

**CURRICULUM VITAE**

<b>NAME</b>	Blake M. Hanson, M.S., Ph.D.
<b>PRESENT TITLE(s)</b>	Assistant Professor Center for Infectious Diseases Department of Epidemiology, UTHealth School of Public Health at Houston University of Texas Health Science Center at Houston  Faculty, Regular Member Graduate School of Biomedical Sciences University of Texas Health Science Center at Houston  Adjunct Assistant Professor of Medicine Department of Medicine Houston Methodist Academic Institute
<b>ADDRESS</b>	RAS E739 1200 Pressler Street Houston, TX 77030
<b>UNDERGRADUATE EDUCATION</b>	Augustana College Rock Island, IL September 2004 - May 2008 Degree: Bachelor of Arts
<b>GRADUATE EDUCATION</b>	The University of Iowa, College of Public Health Iowa City, IA May 2010 – December 2013 Degree: Doctorate of Philosophy, Epidemiology Thesis: “Determination of the prevalence and incidence, molecular characterization, and nasal and pharyngeal colonization patterns of <i>Staphylococcus aureus</i> among urban and rural Iowans”  The University of Iowa, College of Education Iowa City, IA September 2011 - May 2013 Degree: Graduate Certificate in College Teaching  The University of Iowa, College of Public Health Iowa City, IA September 2008 - May 2010 Degree: Master of Science, Epidemiology
<b>POSTGRADUATE TRAINING</b>	Postdoctoral Associate Weinstock Laboratory The Jackson Laboratory for Genomic Medicine

January 2014 – August 2017

**ACADEMIC  
APPOINTMENTS**

- October 2017 – Present  
Assistant Professor  
Center for Infectious Diseases  
Department of Epidemiology, UTHealth School of Public Health at Houston  
University of Texas Health Science Center at Houston
- March 2024 – Present  
Adjunct Assistant Professor of Medicine  
Houston Methodist Academic Institute  
Houston, Texas
- April 2022 – Present  
Faculty, Regular Member  
Graduate School of Biomedical Sciences  
University of Texas Health Science Center at Houston
- September 2021 – September 2024  
Assistant Professor  
Division of Pediatric Infectious Diseases, McGovern Medical School  
University of Texas Health Science Center at Houston
- October 2017 – September 2024  
Associate Director of CARMiG  
The University of Texas Health Sciences Center  
Houston, Texas
- October 2017 – September 2021  
Assistant Professor  
Division of Infectious Diseases, McGovern Medical School  
University of Texas Health Science Center at Houston
- January 2014 – August 2017  
Postdoctoral Associate  
Mentor: Dr. George Weinstock  
Weinstock Laboratory  
The Jackson Laboratory for Genomic Medicine  
Farmington, Connecticut
- October 2008 – December 2013  
Graduate Research Assistant  
Center for Emerging Infectious Diseases, The University of Iowa  
College of Public Health  
Iowa City, Iowa
- October 2008 – December 2010  
Graduate Research Assistant  
Health Effectiveness Research Center, The University of Iowa  
College of Public Health

Blake M. Hanson, M.S., Ph.D.

Iowa City, Iowa

**PROFESSIONAL  
ORGANIZATIONS**

Local

- Iowa Academy of Science (2009-2010)

National

- American Heart Association (2024 – Current)
- American Society of Microbiology (2009 – Current)
- American Public Health Association (2009, 2024 – Current)
- Infectious Disease Society of America (2013, 2023 – Current)

**HONORS AND AWARDS**

September 2020 – November 2023, Antimicrobial Resistance Leadership Group (ARLG) Trialist in Training

January 2015 – March 2017, JAX Scholar Postdoctoral Fellow  
The Jackson Laboratory for Genomic Medicine

April 2013, Oral Presentation Student Award Winner  
2<sup>nd</sup> Great Plains Emerging Infectious Diseases Conference, Iowa City, Iowa, 2013

April 2013, Poster Session Award Winner  
The University of Iowa Health Sciences Research Week  
The University of Iowa

September 2011, ASM student presenter travel grant  
Funded attendance to the 2<sup>nd</sup> ASM conference on methicillin-resistant staphylococci in animals, Boston, USA, 2011

May 2009 – May 2010, Elected Vice President  
The College of Public Health Student Association  
College of Public Health  
The University of Iowa

May 2009 – May 2010, Elected President  
Epidemiology Student Association  
College of Public Health  
The University of Iowa

September 2009, ASM student presenter travel grant  
Funded attendance to the ASM conference on methicillin-resistant staphylococci in animals, London, England, 2009

**PAST AND CURRENT  
TEACHING ACTIVITIES**

Course Instructor, Epidemiology and Control of Infectious Diseases

January 2020 – Current  
Course, design, discussion facilitation, instruction  
The University of Texas School of Public Health

Course Instructor, Epidemiology III  
June 2018 – Current

Blake M. Hanson, M.S., Ph.D.

