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Filling the gaps in clinical proteomics: a do-it-yourself guide for the identification of the emerging pathogen Arcobacter by matrix-assisted laser desorption ionization-time of flight mass spectrometry



Anne-Marie Van den Abeele^{a,*}, Dirk Vogelaers^b, Peter Vandamme^c, Elke Vanlaere^a, Kurt Houf^d

^a Microbiology Laboratory, Saint-Lucas Hospital, Groenebriel 1, 9000 Ghent, Belgium

^b Department of General Internal Medicine, Ghent University Hospital, De Pintelaan 185, Ghent, Belgium

^c Department of Biochemistry and Microbiology, Ghent University, Faculty of Sciences, Ghent, Belgium

^d Department of Veterinary Public Health and Food Safety, Faculty of Veterinary Medicine, Ghent University, Salisburylaan 133, 9820 Merelbeke, Belgium

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ABSTRACT

Arcobacters are considered emerging gastrointestinal pathogens. Rapid, reliable and species-specific identification of these bacteria is important. Biochemical tests commonly yield negative or variable results. Molecular methods prove more reliable but are time consuming and lack specificity. Matrix assisted laser desorption/ ionization-time of flight mass spectrometry (MALDI-TOF MS) is a fast, cheap and robust technique that has revolutionized genus and species identification in clinical microbiology.

The performance of an in vitro diagnostic (RUO) spectral database of MALDI-TOF MS for the identification of human clinically relevant Arcobacter isolates was validated and compared to an in house created Reference Spectral database (RS) containing a representative set of deposited Arcobacter strains of zoonotic interest. A challenge panel of clinical, human and veterinary, unique Campylobacteraceae strains was used to test accuracy.

Using direct colony transfer, sensitivity with RS was significantly better than with RUO for A. butzleri and A. cryaerophilus identification (100% and 92% versus 74% and 16%). For A. skirrowii, sensitivity remained low (21% versus 0%). Reanalysis using formic acid overlay (on-target extraction) augmented sensitivity for the latter species to 64%. Specificity of RS database remained excellent without any misidentifications of human clinical strains including Campylobacter fetus and C. jejuni/coli.

The use of an enriched database for MALDI-TOF MS identification of Arcobacter spp. of human interest produced high-confidence identifications to species level resulting in a significantly improved sensitivity with conservation of excellent specificity. Misidentifications, which can have therapeutic and public health consequences, were not encountered.

1. Introduction

Arcobacters are emerging bacterial zoonotic agents, causing predominantly gastrointestinal infections in humans. Initially, two of the human relevant species were taxonomically classified in the genus Campylobacter: Campylobacter cryaerophila (Neill et al., 1985) and Campylobacter butzleri (Kiehlbauch et al., 1991). In the latter year, both species were transferred into a new genus, Arcobacter, within the family Campylobacteraceae (Vandamme et al., 1991). Since then, 25 additional species and one Candidatus species have been identified and new species are pending (Figueras et al., 2017). Arcobacters inhabit a wide variety of ecological niches and have been isolated from various hosts and environments (Collado and Figueras, 2011). Some Arcobacter

species are commonly present in the digestive tract of healthy farm animals (Van Driessche et al., 2003). Although their prevalence varies geographically, the species most frequently reported are Arcobacter butzleri, A. cryaerophilus and A. skirrowii (On et al., 2002). They have been associated with enteritis and mastitis in animals, and may play a role in animal reproductive disorders, such as infertility and late term abortions in cattle, pigs and sheep (Ferreira et al., 2016). These same species are also reported in human infections throughout the world. An association of A. butzleri and A. cryaerophilus with enteritis, colitis and septicemia has been put forward in epidemiological studies, outbreaks and case reports, though A. cryaerophilus has also been recovered from stool of asymptomatic humans (Houf and Stephan, 2007). A. skirrowii has been implicated as causal agent of colitis in individual case reports

* Corresponding author. E-mail address: annemarie.vandenabeele@azstlucas.be (A.-M. Van den Abeele).

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Table 1

Main spectra present in the 3 spectral libraries: manufacturer-cleared IVD-RUO, in house RS- large and in house RS-small.

