



EDITORIAL

BOSC 2022: the first hybrid and 23rd annual Bioinformatics Open Source Conference [version 1; peer review: not peer reviewed]

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Abstract

The 23rd annual Bioinformatics Open Source Conference (BOSC 2022) was part of this year's conference on Intelligent Systems for Molecular Biology (ISMB). Launched in 2000 and held every year since, BOSC is the premier meeting covering open source bioinformatics and open science. ISMB 2022 was, for the first time, a hybrid conference, with the in-person component hosted in Madison, Wisconsin (USA). About 1000 people attended ISMB 2022 in person, with another 800 online. Approximately 200 people participated in BOSC sessions, which included 28 talks chosen from submitted abstracts, 46 posters, and a panel discussion, "Building and Sustaining Inclusive Open Science Communities". BOSC 2022 included joint keynotes with two other COSIs. Jason Williams gave a BOSC / Education COSI keynote entitled "Riding the bicycle: Including all scientists on a path to excellence". A joint session with Bio-Ontologies featured a keynote by Melissa Haendel, "The open data highway: turbo-boosting translational traffic with ontologies."

Keywords

bioinformatics; open source; open science; Diversity, Equity and Inclusion

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This article is an Editorial and has not been subject to external peer review.

Any comments on the article can be found at the end of the article.



This article is included in the **Bioinformatics** gateway.



This article is included in the **Bioinformatics** **BOSC** Open Source Conference (BOSC) collection.

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Introduction

The 23rd annual **Bioinformatics Open Source Conference**, BOSC 2022, was held as a Community of Special Interest, or COSI, which are tracks of the conference on Intelligent Systems in Molecular Biology, ISMB 2022. Occurring annually since 2000, BOSC has been part of ISMB in all but two occasions, in 2018 and 2020, when it partnered with the Galaxy Community Conference (GCC). BOSC's parent organization is the Open Bioinformatics Foundation (OBF), a non-profit, volunteer-run organization (Figure 1) that promotes open source software and Open Science in the biological research community.

Similar to BOSC 2021, which included a joint session with the Function COSI, BOSC 2022 featured joint keynotes with two other COSIs at ISMB: the Education and Bio-Ontologies tracks. It also featured the return of a panel, which we had not attempted to offer during the two years (2020 and 2021) when the meeting was held virtually.

Geraldine van der Auwera spoke for many when she observed, “[BOSC] is my favorite conference because it brings together such a welcoming group of people, and the lineup of talks typically offers a great balance of technical learning and community-building insights.”

The first hybrid BOSC

Due to the continued risk of COVID-19 worldwide, ISMB held the conference in a hybrid format, welcoming attendees in person and online via a virtual conference platform. The in-person component was hosted at the beautiful Monona Terrace Community and Convention Center (designed by Frank Lloyd Wright) in Madison, Wisconsin, USA. There were 997 in-person and 776 online participants. Overall, the conference was successful, although running a hybrid meeting remains more difficult than running an all in-person or all online event. Unfortunately, some sessions were plagued with technical issues, and the interaction between in-person and online participants was minimal. Regardless, hybrid meetings are probably here to stay, given the major advantage of greater inclusivity that online meetings offer, and taking into account the financial, environmental, and public health considerations associated with in-person attendance. As we all adjust to this new hybrid format, lessons learned from this year's experience will help improve the social and technical aspects of future hybrid meetings.



Figure 1. Some of the BOSC organizers (Nicole Vasilevsky, Nomi Harris, and Monica Munoz-Torres) with OBF Secretary Chris Fields. Other BOSC organizing committee members not shown: Jason Williams, Deepak Unni, Hervé Ménager, Karsten Hokamp. Shared under a [CC-BY-SA](https://creativecommons.org/licenses/by-sa/4.0/) license.



Figure 2. BOSC participants Scott Cain and Drew Hasley chatted while Drew's service dog, Shade, rested at his feet. Shared under a [CC-BY-SA license](#).