Course design, discussion facilitation, instruction  
The University of Texas School of Public health

Journal Club Coordinator, Summer Student Journal Club  
June 2014 – August 2014  
Course design, discussion facilitation  
The Jackson Laboratory for Genomic Medicine

Teaching Assistant, Epidemiology III: Theories  
August 2013 – December 2013  
Course design, discussion facilitation, teach select lectures  
The University of Iowa

Teaching Assistant, Epidemiology of Infectious Diseases  
August 2012 – December 2012  
Assist in course design and teach select lectures  
The University of Iowa

Teaching Assistant, Epidemiology II: Advanced Methods  
January 2012 – May 2012  
Assist in course design and teach select lectures  
The University of Iowa

Journal Club Coordinator, Department of Epidemiology  
October 2011 – May 2013  
Bi-weekly journal club  
The University of Iowa

Teaching Assistant, Zoonotic Diseases  
June 2011  
Two-week summer course  
The University of Iowa

Video Session Coordinator, Epidemiology of Infectious Diseases  
August 2010 – December 2010; August 2011 – December 2011  
Video Selection, Core Curriculum  
The University of Iowa

Seminar Coordinator, Department of Epidemiology  
May 2010 – May 2012  
Weekly seminar course  
The University of Iowa

**SERVICE – THE  
UNIVERSITY OF TEXAS  
HEALTH SCIENCES  
CENTER**

Co-Coordinator for The Infectious Disease Epidemiology  
Certificate  
August 2023 - Current  
School of Public Health

Texas Epidemic Public Health Institute Wastewater Committee  
August 2022 – Current  
School of Public Health

Blake M. Hanson, M.S., Ph.D.

Texas SARS-CoV-2 Variant Detection Network Steering Committee  
University of Texas Health Sciences Center  
October 2021 – Current

UTHealth COVID-19 Clinical Research Workgroup  
July 2020 – January 2022

SPH Preliminary Exam Committee  
August 2019 – June 2022  
Department of Epidemiology  
School of Public Health

SPH Center for Infectious Disease Faculty Search Committee  
August 2019 – September 2021  
Department of Epidemiology  
School of Public Health

SPH Human Genetics Center Faculty Search Committee  
August 2019 – July 2020  
Department of Epidemiology  
School of Public Health

SPH IT Faculty Advisory Committee  
January 2019 – current  
School of Public Health

#### **SERVICE – EXTERNAL**

Early Career Reviewer – NIH Etiology, Diagnostic, Intervention and Treatment (EDIT) study section  
June 2024

American Heart Association Fellowship Reviewer – Genomics and Translational Biology committee  
2023 Cycle – Current

Antimicrobial Resistance Leadership Group Diagnostics Committee  
February 2022 – Current

American Heart Association Career Development Award Reviewer – Population committee  
2022 Cycle

Antimicrobial Agents and Chemotherapy – Editorial Board  
January 2021 – Current

#### **SERVICE – THE JACKSON LABORATORY**

Postdoctoral Association Co-Founder  
February 2014 – August 2017  
The Jackson Laboratory for Genomic Medicine

**SERVICE – THE  
UNIVERSITY OF IOWA**

Seminar Coordinator  
December 2011 – May 2013  
Department of Epidemiology  
College of Public Health

Journal Club Coordinator  
May 2010 – May 2012  
Department of Epidemiology  
College of Public Health

**SPONSORSHIP OF  
CANDIDATES FOR  
GRADUATE DEGREE**

Jesus Duran Ramirez  
PhD Student in Microbiology and Molecular Genetics  
2022 – Current

Hossaena Ayele  
PhD Candidate in Epidemiology  
2021 – Current

An Q. Dinh  
PhD Student in Epidemiology  
2021 – Current

Shezeen Gillani  
MPH degree conferred 2023  
PhD Student in Epidemiology  
2020 – Current

Shelby R. Simar, MPH, PhD  
MPH degree conferred 2019  
PhD conferred 2024  
2017 – 2024

William C. Shropshire, MPH, PhD  
PhD conferred 2020  
2017 – 2020

**SERVICE ON STUDENT  
THESIS AND EXAM  
COMMITTEES**

Jesus Duran Ramirez, MPH  
PhD Thesis  
Role: Committee Member  
Institution: MD Anderson-UTHealth Houston Graduate School of Biomedical Sciences  
Program: Microbiology and Molecular Genetics  
Fall 2023 – Current

Jana Gomez, MS  
PhD Thesis  
Role: Committee Member  
Institution: MD Anderson-UTHealth Houston Graduate School of Biomedical Sciences  
Program: Microbiology and Molecular Genetics  
Summer 2022 – Current

Gretchen Morrison  
PhD Thesis  
Role: Committee Member  
Institution: MD Anderson-UTHealth Houston Graduate School of Biomedical Sciences  
Program: Microbiology and Molecular Genetics  
Spring 2022 – Current

