

A new method of screening cooling water and process water for Legionella pneumophila

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Colophon

Title

A new method of screening cooling water and process water for Legionella pneumophila.

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Client

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This report is public; all names of companies and laboratories involved are made anonymous.

Summary

The industry needs a fast and reliable analytical method for *Legionella* detection or monitoring, to enable optimal management and control of *Legionella* concentrations in cooling water and process water systems.

Taking the latest developments in the field of analysis as the starting point, over a 6-week period this study used 10 different analytical techniques to examine a total of 52 samples from 15 cooling water systems and 4 process water systems for the presence of *Legionella* and/or *Legionella pneumophila*. The main aim of the study was to test and evaluate the usefulness of an analytical scheme for cooling water and process water samples, which involved using Q-PCR analysis for *Legionella pneumophila* (in accordance with draft NEN 6254) and following this in the case of a positive result with a specific cultivation method for *Legionella pneumophila* (based on NEN 6253 with MWY medium). KWR developed both methods.

The other project objectives were:

- Development of a standard monitoring protocol that enables a representative sample to be taken of a complex cooling water or process water system.
- Comparison of the analysis results obtained using the aforementioned analytical techniques with the results of similar Q-PCR techniques (Applied Biosystems, according to protocol v.2.0; Pall Genesystems according to AFNOR XPT90-471) and other cultivation methods for *Legionella* (according to NEN 6265, NEN 6265:2007 with MWY medium and ISO 11731) and *Legionella pneumophila* (according to draft NEN 6253).
- Testing and evaluating the accuracy of a qualitative field test on the basis of an immunochromatographic assay for *Legionella pneumophila* sero-group 1 (Nalco, FastpathTM).

The sampled process water systems were systems in the paper industry. This process water has a high concentration of undissolved substances, organic matter and micro-organisms. This makes performing a specific microbiological assessment in the field particularly difficult, for example one involving a *Legionella* sp. cultivation method. As the conditions in a paper factory appear to be favourable for the growth of legionella bacteria, the paper industry also needs a robust analytical technique to provide greater clarity about the presence of these micro-organisms in process water.

The following method of working was adopted in this study:

- Each participating company selected 3 cooling water or process water systems with a demonstrable 'legionella history' (select sample).
- Each participating company assigned responsibility to one or more people for taking samples in accordance with the protocol, which KWR provided (appendix I), and for registering the current operating data for the selected systems (appendix II).
- All the participating companies were given 27 sterilised 1-litre bottles with a standard content comprising a thiosulphate solution for neutralising oxidising biocides and an NTA solution for complexing heavy metals.
- The company then sampled each cooling water or process water system on three different days, after which it sent the samples to KWR by courier. Three sample bottles were filled for each system. A note of the current operating data for each system was made on the information sheet.
- The contents of the three bottles with samples from the same system were homogenised by KWR on the same day and divided into new sample bottles. These sample bottles were sent to the participating laboratories by courier on the same day (see table 2).
- Analysis started the day after sampling.
- The participating laboratories sent the analysis results to KWR for further processing.

The analysis results have been processed and partially tested statistically. This has led to the following conclusions and recommendations:

- This study has demonstrated the usefulness and added value for analysing cooling water and process water samples in accordance with the analysis scheme based on using Q-PCR analysis for *Legionella pneumophila*, followed in the case of a positive result by a specific cultivation method for *Legionella pneumophila*. Besides providing managers of cooling water and process water systems with information faster, this new screening methodology also provides them with more specific information on the presence of *Legionella pneumophila* in the system. The choice of the specific Q-PCR for *Legionella pneumophila* in the scheme does not affect the result. The choice of the specific cultivation method should be based on the recommendations of this study and possibly the results of a further study.
- Variance analysis showed for the cooling water and process water samples used in this study that there were no significant statistical differences between the results of three Q-PCR techniques for detecting *Legionella pneumophila* performed by three different laboratories in accordance with three different standards.
- In most cases, the results of the Q-PCR techniques for the cooling water samples studied meet the requirement for a detection limit of at least 1,000 units/litre, based on table 12 of Health & Safety Information Sheet AI-32. If using Q-PCR technology systematically leads to more units per litre being detected than the number detected using the cultivation method, table 12 of the Health & Safety Information Sheet will have to be expanded for the interpretation of Q-PCR results. With regard to the specific situation at a company, when switching to Q-PCR, the method should preferably be used in parallel for a while with the cultivation method normally used up to that point, so that the results of the old and new method can be correlated. This can then be used as a basis for adjusting the action levels in the legionella control plan.
- The practical usefulness of the analysis scheme based on Q-PCR, followed by a cultivation method specifically for *Legionella pneumophila*, may be limited by the observation that using Q-PCR for the specific detection of *Legionella pneumophila* in the cooling water samples analysed does not lead to less positive samples than detection based on a cultivation method for *Legionella* in total. In comparison with drinking water systems, *Legionella pneumophila* is apparently fairly generally present in cooling water systems.
- This study has shown that applying NEN 6265 with BCYE medium for cooling water and process water samples leads to unsatisfactory results, owing to disruptive additional growth on the culture medium. NEN 6265 was revised for this in 2007 by also describing the MWY medium for samples with excessive additional growth.
- This study has shown that pretreating cooling water and process water samples with acid in accordance with ISO 11731:1998(E) and counting more frequently on day 3 and day 5 after incubation, leads to fewer outcomes for which no result can be stated on account of additional growth on the culture medium.
- Variance analysis has shown for the cooling water and process water samples used in this study that there are no significant statistical differences between the results of the cultivation method according to ISO 11731 and the cultivation method according to NEN 6265:2007 using an MWY culture medium, provided that the agar plates are also assessed on day 3 and day 5.
- Taking into account the method's limitations, the FastpathTM method is a useful addition to the set of instruments available to the process operator responsible for the daily management of cooling water systems. The method's main benefits are speed, simplicity and the possibility of on-site implementation. However, it is important to realise that specificity in respect of *Legionella pneumophila* sero-group 1 means that the other sero-groups are not taken into account. (Nevertheless *Legionella pneumophila* sero-group 1 is the main cause of legionella pneumonia; up to 90% of registered cases worldwide). Moreover, the information is of such a qualitative nature that it cannot replace the measurements that are normally made.
- This study has not provided any indications of the presence of high concentrations of legionella bacteria in process water samples from the paper and cardboard industry. Nevertheless, the results indicate that *Legionella* or *Legionella pneumophila* can be detected, at least in a number of samples.
- It was notable for practically every cooling water system examined in this study that disinfection hardly ever consistently resulted in a legionella bacteria concentration of less than 1,000 cfu/l.
- The laboratories participating in this study have demonstrated that it is possible to achieve reliable results using a cultivation method for *Legionella* and/or *Legionella pneumophila*. A contributory factor

to this was that these laboratories were experienced and did everything possible to achieve good analytical results, without attempting to cut costs or avoid difficulties. The interlaboratory comparisons organised by KWR also confirmed that a certain degree of reliability can be achieved using a cultivation method.

Recommendations:

- It is recommended on the basis of the experiences from this study that the specific cultivation method for *Legionella pneumophila*, with an MWY cultivation medium based on NEN 6265:2007 and the draft version of NEN 6253, should be further optimised by means of pretreatment with acid and more frequent counting.
- As good results in this study were obtained using the cultivation method according to ISO 11731, it would be advisable to examine the extent to which the method could also be made suitable for the specific cultivation of *Legionella pneumophila*.
- A further recommendation assuming a sufficiently large non-select sample of industrial cooling water systems is that the two specific cultivation methods for detecting *Legionella pneumophila* should be checked both separately and in combination with Q-PCR assessment to determine their performance in terms of yield and cost.
- It would be advisable to set up interlaboratory comparisons for the Q-PCR techniques, as is done for the cultivation method.
- The cultivation method in accordance with ISO 11731 is recommended for detecting *Legionella* in process water from the paper and cardboard industry, subject to pretreating the sample with acid and reading the culture medium counts more frequently after incubation.
- A further examination of the role that 'full stream' or 'side stream' filtration plays in the effectiveness of disinfection in cooling water systems is recommended. Relevant variables in this are the type of disinfectant and the dose and method of dosing (continuous/discontinuous).

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1 Introduction

1.1 Background

Employer's obligations under the Dutch Working Conditions Act include the requirement to take appropriate measures to prevent employees from being exposed to legionella bacteria. The introduction of health and safety policy rule 4.87-1 obliged employers to conduct a hazard identification and risk assessment (HIRA) for possible growth of *Legionella* in their cooling water and process water systems. In practice, the HIRA often results in management and control measures in the systems concerned. Experiences at Corus and Dow in recent years showed that the complex matrix of cooling water makes it particularly difficult to monitor *Legionella* in cooling water using the existing cultivation method in accordance with NEN 6265 (as described in the policy rule, as well as the alternative cultivation method according to ISO 11731). Moreover, there are major differences between the analysis results of different laboratories and the result can only be determined after at least 7 days.

The industry clearly needs a fast and reliable analytical method for *Legionella* detection or monitoring, to enable optimal management and control of *Legionella* concentrations in cooling water and process water systems. On the grounds of the limit values stipulated in Health & Safety Information Sheet (AI-32)⁵ for *Legionella* in cooling water, an analytical method of this kind should have a detection limit of at least 1,000 legionella bacteria per litre.

Three recent developments at KWR concerning the analysis of legionella bacteria were recently extensively reported in the journal $H_2O^{[1][2][3]}$:

- The newly developed quantitative real-time PCR method (Q-PCR) enables the concentration of *Legionella pneumophila* in a water sample to be measured within a few hours^[1]. The Q-PCR method's reproducibility is better than that of the standard cultivation method. Chemical disinfection reduced the concentration of *L. pneumophila*, by more than 90 percent but thermally disinfecting mains water had hardly any effect on the results of the Q-PCR method. Correctly interpreting the results obtained therefore requires information on the water sample's origin/history.
- A cultivation method was developed at KWR that specifically demonstrates the presence of *Legionella pneumophila* on the basis of an increase in pH in the culture medium and a higher incubation temperature^[2].
- Existing information has shown that *Legionella pneumophila* is by far the most important cause of legionella pneumonia. The researchers therefore concluded in a third article that the policy in the Netherlands should primarily focus on combating *Legionella pneumophila* in mains water/water systems^[3].

A fourth development concerns an amendment of the existing standard NEN 6265 for the analysis of *Legionella*. A revised version of NEN 6265 was published in October 2007, and included a modified method of working for "isolating *Legionella* in a sample expected to contain disruptive flora" (such as cooling water and process water). In addition to making use of the usual BCYE medium, this method of working uses plates with an MWY medium ¹.

The selective detection methods for *Legionella pneumophila* (Q-PCR and cultivation) make it possible to specifically tackle this organism in water systems. The most obvious use of these detection methods in practice is for rapidly screening a series of samples (e.g. from cooling tower systems) using Q-PCR, possibly followed by the selective cultivation method for *Legionella*

¹ MWY = Modified Wadowsky Yee Agar; BCYE = Buffered Charcoal Yeast Extract Agar

pneumophila, if *Legionella pneumophila* is detected in excess of a certain concentration (see scheme below).



Figure 1. Proposed scheme for Legionella pneumophila screening in cooling and process water.

If the scheme shown in figure 1 proves to be economically and technically successful, it could provide an alternative to present cultivation methods referred to in documents such as the Health & Safety Information Sheet⁵ (also referred to as AI-32) as standard methods for sampling and analysing *Legionella* in cooling water systems. It should be noted in connection with this that the government has not yet taken the step from detection of all viable varieties of legionella to specific detection of *Legionella pneumophila*.

The results of an analysis to detect *Legionella* or *Legionella pneumophila* may have major consequences that affect a factory's business operations and the environment. It is essential in connection with this to have a representative water sample from the water system being investigated. In practice, the analysis and analysis results receive a great deal of attention but not enough attention is paid to the quality of sampling. It is difficult to take a representative sample in a large cooling water or process water system. A standard sampling protocol is therefore required, in which the best options for the time, place and method of sampling are laid down. Moreover, to enable interpretation of the analysis results, it is important to keep a meticulous record of the conditions under which samples were taken.

Besides a reliable analytical technique that can be used for testing against statutory frameworks and company-specific guidelines, operators in the industry also need a fast, simple and cheap method for daily system monitoring. In practice, ATP measurements and colony number measurements (dip slides for example) are often used for this, both of which provide an idea of the system's general microbiological state. The immunochromatographic assay is a new development in this field and can be used for qualitative detection (present/absent) of *Legionella pneumophila* sero-group 1 (*Lp* SG1). This method is fast, simple and can be used on site. A major advantage of using this method is that it provides specific information on the presence of the most relevant pathogen of the *Legionella* genus. The immunochromatographic assay thereby offers a potentially useful addition to the operator's set of instruments.

1.2 Project objectives

The project objectives are:

- Development of a standard monitoring protocol that enables a representative sample to be taken of a complex cooling water or process water system.
- Testing and evaluating the usefulness of the scheme in figure 1 for cooling water and process water samples.
- Comparison (at KWR) of the analysis results obtained using the analytical techniques shown in figure 1 with the results of similar Q-PCR techniques and cultivation methods at other laboratories.
- Testing and evaluating the accuracy of the qualitative immunochromatographic assay for *Legionella pneumophila* SG 1 in cooling water and process water.

1.3 Companies and laboratories involved

This project was initiated as a multi-client project in which the aim was for all the project's industrial participants to supply cooling water and process water samples from their own systems. Table 1 provides a list of companies that provided samples for this project.

Company	Sample types
А	cooling water from 2 systems and process water from 1 system at the same site
В	cooling water from 3 systems at the same site
С	cooling water from 3 systems at the same site
D	cooling water from 4 systems at various sites
Е	cooling water from 3 systems at the same site
F	process water from 3 different systems at 3 different sites

Table 1 List of companies involved in this study

The Royal Association of Dutch Paper and Cardboard Manufacturers (Koninklijke VNP) represents 21 paper and cardboard manufacturers in the Netherlands. These factories make repeated use of their process water. This results in water with a high concentration of undissolved substances, organic matter and micro-organisms. It is therefore particularly difficult to perform a specific microbiological assessment, for *Legionella*, for instance. Besides it being difficult to filter sufficient water for detection purposes, there is also often excessive additional growth on the culture medium. Conditions in a paper factory (high temperatures, much aerosol formation, high humidity) constitute all the reasons for monitoring the process water for *Legionella*. VNP is consequently extremely interested in evaluating new methods, such as the ones that are the subject of this project.

Table 2 provides a list of laboratories participating in this study. Chapter 2 provides a more detailed description of the methods used.

Laboratory	Method(s) used
Lab I	cultivation methods related to NEN 6265 and ISO 11731
Lab II	Q-PCR Pall Genesystems
Lab III	Q-PCR
Lab IV	Q-PCR
	cultivation methods related to NEN 6265
	Nalco Fastpath™ (immunochromatographic assay)

Table 2 List of laboratories involved in this project

2 Methods

2.1 Sampling protocol

A small committee of representatives from the industry and cooling water technology experts drew up a sampling protocol for sampling cooling water and process water systems in aid of legionella analysis. The complete protocol is included in appendix I. The protocol includes an information sheet (list for completion) for recording current data on the sampled system. This information sheet is included in appendix II.

The protocol was primarily intended for sampling within the scope of this project and was distributed to the participating companies. It can also be seen as a general sampling protocol for cooling water and process water systems.

