

# Practical Management of HAP/VAP



전 경 만

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# Korean HAP/VAP Study

Multicenter Retrospective Cohort, Koran HAP/VAP Study Group

**서울**

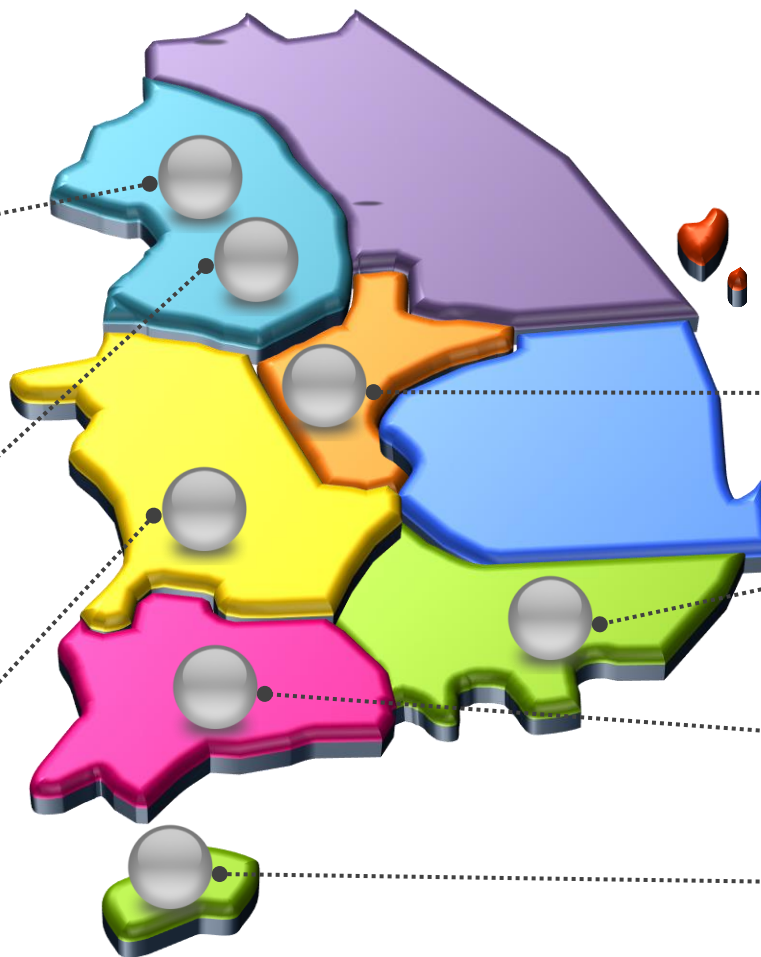
- 삼성서울병원
- 서울아산병원
- 건국대학병원
- 고려대학 구로병원
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**경기**

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**전북**

- 전북대학병원



**대전 충남**

- 충남대학병원

**부산 경남**

- 양산부산대학병원
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**광주 전남**

- 전남대학병원

**제주**

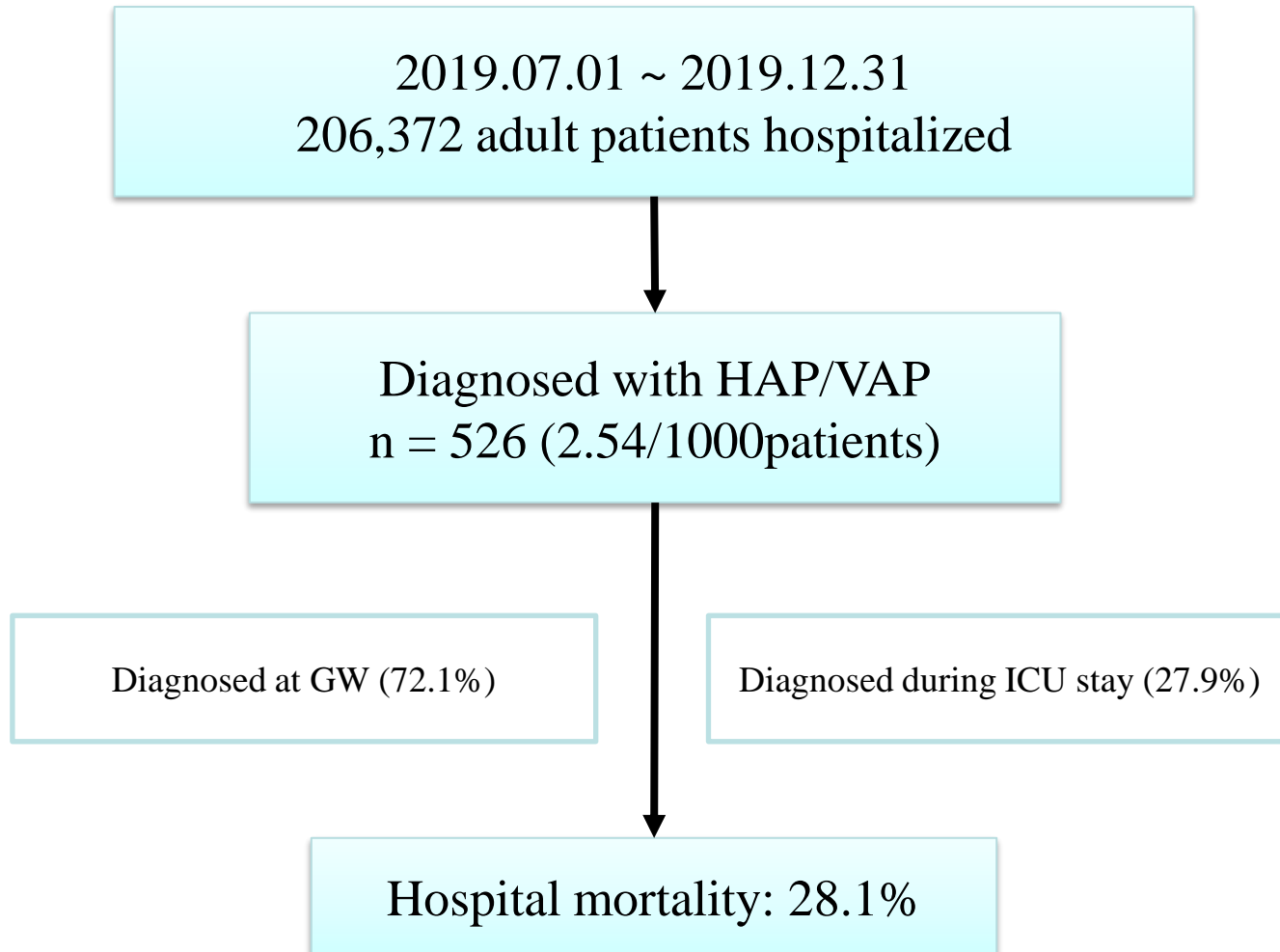
- 제주대학병원

# Study Population



- All consecutive adult patients who admitted to the **13** participating hospitals during a **six-month** (July 1 through December 31, 2019) **period** were screened for eligibility.
- Diagnosis of HAP/VAP with IDSA/ATS diagnostic criteria
  - **New lung infiltrate plus clinical evidence** that the infiltrate is of an infectious origin, which include
    - New onset of fever
    - Purulent sputum
    - Leukocytosis
    - Decline in oxygenation
  - HAP is defined as a pneumonia not incubating at the time of hospital admission and occurring 48 hours or more after admission.
  - VAP is defined as a pneumonia occurring >48 hours after endotracheal intubation.

# Prevalence of HAP/VAP



# Baseline Characteristics



Variables	No. of patients or Median (IQR)
Age, year	71.0 (62.0–79.0)
Sex, male	360 (68.4)
Reason for admission	
Medical	
Diagnostic work up	303 (57.6)
Medical disease treatment	52 (9.9)
Surgical	
Elective operation	106 (20.2)
Emergency operation	52 (9.9)
Other reasons than operation	13 (2.5)
Charlson comorbidity index	5.0 (3.0–6.0)
Clinical frailty scale	5.0 (3.0–7.0)
Risk for aspiration	333 (63.3)
Impaired swallowing	223
Impaired consciousness	171
Increased change of gastric contents reaching the lung	202
Impaired cough reflex	156

# Microorganism Isolated (n=211, 40.1%)

Microorganism	No. of patients
Gram-positive pathogens	
<i>Staphylococcus aureus</i>	24
<i>Streptococcus pneumoniae</i>	7
<i>Enterococcus faecium</i>	5
<i>Nonstaphylococcus aureus Staphylococcus species</i>	4
Others*	5
Gram-negative pathogens	
<i>Acinetobacter baumannii</i>	68
<i>Pseudomonas aeruginosa</i>	36
<i>Klebsiella pneumoniae</i>	35
<i>Escherichia coli</i>	11
<i>Sternotrophomonas maltophilia</i>	11
<i>Enterobacter cloacae</i>	10
<i>Serratia marcescens</i>	5
<i>Proteus species</i>	4
<i>Klebsiella aerogenes</i>	3
<i>Moraxella catarrhalis</i>	2
<i>Burkholderia cephalica</i>	2
<i>Citrobacter species</i>	2
Others**	2

\*Others included one *Streptococcus pyogenes*, one *Streptococcus mitis*, one *Streptococcus agalactiae*, one *Streptococcus angiosus*, and one *Listeria monocytogenes*.

\*\*Others included one *Klebsiella oxytoca* and one *Raoultella planticola*.

# Clinical Findings



Variables	No. of patients or Median (IQR)
Initial empirical antibiotics	
Extended-spectrum penicillin/beta-lactamase inhibitor	312 (59.3)
Respiratory fluoroquinolone	169 (32.1)
3 <sup>rd</sup> cephalosporin	53 (10.1)
Cefepime	35 (6.7)
Aminoglycoside	13 (2.5)
Glycopeptide	79 (15.0)
Carbapenem	107 (20.3)
Colistin	8 (1.5)
Macrolide	5 (1.0)
Others	44 (8.4)
More than two drugs	249 (47.3)
Adjunctive corticosteroid treatment	91 (17.3)
Classification of infection	
Microbiologically documented	213 (40.3)
Clinically documented	297 (56.2)
Possible infection	18 (3.4)
Inappropriate antibiotic treatment (n = 211)	102 (48.3)
Occurrence of MDR pathogen (n=211)	138 (70.4)
Staphylococcus aureus	67
Enterococcus spp.	30
Enterobacteriaceae	27
Pseudomonas aeruginosa	19
Acinetobacter spp.	4
Duration of antibiotic treatment, day	13.0 (7.0–24.0)

# Outcomes of HAP/VAP



Variables	No. of patients or Median (IQR)
Clinical response	
Clinical cure	344 (65.4)
Clinical Failure	171 (32.5)
Recurrence	11 (2.1)
Microbiological response (n=211)	
Microbiological eradication	106 (57.9)
Colonization	23 (12.6)
Microbiological failure	50 (27.3)
Microbiological recurrence	4 (2.2)
Additional ICU admission	107 (28.2)
Hospital mortality	28.1%
Hospital length of stay, day	30 [18–53]
Discharge destination (n=378)	
Home	205 (54.2)
Transfer	173 (45.8)
Step-down referral	158
Step-up referral	15
Limitation of life-sustaining treatment at any time	157 (29.8)

# Summary of Korean HAP/VAP Study



- The prevalence of HAP/VAP in adult patients hospitalized during a six-month period in Korea was **2.54/1000 hospitalized patients**.
- **Overall in-hospital mortality was 28.1%** in adult patients with HAP/VAP in Korea.
- Patients with HAP/VAP from tertiary hospitals in Korea were **elderly, had a risk of aspiration, and were often referred to step-down centers**.

KATRDIC 2020



# 66 Male with AGC

## Progression of New Lung Infiltrates



POD#1



POD#2



POD#3

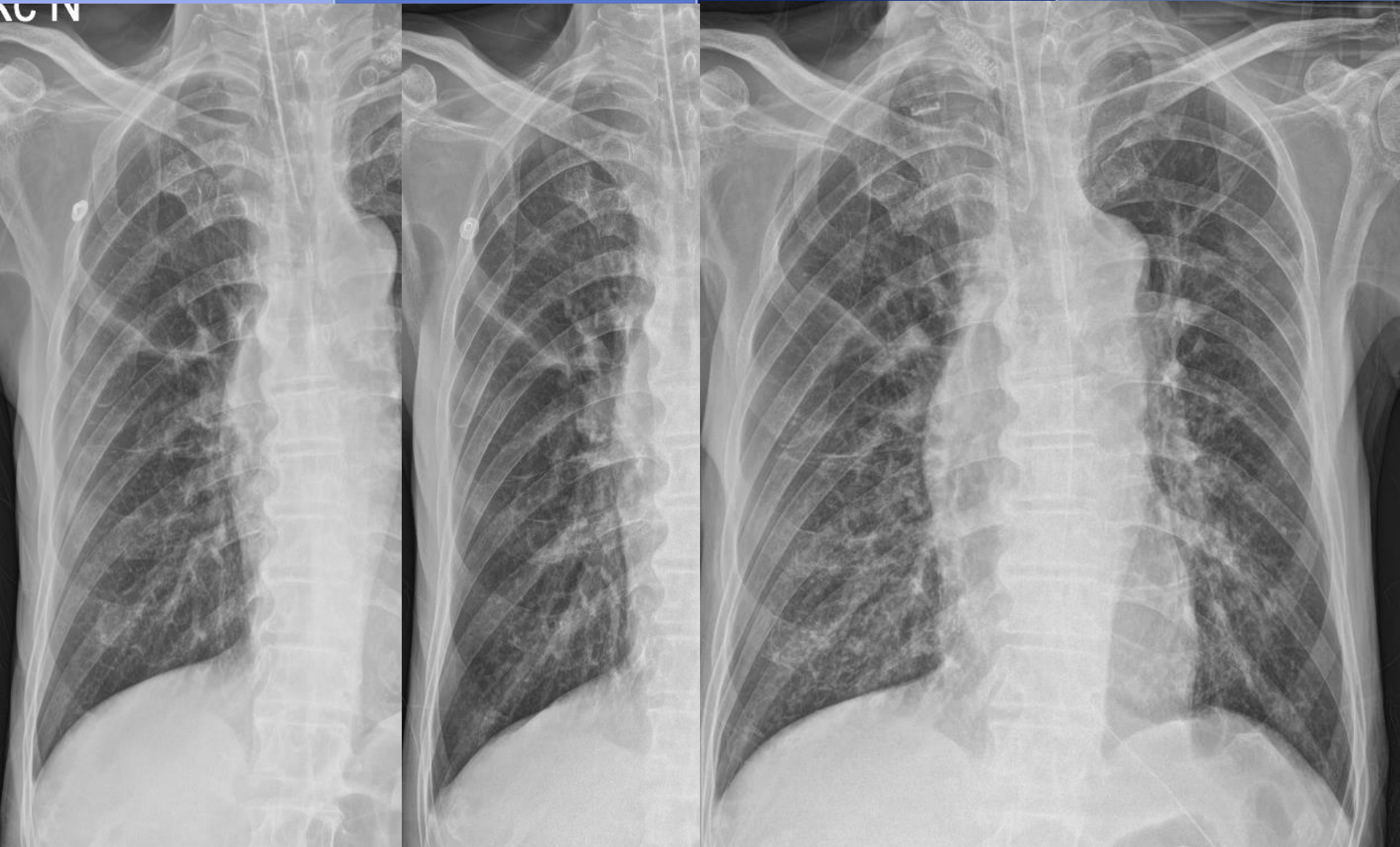
Leukocytosis (+, 21,290), purulent sputum (+),  
severe hypoxemia (+), new onset of fever (-)

