

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

PI: Dr. Henry W Boom. 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.

PI: Dr. Nicholas Schork, Professor

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 lymphocytes and harvesting DNA.

PI: Dr. Aravinda Chakravarti, Professor

1993

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

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

1991-1994

Technologist B The Cleveland Clinic Foundation, Department of Immunology.

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

PI: Dr. Dennis Stuehr, Staff

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.

Supervisor: Dr. Barbara Hempstead, Professor

SKILLS

Statistical: 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.

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

Software Tools: REDCap, OpenClinica, GraphPad, Powerpoint, Access, Word

Additional skills: 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.
<https://doi.org/10.1371/journal.pone.0249477>

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

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.
<https://doi.org/10.1016/j.tube.2019.101859>

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. <https://doi.org/10.1164/rccm.201711-2340OC>

Duffy FJ, Thompson E, Downing K, Suliman S, Mayanja-Kizza H, Boom WH, **Thiel B**, Weiner Iii J, Kaufmann SHE, Dover D, Tabb DL, Dockrell HM, Ottenhoff THM, Tromp G, Scriba TJ, Zak DE, Walzl G, et al. A Serum Circulating miRNA Signature for Short-Term Risk of Progression to Active Tuberculosis Among Household Contacts. *Frontiers In Immunology*. 2018;9:661. <https://doi.org/10.3389/fimmu.2018.00661>

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. <https://doi.org/10.5588/ijtld.17.0655>

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, Hawkrigde 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](https://doi.org/10.1016/S0140-6736(15)01316-1)

Furin JJ, Du Bois J, van Brakel E, Chheng P, Venter A, Peloquin CA, Alsultan A, **Thiel BA**, Debanne SM, Boom WH, Diacon AH, Johnson JL. Early bactericidal activity of AZD5847 in patients with pulmonary tuberculosis. *Antimicrobial Agents and Chemotherapy*. 2016;60:6591. <https://doi.org/10.1128/AAC.01163-16>

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>

Benghiac AG, **Thiel BA**, Haba D. Reliability of the frontal sinus index for sex determination using CBCT. *Romanian Journal of Legal Medicine*. 2015;23:275. <https://doi.org/10.4323/rjlm.2015.275>

Chen J, Leskov IL, Jr AY, **Thiel B**, Kevil CG, Stokes KY, Orr AW. Recruitment of the adaptor protein Nck to PECAM-1 couples oxidative stress to canonical NF- κ B signaling and inflammation. *Sci Signal*. 2015;8:ra20. <https://doi.org/10.1126/scisignal.2005648>

Revsbech IG, Shen X, Chakravarti R, Jensen FB, **Thiel B**, Evans AL, Kindberg J, Frobert, O, Stuehr DJ, Kevil, CG, Fago A. Hydrogen sulfide and nitric oxide metabolites in the blood of free-ranging brown bears and their potential roles in hibernation. *Free Radic Biol Med*. 2014;73:349. <https://doi.org/10.1016/j.freeradbiomed.2014.05.025>

Ma N, Zalwango S, Malone LL, Nsereko M, Wampande EM, **Thiel BA**, Okware B, Igo RP Jr, Joloba ML, Mupere E, Mayanja-Kizza H, Boom WH, Stein CM. Clinical and epidemiological characteristics of individuals resistant to M tuberculosis infection in a longitudinal TB household contact study in Kampala, Uganda. *BMC Infect Dis*. 2014;14:352. <https://doi.org/10.1186/1471-2334-14-352>

Johnson DF, Malone LL, Zalwango S, Mukisa Oketcho J, Chervenak KA, **Thiel B**, Mayanja-Kizza H, Stein CM, Boom WH, Lancioni CL, Tuberculosis Research Unit. Tuberculin skin test reversion following isoniazid preventive therapy reflects diversity of immune response to primary *Mycobacterium tuberculosis* infection. *PLoS ONE* 2014;9(5):e96613. <https://doi.org/10.1371/journal.pone.0096613>

Johnson JL, Geldenhuys H, **Thiel BA**, Toefy A, Suliman S, Pienaar B, Chheng P, Scriba T, Boom WH, Hanekom W, Hatherill M. Effect of isoniazid therapy for latent TB infection on QuantiFERON-TB Gold In-tube responses in tuberculin skin test positive adults in a high TB incidence area: A controlled study. *Chest* 2014;145:612. <https://doi.org/10.1378/chest.13-1232>

