

1 *Supplemental data*

2 **Rapid diagnostics of orthopaedic implant-associated infections using nanopore shotgun metagenomics sequencing on tissue**
3 **biopsies**

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17 **Table S1. Results of microbial identification across conventional microbiological methods and nanopore shotgun metagenomic**
18 **sequencing for all OIAI patients' biopsies.** Culture growth densities are termed as follows: rich, moderate (mod), sparse, spread
19 single colonies (SSC), and requiring pre-cultivation in enrichment broth (broth). V* indicates the name of the individual tissue
20 biopsy, NA (not applicable), NTC (no template control), and EN (extraction negative control).

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Patient ID	Biopsy/control ID	Nanopore results: putative pathogens, reads	Culture results, growth	Nanopore results: putative background (species, reads)	Nanopore results: human DNA reads
101	VB	Negative	Culture negative	<i>E. coli</i> , 4; <i>Arthrobacter sp.</i> , 2; <i>Mucilaginibacter sp.</i> , 1	6
101	V2	Negative	Culture negative	<i>E. coli</i> , 15; <i>Arthrobacter sp.</i> , 1;	3,699

				<i>Mucilaginibacter</i> sp., 1	
101	V3	Negative	Culture negative	<i>E. coli</i> , 2; <i>Arthrobacter</i> sp., 1	161
101	V4	Negative	Culture negative	<i>E. coli</i> , 5	46
101	V5	Negative	Culture negative	<i>E. coli</i> , 12	60
101	EN	Negative	NA	<i>E. coli</i> , 3; <i>Arthrobacter</i> sp., 1	18
101	NTC	Negative	NA	<i>E. coli</i> , 5; <i>Arthrobacter</i> sp., 2;	2
				<i>Mucilaginibacter</i> sp., 1	
104	VB	<i>S. lugdunensis</i> , 44; <i>S. epidermidis</i> , 6; <i>C. acnes</i> , 2; others <2	<i>S. Lugdunensis</i> , sparse	<i>E. coli</i> , 8; others <2	89
104	V2	Negative	<i>S. Lugdunensis</i> , sparse	Negative	
104	V3	<i>S. lugdunensis</i> , 306; <i>S. epidermidis</i> , 5; <i>C. acnes</i> , 1	<i>S. Lugdunensis</i> , sparse	<i>E. coli</i> , 10; others, 1	308
104	V4	<i>S. lugdunensis</i> , 24;	<i>S. Lugdunensis</i> , ssc;	<i>E. coli</i> , 5; others, <2	168

		<i>C. acnes</i> , 2	<i>C. acnes</i> , ssc		
104	V5	<i>S. lugdunensis</i> , 58; <i>S. epidermidis</i> , 2	<i>S. Lugdunensis</i> , sparse	<i>M. globosa</i> , 160; <i>E. coli</i> , 2; others, <2	46283
104	V6	Negative	<i>S. Lugdunensis</i> , sparse; <i>C. acnes</i> ssc	<i>E. coli</i> , 6; others, 1	36
104	NTC	Negative	NA	Negative	
105	VB	Negative	Culture negative	<i>E. coli</i> , 4	821
105	V2	NA	<i>S. Aureus</i> , broth; <i>S. epidermidis</i> , broth	NA	
105	V3	NA	<i>S. Aureus</i> , broth; <i>S. epidermidis</i> , broth	NA	
105	V4	<i>S. aureus</i> , 49	<i>S. Aureus</i> , broth; <i>S. epidermidis</i> , broth	<i>E. coli</i> , 24	67150
105	V5	Negative	<i>S. Aureus</i> , ssc; <i>S. epidermidis</i> , broth	<i>E. Coli</i> , 16	10235
105	EN	Negative	NA	<i>E. coli</i> , 8;	1

				<i>Mycobacterium abscessus</i> ,2	
105	NTC	Negative	NA	<i>E. coli</i> ,1;	1
107	VB	<i>S. epidermidis</i> , 33; <i>S. aureus</i> , 2	<i>S. epidermidis</i> , ssc	<i>E. coli</i> , 32; <i>B. subtilis</i> , 5; <i>Arthrobacter sp.</i> , 5; <i>M. globosa</i> , 4; <i>Mucilaginibacter sp.</i> , 2	1944
107	V2	Negative	<i>S. epidermidis</i> , broth	<i>E.coli</i> , 31; <i>Arthrobacter sp.</i> , 2; <i>Bacillus mycoides</i> , 2	479
107	V3	Negative	<i>S. epidermidis</i> , ssc	<i>E.coli</i> , 31; <i>Arthrobacter sp.</i> , 3; <i>B. subtilis</i> , 3; <i>Mucilaginibacter sp.</i> , 2	1173
107	V4	Negative	<i>S. epidermidis</i> , broth	<i>E. coli</i> , 16; <i>Arthrobacter sp.</i> , 3; <i>Mucilaginibacter sp.</i> , 3	15
107	V5	Negative	Culture negative	<i>E coli</i> , 97; <i>B. subtilis</i> , 4; <i>Mucilaginibacter sp.</i> , 4; <i>M. globosa</i> , 2; <i>Arthrobacter sp.</i> , 4;	97

				<i>Bacillus mycoides</i> , 2	
107	EN	Negative	NA	<i>E coli</i> , 15; <i>Arthrobacter sp.</i> , 2; <i>C. acnes</i> , 2; <i>B. mycoides</i> , 2;	4
107	NTC	Negative	NA	<i>E coli</i> , 13; <i>B. subtilis</i> , 4; <i>Mucilaginibacter sp.</i> , 2; <i>Arthrobacter sp.</i> , 2; <i>Bacillus mycoides</i> , 1	
108	VB	<i>S. aureus</i> , 2338	<i>S. aureus</i> , ssc	<i>M. Globosa</i> , 42; <i>E. Coli</i> , 5; others, <5	11400
108	V2	<i>S. aureus</i> , 80	<i>S. aureus</i> , ssc	<i>Arthrobacter sp.</i> , 6; others, <6	2351
108	V3	<i>S. aureus</i> , 369	<i>S. aureus</i> , sparse	<i>M. Globosa</i> , 77; <i>Arthrobacter sp.</i> , 3; Others, <3	38980
108	V4	<i>S. aureus</i> , 4;	<i>S. aureus</i> , ssc	<i>E. Coli</i> , 25 <i>C. Acnes</i> , 4; others, <4	59
108	V5	<i>S. aureus</i> , 182	<i>S. aureus</i> , sparse	<i>E. Coli</i> , 33; <i>B. Subtilis</i> , 2; others, 1	83
108	EN	Negative	NA	Negative	