CRP 13.54 mg/dL, PCT 6.48 ng/mL

# Sensitivity & Specificity of Clinical Variables In Diagnosis of VAP

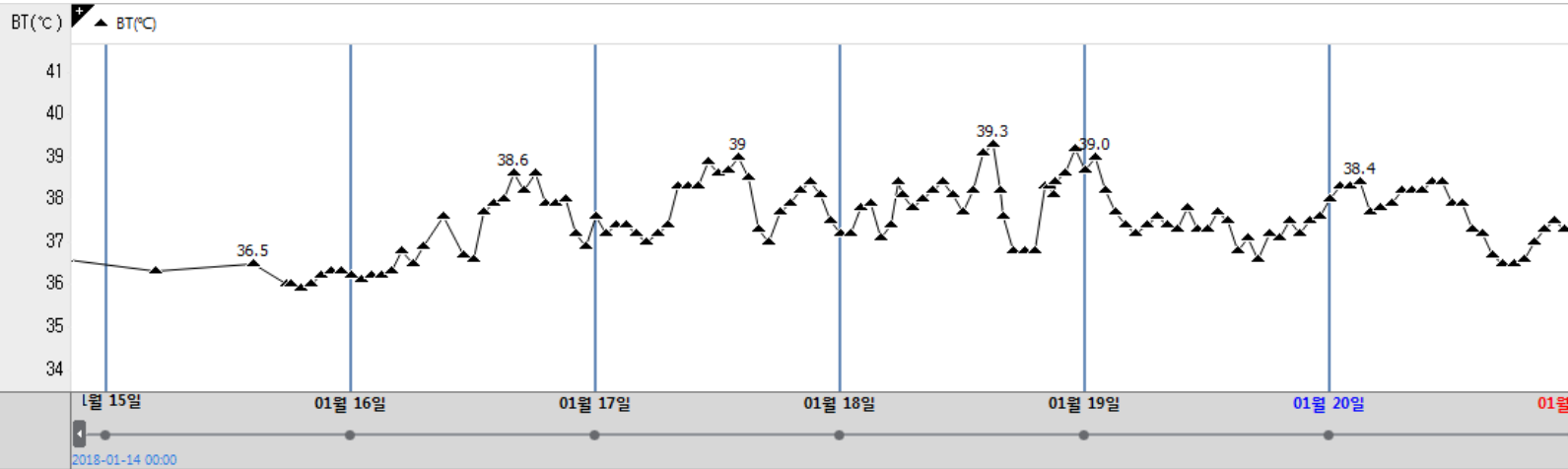
Finding and Source	Gold Standard	Sensitivity, %	Specificity, %	LR (95% CI)	
				Positive	Negative
<b>Fever</b>					
Independent					
Chastre et al, <sup>39</sup> 1984	Histology alone	67	65	1.9 (0.84-4.3)	0.51 (0.16-1.7)
Torres et al, <sup>41</sup> 1994	Histology alone	56	58	1.3 (0.61-2.9)	0.76 (0.38-1.5)
Fàbregas et al, <sup>45</sup> 1999	Histology and culture	46	42	0.79 (0.37-1.7)	1.3 (0.56-3.0)
Summary				<u>1.2 (0.76-1.9)</u>	<u>0.86 (0.54-4.1)</u>
<b>Abnormal WBC count</b>					
Independent					
Chastre et al, <sup>39</sup> 1984	Histology alone	50	45	0.91 (0.37-2.2)	1.1 (0.44-2.8)
Fàbregas et al, <sup>45</sup> 1999	Histology and culture	77	58	1.8 (0.89-3.8)	0.40 (0.13-1.2)
Summary				<u>1.3 (0.76-2.4)</u>	<u>0.74 (0.34-1.6)</u>
<b>Sputum purulence, macroscopic</b>					
Independent					
Torres et al, <sup>41</sup> 1994	Histology alone	83	33	1.3 (0.80-2.0)	0.50 (0.14-1.8)
Fàbregas et al, <sup>45</sup> 1999	Histology and culture	69	42	1.2 (0.65-2.2)	0.74 (0.26-2.1)
Summary				<u>1.3 (0.88-1.8)</u>	<u>0.63 (0.28-1.4)</u>
<b>Crepitation on auscultation</b>					
Nonindependent					
Petersen et al, <sup>10</sup> 1999	Histology alone	73	40	<u>1.2 (0.75-2.0)</u>	<u>0.68 (0.27-1.7)</u>
<b>Hypoxemia</b>					
Nonindependent					
Petersen et al, <sup>10</sup> 1999	Histology alone	64	40	<u>1.1 (0.63-1.8)</u>	<u>0.91 (0.40-2.1)</u>
<b>New infiltrate on radiograph</b>					
Independent					
Chastre et al, <sup>39</sup> 1984	Histology alone	100	75	3.5 (1.7-7.5)	0.10 (0.01-1.4)
Torres et al, <sup>41</sup> 1994	Histology alone	78	42	1.3 (0.78-2.3)	0.53 (0.18-1.6)
Fàbregas et al, <sup>45</sup> 1999	Histology and culture	92	33	1.4 (0.88-2.2)	0.24 (0.03-1.8)
Summary				<u>1.7 (1.1-2.5)</u>	<u>0.35 (0.14-0.87)</u>

# 67 YO Male on MV for Respiratory Failure



# Delayed Diagnosis of VAP

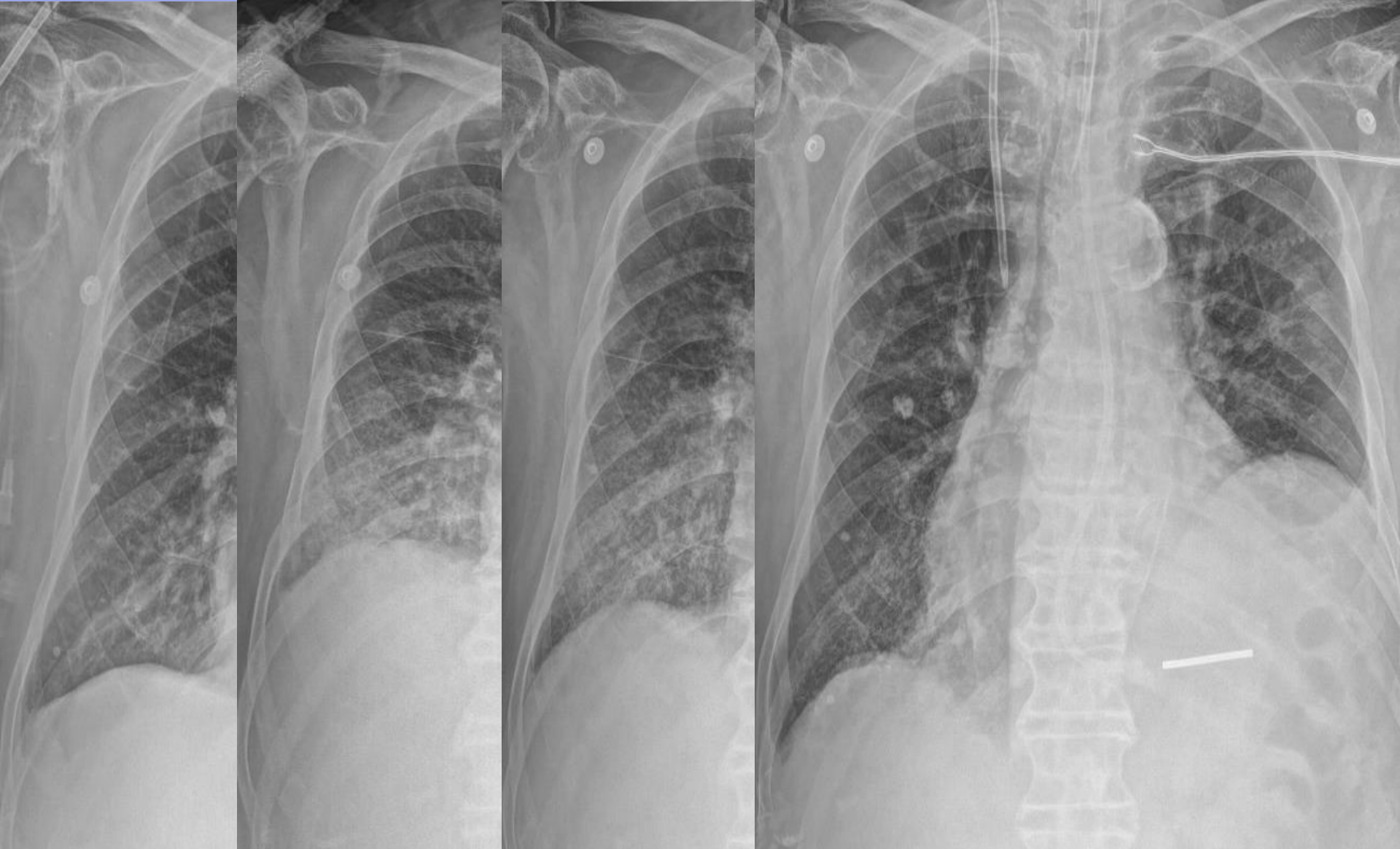
## Slow Progression on CXR



그룹명	항목명	01월 15일	01월 16일	01월 17일	01월 18일	01월 19일	01월 20일	01월 21일
Vital Sign								
BT	BT(°C)	36.5		38.6	39	39.3	39	38.4
검사결과								
	WBC Count, Blood (x10 <sup>3</sup> /μL)			12.74	17.38	12.76	8.57	8.16
	ESR (Erythrocyte Sedimentation Rate)							84
	CRP, Quantitative (High Sensitivity) (mg)		0.77		14.5	29.35	22.12	14.45
	Procalcitonin, quantitative (ng/mL)				0.64			1.36



# 88 YO Male with AE of COPD



# Difficulty in Clinical Diagnosis of HAP/VAP

- Prospective observation with suspected VAP (n = 84)
  - New & persistent infiltration on CXR & purulent tracheal aspirates
  - Clinical/histologic/microbiologic diagnosis

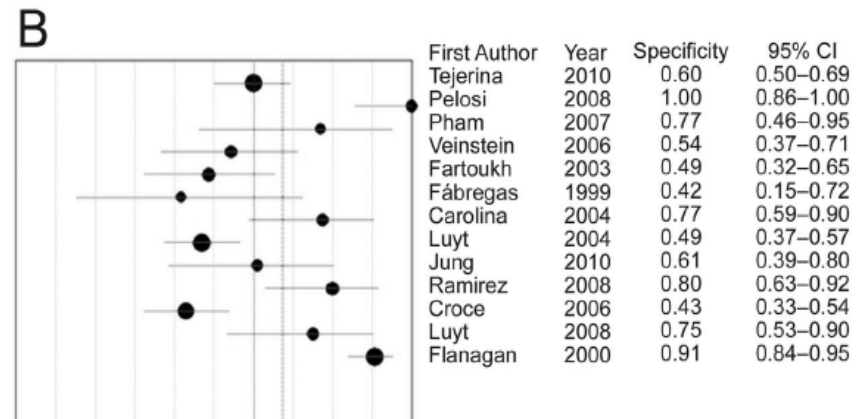
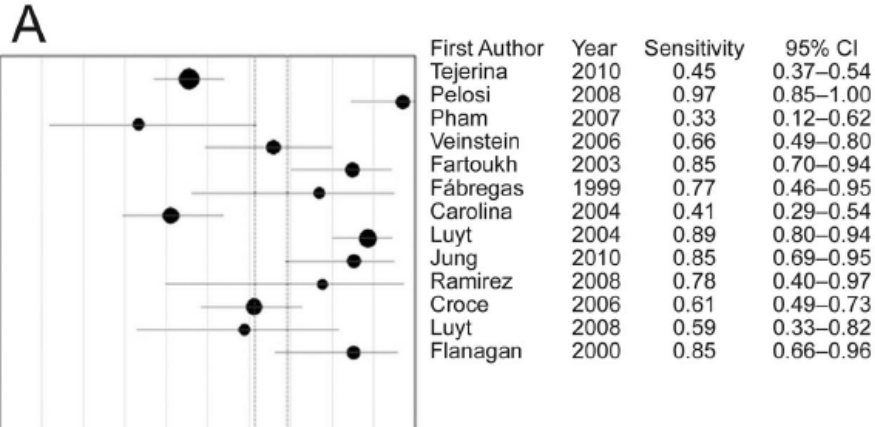
Have pneumonia		Not have pneumonia	
Definite	Probable	Definite	Probable
17 (20%)	10 (12%)	34 (41%)	23 (27%)

- Evaluation of clinical diagnosis ► **not permit the accurate diagnosis**

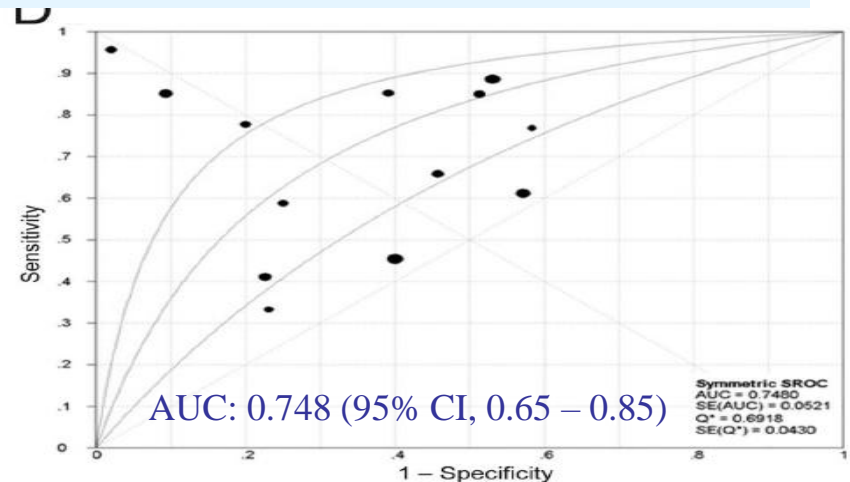
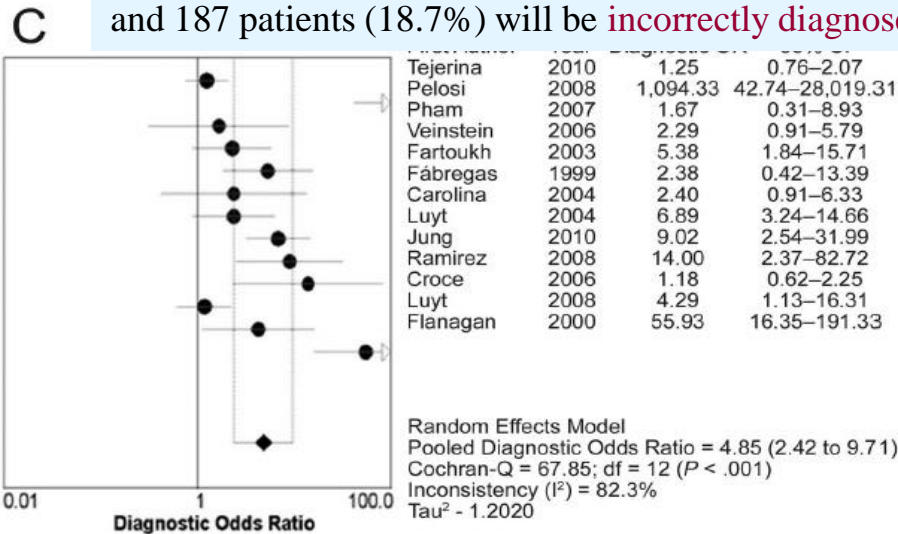
Predictions	Percentage of Accurate Prediction		
	All patients (n = 84)	Patients Who Had Pneumonia (n = 27)	Patients Who Did Not Have Pneumonia (n = 57)
All predictions	77	62	84
Predictions of			
Senior consultants (n = 110)	77	57	84
Staff physicians (n = 186)	72	56	81
Residents (n = 112)	78	70	81
Predictions			
Of best predictor	82	72	86
Of worst predictor	71	50	83
When decision was unanimous (n = 49)	90	79	94

# Diagnostic Accuracy of CPIS

## Systematic Review and Meta-analysis



In a prevalence of VAP of 48%,  
 For every 1000 patients tested, 168 patients (16.8%) will be **incorrectly diagnosed as not having HAP/VAP**  
 and 187 patients (18.7%) will be **incorrectly diagnosed as having VAP**



# Biomarkers for Dx of HAP/VAP

## Results from Meta-analysis



Serum Procalcitonin				
Sensitivity	Specificity	PLR	NLR	Diagnostic OR
67% (53%–79%)	83% (43%–97%)	3.9 (9–17.5)	0.4 (.25–.62)	10 (2–59)
AUC: 0.76 (95% CI, .72–.79)				

In a prevalence of HAP/VAP of 50%,  
For every 1000 patients with suspected HAP/VAP who are evaluated with serum PCT plus clinical criteria, 165 patients (16.5%) would be incorrectly diagnosed as not having HAP/VAP and 85 patients (8.5%) will be incorrectly diagnosed as having HAP/VAP.

