

# External pH modulation during the growth of Vibrio tapetis , the aetiological agent of brown ring disease

A. Rahmani, C. Mathien, Adeline Bidault, Nelly Le Goïc, V. Pichereau,

Christine Paillard

### ▶ To cite this version:

A. Rahmani, C. Mathien, Adeline Bidault, Nelly Le Goïc, V. Pichereau, et al.. External pH modulation during the growth of Vibrio tapetis , the aetiological agent of brown ring disease. Journal of Applied Microbiology, 2020, 129 (1), pp.3-16. 10.1111/jam.14699 . hal-03083014

### HAL Id: hal-03083014 https://hal.science/hal-03083014

Submitted on 4 Jan 2021

**HAL** is a multi-disciplinary open access archive for the deposit and dissemination of scientific research documents, whether they are published or not. The documents may come from teaching and research institutions in France or abroad, or from public or private research centers. L'archive ouverte pluridisciplinaire **HAL**, est destinée au dépôt et à la diffusion de documents scientifiques de niveau recherche, publiés ou non, émanant des établissements d'enseignement et de recherche français ou étrangers, des laboratoires publics ou privés.

1	External pH modulation during the growth of Vibrio tapetis, the etiological agent of							
2	Brown Ring Disease							
3								
4	Running head: External pH modulations by Vibrio tapetis							
5								
6	Alexandra Rahmani <sup>1#</sup> , Clémentine Mathien <sup>1</sup> , Adeline Bidault <sup>1</sup> , Nelly Le Goïc <sup>1</sup> , Christine							
7	Paillard <sup>1#†</sup> , Vianney Pichereau <sup>1#†</sup>							
8	<sup>1</sup> Univ Brest, CNRS, IRD, Ifremer, UMR 6539 LEMAR, F-29280 Plouzané, France							
9	<sup>+</sup> These authors contributed equally							
10	<sup>#</sup> Corresponding authors : alexandra.rahmani@univ-brest.fr ; vianney.pichereau@univ-							
11	brest.fr ; christine.paillard@univ-brest.fr							
12								
13	<u>Abbreviations</u>							
14	BRD: Brown Ring Disease							
15	d.p.i.: Days Post-Infection							
16	EPFs: Extrapallial Fluids							
17	FSSW: Filtered Sterilized Seawater							
18	OD: Optical Density							
19	Keywords: BRD, V. tapetis, R. philippinarum, pH, modulation effect, pH modifications,							

20 neutralizing activity

#### 21 Abstract

22 <u>Aims</u>

Brown Ring Disease (BRD) is an infection of the Manila clam *Ruditapes philippinarum* due to
the pathogen *Vibrio tapetis*. During BRD, clams are facing immunodepression and shell
biomineralization alteration. In this paper, we studied the role of pH on the growth of the
pathogen and formulated hypothesis on the establishment of BRD by *V. tapetis*.

#### 27 Methods and Results

In this study, we monitored the evolution of pH during the growth of *V. tapetis* in a range of
pH and temperatures. We also measured the pH of Manila clam hemolymph and extrapallial
fluids during infection by *V. tapetis*. We highlighted that *V. tapetis* modulates the external pH
during its growth, to a value of 7.70. During the development of BRD, *V. tapetis* also influences
extrapallial fluids and hemolymph pH *in vitro* in the first hours of exposure and *in vivo* after 3
days of infection.

#### 34 Conclusions

Our experiments have shown a close interaction between *V. tapetis* CECT4600, a pathogen of Manila clam that induces BRD, and the pH of different compartments of the animals during infection. These results indicate that that the bacterium, through a direct mechanism or as a consequence of physiological changes encountered in the animal during infection, is able to interfere with the pH of Manila clam fluids. This pH modification might promote the infection process or at least create an imbalance within the animal, that would favor its persistence. This last hypothesis should be tested in future experiment.

#### 42 <u>Significance and Impact of Study</u>

43 This study is the first observation of pH modifications in the context of BRD and might orient

44 future research on the fine mechanisms of pH modulation associated to BRD.

#### 45 **INTRODUCTION**

46

The Manila clam *Ruditapes philippinarum* is a bivalve mollusc species of significant economic 47 interest with a world production mainly from China, followed by European countries such as 48 Italy, Spain and France (Paillard 2004b). It was imported in France in the early 70's for 49 50 aquaculture. Ten years later, mortalities have been observed in clam's cultures, caused by an 51 infection called the Brown Ring Disease or BRD (Paillard et al. 1989). Indeed, infected clams 52 were characterized by a brown deposit in the inner face of the shell between the pallial line and the edge of the shell (Paillard et al. 1994). BRD is caused by a Gram negative bacterium, 53 54 Vibrio tapetis (Paillard and Maes 1990; Borrego et al. 1996). BRD has caused significant 55 economic losses in livestock farms on the Northern European coasts because the pathogen 56 has an optimum infection rate in cold waters at around 14°C. For example, in France, Manila 57 clam production decreased from 500 tons in 1987 to 200 tons in 1989, the aquaculture of clams production being mainly located in Brittany (France), an area highly sensitive to BRD 58 (Paillard 2004b). 59

*V. tapetis* is a microparasite acting in two modes. In the case of BRD, the bacterium is referred
to as an external microparasite because it degrades the periostracal lamina of the clam to
enter the extrapallial compartment without invading the tissues. In this case, the pathogen
induces BRD in Manila clams. In some cases, when tissues are damaged, *V. tapetis* acts as an
internal microparasite by infiltrating the animal's tissues, resulting in mortality within days
(Paillard 2004a).

66

67 During infection by *V. tapetis*, the pathogen therefore colonizes the periostracum and 68 degrades it in order to infiltrate peripheral or even central extrapallial fluids (EPFs). As a

result, it inhibits the biomineralization of the shell, preventing the formation of calcium
carbonate crystals of aragonite and thus leads to the accumulation of the organic brown
deposit (Paillard et al. 1994).

Furthermore, infected clams are facing an immunodepression caused by *V. tapetis*. Hemocytes, which are key effectors of the immune defenses of clams, are altered in BRD by actin cytoskeleton disorganization causing rounding of hemocytes and downregulation of pathogen recognition effectors. Consequently, infected hemocytes are no more able to ensure phagocytic progress to degrade the pathogen (Choquet et al. 2003; Allam et al. 2014; Rahmani et al. 2019).

