

# Eagle Genomics

## Microbiome Insights: Pioneering the Future of Precision Livestock Farming

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14<sup>th</sup> September 2023  
Animal Microbiome Webinar

[ Networking Microbiome Science ]

eagle genomics

# My journey



## Cancer

## Food & Nutrition

## Microbiome

UEA University of East Anglia



Exploring the relationship between food and health

Delivering healthier lives through innovation in gut health, microbiology and food

Mastering the microbiome, transforming lives®

Unlocking the microbiome

Microbiome network science

Scientist

Head of Business Development

Head of Business Development

Chief Business Officer

Director, Strategic Alliances

Director, Microbiome Solutions



# About Eagle Genomics



## Deep Tech scale-up business in the TechBio space

- Driving the Bioeconomy through the digitization of science



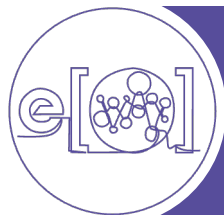
## Applying Microbiome Network Science

- Changing the face of One Health across various players in the ecosystem



## Driving sustainable systems change

- Promoting and protecting the health of humans, animals and the environment



## e[datascientist]™

- Award-winning AI-augmented knowledge discovery platform
- Leverages AI and network science, surfacing scientific connections and exploring multi-causal relationships

# Challenges for future food production

## Growing demand

- World population growth: 10 billions in 2050
- Increasing demand from developing countries

## Limited resources

- >25% of food wasted in production, processing and consumption
- Water and arable land are becoming scarce resources

## Lack of consumer trust

- Missing transparency in food production and processing
- Insufficient food safety in some region

## Lack of integration along the value chain

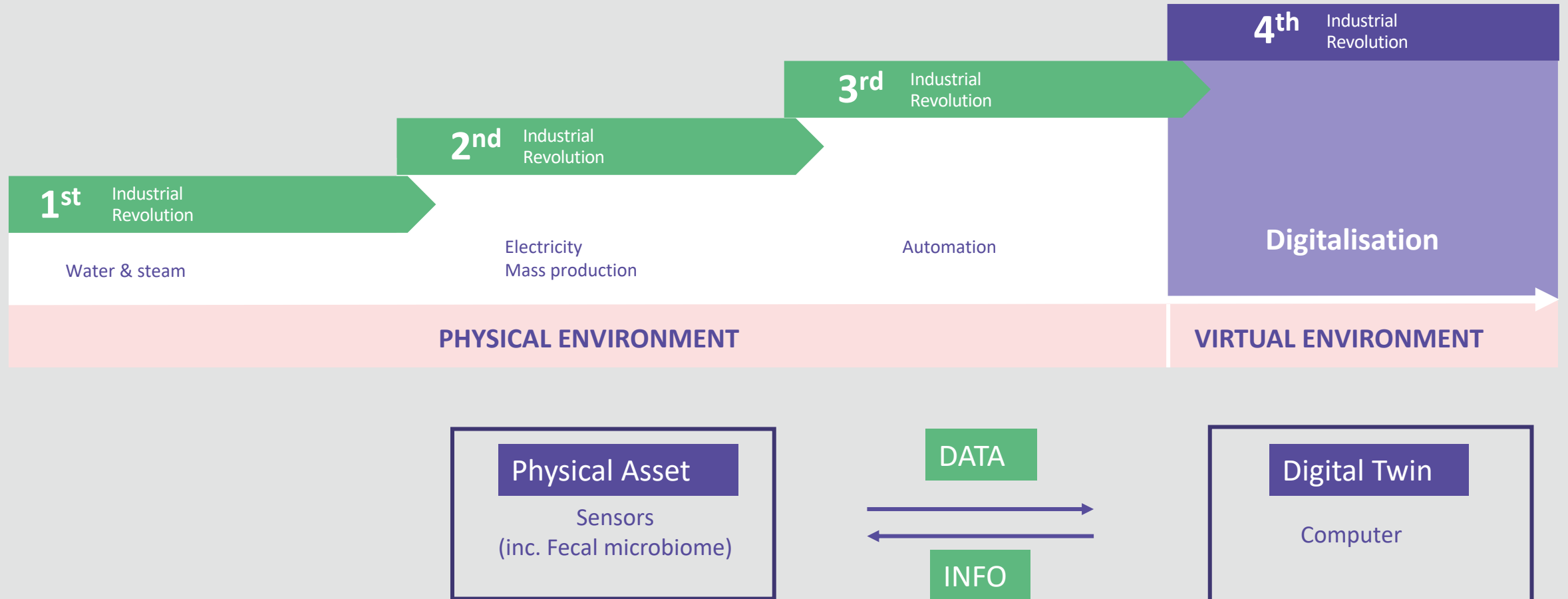
- Fragmented market

**Food  
production  
and  
processing  
must  
become  
more ...**

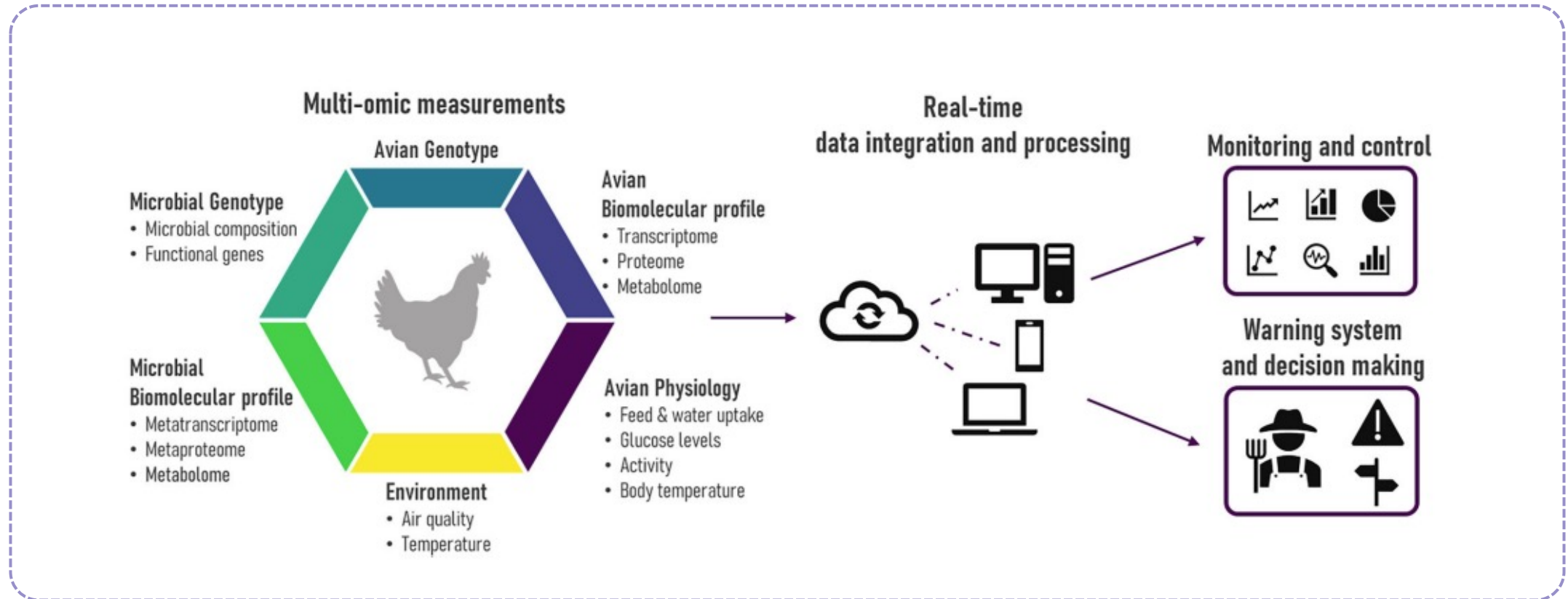
- Efficient
- Sustainable
- Transparent
- Personalised
- Integrated

