

D6.15: Report on outputs of the short-term missions Work Package 6

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REPORT ON SHORT TERM MISSIONS OUTPUTS

Overview of the One Health EJP Short Term Missions

The One Health EJP (OHEJP) Short Term Missions (STMs) were small co-funded travel grants which facilitated the exchange of scientific expertise, methodologies, equipment, and facilities to our consortium members. The aim of these missions was to harmonise the existing approaches and methodologies within the OHEJP.

These STMs drove research forward in a collaborative and non-duplicative fashion to strengthen the scientific capacity within the OHEJP, and to also contribute to the future prevention, preparedness, detection, and response of the EU to foodborne and other emerging threats across human-animal-environmental sectors.

The OHEJP co-funded a total of 19 STMs (each funded 44% by the EU), which involved 22 researchers and a total of 27 institutes within the OHEJP consortium. The Education and Training activities team from Work Package 6 (WP6) was responsible for co-ordinating the calls, selection of applicants for funding and reporting on the short term missions funded each year.

Table 1: All OHEJP short-term missions conducted between 2019 and 2023, with their titles, home and hosting institutes.

STM title	Home institute	Hosting institute
Training on the application of source attribution	IZLER, Italy	<u>RIVM</u> ,
models.		The Netherlands
Application of advanced epidemiological	VISAVET-UCM,	Hasselt University,
analytical methods for antimicrobial resistance	Spain	Belgium
data in <i>Salmonella</i> in pigs.		
Training in bioinformatics to study the dynamics	VISAVET-UCM,	<u>DTU</u> , Denmark
of <i>E. coli</i> in laying hens.	Spain	
Skills development focused on development of a	FHI, Norway	<u>SSI</u> , Denmark
framework for reporting outbreak investigations		
using consumer purchase data.		
Study of the interactions between STEC and	<u>ISS</u> , Italy	<u>UCA</u> , France
human gut microbiota in the ARCOL model.		
Cross-domain and cross-country collaboration	<u>SVA</u> , Sweden	<u>NIPH</u> , Norway
to develop multivariate syndromic surveillance.		
Startup of an efficient sequencing facility.	<u>NVI</u> , Norway	<u>SSI</u> , Denmark
CarbaPlasmid – Tracking endemic	<u>NUI Galway</u> ,	VISAVET-UCM,
carbapenemase plasmids in human, animal and	Ireland	Spain
environmental isolates.		
Zoonotic pathogen detection in rats.	<u>RIVM</u> ,	<u>FLI</u> , Germany
	The Netherlands	
Surveillance and source-attribution of AMR	<u>INIAV</u> , Portugal	DTU Food,
based on metagenomic analysis.		Denmark





		1
Validation and exchange of modelling tools to	<u>UoS</u> , UK	<u>RIVM</u> ,
assess the risk of human salmonellosis based on		The Netherlands
environmental factors using multiple sources of		
data.		
Application of spatial models to identify new	<u>INIA</u> , Spain	<u>RIVM</u> ,
environmental surveillance indicators of		The Netherlands
Salmonella and Campylobacter in pig and		
poultry.		
Construction of double-labelled <i>E. coli</i> strains to	<u>WUR</u> ,	University of
study the effect of antibiotics and interventions	The Netherlands	<u>Copenhagen</u> ,
on horizontal ESBL genes transfer in the		Denmark
chicken's caecal microbiome.		
Microbiological and molecular techniques for	<u>NDVRI</u> , Bulgaria	ANSES, France
Brucella identification from strains.		
Tolerance of biofilm forming bacteria to	<u>APHA</u> , UK	<u>NVI</u> , Norway
disinfectants after repeated disinfectant		
exposure.		
Metagenomic Tools for the Study of SARS-CoV-	<u>INIA</u> , Spain	<u>INSA</u> , Portugal
2 in Animals.		
Genotypic characterisation of antimicrobial	ANSES, France	<u>BfR</u> , Germany
susceptibility and isolation of Aeromonas and		
Vibrio phages from water samples.		
Stabile Intra-specific genomic bacterial	<u>NVI</u> , Norway	Pasteur Institute,
classification systems.		France
Application of Single-Cell Genomics for the	<u>UCM</u> , Spain	KIT, Germany
study of the bacterial reservoirs of plazomicin		
resistance determinants.		

Short Term Missions calls and selection

During the five years of the OHEJP, five STM calls were launched: one for completion of the STMs in 2019, one for 2020, one for 2021, and two for 2022. For all calls, a promotional email marketing strategy was used to disseminate the call information to the Scientific Steering Board (SSB), Project Management Team (PMT), Progamme Managers Committee (PMC), Project Leaders, PhD teams, and Communication Contact Persons. The branded and visually attractive emails contained instructions on how to access the guidelines, application form and templates. The calls were also disseminated in the Education and Training activities monthly bulletins, the OHEJP consortium newsletters, and social media channels, Twitter and Linkedin.

The WP6 Team co-ordinated the calls and selection of the STMs following the validated procedure. STMs application forms, work plans, financial budget templates as well as letters of recommendation from the home and hosting institute and the applicants' *Curriculum Vitae* were sent to three independent reviewers. The reviewers were nominated by the SSB.

The WP6 Team compiled the scores and validated the selection with the Project Management Team (PMT), and the final decision was then communicated to the SSB and applicants. A total of 25 applications were received: five for 2019, five for 2020, four for 2021, and eleven for 2022





(six from the first 2022 call and five from the additional call) and 23 were approved for funding: four for 2019, five for 2020, four for 2021, and ten for 2022 (six from the first 2022 call and four from the additional 2022 call).

Impact of COVID-19

The COVID-19 pandemic affected the success of STMs scheduled in 2020 and 2021 due to the disruption to travel plans and local restrictions. This resulted in the cancellation of four STMs: three in 2020 and one in 2021. One of the four STMs funded for 2021 was post-poned for travelling reasons and took place early in 2022. Therefore, due to the COVID-19 pandemic, only 19 of the 23 awarded STMs took place: four in 2019, two in 2020, two in 2021, and eleven in 2022.

Short Term Missions – Call 2019

Four STMs were funded for 2019, they all took place as scheduled.

1. Short Term Mission 1

STM Researcher: Virginia Filipello
Title of the STM: Training on the application of source attribution models
Home Institute: IZLER, Italy
Mission Hosting Institute: RIVM, the Netherlands
Duration of mission: 1 week

Aim of the mission: The aim of this mission was to enhance source attribution skills through an ad hoc structured and supervised training, which included exercises with real data and validation of learnt competences.







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Training on the application of source attribution models



...I am thoroughly grateful to the OHEJP for having granted such a stimulating and professionalizing experience, and I am confident that the newly learnt competence will advance the positive impact of our ongoing surveillance activities." Virginia Filipello, IZLER

Theme:

Home Institute: Mission Hosting Institute: Duration of mission: Integration of microbiological; Risk assessment and surveillance activities IZLER, Italy RIVM, the Netherlands 1 week

The aim of this mission was to enhance source attribution skills through an ad hoc structured and supervised training, which included exercises with real data and validation of learnt competences.

The need for this mission fell within the context of an ongoing process of updating the current routine surveillance of foodborne pathogens with sequence based techniques at IZSLER.

IZSLER carries out surveillance plans for the most populous region of Italy, Lombardy, producing large amounts of data. Therefore, the integration of surveillance data with source attribution analyses, could be of great usefulness to improve the positive outcomes on the regional territory.

One Health EIP has received funding from the European Union's Horizon 2020 research and Innovation programme under grant agreement No 773830.



Figure 1: This case study can be found on the OHEJP website: https://onehealthejp.eu/community/education-andtraining/short-term-missions-2019.





STM Researchers: Kendy Tzu-Yun Teng

Title of the STM: Application of advanced epidemiological analytical methods for antimicrobial resistance data of *Salmonella* in pigs.

Home Institute: VISAVET-UCM, Spain.

Mission Hosting Institute: Centre for Statistics, Hasselt University, Belgium.

Duration of mission: 7 weeks.

