

Day 3: Check up

MMB-114

Schedule

Day 1: Basics of UNIX and working with the command line

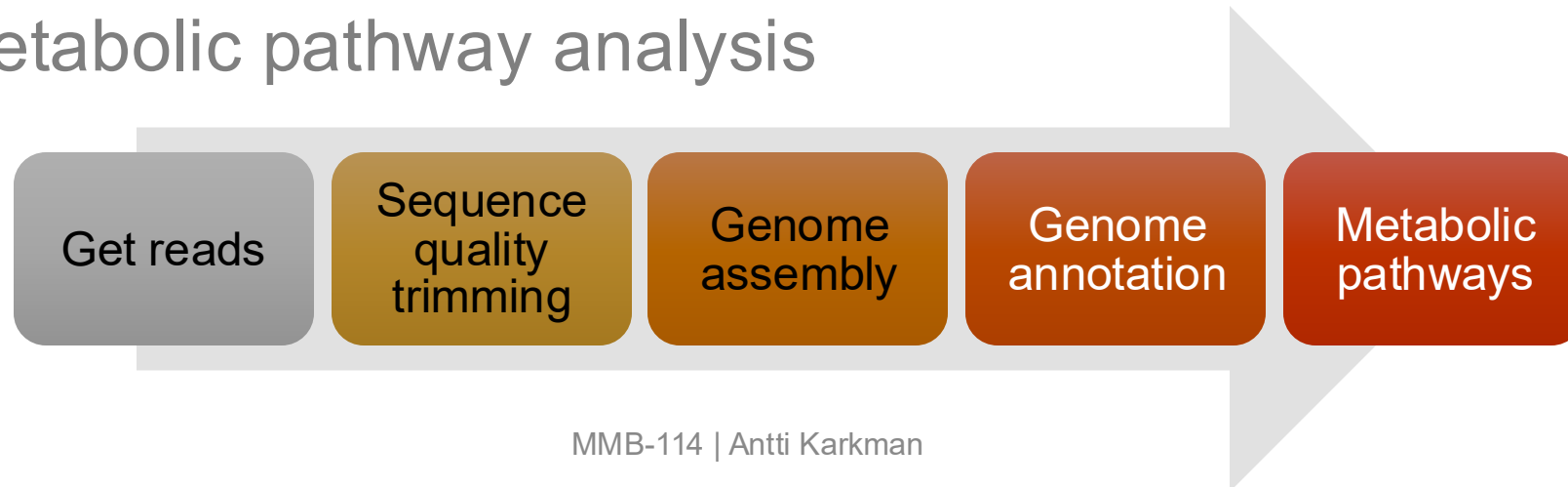
Day 2: Handling of Nanopore/Illumina data

Day 3: Check-up

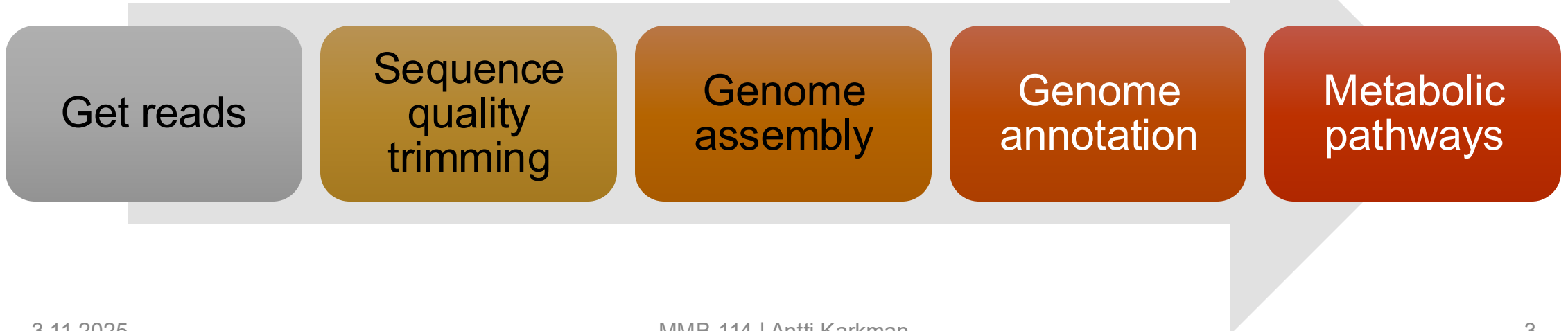
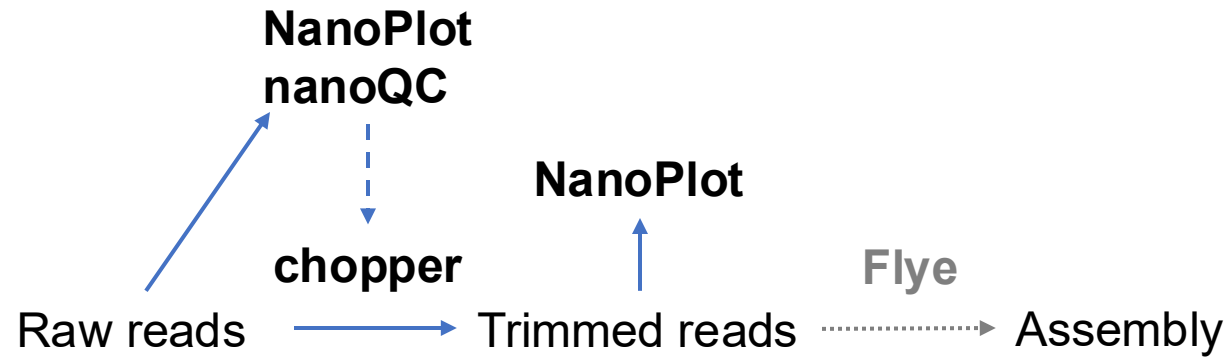
Day 4: Genome assembly

Day 5: Genome annotation

Day 6: Metabolic pathway analysis



Workflow so far



Questions about QC and trimming?

NanoPlot

- Which trimming parameters were set based on NanoPlot?

nanoQC

- Which trimming parameters were set based on nanoQC?

chopper

- What was done with chopper?

Illumina data

- How was the quality compared to Illumina?
- How much sequencing data there was?
- What kind of trimming was necessary?

Questions | Comments?