Sutherland JS, Loxton AG, Haks MC, Kassa D, Ambrose L, Lee JS, Ran L, van Baarle D, Maertzdorf J, Howe R, Mayanja-Kizza H, Boom WH, **Thiel BA**, Crampin AC, Hanekom W, Ota MO, Dockrell H, Walzl G, Kaufmann SH, Ottenhoff TH; GCGH Biomarkers for TB consortium. Differential gene expression of activating Fc γ receptor classifies active tuberculosis regardless of human immunodeficiency virus status or ethnicity. *Clin Microbiol Infect* 2014;20(4):O230-8. <https://doi.org/10.1111/1469-0691.12383>

Sutherland JS, Lalor MK, Black GF, Ambrose LR, Loxton AG, Chegou NN, Kassa D, Mihret A, Howe R, Mayanja-Kizza H, Gomez MP, Donkor S, Franken K, Hanekom W, Klein MR, Parida SK, Boom WH, **Thiel BA**, Crampin AC, Ota M, Walzl G, Ottenhoff

TH, Dockrell HM, Kaufmann SH, GCGH Biomarkers for TB consortium. Analysis of Host Responses to *Mycobacterium tuberculosis* Antigens in a Multi-Site Study of Subjects with Different TB and HIV Infection States in Sub-Saharan Africa. *PLoS ONE* 8(9): e74080. <https://doi.org/10.1371/journal.pone.0074080>

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. <https://doi.org/10.4269/ajtmh.12-0670>

Bark CM, Gitta P, Ogwang S, Nsereko M, **Thiel BA**, Eisenach KD, Boom WH, Joloba ML, Johnson JL. Comparison of time to positive and colony counting in an early bactericidal activity study of anti-tuberculosis treatment *Int J Tuberc Lung Dis* 2013;11:1448. <https://doi.org/10.5588/ijtld.13.0063>

Johnson JL and **Thiel BA**. Time until relapse in tuberculosis treatment trials: Implication for phase 3 trial design *American Journal of Respiratory and Critical Care Medicine* 2012;186:464. <https://doi.org/10.1164/ajrccm.186.5.464>

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>

Bark CM, **Thiel BA** and Johnson JL. Pretreatment time to detection of *Mycobacterium tuberculosis* in liquid culture is associated with relapse after therapy *J Clin Microbiol*. 2012;50:538. <https://doi.org/10.1128/JCM.06193-11>

Bark CM, Dietze R, Okwera A, Quelapio MI, **Thiel BA** and Johnson JL. Clinical symptoms and microbiological outcomes in tuberculosis treatment trials. *Tuberculosis*. 2011;91:601. <https://doi.org/10.1016/j.tube.2011.05.007>

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>

Toossi Z, Hirsch CS, Wu M, Mayanja-Kizza H, Baseke J and **Thiel B**. Distinct cytokine and regulatory T cell profile at pleural sites of dual HIV/tuberculosis infection compared to that in the systemic circulation. *Clin Exp Immunol* 2011;163:333. <https://doi.org/10.1111/j.1365-2249.2010.04269.x>

Lancioni CL, Li Q, Thomas JJ, Ding X, **Thiel B**, Drage MG, Pecora ND, Ziady AG, Shank S, Harding CV, Boom W and Rojas RE. *Mycobacterium tuberculosis* lipoproteins directly regulate human memory CD4(+) T cell activation via Toll-like

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

Black GF, **Thiel BA**, Ota MO, Parida SK, Adegbola R, Boom WH, Dockrell HM, Franken KL, Friggen AH, Hill PC, Klein MR, Lalor MK, Mayanja H, Schoonik G, Stanley K, Weldingh K, Kaufmann SH, Walzl G, Ottenhoff TH; GCGH Biomarkers for TB Consortium. Immunogenicity of novel DosR regulon-encoded candidate antigens of *Mycobacterium tuberculosis* in three high-burden populations in Africa *Clin Vaccine Immunol* 2009;16:1203. <https://doi.org/10.1128/CVI.00111-09>

