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# Revisiting barriers to implementation of bioinformatics into life sciences education

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**Introduction:** Bioinformatics is an interdisciplinary field at the intersection of computational and biological sciences that focuses on the analysis and interpretation of large biological data sets. Although recognized as essential in the life sciences, bioinformatics is not commonly integrated in undergraduate life science education programs. Based on a national survey in 2016, the Network for Integrating Bioinformatics into Life Sciences Education (NIBLSE) published a community-sourced set of core competencies in bioinformatics education. The survey also identified barriers that prevent incorporation of these competencies into the curriculum. In the current study, the NIBLSE group reports the findings of a new survey to 509 life science educators across the US in 2022 to identify current barriers of bioinformatics integration and to determine if the landscape of bioinformatics education has changed since the 2016 survey.

**Results:** Similar to previous results, a majority of respondents who currently teach bioinformatics or plan to teach bioinformatics report barriers. The top two barriers reported are students lacking prerequisite skills/knowledge and instructors lacking time to restructure course content. As in 2016, women reported experiencing barriers to bioinformatics teaching significantly more often than men; faculty from underrepresented minority backgrounds reported barriers more often than non-URM faculty; and educators at minority-serving institutions (MSIs) reported barriers more frequently than colleagues at non-MSIs. For additional insight into the barriers facing these educators, we conducted focus groups which provided qualitative data that supported the survey findings and revealed common themes including faculty perceptions of the relevance of bioinformatics in the curriculum. Despite the perceived value of bioinformatics education, many focus group members cited lack of student preparation and interest, and technological access as barriers. Participants also discussed how professional development and community support would enhance and sustain bioinformatics teaching.

**Discussion:** Taken all together, this study indicates that challenges remain, which vary among faculty types and settings, but that more educators are attempting to integrate bioinformatics into life sciences education. In summary, our results suggest that redoubled efforts to provide training and community support to life sciences faculty is necessary.

## KEYWORDS

undergraduate education, stem education, bioinformatics, barriers to integration, education reform

## Introduction

Bioinformatics is a rapidly expanding interdisciplinary field that encompasses computer science, mathematics, statistics, and biological data (Gauthier et al., 2019). Bioinformatics has greatly influenced the types of questions addressed by contemporary researchers, and given the sheer scale of datasets, how biological research is done. Typically, bioinformatics involves retrieving, processing, filtering, visualizing, and analyzing omics-focused datasets. The rapid advances in inexpensive high-throughput technology driving the sustained torrent of “big data” makes it critical for today’s life science trainees to learn the skills necessary to analyze biological data and make appropriate inferences. Furthermore, many of these disciplinary data science skills may be applied to datasets beyond the scope of biology. Consequently, providing early skill development in bioinformatics helps emerging scientists advance their research potential and primes them to seek out opportunities in higher education and the technical workforce (Porter and Smith, 2019). The field of bioinformatics has grown from early studies on protein sequences in the 1950’s. With the ability to sequence DNA, it has become easier to gather large numbers of sequences; at the same time, computer technology (both hardware and software) have advanced in order to analyze these sequences. It has been suggested that the term “bioinformatics” may soon become completely synonymous with biology (Gauthier et al., 2019).

Currently, training in the use of bioinformatics tools is a major bottleneck in this fast-moving field. In response to this bottleneck, there have been numerous calls for professional development in this area (Carvalho and Rustici, 2013; Attwood et al., 2019) and integration into educational frameworks (Altman, 1998; Machluf and Yarden, 2013; Wilson Sayres et al., 2018; Attwood et al., 2019; Niepielko and Shumskaya, 2021; Rocha et al., 2022; Isik et al., 2023). Integration of bioinformatics learning objectives across undergraduate life science degree programs can theoretically address the issue of bioinformatics literacy (Dinsdale et al., 2015). Further, bioinformatics-driven undergraduate curricula, when carefully designed, also have the potential to address traditional STEM access and diversity barriers (Elgin et al., 2021; Handelsman et al., 2022; The Genomic Data Science Community Network, 2022).

The Network for Integrating Bioinformatics into Life Sciences Education (NIBLSE) began in 2014 as an NSF-funded research coordination network focused on fostering community efforts to promote the integration of bioinformatics within undergraduate biological sciences (Dinsdale et al., 2015; Toby et al., 2022).<sup>1</sup> The network’s overarching goals are to identify best practices for (1) preparation of students for bioinformatics instruction; (2) integration

of bioinformatics into life science curriculum at all levels; (3) assessment of outcomes; and (4) preparation of faculty trained in life sciences to deliver bioinformatics-based curricula.

One of NIBLSE’s primary accomplishments was to develop a community-driven set of competencies for undergraduate bioinformatics instruction (Wilson Sayres et al., 2018). These competencies were based in part on a national survey of life sciences undergraduate faculty conducted in 2016, which gathered information about the nature of bioinformatics education in the US. The survey also provided the first deep investigation of the barriers that undergraduate life sciences faculty face in efforts to integrate bioinformatics (Williams et al., 2019). The barriers questions were open-response items that were then investigated by keyword analysis to quantify thematic responses. We grouped the reported barriers into several “super-categories”: faculty issues (e.g., lack of training, competing time commitments), student issues (e.g., lack of adequate math preparation, lack of adequate coding experience, lack of student interest), curriculum issues (e.g., lack of time for course development, lack of time for inclusion in an overfilled curriculum, speed at which materials and tools change) and less frequently, facilities, resource, and institutional issues (Williams et al., 2019).

The results from the 2016 survey included several key findings (Williams et al., 2019). Overall, lack of expertise and training at the faculty level was identified as the most common barrier among respondents followed by a lack of prerequisite skills and knowledge at the student level. Furthermore, the reported prevalence of barriers varied by demographic groups. In particular, women and faculty identifying as under-represented minorities (URM) reported barriers at higher rates than their comparison groups, men and non-URM faculty.

In the current study, we investigated whether the landscape of barriers for implementing bioinformatics in undergraduate curricula has changed since 2016 by conducting a new national survey followed by focus group discussions. Results from the 2016 survey were used to design Likert-scale items to explore faculty perceptions of specific barriers. For example, in the keyword analysis approach in 2016, many respondents included the word “time” as a barrier but upon reading the responses, time meant different things (time for course development, time/space in the curriculum, time for training), so we used more specific language to explore these issues further in the current survey. One impetus for the study was to further investigate the extent to which barriers to integration of bioinformatics were perceived by faculty who are members of under-represented minority groups and/or faculty at minority-serving institutions (MSIs). Because the number of survey respondents from these faculty was limited, we augmented the information collected in the new survey with a series of focus groups to understand these faculty experiences in more detail. In summary, many of the same barriers still exist although a greater proportion of faculty are in fact attempting to incorporate bioinformatics. Their efforts can be supported by faculty training and

<sup>1</sup> <https://qubeshub.org/community/groups/niblse>

up-to-date curricular materials (Ryder et al., 2020; Drew et al., 2021; Kleinschmit et al., 2021, 2023), efforts in which NIBLSE continues to engage.

## Materials and methods

### Solicitation of survey participants

A mailing list of approximately 13,000 life sciences faculty at 2- and 4-year US institutions was purchased from [ExactData.com](https://www.exactdata.com), a targeted, direct-marketing company that maintains lists of U.S. college and university faculty. As we were particularly interested in responses from faculty from historically underrepresented minority (URM) groups in STEM (e.g., African Americans, Hispanics, Native Americans and Native Alaskans, and Pacific Islanders), we included all available URM biology faculty records from ExactData, approximately 3,000 individual entries. The balance of the records was non-URM faculty. Among other information, each record contained the name, e-mail address, ethnicity, gender, and current institution of the individual.

