

This is the first ENGAGE status newsletter in the current form, and has been produced 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 which is regularly sent by email to all institute partners.

In addition to the milestones and deliverables described below, the [Sharepoint](#) for sharing documents between partner institutions will soon be available. All partners will soon receive an invitation email with further information on how to access and upload documents relevant to other ENGAGE partners. The Sharepoint document sharing site allows partner institutions to share documents in a closed forum.

ENGAGE outside users and additional stakeholders will have access to further information via the [website](#). The website set-up is currently being finalized and the link will be circulated to a.s.a.p.

## Update regarding milestones

An overview of milestones for 2016 is listed in *Appendix 1*.

In summary, the milestones reached or awaiting other tasks are as follows:

- [Milestone M11.1](#) (kick off, interim and final meetings with EFSA) is reached
- [Milestone M2.1](#) (screening for relevant collected and available strains for WGS) is pending as we have been awaiting the completion of Task 2.1
- [Milestone M3.1](#) (Ongoing WGS) is pending as we have been awaiting the completion of Task 2.1

Task 2.1 (establish selective criteria for relevance of included strains and genomes) has now been completed, as the document has been finalized and circulated.

Task 8.5 (identify and contact potential twinning participants among partners, affiliated partners, authorities and research organizations) has been ongoing, leading to a number of persons already having participated in twinning programmes.

Task 8.6 (conduct twinning programs) has been ongoing. At DTU, 5 persons have participated in twinning programmes. One more twinning programme will be conducted in June 2016 at DTU and also, one will run at PHE in June 2016.

## Update regarding deliverables

An overview of deliverables for 2016 is listed in *Appendix 2*.

In summary, the deliverables completed or awaiting other tasks are as follows:

- [Deliverable D1.1](#) (code-of-conduct finalized and circulated) has been completed

- Deliverable D4.1 (create download tools for consortium partners) has been completed
- Deliverable D5.1 (identify bioinformatic tools and software for the analysis and benchmarking) has been completed
- Deliverable D10.1 (dissemination through newsletters to partners, affiliated partners, authorities and research organizations) has been completed [updates were sent by email]
- Deliverable D7.3 (analysis/benchmarking report - "data results") is pending as we are awaiting the completion of Task 5.2

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**Appendix 1: Overview of milestones which are due until October 2016; sorted by date**

<b>Milestone</b>	<b>Task</b>	<b>Description of milestones (Milestones reached are marked grey)</b>	<b>Date (Gantt chart)</b>	<b>Status</b>
M11.1	Task 11.3	Kick off, interim and final meetings with EFSA	2016-Jan	Complete
M2.1	Task 2.5	Screening for relevant collected and available strains for WGS	2016-Feb	Awaiting Task 2.1
M2.1	Task 2.5	Screening for relevant collected and available strains for WGS	2016-Mar	Awaiting Task 2.1
M3.1	Task 3.1	Ongoing WGS	2016-Mar	Awaiting Task 2.1
M2.1	Task 2.5	Screening for relevant collected and available strains for WGS	2016-Apr	Awaiting Task 2.1
M3.1	Task 3.1	Ongoing WGS	2016-Apr	Awaiting Task 2.1
M2.1	Task 2.5	Screening for relevant collected and available strains for WGS	2016-May	
M3.1	Task 3.1	Ongoing WGS	2016-May	
M2.1	Task 2.5	Screening for relevant collected and available strains for WGS	2016-Jun	
M3.1	Task 3.1	Ongoing WGS	2016-Jun	
M5.1	Task 5.2	Pilot comparable benchmarking analysis	2016-Jun	
M5.2	Task 5.3	Interpretation of benchmarking analysis of the sequencing data	2016-Jun	
M2.1	Task 2.5	Screening for relevant collected and available strains for WGS	2016-Jul	
M3.1	Task 3.1	Ongoing WGS	2016-Jul	
M5.2	Task 5.3	Interpretation of benchmarking analysis of the sequencing data	2016-Jul	
M10.1	Task 10.3	Dissemination of output and outcome through GMI	2016-Jul	
M2.1	Task 2.5	Screening for relevant collected and available strains for WGS	2016-Aug	
M3.1	Task 3.1	Ongoing WGS	2016-Aug	
M5.2	Task 5.3	Interpretation of benchmarking analysis of the sequencing data	2016-Aug	
M8.1	Task 8.1	Develop one week hands on training course curricula	2016-Aug	
M2.1	Task 2.5	Screening for relevant collected and available strains for WGS	2016-Sep	
M3.1	Task 3.1	Ongoing WGS	2016-Sep	
M5.2	Task 5.3	Interpretation of benchmarking analysis of the sequencing data	2016-Sep	
M10.2	Task 10.3	Dissemination of output and outcome through GMI	2016-Sep	
M2.1	Task 2.5	Screening for relevant collected and available strains for WGS	2016-Oct	
M3.1	Task 3.1	Ongoing WGS	2016-Oct	
M5.2	Task 5.3	Interpretation of benchmarking analysis of the sequencing data	2016-Oct	
M8.3	Task 8.3	Conduct hands on training courses	2016-Oct	
M9.1	Task 9.1	Workshops	2016-Oct	

**Appendix 2: Overview of deliverables which are due in 2016, sorted by date**

<b>Deliverable</b>	<b>Task</b>	<b>Description of deliverables (COMPLETED deliverables are marked grey)</b>	<b>Date (Gantt chart)</b>	<b>Status</b>
D1.1	Task 1.5	Create and sign NDA, MoU and MTA among partners, affiliated partners, authorities and research organizations	2016-Feb	Completed
D5.1	Task 5.1	Identify bioinformatic tools and software for the analysis and benchmarking	2016-Feb	Completed
D4.1	Task 4.4	Create download tools for consortium partners	2016-Mar	Completed
D10.1	Task 10.2	Dissemination through newsletters to partners, affiliated partners, authorities and research organizations	2016-Mar	Completed
D7.3	Task 7.3	Analysis / benchmarking report - "Data results"	2016-May	Awaiting Task 5.2
D10.2	Task 10.2	Dissemination through newsletters to partners, affiliated partners, authorities and research organizations	2016-Jun	
D7.10	Task 7.4	Protocols / SOPs for DNA prep, library prep, and sequencing	2016-Jul	
D7.11	Task 7.5	Guideline to bioinformatic tools - "Availability, what to use - and when"	2016-Jul	
D7.4	Task 7.3	Analysis / benchmarking report - "Data results"	2016-Aug	
D7.2	Task 7.2	best practise guidelines - "How to get started, what to do and use and what not to do"	2016-Sep	
D10.3	Task 10.2	Dissemination through newsletters to partners, affiliated partners, authorities and research organizations	2016-Sep	
D6.1	Task 6.1	Develop and execute the PT including documentation, reference material and analysis - following the GMI PT	2016-Oct	
D7.5	Task 7.3	Analysis / benchmarking report - "Data results"	2016-Nov	
D7.1	Task 7.1	Develop quality assurance guideline / bioinformatics quality marker tester tool. - "what is good sequence quality"	2016-Dec	
D8.1	Task 8.4	Production and distribution of E-learning lectures and quizzes	2016-Dec	
D10.4	Task 10.2	Dissemination through newsletters to partners, affiliated partners, authorities and research organizations	2016-Dec	
D11.1	Task 11.4	Reporting on financial and technical progress	2016-Dec	