

# Comparative evaluation of Accelerate Pheno™ and Vitek® 2 systems for rapid antibiotic susceptibility testing of pathogens in positive blood cultures

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## Introduction

Bloodstream infection (BSI) remains a major public health concern because of its high incidence and consequences in terms of mortality, morbidity and cost — particularly in the case of nosocomial infection. While timely administration of effective antimicrobial therapy may reduce hospital length of stay and mortality of patients with BSI, a delayed (and potentially less effective) treatment often results in more severe stages of BSI-related disease.

Antimicrobial susceptibility testing (AST) of organisms causing BSI is an undisputed prerequisite for optimal antimicrobial therapy. In contrast to conventional automated methods, e.g. the widely used VITEK® 2 system (bioMérieux, Marcy-l’Étoile, France), recent efforts led to developing new generation methods for AST. Among them, the Accelerate Pheno™ system (Accelerate Diagnostics, Tucson, AZ, USA) is an automated microscopy platform that uses fluorescence *in situ* hybridization for identification and morphokinetic cellular analysis to provide AST results directly from positive blood cultures (PBCs).

## Objective

The objective of this study was to compare the performance of the Accelerate Pheno™ system (Accelerate Diagnostics, Inc.) with that of the conventional phenotypic VITEK® 2 system for rapid antimicrobial susceptibility testing of bacterial pathogens from positive blood culture samples, based on the reference broth microdilution (BMD) method.

## Materials and Methods

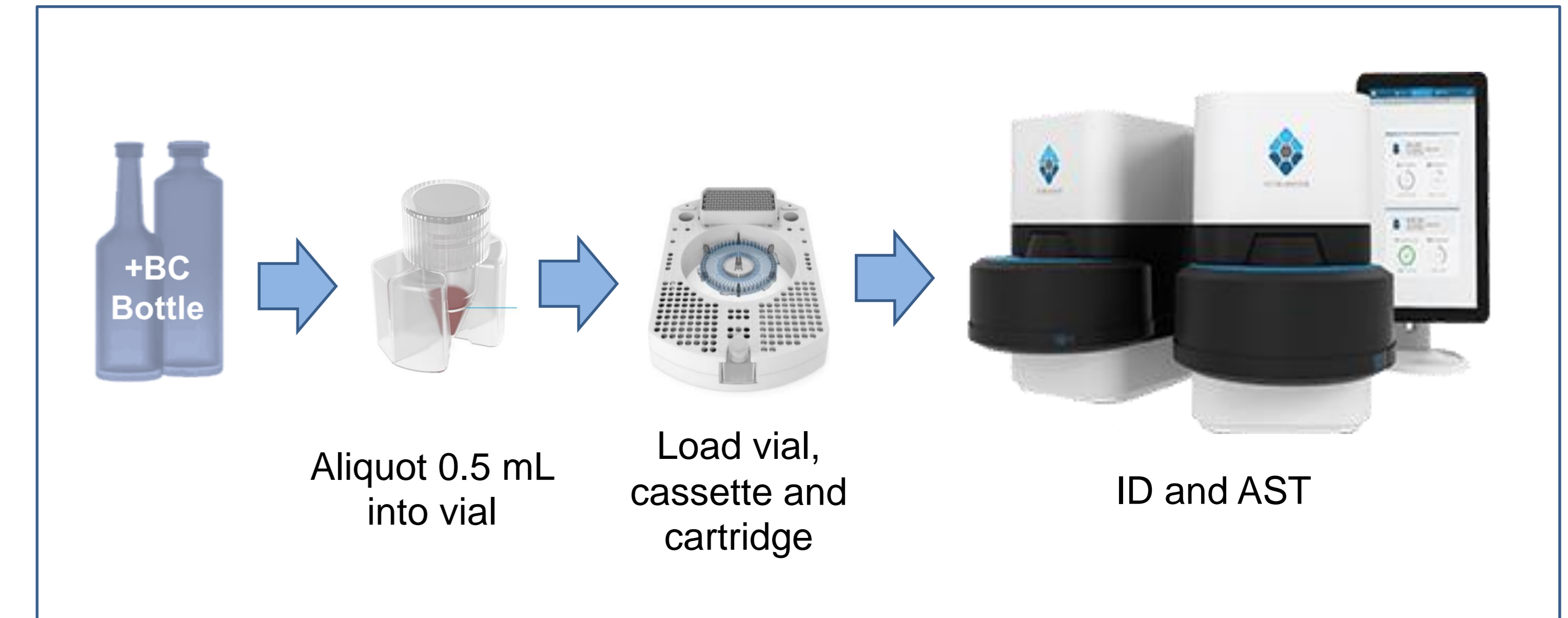
Prospectively collected blood cultures that represented single bloodstream infection patient/episodes over a 6-month period in 2017 were included. Blood cultures were collected in BacT/ALERT® FA and FN PLUS bottles (bioMérieux, Marcy-l’Étoile, France) and incubated until growth detection using the BacT/ALERT® VIRTUO® system (bioMérieux). For PBC samples showing monomicrobial growth (*n* = 86), AST was performed directly using both Accelerate Pheno™ and VITEK® 2 systems with PBC broth. Thereafter, colony isolates derived from subculture of PBC broth were used for the BMD method. AST results were interpreted according to 2017 EUCAST breakpoints.

