

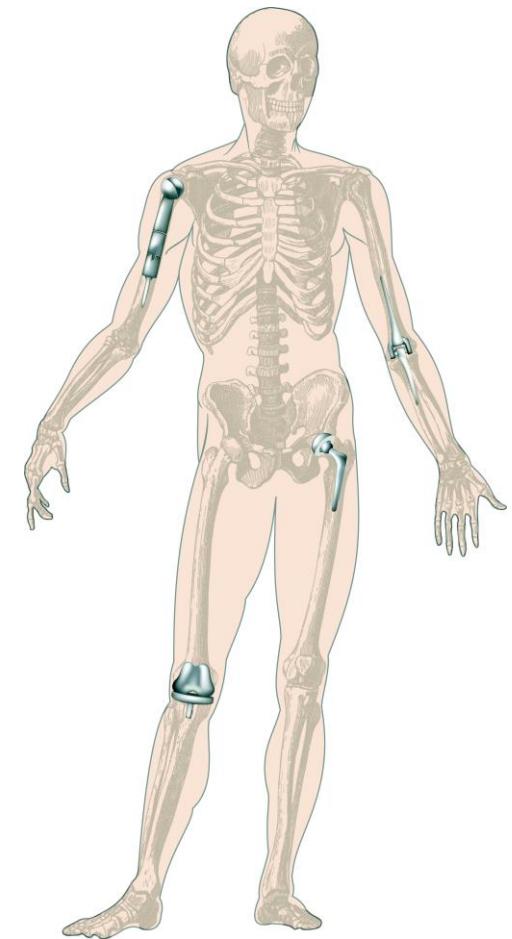
New Approaches to the Diagnosis of Prosthetic Joint Infection

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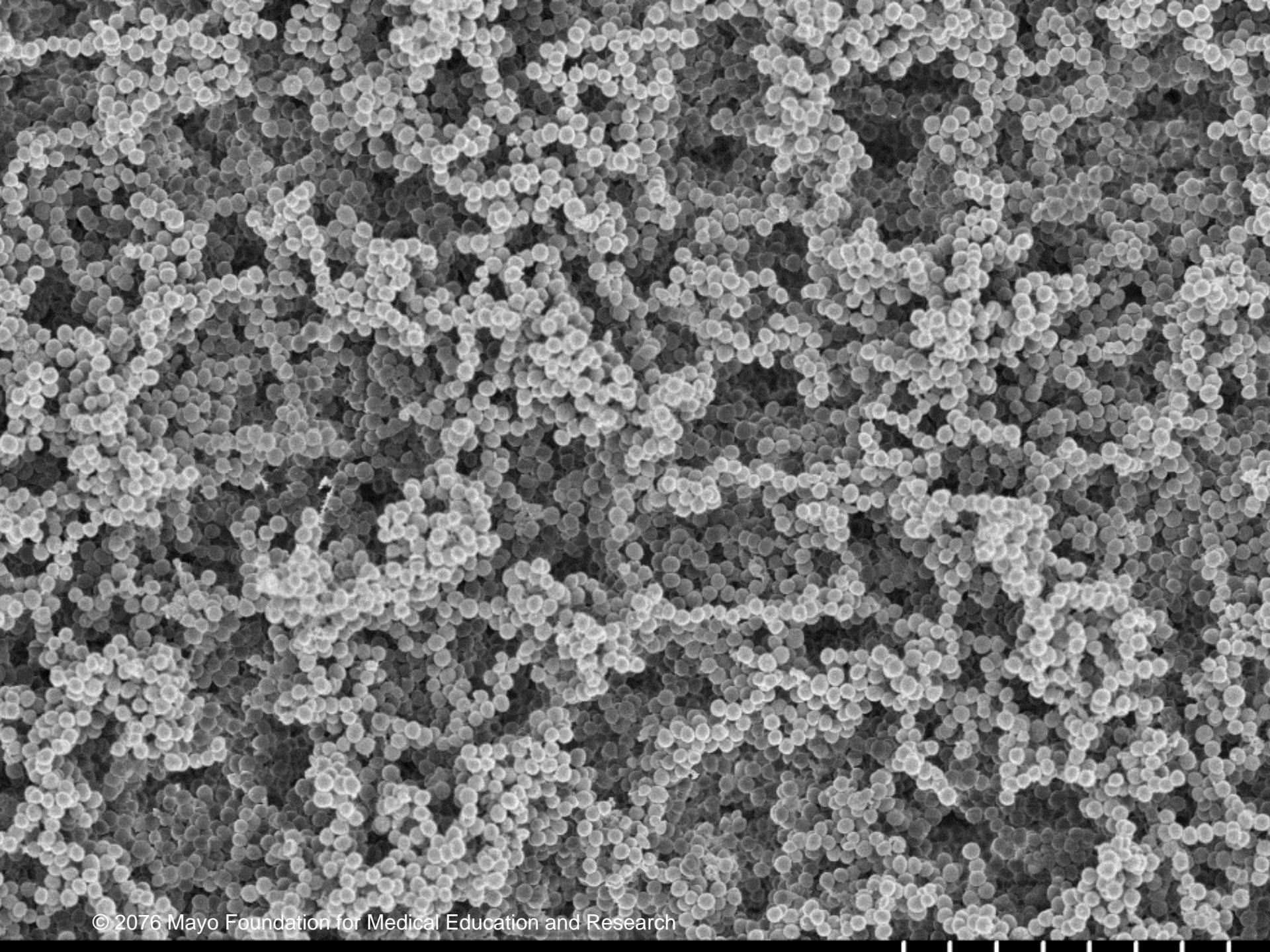


Disclosures

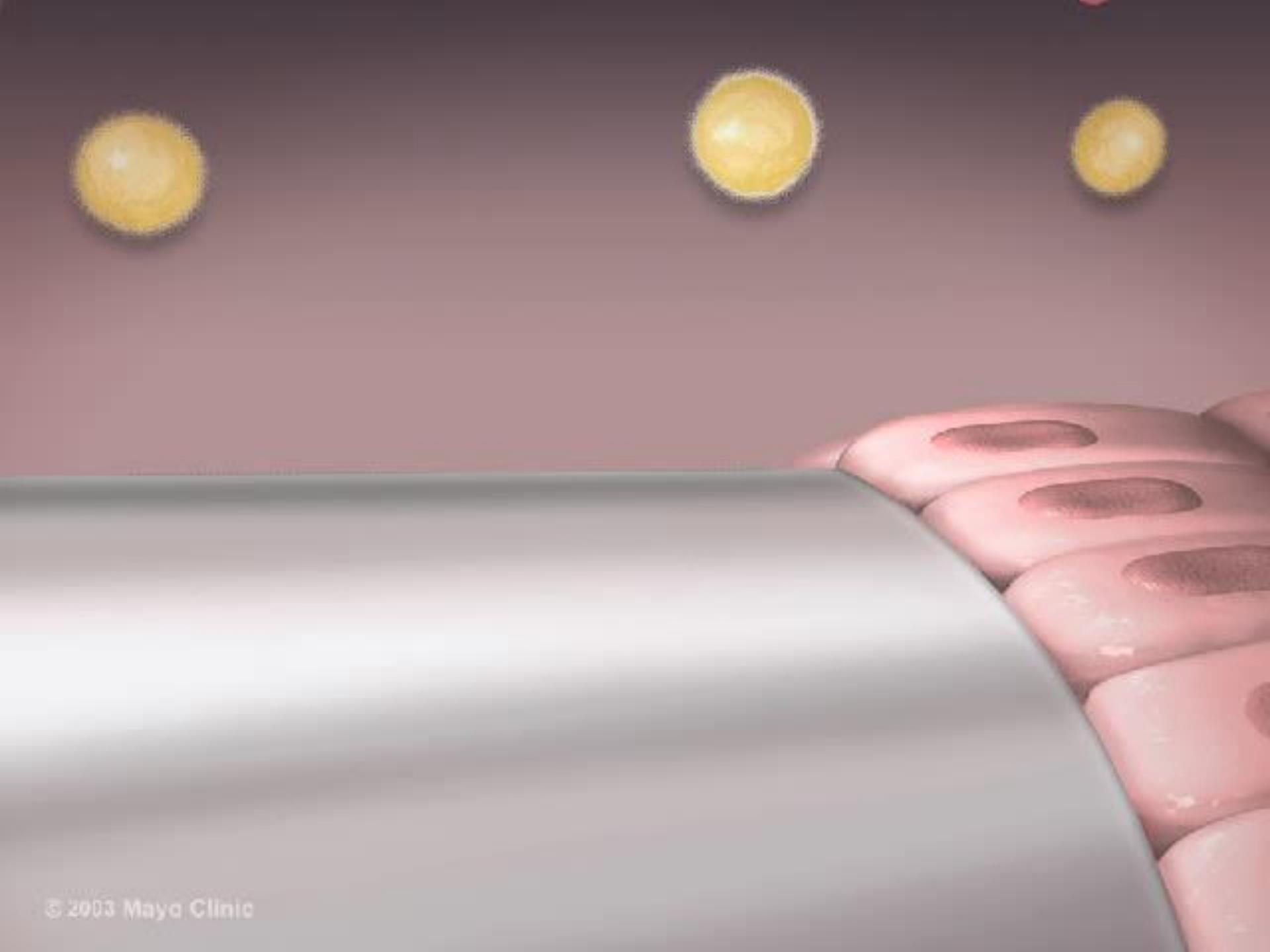
- Employee: Mayo Clinic
- Grants: CD Diagnostics, BioFire, Check-Points, Curetis, 3M, Merck, Hutchison Biofilm Medical Solutions, Accelerate Diagnostics, Allergan, The Medicines Company, Zimmer, Specific Technologies, EnBiotix
- Consultant: Curetis and Qvella (paid to Mayo Clinic)
- Patents: *Bordetella pertussis/parapertussis* PCR issued, device/method for sonication with royalties (paid by Samsung to Mayo Clinic), anti-biofilm substance issued
- Data safety monitoring board: Actelion
- Travel reimbursement/editor's stipends: ASM & IDSA
- Honoraria: NBME, Up-to-Date, Infectious Diseases Board Review Course

Objectives

1. Learn the epidemiology, pathogenesis, and microbiology of prosthetic joint infection
2. Interpret synovial fluid biomarkers in patients with a prosthetic joint
3. Determine appropriate specimen types for microbial diagnostics for prosthetic joint infection

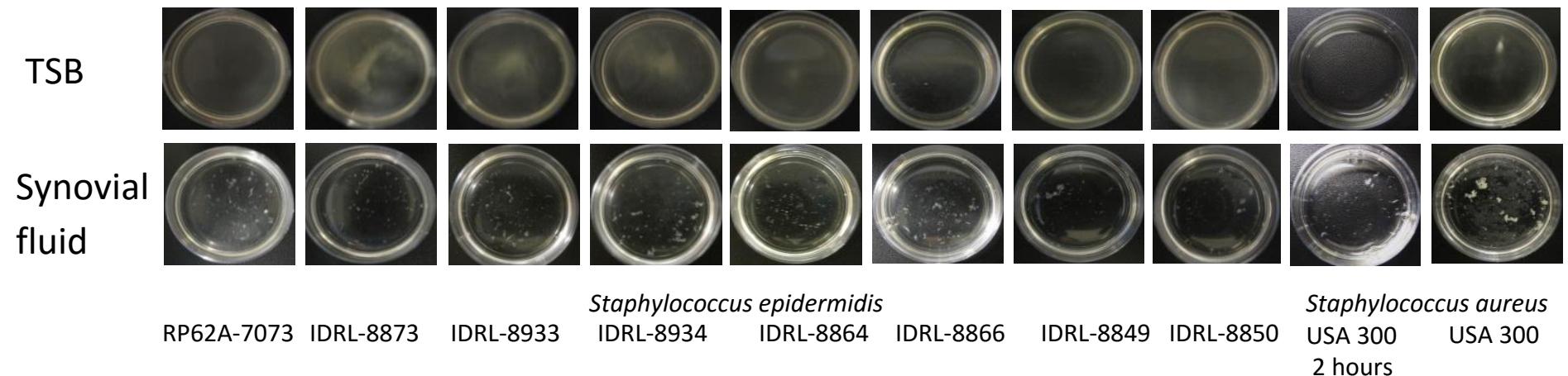


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Biofilm-Like Aggregation

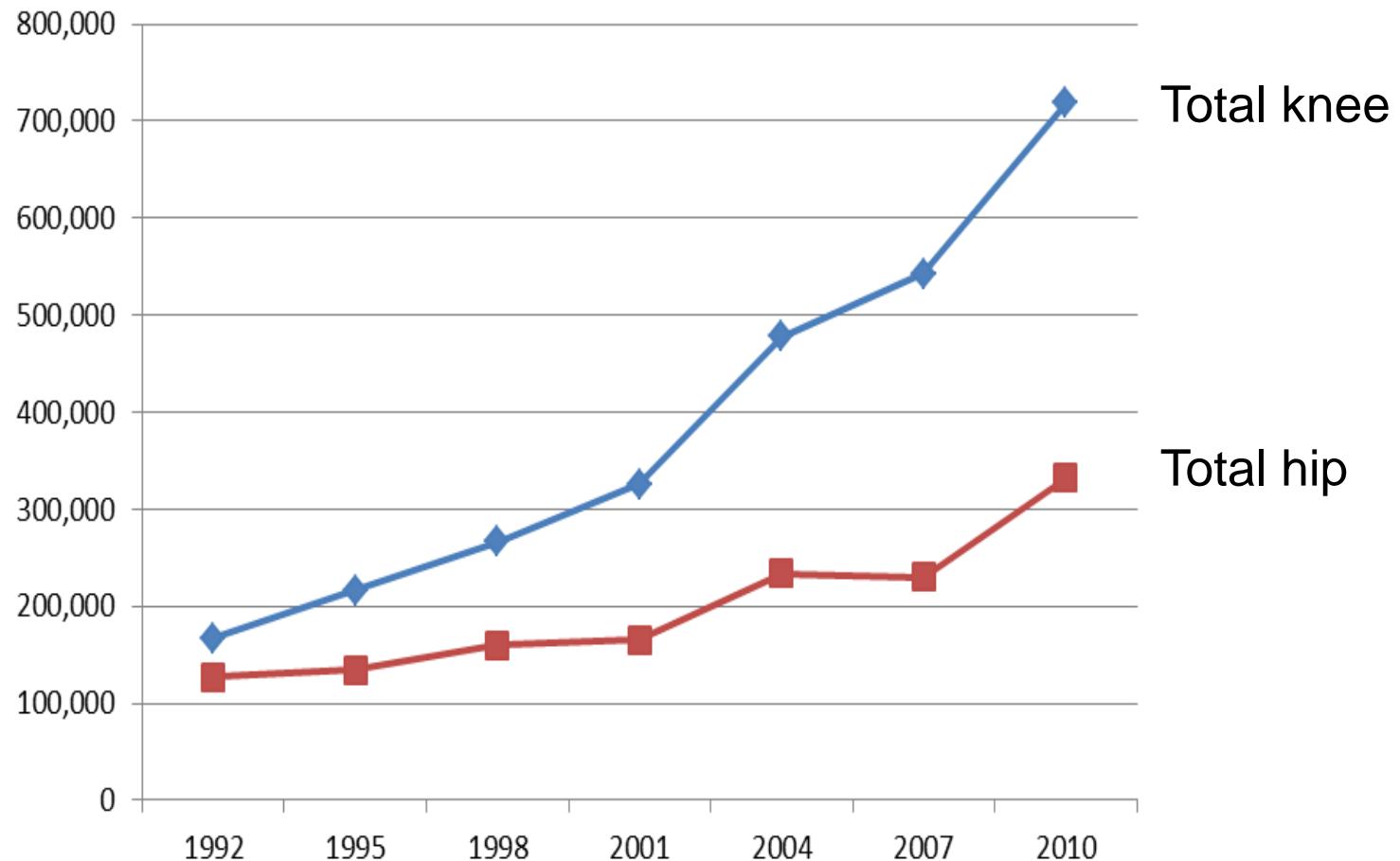
Staphylococcus epidermidis,
Staphylococcus aureus in Synovial Fluid