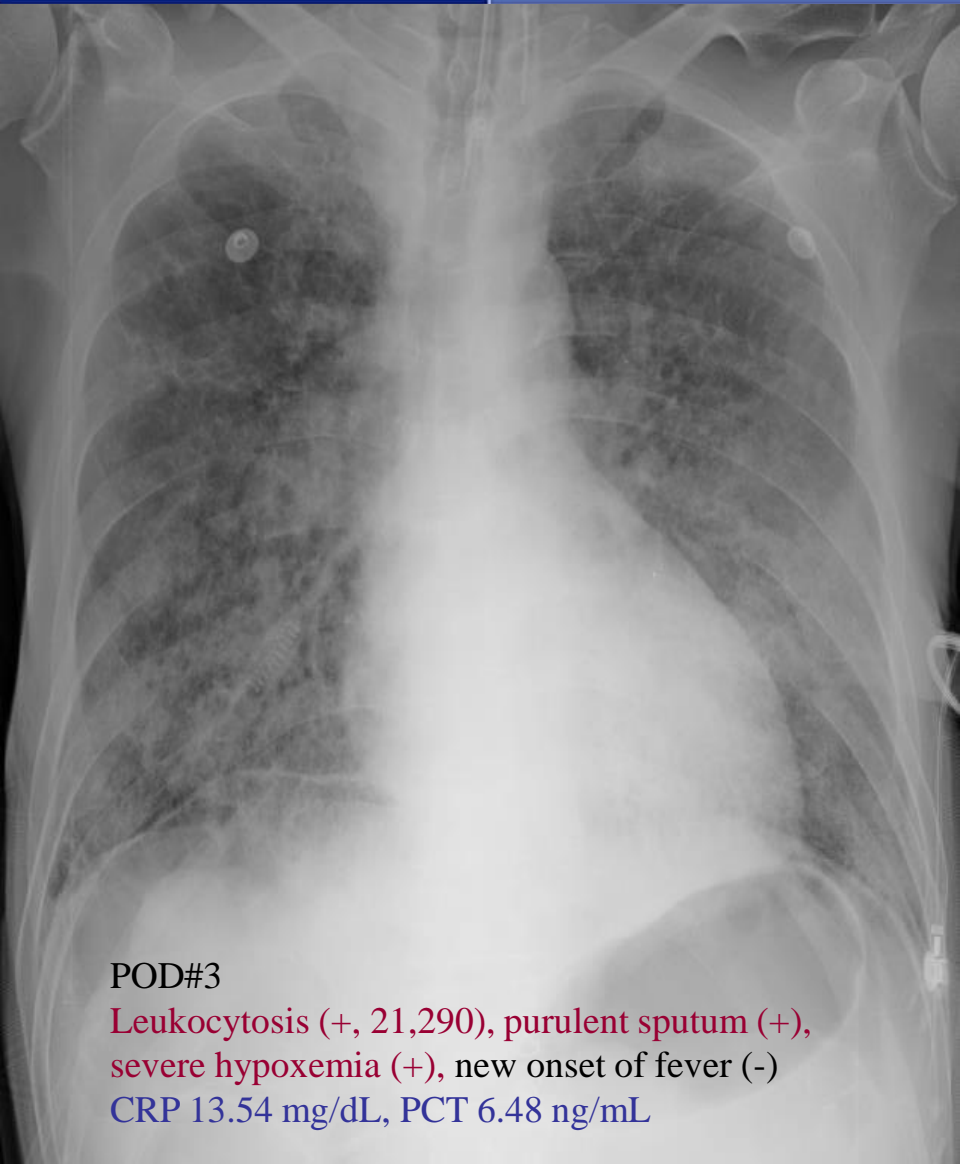
BAL fluid sTREM-1				
Sensitivity	Specificity	PLR	NLR	Diagnostic OR
84% (63%–94%)	49% (18%–81%)	1.6 (.8%–3.3)	0.33 (.12–0.93)	5 (1–24),
AUC: 0.78 (95% CI, .75–.82)				

In a prevalence of HAP/VAP of 50%,  
For every 1000 patients tested, 80 patients (8%) will be incorrectly diagnosed as not having HAP/VAP and 255 patients (25.5%) will be incorrectly diagnosed as having HAP/VAP.

► For patients with suspected HAP/VAP, **the guideline recommends using clinical criteria alone**, rather than using **serum PCT** or **BAL fluid sTREM-1** plus clinical criteria, to decide whether or not to initiate antibiotic therapy (strong recommendation, moderate-quality evidence).

# 66 Male with AGC

## Diagnosis of HAP/VAP



# Classification of HAP/VAP by Time of Onset

## Early vs. Late

- Rationale of prior classification
  - Upper airway colonization was an independent predictor of subsequent tracheobronchial colonization
  - Colonization patterns in the upper and lower airways changed within the first 3–4 days from a community-like to a typical nosocomial pattern

Early onset	Late onset
Within < 5 days	After $\geq$ 5 days
<ul style="list-style-type: none"><li>• Antibiotic sensitive bacteria<ul style="list-style-type: none"><li>- <i>Streptococcus pneumoniae</i></li><li>- <i>Haemophilus influenzae</i></li><li>- <i>Staphylococcus aureus</i></li></ul></li><li>• Better prognosis</li></ul>	<ul style="list-style-type: none"><li>• <b>Multidrug-resistant (MDR) pathogens</b><ul style="list-style-type: none"><li>- <i>P. aeruginosa</i></li><li>- <i>Acinetobacter</i> species</li><li>- MRSA</li></ul></li><li>• Associated with increased mortality and morbidity</li></ul>

# Early- and Late-Onset Pneumonia

## Is This Still a Useful Classification?



TABLE 3. Isolates per 100 pneumonia cases for the most frequent microorganisms, together with the order of frequency of the four most frequent pathogens<sup>a</sup>

Pathogen	No. of isolates per 100 pneumonia cases (order of frequency)					
	“Early-onset” pneumonia			“Late-onset” pneumonia		
	1–4 days	1–5 days	1–7 days	>4th day	>5th day	>7th day
<i>S. aureus</i>	25.7 (1st)	26.8 (1st)	26.9 (1st)	23.7 (1st)	23.0 (1st)	21.0 (1st)
MSSA	21.4	22.9	22.7	16.8	13.8	14.5
MRSA	4.3	4.0	4.3	6.9	6.5	6.5
<i>P. aeruginosa</i>	11.6 (2nd)	11.6 (2nd)	11.9 (2nd)	17.4 (2nd)	16.1 (2nd)	19.9 (2nd)
<i>K. pneumoniae</i>	10.8 (3rd)	10.7 (4th)	11.1 (3rd)	11.8 (3rd)	10.6 (3rd)	12.6 (3rd)
<i>E. coli</i>	10.0 (4th)	10.8 (3rd)	10.6 (4th)	10.1 (4th)	8.7 (4th)	10.1 (4th)
<i>S. pneumoniae</i>	9.3	8.9	8.3	5.1	4.2	4.3
<i>Enterobacter</i> spp.	6.4	6.7	7.5	8.8	7.9	9.4
<i>Haemophilus</i> spp.	6.9	6.9	6.7	3.9	3.1	2.9
<i>Acinetobacter</i> spp.	2.6	2.6	3.2	4.8	4.5	5.5
<i>S. maltophilia</i>	1.3	1.3	1.4	3.1	3.0	3.8

<sup>a</sup> From KISS, 1997 to 2004.

Thus, the predictabilities of the occurrence of pathogens were similar for the earlier (1997-to-2004) and later (2005-to- 2006) time frames. This classification is no longer helpful for empirical antibiotic therapy, since the pathogens are the same for both groups.

# Risk Factors for MDR Pathogens

## ATS/IDSA, 2016



### Risk factors for MDR VAP

Prior intravenous antibiotic use within 90 d

Septic shock at time of VAP

ARDS preceding VAP

Five or more days of hospitalization prior to the occurrence of VAP

Acute renal replacement therapy prior to VAP onset

### Risk factors for MDR HAP

Prior intravenous antibiotic use within 90 d

### Risk factors for MRSA VAP/HAP

Prior intravenous antibiotic use within 90 d

### Risk factors for MDR Pseudomonas VAP/HAP

Prior intravenous antibiotic use within 90 d

### Factors **not** associated with MDR risk

Re-intubation

Immunosuppression

Chronic respiratory failure

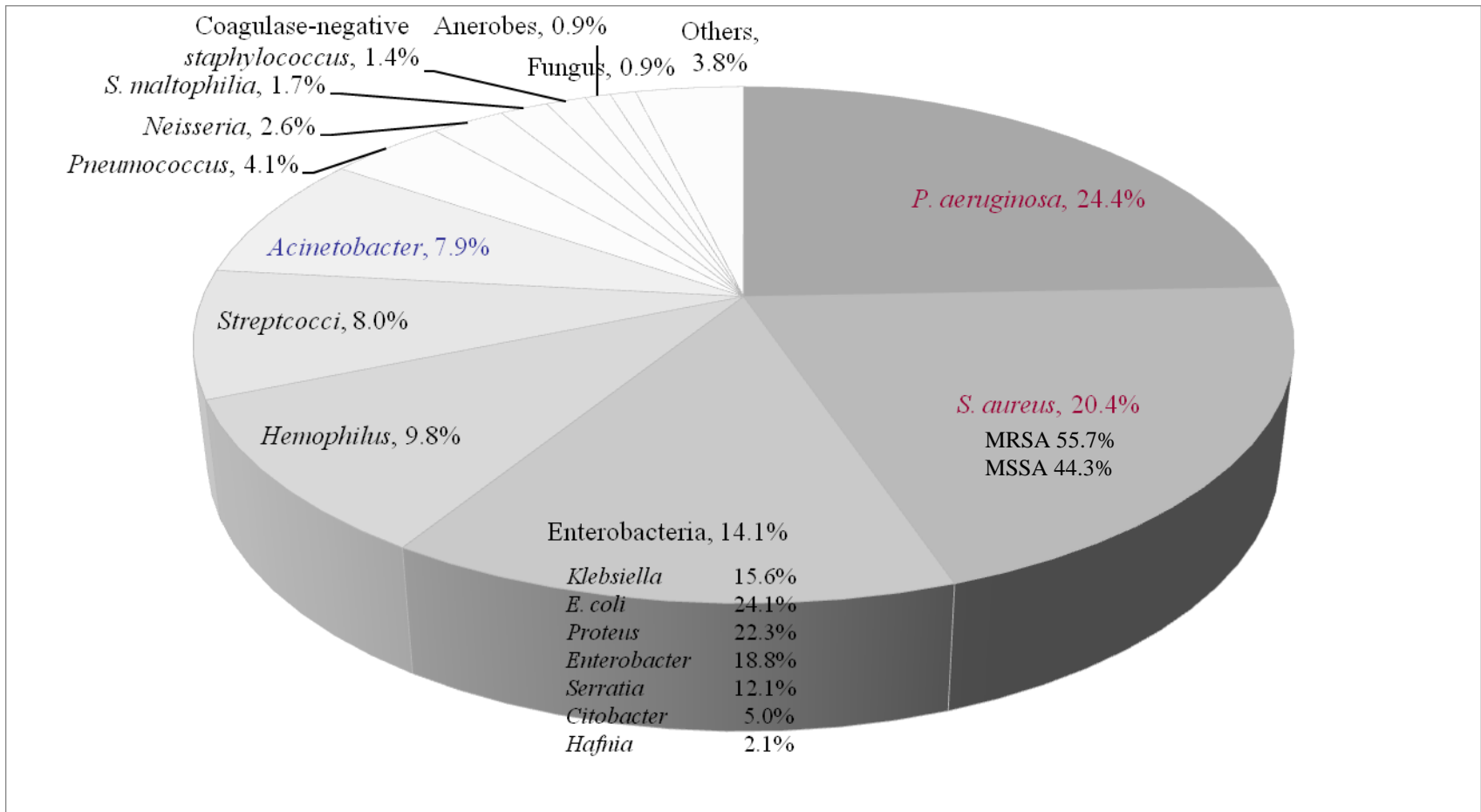
Tracheostomy

Diabetes

Recent use of corticosteroids

# Microbiologic Epidemiology of VAP

## Data from 24 Studies in US



# Microbiologic Epidemiology of VAP

National Healthcare Safety Network (NHSN), U.S., 2009-2010

Pathogen	Frequency	Antimicrobial Resistance Rates
<i>Staphylococcus aureus</i>	24.1%	Methicillin / oxacillin resistant – 48%
<i>Pseudomonas aeruginosa</i>	16.6%	Ciprofloxacin / levofloxacin resistant – 33% Imipenem / meropenem resistant – 30% Cefepime / ceftazidime resistant – 28% Piperacillin-tazobactam resistant – 19% Aminoglycoside resistant – 11% Resistant to $\geq 3$ of the above classes – 18%
<i>Klebsiella species</i>	10.1%	Cefepime / ceftazidime / cefotaxime resistant – 24% Imipenem / meropenem resistant – 11% Resistant to $\geq 3$ classes – 13%
<i>Enterobacter species</i>	8.6%	Cefepime / ceftazidime / ceftriaxone resistant – 30% Imipenem / meropenem resistant – 4% Resistant to $\geq 3$ classes – 1%
<i>Acinetobacter baumannii</i>	6.6%	Imipenem / meropenem resistant – 61% Resistant to $\geq 3$ classes – 63%
<i>Escherichia coli</i>	5.9%	Ciprofloxacin / levofloxacin resistant – 35% Cefepime / ceftazidime / ceftriaxone resistant – 16% Imipenem / meropenem resistant – 4% Resistant to $\geq 3$ classes – 3%

# Differences in Causative Pathogens



Pathogen	Percentage of cases			
	United States ( <i>n</i> = 2585)		All regions ( <i>n</i> = 7496)	
	HABP	VABP	HABP	VABP
<i>Staphylococcus aureus</i>	36.5 <sup>a</sup>	31.9 <sup>a</sup>	26.6 <sup>a</sup>	19.5 <sup>a</sup>
<i>Pseudomonas aeruginosa</i>	19.0 <sup>a</sup>	21.4 <sup>a</sup>	22.4 <sup>a</sup>	26.6 <sup>a</sup>
<i>Enterobacter</i> species	8.6	8.8	7.5	7.0
<i>Klebsiella</i> species	8.0	6.6	10.5	10.2
<i>Serratia</i> species	5.5	6.5	4.1	4.1
<i>Acinetobacter</i> species	4.4	5.3	8.3 <sup>a</sup>	14.3 <sup>a</sup>
Top 6 species	80.4	80.5	79.4	81.7
Pathogens causing CABP <sup>b</sup>	3.3	6.6	2.6	4.1