Aim of the mission: The aim of this mission was to develop skills in analysing data on phenotypic antimicrobial resistance (AMR), to identify patterns among the occurrences of resistance of different antimicrobials and to better understand the interactions and dynamics of *Salmonella* in pigs.







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Application of advanced epidemiological analytical methods for antimicrobial resistance data of Salmonella in pigs



...This short-term mission allowed us to extend connection to institutes external to OHEJP and to form new collaboration. Besides the benefits at the project and institute level, I as a researcher also benefited greatly... Thanks to OHJEP for giving me this opportunity!" Kendy Tzu-Yun Teng, **VISAVET-UCM**

Theme: Home Institute: Mission Hosting Institute: Duration of mission: Skills Development Missions <u>VISAVET-UCM</u>, Spain Centre for Statistics, Hasselt University 7 weeks

The aim of this mission was to develop skills in analysing data on phenotypic antimicrobial resistance (AMR) to identify patterns among the occurrences of resistance of different antimicrobials to better understand the interactions and dynamics in Salmonella in pigs.

During this STM, analysis of AMR data for Salmonella isolates collected between 2001-2013 through the Spanish Veterinary Antimicrobial Resistance Surveillance Network programme was performed. The antimicrobials included Cefotaxime, Chloramphenicol, Ciprofloxacin, Florfenicol, Gentamicin, Nalidixic acid and Tetracycline.

Multiple techniques, including principal component analysis, multiple correspondence analysis, hierarchical clustering, and latent class analysis, were performed to detect potential patterns and clusters among the AMR data. Generalised estimating equations were conducted to examine the evolution of the proportion of the resistant strains of each of the seven antimicrobials. Additionally, the structures of relationships among the antimicrobials were examined by Bayesian network analysis. Efforts to develop new approaches to Bayesian network analysis in Stan also took place.

The work performed during this STM demonstrated useful analytical techniques to explore the interactions among AMR and associations between AMR and Salmonella serotypes. On the base of this action, development to explore AMR phenotypes continues to be carried on.

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Figure 2: This case study can be found on the OHEJP website: https://onehealthejp.eu/community/education-andtraining/short-term-missions-2019.





STM Researcher: Irene Aldea Ramos
Title of the STM: Training in bioinformatics to study the dynamics of *E. coli* in laying hens: Training on the application of source attribution models.
Home Institute: VISAVET-UCM, Spain
Mission Hosting Institute: DTU, Denmark
Duration of mission: 4 weeks.

Aim of the mission: The aim of this mission was to improve the participants' existing bioinformatics knowledge. The National Food Institute of DTU is part of the Centre of Genomic Epidemiology which has several web services for the analysis of whole genome sequences.







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Training in bioinformatics to study the dynamics of E. coli in laying hens: Training on the application of source attribution models



... I not only learned a lot, but I also met colleagues from other parts of the world who work on similar tasks. We were able to share ideas. It was a motivating experience both academically and personally to know other places and cultures, and different way of working. I want to thank OHEJP and members of the DTU for making me feel comfortable in an unknown place ... " Irene Aldea Ramos, VISAVET-UCM

Name: Home Institute: Mission Hosting Institute: Duration of mission: Skills development missions <u>VISAVET-UCM</u>, Spain <u>DTU</u>, Denmark 4 weeks

The aim of this mission was to improve the participants' existing bioinformatics knowledge. The National Food Institute of DTU is part of the Centre of Genomic Epidemiology which has several web services for the analysis of whole genome sequences.

During this month, the participant learned to perform phylogenetic analysis using the complete DNA sequence and worked in the construction of phylogenetic trees with both programs of the Centre of Genomic Epidemiology and with other web-available programs.

They also analysed resistance genes and how they move between bacteria, focusing on mobile genetic elements. The participant also had the opportunity to further analyse their own research data and participated in collaborative discussions where they were given scientific advice and ideas for the work that they can do next.

During their stay, the participant also attended some sessions of a course where they were taught to use some tools for the analysis of WGS data.

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Figure 3: This case study can be found on the OHEJP website: https://onehealthejp.eu/community/education-andtraining/short-term-missions-2019.





STM Researcher: Solveig Jore
Title of the STM: Skills development focused on development of a framework for reporting outbreak investigations using consumer purchase data.
Home Institute: FHI, Norway.
Mission Hosting Institute: SSI, Denmark.
Duration of mission: 1 week.

Aim of the mission: The aim of this mission was to work towards developing a framework for reporting outbreak investigations using consumer purchase data. During this mission, the participant also reviewed the current and previous use of purchase data and the potential of its future use.

Dissemination and communication activities: The researcher published results obtained using the framework developed during the STM, in two research articles:

- Jore, S., Braae, U. C., Møller, F. T., Friesema, I., Paranthaman, K., Jalava, K., Jourdan-DaSilva, N., Löf, E., Rehn M. & Ethelberg S. (2022). A common framework for using and reporting consumer purchase data (CPD) in foodborne outbreak investigations in Europe, Infection Ecology & Epidemiology, 12:1.
 DOI: https://doi.org/10.1080/20008686.2021.2007828
- Dougherty, P.E., Møller, F.T., Ethelberg, S., Isaksson Rø G. Ø. & Jore S. (2022). Simulation and identification of foodborne outbreaks in a large supermarket consumer purchase dataset. Sci Rep 12, 11491.
 DOI: https://doi.org/10.1038/s41598-022-15584-x







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- Contributing to future prevention, preparedness, detection and response of the EU to foodborne and other emerging threats across human-animal-environmental sectors.

Skills development focused on development of a framework for reporting outbreak investigations using consumer purchase data



...SSI has previous experience with using consumer purchase data in several outbreak investigations... it was very useful to become a part of their team for a week and work together with the highly skilled epidemiologists on developing the use of this tool further ..." Solveig Jore, FHI

Name:

Home Institute: Mission Hosting Institute: Duration of mission: Skills development missions; Integration of microbiological; Risk assessment and surveillance activities; Harmonisation of diagnostics tests, platforms and research tools EHI, Norway SSI, Denmark 1 week

The aim of this mission was to work towards developing a framework for reporting outbreak investigations using consumer purchase data.

During this mission, the participant also reviewed the current and previous use of purchase data and the potential of its future use. The participant worked with collaborative colleagues at SSI on a description of "best practice" for using purchase data with the aim of harmonising the use of this kind of surveillance data amongst the European countries.

In addition, the collaborative team started drafting an opinion paper describing the method and existing barriers for the benefit of new users and with the hope that these barriers eventually can be overcome.

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Figure 4: This case study can be found on the OHEJP website: https://onehealthejp.eu/community/education-and-training/short-term-missions-2019.





Short Term Missions - Call 2020

Five STMs were awarded for 2020 but the onset of the COVID-19 pandemic affected the success of these missions considerably due to the disruption to travel plans. Two of the five STMs took place before travel was disrupted in March 2020. Unfortunately, the remaining three STMs were cancelled.

1. Short Term Mission 1

STM Researcher: Paola Chiani
Title of the STM: Study of the interactions between STEC and human gut microbiota in the ARtificial COLon (ARCOL) model.
Theme: Skills Development Missions
Home Institute: Istituto Superiore di Sanità (ISS), Italy

Mission Hosting Institute: Université d'Auvergne Clermont, France

Duration of mission: 5 weeks

Aim of the mission: The aim of this study was to investigate the interactions between Shiga toxin-producing *E. coli* (STEC) and human intestinal microbiota, using the ARtificial COLon model (ARCOL), which simulates the human large intestine functionality. In details, the aim of the STM was to perform *in vitro* infection experiments with STEC in presence of a normal human intestinal microbiota using this innovative model to understand the role played by the human intestinal microflora upon STEC infection, and to investigate changes occurring in the gut microbiota composition in presence of infecting STEC strains.







