

Antibiogram of Organisms Commonly Isolated from Outpatient Urinary Tract Specimens in Clark County, Nevada – January 2004 through March 2006

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Introduction

The increasing occurrence of antibiotic resistance (AR) has become an urgent public health problem. Although it has long been common in nosocomial infections, it is seen with increasing frequency in infections with no healthcare association. In an effort to understand this issue more clearly, the Southern Nevada Health District (SNHD) has been working with Quest Laboratories to examine the local occurrence of antibiotic resistance. This report will focus on the most common bacterial organisms causing outpatient urinary tract infections (UTI) in Clark County.

Methods

SNHD electronically receives culture and sensitivity results for all bacterial isolates processed by Quest laboratories in Clark County. This information is transmitted on a monthly basis. It includes the name of the organism isolated, its antibiotic sensitivity, patient identification information, requesting facility/physician and specimen source. Software programming and manual sorting through organism descriptions to assign consistent names were necessary to extract organisms associated with urinary tract infections from the Quest data. Organism names are not uniformly entered at the laboratory and as such the same organism may have as many as twenty or more descriptions. Therefore, preexisting knowledge of organisms most frequently associated with UTIs was applied and a broad spectrum of organisms was predefined so that AR data associated with those selected organisms could be filtered out from the rest of Quest data. This was done using keywords “urine” or “urethra” as filtering criteria in the source of culture or test name fields.

A computer program was also written to remove duplicates. When a patient had more than one isolate of the same organism within a 30 day period it was removed. To determine outpatient data, facilities in Clark County such as general medical-surgical hospitals, rehabilitation and specialty hospitals,

nursing homes and hospices were removed. In addition, only those bacterial species isolated ≥ 500 times since January 2004 were included in the analysis. Since *Serratia* spp. were isolated only 96 times, they will not be included in this report.

Results

Numerous bacterial species were isolated from urine and urethra sources. Some of these likely represent contaminants. The most common organism reported is *Escherichia coli* followed by *Klebsiella pneumoniae*, *Enterococcus* spp., *Proteus mirabilis*, Coagulase negative *Staphylococcus* spp., *Pseudomonas aeruginosa*, and *Staphylococcus aureus*. Box 1 illustrates the number of isolates identified by their degree of susceptibility to various antibiotics in quarter 1 of 2006 for the three most frequently isolated organisms. The percent susceptibilities are indicated within the bar charts for each organism. Table 1 lists the number of isolates tested for each bacterial species isolated ≥ 500 times from 1/1/2004 to 3/31/2006 and their percent susceptibilities. It is important to note that not every isolate is tested for sensitivity to every antibiotic in the tables.

Discussion

These data include the best approximation of local outpatient urine culture results. In some instances, the raw data might not indicate the specimen source or there is inconsistency in naming the source and thus may have been excluded. Additionally, data may include outpatients that are catheterized. Locally, *E. coli* is isolated from the urinary tract more than seven times greater than all other organisms in this report.

Antimicrobials commonly recommended for empiric treatment of uncomplicated urinary tract infections (uUTIs) include but are not limited to trimethoprim-sulfamethoxazole (TMP-SMX), ciprofloxacin or fluoroquinolones, nitrofurantoin, fosfomycin, cephalexin or other cephalosporins, amoxicillin, and amoxicillin-clavulanic acid (1, 2, 3). If community resistance to TMP-SMX is $\geq 20\%$ or the patient has a history of

allergy to the drug, it is necessary to replace this inexpensive and effective drug with one of the alternatives (3). Urinary tract isolates of *E. coli* in our community show greater than 20% resistance to TMP-SMX. Several antimicrobials to which this organism is greater than 90% susceptible are recommended when empiric treatment is chosen. Although fluoroquinolones are very effective and have an important role in treating uncomplicated cystitis, rising fluoroquinolone resistance is a serious public health threat and routine use for initial therapy is discouraged (3,4,5).

Some experts suggest that routine urine cultures are not necessary for UTIs because of the predictable nature of the causative organism (3). Others counter that pre-therapy cultures are important especially from a public health viewpoint because this information both confirms the diagnosis and reveals the antibiotic susceptibility patterns in the community (4). In an age where bacteria are becoming more resistant and antibiotic treatment alternatives are becoming less abundant and more expensive, the clinician should consider obtaining a urine culture and sensitivity to ascertain the most appropriate first line antibiotic for treatment.

Ideally, a specimen would be collected for culture and the patient could be placed on the most appropriate empiric choice antibiotic while awaiting culture results. Since it would be difficult to assume that the UTI was caused by another organism, without culture evidence, choosing an antibiotic that would be effective against the most likely organisms may be a viable treatment option, with any later modification in therapy based on subsequent culture results. When prescribing antibiotics clinicians should always speak with the patients about the use and non-use of antibiotics, the rationale for taking them and the importance of adhering to the instructions.