Using data from the Carnegie Classification of Institutions of Higher Education,<sup>2</sup> information about an individual's current institution was added to each record, including basic Carnegie classification, size and setting, "control" (whether the institution is public or private), and whether the institution is an HBCU, Tribal, Hispanic-serving, Minority-serving, or women-only institution. For ease of data analysis, the 34 basic Carnegie classifications were collapsed into nine: (1) Not classified; (2) Associate's Colleges; (3) Baccalaureate/Associate's Colleges: Associate's Dominant; (4) Doctoral/Professional Universities; (5) Master's Colleges and Universities; (6) Baccalaureate Colleges; (7) Baccalaureate/Associate's Colleges: Mixed Baccalaureate/Associate's; (8) Special Focus Four-Year; and (9) Tribal Colleges. Similarly, the 19 Carnegie size and setting classifications were binned into five: (1) Not classified; (2) Small or Very Small; (3) Medium; (4) Large or Very Large; and (5) Exclusively graduate/professional.

### The 2022 barriers survey

A 24-item survey instrument, informed by the previous 2016 NIBLSE survey (Williams et al., 2019), was developed to probe the barriers faculty face in integrating bioinformatics into their teaching (See [Supplementary material 1](#)). The survey began with a consent statement, then contained several types of items (multiple-choice, free-response write-ins, and Likert-scale) organized into three blocks: (1) questions that asked how much bioinformatics the respondent currently included in their teaching, (2) the barriers they encountered when doing so, and (3) self-reported demographics including degree(s), current institution(s) and academic role, and extent of bioinformatics training. To streamline the survey for respondents, branching logic was used when appropriate to filter questions (for example, respondents who

indicated no barriers to integrating bioinformatics did not see follow up questions about specific barriers).

The survey was administered using Qualtrics (Qualtrics, Provo, UT, 2020), after approval from the University of Florida's Institutional Review Board (UF IRB#202102370). Following recommendations from Dillman et al. (2014), a link to the survey was distributed by email to the 13,398 individuals in the mailing list on November 4, 2021 and remained open until January 10, 2022 with three reminders sent throughout that period. There were 563 responses to the survey. All potentially identifying information was removed from the data before analysis.

### Survey analysis methods

Data from respondents who did not teach undergraduate courses were removed from the data set, as was data from individuals who did not provide consent, even if they completed other questions in the survey. As in Williams et al. (2019), the barriers reported by respondents were analyzed with respect to several demographic criteria including sex, race/ethnicity, year of highest degree, level of bioinformatics training, extent of current bioinformatics teaching, institutional Carnegie Classification, and MSI status. For a given demographic, respondents who did not answer were not included in that demographic category.

Data analysis was performed in R version 4.1.2 (R Core Team, 2021) (Refer to our [Supplementary material](https://github.com/niblse/barriers_survey_2022) at [https://github.com/niblse/barriers\\_survey\\_2022](https://github.com/niblse/barriers_survey_2022) for all R scripts used in this analysis). Data preparation, statistical summation, and graph preparation were performed using tidyverse (Wickham et al., 2019). Statistical tests (chi-squared, difference in proportions) were performed using infer R package (Couch et al., 2021). The false discovery rate method of Benjamini and Hochberg (1995) was used to adjust  $p$ -values for multiple hypothesis testing when appropriate.

### Focus groups

Five focus groups, each 60–90 min long (Saldana and Omasta, 2022), were conducted with life sciences faculty about the perceived barriers to integrating bioinformatics into their courses. IRB approval for the focus groups was sought from Georgetown University's Institutional Review Board (GU IRB# STUDY00005685), and the study was deemed exempt. In order to volunteer, participants were asked to complete a secure electronic application form ([Supplementary material 2](#)), which explicitly served as consent. A link to the form was sent to individuals on the mailing list. In addition, a flier containing a QR code that linked to the application form was disseminated by social media (specifically, Twitter) and at the 2022 Society for Advancement of Chicanos/Hispanics & Native Americans in Science meeting to elicit additional volunteers. All individuals who completed the form were encouraged to send the form to others who might also be interested ("snow-ball" sampling). A total of 118 individuals responded to the form. Of these, 35 individuals were invited to participate based on their availability and the type of institution at which they taught (as listed in [Table 1](#)) as we aimed for faculty at a diverse set of institutions. As an incentive for participation, those who participated were sent a NIBLSE-branded beverage set.

<sup>2</sup> <https://carnegieclassifications.acenet.edu/>

TABLE 1 Demographic characteristics of the focus group participants.

Group	Focus institution category	Participant	Carnegie classification	Minority status	Self-identification
1	AANAPSI*	1	Doctoral University	AANAPSI	Asian/White, Male
		2	Master's University	(Most students are from URM, high % of Native Americans)	Hispanic/Latino, Female
		3	Associate's College	AANAPSI, HSI	White, Male
2	HSI	1	Baccalaureate College	HSI	White, Female
		2	Doctoral University	HSI	White, Male
		3	Master's University	HSI	Asian, Female
		4	Associate's College	HSI	White, Female
		5	Associate's College	AANAPSI, HSI	Black/African-American, Male
3	HSI	1	Baccalaureate College	HSI	Hispanic/Latino, Male
		2	Baccalaureate/Associate's College	HSI	Hispanic/Latino, Female
		3	Baccalaureate College	HSI	White, Female
		4	Doctoral University	HSI	White, Male
		5	Associate's College	HSI	Black/African-American, Female
		6	Associate's College	HSI	White, Female
4	HBCU/PBI	1	Doctoral University	HBCU	Hispanic/Latino, Male
		2	Baccalaureate College	(Not an official PBI but has largest number of African-American students in peer group)	White, Male
5	Primarily Majority Serving	1	Doctoral University	Predominantly white	Hispanic/Latino, Male
		2	Associate's College	Predominantly white	White, Female
		3	Associate's College	HSI	White, Female
		4	Associate's College	HSI	Hispanic/Latino, Male
		5	Baccalaureate College	Predominantly white	White, Male
		6	Baccalaureate College	Predominantly white	Hispanic/Latino, Male
		7	Baccalaureate College	HSI	White, Female
		8	Master's University	Predominantly white	White, Female

\*AANAPSI, Asian American Native American Pacific Islander Serving Institution; HBCU, Historically Black College or University; HSI, Hispanic-Serving Institution; PBI, Primarily Black Institution; URM, Under-represented minority.

Ultimately, 23 faculty participated in the focus groups, the majority of whom were from Minority-Serving Institutions (Table 1). One focus group was comprised primarily of faculty at Asian-American and Native Pacific Islander serving institutions (AANPI), two groups were comprised of faculty at Hispanic-Serving Institutions (HSIs), one group was comprised of mostly faculty at Historically Black Colleges and Universities (HBCUs) or Primarily Black Institutions (PBIs). The final focus group was composed of faculty from majority-serving institutions, several of whom identified as either African-American or Hispanic (Table 1). The focus group participants also taught at a variety of institution types, including community colleges (Table 1).

The semi-structured focus groups were conducted by a facilitator from Georgetown's Center for New Designs in Learning and Scholarship. An interview guide [written with reference to Saldana and Omasta (2022)] was used by the facilitator (Supplementary material 3) but spontaneous conversations among the participants were encouraged. At least one member of the NIBLSE Leadership Team sat in on each focus group and, as disciplinary experts, occasionally interjected with questions. The focus groups were conducted on Zoom and recorded. The Zoom-derived transcripts of the recordings were analyzed using simple thematic analysis (Harding, 2013) to identify quotes that illustrated themes uncovered by the survey and the focus groups themselves. The quotes were edited to remove stammers, repetitions, and grammatical errors.