Olivia Morris  
MPH Thesis  
Role: Thesis Chair  
Institution: UTHealth Houston  
Program: Epidemiology  
Fall 2022 – Summer 2023

William Shutze, MPH  
Integrated Learning Experience  
Role: Integrated Learning Experience Preceptor  
Institution: UTHealth Houston  
Program: Epidemiology  
Fall 2022 – Spring 2023

Rachel Atterstrom, MPH  
MPH Thesis  
Role: Thesis Chair  
Institution: UTHealth Houston  
Program: Epidemiology  
Fall 2022 – Spring 2023

Sreyya Sudharsan Raamasamy, MS  
MS Thesis  
Role: External Committee Member  
Institution: MD Anderson  
Program: Diagnostic Genetics  
Summer 2022 – Summer 2023

Jesus Duran Ramirez, MPH  
MPH Thesis  
Role: Committee Member  
Institution: UTHealth Houston  
Program: Epidemiology  
Fall 2021 – Spring 2022

Shelby R. Simar, MPH, PhD  
PhD Thesis  
Role: Thesis Chair  
Institution: UTHealth Houston  
Program: Epidemiology  
Spring 2021 – Summer 2024

Shezeen Gillani, MPH  
MPH Thesis

Blake M. Hanson, M.S., Ph.D.

Role: Thesis Supervisor  
Institution: UTHealth Houston  
Program: Epidemiology  
Spring 2021 – Fall 2021

Fernanda Valasco-Huerta, MPH  
MPH Thesis  
Role: Committee Member  
Institution: UTHealth Houston  
Program: Epidemiology  
Spring 2021 – Fall 2021

Anika Stakov, MPH  
MPH Thesis  
Role: Committee Member  
Institution: UTHealth Houston  
Program: Epidemiology  
Spring 2021 – Summer 2021

Katherine Bay, MPH, PhD  
PhD Thesis  
Role: Committee Member  
Institution: UTHealth Houston  
Program: Epidemiology  
Spring 2021 – Spring 2022

Marissa Gagliardo, MPH  
MPH Thesis  
Role: Committee Member  
Institution: UTHealth Houston  
Program: Epidemiology  
Fall 2020 – Summer 2021

Yasmine Hoballah, MS  
MS Thesis  
Role: External Committee Member  
Institution: MD Anderson  
Program: Diagnostic Genetics  
Summer 2020 – Summer 2021

Chioma Odo, PhD  
PhD Thesis  
Role: External Committee Member  
Institution: MD Anderson-UTHealth Houston Graduate School of Biomedical Sciences  
Program: Microbiology and Molecular Genetics  
Fall 2019 – Spring 2024

Chioma Odo, PhD  
PhD Comprehensive Exam  
Role: External Committee Member  
Institution: MD Anderson-UTHealth Houston Graduate School of Biomedical Sciences

Blake M. Hanson, M.S., Ph.D.

Program: Microbiology and Molecular Genetics  
Fall 2019 – Spring 2020

Camden Hallmark, MPH, PhD  
PhD Thesis  
Role: Thesis Chair  
Institution: UTHealth Houston  
Program: Epidemiology  
Spring 2019 – Spring 2021

William C. Shropshire, MPH, PhD  
PhD Thesis  
Role: Thesis Chair, Thesis Supervisor  
Institution: UTHealth Houston  
Program: Epidemiology  
Fall 2018 – Fall 2020

Kirsten Bevan Rydell, MPH  
Master's Thesis  
Role: Thesis Chair, Thesis Supervisor  
Institution: UTHealth Houston  
Program: Epidemiology  
Fall 2018 – Spring 2019

Shelby R. Simar, MPH, PhD  
MPH Thesis  
Role: Thesis Chair, Thesis Supervisor  
Institution: UTHealth Houston  
Program: Epidemiology  
Fall 2018 – Spring 2019

**MENTEES WHO HAVE  
SECURED COMPETATIVE  
FELLOWSHIP FUNDING  
SUPPORT**

Mentee: Henok Tegegne  
Fellowship Type: Postdoctoral  
Agency: NIH NIAID  
Type of Grant: T32  
Fellowship Title: *Texas Medical Center Training Program in Antimicrobial Resistance (TPAMR)*  
Role: Co-Mentor  
Dates of Award: September 2023 – August 2025

Mentee: Mahesh Prasad Bekal  
Fellowship Type: Postdoctoral  
Agency: Texas CPRIT  
Type of Grant: Fellowship  
Fellowship Title: *McWilliams School of Biomedical Informatics at UTHealth Houston, Genomics and Translational Cancer Research Training Program (BIG-TCR) Fellowship*  
Role: Co-Mentor  
Dates of Award: May 2023 – April 2025

Mentee: Taryn Eubank, PharmD  
Fellowship Type: Postdoctoral  
Agency: NIH NIAID

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Type of Grant: T32  
Fellowship Title: *Texas Medical Center Training Program in Antimicrobial Resistance (TPAMR)*  
Role: Co-Mentor  
Dates of Award: July 2021 – June 2023

