

COLLEGE of AMERICAN PATHOLOGISTS

Microbiome Applications in Pathology

CAP Webinar



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• All lines are muted during the presentation

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• Dr. Bry has no disclosures.



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The Microbiome



 Communities of microbes that colonize all body surfaces.

- 10X more microbial cells in the human body than those of the host.
- Important in health and disease.
- Exponential growth in publications and commercial opportunities.

Grice EA, Segre JA. The human microbiome: our second genome. Annu Rev Genomics Hum Genet. 2012;13:151-70. doi: 10.1146/annurev-genom-090711-163814.

Impact of host-microbiome interactions

100% of the US population has been or will be affected by microbiome-promoted diseases and conditions.

Condition	US residents/yr	Role of the Microbiome
Periodontitis, dental caries	>85,000,000	Confirmed
Take medications where microbial biotransformation causes side effects	>50,000,000	Confirmed
Inflammatory Bowel Diseases	1,300,000	Confirmed
Have had <i>C. difficile</i> colitis	500,000	Confirmed
Pre-term birth	400,000	Strong evidence
GI and Oral Cancers	2,500,000	Strong evidence
Food allergies	6,000,000	Strong evidence
Type II Diabetes and pre-diabetic conditions	>70,000,000	Hypothesized
Cardiovascular disease	>84,000,000	Hypothesized

Data from the US Census, MA DPH, CDC, CCFA, AHF, NIH, and Partners Healthcare's Research Patient Data Registry

CAF

Growing interest from industry

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Mining the Microbiome for Transformative Products



Janssen Global Services. Mining the Microbiome for Transormative Products. Available at http://www.janssen.com/human-microbiome-institute. Accessed October 6, 2016.

Pathology applications

Current applications

- *C. difficile* colitis diagnosis and FMT donor screening
- Infectious disease applications using NGS/WGS metagenomic reads
 - Unbiased methods for pathogen detection
- Women's health
- Oral health: periodontitis, dental caries
 - Impact of oral health to other disease conditions: diabetes, pregnancy

Future applications

- Will evolve as we have better mechanistic understanding of microbiota-mediated contributions to disease and targeted therapies
- May leverage existing AP/CP testing (eg, bile acids, immunology, targeted organisms)
- Community effects vs defined biomarkers
- Not limited to effects on colonized surfaces
 - Endocrine, cardiovascular, non-IBD/auto-immune, CNS
 - Carcinogenesis and cancer treatment/prognosis



Bedside Bench Bedside Infrastructure

Move the field from descriptive associations to causal effects of the microbiota *in vivo*







Bucci V, Tzen B, Li N, Simmons M, Tanoue T, Bogart E, Deng L, Yeliseyev V, Delaney ML, Liu Q, Olle B, Stein RR, Honda K, Bry L, Gerber GK. MDSINE: Microbial Dynamical Systems INference Engine for microbiome time-series analyses. Genome Biol. 2016 Jun 3;17(1):121. doi: 10.1186/s13059-016-0980-6.

Methods for studying the microbiome

Next generation sequencing

- 16S rRNA gene phylotyping
- Metagenomics
- Virome
- Eukaryotic colonizers (fungi, protozoa, parasites)
- Other molecular methods
 - Targeted probes
 - Hybrid methods: immune or metabolite capture
- Metabolomics
 - Microbial metabolites in directed vs undirected fashion
 - SCFA GC/LC



– Mass-spec profiles

Methods for studying the microbiome, continued

- Microbiologic
 - Culture-based methods
 - Antigen detection
 - Microbial genetics and synthetic biologic techniques
- Computational
 - Bioinformatic tools: OTU clustering, metagenomics
 - Longitudinal dynamics, outcomes prediction, principled models
 - Essential to make sense of complex datasets and distinguish signal from noise



16S rRNA gene phylotyping

- Amplification over conserved regions of 16S rRNA (ITS, other conserved targets)
- NGS of amplified product (MiSeq, IonTorred)
 - Short read platforms limit to V4, V1-3, V3-4 or 3-5
- Bioinformatic/computational methods to get to operational taxonomic units

(OTU)

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16S rRNA gene phylotyping

- Has revolutionized evaluation of complex microbial ecosystems
- Limits with resolution and detection of ecosystem members
- Research use only assay





Before phylotyping methods

16S phylotyping on short-read platforms..