## Results

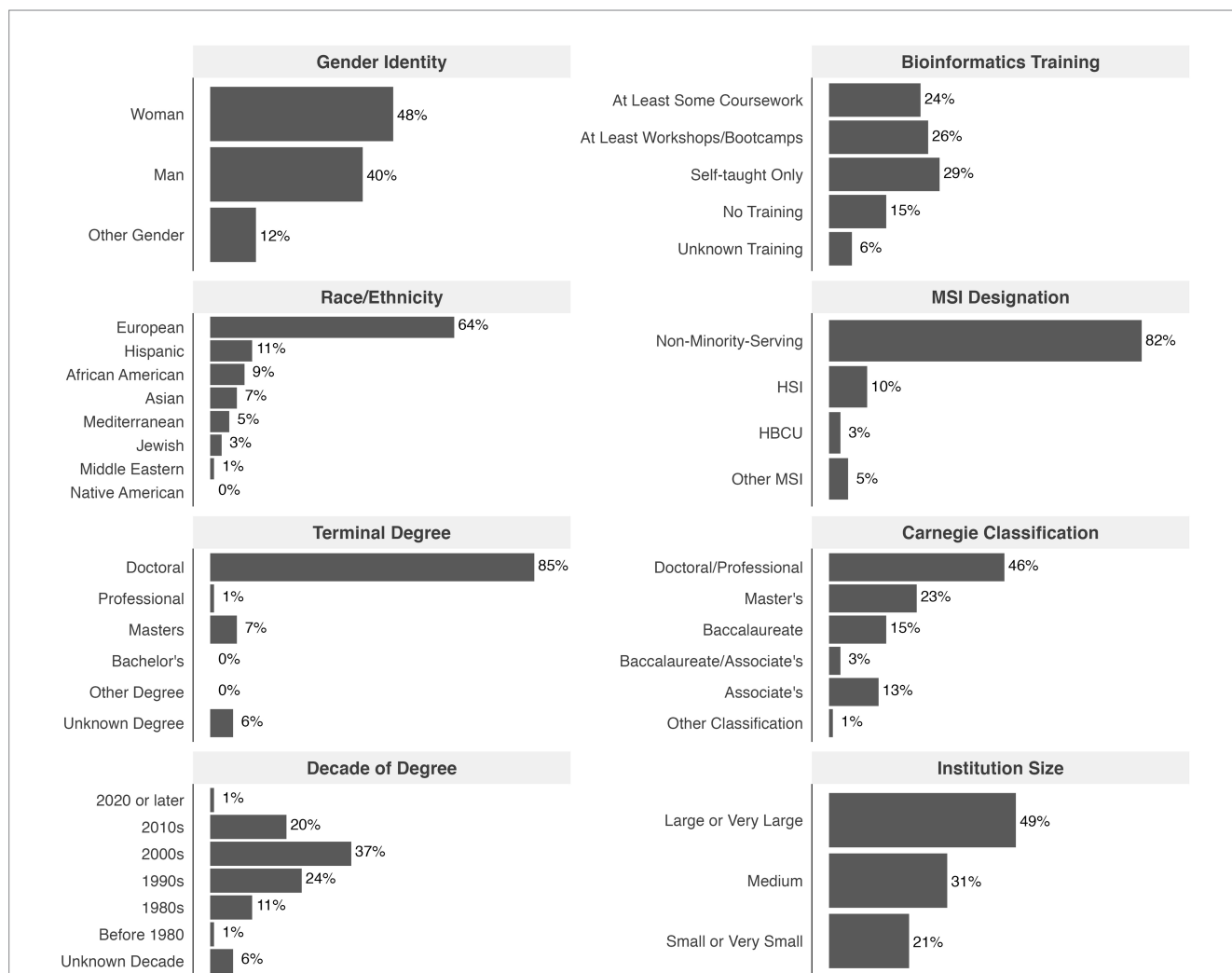
### The survey

#### A broad spectrum of educators participated in the 2022 NIBLSE survey

Altogether, 556 individuals responded to the current NIBLSE survey on barriers to integrating bioinformatics in undergraduate life science education. The 509 educators who agreed to participate in the study and taught undergraduate students were from diverse demographic groups and institution types, most often with some training in bioinformatics (Figure 1). The composite survey respondent is a woman or man of European descent with a PhD earned in 2000–2009, with some bioinformatics experience, working at a non-minority-serving, doctoral-granting, large institution.

#### More undergraduate bioinformatics education is needed, but instructors face barriers

While 90% of those who answered (435 of 484 respondents) agreed that more bioinformatics content is needed in undergraduate courses at their institutions, over 20% ( $n = 119$ ) of the 509 survey participants did not plan to integrate bioinformatics into their courses. The most frequently cited reasons were because teaching bioinformatics was not their responsibility or appropriate for their course ( $n = 51$ ), but many others said there were too many barriers ( $n = 31$ ). On the other hand, over 60% ( $n = 315$ ) of the survey participants currently teach some bioinformatics in a life science course (twice the rate seen in the 2016 survey), and 30% of those ( $n = 100$ ) within a classroom-based undergraduate research experience (CURE) or a summer undergraduate research experience (SURE).



**FIGURE 1** Summary of respondent demographics. Summary demographics shown as percentages of participants ( $n = 509$ , the total number of respondents teaching undergraduates who agreed to participate). Participants self-identified gender and race/ethnicity; European included Western European, Eastern European, and Scandinavian; Asian included East Asian, South Asian, and Southeast Asian. Participants could select multiple items for bioinformatics training and were categorized based on most formal training (e.g., if coursework and workshops/bootcamps marked, then indicated as "At Least Some Coursework").

Nevertheless, integrating bioinformatics into undergraduate life science courses continues to be a challenge. Over 70% (259 of 362 respondents) of the instructors teaching bioinformatics or who plan to do so reported facing barriers. In particular, instructors identified students lacking prerequisite skills and lack of time to restructure course content as minor to severe challenges more frequently than the other barriers (adj. value of  $p < 0.01$ ; Figure 2). In contrast, instructors cited not having the autonomy to add course content and lack of student interest less frequently as challenges (adj. value of  $p < 0.01$ ).

