

HADLEY STEVENS SMITH, PhD, MPSA

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she, her, hers
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EDUCATION

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| Baylor College of Medicine Center for Medical Ethics and Health Policy Postdoctoral Fellow, Health Policy Mentor: Amy McGuire, JD, PhD | 2019-Present |
| The University of Texas Health Science Center at Houston School of Public Health PhD in Public Health, Health Economics track Minor fields: Epidemiology and Biostatistics Dissertation: "Clinical Diagnostic Whole Exome Sequencing for Infants in Intensive Care Settings: Outcomes Analysis and Economic Evaluation" Chair: J. Michael Swint, PhD | 2015-2019 |
| Texas A&M University The Bush School of Government and Public Service MPSA, Master of Public Service and Administration Concentration: Health Policy Analysis Dean's Certificate in Leadership, Public Service Leadership Program | 2013-2015 |
| BS, Political Science (3+2 program with MPSA) Minor: Genetics Summa Cum Laude | 2010-2014 |

CONTINUING EDUCATION

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| Decision Modelling for Health Economic Evaluation – London School of Hygiene and Tropical Medicine and University of York | 2021 |
| Measurement, Design, and Analysis Methods for Health Outcomes Research – Harvard T.H. Chan School of Public Health, Executive and Continuing Professional Education | 2019 |
| Genetics and Genomics – Stanford University, Center for Professional Development | 2018 |

RESEARCH INTERESTS

I am a health economist working at the intersection of health economics and the ethical, legal, and social implications (ELSI) of genomics. My research examines precision medicine applications from patient-centered, clinical, and economic perspectives. I am interested in evaluation of the implementation and delivery of clinical genomics, specifically for newborn and pediatric patients, and assessment of family-level impacts of genomic sequencing. I aim to inform policies that advance equitable and efficient implementation of genomic medicine to improve population health.

Fields: Health economics and outcomes research, policy analysis, empirical bioethics

Methods: Applied econometrics; cost-effectiveness analysis; preference elicitation; psychometrics; qualitative methods

Issues: Clinical integration of precision medicine; translational research; patient-centered outcomes measurement; valuation of health and non-health outcomes; spillover effects; access to care; health equity; child health and well-being

Software: Stata, REDCap, Epic electronic medical record

ACADEMIC APPOINTMENTS

The University of Texas Health Science Center at Houston

August 2021 –

School of Public Health, Department of Management, Policy and Community Health

Adjunct Instructor

RESEARCH AND TEACHING ASSISTANTSHIPS

Baylor College of Medicine

Sept 2016 – May 2019

Department of Molecular and Human Genetics *Brendan Lee Lab*
& Center for Medical Ethics and Health Policy *Mentor: Amy McGuire*
Research Associate

The University of Texas Health Science Center

Sept 2015 – Aug 2016

Department of Management, Policy and Community Health *Frances Lee Revere*
Graduate Assistant for Research/Teaching Assistant

Texas A&M Health Science Center

Summer 2014

Center for Innovation in Advanced Development and Manufacturing
Policy Intern, Public Health Preparedness and Response

Texas A&M University

Nov 2013 – May 2015

Department of Political Science *Erica Owen Palmer*
Graduate Assistant for Research, Political Economy

Texas A&M University

Fall 2013 – Spring 2014

Department of Political Science *Erica Owen Palmer*
Teaching Assistant, Polimetrics

Texas A&M University

Sept 2012 – May 2013

Project for Equity, Representation, and Governance, Department of Political Science
Research Technician

Personal Genetics Education Project (pgEd)

Summer 2013

Harvard University Medical School *Wu Lab*
Research Intern

MANUSCRIPTS IN PROGRESS

1. **Smith HS**, Pereira SP, Morain SR. Promoting justice in genomics: Stakeholder perspectives and feasibility of a genomics-enabled learning health system
2. **Smith HS** et al. Validating the PedsQL measure of pediatric quality of life in the context of clinical genome sequencing: Results from the CSER Consortium
3. **Smith HS** et al. Timeline and clinical context of exome sequencing for infants in intensive care
4. **Smith HS**, Sanchez C, Maag R, et al. Patient and clinician perceptions of precision cardiology care: Findings from the HeartCare Study
5. **Smith HS** et al. Impact of exposure to genomic sequencing on attitudes and perceived importance among parents and physicians in the BabySeq Project
6. Uveges MK, **Smith HS**, Pereira S, McGuire AL, Beggs AH, Green RC, Holm IA. Sharing of infants' genetic risk information among family members in the BabySeq Project
7. Rubanovich CK., Rajagopalan RM, **Smith HS**, Cheema JS, Kwaning A, Myers MI, Honcharov V, Bloss CS. A scoping review of the literature on attitudes, interest, and impacts of genetic ancestry testing in the United States

