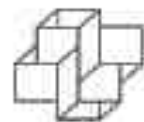
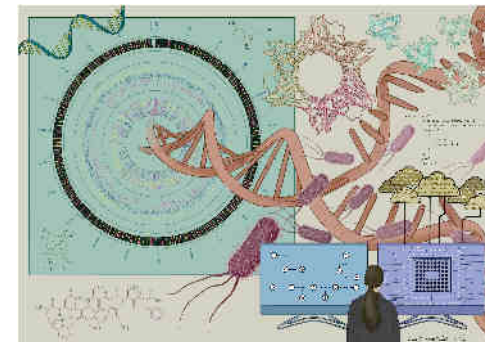


# OMICS AND BIOINFORMATICS: STRATEGIC VISION OF THE PAST AND CHALLENGES FOR THE FUTURE

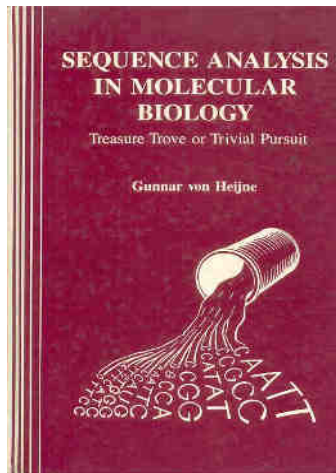
Ana Tereza Vasconcelos

Laboratório Nacional de Computação  
Científica LNCC/MCTI



Laboratório  
Nacional de  
Computação  
Científica

# 1988 - Biofísica UFRJ/LNCC



*G. von Heijne*  
(1987)



Prof. Darcy F. de Almeida - UFRJ

Bioinformatics : research *in silico*

(A. Danchin)


54. **VASCONCELOS, A. T. R.**; ALMEIDA, Darcy Fontoura de ; COIMBRA, C. A. . Prokaryote DNA damage-inducible SOS system genes detected by nucleotide sequence analyses. In: Academia Brasileira de Ciências, 1994, Rio de Janeiro. An. Academia Brasil G, v. 66. p. 127.



10:20 - 11:00	<p><b>Conferência: Genomics – a Brazilian vocation</b></p> <p>Andrew Simpson, Origen Biotecnologia, São Paulo, SP</p>
11:00 - 12:30	<p><b>Painel: Discussões com empreendedores do ramo da genômica e bioinformática</b></p> <p>Moderador: Maria Sueli Soares Felipe, UNB, Brasília, DF          Gonçalo Amarante Guimarães Pereira, Unicamp, Campinas, SP          João Bosco Oliveira, Genonilla Diagnostico          Luiz Felipe Valtor de Oliveira, BiomeHub e Neoprospecta          Michael Butterfield, Centro de Tecnologia Canavieira, Piracicaba, SP          Paulo Arruda, Unicamp, Campinas, SP</p>
12:30 - 14:30	<p><b>Almoço</b></p>
14:30 - 15:00	<p><b>Conferência: Visão geral de CT&amp;I no Brasil</b></p> <p>Ilduê do Castro Moreira, Presidente da SBPC, São Paulo, SP</p>
15:30 - 17:00	<p><b>Painel: Discussão com líderes de agências e ministros</b></p> <p>Moderador: Augusto Schrank, UFRGS, Porto Alegre, RS          Marcelo Marcos Moraes - Representante do MCTIC          Marcela Caloguesi de Sa - Coordenadora COBRG - CNPq          Marie-Anne Van Sluys - Representante da FAPESP          André Marco de Oliveira Gomes - Representante da FAPERJ          Paulo Sérgio Lacerda Beltrão - Diretor da FAPEMIG, Belo Horizonte, MG          Marco Aurélio Krieger - Vice-Presidente da Fiocruz          Carlos Frederico Martins Menck - Coordenador da CAPES/CB 1, USP São Paulo, SP</p>

	14 Dezembro	15 Dezembro	16 Dezembro
9:30 - 11:00			<p><b>Painel: Tópicos Avançados em genômica e bioinformática</b></p> <p>Moderador: Emmanuel Dias-Neto, AC Camargo Câncer Center, São Paulo, SP          Ândrea Kely Campos Ribeiro dos Santos, UFPA, Belém, PA          Anete Pereira Souza, Unicamp, Campinas, SP          Arnaldo Zaha, UFRGS, Porto Alegre, RS          Célia Maria de Almeida Soares, UFG, Goiânia, GO          Roberto Lins, IAM FIOCRUZ, Recife, PE          Lygia da Veiga Pereira, USP, São Paulo, SP</p>
9:30 - 11:00			<p><b>Painel: Recursos humanos em genômica e bioinformática</b></p> <p>Moderadora: Glória Regina Franco, UFMG, Belo Horizonte, MG          Ana Carolina Ramos Guimarães, Fiocruz, Rio de Janeiro, RJ          Ana Tereza Ribeiro de Vasconcelos, LNCC, Petrópolis, RJ          Glória Regina Franco, UFMG, Belo Horizonte, MG          João Paulo Matos Santos Lima, UFRN, Natal, RN          Alan Mitchell Durham, IME/USP, São Paulo, SP          Pedro Geraldo Pascutti, UFRJ, Rio de Janeiro, RJ</p>
11:00 - 12:30			<p><b>Painel: Bioinformática: Oportunidades e desafios</b></p> <p>Moderador: Sandro José de Souza, UFRN          Helder Takashi Imoto Nakaya, USP          João Paulo Kitajima, Mendelica Análise Genômica, São Paulo, SP          Paulo Costa Carvalho, Fiocruz, Curitiba, PR          Richard Charles Garrat, USP/São Carlos</p>
12:30 - 14:00			<p><b>Almoço</b></p>
14:00 - 14:30			<p><b>Conferência: Percepção em relação à bioinformática e genômica</b></p> <p>Luisa Massarani, Fiocruz, Rio de Janeiro, RJ</p>
14:30 - 15:00			<p><b>Conferência: A Revolução Genômica na Medicina</b></p> <p>Sérgio Danilo Junho Pena, UFMG, Belo Horizonte, MG</p>
15:00 - 16:30			<p><b>Painel: O impacto da genômica no cenário de C&amp;T no Brasil</b></p> <p>Moderador: Ana Tereza Ribeiro de Vasconcelos, LNCC, Petrópolis, RJ          Ana Maria Benko Iseppon, UFPE, Recife, PE          Marie-Anne Van Sluys, USP São Paulo, SP          Santuzia Maria Ribeiro Teixeira, UFMG, Belo Horizonte, MG</p>
			<p><b>Painel: Futuro da genômica e bioinformática</b></p> <p>Moderador: Lucymara F. Agnez Lima, UFRN, Natal, RN          Eduardo Emrich Soares, Biomina, Belo Horizonte, MG          Elizabeth Pacheco Batista Fontes, UFV, Viçosa, MG          Guilherme Corrêa de Oliveira, ITV, Belém, PA          Renato Santana de Aguiar, UFMG, Belo Horizonte, MG</p>
			<p><b>Encerramento</b></p>

