

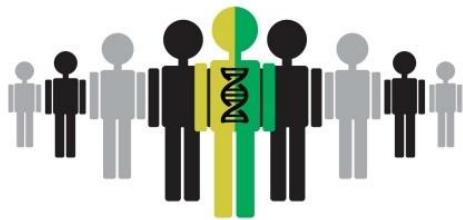
## **BIPMed – New Projects and Advances**

**[www.bipmed.org](http://www.bipmed.org)**



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**Project Manager**  
**University of Campinas – UNICAMP**

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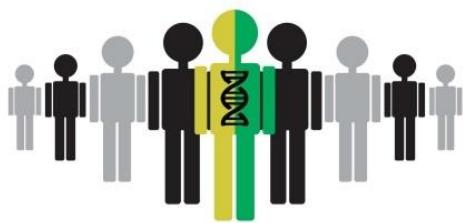


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# Databases



- Lieden Open Variation Database 3.0
- Web-based gene sequence variation database
- API enables simple queries or listing of gene and variant data
- Freely available tool for Gene-centered collection and display of DNA variations.



## List of public LOVD installations



If you are looking for a specific gene database, please also check our [list of LSDBs](#).

Please note that the latest available build is always installed on our [Leiden server](#). We offer free use of this LOVD installation interested, e.g. to maintain/start a LSDB. [Contact us](#) for more information.

Show only LOVDs with gene symbol:  Show 

You can also [download this list](#) to see the complete list of genes in the databases.

**In total: 3,913,937 variants (2,943,353 unique) in 424,351 individuals in 85 LOVD installations.**

[http://databases.lovd.nl/whole\\_genome/](http://databases.lovd.nl/whole_genome/)

LOVD 3.0-20a

22002 genes

Whole genome datasets

A1BG, A1BG-AS1, A1CF, A2M, A2M-AS1, A2ML1, A2MP1, A4GALT, A4GNT, A...

<http://bipmed.igm.unicamp.br/>

LOVD 3.0-17

20930 genes

BIPMed WES

A1BG, A1BG-AS1, A1CF, A2M, A2M-AS1, A2ML1, A2MP1, A3GALT2, A4GALT...

<https://databases.lovd.nl/shared/>

LOVD 3.0-20b

229 // genes

Shared database

A1BG, A1BG-AS1, A1CF, A2LD1, A2M, A2M-AS1, A2ML1, A2MP1, A3GALT2,...

<http://proteomics.bio21.unimelb.edu.au/lovd/>

LOVD 3.0-07

14772 genes

LOVD - Leiden Open Variation Database

A1BG, A1BG-AS1, A1CF, A2M, A2ML1, A4GALT, A4GNT, AAAS, AACSD, AACSP...

<http://bipmed.igm.unicamp.br/cfa/>

LOVD 3.0-20

41 genes

Craniofacial anomalies

APOC2, APOC4-APOC2, AXIN2, BCL3, BMP4, CLPTM1, DVL2, ERBB2, FGF22...

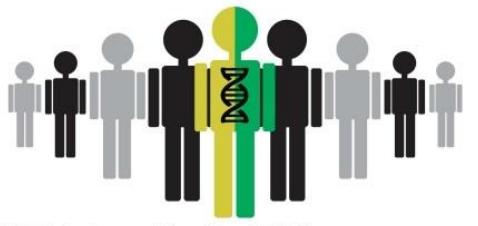
<https://ab-openlab.csir.res.in/mitolsdb/>

LOVD 2.0-35

3 / genes

MitoLSDB

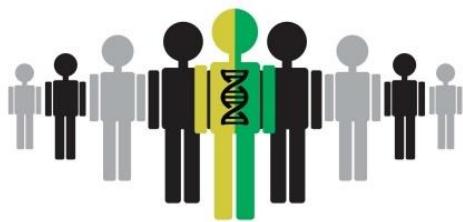
MTATP6, MTATP8, MTCO1, MTCO2, MTCO3, MTCYB, MTND1, MTND2, MTND3, M...