In summary, the following recommendations apply to sampling cooling water systems:

- Discontinuously disinfected cooling water systems should be sampled at the end of the disinfection cycle, which means just before introducing a new dose of disinfectant. The time of sampling is irrelevant in systems that are continuously disinfected. However, the position where samples are taken is important. Samples should preferably be taken just before the location of the dosing point.
- The following preferred order for the position where samples are taken should be used when sampling cooling water systems:
 - Sampling of the falling water in the free space above the cooling tower basin (*at least 1 metre from the side and using special tools*).
 - Sampling of the water in the cooling tower basin from a position as close as possible to the recirculation pump's suction pipe (*at least 1 metre from the side and within 10 to 20 cm below the water's surface*).
 - Sampling from a sampling tap in a main pipe on the outlet side of the recirculation pump, as close as possible to the cooling tower (*tap-out pipe should preferably not be made of rubber or plastic, should be shorter than 3 metres and should be flushed for 3 minutes before samples are taken*).

The starting point for the preferred order is that the sample should represent the quality of the cooling water as far as possible at the point where it is sprayed and could be spread into the surrounding area as an aerosol. Spraying of cooling water takes place above the cooling section, which is generally an unsuitable place for sampling. The positions suggested here are intended to provide samples that are as close as possible to the aforementioned representative quality. Wall effects, namely factors which affect the quality of the sample owing to a biofilm or sediment (in the basin or in a pipe), are a key factor here and should be prevented as far as possible. This is the reason for the chosen distances to the wall of the cooling tower and the flushing time for the sampling point.

- It is particularly important to ensure homogeneous water distribution across the cooling tower when directly sampling falling water. This is also a standard precondition for a cooling tower to operate properly and is therefore in practice an important point for the operator's attention. Poor distribution is an indicator of local clogging in the cooling section, which reduces the system's cooling capacity and increases the likelihood of legionella bacteria growing.
- It is important to keep a record of the conditions under which samples were taken and especially of the disinfection regime.

In the case of process water systems, large variations in the types of systems make it impossible to provide standard recommendations for the best place to take samples. A few recommendations are provided in appendix II for the paper and cardboard industry and for the steel industry. The following general recommendations apply for process water flows:

• Discontinuously disinfected process water systems should be sampled at the end of the disinfection cycle. The time of sampling is irrelevant in systems that are continuously disinfected. However, the position where samples are taken is important. Samples should preferably be taken just before the location of the dosing point.

• It is important to keep a record of the conditions under which samples were taken and especially of the disinfection regime.

2.2 Analytical methods

2.2.1 Summary of analytical methods used in this study

Table 3 provides a list of the analytical methods and associated standards used in this study.

Analytical technique	Standard	Laboratory that performed the work in this study
cultivation method for <i>Legionella</i> with BCYE medium	in accordance with NEN 6265	Lab IV
cultivation method for Legionella	in accordance with ISO 11731	Lab I
cultivation method for <i>Legionella</i> with MWY medium	in accordance with NEN 6265:2007	Lab IV
	based on NEN 6265:2007 (with more frequent reading)	Lab I
cultivation method for <i>Legionella pneumophila</i> with BCYE medium	in accordance with draft version of NEN 6253	Lab IV
cultivation method for <i>Legionella pneumophila</i> with MWY medium	based on NEN 6265:2007 and draft version of NEN 6253	Lab IV
Q-PCR for Legionella pneumophila	in accordance with draft version of NEN 6254; growth of a fragment of the <i>mip</i> gene (DNA) with PCR and specific primers.	Lab IV
Q-PCR for Legionella pneumophila	Applied Biosystems. Detection and Quantification of <i>Legionella spp</i> . and <i>Legionella pneumophila</i> . In accordance with protocol v.2.0	Lab III
Q-PCR for Legionella pneumophila	Pall Genesystems. Quantitative PCR for <i>Legionella pneumophila</i> . Validated by AFNOR in accordance with XPT90-471 standard. Additional interpretation of the PCR result in accordance with draft version of NEN 6254.	Lab II
Fastpath TM	Nalco	Lab IV

Table 3 List of the analytical methods used in this project

FastpathTM is a qualitative field test especially for detecting *Legionella pneumophila* sero-group 1. The method is based on a chromatographic test that uses an immunoassay in which a reaction occurs between antibodies on the detection equipment and antigens of *Lp* SG1. The method uses a test strip on which the preconcentrated sample has to be placed. The sample has to be pretreated by filtration to obtain a sufficiently low detection limit. The test strip has a test band and a control band. The control band always indicates a red line; the test band shows a second red line for the presence of *Legionella pneumophila* sero-group 1 in excess of the detection limit. The detection limit is 100 cfu/l following pretreatment of 250 ml of the sample by filtration and 100,000 cfu/l without filtration. All pretreatment and detection requirements were provided by Nalco. Laboratory technicians have been trained to use the method.

2.2.2 Differences in performing the various cultivation methods

The analytical method described in NEN 6265 is generally used for *Legionella* detection in water/drinking water samples in the Netherlands. The method requires use of a specific culture medium (BCYE = Buffered Charcoal Yeast Extract). This culture medium is known to lead to problems when used for water samples with a high concentration of disruptive flora.

The internationally used ISO 11731 method is based on the same type of medium but with the addition of various other substances, the names of which are indicated in the name of the medium (GVPC = Glycine Vancomycine Polymixine Cycloheximide agar).

The KWR study of 2005^[4] showed that for samples with a high concentration of disruptive flora – such as samples of cooling water and process water – additional growth is best inhibited by using MWY medium (MWY = Modified Wadowsky Yee agar) as the culture medium. This medium was consequently introduced in 2007, in the revised version of NEN 6265.

Besides the culture medium's composition, the pH value and incubation temperature are also important parameters, in connection with the *Legionella* yield in the analysis. Research has established that the variation in pH and incubation temperature can be used to make the cultivation method specific for detecting *Legionella pneumophila*^[2]. With respect to NEN 6265:2007, the culture medium's pH is increased from 6.9 ± 0.1 to 7.3 ± 0.5 and the incubation temperature from 36 ± 2 °C to 40 ± 0.5 °C. The method for the specific analysis of *Legionella pneumophila* has been set out in the draft version of NEN 6253.

When performing the method set out in NEN 6265:2007, using MWY medium, lab I adopted a method that differed in the following way from the usual standard:

• The culture media were assessed after 3, 5, 7 and 10 days. Standard practice is to count the colonies on a culture medium after an incubation period of at least 7 days. Reading the culture media more frequently makes the method more labour intensive and more expensive but may offer the advantage of better assessment, as additional growth is still limited.

2.2.3 Differences in performing the three Q-PCR techniques

The quantitative PCR method, referred to simply as Q-PCR, is based on the Polymer Chain Reaction, in which a specific DNA fragment can be amplified to produce large numbers of copies under the influence of thermal cycles and with the aid of enzymatic reactions. Specific amplification is made possible by using short synthetic DNA molecules, known as primers. The base sequence or DNA sequence in the primers is chosen so that it binds selectively to the DNA, in this case of *Legionella pneumophila*. An enzymatic chain reaction occurs in the PCR once the primers bind to the DNA obtained from the water sample and the DNA fragment is amplified between the primers during each temperature cycle. Amplification can only occur if the sample contains this specific *L. pneumophila* DNA. In Real-time PCR, the formation of the amplified DNA is measured during each cycle (real-time). A synthetic DNA molecule known as the probe is used for this and is labelled with a fluorescent dye. The fluorescence occurs after the DNA has formed. Quantification is possible because there is a clear link between the time in the reaction at which it becomes possible to detect the DNA fragment that has formed (C_T value) and the *L. pneumophila* DNA concentration in the sample.

2.2.3.1 Q-PCR according to the draft version of NEN 6254

The Q-PCR method according to the draft version of NEN 6254 was developed and validated at KWR. Detection of *L. pneumophila* is carried out by specific primers that amplify a small fragment of the *mip* gene. The possibility of inhibiting the PCR and yield of DNA isolation is quantified by adding control DNA to every sample. The control is quantified at the same time in a multiplex PCR. The quantitative result of *L. pneumophila* detection is corrected using the yield produced by the control. Around 20 samples can be analysed per analysis.

2.2.3.2 Q-PCR according to Pall Genesystems Technology

The Q-PCR method is based on a method that uses a completely standardised PCR system comprising a DNA extraction, a specific PCR machine and a gene disc that includes everything necessary for the specific DNA amplification. Besides detection of *L. pneumophila* and/or *Legionella* spp., the detection

method also includes various quality controls, such as an external quantitative positive control. No information is available on the DNA fragment that is amplified, the DNA sequence of primers or the probe. Six or 12 samples can be analysed simultaneously per analysis run. The method was validated by AFNOR. The interpretation was also performed (limit of detection (LOD) and lowest quantity determinable (LQD)) in accordance with the draft version of NEN 6254 when the results in this study were evaluated.

2.2.3.3 Q-PCR according to the Applied Biosystems detection kit

Applied Biosystems developed and marketed the Q-PCR method. The Q-PCR detection and quantification kit for *L. pneumophila* includes an internal positive control and Amperase UNG. Amperase UNG treatment prevents the reamplification of PCR DNA fragments that were formed in earlier PCR experiments. This method also lacks information on the DNA sequence of primers, probes and controls.

2.3 Method used in the study

As described in section 1.3, the project's participants supplied samples form 15 cooling water systems and 4 process water systems. KWR collected and homogenised the water samples and then distributed them to the participating laboratories for further analysis.

The following method of working was used for this:

- Each participating company selected 3 cooling water or process water systems with a demonstrable 'legionella history' (select sample).
- Each participating company assigned responsibility to one or more people for taking samples in accordance with the protocol, which KWR provided (appendix I), and for registering the current operating data for the selected systems (appendix II).
- All the participating companies were given 27 sterilised 1-litre bottles with a standard content comprising a thiosulphate solution for neutralising oxidising biocides and an NTA solution for complexing heavy metals.
- The company then sampled each cooling water or process water system on three different days, after which the samples were sent to KWR by courier. Three sample bottles were filled for each system. A note of the current operating data for each system was made on the information sheet.
- The contents of the three bottles with samples from the same system were homogenised by KWR on the same day and divided into new sample bottles. These sample bottles were sent to the participating laboratories by courier on the same day (see table 2).
- Analysis started the day after sampling.
- The participating laboratories sent the analysis results to KWR for further processing.

The samples were consecutively coded from OPIW1 to OPIW54. The contents of sample bottles OPIW 1, OPIW 2 and OPIW 3 from company D were mistakenly mixed with each other while the work was underway. The resulting sample was further analysed as OPIW 1 and codes OPIW 2 and OPIW 3 were removed from the overviews.

The study led to a results matrix for 52 samples analysed using 10 analytical techniques, which produced 520 results. The matrix is shown in appendix III.

2.4 Comparison of the analysis results of the various analytical methods

A more in-depth statistical examination of some parts of the study was conducted by a statistician with the aid of variance analysis (ANOVA method, see appendix VI).

International standard ISO 16140 was also used for a qualitative comparison of the analysis results. The aforementioned ISO standard includes a protocol which describes how alternative microbiological

analytical methods can be compared with a reference method. The following parameters and definitions are important for this:

- Relative accuracy (AC). Degree of accordance between the result of the reference method and the result obtained with the alternative model using the same samples.
- Relative sensitivity (SE). Ability of the alternative method to analyse for the parameter concerned when it is detected by the reference method.
- Relative specificity (SP). Ability of the alternative method <u>not</u> to analyse for the parameter concerned when it is <u>not</u> detected by the reference method.

These parameters can be calculated using a results matrix, as shown in table 4, and using the following formulas:

- AC = (PA + NA)/N * 100%
- SE = $PA/N_{+} * 100\% = PA/(PA + ND) * 100\%$
- $SP = NA/N_* 100\% = NA/(NA + PD) * 100\%$

Table 4. Results matrix for the qualitative comparison of various analytical methods

Results	Reference method	Reference method
(total N samples)	positive (N+ samples)	negative (N. samples)
Alternative method	+/+ number of samples with a	-/+ number of samples with a
Positive	positive accordance (PA)	positive deviation (PD)
Alternative method	+/- number of samples with a	-/- number of samples with a
Negative	negative deviation (ND)	negative accordance (NA)

3 Results and discussion

3.1 General

The main aim of this study was to test and evaluate the usefulness of the scheme in figure 1 for cooling water and process water samples. It essentially involves using Q-PCR for *Legionella pneumophila*, possibly followed by a cultivation method specifically for *Legionella pneumophila* to confirm the viability of the bacteria.

Appendix IV includes a summary of the results achieved after applying the analytical methods referred to in figure 1, on the understanding that the specific cultivation method according to the draft version of NEN 6253 was used in every case and not only when Q-PCR produced a positive result. The results can be summarised as follows:

33 of the 52 samples (63.5%) showed a positive result after using Q-PCR in accordance with the draft version of NEN 6254.

- A positive result was also obtained for 16 of the 33 corresponding samples when the specific cultivation method was used.
- Excessive additional growth on the agar plate meant that no result could be stated for 9 of the 33 corresponding samples when the specific cultivation method was used.
- However, DNA copies were found for 8 of the 33 corresponding samples when Q-PCR was used but no viable bacteria were detected using the specific cultivation method (< detection limit).
- The result for both Q-PCR and the cultivation method was below the detection limit for 11 of the 52 samples (21%).
- No Q-PCR could be stated for 5 of the 52 samples (9.6%) because the yield of the internal control was below the limit value for reliable analysis of the sample.
- For 3 of the 52 samples (5.8%), the result using Q-PCR was below the detection limit and too much additional growth on the culture medium meant that no result could be stated for the cultivation method.
- No false negative results were obtained using the Q-PCR method, which means that no DNA copies were found using Q-PCR but colonies were found on the culture medium when using the cultivation method.

The case in which 8 of the 33 results were positive using Q-PCR and negative using the cultivation method indicates that, although there may not have been any viable *Legionella pneumophila* bacteria in the sample at the time of sampling, it nevertheless contained insufficiently lysed legionella bacteria which would still be detected. (The DNA of dead legionella bacteria that has completely disintegrated or lysed is fairly present in the solution, so it passes through the filter during filtration and no longer affects the PCR result.) This is a typical result after using the scheme in figure 1 when thermal or chemical disinfection has been used in the sampled system. In such situations it indicates effective disinfection but also the presence of legionella bacteria during certain periods (such as between disinfection operations) or in certain parts of the system. The latter may apply in the case of continuous disinfection, for example. The overview in appendix IV, which also includes a summary of the disinfection method used per system, shows that this is indeed the case in systems that undergo discontinuous disinfection by means of shock dosing and in systems that are continuously disinfected. Table 5 shows a simplified scheme with a proposal for a general qualitative interpretation of the screening according to figure 1.

Q-PCR	specific cultivation	interpretation
	method	
< detection limit	< detection limit	The sampled water contains no detectable <i>Legionella</i> <i>pneumophila</i> or its remains. Depending on the specific circumstances (size of the system; representativeness of the sample) this is an indication of a microbiologically stable system in which no <i>Legionella</i> growth occurs.
positive result	< detection limit	The sampled water contains no viable <i>Legionella</i> <i>pneumophila</i> in excess of the detection limit but there is an indication that growth can nevertheless occur in the system at certain times (periods between two shock doses, for example) or at certain positions (outside the range of the sampling point and outside the range of continuous disinfection). Situation requires further study.
positive result	positive result	 The system contains viable <i>Legionella pneumophila</i> bacteria. The situation has to be assessed on the basis of established concentration levels according to applicable protocols. If disinfection is carried out, its effectiveness should be further ascertained.