Total Hip & Knee Replacement Procedures

United States

National Hospital Discharge Survey <http://www.cdc.gov/nchs/about/major/hdasd/listpubs.htm>



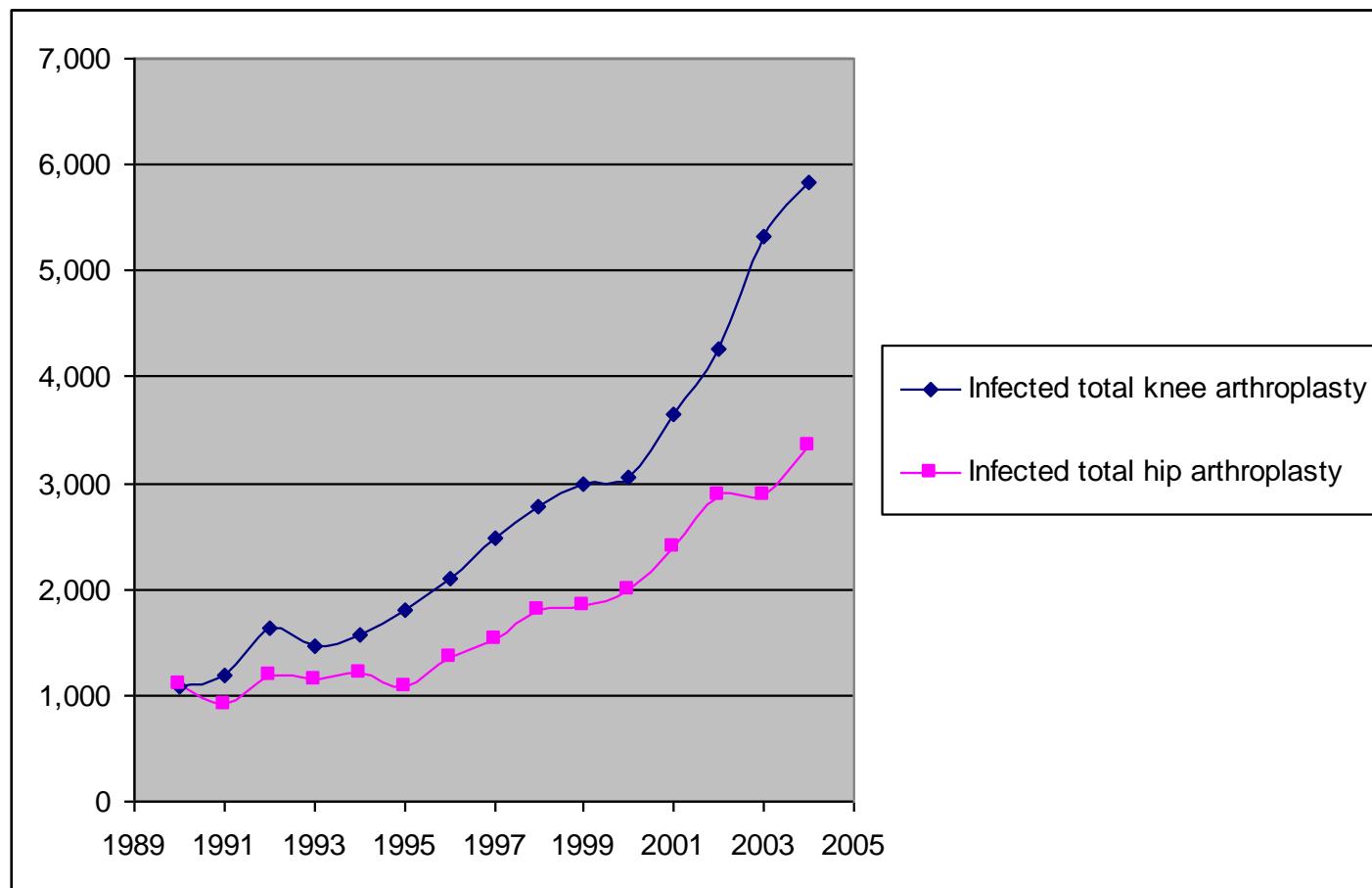
Prosthetic Joint Infection Rates

	Years*	
	0-2	2-10
Prosthetic knee infection	1.55%	0.46%
Prosthetic hip infection	0.78%	0.33%

*Medicare Population
Primary Elective Arthroplasty
1997-2006

Kurtz et al. Clin Orthop Relat Res 2010;468:52
Ong et al. J Arthroplasty 2009;24:105

Prosthetic Hip and Knee Infections United States



Kurtz et al. J Arthroplasty 2008;23:984



Prosthetic Joint Infection Microbiology

	Hip and Knee		Hip	Knee	Shoulder	Elbow
	All time periods	Early				
Number of joints	2435	637	1979	1427	199	110
<i>Staphylococcus aureus</i>	27	38	13	23	18	42
Coagulase negative staphylococci	27	22	30	23	41	41
<i>Streptococcus</i> species	8	4	6	6	4	4
<i>Enterococcus</i> species	3	10	2	2	3	0
Aerobic gram negative bacilli	9	24	7	5	10	7
Anaerobic bacteria	4	3	9	5		
<i>Propionibacterium acnes</i>					24	1
Other anaerobes					3	0
Culture negative	14	10	7	11	15	5
Polymicrobial	15	31	14	12	16	3
Other	3					

Tande and Patel. Clin Microbiol Rev 2014;27:302





PJI Definitions

	Definitions of Prosthetic Joint Infection					
	2011 Musculoskeletal Infection Society ¹		2013 Infectious Diseases Society of America ²		2013 International Consensus ³	
	Definitive evidence	Supportive evidence	Definitive evidence	Supportive evidence	Definitive evidence	Supportive evidence
Sinus tract communicating with the prosthesis	X		X		X	
Identical microorganism isolated from ≥2 cultures	X		X		X	
Purulence surrounding the prosthesis		X	X			
Acute inflammation of periprosthetic tissue		X		X		X
A single culture with any microorganism		X				X
A single culture with a virulent microorganism				X		
Elevated synovial fluid leukocyte count		X				X
Elevated synovial fluid neutrophil percentage		X				X
Elevated serum ESR and CRP		X				X

¹Parvizi et al. Clin Orthop Relat Res 2011;469:2992

²Parvizi & Gehrke. Proceedings of the International Consensus Meeting on Periprosthetic Joint Infection 2013

³Osmon et al. Clin Infect Dis 2013;56:e1

Musculoskeletal Infection Society (MSIS) and Infectious Diseases Society of America (IDSA) Diagnostic Criteria for Prosthetic Joint Infection



MSIS

Aseptic
Failure

Prosthetic
Joint Infection

Aseptic
Failure

IDSA Prosthetic
Joint
Infection

297 1

4 69

p=0.56

Musculoskeletal Infection Society (MSIS) and International Consensus Meeting (ICM) Diagnostic Criteria for Prosthetic Joint Infection



MSIS

Aseptic
Failure

Prosthetic
Joint Infection

Aseptic
Failure

298 0

ICM Prosthetic
Joint
Infection

3 70

p=0.65

Infectious Diseases Society of America (IDSA) and International Consensus Meeting (ICM) Diagnostic Criteria for Prosthetic Joint Infection



IDSA

Aseptic
Failure

Prosthetic
Joint Infection

Aseptic
Failure

294 4

ICM Prosthetic
Joint
Infection

4 69
p=0.9

Test Characteristics and Relative Costs of Preoperative Tests for PJI Diagnosis



Test	Joint	Threshold value	Sensitivity	Specificity	+ LR	- LR	Cost	Comments
PERIPHERAL BLOOD								
WBC	Hip and knee	11,000 x 10 ⁹ /L	45	87	3.5	0.6	\$	1796 patients in 15 studies
CRP		10 mg/L	88	74	3.4	0.2	\$	3225 patients in 23 studies
ESR		30 mm/hr	75	70	2.5	0.4	\$	3370 patients in 25 studies
IL-6		10 pg/mL	97	91	10.8	0.0	\$	432 patients in 3 studies
Procalcitonin		0.3 ng/mL	33	98	16.5	0.7	\$	78 patients in single study
IMAGING								
Plain radiograph	Hip	Lucency or periosteal new bone formation	75	28	1.0	0.9	\$	65 patients in single study
Triple phase bone scan	Late hip	Increased uptake on all 3 phases	88	90	8.8	0.1	\$\$\$	46 patients in single study
Bone scan/labeled leukocyte scan	Late hip and knee	Incongruent images	64	70	2.1	0.5	\$\$\$	166 patients in single study
FDG-PET scan	Hip and knee		82	87	6.1	0.2	\$\$\$\$\$	635 patients in 11 studies
SYNOVIAL FLUID								
Cell count	Knee	1100 cells/ μ L	91	88	7.6	0.1	\$\$	429 patients in single study
Neutrophil percentage		64%	95	95	17.9	0.1	\$\$	
Cell count	Hip	4200 cells/ μ L	84	93	12.0	0.2	\$\$	201 patients in single study
Neutrophil percentage		80%	84	82	4.7	0.2	\$\$	
Cell count	Knee (<6 weeks)	27,800 cells/ μ L	84	99	84.0	0.2	\$\$	146 patients in single study
Neutrophil percentage		89%	84	69	2.7	0.2	\$\$	
Culture	Hip and knee		72	95	14.4	0.3	\$\$	3332 patients in 34 studies

Multiplex Antibody Detection

- Developed multiplex antibody detection-based immunoassay by “comparative immunoproteomics” – compared antibody response profile of arthroplasty patients with and without PJI

	Sensitivity	Specificity
Staphylococci targeted	68/94 (72%)	213/264 (81%)
<i>S. aureus</i>	36/54 (67%)	
<i>S. epidermidis</i>	26/35 (74%)	
<i>S. lugdunensis</i>	9/9 (100%)	
<i>S. agalactiae</i>	6/8 (75%)	250/270 (93%)
<i>P. acnes</i>	5/13 (39%)	235/277 (85%)

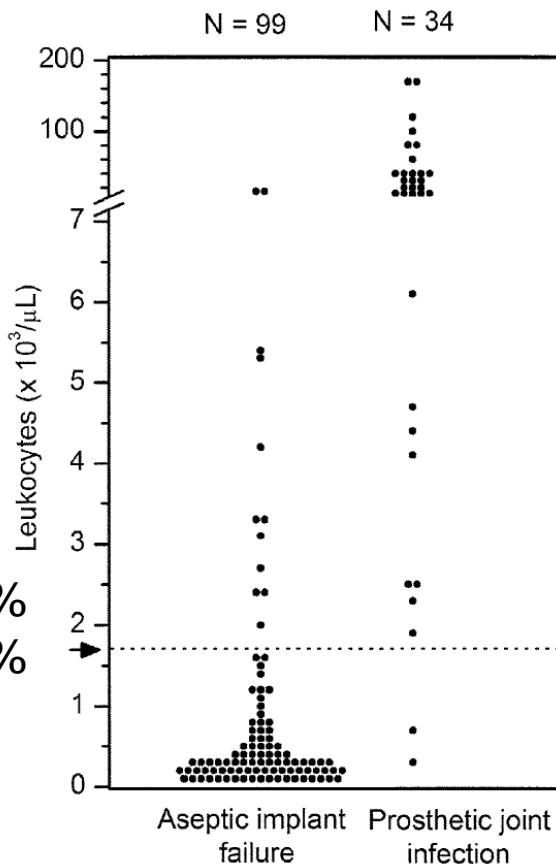
Synovial Fluid Should be Cultured in Blood Culture Bottles

- 219 subjects, revision hip or knee arthroplasty
- Intraoperative fluid inoculated into aerobic and anaerobic blood culture bottles (15 ml each)

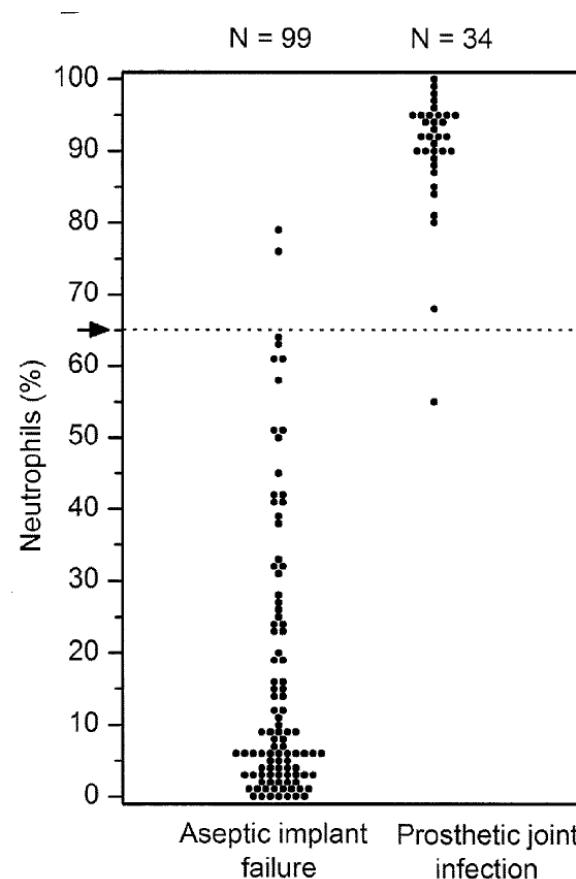
	Sensitivity	Specificity	Positive predictive value	Negative predictive value
Synovial fluid culture in blood culture vial	58	100	100	71
Synovial fluid culture standard	47	100	100	61



Synovial Fluid Leukocyte/Differential Prosthetic Knee (>6 months from index surgery)



Sensitivity, 94%
Specificity, 88%



Sensitivity, 97%
Specificity, 98%



Trampuz et al. Am J Med 2004;117:556

Synovial Fluid Leukocyte/Differential

	WBC (cells/ μ l)	Sensitivity	Specificity	Neutrophil percentage	Sensitivity	Specificity	Joint type	Time from surgery
Trampuz Am J Med 2004;117:556	1,700	94	88	65	97	98	Knee (133)	>6 months
Ghanem JBJS 2008;90:1637	1,100	91	88	64	95	95	Knee (429)	Varied
Schinsky JBJS 2008;90:1869	4,200	84	93	80	84	82	Hip (201)	Varied
Bedair CORR 2011;469:34	10,700	95	91	89	84	69	Knee (146)	<6 weeks
Cipriano JBJS 2012;94:594 Non-inflammatory arthritis	3,450	91	93	78	96	87	Hip/Knee (810)	Unknown
Cipriano JBJS 2012;94:594 Inflammatory arthritis	3,444	88	80	75	100	82	Hip/Knee (61)	Unknown
Zmistowski J Arthroplasty 2012;27:1589	3,000	93	94	75	93	83	Knee (150)	Unknown
Dinneen Bone & Joint J 2013;95:554	1,590	90	91	65	90	87	Hip/Knee (75)	Unknown

