# BONNIE THIEL

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#### SUMMARY

Data scientist with 15+ years of experience in statistical modeling of experimental and population data. Expert in building statistical models in R and SAS, hypothesis testing and power/sample size estimates. Managed projects and data collection processes for tuberculosis research projects including database design and construction for a large TB consortium. Interested in applying statistical and bioinformatics tools to answer questions in biology and medicine.

# **EDUCATION**

Case Western Reserve University, Cleveland, OH Ph.D. Systems Biology and Bioinformatics, 2021

Case Western Reserve University, Cleveland, OH M.S. Epidemiology and Biostatistics, 1997

University of Colorado, Boulder, CO B.S. Molecular, Cellular and Developmental Biology and Biochemistry, 1988

#### **EXPERIENCE**

2003 – present

**Specialist III** Case Western Reserve University, Tuberculosis Research Unit, Department of Medicine

- Statistical analysis of epidemiologic and experimental data relating to tuberculosis infection and disease
- Bioinformatics analysis including proteomics, transcriptomics and genomics
- Project and data management for a large international research consortium
- Production of manuscripts, reports, protocols and procedures

Contact: Dr. Henry W Boom. 216-368-4847. whb@case.edu.

2001-2003

**Specialist II** Case Western Reserve University, Department of Epidemiology and Biostatistics

1996-2001

**Specialist I** Case Western Reserve University, Department of Epidemiology and Biostatistics

- Statistical analysis and modeling of data in molecular and genetic epidemiology.
- Synthesis and summary of analysis in the form of reports and publications.

Contact: Dr. Nicholas Schork, Professor. <a href="mailto:nschork@ucsd.edu">nschork@ucsd.edu</a>.

#### 1994-1996

**Research Assistant II** Case Western Reserve University, Department of Genetics

- Lab manager duties included maintaining equipment and ordering supplies.
- Performed molecular genetic experiments using human and animal DNA and RNA including genotyping and mutation analysis.
- Tissue culture experience included growing and transforming human lympocytes and harvesting DNA.

Contact: Dr. Aravinda Chakravarti, Professor. aravinda@jhmi.edu.

#### 1993

**Student Assistant** Case Western Reserve University, Department of Epidemiology and Biostatistics

- Involved in primary data collection for a study on periphreal vascular disease
- Interviewed patients and tracked data collection.

# 1991-1994

**Technologist B** The Cleveland Clinic Foundation, Department of Immunolgy.

- Worked on projects involving the molecular biology and protein chemistry of nitric oxide synthase.
- Gained experience in techniques including expression vector systems, transient and stables transfection and RNA and DNA purification.

Contact: Dr. Dennis Stuehr, Staff. 216-445-6950. stuehrd@ccf.org.

#### 1989-1990

**Technician II** Cornell University Medical College, Department of Cell Biology.

Contributed to a study on the nerve growth factor receptor.

 Performed receptor-ligand binding assays, Scatchard analysis, crosslinking reactions, iodinations, transfection, nick translation and western and northern blotting.

Contact: Dr. Barbara Hempstead, Professor. blhempst@med.cornell.edu

### **SKILLS**

<u>Statistical:</u> Expert level in modeling and prediction applied to data derived from experimental settings, clinical studies, large population studies and data repositories. Working knowledge of machine learning techniques.

<u>Programming:</u> High level SAS and R programming for statistical analysis and data mangagement, experience with Unix/Linux computing systems and scripting

<u>Software Tools:</u> REDCap, OpenClinica, GraphPad, Powerpoint, Access, Word <u>Additional skills:</u> Study design, power and sample size calculation, data management, database systems, data collection design and process, quality control/ assessment.

## **PUBLICATIONS**

**Thiel BA**, Worodria W, Nalukwago S, Nsereko M, Sanyu I, Rejani L, Zawedde J, Canaday DH, Stein CM, Chervenak KA, Malone LL, Kiyemba R, Silver RF, Johnson JL, Mayanja-Kizza H, Boom WH. Immune cells in bronchoalveolar lavage fluid of Ugandan adults who resist versus those who develop latent Mycobacterium tuberculosis infection. *PLoS ONE*. 2021;16. <a href="https://doi.org/10.1371/journal.pone.0249477">https://doi.org/10.1371/journal.pone.0249477</a>