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Study of the interactions between STEC and human gut microbiota in the ARtificial COLon (ARCOL) model



... The experience was very formative for my career. It involved the use of the innovative ARCOL system, which is not present in Italy, and only present in a few institutes world wide. I'm grateful to have visited this excellence centre and to acquire a skill in the use of such model and of a holistic approach..." Paola Chiani, ISS

Theme: Home Institute: Mission Hosting Institute: Duration of mission: Skills Development Missions Istituto Superiore di Sanità (ISS), Italy Université Clermont Auvergne, France 5 weeks

Shiga toxin-producing E. coli (STEC) are zoonotic pathogens, causing severe afflictions in humans. Upon STEC infection, the host can present a wide range of symptoms including uncomplicated diarrhoea, haemorrhagic colitis and the life-threatening haemolytic uremic syndrome.

The aim of this mission was to investigate the interactions between STEC and human intestinal microbiota, during experimental infection using the ARtificial COLon (ARCOL) model, which simulates the human large intestine functionality. During the STM all the parameters to simulate the proper conditions for the ARCOL model of the intestine of subjects belonging to the considered age groups were determined and all the samples were collected for the following analyses.

The realisation of this project was possible thanks to the collaboration with colleagues from the Université Clermont Auvergne, in Clermont-Ferrand who developed the model, and hosted the mission.

One Health EJP has received funding from the European Union's Horizon 2020 research and innovation programme under grant agreement No 773830.



Figure 5: This case study can be found on the OHEJP website: <u>https://onehealthejp.eu/community/education-and-</u> <u>training/short-term-missions-2020</u>





STM Researcher: Wiktor Gustafsson
Title of the STM: Cross-domain and cross-country collaboration to develop multivariate syndromic surveillance
Theme: Skills Development Missions
Home Institute: Swedish National Veterinary Institute (SVA), Sweden Mission
Hosting Institute: Norwegian Institute of Public Health (NIPH), Norway
Duration of mission: 2 weeks

Aim of the mission: The aim of the STM was collaboration and knowledge exchange between Norway and Sweden in the development of One Health systems for multivariate syndromic surveillance of veterinary and public health data combined. Two separate systems are in development in Sweden at the SVA and Norway at the NIPH. The Swedish system is explanatory, aiming to combine several data sources to explain outbreaks and improve the accuracy of detection. Norway, on the other hand, is developing a predictive system, which aims to use some source(s) of data to predict the outcome of others.

Dissemination and communication activities: The results obtained during the STM contributed to the OHEJP NOVA project's work on the development of a multivariate syndromic surveillance engine, and a dashboard showcasing the surveillance results from the *Campylobacter* test case. This has been presented as a poster at the OHEJP Annual Scientific Meeting 2020.







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Cross-Domain and Cross-Country Collaboration to Develop Multivariate Syndromic Surveillance



...I am very grateful to the OHEJP for this opportunity. It gave me the chance to form new connections, exchange thoughts and ideas with new people. **Colleagues were very** welcoming: I look forward to working with them in the future. It was a rewarding experience to visit a public health institute at the doorstep of a global pandemic..."

Name: Home Institute: Mission Hosting Institute: Duration of mission: Skills Development Missions National Veterinary Institute (SVA), Sweden Norwegian Institute of Public Health (NIPH), Norway 2 weeks

The aim of the STM was a collaboration and knowledge exchange between Norway and Sweden in the development of One Health systems for multivariate syndromic surveillance (SyS) of veterinary and public health data combined.

Two separate systems are in development in Sweden and Norway by the SVA and the NIPH, respectively. The Swedish system is explanatory, aiming to combine several data sources to explain outbreaks and improve the accuracy of detection. Norway, on the other hand, is developing a predictive system, which aims to use some source(s) of data to predict the outcome of others.

As a test case, both systems were evaluated on the surveillance and outbreak detection of Campylobacter in humans, using data from public health, broiler chicken and weather.

Wiktor Gustafsson, SVA One Health EJP has received funding from the European Union's Horizon 2020 research and innovation programme under grant agreement No 773830.

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Figure 6: This case study can be found on the OHEJP website: <u>https://onehealthejp.eu/community/education-and-</u> training/short-term-missions-2020





Short Term Missions – Call 2021

Four STMs were awarded for 2021 but the COVID-19 pandemic continued to affect the success of these missions due to the disruption to travel plans. Two of the four STMs took place, one was post-poned to 2022 and one was cancelled.

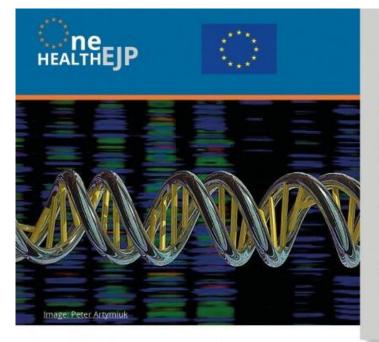
1. Short Term Mission 1

STM Researcher: Catherine Arnason Boe
Title of the STM: Start-up of an efficient sequencing facility
Theme: Skills Development Missions
Home Institute: The Norwegian Veterinary Institute, Norway
Mission Hosting Institute: Statens Serum Institut (SSI), Denmark
Duration of mission: 1 week

Aim of the mission: The project SEQ-TECH at The Norwegian Veterinary Institute (NVI) has acquired a platform of sequencing machines, pipetting robots and other equipment through 2019-2021 and now set-up for an automated flow-through of several hundred high throughput sequencing samples per week. Although the equipment was there, the routine workflow was not fully established, and the potential of the different machines not exploited. The aim of this mission was to develop skills on the management of a sequencing facility using an automated workflow.







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Start-up of an efficient sequencing facility



I will definitely use the new knowledge to advance and streamline several steps of our ongoing sequencing projects and routine workflows_The connections made with researchers and technical staff will be valuable for the future. I am very grateful to the OHEJP for granting this opportunity. I would also like to express gratitude to SSI and the kind staff for allowing me into their labs."

Cathrine Arnason Bøe,

Theme: Home Institute: Mission Hosting Institute: Duration of Mission:

Foodborne Zoonoses, AMR and Emerging threats Norwegian Veterinary Institute, Norway SSI, Denmark 1 week

The aim of this mission was to develop skills on the management of a sequencing facility using an automated workflow. The project SEQ-TECH at the Norwegian Veterinary Institute (NVI) has recently acquired equipment to set-up an automated high throughput sequencing platform. However, the routine workflow is not yet entirely established, and the potential of the different machines not fully exploited. By visiting the SSI and following their weekly routine for library preparation and high-throughput sequencing (HTS), the participant aimed to learn how to improve the NVI workflow.

During this mission, routine whole genome sequencing (WGS) of bacteria and virus (SAR5-CoV-2) were performed, the data quality control process was described and one-on-one meetings with key personnel were held. Inspiration and several ideas of how to improve routine sequencing and start novel sequencing tasks was gained by observing the SSI workflow and quality control pipeline. In addition, sharing of experiences regarding Nanopore sequencing and WGS will be valuable going forward.

The STM was a revelation for all the possibilities available at NVI, using the new equipment, to implement higher throughput sequencing for surveillance, preparedness, diagnostics and research in the near future.

Norwegian Veterinary Institute
One Health EJP has received funding from the European Union's Horizon 2020 research and innovation programme under grant agreement No 773830.



Figure 7: This case study can be found on the OHEJP website: https://onehealthejp.eu/community/education-andtraining/short-term-missions-2021





STM Researcher: Liam Burke
Title of the STM: CarbaPlasmid – Tracking endemic carbapenemase plasmids in human, animal and environmental isolates
Theme: One Health, AMR
Home Institute: NUI Galway, Ireland
Mission Hosting Institute: VISAVET, Universidad Complutense Madrid, Spain
Duration of mission: 1 month

Aim of the mission: The aim of this mission was to develop skills in nanopore sequencing and hybrid sequence analysis, to characterise antimicrobial resistance plasmids for One Health epidemiological investigations.







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CarbaPlasmid – Tracking endemic carbapenemase plasmids in human, animal and environmental isolates



This STM allowed me to learn cutting edge techniques in bacterial sequence analysis, which will greatly benefit researchers within my lab and our collaborators. I had an amazing experience in Madrid: everyone at the ARU made me feel like one of the team! This mission has paved the way for further scientific collaboration and intercultural opportunities..."

