# The Use of Plasma Next-Generation Sequencing Test in the Management of Immunocompetent and Immunocompromised Patients – A Single Center Retrospective Study

# Baylor Collegeof Medicine



Denise Marie A. Francisco, MD<sup>1</sup>, Laila Woc-Colburn, MD<sup>1</sup>, Travis Carlson, PharmD, RPh<sup>2</sup>, Todd Lasco, PhD, D(ABMM)<sup>3</sup>, Miriam Barrett, MBA, PMP<sup>3</sup>, Mayar Al-Mohajer, MD<sup>1</sup> <sup>1</sup>Section of infectious Diseases, Baylor College of Medicine, Houston, Texas; <sup>2</sup>Department of Pathology and Clinical Microbiology, CHI St. Luke's Health – Baylor St. Luke's Health – – Baylor St. Luke's Medical Center, Houston, Texas

# Abstract

# BACKGROUND

Microbiological culture data is a longstanding gold standard in diagnostics. Unfortunately, yield from cultures have been inconsistent and slow, prompting the need for newer tests including the plasma-based next generation sequencing (NGS) tool.

This study aims to describe the use of NGS and the corresponding change in management.

## METHODS

A descriptive retrospective study was done on hospitalized adults at CHI-Baylor St. Luke's in Houston, Texas with NGS tests from Jan 1, 2017 to Dec 31, 2018.

### RESULTS

There were 167 NGS tests performed. Most patients were non-Hispanic (n=129) Caucasian (n=106) males (n=116) with a mean age of 52. Furthermore, 61 were immunocompromised patients [solid organ transplant (n=30), HIV-AIDS (n=14) and rheumatology patients on immunosuppression (n=12)].

During the study, the hospital staff prepared a list of indications for NGS testing including: systemic or deep seated infection where a biopsy or other workup is negative or not possible (n=50), fever of unknown origin (n=26), culture negative endocarditis (n=15), HIV/AIDS with fever (n=10), transplant patient with fever (n=5). There were 60 cases where the indications were not on this list (36%).

Results showed that 118/167 (71%) were positive. The most common organisms identified were gram negative bacteria (54/118; 46%) followed by viruses (49/118; 42%), gram-positive bacteria (48/118; 41%), fungi (16/118; 14%), atypical bacteria (9/118; 8%), mycobacterium (4/118; 3%), and parasites (4/118; 3%). Blood cultures were concurrently obtained in 148/167 (89%) of the cases and returned negative in 137/148 (93%) of cases.

In terms of change of management, the largest change was found in glycopeptide use (36 fewer patients after NGS results). Next was on anti-mycobacterial drugs where 27 were added among 8 instances. Only 36 patients were taken off antibiotics, even though 49 patients had negative results. In total, 120 out of 160 cases had antibiotic changes.

### CONCLUSION

We observed a large decrease in glycopeptide use after NGS results which suggests physicians' comfort in withdrawing MRSA coverage. In addition, antimycobacterial coverage increased corresponding to early mycobacterial detection with NGS. This study highlights the importance of clinical judgement in the age of rapid diagnostics.

- To describe change in management in antibiotic choice before and after a NGS test
- Exclusion:
- Less than 18 years old have a result due to NGS tests that did not inadequate/inappropriate sample

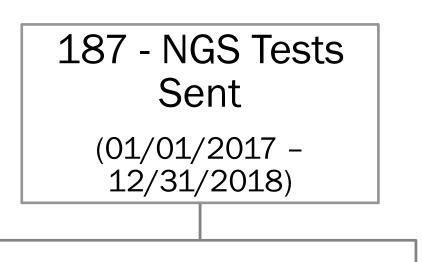
Figure 1. Next Generation Sequencing Test Breakdown

<u>Included</u> 167 - Had Results

	Variable
Age (mean, S	SD)
Gender	
Male	
Female	
Race	
Caucasiar	
Black or A	Trican Am
Asian Others	
Ethnicity Hispanic	
Not Hispa	nic
Unable to	
Charlson's C	
(mean, SD)	
Immune Sys	tem Statu
Immunoco	ompetent
Immunosi	• •
HIV-AIC	
Neutro	•
	rgan Tran opoietic S <sup>.</sup>
Transplant	Spoletic S
•	atological
Others	0
Approved Ind	dications f
Culture N	egative Er
Fever of L	Jnknown (
•	with Feve
	nt with Fev
Systemic/ Where Biops	Deep Sea
Negative or l	-
Others	1011 0331
Waiting Perio	od (mean,
-	veen Colle
Received	
Days betw	veen Colle
Reported	

# **Aims and Methods**

- Number of antibiotics used
  - Change in treatment (based on pathogen coverage)