PEER-REVIEWED PUBLICATIONS

Includes manuscripts under review; * denotes equal authorship contribution

1. Murali CN, Lalani SR, Azamian M, Miyake C, **Smith HS**. Quality of Life, illness perceptions, and parental lived experiences in *TANGO2*-related metabolic encephalopathy and arrhythmias: Insights from family engagement (submitted)
2. Halley MC, **Smith HS**, Ashley EA, Goldenberg AJ, Tabor HK. Reframing translational research in rare diseases in the United States: A mosaic approach. (under review)
3. **Smith HS**, Morain SR, Robinson JO, et al. Perceived utility of genomic sequencing: Qualitative analysis and synthesis of a conceptual model to inform instrument development. (under review)
4. Gutierrez AM, Robinson JO, Outram SM, **Smith HS**, et al. Examining access to care in clinical genomic research and medicine: Experiences from the CSER consortium. (R&R)
5. **Smith, HS**, Brothers KB, Knight SJ, et al. Conceptualization of utility in translational clinical genomics research. Forthcoming in *American Journal of Human Genetics*.
6. Phillips KA, Trosman JR, Douglas MP...**Smith HS**. US private payers' perspectives on insurance coverage for genome sequencing versus exome sequencing: A study by the Clinical Sequencing Evidence-generating Research Consortium (CSER). Forthcoming in *Genetics in Medicine*.
7. Pereira S,* **Smith HS**,* Frankel L, et al. Psychosocial effect of newborn genomic sequencing on families in the BabySeq Project: A randomized clinical trial. *JAMA Pediatrics*. 2021. Published online 23 August 2021.
8. Fowler LR, Schoen L, **Smith HS**, Morain SR. Sex education on TikTok: A content analysis of themes. *Health Promotion Practice*. 2021. Published online 21 August 2021.
9. Murdock D, Venner E, Munzy DM,...**Smith HS**... et al. Genetic testing in ambulatory cardiology clinics reveals high rate of findings with clinical management implications. *Genetics in Medicine*. 2021. Published online 06 August 2021.
10. **Smith HS**. How Should Economic Value Be Considered in Treatment Decisions for Individual Patients? *AMA Journal of Ethics*. 2021;23(8):E607-612.
11. Makhnoon S, **Smith HS**, Bednar EM, et al. Disclosure of familial implications of pathogenic variants in breast-cancer genes to patients: Opportunity for prompting family communication. *J Community Genet*. 2021;12(3):439-447. PMID: 33481188.
12. **Smith HS**, Franciskovich R, Lewis AM, Gerard A, Littlejohn RJ, Nugent K, Rodriguez J, Streff H. Outcomes of prior authorization requests for genetic testing in outpatient pediatric genetics clinics. *Genetics in Medicine*. 2021;23(5):950-955.
13. Guzauskas GF, Garbett S, Zhou Z, Spencer SJ, **Smith HS**, Hao J, Hassen D, Snyder SR, Graves JA, Peterson JF, Williams MS, Veenstra DL. Cost-effectiveness of population-wide genomic screening for hereditary breast and ovarian cancer in the United States. *JAMA Network Open*. 2020;3(10):e2022874-e2022874. PMID: 33119106.
14. Lázaro-Muñoz G, Torgerson L, **Smith HS**, Pereira S. Perceptions of best practices for return of results in an international survey of psychiatric genetics researchers. *European Journal of Human Genetics*. 2021;29(2):231-240. PMID: 33011736.

