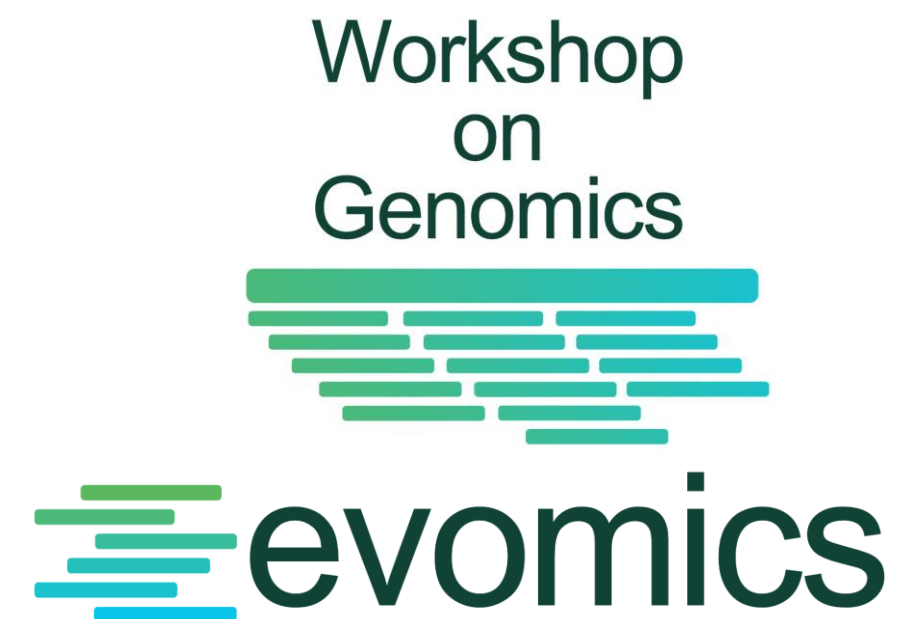


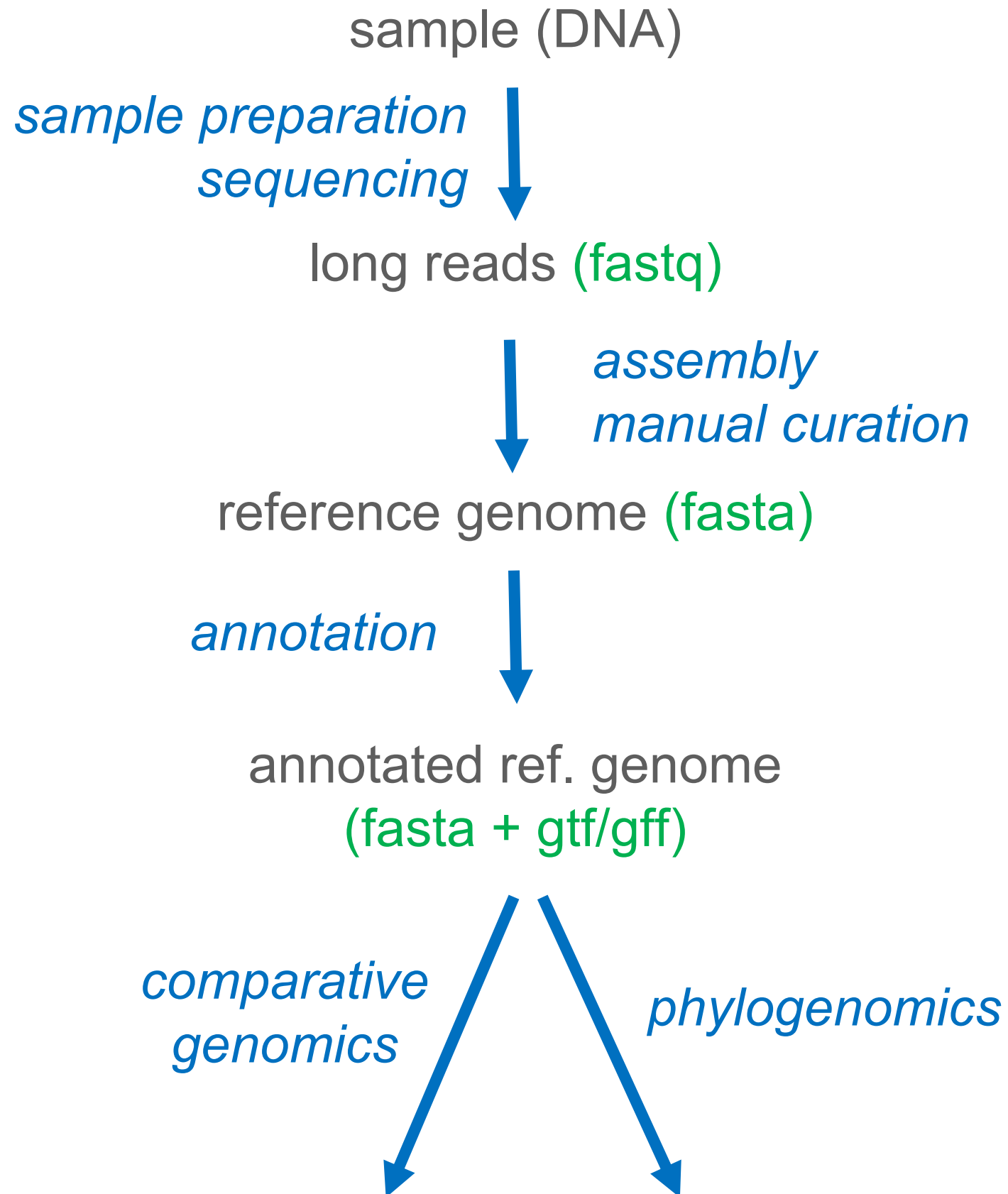


Workshop on Genomics 2026

Český Krumlov, 11th - 24th January

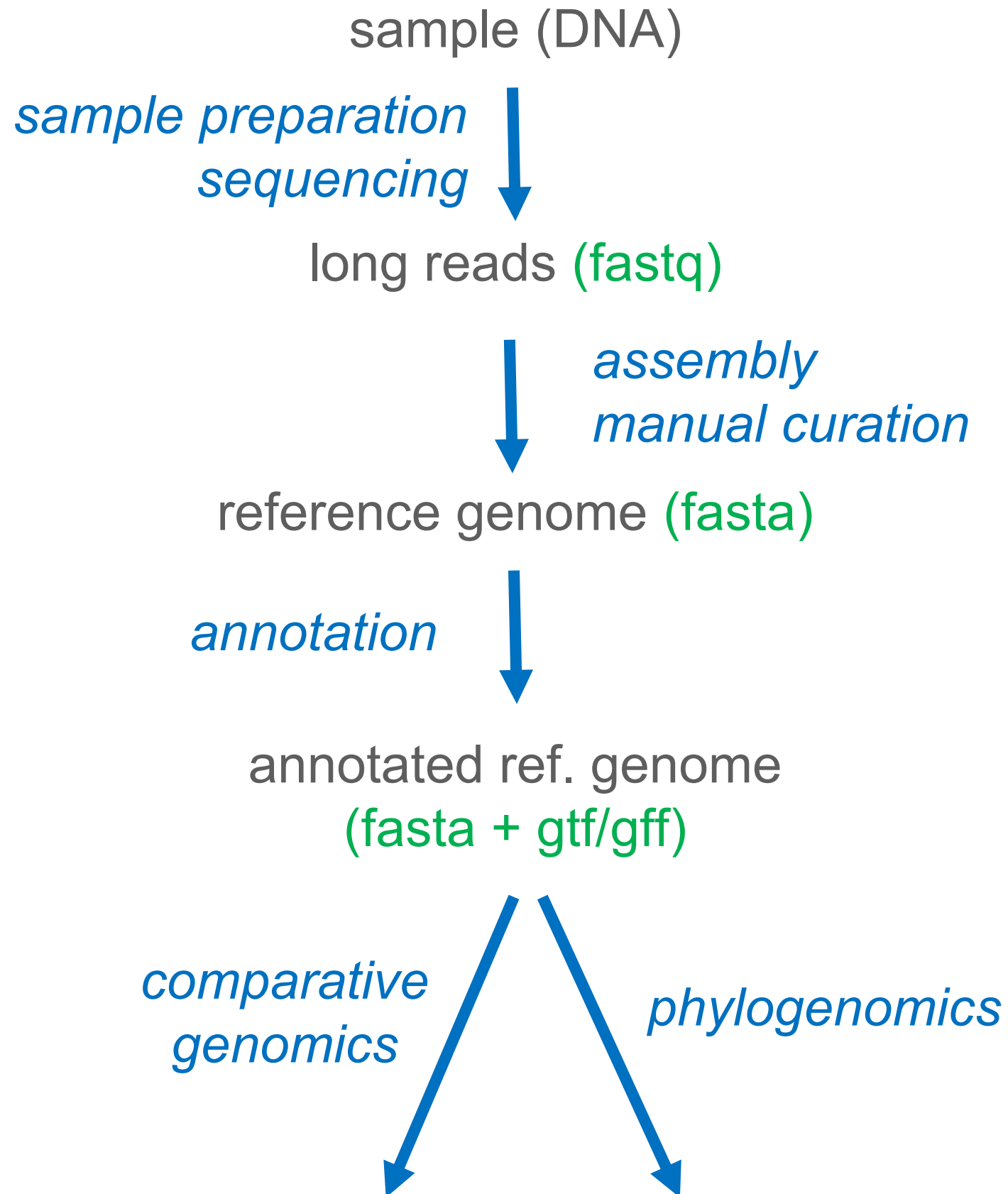


- Wrap-up
- Feedback (~20 mins)
- Round Table
- Prizes!
- T-shirts & Certificates





Olga Vinnere
Pettersson





Olga Vinnere
Pettersson

sample (DNA)
*sample preparation
sequencing* ↓
long reads (*fastq*)



Camille Marchet &
Antoine Limasset



Marcela Uliano-Silva

↓ *assembly
manual curation*
reference genome (*fasta*)

annotation

↓
annotated ref. genome
(*fasta + gtf/gff*)



Camilla Santos

*comparative
genomics*

phylogenomics



Olga Vinnere
Pettersson

sample (DNA)
*sample preparation
sequencing* ↓
long reads (*fastq*)



Camille Marchet &
Antoine Limasset



Marcela Uliano-Silva



Katharina Hoff

*assembly
manual curation* ↓
reference genome (*fasta*)

annotation ↓

annotated ref. genome
(*fasta + gtf/gff*)



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*comparative
genomics*

phylogenomics



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(*fasta + gtf/gff*)



Camilla Santos

*comparative
genomics*

phylogenomics



Nathan Clark



Olga Vinnere
Pettersson

sample (DNA)
*sample preparation
sequencing* ↓
long reads (*fastq*)



Camille Marchet &
Antoine Limasset



Marcela Uliano-Silva



Katharina Hoff

*assembly
manual curation* ↓
reference genome (*fasta*)

annotation ↓

annotated ref. genome
(*fasta + gtf/gff*)



Nathan Clark

*comparative
genomics*

phylogenomics



Rosa Fernández

sample (DNA)

*sample preparation
sequencing*



short reads (fastq)

reference genome (fasta)



alignment/mapping



alignment (sam/bam)