Liam Burke, NUI Galway Theme: Home Institute: Mission Hosting Institute: Duration of Mission: One Health, AMR <u>NUI Galway</u>, Ireland <u>VISAVET</u>, Universidad Complutense Madrid, Spain 1 month

The aim of this mission was to develop skills in nanopore sequencing and hybrid sequence analysis in order to characterise antimicrobial resistance plasmids for One Health epidemiological investigations.

In this project, Carbapenemase-producing Enterobacterales (CPE) isolated from the natural environment, hospital wastewater, the hospital environment and hospital patients in Galway, Ireland, were analysed by hybrid sequencing analysis. Several techniques and applications for analysis of hybrid bacterial sequence data were learned. A harmonised hybrid sequence analysis pipeline was successfully transferred between One Health EJP partners Universidad Complutense Madrid (UCA) and NUI Galway.

This short term mission developed capacity for long-read sequencing and hybrid sequence assembly in Irish OHEJP partner NUI Galway, which will also be used to support the surveillance function of the Irish National CPE Reference Laboratory. This expertise will be fundamental in ongoing and future collaborative AMR research projects with OHEJP partners and in the surveillance of endemic carbapenemase plasmids in a One-Health context in Ireland.

The data will contribute to bridging the knowledge gaps that exist in our understanding of the role of the environment in the persistence, evolution and transmission of AMR plasmids and the propensity for carbapenemase plasmid transfer between different bacterial species and strains in distinct ecological niches. It may also identify emerging AMR threats from a One Health perspective.

The training and harmonisation elements of the STM were successful, resulting in successful training of Dr Burke in nanopore sequencing and hybrid sequence analysis methodology and transfer of the protocol to his home Institution. The research outcome of the STM was also highly successful, resulting in 39 complete hybrid-assembled CPE genomes including 38 fully circularised CPE plasmids. Further analysis will result in a high quality scientific publication. Furthermore, the STM has strengthened links between NUI Galway's ARME lab and UCM's ARU lab, paving the way for further One Health collaborative research on antimicrobial resistance.

One Health EJP has received funding from the European Union's Horizon 2020 research and innovation programme under grant agreement No 773830.

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Figure 8: This case study can be found on the OHEJP website: https://onehealthejp.eu/community/education-andtraining/short-term-missions-2021





Short Term Missions – Call 2022

Ten STMs were awarded for 2022, six from the first call and four from the additional call. One of the STMs awarded in the 2021 call also took place early 2022. A total of eleven STMs were conducted in 2022.

1. Short Term Mission 1

STM Researcher: Ana Christina Ferreira

Title of the STM: Surveillance and source-attribution of AMR based on metagenomic analysis. **Theme:** Skills Development Missions.

Home Institute: National Institute for Agrarian and Veterinary Research (INIAV), Portugal. **Mission Hosting Institute:** Technical University of Denmark (DTU Food), Denmark. **Duration of mission:** 2 weeks.

Aim of the mission: The aim of this STM was to learn novel approaches and models based on metagenomic data for surveillance and to infer source attribution of AMR determinants. INIAV is the National Reference Laboratory for AMR in animals and food of animal origin and is in need of scientists with the skills to analyse and interpret metagenomics data.

Dissemination and communication activities:

The following conference talk was presented: Nogueira, T., Ferreira, A.C., Botelho, A. (2021). Comparative analysis of the resistome and the mobilome of estuarine aquacultures and other aquatic environments. Oral presentation, One Health EJP Annual Scientific Meeting 2021 (Online Event).

A research publication is also being prepared by the STM researcher, in collaboration with DTU colleagues.







Short Term Missions (STMs) are small travel grants with the aim of:

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- Contributing to the future prevention, preparedness, detection and response of the EU to foodborne and other emerging threats across human-animal-environmental sectors.

Surveillance and source-attribution of AMR based on metagenomic analysis



...it was very useful to become part of the DTU group and work together for two weeks. This collaboration allowed me to increase my knowledge and to improve skills. The information and tools acquired will be applied to the analysis of metagenomic data for source tracking in aquaculture environments and in future collaborative projects in this area " Ana Cristina Ferreira,

INIAV, Portugal

Theme: Home Institute:

Mission Hosting Institute: Duration of Mission: One Health, AMR National Institute for Agrarian and Veterinary Research (INIAV), Portugal Technical University of Denmark (DTU Food), Denmark 2 weeks

The aim of this STM was to learn novel approaches and models based on metagenomic data for surveillance and to infer source attribution of AMR determinants.

INIAV is the National Reference Laboratory for AMR in animals and food of animal origin, and is in need of scientists with the skills to analyse and interpret metagenomics data.

The STM gave Ana the opportunity to improve her skills for analysing metagenomics data for the purpose of surveillance and source attribution of AMR. Ana was engaged in the whole process from when the samples arrived in the lab to the final epidemiological data analyses, although the main focus was on the bioinformatic analyses and epidemiologically modelling.

Moreover, this mission will contribute to the development of skills in metagenomics, bioinformatics, and AMR source-attribution approaches, needed in the frame of the OHEJP consortium and in future collaborations. Use of data from omics technologies (WGS and metagenomics) for sourceattribution WGS and metagenomics based surveillance allows the characterisation of antimicrobial resistance determinants, making it possible to identify the potential for their dissemination.

The training will improve the participation of INIAV in the OHEJP DISCOVER project methods: assessment/improvement- related with source attribution of AMR based on metagenomics.

Sharing of scientific expertise and methodologies will allow the harmonisation of approaches within the OHEJP network. Furthermore, it will strengthen the collaboration and scientific capacity of partners contributing to future preparedness in the surveillance, detection and response of the EU to foodborne, AMR and emerging threats.

One Health EJP has received funding from the European Union's Horizon 2020 research and innovation programme under grant agreement No 773830.



Figure 9: This case study can be found on the OHEJP website: <u>https://onehealthejp.eu/community/education-and-</u> <u>training/short-term-missions-2022</u>.





STM Researcher: Marieke de Cock
Title of the STM: Zoonotic pathogen detection in rats (funded through the 2021 STM call, postponed to 2022 due to COVID-19).
Theme: Emerging threats.
Home Institute: National Institute of Public Health and the Environment (RIVM).
Mission Hosting Institute: Friedrich-Loeffler-Institut (FLI).
Duration of mission: 1month.

Aim of the mission: The aim of the mission was for the PhD student to test rat and mouse samples for different zoonotic pathogens. The samples were tested for Cowpox virus, rat Hepatitis E virus (rat HEV), Lymphocytic choriomeningitis virus (LCMV) and *Streptobacillus moniliformis*.

Dissemination and communication activities: The STM led to the publication of a joint research article between the institutes: Heuser, E., Drewes, S., Trimpert, J., Kunec, D., Mehl, C., de Cock, M.P., de Vries, A., Klier, C., Oskamp, M., Tenhaken, P., et al. (2023) Pet Rats as the Likely Reservoir for Human Seoul Orthohantavirus Infection. Viruses. 15(2):467. DOI: https://doi.org/10.3390/v15020467







Short Term Missions (STMs) are small travel grants with the aim of: Sharing scientific expertise,

- sharing scientific expertise, methodologies, equipment and facilities to harmonise the existing approaches and methodologies within the large
- OHEJP European network Driving the research forward in a collaborative and non-duplicative fashion to strengthen both the scientific capacity within the OHEJP
- Contributing to the future prevention, preparedness, detection and response of the EU to foodborne and other emerging threats across human-animal-environmental sectors.

Zoonotic pathogen detection in rats



Not only did this STM result in new data for my PhD project, it also strengthened the collaboration between FLI and RIVM, both in the form of sample/data sharing and publishing papers together. I can recommend all my fellow PhDs to work at another research institute for a while, not only to broaden your horizon, but also to have an unforgettable experience and to meet other researchers!"