Many factors are involved in BRD. Indeed, some alterations on the inner shell have significant impacts on the functioning of the affected clams. Impeding the functioning of adductor muscles and siphons, they result in abnormal positioning of the clam in the sediment and prevent the proper closure of valves that can lead to mortality when fishing or directly in the sediment, as clams can no longer retain their inter-valve water (Goulletquer et al. 1989; Paillard 2016).

Shell biomineralization is characterized by the formation of Calcium-Carbonate crystals (CaCO<sub>3</sub>) from calcium (Ca<sup>2+</sup>) and carbonate (CO<sub>3</sub><sup>2-</sup>) ions. CaCO<sub>3</sub> crystals are the main constituent of shellfish. In the Manila clam, the shell is composed of aragonitic crystals (Taylor, J D et al. 1973; Trinkler et al. 2011). The saturation state of this mineral in seawater determines whether the biomineralization reaction is favored in the direction of crystal formation or dissolution. It is defined by the concentrations of Ca<sup>2+</sup> and CO<sub>3</sub><sup>2-</sup> ions.

Biomineralization process is often studied in the context of ocean acidification because pH is
a factor that considerably influences the physiology of marine calcified organisms, and
especially this crystals formation. Indeed, acidification has the effect of reducing the

saturation rate of calcium carbonate crystals through the reduction in available  $CO_3^{2-}$  ions, 93 thus promoting the dissolution of the shell (Gattuso and Hansson 2011). In the mussel Mytilus 94 edulis, decrease in pH leads to a decrease in the net calcification rate, Ca<sup>2+</sup> ion content and, in 95 general, in a change in the ultrastructure of the shell and a decrease in amino acid content (Li 96 et al. 2015). Combination of temperature and pCO2 increase results in a decreased shell 97 hardness in the clam Mercenaria mercenaria and in the American oyster Crassostrea virginica, 98 indicating major changes in the biomineralization process (Ivanina et al. 2013). In the brittle 99 100 stars Amphiura filiformis, authors showed that the animal was able to compensate, in the short term, for the effects of pH decline by accelerating calcification and metabolism at the 101 102 expense of certain functions such as growth and reproduction because of the high energy cost required for this mechanism (Wood et al. 2008). In summary, most of the studies reflect 103 104 that ocean acidification reduces the calcification rates of animals composed of calcitic or 105 aragonitic crystal shells by promoting the dissolution reaction of these crystals.

106 In the Manila clams, calcification, i.e. shell biomineralization reaction, occurs in extrapallial fluids: ie the mantle secretes the shell components, the organic matrix (conchiolin) and the 107 108 calcium carbonate that form the aragonitic crystal shell (Paillard and Le Pennec 1993; Trinkler 109 2009; Trinkler et al. 2011). The periostracum (outer part of the shell) is made of a sclerotinized organic matrix. Shell organic matrix is mainly composed under normal conditions of aspartic 110 111 acid, glycine and serine, responsible for binding with calcium ions: chelation by sulphate ester 112 groups on polysaccharide chains would initiate biomineralization. When BRD develops, affected clams have deficiencies in amino acid such as glutamic acid, aspartic acid, serine, 113 114 and alanine, which can lead to poor binding to Calcium - Carbonate (Goulletguer et al. 1989). 115 Given the importance of pH homeostasis in the biomineralization process, it seems likely that pH might play a significant role in the pathogenicity of V. tapetis in the case of BRD. 116

Nevertheless, the influence of pH on the development of V. tapetis, as well as on the 117 development of BRD, have never been investigated. In this paper, we chose to 118 119 investigate the influence of pH on the development of the pathogen and the modulation 120 of pH associated to this bacterium in clams' fluids. Furthermore, we monitored for the first time the pH modification in clam fluids during *in vivo* challenge by V. tapetis allowing 121 us to formulate hypothesis that pH variations due to V. tapetis might be a virulence 122 strategy of this pathogen linked to pathogenicity and to its ability to alters 123 biomineralization. This study is the first investigation of pH modifications in the context 124 of BRD. 125

#### 127 Materials and methods

1) Monitoring of the pH modifications during growth of V. tapetis in culture media 128 129 <u>Culture media and bacterial strain</u>: The bacterial strain used in these experiments is V. tapetis CECT4600, grown in Zobell medium (Zobell 1941) and incubated at 18°C. The first objective 130 131 was to monitor the growth of this strain at 7 different pH values and 4 different temperatures. The 7 liquid culture media used in this experiment were obtained from Zobell media 132 133 supplemented with NaOH 1 M or HCl 0.1 M until the expected pH is obtained. After autoclave 134 sterilization, the pHs of the 7 media were 6.00, 6.45, 7.00, 7.55 (not modified, control medium, CM), 7.86, 8.12 and 8.70, respectively. pH of the cultures/media were measured by using a 135 Mettler Toledo© InLab Micro electrode after calibration of the electrode with pH buffers 4, 7 136 and 9 at the beginning and the end of each experiment. The 4 temperatures tested were 14°C 137 (optimal infection temperature), 18°C (optimal growth temperature), 21°C (temperature at 138 which the recovery process occurs mainly) and finally 27°C (temperature at which the bacteria 139 140 die) (Borrego et al. 1996; Paillard et al. 2004; Trinkler 2009; Paillard 2016).

<u>Growth experiments:</u> colonies of strain CECT4600 were resuspended in 2 ml of Zobell medium 141 (unmodified pH, control medium CM). The Optical Density (OD) was measured at a 142 143 wavelength of 492 nm and the suspension concentration was adjusted to an OD of about 0.1, i.e. to a cell density of 9.4 x 10<sup>7</sup> CFU.ml<sup>-1</sup>. The suspension was diluted at 1:100 in 3 ml of Zobell 144 medium at the desired pH. The control samples were the same sterile media. 400 µL of 145 inoculated media or controls were distributed in a Bioscreen© 100-wells plate (5 replicates 146 per tested condition), allowing a sterile, thermally controlled culture with an OD 147 measurement at regular intervals under stirring. This protocol was repeated twice; each 148 experiment was run for 24 hours (except at 14°C where the experiment ran for 30 h to take 149

into account the reduced bacterial growth), in a shaking mode, with measurement of OD
every 20 minutes at a wavelength of 492 nm; temperature was set up at either 14, 18, 21 or
27°C. To represents the pH measurement at the beginning of the growth experiment (To) and
TF (TFinal) represent the pH measurement at the end of experiment, depending on the
different conditions as detailed above.

