Manuel Ariel Cáceres Reyes

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About me

I am a Doctoral Student at University of Helsinki. I work under the supervision of Alexandru I. Tomescu in the Graph Algorithms team, part of the Algorithmic Bioinformatics group at the Department of Computer Science. My research focuses on graph algorithms, algorithmic bioinformatics and compressed data structures. I consider myself a person responsible and devoted to his job. I enjoy learning new skills and I am always in search of new knowledge that I can apply to my job and life.

Education

University of Helsinki Helsinki, Finland

PhD. IN COMPUTER SCIENCE

2020 - Now

• Thesis Working Title: Safe, Complete and Parameterized Graph Algorithms for Bioinformatics.

University of Chile Santiago, Chile

MSc. in Computer Science, dual degree program with Computer Science Engineering.

2012 - 2019

Thesis Topic: Compressed Suffix Trees using Block Trees.
 Design and implementation of suffix trees, a data structure widely use in string processing. The structure was adapted to work in compressed space, specially for repetitive inputs. To do so I used and modified Block Trees, a modern data structure capable of capturing the repetitions.

Distinctions & Scholarships

THIRD PLACE IN THE Contest of Master Thesis in Computer Science of Latin America (CLEI 2020), ECUADOR 2020

Outstanding International Research, Department of Computer Science, University of Chile, Chile 2019

Scholarship to cover the 2nd semester of MSc. in Computer Science, University of Chile, Chile

2012-2015

MAXIMUM SCORE IN THE NATIONAL UNIVERSITY SELECTION TEST, MATHEMATICS, CHILE

PART OF THE HONOR LIST (TOP 6% OF STUDENTS), UNIVERSITY OF CHILE, CHILE

2011

2018

Publications

- Safety in multi-assembly via paths appearing in all path covers of a DAG, Manuel Cáceres, Brendan Mumey, Edin Husić, Romeo Rizzi, Massimo Cairo, Kristoffer Sahlin and Alexandru I. Tomescu. Submitted to the IEEE/ACM Transactions on Computational Biology and Bioinformatics (TCBB).
- Faster Repetition-Aware Compressed Suffix Trees based on Block Trees, Manuel Cáceres & Gonzalo Navarro. Submitted to the Information and Computation (I&C).
- A linear-time parameterized algorithm for computing the width of a DAG, Manuel Cáceres, Massimo Cairo, Brendan Mumey, Romeo Rizzi and Alexandru I. Tomescu. Submitted to the International Workshop on Graph-Theoretic Concepts in Computer Science (WG) 2021. A preliminary version can be found at https://arxiv.org/abs/2007.07575.
- Block Trees, Djamal Belazzougui, **Manuel Cáceres**, Travis Gagie, Pawel Gawrychowski, Juha Kärkkäinen, Gonzalo Navarro, Alberto Ordóñez, Simon J. Puglisi, and Yasuo Tabei. Accepted to the *Journal of Computer and System Sciences* (JCSS) 2021. A preliminary version can be found at https://me.ariel.computer/files/jcss20.pdf.
- Fast Indexes for Gapped Pattern Matching, Manuel Cáceres, Simon J. Puglisi and Bella Zhukova. Accepted to the International Conference on Current Trends in Theory and Practice of Informatics (SOFSEM) 2020 conference. A preliminary version can be found at https://arxiv.org/abs/2002.12662.
- Faster Repetition-Aware Compressed Suffix Trees based on Block Trees, Manuel Cáceres and Gonzalo Navarro. Accepted to the String Processing and Information Retrieval (SPIRE) 2019 conference. A preliminary version can be found at https://arxiv.org/abs/1902.03274.

Research visits & Presentations

- Talk "Safety in multi-assembly via paths appearing in all path covers of a DAG" in the Workshop Data Structures in Bioinformatics (DSB) 2021, Milano, Italy.
- Talk "Faster Minimum Path Covers of dense DAGs of small width" in the *Bioinformatics Research and Education Workshop* (BREW) 2020, Tartu, Estonia.
- Talk of my master thesis results in the X Workshop CeBiB Biotechnology, Metabolic Engineering, Bioinformatics, and Important Applications 2019, Santiago, Chile.
- Talk of my master thesis work to the research group: Database Laboratory (2018) of the University of A Coruña, Coruña, Spain.
- Research stay (2018, one month) at University of A Coruña, Coruña, Spain, granted by the BIRDS project. During the stay I worked with PhD. student Adrián Gómez on mt master thesis chapter: Compression of Repetitive Differential Arrays.
- Talk of my master thesis work to the research group: Genome Scale Algorithmics (2018) of the University of Helsinki, Helsinki, Finland.
- Research stay (2018, two months) at University of Helsinki, Helsinki, Finland granted by the BIRDS project. During the stay I worked with Associate Professor Simon Puglisi and PhD. student Bella Zhukova on the problem *Variable-Length Pattern Matching*.
- Poster presentation of my master thesis proposal on the VIII Workshop CeBiB Biotechnology, Metabolic Engineering, Bioinformatics, and Important Applications 2018, Santiago, Chile.

Teaching

University of Helsinki Helsinki, Finland

TEACHING ASSISTANT 2020-Now

- Depending on the course I prepared and presented model solutions to typical problems, graded exams, exercises and assignments, guided students with their projects or assignments, and took exams of the following graduate level courses:
 - Elements of Bioinformatics.
 - Design and Analysis of Algorithms.

University of Chile
Santiago, Chile

LECTURER 2015

• Undergraduate course Computer Tools for Engineering & Science, including text processors, spreadsheets, Matlab, R, Maple and basic Python. I prepared and taught classes and helped students with in-class exercises.

University of Chile Santiago, Chile

TEACHING ASSISTANT 2014 - 2018

- Depending on the course I prepared and presented model solutions to typical problems, graded exams, exercises and assignments, guided students with their projects or assignments, and took exams of the following graduate and undergraduate level courses:
 - Design and Analysis of Algorithms.
 - Algorithms and Data Structures.
 - Theory of computation.
 - Advanced Analysis of Algorithms.
 - Computer Tools for Engineering & Science.
 - Introduction to Programming.
 - Development of Software for Lego Robots.

Service

- Subreferee of the International Conference on Research in Computational Molecular Biology (RECOMB) 2021.
- Subreferee of the Data Compression Conference (DCC) 2021.
- Subreferee of the Workshop on Algorithms in Bioinformatics (WABI) 2020.
- Reviewer of the Bioinformatics Research and Education Workshop (BREW) 2020.
- Subreferee of the SIAM Symposium on Algorithm Engineering and Experiments (ALENEX) 2020.
- Subreferee of the *Theoretical Computer Science* (TCS) journal, in 2019.
- Subreferee of the European Symposia on Algorithms (ESA) 2018 track B.

Other Professional Experience

SimpliRoute Santiago, Chile

FULL-STACK DEVELOPER 2019

• Develop new features, optimize features, detect and fix bugs of the mobile and web apps "SimpliRoute" www.simpliroute.com, the number one software of logistic intelligence in Latin America.

Adereso Santiago, Chile

FULL-STACK DEVELOPER 2019

• Develop new features, optimize features, detect and fix bugs of the web app "PostCenter" www.postcenter.io, which centralizes and optimizes customer support of companies.