# BABU ROSHAN PADMANABHAN

#### **Bioinformatics Scientist**

University Hospitals, Cardiovascular Research Institute Wolstein Research Building, Cleveland Ohio 44106

## **EMPLOYMENT**

May 2020 -

#### Bioinformatics Scientist

University Hospitals

Oleveland, Ohio

o Working on epigenomics, single-cell genomics and transcriptomics.

2016-2020

#### Postdoctoral Bioinformatics Fellow

Cleveland Clinic

Oleveland, Ohio

#### **Accomplishments:**

- o Used machine learning and genomic approaches to understand the role of microbiome in cancer.
- o Identified highly correlated multi-omic signatures between metabolomics and microbial datasets.
- o Worked on microbial associations in patients with autism spectrum disorder, vasculitis, and breast cancer.
- o Utilized ChIP-Seq and RNA-Seq methods to analyze the role of PTEN in transcription.
- o GermLine variant analysis of Children, Adolescents, and Young Adults (C-AYA) with SolidTumor.

2015 – 2016

#### Bioinformatics Engineer

Institut des Maladies Métaboliques et Cardiovasculaires

◆ Toulouse, France

#### Accomplishments:

- o Developed pipelines for Metagenomic (16s Amplicon as well as Shotgun) and Bulk RNASeq datasets.
- o Analyzed several Metagenomic datasets.
- o Applied machine learning algorithms and Network analysis on clinical and metagenomic datasets.

2006 – 2011

#### Senior Research Fellow

Hyderabad Central University

Hyderabad, India

#### Accomplishments:

- o Poster presentation accepted in the European Conference on Computational Biology at Ghent, Belgium, 2010.
- o Learned embryonic stem cell culture techniques

2004 -2006

#### Bioinformatician

Jawaharlal Nehru Centre for Advanced Scientific Research

Pangalore, India

#### Accomplishments:

- o Assembled, Annotated and did comparative genomics of several HIV genomes.
- o Developed lab website.



### CONTACT INFO

- github.com/rosaak
- **J** +1 216-280-7550

#### **SKILLS**

- Experience in design and development of next generation sequencing (NGS) data processing pipelines
- Experience in analyzing Metagenomics, Transcriptomics, Epigenomics datasets
- Experience in multi-omics integration
- Experience in statistical analysis, machine learning
- Proficient in R, Bash and Python
- Good communication skills and a track record of successful collaborations with cross-functional teams

### **EDUCATION**

2011- 2014		Ph.D. Aix-Marseille University	<b>♥</b> Marseilles, France
2003- 2004	•	Advanced Post Graduate Diploma in Bioinformatics Jawaharlal Nehru University	♦ New Delhi, India
2000- 2002	•	M.Sc. in Marine Biotechnology Goa University	<b>♀</b> Goa, India
1997- 2000	•	B.Sc. in Biotechnology Kerala University	<b>♥</b> Kerala, India
		HONORS AND SEMINARS / WORKSHOPS ATTENDED	
2019	•	Artificial Intelligence in Oncology : Advancements and Policy Case Western Reserve University	<b>♀</b> Cleveland, OH
2019	•	International PTEN Symposium From Patient-Centered Research to Clinical Care	<b>♀</b> Cleveland, OH
2013		Galaxy Training Galaxy Community Conference	Oslo, Norway
2013		3-days workshop on Scientific Writing Practice Aix-Marseille University	
2006	•	EMBO Workshop Center for Cellular and Molecular Biology	<b>♀</b> Hyderabad, India
2010	•	Molecular evolution, structure and functional divergence of Lipoxygenase gene to Poster Presentation	•
2002	•	9th European Conference on Computational Biology  National workshop on Bioinformatics: Principles, Algorithms and Tools	<b>♀</b> Ghent, Belgium
		C-DIT and TBGRI	<b>♥</b> Kerala, India
2001,2004		National Eligibility Test (NET) for Junior Research Fellowship (JRF) The Council of Scientific & Industrial Research (CSIR) and University Grants Commission	<b>♀</b> India

## ▼ DOMAIN SPECIFIC KNOWLEDGE

### Programming Related

- ٠R
- Python
- Perl
- JavaScript / HTML / CSS
- Bash, Awk, Sed
- DataBase: MySQL, SQLite, Mongodb
- Web: Django, Dash (plotly), Shiny
- Linux Administration
- Experience working in HPC, Docker, Snakemake, Jupyter Notebook

#### Proficient in

- Data wrangling and visualization: pandas, numpy, dplyr, ggplot2, matplotlib and seaborn
- Machine Learning libraries : scikit-learn, caret, randomforest, xgboost, t-sne
- · Bioconductor libraries for NGS: Rsamtools, Rsubreads, DESeq2, edger, phyloseq, maftools, limma
- Analysis of Next Generation Sequence Data
- · Microbial whole genome assembly
- Variant detection related [GATK, maftools, XHMM]
- Transcriptomics [STAR,Kallisto,Sleuth,DESeq2]
- ChIPSeg / ATAC-Seg [macs2, HOMER, diffbind, bicycle]
- · Amplicon and Shotgun Metagenomics, Metabolomics and Pan-genomics [Qiime, Micca, mothur, MOCAT, Kraken, mixOmics1
- Integrative systems biology
- · Setting up and maintaining local database for Handling NGS Data, Galaxy local server etc.