Diversity, equity and inclusion

As in previous years, BOSC prioritized its time and funds to ensure our meeting was more diverse and inclusive. With help from our sponsors, we granted free registration to 18 participants, including 16 attendees from groups that are underrepresented in bioinformatics, and covered travel expenses for several of the panelists and keynote speakers. Diversity, Equity and Inclusion was a focus of several of the BOSC sessions, including the panel and the session on Inclusion & Open Science (see below for details).

As far as we are aware, BOSC 2022 was the first BOSC, possibly even the first ISMB, to include a service dog ([Figure 2](#)), demonstrating tangible progress in our efforts to make BOSC more inclusive and accessible.

Conference program

Keynotes

This year's program included joint keynotes with two other COSIs ([Figure 3](#)). Jason Williams ([Figure 4](#)) gave a BOSC/ Education COSI keynote entitled "Riding the bicycle: Including all scientists on a path to excellence". Jason drew on his extensive experience as an educator at the Cold Spring Harbor DNA Learning Center, discussing the newly coined "bicycle principles" for making bioinformatics short-format training more effective, persistent, and inclusive. The principles are centered around two cyclic – hence "bi-cycle" – and iterative processes (see bikeprinciples.org, where you can offer feedback on the proposed principles). One "wheel" is made up of the Core Principles, which apply to all short-format training:

- Use best evidence
- Promote catalytic learning
- Be effective; provide evidence of learning
- Be inclusive

BOSC 2022 Invited Speakers

Keynotes

Jason Williams
Cold Spring Harbor Laboratory

Melissa Haendel
University of Colorado Anschutz
Medical Campus

Panelist

Jenea Adams
University of Pennsylvania

Panelist

Andrew Hasley
North Carolina State University

Panelist

Monica Munoz-Torres
University of Colorado Anschutz Medical Campus

Panelist

Gary McDowell
Lightoller LLC

Panelist

Rachel Torchet
Institut Pasteur

Figure 3. BOSC 2022 keynotes and panelists.

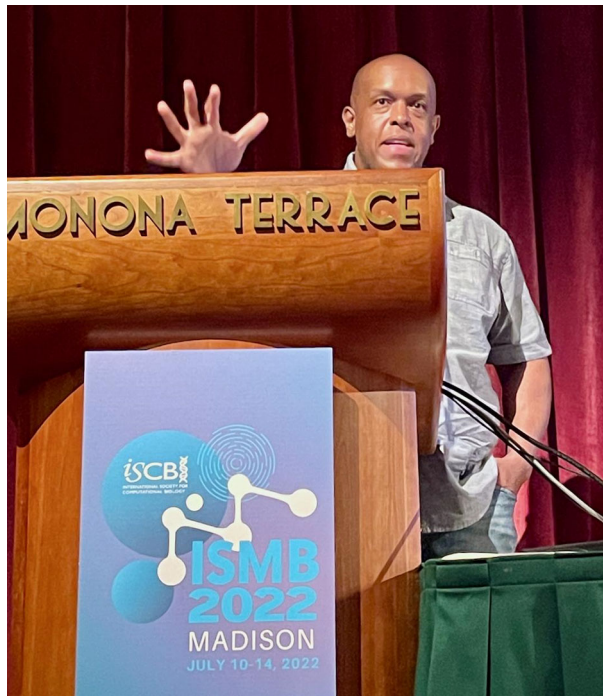


Figure 4. Keynote speaker Jason Williams discussed the “bicycle principles” for bioinformatics short-format training in a joint BOSC/Education COSI talk. Shared under a [CC-BY-SA license](#).

