PIERO PALACIOS BERNUY

■ PROFESSIONAL EXPERIENCE

Current

Bioinformatician scientist

Remote



- · Provide statistical computational tools for biologically based activities like genetic analysis, measurement of gene expression, and also gene sets analysis.
- · Analyze large molecular datasets like Next Generation Sequencing (NGS) data, microarray data, genomic sequence data, and also proteomics data for clinical or basic research purposes.
- · Develop new software applications or customize existing applications to meet specific project needs.

Current 2023

Consultancy for Statistical Support - Seed System / Agrobiodiversity / Timeline datasets

International Potato Center

Remote

- · Conduct statistical analysis as agreed with the team.
- · Generate graphs, tables, and visuals to best represent results.
- · Contribute to science article development.

2023

2022

Bioinformatician intern

Rayca Precision

• Remote

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

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

2019 2018 Professional practitioner

International Potato Center

O 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".



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

TEACHING

Rstudio and

LANGUAGES

Spanish: Native

English: Intermediate

R package pagedown and datadrivency.

Code available on

GitHub.



resume is available here.

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



COURSES

Machine Learning with Python-From Linear Models to Deep 2023 Learning

MITx (Massachusetts Institute of Technology)

• Remote

· Certificate

2022 Fundamentals of Statistics

MITx (Massachusetts Institute of Technology)

• Remote

· Certificate

Probability - The Science of Uncertainty and Data 2022

MITx (Massachusetts Institute of Technology)

• Remote

· Certificate

Introduction to Computer Science and Programming Using 2021

MITx (Massachusetts Institute of Technology)

• Remote

· Certificate

Data analysis for Genomics 2020

HARVARDx

Remote

- · Professional Program
- · Certificate

Advanced Bioconductor 2020

HARVARDx

• Remote

· Certificate

2020

Case studies in functional genomics

HARVARDx

Remote

· Certificate

Selected activities:

PROGRAMMING SKILLS

₽ R

tidyverse

M+ RMarkdown / Quarto

GitHub

▼ Bioconductor

■ HPC

Python

Introduction to Bioconductor 2020 **♀** Remote HARVARDX · Certificate High-dimensional data analysis 2020 **♀** Remote HARVARDx · Certificate Statistical Inference and Modeling for High Performance 2020 Experiments **♀** Remote HARVARDx · Certificate Statistical analysis in bioinformatics 2019 **♀** Remote USMx (University System of Maryland) · Certificate