108	NTC	Negative	NA	Negative	
109	VB	<i>S. aureus</i> , 70	<i>S. aureus</i> , ssc	<i>M. globosa</i> , 22; <i>E.coli</i> , 17; <i>C. acnes</i> , 7; <i>Arthrobacter sp.</i> , 4; <i>B. subtilis</i> , 2	16781
109	V2	<i>S. aureus</i> , 3	<i>S. aureus</i> , ssc	<i>M. globosa</i> , 27; <i>E. coli</i> , 22; <i>B. subtilis</i> , 10; <i>Arthrobacter sp.</i> , 4	21171
109	V3	<i>S. aureus</i> , 2	<i>S. aureus</i> , ssc	<i>E. coli</i> , 16; <i>Arthrobacter sp.</i> , 5; <i>M. globosa</i> , 16; <i>C. acnes</i> , 3; <i>B. subtilis</i> , 3	25931
109	V4	<i>S. aureus</i> , 1	<i>S. aureus</i> , ssc	<i>E. coli</i> , 16; <i>Arthrobacter sp.</i> , 3; <i>B. subtilis</i> , 2	323
109	V5	Negative	<i>S. aureus</i> , ssc	<i>E. coli</i> , 9; <i>Arthrobacter sp.</i> , 2; <i>B. subtilis</i> , 3	56
109	EN	Negative	NA	<i>E coli</i> , 7; <i>B. subtilis</i> , 5	3
109	NTC	Negative	NA	Negative	5

110	VB	Negative	<i>S. aureus</i> , ssc	<i>M. globosa</i> , 2; <i>E. coli</i> , 2; others, <2	38222
110	V2	Negative	<i>S. aureus</i> , ssc	<i>M. globosa</i> , 56; <i>E. coli</i> , 9; others, 1	1029
110	V3	Negative	<i>S. aureus</i> , ssc	<i>M. globosa</i> , 9; <i>E. coli</i> , 5; others, 1	2043
110	V4	Negative	<i>S. aureus</i> , ssc	<i>E. coli</i> , 8; others, 1	28
110	V7	Negative	<i>S. aureus</i> , ssc	<i>E. coli</i> , 7; other, 1	1
110	NTC	Negative	NA	Negative	
111	VB	<i>S. aureus</i> , 537	<i>S. aureus</i> , rich	<i>M. globosa</i> , 12; others, <5	945
111	V2	<i>S. aureus</i> , 5683	<i>S. aureus</i> , rich	<i>M. globosa</i> , 141; <i>E. coli</i> , 5; others, <5	75837
111	V4	<i>S. aureus</i> , 217057; <i>S. argenteus</i> 68; <i>S. lugdunensis</i> , 53; <i>S. epidermidis</i> , 42; <i>S. pseudointermedius</i> , 30	<i>S. aureus</i> , rich	<i>E. coli</i> , 19; others <19	1799
111	V5	<i>S. aureus</i> , 55397;	<i>S. aureus</i> , rich	<i>M. globosa</i> , 24; <i>E. coli</i> , 16;	15138

		<i>S. lugdunensis</i> , 17		others <16	
111	V6	<i>S. aureus</i> , 2875	<i>S. aureus</i> , rich	<i>C. acnes</i> , 15; <i>E.coli</i> , 11; <i>M. globosa</i> , 4; others, <4	2064
111	NTC	Negative	NA	<i>E. coli</i> , 1; <i>B. mycoides</i> , 1	
112	VB	<i>S. aureus</i> , 19	<i>S. aureus</i> , sparse	others, 1	29
112	V2	<i>S. aureus</i> , 6150	<i>S. aureus</i> , sparse	<i>M. globosa</i> , 15; <i>E. coli</i> , 8; others <3	8022
112	V3	<i>S. aureus</i> , 3326; <i>S. lugdunensis</i> , 4	<i>S. aureus</i> , rich	others, <4	154
112	V4	<i>S. aureus</i> , 143	<i>S. aureus</i> , sparse	<i>M. globosa</i> , 9; <i>E. coli</i> , 4; others <4	2334
112	NTC	Negative	NA	Negative	
114	V2	<i>S. aureus</i> , 190119; <i>S. argenteus</i> , 64; <i>S. lugdunensis</i> , 27	<i>S. aureus</i> , mod	many others, <25	3524
114	V3	<i>S. aureus</i> , 582103; <i>S. argenteus</i> , 113; <i>S. lugdunensis</i> , 42;	<i>S. aureus</i> , rich	many others <25	449

		<i>S. epidermidis</i> , 29			
114	V4	<i>S. aureus</i> , 360674; <i>S. argenteus</i> , 56; <i>S. lugdunensis</i> , 18	<i>S. aureus</i> , mod	many others <25	70
114	V5	<i>S. aureus</i> , 52	<i>S. aureus</i> , rich	many others <3	5397
114	V6	<i>S. aureus</i> , 526710; <i>S. argenteus</i> , 150; <i>S. lugdunensis</i> , 68; <i>S. epidermidis</i> , 39	<i>S. aureus</i> , mod	many others <30	142
114	EN	<i>S aureus</i> , 29	NA	many others <3	11
114	NTC	Negative	NA	<i>S. aureus</i> , 7; <i>M. oryzae</i> , 1	
115	VB	<i>C. acnes</i> , 23	<i>C. acnes</i> , ssc, anaerobic	<i>E. coli</i> , 4; others 1	880
115	V2	<i>C. acnes</i> , 11	Culture negative	<i>S. equorum</i> , 3; others, <3	3722
115	V3	<i>C. acnes</i> , 31	Culture negative	<i>M. globosa</i> , 22; <i>E. coli</i> , 10; <i>Klebsiella pneumoniae</i> , 4; others, <4	9722

115	V4	<i>C. acnes</i> , 30	<i>C. acnes</i> , ssc, anaerobic	<i>E. coli</i> , 15; <i>Arthrobacter sp.</i> , 3; others, <3	44
115	V5	<i>C. acnes</i> , 60	<i>C. acnes</i> , ssc, anaerobic	<i>E. coli</i> , 19; <i>M. globosa</i> , 3; others, <3	1105
115	EN	Negative	NA	<i>E. coli</i> , 7; <i>C. jeikeium</i> , 1	
115	NTC	Negative	NA	Negative	
116	VB	<i>S. aureus</i> , 18; <i>S. epidermidis</i> , 4;	<i>S. aureus</i> , ssc	<i>E. coli</i> , 37; <i>B. subtilis</i> , 6	684
116	V2	<i>S. aureus</i> , 13277; <i>S. argenteus</i> , 15; <i>S. nepalensis</i> , 12; <i>S. epidermidis</i> , 11; <i>S. agnetis</i> , 10; <i>S. haemolyticus</i> , 6	<i>S. aureus</i> , sparse	Negative	34
116	V3	<i>S. aureus</i> , 10	<i>S. aureus</i> , ssc	<i>E. coli</i> , 13;	176

				<i>M. osloensis</i> , 3; <i>B. subtilis</i> , 2	
116	V4	<i>S. aureus</i> , 1	<i>S. aureus</i> , ssc	<i>E. coli</i> , 24; <i>B. subtilis</i> , 5	875
116	R4	<i>S. aureus</i> , 450 <i>S. epidermidis</i> , 1	<i>S. aureus</i> , sparse	<i>E. coli</i> , 23; <i>B. subtilis</i> , 7	3226
116	EN	<i>S. epidermidis</i> , 1	NA	<i>E. coli</i> , 8; others, 1	1
116	NTC	Negative	NA	Negative	1
117	VB	<i>S. aureus</i> , 150	<i>S. aureus</i> , mod	Others, 2	124
117	V2	<i>S. aureus</i> , 8	<i>S. aureus</i> , sparse	Others, 2	1634
117	V3	<i>S. aureus</i> , 70; <i>S. epidermidis</i> , 2	<i>S. aureus</i> , sparse	Others, 2	410
117	V4	<i>S. aureus</i> , 6	<i>S. aureus</i> , ssc	Others, 2	102
117	V5	<i>S. aureus</i> , 156; <i>S. epidermidis</i> , 4	<i>S. aureus</i> , sparse	Negative	66
117	NTC	Negative	NA	Others, 1	20
118	VB	Negative	Culture negative	Negative	1821