Natural Progression of Synovial Fluid Leukocytes in TKA

WBC, Neutrophil Percentage and Total Neutrophil Count by Time Period*

	First 45 Days	Days 46 to 90	3 Months to 1 Year	1 to 2 Years
WBC count/ μ L	3037 \pm 3786	1119 \pm 1325†	474 \pm 894‡	428 \pm 706
Neutrophil %	69 \pm 27	49 \pm 23†	41 \pm 27	28 \pm 32‡
Neutrophils μ L	2533 \pm 3483	649 \pm 871†	270 \pm 598‡	241 \pm 552

*Values are means and standard deviations

†Significantly different from the first 45 days ($p<0.05$)

‡Significantly different from days 46 to 90 ($p<0.05$)

Synovial Fluid CRP

CRP (mg/L)	Aseptic Failure	PJI	
Knee	N = 11	N = 10	Ronde-Oustau et al. Orthopaed & Traumatol: Surg and Res 2014;100:217
Serum	4.69 (0.94-66.50)	52.00 (19.50-225.00)	
Synovial Fluid	0.69 (0.17-30.80)	13.95 (4.39-92.10)	
Knee and Hip	N = 43	N = 20	Parvizi et al. J Arthroplasty 2012;27:12
Serum	1.1 (0-125)	98 (3-263)	
Synovial Fluid	2 (0-13)	40 (0-103)	
Knee			Parvizi et al. Clin Orthop Relat Res 2012;470:54
Serum	9.6 (N = 30)	91.7 (N = 25)	
Synovial Fluid (Individual ELISA)	0.002 (N = 5)	0.322 (N = 10)	
Synovial Fluid (Multiplex ELISA)	1.19 (N = 34)	22.49 (N = 25)	
Knee			Vanderstappen et al. Acta Orthop Belg 2013;79:326
Serum	3.38 (0.50-10.10) (N = 17)	32.70 (6.70-145.10) (N = 7)	
Synovial Fluid	1.08 (0.30-5.00) (N = 33)	11.98 (2.00-30.20) (N = 11)	
Hip	(N = 27)	(N = 38)	Buttarro et al. Clin Orthop Rel Res 2015;473:3876
Synovial Fluid	Cutoff 9.5 mg/L, sensitivity 90%, specificity 94%		

Novel Synovial Fluid Biomarkers

Biomarker	Cutoff	Sens-itivity (%)	Spec-ifity (%)	
α -Defensin	4.8 $\mu\text{g}/\text{mL}$	100	100	Deirmengian et al. Clin Orthop Rel Res 2014;472:3254 (95 subjects; 29 PJI; hip/knee)
Neutrophil elastase 2	2.0 $\mu\text{g}/\text{mL}$	100	100	
Bactericidal/permeability-increasing protein	2.2 $\mu\text{g}/\text{mL}$	100	100	
Neutrophil gelatinase-associated lipocalin	2.2 $\mu\text{g}/\text{mL}$	100	100	
Lactoferrin	7.5 $\mu\text{g}/\text{mL}$	100	100	
IL-8	6.5 ng/mL	100	95	
CRP	12.2 mg/L	90	97	
Resistin	340 ng/mL	97	100	
α -Defensin	5.2 mg/L	97	96	Deirmengian et al. JBJS 2014;96:1439 (149 subjects; 37 PJI; hip/knee; did not report WBC)
CRP	3 mg/L	97	79	
α -Defensin → if positive, CRP		97	100	
α -Defensin	5.2 mg/L	100	100	Deirmengian et al. CORR 2015;473:198 (46 subjects; 23 PJI; hip/knee; did not report WBC performance)
α -Defensin	7720 ng/mL	95	100	Bingham et al. CORR 2014;472:4006 (57 subjects; 19 PJI; hip/knee; did not report %neutrophils)
α -Defensin	0.48 S/CO	63	95	Frangiamore et al. J Shoulder Elbow Surg 2015;24:1021 (33 cases; 11 PJI; shoulder)
α -Defensin	1 S/CO	99		Shahi et al. CORR 2016;474:1610 (106 PJI subjects; hip/knee)
α -Defensin	1 S/CO	97	97	Bonanzinga et al. CORR 2017;475:408 (156 subjects; 33 PJI; hip/knee)
α -Defensin		77	82	Suda et al. Int Ortho 2017 In Press (28 subjects – 30 joints; hip/knee; 13 PJI)
IL-6	359.3 pg/mL	87	90	Frangiamore et al. J Bone Joint Surg Am 2015;97:63 (35 subjects; 15 PJI; shoulder)
IL-1 β IL-6	8.26 pg/ml 8.7 ng/ml	90 81	87 96	Frangiamore et al. Clin Orthop Relat Res 2016;474:1630 (90 subjects; 31 PJI; hip or knee)

Intraoperative Frozen Section Histopathology

Reference	Specimen	Joint	#PMN*	n	Sensitivity	Specificity	PPV**
Feldman et al. JBJS(Am) 1995;77:1807	JC, IM	Hip/knee	5	33	100	100	100
Athanasou et al. JBJS(Br) 1995;77:28	JC, IM	Hip/knee	1	106	90	96	88
Lonner et al. JBJS(Am) 1996;78:1553	JC, IM, ASPI	Hip/knee	5	175	84	96	70
			10		84	99	89
Pace et al. J Arthroplasty 1997;12:64	JC, IM	Hip/knee	5	18	82	93	82
Abdul-Karim et al. Mod Pathol 1998;11:427	IM, ST, UDT	Hip/knee	5	64	43	97	60
Banit et al. CORR 2002;401:230	JC, ASPI	Knee	10	55	100	96	82
		Hip	10	63	45	92	55
Musso et al. Postgrad Med J 2003;79:590	JC, IM, ASPI	Hip/knee	5	45	50	95	60
Wong et al. J Arthroplasty 2005;20:1015	JC, IM, SS	Hip/knee	5	33	93	77	68
			10		86	85	75
Ko et al. J Arthroplasty 2005;20:189	JC, IM, ASPI	Hip/knee	5	40	67	97	86
Frances Borrego et al. Int Orthop 2007;31:33	PST	Hip	10	63	67	90	80
		Knee		83	50	100	100
Nunez et al. Acta Orthop 2007;78:226	JC, IM, ASPI	Hip	5	136	86	87	79
Tohtz et al. CORR 2010;468:762	IM	Hip	10	52	87	100	100

*Some studies used >, others ≥ the number shown, **Positive predictive value

JC, joint pseudocapsule; IM, interface membrane; ASPI, any area that appears suspicious for possible infection; ST, synovial tissue; SS, synovial surface; UDT, unusually discolored tissue; PST, periprosthetic soft tissue

Periprosthetic Tissue and Device Culture

- **IDSA Guidelines** - At least 3 and optimally 5 or 6* periprosthetic intraoperative tissue samples or the explanted prosthesis itself should be submitted for aerobic and anaerobic culture at the time of surgical débridement or prosthesis removal to maximize the chance of obtaining a microbiologic diagnosis (B-II)

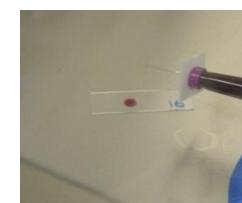
Osmon et al. Clin Infect Dis 2013;56:1

*Atkins et al. J Clin Microbiol 1998;36:2932

- **ESCMID Guidelines** - Peri-implant tissue (AIII) and device/prosthesis sonication (BII)

Høiby et al. Clin Microbiol Infect 2015;21:S21

Periprosthetic Tissue Culture Blood Culture Bottles



Periprosthetic Tissue Culture Blood Culture Bottle Study (Oxford #1)

- Tissue + sterile glass beads (Ballotini) 5 ml saline (shaking)
 - 1 ml each into Robertson's cooked meat broth, fastidious anaerobic broth, BACTEC Anaerobic/F & Aerobic/F bottles
 - 0.25 ml each onto chocolate, aerobic & anaerobic (2) blood agars
 - 5 day incubation
 - 141 elective joint revisions (mean, 4.9 specimens/case)
 - 23 PJI cases

Sensitivity/ Specificity (≥2 positive)	Direct plates	Fastidious anaerobic broth	Cooked meat broth	BACTEC blood culture bottles
Sensitivity, %	39	57	83	87
Specificity, %	100	100	97	98

Periprosthetic Tissue Culture Blood Culture Bottle Study (Oxford #2)

- Tissue + 3 ml saline + sterile glass beads → vortexed 15 sec
 - 0.5 ml BACTEC Lytic/10 Anaerobic/F & Plus Aerobic/F bottles
 - 14 day incubation
- 322 patients (mean, 4 specimens/case)
- 66/79 PJI culture-positive (sensitivity, 84%)
 - Positive within 3 days of culture in all cases except 1 (detected at 8 days)
- 7/243 non-PJI culture-positive (specificity, 97%)
- *Propionibacterium* species
 - Sub-culturing negative bottles after 14-days' incubation detected a single additional *Propionibacterium* isolate from 1000 bottles



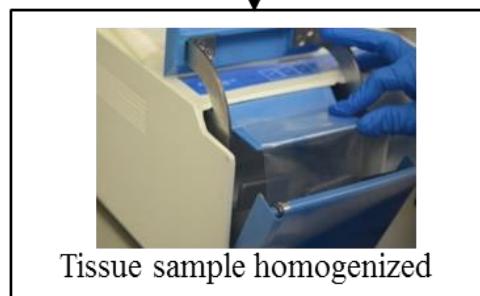
Periprosthetic Tissue Blood Culture Bottle Study (Mayo Clinic)



- Patients undergoing revision arthroplasty, Mayo Clinic, 8/2013–4/2014
 - 369 subjects - 138 hip, 160 knee, 49 shoulder, 22 elbow arthroplasties
 - 117 (32%) met IDSA diagnostic criteria for PJI



Peri-prosthetic tissue sample



Tissue sample homogenized

Conventional Culture Technique:

Homogenate inoculated onto agars and broth



New Technique:

Homogenate inoculated into blood culture bottles and placed onto the semi-automated machine



Media manually checked daily for growth and sub-cultured if growth detected

Bottles continuously monitored for growth and sub-cultured if flagged positive



Periprosthetic Tissue Blood Culture Bottle Study (Mayo Clinic) Bayesian Latent Class Modeling

Media Combination	Sensitivity	Specificity
Aerobic and anaerobic agars	48.9	99.7
Aerobic and anaerobic agars and thioglycollate broth	62.6	98.1
Aerobic and anaerobic blood culture bottles	92.1	99.7
Aerobic and anaerobic blood culture bottles and thioglycollate broth	92.1	98.8
Aerobic and anaerobic blood culture bottles and aerobic agar	94.6	99.7
Aerobic and anaerobic blood culture bottles and anaerobic agar	96.8	99.8
Aerobic and anaerobic blood culture bottles and aerobic and anaerobic agars	99.1	99.7
All media combined	99.1	97.3

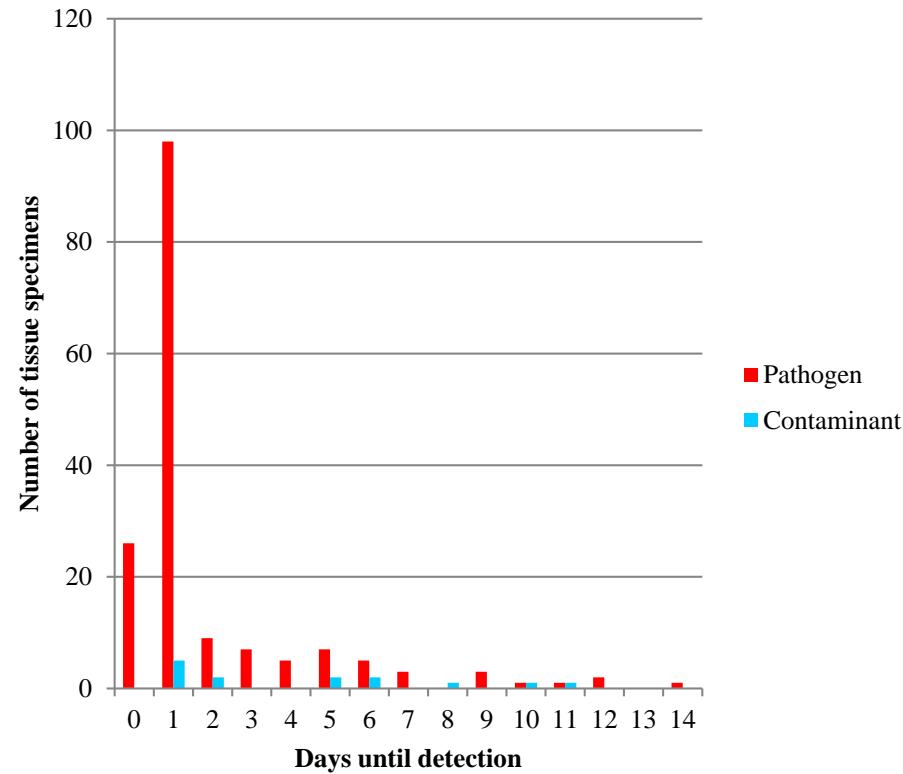
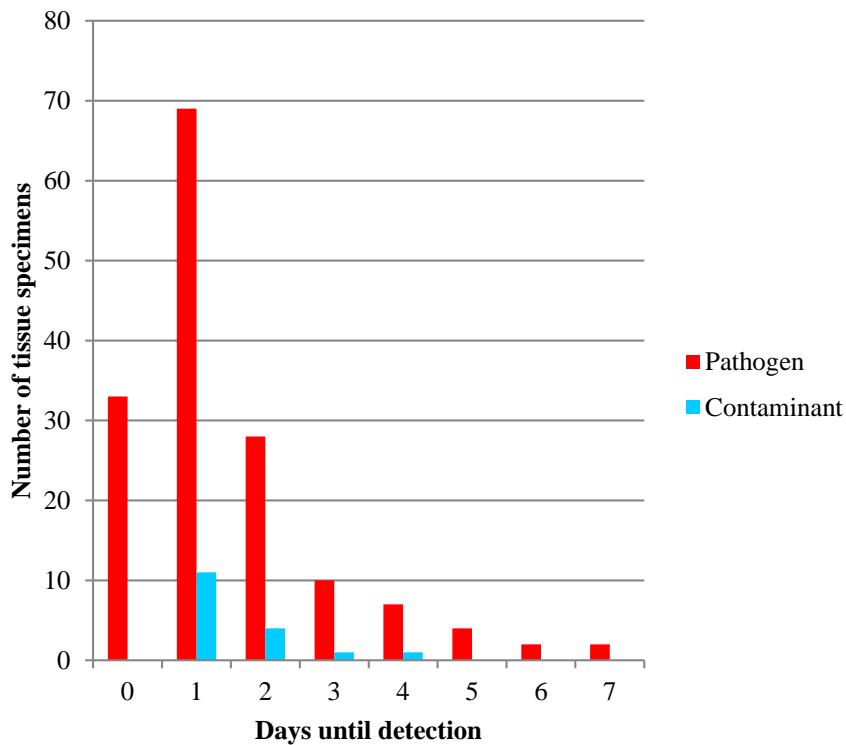