Marieke de Cock, RIVM, The Netherlands Theme: Home Institute: Mission Hosting Institute: Duration of Mission: One Health, Emerging Threats RIVM, The Netherlands FLI, Germany 1 month

The Short Term Mission opened up the opportunity to compare and harmonise pathogen detection methods between RIVM and FLI, share reference material, and test new detection methods.

The mission focused on detection methods for specific zoonotic pathogens carried and transmitted by rats, to gain more insight in the prevalence and transmission of such infective agents that are able to cross species barriers.

The laboratory at FLI has experience in detecting specific rat-borne pathogens such as Seoul orthohantavirus, cowpox virus and rat hepatitis E virus, and it has the facilities to perform those analyses, currently not available at the RIVM. Marieke was able to learn these detection techniques, with the aim to set up these methods at RIVM.

The detection of specific rodent-borne pathogens will be included in a paper, expected towards the end of 2022, about the effect of urban greening on rodent abundance, pathogen prevalence, and diversity in wild rats from urban areas. During the STM, the team detected lymphocytic choriomeningitis virus (LCMV) in one house mouse, which is an exciting result that will be included in a separate paper about LCMV in collaboration with FLI.

As well as producing valuable test results, the mission strengthen the current collaboration between FLI and RIVM, opened up interesting discussions about rodent-borne (zoonotic) pathogen research, and led to new, unforeseen opportunities for sharing of data and the writing of papers.

The outcomes of this STM are a step forward in the harmonisation of pathogen detection methods and responses to potential emerging threats, in line with the One Health approach and vision.

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Figure 10: This case study can be found on the OHEJP website: <u>https://onehealthejp.eu/community/education-and-</u> <u>training/short-term-missions-2021</u>.





STM Researcher: Laura Gonzalez Villeta

Title of the STM: Validation and exchange of modelling tools to assess the risk of human Salmonellosis based on environmental factors using multiple sources of data.

Theme: One Health, skills development.

Home Institute: University of Surrey, UK.

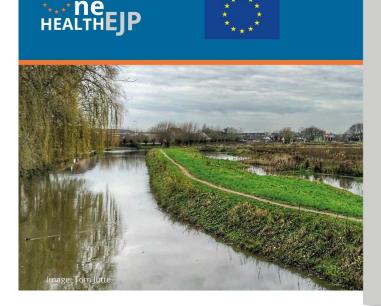
Mission Hosting Institute: Dutch National Institute for Health and Environment (RIVM). **Duration of mission:** 1 month.

Aim of the mission: The aim of this mission was to investigate whether the effect of weather on human salmonellosis cases is similar regardless of the country under study, using a novel statistical modelling approach developed at the University of Surrey, UK.

Dissemination and communication activities: The PhD student drafted a research article that includes results obtained during the STM. This work should be published soon.







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- Contributing to the future prevention, preparedness, detection and response of the EU to foodborne and other emerging threats across human-animal-environmental sectors.

Validation and exchange of modelling tools to assess the risk of human Salmonellosis based on environmental factors using multiple sources of data



The mission has not only helped me to accomplish the main objective of validating the model for my PhD but has also helped me to enhance my networking and communication capacities, opening my mind to different ways of communicating and interpreting each other. I cannot thank the OHEJP consortium enough. It was inspiring to see such motivated and skilled scientists in action applying in practice what I am learning.

Laura Gonzalez Villeta University of Surrey, UK Theme: Home Institute: Mission Hosting Institute: Duration of Mission: One Health Missions - Foodborne Zoonoses University of Surrey, UK Dutch National Institute for Health and Environment (RIVM) 1 month

The aim of this mission was to investigate whether the effect of weather on human salmonellosis cases is similar regardless of the country under study, using a novel statistical modelling approach developed in the University of Surrey, UK. The model was built on 30 years of daily epidemiological data from the UK Health Security Agency (UKHSA) and a high resolution spatio-temporal matching weather database from the MetOffice. The model first estimates the probability to observe salmonellosis cases conditional on a given combination of weather factors. Based on this information together with local weather and demographic data, the model reproduces relatively well the empirical patterns from epidemiological surveillance data for England and Wales. It also points to maximum air temperature, relative humidity, and day length as one of the most relevant combinations that influence the incidence of the disease.

During this mission, the model was applied to the Dutch setting, using 5 years of national surveillance data on salmonellosis cases reported to the Dutch National Institute of Public Health at a daily resolution and the weather variables of interest for the same period of time obtained from the open-resource Royal Netherlands Meteorological Institute (KNMI) website. To assess the universal component of the model to identify the weather-disease relationship regardless of geographical area, the probability of finding a salmonellosis case calculated for England and Wales were used. The model's results were then compared with salmonellosis disease records from The Netherlands. The preliminary results indicate that the model captures the magnitude and key seasonal patterns of the Dutch data. However, the model also results in some secondary peaks in the incidence in early spring not observed in the real data, perhaps related to differences in notification and health seeking behaviour.

The STM has improved on the relationship between two existing One Health EJP partners and enhanced both current and future collaborations between the partner institutes. A joint publication of the collaborative results should be published soon and this STM should result into a unified programme of research integrating the two approaches over diverse geographic and socio-economics settings.

One Health EJP has received funding from the European Union's Horizon 2020 research and innovation programme under grant agreement No 773830.

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Figure 11: This case study can be found on the OHEJP website: <u>https://onehealthejp.eu/community/education-and-</u> <u>training/short-term-missions-2022</u>.





STM Researcher: Antonio Rodriguez

Title of the STM: Application of spatial models to identify new environmental surveillance indicators of *Salmonella* and *Campylobacter* in pig and poultry.

Theme: One Health, skills development.

Home Institute: The National Institute for Agricultural and Food Research and Technology (INIA), Spain.

Mission Hosting Institute: National Institute for Public Health and the Environment (RIMV), The Netherlands.

Duration of mission: 1 month.

Aim of the mission: The aim of this mission was to develop a procedure to model surveillance data of zoonotic diseases using machine learning methods. The model was tested on the Wild Boar Surveillance Plan of Andalucía (Spain) data, with the aim to determine the most important risk factors that shape wild boar diseases in Andalucía and study their interaction between each other.

Dissemination and communication activities: The STM researcher trained their team members on the methods learnt during the STM. They also organised a teleconference with people in charge of the Wildlife Epidemiological Surveillance Program of Andalusia (Spain) at the Water and Environment Agency from Andalusia and the Animal Health Research Centre, to give them feedback about the data in order to strengthen the surveillance program.







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- Contributing to the future prevention, preparedness, detection and response of the EU to foodborne and other emerging threats across human-animal-environmental sectors.

Application of spatial models to identify new environmental surveillance indicators of Salmonella and Campylobacter in pig and poultry



I had the chance to meet a new research centre and to collaborate with and take advice from experts on the field. We obtained results that we could not have obtained working independently. I carried out an important advance in my research project and acquired new skills that we will use in future projects. I would like to highlight that thanks to my shortterm mission I pushed my data analysis skills to an upper level."

Antonio Rodríguez The National Institute for Agricultural and Food Research and Technology Spain Theme: Home Institute:

Mission Hosting Institute:

Duration of Mission:

One Health Missions - Skills development missions The National Institute for Agricultural and Food Research and Technology (INIA), Spain National Institute for Public Health and the Environment (RIMV), The Netherlands 1 month

The aim of this mission was to develop a procedure to model surveillance data of zoonotic diseases using machine learning methods. The model was tested on the Wild Boar Surveillance Plan of Andalucía (Spain) data, with the aim to determine the most important risk factors that shape wild boar diseases in Andalucía and study their interaction between each other.