Rojas RE, Chervenak KA, Thomas J, Morrow J, Nshuti L, Zalwango S, Mugerwa RD, **Thiel BA**, Whalen CC, Boom WH. Vδ2⁺ γδ T Cell Function in *Mycobacterium tuberculosis*- and HIV-1-Positive Patients in the United States and Uganda: Application of a Whole-Blood Assay. *Journal of Infectious Disease* 2005;192:1806.
<https://doi.org/10.1086/497146>

Visscher PM, Haley CS, Ewald H, Mors O, Egeland J, **Thiel B**, Ginns E, Muir W, Blackwood DH. Joint multi-population analysis for genetic linkage of bipolar disorder or “wellness” to chromosome 4p. *Am J Med Genet B Neuropsychiatr Genet.* 2005;133:18.
<https://doi.org/10.1002/ajmg.b.30108>

Koivukoski L, Fisher SA, Kanninen T, Lewis CM, von Wöhrn F, Hunt S, Kardina SLR, Levy D, Perola M, Rankinen 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. <https://doi.org/10.1093/hmg/ddh237>

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.
<https://doi.org/10.1002/pros.10304>

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.
[https://doi.org/10.1016/S0895-7061\(02\)03246-6](https://doi.org/10.1016/S0895-7061(02)03246-6)

Province MA, Kardina SLR, Ranade K, Rao DC, **Thiel BA**, Cooper RS, Risch N, Turner ST, Cox DR, Hunt SC, Weder AB, Boerwinkle E. A meta-analysis of genome-wide linkage scans for hypertension: The National Heart Lung and Blood Institute Family Blood Pressure Program *Amer J Hypertens.* 2003;16:144-147.
[https://doi.org/10.1016/S0895-7061\(02\)03248-X](https://doi.org/10.1016/S0895-7061(02)03248-X)

Schork NJ, Gardner JP, Zhang L, Fallin D, **Thiel B**, Jakubowski H, Aviv A. Genomic association/linkage of sodium lithium countertransport in CEPH pedigrees *Hypertension.* 2002;40:619. <https://doi.org/10.1161/01.HYP.0000037131.41957.A8>

Hall MA, Norman P, **Thiel B**, Tiwari H, Peiffer A, Vaughan RW, Prescott S, Leppert M, Schork NJ, Lanchbury JS. Quantitative trait loci on chromosomes 1, 2, 3, 4, 8, 9, 11, 12 and 18 control variation in levels of T and B lymphocyte sub-populations *Am J Hum Genet.* 2002;70:1172. <https://doi.org/10.1086/340090>

Morrison SJ, Qian D, Jerabek L, **Thiel BA**, Park IK, Ford PS, Kiel JM, Schork NJ, Weissman IL, Clark MF. A genetic determinant that specifically regulates the frequency of hematopoietic stem cells *J Immunol.* 2002;168:635. <https://doi.org/10.4049/jimmunol.168.2.635>

Philibert RA, Cheung D, Welsh N, Damschroder-Williams P, **Thiel B**, Ginns EI, Gershenfeld HK. Absence of a significant linkage between Na⁺, K⁺-ATPase subunit (*ATP1A3* and *ATP1B3*) genotypes and bipolar affective disorder in the Old-Order Amish. *Amer J Med Gen.* 2001;105:291. <https://doi.org/10.1002/ajmg.1322>

Schork NJ, Fallin D, **Thiel B**, Xu X, Broeckel U, Jacob H, and Cohen D. The future of genetic case/control studies. *Advances in Genetics* 2000;42:191. [https://doi.org/10.1016/S0065-2660\(01\)42023-2](https://doi.org/10.1016/S0065-2660(01)42023-2)

Thiel B and Weder AB. Genes for Essential Hypertension: Hype, Help, or Hope? *J Clin Hypertens.* 2000;2:187. http://www.lejacq.com/articleDetail.cfm?pid=J%20Clin%20Hypertens_2;3:187

Schork, NJ, Chakravarti A, **Thiel B**, Fornage M, Jacob HJ, Cai R, Rotimi CN, Cooper RS, Weder, AB Lack of Association between a biallelic polymorphism in the adducin gene and blood pressure in Whites and African Americans *Amer J Hum Gen.* 2000;13:693. [https://doi.org/10.1016/S0895-7061\(00\)00237-5](https://doi.org/10.1016/S0895-7061(00)00237-5)

