

# Highlights of the 1st Argentine Symposium of Young Bioinformatics Researchers (1SAJIB) organized by the ISCB RSG-Argentina

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## 29 Abstract

- 30 The 1st Argentine Symposium of Young Bioinformatics Researchers took place on 9-10
- 31 May 2016 at Universidad de Buenos Aires, Buenos Aires, Argentina. This event evolved
- 32 from a previous meeting series known as Argentine Student Council Symposium that
- 33 have been successfully organized since 2012 by the Regional Student Group of
- 34 Argentina (RSG-Argentina). The change in name reflects the new vision of the
- organizing committee to gather together all students at Bachelor, Master and PhD levels, postdocs and researchers that are still not Principal Investigator. Here we
- 37 summarize the main activities and outcomes from this year's meeting and offer some
- 38 insights into our future plans.



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## 41 Introduction

The Regional Student Group of Argentina (RSG-Argentina) was officially recognized as the local branch of the International Society for Computational Biology Student Council (ISCB-SC) in 2012. Ever since, it has steadily grown as a nonprofit, nationwide network of undergraduate students, PhD candidates and postdoctoral researchers working or interested in bioinformatics and computational biology. It aims to break the limits imposed by geography and science fields to provide a space of interaction among students and professionals, encourage the exchange of ideas and knowledge to develop opportunities for research and education, and assist in the growth of the bioinformatics community in the country and Latin America. Led by proactive students and with constant support from ISCB-SC and the Argentine Association of Bioinformatics and Computational Biology (A2B2C), the RSG-Argentina has been offering opportunities for learning new skills, sharing research outcomes and professional development in general for the last five years.

## Hightlights from 1st Argentine Symposium of Young Bioinformatics Researchers

The 1st Argentine Symposium of Young Bioinformatics Researchers (1st SAJIB) was held on 9-10 May 2016 at Universidad de Buenos Aires, Buenos Aires, Argentina. The symposium followed the format from previously organized events by RSG-Argentina, consisting of a first day of talks and poster presentations and a second day of practical workshops. We had 3 keynotes speakers and 6 short talks given by students in the first day, and 2 workshops in the second one.

## Scope and format of the symposium

The local computational biology and bioinformatics community has continuously grown over the last decade, especially since the Argentine Association of Bioinformatics and Computational Biology (A2B2C) was created in 2009. In total 6 annual Argentine conferences have been organized by the A2B2C in different cities across the country. Since its creation in 2012, RSG-Argentina has been supporting these conferences by gathering together the students community, engaging universities and organizing specific complementary workshops as satellite events at the venues. However, there are two main problems that constitute a barrier for young students to integrate into this community. As the Argentine annual conference on bioinformatics is held on different geographical places across the country, this makes it difficult for many people, specially students, to attend, both for economic and time reasons. Also, since internationally recognized keynotes are invited to the conference, the official language is English and this constitutes an idiomatic barrier for some undergraduate or PhD students in the early



- 76 years seeking to participate actively and get the most of the event by interacting with 77 senior scientists or presenting their work.
- SAJIB aims to complement the annual Argentine conference by solving the 78 aforementioned problems. The Symposium is planned to be organized annually in 79 80 Buenos Aires with Spanish as its official language. Since Buenos Aires is centrally located and well-connected it is easy and cheap to reach from most places in the 81 82 country. Following the structure of the symposia organized by the ISCB Student Council
- (Francescatto et al., 2015; Parra et al., 2015; Wilkins et al., 2016) and other RSGs 83
- 84 (White et al., 2016), this event is held on a more relaxed atmosphere than the main 85 conference, encouraging students to feel more comfortable presenting their work and
- interacting with each other. 86
- 87 Securing funding for this year's symposium allowed us to recognize the work and effort
- of some participants. Andrés Rabinovich, PhD student from Fundación Instituto Leloir 88
- 89 and CONICET fellow, was selected by the organizers to receive the Best Presentation
- award. All invited Keynote speakers and Workshop lecturers received a desktop gift. 90
- Plus, coffee-break refreshments and RSG-Argentina pins were given for free to all 91
- participants. 92

