1	Brief report

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3	A comparative study between Real-time PCR and LAMP to detect carbapenemase and/or
4	extended-spectrum- $\beta$ -lactamase genes in Enterobacterales directly from bronchoalveolar
5	lavage fluid samples.
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25 **Objectives**: To evaluate and compare the efficacy of Real-time PCR (Xpert Carba-R) and LAMP 26 (Eazyplex<sup>®</sup> SuperBug CRE) for detecting carbapenemase carriage in *Enterobacterales* directly from 27 broncoalveolar lavage (BAL).

Methods: Negative BAL samples were spiked with 21 well-characterized carbapenemase-producing Enterobacterales strains to a final concentration of  $10^2$  to  $10^4$  CFU/mL. Xpert Carba-R, which detects five targets ( $bla_{KPC}$ ,  $bla_{NDM}$ ,  $bla_{VIM}$ ,  $bla_{OXA-48}$ , and  $bla_{IMP-1}$ ), and Eazyplex<sup>®</sup> SuperBug CRE system (Amplex-Diagnostics GmbH, Germany) that detects seven genes ( $bla_{KPC}$ ,  $bla_{NDM}$ ,  $bla_{VIM}$ ,  $bla_{OXA-48}$ ,  $bla_{OXA-181}$ ,  $bla_{CTXM-1}$ , and  $bla_{CTXM-9}$ ), were evaluated to detect these genes directly from BAL samples.

Results: Xpert Carba-R showed 100% agreement with carbapenemase characterization by PCR and sequencing for all final bacteria concentration. Eazyplex® SuperBug CRE showed 100%, 80% and 27% agreement with PCR and sequencing when testing 10<sup>4</sup>, 10<sup>3</sup> and 10<sup>2</sup> CFU/mL, respectively. False negative results for Eazyplex® SuperBug CRE matched with highest Ct values for Xpert Carba-R. Hands-on time for both assays was about 15 minutes, but Eazyplex® SuperBug CRE results were available within 30 minutes, whereas Xpert Carba-R took around 50 minutes.

41 R and Eazyplex<sup>®</sup> SuperBug CRE, to detect bacterial carbapenem resistance genes directly in lower 42 respiratory tract samples. Our results could be used as proof-of-concept data for validation of these 43 tests for this indication.

#### 44 Introduction

Ventilator-associated pneumonia (VAP), the hospital-acquired pneumonia that develops in patients with tracheal intubation or on mechanical ventilation for at least 48 hours, is one of the leading causes of infection and death in the healthcare setting. Patients with VAP present longer periods with mechanical ventilation, as well as longer stay in the ICU and in the hospital.<sup>1</sup>

Unfortunately, the diagnosis of VAP is complicated and there is no a reliable reference test.<sup>2</sup> Microbiological diagnosis is necessary to confirm the clinical suspicion of VAP, but it is not exempt from limitations (negative culture due to empiric treatment, expensive molecular tests, difficulty to differentiate colonization from infection). Early and adequate treatment decreases the mortality,<sup>3</sup> but empirical treatment is usually initiated in patients with suspected VAP before having the definitive diagnosis,<sup>4</sup> which could cause unnecessary health care costs and risk of antibiotic resistance appearance.

56 *Enterobacterales*, such as *Escherichia coli* and *Klebsiella pneumoniae*, are among the most 57 frequent pathogens causing VAP.<sup>5</sup> These microorganisms are also in the WHO priority pathogens 58 list due to the carbapenem and 3<sup>rd</sup> generation cephalosporin resistance.<sup>6</sup>

59 The application of rapid diagnostic techniques to identify microbial pathogens seems to 60 have a huge impact in the management of patients with VAP, reducing inappropriate or 61 unnecessary antimicrobial treatments and mortality in these patients.<sup>7,8,9</sup>