118	V2	Negative	Culture negative	Negative	417
118	V3	Negative	Culture negative	Negative	1077
118	V4	Negative	Culture negative	<i>Ralstonia insidiosa</i> , 1	1835
118	V5	Negative	Culture negative	<i>Yersinia enterocolitica</i> , 1	8570
118	EN	Negative	Culture negative	IA	IA
118	NTC	Negative	NA	<i>E. coli</i> , 1	0
120	VB	Negative	<i>S. caprae</i> , ssc	<i>E. coli</i> , 85; <i>B. subtilis</i> , 18; <i>Arthrobacter sp.</i> , 13; <i>Mucilaginibacter sp.</i> , 6; <i>B. mycoides</i> , 4; others < 4	9660
120	V2	Negative	<i>S. caprae</i> , ssc	<i>E. coli</i> , 66; <i>B. subtilis</i> , 18; <i>Arthrobacter sp.</i> , 11; <i>Mucilaginibacter sp.</i> , 3; <i>B. mycoides</i> , 2; others, < 2	28
120	V3	Negative	<i>S. caprae</i> , ssc	<i>E. coli</i> , 66; <i>B. subtilis</i> , 18; <i>Arthrobacter sp.</i> , 11;	2848

				<i>Mucilaginibacter</i> sp., 3; <i>B. mycoides</i> , 2; others < 2	
120	V4	Negative	<i>S. caprae</i> , ssc	<i>E. coli</i> , 44; <i>B. subtilis</i> , 14; <i>Arthrobacter</i> sp., 8; <i>Mucilaginibacter</i> sp., 5; <i>B. mycoides</i> , 3	220
120	V5	Negative	<i>S. caprae</i> , ssc	<i>E. coli</i> , 57; <i>B. subtilis</i> , 18; <i>Arthrobacter</i> sp., 5; <i>Mucilaginibacter</i> sp., 6; <i>B. mycoides</i> , 4; others < 4	34
120	EN	Negative	NA	<i>E. coli</i> , 7; <i>Arthrobacter</i> sp., 5; <i>C. acnes</i> 2; <i>B. subtilis</i> 1; <i>B. mycoides</i> 1, others, 1	4
120	NTC	Negative	NA		1
121	V2	Numerous GPAC;	<i>S. aureus</i> , ssc; <i>Peptoniphilus</i>	>2000 distinct species/subspecies, apparent	4998

		<i>S. aureus</i> , 984	<i>harei</i> , mod	2nd infection	
121	V3	Numerous GPAC; <i>S. aureus</i> , 1	<i>S. aureus</i> , broth; <i>Peptoniphilus harei</i> , sparse	>2000 distinct species/subspecies, apparent 2nd infection	144506
121	V4	Numerous GPAC	<i>S. aureus</i> , broth; <i>Peptoniphilus harei</i> , ssc	>2000 distinct species/subspecies, apparent 2nd infection	584
121	V6	Numerous GPAC	<i>S. aureus</i> , broth; <i>Peptoniphilus harei</i> , ssc	>2000 distinct species/subspecies, apparent 2nd infection	305
121	V8	Numerous GPAC; <i>S. aureus</i> , 1	Culture negative	>2000 distinct species/subspecies, apparent 2nd infection	27903
121	EN	Negative	NA	<i>E. coli</i> , 14; <i>S. mitis</i> , 5; <i>S. oralis</i> , 3; <i>B. subtilis</i> , 3; <i>C. acnes</i> , 3; others <3	67
121	NTC	Negative	NA	<i>E. coli</i> , 5; <i>B. subtilis</i> , 1	5
122	VB	Negative	<i>S. aureus</i> , sparse	<i>M. globosa</i> , 181; others <1	10187
122	V2	<i>S. aureus</i> , 117	<i>S. aureus</i> , ssc	<i>S. aureus</i> , 117; <i>S. parasanguinis</i> , 1	9

122	V3	<i>S. aureus</i> , 3480	<i>S. aureus</i> , rich	<i>M. globosa</i> , 154; <i>E. coli</i> , 9; <i>Mucilaginibacter sp.</i> , 5; others <5	31949
122	V4	<i>S. aureus</i> , 1269	<i>S. aureus</i> , ssc	<i>E. coli</i> , 4; others <4	1589
122	V5	<i>S. aureus</i> , 99	<i>S. aureus</i> , sparse	<i>M. globosa</i> , 27; <i>E. coli</i> , 5; others <5	2398
122	EN	Negative	NA	<i>E. coli</i> , 10; <i>M. osloensis</i> , 3; others <3	9
122	NTC	Negative	NA	Negative	
123	VB	<i>S. aureus</i> , 15	<i>S. aureus</i> , ssc	<i>E. coli</i> , 10; <i>M. globosa</i> , 14; others, 1	56
123	V2	Negative	<i>S. aureus</i> , ssc; β -hemolytic streptococci, broth	<i>M. globosa</i> , 14; <i>E. coli</i> , 2; others, 1	6646
123	V3	<i>S. aureus</i> , 2	<i>S. aureus</i> , ssc; β -hemolytic streptococci, broth	<i>E. coli</i> , 6; others, 1	36
123	V4	<i>S. aureus</i> , 8	<i>S. aureus</i> , ssc	<i>E. coli</i> , 2; <i>S. oralis</i> 2; others, 1	70
123	V5	Negative	<i>S. aureus</i> , ssc	Negative	2

123	EN	Negative	NA	<i>E. coli</i> , 2; <i>C. acnes</i> , 2; others, 1	56
123	NTC	Negative	NA	Negative	0
124	VB	Negative	<i>S. epidermidis</i> , broth; <i>Corynebacterium amycolatum</i> and <i>Neisseria subflava</i> , broth (both presumed contamination)	<i>C. acnes</i> , 2; <i>E. coli</i> , 2	12
124	V2	Negative	Culture negative	<i>C. acnes</i> , 11	7
124	V3	Negative	Culture negative	<i>C. acnes</i> , 3; <i>Propionibacterium sp.</i> , 2; <i>C. granulosum</i> , 2	0
124	V4	Negative	<i>S. epidermidis</i> , broth	<i>C. acnes</i> , 6	33
124	V5	Negative	Culture negative	Negative	0
124	EN	Negative	NA	<i>C. acnes</i> , 2	10
124	NTC	Negative	NA	Negative	NA
125	VB	Negative	<i>S. aureus</i> , ssc	<i>E. coli</i> , 2; others, 1	1
125	V2	<i>S. aureus</i> , 2	Culture negative	<i>E. coli</i> , 6; <i>M. globosa</i> , 4; others, 1	222

