

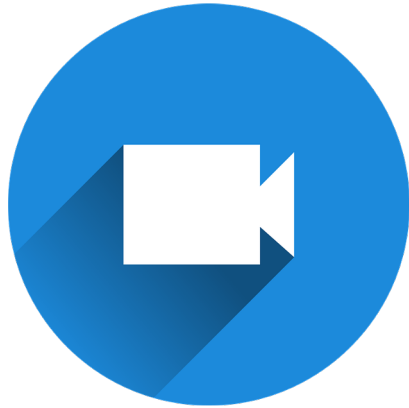
# Submitting, Finding and Downloading Raw Sequencing Data with ENA

Geert van Geest

Interfaculty Bioinformatics Unit, UniBe  
Training group, SIB

# Course etiquette

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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

- **Valeria di Cola:** Training coordinator at SIB
- **Maira Ihsan:** Guest lecturer from EBI
- **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

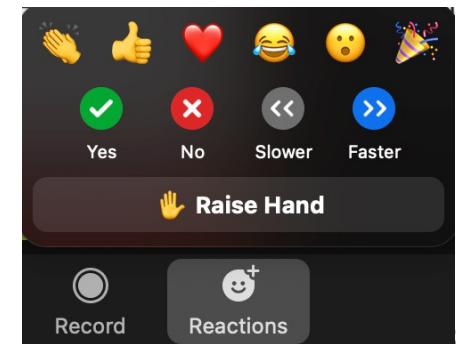
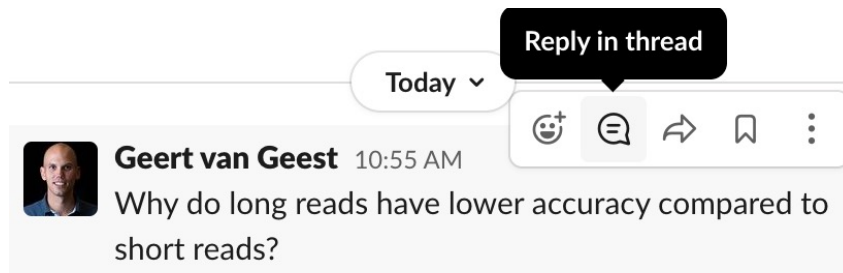
- Course website:

<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