Annals of Clinical Microbiology and Antimicrobials



Research Open Access

Verification of the ProPneumo-I assay for the simultaneous detection of Mycoplasma pneumoniae and Chlamydophila pneumoniae in clinical respiratory specimens

Rachel R Higgins¹, Ernesto Lombos¹, Patrick Tang¹, Karl Rohoman¹, Anne Maki¹, Shirley Brown¹, Frances Jamieson^{1,2} and Steven J Drews*^{1,2,3}

Address: ¹Ontario Public Health Laboratories, Ministry of Health and Long-Term Care, Toronto, Ontario, Canada, ²Department of Pathobiology and Laboratory Medicine, University of Toronto, Toronto, Ontario, Canada and ³Mount Sinai Hospital, Toronto, Ontario, Canada

Email: Rachel R Higgins - rachel.higgins@ontario.ca; Ernesto Lombos - ernesto.lombos@ontario.ca; Patrick Tang - patrick.tang@ontario.ca; Karl Rohoman - karl.rohoman@ontario.ca; Anne Maki - anne.maki@ontario.ca; Shirley Brown - shirley.brown@ontario.ca; Frances Jamieson - frances.jamieson@ontario.ca; Steven J Drews* - steven.drews@ontario.ca

Published: 10 March 2009

Annals of Clinical Microbiology and Antimicrobials 2009, 8:10 doi:10.1186/1476-0711-8-10

This article is available from: http://www.ann-clinmicrob.com/content/8/1/10

© 2009 Higgins et al; licensee BioMed Central Ltd.

This is an Open Access article distributed under the terms of the Creative Commons Attribution License (http://creativecommons.org/licenses/by/2.0), which permits unrestricted use, distribution, and reproduction in any medium, provided the original work is properly cited.

Received: 28 November 2008 Accepted: 10 March 2009

Abstract

Background: Mycoplasma pneumoniae and Chlamydophila pneumoniae are major causes of lower and upper respiratory infections that are difficult to diagnose using conventional methods such as culture. The ProPneumo-I (Prodesse, Waukesha, WI) assay is a commercial multiplex real-time PCR assay for the simultaneous detection of M. pneumoniae and/or C. pneumoniae DNA in clinical respiratory samples.

Objective: The aim of this study was to evaluate the sensitivity and specificity of the ProPneumo-I, a newly developed commercial multiplex real-time PCR assay.

Methods: A total of 146 clinical respiratory specimens, collected from 1997 to 2007, suspected of *C. pneumoniae* or *M. pneumoniae* infections were tested retrospectively. Nucleic acid was extracted using an automated NucliSense easyMag (bioMerieux, Netherlands). We used a "Homebrew" monoplex real-time assay as the reference method for the analysis of *C. pneumoniae* and culture as the reference method for the analysis of *M. pneumoniae*. For discordant analysis specimens were re-tested using another commercial multiplex PCR, the PneumoBacter-I assay (Seegene, Korea).

Results: Following discordant analysis, the sensitivity of the ProPneumo-I assay for pathogens, *C. pneumoniae* or *M. pneumoniae*, was 100%. The specificity of the ProPneumo-I assay, however, was 100% for *C. pneumoniae* and 98% for *M. pneumoniae*. The limits of detection were I genome equivalent (Geq) per reaction for pathogens, *M. pneumoniae* and *C. pneumoniae*. Due to the multipex format of the ProPneumo-I assay, we identified 5 additional positive specimens, 2 *C. pneumoniae* in the *M. pneumoniae*-negative pool and 3 *M. pneumoniae* in the *C. pneumoniae*-negative pool.

Conclusion: The ProPneumo-I assay is a rapid, sensitive and effective method for the simultaneous detection of *M. pneumoniae* and *C. pneumoniae* directly in respiratory specimens.

