

PATH-SAFE

Pathogen Surveillance in Agriculture, Food and Environment (PATH-SAFE) Programme

PATH-SAFE is a 4-year, cross government and UK wide programme, led by the Food Standards Agency (FSA) and utilising £24m funding from the HMT Shared Outcomes Fund (SOF) and match funding from a range of government and academic delivery partners. The programme is piloting the development of a national surveillance network, using the latest DNA-sequencing technology and environmental sampling, to improve the detection and tracking of foodborne human pathogens and associated AMR through the whole agri-food system from farm-to-fork.

Programme background

Foodborne disease (FBD) is a major public health risk with 2.4 million individual illnesses and more than 16,000 hospitalisations per year. Most human disease is caused by a handful of bacteria which enter the food chain from farmed animals or the environment.

In addition to FBD, the agri-food supply chain also poses a risk for the transmission of antimicrobial resistance (AMR) through food, animals, humans, and water. Whilst the UK has made progress in reducing its use of antibiotics in humans and animals in the last five years, [drug-resistant bloodstream infections in humans have increased by 32% from 2015 to 2019](#). The rise and spread of AMR is creating a new generation of 'superbugs' that cannot be treated with existing medicines.

For these reasons, government departments already undertake surveillance activities by analysing samples from food, livestock, and humans. Advances in technology and data management offer the opportunity to create a step change in surveillance, to protect public health. The PATH-SAFE programme is establishing a new data platform that will allow for the analysis, storage and sharing of pathogen sequence and source data, collected from multiple locations across the UK by government departments and public organisations. This single system will enable rapid identification and tracking of FBD and AMR. This will improve public health and minimise the economic and environmental impact of outbreaks.

Aims of the programme

1. To pilot a better national surveillance system for the monitoring and tracking of FBD and AMR in the environment and agri-food system
2. To bring together and build on existing initiatives across the UK and to understand what the end-user needs to improve how they work in this space
3. To provide better data to identify the prevalence, source and pathways of FBD and AMR, helping to prevent spread by enhanced targeting of interventions

Benefits of the programme

1. Generation of information and knowledge that support enhanced identification of FBD and AMR sources
2. Providing improved user access to relevant data so they can make more informed, evidence-based decisions
3. Data that underpins the prevention and prediction of FBD and AMR spread by supporting improved, cost-effective targeting of interventions, providing economic savings both for government and industry
4. Improved surveillance system that supports better identify and quicker reaction to emergent diseases (or diseases of increasing concern).
5. Potential to reduce the number and cost of FBDs and AMR, lower commercial losses, strengthen UK Science Excellence, and enhance the UK food sector's reputation

Programme Partners

The PATH-SAFE programme is led by the FSA, supported by several core partners:

- Food Standards Scotland (FSS)
- Department of Environment, Food and Rural Affairs (Defra)
- Animal and Plant Health Agency (APHA)
- Centre for Environment, Fisheries and Aquaculture Science (Cefas)
- Environment Agency (EA)
- Veterinary Medicines Directorate (VMD)
- Department of Health and Social Care (DHSC)
- UK Health Security Agency (UKHSA)

The programme has been delivered through a one-health, four-nation collaborative approach, both at the delivery level and at the strategic level.

More than 50 organisations, from across government, academia and industry, have been involved in delivery and advisory roles. These have included:

Funding source and timeline

PATH-SAFE is funded via the [Shared Outcomes Fund \(SOF\)](#).

Spending Round 2019 announced £200 million for the Shared Outcomes Fund to fund pilot projects to test innovative ways of working across the public sector. The first round of the SOF funded a wide range of projects to be run in 2020-21 and 2022-23.

Spending Review 2020 announced a further £200 million was to be made available for a second round of the Shared Outcomes Fund between 2021-22 and 2023-24.

In 2021 the programme successfully secured £19.2m in phase 2 of the SOF, to run a three-year programme concluding in March 2024.