# Microbiologic Epidemiology of VAP in Asia

## Asian HAP Working Group, 2008



	India	Thailand	Malaysia	China	Korea	Taiwan
<i>Acinetobacter</i> spp	38%	30%	23%	16%	26%	20%
<i>P. aeruginosa</i>	20%	18%	18%	18%	18%	21%
MRSA	5%	8%	12%	16%	34%	18%
<i>Klebsiella</i> spp	23%	7%	6%	14%	9%	9%
<i>E. coli</i>	-	-	-	6%	-	4%
Enterobacteriaceae	-	-	-	8%	-	3%
<i>S. maltophilia</i>	2%	-	11%	-	6%	3%

Am J Infect Control 2008;36:S83

# Microbiologic Epidemiology of VAP in SMC MICU, 2017.01



접수일자	보고일자	검사코드	검사명	검체명	미생물명
2017-01-01 13:50:48	2017-01-03 10:12:48	BL4020	Gram Stain/Culture	(Endo)Tracheal Aspir	Acinetobacter baumannii complex (CRGNB)
2017-01-03 11:24:00	2017-01-06 11:02:50	BL4020	Gram Stain/Culture	(Endo)Tracheal Aspir	Staphylococcus aureus (MRSA)
2017-01-06 22:10:53	2017-01-10 11:52:35	BL4020	Gram Stain/Culture	(Endo)Tracheal Aspir	Acinetobacter baumannii complex (CRGNB)
2017-01-08 00:20:04	2017-01-10 11:58:22	BL4020	Gram Stain/Culture	(Endo)Tracheal Aspir	Stenotrophomonas maltophilia
2017-01-08 18:57:24	2017-01-11 11:08:04	BL4020	Gram Stain/Culture	(Endo)Tracheal Aspir	Pseudomonas aeruginosa
2017-01-16 11:32:36	2017-01-18 11:43:07	BL4020	Gram Stain/Culture	(Endo)Tracheal Aspir	Serratia marcescens
2017-01-19 10:13:24	2017-01-21 10:27:46	BL4020	Gram Stain/Culture	(Endo)Tracheal Aspir	Pseudomonas aeruginosa
2017-01-24 18:25:52	2017-01-27 18:05:34	BL4020	Gram Stain/Culture	(Endo)Tracheal Aspir	Pseudomonas aeruginosa

# Microbiologic Epidemiology of VAP in SMC MICU, 2017.09~11

Pathogen for VAP	Number (%)	Drug resistance
<i>Staphylococcus aureus</i>	11 (24%)	MRSA
<i>Acinetobacter baumannii</i>	10 (22%)	CR 9 (90%)
<i>Pseudomonas aeruginosa</i>	8 (18%)	CR 6 (75%)
<i>Stenotrophomonas maltophilia</i>	7 (16%)	
Enterobacteriaceae	6 (13%)	
Others	3	

► Individual hospitals should determine their most appropriate antimicrobial regimens for the empiric treatment of VAP **base on the hospital specific microbiologic information**

Chest 2000;117:1434

Infect Control Hosp Epidemiol 2007; 28:389

# Microorganism Isolated (n=211, 40.1%) Korean HAP/VAP Study



Microorganism	No. of patients
Gram-positive pathogens	
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Others*	5
Gram-negative pathogens	
<i>Acinetobacter baumannii</i>	68
<i>Pseudomonas aeruginosa</i>	36
<i>Klebsiella pneumoniae</i>	35
<i>Escherichia coli</i>	11
<i>Sternotrophomonas maltophilia</i>	11
<i>Enterobacter cloacae</i>	10
<i>Serratia marcescens</i>	5
<i>Proteus species</i>	4
<i>Klebsiella aerogenes</i>	3
<i>Moraxella catarrhalis</i>	2
<i>Burkholderia cephalica</i>	2
<i>Citrobacter species</i>	2
Others**	2

\*Others included one *Streptococcus pyogenes*, one *Streptococcus mitis*, one *Streptococcus agalactiae*, one *Streptococcus angiosus*, and one *Listeria monocytogenes*.

\*\*Others included one *Klebsiella oxytoca* and one *Raoultella planticola*.

# Initial Management of HAP/VAP in Practice

## ATS/IDSA Guideline, 2016

Suspected HAP/VAP

- Presence of **new or progressive infiltrate**
- At least 2 of **fever, increase in WBC count, and purulence**



**Obtain lower respiratory tract specimen** using noninvasive sampling for culture

- Antibiotic regimen can be adjusted on the basis of those results
- Opportunity to de-escalate antibiotic coverage based on the results



**Empiric treatment regimens** informed by the **local distribution of pathogens** associated with HAP/VAP and their antimicrobial susceptibilities

# Rapid Diagnostic Tests

## Approved by US FDA



System	FilmArray (BioMérieux)	BD MAX (BD)	eSensor (GenMark)	Verigene (Luminex)	NxTAG (Luminex)	GeneXpert (Cepheid)
Method	Multiplex PCR with melt curve analysis	Real-time PCR	DNA hybridization	Microarray using nanoparticle probes	PCR with liquid phase bead array	On-demand PCR
Turnaround time	~ 1 h	~ 2 h	~ 6 hr	~ 2.5 h	< 4 hr	0.5-3 h
CLIA complexity	Moderate	Moderate	High	Moderate	High	Moderate/ waived
Panels	Resp, BC, GI, meningitis	MRSA, C. diff, GI, STD	Resp, BC	Resp, BC, GI	Resp	MRSA, CRE, VRE, C. diff

# BioFire® FilmArray® System



FILMARRAY



Sample Prep

**Simple:**  
Only 2 minutes of hands-on time



Amplification

**Easy:**  
No precise pipetting required



Detection

**Fast:**  
Run time of about 1 hour

# BioFire® FilmArray® Pneumonia *plus* Panel

Bacteria (semi quantitative)	Antibiotic Resistance Genes	Viruses
<ul style="list-style-type: none"> <li>• <b><i>Acinetobacter calcoaceticus-baumannii</i> complex</b></li> <li>• <i>Serratia marcescens</i></li> <li>• <i>Proteus</i> spp.</li> <li>• <b><i>Klebsiella pneumoniae</i> group</b></li> <li>• <b><i>Enterobacter aerogenes</i></b></li> <li>• <b><i>Enterobacter cloacae</i></b></li> <li>• <i>Escherichia coli</i></li> <li>• <i>Haemophilus influenzae</i></li> <li>• <i>Moraxella catarrhalis</i></li> <li>• <b><i>Pseudomonas aeruginosa</i></b></li> <li>• <b><i>Staphylococcus aureus</i></b></li> <li>• <i>Streptococcus pneumoniae</i></li> <li>• <i>Klebsiella oxytoca</i></li> <li>• <i>Streptococcus pyogenes</i></li> <li>• <i>Streptococcus agalactiae</i></li> <li>• <del><i>Stenotrophomonas maltophilia</i></del></li> </ul>	<p><b>ESBL</b> CTX-M</p> <p><b>Carbapenemases</b> KPC NDM Oxa48-like <del>Oxa23</del> <del>Oxa51</del></p> <p>VIM IMP</p> <p><b>Methicilin Resistance</b> mecA/mecC and MREJ</p>	<ul style="list-style-type: none"> <li>• Influenza A</li> <li>• Influenza B</li> <li>• Adenovirus</li> <li>• Coronavirus</li> <li>• Parainfluenza virus</li> <li>• Respiratory Syncytial virus</li> <li>• Human Rhinovirus/Enterovirus</li> <li>• Human Metapneumovirus</li> <li>• Middle East Respiratory Syndrome Coronavirus (MERS-CoV)</li> </ul>

# Diagnostic Performance

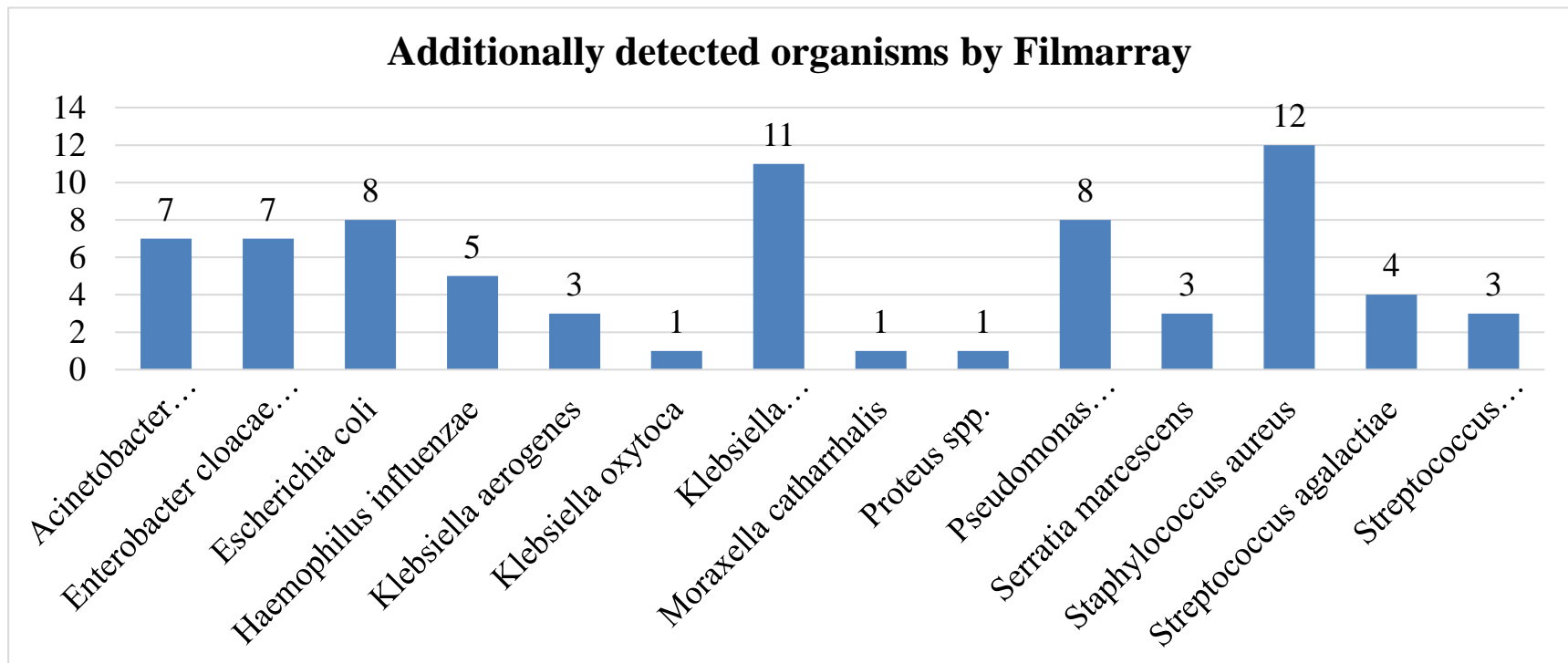
## SMC Experience



Evaluation of the BioFire FilmArray Pneumonia Panel for rapid detection of respiratory bacterial pathogens and antibiotic resistance genes in sputum and endotracheal aspirate specimens

In Young Yoo<sup>a,1</sup>, Kyungmin Huh<sup>b,c,1</sup>, Hyang Jin Shim<sup>d</sup>, Sun Ae Yun<sup>d</sup>, Yoo Na Chung<sup>a</sup>, On Kyun Kang<sup>a</sup>, Hee Jae Huh<sup>a,\*</sup>, Nam Yong Lee<sup>a,\*</sup>

International Journal of Infectious Diseases 95 (2020) 326–331



# Potential Tx Changes by FilmArray (n=46)

## SMC Experience

Potential treatment changes based on FilmArray Pneumonia Panel *plus* for 46 cases of suspected pneumonia.

Treatment modification	Cases, number (%)
<b>No modification</b>	<b>23 (50.0%)</b>
<b>Treatment changed</b>	<b>23 (50.0%)</b>
<b>Antibiotic escalation</b>	<b>13 (28.3%)</b>
Vancomycin	3
Carbapenem	2
Agents covering CRGNB <sup>a</sup>	8
<b>Antibiotic de-escalation</b>	<b>10 (21.7%)</b>
Vancomycin	3
Meropenem	3
Piperacillin/tazobactam	2
Vancomycin and meropenem	1
Vancomycin and amikacin	1

<sup>a</sup> Carbapenem-resistant gram-negative bacilli.

# Multicenter Evaluation of BioFire FilmArray French FA-PP Study



## A B S T R A C T

*Objectives:* To evaluate performances of the rapid multiplex PCR assay BioFire FilmArray Pneumonia Panel (FA-PP) for detection of bacterial pathogens and antibiotic resistance genes in sputum, endotracheal aspirate (ETA) and bronchoalveolar lavage (BAL) specimens.

*Methods:* This prospective observational study was conducted in 11 French university hospitals (July to December 2018) and assessed performance of FA-PP by comparison with routine conventional methods.

*Results:* A total of 515 respiratory specimens were studied, including 58 sputa, 217 ETA and 240 BAL. The FA-PP detected at least one pathogen in 384 specimens, yielding an overall positivity rate of 74.6% (384/515). Of them, 353 (68.5%) specimens were positive for typical bacteria while eight atypical bacteria and 42 resistance genes were found. While identifying most bacterial pathogens isolated by culture (374/396, 94.4%), the FA-PP detected 294 additional species in 37.7% (194/515) of specimens. The FA-PP demonstrated positive percentage agreement and negative percentage agreement values of 94.4% (95% CI 91.7%–96.5%) and 96.0% (95% CI 95.5%–96.4%), respectively, when compared with culture. Of FA-PP false-negative results, 67.6% (46/68) corresponded to bacterial species not included in the panel. At the same semi-quantification level (in DNA copies/mL for FA-PP versus in CFU/mL for culture), the concordance rate was 43.4% (142/327) for culture-positive specimens with FA-PP reporting higher semi-quantification of  $\geq 1 \log_{10}$  in 48.6% (159/327) of cases. Interestingly, 90.1% of detected bacteria with  $\geq 10^6$  DNA copies/mL grew significantly in culture.

*Conclusions:* FA-PP is a simple and rapid molecular test that could complement routine conventional methods for improvement of diagnosis accuracy of pneumonia. **Nabil Gastli, Clin Microbiol Infect 2020;■:1**

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# Unyvero LRT Panel, Curetis



FDA-cleared Unyvero uniquely and accurately detects the most clinically relevant pathogens and antibiotic resistance markers associated with pneumonia.

