

# Focus on the Joint Research Activities

- Improving measurement methods and ethics, developing new methods, enhancing phenotyping capacity
- Reducing experimental constraints on animals

WP5: *In vivo*  
“gold standard”  
measurements



WP6: Proxies and  
less-invasive  
methods



WP7: Sensor  
based  
phenotyping



## WP5

# Evaluation and standardization of nutrient use efficiency and emission measurement techniques

**Leader WP5.** Chris Reynolds (UREAD)

**Task 5.1.** Chris Reynolds (UREAD)

**Task 5.2.** Jan Dijkstra (WU)

**Involved partners** INRA, SRUC, WU, UREAD, FBN, AU



# WP5 Objectives

- To address **sources of variation in key in vivo measurements** of dietary nutrient use efficiency and associated emissions of methane and nitrogen by cattle (dairy and beef)
- To improve the accuracy and precision of measurements
- To unify the methods used across SmartCow RIs



## Task 5.1 Optimising measurements of **diet digestion and N balance** using *total collection of faeces and urine*

- 1/ **Historical databases (WP3) of digestion trials** and N balance analyzed for sources of variation due to:
  - Location (experimental error) and animal
- 2/ **New methods** of collection developed as needed
  - Non-invasive, automated collection of urine and faeces
- 3/ **Test** of optimum sampling period and procedures
  - Collections over 10 days

### Outputs:

- Optimised protocols for measuring digestion and N balance that minimise variation and measurement error, irrespective of animals (gender, physiological state) and diet types



## Task 5.2 Reducing uncertainty in methane (CH<sub>4</sub>) emission measurements using *Respiration Chambers*

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- 1/ **Historical databases** (WP3) of CH<sub>4</sub> emissions analyzed for sources of variation due to :
  - Location (experimental error) and animal
- 2/ **Ring test** of CH<sub>4</sub> recovery for chambers
  - Methane recovery as described by Gardiner et al. 2015
  - Identify sources of variation and options for improving recovery

### Outputs

- Optimised protocols for measuring CH<sub>4</sub> emissions that minimise variation and measurement error



## WP6

# Evaluating proxies to quantify feed efficiency and its determinants in cattle

**Leader WP6.** Cécile Martin (INRA)

**Task 6.1.** Gonzalo Cantalapiedra (INRA)

**Task 6.2.** Frédéric Dehareng (CRA-W)

**Involved partners** SRUC, UREAD, AU, CRA-W, INRA



# WP6 Objectives

- To evaluate **proxies\*** of feed efficiency and its determinants in cattle (dairy and beef)
- To identify their **range of applicability** across **diets** and **individuals**

\* define as indicators measurable frequently in body matrices easy to access; less invasive and costly, easier to implement



# WP6 Approach and Methodology

1/ Descriptive analysis of **existing [database & samples bank]** from SmartCow partners

- Identification of proxies for which new samples analysis are needed

2/ Collection of **new and relevant [data & samples]** of body matrices from animal experiments (**WP5, SmartCow RIs**)

- to complete the database (animals, diets)

3/ Assessment of potential and limits of each prediction model

- Validation of proxies with a high potential of prediction
- Exploration of new proxies (single or in combination)





## Task 6.1 Evaluation of proxies to predict digestibility, N partitioning and animal feed efficiency

Parameter	Proxies	Matrices	Status
<ul style="list-style-type: none"> <li>Total tract digestibility</li> <li>N partitioning</li> <li>Animal feed efficiency</li> </ul>	<ul style="list-style-type: none"> <li>NIR</li> <li>Urea-N concentration</li> <li>15N natural abundance</li> <li>15N natural abundance</li> <li>Urea-N</li> <li>Metabolites</li> </ul>	<ul style="list-style-type: none"> <li>Faeces</li> <li>Milk, blood, urine, faeces</li> <li>Blood</li> </ul>	<ul style="list-style-type: none"> <li>Solid</li> <li>New</li> </ul>

Open to collaboration

## Task 6.2 Evaluation of proxies to investigate rumen fermentation parameters : CH<sub>4</sub> emission and VFA, ammonia, pH