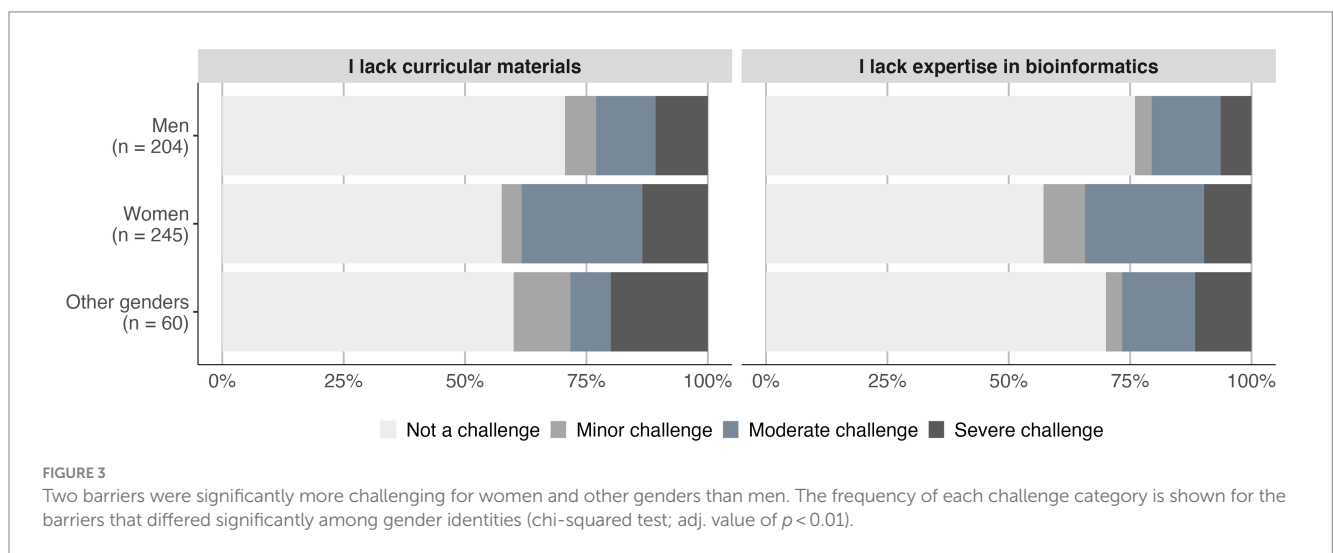
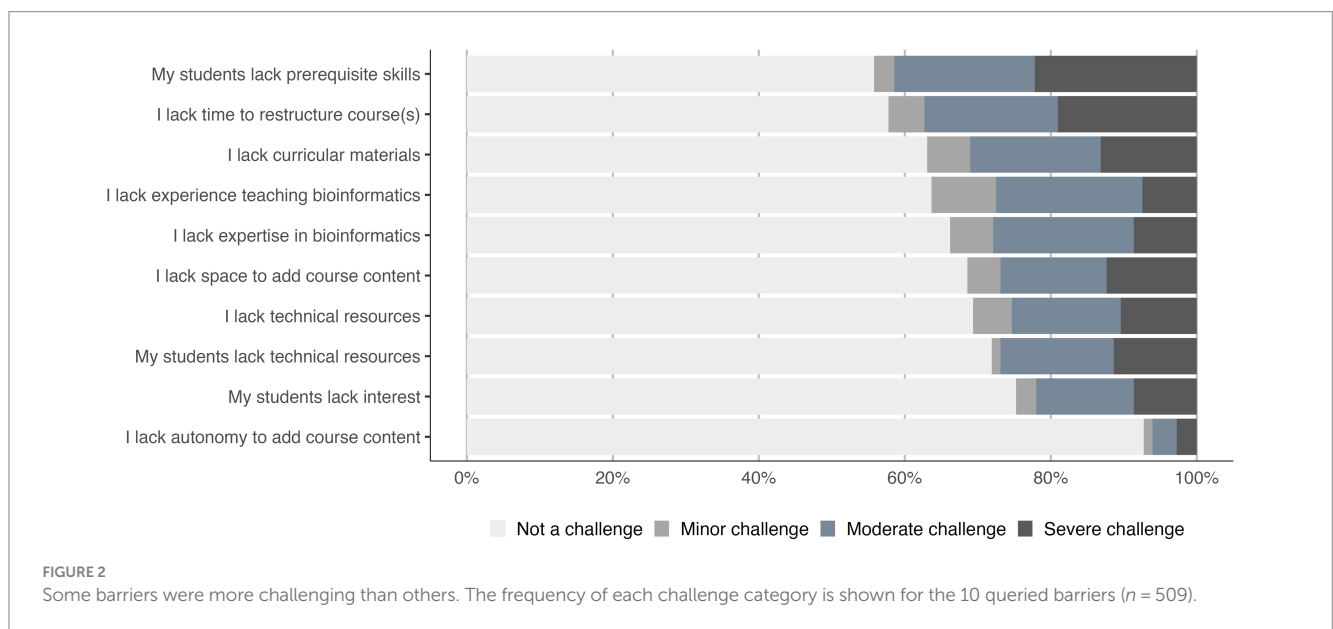
### Women experience more barriers than men

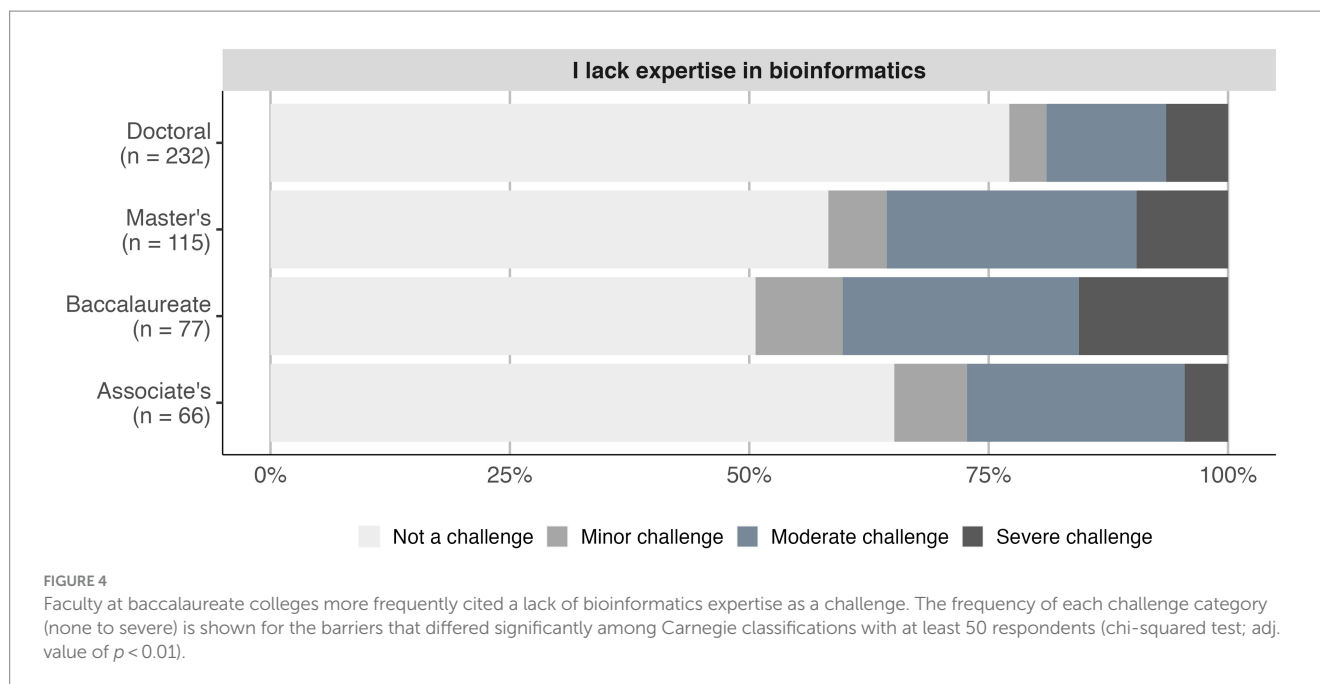
We previously reported in 2016 that women engaged in undergraduate life science education face barriers more frequently than men (Williams et al., 2019). We saw a similar pattern in this 2022 survey with women (and other gender identities) more likely to report barriers to integrating bioinformatics into their teaching than men (Men = 61%, Women = 78%, Others = 77%; value of  $p < 0.01$ ). For the

10 queried barriers, women and other gender identities more often identified lack of bioinformatics expertise and lack of curricular materials as challenges than men did (adj. value of  $p < 0.01$ ; Figure 3).

### The challenge of integrating bioinformatics varies with Carnegie classification

We reported in 2016 that the frequency of identified barriers diverged between instructors at different institution types (Williams et al., 2019). Similarly, in the 2022 survey we found a significant association between Carnegie classification and encountering barriers (value of  $p = 0.036$ ). Faculty at baccalaureate colleges reported facing barriers more frequently than the overall faculty average (82.3% vs. 71.4%), while faculty at doctoral institutions reported facing barriers less frequently (64.5%). In particular, the frequency that a lack of expertise in bioinformatics was identified as a challenge varied between faculty at different institution types (adj. value of  $p < 0.01$ ; Figure 4). No significant difference between institution types was observed for the other queried barriers.





### Some MSI faculty report a higher frequency of barriers than non-MSI faculty

Similar to findings in the 2016 survey, we found in the 2022 survey that while faculty at historically-black colleges and universities (HBCUs) or Hispanic-serving institutions (HSIs) more frequently reported barriers to integrating bioinformatics into their teaching than faculty elsewhere (HBCU = 82%,  $n = 17$ ; HSI = 81%,  $n = 37$ ; other-HSI = 65%,  $n = 20$ ; non-MSI = 73%,  $n = 306$ ), the differences were not statistically significant.