**Table 1:** Performance of the Accelerate Pheno™ and VITEK® 2 systems compared to BMD for Gram-negative bacterial species (*n* = 62)

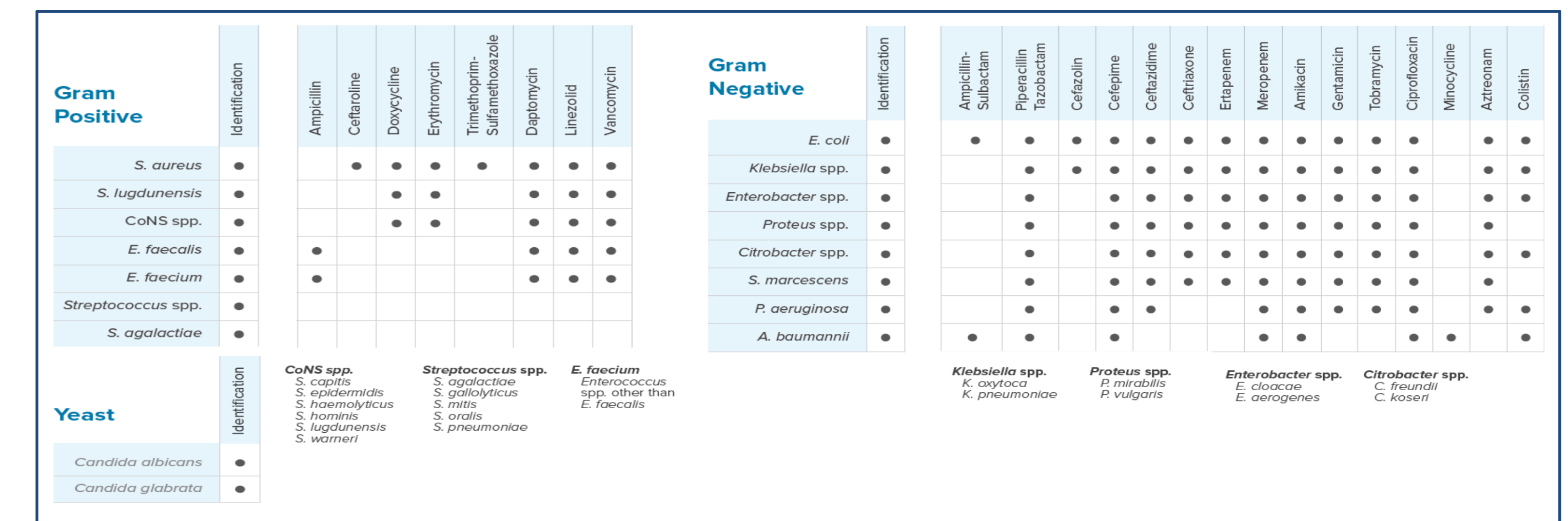
Antibiotic	Accelerate Pheno™ system							VITEK® 2 system								
	CA	VME	ME	mE	S	I	R	CA	VME	ME	mE	S	I	R		
<b>Enterobacteriaceae (n=46)</b>																
Amikacin	40/46	87.0%	0	0	6	40	5	1	37/46	80.4%	0	2	7	40	5	1
Cefepime	40/45	88.9%	0	0	5	28	2	15	39/46	84.8%	0	0	6	28	2	16
Ceftazidime	43/46	93.5%	0	0	3	24	4	18	42/46	91.3%	0	0	4	24	4	18
Ciprofloxacin	44/46	95.7%	1	0	1	25	1	20	45/46	97.8%	0	0	1	25	1	20
Colistin	39/41	95.1%	0	2	0	39	0	2	40/41	97.6%	1	0	0	39	0	2
Ertapenem	45/46	97.8%	1	0	0	36	1	9	45/46	97.8%	0	0	1	36	1	9
Gentamicin	45/46	97.8%	0	0	1	37	0	9	40/46	87.0%	0	0	6	37	0	9
Meropenem	44/46	95.7%	2	0	0	37	0	9	45/46	97.8%	0	0	1	37	0	9
Piperacillin-tazobactam	41/46	89.1%	1	1	3	29	3	14	40/46	87.0%	0	0	7	29	3	14
TOTAL	381/408	93.4%	5	3	19	295	16	97	373/409	91.2%	1	2	33	295	16	98
<b>Acinetobacter baumannii (n=8)</b>																
