



Identification of Pasteurellaceae by Matrix-Assisted Laser Desorption Ionization-Time of Flight Mass Spectrometry

S. Zimmermann^{1*}, E. Gillmann¹, R. Mutters² and I. Burckhardt¹

¹ Department of Infectious Diseases, Microbiology and Hygiene, University of Heidelberg, Germany, ²Institute of Medical Microbiology and Hygiene, Philipps University Marburg, Germany

*corresponding author: stefan.zimmermann@med.uni-heidelberg.de

Objectives

The taxonomic systematic of Pasteurellaceae is a difficult topic. Some species has been reclassified from the genus Haemophilus to Aggregatibacter (e.g. Aggregatibacter aphrophilus and paraaphrophilus). Other species left the genus Pasteurella and new genus were defined (e.g. Avibacterium or Gallibacterium). In daily routine in the microbiology lab many of them are difficult to identify in commercially available biochemical systems using parameters. Matrix-assisted laser desorption ionization-time of flight (MALDI-TOF) mass spectrometry is a rapid and accurate tool for the identification of many microorganisms. We assessed this technology for more than 30 Pasteurellaceae species from the Medical Culture Collection Marburg (MCCM). In addition we determined the presence of ßlactamases in the strains, an important marker for effective treatment, recommendations,

Materials and Methods

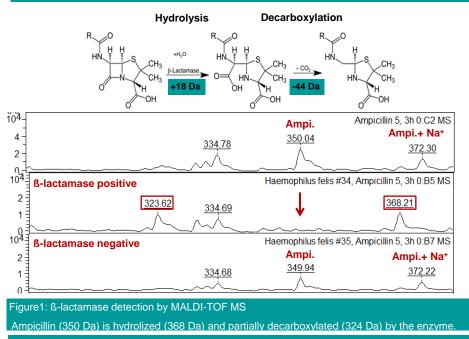
Pasteurellaceae strains were grown on blood chocolate under aerobic or or agar microaerophilic conditions between 24 to 72 hours. Identification with mass spectrometry was done by smear preparations (direct inoculation of the target from the plate) and after a protein extraction protocol. The results were compared to the MCCM database and all identifications were confirmed by 16SrDNA sequencing. In addition the presence of ß-lactamases was determined by MALDI-TOF using a modification of our recently described protocol for carbapenemase detection (Burckhardt et al., JCM 2011).

Results							
No	Strain	мссм	Sco	MS result (db-2010)		DNA Sequencing	MS result 2"(db-2012)
2	Pasteurella stomatis	0104	1.7	Pasteurella multocida (Genus)	ndb	Pasteurella stomatis	Pasteurella stornatis
3	Pasteurella pneumotropica	3303	1.8	Haemophilus influenzae (Genus)	ndb	Pasteurella pneumotropica	Pasteurella pneumotropica
4	Pasteurella pneumotropica	3409	1.8	Haemophilus influenzae	ndb	Pasteurella pneumotropica	Pasteurella pneumotropica
5	Pasteurella multocida septica	0378	2.3	Pasteurella multocida		Pasteurella multocida	Pasteurella multocida
6	Pasteurella multocida	3024	2.2	Pasteurella multocida		Pasteurella multocida	Pasteurella multocida
7	Pasteurella multocida gallicida	3280	2.4	Pasteurella multocida		Pasteurella multocida	Pasteurella multocida
9	Avibacterium gallinarum	0033	1.8	Avibacterium gallinarum		Avibacterium gallinarum	Avibacterium gallinarum
10	Avibacterium gallinarum	0003	1.5	Avibacterium gallinarum		Avibacterium gallinarum	Avibacterium gallinarum
11	Pasteurella dagmatis	3139	2.0	Pasteurella multocida (Genus)	ndb	Pasteurella pneumotropica	
12	Pasteurella dagmatis	3191	1.6	Pasteurella multocida	ndb	Pasteurella dagmatis	Pasteurella dagmatis
13	Pasteurella canis	0045 ^T	1.5	Pasteurella multocida	ndb	Pasteurella canis	Pasteurella canis
14	Pasteurella canis	3430	1.3	Pasteurella multocida	ndb	Pasteurella canis	Pasteurella canis
15	Pasteurella bettyae	0450 ⁺	1.6	Pasteurella multocida (Genus)	ndb	Pasteurella bettyae	Pasteurella bettyae
17	Pasteurella avium	0272 [™]	2.3	Pasteurella avium		Pasteurella avium	Pasteurella avium
18	Pasteurella anatis	0140 ⁷	2.1	Pasteurella anatis		Pasteurella anatis	Pasteurella anatis
19	Pasteurella anatis	0141	2.2	Pasteurella anatis		Pasteurella anatis	Pasteurella anatis
22	Mannheimia varigena	2528	2.0	Mannheimia haemolytica		M. glucosida/M. haemolytica	
23	Mannheimia glucosida	0575	2.1	Mannheimia haemolytica	ndb	Mannheimia glucosida	Mannheimia glucosida
24	Mannheimia glucosida	0388	1.8	Mannheimia haemolytica	ndb	Mannheimia glucosida	Mannheimia glucosida
25	Histophilus ovis	0338	2.0	Histophilus somni		Histophilus somni	
26	Histophilus ovis	1381	1.5	Mannheimia glucosida		Mannheimia glucosida	
27	Haemophilus parainfluenzae	3215	2.3	Haemophilus parainfluenzae		Haemophilus parainfluenzae	Haemophilus parainfluenzae
28	Haemophilus parainfluenzae	2641	2.4	Haemophilus parainfluenzae		Haemophilus parainfluenzae	Haemophilus parainfluenzae
	Haemophilus parahaemolyticus	3212		Haemophilus parainfluenzae		Haemophilus parainfluenzae	
	Haemophilus influenzae	0617		Haemophilus influenzae		Haemophilus influenzae	Haemophilus influenzae
	Haemophilus haemolyticus	0352		Haemophilus influenzae	ndb	Haemophilus sp.	Haemophilus haemolyticus
32	Haemophilus haemolyticus	2169	2.0	Haemophilus influenzae	ndb	Haemophilus haemolyticus	Haemophilus haemolyticus
33	Haemophilus haemoglobinophilus	0723 [™]	1.5	Haemophilus influenzae	ndb	Haemophilus haemoglobinophilus	Haemophilus haemoglobinophilus
34	Haemophilus felis	2738	1.5	Pasteurella multocida	ndb	Haemophilus felis	Haemophilus felis
35	Haemophilus felis	3143	1.8	Pasteurella multocida	ndb	Haemophilus felis	Haemophilus felis
	Actinobacillus muris	821	1.3	Haemophilus influenzae	ndb	Actinobacillus muris	Actinobacillus muris
	Actinobacillus lignieresii	2712	2.1	Aggregatibacter aphrophilus		Aggregatibacter aphrophilus	
43	Aggregatibacter actinomycetemcomitans	0135	2.3	Aggregatibacter actinomycetemc.		Aggregatibacter actinomycetemc.	Aggregatibacter actinomycetemc.

 Table1: Identification of different Pasteurellaceae strains by MALI-TOF MS

 27 of 33 strains were identified on species level (4 on genus level, 2 unsolved).

 IMCCM: Medical Culture Collection Marburg; ndb: not in database]



Conclusions

1. MALDI-TOF is a fast and reliable method to identify Pasteurellaceae in daily lab routine.

2. The presence of ß-lactamases can be investigated on the mass spectrometer.