Table 5 Scheme for interpreting results of the screening method according to figure 1

3.2 Comparison of Fastpath[™] with other techniques

The FastpathTM method was executed at lab IV using test kits provided by the supplier. During execution, it emerged that reading the test strip can actually be a point for discussion, in the sense that interpretation of some results involves a certain degree of subjectivity.

The results of FastpathTM - as an alternative method - were compared with those of the Q-PCR carried out by lab IV (see appendix V). As Q-PCR focuses on *Legionella pneumophila* and the immunoassay only on *Legionella pneumophila* sero-group 1, the interpretation also took into account the results of serotyping carried out by lab III, in the case of the ISO 11731 and NEN 6265 analysis. This makes it possible to determine whether, for example, a negative result of the FastpathTM method is correct.

The results can be summarised as follows:

- A total of 47 samples were assessed (there was no result for 5 samples when Q-PCR was used as the reference method).
- The Fastpath method gave a false positive result 4 times, a false negative result 5 times and a correct negative result 6 times. It should be noted in connection with this that two false positive results were obtained for samples from the paper industry, whereby the detection limit for Q-PCR was much higher.
- The relative accuracy (AC) of Fastpath[™] was 81% of that of Q-PCR.
- The method's relative sensitivity (SE) was 85%.
- The relative specificity (SP) was 71%.
- When determining these parameters, a correct negative result of FastpathTM was interpreted as a positive accordance (PA). A correct negative result arises when *Legionella pneumophila* is detected using Q-PCR, for example, but the serotyping indicates that sero-group 1 has not been found in the sample concerned.

In view of the good results obtained using the ISO method (see section 3.4), this cultivation method's results were also compared with those of FastpathTM (see appendix V).

The results can be summarised as follows:

- A total of 49 samples were assessed (no results for 3 samples with ISO method as reference method).
- The Fastpath method gave a false positive result 5 times, a false negative result 5 times and a correct negative result 10 times. The sample codes for the 5 false negative results correspond exactly with those obtained from the comparison with Q-PCR. In two of the 5 false negative results, relatively low concentrations were measured using the ISO method (200 and 530 cfu/l) but in the three other cases the concentrations were around 60,000 cfu/l.
- The relative accuracy (AC) of FastpathTM was 80% of that of ISO.
- The method's relative sensitivity (SE) was 86%.
- The relative specificity (SP) was 58%.
- When determining these parameters, a correct negative result of FastpathTM was interpreted as a positive accordance (PA).

On the basis of these results, the degree of accordance of Fastpath[™] with the two reference methods is approximately 80%. It is ultimately up to the users to determine according to their own criteria whether this level of accuracy is sufficient for using this fast screening method in practice. It is also important to realise that the method is limited to *Legionella pneumophila* sero-group 1; however, this is the main cause of legionella pneumonia (up to 90% of registered cases worldwide).

3.3 Comparison of Q-PCR techniques

Appendix VI provides an overview of the Q-PCR analysis results obtained by three different laboratories using three different methods. The main characteristics are summarised in the table below.

	Q-PCR lab IV	Q-PCR lab III	Q-PCR lab II
number of positive samples	33	26	47
• which were > 100,000	• 11	• 5	• 3
• which were > 1,000	• 22	• 19	• 30
• which were $\leq 1,000$	• 3	• 2	• 14
number of negative samples	14	22	5
(< detection limit)			
number of samples with no result	5	4	0

Table 6. Main characteristics of the results of the three Q-PCR techniques

No value could be determined for 5 of the Q-PCRs performed by lab IV, as the yield of the internal control was below the limit value for reliable quantitative analysis. However, in 4 of the 5 cases *Legionella pneumophila* was detected. In 4 samples in the Q-PCR performed by lab III the level of inhibition produced an unreliable result. No value is stated for these cases.

Figure 2 sets out the Q-PCR results of each of the three laboratories against each other in a double logarithmic diagram. An analysis result below the detection limit is shown here as the logarithm of half of the detection limit concerned (log{detection limit/2}). The good accordance between the three methods is striking. It is also notable that the Q-PCR at lab IV detected a higher number of copies more often than the other two methods. No immediate explanation is available for this but it may be an effect of the yield correction that was made.



Table 7 Results of the qualitative assessment of the three Q-PCR techniques vis-à-vis ISO 11731 as the reference method.

	Q-PCR lab IV	Q-PCR lab III	Q-PCR lab II
relative accuracy (AC)	91%	78%	82%
relative specificity (SP)	89%	82%	33%
relative sensitivity (SE)	92%	76%	97%

A comparison was made in accordance with ISO 16140 (see section 2.4) to enable an initial qualitative assessment of the results of the three Q-PCR methods and the results of the cultivation method according to ISO 11731 as a reference. When making comparisons it must be remembered that the Q-PCR is concerned with *Legionella pneumophila* and the ISO method with *Legionella as a whole*. The ISO method's serotyping was therefore also taken into account, to ascertain whether a negative Q-PCR result was correct. A correct negative result was then deemed to be a positive accordance (PA) in the qualitative comparison. Table 7 shows a summary of this qualitative comparison's results. The calculations shown in

tabular form in appendix VI only included those samples for which both analytical methods produced a result. Table 7 generally shows a high level of accordance between the Q-PCR results and the results obtained in accordance with the method of ISO 11731. Only the specificity of the Pall Genesystems Q-PCR was relatively low. This was because this method detected a quantifiable number of legionella cells in 92% of the samples examined, which is a relatively high figure (8% of the samples were below the detection limit).

A variance analysis was used to compare the results of the three different Q-PCR techniques with each other (see appendix VII). The zero hypothesis was tested in this analysis that the mean of the analysis results is the same for the three different Q-PCR methods. It is 95% certain that this hypothesis cannot be rejected. Statistically, this means that there are no significant differences between the three methods.

The detection limit of the various Q-PCR techniques was generally lower than 1,000 copies/litre for cooling water samples. It was only necessary to state a higher detection limit than that on a number of occasions when using the Q-PCR of lab III. The 1,000 cfu/l requirement stated in the Health & Safety Information Sheet could not be met in those cases. However, the question is whether the table in question, which was drawn up on the basis of the interpretation of results of the cultivation method, can still be used for interpreting Q-PCR results. The main point is whether using Q-PCR is a reason for significantly higher numbers of copies per litre with respect to the number of colony-forming units per litre. This is examined in figure 3 by setting out the average results of the Q-PCR methods against those of the specific cultivation method for *Legionella pneumophila* using MWY medium (in accordance with the draft version of NEN 6253). In both cases, values below the detection limit have been included in the graph with a value corresponding with half the value of that detection limit.



Figure 3. Comparison of the specific cultivation results for Legionella pneumophila, according to the draft version of NEN 6253 using MWY medium, and the average of the results of the three Q-PCR techniques. The finely dotted line shows the relationship between the numbers detected with the two methods that have been compared. The other line is a linear trend line.

An initial glance at figure 3 shows that there is a wide spread in the analysis results. The spread is partially accounted for by disinfection in the systems examined (see explanation in section 3.1). In any case, it is clear from figure 3 that in 94% of the samples DNA copies were found in larger numbers when using Q-PCR than the numbers of colony-forming units that were found when using the cultivation method. This is possibly explained by the fact that, unlike with the cultivation method, corrections are made for the detection yield when Q-PCR is used. Poor correlation makes it impossible to establish a fixed relationship.

3.4 Comparison of cultivation methods

A total of five different cultivation techniques were used in this study (see table 3 and appendix VIII). The method according to NEN 6265:2007 using MWY medium was applied by two laboratories (lab IV and lab I).

The results of the 'normal' NEN 6265 using BCYE medium were disappointing because no result could be provided for 31 of the 52 samples (60%) owing to excessive additional growth on the culture medium. This is as expected, given the complex matrix of the cooling water and process water samples and is also the reason why the MWY medium was included in the NEN standard. Poor results were also obtained with the cultivation method using BCYE medium with a higher pH culture medium (pH 7.3) and samples at a higher incubation temperature (40 °C), which was specifically for *Legionella pneumophila* analysis. In that case, no result could be stated for 34 of 52 samples (65%) owing to excessive additional growth on the culture medium.

Using MWY medium in accordance with NEN 6265:2007 considerably reduced the amount of disruptive additional growth. This analytical method was used by lab IV as well as lab I, whereby it should be noted that lab I also made counts on day 3 and 5. The summary in appendix VIII shows that there was too much additional growth on the culture medium for respectively 15 (29%) and 16 (31%) of the 52 samples. That is half of the figure vis-à-vis the results when BCYE medium was used. Figure 4 shows the results of the two assessment methods using MWY medium alongside each other. It should be noted here that in the results of lab IV a figure of "< 100 cfu/l" is shown as 50 cfu/l (detection limit with a log value of 1.7) and in the results of lab I a figure of "< 500 cfu/l" is shown as 250 (log value 2.7).



Figure 4. Comparison of the results of lab IV and lab I obtained using NEN 6265:2007 (MWY). The dotted lines show the relationship between the numbers detected using the two methods that have been compared (respectively a factor of 10 higher and a factor of 10 lower).

Two of the results shown in the graph differ markedly because they were below the detection limit in the analysis of lab IV whereas lab I stated figures of 4,000 and 65,000 cfu/l. These higher figures for these two specific samples appear to be more in line with the corresponding results obtained using the Q-PCR and ISO 11731 methods. Moreover, there is exceptionally good accordance between the results, whereby the yields were generally slightly higher for the detection method used by lab I. This may be a consequence of the extra treatment with acid applied during sample pretreatment.

Table 8 provides details of a qualitative assessment carried out in accordance with the ISO 16140 method (see section 2.4) of the results of the two NEN 6265:2007 methods using MWY medium vis-à-vis ISO 11731 as the reference method. This shows a high level of accordance although in the case of lab I the specificity of the detection method using MWY medium was relatively low. Low specificity may point to

a benefit of using MWY medium, as in cases in which the reference method is unable to detect anything, NEN6265:2007 using MWY medium is able to do so.

nie			
	NEN 6265:2007 (MWY)	NEN 6265:2007 (MWY)	
	Lab IV	Lab I	
relative accuracy (AC)	81%	83%	
relative specificity (SP)	89%	50%	
relative sensitivity (SE)	79%	90%	

Table 8 Results of the qualitative assessment of the NEN 6265:2007 (MWY) vis-à-vis ISO 11731 as the reference method.

Appendix VIII's summary of the cultivation methods shows that the cultivation method according to ISO 11731 performed best. In that case only 3 samples (5.7%) displayed excessive additional growth on the culture medium, so that no value could be stated. In the ISO method, lab I applied pretreatment with acid (in accordance with the standard) and also made counts more frequently after 3, 5, 7 and 10 days. This makes the method more laborious but produces a better result. In 37 of the 52 samples (71%) *Legionella* was detected and in 18 of those cases it was in concentrations of 10,000 cfu/l or higher. The comparison in figure 5 of the results from lab I obtained with ISO 11731 and those obtained with NEN 6265:2007 (MWY) indicates a turning point at 1,000 cfu/l. The ISO method's yield is higher when the sample contains higher numbers of legionella bacteria; the yield is lower when the numbers are lower. The significance of the measurement is lower for a figure below 1,000 cfu/litre.



Figure 5. Comparison of the results from lab I obtained using ISO 11731 and NEN 6265:2007 (MWY). The dotted lines show the relationship between the numbers detected using the two methods that have been compared (respectively a factor of 10 higher and a factor of 10 lower).

The relationship between the two methods was also examined using variance analysis (see appendix VII). The zero hypothesis was tested in this analysis that the mean of the analysis results is the same for both methods. It is 95% certain that this hypothesis cannot be rejected. Statistically, this means that there were no significant differences between the results of the two methods (noting that the plates were also assessed for both methods after 3 and 5 days).

Finally, this study also applied a cultivation method specifically for *Legionella pneumophila* analysis in accordance with figure 1 (see also section 3.1). The method is based on the draft version of NEN6253 using MWY medium, with a different culture medium pH and incubation temperature to inhibit growth

of non-pneumophila varieties. The statements below are based on a comparison of results of this cultivation method and those lab I obtained with NEN 6265:2007 using MWY medium and the associated serotyping:

- The results of both analyses were positive for 13 samples and in accordance with serotyping in every case (i.e. the presence of *Legionella pneumophila* was confirmed by serotyping).
- In 6 samples *Legionella* was detected and the concentration of *Legionella pneumophila* was below the detection limit. However, only *Legionella pneumophila* sero-group 1 or sero-groups 2 14 were confirmed by serotyping, so this might point to a discrepancy.
- In 5 samples *Legionella* was detected and the concentration of *Legionella pneumophila* was below the detection limit, and serotyping did indeed point to the presence of non-pneumophila varieties.
- The results of both methods of analysis were below the detection limit for 6 of the samples.
- No result was stated for the remaining 22 samples, as there was too much additional growth on the culture medium in 13 cases for one of the methods and in 9 cases for both of the methods.

In any case, it follows from this that there was accordance between the analytical techniques for 24 of the 30 samples. Serotyping indicated possible poor accordance between the two methods for the remaining 6 samples but it should be noted here that serotyping always involves making a selection from the colonies on the culture medium.

3.5 Effect of cooling water disinfection on the analysis results

Cooling water disinfection is a normal part of conditioning cooling water systems. The main aim of disinfection is to prevent biofilm formation in cooling water systems. Biofilm formation in heat-exchangers reduces thermal transfer and can reduce the cooling capacity of cooling water sections, as it restricts air flow. Disinfection is also intended to inhibit *Legionella sp.* growth in order to reduce exposure risks in the surrounding area.

A distinction should be made in cooling water disinfection between continuous application of a disinfectant and discontinuous application by means of shock dosing. Extensive use is still made of sodium hypochlorite (NaOCl) for disinfection but a move is underway towards more sustainable disinfection, using hydrogen peroxide, ozone or physical techniques such as ultrasound or cavitation, for example. It is also relevant to distinguish between systems with and without filtration (full stream filtration or side stream filtration). Filtration is usually used to remove undissolved substances in the circulating cooling water. This can have a positive as well as a negative effect. On the one hand, removing undissolved substances may reduce the likelihood of the cooling system becoming microbiologically contaminated. On the other hand, microbiological accumulation in the actual filter leads to increased use of sodium hypochlorite and can reduce the effectiveness of disinfection.

In this study, samples were taken from 15 different cooling water systems in which the following disinfection strategies are used:

- No disinfection (1 system).
- Use of Sonoxide (1 system).
- Continuous addition of sodium hypochlorite (NaOCl) (3 systems).
- Discontinuous addition of sodium hypochlorite (NaOCl) / shock dosing (7 systems).
- Continuous addition of hydrogen peroxide (2 systems).
- Continuous addition of ozone (1 system).

Three systems also used filtration in a side stream (1 system) or in the entire circulating water flow (2 systems).

The limited scope of this study cannot justify any statistically sound statements on the effect of a particular disinfection strategy on the presence of legionella bacteria in the cooling water. Moreover, the participating companies selected cooling tower systems with a known legionella history, so the sample of industrial cooling water systems was emphatically not non-select.