Sensitivity and Specificity using PJI (IDSA Criteria) as the Gold Standard

Media Combination	Sensitivity	Specificity
Aerobic and anaerobic agars	33.3	100.0
Aerobic and anaerobic agars and thioglycolate broth	44.4	98.8
Aerobic and anaerobic blood culture bottles	60.7	98.8
Aerobic and anaerobic blood culture bottles and thioglycolate broth	63.3	98.8
Aerobic and anaerobic blood culture bottles and aerobic agar	62.4	98.8
Aerobic and anaerobic blood culture bottles and anaerobic agar	62.3	98.0
Aerobic and anaerobic blood culture bottles and aerobic and anaerobic agars	64.1	98.0
All media combined	67.5	96.8



Periprosthetic Tissue Blood Culture Bottle Study (Mayo Clinic) Time-to-Detection

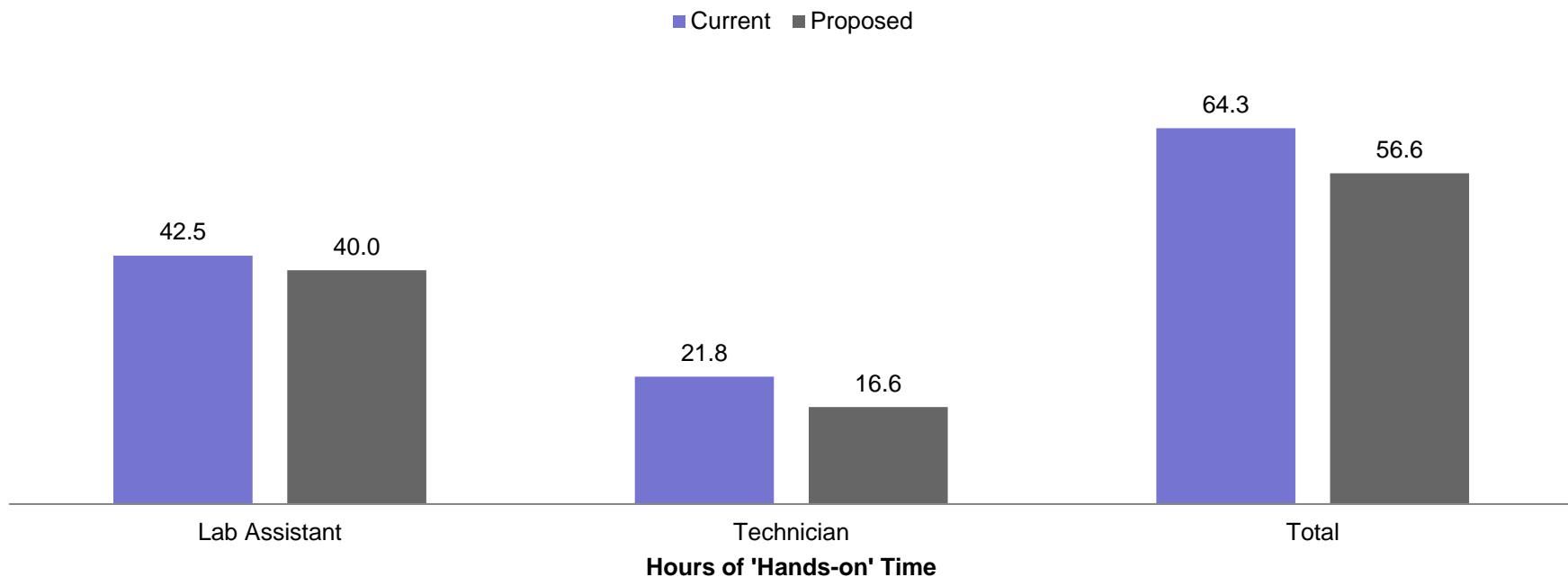


Aerobic bottles

Anaerobic bottles

Laboratory Workflow Analysis of Culture of Periprosthetic Tissues in Blood Culture Bottles

- 10-11% reduction in total staff time
- 20-24% reduction in technician time



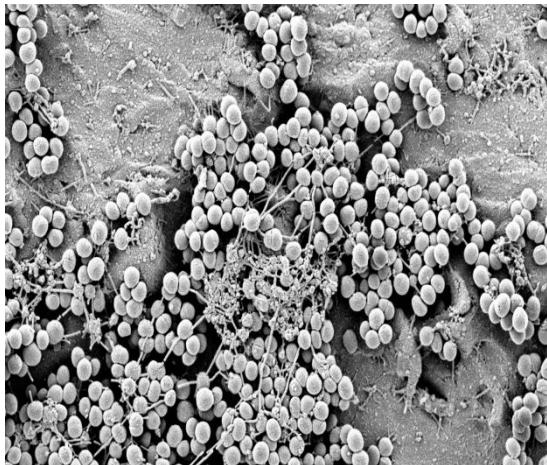
Prospective Multicenter Trial Assessing Sample Number

- 215 PJI subjects
 - 5 samples/patient - assessed using random selection 2, 3, 4 or 5 samples
 - Results with individual culture media compared
 - Aerobic bacteria: Blood agar, 70%; chocolate agar, 69%; Schaedler broth, 69%; pediatric blood culture bottle, 83%
 - Anaerobic bacteria: Blood agar, 31%; chocolate agar, 53%; Schaedler broth, 75%; pediatric blood culture bottle, 54%
 - Agreement of 98 and 100%, respectively, for bacteriological criterion and confirmed PJI diagnosis obtained with 4 perioperative samples
 - Highest percentages of agreement obtained with association of 3 culture media, blood culture bottle, chocolate agar and Schaedler broth, incubated for 5, 7 and 14 days, respectively
- **4 samples seeded in/on 3 culture media sufficient for diagnosing PJI**

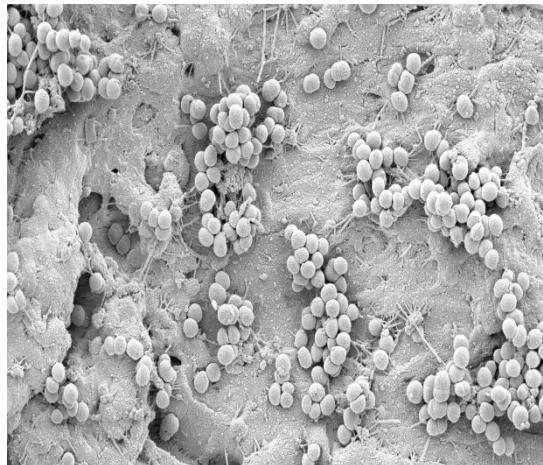
Periprosthetic Tissue Culture

- 499 consecutive patients undergoing revision arthroplasty (August 2013 through April 2014)
- 1,437 periprosthetic tissue samples
- Conventional culture techniques (aerobic and anaerobic sheep blood agars, thioglycolate broth)
 - Greatest accuracy with four specimens (91%; 95% Credible Interval: 77-100%)
- Inoculation of tissues into aerobic and anaerobic blood culture bottles
 - Greatest accuracy with three specimens (92%; 95% Credible Intervals: 79-100%)

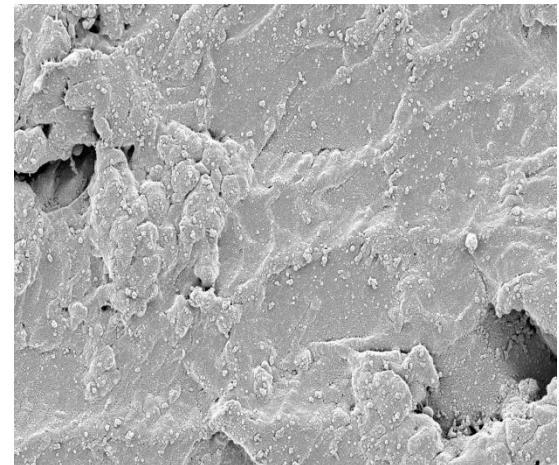
Staphylococcus epidermidis Biofilm on Polycarbonate Coupons Scanning Electron Microscopy



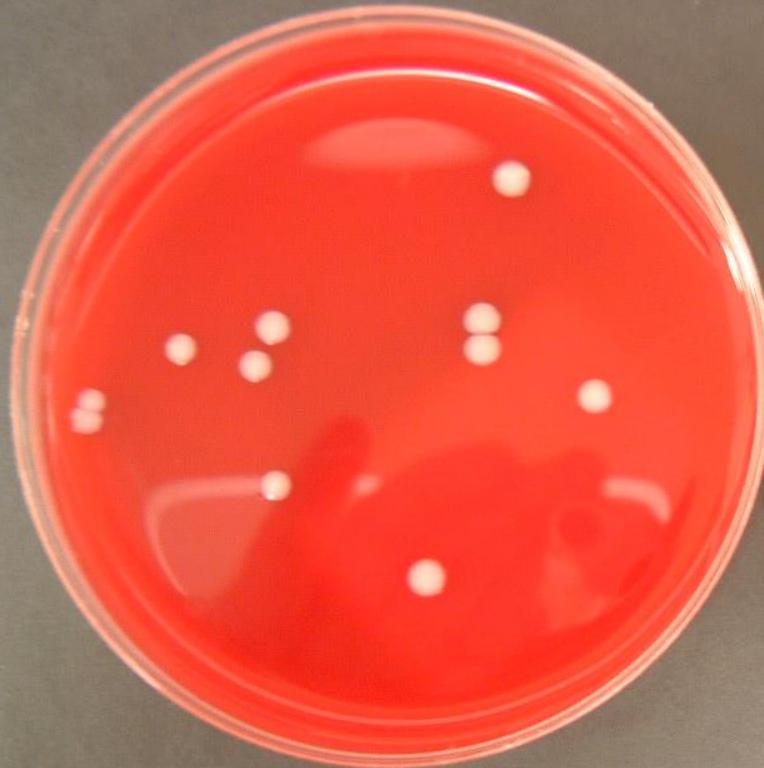
Soaking



Scraping



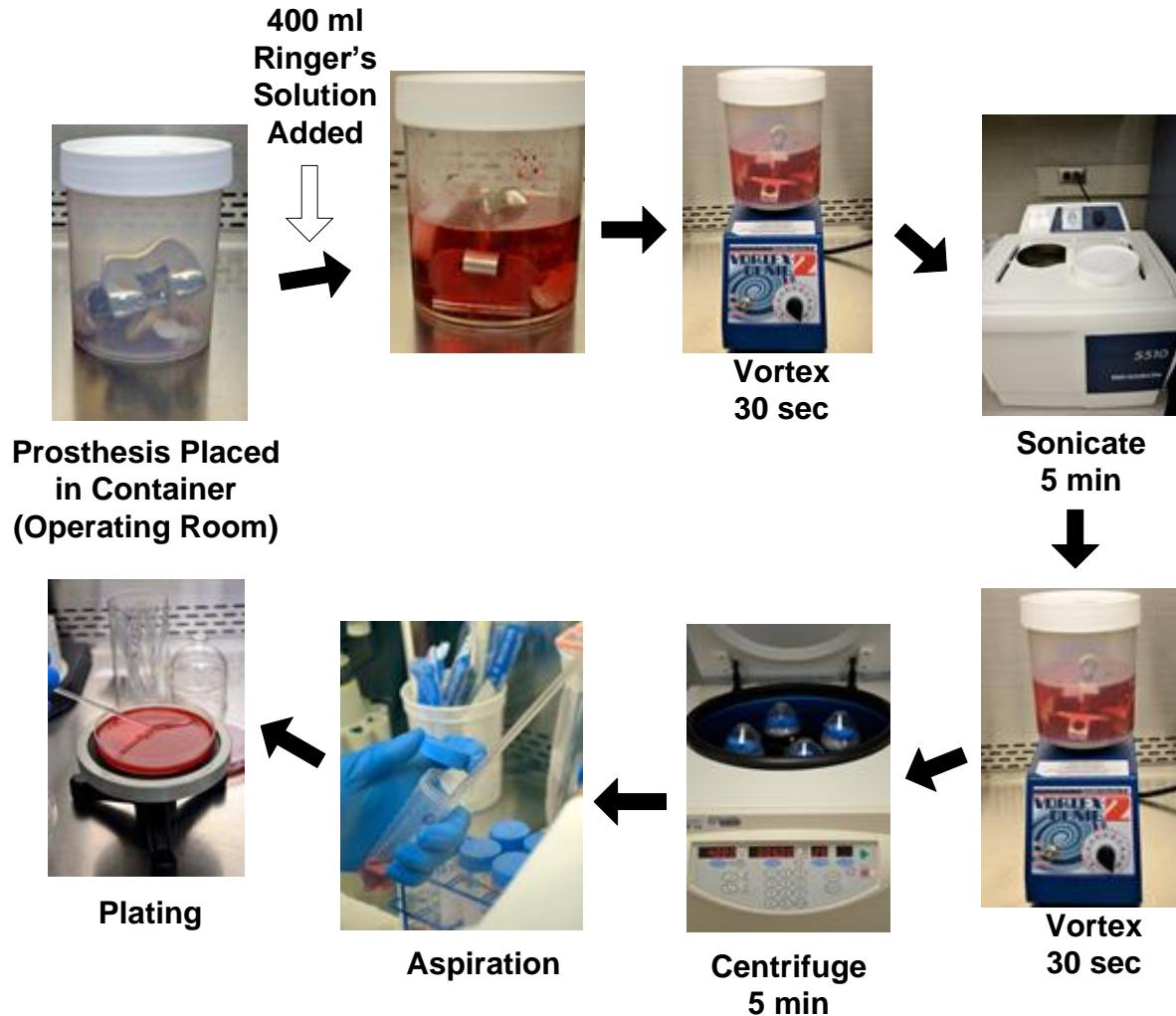
Sonication



Tissue

Sonicate

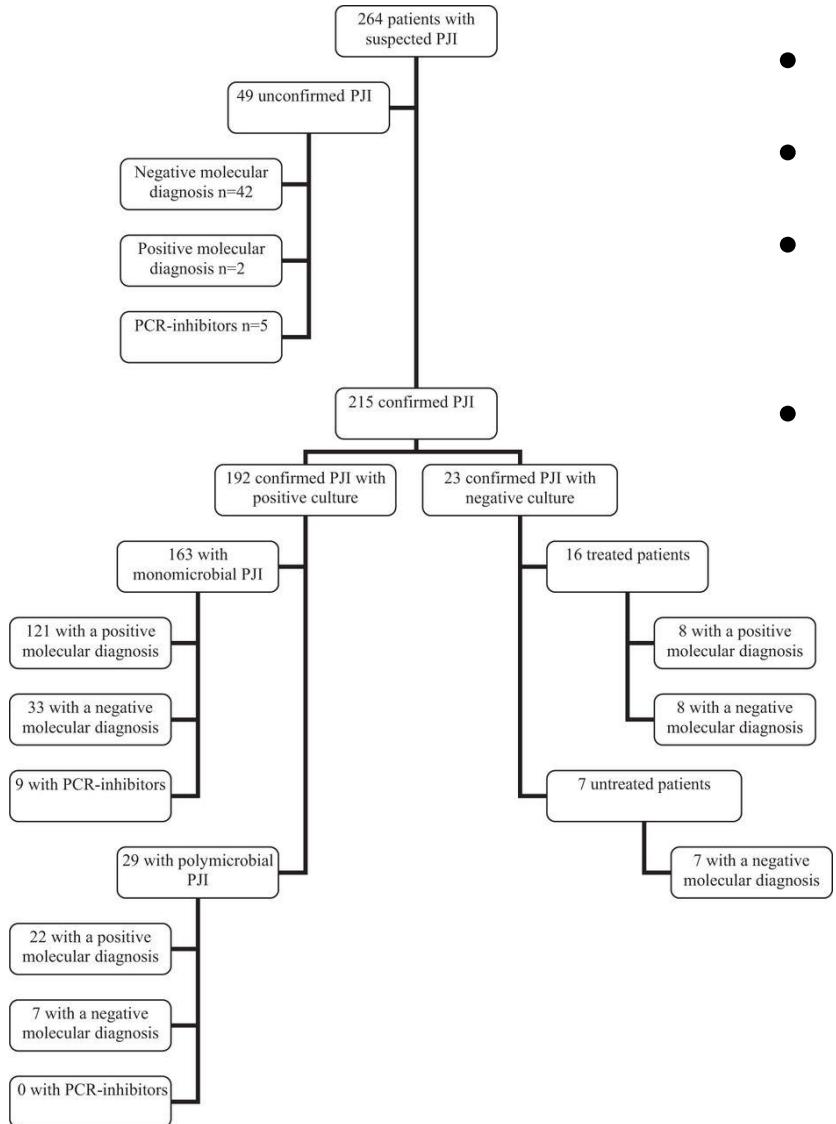
Current Orthopedic Implant Processing - Mayo Clinic



