#### Joint Commission

EP 20 (revised): The antibiotic stewardship program collects, analyzes, and reports data to hospital leadership and prescribers.

Note: Examples of antibiotic stewardship program data include antibiotic resistance patterns, antibiotic prescribing practices, or an evaluation of antibiotic stewardship activities.

#### College of American Pathologists

#### MIC.21946 Cumulative Susceptibility Data

Phase 1

Requirement For hospital based microbiology laboratories, cumulative antimicrobial susceptibility test data are maintained and

reported to the medical staff at least yearly.

The minimum requirements: **Options:** Each hospital Whole hospital Once a year Unit type Reported to medical staff Unit specific

The minimum requirements:

**Options:** 

Each hospital

More frequently

Once a year

Rolling updates

Reported to medical staff

The minimum requirements:

Each hospital

Printed

Once a year

Electronic

Reported to medical staff

#### Antimicrobic % Interpretive Report

NORTON HEALTHCARE

2021 Audubon

Organism	(# of iso)																				
Organism	(# 01 150)	CFT/CA	CFTE	CFX	CFZ	CL	CN	CP	CP-S	CPD	CPE	CRM	CTN	CZ	CZA	DOR	ECAZ	ESA	ESB	ETP	FD
C. farmeri	(2) S	-		-	0%			100%	-		100%	0%	100%	-						100%	100
01,141111011	(_, _			-	0%			0%			0%	0%	0%							0%	09
	R				100%			0%			0%	100%	0%							0%	09
		0	0	0	2	0	0	2	0	0	2	1	1	0	0	0	0	0	0	2	1
C. freundii	(16) S			0%	0%			100%	-		100%	100%	92%		100%	100%				100%	93
	1			0%	0%			0%			0%	0%	8%		0%	0%				0%	7
	R			100%	100%			0%			0%	0%	0%		0%	0%				0%	0
		0	0	2	16	0	0	16	0	0	16	4	12	0	11	1	0	0	0	16	1
C. freundii cplx	(17) S			11%	0%			100%			100%	80%	86%		100%	100%				94%	10
	_!			11%	0%			0%			0%	0%	0%		0%	0%				6%	0
	R			78%	100%			0%			0%	20%	14%		0%	0%				0%	0
		0	0	9	17	0	0	17	0	0	17	10	7	0	9	9	0	0	0	17	8
C. indologenes	(2) S							100%			0%										-
	1							0%			0%							-			
	R							0%			100%										
		0	0	0	0	0	0	2	0	0	2	0	0	0	0	0	0	0	0	0	
C. koseri	(30) S			86%	100%			100%			100%	78%	100%		100%	100%				100%	50
	1			14%	0%			0%			0%	11%	0%		0%	0%				0%	45
	R		-	0%	0%			0%			0%	11%	0%	-	0%	0%	-			0%	5
		0	0	7	30	0	0	30	0	0	30	9	21	0	7	7	0	0	0	30	2