variant calling



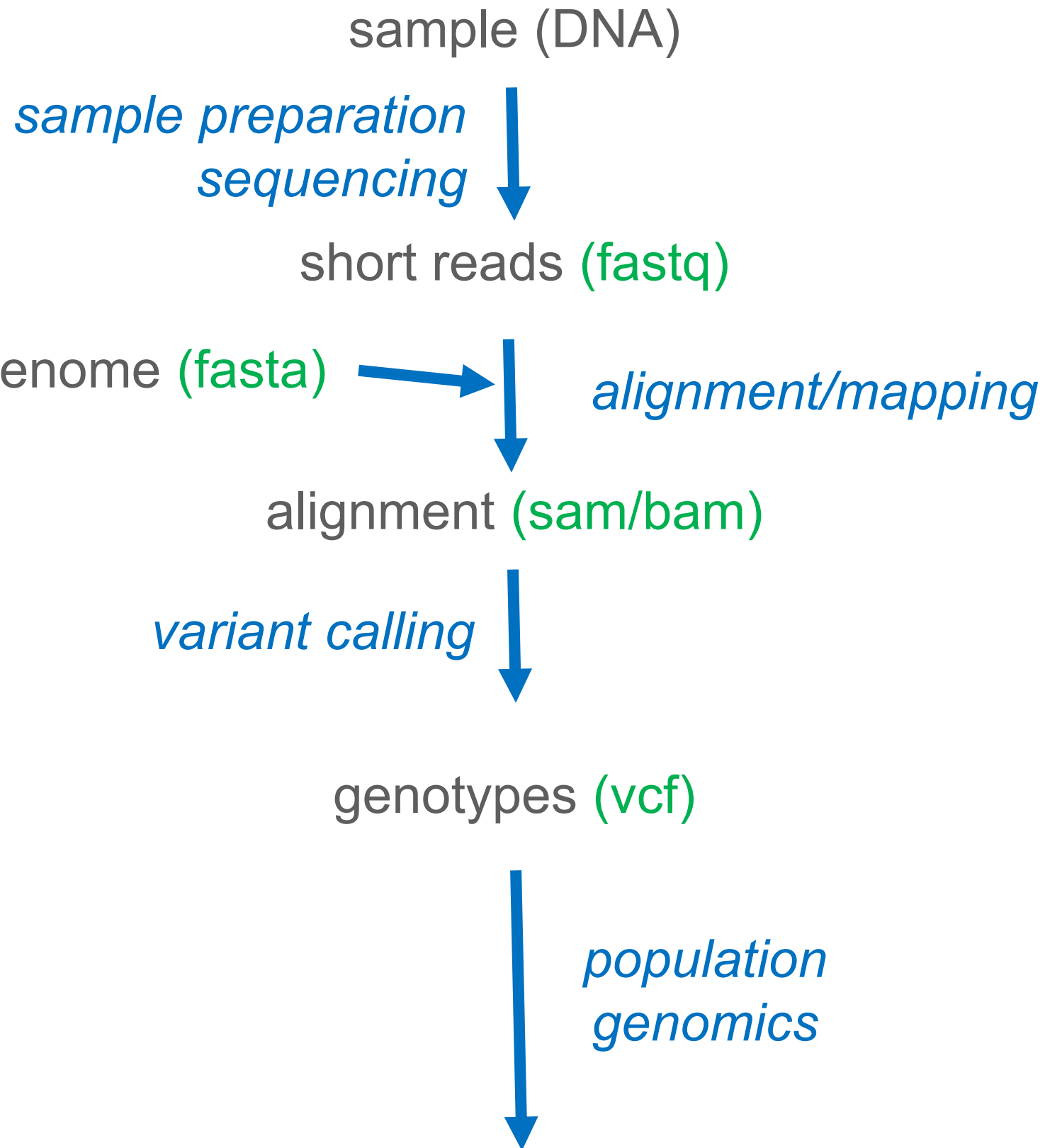
genotypes (vcf)

*population
genomics*



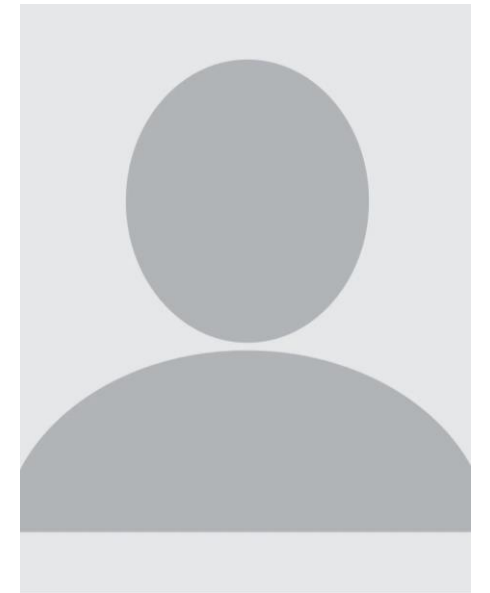
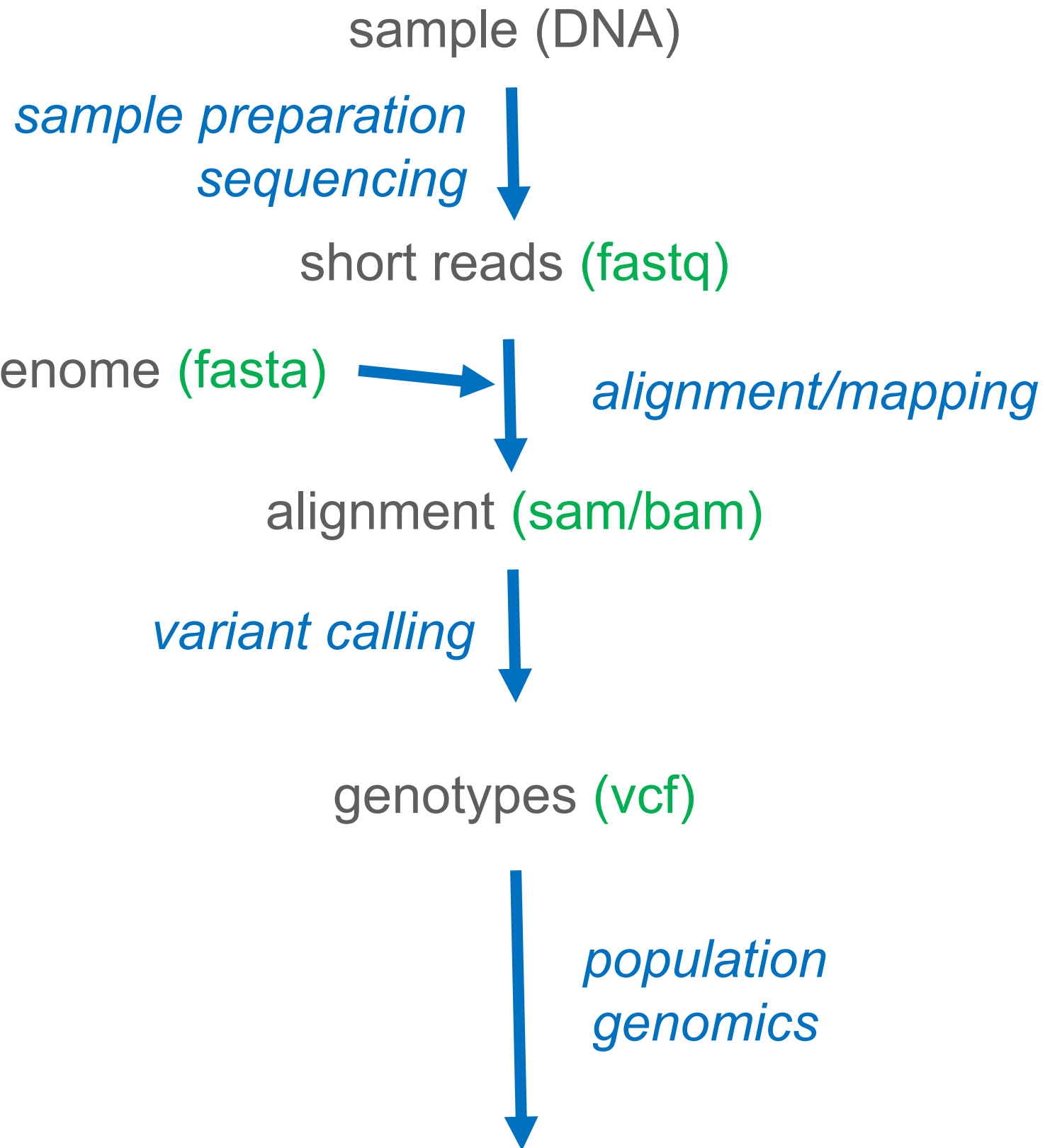