During this mission, the model validation was performed in two main steps. First, a single response variable was selected for the study: the *Salmonella* serology of hunted wild boars. In the first week of the STM, the data was organised in an appropriate way for training machine learning algorithms. Additionally, climate and wild boar distribution information from published maps were collected, to include them as potential drivers of the pathogens. The results being conclusive, a modelling procedure valid for all the wild boar diseases in the dataset was designed. This includes the processing of the dataset before the modelling (pre-processing step), the model application itself and its validation, and the study of the model output. The best way to compensate the unbalance of the positives and negatives in the dataset was explored, to avoid reducing the sensitivity or specificity of the model. A script was prepared to provide several validation statistics based on the comparison between the predictions on the test data and the real positive and negative values. The variable importance was assessed on the training and test data, partial dependence plots based on these data were created and detection of interactions between predictors on the test data was conducted. The developed procedure successfully determined the most important risk factors that shape wild boar diseases in Andalucía and studied their interaction between each other. Using the data from previous hunting seasons, it also allowed to identify areas with higher and lower probabilities of boar disease and to represent the predictions in a map.

The STM has improved on the relationship between two existing One Health EJP partners and enhanced both current and future collaborations between the partner institutes. Additionally, this procedure has been automatised and can be easily applied to the data from the *Salmonella* and *Campylobacter* surveillance in the food chain by the Spanish Agency of Food Security and Nutrition (AESAN) and the *Salmonella* and *Campylobacter* Spanish surveillance plan in poultry farm.

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Figure 12: This case study can be found on the OHEJP website: <u>https://onehealthejp.eu/community/education-and-</u> training/short-term-missions-2022.





STM Researcher: Ingrid Cardenas Rey

Title of the STM: Construction of dual labelled *E. coli* strains to study the effect of antibiotics and microbiota interventions on the horizontal transfer of ESBL genes in the *in vitro* chicken caecal microbiota.

Theme: One Health Missions - Antimicrobial Resistance (AMR).

Home Institute: Wageningen Bioveterinary Research (WUR), The Netherlands.

Mission Hosting Institute: University of Copenhagen, Denmark.

Duration of mission: 2 months.

Aim of the mission: The aim of this mission was for the PhD student to learn bacterial cloning methods using fluorescent reporter proteins, to produce dual labelled *E. coli* strains. Dual labelling (fluorescent tagging of the chromosome and AMR-gene carrying plasmids) of bacteria is a powerful tool to study plasmid-mediated antimicrobial resistance among complex *in vitro* microbial communities simulated on *in vitro* gut systems like the chicken caeca.

Dissemination and communication activities: The PhD student presented the results obtained thanks to the STM at a poster carousel of the microbial ecology group in June 2022 at a workshop in Nunspet (The Netherlands) and at the OHEJP side event "Lessons Learnt from a European Multidisciplinary Initiative" at the ONE2022 Conference in June 2022 in Brussels (Belgium).







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Construction of dual labelled E. coli strains to study the effect of antibiotics and microbiota interventions on the horizontal transfer of ESBL genes in the in vitro chicken caecal microbiota



A short term mission should be a must-do for all young researchers! It was a priceless experience that enriched my knowledge and skillsets and expanded my network. I built up and exercised my lab skills, critical thinking and cooperation in a multidisciplinary group. I was challenged daily with new knowledge and experimental work but also strongly supported by a wonderful work team!"

Theme: Home Institute: Mission Hosting Institute: Duration of Mission: One Health Missions - Antimicrobial Resistance (AMR) Wageningen Bioveterinary Research (WUR), The Netherlands University of Copenhagen, Denmark 2 months

The aim of this mission was for the PhD student to learn bacterial cloning methods using fluorescent reporter proteins, to produce dual labelled *E. coli* strains. Dual labelling (fluorescent tagging of the chromosome and AMR-gene carrying plasmids) of bacteria is a powerful tool to study plasmid -mediated antimicrobial resistance among complex *in vitro* microbial communities simulated on *in vitro* gut systems like the chicken caeca. This STM enabled the PhD student to reach the objectives proposed in the OHEJP PhD project VIMOGUT, which studies the chicken gut microbiota and microbiota interventions to reduce horizontal transmission of Extended Spectrum ß-Lactam (ESBL) genes.

During this mission, five commensal ESBL *E. coli* strains that originated from chicken broiler caeca and belonged to the collection of The Dutch National Reference Lab were used for bacterial cloning experiments. All strains carried an ESBL (blaCTXM-1 and blaSHV-12) or AmpC β-lactamase (blaCMY-2) gene on plasmids highly prevalent in the broilers production and were susceptible to three antimicrobials. Two fluorescent reporter proteins were used for chromosome and plasmid tagging, namely, mCherry and Green fluorescent protein (GFP). Bacterial cloning is a challenging research area that requires time and repeatability of the experimental work. Technical issues and challenges were encountered during this multi-step process. However, the PhD student learned the bacterial cloning techniques needed for the completion of future *in vitro* experiments and developed and strengthened her lab skills.

The STM opened significant cooperation channels between the AMR group at Wageningen Bioveterinary Research and the One Health Antimicrobial Resistance (OHAR) research group at the University of Copenhagen. The output of the work performed during the STM, and upcoming in vitro experiments are expected to be published as a collaborative research article in a scientific journal.

Ingrid Cardenas Rey Wageningen Bioveterinary Research, The Netherlands

arch, One Health EJP has received funding from the European Union's Horizon 2020 research and innovation programme under grant agreement No 773830.

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Figure 13: This case study can be found on the OHEJP website: <u>https://onehealthejp.eu/community/education-and-</u> <u>training/short-term-missions-2022</u>.





STM Researchers: Albena Dimitrova Angelova, Gergana Mateva et Mihail Vladimirov Milanov
Title of the STM: Microbiological and molecular techniques for *Brucella* identification.
Theme: One Health Missions.
Home Institute: NDVRI, Bulgaria.
Mission Hosting Institute: Anses, France.
Duration of mission: 1 week.

Aim of the mission: The aim of this mission was for the three researchers to learn standardised microbiological and molecular techniques, including DNA extraction and real-time PCR, for the identification of *Brucella spp*. As Brucellosis is the among the leading lab-acquired infections, learning effective biosafety/ biosecurity whilst working with the causative agent is paramount. This STM enabled the researchers to complete objectives in the OHEJP IDEMBRU project, which studies the zoonotic potential, virulence, and persistence markers in isolated strains of *Brucella spp*.







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Microbiological and molecular techniques for Brucella identification

Theme: Home Institute: Mission Hosting Institute: Duration of Mission: STM Researchers: One Health missions NDVRI, Bulgaria ANSES, France 1 week Albena Dimitrova Angelova Gergana Mateva Mihail Vladimirov Milanov

The aim of this mission was for the three researchers to learn standardised microbiological and molecular techniques, including DNA extraction and real-time PCR, for the identification of *Brucella spp*. As Brucellosis is the among the leading lab-acquired infections, learning effective biosafety/ biosecurity whilst working with the causative agent is paramount. This STM enabled the researchers to complete objectives in the OHEJP IDEMBRU project, which studies the zoonotic potential, virulence and persistence markers in isolated strains of *Brucella spp*.

During this mission, theoretical training was provided by the host institute on the critical steps for *Brucella* culture followed by principles of molecular approaches. Prior to the researchers undertaking any laboratory activities, practical demonstrations on bacterial isolation, culture and typing were provided, followed by demonstrations of molecular techniques including DNA extraction, real-time PCR protocols, gel electrophoresis and High-Resolution Melting PCR. Each day, round table discussions provided additional insight into each step of the training. A final presentation on molecular epidemiology afforded further knowledge on applications, perspectives and interpretation of data. The researchers strengthened existing and developed new laboratory skills that will assist their future research.

The STM provided valuable training, allowing for the isolation and identification of *Brucella spp.*; further experimentation will be performed on *Brucella spp.* strains, under rigorous biosafety standards, at the NDVRI laboratories in Bulgaria.

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Figure 14: This case study can be found on the OHEJP website: <u>https://onehealthejp.eu/community/education-and-</u><u>training/short-term-missions-2022</u>.

