

# Giuliano Netto Flores Cruz

Pharmacist Bioinformatician

Vancouver, British Columbia - Canada

[cruz.gnf@gmail.com](mailto:cruz.gnf@gmail.com)

[Personal web page](#)

## EDUCATION

### **The University of British Columbia (UBC), MSc. in Bioinformatics**

SEPT 2021 — AUG 2023 (Vancouver, BC - Canada)

Advisor: Dr. Keegan Korthauer

### **Federal University of Rio Grande do Sul (UFRGS), Title of Pharmacist**

FEB 2012 — DEC 2018 (Porto Alegre, RS - Brazil)

Advisors: MSc. Paulo Saraiva & Dr. Alexandre Fuentefria

## EXPERIENCE

### **BC Children's Hospital Research Institute, Research Assitant**

SEP 2021 — PRESENT (Vancouver, BC - Canada)

- RA at [Keegan Korthauer's lab](#).
- Development of statistical methodology and computational tools for the analysis of high-dimensional genomics data.

### **BiomeHub, Bioinformatician**

SEP 2019 — PRESENT (Florianópolis, SC - Brazil)

- Focus on hospital environments and human health.
- Research: development of statistical models to estimate bacterial colony-forming units (CFU) using next-generation sequencing data.

### **Neoprospecta, Bioinformatician**

JUN 2018 — AUG 2019 (Florianópolis, SC - Brazil)

- Develop, implement, and maintain bioinformatics pipelines.
- Microbiome data analysis consultancy for academia and industry.
- Research: hospital microbiome and infection control.

### **Immunology Lab (School of Pharmacy - UFRGS), Research and Teaching Assistant**

JAN 2016 — MAY 2018 (Porto Alegre, RS - Brazil)

- TA: Immunology, Clinical Immunology, and Immunodiagnostics.
- RA: idiopathic pulmonary fibrosis using gene expression data.

### **Rohwer Lab (San Diego State University), Research Assistant**

MAY 2015 — AUG 2015 (San Diego, CA - USA)

- Internship as part of the *Science without Borders* program.
- RA: laboratory protocols for endotoxin-free bacteriophage stock preparation.

## SKILLS

\* Superscripts index related publications.

### **R & Python**

- Tidyverse, Bioconductor, brms, rstan
- Django, common python data science libraries

### **Bayesian modeling<sup>1, 2</sup>**

### **Machine Learning<sup>3</sup>**

### **Data types and wet lab:**

- Microbiome<sup>1, 6</sup>
- Cell image<sup>3</sup>
- Gene expression<sup>4</sup>
- Immunology<sup>4</sup>
- Microbiology<sup>5, 2</sup>

### **PostgreSQL, BigQuery**

### **Django REST API & Flask**

### **Google Cloud Platform**

### **Unix & shell scripting**

## COMPLEMENTARY EDUCATION

### **MITx, Micromasters in Statistics and Data Science**

MAY 2019 — DEC 2020 (Boston, MA - USA)

Delivered online by MIT faculty through the EdX platform. Comprising advanced, in-depth courses on probability theory, statistics, machine learning, and data analysis – plus a capstone (proctored exam) for each discipline.

### **Harvardx, Professional Certificate in Data Analysis for the Life Sciences**

FEB 2018 — JUL 2018 (Cambridge, MA - USA)

Delivered online by Harvard faculty through the EdX platform. Comprising four courses on statistical computing and high-dimensional data analysis.

## AWARDS

### **Concord University, Science without Borders (CAPES)**

JUL 2014 — MAY 2015 (Athens, WV - USA)

Student exchange program sponsored by the Coordenação de Aperfeiçoamento de Pessoal de Nível Superior (CAPES), a federal foundation within the Brazilian Ministry of Education.

## SELECTED PUBLICATIONS

\* Indices relate to superscripts from the skills section. See the complete list at [Google Scholar](https://scholar.google.com/).

- 1) **Cruz GNF**<sup>+</sup>, Christoff AP<sup>+</sup>, Oliveira LFV. 2020. “Equivolumetric protocol generates library sizes proportional to total microbial load in next-generation sequencing”. *BioRxiv*. <https://doi.org/10.1101/2020.02.03.932301>
- 2) Christoff AP<sup>+</sup>, **Cruz GNF**<sup>+</sup> et al. 2020. “Swab pooling for large-scale RT-qPCR screening of SARS-CoV-2”. *MedRxiv*. <https://doi.org/10.1101/2020.09.03.20187732>
- 3) Lopes W<sup>+</sup>, **Cruz GNF**<sup>+</sup>, Rodrigues ML, Vainstein MH, Kmetzsch L, Staats CC, Vainstein MH, Schrank A. 2020. “Scanning electron microscopy and machine learning reveal heterogeneity in capsular morphotypes of the human pathogen *Cryptococcus* spp.”. *Scientific Reports*. <https://doi.org/10.1038/s41598-020-59276-w>
- 4) **Cruz GNF**. 2018. “Genomic analysis of macrophage gene signatures during idiopathic pulmonary fibrosis development”. *UFRGS*. Final undergraduate monograph. <http://hdl.handle.net/10183/195681>
- 5) Bonilla N, Rojas MI, **Cruz GNF**, Hung S, Rohwer F, Barr JJ. 2016. “Phage on tap—a quick and efficient protocol for the preparation of bacteriophage laboratory stocks”. *PeerJ*. <https://doi.org/10.7717/peerj.2261>
- 6) Christoff AP, Sereia AFR, **Cruz GNF**, et al. 2020. “One-year cross-sectional study in adult and neonatal intensive care units reveals the bacterial and antimicrobial resistance genes profiles in patients and hospital surfaces”. *Plos One*. <https://doi.org/10.1371/journal.pone.0234127>
- 7) Saraiva PJ, **Cruz GNF**. “Diagnóstico e tratamento da sífilis”. In: Associação Brasileira de Ciências Farmacêuticas; Pereira LRL, Farias RF, Castro MS, organizadores. PROFARMA: Programa de Atualização em Ciências Farmacêuticas: da Assistência Farmacêutica à Farmácia Clínica. Ciclo 1. Porto Alegre: Artmed Panamericana; 2019. p. 129–64. (Sistema de Educação Continuada a Distância, v.3).

<sup>+</sup>These authors contributed equally to the corresponding work.