Comparison of Sonicate Fluid and Tissue Culture Diagnosis of Orthopedic Foreign Body Infection

Implant type		Sonicate fluid	Periimplant tissue	p value	Reference
Prosthetic hip/knee joint	Sensitivity	79%	61%	<0.001	Trampuz et al, NEJM 2007 Vol 357;654
	Specificity	99%	99%		
Prosthetic shoulder joint	Sensitivity	67%	55%	0.046	Piper et al, JCM 2009 Vol 47:1878
	Specificity	98%	95%		
Spine implant	Sensitivity	91%	73%	0.046	Sampedro et al, Spine 2010 Vol 25:1218
	Specificity	97%	93%		
Prosthetic elbow joint	Sensitivity	89%	55%	0.18	Vergidis et al, JSES 2011 Vol 20;1275
	Specificity	100%	93%		
Prosthetic hip/knee/shoulder joint, mega-prostheses, osteosynthesis, spine implant	Sensitivity	83%	61%	0.034	Holinka et al. J Orthop Res 2011 Vol 29:617
	Specificity	95%	95%		
Prosthetic hip joint	Sensitivity	75	69		Bogut et al. Polish J Microbiol 2014;63:299
	Specificity	97	100		
Prosthetic hip/knee joint, internal device	Sensitivity	77	34	<0.002	Scorzolini et al. New Microbiol 2014;37:321
Fracture fixation implant (plate and screws, spine implant, intramedullary nail)	Sensitivity	90	57	<0.001	Yano et al. J Clin Microbiol 2014;52:4176
	Specificity	91	96		

Lack of Sensitivity of Periprosthetic Tissue Broad-Range Bacterial PCR



- 2-year period
- 5 tissues/patient
- Culture, 16S rRNA gene PCR/sequencing
- 264 suspected cases of PJI
 - PJI confirmed (215)
 - Culture positive, 192 (89%)
 - PCR positive, 151 (73%)
 - Non-PJI (49)
 - PCR positive, 2 (specificity, 96%)

Sonication and Broad-Range PCR Hip/Knee Prostheses



- Sonication fluid more sensitive than periprosthetic tissue

Rak et al. Acta Orthopaedica 2016;87:339

Test	Sensitivity 135 PJI	Specificity 231 Aseptic Failure	Accuracy
% (95% Confidence Interval)			
Tissue culture	70.4 (64.5-76.3)	98.7 (97.2-100)	88.3 (84.2-92.4)
Sonicate fluid culture	72.6 (66.8-78.4)	98.3 (96.6-100)	88.8 (84.7-92.9)
Sonicate fluid broad-range PCR	70.4 (64.5-76.3)	97.8 (95.9-99.7)	87.7 (83.5-91.9)
Combination of two tests above	83.0 (78.2-87.8)	95.7 (93.1-98.3)	91.0 (87.3-94.7)
Sonicate fluid culture plus PCR	78.5 (73.2-83.8)	97.0 (94.8-99.2)	90.2 (86.4-94.0)
Synovial fluid culture	64.7 (56.5-72.9)	96.9 (93.9-99.9)	84.1 (77.8-90.4)
Sonicate fluid PCR - lower cutoff (CP <27.59 cycles)	80.0 (74.8-85.2)	90.9 (87.2-94.6)	86.8 (82.5-91.3)



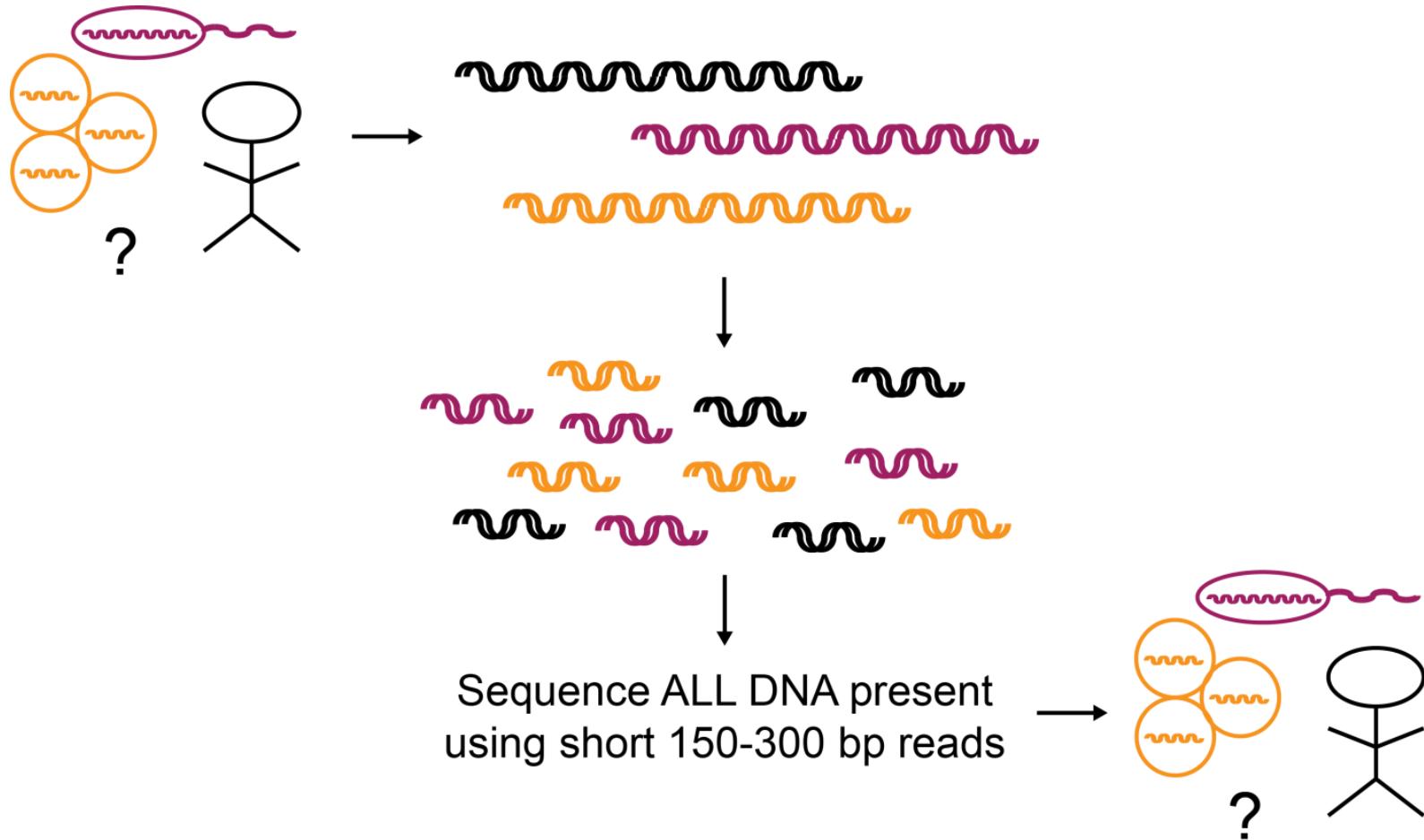
Sonication Studies – PJI PCR Panel Hip/Knee Prostheses

Test	Aseptic failure (290)	PJI (144)	Sensitivity	Specificity	PPV	NPV
	No. of patients with positive specimens		% (95% confidence interval)			
Synovial-fluid culture	5/161	59/89	66.3 (55.5-76.0)	96.9 (92.9-99.0)	92.2 (82.7-97.4)	83.9 (77.8-88.8)
Tissue culture						
Any growth	45	119	82.6 (75.4-88.4)	84.5 (79.8-88.5)	72.6 (65.1-79.2)	90.7 (86.6-93.9)
≥2 positive tissues (same organism)	6	101	70.1 (62.0-77.5)	97.9 (95.6-99.2)	94.4 (88.2-97.9)	86.9 (82.7-90.3)
Sonicate fluid culture	5	105	72.9 (64.9-80.0)	98.3 (96.0-99.4)	95.5 (89.7-98.5)	88.0 (83.9-91.3)
Sonicate fluid PCR (10 assay panel)			77.1 (69.3-83.7)	97.9 (95.6-99.2)	94.9 (89.2-98.1)	89.6 (85.7-92.7)
Any positive result	6	111				
<i>Staphylococcus</i> sp	2	75				
<i>S. aureus</i>	0	28				
Coagulase-negative staphylococci	2	47				
<i>Streptococcus</i> sp	3	11				
<i>Enterococcus/Granulicatella/Abiotrophia</i> sp	0	11				
Enterobacteriaceae	1	8				
Gram-positive anaerobic cocci	0	8				
<i>Propionibacterium</i> sp	0	8				
<i>P. aeruginosa</i>	0	5				
<i>Corynebacterium</i> sp	0	4				
<i>C. jeikeium/urealyticum</i>	0	0				
Non- <i>jeikeium</i> sp	0	4				
<i>Proteus</i> sp	0	1				
<i>B. fragilis</i> group	0	0				

Commercial Multiplex PCR Assays

- Unyero i60 implant and tissue infection (ITI) multiplex PCR (Curetis)
 - 28 subjects; 30 hip/knee arthroplasties, 13 PJI
 - **Tissue** “synovia” (2 mm punch press)
 - Sensitivity, 31%; specificity, 100%
Suda et al. Int Ortho 2017 In Press
 - 28 subjects, 79 samples
 - **Tissue** (3 mm)³
 - 7 culture positive – 3 concordant positive
 - 21 culture negative – 20 concordant negative
Borde et al. Infection 2015;43:551
 - 31 subjects
 - **Sonication fluid and synovial fluid**
 - Sensitivity, 67%; specificity 100%
Hischebeth et al. J Microbiol Meth 2016;121:27

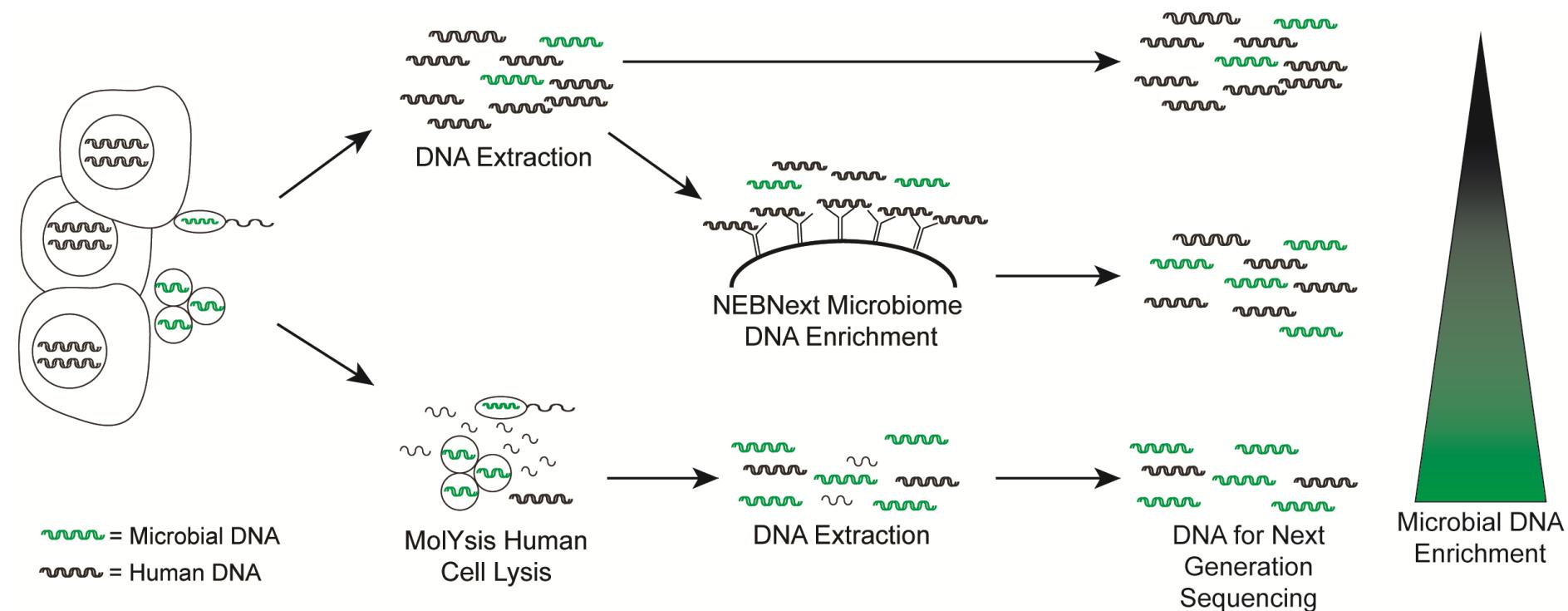
Metagenomic Shotgun Whole Genome Sequencing



Methods

- Paired-end libraries prep: **NEBNext Ultra DNA Library Prep Kit**
- Sequencing: **Illumina HiSeq 2500** in rapid run mode with 2x250 bp reads, as single samples or multiplexed with 2 or 6 samples per lane, based on number of samples available
- Paired-end reads pre-processing: **seqtk**, **Trimmomatic** & **Livermore Metagenomics Analysis Toolkit** with pre-processing scripts
- Pre-filter human and PhiX reads: **BioBloom tools**
- Taxonomy assignment: **LMAT with kML + Human.v4-14.20.g10.db database**

Comparison of Microbial DNA Enrichment Tools for Metagenomic Whole Genome Sequencing



Effect of Enrichment Methods by Metagenomic Whole Genome Sequencing

	No Enrichment	NEBNext Microbiome DNA Enrichment	MolYsis Enrichment
<i>S. aureus</i> PJI			
% of reads	0.02% (4,158 of 25,609,460)	0.2% (350,625 of 169,981,133)	7.7% (2,286,890 of 29,530,730)
Enrichment factor		13X	481X
<i>S. epidermidis</i> PJI			
% of reads	0.007% (1,682 of 23,606,476)	0.2% (133,680 of 74,544,475)	7.0% (2,268,087 of 32,184,381)
Enrichment factor		25X	986X
<i>E. faecalis</i> PJI			
% of reads	0.006% (1,671 of 26,949,030)	0.5% (497,206 of 94,522,959)	59.4% (16,407,878 of 27,643,294)
Enrichment factor		85X	9,580X

