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IMD-W™ Microbial Challenge

Test Results

Introduction

This fact sheet provides a summary of microbial test results obtained with the IMD-W instantaneous microbial detection™ system for water. The IMD-W system is a pharmaceutical water quality monitoring and risk management tool. This system has been designed for the routine monitoring of Purified Water (PW) and Water for Injection (WFI) systems, including distribution loops, storage tanks and points-of-use (POU). The IMD-W system can be used to continuously monitor a pharmaceutical water loop and in conventional point sampling applications. Challenge organisms were chosen based on guidance provided in the United States and European Pharmacopeias, and by the Online Water Bioburden Analyzer (OWBA) working group.^A Testing was designed to show the capability and sensitivity of the IMD-W system in detecting planktonic (single, free-floating) microorganisms through the assessment of thirteen industry relevant organisms.

Background

Water plays a predominant role in the formulation and manufacture of pharmaceutical products. The traditional culture-based methods commonly used to ensure water quality, however, provide an episodic view of quality at best. Other process monitoring tools such as those for Total Organic Carbon (TOC) and conductivity also play a predominant role, but sensitivity on the order of a single microorganism is outside of the target range for such systems. The pharmaceutical industry continues to recognize a need to implement modern technologies to drive risk reduction and process control, as evident in guidance such as the FDA's 2004 Guidance for Industry document on Process Analytical Technology (PAT), ICH Guidelines Q8, Q9 and Q10, and the FDA's Pharmaceutical cGMPs for the 21st Century, which encourage the adoption of Quality by Design (QbD) principles and new technologies. The OWBA working group, composed of representatives from key pharmaceutical companies, has articulated the industry's need for a real-time system for water quality assessment, with an overall goal of encouraging the development and use of such new technologies.^A

The fundamental method of microbial detection is different between traditional culture-based methods and bio-fluorescent particle (BFP) systems based on light-induced fluorescence (LIF) such as the IMD-W system. Traditional culture-based methods require cell

proliferation, leading to the formation of a visually detectable colony-forming unit (CFU) to indicate microorganism presence. Yet, with common media such as Tryptic Soy Agar (TSA), and typical incubation parameters, not all organisms are culturable. POU sampling with traditional culture-based methods is a currently accepted and primarily practiced method for assessing pharmaceutical water quality. POU testing may be performed as infrequently as once every two weeks at each sample point, resulting in limited sampling data and retrospective culture-based results. A LIF-based detection method does not require cell growth and is not restricted by limitations such as an incompatible media or incubation conditions. The IMD-W system offers the ability to perform POU testing and continuous monitoring, making the system an excellent tool for use in trending, risk reduction and process control. To further evaluate the system's microbial detection capability, its sensitivity was assessed.

Test Parameters

I. Microbial Species Tested

Thirteen industry relevant organisms were utilized to challenge the IMD-W system and assess its detection performance. These organisms include *Burkholderia cepacia* (NBRC 14595), *Brevundimonas diminuta* (ATCC 19146), *Escherichia coli* (ATCC 8739), *Methylobacterium extorquens* (NBRC 15911), *Pseudomonas aeruginosa* (ATCC 9027), *Pseudomonas fluorescens* (ATCC 17386), *Ralstonia pickettii* (ATCC 27511), *Salmonella enterica* (NCTC 6017), *Stenotrophomonas maltophilia* (ATCC13637), *Bacillus subtilis* (ATCC 6633), *Staphylococcus aureus* (ATCC 6538), *Aspergillus brasiliensis* (ATCC 16404), and *Candida albicans* (ATCC 10231). *A. brasiliensis*, *B. cepacia*, *C. albicans*, *E. coli*, *M. extorquens*, *P. aeruginosa*, *P. fluorescens*, *R. pickettii*, *S. aureus*, *S. enterica*, and *S. maltophilia* were purchased from the National Institute of Technology and Evaluation (NITE) biological resource center. *B. diminuta* was purchased from the RIKEN BioResource Center, and the *B. subtilis* spore suspension was obtained from MesaLabs (Ref: SUS-1A-6). *A. brasiliensis* and *B. subtilis* were in the spore state, while all other microorganisms were tested as vegetative cells.