At the start of 2024, PATH-SAFE successfully secured an additional £4.7m of funding to continue for an additional year (April 2024 to March 2025), with HMT SOF providing £2.2m of the continuation funding, leveraged by £2.5m of match funding from PATH-SAFE core government partner departments and academic delivery partners.

Programme Themes

The programme is organised thematically for governance, oversight, reporting and coordination purposes.

Further information about each of the themes, including overviews, delivery partners, status, key successes and outputs can be found on the following pages:

- [National foodborne disease genomic data platform](#)
- [Onsite Diagnostics](#)
- [Novel Foodborne Pathogen, and associated antimicrobial resistance, knowledge and surveillance approaches](#)
- [Programme Evaluation](#)

The programme has developed new systems and tools, explored novel surveillance approaches and generated knowledge, to improve the detection and tracking of foodborne human pathogens and antimicrobial resistance, through the whole agri-food system from farm-to-fork.

Alongside the technical advances the programme has driven forward, PATH-SAFE has also established and promoted a collaborative, four nation, One Health approach to surveillance, which has been referenced as an exemplar within initiatives such as the Biological Surveillance Strategy and the UK AMR National Action plan (noted within as a key achievement).

The programme has influenced, informed, and supported programmes, strategies and initiatives across government and beyond, including:

- Biothreats Radar and National Biosurveillance Network (NBN)
- UK Microbial Forensics Consortium (UKMFC)
- Genomics of Animal and Plant Disease Centre Phase 2 (GAP-DC2)
- AMR National Action plan (NAP) 2019-24 and 2024-29
- UKRI Tackling Infections Strategic Theme
- Government Office for Technology Transfer (GOTT)
- Microbiology Society – Knocking Out AMR initiative
- ONS Integrated Data Service (IDS) Programme
- Food Safety Research Network (FSRN)

Events and communications

Throughout the programme, a range of events and communication activities have been undertaken including a webinar series, a conference and publication of quarterly newsletters.

Further information on these events and activities can be found on the following pages:

- [Webinar series and newsletters](#)
- [PATH-SAFE biosurveillance conference \(February 2024\)](#)

Get in touch

If you would like to get in touch with the PATH-safe team, you can email them at pathsafe@food.gov.uk

PATH-SAFE programme National foodborne disease genomic data platform

At the heart of PATH-SAFE programme is the Genomic Data Platform, a user-friendly platform for the analysis and visualisation of bacterial genomes and associated metadata.

Overview

The UK is recognised global leader in genomic database systems. We are utilising this existing expertise, working with academic colleagues and major 'big data' stakeholders, to create a user-friendly platform for the rapid interrogation and contextualisation of genomic data.

While the rapid detection, identification and tracking of pathogens has always been fundamental to public health, it is only in recent years, with advances in DNA-based methods, that we now have the means to do this. As noted in the UK National Data Strategy, innovation has brought with it an exponential growth in data; in its generation and use, and in the world's increasing reliance upon it.

By embracing data and the benefits its use brings, the UK now faces tangible opportunities to improve our society and grow our economy. It can transform our public services and dramatically improve health outcomes nationally. Modern genome sequencing offers the potential to do just that; by identifying pathogen strains rapidly and cheaply with a high resolution and from this reconstruct chains of transmission and trace outbreaks to their source. However, we need the data infrastructure to properly enable this.

The PATH-SAFE genomic data platform ultimately seeks to address this need at a national level by providing a cross-government genomics capability which can perform analysis of genomic data and associated metadata to facilitate rapid identification of pathogen strains of interest and support elucidation of transmission pathways.