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15. **Smith HS**, Swint JM, Lalani SR, Russell, HV, Lee BH. Exome sequencing compared with standard genetic tests for critically ill infants with suspected genetic conditions. *Genetics in Medicine*. 2020;22(8):1303-1310. PMID: 32336750
 16. Goddard KA, Angelo FA, Ackerman SL,...**Smith HS**, Waltz M, Lee SS-J. Lessons learned about harmonizing survey measures for the CSER consortium. *Journal of Clinical and Translational Science*. 2020;4(6):537-546. PMID: 33948230
 17. **Smith HS**, Russell HV, Lee BH, Morain SR. Using the Delphi method to identify clinicians' perceived importance of pediatric exome sequencing results. *Genetics in Medicine*. 2020;22(1):69-76. PMID: 31273346
 18. **Smith HS**, Swint JM, Lalani SR, et al. Clinical application of genome and exome sequencing as a diagnostic tool for pediatric patients: A scoping review of the literature. *Genetics in Medicine*. 2019;21(1):3-16. PMID: 29760485
 19. Meng L, Pammi M, Saronwala, A,...**Smith HS**...Lalani, SR. Use of exome sequencing for infants in intensive care units: Ascertainment of severe single-gene disorders and effect on medical management. *JAMA Pediatrics*. 2017;171(12):e173438. PMID: 28973083

EDITORIALS & COMMENTARIES

1. **Smith HS**, Morain SR. Appropriate Care for Adolescent Eating Disorders in Isolating and Disruptive Times. *J Cognitive Psychotherapy*. 2021;35(3):1-3.
2. **Smith HS**, McGuire AL, Wittenberg E, Lavelle TA. Family-level impact of genetic testing: Integrating health economics and ethical, legal, and social implications. *Personalized Medicine*. 2021;18(3):209-212.
3. Rubanovich CK, **Smith HS**, Bloss CS. Commentary on the development of the Clinician-reported Genetic testing Utility InDEx (C-GUIDE). *Genetics in Medicine*. 2020;22(3):665-666.

FUNDED PROJECTS

K99HG011491 (PI: Smith)

12/2020 – 12/2022

An ELSI-Integrated Evaluation of the Family-Level Utility of Pediatric Genomic Sequencing

The goal of the proposed research is to investigate how clinical genomic sequencing impacts families of pediatric patients. This research will develop an empirically informed framework of normative values important to families of pediatric patients, including ethical, legal, and social implications (ELSI), which will then be used to elicit preferences for features of sequencing from a nationally representative sample of parents in the US. The results of this work will lead to a measure of family utility that integrates ELSI and can be used in a decision analytic model to assess the impact of genomic sequencing and the importance of including family-level ELSI considerations in such analyses.

Role: PI

Baylor College of Medicine and Houston Methodist Collaborative Pilot Grants in Alzheimer's Disease and Related Dementias, (PI: McGuire)

08/2020 – 07/2021

Integrating Precision Medicine into Alzheimer's Disease Care

This project seeks to assess the psychosocial impact of precision neurological care for Alzheimer's disease, including biomarker results, to neurology clinic patients.

Role: Postdoctoral Associate

Precision Medicine and Population Health Grant, BCM (PI: Pereira)

07/2019 – 06/2020

Clinical Impact and Perceived Utility of Genomic Cardiovascular Disease Risk Information in a Precision Medicine for Population Health Initiative

This project seeks to assess the clinical impact and perceived utility of genomic cardiovascular disease risk information in a genomics-enabled learning health system project.

Role: Postdoctoral Associate

3U01 HG006485-06S1 (PIs: Plon, Parsons, McGuire)

09/11/2018 – 05/31/2021

Evaluating Utility and Improving Implementation of Genomic Sequencing for Pediatric Cancer Patients in the Diverse Population and Healthcare Settings of Texas – Administrative Supplement

The goal of this supplement project is to develop and validate a novel measure of perceived utility in genomic medicine by exploring perceptions of utility of clinical genome sequencing across individuals participating in Clinical Sequencing Evidence-Generating Research consortium projects.

Role: Research Associate

3U01 HG007292-06S1 (PIs: Goddard, Wilfond)

09/13/2018 – 05/31/2019

Exome Sequencing in Diverse Populations in Colorado and Oregon – Administrative Supplement

This supplement project will validate key harmonized measures being administered across the Clinical Sequencing Evidence-Generating Research consortium.

