

## Alexej Gossmann

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<b>AREAS OF INTEREST</b>	Sparse Regression Models, Mixed Effects Models, Genetics, Spatial Statistics, Computational Statistics, Differential Equations
<b>EDUCATION</b>	<p><i>PhD</i>, Mathematics Tulane University, New Orleans, Louisiana, expected May 2017</p> <p><i>MS</i>, Statistics Tulane University, New Orleans, Louisiana, May 2014 Master's Research Project: Analysis of Bone Growth Data by Mixed-Effects SS ANOVA Methods (supervised by Dr. Lacey) GPA: 3.975</p> <p><i>BS</i>, Mathematics Technische Universität Darmstadt, Darmstadt, Germany, May 2012 Thesis: On disjunction and numerical existence properties of extensions of Heyting arithmetic (supervised by Dr. Kohlenbach) GPA: 4.0</p>
<b>EXPERIENCE</b>	<p><i>Teaching Experience</i></p> <ul style="list-style-type: none"><li>• <i>Instructor</i>: Calculus 1, Tulane University, Department of Mathematics, Fall 2014.</li><li>• <i>Co-Teacher</i>: Statistics for Scientists, Tulane University, Department of Mathematics, Spring 2014.</li><li>• <i>Teaching Assistant</i>: Various Statistics, Calculus, and Real Analysis courses at Tulane University and Technische Universität Darmstadt, Fall 2010 - Spring 2014.</li></ul> <p><i>Internships</i></p> <ul style="list-style-type: none"><li>• <i>Google Summer of Code 2015</i>. Adding Linear Mixed Effects Models Support to SciRuby. Supervised by the Ruby Science Foundation. May – August 2015.</li></ul> <p><i>Service</i></p> <ul style="list-style-type: none"><li>• President of the SIAM student chapter at Tulane University. September 2014 – Present.</li><li>• Main organizer of the Graduate Student Colloquium at the department of Mathematics at Tulane University. September 2014 – Present.</li></ul> <p><i>Research Assistantship</i></p> <ul style="list-style-type: none"><li>• Development of statistical and machine learning methods for imaging-genomics in Dr. Yu-Ping Wang's group. January 2015 – Present.</li><li>• Testing a linear correlation between two <math>L^2</math> spatial fields, supervised by Dr. Gromenko, Tulane University, Department of Mathematics. June – August 2014.</li></ul> <p><i>Other</i></p>

- *SAMSI 2014 Mathematical and Statistical Modeling Workshop for Graduate Students*. Project: Allergy, Asthma and Exposures in the Homes of the US Population (supervised by scientists from Rho Inc.), North Carolina State University. July 2014.
- Kommando 1. Luftwaffendivision, Fürstenfeldbruck. Military service at the department for public relations and press. June 2008 – February 2009.

- PUBLICATIONS**
- [1] Shaolong Cao, Huaizhen Qin, Alexej Gossmann, Hong-Wen Deng, and Yu-Ping Wang. Unified tests for fine scale mapping and identifying sparse high-dimensional sequence associations. *Bioinformatics*, 2015.
  - [2] Mimi C Sammarco, Jennifer Simkin, Alexander J Cammack, Danielle Fassler, Alexej Gossmann, Luis Marrero, Michelle Lacey, Keith Van Meter, and Ken Muneoka. Hyperbaric oxygen promotes proximal bone regeneration and organized collagen composition during digit regeneration. *PloS one*, 10(10), 2015.
  - [3] Shaolong Cao, Huaizhen Qin, Alexej Gossmann, Hong-Wen Deng, and Yu-Ping Wang. Unified tests for fine scale mapping and identifying sparse high-dimensional sequence associations. In *Proceedings of the 6th ACM Conference on Bioinformatics, Computational Biology and Health Informatics*, BCB '15, pages 241–249, New York, NY, USA, 2015. ACM.
  - [4] Alexej Gossmann, Shaolong Cao, and Yu-Ping Wang. Identification of significant genetic variants via slope, and its extension to group slope. In *Proceedings of the 6th ACM Conference on Bioinformatics, Computational Biology and Health Informatics*, BCB '15, pages 232–240, New York, NY, USA, 2015. ACM.

- TALKS AND COLLOQUIA**
- Title: *Identification of Significant Genetic Variants via SLOPE, and its Extension to Group SLOPE*. The 6th ACM Conference on Bioinformatics, Computational Biology, and Health Informatics, Atlanta, GA, 2015.
  - Title: *Reproducing Kernel Hilbert Spaces and Smoothing Spline Regression*. Graduate student colloquium, Tulane University, 2014.

- POSTER PRESENTATIONS**
- A. Gossmann, S. Cao, and Y.-P. Wang, Identification of Significant Genetic Variants via SLOPE, and its Extension to Group SLOPE; (Abstract/Program #1343F). Presented at the 65th Annual Meeting of The American Society of Human Genetics, October 9, 2015, Baltimore, MD.

- SOFTWARE**
- `mixed_models` – Fit statistical (linear) models with fixed and mixed (random) effects in Ruby. Project repository: [https://github.com/agisga/mixed\\_models](https://github.com/agisga/mixed_models)
  - `spitzy` – A toolbox of numerical differential equation solvers written in pure Ruby. Project repository: <https://github.com/agisga/spitzy>

**SKILLS**      *Computer skills:* R, Ruby, C++, Matlab, L<sup>A</sup>T<sub>E</sub>X, Linux.

*Language Knowledge:* Bilingual in German and Russian, fluent in English, basic knowledge of French.