# Mirrors

## BIPMed has 4 installations

- Main site: IQ – Unicamp - Campinas
- Mirror 1: FCM – Unicamp - Campinas
- Mirror 2: USP – Ribeirão Preto
- Mirror 3: UNIFESP – São Paulo



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# BIPMed

## Our Products:

- Genomic databases:
  - BIPMed-WES-db: 256 individuals **REFERENCE POPULATION**
  - BIPMed-Array-db: 264 individuals **REFERENCE POPULATION**
    - DISEASE SPECIFIC DATABASES
- BIPMed Beacon
- GA4GH R client
- .....

# Disease/Phenotype specific projects

- Epilepsy (BRAINN, ILAE-ALADE) – EE
  - DOOSE SYNDROME
  - DRAVET SYNDROME
- Craniofacial Anomalies – Dr. Vera Lucia Gil Da Silva Lopes
  - Cleft lip and palate (BCFP)
- BRCA Challenge (GA4GH, HVP) – Dr. Patricia Prolla and Edenir Palmero



**BRCA  
EXCHANGE**

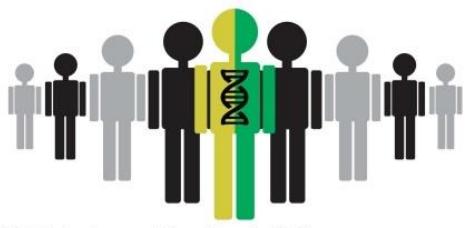
# BRCA Challenge

- Dr. Patricia Prolla, Dr. Edenir Palmero and Dr. Maria Aparecida Folgueira
- Supported by GA4GH, LOVD and the Human Variome Project
  - Understanding of the genetic basis of breast cancer, ovarian cancer and other diseases
  - Pooling data on BRCA1/2 genetic variants and corresponding clinical data from around the world.
- The BRCA Challenge on BIPMed
  - **75 variants in BRCA1**
  - **87 variants in BRCA2**



# Craniofacial Anomalies

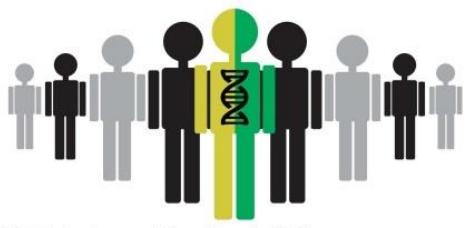
- Dr. Vera Lucia Gil Da Silva Lopes
- 253 variantes in 41 genes
- 358 Healthy individuals with no history of oral clefting in the past three generations
- 157 patients with Non-syndromic Cleft Lip and Palate



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# Under development

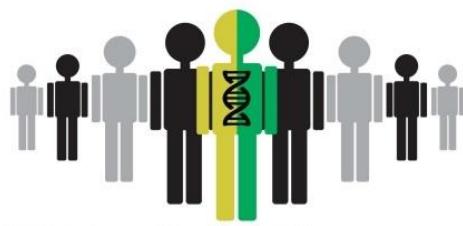
- Copy number databases;
- Craniofacial anomalies SNP Array
  - Affymetrix Genome-Wide Human SNP Array 6.0;
- Stroke (BRAINN, ISGC-Latin American Initiative);
- Pathogenic hemoglobins – Global Globin (HVP) – Dr. Monica Melo;
- ApoE Challenge (BRAINN, ABN) – Dr. Marcio Balthazar;