125	V3	Negative	<i>M. luteus</i> , broth; <i>S. aureus</i> , broth	<i>Arthrobacter sp.</i> , 1	12
125	V4	<i>S. aureus</i> , 1300	<i>M. luteus</i> , broth	<i>E. coli</i> , 5; others, <5	1943
125	EN	Negative	NA	<i>E. coli</i> , 4; others, 1	2
125	NTC	Negative	NA	Negative	0
127	VB	<i>S. aureus</i> , 4	<i>S. aureus</i> , sparse	others, 1	8
127	V2	<i>S. aureus</i> , 922;	<i>S. aureus</i> , rich	others, 2	310
127	V3	<i>S. aureus</i> , 2256	<i>S. aureus</i> , rich	others, 1	221
127	V4	<i>S. aureus</i> , 103	<i>S. aureus</i> , mod		17
127	V5	<i>S. aureus</i> , 734	<i>S. aureus</i> , rich	others, 1	30
128	VB	<i>S. aureus</i> , 9	<i>S. aureus</i> , sparse	others, 1	95
128	V2	<i>S. aureus</i> , 304	<i>S. aureus</i> , mod	others, 1	611
128	V3	<i>S. aureus</i> , 1	<i>S. aureus</i> , mod	<i>M. Osloensis</i> , 1	6
128	V4	<i>S. aureus</i> , 2	<i>S. aureus</i> , sparse	others, 1	19
128	V5	<i>S. aureus</i> , 31618	<i>S. aureus</i> , sparse	<i>S. Lugdunensis</i> , 17; others <17	2240
128	EN	Negative	NA	others, 1	1

128	NTC	Negative	NA	<i>S. Aureus</i> , 2	0
130	VB	<i>S. aureus</i> , 452	<i>S. aureus</i> , broth	<i>M. globosa</i> , 10; others 2 reads	656
130	V2	<i>S. aureus</i> , 12	<i>S. aureus</i> , broth	<i>M. globosa</i> , 18; <i>E. coli</i> , 4; others 1	1019
130	V3	<i>S. aureus</i> , 141	<i>S. aureus</i> , ssc	<i>A. oris</i> , 51; <i>V. parvula</i> , 43; <i>R. dentocariosa</i> , 34; <i>S. sanguinis</i> , 33; <i>M. globosa</i> , 7; <i>Cutibacterium sp.</i> , 31; <i>N. sicca</i> , 21; <i>F. nucleatum</i> , 9; <i>S. gordonii</i> , 5; others <5	8556
130	NTC	Negative	NA	Negative	11
133	VB	Negative	β -hemolytic streptococci, broth	<i>E. coli</i> , 9; <i>Klebsiella sp.</i> , 9; others <2	85
133	V2	<i>S. agalactiae</i> , 57	β -hemolytic streptococci, broth	<i>M. osloensis</i> , 11; <i>E. coli</i> , 5; others <5	275
133	V3	<i>S. agalactiae</i> , 131	β -hemolytic streptococci, rich	others, 1	53

133	V4	<i>S. agalactiae</i> , 30	β -hemolytic streptococci, sparse	<i>M. osloensis</i> , 5; others <5	199
133	V5	Negative	β -hemolytic streptococci, ssc	<i>E. coli</i> , 7; <i>F. nucleatum</i> , 6; others <5	37
133	EN	Negative	NA	<i>C. acnes</i> , 45; <i>S. capitis</i> , 7; others, <3	7
133	NTC	Negative	NA	<i>B. subtilis</i> , 1	0
134	V3	Negative	β -hemolytic streptococci, mod	<i>M. globosa</i> , 271; <i>Arthrobacter sp.</i> , 2; others <2	7556
134	V4	Negative	β -hemolytic streptococci, ssc	<i>E. coli</i> , 2; others <2	90
134	V5	Negative	β -hemolytic streptococci, sparse	<i>M. globosa</i> , 5; <i>E. coli</i> , 1	431
134	V6	<i>S. agalactiae</i> , 33	β -hemolytic streptococci, mod	<i>Arthrobacter sp.</i> , 2; others, <2	715
134	EN	Negative	NA	<i>E. coli</i> , 2; others <2	2
134	NTC	Negative	NA	Negative	0

135	VB	<i>S. dysgalactiae</i> , 17; <i>S. agalactiae</i> , 3; <i>S. mitis</i> , 3; <i>S. pyogenes</i> , 3	β -hemolytic streptococci, ssc	<i>M. globosa</i> , 68; <i>E. coli</i> , 18; <i>M. osloensis</i> , 11; others, 3	35999
135	V2	<i>S. dysgalactiae</i> , 240706, <i>S. pyogenes</i> , 10761; <i>S. agalactiae</i> , 8037; <i>S. anginosus</i> , 2038; <i>S. lutetiensis</i> , 410; <i>S. gallolyticus</i> , 342; <i>S. mutans</i> , 193; <i>S. gordonii</i> , 188	β -hemolytic streptococci, ssc	<i>A. mediterraneensis</i> , 883; <i>M. vaginalis</i> , 452; others, <200	1159
135	V3	Negative	Culture negative	<i>E. coli</i> , 8; <i>M. globosa</i> , 3; <i>B. subtilis</i> , 3; others, <3	631
135	V4	Negative	Culture negative	<i>M. globosa</i> , 338;	207550
135	V5	Negative	β -hemolytic streptococci,	<i>M. globosa</i> , 15; <i>E. coli</i> , 4;	7106

			ssc	<i>B. subtilis</i> , 3; others, <3	
135	EN	Negative	NA	<i>E. coli</i> , 30; <i>M. osloensis</i> , 11; <i>S. mitis</i> , 5; others, <5	475
135	NTC	Negative	NA	Negative	1
136	VB	Mitis group streptococci tot, 4; <i>S. capitis</i> , 3; <i>S. parasanguinis</i> , 2	<i>Granulicatella adiacens</i> , sparse	<i>M. globosa</i> , 68; <i>E. coli</i> , 41; <i>Arthrobacter sp.</i> , 4; others, <4	47624
136	V2	Mitis group streptococci tot, 18; staphylococci tot, 2	<i>Granulicatella adiacens</i> , sparse	<i>M. globosa</i> , 45; <i>E. coli</i> , 19, <i>B. subtilis</i> , 4; others, <4	29168
136	V3	Staphylococci tot, 1	<i>Granulicatella adiacens</i> , sparse; <i>Staphylococcus sp.</i> , broth	<i>E. coli</i> , 10; <i>M. globosa</i> , 7; <i>Arthrobacter sp.</i> , 5; others, <5	2351
136	V4	Mitis group streptococci tot, 4; staphylococci tot, 3	<i>Granulicatella adiacens</i> , ssc	<i>E. coli</i> , 21; <i>M. globosa</i> , 42; <i>Arthrobacter sp.</i> , 3; others, <3	24965
136	V5	Negative	<i>Granulicatella adiacens</i> , sparse	<i>E. coli</i> , 45; <i>M. globosa</i> , 244; <i>Arthrobacter sp.</i> , 3; <i>B. subtilis</i> , 3;	81413

				others, <3;	
136	EN	Negative	NA	<i>E. coli</i> , 4; <i>S. capitis</i> , 4; others, <4	27
136	NTC	Negative	NA	Negative	2
137	VB	Negative	Culture negative	Negative	15
137	V2	Negative	Culture negative	Negative	9
137	V3	Negative	Culture negative	Negative	2
137	V4	Negative	<i>Bacillus sp.</i> , broth (presumed contamination)	Negative	27
137	V5	Negative	Culture negative	Negative	8
137	EN	Negative	NA	Negative	1
137	NTC	Negative	NA	Negative	0
139	VB	<i>S. aureus</i> , 189	<i>S. aureus</i> , ssc	<i>E.coli</i> , 30; others <3	1277
139	V2	<i>S. aureus</i> , 12962; <i>S. argenteus</i> , 6	<i>S. aureus</i> , rich	<i>E. coli</i> , 19; others, <6	234
139	V3	<i>S. aureus</i> , 12423	<i>S. aureus</i> , rich	<i>E. coli</i> , 25; <i>C. acnes</i> , 7; others, <7	638

