Guillermo de Anda-Jáuregui

Systems Pharmacology, Network Biology

Curriculum Vitae

Current position Cátedra CONACYT associate researcher

Institution Consejo Nacional de Ciencia y Tecnología (CONACYT)

commissioned to:

Computational Genomics Division

Instituto Nacional de Medicina Genómica

(INMEGEN)

Periférico Sur 4109

Colonia Arenal Tepepan, Alcaldía Tlalpan,

Ciudad de México, 14610

e-mail gdeanda@inmegen.edu.mx

Github www.github.com/guillermodeandajauregui

Education

Ph.D. Biomedical Sciences

(Doctorado en Ciencias Biomédicas)

Project: Search for therapeutic targets associated to breast

cancer molecular subtypes

Universidad Nacional Autónoma de México

2011-2016

B.Sc. Pharmaceutical and Biological Chemistry

(Química Farmacéutica Biológica)

Facultad de Química

Universidad Nacional Autónoma de México

2005-2011

Work Experience

May 2019 - Today Cátedra CONACYT associate researcher,

Assigned to:

Computational Genomics Division

Instituto Nacional de Medicina Genómica

Nov. 2018 – Apr. 2019 Research associate

Computational Genomics Division

Instituto Nacional de Medicina Genómica

Jan. 2018 - Nov. 2018 Post-doctoral researcher

Project: *Cloud bioinformatics platform for the identification*

of genetic variants, an application for molecular

diagnostics and genomic research.

Instituto Nacional de Medicina Genómica

Nov. 2016 – Dec. 2017 Post-doctoral Research Fellow

Project: Systems Pharmacology

Junguk Hur Lab

School of Medicine and Health Scienes

University of North Dakota

Research Interests Systems pharmacology

Network biology Complex Systems

Biomedical applications of data science

Honors, awards, scholarships

2017 – to date Sistema Nacional de Investigadores (SNI)

CONACYT Candidate

2017 University of North Dakota Provost Award

109th Annual Meeting

North Dakota Academy of Sciences

2011 - 2016 Doctoral Grant

CONACYT No. 324432

2009 Student Exchange Scholarship

UNAM / UCLA

2006 - 2009 High Achievement Academic Program

(Programa de Alta Exigencia Académica)

UNAM

Teaching experience

2014 - 2016 Course: Biopharmacy Laboratory

Program: Pharmaceutical Biological Chemistry

Facultad de Química

UNAM

Publications *Equal Contribution

(Co-) corresponding author

2019 Guillermo de Anda-Jáuregui, Sergio A. Alcala-Corona,

Jesús Espinal-Enríquez, Enrique Hernandez-Lemus Functional and Transcriptional Connectivity of Communities in Breast Cancer Regulatory Networks

Applied Network Science

2019 Guillermo de Anda-Jáuregui, Jesús Espinal-Enríquez,

Santiago Sandoval-Motta, Enrique Hernández-Lemus

	A Boolean network approach to estrogen transcriptional regulation Complexity
2019	Guillermo de Anda-Jáuregui [#] Guideline for comparing functional enrichment of biological network modular structures Applied Network Science
2019	Guillermo de Anda-Jáuregui, Brett A. McGregor, Kai Guo, Junguk Hur A Network Pharmacology Approach for the Identification of Common Mechanisms of Drug-Induced Peripheral Neuropathy CPT: Pharmacometrics and Systems Pharmacology
2019	Guillermo de Anda-Jáuregui, Jesús Espinal-Enriquez, Enrique Hernández-Lemus Spatial Organization of the Gene Regulatory Program: An Information Theoretical Approach to Breast Cancer Transcriptomics Entropy
2019	Guillermo de Anda-Jáuregui, Kai Guo, Junguk Hur Network-Based Assessment of Adverse Drug Reaction Risk in Polypharmacy Using High-Throughput Screening Data International Journal of Molecular Sciences
2019	Guillermo de Anda-Jáuregui, Kai Guo, Brett A McGregor, Eva L Feldman, Junguk Hur Pathway crosstalk perturbation network modeling for identification of connectivity changes induced by diabetic neuropathy and pioglitazone BMC Systems Biology
2019	Guillermo de Anda-Jáuregui [#] , Jesus Espinal-Enriquez, Junguk Hur, Sergio Antonio Alcalá-Corona, Lena Ruiz-Azuara, Enrique Hernández-Lemus [#] Identification of Casiopeina II-gly secondary targets through a systems pharmacology approach Computational biology and chemistry
2018	Brett A McGregor*, Stephanie Eid*, Amy E Rumora, Benjamin Murdock, Kai Guo, Guillermo de Anda-Jáuregui, James E Porter, Eva L Feldman, Junguk Hur Conserved Transcriptional Signatures in Human and Murine Diabetic Peripheral Neuropathy Scientific reports
2018	Sergio Antonio Alcalá-Corona*, Guillermo de Anda- Jáuregui*, Jesús Espinal-Enriquez*, Hugo Tovar, Enrique Hernández-Lemus