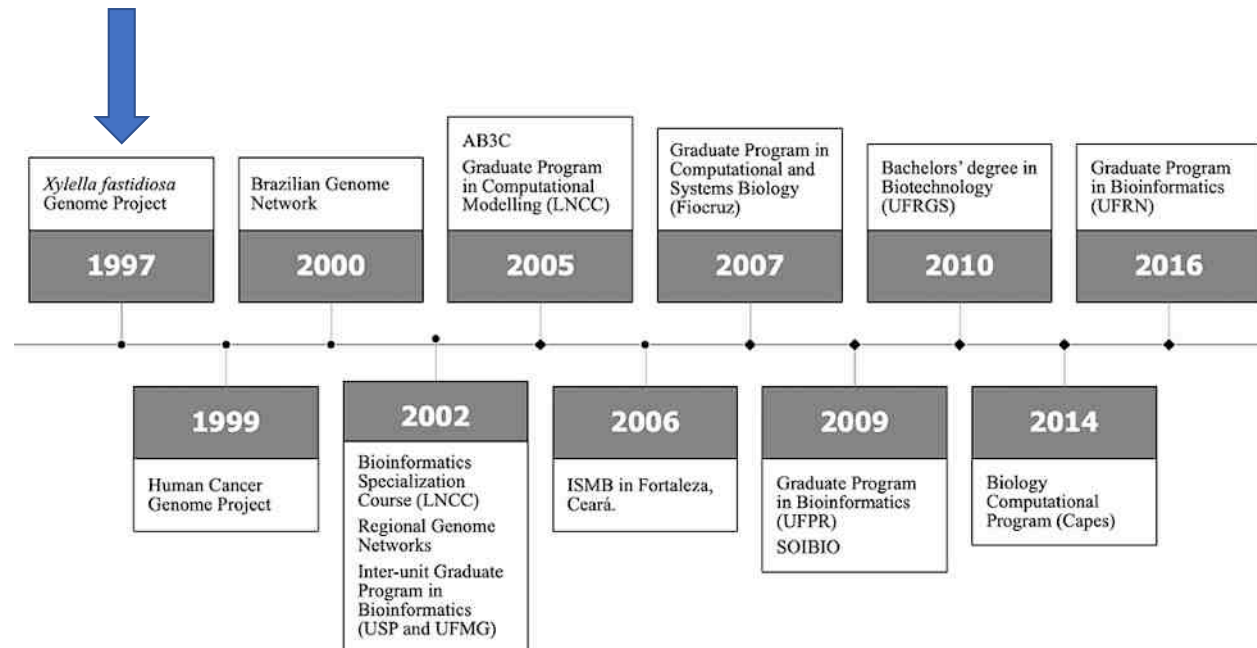
## The past, present and future of genomics and bioinformatics: A survey of Brazilian scientists

Mariana Rocha<sup>1</sup> , Luisa Massarani<sup>2</sup>, Sandro José de Souza<sup>3,4,5</sup> and Ana Tereza R. de Vasconcelos<sup>6</sup>



Sub-questions (SQ)	Objectives
SQ1: What is the profile of the researchers currently working in the field?	To collect and evaluate details about genomics and bioinformatics professionals considering demographic features, educational background, and professional experience
SQ 2: What are the perceptions about the present and future? How does it confront the perceptions about the area in other countries?	To collect and evaluate genomics and bioinformatics professionals' thoughts on the current and future situation of the field both in Brazil and abroad.
SQ 3: What are the milestones achieved in the area, and what are the expert wishes for the future?	To collect and evaluate information about the country's main achievements and what should be considered for future work.