139	V4	<i>S. aureus</i> , 7362	<i>S. aureus</i> , sparse	<i>E. coli</i> , 18; <i>M. globosa</i> , 4; others, < 4	3317
139	V5	<i>S. aureus</i> , 18	<i>S. aureus</i> , sparse	<i>E. coli</i> , 18; <i>C. acnes</i> , 5; others, <5	40
139	EN	Negative	NA	<i>E. coli</i> , 9; others, 1	14
139	NTC	Negative	NA	<i>B. subtilis</i> , 2; <i>E. coli</i> , 1; <i>Arthrobacter sp.</i> , 1	1
140	VB	<i>S. aureus</i> , 714226; <i>S. argenteus</i> , 122; <i>S. epidermidis</i> , 108; <i>S. lugdunensis</i> , 94; <i>Staphylococcus sp.</i> , <50*	<i>S. aureus</i> , rich	<i>Bacillus sp.</i> , 12; others, < 10	2785
140	VB	<i>S. aureus</i> , 661866; <i>S. argenteus</i> , 142; <i>S. epidermidis</i> , 57; <i>S. lugdunensis</i> , 70; <i>Staphylococcus sp.</i> , <50*	<i>S. aureus</i> , rich	<i>B. cerues.</i> , 16; <i>B. thuringiensis</i> , 14; others, < 10	653

140	V2	<i>S. aureus</i> , 776139; <i>S. argenteus</i> , 175; <i>S. epidermidis</i> , 93; <i>S. lugdunensis</i> , 106; <i>Staphylococcus sp.</i> , <50*	<i>S. aureus</i> , rich	<i>B. cereus</i> , 38; others, <10	372
140	V3	<i>S. aureus</i> , 18112; <i>Staphylococcus sp.</i> , <50*	<i>S. aureus</i> , rich	<i>E. coli</i> , 3; others, 1	382
140	V4	<i>S. aureus</i> , 18417; <i>Staphylococcus sp.</i> , <50*	<i>S. aureus</i> , rich	<i>E. coli</i> , 4; <i>M. globosa</i> , 3; others, <2	527
140	EN	Negative	NA	<i>S. aureus</i> , 2; <i>S. argenteus</i> , 5; <i>E. coli</i> , 2; <i>C. acnes</i> , 2; others, <2	1
140	NTC	Negative	NA	<i>S. aureus</i> , 2; <i>B. subtilis</i> , 2; <i>E. coli</i> , 2; <i>Mucilaginibacter sp.</i> , 1	0
141	VB	<i>S. aureus</i> , 2123024; <i>S. lugdunensis</i> , 542; <i>S. epidermidis</i> , 431;	<i>S. aureus</i> , rich	<i>M. globosa</i> , 205; <i>B. thuringiensis</i> , 114; <i>Bacillus sp.</i> , 72;	160982

		<i>S. argenteus</i> , 282; <i>P. pseudointermedius</i> , 252; <i>S. schleiferi</i> , 112; <i>S. stepanovicii</i> , 100; <i>S. simiae</i> , 99; <i>S. capitis</i> , 73		<i>E. coli</i> , 52; many others, <50	
141	V2	<i>S. aureus</i> , 1173	<i>S. aureus</i> , rich	<i>M. globosa</i> , 48; <i>E. coli</i> , 20; <i>C. acnes</i> , 11; others, <10	32184
141	V3	<i>S. aureus</i> , 214625; <i>S. lugdunensis</i> , 63	<i>S. aureus</i> , rich	many, < 50	1820
141	V4	<i>S. aureus</i> , 13941	<i>S. aureus</i> , rich	<i>M. globosa</i> , 58; <i>E. coli</i> , 34; others, <10	23033
141	V5	<i>S. aureus</i> , 31057	<i>S. aureus</i> , rich	<i>E. coli</i> , 24; others < 20	432
141	EN	Negative	NA	<i>S. aureus</i> , 78; <i>E. coli</i> , 16; <i>Arthrobacter sp.</i> , 5; others, <5	12
141	NTC	Negative	NA	<i>S. aureus</i> , 5; <i>E. coli</i> , 1;	0

				<i>Alkalilimnicaola ehrlichii</i> , 1	
142	VB	<i>S. epidermidis</i> , 8; <i>S. aureus</i> , 7	Culture negative	<i>C. acnes</i> , 128; <i>E. coli</i> , 61; <i>M. globosa</i> , 28; <i>B. subtilis</i> , 17; <i>Arthrobacter sp.</i> , 9; <i>Mucilaginibacter sp.</i> , 4; <i>S. capitis</i> , 3; <i>P. ananatis</i> , 2; <i>M. osloensis</i> , 2; others <2	24252
142	V2	<i>S. epidermidis</i> , 3	Culture negative	<i>C. acnes</i> , 56; <i>E. coli</i> , 57; <i>M. globosa</i> , 4; <i>B. subtilis</i> , 10; <i>Arthrobacter sp.</i> , 11; <i>Mucilaginibacter sp.</i> , 4; <i>B. mycoides</i> , 3; <i>S. mitis</i> , 3	959
142	V3	<i>S. oralis</i> , 4	Culture negative	<i>C. acnes</i> , 21; <i>E. coli</i> , 48; <i>M. globosa</i> , 20; <i>B. subtilis</i> , 5; <i>Arthrobacter sp.</i> , 3	10883
142	V4	Negative	Culture negative	<i>C. acnes</i> , 4; <i>E. coli</i> , 76;	518

				<i>M. globosa</i> , 20; <i>B. subtilis</i> , 25; <i>Arthrobacter sp.</i> , 13; <i>B. mycoides</i> , 4; <i>Mucilaginibacter sp.</i> , 2	
142	V5	<i>S. oralis</i> , 4;	Culture negative	<i>C. acnes</i> , 6; <i>E. coli</i> , 46; <i>B. subtilis</i> , 19; <i>Arthrobacter sp.</i> , 15; <i>B. mycoides</i> , 3; <i>Mucilaginibacter sp.</i> , 4	198
142	EN	Negative	NA	<i>C. acnes</i> , 8; <i>E. coli</i> , 22; <i>B. subtilis</i> , 4; <i>Arthrobacter sp.</i> , 6; <i>Mucilaginibacter sp.</i> , 2; <i>B. mycoides</i> , 2	0
142	NTC	Negative	NA		0
200	VB	Negative control	Culture negative	<i>E. coli</i> , 37; <i>Arthrobacter sp.</i> , 13; <i>M. osloensis</i> , 13; <i>B. subtilis</i> , 12;	843

