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Influence of chlorine discontinuities on the bacteria present in biofilm and water of a drinking water distribution system

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Introduction

Water distribution networks seldom function in stationary conditions. According to many parameters (quality of the resource, chlorine rate applied, technical interventions...), networks are constantly subjected to hydraulic, biological as physicochemical discontinuities. This study concerned disinfectant discontinuities (bleach). Few works already published on this topic and limited to discontinuities "all-or-nothing", have thus identified a risk of unsteadiness of the distributed water quality. To our knowledge, no published data exists on discontinuities with a "high and low" sequence of chlorine concentration and none have explored their

Objective

The objective is to appreciate, in drinking water and biofilm, the impact of chlorine discontinuities (changing from 0.1 to 0.4 then 0.1 mg Cl₂/L of residual chlorine) on:

- the total number of bacteria,
- the viability of bacteria (membrane integrity and culturability),
- the phylogenetic composition of bacteria (α, β et γ-proteobacteria).

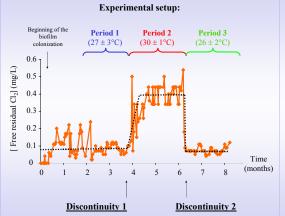
Methods

The experimental distribution system (NanCIE):



Continuous feeding with re-chlorinated drinking water in order to reach 0.1 or 0.4 mg Cl₂/L in the water at the outlet of the distribution system

Cement-lined cast iron Flow rate = 10 L/h240 L HRT = 24 h



(0.1→0.4 mg Cl₂/L)

(0.4→0.1 mg Cl₂/L)

Microbiological analysis:

- · Regular analysis of water during the three periods
- · Punctual analysis of biofilm: - before discontinuity 1 - 24d and 64d after discontinuity 1

 - 18d and 54d after discontinuity 2

Microbial parameters:

- Total number of bacteria (Syto 9 staining)
- Viable bacteria (with a membrane integrity) (kit BacLight)
- Culturable bacteria (PCA, 22°C, 15days)



Phylogenetic composition (FISH): (adapted from Batté et al., 2003)

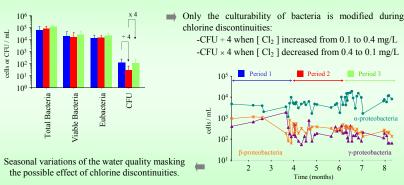
- Eubacteria (probe EUB338)
- α-proteobacteria (probe ALF1b)
- β -proteobacteria (probe BET42a)
- γ-proteobacteria (probe GAM42a)



Characteristics of the drinking water at in inlet of the distribution system (average of the 3 periods)

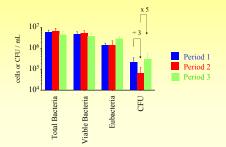
Parameter (/mL)	Mean ± SD	n
Total Bacteria	$1.9\ 10^5\pm1.0\ 10^5$	44
Viable Bacteria	$2.2\ 10^4\pm2.3\ 10^4$	43
CFU (PCA, 22° C, 15d)	$4.7\ 10^{-1}\ \pm\ 8.0\ 10^{-1}$	47
Eubacteria	$3.4\ 10^4\pm 2.3\ 10^4$	41
α -proteobacteria	$1.4\ 10^4\pm 9.8\ 10^3$	39
β -proteobacteria	$3.9\ 10^2\pm 6.0\ 10^2$	45
Y -proteobacteria	$2.1\ 10^2\pm2.6\ 10^2$	45

Effect of the chlorine discontinuity on the bacterial quality of water



→No significant effect of Cl₂ discontinuities on the phylogeny of bacteria present in drinking water

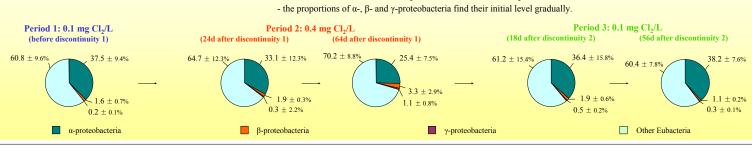
Effect of the chlorine discontinuity on the biofilm



- → The increase of the residual chlorine concentration in distributed drinking water from 0.1 to 0.4 mg Cl₂/L:
 - alters the culturability of biofilm bacteria (CFU ÷ 3 after discontinuity), no modification of the total number of bacteria.
 - changes the phylogenetic composition of biofilms: \triangledown of the β and γ -proteobacteria and \square of the α -proteobacteria.

⇒While remaining a minority sub-group, the proportion of γ-proteobacteria, including many bacteria of sanitary interest (coliforms, Legionella, Pseudomonas, ...), increased by a factor 6

- → This effect is temporary because the signals are reversed when the Cl₂ concentration is again brought back to 0.1 mg Cl₂/L:
 - the culturable bacteria increased by a factor 5 after discontinuity.



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