew & Olsen  
71 2007).

72 *Vibrio* appears to interact differently with different host species as described by Sandlund,  
73 Rødseth, Knappskog, Fiksdal & Bergh (2010). *V. anguillarum* serotype O2 $\alpha$  (HI-610) was  
74 detected in the GI tract, abdominal cavity, gallbladder and kidney of turbot (*Scophthalmus*  
75 *maximus*) larvae. In Atlantic cod (*Gadus morhua*) larvae, this bacterium was visualized only  
76 in the GI tract, whereas necrosis of dermal cells and the presence of these bacteria in sensory  
77 cells of the head region were observed in Atlantic halibut (*Hippoglossus hippoglossus*) larvae.  
78 In another study, a high mortality but little histopathology was observed in cod larvae  
79 challenged with the HI-610 strain; therefore it is supposed that bacterial toxins may have a  
80 vital role in pathogenesis (Engelsen, Sandlund, Fiksdal & Bergh 2008).

81 The aim of the present study was to define the presence and location of pathogenic *V.*  
82 *anguillarum* strain HI-610 in gnotobiotic sea bass larvae at different time intervals after  
83 experimental infection using fluorescence, confocal, light and electron microscopy.

84

## 85 **Materials and methods**

### 86 **Bacterial strains and media**

87 Insertion of the Green Fluorescent Protein (GFP) gene into GFP-HI-610 transconjugants was  
88 obtained by conjugation between rifampicin resistant *V. anguillarum* HI-610, originally  
89 isolated from cod (Samuelsen, Bergh & Ervik 2003) and *Escherichia coli* DH5 $\alpha$  carrying the  
90 suicide plasmide pJBA120 containing a mini-Tn5 transposon with the *gfp* gene (Andersen,  
91 Sternberg, Poulsen, Bjørn, Givskov & Molin 1998) and subsequent selection on Luria Broth  
92 agar containing rifampicin and kanamycin as described by Boon, Goris, De Vos, Verstraete &  
93 Top (2001). *V. anguillarum* GFP-HI-610 and wild type *V. anguillarum* WT-HI-610 were  
94 grown as described by Dierckens *et al.* (2009). To determine the growth curve of strain GFP-

95 HI-610 and strain WT-HI-610, batch cultures were made by inoculating 2 day-old cultures  
96 into 10 ml of liquid media (MB) in 20 ml plastic centrifugation tubes and incubating at 16°C  
97 with agitation. Every 3 hrs, bacterial growth was checked by aseptically removing 1 ml of the  
98 aliquot and measuring the absorbance at a wavelength of 550 nm with a spectrophotometer  
99 (Genesys 20, Thermospectronic, Aalst, Belgium).

100

### 101 **Experimental set-up**

102 Eggs of *Dicentrarchus labrax* were obtained from natural spawning at the hatchery of  
103 Ecloserie Marine de Gravelines (Gravelines, France). To obtain germ-free eggs, a standard  
104 disinfection protocol was used previously described by Dierckens *et al.* (2009). After hatching  
105 (60 hrs), a transparent sterile screw cap vial (total volume: 15 ml) was filled with 10 ml  
106 filtered, autoclaved sea water (FASW) containing 10 mg rifampicin l<sup>-1</sup>. Twelve larvae were  
107 transferred to each vial. A total amount of 100 vials was placed on a rotor turning at 4 rpm  
108 with an axis tangential to the axis of the vials. Throughout the experiment, eggs and larvae  
109 were kept in a temperature controlled room at 16 ± 0.5°C in constant dim light (10 candela  
110 steradian m<sup>-2</sup>) at a salinity of 36 g l<sup>-1</sup>. From day 7 AH until day 13, thirty freshly hatched  
111 germ-free *Artemia franciscana* nauplii (Marques, François, Dhont, Bossier & Sorgeloos  
112 2004) were added to each vial (± 2.5 nauplii/larva), every second day. The experiment was  
113 approved by the ethical committee of Ghent University under the number EC2005/95.