^{*} Corresponding author

Background

Mycoplasma pneumoniae and Chlamydophila pneumoniae are major causes of lower and upper respiratory tract infection in adults and children [1-3] with high incidence occurring among schoolchildren [4,5]. Both pathogens are difficult to diagnose using conventional methods such as culture and serology. Furthermore, respiratory infection may be caused by one of several bacterial or viral pathogens that share similar symptoms and clinical features. To overcome difficulties in the detection of these pathogens, PCR assays have been developed for the rapid detection and identification of pathogens directly in respiratory samples [6]. PCR assays are widely used in the clinical arena with real-time PCR slowly replacing conventional PCR assays. Several "home-brew" single target (monoplex) assays have been utilized for the detection of numerous pathogens, including M. pneumoniae [7-9]. However, there are multiple regulatory issues with the use of "home-brew" assays and controversies exist as to their use in a diagnostic laboratory setting. As commercial assays become more available, there is a growing movement to replace "homebrew" assays with commercially available products [10,11]. As well, the monoplex PCR format is being replaced by a multiplex one. The aim of this study was to evaluate the analytical sensitivity and the diagnostic specificity and sensitivity of the ProPneumo-1 assay, a newly developed commercial multiplex real-time PCR assay for the simultaneous detection of M. pneumoniae and/or C. pneumoniae DNA in clinical respiratory specimens.

Methods

Specimen collection and analysis by the reference method

Historically, suspected *C. pneumoniae* infections were tested using a "home-brew" single target monoplex real-time PCR assay [12] and suspected *M. pneumoniae* infections was tested using culture. As shown in Table 1, we examined a total of 146 clinical respiratory specimens; of which 58 were previously tested for *C. pneumoniae* using a "home-brew" real-time PCR and 88 were previously tested for *M. pneumoniae* using culture. Specimens were collected

in Ontario, between 1997 and 2007, from patients with suspected *C. pneumoniae* or *M. pneumoniae* infections. The types of specimen submitted for testing at the Central Public Health Laboratory in Toronto, Ontario, including patient's sex and age are described in Table 2 for *M. pneumoniae*, and 2B for *C. pneumoniae*. Specimens and extracted nucleic acid were stored at -80°C until use.

DNA for "home-brew"monoplex real-time PCR was extracted from specimens using the MagNA Pure LC (Roche, Mannheim, Germany) as per manufacturer's instructions. Specimens were tested using a "home-brew" single target monoplex real-time PCR assay [12]. Real-time amplification was carried out using the Roche Light Cycler (Roche, Mannheim, Germany), as previously published [12,13].

Mycoplsma pneumoniae

Clinical specimens from patients with suspected *M. pneumoniae* infection were cultured in PPLO medium containing thallium acetate, horse serum, yeast dialysate and supplemented with amphotericin B (0.5 mg/ml), penicillin G (100,000 u/ml) and nystatin (50,000 u/ml). Biphasic culture flasks were inoculated with specimens, incubated at 37°C and inspected daily for 4 weeks. The organism was identified based on typical colonial morphology on the agar medium and the change in the broth color from red to orange then to yellow in the absence of turbidity of the broth.

ProPneumo-I and PneumoBacter-I assays

Nucleic acid, for analysis by the ProPneumo-1 (Prodesse, Waukesha, WI) or the PneumoBacter-1 (Seegene, Korea) assay, was extracted using the easyMag NucliSense magnetic extraction (bioMerieux, Netherlands) [14]. Each nucleic acid extract was amplified by both, the PromPneumo-1 and PneumoBacter-1 multiplex PCR assays. The ProPneumo-1 assay was performed as per manufacturer's instructions. Amplification and detection of the ProPneumo-1 test was performed on the ABI 7500 (Applied

Table I: Summary of results for the evaluation of the ProPneumo-I multiplex real-time PCR assay

	C. pr	C. pneumoniae		M. pneumoniae		
	Monoplex	ProPneumo-I	Culture	ProPneumo-I		
Positive specimens	11	14	42	41		
Negative specimens	47	43	46	48		
Total	58	57	88	89		

Note: The ProPneumo-I assay detected 3 positive M. pneumoniae specimens in the C. pneumoniae-negative pool and 2 positive C. pneumoniae specimens in the M. pneumoniae-negative pool. Accordingly, 57 and 89 specimens were resolved for C. pneumoniae and M. pneumoniae, respectively by the ProPneumo-I assay compared to 58 and 88 by the Monoplex assay and culture, respectively. The total number of specimens tested remains 146.

