

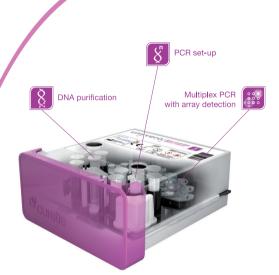
unyvero

Unyvero's sample-to-answer platform provides rapid results for severe infectious diseases in hospitalized patients

Powerful multiplex PCR technology combined with the broadest range of microorganism and resistance targets sets the Unyvero System apart.

The Unyvero System consists of:

- Lysator to lyse and process a variety of native samples
- Cockpit to manage testing process, display, store, and transmit results
- Analyzer to perform DNA testing with random-access, multiplex PCR



A single test handles one patient sample, analyzes over 100 DNA analytes and delivers reliable results within just 4-5 hours





Unyvero is designed to expand with your growing needs

Applications for severe infections:

- Blood Culture BCU
- Hospitalized Pneumonia HPN
- Intra-Abdominal Infection IAI
- Implant & Tissue Infection ITI
- Urinary Tract Infection UTI

The Unyvero System is distribuited on an exclusive basis by A.Menarini Diagnostics in the following countries: Benelux, France, Germany, Greece, Italy, Portugal, Spain, United Kingdom.

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Fast & Simple Syndromic Testing for Severe Infections - Improving Patient Outcomes





In industrialized countries, sepsis is responsible for as many deaths as heart attacks1

Sepsis is a major health issue and its recorded incidence is rising every year.2

Mortality rates can be as high as 50%.

Direct costs per sepsis patient range from € 23,000 to € 29,000 in Europe.

A microbiological analysis is recommended to narrow the empiric antimicrobial therapy as quickly as possible.

> Worldwide, someone dies of sepsis every 3-4 seconds.4

Faster detection enables earlier optimization of therapy

fungi and antibiotic resistance genes.

- Pathogen identification can take days using routine microbiology methods
- Every hour effective antibiotic treatment is delayed, sepsis mortality rate increases up to 8%
- Early identification can help reduce morbidity and mortality, improve patient care and reduce healthcare costs

critical determinant of survival in human septic shock. Crit Care Med. 2006; 34(6):1589-96

- The Unyvero BCU Application simultaneously identifies a large panel of bacteria,

7 Kumar A et al. Duration ot hypotension before

Institut Pasteur, Disease Sheet -Sepsis, 2014.

- 2 Hotchkiss RS et al., Sepsis and septic shock. Nat Rev Dis Primers. 2016; 2:16045.
- 3 Fleischmann C et al., Assessment of Global Incidence and Mortality of Hospital-treated Sepsis. Current Estimates and Limitations. Am J Respir Crit Care Med. 2016; 193(3):259-72.
- 4 WSD Fact Sheet 2018 (www.world-sepsis-day.org).
- ⁵ Burchardi H and Schneider H. Economic aspects of severe sepsis: a review of intensive care unit costs, cost of illness and cost effectiveness of therapy. Pharmacoeconomics. 2004; 22(12):793-813.
- 6 Rhodes A. et al., Surviving Sepsis Campaign: International Guidelines for Management of Sepsis and Septic Shock: 2016. Intensive Care Med. 2017; 43:304–377.

Sample Types

Positively flagged blood cultures.



Easy Workflow



Multiple Sample Types



Results

Unyvero Blood Culture (BCU) Cartridge

bacteria	Enterobacteriaceae	bacteria	Fungi	Resistance	Gene
Staphylococcus aureus Coagulase negative staphylococci Streptococcus spp. Streptococcus agalactiae Streptococcus pneumoniae Streptococcus pyogenes/ dysgalactiae Enterococcus spp. Enterococcus faecalis Listeria monocytogenes	Citrobacter freundii/koseri Escherichia coli Enterobacter cloacae complex Klebsiella aerogenes (E. aerogenes) Klebsiella oxytoca Klebsiella pneumoniae Klebsiella variicola Proteus spp. Serratia marcescens	Acinetobacter baumannii complex Pseudomonas aeruginosa Stenotrophomonas maltophilia Anaerobic bacteria Cutibacterium acnes (P. acnes)	Aspergillus spp. Candida spp. Candida albicans Candida dubliniensis Candida glabrata I. orientalis (C. krusei) Candida parapsilosis Candida tropicalis	Aminoglycoside Macrolide/ Lincosamide Oxacillin Vancomycin 3rd generation Cephalosporins	ermA mecA mecC vanA vanB ctx-M
Corynebacteriaceae	Universal bacteria	Other Gram-negative bacteria	Mycobacteriaceae	Carbapenem	kpc imp ndm oxa-23 oxa-24/40
Corynebacterium spp.	Detection of prokaryotic genetic sequence	Haemophilus influenzae Neisseria meningitidis	Mycobacterium spp.		oxa-48 oxa-58 vim

Clinical evidence demonstrates the benefits provided by the Unyvero solution

Study ¹

Multicenter performance evaluation. Clinical laboratories from HDZ Bad Oeynhausen, UKE Hamburg and OWS Vienna

Number of samples 178 positive blood cultures.

96.8% Sensitivity

99.8% Specificity

Burrack-Lange et al., Multicenter assessment of the rapid Unyvero Blood Culture molecular assay J. Med. Microbiology 2018; 67(9):1294-1301 AST: Antibiotic Susceptibility Testing

Value of resistance genes

- 119 resistance genes detected during study period.
- Important information for earlier targeted therapy and infection control.

Rapid identification is critical for survival

Using Unyvero, the average time to results was:

- Reduced by 11h compared to identification results.
- Reduced by 34h compared to full AST results.

Polymicrobial infections are

- 6/7 polymicrobial infections, with pathogens included in the panel, correctly identified.
- 5 samples with an additional micro-organism were detected using Unyvero.





Study 2

Comparison with routine microbiology. University Hospital Essen, Germany.

100%

Agreement

Number of samples 50 positive blood cultures. 7 spiked blood cultures.

Study population

46 patients.

Age 1 month-45 years.

for resistance markers

Unyvero turnaround time 4-5 hours

95% Agreement

with ID results

Spiked cultures

- 7/7 Spiked bacteria correctly identified.
- 3/3 Resistance markers correctly identified (mecC, vanA and vanB).







Conclusion

The Unyvero BCU Application is a useful tool for the rapid detection of pathogens and resistance markers directly from positive blood cultures.

Schmidt et al., (2018) poster presentation DGHM.