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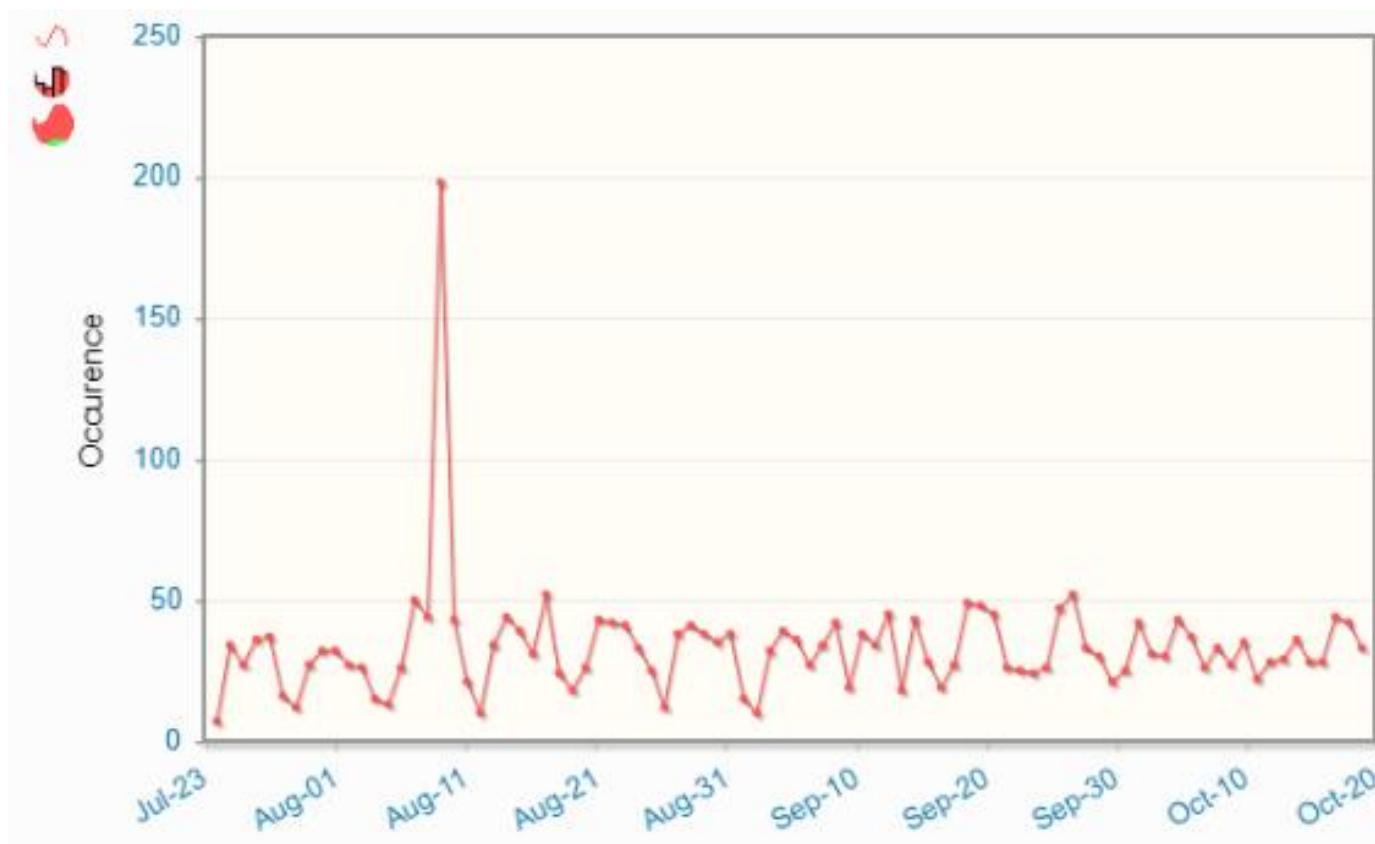
# Under development

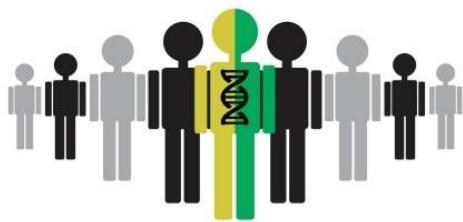
- 40 reference individuals born in Ribeirão Preto region with more than 65 years old – Prof. Wilson Silva Junior
  - Whole Exome Sequencing;
  - Affymetrix Genome-Wide Human SNP Array 6.0
- 40 reference individual born in Barretos region – Prof. Vinicius Vasquez
  - Whole Exome Sequencing;
  - Affymetrix Genome-Wide Human SNP Array 6.0



