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Education & Training

1996-1999 – B.S. Biomedical Science, Rio de Janeiro University, Brazil.

2001-2002 – M.S. Parasitology Biology – Infectious diseases, Oswaldo Cruz Foundation, Brazil.

2003-2009 – Ph.D. Infectious Diseases Clinical Research – Immunogenetics, Oswaldo Cruz Foundation, Brazil.

2009- 2014 – Post doctorate in Vaccine Development, Texas Biomedical Research Institute, San Antonio, USA.

Research Focus

I am an Immunologist with more than 15 years of experience in the clinical research of infectious diseases, predominantly in *Mycobacterium tuberculosis*. My research focused on the immunopathogenesis of tuberculosis, and I was interested in the identification and characterization of susceptibility genes for this disease with the ultimate objective of gaining an insight into the biological pathways involved in disease pathogenesis. I have extensive experience in several immunological, pathological and microbiological technologies and most recently in high-throughput genomic technologies and applying these to help understand the genetic underpinnings of disease. As a Co-I on a several projects, I have continued to significantly enhance my expertise in several other immunological and molecular techniques predominantly using high-throughput omics.

In 2006, I immigrated to the United States to work on the development of a DNA vaccine against tuberculosis in non-human primates at Texas Biomedical Research Institute. I assisted the PI with compiling an NIH R01 grant entitled "TB vaccine development in Nonhuman Primate Model" which was awarded \$3.8 million. I oversaw and worked on all laboratory aspects of this project. I was able to gain insight and expertise in several project aspects: technological

experience, project management and data analysis.

Recently, my research efforts and expertise are in the molecular and functional aspects of gene identification for complex diseases. I am involved in several projects where I manage the molecular components, being responsible for the phenotyping and sequencing of sample from several complex diseases such as cardiovascular disease, type 2 diabetes, bipolar disease and Alzheimer's disease. I have also played a major role in a few large-scale human genetic projects in the STDIOI. I generated quality data for several studies trying to understand the influence of host genetics in susceptibility to complex diseases. I am also working with epigenetic influences on Alzheimer's disease. My role as a PI on this project was being responsible for the overall project including bisulfite conversion of the DNA, running the methylation arrays and performing the statistical analyses.

For the last two and a half years, I have been working with exposome, lipidomics, metabolomics and metallomics. I am focused on the directly measured pollutome component of the Major Depressive disorder (MDD) exposome. Our goal is to use a novel genomic approach to improve the search for environmental determinants of MDD to provide a more complete conceptualization of the illness, facilitating treatment and prevention. We will use our extended Mexican American pedigrees, high-throughput pollutomics/metallomics, and novel statistical analyses for the possible identification and understanding of the pollutant-influenced pathways and mechanisms contributing to MDD risk in Mexican Americans.

Publications

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2. Albuquerque MC, Aleixo AL, Benchimol EI, **Leandro AC**, das Neves LB, Vicente RT, Bonecini-Almeida Mda G, Amendoeira MR (2009) The IFN-gamma +874T/A gene polymorphism is associated with retinochoroiditis toxoplasmosis susceptibility. *Mem Inst Oswaldo Cruz* 104 (3): 451 - 455. doi: 10.1590/s0074-02762009000300009. PMID: 19547871.
3. Martineau AR, **Leandro AC***, Anderson ST, Newton SM, Wilkinson KA, Nicol MP, Pienaar SM, Skolimowska KH, Rocha MA, Rolla VC, Levin M, Davidson RN, Bremner SA, Griffiths CJ, Eley BS, Bonecini-Almeida MG, Wilkinson RJ (2010) Association between Gc genotype and susceptibility to TB is dependent on vitamin D status. *Eur Respir J* 35(5): 1106 - 1112. PMCID: PMC2864196 * **co-first author**. doi: 10.1183/09031936.00087009. PMID: 19797128; PMCID: PMC2864196.
4. **Leandro AC**, Rocha MA, Lamoglia-Souza A, VandeBerg JL, Rolla VC, Bonecini-Almeida Mda G (2013) No association of IFNG+874T/A SNP and NOS2A-954G/C

SNP variants with nitric oxide radical serum levels or susceptibility to tuberculosis in a Brazilian population subset. *Biomed Res Int* 2013: 901740. doi: 10.1155/2013/901740. PMID: 24024215; PMCID: PMC3759278.

5. Cepeda M, Salas M, Folwarczny J, **Leandro AC**, Hodara VL, de la Garza MA, Dick EJ Jr, Owston M, Armitige LY, Gauduin MC (2013) Establishment of a neonatal rhesus macaque model to study Mycobacterium tuberculosis infection. *Tuberculosis (Edinb)* 93 Suppl: S51 - S59. doi: 10.1016/S1472-9792(13)70011-8. PMID: 24388650; PMCID: PMC4051704.
6. Knowles EEM, Curran JE, Göring HHH, Mathias SR, Mollon J, Rodrigue A, Olvera RL, **Leandro AC**, Duggirala R, Almasy L, Blangero J, Glahn DC. Family-based analyses reveal novel genetic overlap between interleukin-8 and risk for suicide attempt. *Brain Behav Immun*. 2019 Aug; 80:292-299. doi: 10.1016/j.bbi.2019.04.004. PMID: 30953777; PMCID: PMC7168352.
7. Rodrigue AL, Knowles EEM, Mollon J, Mathias SR, Koenis M, Peralta JM, **Leandro AC**, Fox PT, Sprooten E, Kochunov P, Olvera RL, Duggirala R, Almasy L, Curran JE, Blangero J, Glahn DC (nd) Evidence for pleiotropy between human cerebral white matter microstructure and inflammation. *Hum Brain Mapp*. 2019 Oct 1;40(14):4180-4191. doi: 10.1002/hbm.24694. PMID: 31187567; PMCID: PMC6707845.
8. Blackburn NB, Michael LF, Meikle PJ, Peralta JM, Mosior M, McAhren S, Bui HH, Bellinger M, Giles C, Kumar S, **Leandro AC**, Almeida M, Weir JM, Nestel PJ, Simes J, Sullivan DR, Tonkin AM, Mahaney MC, Dyer TD, Almasy L, VandeBerg JL, Williams-Blangero S, Glahn DC, Duggirala R, Kowala M, Blangero J, Curran JE (nd) A rare variant in DEGS1 with major effects on the de novo ceramide synthesis pathway. *J Lipid Res*. 2019 Sep;60(9):1630-1639. doi: 10.1194/jlr.P094433. PMID: 31227640; PMCID: PMC6718439.
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