# Animal Health & Welfare

## From analog to digital

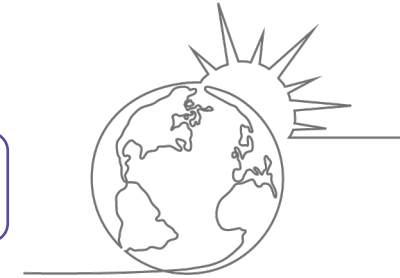


# Future integration of multi-omics measurements



Goossens E, Dehau T, Ducatelle R, Van Immerseel F. Omics technologies in poultry health and productivity - part 2: future applications in the poultry industry. *Avian Pathol.* 2022;51(5):418-423.

# Microbiota, Gut Health and Chicken Productivity



## HOST FACTORS

- Genetic background
  - Bird species
  - Type
  - Breed
  - Sex
- Gut development and maturation
  - Immune system
  - Gut morphology
  - Microbiota acquisition
  - Gut region



## GUT HEALTH

- Nutrient assimilation
- Intestinal barrier integrity
- Immune response efficiency
- Inflammatory balance
- Susceptibility to enteric pathogens (E. coli, Salmonella, Campylobacter, C. perfringens)

## ENVIRONMENTAL FACTORS

- Seasonal climate changes
- Geographical climate changes

## FARM MANAGEMENT

- Stress/Welfare
- Stocking density
  - Temperature
  - Photoperiod
  - Ventilation
  - Access to range

- Health interventions
- Antibiotic growth promoters
  - Therapeutic antibiotics
  - Vaccination
  - Probiotics / Prebiotics
  - Plant bioactives

- Litter management
- Litter components
  - Humidity
  - Reuse / Removal cycles

- Feed management
- Feed access
  - Water access
  - Feed changes
  - Feed quality

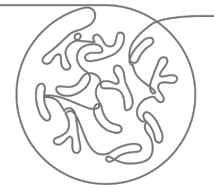
## NUTRITION

- Feed components
- Particle size
- Fiber source
- Protein source
- Micronutrients
- Enzymes

## LITTER MICROBIOTA



## GUT MICROBIOTA

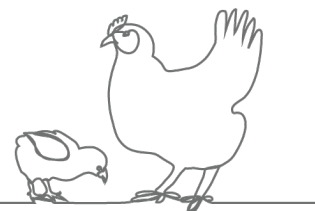


- Functional output
- Production of SCFA
  - Added metabolic potential
  - Competitive exclusion of pathogens

- Composition
- Richness of species
  - Population structure and complexity
  - Balance between commensal and detrimental microorganisms
  - Competitive exclusion of pathogens

## CHICKEN HEALTH & WELFARE

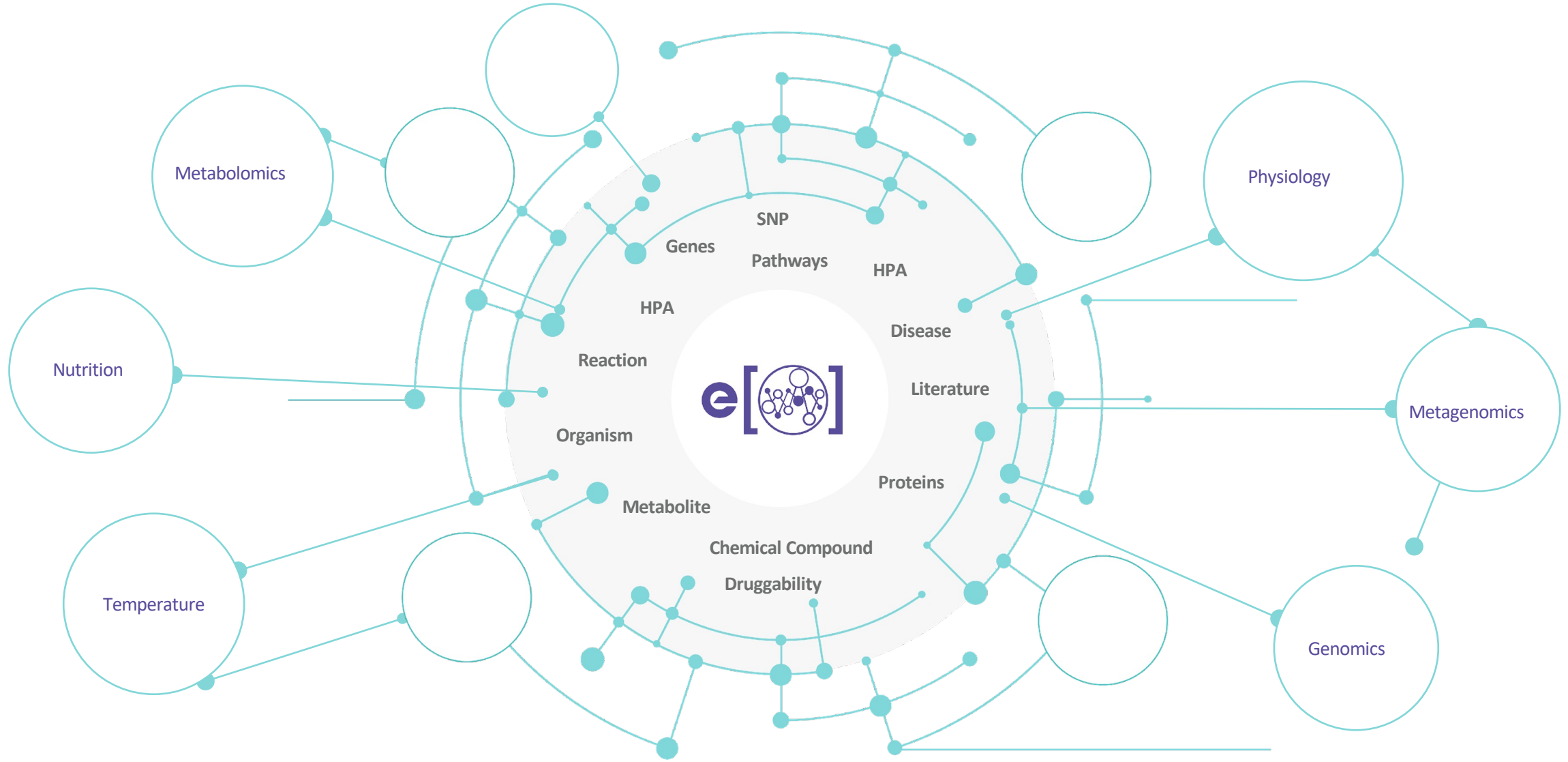
- Energy balance
- Feed efficiency
- Growth rate