Olga Vinnere
Pettersson





Olga Vinnere
Pettersson



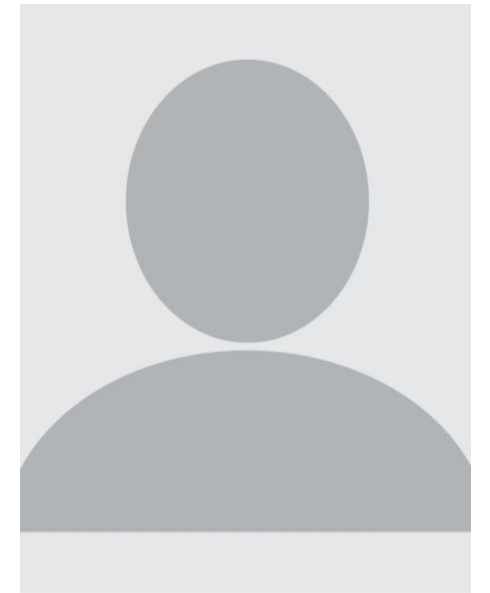
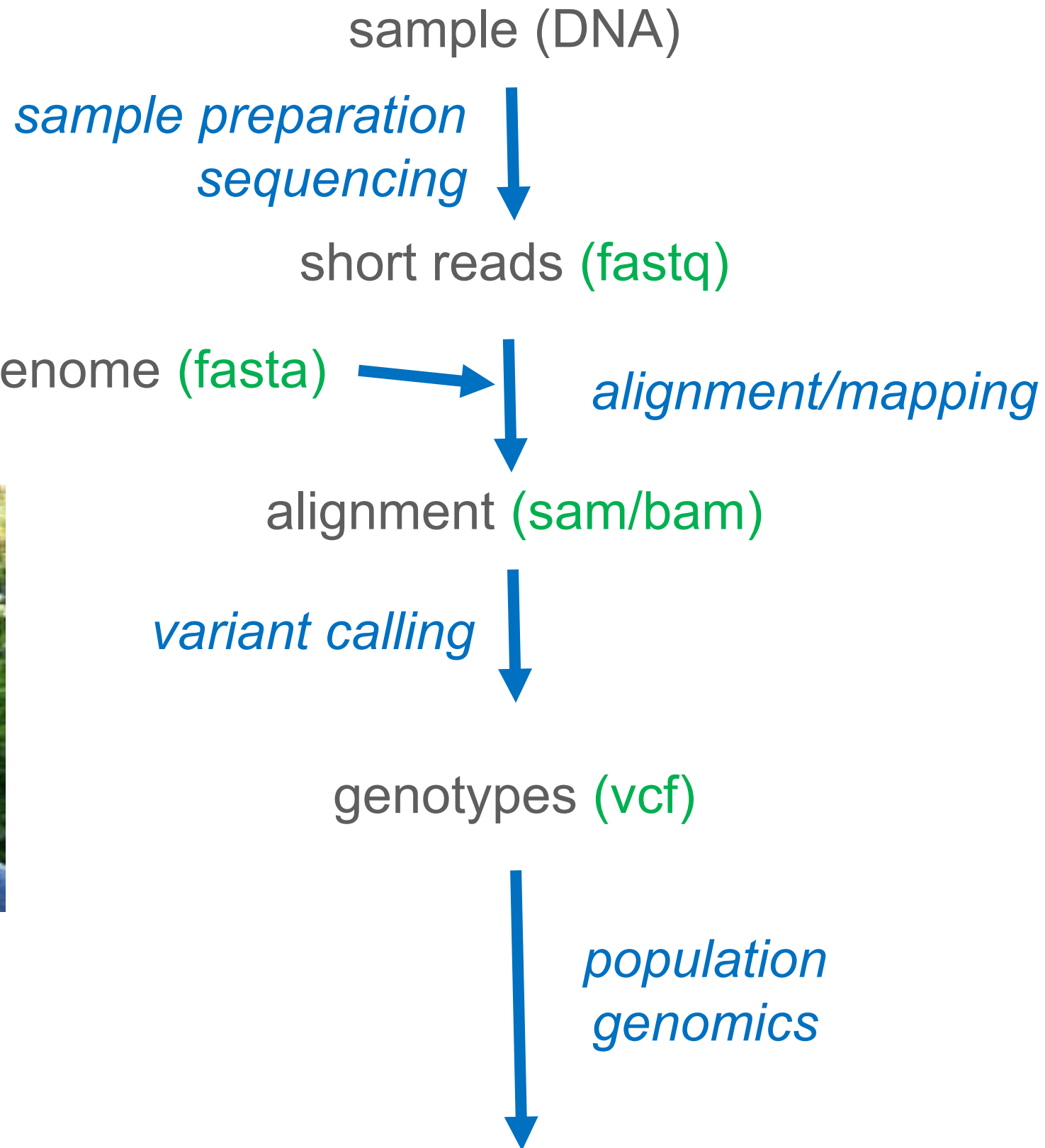
Benjamin Buchfink