### Faculty with more recent degrees have more bioinformatics training but are equally likely to integrate bioinformatics in their courses

We reported in 2016 that faculty receiving their terminal degree more recently were more likely to have had formal bioinformatics training, but less likely to integrate bioinformatics than faculty who obtained their degrees a decade or more earlier (Williams et al., 2019). The 2022 survey also found that faculty with more recent terminal degrees were more likely than more experienced faculty to have some training, including one or more courses in bioinformatics (value of  $p < 0.001$ ; Figure 5A). Despite having more extensive formal training in bioinformatics, faculty with more recent terminal degrees somewhat more often (but not significantly) reported facing challenges to integrating bioinformatics (Figure 5B). Nevertheless, in the 2022 survey we did not see an association between the decade of receiving the terminal degree and integrating at least some bioinformatics instruction in life science classes (Figure 5C). Overall, however, faculty with some bioinformatics training were much more likely to integrate bioinformatics instruction (72%,  $n = 403$ ) than those with no training (12%,  $n = 76$ , value of  $p < 0.001$ ).

### URM and some MSI faculty report a higher frequency of barriers

The 2016 survey suggested that faculty who are members of underrepresented minority (URM) groups and/or at minority-serving

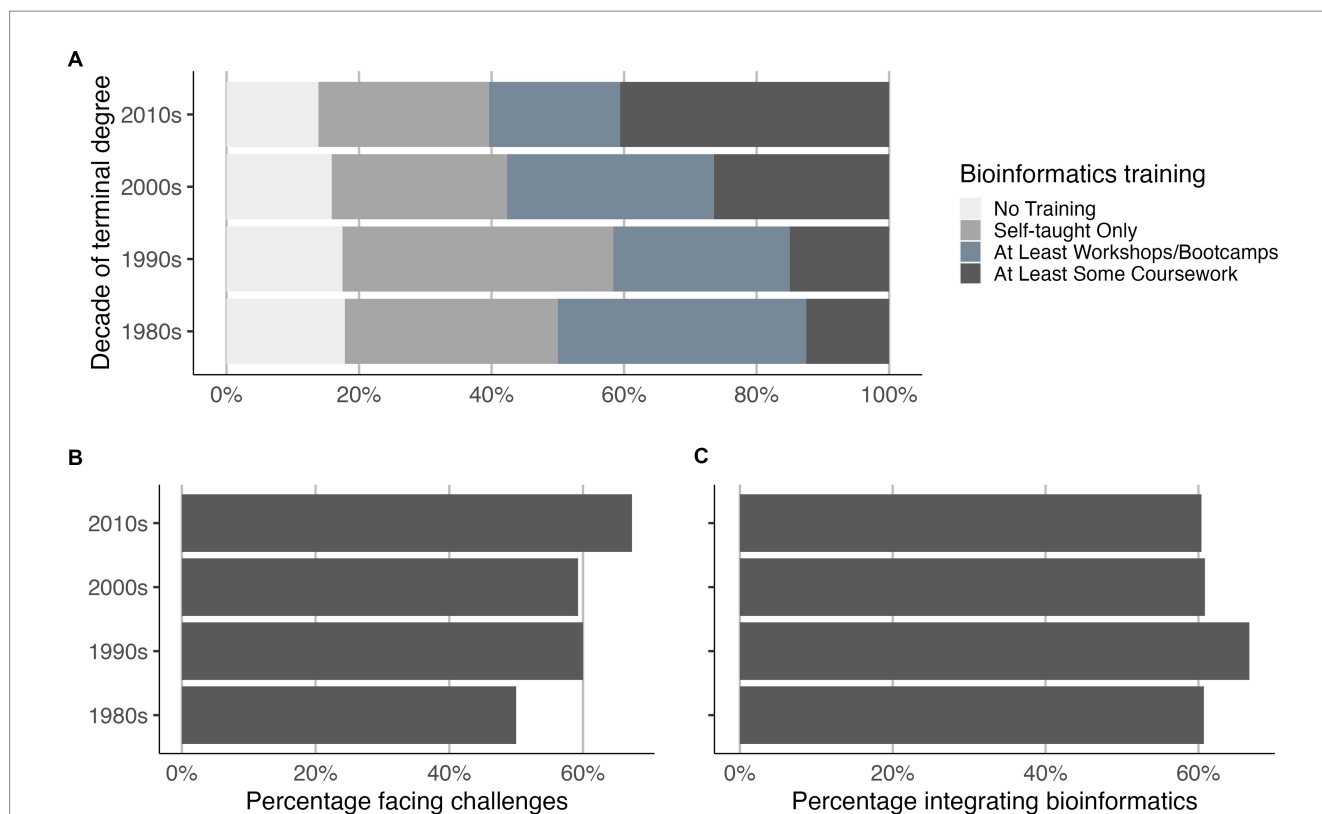
institutions (MSIs) experienced barriers at a higher rate compared to other participants (Williams et al., 2019). While URM faculty more frequently reported barriers to integrating bioinformatics into their teaching than non-URM faculty (URM = 81%,  $n = 77$ ; non-URM = 72%,  $n = 317$ ), the 2022 survey differences were not statistically significant. Similarly, while faculty at historically-black colleges and universities (HBCUs) or Hispanic-serving institutions (HSIs) more frequently reported barriers to integrating bioinformatics into their teaching than faculty elsewhere (HBCU = 82%,  $n = 17$ ; HSI = 81%,  $n = 37$ ; other-HSI = 65%,  $n = 20$ ; non-MSI = 73%,  $n = 306$ ), the differences were also not statistically significant. We explored these trends further in the focus groups (see below).

### The focus groups

In addition to the survey, we conducted 5 different focus groups with faculty who are attempting to integrate bioinformatics into their teaching. Using a semi-structured format, which included a common script (Supplementary material 3) but allowed the conversations to follow the flow of thoughts, we gained insights into the perceived barriers for faculty and their students. Although we initially arranged the focus groups based on the type of institution (minority-serving or not), common themes emerged across all 5 focus groups, and participants in a given focus group were generally in agreement when discussing the various questions.

### Is bioinformatics instruction important for students?

Each session started by asking the participants why they felt it was important to teach bioinformatics to their students. There were a range of responses, but most felt it was important for students to have the data analysis skills necessary for future work or advanced training. Several participants acknowledged they had no hard data regarding the extent to which students were more attractive as candidates for work or higher education, but



**FIGURE 5** Relationship of decade of terminal degree with bioinformatics training, barriers, and teaching bioinformatics. Faculty who received their terminal degrees more recently (A) more frequently had some course work in bioinformatics but were (B) more likely to report barriers to integrating bioinformatics. Nevertheless, respondents integrated bioinformatics at nearly the same frequency regardless of decade of terminal degree (C). The small number of instructors who obtained their terminal degree before 1980 or after 2019 were excluded from this analysis. Participants could select multiple items for bioinformatics training and were categorized based on most formal training (e.g., if coursework and workshops/bootcamps marked, then indicated as “At Least Some Coursework”).

**TABLE 2** Responses to the question, “Why teach bioinformatics?”

Primary theme	Secondary themes	Representative quote(s)
Faculty feel it is important for students to get skills and experience with data analysis.	Important for going into the workforce.	“It really helps them get jobs. That’s for sure.”
	Important for higher education/research.	“If you are a working molecular biologist, you can easily spend half your time (even if you are not doing genomics) on the computer, manipulating sequences and so on. So even a very basic level of bioinformatics, of doing a translation and figuring out a reading frame and then doing a multiple sequence alignment and things like that [are] sort of essential!” “It’s the future.”

their impression was that students with some data analysis skills had more opportunities available to them (Table 2). Several faculty members talked about the changes in professional health fields specifically as an impetus for including bioinformatics in their teaching.