The platform aims to realise the following benefits:

- improve the tracking of FBD pathogens and associated AMR through faster identification of variants of interest, thereby increasing outbreak investigation efficiency
- through analysis, bring together currently unconnected government and public genomic pathogen data to maximise the use of available FBD biosurveillance data and thereby expedite variant identification
- create efficiencies in cross-government working in FBD biosurveillance through data sharing and collaboration via the platform
- reduce public health risks associated with foodborne disease outbreaks by tracing sources of infection faster, potentially facilitating faster removal of those sources
- reduce economic losses associated with FBD by reducing the number of missed, slow or inaccurate assessments of outbreak cause, thereby reducing the costs/resources associated for people affected by FBD (e.g. sick days required) and by businesses (e.g. additional administrative costs of dealing with an outbreak)

Delivery Partners

Our delivery partners for this platform are:

- Digital Epidemiology Services
- University of Oxford
- University of Warwick
- University of Birmingham

Status

The data platform, with analytics for Salmonella, was launched for use by relevant government departments in March 2024 and available for use by relevant government organisations.

Work in 24/25 will focus on the integration of data, as well as the addition of analytics for E. coli and Listeria.

End user engagement with key government partner organisations restarted in May 2024, with discussions about platform requirements for the new pathogens to be added (E. coli and Listeria) continuing over the summer. The delivery team (Digital Epidemiology Services) met with the team from CLIMB BIG-DATA, who are providing the secure infrastructure on which the platform has been built, for an in-person FY24/25 planning session at University of Birmingham.

An evaluation piece being conducted by Enterobase, to assess existing genome assembly pipelines for E. coli and Listeria with the aim of making a recommendation on which should be implemented in the platform, has also begun and will form a key first step in the development

process.

Key successes

Key success of this work include:

- following a 15-month development period, the first iteration of the genomic data platform was made live in March 2024, ready for submission and analysis of Salmonella genomes by participating government partner agencies
- the platform and its functionality has built upon existing and proven surveillance tools and infrastructure, with [CLIMB-BIG-DATA](#) providing the cloud infrastructure used by the platform, [Pathogenwatch](#) tools supplying the platform's analytics capability and a call-out feature to [Enterobase](#) supporting the platform's context genome search functions
- a discovery phase was completed prior to development being initiated which sought to elucidate existing surveillance infrastructure and processes to inform platform design
- extensive end user engagement undertaken during platform development, ahead of release of the first iteration of the platform (another opportunity will be provided ahead of release of the second iteration)
- set-up of four advisory groups (Technical, AMR, Data Standards and International) which provided recommendations on the design and capabilities of the platform and on the application, technical requirements of and best practices for WGS in Salmonella FBD surveillance in general - a white paper summarising the recommendations has been collated and is due for publication in Q3 2024

Outputs

All projects were presented at the PATH-SAFE Biosurveillance Conference on 28 and 29 February in London. [Recordings, slides and posters are available on the conference webpage.](#)

PATH-SAFE: Onsite Diagnostics

Alongside the development of laboratory-based approaches and methodologies, undertaken within a separate theme, this PATH-SAFE theme has specifically considered the application of onsite diagnostics (OSD) for the rapid detection of foodborne pathogens (FBP), outside of a conventional laboratory setting.

Overview

The advancement of rapid and accurate onsite diagnostic methods provides a real opportunity in relation to detection and control of foodborne pathogens.

The first phase of this work undertaken in 22/23 and 23/24 focused on the identification of promising technologies that could be piloted for on-site testing of foodborne pathogens, including piloting selected technologies.

A comprehensive literature review was undertaken to identify technologies under development for on-site testing. This included both technologies proven or presumed to be applicable in detecting targets relevant to food safety, as well as those developed for other sectors. These technologies were assessed through a customised technology readiness level (TRL) framework with each technology assigned a TRL in combination with each in-scope pathogen or matrix.

To understand the testing requirements in real world scenarios, both "strategic" and "operational" stakeholders were mapped and engaged in focus groups and interviews. This provided a broad perspective on on-site testing and supported identification of sectors or processes that are well-suited or not suited for portable detection technologies.

Results from the literature review and end-user study led to the shortlisting of technologies based on relative maturity and suitability for on-site use, mapped against potential applications within the food sector for deployment of the technologies. This decision matrix allowed an evaluation of each of the shortlisted technologies in the context of specific end-user needs, and a final selection was made based on the outcome.