155

156 pH monitoring during the bacterial growth: Two precultures of V. tapetis CECT4600 were performed in Zobell broth during 15 h at 18°C and 24 h at 14°C in order to reach an OD of 157 0.98. Each preculture was diluted 12 times (initial OD = 0.08) with fresh Zobell medium at the 158 different pHs (from 6.00 to 8.70). With these new cultures, bacterial growth was performed 159 under shaking at 14° and 18°C, respectively from the corresponding pre-culture, in three 160 161 replicates. The pH was measured, from 200 µl-samples, at the beginning of the experiment, every hour during the exponential growth phase, and finally in the stationary phase (i.e. 48 h 162 163 of growth at 14°C and 24 h of growth at 18°C).

164

165

# 2) Monitoring of the pH in extrapallial fluids (EPFs) and hemolymph of Manila clams during infection by *V. tapetis*

167

166

168 <u>Animals:</u> Animals used in this study were adult Manila clams (about 40 mm) from Landeda 169 (Finistère, France), kindly provided by the company SATMAR. One pool of 40 clams (for the 170 *in vitro* study of fluids) and 2 pools of 100 clams (for the *in vivo* infection study) were 171 acclimatized for 3 and 2 weeks, respectively, in oxygenated seawater at 14°C.

172

173 *Fluids and shell sampling:* The fluids chosen for this experiment were hemolymph and EPFs. Hemolymph, EPFs and shell were randomly collected extemporaneously from 10 clams. 174 175 Hemolymphs have been collected from the adductor muscle and EPFs from the extrapallial 176 cavity of the animals as previously described (Le Bris et al. 2015). The quality of the hemolymph samples was checked under the microscope (presence of pseudopods, 177 178 unrounded hemocytes, serum and hemocytes without microorganism contaminations) then 179 the samples were pooled and the number of hemocytes was determined by using a Malassez 180 counting grid. Shell parts were broken with a hammer to obtain small pieces and were sterilized by UV (wavelength 254 nm) for 15 minutes. EPFs were also pooled. The hemocytes 181 concentration in the hemolymph pool was  $5.45 \times 10^5$  hemocyte per ml. 182

183

184 For *in vitro* experiments, 150 µl of FSSW, hemolymph or EPFs were placed in 1.5 ml sterile 185 tubes, with supplementation of powdered Manila clam shell, in the same amount in each sample, in order to design an *in vitro* experiment that is as close as possible of natural 186 conditions and to allow ion exchanges between fluids and shell that might be dependent of 187 188 the V. tapetis growth. Then, 200 µL of a V. tapetis CECT4600 suspension in FSSW was added to each tube, in order to reach a final bacteria: hemocytes ratio of  $25:1(1.37 \times 10^7 \text{ CFU.ml}^{-1})$ ; 189 200 µl of FSSW was added for the controls. Each condition was tested in triplicates. pH 190 191 measures were performed at times 0 h, 3 h, 24 h, 48 h and 72 h and tubes were shaken before 192 measurement.

193

For *in vivo* infection experiments, the protocol was followed as previously described (Paillard
and Maes 1994). The animals were first left out of the water during the night before infection
to allow the valves to open spontaneously in water, and then injected with 100 µl of FSSW for

50 control individuals, or 100  $\mu$ l of bacterial suspension (5 x 10<sup>8</sup> CFU.ml<sup>-1</sup> in FSSW) for 50 197 infected individuals, in the pallial cavity. The injected animals were left out of water for 3 198 199 additional hours to allow the pathogen to colonize them, and then were transferred again 200 into their respective seawater tanks. Animals were sampled at different time points after 201 infection, ie. after 15 hours, 1 day, 3 days, 7 days, 14 days and 21 days post injection (d.p.i.). 202 pH of the EPFs and hemolymph samples were measured by using the Mettler Toledo 203 microelectrode calibrated with pH 4, 7 and 9 buffers. The initial sample (To) was taken just 204 before injection. The mode of infection does not induce mortality and allows to reproduce first stage of BRD to see the effect of V. tapetis, even before the development of organic 205 206 deposit in the shell, highlighting the pH variations that occurs in the really first days of V. *tapetis* infection. 207

208

209 <u>Statistical analyses:</u> Statistical analyses were performed by using a Student test or a Mann
210 Whitney Wilcoxon test, depending of data normality.

#### 211 **RESULTS**

#### 212 V. tapetis CECT4600 modulates the pH of Zobell culture medium during its growth.

*V. tapetis* was cultured in Zobell broths which pH were adjusted from 6.00 to 8.70, at 4
different temperatures ranging from 14°C to 27°C. The results obtained are reported in Table
1. First, we observed that the bacterium is able to grow at temperatures ranging from 14°C
to 21°C, in the pH range tested, but not at 27°C. In addition, consistently with the results
previously published by Borrego et al (1996), the optimal growth temperature is 18°C in the
unmodified medium.

V. tapetis has a high tolerance to pH because we observed a growth in all the tested culture 219 media. However, at 14°C, the fastest growth was obtained for the control medium (pH 7.4, 220 DT = 3.15 h), but growths were observed over a pH range from 6.45 to 8.12, with doubling 221 222 times from 3.15 to 3.5 and max ODs higher than 1, while the growth rates were lower at the 223 extreme pHs tested, ie. 6 and 8.7 (Table 1). The best growth at 14°C was observed in the 224 unmodified culture medium (MC, with a DT of 3.15h), and at pH 7.86 at 18°C (DT = 1.79h). Interestingly, at the growth temperatures 18 and 21°C, very good growth parameters were 225 obtained for more acidic pHs, from which the lowest DTs were observed for the pH value 6.45 226 227 (DTs = 1.85 and 1.93 h at 18 and 21°C, respectively).

The pHs of the culture media have been measured at both the beginning (To) and the end (TF) of bacterial growth in all the conditions tested. Results are presented in Figures 1 to 3. In the control samples (no bacteria), the pHs did not change significantly between the measurements made at To and TF (Fig. 1 to 3). By contrast, after growth of *V. tapetis* at 14, 18 and 21°C, the pH changed during growth to reach an average value of 7.90 in all media tested (Fig. 1 and 2). The pH modifications are the same at these three temperatures, allowing us to

show them all in the same figures (Fig. 1 and 2). We did not observe such a pH modification
of the culture medium at 27°C, a condition where no growth of the bacterium was observed
(Fig. 3).