<https://doi.org/10.1590/1678-4685-GMB-2021-0354>

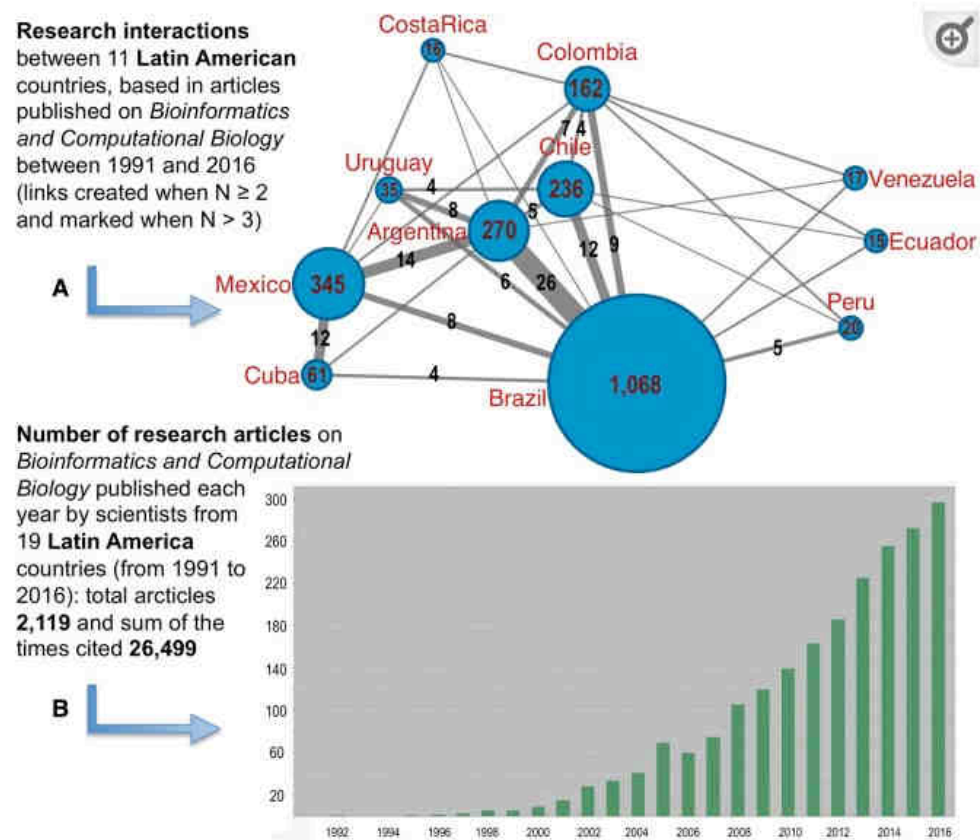


<https://doi.org/10.1590/1678-4685-GMB-2021-0354>





Figure 2

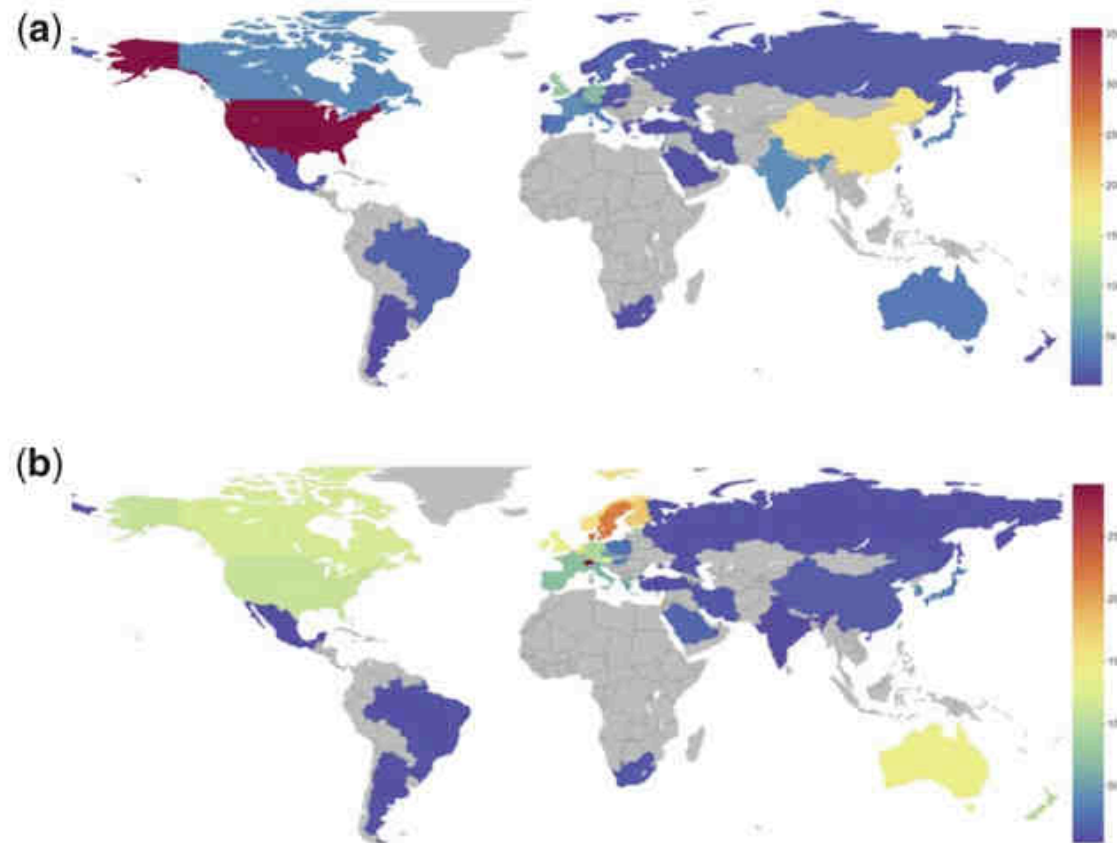


(A) Network presenting the research interactions between 11 LA countries, based in articles published on bioinformatics and computational biology between 1991 and 2016 (links created when the number of papers is  $N \geq 2$  and marked when  $N > 3$ ). (B) Number of research articles on bioinformatics and computational biology published each year by scientists from 19 LA countries (from 1991 to 2016). The total number of articles was 2119 and the total number of times that have been cited: 26 499.

Images in this article

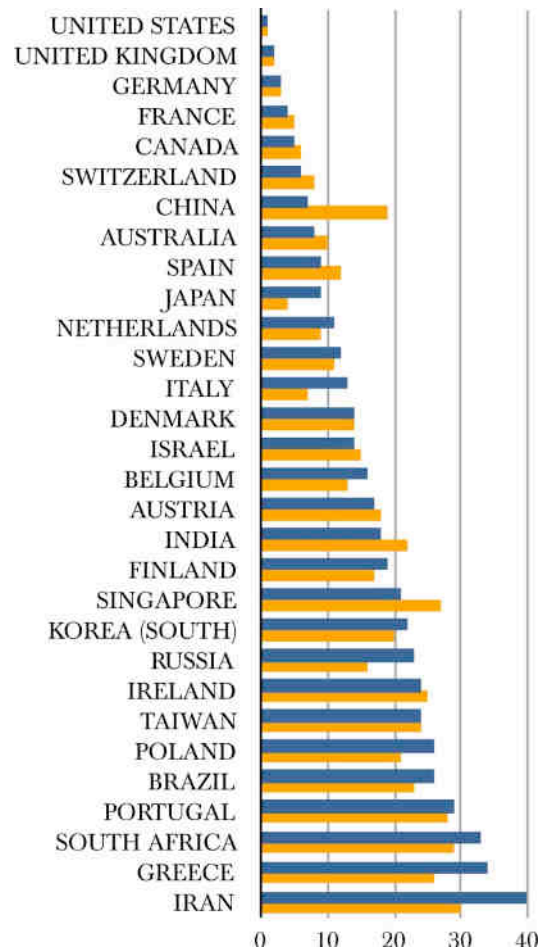
[Brief Bioinform. 2019 Mar; 20\(2\): 390–397.](#)

Fig. 2.



A world map depiction highlighting the top 40 countries in bioinformatics, based on publications output. (a) Absolute numbers of bioinformatics publications (scale provided, right), (b) relative number of bioinformatics publications per capita (million inhabitants). See [Supplementary Table S2](#) for a full list of 40 countries. Figure generated by Displayr ([www.displayr.com](http://www.displayr.com))





Rank listing of leading countries in bioinformatics and top 1% of highly cited papers. Bioinformatics h-index rank (blue), top 1% highly cited papers rank (orange)—lower is better.