				<i>C. acnes</i> , 9 <i>M. globosa</i> , 7; <i>Mucilaginibacter sp.</i> , 3	
200	V2	Negative control	Culture negative	<i>E. coli</i> , 68; <i>M. globosa</i> , 34; <i>Arthrobacter sp.</i> , 17; <i>B. subtilis</i> , 12; <i>Mucilaginibacter sp.</i> , 7; <i>M. osloensis</i> , 5; <i>C. acnes</i> , 5; <i>B. mycoides</i> , 3; <i>Pantoa ananatis</i> , 3; <i>S. gordonii</i> , 3	35266
200	V3	Negative control	Culture negative	<i>E. coli</i> , 40; <i>Arthrobacter sp.</i> , 7; <i>M. osloensis</i> , 3; <i>B. subtilis</i> , 4; <i>C. acnes</i> , 3; <i>M. globosa</i> , 32;	14882
200	V4	Negative control	Culture negative	<i>E. coli</i> , 37; <i>Arthrobacter sp.</i> , 6; <i>B. subtilis</i> , 13; <i>Mucilaginibacter sp.</i> , 7; <i>M. osloensis</i> , 5; <i>B. mycoides</i> , 3;	

200	V5	Negative control	Culture negative	<i>E.coli</i> , 29; <i>Arthrobacter sp.</i> , 20; <i>M. osloensis</i> , 4; <i>B. subtilis</i> , 10; <i>C. acnes</i> , 6; <i>M. globosa</i> , 14; <i>Mucilaginibacter sp.</i> , 4	10020
200	EN	Negative control	NA	Negative	
200	NTC	Negative control	NA	<i>E. coli</i> , 5;	5
201	VB	Negative control	Culture negative	others, <3	254
201	V2	Negative control	Culture negative	<i>M. globosa</i> , 3	450
201	V3	Negative control	Culture negative	others, <3	141
201	V4	Negative control	Culture negative	others, <3	97
201	V5	Negative control	Culture negative	others, <3	41
201	EN	Negative control	NA	others, <3	1
201	NTC	Negative control	NA	Negative	0
202	VB	Negative control	Culture negative	<i>E. coli</i> , 3; <i>M. globosa</i> , 4;	1783

202	V2	Negative control	Culture negative	<i>M. globosa</i> , 7;	510
202	V3	Negative control	Culture negative	<i>E. coli</i> , 12; <i>M. globosa</i> , 21;	12678
202	V4	Negative control	Culture negative	<i>E. coli</i> , 8	217
202	V5	Negative control	Culture negative	<i>E. coli</i> , 13; <i>M. globosa</i> , 40;	25376
202	EN	Negative control	NA	<i>E. coli</i> , 6	4
202	NTC	Negative control	NA	Negative	0
203	VB	Negative control	Culture negative	Negative	5
203	V2	Negative control	Culture negative	Negative	6
203	V3	Negative control	Culture negative	<i>M. globosa</i> , 318;	5599
203	V4	Negative control	Culture negative	<i>M. osloensis</i> , 3	3886
203	V5	Negative control	Culture negative	<i>M. globosa</i> , 23; <i>E. coli</i> , 3	1891
203	EN	Negative control	NA	Negative	0
203	NTC	Negative control	NA	Negative	0
204	VB	Negative control	Culture negative	<i>E. coli</i> , 7; <i>M. globosa</i> , 75; <i>Arthrobacter sp.</i> , 2;	32185

				<i>K. pneumoniae</i> , 1; <i>M. osloensis</i> , 11	
204	V2	Negative control	Culture negative	<i>E. coli</i> , 15; <i>M. globosa</i> , 16; <i>Arthrobacter sp.</i> , 1; <i>C. acnes</i> , 3; <i>S. capitis</i> , 3; <i>Klebsiella pneumoniae</i> 3 <i>M. osloensis</i> , 17; <i>Acinetobacter baumannii</i> , 2	20967
204	V3	Negative control	Culture negative	<i>E. coli</i> , 2; <i>M. globosa</i> , 727; <i>Arthrobacter sp.</i> , 2	13571
204	V4	Negative control	Culture negative	<i>E. coli</i> , 4; <i>M. globosa</i> , 132; <i>Arthrobacter sp.</i> , 2	6102
204	V5	Negative control	Culture negative	<i>E. coli</i> , 4; <i>B. subtilis</i> , 2; <i>Mucilaginibacter sp.</i> , 2	210
204	EN	Negative control	NA	<i>M. osloensis</i> , 10; <i>E. coli</i> , 3	14
204	NTC	Negative control	NA	<i>E. coli</i> , 1; <i>B. subtilis</i> , 1	0

**S. pseudointermedius*, *S. schleiferi*, *S. capitis*, *S. stepanovicii*, *S. simiae*, *S. pasteurii*, *S. simulans*, *S. lutrae*, *S. saprophyticus*, *S. nepalensis*, *S. piscifermentans*, *S. haemolyticus*, *S. pettenkoferi*, *S. sciuri*, *S. xylosus*, *S. argenteus*

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28 **Table S2. Bioinformatic program versions used.**

Program	Version
MinKNOW	1.15.4 to v. 3.6.5
Guppy	3.0.3 to v. 3.2.10

EPI2ME	2.57.1769546 to v. 2019.7.9.2549693
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31 **Table S3. Nanopore sequencing QC metrics, totals for each patient.**

Patient ID	Total reads	Total bases	Mean quality score	Mean read length (bp)
101	4567	14.1 Mb	10.7	3084
104	52607	144.1 Mb	12.84	2739
105	84234	263.3 Mb	12.1	3126
107	9404	17.5 Mb	9.59	1863
108	60125	141.6 Mb	12.51	2354
109	385891	1.3 Gb	12.74	3296
110	44309	124.6 Mb	12.54	2811
111	394935	1.1 Gb	12.46	2779
112	21750	60.9 Mb	12.48	2800
114	44309	124.6 Mb	12.54	2811

115	17165	41.7 Mb	12.17	2430
116	22074	61.4 Mb	11.84	2780
117	1463935	3.8 Gb	8.07	2621
118	73076	89.3 Mb	5.66	1222
120	81223	345.9 Mb	11.77	4258
121	458946	1.9 Gb	12.8	4186
122	56248	122.2 Mb	11.56	2172
123	7750	20.5 Mb	11.84	2647
124	758	560.9 Kb	8.12	739
125	4423	10.8 Mb	11.86	2445
127	71026	235.4 Mb	10.31	3313
128	37382	111.4 Mb	12.38	2980
130	3880728	14.5 Gb	10.53	3738
133	1536	3.2 Mb	11.46	2091
134	10624	22.4 Mb	11.63	2107
135	535666	2 Gb	12.41	3686