If the results of using the ISO 11731 method are taken as the starting point, it is nevertheless clear that disinfection was practically never sufficiently effective in any cooling water system. Two systems of company C and one of company E formed an exception to this but it should be pointed out that extremely high doses of chlorine were used in the systems of company C during the sampling period, owing to a defective chlorine measurement system.

The following statements on the cooling water systems covered by this study apply, if the analysis results obtained using the ISO 11731 method are used as a basis for making a rough qualitative assessment of the performance of the aforementioned disinfection strategies:

- Practically all disinfection strategies that make use of a shock dose without sand filtration perform reasonably.
- All disinfection strategies involving the continuous addition of disinfectant in combination with a sand filter (full stream and/or side stream) perform poorly (ozone with side stream filtration and sodium hypochlorite with full stream filtration).
- However, disinfection strategies involving continuous addition without sand filtration perform reasonably (Sonoxide, sodium hypochlorite, and hydrogen peroxide).

This may point to an unwanted effect of using side stream or full stream filtration in the cooling water systems covered by this study. In practice, it may be assumed that considerable biofilm formation occurs on the large internal area of a sand filter. From the operational point of view that appears to be an advantage because in that case no biofilm forms on the heat-exchanger's pipes or on the cooling section. However, in the case of continuous disinfection or if filtration is not stopped during discontinuous disinfection, the biofilm in the sand bed may consume so much disinfectant that the amount of disinfectant received by the rest of the system after the sand bed is insufficient. Depending on the position where filtration takes place with respect to the point where disinfectant is added, this may lead to inadequate disinfection in parts of the system.

4 Evaluation and conclusions

4.1 General

Over a 6-week period this study used 10 different analytical techniques to examine a total of 52 samples from 15 cooling water systems and 4 process water systems for the presence of *Legionella* or *Legionella pneumophila*.

A standard monitoring protocol was drawn up prior to sampling. The aim of the protocol was to enable sampling to provide the most representative picture possible of legionella concentration in the cooling water or process water system at the point in the system where spraying is most likely to occur and aerosols are spread into the surrounding area. For example, it is therefore clear from this that discontinuously disinfected systems should be sampled at the end of the disinfection cycle. In cooling water systems, samples should preferably be taken of the falling water in the free space above the cooling tower basin. Following this, sampling from the cooling tower basin is preferable to sampling from a sampling tap on the outlet side of the circulation pipe, as close as possible to the basin.

As far as possible, participants in this study took samples in accordance with the monitoring protocol. However, it emerged that most systems were sampled by means of a sampling tap in the circulation pipe. This is understandable, as these sampling points have been fitted especially for that purpose and because this is the least complicated way of taking samples in practice. However, if the flow through the sampling tap is insufficient before starting to take samples, there is a risk of contaminating the sample with water and biofilm from the tap-out pipe that goes to the sampling point.

The results of the individual analytical methods and the comparison of the analytical methods in chapter 3 lead to the conclusion that the screening method for *Legionella pneumophila* shown in figure 1 is a useful method that could be employed in practice. The three Q-PCR techniques are subject to relatively little inhibition from the sample matrix. No statistically significant differences were found between the three different Q-PCR techniques (appendix VII). The methods also displayed a higher degree of accordance with the cultivation method according to ISO 11731, which performed well in this study (table 7). The high performance was partly attributable to the use of sample pretreatment with acid and frequent reading of the culture media during the incubation period. This makes the method considerably more laborious and expensive.

Cooling water systems with a known 'legionella history' were selected for this study. *Legionella* was therefore expected to be detected in a relatively large number of samples. This applies to both the non-specific cultivation methods (for example ISO 11731; 81% of the samples were positive) and the Q-PCR technique for *Legionella pneumophila* (an average of 75% of the cooling water samples were positive). Primarily this means that *Legionella pneumophila* occurs in relatively large amounts in the cooling water systems studied, unlike in the case of drinking water systems, in which non-pneumophila varieties are found more often³. It also means that using Q-PCR to measure *Legionella pneumophila* in cooling water samples does not directly result in fewer positive samples. Moreover, figure 3 shows that the number of copies of DNA per litre is generally higher than the number of colony-forming units per litre. These two aspects make routine use of Q-PCR in cooling water samples less attractive in practical situations. The method is still attractive in situations in which speed of analysis plays a role, such as when determining the effectiveness of measures in emergencies.

In the traditional method of NEN 6265 using BCYE medium, it was established that the culture medium is very susceptible to additional growth. This study has confirmed with regard to this aspect that using an MWY medium leads to a considerable improvement. The good accordance of the results of detection methods applied by lab IV and lab I according to NEN 6265:2007 using MWY medium also indicates that the method is readily reproducible (figure 4). These detection results also showed a high level of accordance with the results using the ISO 11731 method (table 8). The statistical study showed no

significant differences in the results (appendix VII). However, the comparison shows that for higher numbers of legionella bacteria the yield using the ISO method is slightly higher (figure 5) but is slightly lower when the numbers are lower. Finally, the results show that pretreating samples with acid in accordance with ISO 11731 leads to a considerably lower percentage of samples with too much disruptive additional growth on the culture medium (6% for ISO 11731 against 29% for NEN 6265:2007 using MWY medium). This undoubtedly demonstrates the importance of pretreatment with acid.

The Q-PCR method for specifically detecting *Legionella pneumophila* is also in line with a cultivation method for specifically detecting the same variety of legionella. The draft version of the NEN 6253 method with the BCYE medium used for it performed poorly in this study. As with the traditional NEN 6265 method, this method suffers from excessive problems from unwanted flora. However, using an MWY medium leads to a substantial improvement. The specific cultivation method for *Legionella pneumophila* then displays good accordance with the NEN 6265:2007 method using MWY medium and the associated serotyping. It was demonstrated in section 3.1 that the method to confirm a Q-PCR result can provide useful additional information which can be used as a basis for assessing the current situation in a cooling water or process water system (table 5). Pretreating a sample with acid and reading the culture media more often would probably further improve the method's yield – as was the case with the method of working in lab I using the ISO 11731 method.

In view of its high level of accordance with the results of Q-PCR and the ISO 11731 method and given the speed and simplicity of the test, the FastpathTM method would appear to be a useful addition to the set of instruments available to the process operator responsible for the daily management of cooling water systems. The method's specificity makes it a useful addition to ATP measurements, colony number measurements or dip slides that are already used for monitoring water systems. The method's speed and simplicity make it exceptionally interesting for monitoring in emergencies. However, it is important to realise that specificity for *Legionella pneumophila* sero-group 1 means that the other serogroups are not taken into account. Moreover, the information is of such a qualitative nature that it cannot replace the measurements that are normally made. However, as it accounts for 90% of registered cases worldwide, *Legionella pneumophila* sero-group 1 is the main cause of legionella pneumonia.

4.2 Process water in the paper and cardboard industry

As mentioned in section 1.3, Koninklijke VNP is especially interested in developing a detection method for *Legionella* or *Legionella pneumophila* in the process water used by the paper and cardboard industry. It is generally assumed that conditions in the process water are optimal for *Legionella* growth but detecting the organism in this difficult matrix continues to be a problem.

Detecting legionella bacteria in the paper industry's process water proved to be a problem in this study too. For example, when using the FastpathTM method it was difficult to filter the required amount of water. When the Q-PCR technique (lab IV) was used, no results were stated on a number of occasions because the internal control was below the limit value for a reliable result. This indicates that PCR efficiency was too low, owing to an effect connected with the matrix (inhibition). Table 9 shows the results of the four most useful analytical methods for analysing the paper industry's process water. The values of the Pall Genesystems Q-PCR shown in the table have been recalculated in accordance with the draft version of NEN 6254. According to the original AFNOR standard, *Legionella pneumophila* may well have been detected (result above the limit of detection, LOD) for all the results (with the exception of those for OPIW 28 and 37) but the number of DNA copies could not be quantified (result below the limit of quantification, LOQ, as specified in the AFNOR standard). The other Q-PCR methods have higher detection limits, as a result of the small volume that was processed owing to problems with filtering the samples.

In the case of the paper industry's process water, the results obtained using the cultivation method according to ISO 11731 appear to be the most useful. However, the results indicate that the process water contains no *Legionella* or only relatively low concentrations of up to 1,000 cfu/l. Disruptive additional

growth at a level too high to permit a reliable result was only found in samples OPIW 30 and OPIW 38, which, like OPIW 46, came from the overflow of a dust chute. At the moment, although conditions in the process water appear to be optimal for *Legionella* growth, hardly any excessive *Legionella* growth was observed in the samples analysed for this study. Inhibition of *Legionella* growth and/or competition from other micro-organisms may play a role in this.

			6	
sample no.	Q-PCR	Q-PCR	Q-PCR	ISO 11731
	Lab IV	Lab II	Lab III	
	number of copies/l	number of	number of	cfu/l
		copies/1	copies/1	
OPIW 29	no result	3,430	no result	< 50
OPIW 39	406	537	< 1,200	510
OPIW 47	< 6,931	732	< 550	960
OPIW 30	no result	35,100	< 700	no result
OPIW 38	8,133	6,520	< 1,200	no result
OPIW 46	< 2,688	18,600	< 1,000	< 500
OPIW 28	< 2,430	< 680	< 1,400	< 50
OPIW 37	< 2,931	< 680	< 1,200	< 50
OPIW 48	no result	1,510	< 29,000	< 50

Table 9 Analysis results for samples from the paper and cardboard industry

4.3 Pricing

Q-PCR technology is developing rapidly. More and more laboratories are capable of using the method and more and more samples are being presented for analysis. This leads to competition and a reduction in analysis costs. The price charged by a commercial laboratory for Q-PCR analysis of large numbers of samples (n > 20) is expected to work out in the range of \notin 40 to 60 per sample.

In comparison with analysing drinking water samples, the current practice of using the cultivation method on cooling water samples requires more time (+ 50%) for isolation and identification. When cultivating *Legionella pneumophila*, if treating the sample with acid increases the yield and more frequent counting leads to a better result, it will be necessary to take into account that the detection method will be even more labour-intensive and that the current cost of the cultivation method will increase. The price is expected to increase by 50 to 100% with respect to the current price for drinking water samples. However, the latter price can vary considerably in practice and is not always a true reflection of the quality required for a good analysis.

4.4 Conclusions and recommendations

Based on the results of the study, figure 6 is a more detailed version of the scheme shown in figure 1. The Fastpath[™] method is a useful addition to the set of instruments available to the process operator for the daily management of cooling water systems. This method is faster in providing information on the possible presence of *Legionella pneumophila* sero-group 1 in cooling water and in that respect could initiate further analysis in accordance with the scheme in figure 6.

This study has shown that Q-PCR technology provides a fast, reliable and affordable technique for detecting *Legionella pneumophila* in cooling water and process water. (A restriction on using this technique is that the government has not made the step from detecting *Legionella* as a whole to detecting *Legionella pneumophila*.) The speed of the Q-PCR method is a major advantage for the industry. However, if it emerges that *Legionella pneumophila* is fairly common in cooling water systems, it will also be necessary to perform analyses regularly in accordance with the scheme in figure 6, using the cultivation method to confirm the viability of the legionella bacteria. This means that the time gain would be lost for some samples and, moreover, that performing screening in accordance with the scheme in figure 6

would be relatively expensive in comparison with only conducting an analysis using the cultivation method.

However, this study has shown that successfully using a cultivation method on cooling and process water samples will also lead to extra costs. Pretreating a sample with acid and counting more frequently for the specific detection of *Legionella pneumophila* make detection much more labour-intensive and therefore expensive. Of all the cultivation methods used, the cultivation method according to ISO 11731 performed well. However, counting was also more frequent in this case, to prevent additional growth on the culture medium from making it difficult to count colonies.



Figure 6. Modified scheme for screening for Legionella pneumophila in cooling water and process water on the basis of the results of this study.

The advice for selecting a cultivation method from the scheme shown in figure 6 is to

- examine whether the ISO 11731 method could be made suitable for detecting *Legionella pneumophila* and
- to further optimise the specific cultivation method for *Legionella pneumophila* using an MWY culture medium (based on NEN 6265:2007 and the draft version of NEN 6253) by applying pretreatment with acid and more frequent counting.

Furthermore, it would be advisable to check the performance of the two selective cultivation methods separately and in combination with Q-PCR (in terms of their microbiological yield and cost) by using them on a sufficiently large non-select sample of industrial cooling water systems. The results of the study will enable a final decision to be taken on the usefulness of advance screening using Q-PCR in accordance with the scheme in figure 6.

Conclusions:

- This study has demonstrated the usefulness and added value for analysing cooling water and process water samples in accordance with the scheme in figure 6. Besides providing managers of cooling water and process water systems with information faster, this new screening methodology also provides them with more specific information on the presence of *Legionella pneumophila* in the system. The choice of the specific Q-PCR for *Legionella pneumophila* in the scheme does not affect the result. The choice of the specific cultivation method should be based on the recommendations of this study and possibly the results of a further study.
- Variance analysis has shown for the cooling water and process water used in this study that there are no significant statistical differences between the results of three Q-PCR techniques performed by

three different laboratories in accordance with three different standards for detecting *Legionella pneumophila*.

- In most cases, the results of the Q-PCR techniques for the cooling water samples studied meet the requirement for a detection limit of at least 1,000 units/litre, based on table 12 of Health & Safety Information Sheet AI-32. If using Q-PCR technology systematically leads to more units per litre being detected than the number detected using the cultivation method, table 12 of the Health & Safety Information Sheet will have to be expanded for the interpretation of Q-PCR results. With regard to the specific situation at a company, when switching to Q-PCR, the method should preferably be used in parallel for a while with the cultivation method normally used up to that point, so that the results of the old and new method can be correlated. This can then be used as a basis for adjusting the action levels in the legionella control plan.
- The practical usefulness of the scheme in figure 6 may be limited by the observation that using Q-PCR for the specific detection of *Legionella pneumophila* in the cooling water samples analysed, does not lead to less positive samples than detection based on a cultivation method for *Legionella* in total. In comparison with drinking water systems, *Legionella pneumophila* is apparently fairly generally present in cooling water systems.
- This study has shown that applying NEN 6265 with BCYE medium for cooling water and process water samples leads to unsatisfactory results, owing to disruptive additional growth on the culture medium. NEN 6265 was revised for this in 2007 by also describing the MWY medium for samples with excessive additional growth.
- This study has shown that pretreating cooling water and process water samples with acid in accordance with ISO 11731:1998(E) and counting more frequently on day 3 and day 5 after incubation, leads to fewer outcomes for which no result can be stated on account of additional growth on the culture medium.
- Variance analysis has shown for the cooling water and process water used in this study that there are no significant statistical differences between the results of the cultivation method according to ISO 11731 and the cultivation method according to NEN 6265:2007 using an MWY culture medium, provided that the agar plates are also assessed on day 3 and day 5.
- Taking into account the method's limitations, the FastpathTM method is a useful addition to the set of instruments available to the process operator responsible for the daily management of cooling water systems. The method's main benefits are speed, simplicity and the possibility of on-site implementation. However, it is important to realise that specificity in respect of *Legionella pneumophila* sero-group 1 means that the other sero-groups are not taken into account. (Nevertheless *Legionella pneumophila* sero-group 1 is the main cause of legionella pneumonia; up to 90% of registered cases worldwide). Moreover, the information is of such a qualitative nature that it cannot replace the measurements that are normally made.
- This study has not provided any indications of the presence of high concentrations of legionella bacteria in process water samples from the paper and cardboard industry. Nevertheless, the results indicate that *Legionella* or *Legionella pneumophila* can be detected, at least in a number of samples.
- It was notable for practically every cooling water system examined in this study that disinfection hardly ever consistently resulted in a legionella bacteria concentration of less than 1,000 cfu/l.
- The laboratories participating in this study have demonstrated that it is possible to achieve reliable results using a cultivation method for *Legionella* and/or *Legionella pneumophila*. A contributory factor to this was that these laboratories were experienced and did everything possible to achieve good analytical results, without attempting to cut costs or avoid difficulties. The interlaboratory comparisons organised by KWR also confirmed that a certain degree of reliability can be achieved using a cultivation method.

