Milk Bacteriological Analysis Using MALDI-TOF Technology

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Many veterinary and milk quality laboratories are now using MALDI-TOF MS (Matrix Assisted Laser Desorption/Ionization-Time of Flight Mass Spectrometry) for identification of bacteria from milk samples. The aim of this article is to provide information on this new technology and on current practices being followed by diagnostic laboratories for reporting results generated using MALDI-TOF.

What is MALDI-TOF?

MALDI-TOF uses laser excitation of biomolecules, such as proteins and mass spectrometry, to create a unique protein spectrum pattern of an unknown microorganism. Results are compared with thousands of spectra in a reference database, from which a bacterial identification is provided. The procedure is rapid, accurate, and frugal in its use of chemicals and human resources required for testing, while providing species identification for a majority of samples tested (1).

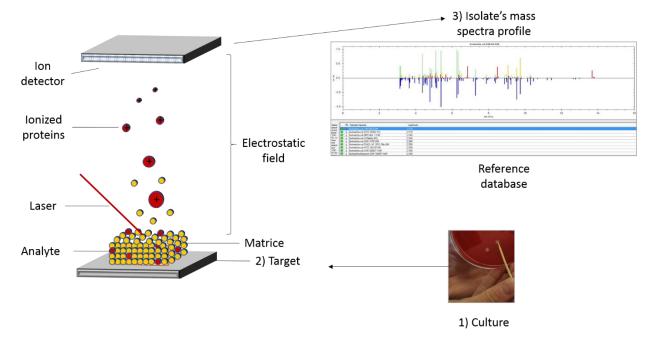


Figure 1. Principles of MALDI-TOF analysis. 1) For bacterial species that can be grown on culture media, an isolated colony is picked and dropped on a metal target. 2) the colony (in red) is covered with a matrice (in yellow). A pulsed laser is then used to ionize the isolate's molecules. The time of flight of the various molecules in an electrostatic field is then captured by an ion detector to generate a mass spectra profile for that isolate. 3) The observed profile is then compared to those of a reference database with spectra from thousands of known bacteria and a score describing confidence in the genus- or species-level identification is reported.

Why use MALDI-TOF instead of traditional microbiological identification methods?

Like traditional microbiological methods, a 24- to 48-h culture is needed before proceeding to MALDI-TOF identification of a given isolate. Therefore, organisms that do not grow on common media cannot be identified using MALDI-TOF. When bacterial colonies can be cultured, however, they can then be identified to genus (e.g. *Staphylococci*) and species level (e.g. *aureus*), in one single and rapid (<60 sec) analysis using MALDI-TOF. In comparison, traditional microbiological methods often require multiple biochemical tests, re-incubation, and an additional day before a final identification can be given (2). So, an important advantage of MALDI-TOF technology is its promptness for obtaining an identification.

The most important advantage of MALDI-TOF technology, however, is that it can provide bacterial identification information more specific than traditionally reported for organisms isolated from milk. For instance, in the past, several expensive tests (e.g. test kits for bacterial identification or gene sequencing) and additional days were needed to identify to the species-level members of closely related bacterial groups, such as non-aureus Staphylococci (NAS, also known as coagulase negative Staphylococci or CNS), and these tests were generally not conducted on a routine basis. MALDI-TOF technology allows a greater number of bacterial isolates to be rapidly and accurately identified to species level. Therefore, many groups of bacteria, such as those traditionally identified as NAS, will now be identified using MALDI-TOF by genus and species (e.g. Staphylococcus chromogenes, Staphylococcus simulans) (3, 4). Though it might be disturbing for a practitioner to receive for the first time a milk culture result reporting growth of an uncommon bacterial species, it is important to remember that these species "appeared" uncommon because they were not identified to species level with traditional microbiological culture. Presence of these pathogen species in milk is not novel; it is the fact that we can now identify them that is novel. If you see unfamiliar bacterial names on your mastitis reports, the diagnostic laboratory can provide further information on these.

A list of many of the bacterial species that may be isolated from milk samples using MALDI-TOF identification is presented in the Appendix. They are grouped under their common classification for mastitis management purposes. Mastitis researchers are currently applying bacterial identification information generated using MALDI-TOF and other advanced technologies to explore the role of specific bacteria present in milk and to understand their epidemiology. This may lead to valuable information that will be used to implement new prevention and control strategies, and refine mastitis treatment protocols in the future. Meanwhile, current knowledge on the group of organisms they belong to should be used for management of intramammary infections caused by these organisms.

How does MALDI-TOF testing compare with PCR testing?

Like traditional bacteriological methods, MALDI-TOF absolutely requires the culture of an organism on culture media (see Figure 1). Therefore, many of the same advantages and disadvantages reported for routine milk analysis would also apply to milk culture, followed by MALDI-TOF identification (2). Readers are encouraged to consult the NMC PCR (polymerase chain reaction) Fact Sheet for an exhaustive description of the advantages and disadvantages of both culture and PCR-based methods. Briefly, by using PCR, milk can be tested directly without culture, which can reduce the time to results by 24 h, compared with culture followed by MALDI-TOF identification. A PCR test, however, will only be able to test for the presence of the specific organisms targeted ahead of time in that test, whereas, with MALDI-TOF, every organism species retrieved on the culture media can potentially be identified.

How confident can we be with MALDI-TOF results?