Amikacin	7/8	87.5%	0	0	1	2	1	5	7/8	87.5%	0	0	1	2	1	5
Ciprofloxacin	8/8	100%	0	0	0	1	0	7	8/8	100%	0	0	0	1	0	7
Colistin	5/8	62.5%	0	3	0	6	0	2	7/8	87.5%	1	0	0	6	0	2
Meropenem	8/8	100%	0	0	0	1	0	7	8/8	100%	0	0	0	1	0	7
TOTAL	28/32	87.5%	0	3	1	10	1	21	30/32	93.8%	1	0	1	10	1	21
<b>Pseudomonas aeruginosa (n=8)</b>																
Amikacin	7/8	87.5%	0	0	1	3	1	4	8/8	100%	0	0	0	3	1	4
Cefepime	8/8	100%	0	0	0	4	0	4	8/8	100%	0	0	0	4	0	4
Ceftazidime	5/8	62.5%	0	3	0	3	0	5	7/8	87.5%	0	1	0	3	0	5
Ciprofloxacin	7/8	87.5%	1	0	0	1	0	7	8/8	100%	0	0	0	1	0	7
Colistin	8/8	100%	0	0	0	8	0	0	8/8	100%	0	0	0	8	0	0
Gentamicin	8/8	100%	0	0	0	2	0	6	7/8	87.5%	1	0	0	2	0	6
Meropenem	7/8	87.5%	0	0	1	3	0	5	7/8	87.5%	0	0	1	3	0	5
Piperacillin-tazobactam	8/8	100%	0	0	0	3	0	5	7/8	87.5%	0	1	0	3	0	5
TOTAL	58/64	90.6%	1	3	2	27	1	36	60/64	93.8%	1	2	1	27	1	36
<b>Overall</b>	<b>467/504</b>	<b>92.7%</b>	<b>6</b>	<b>9</b>	<b>22</b>	<b>332</b>	<b>18</b>	<b>154</b>	<b>463/505</b>	<b>91.7%</b>	<b>3</b>	<b>4</b>	<b>35</b>	<b>332</b>	<b>18</b>	<b>155</b>