Olga Vinnere
Pettersson



Petr Danecek



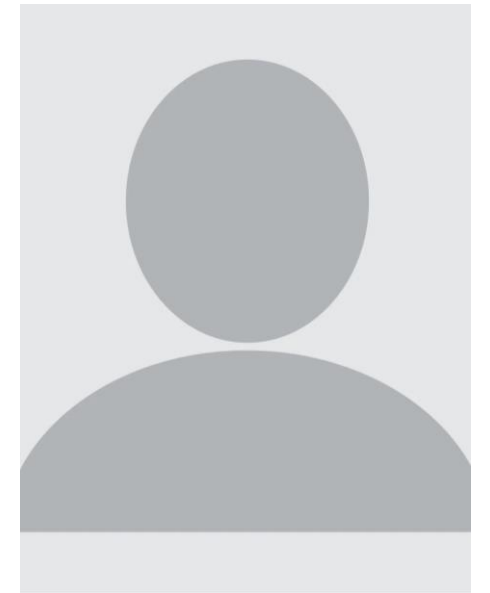
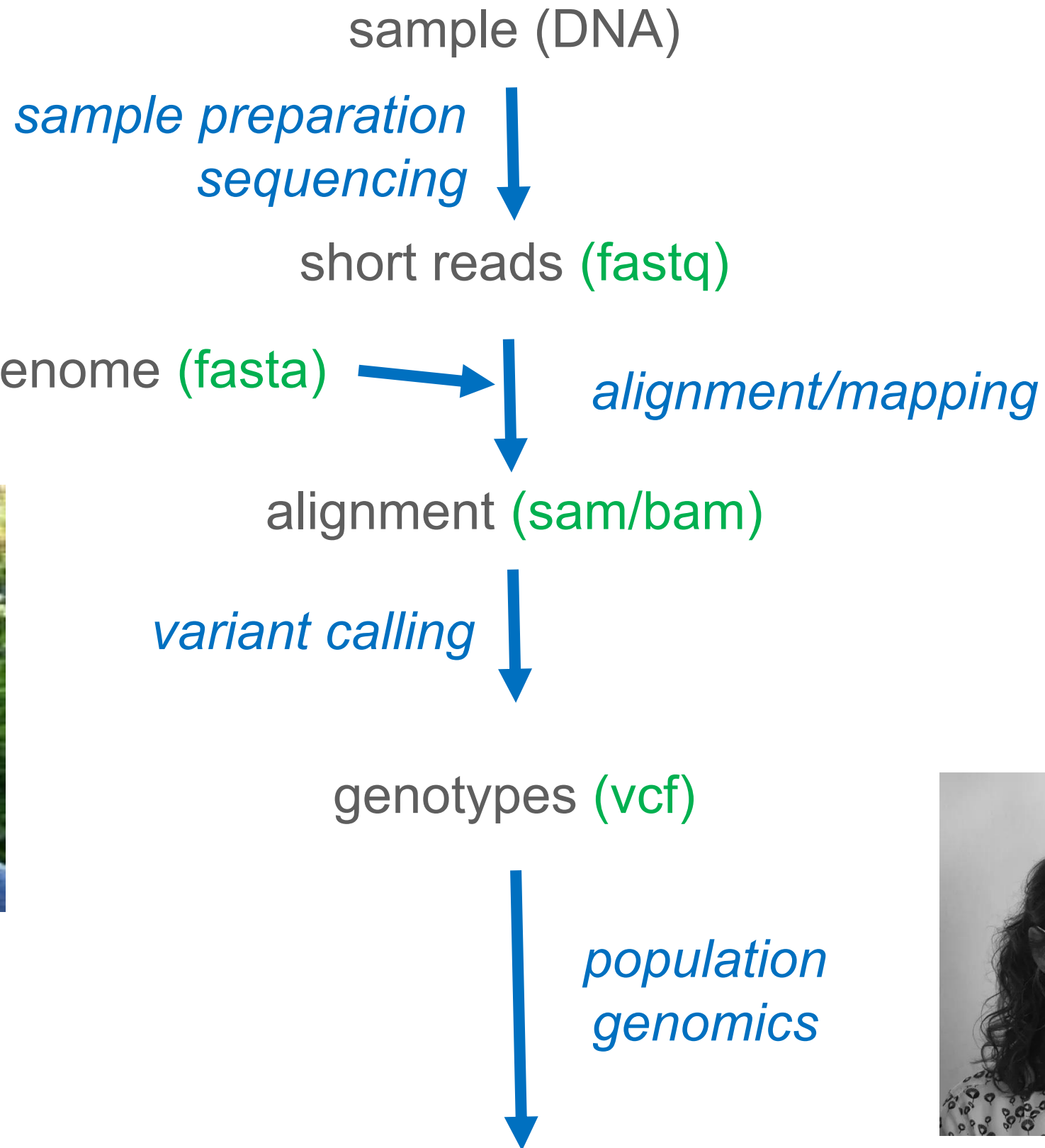
Benjamin Buchfink



Olga Vinnere
Pettersson



Petr Danecek



Benjamin Buchfink



Madeline Chase

sample (DNA)

*sample preparation
sequencing*



long reads (fastq)

reference genome (fasta)



alignment/mapping



alignment (sam/bam)