Mentee: Shelby Simar, MS  
Fellowship Type: Predoctoral  
Agency: NIH NIAID  
Type of Grant: T32  
Fellowship Title: *UTHealth Molecular Basis of Infectious Disease Research Training Fellowship*  
Role: Mentor  
Dates of Award: September 2020 – August 2022

Mentee: Naomi Bier, PhD  
Fellowship Type: Postdoctoral  
Agency: NIH NCADS  
Type of Grant: TL1  
Fellowship Title: *UTHealth Center for Clinical and Translational Sciences TL1 Training Program Fellowship*  
Role: Mentor  
Dates of Award: May 2020 – April 2022

## RESEARCH SUPPORT

### A. ACTIVE RESEARCH SUPPORT

**NIH/NIAID: 2R01AI132354-06 (Krachler)  
Mechanoregulation of EHEC Virulence**

06/21/2024 – 06/20/2029

This project aims to understand how bacteria sense and respond to physical forces as environmental cues, particularly focusing on enterohemorrhagic *E. coli* (EHEC). Key discoveries include identifying the transcriptional regulator GrlA, which controls the LEE pathogenicity island encoded type 3 secretion system in response to mechanical stimuli. GrlA dissociates from the inner membrane under shear force, activating virulence factors. The project will further investigate the mechanoregulation of Shiga toxin and its impact on gut motility and microbial competition within the host.

**Role: Co-Investigator**

**Office of the Governor & US Department of Treasury: 4546301 (Hanson as a sub-project of TEPHI)  
Texas Epidemic Public Health Institute (TEPHI) Antimicrobial Resistance Wastewater Surveillance**  
11/08/2021 – 12/31/2026

This is an independent sub-project of the TEPHI funding. This project aims to utilize wastewater surveillance for monitoring antimicrobial resistance (AMR), offering a novel approach to detect resistant pathogens and their genes in healthcare settings, particularly hospitals. This study will collect wastewater samples from Memorial Hermann hospital to identify carbapenem-resistant (CR) and extended-spectrum beta-lactamase-resistant (ESBL-R) ESKAPE pathogens. By comparing wastewater findings with clinical isolates, the project aims to detect overlapping and unique resistance mechanisms, potentially identifying subclinical pathogens. This approach could serve as an early warning system for emerging resistance, enhancing infection control, and improving patient safety and public health outcomes.

**Role: Principal Investigator**

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**Office of the Governor & US Department of Treasury: 4546301 (Boerwinkle)**

**TEPHI Viral Wastewater Surveillance**

11/08/2021 – 12/31/2026

This is an independent sub-project of the TEPHI funding. This project aims to utilize comprehensive wastewater surveillance to detect and monitor a wide range of human and animal viruses, identify specific viral mutations, and provide early warnings about potential public health threats to enhance timely intervention and resource allocation

**Role: Co-Investigator**

**Texas Department of State Health Services: HHS001089700001/292484 (Boerwinkle)**

**Texas SARS-CoV-2 Variant Network**

08/01/2021 – 07/31/2026

This project aims to develop and implement a statewide network of laboratories collecting and sequencing SARS-CoV-2 positive samples and reporting them to the Texas Department of State Health Services (DSHS).

**Role: Co-Investigator, Co-Director Coordinating Center**

**NIH/NIAID: R01-AI148342 (Arias)**

**VENOUS: A Translational Approach to Enterococcal Bacteremia**

07/01/20-06/30/25

This project hypothesizes a deep understanding of the clinical and microbiological aspects of VRE bloodstream infections and dynamics of the population structure of infecting isolates is crucial to help design novel diagnostic approaches and treatment regimens to improve the outcomes of these difficult-to-treat infections. Using an assembled cohort, we proposed to i) characterize the clinical impact of VRE bacteremia, ii) dissect the population structure of VRE causing bloodstream infections and, iii) develop a new minimal inhibitory concentration (MIC)-independent diagnostic test to assess DAP susceptibility, seeking to guide clinicians with a novel tool to allow accurate identification of DAP-susceptible isolates and improve the use of DAP and combination with β-lactams against these organisms.

**Role: Subcontract Principal Investigator**

**NIH/NIAID: P01- AI152999-01 (Arias)**

**Dynamics of colonization and infection by multidrug-resistant pathogens in immunocompromised and critically ill patients (DYNAMITE)**

07/01/20-06/31/26

This project hypothesizes that patient susceptibility to gut-derived nosocomial colonization and subsequent infection is critically dependent on functional microbiota-pathogen interactions that can be detected via a holistic combination of pathogen, host, and commensal microbiota analyses.