S. aureus PJI

No Enrichment

Read Score	# of Reads	TaxID Taxonomy group
877206	445817	9606 species,Homo sapiens genus,Streptococ
54347.8	29068	1301 <i>cus</i>
25834.1	32491	2759 superkingdom,Eukaryota
9473.81	15978	131567 no rank,cellular organisms species group,Streptococcus
7892.17	4622	671232 <i>anginosus</i> group
4756.18	2901	1279 genus,Staphylococcus species,Staphylococcus
1567.67	1007	1280 <i>aureus</i>
1197.96	755	1338 species,Streptococcus intermedius genus,Pseudomo
843.518	509	286 <i>nas</i>
534.066	346	76860 species,Streptococcus constellatus no rank,Streptococcus <i>tigurinus</i>
367.308	232	1282664 AZ_3a
361.242	530	1 depth=0,taxid=1,ktaxid=1,entries=-1
347.345	228	1282 species,Staphylococcus epidermidis subspecies,Streptococcus
255.733	159	184250 <i>constellatus</i> subsp. <i>pharyngis</i>
252.168	204	1328 species,Streptococcus <i>anginosus</i> no rank,Streptococcus intermedius
223.389	155	857292 F0395 no rank,Streptococcus intermedius
221.627	170	883167 F0413
206.954	130	1248438 no rank,Pseudomonas alcaliphila 34
175.499	102	1743 genus,Propionibacterium
145.828	96	90964 family,Staphylococcaceae no rank,Streptococcus equi subsp.
131.955	141	1051072 <i>zooepidemicus</i> ATCC 35246 species group,Pseudomonas
106.976	75	136841 <i>aeruginosa</i> group no rank,Pseudomonas
79.7075	55	1182590 <i>pseudoalcaligenes</i> CECT 5344 no rank,Streptococcus <i>salivarius</i>
71.3676	49	1074494 M18
71.2256	52	1300 family,Streptococcaceae
70.0294	52	32630 species,synthetic construct species,Streptococcus
69.8811	44	1077464 <i>tigurinus</i>
51.1347	31	101564 species,Pseudomonas alcaliphila no rank,Pseudomonas <i>mendocina</i>
48.2248	35	1390370 EGD-AQ5 phylum,Firmicute
47.8926	86	12396

NEBNext Enrichment

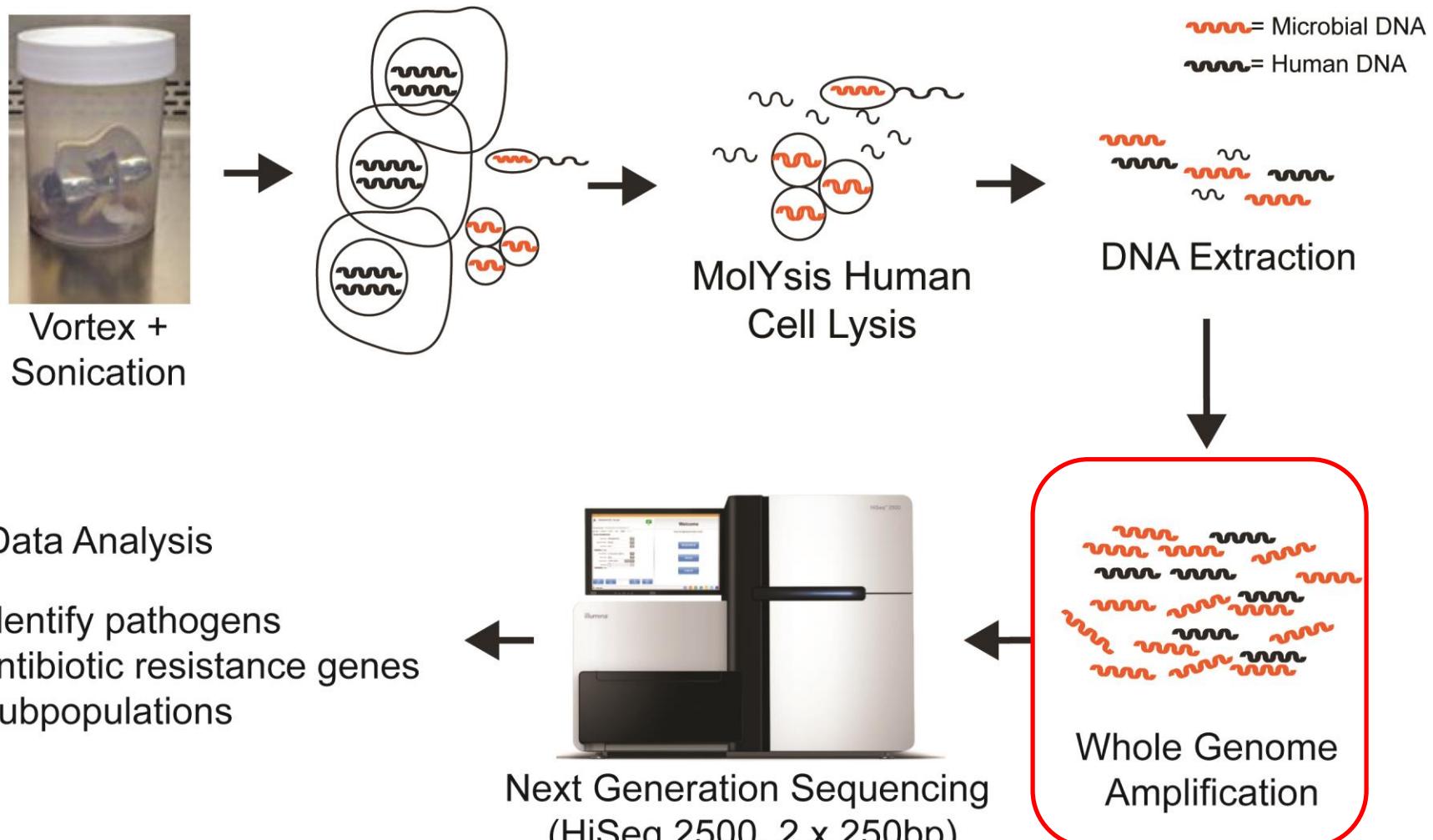
Read Score	# of Reads	TaxID Taxonomy group
8.14E+07	4168479	9606 species,Homo sapiens
	4	genus,Streptococ
807585	1210368	2759superkingdom,Eukaryota
611285	342132	32630species,synthetic construct
580565	336338	1280species,Staphylococcus aureus
268536	481220	131567 no rank,cellular organisms
15809	10833	1279genus,Staphylococcus
13898.3	18574	1depth=0,taxid=1,ktaxid=1,entries=-1
13530.1	9200	286genus,Pseudomonas
2688.34	2762	136843species group,Pseudomonas fluorescens
1816.09	1388	groupno rank,Staphylococcus aureus HIF003_B2N-C
1561.25	959	90964family,Staphylococcaceae
1318.53	776	1743genus,Propionibacterium
1236.36	1069	10003441species,Staphylococcus aureus plasmid PTZ2162
1226.75	1799	294species,Pseudomonas fluorescens
1029.9	1597	1292037no rank,Amycolatopsis vancoresmycin DSM 44592
923.909	719	10003448 species,Staphylococcus aureus plasmid pSA1379
645.914	403	463794no rank,Pseudomonas fluorescens BBc6R8
564.99	561	613genus,Serratia
462.127	1310	2superkingdom,Bacteria
408.322	283	1197727species,Pseudomonas sp. Ag1
256.031	204	1284392species,Pseudomonas sp. FH1
230.898	254	543family,Enterobacteriaceae
217.701	156	10003454 species,Staphylococcus aureus plasmid EDINA
203.746	204	317species,Pseudomonas syringae
196.998	215	1239phylum,Firmicutes
181.019	146	68887species,Torque teno virus
178.088	112	1206777species,Pseudomonas sp. Lz4W
154.956	184	1163065species,Pseudomonas sp. CBZ-4
144.54	125	135621family,Pseudomonadaceae
112.687	87	99158species,Hammondia hammondi
109.857	84	1240676species,Pseudomonas sp. PAMC 26793
94.634	63	1301genus,Streptococcus
93.5661	55	1248438no rank,Pseudomonas alcaliphila 34
91.8243	74	80865genus,Delftia
86.2212	81	308865genus,Elizabethkingia
77.866	45	985010species,Pseudomonas sp. TJI-51
74.2086	46	53335genus,Pantoea
69.9358	45	91459species,Methylobacterium sp. B1

MolYsis Enrichment

Read Score	# of Reads	TaxID Taxonomy group
	219358	1280species,Staphylococcus aureus
3.21E+06	7	9606 species,Homo sapiens
1.26E+06	713346	2759superkingdom,Eukaryota
193781	215573	1279genus,Staphylococcus
96085.8	71730	no rank,Staphylococcus
86648.5	93461	131567 no rank,cellular organisms
		no rank,Staphylococcus aureus
17892.4	15986	1156998 HIF003_B2N-C
4970.16	3686	90964family,Staphylococcaceae
		species,Staphylococcus aureus plasmid
3083.97	2641	10003441 pTZ2162
		species,Staphylococcus aureus plasmid
2110.41	1776	10003448 pSA1379
		1depth=0,taxid=1,ktaxid=1,entries=-1
1768.6	2414	32630species,synthetic construct
1649.2	1185	1239phylum,Firmicutes
958.691	1031	species,Staphylococcus aureus plasmid
867.117	669	10003454 EDINA
716.226	2113	2superkingdom,Bacteria
		species,Staphylococcus aureus plasmid
372.719	290	10003425 pBORa53
307.861	198	286genus,Pseudomonas
200.233	175	68887species,Torque teno virus
192.236	504	1385order,Bacillales
99.5046	91	10003460species,Staphylococcus aureus plasmid
77.8404	173	91061 class,Bacilli
50.2773	45	308865genus,Elizabethkingia
47.0102	71	1282species,Staphylococcus epidermidis
44.0471	40	1301genus,Streptococcus
39.0756	22	1743genus,Propionibacterium
37.8316	24	1248438no rank,Pseudomonas alcaliphila 34
		species group,Pseudomonas aeruginosa
28.2731	18	136841 group
		no rank,Pseudomonas mendocina EGD-
24.6249	18	1390370 AQ5

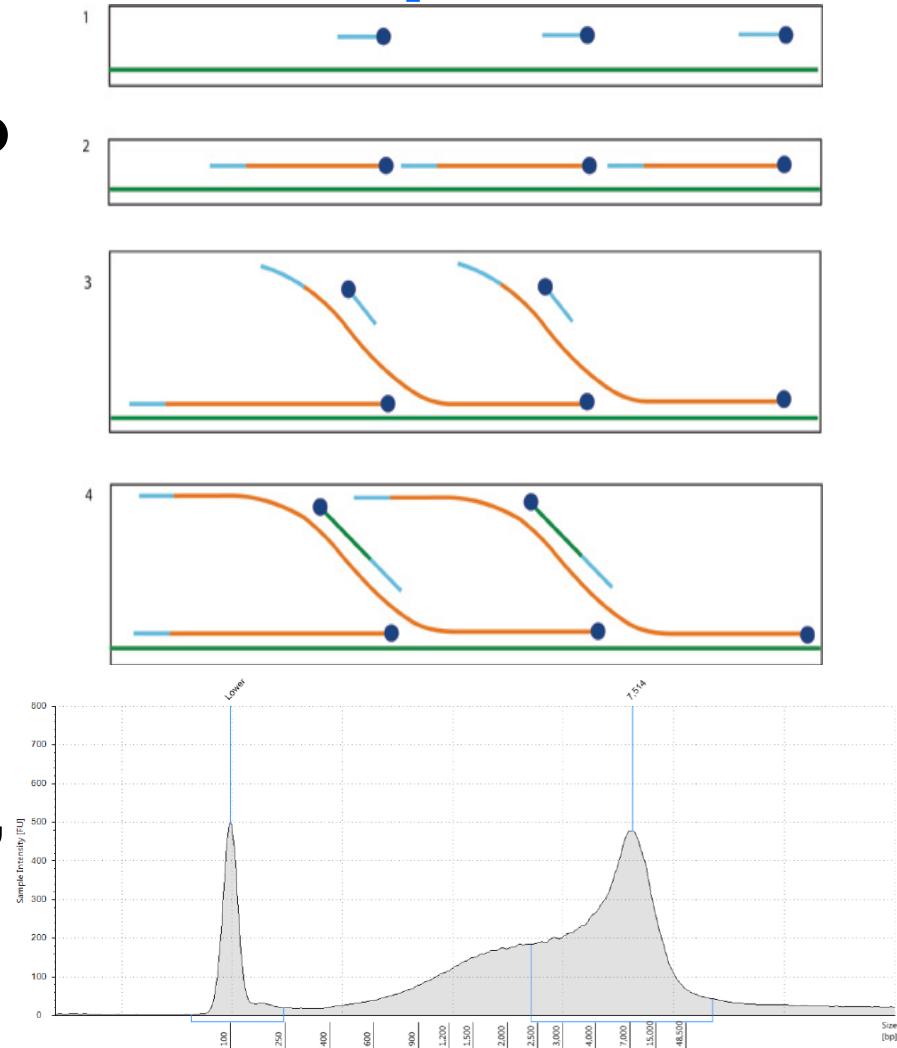
Thoendel et al. J Microbiol Methods
2016;127:141–5

Host DNA Depletion Depletes DNA Quantities for Sequencing



Whole Genome Amplification Using Multiple Displacement Amplification

- Random hexamers bind to template DNA
- Phi29 DNA polymerase amplifies DNA, displacing previous dsDNA
- High-fidelity, isothermal reaction
- Not completely unbiased
 - High GC content DNA, low abundance templates, small DNA fragments



Impact of Contaminating DNA in Whole Genome Amplification Kits on PJI Diagnosis

- 8 sonicate fluids, including culture-positive and –negative PJI, and aseptic failure
- DNA extraction: MoBio Bacteremia DNA isolation kit
- Whole genome amplification:
 1. Qiagen REPLI-g Single Cell kit (Qiagen, Hilden Germany)
 2. Illustra V2 Genomiphi kit (GE Healthcare Bio-Sciences, Pittsburgh PA)
 3. Illustra Single Cell Genomiphi kit
- Amplified DNA purification: Agencourt Ampure XP beads
- Paired-end library preparation: NEBNext Ultra DNA Library Prep Kit
- Sequencing: Illumina HiSeq 2500 rapid run mode, 2x250 bp reads (multiplexed 6 samples/lane)
- Controls (no WGA) - library preparation with NEBnext Ultra II DNA library Prep Kit (4 samples/lane)
- Illumina adapters removal: Trimmomatic
- Human and PhiX sequences prefiltration: BioBloom tools
- Taxonomic assignment of reads: Livermore Metagenomics Analysis Toolkit (LMAT)
- Trimming of low-quality reads: Trimmomatic
- Analysis of trimmed reads: MetaPhlan2

Sonicate Fluid Samples Tested and Relative Read Counts (LMAT)