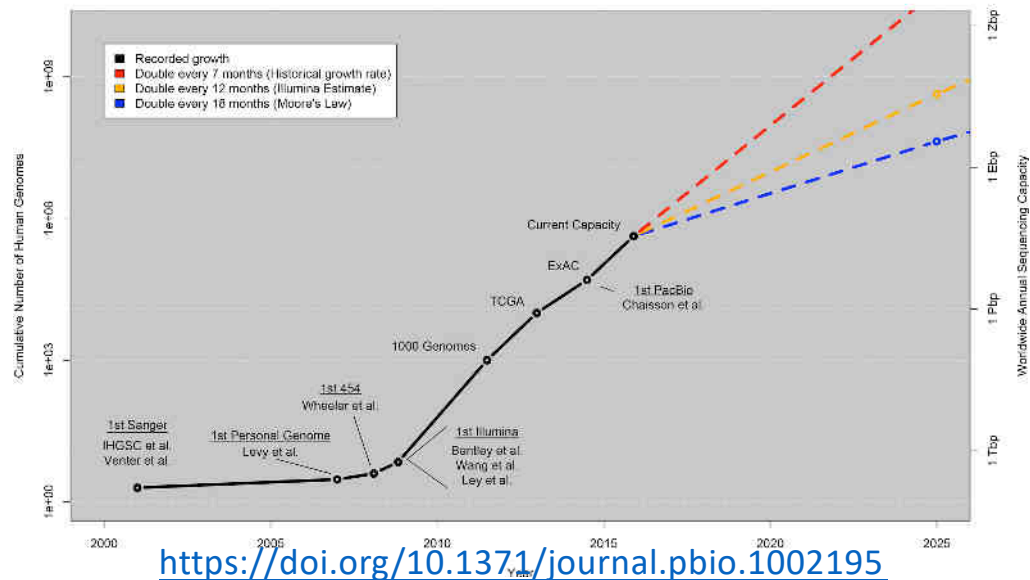
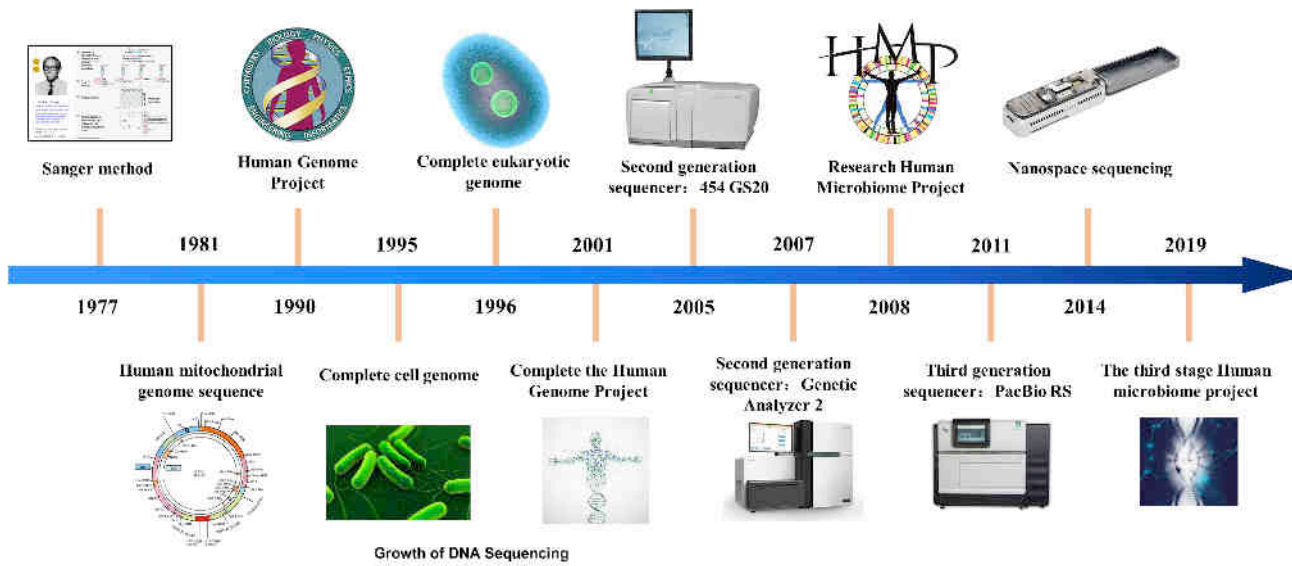
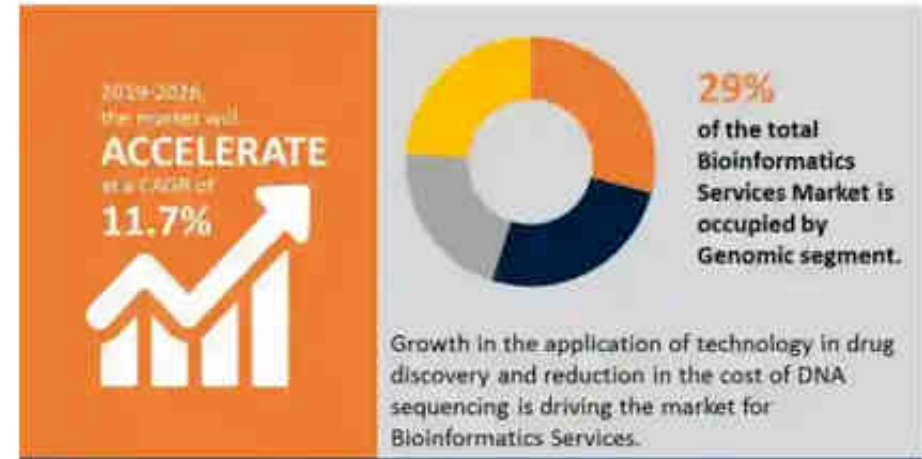
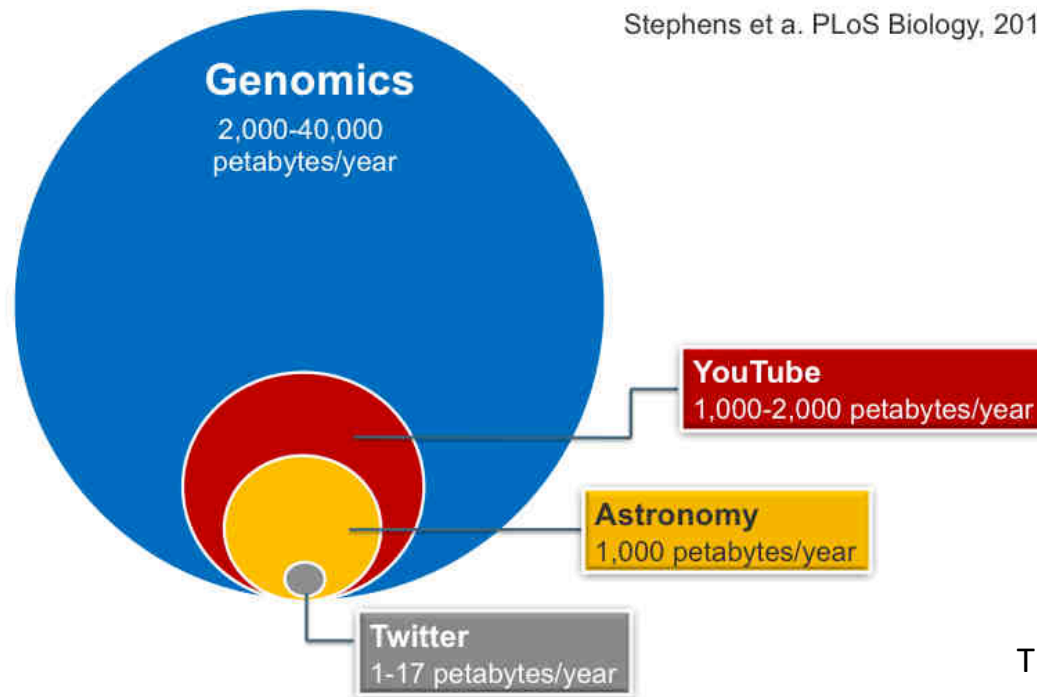


Fig 1. Growth of DNA sequencing. The plot shows the growth of DNA sequencing both in the total number of human genomes sequenced (left axis) as well as the worldwide annual sequencing capacity (right axis: Tera-basepairs (Tbp), Peta-basepairs (Pbp), Exa-basepairs (Ebp), Zetta-basepairs (Zbps)).

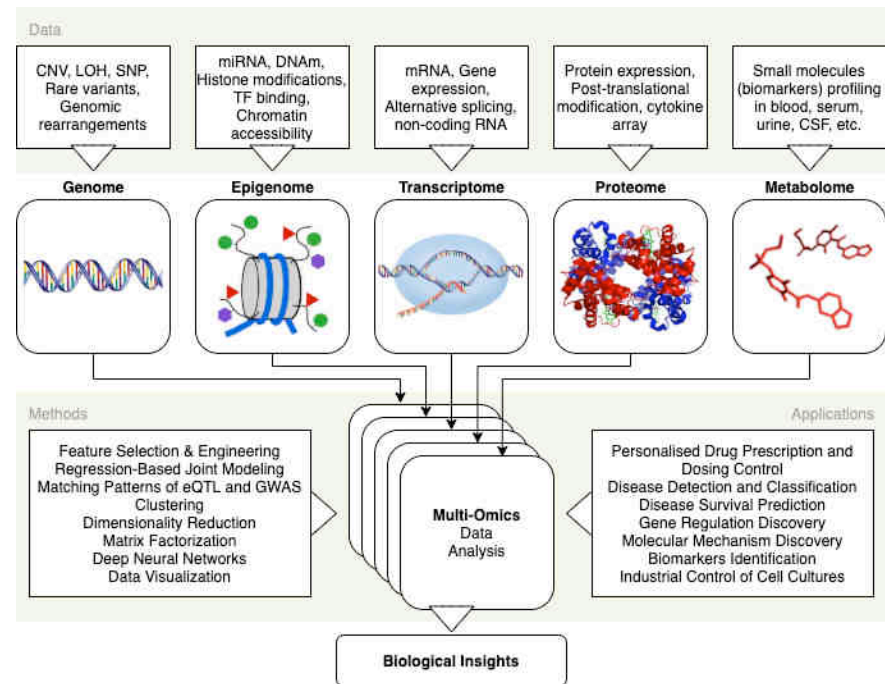
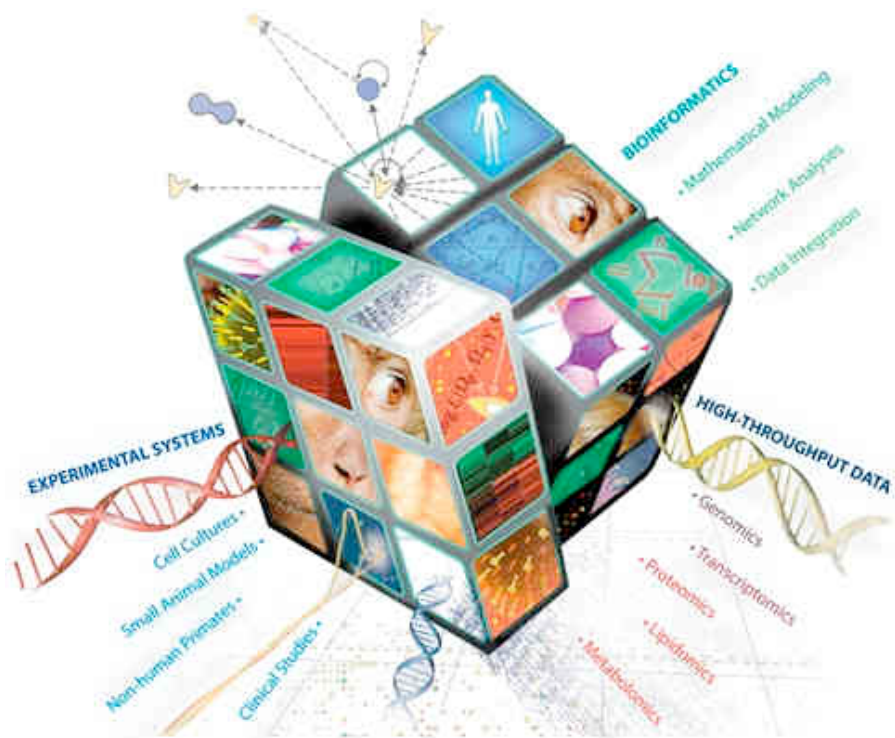
Stephens et al. PLoS Biology, 2015