Two technologies were selected, validated in the laboratory and various performance criteria such as specificity, sensitivity, repeatability, and reproducibility were evaluated. The finalised methods were tested in the hands of end-users, with end-user training initially provided in a controlled environment and subsequently the end users conducted the tests themselves onsite and provided feedback.

A separate, but related piece of work was undertaken by UKHSA and 20/30 Labs to explore the possibility of repurposing of rapid, Covid-19 in-field wastewater diagnostic technology for the detection of foodborne pathogens. This work demonstrated that rapid detection of seven foodborne pathogens in wastewater is feasible using rapid Loop-mediated isothermal amplification (LAMP) based technologies.

This work has demonstrated the potential utility of using onsite diagnostics as inspection tools in the FBD arena, whilst also identifying that there are no "ready-made" solutions on the market that meets the specific needs of end-users but found that there are a wide variety of emerging technologies. It has therefore become apparent that guidance for the 'real world' deployment of onsite diagnostics, is essential. The focus of the 24/25 continuation work is to develop such a set of guidelines, in particular for the use of onsite diagnostics for official controls in the food sector. End user insights and perspectives will be sought, and a case study will be developed to illustrate application of the guidelines. Needs from the policy, legislative and accreditation body perspectives will also be considered (Fera).

Delivery partners

Delivery partners for this work:

- Fera Science
- University of Lincoln

Status

A horizon scanning exercise to look for and assess existing guidelines for the deployment of onsite diagnostic technologies has been undertaken and findings delivered to the programme.

End user engagement is also being planned, with a review of end users that have been involved in the project to date to be conducted and further appropriate end users identified and approached as needed. Work is underway to draft a structure for the guidelines and summarising key points for inclusion.

Key successes

Key successes for this work include:

- horizon scanning identified more than 65 potential onsite diagnostic technologies, which were subsequently assessed using a Technology Readiness Level (TRL) framework developed as part of PATH-SAFE
- two onsite diagnostic technologies successfully piloted by end users in onsite locations and key scenarios
- significant end-user engagement through the completion of an end-user study to understand community needs and to help in identifying sectors or processes that are well-studied, or not, for portable detection technologies

Outputs

All projects were presented at the PATH-SAFE Biosurveillance Conference on 28 and 29 February in London. [Recordings, slides and posters are available on the conference webpage.](#)

PATH-SAFE: Novel Foodborne Pathogen, and associated antimicrobial resistance, knowledge and surveillance approaches

Underpinning any surveillance system, are sample collection and analysis methodologies, baseline data to draw comparison to and data analysis tools.

Overview

The work within this theme has focused on three key areas:

- development of novel surveillance methodologies, approaches and tools
- generation of data on the genomic diversity of a range of foodborne pathogens, and associated AMR, across the four nations of the UK to establish baselines and address knowledge gaps
- exploration of whole genome sequence data to investigate foodborne pathogen and antimicrobial resistance transmission routes

Drawing on the strengths and expertise of a range of delivery partners, and the power of working collaboratively, it has been possible to deliver a large portfolio of projects that have incorporated perspectives and input from across the agri-food system.

Food Standards Scotland (FSS) have sequenced thousands of *Escherichia coli* (E.coli) isolates, from a diverse range of sources across Scotland, to help elucidate transmission dynamics through the agri-food chain and to develop machine learning source attribution models for predicting the source or host of origin for E.coli isolates. The team are now sequencing further agri-food related isolates to verify the application of the E. coli source attribution models, looking to demonstrate proof of concept for use of the model in the investigation of foodborne illness and risk management, and extend the approach to develop a similar model for *Salmonella* Typhimurium (STm).