## 🖵 SOFTWARE, DATABASES AND DATA ANALYSIS

### **Data Analysis**

· Comprehensive germline genomic profiles of children, adolescents and young adults with solid tumors. github link

#### **Data Visualization**

 Sankey Plots Visualization showing drug-target network analysis in children, adolescents, and young adults (C-AYA) with solid tumors github link

2019

#### MAGI

· A pipeline for whole genome comparison, github link

#### **ARG-ANNOT** (Antibiotic Resistance Gene-ANNOTation)

· A database to discover antibiotic resistance genes in bacterial genomes. link 2013



### PROFESSIONAL SERVICE

#### Reviewer for journals

- New Microbes and New Infections (Elsevier)
- Scientific Reports (Nature)
- · Journal of Applied Microbiology
- · Letters in Applied Microbiology
- Cancer Research (AACR)

## ♣☐ EXTRA CURRICULAR

#### Passionate about

- Photography
- Sketching
- Data Visualization
- Online Learning

### PUBLICATIONS

 Germline EGFR variants are over-represented in adolescents and young adults (AYA) with adrenocortical carcinoma

Human Molecular Genetics., 2020 Dec 16, ddaa268 PMID: 33326033

Sara Akhavanfard, Lamis Yehia, Roshan Padmanabhan, Jordan P. Reynolds, Ying Ni and Charis Eng

Exposure to Air Pollution Disrupts Circadian Rhythm through Alterations in Chromatin Dynamics iScience., Volume 23, Issue 11, 101728 PMID: 33241196

Rengasamy Palanivel, Vinesh Vinayachandran, Shyam Biswal, Jeffrey A.Deiuliis, **Roshan, Padmanabhan**, Bongsoo Park, Roopesh SinghGangwar, Jared C.Durieux, Elaine AnnEbreo Cara, Lopa Das, Graham Bevan, Zahi A.Fayad, Ahmed Tawakol, Mukesh K.Jain, Sujata Rao, Sanjay Rajagopalan

2020 • Metabolic stress regulates genome-wide transcription in a PTEN-dependent manner

Human Molecular Genetics., 2020, Vol. 29, No. 16 PMID: 32744308

Ata Abbas, Roshan Padmanabhan, Charis Eng

Bacteriome and mycobiome and bacteriome-mycobiome interactions in head and neck squamous cell carcinoma

Oncotarget. 2020 Jun 23; 11(25): 2375-2386. PMID: 32637029

Elizabeth Shay, Naseer Sangwan, Roshan Padmanabhan, Scott Lundy, Brian Burkey, and Charis Eng

2020 Comprehensive germline genomic profiles of children, adolescents and young adults with solid tumors.

Nature Communications. 2020 May 5; 11(1):2206. PMID: 32371905

Sara Akhavanfard, Roshan Padmanabhan, Lamis Yehia, Feixiong Cheng, Charis Eng.

2020 • Copy Number Variation and Clinical Outcomes in Patients with Germline PTEN Mutations.

JAMA Network Open. 2020;3(1):e1920415. PMID: 32003824

Lamis Yehia, Marilyn Seyfi, Lisa-Marie Niestroj, Roshan Padmanabhan, Ying Ni, Thomas Frazier, Dennis Lal, Charis Eng.

Resveratrol-mediated glycemic regulation is blunted by curcumin and is associated to modulation of gut microbiota.

The Journal of Nutritional Biochemistry. 2019 Oct;72:108218 PMID: 31473511

Navin Sreng, Serge Champion, Jean-Charles Martin, Saber Khelaifia, Jeffrey E Christensen, **Roshan Padmanabhan**, Vincent Azalbert, Vincent Blasco-Baque, Pascale Loubieres, Laurent Pechere, Jean-François Landrier, Rémy Burcelin, Eric Sérée.

PTEN modulates gene transcription by redistributing genome-wide RNA polymerase II occupancy. Hum Mol Genet. 2019 Sep 1;28(17):2826-2834. PMID: 31127935

Ata Abbas, Roshan Padmanabhan, Todd Romigh, Charis Eng.