PUBLIC

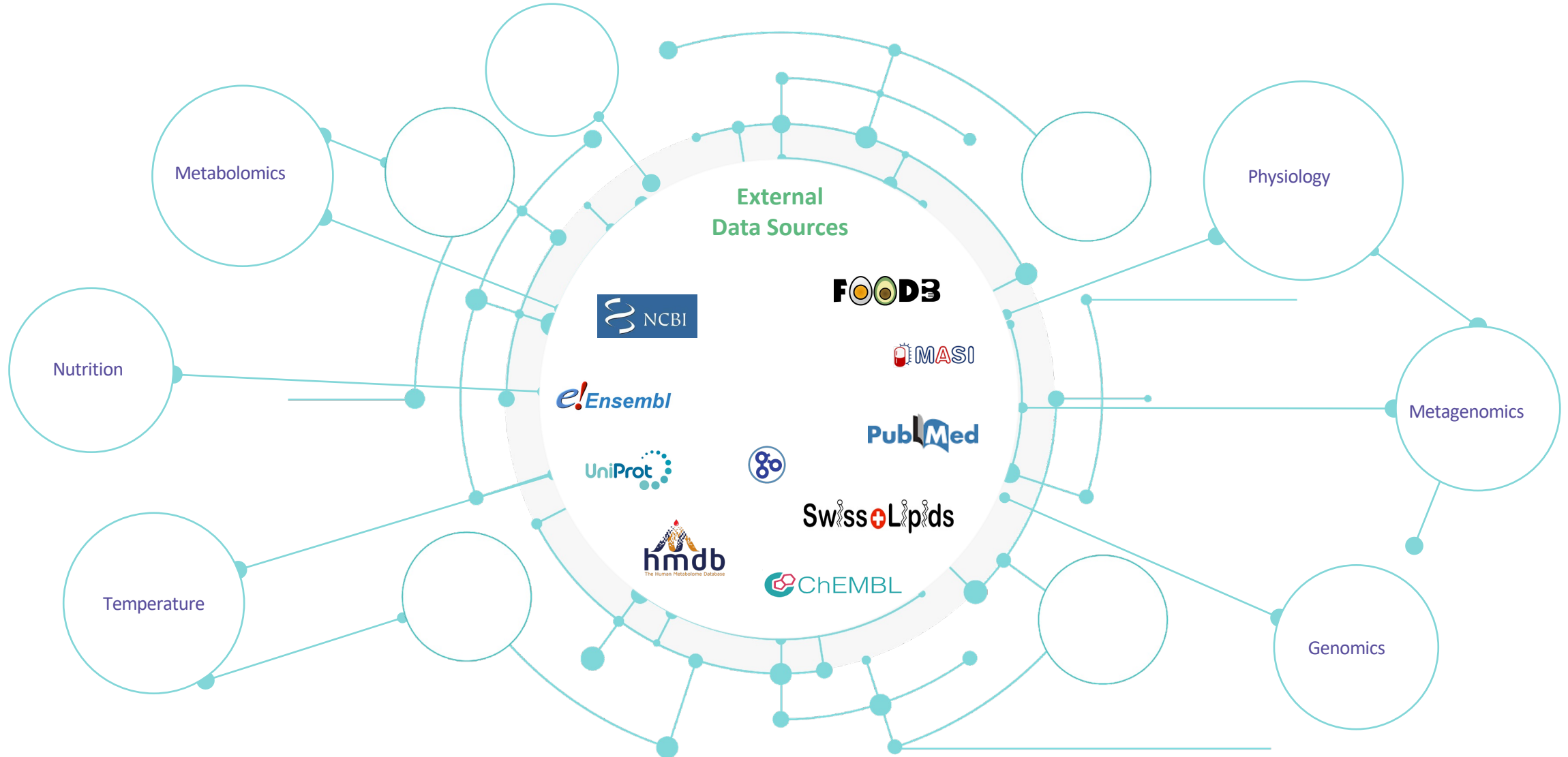


# Our approach: create a **data fabric** from siloed data...

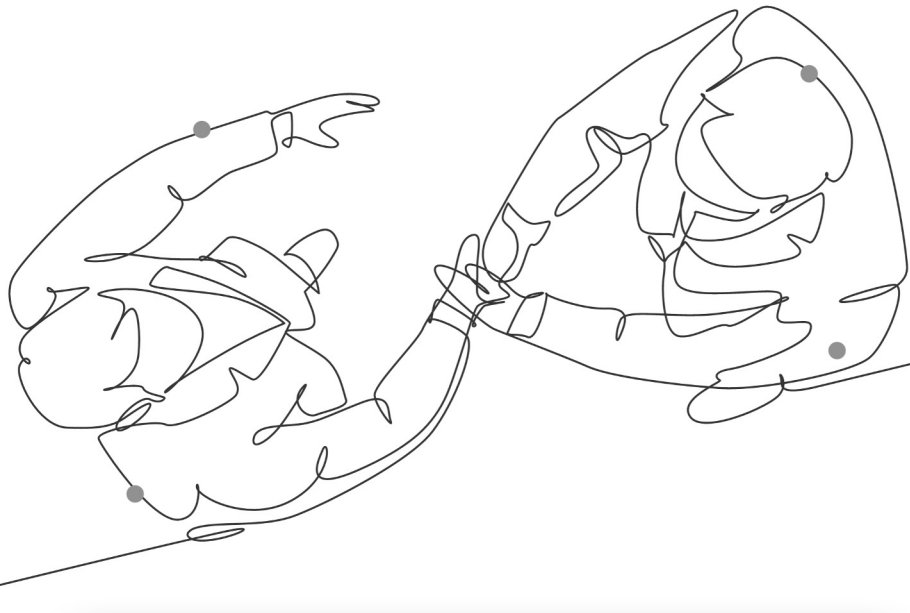




... and **enrich** it with additional data assets...



... including from our existing knowledge & technology partners



### Technology Partners



### Knowledge Partners



### Ecosystem Entry Partners



# Step 1: Onboarding measurement data and knowledge

## Data curation

The screenshot shows a 'Domain management' interface with a tree view on the left containing entities like Study details, Experiment, Subject, Timepoint, and Treatment. On the right, a 'Sample' entity is selected, showing a table of processed data with columns for EDGE ID, DISPLAY NAME, and Attributes.

EDGE ID	DISPLAY NAME	Attributes
EGG-A0001	1_DH_ActivatorConcentration	1 Attributes
EGG-A0002	1_DH_NaphtheneConcentration	1 Attributes
EGG-A0003	1_DH_PhenoanthreneConcentration	1 Attributes
EGG-A0004	11_AmyrinolConcentration	1 Attributes
EGG-A0005	2_DH_FluoreneConcentration	2 Attributes
EGG-A0006	2_DH_NaphtheneConcentration	2 Attributes
EGG-A0007	2_DH_PhenoanthreneConcentration	2 Attributes
EGG-A0008	3_DH_FluoreneConcentration	3 Attributes
EGG-A0009	3_DH_PhenoanthreneConcentration	3 Attributes
EGG-A0010	4_DH_PhenoanthreneConcentration	4 Attributes
EGG-A0011	9_DH_FluoreneConcentration	9 Attributes
EGG-A0012	9_DH_PhenoanthreneConcentration	9 Attributes
EGG-A0013	acetylcholinesteraseConcentration	1 Attributes
EGG-A0014	acetylcholinesteraseConcentration	1 Attributes
EGG-A0015	activeLesions	1 Attributes
EGG-A0016	adherenceScore	1 Attributes
EGG-A0018	anti-oxidantConcentration	1 Attributes
EGG-A0019	anthraceneConcentration	1 Attributes

## Study onboarding & management

The screenshot shows a 'Study Management' interface for a study titled 'Transcription profiling by array of rat vascular smooth muscle'. It features a 'Study Details' section and a 'METADATA UPLOAD' section with a grid of templates for various categories like Beauty & Personal Care, Oral, Skin, Scalp, Deo, Hair, Household & Laundry, Household, Laundry, Nutrition, Food, Beverages, and Ice Cream.