Sample	Site	Microbiology		Qiagen REPLI-g		Illustra Single Cell		Illustra V2	
		Sonicate Fluid	Positive tissue cultures	Total reads		Total reads		Total reads	
				Pathogen reads	Non-pathogen reads	Pathogen reads	Non-pathogen reads	Pathogen reads	Non-pathogen reads
Culture-positive PJI									
980	H	Group C Streptococcus <20 CFUs	1 of 3	27,244,012		36,576,986		25,754,214	
				2,694,906	103	286,577	556	3,075,181	566
982	K	<i>S. epidermidis</i> >100 CFUs	3 of 4	28,320,234		30,284,116		26,508,366	
				3,272,895	591	102,934	203	9,539,147	17,625
986	K	<i>S. aureus</i> <20 CFUs	1 of 5	26,835,306		28,991,289		27,391,148	
				26,397	77,581	424	308	168,046	240,703
996	H	<i>B. fragilis</i> , >100 CFUs	3 of 3	28,388,841		28,745,844		28,904,638	
				155,761	4,853	11,134	1,346	96,154	8,140
1002	K	<i>C. striatum</i> , 51- 100 CFUs	3 of 3	32,165,206		27,925,551		32,338,186	
				536,155	2,594	5,503	422	1,000,810	53,019
Culture-negative PJI									
984	K	Culture negative	0 of 4	31,616,449		26,240,307		36,808,230	
				NA	2,821	NA	305	NA	16,974

Sonicate Fluid Samples Tested and Relative Read Counts (LMAT)

Sample	Site	Microbiology		Qiagen REPLI-g		Illustra Single Cell		Illustra V2	
		Sonicate Fluid	Positive tissue cultures	Total reads	Pathogen reads	Non-pathogen reads	Pathogen reads	Non-pathogen reads	Pathogen reads
Aseptic Failure									
983	K	Culture negative	0 of 3	32,924,419		29,076,492		32,128,178	
				NA	278	NA	522		45,539
987	K	Anaerobic organism, <20 CFUs	1 of 3, <i>Bacillus</i> species	30,316,155		30,587,175		27,955,363	
				NA	7,628	NA	265	NA	655,417
Controls									
<i>C. glutamicum</i>	NA	NA	NA	28,121,822		28,973,351		26, 986,400	
				26,912,706	2,145	22,276,446	856	26,149,302	1,311
Ringers	NA	NA	NA	27,488,254		29,531,289		27,810,322	
				NA	24,140	NA	3,560	NA	835,014
WGA no template	NA	NA	NA	25,669,747		53,626,034		24,512,662	
				NA	6,695	NA	46,759	NA	7,818,383

Taxonomic Identification of Reads (Performed using LMAT, Grouped by Genus)

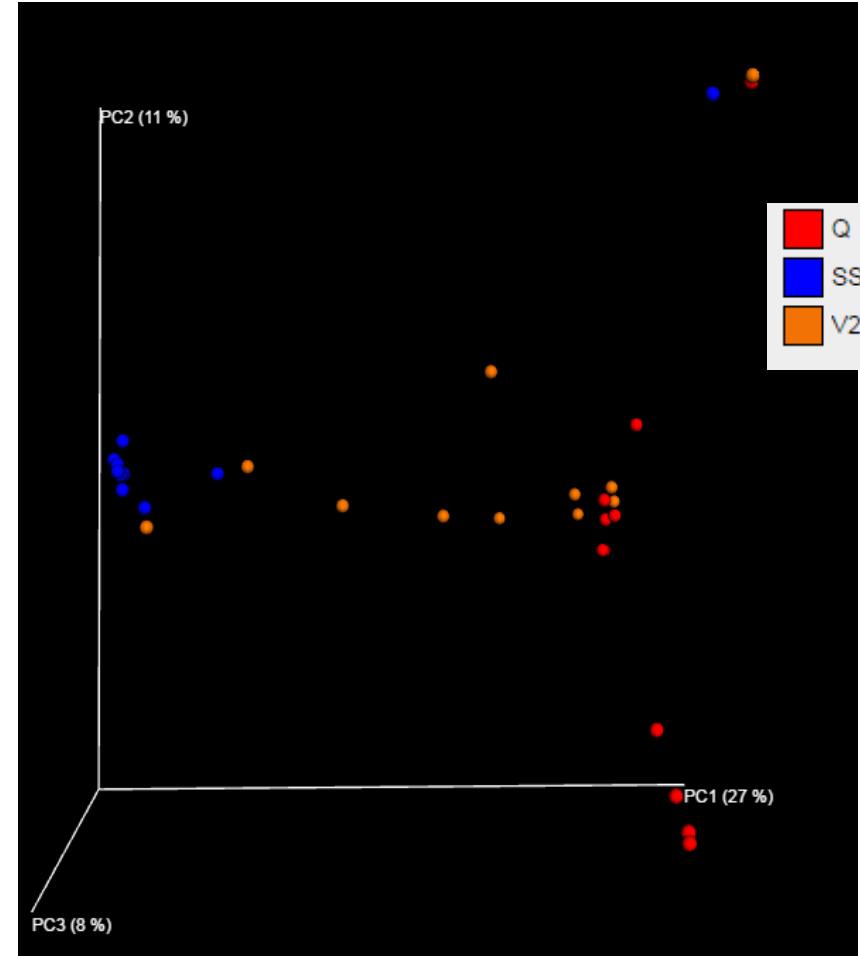
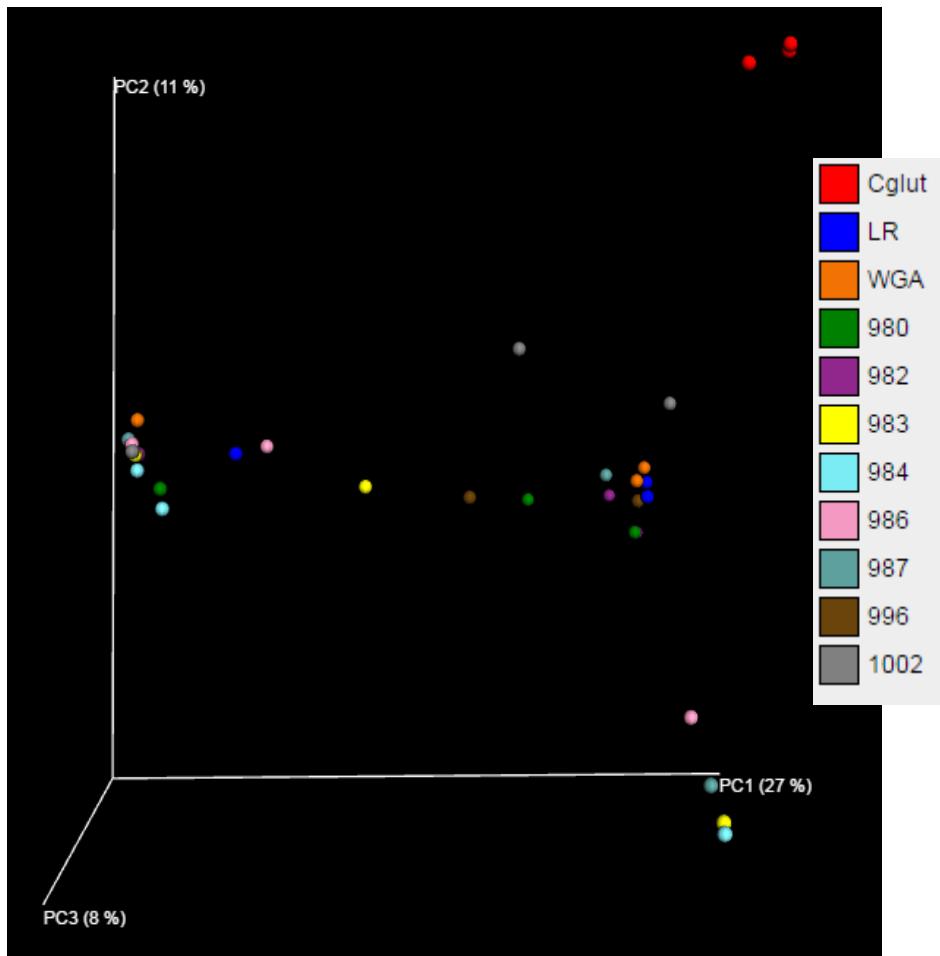
Sample	LMAT Identifications (Number of reads)		
	Qiagen REPLI-g	Illustra Single Cell	Illustra V2
Culture-positive PJI			
980	Streptococcus (2,694,906)	Streptococcus (286,577), Toxoplasma (489)	Streptococcus (3,075,181), Pseudomonas (232), Anaerococcus (135)
982	Staphylococcus (3,272,895), Propionibacterium (195), Mupapillomavirus (137), Finegoldia (115)	Staphylococcus (102,934), Toxoplasma (131)	Staphylococcus (9,539,147), Pseudomonas (15,292), Acinetobacter (739), Pseudoperonospora (346), Streptococcus (303), Elizabethkingia (242), Malassezia (176), Propionibacterium (142)
986	Streptococcus (37,268), Staphylococcus (26,397), Prevotella (13,472), Haemophilus (11,513), Campylobacter (3,344), Propionibacterium (2,298), Gemella (2,045), Alloprevotella (1,756), Chryseobacterium (1,531), Rothia (1,458), Neisseria (913), Capnocytophaga (911), Acinetobacter (297), Corynebacterium (248), Alloiococcus (156)	Staphylococcus (424), Toxoplasma (118)	Staphylococcus (168,046), Haemophilus (69,045), Streptococcus (66,370), Pseudomonas (58,244), Corynebacterium (18,344), Elizabethkingia (12,471), Propionibacterium (7,678), Gemella (2,298), Neisseria (2,092), Rothia (1,186), Empedobacter (835), Acinetobacter (479), Fusobacterium (434), Lactobacillus (237), Prevotella (185), Granulicatella (144), Delftia (130), Achromobacter (126)
996	Bacteroides (155,761), Clostridium (4,115), Parabacteroides (191), Corynebacterium (168), Blautia (143)	Bacteroides (11,134), Toxoplasma (669), Clostridium (454)	Bacteroides (96,154), Clostridium (4,052), Pseudomonas (1,483), Yarrowia (563), Corynebacterium (364), Propionibacterium (331), Staphylococcus (345), Parabacteroides (239), Streptococcus (116), Anaerotruncus (104)
1002	Corynebacterium (536,155), Lactococcus (2,195), Acinetobacter (187), Propionibacterium (108)	Corynebacterium (5,503), Toxoplasma (346)	Corynebacterium (1,000,810), Pseudomonas (32,865), Staphylococcus (5,704), Propionibacterium (5,396), Lactococcus (3,008), Streptococcus (2,121), Acinetobacter (1,293), Dermabacter (634), Malassezia (424), Elizabethkingia (382), Achromobacter (224), Actinobaculum (123), Delftia (108), Debaryomyces (102), Arcanobacterium (101)
Culture-negative PJI			
984	Streptococcus (2,425), Acinetobacter (167)	Toxoplasma (230)	Pseudomonas (12,472), Acinetobacter (1,395), Elizabethkingia (936), Alloiococcus (706), Streptococcus (395), Staphylococcus (264), Achromobacter (205), Propionibacterium (169), Delftia (134), Malassezia (111)

Taxonomic Identification of Reads (Performed using LMAT, Grouped by Genus)

Sample	LMAT Identifications (Number of reads)		
	Qiagen REPLI-g	Illustra Single Cell	Illustra V2
Aseptic Failure			
983	Streptococcus (352), Gloeocapsa (137)	Toxoplasma (442)	Pseudomonas (42,315), Acinetobacter (788), Elizabethkingia (742), Staphylococcus (630), Streptococcus (475), Propionibacterium (289)
987	Acinetobacter (5,396), Streptococcus (1,179), Prevotella (433), Propionibacterium (346), Kurthia (122)	Toxoplasma (173)	Pseudomonas (642,812), Elizabethkingia (5,283), Propionibacterium (3,271), Peptoniphilus (1,742), Acinetobacter (560), Epilithonimonas (372), Pseudoperonospora (353), Anaerococcus (255), Streptococcus (188), Capnocytophaga (172), Malassezia (158)
Controls			
C. glutamicum	Corynebacterium (26,912,706), Rothia (1,070), Streptococcus (545), Neisseria (221), Haemophilus (240)	Corynebacterium (22,276,446), Streptococcus (333), Toxoplasma (126)	Corynebacterium (26,149,302), Pseudomonas (627), Streptococcus (289)
Ringers	Neisseria (10,575), Streptococcus (8,441), Dolosigranulum (3,912), Haemophilus (716), Propionibacterium (385)	Streptococcus (1,192), Toxoplasma (969), Staphylococcus (929), Propionibacterium (127), Corynebacterium (123)	Pseudomonas (301,181), Staphylococcus (286,646), Streptococcus (135,065), Yarrowia (40,273), Propionibacterium (27,201), Achromobacter (15,251), Elizabethkingia (8,617), Acinetobacter (4,813), Haemophilus (4,423), Alloiococcus (2,610), Granulicatella (2,381), Malassezia (2,557), Delftia (681), Fusarium (671), Corynebacterium (593), Veillonella (498), Alloprevotella (384), Chryseobacterium (216), Dolosigranulum (234), Neisseria (177), Aspergillus (165)
WGA no template	Achromobacter (5,072), Rothia (1,031), Micrococcus (451), Delftia (110)	Sphingomonas (35,596), Staphylococcus (6,414), Propionibacterium (3,596), Peptoniphilus (697), Finegoldia (196), Elizabethkingia (149)	Pseudomonas (5,297,561), Propionibacterium (1,557,410), Staphylococcus (407,691), Lactococcus (297,420), Elizabethkingia (134,333), Malassezia (89,791), Delftia (13,668), Acinetobacter (7,858), Phyllobacterium (2,808), Comamonas (3,175), Chryseobacterium (1,732), Achromobacter (1,734), Corynebacterium (1,061), Rhodococcus (1,231), Micrococcus (452), Polynucleobacter (426), Streptococcus (274), Wolbachia (223), Alternaria (220)

PCoA Plots - β -diversity Between Samples

Fastq files depleted of human and PhiX sequences concatenated and HUMAnN2 (version 0.5.0) used to determine gene abundances of organisms present; QIIME used to calculate β -diversity and generate PCoA plots.