Weiner J, 3rd, Maertzdorf J, Sutherland JS, Duffy FJ, Thompson E, Suliman S, McEwen G, **Thiel B**, Parida SK, Zyla J, Hanekom WA, Mohney RP, Boom WH, Mayanja-Kizza H, et al. Metabolite changes in blood predict the onset of tuberculosis. *Nature Communications*. 2018;9. <a href="https://doi.org/10.1038/s41467-018-07635-7">https://doi.org/10.1038/s41467-018-07635-7</a>

Ronacher K, Chegou NN, Kleynhans L, Djoba Siawaya JF, du Pleiss N, Loxton AG, Maasdorp E, Tromp G, Kidd M, Stanley K, Kriel M, Menezes A, Gutschmidt A, van der Spuy GD, Warren RM, Dietze R, Okwera A, **Thiel B**, Belisle JT, Cliff JM, Boom WH, Johnson JL, van Helden PD, Dockrell HM, Walzl G. Distinct serum biosignatures are associated with different tuberculosis treatment outcomes. *Tuberculosis*. 2019:118. <a href="https://doi.org/10.1016/j.tube.2019.101859">https://doi.org/10.1016/j.tube.2019.101859</a>

Duffy FJ , Weiner J , Hansen S, Tabb DL, Suliman S , Thompson E, Maertzdorf J, Shankar S , Tromp G , Parida S , Dover D , Axthelm MK , Sutherland JS , Dockrell HM, Ottenhoff THM, Scriba TJ , Picker LJ , Walzl G , Kaufmann SHE , Zak DE. **The GC6-74 Consortium.** Immunometabolic Signatures Predict Risk of Progression to Active Tuberculosis and Disease Outcome. *Frontiers In Immunology*. 2019:10:527. https://doi.org/10.3389/fimmu.2019.00527

Dousa KM, Hamad A, Albirair M, Al Soub H, Elzouki AN, Alwakeel M, **Thiel BA**, Johnson JL. Impact of diabetes mellitus on the presentation and response to treatment of adults with pulmonary tuberculosis in Qatar. *Open Forum Infectious Disease*. 2019;6. https://doi.org/10.1093/ofid/ofy335

Stein CM, Zalwango S, Malone LL, **Thiel B**, Mupere E, Nsereko M, Okware B, Kisingo H, Lancioni CL, Bark CM, Whalen CC, Joloba ML, Boom WH, Mayanja-Kizza H. Resistance and Susceptibility to Mycobacterium Tuberculosis Infection and Disease in Tuberculosis Households in Kampala, Uganda. *Am J Epidemiol.* 2018;187:1477. https://doi.org/10.1093/aje/kwx380

Suliman S, Thompson E, Sutherland J, Weiner Rd J, Ota MOC, Shankar S, Penn-Nicholson A, **Thiel B**, Erasmus M, Maertzdorf J, Duffy FJ, Hill PC, Hughes EJ, Stanley K, Downing K, Fisher ML, Valvo J, Parida SK, van der Spuy G, Tromp G, Adetifa IMO, Donkor S, Howe R, Mayanja-Kizza H, Boom WH, Dockrell H, Ottenhoff THM, Hatherill M, Aderem A, Hanekom WA, Scriba TJ, Kaufmann SH, Zak DE, Walzl G Four-gene Pan-African Blood Signature Predicts Progression to Tuberculosis. *Am J Respir Crit Care Med*. 2018;197:1198. <a href="https://doi.org/10.1164/rccm.201711-2340OC">https://doi.org/10.1164/rccm.201711-2340OC</a>

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Bark CM, **Thiel BA**, Ogwang S, Sekitoleko G, Muzanyi G, Joloba ML, Johnson JL. Sputum smear-positive, culture-negative state during anti-tuberculosis treatment in the MGIT liquid culture era. *The International Journal of Tuberculosis and Lung Disease*. 2018;33:306. <a href="https://doi.org/10.5588/ijtld.17.0655">https://doi.org/10.5588/ijtld.17.0655</a>

Alsultan A, Furin JJ, Du Bois J, van Brakel E, Chheng P, Venter A, **Thiel B**, Debanne SA, Boom WH, Diacon AH, Johnson JL, Peloquin CA. Population Pharmacokinetics of AZD-5847 in Adults with Pulmonary Tuberculosis. *Antimicrob Agents Chemother*. 2016;60:6591. https://doi.org/10.1128/AAC.01163-16

