# PIERO PALACIOS BERNUY

# ■ PROFESSIONAL EXPERIENCE

Current

Bioinformatician intern

#### Rayca Precision

Q Lima, PE

2022

- · Develop & Deploy NGS data processing pipelines
- · Analyze large amounts of information to discover trends and patterns

2019 2018 Professional practitioner

### International Potato Center

**Q** Lima. PE

- · Management and recovery of the collection in vitro "Low viability of sweet potato"
- · Responsible for training in botanical and physiological knowledge in order to strengthen the methodology of in vitro propagation.
- · Establishment of working methodologies and flowcharts for the management of the collection in vitro "Low viability of sweet potato".

2019

Consultant

#### International Potato Center

Q Lima, PE

- · Research, development, and improvement of an in vitro medium for the propagation of sweet potato compared with media used by the International Potato Centre
- Experience in the area SO Conserving Biodiversity for the Future and procedures ISO17025.
- · Proposal of a new medium of cultivation to generate a better growth of the plants of sweet potato

## ♣■ EDUCATION

2018 2012 B. Sc. Biotechnology

### La Molina National Agrarian University

Q Lima. PE

- Thesis: Impact of high temperature on coffee (Coffea arabica) transcriptome
- · Development of bioinformatics pipelines in HPC (High performance computing) for functional analysis of coffee transcriptome



CONTACT



Hi! My name is Piero, and I am a Biotechnologist from Peru PE 🐂.

I have especializations in data analysis and bioinformatics from Harvard University and MIT.

I have a blog about bioinformatics aimed at people who are enthusiastic about this field

**TFACHING** 

Rstudio and

Languages

Spanish: Native

English: Intermediate

R package pagedown and datadrivency.

Code available on G GitHub.



resume is available here.

### COURSES 2022 Probability - The Science of Uncertainty and Data **♀** Remote MITx (Massachusetts Institute of Technology) ntroduction to Computer Science and Programming Using 2021 Python • Remote MITx (Massachusetts Institute of Technology) Data analysis for Genomics 2020 • Remote **HARVARD**x · Professional Program Analysis of high-dimensional data 2020 **♀** Remote HARVARDX Statistical Inference and Modeling for High Performance 2020 Experiments • Remote **HARVARD**X Statistical analysis in bioinformatics 2019 • Remote USMx (University System of Maryland)

### Selected activities:

### **PROGRAMMING SKILLS**

- **R** R
- ▲ tidyverse
- M RMarkdown / Quarto
- GitHub
- ▼ Bioconductor
- ☐ HPC
- Python
- ⚠ Linux