To ensure that MALDI-TOF identification results are consistent among laboratories, standard procedures are used by all labs. Every laboratory using MALDI-TOF identification can provide information about the

MALDI equipment manufacturer, specific machine, software version, and, most importantly, the use of any custom library for identification and the minimal score required to accept identification at the genus and species levels. Indeed, laboratories have the opportunity to add main spectra in custom databases using well-characterized isolates, which may allow for identification of additional species or improve accuracy for already identifiable species (4). Mentioning use of such custom databases is, therefore, of importance for interpretation of MALDI-TOF results. In 2018, the manufacturers' libraries contained protein profiles for up to 7,311 isolates from 2,500 microbial species (library contents differ by MALDI-TOF used). Like many microbiological identification systems, most of the well-characterized isolates used to generate the reference profiles found in the manufacturer's database are human isolates. For many microbial species, protein profiles of animal isolates will be quite similar to those of human isolates and will be correctly identified. It remains to be determined whether this is true for all species (1).

Conclusions

In summary, MALDI-TOF Mass Spectrometry is now widely used in veterinary microbiology laboratories. It has been proven to provide consistent, accurate, and rapid identification of bacteria, fungi, and mycobacteria. One powerful feature of this technology is the ability to expand and share databases. As information is added to these databases, more bacteria will be identified to the species level in laboratories using this technology and the updated databases.

References

- 1. Kostrzewa, M., and S. Schubert. MALDI-TOF mass spectrometry in microbiology. Caister Academic Press N, UK, 2016.
- 2. Middleton, J.R., L.K. Fox, and G. Pighetti. 2017. Laboratory Handbook on Bovine Mastitis. National Mastitis Council. New Prague, MN.
- 3. Cameron, M., H.W. Barkema, J. De Buck, S. De Vliegher, M. Chaffer, J. Lewis, and G.P. Keefe. Identification of bovine-associated coagulase-negative staphylococci by matrix-assisted laser desorption/ionization time-of-flight mass spectrometry using a direct transfer protocol. J. Dairy Sci. 2017;100(3):2137-47.
- 4. Cameron, M., J. Perry, J.R. Middleton, M. Chaffer, J. Lewis, and G.P. Keefe. Short communication: Evaluation of MALDI-TOF mass spectrometry and a custom reference spectra expanded database for the identification of bovine-associated coagulase-negative staphylococci. J Dairy Sci. 2018;101(1):590-5.

Appendix

Bacterial species that may be identified from milk samples using MALDI-TOF

1) Main Contagious Mastitis Pathogens

The source of infection is mainly the udder from infected animals. Intramammary infections are acquired primarily during milking.

Corynebacterium bovis Mycoplasma bovis Mycoplasma spp.

Staphylococcus aureus Streptococcus agalactiae

2) Main Environmental Mastitis Pathogens

Environmental bacteria are found in the cow's environment. Intramammary infection occurs when bacteria from dirt or manure at the teat end hinder the defense mechanisms of the udder.

Gram positive (environmental Streptococci and Streptococci-like organisms):

Aerococcus spp. Enterococcus hirae Streptococcus canis^a

Aerococcus viridans Enterococcus saccharolyticus Streptococcus dysgalactiae

Enterococcus spp.Lactococcus garvieaeStreptococcus equiEnterococcus casseliflavusLactococcus lactisStreptococcus uberisaEnterococcus faecalisStreptococcus spp.Streptococcus vestibularisEnterococcus faeciumStreptococcus bovis/Streptococcus equinus complexb

Gram negative (coliforms):

Citrobacter sp. Hafnia sp. Pantoea sp.

Enterobacter sp. Klebsiella sp. Pantoea agglomerans

Enterobacter cloacae Klebsiella oxytoca Escherichia coli Klebsiella pneumoniae

3) Other Common Mastitis Pathogens

Non-aureus Staphylococci are the most prevalent bacteria isolated from milk in herds using recommended mastitis control measures. Some NAS species appear to be opportunistic and infect the teat canal and mammary gland from skin sources, or from the environment. Some species, however, can possibly be transmitted from cow to cow.

Non-aureus Staphylococci (NAS):

Staphylococcus agnetis/hyicus²Staphylococcus capitisStaphylococcus chromogenesStaphylococcus cohniiStaphylococcus devrieseiStaphylococcus epidermidisStaphylococcus equorumStaphylococcus fleurettiStaphylococcus gallinarumStaphylococcus haemolyticusStaphylococcus saprophyticusStaphylococcus sciuriStaphylococcus simulansStaphylococcus succinusStaphylococcus warneri

Staphylococcus xylosus other Staphylococcus species^d

^a Certain Streptococcus uberis and Streptococcus canis strains may act as contagious pathogens.

^b *S. bovis/S. equinus* complex includes several Streptococcus species. Your diagnostic laboratory can provide further information.

^c MALDI-TOF cannot differentiate *S. agnetis* from *S. hyicus*.

^d Additional *Staphylococcus* identified by MALDI are likely to be reported to species level as the MALDI-TOF database expands.

4) Uncommon Mastitis Pathogens

Numerous other bacteria can cause intramammary infections (IMI). While most of these IMI are usually sporadic, some may cause severe herd problems.

Other Gram positive:

Trueperella pyogenes Nocardia spp.

Bacillus spp. Listeria monocytogenes

Other Gram negative:

Acinetobacter spp. Proteus spp. Serratia spp.

Acinetobacter baumanii Proteus mirabilis Serratia liquefacians Pasteurella spp. Proteus vulgaris Serratia marcescens

Pasteurella multocida Salmonella spp. Yersinia spp.

Considered as or similar to *Pseudomonas*:

Aeromonas spp. Plesiomonas shigelloides Pseudomonas fulva Flavimonas spp. Pseudomonas spp. Stentrophomonas spp.

Other Non-Bacteria:

Yeast Prototheca spp.