^A Cundell, A., Gordon, O., Haycocks, N., Johnston, J., Luebke, M., Lewis, N., et al. (2013, May/June). *Novel Concept for Online Water Bioburden*

Analysis: Key Considerations, Applications, and Business Benefits for Microbiological Risk Reduction. American Pharmaceutical Review, 26-31

Organisms were chosen based on guidance set forth in USP <61>, USP <62>, USP <71>, EP 2.16.12, EP 2.6.13, and by the OWBA working group. ^A All seven compendial microorganisms listed in the OWBA testing protocol were tested with the IMD-W system and the traditional culture-based method for recovery comparison. ^B According to the testing protocol, the purpose of testing these organisms was to verify that an online water bioburden analyzer system is capable of enumerating the indicator aerobic QC microorganisms listed in USP <61>, USP <62>, and USP <71>. ^B *R. pickettii*, *S. maltophilia*, *P. fluorescens* and *B. cepacia* also are highlighted in the testing protocol as four examples of common waterborne and environmental bacteria that could be tested as part of industrial isolate testing. In an effort to challenge the IMD-W system further, *B. diminuta* was selected for its small size and *M. extorquens* as another waterborne organism of interest.

Table 1 shows the thirteen microorganisms tested and whether or not the organism is specified in the USP, EP and OWBA documents referenced. A checkmark indicates that the tested organism is listed in the referenced document, while a dash indicates that the organism is not. All organisms listed in USP <61> were tested. For USP <62>, EP 2.16.12 and EP 2.6.13, the only listed organisms not yet tested on the IMD-W system are one of the two listed *S. enterica* strains and the anaerobe *Clostridium sporogenes*. With regard to USP <71>, all listed organisms have been tested except the anaerobe *C. sporogenes*, its alternate *Bacteroides vulgatus*, and *Micrococcus luteus*, the alternate for the tested *P. aeruginosa*.

Microorganism Tested	USP <61>	USP <62>	USP <71>	EP 2.16.12	EP 2.6.13	OWBA
<i>B. cepacia</i>	--	--	--	--	--	✓
<i>B. diminuta</i>	--	--	--	--	--	--
<i>E. coli</i>	--	✓	--	✓	✓	✓
<i>M. extorquens</i>	--	--	--	--	--	--
<i>P. aeruginosa</i>	✓	✓	✓	--	✓	✓
<i>P. fluorescens</i>	--	--	--	--	--	✓
<i>R. pickettii</i>	--	--	--	--	--	✓
<i>S. enterica</i>	--	✓	--	✓	✓	✓
<i>S. maltophilia</i>	--	--	--	--	--	✓
<i>B. subtilis</i>	✓	--	✓	--	--	✓
<i>S. aureus</i>	✓	✓	✓	✓	✓	✓
<i>A. brasiliensis</i>	✓	--	✓	--	--	✓
<i>C. albicans</i>	✓	✓	✓	✓	✓	✓

Table 1: Microorganisms tested and their mention in regulatory/guidance documents. A few other organisms listed in referenced guidance have not yet been tested (the second strain on *S. enterica*, the anaerobe *C. sporogenes*, and "alternate" organisms *B. vulgatus* and *M. luteus*.)

II. Microbe Preparation

Vegetative organisms including *B. diminuta*, *E. coli*, *P. aeruginosa*, *S. aureus*, and *S. enterica* were inoculated from their glycerol stock in Tryptic Soy Broth (TSB) and cultured aerobically overnight at 32°C while *B. cepacia* was cultured aerobically for two days at 32°C. The bacteria were then streaked onto TSA and incubated at approximately 32°C for 20 to 30 hours, and 40-48 hours for *B. cepacia*, to achieve the stationary phase. *M. extorquens* was cultured in a laboratory-prepared liquid medium for four days at 32°C, streaked onto R2A, and incubated at approximately 32°C for four days. *R. pickettii* was cultured in a laboratory-prepared nutrient broth for three days at 32°C, streaked onto R2A, and incubated at approximately 32°C for three days. *S. maltophilia* and *P. fluorescens* were cultured in R2A for two days at 32°C, streaked onto R2A, and incubated at approximately 32°C for 40 to 48 hours. *C. albicans* was cultured in Sabouraud glucose broth for 40 to 48 hours at 25°C, streaked onto Sabouraud glucose agar, and incubated at approximately 25°C for 40 to 48 hours. The bacteria were then harvested in sterile, distilled water (DW) and washed through centrifugation at 2,100g for three minutes. The supernatant was removed and the pellet was resuspended in filtered DW. Optical density measurements at 600 nm (OD₆₀₀) then were utilized to estimate concentration, followed by dilution with filtered DW to reach the target microbial concentration. This dilution was utilized in the system testing and plated on TSA to perform a final titer check.