Recommendations:

• It is recommended on the basis of the experiences from this study that the specific cultivation method for *Legionella pneumophila*, with an MWY cultivation medium based on NEN 6265:2007 and the draft version of NEN 6253, should be further optimised by means of pretreatment with acid and more frequent counting.

- As good results in this study were obtained using the cultivation method according to ISO 11731, it would be advisable to examine the extent to which the method could also be made suitable for the specific cultivation of *Legionella pneumophila*.
- A further recommendation assuming a sufficiently large non-select sample of industrial cooling water systems is that the two specific cultivation methods for detecting *Legionella pneumophila* should be checked both separately and in combination with Q-PCR assessment to determine their performance in terms of yield and cost.
- It would be advisable to set up interlaboratory comparisons for the Q-PCR techniques, as is done for the cultivation method.
- The cultivation method in accordance with ISO 11731 is recommended for detecting *Legionella* in process water from the paper and cardboard industry, subject to pretreating the sample with acid and reading the culture medium counts more frequently after incubation.
- A further examination of the role that 'full stream' or 'side stream' filtration plays in the effectiveness of disinfection in cooling water systems is recommended. Relevant variables in this are the type of disinfectant and the dose and method of dosing (continuous/discontinuous).

5 References

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I Appendix: Sampling protocol (partly in English)

Inleiding

Aanleiding

Dit protocol is opgesteld als onderdeel van het project OPIW 15 "Evaluatie van een snelle, betrouwbare en reproduceerbare screeningsmethodiek voor *Legionella*'s in koelwater en proceswater".

Op grond van de ervaringen met de toepassing van het protocol in dit onderzoek kan worden overwogen de methodiek verder te standaardiseren.

De input voor dit protocol is geleverd door een team bestaande uit de volgende personen: Antoine van Hoorn (Corus), Jack Smeets (KEMA Zuid), Jo Savelkoul, Ralph Lindeboom en Frank Oesterholt (allen Kiwa Industrie & Water).

Doelstelling

Dit protocol is bedoeld voor de deelnemers aan OPIW 15 en heeft als doel een monster/staal te verkrijgen dat een zo representatief mogelijk beeld geeft van de concentratie legionellabacteriën in het (circulerende) water van te onderzoeken object. Daarnaast moet het protocol leiden tot een gestandaardiseerde monstername/staalname binnen het project.

Tijdens het startoverleg is afgesproken dat elke projectdeelnemer 3 locaties selecteert (bijvoorbeeld 3 koelwatersystemen respectievelijk 3 proceswatersystemen, zoals in gebruik bij de productie van papier dan wel de productie van ijzer en staal) die vervolgens in de loop van het project 3 keer worden bemonsterd. Totaal 9 monsters per deelnemer.

Protocol

Conditions

Conditions for sampling of cooling water systems

For this research only samples should be taken from cooling water systems that have been continuously working during a period of at least 2 weeks.

On purpose of the research the water distribution in the cooling water tower has to be controlled once before starting the first sampling. A bad water distribution reveals positions with splash streams (e.g. caused by a broken distribution scale) and/or positions with just very small water contents (obstructed nozzles/sprinklers). (This inspection should be part of regular controls as these problems (i) could have negative influences on the capacity of the cooling water systems, (ii) could cause a high chance of pollution of the package and (iii) also the chance of growth of legionella bacteria. Inspection preliminary to this research is of great importance because you can hardly execute/carry out good and representative samples in a cooling water tower with a bad water distribution. In case of bad water distribution the sampler and operator have to decide together to sample the cooling water tower by other methods (see preferences listed below).

Voorwaarden voor bemonstering van proceswatersystemen

Voor dit onderzoek dienen alleen proceswatersystemen te worden bemonsterd die gedurende een periode van ten minste 2 weken continu in bedrijf zijn geweest.

Bij voorkeur wordt een proceswatersysteem of een deel van een proceswatersysteem geselecteerd waaraan geen biocide wordt gedoseerd.

Preparation

The sampling has to be executed by an instructed sampler. Sampler is perfectly aware of the contents of this protocol.

The sampling has to be executed with sterilised bottles provided by KWR (content 1 litre). In a standard way these bottles contain a solution of sodiumthiosulphate for neutralisation of eventually present chlorine (or another oxidation biocide) and a NTA-solution for **complex formation** of eventual presence of heavy metals.

Execution

For this research sampling volumes are required bigger than 1 litre, so more bottles have to be filled. As expected three bottles have to be filled (total 3 litres) directly after each other.

As the bottles contain a small content of sodiumthiosulphate and NTA solution, these bottles preferably should not overflow during the sampling. We advise to fill these bottles for only 90%.

After the last sampling the water temperature has to be measured and registered.

KWR provides each run of sampling bottles with an <u>information sheet</u> for the registration of temperature and other relevant conditions under which circumstances the sampling has been done. The sampler should fill these sheets in (eventually in a mutual agreement with the operator).

Execution sampling cooling water systems

Cooling water systems discontinuously chorinated (disinfected) have to be sampled at the end of the disinfection cycle (i.e. just for the new shock dose).

For cooling water systems continuously disinfected the moment of sampling does not matter.

Sampling a cooling water system next sequence of preferences has to be used to determine the position of sampling.

1. Sampling of <u>the falling water in the free space above the cooling water</u> <u>basin</u>. The position of sampling is minimal 1 metre from the edge of the cooling tower. A special tool adjusted at the circumstances with the help of which a sampling bottle can be positioned at the desired distance in the water curtain, has to be used.

If this way of sampling is not executable for practical reasons (e.g. too small louvres for the sampling bottle) and/or bad water distribution in the cooling tower (see paragraph 1.3) this way of sampling has to be cancelled. Than the next preference will be:

2. Sampling of <u>water in the cooling water basin</u>: The position of sampling has to be 1 metre from the edge, maximum 10 till 20 centimetres under the water surface (avoid contact with the bottom) and on a position in the cooling water basin as close as possible to the suction pipe of the recirculation pump (water has to be in motion).

As for practical reasons this way of sampling is also not possible to execute due to too far distance between the sampling position and the cooling water basin or due to position points having a bad through-flow, this way of sampling has to be cancelled. In this case the next preference will be: Taking samples from a <u>sampling tap in a main water pipe</u> after the recirculation pump (press side of the pump). *This way of sampling has to be carried out as close as possible to the cooling water tower. For this method a supplementary requirement is that the pipe to the sampling point (measured from the main water tap) has to be shorter than 3 metres. Before starting sampling the pipe has to be continuously streamed with full jets during at least 3 minutes in order to avoid wall effects.* In fact such a sampling pipe should be regularly flushed, e.g. as part of the procedure of flushing 'emergency showers'.

Uitvoering bemonstering proceswatersystemen

Proceswatersystemen die discontinue worden gedesinfecteerd dienen te worden bemonsterd aan het einde van de desinfectiecyclus (dat wil zeggen vlak voor de nieuwe dosering van desinfectiemiddel).

Voor proceswatersystemen die continue worden gedesinfecteerd, maakt het moment van monsterneming niet uit.

Papier- en kartonindustrie

In de Handleiding *Legionella* voor de papier- en kartonindustrie die door Kiwa Industrie & Water in opdracht van de Koninklijke VNP is opgesteld [1], is een prioriteitstelling opgenomen voor proceswaterlocaties in de papierfabriek. Deze prioriteitstelling is uitgevoerd op basis van het potentiële risico op blootstelling aan *Legionella* via aërosolen, op basis van de mate van aërosolvorming, de frequentie en duur van aanwezigheid van personeel en de kans op vermeerdering van *Legionella*.

Op grond van deze referentie dient voor de bemonstering van proceswater uit een papier- of kartonproductiebedrijf de volgende **voorkeursvolgorde** te worden gehanteerd voor de keuze van de positie van monsterneming (locaties die los staan van de papierproductieprocessen zoals hogedrukreinigers en schoonmaakhaspels zijn buiten beschouwing gelaten):

- 1. de ontwatering van de zeefpartij;
- open sproeisysteem voorraadkuipen/indikkers proceswater gevoed; open pulper (stofvoorbereiding); gapvormer (fijnpapier/tissue stofomloop) sorteertrommel rejectreiniging (stofvoorbereiding); sproeisysteem viltreiniging proceswater gevoed (perspartij) sproeisysteem vilt- en zeefreiniging proceswater gevoed (zeef/nat-partij)

Bij de selectie van het monsterpunt dienen de volgende overwegingen te worden meegenomen:

- de prioriteit zoals hierboven aangegeven (1,2)
- een vrij vallende waterstroom heeft de voorkeur boven bemonstering via een tappunt (vergelijkbare situatie als bij de koeltoren);
- bij bemonstering van een tappunt dient dat tappunt <u>niet</u> door middel van een (rubber of kunststof) slang te zijn gekoppeld aan de installatie;

- bij bemonstering van een tappunt dient dat tappunt ten minste 5 liter water per minuut te leveren (uitgangspunt inwendige leidingdiameter max. 20 mm);
- bij bemonstering van een tappunt dient de uittapleiding naar het tappunt bij voorkeur zo kort mogelijk te zijn (< 1 meter). Om wandeffecten van de leiding en het tappunt te voorkomen dient de leiding minimaal 1 minuut bij volle straal te worden doorstroomd alvorens met de monsterneming wordt gestart.

<u>IJzer- en staalindustrie</u>

De ervaring heeft geleerd dat binnen de ijzer- en staalindustrie regelmatig relevante concentraties legionellabacteriën in proceswatersystemen aanwezig zijn. Ook hier wordt aan de hand van factoren als de mate van aërosolvorming, de frequentie en duur van aanwezigheid van personeel en de kans op vermeerdering van *Legionella* preventief een standaard monsternameprogramma uitgevoerd op een aantal verschillende locaties.

Enkele overwegingen voor de selectie van de bemonsteringslocatie die hierboven zijn opgenomen voor de papierindustrie gelden ook hier:

- een vrij vallende waterstroom heeft de voorkeur boven bemonstering via een tappunt (vergelijkbare situatie als bij de koeltoren);
- bij bemonstering van een tappunt dient dat tappunt <u>niet</u> door middel van een (rubber of kunststof) slang te zijn gekoppeld aan de installatie;
- bij bemonstering van een tappunt dient dat tappunt ten minste 5 liter water per minuut te leveren (uitgangspunt inwendige leidingdiameter max. 20 mm);
- bij bemonstering van een tappunt dient de uittapleiding naar het tappunt bij voorkeur zo kort mogelijk te zijn (< 1 meter). Om wandeffecten van de leiding en het tappunt te voorkomen dient de leiding minimaal 1 minuut bij volle straal te worden doorstroomd alvorens met de monsterneming wordt gestart.

Persoonlijke bescherming

Personal protection during sampling of cooling tower water.

During the sampling of cooling water systems following preferences 1 and 2 at least the regular, required personal protection outfits have to be used in combination with gloves an a mask of type FFP3. These masks have to meet the requirements following EN 149:2001.

Until recently filtration masks where available only protecting against solids or liquids. At this moment all filtration masks are suitable for the protection against solid particles as well as liquid particles. The out of date qualifications written at the type of masks, were: S means Solids L means Liquids

Persoonlijke bescherming bij bemonstering proceswatersystemen

Bij de bemonstering van proceswatersystemen dienen de reguliere, voorgeschreven persoonlijke beschermingsmiddelen te worden gedragen, maar in ieder geval handschoenen.

Bij bemonstering van een vrij vallende waterstroom waarvoor een afgeschermde positie in het proces moet worden ingenomen, moet – in verband met de kans op blootstelling aan hoge concentraties aërosolen worden overwogen om een gelaatsmasker van klasse FFP3 te dragen (zie opmerking paragraaf 2.4.1.)

Transport of the samples

Immediately after sampling each sample has to be put on ice or icepacks and placed in a cooling box. On the same day of sampling, the samples have to be delivered (if necessary by courier) before 12.00 AM at Kiwa Water Research to the attention of Harm Veenendaal (chief of the microbiology laboratory).

Referenties

[1] Handleiding *Legionella* voor de papier- en kartonindustrie in Nederland. KWR 07.020. Kiwa Industrie & Water, 2007.

II Appendix: Information sheet

Information sheet monstername OPIW 15

This sheet should be filled in after sampling of a cooling water or process water system. It provides information about the sampling itself and the relevant system conditions at the time of sampling. (See other side for footnotes).

General Information	_
Date/time	//2008:hrs
Name sampler/telephone nr	
Company name	
Description cooling water or process water system	
Description used sampling method ¹ (see protocol sampling) Description used codes on sampling bottle for this sample	
Number of filled bottles	
Water temperature after sampling	°C
Visual description samples ²	
Information in detail sampling system	
Description disinfection procedure cooling- or process water system ³	
Expired time after last chlorination	
Actual chloride concentration or last measured chloride concentration + date and time	
Position blow down (open/closed) during sampling	
Last measured ATP value for the system + date/time ⁴	
Used ATP action level ⁴	
Last measured colony count + date/time ⁴	
Used action level for colony count ⁴	
Last measured legionella concentration + date/time ⁴	
Hydraulic residence time in the system = volume system/blow down	
Type of make up water	
Are sand filters being used in the cooling water systems? Side stream of full stream?	

Notes:

- 1) Describe the way the sampling was executed. Therefore we refer to the protocol's sequence of preferences for sampling.
- 2) Describe remarkable things of the taken water samples, e.g. troubled or clear water, many suspended particles, sediment in the bottle, odour and colour, etc. etc.
- 3) What type(s) of disinfection method(s) has (have) been used? Do they disinfect continuously or discontinuously. What doses have been used and how frequently?
- 4) Gather these data only when these are applicable, e.g. when ATP measurements or dip slides have been used as protection of the system. Describe in this case when the last measurement took place (date). In both cases please mention also which action levels have been used.