**What barriers do the students exhibit?**

We then asked participants what barriers they encountered with respect to their students. Many said most students do not really understand what bioinformatics is or appreciate the advantages of knowing something about bioinformatics or data science (Table 3). However, several individuals also said that once students had some exposure, they understood better and “got into” it. On the other hand,

one participant in Focus Group 3 suggested that some of the savvy students are aware of bioinformatics and data science and are pushing for access to courses.

[There is] a lot of interest from around the university, but my classes are [currently] over full. [Some of] my students have that drive to really push their computer skills and have already recognized the economic need. But there is a huge proportion of our students who are graduating without ever having taken a formal computer-focused course. And so the phrase that I keep bringing up in meetings with administrators is, do you want our graduates to be able to visualize data in spreadsheets and have installed programs on a computer before? [There are] some



essential basic skills that are easily integrated into a more formalized education in my field, biology, and how [they] interact with computer systems.

Thus, some biology students are aware of the advantages of learning about bioinformatics, but such students appear to be in the minority, as this statement was not echoed by other members of this group or members of the other groups.

Some participants talked about technical and logistical barriers that students face, such as the difficulty students at their institutions have with regular access to the internet or computers. Reasons for this

difficulty included geographic location and socioeconomic status of the students.

### What prevents you from teaching bioinformatics?

In the course of listening to the focus group participants, we learned that many faculty, in fact, have some experience teaching bioinformatics but they still encountered a number of difficulties (Table 4).

First, many faculty are alone in trying to teach bioinformatics at their institution. On the other hand, some participants are working to develop programs at their institutions and are involved in training

TABLE 3 Responses to the question, “What barriers do the students exhibit?”

Primary theme	Secondary themes	Representative quote(s)
Many students lack understanding of what bioinformatics is or the advantages of knowing it	Students avoid math and computers.	<p>“The so-called typical biology student, their skills in anything related to math, anything related to numbers, [are] pretty weak. I see that all the way [from incoming undergraduates to] the graduate level. [There seems to be a] systematic avoidance of math [and] statistics. That’s why I like to tell them ‘Math is your friend,’ which I really believe, ‘if I can do it, you can do it.’”</p> <p>“It’s been hard to drag biology students into it, partially, because there’s very low interest. There’s also very low knowledge of what it is. It’s not something they have heard in high school. It’s not something we really are able to integrate into the curriculum with our limited faculty at an introductory level. So if they do see these sorts of tools, they are seeing it in their junior [year], maybe their senior [year], maybe their last semester.”</p>
	Students find it challenging to learn biology and bioinformatics “languages” at the same time.	Especially for students who are not native English speakers: “The regular language is a barrier, plus all the technical language.”
	Students have variable backgrounds in math and statistics.	“One of the barriers for me is the disparity in readiness of students. [Some are] maybe more interested in it, too. And then the ones who are more motivated, obviously are a lot more successful. They’ll work through the barriers of finding stuff. But it does become challenging, trying to keep everyone on the same page and at the same level.”
	Students may have difficulty in obtaining regular access to a computer or the internet	<p>Geographic location: “Being in basically a rural area.”</p> <p>Socioeconomic status: “Some students do not have a computer at home or an internet connection. There is a computer lab at their school and there are computer loans through school, [but this is] not fair compared to other students.”</p>

TABLE 4 Responses to the question, “What barriers do you have in attempts to incorporate bioinformatics into your teaching?”

Primary theme	Secondary themes	Representative quote(s)
Need for professional development	Many faculty are alone in trying to incorporate bioinformatics into their teaching at their institution. Others are working to develop programs at their institution and are involved in training others.	<p>“Some of what I’ve done has been for survival and trying to figure out methods I can use to be productive research-wise, while also teaching my students skills. But out of most of the faculty I’m the only one with any real bioinformatics training”</p> <p>“We created a community of 4 or 5 faculty members who are implementing CURE-like experiences that involve bioinformatics tools. [It’s] kind of organic and not super-organized, but certainly we create a little bit of a critical group there that is moving that forward, so that’s exciting for me.”</p>
	Keeping up with a fast-moving field.	<p>“But I know I’m behind and so, in order for me to get ahead and find out new things that I can do, and incorporate into both Gen Bio and advanced courses, I think professional development is going to be key. We need professional development training. And maybe that should also include staff that come with us for professional development training, because that way there’s more hands in the classroom to help”</p> <p>“[It’s hard to] know more or less the standard of what is currently done in bioinformatics; the tools [I used before] probably aren’t used anymore. Basically, I need an update.”</p>
Curriculum issues	Time in existing curriculum for bioinformatics content	“Where are you going to make room? What are you going to cut? You almost need a curriculum overhaul. If you have it in one class, great, so they do it for one semester, but then there’s nothing to build on. They might lose interest, or forget the skills.”

others. Second, although a number of people said they incorporated some bioinformatics, virtually everyone said that they found it difficult to keep up with this fast-moving field. Instructors have a range of abilities to teach bioinformatics; for example, some have programming expertise and can convey these skills to their students, while others incorporate only web-based tools that require no coding skills. In short, professional development is still important for many. Third, many expressed the thought that there are too many things to cover in the curriculum already. Finally, several participants mentioned the need to get administrative support in efforts to implement bioinformatics more broadly as well as the need for technical support, including computers with operating systems capable of running the necessary software.

## Discussion

We conducted a new survey to understand the barriers undergraduate life sciences faculty face in efforts to bring bioinformatics to the classroom and to determine to what extent the barriers observed in our 2016 study (Williams et al., 2019) are still present. We augmented the findings in the 2022 survey with focus groups composed of faculty from minority- and majority-serving institutions of different sizes and Carnegie classifications. The qualitative information gleaned from the focus groups corroborated the survey data.

As before, the vast majority of respondents agreed that bioinformatics is needed in life sciences curricula. We had more than 500 respondents in the current (2022) survey, about 40% of the number of respondents in the 2016 survey. Importantly, in the 2016 survey, only about 30% of participants indicated they were incorporating bioinformatics, while in this survey more than 60% of participants were doing so. This is a significant increase (chi-squared;  $p < 0.01$ ) in the ensuing 6 years. Nevertheless, barriers are still apparent. The majority of respondents reported facing barriers to integrating bioinformatics, although only a handful of barriers were identified as presenting moderate or severe challenges (Figure 2). In particular, instructors more frequently cited lack of experience teaching bioinformatics, time to restructure course content, and lack of student prerequisite skills. These sentiments were echoed in the focus group conversations (Tables 3, 4).

As in the 2016 survey (Williams et al., 2019), in the 2022 survey, women experienced more barriers than men (Figure 3). These findings suggest that women in particular would benefit from targeted professional development opportunities. In cooperation with QUBES,<sup>3</sup> NIBLSE will continue to offer bioinformatics-focused Faculty Mentoring Networks (FMNs) (Ryder et al., 2020; Kleinschmit et al., 2023). Facilitated by faculty with expertise in the topic under consideration, FMNs support groups of educators working over the course of a semester on a defined task. For example, NIBLSE-led FMNs have assisted faculty with integrating and adapting an existing bioinformatics exercise to their own classroom learning goals.