Role: Research Associate

U19 HD077671 (PIs: Green, Beggs)

9/5/2013 – 9/30/2019

Genome sequence-based screening for childhood risk and newborn illness

The major goals of this study are to explore the implications, challenges, and opportunities associated with the use of genomic sequence information in the newborn period by generating high-quality exome sequencing data and returning the results of this data to research participants.

Role: Research Associate

INVITED TALKS AND CONFERENCE PRESENTATIONS

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| Promoting justice in genomics: Stakeholder perspectives and feasibility of a genomics-enabled learning health system | 2021 |
| American Society for Bioethics & Humanities Annual Conference (scheduled for October, virtual) | |
| The intersection of ethics and health economics: Perspectives on the ethical assessment of value in clinical genomics | 2021 |
| American Society for Bioethics & Humanities Annual Conference (scheduled for October, virtual) | |
| The Impact of Newborn Genomic Sequencing on Families: Findings from the BabySeq Project | 2021 |
| Clinical Sequencing Evidence-Generating Research (CSER) Consortium Junior Investigator Presentation at consortium-wide virtual meeting | |
| Genetic Testing and Vulnerable Populations: Race, Ethnicity, and Identify in Neonatal Clinical Care | 2021 |
| Texas Children's Hospital Neonatology Ethics Conference | |
| Eating Disorders in COVID Times: Implications for Primary Care Physicians | 2020 |
| UTHealth and Harris Health System Ambulatory Care Services Grand Rounds | |
| Appropriate Care for Adolescent Eating Disorders in Isolating and Disruptive Times | 2020 |
| Affinity Group presentation at American Society of Bioethics and Humanities Annual Meeting | |
| The Development of a Model and Measure of Patient Perceived Utility of Genomic Sequencing | 2020 |

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| Panel presentation with Morain SR, Rubanovich CK, and Wilfond B ELSI Congress <i>Conference canceled due to COVID-19 pandemic</i> | |
| Health Care Utilization And Cost For Infants Receiving Exome Sequencing Platform Presentation, <i>Recorded for online conference during COVID-19 pandemic</i> American College of Medical Genetics and Genomics Annual Meeting | 2020 |
| Utility of Genomic Testing: A Multidisciplinary Perspective Panel Discussion with McGuire AL, Malek J, Brothers KB American Society for Bioethics & Humanities Annual Conference | 2019 |
| When There is No Comparison: Leveraging the Electronic Medical Record to Compare Genetic Tests for Critically Ill Infants Houston VA Center for Innovations in Quality, Effectiveness and Safety (IQuES) Health Services Research Grand Rounds | 2019 |
| Family Matters: Evaluating Genomic Sequencing at the Family Level Baylor College of Medicine Department of Molecular and Human Genetics Grand Rounds | 2019 |
| Promoting Engagement of Families in Research for Finding Better Treatment for TANGO2-Related Disorder Patient engagement workshop at TANGO2 Family Conference (with Murali C) | 2019 |
| Impact of Sequencing on Outcomes and Costs for Critically Ill Infants Clinical Sequencing Evidence-Generating Research (CSER) Consortium Junior Investigator Presentation at consortium-wide virtual meeting | 2019 |
| Economic Evaluation of Exome Sequencing in Newborns Baylor College of Medicine Genetics Grand Rounds | 2019 |
| Beyond Diagnosis – It’s Not What You Know, It’s What You Do With It HudsonAlpha Genomic Medicine Conference | 2018 |
| The Future of Public Health Invited student on panel featuring the Executive Director of the American Public Health Association | 2017 |
| Knowing Your Genome: Introduction to the Science and Ethical, Legal, and Social Implications Informational, community engagement talk at Bryan-College Station, TX Adult Forum | 2013 |

TEACHING EXPERIENCE

Instructor or Small Group Leader - Institution and Course

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| Baylor College of Medicine | |
| Methods & Analysis in Health Policy (MEETH-415), MS2 and MS3 students Assistant Course Director | Fall 2021 |
| Capstone Research Development Small Group Leader | Summer 2020, Fall 2020 |
| Medical Ethics (MBETH-MAIN), MS1 and health professions students Small Group Leader | Spring 2020, 2021 |