	Network Modularity and Hierarchical Structure in Breast Cancer Molecular Subtypes International Conference on Complex Systems (Proceedings)
2018	Guillermo de Anda-Jáuregui, Kai Guo, Brett A McGregor, Junguk Hur Exploration of the anti-inflammatory drug space through network pharmacology: applications for drug repurposing Frontiers in physiology
2018	Sergio Antonio Alcalá-Corona*, Jesús Espinal-Enríquez*, Guillermo De Anda Jáuregui, Enrique Hernández-Lemus The Hierarchical Modular Structure of HER2+ Breast Cancer Network Frontiers in physiology
2018	Guillermo de Anda-Jáuregui*, Jesús Espinal-Enríquez*, Diana Drago-García, Enrique Hernández-Lemus Nonredundant, Highly Connected MicroRNAs Control Functionality in Breast Cancer Networks International Journal of Genomics
2017	Sergio Antonio Alcalá-Corona*, Guillermo de Anda- Jáuregui*, Jesús Espinal-Enríquez, Enrique Hernández- Lemus Network Modularity in Breast Cancer Molecular Subtypes Frontiers in physiology
2017	Jesus Espinal-Enriquez, Cristóbal Fresno, Guillermo Anda- Jáuregui, Enrique Hernandez-Lemus RNA-Seq based genome-wide analysis reveals loss of inter-chromosomal regulation in breast cancer Scientific Reports
2016	Guillermo de Anda-Jáuregui, Tadeo E Velázquez-Caldelas, Jesús Espinal-Enríquez, Enrique Hernández-Lemus. Transcriptional network architecture of breast cancer molecular subtypes Frontiers in physiology
2015	Guillermo de Anda-Jáuregui, Raúl A Mejía-Pedroza, Jesús Espinal-Enríquez, Enrique Hernández-Lemus Crosstalk events in the estrogen signaling pathway may affect tamoxifen efficacy in breast cancer molecular subtypes
2012	Computational biology and chemistry Helgi Jung Cook, Guillermo de Anda-Jáuregui, Kenneth Rubio Carrasco, Lourdes Mayet Cruz Comparación de perfiles de disolución: Impacto de los

criterios de diferentes agencias regulatorias en el cálculo de f2

Revista mexicana de ciencias farmacéuticas

Preprints

2018 Enrique Hernández-Lemus*, Jesús Espinal-Enríquez*,

Guillermo de Anda-Jáuregui.*

Probabilistic Multilayer Networks.

arXiv

2018 Diana García-Cortés, Guillermo de Anda-Jáuregui,

Cristobal Fresno, Enrique Hernandez-Lemus, Jesús Espinal-

Enriquez

Loss of trans regulation in breast cancer molecular

subtypes. bioRxiv

Conference presentations and Invited Talks

2018 Big Data: Networks and Genomics

Guillermo de Anda-Jáuregui & Jesús Espinal-Enríquez

Invited talk/workshop

Feria de Ciencias y Humanidades, UNAM

Ciudad de México, México

2018 Emergence of highly connected, non-redundant micro-

RNAs in breast cancer transcriptional networks and

their role in the control of biological functions

Guillermo de Anda-Jauregui, Jesus Espinal-Enriquez, Diana

Drago-Garcia and Enrique Hernandez-Lemus

Short talk

Conference n Complex Systems 2018

Thessaloniki, Greece

2018 Loss of trans-regulation in breast cancer

Jesús Espinal-Enriquez, Cristobal Fresno, Guillermo de

Anda-Jáuregui and Enrique Hernandez-Lemus

Short talk

Conference n Complex Systems 2018

Thessaloniki, Greece

8 Mexican corruption network: Using journalistic big data

to discover major political crimes

Jesús Espinal-Enriquez, Guillermo de Anda-Jáuregui, Rodrigo Garcia Herrera, Sergio Antonio Alcala-Corona,