119 Culture-positive PJs

102 identical results

11 samples additional pathogens detected

- 4 Polymicrobial mixes
- 2 *S. aureus*
- 1 *S. epidermidis*
- 1 *S. agalactiae*
- 1 *Clostridium* sp.
- 1 *Peptoniphilus harei*
- 1 *P. acnes*

6 samples pathogens not detected

- 1 *Bacillus* sp.
- 1 *M. abscessus*
- 1 *P. acnes* (sonicate fluid no growth)
- 1 *Porphyromonas* sp.
- 1 *S. epidermidis*
- 1 *P. aeruginosa*

51 Culture-negative PJs

39 No pathogens detected

12 Potential pathogens detected

- 2 *S. dysgalactiae*
- 2 *Corynebacterium* sp.
- 2 *S. aureus*
- 1 Polymicrobial mix
- 1 *C. albicans*
- 1 *S. agalactiae*
- 1 *S. epidermidis*
- 1 *Mycoplasma salivarium*

162 Aseptic Failures

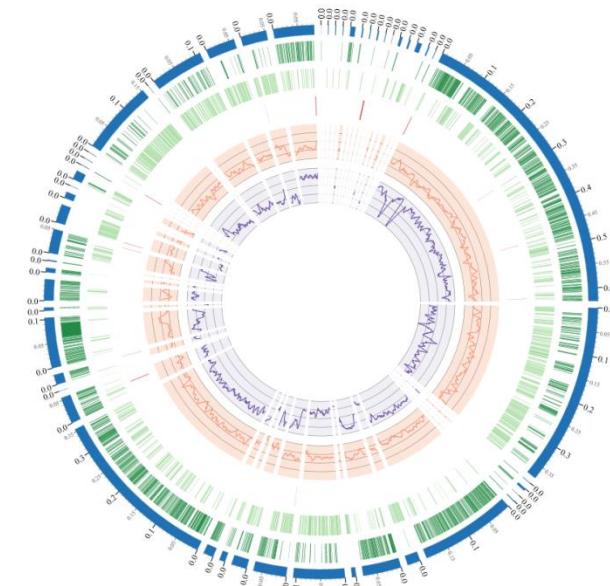
158 No pathogens detected

4 Potential pathogens detected

- 3 *S. aureus*
- 1 *Streptococcus sanguinis*

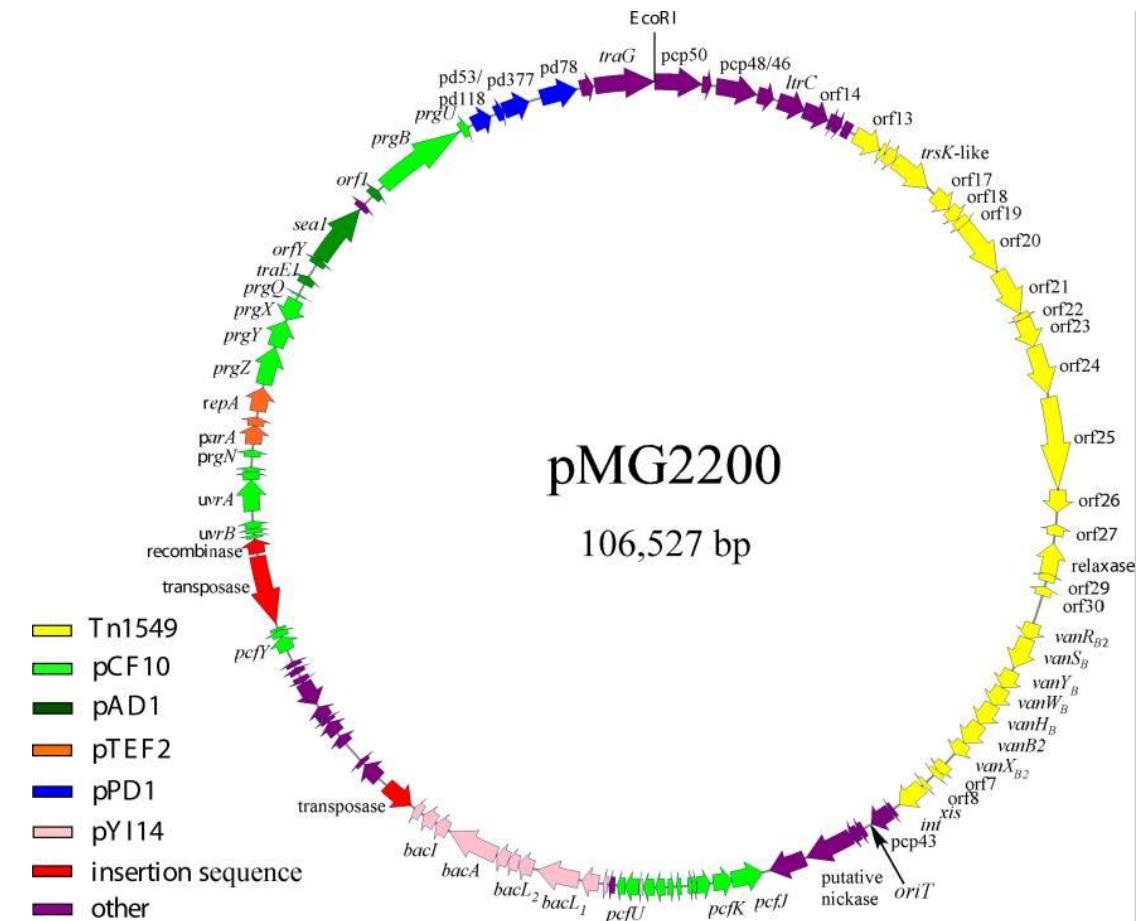
S. aureus genome assembly

- Used PATRIC online tools to assemble and annotate genome
- 2,822,458 bp genome
 - Typically 2.7 to 3.0 Mb
- Able to identify antibiotic resistance genes (*mecA*, *ermB*, *norA*)



E. faecalis plasmid analysis

- Reads identified mapping to multiple plasmids, pMG2200 most common
- pMG2200: plasmid with vancomycin resistance genes

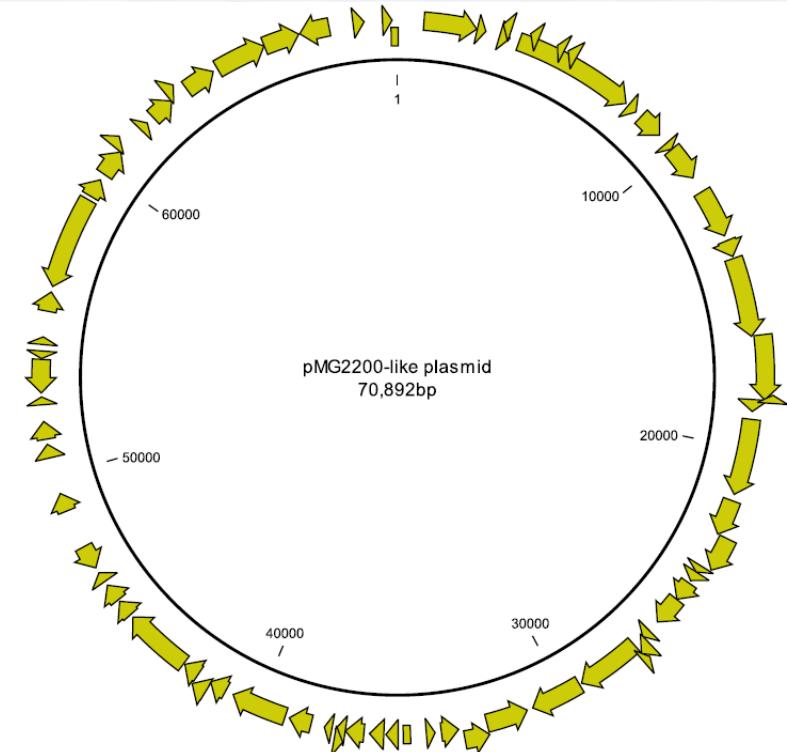


Analysis of pMG2200-Like Plasmid

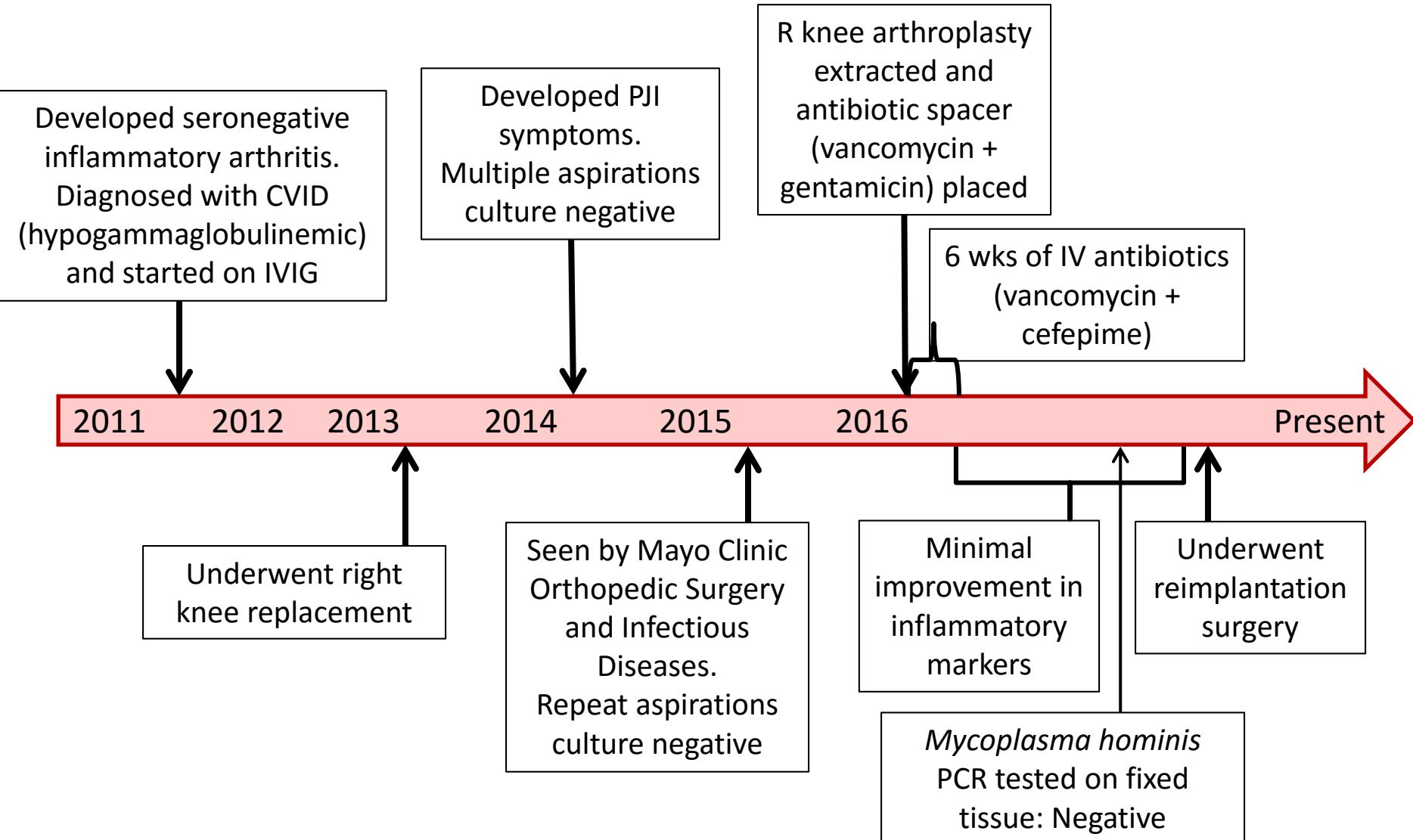
Alignment of sequences to pMG2200



- If pull sequences labeled as plasmids, can assemble into single 70,892 bp fragment
- High homology to pMG2200 with transposon-like element absent



53 yo Man – Right Knee PJI



Metagenomic Analysis From Resected Arthroplasty

Number of Reads (out of 27,984,652)	Taxonomic Identification
1,796	<i>Mycoplasma salivarium</i>
58	<i>Acinetobacter junii*</i>
19	<i>Propionibacterium acnes*</i>
5	<i>Plasmodium gaboni*</i>
5	<i>Salpingoeca rosetta*</i>
3	<i>Aspergillus</i> sp.*

Follow-up Studies

- Second sonicate fluid aliquot from resection surgery:
 - Metagenomic analysis without MolYsis enrichment
 - 16S rRNA gene PCR of sonicate fluid: *M. salivarium*

Metagenomic Analysis without MolYsis

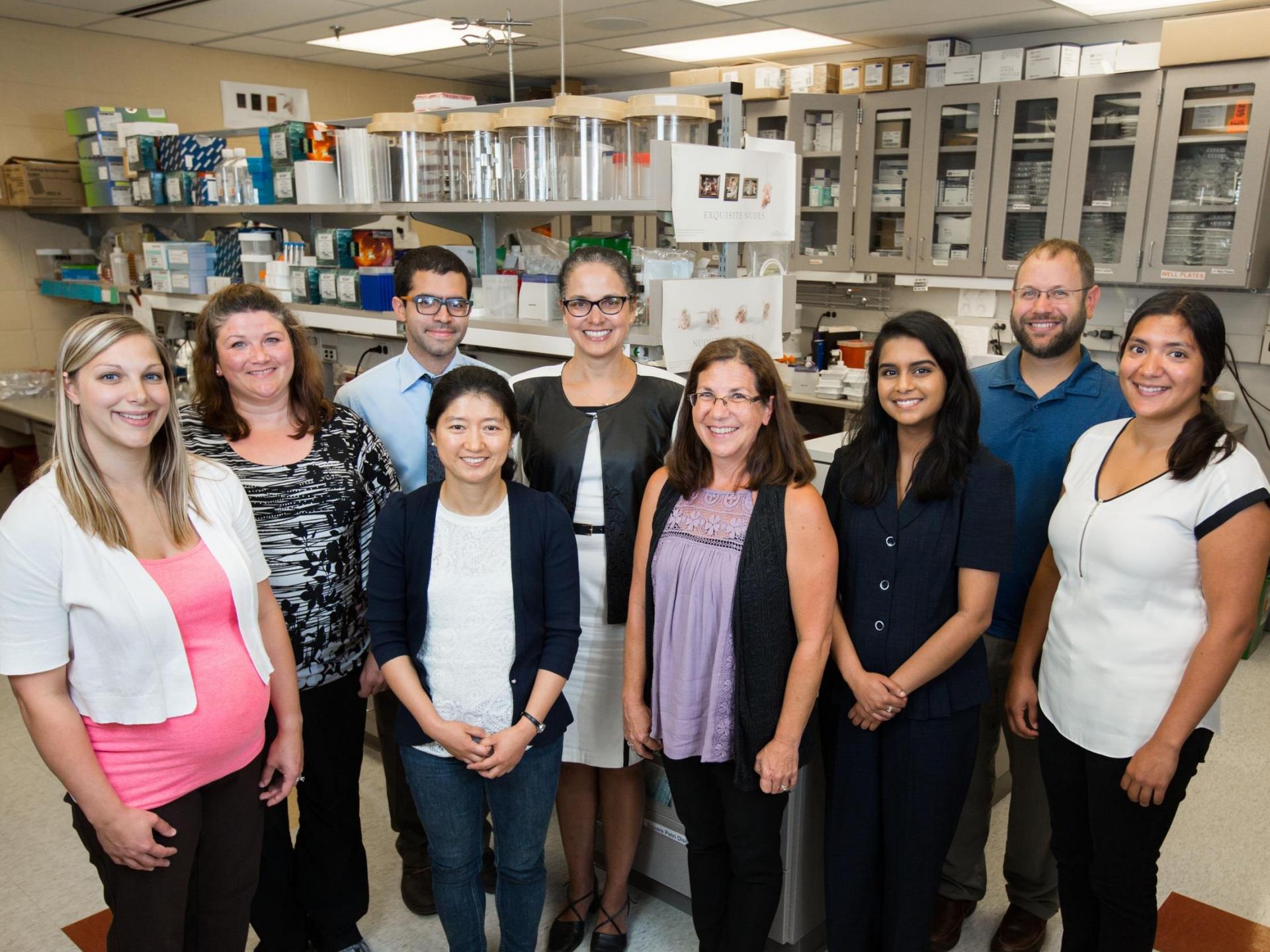
Number of Reads (out of 234,088,137)	Taxonomic Identification
19,642	<i>Mycoplasma salivarium</i>
371	<i>Propionibacterium acnes*</i>
247	<i>Pseudomonas stutzeri*</i>
143	<i>Toxoplasma gondii*</i>
86	<i>Corynebacterium sp.*</i>
73	<i>Acinetobacter junii*</i>

* Known background

Take-Home Points

“Prosthetic Joint Infection - Think Biofilms”

- Device cultures
 - That's where the biofilms are
- Culture in blood culture bottles
 - Synovial fluid
 - Periprosthetic tissues (just 3 needed!)
- Molecular diagnostics
 - Specimen type matters (tissue poor specimen)
 - Not first-line (? role as “rapid” diagnostics)
 - ?Challenging cases



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