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# Keynote presentations

- We had three keynote speakers representing different areas of research who delivered interesting talks on their current projects related with Bioinformatics:
  - 1. Dr. Adrian Turjanski leads the Structural Bioinformatics Group at UBA-CONICET-IQUIBICEN. He works on the application of computational biology techniques to find potential drug targets in pathogen genomes, with a particular interest in fighting Mycobacterium tuberculosis.
  - 2. Dr. Ernesto Roman is a biochemist and computational biophysicist whose main interest is on the sequence-structure-dynamic-function relationships in proteins. He studies mutations on the human frataxin protein responsible for the Friedreich's ataxia syndrome.
  - 3. Dr. Pablo Turjanski is a computer scientist developing computational methods and algorithms to analyze biological data under a framework of Big Data science. He currently works on parameter-free and alignment-free methods for studying protein repeats.

## Short talks

- 109 The symposium offered 6 short talks by current undergraduate and PhD students, along
- 110 with 25 posters presented by undergraduate students, PhD candidates and postdoctoral
- 111 researchers.
- 112 Jorgelina Moreiras Clemente, from Universidad Nacional de la Plata (UNLP), gave the
- 113 first talk, where she presented her work on metagenomic analysis of soils from extreme



- environments, focusing on the search for microorganisms with potential use for soil
- 115 bioremediation.
- 116 Ariel Aptekmann, from Universidad de Buenos Aires (UBA), gave an insight into the
- 117 relation between optimal growth temperature and information content in the core
- 118 promoter region of archeal genomes.
- Andrés Rabinovich, from Fundación Instituto Leloir (FIL), presented an heuristic method
- to find group of genes that are similarly expressed with high biological congruence,
- using both gene expression profiles and gene ontology information.
- Julia Marchetti, from Universidad Nacional de Quilmes (UNQ), discussed the possible
- random origin of proteins through the analysis of enzymes with reduced numbers of
- amino acids in their composition, highlighting their capability to fold and sustain their
- 125 biological functions.
- 126 Victoria Dumas, from UBA, presented a study on the bacterial p450s family. It integrates
- information obtained from both the genetic and structural context of those proteins in
- order to develop a method to predict the reaction catalyzed by each enzyme.
- 129 Finally, Osvaldo Burastero, from UBA, shared his work on the Histidine Kinase CpxA
- autophosphorylation mechanism, for which he used a hybrid QM/MM approach.

# 131 Workshops

- Dr. Elin Teppa (FIL) and BSc/MSc Marcia Hasenahuer (UNQ) gave a 10-hour course
- 133 called 'Visualization, Analysis and Molecular Modelling'. This workshop aimed to
- introduce basic and advanced tools to represent, manipulate and model biomolecular
- 135 structures, using methods considered nowadays as standard tools. The course made
- 136 special focus on guided demonstration and hands-on activities to develop analysis
- 137 skills.
- 138 Dr. Diego Javier Zea (FIL) delivered the 8-hour course 'A day with (the) Julia
- 139 (language)'. This course introduces the basics of Julia, a high level programming
- language designed for scientific computing with a close to C performance. The
- workshop covers from the basic scripting tools needed in bioinformatics pipelines to the
- 142 available tools for data analysis.

## 143 **Feedback**

- 144 The 1st SAJIB was generally well-received. The presented posters and talks covered a
- wide range of topics and most participants agreed that the scope of the meeting suited
- their particular interests. As a networking opportunity, SAJIB favoured discussions and
- ideas were proposed for collaborations among students from different parts of the
- country. As seen in previous events, many students attending for the first time were
- 149 generally interested in participating more actively in the organisation and planning
- activities of RSG-Argentina. This provides an excellent opportunity for increasing the
- diversity of the organizing committee. Participants proposed the organization of similar



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- meetings in other cities in the country next year. It is difficult for a relatively small and
- self-organized group to reach sponsors, so our funding capability for travel costs and
- remote assistance is limited. However, as we are continuously growing our list of
- members and consolidating our presence in courses and meetings nationwide, this
- should become a priority. Overall, the feedback that we received was remarkably
- positive and encouraged us to continue organising events of this magnitude.