42 Xpert Carba-R (Cepheid, Sunnyvale, USA) is an integrated commercial technique (sample 43 extraction, amplification by PCR and detection carried out within a self-contained cartridge) that 44 detects five targets for carbapenemase-producing organisms (*bla*<sub>KPC</sub>, *bla*<sub>NDM</sub>, *bla*<sub>VIM</sub>, *bla*<sub>OXA-48</sub>, and 45 *bla*<sub>IMP-1</sub>). On the other hand, the Eazyplex<sup>®</sup> SuperBug CRE system (Amplex-Diagnostics GmbH, 46 Germany) combines loop-mediated isothermal amplification (LAMP) of the target and real-time 47 photometric detection of amplified material for rapid and simple detection of extended-spectrum

68 beta-lactamase (ESBL) and carbapenemase-encoding genes (*bla*<sub>CTXM-1</sub>, *bla*<sub>CTXM-9</sub>, *bla*<sub>KPC</sub>, *bla*<sub>NDM</sub>,

69  $bla_{VIM}$ ,  $bla_{OXA-48}$ , and  $bla_{OXA-181}$ ).

The aim of this study was to evaluate and compare the efficacy of Xpert Carba-R and Eazyplex<sup>®</sup> SuperBug CRE for detecting ESBL and carbapenemase carriage directly from broncoalveolar lavage (BAL) samples inoculated with a well-defined collection of *Enterobacterales* clinical isolates.

#### 74 Material and Methods

# 75 Bacterial isolates and samples

76 Twenty-one Enterobacterales strains producing the following carbapenemases were included: four strains producing NDM, five strains producing OXA-48, one strain producing both 77 78 NDM and OXA-48, one strain producing OXA-181, five strains producing KPC and five strains 79 producing VIM (Table 1). Some of the strains were also carrying ESBLs (CTXM-1 or CTXM-9). These 80 organisms were identified using MALDI-TOF (Bruker Daltonics, Bremen, Germany). The presence of 81 genes encoding carbapenemases and ESBL (*bla*<sub>CTXM-1</sub>, *bla*<sub>CTXM-9</sub>, *bla*<sub>KPC</sub>, *bla*<sub>NDM</sub>, *bla*<sub>VIM</sub>, *bla*<sub>OXA-48</sub>, 82 bla<sub>IMP-1</sub> and bla<sub>OXA-181</sub>) was confirmed by PCR amplification with specific primers and Sanger DNA 83 sequencing.<sup>10</sup> Negative BAL samples by bacterial culture were collected from the Clinical 84 Microbiology Laboratory at the Hospital Clinic of Barcelona (Spain) after being processed for 85 routine techniques. They were mixed to obtain a homogeneous matrix and stored at -80°C until use. Negative samples were homogenized by vortexing and spiked with a dilution of each of the 86 selected strains to a final concentration of 10<sup>2</sup> to 10<sup>4</sup> CFU/mL. Two protocols depending on the 87 88 method used are described in Figure 1.

### 89 Sample preparation for Xpert Carba-R

Spiked samples were mixed with sample reagent of the Xpert Carba-R assay (vol:vol), vortexed and incubated at room temperature for ten minutes, vortexing each five minutes. Then, 1.7 mL of the sample reagent was transferred into the cartridge and the assay was run on the GeneXpert platform (Cepheid, Sunnyvale, USA) according to the manufacturer's instructions.

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## 97 Sample preparation for Eazyplex<sup>®</sup> SuperBug CRE

98 Same volume of spiked sample (850  $\mu$ L) was centrifuged at 14,000 x g for 5 min., the pellet 99 was resuspended in 500 µL of resuspension and lysis fluid (RALF), and incubated in a thermal block 100 (99°C) for 2 min. After centrifugation, 25 µL of the supernatant were added to each tube of the 101 strip containing the ready-to-use mastermix. The strip was immediately placed into the Genie II 102 instrument (OptiGene, Horsham, United Kingdom). The reaction mixtures were incubated at 66°C 103 for 30 min with fluorescence monitoring. The Eazyplex® SuperBug CRE test strip contains eight 104 wells with six oligonucleotide primers in each well, allowing the simultaneous specific amplification 105 of seven different resistance genes, plus an internal control.