The Animal and Plant Health Agency (APHA) and **Veterinary Medicines Directorate (VMD)**, have generated new data and knowledge on AMR presence, diversity, and transmission by applying genomics to existing surveillance sampling and initiating novel sampling in understudied compartments of the agri-food chain, particularly ruminants (via bulk milk, and cattle and sheep at slaughter) and animal feed. Whole genome sequence data produced by these projects is now

undergoing enhanced genomic to enhance understanding of AMR dynamics, transmission pathways and mobile genetic elements dissemination. Further novel sampling is being conducted to determine the prevalence of AMR bacteria in imported Raw Pet Food and to explore the application of novel abattoir environment sampling approaches, as a potential alternative surveillance method that could act as a proxy for existing, labour intensive, livestock sampling methods.

The Environment Agency (EA) have explored surveillance options for AMR in the environment including river water and bathing waters, wild flora and fauna, shellfish and bioaerosols, and developed a pilot environmental surveillance system This work is continuing as part of the Environment Agency research portfolio.

The Centre for Environment, Fisheries and Aquaculture Science (Cefas) undertook a river catchment pilot to compare wastewater, river water and shellfish for monitoring of a range of foodborne pathogens and have worked with UKHSA and the Public Analyst laboratories in Scotland to demonstrate the utility of wastewater for monitoring norovirus and salmonella at a national scale. The team is now continuing the application and advancements of methodologies developed in PATH-SAFE to demonstrate the utility of bivalve molluscan shellfish (BMS), a species routinely sampled through pre-existing surveillance frameworks, as a sentinel in a 'sample once test many' approach.

Bangor University are examining the flow, dynamics, and risk posed by AMR and FBP in wastewater discharges, particularly those originating from healthcare settings. Further work is ongoing to investigate seasonal patterns and potential wastewater treatment opportunities and barriers.

Food Standard Agency (FSA), UK Health Security Agency (UKHSA), NHS Lothian and Cefas are undertaking a project to generate FBD prevalence and WGS data from business-as-usual UKHSA polio wastewater monitoring, utilising methodologies developed by Cefas within their PATH-SAFE project. This data will subsequently be compared to data from clinical samples collected through [the third study of Infectious Intestinal Disease in the UK](#), presenting an exciting opportunity to align two large surveillance efforts to generate insights into the effectiveness of wastewater monitoring for FBP.

Queen's University Belfast (QUB) have been investigating the application of air and near-source wastewater monitoring for AMR and FBP in a care home setting in Northern Ireland. This work is continuing to explore seasonal trends, increasing understanding of AMR and foodborne pathogens prevalence and transmission and further evidencing the utility of the novel surveillance methods.

University of Oxford have sequenced thousands of Campylobacter isolates, from a diverse range of collaborators, to help elucidate source attribution and transmission dynamics through the agri-food chain, and the rise in AMR. Sequencing and analysis of further human and agri-food Campylobacter isolates is on ongoing, to determine levels of AMR in Campylobacter isolates currently circulating in the UK and enhance understanding of AMR gene transmission.

Agri-food and Biosciences Institute (AFBI) in Northern Ireland selected and sequenced archived isolates of Salmonella and Listeria monocytogenes from food and animal sources, to provide important historical context against which future outbreak sequences can be compared.

Delivery Partners

In addition to the lead delivery partners highlighted in the overview section above, a number of other collaborators have supported the delivery of these projects, including:

- Scottish Environment Protection Agency
- Public Health Scotland
- Scotland's Rural College
- Moredun Research Institute
- The University of Edinburgh
- Northern Ireland Environment Agency
- Department of Health (NI)
- Agriculture, Environment and Rural Affairs (NI)
- Department of Infrastructure (NI)
- HSC Public Health Agency
- Northern Ireland Water
- Welsh Government
- Public Health Wales
- Welsh Water
- Cardiff University
- South West Water
- Ribble Rivers Trust
- Biosphere North Devon
- Virosafety
- Resistomap
- Quadram Institute
- University of Exeter
- Cranfield University
- UK Pet Food
- Agricultural Industries Confederation
- National Milk Laboratories
- Arla
- UK Centre for Ecology and Hydrology
- Aecom
- Arup
- Capita
- Deloitte
- MicrobesNG

Status

Project teams across all projects have been very busy making the necessary arrangements to continue the projects for an additional year. This has included establishing agreements and contracts, developing and refining project brief (including defining milestones and deliverables), engaging with key internal and external stakeholders, establishing extended / new sampling plans and analysis workflows. Several projects have initiated sample collection and data analysis, with others due to start in the coming weeks in line with their project plans.