114

### 115 **Exposure study**

116 Two trials were carried out in this study: trial 1 was performed according to the study of  
117 Dierckens *et al.* (2009). A standard suspension of 10<sup>5</sup> cfu *V. anguillarum* ml<sup>-1</sup> was  
118 administered via immersion to day 3 AH larvae in order to test the virulence of WT-HI-610  
119 and GFP-HI-610 *Vibrio anguillarum*. On day 7, 9, 11 and 13 AH (4 days to 10 days post

120 exposure; p.e.), 5 replicate vials (60 larvae) were used for survival counts, and for  
121 fluorescence, confocal and light microscopy. In trial 2, day 4 AH larvae were exposed to a  
122 density of  $10^8$  cfu ml<sup>-1</sup> GFP-HI-610 in order to track early pathogenic invasion of the gut.  
123 Whole larval bodies were sampled for fluorescence, confocal, light microscopy and TEM at  
124 2-6-12-24-48-72-120-168 hrs p.e. (2 hrs to 7 days p.e. until day 11 AH). Day 4 AH larvae  
125 were used to ensure possible uptake of bacteria by drinking as their mouth is supposed to be  
126 already open from day 3 AH, as the bacteria were tracked already after 2 hrs p.e. For both  
127 trials, control germ-free larvae were kept under similar conditions and sampled at the same  
128 time points as challenged larvae. All larvae were euthanized by 0.1% benzocain immersion  
129 before sampling except when mentioned otherwise.

130

### 131 **Testing germ-free status and quantification of bacterial density inside the fish larvae**

132 After 24 hrs of incubation, 30 sea bass eggs were aseptically removed from the incubation  
133 bottles and processed as described by Rekecki *et al.* (2009). In addition, 1 ml water from each  
134 incubation bottle and from all sampled vials was added to a tube containing 9 ml of 10%  
135 sterile MB from the first time point until the end of the experiment. No colony forming  
136 bacteria and no turbidity were detected after 72 hours incubation of eggs and larvae during the  
137 course of the trial.

138 In order to determine the density of gut microbiota in sea bass larvae in trial 1, live challenged  
139 and control germ-free larvae were anesthetized with 0.1% benzocain for 10 s. After surface  
140 disinfection with 0.1% benzalconium chloride for 10 s the larvae were washed 3 times in  
141 sterile NSS for 10 s, before being homogenized and plated on 10% Marine Agar (MA) with  
142 10 mg l<sup>-1</sup> rifampicin. Agar plates and the marine broth tubes were incubated at an increasing  
143 temperature from 20°C to 28°C over a 72 hrs period and were checked for bacterial growth.

144

## 145 **Sampling procedure and microscopy**

146 Fluorescence and confocal microscopy: for every time point, 5 challenged larvae were  
147 anesthetized with 0.1% benzocain for 10 s, the external body surface was disinfected with  
148 0.1% benzalconium chloride for 10 s, and washed in FASW for 10 s, before fixation in 4%  
149 paraformaldehyde for 2 hrs. The whole larval body was individually mounted on glass slides  
150 with a solution of glycerine and 1,4-diaza-bicyclo[2,2,2]-octane (DABCO) (ACROS  
151 Organics, Geel, Belgium). To examine bacterial colonization of the skin, 3 additional larvae  
152 were sampled and treated at every time point in the same way but without the disinfection  
153 step. Serial optical sections of larvae were acquired using a LEICA TCS SP2 confocal  
154 microscope using the Argon 488 nm excitation laser line (Leica Microsystems Belgium  
155 BVBA, Groot-Bijgaarden, Belgium) and Olympus BX61 fluorescence microscope (Olympus  
156 Belgium N.V., Aartselaar, Belgium) using the filter cube U-MWIB2.

157

158 Light microscopy: five sea bass larvae were fixed for 3 hours in Carnoy's fixative (6 parts  
159 absolute alcohol, 3 parts chloroform, 1 part glacial acetic acid 100%) for Giemsa staining and  
160 processed and cut as described by Rekecki *et al.* (2009).

161 Transmission electron microscopy (TEM): whole larvae were fixed in Karnovsky fixative  
162 (2% formaldehyde and 2.5% glutaraldehyde in 0.1M sodium cacodylate buffer pH 7.4)  
163 overnight at 4°C, washed in sodium cacodylate buffer (pH 7.4), post-fixed overnight with 1%  
164 osmium tetroxide, dehydrated in a graded series of alcohol and embedded in SPURR's resin.  
165 Serial ultrathin sections of 60 nm ranging from the ventral border of the yolk sac up to the  
166 swim bladder of the larval GI tract were made at intervals of 7 µm, using the Leica EM UC6  
167 ultramicrotome (Leica Microsystems Belgium BVBA), mounted on formvar coated single  
168 slot copper grids (Laborimpex N.V., Brussels, Belgium) and post-stained with uranyl acetate  
169 and lead citrate (Leica EM stain, Leica Microsystems Belgium BVBA). Specimens were

170 viewed on a Jeol 1200 EXII transmission electron microscope (JEOL Europe BV, Zaventem,  
171 Belgium) at 80kV accelerating voltage. Different segments of the gut, namely oesophagus,  
172 gastric region, mid- and hindgut of sea bass larvae were investigated by TEM.