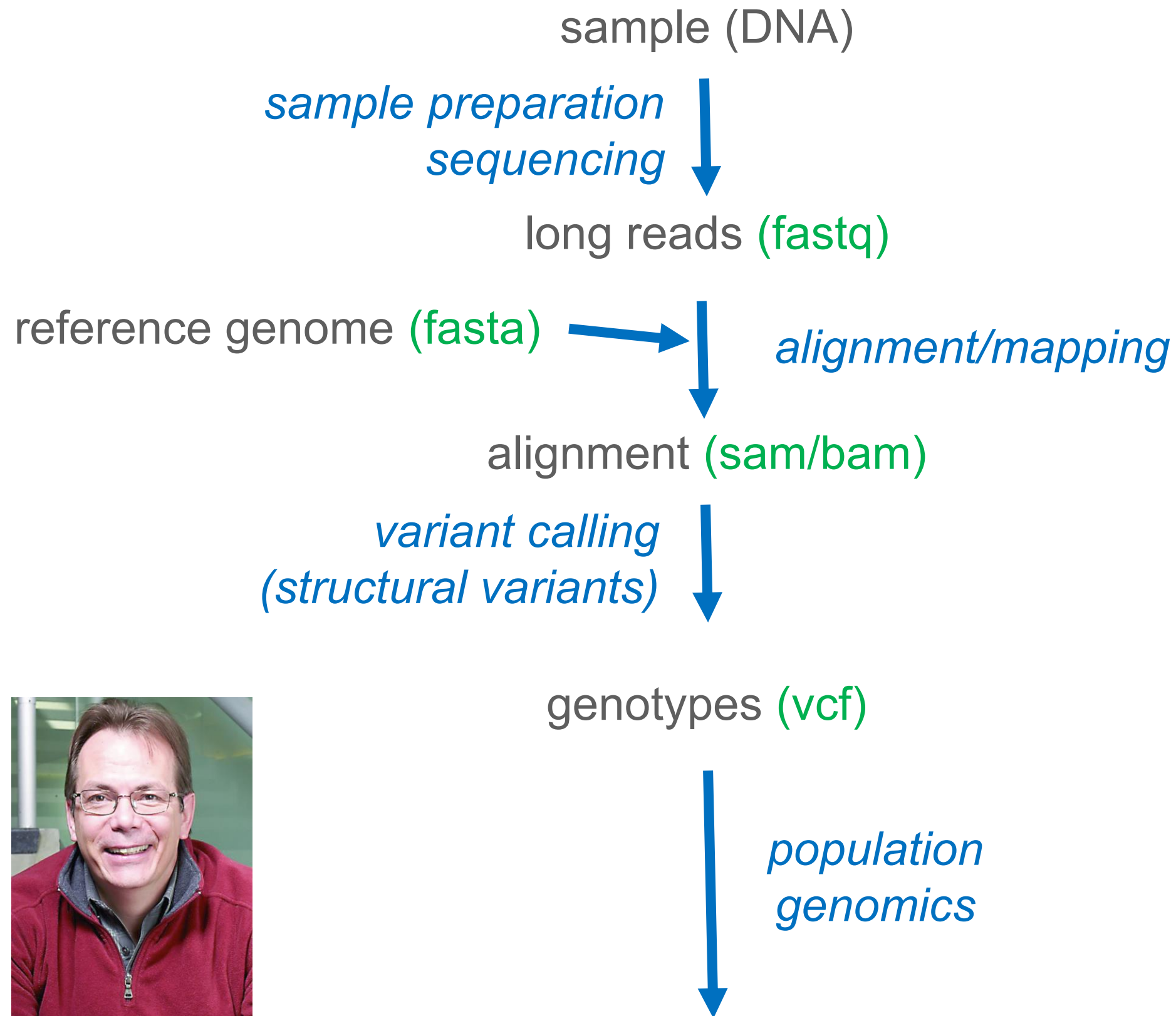
*variant calling
(structural variants)*



genotypes (vcf)