Table 2: Overview of test results for a subset of M. pneumoniae patients

Patient No	Sex	Age	Specimen	Pathogen	Culture	Prodesse PP-I	Seegene PB-I
01	М	0.75	ETA	Mpn	+	+	+
02	М	32	NPS	Mpn	+	+	+
03	F	36	Sputum	Mpn	+	+	+
04	F	40	ETA	Mpn	+	+	+
05	М	26	Sputum	Mpn	+	+	+
06	F	56	Sputum	Mpn	+	+	+
07	F	72	Sputum	Mpn	+	+	+
08	F	26	BR wash	Mpn	+	+	+
09	F	7	TS	Mpn	+	+	+
10	М	NA	BAL	Mpn	+	+	+
П	F	П	NPS	Mpn	+	+	+
12	М	9	NPS	Mpn	+	+	+
13	NA	NA	NA	Mpn	+	+	+
14	М	3	NPS	Mpn	+	+	+
15	М	3	NPS	Mpn	+	+	+
16	М	6	NPS	Mpn	+	+	+
17	М	2	NPA	Mpn	+	+	+
18	М	13	NPS	Mpn	+	+	+
19	М	13	Sputum	Mpn	+	+	+
20	F	17	NPS	Mpn	+	+	+
21	F	13	NPS	Mpn	+	+	+
22	М	47	Sputum	Mpn	+	+	+
23	F	14	NPS	Mpn	+	+	+
24	F	5	NPS	Mpn	+	+	+
25	F	70	Sputum	Mpn	+	+	+
26	F	5	NPS	Mpn	+	+	+
27	М	44	Sputum	Mpn	+	+	+

Table 2: Overview of test results for a subset of M. pneumoniae patients (Continued)

28	F	43	TS	Mpn	+	+	+
29	М	4	Sputum	Мрп	+	+	+
30	F	7	TS	Мрп	+	+	+
31	М	5	TS	Mpn	+	+	+
32	М	15	ETA	Mpn	+	+	+
33	М	10	TS	Мрп	+	+	+
34	М	8	TS	Mpn	+	+	+
35	F	23	Pharyngeal	Мрп	+	+	+
36	М	50	TS	Мрп	+	+	+
37	F	7	BAL	Mpn	+	+	+
38	NA	NA	NA	Мрп	+	+	+
39	F	61	Sputum	Мрп	+	-	-
40	F	7	NPS	Mpn	+	-	-
41	М	6	NPS	Mpn	+	-	-
42	М	18	TS	Mpn	+	-	-
43	М	9	NPS	Мрп	-	+	-
*44	NA	NA	NPS			+	+
*45	NA	NA	NPS			+	+
*46	NA	NA	NPS			+	+

Biosystems, Foster City, CA). For discordant analysis all specimens, including concordant specimens, were retested using the PneumoBacter-1 assay (Seegene, Korea). PneumoBacter-1 is an agarose gel-based assay, amplification was performed on the iCycler (Bio-Rad, Milpitas, CA) and amplicons were imaged, following gel electrophoresis, as per manufacturer's protocols.

Experimental controls: external, internal and extraction controls

M. pneumoniae and C. pneumoniae quantified DNA controls (Advanced Biotechnologies Inc., Columbia, MD) were included in each assay as external positive controls. For internal DNA control and to monitor for potential PCR inhibitors, exogenous DNA was spiked into each specimen. To monitor for the integrity of extracted nucleic acid, all specimens were also tested for the presence of the

housekeeping gene, human glyceraldehyde phosphate dehydrogenase (GAPDH) (Applied Biosystems. Foster City, CA). Real-time amplification and detection of the GAPDH gene sequence was performed on the ABI 7900HT (Applied Biosystems, Foster City, CA), as previously described [15,16]. This test was required to ascertain the presence of nucleic acid in specimens, which tested negative for *C. pneumoniae* or *M. pneumoniae*. For negative controls, target nucleic acid and specimens were replaced by PCR grade water in each PCR assay or nucleic acid extraction run, respectively.

Limits of detection

The limits of detection for the assays described in this study, ProPneumo-1 and PneumoBacter-1, were determined using 10 fold serial dilutions of quantified *M. pneumoniae* and *C. pneumoniae* DNA (Advanced

Biotechnologies Inc., Columbia, MD) in PCR-grade water. Data analysis was carried out using SPSS 15.0 (SSPS Inc., Chicago, IL). Turn-around-times per assay were estimated from a review of laboratory procedures and include time from specimen reception to results reporting.

