OMB No.: 0925-0642
Expiration Date: 05/31/2020

Public reporting burden for this collection of information is estimated to average 5 minutes per response, including the time for reviewing instructions, searching existing data sources, gathering and maintaining the data needed, and completing and reviewing the collection of information. An agency may not conduct or sponsor, and a person is not required to respond to, a collection of information unless it displays a currently valid OMB control number. Send comments regarding this burden estimate or any other aspect of this collection of information, including suggestions for reducing this burden to: NIH, Project Clearance Branch, 6705 Rockledge Drive, MSC 7974, Bethesda, MD 20892-7974, ATTN: PRA (0925-0642). Do not return the completed form to this address.

Survey Questions

1. What is your field of research? Please check all that apply.

- [ ] Cancer Genomics
- [ ] Bioinformatics
- [ ] Genetic/Genomic Epidemiology
- [ ] Statistical Genomics

Other (please specify)

2. What were your reasons for attending past SeqSPACE webinars? Please check all that apply.

- [ ] I was interested in the webinar speaker
- [ ] I conducted research in the topic highlighted in the webinar
- [ ] I wanted to know more about the topic highlighted in the webinar

Other (please specify)

3. On average, how valuable were all the SeqSPACE webinars you attended? 1=Not at all valuable to 5=Extremely valuable

- ○ 1 - Not at all valuable
- ○ 2 - Not so valuable
- ○ 3 - Somewhat valuable
- ○ 4 - Very valuable
- ○ 5 - Extremely valuable

4. Have you used the information from these webinars in your work?

- ○ Yes
- ○ No

If YES, please explain:

5. Have you ever contacted a presenter from a SeqSPACE webinar after their presentation?

- ◯ Yes
- ◯ No

6. How likely are you to recommend this series to colleagues in your field? 0 (Not likely at all) to 10 (Extremely Likely)

0	1	2	3	4	5	6	7	8	9	10
◯	◯	◯	◯	◯	◯	◯	◯	◯	◯	◯

7. What topics would you like to see addressed in future SeqSPACE webinars?

8. Do you have any additional comments about the SeqSPACE webinar series?