Figure 2 shows the statistical distribution of pH values measured at To and TF in all media. The mean and variance of this distribution differed significantly during the experiment only when the bacterium was able to grow (*pvalue* < 0.0001). Overall, these results show that *V. tapetis* is able to modify the pH of its culture medium during its growth towards a pH close to 7.90.

We measured the evolution of medium pH all along the exponential growth phase of V. 242 tapetis at 14°C (Fig. 4) and 18°C (Fig. 5). In the control samples, the pH changed only slightly 243 244 during the experiment, resulting in very low slope curves (data not shown). By contrast, in the experimental samples, during the growth of V. tapetis, the pHs of the media (which were 245 246 initially 6 to 8.7) converged to an average value of 7.71, whatever the temperature of growth 247 (Fig. 4 and 5). Therefore, our experiments show that V. tapetis is able, during its growth, to modify the pH of its culture medium. More precisely, V. tapetis tends to 'neutralize' culture 248 249 media during its growth, by acidifying alkaline media and alkalinizing acidic media, to make 250 it reach a mean value of 7.7-7.9 whatever the initial pH of the culture medium. However, this 251 effect can be due to either a passive mechanism as a consequence of the bacterium's growth or an active mechanism that allow the pathogen to infect the Manila clam. 252

# 253 <u>External pH modulation during the in vitro growth of V. tapetis CECT4600 in clams</u> 254 <u>extrapallial fluids or hemolymph.</u>

255 We aimed at determining whether the bacterium can also modulate the pH of clam 256 biological fluids in which *V. tapetis* is known to proliferate during the infection, ie. the EPFs

257 and the hemolymph. In these experiments, FSSW was used as the reference fluid because it does not allow any growth of *V. tapetis*. The same amount of powdered shell fragments was 258 259 added to each sample in order to design an *in vitro* experiment that is as close as possible of 260 natural conditions and to allow ions exchanges between fluids and shell that might influence V. tapetis growth. The obtained results are shown in the Figure 6. Our experiments showed 261 262 that in FSSW, the presence of V. tapetis did not lead to any significant change in pH as 263 compare to the control, at each time point tested. By contrast, in EPFs and hemolymph, 264 exposure to V. tapetis led to a significant decrease of pH as compared to the initial conditions 265 after 3 hours of exposure (*pvalue < 0.001*). Indeed, for both clam biological fluids (ie. EPFs and 266 hemolymph), the pH significantly dropped from 7.6 to 7.4 after three hours of Vibrio exposure, and then stabilized at approx. 7.25. However, for longer times, the pH also tended to decrease 267 268 in the controls, so as no significant difference appears between the control samples and the 269 samples exposed to *V. tapetis* for the last time points, thus revealing an interaction only in 270 the first hours of exposure.

# 271 <u>Extrapallial fluids and hemolymph pH modulation during in vivo challenge of the Manila</u> 272 <u>clams by V. tapetis CECT4600.</u>

273 The previous demonstration that the presence of V. tapetis can interfere in vitro with the pH 274 of these clam biological fluids led us to hypothesize that this could happen in vivo, and that 275 the modulation of pH of these fluids could be an important aspect of the infection process. 276 Monitoring of pH evolution in EPFs and hemolymph after Manila clams infection by V. tapetis 277 is presented in Figures 7 and 8, respectively. Both fluids displayed a relatively similar pH 278 evolution following the injection of FSSW or V. tapetis in the animals. Considering the whole 279 in vivo infection experiment (22 days), we did not observe any significant difference in pH 280 between the control and infected animals for both fluids. However, we observed a weak but

significant difference between the pHs of the EPFs and hemolymph from control and infected
clams at 3 days post injection (dpi). Indeed, the pH of these fluids were higher (of approx. o.2
pH unit) in both cases in clams injected with a suspension of *V. tapetis*, as compared to clams
injected with FSSW (*pvalue < o.o2*). Our results then tend to suggest that exposure to *V. tapetis* induce physiological changes in challenged Manila clam that influence the pH of EPF
and hemolymph at 3 days of exposure to the pathogen. This also suggests that this particular
period should be more precisely investigated during the study of host-pathogen interaction.

#### 289 **DISCUSSION**

pH homeostasis is an essential aspect of cell physiology. It is also one of the main 290 291 environmental factors controlling the biomineralization of calcified organisms. This explains, at least in part, the current abundance of studies dealing with the impact of ocean 292 293 acidification on marine mollusks and other calcified marine organisms (Gattuso and Hansson 294 2011). The relation between decrease of pH and an increase of temperature has been studied 295 in several bivalves species, showing that these two factors can affect the immune response 296 as in the Mediterranean mussel Mytilus galloprovincialis (Matozzo et al. 2012) and was also 297 shown to influence the abundance of parasites and incidence of bacteria as in the blue mussel 298 Mytilus edulis (Mackenzie et al. 2014). Finally, pH adaptation is a crucial component of virulence of many pathogenic bacteria. In addition to having to survive the acid digestive 299 300 barriers in some hosts, they must be able to tolerate acid stress resulting from lysosomal 301 activity after being phagocytized by immune system cells such as macrophages or hemocytes 302 (Asplund et al. 2014).

303

304 EPFs and hemolymph are involved in shell biomineralization and in immunity of the Manila clam, respectively, and have therefore a crucial importance in the development of BRD. BRD 305 is typically a disease of clam biomineralization (Paillard et al. 1994). It is also a bacterial 306 307 disease, in which the phenomenon of phagocytosis, and more specifically the phagosome-308 lysosome fusion, has a particular importance in the animal's immunity (Paillard 2004; Rahmani et al., 2019). For all these reasons, it is surprising that the effect of pH on V. tapetis 309 310 has never been considered to date in the context of the BRD. In this study, we first aim to 311 characterize the relationship between external medium pH and the growth of the pathogen V. tapetis. 312

313 We observed that V. tapetis, the bacterial etiological agent of BRD, is able to grow over a wide range of pHs, including at a cold temperature (14°C) which is an optimum for BRD 314 315 development (Paillard et al. 2004). Interestingly, we showed that the bacterium, during its 316 growth, modifies the pH of its environment, increasing it in acidic media, or acidifying it in 317 alkaline ones, and systematically replaced the external pH to a value close to 7.7 during its 318 growth in a kind of 'neutralizing activity' in the tested range. We further characterized this 319 last point, by measuring *in vitro* the impact of the bacterial growth on the pH of the main 320 biological fluids of the Manila clams, ie. extrapallial fluids and hemolymph, and then in vivo, the pH of these fluids in a context of infection by inducing the main mode of infection of V. 321 322 *tapetis*, allowing the pathogen to colonize extrapallial fluids. We observed that at three hours after exposure to V. tapetis, the pH of the two fluids decreased significantly more than that 323 324 of the control in presence of powdered Manila clam shell. EPFs and hemolymph contain 325 hemocytes, which are the main cellular actors of the immune system and one of the main 326 targets of the pathogen.