106 Analysis

107 We evaluated the agreement between the results obtained by conventional methods and 108 rapid test applied directly to the clinical sample. Time to obtain the results and agreement were 109 compared between Xpert Carba-R and Eazyplex<sup>®</sup> SuperBug CRE. Statistical analysis was performed 110 with the software R (version 3.4.4).

### 111 **Results and discussion**

112 Considering the Xpert Carba-R, 100% agreement with PCR and sequencing was observed 113 with all concentrations (Table 1). Regarding Eazyplex<sup>®</sup> SuperBug CRE, the concordance was 100%, 80% and 27% when testing 10<sup>4</sup>, 10<sup>3</sup> and 10<sup>2</sup> CFU/mL, respectively. False negative results for 114 Eazyplex<sup>®</sup> SuperBug CRE matched with highest Ct values for Xpert Carba-R (median Ct of 33.2 and 115 116 28.3 for negative and positive Eazyplex SuperBugCRE resultls, respectively; p<0.001). At 10<sup>2</sup> 117 CFU/mL, most of the genes were not detected in contrast to Xpert Carba-R. This is an integrated 118 assay that includes a DNA extraction step, which increases the amount of DNA for the amplification 119 step compared to LAMP assay.

Hands-on time for both assays was about 15 minutes, but with an easier-to-use sample preparation for Xpert Carba-R. Regarding time-to-results, Eazyplex SuperBugCRE results were available within 30 minutes, whereas Xpert Carba-R took around 50 minutes.

123 In order to distinguish between infection and colonization, it is established that a 124 concentration of microorganisms in BAL  $\geq 10^4$  CFU/mL is indicative of infection,<sup>12</sup> although several 125 factors can modify this interpretation. In a previous study evaluating Xpert Carba-R on BAL 126 samples, a cutoff value to distinguish between colonization (Ct ≤24.7 corresponded to a count of 127  $\geq 10^5$  CFU/mL) and infection (Ct >26.9 corresponded to a count of <10<sup>4</sup> CFU/mL) was proposed.<sup>11</sup> 128 Although we observed a good correlation between the Ct value for Xpert Carba-R and the bacteria 129 concentration (r=-0.81, p<0.001), due to the limited number of samples we could not establish such a cutoff. This was not observed for Eazyplex<sup>®</sup> SuperBug CRE results (r=0.45, p<0.001). 130 131 Although both assays were able to detect the target genes at bacterial concentrations below the 132 diagnostic limit for infection in BAL, the Xpert Carba-R showed a limit of detection 10-fold lower 133 than that of the Eazyplex<sup>®</sup> SuperBug CRE.

We here describe the successful use of two commercial diagnostic tests to detect bacterial
 carbapenem resistance genes directly in lower respiratory tract samples: Xpert Carba-R designed to

- 136 identify patients showing gastrointestinal colonization with carbapenemase-producing organisms,
- 137 and Eazyplex<sup>®</sup> SuperBug CRE designed to detect carbapenemases and ESBL from bacteria strains or
- 138 direct from positive blood culture. Our results could be used as proof-of-concept for the
- 139 development of studies with larger sample size to validate these tests for this indication.
- 140

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# 153 Transparency declarations

154 None to declare.

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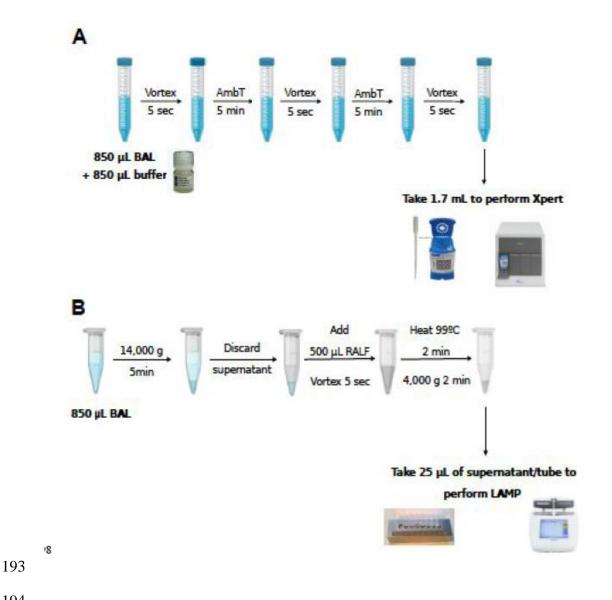
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- 187 **Table 1**. Results of Xpert Carba-R and Eazyplex<sup>®</sup> SuperBug CRE for each strain and concentration
- 188 expressed in Ct values and time, respectively.