2019

# Microbiomes of Inflammatory Thoracic Aortic Aneurysms Due to Giant Cell Arteritis and Clinically Isolated Aortitis Differ From Those of Non-Inflammatory Aneurysms.

Pathogens and Immunity. 2019 Mar 15;4(1):105-123. PMID: 30993253

Getz TM\*, Hoffman GS\*, **Padmanabhan R**, Villa-Forte A, Funchain P, Roselli EE, Blackstone E, Johnston D, Petterson G, Soltesz, Svensson LG, Calabrese LH, Clifford AH, Eng C.

#### 2019 • The Microbiome of Temporal Arteries.

Pathogens and Immunity. 2019;4(1):61-78. PMID: 30993251

Hoffman GS, Getz TM, **Padmanabhan R**, Villa-Forte A, Clifford AH, Funchain P, Sankunny M, Perry JD, Blandford A, Kosmorsky G, Lystad L, Calabrese LH, Eng C.

### 2019 • Microbiome in PTEN hamartoma tumor syndrome.

Endocr Relat Cancer. 2017, PMID: 29233840

Victoria Byrd, Ted M Getz, Roshan Padmanabhan, Hans Arora and Charis Eng.

#### 2017 • Transfer of dysbiotic gut microbiota has beneficial effects on host liver metabolism.

Mol Syst Biol. 2017 Mar 16;13(3):921, PMID: 28302863

Simon Nicolas, Vincent Blasco-Baque, Audren Fournel, Jerome Gilleron, Pascale Klopp, Aurelie Waget, Franck Ceppo, Alysson Marlin, **Roshan Padmanabhan**, Jason Siacovoni, François Tercé, Patrice D Cani, Jean-François Tanti, Remy Burcelin, Claude Knauf, Mireille Cormont, Matteo Serino.

# Microbial Culturomics to Map Halophilic Bacterium in Human Gut: Genome Sequence and Description of Oceanobacillus Jeddahense sp. nov.

OMICS, 2016, 20 (4), 248-258. PMID: 27093109

Saber Khelaifia, Jean-Christophe Lagier, Fehmida Bibi, Esam Ibraheem Azhar, Olivier Croce, **Roshan Padmanabhan**, Asif Ahmad Jiman-Fatani, Muhammad Yasir, Catherine Robert, Claudia Andrieu, Pierre-Edouard Fournier, Didier Raoult.

#### Periodontal dysbiosis linked to periodontitis is associated with cardiometabolic adaptation to highfat diet in mice.

Am J Physiol Gastrointest Liver Physiol. 2016,1;310(11). PMID: 27033119

Maxime Branchereau, François Reichardt, Pascale Loubieres, Pauline Marck, Aurélie Waget, Vincent Azalbert, André Colom, **Roshan Padmanabhan**, Jason Iacovoni, Anaïs Giry, François Tercé, Christophe Heymes, Remy Burcelin, Matteo Serino, Vincent Blasco-Baque.

#### 2015 Genome sequence and description of Pantoea septica strain FF5.

Stand Genomic Sci. 2015; 10: 103. PMID: 26568786

2015

Cheikh Ibrahima Lo, **Roshan Padhmanabhan**, Oleg Mediannikov, Thi Tien Nguyen, Didier Raoult, Pierre-Edouard Fournier, and Florence Fenollar.

# High-quality genome sequence and description of Bacillus dielmoensis strain FF4(T) sp. nov. Stand Genomic Sci. 2015 22; 10:41. PMID: 26221422

Cheikh Ibrahima Lo, **Roshan Padhmanabhan**, Oleg Mediannikov, Jérôme Terras, Catherine Robert, Ngor Faye, Didier Raoult, Pierre-Edouard Fournier, and Florence Fenollar.

## Noncontiguous finished genome sequence and description of Necropsobacter massiliensis sp. nov. New Microbes New Infect. 2015 Nov; 8: 41–50. PMID: 26587237

C.I. Lo, **R. Padhamanabhan**, B. Fall, B. Sambe-Ba, O. Mediannikov, T.-T. Nguyen, E. Prudent, N. Faye, Wade, D. Raoult, P.-E. Fournier, and F. Fenollar.

# • ARG-ANNOT (Antibiotic Resistance Gene-ANNOTation), a new bioinformatic tool to discover antibiotic resistance genes in bacterial genomes.

Antimicrob Agents Chemother, 2014;58(1):212-20. PMID: 24145532

Gupta SK, **Padmanabhan BR**, Diene SM, Lopez- Rojas R, Kempf M, Landraud L, Rolain JM.