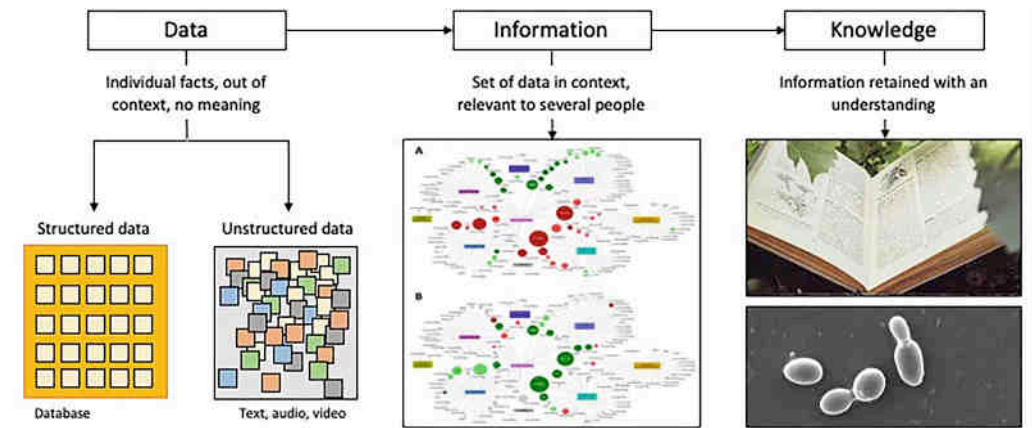
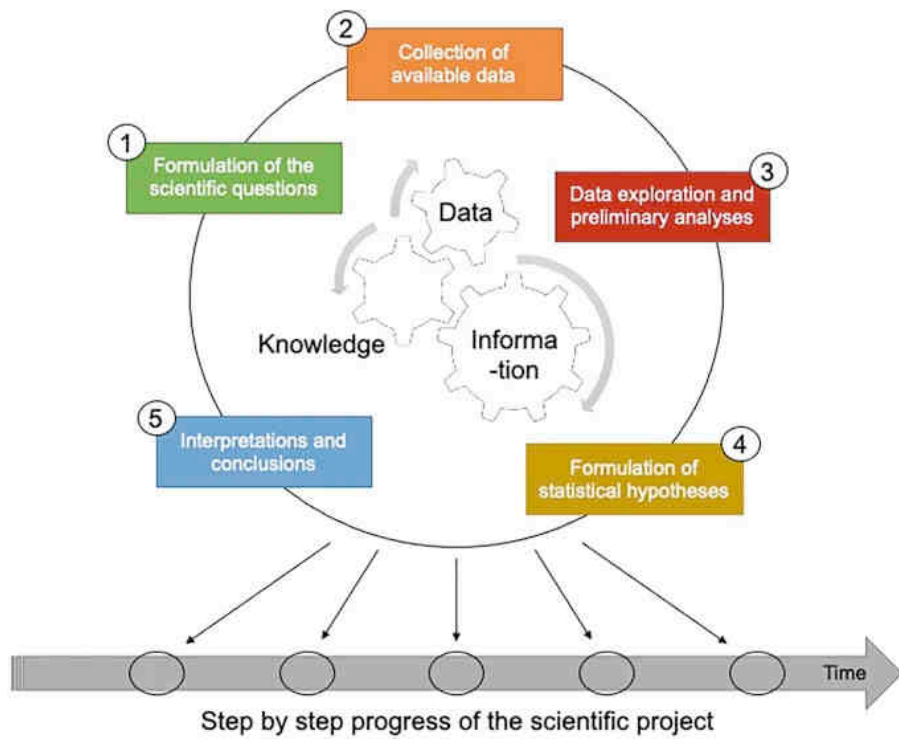
The Global Bioinformatics Services Market is forecast to reach USD 6.44 Billion by 2028.

<https://www.reportsanddata.com/report-detail/bioinformatics-services-market>

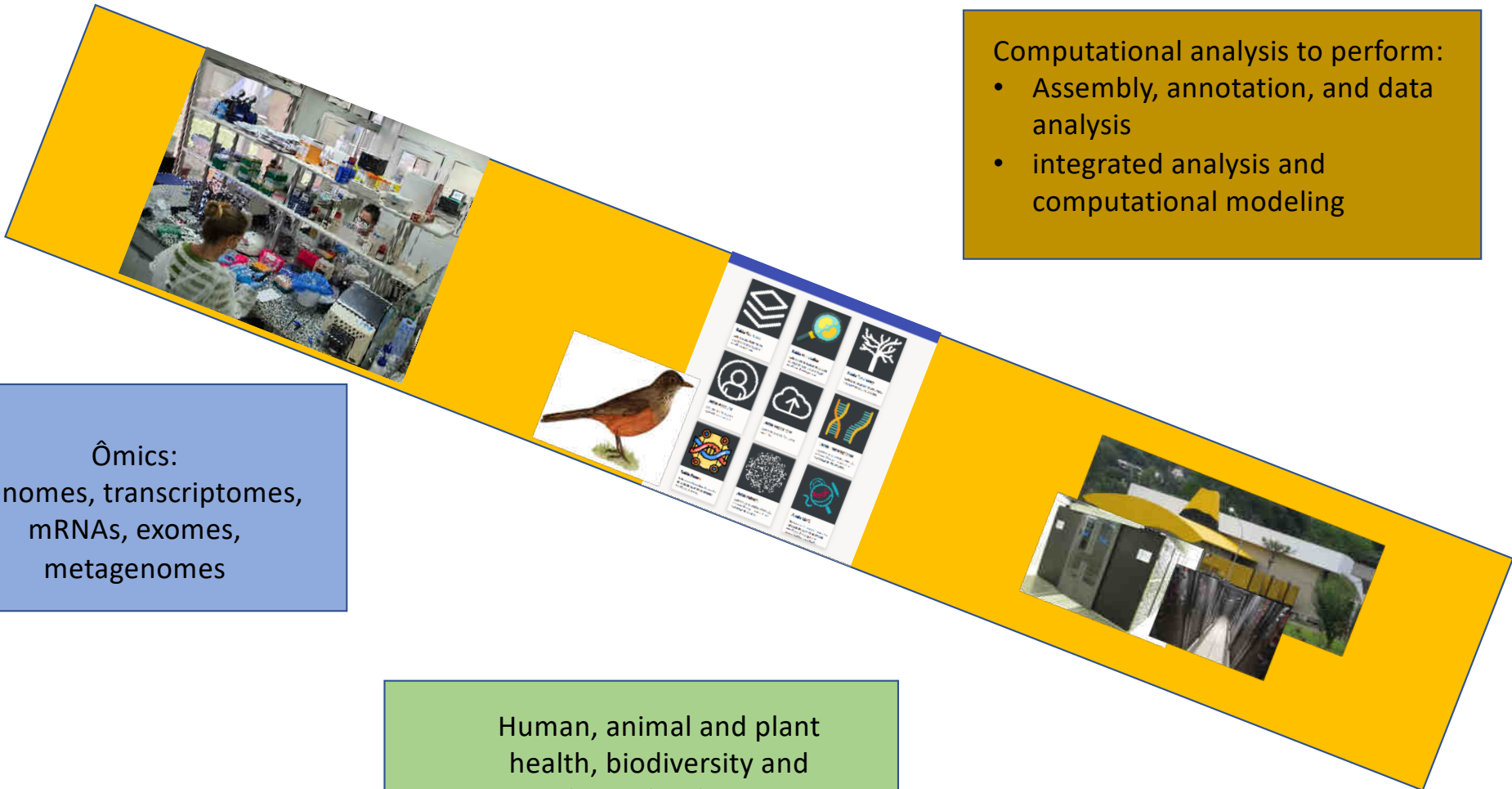
“Projecting to the year 2025, we compared genomics with three other major generators of Big Data: astronomy, YouTube, and Twitter. Our estimates show that genomics is a “four-headed beast”—it is either on par with or the most demanding of the domains analyzed here in terms of data acquisition, storage, distribution, and analysis.”