Parameter	Proxies	Matrices	Status
<ul style="list-style-type: none"><li>• CH<sub>4</sub> emission</li></ul>	<ul style="list-style-type: none"><li>• MIR</li><li>• NIR</li></ul>	<ul style="list-style-type: none"><li>• Milk</li><li>• Faeces</li></ul>	<ul style="list-style-type: none"><li>• Solid</li><li>• New</li></ul>
<ul style="list-style-type: none"><li>• VFA, ammonia, pH</li></ul>	<ul style="list-style-type: none"><li>• MIR</li><li>• Volatile metabolome</li></ul>	<ul style="list-style-type: none"><li>• Milk</li><li>• Breath gas</li></ul>	

**Open to collaboration**

### Outputs Tasks 6.1 and 6.2

- Improvement of existing prediction models (solid)
- Development of new equations (single or combination)
- Standard Guidelines for using most promising proxies



## WP7

Using sensor data for a multivariate approach to phenotype behavioural traits, health and feed efficiency

Leader **WP7**. Lene Munksgaard (AU)

Task **7.1** Lene Munksgaard (AU)

Task **7.2** Kees Van Reenen (WUR-DLO)

Involved partners AU, INRA, IRTA, WUR-DLO



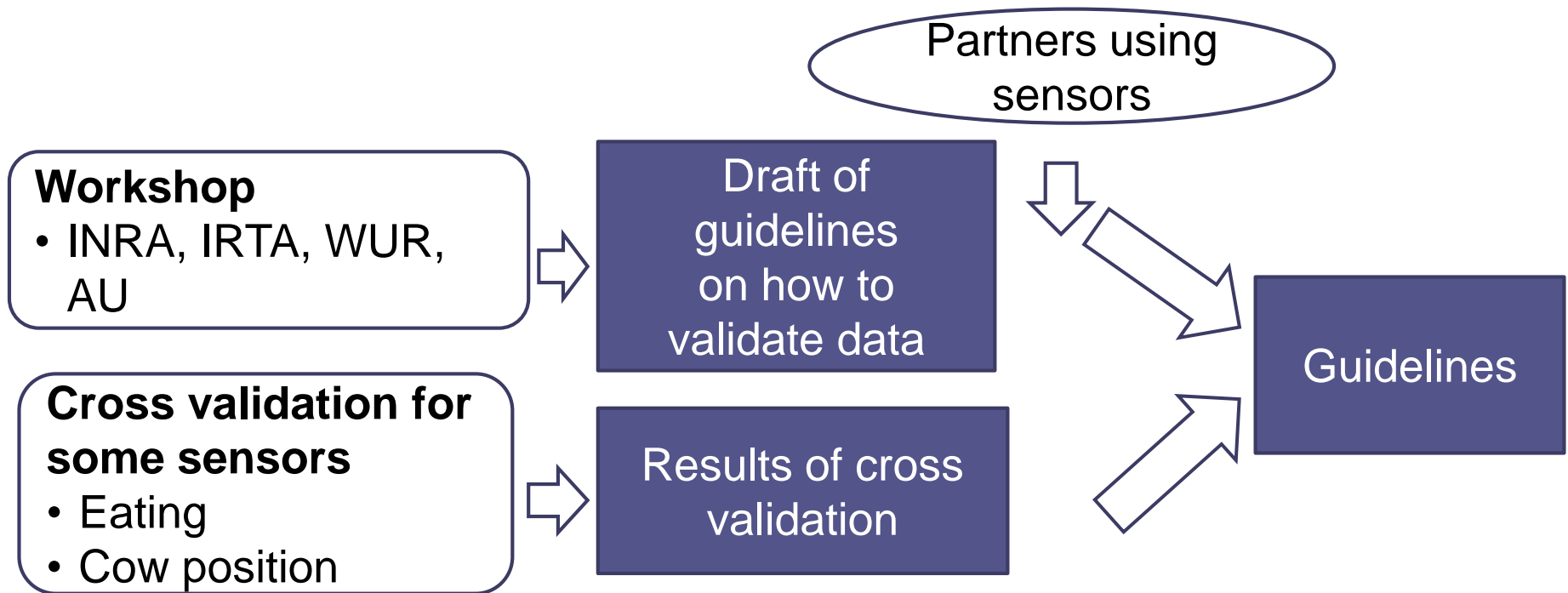
# WP7 Objectives

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- To develop and test **uniform protocols** for validating the quality of data from **smart technologies**
- To develop new ways of **predicting** animals' characteristics based on their **behaviour**
- To implement methods for the analysis of combined data-sets in order to **phenotype cows**



## Task 7.1 Development and test of uniform guidelines for validation of sensors output on animal behaviour

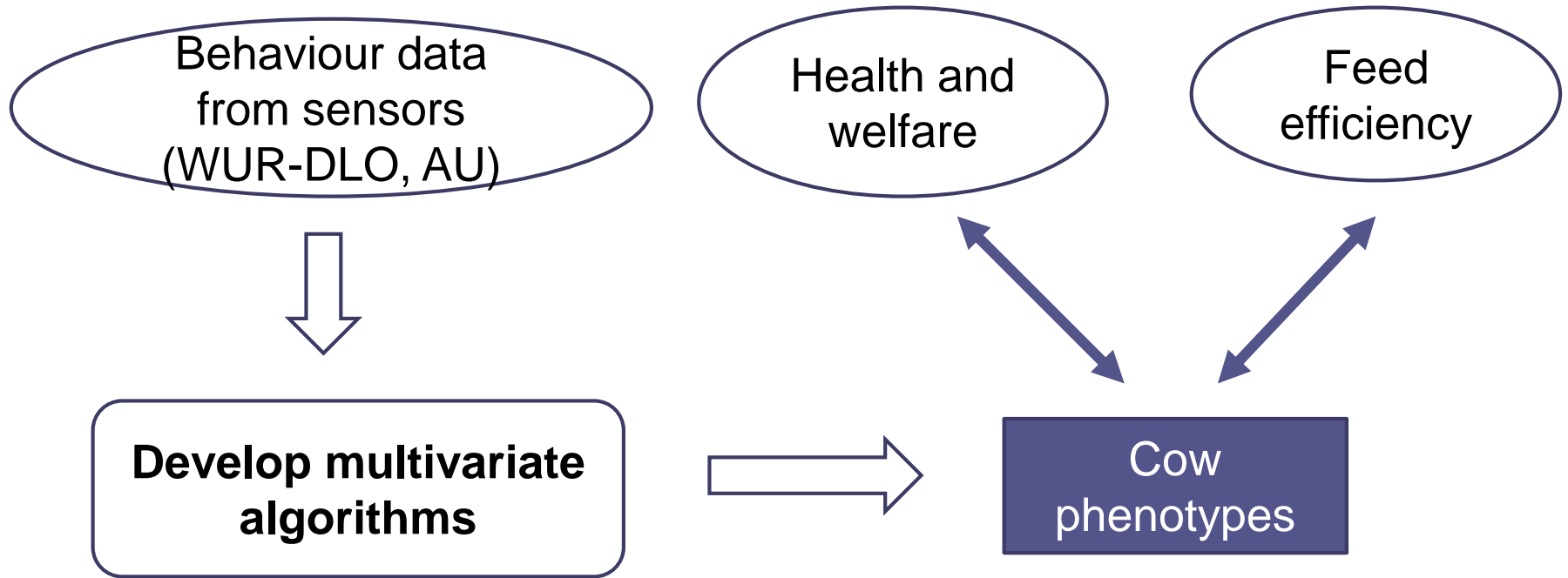


### Outputs

- Uniform methods and guidelines for validating sensors



## Task 7.2 Development of novel algorithms for prediction of complex cow characteristics



### Outputs

- Algorithms for phenotyping cows based on automatic recording of behaviour and the relationship to health, welfare and efficiency





# Thank you for your attention



Further information on: [www.smartcow.eu](http://www.smartcow.eu)