BACTERIA		RESISTANCE	GENES	
<i>Acinetobacter</i> spp.	<i>Moraxella catarrhalis</i>	Carbapenems	<i>kpc</i>	<i>oxa-48</i>
<i>Chlamydia pneumoniae</i>	<i>Morganella morganii</i>		<i>ndm</i>	<i>oxa-58</i>
<i>Citrobacter freundii</i>	<i>Mycoplasma pneumoniae</i>		<b><i>oxa-23</i></b>	<i>vim</i>
<i>Enterobacter cloacae</i> complex	<i>Proteus</i> spp.		<i>oxa-24</i>	
<i>Escherichia coli</i>	<i>Pseudomonas aeruginosa</i>	3rd Generation Cephalosporins		
<i>Haemophilus influenzae</i>	<i>Serratia marcescens</i>		<i>ctx-M</i>	
<i>Klebsiella oxytoca</i>	<i>Staphylococcus aureus</i>	Oxacillin/Cefoxitin		
<i>Klebsiella pneumoniae</i>	<b><i>Stenotrophomonas maltophilia</i></b>		<i>mecA</i>	
<i>Klebsiella variicola</i>	<i>Streptococcus pneumoniae</i>	Penicillin		
<i>Legionella pneumophila</i>			<i>tem</i>	
FUNGI				
<i>Pneumocystis jirovecii</i> *				



\* included on the Unyvero LRT BAL panel.

- Rapid, sample to answer direct from native specimen
- Simple and clear qualitative results based on quantitative algorithms
- Critical information for life-saving treatment decisions

## Specimen Types:

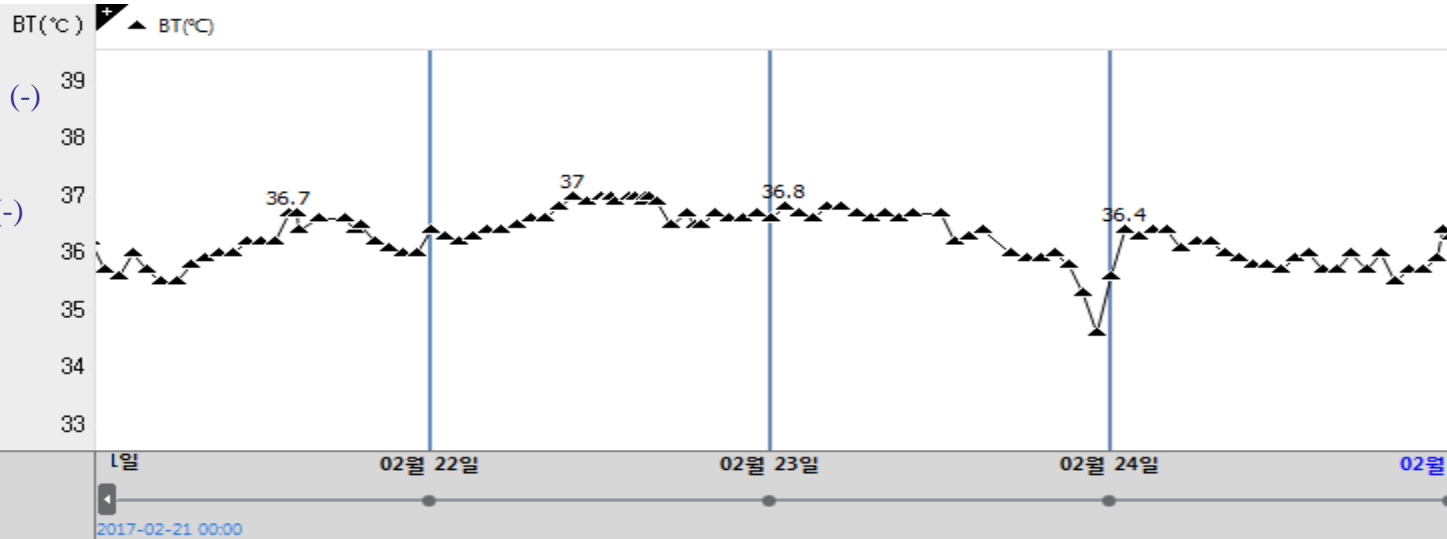
- Endotracheal Aspirate
- Bronchoalveolar Lavage (including mini-BAL)

# 66 Male with AGC

## Empiric Therapy with Cefepime 2g q 12hr



- Prior intravenous antibiotic (-)
- Septic shock (-)
- ARDS (-)
- ≥ 5 days of hospitalization (-)
- RRT (-)



그룹명	항목명	02월 21일	02월 22일	02월 23일	02월 24일
<b>▶ Vital Sign</b>					
BT	BT(°C)	36.7	37	36.8	36.4
<b>▶ 검사결과</b>					
WBC Count, Blood (x10 <sup>3</sup> /μL)		21.29	14.38	8.89	10.61
CRP, Quantitative (High Sensitivity) (mg/L)		13.54	11.97	13.99	16.08
Procalcitonin, quantitative (ng/mL)		6.48			
<b>▶ 항생제</b>					
Cefepime(q)		<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>
Vancomycin HCl(q)				<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>
Levofloxacin(mg)				<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>
Cefotetan(q)					<input checked="" type="checkbox"/>
<b>▶ 미생물검사</b>					
Gram Stain and Cultur...	Nasal-MRSA (Endo)Tracheal Aspir	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>		
Funus C...	(Endo)Tracheal Aspir		<input checked="" type="checkbox"/>		

# Dosing of Cefepime in Adults

## Clinical Drug Information, Lexicomp®



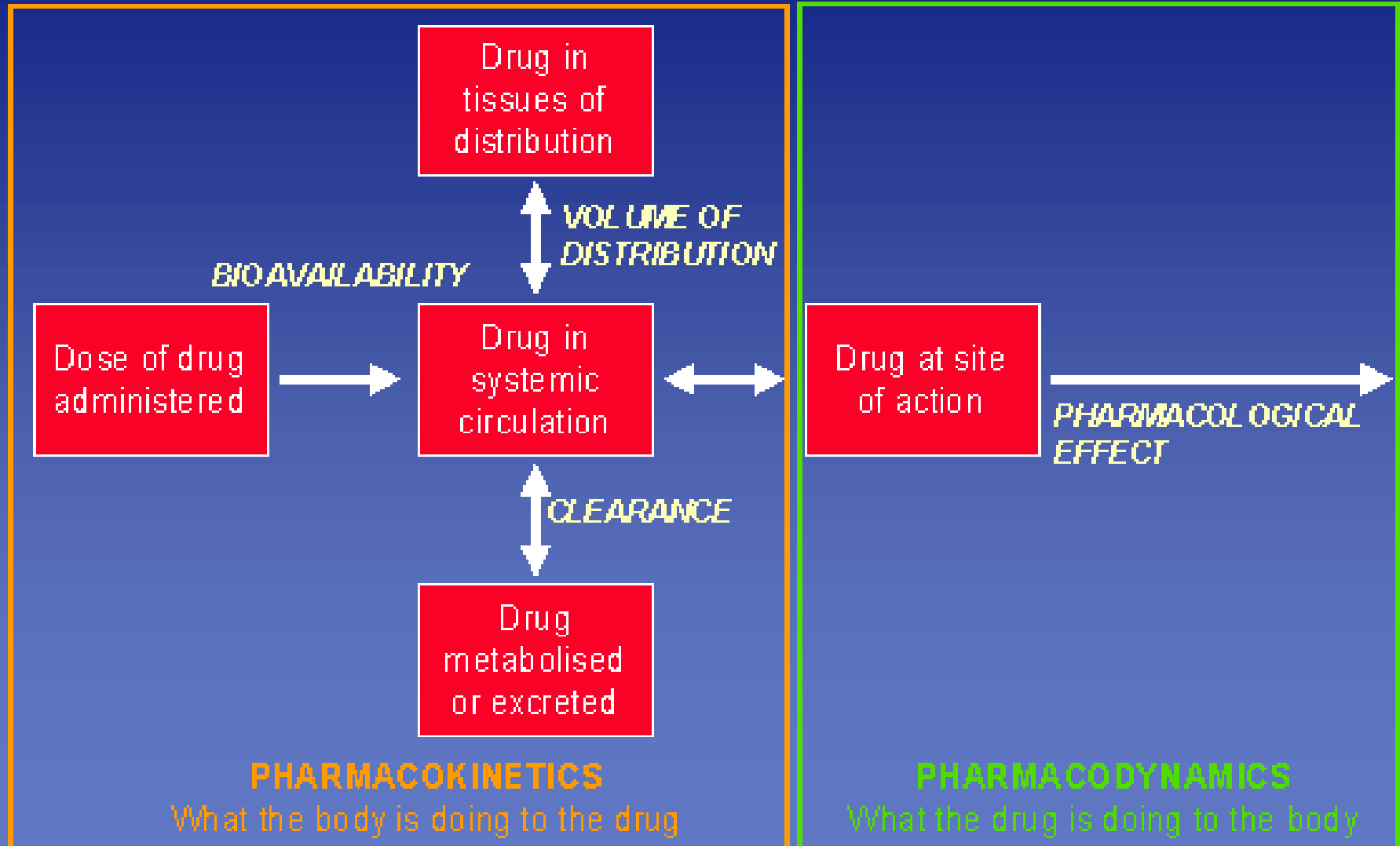
- **Pneumonia: IV:**
- *Manufacturer's labeling:*
  - Due to *P. aeruginosa*: 2 g every 8 hours for 10 days
  - Not due to *P. aeruginosa*: 1 to 2 g every 8 to 12 hours for 10 days
- *Alternate dosing:* Hospital-acquired or ventilator-associated: **2 g every 8 hours** for 7 days; may consider shorter or longer duration depending on rate of clinical improvement. Administration as an extended infusion may be considered. When used as empiric therapy, use in combination with an agent active against MRSA (unless coverage of MSSA only is appropriate) with or without an additional antipseudomonal agent (dependent on patient and institution-specific risk factors) (Kalil 2016)

# Initial Empiric Antibiotic Therapy for VAP

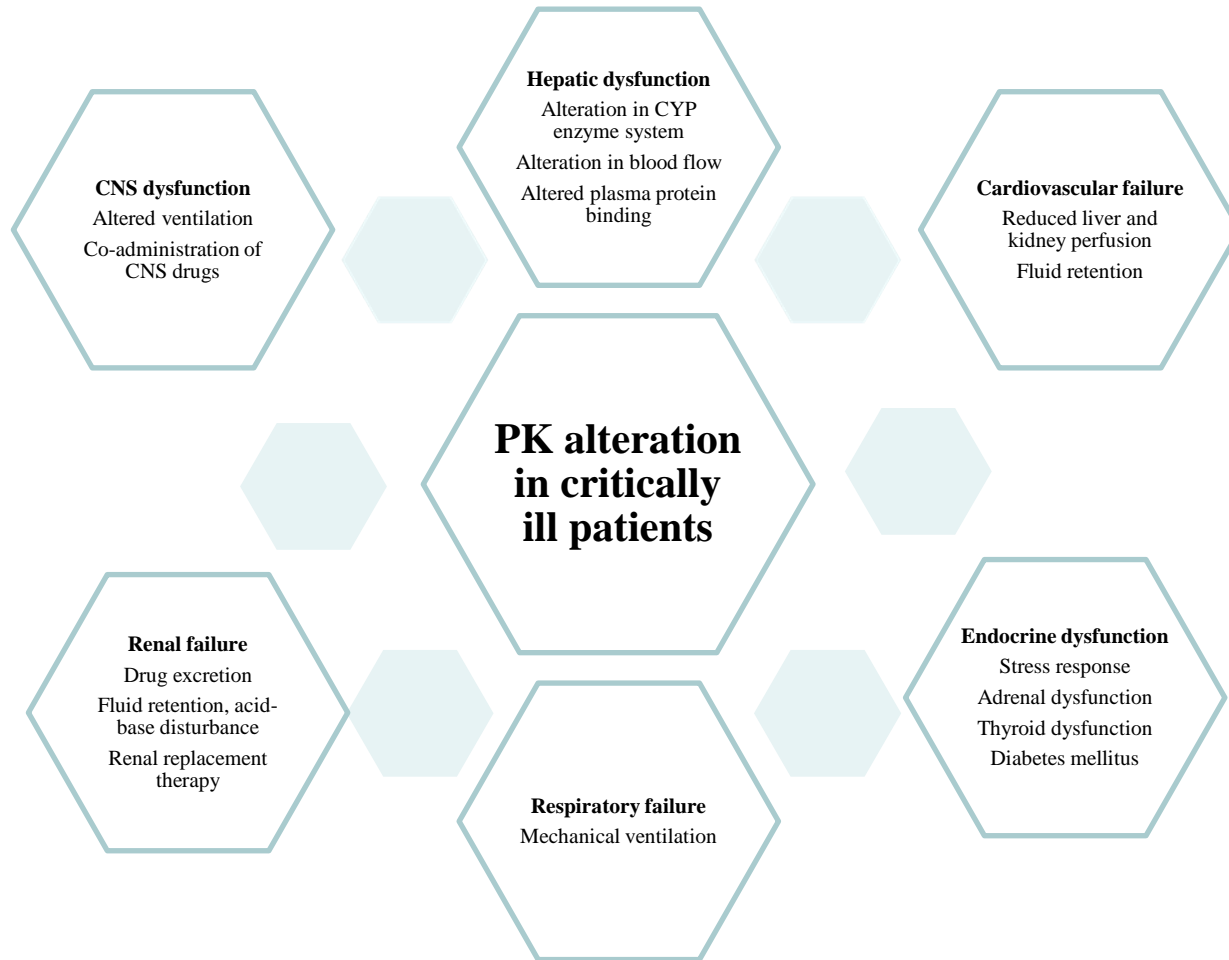
## Antibiotic Dosing

A. Gram-Positive Antibiotics With MRSA Activity	B. Gram-Negative Antibiotics With Antipseudomonal Activity: $\beta$ -Lactam-Based Agents	C. Gram-Negative Antibiotics With Antipseudomonal Activity: Non- $\beta$ -Lactam-Based Agents
Vancomycin 15 mg/kg IV q8–12h (consider a loading dose of 25–30 mg/kg $\times$ 1 for severe illness)	Piperacillin-tazobactam 4.5 g IV q6h	Ciprofloxacin 400 mg IV q8h Levofloxacin 750 mg IV q24h
OR	OR	OR
Linezolid 600 mg IV q12h	Cefepime 2 g IV q8h Ceftazidime 2 g IV q8h	Amikacin 15–20 mg/kg IV q24h Gentamicin 5–7 mg/kg IV q24h Tobramycin 5–7 mg/kg IV q24h
	OR	OR
	Imipenem 500 mg IV q6h Meropenem 1 g IV q8h	Colistin 5 mg/kg IV $\times$ 1 (loading dose) followed by 2.5 mg $\times$ (1.5 $\times$ CrCl + 30) IV q12h (maintenance dose) Polymyxin B 2.5–3.0 mg/kg/d divided in 2 daily IV doses
	OR	
	Aztreonam 2 g IV q8h	
Choose one gram-positive option from column A, one gram-negative option from column B, and one gram-negative option from column C. Note that the initial doses suggested in this table may need to be modified for patients with hepatic or renal dysfunction.		