Genome Sequence of Borrelia crocidurae Strain 03-02, a Clinical Isolate from Senegal. 2014 Genome Announc. 2014 Nov 6;2(6):e01150-14. PMID: 25377717 Aurélien Fotso Fotso, Oleg Mediannikov, Roshan Padmanabhan, Catherine Robert, Pierre-Edouard Fournier, Didier Raoult, Michel Drancourt. High quality draft genome sequence and description of Occidentia massiliensis gen. nov., sp. nov., 2014 a new member of the family Rickettsiaceae. Stand Genomic Sci. 2014 Dec 8;9:9. PMID: 25780502 Oleg Mediannikov, Thi-Thien Nguyen, Lesley Bell-Sakyi, Roshan Padmanabhan, Pierre-Edouard Fournier, and Didier Raoult. Genome sequence and description of Corynebacterium ihumii sp. nov. 2014 Stand. Genomic Sci. 2014 Mar 15;9(3):1128-43. PMID: 25197488 Roshan Padmanabhan, Grégory Dubourg, Jean-Christophe Lagier, Carine Couderc, Caroline Michelle, Didier Raoult, Pierre- Edouard Fournier. Non-contiguous finished genome sequence and description of Collinsella massiliensis sp. nov. 2014 Stand. Genomic Sci. 2014 Mar 15;9(3):1144-58. PMID: 25197489 Roshan Padhmanabhan, Gregory Dubourg, Jean-Christophe Lagier, Thi-Thien Nguyen, Carine Couderc, Morgane Rossi-Tamisier, Aurelia Caputo, Didier Raoult, Pierre-Edouard Fournier. **Draft Genome Sequence of Necropsobacter rosorum Strain P709T** 2014 Genome Announc. 2014 Oct 9;2(5):e00913-14. PMID: 25301642 Roshan Padmanabhan, Catherine Robert, Florence Fenollar, Didier Raoulta, Pierre-Edouard Fournier. Non-contiguous-finished genome sequence and description of Paenibacillus gorillae sp. nov. 2014 Stand. Genomic Sci. 2014 Mar 15;9(3):1031-45 PMID: 25197481 Mamadou Bhoy e Keita, Roshan Padhmanabhan, Aurelia Caputo, Catherine Robert, Eric Delaporte, Didier Raoult, Pierre-Edouard Fournier, Fadi Bittar. A polyphasic strategy incorporating genomic data for the taxonomic description of novel bacterial 2014 species. [Review] Int J Syst Evol Microbiol. 2014 Feb; 64(Pt 2):384-391 PMID: 24505076 Dhamodharan Ramasamy, Ajay Kumar Mishra, Jean-Christophe Lagier, Roshan Padhmanabhan, Morgane Rossi, Erwin Sentausa, Didier Raoult and Pierre-Edouard Fournier. Non-contiguous-finished genome sequence and description of Gorillabacterium massiliensis gen. 2014 nov, sp. nov., a new member of the Family Paenibacillaceae. Stand. Genomic Sci. 2014 Mar 15;9(3):807-20. PMID: 25197465 Mamadou Bhoye Keita, Roshan Padhmanabhan, Aurelia Caputo, Catherine Robert, Eric Delaporte, Didier Raoult, Pierre-

Genomics and Metagenomics in Medical Microbiology. [Review]

Journal of Microbiological Methods 2013 Dec;95(3):415-24 PMID: 24200711

Edouard Fournier, Fadi Bittar.

2013

Roshan Padmanabhan, Ajay Kumar Mishra, Didier Raoult, Pierre-Edouard Fournier.

Non-contiguous finished genome sequence and description of Megasphaera massiliensis sp. nov Stand. Genomic Sci. 2013 Aug 7;8(3):525-38. PMID: 24501636

Roshan Padmanabhan, Jean-Christophe Lagier, Nicole Prisca Makaya Dangui, Caroline Michelle, Carine Couderc, Didier Raoult, Pierre-Edouard Fournier.

Non-contiguous finished genome sequence and description of Clostridium dakarense sp. nov Stand. Genomic Sci. 2013 Sep 30;9(1):14-27. PMID: 24501642

Cheikh Ibrahima Lo, Ajay Kumar Mishra, **Roshan Padhmanabhan**, Bissoume Samb, Ba, Amy Gassama Sow, Catherine Robert, Carine Couderc, Ngor Faye, Didier Raoult, Pierre-Edouard Fournier, Florence Fenollar.

Exceptional molecular and coreceptor-requirement properties of molecular clones isolated from an Human Immunodeficiency Virus Type-1 subtype C infection.

Retrovirology 2008 Mar 7;5:25. PMID: 18328091

2008

Dash P K, Siddappa N B, Mangaiarkarasi A, Mahendarkar A, **Roshan P B**, Anand K K, Mahadevan A, Shankar S K, Satishchandra P, Prasad V R, Ranga U.