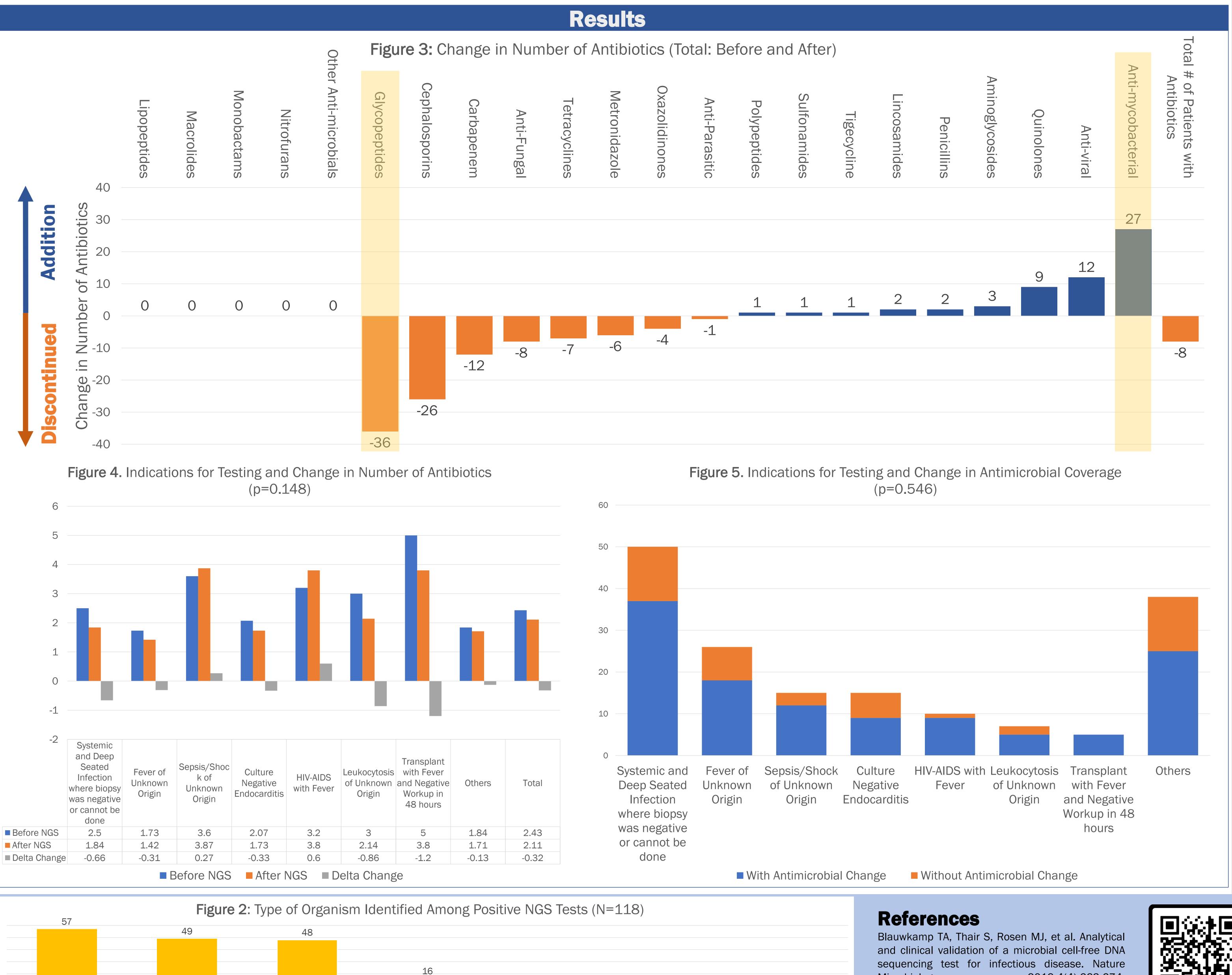


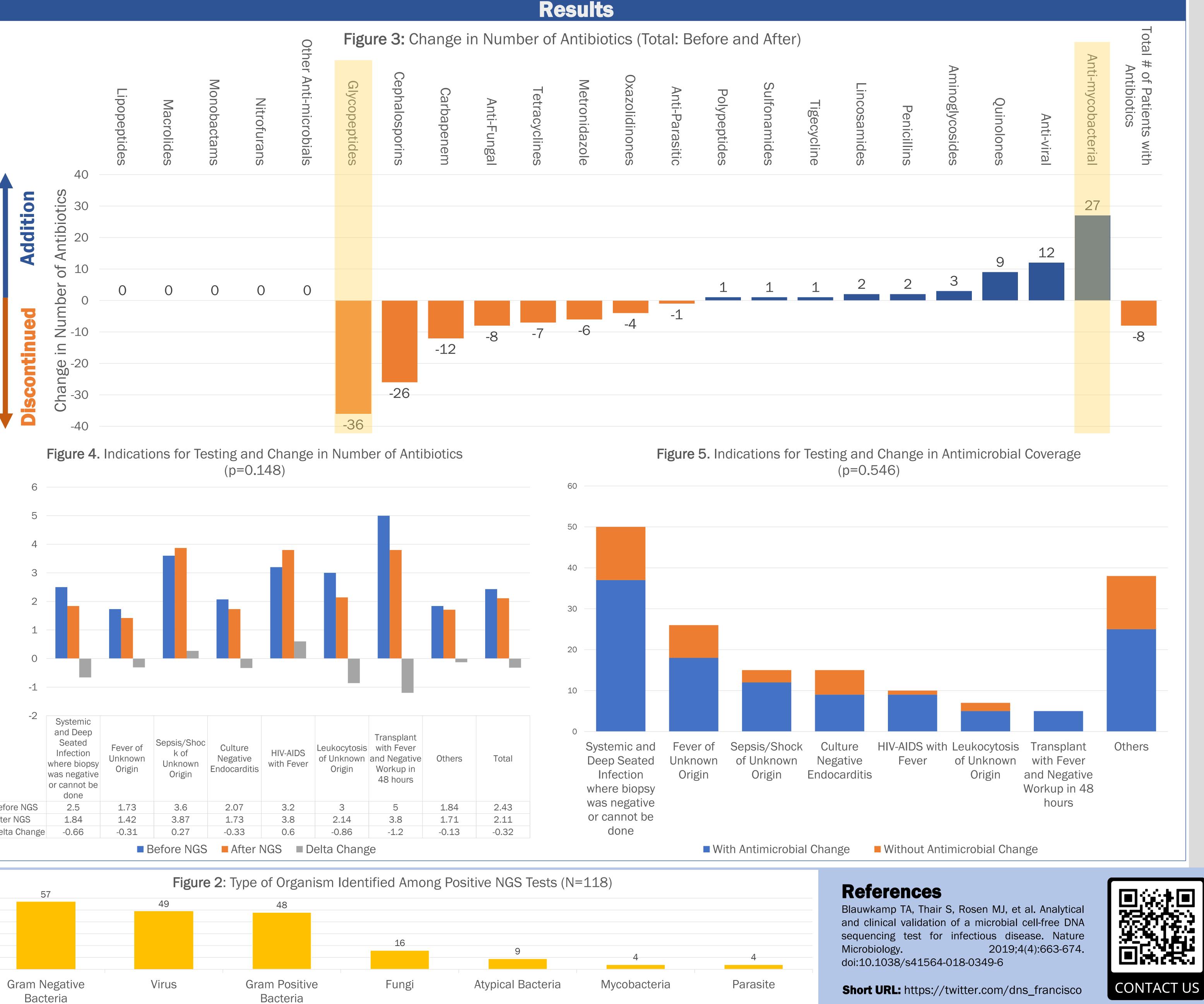
<u>Excluded</u> 20 - Inadequate Samples / No results