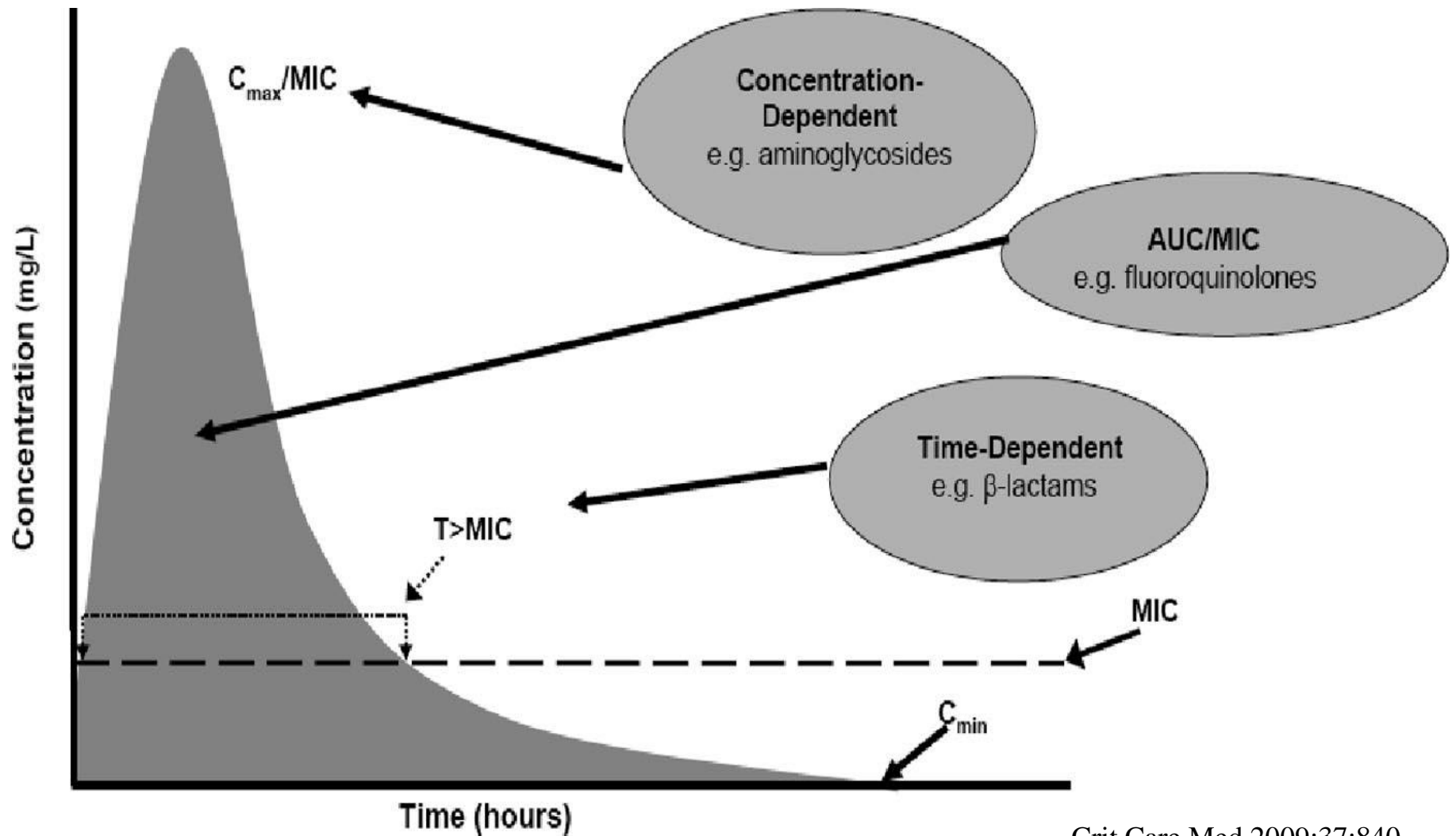
# Pharmacokinetic/Pharmacodynamic Optimization



# PK Alteration in Critically Ill Patients



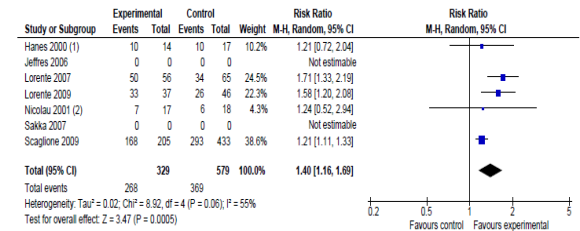
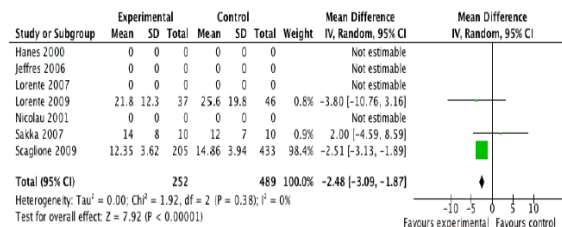
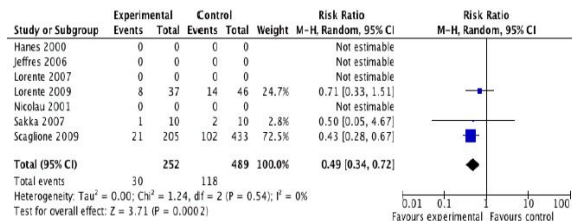
# PK Profile of Antibiotics



# Antibiotic Dosing Determined by PK/PD Data

## ATS/IDSA Guideline, 2016

- PK/PD optimized dosing refers to the use of
  - Antibiotic blood concentrations
  - Extended and continuous infusions
  - Weight-based dosing



► Decreased mortality, decreased ICU length of stay, and increased clinical cure rate

Clin Infect Dis 2016;63(5):e61

\* No published studies describing the PK/PD of piperacillin/tazobactam or polymyxins (colistin or polymyxin B) in patients with HAP/VAP are available at this time and so these drugs were not included in this recommendation.

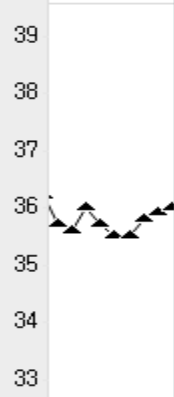
# 66 Male with AGC

## Isolation of MRSA



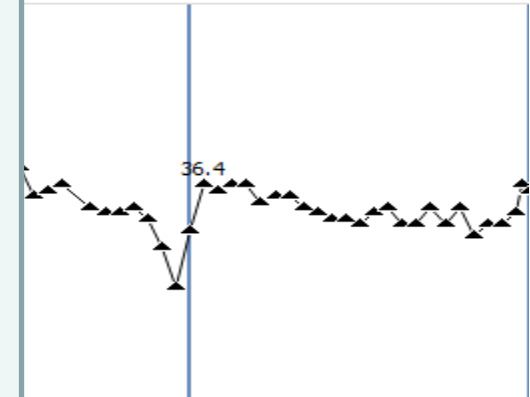
- Prior intravenous antibiotic (-)
- Septic shock (-)
- ARDS (-)
- ≥ 5 days of hospitalization (-)
- RRT (-)

BT(°C) ▲ BT(°C)



Isolate : #01  
Organism : Staphylococcus aureus, moderate.

Antibiotics	MIC	Susceptibility
Cefoxitine screen	Pos	+
Penicillin-G	>=0.5	R
Oxacillin	>=4	R
Gentamicin	<=0.5	S
Habekacin	<=1	S
Ciprofloxacin	<=0.5	S
Inducible Clindamycin Resistance	Pos	+
Erythromycin	>=8	R
Telithromycin	<=0.25	S
Clindamycin	<=0.25	R
Quinupristin/Dalfopristin	<=0.25	S
Linezolid	2	S
Teicoplanin	<=0.5	S
<b>Vancomycin</b>	<b>&lt;=0.5</b>	<b>S</b>
Tetracycline	<=1	S
Tigecycline	<=0.12	S
Nitrofurantoin	<=16	S
Fusidic acid	<=0.5	S
Mupirocin	<=2	S
Rifampin	<=0.5	S
Trimethoprim/Sulfa	<=10	S



그룹명	항목명	1일
2017-02-21 00		
Vital Sign		
BT	BT(°C)	
검사결과		
WBC Count, Blood (x10 <sup>3</sup> /μL)		
CRP, Quantitative (High Sensitivity) (mg/dL)		
Procalcitonin, quantitative (ng/mL)		
항생제		
Cefepime(q)		
Vancomycin HCl(q)		
Levofloxacin(mg)		
Cefotetan(q)		
미생물검사		
Gram Stain and Cultur...	Nasal-MRSA (Endo)Tracheal Aspir	
Funus C...	(Endo)Tracheal Aspir	

02월 24일	02월
36.4	36.4
39	10.61
99	16.08

# Vancomycin for HAP/VAP

## Optimal Target Serum Concentration



- Optimal target serum concentration for complicated MRSA infection: trough serum vancomycin concentrations of 15–20 mg/L
  - Bacteremia
  - Endocarditis
  - Osteomyelitis
  - Meningitis
  - Hospital-acquired pneumonia
  - ▶ Achieve an AUC/MIC of >400 for most patients if the MIC is <1 mg/L
  - ▶ To achieve rapid attainment of this target concentration for seriously ill patients, a loading dose of 25–30 mg/kg (based on actual body weight) can be considered.

Am J Health-Syst Pharm 2009; 66:82

# Vancomycin Loading Dose in HAP/VAP

## SMC Experience



**Table 2** Vancomycin dose and  $C_{trough}$  measurement between loading dose (LD) and non-LD groups

Characteristics	LD group (n=22)	Non-LD group (n=59)	P value
First dose of vancomycin			
Median dose per weight, mg/kg	24.4 (21.6–25.5)	15.2 (12.1–17.5)	<0.001
Groups by median dose per weight, mg/kg			<0.001
<15	0 (0.0)	29 (49.2)	
15–20	2 (9.1)	23 (39.0)	
20–25	14 (63.6)	7 (11.9)	
25–30	5 (22.7)	0 (0.0)	
>30	1 (4.5)	0 (0.0)	

**Table 3** Proportion of pharmacokinetic target (trough concentration >15 mg/L) attainment between vancomycin loading dose (LD) and non-LD groups

Characteristics	LD group (n=22)	Non-LD group (n=59)	OR (95% CI)	P value
Time to target attainment, h	54 (24–110)	60 (24–93)	NA	0.981
Target attainment within 24 h	6 (28.6)	17 (30.9)	0.89 (0.30–2.70)	0.843
Target attainment within 48 h	10 (47.6)	22 (40.0)	1.36 (0.50–3.75)	0.547

Values are median with interquartile range or n (%). OR, odds ratio; CI, confidence interval

# Effect of Vancomycin Loading Dose

## SMC Experience



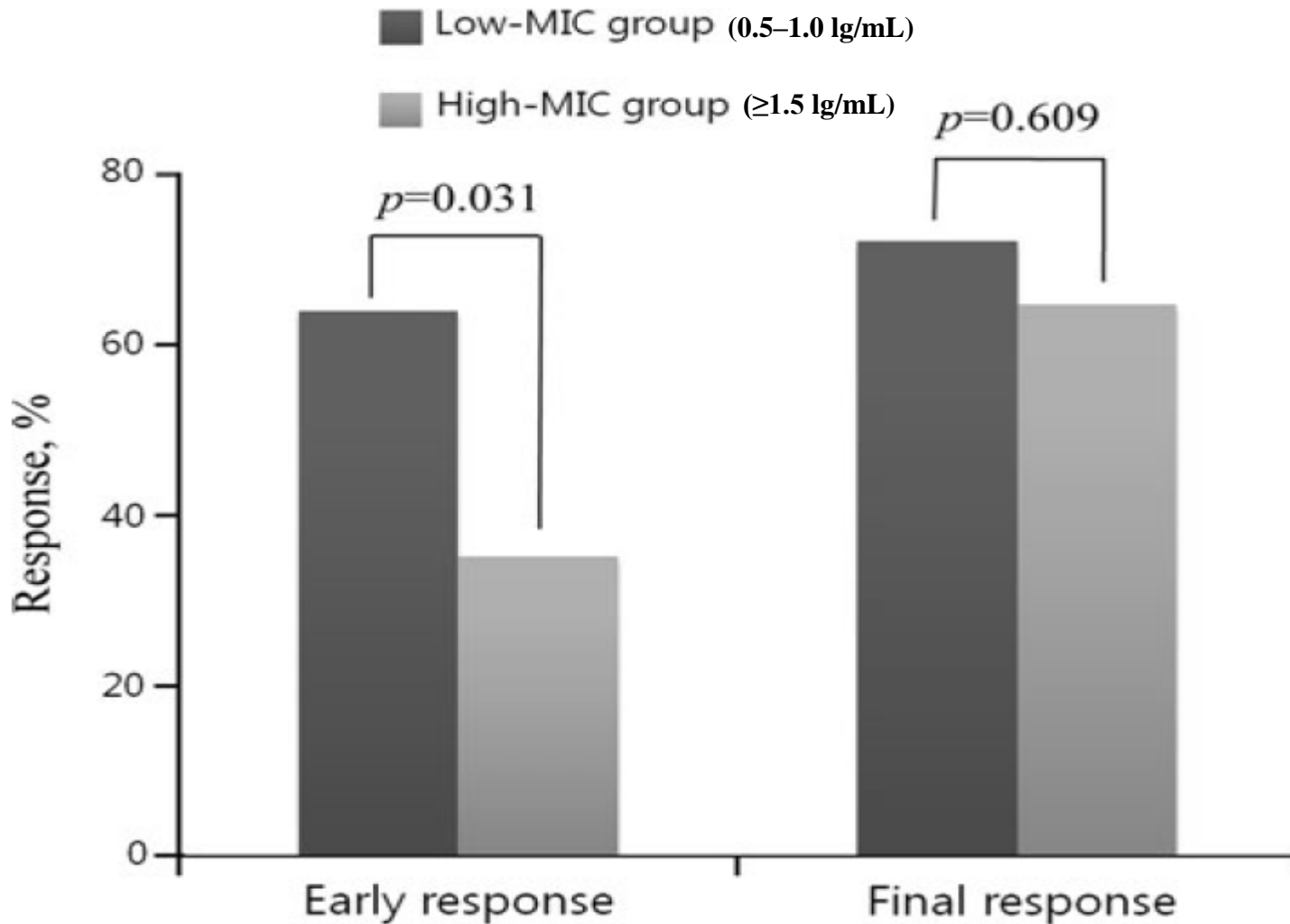
**Table 5** Univariable and multivariable analyses with logistic regression model for probability of clinical cure

Variables	Univariable		Multivariable	
	OR (95% CI)	P value	Adjusted OR (95% CI)	P value
Age, per year	N/A	0.486	0.97 (0.93–1.01)	0.109
Sex, female	1.20 (0.43–3.40)	0.727	1.52 (0.47–4.94)	0.484
Vancomycin LD	1.10 (0.39–3.13)	0.860	0.91 (0.27–2.99)	0.870
Target attainment in 48 h	0.83 (0.30–2.24)	0.706		
Cardiovascular disease	2.80 (0.73–10.76)	0.123	4.79 (0.99–23.17)	0.051
Mechanical ventilation	1.00 (0.87–11.54)	>0.999		
CRRT	0.33 (0.12–0.93)	0.031	0.31 (0.09–1.04)	0.058
Status of post-CPR	0.48 (0.06–3.61)	0.597		
ECMO	0.46 (0.11–2.00)	0.431		
<b>Vancomycin MIC</b>				
≤1	Reference		Reference	
2	0.21 (0.05–0.90)	0.054	0.19 (0.04–1.00)	0.050
Bacteremia	0.28 (0.07–1.10)	0.076	0.31 (0.07–1.40)	0.128

OR, odds ratio; CI, confidence interval; LD, loading dose; CRRT, continuous renal replacement therapy; CPR, cardiopulmonary resuscitation; ECMO, extracorporeal membrane oxygenation; MIC, minimum inhibitory concentration

# MIC of Vancomycin and Clinical Outcome

## AMC Data



# Vancomycin: We Can't Get There From Here

Nimish Patel,<sup>1</sup> Manjunath P. Pai,<sup>1</sup> Keith A. Rodvold,<sup>5</sup> Ben Lomaestro,<sup>3,4</sup> George L. Drusano,<sup>2</sup> and Thomas P. Lodise<sup>1,2</sup>

<sup>1</sup>Albany College of Pharmacy and Health Sciences, <sup>2</sup>Ordway Research Institute, <sup>3</sup>Albany Medical Center Hospital; <sup>4</sup>Albany Medical College, Albany, New York; and <sup>5</sup>University of Illinois at Chicago, Chicago, Illinois

