

Examples of teaching approaches used in the Train-the-Trainer course on Capacity Building for Genomic Surveillance of AMR in low- and middle income countries

Strategy	Description	Implementation examples
<b>challenge and problem-based tasks</b>	problem solving and decision making exercises	<ul style="list-style-type: none"> <li>• Use card-based blueprint to order processes and procedures in a typical WGS for AMR surveillance pipeline;</li> <li>• Use of fictive case scenarios involving training others in genomics aspects.</li> </ul>
<b>case studies</b>	use of real cases, as published in the literature	<ul style="list-style-type: none"> <li>• Identify MDR <i>Salmonella enterica</i> subsp. <i>enterica</i> serotype Kentucky lineage expansion (Hawkey, Le Hello et al. 2019)</li> <li>• Investigation of a Brazilian epidemic of <i>Mycobacterium abscessus</i> species complex (Everall, Nogueira et al. 2017)</li> <li>• <i>Klebsiella pneumoniae</i> outbreak investigation (Wilson, Khokhar et al. 2018)</li> </ul>
<b>historical and chronological presentation of processes</b>	demonstration of lab procedures, visual tours, lab visits, exhibition tour.	<ul style="list-style-type: none"> <li>• Visit to the sequencing facilities at the Sanger Institute.</li> <li>• Demonstration of Library Preparation for WGS.</li> <li>• Demonstration of best practices for Antimicrobial Susceptibility testing</li> </ul>
<b>comparative analysis of tools and platforms</b>	presenting alternative ways of achieving the same/similar result	<ul style="list-style-type: none"> <li>• Comparison of data merging and consolidation techniques: interactively with spreadsheets versus programmatically with R and Python.</li> <li>• Comparison of different databases and software tools used to determine the AMR phenotype based on genomic data.</li> </ul>
<b>project planning and management techniques</b>	considering standards and quality control, constraints and limitations, time management; project planning and constraints [financial, scaling and other]	<ul style="list-style-type: none"> <li>• Exercise on how to set up a WGS laboratory</li> <li>• QC assessment of libraries and sequencing</li> </ul>
<b>provision of resources</b>	provision of links to resources and portals, quality control standards, virtual and cloud tools, toolkits	<ul style="list-style-type: none"> <li>• Software tools: <ul style="list-style-type: none"> <li>• AMR prediction tools: ARIBA (Hunt, Mather et al. 2017), Pathogen.watch, AMRfinder (Feldgarden, Brover et al. 2019)</li> <li>• AMR databases: CARD (Alcock, Raphenya et al. 2020), ResFinder (Bortolaia, Kaas et al. 2020)</li> <li>• Nucleotide archives: Genbank (Sayers, Cavanaugh et al. 2020), European Nucleotide Archive (Leinonen, Akhtar et al. 2011)</li> </ul> </li> </ul>
<b>analysis and evaluation</b>	Participants' specific contexts consideration and evaluation	<ul style="list-style-type: none"> <li>• Big picture exercise where participants evaluate their own circumstances.</li> </ul>