The other “wheel” is the Community Principles, which apply when short-format training is organized by communities:

- Reach
- Scale
- Sustain

The second keynote speaker, Melissa Haendel (Figure 5), headlined a joint session with Bio-Ontologies, In her talk, “The open data highway: turbo-boosting translational traffic with ontologies.” Melissa described some of the problems involved in integrating, harmonizing, and mapping biomedical data between multiple resources and standards. For example, there are many different terminologies and definitions of diseases, so how can we tell whether two diseases



Figure 5. In a joint BOSC/Bio-Ontologies session, keynote speaker Melissa Haendel took attendees on a fast ride down the translational highway with the help of ontologies. Shared under a [CC-BY-SA license](#).

discussed in different studies are in fact equivalent? Melissa leads projects such as the Mondo disease ontology, the Human Phenotype Ontology (HPO), and the LinkML data modeling language, which aim to help address those challenges. Melissa discussed ways to bridge the translational divide by leveraging semantics and ontologies — with the caveat that in most cases, if you think you need to invent a new ontology, you actually don't.

Panel

BOSC 2022 ended with a panel (Figure 6) entitled “[Building and Sustaining Inclusive Open Science Communities](#)” with panelists who not only are known for supporting diversity and inclusion, but also themselves are members of groups that are underrepresented in our field. As most of us are well aware, building great software is just a starting point; to maximize the impact, it's also necessary to put effort into maintaining it. Similarly, open source and open science communities need to be maintained and expanded, and initial efforts at inclusion and diversity need to be sustained and extended. This lively, interactive panel focused on the “what next?” part -- what needs to come after the initial steps to diversify an open source project or community.

Led by Jason Williams, who is well known for his work in diversity and inclusion, the panel included Jenea Adams, founder of the Black Women in Computational Biology (BWCB) Network; Andrew Hasley, a blind biologist with an interest in Universal Design for Learning, a framework for creating inclusive instructional environments; Monica Munoz-



Figure 6. Jason Williams moderating the panel on “[Building and Sustaining Inclusive Open Science Communities](#)”. Shared under a [CC-BY-SA license](#).

Torres, who works to improve diversity in the bioinformatics workforce; Gary McDowell, whose focus is on early-career researchers; and Rachel Torchet, who co-founded an organization to promote diversity, equality, and inclusion at the Institut Pasteur.

The panelists discussed challenges and opportunities that we can all tackle as a community to be more inclusive. As audience member Hilmar Lapp – who, by the way, is the only person who has attended all 23 BOSCs! – observed, “The panel reminded us that we all have much more agency than we think in making spaces welcoming to everyone who wants to enter.”

Talks and posters

As usual, BOSC included topical sessions with short and long talks chosen from submitted abstracts. The complete talk schedule can be found [here](#). This year’s session topics were:

- Inclusion & Open Science
- Analysis tools & approaches
- Data access and visualization
- Standards for Open Science
- Workflow Management Systems
- Open ontologies and tools (joint session with Bio-Ontologies)

The new session on Inclusion & Open Science featured a talk by Rachel Torchet on gender-based disparities and biases in science, with a quantitative analysis based on observational study of the French Open Days in Biology, Computer Science and Mathematics (JOBIM for its initials in French) bioinformatics conference. In one notable finding, the online JOBIM attracted more female attendees than previous in-person conferences, but even so, women attendees asked half as many questions as men.

The always popular Workflow Management Systems session ([Figure 7](#)) included a number of talks related to Common Workflow Language (CWL), which grew out of a BOSC CodeFest (as it was then known) in 2014. The hero of this session was Geraldine van der Auwera ([Figure 8](#)), who compressed her 20-minute talk about deciphering Workflow



Figure 7. Matthew Gazzara and Farica Zhuang presented “A cloud based international community effort for reproducible benchmarking of genomic tools” during the Workflows session. Shared under a [CC-BY-SA license](#).



Figure 8. Geraldine van der Auwera heroically compressed her 20-minute talk about deciphering WDL workflows into a 5-minute lightning presentation. Shared under a [CC-BY-SA license](#).