MIC value	AUC/MIC ratio $\geq 400$			Nephrotoxic event	
	0.5mg/L (%)	1.0mg/L (%)	2.0mg/L (%)	Non-ICU (%)	ICU (%)
500 mg IV Q12H	57	15	0.7	3	10
1000 mg IV Q12H	90	57	15	6	16
1500 mg IV Q12H	97	79	38	9	25
2000 mg IV Q12H	98	90	57	14	34

# New Guideline for TDM of Vancomycin, 2020

Clinical Infectious Diseases

IDSA FEATURES

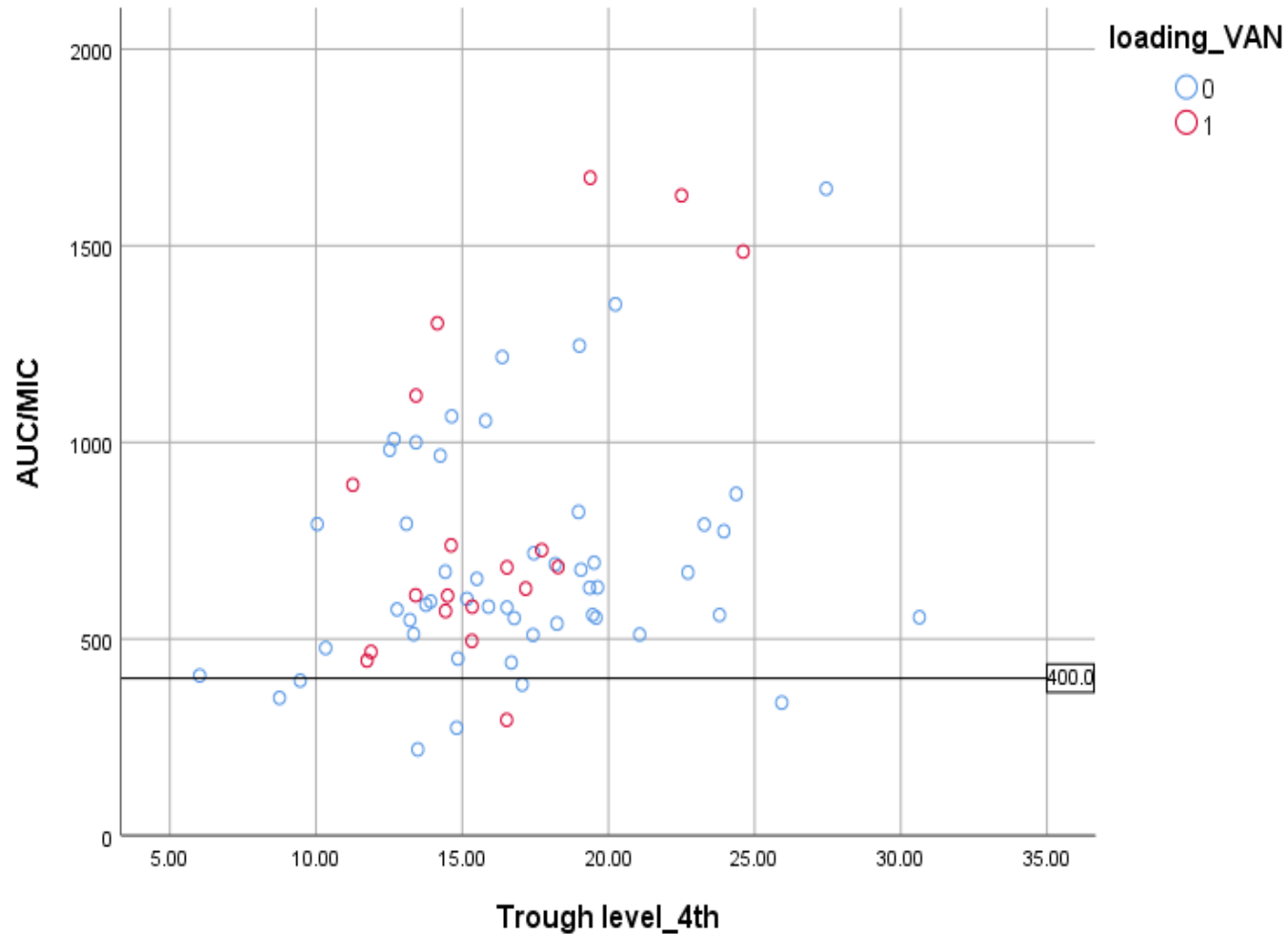


## Therapeutic Monitoring of Vancomycin for Serious Methicillin-resistant *Staphylococcus aureus* Infections: A Revised Consensus Guideline and Review by the American Society of Health-system Pharmacists, the Infectious Diseases Society of America, the Pediatric Infectious Diseases Society, and the Society of Infectious Diseases Pharmacists

1. In patients with suspected or definitive serious MRSA infections, an individualized target of the AUC/MIC<sub>BMD</sub> ratio of 400 to 600 (assuming a vancomycin MIC<sub>BMD</sub> of 1 mg/L) should be advocated to achieve clinical efficacy while improving patient safety (**A-II**).
3. Trough-only monitoring, with target between 15 and 20 mg/L, is no longer recommended based on efficacy and nephrotoxicity data in patients with serious infections due to MRSA (**A-II**). There is insufficient evidence to provide recommendations on whether trough-only or AUC-guided vancomycin monitoring should be used among patients with noninvasive MRSA or other infections.

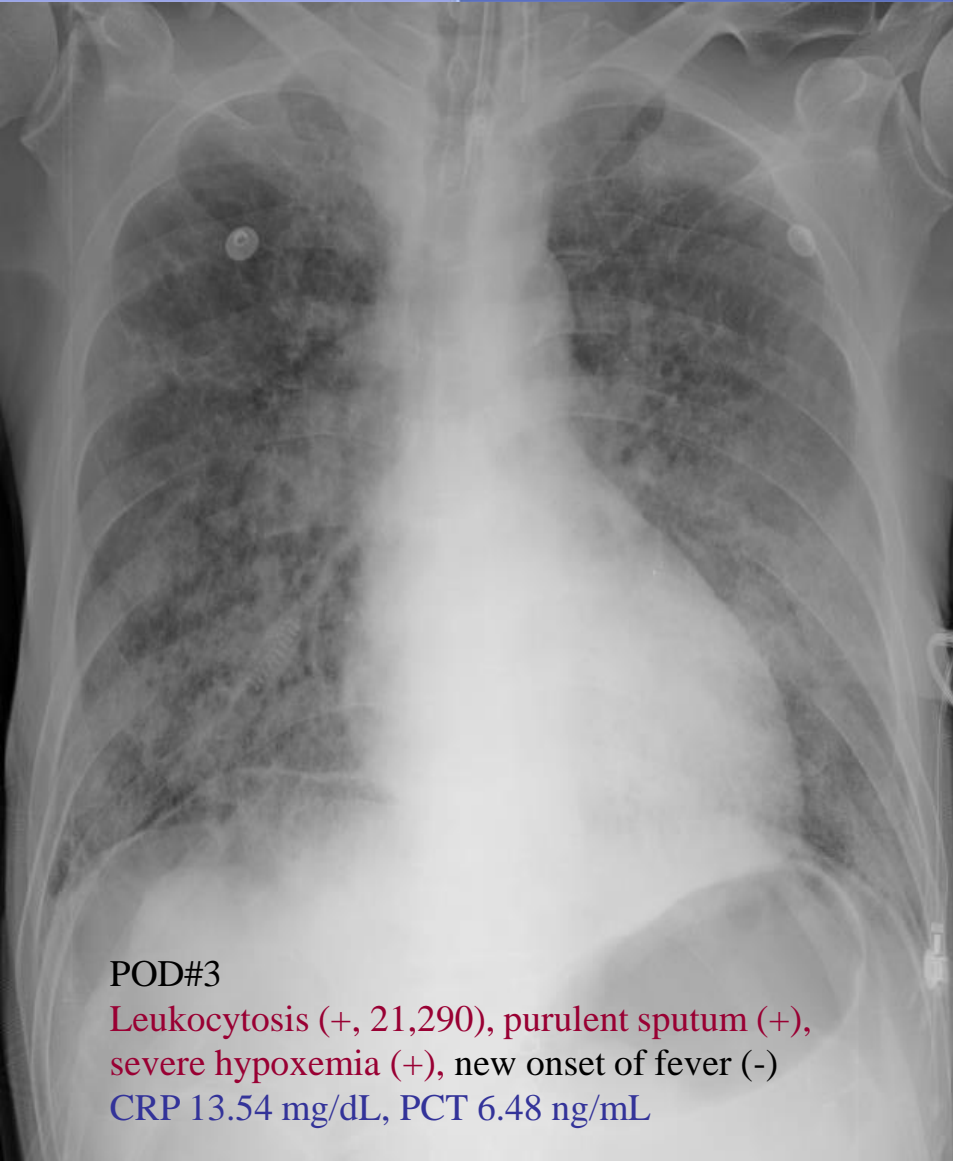
# AUC/MIC Based on Trough 15–20 mg/L

## SMC Experience



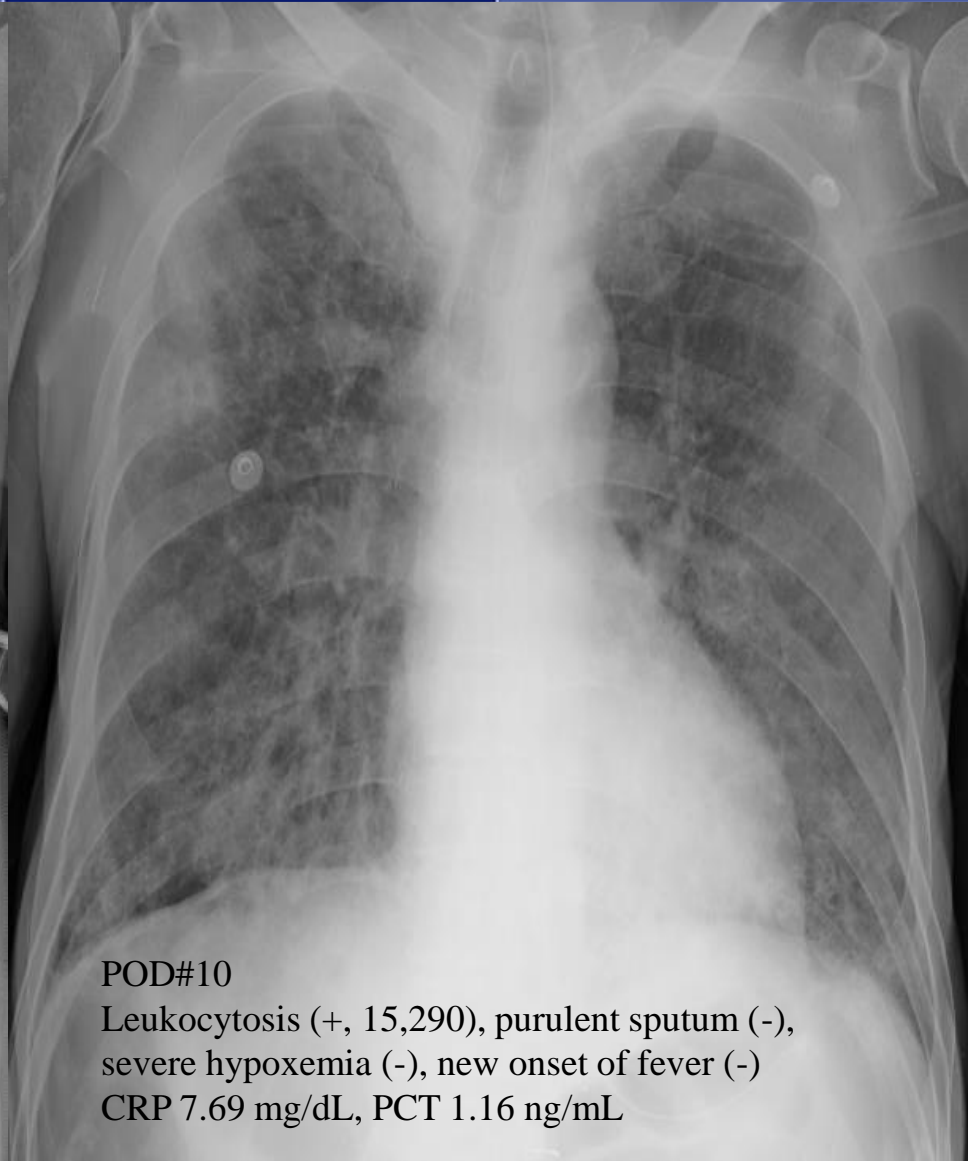
# 66 Male with AGC

## Vancomycin for 7 days



POD#3

Leukocytosis (+, 21,290), purulent sputum (+),  
severe hypoxemia (+), new onset of fever (-)  
CRP 13.54 mg/dL, PCT 6.48 ng/mL



POD#10

Leukocytosis (+, 15,290), purulent sputum (-),  
severe hypoxemia (-), new onset of fever (-)  
CRP 7.69 mg/dL, PCT 1.16 ng/mL

# Carbapenem Resistant GNB



*Acinetobacter baumannii*



*Pseudomonas aeruginosa*



*Klebsiella pneumoniae*



The key elements that define the threat of carbapenem-resistant gram-negative pathogens

- Increasing incidence of these pathogens worldwide since the turn of the century
- Lack of safe and efficacious agents for treatment once the efficacy of carbapenems is lost due to resistance
- High mortality rates associated with carbapenem-resistant gram-negative infections