## Analyzed measurements e.g., signatures

The screenshot shows a 'Signature Analysis' interface with a table of gene signatures and a 'Signature Metadata' panel on the right.

Gene ID	Gene Name	base_Mean	logFC	l	lfcSE	pvalue
ENSG0000025040	HEBP	1153.871702	-0.61664701	0.20772247	2.86E-07	
ENSG0000024972	KHRL2	1538.219483	0.71389706	0.34717337	1.31E-06	
ENSG0000024757	ELF3	1875.806548	-0.4172724	0.19275421	1.16E-05	
ENSG0000019430	KCHD3	183.540154	-0.56134881	0.40341482	1.61E-05	
ENSG0000010649	KCHD3	187.4133885	-0.55899448	0.19781739	4.11E-05	
ENSG00000189324	SOX40	16772.30204	-0.33799479	0.16617662	4.48E-05	
ENSG00000130864	HEBP	1153.871702	-0.51793291	0.15581563	6.66E-05	
ENSG00000183444	KHRL2	166.3402151	-0.29261309	0.32893965	6.68E-05	
ENSG00000183444	KHRL2	166.3402151	-0.30317509	0.15438547	8.76E-05	
ENSG00000183444	KHRL2	166.3402151	-0.30710507	0.16927965	0.000305	
ENSG00000183444	KHRL2	166.3402151	-0.29915124	0.17549796	9.86E-05	
ENSG00000183444	KHRL2	166.3402151	-0.30261586	0.24882056	0.000371	
ENSG00000183444	KHRL2	166.3402151	-0.30129744	0.11541353	0.00017443	
ENSG00000183444	KHRL2	166.3402151	-0.26403242	0.22237974	0.00017443	
ENSG00000183444	KHRL2	166.3402151	-0.29058675	0.23947083	0.00017443	
ENSG00000183444	KHRL2	166.3402151	-0.29491519	0.17549796	9.86E-05	
ENSG00000183444	KHRL2	166.3402151	-0.30261586	0.24882056	0.000371	
ENSG00000183444	KHRL2	166.3402151	-0.30129744	0.11541353	0.00017443	
ENSG00000183444	KHRL2	166.3402151	-0.26403242	0.22237974	0.00017443	
ENSG00000183444	KHRL2	166.3402151	-0.29058675	0.23947083	0.00017443	

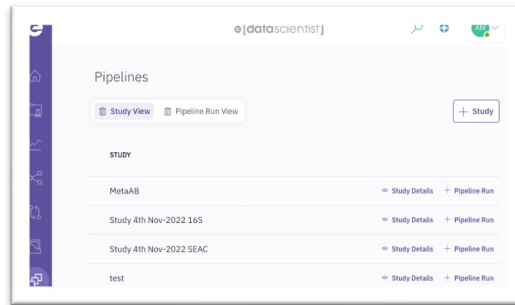
## Onboarding public & premium datasets

The screenshot shows an 'Onboarding public & premium datasets' interface with a list of datasets and a selection panel on the right.



