



Etest



The principle behind the VITEK® MS for identification of microorganisms is more than 20 years old. Working directly with AnagnosTec and Shimadzu, two pioneers in the field of bacterial identification using mass spectrometry, bioMérieux offers VITEK® MS with the same high standards you expect

- 1988: first commercially available MALDI-TOF system from Shimadzu
- 1998: AnagnosTec develops the SARAMIS™ Database
- 2000: European patent for the SARAMIS™ Database
- 2002: Koichi Tanaka (Shimadzu) wins the Nobel Prize for Soft Laser Desorption*
- * Desorption of large molecules that results in ionization without the formation of fragment ions.

SPECIFICATIONS

Dimension

VITEK® 2

- Size (w h d) 0.7 m x 1.92 m x 0.85 m minimum distance to wall at back is 100 mm
- Weight 330 kg excluding data system

VITEK® MS

Installation Requirements

- Electrical 200 VAC, 50/60 Hz, 1000 VA single phase OR 230 VAC, 50/60 Hz, 1000 VA single phase
- A "clean", stable and continuous mains supply is required for reliable operation
- Temperature ambient 18° to 26° Celsius
- Relative humidity less than 70% non condensing
- Vibration free, firm, level floor, at least 330 kg supported at four points

Laser

- 337 nm nitrogen laser, fixed focus
- 3 ns pulse rate 50Hz (50 laser shots per second)
- Near normal (on-axis) incidence of the laser beam to the sample
 Laser power and laser aim under software control

Analyzer

- Linear flight tube of 1.2 m drift length
- Vacuum maintained by two turbomolecular pumps (nominal 250 l/s) with rotary backing
- Beam blanking to deflect unwanted high intensity signals e.g. matrix ions

Mass range

• 1 to 500 kDa

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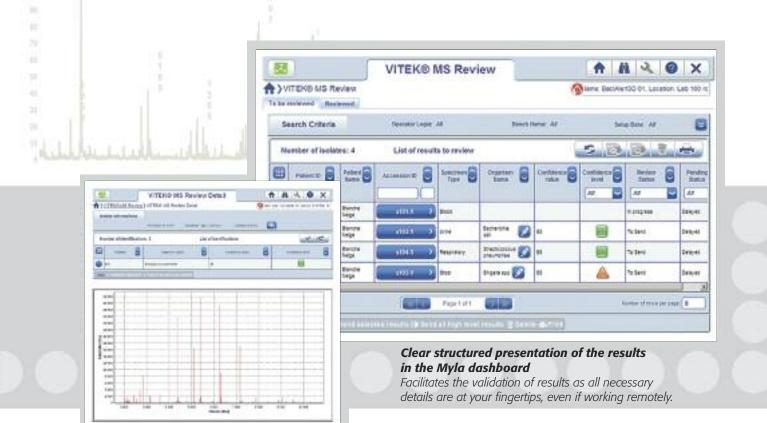


IDENTIFICATION & SUSCEPTIBILITY TESTING VITEK MS **Fast Flexible Innovative** VITEK 2 BIOMÉRIEUX



VITEK MS[™] The best and most complete **ID/AST solution**





Fast Identification, Flexibility and Innovative solution

A clinically RELEVANT database

Microbial identification is achieved by obtaining spectra using MALDI-TOF technology (Matrix Assisted Laser Desorption Ionization Time-of-Flight) and analyzing the spectra with the VITEK® MS database.

- Comprised of clinically relevant species with more than 25 000 spectra. Robust validation using an Advanced Spectra Classifier for reliable identification
- A large number of strains tested for each species in the database



Traceability and Flexibility

VITEK® MS includes the VITEK® MS Prep Station to securely link specimen information with each spot on the target slide and to the VITEK® 2 cassette position

- Traceability: Connect multiple VITEK® MS PREP stations facilitating traceability of all ID/ASTs
- Flexibility: Disposable target slides with unique barcodes eliminate manual data entry and workflow is enhanced on independent workstations
- Generation of electronic worksheets during set up

The VITEK® MS Advantage

- Full Integration with ID from VITEK® MS to AST from VITEK® 2 by a single provider. A connection made by us and managed by us!
- Optimized sample loading: Simply deposit the organisms onto the target slide, add matrix and run the mass spec.
- On target extraction: Protein extraction, if needed can be performed directly on the target slide.
- Ready-to-use consumables: The VITEK® MS comes with ready-to-use, light stable matrix solution saving time on reagent preparation.
- Efficiency: Up to four target slides with 48 positions each can be analyzed in parallel in the system allowing testing of 192 isolates in one run.
- Confidence: Disposable slides eliminate the need for cleaning and potential sample contamination.
- High resolution: Highly reproducible identification from microorganism protein mass spectra in the >10k Dalton range. Ability to scan up to 500k Dalton enabling the possibility for future applications.
- Convenience: With Myla™ you can easily access results and system information through a networked PC.

All information available at your fingertips, when you need it and where you need it!