**Role: Subcontract Principal Investigator, Co-Core Director**

**B. PENDING RESEARCH SUPPORT**

**NIH/NIAID (Hanson)**

**A Clinical Application for Pathogen Genomic Insights in Healthcare Excellence: Enhancing GENO-STELLAR for Hospital-wide surveillance**

04/01/2025 – 03/31/2030

This project aims to expand GENO-STELLAR's support to additional pathogens and enhance its current features and functionality. By implementing antimicrobial susceptibility predictive models and improving integration with existing systems, we seek to create a rapid, accessible, and clinically actionable tool. Ultimately, we plan to conduct a clinical trial to evaluate GENO-STELLAR's effectiveness in reducing the time from empiric treatment to definitive therapy.

**Role: Principal Investigator**

**NIH/NIAID (McNeil)**

Blake M. Hanson, M.S., Ph.D.

**The Relationship between Antibiotic Susceptibility, Microbial Genomics and Clinical Outcome in Pediatric *Staphylococcus aureus* Infections**

04/01/2025 – 03/31/2030

This project aims to generate a multicenter prospective cohort study assessing the influence of different antimicrobial susceptibility and microbial genetic backgrounds on patient outcomes of invasive methicillin-susceptible *Staphylococcus aureus* infections.

**Role:** Subcontract Principal Investigator

**NIH/DHHS (Wu)**

**Forecasting viral transmission dynamics and evolution at the community level**

12/01/2024 - 11/30/2029

This proposal aims to utilize the SARS-CoV-2 Omicron variant as a model for exploring and quantifying/predicting the emergence and establishment of a novel SARS-CoV-2 variant. This proposal will also observe novel emergence of new variants as well as assess the impact of these models to other pathogen trends such as norovirus.

**Role:** Co-Investigator

**C. COMPLETED RESEARCH SUPPORT**

**NIH/NIAID: K01AI148593-01 (Hanson)**

**Mobile Genetic Elements and Clinical Outcomes in *Staphylococcus aureus* Bacteremia**

07/01/20 – 06/30/24

This project aims to clarify the role mobile genetic elements play in transmitting antimicrobial resistance and driving genetic diversity within endemic *Staphylococcus aureus* lineages causing bacteremia. Additionally, this research will provide the platform for the career development of a young scientist focused on bacterial phylogenetics, clinical epidemiology, and machine learning methods to tackle the dire consequences of AMR

**Role:** Principal Investigator

**NIH/NIAID: UM1AI104681 (Hanson as a sub-project of the Antimicrobial Resistance Leadership Group) GENOmics, Sequencing-based Typing, Epidemiology, Linkage, and Antimicrobial Resistance (GENOSTELLAR) Tool**

7/2020-6/2024

This is an independent sub-project of the Antimicrobial Resistance Leadership Group funding. This project aims to develop an online tool for clinicians and scientists to integrate bacterial whole genome sequencing data with centralized phenotypic susceptibility results and clinical outcomes to generate genomic insights into antimicrobial resistance mechanism in clinical isolates that could aid in the management of patients affected with multi-drug resistant bacteria.

**Role:** Principal Investigator

**UTHealth CCTS COVID-19 Pilot Funding (Hanson)**

**COVID-19 and SARS-CoV-2 infection surveillance of high-risk community members in the Houston area**

07/01/2020 – 06/30/2021

This project aims to collect nasal samples and blood to assess for active SARS-CoV-2 infection and evidence of previous infection, respectively. Focusing in on high-risk communities in the Houston area, we are assessing the impact of the COVID-19 pandemic on our local communities.

**Role:** Principal Investigator

**School of Public Health PRIME Award, The University of Texas Health Science Center (Hanson)**

**Elucidating the antimicrobial resistance mechanisms and mapping the mobilome of *Staphylococcus aureus***

09/01/2018 – 08/31/2019

Blake M. Hanson, M.S., Ph.D.

This project aims to characterize the diversity of mobile genetic elements present within *S. aureus* causing bacteremia in a single hospital in Latin America.

**Role:** Principal Investigator

## CONFERENCE ABSTRACTS AND PUBLICATIONS

### A. SELECTED INVITED CONFERENCE TALKS AND PANELS

1. **Hanson BM**, Treangen T, Ajami N, Harrison L. Career mentoring: careers in computational genomics applied to infectious diseases. Texas Medical Center Antimicrobial Resistance and Stewardship Conference 2023. Houston, TX (2023). Invited Panel Speaker
- 2.
3. **Hanson BM**, Tran TT, Miller W. Career mentoring: career development awards in antimicrobial resistance. Texas Medical Center Antimicrobial Resistance and Stewardship Conference 2021. Houston, TX (2021). Invited Panel Speaker
4. **Hanson BM**. Nanopore sequencing to understand AMR. Oxford Nanopore Day Houston. Houston, TX (2019). Invited Talk
5. **Hanson BM**. The application of genomic sequencing in pathogen identification and antimicrobial susceptibility detection. Texas Department of State Health Services Healthcare Safety Conference 2018. Austin, TX (2018). Invited Keynote Talk
6. **Hanson BM**, Weinstock G, Sodergren E, Leopold S, Nguyen H, Brown JS, van Belkum A, Dunne W, Schechter L, Creely D, Humphries R, Yang S, Hemarajata P, Miller W, Kahn A, Arias C. Determining novel antimicrobial resistance mechanisms with Oxford Nanopore long-read sequencing data. Oxford Nanopore London Calling 2018. London, UK (2018). Invited Talk

