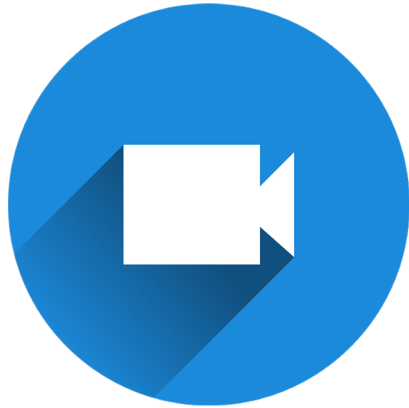


Submitting, Finding and Downloading Raw Sequencing Data with ENA

Geert van Geest

Interfaculty Bioinformatics Unit, UniBe
Training group, SIB

Course etiquette



Video on when
possible



Mute when
not speaking

Use your **first** and **last** name as your Zoom ID
Right-click on yourself > Rename..

Trainers/organisers

- **Patricia Palagi:** Manager training group at SIB
- **Geert van Geest:** trainer at SIB/bioinformatician at IBU Bern

Learning outcomes

- Understand the **main data structures** underlying ENA
- Find an **appropriate portal** for submitting and finding raw NGS data
- **Find** and **download** raw sequencing **data** from SRA
- Successfully **submit** raw sequencing data to ENA

Learning experiences

- Lectures
- Polls
- Exercises

Questions

Communication

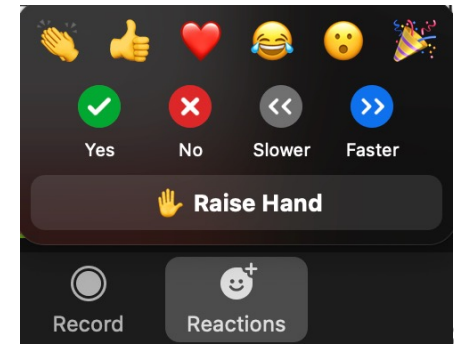
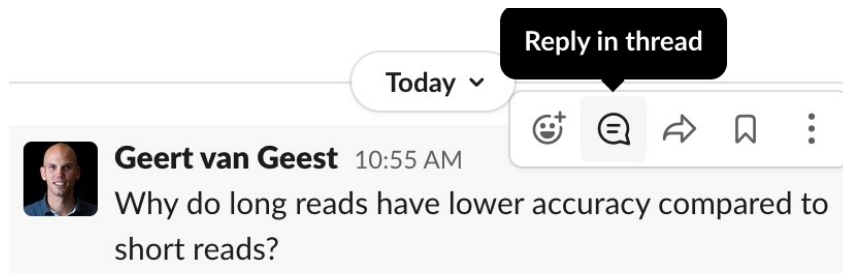
- Course website:

<https://sib-swiss.github.io/ena-seqdata-training/>

- Google docs
- Slack

Asking questions

- During lectures: zoom functionality
- Personal interest questions: [#background](#)
- During exercises: [#q-and-a](#) on slack



Introduction round

- Current job/study
- Why you are joining this course