# Step 2: Analyzing and classifying raw measurement data

## Selecting bioinformatics pipelines



## Configuring pipelines

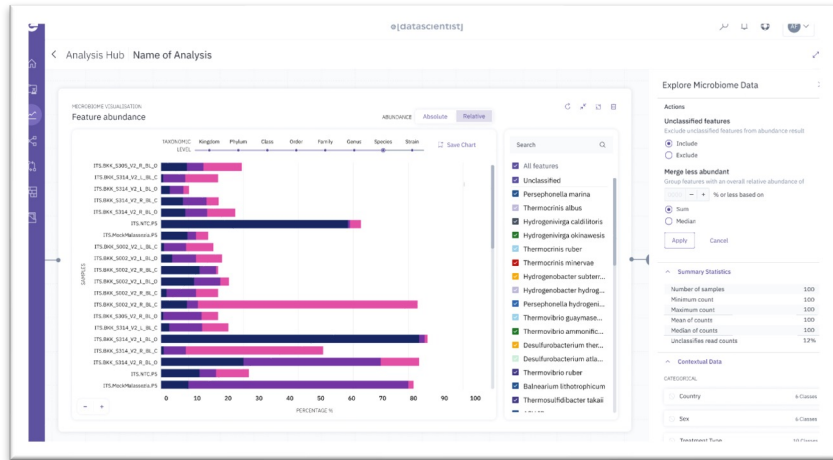


Name	Description
<b>Metagenomics (WGS) assembly based</b>	Processes metagenomic paired-end whole genome sequencing data to generate contig assemblies, gene predictions, taxonomic classification and functional annotation.
<b>Metagenomics (WGS) read based</b>	Processes metagenomic paired-end whole genome sequencing data and matches reads against references to generate taxonomic classification and functional annotation.
<b>Illumina 16S</b>	Processes 16S targeted sequencing data to generate prokaryotic taxonomic classifications and phylogeny.
<b>Illumina 18S</b>	Processes 18S targeted sequencing data to generate eukaryotic taxonomic classifications & phylogeny.
<b>ITS marker gene</b>	Processes ITS targeted sequencing data to generate fungal taxonomic classifications & phylogeny.
<b>Illumina QC visualisation</b>	Marker gene QC visualisation. Processes targeted sequencing data and performs quality assessment of the inputs.
<b>Single isolate / Clonal Assembly</b>	Processes single isolate genome sequences and generates complete whole genomes for individual bacterial strains. Provides characterisation including functional annotation, prediction of secondary metabolite gene clusters & antimicrobial resistance genes