Definitions

The following definitions were used for analysis. Concordant specimen: agreement between "home-brew" assay result and Prodesse ProPneumo-1 assay result. Discordant specimen: disagreement between "home-brew" assay result and Prodesse ProPneumo-1 assay result. Resolved specimen: agreement in result, either positive or negative between any two of the following assays; 1) "home-brew" or culture, 2) ProPneumo-1, and 3) PneumoBacter-1.

Results

Table 1 summarizes the results obtained for the evaluation of the ProPneumo-1 real time multiplex PCR assay. Overviews of the results obtained for subsets of M. pneumoniae and *C. pneumoniae* patients are presented in Table 2 and Table 3, respectively. Among patients who tested positive for either pathogen, the ratio of male to female is approximately the same and, with the majority of positive patients being under the age of 20 years. Table 2 also displays the results obtained for each of the three tests used to identify positive M. pneumoniae (A) or C. pneumoniae (B) patients. Using a monoplex "home-brew" real-time PCR as a reference in the analysis of C. pneumoniae and culture as a reference in the analysis of M. pneumoniae, 14 and 41 specimens tested positive for C. pneumoniae and M. pneumoniae, respectively by the ProPneumo-1 assay. As shown in Table 1, 11 patients tested positive for C. pneumoniae by the "home-brew" monoplex assay and 42

Table 3: Overview of test results for a subset of C. pneumoniae patients

Patient No	Sex	Age	Specimen	Pathogen	Monoplex PCR	Prodesse PP-I	Seegene PB-I
01	М	13	BAL	Cpn	+	+	+
02	М	0.2	NPS	Cpn	+	+	+
03	F	10	BAL	Cpn	+	+	+
04	F	10	ETA	Cpn	+	+	+
05	F	2	BAL	Cpn	+	+	+
06	F	10	BAL	Cpn	+	+	+
07	F	63	NPS	Cpn	+	+	+
08	М	32	NPS	Cpn	+	+	+
09	F	63	TS	Cpn	+	+	+
10	F	19	NA	Cpn	+	+	+
11	М	48	BAL	Cpn	+	-	-
12	NA	NA	Autopsy	Cpn	-	+	+
13	NA	NA	Sputum	Cpn	-	+	+
** 4	F	25	NPS			+	+
**15	F	44	NPS			+	+

Abbreviations: NA, not available; M, male; F, Female; NPS, nasopharyngeal swab; NPA, nasopharyngeal aspirate; BAL, bronchoalveolar lavage; ETA, endotracheal aspirate; TS, throat swab; BR, bronchial; Cpn, C. pneumoniae; Mpn, M. pneumoniae; PP-I, ProPneumo-I; PB-I, PneumoBacter-I. *, These specimens tested negative for C. pneumoniae in the monoplex assay but were defined positive for M. pneumoniae by the ProPneumo-I and PneumoBacter-I assays. **, These specimens tested negative for M. pneumoniae in culture but were defined positive for C. pneumoniae by the ProPneumo-I and PneumoBacter-I assays.

patients tested positive for *M. pneumoniae* by culture. Similarly, 48 and 43 specimens were defined as negative in the ProPneumo-1 assay, for *M. pneumoniae* and *C. pneumoniae*, respectively. Although, 42 positive *M. pneumoniae* specimens were identified by culture and 41 by the ProPneumo-1assay, the identity of 8 positive specimens however, varies between the two methods (see Table 2). Curiously, both assays the ProPneumo-1 and Pneumo-Bacter-1 detected 5 additional positive specimens, 3 for *M. pneumoniae* and 2 for *C. pneumoniae* in the *C. pneumoniae*-negative and *M. pneumoniae*-negative pool, respectively.

The limit of detection of the ProPneumo-1 assay as determined by Probit regression (95% probability) was estimated as follows: 1 genome equivalent (Geq)/reaction for *M. pneumoniae* and 1 Geq/reaction for *C. pneumoniae*. The limit of detection for PneumoBacter-1 was 1 Geq/reaction for *M. pneumoniae* and 1 Geq/reaction for *C. pneumoniae*. The turn-around-times, from specimen reception to reporting, are as follows: ProPneumo-1, 6 hours; "homebrew" monoplex PCR, 10 hours; PneumoBacter-1, 12 hours, and Mycoplasma culture, 4 weeks.