# Which Antibiotic Should Be Used to Treat Patients With HAP/VAP Due to Carbapenem-Resistant Pathogens?

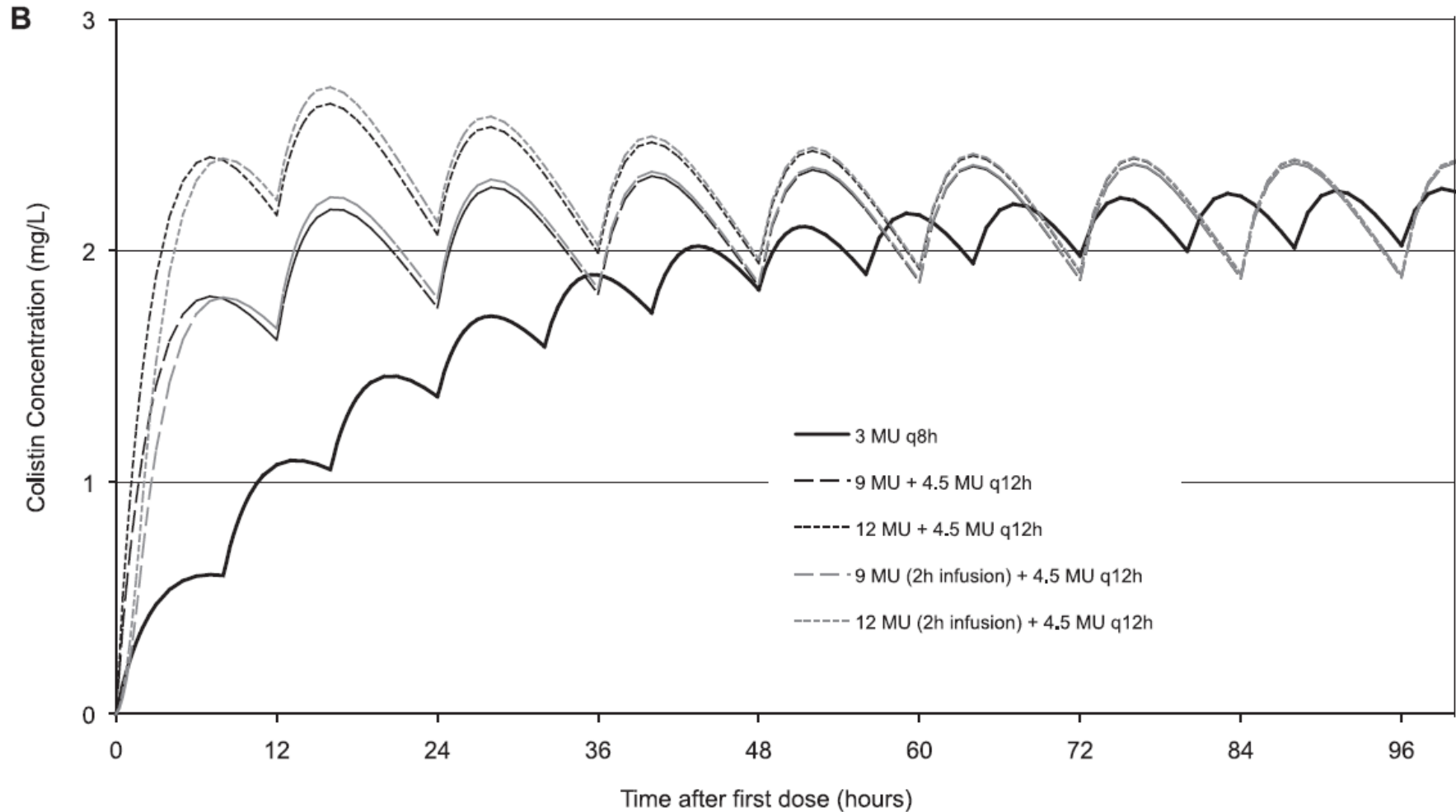
- Suggested drugs for carbapenem-resistant Gram-negative bacterial pneumonia
  - Backbone:  $\beta$ -Lactam
  - Accompanying drug: Fluoroquinolone
  - Alternative accompanying drug: Polymyxin or aminoglycoside
  - Other alternatives: Fosfomycin or tigecycline

Antimicrob Agents Chemother 2020;64:e02183

- IDSA/ATS Guideline, 2016
  - In patients with HAP/VAP caused by a carbapenem-resistant pathogen that is sensitive only to polymyxins, we recommend intravenous polymyxins (colistin or polymyxin B) (strong recommendation, moderate-quality evidence), and we suggest adjunctive inhaled colistin (weak recommendation, low-quality evidence).

Clin Infect Dis 2016;63(5):e61

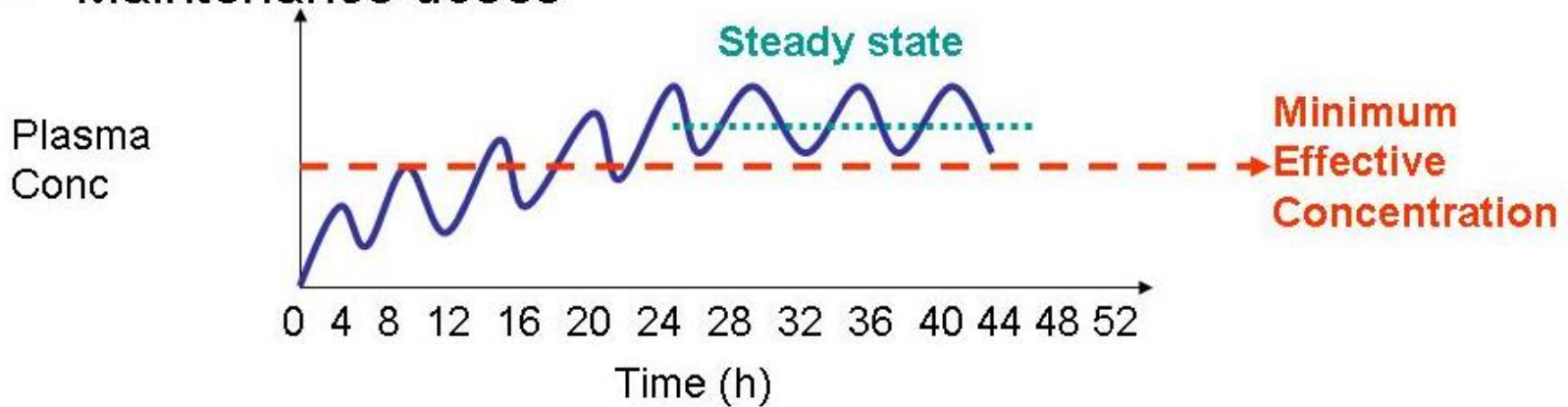
# Concentrations of Colistin by Different Dosing



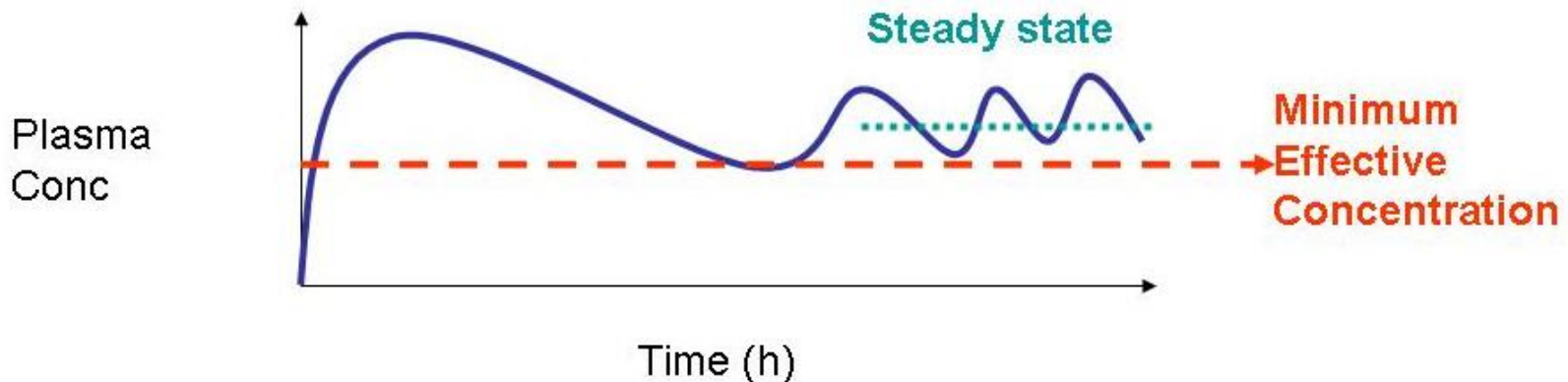
# Loading Dose of Colistin



- Maintenance doses



- Loading dose and Maintenance doses

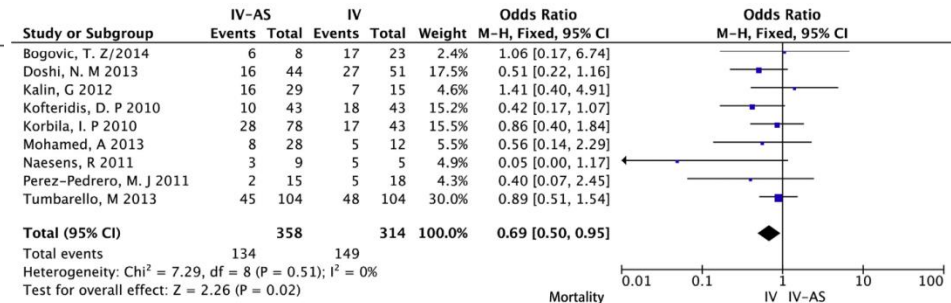
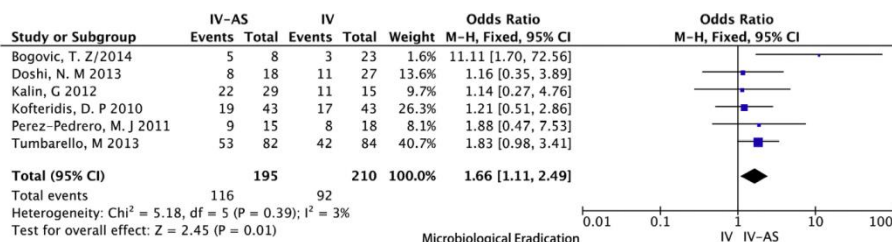
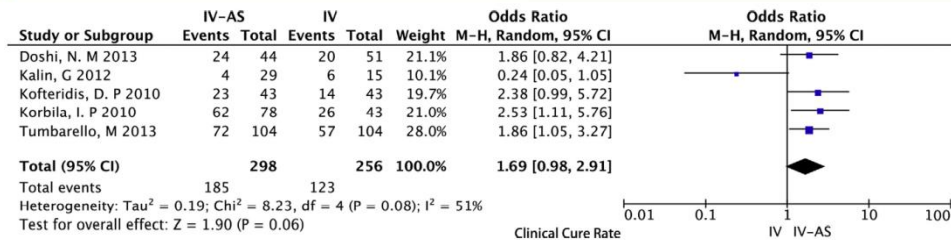
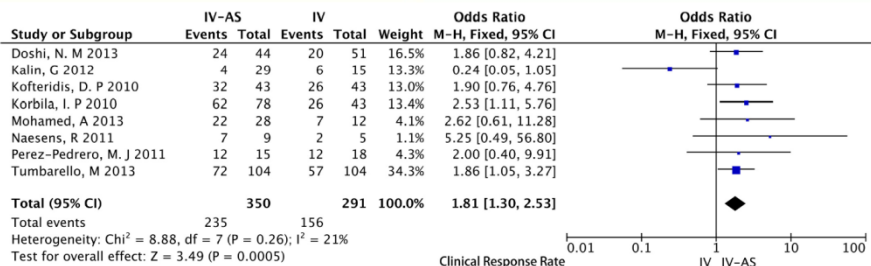


# New Dosing Guidance of IV Colistin



Dose	Category of Critically Ill Patient	Dosing Suggestions <sup>a</sup>
Loading dose	All patient categories	Equation 1: Loading dose of CBA (mg) = $C_{ss,avg}$ target (mg/L) × 2.0 × ideal body weight (kg) To achieve a $C_{ss,avg}$ of 2 mg/L in a patient with an ideal body weight of 75 kg, the loading dose would be 300 mg CBA (9 million IU), the suggested maximum loading dose. The 1st regular daily dose should be administered 12 h later.
Daily dose <sup>b</sup>	Not receiving RRT	Equation 2 <sup>c</sup> : Daily dose of CBA (mg) = $C_{ss,avg}$ target (mg/L) × $10^{(0.0048 \times CrCl + 1.825)}$ See <a href="#">Table 3</a> ("look-up" table) for the daily dose to target a plasma colistin $C_{ss,avg}$ of 2 mg/L, depending on the patient's creatinine clearance.
	Receiving RRT	The baseline daily dose of colistimethate for a $C_{ss,avg}$ of 2 mg/L in a patient with creatinine clearance of 0 mL/min is 130 mg/d of CBA (3.95 million IU/d) (see <a href="#">Table 3</a> ) <sup>d</sup> ; the supplement to the baseline daily dose needed during receipt of RRT is 10% of the baseline dose per 1 h of RRT.
	Intermittent hemodialysis	Nondialysis day: CBA dose of 130 mg/d (3.95 million IU/d), ie, baseline dosing for a $C_{ss,avg}$ of 2 mg/L; dialysis day supplement: add 30% or 40% to baseline daily dose after a 3- or 4-h session, respectively. <sup>e</sup> The dialysis session should occur toward the end of a colistimethate dosing interval, and the supplement to the baseline (nondialysis) daily dose should be administered with next regular dose, after the dialysis session has ended.
	SLED	During SLED: add 10% per 1 h of SLED replacement to baseline daily dose for a $C_{ss,avg}$ of 2 mg/L <sup>f</sup> ; for a patient receiving a 10-h nocturnal SLED session each day and receiving colistimethate every 12 h, the dose would be (baseline CBA dose of 130 mg/d for a patient with creatinine clearance of 0 mL/min + supplemental dose comprising 10% of the baseline dose per h × 10 h); ie, for this case the CBA dose would be 260 mg/d (7.9 million IU/d). It is suggested that the SLED session begin 1–2 h after the afternoon/evening dose; in such a case, it may be most convenient and safe to administer 130 mg CBA (3.95 million IU) every 12 h.
	CRRT	During CRRT: add 10% per 1 h of CRRT to the baseline daily dose for a $C_{ss,avg}$ of 2 mg/L <sup>g</sup> ; the suggested CBA dose is 440 mg/d (~13 million IU/d).

# Adjunctive Inhaled Colistin Meta-analysis



Int J Antimicrob Agents 2015;46:603

	Relative risk	95% CI
Mortality	0.84	0.63 – 1.12
Clinical cure	1.29	1.13 – 1.47
Nephrotoxicity	1.11	0.78 – 1.57

► The guidelines suggests both inhaled and systemic antibiotics, rather than systemic antibiotics alone

Clin Infect Dis 2016;63(5):e61

# IV Loading and Inhaled Colistin

## SMC Experience



**Table 3.** Clinical outcomes of 191 patients with HAP or VAP caused by CRGNB, who were treated with colistin in the ICU.

Variables	Non-LD IV (n=70)	LD IV (n=86)	AS-LD (n=35)	p value
Clinical response				
Clinical cure	32 (46)	36 (42)	17 (49)	0.764
Recurrence	4 (6)	10 (12)	4 (11)	0.414
Clinical failure	34 (49)	40 (47)	14 (40)	0.724
Microbiological response				
Eradication	21/67 (31)	27/81 (33)	21/35 (60)	0.010*‡
Recurrence	9/67 (13)	12/81 (15)	7/35 (20)	0.663
Colonization	20/67 (30)	22/81 (27)	3/35 (9)	0.047‡
Microbiological failure	17/67 (25)	20/81 (25)	4/35 (11)	0.222
Duration of ICU stay, days	13 (8–21)	12 (8–18)	20 (10–33)	0.013‡
Mortality				
30-day mortality	32 (46)	42 (49)	8 (23)	0.027‡
90-day mortality	41 (59)	50 (58)	16 (46)	0.396
Nephrotoxicity	27/50 (54)	23/61 (38)	16/27 (59)	0.100
Initiation rates of RRT	8/50 (16)	5/61 (8)	6/27 (22)	0.151

AS, aerosolized; CRGNB, carbapenem-resistant gram-negative bacteria; HAP, hospital-acquired pneumonia; ICU, intensive care unit; IV, intravenous; LD, loading dose; RRT, renal replacement therapy; VAP, ventilator-associated pneumonia.

\*, ‡, § indicate significant differences ( $p < 0.017$ ) between the non-LD IV group and the LD IV group, LD IV group and AS-LD group, and the non-LD IV group and the AS-LD group, respectively.

# Practical Management of HAP/VAP

## Summary



- For patients with suspected HAP/VAP,
  - Diagnose with **clinical criteria alone**
  - Microbiologic examination with **non-invasive sampling** (with semi-quantitative cultures) and **rapid diagnostic techniques**
  - Empiric antibiotics based on **the local distribution of pathogens and antimicrobial susceptibilities**.
    - In case where local antimicrobial susceptibility rates are not available
      - High risk of mortality
      - MDR risk: Prior intravenous antibiotic use within 90 d
  
- Pharmacokinetic/Pharmacodynamic optimization of antibiotic therapy
  - **Antibiotic dosing be determined using PK/PD data**, rather than the manufacturer's prescribing information
    - Accurate dosing and monitoring of drug level
    - Loading dose
    - Inhaled antibiotics