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# Access Statistics

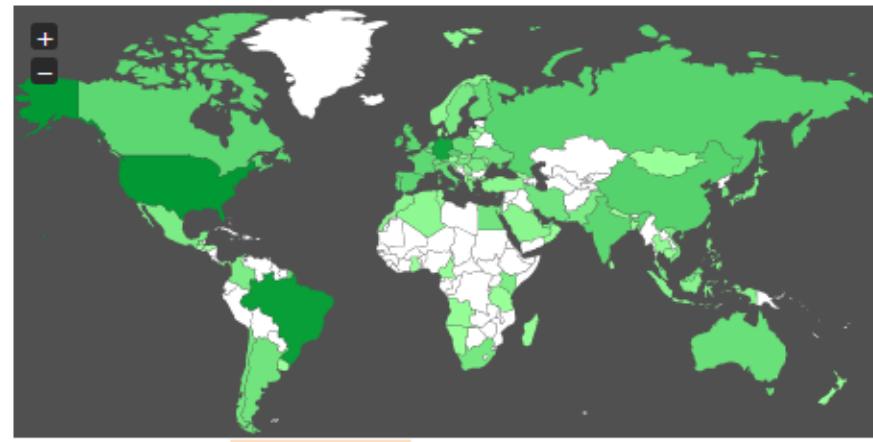
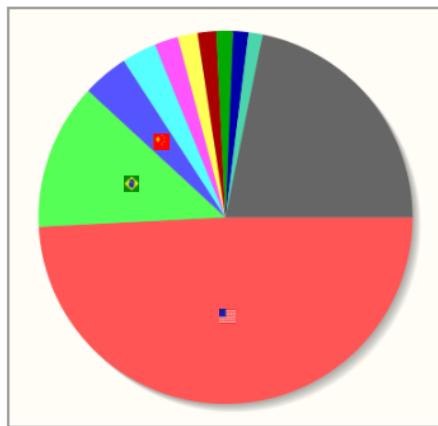




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# Access Statistics

Countries		
United States - 🇺🇸	889	49.1 %
Brazil - 🇧🇷	229	12.6 %
China - 🇨🇳	72	3.9 %
India - 🇮🇳	54	2.9 %
Russian Federation - 🇷🇺	37	2.0 %
France - 🇫🇷	31	1.7 %
Great Britain - 🇬🇧	29	1.6 %
Germany - 🇩🇪	26	1.4 %
Philippines - 🇵🇭	24	1.3 %
Canada - 🇨🇦	22	1.2 %
Others	394	21.8 %



Requests    Accesses    Hosts    Traffic

	Hosts	Percentage	Requests	Percentage	Accesses	Percentage	Traffic (Mb)	Percentage
Europe	305	16.9 %	16786	32.4 %	12162	47.4 %	232.7	45.2 %
South-America	252	13.9 %	14412	27.8 %	7307	28.5 %	97.9	19.0 %
North-America	918	50.8 %	18132	35.0 %	5350	20.9 %	152.1	29.5 %
Central-America	7	0.4 %	7	0.0 %	7	0.0 %	0.3	0.1 %
Africa	40	2.2 %	245	0.5 %	70	0.3 %	4.0	0.8 %
Oceania	17	0.9 %	193	0.4 %	69	0.3 %	1.9	0.4 %
Asia	267	14.8 %	2067	4.0 %	1686	2.7 %	26.0	5.0 %
Misc	1	0.1 %	1	0.0 %	1	0.0 %	0.0	0.0 %

# LatinGen



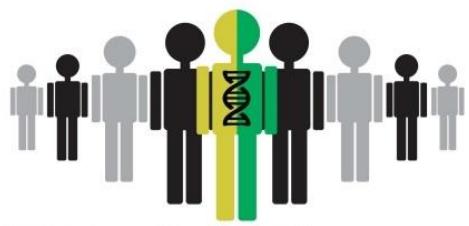
- Development of the portal [latingen.org](http://latingen.org)
- Disseminate the idea of federated databases
- Act as a centralizer of genomic information in Latin America
- Provide assistance for other research groups

[contact@latingen.org](mailto:contact@latingen.org)

# LatinGen



- Future projects:
  - Develop integration of participants databases
    - Projects using the LOVD interface provide easy integration
    - Projects with other database structure can be challenging
  - Provide bioinformatics training in Latin America



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# Acknowledgment





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