

Biostatistics Shared Resource

Joseph W Hogan, Director



Overview

The Biostatistics SR offers faculty-level biostatistics and data science collaboration on projects and grant proposals; develops reproducible data pipelines; generates innovative methods to address key analytic and computational challenges; facilitates scientific monitoring of studies and trials; ensures use of rigorous and modern methods for study design, analysis and reporting.

The services ensure that LCC members can have access and utilize rigorous and innovative analytic methods that make optimal use of data associated with investigator-initiated research projects.

Key Services

- Project-specific faculty-level collaboration on new projects and grant proposals
- Bioinformatics and computational biology support for experimental design, data acquisition, preprocessing, visualization, and pipeline development for mapping and preprocessing in -omics platforms
- Ensure quality control and reproducibility of pipelines and data analyses by maintaining archives for data and code
- Consultation and collaboration on analysis of observational data (e.g. EHR) and cohort studies
- Facilitate access to Brown computing resources
- Provide staffing for SRMC

Key Personnel

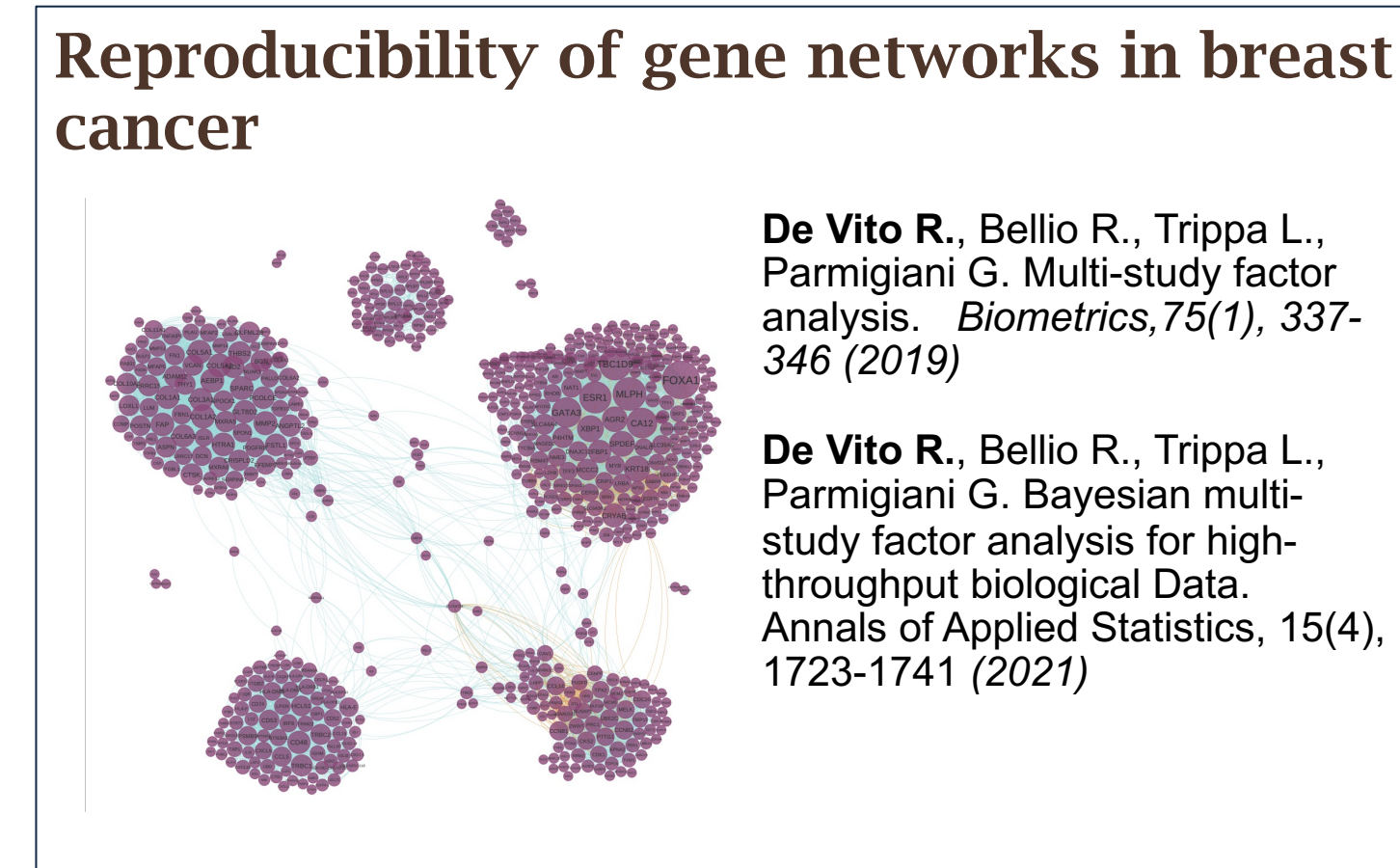
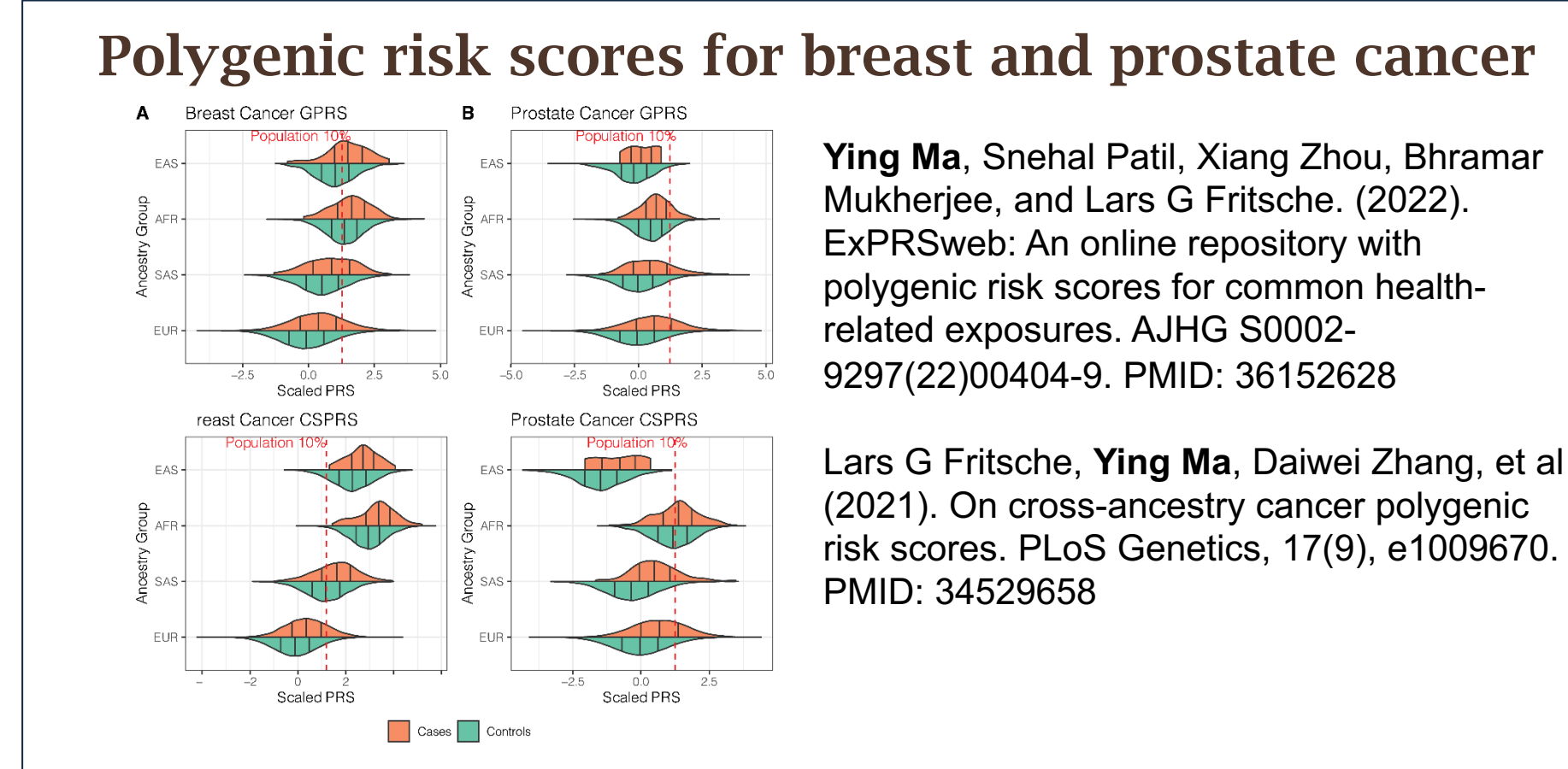
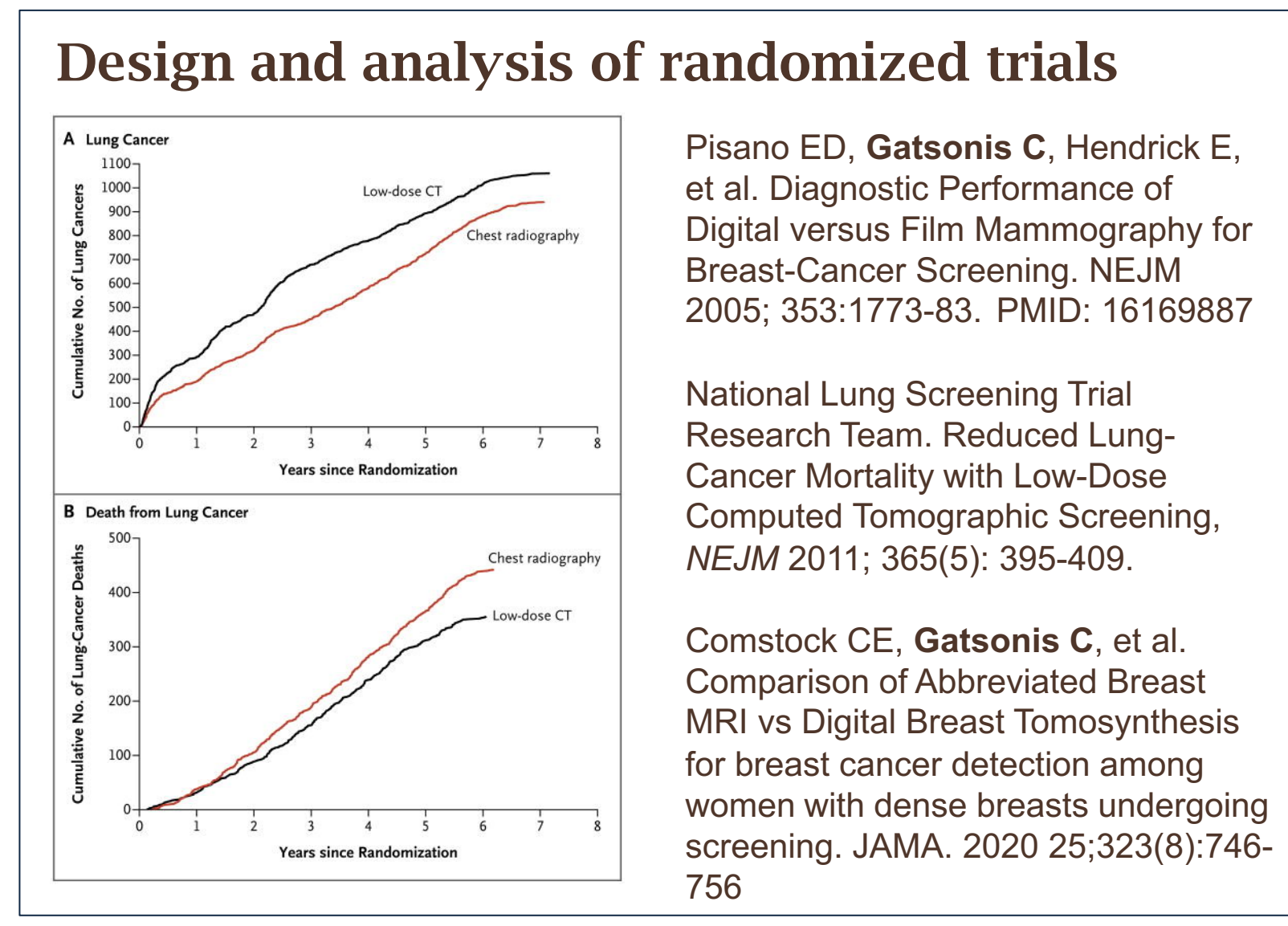
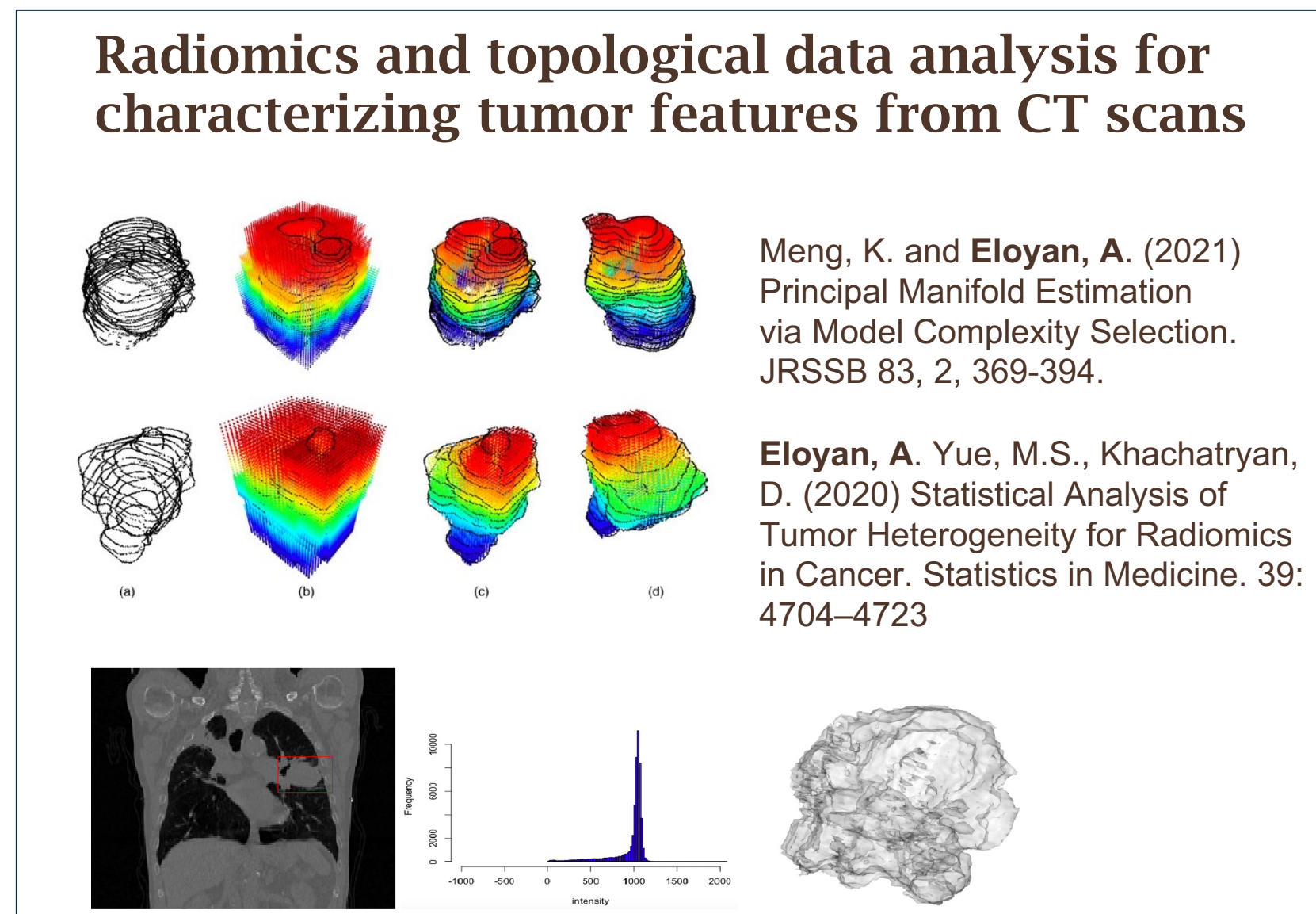
Faculty