III Appendix: Overview research results

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	th (Lp1) Ico	volume analysed (ml)	250	250	250	250	250	200	250	230	250	250	250	250	250	250	250	250	022	250	250	250	250	250	250	250	250	250	250	250	250	250	250	250	150	150	250	250	02C	250	250	75	250	250	40	40	25 E0	00	t 0
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	PCR (Lp) concept o-NEN 6254	kopieën/l	223.798	196.796	87.914	79.863	133.443	00.130 7.000	020.0 1 164	1.104	504.1 772 778	335.753	127.073	8.800	1.691	2.063		1.633.306	03/.144	<591 2554	100/	22.717	664	2.736	<549	<612	711	32.696	75.855	< 699 5 060	19.763	1.786	<508	29.762		161.853	<848	<844	12 857	174.733	117.841	ľ	406	<6931		8.133	<2688	<2931	
KWR	MWY pneu pH 7,3 & temp 40C	cfu/l	1.600	(1)	6.500	<100	27.920	00.5.00	0012	.040		600	500	1.040	960	<100	(1)	<100	(I)	<100	2100 2100	16.800	<100	700	(1)	<100	<100	120.000	8.900		<100	2.233	<100	25.333	<100	(1)	<100	<100	(1)	00	(1)	(1)	(1)	<100	(1)	(1)	(1)	2100 2100	<100
	BCYE pnue pH 7,3 & temp 40C	cfu/l	5.400	(3)	3.400	500	(3)	000.12	(1)	(9)	(1) EOO	(3)	<100	(1)	(3)	(1)	(1)	(3)	(I)	<100	2100 2100	(1)	200	300	(1)	<100	<100	(1)	(3)		(3)	(1)	(3)	(1)	(1)	(1)	(1)	<100		(1)	(1)	(1)	(1)	(1)	(1)	(1)	(1)	2100 2100	<100
	MWY NEN 6265:2007	cfu/l	7.800	38.500	15.600	300	146.200	40.000			3 500	1.200	300	1.900	480	<100	Ð	100	(1)	<100	< 100	28.100	1.700	1.800	100	200	100	(1)	50.000	(1)	(1)	(1)	<100	6.400	<100	6.100	<100	<100	(1)	<100	<100	(1)	(1)	<100	(1)	(1)	(1)	(1)	<100
	BCYE + AB NEN 6265	cfu/l	1.800	10.200	(1)	300	(1)	008.76	(L)		(1)	600	100	(1)	(1)	(1)	<100	4.080	(1)	<100	<100	(1)	500	1.375	100	633	400	(1)	(1)	(1)	(1)	(1)	(1)	(1)	(1)	(1)	<100	<100	(1)	(1)	(1)	(1)	(1)	(1)	(1)	(1)	(1)	(1)	(1)
	sample code		OPIW 7	OPIW 25	OPIW 44	OPIW 8	OPIW 26	OPIW 43		OPIN/ 45	OPIN 43	OPIW 22	OPIW 31	OPIW 6	OPIW 23	OPIW 33	OPIW 5	OPIW 24	0PIW 32	OPIW 15		OPIW 13	OPIW 41	OPIW 50	OPIW 14	OPIW 42	OPIW 49	OPIW 10	OPIW 19	OPIW 1	OPIW 20	OPIW 12	OPIW 21	OPIW 18	OPIW 35	OPIW 53	OPIW 16	OPIW 36	OPIW 54	OPIW 34	OPIW 52	OPIW 29	OPIW 39	OPIW 47	OPIW 30	OPIW 38	OPIW 46	OPIM 20	OPIW 48
	company		A	A	۷	A	A	4	× <	4	τ α		В	В	В	в	в	8	<u>م</u>	ບບ	50		c	S	U	U	υ	۵	٥			٥	Δ	ш	ш	ш	ш	ш	ц	ш	ıш	L	L	ц	ш	ш		L U	<u> </u>
	date 2008		26-aug	16-sep	29-sep	26-aug	16-sep	Z9-Sep	26-aug	10-Sep	Je-ep	16-sep	23-sep	26-aug	16-sep	23-sep	26-aug	16-sep	z3-sep	Z-sep	E-Sep	2-sen	29-sep	6-okt	2-sep	29-sep	6-okt	2-sep	16-sep	26-aug	16-sep	2-sep	16-sep	2-sep	23-sep	6-okt	2-sep	23-sep	0-0KI	23-sen	6-okt	23-sep	29-sep	6-okt	23-sep	29-sep	6-okt	20-sep	6-okt

- No Legionella found, result not reliable due to too much side growth on the agar medium Estimated value, real number possibly higher due to much side growth on the agar medium By mistake inoculated on BCYE-AB, pH 7.3. No Legionella found, result not reliable due to too much side growth on the agar medium

 - Analysis inhibited by the samples matrix. Real number of *Legionella pneumophila* is probably higher No *Legionella pneumophila* found. Analysis inhibited by the samples matrix; not reliable result. Difficult to visibly detect line; line is very thin

- Legionella preumophila found, but result can not be quantified. Colonies (partity) confirmed with PCR Though negativ at confirmation on plate, presence of *Legionella* is suspected The output of the internal control sample is too low to give a reliable quantitative result *Legionella pneumophia* found, but the output of the internal control sample is too low to give a reliable quantitative result *Legionella pneumophia* concentration is indicative, below the lower limit for a reliable quantitative result (1) No Le Stima (2) Estima (3) By may (5) No Le (5) No Le (6) Difficu (6) Difficu (6) Doug (6) Thoug (10) The or (11) Legior (12) Legior (12) Serotypes: a = Serotypes: a = Coloni
- Legionella pneumophila serogroep 1. Legionella pneumophila serogroep 2-14. Legionella bacteria non pneumophila.

IV Appendix: Q-PCR *Lp* followed by a specific cultivation method for *Lp*

			full stream sand filtration	full stream sand filtration	full stream sand filtration	full stream sand filtration	full stream sand filtration	full stream sand filtration							autom. chloringssysteem 2 * 2 hour/day 2 ppm	autom. chloringssysteem 2 * 2 hour/day 2 ppm	autom. chloringssysteem 2 * 2 hour/day 2 ppm	shock dose op basis van hoge TBC/ATP	shock dose op basis van hoge TBC/ATP	shock dose op basis van hoge TBC/ATP	real concentration: 0 ppm	real concentration: 32 ppm	real concentration: 13 ppm	real concentration: 0,13 ppm	real concentration: 4,5 ppm	real concentration: 4,2 ppm	real concentration: 0,2 ppm	real concentration: 15 ppm	real concentration: 1,5 ppm	continuous chloring 1,25 ppm free chlorine	continuous chloring 1,25 ppm free chlorine	shock dose once every 2 weeks	continuous chloring 1,25 ppm tree chlorine	continuous critoring 1,20 pprintee critorine	stituce doss last i ween ago (20/0)							side stream filtration	side stream filtration	side stream filtration									
desinfectie			0,1 - 0,2 Cl cont + 1 ppm ClO2 1 u/day	0,1 - 0,2 Cl cont + 1 ppm ClO2 1 u/day	0,1 - 0,2 Cl cont + 1 ppm ClO2 1 u/day	0,4 - 0,5 ppm sodiumhypochlorite cont	0,4 - 0,5 ppm sodiumhypochlorite cont	0,4 - 0,5 ppm sodiumhypochlorite cont	Sonoxide B35	Sonoxide B35	Sonoxide B35	no disinfection allowed	no disinfection allowed	no disinfection allowed	0,10 ppm actual Cl2 conc.	0,10 ppm actual Cl2 conc.	0,15 ppm actual Cl2 conc.	last shock dose 4 mnd ago; 2 ppm	last shock dose 5 mnd ago	last shock dose 5 mnd ago	0,5 ppm Cl 15 min/24 hour	0,5 ppm Cl 15 min/24 hour	0,5 ppm Cl 15 min/24 hour	0,5 ppm Cl2 15 min per 8 hour	0,5 ppm Cl2 15 min per 8 hour	0,5 ppm Cl2 15 min per 8 hour	0,5 ppm Cl 15 min/24 hour	0,5 ppm Cl 15 min/24 hour	0,5 ppm Cl 15 min/24 hour	disinfection with NaOCI	disinfection with NaOCI	disinfection with NaOCI	disinfection with NaOCI				continu H2O2 continu H2O2		continu H2O2	continu H2O2	continu H2O2	continu ozone dose	continu ozone dose	continu ozone dose	none	none	none	none	none	none	hypochlorite during cleaning; 5 days ago	hypochlorite during cleaning; 8 days ago	hypochlorite during cleaning; 6 days ago
	neu	sd	1746		1458		9008	4192		876		652	652	354	727	699								8362		1095					10071			1365	0001	1776	41 20																
	d YWM σt s c τ ⊔ c	cfu/l	1600 (2)	(1)	6500	<100	27920 (8)	50300	<100	1040 (8)	<100	600 (8)	600	500	1040	960 (8)	<100	(1)	<100	(1)	<100	<100	<100	16800	<100	700	(1)	<100	<100	120000 (2)	8900 (8)	(1)	(1)	~100 2232 (2)	<100	25332 (0)	(2) 0012 20000	(1)	<100	<100	<100	(1)	(1)	(1)	(1)	(1)	<100	(1)	(1)	(1)	<100	<100	<100
lab IV	(Lp)	sd scot	20410,91	21749,99	2321,93	4183,07	16380,87	2834,72	1693,57	256,61	1333,99	75572,04	38319,38	16420,04	478,06	195,14	1497,68		8173,55	10501,85				2215,60	379,95	627,12			696,19	770,64	4193,87	1	448,97 4226 05	4.020,30 8.28	0,20	0661 27	3001,27	3392 34	-014000			3147,68	20725,33	11670,29		182,27			479,00				
	PCR	concept o-	,24E+05	,0E+05	,79E+04	,99E+04	,3E+05	;51E+04	,63E+03	,2E+03	,47E+03	(,22E+05	(,4E+05	,27E+05	(,80E+03	,7E+03	1,06E +03	(11	,6E+06	;,37E+05	:591	:551	:655	,27E+04	;64E+02	,74E+03	549	:612	,11E+02 (12	(,27E+04	,6E+04	: 699	,06E +03	, UET 04 70E ±03	508		, 30E TU4	R+05	848	844	:905	.29E+04	,75E+05	,2E+05	(11	.06E+02 (12	:6931	(11	,13E+03 (12	:2688	2430	:2931	0 L)
		Alliple code	DPIW 7	DPIW 25	DPIW 44 8	DPIW 8 7	DPIW 26	DPIW 43 8	DPIW 9	DPIW 27	DPIW 45	DPIW 4 8	DPIW 22	DPIW 31 1	DPIW 6 8	DPIW 23	DPIW 33 2	PIW 5 -	DPIW 24	DPIW 32 6	DPIW 15 <	DIW 40	DPIW 51 <	DPIW 13 2	DPIW 41 6	DPIW 50 2	DPIW 14 <	0PIW 42 <	DPIW 49 7	DPIW 10	DPIW 19 7	DPIW 1						DPIW 53	DIW 16	DPIW 36	DPIW 54 <	DPIW 17	DPIW 34	DPIW 52 1	DPIW 29 -	DPIW 39 4	DIW 47 <	DPIW 30	0PIW 38	DIW 46 <	DPIW 28 <	DPIW 37	DIW 48 -
		Junpany &						5	5	7		<u> </u>	<u> </u>	0	<u> </u>	<u> </u>	0	3	3	<u>د</u>	0	0	0	0	0	0		0	0	0										, 0													<u>_</u>
			26-aug	16-sep /	29-sep	26-aug /	16-sep /	29-sep /	26-aug /	16-sep /	29-sep /	26-aug E	16-sep E	23-sep E	26-aug E	16-sep E	23-sep E	26-aug E	16-sep E	23-sep E	2-sep (29-sep (6-okt (2-sep (29-sep (6-okt (2-sep	29-sep (6-okt (2-sep [16-sep [26-aug	Z-sep	10-sep	16-cen	dee o	23-sep	6-0kt	2-sen	23-sep	6-okt E	2-sep E	23-sep E	6-okt E	23-sep	29-sep	6-okt F	23-sep F	29-sep F	6-okt F	23-sep	29-sep	6-okt

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- No Legionella found, result not reliable due to too much side growth on the agar medium
- Estmated value, real number possibly higher due to much side growth on the agar medium Analysis inhibited by the samples matrix. Real number of *Legionella pneumophila* is probably higher
 - - No Legionella pneumophila found. Analysis inhibited by the samples matrix; not reliable result.
- Colonies (partly) confirmed with PCR $\begin{array}{c} (12) \\ ($
- The output of the internal control sample is too low to give a reliable quantitative result
- Legionella pneumophila found, but the output of the internal control sample is too low to give a reliable quantitative result Legionella pneumophila concentration is indicative; below the lower limit for a reliable quantitative result

V Appendix: Comparison Fastpath and Q-PCR/ISO 11731

					A	1 - 41		
sam	ple code	concep	ot o-NEI) N 6254	rastpatr Nalo	n (Lp1) co	ISO	NEN 6265
						volume analysed		
Č	- IV	kopieën/l	ы С	d 0110.01	result	(ml)	Serotype	Serotype
	W / W 25	2,24E+U5 2 0F+05	20	0410,91 1749.99	positive (6) positive (6)	250	a+b+c a+b+c	a+b+c a+b+c
Ыd	V 44	8,79E+04		321,93	negative	250	a+b+c	a+b+c
Ido	W 8	7,99E+04	4	183,07	positive (6)	250	q	a+b
lЧО	W 26	1,3E+05	7	6380,87	positive (6)	250	p	q
G	W 43	8,51E+04	Ñ	834,72	negative	250	q	q.
e o	W 9	5,63E+03	÷ 0	693,57	positive (6)	250	D+C	a+b
Ъб	W 27	1,2E+03	N T	56,61	negative	250	D+C	D+C
5	W 45	1,4/E+03	- 1	333,99	negative	250	D+C	D+C
ЪĞ	W 4	8,22E+05	~ 7	55/2,04	negative	250	Ω.	۵
ЪG	77. M	3,4E+05	τ̈́ τ	8319,38	negative	250	q	
	W 31	1,2/E+U5		042U,U4	negative	750	a+b	0
	0 10	0,00E+U3	7 7	10,00 05.11	positivo	090	a+b	
	VN 22	2 DEFLO3		407.68	positiva	250	4TC	
	W 5		(11)	00,101	positive (6)	250	a+b +b	
OPI	W 24	1.6E+06	8	173.55	negative	250	d+b	e
0 PI	W 32	6,37E+05	I	0501,85	positive (6)	250	a+b	
IdO	W 15	<591			negative	250	v	
IdO	W 40	<551			negative	250		
IdO	W 51	<655			negative	250		
lЧО	W 13	2,27E+04	5	215,60	negative	250	a+b	a+b
OPI	W 41	6,64E+02	ŝ	79,95	positive (6)	250	q.	a+b
	W 50	2,74E+03	6	27,12	positive (6)	250	a+b	a+b
36	W 14	<549			negative	250	Q	0
36	IV 42	< 11 - 00 - 21.02		10	negative	250	0	D+C
56	W 49	7,11E+UZ	6 (ZL)	90,19	negative	720	D+C	a+b+c
56	01 10	3,2/E+04	< <	10,04	positive	022	a+b	9450
56	IV 18	/ 0E+04	4	193,07	positivo	020	a+b+c	a+D
5 6	111	< 089 5 06F±03	4	48 97	positive	250	n 1	4
5 6	02 MIC	2,005-03	4	326.95	positive	250	a+b	a+b
ō	PIW 12	1,79E+03	8	.28	positive	250	a+b	1
Ö	olW 21	<508			positive	250		a
ō	JW 18	2,98E+04	ō	661,27	positive (6)	250	q	b+c
Ō	PIW 35	-	(11)	1000	positive	150		-
5 6	1W 33	<848	Ó	332,34	positive negative	250	a+D+C	a+0 h+c
ö	PIW 36	<844			negative	250	0	2
P	IW 54	<905			negative	250		q
g	IW 17	1,29E+04	33	147,68	positive	250		
ğ	PIW 34	1,75E+05	2	0725,33	positive	250	a+b	p
Ğ	PIW 52	1,2E+05	1	1670,29	positive	250	a+b	a+b
Ğ	JW 29	-	(11)		positive	75		
Ъ	1W 39	4,06E+02	(12) 1	82,27	negative	250	U	
Ö	PIW 47	<6931	4		positive	250	b+c	
Ъ	N 30		(11)	000	positive	40		
56	IW 38	8,13E+03 <7688	(12) 4	/9,00	positive	40 25		
0	W 28	<2430			negative	50		
9	IW 37	<2931	Η		negative	40		
ldО	W 48		(10)		negative	10		

000	202	2020
Fathpath 52 samples	Jumes talse	

47 samples PCR = reference mei astpath = alternatiev	thod ve method		
PCR + PCR -	+ Fastp 28 4	ath - 5 10	
io result PCR		5	
elative accuracy elative specificy elative sensitivity	AC S P S P		81% 71% 85%

Attention! Truly negative results with Fastpath are judged as Fastpath+ This means that during serotyping no Lp SG1 was found.