Diagnostic specificity and sensitivity

To determine the diagnostic specificity and sensitivity of the ProPneumo-1 assay, results were scored against the reference method ["home-brew" monoplex real-time PCR for *C. pneumoniae* (see Table 4) and culture for *M. pneumoniae* (Table 5)]. To resolve discordant results, specimens were re-tested using the multiplex PneumoBacter-1 method (Tables 4 and Table 5).

The ProPneumo-1 assay detected 14 positive *C. pneumoniae* specimens compared to 11 in the monoplex "homebrew" assay, of which 1 specimen was false positive. Similarly the ProPneumo-1 detected 43 negative *C. pneumoniae* specimens compared to 47 in the monoplex "homebrew" assay, of which 2 specimens were false negative (Table 1 and Table 3). Prior to discordant analysis, the sensitivity and specificity of the ProPneumo-1 assay for *C. pneumoniae* was 91% and 96%, respectively (Table 4).

In the case of *M. pneumoniae*, ProPneumo-1 and culture detected 42 and 41 positive specimens, respectively. However, 4 specimens scored positive by the culture method (Table 1 and Table 2) and negative by the PCR method. Likely, due to repeated freeze-thaw cycles and the long storage period at -80 °C (up to 10 years), these 4 specimens are devoid of intact bacterial DNA and yielded neg-

Table 4: Specificity and sensitivity of the ProPneumo-I assay for C. pneumoniae.

Before discordant analysis				
C. pneumoniae	Monoplex "Home-Brew" Results			
ProPneumo-I Results		Positive	Negative	
	Positive	10	2	
	Negative	1	45	
	Total	10/11	45/47	
		Sensitivity = 91%	Specificity = 96%	
Following discordant analysis with Pneum	noBacter-I			
C. pneumoniae		Reference Me	thod Results	
ProPneumo-I Results		Positive	Negative	
	Positive	14	0	
	Negative	0	43	
		0	43	

Table 5: Specificity and sensitivity of the ProPneumo-I assay for M. pneumoniae,

Before discordant analysis					
M. pneumoniae	Culture Results				
ProPneumo-I Results		Positive	Negative		
	Positive	38	I		
	Negative	2	47		
	Total	38/40	47/48		
		Sensitivity = 95%	Specificity = 98%		
Following discordant analysis with Pneumo	Bacter-I				
M. pneumoniae	Reference Method Results				
ProPneumo-I Results		Positive	Negative		
	Positive	41	I		
	Negative	0	47		
	Total	41/41	47/48		
		Sensitivity = 100%	Specificity = 98%		

ative results in both PCR tests. Similarly, one specimen appeared false positive by the ProPneumo-1 assay. We suggest that this specimen is likely a true positive which was not detected by the PneumoBacter-1 assay, likely due to its gel-based format.

Prior to discordant analysis the sensitivity and specificity of the ProPneumo-1 assay for *M. pneumoniae* was 95% and 98%, respectively (Table 5). Following discordant analysis by the PneumoBacter-1 multiplex assay and classification of specimens as per the definitions described above, the sensitivity and specificity of the ProPneumo-1 assay for *C. pneumoniae* was 100% and 100%, respectively (Table 4). The sensitivity and specificity of the ProPneumo-1 assay for *M. pneumoniae*, however, was 100% and 98%, respectively (Table 5). Consistent with its multiplex format, the ProPneumo-1 assay detected 5 additional positive specimens, 2 *C. pneumoniae* in the *M. pneumoniae*-negative pool and 3 *M. pneumoniae* in the *C. pneumoniae*-negative pool.

Discussion

Early work in molecular diagnostics utilized technologies that were considered to be site-developed or "home-brew" in nature. Although these assays were considerable

advances at their time, they were often plagued by requirements for up-front research and development, quality assurance, and laboratory accreditation issues [10,11]. As a result, past molecular diagnostics for the clinical microbiology laboratory were often restricted to laboratories that had the monetary and human resources to meet these demands. In the last five years however, commercial products such as those identified in this manuscript have allowed for the implementation of molecular diagnostics without the up-front requirement for research and development as well as simplified quality assurance steps.