Key successes

Across the theme:

- more than 7,000 samples have been collected, across a range of sample sources and four nations
- additional analysis has been undertaken on over 2,300 samples collected as part of existing surveillance activities
- exploration of more than 3,500 archived samples and 4700 archived pathogen isolates
- in excess of 18,000 pathogen isolates have been sequenced

- increased understanding of methods suitable for environmental monitoring across a range of different media and development of a pilot environmental surveillance system

Novel surveillance methods, approaches and tools developed and tested using real world samples and data for the first time, including:

- protocols for isolation and testing of foodborne pathogens from river water, wastewater and shellfish
- wastewater-based AMR and FBD surveillance sample collection schedules, sample analysis and data analysis pipelines
- machine learning models to determine source/host attribution of pathogens

Contamination and transmission models:

- demonstrable benefit of wastewater-based surveillance inspiring nursing homes in Northern Ireland to consider wastewater systems design in future builds to support wastewater-based surveillance
- large sample volume, broad scope and high throughput nature of PATH-SAFE projects have accelerated development and refinement of existing and established methods, including long read sequencing methodologies used by APHA for routine AMR surveillance
- utility of routine whole genome sequencing demonstrated through patterns, trends and insight gained across the programme, including the identification of resistant bacterial isolates of concern within the isolates sequenced for PATH-SAFE
- strengthened cross government and academia collaborative approach to identifying and pursuing surveillance opportunities, methods development and results analysis
- industry actively involved in the design and conduct of government surveillance activities, and promoting benefits of such collaborations, including milk producers and an animal feed mill, in collaboration with National Milk Record and The Agricultural Industries Confederation respectively

Outputs

All projects were presented at the PATH-SAFE Biosurveillance Conference on 28 and 29 February in London. [Recordings, slides and posters are available on the conference webpage.](#)

Environment Agency Publications

[Scoping review into environmental selection for antifungal resistance and testing methodology.](#) Published on 22 July 2022 on gov.uk, this report reviews the current understanding of the mechanism for selection for antifungal resistance in fungal species following exposure to antifungals.

[Antimicrobial resistance surveillance pilot site selection and database extension.](#) Published on 22 July 2022 on gov.uk, this project developed selection criteria to identify suitable river catchments for piloting a surveillance programme for environmental antimicrobial resistance.

[Sampling strategy and assessment options for environmental antimicrobial resistance in airborne microorganisms.](#) Published on 22 July 2022 on gov.uk, this report reviews the available sampling options for antimicrobial resistant microorganisms, including their antimicrobial resistance genes, from the atmosphere.

[Antifungal medicines in the terrestrial environment: Levels in biosolids from England and Wales.](#) Published on 02 February 2023 in Science of The Total Environment, this article reports on biosolids from ten sites in England and one in Wales that were sampled and analysed for clinical

antifungals.

[Environmental surveillance of AMR, perspective from a national environmental regulator in 2023.](#) Published 30 March 2023 in Eurosurveillance, this is a perspective paper discussing some of the issues and choices that arise in selecting the most appropriate methods for surveillance for AMR in the environment from the perspective of a government agency that conducts routine environmental monitoring for other health related purposes.

[Environmental antimicrobial resistance: A review of biological methods.](#) Published 26 October 2023 on gov.uk, this report reviews biological methods for the detection of environmental antimicrobial resistance.

[Antimicrobial resistance surveillance strategies within wild flora and fauna of England.](#) Published 26 October 2023 on gov.uk, this report reviews potential antimicrobial resistance surveillance strategies for wild flora and fauna.

[Shellfish as bioindicator for coastal antimicrobial resistance.](#) Published 26 October 2023 on gov.uk, this report covers development of testing approaches to assess the presence and identity of antimicrobials and resistant microorganisms in marine shellfish.

