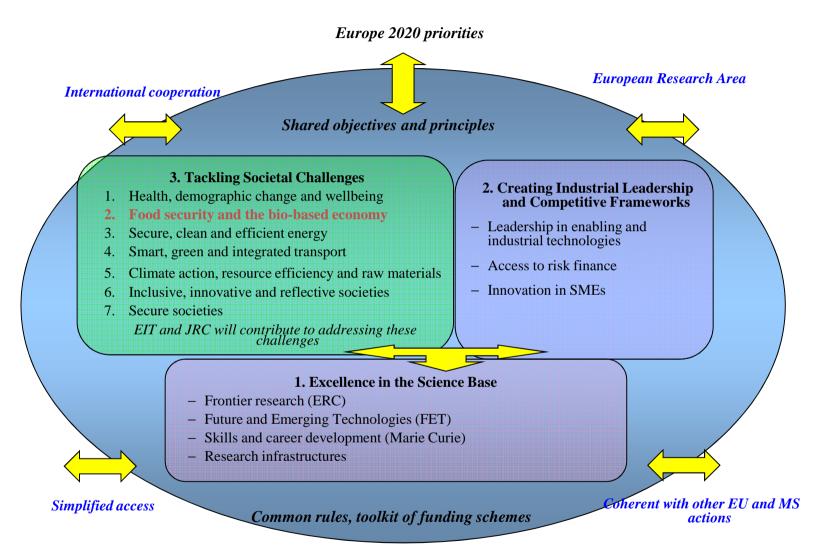




Involvement of EURL Lm in the European project COMPARE

Sophie ROUSSEL, SEL Unit

Horizon 2020 - Objectives and structure



COMPARE project :

-COllaborative Management Platform for detection and Analyses of (Re-)emerging and foodborne outbreaks in Europe

-Coordinator: DTU-Food, (Franck Aarestrup's team) 01/12/2014-30/11/2019 (60 months)

- -29 European partners:
- -Anses: ; Unit SEL (EURL for *Listeria monocytogenes , Salmonella, E. coli*); Anses WGS platform (Ploufragan, Brittany)
- -15 Work packages
- Face-to face meeting: 11-12 March 2015, Copenhague

Similar aims to those targeted by GMI (Global Microbial identifier) www.globalmicrobialidentifier.org

COMPARE - aims :

- □ To promote development and deployment of novel applications, data sharing and analysis systems across the diversity of pathogens, domains and sectors.
- □ To improve rapid identification, containment and mitigation of emerging infectious diseases and foodborne outbreaks,
- To develop a cross-sector and cross-pathogen analytical framework and globally linked data and information sharing platform,
- To integrate state-of-the-art strategies, tools, technologies and methods for collecting, processing and analyzing sequence-based pathogen data in combination with associated (clinical, epidemiological and other) data,
- □ To generate actionable information to relevant authorities in the human health, animal health and food safety domains;

Items covered by COMPARE - Partners

Human Health; Animal health; Food safety; Viruses; Bacteria; Protozoa

- 1.DTU, DK: Veterinary /food bacteriology, Bioinformatics Meta-genomics Foodborne pathogens, Risk Assessment, Risk modeling, Foodborne virus
- 2.EMC , NL: Public health epidemiology, Molecular virology, Comparative pathology, bioinformatics
- 3.SSI, DK: Public health epidemiology, Foodborne pathogens
- 4.FLI, DE: Veterinary Virology/Zoonoses, Epidemiology, Databases, Data analysis and curation, Next generation sequencing
- 5. ANSES, FR: Molecular bacteriology, Foodborne pathogens, Epidemiology
- 6.RKI, DE: Molecular diagnostics, epidemiology, biochemistry
- 7.ISS , IT : Molecular biology and epidemiology, Parasitology
- 8.RIVM, NL: Veterinary virology, epidemiology
- 9.AHVLA, UK: Veterinary bacteriology and virology, bioinformatics, epidemiology, risk assessment modeling

Partners

- 11. UEDIN, UK: Epidemiology, pathogen evolution
- 12. UK- Bonn (DE) Virology, molecular diagnostics, virus evolution
- 13. AMC, NL: Microbiology, bacteriology, clinical diagnostics
- 14. UA, BE: Microbiology, bacteriology, clinical diagnostics
- 15. Artemis, NL: Virology, wildlife health, virus discovery
- 16. UCAM, UK: Mathematical modeling, pathogen evolution
- 17. TIHO , DE : Veterinary pathology
- 18. UCLM, ES , : Wildlife health
- 19. FMER, FR: Microbiology, pathogen discovery
- 20. AUTH, GR Microbiology, molecular epidemiology, veterinary virology
- 21. IFREMER, FR: Molecular bacteriology
- 22. EUR , NL : Epidemiology
- 23. ANU , AU : Epidemiology
- 27. UNIBO, IT: Veterinary & food bacteriology
- 28. DSMZ, DE: Bacterial population genomics
- 29. Sanger, UK: Genomics, molecular microbiology

WP2 – Harmonised standards for sample processing and sequencing

To develop harmonized analytical workflows for generation of high quality NGS data in combination with relevant metadata for pathogen detection and typing across sample types, pathogens and domains

- -sample handling, sample processing, sequencing protocols, sequence curation, storage...
- Historic and prospective biobanks as reference
- Develop a scheme for ring trials and external quality assurance systems