

JOINT INTEGRATIVE PROJECTS (JIP)	INTEGRATIVE STRATEGIC ACTIVITIES	JOINT RESEARCH PROJECTS (JRP)		
		FOODBORNE ZONOOSES	ANTIMICROBIAL RESISTANCE (AMR)	EMERGING THREATS
<a href="#">MATRIX</a> : solutions to support and advance One Health surveillance	Design and implementation of surveillance activities	<a href="#">AIR-SAMPLE</a> : air filters to detect <i>Campylobacter</i> in broiler houses		
<a href="#">COHESIVE</a> : pathway analysis of detection of outbreaks		<a href="#">NOVA</a> : code to model disease spread and explore disease surveillance options		
<a href="#">OH-HARMONY-CAP</a> : diagnostics, laboratories capabilities, capacities and interoperability collection tool	Laboratory methods	<a href="#">METASTAVA</a> : <a href="#">Guidelines</a> for sequence based metagenomics disease surveillance	<a href="#">IMPART</a> : updated and improved detection protocols multicentre evaluation study <a href="#">results</a> . New <a href="#">ECOFEs</a> of veterinary antibiotics	<a href="#">Tox-Detect</a> : database of protein profiles of foodborne toxogenic bacteria
		<a href="#">TOXOSOURCES</a> : <a href="#">Harmonized Methods</a> for detecting <i>Toxoplasma gondii</i> contamination in fresh produce	<a href="#">FARMED</a> : <a href="#">Detection and Characterisation</a> of unauthorised genetically modified microorganisms	<a href="#">MAD-Vir</a> : <a href="#">Tool</a> to detect known viruses and discover new viruses
			<a href="#">WORLDCOM</a> : predict/detect AMR from microbial samples and genomic sequences	<a href="#">TELE-Vir</a> : portable toolbox for identification and characterisation of emerging virus threats
			<a href="#">MedVetKlebs</a> : <a href="#">The ZKIR Assay</a> , a Real-Time PCR method for the detection of <i>Klebsiella pneumoniae</i> in environmental samples	<a href="#">IDEMBRU</a> : toolbox for rapid detection identification of emerging <i>Brucella</i> species
				<a href="#">MEEmE</a> : detection tools standardisation and data collection tools on <i>Echinococcus multilocularis/granulosus</i> in the food chain
				<a href="#">PARADISE</a> : novel genotyping schemes and detection strategies for <i>Cryptosporidium</i> and <i>Giardia</i> detection
<a href="#">CARE</a> : <a href="#">database of strains and genomes</a> for quality control analysis in food safety	Reference material and data	<a href="#">LISTADAPT</a> : <a href="#">Algorithm</a> for selecting strains to explore the diversity of strains circulating	<a href="#">ARDIG</a> : collection of large number of genomes that can be used as reference material for AMR confirmation	
<a href="#">ORION</a> : framework for understanding and information exchange - <a href="#">One Health Surveillance Codex</a>	Interpretation of surveillance data	<a href="#">ADONIS</a> : decision making tool to determine causes and best interventions in human <i>S. Enteritidis</i> infections	<a href="#">ARDIG</a> : <a href="#">Comparability between antimicrobial usage and AMR data to improve AMR surveillance</a>	
<a href="#">COVRIN</a> : models for risk assessment of SARS-CoV-2		<a href="#">BeONE</a> : integrative solutions for foodborne pathogens surveillance	<a href="#">FULL-FORCE</a> : data on plasmid structure and variability of drug-resistant organisms	
<a href="#">COHESIVE</a> : information system that stores genomics data and metadata of pathogens from different countries ( <a href="#">demo</a> )		<a href="#">DISCoVeR</a> : models and methods for attributing human foodborne infections to animal, food and environment sources		
		<a href="#">TOXOSOURCES</a> : methods to evaluate the relative contribution of different sources of <i>Toxoplasma gondii</i> infections		
		<a href="#">MedVetKlebs</a> : <a href="#">Multicentric Study</a> of <i>Klebsiella pneumoniae</i> in European food products		
<a href="#">COHESIVE</a> : <a href="#">Risk Analysis System</a> for zoonoses; <a href="#">FoodChain-Lab</a> web application to trace suspicious food items; quantitative <a href="#">shiny Risk</a> application assessment toolbox; risk assessment <a href="#">Decision Support Tool</a>	Cross-sector communication of data	<a href="#">BIOPIGEE</a> : education and training activities	<a href="#">FULL-FORCE</a> : tool box for Single Molecule Real Time sequencing for AMR surveillance	
		<a href="#">NOVA</a> : mathematic models for data combination and analysis for One Health syndromic surveillance systems	<a href="#">FED-AMR</a> : new data on the role of extracellular DNA as an AMR source and on AMR spread in agricultural environment	
<a href="#">ORION</a> : solutions for interoperability to improve data FAIRness - <a href="#">OHEJP Glossary</a> , <a href="#">One Health Linked Data Toolbox</a> , <a href="#">Health Surveillance Ontology</a>			<a href="#">RaDAR</a> : modelling methodology for AMR specific source attribution, disease burden	
<a href="#">COHESIVE</a> : review on economic <a href="#">Economic Analysis of Foodborne Zoonoses</a>	Action (prevention and response)	<a href="#">MoMIR-PPC</a> : <a href="#">Prevention &amp; Control Measures</a> against <i>Salmonella</i> at the poultry production level		
		<a href="#">BIOPIGEE</a> : biosecurity measures for the control of <i>Salmonella</i> and HEV in primary pig production and abattoir		