327

328 In a previous transcriptomic analysis, we showed that V. tapetis is able to induce 329 deregulations of clam hemocytes physiology as reorganization of actin cytoskeleton, reduction of lysosomal activity and down regulation of genes related to the complement 330 331 pathway (Rahmani et al. 2019). In another previous study, our team had demonstrated that 332 the cytotoxicity of the bacterium to hemocytes was maximal, in vitro, after three hours of exposure to the pathogen (Choquet et al. 2003). This phenomenon is reproduced by exposing 333 334 Manila clam hemocytes to V. tapetis during precisely three hours. The decrease in pH in the 335 first three hours of exposure could be due to the growth of the pathogen and could be related

(either as a cause or a consequence) to its cytotoxic effect on clam hemocytes where theinfluence of the bacterium is maximum.

338

The relationship between pH regulation and virulence is not always clear since the regulation 339 of the internal pH is a necessity for all cell types. Bacteria regulate their intracellular pH by 340 341 using proton pumps, or by transporting and/or metabolizing several acid or base compounds. For example, Escherichia coli response to acid stress involves amino acid decarboxylase 342 343 antiporter pairs and proton-pumping respiratory chains complexes causing H+ ion efflux outside 344 the cell. On the contrary, during alkaline stress, *E. coli* was found to have a strong activation of NhaA Na<sup>+</sup>/ $H^+$  antiporter, an up-regulation of ATP synthetase leading to a strong entry of 345 H+ ion into the cell (Krulwich et al. 2011). However, intestinal bacteria are facing larger pH 346 347 ranges than marine ecosystems. NhaA antiporter is also present in the genome of V. tapetis CECT4600 according to the recently published annotation (Dias et al. 2018), as well as 348 349 calcium or potassium proton antiporters. Furthermore, the presence of an enzyme such as arginine deiminase in the genome of this pathogen might explain alkalinization of external 350 351 pH observed during acidic stress as already shown in another bacterium (Budin-Verneuil et 352 al. 2006).

Regardless of this internal pH regulation, many bacteria modify the pH of their environment as a result of their energy metabolism. This is particularly the case for many fermentative bacteria producing large quantities of organic acids (e.g. lactic acid, propionic acid, formic acid, etc.) leading to an acidification of their environment (Nuryana et al. 2019). Other bacteria are known to alkalinize their environment, by using organic acids (e.g. lactate, aspartate, glutamate) as carbon and energy sources (Stancik et al. 2002). Nevertheless, these metabolisms are very dependent on the nature of the substrates used by the bacteria, and

there are no case, to our knowledge, of bacteria capable of being both acidifying andalkalinizing in the same culture medium, only depending on the initial pH of that medium.

362 This discovery raises many questions about the V. tapetis's adaptation mechanisms to pH stress, as well as about the energy metabolism(s) it uses, and the substrates it degrades from 363 the Zobell medium, at the different pHs. This also raises important questions regarding the 364 involvement of this 'neutralizing activity' in the expression of its pathogenicity, in the context 365 366 of BRD development. The mechanisms related to this activity are for now unknown and 367 might be due to several type of mechanisms as described above. Nevertheless, this study has revealed new, previously unsuspected problematics related to pH modulation, highlighting 368 369 the need for further analysis of genes expressed during growth in order to elucidate the mechanisms linked to this phenomenon. 370

371

372 To better understand the close interaction between the clam and the pathogen in this context, we need to refer to BRD dynamics. BRD is a chronic infection, as the pathogen does 373 not induce mortalities in the classical way of infection and can persist in infected animals 374 375 (Paillard 2016). Indeed, the acute phase of the infection is divided into two parts: a first one 376 characterized by an increase of V. tapetis concentration followed by the production of a brown deposit on the inner part of the shell, and a second one, which is not always present, 377 378 characterized by shell repair that can lead to the complete remission of clams (Paillard 2016). 379 In this study, we have performed injection of V. tapetis in the pallial cavity. This mode of infection allows recreating the first stages of BRD by allowing the pathogen to reproduce all 380 381 the early steps of BRD and then induce an acute phase of infection until formation of the 382 brown deposit. Dynamics of BRD and kinetics of V. tapetis in EPFs have previously been well characterized. In the first hours of infection, the pathogen colonizes and degrades the 383

periostracal lamina in order to enter extrapallial fluids, and then colonizes the shell secretion
(Paillard and Maes 1995), leading to an increase of *V. tapetis* concentration in EPFs. Between
2 days and 7 d.p.i. the EPFs concentration of *V. tapetis* reaches its maximum value (Bidault et
al. 2015).

388 This particular period corresponds to an increase of pH in EPFs according to our study where 389 significant differences of pH have been observed in infected animals after 3 days (*in vivo*). It 390 should also be noted that an increase of pH has also been reported in hemolymph at 3 d.p.i., 391 thus probably revealing the changes that occurs in this compartment at the biochemical level 392 during BRD (Allam et al. 2006).

393

In addition, we know that during infection, major changes occur in EPFs at the enzymatic 394 395 level with, for example, an increase in the activity of phenoloxidase, an enzyme involved in 396 the humoral response and melanization which leads to the brown deposit production (Söderhäll and Cerenius 1998). Previous studies showed that phenoloxidase activity is 397 sensitive to abiotic factors such as pH (Le Bris 2013). Indeed, phenoloxidase activity decrease 398 399 with pH in a range of physiological values in the Atlantic blue crab *Callinectes sapidus*, mostly 400 in the hemolymph compartment (Tanner et al. 2006). Thus, increasing the pH in a range of physiological values in both hemolymph and EFPs could modulate phenoloxidase activity in 401 402 these two compartments. As phenoloxidase is related to melanization process, the increase 403 of pH can also modulate melanization, associated to the bacterial embedding within shell 404 matrix.

