

2020 Kansas Antibiogram

Methodology

Collection Process and State Representation: Antibiotic susceptibility data was collected amongst 66 Kansas healthcare facilities. One hundred and twenty-three healthcare facilities were contacted regarding institutional antibiograms, of which 66 responded with results. Of the 82 critical access hospitals contacted, 47 provided results, 14 had no antibiogram, 1 had an antibiogram in process not provided, and 20 did not report back to which a second attempt was made. Amongst 51 acute care hospitals, 30 provided antibiograms and 21 did not provide even with re-requests sent out. Of these 77 received, 13 were redundant (i.e., the critical access hospital used the nearby acute care hospital's antibiogram). Two clinics also were able to provide data. The state was well represented, with facilities in each region: 10 northwest, 7 northcentral, 11 northeast (excluding Kansas City metro hospitals), 6 Kansas City metro facilities, 9 southeast, 11 southcentral, 4 Wichita metro, and 8 southwest facilities. Given a relative lack of antibiograms in many critical access and rural clinics compared to their urban peers, we attempted as much as possible to provide regional antibiotic susceptibility patterns.

Antibiogram development: The CLSI guidelines were followed in the aggregation of data from all reported hospital antibiograms. Antibiotic and organism combinations intrinsically resistant or clinically irrelevant were censored or grayed in the antibiogram.

Limitations: The majority of data provided was from reference labs in alignment with Clinical and Laboratory Standards Institute (CLSI) guidelines. However, 31 facilities reported back institutional antibiogram forms. Of these, 15 were from tertiary and large acute care hospitals whose labs were confirmed to be in alignment with CLSI. Sixteen forms were from critical access hospitals, of which the reference lab to which this data was acquired were unable to be confirmed was CLSI guidelines. An internal assessment of outliers or implausible data was conducted. Attempts were made to confirm outlying data. As confirmation could not be acquired, this data was excluded. This was a rare event, occurring no more than 4 or 5 times out of the thousands of susceptibility profiles.

Acknowledgements

We would like to acknowledge the clinical microbiologists who submitted antibiogram data on behalf of their healthcare facility. We would also like to thank our academic partners at the University of Kansas whose infectious disease physicians and infectious disease pharmacists contributed directly to the creation and clinical content of this antibiogram:

Kellie Wark, MD MPH

Rachel Weihe, MD

Nicole Wilson, PharmD, BCIDP

The KDHE HAI/AR Program is a resource for developing and strengthening Kansas healthcare facilities stewardship activities. Visit <https://www.kdheks.gov/epi/hai.htm> for more information.

Gram positive table key

Gray = Insufficient data to compile total susceptibility, not tested, or inappropriate for clinical use because of intrinsic use	
1	Rifampin to be used in combination only (rapid resistance emergence on single-agent therapy)
2	Ampicillin predicts amoxicillin susceptibility
3	Oxacillin predicts nafcillin, cefazolin, cephalexin susceptibility
4	Erythromycin predicts azithromycin susceptibility
5	Nitrofurantoin only for urinary infections
6	Gentamicin for synergy
7	Northeast region excluding KC metro
8	<i>E. faecalis</i> KC-metro linezolid susceptibility is lower likely as a reflection of greater VRE- <i>faecalis</i> only isolates linezolid susceptibilities reported at 1 over-represented institution
9	<i>S. pneumoniae</i> meningeal / non-meningeal breakpoint rates
10	<i>C. striatum</i> develops rapid daptomycin resistance while on treatment
11	Low numbers of <i>S. hominis</i> and <i>S. pyogenes</i> , numbers <30 may be statistically unreliable
12	Strep anginosus group (anginosus, constellatus, intermedius)
13	Strep mitis group (mitis, oralis, mutans, bovis, sanguinis)

Gram negative / yeast table key

Gray = Insufficient data to compile total susceptibility, or not tested, or inappropriate in clinical use because of intrinsic use	
1	Ampicillin predicts amoxicillin susceptibility
2	Ampicillin/sulb. generally predicts amoxicillin/clav. susceptibility (except for <i>A. baumannii</i> which is intrinsically resistant)
3	Nitrofurantoin only for urinary infections
4	Northeast region excluding KC metro
5	<i>C. glabrata</i> has no established voriconazole breakpoint (i.e. current data are insufficient to demonstrate a correlation between in vitro susceptibility and clinical outcome for <i>C. glabrata</i> and voriconazole)
6	Isolate numbers of <i>C. tropicalis</i> <30, and may be statistically unreliable
7	For yeasts, data are susceptible-dose dependent; in vivo susceptibility is dependent upon achieving the maximal possible blood level for mucosal and invasive infections