### B. SELECTED PEER REVIEWED CONFERENCE ABSTRACTS

7. Troisi C, Bauer C, **Hanson BM**, Deegan J. From sewage to solutions: The possibilities and implications of wastewater monitoring. American Public Health Association 2024. Minneapolis, MN (2024).
8. **Hanson BM**, Walker JN, Hunter T, Simar SR, Duran Ramirez J, Myckatyn T. A prospective randomized clinical trial demonstrates prophylactic antibiotic administration does not reduce microbial burden or alter the breast microbiome during implant-based breast reconstruction. ISSSI 2024. Perth, Australia (2024).
9. **Hanson BM**, Gillani S, Al Rustem H, Bobier K, Hardy M, Freeland M, Rai S, Kubin G, Milleron R, Boerwinkle E. Evolving with Evolution: Utilizing Dynamic Dashboards to Track Sars-Cov-2 Variants in Texas in a Time of Expanding Genomic Diversity. CSTE 2024. Pittsburgh, PA (2024).
10. Troisi C, Bauer C, **Hanson BM**, Deegan J. From sewage to solutions: The possibilities and implications of wastewater monitoring. Preparedness Summit 2024. Cleveland, OH (2024).
11. **Hanson BM**, Gillani S, Al Rustem H, Bobier K, Hardy M, Freeland M, Rai S, Kubin G, Milleron R, Boerwinkle E. SARS-CoV-2 genomic surveillance system development as a model for other emerging pathogens in Texas. IDWeek 2023. Boston, MA (2023).

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12. **Hanson BM**. Genomics of Carbapenem-resistant *Enterobacteriales* in the USA. Texas Medical Center Antimicrobial Resistance and Stewardship Conference 2021. Houston, TX (2021).
13. **Hanson BM**. Nanopore Sequencing to Understand AMR. ASM Conference on Rapid Applied Microbial Next-Generation Sequencing and Bioinformatic Pipelines 2018. Tysons Corner, VA (2018).
14. **Hanson BM**. Nanopore Sequencing to Understand AMR. Texas Medical Center Antimicrobial Resistance and Stewardship Conference 2018. Houston, TX (2018).
15. **Hanson BM**, Weinstock G, Shannon W, Kwambana B, Deych E, Betts M, Worwui A, Carter S, Brown JS, Ceesay F, Tientcheu PE, Phillips D, Sambou E, Sodergren E, Antonio M. Early Life Colonization of the Nasopharyngeal Microbiome: A Longitudinal Study in The Gambia. *The Human Microbiome: Emerging Themes at the Horizon of the 21<sup>st</sup> Century*. Bethesda, Maryland (2017).
16. **Hanson BM**, Leopold S, Rojas L, Nguyen H, Chen L, Lek S, Spakowicz D, Johnson J, Brown J, Bonomo R, Arias C, Weinstock G. Applications of long-read sequencing in infectious disease genomics. *Infectious Disease Genomics* 2015. Cambridge, London (2015).
17. **Hanson BM**, Kwambana B, Zhou Y, Worwui A, Sodergren E, Ndukum-Ndonwi MB, Kinteh MA, Clark E, Bah S, Bah E, Sanneh K, Kijera M, Gibba I, Bojang L, Haffner M, Cox M, Faal A, Sillah L, Said B, Ezedimbu-Michael P, Tientcheu O, Antonio M, Weinstock G. Pneumococcal vaccination and the nasopharyngeal microbiome: a longitudinal study in The Gambia. *ASM General Meeting*. New Orleans, Louisiana (2015).
18. **Hanson BM**, Kwambana B, Zhou Y, Worwui A, Sodergren E, Ndukum-Ndonwi MB, Kinteh MA, Clark E, Bah S, Bah E, Sanneh K, Kijera M, Gibba I, Bojang L, Haffner M, Cox M, Faal A, Sillah L, Said B, Ezedimbu-Michael P, Tientcheu O, Antonio M, Weinstock G. Pneumococcal vaccination and the nasopharyngeal microbiome: a longitudinal study in The Gambia. *5<sup>th</sup> International Human Microbiome Congress*. Luxembourg City, Luxembourg (2015).
19. **Hanson BM**, Kates A, O'Malley S, Farina S, Klostermann C, Wu J, Meirick T, Quick M, Forshey B, Smith T. Longitudinal study of Iowa families to determine prevalence of *Staphylococcus aureus* and observe colonization incidence and familial transmission. *IDWeek* 2013. San Francisco, California (2013).
20. **Hanson BM**, Kates A, O'Malley S, Farina S, Klostermann C, Wu J, Meirick T, Quick M, Forshey B, Smith T. Longitudinal study of Iowa families to determine prevalence of *Staphylococcus aureus* and observe colonization incidence and familial transmission. *Great Plains Emerging Infectious Diseases Conference*. Iowa City, Iowa (2013).
21. **Hanson BM**, Kates A, O'Malley S, Farina S, Klostermann C, Wu J, Meirick T, Quick M, Forshey B, Smith T. Longitudinal study of Iowa families to determine prevalence of *Staphylococcus aureus* and observe colonization incidence and familial transmission. *University of Iowa Health Sciences Research Week*. Iowa City, Iowa (2013).
22. Beahm A, **Hanson BM**, Karriker L, Ramirez A, Smith TC, Frana T. Molecular characterization of *Staphylococcus aureus* and methicillin-resistant *Staphylococcus aureus* isolated from veterinary students, swine, and their environment. *Great Plains Emerging Infectious Diseases Conference*. Iowa City, Iowa (2012).