405

406 Considering these informations, the fact that *V. tapetis* is able to modulate the pH of such
407 compartments of the Manila clam in the context of BRD then questioned the impact of this

408 factor in the development of BRD itself. Indeed, pH is known to modulate the virulence of many pathogenic bacteria. In *Vibrio cholerae*, for example, it has been shown that the protein 409 410 ToxR, which is responsible for the regulation of many virulence genes, is also strongly 411 involved in the response to acid stress, suggesting a close relationship between the virulence 412 and the acid stress response in this pathogen (Merrell and Camilli 2002; Lund et al. 2014). In 413 Pseudomonas aeruginosa, it has been shown that the infection was attenuated at pH 6 rather 414 than pH 7.6, and that the bacterium expresses genes that allow it to alkalinize the medium at 415 acidic pH, while at neutral pH, it preferentially expresses iron metabolism-related genes 416 (Romanowski et al. 2011). In the case of *V. tapetis*, we don't know yet if pH variations are due 417 to sensing of external pH by V. tapetis that induce a change in response, maybe in relation with its pathogenic activity or to the growth of the pathogen. This mechanism should be 418 419 investigated in a future study.

420

To summarize, after investigation for pH modulation during interaction between the Manila 421 clam and its pathogen V. tapetis, our results indicate that V. tapetis is able to modulate the 422 423 pH by a "neutralizing activity" during its growth. For the first time, we highlighted this activity 424 in both *in vitro* challenge and *in vivo* infection of Manila clam. We have determined that this interaction occurs at a very precise time (3 hours exposure in vitro and 3 days in vivo). These 425 426 special time periods correspond to the main interactions between the pathogen and the clam 427 hemocytes. It is then likely that these close interactions might play a role in the first steps of BRD development. Nevertheless, the mechanisms related to this phenomenon are, for now, 428 429 unknown. Our study is the first evidence that pH modulations might be a novel, and since 430 now undiscovered, mechanism that will help us to better understand host-pathogen interaction in the context of BRD. Moreover, it allows to focus on really precise time period 431

432 by both *in vitro* and *in vivo* challenges in order to better characterize these particular 433 interactions and to understand the mechanisms and the importance of pH modulation in the 434 pathogenic activity of *V. tapetis*.

435

436

#### 437 AUTHORS CONTRIBUTIONS

AR designed the protocols with help of AB, NLG and CP and VP. AR performed *in vitro* experiments on *V. tapetis*. AR and CM performed *in vitro* and *in vivo* experiment on Manila clams challenged by *V. tapetis*. AR, VP and CP wrote the article (the original draft was written by AR). This article was carefully reviewed by other co-authors, who all approved the final version.

443

#### 444 FUNDINGS

This project received grants from the H2020 European project "VIVALDI" (grant agreement
N°678589). This work was also supported by the "Université de Bretagne Occidentale" (UBO,
France), and the "investment for the future" programs LabexMER (ANR-10-LABX-19) and
ISblue (ANR-17-EURE-0015).

449

#### 450 ACKNOWLEDGMENTS

We warmly thank Jean François AUVRAY from the SATMAR company of Landeda (Finistère,
France) for providing the clams. We also warmly thank Eric DABAS for helping with
zootechnical support.

### **CONFLICT OF INTEREST**

456 No conflict of interest declared.

#### 459 **REFERENCES**

Allam B, Paillard C, Auffret M, Ford SE (2006) Effects of the pathogenic Vibrio tapetis on 460 461 defence factors of susceptible and non-susceptible bivalve species: II. Cellular and 462 biochemical changes following in vivo challenge. Fish Shellfish Immunol 20:384-397 463 Allam B, Pales Espinosa E, Tanguy A, Jeffroy F, Le Bris C, Paillard C (2014) Transcriptional changes in Manila clam (Ruditapes philippinarum) in response to Brown Ring 464 Disease. Fish Shellfish Immunol 41:2–11 . https://doi.org/10.1016/j.fsi.2014.05.022 465 466 Asplund ME, Baden SP, Russ S, Ellis RP, Gong N, Hernroth BE (2014) Ocean acidification and host-pathogen interactions: blue mussels, Mytilus edulis, encountering Vibrio 467 tubiashii. Environ Microbiol 16:1029–1039. https://doi.org/10.1111/1462-2920.12307 468 Bidault A, Richard GG, Le Bris C, Paillard C (2015) Development of a Tagman real-time PCR 469 470 assay for rapid detection and quantification of Vibrio tapetis in extrapallial fluids of 471 clams. PeerJ 3:e1484 . https://doi.org/10.7717/peerj.1484 472 Borrego JJ, Castro D, Luque A, Paillard C, Maes P, Garcia MT, Ventosa A (1996) Vibrio 473 tapetis sp. nov., the causative agent of the brown ring disease affecting cultured clams. Int J Syst Evol Microbiol 46:480-484 474 475 Budin-Verneuil A, Maguin E, Auffray Y, Ehrlich DS, Pichereau V (2006) Genetic structure and transcriptional analysis of the arginine deiminase (ADI) cluster in Lactococcus 476 477 lactis MG1363. Can J Microbiol 52:617-622. https://doi.org/10.1139/wo6-009 478 Choquet G, Soudant P, Lambert C, Nicolas J-L, Paillard C (2003) Reduction of adhesion 479 properties of *Ruditapes philippinarum* hemocytes exposed to *Vibrio tapetis*. Dis Aquat Organ 57:109–116 480 Dias GM, Bidault A, Le Chevalier P, Choquet G, Der Sarkissian C, Orlando L, Medique C, 481 Barbe V, Mangenot S, Thompson CC, Thompson FL, Jacq A, Pichereau V, Paillard C 482 (2018) Vibrio tapetis Displays an Original Type IV Secretion System in Strains 483 Pathogenic for Bivalve Molluscs. Front Microbiol 9:227 484 Gattuso J-Pierre, Hansson Lina (2011) Ocean acidification. Oxford University Press, 485 Oxford ; 486 487 Goulletquer P, Héral M, Béchemin C, Richard P (1989) Anomalies de calcification chez la 488 palourde japonaise Ruditapes philippinarum: caractérisation et comparaison des 489 compositions en acides aminés de différentes parties de la coquille analysées par HPLC. Aquaculture 81:169–183 490 491 Ivanina AV, Dickinson GH, Matoo OB, Bagwe R, Dickinson A, Beniash E, Sokolova IM 492 (2013) Interactive effects of elevated temperature and CO 2 levels on energy 493 metabolism and biomineralization of marine bivalves Crassostrea virginica and Mercenaria mercenaria. Comp Biochem Physiol A Mol Integr Physiol 166:101–111 494 Krulwich TA, Sachs G, Padan E (2011) Molecular aspects of bacterial pH sensing and 495 496 homeostasis. Nat Rev Microbiol 9:330-343 497 Le Bris C (2013) Le système phénoloxydase: caractérisation biochimique et rôle dans la réponse immunitaire chez la palourde japonaise Venerupis philippinarum exposée à 498 Vibrio tapetis. Université de Bretagne occidentale-Brest 499 Le Bris C, Richard G, Paillard C, Lambert C, Seguineau C, Gauthier O, Pernet F, Guérard F 500