The ProPneumo-1 assay demonstrated several strengths as a commercial product for use in a wider range of clinical microbiology laboratory settings outside of the reference or highly resourced setting. First, this assay is a rapid, sensitive and specific method that allows for the diagnosis of *C. pneumoniae* and *M. pneumoniae* in a variety of clinical specimens. Second, the fact that the Pro-Pneumo-1 assay is a commercial product allows for simpler implementation in the diagnostic laboratory, and continued validation (e.g. quality assurance) when compared to "homebrew" assays [10,11]. Third, the ProPneumo-1 assay is a multiplex assay, which allows for multiple primer combinations in one reaction, instead of multiple monoplex

reactions. This not only makes the assay easier to perform in a laboratory but also has the potential of utilizing less patient specimen [17]. Indeed, using the ProPneumo-1 multiplex assay we detected 5 additional positive specimens, 2 *C. pneumoniae* in the *M. pneumoniae*-negative pool and 3 *M. pneumoniae* in the *C. pneumoniae*-negative pool.

There are several limitations to this study including its retrospective nature. A further limitation of this study is the relatively low number of *C. pneumonia*-positive specimens analyzed. However, the number of specimens analyzed represents the total complement of specimens available from this region over a period of ten years. The authors hope that further prospective analysis will allow for the further characterization of this commercial assay.

Several studies have evaluated the use of multiplex PCR for the detection of M. pneumoniae and C. pneumoniae in clinical specimens [18-20]. These studies have used a variety of assays, patient populations and sample types, thus proper validation and standardization are often lacking. These factors make it difficult to compare different studies although the finding has been generally consistent. A study by Khanna and colleagues [19] described the Pneumoplex assays (standard and real-time PCR) that can detect 5 pathogens, including M. pneumoniae and C. pneumoniae, in a single reaction. The diagnostic sensitivity of the Pneumoplex assay was 100% and was defined by spiking negative BAL specimens with bacterial DNA. The diagnostic specificity and sensitivity of the ProPneumo-1 assay, on the other hand, were determined in this study directly from patient's specimens. Another study by Ginevra and colleagues [20] described the triplex Chlamylege assay, for the detection of Legionella, M. pneumoniae and C. pneumoniae. The diagnostic specificity of the Chlamylege assay was also 100% and it was defined by sequencing the DNA of discordant specimens. Compared to our ProPneumo-1 study, fewer positive specimens were used for the evaluation of the Chlamylege assay (2 C. pneumoniae and 9 M. pneumoniae). However, the lack of an of appropriate "gold standard" for the quantitative analysis of intracellular pathogens makes comparisons of the sensitivities of these different assays difficult.

In conclusion, the ProPneumo-1 real-time multiplex PCR assay is a sensitive, specific, convenient and reliable tool for the detection of atypical respiratory infection pathogens directly from respiratory tract specimens. The authors believe that this product can be effectively implemented in a wide variety of clinical microbiology settings outside of reference and highly resourced laboratory settings.

Competing interests

The authors declare that they have no competing interests.

Authors' contributions

RRH undertook data analysis, assisted in validation study planning and drafted a manuscript. EL and PT participated in validation study planning and work. KR, AM, and SB undertook specimen collection and reference method analysis. FJ assisted in planning of experiments. SJD planned and coordinated study and drafted manuscript.