<http://www.a2idea.com/omics-analyses>



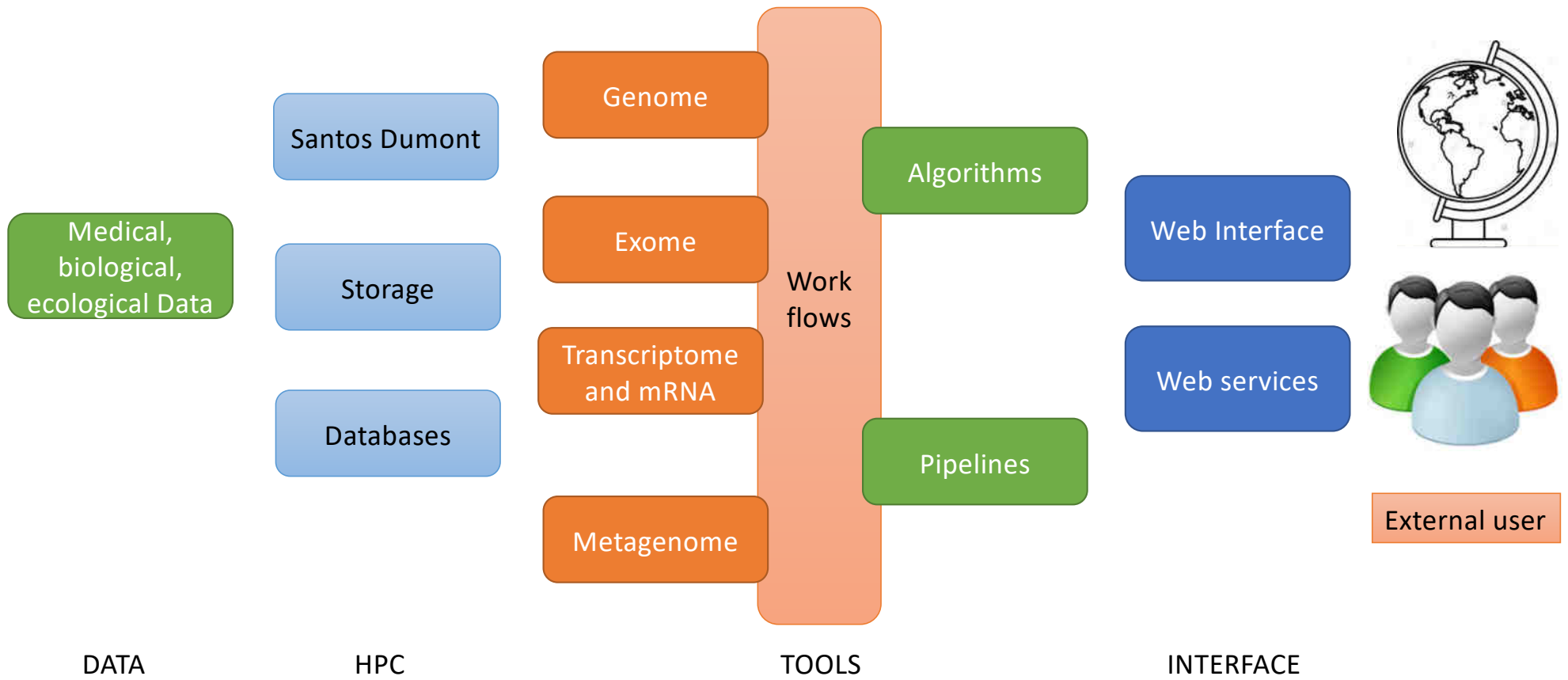




- Computational analysis to perform:
- Assembly, annotation, and data analysis
  - integrated analysis and computational modeling

Ômics:  
Genomes, transcriptomes,  
mRNAs, exomes,  
metagenomes

Human, animal and plant  
health, biodiversity and  
biotechnology

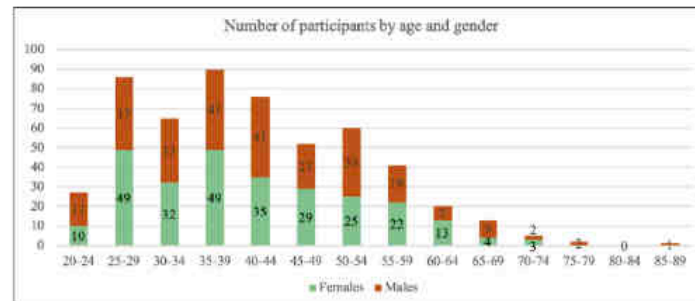


## The past, present and future of genomics and bioinformatics: A survey of Brazilian scientists

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The data was collected through an online questionnaire containing 25 questions, 19 closed-ended and six open-ended. The first closed-ended question asks how long the respondent is working in the area of genomics and bioinformatics. If the participant selected the option that states s/he is not working in the area, the questionnaire was then ended.