Where methods need to go

Sequence vs culture or probe-based methods Complex ecosystems >10¹⁰ CFU/g



Organisms Present

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Credit: Felipe Esquivel Reed



- Component members and gene content
- More expensive and computationally intensive; needs for curated reference data
- Clinical applications for unbiased pathogen detection
 - CNS other sterile body sites

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- Immunocompromised patients (urine in kidney transplant patients)





Delwart E. A roadmap to the human virome. PLoS Pathog. 2013 Feb;9(2):e1003146. doi: 10.1371/journal.ppat.1003146.

Metabolomics/metabolite phenotyping

- MassSpec, GC/LC: Microbial ID via MALDI-TOF, SCFA profiles
- Important for defining "dysbiosis" and microbial factors that can be used diagnostics or to predict patient outcomes
 - Incorporate with host-makers



Institute for Molecular Medicine Finland (FIMM). FIMM Metabolomics Work Flow. Available at https://www.fimm.fi/en/services/ technologycentre/metabolomics. Accessed October 6, 2016.

Pathogen genome sequencing

NGS-platforms, commonly MiSeq

- HIV/HCV genotyping: used in other countries. FDA-approved tests not yet available in the US
- Hospital and national surveillance activities, difficult to culture species



Genomic analysis pipeline

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CRE: Epidemiologic Analyses



Plasmid	MLST	Strain
P34399-43.500kb	ST-258	BWH-C10
P34399-43.500kb	ST-258	BWH-C8
pKPC-484	ST-258	BWH-C1
pKPC-484	ST-258	BWH-C2
chromosomal	ST-258	BWH-C5
chromosomal, pBK15692	ST-258	BWH-C6
pBWH-C7-KPC	ST-258	BWH-NC12
No KPC	ST-258	BWH-C7

B) E. cloacae



Plasmid	MLST	Strain
pBWH-C13-KPC	ST-171	BWH-C14
pBWH-C13-KPC	ST-171	BWH-C15
pBWH-C13-KPC	ST-171	BWH-C13
No KPC	ST-78	BWH-NC28
pBWH-C16-KPC	ST-78	BWH-C16
pBWH-C16-KPC	ST-78	BWH-C17
ME-BWH-C11-KPC	ST-78	BWH-C11
ME-BWH-C11-KPC	ST-78	BWH-C12
No KPC	ST-78	BWH-NC16



Nicole D. Pecora et al. *mBio* 2015; doi:10.1128/mBio.01030-15 **mBio**

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Case Studies: Host-Microbiome Systems

- (1) <u>Infectious Disease:</u> C. difficile colitis: development of therapeutic microbiota
- (2) Food allergies: Integrated systems identifying therapeutic human microbiota
- (1) <u>Dysbiosis in disease</u>: Therapeutic microbiota and small molecule targets in IBD
- (2) <u>Microbiota-mediated effects in cancer</u> development and treatment
- (1) <u>Drug and xenobiotic metabolism</u>: IBD and cancer therapeutics; other drug classes

Clostridium difficile colitis

Pseudomembranous colitis

- 5-10% population colonized with *C. difficile*
 - Not all strains are toxigenic
- Substantive morbidity and mortality, particularly in immunocompromised patients
- 10-20% of patients fail antibiotic treatment to develop recurrent C. diff.
- Microbiota-mediated protection
 - Positive and negative effects from primary and secondary bile acids on C. diff germination
 - Competition for nutrients, colonization niches



Yu J, Kim NY, Lee HM, Lee HN, Ahn HJ, Kim SW, Choi KY. A Case of Pseudomembranous Colitis in a Juvenile Rheumatoid Arthritis Patient Taking Methotrexate. Korean J Gastroenterol. 2010 Dec;56(6):387-390. https://doi.org/10.4166/kjg.2010.56.6.387



Clostridium difficile colitis, continued

- Therapy for recurrent infection: Fecal Microbiota
 - **Transplant/FMT**
 - Colonoscope vs oral capsule
 - OpenBiome
 - Seres Therapeutics, others
- Target therapy per missing microbial activities



Yu J, Kim NY, Lee HM, Lee HN, Ahn HJ, Kim SW, Choi KY. A Case of Pseudomembranous Colitis in a Juvenile Rheumatoid Arthritis Patient Taking Methotrexate. Korean J Gastroenterol. 2010 Dec;56(6):387-390. https://doi.org/10.4166/kjg.2010.56.6.387





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Infectious Disease Models: C. difficile colitis



- Defined formulations will be available for treatment of recurrent *C. diff*

Pathology applications:

- Biomarkers to assess patients at risk of recurrence, predict successful therapy
 - Stool-based analytes vs microbial community signatures
- As blood banks manage blood transfusion will micro labs manage therapeutic microbial communities, whether from vendor or in-house sources



Bucci V, Tzen B, Li N, Simmons M, Tanoue T, Bogart E, Deng L, Yeliseyev V, Delaney ML, Liu Q, Olle B, Stein RR, Honda K, Bry L, Gerber GK. MDSINE: Microbial Dynamical Systems INference Engine for microbiome time-series analyses. Genome Biol. 2016 Jun 3;17(1):121. doi: 10.1186/s13059-016-0980-6.