New screening method for *Legionella pneumophila* in cooling water and process water © KWR

May 2009

			labl	Ν		lab I	
date 2008	company	sample code	Fastpath Nalo	i (Lp1) co	ISO 117	731	NEN 6265
				volume analvsed			
			result	(ml)	cfu/l	Serotype	Serotype
26-aug	A	OPIW 7	positive (6)	250	70.000	a+b+c	a+b+c
16-sep	A	OPIW 25	positive (6)	250	130.000	a+b+c	a+b+c
29-sep	A	OPIW 44	negative	250	60.000	a+b+c	a+b+c
26-aug	A	OPIW 8	positive (6)	250	1.800 222 222	<u>а</u> .	a+b
16-sep	A <	OPIW 26	positive (6)	250	320.000	a 4	0
29-Sep	₹ <		negative (6)	250		0 	440
20-aug	τ <		pusitive (u)	230	2.900	D+C	a T U
des-oi	× <	OPIW 2/	negative	750	008.1	D+C	D+C
29-sep	A	OPIW 45	negative	250	2.300	D+C	D+C
26-aug	в	OPIW 4	negative	250	18.000	p	q
16-sep	в	OPIW 22	negative	250	870	p	p
23-sep	в	OPIW 31	negative	250	530	a+b	p
26-aug	в	OPIW 6	positive	250	4.100	a+b	
16-sep	8	OPIW 23	positive	250	< 50		
23-sep	חת	OPIW 33	positive	250	3.600	a+b	
Zo-aug	ממ		positive (o)	750	11.000 rr 000	a+b	
des-oi	ממ		negative (6)	092	000.65	a+b	UD UD
ca-sep	0.0		positive (o)	230	00.00	a+b	
Z-Sep	ں در		negative	250	20	υ	
29-sep	5	OPIW 40	negative	092	< 50		
6-okt	<u>.</u>		negative	250	< 50	-	-
Z-sep	<u>.</u>	OPIW 13	negative	250	65.000 555	a+b	a+b
Z9-Sep		OPIW 41	positive (6)	250	530	a	a+b
6-okt	0	OPIW 50	positive (6)	250	420	a+b	a+b
Z-sep	<u> </u>	OPIW 14	negative	250	500 540	q	0
Z9-Sep	5	OPIW 42	negative	0920	210	0	D+C
0-OKI		OPIW 49	negative	067	200	D+C	a+b+c
2-sep		OPIW 10	positive	250	250.000	a+b	0
16-sep		OPIW 19	positive	250	210.000	a+b+c	a+b
26-aug		OPIW 1	positive	250	540 52 525	р.	
Z-Sep			positive	750	000.23	a+b	0
16-sep			positive	250	57.000	a+b	a+b
z-sep			positive	020	002 1	a+D	
10-sep			positive (6)	027	< 50	4	
23-55D			positive (0)	150	/ FD	a	0+0
6-okt	ц		positive	150	64 000	0+4+0	4+0
2-sen	1 11	OPIW 16	negative	250	< 50	a. n. c	0+0
23-sep	ш	OPIW 36	negative	250	96	C	
6-okt	ш	OPIW 54	negative	250	< 50		q
2-sep	ш	OPIW 17	positive	250	(1)		
23-sep	ш	OPIW 34	positive	250	70.000	a+b	q
6-okt	ш	OPIW 52	positive	250	69.000 (2)	a+b	a+b
23-sep	ш	OPIW 29	positive	75	< 50		
29-sep	L	OPIW 39	negative	250	510	c	
6-okt	L	OPIW 47	positive	250	960	b+c	
23-sep	ш	OPIW 30	positive	40	(1)		
29-sep	ш	OPIW 38	positive	40	(1)		
6-okt	ш	OPIW 46	positive	25	< 500		
23-sep	ш	OPIW 28	negative	50	< 50		
29-sep	<u></u>	OPIW 37	negative	40	< 50		
6-okt	L	OPIW 48	negative	10	< 50		

	Fathpath	52 samples	5 times false negative	5 times false positive	
	ш	ŝ	ŝ	ŝ	

49 samples			
ISO = reference met	thod		
Fastpath = alternatie	eve methoo	-	
Fastpath	+ Fastpat	- -	
. + OSI	. 22	5	
- OSI	5	7	
no result ISO		e	
relative accuracy	AC		80%
relative specificy	SP		58%
relative sensitivity	SE		86%

Attention! Truly negative results with Fastpath are judged as Fastpath+ This means that during serotyping no Lp SG1 was found.

New screening method for *Legionella pneumophila* in cooling water and process water © KWR

May 2009

VI Appendix: overview Q-PCR results

					(2)										(4)		(2)				Ţ	Ţ				T	Ι	Γ	Ι							(2)			T	Ţ		(4)	(4)	į	(ç)	Ţ	I	I	L			_
lab III	PCR (Lp)	kopieën/l	3,10E+03	1,70E+04		9,00E+03	1,63E+05	7,92E+04	<480	6,70E+03	<720	4,00E+05	2,69E+05	1,32E+04	1,00E+04	<1700		8,80E+04	7,53E+05	3,52E+05	<800	<720	9,50E+02	5,70E+03	<1200	1,50E+03		1 58F+03	1 226100	1,33E+04 8.50E+02	<480	2,14E+04	2,74E+04	<800	2,00E+03		<1500	006>	<800	2,90E+03	<1600	3,10E+03	4,80E+04	1,28E+05	-	< 1200 7660		<1200	<1000	<1400	<1200	<29000
		bol	3,5	4,2		4,0	5,2	4,9	2,4	3,8	2,6	5,6	5,4	4,1	4,0	2,9		4,9	5,9	5,5	2,6	2,6	3,0	3,8	2,8	3,2 7 7	ν, 2 Ο Ω	3 0 3 0	4.4	- 6	2.4	4,3	4,4	2,6	3,3		2,9	2,6	2,6 2 F	0'0 0	2,9	3,5	4,7	5,1	c	2'A	2.5 2.5	2 8 0	2.7	2,9	2,8	4,2
		log sd	2,4	2,9	3,8	1,2	3,3	4,1	2,6	1,9	3,6	3,0	3,5	3,9	3,4		2,8	4,7	4,1	4,6	3,1	2,3		4,1	2,2	2,0	0,0	0,4 0 7	ч, ч	4.3	2	3,5	2,0	2,5	3,2	3,5	2,8	4,2	• •	4, 1 1	7,5 1	C, 2	3,8 1	3,5	3,2	2,2	2 A 2 A	3.0	4.0	2		2,6
	to NEN	sd	273	266	6214	15	1791	12423	369	71	3616	965	2927	7891	2266		694	54747	13193	38684	1373	194		11393	148	400	341 2776	552	2002	20165		2829	102	354	1568	3270	703	15729	090	807	303	308	6625	30/8	1/6/	1/4	6603	1457	10284			416
lab II	d according	bc	3,0	1,3	1,4	2,4	1,4	1,8	8,	8,0	1,4	1,5	8'1	1,4	1,5	3,0	2,9	5,9	1,8	5,1	3,5	8	3,4	9	80	0,0	ς a	0, -		17	5	3,8	2,3	3,0	3,7	3,8	3,2	5,1 -	1,5 0		0	χı	15	5 A	2'D		15	2.5	5	2,5	2,5	3,2
	CR recalculate	pieën/l	02E+03	32E+04	75E+04 ²	59E+02	53E+04 ²	39E+04 ²	31E+02	38E+02	32E+04 ²	20E+04	70E+04	41E+04 ²	04E+04 4	2.13E+03	39E+02	16E+05 {	00E+04 ²	23E+05	16E+03	95E+02	32E+03	21E+04 4	91E+02			34F+03		51F+04	380	36E+03	31E+02	38E+03	51E+03	39E+03	56E+03	25E+05	80 201-00		0/E+03	01E+02	20E+04	04E+03	13E+03		51E+04	50E+03	36F+04	5.80E+02	3.80E+02	51E+03
	ă	log sd kc	4,3 1,	4,3 1,	3,4 <mark>2,</mark>	3,6 <mark>2,</mark>	4,2 <mark>2,</mark>	3,5 <mark>6,</mark>	3,2 6,	2,4 <mark>6,</mark>	3,1 <mark>2,</mark>	4,9 <mark>3,</mark>	4,6 <mark>5,</mark>	4,2 <mark>2,</mark>	2,7 <mark>3,</mark>	2,3 <	3,2 <mark>8,</mark>	8,	3,9 <mark>6,</mark>	4,0 1,	က်	Û.	<mark>. 1</mark> 2	3,3	2,6	γ <mark>α</mark>	<u>-</u> ע	0 8 1		3.6	V	2,7 <mark>6,</mark>	3,6 1,	0,9 <mark>1,</mark>	<u></u> 2,	4,0 <mark>5,</mark>	- - -	3,5	V u	Ó T	1 — 1	3,5	4,3	4,1	າ ເ	2'0 7	- ~	0 1 0	- <mark>-</mark>	v	v	_
	54	sd	20410,91	21749,99	2321,93	4183,07	16380,87	2834,72	1693,57	256,61	1333,99	75572,04	38319,38	16420,04	478,06	195,14	1497,68		8173,55	10501,85				2215,60	379,95	021,12		696 19	770.64	4193.87		448,97	4326,95	8,28		9661,27		3392,34			00 11 10	314/,68	20725,33	116/0,29	1000	12,201		479.00	00.01			
	o) N 62																	(11)									Τ	(12)	1-1								(11)								(11)	(12)	(11)	(1)			Π	(10)
lab IV	PCR (L	log	5,3	5,3	4,9	4,9	5,1	4,9	3,8	3,1	3,2	5,9	5,5	5,1	3,9	3,2	3,3		6,2	5,8	2,5	2,4	2,5	4,4	2,8	3,4 2 4	с,4 л л	C'7	1 1 1	4.9	2.5	3,7	4,3	3,3	2,4	4,5		5,2	2,6 2,6	ο, o	2,6		5,2	5,1	0	2'0 7	0,0	9 0	3.1	3,1	3,2	
	COL	kopieën/l	2,24E+05	2,0E+05	8,79E+04	7,99E+04	1,3E+05	8,51E+04	5,63E+03	1,2E+03	1,47E+03	8,22E+05	3,4E+05	1,27E+05	8,80E+03	1,7E+03	2,06E+03		1,6E+06	6,37E+05	<591	<551	<655	2,27E+04	6,64E+02	Z,/4E+U3	V349	<012 7 11F+02	207E±04	7.6F+04	< 699	5,06E+03	2,0E+04	1,79E+03	<508	2,98E+04		1,6E+05	<848	<044	<906>	1,29E+04	1,75E+05	1,2E+05	-	4,00E+02	- 020/	8 13E+03	<2688	<2430	<2931	,
	sample code		2PIW 7	JPIW 25	DPIW 44	DPIW 8	DPIW 26	DPIW 43	DPIW 9	DPIW 27	DPIW 45	DPIW 4	DPIW 22	DPIW 31	DPIW 6	DPIW 23	DPIW 33	DPIW 5	DPIW 24	DPIW 32	JPIW 15	DPIW 40	DPIW 51	DPIW 13	DPIW 41			DPIN/ 49		DPIW 19	DPIW 1	DPIW 11	DPIW 20	DPIW 12	DPIW 21	DPIW 18	DPIW 35	DPIW 53	DPIW 16		JPIW 54	71 MI40	DPIW 34	JPIW 52	DPIW 29	7 1/1 39	DPIM 30	DPIM 38	DPIW 46	DPIW 28	DPIW 37	DPIW 48
	s vneamos	·	A	A C	A	A	A	A	5	V	A	8	8	В	8	<u>е</u>	В	8	<u>о</u>	9	ບ ບ	<u>с</u>	0	0))) (0		<u> </u>	Ш															
	date 2008		26-aug	16-sep	29-sep	26-aug	16-sep	29-sep	26-aug	16-sep	29-sep	26-aug	6-sep	23-sep	26-aug	16-sep	23-sep	26-aug	6-sep	23-sep	e-sep	9-sep	5-okt	-sep	9-sep	POKI	n-sep	-okt		6-sen	6-aud	-sep	6-sep	-sep	6-sep	-sep	3-sep	Fokt	-sep	co-sep	-okt	-sep	3-sep	-okt	3-sep	ca-sep	2-cen	-0-sep	3-okt	23-sep	29-sep	i-okt

Attention: truly negativ results of Q-PCR are judged as + This means that no Lp was found during serotyping

45 samples ISO = reference method QPCR KWR = alternatieve metho		
ISO + ISO + ISO QPCR KWR + 33 QPCR KWR - 3	د ۵	
no result	7	
relative accuracy AC relative specificy SP relative sensitivity SE		91% 89% 92%

ISO vs D

45 samples

ISO = reference method QPCR D = alternatieve method	
ISO + ISO QPCR Labore+ 26 QPCR Labore- 8	N 6
no result	7
relative accuracy AC relative specificy SP relative sensitivity SE	78% 82% 76%
ISO vs GS	

	8 4	3	82% 33% 97%
amples hod ttieve method	36 + 1SO - 36 1		AC SP SP
49 s; GS = reference met QPCR GS = alterna	IS QPCR GS + QPCR GS -	no result	relative accuracy relative specificy relative sensitivity

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May 2009 New screening method for *Legionella pneumophila* in cooling water and process water © KWR - 58 - May 3

- Analysis inhibited by the samples matrix. Real number of Legionella pneumophila is probably higher
 - No Legionella pneumophila found. Analysis inhibited by the samples matrix; not reliable result.
 - Legionella pneumophila found, but result can not be quantified.

 $\begin{array}{c} (4) \\ (5) \\ (12$

- The output of the internal control sample is too low to give a reliable quantitative result
- Legionella pneumophila found, but the output of the internal control sample is too low to give a reliable quantitative result Legionella pneumophila concentration is indicative; below the lower limit for a reliable quantitative result

VII Appendix: Details statistical analysis

Test on differences between methods of Q-PCR analysis

Three laboratories – laboratory II, III and IV - each use a different method to analyse on Q-PCR. To examine the differences between the results of the three methods, we took 52 samples of cooling water and process water, coming from 20 locations at 6 sites (plants). These were all locations with a history of Legionella pneumophila occurrences. Most locations were sampled at three different periods, some at two different periods and some at only one period. Each sample was analysed by all three labs.