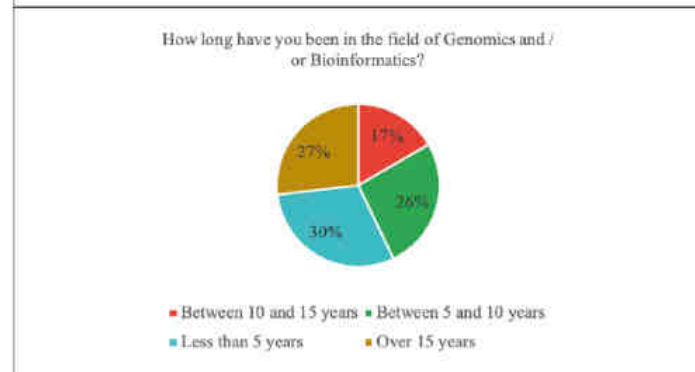
A



B

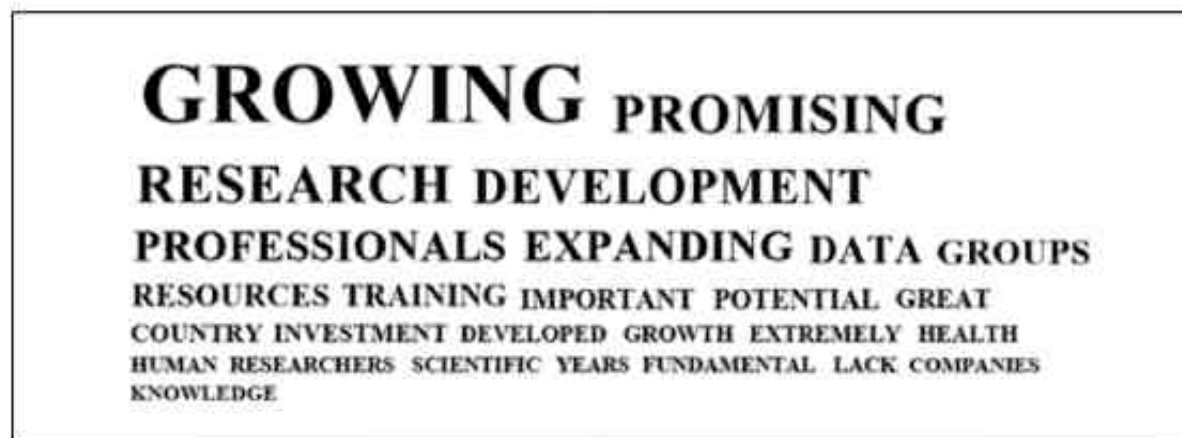


C



**Table 2** - Answer to the question: “Mark in the table below, for all levels of education, if you are attending or have already completed and in case you have already completed what was the year of completion”.

Time since completion	Bachelor's degree	Specialisation	Master's degree	PhD	Post-doctorate
Over than 10 years ago	64.5%	18.9%	44.9%	41.0%	22.6%
Between 05 and 10 years	15.5%	2.8%	13.3%	14.6%	12.2%
Up to 05 years	17.2%	3.5%	19.2%	14.4%	12.8%
Over 3 years ago	0.2%	0.0%	0.0%	0.2%	0.0%
Studying	0.9%	1.1%	7.8%	16.1%	10.2%



**Figure 2** - Word cloud illustrating the most frequent words occurring in the participants' responses referring to the current situation of the field of genomics and/or bioinformatics in Brazil.



Table 6 - Examples of sentiment analysis classification results performed on the future situation of the field in Brazil.



Sentiment classification	%	Example
Positive	46.0%	Case 38 - Extremely promising, with the emergence of new research and investment groups from the private sector in the creation of new companies specialised in the area.
Neutral	41.0%	Case 453 - I believe that all professionals will have to know the minimum bioinformatics for the development of their research in the near future.
Negative	13.0%	Case 107 - Difficult, resources are still limited and more collaboration is lacking, including data availability

Table 9 - Categories adopted to classify the milestones of genomics and/or bioinformatics in Brazil.



Category	Description	Percentage
Programs and projects	Implementation of programs and projects in the area, such as the Human Genome Project.	87.0%
Sequencing of organisms	Highlights of the main organisms sequenced, such as those relevant to areas like agriculture and health	66.0%
Technology	Development of new technologies, such as sequencers, high-performance computers and servers.	60.0%
Funding	More financial investment on research in the area	4.0%
Education and training	Development of formal education courses (bachelor's and post-graduation degree) and training programs.	13.0%
Industry and institutions	Creation of new institutions and companies /start-ups in the area	15.0%
Techniques	Development of new sequencing techniques.	10.0%

**Table 10** - Categories adopted on the classification of the respondents' wish lists.

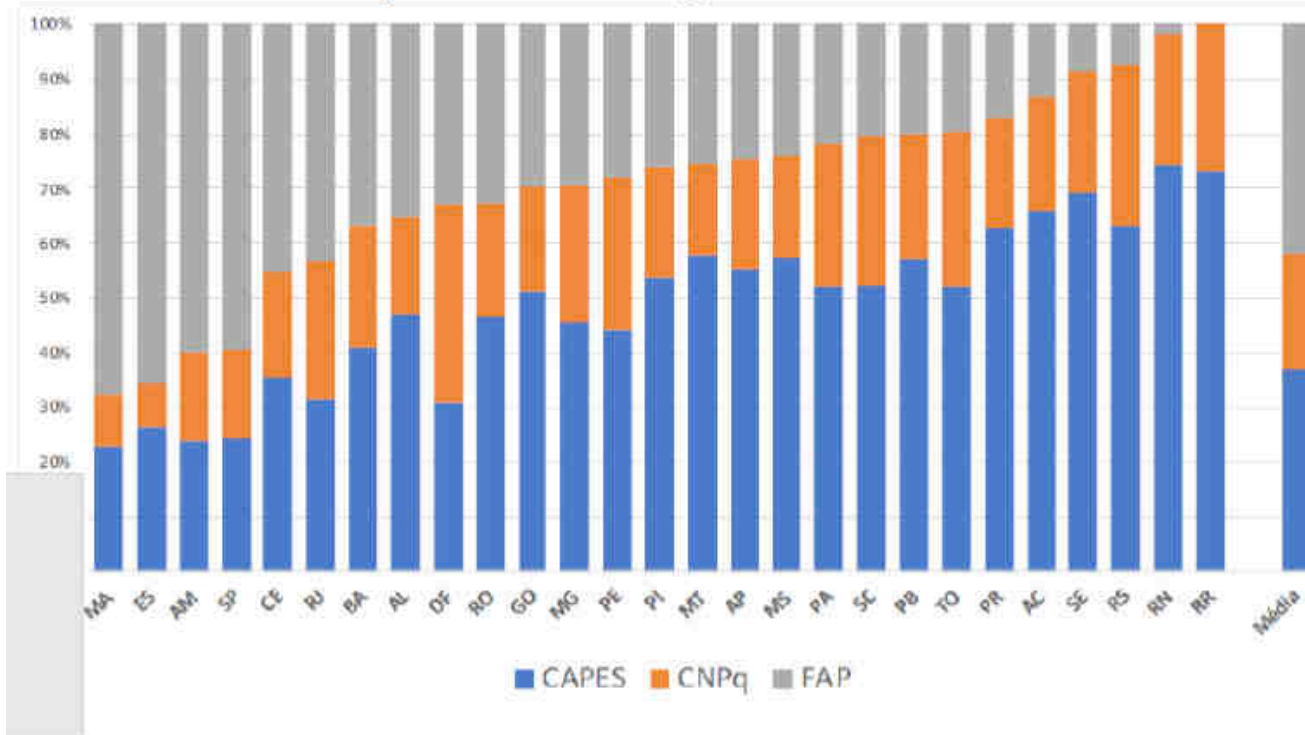
Category	Description	Percentage
Funding	Responses state the need to funding for research and development in the area. It also includes the reduction of the price of research inputs and equipment, besides investment in the industry providing larger teams and start-ups creation.	56.8%
Formal education	Responses related to the need of implementing new bachelor's, master's and PhD degrees, besides training for specific technologies.	48.2%
Collaboration	States the wishes for more collaboration between research groups, institutions and private companies/industry.	33.3%
Infra-structure	Comments about the need of new equipment researchers would like to have in their hosting university or company, such as supercomputers or research inputs.	32.5%
Career	A higher number of job positions for researchers and professionals in the area, higher salaries etc.	9.6%
Communication	Better communication about the area to the general public, besides including aspects related to the area to the basic school curriculum.	5.7%