Ginns EI, St Jean P, Philibert RA, Galdzicka M, Damschroder-Williams P, **Thiel B**, Long RT, Ingraham LJ, Dalwaldi H, Murray MA, Paul S, Remortel BG, Patel AP, Anderson MCH, Shaio C, Lau E, Dymarskaia I, Martin BM, Stubblefield B, Falls KM, Carulli JP, Keith TP, Fann CSJ, Lacy LG, Allen CR, Hostetter AM, Elston RC, Schork NJ, Egeland JA, Paul SM. A genome-wide search for chromosomal loci linked to mental health wellness in relatives at high risk for bipolar affective disorder among the Old Order Amish. *Proc Natl Acad Sci USA.* 1998;95:15531. <https://doi.org/10.1073/pnas.95.26.15531>

Bix M, Wang ZE, **Thiel B**, Schork NJ, Locksley RM. Genetic regulation of commitment to interleukin 4 production by a CD4(+) T cell-intrinsic mechanism *J Exp Med* 1998;188:2289. <https://doi.org/10.1084/jem.188.12.2289>

Faust C, Lawson KA, Schork NJ, **Thiel B**, Magnuson T. The Polycomb-group gene *ee* is required for normal morphogenic movements during gastrulation in the mouse embryo *Development* 1998;125:4495. <https://doi.org/10.1242/dev.125.22.4495>

Allison DB, **Thiel B**, St Jean, P Elston, RC Infante MC, Schork, NJ. Multiple Phenotype modeling in gene-mapping studies of quantitative traits: power advantages *Am J Hum Genet* 1998;63:11990. <https://doi.org/10.1086/302038>

Schork, NJ, **Thiel B**, St Jean P. Linkage analysis, kinship, and the short-term evolution of chromosomes *J Exp Zool* 1998;282:133.
[https://doi.org/10.1002/\(SICI\)1097-010X\(199809/10\)282:1/2<133::AID-JEZ15>3.0.CO;2-V](https://doi.org/10.1002/(SICI)1097-010X(199809/10)282:1/2<133::AID-JEZ15>3.0.CO;2-V)

Schork NJ, Allison DB and **Thiel B**. Mixture distributions in human genetics research *Stat Methods Med Res* 1996;5:155. <https://doi.org/10.1177/096228029600500204>

Angrist M, Bolk S, **Thiel B**, Puffenberger EG, Hofstra RM, Buys CHCM, Cass DT, Chakravarti A. Mutation analysis of the RET receptor tyrosine kinase in Hirschsprung disease *Hum Mol Gen* 1995;4:821. <https://doi.org/10.1093/hmg/4.5.821>

Shaw SH, Farr JEW, **Thiel BA**, Matise TC, Weissenbach J, Chakravarti A, Richard CW. A radiation map of 95 STS's spanning human chromosome 13q *Genomics*. 1995;27:502. <https://doi.org/10.1006/geno.1995.1083>

Baek KJ, **Thiel BA**, Lucas S, Stuehr DJ. Macrophage nitric oxide synthase subunits: Purification, characterization, and role for prosthetic groups and substrate in regulating their assembly into a dimeric enzyme *J Biol Chem* 1993;268:21120. [https://doi.org/10.1016/S0021-9258\(19\)36901-7](https://doi.org/10.1016/S0021-9258(19)36901-7)

Deng WD, **Thiel B**, Tannenbaum CS, Hamilton TA, Stuehr DJ. Synergistic cooperation between T-cell lymphokines for induction of the nitric oxide synthase gene in murine peritoneal macrophages *J Immunol* 1993;151:322.
<http://www.jimmunol.org/content/151/1/322>

Hempstead BL, Patil N, **Thiel B**, Chao MV. Deletion of cytoplasmic sequences of NGF receptor leads to loss of high affinity ligand binding *J Biol Chem* 1990;265:9595. [https://doi.org/10.1016/S0021-9258\(19\)38707-1](https://doi.org/10.1016/S0021-9258(19)38707-1)

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](https://doi.org/10.1016/0006-291X(89)92615-6)