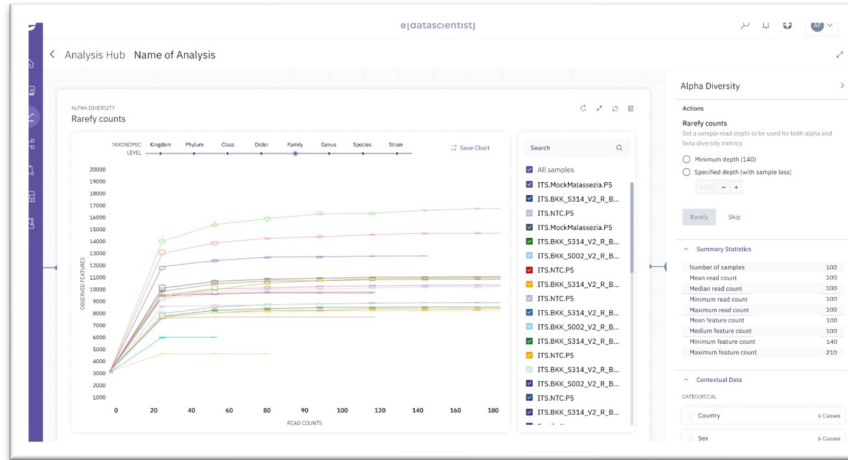


# Step 3: Carry out foundational microbiome analyses in Analysis Hub

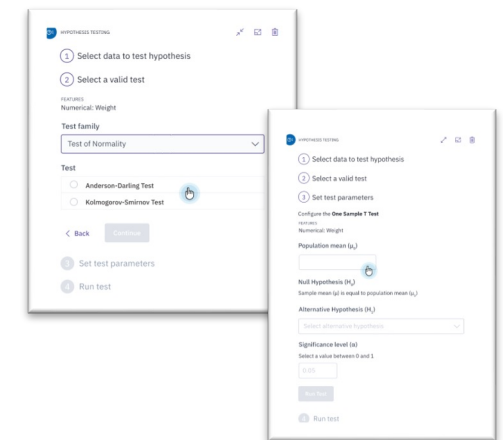
## Feature Abundance



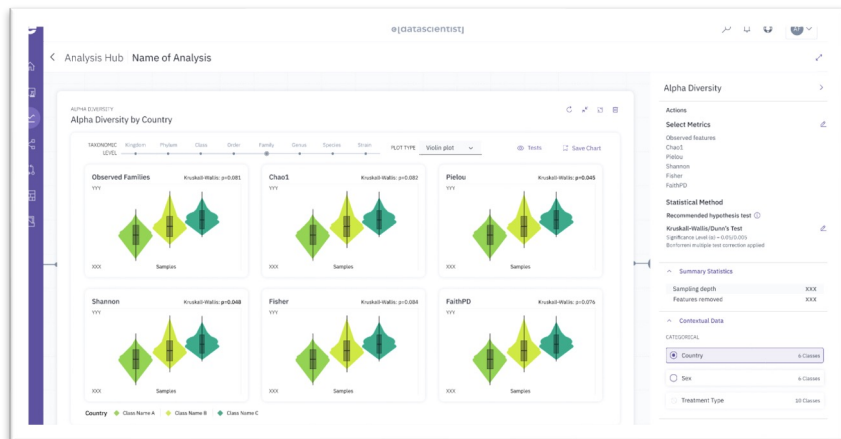
## Rarefaction



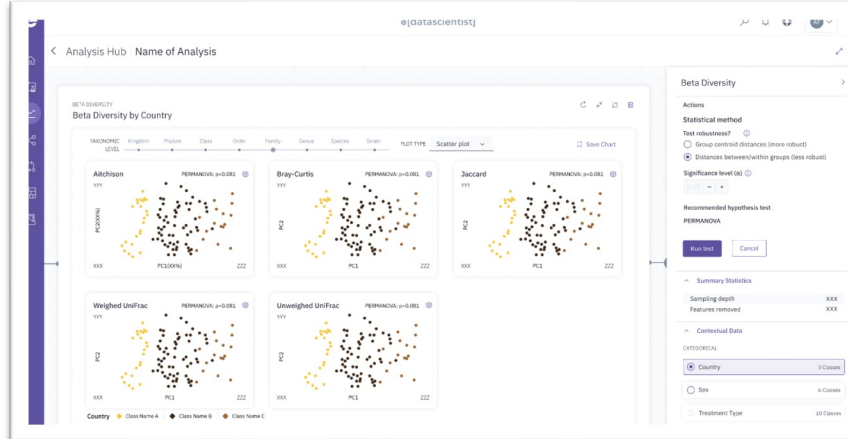
## Hypothesis Testing



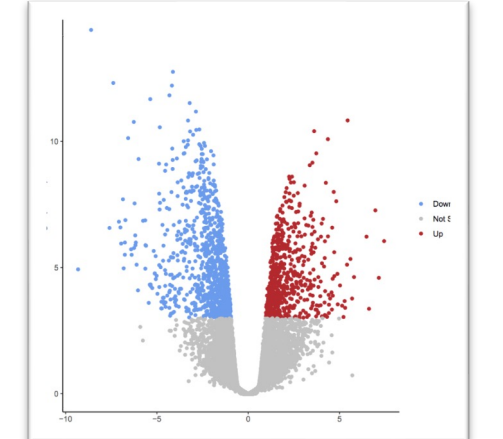
## Alpha Diversity



## Beta Diversity



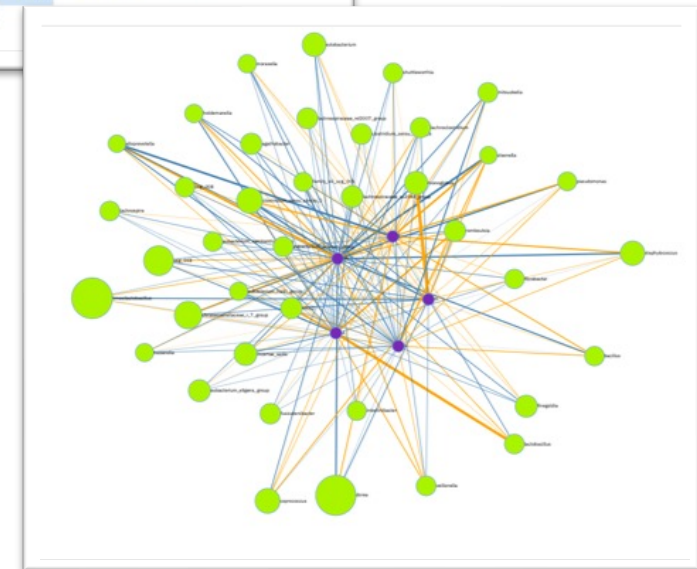
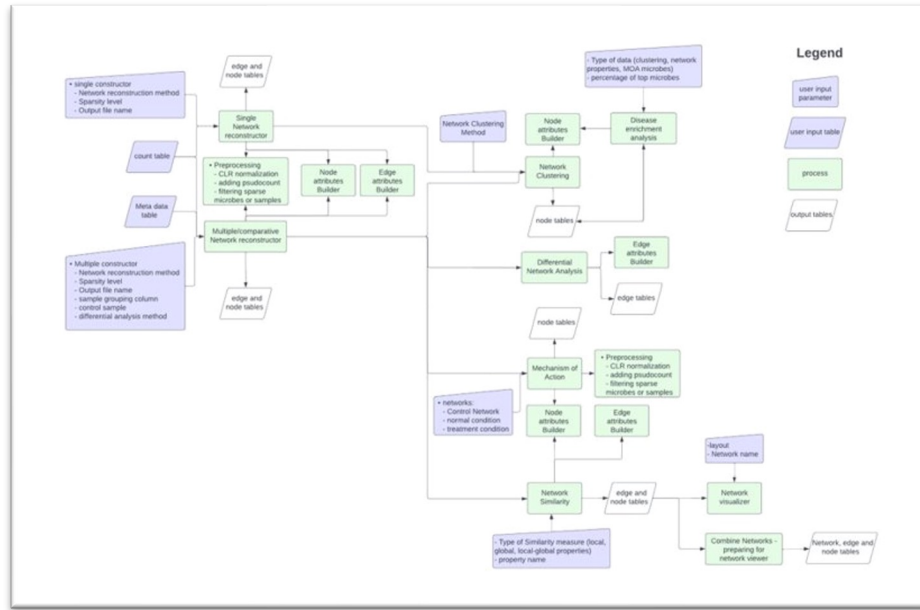
## Differential Abundance