Health Policy Journal Club (MEERM-628) Fall 2021
Teaching faculty

Texas A&M EnMed

Health Economics, Elective course for MS2 students Fall 2020
Instructor

Single Lectures - Institution, Course, and Lecture title(s)

Baylor College of Medicine

Introduction to Health Policy (MEETH-413), MS1 students Summer 2020,
Affordability, Access, and Demand for Health 2021
Cost, Value, and Payment in Health Care

Methods & Analysis in Health Policy (MEETH-415), MS2 and MS3 students Fall 2020
Stakeholder Engagement

Thesis Research Methods I (GCTHE 81001), Genetic Counseling Master's students Spring 2020,
Research Specific Aims 2021
Building on Qualitative Research
Electronic Medical Record Research

Ethics, Professionalism, and Policy Program (EP3), Online Graduate Medical Education 2019
Healthcare Economics Module, Contributor

Texas A&M EnMed

Practice of Medicine, MS2 students Fall 2020, 2021
Health Economics

University of Texas Health Science Center, School of Public Health

Introduction to Health Economics (PHD 3910), PhD and MPH students Spring 2018
Statistical Methods in Health Economics

Methods for Economic Evaluation of Health Care (PHD 3915), PhD and MPH students Fall 2018,
Uncertainty Analysis and Bootstrapping 2019, 2020,
Net Benefit Regression 2021

Doctoral Dissertation Proposal Development course, PhD students Spring 2017,
Measurement Model Development (x 3) 2018, 2019
Preparing for Dissertation Stage Panel (x1)

Texas A&M University

Bush School Health Economics and Policy (PSAA 638), Masters students Spring 2019
Health Care Quality

Genetic Expression in the Context of Modern Medicine (BIMS 289), undergraduate students Fall 2014
Bioethics, Science, and Public Policy

MENTORING

‡Thesis committee chair; *Co-Mentor with faculty member

| Student | Program | Thesis/Capstone Project | Year |
|------------------|---|---|-------------|
| Julie Stefka‡ | Master of Science in Genetic Counseling, BCM School of Health Professions | Cascade testing following exome sequencing | 2021- |
| Sowmya Yennam* | BCM School of Medicine, Health Policy and Ethics Pathways | Cost conversations in ophthalmology | 2020- |
| Brittan Elliott* | BCM School of Medicine, Health Policy Pathway | Policy considerations of newborn genomic screening versus standard newborn screening | 2020- |
| Adasia Ritenour | Master of Science in Genetic Counseling, BCM School of Health Professions | Parental experiences and satisfaction throughout the genetic diagnostic experience: Exploring the impact of clinical setting in the context of COVID-19 | 2019-2021 |
| Hannah Helber | Master of Science in Genetic Counseling, BCM School of Health Professions | Assessment of Understanding and Satisfaction of Exome Sequencing: Video vs Telephone Consent | 2018-2020 |

PROFESSIONAL MEMBERSHIPS

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| International Health Economics Association | 2021 – present |
| American Society for Bioethics and Humanities | 2019 – present |
| Society for Medical Decision Making | 2019 – present |
| American Society of Human Genetics | 2019 – present |
| AcademyHealth | 2016 – 2019 |

WORKING GROUPS

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| Econ-Omics , International Health Economics Association Convener for Research and International Collaboration | 2021 – present |
| Global Economics and Evaluation of Clinical Genomics Sequencing Working Group (GEECS) | 2021 – present |
| Clinical Sequencing Evidence-Generating Research (CSER) Consortium , National Institutes of Health; Clinical Utility, Health Economics and Policy (CUHEP) | 2017 – present |
| Undiagnosed Diseases Network (UDN) , National Institutes of Health; U3 (UDN Utility and Utilization) | 2017 – 2019 |

PROFESSIONAL SERVICE

Institutional Review Board Member, Rice University (August 2020-present)

Baylor College of Medicine Social Determinants of Health, Medicine, and Ethics Journal Club, co-founder and co-leader (June 2020-present)

Baylor College of Medicine Policy Statement Contributor: Strengthening Our Commitment to Racial Justice to Improve Public Health (August 2020)

Baylor College of Medicine Ethics, Policy & Implementation committee member for strategic planning initiative (2018)

Peer reviewer: *Genetics in Medicine, Pediatrics, Social Science & Medicine, Value in Health*