501 (2015) Immune responses of phenoloxidase and superoxide dismutase in the manila 502 clam Venerupis philippinarum challenged with Vibrio tapetis – Part I: Spatiotemporal evolution of enzymes' activities post-infection. Fish Shellfish Immunol 503 504 42:16–24 . https://doi.org/10.1016/j.fsi.2014.10.021 505 Li S, Liu C, Huang J, Liu Y, Zheng G, Xie L, Zhang R (2015) Interactive effects of seawater acidification and elevated temperature on biomineralization and amino acid 506 metabolism in the mussel Mytilus edulis. J Exp Biol 218:3623-3631. 507 508 https://doi.org/10.1242/jeb.126748 509 Lund P, Tramonti A, De Biase D (2014) Coping with low pH: molecular strategies in 510 neutralophilic bacteria. FEMS Microbiol Rev 38:1091-1125. 511 https://doi.org/10.1111/1574-6976.12076 512 Mackenzie CL, Lynch SA, Culloty SC, Malham SK (2014) Future Oceanic Warming and 513 Acidification Alter Immune Response and Disease Status in a Commercial Shellfish 514 Species, Mytilus edulis L. PLOS ONE 9:e99712. 515 https://doi.org/10.1371/journal.pone.0099712 516 Matozzo V, Chinellato A, Munari M, Finos L, Bressan M, Marin MG (2012) First Evidence of 517 Immunomodulation in Bivalves under Seawater Acidification and Increased 518 Temperature. PLOS ONE 7:e33820 . https://doi.org/10.1371/journal.pone.oo33820 Merrell D, Camilli A (2002) Acid tolerance of gastrointestinal pathogens. Curr Opin 519 520 Microbiol 5:51-55. https://doi.org/10.1016/S1369-5274(02)00285-0 Nuryana I, Andriani A, Lisdiyanti P, Yopi (2019) Analysis of organic acids produced by lactic 521 522 acid bacteria. IOP Conf Ser Earth Environ Sci 251:012054. 523 https://doi.org/10.1088/1755-1315/251/1/012054 524 Paillard C, Allam B, Oubella R (2004) Effect of temperature on defense parameters in Manila clam Ruditapes philippinarum challenged with Vibrio tapetis. Dis Aquat 525 526 Organ 59:249–262 Paillard C (2004a) A short-review of brown ring disease, a vibriosis affecting clams, 527 Ruditapes philippinarum and Ruditapes decussatus. Aquat Living Resour 17:467–475. 528 529 https://doi.org/10.1051/alr:2004053 530 Paillard C (2004b) Rôle de l'environnement dans les interactions hôtes-pathogènes; développement d'un modèle de vibriose chez les bivalves. Habilit À Dir Rech HDR 531 Univ Bretagne Occident Brest 532 533 Paillard C (2016) An ecological approach to understanding host-pathogen-environment 534 interactions: the case of Brown Ring Disease in clams. In: Oysters and Clams: 535 Cultivation, Habitat Threats and Ecological Impact 536 Paillard C, Le Pennec M (1993) Ultrastructural studies of the mantle and the periostracal 537 lamina in the manila clam, Ruditapes philippinarum. Tissue Cell 25:183–194. https://doi.org/10.1016/0040-8166(93)90018-G 538 539 Paillard C, Maes P (1990) Aetiology of brown ring disease in *Tapes philippinarum*: 540 pathogenicity of a Vibrio sp. Comptes Rendus Académie Sci Ser 3 Sci Vie 310:15–20 541 Paillard C, Maes P (1994) Brown ring disease in the Manila clam *Ruditapes philippinarum*: 542 establishment of a classification system. Dis Aquat Organ 19:137–146

543 544 545	Paillard C, Maes P (1995) The Brown Ring Disease in the Manila Clam, <i>Ruditapes philippinarum</i> . Part 1. Ultrastrucural alterations of the periostracal lamina. J Invertebr Pathol 65:91–100 . https://doi.org/10.1006/jipa.1995.1015
546	Paillard C, Maes P, Oubella R (1994) Brown ring disease in clams. Annu Rev Fish Dis 4:219–
547	240 . https://doi.org/10.1016/0959-8030(94)90030-2
548	Paillard C, Percelay L, Le Pennec M, Le Picard D (1989) Origine pathogène de l'«anneau
549	brun» chez <i>Tapes philippinarum</i> (Mollusque, bivalve). Comptes Rendus Académie
550	Sci Sér 3 Sci Vie 309:235–241
551	Rahmani A, Corre E, Richard G, Bidault A, Lambert C, Oliveira L, Thompson C, Thompson
552	F, Pichereau V, Paillard C (2019) Transcriptomic analysis of clam extrapallial fluids
553	reveals immunity and cytoskeleton alterations in the first week of Brown Ring
554	Disease development. Fish Shellfish Immunol 93:940–948 .
555	https://doi.org/10.1016/j.fsi.2019.08.025
556	Romanowski K, Zaborin A, Fernandez H, Poroyko V, Valuckaite V, Gerdes S, Liu DC,
557	Zaborina OY, Alverdy JC (2011) Prevention of siderophore- mediated gut-derived
558	sepsis due to <i>P. aeruginosa</i> can be achieved without iron provision by maintaining
559	local phosphate abundance: role of pH. BMC Microbiol 11:212.
560	https://doi.org/10.1186/1471-2180-11-212
561	Söderhäll K, Cerenius L (1998) Role of the prophenoloxidase-activating system in
562	invertebrate immunity. Curr Opin Immunol 10:23—28 .
563	https://doi.org/10.1016/S0952-7915(98)80026-5
564	Stancik LM, Stancik DM, Schmidt B, Barnhart DM, Yoncheva YN, Slonczewski JL (2002)
565	pH-Dependent Expression of Periplasmic Proteins and Amino Acid Catabolism in
566	<i>Escherichia coli.</i> J Bacteriol 184:4246–4258 . https://doi.org/10.1128/JB.184.15.4246-
567	4258.2002
568 569 570	Tanner CA, Burnett LE, Burnett KG (2006) The effects of hypoxia and pH on phenoloxidase activity in the Atlantic blue crab, <i>Callinectes sapidus</i> . Comp Biochem Physiol A Mol Integr Physiol 144:218–223
571	Taylor, J D, Kennedy, W J, Hall, A (1973) The Shell Structure and Mineralogy of the Bivalvia.
572	II. <i>Lucinacea-Clavagellacea</i> . Conclusions. Bull Br Mus Nat Hist Zool Lond 22:253–
573	294
574	Trinkler N (2009) La guérison coquillière: un mécanisme de défense de la palourde
575	japonaise <i>Ruditapes philippinarum</i> face au <i>Vibrio tapetis</i> dans le cadre de la maladie
576	de l'anneau brun. Université de Bretagne occidentale-Brest
577 578 579	Trinkler N, Bardeau J, Marin F, Labonne M, Jolivet A, Crassous P, Paillard C (2011) Mineral phase in shell repair of Manila clam <i>Venerupis philippinarum</i> affected by brown ring disease. Dis Aquat Organ 93:149–162 . https://doi.org/10.3354/da002288
580 581	Wood HL, Spicer JI, Widdicombe S (2008) Ocean acidification may increase calcification rates, but at a cost. Proc R Soc Lond B Biol Sci 275:1767–1773
582 583	Zobell C (1941) Studies on marine bacteria. I. The cultural requirements of heterotrophic aerobes. J Mar Res 4:41–75
584	