# Step 4: Network based analysis of measurement data

## Network-based analysis pipelines

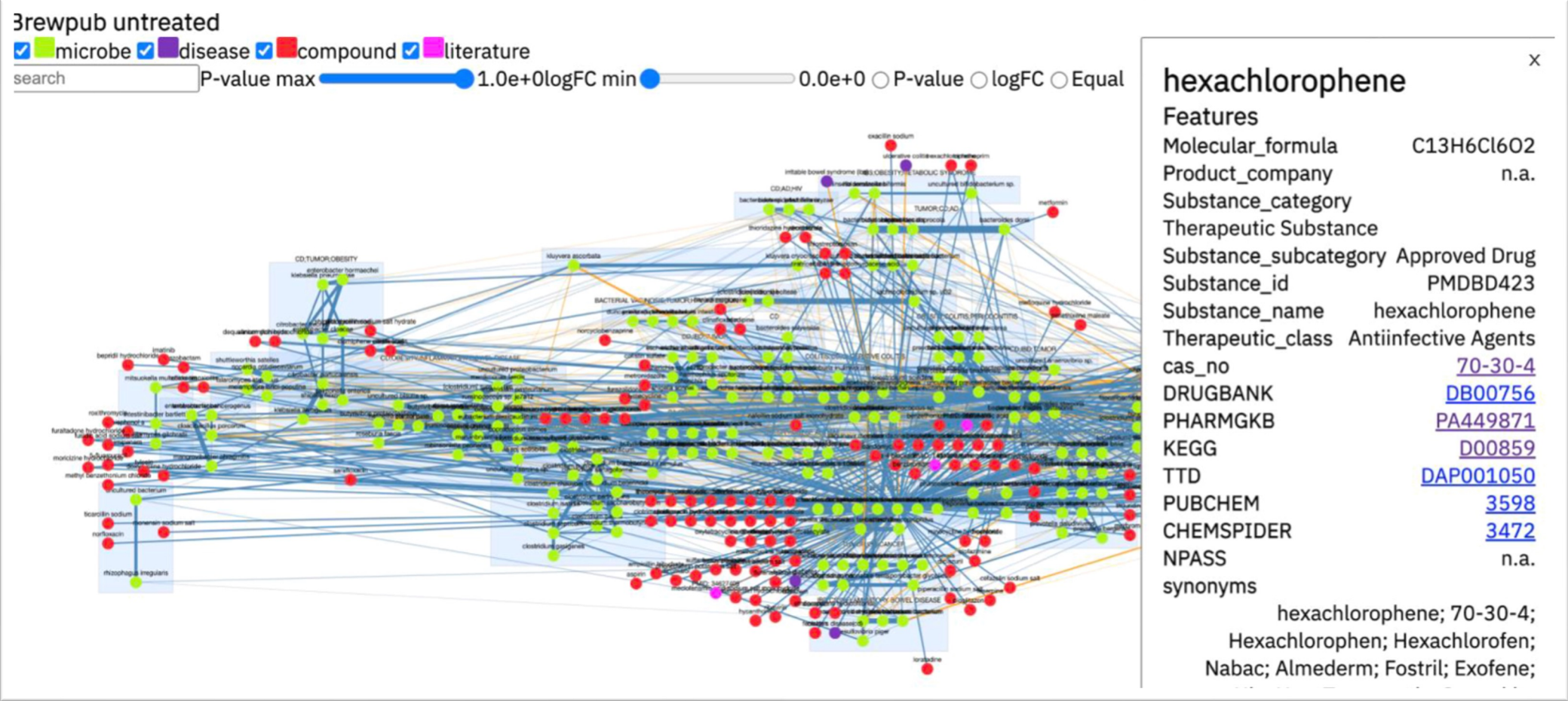
## Networked omics data exploration



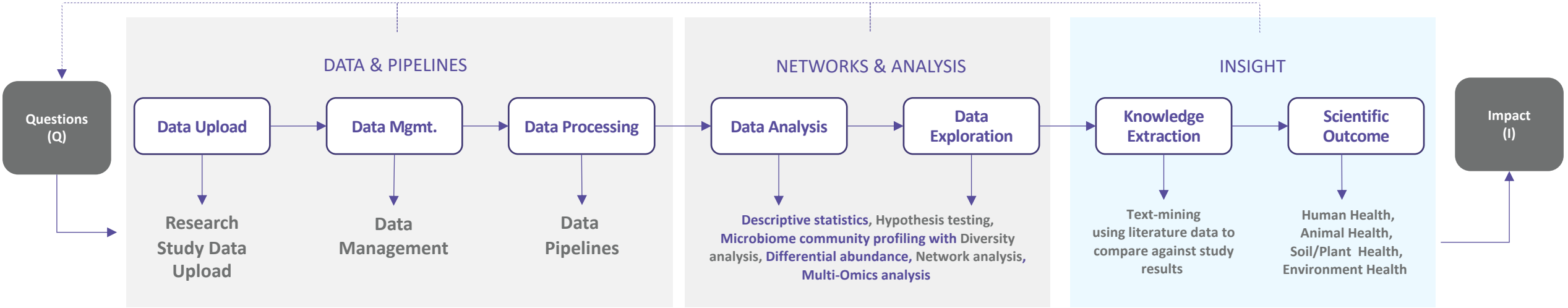
Identify hubs, biomarkers and controllers by exploring networked properties and functional clusters

# Step 5: Enriching networks with information from the Knowledge Graph

Connecting known links to diseases, KEGG compounds and literature references



# e[datascientist]<sup>TM</sup>: Workflows enabling insight generation and impact...

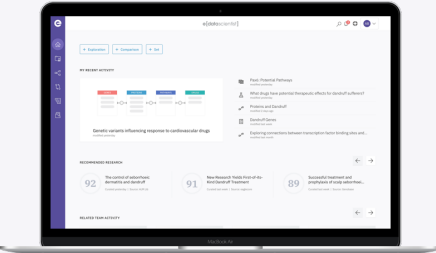


Semi-Automated Curation  
Common Data Model  
Bio-Data Processing Pipelines

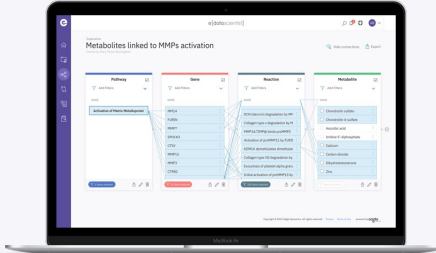
Multi-Layer Hypergraph,  
Network-, and Multi-Omics  
Analysis

Knowledge Extraction,  
Decision Support, Recommendations /  
Reporting

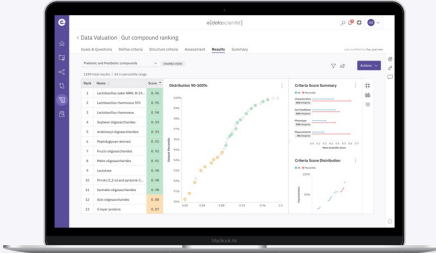
## e[datascientist]<sup>TM</sup> Applications



