

Supplementary Table 2. Number and percentage of GISP isolates¹ reported from 2018 encoding susceptibility associated genotypes by sexual behavior and race/ethnicity. Dash indicates that no isolates were collected from the demographic group indicated.

	PCN Susceptible Genotype			TET Susceptible Genotype		
	MSM	MSMW	MSW	MSM	MSMW	MSW
American Indian	0 (0.0%)	-	0 (0.0%)	0 (0.0%)	-	2 (40.0%)
Asian	0 (0.0%)	2 (5.1%)	1 (5.0%)	1 (5.3%)	0 (0.0%)	5 (25.0%)
Black	20 (5.0%)	0 (0.0%)	33 (5.7%)	13 (10.8%)	3 (7.7%)	32 (22.9%)
Hispanic	1 (0.9%)	0 (0.0%)	1 (0.9%)	13 (11.8%)	1 (7.7%)	13 (11.6%)
Multi-racial	0 (0.0%)	0 (0.0%)	2 (10.0%)	1 (6.25%)	0 (0.0%)	4 (20.0%)
Native Hawaiian	0 (0.0%)	0 (0.0%)	0 (0.0%)	1 (11.1%)	0 (0.0%)	0 (0.0%)
White	0 (0.0%)	0 (0.0%)	7 (6.8%)	14 (8.0%)	3 (11.5%)	17 (16.5%)
p-value (χ^2 test)	0.073		0.96	0.52		0.19

References.

- 1 Reimche JL, Chivukula VL, Schmerer MW, *et al.* Genomic analysis of the predominant strains and antimicrobial resistance determinants within 1479 *Neisseria gonorrhoeae* isolates from the U.S. Gonococcal Isolate Surveillance Project in 2018. *Sex Transm Dis* 2021; published online May 14. DOI:10.1097/OLQ.0000000000001471.