# Manual

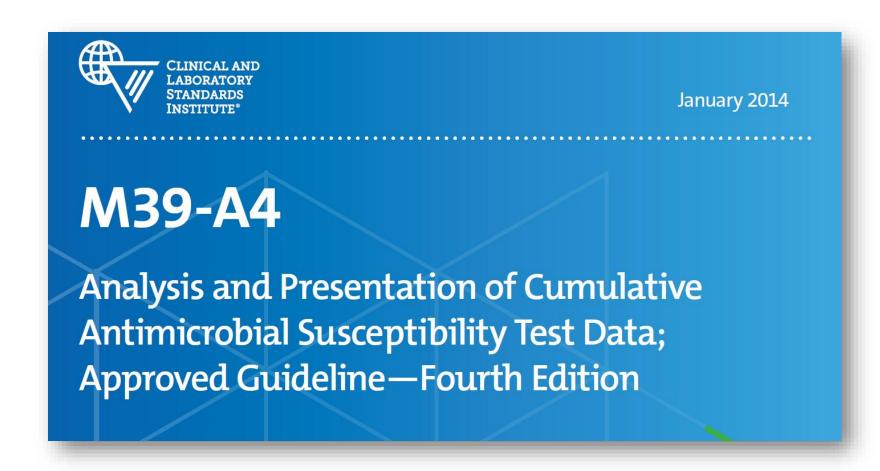


# Automated

#### The microbiology laboratory database software

WHONET is a free desktop Windows application for the management and analysis of microbiology laboratory data with a particular focus on antimicrobial resistance surveillance developed and supported by the WHO Collaborating Centre for Surveillance of Antimicrobial Resistance at the Brigham and Women's Hospital in Boston, Massachusetts. WHONET, available in 44 languages, supports local, national, regional, and global surveillance efforts in over 2,300 hospital, public health, animal health, and food laboratories in over 130 countries worldwide.

### The How-To Rules



## The How-To Rules

- Which isolates to include
  - Those collected from human specimens for diagnostic purposes
    - No surveillance cultures (MRSA, VRE, CRE, etc.)
    - No environmental cultures
    - No non-human critters
  - Obviously only those you do susceptibility testing on



### M39-A4

Analysis and Presentation of Cumulative Antimicrobial Susceptibility Test Data; Approved Guideline—Fourth Edition

 Only the first isolate of a given species per patient, per analysis period (eg, one year), irrespective of body site, antimicrobial susceptibility profile, or other phenotypic characteristics

Date of Collection	Isolate	Specimen Type	In the Antibiogram?
2/14/2021	Klebsiella pneumoniae	BAL	YES
2/15/2021	Klebsiella pneumoniae	Blood	NO
2/20/2021	Klebsiella pneumoniae	Sputum	NO
2/20/2021	Staphylococcus aureus	Sputum	YES
4/8/2021	Citrobacter freundii	Wound	YES
4/8/2021	Staphylococcus aureus	Wound	NO
7/7/2021	Klebsiella pneumoniae	Sputum	NO
12/31/2021	Staphylococcus aureus	Blood	NO
1/2/2022	Staphylococcus aureus	Blood	YES
1/5/2022	Staphylococcus aureus	Wound	NO

Mrs. Winter\*

In a laboratory that creates one antibiogram for each calendar year

What's your best source of information in deciding on an empiric therapy for these isolates?

Well, what about this one?

- Which antibiotics to include
  - Only those you routinely <u>test</u> on all isolates of this species, even if you don't routinely <u>report</u> them
  - None that are "supplemental" testing



### M39-A4

Analysis and Presentation of Cumulative Antimicrobial Susceptibility Test Data; Approved Guideline—Fourth Edition

 Only antibiotics that may be appropriate or recommended for empiric therapy

Drug Tested	Reporting Rules	%S of all isolates*	%S of reported isolates*
Cefazolin	Always reported	80%	80%
Ceftriaxone	Only reported if cefazolin NS	92%	50%
Ceftazidime/avibactam	Only reported if CRE	92%	66%

## The How-To Rules

- What to report in the antibiogram
  - Only those species with 30 or more isolates (okay, maybe 10)
  - Group similar organisms? Such as
    - Viridans streptococci
    - Coagulase-negative staphylococci
    - Citrobacter freundii complex
  - For those with lower numbers, include a longer time frame?



### M39-A4

Analysis and Presentation of Cumulative Antimicrobial Susceptibility Test Data; Approved Guideline—Fourth Edition

- Report as percent susceptible
  - That's fully susceptible, not I, not R
- For those antibiotics with susceptible dose-dependent interpretations, report that percentage separately