Abstract reviewer: Society for Medical Decision Making (2020)

Social Media Reporter, Society for Medical Decision Making 2020 Annual Conference (#SMDM20)

AWARDS AND HONORS

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| Dean's Award of Excellence, Baylor College of Medicine | 2021 |
| Best Poster, Baylor College of Medicine Health and Science Policy Research Day | 2019 |
| Delta Omega Honorary Society in Public Health | 2019 |
| Finalist, AcademyHealth HSRProj Research Competition for Students | 2018 |
| Dean's Excellence Scholarship, UTHealth | 2016 |
| Outstanding New Doctoral Student Merit Scholarship, UTHealth | 2015 |
| Pi Alpha Alpha (honor society for schools of public affairs and administration) | 2015 |
| The George and Barbara Bush Fellowship, Texas A&M University | 2013 |
| Phi Beta Kappa | 2013 |
| Coca-Cola Scholar, Coca-Cola Scholars Foundation | 2010 |
| Texas A&M Lechner Honors Scholarship | 2010 |

ABSTRACTS AND POSTER PRESENTATIONS

‡Mentored student

1. Ritenour A, ‡ **Smith HS**; Murali CN, Mizerik E, Spector AF, Fernbach SD, Magoulas PL. Parental experiences and satisfaction throughout the genetic diagnostic experience: Exploring the impact of clinical setting in the context of COVID-19. National Society of Genetic Counselors Annual Meeting. 2021.
2. **Smith HS**, Pereira S, Lee B, Veenstra DL, McGuire AL. Family-level impacts of pediatric genomic sequencing: A qualitative analysis and attribute framework. National Human Genome Research Institute Research Training and Career Development Annual Meeting. April 19 2021. Virtual Meeting hosted by the UCLA T32 Program.
3. Uveges MK, **Smith HS**, Pereira S, McGuire AL, Beggs AH, Green RC, Holm IA. Sharing of Infants' Genetic Risk Information Among Family Members in the BabySeq Project. Paper presentation at American Society of Bioethics and Humanities. October 2020. *Conference held online due to COVID-19 pandemic.*
4. **Smith HS**, Rubanovich CK, McGuire, AL. Family Matters: Integrating ELSI Considerations into Evaluations of Family-Level Impact of Genomic Sequencing. ELSI Congress. June 15—17, 2020, New York, NY. *Conference canceled due to COVID-19 pandemic.*
5. Malek J, **Smith HS**, Islam R, Robinson JO, Hsu RL, Canfield I, Raes-Martinez R, Scollon S, Recinos A, Majumder M, Parsons DW, Plon SE, McGuire AL. Great Expectations: Parents' Expected Utility of Genomic Sequencing for Pediatric Cancer Patients in the Texas KidsCanSeq Study. ELSI Congress. June 15—17, 2020, New York, NY. *Conference canceled due to COVID-19 pandemic*
6. Malek J, **Smith HS**, Islam R, Robinson JO, Wang T, Hilsenbeck SG, Hsu RL, Canfield I, Raes-Martinez R, Scollon S, Majumder M, Parsons DW, Plon SE, McGuire AL. Parents' Expected Utility of Genomic Sequencing For

Pediatric Cancer Patients In The Texas KidsCanSeq Study. Platform presentation at American College of Medical Genetics Annual Clinical Genetics Meeting. *Conference held online due to COVID-19 pandemic*