Bioinformatics Hub of Kenya Initiative: Connecting bioinformatics researchers and students in East Africa

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1. International Institute of Tropical Agriculture (IITA)
2. International Centre for Insect Physiology and Ecology (ICIPE)
3. Bioinformatics Hub of Kenya Initiative (BHKi)
4. Institute of Primate Research (IPR)
5. University of Nairobi

Abstract

In recent years there has been a significant uptake of bioinformatics projects within Sub-Saharan Africa (SSA), but only a few individuals are skilled in analyzing bioinformatics data. The bioinformatics services are mostly outsourced, thus neglecting capacity building and empowering the SSA community with bioinformatics skills. The Bioinformatics Hub of Kenya Initiative (BHKi) (<https://bhki.org/>) is a community-based organization aiming to train and empower students and researchers in SSA. BHKi also serves as a platform for networking and creating collaboration between researchers.

We provide different models of learning and training through workshops, hackathons, talks, seminars, science cafes, and conferences. Our approach involves a train-the-trainer model where we teach participants core skills and assign them mini-projects to solidify their skills. Participants present their mini-project outputs at BHKi events, and we urge them to train colleagues in their affiliated institutions. We have also collaborated with the Carpentries organization to train BHKi members to be instructors and lead different workshops.

From the initiative, BHKi has organized multiple workshops on various computational topics and programming languages. BHKi managed to sensitize on Open Science, conduct a genomic data analysis workshop, train-the-trainer session, hackathon challenges, and a conference themed 'Empowering researchers with Bioinformatics and Open Science Skills (BOSS)'. The BOSS events' had a good representation through the region and career levels.

The BHKi success has been achieved through the strong dedication of the organizing team, leadership team, and strong support from other organizations. We plan on university and high school outreach to further sensitize the bioinformatics field and work on open source projects.

Multi community approach

Open Life Science
Project training
Project management

Open Science Kc
Open data
Open Science

Carpentries
Training materials
Instructor training

BHKi

TCC Africa
Communication skills

Achievements

- Won our first grant from the Open Science Society grant for "Training and Open Science"
- OBF fund for a training program for research groups and researchers
- Successfully completed Hack, Collaborate
- Conducted our first social media outreach activities in Undergraduate schools
- Organized multiple events
- Registered on the Open Science Society

Communities helping us achieve goals

Open Science Kc
High schools
Open Science Kc
Communities
Colleges
BHKi events and target communities/organizations
Self-organized communities
Private community groups
Research institutions

Approach

Train the trainers
Engage the trainers with training materials and assign them to trainees in the community

Outreach
Reach out to various community, institutional and other events

Train the learners
Training the learners to become self-organizing communities and open science

Sensitize
Influence these communities about bioinformatics skills and importance of open science

Impact

Attendance by Career Level

Attendance by Country

Festus Nyasimi

Figure 9. Poster presenter Festus Nyasimi, who received an OBF Event Fellowship to help defray his expenses in attending ISMB/BOSC. Shared under a [CC-BY-SA license](#).



Figure 10. Some of the in-person CoFest 2022 participants working together at the Madison Public Library. Shared under a [CC-BY-SA license](#).

Description Language (WDL) workflows into a 5-minute lightning presentation to help make up for an otherwise catastrophic delay earlier in the session due to technical issues.

46 posters were presented at BOSC 2022, 29 in person (Figure 9) and 17 in the online platform.

CoFest

As usual, BOSC/ISMB were followed by a two-day **CollaborationFest** (CoFest). A hands-on-keyboards event that has been held after BOSC for the last twelve years, CoFest was originally called CodeFest, and was renamed “CollaborationFest” in 2018 to acknowledge the importance of activities besides coding, such as working on documentation, training materials, and use cases. Like ISMB, CoFest 2022 was a hybrid meeting, with some participants in person (Figure 10) and others online. Participants worked together on six collaborative projects, three primarily involving in-person participants and three online. Bhavesh Patel, a first-time BOSC and CoFest attendee, observed, “The CoFest was my favorite part as it provided a very informal setting to get to know all the participants on a personal level. It also gave me a unique opportunity to discuss personally with experts of the field and get one-on-one feedback on my work.”

Data availability

No data are associated with this article.

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