

# Fábio Cassarotti Parronchi Navarro

---

New Haven - CT - United States of America

fabio.navarro@yale.edu – +1 (475) 227 5975

## Education

- **Universidade de São Paulo**  
Ph.D. in sciences (biochemistry) (2011-current)  
Advisor: Anamaria A. Camargo, Ph.D.  
Co-advisor: Pedro A. F. Galante, Ph.D.
- **Universidade Federal de São Carlos**  
Computer Engineering (2004-2009)

## Honours and Awards

- Best poster award - X-meeting 2012 (Genomics).
- Prêmio Viagem (2013/1) - Biochemistry Program.
- Best poster award - X-meeting 2013 (Genomics).
- AWS in Education Research Grant (2014).
- EMBO Conference Travel Grant (2014).

## Research Interest

- mRNA retroduplication on primates, human populations and cancer genomes.
- Large-scale analysis of structural variations.
- Pseudogene neofunctionalization.

## Research Experience

- **Yale University**  
Post-doctoral associate at Mark Gerstein Lab (2014-Current)
- **Hospital Sírio-Libanês**  
Bioinformatics Lab. (<http://www.bioinfo.mochsl.org.br>) (2011-2014)
- **Ludwig Institute of Cancer Research**  
Compbio Lab. Technician (2009-2010)
- **Embrapa - CNPDIA**  
Developer (2005-2007)

## Teaching Experience

- **Universidade de São Paulo**
    - Molecular Biology for Medicine (2012)  
Teaching assistant
    - Computational Molecular Biology for Chemistry (2011)  
Teaching assistant
-

## Computational Biology Experience

- **Developing computational pipelines**
  - Programming proficiency: Perl\*, Python, C++ and Java
- **Cancer Genomics**
  - Single Nucleotide Variation analysis
  - Structural Variation analysis
- **Second Generation Sequencing**
  - SNP/SNV calling
  - Structural variation calling\*
  - RNA-seq analysis
    - Differential expression and Pathway enrichment
  - 1KGP - [DBvar](#)
- **Comparative genomics**
  - Primate/Murine orthology
  - Repetitive elements
  - Purifying selection
- **Web development and Lab. Infrastructure**
  - Basic CakePHP and MySQL
  - Design and architecture of [Multiple User Equipament](#) cluster

## Peer-reviewed Publications

- The PsychENCODE project.  
**Psych ENCODE consortium**, Nature Neuroscience, 2015.  
doi:10.1038/nn.4156
- A genome-wide landscape of retrocopied protein-coding genes in primates genomes.  
**Navarro FCP**, Galante PAF., Genome Biology and Evolution, 2015.  
doi:10.1093/gbe/evv142
- Gene Copy-Number Polymorphism Caused by Retrotransposition in Humans.  
Schrider DR\*, **Navarro FCP\***, et al. Plos Genetics, 2013.  
doi:10.1371/journal.pgen.1003242
- RCPedia: a database of retrocopied genes.  
**Navarro FCP**, Galante PAF, Bioinformatics, 2013.  
doi:10.1093/bioinformatics/btt104
- SPLOOCE: a new portal for the analysis of human splicing variants.  
Kroll JE, Galante PAF, Ohara DT, **Navarro FCP**, et al., RNA Biology, 2012  
doi: 10.4161/rna.22182
- Distinct patterns of somatic alterations in a lymphoblastoid and a tumor genome derived from the same individual.  
Galante PAF, Parmigiani RB, Zhao Q, Caballero OL, Souza JE, **Navarro FCP**, et al., Nucleic Acid Research, 2011.  
doi:10.1093/nar/gkr221

## Work in progress

- Evolutionary and functional impact of a primate specific retrocopy into the Transferrin locus.  
**Navarro FCP**, Urllass AS, Smith VS, Camargo AA, Galante PAF, MacGillivray RTA.