NGS - variant analysis

Geert van Geest Interfaculty Bioinformatics Unit, UniBe Training group, SIB

Learning outcomes

- Understand important aspects of NGS and read alignment for variant analysis
- Perform a read alignment ready for variant analysis
- Perform variant calling according to GATK best practices
- Perform a variant annotation

Learning experiences

- Lectures
- Quiz questions
- Exercises

Communication

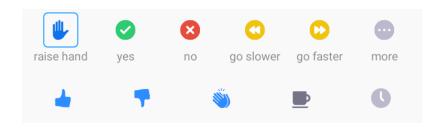
Course website:

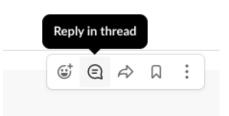
https://sib-swiss.github.io/NGS-variants-training/

- Slack
- Google docs

Asking questions

- During lectures: zoom functionality
- Personal interest questions: #background
- During exercises:
 - #q_and_a on slack
 - if really stuck: no button in zoom
 - if finished: yes button in zoom





Get to know each other

- Write in the google doc (5 minutes):
 - Three keywords about yourself
 - Why you are joining this course, and what you want to learn
- You will discuss them in breakout rooms afterwards (15 minutes)
 - Introduce yourself based on what you've written in the doc