Acknowledgements

We thank Raymond Poon and Mary Morrison for help and technical exper-

References

- Stralin K, Tornqvist E, Kaltoft MS, Olcen P, Holmberg H: Etiologic diagnosis of adult bacterial pneumonia by culture and PCR applied to respiratory tract samples. J Clin Microbiol 2006, 44:643-645.
- Loens K, Ursi D, Goossens H, leven M: Molecular diagnosis of Mycoplasma pneumoniae respiratory tract infections. J Clin Microbiol 2003, 41:4915-4923.
- Waites KB, Talkington DF: Mycoplasma pneumoniae and its role as a human pathogen. Clin Microbiol Rev 2004, 17:697-728.
- Heath PT: Epidemiology and bacteriology of bacterial pneumonias. Paediatr Respir Rev 2000, 1:4-7.
- McIntosh K: Community-acquired pneumonia in children. N Engl J Med 2002, 346:429-437.
- Daxboeck F, Krause R, Wenisch C: Laboratory diagnosis of Mycoplasma pneumoniae infection. Clin Microbiol Infect 2003, 9:263-273.
- Loens K, leven M, Ursi D, Beck T, Overdijk M, Sillekens P, Goossens H: Detection of Mycoplasma pneumoniae by real-time nucleic acid sequence-based amplification. J Clin Microbiol 2003, 41:4448-4450
- 3. Winchell JM, Thurman KA, Mitchell SL, Thacker WL, Fields BS: Evaluation of three real-time PCR assays for detection of Mycoplasma pneumoniae in an outbreak investigation. J Clin Microbiol 2008, 46:3116-3118.
- Dumke R, Schurwanz N, Lenz M, Schuppler M, Luck C, Jacobs E: Sensitive detection of Mycoplasma pneumoniae in human respiratory tract samples by optimized real-time PCR approach. | Clin Microbiol 2007, 45:2726-2730.
- Gammie AJ: For the proposition: for the diagnosis of viral infections, commercial assays provide more reliable results than do in-house assays. Rev Med Virol 2008, 18:73-76.
- James V: Against the proposition: for the diagnosis of viral infections, commercial assays provide more reliable results than do in-house assays. Rev Med Virol 2008, 18:77-80.
- Tondella ML, Talkington DF, Holloway BP, Dowell SF, Cowley K, Soriano-Gabarro M, Elkind MS, Fields BS: Development and evaluation of real-time PCR-based fluorescence assays for detection of Chlamydia pneumoniae. J Clin Microbiol 2002, 40:575-583.
- Ursi D, Dirven K, Loens K, leven M, Goossens H: Detection of Mycoplasma pneumoniae in respiratory samples by real-time PCR using an inhibition control. J Microbiol Methods 2003, 55:149-153.
- Loens K, Ursi D, Goossens H, leven M: Evaluation of the Nucli-Sens miniMAG RNA extraction and real-time NASBA applications for the detection of Mycoplasma pneumoniae and Chlamydophila pneumoniae in throat swabs. J Microbiol Methods 2008. 72:217-219.
- Drews SJ, Blair J, Lombos E, DeLima C, Burton L, Mazzulli T, Low DE: Use of the Seeplex RV Detection kit for surveillance of respiratory viral outbreaks in Toronto, Ontario, Canada. Ann Clin Lab Sci 2008, 38:376-379.
- 16. Eshaghi A, Blair J, Burton L, Choi KW, De LC, Duncan C, Guyard C, Higgins R, Lombos E, Low DE, Mazzulli T, Drews SJ: Characterization of an influenza A and influenza B co-infection of a patient in a long-term care facility with co-circulating influenza A and influenza B. Int J Infect Dis 2008 in press.

- Brice G, Drews SJ, Low DE: Respiratory virus panels for global surveillance of emerging infectious diseases. J Clin Virol 2007, 40:S58-S60.
- Boman J, Gaydos CA, Quinn TC: Molecular diagnosis of Chlamydia pneumoniae by infection. J Clin Microbiol 1999, 37:3791-3799.
- Khanna M, Fan J, Pehler-Harrington K, Waters C, Douglass P, Stallock J, Kehl S, Henrickson KJ: The pneumoplex assays, a multiplex PCR-enzyme hybridization assay that allows simultaneous detection of five organisms, Mycoplasma pneumoniae, Chlamydia (Chlamydophila) pneumoniae, Legionella micadadei, and Bordetella pertussis, and its real-time counterpart. J Clin Microbiol 2005, 43:565-571.
- Ginevra C, Baranger C, Ros A, Mory O, Stephan J-L, Freymuth F, Joannes M, Pozzeto B, Grattard F: Development and evaluation of chlamylege, a new commercial test allowing simultaneous detection and identification of Legionella, Chalmydophila pneumoniae, and Mycoplasma pneumoniae in clinical respiratory specimens by multiplex PCR. J Clin Microbiol 2005, 43:3247-3254.

Publish with **Bio Med Central** and every scientist can read your work free of charge

"BioMed Central will be the most significant development for disseminating the results of biomedical research in our lifetime."

Sir Paul Nurse, Cancer Research UK

Your research papers will be:

- available free of charge to the entire biomedical community
- peer reviewed and published immediately upon acceptance
- cited in PubMed and archived on PubMed Central
- \bullet yours you keep the copyright

Submit your manuscript here: http://www.biomedcentral.com/info/publishing_adv.asp