136	195174	604.9 Mb	12.53	3099
137	13690	14.6 Mb	4.46	1069
139	41736	139.9 Mb	12.54	3351
140	2273296	9.8 Gb	12.87	4324
141	2744512	11.9 Gb	12.89	4335
142	30010	103.3 Mb	12.44	3441
200	67637	162.5 Mb	12.59	2402
201	2288	2.1 Mb	9.99	931
202	43111	97.8 Mb	12.39	2267
203	15396	30.8 Mb	11.42	2002
204	79478	198.7 Mb	12.26	2499

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34 **Table S4. Overview of all patients whose biopsies were positive for either an antimicrobial resistance (AMR) phenotype from**
 35 **conventional antibiotic susceptibility testing (AST) or AMR genotype from shotgun metagenomic sequencing.** Definitions used
 36 to categorize the isolates' AST results were (S) sensitive, intermediate (I) or resistant (R). Identity cut-off for AMR genotype match
 37 was set at 90%. *Other* indicates detected AMR genes not reflected in AST phenotype, or putative AMR genes of uncertain function.
 38 *None detected* indicates that patient's biopsies were sequencing positive, but no AMR genes were detected. *Sequencing negative*
 39 indicates that sequencing failed to detect the cultured pathogen, and hence its AMR gene(s).

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ID	AST (phenotype)	AMR Genotype (reads)
104	Find 1: <i>S. lugdunensis</i>	None detected
	Ciprofloxacin (S)	
	Erythromycin (S)	
	Fusidic acid (S)	
	Gentamicin (S)	
	Clindamycin (S)	
	Chloramphenicol (S)	
	Linezolid (S)	
	Oxacillin (S)	
	Rifampicin (S)	
	Penicillin (R)	
	Tetracycline (S)	
	Trimethoprim/sulfamethoxazole (S)	
	Find 2: <i>C. acnes</i>	

105	Clindamycin (S)	
	Chloramphenicol (S)	
	MeroPenicillin em (S)	
	Metronidazole (R)	
	Penicillin	
	Find 1: <i>S. aureus</i>	Sequencing negative
	Ciprofloxacin (S)	
	Erythromycin (S)	
	Fusidic acid (R)	
	Gentamicin (S)	
	Clindamycin (S)	
	Chloramphenicol (S)	
	Linezolid (S)	
	Oxacillin (S)	
	Rifampicin (S)	
	Penicillin (R)	
	Teicoplanin (S)	
	Tetracycline (S)	
	Trimethoprim/sulfamethoxazole (S)	
	Vancomycin (S)	
	Find 2: <i>S. epidermidis</i>	
	Ciprofloxacin(S)	
	Erythromycin (S)	

	Fusidic acid (R)	
	Gentamicin (S)	
	Clindamycin (S)	
	Chloramphenicol (S)	
	Linezolid (S)	
	Oxacillin (S)	
	Rifampicin (S)	
	Teicoplanin (S)	
	Tetracycline (S)	
	Trimethoprim/sulfamethoxazole (S)	
	Vancomycin (S)	
107	Find: <i>S. epidermidis</i>	Sequencing negative
	Ciprofloxacin (R)	
	Erythromycin (R)	
	Fusidic acid (R)	
	Gentamicin (S)	
	Clindamycin (R)	
	Chloramphenicol (S)	
	Linezolid (S)	
	Oxacillin (R)	
	Rifampicin (S)	
	Teicoplanin (S)	
	Tetracycline (S)	

	Trimethoprim/sulfamethoxazole (R)	
	Vancomycin (S)	
108	Find: <i>S. aureus</i>	
	Ciprofloxacin(S)	
	Erythromycin (S)	
	Fusidic acid (S)	
	Gentamicin (S)	
	Clindamycin (S)	
	Chloramphenicol (S)	
	Linezolid (S)	
	Oxacillin (S)	
	Rifampicin (S)	
	Penicillin (R)	<i>blaZ</i> (26)
	Tetracycline (S)	
	Trimethoprim/sulfamethoxazole (S)	
	other	<i>sav1866</i> (7)
109	Find: <i>S. aureus</i>	None detected
	Ciprofloxacin (S)	
	Erythromycin (S)	
	Fusidic acid (S)	
	Gentamicin (S)	
	Clindamycin (S)	
	Chloramphenicol (S)	

	Linezolid (S)	
	Oxacillin (S)	
	Rifampicin (S)	
	Penicillin (R)	
	Tetracycline (S)	
	Trimethoprim/sulfamethoxazole (S)	
111	Find: <i>S. aureus</i> (S)	
	Ciprofloxacin (S)	
	Erythromycin (S)	
	Fusidic acid (S)	
	Gentamicin (S)	
	Clindamycin (S)	
	Chloramphenicol (S)	
	Linezolid (S)	
	Oxacillin (S)	
	Rifampicin (S)	
	Penicillin (R)	<i>blaZ</i> (3390)
	Tetracycline (S)	
	Trimethoprim/sulfamethoxazole (S)	
	other	<i>tet38</i> (480), <i>sav1866</i> (394), <i>mepA</i> (376), <i>mgrA</i> (296), <i>arlS</i> (281), <i>mepR</i> (94)
112	Find: <i>S. aureus</i>	
	Ciprofloxacin (S)	

	Erythromycin (S)	
	Fusidic acid (S)	
	Gentamicin (S)	
	Clindamycin (S)	
	Linezolid (S)	
	Oxacillin (S)	
	Rifampicin (S)	
	Penicillin (R)	
	Tetracycline (S)	
	Trimethoprim/sulfamethoxazole (S)	
	other	<i>sav1866</i> (15), <i>tet38</i> (15), <i>mgrA</i> (10), <i>arlS</i> (7), <i>mepA</i> (5), <i>mepR</i> (5)
114	Find: <i>S. aureus</i>	
	Ciprofloxacin (S)	
	Erythromycin (S)	
	Fusidic acid (S)	
	Gentamicin (S)	
	Clindamycin (S)	
	Linezolid (S)	
	Oxacillin (S)	
	Rifampicin (S)	
	Penicillin (R)	<i>blaZ</i> (943)
	Tetracycline (S)	

	Trimethoprim/sulfamethoxazole (S)	
	Other	<i>blaZ</i> (943) <i>tet38</i> (2403), <i>sav1866</i> (1622), <i>mgrA</i> (1257), <i>arlS</i> (1154), <i>arlR</i> (344), <i>mepR</i> (338)
116	Find: <i>S. aureus</i>	
	Ciprofloxacin(S)	
	Erythromycin (S)	
	Fusidic acid (S)	
	Gentamicin (S)	
	Clindamycin (S)	
	Linezolid (S)	
	Oxacillin (S)	
	Rifampicin (S)	
	Penicillin (R)	<i>blaZ</i> (156), <i>bla</i> _{TEM-4} (55)
	Tetracycline (S)	
	Trimethoprim/sulfamethoxazole (S)	
	Other	<i>sav1866</i> (17), <i>tetC</i> (813) <i>bla</i> _{TEM-4} (55), <i>mepA</i> (10), <i>arlS</i> (19), <i>mgrA</i> (8)
117	Find: <i>S. aureus</i>	
	Ciprofloxacin (R)	<i>mgrA</i> , 1 (87% match)
	Erythromycin (S)	
	Fusidic acid (S)	
	Gentamicin (S)	