**Table 2:** Performance of Accelerate Pheno™ and VITEK® 2 systems compared to BMD for Gram-positive bacterial species (*n* = 24)

Antibiotic	Accelerate Pheno™ system							VITEK® 2 system								
	CA	VME	ME	mE	S	I	R	CA	VME	ME	mE	S	I	R		
<b>Staphylococcus aureus (n=10)</b>																
Daptomycin	10/10	100%	0	0	0	10	0	0	10/10	100%	0	0	0	10	0	0
Erythromycin	10/10	100%	0	0	0	7	0	3	10/10	100%	0	0	0	7	0	3
Linezolid	10/10	100%	0	0	0	10	0	0	10/10	100%	0	0	0	10	0	0
Trimethoprim-Sulfamethoxazole	9/10	90.0%	0	0	1	10	0	0	10/10	100%	0	0	0	10	0	0
Vancomycin	10/10	100%	0	0	0	10	0	0	10/10	100%	0	0	0	10	0	0
TOTAL	49/50	98.0%	0	0	1	47	0	3	50/50	100%	0	0	0	47	0	3
<b>Coagulase-Negative Staphylococcus spp. (n=6)</b>																
Daptomycin	6/6	100.0%	0	0	0	6	0	0	6/6	100%	0	0	0	6	0	0
Erythromycin	5/5	100.0%	0	0	0	1	0	4	6/6	100%	0	0	0	1	0	5
Linezolid	6/6	100%	0	0	0	6	0	0	6/6	100%	0	0	0	6	0	0
Vancomycin	6/6	100%	0	0	0	6	0	0	6/6	100%	0	0	0	6	0	0
TOTAL	23/23	100%	0	0	0	19	0	4	24/24	100%	0	0	0	19	0	5
<b>Enterococcus spp. (n=8)</b>																
Ampicillin	8/8	100%	0	0	0	5	0	3	8/8	100%	0	0	0	5	0	3
Linezolid	8/8	100%	0	0	0	8	0	0	8/8	100%	0	0	0	8	0	0
Vancomycin	7/7	100%	0	0	0	5	0	2	7/8	87.5%	1	0	0	5	0	3
TOTAL	23/23	100%	0	0	0	18	0	5	23/24	95.8%	1	0	0	18	0	6
<b>Overall</b>	<b>95/96</b>	<b>99.0%</b>	<b>0</b>	<b>0</b>	<b>1</b>	<b>84</b>	<b>0</b>	<b>12</b>	<b>97/98</b>	<b>99.0%</b>	<b>1</b>	<b>0</b>	<b>0</b>	<b>84</b>	<b>0</b>	<b>14</b>



**Figure 1:** Accelerate Pheno™ system workflow.



**Figure 2:** Accelerate Pheno™ system blood culture identification and AST panel.

## Results

The overall categorical agreement between Accelerate Pheno™ system and BMD was 92.7% (467/504) for Gram-negative organisms and 99.0% (95/96) for Gram-positive organisms, with rates for very major errors of 3.6% (6/166), major errors 2.2% (9/416) and minor errors 3.8% (23/600).

The overall categorical agreement between the VITEK® 2 system and BMD was 91.7% (463/505) for Gram-negative organisms and 99.0% (97/98) for Gram-positive organisms, with rates of very major errors of 2.4% (4/169), major errors 1.0% (4/416) and minor errors 5.8% (35/603).

Importantly, in contrast with the VITEK® 2 system, no false-susceptible results occurred with two colistin-resistant organism-growing PBCs tested using the Accelerate Pheno™ system.

## Conclusions

Based on these findings, the Accelerate Pheno™ system can be a valid alternative for the rapid AST of Gram-negative and Gram-positive bacteria in bloodstream infections.

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