- Figure 1. Sample preparation workflow for Xpert Carba-R (Figure 1A) and Eazyplex<sup>®</sup> SuperBug CRE
- (Figure 1B).
- BAL: bronchoalveolar lavage; AmbT: ambient temperature; RALF: resuspension and lysis buffer.



**Table 1**. Results of Xpert Carba-R and Eazyplex<sup>®</sup> SuperBug CRE for each strain and concentration

196 expressed in Ct values and time, respectively.

Bacterial species	Resistance mechanism	Concentration (ufc/mL)	Xpert Carba-R (Target Ct)	Eazyplex <sup>®</sup> SuperBug CRE (Target min:sec)
Klebsiella	ebsiella		NDM 26.3	NDM 9:45; CTXM-1 10:00
pneumoniae	NDM CTXM-1	10 <sup>3</sup>	NDM 28.4	NDM 14:15; CTXM-1 15:30
16-516		10 <sup>2</sup>	NDM 29.3	NDM NEG; CTXM-1 NEG
K nneumoniae	NDM CTXM-1	10 <sup>4</sup>	NDM 26.6	NDM 11:45; CTXM-1 10:15
K. pneumoniae 16-377		10 <sup>3</sup>	NDM 30.5	NDM 17:00; CTXM-1 13:30
		10 <sup>2</sup>	NDM 35.4	NDM NEG; CTXM-1 16:30
Escherichia coli	scherichia coli NDM VR22 CTXM-1	104	NDM 24.2	NDM 8:30; CTXM-1 10:00
		10 <sup>3</sup>	NDM 27.5	NDM 13:30; CTXM-1 11:00
υνκζζ		10 <sup>2</sup>	NDM 30.5	NDM NEG; CTXM-1 22:15
E. coli		10 <sup>4</sup>	NDM 25.7	NDM 8:15; CTXM-1 6:00
16-543	NDM CTXM1	10 <sup>3</sup>	NDM 27.2	NDM 9:30; CTXM-1 8:30
10 343	CIXINI	10 <sup>2</sup>	NDM 30.6	NDM 15:15; CTXM-1 10:00
K. pneumoniae HR4	NDM OXA-48 CTXM1	10 <sup>4</sup>	NDM 27.0; OXA-48 26.7	NDM 12:00; OXA-48 17:15; CTXM-1 11:45
		10 <sup>3</sup>	NDM 31.4; OXA-48 31.2	NDM NEG; OXA-48 NEG; CTXM-1 12:00
		10 <sup>2</sup>	NDM 33.4; OXA-48 33.4	NDM NEG; OXA-48 NEG; CTXM-1 19:00
K. pneumoniae	OXA-48	10 <sup>4</sup>	OXA-48 28.2	OXA-48 11:30
16-420		10 <sup>3</sup>	OXA-48 29.8	OXA-48 18:00
		10 <sup>2</sup>	OXA-48 30.4	NEG
K. pneumoniae	0.44	10 <sup>4</sup>	OXA-48 26.3	OXA-48 10:45; CTXM1 9:45
45-425	OXA-48 CTXM1	10 <sup>3</sup>	OXA-48 30.5	OXA-48 13:15; CTXM1 11:30
		10 <sup>2</sup>	OXA-48 33.1	NEG
E. coli		104	OXA-48 27.0	OXA-48 12:30
16-476 OXA-48	OXA-48	10 <sup>3</sup>	OXA-48 28.3	OXA-48 20:00
		10 <sup>2</sup>	OXA-48 31.3	NEG
K. pneumoniae		104	OXA-48 29.7	OXA-48 13:30; CTXM1 10:30
66134	OXA-48 CTXM1	10 <sup>3</sup>	OXA-48 31.7	OXA-48 17:00; CTXM1 13:30
	CIMUL	10 <sup>2</sup>	OXA-48 35.3	OXA-48 NEG; CTXM1 NEG
E. coli	OXA-48	10 <sup>4</sup>	OXA-48 29.4	OXA-48 13:45
65273		10 <sup>3</sup>	OXA-48 29.9	OXA-48 14:30
		10 <sup>2</sup>	OXA-48 30.3	OXA-48 NEG