[A review of approaches to monitoring and surveillance of antimicrobial resistance in bathing waters.](#) Published 26 October 2023 on gov.uk, this report improves understanding of antimicrobial resistance in relation to human exposures via bathing waters.

[Antimicrobial resistance in bioaerosols: towards a national surveillance strategy.](#) Published 26 October 2023 on gov.uk, this report considers the decisions that must be made in designing a national surveillance strategy for antimicrobial resistance in bioaerosols.

Cefas Publications

[Development and validation of a duplex RT-qPCR assay for norovirus quantification in wastewater samples.](#) Published November 2023 this article covers development of a duplex RT-qPCR assay that [enables the sensitive detection of both NoVGI and NoVGII](#) in wastewater-derived RNA eluents, in a time and cost-effective way and may be used for surveillance to monitor public and [environmental health](#).

[Long Amplicon Nanopore Sequencing for Dual-Typing RdRp and VP1 Genes of Norovirus Genogroups I and II in Wastewater.](#) Published on 27 March 2024 on bioRxiv, this article covers the development and optimisation of a long amplicon nanopore-based method for dual-typing the [RNA-dependent RNA polymerase \(RdRp\)](#) and major structural protein (VP1) regions of norovirus, with wastewater as the sample matrix.

[Realising a global One Health disease surveillance approach: insights from wastewater and beyond.](#) Published 22 June 2024 in Nature Communications, this article presents case studies centred around the recent global approach to tackle antimicrobial resistance and the current interest in wastewater testing, with the concept of “one sample many analyses” to be further explored as the most appropriate means of initiating this endeavour.

[Piloting wastewater-based surveillance of norovirus in England.](#) Published 1 October 2024 in Water Research, this study took approximately 3,200 samples of wastewater from across England, previously collected for quantification of SARS-CoV-2, and re-analysed them for the quantification of norovirus genogroup I (GI) and II (GII). [Comparisons of national average norovirus concentrations in wastewater against concomitant norovirus reported case numbers showed a significant linear relationship.](#)

Methodology publications

Cefas: [Long Amplicon Nanopore Sequencing for Dual-Typing RdRp and VP1 Genes of Norovirus Genogroups I and II in Wastewater V.2](#)

Data

FSS: [Scottish E.coli](#) (locate sequencing by searching the 'Comment' field for 'PATH-SAFE' and 'PATH-SAFE project')

Cefas: [Norovirus, ammonia and orthophosphate in wastewater samples collected from across England 2021 - 2022](#)

Cefas: [Whole Genome Sequencing Salmonella in Scottish Wastewater](#)

University of Oxford: [Campylobacter isolate collection on PubMLST](#)

AFBI: [Listeria Sequences](#)

AFBI: [Salmonella sequences](#) (locate sequencing by searching the 'Comment' field for 'PATH-SAFE' and 'PATH-SAFE_23_24')

PATH-SAFE: Programme Evaluation

An independent programme evaluation has been conducted, alongside the programme conduct.

Overview

The evaluation aims to identify areas for learning and improvement and the outcomes of PATH-SAFE that improve surveillance infrastructure.

There are three main objectives of the evaluation:

- design and framing: to design a programme ToC for the programme, articulating the change that is intended to be achieved by PATH-SAFE - this includes outlining assumptions, external factors and managing risks associated with the programme
- formative process evaluation: to utilise bespoke evaluation methods for assessing the processes underpinning the programme and delivering the outputs of the four Ws to identify areas for learning and improvement
- understanding impact: to identify and prioritise outcome indicators and data sources to validate the outcomes of the ToC and develop an assessment of PATH-SAFE's contribution

Delivery Partner

RAND Europe

Status

The phase 1 evaluation report is being finalised and is due to be published early Autumn.

In tandem, work on phase 2 of the evaluation has begun. The first step is to update the outcome measurement framework to reflect the renewed focus of phase 2 of the PATH-SAFE programme. A workshop will be held with the programme team and partners to discuss and agree the revised anticipated outcomes and impacts.

