## Staphylococcus aureus - NRS383

Other designations (USA200; 96758)

	Demographic Informa	ition:	
Country/State:	United States / North Carolina	Date of Isolation:	N/A
Patient Age:	N/A	Gender:	N/A
Patient Location:	N/A	Patient Service:	N/A
Culture Source:			•
Phenotypic Properties:	MRSA, also resistant to ERY, CLIN, T	ET, SXT, LEV, GM	
Genotypic Properties:	PFT is USA200, MLST is ST36, spa ty group III; tsst+, sea+	pe is WGKAKAOMQQQ, SCCmo	ec II, agr
Literature Citation:	JCM, 2003, 41:5113-5120.		
Comment:	N/A		
Description:	Second most common U.S. health care-associated PFT; EMRSA16 clone		
Repositories:	Known Clinically Associated Strains - Molecular Typing Control Strains	Others	

S = Susceptible; R = Resistant; NS = Not Susceptible; I = Intermediate; N/A = Not Available P = Pending results

NARSA Antimicrobial Profile for Other Antimicrobial Agents				Molecular Characterization Data	
Drug	μg/mL MIC <del> (pg/ml)</del>	CLSI		PCR	Result
Drug		Interptn		mecA	POSITIVE
Ciprofloxacin	> 16	R	1		
Clindamycin	> 4	R			
Daptomycin	= 0.5	N/A	1		
Erythromycin	> 8	R	1		
Gentamicin	> 16	R	1		
Linezolid	= 2	S	1		
Oxacillin	> 16	R	1		
Penicillin	> 2	R	1		
Quinupristin/dalfopristin	= 1	S			
Teicoplanin	= 1	S	1		
Tetracycline	= 4	S			
Trimeth/sulfa	= 2/38	S			
Vancomycin	= 0.5	S			

Genotype Data					
Genotype	Result				
eGenomic <i>spa</i> type	16				
eGenomic <i>spa</i> repeats	WGKAKAOMQQQ				
Ridom <i>spa</i> type	t018				
SCC <i>mec</i> type	II .				
Clonal Complex (CC)	30 (36)				
Sequence Type (ST)	36 - SLV				
Genotype Data file	•				