

**iDASH Privacy & Security Workshop 2021 - Secure Genome Analysis Competition**  
**Track 1**  
**Frequently Asked Questions**

**NOTE:** Please send your questions to [tskuo@ucsd.edu](mailto:tskuo@ucsd.edu), and we will reply here as soon as we can. Since we may post your questions here, please try to generalize your questions and do not reveal your identity and methods in the questions. We may also modify or merge your questions to increase the generalizability.

1. Are you going to (1) only insert records for a time, and then only query them (which is written in the subject), or (2) have a mixed workload of insertions and queries?

We are planning to evaluate using (1).

2. Is it safe to assume that preferences always appear in the same order in the records?

Yes, the preferences in the test data will appear in the same order.

3. Would it be ok to return an array that contains patient IDs equal 0 at its end (which should then be ignored)?

Unfortunately no, the returned patient IDs would need to be exactly the same as expected (i.e., 0 or other “null” IDs should be excluded).

4. Do you expect patient IDs to be returned in a precise order?

No, the returned patient IDs can be in any order.