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TABLE E-1 Demographic and Clinical Data for Spinal Instrumentation to Treat Scoliosis

Data	Center 1* (N = 284)	Center $2* (N = 464)$	Center $3* (N = 599)$	P Value
Demographic				
Sex				0.011
Male	111 (39.1%)	180 (38.8%)	282 (47.1%)	
Female	173 (60.9%)	284 (61.2%)	317 (52.9%)	
Age† (yr)	11.5 ± 4.7	11.4 ± 4.8	11.5 ± 4.7	0.953
Race				< 0.001
Black	13 (4.6%)	97 (20.9%)	38 (6.3%)	
White	186 (65.5%)	314 (67.7%)	193 (32.2%)	
Asian	13 (4.6%)	12 (2.6%)	59 (9.8%)	
Hispanic	26 (9.2%)	35 (7.5%)	240 (40.1%)	
Other or not specified	46 (16.2%)	6 (1.3%)	69 (11.5%)	
Clinical				
Etiology				0.031
Idiopathic	107 (37.7%)	203 (43.8%)	199 (33.2%)	
Non-idiopathic				
Neuromuscular	80 (28.2%)	129 (27.8%)	182 (30.4%)	
Syndromic	30 (10.6%)	50 (10.8%)	79 (13.2%)	
Congenital	50 (17.6%)	53 (11.4%)	102 (17.0%)	
Other	17 (6.0%)	29 (6.3%)	37 (6.2%)	
Type of procedure				< 0.001
Primary arthrodesis	176 (62.0%)	261 (56.3%)	339 (56.6%)	
Revision arthrodesis	18 (6.3%)	28 (6.0%)	81 (13.5%)	
Growing construct insertion	24 (8.5%)	29 (6.3%)	30 (5.0%)	
Growing construct lengthening	41 (14.4%)	119 (25.6%)	84 (14.0%)	
Growing construct revision	25 (8.8%)	27 (5.8%)	65 (10.9%)	

^{*}The values are given as the number of procedures, with the percentage in parentheses. †The values are given as the mean and the standard deviation.

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TABLE E-2 Cultured Organisms (N = 140) from Seventy-one Culture-Positive Infections

Organisms	Frequency*	
Gram-positive	80 (57.1%)	
Staphylococcus aureus	35 (25.0%)	
Methicillin-susceptible	20 (14.3%)	
Methicillin-resistant	15 (10.7%)	
Coagulase-negative staphylococci	24 (17.1%)	
Enterococcal species	6 (4.3%)	
Group A beta-hemolytic streptococcus	3 (2.1%)	
Propionibacterium acnes	3 (2.1%)	
Viridians streptococci	3 (2.1%)	
Other†	6 (4.3%)	
Gram-negative	60 (42.9%)	
Pseudomonas aeruginosa	15 (10.7%)	
Escherichia coli	13 (9.3%)	
Enterobacter species	9 (6.4%)	
Klebsiella pneumoniae	6 (4.3%)	
Bacteroides species	5 (3.6%)	
Proteus species	4 (2.9%)	
Acinetobacter species	3 (2.1%)	
Other‡	5 (3.6%)	

^{*}The values are given as the number of organisms, with the percentage in parentheses. †Other gram-positive organisms included Bacillus circulans, Peptostreptococcus asaccharolyticus, Peptostreptococcus magnus, Clostridium sporogenes, and Corynebacterium species. ‡Other gram-negative organisms included Serratia marcescens, Citrobacter koseri, and Citrobacter freundii complex.