Sofia Ramirez and Issa Luna Pla

Short talk

Conference n Complex Systems 2018

Thessaloniki, Greece

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2018

(presented by Jesús Espinal-Enríquez) 2018 **Large-Scale DNA Methylation Profiling of Human** Diabetic Peripheral Neuropathy in Subjects with Type 2 **Diabetes Mellitus.** Kai Guo, Sarah Elzinga, Stephanie Eid, Claudia Figueroa-Romero, Brett A Mcgregor, Guillermo De Anda-Jáuregui, Crystal Pacut, Eva L Feldman, Junguk Hur. Poster American Diabetes Association 78 Scientific Sessions Orlando, Florida, USA (presented by Kai Guo) 2018 Biomedical network analysis: drugs and cancer Short talk useR! MEX 2018 Ciudad de México, México 2018 **Network Pharmacology: from Big Data to Precision** Medicine Invited talk Institutional Seminar, Instituto Nacional de Medicina Genómica Ciudad de México, México 2017 The use of a drug-gene perturbation network for the study of drug side effects: The case of drug-induced peripheral neuropathy Guillermo de Anda-Jáuregui, Kai Guo, Brett McGregor, Junguk Hur Short talk 1st Latin American Conference on Complex Networks Puebla, Puebla, México Changes in pathway connectivity induced by disease and 2017 therapeutic treatment: the case of diabetic neuropathy and pioglitazone Guillermo de Anda-Jáuregui, Kai Guo, Brett McGregor, Junguk Hur Short talk Conference on Complex Systems 2017 Cancún, Quintana Roo, México 2017 Network pharmacology exploration of the antiinflammatory drug space Guillermo de Anda-Jáuregui, Kai Guo, Brett McGregor, Junguk Hur Poster 25th Conference on Intelligent Systems for Molecular Biology and 16th European Conference on Computational

Biology

Prague, Czech Republic

2017

Pathway-level effects of pioglitazone on neuropathy in the BKS-db/db Mouse Model of Type 2 diabetes

Guillermo de Anda-Jáuregui, Kai Guo, Brett McGregor,

Junguk Hur

Poster

American Diabetes Association 77 Scientific Sessions

San Diego, California, USA

2017

Using systems pharmacology to identify common mechanisms of drug-induced peripheral neuropathy Guillermo de Anda-Jáuregui, Kai Guo, Brett McGregor,

Junguk Hur

Short talk

109th Annual Meeting, North Dakota Academy of Sciences

Grand Forks, North Dakota, USA

2015

Effects of pathway crosstalk in the therapeutic efficacy

of antiestrogenic treatment for breast cancer

Invited talk

Symposium: Advances and perspectives in medical

chemistry

Taxco, Guerrero, México

2015

Crosstalk events in the estrogen signaling pathway that affect tamoxifen efficacy in different subtypes of breast

cancer

Invited talk

Institutional Seminar, Instituto Nacional de Medicina

Genómica

Ciudad de México, México

Programming languages & Computational Skills

R

General scripting

Network Analysis:

Igraph

Data Analysis:

Tidyverse data.table

General Bioinformatics:

BioMart annotation

Microarray preprocessing pipeline RNAseq preprocessing pipeline

limma

Pathway analysis:

GSEA

GAGE

Over-representation analysis

Pathifier

Python General scripting

Data Analysis: Pandas

Network Analysis:

NetworkX

Linux Bash scripting

Command line environment

Desktop environment

HTCondor distributed computing

Windows Desktop environment

General tools LaTeX typesetting

Microsoft Office Suite LibreOffice Suite

GIMP (image processing)
Inkscape (image processing)
ImageMagick (image processing)

Git version control

Languages

Spanish, native

English, fluent

Cambridge Certificate in Advanced English (CEFR Level 1)

French, basic