*population
genomics*

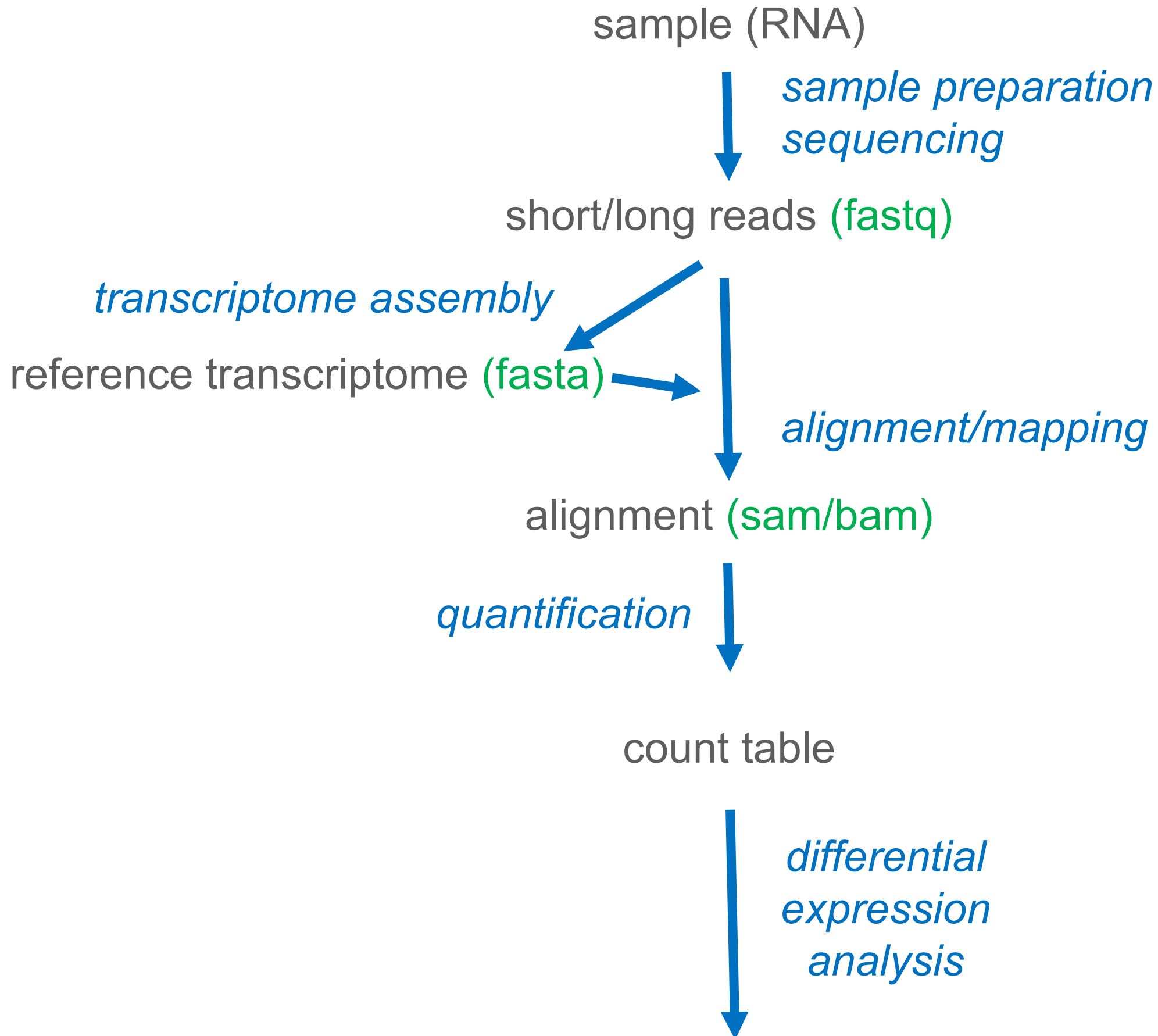


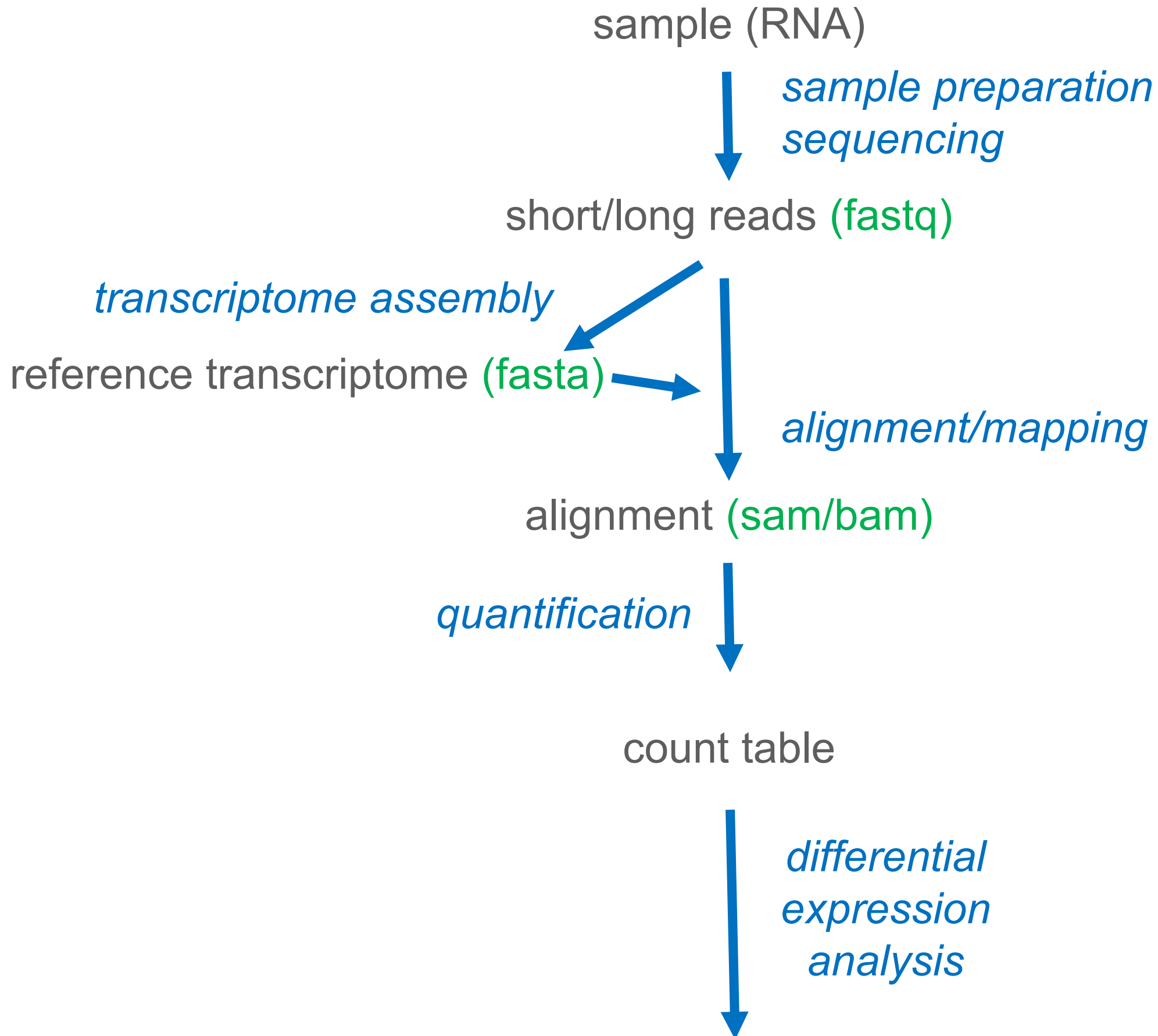


Evan Eichler

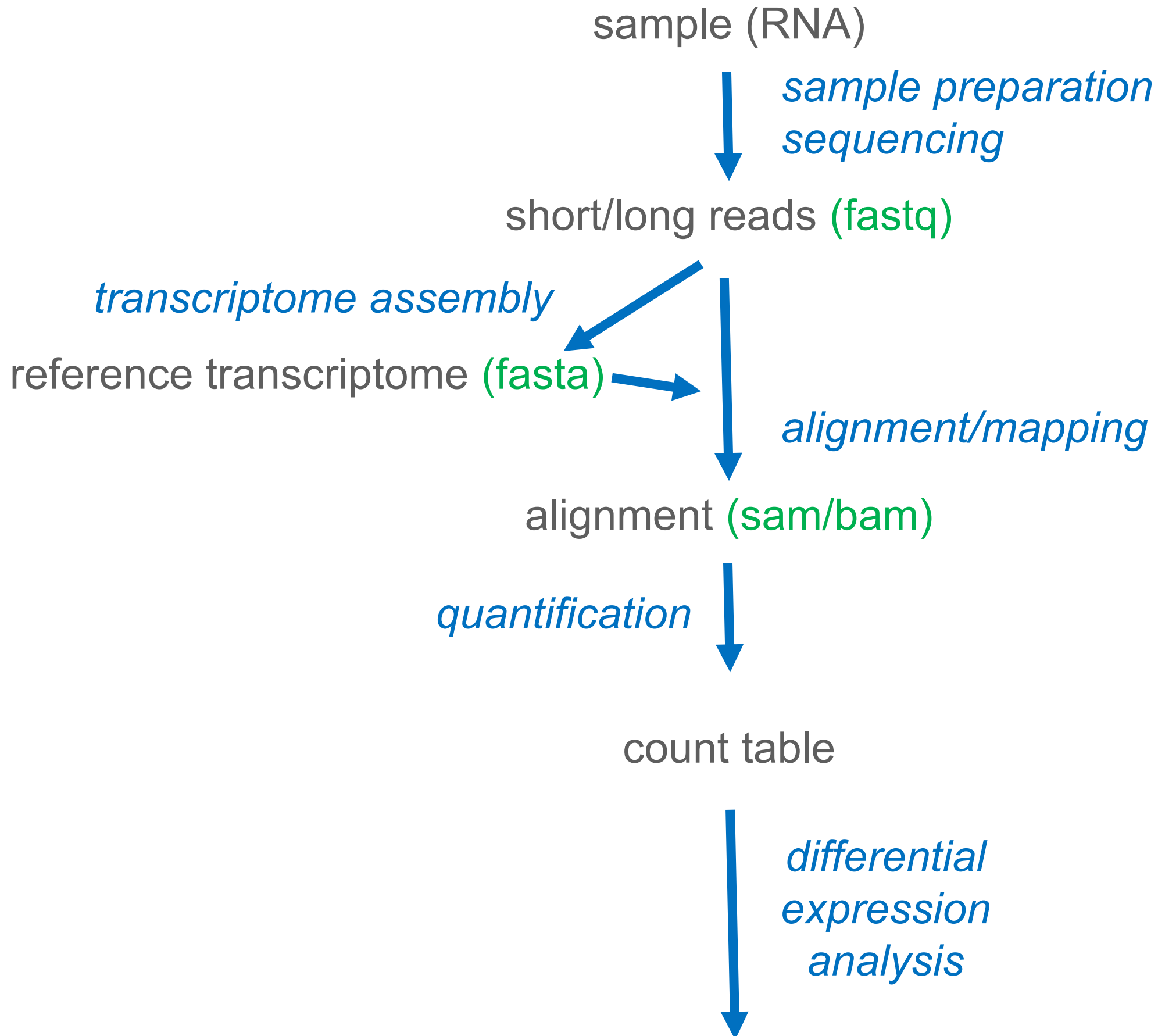


Claire Mérot





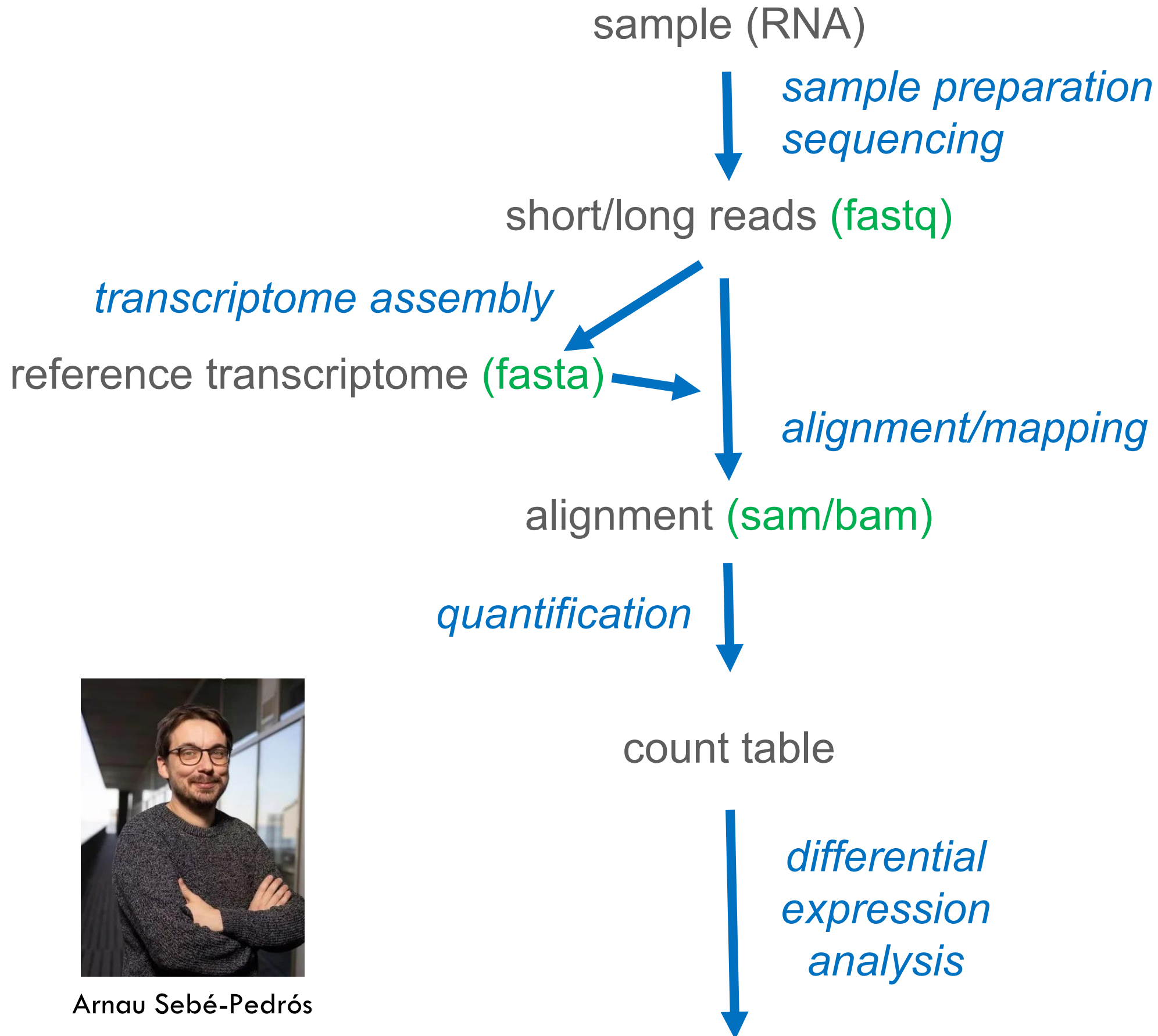
Brian Haas



Brian Haas



Rachel Steward



Brian Haas



Rachel Steward



Arnau Sebé-Pedrós



Maliheh Mehrshad



Guy Leonard



Mercè
Montoliu Nerin



Miloš Duchoslav



Petr Danecek



Dag Ahren



Scott Handley



Chris Wheat







Time for feedback!



The 2026 Workshop on Genomics takes place from the **11th-24th of January, 2026** in Český Krumlov, Czechia.

Important information:

- [Group photo!](#)

Feedback

- [Pre-workshop feedback](#)
- [Feedback week 1](#)
- [Feedback week 2](#)
- For sending feedback during the workshop, direct message [evomics.workshops](#) on slack, or email evomics.workshop@gmail.com

Find your way around Český Krumlov

- Book your town activities [here](#)
- [Map](#) of the **main locations in the schedule** (Save to your Google Maps!)
- [Map](#) of some of the main Český Krumlov **restaurants** (Save to your Google Maps!)

