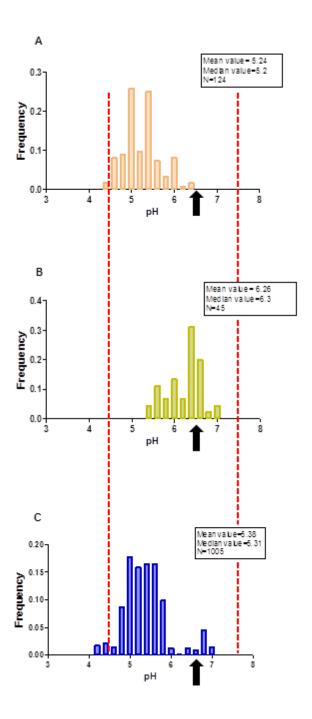
Supplementary Material SM1-to-SM3

SM1 . **Distributions in the** pH**-values among aquatic environments.** Distributions are given for three BU endemic areas: (A) Akonolinga, Cameroon; (B) Bankim, Cameroon and (C) French Guiana. (From the data published in Garchitorena *et al.* 2015a and Morris *et al.* 2014, 2016). Dotted lines delineate the range of pH values used in our experiments. The black arrow indicates the pH value of control medium 7H9.



SM 2. Statistics driven from the maximal (A) and minimal (B) linear mixed models used to fit the number of *Mycobacterium ulcerans* cells. Models are transcribed in *R* language. In both cases, samples were declared as explanatory variables with random effects. Statistics in bold characters are those that are significant at 5% C.I.

(A) Maximal model:

Ime (number of bacteria ~ pH×Chitin×Time+ pH×Chitin+pH×Time+ Chitin×Time + Chitin + Time + pH, random=~1|Sample,data=data,method="ML)

Explanatory variable	F- value	<i>p</i> -value
Time	13.077	<0.001
рН	1.632	0.161
Chitin	4.305	0.046
<i>pH</i> ×Chitin	1.980	0.088
<i>pH</i> ×Time	1.820	0.026
Chitin×Time	9.724	<0.001
<i>pH</i> ×Chitin×Time	0.899	0.593

(B) Minimal model:

 $Ime(Numberbacteria^{\sim} pH \times Chitin + pH \times Time + Chitin \times Time + Chitin + Time + pH, random=^1|Sample, data=data, method="ML")$

Explanatory variable	F- value	p-value
Time	13.261	<0.001
рН	1.610	0.167
Chitin	4.679	0.038
<i>pH</i> ×Chitin	2.008	0.084
<i>pH</i> ×Time	1.845	0.021
Chitin×Time	9.861	<0.001

SM 3. Changes in the number of *Mycobacterium ulcerans* cells within three pH groups. The three pH groups are defined as follows: pH low (4.5, 5.0, 5.5); pH middle (6.0, 6.5) and pH high (7.0, 7.5). Each color line represents a different time (t = 0, 25, 35 and 50).

