# Alberto Gil-de-la-Fuente

alberto.gilf@gmail.com GitHub Ceu Mass Mediator

## Research, education, code

#### Profile

Alberto Gil de la Fuente was born in Madrid in 1990. He studied Computer Engineering in San Pablo-CEU Universities. He graduated in 2013 obtaining the award to the best academic record. In 2016 he finished his MsC in Computer Engineering in Complutense University, in Madrid. During the MsC he did a period abroad of 6 months in Linköping University, Sweden. I started my PhD in Life and Science Technologies in San Pablo-CEU University in 2015. The project consists in an on-line tool to help with the metabolite identification. I also did a researching stay in the University of Alberta (Wishart Lab) for 6 months during 2018. You can see the results of the PhD development in the web page listed in the header. During his studies, Alberto combined both work and education. He worked as a database administrator in one of the biggest Spanish IT companies (Informatica El Corte Ingles: IECISA). He obtained some of the certifications from Oracle, DB2, MySQL or MongoDB, some of the main database technologies used nowadays. He is currently finishing the project and looking forward to continuing the research in the Metabolomics field, which he believes that it will bring the most significant advances in the systems biology field. At the same time, he recently started to enjoy the synergies of teaching. He thinks that the education is key in the formation of independent and autonomous people.

Particularly, he really enjoys High-Performance Computing. We should help the computers to make their work, and he has developed some projects to apply the High-Performance computing to life sciences.

Personally, he likes to create products that help people. He enjoys his free time doing sports, socializing with the family and friends and learning new technologies for a subsequent application solving real problems. he regularly does sports, he is part of the futsal team of the CEU-San Pablo University, and he actively participates in different event races, both running or biking. During his university studies he was football trainer of U-12 and U-15 teams, leading the players to become adults and having a lot of fun:)

#### Scientific Publications

Under Review. LAS: A Lipid Annotation Service Capable of Explaining the Annotations it

Generates Under Review

Fernández-López, M., Gil-de-la-Fuente, A., Godzien, J., Rupérez, F.J., Barbas, C. and Otero, A. Computational and Structural Biotechnology Journal. Impact Factor: 4.720 (Q1)

BioTransformer: a comprehensive computational tool for small molecule metabolism

prediction and metabolite identification

Djoumbou-Feunang, Y., Fiamoncini, J., Gil-de-la-Fuente, A., Greiner, R., Manach, C. and Wishart, D.S. Journal of cheminformatics. Impact Factor: 3.893 (Q1)

CEU Mass Mediator 3.0: a metabolite annotation tool

Gil-de-la-Fuente, A., Godzien, J., Saugar, S., Garcia-Carmona, R., Badran, H., Wishart, D.S., Barbas, C. and Otero, A. Journal of proteome research. Impact Factor: 3.950 (Q1)

Characterization and annotation of oxidized glycerophosphocholines for non-targeted

metabolomics with LC-QTOF-MS data

Gil-de-la-Fuente, A., Traldi, F., Siroka, J., Kretowski, A., Ciborowski, M., Otero, A., Barbas, C. and Godzien, J. Analytica Chimica Acta. Impact Factor: 5.123 (Q1) Under review

Knowledge-based metabolite annotation tool: CEU Mass Mediator

Gil-de-la-Fuente, A., Godzien, J., Fernández-López, M., Rupérez, F.J., Barbas, C. and Otero, A. Journal of Pharmaceutical and Biomedical Analysis Impact Factor: 2.831 (Q1)

Differentiating signals to make biological sense - a guide through databases for MS-based

non-targeted metabolomics

Gil-de-la-Fuente, A., Armitage, E.G., Otero, A. Barbas, C. and Godzien, J. Electrophoresis. Impact Factor: 2.744 (Q2)

GPU-based Acceleration of ECG Characterization using High-Order Hermite Polynomials

Gil-de-la-Fuente, A., Márquez, D.G., Caffarena, G., Iriarte, A. and Otero, A. Current Bioinformatics. Impact Factor: 0.770 (Q4)

### Book Chapters

Processing Metabolomics and Proteomics Data with Open Software. Chapter 11: Metabolite

annotation with CEU Mass Mediator

Gil-de-la-Fuente, Godzien, J., A., Otero and A. Barbas, C. Royal Society of Chemistry. Under review

Processing Metabolomics and Proteomics Data with Open Software. Chapter 12: Metabolite

annotation using in-silico generated compounds: MINE and BioTransformer

Gil-de-la-Fuente, A., Godzien, J., Otero and A. Barbas, C. Royal Society of Chemistry. Under review

Processing Metabolomics and Proteomics Data with Open Software. Chapter 3: Metabolomics

Wishart D.S., Godzien, J., Gil-de-la-Fuente, A., Otero and A. Barbas, C.

Under review

Data Analysis for Omic Sciences: Methods and Applications. Chapter 15: Metabolite

Annotation and Identification

Royal Society of Chemistry. Under review

Godzien, J., Gil-de-la-Fuente, A., Otero, A. and Barbas, C.