IVD-RUO Bruker reference library	RS-large in house reference Strains library	RS-small dereplicated in house reference Strains library
A. butzleri 347 98 NVU		
A. butzleri 460 98 NVU		
	A. butzleri LMG 15577	
A. butzleri CCUG 10373 A. butzleri DSM 8739 ^T	A. butzleri LMG 6620	
	A. butzleri LMG 10223	
	A. butzleri LMG 9906	A. butzleri LMG 9906
	A. butzleri LMG 9869	
	A. butzleri LMG 14714	A. butzleri LMG 14714
	A. butzleri LMG 10828 ^T	A. butzleri LMG 10828 ^T
	A. butzleri LMG 9910	
	A. butzleri LMG 9939	
	A. butzleri LMG 11120	
	A. butzleri LMG 10828 ^T	
	A. butzleri LMG 11632 A. butzleri LMG 10900	
A. cibarius DSM 17680 ^T	A. Dutzteri Livig 10900	
A. cryaerophilus QBR 3911 QBC		
A. cryaerophilus QBR 3998 QBC		
A. cryaerophilus CBR 3998 CBC A. cryaerophilus T277 CPB		
A. cryaerophilus V441 CPB		
A. cryaerophilus CCUG 1780 T NVU		
	A. cryaerophilus LMG 9947	A. cryaerophilus LMG 9947
	A. cryaerophilus LMG 10228	A. cryaerophilus LMG 10228
	A. cryaerophilus LMG 9865	A. cryaerophilus LMG 9865
	A. cryaerophilus LMG 9867	
	A. cryaerophilus LMG 10216	
	A. cryaerophilus LMG 10209	
A. cryaerophilus DSM 7289 ^T	A. cryaerophilus LMG 24291 ^T	A. cryaerophilus LMG 24291 ^T
	A. cryaerophilus LMG 10242	•
	A. cryaerophilus LMG 10244	
	A. cryaerophilus LMG 10221	
	A. cryaerophilus LMG 10224	A. cryaerophilus LMG 10224
	A. cryaerophilus LMG 10216	
	A. cryaerophilus	
	A. cryaerophilus LMG 10233	
	A. cryaerophilus LMG 10215	
	A. cryaerophilus	
	A. cryaerophilus LMG 10217	A. cryaerophilus LMG 10217
	A. cryaerophilus LMG 10222	
	A. cryaerophilus LMG 10218	
	A. cryaerophilus	
	A. cryaerophilus LMG 10212	
	A. cryaerophilus LMG 10213	
	A. cryaerophilus LMG 10244	A museum titus IMC 0060
	A. cryaerophilus LMG 9863	A. cryaerophilus LMG 9863
	A. cryaerophilus LMG 9948 A. cryaerophilus LMG 9065	
	A. cryaerophilus LMG 10210	
	A. cryaerophilus LMG 10210 A. cryaerophilus LMG 10211	
	A. cryaerophilus LMG 10228	
A. skirrowii DSM 7302 ^T	A. skirrowii LMG 6621^{T}	
A. skirrowii ATCC 51132 ^T	A. skirrowii LMG 6621^{T}	A. skirrowii LMG 6621^{T}
AL SKOWE MICE 51152	A. skirrowii LMG 9801	A. skirtowit LMG 9801
	A. skirrowii LMG 14985	A. skirowii LMG 14985
		A. skirrowii LMG 9912 ^a
		A. skirrowii LMG 10234 ^a
Total number of strains in spectral databases		
13	46	15

Data represent MSPs of strains from the Bruker Reference Library 3.2.1.0 (n = 13), MSPs of strains from the home made spectral database RS-large (n = 46) and MSPs of the dereplicated home made spectral database RS-small (n = 15), ^a two strains added after dereplication from the large in house reference library, species Type strains are designated by superscript T. The name of the MSP consists of the abbreviation of the culture collection and strain number. For the in house libraries, RS large and small, LMG reference numbers are used throughout the list. Corresponding reference numbers from other culture collections can be retrieved from www. straininfo.net.

CCUG, Culture Collection of the University of Göteborg;

ATCC, American Type Culture Collection;

DSM, Deutsche Sammlung von Mikroorganismen und Zellkulturen;

(Wybo et al., 2001). *Arcobacter* is presumably transmitted to humans by contaminated food and water, with poultry and pork products identified as important reservoirs (Collado and Figueras, 2011). Two extensive studies identified *Arcobacter* as the fourth most common *Campylobacter*-like organism in stool of patients with gastrointestinal

disease (Vandenberg et al., 2004, Prouzet-Mauléon et al., 2006) and one study ranked *Arcobacter* fourth as bacterial pathogen causing diarrheal illness (Van den Abeele et al., 2014). The three *Arcobacter* species of human relevance are Gram-negative, slender, spirally curved rods, able to grow in air and at temperatures ranging from 25 up to Download English Version:

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