173

#### 174 **Statistics**

175 Statistical analysis was performed by means of SPSS 15.0 software. Survival was reported as  
176 mean  $\pm$  standard error of the mean (S.E.M.). Data were tested for normality and in case of  
177 normal distribution, they were subjected to one-way analysis of variance (ANOVA).  
178 Bonferroni and Tukey test was used for multiple comparisons among means in case of non-  
179 homogeneity. For non-parametric tests, Kruskal Wallis was used to compare response  
180 variables from three treatment groups. Mann-Whitney U test was used for pair-wise  
181 comparisons. Significance was accepted at  $p < 0.05$ .

182

#### 183 **RESULTS**

184 Mortality and bacterial density - trial 1

185 *In vitro*, the growth curves of both strains GFP-HI-610 and WT-HI-610 were similar (data not  
186 shown). A significantly higher mortality was demonstrated in both wild-type and germ-free  
187 challenged groups ( $76 \pm 19\%$  and  $44 \pm 6\%$ ) as compared to the control group ( $4 \pm 2\%$ ) from  
188 day 11 AH onwards. This difference in mortality became more apparent by the end of the  
189 experiment on day 13 AH, being  $100\%$ ,  $89 \pm 9\%$  and  $11 \pm 2\%$ , respectively. On day 7 AH,  
190 the bacterial density within the sea bass larvae exposed to WT-HI-610 and GFP-HI-610  
191 strains was  $6.0 \times 10^4$  cfu larva<sup>-1</sup> and  $1.8 \times 10^3$  cfu larva<sup>-1</sup>, respectively. On day 9 AH, the strain  
192 WT-HI-610 density ( $7.1 \times 10^4$  cfu larva<sup>-1</sup>) had not increased compared to day 7 AH, whereas  
193 the strain GFP-HI-610 density had increased by 1 log ( $1.0 \times 10^4$  cfu larva<sup>-1</sup>). By day 11 AH,



194 the density of both WT-HI-610 and GFP-HI-610 strains became similar ( $1.8 \times 10^5$  cfu larva<sup>-1</sup>  
195 <sup>1</sup>and  $3.4 \times 10^5$  cfu larva<sup>-1</sup>), respectively.

196

## 197 Microscopy

### 198 Fluorescence, confocal and light microscopic findings

199 In trial 1, five larvae sampled per time point were examined for the presence of green  
200 fluorescent bacteria. The first indication of bacterial adhesion in the gut and the localization of  
201 bacteria within the swim bladder occurred both from day 9 AH (6 days p.e.) onwards .

202 During trial 2, samples were taken at 2-6-12-24-48-72-120-168 hrs p.e. After 2 hrs p.e.,  
203 bacteria were localized in the oesophagus, mid- and hindgut of day 4 AH larvae. Light  
204 microscopic study revealed bacterial adhesion to gut enterocytes already at 2 hrs p.e. as  
205 confirmed by Giemsa staining, while at 48 hrs p.e. bacteria, single or grouped, were clearly  
206 seen in close contact to the apical brush border of midgut and hindgut enterocytes visualized  
207 by fluorescence and confocal microscopy (Figs. 1.1-1.2). At 12 hrs p.e., a few bacteria could  
208 be detected in the lumen of the ureter. Starting from 48 hrs p.e. onwards (day 6 AH larvae),  
209 bacteria were visible within the swim bladder (Figs. 1.3-1.4). In total, bacteria were present in  
210 the swim bladders of 10% (6 out of 60 larvae) of the investigated sea bass larvae. Bacterial  
211 presence on the skin was observed sporadically at the yolk sac and the cranial regions in 13%  
212 of larvae. During the whole experiment, the number of bacteria present on the skin never  
213 reached the initial levels of bacteria observed in the GI tract at 2 hrs p.e.. Control germ-free  
214 larvae did not contain any GFP-labelled bacteria.