B. subtilis and *A. brasiliensis* spore suspensions were prepared following a different procedure. *B. subtilis* spore suspensions were diluted with filtered DW directly from the stock suspension. *A. brasiliensis* spores were inoculated from the stock culture onto Sabouraud glucose agar at 25°C for approximately seven days. Phosphate buffer supplemented with 0.05% Tween 80 solution was utilized to recover *A. brasiliensis* spores from the culture plate. The spore suspension was then filtered through eight layers of sterile gauze to remove hyphae and centrifuged at 1,600g for ten minutes to wash the spores. The supernatant was removed, the pellet was resuspended in filtered DW, and the suspension was centrifuged at 1,400g for ten minutes to continue washing. The second centrifugation step was repeated three times. The supernatant was again removed and the pellet was resuspended in filtered DW. Microscopy was utilized to confirm the absence of hyphae and determine the suspension concentration. Dilution to the desired concentration was performed with filtered DW and a final titer check was performed.

Five target concentrations were tested for each microorganism, except *B. diminuta* which was tested at three concentrations, including 0.1 CFU/mL, 1 CFU/mL, 10 CFU/mL, 100 CFU/mL and 1000 CFU/mL.

^B Cundell, A., Luebke, M., Gordon, O., Mateffy, J., Haycocks, N., Weber, J. W., et al. (2013, April 24). On-Line Water Bioburden Analyzer Testing Protocol. Document ID OWBA-TP-2013-v1.5.

Six replicates were performed at each concentration tested. Concentrations were chosen to ensure that the IMD-W system has an appropriate sensitivity to assess the current compendial limit of 10 CFU/mL for WFI.

III. Test Systems and Apparatus

Testing was completed with two IMD-W systems, a Rion KS-42B liquid particle counter, and a Pall MicroFunnel manifold with 0.45µm disposable filter for water sampling and sample culture on TSA plates.

A water loop shown in **Figure 1** was designed specifically for this testing in order to obtain a very clean background adequate for low-level microbial injections. An ultrapure water supply and inline 0.05µm filter permitted extremely low background particulate counts in the loop such that microorganism testing down to 0.1 CFU/mL was possible. A sample injector was utilized to introduce small microbial samples into the loop. The Rion liquid particle counter was utilized as a reference system to confirm particulate counts within the loop during microbe sampling. The sample preparation and static mixer were used to create a homogeneously-dispersed, planktonic sample within the water loop before concurrent sampling by the four instruments. Water samples were filtered and plated on TSA to obtain traditional culture-based results for comparison to IMD-W data.

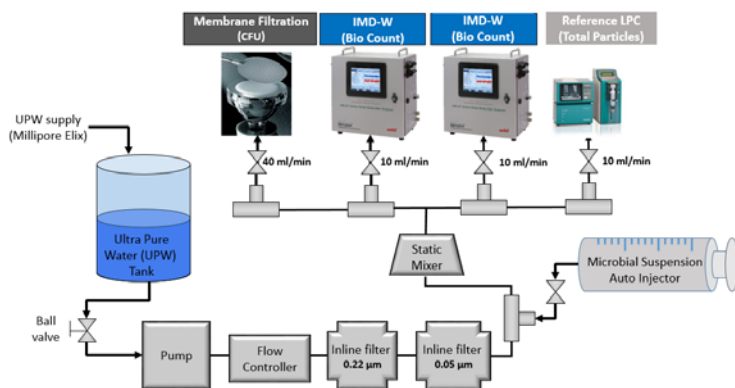


Figure 1: Custom-built water loop for microbial testing.

Test Results

The IMD-W system was challenged with thirteen microorganisms. Testing was designed such that single cells, as opposed to agglomerates, were sampled by the IMD-W system to ensure sensitivity down to the level of intrinsic fluorescence emitted by planktonic microbes. **Figure 2** shows a summary of microorganism results for the thirteen organisms tested on the IMD-W system and the traditional culture-based method, separated based on waterborne versus non-waterborne microorganisms. IMD-W *S. maltophilia* biologic count results were lower than expected based on other testing performed with this organism. Additional testing and investigation is planned to study this organism further.