Elsevier. ISBN: 978-0-444-64044-4

2018

2019

2018

2019

2018

2017

Intornations	1 Conferences	
International	I Conferences  BioTransformer: a comprehensive computational tool for small molecule metabolism	
	BioTransformer: a comprehensive computational tool for small molecule metabolism prediction and metabolite identification  Discumbou-Feynage Y. Fiamoncini, J. Gil-de-la-Fuente, A. Greiner, R. Manach, C. and Wishart D.S.	2018
	Djoumbou-Feunang, Y., Fiamoncini, J., Gil-de-la-Fuente, A., Greiner, R., Manach, C. and Wishart D.S. The Second Annual Canadian Metabolomics Conference, Canmore	2018
	A fast and reliable spectral quality assessment in metabolomics studies  Gil de la Fuente, A., Traldi, F., Kowalczyk, T., Ciborowski, M., Otero, A., Barbas, C. and Godzien, J.  14th Annual Conference of the Metabolomics Society	
	CMM workshop: Improving Metabolit ID Approaches — Confidence and CASMI S., Sumner, L.W., Gil de la Fuente, A., Godzien, J. and Barbas, C. 14th Annual Conference of the Metabolomics Society	2018
	Ceu Mass Mediator: a knowledge-based tool for metabolite annotation  Gil de la Fuente, A., Godzien, J., Barbas, C. and Otero, A.  28th Pharmaceutical and Biomedical Analysis Conference	2017
	Low-power, Low-Latency Hermite Polynomial Characterization of Heartbeats Using a	
	Programmable Gate Array  Lakhotia, K., Caffarena, G., Gil-de-la-Fuente, A. Márquez, D.G., Otero, A. and Desai, M.P.  International Conference on Bioinformatics and Biomedical Engineering	2016
	Polynomial Characterization of Heartbeats with Graphics Processing Units Gil-de-la-Fuente, A., Caffarena, G., Marquez, D.G. and Otero, A. International Conference on Bioinformatics and Biomedical Engineering	2014
Official Educ	cation	
	San Pablo-CEU University PhD in Life Sciences and Technologies.	2015-current
	University of Alberta Researching period in Wishart Lab.	2018
	Linköping University  Erasmus scholarship.	2015-2016
	Complutense University  MsC in computer Engineering. GPA 8.69/10.	2014-2016
	Alfonso X el Sabio University  Degree in Information Systems Engineering (Grado en Ingeniería de Sistemas de Información). GPA 8.81/10.	2013-2014
	San Pablo-CEU University  Computer engineering (Ingeniería Informática de Sistemas). GPA 8.68/10.	2009-2012
Honors	Santander Scholarships Scholarship to help the mobility of researchers. Funding to do a period abroad in the University of Alberta	2018
	San Pablo-CEU university	2015
	FPI scholarship for PhD studies  Spanish ministry of Education and Science	2015
	Erasmus Scholarship  Alfonso X el Sabio University	2014
	Distinction in grade Thesis called "Heartbeats clustering using GPUs"  San Pablo-CEU University	2013
	Best Academic Record of the year	2013
	San Pablo-CEU University  Distinction in grade Thesis called "Characterization of heartbeats by Hermite Polynomials using GPUs"	
	Comunidad de Madrid  Excellent Academic Performance.  Award for students with a GPA higher than 9.25 during an academic year	2013
~		2015-curren
Career	PhD Student San Pablo-CEU University Development a software tool called Ceu Mass Mediator The title of the thesis project is design, validation and	

Development a software tool called Ceu Mass Mediator The title of the thesis project is design, validation and implementation of a software tool for metabolite identification. New strategies to aid the researchers during the metabolite annotation and identification have been developed. The development of these thesis has been published in JCR Q1 journals (see publications) and international conferences (see conferences). The framework is relatively used by the metabolomics community, having users in different countries like USA, Canada, China, Australia, Spain, France or Poland, among others.

Informatica El Corte Ingles (IECISA) The job in IECISA was my first experience related with IT. I started at the same time than the university since the first year. I was system administrator for different database engines such as Oracle, IBM DB2, MySQL, MongoDB or IBM InfoSphere Change Data Capture, among others. High performance architectures were installed and maintained by me (Oracle RAC 10g and 11g) as well as high availability architectures (Oracle DataGuard). We used both Oracle RDBMS and Oracle ASM. I was in charge of providing backup and recovery databases in case of disasters using Recovery Manager. Official Courses 2016 **Agilent Technologies** MS multiplatform Metabolomics Workshop 2015 Oracle Database 12c: New features for Administrators 2014 IESE Business School Leadership and Management Program 2014 NVIDIA CUDA and OpenAcc 2013 Oracle MySQL for Database Administrators Ed 3.1 2013 **NVIDIA** CUDA and OpenACC **IBM** DB2 IBM BootCamp. DB210 2010 Universidad a Distancia Española (UNED) Web programming using PHP, JavaScript and HTML 2010 Universidad a Distancia Española (UNED) .net programming 2009 Universidad a Distancia Española (UNED) Server Administration 2009 Universidad a Distancia Española (UNED) Java programming Certifications 2015 Oracle Database 11g Administrator Certified Professional SQL administration II (1Z0-053) 2015 Oracle Database 11g Administrator Certified Associate SQL Fundamentals, Administration I (1Z0-051 - 1Z0-052) 2013 Oracle MySQL Certified Professional. 1Z0-873 - 1Z0-874 MySQL 5 Database Administrator part I, II (1Z0-873 - 1Z0-874) 2013 Information Management DB2 10 Technichal Proffesional v3 2013 Certified Database Associate — DB2 10.1 Fundamentals IBM 000-N18 Adaptability Skills Multi-task Teaching **CUDA** Java C.C++ PrimeFaces **AWS** HTML SQL PL/SQL ORM LaTeX

Scholarship to integrate new knowledge into the expert system developed during the thesis in CEU-San Pablo University. I joined a new international environment and I obtained experimental knowledge that could be subsequently applied for the metabolites annotation and identification. I supported the younger researchers regarding the IT architecture and the

Research Assistant
University of Alberta

technological issues occurred.

Database Administrator

2018

2009-2015