

## Detailed programme overview

WEEK 1: Morning (9-12)	Monday	Tuesday	Wednesday	Thursday	Friday
	<p><b>Course intro</b></p> <p><i>Lecture (9-9.30):</i> "Welcome and introduction by ECDC"</p> <p><i>Lecture (9.45-10.30):</i> "The role of genome sequencing in pathogen surveillance"</p> <p><i>Short talk (10.40-11):</i> "Overview of course and ILOs"</p> <p><i>Group talk (11-12):</i> "Getting to know your fellow course participants"</p>	<p><b>Getting started with Python</b></p> <p><i>Practical (9.30-10.15):</i> "Python introduction"</p> <p><i>Lecture (10.30-10.50):</i> "Python as a bioinformatical tool"</p> <p><i>Practical (10.50-12):</i> "Getting started with python"</p>	<p><b>Sequencing and raw-data QC</b></p> <p><i>Lecture (9.00 - 9.30):</i> "Introduction to sequencing"</p> <p><i>Lecture (9.40 – 10.45):</i> "Overview of sequencing technologies"</p> <p><i>Interactive Lecture/Practical (11.00 – 12.00):</i> "Sequencing Quality"</p>	<p><b>Genome assembly</b></p> <p><i>Lecture (9-9.45):</i> "Genome assembly strategies"</p> <p><i>Practical (10-11):</i> "Bacterial genome assembly with short reads"</p> <p><i>Practical (11-12):</i> "Basic assembly QC stats"</p>	<p><b>Genome assembly</b></p> <p><i>Lecture (9-9.30):</i> "Intro to long-read and hybrid genome assembly"</p> <p><i>Practical (9.30-11):</i> "Long-read and hybrid assembly"</p> <p><i>Practical (11-12):</i> "Genome assembly comparison"</p>

WEEK 1: Afternoon (13-16.30)	Monday	Tuesday	Wednesday	Thursday	Friday
	<p><b>Getting started with the command-line and Bash</b></p> <p><i>Lecture (13-13.20): “Command-line introduction”</i></p> <p><i>Practical (13.20-14.20): “Getting started with the Command-line”</i></p> <p><i>Lecture (14.30-14.45): “Bash Introduction”</i></p> <p><i>Practical (14.45-16.30): “Getting started with Bash”</i></p>	<p><b>Advancing in programming</b></p> <p><i>Lecture (13-13.20): “Programming recap and Biopython deep-dive”</i></p> <p><i>Coding session (13.20-16.20): Free</i></p> <p><i>Lecture (16.20-16.30): “Programming wrap-up”</i></p>	<p><b>Sequencing and raw-data QC</b></p> <p><i>Lecture/Practical (13.00-15.00): “Quality control of sequencing data”</i></p> <p><i>Lecture/Practical (15:00-16.30): “Contamination control, sample aggregation, and wrap-up”</i></p>	<p><b>Genome assembly</b></p> <p><i>Lecture/practical (13-14.30): “Contamination”</i></p> <p><i>Coding session (14.30-16.30): Biopython for assembly analysis</i></p>	<p><b>Genome analysis intro</b></p> <p><i>Lecture/Practical (13-15): “Blast, in-silico MLST and gene annotation”</i></p> <p><i>Coding session (15-16.30): Free</i></p>

WEEK 2: Morning	Monday	Tuesday	Wednesday	Thursday	Friday
	<p><b>Getting started with phylogenies</b></p> <p><i>Lecture (9-10):</i> “Evolution and phylogenies”</p> <p><i>Practical (10.20-12):</i> “Sequence handling and alignment”</p>	<p><b>Outbreaks, typing and AMR</b></p> <p><i>Lecture (9-9.25):</i> Intro to WGS-based FWD surveillance</p> <p><i>Practical (9.25-12):</i> Listeria outbreak detection</p>	<p><b>Pipeline development</b></p> <p><i>Lecture (9-9.45):</i> “Basic principles of automation”</p> <p><i>Practical (10.00-11.45):</i> “Pipeline development – scaling number of tools and samples”</p>	<p><b>Data sharing</b></p> <p><i>Lecture (9-9.45):</i> “Introduction to sequence databases and data sharing”</p> <p><i>Demo/practical (10.00-11.15):</i> “Uploading raw sequence data to ENA”</p> <p><i>Practical (11.15-12):</i> “Data exploration and retrieval using ENA resources”</p>	<p><b>Automation – going large-scale</b></p> <p><i>Lecture (9-9.45):</i> ”Automating the lab for large-scale testing and sequencing”</p> <p><i>Site-visit (9.45-10.15):</i> “Visit to the Danish Biobank”</p> <p><b>What ´s next?</b> <i>Lecture (10.45-11.30):</i> “Outlook – how to keep learning”</p> <p><b>Evaluation</b> Evaluation (11.30-12): “Course evaluation: formal”</p>

WEEK 2: Afternoon	Monday	Tuesday	Wednesday	Thursday	Friday
	<b>Getting started with phylogenies</b>  <i>Lecture (13-14):</i> “Phylogenetic methods”  <i>Practical (14.30-16.30):</i> “Inferring phylogenies”	<b>Outbreaks, typing and AMR</b>  <i>Practical (13-14.30):</i> Serotyping and virulence typing  <i>Practical (14.30-16):</i> AMR and point mutations detection in Salmonella  <i>Wrap-up/Q&amp;A (16-16.30)</i>	<b>Pipeline development</b>  <i>Practical (12.45-15.30):</i> “Pipeline development – from raw reads to data sharing”  <i>Free session (15.30-16.30):</i> “Setup at SSI and how automation could be introduced in participants’ workflows”	<b>Data sharing</b>  <i>Lecture (13-13.45):</i> “Intro to online tools for genome analysis, data sharing and visualization, developed by ECDC)  <i>Practical (14-16.30):</i> “Exploring Microreact, EpiPulse and Tessy”	<b>Evaluation</b>  <i>Evaluation (13-13.30):</i> “Course evaluation: (in)formal”  <b>What ´s next?</b>  <i>Free session (13.30-15.30):</i> “Q and A with your trainers”