	Clindamycin (S)	
	Linezolid (S)	
	Oxacillin (S)	
	Rifampicin (S)	
	Penicillin (R)	
	Tetracycline (S)	
	Trimethoprim/sulfamethoxazole (S)	
120	Find: <i>S. caprae</i>	Sequencing negative
	Ciprofloxacin (S)	
	Erythromycin (S)	
	Fusidic acid (R)	
	Gentamicin (S)	
	Clindamycin (S)	
	Linezolid (S)	
	Oxacillin (S)	
	Rifampicin (S)	
	Teicoplanin (S)	
	Tetracycline (S)	
	Trimethoprim/sulfamethoxazole (S)	
	Vancomycin (S)	
121	Find: <i>S. aureus</i>	None detected
	Ciprofloxacin (S)	
	Erythromycin (S)	

	Fusidic acid (S)	
	Gentamicin (S)	
	Clindamycin (S)	
	Linezolid (S)	
	Oxacillin (S)	
	Rifampicin (S)	
	Penicillin (R)	
	Tetracycline (S)	
	Trimethoprim/sulfamethoxazole (S)	
122	Find: <i>S. aureus</i>	
	Ciprofloxacin (S)	
	Erythromycin (S)	
	Fusidic acid (S)	
	Gentamicin (S)	
	Clindamycin (S)	
	Linezolid (S)	
	Oxacillin (S)	
	Rifampicin (S)	
	Penicillin (R)	<i>blaZ</i> (8)
	Tetracycline (S)	
	Trimethoprim/sulfamethoxazole (S)	
	other	<i>tet38</i> (12), <i>sav1866</i> (9), <i>mgrA</i> (6)
123	Find 1: <i>S. aureus</i>	None detected

	Ciprofloxacin (S)	
	Erythromycin (S)	
	Fusidic acid (S)	
	Gentamicin (S)	
	Clindamycin (S)	
	Linezolid (S)	
	Oxacillin (S)	
	Rifampicin (S)	
	Penicillin (R)	
	Tetracycline (S)	
	Trimethoprim/sulfamethoxazole (S)	
	Find 2: β-hemolytic streptococci	
	Erythromycin (S)	
	Clindamycin (S)	
	Oxacillin (S)	
	Penicillin (S)	
	Trimethoprim/sulfamethoxazole (S)	
	Vancomycin (S)	
124	Find: <i>S. epidermidis</i>	Sequencing negative
	Ciprofloxacin (R)	
	Erythromycin (R)	
	Fusidic acid (S)	
	Gentamicin (R)	

	Clindamycin (R)	
	Linezolid (S)	
	Oxacillin (R)	
	Rifampicin (S)	
	Teicoplanin (S)	
	Tetracycline (S)	
	Trimethoprim/sulfamethoxazole (R)	
	Vancomycin (S)	
125	Find: <i>S. aureus</i>	
	Ciprofloxacin (S)	
	Erythromycin (S)	
	Fusidic acid (S)	
	Gentamicin (S)	
	Clindamycin (S)	
	Linezolid (S)	
	Oxacillin (S)	
	Rifampicin (S)	
	Penicillin (R)	<i>blaZ</i> (2)
	Tetracycline (S)	
	Trimethoprim/sulfamethoxazole (S)	
	other	<i>mepA</i> (5), <i>sav1866</i> (3), <i>tet38</i> (2)
127	Find: <i>S. aureus</i>	
	Ciprofloxacin (R)	<i>arlS</i> (11), <i>mgrA</i> (6),

	Erythromycin (S)	
	Fusidic acid (S)	
	Gentamicin (S)	
	Clindamycin (S)	
	Linezolid (S)	
	Oxacillin (S)	
	Rifampicin (S)	
	Penicillin (S)	
	Tetracycline (S)	
	Trimethoprim/sulfamethoxazole (S)	
	other	<i>sav1866</i> (8)
128	Find: <i>S. aureus</i>	
	Ciprofloxacin (I)	<i>mgrA</i> (34), <i>arlS</i> (40), <i>arlR</i> (9)
	Erythromycin (S)	
	Fusidic acid (S)	
	Gentamicin (S)	
	Clindamycin (S)	
	Linezolid (S)	
	Oxacillin (S)	
	Rifampicin (S)	
	Penicillin (R)	<i>blaZ</i> (28)

	Tetracycline (R)	<i>tetK</i> (28), <i>tet38</i> (50), <i>sav1866</i> (52), <i>mepR</i> (12), <i>mepA</i> (36), <i>mgrA</i> (34)
	Trimethoprim/sulfamethoxazole (S)	
130	Find: <i>S. aureus</i>	
	Ciprofloxacin (S)	
	Erythromycin (S)	
	Fusidic acid (S)	
	Gentamicin (S)	
	Clindamycin (S)	
	Linezolid (S)	
	Oxacillin (S)	
	Rifampicin (S)	
	Penicillin (R)	
	Tetracycline (S)	
	Trimethoprim/sulfamethoxazole (S)	
	other	<i>mepA</i> (2)
135	Find: B-hemolytic streptococci	
	Erythromycin (R) , Clindamycin (R)	<i>lmrP</i> , 620 (88.8% match)
	Penicillin (S)	
139	Find: <i>S. aureus</i>	
	Ciprofloxacin (S)	
	Erythromycin (S)	
	Fusidic acid (S)	

	Gentamicin (S)	
	Clindamycin (S)	
	Linezolid (S)	
	Oxacillin (S)	
	Rifampicin (S)	
	Penicillin (R)	<i>blaZ</i> (45)
	Tetracycline (S)	
	Trimethoprim/sulfamethoxazole (S)	
	other	<i>tet38</i> (95), <i>mepA</i> (56), <i>sav1866</i> (53), <i>arlS</i> (36), <i>mepR</i> (9), <i>arlR</i> (7)
140	Find: <i>S. aureus</i>	
	Ciprofloxacin (S)	
	Erythromycin (S)	
	Fusidic acid (S)	
	Gentamicin (S)	
	Clindamycin (S)	
	Linezolid (S)	
	Oxacillin (S)	
	Rifampicin (S)	
	Penicillin (R)	<i>blaZ</i> (3392)
	Tetracycline (S)	
	Trimethoprim/sulfamethoxazole (S)	

	other	<i>tet38</i> (5251); <i>sav1866</i> (4451); <i>mepA</i> (4397); <i>mgrA</i> (3580); <i>arlS</i> (3521); <i>arlR</i> (856); <i>mepR</i> (773)
141	Find: <i>S. aureus</i>	
	Ciprofloxacin (S)	
	Erythromycin (S)	
	Fusidic acid (S)	
	Gentamicin (S)	
	Clindamycin (S)	
	Linezolid (S)	
	Oxacillin (S)	
	Rifampicin (S)	
	Penicillin (R)	<i>blaZ</i> (3172)
	Tetracycline (S)	
	Trimethoprim/sulfamethoxazole (S)	
	other	<i>tet38</i> (6479), <i>sav1866</i> (4362), <i>mepA</i> (5456), <i>mgrA</i> (3431), <i>arlS</i> (3106), , <i>arlR</i> (754), <i>mepR</i> (1010)