## **Conclusions and future directions**

159 Almost a decade after the first article on the emerging landscape of computational biology in Argentina (Bassi, González and Parisi, 2007), we have already reached many 160 milestones as a scientific community. The Argentine Association of Bioinformatics and 161 Computational Biology was created and with it, the Argentine conference in these fields 162 163 has been held annually for the last 7 years. The student community got organized and 164 created the Argentine branch of the ISCB Regional Student Groups. Bioinformatics and 165 computational biology have steadily grown in the last years and this is evident in the amount of research teams spread all over the country that are working purely on these 166 167 disciplines or using them as accessory tools for the wet lab. This growth is also seen in 168 the amount of publications in top tier journals in these fields and the emergence of 169 specific undergraduate and Masters programs. This offers an unprecedented number of 170 opportunities for young students looking for a place in academia and a career in 171 bioinformatics research.

As is commonplace for most scientific disciplines in developing countries like Argentina, there are many economic, social and geographic challenges preventing the reach of the full potential that the Argentine bioinformatics and computational communities have to offer. To overcome those difficulties, interaction among the students, senior researchers, A2B2C, the universities, the Ministry of Science and Technology and the National Scientific and Technical Research Council of Argentina has been and still is fundamental for the continuous development that we have experienced as a scientific community. The RSG-Argentina has the extremely important responsibility of bridging between the needs of students for improving their skills and future potential, and the academic and research strategies that are instrumented by senior investigators and organizations. Our organization promotes the nurturing of the youngest players in bioinformatics and computational biology, their involvement in specialized and high-quality meetings and the acquisition of soft skills and leadership capabilities to become active players in Argentina and internationally.

Social interaction and science communication among peers is the rosetta stone upon which we can build a competitive and solid community that will endure in time. In 10 years we moved from knowing about a disperse number of groups scattered along the country, with almost no connection among them, to have a solid scientific organization at different levels. The 1st SAJIB brought together more than 40 students from all over



- the country and from all academic levels, from undergraduates to postdocs and early
- 192 career researchers. The organization of this event by the students collective, gathered
- together in the RSG-Argentina and with the strong support of the senior researchers, is
- 194 proof that we are walking in the right direction and confirmation that many good things
- are still to come if we continue towards this path.
- 196 For more information regarding our activities please check our website
- 197 <a href="http://www.rsgargentina.com.ar">http://www.rsgargentina.com.ar</a>. We are also active on Twitter (@RSGArgentina) and
- 198 Facebook (RSG Argentina).

## 199 Author contributions

- 200 R. Gonzalo Parra and Nicolas Palopoli drafted the manuscript. All authors completed
- and reviewed the manuscript. All authors contributed to the organization of the
- 202 symposium.

## 203 Grant information

- 204 Universidad de Buenos Aires and the International Society for Computational Biology
- 205 Student Council (ISCB-SC) sponsored the event. Universidad de Buenos Aires provided
- 206 the venue for the meeting and computational support for the workshops. ISCB-SC
- 207 sponsored awards and event expenditures. The funders had no role in study design,
- 208 data collection and analysis, decision to publish, or preparation of the manuscript.

## 209 Acknowledgements

- 210 We are very grateful to all RSG-Argentina members who helped with students
- 211 arrangements, revision of abstracts and other organizational tasks related with SAJIB.
- 212 We would like to thank our PhD advisors and group leaders for their patience and
- support in allowing us to take part in this event and other RSG-Argentina activities. In
- 214 particular, we are very thankful to Prof. Marcelo Marti and Prof. Adrián Turjanski who
- 215 helped us with arranging the symposium facilities at Universidad de Buenos Aires.

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