#### FIGURES AND TABLES

Figure 1: pH changes in a V. tapetis CECT4600 culture after growth at 14°C, 18° and 21°C in Zobell media which pH were initially adjusted from
6.00 (M600) to 8.70 (M870) (15 replicates)
Figure 2: pH variance of a V. tapetis CECT4600 culture after growth at 14°, 18° and 21°C in pH adjusted Zobell media from 6.00 (M600) to 8.70
(M870) (15 replicates). Left : To ; Right : TF or TFinal
Figure 3: pH changes in a V. tapetis CECT4600 culture after growth at 27° in Zobell media which pH were initially adjusted from 6.00 (M600) to
8.70 (M870) (15 replicates)
Figure 4: pH change in a culture of <i>V. tapetis</i> during the exponential growth phase at 14°C, in Zobell media which pH were initially adjusted
from 6.oo (M6oo) to 8.70 (M87o) ; CM is the original medium where pH was not modified
Figure 5: pH change in a culture of V. tapetis during the exponential growth phase at 18°C, Zobell media which pH were initially adjusted from
6.oo (M6oo) to 8.7o (M87o) ; CM is the original medium where pH was not modified
Figure 6: pH changes after V. tapetis exposure in FSSW (Filtered sterilized Seawater), EPF (extrapallial fluids) and Hemolymph (He) mixed with
powdered Manila clam shell. ** : pvalue=0.0017 ; ***=pvalue=0.0007, Student test
Figure 7: pH changes in EPFs (extrapallial fluids) of the Manila clam <i>Ruditapes philippinarum</i> after infection by <i>V. tapetis</i> . Control: injected by
FSSW ; Infected: injected by <i>V. tapetis</i>
Figure 8: pH changes in Hemolymph of the Manila clam Ruditapes philippinarum after infection by V. tapetis. Control: injected by FSSW ;
Infected: injected by <i>V. tapetis</i>



Figure 1: pH changes in a *V. tapetis* CECT4600 culture after growth at 14°C, 18° and 21°C in Zobell media which pH were initially adjusted from 6.00 (M600) to 8.70 (M870) (15 replicates)



Figure 2: pH variance of a *V. tapetis* CECT4600 culture after growth at 14°, 18° and 21°C in pH adjusted Zobell media from 6.00 (M600) to 8.70 (M870) (15 replicates). Left : To ; Right : TF or TFinal

\*\*\*\* : pvalue < 0.0001, Student test



Figure 3: pH changes in a *V. tapetis* CECT4600 culture after growth at 27° in Zobell media which pH were initially adjusted from 6.00 (M600) to 8.70 (M870) (15 replicates)







Figure 6: pH changes after *V. tapetis* exposure in FSSW (Filtered sterilized Seawater), EPF (extrapallial fluids) and Hemolymph (He) mixed with powdered Manila clam shell. \*\* : pvalue=0.0017 ; \*\*\*=pvalue=0.0007, Student test



Figure 7: pH changes in EPFs (extrapallial fluids) of the Manila clam *Ruditapes philippinarum* after infection by *V. tapetis*. Control: injected by FSSW ; Infected: injected by *V. tapetis* 



Figure 8: pH changes in Hemolymph of the Manila clam *Ruditapes philippinarum* after infection by *V. tapetis*. Control: injected by FSSW ; Infected: injected by *V. tapetis* 

Table 1: growth parameters of *V. tapetis* in a range of pH media and different temperatures. DT : doubling Time ; OD : Optic Density, T° :

Temperature; h: hour. M=Medium and the number is the pH (600= pH 6.00); MC is the original medium (unmodified pH).

		<i>V. tapetis</i> CECT4600 growth													
	M6oo		M645		М700		МС		M786		M812		M870		
T٩	Time of growth	DT (h)	Maximum OD <sub>492</sub>	DT (h)	Maximum OD <sub>492</sub>	DT (h)	Maximum OD <sub>492</sub>	DT (h)	Maximum OD <sub>492</sub>	DT (h)	Maximum OD <sub>492</sub>	DT (h)	Maximum OD <sub>492</sub>	DT (h)	Maximum OD <sub>492</sub>
14°C	31H	5,00	0,945	3,31	1,121	3,39	1,155	3,15	1,047	3,50	1,128	3,16	1,143	7,03	0,946
18°C	24H	2,37	1,201	1,85	1,220	1,95	1,214	2,00	1,120	1,79	1,178	2,42	1,245	3,54	1,010
21°C	24H	2,19	1,231	1,93	1,204	2,30	1,207	2,50	1,167	2,83	1,201	2,52	1,223	3,97	1,061
27°C	24H	0	0,012	0	0,024	0	0,033	0	0,06	0	0,021	0	0,021	0	0,031