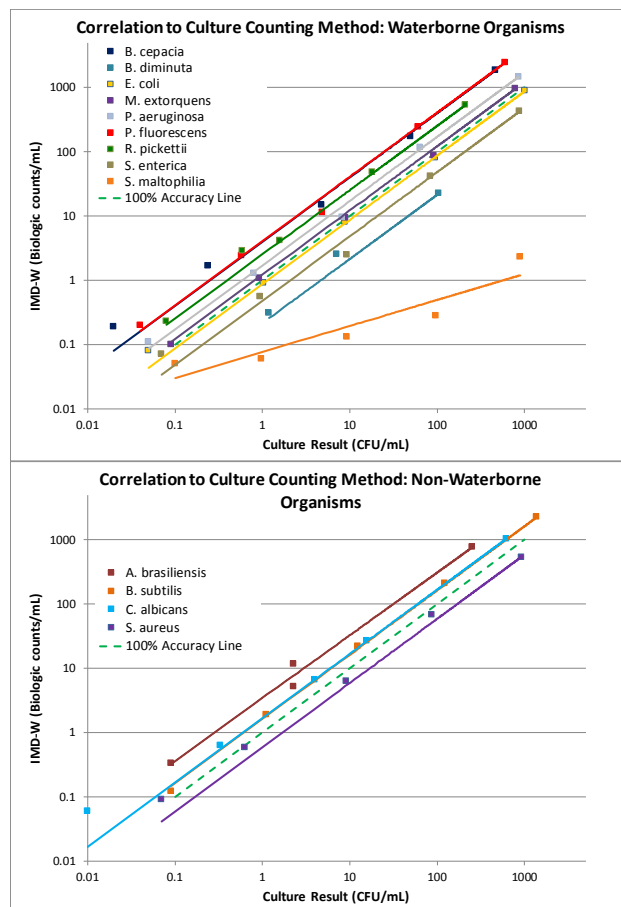


Figure 2: IMD-W system and culture plate results for nine waterborne and four non-waterborne industry-relevant microorganisms. The dashed green line represents the 100% Accuracy Line. Organisms above this line showed higher counts on the IMD-W system than on the reference culture-based method.

IMD-W *B. diminuta* biologic counts versus traditional method recoveries were also lower than seen for the other organisms tested, in this case, believed to be the result of its very small size. Note that the lowest concentration data point is not indicative of the IMD-W system's limit of detection (LOD), but is instead based on the minimum concentration tested in this challenge testing. The testing of very low microbial concentrations is quite difficult due to the necessity for an extremely clean background. As shown in **Figure 1**, a custom-built water loop with a 0.05µm in-line filter was required for testing down to even 0.1 CFU/mL.

IMD-W biologic count results correlate well with culture-based results for the organisms tested. This is shown by the R^2 values in **Table 2** that are close to a value of one, indicating a high correlation in the results from both techniques across the concentration range tested. The variability in recovery seen across organisms in **Figure 2**, however, make it such that a conversion factor between CFU and biologic counts cannot be applied to these fundamentally different detection methods. This is due to differences in size, fluorescence and culturability of these organisms, for example. Although there is variability in recovery, all thirteen organisms were detected by the IMD-W system down to very low concentration levels, as shown by the data.

Microorganism Tested	Coefficient of Determination (R^2)
WATERBORNE	
B. cepacia	0.9998
B. diminuta	0.9982
E. coli	1.0000
M. extorquens	0.9999
P. aeruginosa	0.9978
P. fluorescens	0.9989
R. pickettii	0.9993
S. enterica	0.9974
S. maltophilia	0.9370
NON-WATERBORNE	
A. brasiliensis	1.0000
B. subtilis	1.0000
C. albicans	0.9999
S. aureus	0.9991

Table 2: Coefficient of determination (R^2) values are shown for the relationship between IMD-W biologic counts and culture-based CFU results. A value close to one shows a high level of correlation in the results from both methods.

Conclusion

- The IMD-W system is capable of continuous and real-time bioburden monitoring, and offers end users the ability to monitor water system control and react to out-of-specification events in a much timelier manner than with episodic/traditional methods alone.
- This microbial testing challenged the IMD-W system with thirteen industry relevant microorganisms to determine the system's ability to detect a range of organisms down to the single cell level.
- The IMD-W system is capable of single-cell detection for all organisms tested.
- With a focus on sensitivity and the ability to monitor water systems continuously and in real-time, the IMD-W system is a powerful monitoring and trending tool capable of increasing product quality assurance and process understanding.

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