E. coli		<b>10</b> <sup>4</sup>	OXA-48 28.0	OXA-181 14:15; CTXM1 12:00
15-288	OXA-181 CTXM1	10 <sup>3</sup>	OXA-48 31.0	OXA-181 23:30; CTXM1 NEG
13 200	CIANI	10 <sup>2</sup>	OXA-48 33.4	OXA-181 NEG; CTXM1 NEG
K		10 <sup>4</sup>	KPC 28.3	KPC 19:30
K. pneumoniae	КРС	10 <sup>3</sup>	KPC 31.1	NEG
MB545		10 <sup>2</sup>	KPC 35.3	NEG
K. pneumoniae	КРС	10 <sup>4</sup>	KPC 28.5	KPC 17:15
		10 <sup>3</sup>	KPC 32.6	KPC 21:00
45872		10 <sup>2</sup>	KPC 35.9	NEG
E. coli		10 <sup>4</sup>	KPC 28.6	KPC 16:45
182046440	КРС	10 <sup>3</sup>	KPC 31.2	KPC 22:50
		10 <sup>2</sup>	KPC 35.5	NEG
Enterobacter		10 <sup>4</sup>	KPC 28.6	KPC 18:15; CTXM-1 13:15
asburiae	KPC CTXM1	10 <sup>3</sup>	KPC 30.4	KPC 18:45; CTXM-1 14:00
	CIXIII	10 <sup>2</sup>	KPC 33.1	KPC NEG; CTXM-1 NEG
K. pneumoniae	KPC CTXM1	10 <sup>4</sup>	KPC 28.1	KPC 16:00; CTXM-1 11:00
		10 <sup>3</sup>	KPC 31.0	KPC 21:00; CTXM-1 11:45
19-684		10 <sup>2</sup>	KPC 34.8	KPC NEG; CTXM-1 NEG
K. pneumoniae		10 <sup>4</sup>	VIM 22.3	VIM 13:00
18-87	VIM	10 <sup>3</sup>	VIM 28.3	VIM 19:45
		10 <sup>2</sup>	VIM 31.1	NEG
E. coli		10 <sup>4</sup>	VIM 24.1	VIM 9:30
18-38	VIM	10 <sup>3</sup>	VIM 26.7	VIM 12:15
		10 <sup>2</sup>	VIM 30.6	VIM 19:45
K. pneumoniae		10 <sup>4</sup>	VIM 25.1	VIM 15:30
MC-32-145	VIM	10 <sup>3</sup>	VIM 28.9	VIM 18:00
		10 <sup>2</sup>	VIM 31.3	VIM 19:30
Enterobacter		10 <sup>4</sup>	VIM 27.2	VIM 12:15; CTXM9 14:15
cloacae	VIM	10 <sup>3</sup>	VIM 32.5	VIM 16:15; CTXM9 18:30
MC-13-3	CTXM9	10 <sup>2</sup>	VIM 36.7	VIM NEG; CTXM9 19:45
E. coli	VIM CTXM9	10 <sup>4</sup>	VIM 25.8	VIM 12:45; CTXM9 17:30
		10 <sup>3</sup>	VIM 29.9	VIM 18:20; CTXM9 18:15
MC-4-19				