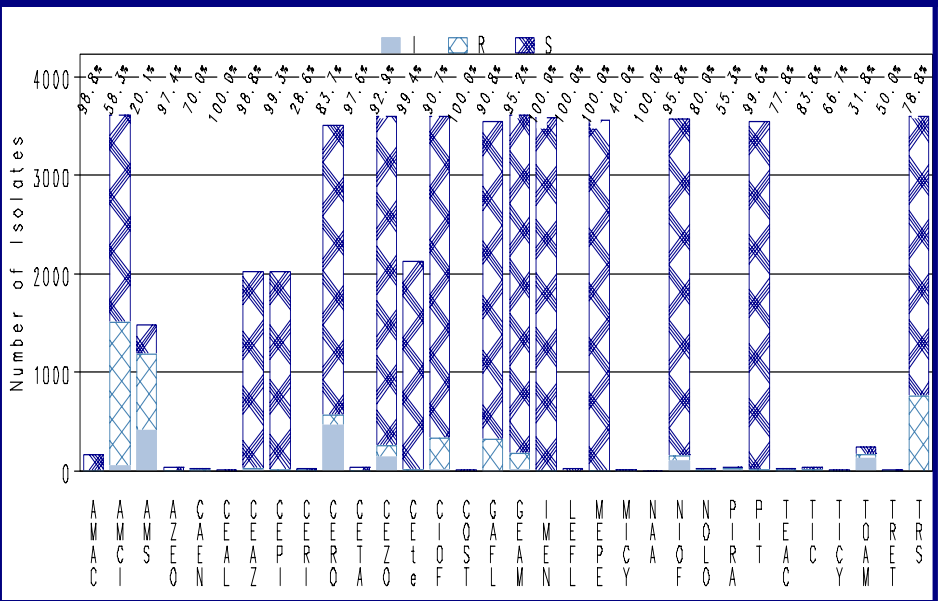
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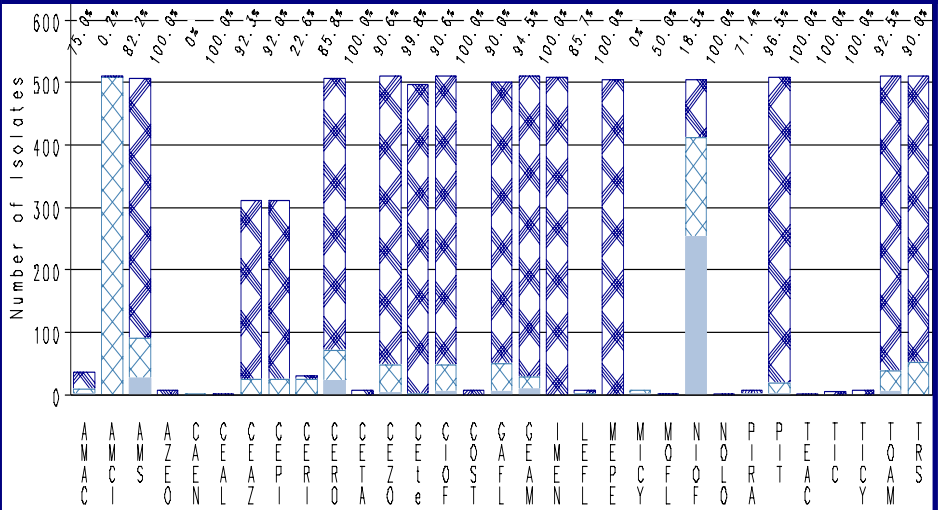
Box 1. Number of Isolates Tested by AR Result (Qtr1, 2006)