- Joseph Hogan (Director)
- Zhijin Wu (Associate Director)
- Consulting faculty in biostatistics and bioinformatics

Technical staff

- Technical Director/Manager - TBN
- Data manager (100%)
- Scientific programmer (100%)
- Software engineer / developer (50%)

Examples of Scientific Impact and Expertise



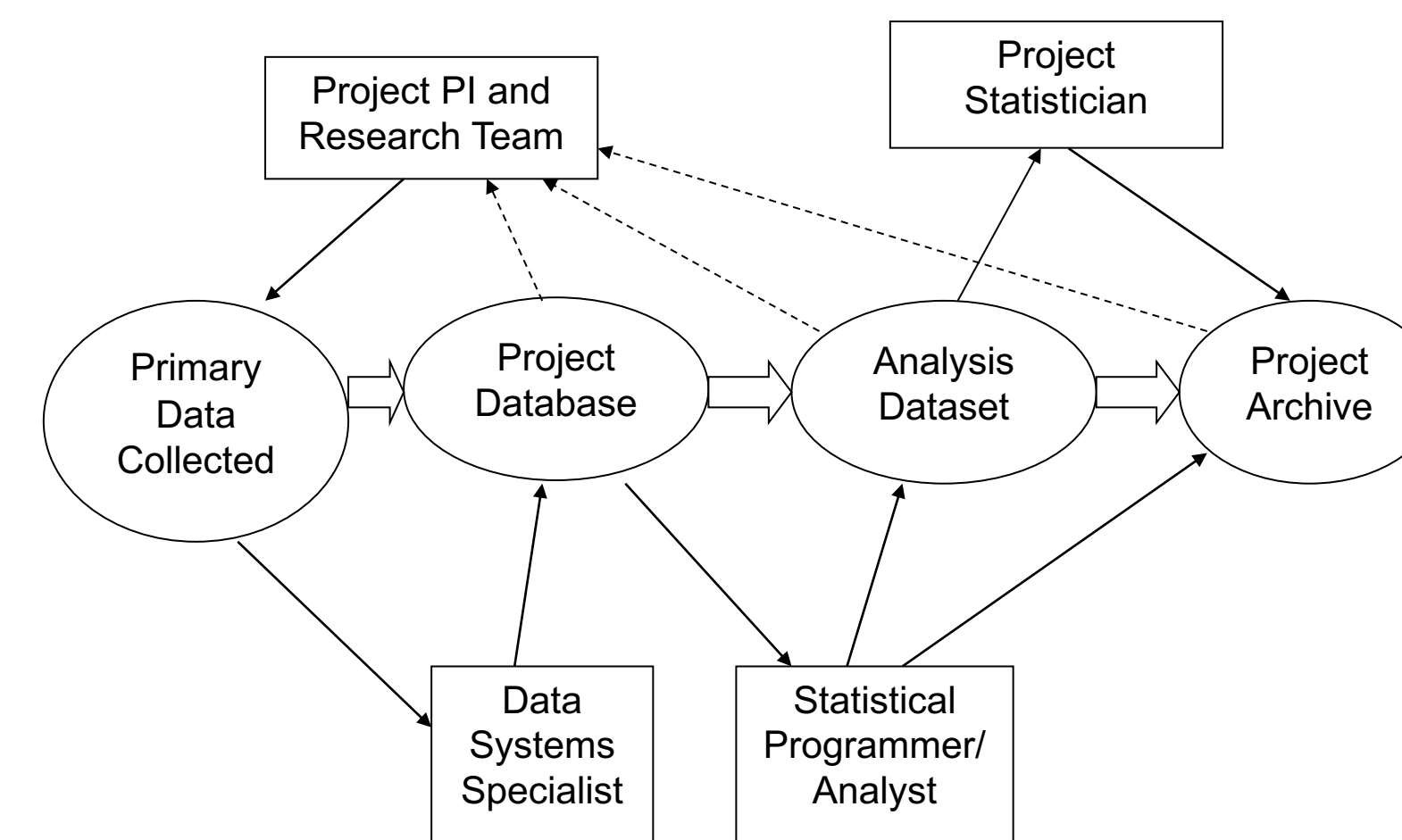
Value Added

- Access to specialized biostatistics faculty expertise
- Ensure optimal study design for expensive data
- Ensure rigorous and cutting edge analytic methods for grant proposals and manuscripts
- Ensure reproducibility of analyses

Key areas of expertise

- Design and analysis of randomized trials
- rNA-seq and single-cell rNA methods
- Spatial transcriptomics
- Causal inference for observational cohort data
- Machine learning and prediction
- Radiomics and image analysis
- Design and analysis for pre-clinical studies and in-vivo experiments