Therapeutic microbiota

NegCC



Therapeutic microbiota for treat-to-prevent and treat-to-cure regimens

Pathology applications:

- Identify dysbiotic communities vs. absence of critical members
- Causative microbial pathways and/or products and metabolites
- Follow response to treatment (microbiome and routine immunologic CAP analyses)



Mast Cell Responses - tissue localization and MMCP-1 production



Dysbiosis in Disease



Pathology Applications:

- Specific biomarkers that are predictive of dysbiotic conditions
 - Microbial metabolites or products
 - NASH, subsets of diabetic/pre-diabetic patients
 - Host markers
 - Metabolic, immune, hormonal
 - IBS, auto-immune and allergic diseases
 - **Computational modeling of patient-microbial dynamics**

Microbiome in Cancer

Direct and indirect effects on carcinogenesis

- Microbial biotransformations: carcinogens, ingested cyanogenic glycosides, environmental toxicants.
- Chronic inflammation: *H pylori*, HCV
- Epithelial cell turnover -> commensal interactions with APC/MIN and other cell division pathways

Impact on drug efficacy and novel drug targets

- Immunomodulatory functions => efficacy of checkpoint inhibitors
- Direct biotransformation of oral and IV-administered anti-neoplastics: Irinotecan => microbiota-mediated toxicities
- Gut commensal communities and host susceptibility to GVHD post-SCT

Pathology Applications:

- Assessment of microbial communities impacting drug efficacy or toxic profiles
- Identify pathobionts -> pathogens important in diagnosis and management
- Small molecule targets on host or
- microbial side important for dx and
- CAP therapeutic implications



Ohaegbulam KC, Assal A, Lazar-Molnar E, Yao Y, Zang X. Human cancer immunotherapy with antibodies to the PD-1



Louis, P., et al. (2014). "The gut microbiota, bacterial metabolites and colorectal cancer." Nat Rev Micro 12(10): 661-672.

Drug metabolism

Microbiota-mediated biotransformation of drugs
Microbiota-mediated Toxicities
Altered or Reduced Efficacy







NSAIDS

Irinotecan

Sulfasalazine



Pathology Applications:

- Assay for microbial drug-transforming activities
 - Microbial beta-glucuronidases, sterol metabolizing enzymes
- Contributions to dysbiosis
 - Immunomodulators, anti-microbials, altered host factors (bile acids, antibodies)
- Immunologic competence for therapeutics to act
 - Checkpoint inhibitors necessary microbial activities or postintervention to boost activities (microbial and immunologic markers)
- Incorporate with therapeutic drug monitoring (TDM)



Pathology/Pathologists and the Microbiome

- Sanity check for moving things to productive clinical use
 - Assessment of the literature
 - Assuring robust evidence for clinical use
 - Participating in and providing infrastructure for clinical trials
- New and existing methods and resources will be used
 - Microbiology, chemistry, TDM, immunology, AP services, others
 - Need for incorporation of computational models
 - Warehousing of microbiome and pathogen genomic information



"Bugs as drugs" – oversight and quality programs

BWH

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Credit: Anthony Howe



Exploring Korea. Modo Sculpture Park. Available at http://www.exploringkorea.com/modosculpture-park-full-house-set/. Accessed October 6, 2016.

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B 16, 15	American Association Of Forensic Sciences (AAFS) Hyst Regency Ortanos, PL	2015 Policy Meeting
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Short Presentations on Emerging Concepts (SPECS)

- Pathology SPECs are:
 - short PowerPoints, created for pathologists
 - Focused on diseases where molecular tests play a key role in patient management
- New topics are Renal Tumors, cell free DNA (cfDNA), and PD-L1 as well as other emerging topics
- Access them <u>www.cap.org</u> > Resources and Publications





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 - The digital copy of the Resource Guides are a complimentary member benefit
 - Access them <u>www.cap.org</u> > Resources and Publications







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