7. Streff H, Franciskovich R, Gerard A, Lewis A, Valdez A, **Smith HS**. Outcomes Of Insurance Pre-authorization Requests For Genomic Testing In An Outpatient Pediatric Genetics Clinic. American College of Medical Genetics Annual Clinical Genetics Meeting. 2020. *Conference held online due to COVID-19 pandemic*
8. Chau J, Bensen JT, Desrosiers LR, Finnila CR, Hindorff LA, Hoban H, Kelly NR, Lynch FL, Rolf B, Smith HS, Wasserstein MP, Yip T, Hassmiller-Lich K. A Time And Motion Study From The Clinical Sequencing Evidence-Generating Research (CSER) Consortium. Poster presentation at American College of Medical Genetics Annual Clinical Genetics Meeting. 2020. *Conference held online due to COVID-19 pandemic*
9. **Smith H**, Pereira S, Robinson J, Frankel L, Christensen K, Parad R, Agrawal P, Yu T, Holm I, Beggs A, Green R, McGuire A. Physician Attitudes, Confidence, and Perceived Utility Following Experience with Newborn Genomic Sequencing in the BabySeq Project. American Society of Human Genetics Annual Conference. October 2019, Houston TX. *Reviewers' Choice Award*
10. Dai H, Murdock DR, **Smith H**, Burrage LC, Rosenfeld JA, Lee B, Undiagnosed Diseases Network. Game of Exomes: Comparing sequencing of commercial laboratories in the Undiagnosed Diseases Network. American Society of Human Genetics Annual Conference. October 2019, Houston TX.
11. Pereira S, **Smith HS**, Frankel L, Robinson JO, Islam R, Christensen KD, Genetti CA, Blout CL, Parad R, Agrawal P, Waisbren SE, Yu T, Holm IA, Beggs AH, Green RC, McGuire AL. BabySeq abstract. The Impact of Newborn Genomic Sequencing on the Parent-Child Relationship in the BabySeq Project. American Society for Bioethics & Humanities Annual Conference. October, 24-27, 2019, Pittsburgh, PA.
12. **Smith HS**, Morain SR. Public Support for Novel Strategies to Reduce Alcohol-Related Motor Vehicle Injuries. Poster presentation at the AcademyHealth 2019 Annual Research Meeting, June 2019, Washington, DC.
13. **Smith HS**, Morain SR. Public Support for Strategies to Reduce Youth Tobacco Use. Poster presentation at the AcademyHealth 2019 Annual Research Meeting, June 2019, Washington, DC.
14. **Smith HS**, Russell HV. Cost-Effectiveness of Exome Sequencing Compared to Standard Diagnostics for Newborns and Infants in Intensive Care. Poster presentation at the Baylor College of Medicine 2019 Health and Science Policy Research Day, April 2, 2019, Houston, TX.
15. **Smith HS**, Canfield I, Robinson JO, Morain SR, Malek J, McGuire AL. Utility of Genomic Testing: A Multi-Disciplinary Perspective. Poster presentation at the Baylor College of Medicine 2019 Health and Science Policy Research Day, April 2, 2019, Houston, TX.
16. Blout C, Christensen K, **Smith H**, Pereira S, Robinson J, Genetti C, Fayer S, Betting W, Schwarts T, Holm I, Beggs A, McGuire A, Green R. Physician Perceived Utility and Preparedness for Newborn Genomic Sequencing: Findings from the BabySeq Project. Platform Presentation at the American College of Medical Genetics Annual Clinical Genetics Meeting, April 2-6, 2019, Seattle, Washington.
17. Murdock DR, **Smith H**, Chen S, Dawson B, Burrage L, Rosenfeld J, Lee B, UDN members, Comparative Quality of Whole Exome Sequencing Among Commercial Laboratories for Patients in the Undiagnosed Diseases Network. Poster presentation at the American Society of Human Genetics Annual Meeting, October 2018, San Diego, California.

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18. **Smith HS**, Russell HV, Magoulas P, Meng L, Bostwick B, Eng C, Kulkarni S, Dominguez-Vidana R, Shaw C, Lalani S, Lee BH. Operational Features from Implementation of Diagnostic Whole Exome Sequencing for Infants in Intensive Care Settings. Poster Presentation at the American College of Medical Genetics and Genomics Annual Meeting, April 12, 2018, Charlotte, NC.

 19. Meng L, Pammi M, Saronwala A, Magoulas P, Ghazi A, Vetrini F, He W, Dharmadhikari A, Qu C, Ge X, Tokita M, Santiago-Sim T, Dai H, **Smith H**, Azamian M, Wangler M, Scott D, Belmont J, Wang X, Leduc M, Xiao R, Liu P, Shaw C, Walkiewicz M, Bi W, Xia F, Lee B, Eng C, Yang Y, Lalani S. Utility of exome sequencing for infants in intensive care units: Ascertainment of severe single-gene disorders and impact on medical management. Presented at the 67th Annual Meeting of The American Society of Human Genetics, October 19, 2017, Orlando, FL.