Outputs

Evaluation Framework Report - [Evaluation of the PATH-SAFE programme](#).

All projects were presented at the PATH-SAFE Biosurveillance Conference on 28 and 29 February in London. [Recordings, slides and posters are available on the conference webpage.](#)

PATH-SAFE: Events and communications

Throughout the programme, a range of events and communication activities have been undertaken.

Newsletters

For more information and updates on the PATH-SAFE programme, read the latest PATH-SAFE newsletter:

- [PATH-SAFE newsletter September 2024](#)
- [PATH-SAFE newsletter July 2024](#)
- [PATH-SAFE newsletter March 2024](#)
- [PATH-SAFE newsletter December 2023](#)
- [PATH-SAFE newsletter September 2023](#)
- [PATH-SAFE newsletter June 2023](#)
- [PATH-SAFE newsletter March 2023](#)

PDF

[Gweld PATH-SAFE newsletter, January 2023 as PDF\(Open in a new window\)](#) (273.2 KB)

PATH-SAFE Biosurveillance Conference

On Wednesday 28 and Thursday 29 February 2024, [a two-day PATH-SAFE Biosurveillance Conference was held in London](#). The aim of the conference was to facilitate knowledge exchange within the biosurveillance community, by showcase the biosurveillance work that has been undertaken within the PATH-SAFE programme initiatives, whilst and also creating an opportunity for people to connect and expand their professional networks.

You can access further information, including the presentation recordings, slides and posters on [the PATH-SAFE Biosurveillance Conference page](#).

Innovation in Biosurveillance Event

In November 2023, PATH-SAFE instigated and hosted a biosurveillance event that brought together over 100 key biosurveillance stakeholders from across government, academia and industry. The event actively promoted information sharing. Within facilitated workshops sessions systematic barriers were discussed, potential solutions identified, biosurveillance initiatives mapped and future opportunities explored.

Webinars

PATH-SAFE ran a series of webinar between March 2023 and January 2024 which to showcase results from the programme, as well topics relevant to programme partners.

15 March 2023 - Understanding source attribution, infection threat and level of AMR of E. coli in Scotland using whole genome sequencing

With Dr Adriana Vallejo-Trujillo (Research Fellow, Food Standards Scotland).

Please note the following resources are not accessible. If you require an accessible alternative please contact fsa.communications@food.gov.uk.

Slides available on request.

27 April 2023 - Genomic epidemiology of foodborne pathogens and antimicrobial resistance

A talk from Professor Alison Mather.

Unfortunately, due to the sensitivity of content, we are unable to share a recording or slides.

18 May 2023 - PATH-SAFE and The Food Safety Research Network Showcase

Due to a technical issue it was not possible to record this webinar. Slides available on request.

15 June 2023 - Rapid diagnostic technologies for Foodborne Pathogens

Please note the following resources are not accessible. If you require an accessible alternative, please contact fsa.communications@food.gov.uk.

Slides available on request.

13 July 2023 - First steps towards an environmental surveillance system for Antimicrobial Resistance (AMR) in England

Unfortunately, due to the sensitivity of content, we are unable to share a recording or slides.

25 September 2023 - AMR Surveillance Pilots: Dairy Cattle and Animal Feed Surveys webinar slides

Slides available on request.

17 October 2023 - PATH-SAFE Process Evaluation

Please note the following resources are not accessible. If you require an accessible alternative, please contactfsa.communications@food.gov.uk.

Slides are available on request.

9 November 2023 - Genomics of Animal and Plant Health Disease Centre II (GAP-DC2)

Unfortunately, due to the sensitivity of content, we are unable to share a recording or slides.

17 January 2024 - Genomics of antimicrobial resistant Campylobacter transmission through UK Agri-Food systems

Unfortunately, due to the sensitivity of content, we are unable to share a recording or slides.

[Yn ôl i'r brig](#)