The 2016 survey suggested that URM faculty experience barriers at higher rates than non-URM faculty; however, the number of URM respondents was too low to detect statistical significance. Therefore, for this survey (as well as the focus groups), we targeted URM faculty (who composed approximately 23% of the names on the mailing list). In the

2022 study, we had about the same absolute number of URM respondents as in the 2016 study, but they were a larger percentage of the total. While we observed the same trend in the current survey, suggesting that URM faculty might experience barriers at higher rates, the results were not statistically significant again this time due to insufficient sample size. In addition, in the 2016 survey (Williams et al., 2019), faculty at MSIs showed a slight trend toward fewer barriers than faculty at majority institutions. When we examined this issue with the current survey data, there was a slight trend in the opposite direction, that MSI faculty experienced barriers at a slightly higher rate than faculty at majority institutions, although again, the results were not statistically significant. Nevertheless, making opportunities for professional development and curricular support available to these faculty would also be advantageous. For example, it was previously reported that faculty at MSIs are disadvantaged in efforts to seek federal funding (Chavela Guerra and Wilson, 2021), yet a recent report from the National Academies of Science, Engineering, and Medicine documents that MSIs are an underutilized resource for strengthening STEM fields in the US (National Academies of Sciences, Engineering, and Medicine, 2019).

When comparing faculty at different institution types, faculty at baccalaureate institutions experienced the most barriers, followed by those at master's institutions (Figure 4). Based on the focus groups, such faculty are likely to be alone in their efforts to incorporate bioinformatics into their teaching and would benefit from participating in networks outside of their home institution. One opportunity for network participation by undergraduate instructors is through FMNs (see above); others are cross-institutional course-based undergraduate research experience (CURE) and summer undergraduate research experience (SURE) initiatives incorporating bioinformatics components, such as the Genomics Education Partnership (GEP) (Elgin et al., 2017), Genome Solver (Mathur et al., 2019), Prevalence of Antibiotic-Resistance in the Environment (PARE) (Fuhrmeister et al., 2021; Bliss et al., 2023), SEA-PHAGES (Hanauer et al., 2017), Small World Initiative (SWI,<sup>4</sup>) and Tiny Earth (Hurley et al., 2021). These programs provide centralized support for faculty, including training and professional development, as well as access to computational resources, all of which address barriers identified in the survey and articulated in the focus groups, and may be of particular help to faculty at smaller institutions (Muth and McEntee, 2014). In addition, several of these projects promote development of bioinformatics skills in students as early as freshman year, allowing for growth of skills over the ensuing undergraduate years. While the finding that one-third of faculty integrating bioinformatics do so through some type of undergraduate research experience indicates that many faculty already perceive the value of this approach, expansion of such programs would further promote efforts to support bioinformatics education. Somewhat surprisingly, faculty at associate's colleges said they experienced fewer barriers than faculty at baccalaureate or master's institutions. The reasons for this are presently unclear but might reflect differences in content delivered, clientele served, and/or emphasis on practical skills for job seekers.

We found in the 2016 survey (Williams et al., 2019) that faculty who had received their terminal degree more recently had more formal training in bioinformatics than faculty who received their degrees in previous decades. At the same time, more recently awarded

<sup>3</sup> <https://qubeshub.org>

<sup>4</sup> <https://www.smallworldinitiative.org>

degree recipients experienced more barriers in attempts to implement bioinformatics into their teaching. These findings were replicated in the current survey (Figure 5A) and may reflect the more limited teaching experience of newer professors. Alternatively, these results might suggest that there are structural barriers, not explicitly investigated in this study, that hinder progress. It will be necessary to gather information from administrators and other stake-holders to examine such barriers in more detail, but this echoed comments from the focus group conversations that stated administrative support was necessary for expansion of bioinformatics offerings to students.

Arguably, the most difficult barrier to address and overcome for the integration of bioinformatics is lacking the autonomy to add course content (Figure 2). Addressing this issue is typically beyond the reach of individual instructors as changing departmental culture and acquiring administrative approval is necessary. Interestingly, however, respondents cited this potential barrier least frequently. The other barriers identified were those over which individual faculty have at least some autonomy whether it be seeking external assistance with curricula and/or professional development or reprioritizing individual commitments. Information from the focus groups suggested that students often believe they are not interested in bioinformatics upon initial introduction, but after they are exposed to the power of these concepts to address biological questions, as well as the appropriate tools and techniques, they tend to be very interested in further exploration. Likewise, although prerequisite student skills are perceived to be a barrier, incremental exposure to introductory quantitative techniques through a bioinformatics lens by individual instructors could be an effective mechanism to build needed skills.

In summary, in the ensuing interval since our first survey in 2016, notable progress has been made. About twice as many faculty are incorporating at least some bioinformatics learning into their teaching. However, as with our previous results, undergraduate faculty nearly uniformly signaled the need for training in bioinformatics and help in developing curricular resources to teach bioinformatics as revealed in both the survey data and the conversations in the focus groups. NIBLSE has been working in both arenas to address these barriers (Ryder et al., 2020; Kleinschmit et al., 2023). As mentioned previously, other groups have also made efforts to address these barriers, including GEP, SEA-PHAGES, SWI, Tiny Earth, PARE, etc. These groups also have curriculum available for use. The combination of these ongoing initiatives provides a robust framework for future efforts to ensure that tomorrow's researchers possess the necessary skills and attitudes for conducting research in the 21st century.

## Data availability statement

The datasets presented in this study can be found in online repositories. The names of the repository/repositories and accession number(s) can be found in the article/Supplementary material.

## Ethics statement

The studies involving humans were approved by the University of Florida IRB and the Georgetown University IRB. The studies were conducted in accordance with the local legislation and institutional

requirements. The participants provided their written informed consent to participate in this study.

## Author contributions

JD: Conceptualization, Data curation, Methodology, Writing – original draft, Writing – review & editing. WM: Data curation, Formal analysis, Funding acquisition, Methodology, Visualization, Writing – original draft, Writing – review & editing. SG: Conceptualization, Methodology, Writing – review & editing. AK: Conceptualization, Methodology, Writing – original draft, Writing – review & editing. MM: Investigation, Writing – review & editing. MP: Conceptualization, Funding acquisition, Methodology, Writing – original draft, Writing – review & editing. ET: Conceptualization, Funding acquisition, Writing – review & editing. JW: Conceptualization, Formal analysis, Methodology, Software, Visualization, Writing – original draft, Writing – review & editing. BM: Writing – review & editing. AR: Conceptualization, Funding acquisition, Investigation, Methodology, Supervision, Writing – original draft, Writing – review & editing.

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## Conflict of interest

The authors declare that the research was conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest.

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## Supplementary material

The Supplementary material for this article can be found online at: <https://www.frontiersin.org/articles/10.3389/feduc.2023.1317191/full#supplementary-material>