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23. O'Brien AM, **Hanson BM**, Farina SA, Wu JY, Simmering JE, Wardyn SE, Kulick ME, Wallinga DB, Smith TC. *Staphylococcus aureus* and methicillin-resistant *Staphylococcus aureus* (MRSA) in retail pork. 2<sup>nd</sup> ASM-ESCMID Conference on methicillin-resistant *Staphylococci* in Animals. Boston, Maryland (2011).
24. **Hanson BM**, Harper, AL, Wardyn SE, Dressler AE, Smith TC. Characterizing *Staphylococcus aureus* strains presenting at a Mongolian Hospital. University of Iowa Epidemiology Poster Session. Iowa City, Iowa (2010).
25. **Hanson BM**, Harper, AL, Scheibel, RP, Roberts LK, Smith TC. Prevalence of *Staphylococcus aureus* and methicillin-resistant *Staphylococcus aureus* in retail meat in Iowa. Iowa Academy of Science Poster Session. Des Moines, Iowa (2009).
26. **Hanson BM**, Harper, AL, Scheibel, RP, Roberts LK, Smith TC. Prevalence of *Staphylococcus aureus* and methicillin-resistant *Staphylococcus aureus* in retail meat in Iowa. Midwest Rural Agricultural Safety & Health Forum. Johnston, Iowa (2009).
27. **Hanson, BM**, Harper, AL, Scheibel, RP, Roberts LK, Smith TC. Prevalence of *Staphylococcus aureus* and methicillin-resistant *Staphylococcus aureus* in retail meat in Iowa. ASM-ESCMID Conference on methicillin-resistant *Staphylococci* in Animals Poster Session. London, England (2009).
28. **Hanson, BM**, Harper, AL, Scheibel, RP, Roberts LK, Smith TC. Prevalence of *Staphylococcus aureus* and methicillin-resistant *Staphylococcus aureus* in retail meat in Iowa. University of Iowa Research Week Poster Session. Iowa City, Iowa (2009).

**C. PREPRINT PUBLICATIONS** (\*Corresponding author)

**D. PEER REVIEWED PUBLICATIONS** (\*Corresponding author)

1. **Hanson BM**<sup>^</sup>, Tisza MJ<sup>^</sup>, Clark JR, Wang L, Payne K, Ross MC, Mena KD, Gitter A, Javornik Cregeen SJ, Cormier J, Avadhanula V, Terwilliger A, Balliew J, Wu F, Rios J, Deegan J, Piedra PA, Petrosino JF, Boerwinkle E, Maresso AW. Sequencing-Based Detection of Avian Influenza A(H5N1) Virus in Wastewater in Ten Cities. *N Engl J Med.* 2024 Sep 26;391(12):1157-1159. doi: 10.1056/NEJMc2405937. Epub 2024 Sep 11. PMID: 39259887.  
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2. Chen X, Balliew J, Bauer CX, Deegan J, Gitter A, **Hanson BM**, Maresso AW, Tisza MJ, Troisi CL, Rios J, Mena KD, Boerwinkle E, Wu F. Revealing patterns of SARS-CoV-2 variant emergence and evolution using RBD amplicon sequencing of wastewater. *J Infect.* 2024 Sep 26;89(5):106284. doi: 10.1016/j.jinf.2024.106284. PMID: 39341403.
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## E. CONTINUING MEDICAL EDUCATION

1. Myckatyn TM, Duran Ramirez JM, Walker JN, **Hanson BM**. Management of biofilm with breast implant surgery. *Plast Reconstr Surg*. 2023 Nov 1;152(5):919e-942e. doi: 10.1097/PRS.00000000000010791. Epub 2023 Oct 23. PMID: 37871028

## F. CONFERENCE PROCEEDINGS (\*Corresponding author)

1. Hourcade JP, Chrischilles EA, Gryzlak BM, **Hanson BM**, Dunbar DE, Eichmann DA, Lorentzen RR. Design lessons for older adult personal health records software from older adults. Proceedings of 6<sup>th</sup> International Conference on Universal Access in Human-Computer Interaction, held as part of HCI International. Lecture Notes in Computer Science, 6766, 176-85.