Data Workflow and Pipeline



Key Publications

- Huntington, K.E., Louie, A.D., Srinivasan, P.R., Schorl, C., Lu, S., Silverberg, D., Newhouse, D., Wu, Z., Zhou, L., Borden, B.A. & Giles, F.J. (2023). GSK-3 inhibitor elraglisib enhances tumor-infiltrating immune cell activation in tumor biopsies and synergizes with anti-PD-L1 in a murine model of colorectal cancer. *International Journal of Molecular Sciences*, 24(13), 10870.
- De Souza A.L., Mega A.E., Douglass J., Olszewski A.J., Gamsiz Uzun E.D., Uzun A., Chou C., Duan F., Wang J., Ali A., Golijanin D.J., Holder S.L., Lagos G.G., Safran H., El-Deiry W., Carneiro B.A. (2023). Clinical features of patients with MTAP-deleted bladder cancer. *Am J Cancer Res*.
- Bocage A., Orang'o O., Liu T., Itsura P., Tonui P., Muthoka K., Stephen K., Sam S.S., Caliendo A., Cu-Uvin S (2022). HIV-1 RNA genital tract shedding after cryotherapy for visual inspection with acetic acid-positive cervical lesions in western Kenya. *Am Journal of Obstetrics & Gynecology* 226, 291-292.
- Ma, Y., & Zhou, X. (2022). Spatially informed cell-type deconvolution for spatial transcriptomics. *Nature Biotechnology*, 40, 1349-1359.
- Comstock, C. E., Gatsonis, C., et al. (2020). Comparison of Abbreviated Breast MRI vs Digital Breast Tomosynthesis for breast cancer detection among women with dense breasts undergoing screening. *JAMA*, 323(8), 746-756.
- Steingrimsson, J. A., Gatsonis, C., Li, B., Dahabreh, I.J. (2023). Transporting a Prediction Model for Use in a New Target Population. *Am J Epidemiol*, 192(2), 296-304.
- Morrison, S., Gatsonis, C., Eloyan, A., Steingrimsson, J.A (2023). Survival analysis using deep learning with medical imaging *Int J Biostat*.
- Carlos, R. C., Obeng-Gyasi, S., Cole, S. W., Zebrack, B. J., Pisano, E. D., Troester, M. A., Timsina, L., Wagner, L. I., Steingrimsson, J. A., Lee, C. I., Adams, A., & Wilkins, C. H. (2022). Linking Structural Racism and Discrimination and Breast Cancer Outcomes: A Social Genomics Approach. *Journal of Clinical Oncology*, 40(13), 1407-1413.
- Partridge, S.C., Steingrimsson, J.A., Newitt, D.C., Gibbs, J.E., Marques, H.S., Bolan, P.J., Chenevert, T.L., Rosen, M.A., & Hylton, N.M. (2022). Impact of Alternate b-Value Combinations and Metrics on Predictive Performance and Repeatability of Diffusion-Weighted MRI in Breast Cancer Treatment: Results from the ECOG-ACRIN A6698 Trial. *Tomography*, 8(2), 701-717.
- Xu K., Diaz A.A., Duan F., Lee M., Xiao X., Liu H., Liu G., Cho M.H., Gower A.C., Alekseyev Y.O., Spira A., Aberle D.R., Washko G.R., Billatos E., Lenburg M.E.; DECAMP investigators (2023). Bronchial Gene Expression Alterations Associated with Radiographic Bronchiectasis. *Eur Respir J*, 61(1):2200120.
- Hu L, Ji J, Ennis R, Hogan JW (2022). A flexible approach for causal inference with multiple treatments and clustered survival outcomes. *Statistics in Medicine* 41, 4982-4999.
- Stack B.C Jr, Duan F, Romanoff J, Sicks J.D., Subramaniam R.M., Lowe V.J., (2023). Impact of Neck PET/CT Positivity on Survival Outcomes-Visual and Quantitative Assessment: Results From ACRIN 6685. *Clin Nucl Med*, 48(2):126-131.
- Black, W. C., Gareen, I. F., ..., Gatsonis, C. (2014). Cost-effectiveness of CT screening in the National Lung Screening Trial. *New England Journal of Medicine*, 371(19), 1793-1802.

Future Plans

- Recruit faculty with SPH / LCC support
- Build user base
- Formalize Shared Resource operations and budgetary support
- Program of research focused on developing innovative statistical and data science methods