## References

- Altman, R. B. (1998). A curriculum for bioinformatics: the time is ripe. *Bioinformatics* 14, 549–550. doi: 10.1093/bioinformatics/14.7.549
- Attwood, T. K., Blackford, S., Brazas, M. D., Davies, A., and Schneider, M. V. (2019). A global perspective on evolving bioinformatics and data science training needs. *Brief. Bioinform.* 20, 398–404. doi: 10.1093/bib/bbx100
- Benjamini, Y., and Hochberg, Y. (1995). Controlling the false discovery rate: a practical and powerful approach to multiple testing. *J. R. Stat. Soc. Ser. B* 57, 289–300.
- Bliss, S. S., Abraha, E. A., Fuhrmeister, E. R., Pickering, A. J., and Bascom-Slack, C. A. (2023). Learning and stem identity gains from an online module on sequencing-based surveillance of antimicrobial resistance in the environment: an analysis of the Pare-Seq curriculum. *PLoS One* 18:E0282412. doi: 10.1371/journal.pone.0282412
- Carvalho, B. S., and Rustici, G. (2013). The challenges of delivering bioinformatics training in the analysis of high-throughput data. *Brief. Bioinform.* 14, 538–547. doi: 10.1093/bib/bbt018
- Chavela Guerra, R. C., and Wilson, C. (2021). From lack of time to stigma: barriers facing faculty at minority-serving institutions pursuing federally funded research. *ASEE Virtual Ann. Conf.* Available at: <https://par.nsf.gov/biblio/10304434>
- Couch, S. P., Bray, A. P., Ismay, C., Chasnovski, E., Baumer, B. S., and Çetinkaya-Rundel, M. (2021). Infer: an R package for tidyverse-friendly statistical inference. *J. Open Source Softw.* 6:3661. doi: 10.21105/joss.03661
- Dillman, D. A., Smyth, J. D., and Christian, L. M. (2014). *Internet, Phone, Mail, and Mixed Mode Surveys: the Tailored Design Method*. Hoboken, NJ: John Wiley & Sons, Inc.
- Dinsdale, E., Elgin, S. C., Grandgenett, N., Morgan, W., Rosenwald, A., Tappich, W., et al. (2015). NIBLSE: a network for integrating bioinformatics into life sciences education. *CBE Life Sci. Educ.* 14, 1–4. doi: 10.1187/cbe.15-06-0123
- Drew, J. C., Grandgenett, N., Dinsdale, E. A., Vazquez Quinones, L. E., Galindo, S., Morgan, W. R., et al. (2021). There is more than multiple choice: crowd-sourced assessment tips for online, hybrid, and face-to-face environments. *J. Microbiol. Biol. Educ.* e00205-21:22. doi: 10.1128/jmbe.00205-21
- Elgin, S. C. R., Hauser, C., Holzen, T. M., Jones, C., Kleinschmit, A., Leatherman, J., et al. (2017). The Gep: crowd-sourcing big data analysis with undergraduates. *Trends Genet.* 33, 81–85. doi: 10.1016/j.tig.2016.11.004
- Elgin, S. C., Hays, S., Mingo, V., Shaffer, C. D., and Williams, J. (2021). Building back more equitable stem education: teach science by engaging students in doing science. *bioRxiv*. doi: 10.1101/2021.06.01.446616
- Fuhrmeister, E. R., Larson, J. R., Kleinschmit, A. J., Kirby, J. E., Pickering, A. J., and Bascom-Slack, C. A. (2021). Combating antimicrobial resistance through student-driven research and environmental surveillance. *Front. Microbiol.* 12:577821. doi: 10.3389/fmicb.2021.577821
- Gauthier, J., Vincent, A. T., Charette, S. J., and Derome, N. (2019). A brief history of bioinformatics. *Brief. Bioinform.* 20, 1981–1996. doi: 10.1093/bib/bby063
- Hanauer, D. I., Graham, M. J., Sea, P., Betancur, L., Bobrownicki, A., Cresawn, S. G., et al. (2017). An Inclusive Research Education Community (IREC): impact of the SeaPhages program on research outcomes and student learning. *Proc. Natl. Acad. Sci. U. S. A.* 114, 13531–13536. doi: 10.1073/pnas.1718188115
- Handelsman, J., Elgin, S., Estrada, M., Hays, S., Johnson, T., Miller, S., et al. (2022). Achieving stem diversity: fix the classrooms. *Science* 376, 1057–1059. doi: 10.1126/science.abn9515
- Harding, J. (2013). *Qualitative data analysis from start to finish*. Thousand Oaks, CA: Sage Publications.
- Hurley, A., Chevrette, M. G., Acharya, D. D., Lozano, G. L., Garavito, M., Heinritz, J., et al. (2021). Tiny earth: a big idea for stem education and antibiotic discovery. *mBio* 12:e03432-20. doi: 10.1128/mBio.03432-20
- Isik, E. B., Brazas, M. D., Schwartz, R., Gaeta, B., Palagi, P. M., Van Gelder, C. W. G., et al. (2023). Grand challenges in bioinformatics education and training. *Nat. Biotechnol.* 41, 1171–1174. doi: 10.1038/s41587-023-01891-9
- Kleinschmit, A. J., Rosenwald, A., Ryder, E. F., Donovan, S., Murdoch, B., Grandgenett, N. F., et al. (2023). Accelerating stem education reform: linked communities of practice promote creation of open educational resources and sustainable professional development. *IJ Stem Ed* 10:16. doi: 10.1186/s40594-023-00405-y
- Kleinschmit, A. J., Ryder, E. F., Kerby, J. L., Murdoch, B., Donovan, S., Grandgenett, N. F., et al. (2021). Community development, implementation, and assessment of a NIBLSE bioinformatics sequence similarity learning resource. *PLoS One* 16:E0257404. doi: 10.1371/journal.pone.0257404
- Machluf, Y., and Yarden, A. (2013). Integrating bioinformatics into senior high school: design principles and implications. *Brief. Bioinform.* 14, 648–660. doi: 10.1093/bib/bbt030
- Mathur, V., Arora, G. S., McWilliams, M., Russell, J., and Rosenwald, A. G. (2019). The genome solver project: faculty training and student performance gains in bioinformatics. *J. Microbiol. Biol. Educ.* 20:20.1.4. doi: 10.1128/jmbe.v20i1.1607
- Muth, T. R., and Mcentee, C. M. (2014). Undergraduate urban metagenomics research module. *J. Microbiol. Biol. Educ.* 15, 38–40. doi: 10.1128/jmbe.v15i1.645
- National Academies of Sciences, Engineering, and Medicine. (2019). *Minority serving institutions: America's underutilized resource for strengthening the stem workforce*. Washington, DC: The National Academies Press.
- Niepielko, M. G., and Shumskaya, M. (2021). Early requirement for bioinformatics in undergraduate biology curricula. *Front. Bioinform.* 1:656531. doi: 10.3389/fbinf.2021.656531
- Porter, S. G., and Smith, T. M. (2019). Bioinformatics for the masses: the need for practical data science in undergraduate biology. *OMICS* 23, 297–299. doi: 10.1089/omi.2019.0080
- R Core Team (2021). *R: a language and environment for statistical computing*. Vienna, Austria: R Foundation For Statistical Computing
- Rocha, M., Massarani, L., Souza, S. J., and Vasconcelos, A. T. R. (2022). The past, present and future of genomics and bioinformatics: a survey of Brazilian scientists. *Genet. Mol. Biol.* 45:E20210354. doi: 10.1590/1678-4685-gmb-2021-0354
- Ryder, E. F., Morgan, W. R., Sierk, M., Donovan, S. S., Robertson, S. D., Orndorf, H. C., et al. (2020). Incubators: building community networks and developing open educational resources to integrate bioinformatics into life science education. *Biochem. Mol. Biol. Educ.* 48, 381–390. doi: 10.1002/bmb.21387
- Saldana, J., and Omasta, M. (2022). *Qualitative research: analyzing life*. Thousand Oaks, CA: Sage Publications
- The Genomic Data Science Community Network (2022). Diversifying the genomic data science research Community. *Genome Res.* 32, 1231–1241. doi: 10.1101/gr.276496.121
- Toby, I., Williams, J. J., Lu, G., Cai, C., Crandall, K. A., Dinsdale, E. A., et al. (2022). Making change sustainable: Network for Integrating Bioinformatics into Life Sciences Education (NIBLSE) meeting review. *Coursesource* 9. doi: 10.24918/cs.2022.10
- Wickham, H., Averick, M., Bryan, J., Chang, W., McGowan, L. D., Francois, R., et al. (2019). Welcome to the tidyverse. *J. Open Source Softw.* 4:1686. doi: 10.21105/joss.01686
- Williams, J. J., Drew, J. C., Galindo-Gonzalez, S., Robic, S., Dinsdale, E., Morgan, W. R., et al. (2019). Barriers to integration of bioinformatics into undergraduate life sciences education: a national study of us life sciences faculty uncover significant barriers to integrating bioinformatics into undergraduate instruction. *PLoS One* 14:E0224288. doi: 10.1371/journal.pone.0224288
- Wilson Sayres, M. A., Hauser, C., Sierk, M., Robic, S., Rosenwald, A. G., Smith, T. M., et al. (2018). Bioinformatics core competencies for undergraduate life sciences education. *PLoS One* 13:e0196878. doi: 10.1371/journal.pone.0196878