### 215 Ultrastructural findings

216 Ciliated cells were observed in fore-, mid- or hindguts in both uninfected and challenged  
217 larvae. At 48 hrs p.e., numerous intraluminal bacteria were visualized in close contact with  
218 ciliated cells of the oesophageal mucosa (Fig. 2.1a). Bacterial adhesion to the oesophageal

219 epithelium characterised by electron dense area between the bacterial outer membrane and the  
220 host cell membrane was observed (Fig. 2.1b). Shedding of epithelial cells into the gut lumen  
221 was regularly seen. Bacterial-like structures within the gut lumen and close contact  
222 (interspace of 100-200 nm) of bacteria to the microvilli of midgut and hindgut enterocytes  
223 were visualized during the course of the experiment (Figs. 2.2-2.3). Bacterial clusters within  
224 the midgut lumen were surrounded by and in direct physical contact with membrane-like  
225 structures at 120 hrs p.e. (Fig. 2.4). A putative dendritic cell with phagosomes protruding with  
226 cytoplasmic extensions towards the midgut lumen was visualized at 168 hrs p.e. (Fig. 2.5). In  
227 both challenged and control animals, intraepithelial macrophages situated at the basal portions  
228 of midgut and hindgut enterocytes were visualized at 48 hrs p.e. (Fig. 2.6).

229

## 230 **DISCUSSION**

231 Although *in vitro* the growth curves of both bacterial strains were similar, *in vivo* their growth  
232 seemed to be delayed. The sea bass larvae rely only upon endogenous yolk reserves during  
233 the first days of development and hence the larval gut is a nutrient-poor environment for  
234 bacteria. In this respect, the GFP production may lead to slower bacterial growth and slightly  
235 delayed larval mortality. Incorporation of GFP can be a metabolic burden for the bacterial  
236 cell, as more energy is required for this extra protein (Allison & Sattenstall 2007). However,  
237 at day 11 AH, when the larvae were fed already for several days the bacterial density in both  
238 groups became similar. As an alternative we put forward the controversial hypothesis that the  
239 growth of strain GFP-HI-610 in the larval GI tract may be delayed due to down regulation  
240 and/or inactivation of a virulence factor due to chromosomal GFP insertion.

241 Engelsen and colleagues (2008) did not detect bacteria associated with the skin of cod larvae  
242 during immersion challenge with *V. anguillarum* HI-610. O'Toole, von Hofsten, Rosqvist,  
243 Olsson & Wolf-Watz (2004) localised GFP-labelled *V. anguillarum* firstly in the gut of

244 zebrafish larvae, followed by the fish surface. In our study, only a few bacteria were noted on  
245 the skin in 13% of larvae, thus it can be concluded that no significant colonisation of the skin  
246 was observed. An increasing number of GFP-labelled *V. anguillarum* was localised in the GI  
247 tract lumen in the present study. Already from 2 hrs p.e. onwards, putative adhesion to and  
248 colonisation of gut enterocytes was observed. At 48 hrs p.e. the presence of bacteria in the  
249 swimbladder was demonstrated by fluorescence and confocal microscopy. Sea bass larvae  
250 belong to the transient physostomous group using a functional pneumatic duct for a limited  
251 time in order to inflate their swim bladder (Chatain 1986; Kitajima, Watanabe, Tsukashima &  
252 Fujita 1994). Inflation occurs when sea bass larvae switch from endogenous to exogenous  
253 feeding, whereafter a gas gland, located ventrally in the swim bladder, will be responsible for  
254 buoyancy control (Chatain 1986; Battaglione & Talbot 1990; Trotter, Pankhurst & Battaglione  
255 2005). Immunosuppressed larvae may become vulnerable to bacterial aerocystitis, an  
256 inflammation of the swim bladder epithelium that causes dysfunction of the gas gland and  
257 collapse of a partially inflated swim bladder (Summerfelt 1996). Similar findings were  
258 reported in walleye (*Stizostedion vitreum*) larvae (Kindschi & MacConnell 1989; Marty,  
259 Hinton & Summerfelt 1995) and in zebrafish (*Danio rerio*) larvae (Kent, Matthews, Laver &  
260 Schech 2007). Inflation failure is considered to be irreversible, forming one of the major  
261 bottlenecks in larviculture as buoyancy control, swimming ability and feeding success  
262 become jeopardized (Peruzzi, Westgaard & Chatain 2007). It needs to be taken into account  
263 that in commercial hatcheries, both allochthonous and autochthonous gut bacteria, can cause  
264 disease under certain conditions (Ringø *et al.* 2007) and can also enter and cause collapse and  
265 noninflation of the swim bladder, eventually leading to starvation and death. Therefore,  
266 application of probiotics and their putative beneficial effect as ecological competitor in the gut  
267 may be questioned when larval survival is endangered by bacterial aerocystitis.