		Penicillins								ohalo	spori	ins	Monobactam	Carbap	arbapenems Aminoglycosides					G	ram	+ Co	vera	ge	Others					
Norton Hospital 2021	Number Tested	Amoxicillin/Clavulanate	Ampicillin	Ampicillin/Sulbactam	Oxacillin	Penicillin	Piperacillin/Tazobactam	Oral cephalosporins for uncomplicated UTI	Cefazolin	Cefepime	Ceftazidime	Ceftriaxone	Aztreonam	Ertapenem	Meropenem	Amikacin	Gentamicin	Gentamicin Synergy	Tobramycin	Clindamycin [1, 2]	Erythromycin [2]	Vancomycin	Linezolid	Daptomycin	Ciprofloxacin	Levofloxacin	Nitrofurantoin [2]	Tetracydine	Trimeth/Sulfa	
Acinetobacter baumannii complex [3]	20	0	0	75	0	0			0	65	80		0	0	75	85	75		85	0	0	0	0	0	85	75			75	
Citrobacter freundii complex [4]	29	0	0	0	0	0	93		0	97	83	79	83	97	97	100	90		93	0	0	0	0	0	90	93	89		83	
Citrobacter koseri	19	89	0	84	0	0	95		89	100	100	89	95	100	100		100		100	0	0	0	0	0	89	100			95	
Enterobacter cloacae complex [5]	89	0	0	0	0	0	79		0	88	71	56	65	82	97	100	99		98	0	0	0	0	0	97	100			94	
Escherichia coli	965	85	47	55	0	0	97	86	66	92	91	87	88	99	99	99	91		91	0	0	0	0	0	74	75	97		72	
Klebsiella aerogenes	41	0	0	0	0	0	88		0	98	85	80	83	95	100	100	100		100	0	0	0	0	0	95	95	18		98	
Klebsiella oxytoca	53	98	0	81	0	0	100		15	98	94	96	94	100	100		100		96	0	0	0	0	0	98	98	80		96	
Klebsiella pneumoniae	224	88	0	77	0	0	94	88	79	94	90	89	90	98	99	100	96		95	0	0	0	0	0	92	95	37		89	
Morganella morganii	17	0	0	0	0	0	100		0	94	53	53	71	100	100	100	94		94	0	0	0	0	0	88	94	0		94	
Proteus mirabilis	125	92	79	90	0	0	99	87	66	94	95	92	89	99	100	100	90		90	0	0	0	0	0	75	78	0	0	79	
Proteus vulgaris	12	92	0	92	0	0	100		0	100	92	50	33	100	100		100		100	0	0	0	0	0	100	100	0	0	100	
Pseudomonas aeruginosa	166	0	0	0	0	0	84		0	83	80	0	70	0	89	96	81		94	0	0	0	0	0	81	83	0	0	0	
Serratia marcescens	47	0	0	0	0	0	57		0	98	43	53	38	100	100	98	96		91	0	0	0	0	0	100	100	0		98	
Stenotrophomonas maltophilia	33	0	0	0	0	0	0		0		45	0	0	0	0	0	0		0	0	0	0	0	0		88		0	94	
Staphylococcus aureus	682	47	<del>                                     </del>	-	47				47				0							77	33	100	100	99	_	-	$\vdash$	91	98	
Methicillin-resistant S. aureus	391	0	$\vdash$		0				0		$\vdash$		0							74		100	100	99		$\vdash$	+	94	97	
Methicillin-resistant 3. aureus	331	100			100				100				0							81	56	100	100	100		<del>                                     </del>	$\vdash$	89	99	
Staphylococcus capitis	10	100	$\vdash$		100				100				0							*	*	100	100		$\vdash$	$\vdash$	*	100	100	
Staphylococcus epidermidis	126	37	$\vdash$	$\vdash$	37				37				0			$\vdash$				56	34	100	100	100	$\vdash$	$\vdash$	100	81	65	
Staphylococcus haemolyticus	15	27		$\vdash$	27				27				0							*	*	100	100	100		$\vdash$	*	93	60	
Staphylococcus lugdunensis	36	83		$\vdash$	83				83				0			$\vdash$				71	63	100	100	100		$\vdash$	*	92	97	
Other coagulase-negative staphylococci	25	50			50				50				0							46	40	100	100	100			*	80	84	
Enterococcus faecalis	119		99			99			0	0	0	0	0			0	0	71	0	0	26	91	100	99			*	33	0	
Enterococcus faecium	38		29			29			0	0	0	0	0			0	0	79	0	0	3	39	100	97		<del>                                     </del>	*	29	0	
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