

This ENGAGE status newsletter is intended to give an outline of the status and an overview of the tasks performed and ongoing as part of the ENGAGE project.

Update regarding ongoing tasks

All ongoing tasks are listed in the Action Plan.

In addition to the milestones and deliverables described below, the Sharepoint for sharing documents between partner institutions is available (via Share DTU; <https://share.dtu.dk>). All partners have received an invitation email with further information on how to access and upload documents relevant to other ENGAGE partners (sharing of documents in a closed forum).

ENGAGE outside users and additional stakeholders have access to further information via the website (<http://www.engage-europe.eu/>).

Update regarding milestones

An overview of currently relevant milestones is listed in *Appendix 1*.

A brief extract of the milestones reached since December 2016:

M11.2, Task 11.3 – Interim meeting with EFSA – conducted in January 2017

M3.1, Task 3.1 – Ongoing WGS – conducted continuously

Update regarding deliverables

An overview of currently relevant deliverables listed in *Appendix 2*.

A brief extract of the deliverables completed or pending since December 2016:

Completed deliverables:

- **D6.1** (Task 6.1, Develop and execute the PT including documentation, reference material and analysis - following the GMI PT)
- **D7.5** (Task 7.3, Analysis / benchmarking report - "Data results" Report 3 (MOST vs SeqSero vs SISTR))
- **D8.1** (Task 8.4, Production and distribution of E-learning lectures and quizzes)
- **D11.1** (Task 11.4, Reporting on financial and technical progress)
- **D7.6** (Task 7.3, Analysis / benchmarking report - "Data results" Report 4 AMR genes)

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Appendix 1: Overview of milestones which are due from December 2016 until September 2017; sorted by date

Milestone	Task	Description of milestones (Milestones reached are marked grey)	Date (Gantt chart)	Status
M2.1	Task 2.5	Screening for relevant collected and available strains for WGS	2016-Dec	Complete
M3.1	Task 3.1	Ongoing WGS	2016-Dec	Complete
M5.2	Task 5.3	Interpretation of benchmarking analysis of the sequencing data	2016-Dec	Complete
M10.3	Task 10.3	Dissemination of output and outcome through GMI	2016-Dec	Complete
M11.2	Task 11.3	Kick off, interim and final meetings with EFSA	2016-Dec	Complete
M2.1	Task 2.5	Screening for relevant collected and available strains for WGS	2017-Feb	Complete
M3.1	Task 3.1	Ongoing WGS	2017-Feb	Complete
M5.2	Task 5.3	Interpretation of benchmarking analysis of the sequencing data	2017-Feb	Complete
M2.1	Task 2.5	Screening for relevant collected and available strains for WGS	2017-Mar	Complete
M3.1	Task 3.1	Ongoing WGS	2017-Mar	Complete
M5.2	Task 5.3	Interpretation of benchmarking analysis of the sequencing data	2017-Mar	Complete
M2.1	Task 2.5	Screening for relevant collected and available strains for WGS	2017-Apr	Complete
M3.1	Task 3.1	Ongoing WGS	2017-Apr	Complete
M5.2	Task 5.3	Interpretation of benchmarking analysis of the sequencing data	2017-Apr	Complete
M2.1	Task 2.5	Screening for relevant collected and available strains for WGS	2017-May	
M3.1	Task 3.1	Ongoing WGS	2017-May	
M5.2	Task 5.3	Interpretation of benchmarking analysis of the sequencing data	2017-May	
M2.1	Task 2.5	Screening for relevant collected and available strains for WGS	2017-Jun	
M3.1	Task 3.1	Ongoing WGS	2017-Jun	
M5.2	Task 5.3	Interpretation of benchmarking analysis of the sequencing data	2017-Jun	
M8.6	Task 8.6	Conduct twinning programs	2017-Jun	
M2.1	Task 2.5	Screening for relevant collected and available strains for WGS	2017-Jul	
M3.1	Task 3.1	Ongoing WGS	2017-Jul	
M5.2	Task 5.3	Interpretation of benchmarking analysis of the sequencing data	2017-Jul	
M2.1	Task 2.5	Screening for relevant collected and available strains for WGS	2017-Aug	
M3.1	Task 3.1	Ongoing WGS	2017-Aug	
M5.2	Task 5.3	Interpretation of benchmarking analysis of the sequencing data	2017-Aug	
M2.1	Task 2.5	Screening for relevant collected and available strains for WGS	2017-Sep	
M3.1	Task 3.1	Ongoing WGS	2017-Sep	

Appendix 2: Overview of deliverables which are due from January 2017, sorted by date

Deliverable	Task	Description of deliverables (COMPLETED deliverables are marked grey)	Date (Gantt chart)	Status
D6.1	Task 6.1	Develop and execute the PT including documentation, reference material and analysis - following the GMI PT	2017-Jan	Complete
D7.5	Task 7.3	Analysis / benchmarking report - "Data results" Report 3 (MOST vs SeqSero vs SISTR)	2017-Jan	Complete
D7.1	Task 7.1	Develop quality assurance guideline / bioinformatics quality marker tester tool. - "what is good sequence quality"	2017-Jan	Complete
D8.1	Task 8.4	Production and distribution of E-learning lectures and quizzes	2017-Jan	Complete
D10.4	Task 10.2	Dissemination through newsletters to partners, affiliated partners, authorities and research organizations	2017-Jan	Complete
D11.1	Task 11.4	Reporting on financial and technical progress	2017-Jan	Complete
D7.6	Task 7.3	Analysis / benchmarking report - "Data results" Reprot 4 AMR genes	2017-Mar	Complete
D10.5	Task 10.2	Dissemination through newsletters to partners, affiliated partners, authorities and research organizations	2017-Apr	Complete
D7.7	Task 7.3	Analysis / benchmarking report - "Data results"	2017-Jun	
D10.6	Task 10.2	Dissemination through newsletters to partners, affiliated partners, authorities and research organizations	2017-Jul	
D7.8	Task 7.3	Analysis / benchmarking report - "Data results"	2017-Sep	
D10.7	Task 10.2	Dissemination through newsletters to partners, affiliated partners, authorities and research organizations	2017-Oct	
D6.2	Task 6.1	Develop and execute the PT including documentation, reference material and analysis - following the GMI PT	2017-Nov	
D7.9	Task 7.3	Analysis / benchmarking report - "Data results"	2017-Dec	
D8.2	Task 8.4	Production and distribution of E-learning lectures and quizzes	2018-Jan	
D10.8	Task 10.4	Submission of generated genomes to ENA	2018-Jan	
D11.2	Task 11.4	Reporting on financial and technical progress	2018-Jan	