UF	Média 2018-2020		Média 2018-2020		Média 2018-2020		Total (Soma CAPES+CNPq+FAP)	PIB em 2018 (1.000.000 R\$)	% do PIB	
	CNPq		CAPES		FAPs					
AC	R\$	1.379.359,89	R\$	4.370.800,00	R\$	861.205,50	R\$	6.611.365,39	15.331	0,043%
AL	R\$	6.957.728,68	R\$	18.128.400,00	R\$	13.469.500,15	R\$	38.555.628,83	54.413	0,071%
AM	R\$	14.909.830,35	R\$	22.375.200,00	R\$	55.226.347,14	R\$	92.511.377,49	100.109	0,092%
AP	R\$	1.004.960,66	R\$	2.779.600,00	R\$	1.247.455,99	R\$	5.032.016,66	16.795	0,030%
BA	R\$	36.646.609,68	R\$	68.047.200,00	R\$	61.242.543,41	R\$	165.936.353,09	286.240	0,058%
CE	R\$	31.401.020,40	R\$	57.749.200,00	R\$	73.033.453,66	R\$	162.183.674,06	155.904	0,104%
DF	R\$	64.772.071,68	R\$	55.588.400,00	R\$	59.330.797,59	R\$	179.691.269,28	254.817	0,071%
ES	R\$	9.500.469,30	R\$	31.893.600,00	R\$	78.194.349,20	R\$	119.588.418,51	137.020	0,087%
GO	R\$	17.252.596,73	R\$	46.105.600,00	R\$	26.716.504,47	R\$	90.074.701,20	195.682	0,046%
MA	R\$	5.415.352,60	R\$	13.118.800,00	R\$	38.926.358,89	R\$	57.460.511,49	98.179	0,059%
MG	R\$	126.631.095,86	R\$	231.575.200,00	R\$	148.493.413,14	R\$	506.699.709,00	614.876	0,082%
MS	R\$	9.707.091,55	R\$	30.196.400,00	R\$	12.533.147,23	R\$	52.436.638,78	106.969	0,049%
MT	R\$	7.373.514,78	R\$	24.840.400,00	R\$	10.839.015,00	R\$	43.052.929,78	137.443	0,031%
PA	R\$	27.145.510,65	R\$	53.408.000,00	R\$	22.308.096,18	R\$	102.861.606,82	161.350	0,064%
PB	R\$	27.438.942,33	R\$	68.868.400,00	R\$	24.157.379,02	R\$	120.464.721,35	64.374	0,187%
PE	R\$	51.746.265,08	R\$	82.034.000,00	R\$	51.641.429,77	R\$	185.421.694,85	186.352	0,100%
PI	R\$	5.783.512,29	R\$	15.434.400,00	R\$	7.452.378,57	R\$	28.670.290,85	50.378	0,057%
PR	R\$	55.863.983,16	R\$	175.161.600,00	R\$	47.511.898,77	R\$	278.537.481,92	440.029	0,063%
RJ	R\$	222.297.441,18	R\$	273.772.800,00	R\$	375.992.133,33	R\$	872.062.374,51	758.859	0,115%
RN	R\$	16.595.125,62	R\$	50.948.000,00	R\$	1.147.320,00	R\$	68.690.445,62	66.970	0,103%
RO	R\$	2.196.453,31	R\$	4.944.400,00	R\$	3.465.914,96	R\$	10.606.768,28	44.914	0,024%
RR	R\$	984.256,60	R\$	2.708.400,00	R\$	-	R\$	3.692.656,60	13.370	0,028%
RS	R\$	120.027.993,00	R\$	256.588.400,00	R\$	29.500.000,00	R\$	406.116.393,00	457.294	0,089%
SC	R\$	51.666.115,22	R\$	98.626.400,00	R\$	38.097.839,27	R\$	188.390.354,49	298.227	0,063%
SE	R\$	7.162.072,75	R\$	22.282.000,00	R\$	2.759.182,31	R\$	32.203.255,06	42.018	0,077%
SP	R\$	375.513.871,46	R\$	576.839.600,00	R\$	1.402.209.850,67	R\$	2.354.563.322,13	2.210.562	0,107%
TO	R\$	3.359.402,19	R\$	6.216.400,00	R\$	2.355.903,74	R\$	11.931.705,93	35.666	0,033%
x-Total	R\$	1.300.732.646,99	R\$	2.294.601.600,00	R\$	2.588.713.417,96	R\$	6.184.047.664,95	7.004.141	0,088%

Fonte : Odir Dellagostin

## Contribuição de cada agência em cada Estado



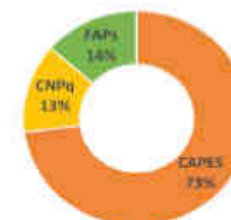
## Participação das agências no auxílio à pesquisa - dados de 2020

Agência	Auxílio à Pesquisa em 2020
Capes	R\$ 42.495.502,00
CNPq	R\$ 228.235.575,91
FAPs	R\$ 1.271.326.649,78



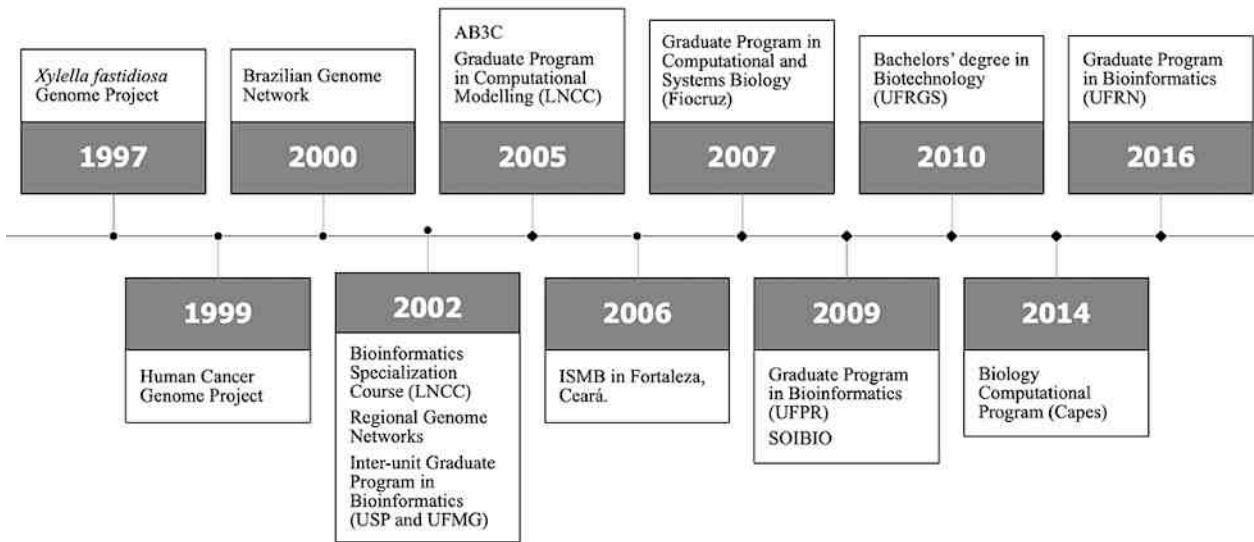
## Bolsas de pós-graduação em 2020

Agência	Mestrado	Doutorado
CAPES	43.497	46.105
CNPq	7.906	8.434
FAPs	8.737	8.101
<b>Total</b>	<b>60.140</b>	<b>62.640</b>
Bolsistas	44,2%	50,3%



Obrigado: Odir Dellagostin!







# Acknowledgment