C The STM was an amazing

experience and has provided newfound knowledge regarding the skill and requirements for microbiological manipulation and molecular diagnostics, especially when working with potentially infectious bacteria, such as Brucella. Disseminating this knowledge to other colleagues will further help and enhance everyday tasks in our laboratories'

Albena Dimitrova Angelova, NDVRI, Bulgaria





STM Researcher: Emma Brook

Title of the STM: Tolerance of biofilm forming bacteria to disinfectants after repeated disinfectant exposure.

Theme: One Health Missions- Veterinary, Food, Medical and or Environmental research. **Home Institute:** Animal and Plant Health Agency (APHA), UK. **Mission Hosting Institute:** Norwegian Veterinary Institute (NVI), Norway.

Duration of mission: 3 weeks.

Aim of the mission: The aim of this mission was to test methods for measuring disinfectants efficacy, after repeated use on bacteria that survived treatment due to the formation of biofilms. Methods for disinfectant efficacy testing against planktonic foodborne bacteria are well established, yet methods for efficacy of disinfectants for biofilms are still novel, particularly the effect of repeated exposure to disinfectants on the survivability of bacteria in biofilms. This STM aimed to promote harmonisation of current test methods and models used in biofilm and disinfectant testing.







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- Contributing to the future prevention, preparedness, detection and response of the EU to foodborne and other emerging threats across human-animal-environmental sectors.

Tolerance of biofilm forming bacteria to disinfectants after repeated disinfectant exposure



My personal and professional confidence grew during the STM. The information shared has been invaluable to my learning and development. I have gained experience in a wider range of methods used in biofilm and disinfectant research, as well as with a species of bacteria. We exchanged skills and to allow harmonisation of approaches. The methods and techniques learnt will be applied to ongoing and future projects back at my home institute...." Emma Brook APHA, UK

Theme:

Home Institute: Mission Hosting Institute: Duration of Mission: One Health Missions- Veterinary, Food, Medical and/or Environmental research Animal and Plant Health Agency (APHA), UK Norwegian Veterinary Institute (NVI), Norway 3 weeks

The aim of this mission was to test methods for measuring disinfectants efficacy, after repeated use on bacteria that survived treatment due to the formation of biofilms. Methods for disinfectant efficacy testing against planktonic foodborne bacteria are well established, yet methods for efficacy of disinfectants for biofilms are still novel, particularly the effect of repeated exposure to disinfectants on the survivability of bacteria in biofilms. This STM aimed to promote harmonisation of current test methods and models used in biofilm and disinfectant testing.

During this mission, biofilms of Staphylococcus aureus were exposed to different concentrations of Benzalkonium chloride disinfectant. Following neutralisation of the disinfectant, biofilm was removed via mechanical action. Persisting bacteria were enumerated and isolated to prepare next-generation biofilms. These were then exposed to further disinfection using the same concentration to which they were previously exposed. This was repeated over nine consecutive experiments. The results have shown that increased tolerance to biocides by biofilm forming bacteria can occur. However, conflicting statistical results suggest the need for further investigation. Future studies should expand on the data collected in this study, to help develop knowledge on the role of biofilm producing bacteria in the dissemination of biocide resistance and factors affecting the variability in biofilm behaviour.

The STM has improved on the relationship between two existing One Health EJP partners and enhanced both current and future collaborations between the partner institutes. The STM has supported collaborative research in the current project (BIOPIGEE) through improvements of working arrangements and understanding of the used methods.

One Health EJP has received funding from the European Union's Horizon 2020 research and innovation programme under grant agreement No 773830.

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Figure 15: This case study can be found on the OHEJP website: <u>https://onehealthejp.eu/community/education-and-</u> <u>training/short-term-missions-2022</u>.





STM Researcher: Carlos Sacristán Yagüe
Title of the STM: Metagenomics tools for the study of SARS-COV-2 in animals.
Theme: One Health, Skills Development Missions.
Home Institute: INIA, Spain.
Mission Hosting Institute: INSA, Portugal.
Duration of mission: 2 weeks.

Aim of the mission: The aim of this mission was to expand the researcher's training to a higher and more practical level by learning the application of metagenomics and completing sequencing within a One Health context. These tools are essential for modern surveillance and diagnostics of infectious diseases and are key to shed light on the epidemiology of novel viruses.







Short Term Missions (STMs) are small travel grants with the aim of:

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- OHEJP European network Driving the research forward in a collaborative and non-duplicative fashion to strengthen both the scientific capacity within the OHEJP
- Contributing to the future prevention, preparedness, detection and response of the EU to foodborne and other emerging threats across human-animal-environmental sectors.

Metagenomic Tools for the Study of SARS-CoV-2 in Animals



...a great personal and professional experience. I closely followed the group's daily activities, and performed specific training activities designed to maximize my learning experience. I had the opportunity of getting acquainted with the National Health System of Portugal, and participated in several cultural and scientific activities carried out at INSA during my stay. I gained valuable skills, that will allow me to use metagenomics in my future research. Carlos Sacristán Yagüe INIA, Spain

Theme: Home Institute: Mission Hosting Institute: Duration of Mission: One Health, Skills Development Missions INIA, Spain INSA, Portugal 2 weeks

The aim of this mission was to expand the Carlos' training to a higher and more practical level by learning the application of metagenomics and completing sequencing within a One Health context. These tools are essential for modern surveillance and diagnostics of infectious diseases and are key to shed light on the epidemiology of novel viruses.

During the mission, Carlos had the opportunity to learn how to use the INSaFLUTELE-VIR suite: an easy-to-use open web-based bioinformatics suite that was designed for the genomic surveillance of human seasonal influenza and SARS-CoV-2, and recently adapted to monkey pox virus (now renamed Mpox). This platform was proven to be an excellent surveillance tool during the SARS-CoV-2 pandemic and a new module has been designed by researchers in the mission hosting institute to simplify metagenomic analyses and identify new viruses. Carlos also learned about the ReporTree tool, a surveillance-oriented tool to strengthen the linkage between pathogen genetic clusters and epidemiological data and was trained on the use of nanopore sequencing with MinION. The MinION was used during the mission for the complete sequencing of SARS-CoV-2, the diagnostics of MPox virus and sequencing of a poliovirus. The "Young Researcher Day" and "INSA Day" took place during the mission, showcasing scientific and cultural activities. It allowed Carlos to expand his professional network.

The mission provided Carlos the training to diagnose and characterize future SARS-CoV-2 cases in animals using metagenomics, which is key to determine whether a virus has changed/mutated during a spill over event. One of the outputs of the project will be the adaptation of modern metagenomics and bioinformatics tools in the COVRIN project. The mission also strengthened further the collaboration between the institutes, with a new collaboration established to research on hepatitis E virus in wildlife.

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Figure 16: This case study can be found on the OHEJP website: <u>https://onehealthejp.eu/community/education-and-</u> <u>training/short-term-missions-2022</u>.





STM Researchers: Sandrine Baron and Laetitia Le Devendec
Title of the STM: Genotypic characterisation of antimicrobial susceptibility and isolation of *Aeromonas* and *Vibrio* phages from water samples.
Theme: One Health, Skills Development Missions.
Home Institute: Anses, France.
Mission Hosting Institute: BfR, Germany.
Duration of mission: 4 weeks.

Aim of the mission: The objective of the mission was to share technical knowledge on the detection and characterisation of *Aeromonas* and *Vibrio* genus and to establish a collaboration between the institutes, to study of the dissemination of antibiotic resistance in aquatic environments.







Short Term Missions (STMs) are small travel grants with the aim of:

- Sharing scientific expertise, methodologies, equipment and facilities to harmonise the existing approaches and methodologies within the large
- OHEJP European network Driving the research forward in a collaborative and non-duplicative fashion to strengthen both the scientific capacity within the OHEJP
- Contributing to the future prevention, preparedness, detection and response of the EU to foodborne and other emerging threats across human-animal-environmental sectors.

Genotypic characterisation of antimicrobial susceptibility and isolation of Aeromonas and Vibrio phages from water samples

This mission was a great opportunity to discuss the dissemination of antibiotic resistance in the aquatic environment in connection with aquaculture activities. In addition to the improvement of the technical skills of both teams, the diversity of our fields of study (water, fish farming vs food) opened up perspectives for new collaborations, using a One Health approach."