Escherichia coli: Susceptibility data reported in Qtr1, 2006

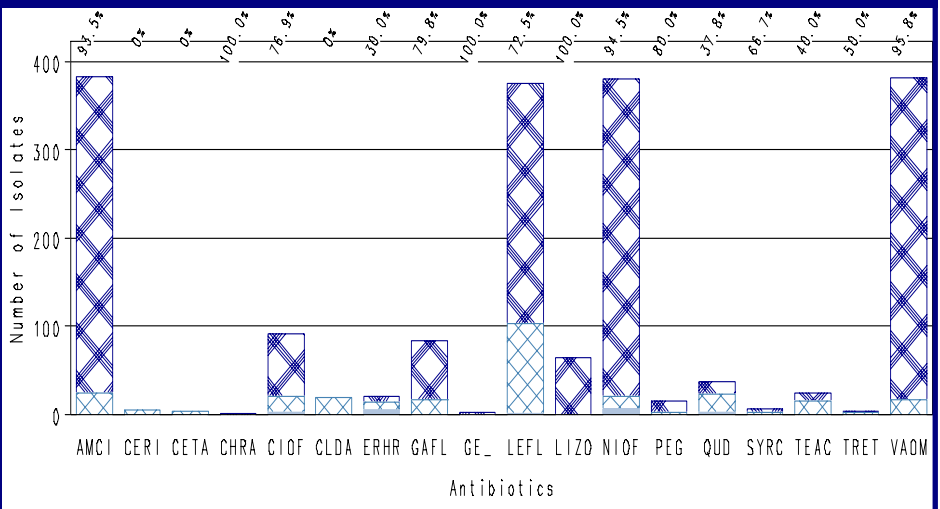
	Enterococcus spp.				Escherichia coli				Klebsiella pneumoniae			
	I	R	S	All	I	R	S	All	I	R	S	All
AMAC	0	0	0	0	1	1	168	170	2	7	27	36
AMCI	0	25	358	383	49	1459	2104	3612	0	509	1	510
AMS	0	0	0	0	412	773	298	1483	26	64	416	506
AZEO	0	0	0	0	0	1	37	38	0	0	8	8
CAEN	0	0	0	0	1	5	14	20	0	1	0	1
CEZO	0	0	0	0	140	117	3343	3600	4	44	462	510
CEPI	0	0	0	0	0	14	2003	2017	0	25	286	311
CETA	0	4	0	4	0	1	40	41	0	0	7	7
CEAZ	0	0	0	0	8	17	1992	2017	0	24	287	311
CERI	0	5	0	5	0	15	6	21	0	24	7	31
CERO	0	0	0	0	459	111	2934	3504	23	49	434	506
CEAL	0	0	0	0	0	0	5	5	0	0	1	1
CHRA	0	0	1	1	0	0	0	0	0	0	0	0
CIOF	2	19	70	91	4	331	3258	3593	5	43	463	511
CLDA	0	19	0	19	0	0	0	0	0	0	0	0
COST	0	0	0	0	0	0	5	5	0	0	8	8
CEte	0	0	0	0	6	6	2109	2121	0	1	495	496
ERHR	5	9	6	20	0	0	0	0	0	0	0	0
G AFL	0	17	67	84	2	322	3215	3539	6	44	451	501
GEAM	0	0	0	0	7	167	3438	3612	9	19	483	511
GE_	0	0	2	2	0	0	0	0	0	0	0	0
IMEN	0	0	0	0	0	1	3584	3585	0	0	509	509
LEFL	1	102	272	375	0	0	18	18	0	1	6	7
LIZO	0	0	64	64	0	0	0	0	0	0	0	0
MEPE	0	0	0	0	0	1	3556	3557	0	0	504	504
MICY	0	0	0	0	1	2	2	5	2	6	0	8
MOFL	0	0	0	0	0	0	0	0	0	1	1	2
NAA	0	0	0	0	0	0	1	1	0	0	0	0
NIOF	6	15	359	380	99	52	3420	3571	253	158	93	504
NOLO	0	0	0	0	0	4	16	20	0	0	1	1
PEG	0	3	12	15	0	0	0	0	0	0	0	0
PIRA	0	0	0	0	3	14	21	38	2	0	5	7
PIT	0	0	0	0	13	2	3530	3545	2	16	491	509
QUD	2	21	14	37	0	0	0	0	0	0	0	0
SYRC	0	2	4	6	0	0	0	0	0	0	0	0
TEAC	0	15	10	25	1	5	21	27	0	0	1	1
TIC	0	0	0	0	5	1	31	37	0	0	6	6
TICY	0	0	0	0	0	1	2	3	0	0	8	8
TOAM	0	0	0	0	123	40	76	239	5	33	472	510
TRET	0	2	2	4	0	2	2	4	0	0	0	0
TRS	0	0	0	0	0	760	2830	3590	0	51	459	510
VAOM	0	16	366	382	0	0	0	0	0	0	0	0



Klebsiella pneumoniae: Susceptibility data reported in Qtr1, 2006



Enterococcus spp.: Susceptibility data reported in Qtr1, 2006



See below for antibiotic codes, %susceptible noted on charts.

Antibiotic codes: AMAC:amikacin AMCI:ampicillin AMS:ampicillin_sulbactam AZEO:aztreonam CAEN:carbenicillin CEAL:cephalothin CEAZ:ceftazidime CEPI:cefepime CERI:ceftriaxone CERO:cefuroxime CETA:cefotaxime CEte:cefotetan CEZO:cefazolin CHRA:chloramphenicol CIOF:ciprofloxacin CLDA:clindamycin COST:colistin ERHR:erythromycin GAFL:gatifloxacin GEAM:gentamicin GE_gent_synergy IMEN:imipenem LEFL:levofloxacin LIZO:linezolid MEPE:meropenem MICY:minocycline MOFL:moxifloxacin NAA:maladixic_acid NIOF:nitrofurantoin NOLO:norfloxacine PEG:penicillin_g PIRA:piperacillin PIT:piperacillin_tazobactam QUD:quinupristin_dalfopristin SYRC:synergicid TEAC:tetracycline TIC:ticarcillin_clavulanic_acid TICY:tigecycline TOAM:tobramycin TRET:trimethoprim TRS:trimethoprim_sulfamethoxazole VAOM:vancomycin