Zak DE, Penn-Nicholson A, Scriba TJ, Thompson E, Suliman S, Amon LM, Mahomed H, Erasmus M, Whatney W, Hussey GD, Abrahams D, Kafaar F, Hawkridge T, Verver S, Hughes EJ, Ota M, Sutherland J, Howe R, Dockrell HM, Boom WH, **Thiel B**, Ottenhoff THM, Mayanja-Kizza H, Crampin AC, Downing K, Hatherill M, Valvo J, Shankar S, Parida SK, Kaufmann SHE, Walzl G, Aderem A, Hanekom WA. A blood RNA signature for tuberculosis disease risk: a prospective cohort study. *Lancet*. 2016;387:2312. https://doi.org/10.1016/S0140-6736(15)01316-1

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**Thiel BA**, Bark CM, Nakibali JG, van der Kuyp F, Johnson JL. Reader variability and validation of the Timika X-ray score during treatment of pulmonary tuberculosis. *IJTLD* 2016; 20:1358. https://doi.org/10.5588/ijtld.16.0186

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Tao L, Zalwango S, Chervenak K, **Thiel B**, Malone LL, Qiu F, Mayanja-Kizza H, Boom WH, Stein CM; Tuberculosis Research Unit. Genetic and shared environmental influences on interferon-γ production in response to *Mycobacterium tuberculosis* antigens in a Ugandan population. *Am J Trop Med Hyg* 2013;89:169. <a href="https://doi.org/10.4269/ajtmh.12-0670">https://doi.org/10.4269/ajtmh.12-0670</a>

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Mahan CS, Zalwango S, **Thiel BA**, Malone LL, Chervenak KA, Baseke J, Dobbs D, Stein CM, Mayanja H, Joloba M, Whalen CC, Boom WH. Innate and adaptive immune responses during acute M tuberculosis infection in adult household contacts in Kampala, Uganda. *Am J Trop Med Hyg.* 2012;86:690. https://doi.org/10.4269/ajtmh.2012.11-0553

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Bark CM Okwera A, Joloba ML, **Thiel BA**, Nakibali JG, Debanne SM, Boom WH, Eisenach KD and Johnson JL. Time to detection of *Mycobacterium tuberculosis* as an alternative to quantitative cultures. *Tuberculosis*. 2011;91:257. https://doi.org/10.1016/j.tube.2011.01.004

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receptors 1 and 2 *Infection and immunity* 2011;79:663. https://doi.org/10.1128/IAI.00806-10

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Koivukoski L, Fisher SA, Kanninen T, Lewis CM, von Wowern F, Hunt S, Kardia SLR, Levy D, Perola M, Rankinin T, Rao DC, Rice T, **Thiel BA**, Melander O. Meta-analysis of genome-wide scans for hypertension and blood pressure in Caucasians shows evidence of susceptibility regions on chromosomes 2 and 3. *Human Molecular Genetics*. 2004;13:2325. <a href="https://doi.org/10.1093/hmg/ddh237">https://doi.org/10.1093/hmg/ddh237</a>

Witte JS , Suarez BK, **Thiel B**, Lin J, Yu A, Banerjee TK, Burmester JK, Casey G, Catelona WJ. Genome-wide scan of brothers: replication and fine mapping of prostate cancer susceptibility and aggressiveness loci *Prostate*. 2003;1:298. <a href="https://doi.org/10.1002/pros.10304">https://doi.org/10.1002/pros.10304</a>

**Thiel B**, Chakravarti A, Cooper R, Lewis S, Lynn A, Tiwari H, Schork NJ and Weder AB. A genome wide linkage analysis investigating the determinants of blood pressure in Caucasians and African Americans. *Amer J Hypertens* 2003;16:151. <a href="https://doi.org/10.1016/S0895-7061(02)03246-6">https://doi.org/10.1016/S0895-7061(02)03246-6</a>

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Stuehr DJ, Kwon NS, Gross SS, **Thiel BA**, Levi R, Nathan CF. Synthesis of nitrogen oxides from L-Arginine by macrophage cytosol: requirement for inducible and constitutive components *Biochem Biophys Res Comm* 1989;161:420. https://doi.org/10.1016/0006-291X(89)92615-6