Sandrine Baron and Laetitia Le Devendec ANSES, France Theme: Home Institute: Mission Hosting Institute: Duration of Mission: One Health, Skills Development Missions ANSES, France BfR, Germany 4 weeks

The objective of the mission was to share technical knowledge on the detection and characterisation of *Aeromonas* and *Vibrio* genus and to establish a collaboration between the institutes, to study of the dissemination of antibiotic resistance in aquatic environments.

During this mission, Sandrine and Laetitia were trained on phages' cultivation, enumeration and conservation methods. They now have in hands the detailed protocols, as well as phages and competent bacteria cultures, provided by the BfR team. They will be able to put the methods into practice in Anses. Sandrine and Laetitia travelled to BfR with culture media for the detection of *Aeromonas sp.*, from water and fish samples. Water samples have been collected nearby the BfR laboratory and the German team performed the entire handling: filtration, cultivation and presomptive identification. The partner teams established common protocols for the study of the *Aeromonas* and *Vibrio* genera and initiated the sharing of the Maldi-Tof databases, for the improvement of the species identification of bacteria from the *Aeromonas* and *Vibrio* genera. The two institutes are also now both involved in a research group, which aims to improve the methodology of antimicrobial susceptibility testing in aquatic bacteria and determine Epidemiological cut off values (Ecoff). Meetings and video conferences with French and Canadian researchers took place, to discuss research ideas. At the end of this STM, two research project proposals were elaborated.

This short-term mission strengthened the emerging collaboration between the two teams from the One Health EJP consortium. It has also led to a reflection on the use of phages to improve and reduce the use of antibiotics in aquaculture, a farming sector from which a One Health approach is absolutely needed.

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Figure 17: This case study can be found on the OHEJP website: <u>https://onehealthejp.eu/community/education-and-</u> <u>training/short-term-missions-2022</u>.



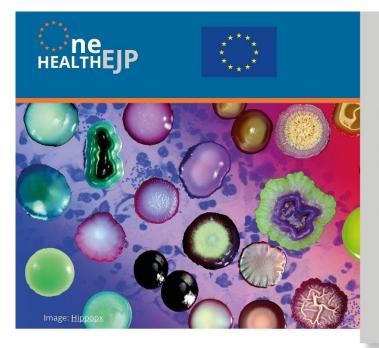


STM Researcher: Eve Zeyl Fiskebeck
Title of the STM: Stable Intra-specific genomic bacterial classification systems.
Theme: One Health Missions.
Home Institute: Norwegian Veterinary Institute (NVI), Norway.
Mission Hosting Institute: Institut Pasteur, France.
Duration of mission: 4 days.

Aim of the mission: The aim of this mission was for the researcher to gain knowledge on the usage and design of a stable classification system and nomenclature approach for bacterial strains and to understand advantages and limitations of its use in genomic epidemiology. Previous isolate nomenclature/classification systems including serotyping and MLST typing do not always satisfy two major criteria: long-term stability and congruence with phylogenetic relationships, which are essential for routine epidemiological surveillance.







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Stable Intra-specific genomic bacterial classification systems



The STM allowed me to understand and practice applying the newly developed cgLIN bacterial nomenclature system. I now have a better understanding on how the classification system can be used for genomic epidemiology of zoonotic and infectious diseases and able to disseminate this knowledge to other scientists. This will strengthen the activities in One Health to address existing and emerging threats of foodborne and zoonotic agents as well as animal pathogens" Eve Zeyl Fiskebeck NVI, Norway

Theme: Home Institute: Mission Hosting Institute: Duration of Mission: One Health <u>Norwegian Veterinary Institute</u> (NVI), Norway <u>Institut Pasteur</u>, France 5 days

The aim of this mission was for the researcher to gain knowledge on the usage and design of a stable classification system and nomenclature approach for bacterial strains and to understand advantages and limitations of its use in genomic epidemiology.

Previous isolate nomenclature/classification systems including serotyping and MLST typing do not always satisfy two major criteria: long-term stability and congruence with phylogenetic relationships, which are essential for routine epidemiological surveillance. During this mission, theoretical training was provided by the host institute, to gain deep understanding and practical experience of a novel classification system and nomenclature approach for bacterial strains (cgLIN codes). Training also allowed further understanding of the advantages and limitations of its use in genomic epidemiology.

The STM provided valuable training, allowed knowledge transfer and strengthened the collaboration between One Health partners. Further funding will be sought to continue this valuable collaboration, in order to further develop and expand the nomenclature system for additional bacterial pathogens.

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Figure 18: This case study can be found on the OHEJP website: <u>https://onehealthejp.eu/community/education-and-</u><u>training/short-term-missions-2022</u>.





STM Researcher: Bosco Rodríguez Matamoros
Title of the STM: Application of 'Single-Cell Genomics' for the study of the bacterial reservoirs of plazomicin resistance determinants.
Home Institute: Universidad Complutense de Madrid, Spain.
Mission Hosting Institute: Karlsruhe Institute of Technology, Germany.
Duration of mission: 3 months.

Aim of the mission: The aim of this mission was the development of a single cell workflow, for the detection of low abundant reservoirs of next-generation aminoglycoside resistance mechanisms. The main objective was to label single cells from complex environments with fluorescence markers specific for the genes of interest. This Short-Term Mission was framed within the OHEJP PhD METAPRO project and was expected to produce results complementing the ones already produced with the metagenomic analyses.







Short Term Missions (STMs) are small travel grants with the aim of: Sharing scientific expertise,

- Sharing scientific expertise, methodologies, equipment and facilities to harmonise the existing approaches and methodologies within the large
- OHEJP European network Driving the research forward in a collaborative and non-duplicative fashion to strengthen both the scientific capacity within the OHEJP
- Contributing to the future prevention, preparedness, detection and response of the EU to foodborne and other emerging threats across human-animal-environmental sectors.

Application of 'Single-Cell Genomics' for the study of the bacterial reservoirs of plazomicin resistance determinants



Thanks to the One Health EJP I have had the chance to expand my skill sets and learn new methods. Having the opportunity to know a new research centre and discuss my project with very talented people has helped me expand my critical thinking and strengthen my professional confidence. We built a long-lasting collaborative network that I trust can give promising results in the close future.

Bosco Rodríguez Matamoros Universidad Complutense de Madrid, Spain Theme:

Home Institute: Mission Hosting Institute: Duration of Mission: One Health Missions , Skills Development missions, Antimicrobial Resistance (AMR) <u>Universidad Complutense de Madrid</u>, Spain <u>Karlsruhe Institute of Technology</u>, Germany 3 months

The aim of this mission was for the PhD student to learn about the development of single cell workflows, for the detection of low abundance reservoirs of next-generation aminoglycoside (such as plazomicin) resistance mechanisms. The main objective was to label single cells from complex environments with fluorescence markers specific for the genes of interest. This STM enabled the PhD student to produce results that complemented the ones already produced within the OHEJP PhD METAPRO project using metagenomic analyses.

During this mission, three different labelled probes were designed for the detection of the aminoglycoside resistance gene *npmA*, a gene that confers high level of resistance to all known aminoglycosides, including plazomicin, apramycin and other next-generation aminoglycosides. To test the probes, the *npmA* resistance gene was introduced in two plasmids with different copy numbers to use as positive controls, and the same plasmids without the resistance gene were used as negative controls. All these control plasmids were introduced in *E. coli* cells and a classical fluorescence in situ hybridisation (FISH) protocol was performed with the three probes independently to check their labelling efficiency. Two of the designed probes showed promising results and have the potential to be used with environmental samples. Samples have been taken to evaluate the potential sorting of *npmA* positive cells and is planned to be performed in the upcoming months.

This STM has opened a new collaboration channel between two research groups with different scopes for the study of antimicrobial resistance with a One Health approach. The collaboration between the partner institutes is expected to last longer than the extend of the mission and we expect to produce interesting results than could potentially be published as a research article in a scientific journal.

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