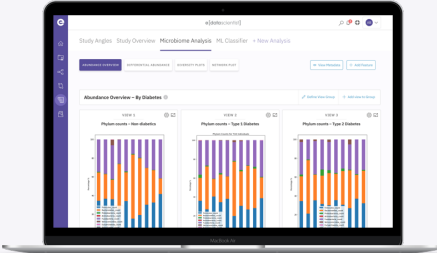
Conversational Interface



Data Exploration



Valuation & Decision Framework



Data Analysis

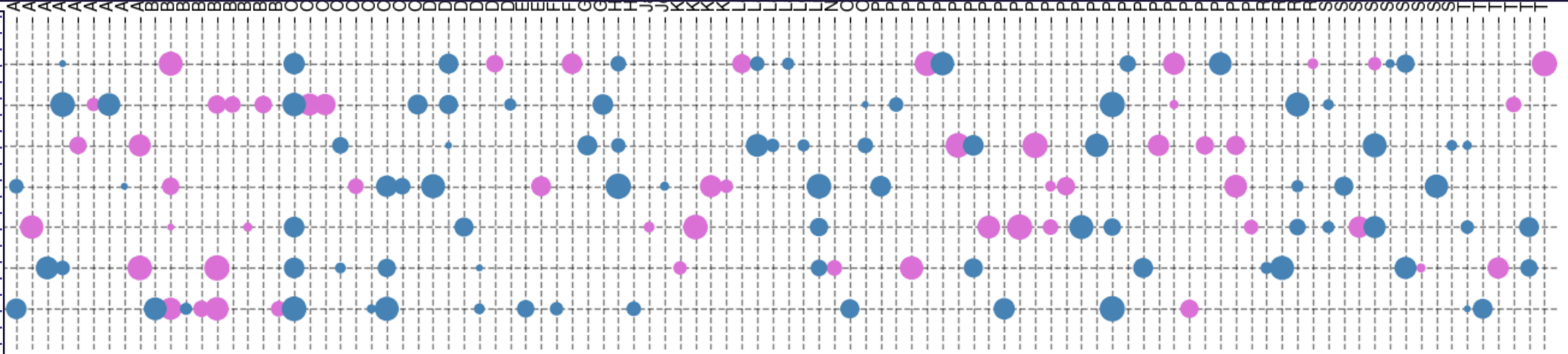




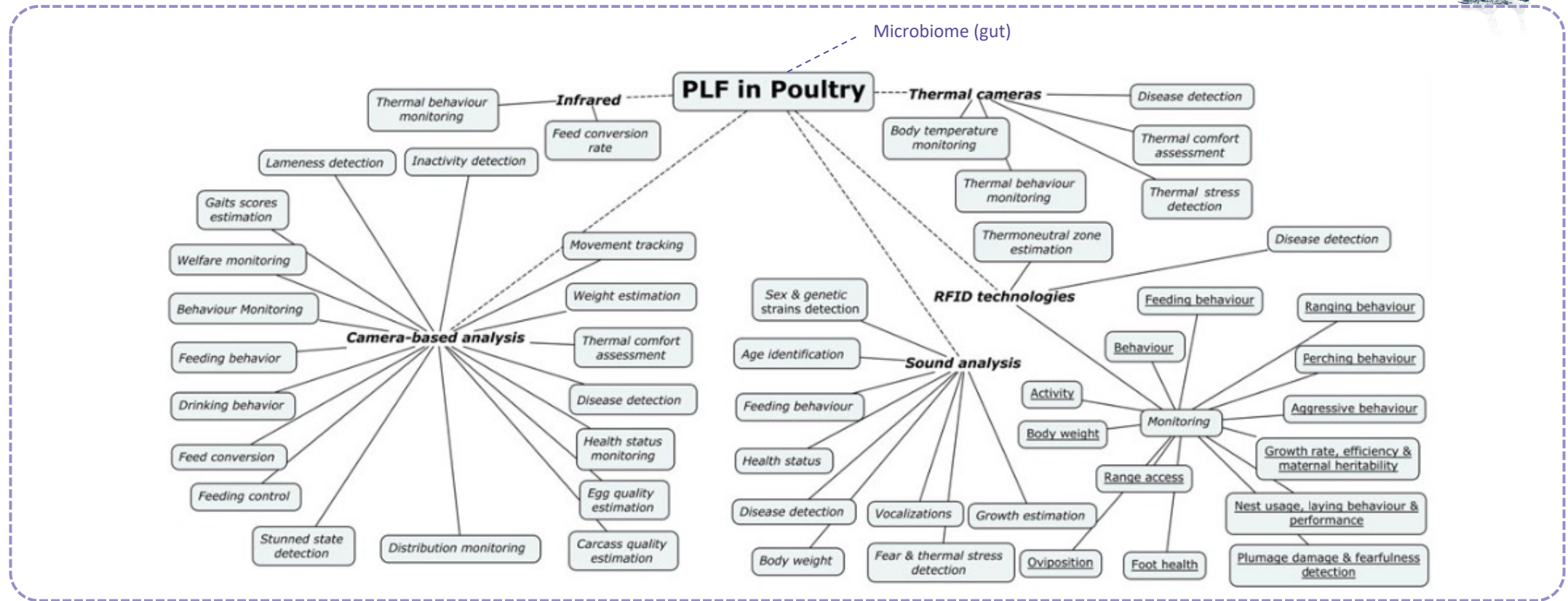
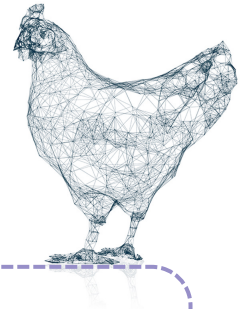
# Feature importance for 21-days old chickens

Redacted

Redacted

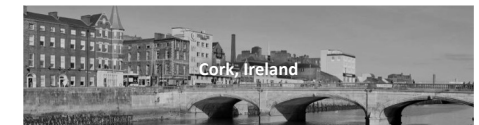
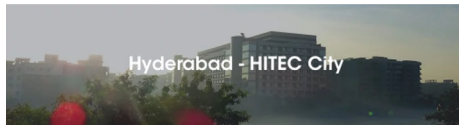
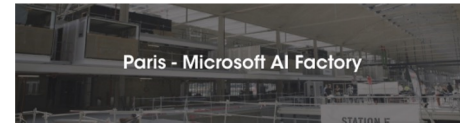


# The microbiome is the next frontier in PLF



Tzanidakis, C., Simitzis, P., Panagakis, P. (2023). Precision Livestock Farming (PLF) Systems: Improving Sustainability and Efficiency of Animal Production. In: García Márquez, F.P., Lev, B. (eds) Sustainability. International Series in Operations Research & Management Science, vol 333. Springer, Cham.

# Thank you



[www.EagleGenomics.com](http://www.EagleGenomics.com)



[Christian.Roghi@EagleGenomics.com](mailto:Christian.Roghi@EagleGenomics.com)



# Collaboration with **livestock**

- The health and wellness of livestock is critical to the economic success of the livestock industry
- There is no vehicle to address livestock health problems quickly.



## **Balanced Gut improves Livestock Health**

A balanced and diverse gut microbiome promotes efficient digestion, better growth, and overall health.



## **Optimize Livestock Performance**

A well-balanced livestock microbiome can improve production efficiency, allowing for body weight gain and increased production of eggs, milk, etc.



## **Efficiently Manage Disease**

Identifying the specific microorganisms that are associated with disease could help prevent or treat disease outbreaks in livestock.



## **Enhance Feed Efficiency**

Monitoring the effects of ingredients on gut health could help identify ways to enhance feed efficiency, leading to cost savings and improved sustainability.