## G. REVIEWS AND EDITORIALS (\*corresponding author)

1. Darkoh C, DuPont HL, Arias CA, Oyaro M, Brown EL, **Hanson BM**. Reply to Lutgring et al. *Clin Infect Dis*. 2022 Jul 12;. doi: 10.1093/cid/ciac376. PMID: 35818862.
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3. Morales-Marroquin E, **Hanson B**, Greathouse L, de la Cruz-Munoz N, Messiah SE. Comparison of methodological approaches to human gut microbiota changes in response to metabolic and bariatric surgery: A systematic review. *Obes Rev*. 2020 Apr 6. PMID: 32249534
4. Walker JN, **Hanson BM**, Myckatyn TM. Commentary on: optimizing breast pocket irrigation: the breast implant-associated anaplastic large cell lymphoma (BIA-ALCL) era. *Aesthet Surg J*. 2020 May 16;40(6):626-629. PMID: 31740924
5. Galloway-Peña J, **Hanson B**. Tools for analysis of the microbiome. *Dig Dis Sci*. 2020 Mar;65(3). PMID: 32002757
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## H. BOOKS AND BOOK CHAPTERS

1. Thapalia D, **Hanson BM**, Kates AE, Klostermann CA, Nair R, Wardyn SE, Smith TC. Zoonoses – Infections Affecting Humans and Animals. 1<sup>st</sup> ed. Zoonotic Diseases of Swine: Food-borne and Occupational Aspects of Infection. 2015: Springer Netherlands.

## RELEVANT MEDIA EXPOSURE

### A. PRINT AND WEBSITES

1. Date: September 19, 2024  
Publication: Medpage Today  
Website: <https://www.medpagetoday.com/infectiousdisease/surveillance/111909>
2. Date: July 16, 2024  
Publication: ARLG Quarterly Newsletter Q3 2024  
Website: <https://arlq.org/geno-stellar-an-interview-with-blake-hanson-phd/>
3. Date: May 23, 2024  
Publication: Bloomberg  
Website: <https://www.bloomberg.com/news/articles/2024-05-23/bird-flu-outbreak-in-dairy-cows-is-more-widespread-sewage-tests-suggest>
4. Date: May 22, 2024  
Publication: CNN  
Website: <https://www.cnn.com/2024/05/22/health/bird-flu-wastewater-monitoring-texas/index.html>
5. Date: May 21, 2024  
Publication: Clinical & Engineering News  
Website: <https://cen.acs.org/biological-chemistry/US-CDC-begins-tracking-influenza/102/web/2024/05>
6. Date: May 16, 2024  
Publication: MedPage Today  
Website: <https://www.medpagetoday.com/special-reports/exclusives/110169>
7. Date: May 16, 2024

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Publication: LA Times

Website: <https://www.latimes.com/environment/story/2024-05-16/texas-researchers-identify-bird-flu-in-wastewater-of-nine-cities-filling-in-cdc-data-gaps>

8. Date: April 15, 2021

Publication: Texas Monthly – A DNA zoo maps the mysteries of all creatures great and small

Website: <https://www.texasmonthly.com/news-politics/dna-zoo-baylor-college-medicine-houston/>

9. Date: May 2, 2020

Publication: ASM-TMC COVID-19 101 – Debunking Myths

Website: <https://www.youtube.com/watch?v=RUKucmvWoqY&list=LLf-MUHEm4PDyCcodDvGaaqw&index=2&t=1898s>

10. Date: May 24, 2018

Publication: Oxford Nanopore News

Website: <https://nanoporetech.com/about-us/news/london-calling-day-1-morning-writeups>

Website: <https://nanoporetech.com/resource-centre/talk/determining-novel-antimicrobial-resistance-mechanisms-oxford-nanopore-long>

11. Date: February 11, 2016

Publication: YouTube – Oxford Nanopore Technologies

Website: <https://www.youtube.com/watch?v=AWEIbgV9uAg>

12. Date: January 14, 2016

Publication: we-ha.com

Website: <http://we-ha.com/jackson-laboratories-will-kick-off-discussion-series-on-family-health/>

13. Date: January 7, 2016

Publication: Hartford Courant

Website: <http://www.courant.com/community/west-hartford/hc-west-hartford-childrens-museum-partnership-0108-20160107-story.html>

14. Date: January 4, 2016

Publication: West Hartford Patch

Website: <http://patch.com/connecticut/westhartford/jackson-laboratory-lead-public-discussion-human-microbiome-family-health>

## B. TELEVISION

1. Date: June 14, 2024

Publication: Fox 26 Morning News

Topic: Interpretation and implications of H5N1 detection in wastewater in Texas