# Results

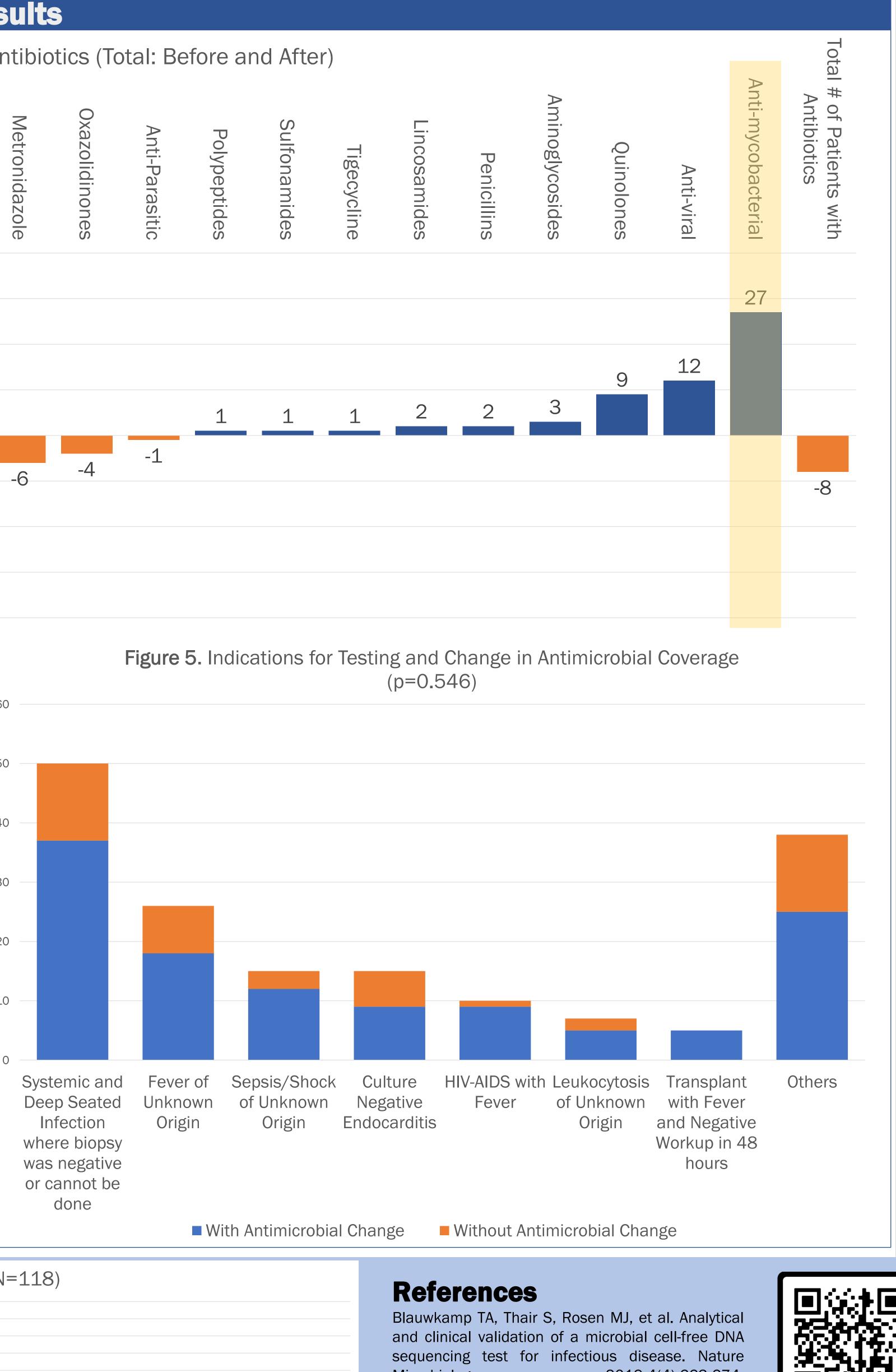
**Table 1.** Demographic and Laboratory Characteristics

logiaphic	lographic and Laboratory Characteristics							
	Total N (%/SD) N=167	NGS + (%/SD) N=118	NGS - (%/SD) N=49	P-Value				
	52 (16)	51 (16)	54 (15)	0.26				
	116 (69%) 51 (31%)	83 (70%) 35 (30%)	33 (67%) 16 (33%)	0.70				
e nerican	106 (63%) 38 (23%) 13 (8%) 10 (6%)	75 (64%) 28 (24%) 9 (8%) 6 (5%)	31 (63%) 10 (20%) 4 (8%) 4 (8%)	0.87				
ne	36 (22%) 129 (77%) 2 (1%)	27 (23%) 89 (75%) 2 (2%)	9 (18%) 40 (82%) 0 (0%)	0.44				
ity Index	3.70 (2.79)	3.74 (2.76)	3.61 (2.87)	0.79				
us c d hsplant Stem Cell	106 (64%) 61 (36%) 14 (23%) 3 (5%) 30 (49%) 1 (2%)	75 (64%) 43 (36%) 13 (30%) 2 (5%) 21 (49%) 0 (0%)	31 (63%) 18 (37%) 1 (6%) 0 (0%) 9 (50%) 1 (6%)	0.97				
al	12 (20%) 1 (2%)	7 (16%) 0 (0%)	5 (28%) 1 (6%)					
for Testing Indocarditis Origin er ever eated Infection er Workup is ible	107 (64%) 15 (9%) 26 (16%) 10 (6%) 5 (3%) 50 (30%) 60 (36%)	10 (8%) 20 (17%) 9 (8%) 4 (3%)	5 (10%) 6 (12%) 1 (2%) 1 (2%)	0.23				
<b>, SD)</b> lected and lected and	2 (1) 3 (1)	2 (1) 3 (1)	2 (1) 3 (1)					





/Shoc of Iown gin	Culture Negative Endocarditis	HIV-AIDS with Fever	Leukocytosis of Unknown Origin	Transplant with Fever and Negative Workup in 48 hours	Others	Total
6	2.07	3.2	3	5	1.84	2.43
37	1.73	3.8	2.14	3.8	1.71	2.11
27	-0.33	0.6	-0.86	-1.2	-0.13	-0.32
NGS After NGS Delta Change						



CONTACT US: Denise Marie A. Francisco, MD dafrancisco@mdanderson.org dns.a.francisco@gmail.com 1515 Holcombe Boulevard, Unit 1460 Houston, Texas 77030 Phone: 713-792-6830 Fax: 713-745-6839 TWITTER: @dns\_francisco