We tested statistically on differences between the results of the three labs, using analysis of variance (anova). If a sample result of one or more of the labs was missing, the results of that sample were not used. This was the case for 8 samples, so 44 samples remained for the analysis. Censored data were set at half the reporting limit. If the sample results of two or more labs were censored, they were set at half the lowest reporting limit for that sample, to avoid artificial differences between the labs. After that, each result was transformed by taking its logarithm. This was done to better meet the underlying assumption of analysis of variance that the residuals of the anova-model come from a normal distribution.

Figure 1 shows the boxplots of the logarithms of the results of the 44 samples for each lab.



Figure 1: Boxplots of the logarithms of the results of the 44 samples for each lab.

As can be seen from figure 1 the centres of the three boxplots do not differ much. However, the upper tails of the boxplots of lab IV and lab III are longer than that of lab II. This means that the high results of lab II tend to be lower than the high results of lab III and IV.

Using analysis of variance, we tested the null hypothesis that the mean of the logarithm of the results is the same for the three labs. The alternative hypothesis is that these means are not the same. We declared site, location and sample to be random factors and lab to be a fixed factor. Location was declared nested within site and sample nested within location. The F-test on the significance of the factor lab resulted in a p-value of 0.083, which means that the null hypothesis is not rejected (with 95% confidence). **Therefore**,

we did not find statistical significant differences between the means of the logarithms of the results of the three labs.

The residuals of the anova model conform to normality (with 95% confidence), as the Shapiro-Wilk test on normality of these residuals has a p-value of 0.321. So this condition for the use of anova was met.

Table 1 shows for each lab the estimated mean of the logarithms of the results and the lower and upper bound of the 95% confidence interval of the estimated mean.

Table 1: For each lab the estimated mean of the logarithms of the results and the lower and upper bound of the 95% confidence interval of the estimated mean.

Dependent Variable:log_PCR

			95% Confide	ence Interval
Lab	Mean	Std. Error	Lower Bound	Upper Bound
II	3.584 ^a	.092	3.402	3.766
IV	3.789 ^a	.092	3.607	3.971
	3.505 ^a	.092	3.323	3.687

a. Based on modified population marginal mean.

As can be seen from table 1 the confidence intervals of the three means show overlap.

Test on differences between methods of total Legionella analysis

To examine the differences between two methods to determine the concentration of total Legionella – ISO 11731 and NEN 6265:2007 (MWY) - we used the same 52 samples of cooling water and process water as described before. Each sample was analysed by both methods.

We tested statistically on differences between the results of the two methods, using analysis of variance (anova). If a sample result of one method was missing, the results of that sample were not used. This was the case for 16 samples, so 36 samples remained for the analysis. Censored data were set at half the reporting limit. If the sample results of both methods were censored, they were set at half the lowest reporting limit for that sample, to avoid artificial differences between the methods. After that, each result was transformed by taking its logarithm. This was done to better meet the underlying assumption of analysis of variance, that the residuals of the anova-model come from a normal distribution.

Figure 2 shows the boxplots of the logarithms of the results of the 36 samples for each method.





As can be seen from figure 2 the centres of the two boxplots do not differ much.

Using analysis of variance, we tested the null hypothesis that the mean of the logarithm of the results is the same for the two methods. The alternative hypothesis is that these means are not the same. We declared site, location and sample to be random factors and method to be a fixed factor. Location was declared nested within site and sample nested within location. The F-test on the significance of the factor method resulted in a p-value of 0.070, which means that the null hypothesis is not rejected (with 95% confidence). Therefore, we did not find a statistical significant difference between the means of the logarithms of the results of the two methods.

The residuals of the anova model conform to normality (with 95% confidence), as the Shapiro-Wilk test on normality of these residuals has a p-value of 0.909. So this condition for the use of anova was met.

Table 2 shows for each method the estimated mean of the logarithms of the results and the lower and upper bound of the 95% confidence interval of the estimated mean.

Table 2: For each method the estimated mean of the logarithms of the results and the lower and upper bound of the 95% confidence interval of the estimated mean.

			95% Confide	ence Interval
Method	Mean	Std. Error	Lower Bound	Upper Bound
ISO	3.475 ^a	.067	3.340	3.611
NEN	3.299 ^a	.067	3.164	3.435

Dependent Variable:log_Legion

a. Based on modified population marginal mean.

As can be seen from table 2 the confidence intervals of the two means show overlap.

VIII Appendix: Comparison cultivation methods

					(5)									(4)	151	(c)																		(2)						(4)	F	(2)	ÿ							
	۲ 5-2007	Serotype	a+b+c	a+b+c	a+b+c	a+b	_ q	a+b	b+c	b+c	p	þ	p					ט				a+b	a+b	a+b	U	b+c	a+b+c	9 	a+b	٤	a+b	•	a	b+c		a+b	b+c	4	α	2	a+h	2								
ab I	MM NEN 626	cfu/l	16.000	79.000	14.000	1.900 1 00 000	54.000	1.900	2.200	2.000	1.500	380	1.100	()				.1)	<500	< 500	< 500	30.000	980	360	98	320	190	10.000	49.000		29.000	(1)	38	11.000	< 500	25.000	1 <u>90</u>	< 500	0/7		55 000 (2)	(1)		< 500	(1)	Ē	(1)	(1)	(1)	(1)
	731	Serotype	a+b+c	a+b+c	a+b+c	<u>ہ</u>	2 0	b+c	b+c	b+c	q	q	a+b	a+b	4	a+b	a+b	a+b	0			a+b	q	a+b	A	υ.	b+c	a+b	a+b+c	4+0	a+b	a+b		q	Ì	a+b+c		0		47.0	a+5 4+	2			2					Ĭ
	150 11	cfu/l	70.000	130.000	60.000	1.800 320.000	220.000	2.900	1.900	2.300	18.000	870	530	4.100	< 50	3.000	11.000 55.000	55.000 65.000	98	< 50	< 50	65.000	530	420	500	210	200	250.000	210.000		57.000	1.500	< 50	10.000	< 50	64.000	< 50	96	< 50		69 000 (2)	< 50	510 510	960	(1)	(1)	< 500	< 50	< 50	< 50
	oneu	por de	1746		1458	9008	4192		876		652	652	354	121	609							8362		1095				12001	100/1			1365		4726																
	MWY p PH 7 3 & to	cfu/l	1600 (2)	(1)	6500	<100 27020 (8)	50300	<100	1040 (8)	<100	600 (8)	600	500 1010	1040 555 (5)	900 (8)	<100	(1)	(1)	<100	<100	<100	16800	<100	700	(1)	<100	<100	<u>120000 (2)</u>	8900 (8)		<100	2233 (2)	<100	25333 (2)	<100	(1)	<100	<100	<100	(1)		(1)	(1)	<100	(1)	(1)	(1)	<100	<100	<100
	pnue	sd by	2966		3029	612	20905				612						I						447	314																										
ab IV	BCYE	cfu/l	5400 (8)	(3)	3400	500	27000 (9)	(1)	(3)	(1)	500 (8)	(3)	<100	(1) (0)	(3)		(-)	(1)	<100	<100	<100	(1)	200	300	(1)	<100	<100	(1)	(3)	(1)	(1)	(1)	(3)	(1)	(1)	(1)	(1)	<100		(7) 00001.	(1)	(1)	(1)	(1)	(1)	(1)	(1)	<100	<100	<100
-	WY 865-2007	sd	2439	6124	2104	447 32823	9562		489	548	2598	1304	447	1245	QL /		1 2 7	101				8073	1823	1304	224	274	224							3525		3380														
	MEN 63	cfu/l	7800	38500	15600	300 146200	46600	(1)	300 (8)	006	3500	1200	300	1900 100 (0)	480 (8)	100	100 (0)	(1)	<100	<100	<100	28100	1700	1800	100	200	100	(1)	50000 (2)	(1)	()	(1)	<100	6400 (2)	<100	6100 (8)	<100	<100	001.×	(1)	<100	(1)	(1)	<100	(1)	(1)	(1)	<100	(1)	<100
	E + AB L 6265	sd	1151	5461		274	15441				179	1342	137				0760	P 040					600	1031	224	351	418																							
	BCY	cfu/l	1800 (2)	10200	(1)	300 (1)	37900 (2)	(1)	(1)	(1)	720 (8)	600 (2)	100 (2)	(L)	(1)	(1)		(1)	<100	<100	<100	(1)	500 (2)	1375	100 (2)	633 (2)	400	E) 5	(1)		()	(1)	(1)	(1)	(1)	(1)	<100	<100	00 L>	(E)	Ē	(1)	(1)	(1)	(1)	(1)	(1)	<100	(1)	(1)
	ebos elames		7 WIPC	DPIW 25	DPIW 44	DPIW 8	DPIW 43	6 MIdC	DPIW 27	DPIW 45	2PIW 4	DPIW 22	DPIW 31	OPIW 6				DPIW 24	DPIW 15	DPIW 40	DPIW 51	DPIW 13	DPIW 41	DPIW 50	DPIW 14	DPIW 42	DPIW 49	DPIW 10	DPIW 19		DPIW 20	DPIW 12	DPIW 21	DPIW 18	DPIW 35	DPIW 53	DPIW 16	DPIW 36	UPIW 54		DPIW 52	20 MIdC		DPIW 47	DPIW 30	DPIW 38	DPIW 46	DPIW 28	DPIW 37	DPIW 48
	, vieu mor		4	4	4			4	4	4	ő	ص ص							0			0	, U	0			۔ باری						0	U1	ЦП	<u>- п</u>														
	date 2008		26-aug	16-sep	29-sep	26-aug	29-sep	26-aug	16-sep	29-sep	26-aug	16-sep	23-sep	26-aug	10-sep	23-Sep	20-aug	23-sep	2-sep	29-sep	6-okt	2-sep	29-sep	6-okt	2-sep	29-sep	6-okt	2-sep	16-sep	2 con	2-sep 16-sep	2-sep	16-sep	2-sep	23-sep	6-okt	2-sep	23-sep	0-OKI	Z-Sep	6-okt	23-sen	20-sen	6-okt	23-sep	29-sep	6-okt	23-sep	29-sep	6-okt

- No Legionella found, result not reliable due to too much side growth on the agar medium
- Estmated value, real number possibly higher due to much side growth on the agar medium
- By mistake inoculated on BCYE-AB, pH 7.3. No Legionella found, result not reliable due to too much side growth on the agar medium
- Analysis inhibited by the samples matrix. Real number of *Legionella pneumophila* is probably higher No *Legionella pneumophila* found. Analysis inhibited by the samples matrix; not reliable result. (1) No Le
 (2) Estm:
 (3) By mi
 (4) Analy
 (5) Diffici
 (6) Diffici
 (7) Legio
 (8) Color
 (9) Thou
 (7) Serotypes: a =
 c =

 - Difficult to visibly detect line; line is very thin Legionella pneumophila found, but result can not be quantified.
- Colonies (partly) confirmed with PCR Though negativ at confirmation on plate, presence of *Legionella* is suspected
 - Legionella pneumophila serogroep 1.
- Legionella pneumophila serogroep 2-14. Legionella bacteria non pneumophila.

May 2009

				lab IV			lab I/labl	_	
				ΥWΜ			ΥWΜ		
date 2008	company	sample code	N مواريا	EN 6265:20	07 ed	ofiu/I	<u>IN 6265:20(</u>	07 Sarotuna	
26-aug	Δ	OPIW 7	7800	3.9	5439	16,000	4.2	athtc	
16-sep	A	OPIW 25	38500	4.6	6124	79.000	4.9	a+b+c	
29-sep	A	OPIW 44	15600	4,2	2104	14.000	4,1	a+b+c	(2)
26-aug	A	OPIW 8	300	2,5	447	1.900	3,3	a+b	
16-sep	A	OPIW 26	146200	5,2	32823	190.000	5,3	р	
29-sep	A	OPIW 43	46600	4,7	9562	64.000	4,8	р	
26-aug	A	OPIW 9	(1)			1.900	3,3	a+b	
16-sep	A	OPIW 27	300	2,5	489	2.200	3,3	b+c	
29-sep	A	OPIW 45	006	3,0	548	2.000	3,3	b+c	
26-aug	В	OPIW 4	3500	3,5	2598	1.500	3,2	р	
16-sep	В	OPIW 22	1200	3,1	1304	380	2,6	p	
23-sep	В	OPIW 31	300	2,5	447	1.100	3,0	q	
26-aug	в	OPIW 6	1900	3,3	1245	(1)			(4)
16-sep	в	OPIW 23	480	2,7	716	(1)			į
23-sep	в	OPIW 33	50	1,7		(1)			(2)
26-aug	я		(1)	000		(1)	0		
10-sep	חם	OPIW 24	100	2,0	13/	1.900	3,3	ø	
z3-sep	в В	OPIW 32	(1)	1		(1)			
Z-sep	5	OPIW 15	09	1,/		250	2,4		
29-sep	0	OPIW 40	50	1,7		250	2,4		
6-okt	: :	OPIW 51	50	1,7		250	2,4		
2-sep	0	OPIW 13	28100	4,4	8073	60.000	4,8	a+b	
29-sep	0	OPIW 41	1700	3,2	1823	980 365	3,0	a+b	
6-okt			1800	3,3	1304	86U 2 2	2,9	a+b	
2-sep	ບ	OPIW 14	100	2,0	224	98 262	2,0	с .	
Z9-sep	50		200	2,3	2/4	320	9 1 2	D+C	
0-OKI	נ נ	OPIW 49	001	z,U	724	490	2,1	a+b+c	
Z-sep			(1)	1		40.000	4,6	م	
16-sep		OPIW 19	20000	4,7		49.000	4,7	a+b	
26-aug		OPIW 1	(1)			(1)	1	_	
Z-sep	בו		(1)			4.900 60.000	3,7	<u>م</u>	
16-sep		OPIW 20	(1)			29.000	4,5	a+b	
Z-sep			(1)	1		(1)	4	,	
To-sep			00,00	1,/	1010	88	9. 1. 2.		í.
Z-sep	ц	OPIW 18	6400	3,8	3525	11.000	4,0	D+C	(၎)
23-sep	ш	OPIW 35	50	1,7	0000	250	2,4	-	
6-okt	ш	OPIW 53	<u>6100</u>	3,8 -	3380	25.000	4,4 0.0	a+b	
2-sep	ш	OPIW 16	50	1,7		1 <u>90</u>	2,3	b+c	
23-sep	ш	OPIW 36	50	1,/		250	2,4	4	
0-0KT	ш	OPIW 54	09	1,/		2/0	Z,4	۵	141
23 con	ц		ED	17			2 G	4	
5 01+	1 11		202	1,1		4.000	0,0	470	t)
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23-sep	ĿЦ		(1)			(1)			(G)
za-sep		OPIN 39		1		(1)			
6-okt	ш		50	1,/		250	2,4		
zo-sep						(1)			
Z9-Sep			(1)						
0-UKI	LU			1 7		(1)			
23-Sep	<u>. u</u>	0FIVV 20	00	1,1					
Z9-sep	<u> </u>	OFIV 3/	(_)	L .		(1)	_		
6-okt	LL.	OPIW 48	50	1.7		(1)			

conclusions

52 samples

29% MWY KWR 15 times (1) too much side growth

MWY D 16 times (1) too much side growth

31%

comparance

17%

 11 times results differ
 21%

 9 times too much side growth for at least 1 of 2 methods
 2 times < 100 versus > 100

In graphic all WTCB results < 500 as 250 ; KWR results < 100 as 50



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