268 Ciliated cells were observed in the different segments of the GI tract in both challenged and  
269 control germ-free larvae of different ages. In the midgut of two weeks old herring (*Clupea*  
270 *harengus*) larvae, ciliated cells have been described which are supposed to propel bacteria  
271 towards the hindgut where antigen uptake occurs (Hansen, Strøm & Olafsen 1992). The  
272 function of ciliated cells could be a cleansing mechanism, preventing organisms to make  
273 contact with binding sites on mucosal surfaces. Bacteria must adhere first to intestinal mucosa  
274 in order to withstand this action (Lu & Walker 2001). In our study, bacterial adhesion to the  
275 oesophageal mucosa was demonstrated at 48 hrs p.e.. Electron dense contact between the  
276 bacterial outer membrane and an invaginated host cell membrane is considered to be an  
277 attachment site (Neutra 1984). Next to avoiding cleansing mechanisms, adhesion is also  
278 important for infection resulting in other benefits for the bacteria such as growth advantages,  
279 enhanced toxicity to the host, and increased resistance against antimicrobial or antitoxin  
280 activities (Ofek & Doyle 1994). Our study additionally demonstrated scattered bacteria in the  
281 lumen of mid- and hindgut enterocytes in close contact with the brush border layer (Fig. 2.4).  
282 Whether this implies that bacteria attached to the brush border effectively enter the fish by  
283 this intestinal route, is not certain as suggested by Engelsen *et al.* (2008).

284 In general, the hindgut is considered to be mainly involved in phagocytotic activity and less  
285 involved in nutrient absorption (Buddington & Diamond 1987). In both the challenged and  
286 germ-free groups, engulfment of particles, but not of bacteria was observed in both mid- and  
287 hindgut intraepithelial macrophages. The exact nature of these particles hitherto is not clear.

288 In *in vivo* studies with *V. anguillarum*, the pathogen did not penetrate turbot juvenile  
289 epithelial cells nor was it taken up intracellularly (Olsson, Jöborn, Westerdahl, Blomberg,  
290 Kjelleberg & Conway 1996). In contrast, immunohistochemical studies evidenced a release of  
291 free *V. anguillarum* from an enterocyte endosome into the lamina propria in turbot larvae  
292 (Grisez, Chair, Sorgeloos, & Ollevier 1996). In epithelial cell lines, *V. anguillarum* strains can

293 survive intracellularly as a result of uptake by phagocytosis or endocytosis induced by  
294 bacteria (Wang, Oon, Ho, Wong, Lim & Leung 1998). In a preliminary TEM study on  
295 conventional sea bass juveniles, internalized bacterial colonies within the gut enterocytes  
296 were described (Rekecki, unpublished results). However, in the current study, no intra- or  
297 intercellular bacterial translocation from the GI tract was observed. We observed expulsion of  
298 enterocytes into the gut lumen which can be either due to natural shedding of gut enterocytes  
299 needed for tissue homeostasis, or a major host defense mechanism of the gut epithelium. The  
300 latter involves rapid exfoliation of damaged cells infected by bacterial pathogens in order to  
301 prevent bacterial colonisation (Kim, Ashida, Ogawa, Yoshikawa, Mimuro & Sasakawa 2010).  
302 Whether these cells in the gut lumen are part of the hosts defense mechanism, is unknown.  
303 In conclusion, by using the established sea bass model and by labelling the pathogenic  
304 bacterium with GFP, we managed to study bacteria *in situ* during early colonisation of the  
305 host. The present findings provide a baseline for investigating the effects of probiotica (e.g.  
306 quorum sensing inhibitors), antimicrobial peptides, prebiotica, immunostimulants and other  
307 substances affecting virulence factors, hence colonisation and infection. Finally, whether  
308 probiotics can exert a protective role in fish larvae once their swim bladder is infected, is a  
309 topic for further research.

310

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